



Full wwPDB X-ray Structure Validation Report ⓘ

May 16, 2020 – 09:20 pm BST

PDB ID : 4V6F
Title : Elongation complex of the 70S ribosome with three tRNAs and mRNA.
Authors : Jenner, L.B.; Yusupova, G.; Yusupov, M.
Deposited on : 2009-07-09
Resolution : 3.10 Å(reported)

This is a Full wwPDB X-ray Structure Validation Report for a publicly released PDB entry.

We welcome your comments at validation@mail.wwpdb.org

A user guide is available at

<https://www.wwpdb.org/validation/2017/XrayValidationReportHelp>

with specific help available everywhere you see the ⓘ symbol.

The following versions of software and data (see [references ⓘ](#)) were used in the production of this report:

MolProbity : 4.02b-467
Mogul : 1.8.5 (274361), CSD as541be (2020)
Xtriage (Phenix) : 1.13
EDS : 2.11
Percentile statistics : 20191225.v01 (using entries in the PDB archive December 25th 2019)
Refmac : 5.8.0158
CCP4 : 7.0.044 (Gargrove)
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.11

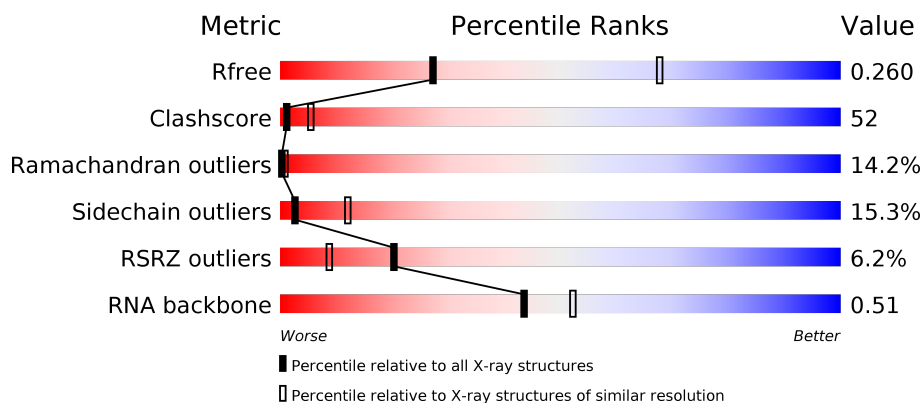
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

The reported resolution of this entry is 3.10 Å.

Percentile scores (ranging between 0-100) for global validation metrics of the entry are shown in the following graphic. The table shows the number of entries on which the scores are based.



Metric	Whole archive (#Entries)	Similar resolution (#Entries, resolution range(Å))
R_{free}	130704	1094 (3.10-3.10)
Clashscore	141614	1184 (3.10-3.10)
Ramachandran outliers	138981	1141 (3.10-3.10)
Sidechain outliers	138945	1141 (3.10-3.10)
RSRZ outliers	127900	1067 (3.10-3.10)
RNA backbone	3102	1116 (3.40-2.80)

The table below summarises the geometric issues observed across the polymeric chains and their fit to the electron density. The red, orange, yellow and green segments on the lower bar indicate the fraction of residues that contain outliers for ≥ 3 , 2, 1 and 0 types of geometric quality criteria respectively. A grey segment represents the fraction of residues that are not modelled. The numeric value for each fraction is indicated below the corresponding segment, with a dot representing fractions $\leq 5\%$. The upper red bar (where present) indicates the fraction of residues that have poor fit to the electron density. The numeric value is given above the bar.

Mol	Chain	Length	Quality of chain
1	AA	2909	<div> <div>3%</div> <div>22%</div> <div>49%</div> <div>24%</div> <div>5%</div> </div>
2	AB	122	<div> <div>2%</div> <div>27%</div> <div>48%</div> <div>24%</div> <div>.</div> </div>
2	DB	122	<div> <div>%</div> <div>34%</div> <div>44%</div> <div>20%</div> <div>.</div> </div>
3	AD	276	<div> <div>%</div> <div>27%</div> <div>53%</div> <div>16%</div> <div>..</div> </div>

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Mol	Chain	Length	Quality of chain
3	DD	276	
4	AE	206	
4	DE	206	
5	AF	210	
5	DF	210	
6	AG	182	
6	DG	182	
7	AH	180	
7	DH	180	
8	AK	148	
8	DK	148	
9	AM	140	
9	DM	140	
10	AN	122	
10	DN	122	
11	AO	150	
11	DO	150	
12	AP	141	
12	DP	141	
13	A0	118	
13	D0	118	
14	AQ	112	
14	DQ	112	
15	AR	146	
15	DR	146	

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Mol	Chain	Length	Quality of chain
16	A1	118	
16	D1	118	
17	A2	101	
17	D2	101	
18	AS	113	
18	DS	113	
19	AT	96	
19	DT	96	
20	AU	110	
20	DU	110	
21	AV	206	
21	DV	206	
22	A3	85	
22	D3	85	
23	AZ	98	
23	DZ	98	
24	AW	72	
24	DW	72	
25	AX	60	
25	DX	60	
26	A4	71	
26	D4	71	
27	A5	60	
27	D5	60	
28	A6	54	

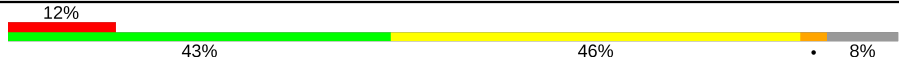
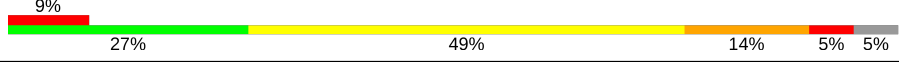
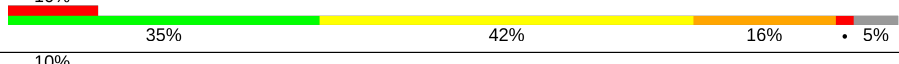
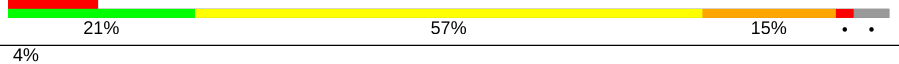
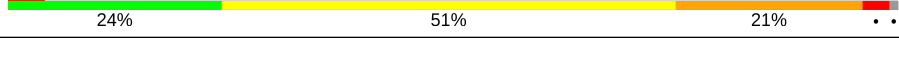
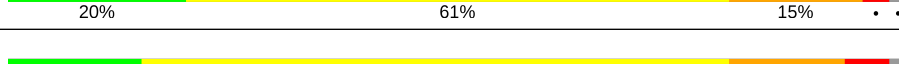
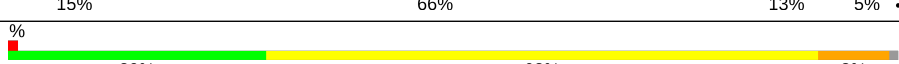
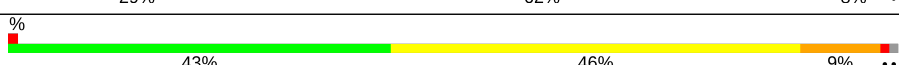
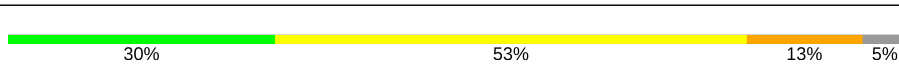

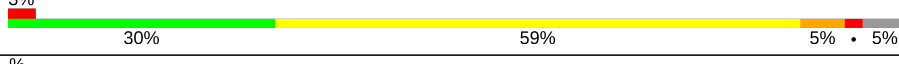
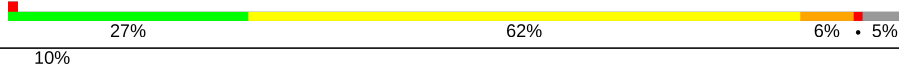
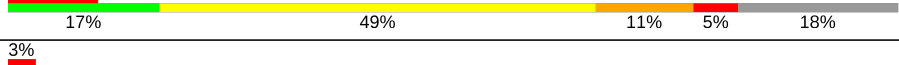

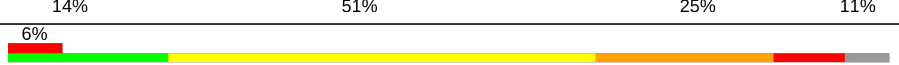
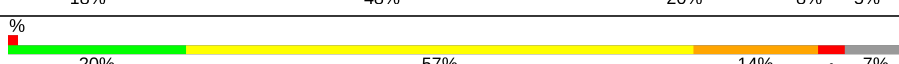
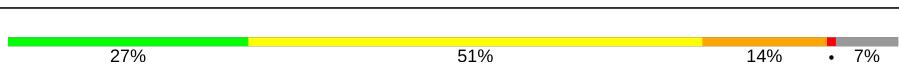
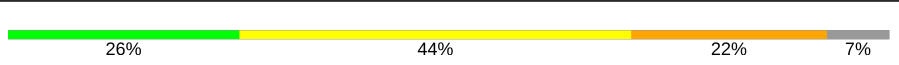
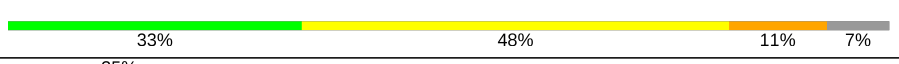
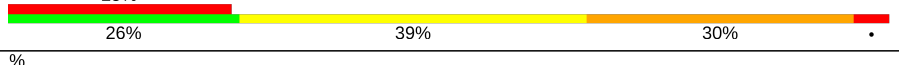
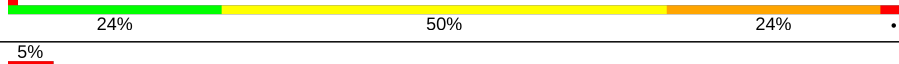

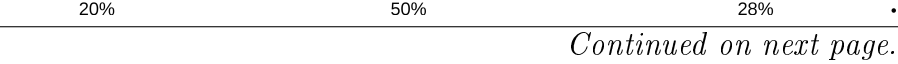


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Mol	Chain	Length	Quality of chain
28	D6	54	
29	A7	49	
29	D7	49	
30	A8	65	
30	D8	65	
31	BA	1516	
32	BE	256	
32	CE	256	
33	BF	239	
33	CF	239	
34	BG	209	
34	CG	209	
35	BH	162	
35	CH	162	
36	BI	101	
36	CI	101	
37	BJ	156	
37	CJ	156	
38	BK	138	
38	CK	138	
39	BL	128	
39	CL	128	
40	BM	105	
40	CM	105	
41	BN	129	

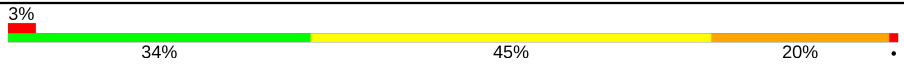
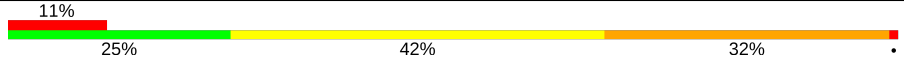
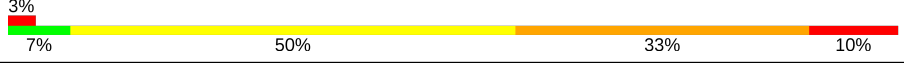

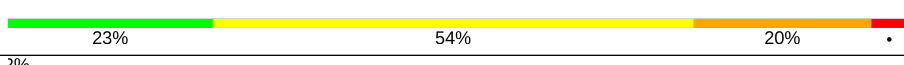
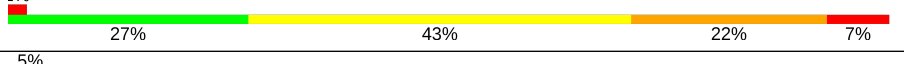
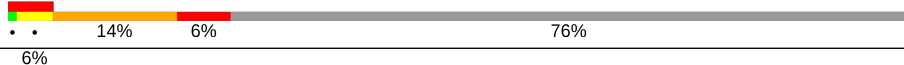


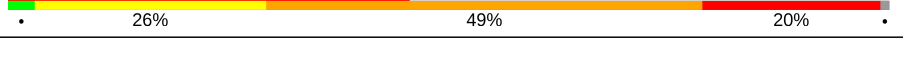
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Mol	Chain	Length	Quality of chain
41	CN	129	
42	BO	132	
42	CO	132	
43	BP	126	
43	CP	126	
44	BQ	61	
44	CQ	61	
45	BR	89	
45	CR	89	
46	BS	88	
46	CS	88	
47	BT	105	
47	CT	105	
48	BU	88	
48	CU	88	
49	BV	93	
49	CV	93	
50	BW	106	
50	CW	106	
51	BX	27	
51	CX	27	
52	BB	76	
52	BC	76	
52	BD	76	
52	CB	76	

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Mol	Chain	Length	Quality of chain
52	CC	76	
52	CD	76	
53	B1	30	
53	C1	30	
54	CA	1515	
55	DA	2912	
56	DI	125	
56	DJ	125	
57	DY	173	
58	DL	147	

The following table lists non-polymeric compounds, carbohydrate monomers and non-standard residues in protein, DNA, RNA chains that are outliers for geometric or electron-density-fit criteria:

Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
59	MG	A7	102	-	-	-	X
59	MG	AA	3116	-	-	-	X
59	MG	AA	3333	-	-	-	X
59	MG	AA	3418	-	-	-	X
59	MG	AA	3423	-	-	-	X
59	MG	AA	3428	-	-	-	X
59	MG	AA	3717	-	-	-	X
59	MG	AA	3719	-	-	-	X
59	MG	AA	3743	-	-	-	X
59	MG	AA	3775	-	-	-	X
59	MG	AA	3852	-	-	-	X
59	MG	AA	3898	-	-	-	X
59	MG	AA	3908	-	-	-	X
59	MG	AA	3922	-	-	-	X
59	MG	AA	3990	-	-	-	X
59	MG	AA	4044	-	-	-	X
59	MG	AA	4075	-	-	-	X
59	MG	AA	4093	-	-	-	X
59	MG	AA	4097	-	-	-	X
59	MG	AA	4107	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
59	MG	AA	4133	-	-	-	X
59	MG	AO	206	-	-	-	X
59	MG	AQ	205	-	-	-	X
59	MG	AS	203	-	-	-	X
59	MG	AW	101	-	-	-	X
59	MG	BA	1612	-	-	-	X
59	MG	BA	1621	-	-	-	X
59	MG	BA	1698	-	-	-	X
59	MG	BA	1772	-	-	-	X
59	MG	BA	1782	-	-	-	X
59	MG	BA	1828	-	-	-	X
59	MG	BA	1859	-	-	-	X
59	MG	BA	1956	-	-	-	X
59	MG	BA	1968	-	-	-	X
59	MG	BA	2068	-	-	-	X
59	MG	BA	2104	-	-	-	X
59	MG	BA	2198	-	-	-	X
59	MG	BA	2213	-	-	-	X
59	MG	BA	2243	-	-	-	X
59	MG	BA	2244	-	-	-	X
59	MG	BA	2276	-	-	-	X
59	MG	BD	126	-	-	-	X
59	MG	CA	1925	-	-	-	X
59	MG	CA	1942	-	-	-	X
59	MG	CA	1946	-	-	-	X
59	MG	CA	1997	-	-	-	X
59	MG	CA	2010	-	-	-	X
59	MG	CA	2065	-	-	-	X
59	MG	CA	2105	-	-	-	X
59	MG	CA	2124	-	-	-	X
59	MG	CA	2304	-	-	-	X
59	MG	CD	104	-	-	-	X
59	MG	CD	108	-	-	-	X
59	MG	CD	118	-	-	-	X
59	MG	CW	201	-	-	-	X
59	MG	D0	203	-	-	-	X
59	MG	D6	103	-	-	-	X
59	MG	DA	3183	-	-	-	X
59	MG	DA	3262	-	-	-	X
59	MG	DA	3274	-	-	-	X
59	MG	DA	3332	-	-	-	X
59	MG	DA	3344	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
59	MG	DA	3367	-	-	-	X
59	MG	DA	3575	-	-	-	X
59	MG	DA	3576	-	-	-	X
59	MG	DA	3598	-	-	-	X
59	MG	DA	3611	-	-	-	X
59	MG	DA	3647	-	-	-	X
59	MG	DA	3676	-	-	-	X
59	MG	DA	3678	-	-	-	X
59	MG	DA	3741	-	-	-	X
59	MG	DA	3757	-	-	-	X
59	MG	DA	3770	-	-	-	X
59	MG	DA	3873	-	-	-	X
59	MG	DA	3984	-	-	-	X
59	MG	DA	4015	-	-	-	X
59	MG	DA	4082	-	-	-	X
59	MG	DA	4185	-	-	-	X
59	MG	DA	4338	-	-	-	X
59	MG	DA	4454	-	-	-	X
59	MG	DA	4474	-	-	-	X
59	MG	DA	4488	-	-	-	X
59	MG	DA	4512	-	-	-	X
59	MG	DA	4545	-	-	-	X
59	MG	DA	4572	-	-	-	X
59	MG	DA	4641	-	-	-	X
59	MG	DA	4668	-	-	-	X
59	MG	DA	4681	-	-	-	X
59	MG	DA	4688	-	-	-	X
59	MG	DA	4698	-	-	-	X
59	MG	DA	4728	-	-	-	X
59	MG	DA	4771	-	-	-	X
59	MG	DA	4779	-	-	-	X
59	MG	DA	4809	-	-	-	X
59	MG	DA	4827	-	-	-	X
59	MG	DA	4872	-	-	-	X
59	MG	DA	4874	-	-	-	X
59	MG	DA	4879	-	-	-	X
59	MG	DA	4886	-	-	-	X
59	MG	DA	4892	-	-	-	X
59	MG	DA	4915	-	-	-	X
59	MG	DA	4929	-	-	-	X
59	MG	DA	4943	-	-	-	X
59	MG	DA	4948	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
59	MG	DA	4956	-	-	-	X
59	MG	DA	4973	-	-	-	X
59	MG	DA	4976	-	-	-	X
59	MG	DA	4980	-	-	-	X
59	MG	DA	4985	-	-	-	X
59	MG	DA	5015	-	-	-	X
59	MG	DA	5053	-	-	-	X
59	MG	DA	5054	-	-	-	X
59	MG	DA	5056	-	-	-	X
59	MG	DA	5058	-	-	-	X
59	MG	DA	5077	-	-	-	X
59	MG	DB	209	-	-	-	X
59	MG	DB	211	-	-	-	X
59	MG	DB	231	-	-	-	X
59	MG	DB	269	-	-	-	X
59	MG	DE	308	-	-	-	X
59	MG	DM	202	-	-	-	X
59	MG	DU	215	-	-	-	X
60	ZN	A4	101	-	-	-	X

2 Entry composition

There are 61 unique types of molecules in this entry. The entry contains 307345 atoms, of which 0 are hydrogens and 0 are deuteriums.

In the tables below, the ZeroOcc column contains the number of atoms modelled with zero occupancy, the AltConf column contains the number of residues with at least one atom in alternate conformation and the Trace column contains the number of residues modelled with at most 2 atoms.

- Molecule 1 is a RNA chain called 23S RIBOSOMAL RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
1	AA	2909	Total	C	N	O	P	0	0	0
			62647	27884	11716	20139	2908			

There are 8 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
AA	?	-	U	DELETION	GB AP008226.1
AA	?	-	U	DELETION	GB AP008226.1
AA	654A	A	G	CONFLICT	GB AP008226.1
AA	654E	C	G	CONFLICT	GB AP008226.1
AA	654P	G	C	CONFLICT	GB AP008226.1
AA	654T	A	C	CONFLICT	GB AP008226.1
AA	1058	U	G	CONFLICT	GB AP008226.1
AA	1080	A	C	CONFLICT	GB AP008226.1

- Molecule 2 is a RNA chain called 5S RIBOSOMAL RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
2	AB	122	Total	C	N	O	P	0	0	0
			2617	1166	486	844	121			
2	DB	122	Total	C	N	O	P	0	0	0
			2617	1166	486	844	121			

There are 2 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
AB	1M	A	-	INSERTION	GB X01554.1
DB	1M	A	-	INSERTION	GB X01554.1

- Molecule 3 is a protein called 50S RIBOSOMAL PROTEIN L2.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
3	AD	272	Total	C	N	O	S	0	0	0
			2115	1335	420	357	3			
3	DD	272	Total	C	N	O	S	0	0	0
			2115	1335	420	357	3			

- Molecule 4 is a protein called 50S RIBOSOMAL PROTEIN L3.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
4	AE	205	Total	C	N	O	S	0	0	0
			1568	991	300	271	6			
4	DE	205	Total	C	N	O	S	0	0	0
			1568	991	300	271	6			

- Molecule 5 is a protein called 50S RIBOSOMAL PROTEIN L4.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
5	AF	208	Total	C	N	O	S	0	0	0
			1627	1037	304	283	3			
5	DF	202	Total	C	N	O	S	0	0	0
			1585	1011	297	275	2			

- Molecule 6 is a protein called 50S RIBOSOMAL PROTEIN L5.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
6	AG	181	Total	C	N	O	S	0	0	0
			1474	942	268	260	4			
6	DG	181	Total	C	N	O	S	0	0	0
			1474	942	268	260	4			

- Molecule 7 is a protein called 50S RIBOSOMAL PROTEIN L6.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
7	AH	170	Total	C	N	O	S	0	0	0
			1307	829	245	232	1			
7	DH	170	Total	C	N	O	S	0	0	0
			1307	829	245	232	1			

- Molecule 8 is a protein called 50S RIBOSOMAL PROTEIN L9.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
8	AK	146	Total	C	N	O	S	0	0	0
			1136	726	201	208	1			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
8	DK	146	Total	C	N	O	S	0	0	0
			1136	726	201	208	1			

- Molecule 9 is a protein called 50S RIBOSOMAL PROTEIN L13.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
9	AM	138	Total	C	N	O	S	0	0	0
			1104	712	206	182	4			
9	DM	138	Total	C	N	O	S	0	0	0
			1104	712	206	182	4			

- Molecule 10 is a protein called 50S RIBOSOMAL PROTEIN L14.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
10	AN	122	Total	C	N	O	S	0	0	0
			933	588	171	170	4			
10	DN	122	Total	C	N	O	S	0	0	0
			933	588	171	170	4			

- Molecule 11 is a protein called 50S RIBOSOMAL PROTEIN L15.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
11	AO	150	Total	C	N	O	S	0	0	0
			1145	712	232	198	3			
11	DO	150	Total	C	N	O	S	0	0	0
			1145	712	232	198	3			

- Molecule 12 is a protein called 50S RIBOSOMAL PROTEIN L16.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
12	AP	141	Total	C	N	O	S	0	0	0
			1122	715	212	188	7			
12	DP	141	Total	C	N	O	S	0	0	0
			1122	715	212	188	7			

- Molecule 13 is a protein called 50S RIBOSOMAL PROTEIN L17.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
13	A0	117	Total	C	N	O		0	0	0
			960	599	202	159				
13	D0	118	Total	C	N	O	S	0	0	0
			968	604	203	160	1			

- Molecule 14 is a protein called 50S RIBOSOMAL PROTEIN L18.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
14	AQ	111	Total	C	N	O	0	0	0
			882	556	176	150			
14	DQ	111	Total	C	N	O	0	0	0
			882	556	176	150			

- Molecule 15 is a protein called 50S RIBOSOMAL PROTEIN L19.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
15	AR	137	Total	C	N	O	S	0	0	0
			1141	710	234	196	1			
15	DR	137	Total	C	N	O	S	0	0	0
			1141	710	234	196	1			

- Molecule 16 is a protein called 50S RIBOSOMAL PROTEIN L20.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
16	A1	117	Total	C	N	O	S	0	0	0
			964	610	202	151	1			
16	D1	117	Total	C	N	O	S	0	0	0
			964	610	202	151	1			

- Molecule 17 is a protein called 50S RIBOSOMAL PROTEIN L21.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
17	A2	101	Total	C	N	O	S	0	0	0
			779	501	142	135	1			
17	D2	101	Total	C	N	O	S	0	0	0
			779	501	142	135	1			

- Molecule 18 is a protein called 50S RIBOSOMAL PROTEIN L22.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
18	AS	113	Total	C	N	O	S	0	0	0
			900	566	177	155	2			
18	DS	113	Total	C	N	O	S	0	0	0
			900	566	177	155	2			

- Molecule 19 is a protein called 50S RIBOSOMAL PROTEIN L23.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
19	AT	92	Total	C	N	O	0	0	0
			725	471	131	123			
19	DT	92	Total	C	N	O	0	0	0
			725	471	131	123			

- Molecule 20 is a protein called 50S RIBOSOMAL PROTEIN L24.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
20	AU	102	Total	C	N	O	S	0	0	0
			785	505	150	125	5			
20	DU	102	Total	C	N	O	S	0	0	0
			785	505	150	125	5			

- Molecule 21 is a protein called 50S RIBOSOMAL PROTEIN L25.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
21	AV	187	Total	C	N	O	S	0	0	0
			1489	949	264	273	3			
21	DV	200	Total	C	N	O	S	0	0	0
			1582	1008	279	292	3			

- Molecule 22 is a protein called 50S RIBOSOMAL PROTEIN L27.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
22	A3	84	Total	C	N	O	S	0	0	0
			662	410	140	111	1			
22	D3	84	Total	C	N	O	S	0	0	0
			662	410	140	111	1			

- Molecule 23 is a protein called 50S RIBOSOMAL PROTEIN L28.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
23	AZ	97	Total	C	N	O	S	0	0	0
			763	481	150	131	1			
23	DZ	97	Total	C	N	O	S	0	0	0
			763	481	150	131	1			

- Molecule 24 is a protein called 50S RIBOSOMAL PROTEIN L29.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
24	AW	69	Total	C	N	O	S	0	0	0
			581	358	118	104	1			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
24	DW	69	Total	C	N	O	S	0	0	0
			581	358	118	104	1			

- Molecule 25 is a protein called 50S RIBOSOMAL PROTEIN L30.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
25	AX	59	Total	C	N	O	S	0	0	0
			469	298	90	81				
25	DX	59	Total	C	N	O	S	0	0	0
			469	298	90	81				

- Molecule 26 is a protein called 50S RIBOSOMAL PROTEIN L31.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
26	A4	71	Total	C	N	O	S	0	0	0
			581	364	108	104	5			
26	D4	71	Total	C	N	O	S	0	0	0
			581	364	108	104	5			

- Molecule 27 is a protein called 50S RIBOSOMAL PROTEIN L32.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
27	A5	59	Total	C	N	O	S	0	0	0
			459	288	90	76	5			
27	D5	59	Total	C	N	O	S	0	0	0
			459	288	90	76	5			

- Molecule 28 is a protein called 50S RIBOSOMAL PROTEIN L33.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
28	A6	45	Total	C	N	O	S	0	0	0
			389	241	79	65	4			
28	D6	45	Total	C	N	O	S	0	0	0
			389	241	79	65	4			

- Molecule 29 is a protein called 50S RIBOSOMAL PROTEIN L34.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
29	A7	49	Total	C	N	O	S	0	0	0
			430	263	108	57	2			
29	D7	49	Total	C	N	O	S	0	0	0
			430	263	108	57	2			

- Molecule 30 is a protein called 50S RIBOSOMAL PROTEIN L35.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
30	A8	64	Total	C	N	O	S	0	0	0
			517	331	102	82	2			
30	D8	64	Total	C	N	O	S	0	0	0
			517	331	102	82	2			

- Molecule 31 is a RNA chain called 16S ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
31	BA	1516	Total	C	N	O	P	0	0	0
			32571	14499	6024	10533	1515			

- Molecule 32 is a protein called 30S ribosomal protein S2.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
32	BE	237	Total	C	N	O	S	0	0	0
			1924	1228	344	347	5			
32	CE	237	Total	C	N	O	S	0	0	0
			1924	1228	344	347	5			

- Molecule 33 is a protein called 30S ribosomal protein S3.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
33	BF	206	Total	C	N	O	S	0	0	0
			1612	1016	314	281	1			
33	CF	205	Total	C	N	O	S	0	0	0
			1605	1011	313	280	1			

- Molecule 34 is a protein called 30S ribosomal protein S4.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
34	BG	208	Total	C	N	O	S	0	0	0
			1703	1066	339	291	7			
34	CG	208	Total	C	N	O	S	0	0	0
			1703	1066	339	291	7			

- Molecule 35 is a protein called 30S ribosomal protein S5.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
35	BH	151	Total	C	N	O	S	0	0	0
			1155	729	218	204	4			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
35	CH	151	Total	C	N	O	S	0	0	0
			1155	729	218	204	4			

- Molecule 36 is a protein called 30S ribosomal protein S6.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
36	BI	101	Total	C	N	O	S	0	0	0
			843	531	155	154	3			
36	CI	101	Total	C	N	O	S	0	0	0
			843	531	155	154	3			

- Molecule 37 is a protein called 30S ribosomal protein S7.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
37	BJ	155	Total	C	N	O	S	0	0	0
			1257	781	252	218	6			
37	CJ	155	Total	C	N	O	S	0	0	0
			1257	781	252	218	6			

- Molecule 38 is a protein called 30S ribosomal protein S8.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
38	BK	138	Total	C	N	O	S	0	0	0
			1116	705	215	193	3			
38	CK	138	Total	C	N	O	S	0	0	0
			1116	705	215	193	3			

- Molecule 39 is a protein called 30S ribosomal protein S9.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
39	BL	127	Total	C	N	O	0	0	0
			1010	639	197	174			
39	CL	127	Total	C	N	O	0	0	0
			1010	639	197	174			

- Molecule 40 is a protein called 30S ribosomal protein S10.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
40	BM	99	Total	C	N	O	S	0	0	0
			801	504	157	139	1			
40	CM	99	Total	C	N	O	S	0	0	0
			801	504	157	139	1			

- Molecule 41 is a protein called 30S ribosomal protein S11.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
41	BN	119	Total	C	N	O	S	0	0	0
			885	549	168	165	3			
41	CN	119	Total	C	N	O	S	0	0	0
			885	549	168	165	3			

- Molecule 42 is a protein called 30S ribosomal protein S12.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
42	BO	125	Total	C	N	O	S	0	0	0
			975	614	196	164	1			
42	CO	125	Total	C	N	O	S	0	0	0
			975	614	196	164	1			

- Molecule 43 is a protein called 30S ribosomal protein S13.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
43	BP	121	Total	C	N	O	S	0	0	0
			964	597	199	166	2			
43	CP	125	Total	C	N	O	S	0	0	0
			997	617	207	171	2			

- Molecule 44 is a protein called 30S ribosomal protein S14 type Z.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
44	BQ	60	Total	C	N	O	S	0	0	0
			492	312	104	72	4			
44	CQ	60	Total	C	N	O	S	0	0	0
			492	312	104	72	4			

- Molecule 45 is a protein called 30S ribosomal protein S15.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
45	BR	88	Total	C	N	O	S	0	0	0
			734	459	147	126	2			
45	CR	88	Total	C	N	O	S	0	0	0
			734	459	147	126	2			

- Molecule 46 is a protein called 30S ribosomal protein S16.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
46	BS	84	Total	C	N	O	S	0	0	0
			705	446	140	118	1			
46	CS	84	Total	C	N	O	S	0	0	0
			705	446	140	118	1			

- Molecule 47 is a protein called 30S ribosomal protein S17.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
47	BT	100	Total	C	N	O	S	0	0	0
			834	534	155	143	2			
47	CT	100	Total	C	N	O	S	0	0	0
			834	534	155	143	2			

- Molecule 48 is a protein called 30S ribosomal protein S18.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
48	BU	72	Total	C	N	O		0	0	0
			591	376	117	98				
48	CU	72	Total	C	N	O		0	0	0
			591	376	117	98				

- Molecule 49 is a protein called 30S ribosomal protein S19.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
49	BV	83	Total	C	N	O	S	0	0	0
			665	424	122	117	2			
49	CV	88	Total	C	N	O	S	0	0	0
			702	447	131	122	2			

- Molecule 50 is a protein called 30S ribosomal protein S20.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
50	BW	99	Total	C	N	O	S	0	0	0
			763	470	162	129	2			
50	CW	99	Total	C	N	O	S	0	0	0
			763	470	162	129	2			

- Molecule 51 is a protein called 30S ribosomal protein Thx.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
51	BX	25	Total	C	N	O	0	0	0
			217	134	52	31			

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Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
51	CX	25	Total	C	N	O	0	0	0
			217	134	52	31			

- Molecule 52 is a RNA chain called TRNA-PHE WITH UNMODIFIED NUCLEOTIDES EXCEPT FOR MIA37.

Mol	Chain	Residues	Atoms						ZeroOcc	AltConf	Trace
52	BD	76	Total	C	N	O	P	S	0	0	0
			1626	729	290	531	75	1			
52	BB	76	Total	C	N	O	P	S	0	0	0
			1626	729	290	531	75	1			
52	BC	76	Total	C	N	O	P	S	0	0	0
			1626	729	290	531	75	1			
52	CD	76	Total	C	N	O	P	S	0	0	0
			1626	729	290	531	75	1			
52	CB	76	Total	C	N	O	P	S	0	0	0
			1626	729	290	531	75	1			
52	CC	76	Total	C	N	O	P	S	0	0	0
			1626	729	290	531	75	1			

- Molecule 53 is a RNA chain called MRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
53	B1	30	Total	C	N	O	P	0	0	0
			621	279	88	225	29			
53	C1	30	Total	C	N	O	P	0	0	0
			621	279	88	225	29			

- Molecule 54 is a RNA chain called 16S ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
54	CA	1515	Total	C	N	O	P	0	0	0
			32551	14490	6022	10525	1514			

- Molecule 55 is a RNA chain called 23S RRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
55	DA	2912	Total	C	N	O	P	0	0	0
			62707	27911	11722	20163	2911			

There are 7 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
DA	161	U	-	INSERTION	GB AP008226.1
DA	654A	A	G	CONFLICT	GB AP008226.1
DA	654E	C	G	CONFLICT	GB AP008226.1
DA	654P	G	C	CONFLICT	GB AP008226.1
DA	654T	A	C	CONFLICT	GB AP008226.1
DA	1058	U	G	CONFLICT	GB AP008226.1
DA	1080	A	C	CONFLICT	GB AP008226.1

- Molecule 56 is a protein called 50S ribosomal protein L7/L12.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
56	DI	30	Total	C	N	O	S	0	0	0
			237	150	38	48	1			
56	DJ	30	Total	C	N	O	S	0	0	0
			237	150	38	48	1			

- Molecule 57 is a protein called 50S ribosomal protein L10.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
57	DY	145	Total	C	N	O	S	0	0	0
			1107	708	193	204	2			

- Molecule 58 is a protein called 50S ribosomal protein L11.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
58	DL	145	Total	C	N	O	S	0	0	0
			1071	681	188	197	5			

- Molecule 59 is MAGNESIUM ION (three-letter code: MG) (formula: Mg).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	CA	724	Total	Mg	0	0
			724	724		
59	AB	36	Total	Mg	0	0
			36	36		
59	CV	4	Total	Mg	0	0
			4	4		
59	DO	18	Total	Mg	0	0
			18	18		
59	AW	2	Total	Mg	0	0
			2	2		
59	DZ	5	Total	Mg	0	0
			5	5		

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Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
59	BI	1	Total Mg 1 1	0	0
59	BT	2	Total Mg 2 2	0	0
59	D3	7	Total Mg 7 7	0	0
59	AA	1166	Total Mg 1166 1166	0	0
59	CQ	3	Total Mg 3 3	0	0
59	AR	5	Total Mg 5 5	0	0
59	BC	16	Total Mg 16 16	0	0
59	CJ	1	Total Mg 1 1	0	0
59	D4	2	Total Mg 2 2	0	0
59	DE	15	Total Mg 15 15	0	0
59	AQ	5	Total Mg 5 5	0	0
59	DP	4	Total Mg 4 4	0	0
59	BS	9	Total Mg 9 9	0	0
59	CE	6	Total Mg 6 6	0	0
59	A3	4	Total Mg 4 4	0	0
59	AF	7	Total Mg 7 7	0	0
59	DK	2	Total Mg 2 2	0	0
59	AK	3	Total Mg 3 3	0	0
59	DF	25	Total Mg 25 25	0	0
59	BE	5	Total Mg 5 5	0	0
59	DU	19	Total Mg 19 19	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	BP	2	Total 2	Mg 2	0	0
59	AE	4	Total 4	Mg 4	0	0
59	DL	1	Total 1	Mg 1	0	0
59	AV	1	Total 1	Mg 1	0	0
59	DV	5	Total 5	Mg 5	0	0
59	BU	1	Total 1	Mg 1	0	0
59	CN	2	Total 2	Mg 2	0	0
59	D0	11	Total 11	Mg 11	0	0
59	CC	27	Total 27	Mg 27	0	0
59	CP	3	Total 3	Mg 3	0	0
59	DA	2077	Total 2077	Mg 2077	0	0
59	AU	6	Total 6	Mg 6	0	0
59	BO	1	Total 1	Mg 1	0	0
59	CI	1	Total 1	Mg 1	0	0
59	A7	4	Total 4	Mg 4	0	0
59	D5	8	Total 8	Mg 8	0	0
59	A8	4	Total 4	Mg 4	0	0
59	AO	7	Total 7	Mg 7	0	0
59	CS	6	Total 6	Mg 6	0	0
59	DB	76	Total 76	Mg 76	0	0
59	AP	1	Total 1	Mg 1	0	0

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Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
59	BA	676	Total Mg 676 676	0	0
59	DQ	2	Total Mg 2 2	0	0
59	BL	2	Total Mg 2 2	0	0
59	C1	6	Total Mg 6 6	0	0
59	CD	30	Total Mg 30 30	0	0
59	A2	1	Total Mg 1 1	0	0
59	D6	3	Total Mg 3 3	0	0
59	DH	5	Total Mg 5 5	0	0
59	DG	5	Total Mg 5 5	0	0
59	BF	2	Total Mg 2 2	0	0
59	DR	4	Total Mg 4 4	0	0
59	BQ	3	Total Mg 3 3	0	0
59	CG	11	Total Mg 11 11	0	0
59	A1	5	Total Mg 5 5	0	0
59	AD	13	Total Mg 13 13	0	0
59	CT	3	Total Mg 3 3	0	0
59	DM	6	Total Mg 6 6	0	0
59	DX	2	Total Mg 2 2	0	0
59	AZ	3	Total Mg 3 3	0	0
59	BK	6	Total Mg 6 6	0	0
59	DW	7	Total Mg 7 7	0	0

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Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
59	BV	1	Total Mg 1 1	0	0
59	CM	3	Total Mg 3 3	0	0
59	D1	12	Total Mg 12 12	0	0
59	CB	21	Total Mg 21 21	0	0
59	CW	1	Total Mg 1 1	0	0
59	DN	2	Total Mg 2 2	0	0
59	AT	4	Total Mg 4 4	0	0
59	BH	5	Total Mg 5 5	0	0
59	CH	6	Total Mg 6 6	0	0
59	A6	2	Total Mg 2 2	0	0
59	D2	10	Total Mg 10 10	0	0
59	B1	4	Total Mg 4 4	0	0
59	AN	1	Total Mg 1 1	0	0
59	CR	3	Total Mg 3 3	0	0
59	AS	3	Total Mg 3 3	0	0
59	BB	13	Total Mg 13 13	0	0
59	BM	3	Total Mg 3 3	0	0
59	BX	1	Total Mg 1 1	0	0
59	D8	9	Total Mg 9 9	0	0
59	CK	11	Total Mg 11 11	0	0
59	A5	3	Total Mg 3 3	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	D7	5	Total 5	Mg 5	0	0
59	DD	14	Total 14	Mg 14	0	0
59	BG	7	Total 7	Mg 7	0	0
59	DS	8	Total 8	Mg 8	0	0
59	CF	3	Total 3	Mg 3	0	0
59	A0	1	Total 1	Mg 1	0	0
59	AG	3	Total 3	Mg 3	0	0
59	AH	1	Total 1	Mg 1	0	0
59	DY	4	Total 4	Mg 4	0	0
59	BD	26	Total 26	Mg 26	0	0
59	DT	7	Total 7	Mg 7	0	0
59	BW	8	Total 8	Mg 8	0	0
59	CL	1	Total 1	Mg 1	0	0

- Molecule 60 is ZINC ION (three-letter code: ZN) (formula: Zn).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
60	BG	1	Total 1	Zn 1	0	0
60	BQ	1	Total 1	Zn 1	0	0
60	CQ	1	Total 1	Zn 1	0	0
60	A4	1	Total 1	Zn 1	0	0
60	CG	1	Total 1	Zn 1	0	0
60	D4	1	Total 1	Zn 1	0	0

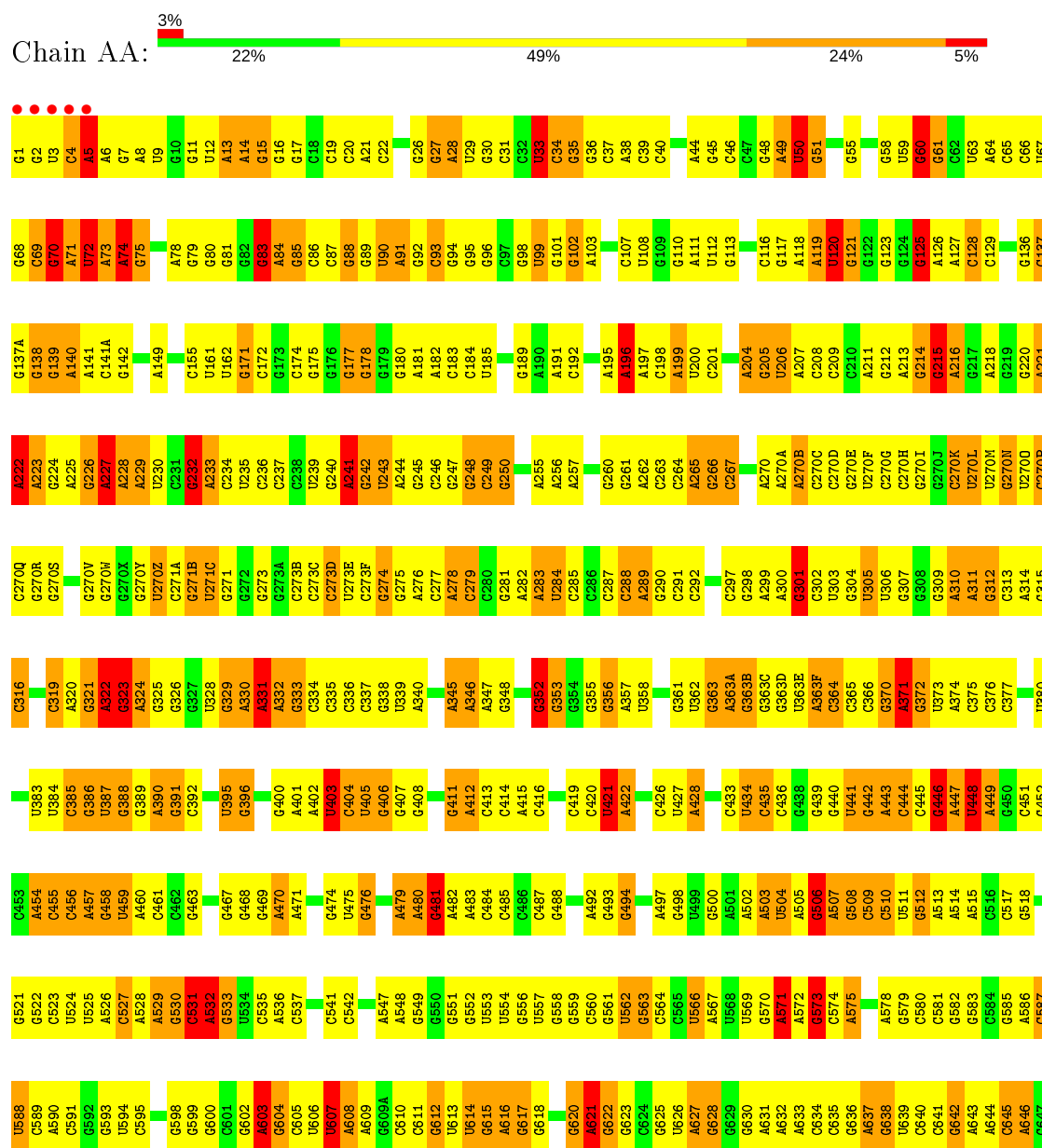
- Molecule 61 is water.

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
61	BA	3	Total 3	O 3	0	0
61	B1	3	Total 3	O 3	0	0

3 Residue-property plots

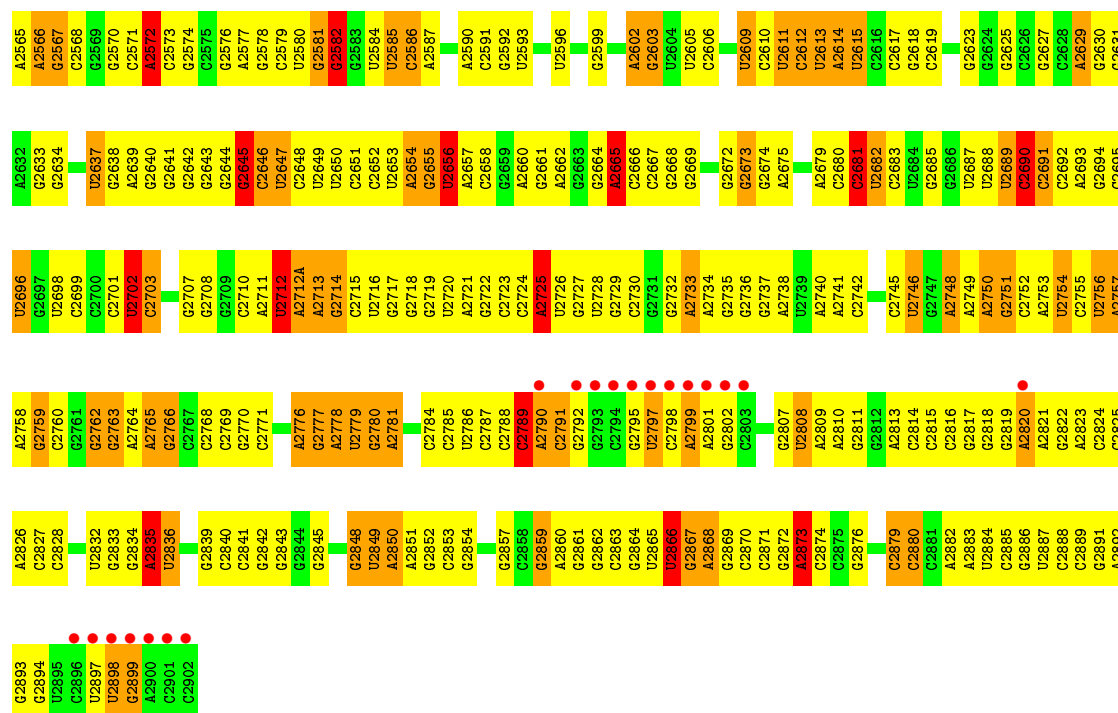
These plots are drawn for all protein, RNA and DNA chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and electron density. Residues are color-coded according to the number of geometric quality criteria for which they contain at least one outlier: green = 0, yellow = 1, orange = 2 and red = 3 or more. A red dot above a residue indicates a poor fit to the electron density ($RSRZ > 2$). Stretches of 2 or more consecutive residues without any outlier are shown as a green connector. Residues present in the sample, but not in the model, are shown in grey.

• Molecule 1: 23S RIBOSOMAL RNA

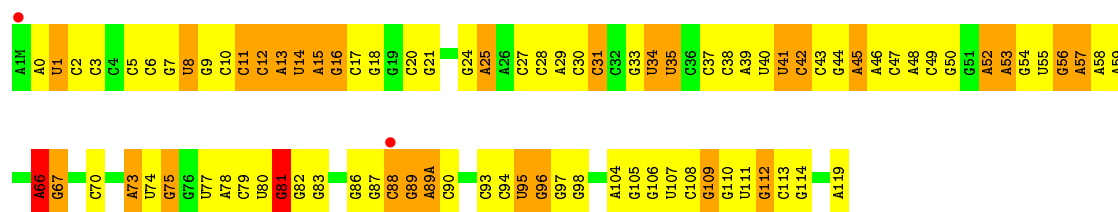


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G1465	G1401	G1338	A1275	G1151	G1151	A1088	A1020	U958	C894	A829	G763	C698	C650
G1466	C1402	G1339	A1276	A1214	C1152	G1089	A1021	A959	U895	G830	A764	C698	G651
C1467	C1403	U1340	G1277	G1215	C1153	U1090	G1022	A960	A896	G831			
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U1405	U1405	A1342	G1279	A1155	A1155	G1092	G1024	G962	C898	U833	G769		A654A
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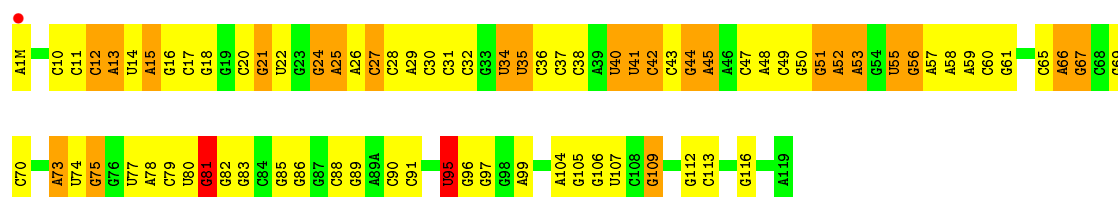
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• Molecule 2: 5S RIBOSOMAL RNA

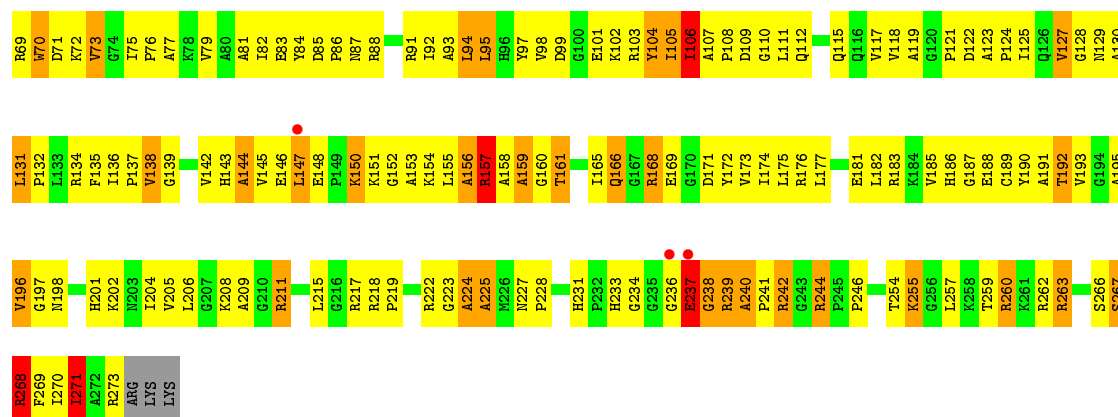


• Molecule 2: 5S RIBOSOMAL RNA

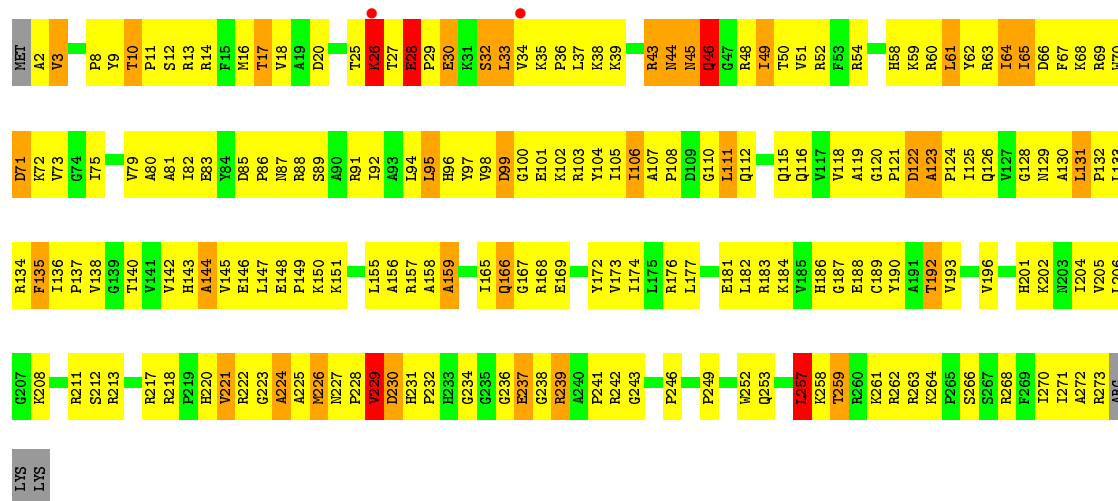


• Molecule 3: 50S RIBOSOMAL PROTEIN L2

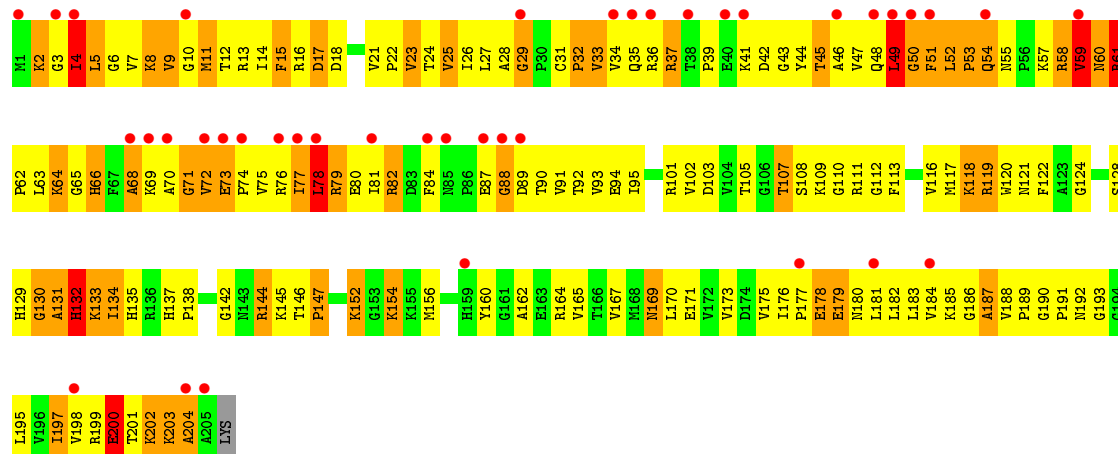




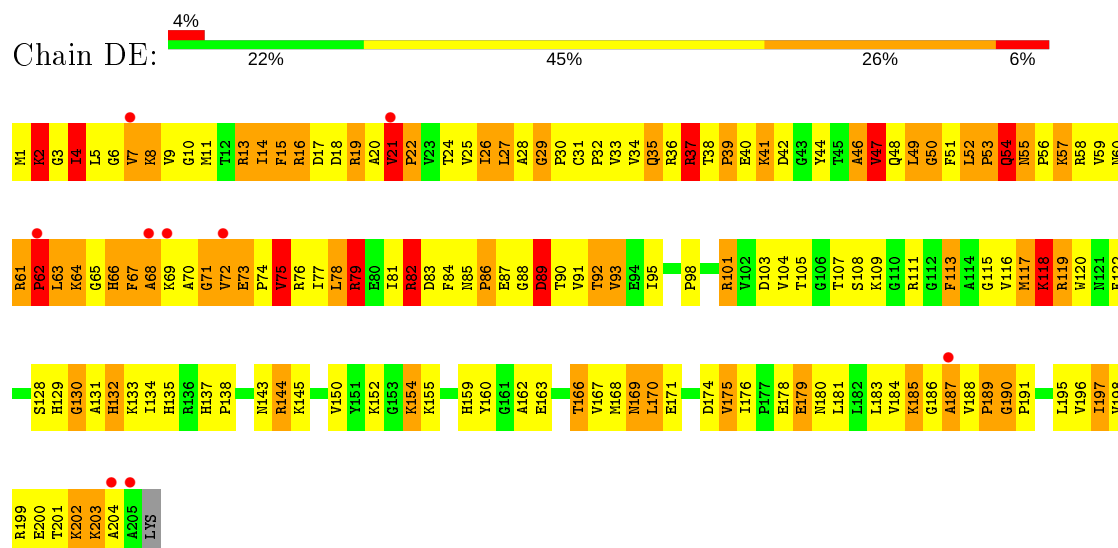
• Molecule 3: 50S RIBOSOMAL PROTEIN L2



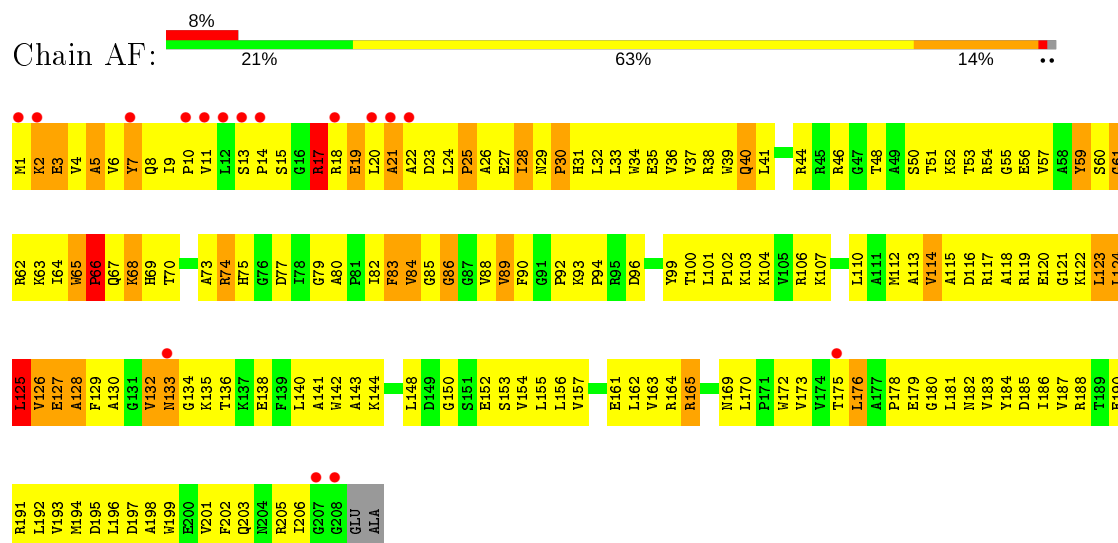
• Molecule 4: 50S RIBOSOMAL PROTEIN L3



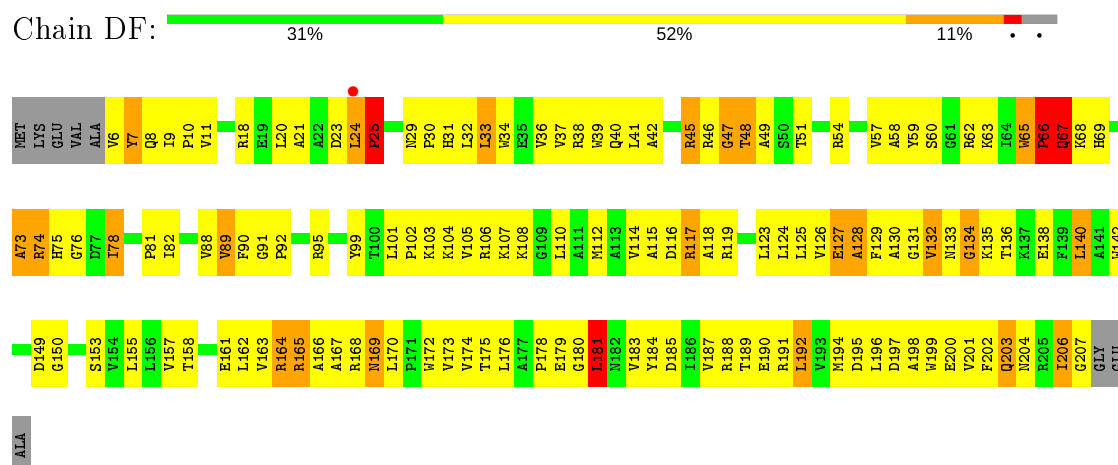
- Molecule 4: 50S RIBOSOMAL PROTEIN L3



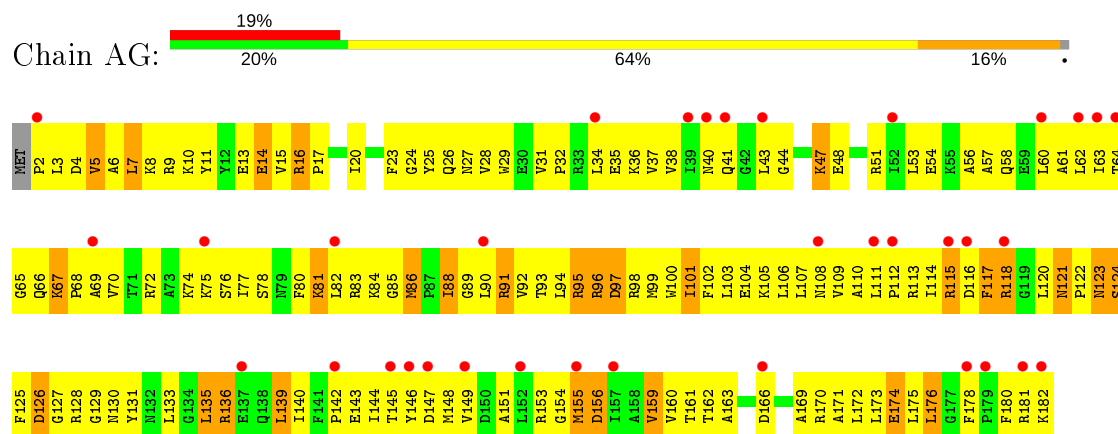
• Molecule 5: 50S RIBOSOMAL PROTEIN L4

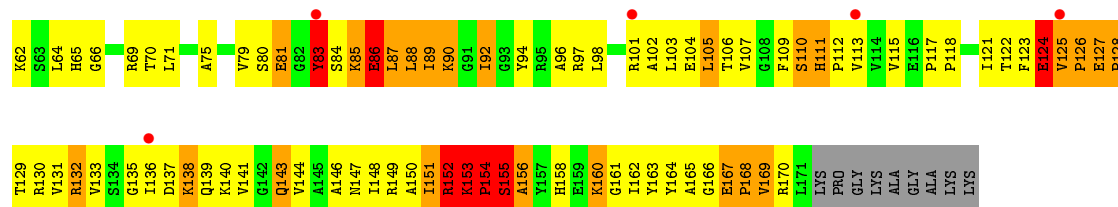


• Molecule 5: 50S RIBOSOMAL PROTEIN L4

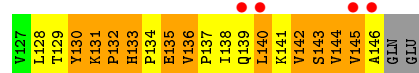
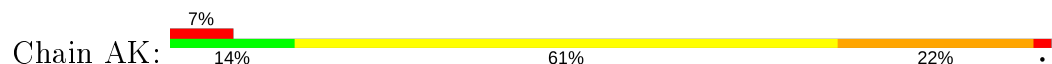


● Molecule 6: 50S RIBOSOMAL PROTEIN L5

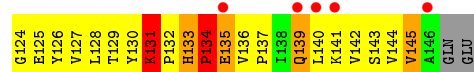
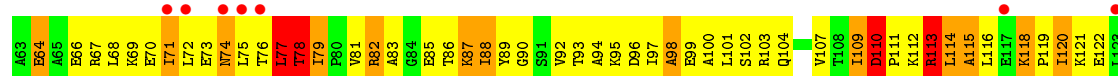
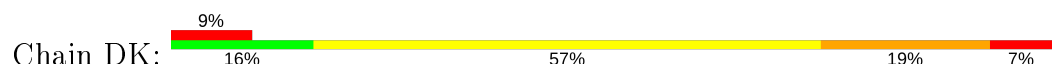




• Molecule 8: 50S RIBOSOMAL PROTEIN L9



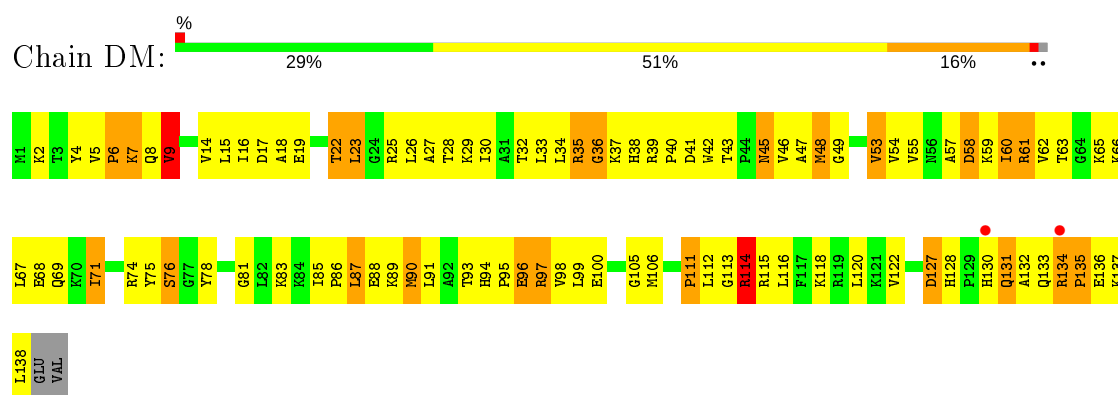
• Molecule 8: 50S RIBOSOMAL PROTEIN L9



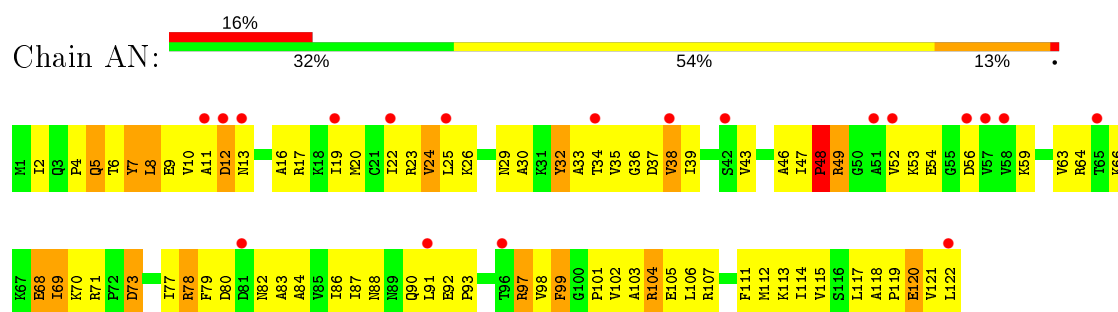
• Molecule 9: 50S RIBOSOMAL PROTEIN L13



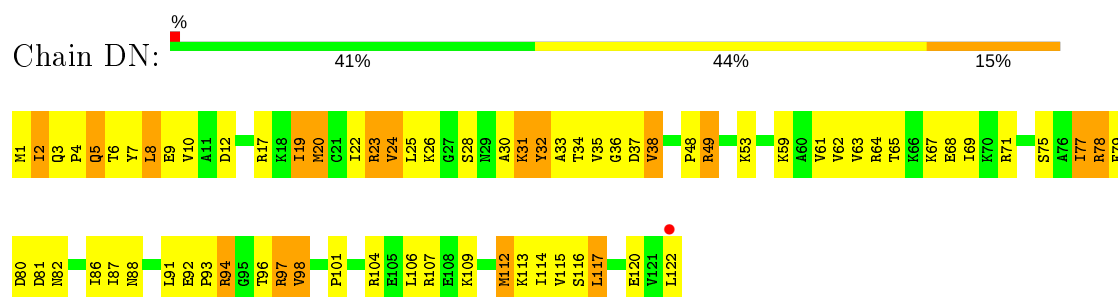
• Molecule 9: 50S RIBOSOMAL PROTEIN L13



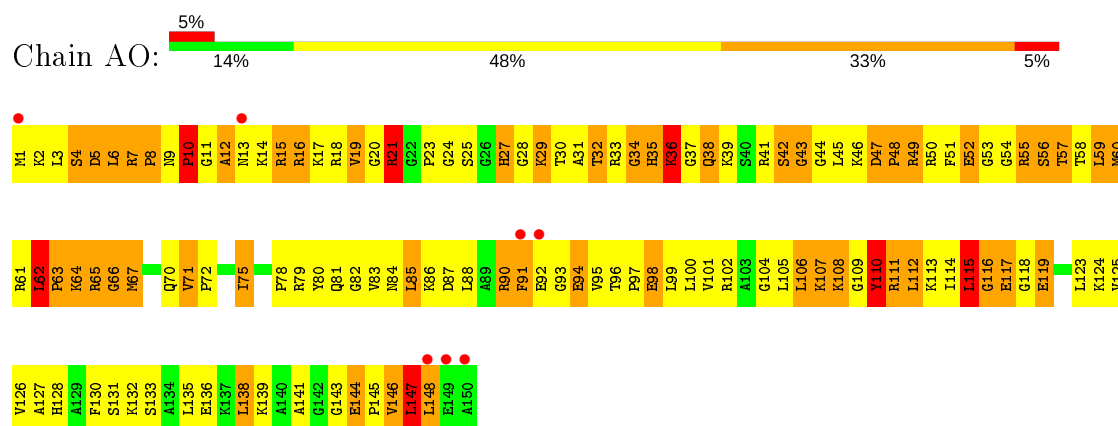
● Molecule 10: 50S RIBOSOMAL PROTEIN L14



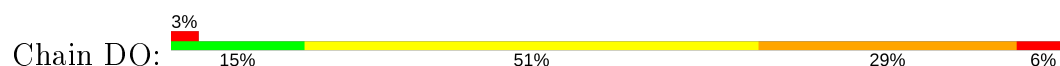
- Molecule 10: 50S RIBOSOMAL PROTEIN L14

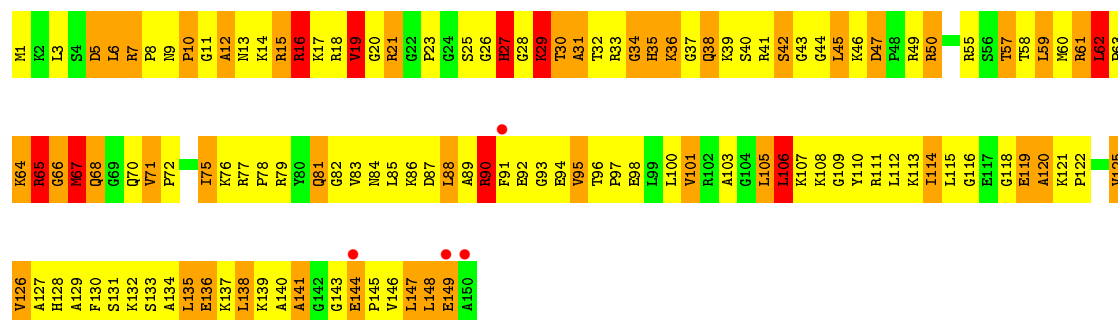


- Molecule 11: 50S RIBOSOMAL PROTEIN L15

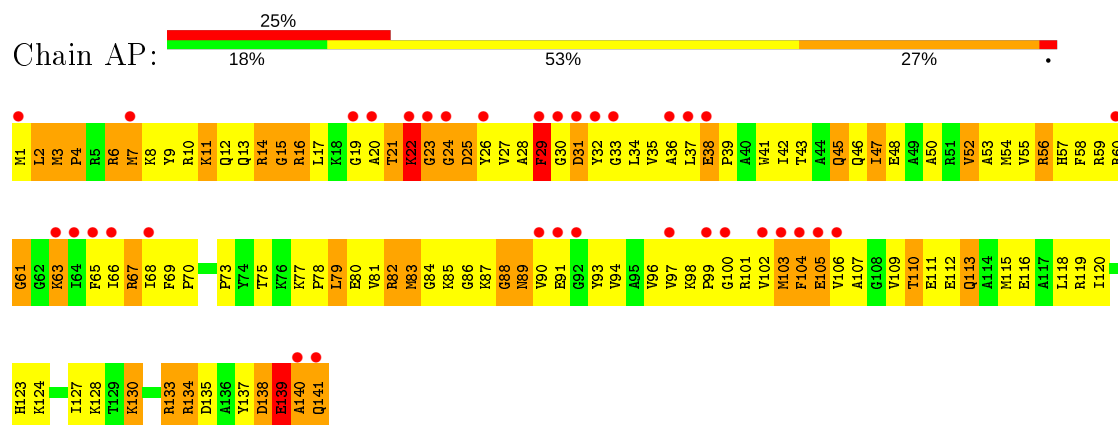


- Molecule 11: 50S RIBOSOMAL PROTEIN L15

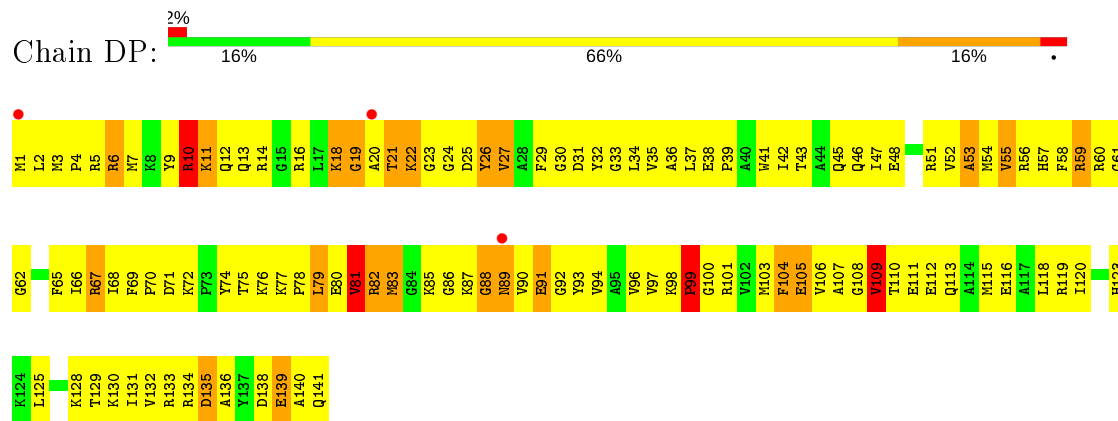




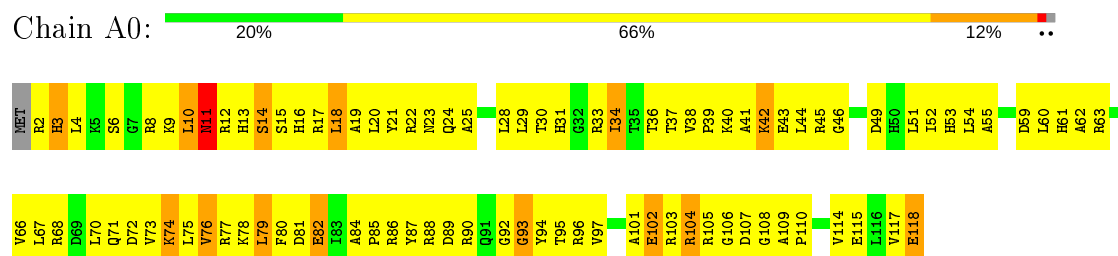
• Molecule 12: 50S RIBOSOMAL PROTEIN L16



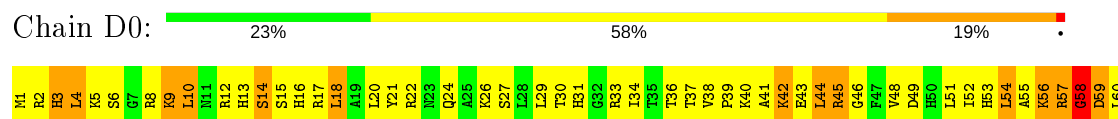
• Molecule 12: 50S RIBOSOMAL PROTEIN L16



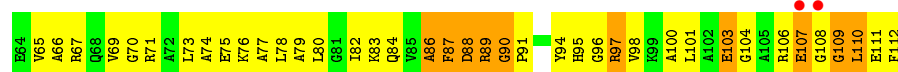
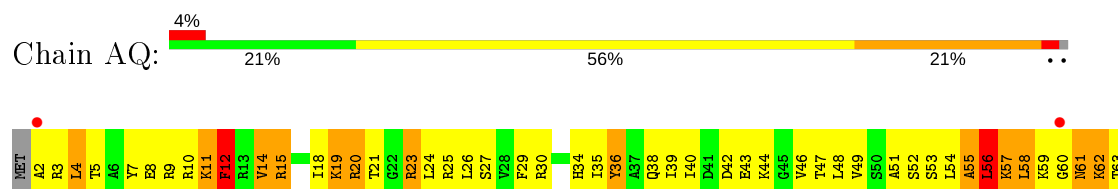
• Molecule 13: 50S RIBOSOMAL PROTEIN L17



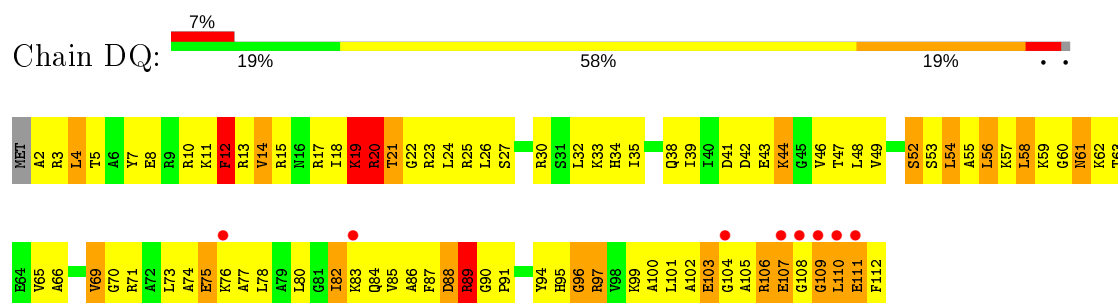
• Molecule 13: 50S RIBOSOMAL PROTEIN L17



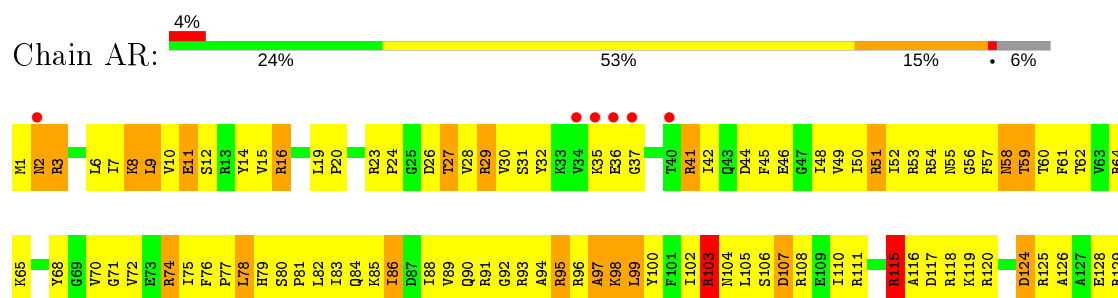
• Molecule 14: 50S RIBOSOMAL PROTEIN L18



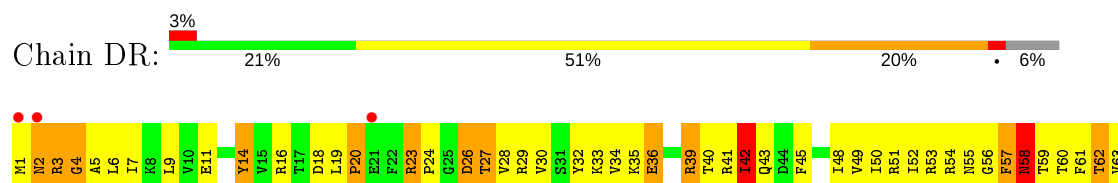
• Molecule 14: 50S RIBOSOMAL PROTEIN L18

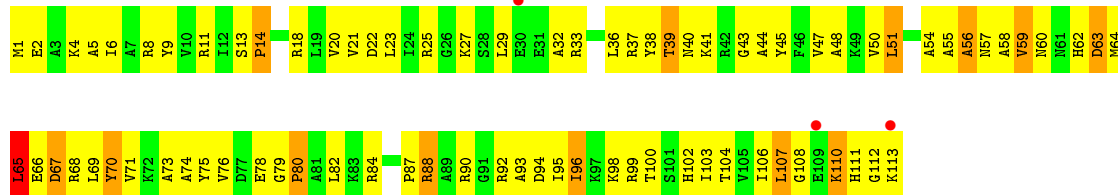


• Molecule 15: 50S RIBOSOMAL PROTEIN L19

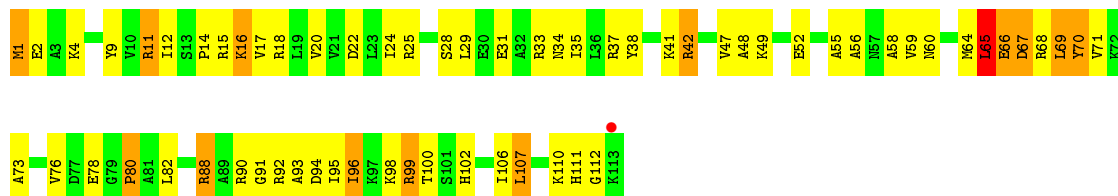


• Molecule 15: 50S RIBOSOMAL PROTEIN L19

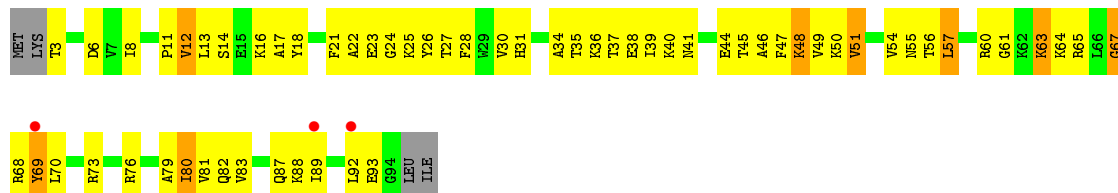




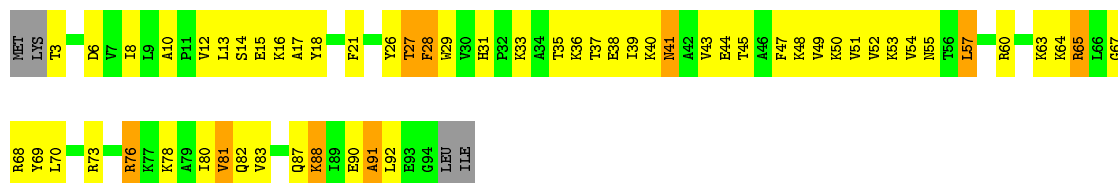
• Molecule 18: 50S RIBOSOMAL PROTEIN L22



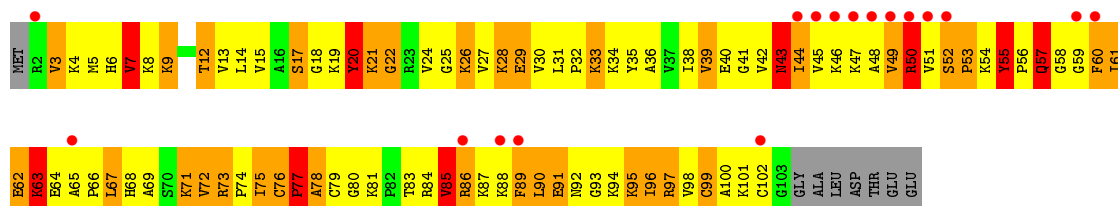
• Molecule 19: 50S RIBOSOMAL PROTEIN L23



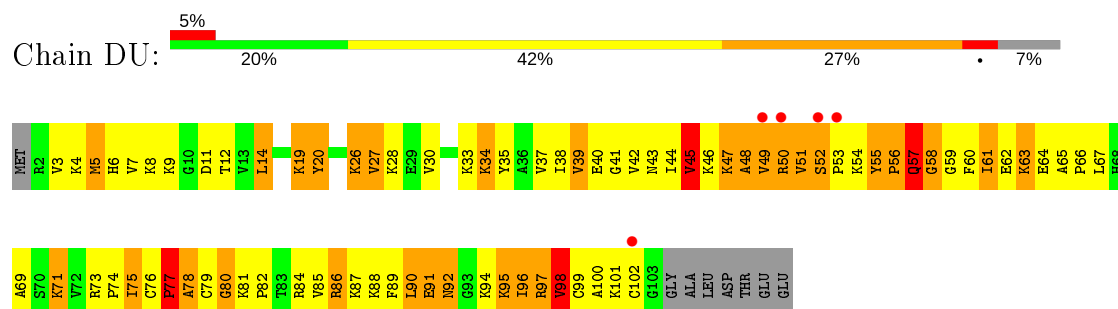
• Molecule 19: 50S RIBOSOMAL PROTEIN L23



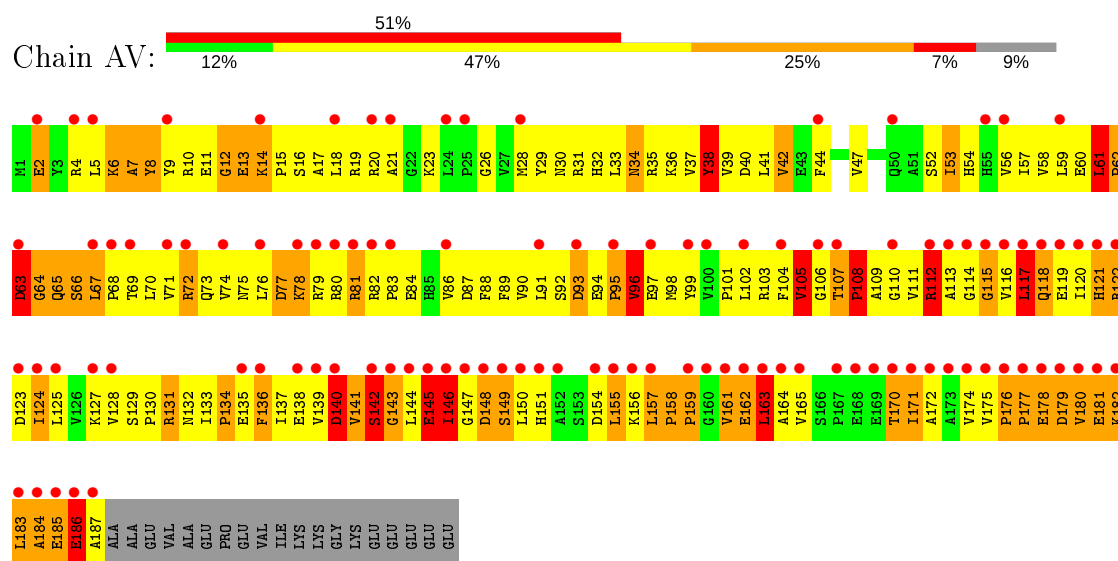
• Molecule 20: 50S RIBOSOMAL PROTEIN L24



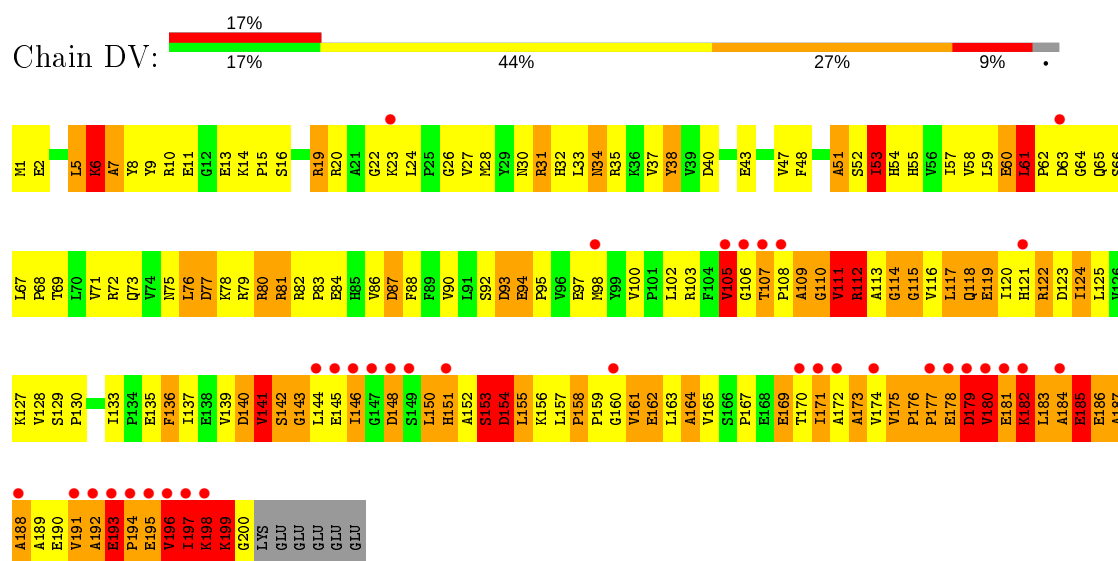
- Molecule 20: 50S RIBOSOMAL PROTEIN L24



- Molecule 21: 50S RIBOSOMAL PROTEIN L25

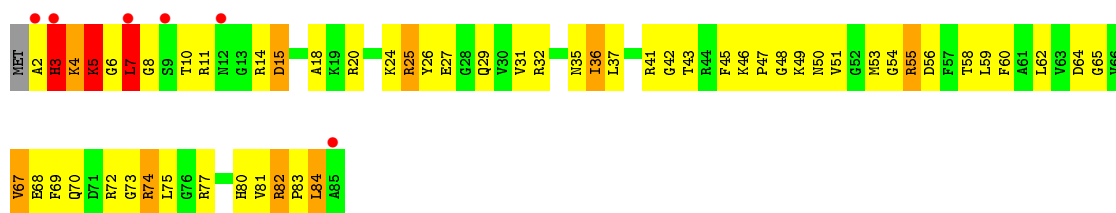


- Molecule 21: 50S RIBOSOMAL PROTEIN L25

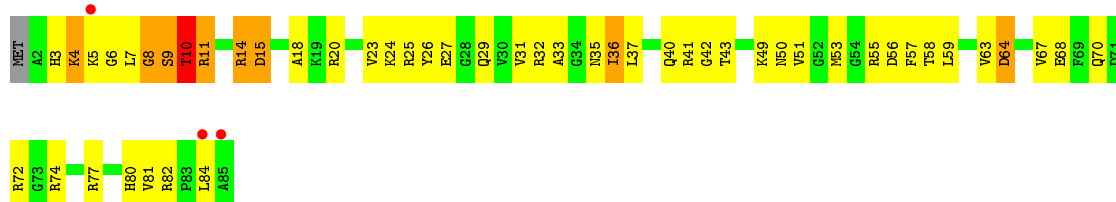


- Molecule 22: 50S RIBOSOMAL PROTEIN L27

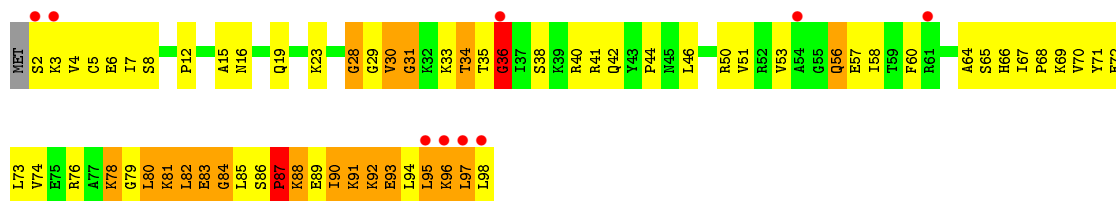




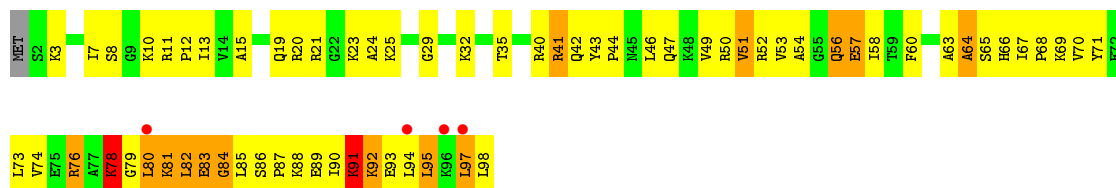
• Molecule 22: 50S RIBOSOMAL PROTEIN L27



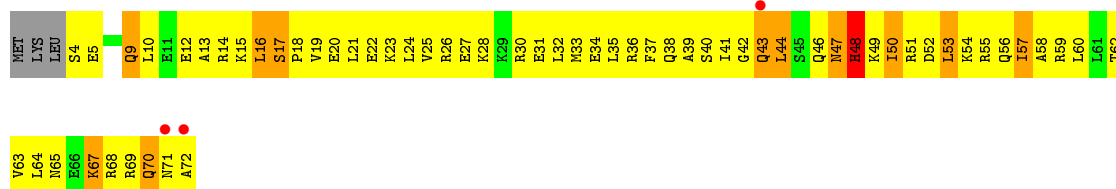
• Molecule 23: 50S RIBOSOMAL PROTEIN L28



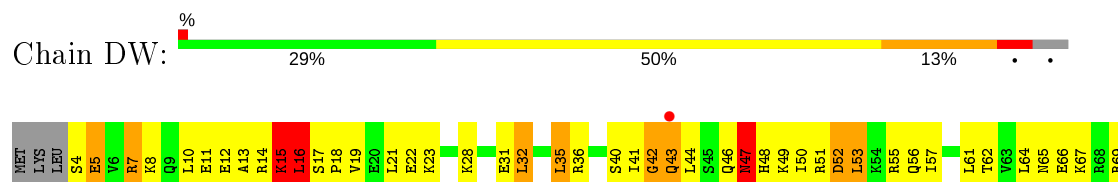
• Molecule 23: 50S RIBOSOMAL PROTEIN L28



• Molecule 24: 50S RIBOSOMAL PROTEIN L29

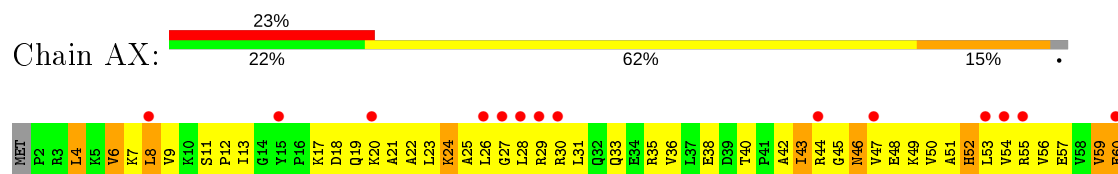


• Molecule 24: 50S RIBOSOMAL PROTEIN L29

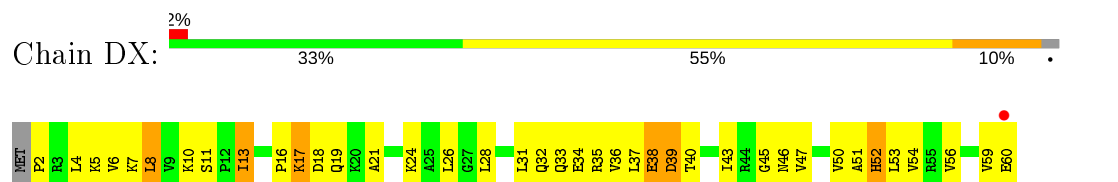


Q70
N71
A72

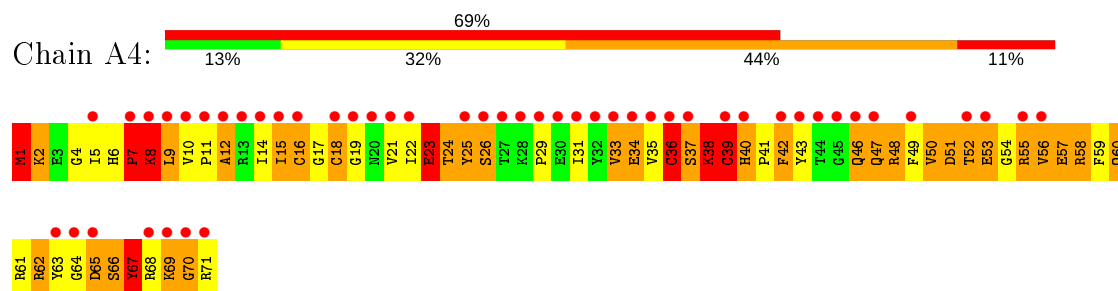
• Molecule 25: 50S RIBOSOMAL PROTEIN L30



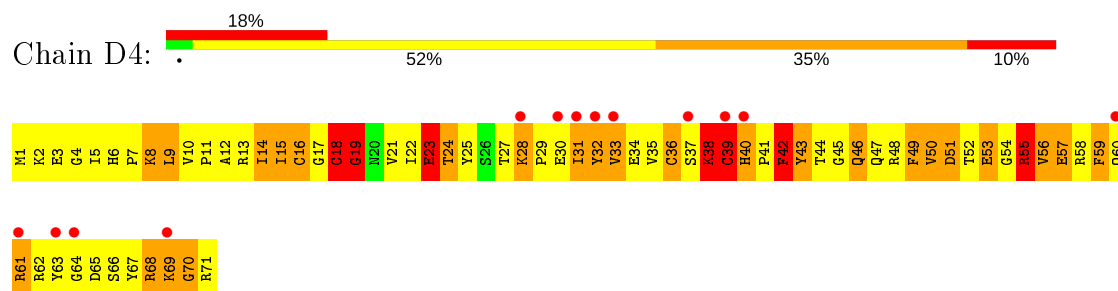
• Molecule 25: 50S RIBOSOMAL PROTEIN L30



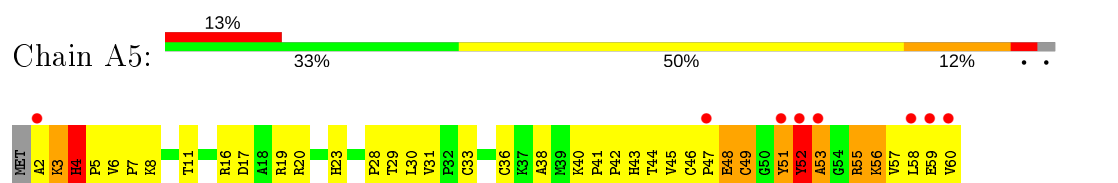
• Molecule 26: 50S RIBOSOMAL PROTEIN L31



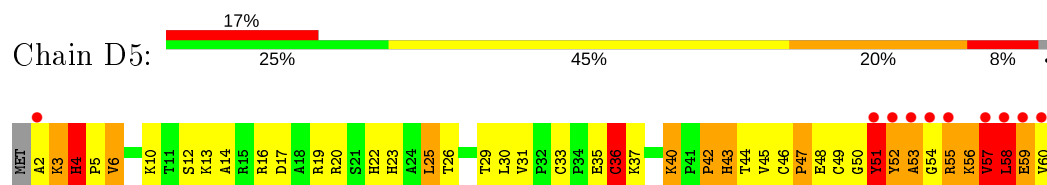
• Molecule 26: 50S RIBOSOMAL PROTEIN L31



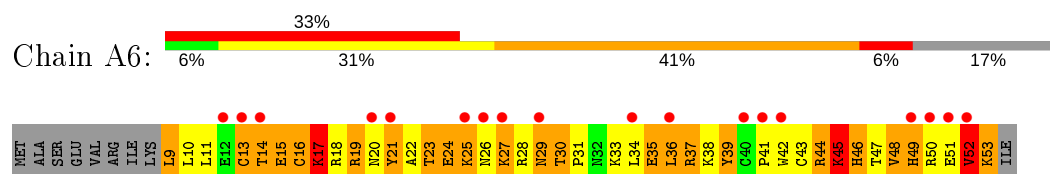
• Molecule 27: 50S RIBOSOMAL PROTEIN L32



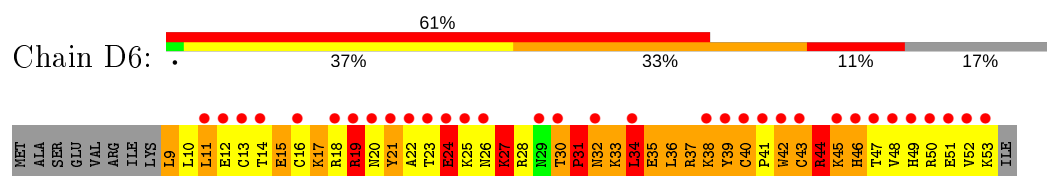
- Molecule 27: 50S RIBOSOMAL PROTEIN L32



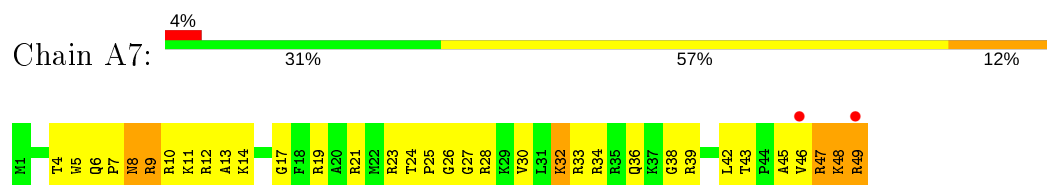
- Molecule 28: 50S RIBOSOMAL PROTEIN L33



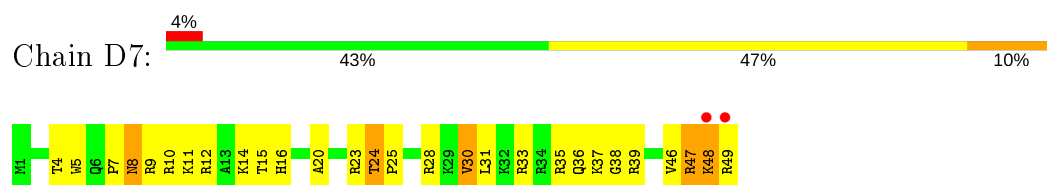
- Molecule 28: 50S RIBOSOMAL PROTEIN L33



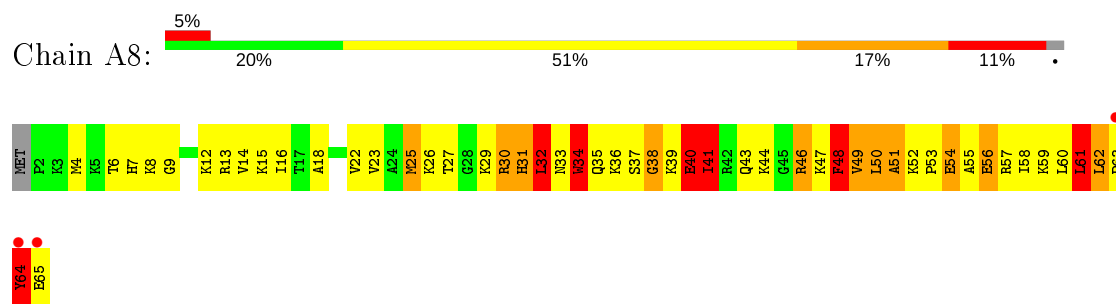
- Molecule 29: 50S RIBOSOMAL PROTEIN L34



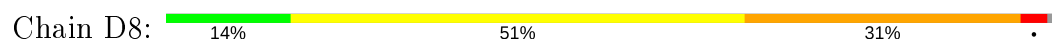
- Molecule 29: 50S RIBOSOMAL PROTEIN L34

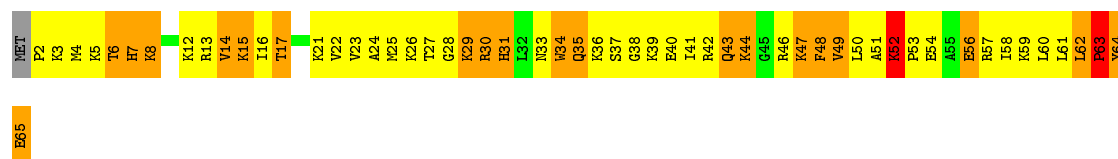


- Molecule 30: 50S RIBOSOMAL PROTEIN L35

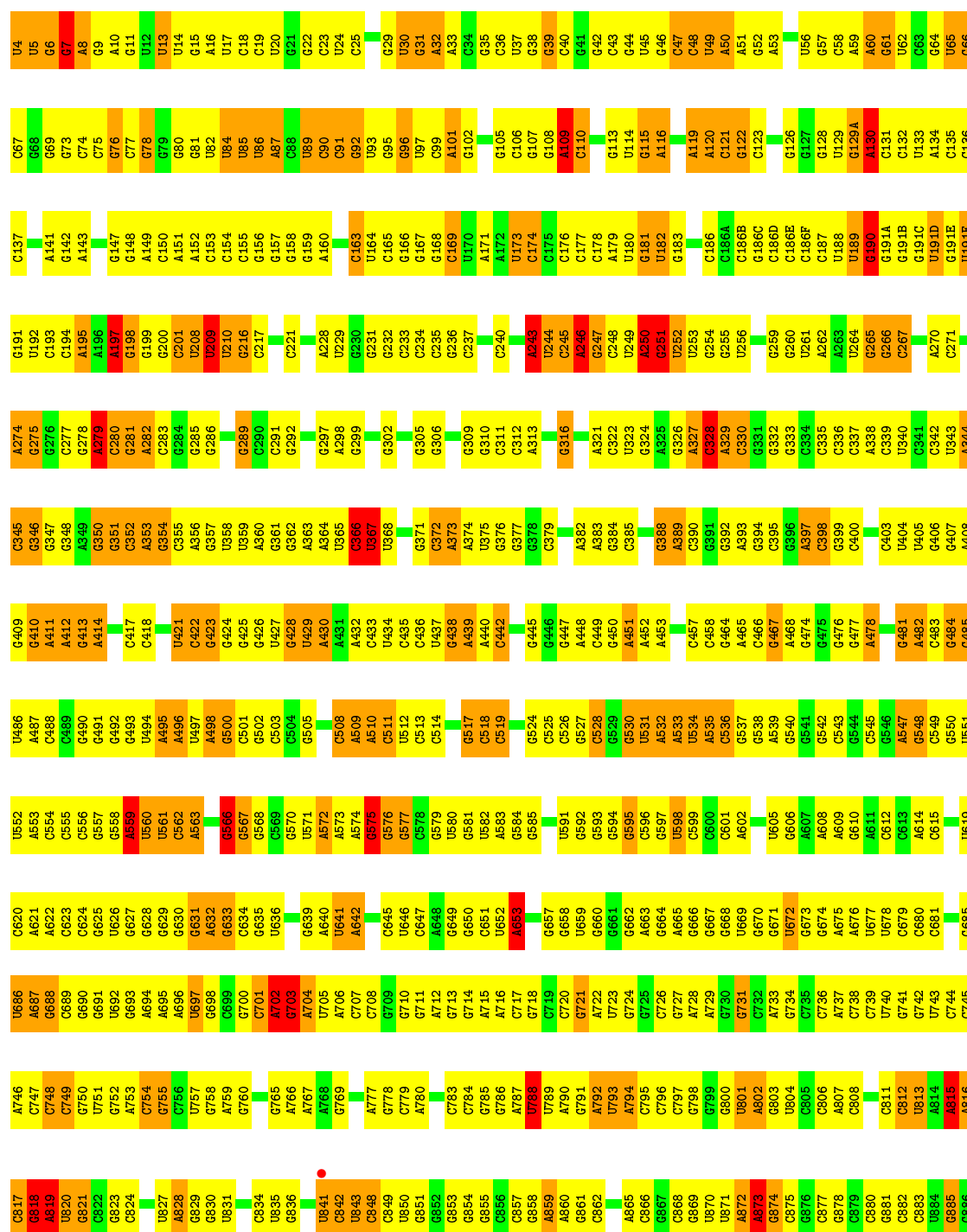
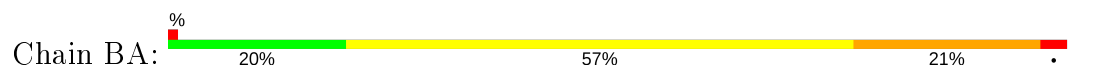


- Molecule 30: 50S RIBOSOMAL PROTEIN L35

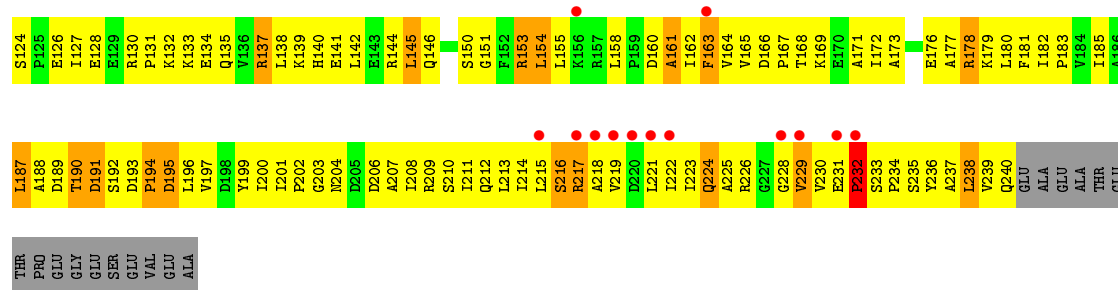




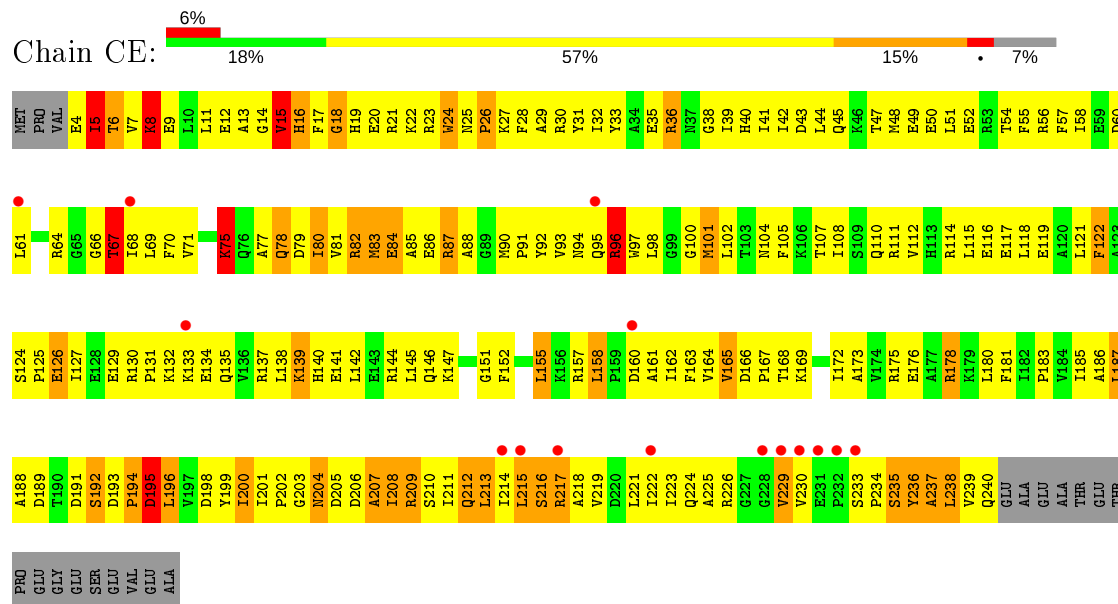
• Molecule 31: 16S ribosomal RNA



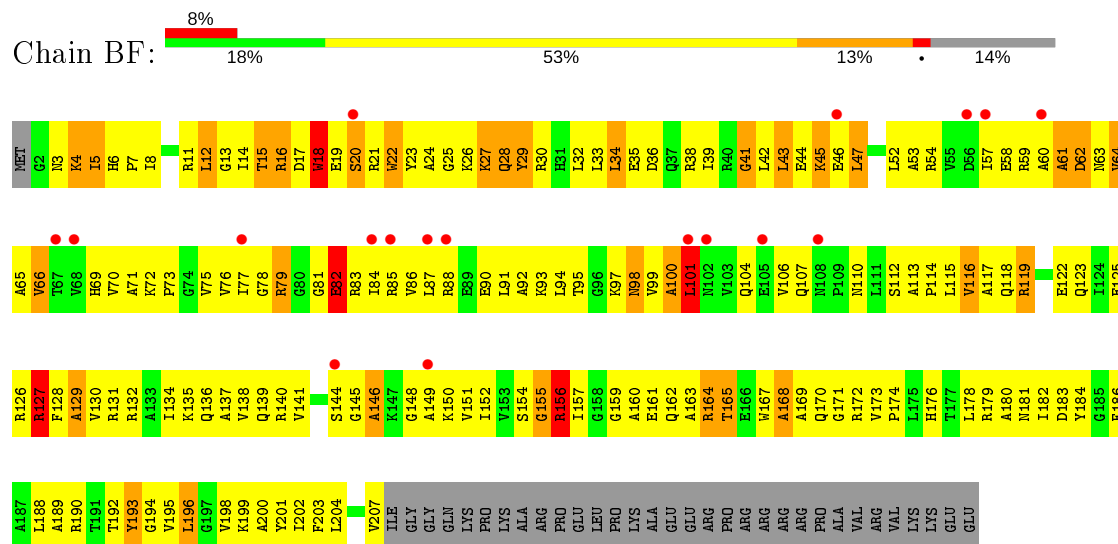




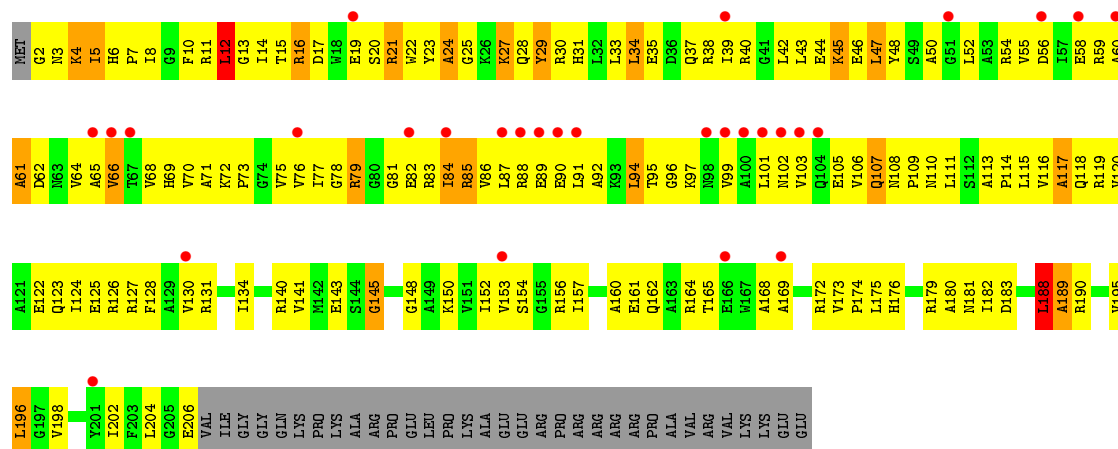
• Molecule 32: 30S ribosomal protein S2



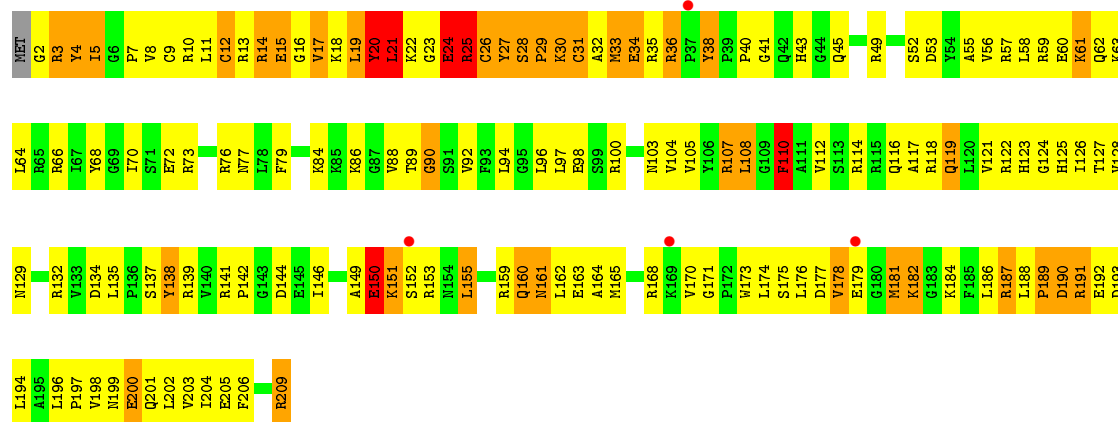
• Molecule 33: 30S ribosomal protein S3



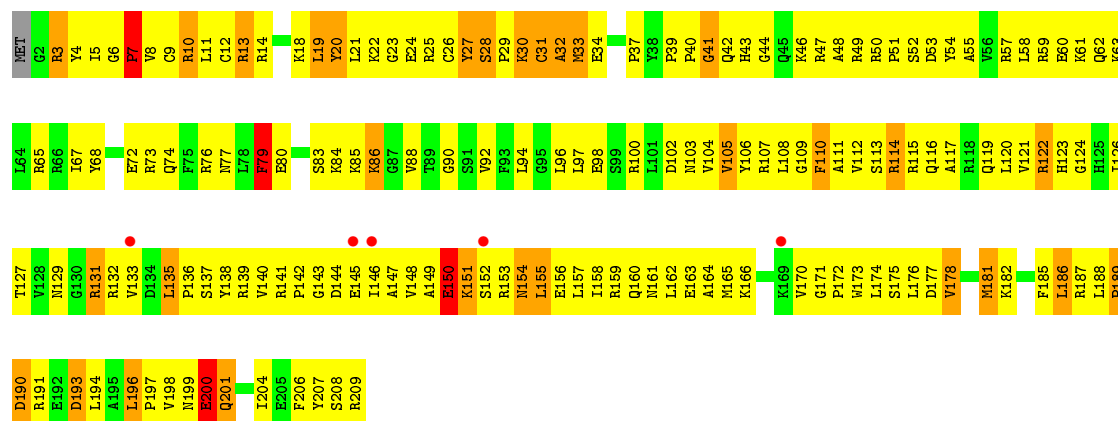
• Molecule 33: 30S ribosomal protein S3



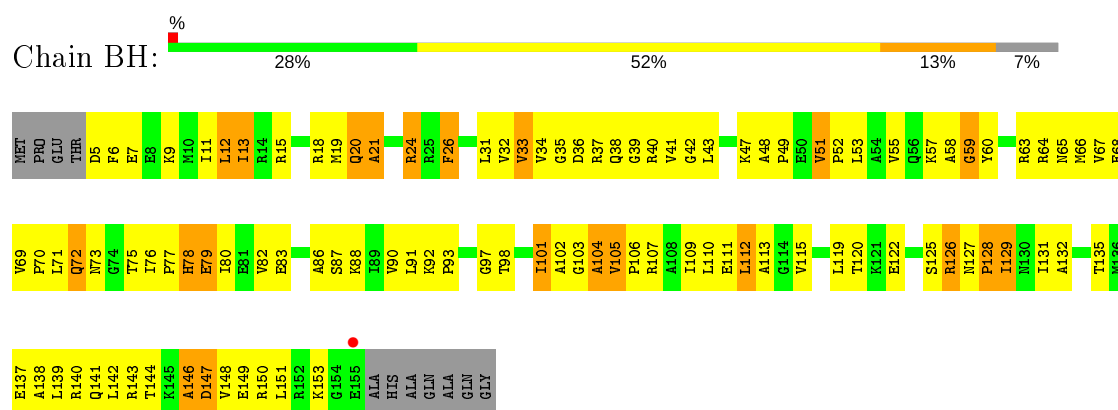
• Molecule 34: 30S ribosomal protein S4



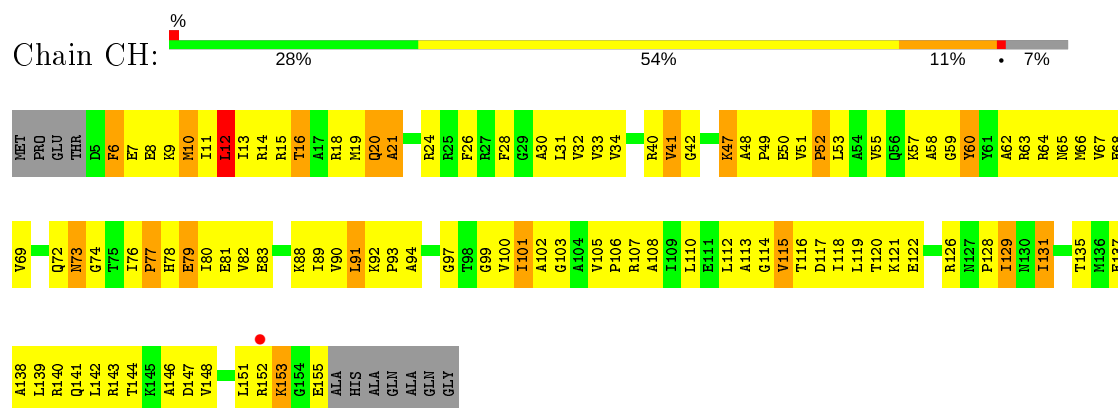
• Molecule 34: 30S ribosomal protein S4



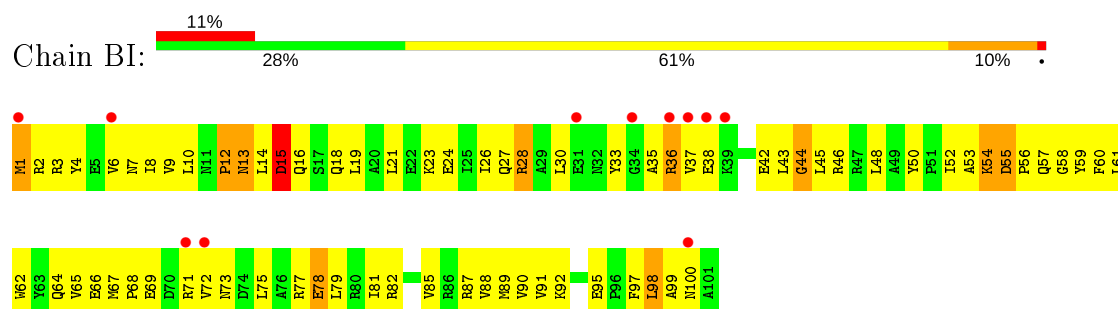
• Molecule 35: 30S ribosomal protein S5



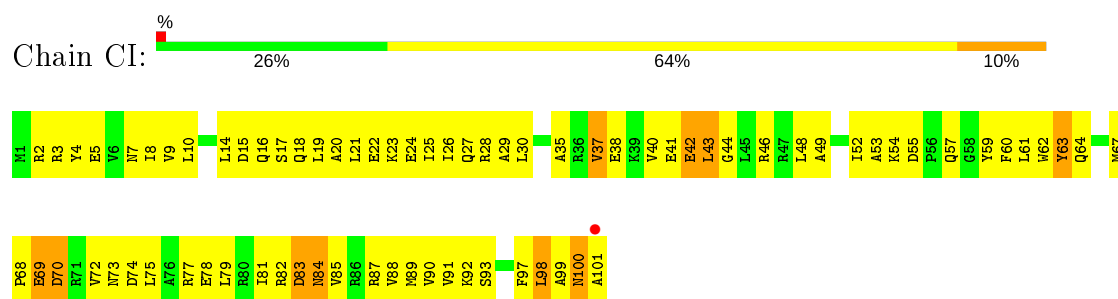
• Molecule 35: 30S ribosomal protein S5



• Molecule 36: 30S ribosomal protein S6

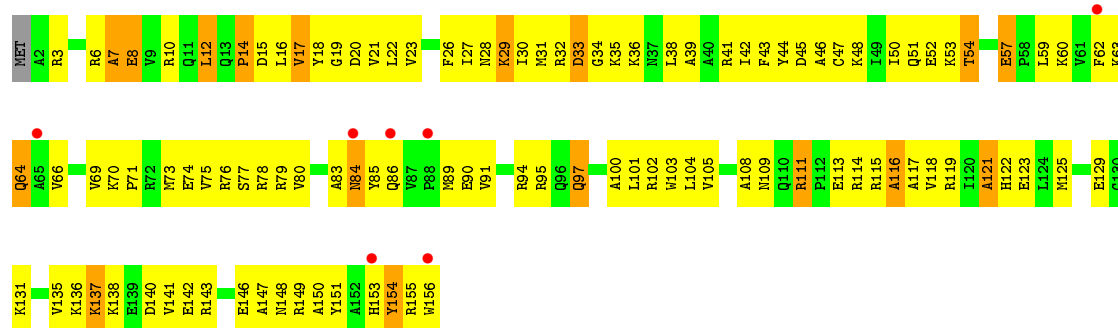


• Molecule 36: 30S ribosomal protein S6

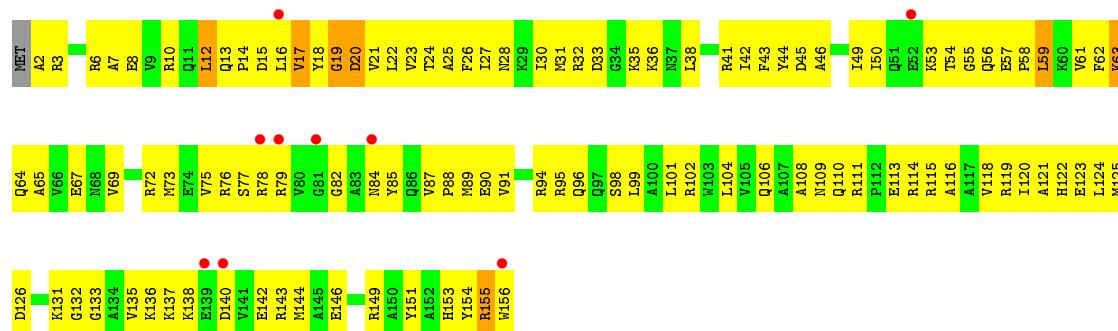


• Molecule 37: 30S ribosomal protein S7





• Molecule 37: 30S ribosomal protein S7



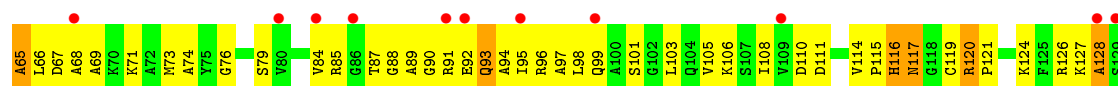
• Molecule 38: 30S ribosomal protein S8



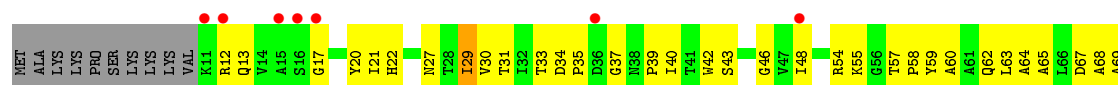
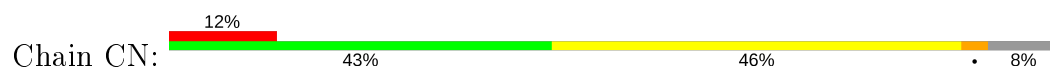
• Molecule 38: 30S ribosomal protein S8



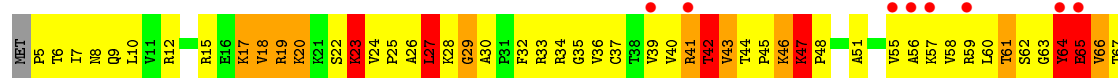
• Molecule 39: 30S ribosomal protein S9



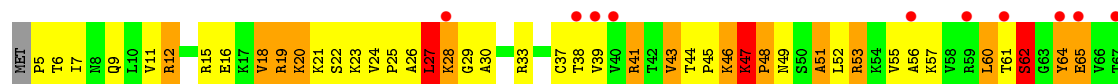
- Molecule 41: 30S ribosomal protein S11



- Molecule 42: 30S ribosomal protein S12



- Molecule 42: 30S ribosomal protein S12

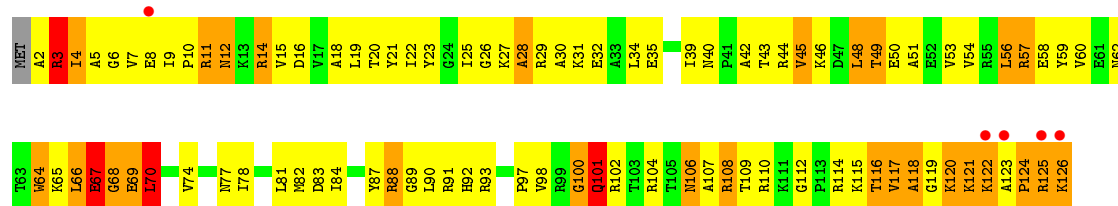


- Molecule 43: 30S ribosomal protein S13

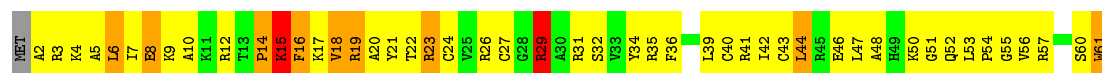


- Molecule 43: 30S ribosomal protein S13





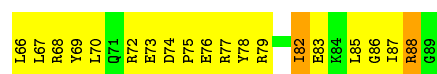
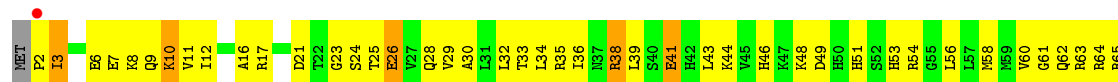
- Molecule 44: 30S ribosomal protein S14 type Z



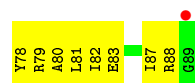
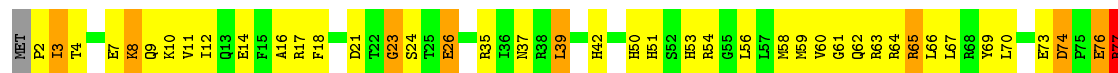
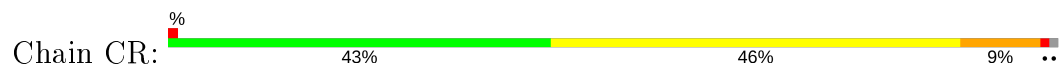
- Molecule 44: 30S ribosomal protein S14 type Z



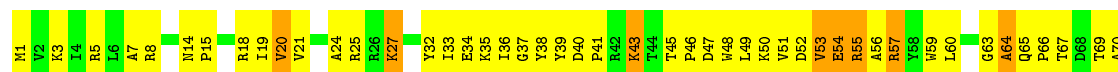
- Molecule 45: 30S ribosomal protein S15



- Molecule 45: 30S ribosomal protein S15

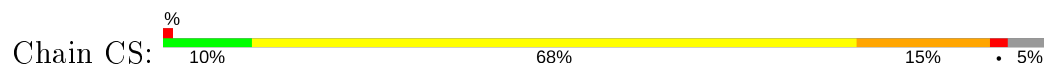


- Molecule 46: 30S ribosomal protein S16





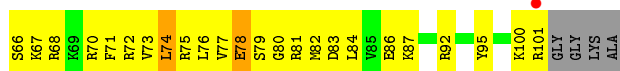
- Molecule 46: 30S ribosomal protein S16



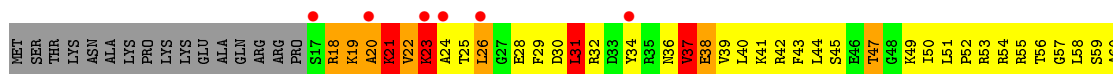
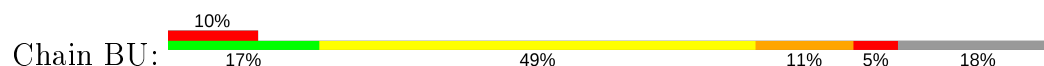
- Molecule 47: 30S ribosomal protein S17



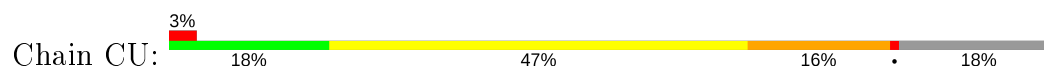
- Molecule 47: 30S ribosomal protein S17



- Molecule 48: 30S ribosomal protein S18

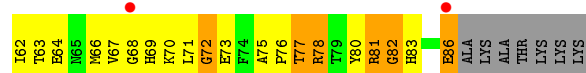
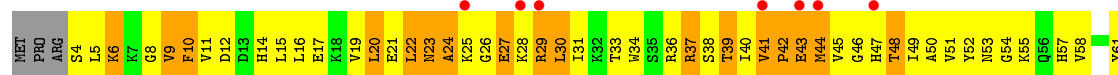
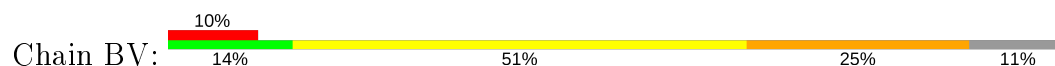


- Molecule 48: 30S ribosomal protein S18

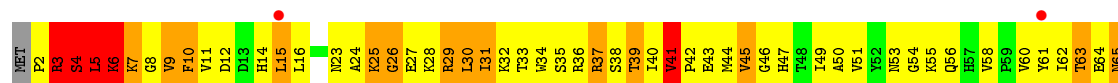
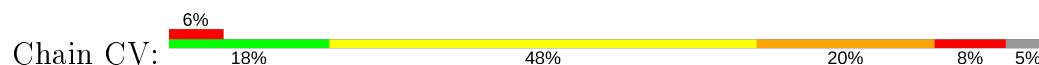




• Molecule 49: 30S ribosomal protein S19



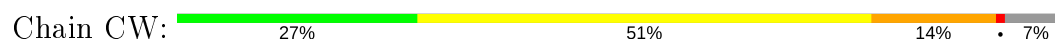
• Molecule 49: 30S ribosomal protein S19



• Molecule 50: 30S ribosomal protein S20

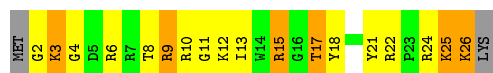


• Molecule 50: 30S ribosomal protein S20



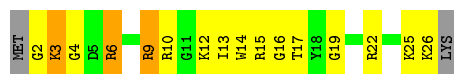
• Molecule 51: 30S ribosomal protein Thx

Chain BX: 



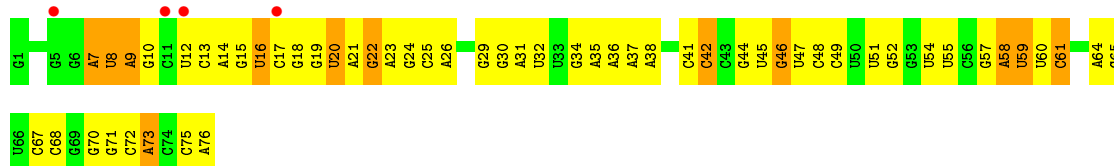
- Molecule 51: 30S ribosomal protein Thx

Chain CX: 



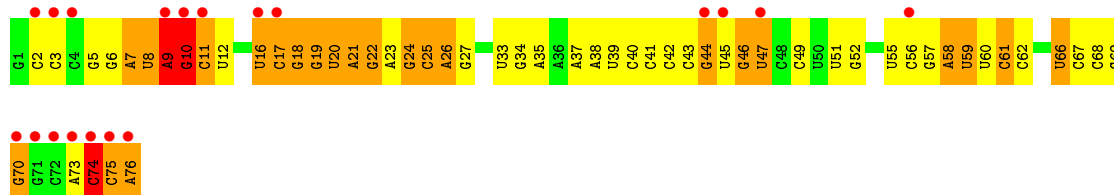
- Molecule 52: TRNA-PHE WITH UNMODIFIED NUCLEOTIDES EXCEPT FOR MIA37

Chain BD: 



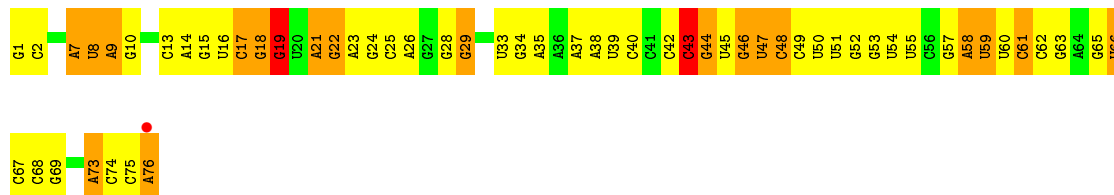
- Molecule 52: TRNA-PHE WITH UNMODIFIED NUCLEOTIDES EXCEPT FOR MIA37

Chain BB: 



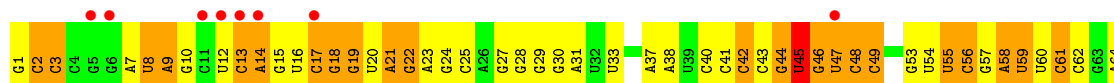
- Molecule 52: TRNA-PHE WITH UNMODIFIED NUCLEOTIDES EXCEPT FOR MIA37

Chain BC: 



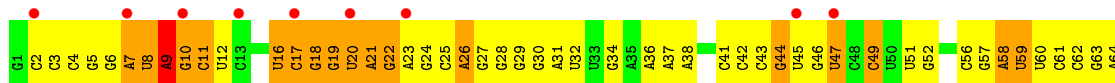
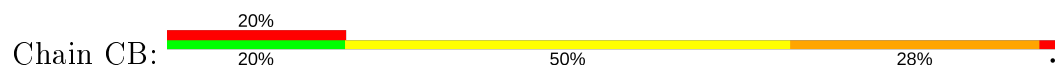
- Molecule 52: TRNA-PHE WITH UNMODIFIED NUCLEOTIDES EXCEPT FOR MIA37

Chain CD: 





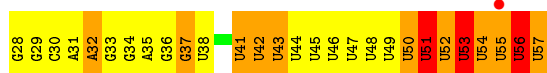
- Molecule 52: TRNA-PHE WITH UNMODIFIED NUCLEOTIDES EXCEPT FOR MIA37



- Molecule 52: TRNA-PHE WITH UNMODIFIED NUCLEOTIDES EXCEPT FOR MIA37



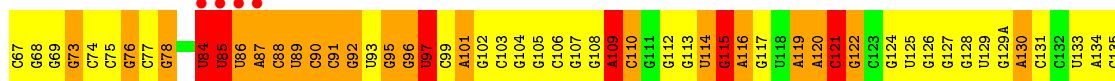
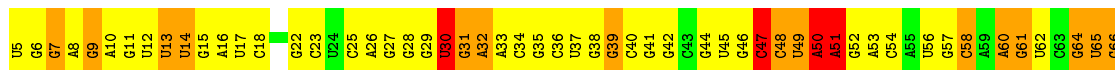
- Molecule 53: MRNA



- Molecule 53: MRNA



- Molecule 54: 16S ribosomal RNA

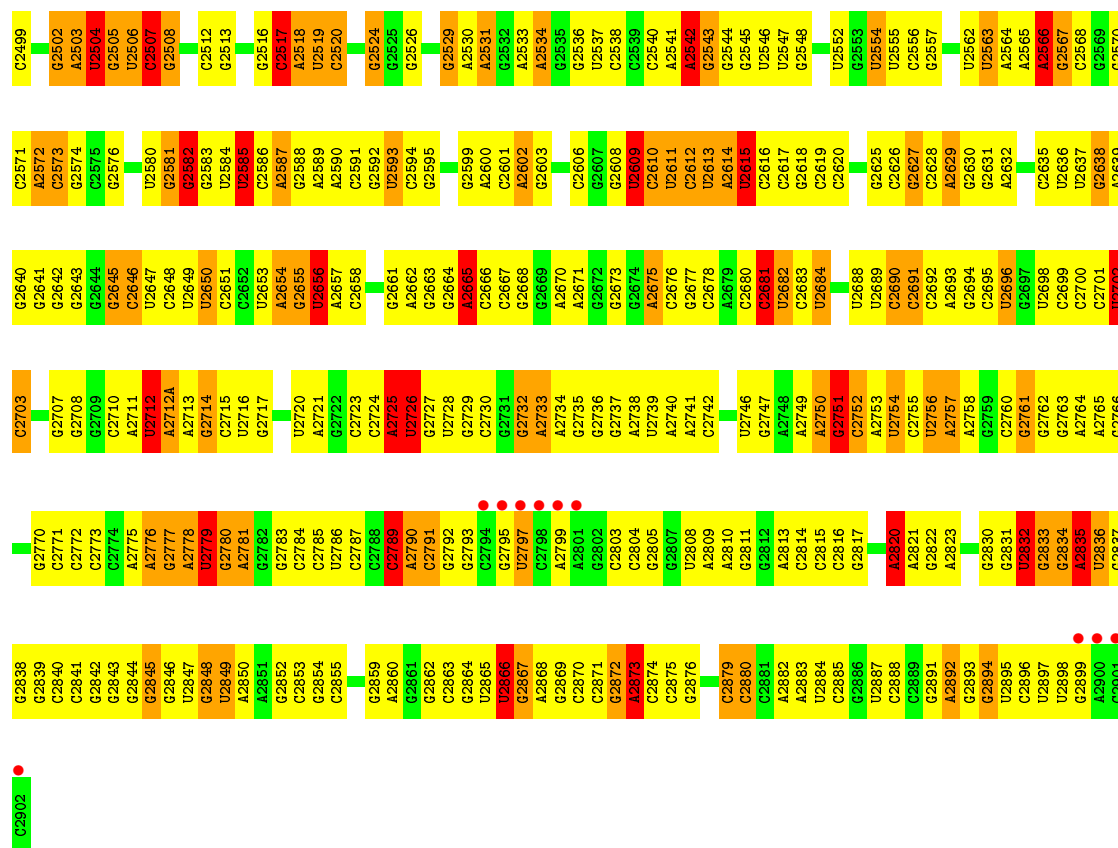


U1086	G1024	A959	A889	A815	G742	G680	G617	A547	G481	G407	C341	C268	G191E	C136
G1087	U1025	U950	G890	A816	U743	G681	G618	G548	A482	A408	C342	C269	U191F	C137
G1088	G1026	U961	G889	G817	G744	G682	U619	G549	A483	G409	U343	A270	G191	G138
U1089	C1027	A892	A892	G818	G745	G683	G620	G550	G484	G410	A344		U192	G139
U1090	G1028	G963	G893	A819	A746	A684	G621	U551	G485	A411	C345	A274	C193	A140
U1091	C1028A	A964	G894	U820	G747	G685	A622	U552	U486	A412	G346	G275	C194	A141
A1092	G1028B	A885	G895	G821	C748	U686	G623	A553	A487	G413	C347	G276	A195	G142
A1093	G1029	G966	G896		C749	A687	G624	G554	C488	A414	G348	C277	A196	A143
U1094	C1030	G967		G825	G750	G688	G625	G555	C489			G278	A197	G144
U1095	G1031	A968	A900	C826	U751	G689	G626	G556	G490	C418	G351	A279	G198	
U1096	A1032	A969	A901	U827	G752	G690	U627	G557		C419	C352	C280	G199	G147
C1097	G1032A	C970	G902	A828	A753	G691	G628		G493	U420	A353	G281	G200	G148
U1098	G1032B	G971	G903	G829	C764	U692	G629	A559	U494	U421	G354	A282	C201	A149
G1099	G1033	C972		G830	G755	G693	G630	U560	A495	C422		C283	U208	
C1100	G1034	G973	G906	U831		A694	G631	U561	A496	C423	G357		U209	A152
A1101	A974	A974		C832	G758	A695	A632	U562	U497	G424	U358	A288	U210	C153
A1102	G1036	A975	U911	U833	A759	A696	G633	A563	A498	C425	U359	G289	G216	C154
G1103	C1037	G976	C912	C834	G760	U697	G634	C564	G500	C426		C290	G220	
G1104	G1038	A977	A913	U835	A761	G698	G635	U565	C501	U427	A363	C291	G221	G157
A1105	G1039	A978	A914	G836	C764	C699	U636	G566	G502	C428	U364	G292	C221	G158
G1106	U1040	C979	A915	G837	G765	G700	G637	G567	C503	U429	U365		U222	G159
C1107		C980	A918	G838	A766	C701	G638	G568	C504	A430	C366	G297	U223	A160
G1108	A1046	U981	U918	U841	A767	A702	G639		C505	A431	U367	A298	C224	A161
	G1047	U982	A919	C842	A768	G703	A640	A572	G506	A432	U368	G299	C225	A162
C1112	G1048	A983	U920	U843		A704	U641	A573	C507	C433	U370	A300	G226	C163
	U1049		U921	C846	G773	U765	A642	A574	C508	U434	C371	G301		U164
C1116	G1050	A986	G922	C849		A706	G643	G575	A509	C435	G372	G302	U229	G165
G1117	C1051	G987	A923	U850	G776	G707	G650	G576	A510	C436	C373		G230	G166
C1118	U1052		C924	G851	A777	G708	G645	G577	C511	U437	A373	G305	G231	G167
G1119	G1053	C990	G925	G852	G778	G709	U646	C578	U512	C438	A374	G306	G232	G168
G1120	G1054	U991	G926	G853	G779	G710	C647	G579		A439	U375	G309	C233	C169
U1121	A1055	U992	G927	G854	C780	G711	A648	U580	G517	C440	G376		C234	U170
U1122	U1056	G993			A781	A712	G649		C518	C442	G377		C235	A171
G1123	G1057	A994	C930	C857	A782	G713	G650	G585	C519	G443			C241	U172
U1125	C1058		C931	G858	C783	G714	G651	C586		C444	C381	C312	C242	U173
U1126	C1059	G998	G932	A859	C784	A715	U652	A523	A523	G445	A382	A313	A243	C174
U1127	G1060	U999	G933	A860	G785	A716	G653	G524	G524	G446	A383	C314	C244	C175
C1128	U1062	A1000	A935	U863		G717	G654	G595			C384	A315	U244	C176
C1129	C1063	G1001		A864	U788	G718	A655	G596			C385	G316	C245	C177
A1130	G1064	G1002	G939	A865	A790	C720	G657	G597	C528	A451	U387	A321	A246	C178
G1131	U1065	G1003	C940	C866	G791	G721	G658	U598	G529	A452	G388	G322	G247	A179
C1132	A1066	A1004	G941	G869	A792	A722	U659	C599	G530	A453	C389	U323	A250	U180
G1133	C1067	A1005	G942	U870	U793	G723	G660	C600	U531	C454	C390	G324	G251	G181
U1135	C1068	C1006	U943	G871	C795	G724	G661	C601	A532		G391	A325	U252	U182
C1136	U1070	C1008	G945	A872		G725	A662	A602	U533	G456	A393	G326	U253	G183
C1137	C1071	G1009	A946	A873	G800	G726	A663	U603	U534		A394	A327	G254	A185
G1138	G1072	G947	G947	G874	U801	A728	G664	G604	A535	C458	C395	C328	G255	C186
G1139	U1073	C948		G875	A802	A729	A665	G606	C536	A465	G396	A329	U256	C186A
C1140	C1074	A949		C877	G803	G730	G666	A607	G537	C466	A397	C330	G257	C186B
C1141	A1015	U950		G878	U804	G731	G671	A608	G538	G467	C398	G332	G258	G186C
G1142	A1016	G879		C878	C805	C732	U672	A609	A539	A468	G399	G333	G260	C186D
G1143	G1017	U952		C880	C806	G736	G673	G610	G540	C474	C400	C334	U261	U188
G1144	C1018	G953		G881	A807	G737	G674	A611	G541	G475	C401	C335	A262	U189
C1145	U1019	G954		C882	C808	G738	A675	G612	G542	G476	G402	C336	A263	G190
A1146	G1081	U1020		C883		G739	A676	G613	C543	G477	C403	C337	U264	G191A
C1147	G1082	G1021		U812	G812	U740	G677	A614	G544	A478	U404	A338	G265	G191B
G1148		U957		G885	U813		U678	C615	C545	C479	U405	C339	G266	G191C
C1149	U1085	A958		A814	G741		G679	G616	G546	U480	G406	U340	C267	U191D

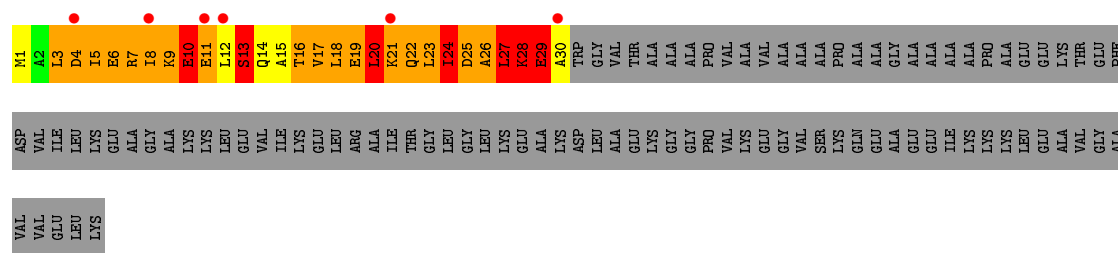




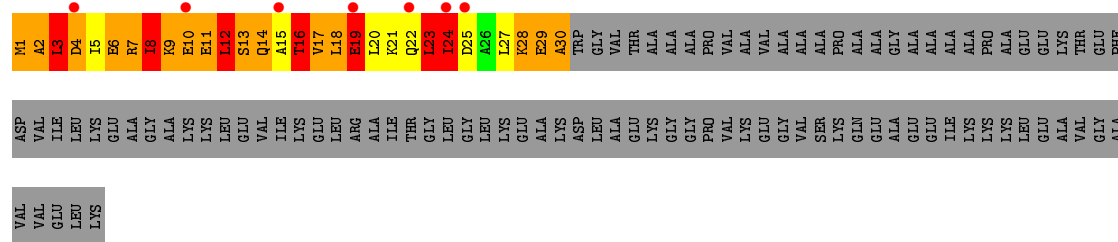




- Molecule 56: 50S ribosomal protein L7/L12

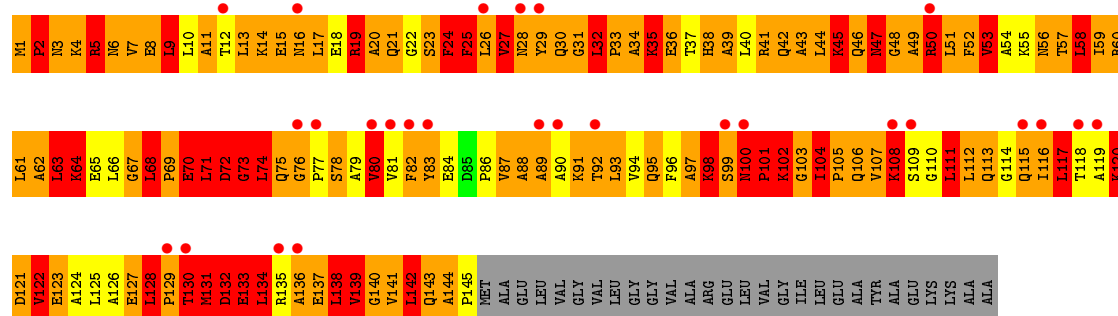


- Molecule 56: 50S ribosomal protein L7/L12



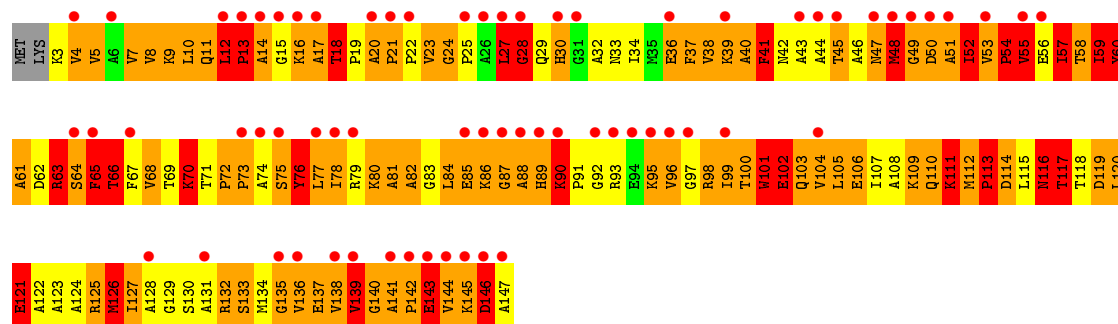
- Molecule 57: 50S ribosomal protein L10

Chain DY: 16% 16% 43% 24% 16%



• Molecule 58: 50S ribosomal protein L11

Chain DL: 45% 26% 49% 20%



4 Data and refinement statistics

Property	Value	Source
Space group	P 21 21 21	Depositor
Cell constants a, b, c, α , β , γ	209.46 Å 446.20 Å 623.05 Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	300.00 – 3.10 223.10 – 3.00	Depositor EDS
% Data completeness (in resolution range)	96.5 (300.00-3.10) 99.8 (223.10-3.00)	Depositor EDS
R_{merge}	0.19	Depositor
R_{sym}	0.27	Depositor
$\langle I/\sigma(I) \rangle$ ¹	1.72 (at 3.01 Å)	Xtriage
Refinement program	CNS	Depositor
R, R_{free}	0.215 , 0.255 0.222 , 0.260	Depositor DCC
R_{free} test set	34306 reflections (2.99%)	wwPDB-VP
Wilson B-factor (Å ²)	61.5	Xtriage
Anisotropy	0.212	Xtriage
Bulk solvent k_{sol} (e/Å ³), B_{sol} (Å ²)	0.27 , 75.2	EDS
L-test for twinning ²	$\langle L \rangle = 0.43$, $\langle L^2 \rangle = 0.25$	Xtriage
Estimated twinning fraction	No twinning to report.	Xtriage
F_o, F_c correlation	0.93	EDS
Total number of atoms	307345	wwPDB-VP
Average B, all atoms (Å ²)	77.0	wwPDB-VP

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 1.45% of the height of the origin peak. No significant pseudotranslation is detected.*

¹Intensities estimated from amplitudes.

²Theoretical values of $\langle |L| \rangle$, $\langle L^2 \rangle$ for acentric reflections are 0.5, 0.333 respectively for untwinned datasets, and 0.375, 0.2 for perfectly twinned datasets.

5 Model quality ⓘ

5.1 Standard geometry ⓘ

Bond lengths and bond angles in the following residue types are not validated in this section: MIA, ZN, MG

The Z score for a bond length (or angle) is the number of standard deviations the observed value is removed from the expected value. A bond length (or angle) with $|Z| > 5$ is considered an outlier worth inspection. RMSZ is the root-mean-square of all Z scores of the bond lengths (or angles).

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
1	AA	0.55	6/70167 (0.0%)	0.87	170/109541 (0.2%)
2	AB	0.42	1/2928 (0.0%)	0.80	0/4568
2	DB	0.58	3/2928 (0.1%)	0.85	5/4568 (0.1%)
3	AD	0.45	0/2165	0.80	1/2919 (0.0%)
3	DD	0.54	0/2165	0.87	1/2919 (0.0%)
4	AE	0.39	0/1601	0.77	1/2160 (0.0%)
4	DE	0.50	0/1601	0.89	3/2160 (0.1%)
5	AF	0.39	0/1662	0.74	1/2249 (0.0%)
5	DF	0.50	0/1620	0.76	0/2194
6	AG	0.31	0/1499	0.58	0/2016
6	DG	0.38	0/1499	0.66	0/2016
7	AH	0.28	0/1332	0.62	0/1802
7	DH	0.41	0/1332	0.89	2/1802 (0.1%)
8	AK	0.33	0/1151	0.74	0/1558
8	DK	0.35	0/1151	0.74	1/1558 (0.1%)
9	AM	0.34	0/1131	0.66	0/1525
9	DM	0.46	0/1131	0.81	1/1525 (0.1%)
10	AN	0.40	0/943	0.67	0/1269
10	DN	0.49	0/943	0.75	0/1269
11	AO	0.39	0/1162	0.80	2/1544 (0.1%)
11	DO	0.51	0/1162	0.95	3/1544 (0.2%)
12	AP	0.39	0/1143	0.72	0/1527
12	DP	0.53	0/1143	0.80	1/1527 (0.1%)
13	A0	0.38	0/974	0.67	0/1302
13	D0	0.45	0/982	0.79	1/1312 (0.1%)
14	AQ	0.34	0/892	0.70	0/1187
14	DQ	0.41	0/892	0.84	1/1187 (0.1%)
15	AR	0.38	0/1155	0.68	0/1542
15	DR	0.44	0/1155	0.75	1/1542 (0.1%)
16	A1	0.38	0/982	0.67	0/1306
16	D1	0.49	0/982	0.75	0/1306
17	A2	0.38	0/790	0.75	0/1057

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
17	D2	0.45	0/790	0.80	0/1057
18	AS	0.42	0/911	0.70	0/1220
18	DS	0.45	0/911	0.73	0/1220
19	AT	0.49	0/739	0.73	0/993
19	DT	0.52	0/739	0.79	0/993
20	AU	0.44	0/798	0.78	1/1064 (0.1%)
20	DU	0.45	0/798	0.77	0/1064
21	AV	0.35	0/1521	0.74	3/2064 (0.1%)
21	DV	0.59	4/1615 (0.2%)	0.93	7/2191 (0.3%)
22	A3	0.41	0/671	0.74	0/892
22	D3	0.46	0/671	0.76	0/892
23	AZ	0.40	0/770	0.79	1/1022 (0.1%)
23	DZ	0.45	0/770	0.76	0/1022
24	AW	0.43	0/583	0.73	0/771
24	DW	0.49	0/583	0.78	0/771
25	AX	0.31	0/474	0.67	0/635
25	DX	0.39	0/474	0.71	0/635
26	A4	0.43	1/594 (0.2%)	0.82	1/795 (0.1%)
26	D4	0.43	0/594	1.03	6/795 (0.8%)
27	A5	0.38	0/473	0.68	0/639
27	D5	0.48	0/473	0.75	0/639
28	A6	0.37	0/396	0.87	0/529
28	D6	0.37	0/396	0.88	0/529
29	A7	0.45	0/438	0.71	0/575
29	D7	0.52	0/438	0.79	0/575
30	A8	0.51	0/525	0.97	1/691 (0.1%)
30	D8	0.61	0/525	0.95	0/691
31	BA	0.45	2/36457 (0.0%)	0.80	41/56899 (0.1%)
32	BE	0.31	0/1959	0.59	0/2642
32	CE	0.32	0/1959	0.61	0/2642
33	BF	0.31	0/1636	0.57	0/2205
33	CF	0.35	0/1629	0.59	0/2195
34	BG	0.39	0/1733	0.74	5/2318 (0.2%)
34	CG	0.40	0/1733	0.69	3/2318 (0.1%)
35	BH	0.35	0/1171	0.66	0/1576
35	CH	0.39	0/1171	0.67	0/1576
36	BI	0.39	0/856	0.65	0/1154
36	CI	0.39	0/856	0.66	0/1154
37	BJ	0.32	0/1276	0.57	0/1709
37	CJ	0.34	0/1276	0.57	0/1709
38	BK	0.31	0/1136	0.61	0/1527
38	CK	0.36	0/1136	0.65	0/1527
39	BL	0.32	0/1029	0.59	0/1379

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
39	CL	0.33	0/1029	0.63	0/1379
40	BM	0.31	0/814	0.61	0/1095
40	CM	0.34	0/814	0.61	0/1095
41	BN	0.35	0/900	0.61	0/1213
41	CN	0.38	0/900	0.63	0/1213
42	BO	0.38	0/991	0.67	0/1327
42	CO	0.44	0/991	0.84	1/1327 (0.1%)
43	BP	0.32	0/974	0.66	1/1303 (0.1%)
43	CP	0.40	0/1008	0.74	0/1347
44	BQ	0.34	0/501	0.59	0/664
44	CQ	0.40	0/501	0.69	1/664 (0.2%)
45	BR	0.35	0/745	0.56	0/992
45	CR	0.39	0/745	0.63	0/992
46	BS	0.37	0/721	0.61	0/970
46	CS	0.35	0/721	0.66	0/970
47	BT	0.35	0/847	0.62	0/1131
47	CT	0.36	0/847	0.60	0/1131
48	BU	0.42	0/596	0.75	1/790 (0.1%)
48	CU	0.38	0/596	0.69	0/790
49	BV	0.35	0/679	0.67	0/913
49	CV	0.50	0/717	0.97	3/963 (0.3%)
50	BW	0.32	0/765	0.62	0/1007
50	CW	0.31	0/765	0.64	0/1007
51	BX	0.33	0/221	0.58	0/288
51	CX	0.38	0/221	0.51	0/288
52	BB	0.35	0/1783	0.77	4/2776 (0.1%)
52	BC	0.43	0/1783	0.78	1/2776 (0.0%)
52	BD	0.31	0/1783	0.74	0/2776
52	CB	0.41	0/1783	1.00	6/2776 (0.2%)
52	CC	0.58	0/1783	0.93	3/2776 (0.1%)
52	CD	0.38	0/1783	0.83	3/2776 (0.1%)
53	B1	0.47	0/689	0.94	3/1069 (0.3%)
53	C1	0.48	0/689	0.91	2/1069 (0.2%)
54	CA	0.50	1/36435 (0.0%)	0.82	58/56865 (0.1%)
55	DA	0.70	12/70233 (0.0%)	0.96	262/109643 (0.2%)
56	DI	1.11	2/236 (0.8%)	1.41	4/315 (1.3%)
56	DJ	1.82	5/236 (2.1%)	1.78	9/315 (2.9%)
57	DY	0.98	5/1123 (0.4%)	1.55	24/1520 (1.6%)
58	DL	0.72	1/1091 (0.1%)	1.34	16/1479 (1.1%)
All	All	0.53	43/328085 (0.0%)	0.85	666/490785 (0.1%)

Chiral center outliers are detected by calculating the chiral volume of a chiral center and verifying if the center is modelled as a planar moiety or with the opposite hand. A planarity outlier is detected

by checking planarity of atoms in a peptide group, atoms in a mainchain group or atoms of a sidechain that are expected to be planar.

Mol	Chain	#Chirality outliers	#Planarity outliers
1	AA	0	109
2	AB	0	4
2	DB	0	4
3	AD	0	2
31	BA	0	34
52	BB	0	1
52	BC	0	3
52	BD	0	1
52	CB	0	1
52	CD	0	2
53	B1	0	4
53	C1	0	3
54	CA	0	57
55	DA	0	170
All	All	0	395

All (43) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
31	BA	889	A	O3'-P	-25.82	1.30	1.61
56	DJ	17	VAL	CB-CG2	-16.90	1.17	1.52
57	DY	80	VAL	CB-CG1	-13.58	1.24	1.52
56	DJ	17	VAL	CA-CB	-9.21	1.35	1.54
57	DY	139	VAL	CB-CG2	-8.82	1.34	1.52
56	DI	24	ILE	CB-CG2	-8.49	1.26	1.52
56	DJ	17	VAL	CB-CG1	-7.93	1.36	1.52
2	DB	81	G	C6-N1	-7.89	1.34	1.39
55	DA	2665	A	C6-N6	-7.64	1.27	1.33
1	AA	1342	A	O3'-P	-7.64	1.51	1.61
1	AA	1359	A	C5-C6	-7.33	1.34	1.41
2	DB	95	U	C2-O2	7.04	1.28	1.22
55	DA	621	A	C6-N6	-7.02	1.28	1.33
58	DL	52	ILE	CB-CG1	-7.01	1.34	1.54
55	DA	945	A	N9-C4	7.00	1.42	1.37
57	DY	111	LEU	C-O	-6.85	1.10	1.23
54	CA	788	U	N3-C4	6.75	1.44	1.38
31	BA	788	U	N3-C4	6.72	1.44	1.38
56	DI	26	ALA	CA-CB	-6.68	1.38	1.52
55	DA	897	C	C4-C5	-6.59	1.37	1.43
55	DA	383	U	N1-C2	6.30	1.44	1.38
55	DA	74	A	N9-C4	-6.05	1.34	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
57	DY	51	LEU	CG-CD1	-5.94	1.29	1.51
56	DJ	2	ALA	CA-CB	-5.93	1.40	1.52
55	DA	2665	A	C5-C6	-5.93	1.35	1.41
21	DV	200	GLY	CA-C	5.86	1.61	1.51
21	DV	61	LEU	CA-CB	5.86	1.67	1.53
55	DA	383	U	C2-O2	5.78	1.27	1.22
1	AA	2665	A	C6-N6	-5.66	1.29	1.33
56	DJ	17	VAL	CA-C	-5.57	1.38	1.52
1	AA	654(M)	C	N1-C2	5.56	1.45	1.40
2	DB	95	U	N1-C2	5.55	1.43	1.38
55	DA	1612	C	N1-C2	-5.45	1.34	1.40
55	DA	654(H)	G	C5-C6	-5.44	1.36	1.42
55	DA	654(M)	C	N1-C2	5.43	1.45	1.40
1	AA	621	A	C6-N6	-5.26	1.29	1.33
1	AA	654(H)	G	C5-C6	-5.24	1.37	1.42
21	DV	196	VAL	CB-CG1	-5.23	1.41	1.52
26	A4	1	MET	SD-CE	-5.19	1.48	1.77
55	DA	896	A	C2-N3	-5.17	1.28	1.33
21	DV	196	VAL	CA-CB	-5.15	1.44	1.54
57	DY	73	GLY	C-O	-5.11	1.15	1.23
2	AB	81	G	C6-N1	-5.05	1.36	1.39

All (666) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
31	BA	889	A	P-O3'-C3'	-30.02	83.67	119.70
52	CB	74	C	C1'-O4'-C4'	-20.78	93.28	109.90
31	BA	889	A	O3'-P-O5'	-20.67	64.72	104.00
1	AA	4	C	OP1-P-O3'	-19.04	63.31	105.20
52	CB	74	C	O4'-C1'-N1	18.78	123.23	108.20
55	DA	4	C	OP1-P-O3'	-18.47	64.56	105.20
55	DA	4	C	OP2-P-O3'	-18.18	65.21	105.20
1	AA	4	C	OP2-P-O3'	-18.01	65.57	105.20
1	AA	1342	A	P-O3'-C3'	17.11	140.24	119.70
55	DA	2286	A	C1'-O4'-C4'	-16.67	96.56	109.90
52	CC	20	U	C1'-O4'-C4'	-16.02	97.09	109.90
55	DA	945	A	C1'-O4'-C4'	-15.95	97.14	109.90
55	DA	2468	G	C1'-O4'-C4'	-15.21	97.73	109.90
7	DH	125	VAL	C-N-CD	-15.16	87.25	120.60
57	DY	51	LEU	CB-CG-CD2	-14.72	85.97	111.00
31	BA	889	A	OP1-P-O3'	14.69	137.53	105.20
1	AA	1379	A	C1'-O4'-C4'	-14.66	98.17	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	AA	2286	A	C1'-O4'-C4'	-13.73	98.92	109.90
1	AA	1762	A	C1'-O4'-C4'	-13.70	98.94	109.90
55	DA	945	A	N9-C4-C5	-13.70	100.32	105.80
55	DA	1379	A	C1'-O4'-C4'	-13.43	99.16	109.90
1	AA	2311	A	C1'-O4'-C4'	-13.25	99.30	109.90
56	DJ	3	LEU	CB-CG-CD2	-12.88	89.11	111.00
55	DA	1372	U	C5-C6-N1	12.46	128.93	122.70
1	AA	1342	A	OP1-P-O3'	12.41	132.50	105.20
55	DA	1372	U	C4-C5-C6	-12.38	112.27	119.70
57	DY	76	GLY	C-N-CD	-12.28	93.59	120.60
55	DA	1544	C	N1-C1'-C2'	12.07	129.69	114.00
55	DA	607	U	N3-C4-O4	-11.94	111.04	119.40
55	DA	1925	C	N1-C1'-C2'	-11.79	98.68	114.00
52	CC	20	U	N1-C1'-C2'	11.77	129.30	114.00
55	DA	945	A	N9-C1'-C2'	11.60	129.07	114.00
1	AA	322	A	C1'-O4'-C4'	-11.53	100.68	109.90
1	AA	5	A	O5'-P-OP1	-11.27	95.56	105.70
52	CB	74	C	C2-N1-C1'	10.88	130.77	118.80
52	CB	74	C	C6-N1-C1'	-10.68	107.98	120.80
55	DA	607	U	C5-C4-O4	-10.59	119.55	125.90
1	AA	1359	A	C3'-C2'-C1'	-10.54	93.07	101.50
52	CC	20	U	O4'-C1'-N1	10.46	116.57	108.20
56	DJ	17	VAL	CA-CB-CG2	-10.16	95.66	110.90
52	CD	45	U	N1-C1'-C2'	10.11	127.14	114.00
57	DY	73	GLY	N-CA-C	-9.87	88.44	113.10
55	DA	945	A	C6-C5-N7	-9.63	125.56	132.30
55	DA	2447	G	N9-C1'-C2'	9.59	126.46	114.00
57	DY	19	ARG	NE-CZ-NH1	-9.51	115.55	120.30
52	CD	45	U	O4'-C1'-N1	9.46	115.77	108.20
55	DA	2311	A	C1'-O4'-C4'	-9.33	102.44	109.90
58	DL	24	GLY	N-CA-C	-9.23	90.02	113.10
55	DA	1616	A	N9-C1'-C2'	9.20	125.96	114.00
1	AA	2789	C	O4'-C1'-N1	9.17	115.53	108.20
55	DA	1828	G	N9-C1'-C2'	9.16	125.90	114.00
55	DA	2656	U	N3-C4-O4	-9.12	113.01	119.40
56	DI	24	ILE	CG1-CB-CG2	-9.08	91.43	111.40
21	DV	193	GLU	C-N-CD	-9.08	100.63	120.60
55	DA	1340	U	N1-C1'-C2'	9.04	125.75	114.00
1	AA	322	A	N9-C1'-C2'	9.01	125.71	114.00
55	DA	508	G	N9-C1'-C2'	9.00	125.70	114.00
1	AA	607	U	C5-C4-O4	-8.95	120.53	125.90
1	AA	2490	G	C1'-O4'-C4'	-8.90	102.78	109.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
55	DA	1372	U	N1-C2-N3	-8.81	109.61	114.90
55	DA	2307	G	O4'-C1'-N9	8.73	115.19	108.20
55	DA	1698	A	N9-C1'-C2'	8.72	125.34	114.00
55	DA	2789	C	O4'-C1'-N1	8.71	115.17	108.20
55	DA	945	A	C8-N9-C1'	-8.67	112.10	127.70
55	DA	1929	G	N9-C1'-C2'	8.62	125.20	114.00
21	DV	61	LEU	CA-CB-CG	8.61	135.09	115.30
55	DA	1131	G	N9-C1'-C2'	8.56	125.13	114.00
49	CV	6	LYS	CA-C-N	-8.55	98.39	117.20
55	DA	2665	A	C5-C6-N6	-8.54	116.87	123.70
55	DA	654(M)	C	N1-C1'-C2'	8.50	125.05	114.00
57	DY	51	LEU	CA-CB-CG	-8.50	95.76	115.30
1	AA	788	A	N9-C1'-C2'	8.49	125.04	114.00
55	DA	2609	U	N1-C1'-C2'	8.47	125.01	114.00
55	DA	945	A	N3-C4-N9	8.45	134.16	127.40
57	DY	111	LEU	CA-CB-CG	8.45	134.73	115.30
26	D4	18	CYS	CA-CB-SG	8.41	129.14	114.00
55	DA	2032	G	N9-C1'-C2'	8.38	124.90	114.00
57	DY	51	LEU	CB-CG-CD1	8.36	125.22	111.00
55	DA	945	A	C4-C5-N7	8.32	114.86	110.70
1	AA	654(M)	C	N1-C1'-C2'	8.26	124.74	114.00
55	DA	421	U	N1-C1'-C2'	8.24	124.71	114.00
1	AA	2311	A	N9-C1'-C2'	8.18	124.63	114.00
55	DA	1359	A	C3'-C2'-C1'	-8.17	94.96	101.50
57	DY	35	LYS	N-CA-C	-8.16	88.98	111.00
55	DA	371	A	N9-C1'-C2'	8.15	124.59	114.00
1	AA	323	G	O4'-C1'-N9	8.14	114.71	108.20
55	DA	654(I)	C	N1-C1'-C2'	8.12	124.56	114.00
55	DA	1992	G	C2'-C3'-O3'	8.10	127.33	109.50
54	CA	575	G	N9-C1'-C2'	8.09	124.52	114.00
55	DA	945	A	C4-N9-C1'	8.09	140.86	126.30
56	DJ	17	VAL	CA-CB-CG1	8.06	123.00	110.90
54	CA	1003	G	N9-C1'-C2'	-8.06	103.14	112.00
55	DA	2311	A	N9-C1'-C2'	8.05	124.47	114.00
1	AA	801	G	N9-C1'-C2'	8.04	124.45	114.00
1	AA	222	A	N9-C1'-C2'	8.03	124.44	114.00
1	AA	1372	U	C3'-C2'-C1'	-8.03	95.08	101.50
55	DA	2307	G	C1'-O4'-C4'	-8.01	103.49	109.90
1	AA	70	G	N9-C1'-C2'	8.00	124.41	114.00
54	CA	889	A	N9-C1'-C2'	8.00	124.40	114.00
1	AA	1247	A	N9-C1'-C2'	7.99	124.38	114.00
1	AA	1372	U	O4'-C1'-N1	-7.98	101.81	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
54	CA	1502	A	N9-C1'-C2'	7.98	124.37	114.00
54	CA	511	C	N1-C1'-C2'	7.98	124.37	114.00
1	AA	1252	G	N9-C1'-C2'	7.98	124.37	114.00
1	AA	1992	G	N9-C1'-C2'	7.94	124.33	114.00
55	DA	2656	U	C5-C4-O4	-7.93	121.14	125.90
1	AA	573	G	N9-C1'-C2'	7.92	124.30	114.00
55	DA	788	A	N9-C1'-C2'	7.91	124.29	114.00
1	AA	895	U	N3-C4-O4	-7.89	113.88	119.40
55	DA	2517	C	N1-C1'-C2'	7.88	124.24	114.00
55	DA	1781	C	N1-C1'-C2'	7.85	124.21	114.00
55	DA	249	C	N1-C1'-C2'	7.84	124.20	114.00
54	CA	1542	U	C4'-C3'-O3'	7.83	128.66	113.00
55	DA	2286	A	N9-C1'-C2'	7.82	124.17	114.00
55	DA	1544	C	O4'-C1'-N1	7.79	114.43	108.20
58	DL	140	GLY	N-CA-C	-7.77	93.68	113.10
31	BA	367	U	C1'-O4'-C4'	-7.75	103.70	109.90
21	AV	61	LEU	C-N-CD	-7.75	103.55	120.60
55	DA	1252	G	N9-C1'-C2'	7.71	124.02	114.00
55	DA	1380	G	O4'-C1'-N9	-7.71	102.03	108.20
55	DA	1372	U	O4'-C1'-N1	-7.70	102.04	108.20
55	DA	1397	U	C2'-C3'-O3'	7.69	126.42	109.50
55	DA	301	G	N9-C1'-C2'	7.67	123.97	114.00
55	DA	70	G	C2'-C3'-O3'	7.67	126.37	109.50
21	DV	176	PRO	C-N-CD	-7.62	103.84	120.60
54	CA	246	A	N9-C1'-C2'	7.61	123.89	114.00
54	CA	367	U	N1-C1'-C2'	7.59	123.87	114.00
55	DA	1397	U	N1-C1'-C2'	7.59	123.86	114.00
55	DA	529	A	N9-C1'-C2'	7.57	123.84	114.00
56	DJ	3	LEU	CB-CG-CD1	7.54	123.81	111.00
57	DY	138	LEU	N-CA-C	-7.54	90.66	111.00
54	CA	1064	G	N9-C1'-C2'	7.50	123.74	114.00
49	CV	6	LYS	O-C-N	7.49	134.68	122.70
53	B1	53	U	O4'-C1'-N1	7.45	114.16	108.20
54	CA	13	U	N1-C1'-C2'	7.43	123.66	114.00
1	AA	2835	A	N9-C1'-C2'	7.42	123.64	114.00
55	DA	70	G	N9-C1'-C2'	7.41	123.63	114.00
55	DA	199	A	N9-C1'-C2'	7.41	123.63	114.00
54	CA	1542	U	N1-C1'-C2'	-7.40	103.86	112.00
1	AA	83	G	N9-C1'-C2'	7.39	123.60	114.00
55	DA	383	U	N1-C2-O2	7.37	127.96	122.80
55	DA	654(S)	G	N9-C1'-C2'	-7.36	103.91	112.00
55	DA	1197	G	C5-C6-O6	7.36	133.02	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
48	BU	18	ARG	N-CA-C	-7.30	91.30	111.00
1	AA	1379	A	N9-C1'-C2'	7.29	123.47	114.00
55	DA	227	A	N9-C1'-C2'	7.28	123.46	114.00
55	DA	913	U	N1-C1'-C2'	7.28	123.46	114.00
1	AA	2681	C	N1-C1'-C2'	7.27	123.45	114.00
55	DA	139	G	N9-C1'-C2'	7.27	123.45	114.00
54	CA	1054	C	N1-C1'-C2'	7.26	123.44	114.00
55	DA	1372	U	N3-C4-C5	7.26	118.96	114.60
54	CA	702	A	N9-C1'-C2'	7.25	123.43	114.00
56	DI	27	LEU	CA-CB-CG	7.25	131.98	115.30
55	DA	801	G	N9-C1'-C2'	7.24	123.41	114.00
55	DA	2585	U	N1-C1'-C2'	7.24	123.41	114.00
57	DY	67	GLY	N-CA-C	7.23	131.17	113.10
2	DB	95	U	N1-C2-O2	7.21	127.84	122.80
54	CA	794	A	C4-N9-C1'	7.21	139.27	126.30
55	DA	1634	A	N9-C1'-C2'	7.20	123.36	114.00
1	AA	1397	U	C2'-C3'-O3'	7.19	125.32	109.50
31	BA	1504	G	N9-C1'-C2'	7.17	123.32	114.00
31	BA	246	A	N9-C1'-C2'	7.17	123.31	114.00
1	AA	2566	A	N9-C1'-C2'	7.16	123.31	114.00
55	DA	2345	G	N9-C1'-C2'	7.16	123.31	114.00
55	DA	1653	G	C2'-C3'-O3'	7.14	125.20	109.50
54	CA	1201	A	N9-C1'-C2'	7.13	123.27	114.00
1	AA	1652	A	C2'-C3'-O3'	7.12	125.18	109.50
55	DA	1341	U	N1-C1'-C2'	7.12	123.25	114.00
55	DA	526	A	N9-C1'-C2'	7.11	123.24	114.00
55	DA	945	A	O4'-C1'-N9	7.11	113.89	108.20
55	DA	1992	G	N9-C1'-C2'	7.11	123.24	114.00
1	AA	1992	G	C2'-C3'-O3'	7.09	125.10	109.50
54	CA	1498	U	C2'-C3'-O3'	7.08	125.08	109.50
55	DA	739	G	N9-C1'-C2'	7.07	123.19	114.00
57	DY	51	LEU	CB-CA-C	-7.06	96.79	110.20
55	DA	1212	G	C2'-C3'-O3'	7.04	125.00	109.50
55	DA	2391	G	N9-C1'-C2'	7.04	123.15	114.00
1	AA	1566	A	N9-C1'-C2'	7.02	123.13	114.00
31	BA	575	G	N9-C1'-C2'	7.02	123.12	114.00
55	DA	945	A	C3'-C2'-C1'	-7.02	95.89	101.50
1	AA	1329	U	N1-C1'-C2'	7.00	123.11	114.00
55	DA	1272	A	O4'-C1'-N9	7.00	113.80	108.20
1	AA	352	G	N9-C1'-C2'	7.00	123.10	114.00
4	DE	21	VAL	C-N-CD	-6.99	105.23	120.60
54	CA	794	A	C8-N9-C1'	-6.98	115.14	127.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
31	BA	1201	A	N9-C1'-C2'	6.96	123.05	114.00
31	BA	7	G	N9-C1'-C2'	6.95	123.03	114.00
55	DA	685	A	N9-C1'-C2'	6.94	123.02	114.00
31	BA	1529	G	O4'-C1'-N9	6.94	113.75	108.20
54	CA	518	C	N1-C1'-C2'	6.93	123.02	114.00
57	DY	19	ARG	NE-CZ-NH2	6.92	123.76	120.30
1	AA	1609	A	N9-C1'-C2'	6.90	122.97	114.00
1	AA	2238	G	N9-C1'-C2'	6.89	122.95	114.00
49	CV	85	LYS	N-CA-C	-6.88	92.42	111.00
1	AA	1966	A	N9-C1'-C2'	6.88	122.94	114.00
1	AA	322	A	O4'-C1'-N9	6.87	113.70	108.20
1	AA	654(S)	G	N9-C1'-C2'	-6.86	104.46	112.00
1	AA	371	A	N9-C1'-C2'	6.84	122.89	114.00
55	DA	1022	G	N9-C1'-C2'	6.83	122.88	114.00
55	DA	1385	G	N9-C1'-C2'	6.81	122.86	114.00
1	AA	2345	G	N9-C1'-C2'	6.80	122.85	114.00
31	BA	794	A	C4-N9-C1'	6.78	138.51	126.30
54	CA	788	U	C2-N3-C4	-6.78	122.94	127.00
1	AA	1428	C	N1-C1'-C2'	6.76	122.79	114.00
52	CB	74	C	P-O3'-C3'	6.75	127.80	119.70
55	DA	1250	G	N9-C1'-C2'	6.75	122.78	114.00
1	AA	1780	A	N9-C1'-C2'	6.75	122.77	114.00
54	CA	717	C	N1-C1'-C2'	6.75	122.77	114.00
55	DA	72	U	N1-C1'-C2'	6.74	122.76	114.00
55	DA	2275	C	N1-C1'-C2'	6.73	122.75	114.00
1	AA	1800	C	N1-C1'-C2'	6.69	122.69	114.00
1	AA	2866	U	N1-C1'-C2'	6.69	122.69	114.00
1	AA	1378	A	P-O3'-C3'	6.68	127.72	119.70
1	AA	2448	A	N9-C1'-C2'	6.67	122.67	114.00
55	DA	2286	A	O4'-C1'-N9	6.67	113.53	108.20
1	AA	2447	G	N9-C1'-C2'	6.66	122.65	114.00
1	AA	60	G	N9-C1'-C2'	6.65	122.65	114.00
55	DA	1647	G	N9-C1'-C2'	6.65	122.64	114.00
31	BA	197	A	N9-C1'-C2'	6.65	122.64	114.00
57	DY	111	LEU	CA-C-N	6.65	131.82	117.20
55	DA	99	U	N1-C1'-C2'	6.64	122.64	114.00
57	DY	128	LEU	C-N-CD	-6.64	105.99	120.60
55	DA	856	C	C2'-C3'-O3'	6.64	124.32	113.70
54	CA	1498	U	N1-C1'-C2'	6.62	122.61	114.00
1	AA	421	U	N1-C1'-C2'	6.62	122.61	114.00
55	DA	2060	A	C5'-C4'-O4'	-6.61	101.16	109.10
58	DL	23	VAL	N-CA-C	-6.61	93.15	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
55	DA	1786	A	N9-C1'-C2'	6.61	122.59	114.00
31	BA	8	A	N9-C1'-C2'	6.59	122.57	114.00
1	AA	2656	U	C5-C4-O4	-6.59	121.95	125.90
21	AV	176	PRO	C-N-CD	-6.59	106.11	120.60
1	AA	571	A	N9-C1'-C2'	6.57	122.54	114.00
54	CA	653	A	N9-C1'-C2'	6.56	122.53	114.00
1	AA	120	U	N1-C1'-C2'	6.54	122.51	114.00
55	DA	793	A	N9-C1'-C2'	6.54	122.50	114.00
31	BA	794	A	C8-N9-C1'	-6.54	115.93	127.70
1	AA	607	U	N3-C4-O4	-6.53	114.83	119.40
55	DA	2820	A	N9-C1'-C2'	6.52	122.47	114.00
54	CA	1159	U	N1-C1'-C2'	6.51	122.47	114.00
55	DA	1838	C	N1-C1'-C2'	6.51	122.47	114.00
55	DA	455	C	N1-C1'-C2'	6.50	122.45	114.00
56	DI	29	GLU	CA-CB-CG	-6.48	99.15	113.40
55	DA	1954	G	N9-C1'-C2'	6.46	122.40	114.00
1	AA	2060	A	N9-C1'-C2'	6.46	122.40	114.00
58	DL	47	ASN	N-CA-C	-6.46	93.56	111.00
55	DA	97	C	C5'-C4'-C3'	-6.46	105.67	116.00
54	CA	47	C	N1-C1'-C2'	6.45	122.38	114.00
55	DA	829	A	N9-C1'-C2'	6.44	122.38	114.00
55	DA	1925	C	C2-N3-C4	-6.44	116.68	119.90
54	CA	872	A	O4'-C1'-N9	6.44	113.35	108.20
1	AA	1786	A	N9-C1'-C2'	6.44	122.37	114.00
1	AA	1698	A	N9-C1'-C2'	6.40	122.32	114.00
15	DR	59	THR	N-CA-C	-6.39	93.74	111.00
31	BA	190	G	N9-C1'-C2'	6.37	122.28	114.00
1	AA	2286	A	N9-C1'-C2'	6.35	122.25	114.00
1	AA	2645	G	N9-C1'-C2'	6.34	122.24	114.00
55	DA	2726	U	N1-C1'-C2'	6.33	122.23	114.00
1	AA	1380	G	O4'-C1'-N9	-6.33	103.14	108.20
31	BA	1502	A	N9-C1'-C2'	6.33	122.22	114.00
1	AA	1818	U	N1-C1'-C2'	6.32	122.22	114.00
58	DL	27	LEU	CA-CB-CG	6.31	129.82	115.30
34	BG	33	MET	N-CA-C	-6.31	93.96	111.00
55	DA	434	U	N1-C1'-C2'	6.30	122.20	114.00
1	AA	2873	A	N9-C1'-C2'	6.30	122.19	114.00
54	CA	993	G	N9-C1'-C2'	6.29	122.18	114.00
55	DA	1372	U	C5-C4-O4	-6.28	122.13	125.90
55	DA	2725	A	N9-C1'-C2'	6.28	122.16	114.00
1	AA	1342	A	O3'-P-O5'	-6.27	92.08	104.00
55	DA	1934	C	C5'-C4'-O4'	-6.27	101.58	109.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	AA	2665	A	C5-C6-N6	-6.26	118.69	123.70
55	DA	1698	A	C2'-C3'-O3'	6.25	123.71	113.70
55	DA	457	A	N9-C1'-C2'	6.25	122.13	114.00
55	DA	1534	G	N9-C1'-C2'	-6.25	105.12	112.00
55	DA	897	C	C2-N1-C1'	6.25	125.67	118.80
1	AA	1385	G	N9-C1'-C2'	6.24	122.12	114.00
31	BA	717	C	N1-C1'-C2'	6.24	122.11	114.00
1	AA	2458	G	N9-C1'-C2'	6.24	122.11	114.00
55	DA	654(H)	G	C4-N9-C1'	6.22	134.59	126.50
31	BA	328	C	N1-C1'-C2'	6.21	122.08	114.00
55	DA	2665	A	C6-N1-C2	-6.21	114.87	118.60
55	DA	222	A	N9-C1'-C2'	6.21	122.07	114.00
55	DA	2665	A	C5-C6-N1	6.20	120.80	117.70
55	DA	5	A	OP1-P-OP2	6.19	128.88	119.60
55	DA	1950	G	O4'-C1'-N9	6.19	113.15	108.20
55	DA	654(H)	G	O4'-C1'-N9	-6.19	103.25	108.20
55	DA	621	A	C5-C6-N6	-6.19	118.75	123.70
55	DA	1955	U	N1-C1'-C2'	6.18	122.03	114.00
55	DA	1544	C	O4'-C1'-C2'	6.17	113.15	107.60
31	BA	653	A	N9-C1'-C2'	6.17	122.02	114.00
1	AA	2690	C	N1-C1'-C2'	6.16	122.01	114.00
55	DA	1799	G	N9-C1'-C2'	6.15	122.00	114.00
1	AA	1372	U	O3'-P-O5'	6.14	115.67	104.00
58	DL	52	ILE	CB-CA-C	-6.14	99.33	111.60
1	AA	323	G	C1'-O4'-C4'	-6.12	105.00	109.90
55	DA	2789	C	C1'-O4'-C4'	-6.12	105.00	109.90
57	DY	80	VAL	CG1-CB-CG2	-6.12	101.11	110.90
20	AU	20	TYR	N-CA-C	-6.11	94.49	111.00
55	DA	1081	U	N1-C1'-C2'	6.11	121.95	114.00
55	DA	1966	A	N9-C1'-C2'	6.11	121.94	114.00
11	DO	67	MET	N-CA-C	-6.11	94.52	111.00
54	CA	328	C	N1-C1'-C2'	6.10	121.93	114.00
3	DD	229	VAL	CB-CA-C	-6.10	99.81	111.40
55	DA	1307	A	C5'-C4'-C3'	-6.09	106.26	116.00
55	DA	2542	A	N9-C1'-C2'	6.09	121.91	114.00
1	AA	621	A	C5-C6-N6	-6.08	118.83	123.70
1	AA	2873	A	O4'-C1'-N9	6.08	113.07	108.20
55	DA	621	A	C6-N1-C2	-6.08	114.95	118.60
55	DA	2665	A	N1-C6-N6	6.07	122.24	118.60
55	DA	793	A	C4'-C3'-O3'	-6.07	96.66	109.40
21	DV	153	SER	N-CA-CB	-6.07	101.40	110.50
1	AA	2712	U	O4'-C1'-N1	6.07	113.05	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
55	DA	1567	A	N9-C1'-C2'	6.06	121.88	114.00
54	CA	509	A	C2'-C3'-O3'	6.05	123.39	113.70
1	AA	1302	A	N9-C1'-C2'	6.05	121.87	114.00
1	AA	2789	C	C1'-O4'-C4'	-6.04	105.06	109.90
1	AA	301	G	N9-C1'-C2'	6.04	121.85	114.00
56	DJ	23	LEU	N-CA-C	-6.04	94.70	111.00
1	AA	5	A	OP1-P-OP2	6.03	128.65	119.60
4	DE	118	LYS	N-CA-C	-6.03	94.72	111.00
23	AZ	36	GLY	N-CA-C	6.03	128.17	113.10
34	BG	21	LEU	CA-CB-CG	-6.02	101.45	115.30
55	DA	2702	U	N1-C1'-C2'	6.02	121.83	114.00
55	DA	2681	C	C2'-C3'-O3'	6.01	123.31	113.70
55	DA	896	A	N9-C1'-C2'	6.00	121.80	114.00
55	DA	1602	U	N1-C1'-C2'	6.00	121.80	114.00
55	DA	1427	A	N9-C1'-C2'	5.99	121.79	114.00
55	DA	573	G	N9-C1'-C2'	5.98	121.78	114.00
1	AA	2702	U	N1-C1'-C2'	5.98	121.77	114.00
1	AA	446	G	N9-C1'-C2'	5.98	121.77	114.00
55	DA	479	A	N9-C1'-C2'	5.98	121.77	114.00
26	D4	36	CYS	CA-CB-SG	5.98	124.76	114.00
1	AA	199	A	N9-C1'-C2'	5.97	121.77	114.00
26	D4	39	CYS	CA-CB-SG	5.97	124.74	114.00
7	DH	124	GLU	N-CA-C	-5.97	94.89	111.00
54	CA	50	A	N9-C1'-C2'	5.96	121.75	114.00
31	BA	243	A	N9-C1'-C2'	5.96	121.74	114.00
57	DY	39	ALA	N-CA-C	-5.95	94.93	111.00
55	DA	1800	C	N1-C1'-C2'	5.95	121.73	114.00
1	AA	2285	C	P-O3'-C3'	-5.94	112.57	119.70
31	BA	1528	U	N1-C1'-C2'	5.94	121.72	114.00
1	AA	2051	A	N9-C1'-C2'	5.94	121.72	114.00
21	DV	109	ALA	N-CA-C	-5.93	94.98	111.00
55	DA	995	C	N1-C1'-C2'	5.93	121.71	114.00
1	AA	930	U	N1-C1'-C2'	5.92	121.70	114.00
55	DA	27	G	C5'-C4'-O4'	-5.92	102.00	109.10
55	DA	2468	G	O4'-C1'-N9	5.92	112.93	108.20
1	AA	1970	A	N9-C1'-C2'	5.91	121.69	114.00
52	BB	74	C	O4'-C1'-N1	5.90	112.92	108.20
55	DA	1359	A	C4-C5-N7	5.89	113.64	110.70
1	AA	603	A	N9-C1'-C2'	5.88	121.65	114.00
55	DA	791	C	N1-C1'-C2'	5.88	121.65	114.00
1	AA	2656	U	N1-C1'-C2'	-5.88	105.53	112.00
53	C1	48	U	N1-C1'-C2'	5.88	121.64	114.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
55	DA	930	U	N1-C1'-C2'	5.87	121.63	114.00
1	AA	989	G	N9-C1'-C2'	5.87	121.63	114.00
55	DA	2681	C	N1-C1'-C2'	5.87	121.63	114.00
34	CG	31	CYS	N-CA-C	-5.86	95.17	111.00
55	DA	654(H)	G	C8-N9-C1'	-5.86	119.38	127.00
1	AA	739	G	N9-C1'-C2'	5.86	121.62	114.00
52	BC	43	C	C2'-C3'-O3'	5.86	123.07	113.70
55	DA	1025	G	N9-C1'-C2'	5.86	121.61	114.00
55	DA	2490	G	N9-C1'-C2'	5.86	121.61	114.00
1	AA	1762	A	N9-C1'-C2'	5.85	121.61	114.00
1	AA	125	G	N9-C1'-C2'	5.85	121.60	114.00
55	DA	2426	A	N9-C1'-C2'	5.84	121.59	114.00
55	DA	2205	C	C5'-C4'-C3'	-5.84	106.66	116.00
55	DA	196	A	O4'-C1'-N9	5.83	112.86	108.20
55	DA	932	G	O4'-C1'-N9	5.82	112.86	108.20
55	DA	2866	U	C2'-C3'-O3'	5.80	122.99	113.70
31	BA	595	G	N9-C1'-C2'	5.79	121.53	114.00
1	AA	323	G	N9-C1'-C2'	5.79	121.53	114.00
54	CA	1124	G	N9-C1'-C2'	5.79	121.53	114.00
55	DA	1359	A	N7-C8-N9	5.79	116.69	113.80
11	AO	115	LEU	CA-CB-CG	5.79	128.60	115.30
1	AA	2873	A	C1'-O4'-C4'	-5.78	105.27	109.90
55	DA	1566	A	N9-C1'-C2'	5.78	121.52	114.00
55	DA	60	G	N9-C1'-C2'	5.78	121.51	114.00
1	AA	1385	G	O4'-C1'-N9	5.78	112.82	108.20
1	AA	2572	A	N9-C1'-C2'	5.77	121.50	114.00
55	DA	458	G	N9-C1'-C2'	5.76	121.49	114.00
55	DA	1607	C	N1-C1'-C2'	5.76	121.49	114.00
1	AA	913	U	N1-C1'-C2'	5.75	121.48	114.00
55	DA	2198	A	N9-C1'-C2'	5.75	121.47	114.00
58	DL	76	TYR	CA-CB-CG	5.75	124.32	113.40
1	AA	1359	A	C6-C5-N7	-5.74	128.28	132.30
55	DA	624	C	C5'-C4'-C3'	-5.74	106.81	116.00
53	B1	56	U	N1-C1'-C2'	5.73	121.45	114.00
55	DA	2507	C	C5'-C4'-O4'	-5.73	102.22	109.10
58	DL	132	ARG	NE-CZ-NH2	-5.73	117.44	120.30
1	AA	15	G	N9-C1'-C2'	-5.72	105.71	112.00
1	AA	2275	C	N1-C1'-C2'	5.72	121.44	114.00
1	AA	2490	G	O4'-C1'-N9	5.72	112.77	108.20
54	CA	1190	G	N9-C1'-C2'	5.71	121.43	114.00
54	CA	243	A	N9-C1'-C2'	5.71	121.43	114.00
1	AA	2346	A	N9-C1'-C2'	5.71	121.42	114.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	AA	2384	G	N9-C1'-C2'	5.71	121.42	114.00
55	DA	2490	G	O4'-C1'-N9	5.71	112.77	108.20
4	DE	58	ARG	N-CA-C	-5.71	95.59	111.00
1	AA	562	U	N1-C1'-C2'	5.70	121.41	114.00
1	AA	2506	U	N1-C1'-C2'	5.70	121.42	114.00
1	AA	322	A	N9-C4-C5	5.70	108.08	105.80
54	CA	818	G	N9-C1'-C2'	5.70	121.41	114.00
31	BA	1064	G	N9-C1'-C2'	5.70	121.41	114.00
55	DA	830	G	C5'-C4'-O4'	-5.70	102.26	109.10
1	AA	955	C	C5'-C4'-C3'	-5.69	106.89	116.00
2	DB	95	U	N3-C4-O4	-5.69	115.42	119.40
58	DL	70	LYS	N-CA-C	-5.68	95.66	111.00
54	CA	559	A	N9-C1'-C2'	5.68	121.38	114.00
2	DB	81	G	N1-C6-O6	-5.67	116.50	119.90
55	DA	1693	U	C5'-C4'-C3'	-5.67	106.93	116.00
1	AA	448	U	N1-C1'-C2'	5.66	121.36	114.00
54	CA	630	G	N9-C1'-C2'	-5.66	105.77	112.00
55	DA	654(H)	G	C5'-C4'-O4'	5.66	115.89	109.10
21	DV	200	GLY	CA-C-O	-5.65	110.43	120.60
1	AA	654(M)	C	C2-N1-C1'	5.64	125.01	118.80
55	DA	687	C	C5'-C4'-C3'	-5.64	106.97	116.00
1	AA	5	A	O5'-P-OP2	-5.64	100.62	105.70
13	D0	58	GLY	N-CA-C	5.64	127.20	113.10
1	AA	1397	U	N1-C1'-C2'	5.64	121.33	114.00
57	DY	50	ARG	NE-CZ-NH1	-5.64	117.48	120.30
1	AA	2391	G	N9-C1'-C2'	5.63	121.33	114.00
58	DL	52	ILE	CB-CG1-CD1	-5.63	98.13	113.90
52	BB	9	A	C2'-C3'-O3'	5.63	122.71	113.70
54	CA	801	U	C5'-C4'-O4'	-5.62	102.36	109.10
1	AA	2062	A	N9-C1'-C2'	5.61	121.30	114.00
55	DA	989	G	O4'-C1'-N9	5.61	112.69	108.20
31	BA	251	G	N9-C1'-C2'	5.61	121.29	114.00
55	DA	2425	A	O4'-C1'-N9	5.61	112.68	108.20
55	DA	1324	G	N9-C1'-C2'	5.60	121.28	114.00
54	CA	115	G	N9-C1'-C2'	5.60	121.28	114.00
54	CA	97	U	N1-C2-O2	5.59	126.72	122.80
11	DO	34	GLY	N-CA-C	5.59	127.07	113.10
55	DA	2311	A	O4'-C1'-C2'	-5.58	100.22	105.80
1	AA	1372	U	C1'-O4'-C4'	-5.58	105.44	109.90
55	DA	196	A	C1'-O4'-C4'	-5.58	105.44	109.90
57	DY	130	THR	N-CA-C	-5.58	95.94	111.00
31	BA	815	A	N9-C1'-C2'	5.57	121.25	114.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
55	DA	323	G	O4'-C1'-N9	5.57	112.66	108.20
31	BA	109	A	N9-C1'-C2'	5.57	121.24	114.00
55	DA	654(M)	C	C2-N1-C1'	5.55	124.91	118.80
55	DA	2311	A	C3'-C2'-C1'	-5.55	97.06	101.50
56	DJ	30	ALA	CA-C-O	5.55	131.76	120.10
4	AE	61	ARG	N-CA-C	-5.55	96.02	111.00
55	DA	896	A	C4-N9-C1'	-5.55	116.32	126.30
55	DA	2517	C	C2'-C3'-O3'	5.55	122.58	113.70
55	DA	311	A	N9-C1'-C2'	5.54	121.21	114.00
1	AA	2406	U	N1-C1'-C2'	5.54	121.21	114.00
55	DA	74	A	C4-N9-C1'	-5.54	116.34	126.30
55	DA	1128	A	N9-C1'-C2'	5.54	121.20	114.00
1	AA	829	A	N9-C1'-C2'	5.53	121.19	114.00
56	DI	10	GLU	N-CA-C	-5.53	96.07	111.00
57	DY	134	LEU	CA-CB-CG	5.53	128.02	115.30
1	AA	1342	A	OP2-P-O3'	-5.52	93.05	105.20
1	AA	1372	U	N1-C2-N3	-5.52	111.59	114.90
26	D4	19	GLY	N-CA-C	-5.52	99.29	113.10
1	AA	2656	U	N3-C4-O4	-5.52	115.54	119.40
31	BA	818	G	N9-C1'-C2'	5.52	121.17	114.00
52	CD	45	U	C1'-O4'-C4'	-5.51	105.49	109.90
55	DA	2469	A	C1'-O4'-C4'	-5.51	105.49	109.90
55	DA	74	A	C8-N9-C1'	5.51	137.61	127.70
1	AA	72	U	N1-C1'-C2'	5.50	121.15	114.00
1	AA	1272	A	O4'-C1'-N9	5.50	112.60	108.20
55	DA	125	G	C5'-C4'-C3'	-5.50	107.20	116.00
54	CA	630	G	C3'-C2'-C1'	-5.50	97.10	101.50
31	BA	250	A	N9-C1'-C2'	5.49	121.14	114.00
55	DA	2346	A	N9-C1'-C2'	5.49	121.14	114.00
55	DA	1407	C	C5'-C4'-C3'	-5.49	107.22	116.00
55	DA	2307	G	C8-N9-C4	-5.49	104.20	106.40
55	DA	1302	A	N9-C1'-C2'	5.48	121.13	114.00
56	DJ	6	GLU	OE1-CD-OE2	5.48	129.88	123.30
31	BA	559	A	N9-C1'-C2'	5.48	121.12	114.00
31	BA	566	G	N9-C1'-C2'	5.48	121.12	114.00
55	DA	83	G	N9-C1'-C2'	5.48	121.12	114.00
1	AA	1598	C	C5'-C4'-C3'	-5.47	107.24	116.00
2	DB	81	G	C5-C6-O6	5.47	131.88	128.60
1	AA	1380	G	P-O5'-C5'	-5.47	112.15	120.90
12	DP	10	ARG	N-CA-C	-5.47	96.24	111.00
54	CA	1502	A	O4'-C1'-N9	5.45	112.56	108.20
55	DA	1379	A	P-O3'-C3'	5.45	126.24	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	AA	2311	A	C3'-C2'-C1'	-5.45	97.14	101.50
1	AA	531	C	C1'-O4'-C4'	-5.44	105.55	109.90
55	DA	67	U	C2-N3-C4	5.44	130.26	127.00
1	AA	215	G	N9-C1'-C2'	5.44	121.07	114.00
55	DA	2266	A	N9-C1'-C2'	5.44	121.07	114.00
54	CA	960	U	N1-C1'-C2'	5.43	121.06	114.00
1	AA	249	C	O4'-C1'-N1	5.42	112.54	108.20
55	DA	2320	A	N9-C1'-C2'	5.41	121.04	114.00
1	AA	1962	C	N1-C1'-C2'	5.41	121.04	114.00
1	AA	654(H)	G	C4-N9-C1'	5.41	133.53	126.50
55	DA	805	G	O4'-C1'-N9	5.41	112.53	108.20
42	CO	47	LYS	C-N-CD	-5.41	108.70	120.60
57	DY	72	ASP	CB-CG-OD1	5.41	123.17	118.30
55	DA	2249	U	N1-C1'-C2'	5.40	121.03	114.00
1	AA	403	U	N1-C1'-C2'	5.40	121.02	114.00
2	DB	44	G	N9-C1'-C2'	5.39	121.01	114.00
1	AA	856	C	C2'-C3'-O3'	5.38	122.31	113.70
1	AA	1344	G	N9-C1'-C2'	5.38	121.00	114.00
54	CA	752	G	N9-C1'-C2'	5.38	120.99	114.00
55	DA	74	A	C5-C6-N6	5.38	128.00	123.70
55	DA	1558	A	N9-C1'-C2'	5.37	120.99	114.00
1	AA	1372	U	N3-C4-C5	5.37	117.82	114.60
34	BG	34	GLU	N-CA-C	-5.37	96.50	111.00
55	DA	784	A	N9-C1'-C2'	5.36	120.97	114.00
55	DA	897	C	O4'-C1'-N1	-5.36	103.91	108.20
54	CA	47	C	O4'-C1'-N1	5.36	112.48	108.20
55	DA	896	A	C8-N9-C1'	5.36	137.34	127.70
55	DA	2566	A	OP2-P-O3'	5.35	116.97	105.20
57	DY	30	GLN	N-CA-C	-5.35	96.56	111.00
54	CA	1322	C	N1-C1'-C2'	5.35	120.95	114.00
14	DQ	54	LEU	CA-CB-CG	5.35	127.60	115.30
58	DL	116	ASN	N-CA-C	-5.35	96.56	111.00
1	AA	249	C	C2'-C3'-O3'	-5.34	97.75	109.50
55	DA	1300	U	N1-C1'-C2'	5.34	120.95	114.00
55	DA	1616	A	O4'-C1'-N9	5.34	112.47	108.20
31	BA	960	U	N1-C1'-C2'	5.34	120.94	114.00
1	AA	1360	A	C5'-C4'-C3'	-5.34	107.46	116.00
54	CA	794	A	C6-N1-C2	-5.34	115.40	118.60
58	DL	121	GLU	N-CA-C	-5.34	96.59	111.00
1	AA	531	C	O4'-C1'-N1	5.34	112.47	108.20
58	DL	28	GLY	N-CA-C	-5.34	99.76	113.10
1	AA	2311	A	O4'-C1'-C2'	-5.33	100.47	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
54	CA	305	G	N9-C1'-C2'	5.33	120.92	114.00
1	AA	310	A	N9-C1'-C2'	5.32	120.92	114.00
34	BG	12	CYS	CA-CB-SG	5.32	123.58	114.00
55	DA	278	A	C2'-C3'-O3'	5.32	122.22	113.70
55	DA	788	A	OP2-P-O3'	5.32	116.91	105.20
55	DA	474	G	C2'-C3'-O3'	5.32	122.21	113.70
55	DA	1519	G	C5'-C4'-C3'	-5.32	107.49	116.00
54	CA	1003	G	C3'-C2'-C1'	-5.31	97.25	101.50
1	AA	4	C	O3'-P-O5'	5.31	114.09	104.00
34	CG	31	CYS	CA-CB-SG	5.31	123.56	114.00
55	DA	2524	G	C5'-C4'-C3'	-5.31	107.50	116.00
54	CA	1177	G	N9-C1'-C2'	5.31	120.90	114.00
55	DA	1281	G	C5'-C4'-O4'	-5.30	102.74	109.10
1	AA	1822	G	C5'-C4'-O4'	-5.30	102.74	109.10
54	CA	1299	A	N9-C1'-C2'	5.29	120.88	114.00
54	CA	410	G	C2'-C3'-O3'	5.29	122.16	113.70
1	AA	1372	U	C6-N1-C2	5.28	124.17	121.00
54	CA	47	C	C1'-O4'-C4'	-5.28	105.67	109.90
55	DA	503	A	N9-C1'-C2'	5.28	120.87	114.00
55	DA	101	G	N9-C1'-C2'	5.28	120.87	114.00
55	DA	2060	A	OP1-P-O3'	5.27	116.80	105.20
58	DL	12	LEU	CA-CB-CG	-5.27	103.19	115.30
55	DA	752	A	OP2-P-O3'	5.27	116.79	105.20
55	DA	829	A	C4'-C3'-O3'	-5.27	98.34	109.40
55	DA	2245	U	C5'-C4'-C3'	-5.27	107.57	116.00
21	DV	179	ASP	N-CA-C	-5.26	96.79	111.00
31	BA	819	A	N9-C1'-C2'	5.26	120.84	114.00
55	DA	897	C	C5'-C4'-O4'	5.26	115.41	109.10
55	DA	1359	A	C5-N7-C8	-5.26	101.27	103.90
58	DL	84	LEU	N-CA-C	-5.26	96.81	111.00
55	DA	926	A	C5'-C4'-C3'	-5.25	107.59	116.00
54	CA	279	A	N9-C1'-C2'	5.24	120.81	114.00
55	DA	1543	A	N9-C1'-C2'	5.24	120.81	114.00
55	DA	1791	A	O5'-P-OP1	-5.24	100.98	105.70
55	DA	1178	C	C2'-C3'-O3'	5.24	122.08	113.70
55	DA	1344	G	N9-C1'-C2'	5.23	120.80	114.00
11	DO	59	LEU	N-CA-C	-5.23	96.88	111.00
52	BB	10	G	C2'-C3'-O3'	5.23	122.07	113.70
54	CA	315	A	N9-C1'-C2'	5.23	120.80	114.00
8	DK	135	GLU	N-CA-C	5.22	125.10	111.00
55	DA	1701	A	C5'-C4'-C3'	-5.22	107.65	116.00
58	DL	38	VAL	N-CA-C	-5.21	96.93	111.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	AA	232	G	N9-C1'-C2'	5.20	120.77	114.00
44	CQ	40	CYS	CA-CB-SG	5.20	123.36	114.00
55	DA	1325	G	N9-C1'-C2'	5.20	120.76	114.00
55	DA	1900	A	N9-C1'-C2'	5.20	120.76	114.00
55	DA	788	A	C2'-C3'-O3'	5.20	122.02	113.70
57	DY	32	LEU	CA-CB-CG	5.19	127.25	115.30
1	AA	687	C	C5'-C4'-C3'	-5.19	107.69	116.00
1	AA	2426	A	N9-C1'-C2'	5.19	120.75	114.00
31	BA	1157	A	N9-C1'-C2'	5.19	120.75	114.00
1	AA	2346	A	C1'-O4'-C4'	-5.19	105.75	109.90
30	A8	32	LEU	CA-CB-CG	5.19	127.23	115.30
1	AA	241	A	N9-C1'-C2'	5.18	120.74	114.00
55	DA	2033	A	N9-C1'-C2'	5.18	120.74	114.00
1	AA	2250	G	N9-C1'-C2'	5.18	120.74	114.00
53	B1	50	U	C5'-C4'-C3'	-5.18	107.71	116.00
55	DA	1015	G	C5'-C4'-C3'	-5.18	107.71	116.00
34	BG	20	TYR	C-N-CA	5.17	134.64	121.70
55	DA	1791	A	C5'-C4'-C3'	-5.17	107.72	116.00
55	DA	474	G	N9-C1'-C2'	5.17	120.72	114.00
57	DY	136	ALA	N-CA-C	-5.17	97.03	111.00
31	BA	702	A	N9-C1'-C2'	5.17	120.72	114.00
55	DA	4	C	O3'-P-O5'	5.17	113.82	104.00
55	DA	1786	A	O4'-C1'-N9	5.17	112.33	108.20
55	DA	945	A	C4-C5-C6	5.16	119.58	117.00
1	AA	1359	A	C4-C5-N7	5.16	113.28	110.70
1	AA	1385	G	C1'-O4'-C4'	-5.16	105.77	109.90
55	DA	1694	C	C5'-C4'-C3'	-5.16	107.75	116.00
31	BA	209	U	N1-C1'-C2'	5.15	120.70	114.00
52	BB	74	C	C1'-O4'-C4'	-5.15	105.78	109.90
56	DJ	28	LYS	N-CA-C	-5.15	97.08	111.00
1	AA	2275	C	C2'-C3'-O3'	5.15	121.94	113.70
55	DA	2832	U	C2'-C3'-O3'	5.14	121.92	113.70
55	DA	454	A	C5'-C4'-C3'	-5.14	107.78	116.00
55	DA	539	G	C5'-C4'-C3'	-5.14	107.78	116.00
11	AO	65	ARG	N-CA-C	-5.14	97.13	111.00
52	CB	9	A	C2'-C3'-O3'	5.14	121.92	113.70
55	DA	1773	A	N9-C1'-C2'	-5.14	106.35	112.00
1	AA	532	A	N9-C1'-C2'	5.13	120.67	114.00
57	DY	100	ASN	N-CA-C	-5.13	97.14	111.00
26	A4	16	CYS	CA-CB-SG	5.13	123.23	114.00
31	BA	982	U	N1-C1'-C2'	5.13	120.67	114.00
26	D4	70	GLY	N-CA-C	5.12	125.91	113.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
55	DA	1925	C	O4'-C4'-C3'	-5.12	98.88	104.00
1	AA	2542	A	N9-C1'-C2'	5.12	120.66	114.00
55	DA	1143	A	N9-C1'-C2'	5.12	120.66	114.00
1	AA	1939	U	N1-C1'-C2'	5.12	120.65	114.00
3	AD	224	ALA	N-CA-C	-5.12	97.18	111.00
55	DA	1359	A	O4'-C1'-C2'	-5.12	100.68	105.80
26	D4	45	GLY	N-CA-C	-5.12	100.31	113.10
31	BA	366	C	N1-C1'-C2'	5.12	120.65	114.00
1	AA	2061	G	N9-C1'-C2'	5.11	120.65	114.00
55	DA	530	G	N9-C1'-C2'	5.11	120.65	114.00
1	AA	1694	C	N1-C1'-C2'	5.11	120.64	114.00
55	DA	1294	U	C5'-C4'-C3'	-5.11	107.83	116.00
31	BA	130	A	N9-C1'-C2'	5.10	120.64	114.00
54	CA	1506	U	N1-C1'-C2'	5.10	120.64	114.00
1	AA	50	U	N1-C1'-C2'	5.10	120.63	114.00
55	DA	1609	A	C4'-C3'-O3'	-5.10	98.69	109.40
55	DA	141	A	N9-C1'-C2'	5.10	120.63	114.00
55	DA	728	G	C5'-C4'-O4'	-5.10	102.98	109.10
55	DA	2497	A	C2'-C3'-O3'	-5.10	98.28	109.50
54	CA	788	U	C5-C4-O4	-5.10	122.84	125.90
9	DM	114	ARG	N-CA-C	-5.10	97.24	111.00
1	AA	1372	U	C6-N1-C1'	-5.09	114.07	121.20
55	DA	989	G	N9-C1'-C2'	5.09	120.62	114.00
54	CA	109	A	N9-C1'-C2'	5.08	120.61	114.00
55	DA	387	U	O5'-P-OP1	-5.08	101.12	105.70
55	DA	1947	C	C5'-C4'-C3'	-5.08	107.87	116.00
1	AA	226	G	N9-C1'-C2'	5.07	120.59	114.00
56	DJ	12	LEU	CB-CG-CD2	5.07	119.62	111.00
1	AA	1544	C	N1-C1'-C2'	5.07	120.59	114.00
54	CA	760	G	N9-C1'-C2'	-5.07	106.43	112.00
1	AA	1786	A	C1'-O4'-C4'	-5.07	105.85	109.90
55	DA	2035	G	N9-C1'-C2'	5.07	120.59	114.00
53	C1	33	G	O4'-C1'-N9	5.06	112.25	108.20
55	DA	196	A	N9-C1'-C2'	5.06	120.58	114.00
21	AV	117	LEU	CA-CB-CG	5.06	126.94	115.30
1	AA	74	A	C4-N9-C1'	-5.05	117.20	126.30
31	BA	1159	U	N1-C1'-C2'	5.05	120.57	114.00
43	BP	84	ILE	N-CA-C	-5.05	97.36	111.00
31	BA	279	A	N9-C1'-C2'	5.05	120.57	114.00
34	CG	13	ARG	N-CA-C	5.05	124.63	111.00
54	CA	327	A	N9-C1'-C2'	5.04	120.56	114.00
55	DA	1249	U	C2-N3-C4	-5.04	123.98	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
55	DA	1653	G	N9-C1'-C2'	5.04	120.55	114.00
1	AA	2032	G	N9-C1'-C2'	5.04	120.55	114.00
55	DA	1615	C	N1-C1'-C2'	5.04	120.55	114.00
1	AA	1372	U	N1-C1'-C2'	-5.03	106.47	112.00
55	DA	2754	U	C5'-C4'-C3'	-5.03	107.95	116.00
1	AA	227	A	N9-C1'-C2'	5.03	120.53	114.00
55	DA	1371	G	P-O3'-C3'	5.03	125.73	119.70
55	DA	2458	G	C2'-C3'-O3'	-5.02	98.45	109.50
55	DA	2447	G	C4'-C3'-O3'	-5.02	98.86	109.40
1	AA	1762	A	O4'-C1'-C2'	-5.02	100.78	105.80
5	AF	83	PHE	N-CA-C	5.02	124.56	111.00
1	AA	654(H)	G	C8-N9-C1'	-5.02	120.47	127.00
55	DA	1686	C	C5'-C4'-C3'	-5.02	107.97	116.00
55	DA	1396	U	N1-C1'-C2'	5.01	120.52	114.00
31	BA	7	G	C1'-O4'-C4'	-5.01	105.89	109.90
55	DA	2094	G	C5'-C4'-C3'	-5.01	107.99	116.00
55	DA	2835	A	N9-C1'-C2'	5.00	120.50	114.00
1	AA	74	A	C8-N9-C1'	5.00	136.70	127.70
55	DA	1143	A	C4'-C3'-C2'	5.00	107.60	102.60

There are no chirality outliers.

All (395) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
1	AA	1025	G	Sidechain
1	AA	1082	U	Sidechain
1	AA	1143	A	Sidechain
1	AA	1159	U	Sidechain
1	AA	1215	G	Sidechain
1	AA	1242	A	Sidechain
1	AA	1247	A	Sidechain
1	AA	1249	U	Sidechain
1	AA	1250	G	Sidechain
1	AA	1252	G	Sidechain
1	AA	1294	U	Sidechain
1	AA	1298	C	Sidechain
1	AA	1302	A	Sidechain
1	AA	1308	A	Sidechain
1	AA	1312	U	Sidechain
1	AA	1357	U	Sidechain
1	AA	1397	U	Sidechain
1	AA	1427	A	Sidechain

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Mol	Chain	Res	Type	Group
1	AA	1455	G	Sidechain
1	AA	1564	C	Sidechain
1	AA	1566	A	Sidechain
1	AA	1671	U	Sidechain
1	AA	1693	U	Sidechain
1	AA	1751	C	Sidechain
1	AA	1772	G	Sidechain
1	AA	1773	A	Sidechain
1	AA	1774	C	Sidechain
1	AA	1783	A	Sidechain
1	AA	1807	G	Sidechain
1	AA	1818	U	Sidechain
1	AA	1902	C	Sidechain
1	AA	1910	G	Sidechain
1	AA	1940	U	Sidechain
1	AA	1946	U	Sidechain
1	AA	196	A	Sidechain
1	AA	1966	A	Sidechain
1	AA	1991	U	Sidechain
1	AA	201	C	Sidechain
1	AA	2025	C	Sidechain
1	AA	2031	A	Sidechain
1	AA	2034	U	Sidechain
1	AA	2049	G	Sidechain
1	AA	2074	U	Sidechain
1	AA	2227	A	Sidechain
1	AA	2248	C	Sidechain
1	AA	2257	U	Sidechain
1	AA	227	A	Sidechain
1	AA	2307	G	Sidechain
1	AA	2312	U	Sidechain
1	AA	232	G	Sidechain
1	AA	2345	G	Sidechain
1	AA	2346	A	Sidechain
1	AA	2401	U	Sidechain
1	AA	2406	U	Sidechain
1	AA	2427	C	Sidechain
1	AA	2447	G	Sidechain
1	AA	2458	G	Sidechain
1	AA	2497	A	Sidechain
1	AA	250	G	Sidechain
1	AA	2506	U	Sidechain

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Mol	Chain	Res	Type	Group
1	AA	2542	A	Sidechain
1	AA	2582	G	Sidechain
1	AA	2596	U	Sidechain
1	AA	2605	U	Sidechain
1	AA	2609	U	Sidechain
1	AA	2625	G	Sidechain
1	AA	2647	U	Sidechain
1	AA	2656	U	Sidechain
1	AA	2681	C	Sidechain
1	AA	2696	U	Sidechain
1	AA	2725	A	Sidechain
1	AA	2746	U	Sidechain
1	AA	2789	C	Sidechain
1	AA	2835	A	Sidechain
1	AA	2899	G	Sidechain
1	AA	305	U	Sidechain
1	AA	319	C	Sidechain
1	AA	33	U	Sidechain
1	AA	331	A	Sidechain
1	AA	352	G	Sidechain
1	AA	371	A	Sidechain
1	AA	403	U	Sidechain
1	AA	441	U	Sidechain
1	AA	463	G	Sidechain
1	AA	476	G	Sidechain
1	AA	481	G	Sidechain
1	AA	50	U	Sidechain
1	AA	506	G	Sidechain
1	AA	510	C	Sidechain
1	AA	566	U	Sidechain
1	AA	607	U	Sidechain
1	AA	608	A	Sidechain
1	AA	642	G	Sidechain
1	AA	654(G)	C	Sidechain
1	AA	654(M)	C	Sidechain
1	AA	670	A	Sidechain
1	AA	682	G	Sidechain
1	AA	683	C	Sidechain
1	AA	70	G	Sidechain
1	AA	72	U	Sidechain
1	AA	74	A	Sidechain
1	AA	788	A	Sidechain

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Mol	Chain	Res	Type	Group
1	AA	816	C	Sidechain
1	AA	817	C	Sidechain
1	AA	859	G	Sidechain
1	AA	895	U	Sidechain
1	AA	913	U	Sidechain
1	AA	930	U	Sidechain
1	AA	943	U	Sidechain
2	AB	1	U	Sidechain
2	AB	66	A	Sidechain
2	AB	81	G	Sidechain
2	AB	95	U	Sidechain
3	AD	104	TYR	Sidechain
3	AD	9	TYR	Sidechain
53	B1	41	U	Sidechain
53	B1	48	U	Sidechain
53	B1	51	U	Sidechain
53	B1	53	U	Sidechain
31	BA	1049	U	Sidechain
31	BA	1064	G	Sidechain
31	BA	1065	U	Sidechain
31	BA	1077	G	Sidechain
31	BA	114	U	Sidechain
31	BA	1149	C	Sidechain
31	BA	1159	U	Sidechain
31	BA	1201	A	Sidechain
31	BA	1205	U	Sidechain
31	BA	1235	U	Sidechain
31	BA	130	A	Sidechain
31	BA	1341	U	Sidechain
31	BA	1380	U	Sidechain
31	BA	1526	G	Sidechain
31	BA	1529	G	Sidechain
31	BA	1540	U	Sidechain
31	BA	1541	U	Sidechain
31	BA	190	G	Sidechain
31	BA	249	U	Sidechain
31	BA	250	A	Sidechain
31	BA	251	G	Sidechain
31	BA	328	C	Sidechain
31	BA	366	C	Sidechain
31	BA	528	C	Sidechain
31	BA	566	G	Sidechain

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Mol	Chain	Res	Type	Group
31	BA	571	U	Sidechain
31	BA	575	G	Sidechain
31	BA	672	U	Sidechain
31	BA	686	U	Sidechain
31	BA	703	G	Sidechain
31	BA	788	U	Sidechain
31	BA	82	U	Sidechain
31	BA	873	A	Sidechain
31	BA	974	A	Sidechain
52	BB	66	U	Sidechain
52	BC	19	G	Sidechain
52	BC	66	U	Sidechain
52	BC	73	A	Sidechain
52	BD	7	A	Sidechain
53	C1	38	U	Sidechain
53	C1	41	U	Sidechain
53	C1	56	U	Sidechain
54	CA	1054	C	Sidechain
54	CA	1064	G	Sidechain
54	CA	1122	U	Sidechain
54	CA	1124	G	Sidechain
54	CA	1128	C	Sidechain
54	CA	1129	C	Sidechain
54	CA	114	U	Sidechain
54	CA	1143	G	Sidechain
54	CA	1150	U	Sidechain
54	CA	1159	U	Sidechain
54	CA	1190	G	Sidechain
54	CA	1201	A	Sidechain
54	CA	1205	U	Sidechain
54	CA	121	C	Sidechain
54	CA	1225	A	Sidechain
54	CA	1226	C	Sidechain
54	CA	1285	A	Sidechain
54	CA	1380	U	Sidechain
54	CA	1400	C	Sidechain
54	CA	1495	U	Sidechain
54	CA	1502	A	Sidechain
54	CA	1535	C	Sidechain
54	CA	1541	U	Sidechain
54	CA	259	G	Sidechain
54	CA	30	U	Sidechain

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Mol	Chain	Res	Type	Group
54	CA	367	U	Sidechain
54	CA	387	U	Sidechain
54	CA	388	G	Sidechain
54	CA	484	G	Sidechain
54	CA	50	A	Sidechain
54	CA	51	A	Sidechain
54	CA	512	U	Sidechain
54	CA	518	C	Sidechain
54	CA	528	C	Sidechain
54	CA	532	A	Sidechain
54	CA	575	G	Sidechain
54	CA	58	C	Sidechain
54	CA	620	C	Sidechain
54	CA	672	U	Sidechain
54	CA	681	C	Sidechain
54	CA	686	U	Sidechain
54	CA	693	G	Sidechain
54	CA	697	U	Sidechain
54	CA	73	G	Sidechain
54	CA	749	C	Sidechain
54	CA	760	G	Sidechain
54	CA	773	G	Sidechain
54	CA	794	A	Sidechain
54	CA	84	U	Sidechain
54	CA	85	U	Sidechain
54	CA	873	A	Sidechain
54	CA	879	C	Sidechain
54	CA	900	A	Sidechain
54	CA	960	U	Sidechain
54	CA	965	A	Sidechain
54	CA	97	U	Sidechain
54	CA	974	A	Sidechain
52	CB	49	C	Sidechain
52	CD	45	U	Sidechain
52	CD	66	U	Sidechain
55	DA	1025	G	Sidechain
55	DA	104	U	Sidechain
55	DA	1060	U	Sidechain
55	DA	1078	U	Sidechain
55	DA	1082	U	Sidechain
55	DA	1086	A	Sidechain
55	DA	1092	C	Sidechain

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Mol	Chain	Res	Type	Group
55	DA	1099	G	Sidechain
55	DA	1126	A	Sidechain
55	DA	113	G	Sidechain
55	DA	1130	U	Sidechain
55	DA	1132	A	Sidechain
55	DA	1133	U	Sidechain
55	DA	1156	A	Sidechain
55	DA	1161	C	Sidechain
55	DA	1188	U	Sidechain
55	DA	119	A	Sidechain
55	DA	12	U	Sidechain
55	DA	1249	U	Sidechain
55	DA	1252	G	Sidechain
55	DA	1253	A	Sidechain
55	DA	1287	A	Sidechain
55	DA	1288	U	Sidechain
55	DA	1294	U	Sidechain
55	DA	1300	U	Sidechain
55	DA	1302	A	Sidechain
55	DA	1323	U	Sidechain
55	DA	1340	U	Sidechain
55	DA	1357	U	Sidechain
55	DA	138	G	Sidechain
55	DA	139	G	Sidechain
55	DA	1425	G	Sidechain
55	DA	1503	U	Sidechain
55	DA	1534	G	Sidechain
55	DA	1535	U	Sidechain
55	DA	1537	C	Sidechain
55	DA	1558	A	Sidechain
55	DA	1564	C	Sidechain
55	DA	1607	C	Sidechain
55	DA	1610	A	Sidechain
55	DA	1619	G	Sidechain
55	DA	1647	G	Sidechain
55	DA	1651	G	Sidechain
55	DA	1664	A	Sidechain
55	DA	1675	C	Sidechain
55	DA	1693	U	Sidechain
55	DA	1698	A	Sidechain
55	DA	1772	G	Sidechain
55	DA	1773	A	Sidechain

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Mol	Chain	Res	Type	Group
55	DA	1774	C	Sidechain
55	DA	1777	U	Sidechain
55	DA	1779	U	Sidechain
55	DA	1799	G	Sidechain
55	DA	1801	G	Sidechain
55	DA	1808	U	Sidechain
55	DA	1828	G	Sidechain
55	DA	1833	U	Sidechain
55	DA	1834	U	Sidechain
55	DA	1925	C	Sidechain
55	DA	1926	U	Sidechain
55	DA	1929	G	Sidechain
55	DA	1934	C	Sidechain
55	DA	1940	U	Sidechain
55	DA	1946	U	Sidechain
55	DA	1951	U	Sidechain
55	DA	1964	G	Sidechain
55	DA	1970	A	Sidechain
55	DA	1991	U	Sidechain
55	DA	2009	G	Sidechain
55	DA	201	C	Sidechain
55	DA	2025	C	Sidechain
55	DA	2028	U	Sidechain
55	DA	2031	A	Sidechain
55	DA	2034	U	Sidechain
55	DA	2078	C	Sidechain
55	DA	2086	U	Sidechain
55	DA	2098	U	Sidechain
55	DA	2110	G	Sidechain
55	DA	2252	G	Sidechain
55	DA	2257	U	Sidechain
55	DA	227	A	Sidechain
55	DA	2275	C	Sidechain
55	DA	2282	G	Sidechain
55	DA	229	A	Sidechain
55	DA	2313	C	Sidechain
55	DA	2319	G	Sidechain
55	DA	2334	G	Sidechain
55	DA	2345	G	Sidechain
55	DA	240	G	Sidechain
55	DA	2401	U	Sidechain
55	DA	2426	A	Sidechain

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Mol	Chain	Res	Type	Group
55	DA	2427	C	Sidechain
55	DA	243	U	Sidechain
55	DA	2434	A	Sidechain
55	DA	2447	G	Sidechain
55	DA	2452	C	Sidechain
55	DA	2462	U	Sidechain
55	DA	2489	G	Sidechain
55	DA	249	C	Sidechain
55	DA	250	G	Sidechain
55	DA	2504	U	Sidechain
55	DA	2516	G	Sidechain
55	DA	2542	A	Sidechain
55	DA	2563	U	Sidechain
55	DA	2582	G	Sidechain
55	DA	2587	A	Sidechain
55	DA	2593	U	Sidechain
55	DA	2595	G	Sidechain
55	DA	2615	U	Sidechain
55	DA	2627	G	Sidechain
55	DA	265	A	Sidechain
55	DA	2650	U	Sidechain
55	DA	2656	U	Sidechain
55	DA	2665	A	Sidechain
55	DA	2684	U	Sidechain
55	DA	2696	U	Sidechain
55	DA	270(Z)	U	Sidechain
55	DA	2712	U	Sidechain
55	DA	2725	A	Sidechain
55	DA	2751	G	Sidechain
55	DA	2779	U	Sidechain
55	DA	2789	C	Sidechain
55	DA	2873	A	Sidechain
55	DA	323	G	Sidechain
55	DA	339	U	Sidechain
55	DA	370	G	Sidechain
55	DA	371	A	Sidechain
55	DA	385	C	Sidechain
55	DA	387	U	Sidechain
55	DA	4	C	Sidechain
55	DA	411	G	Sidechain
55	DA	43	G	Sidechain
55	DA	441	U	Sidechain

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Mol	Chain	Res	Type	Group
55	DA	463	G	Sidechain
55	DA	467	G	Sidechain
55	DA	476	G	Sidechain
55	DA	487	C	Sidechain
55	DA	488	G	Sidechain
55	DA	508	G	Sidechain
55	DA	52	A	Sidechain
55	DA	56	A	Sidechain
55	DA	566	U	Sidechain
55	DA	567	A	Sidechain
55	DA	575	A	Sidechain
55	DA	590	A	Sidechain
55	DA	602	G	Sidechain
55	DA	606	U	Sidechain
55	DA	607	U	Sidechain
55	DA	621	A	Sidechain
55	DA	630	G	Sidechain
55	DA	654(G)	C	Sidechain
55	DA	654(M)	C	Sidechain
55	DA	670	A	Sidechain
55	DA	675	A	Sidechain
55	DA	682	G	Sidechain
55	DA	683	C	Sidechain
55	DA	70	G	Sidechain
55	DA	72	U	Sidechain
55	DA	730	C	Sidechain
55	DA	739	G	Sidechain
55	DA	792	G	Sidechain
55	DA	807	U	Sidechain
55	DA	829	A	Sidechain
55	DA	859	G	Sidechain
55	DA	895	U	Sidechain
55	DA	943	U	Sidechain
55	DA	956	G	Sidechain
55	DA	980	A	Sidechain
55	DA	990	A	Sidechain
55	DA	995	C	Sidechain
2	DB	51	G	Sidechain
2	DB	55	U	Sidechain
2	DB	81	G	Sidechain
2	DB	95	U	Sidechain

5.2 Too-close contacts ⓘ

In the following table, the Non-H and H(model) columns list the number of non-hydrogen atoms and hydrogen atoms in the chain respectively. The H(added) column lists the number of hydrogen atoms added and optimized by MolProbity. The Clashes column lists the number of clashes within the asymmetric unit, whereas Symm-Clashes lists symmetry related clashes.

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	AA	62647	0	31576	3579	1
2	AB	2617	0	1328	187	0
2	DB	2617	0	1328	116	0
3	AD	2115	0	2195	312	0
3	DD	2115	0	2195	370	0
4	AE	1568	0	1634	355	0
4	DE	1568	0	1634	274	0
5	AF	1627	0	1679	258	0
5	DF	1585	0	1631	192	0
6	AG	1474	0	1535	252	0
6	DG	1474	0	1535	277	0
7	AH	1307	0	1382	233	0
7	DH	1307	0	1382	255	0
8	AK	1136	0	1223	199	0
8	DK	1136	0	1221	222	0
9	AM	1104	0	1180	164	0
9	DM	1104	0	1180	178	0
10	AN	933	0	996	107	0
10	DN	933	0	996	114	0
11	AO	1145	0	1228	255	0
11	DO	1145	0	1227	265	0
12	AP	1122	0	1179	242	0
12	DP	1122	0	1179	180	0
13	A0	960	0	1021	133	0
13	D0	968	0	1033	146	0
14	AQ	882	0	943	142	0
14	DQ	882	0	943	144	0
15	AR	1141	0	1202	171	0
15	DR	1141	0	1202	183	0
16	A1	964	0	1022	169	1
16	D1	964	0	1022	144	0
17	A2	779	0	852	210	0
17	D2	779	0	852	148	0
18	AS	900	0	964	97	0
18	DS	900	0	963	89	0
19	AT	725	0	778	92	0
19	DT	725	0	778	78	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
20	AU	785	0	878	217	0
20	DU	785	0	878	171	0
21	AV	1489	0	1515	418	0
21	DV	1582	0	1613	447	0
22	A3	662	0	688	103	0
22	D3	662	0	688	87	0
23	AZ	763	0	848	107	0
23	DZ	763	0	848	92	0
24	AW	581	0	629	125	0
24	DW	581	0	629	76	0
25	AX	469	0	518	62	0
25	DX	469	0	518	46	0
26	A4	581	0	573	181	0
26	D4	581	0	573	174	0
27	A5	459	0	480	62	0
27	D5	459	0	480	81	0
28	A6	389	0	404	119	0
28	D6	389	0	404	126	0
29	A7	430	0	480	49	0
29	D7	430	0	480	35	0
30	A8	517	0	582	153	0
30	D8	517	0	582	103	0
31	BA	32571	0	16441	1933	1
32	BE	1924	0	1975	295	0
32	CE	1924	0	1975	286	0
33	BF	1612	0	1677	257	0
33	CF	1605	0	1668	217	0
34	BG	1703	0	1763	257	0
34	CG	1703	0	1763	235	0
35	BH	1155	0	1212	133	0
35	CH	1155	0	1213	151	0
36	BI	843	0	857	96	0
36	CI	843	0	857	101	0
37	BJ	1257	0	1296	139	0
37	CJ	1257	0	1296	138	0
38	BK	1116	0	1177	121	0
38	CK	1116	0	1177	164	0
39	BL	1010	0	1037	187	0
39	CL	1010	0	1037	152	0
40	BM	801	0	849	165	0
40	CM	801	0	849	149	0
41	BN	885	0	904	73	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
41	CN	885	0	904	76	0
42	BO	975	0	1062	140	0
42	CO	975	0	1062	138	0
43	BP	964	0	1034	181	0
43	CP	997	0	1072	192	0
44	BQ	492	0	529	84	0
44	CQ	492	0	529	82	0
45	BR	734	0	771	72	0
45	CR	734	0	771	75	0
46	BS	705	0	725	80	0
46	CS	705	0	725	131	0
47	BT	834	0	904	73	0
47	CT	834	0	904	92	0
48	BU	591	0	662	94	0
48	CU	591	0	662	100	0
49	BV	665	0	684	174	0
49	CV	702	0	728	156	0
50	BW	763	0	861	119	0
50	CW	763	0	861	113	0
51	BX	217	0	234	37	0
51	CX	217	0	234	23	0
52	BB	1626	0	832	95	0
52	BC	1626	0	833	86	0
52	BD	1626	0	833	89	0
52	CB	1626	0	832	97	0
52	CC	1626	0	832	62	0
52	CD	1626	0	832	118	0
53	B1	621	0	312	68	0
53	C1	621	0	312	76	0
54	CA	32551	0	16431	1895	0
55	DA	62707	0	31590	3338	1
56	DI	237	0	257	165	0
56	DJ	237	0	256	208	0
57	DY	1107	0	1166	947	0
58	DL	1071	0	1113	696	0
59	A0	1	0	0	0	0
59	A1	5	0	0	0	0
59	A2	1	0	0	0	0
59	A3	4	0	0	0	0
59	A5	3	0	0	0	0
59	A6	2	0	0	0	0
59	A7	4	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
59	A8	4	0	0	0	0
59	AA	1166	0	0	0	0
59	AB	36	0	0	0	0
59	AD	13	0	0	0	0
59	AE	4	0	0	0	0
59	AF	7	0	0	0	0
59	AG	3	0	0	0	0
59	AH	1	0	0	0	0
59	AK	3	0	0	0	0
59	AN	1	0	0	0	0
59	AO	7	0	0	0	0
59	AP	1	0	0	0	0
59	AQ	5	0	0	0	0
59	AR	5	0	0	0	0
59	AS	3	0	0	0	0
59	AT	4	0	0	0	0
59	AU	6	0	0	0	0
59	AV	1	0	0	0	0
59	AW	2	0	0	0	0
59	AZ	3	0	0	0	0
59	B1	4	0	0	0	0
59	BA	676	0	0	0	0
59	BB	13	0	0	0	0
59	BC	16	0	0	0	0
59	BD	26	0	0	0	0
59	BE	5	0	0	0	0
59	BF	2	0	0	0	0
59	BG	7	0	0	0	0
59	BH	5	0	0	0	0
59	BI	1	0	0	0	0
59	BK	6	0	0	0	0
59	BL	2	0	0	0	0
59	BM	3	0	0	0	0
59	BO	1	0	0	0	0
59	BP	2	0	0	0	0
59	BQ	3	0	0	0	0
59	BS	9	0	0	0	0
59	BT	2	0	0	0	0
59	BU	1	0	0	0	0
59	BV	1	0	0	0	0
59	BW	8	0	0	0	0
59	BX	1	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
59	C1	6	0	0	0	0
59	CA	724	0	0	1	0
59	CB	21	0	0	0	0
59	CC	27	0	0	0	0
59	CD	30	0	0	0	0
59	CE	6	0	0	0	0
59	CF	3	0	0	0	0
59	CG	11	0	0	0	0
59	CH	6	0	0	0	0
59	CI	1	0	0	0	0
59	CJ	1	0	0	0	0
59	CK	11	0	0	0	0
59	CL	1	0	0	0	0
59	CM	3	0	0	0	0
59	CN	2	0	0	0	0
59	CP	3	0	0	0	0
59	CQ	3	0	0	0	0
59	CR	3	0	0	0	0
59	CS	6	0	0	0	0
59	CT	3	0	0	0	0
59	CV	4	0	0	0	0
59	CW	1	0	0	0	0
59	D0	11	0	0	0	0
59	D1	12	0	0	0	0
59	D2	10	0	0	0	0
59	D3	7	0	0	0	0
59	D4	2	0	0	0	0
59	D5	8	0	0	0	0
59	D6	3	0	0	0	0
59	D7	5	0	0	0	0
59	D8	9	0	0	0	0
59	DA	2077	0	0	0	0
59	DB	76	0	0	0	0
59	DD	14	0	0	0	0
59	DE	15	0	0	0	0
59	DF	25	0	0	0	0
59	DG	5	0	0	0	0
59	DH	5	0	0	0	0
59	DK	2	0	0	0	0
59	DL	1	0	0	0	0
59	DM	6	0	0	0	0
59	DN	2	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
59	DO	18	0	0	0	0
59	DP	4	0	0	0	0
59	DQ	2	0	0	0	0
59	DR	4	0	0	1	0
59	DS	8	0	0	0	0
59	DT	7	0	0	0	0
59	DU	19	0	0	0	0
59	DV	5	0	0	0	0
59	DW	7	0	0	0	0
59	DX	2	0	0	0	0
59	DY	4	0	0	0	0
59	DZ	5	0	0	0	0
60	A4	1	0	0	0	0
60	BG	1	0	0	0	0
60	BQ	1	0	0	0	0
60	CG	1	0	0	0	0
60	CQ	1	0	0	0	0
60	D4	1	0	0	0	0
61	B1	3	0	0	0	0
61	BA	3	0	0	0	0
All	All	307345	0	204878	26272	2

The all-atom clashscore is defined as the number of clashes found per 1000 atoms (including hydrogen atoms). The all-atom clashscore for this structure is 52.

All (26272) close contacts within the same asymmetric unit are listed below, sorted by their clash magnitude.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2820:A:C8	4:AE:109:LYS:HE3	1.23	1.64
58:DL:7:VAL:HG12	58:DL:57:ILE:CD1	1.29	1.63
57:DY:71:LEU:CB	57:DY:113:GLN:HB3	1.32	1.60
21:AV:175:VAL:HG12	21:AV:177:PRO:CD	1.26	1.56
58:DL:7:VAL:CG1	58:DL:57:ILE:HD12	1.21	1.55
56:DJ:13:SER:CB	56:DJ:17:VAL:HG21	1.30	1.54
57:DY:51:LEU:HD21	57:DY:82:PHE:C	1.22	1.53
57:DY:29:TYR:N	57:DY:81:VAL:CG1	1.69	1.52
57:DY:27:VAL:CB	57:DY:110:GLY:HA3	1.11	1.52
57:DY:27:VAL:CG2	57:DY:110:GLY:CA	1.83	1.52
57:DY:89:ALA:HB3	56:DJ:15:ALA:CB	1.41	1.50
57:DY:27:VAL:CG2	57:DY:110:GLY:HA3	1.01	1.49
56:DJ:12:LEU:CA	56:DJ:13:SER:HB2	1.36	1.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:AV:175:VAL:C	21:AV:177:PRO:HD2	1.20	1.47
58:DL:20:ALA:N	58:DL:25:PRO:HD2	1.31	1.46
56:DJ:12:LEU:N	56:DJ:13:SER:HB2	1.29	1.45
21:DV:191:VAL:HG11	21:DV:197:ILE:CG1	1.42	1.44
1:AA:2820:A:N6	4:AE:192:ASN:H	1.06	1.44
57:DY:25:PHE:CD1	57:DY:82:PHE:CD2	2.03	1.44
57:DY:51:LEU:CD2	57:DY:82:PHE:O	1.63	1.44
57:DY:71:LEU:HB3	57:DY:113:GLN:CB	1.44	1.43
58:DL:20:ALA:H	58:DL:25:PRO:CG	1.31	1.43
43:CP:124:PRO:HB3	43:CP:125:ARG:CG	1.48	1.42
58:DL:18:THR:HB	58:DL:19:PRO:CD	1.48	1.42
57:DY:51:LEU:HD13	57:DY:82:PHE:N	1.15	1.42
57:DY:27:VAL:HB	57:DY:110:GLY:CA	1.45	1.41
21:AV:175:VAL:HG13	21:AV:176:PRO:CD	1.48	1.41
21:DV:189:ALA:CB	21:DV:190:GLU:HG2	1.47	1.41
21:DV:191:VAL:CG1	21:DV:197:ILE:HG21	1.51	1.41
49:BV:41:VAL:CG1	49:BV:42:PRO:HD2	1.48	1.40
56:DJ:13:SER:CA	56:DJ:17:VAL:HG21	1.52	1.40
57:DY:43:ALA:HB3	57:DY:47:ASN:ND2	1.36	1.39
58:DL:104:VAL:O	58:DL:107:ILE:CG2	1.68	1.39
58:DL:8:VAL:O	58:DL:57:ILE:CG1	1.69	1.39
57:DY:27:VAL:HB	57:DY:110:GLY:C	1.36	1.38
57:DY:21:GLN:HE21	57:DY:22:GLY:N	1.19	1.38
58:DL:7:VAL:HG11	58:DL:58:THR:N	1.33	1.38
57:DY:43:ALA:CB	57:DY:47:ASN:HD22	1.37	1.37
56:DJ:13:SER:CB	56:DJ:17:VAL:CG2	2.03	1.36
21:DV:191:VAL:CG1	21:DV:197:ILE:CG2	2.03	1.36
21:AV:115:GLY:CA	21:AV:177:PRO:HG2	1.55	1.35
49:BV:41:VAL:HG12	49:BV:42:PRO:CD	1.53	1.34
55:DA:1060:U:OP1	58:DL:54:PRO:HG3	1.28	1.34
55:DA:1359:A:H3'	55:DA:1359:A:C8	1.62	1.33
21:AV:175:VAL:CG1	21:AV:177:PRO:HD3	1.56	1.33
58:DL:7:VAL:HG13	58:DL:58:THR:O	1.19	1.33
56:DJ:13:SER:HB3	56:DJ:17:VAL:CG1	1.59	1.33
57:DY:19:ARG:CZ	57:DY:84:GLU:OE1	1.76	1.33
57:DY:25:PHE:CZ	57:DY:82:PHE:HB3	1.62	1.33
21:DV:189:ALA:CA	21:DV:190:GLU:HG2	1.56	1.32
21:AV:115:GLY:HA2	21:AV:177:PRO:CG	1.58	1.32
28:A6:41:PRO:CG	28:A6:45:LYS:O	1.77	1.32
58:DL:20:ALA:N	58:DL:25:PRO:CD	1.91	1.32
57:DY:27:VAL:CB	57:DY:110:GLY:CA	1.91	1.31

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:27:VAL:HB	57:DY:111:LEU:N	1.44	1.31
1:AA:1899:G:N2	1:AA:1902:C:H41	1.29	1.31
57:DY:112:LEU:N	57:DY:112:LEU:HD23	1.38	1.30
26:D4:68:ARG:CZ	26:D4:68:ARG:HA	1.61	1.30
1:AA:384:U:C2'	1:AA:385:C:H5'	1.62	1.30
43:BP:80:ARG:O	43:BP:83:ASP:HB3	1.23	1.30
56:DJ:12:LEU:N	56:DJ:13:SER:CB	1.95	1.30
57:DY:29:TYR:CE2	57:DY:32:LEU:HD11	1.64	1.30
58:DL:52:ILE:CG2	58:DL:75:SER:HB2	1.61	1.30
54:CA:1313:U:P	49:CV:6:LYS:HB2	1.71	1.29
1:AA:1359:A:H3'	1:AA:1359:A:C8	1.54	1.29
58:DL:14:ALA:CB	58:DL:50:ASP:HB2	1.60	1.29
1:AA:946:G:O2'	1:AA:947:G:H5'	1.29	1.28
57:DY:89:ALA:CB	56:DJ:15:ALA:HB1	1.62	1.28
57:DY:26:LEU:HD22	57:DY:121:ASP:OD2	1.12	1.27
55:DA:1899:G:H22	55:DA:1902:C:N4	1.31	1.27
1:AA:946:G:O2'	1:AA:947:G:C5'	1.83	1.26
57:DY:16:ASN:HB2	57:DY:19:ARG:NH1	1.50	1.26
54:CA:1313:U:OP2	49:CV:6:LYS:HB2	1.22	1.26
55:DA:1372:U:H5'	55:DA:1372:U:C6	1.70	1.26
57:DY:130:THR:O	57:DY:134:LEU:HD13	1.27	1.26
28:A6:41:PRO:HG2	28:A6:45:LYS:O	1.26	1.25
57:DY:16:ASN:CB	57:DY:19:ARG:HH12	1.48	1.25
57:DY:130:THR:CG2	56:DJ:14:GLN:HE22	1.46	1.25
55:DA:1057:A:N7	55:DA:1086:A:H2'	1.50	1.25
1:AA:2820:A:C8	4:AE:109:LYS:CE	2.20	1.25
42:CO:47:LYS:O	42:CO:49:ASN:N	1.68	1.25
21:AV:175:VAL:CG1	21:AV:177:PRO:CD	2.12	1.24
21:DV:191:VAL:HG11	21:DV:197:ILE:CB	1.66	1.24
58:DL:52:ILE:CD1	58:DL:76:TYR:HB3	1.65	1.24
21:AV:106:GLY:C	21:AV:108:PRO:HD2	1.55	1.24
57:DY:51:LEU:CD1	57:DY:82:PHE:N	1.99	1.24
55:DA:1075:C:H4'	21:DV:195:GLU:CG	1.68	1.24
1:AA:1899:G:H22	1:AA:1902:C:N4	1.35	1.24
55:DA:1899:G:N2	55:DA:1902:C:H41	1.36	1.24
1:AA:2820:A:N6	4:AE:192:ASN:N	1.84	1.23
21:AV:175:VAL:CG1	21:AV:176:PRO:HD2	1.68	1.23
58:DL:52:ILE:HG21	58:DL:75:SER:CB	1.67	1.23
57:DY:21:GLN:NE2	57:DY:22:GLY:H	1.33	1.23
21:DV:116:VAL:HB	21:DV:175:VAL:O	1.14	1.23
55:DA:1301:A:O2'	55:DA:1302:A:H3'	1.36	1.22

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:AD:43:ARG:HH11	3:AD:44:ASN:ND2	1.36	1.22
58:DL:135:GLY:O	58:DL:136:VAL:HG13	1.40	1.22
58:DL:19:PRO:HA	58:DL:25:PRO:CG	1.70	1.21
21:AV:108:PRO:CB	21:AV:141:VAL:O	1.87	1.21
21:DV:190:GLU:O	21:DV:191:VAL:HG23	1.41	1.21
56:DI:9:LYS:O	56:DI:11:GLU:N	1.73	1.21
56:DJ:12:LEU:CB	56:DJ:13:SER:HB2	1.68	1.21
57:DY:23:SER:CB	57:DY:68:LEU:HB2	1.69	1.21
57:DY:75:GLN:HB3	57:DY:110:GLY:O	1.41	1.20
57:DY:71:LEU:HB2	57:DY:112:LEU:O	1.33	1.20
21:DV:189:ALA:HB1	21:DV:190:GLU:CG	1.72	1.20
58:DL:14:ALA:HA	58:DL:49:GLY:HA3	1.21	1.20
57:DY:29:TYR:CA	57:DY:81:VAL:HG12	1.67	1.20
57:DY:25:PHE:CD1	57:DY:82:PHE:CG	2.29	1.20
54:CA:630:G:C2'	54:CA:631:G:H5''	1.70	1.20
54:CA:73:G:N2	54:CA:74:C:H41	1.38	1.20
57:DY:40:LEU:CD2	57:DY:50:ARG:HH12	1.55	1.19
57:DY:132:ASP:O	57:DY:134:LEU:N	1.74	1.19
57:DY:23:SER:OG	57:DY:114:GLY:HA2	1.42	1.19
32:CE:8:LYS:H	32:CE:8:LYS:HD3	1.04	1.19
31:BA:887:G:C2'	31:BA:888:G:H5'	1.72	1.19
57:DY:25:PHE:CE1	57:DY:82:PHE:CG	2.31	1.19
1:AA:2820:A:H8	4:AE:109:LYS:CE	1.55	1.19
3:DD:25:THR:HG21	3:DD:81:ALA:HB1	1.19	1.19
58:DL:87:GLY:HA2	58:DL:96:VAL:HG21	1.19	1.19
3:AD:44:ASN:HB3	3:AD:49:ILE:HA	1.21	1.18
57:DY:50:ARG:O	57:DY:83:TYR:CA	1.91	1.18
56:DJ:14:GLN:HG2	56:DJ:16:THR:O	1.42	1.18
22:A3:32:ARG:H	22:A3:35:ASN:ND2	1.40	1.18
57:DY:73:GLY:O	57:DY:119:ALA:HA	1.43	1.18
57:DY:27:VAL:HG21	57:DY:110:GLY:CA	1.59	1.18
43:BP:22:ILE:HB	43:BP:25:ILE:HG12	1.21	1.18
55:DA:1058:U:H2'	55:DA:1059:G:C8	1.78	1.18
57:DY:73:GLY:O	57:DY:119:ALA:CA	1.90	1.18
1:AA:1378:A:O2'	1:AA:1379:A:H5''	1.42	1.17
58:DL:7:VAL:CG1	58:DL:57:ILE:CD1	1.98	1.17
23:DZ:91:LYS:HA	23:DZ:91:LYS:HE3	1.24	1.17
54:CA:792:A:H2'	54:CA:794:A:N6	1.57	1.17
58:DL:14:ALA:HB2	58:DL:50:ASP:HB2	1.21	1.17
21:AV:175:VAL:C	21:AV:177:PRO:CD	2.13	1.16
58:DL:141:ALA:HB1	58:DL:143:GLU:N	1.58	1.16

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:30:GLN:OE1	57:DY:79:ALA:O	1.62	1.16
1:AA:1359:A:C8	1:AA:1359:A:C3'	2.29	1.16
15:AR:16:ARG:HH21	15:AR:19:LEU:HD21	1.05	1.16
1:AA:1887:C:H2'	1:AA:1888:G:H5''	1.28	1.16
56:DI:24:ILE:HG12	56:DI:25:ASP:H	1.04	1.16
21:AV:144:LEU:O	21:AV:174:VAL:HG21	1.44	1.16
58:DL:18:THR:CB	58:DL:19:PRO:HD3	1.76	1.16
21:DV:189:ALA:CB	21:DV:190:GLU:CG	2.23	1.16
56:DI:3:LEU:HD23	56:DI:7:ARG:HD2	1.23	1.16
54:CA:792:A:C2'	54:CA:794:A:H62	1.57	1.16
49:CV:41:VAL:HB	49:CV:42:PRO:HA	1.20	1.16
13:D0:33:ARG:HH22	27:D5:55:ARG:HG2	1.05	1.16
58:DL:103:GLN:O	58:DL:107:ILE:HB	1.44	1.15
57:DY:80:VAL:HG12	57:DY:81:VAL:H	0.99	1.15
30:A8:32:LEU:HD23	30:A8:34:TRP:H	1.08	1.15
40:BM:4:ILE:HB	40:BM:74:ILE:HD11	1.22	1.15
57:DY:16:ASN:CA	57:DY:19:ARG:NH1	2.09	1.15
57:DY:50:ARG:O	57:DY:83:TYR:HA	0.98	1.15
56:DI:29:GLU:HA	56:DJ:2:ALA:HB1	1.16	1.15
57:DY:25:PHE:HB3	57:DY:82:PHE:CZ	1.81	1.15
30:D8:52:LYS:H	30:D8:53:PRO:CD	1.57	1.15
55:DA:1075:C:C4'	21:DV:195:GLU:HG2	1.76	1.15
21:AV:110:GLY:H	21:AV:143:GLY:HA2	1.06	1.15
1:AA:242:G:H5''	30:A8:62:LEU:HD13	1.18	1.14
55:DA:2519:U:H4'	55:DA:2520:C:OP1	1.43	1.14
57:DY:72:ASP:O	57:DY:74:LEU:N	1.80	1.14
43:CP:124:PRO:CB	43:CP:125:ARG:HG2	1.77	1.14
58:DL:8:VAL:O	58:DL:57:ILE:HG13	1.24	1.14
1:AA:1372:U:C5'	1:AA:1372:U:C6	2.30	1.14
54:CA:1002:G:H2'	54:CA:1003:G:H8	1.13	1.14
55:DA:1372:U:C6	55:DA:1372:U:C5'	2.30	1.14
57:DY:144:ALA:HB1	57:DY:145:PRO:HD2	1.15	1.14
54:CA:1101:A:H4'	54:CA:1102:A:O5'	1.39	1.14
55:DA:483:A:H4'	20:DU:49:VAL:HA	1.22	1.14
56:DJ:12:LEU:HB3	56:DJ:13:SER:CB	1.77	1.14
55:DA:1359:A:C3'	55:DA:1359:A:C8	2.29	1.13
21:DV:194:PRO:HG2	21:DV:196:VAL:CG1	1.77	1.13
54:CA:1226:C:H4'	54:CA:1227:A:OP1	1.48	1.13
28:A6:41:PRO:CD	28:A6:45:LYS:O	1.95	1.13
43:CP:126:LYS:OXT	52:CC:27:G:N7	1.82	1.13
55:DA:2636:U:OP1	4:DE:79:ARG:HA	1.45	1.13

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:AK:142:VAL:HG22	8:AK:143:SER:H	1.13	1.13
46:CS:53:VAL:HG12	46:CS:79:VAL:HG22	1.26	1.13
8:DK:131:LYS:HB3	8:DK:132:PRO:HA	1.25	1.13
14:DQ:106:ARG:NH1	14:DQ:106:ARG:HB2	1.62	1.13
55:DA:1083:U:H5'	57:DY:47:ASN:OD1	1.44	1.13
31:BA:64:G:H4'	31:BA:65:U:H5''	1.25	1.12
43:BP:10:PRO:HB2	43:BP:18:ALA:HB1	1.26	1.12
55:DA:1484:G:H2'	55:DA:1485:G:H5''	1.19	1.12
55:DA:1378:A:O2'	55:DA:1379:A:H5''	1.47	1.12
58:DL:57:ILE:CD1	58:DL:58:THR:H	1.61	1.12
54:CA:1007:C:H2'	54:CA:1008:C:H5''	1.29	1.12
55:DA:1178:C:H2'	55:DA:1179:C:C6	1.82	1.12
56:DI:28:LYS:HA	56:DI:28:LYS:HE3	1.18	1.12
58:DL:12:LEU:HB3	58:DL:13:PRO:HA	1.24	1.12
57:DY:118:THR:HG23	57:DY:119:ALA:H	1.11	1.12
58:DL:20:ALA:N	58:DL:25:PRO:CG	2.06	1.12
1:AA:387:U:O2'	1:AA:388:G:H5''	1.50	1.12
15:DR:90:GLN:HE21	15:DR:90:GLN:HA	1.09	1.12
21:DV:150:LEU:CD2	21:DV:151:HIS:H	1.62	1.12
57:DY:90:ALA:O	57:DY:94:VAL:HB	1.50	1.12
54:CA:547:A:H4'	54:CA:548:G:O5'	1.42	1.12
57:DY:50:ARG:HA	57:DY:83:TYR:CD1	1.84	1.12
57:DY:40:LEU:HD23	57:DY:50:ARG:HH12	1.05	1.12
57:DY:28:ASN:HB3	57:DY:81:VAL:HG13	1.31	1.11
28:D6:15:GLU:HG2	28:D6:16:CYS:H	1.15	1.11
55:DA:1371:G:O2'	55:DA:1372:U:C5	1.99	1.11
57:DY:28:ASN:OD1	57:DY:83:TYR:HE2	1.28	1.11
58:DL:20:ALA:H	58:DL:25:PRO:HG2	1.15	1.11
54:CA:1234:C:H4'	54:CA:1364:U:O2'	1.48	1.11
3:DD:27:THR:HG23	3:DD:28:GLU:H	1.13	1.11
57:DY:104:ILE:HG13	57:DY:105:PRO:HD2	1.15	1.11
57:DY:36:GLU:O	57:DY:38:HIS:ND1	1.82	1.11
9:AM:39:ARG:HH21	9:AM:41:ASP:HB2	1.13	1.11
58:DL:132:ARG:HG2	58:DL:137:GLU:OE2	1.49	1.11
27:A5:4:HIS:HB3	27:A5:5:PRO:HD3	1.24	1.11
30:A8:49:VAL:HG12	30:A8:50:LEU:N	1.55	1.11
8:AK:79:ILE:HB	8:AK:142:VAL:HG11	1.17	1.11
57:DY:73:GLY:HA3	57:DY:112:LEU:CD1	1.80	1.11
1:AA:1924:C:C4	1:AA:1925:C:C5	2.39	1.11
1:AA:1341:U:H5''	19:AT:57:LEU:CB	1.80	1.11
4:DE:21:VAL:HB	4:DE:22:PRO:HB3	1.22	1.11

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2753:A:H2'	1:AA:2754:U:H5''	1.24	1.11
11:DO:62:LEU:O	11:DO:62:LEU:HD23	1.51	1.11
20:DU:76:CYS:SG	20:DU:77:PRO:HD2	1.90	1.11
57:DY:16:ASN:HA	57:DY:19:ARG:HH11	1.03	1.11
57:DY:18:GLU:HG2	57:DY:66:LEU:HD13	1.22	1.11
1:AA:458:G:N2	1:AA:470:A:OP2	1.82	1.10
43:CP:124:PRO:CB	43:CP:125:ARG:CG	2.29	1.10
55:DA:1083:U:H1'	55:DA:1086:A:H61	1.11	1.10
57:DY:50:ARG:C	57:DY:51:LEU:HG	1.72	1.10
1:AA:2701:C:H3'	1:AA:2702:U:C5'	1.81	1.10
1:AA:1484:G:H2'	1:AA:1485:G:H5''	1.20	1.10
43:CP:124:PRO:HB2	43:CP:125:ARG:HB2	1.15	1.10
56:DJ:12:LEU:CA	56:DJ:13:SER:CB	2.27	1.10
54:CA:630:G:O2'	54:CA:631:G:OP1	1.65	1.10
57:DY:130:THR:HG21	56:DJ:14:GLN:HE22	1.08	1.10
55:DA:1085:A:H2'	55:DA:1086:A:C8	1.84	1.10
21:DV:189:ALA:HA	21:DV:190:GLU:HG2	1.23	1.10
1:AA:611:C:H2'	1:AA:612:G:H5''	1.14	1.10
55:DA:2701:C:H3'	55:DA:2702:U:H5''	1.31	1.10
21:DV:191:VAL:HG11	21:DV:197:ILE:CG2	1.76	1.10
1:AA:458:G:H1'	1:AA:459:U:H5	1.14	1.10
52:BB:74:C:O2'	52:BB:75:C:H5'	1.48	1.10
58:DL:7:VAL:HG12	58:DL:57:ILE:CG1	1.81	1.10
2:AB:42:C:H4'	6:AG:67:LYS:HD3	1.32	1.10
54:CA:1305:G:H22	54:CA:1331:G:H2'	1.12	1.10
55:DA:654(M):C:H3'	55:DA:654(N):G:N7	1.64	1.10
57:DY:25:PHE:HB3	57:DY:82:PHE:CE1	1.86	1.10
58:DL:52:ILE:CG1	58:DL:76:TYR:HB3	1.81	1.10
43:CP:124:PRO:CB	43:CP:125:ARG:CB	2.30	1.09
50:CW:71:THR:HG22	50:CW:72:LEU:H	1.12	1.09
4:DE:61:ARG:CB	4:DE:62:PRO:HD2	1.81	1.09
34:BG:25:ARG:HB3	34:BG:25:ARG:HH11	1.10	1.09
57:DY:111:LEU:C	57:DY:112:LEU:HD23	1.73	1.09
57:DY:13:LEU:HD23	57:DY:62:ALA:HB1	1.10	1.09
21:AV:185:GLU:O	21:AV:186:GLU:HB2	1.52	1.09
55:DA:1179:C:H2'	55:DA:1180:C:H5''	1.33	1.09
17:A2:80:GLN:HA	17:A2:80:GLN:NE2	1.60	1.09
17:A2:80:GLN:CA	17:A2:80:GLN:HE21	1.64	1.09
32:CE:178:ARG:HB2	32:CE:178:ARG:HH11	1.16	1.09
55:DA:1077:A:C3'	55:DA:1078:U:H5'	1.83	1.09
17:A2:85:LYS:HG3	17:A2:87:HIS:N	1.68	1.09

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:A8:49:VAL:CG1	30:A8:50:LEU:HD23	1.83	1.09
56:DI:9:LYS:O	56:DI:10:GLU:C	1.91	1.09
49:CV:31:ILE:HG23	49:CV:49:ILE:HA	1.34	1.09
21:DV:128:VAL:HA	21:DV:161:VAL:CG2	1.82	1.09
57:DY:93:LEU:HD21	57:DY:126:ALA:HB1	1.24	1.09
1:AA:2306:C:H3'	1:AA:2307:G:H5''	1.30	1.09
1:AA:2893:G:H5'	1:AA:2894:G:H5'	1.10	1.09
58:DL:141:ALA:CB	58:DL:143:GLU:H	1.66	1.09
9:DM:115:ARG:HA	9:DM:118:LYS:HE3	1.34	1.09
57:DY:53:VAL:O	57:DY:58:LEU:HD21	1.53	1.09
57:DY:87:VAL:HG13	57:DY:91:LYS:HB2	1.20	1.09
22:D3:32:ARG:H	22:D3:35:ASN:ND2	1.51	1.09
31:BA:1129:C:H4'	31:BA:1130:A:H5'	1.15	1.08
49:CV:88:LYS:HA	49:CV:88:LYS:HE2	1.25	1.08
14:DQ:106:ARG:HH11	14:DQ:106:ARG:HB2	1.10	1.08
57:DY:25:PHE:CD1	57:DY:82:PHE:CE2	2.41	1.08
57:DY:26:LEU:HA	57:DY:112:LEU:HA	1.15	1.08
54:CA:1003:G:H2'	54:CA:1004:A:H5'	1.22	1.08
7:DH:89:ILE:HD11	7:DH:129:THR:HB	1.27	1.08
57:DY:16:ASN:CB	57:DY:19:ARG:NH1	2.10	1.08
56:DJ:15:ALA:O	56:DJ:16:THR:HG23	1.52	1.08
21:DV:191:VAL:HG13	21:DV:197:ILE:HG21	1.24	1.08
1:AA:2746:U:H4'	7:AH:138:LYS:HG3	1.36	1.08
55:DA:1077:A:H3'	55:DA:1078:U:C5'	1.82	1.08
55:DA:49:A:N7	55:DA:120:U:H5	1.50	1.08
57:DY:90:ALA:N	56:DJ:15:ALA:HB2	1.67	1.08
14:DQ:59:LYS:HG2	14:DQ:60:GLY:H	1.18	1.08
57:DY:71:LEU:CB	57:DY:112:LEU:O	2.00	1.08
54:CA:792:A:N9	54:CA:794:A:N6	2.01	1.08
2:DB:74:U:H2'	2:DB:75:G:H5''	1.36	1.08
57:DY:142:LEU:HD13	57:DY:143:GLN:H	1.08	1.08
4:DE:61:ARG:HB3	4:DE:62:PRO:CD	1.82	1.08
57:DY:58:LEU:H	57:DY:58:LEU:HD23	1.12	1.08
7:AH:22:GLY:HA2	7:AH:37:VAL:HG12	1.34	1.08
30:A8:49:VAL:HG13	30:A8:50:LEU:HD23	1.30	1.08
1:AA:1928:A:H2'	1:AA:1929:G:C5'	1.84	1.08
21:AV:176:PRO:N	21:AV:177:PRO:HD2	1.50	1.07
31:BA:279:A:H4'	31:BA:280:C:H5''	1.31	1.07
54:CA:1449:C:H2'	54:CA:1450:U:H5''	1.36	1.07
1:AA:1929:G:H4'	1:AA:1930:G:OP1	1.47	1.07
4:AE:60:ASN:HD22	4:AE:63:LEU:HB2	1.15	1.07

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AP:75:THR:HA	12:AP:88:GLY:HA2	1.33	1.07
54:CA:1399:C:H4'	54:CA:1400:C:O5'	1.50	1.07
56:DJ:12:LEU:HB3	56:DJ:13:SER:OG	1.54	1.07
57:DY:92:THR:HG22	57:DY:93:LEU:HD23	1.29	1.07
30:D8:52:LYS:H	30:D8:53:PRO:HD2	1.08	1.07
57:DY:7:VAL:HG22	57:DY:8:GLU:H	1.00	1.07
1:AA:2519:U:H4'	1:AA:2520:C:OP1	1.53	1.07
17:D2:58:VAL:HB	17:D2:98:GLU:HB2	1.28	1.07
55:DA:1061:U:H4'	55:DA:1070:A:H1'	1.09	1.07
56:DI:3:LEU:CD2	56:DI:7:ARG:HD2	1.83	1.07
56:DJ:13:SER:CA	56:DJ:17:VAL:CG2	2.31	1.07
58:DL:14:ALA:CB	58:DL:50:ASP:CB	2.31	1.07
57:DY:32:LEU:HB2	57:DY:33:PRO:CD	1.84	1.07
57:DY:9:LEU:HD13	57:DY:10:LEU:H	0.99	1.07
21:AV:131:ARG:HH11	21:AV:131:ARG:HG2	0.94	1.07
58:DL:52:ILE:HD11	58:DL:76:TYR:HB3	1.10	1.07
21:DV:128:VAL:HA	21:DV:161:VAL:HG21	1.08	1.07
1:AA:2503:A:H4'	1:AA:2504:U:OP1	1.50	1.07
54:CA:1124:G:H3'	54:CA:1145:C:H41	1.10	1.07
7:DH:126:PRO:HD2	7:DH:127:GLU:H	1.07	1.07
33:CF:70:VAL:HG12	33:CF:72:LYS:H	1.20	1.07
17:A2:71:LEU:H	17:A2:86:GLY:HA3	1.07	1.07
4:DE:61:ARG:HB3	4:DE:62:PRO:HD2	1.09	1.07
58:DL:3:LYS:O	58:DL:4:VAL:HG23	1.52	1.07
57:DY:50:ARG:CG	57:DY:51:LEU:H	1.66	1.07
57:DY:76:GLY:O	57:DY:111:LEU:HB3	1.54	1.07
55:DA:1371:G:O2'	55:DA:1372:U:H5	1.33	1.07
5:DF:32:LEU:HD11	5:DF:105:VAL:HG13	1.35	1.07
1:AA:2533:A:H2'	1:AA:2534:A:H5''	1.36	1.07
20:AU:15:VAL:HB	20:AU:22:GLY:HA3	1.35	1.07
49:BV:63:THR:H	49:BV:66:MET:HE3	1.10	1.07
21:DV:116:VAL:HG13	21:DV:117:LEU:HD12	1.11	1.07
57:DY:27:VAL:HG22	57:DY:28:ASN:N	1.61	1.07
57:DY:2:PRO:HG2	57:DY:3:ASN:H	1.19	1.07
2:AB:74:U:H2'	2:AB:75:G:H5''	1.32	1.06
32:CE:84:GLU:HB3	32:CE:219:VAL:HG21	1.36	1.06
11:DO:75:ILE:H	11:DO:75:ILE:HD13	1.15	1.06
21:DV:191:VAL:CG1	21:DV:197:ILE:CG1	2.34	1.06
54:CA:1129:C:H4'	54:CA:1130:A:H5'	1.32	1.06
1:AA:2091:U:H3'	1:AA:2092:U:C5'	1.86	1.06
31:BA:1003:G:H2'	31:BA:1004:A:H5''	1.32	1.06

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:DJ:13:SER:HB3	56:DJ:17:VAL:HG11	1.21	1.06
57:DY:93:LEU:HD22	57:DY:97:ALA:HB3	1.07	1.06
1:AA:1924:C:C2	1:AA:1925:C:C6	2.44	1.06
1:AA:266:G:H2'	1:AA:267:C:H5''	1.36	1.06
11:AO:19:VAL:HG22	11:AO:20:GLY:H	1.13	1.06
40:BM:44:VAL:HG22	40:BM:66:ARG:HG2	1.37	1.06
45:CR:87:ILE:HG22	45:CR:88:ARG:H	1.21	1.06
58:DL:112:MET:N	58:DL:113:PRO:HD2	1.69	1.06
57:DY:27:VAL:HG22	57:DY:28:ASN:H	1.03	1.06
57:DY:63:LEU:HD21	57:DY:65:GLU:OE1	1.53	1.06
57:DY:29:TYR:N	57:DY:81:VAL:HG12	0.74	1.06
22:A3:5:LYS:HE2	52:BC:73:A:O2'	1.55	1.06
39:BL:65:VAL:HG22	39:BL:66:ARG:H	1.19	1.06
6:DG:56:ALA:HB2	6:DG:153:ARG:HE	1.17	1.06
52:CB:57:G:H5''	21:DV:182:LYS:HZ1	1.18	1.06
57:DY:130:THR:HG21	56:DJ:14:GLN:NE2	1.71	1.06
21:DV:117:LEU:HD13	21:DV:118:GLN:H	0.90	1.06
39:BL:16:ARG:HB2	39:BL:16:ARG:HH11	1.09	1.06
1:AA:611:C:C2'	1:AA:612:G:H5''	1.85	1.06
3:AD:147:LEU:HD22	3:AD:155:LEU:HD11	1.37	1.06
54:CA:792:A:C8	54:CA:794:A:N6	2.23	1.06
58:DL:104:VAL:O	58:DL:107:ILE:HG22	1.29	1.06
57:DY:51:LEU:CD2	57:DY:82:PHE:C	2.12	1.06
40:BM:8:LEU:HG	40:BM:96:ILE:HG22	1.33	1.05
32:CE:7:VAL:HG21	32:CE:217:ARG:NH1	1.71	1.05
8:DK:115:ALA:HB3	8:DK:128:LEU:HD11	1.12	1.05
55:DA:1076:C:H2'	55:DA:1077:A:H5''	1.06	1.05
55:DA:1082:U:C4'	58:DL:117:THR:HG21	1.85	1.05
58:DL:42:ASN:O	58:DL:46:ALA:CB	2.05	1.05
34:CG:187:ARG:HH21	34:CG:190:ASP:HB2	1.12	1.05
46:CS:45:THR:HG22	46:CS:47:ASP:H	1.19	1.05
57:DY:60:ARG:HE	57:DY:60:ARG:HA	0.88	1.05
57:DY:19:ARG:NH1	57:DY:84:GLU:OE1	1.87	1.05
2:AB:39:A:H2'	26:A4:1:MET:CE	1.84	1.05
13:D0:33:ARG:NH2	27:D5:55:ARG:HG2	1.71	1.05
12:DP:60:ARG:HG3	21:DV:181:GLU:OE2	1.55	1.05
49:CV:87:ALA:O	49:CV:88:LYS:HD2	1.56	1.05
20:DU:95:LYS:HB3	20:DU:100:ALA:HA	1.34	1.05
21:DV:116:VAL:HG13	21:DV:117:LEU:CD1	1.84	1.05
57:DY:73:GLY:HA3	57:DY:112:LEU:HD11	1.36	1.05
40:CM:48:THR:HA	40:CM:62:HIS:HB3	1.33	1.05

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:637:A:H4'	55:DA:638:G:O5'	1.53	1.05
58:DL:141:ALA:HB1	58:DL:143:GLU:H	0.92	1.05
58:DL:7:VAL:CG1	58:DL:58:THR:N	2.19	1.05
57:DY:16:ASN:HA	57:DY:19:ARG:NH1	1.70	1.05
55:DA:2015:A:H1'	27:D5:2:ALA:HA	1.37	1.05
57:DY:135:ARG:HH11	57:DY:138:LEU:HG	1.21	1.05
21:AV:145:GLU:HA	21:AV:174:VAL:HG11	1.37	1.05
52:BD:48:C:C5	52:BD:59:U:H1'	1.90	1.05
21:DV:187:ALA:HB2	21:DV:193:GLU:HG2	1.38	1.05
57:DY:25:PHE:CB	57:DY:82:PHE:CZ	2.40	1.05
57:DY:25:PHE:CE1	57:DY:82:PHE:CB	2.40	1.05
56:DI:29:GLU:HG3	56:DJ:6:GLU:OE1	1.55	1.04
1:AA:925:C:H2'	1:AA:926:A:H5''	1.37	1.04
31:BA:1528:U:O2'	31:BA:1529:G:H5''	1.58	1.04
33:BF:150:LYS:HE2	33:BF:152:ILE:HD11	1.37	1.04
55:DA:1142(A):A:O2'	55:DA:1143:A:H3'	1.57	1.04
56:DJ:18:LEU:O	56:DJ:21:LYS:N	1.89	1.04
21:DV:61:LEU:CD1	21:DV:65:GLN:HB2	1.85	1.04
1:AA:2599:G:N7	3:AD:236:GLY:O	1.91	1.04
31:BA:696:A:H2'	31:BA:697:U:H5''	1.37	1.04
4:DE:170:LEU:HD22	4:DE:184:VAL:HG12	1.39	1.04
7:DH:153:LYS:HB3	7:DH:154:PRO:HD2	1.06	1.04
38:BK:30:ARG:HH11	38:BK:30:ARG:HB3	1.18	1.04
4:DE:14:ILE:CG2	4:DE:15:PHE:N	2.20	1.04
1:AA:2529:G:H5'	1:AA:2530:A:H5''	1.34	1.04
8:AK:79:ILE:N	8:AK:142:VAL:HG21	1.71	1.04
21:AV:163:LEU:HD23	21:AV:163:LEU:H	1.14	1.04
34:BG:12:CYS:HA	34:BG:21:LEU:HD23	1.39	1.04
54:CA:792:A:O2'	54:CA:794:A:N7	1.89	1.04
13:D0:117:VAL:HG22	13:D0:118:GLU:H	1.13	1.04
24:DW:16:LEU:HG	24:DW:16:LEU:O	1.53	1.04
57:DY:127:GLU:HG3	57:DY:128:LEU:H	1.17	1.04
1:AA:2820:A:N7	4:AE:191:PRO:HB3	1.73	1.04
42:BO:41:ARG:HB3	42:BO:41:ARG:HH11	1.18	1.04
42:BO:47:LYS:HB3	42:BO:48:PRO:CD	1.86	1.04
56:DI:24:ILE:CG1	56:DI:25:ASP:H	1.67	1.04
57:DY:93:LEU:CD2	57:DY:126:ALA:HB1	1.86	1.04
1:AA:674:G:H1'	5:AF:74:ARG:HD3	1.39	1.04
1:AA:1341:U:H5''	19:AT:57:LEU:HB3	1.05	1.04
21:AV:146:ILE:HG23	21:AV:147:GLY:H	1.22	1.04
31:BA:1226:C:H4'	31:BA:1227:A:OP1	1.56	1.04

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:87:VAL:HG13	57:DY:91:LYS:CB	1.86	1.04
54:CA:792:A:C1'	54:CA:794:A:H62	1.71	1.04
58:DL:108:ALA:C	58:DL:111:LYS:HD3	1.78	1.04
57:DY:73:GLY:O	57:DY:119:ALA:C	1.96	1.04
57:DY:29:TYR:HE2	57:DY:32:LEU:HD11	1.02	1.04
53:C1:53:U:C2'	53:C1:54:U:H5'	1.87	1.04
54:CA:630:G:H3'	54:CA:630:G:C8	1.92	1.04
54:CA:73:G:N2	54:CA:74:C:N4	2.05	1.04
55:DA:1484:G:C2'	55:DA:1485:G:H5''	1.88	1.04
31:BA:168:G:H2'	31:BA:169:C:H5''	1.39	1.04
52:CD:8:U:H2'	52:CD:13:C:H41	1.15	1.04
21:DV:117:LEU:HD13	21:DV:118:GLN:N	1.72	1.04
55:DA:896:A:N1	21:DV:178:GLU:OE2	1.89	1.04
1:AA:33:U:H4'	1:AA:34:C:OP1	1.58	1.03
31:BA:792:A:H2'	31:BA:794:A:N6	1.73	1.03
16:D1:90:VAL:HG12	16:D1:91:ASP:H	1.17	1.03
26:D4:38:LYS:O	26:D4:40:HIS:N	1.90	1.03
52:CB:57:G:H5''	21:DV:182:LYS:NZ	1.73	1.03
57:DY:26:LEU:CD2	57:DY:121:ASP:OD2	2.06	1.03
3:AD:236:GLY:O	3:AD:237:GLU:HB2	1.56	1.03
57:DY:50:ARG:HG3	57:DY:51:LEU:H	1.15	1.03
57:DY:60:ARG:NE	57:DY:60:ARG:HA	1.72	1.03
35:BH:31:LEU:HD21	35:BH:43:LEU:HD11	1.35	1.03
58:DL:19:PRO:HA	58:DL:25:PRO:CD	1.87	1.03
58:DL:21:PRO:HG2	58:DL:24:GLY:HA3	1.35	1.03
12:DP:76:LYS:N	12:DP:88:GLY:HA3	1.71	1.03
57:DY:7:VAL:HG22	57:DY:8:GLU:N	1.73	1.03
48:BU:18:ARG:O	48:BU:19:LYS:HB3	1.55	1.03
56:DJ:14:GLN:HA	56:DJ:15:ALA:C	1.75	1.03
58:DL:9:LYS:HD2	58:DL:9:LYS:H	0.90	1.03
57:DY:112:LEU:N	57:DY:112:LEU:CD2	2.16	1.03
57:DY:142:LEU:CD1	57:DY:143:GLN:H	1.70	1.03
55:DA:1083:U:C5'	57:DY:47:ASN:OD1	2.07	1.03
1:AA:1963:U:H4'	1:AA:1964:G:OP1	1.57	1.03
1:AA:893:C:H2'	1:AA:894:C:C6	1.92	1.03
8:DK:77:LEU:HD11	8:DK:140:LEU:HB2	1.37	1.03
58:DL:11:GLN:HG3	58:DL:12:LEU:H	1.20	1.03
1:AA:1928:A:C2'	1:AA:1929:G:H5'	1.88	1.03
21:AV:108:PRO:HB3	21:AV:141:VAL:O	1.53	1.03
58:DL:101:TRP:HA	58:DL:104:VAL:HB	1.39	1.03
58:DL:93:ARG:NH1	58:DL:135:GLY:HA2	1.73	1.03

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:25:PHE:CE1	57:DY:82:PHE:CD2	2.43	1.03
55:DA:2760:C:C2'	55:DA:2761:G:H5''	1.89	1.03
56:DI:30:ALA:HA	56:DJ:3:LEU:HD21	1.38	1.03
21:DV:112:ARG:HG3	21:DV:112:ARG:HH11	1.22	1.03
57:DY:130:THR:O	57:DY:134:LEU:CD1	2.06	1.03
17:A2:85:LYS:HG3	17:A2:87:HIS:H	0.89	1.03
23:AZ:87:PRO:HA	23:AZ:90:ILE:HG22	1.36	1.03
56:DJ:13:SER:OG	56:DJ:17:VAL:HG22	1.59	1.03
17:A2:71:LEU:H	17:A2:86:GLY:CA	1.70	1.03
54:CA:1139:G:H22	54:CA:1144:G:H1	1.03	1.03
55:DA:1332:G:N2	55:DA:1609:A:H2'	1.74	1.03
55:DA:889:C:H2'	55:DA:890:A:O4'	1.58	1.03
21:AV:97:GLU:HB3	21:AV:125:LEU:HD11	1.37	1.03
53:C1:53:U:H2'	53:C1:54:U:H5'	1.39	1.02
54:CA:77:C:H2'	54:CA:78:G:H5''	1.40	1.02
56:DI:7:ARG:HE	56:DI:8:ILE:CG1	1.71	1.02
56:DJ:12:LEU:CB	56:DJ:13:SER:CB	2.36	1.02
55:DA:1190:G:H5'	11:DO:32:THR:HA	1.40	1.02
58:DL:105:LEU:HD12	58:DL:106:GLU:H	0.91	1.02
58:DL:7:VAL:CG1	58:DL:58:THR:O	2.07	1.02
57:DY:104:ILE:CG1	57:DY:105:PRO:HD2	1.87	1.02
57:DY:51:LEU:HD13	57:DY:81:VAL:C	1.77	1.02
54:CA:1178:G:H5'	39:CL:93:ARG:HH21	1.18	1.02
56:DI:7:ARG:HE	56:DI:8:ILE:HG12	1.23	1.02
58:DL:120:LEU:O	58:DL:121:GLU:HB2	1.59	1.02
58:DL:52:ILE:CG1	58:DL:76:TYR:CB	2.37	1.02
31:BA:1322:C:O2'	31:BA:1323:G:H5'	1.58	1.02
17:D2:35:LEU:HD21	17:D2:57:VAL:HG22	1.39	1.02
12:DP:79:LEU:O	22:D3:4:LYS:NZ	1.92	1.02
55:DA:1061:U:H4'	55:DA:1070:A:C1'	1.88	1.02
4:DE:170:LEU:HD23	4:DE:185:LYS:HB2	1.41	1.02
57:DY:112:LEU:CD1	57:DY:121:ASP:HB2	1.88	1.02
1:AA:2820:A:C5	4:AE:191:PRO:HB2	1.95	1.02
47:CT:67:LYS:HA	47:CT:70:ARG:HH12	1.20	1.02
42:BO:23:LYS:HD3	42:BO:23:LYS:H	1.21	1.02
12:AP:77:LYS:HZ3	12:AP:82:ARG:HA	1.23	1.02
56:DI:30:ALA:HA	56:DJ:3:LEU:CD2	1.89	1.02
12:DP:76:LYS:H	12:DP:88:GLY:HA3	0.86	1.02
57:DY:112:LEU:HD13	57:DY:121:ASP:OD2	1.59	1.02
1:AA:1928:A:H2'	1:AA:1929:G:H5'	1.05	1.02
8:AK:79:ILE:H	8:AK:142:VAL:HG21	0.85	1.02

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:DL:83:GLY:H	58:DL:99:ILE:CG2	1.72	1.02
58:DL:99:ILE:HG13	58:DL:138:VAL:HG21	1.38	1.02
57:DY:27:VAL:CB	57:DY:111:LEU:N	2.23	1.02
17:A2:5:VAL:HG23	17:A2:37:VAL:HG11	1.40	1.02
1:AA:2645:G:H4'	1:AA:2732:G:O2'	1.59	1.02
11:AO:75:ILE:HD13	11:AO:75:ILE:H	1.20	1.02
31:BA:1116:C:H2'	31:BA:1117:G:H5''	1.34	1.02
54:CA:562:C:O2'	42:CO:15:ARG:HB3	1.59	1.02
4:DE:24:THR:HG21	4:DE:188:VAL:HG11	1.41	1.02
5:DF:66:PRO:O	5:DF:67:GLN:HB3	1.56	1.02
1:AA:1024:G:H3'	1:AA:1025:G:H5''	1.40	1.02
1:AA:2820:A:H62	4:AE:192:ASN:N	1.51	1.02
4:AE:60:ASN:C	4:AE:62:PRO:HD2	1.81	1.01
49:BV:42:PRO:O	49:BV:45:VAL:N	1.93	1.01
7:DH:4:ILE:HG13	7:DH:6:ARG:NE	1.75	1.01
58:DL:105:LEU:HD12	58:DL:106:GLU:N	1.75	1.01
21:DV:191:VAL:HG11	21:DV:197:ILE:HG12	1.41	1.01
8:AK:109:ILE:H	8:AK:109:ILE:HD13	1.24	1.01
52:CD:21:A:H2'	52:CD:22:G:H5''	1.42	1.01
38:CK:6:ILE:HB	38:CK:85:ARG:HH12	1.19	1.01
21:DV:191:VAL:HG11	21:DV:197:ILE:HG13	1.37	1.01
43:CP:88:ARG:HB3	43:CP:88:ARG:HH11	1.19	1.01
58:DL:9:LYS:N	58:DL:9:LYS:HD2	1.75	1.01
57:DY:71:LEU:HB2	57:DY:112:LEU:C	1.80	1.01
57:DY:49:ALA:N	57:DY:84:GLU:HB2	1.75	1.01
57:DY:18:GLU:HG2	57:DY:66:LEU:CD1	1.89	1.01
1:AA:1484:G:C2'	1:AA:1485:G:H5''	1.90	1.01
31:BA:56:U:H2'	31:BA:57:G:C8	1.95	1.01
31:BA:1329:A:H5''	43:BP:25:ILE:O	1.58	1.01
58:DL:18:THR:CG2	58:DL:38:VAL:CG1	2.38	1.01
31:BA:547:A:H4'	31:BA:548:G:O5'	1.54	1.01
49:CV:84:GLY:HA2	49:CV:87:ALA:HB3	1.42	1.01
55:DA:2635:C:H5''	4:DE:78:LEU:HA	1.42	1.01
55:DA:603:A:H4'	55:DA:604:G:O5'	1.55	1.01
55:DA:762:U:H4'	55:DA:763:G:O5'	1.59	1.01
21:AV:141:VAL:HG21	21:AV:144:LEU:HD23	1.41	1.01
31:BA:77:C:H2'	31:BA:78:G:H5''	1.38	1.01
55:DA:1082:U:H4'	58:DL:117:THR:HG21	1.02	1.01
5:DF:107:LYS:HD2	5:DF:206:ILE:HD13	1.43	1.01
58:DL:73:PRO:HB3	58:DL:77:LEU:HD13	1.41	1.01
21:AV:69:THR:HG22	21:AV:90:VAL:HA	1.41	1.01

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1363:A:H4'	54:CA:1364:U:H5''	1.37	1.01
24:DW:41:ILE:HD11	24:DW:44:LEU:HB2	1.37	1.01
7:AH:153:LYS:HB3	7:AH:161:GLY:HA2	1.39	1.01
20:AU:97:ARG:HH21	20:AU:98:VAL:HB	1.26	1.01
17:D2:35:LEU:HB2	17:D2:37:VAL:HG22	1.37	1.01
57:DY:54:ALA:HB1	57:DY:57:THR:HB	1.38	1.01
55:DA:896:A:C2	21:DV:178:GLU:OE2	2.14	1.01
31:BA:792:A:C2'	31:BA:794:A:H62	1.74	1.01
56:DI:16:THR:HG23	56:DI:17:VAL:H	1.23	1.01
58:DL:19:PRO:C	58:DL:25:PRO:HD2	1.80	1.01
58:DL:57:ILE:HD13	58:DL:58:THR:N	1.74	1.01
21:DV:150:LEU:HD22	21:DV:151:HIS:N	1.76	1.01
57:DY:92:THR:CG2	57:DY:93:LEU:HD23	1.90	1.01
1:AA:481:G:OP2	20:AU:47:LYS:HB2	1.60	1.00
12:AP:82:ARG:HG2	12:AP:82:ARG:HH11	1.26	1.00
23:AZ:78:LYS:HD2	23:AZ:80:LEU:HD21	1.39	1.00
55:DA:1063:G:H1'	58:DL:134:MET:HE1	1.42	1.00
55:DA:1934:C:H5'	55:DA:1934:C:H6	1.23	1.00
3:DD:35:LYS:HG2	3:DD:64:ILE:N	1.73	1.00
4:DE:4:ILE:HD12	4:DE:28:ALA:HB1	1.40	1.00
58:DL:14:ALA:HB1	58:DL:50:ASP:HB2	1.41	1.00
21:DV:150:LEU:HD21	21:DV:154:ASP:CB	1.90	1.00
10:AN:4:PRO:O	10:AN:5:GLN:HB2	1.57	1.00
35:CH:40:ARG:HH11	35:CH:40:ARG:HB3	1.25	1.00
43:CP:124:PRO:CB	43:CP:125:ARG:HB2	1.90	1.00
55:DA:1061:U:C4'	55:DA:1070:A:H1'	1.91	1.00
56:DI:24:ILE:N	56:DI:27:LEU:HD12	1.75	1.00
21:DV:116:VAL:O	21:DV:174:VAL:HA	1.61	1.00
21:DV:187:ALA:CB	21:DV:193:GLU:HG2	1.92	1.00
6:DG:112:PRO:HB3	26:D4:37:SER:H	1.25	1.00
1:AA:384:U:H2'	1:AA:385:C:H5'	1.01	1.00
40:BM:40:LEU:HB3	40:BM:69:ASN:HB3	1.39	1.00
56:DJ:13:SER:HB3	56:DJ:17:VAL:CG2	1.73	1.00
57:DY:71:LEU:HD22	57:DY:72:ASP:H	1.24	1.00
55:DA:1533:C:H2'	55:DA:1534:G:N7	1.77	1.00
21:DV:116:VAL:HG11	21:DV:118:GLN:OE1	1.61	1.00
53:C1:52:U:O2'	53:C1:53:U:H5''	1.60	1.00
45:CR:82:ILE:HD11	45:CR:88:ARG:HB2	1.44	1.00
30:D8:36:LYS:HB3	30:D8:40:GLU:HG2	1.44	1.00
57:DY:134:LEU:HD23	56:DJ:19:GLU:OE1	1.61	1.00
1:AA:585:G:N1	1:AA:1253:A:OP1	1.94	1.00

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:894:C:H3'	1:AA:895:U:H6	1.22	1.00
52:BD:35:A:N6	52:BD:37:MIA:H153	1.77	1.00
43:CP:124:PRO:HB2	43:CP:125:ARG:CB	1.88	1.00
27:D5:33:CYS:HB2	27:D5:40:LYS:HD3	1.40	1.00
55:DA:1064:C:H4'	58:DL:89:HIS:HA	1.43	1.00
21:DV:116:VAL:CB	21:DV:175:VAL:O	2.09	1.00
1:AA:458:G:H1'	1:AA:459:U:C5	1.96	1.00
31:BA:1443:G:H3'	31:BA:1446:A:H5''	1.43	1.00
52:CB:74:C:O2'	52:CB:75:C:OP2	1.78	1.00
3:DD:35:LYS:HD2	3:DD:104:TYR:CD1	1.96	1.00
57:DY:24:PHE:CZ	57:DY:88:ALA:HB2	1.96	1.00
1:AA:1899:G:N2	1:AA:1902:C:N4	1.99	1.00
55:DA:1061:U:H5	58:DL:54:PRO:HB3	1.26	1.00
57:DY:93:LEU:CD2	57:DY:97:ALA:HB3	1.91	1.00
1:AA:1212:G:H1'	1:AA:1237:A:N6	1.77	0.99
21:AV:175:VAL:CG1	21:AV:176:PRO:CD	2.34	0.99
31:BA:872:A:H4'	31:BA:873:A:OP1	1.61	0.99
39:BL:16:ARG:NH1	39:BL:16:ARG:HB2	1.77	0.99
43:BP:83:ASP:OD1	43:BP:84:ILE:CD1	2.09	0.99
55:DA:2131:G:H5'	55:DA:2132:U:H5''	1.42	0.99
54:CA:1028:C:H2'	54:CA:1028(A):C:H5''	1.42	0.99
54:CA:630:G:H2'	54:CA:631:G:C5'	1.92	0.99
58:DL:107:ILE:CG2	58:DL:108:ALA:H	1.74	0.99
56:DJ:18:LEU:HA	56:DJ:21:LYS:HB2	1.43	0.99
55:DA:265:A:O2'	55:DA:266:G:H4'	1.61	0.99
58:DL:20:ALA:H	58:DL:25:PRO:CD	1.62	0.99
54:CA:1002:G:H2'	54:CA:1003:G:C8	1.96	0.99
4:DE:21:VAL:HB	4:DE:22:PRO:CB	1.91	0.99
13:A0:37:THR:HG22	13:A0:39:PRO:HD2	1.39	0.99
5:AF:24:LEU:HB3	5:AF:25:PRO:HD2	1.44	0.99
43:BP:3:ARG:HD2	43:BP:9:ILE:HD11	1.41	0.99
54:CA:980:C:H5'	54:CA:981:U:OP2	1.62	0.99
9:DM:134:ARG:H	9:DM:135:PRO:HD3	1.26	0.99
23:AZ:91:LYS:HA	23:AZ:91:LYS:HE3	1.45	0.99
31:BA:887:G:H2'	31:BA:888:G:H5'	1.01	0.99
54:CA:188:U:H2'	54:CA:189:U:H5''	1.44	0.99
26:D4:58:ARG:HA	26:D4:62:ARG:HB3	1.45	0.99
58:DL:57:ILE:HD13	58:DL:58:THR:H	0.88	0.99
1:AA:2820:A:N7	4:AE:191:PRO:CB	2.25	0.99
20:AU:61:ILE:HG22	20:AU:62:GLU:H	1.28	0.99
32:CE:54:THR:HG21	32:CE:201:ILE:HD11	1.43	0.99

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:DI:28:LYS:HA	56:DI:28:LYS:CE	1.89	0.99
58:DL:69:THR:HG22	58:DL:70:LYS:H	1.26	0.99
11:DO:122:PRO:HB3	11:DO:141:ALA:HB1	1.44	0.99
54:CA:630:G:H2'	54:CA:631:G:H5''	1.01	0.99
21:AV:145:GLU:O	21:AV:146:ILE:HD13	1.61	0.99
49:BV:5:LEU:HD22	49:BV:6:LYS:H	1.27	0.99
39:CL:8:GLY:HA2	39:CL:79:LEU:HD12	1.44	0.99
56:DJ:13:SER:HB3	56:DJ:17:VAL:HG21	1.10	0.99
58:DL:9:LYS:CD	58:DL:9:LYS:H	1.76	0.99
21:DV:150:LEU:HD21	21:DV:154:ASP:CG	1.81	0.99
20:AU:63:LYS:HZ2	20:AU:64:GLU:H	1.09	0.98
26:D4:69:LYS:HD3	26:D4:70:GLY:N	1.76	0.98
55:DA:1026:U:H4'	55:DA:1027:A:OP1	1.61	0.98
57:DY:25:PHE:CZ	57:DY:82:PHE:CB	2.46	0.98
57:DY:50:ARG:HG3	57:DY:51:LEU:N	1.76	0.98
1:AA:654(B):C:H2'	1:AA:654(C):G:O4'	1.61	0.98
58:DL:11:GLN:HG2	58:DL:41:PHE:HZ	1.27	0.98
57:DY:51:LEU:HD21	57:DY:82:PHE:O	0.81	0.98
1:AA:2572:A:C8	4:AE:144:ARG:HD2	1.97	0.98
7:DH:153:LYS:CB	7:DH:154:PRO:HD2	1.93	0.98
57:DY:75:GLN:HE21	57:DY:76:GLY:H	1.04	0.98
55:DA:1060:U:OP1	58:DL:54:PRO:CG	2.11	0.98
55:DA:1318:C:H2'	55:DA:1319:G:H5''	1.42	0.98
58:DL:112:MET:HG3	58:DL:118:THR:O	1.64	0.98
21:DV:118:GLN:HA	21:DV:118:GLN:NE2	1.77	0.98
57:DY:9:LEU:HD13	57:DY:10:LEU:N	1.78	0.98
31:BA:1028:C:H2'	31:BA:1028(A):C:H5''	1.42	0.98
57:DY:9:LEU:CD1	57:DY:10:LEU:H	1.75	0.98
1:AA:1341:U:O2'	1:AA:1397:U:O2'	1.79	0.98
1:AA:1924:C:N3	1:AA:1925:C:C5	2.30	0.98
32:BE:224:GLN:HA	32:BE:229:VAL:HG22	1.43	0.98
43:BP:83:ASP:OD1	43:BP:84:ILE:HD12	1.61	0.98
55:DA:2760:C:H2'	55:DA:2761:G:H5''	1.45	0.98
2:DB:74:U:C2'	2:DB:75:G:H5''	1.93	0.98
58:DL:14:ALA:HB2	58:DL:50:ASP:CB	1.93	0.98
20:DU:97:ARG:HH21	20:DU:98:VAL:HB	1.27	0.98
57:DY:27:VAL:HG23	57:DY:110:GLY:HA3	1.46	0.98
1:AA:2701:C:H3'	1:AA:2702:U:H5''	0.98	0.98
9:AM:133:GLN:HG2	9:AM:135:PRO:HD3	1.45	0.98
55:DA:1798:U:H5'	3:DD:259:THR:HG22	1.41	0.98
55:DA:905:U:H2'	55:DA:906:G:H5''	1.45	0.98

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:DI:7:ARG:NE	56:DI:8:ILE:HG12	1.77	0.98
58:DL:20:ALA:N	58:DL:25:PRO:HG2	1.71	0.98
30:A8:49:VAL:HG12	30:A8:50:LEU:H	1.12	0.98
20:AU:91:GLU:HG3	20:AU:92:ASN:H	1.28	0.98
8:DK:92:VAL:HG13	8:DK:120:ILE:HG23	1.44	0.98
58:DL:105:LEU:CD1	58:DL:106:GLU:H	1.75	0.98
12:DP:76:LYS:H	12:DP:88:GLY:CA	1.75	0.98
57:DY:71:LEU:CA	57:DY:113:GLN:HB3	1.92	0.98
57:DY:132:ASP:C	57:DY:134:LEU:H	1.66	0.98
21:AV:175:VAL:O	21:AV:177:PRO:HG2	1.62	0.98
35:BH:43:LEU:H	35:BH:65:ASN:HD22	1.09	0.98
56:DI:24:ILE:HA	56:DI:27:LEU:HD13	1.45	0.98
58:DL:95:LYS:HB3	58:DL:136:VAL:HG21	1.42	0.98
1:AA:2756:U:O2'	1:AA:2757:A:H5''	1.64	0.98
1:AA:387:U:O2'	1:AA:388:G:C5'	2.11	0.98
1:AA:896:A:H5'	1:AA:897:C:OP2	1.63	0.98
52:CC:58:A:H4'	52:CC:59:U:OP1	1.63	0.98
57:DY:29:TYR:CE2	57:DY:32:LEU:CD1	2.46	0.98
21:AV:146:ILE:CG2	21:AV:147:GLY:H	1.77	0.97
55:DA:654(B):C:H2'	55:DA:654(C):G:O4'	1.61	0.97
5:DF:103:LYS:HA	5:DF:106:ARG:HG3	1.45	0.97
58:DL:138:VAL:O	58:DL:139:VAL:HB	1.62	0.97
30:A8:32:LEU:HD23	30:A8:34:TRP:N	1.79	0.97
1:AA:1359:A:H8	1:AA:1359:A:C3'	1.73	0.97
1:AA:90:U:H2'	1:AA:90:U:O2	1.64	0.97
31:BA:887:G:H2'	31:BA:888:G:C5'	1.93	0.97
55:DA:1083:U:H2'	55:DA:1085:A:OP2	1.62	0.97
6:DG:107:LEU:O	26:D4:38:LYS:HG2	1.64	0.97
11:DO:64:LYS:C	11:DO:66:GLY:H	1.66	0.97
57:DY:7:VAL:HG13	57:DY:8:GLU:N	1.77	0.97
26:A4:53:GLU:HG3	26:A4:54:GLY:H	1.25	0.97
3:AD:35:LYS:HG2	3:AD:64:ILE:N	1.77	0.97
20:AU:38:ILE:HG22	20:AU:66:PRO:HA	1.41	0.97
21:AV:108:PRO:CG	21:AV:141:VAL:O	2.11	0.97
58:DL:25:PRO:HA	58:DL:27:LEU:HG	1.43	0.97
57:DY:24:PHE:O	57:DY:25:PHE:O	1.82	0.97
1:AA:2392:A:H2	1:AA:2424:C:H42	1.08	0.97
54:CA:1363:A:H1'	54:CA:1365:G:N7	1.79	0.97
57:DY:80:VAL:HG12	57:DY:81:VAL:N	1.73	0.97
13:A0:73:VAL:O	13:A0:76:VAL:HG12	1.64	0.97
17:A2:69:LYS:HD3	17:A2:85:LYS:HD3	1.46	0.97

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2091:U:H3'	1:AA:2092:U:H5'	1.44	0.97
1:AA:434:U:H4'	1:AA:435:C:OP1	1.63	0.97
4:AE:60:ASN:ND2	4:AE:63:LEU:HB2	1.78	0.97
58:DL:11:GLN:HG3	58:DL:12:LEU:N	1.77	0.97
58:DL:52:ILE:HD11	58:DL:76:TYR:CB	1.93	0.97
14:DQ:88:ASP:O	14:DQ:89:ARG:HB3	1.61	0.97
1:AA:1267:U:C4	1:AA:2012:G:C2	2.53	0.97
5:AF:8:GLN:HG3	5:AF:126:VAL:HA	1.45	0.97
1:AA:1341:U:C5'	19:AT:57:LEU:HB3	1.94	0.97
43:CP:124:PRO:HB3	43:CP:125:ARG:CB	1.93	0.97
46:CS:4:ILE:HD11	46:CS:64:ALA:HB1	1.47	0.97
55:DA:2712:U:HO2'	55:DA:2712(A):A:H8	0.98	0.97
3:AD:35:LYS:NZ	3:AD:104:TYR:HB2	1.79	0.97
19:AT:34:ALA:HB1	19:AT:39:ILE:HD11	1.45	0.97
31:BA:251:G:C6	31:BA:266:G:O6	2.18	0.97
31:BA:872:A:O2'	31:BA:873:A:H3'	1.63	0.97
50:CW:48:LYS:HB3	50:CW:51:GLU:HG3	1.44	0.97
55:DA:1076:C:C2'	55:DA:1077:A:H5''	1.93	0.97
57:DY:111:LEU:C	57:DY:112:LEU:CD2	2.32	0.97
57:DY:60:ARG:HE	57:DY:60:ARG:CA	1.77	0.97
1:AA:27:G:HO2'	1:AA:28:A:H8	1.08	0.97
19:AT:18:TYR:HA	19:AT:21:PHE:HD2	1.30	0.97
37:BJ:73:MET:HG2	37:BJ:90:GLU:HA	1.44	0.97
55:DA:593:G:O2'	30:D8:61:LEU:HD13	1.65	0.97
57:DY:132:ASP:O	57:DY:134:LEU:HD22	1.65	0.97
1:AA:2111:C:H41	1:AA:2147:G:N2	1.61	0.97
31:BA:1502:A:H2	31:BA:1505:G:H1	1.11	0.97
50:CW:100:ILE:HG13	50:CW:102:GLY:H	1.27	0.97
55:DA:887:A:HO2'	55:DA:889:C:H5	1.13	0.97
56:DI:21:LYS:O	56:DI:26:ALA:CB	2.12	0.97
57:DY:90:ALA:H	56:DJ:15:ALA:HB2	1.20	0.97
21:DV:190:GLU:O	21:DV:191:VAL:CG2	2.12	0.97
57:DY:27:VAL:HA	57:DY:111:LEU:HD13	1.42	0.97
7:AH:152:ARG:HG3	7:AH:153:LYS:HG2	1.46	0.96
12:AP:75:THR:HG22	12:AP:88:GLY:HA3	1.44	0.96
31:BA:410:G:OP2	34:BG:25:ARG:HG2	1.65	0.96
55:DA:1454:U:H4'	55:DA:1455:G:OP1	1.62	0.96
55:DA:890:A:H3'	55:DA:892:G:H8	1.30	0.96
56:DJ:5:ILE:CG2	56:DJ:9:LYS:HG3	1.95	0.96
58:DL:18:THR:HG22	58:DL:38:VAL:CG1	1.94	0.96
55:DA:2419:U:H4'	28:D6:23:THR:HG21	1.43	0.96

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:DH:126:PRO:CD	7:DH:127:GLU:H	1.76	0.96
27:A5:4:HIS:HB3	27:A5:5:PRO:CD	1.94	0.96
8:AK:79:ILE:H	8:AK:142:VAL:CG2	1.78	0.96
21:AV:145:GLU:OE1	21:AV:174:VAL:CG1	2.13	0.96
31:BA:1129:C:C4'	31:BA:1130:A:H5'	1.95	0.96
27:D5:4:HIS:HB3	27:D5:5:PRO:CD	1.94	0.96
3:DD:32:SER:O	3:DD:33:LEU:HB2	1.65	0.96
21:DV:61:LEU:HD13	21:DV:62:PRO:O	1.65	0.96
57:DY:138:LEU:O	57:DY:139:VAL:HG12	1.65	0.96
57:DY:14:LYS:HA	57:DY:14:LYS:HE3	1.48	0.96
5:AF:161:GLU:HG2	5:AF:164:ARG:HH22	1.28	0.96
54:CA:1182:G:H4'	54:CA:1183:A:H5''	1.46	0.96
55:DA:2287:A:N6	55:DA:2344:U:H3	1.62	0.96
55:DA:2469:A:C2	55:DA:2481:G:N2	2.33	0.96
56:DI:29:GLU:CA	56:DJ:2:ALA:HB1	1.94	0.96
57:DY:118:THR:CG2	57:DY:119:ALA:H	1.71	0.96
55:DA:1077:A:H3'	55:DA:1078:U:H5'	0.97	0.96
56:DI:19:GLU:O	56:DI:20:LEU:O	1.83	0.96
57:DY:27:VAL:HG21	57:DY:110:GLY:N	1.79	0.96
6:DG:67:LYS:HE2	26:D4:6:HIS:NE2	1.79	0.96
55:DA:2503:A:H4'	55:DA:2504:U:OP1	1.62	0.96
58:DL:50:ASP:H	58:DL:53:VAL:HG21	1.30	0.96
57:DY:8:GLU:OE1	57:DY:52:PHE:HD1	1.48	0.96
26:A4:60:GLN:HE21	26:A4:60:GLN:N	1.62	0.96
54:CA:1053:G:H5'	54:CA:1054:C:H5'	1.46	0.96
48:CU:18:ARG:H	48:CU:18:ARG:CD	1.79	0.96
55:DA:1083:U:C4'	57:DY:41:ARG:HD3	1.95	0.96
55:DA:84:A:H4'	55:DA:85:G:O5'	1.65	0.96
57:DY:93:LEU:HG	57:DY:126:ALA:C	1.86	0.96
57:DY:23:SER:HG	57:DY:114:GLY:HA2	1.22	0.96
29:A7:12:ARG:HD3	29:A7:46:VAL:HG21	1.47	0.96
1:AA:2758:A:H2'	1:AA:2759:G:H5''	1.45	0.96
19:AT:63:LYS:HE3	19:AT:63:LYS:H	1.30	0.96
31:BA:56:U:H2'	31:BA:57:G:H8	1.30	0.96
55:DA:51:G:O2'	55:DA:119:A:N1	1.97	0.96
58:DL:54:PRO:HD2	58:DL:72:PRO:HA	1.46	0.96
15:DR:50:ILE:HD11	15:DR:102:ILE:HD11	1.46	0.96
16:A1:95:LEU:C	16:A1:97:ASP:H	1.60	0.96
3:AD:255:LYS:O	3:AD:255:LYS:HD2	1.65	0.96
8:AK:5:LEU:HD11	8:AK:19:VAL:HG12	1.45	0.96
31:BA:1363:A:H1'	31:BA:1365:G:N7	1.81	0.96

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:DD:35:LYS:NZ	3:DD:104:TYR:HB2	1.80	0.96
57:DY:40:LEU:CD2	57:DY:50:ARG:NH1	2.29	0.96
1:AA:1225:C:H4'	17:A2:85:LYS:HB2	1.48	0.95
11:AO:64:LYS:HE3	30:A8:30:ARG:CZ	1.96	0.95
1:AA:2394:C:OP1	11:AO:63:PRO:HD2	1.64	0.95
20:AU:87:LYS:HB3	20:AU:92:ASN:HB3	1.46	0.95
2:DB:20:C:H2'	2:DB:21:G:H5''	1.48	0.95
58:DL:86:LYS:HE2	58:DL:86:LYS:CA	1.94	0.95
57:DY:50:ARG:O	57:DY:51:LEU:HG	1.65	0.95
31:BA:562:C:O2'	42:BO:15:ARG:HB3	1.67	0.95
55:DA:2147:G:H2'	55:DA:2148:G:O4'	1.66	0.95
24:DW:50:ILE:HD12	24:DW:51:ARG:H	1.31	0.95
3:AD:242:ARG:H	3:AD:242:ARG:HD2	1.31	0.95
21:DV:192:ALA:O	21:DV:194:PRO:HD3	1.66	0.95
57:DY:70:GLU:O	57:DY:71:LEU:HD12	1.64	0.95
3:AD:32:SER:HA	3:AD:36:PRO:HD2	1.45	0.95
55:DA:1169:G:H2'	55:DA:1170:G:H5''	1.48	0.95
3:DD:43:ARG:NH1	3:DD:44:ASN:OD1	1.99	0.95
21:DV:146:ILE:HA	21:DV:174:VAL:HB	1.45	0.95
4:AE:48:GLN:HG2	4:AE:78:LEU:HD12	1.48	0.95
24:AW:14:ARG:HG2	24:AW:15:LYS:HE3	1.47	0.95
25:AX:59:VAL:HG12	25:AX:60:GLU:H	1.28	0.95
6:DG:67:LYS:HE2	26:D4:6:HIS:CE1	2.00	0.95
29:D7:8:ASN:ND2	29:D7:11:LYS:H	1.64	0.95
56:DI:20:LEU:C	56:DI:24:ILE:CG2	2.35	0.95
58:DL:7:VAL:HG13	58:DL:57:ILE:HD12	1.45	0.95
57:DY:134:LEU:CA	57:DY:137:GLU:HG2	1.96	0.95
57:DY:28:ASN:OD1	57:DY:83:TYR:CE2	2.18	0.95
1:AA:673:C:H5'	5:AF:54:ARG:HH12	1.31	0.95
3:AD:108:PRO:HG2	3:AD:111:LEU:HB2	1.46	0.95
21:AV:146:ILE:CG2	21:AV:147:GLY:N	2.30	0.95
35:BH:51:VAL:HB	35:BH:52:PRO:HD3	1.46	0.95
55:DA:792:G:H5''	55:DA:793:A:H5'	1.46	0.95
8:DK:78:THR:HA	8:DK:141:LYS:HB2	1.46	0.95
21:DV:118:GLN:HE21	21:DV:118:GLN:HA	1.28	0.95
21:DV:150:LEU:HD22	21:DV:151:HIS:H	1.25	0.95
57:DY:27:VAL:HG21	57:DY:110:GLY:HA3	1.19	0.95
57:DY:141:VAL:HG13	57:DY:142:LEU:H	1.31	0.95
57:DY:16:ASN:CA	57:DY:19:ARG:HH11	1.73	0.95
57:DY:50:ARG:CA	57:DY:83:TYR:CD1	2.49	0.95
7:AH:92:ILE:HD12	7:AH:92:ILE:H	1.31	0.95

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:49:A:H5''	55:DA:50:U:H3'	1.48	0.95
56:DI:17:VAL:HA	56:DI:20:LEU:HD12	1.48	0.95
56:DJ:4:ASP:O	56:DJ:8:ILE:HB	1.64	0.95
9:AM:73:THR:HG22	9:AM:84:LYS:HB3	1.49	0.95
52:BB:69:G:H2'	52:BB:70:G:H5''	1.49	0.95
17:D2:52:VAL:HG21	17:D2:55:ALA:HB3	1.48	0.95
4:DE:78:LEU:HD23	4:DE:79:ARG:HB2	1.49	0.95
24:DW:65:ASN:HD22	24:DW:69:ARG:NH2	1.65	0.95
57:DY:118:THR:HG23	57:DY:119:ALA:N	1.70	0.95
1:AA:1928:A:C2'	1:AA:1929:G:C5'	2.45	0.95
1:AA:654(M):C:H3'	1:AA:654(N):G:N7	1.81	0.95
1:AA:90:U:HO2'	1:AA:91:A:H8	0.98	0.95
15:AR:16:ARG:NH2	15:AR:19:LEU:HD21	1.82	0.95
43:BP:92:HIS:CE1	43:BP:98:VAL:HG21	2.02	0.95
15:DR:26:ASP:HB3	15:DR:91:ARG:HA	1.47	0.95
57:DY:73:GLY:O	57:DY:120:LYS:N	1.98	0.95
17:A2:71:LEU:N	17:A2:86:GLY:HA3	1.81	0.95
1:AA:2701:C:C3'	1:AA:2702:U:H5''	1.95	0.95
7:AH:106:THR:HG22	7:AH:112:PRO:HB3	1.48	0.95
21:AV:146:ILE:C	21:AV:148:ASP:H	1.71	0.95
21:AV:144:LEU:C	21:AV:146:ILE:H	1.69	0.95
27:D5:55:ARG:HG3	27:D5:57:VAL:H	1.32	0.95
57:DY:107:VAL:HG12	57:DY:108:LYS:N	1.79	0.95
57:DY:26:LEU:N	57:DY:82:PHE:CZ	2.35	0.95
53:B1:36:G:H2'	53:B1:37:G:H5''	1.48	0.94
31:BA:1007:C:H2'	31:BA:1008:C:H5''	1.49	0.94
49:BV:9:VAL:HG12	49:BV:10:PHE:H	1.32	0.94
54:CA:1313:U:OP2	49:CV:6:LYS:CB	2.14	0.94
54:CA:1313:U:P	49:CV:6:LYS:CB	2.55	0.94
49:CV:40:ILE:HG12	49:CV:41:VAL:HG22	1.49	0.94
55:DA:1082:U:H4'	58:DL:117:THR:CG2	1.96	0.94
55:DA:676:A:H8	55:DA:2069:G:N2	1.64	0.94
28:A6:41:PRO:HD2	28:A6:45:LYS:O	1.63	0.94
1:AA:458:G:O2'	1:AA:459:U:OP2	1.85	0.94
52:BD:21:A:H2'	52:BD:22:G:H5''	1.47	0.94
52:CD:41:C:H2'	52:CD:42:C:H5''	1.49	0.94
49:CV:88:LYS:HA	49:CV:88:LYS:CE	1.91	0.94
16:D1:83:LEU:HA	16:D1:88:ILE:HD11	1.46	0.94
26:D4:7:PRO:HB2	26:D4:27:THR:HG21	1.46	0.94
55:DA:2134:A:H62	55:DA:2157:G:H1'	1.31	0.94
8:DK:95:LYS:HA	8:DK:111:PRO:HG3	1.49	0.94

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:A6:36:LEU:HD23	28:A6:50:ARG:HD2	1.48	0.94
20:AU:13:VAL:HG23	20:AU:73:ARG:O	1.66	0.94
20:AU:75:ILE:HG13	20:AU:79:CYS:HA	1.46	0.94
24:AW:70:GLN:HG2	24:AW:71:ASN:H	1.28	0.94
54:CA:266:G:H5''	54:CA:268:C:H41	1.31	0.94
54:CA:579:G:H5'	54:CA:728:A:H1'	1.47	0.94
55:DA:1869:G:H5'	55:DA:1870:C:OP2	1.65	0.94
55:DA:803:U:H6	55:DA:803:U:H5'	1.29	0.94
58:DL:113:PRO:O	58:DL:114:ASP:HB3	1.65	0.94
9:DM:96:GLU:C	9:DM:98:VAL:H	1.68	0.94
21:DV:191:VAL:HG12	21:DV:197:ILE:CG2	1.95	0.94
28:A6:48:VAL:HG13	28:A6:49:HIS:H	1.32	0.94
1:AA:1212:G:H1'	1:AA:1237:A:H61	1.29	0.94
1:AA:265:A:O2'	1:AA:266:G:H4'	1.67	0.94
4:AE:200:GLU:HG2	4:AE:201:THR:H	1.32	0.94
31:BA:1534:A:H2'	31:BA:1535:C:C6	2.02	0.94
54:CA:38:G:C2	54:CA:397:A:H2	1.84	0.94
26:D4:68:ARG:CA	26:D4:68:ARG:CZ	2.45	0.94
4:DE:14:ILE:HG22	4:DE:15:PHE:H	1.30	0.94
58:DL:86:LYS:HE2	58:DL:86:LYS:HA	1.49	0.94
24:DW:50:ILE:CD1	24:DW:51:ARG:H	1.80	0.94
1:AA:608:A:C4	1:AA:621:A:N6	2.34	0.94
8:AK:142:VAL:HG22	8:AK:143:SER:N	1.83	0.94
24:AW:50:ILE:HD12	24:AW:51:ARG:H	1.31	0.94
55:DA:1019:U:H3	55:DA:1142(A):A:H62	1.01	0.94
58:DL:19:PRO:CA	58:DL:25:PRO:CG	2.46	0.94
57:DY:27:VAL:HG21	57:DY:109:SER:C	1.88	0.94
22:A3:82:ARG:HG3	22:A3:84:LEU:HD13	1.46	0.94
30:D8:43:GLN:C	30:D8:44:LYS:HD2	1.88	0.94
55:DA:1083:U:H1'	55:DA:1086:A:N6	1.82	0.94
52:CD:72:C:H2'	52:CD:73:A:H5''	1.49	0.94
56:DI:9:LYS:O	56:DI:12:LEU:N	1.99	0.94
58:DL:18:THR:HB	58:DL:19:PRO:HD3	0.97	0.94
57:DY:75:GLN:HE21	57:DY:76:GLY:N	1.65	0.94
17:A2:80:GLN:HE21	17:A2:80:GLN:HA	0.78	0.94
1:AA:2131:G:H5'	1:AA:2132:U:H5''	1.48	0.94
1:AA:791:C:H4'	1:AA:792:G:OP1	1.67	0.94
8:AK:131:LYS:HB3	8:AK:132:PRO:HA	1.46	0.94
31:BA:1305:G:H22	31:BA:1331:G:H2'	1.33	0.94
34:BG:12:CYS:HA	34:BG:21:LEU:CD2	1.98	0.94
52:CD:20:U:C2'	52:CD:21:A:H5'	1.97	0.94

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1771:C:HO2'	55:DA:1786:A:H8	1.15	0.94
55:DA:1925:C:C6	55:DA:1925:C:H3'	2.02	0.94
55:DA:2448:A:H4'	55:DA:2449:U:OP2	1.64	0.94
58:DL:108:ALA:CA	58:DL:111:LYS:HD3	1.98	0.94
21:DV:189:ALA:HB1	21:DV:190:GLU:HG3	1.50	0.94
21:DV:61:LEU:HD11	21:DV:65:GLN:HB2	1.48	0.94
1:AA:93:C:H5'	1:AA:94:G:OP2	1.67	0.94
4:AE:37:ARG:HG3	4:AE:46:ALA:HB3	1.47	0.94
21:AV:107:THR:N	21:AV:108:PRO:HD2	1.80	0.94
31:BA:1116:C:C2'	31:BA:1117:G:H5''	1.97	0.94
54:CA:1007:C:C2'	54:CA:1008:C:H5''	1.97	0.94
49:CV:41:VAL:HB	49:CV:42:PRO:CA	1.94	0.94
50:CW:82:SER:O	50:CW:86:ARG:HB2	1.66	0.94
5:DF:46:ARG:HH11	5:DF:46:ARG:HG2	1.32	0.94
58:DL:42:ASN:O	58:DL:46:ALA:HB3	1.66	0.94
57:DY:112:LEU:HD11	57:DY:121:ASP:HB2	1.48	0.94
26:D4:70:GLY:O	26:D4:71:ARG:HB2	1.65	0.94
55:DA:1085:A:H2'	55:DA:1086:A:N7	1.82	0.94
55:DA:2469:A:H2	55:DA:2481:G:H21	0.97	0.94
58:DL:73:PRO:HB3	58:DL:77:LEU:CD1	1.97	0.94
30:A8:50:LEU:HG	30:A8:51:ALA:H	1.30	0.94
1:AA:2898:U:H2'	1:AA:2899:G:H8	1.32	0.94
54:CA:1502:A:H2	54:CA:1505:G:H1	1.03	0.94
9:DM:137:LYS:HG3	9:DM:138:LEU:H	1.32	0.94
57:DY:127:GLU:O	57:DY:128:LEU:HD23	1.65	0.94
57:DY:19:ARG:C	57:DY:21:GLN:H	1.71	0.94
57:DY:55:LYS:HD2	57:DY:79:ALA:HA	1.49	0.94
1:AA:1569:A:O2'	3:AD:38:LYS:HE2	1.68	0.93
55:DA:1058:U:H2'	55:DA:1059:G:N7	1.83	0.93
55:DA:1179:C:C2'	55:DA:1180:C:H5''	1.97	0.93
8:DK:38:LEU:H	8:DK:38:LEU:HD12	1.30	0.93
1:AA:2873:A:H8	13:A0:6:SER:H	1.08	0.93
42:CO:18:VAL:HG23	42:CO:19:ARG:H	1.30	0.93
55:DA:1925:C:N4	55:DA:1926:U:N1	2.16	0.93
56:DI:29:GLU:HA	56:DJ:2:ALA:CB	1.98	0.93
21:DV:150:LEU:CD2	21:DV:151:HIS:N	2.30	0.93
1:AA:914:C:H2'	1:AA:915:C:H5'	1.49	0.93
20:AU:94:LYS:HD2	20:AU:101:LYS:HZ3	1.31	0.93
20:AU:76:CYS:SG	20:AU:77:PRO:HD2	2.08	0.93
34:BG:12:CYS:CB	34:BG:21:LEU:HD22	1.98	0.93
28:D6:22:ALA:HB2	28:D6:42:TRP:HZ2	1.33	0.93

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1359:A:H3'	55:DA:1359:A:H8	1.19	0.93
58:DL:126:MET:H	58:DL:126:MET:HE3	1.34	0.93
2:AB:81:G:N7	2:AB:96:G:C2	2.37	0.93
34:CG:114:ARG:HH11	34:CG:114:ARG:HG3	1.31	0.93
16:D1:34:LYS:HA	16:D1:34:LYS:HE2	1.50	0.93
58:DL:107:ILE:HG23	58:DL:108:ALA:N	1.81	0.93
57:DY:93:LEU:HA	57:DY:96:PHE:O	1.67	0.93
1:AA:1019:U:H3	1:AA:1142(A):A:N6	1.64	0.93
54:CA:612:C:O2	54:CA:629:G:N2	2.02	0.93
54:CA:628:G:N2	54:CA:629:G:N3	2.17	0.93
32:CE:122:PHE:HD1	32:CE:139:LYS:HZ1	1.17	0.93
58:DL:89:HIS:O	58:DL:90:LYS:HB2	1.67	0.93
21:DV:190:GLU:C	21:DV:191:VAL:HG23	1.89	0.93
57:DY:75:GLN:NE2	57:DY:76:GLY:H	1.66	0.93
9:AM:91:LEU:HA	9:AM:95:PRO:HB3	1.50	0.93
23:AZ:7:ILE:HG23	23:AZ:95:LEU:HD11	1.50	0.93
5:DF:136:THR:HG22	5:DF:166:ALA:O	1.69	0.93
57:DY:29:TYR:HE2	57:DY:32:LEU:CD1	1.79	0.93
4:AE:8:LYS:HE3	4:AE:188:VAL:HG13	1.46	0.93
54:CA:1449:C:C2'	54:CA:1450:U:H5''	1.98	0.93
37:CJ:113:GLU:HB2	37:CJ:119:ARG:HG2	1.50	0.93
54:CA:1128:C:H5'	39:CL:16:ARG:HH22	1.31	0.93
55:DA:1689:A:H62	55:DA:1698:A:H2	1.02	0.93
58:DL:144:VAL:HG13	58:DL:145:LYS:H	1.34	0.93
15:DR:39:ARG:HG2	15:DR:40:THR:H	1.33	0.93
57:DY:142:LEU:HD13	57:DY:143:GLN:N	1.82	0.93
1:AA:1372:U:C6	1:AA:1372:U:C4'	2.49	0.93
31:BA:1305:G:HO2'	31:BA:1306:A:H8	1.08	0.93
56:DI:23:LEU:C	56:DI:24:ILE:HG22	1.89	0.93
56:DJ:5:ILE:HG22	56:DJ:9:LYS:HB2	1.51	0.93
58:DL:8:VAL:O	58:DL:57:ILE:HG12	1.66	0.93
3:AD:35:LYS:HZ1	3:AD:104:TYR:HB2	1.30	0.93
31:BA:134:A:H61	46:BS:25:ARG:NH1	1.66	0.93
55:DA:1925:C:N4	55:DA:1926:U:C6	2.37	0.93
38:BK:42:GLU:HG3	38:BK:109:ILE:HD12	1.51	0.93
40:BM:79:ARG:H	40:BM:79:ARG:HD3	1.34	0.93
42:BO:47:LYS:HB3	42:BO:48:PRO:HD2	1.51	0.93
49:CV:10:PHE:N	49:CV:10:PHE:CD1	2.32	0.93
50:CW:26:ASN:HB2	50:CW:71:THR:HG23	1.49	0.93
56:DI:23:LEU:O	56:DI:24:ILE:HG22	1.69	0.93
57:DY:23:SER:HB3	57:DY:68:LEU:HB2	1.46	0.93

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:AF:3:GLU:HA	5:AF:24:LEU:HG	1.48	0.92
21:AV:175:VAL:O	21:AV:177:PRO:HD2	1.69	0.92
31:BA:168:G:C2'	31:BA:169:C:H5''	2.00	0.92
37:BJ:113:GLU:HB2	37:BJ:119:ARG:HG2	1.50	0.92
49:BV:41:VAL:H	49:BV:44:MET:HE3	1.33	0.92
43:CP:10:PRO:HB2	43:CP:18:ALA:HB1	1.51	0.92
49:CV:10:PHE:H	49:CV:10:PHE:HD1	1.14	0.92
11:DO:83:VAL:HG11	11:DO:112:LEU:HD21	1.51	0.92
15:DR:24:PRO:HA	15:DR:49:VAL:HG13	1.49	0.92
5:AF:178:PRO:HB2	5:AF:201:VAL:HG11	1.51	0.92
55:DA:1078:U:H1'	55:DA:1088:A:H2	1.33	0.92
55:DA:1434:A:H61	55:DA:1558:A:N6	1.67	0.92
57:DY:130:THR:HG22	56:DJ:14:GLN:HE22	1.32	0.92
31:BA:792:A:C1'	31:BA:794:A:H62	1.83	0.92
54:CA:1348:U:H3	54:CA:1374:A:H2	1.15	0.92
35:CH:40:ARG:HB3	35:CH:40:ARG:NH1	1.84	0.92
57:DY:93:LEU:HD22	57:DY:97:ALA:CB	1.99	0.92
1:AA:1454:U:H4'	1:AA:1455:G:OP1	1.68	0.92
31:BA:409:G:OP1	34:BG:24:GLU:HG3	1.68	0.92
37:CJ:15:ASP:H	37:CJ:20:ASP:H	1.17	0.92
58:DL:7:VAL:HG11	58:DL:57:ILE:C	1.89	0.92
15:AR:132:LYS:HG2	15:AR:136:GLN:HE22	1.35	0.92
54:CA:1003:G:C2'	54:CA:1004:A:H5'	1.98	0.92
38:CK:102:ARG:HH11	38:CK:105:ARG:HH22	0.97	0.92
6:DG:47:LYS:HD3	6:DG:81:LYS:HB2	1.50	0.92
7:DH:153:LYS:HB3	7:DH:154:PRO:CD	1.97	0.92
12:DP:66:ILE:HA	12:DP:104:PHE:HA	1.51	0.92
6:AG:67:LYS:HG3	26:A4:6:HIS:HB3	1.50	0.92
23:AZ:51:VAL:HG11	23:AZ:74:VAL:HG21	1.49	0.92
54:CA:973:G:H1'	40:CM:55:LYS:HE2	1.51	0.92
54:CA:690:G:H22	41:CN:55:LYS:NZ	1.67	0.92
55:DA:1175:U:O2'	55:DA:1176:G:H4'	1.69	0.92
55:DA:1236:G:H4'	55:DA:1237:A:OP1	1.70	0.92
57:DY:62:ALA:O	57:DY:63:LEU:HD23	1.69	0.92
1:AA:1236:G:H4'	1:AA:1237:A:OP1	1.67	0.92
8:AK:79:ILE:CB	8:AK:142:VAL:HG11	2.00	0.92
40:CM:24:VAL:HG22	40:CM:72:VAL:HG11	1.51	0.92
41:CN:127:LYS:HE2	41:CN:127:LYS:HA	1.50	0.92
1:AA:2732:G:H3'	1:AA:2733:A:H5'	1.51	0.92
31:BA:991:U:H3	31:BA:1213:A:H62	1.17	0.92
31:BA:579:G:H5'	31:BA:728:A:H1'	1.50	0.92

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:BG:25:ARG:HB3	34:BG:25:ARG:NH1	1.85	0.92
48:BU:22:VAL:O	48:BU:23:LYS:HG3	1.69	0.92
55:DA:2458:G:H2'	55:DA:2490:G:O6	1.69	0.92
4:DE:14:ILE:HG23	4:DE:15:PHE:N	1.82	0.92
4:DE:50:GLY:HA2	4:DE:77:ILE:HA	1.51	0.92
57:DY:89:ALA:CB	56:DJ:15:ALA:CB	2.34	0.92
8:DK:77:LEU:HD11	8:DK:140:LEU:CB	2.00	0.92
21:DV:189:ALA:HA	21:DV:190:GLU:CG	2.00	0.92
31:BA:406:G:H21	34:BG:119:GLN:HE22	1.12	0.92
31:BA:827:U:H3	31:BA:872:A:N6	1.68	0.92
39:BL:95:LYS:HD3	39:BL:96:LEU:N	1.84	0.92
21:DV:184:ALA:O	21:DV:185:GLU:HB2	1.70	0.92
57:DY:15:GLU:O	57:DY:15:GLU:HG3	1.66	0.92
9:AM:39:ARG:NH2	9:AM:41:ASP:HB2	1.85	0.92
15:AR:24:PRO:HA	15:AR:49:VAL:HG13	1.52	0.92
21:AV:67:LEU:HD23	21:AV:68:PRO:HD2	1.49	0.92
54:CA:820:U:H4'	54:CA:821:G:OP2	1.69	0.92
57:DY:27:VAL:CB	57:DY:111:LEU:H	1.81	0.92
57:DY:130:THR:CG2	56:DJ:14:GLN:NE2	2.30	0.92
57:DY:32:LEU:CB	57:DY:33:PRO:CD	2.48	0.92
1:AA:1056:G:H4'	1:AA:1086:A:H1'	1.52	0.91
31:BA:1003:G:C2'	31:BA:1004:A:H5''	1.99	0.91
44:BQ:12:ARG:HG2	44:BQ:14:PRO:HD3	1.49	0.91
55:DA:1826:G:H4'	3:DD:242:ARG:HH21	1.34	0.91
8:DK:115:ALA:CB	8:DK:128:LEU:HD11	1.99	0.91
57:DY:26:LEU:H	57:DY:82:PHE:HE2	1.14	0.91
1:AA:71:A:H4'	1:AA:72:U:O5'	1.69	0.91
5:AF:155:LEU:HD23	5:AF:186:ILE:HD13	1.50	0.91
20:AU:89:PHE:HD1	20:AU:90:LEU:HD23	1.34	0.91
21:AV:110:GLY:N	21:AV:143:GLY:HA2	1.84	0.91
23:AZ:44:PRO:HG2	23:AZ:46:LEU:HD13	1.51	0.91
53:C1:32:A:H2'	53:C1:33:G:O4'	1.68	0.91
58:DL:132:ARG:CG	58:DL:137:GLU:OE2	2.17	0.91
55:DA:631:A:OP1	11:DO:64:LYS:HE2	1.70	0.91
57:DY:91:LYS:NZ	57:DY:95:GLN:NE2	2.18	0.91
55:DA:2394:C:OP1	11:DO:63:PRO:HD2	1.70	0.91
2:DB:75:G:H5'	2:DB:75:G:H8	1.35	0.91
58:DL:18:THR:CG2	58:DL:38:VAL:HG12	1.98	0.91
11:DO:85:LEU:HA	11:DO:88:LEU:HD22	1.52	0.91
57:DY:32:LEU:HB2	57:DY:33:PRO:HD2	1.50	0.91
17:A2:14:VAL:HB	17:A2:96:ILE:HG13	1.51	0.91

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:532:A:H2	54:CA:1206:G:H21	1.16	0.91
57:DY:127:GLU:CG	57:DY:128:LEU:H	1.81	0.91
23:AZ:82:LEU:HG	23:AZ:83:GLU:H	1.35	0.91
16:D1:92:ARG:O	16:D1:94:ASN:N	2.04	0.91
55:DA:1026:U:O2'	55:DA:1027:A:H5''	1.70	0.91
4:DE:6:GLY:HA3	4:DE:26:ILE:HD11	1.52	0.91
4:DE:7:VAL:HG23	4:DE:8:LYS:H	1.34	0.91
58:DL:107:ILE:CG2	58:DL:108:ALA:N	2.28	0.91
58:DL:19:PRO:CA	58:DL:25:PRO:HG2	1.98	0.91
21:DV:150:LEU:HD21	21:DV:154:ASP:HB2	1.52	0.91
21:DV:191:VAL:HG21	21:DV:197:ILE:HG12	1.51	0.91
57:DY:134:LEU:HA	57:DY:137:GLU:HG2	1.50	0.91
1:AA:848:G:H2'	1:AA:849:A:C8	2.06	0.91
39:BL:26:VAL:HG22	39:BL:61:ALA:HB3	1.52	0.91
54:CA:963:G:H21	40:CM:55:LYS:HD3	1.36	0.91
49:CV:41:VAL:HG13	49:CV:44:MET:HB2	1.52	0.91
57:DY:71:LEU:HD22	57:DY:72:ASP:N	1.86	0.91
26:A4:1:MET:SD	26:A4:1:MET:N	2.42	0.91
28:A6:41:PRO:CG	28:A6:45:LYS:C	2.38	0.91
29:A7:24:THR:HG23	29:A7:27:GLY:H	1.35	0.91
30:A8:40:GLU:HA	30:A8:43:GLN:HB2	1.51	0.91
1:AA:2420:C:H41	30:A8:31:HIS:HB3	1.33	0.91
31:BA:279:A:H4'	31:BA:280:C:C5'	1.99	0.91
42:BO:8:ASN:HD22	47:BT:34:LYS:HE2	1.34	0.91
16:D1:64:ARG:HG2	16:D1:64:ARG:HH21	1.36	0.91
55:DA:1022:G:H22	55:DA:1142(A):A:H2	1.18	0.91
55:DA:860:U:H5	55:DA:917:A:N1	1.68	0.91
4:DE:14:ILE:O	4:DE:15:PHE:CD2	2.22	0.91
8:DK:74:ASN:ND2	8:DK:75:LEU:H	1.69	0.91
1:AA:2755:C:H4'	1:AA:2756:U:H5	1.33	0.91
1:AA:900:A:H3'	1:AA:901:A:H8	1.34	0.91
21:AV:131:ARG:NH1	21:AV:131:ARG:HG2	1.74	0.91
34:BG:100:ARG:HH12	34:BG:137:SER:HB3	1.36	0.91
54:CA:1067:A:HO2'	54:CA:1068:G:H8	1.12	0.91
55:DA:894:C:H2'	55:DA:895:U:C6	2.05	0.91
58:DL:95:LYS:N	58:DL:136:VAL:HG11	1.85	0.91
11:DO:75:ILE:H	11:DO:75:ILE:CD1	1.84	0.91
14:DQ:71:ARG:HG2	14:DQ:104:GLY:HA2	1.48	0.91
15:DR:27:THR:HG23	15:DR:90:GLN:HB3	1.51	0.91
57:DY:13:LEU:CD2	57:DY:62:ALA:HB1	1.98	0.91
1:AA:1924:C:N3	1:AA:1925:C:C6	2.39	0.91

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1267:U:C5	1:AA:2012:G:C2	2.58	0.91
20:AU:81:LYS:HD3	20:AU:97:ARG:NE	1.86	0.91
21:AV:175:VAL:HG13	21:AV:176:PRO:N	1.81	0.91
31:BA:255:G:H1'	47:BT:16:GLN:NE2	1.85	0.91
31:BA:351:G:H4'	31:BA:352:C:OP1	1.70	0.91
35:BH:48:ALA:HB1	35:BH:49:PRO:HD2	1.50	0.91
52:CD:9:A:H62	52:CD:23:A:H62	1.18	0.91
55:DA:1725:G:H8	55:DA:1725:G:H5'	1.35	0.91
55:DA:2469:A:H2	55:DA:2481:G:N2	1.68	0.91
55:DA:2580:U:H4'	4:DE:130:GLY:HA3	1.53	0.91
55:DA:2734:A:H5'	55:DA:2735:G:OP2	1.69	0.91
58:DL:19:PRO:C	58:DL:25:PRO:CD	2.38	0.91
57:DY:25:PHE:CE1	57:DY:82:PHE:HB3	2.01	0.91
9:AM:15:LEU:HG	9:AM:134:ARG:HE	1.33	0.91
10:AN:104:ARG:HB3	10:AN:104:ARG:NH1	1.86	0.91
52:BD:41:C:H2'	52:BD:42:C:H5''	1.50	0.91
40:BM:10:GLY:HA3	40:BM:16:LEU:HD21	1.51	0.91
32:CE:204:ASN:ND2	32:CE:206:ASP:H	1.69	0.91
40:CM:38:ILE:HD11	40:CM:71:LEU:HD23	1.51	0.91
43:CP:120:LYS:HD3	43:CP:120:LYS:N	1.84	0.91
49:CV:15:LEU:H	49:CV:15:LEU:HD23	1.36	0.91
55:DA:1372:U:H6	55:DA:1372:U:H5'	1.12	0.91
58:DL:12:LEU:HB3	58:DL:13:PRO:CA	2.01	0.91
58:DL:8:VAL:C	58:DL:57:ILE:HG13	1.90	0.91
57:DY:75:GLN:CB	57:DY:110:GLY:O	2.18	0.91
57:DY:43:ALA:CB	57:DY:47:ASN:ND2	2.11	0.91
1:AA:1086:A:H4'	1:AA:1103:A:H61	1.35	0.90
1:AA:2898:U:H2'	1:AA:2899:G:C8	2.06	0.90
10:AN:47:ILE:HG13	10:AN:48:PRO:HD2	1.51	0.90
21:AV:183:LEU:HD23	21:AV:183:LEU:H	1.33	0.90
43:BP:80:ARG:O	43:BP:83:ASP:CB	2.17	0.90
55:DA:259:G:H21	55:DA:621:A:H8	1.19	0.90
6:DG:37:VAL:HG22	6:DG:159:VAL:HA	1.53	0.90
56:DJ:12:LEU:HB3	56:DJ:13:SER:HB2	1.43	0.90
58:DL:106:GLU:HG2	58:DL:109:LYS:HB2	1.53	0.90
57:DY:70:GLU:C	57:DY:71:LEU:HD12	1.90	0.90
17:A2:48:GLY:HA3	17:A2:52:VAL:HG22	1.52	0.90
1:AA:925:C:C2'	1:AA:926:A:H5''	1.99	0.90
11:AO:146:VAL:HG22	11:AO:147:LEU:H	1.36	0.90
18:AS:59:VAL:HG23	18:AS:65:LEU:H	1.37	0.90
26:A4:52:THR:HG21	43:BP:65:LYS:HD3	1.54	0.90

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:D3:51:VAL:HG23	22:D3:81:VAL:HG23	1.53	0.90
11:DO:62:LEU:HD21	30:D8:25:MET:HB2	1.52	0.90
55:DA:1033:U:H4'	55:DA:1034:G:OP1	1.70	0.90
55:DA:1061:U:C5	58:DL:54:PRO:HB3	2.07	0.90
55:DA:1803:A:O2'	3:DD:259:THR:HG21	1.71	0.90
6:DG:7:LEU:HD21	6:DG:176:LEU:HD22	1.52	0.90
7:DH:13:LYS:HA	7:DH:13:LYS:HE2	1.51	0.90
57:DY:62:ALA:C	57:DY:63:LEU:HD23	1.91	0.90
1:AA:2580:U:H4'	4:AE:130:GLY:HA2	1.51	0.90
19:AT:18:TYR:HA	19:AT:21:PHE:CD2	2.07	0.90
56:DI:20:LEU:CA	56:DI:24:ILE:HG21	2.02	0.90
21:DV:105:VAL:CG1	21:DV:140:ASP:HB3	2.02	0.90
55:DA:1075:C:H4'	21:DV:195:GLU:HG2	0.92	0.90
1:AA:1281:G:H5'	1:AA:1281:G:H8	1.36	0.90
1:AA:2753:A:C2'	1:AA:2754:U:H5''	2.01	0.90
54:CA:96:G:H2'	54:CA:97:U:H5'	1.53	0.90
30:D8:29:LYS:HB2	30:D8:44:LYS:HG2	1.54	0.90
21:DV:189:ALA:CA	21:DV:190:GLU:CG	2.48	0.90
57:DY:50:ARG:C	57:DY:83:TYR:HA	1.90	0.90
3:AD:43:ARG:HH11	3:AD:44:ASN:HD22	1.17	0.90
12:AP:31:ASP:H	12:AP:107:ALA:HB2	1.35	0.90
31:BA:6:G:H4'	31:BA:298:A:H4'	1.52	0.90
31:BA:939:G:H5''	37:BJ:102:ARG:NH2	1.85	0.90
17:D2:89:GLN:HA	17:D2:89:GLN:HE21	1.37	0.90
28:D6:27:LYS:HB2	28:D6:27:LYS:HZ2	1.35	0.90
55:DA:1056:G:OP1	57:DY:35:LYS:HD3	1.72	0.90
56:DI:21:LYS:O	56:DI:26:ALA:HB2	1.72	0.90
57:DY:138:LEU:HD21	56:DJ:22:GLN:OE1	1.71	0.90
15:DR:102:ILE:HB	15:DR:110:ILE:HD13	1.50	0.90
21:AV:175:VAL:HG13	21:AV:176:PRO:HD2	0.90	0.90
40:BM:48:THR:HA	40:BM:62:HIS:HB3	1.54	0.90
34:CG:187:ARG:NH2	34:CG:190:ASP:HB2	1.86	0.90
8:DK:87:LYS:HA	8:DK:122:GLU:HA	1.52	0.90
8:AK:88:ILE:HG22	8:AK:89:TYR:H	1.36	0.90
31:BA:1053:G:H5'	31:BA:1054:C:H5'	1.51	0.90
34:CG:196:LEU:HD12	34:CG:196:LEU:H	1.36	0.90
58:DL:112:MET:HE1	58:DL:123:ALA:HB3	1.53	0.90
24:DW:42:GLY:O	24:DW:44:LEU:N	2.05	0.90
24:AW:22:GLU:O	24:AW:26:ARG:HG3	1.72	0.90
49:BV:49:ILE:HD12	49:BV:49:ILE:H	1.34	0.90
44:CQ:12:ARG:C	44:CQ:14:PRO:HD2	1.92	0.90

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1024:G:H3'	55:DA:1025:G:H5''	1.54	0.90
6:DG:67:LYS:HG2	26:D4:5:ILE:HG22	1.51	0.90
31:BA:1014:A:H2	31:BA:1219:U:H1'	1.37	0.90
42:BO:60:LEU:HB2	42:BO:64:TYR:HB2	1.54	0.90
52:CC:44:G:H3'	52:CC:45:U:C6	2.07	0.90
43:CP:90:LEU:HD22	49:CV:78:ARG:HH21	1.31	0.90
50:CW:89:ARG:HH21	50:CW:104:LEU:HD21	1.35	0.90
28:D6:41:PRO:HD2	28:D6:46:HIS:N	1.87	0.90
12:DP:65:PHE:O	12:DP:66:ILE:HG12	1.72	0.90
55:DA:297:C:H5''	20:DU:85:VAL:HG21	1.51	0.90
57:DY:87:VAL:O	57:DY:91:LYS:HB2	1.71	0.90
30:A8:49:VAL:CG1	30:A8:50:LEU:H	1.82	0.89
31:BA:954:G:H21	31:BA:1227:A:H62	1.16	0.89
43:BP:10:PRO:CB	43:BP:18:ALA:HB1	2.01	0.89
17:D2:35:LEU:H	17:D2:35:LEU:HD22	1.35	0.89
5:DF:9:ILE:HD11	5:DF:125:LEU:HG	1.52	0.89
6:DG:56:ALA:CB	6:DG:153:ARG:HE	1.84	0.89
58:DL:87:GLY:CA	58:DL:96:VAL:HG21	2.02	0.89
21:DV:73:GLN:HB3	21:DV:87:ASP:OD1	1.71	0.89
57:DY:46:GLN:O	57:DY:47:ASN:HB2	1.70	0.89
1:AA:1332:G:N2	1:AA:1609:A:H2'	1.87	0.89
12:AP:16:ARG:HG3	12:AP:17:LEU:H	1.37	0.89
49:BV:5:LEU:HG	49:BV:9:VAL:HA	1.53	0.89
31:BA:192:U:H4'	50:BW:102:GLY:O	1.73	0.89
54:CA:1124:G:H3'	54:CA:1145:C:N4	1.88	0.89
54:CA:1534:A:C2	54:CA:1535:C:N4	2.39	0.89
43:CP:39:ILE:HD12	43:CP:56:LEU:HD23	1.52	0.89
55:DA:776:G:H4'	55:DA:777:A:O5'	1.72	0.89
3:DD:58:HIS:HD2	3:DD:59:LYS:O	1.55	0.89
3:DD:35:LYS:HG2	3:DD:64:ILE:H	1.36	0.89
56:DJ:15:ALA:C	56:DJ:16:THR:HG23	1.86	0.89
57:DY:16:ASN:HB2	57:DY:19:ARG:HH12	0.73	0.89
1:AA:1698:A:O2'	1:AA:1699:G:H5''	1.71	0.89
12:AP:42:ILE:HD12	12:AP:42:ILE:H	1.36	0.89
20:AU:84:ARG:NH2	20:AU:97:ARG:HB2	1.87	0.89
21:AV:61:LEU:HB3	21:AV:62:PRO:HD2	1.53	0.89
17:D2:38:LEU:HD12	17:D2:56:SER:HA	1.54	0.89
55:DA:897:C:C5	55:DA:897:C:P	2.64	0.89
19:DT:65:ARG:HD3	19:DT:65:ARG:N	1.88	0.89
17:A2:49:THR:HB	17:A2:50:PRO:HD3	1.54	0.89
1:AA:1372:U:H5'	1:AA:1372:U:H6	1.37	0.89

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AB:39:A:H2'	26:A4:1:MET:HE3	1.54	0.89
31:BA:1351:U:H4'	37:BJ:33:ASP:OD2	1.73	0.89
32:CE:80:ILE:HD11	32:CE:208:ILE:HG23	1.54	0.89
26:D4:46:GLN:HG3	26:D4:48:ARG:HG2	1.54	0.89
58:DL:20:ALA:O	58:DL:25:PRO:O	1.90	0.89
18:DS:64:MET:O	18:DS:65:LEU:HB2	1.73	0.89
57:DY:122:VAL:HG12	57:DY:126:ALA:CB	2.01	0.89
57:DY:51:LEU:HD13	57:DY:82:PHE:H	1.07	0.89
15:AR:115:ARG:HD3	15:AR:115:ARG:H	1.35	0.89
36:BI:6:VAL:HG12	36:BI:8:ILE:HD11	1.52	0.89
29:D7:8:ASN:C	29:D7:8:ASN:HD22	1.76	0.89
21:DV:67:LEU:HD23	21:DV:68:PRO:HD2	1.52	0.89
1:AA:2580:U:H4'	4:AE:130:GLY:CA	2.03	0.89
53:B1:36:G:C2'	53:B1:37:G:H5''	2.02	0.89
32:CE:185:ILE:HG22	32:CE:199:TYR:HB2	1.55	0.89
14:DQ:26:LEU:HB3	14:DQ:87:PHE:HA	1.54	0.89
28:A6:41:PRO:HG2	28:A6:45:LYS:C	1.93	0.89
18:AS:110:LYS:HG3	18:AS:111:HIS:ND1	1.87	0.89
31:BA:1244:C:H2'	31:BA:1245:A:C8	2.08	0.89
31:BA:517:G:O2'	31:BA:518:C:OP2	1.91	0.89
49:BV:42:PRO:O	49:BV:44:MET:N	2.06	0.89
55:DA:1482:U:H5'	55:DA:1483:G:OP2	1.73	0.89
55:DA:1964:G:H4'	55:DA:1965:C:OP2	1.73	0.89
1:AA:571:A:H1'	1:AA:573:G:C8	2.07	0.89
1:AA:90:U:O2'	1:AA:91:A:H5''	1.72	0.89
1:AA:946:G:H2'	1:AA:947:G:H8	1.37	0.89
32:BE:44:LEU:HD12	32:BE:45:GLN:H	1.34	0.89
31:BA:1106:G:H5''	33:BF:172:ARG:HG2	1.55	0.89
49:BV:42:PRO:C	49:BV:45:VAL:HG22	1.94	0.89
54:CA:982:U:H4'	54:CA:983:A:O5'	1.71	0.89
48:CU:18:ARG:N	48:CU:18:ARG:CD	2.30	0.89
28:D6:22:ALA:HB2	28:D6:42:TRP:CZ2	2.08	0.89
28:D6:9:LEU:HD13	28:D6:11:LEU:HD21	1.55	0.89
55:DA:1249:U:H2'	55:DA:1249:U:O2	1.70	0.89
58:DL:101:TRP:HD1	58:DL:101:TRP:H	1.17	0.89
57:DY:129:PRO:HD2	57:DY:130:THR:H	1.34	0.89
57:DY:26:LEU:HA	57:DY:112:LEU:CA	2.00	0.89
57:DY:25:PHE:CG	57:DY:82:PHE:CE1	2.60	0.89
1:AA:479:A:O2'	1:AA:481:G:H5''	1.72	0.89
3:AD:43:ARG:NH1	3:AD:44:ASN:ND2	2.20	0.89
52:BB:7:A:H5'	52:BB:8:U:OP2	1.71	0.89

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:BM:4:ILE:HD13	40:BM:82:ILE:HD11	1.55	0.89
40:BM:96:ILE:H	40:BM:96:ILE:HD13	1.38	0.89
42:BO:64:TYR:O	42:BO:65:GLU:HB2	1.72	0.89
55:DA:1906:G:C5	55:DA:1929:G:N2	2.41	0.89
56:DJ:13:SER:HB3	56:DJ:17:VAL:CB	2.02	0.89
14:DQ:106:ARG:HA	14:DQ:110:LEU:HD21	1.54	0.89
57:DY:23:SER:HB2	57:DY:68:LEU:HB2	1.53	0.89
57:DY:26:LEU:N	57:DY:82:PHE:CE2	2.40	0.89
16:A1:92:ARG:HG3	16:A1:94:ASN:HB3	1.53	0.89
37:BJ:97:GLN:HE21	37:BJ:101:LEU:HD11	1.38	0.89
34:CG:47:ARG:NH2	53:C1:57:U:H5	1.70	0.89
22:D3:11:ARG:HB2	22:D3:11:ARG:NH1	1.88	0.89
27:D5:56:LYS:H	27:D5:56:LYS:HD2	1.36	0.89
55:DA:1884:A:H2'	55:DA:1885:A:H5''	1.54	0.89
57:DY:90:ALA:H	56:DJ:15:ALA:CB	1.86	0.89
58:DL:59:ILE:HG22	58:DL:60:TYR:N	1.86	0.89
58:DL:83:GLY:H	58:DL:99:ILE:HG23	1.37	0.89
57:DY:123:GLU:O	57:DY:127:GLU:HB3	1.72	0.89
1:AA:1372:U:C5'	1:AA:1372:U:H6	1.86	0.88
1:AA:2820:A:C5	4:AE:191:PRO:CB	2.56	0.88
1:AA:896:A:C2	21:AV:178:GLU:HG2	2.08	0.88
4:AE:36:ARG:NH2	4:AE:88:GLY:HA3	1.86	0.88
9:AM:15:LEU:HB2	9:AM:134:ARG:HG2	1.53	0.88
31:BA:191(C):G:H3'	31:BA:191(D):U:H5''	1.55	0.88
35:CH:148:VAL:HG21	38:CK:107:LEU:HD22	1.54	0.88
7:DH:89:ILE:CD1	7:DH:129:THR:HB	2.02	0.88
14:DQ:83:LYS:C	14:DQ:109:GLY:HA3	1.93	0.88
24:DW:47:ASN:O	24:DW:49:LYS:N	2.06	0.88
57:DY:141:VAL:HG22	57:DY:142:LEU:N	1.87	0.88
2:AB:74:U:C2'	2:AB:75:G:H5''	2.03	0.88
20:AU:20:TYR:C	20:AU:22:GLY:H	1.76	0.88
21:AV:120:ILE:HG21	21:AV:170:THR:HB	1.55	0.88
43:BP:22:ILE:HB	43:BP:25:ILE:CG1	2.01	0.88
6:DG:16:ARG:HG2	6:DG:16:ARG:HH11	1.37	0.88
58:DL:83:GLY:O	58:DL:97:GLY:HA3	1.73	0.88
57:DY:25:PHE:CB	57:DY:82:PHE:CE1	2.57	0.88
57:DY:25:PHE:HD1	57:DY:82:PHE:CD2	1.84	0.88
1:AA:1799:G:H4'	1:AA:1800:C:O5'	1.74	0.88
1:AA:2296:U:H4'	1:AA:2297:C:OP1	1.73	0.88
1:AA:611:C:H2'	1:AA:612:G:C5'	2.02	0.88
4:AE:8:LYS:O	4:AE:9:VAL:HG22	1.72	0.88

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:AV:144:LEU:HD12	21:AV:146:ILE:O	1.73	0.88
39:BL:24:GLY:HA2	39:BL:59:PHE:O	1.73	0.88
54:CA:1446:A:H1'	15:DR:125:ARG:HH22	1.36	0.88
55:DA:1103:A:H2'	55:DA:1104:C:H5'	1.56	0.88
55:DA:2287:A:H2	55:DA:2346:A:N1	1.71	0.88
55:DA:860:U:H5	55:DA:917:A:C2	1.91	0.88
7:DH:92:ILE:HD12	7:DH:92:ILE:H	1.38	0.88
9:DM:67:LEU:O	9:DM:88:GLU:HG3	1.72	0.88
57:DY:144:ALA:CB	57:DY:145:PRO:HD2	1.97	0.88
57:DY:25:PHE:CG	57:DY:82:PHE:CD1	2.61	0.88
1:AA:1332:G:H22	1:AA:1609:A:H2'	1.38	0.88
4:AE:201:THR:HG22	4:AE:202:LYS:H	1.38	0.88
4:AE:58:ARG:CZ	4:AE:58:ARG:HA	2.03	0.88
5:AF:192:LEU:HD23	5:AF:193:VAL:N	1.87	0.88
32:CE:8:LYS:N	32:CE:8:LYS:HD3	1.89	0.88
33:CF:58:GLU:HB2	33:CF:65:ALA:HB3	1.56	0.88
41:CN:21:ILE:HB	41:CN:84:VAL:HG12	1.55	0.88
16:D1:90:VAL:O	16:D1:92:ARG:N	2.05	0.88
58:DL:36:GLU:O	58:DL:39:LYS:HB2	1.72	0.88
20:DU:63:LYS:HZ2	20:DU:64:GLU:H	0.89	0.88
1:AA:1069:A:H4'	1:AA:1070:A:H5''	1.53	0.88
1:AA:2051:A:H61	1:AA:2614:A:H2'	1.37	0.88
1:AA:2665:A:O2'	1:AA:2666:C:H5'	1.72	0.88
1:AA:458:G:O2'	1:AA:459:U:P	2.31	0.88
20:AU:84:ARG:HH21	20:AU:97:ARG:HB2	1.36	0.88
31:BA:792:A:H2'	31:BA:794:A:H62	1.32	0.88
52:BB:74:C:O2'	52:BB:75:C:C5'	2.20	0.88
52:BD:23:A:H2'	52:BD:24:G:C8	2.09	0.88
55:DA:2111:C:H41	55:DA:2147:G:N2	1.69	0.88
55:DA:670:A:H4'	55:DA:671:C:O5'	1.71	0.88
12:DP:2:LEU:HB3	12:DP:70:PRO:HG2	1.56	0.88
30:A8:49:VAL:CG1	30:A8:50:LEU:N	2.30	0.88
4:AE:52:LEU:HD12	4:AE:76:ARG:HB2	1.56	0.88
11:AO:85:LEU:HA	11:AO:88:LEU:HB3	1.53	0.88
54:CA:1321:C:C5'	54:CA:1322:C:H5''	2.03	0.88
35:CH:50:GLU:HG3	35:CH:52:PRO:HD2	1.56	0.88
55:DA:2475:C:H42	55:DA:2529:G:H1	1.21	0.88
57:DY:27:VAL:HG23	57:DY:110:GLY:CA	1.98	0.88
57:DY:13:LEU:HD22	57:DY:13:LEU:O	1.72	0.88
28:D6:12:GLU:HB3	28:D6:23:THR:HG22	1.54	0.88
3:DD:25:THR:CG2	3:DD:82:ILE:H	1.87	0.88

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DB:42:C:O3'	6:DG:67:LYS:HE3	1.74	0.88
1:AA:266:G:C2'	1:AA:267:C:H5''	2.02	0.88
31:BA:498:A:O2'	31:BA:500:G:C8	2.25	0.88
32:CE:69:LEU:HB3	32:CE:162:ILE:HG22	1.56	0.88
43:CP:57:ARG:HB2	43:CP:57:ARG:HH11	1.37	0.88
7:DH:37:VAL:HG12	7:DH:38:SER:H	1.38	0.88
8:DK:131:LYS:HB3	8:DK:132:PRO:CA	2.03	0.88
58:DL:18:THR:HG23	58:DL:42:ASN:HD21	1.37	0.88
9:DM:89:LYS:O	9:DM:93:THR:HG22	1.74	0.88
57:DY:19:ARG:NH2	57:DY:84:GLU:CD	2.26	0.88
1:AA:1209:G:H21	1:AA:1210:A:H62	1.22	0.88
1:AA:271(B):G:H4'	1:AA:271(C):U:O5'	1.74	0.88
1:AA:620:G:H5'	1:AA:621:A:OP1	1.74	0.88
3:AD:30:GLU:HG3	3:AD:63:ARG:NE	1.88	0.88
9:AM:62:VAL:HG22	9:AM:66:LYS:HG3	1.55	0.88
34:BG:16:GLY:HA2	34:BG:33:MET:HE1	1.56	0.88
54:CA:630:G:C8	54:CA:630:G:C3'	2.57	0.88
42:CO:47:LYS:C	42:CO:49:ASN:H	1.75	0.88
55:DA:1291:C:H5'	55:DA:1536:A:H5'	1.54	0.88
55:DA:2832:U:H4'	55:DA:2833:G:H5''	1.53	0.88
2:DB:56:G:H5'	6:DG:27:ASN:ND2	1.89	0.88
56:DJ:13:SER:C	56:DJ:17:VAL:HG21	1.93	0.88
58:DL:104:VAL:O	58:DL:107:ILE:CB	2.22	0.88
58:DL:104:VAL:O	58:DL:107:ILE:HG21	1.72	0.88
21:DV:128:VAL:CA	21:DV:161:VAL:HG21	2.02	0.88
57:DY:72:ASP:C	57:DY:74:LEU:N	2.25	0.88
23:DZ:56:GLN:HE21	23:DZ:56:GLN:N	1.72	0.88
8:AK:129:THR:HG22	8:AK:137:PRO:HB3	1.54	0.88
21:AV:144:LEU:O	21:AV:144:LEU:HD12	1.74	0.88
31:BA:1067:A:H1'	31:BA:1068:G:O4'	1.74	0.88
28:D6:15:GLU:HG2	28:D6:16:CYS:N	1.89	0.88
55:DA:1653:G:O6	13:D0:9:LYS:O	1.91	0.88
58:DL:104:VAL:HG12	58:DL:105:LEU:N	1.88	0.88
55:DA:483:A:H4'	20:DU:49:VAL:CA	2.03	0.88
57:DY:50:ARG:O	57:DY:51:LEU:CG	2.22	0.88
6:AG:115:ARG:HB3	43:BP:7:VAL:HG11	1.55	0.87
54:CA:1278:U:H5''	54:CA:1279:A:O4'	1.74	0.87
16:D1:92:ARG:HD2	17:D2:11:GLN:NE2	1.89	0.87
11:DO:61:ARG:HH11	30:D8:14:VAL:HG23	1.37	0.87
3:DD:44:ASN:HB2	3:DD:49:ILE:HA	1.56	0.87
1:AA:1887:C:C2'	1:AA:1888:G:H5''	2.02	0.87

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:BI:23:LYS:O	36:BI:27:GLN:HG3	1.75	0.87
40:BM:34:VAL:HG22	40:BM:74:ILE:HG22	1.55	0.87
55:DA:1379:A:O2'	55:DA:1380:G:P	2.31	0.87
4:DE:7:VAL:HG23	4:DE:8:LYS:N	1.88	0.87
9:DM:65:LYS:HB2	9:DM:69:GLN:HE21	1.38	0.87
21:DV:60:GLU:HG3	21:DV:61:LEU:H	1.38	0.87
52:CB:19:G:N2	52:CB:56:C:H42	1.72	0.87
41:CN:79:SER:HB2	41:CN:106:LYS:HD2	1.56	0.87
55:DA:1043:C:H2'	55:DA:1044:G:H5''	1.53	0.87
55:DA:1062:G:H2'	55:DA:1063:G:C8	2.09	0.87
58:DL:100:THR:C	58:DL:102:GLU:H	1.77	0.87
20:DU:97:ARG:H	20:DU:97:ARG:HD3	1.36	0.87
1:AA:1964:G:H4'	1:AA:1965:C:OP2	1.70	0.87
1:AA:322:A:H5''	5:AF:169:ASN:HD22	1.37	0.87
31:BA:1256:A:H5'	31:BA:1257:U:OP1	1.74	0.87
6:AG:115:ARG:HH12	43:BP:7:VAL:HG21	1.37	0.87
55:DA:2298:A:H62	55:DA:2318:G:H8	1.15	0.87
56:DI:16:THR:HG23	56:DI:17:VAL:N	1.87	0.87
58:DL:77:LEU:O	58:DL:107:ILE:HD11	1.74	0.87
57:DY:134:LEU:CD2	56:DJ:19:GLU:OE1	2.22	0.87
57:DY:19:ARG:C	57:DY:21:GLN:N	2.24	0.87
1:AA:2468:G:H5'	12:AP:120:ILE:HD11	1.53	0.87
54:CA:1189:C:H5''	33:CF:5:ILE:HG21	1.53	0.87
55:DA:1080:A:H1'	58:DL:126:MET:HA	1.56	0.87
55:DA:608:A:C4	55:DA:621:A:N6	2.42	0.87
2:DB:12:C:H4'	2:DB:13:A:OP1	1.73	0.87
11:DO:61:ARG:O	11:DO:62:LEU:HD22	1.73	0.87
57:DY:24:PHE:CE1	57:DY:88:ALA:HB2	2.09	0.87
1:AA:848:G:H2'	1:AA:849:A:H8	1.36	0.87
31:BA:187:C:H2'	31:BA:188:U:O4'	1.73	0.87
48:BU:84:LYS:HE2	48:BU:84:LYS:HA	1.57	0.87
49:CV:39:THR:HG22	49:CV:40:ILE:H	1.40	0.87
55:DA:1019:U:H3	55:DA:1142(A):A:N6	1.71	0.87
55:DA:1060:U:H4'	55:DA:1061:U:O5'	1.74	0.87
55:DA:1359:A:C3'	55:DA:1359:A:H8	1.76	0.87
55:DA:1405:U:H2'	55:DA:1406:U:H6	1.38	0.87
55:DA:2414:G:H21	11:DO:67:MET:HE1	1.39	0.87
58:DL:135:GLY:O	58:DL:136:VAL:CG1	2.23	0.87
10:DN:68:GLU:HB3	10:DN:78:ARG:NH1	1.90	0.87
19:DT:12:VAL:HG12	19:DT:27:THR:O	1.75	0.87
1:AA:289:A:H5'	1:AA:290:G:OP2	1.74	0.87

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:BD:15:G:N2	52:BD:48:C:H42	1.73	0.87
31:BA:974:A:H1'	44:BQ:31:ARG:HE	1.40	0.87
34:CG:153:ARG:HD3	34:CG:181:MET:SD	2.15	0.87
38:CK:6:ILE:HB	38:CK:85:ARG:NH1	1.89	0.87
50:CW:71:THR:HG22	50:CW:72:LEU:N	1.88	0.87
13:D0:33:ARG:HD3	13:D0:113:LEU:HD11	1.53	0.87
55:DA:1019:U:O2'	55:DA:1021:A:H2	1.57	0.87
56:DI:25:ASP:O	56:DI:28:LYS:HB2	1.74	0.87
56:DJ:10:GLU:C	56:DJ:17:VAL:HG12	1.94	0.87
6:AG:82:LEU:HA	6:AG:86:MET:SD	2.15	0.87
21:AV:127:LYS:O	21:AV:162:GLU:HB2	1.74	0.87
33:BF:32:LEU:HB3	33:BF:59:ARG:HH12	1.40	0.87
55:DA:49:A:N7	55:DA:120:U:C5	2.41	0.87
3:DD:35:LYS:HE3	3:DD:64:ILE:C	1.95	0.87
56:DI:24:ILE:CG1	56:DI:25:ASP:N	2.30	0.87
2:AB:81:G:N2	2:AB:82:G:N7	2.23	0.87
1:AA:662:G:H5'	11:AO:15:ARG:HA	1.57	0.87
25:AX:6:VAL:HG12	25:AX:54:VAL:HG11	1.57	0.87
31:BA:1280:A:H5'	31:BA:1281:U:OP2	1.74	0.87
33:BF:35:GLU:HA	33:BF:38:ARG:NE	1.90	0.87
54:CA:1067:A:O2'	54:CA:1068:G:H8	1.58	0.87
52:CC:35:A:H2'	52:CC:36:A:H5''	1.57	0.87
50:CW:56:MET:HG2	50:CW:84:LEU:HD11	1.57	0.87
55:DA:1138:G:H21	9:DM:106:MET:HE3	1.40	0.87
55:DA:2317:C:H2'	55:DA:2318:G:H5'	1.54	0.87
55:DA:905:U:C2'	55:DA:906:G:H5''	2.05	0.87
21:DV:192:ALA:C	21:DV:194:PRO:HD3	1.94	0.87
57:DY:101:PRO:HG2	57:DY:102:LYS:H	1.37	0.87
22:A3:31:VAL:HB	22:A3:35:ASN:HD22	1.38	0.86
28:A6:25:LYS:HZ2	28:A6:27:LYS:HD2	1.38	0.86
1:AA:1464:C:HO2'	1:AA:1528:A:H8	0.93	0.86
1:AA:654(R):C:H2'	1:AA:654(S):G:H8	1.39	0.86
4:AE:8:LYS:HG2	4:AE:192:ASN:HD22	1.38	0.86
6:AG:64:THR:HG23	6:AG:66:GLN:H	1.39	0.86
31:BA:673:G:H2'	31:BA:674:G:C8	2.10	0.86
55:DA:1187:G:H5''	17:D2:81:TYR:CE2	2.10	0.86
55:DA:2729:G:H1'	4:DE:187:ALA:HB2	1.56	0.86
7:DH:98:LEU:HB2	7:DH:125:VAL:HG11	1.57	0.86
1:AA:2571:C:H5'	1:AA:2572:A:H5''	1.55	0.86
1:AA:1826:G:H4'	3:AD:242:ARG:HH21	1.39	0.86
3:AD:30:GLU:HG3	3:AD:63:ARG:CZ	2.05	0.86

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:BN:29:ILE:HB	41:BN:44:SER:HB3	1.55	0.86
54:CA:250:A:H4'	54:CA:251:G:O5'	1.76	0.86
54:CA:93:U:H2'	54:CA:95:G:H5''	1.56	0.86
38:CK:102:ARG:HH11	38:CK:105:ARG:NH2	1.73	0.86
22:D3:32:ARG:H	22:D3:35:ASN:HD21	0.91	0.86
3:DD:27:THR:HG21	3:DD:83:GLU:HB3	1.57	0.86
56:DJ:14:GLN:CA	56:DJ:15:ALA:C	2.42	0.86
55:DA:483:A:C4'	20:DU:49:VAL:HA	2.04	0.86
21:DV:60:GLU:O	21:DV:61:LEU:HD12	1.75	0.86
57:DY:142:LEU:HD22	57:DY:143:GLN:N	1.89	0.86
57:DY:50:ARG:N	57:DY:83:TYR:HD1	1.73	0.86
1:AA:1784:A:H4'	1:AA:1785:A:O5'	1.74	0.86
5:AF:183:VAL:O	5:AF:187:VAL:HG23	1.75	0.86
40:BM:49:VAL:HG13	44:BQ:41:ARG:HB2	1.58	0.86
26:D4:16:CYS:C	26:D4:18:CYS:H	1.78	0.86
2:DB:20:C:C2'	2:DB:21:G:H5''	2.05	0.86
3:DD:35:LYS:HD3	3:DD:63:ARG:CB	2.05	0.86
8:DK:4:ILE:HG12	8:DK:18:VAL:HG22	1.55	0.86
58:DL:20:ALA:H	58:DL:25:PRO:CB	1.88	0.86
1:AA:468:G:N7	29:A7:39:ARG:NH2	2.24	0.86
1:AA:2319:G:H4'	1:AA:2320:A:O5'	1.76	0.86
1:AA:2571:C:H5'	1:AA:2572:A:C5'	2.05	0.86
8:AK:117:GLU:CD	8:AK:117:GLU:H	1.75	0.86
31:BA:1004:A:O5'	31:BA:1025:U:O4	1.93	0.86
33:CF:181:ASN:ND2	33:CF:204:LEU:HD12	1.89	0.86
38:CK:12:ARG:HH12	38:CK:27:PRO:HD3	1.40	0.86
55:DA:1049:C:H2'	55:DA:1050:A:H5''	1.57	0.86
55:DA:1359:A:H2'	55:DA:1360:A:H5'	1.57	0.86
55:DA:1820:U:H4'	55:DA:1821:A:OP2	1.74	0.86
55:DA:704:G:H2'	55:DA:726:G:H22	1.40	0.86
58:DL:59:ILE:C	58:DL:60:TYR:HD1	1.78	0.86
58:DL:52:ILE:HG13	58:DL:76:TYR:CB	2.05	0.86
24:DW:47:ASN:HD22	24:DW:47:ASN:H	1.24	0.86
2:AB:7:G:H3'	2:AB:8:U:H5''	1.56	0.86
3:AD:70:TRP:CZ3	3:AD:150:LYS:HA	2.10	0.86
10:AN:10:VAL:HG21	10:AN:16:ALA:O	1.76	0.86
28:D6:38:LYS:O	28:D6:38:LYS:HG3	1.75	0.86
55:DA:1177:A:H4'	55:DA:1178:C:H5''	1.55	0.86
2:DB:81:G:N2	2:DB:82:G:N7	2.24	0.86
20:DU:63:LYS:NZ	20:DU:64:GLU:H	1.73	0.86
1:AA:1935:G:H3'	1:AA:1962:C:H42	1.39	0.86

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:654(R):C:H2'	1:AA:654(S):G:C8	2.11	0.86
6:DG:56:ALA:HB2	6:DG:153:ARG:NE	1.91	0.86
21:DV:117:LEU:CD1	21:DV:118:GLN:H	1.85	0.86
1:AA:2602:A:N6	52:BB:76:A:H5'	1.90	0.86
1:AA:310:A:OP1	20:AU:18:GLY:HA2	1.76	0.86
1:AA:612:G:H5'	1:AA:612:G:H8	1.39	0.86
31:BA:1399:C:H4'	31:BA:1400:C:O5'	1.72	0.86
52:BD:41:C:C2'	52:BD:42:C:H5''	2.05	0.86
55:DA:1348:G:H2'	55:DA:1349:A:H5''	1.57	0.86
55:DA:1507:A:H3'	55:DA:1508:A:H5''	1.56	0.86
55:DA:1652:A:O2'	55:DA:1653:G:H5'	1.76	0.86
58:DL:18:THR:HG21	58:DL:38:VAL:CG1	2.05	0.86
58:DL:87:GLY:HA2	58:DL:96:VAL:CG2	2.05	0.86
15:DR:102:ILE:HB	15:DR:110:ILE:CD1	2.05	0.86
25:DX:43:ILE:O	25:DX:47:VAL:HG23	1.74	0.86
4:AE:87:GLU:O	4:AE:87:GLU:HG3	1.75	0.86
21:AV:175:VAL:O	21:AV:177:PRO:CG	2.23	0.86
31:BA:197:A:H1'	31:BA:198:G:O4'	1.75	0.86
55:DA:1173:G:H5''	55:DA:1174:A:OP1	1.74	0.86
4:DE:52:LEU:H	4:DE:52:LEU:HD12	1.36	0.86
56:DJ:13:SER:CB	56:DJ:17:VAL:CG1	2.44	0.86
56:DJ:1:MET:SD	56:DJ:2:ALA:HB3	2.16	0.86
58:DL:136:VAL:O	58:DL:137:GLU:HB2	1.72	0.86
58:DL:52:ILE:HG21	58:DL:75:SER:HB2	0.87	0.86
6:AG:161:THR:HG22	6:AG:163:ALA:H	1.39	0.86
31:BA:1210:C:H4'	31:BA:1214:C:C4	2.10	0.86
32:CE:200:ILE:H	32:CE:200:ILE:HD12	1.41	0.86
32:CE:5:ILE:HG13	32:CE:221:LEU:HD23	1.55	0.86
3:DD:44:ASN:CB	3:DD:49:ILE:HA	2.06	0.86
12:DP:80:GLU:HA	22:D3:4:LYS:NZ	1.89	0.86
57:DY:19:ARG:CZ	57:DY:84:GLU:CD	2.43	0.86
1:AA:2355:C:H5'	22:A3:36:ILE:HD11	1.56	0.86
1:AA:571:A:HO2'	1:AA:573:G:H8	0.86	0.86
21:AV:110:GLY:H	21:AV:143:GLY:CA	1.88	0.86
54:CA:1305:G:N2	54:CA:1331:G:H2'	1.90	0.86
58:DL:112:MET:SD	58:DL:120:LEU:HD13	2.15	0.86
20:DU:81:LYS:HD3	20:DU:97:ARG:NE	1.91	0.86
57:DY:23:SER:CB	57:DY:68:LEU:CB	2.53	0.86
26:A4:38:LYS:O	26:A4:40:HIS:N	2.08	0.85
55:DA:1533:C:H5'	55:DA:1534:G:OP2	1.76	0.85
2:DB:42:C:H4'	6:DG:67:LYS:HD3	1.57	0.85

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:DL:95:LYS:HD3	58:DL:136:VAL:HG21	1.56	0.85
57:DY:40:LEU:HD23	57:DY:50:ARG:NH1	1.90	0.85
57:DY:75:GLN:HB3	57:DY:111:LEU:HA	1.57	0.85
1:AA:260:G:H1'	1:AA:621:A:H8	1.40	0.85
1:AA:752:A:O2'	1:AA:753:C:OP2	1.93	0.85
7:AH:102:ALA:HA	7:AH:117:PRO:HD3	1.58	0.85
31:BA:511:C:O4'	34:BG:43:HIS:NE2	2.09	0.85
54:CA:96:G:H2'	54:CA:97:U:C5'	2.06	0.85
42:CO:60:LEU:HD23	42:CO:60:LEU:H	1.40	0.85
49:CV:88:LYS:HE2	49:CV:88:LYS:CA	2.06	0.85
55:DA:2810:A:O2'	4:DE:61:ARG:HG3	1.76	0.85
4:DE:16:ARG:HG3	4:DE:16:ARG:O	1.76	0.85
55:DA:2304:G:H21	6:DG:156:ASP:CG	1.78	0.85
58:DL:53:VAL:CB	58:DL:72:PRO:HB2	2.06	0.85
11:DO:135:LEU:HD12	11:DO:139:LYS:HD3	1.58	0.85
20:DU:63:LYS:HZ2	20:DU:64:GLU:N	1.73	0.85
57:DY:71:LEU:HB3	57:DY:113:GLN:CG	2.06	0.85
57:DY:50:ARG:HD3	57:DY:51:LEU:O	1.75	0.85
1:AA:444:C:OP2	16:A1:2:PRO:HD3	1.76	0.85
1:AA:2820:A:H61	4:AE:192:ASN:HB2	1.41	0.85
1:AA:800:A:H4'	1:AA:801:G:O5'	1.75	0.85
50:BW:26:ASN:HB2	50:BW:71:THR:HG23	1.58	0.85
52:CD:16:U:H1'	52:CD:60:U:O2	1.75	0.85
13:D0:63:ARG:O	13:D0:67:LEU:HB2	1.75	0.85
16:D1:105:VAL:HA	17:D2:44:LYS:HD3	1.59	0.85
4:DE:201:THR:HG22	4:DE:203:LYS:H	1.39	0.85
5:DF:198:ALA:HA	5:DF:201:VAL:HG12	1.56	0.85
56:DI:20:LEU:C	56:DI:24:ILE:HG21	1.94	0.85
56:DJ:22:GLN:HG3	56:DJ:25:ASP:HB3	1.58	0.85
58:DL:101:TRP:CA	58:DL:104:VAL:HB	2.06	0.85
21:DV:152:ALA:O	21:DV:154:ASP:N	2.10	0.85
31:BA:1129:C:H4'	31:BA:1130:A:C5'	2.04	0.85
31:BA:1537:U:H2'	31:BA:1538:C:C6	2.10	0.85
3:DD:27:THR:HG23	3:DD:28:GLU:N	1.92	0.85
7:DH:152:ARG:O	7:DH:153:LYS:HB2	1.75	0.85
58:DL:7:VAL:HG13	58:DL:58:THR:C	1.96	0.85
20:DU:39:VAL:HG12	20:DU:40:GLU:H	1.41	0.85
57:DY:107:VAL:HG12	57:DY:108:LYS:H	1.40	0.85
57:DY:93:LEU:HD21	57:DY:126:ALA:CB	2.06	0.85
1:AA:1251:C:O2'	1:AA:1252:G:H3'	1.76	0.85
1:AA:2656:U:H5	1:AA:2664:G:H21	1.21	0.85

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:AN:88:ASN:HD21	10:AN:90:GLN:HB2	1.41	0.85
54:CA:1542:U:O3'	54:CA:1542:U:OP2	1.92	0.85
38:CK:103:VAL:CG2	38:CK:110:ALA:HB2	2.06	0.85
55:DA:528:A:C2	55:DA:2042:A:H2'	2.12	0.85
55:DA:2795:G:H3'	55:DA:2797:U:C5'	2.07	0.85
55:DA:890:A:H3'	55:DA:892:G:C8	2.12	0.85
6:DG:67:LYS:HG2	26:D4:5:ILE:CG2	2.06	0.85
58:DL:51:ALA:O	58:DL:52:ILE:HG12	1.77	0.85
57:DY:16:ASN:ND2	57:DY:25:PHE:HZ	1.74	0.85
1:AA:2506:U:H4'	1:AA:2507:C:OP1	1.76	0.85
10:AN:35:VAL:HG11	10:AN:103:ALA:HB3	1.56	0.85
31:BA:1004:A:O4'	31:BA:1036:G:O6	1.94	0.85
43:BP:4:ILE:HG23	43:BP:5:ALA:H	1.40	0.85
43:CP:108:ARG:HD2	43:CP:108:ARG:N	1.91	0.85
55:DA:1092:C:H2'	55:DA:1093:G:C4'	2.07	0.85
55:DA:1286:A:N1	55:DA:1329:U:H2'	1.92	0.85
55:DA:768:G:O2'	55:DA:1379:A:N6	2.09	0.85
3:DD:273:ARG:O	3:DD:273:ARG:HG3	1.76	0.85
4:DE:57:LYS:HE3	4:DE:59:VAL:HB	1.56	0.85
14:DQ:107:GLU:H	14:DQ:110:LEU:HD11	1.41	0.85
15:DR:51:ARG:HG3	15:DR:98:LYS:HG3	1.55	0.85
1:AA:1111:A:H4'	7:AH:3:ARG:HD3	1.57	0.85
4:AE:12:THR:O	4:AE:23:VAL:HG22	1.75	0.85
12:AP:23:GLY:HA2	21:AV:78:LYS:HE3	1.59	0.85
31:BA:718:G:H5'	41:BN:117:ASN:OD1	1.77	0.85
13:D0:97:VAL:HG22	13:D0:114:VAL:HG22	1.59	0.85
55:DA:1142(A):A:H4'	55:DA:1143:A:OP1	1.75	0.85
55:DA:1405:U:H2'	55:DA:1406:U:C6	2.10	0.85
2:DB:56:G:H5'	6:DG:27:ASN:HD21	1.39	0.85
7:DH:126:PRO:HD2	7:DH:127:GLU:N	1.90	0.85
58:DL:125:ARG:O	58:DL:128:ALA:N	2.09	0.85
21:DV:181:GLU:HG2	21:DV:181:GLU:O	1.76	0.85
57:DY:54:ALA:HB1	57:DY:57:THR:CB	2.06	0.85
22:A3:32:ARG:N	22:A3:35:ASN:ND2	2.24	0.85
1:AA:1083:U:H2'	1:AA:1085:A:OP2	1.77	0.85
5:AF:161:GLU:HG2	5:AF:164:ARG:NH2	1.91	0.85
31:BA:366:C:H4'	31:BA:367:U:OP1	1.76	0.85
52:BB:19:G:H1'	52:BB:57:G:N2	1.92	0.85
38:CK:42:GLU:HG3	38:CK:109:ILE:HD12	1.57	0.85
48:CU:29:PHE:HD2	48:CU:29:PHE:H	1.22	0.85
28:D6:20:ASN:ND2	28:D6:21:TYR:H	1.74	0.85

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:DD:30:GLU:HG3	3:DD:63:ARG:CZ	2.06	0.85
58:DL:106:GLU:O	58:DL:109:LYS:HB3	1.75	0.85
21:DV:61:LEU:HD12	21:DV:65:GLN:HB2	1.56	0.85
57:DY:27:VAL:O	57:DY:28:ASN:HB2	1.75	0.85
1:AA:997:G:OP1	16:A1:93:LYS:HD3	1.77	0.85
1:AA:142:G:H1'	19:AT:37:THR:HG21	1.58	0.85
21:AV:116:VAL:HG12	21:AV:117:LEU:H	1.42	0.85
21:AV:132:ASN:C	21:AV:133:ILE:HD12	1.97	0.85
23:AZ:92:LYS:HZ1	23:AZ:97:LEU:HG	1.40	0.85
31:BA:1502:A:H2	31:BA:1505:G:N1	1.74	0.85
32:BE:231:GLU:HB3	32:BE:232:PRO:HD2	1.59	0.85
35:BH:76:ILE:HG23	35:BH:77:PRO:HD2	1.59	0.85
38:BK:30:ARG:NH1	38:BK:30:ARG:HB3	1.92	0.85
52:CD:20:U:H2'	52:CD:21:A:H5'	1.59	0.85
38:CK:41:ARG:HD2	38:CK:41:ARG:O	1.77	0.85
2:DB:81:G:C2	2:DB:82:G:N7	2.44	0.85
7:DH:89:ILE:HD13	7:DH:90:LYS:N	1.92	0.85
58:DL:60:TYR:OH	58:DL:65:PHE:O	1.95	0.85
57:DY:9:LEU:HD22	57:DY:9:LEU:C	1.98	0.85
1:AA:654(C):G:H3'	1:AA:654(D):G:H8	1.41	0.85
1:AA:749:C:O2	1:AA:1618:A:H2'	1.77	0.85
39:BL:65:VAL:HG22	39:BL:66:ARG:N	1.92	0.85
26:A4:63:TYR:OH	49:BV:39:THR:HB	1.76	0.85
40:CM:39:PRO:HB3	40:CM:70:ARG:HH12	1.41	0.85
55:DA:1057:A:N7	55:DA:1086:A:C2'	2.37	0.85
3:DD:69:ARG:HD3	3:DD:105:ILE:HD11	1.59	0.85
9:DM:62:VAL:HG11	9:DM:66:LYS:HB2	1.56	0.85
21:DV:174:VAL:O	21:DV:175:VAL:HG13	1.75	0.85
57:DY:16:ASN:ND2	57:DY:25:PHE:CZ	2.45	0.85
1:AA:586:A:H5'	5:AF:89:VAL:HG21	1.58	0.84
21:AV:144:LEU:CD1	21:AV:146:ILE:O	2.25	0.84
52:CD:21:A:C2'	52:CD:22:G:H5''	2.06	0.84
55:DA:1454:U:O2'	55:DA:1455:G:N7	2.09	0.84
58:DL:14:ALA:HA	58:DL:49:GLY:CA	2.04	0.84
1:AA:1372:U:H5'	1:AA:1372:U:C6	2.07	0.84
1:AA:1689:A:H62	1:AA:1698:A:H2	1.25	0.84
11:AO:61:ARG:O	11:AO:62:LEU:HB3	1.76	0.84
12:AP:75:THR:HA	12:AP:88:GLY:CA	2.06	0.84
20:AU:72:VAL:HG23	20:AU:73:ARG:H	1.41	0.84
21:AV:141:VAL:HG13	21:AV:141:VAL:O	1.73	0.84
21:AV:53:ILE:HG22	21:AV:71:VAL:HG13	1.58	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:B1:53:U:O2'	53:B1:54:U:OP1	1.94	0.84
32:BE:56:ARG:HB3	32:BE:56:ARG:HH11	1.40	0.84
31:BA:1342:C:H1'	39:BL:124:GLN:NE2	1.92	0.84
42:CO:109:GLY:HA3	42:CO:121:GLY:O	1.77	0.84
55:DA:1378:A:H4'	55:DA:1379:A:OP1	1.75	0.84
56:DI:29:GLU:OE1	57:DY:139:VAL:HG21	1.77	0.84
1:AA:1360:A:H5'	1:AA:1361:G:OP2	1.77	0.84
7:AH:41:MET:HG3	7:AH:54:ARG:HA	1.58	0.84
11:AO:71:VAL:HG13	11:AO:72:PRO:HD3	1.58	0.84
49:BV:41:VAL:O	49:BV:45:VAL:HG13	1.77	0.84
54:CA:1126:U:H1'	54:CA:1280:A:C5	2.12	0.84
55:DA:996:A:H4'	16:D1:92:ARG:HE	1.42	0.84
55:DA:1725:G:C8	55:DA:1725:G:H5'	2.11	0.84
55:DA:1925:C:N4	55:DA:1926:U:C2	2.45	0.84
5:DF:127:GLU:O	5:DF:129:PHE:N	2.10	0.84
21:AV:186:GLU:OE2	21:AV:186:GLU:HA	1.75	0.84
33:BF:44:GLU:HG2	33:BF:52:LEU:HD11	1.58	0.84
54:CA:1129:C:C4'	54:CA:1130:A:H5'	2.08	0.84
52:CD:41:C:C2'	52:CD:42:C:H5''	2.07	0.84
43:CP:65:LYS:HD3	26:D4:50:VAL:HG11	1.58	0.84
55:DA:528:A:C3'	55:DA:529:A:H5''	2.08	0.84
5:DF:7:TYR:HB3	5:DF:21:ALA:HB1	1.57	0.84
7:DH:153:LYS:CG	7:DH:162:ILE:H	1.90	0.84
56:DI:3:LEU:HD23	56:DI:7:ARG:CD	2.06	0.84
56:DJ:12:LEU:H	56:DJ:13:SER:CB	1.86	0.84
58:DL:141:ALA:HB1	58:DL:142:PRO:CA	2.08	0.84
1:AA:2690:C:OP2	13:A0:14:SER:HB3	1.78	0.84
17:A2:30:GLY:N	17:A2:61:VAL:HG11	1.92	0.84
1:AA:141:A:H1'	1:AA:1408:C:O4'	1.76	0.84
1:AA:481:G:H1'	1:AA:506:G:H21	1.42	0.84
1:AA:895:U:H2'	1:AA:895:U:O2	1.75	0.84
1:AA:90:U:C2'	1:AA:91:A:H5''	2.07	0.84
1:AA:1006:C:H1'	9:AM:106:MET:HE3	1.59	0.84
1:AA:95:G:O2'	24:AW:48:HIS:HB3	1.76	0.84
31:BA:397:A:N3	31:BA:397:A:H3'	1.91	0.84
33:BF:188:LEU:HD12	33:BF:195:VAL:HG11	1.58	0.84
54:CA:1200:C:H4'	54:CA:1201:A:H5''	1.59	0.84
16:D1:90:VAL:HG22	17:D2:39:LEU:HB3	1.58	0.84
55:DA:69:C:O2'	55:DA:70:G:H5'	1.76	0.84
3:DD:238:GLY:O	3:DD:239:ARG:O	1.95	0.84
56:DJ:13:SER:OG	56:DJ:17:VAL:CG2	2.19	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:DU:47:LYS:HG2	20:DU:60:PHE:CE1	2.12	0.84
27:A5:55:ARG:O	27:A5:56:LYS:HB2	1.77	0.84
1:AA:2517:C:O2'	1:AA:2518:A:H3'	1.76	0.84
1:AA:384:U:C2'	1:AA:385:C:C5'	2.51	0.84
4:AE:154:LYS:HE3	4:AE:154:LYS:HA	1.59	0.84
31:BA:652:U:H1'	31:BA:653:A:H2	1.40	0.84
31:BA:792:A:N9	31:BA:794:A:N6	2.25	0.84
54:CA:84:U:H5'	54:CA:84:U:C6	2.13	0.84
35:CH:137:GLU:HA	35:CH:140:ARG:HH11	1.42	0.84
55:DA:1929:G:H4'	55:DA:1930:G:OP1	1.78	0.84
12:DP:10:ARG:HB2	12:DP:89:ASN:ND2	1.91	0.84
21:DV:121:HIS:NE2	21:DV:169:GLU:OE2	2.10	0.84
1:AA:1359:A:H3'	1:AA:1359:A:H8	0.97	0.84
1:AA:1894:C:O2'	1:AA:1895:C:H5'	1.77	0.84
1:AA:2820:A:C8	4:AE:191:PRO:HB3	2.13	0.84
20:AU:86:ARG:HB2	20:AU:95:LYS:HD2	1.58	0.84
38:BK:12:ARG:NH1	38:BK:27:PRO:HD3	1.92	0.84
55:DA:1086:A:H5'	55:DA:1103:A:H61	1.43	0.84
55:DA:389:G:H1	11:DO:71:VAL:HG12	1.43	0.84
17:A2:49:THR:HB	17:A2:50:PRO:CD	2.06	0.84
26:A4:56:VAL:HA	26:A4:60:GLN:NE2	1.91	0.84
30:A8:50:LEU:HG	30:A8:51:ALA:N	1.92	0.84
1:AA:1342:A:O2'	1:AA:1344:G:OP2	1.96	0.84
1:AA:625:G:O6	11:AO:107:LYS:HD3	1.77	0.84
1:AA:943:U:C2'	1:AA:944:G:H5'	2.08	0.84
31:BA:974:A:H1'	44:BQ:31:ARG:NE	1.92	0.84
52:BC:18:G:H5'	52:BC:19:G:OP2	1.77	0.84
32:BE:80:ILE:HD13	32:BE:211:ILE:HG22	1.60	0.84
30:D8:52:LYS:N	30:D8:53:PRO:HD2	1.91	0.84
55:DA:1111:A:O2'	55:DA:1112:G:H4'	1.76	0.84
55:DA:2653:U:O2'	7:DH:110:SER:HB2	1.78	0.84
55:DA:887:A:O2'	55:DA:888:C:O5'	1.94	0.84
58:DL:140:GLY:O	58:DL:141:ALA:HB2	1.77	0.84
57:DY:51:LEU:HD21	57:DY:82:PHE:CA	2.07	0.84
57:DY:9:LEU:O	57:DY:13:LEU:HB3	1.77	0.84
1:AA:1022:G:N2	1:AA:1142(A):A:H2	1.76	0.84
1:AA:2147:G:H2'	1:AA:2148:G:O4'	1.77	0.84
7:AH:87:LEU:HA	7:AH:163:TYR:O	1.77	0.84
8:AK:125:GLU:HA	8:AK:141:LYS:HB3	1.59	0.84
11:AO:90:ARG:HG2	11:AO:91:PHE:HD1	1.41	0.84
31:BA:429:U:H4'	31:BA:430:A:O5'	1.77	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:C1:33:G:H2'	53:C1:34:G:C8	2.13	0.84
36:CI:91:VAL:HG11	48:CU:72:ARG:HH12	1.42	0.84
55:DA:1568:G:H5'	3:DD:60:ARG:HA	1.59	0.84
7:DH:106:THR:HG22	7:DH:112:PRO:HB3	1.58	0.84
58:DL:109:LYS:HG2	58:DL:110:GLN:N	1.89	0.84
2:DB:91:C:OP1	12:DP:19:GLY:HA2	1.77	0.84
21:DV:194:PRO:C	21:DV:196:VAL:HG12	1.98	0.84
26:A4:36:CYS:O	26:A4:39:CYS:HB2	1.78	0.84
1:AA:1372:U:O5'	1:AA:1372:U:C5	2.30	0.84
4:AE:35:GLN:HG3	4:AE:64:LYS:HZ2	1.43	0.84
6:AG:28:VAL:O	6:AG:31:VAL:HG12	1.78	0.84
21:AV:115:GLY:HA2	21:AV:175:VAL:O	1.78	0.84
31:BA:96:G:H2'	31:BA:97:U:O4'	1.77	0.84
43:CP:116:THR:HG22	43:CP:117:VAL:N	1.92	0.84
55:DA:1079:C:H3'	55:DA:1080:A:C8	2.12	0.84
55:DA:1778:U:H2'	55:DA:1784:A:N6	1.92	0.84
3:DD:183:ARG:HG2	3:DD:183:ARG:HH11	1.42	0.84
58:DL:8:VAL:N	58:DL:57:ILE:HG13	1.93	0.84
57:DY:138:LEU:HD12	57:DY:140:GLY:N	1.93	0.84
57:DY:2:PRO:CG	57:DY:3:ASN:H	1.84	0.84
57:DY:42:GLN:O	57:DY:42:GLN:HG3	1.78	0.84
1:AA:1536:A:H3'	1:AA:1537:C:C6	2.12	0.83
8:AK:142:VAL:CG2	8:AK:143:SER:H	1.90	0.83
34:BG:150:GLU:O	34:BG:152:SER:N	2.10	0.83
47:BT:57:VAL:HA	47:BT:77:VAL:HG23	1.59	0.83
47:CT:4:LYS:HE3	47:CT:6:LEU:HD21	1.60	0.83
4:DE:13:ARG:HB3	4:DE:21:VAL:HG12	1.60	0.83
8:DK:13:GLY:HA3	8:DK:17:GLN:OE1	1.78	0.83
58:DL:50:ASP:H	58:DL:53:VAL:CG2	1.90	0.83
57:DY:129:PRO:HD2	57:DY:131:MET:H	1.42	0.83
23:DZ:92:LYS:HA	23:DZ:95:LEU:HB2	1.56	0.83
26:A4:2:LYS:HD2	26:A4:6:HIS:NE2	1.93	0.83
1:AA:603:A:H1'	1:AA:604:G:O4'	1.77	0.83
6:AG:136:ARG:O	6:AG:154:GLY:HA2	1.78	0.83
14:AQ:59:LYS:HG2	14:AQ:60:GLY:H	1.42	0.83
52:BD:15:G:H22	52:BD:48:C:H42	1.24	0.83
38:CK:41:ARG:HH11	38:CK:41:ARG:CG	1.90	0.83
26:D4:15:ILE:H	26:D4:15:ILE:HD13	1.43	0.83
55:DA:1047:G:H2'	55:DA:1110:G:N2	1.92	0.83
27:A5:16:ARG:HG2	27:A5:16:ARG:HH11	1.43	0.83
1:AA:1846:G:H5'	1:AA:1847:A:OP2	1.77	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:AG:109:VAL:HG13	26:A4:33:VAL:HG21	1.60	0.83
1:AA:2562:U:H1'	10:AN:23:ARG:NH1	1.93	0.83
15:AR:96:ARG:NH1	15:AR:96:ARG:HB2	1.94	0.83
3:DD:25:THR:HG21	3:DD:81:ALA:CB	2.05	0.83
4:DE:14:ILE:CG2	4:DE:15:PHE:H	1.84	0.83
8:DK:93:THR:HG22	8:DK:119:PRO:HB3	1.60	0.83
10:DN:113:LYS:O	10:DN:117:LEU:HD12	1.78	0.83
12:DP:83:MET:HB2	22:D3:7:LEU:HD12	1.60	0.83
14:DQ:106:ARG:HA	14:DQ:110:LEU:HD11	1.59	0.83
15:DR:90:GLN:NE2	15:DR:90:GLN:HA	1.93	0.83
21:DV:130:PRO:HA	21:DV:133:ILE:HD11	1.59	0.83
1:AA:1249:U:H2'	1:AA:1249:U:O2	1.78	0.83
3:AD:34:VAL:HG21	3:AD:103:ARG:HA	1.60	0.83
1:AA:2820:A:N7	4:AE:109:LYS:HE3	1.90	0.83
8:AK:41:GLU:O	8:AK:45:LYS:HG2	1.77	0.83
31:BA:1175:G:H2'	31:BA:1176:A:C8	2.13	0.83
32:BE:40:HIS:HB3	32:BE:190:THR:HG21	1.59	0.83
37:BJ:79:ARG:HA	37:BJ:83:ALA:O	1.78	0.83
49:BV:29:ARG:O	49:BV:30:LEU:HB2	1.78	0.83
44:CQ:13:THR:N	44:CQ:14:PRO:HD2	1.93	0.83
49:CV:10:PHE:HD1	49:CV:10:PHE:N	1.71	0.83
22:D3:32:ARG:N	22:D3:35:ASN:ND2	2.27	0.83
55:DA:13:A:O2'	55:DA:15:G:N7	2.12	0.83
7:DH:124:GLU:HB2	7:DH:132:ARG:HD2	1.60	0.83
58:DL:50:ASP:N	58:DL:53:VAL:HG21	1.93	0.83
26:A4:2:LYS:HB3	26:A4:6:HIS:NE2	1.93	0.83
1:AA:1829:A:C8	1:AA:1830:C:C5	2.66	0.83
32:BE:44:LEU:HD12	32:BE:45:GLN:N	1.94	0.83
31:BA:939:G:H5''	37:BJ:102:ARG:HH22	1.44	0.83
54:CA:1175:G:H2'	54:CA:1176:A:C8	2.14	0.83
28:D6:12:GLU:HA	28:D6:23:THR:HA	1.58	0.83
57:DY:28:ASN:ND2	57:DY:83:TYR:HD2	1.77	0.83
1:AA:387:U:C5	1:AA:387:U:OP2	2.30	0.83
12:AP:98:LYS:HB3	12:AP:99:PRO:HD2	1.60	0.83
34:BG:49:ARG:HH22	53:B1:57:U:H1'	1.43	0.83
38:BK:97:VAL:HA	38:BK:100:ILE:HD11	1.59	0.83
27:D5:49:CYS:HA	27:D5:58:LEU:HB3	1.58	0.83
55:DA:1526:G:C2'	55:DA:1527:G:H5'	2.08	0.83
55:DA:2176:A:H2'	55:DA:2177:C:C6	2.13	0.83
55:DA:2414:G:H21	11:DO:67:MET:CE	1.90	0.83
55:DA:50:U:H4'	55:DA:51:G:OP2	1.77	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:796:C:H2'	55:DA:797:C:C6	2.13	0.83
55:DA:442:G:N3	5:DF:48:THR:HG21	1.93	0.83
56:DI:24:ILE:HG12	56:DI:25:ASP:N	1.87	0.83
56:DI:7:ARG:CD	56:DI:8:ILE:HG12	2.07	0.83
21:DV:174:VAL:O	21:DV:175:VAL:HG22	1.77	0.83
24:DW:41:ILE:HD11	24:DW:44:LEU:CB	2.07	0.83
57:DY:130:THR:O	57:DY:131:MET:C	2.17	0.83
1:AA:857:C:H5'	22:A3:77:ARG:HH22	1.43	0.83
3:AD:35:LYS:HB3	3:AD:63:ARG:HA	1.61	0.83
4:AE:66:HIS:HE1	4:AE:73:GLU:HB2	1.43	0.83
1:AA:608:A:OP1	5:AF:100:THR:HG21	1.78	0.83
52:BD:72:C:H2'	52:BD:73:A:H5''	1.58	0.83
42:CO:60:LEU:CD2	42:CO:60:LEU:H	1.91	0.83
17:D2:34:GLU:HG3	17:D2:58:VAL:HG22	1.61	0.83
55:DA:1084:A:H1'	57:DY:53:VAL:CG1	2.08	0.83
55:DA:1734:C:H2'	55:DA:1735:C:H5''	1.58	0.83
58:DL:101:TRP:HA	58:DL:104:VAL:CB	2.08	0.83
21:DV:150:LEU:HD23	21:DV:151:HIS:H	1.42	0.83
57:DY:9:LEU:HD22	57:DY:10:LEU:N	1.93	0.83
57:DY:28:ASN:ND2	57:DY:83:TYR:CD2	2.46	0.83
27:A5:36:CYS:SG	27:A5:49:CYS:HB3	2.19	0.83
1:AA:1925:C:O2	1:AA:1925:C:H2'	1.76	0.83
20:AU:89:PHE:CD1	20:AU:90:LEU:HD23	2.14	0.83
37:BJ:26:PHE:O	37:BJ:30:ILE:HG12	1.78	0.83
39:BL:85:LEU:HD12	39:BL:86:VAL:N	1.93	0.83
48:BU:50:ILE:HD11	48:BU:70:ILE:HG21	1.61	0.83
54:CA:429:U:H4'	54:CA:430:A:O5'	1.76	0.83
38:CK:12:ARG:NH1	38:CK:27:PRO:HD3	1.94	0.83
29:D7:5:TRP:NE1	29:D7:7:PRO:HG3	1.94	0.83
3:DD:181:GLU:HA	3:DD:272:ALA:HB3	1.59	0.83
58:DL:112:MET:HE1	58:DL:123:ALA:CB	2.09	0.83
57:DY:27:VAL:CG2	57:DY:28:ASN:N	2.36	0.83
1:AA:1141:U:H4'	1:AA:1142(A):A:O4'	1.78	0.83
1:AA:2656:U:H5	1:AA:2664:G:N2	1.77	0.83
1:AA:84:A:H4'	1:AA:85:G:O5'	1.78	0.83
52:BB:10:G:H3'	52:BB:11:C:H5	1.43	0.83
37:CJ:21:VAL:HG23	37:CJ:22:LEU:H	1.44	0.83
6:DG:53:LEU:HD23	6:DG:54:GLU:N	1.94	0.83
58:DL:9:LYS:C	58:DL:10:LEU:HD23	1.99	0.83
58:DL:115:LEU:O	58:DL:116:ASN:HB2	1.76	0.83
58:DL:14:ALA:HB1	58:DL:50:ASP:CB	2.03	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:DT:50:LYS:H	19:DT:87:GLN:HE22	1.24	0.83
21:DV:191:VAL:O	21:DV:192:ALA:CB	2.27	0.83
57:DY:71:LEU:CD2	57:DY:72:ASP:H	1.92	0.83
1:AA:946:G:H2'	1:AA:947:G:C8	2.14	0.83
1:AA:1614:A:H62	18:AS:93:ALA:HB2	1.43	0.83
20:AU:97:ARG:HD3	20:AU:97:ARG:H	1.43	0.83
43:CP:3:ARG:CZ	43:CP:7:VAL:HG13	2.07	0.83
55:DA:302:C:H2'	55:DA:303:U:H6	1.43	0.83
58:DL:133:SER:HA	58:DL:137:GLU:OE1	1.79	0.83
11:DO:105:LEU:O	11:DO:106:LEU:HB2	1.77	0.83
57:DY:15:GLU:O	57:DY:16:ASN:CB	2.27	0.83
1:AA:1688:U:H1'	1:AA:1701:A:C6	2.14	0.82
1:AA:674:G:O2'	5:AF:74:ARG:HG3	1.79	0.82
1:AA:747:U:O2	1:AA:2014:A:H1'	1.77	0.82
4:AE:34:VAL:HG11	4:AE:64:LYS:HD3	1.60	0.82
31:BA:372:C:H5"	31:BA:373:A:OP1	1.78	0.82
31:BA:920:U:H2'	31:BA:921:U:C6	2.14	0.82
39:BL:16:ARG:CB	39:BL:16:ARG:HH11	1.92	0.82
53:C1:36:G:H3'	53:C1:37:G:H5"	1.58	0.82
54:CA:641:U:H4'	54:CA:642:A:OP1	1.79	0.82
34:CG:23:GLY:HA3	34:CG:112:VAL:CG2	2.09	0.82
42:CO:28:LYS:NZ	42:CO:33:ARG:HH22	1.76	0.82
49:CV:83:HIS:CD2	49:CV:84:GLY:N	2.46	0.82
55:DA:1372:U:C5	55:DA:1372:U:H5'	2.13	0.82
55:DA:654(R):C:H2'	55:DA:654(S):G:H8	1.41	0.82
52:CB:57:G:C5'	21:DV:182:LYS:NZ	2.42	0.82
21:DV:194:PRO:HG2	21:DV:196:VAL:HG11	1.61	0.82
57:DY:2:PRO:O	57:DY:3:ASN:HB2	1.79	0.82
1:AA:1565:C:H5"	3:AD:18:VAL:HG21	1.61	0.82
55:DA:1082:U:H5'	57:DY:45:LYS:O	1.80	0.82
21:DV:146:ILE:HA	21:DV:174:VAL:CB	2.09	0.82
57:DY:50:ARG:CA	57:DY:83:TYR:HD1	1.91	0.82
1:AA:1405:U:H2'	1:AA:1406:U:H6	1.43	0.82
1:AA:2712:U:O2'	1:AA:2712(A):A:O5'	1.97	0.82
21:AV:175:VAL:O	21:AV:177:PRO:CD	2.25	0.82
35:BH:79:GLU:HB3	35:BH:92:LYS:HA	1.59	0.82
26:D4:34:GLU:HG2	26:D4:35:VAL:H	1.43	0.82
27:D5:58:LEU:HD13	27:D5:60:VAL:HB	1.62	0.82
6:DG:109:VAL:HG11	26:D4:33:VAL:HG21	1.60	0.82
58:DL:18:THR:HG21	58:DL:38:VAL:HG12	1.59	0.82
21:DV:150:LEU:O	21:DV:170:THR:O	1.97	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2777:G:H5''	1:AA:2778:A:H5''	1.59	0.82
2:AB:39:A:C6	26:A4:1:MET:HB3	2.15	0.82
3:AD:44:ASN:CB	3:AD:49:ILE:HA	2.06	0.82
5:AF:25:PRO:HG3	5:AF:119:ARG:HD3	1.60	0.82
52:CC:35:A:C2'	52:CC:36:A:H5''	2.10	0.82
33:CF:175:LEU:H	33:CF:175:LEU:HD12	1.44	0.82
55:DA:2287:A:H62	55:DA:2344:U:H3	1.26	0.82
55:DA:2787:C:H1'	4:DE:62:PRO:HD3	1.60	0.82
55:DA:2811:G:P	4:DE:61:ARG:HG2	2.19	0.82
5:DF:34:TRP:CZ2	11:DO:8:PRO:HG3	2.14	0.82
58:DL:65:PHE:HD2	58:DL:65:PHE:C	1.83	0.82
21:DV:117:LEU:HD12	21:DV:117:LEU:H	1.44	0.82
57:DY:123:GLU:O	57:DY:127:GLU:CB	2.27	0.82
1:AA:1340:U:O2'	1:AA:1341:U:P	2.37	0.82
1:AA:2127:G:H3'	1:AA:2128:C:H5''	1.61	0.82
1:AA:879:G:H1	1:AA:898:C:H42	1.24	0.82
4:AE:9:VAL:HG23	4:AE:10:GLY:N	1.94	0.82
31:BA:1214:C:H5''	31:BA:1215:G:OP2	1.77	0.82
49:BV:63:THR:N	49:BV:66:MET:HE3	1.91	0.82
54:CA:405:U:H3'	54:CA:406:G:H5'	1.62	0.82
38:CK:6:ILE:CB	38:CK:85:ARG:HH12	1.92	0.82
40:CM:40:LEU:HB2	40:CM:69:ASN:HB2	1.60	0.82
49:CV:67:VAL:H	26:D4:59:PHE:HE1	1.23	0.82
55:DA:1056:G:N2	55:DA:1087:G:N1	2.28	0.82
55:DA:1526:G:H2'	55:DA:1527:G:H5'	1.60	0.82
55:DA:1924:C:H2'	55:DA:1925:C:O4'	1.80	0.82
52:CD:76:A:H8	55:DA:2394:C:H42	1.27	0.82
56:DI:17:VAL:HA	56:DI:20:LEU:CD1	2.09	0.82
57:DY:104:ILE:HG23	57:DY:105:PRO:N	1.94	0.82
57:DY:28:ASN:HB3	57:DY:81:VAL:CG1	2.08	0.82
27:A5:3:LYS:HA	27:A5:3:LYS:HE3	1.61	0.82
1:AA:1022:G:H22	1:AA:1142(A):A:H2	1.27	0.82
1:AA:762:U:H4'	1:AA:763:G:O5'	1.80	0.82
7:AH:7:LEU:N	7:AH:8:PRO:HD2	1.95	0.82
32:BE:167:PRO:HG3	32:BE:188:ALA:HB2	1.62	0.82
39:BL:66:ARG:HB3	39:BL:66:ARG:NH1	1.94	0.82
54:CA:1139:G:N2	54:CA:1144:G:H1	1.76	0.82
36:CI:27:GLN:HA	36:CI:27:GLN:HE21	1.45	0.82
37:CJ:94:ARG:HH11	37:CJ:94:ARG:HG3	1.45	0.82
55:DA:1378:A:O2'	55:DA:1379:A:C5'	2.28	0.82
55:DA:1943:U:H4'	55:DA:1944:U:O5'	1.79	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2317:C:C2'	55:DA:2318:G:H5'	2.08	0.82
7:DH:55:PRO:HG2	7:DH:61:HIS:ND1	1.95	0.82
58:DL:8:VAL:H	58:DL:57:ILE:HG13	1.44	0.82
57:DY:27:VAL:O	57:DY:81:VAL:O	1.98	0.82
16:A1:90:VAL:HG22	17:A2:39:LEU:HB3	1.60	0.82
3:AD:43:ARG:HH11	3:AD:44:ASN:HD21	1.22	0.82
55:DA:638:G:H2'	55:DA:639:U:C6	2.15	0.82
57:DY:25:PHE:HD1	57:DY:82:PHE:CE2	1.91	0.82
57:DY:43:ALA:H	57:DY:47:ASN:HD21	1.27	0.82
57:DY:50:ARG:CD	57:DY:51:LEU:H	1.92	0.82
57:DY:75:GLN:CB	57:DY:111:LEU:HA	2.10	0.82
57:DY:76:GLY:O	57:DY:111:LEU:CB	2.27	0.82
1:AA:528:A:C2	1:AA:2042:A:H2'	2.14	0.82
31:BA:1062:U:H2'	31:BA:1063:C:C6	2.15	0.82
31:BA:1277:C:HO2'	31:BA:1279:A:H8	1.25	0.82
52:BD:21:A:H3'	52:BD:21:A:N3	1.95	0.82
39:BL:9:ARG:HA	39:BL:13:ALA:O	1.80	0.82
40:BM:40:LEU:HG	40:BM:41:PRO:HD2	1.60	0.82
32:CE:30:ARG:HG3	32:CE:31:TYR:CE1	2.15	0.82
32:CE:18:GLY:H	32:CE:42:ILE:HG22	1.45	0.82
32:CE:8:LYS:H	32:CE:8:LYS:CD	1.88	0.82
33:CF:47:LEU:HD11	33:CF:76:VAL:HG12	1.60	0.82
55:DA:2067:G:H4'	55:DA:2068:U:OP2	1.80	0.82
55:DA:265:A:O2'	55:DA:266:G:C4'	2.27	0.82
3:DD:80:ALA:HB3	3:DD:94:LEU:HD12	1.62	0.82
11:DO:120:ALA:HB2	11:DO:137:LYS:HB3	1.62	0.82
15:DR:91:ARG:O	15:DR:116:ALA:HA	1.78	0.82
21:DV:61:LEU:HD11	21:DV:65:GLN:CB	2.10	0.82
57:DY:27:VAL:CG2	57:DY:110:GLY:HA2	2.05	0.82
57:DY:70:GLU:O	57:DY:71:LEU:CD1	2.28	0.82
57:DY:90:ALA:O	57:DY:94:VAL:CB	2.26	0.82
1:AA:84:A:N6	1:AA:102:G:O2'	2.13	0.82
1:AA:2532:G:H4'	1:AA:2657:A:N1	1.95	0.82
1:AA:2882:A:H5'	13:A0:96:ARG:HG3	1.60	0.82
1:AA:384:U:H2'	1:AA:385:C:C5'	1.98	0.82
4:AE:9:VAL:HG23	4:AE:10:GLY:H	1.45	0.82
12:AP:56:ARG:NH1	12:AP:56:ARG:HB2	1.95	0.82
21:AV:175:VAL:CG1	21:AV:177:PRO:HD2	1.93	0.82
54:CA:351:G:H4'	54:CA:352:C:OP1	1.79	0.82
49:CV:83:HIS:O	49:CV:86:GLU:N	2.13	0.82
55:DA:1058:U:OP1	58:DL:5:VAL:HG22	1.79	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1079:C:H3'	55:DA:1080:A:H8	1.43	0.82
12:DP:20:ALA:HA	12:DP:98:LYS:HB3	1.59	0.82
24:DW:65:ASN:HD22	24:DW:69:ARG:HH21	1.26	0.82
12:AP:16:ARG:HG3	12:AP:17:LEU:N	1.94	0.82
21:AV:127:LYS:HB3	21:AV:162:GLU:HB3	1.61	0.82
1:AA:1913:A:N7	31:BA:1494:G:H4'	1.95	0.82
46:BS:43:LYS:HA	46:BS:48:TRP:HB3	1.61	0.82
31:BA:1313:U:OP1	49:BV:6:LYS:HB3	1.80	0.82
54:CA:1062:U:H2'	54:CA:1063:C:C6	2.15	0.82
54:CA:251:G:H4'	54:CA:252:U:O5'	1.80	0.82
48:CU:86:VAL:HG12	48:CU:87:ARG:HG2	1.62	0.82
26:D4:68:ARG:NH1	26:D4:68:ARG:HA	1.94	0.82
55:DA:2287:A:O2'	55:DA:2288:A:H3'	1.78	0.82
56:DI:16:THR:CG2	56:DI:17:VAL:H	1.92	0.82
11:DO:64:LYS:C	11:DO:66:GLY:N	2.32	0.82
57:DY:141:VAL:HG13	57:DY:142:LEU:N	1.94	0.82
1:AA:2879:C:H4'	1:AA:2880:C:OP1	1.78	0.81
1:AA:2060:A:OP1	5:AF:68:LYS:O	1.97	0.81
14:AQ:19:LYS:O	14:AQ:20:ARG:HB3	1.79	0.81
24:AW:51:ARG:HE	24:AW:55:ARG:NH1	1.78	0.81
31:BA:792:A:O2'	31:BA:794:A:N7	2.13	0.81
52:CD:8:U:H4'	52:CD:9:A:OP1	1.80	0.81
55:DA:2419:U:C4'	28:D6:23:THR:HG21	2.10	0.81
55:DA:270(K):C:C2'	55:DA:270(L):U:H5''	2.09	0.81
6:DG:94:LEU:HD23	6:DG:94:LEU:H	1.45	0.81
58:DL:103:GLN:O	58:DL:107:ILE:CB	2.26	0.81
58:DL:11:GLN:HG2	58:DL:41:PHE:CZ	2.14	0.81
9:DM:47:ALA:HB2	9:DM:112:LEU:HG	1.61	0.81
15:DR:28:VAL:HG23	15:DR:87:ASP:O	1.80	0.81
57:DY:43:ALA:N	57:DY:47:ASN:HD21	1.77	0.81
17:A2:38:LEU:C	17:A2:39:LEU:HD12	2.00	0.81
1:AA:789:A:H3'	1:AA:789:A:OP1	1.80	0.81
10:AN:104:ARG:HB3	10:AN:104:ARG:HH11	1.44	0.81
31:BA:1239:A:H2'	31:BA:1298:C:N4	1.94	0.81
49:BV:23:ASN:HB2	49:BV:43:GLU:OE2	1.79	0.81
54:CA:64:G:H4'	54:CA:65:U:C5'	2.10	0.81
35:CH:101:ILE:HD11	35:CH:119:LEU:HD23	1.61	0.81
35:CH:11:ILE:O	35:CH:12:LEU:HB2	1.78	0.81
38:CK:41:ARG:HG3	38:CK:41:ARG:HH11	1.45	0.81
48:CU:18:ARG:H	48:CU:18:ARG:HD2	1.43	0.81
48:CU:18:ARG:HD2	48:CU:18:ARG:N	1.93	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1049:C:N3	55:DA:2751:G:O6	2.13	0.81
55:DA:1078:U:H1'	55:DA:1088:A:C2	2.14	0.81
55:DA:654(O):G:H2'	55:DA:654(P):G:H8	1.45	0.81
8:DK:3:VAL:HG12	8:DK:38:LEU:HA	1.61	0.81
57:DY:88:ALA:HB1	57:DY:92:THR:OG1	1.79	0.81
29:A7:19:ARG:HH11	29:A7:19:ARG:HG2	1.43	0.81
1:AA:2287:A:N6	1:AA:2344:U:H3	1.78	0.81
3:AD:69:ARG:HD2	3:AD:119:ALA:HB2	1.63	0.81
11:AO:90:ARG:HG2	11:AO:91:PHE:CD1	2.14	0.81
20:AU:6:HIS:O	20:AU:7:VAL:HG13	1.80	0.81
21:AV:144:LEU:O	21:AV:174:VAL:CG2	2.26	0.81
31:BA:1205:U:H1'	33:BF:195:VAL:HG23	1.61	0.81
31:BA:1435:G:H2'	31:BA:1436:U:C6	2.14	0.81
31:BA:689:C:C2'	31:BA:690:G:H5'	2.11	0.81
35:BH:78:HIS:HB2	38:BK:104:ARG:HD2	1.60	0.81
38:BK:82:HIS:HD2	38:BK:138:TRP:HE1	1.28	0.81
40:BM:78:ASN:HB2	40:BM:81:THR:HG23	1.63	0.81
46:BS:7:ALA:HB2	46:BS:20:VAL:HG11	1.59	0.81
46:BS:53:VAL:HG12	46:BS:79:VAL:HG22	1.59	0.81
52:CD:2:C:H5''	52:CD:2:C:H6	1.45	0.81
37:CJ:140:ASP:HA	37:CJ:143:ARG:NH1	1.95	0.81
50:CW:23:ARG:HA	50:CW:26:ASN:HD21	1.45	0.81
55:DA:1734:C:C3'	55:DA:1735:C:H5''	2.10	0.81
55:DA:1784:A:H4'	55:DA:1785:A:O5'	1.80	0.81
2:DB:15:A:H5'	2:DB:16:G:C8	2.14	0.81
7:DH:126:PRO:HD2	7:DH:127:GLU:O	1.80	0.81
7:DH:4:ILE:HD13	7:DH:4:ILE:H	1.45	0.81
56:DJ:10:GLU:OE1	56:DJ:19:GLU:OE2	1.99	0.81
58:DL:112:MET:N	58:DL:113:PRO:CD	2.40	0.81
21:DV:191:VAL:CG1	21:DV:197:ILE:HG23	2.10	0.81
22:A3:68:GLU:HG3	22:A3:80:HIS:HB2	1.60	0.81
1:AA:180:G:OP2	29:A7:32:LYS:HE2	1.80	0.81
2:AB:81:G:O6	2:AB:96:G:C6	2.34	0.81
5:AF:4:VAL:HA	5:AF:19:GLU:CB	2.09	0.81
12:AP:24:GLY:O	12:AP:25:ASP:HB2	1.77	0.81
31:BA:134:A:H61	46:BS:25:ARG:HH12	1.28	0.81
34:CG:12:CYS:HA	34:CG:19:LEU:CD2	2.11	0.81
38:CK:34:GLU:HB3	38:CK:118:VAL:HG21	1.61	0.81
55:DA:1673:U:H2'	55:DA:1674:G:H5'	1.60	0.81
55:DA:2776:A:H4'	55:DA:2777:G:O5'	1.79	0.81
3:DD:35:LYS:HD3	3:DD:63:ARG:CA	2.11	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:389:G:N1	11:DO:71:VAL:HG12	1.94	0.81
24:DW:47:ASN:H	24:DW:47:ASN:ND2	1.78	0.81
1:AA:1324:G:H1'	1:AA:1616:A:H62	1.46	0.81
54:CA:67:C:H2'	54:CA:68:G:C8	2.14	0.81
54:CA:819:A:H5''	54:CA:820:U:OP2	1.80	0.81
55:DA:993:G:OP1	16:D1:50:ARG:NH2	2.14	0.81
2:DB:42:C:H4'	6:DG:67:LYS:CD	2.10	0.81
14:DQ:106:ARG:CA	14:DQ:110:LEU:HD21	2.11	0.81
21:DV:114:GLY:HA3	21:DV:178:GLU:H	1.43	0.81
1:AA:363(A):A:H2'	1:AA:363(B):G:H5''	1.63	0.81
11:AO:19:VAL:HG22	11:AO:20:GLY:N	1.95	0.81
12:AP:56:ARG:HH11	12:AP:56:ARG:HB2	1.46	0.81
21:AV:103:ARG:HB3	21:AV:138:GLU:HA	1.62	0.81
38:BK:82:HIS:HD2	38:BK:138:TRP:NE1	1.78	0.81
51:BX:26:LYS:NZ	51:BX:26:LYS:HA	1.95	0.81
54:CA:22:G:H2'	54:CA:23:C:C6	2.15	0.81
32:CE:183:PRO:HA	32:CE:198:ASP:OD1	1.79	0.81
26:D4:39:CYS:O	26:D4:41:PRO:HD3	1.80	0.81
55:DA:1077:A:H2	58:DL:133:SER:HB3	1.45	0.81
55:DA:1142(A):A:HO2'	55:DA:1143:A:H3'	1.41	0.81
55:DA:2790:A:H2	55:DA:2894:G:H5''	1.45	0.81
55:DA:897:C:H5	55:DA:897:C:P	2.03	0.81
7:DH:150:ALA:C	7:DH:152:ARG:H	1.77	0.81
56:DI:28:LYS:CA	56:DI:28:LYS:HE3	2.08	0.81
56:DJ:13:SER:HA	56:DJ:17:VAL:CG2	2.10	0.81
18:DS:9:TYR:H	18:DS:102:HIS:HD2	1.27	0.81
20:DU:97:ARG:NH2	20:DU:98:VAL:HB	1.95	0.81
57:DY:130:THR:O	57:DY:132:ASP:N	2.14	0.81
57:DY:70:GLU:O	57:DY:71:LEU:CG	2.28	0.81
1:AA:1496:A:H2'	1:AA:1498:C:C5	2.15	0.81
1:AA:2311:A:H3'	1:AA:2312:U:C5	2.15	0.81
4:AE:36:ARG:HH21	4:AE:88:GLY:HA3	1.44	0.81
31:BA:1298:C:C5	37:BJ:114:ARG:HD2	2.16	0.81
31:BA:216:G:O2'	31:BA:217:C:H6	1.61	0.81
22:A3:7:LEU:HA	52:BC:2:C:H5'	1.60	0.81
32:BE:91:PRO:HG3	32:BE:154:LEU:HB2	1.60	0.81
55:DA:1084:A:H1'	57:DY:53:VAL:HG11	1.62	0.81
55:DA:897:C:C6	55:DA:897:C:P	2.74	0.81
5:DF:32:LEU:CD1	5:DF:105:VAL:HG13	2.10	0.81
56:DI:21:LYS:HA	56:DI:24:ILE:HD12	1.63	0.81
56:DJ:7:ARG:HH11	56:DJ:7:ARG:HG2	1.43	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:DL:52:ILE:CG2	58:DL:75:SER:CB	2.40	0.81
21:DV:108:PRO:HG2	21:DV:110:GLY:H	1.45	0.81
30:A8:14:VAL:HG11	30:A8:22:VAL:HG13	1.62	0.81
30:A8:48:PHE:O	30:A8:49:VAL:HG23	1.81	0.81
1:AA:1342:A:N7	1:AA:1345:C:C5	2.48	0.81
2:AB:81:G:C2	2:AB:82:G:N7	2.49	0.81
31:BA:738:C:H5''	36:BI:69:GLU:HB2	1.62	0.81
42:CO:6:THR:H	42:CO:9:GLN:HE21	1.26	0.81
55:DA:654(R):C:H2'	55:DA:654(S):G:C8	2.15	0.81
3:DD:135:PHE:HD2	3:DD:135:PHE:N	1.79	0.81
8:DK:110:ASP:HB2	8:DK:111:PRO:C	2.01	0.81
58:DL:107:ILE:HG22	58:DL:108:ALA:H	1.43	0.81
20:DU:47:LYS:HG2	20:DU:60:PHE:HE1	1.45	0.81
57:DY:28:ASN:CG	57:DY:83:TYR:CE2	2.53	0.81
1:AA:1771:C:HO2'	1:AA:1786:A:H8	1.27	0.81
1:AA:84:A:H61	1:AA:102:G:C2'	1.94	0.81
8:AK:75:LEU:HD22	8:AK:77:LEU:HD22	1.63	0.81
38:BK:103:VAL:HG21	38:BK:110:ALA:HB2	1.60	0.81
48:BU:41:LYS:HD3	48:BU:41:LYS:O	1.80	0.81
53:C1:52:U:C2'	53:C1:53:U:H5''	2.11	0.81
54:CA:1502:A:H2	54:CA:1505:G:N1	1.79	0.81
5:DF:7:TYR:HB3	5:DF:21:ALA:CB	2.11	0.81
7:DH:4:ILE:HG13	7:DH:6:ARG:CZ	2.10	0.81
57:DY:54:ALA:CB	57:DY:57:THR:HB	2.10	0.81
1:AA:1083:U:H1'	1:AA:1086:A:N6	1.95	0.81
1:AA:2402:C:H5	1:AA:2415:G:H22	1.29	0.81
3:AD:65:ILE:HD12	3:AD:66:ASP:N	1.96	0.81
9:AM:125:GLY:HA3	9:AM:126:PRO:O	1.80	0.81
11:AO:112:LEU:HD22	11:AO:113:LYS:N	1.95	0.81
18:AS:73:ALA:HB3	18:AS:106:ILE:HG12	1.62	0.81
23:AZ:82:LEU:HD23	23:AZ:82:LEU:H	1.46	0.81
33:BF:155:GLY:O	33:BF:156:ARG:HB2	1.81	0.81
36:BI:37:VAL:HA	36:BI:65:VAL:HG12	1.63	0.81
54:CA:164:U:H2'	54:CA:165:C:C6	2.16	0.81
54:CA:232:G:H2'	54:CA:233:C:H6	1.46	0.81
54:CA:22:G:H2'	54:CA:23:C:H6	1.44	0.81
35:CH:153:LYS:HE3	35:CH:155:GLU:HB3	1.62	0.81
49:CV:81:ARG:HG3	49:CV:82:GLY:H	1.46	0.81
55:DA:442:G:H21	5:DF:48:THR:HG23	1.43	0.81
58:DL:113:PRO:O	58:DL:114:ASP:CB	2.28	0.81
55:DA:1079:C:O2'	58:DL:129:GLY:HA2	1.81	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:DM:58:ASP:H	9:DM:60:ILE:HD13	1.46	0.81
20:DU:84:ARG:HH12	20:DU:97:ARG:HA	1.44	0.81
21:DV:176:PRO:O	21:DV:178:GLU:HG2	1.80	0.81
28:A6:15:GLU:OE2	28:A6:41:PRO:HB3	1.81	0.81
1:AA:322:A:H5''	5:AF:169:ASN:ND2	1.95	0.81
3:AD:142:VAL:HG23	3:AD:193:VAL:HA	1.62	0.81
49:BV:51:VAL:O	49:BV:58:VAL:HG22	1.81	0.81
43:CP:108:ARG:HD2	43:CP:108:ARG:H	1.45	0.81
45:CR:4:THR:OG1	45:CR:7:GLU:HB2	1.80	0.81
47:CT:67:LYS:HA	47:CT:70:ARG:NH1	1.95	0.81
17:D2:15:GLU:HG3	17:D2:16:PRO:HD2	1.60	0.81
55:DA:70:G:H2'	55:DA:113:G:O2'	1.80	0.81
4:DE:9:VAL:HB	4:DE:25:VAL:HG23	1.62	0.81
21:DV:112:ARG:N	21:DV:112:ARG:HD2	1.95	0.81
21:DV:105:VAL:HG12	21:DV:140:ASP:HB3	1.61	0.81
22:A3:32:ARG:H	22:A3:35:ASN:HD21	1.24	0.80
1:AA:2789:C:H2'	1:AA:2790:A:H5''	1.62	0.80
3:AD:65:ILE:HD11	3:AD:67:PHE:CD2	2.15	0.80
5:AF:10:PRO:HD2	5:AF:13:SER:OG	1.81	0.80
31:BA:1007:C:C2'	31:BA:1008:C:H5''	2.11	0.80
31:BA:216:G:HO2'	31:BA:217:C:H6	0.82	0.80
37:BJ:115:ARG:O	37:BJ:118:VAL:HG22	1.80	0.80
17:D2:16:PRO:HB3	17:D2:99:ILE:HD11	1.63	0.80
55:DA:1372:U:C4'	55:DA:1372:U:C6	2.57	0.80
56:DI:21:LYS:O	56:DI:26:ALA:HB3	1.79	0.80
1:AA:1652:A:H62	13:A0:11:ASN:HD21	1.27	0.80
5:AF:22:ALA:C	5:AF:24:LEU:H	1.83	0.80
20:AU:63:LYS:HZ2	20:AU:64:GLU:N	1.79	0.80
31:BA:1160:G:H1	31:BA:1177:G:H21	1.25	0.80
31:BA:968:A:H4'	31:BA:969:A:OP2	1.80	0.80
54:CA:57:G:H2'	54:CA:58:C:C6	2.16	0.80
49:CV:41:VAL:HG21	49:CV:67:VAL:HG22	1.62	0.80
55:DA:1063:G:C1'	58:DL:134:MET:HE1	2.11	0.80
55:DA:1080:A:H2'	55:DA:1081:U:O4'	1.81	0.80
55:DA:1212:G:O2'	55:DA:1213:A:OP2	1.99	0.80
3:DD:35:LYS:HZ1	3:DD:104:TYR:HB2	1.46	0.80
7:DH:152:ARG:HG3	7:DH:153:LYS:CE	2.10	0.80
11:DO:64:LYS:O	11:DO:66:GLY:N	2.15	0.80
31:BA:192:U:O4'	50:BW:103:GLY:HA2	1.82	0.80
31:BA:559:A:H4'	31:BA:560:U:H5''	1.60	0.80
39:BL:10:ARG:HH21	39:BL:11:LYS:HB2	1.44	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:31:G:N2	54:CA:47:C:H5''	1.97	0.80
33:CF:116:VAL:HG21	33:CF:202:ILE:HD11	1.61	0.80
58:DL:21:PRO:CG	58:DL:24:GLY:HA3	2.10	0.80
58:DL:93:ARG:HH11	58:DL:135:GLY:HA2	1.42	0.80
9:DM:7:LYS:HD2	9:DM:7:LYS:H	1.46	0.80
1:AA:976:C:H5'	1:AA:1156:A:N6	1.97	0.80
1:AA:686:G:N2	1:AA:788:A:H61	1.80	0.80
2:AB:40:U:N3	26:A4:1:MET:SD	2.55	0.80
4:AE:52:LEU:O	4:AE:74:PRO:HA	1.80	0.80
9:AM:39:ARG:HH21	9:AM:41:ASP:CB	1.93	0.80
20:AU:61:ILE:HG22	20:AU:62:GLU:N	1.94	0.80
21:AV:145:GLU:O	21:AV:146:ILE:CD1	2.29	0.80
52:BD:16:U:O2	52:BD:18:G:H5'	1.81	0.80
34:BG:176:LEU:HD12	34:BG:177:ASP:H	1.45	0.80
54:CA:1178:G:H5'	39:CL:93:ARG:NH2	1.96	0.80
33:CF:73:PRO:O	33:CF:76:VAL:HG22	1.82	0.80
43:CP:14:ARG:HA	43:CP:44:ARG:HA	1.61	0.80
44:CQ:2:ALA:O	44:CQ:6:LEU:HD12	1.81	0.80
54:CA:617:G:H4'	46:CS:44:THR:HB	1.63	0.80
49:CV:44:MET:HA	49:CV:47:HIS:HD2	1.47	0.80
55:DA:1077:A:C3'	55:DA:1078:U:C5'	2.51	0.80
55:DA:1673:U:C2'	55:DA:1674:G:H5'	2.12	0.80
58:DL:19:PRO:CA	58:DL:25:PRO:CD	2.59	0.80
1:AA:2893:G:C5'	1:AA:2894:G:H5'	2.03	0.80
1:AA:49:A:H4'	1:AA:50:U:H5'	1.63	0.80
5:AF:66:PRO:O	5:AF:67:GLN:HB3	1.79	0.80
21:AV:103:ARG:O	21:AV:104:PHE:HB2	1.81	0.80
21:AV:60:GLU:O	21:AV:61:LEU:HB2	1.81	0.80
31:BA:518:C:H4'	31:BA:519:C:O5'	1.80	0.80
31:BA:533:A:O2'	31:BA:534:U:H5'	1.81	0.80
34:BG:110:PHE:H	34:BG:110:PHE:HD1	1.30	0.80
50:BW:68:LYS:HD2	50:BW:69:GLY:N	1.96	0.80
52:CD:16:U:H2'	52:CD:17:C:H5'	1.62	0.80
43:CP:124:PRO:HB3	43:CP:125:ARG:HG2	0.83	0.80
44:CQ:39:LEU:HB3	44:CQ:43:CYS:HB2	1.64	0.80
55:DA:1020:A:N1	55:DA:1141:U:H2'	1.97	0.80
56:DJ:18:LEU:CA	56:DJ:21:LYS:HB2	2.10	0.80
1:AA:1164:G:H2'	1:AA:1165:U:C6	2.16	0.80
1:AA:363(F):A:H5'	1:AA:364:C:OP1	1.80	0.80
1:AA:945:A:C4'	1:AA:946:G:OP1	2.30	0.80
2:AB:7:G:C3'	2:AB:8:U:H5''	2.12	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:AD:35:LYS:HD3	3:AD:63:ARG:CB	2.12	0.80
11:AO:124:LYS:NZ	11:AO:143:GLY:HA3	1.96	0.80
31:BA:531:U:H4'	31:BA:532:A:OP1	1.80	0.80
35:BH:148:VAL:HG21	38:BK:107:LEU:HD22	1.64	0.80
49:BV:40:ILE:HG23	49:BV:67:VAL:HA	1.63	0.80
54:CA:1028:C:C2'	54:CA:1028(A):C:H5''	2.10	0.80
54:CA:73:G:H21	54:CA:74:C:H41	1.30	0.80
56:DJ:5:ILE:C	56:DJ:7:ARG:H	1.84	0.80
10:DN:113:LYS:O	10:DN:116:SER:HB3	1.81	0.80
57:DY:21:GLN:HA	57:DY:21:GLN:NE2	1.96	0.80
57:DY:21:GLN:NE2	57:DY:22:GLY:N	2.05	0.80
57:DY:74:LEU:HD13	57:DY:74:LEU:C	2.00	0.80
31:BA:973:G:H3'	31:BA:974:A:H5''	1.62	0.80
54:CA:1126:U:OP2	54:CA:1281:U:H1'	1.81	0.80
52:CD:72:C:C2'	52:CD:73:A:H5''	2.11	0.80
55:DA:1734:C:C2'	55:DA:1735:C:H5''	2.11	0.80
55:DA:897:C:H2'	55:DA:898:C:O4'	1.81	0.80
56:DJ:5:ILE:HG22	56:DJ:9:LYS:CB	2.11	0.80
8:DK:88:ILE:HG12	8:DK:122:GLU:H	1.47	0.80
58:DL:42:ASN:O	58:DL:46:ALA:HB2	1.80	0.80
58:DL:51:ALA:O	58:DL:52:ILE:CG1	2.30	0.80
9:DM:8:GLN:O	9:DM:9:VAL:HG22	1.81	0.80
21:DV:112:ARG:NH1	21:DV:112:ARG:HG3	1.92	0.80
21:DV:196:VAL:O	21:DV:197:ILE:HG13	1.82	0.80
6:AG:108:ASN:HA	26:A4:38:LYS:HB2	1.64	0.80
1:AA:1310:G:OP2	29:A7:9:ARG:NH1	2.14	0.80
1:AA:2348:U:H2'	1:AA:2349:G:H5''	1.63	0.80
1:AA:614:U:H4'	1:AA:615:G:H5''	1.62	0.80
1:AA:943:U:OP2	11:AO:36:LYS:HG3	1.80	0.80
1:AA:955:C:H5'	1:AA:956:G:OP2	1.82	0.80
11:AO:83:VAL:HG12	11:AO:112:LEU:HD21	1.64	0.80
23:AZ:89:GLU:HA	23:AZ:93:GLU:HG3	1.62	0.80
31:BA:1072:G:H2'	31:BA:1073:U:H6	1.47	0.80
54:CA:77:C:C2'	54:CA:78:G:H5''	2.11	0.80
43:CP:126:LYS:O	43:CP:126:LYS:HG2	1.79	0.80
44:CQ:40:CYS:H	44:CQ:43:CYS:HB2	1.44	0.80
55:DA:1899:G:H21	55:DA:1902:C:H5	1.26	0.80
56:DI:24:ILE:O	56:DI:27:LEU:N	2.15	0.80
56:DJ:15:ALA:O	56:DJ:16:THR:CG2	2.30	0.80
12:DP:104:PHE:HE1	12:DP:125:LEU:HD11	1.47	0.80
15:DR:111:ARG:O	15:DR:112:ARG:HG3	1.81	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2815:C:H2'	1:AA:2816:C:H6	1.46	0.80
1:AA:387:U:O2'	1:AA:388:G:P	2.39	0.80
2:AB:15:A:H3'	2:AB:16:G:H5'	1.61	0.80
25:AX:8:LEU:HD13	25:AX:31:LEU:HD12	1.64	0.80
31:BA:631:G:O3'	31:BA:632:A:H8	1.65	0.80
33:BF:15:THR:HG21	33:BF:181:ASN:HA	1.63	0.80
42:BO:59:ARG:HD3	42:BO:65:GLU:HG3	1.63	0.80
54:CA:1536:C:H1'	53:C1:37:G:N2	1.96	0.80
33:CF:59:ARG:HH12	33:CF:97:LYS:HE3	1.47	0.80
54:CA:1117:G:H4'	39:CL:104:ARG:NH2	1.97	0.80
28:D6:9:LEU:HD22	28:D6:11:LEU:HD22	1.63	0.80
55:DA:1071:G:H4'	55:DA:1089:G:OP2	1.82	0.80
55:DA:1078:U:O2'	55:DA:1079:C:H5''	1.82	0.80
55:DA:1165:U:H2'	55:DA:1166:C:C6	2.16	0.80
55:DA:654(L):G:C2	55:DA:654(M):C:H1'	2.16	0.80
56:DJ:10:GLU:CD	56:DJ:19:GLU:OE2	2.19	0.80
56:DJ:5:ILE:HG22	56:DJ:9:LYS:HG3	1.63	0.80
58:DL:8:VAL:O	58:DL:57:ILE:CB	2.30	0.80
58:DL:65:PHE:C	58:DL:65:PHE:CD2	2.54	0.80
11:DO:50:ARG:HH21	11:DO:50:ARG:CB	1.94	0.80
21:DV:152:ALA:HB1	21:DV:163:LEU:CD2	2.11	0.80
57:DY:122:VAL:HA	57:DY:126:ALA:CB	2.11	0.80
57:DY:135:ARG:HB2	56:DJ:19:GLU:OE1	1.80	0.80
57:DY:58:LEU:HD23	57:DY:58:LEU:N	1.95	0.80
26:A4:9:LEU:HD21	26:A4:25:TYR:HB3	1.60	0.80
1:AA:1149:G:H2'	1:AA:1150:C:H6	1.44	0.80
1:AA:2012:G:O3'	18:AS:96:ILE:HG13	1.80	0.80
31:BA:1067:A:HO2'	31:BA:1068:G:H8	1.29	0.80
51:BX:25:LYS:HG2	51:BX:26:LYS:HG2	1.64	0.80
39:CL:83:ARG:O	39:CL:86:VAL:HG12	1.80	0.80
55:DA:1056:G:H1'	55:DA:1086:A:O4'	1.82	0.80
55:DA:1077:A:N3	55:DA:1078:U:H5''	1.96	0.80
55:DA:1899:G:N2	55:DA:1902:C:C5	2.50	0.80
55:DA:2127:G:H3'	55:DA:2128:C:H5''	1.63	0.80
3:DD:176:ARG:HG2	3:DD:176:ARG:HH11	1.47	0.80
57:DY:112:LEU:CD1	57:DY:121:ASP:OD2	2.29	0.80
57:DY:23:SER:O	57:DY:24:PHE:HB3	1.81	0.80
17:A2:24:LYS:HA	17:A2:92:THR:OG1	1.82	0.79
1:AA:1074:G:H2'	1:AA:1075:C:C6	2.17	0.79
1:AA:141:A:H8	1:AA:1595:G:H21	1.27	0.79
1:AA:1645:G:H5''	1:AA:1646:C:H5'	1.64	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1826:G:H4'	3:AD:242:ARG:NH2	1.96	0.79
1:AA:2529:G:H5'	1:AA:2530:A:C5'	2.11	0.79
1:AA:387:U:H5	1:AA:387:U:OP2	1.64	0.79
1:AA:445:C:H2'	1:AA:446:G:O4'	1.82	0.79
1:AA:449:A:H4'	16:A1:3:ARG:NH1	1.96	0.79
1:AA:654(Q):C:H2'	1:AA:654(R):C:C6	2.16	0.79
8:AK:131:LYS:HB3	8:AK:132:PRO:CA	2.11	0.79
8:AK:143:SER:O	8:AK:144:VAL:HG23	1.81	0.79
14:AQ:88:ASP:O	14:AQ:89:ARG:HB3	1.81	0.79
20:AU:94:LYS:CD	20:AU:101:LYS:HZ3	1.95	0.79
31:BA:794:A:C2	31:BA:795:C:N3	2.49	0.79
54:CA:486:U:H2'	54:CA:487:A:H8	1.46	0.79
16:D1:92:ARG:HB3	17:D2:11:GLN:HE22	1.44	0.79
6:DG:112:PRO:HB3	26:D4:37:SER:N	1.97	0.79
58:DL:25:PRO:CA	58:DL:27:LEU:HG	2.11	0.79
57:DY:5:ARG:O	57:DY:7:VAL:HG12	1.82	0.79
16:A1:66:ASN:O	16:A1:70:ARG:HG2	1.81	0.79
1:AA:999:U:C2'	1:AA:1000:A:H5''	2.12	0.79
1:AA:1372:U:O5'	1:AA:1372:U:C6	2.36	0.79
6:AG:16:ARG:HH11	6:AG:16:ARG:HG2	1.47	0.79
21:AV:144:LEU:CD1	21:AV:144:LEU:O	2.30	0.79
25:AX:52:HIS:CD2	25:AX:52:HIS:H	1.99	0.79
31:BA:1002:G:H2'	31:BA:1003:G:C8	2.18	0.79
31:BA:1343:G:H2'	31:BA:1344:C:C6	2.16	0.79
31:BA:1374:A:H2'	31:BA:1375:A:H5'	1.64	0.79
43:BP:27:LYS:HE2	43:BP:31:LYS:HE3	1.62	0.79
27:D5:20:ARG:HA	27:D5:23:HIS:ND1	1.96	0.79
55:DA:1044:G:C2'	55:DA:1045:A:H5''	2.12	0.79
6:DG:145:THR:O	6:DG:146:TYR:HB3	1.81	0.79
8:DK:64:GLU:HG3	8:DK:67:ARG:NH2	1.97	0.79
25:DX:38:GLU:HB3	25:DX:40:THR:HG23	1.62	0.79
57:DY:50:ARG:CG	57:DY:51:LEU:N	2.36	0.79
1:AA:893:C:H2'	1:AA:894:C:C5	2.16	0.79
1:AA:893:C:H2'	1:AA:894:C:H6	1.43	0.79
31:BA:1072:G:H2'	31:BA:1073:U:C6	2.16	0.79
54:CA:1004:A:P	54:CA:1025:U:O4	2.40	0.79
34:CG:30:LYS:C	34:CG:32:ALA:H	1.86	0.79
55:DA:1301:A:C2'	55:DA:1302:A:H3'	2.12	0.79
55:DA:2068:U:H3	55:DA:2430:A:H2	1.29	0.79
56:DI:24:ILE:CA	56:DI:27:LEU:HD13	2.12	0.79
57:DY:71:LEU:CB	57:DY:113:GLN:CB	2.26	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:89:ALA:N	57:DY:92:THR:HB	1.97	0.79
1:AA:1149:G:H2'	1:AA:1150:C:C6	2.17	0.79
1:AA:1405:U:H2'	1:AA:1406:U:C6	2.18	0.79
1:AA:1827:C:C2'	1:AA:1828:G:H5'	2.13	0.79
7:AH:89:ILE:HD11	7:AH:129:THR:HB	1.65	0.79
12:AP:57:HIS:HE1	12:AP:113:GLN:HE21	1.27	0.79
15:AR:16:ARG:HH21	15:AR:19:LEU:CD2	1.93	0.79
31:BA:585:G:H4'	42:BO:8:ASN:HD21	1.48	0.79
40:BM:22:LYS:HD2	40:BM:26:ALA:HB2	1.63	0.79
49:BV:19:VAL:CG1	49:BV:44:MET:HB3	2.11	0.79
54:CA:141:A:H1'	54:CA:182:U:O2	1.83	0.79
54:CA:971:G:H5''	54:CA:972:C:H5''	1.64	0.79
52:CD:71:G:H1'	55:DA:1851:U:O2'	1.81	0.79
32:CE:47:THR:O	32:CE:51:LEU:HG	1.82	0.79
16:D1:108:GLU:HG3	17:D2:44:LYS:HE3	1.64	0.79
55:DA:2044:C:H6	55:DA:2044:C:H5'	1.47	0.79
3:DD:108:PRO:HB3	3:DD:143:HIS:CE1	2.17	0.79
6:DG:60:LEU:O	6:DG:64:THR:HG22	1.82	0.79
58:DL:86:LYS:CE	58:DL:86:LYS:HA	2.10	0.79
57:DY:48:GLY:C	57:DY:84:GLU:HB2	2.02	0.79
57:DY:49:ALA:C	57:DY:83:TYR:HD1	1.86	0.79
16:A1:58:ARG:HA	16:A1:61:TRP:CE3	2.18	0.79
1:AA:1543:A:H1'	1:AA:1545:A:O4'	1.82	0.79
1:AA:1885:A:H5'	1:AA:1886:C:OP2	1.82	0.79
1:AA:2447:G:O2'	1:AA:2448:A:OP2	2.00	0.79
1:AA:2557:G:H2'	1:AA:2558:C:H6	1.48	0.79
1:AA:959:A:N6	12:AP:82:ARG:HH22	1.79	0.79
4:AE:53:PRO:HG2	4:AE:54:GLN:H	1.48	0.79
31:BA:1513:A:H2'	31:BA:1514:C:C6	2.18	0.79
31:BA:274:A:O2'	31:BA:275:G:C8	2.34	0.79
54:CA:629:G:C5'	54:CA:630:G:OP2	2.30	0.79
39:CL:47:LEU:HD22	39:CL:47:LEU:H	1.47	0.79
55:DA:141:A:H8	55:DA:1595:G:H21	1.29	0.79
4:DE:95:ILE:H	4:DE:95:ILE:HD12	1.47	0.79
7:DH:98:LEU:HD12	7:DH:102:ALA:O	1.82	0.79
56:DJ:11:GLU:CA	56:DJ:17:VAL:HG11	2.12	0.79
58:DL:7:VAL:HG11	58:DL:58:THR:H	1.45	0.79
57:DY:16:ASN:HD21	57:DY:25:PHE:HZ	0.90	0.79
13:A0:49:ASP:OD1	13:A0:95:THR:HG22	1.82	0.79
1:AA:1496:A:H8	1:AA:1577:C:HO2'	1.26	0.79
1:AA:1947:C:H2'	1:AA:1948:G:H5''	1.62	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2749:A:H62	1:AA:2750:A:N6	1.81	0.79
1:AA:2758:A:C2'	1:AA:2759:G:H5''	2.12	0.79
9:AM:112:LEU:HD23	9:AM:113:GLY:N	1.96	0.79
12:AP:22:LYS:HD2	12:AP:22:LYS:N	1.98	0.79
52:BB:74:C:C2'	52:BB:75:C:H5'	2.13	0.79
32:BE:82:ARG:HD2	32:BE:92:TYR:HE1	1.46	0.79
39:BL:77:ILE:O	39:BL:81:ILE:HG12	1.82	0.79
54:CA:353:A:H5'	54:CA:353:A:H8	1.47	0.79
38:CK:60:ARG:HG3	38:CK:60:ARG:HH11	1.47	0.79
40:CM:34:VAL:HG22	40:CM:74:ILE:HG22	1.65	0.79
55:DA:518:G:H4'	18:DS:18:ARG:NH1	1.98	0.79
8:DK:128:LEU:HD13	8:DK:129:THR:N	1.97	0.79
58:DL:21:PRO:HB2	58:DL:22:PRO:CD	2.12	0.79
21:DV:116:VAL:HG12	21:DV:118:GLN:HG2	1.65	0.79
57:DY:50:ARG:N	57:DY:83:TYR:CD1	2.50	0.79
30:A8:48:PHE:O	30:A8:49:VAL:CG2	2.30	0.79
4:AE:116:VAL:O	4:AE:117:MET:HB3	1.80	0.79
8:AK:97:ILE:O	8:AK:101:LEU:HD23	1.83	0.79
9:AM:111:PRO:HA	9:AM:114:ARG:NH1	1.97	0.79
12:AP:38:GLU:HB2	12:AP:127:ILE:HG23	1.64	0.79
12:AP:127:ILE:HG22	12:AP:128:LYS:H	1.46	0.79
2:AB:50:G:OP1	14:AQ:63:THR:HG23	1.82	0.79
21:AV:115:GLY:CA	21:AV:177:PRO:CG	2.34	0.79
31:BA:1028:C:C2'	31:BA:1028(A):C:H5''	2.12	0.79
31:BA:1128:C:H2'	31:BA:1139:G:O6	1.83	0.79
52:BB:2:C:H2'	52:BB:3:C:C6	2.18	0.79
54:CA:1135:U:H4'	54:CA:1136:U:H5	1.47	0.79
32:CE:7:VAL:HG21	32:CE:217:ARG:CZ	2.12	0.79
43:CP:125:ARG:HD3	43:CP:126:LYS:H	1.46	0.79
16:D1:88:ILE:HD13	16:D1:88:ILE:H	1.48	0.79
55:DA:1174:A:N7	55:DA:1175:U:H1'	1.97	0.79
55:DA:1209:G:H21	55:DA:1210:A:H62	1.31	0.79
55:DA:1251:C:HO2'	55:DA:1252:G:H3'	1.46	0.79
55:DA:1510:A:OP1	55:DA:1511:A:H5''	1.83	0.79
4:DE:7:VAL:HG22	4:DE:27:LEU:HD23	1.63	0.79
8:DK:40:THR:HG22	8:DK:42:SER:H	1.48	0.79
58:DL:50:ASP:CG	58:DL:51:ALA:H	1.86	0.79
21:DV:127:LYS:O	21:DV:161:VAL:HB	1.82	0.79
21:DV:170:THR:O	21:DV:171:ILE:HB	1.83	0.79
57:DY:40:LEU:HD21	57:DY:50:ARG:NH1	1.95	0.79
7:AH:4:ILE:HB	7:AH:6:ARG:CZ	2.12	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:AK:101:LEU:HG	8:AK:109:ILE:HD12	1.64	0.79
11:AO:75:ILE:N	11:AO:75:ILE:HD13	1.98	0.79
21:AV:175:VAL:CG1	21:AV:176:PRO:N	2.43	0.79
21:AV:5:LEU:HD23	21:AV:47:VAL:HG21	1.64	0.79
31:BA:652:U:H1'	31:BA:653:A:C2	2.17	0.79
54:CA:1296:C:H5'	54:CA:1297:C:OP2	1.83	0.79
54:CA:429:U:H1'	54:CA:430:A:H5''	1.64	0.79
54:CA:1152:A:H5''	40:CM:13:HIS:HD2	1.48	0.79
54:CA:973:G:H1'	40:CM:55:LYS:CE	2.12	0.79
40:CM:46:ARG:HG2	40:CM:64:GLU:HB3	1.63	0.79
55:DA:654(Q):C:H2'	55:DA:654(R):C:C6	2.17	0.79
21:DV:119:GLU:HG3	21:DV:119:GLU:O	1.82	0.79
21:DV:154:ASP:O	21:DV:155:LEU:O	2.01	0.79
1:AA:1019:U:H3	1:AA:1142(A):A:H62	1.30	0.79
1:AA:1885:A:H3'	1:AA:1886:C:H6	1.48	0.79
1:AA:242:G:H5''	30:A8:62:LEU:CD1	2.08	0.79
3:AD:176:ARG:HG2	3:AD:176:ARG:HH11	1.47	0.79
1:AA:2729:G:H1'	4:AE:187:ALA:HB2	1.63	0.79
10:AN:113:LYS:O	10:AN:117:LEU:HD23	1.83	0.79
31:BA:653:A:H1'	38:BK:56:LYS:HE2	1.64	0.79
31:BA:820:U:H4'	31:BA:821:G:OP2	1.83	0.79
31:BA:980:C:H5'	31:BA:981:U:OP2	1.82	0.79
52:BD:8:U:H2'	52:BD:13:C:H41	1.44	0.79
45:BR:87:ILE:HG22	45:BR:88:ARG:H	1.46	0.79
52:CC:36:A:H5'	52:CC:36:A:H8	1.48	0.79
52:CD:45:U:H5'	52:CD:46:G:OP1	1.83	0.79
49:CV:81:ARG:CG	49:CV:82:GLY:N	2.46	0.79
16:D1:92:ARG:HH11	16:D1:95:LEU:CD1	1.94	0.79
55:DA:1360:A:N6	55:DA:1372:U:O4	2.16	0.79
55:DA:363(F):A:H4'	55:DA:364:C:O5'	1.82	0.79
3:DD:35:LYS:HB3	3:DD:63:ARG:HA	1.65	0.79
8:DK:74:ASN:HD22	8:DK:74:ASN:H	1.31	0.79
10:DN:23:ARG:HG3	10:DN:24:VAL:H	1.46	0.79
12:DP:60:ARG:CZ	21:DV:181:GLU:OE1	2.31	0.79
57:DY:127:GLU:CG	57:DY:128:LEU:N	2.45	0.79
16:A1:95:LEU:C	16:A1:97:ASP:N	2.35	0.79
1:AA:1899:G:H21	1:AA:1902:C:H5	1.31	0.79
1:AA:2392:A:H2	1:AA:2424:C:N4	1.79	0.79
1:AA:654(L):G:C2	1:AA:654(M):C:H1'	2.18	0.79
3:AD:131:LEU:HD13	3:AD:136:ILE:HD11	1.65	0.79
4:AE:176:ILE:HB	4:AE:181:LEU:HB2	1.65	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:AU:43:ASN:N	20:AU:43:ASN:HD22	1.81	0.79
31:BA:149:A:H2'	31:BA:150:C:H6	1.48	0.79
34:BG:31:CYS:C	34:BG:33:MET:H	1.84	0.79
49:BV:42:PRO:HA	49:BV:45:VAL:HG13	1.64	0.79
43:BP:90:LEU:HD13	49:BV:78:ARG:HH21	1.48	0.79
34:CG:58:LEU:O	34:CG:62:GLN:HG2	1.83	0.79
17:D2:35:LEU:HB2	17:D2:37:VAL:CG2	2.13	0.79
7:DH:152:ARG:O	7:DH:153:LYS:HD2	1.83	0.79
58:DL:109:LYS:HA	58:DL:120:LEU:HD21	1.64	0.79
15:DR:6:LEU:HA	15:DR:9:LEU:HB2	1.64	0.79
57:DY:73:GLY:C	57:DY:119:ALA:HA	2.03	0.79
16:A1:79:PHE:HE2	16:A1:83:LEU:HD13	1.45	0.78
30:A8:16:ILE:HB	30:A8:65:GLU:HA	1.65	0.78
1:AA:1869:G:H5'	1:AA:1870:C:OP2	1.82	0.78
1:AA:1967:C:H2'	1:AA:1968:G:H5'	1.64	0.78
8:AK:141:LYS:O	8:AK:142:VAL:HB	1.82	0.78
31:BA:1014:A:C2	31:BA:1219:U:H1'	2.16	0.78
43:BP:83:ASP:OD1	43:BP:84:ILE:HD13	1.83	0.78
51:BX:18:TYR:HB3	51:BX:22:ARG:O	1.83	0.78
34:CG:28:SER:HB3	34:CG:29:PRO:HD2	1.63	0.78
54:CA:452:A:H4'	46:CS:72:ARG:NH2	1.97	0.78
55:DA:1598:C:H5'	19:DT:36:LYS:HB2	1.64	0.78
55:DA:2475:C:H2'	55:DA:2475:C:O2	1.83	0.78
55:DA:33:U:O4	55:DA:446:G:O2'	2.01	0.78
55:DA:654(K):C:H2'	55:DA:654(L):G:H8	1.48	0.78
3:DD:142:VAL:HG23	3:DD:193:VAL:HA	1.64	0.78
14:DQ:5:THR:OG1	14:DQ:8:GLU:HG3	1.83	0.78
21:DV:111:VAL:CG2	21:DV:146:ILE:H	1.95	0.78
57:DY:136:ALA:O	57:DY:139:VAL:HB	1.83	0.78
57:DY:84:GLU:HA	57:DY:84:GLU:OE1	1.80	0.78
17:A2:69:LYS:HD3	17:A2:85:LYS:CD	2.13	0.78
22:A3:24:LYS:O	22:A3:25:ARG:HD2	1.83	0.78
1:AA:273(F):C:H3'	1:AA:274:G:H5''	1.65	0.78
1:AA:458:G:C2'	1:AA:459:U:OP2	2.31	0.78
1:AA:881:G:H2'	52:BB:19:G:C6	2.18	0.78
8:AK:6:LEU:HD13	8:AK:36:ALA:HA	1.64	0.78
12:AP:11:LYS:HE2	12:AP:85:LYS:HG2	1.64	0.78
31:BA:632:A:O2'	31:BA:633:G:OP2	2.02	0.78
31:BA:794:A:H2'	31:BA:795:C:C6	2.18	0.78
43:BP:79:LYS:HD3	43:BP:79:LYS:O	1.83	0.78
54:CA:197:A:H4'	54:CA:198:G:O5'	1.83	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:AK:82:ARG:HE	54:CA:56:U:H4'	1.48	0.78
54:CA:920:U:H2'	54:CA:921:U:C6	2.18	0.78
38:CK:102:ARG:NH1	38:CK:105:ARG:HH22	1.78	0.78
48:CU:54:ARG:HH11	48:CU:54:ARG:HB3	1.47	0.78
16:D1:92:ARG:HB3	17:D2:11:GLN:NE2	1.99	0.78
28:D6:27:LYS:HB2	28:D6:27:LYS:NZ	1.98	0.78
55:DA:1699:G:O3'	55:DA:1700:A:H4'	1.84	0.78
8:DK:61:ARG:HA	8:DK:61:ARG:NE	1.97	0.78
58:DL:52:ILE:CD1	58:DL:76:TYR:N	2.46	0.78
58:DL:7:VAL:HG11	58:DL:58:THR:CA	2.12	0.78
58:DL:83:GLY:H	58:DL:99:ILE:HG21	1.48	0.78
11:DO:75:ILE:N	11:DO:75:ILE:HD13	1.96	0.78
21:DV:61:LEU:HD11	21:DV:65:GLN:CG	2.12	0.78
57:DY:32:LEU:CB	57:DY:33:PRO:HD3	2.14	0.78
1:AA:1731:G:H2'	1:AA:1732:A:H5'	1.63	0.78
1:AA:1947:C:C2'	1:AA:1948:G:H5''	2.13	0.78
1:AA:27:G:O2'	1:AA:28:A:H8	1.66	0.78
1:AA:2893:G:H5'	1:AA:2894:G:C5'	2.03	0.78
4:AE:23:VAL:HA	4:AE:184:VAL:O	1.83	0.78
6:AG:135:LEU:HD23	6:AG:140:ILE:HD11	1.64	0.78
23:AZ:92:LYS:NZ	23:AZ:92:LYS:HB3	1.98	0.78
31:BA:737:A:H2'	31:BA:738:C:C6	2.18	0.78
31:BA:244:U:O4	31:BA:906:G:H1'	1.84	0.78
44:BQ:29:ARG:HG2	44:BQ:40:CYS:HB2	1.65	0.78
55:DA:2392:A:H2	55:DA:2424:C:H42	1.32	0.78
26:A4:53:GLU:HG3	26:A4:54:GLY:N	1.98	0.78
29:A7:47:ARG:N	29:A7:47:ARG:HH11	1.81	0.78
1:AA:242:G:C5'	30:A8:62:LEU:HD13	2.07	0.78
1:AA:1171:G:O2'	1:AA:1173:G:O5'	2.01	0.78
1:AA:895:U:C2'	1:AA:895:U:O2	2.31	0.78
1:AA:946:G:N2	1:AA:971:C:O2	2.15	0.78
8:AK:79:ILE:HB	8:AK:142:VAL:CG1	2.06	0.78
15:AR:55:ASN:H	15:AR:59:THR:HB	1.46	0.78
21:AV:140:ASP:O	21:AV:141:VAL:HB	1.84	0.78
52:BD:23:A:H2'	52:BD:24:G:H8	1.48	0.78
54:CA:1443:G:H3'	54:CA:1446:A:H5''	1.64	0.78
40:CM:48:THR:CA	40:CM:62:HIS:HB3	2.14	0.78
55:DA:1044:G:H2'	55:DA:1045:A:H5''	1.65	0.78
55:DA:1083:U:H4'	57:DY:41:ARG:HD3	1.65	0.78
5:DF:66:PRO:O	5:DF:67:GLN:CB	2.31	0.78
56:DI:1:MET:O	56:DI:5:ILE:HD13	1.84	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:DL:10:LEU:O	58:DL:23:VAL:HG13	1.82	0.78
10:DN:75:SER:HB2	15:DR:74:ARG:HH12	1.47	0.78
21:DV:10:ARG:HH21	21:DV:26:GLY:H	1.30	0.78
21:DV:111:VAL:HG23	21:DV:146:ILE:H	1.46	0.78
57:DY:72:ASP:O	57:DY:112:LEU:HD21	1.83	0.78
1:AA:1312:U:HO2'	1:AA:1314:C:H5	1.30	0.78
1:AA:951:C:O2'	1:AA:952:G:H5'	1.82	0.78
6:AG:61:ALA:HB2	6:AG:68:PRO:HD3	1.65	0.78
37:BJ:12:LEU:H	37:BJ:12:LEU:HD12	1.48	0.78
39:BL:46:ALA:O	39:BL:49:PRO:HD2	1.84	0.78
22:D3:11:ARG:HB2	22:D3:11:ARG:HH11	1.47	0.78
55:DA:1826:G:H4'	3:DD:242:ARG:NH2	1.99	0.78
55:DA:205:G:O2'	55:DA:206:U:OP2	2.02	0.78
55:DA:2092:U:H4'	55:DA:2093:G:O5'	1.83	0.78
55:DA:2159:G:H2'	55:DA:2160:G:H8	1.45	0.78
55:DA:747:U:O2	55:DA:2014:A:H1'	1.84	0.78
5:DF:31:HIS:HB2	11:DO:9:ASN:HD21	1.47	0.78
56:DI:3:LEU:HD23	56:DI:4:ASP:N	1.97	0.78
58:DL:86:LYS:O	58:DL:88:ALA:N	2.16	0.78
21:DV:150:LEU:C	21:DV:151:HIS:ND1	2.37	0.78
21:DV:191:VAL:CG2	21:DV:197:ILE:HG12	2.14	0.78
57:DY:136:ALA:HA	56:DJ:6:GLU:OE2	1.82	0.78
55:DA:1084:A:OP1	57:DY:50:ARG:HD2	1.83	0.78
17:A2:22:VAL:HG22	17:A2:23:GLU:H	1.49	0.78
1:AA:1378:A:O2'	1:AA:1379:A:C5'	2.30	0.78
1:AA:1496:A:H8	1:AA:1577:C:O2'	1.66	0.78
1:AA:2134:A:H62	1:AA:2157:G:H1'	1.49	0.78
9:AM:120:LEU:HD21	9:AM:122:VAL:HG23	1.64	0.78
9:AM:55:VAL:HB	9:AM:126:PRO:HB3	1.65	0.78
20:AU:20:TYR:O	20:AU:22:GLY:N	2.15	0.78
21:AV:144:LEU:CG	21:AV:144:LEU:O	2.30	0.78
31:BA:186(B):C:H2'	31:BA:186(C):G:C8	2.19	0.78
31:BA:696:A:C2'	31:BA:697:U:H5''	2.12	0.78
52:BC:43:C:H6	52:BC:43:C:H5'	1.48	0.78
34:BG:61:LYS:NZ	34:BG:62:GLN:HE21	1.81	0.78
54:CA:1211:U:H5'	54:CA:1212:U:OP1	1.82	0.78
32:CE:178:ARG:HH11	32:CE:178:ARG:CB	1.95	0.78
54:CA:1128:C:H4'	39:CL:16:ARG:HH12	1.48	0.78
42:CO:20:LYS:H	42:CO:20:LYS:HD3	1.48	0.78
6:DG:6:ALA:HB3	26:D4:23:GLU:HG3	1.65	0.78
10:DN:34:THR:HG22	10:DN:37:ASP:OD2	1.84	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:DO:65:ARG:HB2	11:DO:65:ARG:HH11	1.47	0.78
15:DR:107:ASP:O	15:DR:110:ILE:HG22	1.83	0.78
30:A8:62:LEU:CB	30:A8:63:PRO:HD3	2.14	0.78
52:BD:8:U:H4'	52:BD:9:A:OP1	1.83	0.78
32:BE:88:ALA:HA	32:BE:226:ARG:HH12	1.48	0.78
42:BO:41:ARG:NH1	42:BO:41:ARG:HB3	1.98	0.78
36:CI:97:PHE:HD2	48:CU:31:LEU:HD21	1.49	0.78
49:CV:2:PRO:O	49:CV:3:ARG:O	2.01	0.78
49:CV:41:VAL:HA	49:CV:44:MET:HG3	1.66	0.78
13:D0:117:VAL:HG22	13:D0:118:GLU:N	1.94	0.78
26:D4:16:CYS:SG	26:D4:18:CYS:N	2.56	0.78
55:DA:649:G:H2'	55:DA:650:C:C6	2.18	0.78
55:DA:897:C:OP1	55:DA:897:C:C5	2.36	0.78
8:DK:133:HIS:HB2	8:DK:134:PRO:HD2	1.66	0.78
57:DY:135:ARG:HH21	56:DJ:18:LEU:HD13	1.49	0.78
13:A0:74:LYS:HE2	13:A0:77:ARG:HH21	1.48	0.78
12:AP:4:PRO:HB2	12:AP:10:ARG:HH22	1.48	0.78
31:BA:1302:U:H5''	31:BA:1303:C:OP2	1.84	0.78
31:BA:411:A:C5	31:BA:413:G:H1'	2.19	0.78
34:BG:139:ARG:HH11	34:BG:139:ARG:HG3	1.48	0.78
49:BV:62:ILE:HA	49:BV:66:MET:HE1	1.65	0.78
54:CA:60:A:H4'	54:CA:61:G:O5'	1.84	0.78
32:CE:14:GLY:O	32:CE:15:VAL:HG13	1.83	0.78
22:D3:32:ARG:N	22:D3:35:ASN:HD21	1.76	0.78
55:DA:1434:A:H61	55:DA:1558:A:H62	1.27	0.78
55:DA:654(C):G:H3'	55:DA:654(D):G:H8	1.49	0.78
56:DI:24:ILE:HA	56:DI:27:LEU:CD1	2.14	0.78
55:DA:2562:U:H1'	10:DN:23:ARG:NH1	1.96	0.78
10:DN:68:GLU:HB3	10:DN:78:ARG:HH11	1.48	0.78
57:DY:27:VAL:CG2	57:DY:80:VAL:HG11	2.14	0.78
28:A6:36:LEU:HA	28:A6:50:ARG:HB3	1.65	0.78
1:AA:1024:G:C3'	1:AA:1025:G:H5''	2.13	0.78
1:AA:1944:U:H1'	1:AA:1955:U:O4'	1.83	0.78
1:AA:607:U:O4	1:AA:608:A:N7	2.16	0.78
1:AA:2820:A:C6	4:AE:191:PRO:HB2	2.18	0.78
20:AU:47:LYS:HA	20:AU:60:PHE:HB3	1.64	0.78
31:BA:251:G:N1	31:BA:266:G:O6	2.17	0.78
31:BA:503:C:OP2	42:BO:116:SER:HB3	1.84	0.78
37:BJ:100:ALA:O	37:BJ:104:LEU:HD23	1.84	0.78
54:CA:1321:C:H5''	54:CA:1322:C:H5''	1.65	0.78
54:CA:644:G:H4'	38:CK:92:ARG:HH21	1.49	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:CE:18:GLY:N	32:CE:42:ILE:HG22	1.98	0.78
13:D0:96:ARG:NH2	13:D0:117:VAL:HG23	1.99	0.78
55:DA:1846:G:H5'	55:DA:1847:A:OP2	1.84	0.78
55:DA:2760:C:H2'	55:DA:2761:G:C5'	2.13	0.78
55:DA:2898:U:H2'	55:DA:2899:G:H8	1.49	0.78
55:DA:860:U:C5	55:DA:917:A:C2	2.72	0.78
58:DL:54:PRO:HD2	58:DL:72:PRO:CA	2.13	0.78
11:DO:106:LEU:O	11:DO:107:LYS:HB2	1.83	0.78
21:DV:192:ALA:C	21:DV:194:PRO:CD	2.51	0.78
57:DY:11:ALA:HB1	57:DY:52:PHE:HE1	1.49	0.78
13:A0:38:VAL:HB	13:A0:39:PRO:HD3	1.65	0.78
1:AA:2306:C:H3'	1:AA:2307:G:C5'	2.13	0.78
6:AG:115:ARG:NH1	6:AG:115:ARG:HB3	1.99	0.78
7:AH:92:ILE:HD13	7:AH:160:LYS:HD3	1.66	0.78
8:AK:88:ILE:HG22	8:AK:89:TYR:N	1.96	0.78
12:AP:106:VAL:HG22	12:AP:118:LEU:HD21	1.65	0.78
14:AQ:5:THR:OG1	14:AQ:8:GLU:HG3	1.83	0.78
31:BA:1281:U:O2'	31:BA:1282:C:OP1	2.02	0.78
31:BA:734:G:H21	48:BU:75:ILE:HD11	1.48	0.78
50:CW:43:LEU:HA	50:CW:46:GLU:HB3	1.65	0.78
28:D6:20:ASN:CG	28:D6:21:TYR:H	1.87	0.78
55:DA:5:A:O2'	55:DA:6:A:H5'	1.83	0.78
8:DK:11:ASN:O	8:DK:12:LEU:HB2	1.82	0.78
8:DK:76:THR:HG23	8:DK:139:GLN:NE2	1.99	0.78
8:DK:99:GLU:HG2	8:DK:103:ARG:NH2	1.98	0.78
58:DL:19:PRO:HD3	58:DL:38:VAL:HG11	1.66	0.78
1:AA:637:A:OP2	11:AO:115:LEU:HD22	1.83	0.77
3:AD:239:ARG:O	3:AD:240:ALA:HB2	1.83	0.77
4:AE:8:LYS:HG2	4:AE:192:ASN:ND2	1.98	0.77
11:AO:64:LYS:C	11:AO:66:GLY:H	1.85	0.77
11:AO:71:VAL:CG1	11:AO:72:PRO:HD3	2.13	0.77
52:BD:8:U:H2'	52:BD:13:C:N4	1.98	0.77
33:BF:86:VAL:O	33:BF:90:GLU:HG2	1.84	0.77
36:BI:79:LEU:O	36:BI:85:VAL:HG11	1.83	0.77
36:BI:91:VAL:HG11	48:BU:72:ARG:NH1	1.99	0.77
37:BJ:84:ASN:HB2	52:BD:37:MIA:H163	1.65	0.77
54:CA:438:G:H4'	34:CG:123:HIS:CE1	2.18	0.77
54:CA:792:A:C2'	54:CA:794:A:N6	2.28	0.77
34:CG:129:ASN:CA	34:CG:145:GLU:HB2	2.14	0.77
49:CV:26:GLY:O	49:CV:27:GLU:HG3	1.84	0.77
55:DA:803:U:H5'	55:DA:803:U:C6	2.18	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1077:A:H4'	58:DL:93:ARG:NH2	1.98	0.77
21:DV:191:VAL:HG12	21:DV:197:ILE:HG23	1.63	0.77
57:DY:138:LEU:CD2	56:DJ:22:GLN:OE1	2.31	0.77
1:AA:1045:A:C2'	1:AA:1046:A:H5''	2.14	0.77
1:AA:1827:C:H2'	1:AA:1828:G:H5'	1.66	0.77
1:AA:2406:U:H5''	1:AA:2408:U:OP2	1.84	0.77
1:AA:566:U:OP1	11:AO:29:LYS:HE2	1.85	0.77
1:AA:196:A:H2'	1:AA:805:G:O6	1.83	0.77
4:AE:201:THR:O	4:AE:202:LYS:HD3	1.84	0.77
5:AF:132:VAL:HG22	5:AF:133:ASN:H	1.49	0.77
11:AO:106:LEU:HD11	11:AO:112:LEU:HD23	1.65	0.77
31:BA:1529:G:H5'	31:BA:1530:G:OP2	1.83	0.77
33:BF:77:ILE:HA	33:BF:84:ILE:HB	1.65	0.77
54:CA:1053:G:C5'	54:CA:1054:C:H5'	2.14	0.77
54:CA:1363:A:H4'	54:CA:1364:U:C5'	2.15	0.77
54:CA:255:G:H4'	47:CT:17:LYS:HD3	1.65	0.77
54:CA:452:A:H62	54:CA:480:U:H3	1.30	0.77
16:D1:69:CYS:HB2	16:D1:74:LEU:HD11	1.65	0.77
28:D6:11:LEU:HD23	28:D6:26:ASN:HB3	1.65	0.77
30:D8:23:VAL:CG1	30:D8:46:ARG:HB3	2.13	0.77
55:DA:2113:U:H5'	55:DA:2114:A:C8	2.19	0.77
55:DA:2340:G:O2'	55:DA:2341:G:H5'	1.83	0.77
1:AA:1045:A:H2'	1:AA:1046:A:H5''	1.66	0.77
1:AA:1047:G:H2'	1:AA:1110:G:H22	1.48	0.77
1:AA:1434:A:H61	1:AA:1558:A:N6	1.80	0.77
31:BA:818:G:H3'	31:BA:819:A:H5'	1.66	0.77
54:CA:438:G:H4'	34:CG:123:HIS:ND1	1.98	0.77
38:CK:69:ARG:HD3	38:CK:75:ARG:O	1.85	0.77
39:CL:13:ALA:HB2	39:CL:68:GLY:HA3	1.67	0.77
49:CV:41:VAL:CB	49:CV:42:PRO:HA	2.09	0.77
17:D2:62:LEU:HD11	17:D2:95:LEU:HB2	1.66	0.77
55:DA:1169:G:C2'	55:DA:1170:G:H5''	2.13	0.77
55:DA:2159:G:H2'	55:DA:2160:G:C8	2.19	0.77
56:DI:21:LYS:O	56:DI:22:GLN:C	2.21	0.77
57:DY:24:PHE:CD2	57:DY:25:PHE:N	2.53	0.77
1:AA:2557:G:H2'	1:AA:2558:C:C6	2.18	0.77
4:AE:200:GLU:HG2	4:AE:201:THR:N	1.98	0.77
31:BA:792:A:C8	31:BA:794:A:N6	2.52	0.77
40:CM:54:PHE:O	40:CM:55:LYS:HG3	1.83	0.77
40:CM:49:VAL:O	40:CM:60:ARG:HB3	1.84	0.77
16:D1:66:ASN:O	16:D1:70:ARG:HB2	1.84	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2055:C:H5'	55:DA:2056:G:OP1	1.85	0.77
55:DA:2061:G:H5''	55:DA:2503:A:N1	1.99	0.77
2:DB:52:A:H62	14:DQ:33:LYS:HG3	1.49	0.77
21:DV:177:PRO:O	21:DV:178:GLU:HB3	1.82	0.77
21:DV:186:GLU:O	21:DV:187:ALA:HB3	1.85	0.77
21:DV:194:PRO:CG	21:DV:196:VAL:CG1	2.60	0.77
57:DY:72:ASP:C	57:DY:112:LEU:HG	2.05	0.77
23:DZ:94:LEU:O	23:DZ:94:LEU:HD23	1.85	0.77
1:AA:1061:U:H4'	1:AA:1070:A:O2'	1.85	0.77
1:AA:1252:G:O2'	1:AA:1253:A:O4'	2.00	0.77
1:AA:1673:U:H2'	1:AA:1674:G:H5'	1.64	0.77
2:AB:87:G:H3'	2:AB:88:C:C5'	2.13	0.77
20:AU:12:THR:CG2	20:AU:26:LYS:HE2	2.14	0.77
20:AU:50:ARG:HB3	20:AU:53:PRO:HG3	1.64	0.77
31:BA:1006:C:H2'	31:BA:1007:C:C6	2.19	0.77
31:BA:827:U:H3	31:BA:872:A:H62	1.33	0.77
40:BM:4:ILE:HA	40:BM:100:THR:HG22	1.67	0.77
54:CA:1322:C:O2'	54:CA:1323:G:H5'	1.85	0.77
46:CS:21:VAL:HG23	46:CS:33:ILE:HB	1.64	0.77
55:DA:1359:A:C4'	55:DA:1359:A:C8	2.67	0.77
55:DA:2795:G:H3'	55:DA:2797:U:H5''	1.64	0.77
55:DA:671:C:O2'	55:DA:672:C:H5'	1.84	0.77
56:DJ:24:ILE:HG22	56:DJ:25:ASP:N	1.98	0.77
10:DN:4:PRO:O	10:DN:5:GLN:HB2	1.84	0.77
15:DR:60:THR:HG22	15:DR:77:PRO:HA	1.66	0.77
21:DV:114:GLY:HA2	21:DV:179:ASP:OD1	1.84	0.77
57:DY:132:ASP:OD2	56:DJ:10:GLU:OE2	2.03	0.77
1:AA:2394:C:H42	52:BD:76:A:H8	1.31	0.77
1:AA:2426:A:H4'	1:AA:2427:C:OP2	1.85	0.77
1:AA:922:U:H2'	1:AA:923:C:C6	2.18	0.77
1:AA:329:G:O6	20:AU:19:LYS:HG2	1.85	0.77
54:CA:826:C:H5'	38:CK:12:ARG:HH21	1.49	0.77
46:CS:45:THR:HG22	46:CS:47:ASP:N	1.99	0.77
55:DA:1066:U:H2'	55:DA:1068:G:OP2	1.84	0.77
55:DA:1179:C:H2'	55:DA:1180:C:C5'	2.15	0.77
55:DA:2645:G:H4'	55:DA:2732:G:O2'	1.84	0.77
55:DA:2752:C:H6	55:DA:2752:C:OP2	1.66	0.77
55:DA:2854:G:H2'	55:DA:2855:C:H6	1.50	0.77
55:DA:582:G:H2'	55:DA:583:G:H8	1.50	0.77
58:DL:77:LEU:H	58:DL:77:LEU:HD12	1.49	0.77
15:DR:3:ARG:HG3	15:DR:7:ILE:HG12	1.67	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:DV:128:VAL:HG22	21:DV:129:SER:H	1.50	0.77
17:A2:22:VAL:HG22	17:A2:23:GLU:N	2.00	0.77
6:AG:104:GLU:HG2	26:A4:23:GLU:CG	2.15	0.77
1:AA:481:G:H1'	1:AA:506:G:N2	1.98	0.77
1:AA:673:C:H5'	5:AF:54:ARG:NH1	2.00	0.77
10:AN:53:LYS:O	10:AN:56:ASP:HB2	1.83	0.77
21:AV:109:ALA:O	21:AV:145:GLU:HB2	1.85	0.77
32:BE:48:MET:HA	32:BE:51:LEU:HD12	1.67	0.77
43:BP:13:LYS:HA	43:BP:44:ARG:HH11	1.50	0.77
44:BQ:14:PRO:HG2	44:BQ:15:LYS:H	1.49	0.77
55:DA:1359:A:H5'	55:DA:1359:A:C8	2.19	0.77
4:DE:65:GLY:HA2	4:DE:70:ALA:HB3	1.67	0.77
58:DL:58:THR:HG21	58:DL:66:THR:OG1	1.85	0.77
11:DO:66:GLY:HA2	11:DO:68:GLN:HE22	1.50	0.77
20:DU:88:LYS:HB3	20:DU:90:LEU:HD23	1.64	0.77
57:DY:127:GLU:HG3	57:DY:128:LEU:N	1.98	0.77
57:DY:22:GLY:O	57:DY:23:SER:HB3	1.84	0.77
57:DY:25:PHE:C	57:DY:82:PHE:CZ	2.58	0.77
57:DY:49:ALA:N	57:DY:84:GLU:O	2.17	0.77
1:AA:2127:G:C3'	1:AA:2128:C:H5''	2.15	0.77
1:AA:2784:C:H4'	4:AE:41:LYS:O	1.85	0.77
1:AA:895:U:H4'	1:AA:896:A:C8	2.18	0.77
1:AA:946:G:C2'	1:AA:947:G:O5'	2.32	0.77
4:AE:78:LEU:N	4:AE:78:LEU:HD23	2.00	0.77
12:AP:42:ILE:CD1	12:AP:42:ILE:H	1.98	0.77
21:AV:120:ILE:O	21:AV:171:ILE:HA	1.84	0.77
21:AV:61:LEU:HB3	21:AV:62:PRO:CD	2.11	0.77
31:BA:1126:U:H3	31:BA:1281:U:C1'	1.98	0.77
32:BE:82:ARG:HD2	32:BE:92:TYR:CE1	2.20	0.77
34:BG:108:LEU:HB3	34:BG:110:PHE:CE1	2.19	0.77
54:CA:789:U:C4	54:CA:792:A:OP2	2.36	0.77
32:CE:33:TYR:HB2	32:CE:43:ASP:HB2	1.67	0.77
33:CF:11:ARG:HB3	33:CF:15:THR:HB	1.66	0.77
33:CF:91:LEU:HB2	33:CF:99:VAL:HG21	1.66	0.77
55:DA:2173:A:H3'	55:DA:2174:C:H6	1.50	0.77
55:DA:34:C:O2'	55:DA:35:G:P	2.43	0.77
55:DA:1111:A:H5'	7:DH:3:ARG:HH11	1.49	0.77
8:DK:72:LEU:HD11	8:DK:101:LEU:HD11	1.66	0.77
57:DY:87:VAL:O	57:DY:88:ALA:O	2.02	0.77
28:A6:41:PRO:HD3	28:A6:47:THR:HG22	1.67	0.77
1:AA:1140:C:H1'	1:AA:1143:A:N7	2.00	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2729:G:H1'	4:AE:187:ALA:CB	2.14	0.77
7:AH:35:VAL:HG11	7:AH:71:LEU:HG	1.66	0.77
1:AA:1341:U:C5'	19:AT:57:LEU:HD22	2.14	0.77
35:BH:76:ILE:HG23	35:BH:142:LEU:HD13	1.66	0.77
54:CA:537:G:H5''	42:CO:113:ARG:NH1	2.00	0.77
36:CI:99:ALA:HB1	48:CU:23:LYS:NZ	2.00	0.77
49:CV:31:ILE:CG2	49:CV:49:ILE:HA	2.14	0.77
50:CW:49:ALA:HB2	50:CW:99:LEU:HD23	1.67	0.77
13:D0:79:LEU:HA	13:D0:83:ILE:HG13	1.66	0.77
55:DA:1077:A:C2	58:DL:133:SER:HB3	2.20	0.77
55:DA:1403:C:H5''	55:DA:1471:A:H1'	1.67	0.77
2:DB:81:G:O6	2:DB:96:G:C5	2.37	0.77
3:DD:135:PHE:CD2	3:DD:135:PHE:N	2.51	0.77
56:DI:20:LEU:O	56:DI:24:ILE:CG2	2.32	0.77
19:DT:3:THR:HA	19:DT:6:ASP:OD2	1.83	0.77
20:DU:40:GLU:HA	20:DU:64:GLU:OE1	1.84	0.77
21:DV:197:ILE:N	21:DV:197:ILE:HD12	2.00	0.77
57:DY:15:GLU:O	57:DY:16:ASN:HB3	1.82	0.77
23:DZ:91:LYS:HG3	23:DZ:92:LYS:H	1.50	0.77
17:A2:78:LYS:O	17:A2:79:VAL:HG13	1.85	0.77
17:A2:80:GLN:CA	17:A2:80:GLN:NE2	2.30	0.77
30:A8:34:TRP:CD1	30:A8:35:GLN:N	2.53	0.77
11:AO:97:PRO:HD3	11:AO:126:VAL:O	1.85	0.77
21:AV:176:PRO:N	21:AV:177:PRO:CD	2.36	0.77
39:BL:65:VAL:CG2	39:BL:66:ARG:H	1.97	0.77
40:CM:4:ILE:HB	40:CM:74:ILE:HG12	1.67	0.77
43:CP:14:ARG:HG2	43:CP:16:ASP:OD2	1.84	0.77
46:CS:20:VAL:HG21	46:CS:32:TYR:CG	2.20	0.77
49:CV:7:LYS:CB	49:CV:7:LYS:NZ	2.45	0.77
55:DA:1359:A:H8	55:DA:1359:A:C5'	1.98	0.77
55:DA:2296:U:H4'	55:DA:2297:C:OP1	1.84	0.77
55:DA:2425:A:H4'	55:DA:2426:A:O5'	1.83	0.77
4:DE:67:PHE:O	4:DE:69:LYS:N	2.16	0.77
56:DI:9:LYS:O	56:DI:11:GLU:CA	2.32	0.77
58:DL:95:LYS:CB	58:DL:136:VAL:HG21	2.14	0.77
10:DN:2:ILE:HD11	10:DN:82:ASN:HD22	1.48	0.77
20:DU:42:VAL:HB	20:DU:67:LEU:HD11	1.67	0.77
27:A5:40:LYS:HZ2	27:A5:46:CYS:H	1.33	0.76
28:A6:31:PRO:HB2	28:A6:33:LYS:HG2	1.66	0.76
3:AD:70:TRP:CD1	3:AD:70:TRP:C	2.58	0.76
12:AP:29:PHE:HB2	12:AP:65:PHE:CE2	2.20	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:BU:22:VAL:C	48:BU:24:ALA:H	1.89	0.76
32:CE:132:LYS:HA	32:CE:135:GLN:HB2	1.66	0.76
32:CE:52:GLU:HG2	32:CE:56:ARG:HH12	1.50	0.76
40:CM:75:ILE:HG13	40:CM:76:ASN:H	1.50	0.76
50:CW:97:ALA:O	50:CW:99:LEU:HD22	1.83	0.76
26:D4:12:ALA:HB1	26:D4:30:GLU:H	1.49	0.76
26:D4:69:LYS:HD3	26:D4:70:GLY:H	1.51	0.76
55:DA:1026:U:O2'	55:DA:1027:A:C5'	2.31	0.76
55:DA:2789:C:H2'	55:DA:2790:A:H5''	1.66	0.76
55:DA:2887:U:H2'	55:DA:2888:C:C6	2.19	0.76
2:DB:20:C:H2'	2:DB:21:G:C5'	2.15	0.76
4:DE:50:GLY:CA	4:DE:77:ILE:HA	2.15	0.76
58:DL:52:ILE:HD12	58:DL:72:PRO:O	1.85	0.76
11:DO:38:GLN:H	11:DO:41:ARG:HG2	1.50	0.76
57:DY:28:ASN:CG	57:DY:83:TYR:HE2	1.88	0.76
1:AA:1460:A:H5''	1:AA:1461:G:OP2	1.84	0.76
1:AA:1944:U:H5''	1:AA:1945:G:OP2	1.84	0.76
1:AA:2518:A:H5''	1:AA:2519:U:OP2	1.85	0.76
1:AA:260:G:H1'	1:AA:621:A:C8	2.19	0.76
1:AA:2784:C:H1'	4:AE:37:ARG:HH22	1.48	0.76
21:AV:145:GLU:OE1	21:AV:145:GLU:CA	2.30	0.76
24:AW:50:ILE:CD1	24:AW:51:ARG:H	1.98	0.76
42:BO:8:ASN:O	42:BO:12:ARG:HG3	1.84	0.76
54:CA:1347:G:N2	54:CA:1373:G:H2'	2.01	0.76
54:CA:498:A:H4'	54:CA:500:G:OP1	1.85	0.76
33:CF:16:ARG:NH1	33:CF:16:ARG:HB2	2.00	0.76
34:CG:175:SER:HB2	34:CG:186:LEU:HD11	1.67	0.76
55:DA:1178:C:H2'	55:DA:1179:C:H6	1.50	0.76
2:DB:81:G:O6	2:DB:96:G:C6	2.37	0.76
58:DL:122:ALA:O	58:DL:126:MET:SD	2.43	0.76
58:DL:19:PRO:HA	58:DL:25:PRO:HD3	1.68	0.76
58:DL:64:SER:O	58:DL:65:PHE:HB3	1.85	0.76
21:DV:178:GLU:OE1	21:DV:180:VAL:HA	1.84	0.76
57:DY:122:VAL:HG12	57:DY:126:ALA:HB2	1.67	0.76
57:DY:139:VAL:HG23	56:DJ:6:GLU:OE2	1.84	0.76
1:AA:1225:C:O3'	17:A2:85:LYS:HB2	1.85	0.76
1:AA:1344:G:H4'	1:AA:1384:A:C5	2.20	0.76
1:AA:1267:U:O4	1:AA:2012:G:N3	2.18	0.76
1:AA:1266:G:O2'	1:AA:2012:G:O6	2.03	0.76
3:AD:35:LYS:HD2	3:AD:104:TYR:CE1	2.20	0.76
3:AD:64:ILE:O	3:AD:64:ILE:HG12	1.85	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:AE:130:GLY:O	4:AE:131:ALA:CB	2.34	0.76
19:AT:65:ARG:HB3	19:AT:70:LEU:HA	1.68	0.76
31:BA:498:A:O2'	31:BA:500:G:O5'	2.04	0.76
32:BE:22:LYS:HZ2	32:BE:22:LYS:H	1.29	0.76
32:BE:21:ARG:HH21	32:BE:38:GLY:HA3	1.50	0.76
34:BG:26:CYS:HA	34:BG:31:CYS:CB	2.16	0.76
54:CA:274:A:H4'	54:CA:275:G:O5'	1.84	0.76
54:CA:96:G:C2'	54:CA:97:U:H5'	2.14	0.76
6:DG:112:PRO:HB3	26:D4:37:SER:HB2	1.65	0.76
3:DD:121:PRO:HB3	3:DD:135:PHE:HE1	1.50	0.76
56:DJ:10:GLU:O	56:DJ:17:VAL:HG12	1.84	0.76
58:DL:138:VAL:HG12	58:DL:139:VAL:N	1.98	0.76
58:DL:49:GLY:HA3	58:DL:50:ASP:HB3	1.65	0.76
58:DL:52:ILE:HD11	58:DL:76:TYR:N	2.00	0.76
21:DV:107:THR:OG1	21:DV:108:PRO:HD3	1.85	0.76
57:DY:115:GLN:HG3	57:DY:115:GLN:O	1.83	0.76
57:DY:18:GLU:CG	57:DY:66:LEU:CD1	2.63	0.76
57:DY:87:VAL:CG1	57:DY:91:LYS:HG3	2.16	0.76
1:AA:999:U:H2'	1:AA:1000:A:H5''	1.66	0.76
1:AA:1175:U:H2'	1:AA:1176:G:H4'	1.67	0.76
1:AA:614:U:H5''	1:AA:615:G:OP1	1.86	0.76
1:AA:888:C:O2'	1:AA:889:C:P	2.44	0.76
12:AP:21:THR:HG21	12:AP:100:GLY:HA3	1.65	0.76
19:AT:35:THR:HG22	19:AT:37:THR:H	1.49	0.76
20:AU:28:LYS:HA	20:AU:28:LYS:HZ3	1.50	0.76
21:AV:141:VAL:HG21	21:AV:144:LEU:CD2	2.14	0.76
1:AA:896:A:H2	21:AV:178:GLU:OE2	1.67	0.76
24:AW:47:ASN:O	24:AW:49:LYS:N	2.19	0.76
31:BA:1152:A:H5''	40:BM:13:HIS:HB2	1.68	0.76
31:BA:495:A:H4'	31:BA:496:A:O5'	1.84	0.76
31:BA:701:C:H1'	31:BA:703:G:C2	2.20	0.76
33:BF:152:ILE:HB	33:BF:199:LYS:HB2	1.65	0.76
33:BF:42:LEU:HA	33:BF:45:LYS:HD2	1.67	0.76
38:BK:51:VAL:HG11	38:BK:60:ARG:HH11	1.50	0.76
53:C1:34:G:H2'	53:C1:35:A:H8	1.50	0.76
54:CA:173:U:H5''	54:CA:197:A:O4'	1.85	0.76
54:CA:677:U:H2'	54:CA:678:U:C6	2.21	0.76
32:CE:96:ARG:H	32:CE:96:ARG:HD2	1.51	0.76
55:DA:1180:C:H2'	55:DA:1181:C:H5'	1.68	0.76
55:DA:229:A:H4'	55:DA:229:A:OP1	1.84	0.76
55:DA:2787:C:O2'	4:DE:61:ARG:HD3	1.86	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:DI:7:ARG:HD3	56:DI:8:ILE:HG12	1.67	0.76
56:DJ:21:LYS:O	56:DJ:24:ILE:HB	1.84	0.76
58:DL:102:GLU:HG2	58:DL:103:GLN:HG2	1.66	0.76
21:DV:196:VAL:C	21:DV:197:ILE:CG1	2.52	0.76
21:DV:191:VAL:CG1	21:DV:197:ILE:HG12	2.07	0.76
57:DY:58:LEU:HA	57:DY:62:ALA:HB2	1.67	0.76
16:A1:98:LEU:C	16:A1:100:VAL:H	1.84	0.76
1:AA:2124:G:H2'	1:AA:2125:G:O4'	1.85	0.76
1:AA:2267:A:H5''	1:AA:2268:A:H5'	1.68	0.76
1:AA:857:C:H5'	22:A3:77:ARG:NH2	2.00	0.76
31:BA:77:C:C2'	31:BA:78:G:H5''	2.13	0.76
32:BE:233:SER:HB2	32:BE:234:PRO:HD2	1.65	0.76
49:BV:42:PRO:HA	49:BV:45:VAL:CG1	2.14	0.76
32:CE:178:ARG:HB2	32:CE:178:ARG:NH1	1.98	0.76
16:D1:60:LEU:HD22	16:D1:64:ARG:HG3	1.67	0.76
55:DA:1698:A:O2'	55:DA:1699:G:H5''	1.86	0.76
3:DD:25:THR:O	3:DD:27:THR:HG22	1.84	0.76
4:DE:117:MET:O	4:DE:118:LYS:HB2	1.86	0.76
7:DH:152:ARG:NE	7:DH:153:LYS:HE3	1.99	0.76
7:DH:153:LYS:HG2	7:DH:162:ILE:H	1.49	0.76
58:DL:10:LEU:HD23	58:DL:10:LEU:N	2.00	0.76
58:DL:99:ILE:CG1	58:DL:138:VAL:HG21	2.14	0.76
55:DA:864:G:OP2	12:DP:22:LYS:HD3	1.85	0.76
21:DV:183:LEU:O	21:DV:184:ALA:HB3	1.84	0.76
21:DV:196:VAL:C	21:DV:197:ILE:HG13	2.06	0.76
1:AA:1074:G:H2'	1:AA:1075:C:H6	1.51	0.76
1:AA:1829:A:N6	1:AA:1976:U:O2	2.18	0.76
1:AA:2131:G:OP1	1:AA:2132:U:H3'	1.84	0.76
1:AA:2176:A:H2'	1:AA:2177:C:C6	2.20	0.76
1:AA:969:U:H2'	1:AA:970:C:C6	2.21	0.76
3:AD:148:GLU:HB2	3:AD:151:LYS:HD2	1.67	0.76
4:AE:2:LYS:HD3	4:AE:95:ILE:HG22	1.65	0.76
5:AF:46:ARG:HG2	5:AF:46:ARG:HH11	1.51	0.76
6:AG:121:ASN:HD22	6:AG:122:PRO:HD2	1.49	0.76
20:AU:87:LYS:HB3	20:AU:92:ASN:CB	2.15	0.76
21:AV:115:GLY:N	21:AV:177:PRO:HB2	2.01	0.76
33:BF:162:GLN:HG2	53:B1:54:U:O2	1.84	0.76
32:BE:165:VAL:HG23	32:BE:166:ASP:H	1.49	0.76
35:BH:43:LEU:H	35:BH:65:ASN:ND2	1.84	0.76
32:CE:5:ILE:HG13	32:CE:221:LEU:CD2	2.15	0.76
49:CV:83:HIS:CD2	49:CV:84:GLY:H	2.03	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:191:G:C4	50:CW:105:SER:HB3	2.21	0.76
6:DG:112:PRO:CB	26:D4:37:SER:HB2	2.15	0.76
55:DA:2128:C:H2'	55:DA:2129:C:C6	2.20	0.76
55:DA:2723:C:OP1	13:D0:3:HIS:HD2	1.67	0.76
3:DD:131:LEU:HD12	3:DD:131:LEU:N	1.99	0.76
3:DD:172:TYR:HB3	3:DD:184:LYS:HG2	1.68	0.76
3:DD:25:THR:HG21	3:DD:82:ILE:H	1.51	0.76
7:DH:21:PRO:HG2	7:DH:22:GLY:H	1.48	0.76
8:DK:115:ALA:HB3	8:DK:128:LEU:CD1	2.06	0.76
57:DY:101:PRO:CG	57:DY:102:LYS:H	1.98	0.76
1:AA:946:G:O2'	1:AA:947:G:O5'	2.03	0.76
6:AG:146:TYR:O	6:AG:149:VAL:HG22	1.86	0.76
7:AH:43:VAL:HG12	7:AH:52:VAL:HG22	1.67	0.76
31:BA:1139:G:N2	31:BA:1143:G:N1	2.34	0.76
52:BB:60:U:H5'	52:BB:61:C:OP2	1.86	0.76
32:BE:12:GLU:HB2	32:BE:16:HIS:ND1	2.01	0.76
39:BL:4:TYR:HB2	39:BL:19:LEU:HB2	1.67	0.76
54:CA:1086:U:H3	54:CA:1099:G:H22	1.30	0.76
54:CA:807:A:H2'	54:CA:808:C:C6	2.21	0.76
42:CO:47:LYS:HB3	42:CO:48:PRO:CD	2.14	0.76
17:D2:98:GLU:C	17:D2:99:ILE:HD13	2.06	0.76
28:D6:34:LEU:H	28:D6:34:LEU:HD13	1.48	0.76
55:DA:1086:A:C5'	55:DA:1103:A:H61	1.99	0.76
55:DA:898:C:H3'	55:DA:899:A:H5'	1.68	0.76
3:DD:121:PRO:HB3	3:DD:135:PHE:CE1	2.21	0.76
7:DH:10:PRO:O	7:DH:11:VAL:HG13	1.86	0.76
10:DN:7:TYR:HE1	10:DN:20:MET:HE3	1.50	0.76
12:DP:74:TYR:CD2	12:DP:91:GLU:HG3	2.20	0.76
25:DX:19:GLN:HE22	25:DX:52:HIS:HE1	1.33	0.76
57:DY:42:GLN:O	57:DY:44:LEU:N	2.17	0.76
57:DY:72:ASP:OD1	57:DY:74:LEU:CA	2.33	0.76
57:DY:91:LYS:HZ3	57:DY:95:GLN:NE2	1.84	0.76
27:A5:40:LYS:HZ2	27:A5:45:VAL:HA	1.50	0.76
1:AA:1613:G:H1	1:AA:1617:C:H2'	1.50	0.76
1:AA:2490:G:H5''	1:AA:2491:U:OP1	1.86	0.76
2:AB:40:U:H6	2:AB:40:U:O5'	1.68	0.76
3:AD:4:LYS:NZ	3:AD:20:ASP:HA	2.00	0.76
1:AA:2787:C:O2'	4:AE:61:ARG:HB3	1.86	0.76
8:AK:4:ILE:HG12	8:AK:18:VAL:HG22	1.66	0.76
39:BL:40:LEU:HD11	39:BL:70:LYS:HG2	1.68	0.76
32:CE:213:LEU:HD21	32:CE:217:ARG:NH1	2.01	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:DL:119:ASP:O	58:DL:122:ALA:HB3	1.86	0.76
58:DL:11:GLN:HB3	58:DL:41:PHE:CZ	2.20	0.76
58:DL:93:ARG:HG2	58:DL:135:GLY:CA	2.14	0.76
15:DR:74:ARG:HG2	15:DR:74:ARG:HH11	1.51	0.76
57:DY:50:ARG:C	57:DY:51:LEU:CG	2.52	0.76
1:AA:1434:A:H61	1:AA:1558:A:H62	1.32	0.76
6:AG:26:GLN:NE2	6:AG:27:ASN:HB2	2.01	0.76
8:AK:12:LEU:HG	8:AK:19:VAL:HG11	1.66	0.76
12:AP:43:THR:OG1	12:AP:46:GLN:HG3	1.84	0.76
31:BA:251:G:H4'	31:BA:252:U:O5'	1.84	0.76
39:BL:8:GLY:HA2	39:BL:79:LEU:HD12	1.67	0.76
54:CA:1499:A:H1'	54:CA:1520:G:H5'	1.66	0.76
54:CA:160:A:H61	54:CA:347:G:H1'	1.49	0.76
32:CE:212:GLN:HG2	32:CE:235:SER:HB2	1.67	0.76
34:CG:33:MET:HE2	34:CG:37:PRO:HA	1.66	0.76
46:CS:28:ARG:HG2	46:CS:28:ARG:HH11	1.51	0.76
55:DA:1095:A:N3	55:DA:1095:A:H2'	2.01	0.76
55:DA:2102:U:H2'	55:DA:2103:C:C6	2.21	0.76
55:DA:582:G:H2'	55:DA:583:G:C8	2.21	0.76
3:DD:124:PRO:HB2	3:DD:126:GLN:NE2	2.00	0.76
58:DL:80:LYS:HE3	58:DL:103:GLN:HB3	1.68	0.76
12:DP:10:ARG:HB2	12:DP:89:ASN:HD21	1.50	0.76
14:DQ:3:ARG:HG2	14:DQ:4:LEU:N	1.99	0.76
19:DT:27:THR:HB	19:DT:80:ILE:HB	1.67	0.76
55:DA:851:U:O2'	25:DX:45:GLY:HA3	1.86	0.76
57:DY:61:LEU:O	57:DY:63:LEU:N	2.19	0.76
57:DY:7:VAL:CG2	57:DY:8:GLU:H	1.68	0.76
13:A0:63:ARG:HB2	13:A0:63:ARG:HH11	1.50	0.76
17:A2:41:GLY:HA3	17:A2:46:VAL:HG11	1.68	0.76
26:A4:12:ALA:HB3	26:A4:24:THR:HG21	1.67	0.76
1:AA:1280:G:H2'	1:AA:1281:G:H5''	1.68	0.76
1:AA:222:A:O2'	1:AA:223:A:OP1	2.04	0.76
1:AA:271(B):G:H4'	1:AA:271(C):U:C5'	2.16	0.76
31:BA:1268:A:H2'	31:BA:1269:A:C8	2.21	0.76
54:CA:686:U:H2'	54:CA:687:A:C8	2.21	0.76
36:CI:37:VAL:HG12	36:CI:38:GLU:H	1.50	0.76
41:CN:34:ASP:HB3	41:CN:40:ILE:HD11	1.67	0.76
45:CR:87:ILE:HG22	45:CR:88:ARG:N	1.98	0.76
55:DA:1058:U:H1'	58:DL:115:LEU:HB2	1.68	0.76
55:DA:1062:G:H2'	55:DA:1063:G:H8	1.49	0.76
55:DA:2468:G:O2'	55:DA:2469:A:P	2.43	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:DE:197:ILE:HD11	4:DE:199:ARG:NH2	2.00	0.76
8:DK:101:LEU:HD21	8:DK:107:VAL:HB	1.66	0.76
8:DK:64:GLU:O	8:DK:67:ARG:HB3	1.85	0.76
58:DL:77:LEU:O	58:DL:80:LYS:HG3	1.86	0.76
9:DM:115:ARG:CA	9:DM:118:LYS:HE3	2.15	0.76
14:DQ:71:ARG:HA	14:DQ:104:GLY:O	1.86	0.76
57:DY:135:ARG:HA	57:DY:138:LEU:HB3	1.66	0.76
1:AA:1212:G:H2'	1:AA:1236:G:N2	2.01	0.75
1:AA:1225:C:H4'	17:A2:85:LYS:CB	2.16	0.75
1:AA:1341:U:C4'	19:AT:57:LEU:HD22	2.16	0.75
10:AN:2:ILE:HD12	10:AN:6:THR:HG21	1.67	0.75
21:AV:163:LEU:CD2	21:AV:163:LEU:H	1.96	0.75
31:BA:1305:G:H22	31:BA:1331:G:C2'	1.99	0.75
31:BA:186(B):C:H2'	31:BA:186(C):G:H8	1.50	0.75
31:BA:741:G:H2'	31:BA:742:G:O4'	1.85	0.75
38:BK:30:ARG:HH11	38:BK:30:ARG:CB	1.97	0.75
48:BU:70:ILE:O	48:BU:74:ARG:HG3	1.85	0.75
54:CA:1053:G:N7	54:CA:1199:U:H3'	1.99	0.75
54:CA:280:C:H4'	54:CA:281:G:OP2	1.87	0.75
34:CG:170:VAL:HG22	34:CG:171:GLY:H	1.50	0.75
49:CV:7:LYS:HZ3	49:CV:7:LYS:CB	1.99	0.75
55:DA:1093:G:H4'	7:DH:170:ARG:HH21	1.50	0.75
55:DA:1359:A:C5'	55:DA:1359:A:C8	2.69	0.75
55:DA:654(I):C:O4'	55:DA:654(I):C:OP1	2.04	0.75
58:DL:77:LEU:HB3	58:DL:107:ILE:CD1	2.16	0.75
9:DM:134:ARG:H	9:DM:135:PRO:CD	1.97	0.75
9:DM:62:VAL:CG1	9:DM:66:LYS:HB2	2.16	0.75
12:DP:87:LYS:O	12:DP:89:ASN:N	2.18	0.75
17:A2:77:ALA:O	17:A2:79:VAL:HG22	1.86	0.75
1:AA:265:A:H2'	1:AA:266:G:O4'	1.87	0.75
1:AA:2735:G:H22	1:AA:2770:G:H1'	1.52	0.75
1:AA:654(S):G:H2'	1:AA:654(T):A:C8	2.21	0.75
8:AK:5:LEU:HD12	8:AK:5:LEU:H	1.49	0.75
14:AQ:26:LEU:O	14:AQ:26:LEU:HD23	1.86	0.75
21:AV:127:LYS:HB3	21:AV:162:GLU:CB	2.16	0.75
23:AZ:86:SER:N	23:AZ:87:PRO:CD	2.49	0.75
34:BG:146:ILE:HD12	34:BG:146:ILE:N	2.01	0.75
48:BU:23:LYS:HA	48:BU:26:LEU:HD11	1.67	0.75
54:CA:1199:U:H4'	40:CM:54:PHE:CE1	2.20	0.75
54:CA:568:G:O6	42:CO:5:PRO:HD3	1.86	0.75
30:D8:29:LYS:NZ	30:D8:44:LYS:HB2	2.01	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1054:A:H2'	55:DA:1055:G:O4'	1.86	0.75
55:DA:2308:G:N2	55:DA:2311:A:H2	1.85	0.75
7:DH:153:LYS:HG2	7:DH:162:ILE:HG13	1.69	0.75
58:DL:20:ALA:CA	58:DL:25:PRO:HD2	2.14	0.75
18:DS:68:ARG:HH21	18:DS:112:GLY:HA3	1.49	0.75
57:DY:55:LYS:HD2	57:DY:79:ALA:CA	2.16	0.75
28:A6:25:LYS:HA	30:A8:34:TRP:CZ3	2.22	0.75
1:AA:2021:C:H5''	1:AA:2022:U:OP2	1.87	0.75
1:AA:1353:A:H4'	3:AD:38:LYS:NZ	2.02	0.75
4:AE:35:GLN:HG3	4:AE:64:LYS:NZ	2.01	0.75
9:AM:15:LEU:HD12	9:AM:136:GLU:HG3	1.68	0.75
21:AV:146:ILE:HG22	21:AV:147:GLY:N	2.00	0.75
24:AW:43:GLN:O	24:AW:44:LEU:HG	1.85	0.75
52:BC:58:A:H1'	52:BC:60:U:C5	2.21	0.75
54:CA:1116:C:H2'	54:CA:1117:G:H5''	1.68	0.75
54:CA:484:G:H4'	54:CA:485:G:O5'	1.85	0.75
54:CA:890:G:O2'	54:CA:891:U:OP2	2.05	0.75
54:CA:946:A:H2'	54:CA:947:G:C8	2.20	0.75
42:CO:60:LEU:CD2	42:CO:60:LEU:N	2.49	0.75
44:CQ:40:CYS:H	44:CQ:43:CYS:CB	2.00	0.75
55:DA:1210:A:H4'	55:DA:1211:U:O5'	1.86	0.75
55:DA:2712:U:C2'	55:DA:2712(A):A:O5'	2.35	0.75
7:DH:135:GLY:HA3	7:DH:141:VAL:HG22	1.69	0.75
56:DJ:18:LEU:O	56:DJ:19:GLU:C	2.22	0.75
8:DK:114:LEU:HD12	8:DK:128:LEU:HD12	1.68	0.75
58:DL:7:VAL:HG11	58:DL:57:ILE:CD1	2.12	0.75
58:DL:56:GLU:O	58:DL:57:ILE:HG22	1.86	0.75
57:DY:30:GLN:O	57:DY:31:GLY:O	2.04	0.75
17:A2:12:TYR:OH	17:A2:22:VAL:HG23	1.87	0.75
26:A4:53:GLU:OE2	26:A4:58:ARG:HB3	1.85	0.75
29:A7:8:ASN:HD22	29:A7:8:ASN:C	1.89	0.75
1:AA:141:A:H5'	1:AA:141(A):C:OP2	1.85	0.75
1:AA:2778:A:H5'	1:AA:2779:U:OP2	1.85	0.75
1:AA:312:G:OP2	1:AA:312:G:H8	1.69	0.75
1:AA:943:U:H2'	1:AA:944:G:H5'	1.67	0.75
2:AB:88:C:H3'	2:AB:89:G:H8	1.50	0.75
3:AD:35:LYS:HD3	3:AD:63:ARG:HB3	1.68	0.75
3:AD:25:THR:HG21	3:AD:82:ILE:H	1.51	0.75
12:AP:81:VAL:O	12:AP:82:ARG:NH1	2.19	0.75
31:BA:1028:C:H2'	31:BA:1028(A):C:C5'	2.16	0.75
31:BA:1117:G:O3'	39:BL:104:ARG:HD3	1.86	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:532:A:H2	54:CA:1206:G:N2	1.85	0.75
54:CA:690:G:H22	41:CN:55:LYS:HZ1	1.33	0.75
54:CA:664:G:H22	54:CA:741:G:H1	1.32	0.75
52:CD:48:C:C5	52:CD:59:U:H1'	2.21	0.75
35:CH:12:LEU:C	35:CH:13:ILE:HD12	2.06	0.75
12:DP:80:GLU:HA	22:D3:4:LYS:CE	2.16	0.75
12:DP:80:GLU:OE2	22:D3:4:LYS:CE	2.34	0.75
55:DA:2712:U:H1'	55:DA:2712(A):A:C8	2.21	0.75
55:DA:2898:U:H2'	55:DA:2899:G:C8	2.21	0.75
6:DG:113:ARG:HH12	6:DG:142:PRO:HA	1.51	0.75
58:DL:111:LYS:HA	58:DL:113:PRO:HG2	1.67	0.75
14:DQ:39:ILE:HD11	14:DQ:73:LEU:HD11	1.68	0.75
20:DU:81:LYS:HB2	20:DU:96:ILE:HG22	1.66	0.75
57:DY:19:ARG:O	57:DY:21:GLN:N	2.19	0.75
57:DY:31:GLY:C	57:DY:32:LEU:HD22	2.07	0.75
1:AA:2820:A:H1'	13:A0:3:HIS:HB3	1.68	0.75
30:A8:22:VAL:HB	30:A8:50:LEU:HD22	1.67	0.75
1:AA:2146:C:H5''	1:AA:2147:G:OP1	1.85	0.75
10:AN:102:VAL:HB	10:AN:106:LEU:HD12	1.69	0.75
15:AR:29:ARG:HG3	15:AR:29:ARG:HH11	1.50	0.75
20:AU:61:ILE:CG2	20:AU:62:GLU:H	1.99	0.75
24:AW:15:LYS:HA	24:AW:67:LYS:HZ1	1.51	0.75
1:AA:1364:G:OP1	23:AZ:3:LYS:HD3	1.86	0.75
31:BA:1004:A:P	31:BA:1025:U:O4	2.45	0.75
31:BA:1118:C:H5'	31:BA:1118:C:H6	1.50	0.75
33:BF:59:ARG:HE	33:BF:64:VAL:HG22	1.52	0.75
31:BA:376:G:H5''	46:BS:5:ARG:HB2	1.67	0.75
36:CI:89:MET:O	36:CI:91:VAL:HG23	1.86	0.75
38:CK:121:ASP:HB2	38:CK:125:ARG:NH2	2.01	0.75
40:CM:24:VAL:HG21	40:CM:37:PRO:HG3	1.67	0.75
55:DA:2150:U:H2'	55:DA:2151:G:C8	2.22	0.75
55:DA:2213:U:H4'	23:DZ:52:ARG:NH1	2.00	0.75
3:DD:35:LYS:CG	3:DD:64:ILE:N	2.49	0.75
58:DL:79:ARG:C	58:DL:81:ALA:H	1.90	0.75
58:DL:78:ILE:HA	58:DL:82:ALA:HB3	1.68	0.75
9:DM:133:GLN:HB2	9:DM:135:PRO:HD3	1.66	0.75
19:DT:50:LYS:H	19:DT:87:GLN:NE2	1.83	0.75
21:DV:152:ALA:HB1	21:DV:163:LEU:HD22	1.67	0.75
21:DV:178:GLU:C	21:DV:180:VAL:H	1.88	0.75
57:DY:138:LEU:C	57:DY:138:LEU:HD12	2.07	0.75
1:AA:1224:G:H5'	1:AA:1225:C:OP2	1.86	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2158:A:H5''	1:AA:2159:G:OP1	1.85	0.75
1:AA:74:A:H4'	1:AA:75:G:O5'	1.86	0.75
21:AV:54:HIS:ND1	21:AV:101:PRO:HD3	2.02	0.75
24:AW:17:SER:OG	24:AW:18:PRO:HA	1.86	0.75
52:BD:20:U:H2'	52:BD:21:A:H5'	1.67	0.75
31:BA:750:G:N3	45:BR:23:GLY:HA3	2.01	0.75
50:BW:77:ALA:O	50:BW:81:LYS:HB2	1.86	0.75
53:C1:56:U:H4'	53:C1:57:U:OP1	1.85	0.75
39:CL:3:GLN:HB3	39:CL:20:ARG:HD3	1.69	0.75
16:D1:92:ARG:HH11	16:D1:95:LEU:HD11	1.50	0.75
17:D2:39:LEU:O	17:D2:40:LEU:HD23	1.85	0.75
6:DG:13:GLU:O	6:DG:14:GLU:HB2	1.85	0.75
7:DH:143:GLN:HE22	7:DH:147:ASN:HD21	1.32	0.75
57:DY:139:VAL:HG23	56:DJ:6:GLU:CD	2.05	0.75
58:DL:11:GLN:O	58:DL:12:LEU:HD22	1.86	0.75
9:DM:71:ILE:N	9:DM:71:ILE:HD13	2.00	0.75
15:DR:91:ARG:HB2	15:DR:121:ILE:HG13	1.69	0.75
1:AA:551:G:H5'	1:AA:1220:A:H1'	1.69	0.75
1:AA:2599:G:C8	3:AD:236:GLY:O	2.39	0.75
1:AA:2657:A:C4	1:AA:2665:A:N6	2.55	0.75
5:AF:143:ALA:HB1	5:AF:148:LEU:HB2	1.67	0.75
31:BA:1301:U:O2	31:BA:1301:U:H2'	1.85	0.75
33:BF:70:VAL:HG12	33:BF:72:LYS:H	1.50	0.75
39:BL:28:VAL:HA	39:BL:63:ILE:O	1.85	0.75
32:CE:42:ILE:HD11	32:CE:202:PRO:HB2	1.67	0.75
33:CF:16:ARG:HH11	33:CF:16:ARG:HB2	1.50	0.75
35:CH:137:GLU:HA	35:CH:140:ARG:NH1	2.01	0.75
36:CI:2:ARG:HD2	36:CI:69:GLU:HB3	1.67	0.75
55:DA:1348:G:C2'	55:DA:1349:A:H5''	2.15	0.75
55:DA:2249:U:H4'	55:DA:2275:C:C5	2.22	0.75
55:DA:27:G:H1'	55:DA:513:A:N6	2.02	0.75
56:DI:4:ASP:HA	56:DI:7:ARG:HD3	1.67	0.75
56:DJ:14:GLN:CG	56:DJ:16:THR:O	2.30	0.75
58:DL:141:ALA:HB1	58:DL:142:PRO:HA	1.68	0.75
55:DA:1064:C:H4'	58:DL:89:HIS:CA	2.16	0.75
11:DO:126:VAL:HG12	11:DO:147:LEU:CD2	2.16	0.75
57:DY:27:VAL:HG23	57:DY:80:VAL:HG11	1.69	0.75
4:AE:117:MET:O	4:AE:118:LYS:HB2	1.85	0.75
6:AG:151:ALA:HB3	6:AG:153:ARG:NH1	2.02	0.75
31:BA:677:U:H2'	31:BA:678:U:C6	2.22	0.75
32:CE:165:VAL:HG23	32:CE:166:ASP:H	1.48	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:CU:26:LEU:HD12	48:CU:29:PHE:CE1	2.22	0.75
16:D1:90:VAL:HG11	17:D2:40:LEU:HG	1.69	0.75
16:D1:90:VAL:HG12	16:D1:91:ASP:N	1.99	0.75
55:DA:2051:A:H61	55:DA:2614:A:H2'	1.51	0.75
58:DL:53:VAL:HB	58:DL:72:PRO:HB2	1.69	0.75
55:DA:483:A:C5'	20:DU:49:VAL:HG13	2.17	0.75
57:DY:130:THR:C	57:DY:132:ASP:N	2.33	0.75
1:AA:361:G:C2	1:AA:362:U:H1'	2.21	0.75
12:AP:21:THR:HG23	12:AP:21:THR:O	1.87	0.75
21:AV:144:LEU:HG	21:AV:144:LEU:O	1.86	0.75
31:BA:60:A:H4'	31:BA:61:G:O5'	1.86	0.75
31:BA:973:G:H1'	40:BM:55:LYS:HE2	1.69	0.75
54:CA:92:G:H2'	54:CA:93:U:O4'	1.87	0.75
33:CF:20:SER:HB2	33:CF:40:ARG:HH22	1.51	0.75
38:CK:6:ILE:HD12	38:CK:6:ILE:H	1.51	0.75
55:DA:1021:A:C8	55:DA:1021:A:H3'	2.22	0.75
55:DA:1678:G:N2	55:DA:1989:G:H22	1.85	0.75
3:DD:25:THR:HG22	3:DD:82:ILE:O	1.85	0.75
3:DD:34:VAL:O	3:DD:34:VAL:HG13	1.87	0.75
3:DD:71:ASP:HB3	3:DD:103:ARG:HH22	1.49	0.75
7:DH:153:LYS:HG3	7:DH:161:GLY:CA	2.17	0.75
56:DI:10:GLU:C	56:DI:14:GLN:HB3	2.06	0.75
16:A1:72:HIS:HE1	16:A1:107:ALA:HA	1.51	0.74
1:AA:1165:U:H2'	1:AA:1166:C:C6	2.21	0.74
6:AG:109:VAL:CG1	26:A4:33:VAL:HG21	2.17	0.74
11:AO:48:PRO:HG2	11:AO:49:ARG:H	1.51	0.74
19:AT:34:ALA:HB1	19:AT:39:ILE:CD1	2.17	0.74
31:BA:1024:G:H2'	31:BA:1025:U:H5''	1.69	0.74
31:BA:17:U:H2'	31:BA:18:C:C6	2.22	0.74
34:BG:92:VAL:O	34:BG:96:LEU:HD23	1.87	0.74
43:BP:79:LYS:HE2	43:BP:82:MET:CE	2.17	0.74
52:CC:20:U:H3'	52:CC:21:A:C5'	2.17	0.74
51:CX:15:ARG:HG2	51:CX:15:ARG:HH11	1.51	0.74
17:D2:38:LEU:HD12	17:D2:56:SER:CA	2.16	0.74
26:D4:37:SER:HB3	26:D4:42:PHE:CE1	2.22	0.74
26:D4:68:ARG:HB3	26:D4:68:ARG:HH11	1.52	0.74
28:D6:52:VAL:HG22	28:D6:53:LYS:H	1.52	0.74
11:DO:61:ARG:NH1	30:D8:14:VAL:HG23	2.02	0.74
55:DA:1112:G:H2'	55:DA:1113:U:H6	1.51	0.74
55:DA:1275:A:H4'	55:DA:1276:A:O5'	1.87	0.74
3:DD:134:ARG:HB2	3:DD:135:PHE:CD2	2.22	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:DH:124:GLU:O	7:DH:126:PRO:HB3	1.87	0.74
7:DH:86:GLU:HG3	7:DH:165:ALA:HB3	1.69	0.74
56:DJ:20:LEU:H	56:DJ:20:LEU:HD12	1.51	0.74
56:DJ:28:LYS:O	56:DJ:30:ALA:N	2.19	0.74
58:DL:112:MET:SD	58:DL:120:LEU:HA	2.27	0.74
12:DP:30:GLY:HA2	12:DP:107:ALA:HB2	1.68	0.74
21:DV:8:TYR:HB2	21:DV:38:TYR:CE2	2.22	0.74
23:DZ:91:LYS:CA	23:DZ:91:LYS:HE3	2.10	0.74
1:AA:140:A:H8	1:AA:1408:C:HO2'	1.28	0.74
1:AA:2656:U:C5	1:AA:2664:G:N2	2.54	0.74
1:AA:945:A:H4'	1:AA:946:G:OP1	1.86	0.74
15:AR:29:ARG:HG3	15:AR:29:ARG:NH1	2.00	0.74
21:AV:125:LEU:HG	21:AV:164:ALA:CB	2.16	0.74
31:BA:1129:C:H5'	31:BA:1130:A:OP1	1.87	0.74
52:BD:20:U:C2'	52:BD:21:A:H5'	2.17	0.74
32:BE:219:VAL:HA	32:BE:222:ILE:HD12	1.70	0.74
43:BP:65:LYS:HE3	43:BP:73:GLU:HG3	1.67	0.74
54:CA:107:G:C2'	54:CA:108:G:H5'	2.18	0.74
55:DA:302:C:H2'	55:DA:303:U:C6	2.22	0.74
55:DA:654:A:N3	55:DA:654:A:H2'	2.02	0.74
55:DA:658:C:H2'	55:DA:659:C:H6	1.50	0.74
56:DI:12:LEU:O	56:DI:13:SER:HB2	1.86	0.74
18:DS:29:LEU:O	18:DS:33:ARG:HG3	1.87	0.74
27:A5:38:ALA:HB3	27:A5:48:GLU:OE2	1.86	0.74
1:AA:1943:U:H4'	1:AA:1944:U:O5'	1.86	0.74
1:AA:529:A:H5''	1:AA:530:G:OP1	1.87	0.74
1:AA:975:G:H1'	1:AA:990:A:C2	2.21	0.74
7:AH:86:GLU:HA	7:AH:132:ARG:HB2	1.67	0.74
9:AM:45:ASN:H	9:AM:45:ASN:HD22	1.35	0.74
9:AM:19:GLU:HA	9:AM:59:LYS:O	1.86	0.74
1:AA:2468:G:H5'	12:AP:120:ILE:CD1	2.18	0.74
31:BA:1101:A:H4'	31:BA:1102:A:O5'	1.87	0.74
31:BA:1363:A:H4'	31:BA:1364:U:OP1	1.85	0.74
31:BA:57:G:H2'	31:BA:58:C:C6	2.22	0.74
32:BE:22:LYS:HZ3	32:BE:40:HIS:CE1	2.06	0.74
54:CA:1095:U:H2'	54:CA:1096:C:C6	2.22	0.74
46:CS:3:LYS:O	46:CS:21:VAL:HA	1.88	0.74
55:DA:2189:U:H2'	55:DA:2190:G:H5''	1.67	0.74
55:DA:2308:G:H22	55:DA:2311:A:H2	1.33	0.74
55:DA:2866:U:O2'	55:DA:2867:G:OP2	2.04	0.74
55:DA:646:A:H2'	55:DA:647:G:O4'	1.86	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:673:C:O2'	5:DF:82:ILE:HD13	1.86	0.74
3:DD:35:LYS:HD2	3:DD:104:TYR:HD1	1.47	0.74
56:DI:24:ILE:HD13	56:DI:26:ALA:H	1.51	0.74
24:DW:13:ALA:HA	24:DW:16:LEU:HD23	1.69	0.74
22:A3:74:ARG:HG3	22:A3:74:ARG:HH11	1.51	0.74
1:AA:1947:C:C3'	1:AA:1948:G:H5''	2.18	0.74
10:AN:111:PHE:O	10:AN:115:VAL:HG23	1.87	0.74
19:AT:11:PRO:HB2	19:AT:13:LEU:HD21	1.69	0.74
20:AU:52:SER:N	20:AU:53:PRO:HD3	2.02	0.74
23:AZ:79:GLY:O	23:AZ:80:LEU:HD13	1.87	0.74
53:B1:51:U:H2'	53:B1:52:U:O4'	1.87	0.74
31:BA:406:G:H5''	34:BG:5:ILE:HG23	1.67	0.74
31:BA:689:C:O2'	31:BA:690:G:H5'	1.87	0.74
54:CA:274:A:HO2'	54:CA:275:G:H8	1.36	0.74
54:CA:630:G:OP1	54:CA:630:G:C4'	2.36	0.74
34:CG:112:VAL:HG12	34:CG:116:GLN:OE1	1.86	0.74
48:CU:21:LYS:O	48:CU:23:LYS:N	2.20	0.74
36:CI:91:VAL:HG11	48:CU:72:ARG:NH1	2.01	0.74
16:D1:98:LEU:C	16:D1:98:LEU:HD23	2.08	0.74
55:DA:1372:U:H6	55:DA:1372:U:C5'	1.83	0.74
55:DA:2067:G:O3'	55:DA:2068:U:H4'	1.87	0.74
55:DA:2146:C:H5''	55:DA:2147:G:OP1	1.87	0.74
14:DQ:83:LYS:O	14:DQ:109:GLY:HA3	1.87	0.74
57:DY:58:LEU:O	57:DY:62:ALA:N	2.17	0.74
57:DY:74:LEU:HD13	57:DY:75:GLN:HG2	1.69	0.74
1:AA:1204:A:O2'	1:AA:1205:U:H5''	1.87	0.74
1:AA:1728:G:N7	1:AA:1731:G:N2	2.34	0.74
1:AA:2393:A:P	30:A8:30:ARG:HB2	2.27	0.74
12:AP:1:MET:HB3	12:AP:69:PHE:HE1	1.50	0.74
19:AT:12:VAL:HG13	19:AT:27:THR:HG23	1.68	0.74
21:AV:145:GLU:OE1	21:AV:174:VAL:HG11	1.85	0.74
31:BA:1148:U:H2'	31:BA:1149:C:O4'	1.87	0.74
31:BA:279:A:O2'	31:BA:280:C:OP2	2.05	0.74
32:BE:54:THR:HG23	32:BE:199:TYR:HB3	1.69	0.74
47:BT:56:VAL:HB	47:BT:78:GLU:HG2	1.67	0.74
54:CA:1182:G:H4'	54:CA:1183:A:C5'	2.17	0.74
54:CA:701:C:H1'	54:CA:703:G:C2	2.22	0.74
39:CL:43:ALA:HA	39:CL:74:ILE:HD13	1.69	0.74
55:DA:718:A:H2'	55:DA:719:C:O4'	1.87	0.74
3:DD:147:LEU:HD11	3:DD:183:ARG:HH12	1.53	0.74
12:DP:1:MET:O	12:DP:2:LEU:HD22	1.88	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:DQ:83:LYS:HG2	14:DQ:109:GLY:H	1.53	0.74
20:DU:50:ARG:HD3	20:DU:53:PRO:HG2	1.69	0.74
12:DP:60:ARG:NE	21:DV:181:GLU:OE1	2.20	0.74
21:DV:20:ARG:HG2	21:DV:20:ARG:HH11	1.51	0.74
57:DY:72:ASP:OD1	57:DY:74:LEU:HA	1.87	0.74
57:DY:74:LEU:HG	57:DY:120:LYS:HA	1.67	0.74
57:DY:51:LEU:HD22	57:DY:82:PHE:H	1.52	0.74
16:A1:92:ARG:HB2	17:A2:11:GLN:NE2	2.02	0.74
1:AA:107:C:H2'	1:AA:108:U:C6	2.23	0.74
1:AA:1342:A:O2'	1:AA:1344:G:P	2.46	0.74
1:AA:1504:C:H2'	1:AA:1505:C:H5'	1.70	0.74
1:AA:2790:A:H1'	1:AA:2893:G:HO2'	1.52	0.74
9:AM:15:LEU:HG	9:AM:134:ARG:NE	2.03	0.74
12:AP:77:LYS:NZ	12:AP:82:ARG:HA	2.02	0.74
24:AW:21:LEU:O	24:AW:25:VAL:HG22	1.87	0.74
24:AW:42:GLY:O	24:AW:44:LEU:N	2.20	0.74
31:BA:376:G:P	46:BS:67:THR:HG21	2.27	0.74
31:BA:736:C:H2'	31:BA:737:A:C8	2.22	0.74
34:BG:189:PRO:HB2	34:BG:194:LEU:HD21	1.69	0.74
35:BH:43:LEU:N	35:BH:65:ASN:HD22	1.85	0.74
36:BI:26:ILE:O	36:BI:30:LEU:HG	1.86	0.74
49:BV:63:THR:HG22	49:BV:66:MET:CE	2.17	0.74
54:CA:198:G:H2'	54:CA:199:G:H8	1.53	0.74
54:CA:942:G:H21	39:CL:124:GLN:NE2	1.86	0.74
37:CJ:38:LEU:HD12	37:CJ:41:ARG:HD2	1.70	0.74
47:CT:45:HIS:NE2	47:CT:47:PRO:HG3	2.03	0.74
55:DA:481:G:HO2'	55:DA:482:A:P	2.11	0.74
3:DD:25:THR:CG2	3:DD:81:ALA:HB1	2.11	0.74
4:DE:68:ALA:O	4:DE:69:LYS:HG3	1.88	0.74
58:DL:52:ILE:HG13	58:DL:76:TYR:HB3	1.66	0.74
20:DU:84:ARG:HH12	20:DU:97:ARG:CA	2.01	0.74
57:DY:72:ASP:O	57:DY:112:LEU:CD2	2.36	0.74
57:DY:51:LEU:CD2	57:DY:82:PHE:H	2.00	0.74
1:AA:1242:A:H5'	1:AA:1243:G:OP2	1.86	0.74
1:AA:1454:U:O2'	1:AA:1455:G:N7	2.21	0.74
1:AA:2080:G:H4'	23:AZ:36:GLY:HA3	1.68	0.74
1:AA:2533:A:C2'	1:AA:2534:A:H5''	2.13	0.74
8:AK:139:GLN:O	8:AK:140:LEU:HB2	1.87	0.74
1:AA:1753:G:H5'	15:AR:95:ARG:HG2	1.68	0.74
53:B1:33:G:H2'	53:B1:34:G:H8	1.50	0.74
31:BA:1139:G:N2	31:BA:1143:G:H1	1.85	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:1149:C:H2'	31:BA:1150:U:C6	2.22	0.74
31:BA:1205:U:H1'	33:BF:195:VAL:CG2	2.18	0.74
35:BH:69:VAL:HG12	35:BH:71:LEU:HD21	1.68	0.74
38:BK:51:VAL:HG11	38:BK:60:ARG:NH1	2.03	0.74
39:BL:97:LYS:HB3	39:BL:98:PRO:HD3	1.69	0.74
54:CA:1322:C:H2'	54:CA:1322:C:O2	1.85	0.74
54:CA:1535:C:C2'	54:CA:1536:C:H5'	2.18	0.74
40:CM:6:ILE:HA	40:CM:97:GLU:O	1.88	0.74
54:CA:1313:U:OP1	49:CV:6:LYS:CB	2.34	0.74
16:D1:95:LEU:HD12	17:D2:11:GLN:HE21	1.53	0.74
55:DA:2712:U:O2'	55:DA:2712(A):A:P	2.46	0.74
55:DA:880:G:H4'	55:DA:880:G:OP1	1.86	0.74
2:DB:48:A:H2'	2:DB:49:C:C6	2.22	0.74
4:DE:103:ASP:OD1	4:DE:201:THR:HA	1.88	0.74
4:DE:78:LEU:HD21	4:DE:79:ARG:HE	1.51	0.74
56:DI:17:VAL:CA	56:DI:20:LEU:HD12	2.17	0.74
56:DI:24:ILE:CA	56:DI:27:LEU:CD1	2.65	0.74
9:DM:96:GLU:C	9:DM:98:VAL:N	2.40	0.74
20:DU:42:VAL:CG1	20:DU:65:ALA:HB3	2.17	0.74
21:DV:120:ILE:HG22	21:DV:121:HIS:CD2	2.22	0.74
21:DV:180:VAL:HG13	21:DV:181:GLU:N	2.03	0.74
57:DY:27:VAL:HG21	57:DY:109:SER:O	1.87	0.74
16:A1:49:HIS:HA	16:A1:52:ARG:HB2	1.69	0.74
26:A4:50:VAL:HG13	26:A4:52:THR:HG23	1.69	0.74
1:AA:1053:C:C3'	1:AA:1054:A:H5''	2.17	0.74
1:AA:1761:C:H5''	1:AA:1762:A:OP2	1.87	0.74
1:AA:2503:A:O2'	1:AA:2505:G:OP2	2.06	0.74
1:AA:654(B):C:C2'	1:AA:654(C):G:O4'	2.36	0.74
3:AD:110:GLY:O	3:AD:112:GLN:HG3	1.87	0.74
4:AE:22:PRO:O	4:AE:23:VAL:HG13	1.86	0.74
20:AU:95:LYS:HB3	20:AU:100:ALA:HA	1.68	0.74
31:BA:1160:G:N1	31:BA:1177:G:N2	2.36	0.74
40:BM:4:ILE:CD1	40:BM:82:ILE:HD11	2.18	0.74
50:BW:51:GLU:HA	50:BW:54:LYS:HB3	1.68	0.74
54:CA:235:C:H5'	47:CT:70:ARG:HG2	1.70	0.74
54:CA:792:A:C4	54:CA:794:A:N6	2.55	0.74
54:CA:406:G:H5''	34:CG:5:ILE:HD13	1.70	0.74
38:CK:91:ARG:HH11	38:CK:91:ARG:HG2	1.53	0.74
43:CP:74:VAL:HA	43:CP:77:ASN:HD22	1.53	0.74
49:CV:5:LEU:HD22	49:CV:10:PHE:CE1	2.22	0.74
27:D5:56:LYS:H	27:D5:56:LYS:CD	2.00	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1050:A:C8	55:DA:2751:G:H2'	2.23	0.74
55:DA:479:A:O2'	55:DA:481:G:H5'	1.87	0.74
43:CP:7:VAL:HB	6:DG:115:ARG:HH22	1.52	0.74
7:DH:30:LYS:HD2	7:DH:81:GLU:H	1.52	0.74
13:A0:86:ARG:NH2	13:A0:118:GLU:HG2	2.02	0.74
26:A4:35:VAL:C	26:A4:37:SER:H	1.90	0.74
1:AA:639:U:H2'	1:AA:640:C:C6	2.23	0.74
1:AA:651:G:H5''	30:A8:18:ALA:HB3	1.69	0.74
3:AD:224:ALA:HB2	3:AD:233:HIS:HB3	1.68	0.74
4:AE:10:GLY:HA3	15:AR:8:LYS:HE2	1.68	0.74
21:AV:116:VAL:HG12	21:AV:117:LEU:N	2.03	0.74
53:B1:37:G:H2'	53:B1:38:U:O4'	1.87	0.74
31:BA:1374:A:O2'	37:BJ:28:ASN:HB3	1.88	0.74
31:BA:210:U:O2	31:BA:210:U:H2'	1.87	0.74
31:BA:274:A:O2'	31:BA:275:G:H8	1.71	0.74
31:BA:481:G:H5''	31:BA:482:A:OP1	1.88	0.74
34:BG:24:GLU:H	34:BG:27:TYR:HB2	1.53	0.74
54:CA:107:G:H2'	54:CA:108:G:H5'	1.70	0.74
55:DA:2420:C:P	30:D8:34:TRP:H	2.11	0.74
55:DA:1534:G:O6	55:DA:1538:G:N2	2.21	0.74
55:DA:2211:G:O2'	55:DA:2212:A:OP2	2.04	0.74
55:DA:2287:A:C2	55:DA:2346:A:N1	2.55	0.74
55:DA:2854:G:H2'	55:DA:2855:C:C6	2.23	0.74
55:DA:2887:U:H2'	55:DA:2888:C:H6	1.51	0.74
8:DK:64:GLU:HG3	8:DK:67:ARG:CZ	2.18	0.74
10:DN:113:LYS:HG2	10:DN:117:LEU:HD11	1.68	0.74
14:DQ:5:THR:OG1	14:DQ:7:TYR:HB3	1.88	0.74
23:DZ:83:GLU:OE1	23:DZ:85:LEU:HD23	1.87	0.74
1:AA:1175:U:C2'	1:AA:1176:G:H4'	2.18	0.74
1:AA:608:A:C5	1:AA:621:A:N6	2.55	0.74
1:AA:644:A:H4'	1:AA:645:C:C5	2.23	0.74
1:AA:832:G:H5''	11:AO:45:LEU:HD11	1.68	0.74
1:AA:879:G:H1	1:AA:898:C:N4	1.84	0.74
5:AF:175:THR:O	5:AF:176:LEU:HB2	1.86	0.74
7:AH:153:LYS:CB	7:AH:161:GLY:HA2	2.18	0.74
31:BA:1020:U:C2'	31:BA:1021:G:H5''	2.18	0.74
52:BC:23:A:H2'	52:BC:24:G:C8	2.23	0.74
54:CA:537:G:H5''	42:CO:113:ARG:HH12	1.53	0.74
36:CI:62:TRP:C	36:CI:63:TYR:HD2	1.91	0.74
39:CL:6:GLY:HA3	39:CL:84:ALA:HB2	1.69	0.74
55:DA:1497:U:H5'	55:DA:1498:C:OP2	1.88	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2308:G:N1	55:DA:2311:A:C2	2.56	0.74
58:DL:95:LYS:CD	58:DL:136:VAL:HG21	2.18	0.74
20:DU:76:CYS:HB3	20:DU:96:ILE:HD13	1.68	0.74
21:DV:196:VAL:HG13	21:DV:196:VAL:O	1.86	0.74
21:DV:51:ALA:HA	21:DV:55:HIS:HD2	1.51	0.74
21:DV:60:GLU:HA	21:DV:66:SER:HA	1.70	0.74
57:DY:138:LEU:C	57:DY:140:GLY:H	1.92	0.74
57:DY:43:ALA:N	57:DY:47:ASN:ND2	2.35	0.74
17:A2:61:VAL:HG13	17:A2:62:LEU:H	1.53	0.73
12:AP:7:MET:HB2	12:AP:10:ARG:NE	2.02	0.73
1:AA:2496:C:OP1	12:AP:81:VAL:HG13	1.86	0.73
15:AR:3:ARG:HG2	15:AR:6:LEU:HB2	1.69	0.73
33:BF:36:ASP:HA	33:BF:39:ILE:HD12	1.70	0.73
43:BP:8:GLU:OE1	43:BP:22:ILE:HA	1.88	0.73
54:CA:33:A:H2'	54:CA:34:C:C6	2.23	0.73
33:CF:130:VAL:O	33:CF:134:ILE:HG12	1.88	0.73
44:CQ:60:SER:O	44:CQ:61:TRP:HB3	1.86	0.73
55:DA:2015:A:C1'	27:D5:2:ALA:HA	2.17	0.73
5:DF:20:LEU:HD12	5:DF:21:ALA:H	1.52	0.73
56:DI:21:LYS:N	56:DI:24:ILE:HD12	2.03	0.73
58:DL:110:GLN:C	58:DL:111:LYS:HE2	2.07	0.73
21:DV:187:ALA:O	21:DV:188:ALA:CB	2.36	0.73
57:DY:26:LEU:O	57:DY:111:LEU:HD13	1.87	0.73
57:DY:26:LEU:N	57:DY:82:PHE:HZ	1.82	0.73
1:AA:1019:U:N3	1:AA:1142(A):A:N6	2.36	0.73
1:AA:2287:A:N6	1:AA:2344:U:N3	2.36	0.73
1:AA:2378:A:H4'	14:AQ:23:ARG:NH1	2.02	0.73
1:AA:273(E):U:O2'	1:AA:273(F):C:H5'	1.88	0.73
1:AA:674:G:C1'	5:AF:74:ARG:HD3	2.17	0.73
1:AA:811:U:H2'	11:AO:21:ARG:HG3	1.69	0.73
31:BA:977:A:C2'	31:BA:978:A:H5'	2.18	0.73
34:BG:19:LEU:HD12	34:BG:21:LEU:HD23	1.69	0.73
33:CF:21:ARG:HD3	33:CF:21:ARG:N	2.03	0.73
54:CA:262:A:H5'	50:CW:74:LYS:HG3	1.70	0.73
55:DA:2115:G:H2'	55:DA:2116:G:C8	2.23	0.73
4:DE:117:MET:O	4:DE:117:MET:HG2	1.87	0.73
7:DH:152:ARG:O	7:DH:153:LYS:CB	2.36	0.73
56:DI:24:ILE:N	56:DI:27:LEU:CD1	2.51	0.73
58:DL:18:THR:HB	58:DL:19:PRO:HD2	1.68	0.73
58:DL:52:ILE:HG13	58:DL:76:TYR:HB2	1.69	0.73
11:DO:36:LYS:HB2	11:DO:40:SER:HB3	1.69	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:DE:7:VAL:HG21	15:DR:1:MET:HE1	1.70	0.73
15:DR:96:ARG:HB2	15:DR:96:ARG:NH1	2.03	0.73
23:DZ:23:LYS:HE3	23:DZ:29:GLY:CA	2.18	0.73
1:AA:2128:C:H2'	1:AA:2129:C:C6	2.23	0.73
1:AA:2394:C:P	11:AO:63:PRO:HD2	2.28	0.73
21:AV:144:LEU:C	21:AV:146:ILE:N	2.37	0.73
31:BA:1453:G:H3'	50:BW:39:LYS:HZ2	1.54	0.73
31:BA:794:A:C2	31:BA:795:C:C4	2.76	0.73
42:BO:32:PHE:HB3	42:BO:84:LEU:HD21	1.69	0.73
43:BP:17:VAL:O	43:BP:18:ALA:C	2.26	0.73
46:BS:5:ARG:HH21	46:BS:24:ALA:HA	1.53	0.73
54:CA:562:C:HO2'	42:CO:15:ARG:HB3	1.52	0.73
54:CA:828:A:H2'	54:CA:829:G:O4'	1.89	0.73
54:CA:960:U:O2	54:CA:960:U:H2'	1.86	0.73
52:CD:41:C:C3'	52:CD:42:C:H5''	2.19	0.73
42:CO:71:PRO:HG3	42:CO:99:HIS:HD2	1.52	0.73
55:DA:1055:G:H2'	55:DA:1056:G:H5'	1.69	0.73
55:DA:1058:U:H5'	58:DL:4:VAL:HB	1.71	0.73
55:DA:2849:U:H2'	55:DA:2866:U:O2	1.88	0.73
56:DJ:17:VAL:O	56:DJ:18:LEU:HB2	1.87	0.73
8:DK:60:GLU:HG3	8:DK:61:ARG:HH22	1.53	0.73
8:DK:72:LEU:HD13	8:DK:107:VAL:HG11	1.69	0.73
58:DL:106:GLU:HG2	58:DL:109:LYS:CB	2.18	0.73
55:DA:1058:U:P	58:DL:5:VAL:HG22	2.28	0.73
12:DP:88:GLY:C	12:DP:90:VAL:H	1.90	0.73
21:DV:116:VAL:CG1	21:DV:118:GLN:OE1	2.36	0.73
21:DV:186:GLU:O	21:DV:186:GLU:CG	2.36	0.73
57:DY:10:LEU:HA	57:DY:13:LEU:HD12	1.70	0.73
1:AA:2331:G:H4'	22:A3:43:THR:H	1.53	0.73
1:AA:1503:U:H2'	1:AA:1504:C:C6	2.23	0.73
1:AA:2537:U:H2'	1:AA:2538:C:C6	2.24	0.73
5:AF:9:ILE:HA	5:AF:15:SER:O	1.89	0.73
7:AH:102:ALA:HB1	7:AH:115:VAL:O	1.87	0.73
7:AH:98:LEU:HD13	7:AH:125:VAL:HG21	1.69	0.73
8:AK:123:LEU:HD22	8:AK:143:SER:HB2	1.71	0.73
33:BF:119:ARG:NH2	33:BF:137:ALA:HA	2.04	0.73
54:CA:1060:C:C4	33:CF:2:GLY:HA3	2.22	0.73
52:CD:8:U:C2'	52:CD:13:C:H41	2.00	0.73
47:CT:21:VAL:HG11	47:CT:59:ILE:HD11	1.68	0.73
51:CX:6:ARG:HE	51:CX:15:ARG:CZ	2.01	0.73
55:DA:1709:U:H2'	55:DA:1710:C:C6	2.22	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1718:G:H2'	55:DA:1725:G:H5''	1.71	0.73
55:DA:387:U:H4'	55:DA:388:G:O5'	1.86	0.73
55:DA:404:C:O2'	55:DA:405:U:OP2	2.06	0.73
55:DA:607:U:OP1	5:DF:102:PRO:HA	1.87	0.73
55:DA:74:A:H4'	55:DA:75:G:O5'	1.86	0.73
3:DD:227:ASN:HB3	3:DD:228:PRO:HD2	1.69	0.73
56:DJ:5:ILE:HG22	56:DJ:9:LYS:CG	2.17	0.73
21:DV:193:GLU:N	21:DV:194:PRO:CD	2.51	0.73
24:DW:16:LEU:O	24:DW:16:LEU:CG	2.34	0.73
2:AB:34:U:H5''	2:AB:35:U:OP1	1.89	0.73
3:AD:106:ILE:HD11	3:AD:196:VAL:HG13	1.69	0.73
4:AE:14:ILE:HG13	15:AR:14:TYR:CZ	2.23	0.73
5:AF:53:THR:HG22	5:AF:56:GLU:CD	2.08	0.73
7:AH:24:VAL:HG21	7:AH:72:ILE:HG23	1.71	0.73
14:AQ:27:SER:HA	14:AQ:88:ASP:HB3	1.70	0.73
33:BF:79:ARG:HE	33:BF:79:ARG:N	1.86	0.73
31:BA:1347:G:H3'	39:BL:108:VAL:O	1.87	0.73
54:CA:1160:G:H1	54:CA:1177:G:H21	1.36	0.73
54:CA:1422:G:H5''	10:DN:48:PRO:HB3	1.71	0.73
54:CA:1434:A:H2'	54:CA:1435:G:O4'	1.89	0.73
32:CE:18:GLY:H	32:CE:42:ILE:CG2	2.00	0.73
33:CF:50:ALA:HB1	33:CF:70:VAL:HG11	1.70	0.73
55:DA:2732:G:H3'	55:DA:2733:A:H5'	1.71	0.73
55:DA:608:A:N9	55:DA:621:A:N6	2.36	0.73
56:DJ:1:MET:CG	56:DJ:2:ALA:H	1.97	0.73
58:DL:53:VAL:HA	58:DL:72:PRO:O	1.88	0.73
11:DO:115:LEU:HA	11:DO:134:ALA:HB2	1.71	0.73
11:DO:120:ALA:HB1	11:DO:138:LEU:HA	1.70	0.73
12:DP:43:THR:OG1	12:DP:46:GLN:HG3	1.89	0.73
18:DS:4:LYS:HB3	18:DS:106:ILE:HG22	1.69	0.73
21:DV:178:GLU:HG3	21:DV:180:VAL:N	2.04	0.73
21:DV:187:ALA:CB	21:DV:193:GLU:CG	2.65	0.73
57:DY:122:VAL:CA	57:DY:126:ALA:HB3	2.18	0.73
17:A2:79:VAL:O	17:A2:80:GLN:CD	2.27	0.73
1:AA:181:A:H5''	29:A7:36:GLN:OE1	1.89	0.73
1:AA:1502:C:H5'	1:AA:1503:U:OP2	1.88	0.73
1:AA:1924:C:C4	1:AA:1925:C:H5	2.02	0.73
1:AA:654(I):C:O2'	1:AA:654(J):A:O5'	2.07	0.73
1:AA:813:U:H2'	1:AA:814:C:C6	2.22	0.73
1:AA:879:G:N2	1:AA:898:C:N3	2.35	0.73
5:AF:79:GLY:HA2	5:AF:86:GLY:HA2	1.70	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:AG:34:LEU:HB2	6:AG:172:LEU:HD21	1.70	0.73
20:AU:94:LYS:HD2	20:AU:101:LYS:NZ	2.04	0.73
21:AV:95:PRO:O	21:AV:96:VAL:HB	1.88	0.73
31:BA:1006:C:H2'	31:BA:1007:C:H6	1.52	0.73
42:BO:75:HIS:HD2	42:BO:77:LEU:HB2	1.54	0.73
44:BQ:29:ARG:HG2	44:BQ:40:CYS:CB	2.17	0.73
45:BR:43:LEU:HD11	45:BR:53:HIS:HA	1.70	0.73
49:BV:29:ARG:HG3	49:BV:48:THR:OG1	1.89	0.73
54:CA:802:A:H2'	54:CA:803:G:O4'	1.88	0.73
46:CS:43:LYS:HG2	46:CS:48:TRP:CE3	2.23	0.73
54:CA:254:G:OP1	47:CT:67:LYS:O	2.06	0.73
49:CV:83:HIS:HD2	49:CV:84:GLY:N	1.85	0.73
30:D8:35:GLN:HA	30:D8:35:GLN:NE2	2.04	0.73
55:DA:1019:U:HO2'	55:DA:1021:A:H2	0.77	0.73
55:DA:414:C:H1'	55:DA:1864:U:O2'	1.87	0.73
55:DA:654(B):C:C2'	55:DA:654(C):G:O4'	2.36	0.73
55:DA:888:C:O2'	55:DA:889:C:H5'	1.89	0.73
4:DE:131:ALA:HB1	4:DE:135:HIS:HE1	1.52	0.73
7:DH:9:ILE:HG22	7:DH:51:ARG:HG2	1.69	0.73
56:DI:11:GLU:HA	56:DI:14:GLN:OE1	1.88	0.73
58:DL:18:THR:CG2	58:DL:38:VAL:HG11	2.18	0.73
9:DM:7:LYS:N	9:DM:7:LYS:HZ2	1.87	0.73
11:DO:15:ARG:O	11:DO:16:ARG:C	2.23	0.73
11:DO:71:VAL:HG13	11:DO:72:PRO:HD3	1.69	0.73
12:DP:60:ARG:HG3	21:DV:181:GLU:CD	2.08	0.73
57:DY:49:ALA:CA	57:DY:84:GLU:O	2.36	0.73
57:DY:49:ALA:N	57:DY:84:GLU:CB	2.51	0.73
17:A2:48:GLY:CA	17:A2:52:VAL:HG22	2.18	0.73
1:AA:1534:G:N2	1:AA:1538:G:O6	2.21	0.73
1:AA:2245:U:H5'	1:AA:2246:G:H5'	1.69	0.73
1:AA:2322:A:H3'	1:AA:2323:G:H8	1.53	0.73
1:AA:865:C:H4'	1:AA:866:A:OP1	1.88	0.73
31:BA:32:A:H2'	31:BA:33:A:C8	2.23	0.73
31:BA:1248:A:H2'	39:BL:70:LYS:HZ1	1.54	0.73
41:BN:67:ASP:OD1	41:BN:71:LYS:HE3	1.87	0.73
54:CA:376:G:O2'	54:CA:377:G:H5'	1.89	0.73
52:CD:7:A:H5'	52:CD:8:U:OP2	1.89	0.73
54:CA:1205:U:H5'	33:CF:190:ARG:HH21	1.53	0.73
34:CG:11:LEU:C	34:CG:13:ARG:N	2.39	0.73
54:CA:690:G:N2	41:CN:55:LYS:HZ1	1.86	0.73
3:DD:60:ARG:HD3	3:DD:87:ASN:OD1	1.89	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:DH:98:LEU:HD13	7:DH:125:VAL:HG21	1.70	0.73
56:DI:7:ARG:NE	56:DI:8:ILE:CG1	2.46	0.73
8:DK:124:GLY:O	8:DK:142:VAL:HG22	1.88	0.73
58:DL:112:MET:HE3	58:DL:118:THR:O	1.87	0.73
55:DA:1079:C:H1'	58:DL:129:GLY:CA	2.18	0.73
21:DV:120:ILE:HB	21:DV:171:ILE:H	1.52	0.73
57:DY:11:ALA:HB1	57:DY:52:PHE:CE1	2.24	0.73
57:DY:23:SER:HB2	57:DY:68:LEU:O	1.88	0.73
1:AA:1653:G:O2'	1:AA:1654:A:OP2	2.04	0.73
1:AA:2183:C:H2'	1:AA:2184:G:H8	1.53	0.73
1:AA:2314:C:O2'	1:AA:2315:G:H5'	1.89	0.73
3:AD:62:TYR:HA	3:AD:87:ASN:ND2	2.04	0.73
4:AE:103:ASP:OD1	4:AE:201:THR:HG23	1.88	0.73
1:AA:1138:G:H21	9:AM:106:MET:CE	2.02	0.73
25:AX:4:LEU:HD21	25:AX:56:VAL:HG13	1.70	0.73
31:BA:1333:A:H2'	31:BA:1334:G:O4'	1.88	0.73
31:BA:186:C:H1'	50:BW:81:LYS:NZ	2.03	0.73
33:BF:25:GLY:C	33:BF:27:LYS:H	1.92	0.73
43:BP:23:TYR:HE1	43:BP:71:ARG:HD3	1.54	0.73
47:BT:78:GLU:OE1	47:BT:81:ARG:HD2	1.89	0.73
49:BV:9:VAL:HG12	49:BV:10:PHE:N	2.03	0.73
54:CA:1535:C:O2'	54:CA:1536:C:H5'	1.89	0.73
54:CA:129(A):G:O2'	54:CA:189:U:H3'	1.88	0.73
38:CK:112:LEU:HA	38:CK:134:ILE:HG12	1.71	0.73
39:CL:59:PHE:HZ	39:CL:88:TYR:HE1	1.36	0.73
43:CP:49:THR:HG22	43:CP:51:ALA:H	1.53	0.73
13:D0:86:ARG:HE	13:D0:118:GLU:HG2	1.53	0.73
55:DA:593:G:O2'	30:D8:61:LEU:CD1	2.37	0.73
55:DA:1934:C:H5'	55:DA:1934:C:C6	2.16	0.73
55:DA:893:C:H3'	55:DA:894:C:C5	2.23	0.73
56:DJ:6:GLU:O	56:DJ:10:GLU:HG2	1.89	0.73
57:DY:8:GLU:O	57:DY:9:LEU:O	2.04	0.73
27:A5:40:LYS:NZ	27:A5:45:VAL:HA	2.04	0.73
1:AA:1331:A:H2'	1:AA:1333:C:H5	1.54	0.73
1:AA:2173:A:H5''	1:AA:2174:C:C5	2.24	0.73
1:AA:2318:G:H22	14:AQ:2:ALA:N	1.85	0.73
1:AA:2422:A:H4'	1:AA:2423:U:OP1	1.88	0.73
1:AA:448:U:O4	1:AA:583:G:H1'	1.88	0.73
6:AG:125:PHE:HB3	6:AG:166:ASP:HB2	1.71	0.73
6:AG:56:ALA:HB2	6:AG:153:ARG:HE	1.52	0.73
7:AH:103:LEU:HD23	7:AH:115:VAL:O	1.88	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AO:52:GLU:HG2	11:AO:55:ARG:O	1.87	0.73
21:AV:140:ASP:O	21:AV:141:VAL:CB	2.37	0.73
21:AV:148:ASP:O	21:AV:149:SER:CB	2.36	0.73
33:BF:95:THR:HG22	33:BF:97:LYS:HG2	1.71	0.73
54:CA:1241:G:H2'	54:CA:1242:C:C6	2.24	0.73
54:CA:232:G:H2'	54:CA:233:C:C6	2.24	0.73
54:CA:566:G:H4'	54:CA:567:G:OP1	1.88	0.73
52:CC:18:G:O6	52:CC:55:U:H1'	1.88	0.73
33:CF:181:ASN:HD21	33:CF:204:LEU:HD12	1.52	0.73
40:CM:6:ILE:HD11	40:CM:72:VAL:HB	1.69	0.73
44:CQ:47:LEU:HA	44:CQ:50:LYS:HG3	1.71	0.73
54:CA:1305:G:H5'	51:CX:4:GLY:HA3	1.71	0.73
55:DA:303:U:H2'	55:DA:304:G:H8	1.54	0.73
55:DA:548:A:H2'	55:DA:549:G:H5'	1.71	0.73
2:DB:74:U:H2'	2:DB:75:G:C5'	2.17	0.73
56:DI:30:ALA:HA	56:DJ:3:LEU:CG	2.19	0.73
58:DL:101:TRP:CD1	58:DL:101:TRP:N	2.57	0.73
58:DL:111:LYS:HD2	58:DL:111:LYS:N	2.03	0.73
55:DA:1077:A:C4'	58:DL:93:ARG:NH2	2.51	0.73
58:DL:95:LYS:HG2	58:DL:136:VAL:HG11	1.70	0.73
55:DA:2562:U:H1'	10:DN:23:ARG:HH11	1.52	0.73
11:DO:126:VAL:HA	11:DO:145:PRO:HB2	1.70	0.73
2:DB:50:G:OP1	14:DQ:63:THR:HG23	1.88	0.73
21:DV:150:LEU:CD2	21:DV:154:ASP:CG	2.57	0.73
21:DV:69:THR:HG22	21:DV:90:VAL:HA	1.70	0.73
57:DY:98:LYS:HG2	57:DY:102:LYS:HA	1.69	0.73
17:A2:58:VAL:HG21	17:A2:100:ARG:HH21	1.53	0.73
26:A4:34:GLU:HB3	43:BP:57:ARG:NH1	2.03	0.73
28:A6:22:ALA:HB3	28:A6:42:TRP:CZ2	2.22	0.73
1:AA:404:C:H1'	1:AA:406:G:C8	2.24	0.73
1:AA:2751:G:C6	7:AH:2:SER:HB3	2.24	0.73
20:AU:69:ALA:O	20:AU:72:VAL:HG22	1.88	0.73
45:BR:17:ARG:HG3	45:BR:17:ARG:HH11	1.54	0.73
48:BU:65:ILE:HD12	48:BU:66:LEU:N	2.04	0.73
53:C1:57:U:H3'	53:C1:57:U:O2	1.88	0.73
52:CD:23:A:H2'	52:CD:24:G:H8	1.52	0.73
32:CE:134:GLU:O	32:CE:138:LEU:HG	1.88	0.73
34:CG:119:GLN:HG3	34:CG:123:HIS:CD2	2.23	0.73
55:DA:1204:A:C2	55:DA:1241:A:N1	2.56	0.73
55:DA:2134:A:N6	55:DA:2157:G:H1'	2.04	0.73
4:DE:53:PRO:HG2	4:DE:54:GLN:H	1.52	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:DH:154:PRO:HD3	7:DH:161:GLY:HA3	1.71	0.73
56:DI:24:ILE:HD13	56:DI:25:ASP:N	2.03	0.73
9:DM:134:ARG:N	9:DM:135:PRO:HD3	2.01	0.73
11:DO:83:VAL:CG1	11:DO:112:LEU:HD21	2.19	0.73
14:DQ:10:ARG:O	14:DQ:12:PHE:N	2.22	0.73
57:DY:46:GLN:O	57:DY:47:ASN:CB	2.36	0.73
28:A6:34:LEU:HD23	28:A6:34:LEU:H	1.52	0.72
5:AF:89:VAL:HG12	5:AF:90:PHE:N	2.03	0.72
1:AA:483:A:H5''	20:AU:49:VAL:HG13	1.70	0.72
20:AU:13:VAL:HG23	20:AU:73:ARG:C	2.10	0.72
31:BA:664:G:H22	31:BA:741:G:H1	1.37	0.72
31:BA:807:A:H2'	31:BA:808:C:C6	2.23	0.72
37:BJ:22:LEU:HG	37:BJ:62:PHE:HE2	1.54	0.72
39:BL:40:LEU:HB2	39:BL:43:ALA:HB2	1.71	0.72
47:BT:9:VAL:HG22	47:BT:56:VAL:HG22	1.70	0.72
54:CA:1036:G:H3'	54:CA:1037:C:C6	2.24	0.72
54:CA:487:A:H2'	54:CA:488:C:O4'	1.89	0.72
54:CA:99:C:H2'	54:CA:101:A:C8	2.24	0.72
37:CJ:73:MET:HG2	37:CJ:90:GLU:HA	1.70	0.72
6:DG:142:PRO:HB2	26:D4:31:ILE:HG21	1.71	0.72
55:DA:2133:G:H2'	55:DA:2157:G:N2	2.03	0.72
55:DA:229:A:O2'	55:DA:230:U:P	2.46	0.72
9:DM:15:LEU:HD13	9:DM:16:ILE:N	2.04	0.72
14:DQ:59:LYS:HG2	14:DQ:60:GLY:N	1.99	0.72
57:DY:116:ILE:HG13	57:DY:117:LEU:HD22	1.71	0.72
57:DY:89:ALA:HA	57:DY:125:LEU:O	1.89	0.72
26:A4:39:CYS:O	26:A4:40:HIS:HB2	1.89	0.72
1:AA:2111:C:H41	1:AA:2147:G:H21	1.37	0.72
5:AF:4:VAL:HA	5:AF:19:GLU:HB2	1.69	0.72
21:AV:145:GLU:OE1	21:AV:145:GLU:HA	1.89	0.72
21:AV:148:ASP:OD2	21:AV:174:VAL:O	2.07	0.72
24:AW:65:ASN:HD22	24:AW:69:ARG:NH2	1.86	0.72
46:BS:14:ASN:N	46:BS:15:PRO:HD3	2.03	0.72
54:CA:57:G:H2'	54:CA:58:C:H6	1.53	0.72
17:D2:49:THR:CB	17:D2:50:PRO:HD2	2.19	0.72
26:D4:68:ARG:NH1	26:D4:68:ARG:HB3	2.04	0.72
2:DB:43:C:P	6:DG:67:LYS:HE3	2.29	0.72
7:DH:127:GLU:OE1	7:DH:128:PRO:HD2	1.89	0.72
56:DI:24:ILE:CD1	56:DI:25:ASP:N	2.52	0.72
8:DK:60:GLU:HG3	8:DK:61:ARG:NH2	2.04	0.72
58:DL:72:PRO:N	58:DL:73:PRO:CD	2.51	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:DO:144:GLU:N	11:DO:145:PRO:HD3	2.04	0.72
57:DY:32:LEU:HB3	57:DY:33:PRO:HD3	1.70	0.72
26:A4:56:VAL:HG12	26:A4:57:GLU:HG3	1.70	0.72
1:AA:1098:A:H3'	1:AA:1099:G:C5'	2.19	0.72
1:AA:226:G:H1'	1:AA:228:A:H61	1.53	0.72
1:AA:583:G:H5''	16:A1:10:ARG:NH1	2.04	0.72
1:AA:654(C):G:H3'	1:AA:654(D):G:C8	2.25	0.72
2:AB:46:A:H2'	2:AB:47:C:C6	2.23	0.72
4:AE:31:CYS:SG	4:AE:51:PHE:HB2	2.29	0.72
8:AK:78:THR:HG21	8:AK:104:GLN:HE22	1.54	0.72
31:BA:8:A:H1'	35:BH:102:ALA:CA	2.18	0.72
42:BO:27:LEU:HD11	42:BO:60:LEU:HB3	1.70	0.72
48:BU:22:VAL:HG12	48:BU:55:ARG:O	1.88	0.72
49:BV:5:LEU:HD22	49:BV:6:LYS:N	2.04	0.72
34:CG:190:ASP:HB3	34:CG:193:ASP:OD1	1.89	0.72
17:D2:52:VAL:CG2	17:D2:55:ALA:HB3	2.20	0.72
28:D6:13:CYS:O	28:D6:21:TYR:HA	1.89	0.72
55:DA:1416:G:HO2'	55:DA:1417:C:H6	1.34	0.72
55:DA:445:C:H2'	55:DA:446:G:O4'	1.89	0.72
55:DA:627:A:O2'	55:DA:628:G:C8	2.42	0.72
55:DA:871:U:H2'	55:DA:871:U:O2	1.89	0.72
3:DD:35:LYS:HE3	3:DD:65:ILE:N	2.03	0.72
6:DG:180:PHE:C	6:DG:182:LYS:H	1.90	0.72
56:DI:1:MET:SD	56:DI:5:ILE:HG21	2.30	0.72
8:DK:94:ALA:HA	8:DK:97:ILE:HG12	1.70	0.72
58:DL:111:LYS:C	58:DL:113:PRO:HD2	2.09	0.72
58:DL:12:LEU:CB	58:DL:13:PRO:HA	2.14	0.72
21:DV:187:ALA:O	21:DV:188:ALA:HB2	1.87	0.72
57:DY:51:LEU:CD1	57:DY:82:PHE:CA	2.67	0.72
1:AA:1053:C:H3'	1:AA:1054:A:H5''	1.72	0.72
1:AA:895:U:H4'	1:AA:896:A:H8	1.54	0.72
4:AE:47:VAL:HG12	4:AE:48:GLN:H	1.55	0.72
21:AV:118:GLN:NE2	21:AV:118:GLN:HA	2.04	0.72
21:AV:150:LEU:HB3	21:AV:172:ALA:HB3	1.72	0.72
21:AV:6:LYS:HD3	21:AV:8:TYR:OH	1.89	0.72
40:BM:22:LYS:HE2	40:BM:90:LEU:HD13	1.70	0.72
49:BV:41:VAL:HG23	49:BV:44:MET:HE3	1.71	0.72
54:CA:1399:C:C4'	54:CA:1400:C:O5'	2.35	0.72
54:CA:376:G:H5''	46:CS:5:ARG:HD2	1.70	0.72
32:CE:172:ILE:HD12	32:CE:172:ILE:H	1.53	0.72
33:CF:84:ILE:O	33:CF:88:ARG:HG3	1.89	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:CJ:113:GLU:CB	37:CJ:119:ARG:HG2	2.19	0.72
43:CP:3:ARG:NH1	43:CP:7:VAL:HG22	2.04	0.72
27:D5:4:HIS:HB3	27:D5:5:PRO:HD3	1.70	0.72
28:D6:28:ARG:HB3	28:D6:28:ARG:HH11	1.54	0.72
55:DA:1510:A:H2	55:DA:1513:C:H42	1.37	0.72
55:DA:1510:A:O2'	55:DA:1511:A:H5'	1.88	0.72
55:DA:1887:C:H3'	55:DA:1888:G:H5''	1.71	0.72
55:DA:879:G:O6	55:DA:898:C:N4	2.21	0.72
56:DI:9:LYS:C	56:DI:11:GLU:N	2.42	0.72
56:DJ:13:SER:HB3	56:DJ:17:VAL:HG13	1.67	0.72
21:DV:151:HIS:CD2	21:DV:169:GLU:O	2.42	0.72
1:AA:1429:G:H2'	1:AA:1430:C:C6	2.25	0.72
1:AA:1750:G:O2'	1:AA:1751:C:H5'	1.88	0.72
1:AA:776:G:H4'	1:AA:777:A:O5'	1.89	0.72
5:AF:178:PRO:HG2	5:AF:179:GLU:OE1	1.90	0.72
11:AO:63:PRO:HB3	30:A8:13:ARG:HG2	1.71	0.72
23:AZ:91:LYS:HG3	23:AZ:92:LYS:H	1.54	0.72
31:BA:1443:G:H3'	31:BA:1446:A:C5'	2.19	0.72
31:BA:407:G:O2'	34:BG:116:GLN:HG3	1.89	0.72
52:BC:7:A:H4'	52:BC:8:U:OP2	1.88	0.72
52:BD:41:C:C3'	52:BD:42:C:H5''	2.19	0.72
33:BF:47:LEU:HD23	33:BF:52:LEU:HD13	1.70	0.72
54:CA:1025:U:HO2'	54:CA:1026:G:H8	1.37	0.72
52:CC:44:G:H3'	52:CC:45:U:H6	1.51	0.72
47:CT:55:ASP:HA	47:CT:79:SER:HA	1.71	0.72
50:CW:50:GLU:HA	50:CW:100:ILE:HG21	1.70	0.72
17:D2:95:LEU:HD13	17:D2:97:LYS:HE3	1.69	0.72
55:DA:1718:G:C2'	55:DA:1725:G:H5''	2.19	0.72
55:DA:1963:U:O2	55:DA:1963:U:H2'	1.87	0.72
55:DA:654(S):G:H2'	55:DA:654(T):A:C8	2.25	0.72
55:DA:701:G:H2'	55:DA:702:G:H5''	1.70	0.72
56:DJ:14:GLN:HA	56:DJ:16:THR:N	2.04	0.72
8:DK:92:VAL:HG13	8:DK:120:ILE:CG2	2.17	0.72
9:DM:22:THR:HG22	9:DM:23:LEU:N	2.03	0.72
55:DA:1012:U:H5	9:DM:28:THR:HG21	1.55	0.72
12:DP:109:VAL:HG13	12:DP:113:GLN:HB3	1.70	0.72
57:DY:135:ARG:NH1	57:DY:138:LEU:HG	2.00	0.72
13:A0:37:THR:CG2	13:A0:39:PRO:HD2	2.19	0.72
1:AA:1926:U:H2'	1:AA:1928:A:OP2	1.89	0.72
1:AA:2129:C:H2'	1:AA:2130:U:H5'	1.70	0.72
1:AA:2693:A:H2'	1:AA:2694:G:H8	1.54	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:247:G:H4'	1:AA:386:G:C5	2.24	0.72
1:AA:886:C:H4'	1:AA:886:C:OP1	1.89	0.72
1:AA:2311:A:C8	6:AG:82:LEU:HD11	2.24	0.72
18:AS:80:PRO:O	18:AS:100:THR:HG22	1.88	0.72
21:AV:115:GLY:N	21:AV:177:PRO:CB	2.52	0.72
31:BA:765:G:H1	31:BA:812:C:H2'	1.55	0.72
50:BW:8:ARG:HH11	50:BW:8:ARG:HG3	1.54	0.72
54:CA:1513:A:H2'	54:CA:1514:C:C6	2.24	0.72
54:CA:557:G:H2'	54:CA:558:G:C8	2.24	0.72
52:CC:47:U:H2'	52:CC:47:U:O2	1.88	0.72
34:CG:129:ASN:N	34:CG:145:GLU:HB2	2.05	0.72
39:CL:111:ARG:HD2	44:CQ:61:TRP:OXT	1.89	0.72
49:CV:81:ARG:CG	49:CV:82:GLY:H	2.02	0.72
55:DA:1535:U:O2	55:DA:1535:U:H3'	1.90	0.72
55:DA:547:A:H2'	55:DA:548:A:C8	2.25	0.72
58:DL:18:THR:HG23	58:DL:42:ASN:ND2	2.03	0.72
21:DV:148:ASP:O	21:DV:173:ALA:HA	1.89	0.72
21:DV:190:GLU:O	21:DV:191:VAL:CB	2.37	0.72
21:DV:194:PRO:CG	21:DV:196:VAL:HG11	2.18	0.72
57:DY:134:LEU:O	57:DY:137:GLU:HG2	1.89	0.72
57:DY:87:VAL:HG13	57:DY:91:LYS:CG	2.19	0.72
1:AA:1225:C:C4'	17:A2:85:LYS:HB2	2.18	0.72
28:A6:38:LYS:HA	28:A6:48:VAL:O	1.90	0.72
1:AA:1171:G:O2'	1:AA:1173:G:O4'	2.08	0.72
1:AA:139:G:N2	1:AA:1596:A:H4'	2.05	0.72
1:AA:686:G:H5'	29:A7:11:LYS:HE2	1.71	0.72
1:AA:910:A:C5	12:AP:13:GLN:HG3	2.24	0.72
3:AD:26:LYS:H	3:AD:26:LYS:HD2	1.53	0.72
6:AG:67:LYS:HD2	6:AG:67:LYS:N	2.05	0.72
9:AM:42:TRP:HA	9:AM:48:MET:HE1	1.69	0.72
20:AU:40:GLU:HA	20:AU:64:GLU:OE1	1.90	0.72
24:AW:17:SER:CB	24:AW:18:PRO:HA	2.20	0.72
31:BA:409:G:OP1	34:BG:24:GLU:CG	2.36	0.72
40:BM:33:GLN:H	40:BM:75:ILE:CD1	2.02	0.72
31:BA:537:G:H5''	42:BO:113:ARG:NH1	2.05	0.72
42:BO:40:VAL:HG21	42:BO:78:GLN:HA	1.71	0.72
46:BS:18:ARG:HD3	46:BS:35:LYS:HD2	1.72	0.72
54:CA:1490:C:O2'	54:CA:1491:G:H5'	1.89	0.72
33:CF:42:LEU:HD11	33:CF:46:GLU:OE2	1.89	0.72
45:CR:74:ASP:HB3	45:CR:77:ARG:HG2	1.71	0.72
54:CA:625:G:H4'	46:CS:16:HIS:CD2	2.23	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:CV:44:MET:O	49:CV:62:ILE:HG21	1.90	0.72
26:D4:69:LYS:CD	26:D4:70:GLY:N	2.52	0.72
55:DA:1496:A:H8	55:DA:1577:C:HO2'	1.35	0.72
55:DA:26:G:H1'	55:DA:514:A:N6	2.05	0.72
2:DB:75:G:H5'	2:DB:75:G:C8	2.24	0.72
56:DI:24:ILE:O	56:DI:27:LEU:HB2	1.89	0.72
56:DI:26:ALA:O	56:DI:27:LEU:C	2.27	0.72
11:DO:138:LEU:C	11:DO:140:ALA:H	1.93	0.72
11:DO:71:VAL:CG1	11:DO:72:PRO:HD3	2.19	0.72
2:DB:52:A:H62	14:DQ:33:LYS:CG	2.02	0.72
20:DU:63:LYS:HA	20:DU:63:LYS:HZ3	1.55	0.72
21:DV:111:VAL:CG2	21:DV:146:ILE:HG13	2.20	0.72
57:DY:52:PHE:C	57:DY:53:VAL:HG22	2.10	0.72
57:DY:94:VAL:O	57:DY:94:VAL:HG12	1.89	0.72
28:A6:26:ASN:O	28:A6:28:ARG:HG2	1.88	0.72
1:AA:1047:G:H2'	1:AA:1110:G:N2	2.05	0.72
1:AA:1309:G:H4'	29:A7:7:PRO:HB2	1.70	0.72
1:AA:1899:G:N2	1:AA:1902:C:C4	2.58	0.72
2:AB:12:C:H2'	22:A3:74:ARG:HB3	1.71	0.72
21:AV:146:ILE:C	21:AV:148:ASP:N	2.38	0.72
23:AZ:53:VAL:HB	23:AZ:58:ILE:HD12	1.72	0.72
31:BA:1278:U:H5''	31:BA:1279:A:O4'	1.90	0.72
34:BG:3:ARG:HB2	34:BG:3:ARG:HH21	1.54	0.72
35:BH:126:ARG:HA	35:BH:131:ILE:HD11	1.69	0.72
44:BQ:15:LYS:HZ3	44:BQ:16:PHE:H	1.37	0.72
49:BV:49:ILE:HD12	49:BV:49:ILE:N	2.04	0.72
54:CA:1027:C:H2'	54:CA:1028:C:C5	2.25	0.72
54:CA:105:G:H2'	54:CA:106:C:C6	2.24	0.72
54:CA:33:A:H2'	54:CA:34:C:H6	1.55	0.72
52:CB:19:G:H22	52:CB:56:C:H42	1.37	0.72
52:CD:23:A:H2'	52:CD:24:G:C8	2.24	0.72
34:CG:129:ASN:HA	34:CG:145:GLU:HB2	1.70	0.72
36:CI:23:LYS:O	36:CI:27:GLN:HG2	1.89	0.72
45:CR:74:ASP:CG	45:CR:77:ARG:HD3	2.10	0.72
55:DA:1813:G:H1'	3:DD:50:THR:OG1	1.89	0.72
55:DA:1956:U:H1'	55:DA:2552:U:OP1	1.90	0.72
55:DA:27:G:H22	55:DA:512:G:H2'	1.53	0.72
5:DF:63:LYS:HA	5:DF:76:GLY:O	1.90	0.72
58:DL:13:PRO:HG2	58:DL:15:GLY:H	1.54	0.72
9:DM:14:VAL:HG13	9:DM:135:PRO:O	1.90	0.72
14:DQ:10:ARG:O	14:DQ:14:VAL:HG12	1.89	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:131:MET:O	57:DY:133:GLU:N	2.22	0.72
13:A0:104:ARG:HH12	13:A0:109:ALA:HB3	1.55	0.72
2:AB:40:U:C6	26:A4:1:MET:HE1	2.24	0.72
30:A8:49:VAL:HG12	30:A8:50:LEU:HD23	1.69	0.72
1:AA:1796:U:H2'	1:AA:1797:C:C6	2.25	0.72
1:AA:2286:A:H5''	1:AA:2287:A:O4'	1.89	0.72
2:AB:95:U:H2'	2:AB:96:G:C8	2.23	0.72
5:AF:4:VAL:HA	5:AF:19:GLU:HB3	1.71	0.72
15:AR:62:THR:CG2	15:AR:75:ILE:HG12	2.20	0.72
31:BA:1158:C:N3	31:BA:1160:G:N7	2.38	0.72
32:BE:16:HIS:CD2	32:BE:209:ARG:HD2	2.24	0.72
36:BI:77:ARG:NH1	36:BI:77:ARG:HB3	2.05	0.72
54:CA:736:C:OP1	48:CU:68:LYS:HE3	1.90	0.72
54:CA:872:A:O2'	54:CA:873:A:H3'	1.89	0.72
38:CK:116:LYS:CA	38:CK:116:LYS:HE2	2.20	0.72
55:DA:1179:C:C3'	55:DA:1180:C:H5''	2.20	0.72
55:DA:2311:A:H3'	55:DA:2312:U:C5	2.25	0.72
55:DA:2656:U:H5	55:DA:2664:G:N2	1.87	0.72
3:DD:70:TRP:CH2	3:DD:150:LYS:HA	2.25	0.72
6:DG:107:LEU:O	26:D4:38:LYS:CG	2.38	0.72
58:DL:60:TYR:OH	58:DL:66:THR:HG22	1.90	0.72
15:DR:90:GLN:HE21	15:DR:90:GLN:CA	1.91	0.72
55:DA:1075:C:H5''	21:DV:195:GLU:OE2	1.89	0.72
57:DY:116:ILE:O	57:DY:117:LEU:HB2	1.88	0.72
1:AA:13:A:H5''	1:AA:14:A:OP1	1.90	0.72
1:AA:1899:G:H22	1:AA:1902:C:H41	0.85	0.72
1:AA:2735:G:H2'	1:AA:2736:G:H8	1.54	0.72
4:AE:108:SER:HB3	4:AE:165:VAL:HG21	1.70	0.72
4:AE:66:HIS:C	4:AE:68:ALA:H	1.93	0.72
24:AW:68:ARG:HH11	24:AW:68:ARG:HG3	1.52	0.72
52:BC:39:U:H2'	52:BC:40:C:H6	1.53	0.72
33:BF:100:ALA:O	33:BF:101:LEU:HB2	1.89	0.72
34:BG:12:CYS:HB3	34:BG:21:LEU:HD22	1.70	0.72
39:BL:43:ALA:HA	39:BL:74:ILE:HD13	1.72	0.72
49:BV:76:PRO:HB2	49:BV:78:ARG:HD3	1.72	0.72
54:CA:1352:C:H2'	54:CA:1353:G:C8	2.25	0.72
54:CA:152:A:H62	54:CA:169:C:N4	1.88	0.72
54:CA:17:U:H2'	54:CA:18:C:C6	2.25	0.72
54:CA:96:G:H5'	54:CA:96:G:H8	1.54	0.72
52:CD:9:A:N6	52:CD:23:A:H62	1.87	0.72
32:CE:5:ILE:HD12	32:CE:224:GLN:HG2	1.72	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:DF:24:LEU:HD23	5:DF:115:ALA:HA	1.71	0.72
55:DA:671:C:OP1	11:DO:42:SER:O	2.08	0.72
21:DV:140:ASP:CG	21:DV:141:VAL:H	1.91	0.72
22:A3:55:ARG:HB3	22:A3:55:ARG:NH1	2.05	0.71
28:A6:9:LEU:HD13	28:A6:11:LEU:HD21	1.72	0.71
1:AA:2189:U:C3'	1:AA:2190:G:H5''	2.20	0.71
18:AS:9:TYR:H	18:AS:102:HIS:HD2	1.37	0.71
31:BA:1394:A:H5''	31:BA:1395:C:OP2	1.90	0.71
32:BE:98:LEU:O	32:BE:101:MET:HG2	1.89	0.71
37:BJ:16:LEU:HD12	39:BL:42:ARG:HA	1.72	0.71
47:BT:3:LYS:HB3	47:BT:61:GLU:HB3	1.72	0.71
35:CH:53:LEU:O	35:CH:57:LYS:HG2	1.89	0.71
36:CI:100:ASN:OD1	48:CU:27:GLY:HA2	1.90	0.71
22:D3:3:HIS:O	22:D3:4:LYS:HB2	1.89	0.71
22:D3:68:GLU:HG2	22:D3:80:HIS:HB2	1.71	0.71
26:D4:60:GLN:C	26:D4:61:ARG:HD2	2.11	0.71
26:D4:58:ARG:CA	26:D4:62:ARG:HB3	2.17	0.71
55:DA:1013:C:O2'	55:DA:1014:U:H5'	1.90	0.71
55:DA:1318:C:C2'	55:DA:1319:G:H5''	2.19	0.71
55:DA:2061:G:H5''	55:DA:2503:A:C2	2.25	0.71
57:DY:142:LEU:CG	57:DY:143:GLN:H	1.96	0.71
57:DY:89:ALA:HB3	56:DJ:15:ALA:HB1	0.72	0.71
22:A3:72:ARG:HH11	22:A3:72:ARG:HG3	1.56	0.71
1:AA:1048:A:H2	1:AA:1112:G:H21	1.35	0.71
1:AA:654(J):A:O2'	1:AA:654(K):C:O5'	2.07	0.71
5:AF:3:GLU:O	5:AF:19:GLU:HB2	1.89	0.71
7:AH:83:TYR:HA	7:AH:134:SER:HB3	1.71	0.71
7:AH:92:ILE:CD1	7:AH:92:ILE:H	2.03	0.71
8:AK:110:ASP:OD2	8:AK:113:ARG:HB2	1.89	0.71
11:AO:75:ILE:CD1	11:AO:75:ILE:H	1.92	0.71
21:AV:14:LYS:O	21:AV:17:ALA:HB3	1.89	0.71
21:AV:60:GLU:HG3	21:AV:61:LEU:N	2.03	0.71
54:CA:1024:G:H3'	54:CA:1025:U:H5''	1.72	0.71
54:CA:1027:C:H2'	54:CA:1028:C:C6	2.24	0.71
34:CG:22:LYS:HB2	34:CG:26:CYS:HB2	1.71	0.71
45:CR:82:ILE:HD11	45:CR:88:ARG:CB	2.18	0.71
48:CU:25:THR:C	48:CU:26:LEU:HD23	2.10	0.71
26:D4:46:GLN:CG	26:D4:48:ARG:HG2	2.20	0.71
55:DA:1084:A:H5'	55:DA:1085:A:OP2	1.90	0.71
55:DA:2475:C:N4	55:DA:2529:G:H1	1.86	0.71
55:DA:481:G:H1'	55:DA:506:G:H21	1.54	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:DD:182:LEU:H	3:DD:272:ALA:HB3	1.55	0.71
3:DD:69:ARG:HH21	3:DD:192:THR:HB	1.55	0.71
4:DE:55:ASN:C	4:DE:57:LYS:H	1.90	0.71
7:DH:87:LEU:HD13	7:DH:148:ILE:HG21	1.72	0.71
58:DL:10:LEU:HD21	58:DL:55:VAL:HG11	1.71	0.71
58:DL:57:ILE:CD1	58:DL:58:THR:N	2.40	0.71
58:DL:76:TYR:O	58:DL:78:ILE:N	2.22	0.71
12:DP:88:GLY:O	12:DP:90:VAL:N	2.23	0.71
18:DS:80:PRO:O	18:DS:100:THR:HG22	1.89	0.71
21:DV:10:ARG:NH2	21:DV:26:GLY:H	1.87	0.71
21:DV:24:LEU:HD21	21:DV:86:VAL:CG2	2.20	0.71
57:DY:25:PHE:CG	57:DY:82:PHE:CZ	2.77	0.71
13:A0:2:ARG:HH11	13:A0:2:ARG:HG3	1.56	0.71
1:AA:1880:C:H6	1:AA:1880:C:H5'	1.54	0.71
1:AA:1786:A:C2	1:AA:2606:C:H1'	2.25	0.71
1:AA:415:A:H2'	1:AA:416:C:C6	2.25	0.71
1:AA:5:A:O4'	1:AA:5:A:P	2.47	0.71
1:AA:959:A:H62	12:AP:82:ARG:NH2	1.88	0.71
14:AQ:23:ARG:HG2	14:AQ:23:ARG:HH11	1.54	0.71
20:AU:39:VAL:HG23	20:AU:40:GLU:H	1.55	0.71
31:BA:216:G:O2'	31:BA:217:C:O5'	2.08	0.71
33:BF:27:LYS:HD3	33:BF:28:GLN:OE1	1.89	0.71
54:CA:1323:G:H2'	54:CA:1324:A:C8	2.25	0.71
54:CA:524:G:H2'	54:CA:525:C:C6	2.26	0.71
54:CA:652:U:H1'	54:CA:653:A:H2	1.54	0.71
35:CH:81:GLU:HG2	35:CH:90:VAL:HG13	1.71	0.71
13:D0:12:ARG:HG3	13:D0:12:ARG:HH11	1.53	0.71
55:DA:1079:C:H2'	55:DA:1080:A:O4'	1.90	0.71
3:DD:66:ASP:OD2	3:DD:69:ARG:HG2	1.90	0.71
7:DH:101:ARG:HG2	7:DH:117:PRO:HG3	1.71	0.71
8:DK:25:TYR:HE2	8:DK:29:TYR:HD2	1.37	0.71
58:DL:124:ALA:O	58:DL:126:MET:HE3	1.90	0.71
9:DM:58:ASP:H	9:DM:60:ILE:CD1	2.03	0.71
55:DA:870:A:OP1	12:DP:6:ARG:HG2	1.90	0.71
57:DY:94:VAL:O	57:DY:95:GLN:HB2	1.90	0.71
1:AA:2103:C:H2'	1:AA:2104:G:C8	2.25	0.71
7:AH:86:GLU:HG3	7:AH:165:ALA:HB2	1.71	0.71
1:AA:2415:G:O3'	11:AO:66:GLY:HA3	1.91	0.71
31:BA:986:A:H2'	31:BA:987:G:C8	2.25	0.71
50:BW:56:MET:HG3	50:BW:84:LEU:CD1	2.21	0.71
51:BX:9:ARG:HH21	51:BX:10:ARG:HE	1.38	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1508:G:H2'	54:CA:1509:C:C6	2.26	0.71
54:CA:486:U:H2'	54:CA:487:A:C8	2.25	0.71
55:DA:2355:C:C4'	22:D3:36:ILE:HD11	2.20	0.71
55:DA:481:G:H1'	55:DA:506:G:N2	2.05	0.71
56:DJ:7:ARG:HH11	56:DJ:7:ARG:CG	2.02	0.71
9:DM:62:VAL:HG13	9:DM:66:LYS:HD2	1.71	0.71
21:DV:177:PRO:O	21:DV:178:GLU:CB	2.36	0.71
1:AA:1251:C:H4'	1:AA:1252:G:OP1	1.91	0.71
1:AA:2267:A:H5''	1:AA:2268:A:C5'	2.20	0.71
1:AA:363(A):A:C2'	1:AA:363(B):G:H5''	2.20	0.71
10:AN:87:ILE:HG21	10:AN:91:LEU:HA	1.72	0.71
11:AO:64:LYS:HE3	30:A8:30:ARG:NH2	2.04	0.71
1:AA:969:U:OP1	25:AX:17:LYS:HG2	1.91	0.71
31:BA:1032:A:H3'	31:BA:1032(A):G:C5'	2.21	0.71
31:BA:366:C:O2'	31:BA:367:U:P	2.48	0.71
31:BA:47:C:H4'	31:BA:48:C:O5'	1.90	0.71
34:BG:63:LYS:HD2	34:BG:198:VAL:HG12	1.73	0.71
37:BJ:23:VAL:HG13	37:BJ:43:PHE:CE2	2.26	0.71
31:BA:875:C:O2'	38:BK:14:ARG:HD2	1.90	0.71
48:BU:31:LEU:HG	48:BU:65:ILE:HD13	1.72	0.71
54:CA:713:G:N2	54:CA:777:A:H1'	2.06	0.71
40:CM:6:ILE:HG22	40:CM:98:ILE:HG13	1.71	0.71
45:CR:7:GLU:O	45:CR:11:VAL:HG23	1.91	0.71
47:CT:77:VAL:O	47:CT:78:GLU:HB2	1.90	0.71
55:DA:1188:U:O2'	55:DA:1189:A:H5'	1.91	0.71
55:DA:1332:G:C2	55:DA:1609:A:H2'	2.25	0.71
55:DA:1955:U:O2'	55:DA:1956:U:H5'	1.90	0.71
55:DA:2308:G:N1	55:DA:2311:A:H2	1.89	0.71
55:DA:607:U:O4	55:DA:608:A:N7	2.24	0.71
55:DA:883:G:O5'	55:DA:883:G:H8	1.70	0.71
6:DG:146:TYR:O	6:DG:149:VAL:HG22	1.90	0.71
7:DH:154:PRO:HG2	7:DH:162:ILE:O	1.91	0.71
12:DP:20:ALA:H	21:DV:79:ARG:HH22	1.36	0.71
12:DP:86:GLY:C	12:DP:88:GLY:H	1.94	0.71
57:DY:138:LEU:HD12	57:DY:139:VAL:N	2.04	0.71
57:DY:8:GLU:O	57:DY:11:ALA:HB3	1.90	0.71
16:A1:92:ARG:HB2	17:A2:11:GLN:HE22	1.53	0.71
30:A8:62:LEU:HB3	30:A8:63:PRO:HD3	1.72	0.71
1:AA:1056:G:H4'	1:AA:1086:A:C1'	2.19	0.71
1:AA:993:G:OP1	16:A1:50:ARG:NH2	2.24	0.71
3:AD:35:LYS:NZ	3:AD:65:ILE:HA	2.04	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:AH:125:VAL:HG22	7:AH:126:PRO:HB3	1.73	0.71
10:AN:47:ILE:CG1	10:AN:48:PRO:HD2	2.21	0.71
18:AS:14:PRO:HB3	18:AS:18:ARG:HH21	1.55	0.71
31:BA:707:C:H2'	31:BA:708:C:H6	1.56	0.71
34:BG:127:THR:HG21	34:BG:149:ALA:HB2	1.72	0.71
38:BK:103:VAL:CG2	38:BK:110:ALA:HB2	2.21	0.71
39:BL:37:PHE:HB3	39:BL:43:ALA:HB1	1.72	0.71
42:BO:46:LYS:HG2	42:BO:47:LYS:N	2.04	0.71
49:BV:17:GLU:O	49:BV:21:GLU:HG2	1.91	0.71
49:BV:76:PRO:HB2	49:BV:78:ARG:NH1	2.05	0.71
54:CA:1363:A:H1'	54:CA:1365:G:C5	2.25	0.71
54:CA:112:G:H5'	54:CA:389:A:H4'	1.72	0.71
54:CA:658:G:H2'	54:CA:659:U:H6	1.55	0.71
54:CA:827:U:H5'	54:CA:828:A:OP2	1.91	0.71
32:CE:40:HIS:C	32:CE:41:ILE:HD12	2.11	0.71
33:CF:105:GLU:HG2	33:CF:106:VAL:H	1.54	0.71
36:CI:69:GLU:O	36:CI:72:VAL:HG12	1.90	0.71
48:CU:18:ARG:H	48:CU:18:ARG:HD3	1.54	0.71
36:CI:99:ALA:HB3	48:CU:29:PHE:HE2	1.55	0.71
16:D1:95:LEU:CD1	17:D2:11:GLN:HE21	2.02	0.71
55:DA:1112:G:H2'	55:DA:1113:U:C6	2.25	0.71
55:DA:654(K):C:H2'	55:DA:654(L):G:C8	2.24	0.71
55:DA:2785:C:O2'	4:DE:64:LYS:HD3	1.90	0.71
6:DG:151:ALA:HB3	6:DG:153:ARG:NH1	2.05	0.71
8:DK:77:LEU:O	8:DK:78:THR:HG23	1.91	0.71
20:DU:20:TYR:CE1	20:DU:42:VAL:HA	2.26	0.71
1:AA:270(E):G:H2'	1:AA:270(F):U:C6	2.26	0.71
1:AA:484:C:H2'	1:AA:485:C:C6	2.24	0.71
1:AA:686:G:H21	1:AA:788:A:H61	1.38	0.71
4:AE:137:HIS:HB3	4:AE:138:PRO:HD2	1.70	0.71
11:AO:52:GLU:CD	11:AO:54:GLY:H	1.94	0.71
1:AA:907:U:O2'	12:AP:101:ARG:NH2	2.24	0.71
21:AV:116:VAL:H	21:AV:177:PRO:HG3	1.55	0.71
1:AA:61:G:H5'	24:AW:50:ILE:HG12	1.72	0.71
23:AZ:86:SER:H	23:AZ:87:PRO:CD	2.04	0.71
23:AZ:7:ILE:HG12	23:AZ:91:LYS:NZ	2.06	0.71
31:BA:1067:A:O2'	31:BA:1068:G:H8	1.73	0.71
31:BA:353:A:H8	31:BA:353:A:H5'	1.54	0.71
31:BA:366:C:C4'	31:BA:367:U:OP1	2.36	0.71
31:BA:887:G:C2'	31:BA:888:G:C5'	2.61	0.71
22:A3:7:LEU:HA	52:BC:2:C:C5'	2.20	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:BI:55:ASP:OD1	36:BI:56:PRO:HD2	1.91	0.71
41:BN:21:ILE:HD13	41:BN:94:ALA:HB1	1.71	0.71
6:AG:115:ARG:NH1	43:BP:7:VAL:HG21	2.05	0.71
49:BV:16:LEU:HA	49:BV:19:VAL:HB	1.73	0.71
53:C1:34:G:H2'	53:C1:35:A:C8	2.26	0.71
38:CK:25:ASP:HA	38:CK:59:LEU:O	1.90	0.71
16:D1:50:ARG:HG2	16:D1:53:ARG:HH21	1.55	0.71
55:DA:1047:G:H2'	55:DA:1110:G:H22	1.54	0.71
55:DA:2469:A:N1	55:DA:2481:G:N3	2.39	0.71
3:DD:62:TYR:HA	3:DD:87:ASN:ND2	2.06	0.71
56:DJ:1:MET:SD	56:DJ:2:ALA:CB	2.79	0.71
58:DL:90:LYS:N	58:DL:91:PRO:HA	2.06	0.71
20:DU:84:ARG:HH12	20:DU:97:ARG:CB	2.04	0.71
21:DV:19:ARG:NH1	21:DV:84:GLU:O	2.23	0.71
1:AA:1286:A:H2'	1:AA:1288:U:OP2	1.90	0.71
1:AA:2344:U:OP1	28:A6:38:LYS:HE3	1.90	0.71
1:AA:669:G:N3	1:AA:669:G:H2'	2.06	0.71
8:AK:72:LEU:HD21	8:AK:107:VAL:HG21	1.72	0.71
10:AN:87:ILE:CG2	10:AN:91:LEU:HA	2.20	0.71
14:AQ:106:ARG:O	14:AQ:107:GLU:HB3	1.90	0.71
31:BA:689:C:H2'	31:BA:690:G:H5'	1.72	0.71
52:BD:15:G:H1	52:BD:48:C:N4	1.89	0.71
33:BF:73:PRO:O	33:BF:76:VAL:HG22	1.90	0.71
48:BU:43:PHE:HA	48:BU:51:LEU:HD12	1.72	0.71
53:C1:29:G:H2'	53:C1:30:C:C4	2.25	0.71
54:CA:1305:G:HO2'	54:CA:1306:A:H8	1.37	0.71
54:CA:1321:C:H5'	54:CA:1322:C:H5''	1.72	0.71
54:CA:188:U:H2'	54:CA:189:U:C5'	2.19	0.71
54:CA:38:G:C2	54:CA:397:A:C2	2.73	0.71
39:CL:45:ALA:O	39:CL:48:GLU:HG2	1.90	0.71
54:CA:523:A:H61	42:CO:92:ASP:HB2	1.55	0.71
45:CR:39:LEU:HD12	45:CR:59:MET:CE	2.20	0.71
48:CU:43:PHE:HE2	48:CU:58:LEU:HD11	1.55	0.71
13:D0:56:LYS:HE2	13:D0:94:TYR:CZ	2.26	0.71
17:D2:49:THR:OG1	17:D2:50:PRO:HD2	1.91	0.71
55:DA:196:A:H2'	55:DA:805:G:O6	1.89	0.71
55:DA:2030:A:H4'	55:DA:2031:A:H8	1.56	0.71
55:DA:2580:U:H4'	4:DE:130:GLY:CA	2.21	0.71
55:DA:631:A:OP2	30:D8:46:ARG:NH2	2.21	0.71
55:DA:658:C:H2'	55:DA:659:C:C6	2.26	0.71
55:DA:676:A:H2	55:DA:802:A:H61	1.37	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DB:81:G:N7	2:DB:96:G:N3	2.38	0.71
3:DD:69:ARG:NH2	3:DD:128:GLY:O	2.20	0.71
4:DE:101:ARG:HG2	4:DE:169:ASN:OD1	1.90	0.71
56:DJ:6:GLU:C	56:DJ:10:GLU:HG2	2.11	0.71
9:DM:43:THR:HG22	9:DM:45:ASN:ND2	2.06	0.71
9:DM:96:GLU:O	9:DM:98:VAL:N	2.24	0.71
57:DY:21:GLN:NE2	57:DY:21:GLN:CA	2.52	0.71
22:A3:53:MET:HB3	22:A3:59:LEU:HD23	1.71	0.71
30:A8:50:LEU:CG	30:A8:51:ALA:H	1.94	0.71
1:AA:1799:G:N2	1:AA:1818:U:O2'	2.24	0.71
1:AA:1948:G:H5'	1:AA:1948:G:C8	2.26	0.71
1:AA:270(Z):U:HO2'	1:AA:271(A):C:H5	1.38	0.71
1:AA:2790:A:H1'	1:AA:2893:G:O2'	1.90	0.71
1:AA:454:A:H4'	1:AA:455:C:OP2	1.90	0.71
1:AA:894:C:H3'	1:AA:895:U:C6	2.14	0.71
1:AA:99:U:O2	1:AA:99:U:H2'	1.89	0.71
3:AD:147:LEU:CD2	3:AD:155:LEU:HD11	2.19	0.71
4:AE:10:GLY:O	4:AE:11:MET:HB2	1.90	0.71
9:AM:112:LEU:HA	9:AM:115:ARG:HB2	1.73	0.71
10:AN:111:PHE:HB3	10:AN:114:ILE:HD12	1.73	0.71
12:AP:10:ARG:O	12:AP:11:LYS:HB2	1.89	0.71
19:AT:67:GLY:O	19:AT:69:TYR:N	2.23	0.71
31:BA:279:A:C4'	31:BA:280:C:H5''	2.16	0.71
37:BJ:108:ALA:O	37:BJ:111:ARG:HG3	1.89	0.71
47:BT:59:ILE:HD13	47:BT:73:VAL:HA	1.73	0.71
54:CA:1028(A):C:H2'	54:CA:1028(B):C:C6	2.26	0.71
54:CA:105:G:H2'	54:CA:106:C:H6	1.54	0.71
54:CA:1218:C:H2'	54:CA:1219:U:C6	2.26	0.71
54:CA:192:U:H4'	50:CW:102:GLY:O	1.90	0.71
54:CA:498:A:O2'	54:CA:500:G:O4'	2.09	0.71
54:CA:686:U:H1'	41:CN:42:TRP:HE1	1.54	0.71
45:CR:82:ILE:HG23	45:CR:83:GLU:N	2.05	0.71
46:CS:19:ILE:HG22	46:CS:36:ILE:HG13	1.71	0.71
28:D6:48:VAL:O	28:D6:49:HIS:HB2	1.89	0.71
55:DA:1359:A:H2'	55:DA:1360:A:C5'	2.20	0.71
55:DA:229:A:N6	55:DA:417:C:O2'	2.19	0.71
55:DA:270(K):C:H2'	55:DA:270(L):U:H5''	1.72	0.71
55:DA:284:U:H2'	55:DA:285:C:C6	2.26	0.71
3:DD:30:GLU:HG3	3:DD:63:ARG:NE	2.05	0.71
4:DE:50:GLY:HA3	4:DE:74:PRO:HG3	1.73	0.71
6:DG:21:ARG:HH11	6:DG:21:ARG:HG2	1.55	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:DJ:15:ALA:C	56:DJ:16:THR:CG2	2.59	0.71
56:DJ:20:LEU:HD12	56:DJ:20:LEU:N	2.06	0.71
15:DR:123:GLN:O	15:DR:125:ARG:N	2.23	0.71
19:DT:40:LYS:O	19:DT:43:VAL:HG12	1.90	0.71
17:A2:5:VAL:HG23	17:A2:37:VAL:CG1	2.20	0.71
6:AG:109:VAL:HA	26:A4:37:SER:CB	2.19	0.71
1:AA:1673:U:C2'	1:AA:1674:G:H5'	2.21	0.71
1:AA:2741:A:H61	1:AA:2763:G:H2'	1.55	0.71
1:AA:845:G:H21	1:AA:933:A:H61	1.38	0.71
1:AA:860:U:O2'	1:AA:861:A:H5'	1.91	0.71
3:AD:76:PRO:HG2	3:AD:98:VAL:CG2	2.20	0.71
5:AF:31:HIS:HB2	11:AO:9:ASN:ND2	2.05	0.71
11:AO:80:TYR:CD1	11:AO:111:ARG:HB3	2.26	0.71
31:BA:448:A:OP2	31:BA:485:G:N2	2.23	0.71
31:BA:965:A:H4'	31:BA:966:G:OP1	1.89	0.71
32:BE:115:LEU:HD13	32:BE:145:LEU:HD12	1.73	0.71
42:BO:89:ARG:HG2	42:BO:90:VAL:H	1.55	0.71
49:BV:19:VAL:HG13	49:BV:44:MET:HB3	1.71	0.71
54:CA:403:C:O2'	54:CA:404:U:H5'	1.91	0.71
52:CB:9:A:O2'	52:CB:10:G:OP1	2.08	0.71
32:CE:60:ASP:O	32:CE:64:ARG:HG2	1.91	0.71
34:CG:173:TRP:CD1	34:CG:174:LEU:HG	2.26	0.71
55:DA:1280:G:H2'	55:DA:1281:G:H5'	1.72	0.71
55:DA:1385:G:H4'	55:DA:1386:C:OP1	1.91	0.71
55:DA:2189:U:C2'	55:DA:2190:G:H5''	2.21	0.71
55:DA:482:A:H4'	20:DU:47:LYS:HD2	1.73	0.71
3:DD:35:LYS:HD3	3:DD:63:ARG:HB3	1.73	0.71
56:DI:30:ALA:CA	56:DJ:3:LEU:HD21	2.20	0.71
8:DK:60:GLU:HG3	8:DK:61:ARG:HH12	1.56	0.71
58:DL:132:ARG:C	58:DL:137:GLU:OE2	2.29	0.71
21:DV:191:VAL:O	21:DV:192:ALA:HB2	1.91	0.71
57:DY:24:PHE:C	57:DY:24:PHE:HD2	1.94	0.71
30:A8:38:GLY:O	30:A8:41:ILE:HG22	1.91	0.70
1:AA:1049:C:N4	7:AH:2:SER:HB2	2.05	0.70
1:AA:1161:C:H1'	17:A2:8:GLY:O	1.90	0.70
1:AA:655:A:H2'	1:AA:656:G:H5'	1.73	0.70
6:AG:121:ASN:HD22	6:AG:122:PRO:CD	2.04	0.70
14:AQ:15:ARG:O	14:AQ:19:LYS:HD3	1.91	0.70
15:AR:11:GLU:N	15:AR:11:GLU:OE1	2.23	0.70
21:AV:28:MET:HG3	21:AV:37:VAL:HG11	1.73	0.70
25:AX:52:HIS:HD2	25:AX:52:HIS:H	1.38	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:628:G:H2'	31:BA:629:G:H8	1.55	0.70
32:BE:188:ALA:HB1	32:BE:192:SER:HB2	1.72	0.70
39:BL:111:ARG:HG2	39:BL:112:LYS:N	2.04	0.70
44:BQ:26:ARG:HD3	44:BQ:43:CYS:HB3	1.71	0.70
53:C1:36:G:C3'	53:C1:37:G:H5''	2.20	0.70
54:CA:1116:C:H2'	54:CA:1117:G:C5'	2.20	0.70
54:CA:794:A:H2'	54:CA:795:C:C6	2.26	0.70
32:CE:121:LEU:O	32:CE:121:LEU:HD23	1.89	0.70
49:CV:64:GLU:O	49:CV:67:VAL:HG23	1.90	0.70
13:D0:67:LEU:CD1	13:D0:76:VAL:HG21	2.21	0.70
6:DG:112:PRO:HA	26:D4:37:SER:HB2	1.73	0.70
28:D6:40:CYS:SG	28:D6:45:LYS:HD3	2.30	0.70
30:D8:23:VAL:HG12	30:D8:46:ARG:HB3	1.71	0.70
55:DA:1058:U:C2'	55:DA:1059:G:C8	2.68	0.70
55:DA:1082:U:O3'	58:DL:117:THR:CG2	2.38	0.70
55:DA:1682:G:H2'	55:DA:1683:C:C6	2.26	0.70
55:DA:639:U:H2'	55:DA:640:C:C6	2.26	0.70
56:DJ:18:LEU:HA	56:DJ:21:LYS:CB	2.19	0.70
56:DJ:20:LEU:H	56:DJ:20:LEU:CD1	2.04	0.70
55:DA:1666:G:H4'	10:DN:6:THR:HG23	1.73	0.70
14:DQ:60:GLY:O	14:DQ:61:ASN:HB2	1.90	0.70
15:DR:66:VAL:HA	15:DR:71:GLY:HA2	1.72	0.70
15:DR:98:LYS:HB3	15:DR:100:TYR:CE1	2.26	0.70
57:DY:134:LEU:C	57:DY:137:GLU:HG2	2.11	0.70
27:A5:40:LYS:NZ	27:A5:46:CYS:H	1.88	0.70
1:AA:1266:G:OP1	1:AA:1266:G:H4'	1.91	0.70
1:AA:1314:C:H5'	1:AA:1314:C:H6	1.56	0.70
1:AA:1899:G:N2	1:AA:1902:C:C5	2.58	0.70
1:AA:185:U:H4'	1:AA:218:A:H4'	1.71	0.70
1:AA:2461:C:H2'	1:AA:2462:U:C6	2.25	0.70
4:AE:8:LYS:HG2	4:AE:192:ASN:HA	1.73	0.70
6:AG:97:ASP:H	6:AG:100:TRP:HD1	1.36	0.70
8:AK:82:ARG:HG3	8:AK:82:ARG:HH11	1.54	0.70
9:AM:35:ARG:HB3	9:AM:42:TRP:HZ3	1.54	0.70
23:AZ:92:LYS:HZ2	23:AZ:92:LYS:HB3	1.56	0.70
31:BA:539:A:H2'	31:BA:540:G:C8	2.26	0.70
32:BE:208:ILE:HA	32:BE:211:ILE:HD12	1.72	0.70
32:BE:80:ILE:CD1	32:BE:211:ILE:HG22	2.20	0.70
32:BE:7:VAL:HG13	32:BE:8:LYS:HD3	1.73	0.70
34:BG:100:ARG:HH12	34:BG:137:SER:CB	2.03	0.70
38:BK:89:PRO:HA	38:BK:92:ARG:HH11	1.55	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:93:U:C2'	54:CA:95:G:H5''	2.20	0.70
34:CG:28:SER:HB3	34:CG:29:PRO:CD	2.20	0.70
40:CM:6:ILE:O	40:CM:6:ILE:HD12	1.91	0.70
42:CO:27:LEU:HD13	42:CO:28:LYS:H	1.54	0.70
49:CV:86:GLU:OE2	49:CV:86:GLU:HA	1.89	0.70
13:D0:33:ARG:HH22	27:D5:55:ARG:CG	1.96	0.70
27:D5:40:LYS:HG2	27:D5:46:CYS:HB3	1.73	0.70
30:D8:36:LYS:HB3	30:D8:40:GLU:CG	2.19	0.70
30:D8:60:LEU:O	30:D8:63:PRO:HD2	1.89	0.70
55:DA:1925:C:C6	55:DA:1925:C:C3'	2.63	0.70
55:DA:226:G:H1'	55:DA:228:A:N6	2.06	0.70
55:DA:270(F):U:H2'	55:DA:270(G):C:C6	2.25	0.70
55:DA:961:C:H5''	55:DA:962:G:OP2	1.91	0.70
8:DK:60:GLU:HG3	8:DK:61:ARG:NH1	2.07	0.70
58:DL:77:LEU:HD21	58:DL:111:LYS:NZ	2.07	0.70
21:DV:15:PRO:O	21:DV:19:ARG:HB2	1.91	0.70
57:DY:132:ASP:O	57:DY:134:LEU:CD2	2.38	0.70
16:A1:5:LYS:HB2	16:A1:5:LYS:NZ	2.07	0.70
1:AA:2092:U:N3	1:AA:2225:A:O2'	2.24	0.70
1:AA:2824:C:H2'	1:AA:2825:C:H5'	1.71	0.70
1:AA:704:G:H2'	1:AA:726:G:H22	1.56	0.70
7:AH:9:ILE:HD12	7:AH:49:VAL:HG11	1.73	0.70
11:AO:57:THR:HG23	11:AO:57:THR:O	1.91	0.70
12:AP:12:GLN:HE21	12:AP:73:PRO:CD	2.03	0.70
15:AR:12:SER:HB3	15:AR:15:VAL:HG13	1.71	0.70
20:AU:38:ILE:HG22	20:AU:66:PRO:CA	2.19	0.70
52:BD:48:C:H5	52:BD:59:U:H1'	1.55	0.70
35:BH:32:VAL:HG12	35:BH:33:VAL:N	2.04	0.70
43:BP:25:ILE:HG22	43:BP:26:GLY:N	2.06	0.70
43:BP:49:THR:HB	43:BP:52:GLU:HG3	1.73	0.70
49:BV:44:MET:HA	49:BV:47:HIS:CD2	2.25	0.70
54:CA:1468:A:H2'	54:CA:1469:G:O4'	1.91	0.70
54:CA:992:U:H1'	54:CA:993:G:C2	2.27	0.70
42:CO:104:VAL:HG12	42:CO:105:TYR:CD1	2.26	0.70
54:CA:375:U:OP1	46:CS:69:THR:HG21	1.90	0.70
28:D6:28:ARG:NH1	28:D6:28:ARG:HB3	2.05	0.70
55:DA:1043:C:C2'	55:DA:1044:G:H5''	2.22	0.70
55:DA:1360:A:C6	55:DA:1372:U:O4	2.43	0.70
55:DA:528:A:H2	55:DA:2043:C:H5'	1.55	0.70
55:DA:479:A:H4'	55:DA:480:A:OP1	1.90	0.70
4:DE:101:ARG:CZ	4:DE:171:GLU:HB2	2.21	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:DF:40:GLN:HE22	5:DF:184:TYR:H	1.37	0.70
7:DH:20:ALA:HB3	7:DH:21:PRO:HD2	1.73	0.70
58:DL:76:TYR:C	58:DL:78:ILE:N	2.41	0.70
57:DY:27:VAL:HA	57:DY:111:LEU:CD1	2.19	0.70
57:DY:27:VAL:CA	57:DY:111:LEU:H	2.03	0.70
57:DY:25:PHE:CG	57:DY:82:PHE:CG	2.79	0.70
1:AA:1928:A:C2'	1:AA:1929:G:H5''	2.20	0.70
1:AA:200:U:H4'	23:AZ:34:THR:HG22	1.72	0.70
1:AA:2123:G:O2'	1:AA:2124:G:H5'	1.92	0.70
1:AA:2354:G:O2'	22:A3:36:ILE:HD12	1.91	0.70
1:AA:654(K):C:H2'	1:AA:654(L):G:H8	1.56	0.70
7:AH:152:ARG:O	7:AH:154:PRO:HD3	1.91	0.70
12:AP:48:GLU:O	12:AP:52:VAL:HG12	1.91	0.70
20:AU:81:LYS:HD3	20:AU:97:ARG:CZ	2.21	0.70
35:BH:51:VAL:HB	35:BH:52:PRO:CD	2.19	0.70
47:BT:68:ARG:HH11	47:BT:68:ARG:HG2	1.56	0.70
54:CA:1128:C:H5'	39:CL:16:ARG:NH2	2.05	0.70
54:CA:1139:G:H4'	54:CA:1140:C:O5'	1.91	0.70
54:CA:1502:A:H5''	54:CA:1503:A:OP2	1.91	0.70
54:CA:194:C:H2'	54:CA:195:A:H5''	1.73	0.70
32:CE:31:TYR:O	32:CE:42:ILE:HG13	1.92	0.70
35:CH:12:LEU:HD23	35:CH:13:ILE:N	2.06	0.70
50:CW:23:ARG:O	50:CW:27:LYS:HB2	1.91	0.70
28:D6:20:ASN:HD22	28:D6:42:TRP:HH2	1.37	0.70
55:DA:1504:C:H5'	55:DA:1505:C:OP2	1.91	0.70
55:DA:1726:G:O2'	55:DA:1727:U:H5'	1.92	0.70
7:DH:117:PRO:HB3	7:DH:123:PHE:CE1	2.26	0.70
58:DL:141:ALA:CB	58:DL:142:PRO:HA	2.21	0.70
10:DN:112:MET:O	10:DN:115:VAL:HG22	1.92	0.70
57:DY:93:LEU:CD2	57:DY:126:ALA:CB	2.64	0.70
57:DY:141:VAL:CG1	57:DY:142:LEU:H	1.92	0.70
17:A2:49:THR:CB	17:A2:50:PRO:HD3	2.22	0.70
30:A8:48:PHE:C	30:A8:49:VAL:HG23	2.12	0.70
1:AA:946:G:HO2'	1:AA:947:G:C5'	2.03	0.70
3:AD:169:GLU:HG2	3:AD:174:ILE:HD11	1.72	0.70
1:AA:2820:A:H61	4:AE:192:ASN:CB	2.04	0.70
1:AA:2531:A:H4'	7:AH:157:TYR:CE2	2.27	0.70
12:AP:21:THR:HA	12:AP:98:LYS:HB2	1.74	0.70
21:AV:177:PRO:C	21:AV:178:GLU:HG3	2.11	0.70
31:BA:1176:A:H2'	31:BA:1177:G:H5'	1.74	0.70
31:BA:1330:U:H4'	43:BP:23:TYR:CE2	2.26	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:579:G:C5'	31:BA:728:A:H1'	2.21	0.70
34:CG:50:ARG:CZ	53:C1:57:U:O2'	2.40	0.70
54:CA:1126:U:H1'	54:CA:1280:A:N7	2.06	0.70
54:CA:702:A:H5'	54:CA:703:G:N7	2.07	0.70
34:CG:114:ARG:NH1	34:CG:114:ARG:HG3	2.07	0.70
45:CR:39:LEU:HD12	45:CR:59:MET:HE1	1.73	0.70
55:DA:2656:U:H5	55:DA:2664:G:H21	1.39	0.70
7:DH:4:ILE:HG13	7:DH:6:ARG:CD	2.21	0.70
58:DL:108:ALA:HA	58:DL:111:LYS:CE	2.21	0.70
10:DN:71:ARG:HG3	10:DN:71:ARG:HH11	1.55	0.70
12:DP:88:GLY:C	12:DP:90:VAL:N	2.43	0.70
14:DQ:27:SER:HA	14:DQ:88:ASP:HB3	1.73	0.70
21:DV:118:GLN:CA	21:DV:118:GLN:NE2	2.45	0.70
21:DV:76:LEU:CD2	21:DV:76:LEU:H	2.03	0.70
17:A2:82:ARG:HH11	17:A2:82:ARG:HG3	1.57	0.70
1:AA:1288:U:O2'	1:AA:1647:G:N2	2.24	0.70
1:AA:676:A:H8	1:AA:2069:G:H21	1.37	0.70
1:AA:265:A:O2'	1:AA:266:G:C4'	2.38	0.70
1:AA:34:C:HO2'	1:AA:35:G:H8	1.39	0.70
1:AA:415:A:H2'	1:AA:416:C:H6	1.56	0.70
3:AD:131:LEU:HD12	3:AD:131:LEU:N	2.07	0.70
12:AP:20:ALA:HB2	21:AV:79:ARG:NH2	2.07	0.70
31:BA:1241:G:H2'	31:BA:1242:C:C6	2.27	0.70
31:BA:399:G:H2'	31:BA:400:C:C6	2.26	0.70
31:BA:686:U:H2'	31:BA:687:A:C8	2.27	0.70
31:BA:922:G:H4'	35:BH:20:GLN:HA	1.74	0.70
38:BK:84:ARG:HH12	38:BK:86:ILE:HD13	1.55	0.70
47:BT:10:VAL:HG13	47:BT:19:VAL:HB	1.74	0.70
54:CA:1225:A:H5"	54:CA:1226:C:OP2	1.91	0.70
33:CF:148:GLY:HA3	33:CF:172:ARG:O	1.91	0.70
33:CF:70:VAL:HG12	33:CF:71:ALA:N	2.07	0.70
40:CM:5:ARG:HH21	40:CM:99:LYS:HD2	1.56	0.70
54:CA:1286:A:OP1	51:CX:26:LYS:HD2	1.90	0.70
6:DG:104:GLU:OE1	26:D4:23:GLU:HB3	1.91	0.70
55:DA:352:G:H5'	55:DA:353:G:OP2	1.92	0.70
6:DG:6:ALA:HB3	6:DG:104:GLU:OE2	1.91	0.70
8:DK:25:TYR:HE2	8:DK:29:TYR:CD2	2.09	0.70
58:DL:59:ILE:C	58:DL:60:TYR:CD1	2.64	0.70
57:DY:142:LEU:C	57:DY:142:LEU:HD22	2.10	0.70
57:DY:23:SER:OG	57:DY:24:PHE:N	2.22	0.70
57:DY:43:ALA:CA	57:DY:47:ASN:ND2	2.55	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:88:ALA:C	57:DY:92:THR:H	1.95	0.70
1:AA:2286:A:OP2	28:A6:28:ARG:CZ	2.39	0.70
1:AA:1829:A:H3'	1:AA:1830:C:H6	1.57	0.70
3:AD:62:TYR:CE1	3:AD:64:ILE:HA	2.27	0.70
4:AE:48:GLN:CG	4:AE:78:LEU:HD12	2.21	0.70
7:AH:85:LYS:O	7:AH:132:ARG:HB2	1.91	0.70
31:BA:1435:G:H2'	31:BA:1436:U:H6	1.56	0.70
31:BA:403:C:O2'	31:BA:404:U:H5'	1.91	0.70
31:BA:706:A:C1'	41:BN:29:ILE:HD11	2.22	0.70
52:BB:74:C:H1'	52:BB:75:C:H5'	1.74	0.70
38:BK:10:LEU:HD22	38:BK:83:ILE:HD11	1.72	0.70
45:BR:16:ALA:HB1	45:BR:21:ASP:HB3	1.72	0.70
54:CA:440:A:H3'	54:CA:442:C:H6	1.57	0.70
54:CA:564:C:H5'	47:CT:32:TYR:CE2	2.27	0.70
54:CA:652:U:H1'	54:CA:653:A:C2	2.27	0.70
52:CB:57:G:H2'	52:CB:58:A:H5''	1.74	0.70
35:CH:126:ARG:HG3	35:CH:126:ARG:HH11	1.56	0.70
40:CM:37:PRO:HA	40:CM:72:VAL:HG22	1.72	0.70
55:DA:1280:G:H2'	55:DA:1281:G:C5'	2.21	0.70
55:DA:1291:C:H2'	55:DA:1292:U:C6	2.26	0.70
3:DD:110:GLY:O	3:DD:112:GLN:HG3	1.91	0.70
4:DE:3:GLY:HA3	4:DE:81:ILE:HG13	1.72	0.70
56:DI:16:THR:HG23	56:DI:17:VAL:HG22	1.72	0.70
10:DN:93:PRO:HB3	10:DN:114:ILE:HD11	1.73	0.70
11:DO:138:LEU:HD12	11:DO:139:LYS:N	2.06	0.70
19:DT:47:PHE:O	19:DT:49:VAL:HG23	1.92	0.70
20:DU:63:LYS:HA	20:DU:63:LYS:NZ	2.07	0.70
20:DU:81:LYS:HB2	20:DU:96:ILE:CG2	2.21	0.70
21:DV:194:PRO:O	21:DV:195:GLU:C	2.29	0.70
57:DY:70:GLU:O	57:DY:71:LEU:CB	2.39	0.70
57:DY:92:THR:HG22	57:DY:93:LEU:N	2.07	0.70
1:AA:1403:C:H5''	1:AA:1471:A:H1'	1.72	0.70
3:AD:159:ALA:H	3:AD:196:VAL:HG11	1.56	0.70
7:AH:143:GLN:HE22	7:AH:147:ASN:ND2	1.89	0.70
7:AH:168:PRO:O	7:AH:169:VAL:HG12	1.91	0.70
21:AV:103:ARG:HE	21:AV:103:ARG:HA	1.57	0.70
21:AV:53:ILE:CG2	21:AV:71:VAL:HG13	2.20	0.70
31:BA:976:G:N2	31:BA:1362(A):C:OP2	2.20	0.70
31:BA:1452:C:O2	31:BA:1452:C:H2'	1.92	0.70
31:BA:448:A:H2'	31:BA:449:C:O2	1.92	0.70
52:BD:37:MIA:H2'	52:BD:38:A:O4'	1.92	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:BF:32:LEU:HB3	33:BF:59:ARG:NH1	2.05	0.70
49:BV:6:LYS:N	49:BV:6:LYS:HD2	2.07	0.70
54:CA:112:G:H4'	54:CA:389:A:H5''	1.73	0.70
54:CA:960:U:O2'	54:CA:961:U:OP2	2.06	0.70
32:CE:187:LEU:HD11	32:CE:204:ASN:O	1.91	0.70
38:CK:1:MET:H3	38:CK:1:MET:HE2	1.56	0.70
43:CP:57:ARG:HH21	26:D4:34:GLU:HB2	1.56	0.70
55:DA:1879:C:H2'	55:DA:1880:C:H5'	1.72	0.70
55:DA:1899:G:N2	55:DA:1902:C:N4	2.11	0.70
55:DA:562:U:O2'	55:DA:572:A:O4'	2.09	0.70
57:DY:88:ALA:O	56:DJ:15:ALA:HB2	1.92	0.70
58:DL:110:GLN:HG3	58:DL:111:LYS:HE3	1.72	0.70
58:DL:98:ARG:N	58:DL:98:ARG:HH11	1.89	0.70
21:DV:175:VAL:HB	21:DV:176:PRO:HA	1.72	0.70
21:DV:192:ALA:C	21:DV:193:GLU:OE1	2.30	0.70
57:DY:43:ALA:HB3	57:DY:47:ASN:HA	1.73	0.70
22:A3:53:MET:HB3	22:A3:59:LEU:CD2	2.22	0.70
29:A7:46:VAL:C	29:A7:47:ARG:HD3	2.12	0.70
4:AE:8:LYS:HB3	4:AE:192:ASN:HA	1.74	0.70
1:AA:2316:C:H1'	6:AG:128:ARG:NH2	2.07	0.70
9:AM:15:LEU:HB2	9:AM:134:ARG:CG	2.21	0.70
1:AA:2849:U:OP1	15:AR:95:ARG:NH1	2.25	0.70
19:AT:3:THR:HA	19:AT:6:ASP:OD2	1.91	0.70
20:AU:28:LYS:HA	20:AU:28:LYS:CE	2.20	0.70
20:AU:28:LYS:HA	20:AU:28:LYS:NZ	2.05	0.70
31:BA:687:A:H4'	31:BA:688:G:O5'	1.90	0.70
31:BA:800:G:H8	31:BA:800:G:O5'	1.75	0.70
34:BG:149:ALA:O	34:BG:153:ARG:HG2	1.92	0.70
54:CA:48:C:H4'	54:CA:49:U:OP2	1.91	0.70
32:CE:178:ARG:HD2	38:CK:71:GLY:O	1.92	0.70
42:CO:39:VAL:HB	42:CO:57:LYS:HB2	1.74	0.70
44:CQ:24:CYS:HB2	44:CQ:40:CYS:N	2.06	0.70
45:CR:8:LYS:O	45:CR:12:ILE:HG13	1.92	0.70
55:DA:607:U:O4	55:DA:608:A:C5	2.44	0.70
8:DK:14:ASP:OD1	8:DK:15:VAL:HG22	1.92	0.70
11:DO:115:LEU:HB2	11:DO:131:SER:HB2	1.74	0.70
57:DY:26:LEU:O	57:DY:111:LEU:CB	2.35	0.70
17:A2:71:LEU:N	17:A2:86:GLY:CA	2.49	0.70
1:AA:2030:A:H4'	1:AA:2031:A:H8	1.56	0.70
1:AA:589:C:H2'	1:AA:590:A:C8	2.26	0.70
1:AA:888:C:O2'	1:AA:889:C:OP2	2.08	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:AF:122:LYS:O	5:AF:123:LEU:HB2	1.92	0.70
9:AM:16:ILE:CD1	9:AM:137:LYS:HB2	2.21	0.70
12:AP:31:ASP:N	12:AP:107:ALA:HB2	2.07	0.70
14:AQ:106:ARG:HA	14:AQ:110:LEU:CD1	2.22	0.70
31:BA:275:G:H5'	47:BT:14:LYS:HD2	1.72	0.70
31:BA:971:G:H5''	31:BA:972:C:H5''	1.73	0.70
34:BG:21:LEU:N	34:BG:21:LEU:HD12	2.07	0.70
39:BL:49:PRO:O	39:BL:53:VAL:HG22	1.92	0.70
26:A4:34:GLU:HB3	43:BP:57:ARG:HH12	1.57	0.70
49:BV:41:VAL:HG23	49:BV:44:MET:CE	2.22	0.70
31:BA:1453:G:O6	50:BW:51:GLU:HB2	1.91	0.70
33:CF:91:LEU:HD11	33:CF:101:LEU:HD12	1.73	0.70
47:CT:59:ILE:CG2	47:CT:71:PHE:HB3	2.22	0.70
28:D6:17:LYS:C	28:D6:19:ARG:H	1.91	0.70
55:DA:1204:A:H2	55:DA:1241:A:N1	1.90	0.70
55:DA:469:G:O6	29:D7:37:LYS:HE2	1.92	0.70
55:DA:654(J):A:O2'	55:DA:654(K):C:O5'	2.08	0.70
6:DG:130:ASN:OD1	6:DG:160:VAL:HA	1.92	0.70
7:DH:8:PRO:HG2	7:DH:69:ARG:NE	2.07	0.70
56:DI:23:LEU:HD12	56:DI:23:LEU:N	2.07	0.70
58:DL:19:PRO:HA	58:DL:25:PRO:HG3	1.72	0.70
58:DL:3:LYS:O	58:DL:4:VAL:CG2	2.35	0.70
12:DP:1:MET:C	12:DP:2:LEU:HD22	2.12	0.70
23:DZ:3:LYS:HD3	23:DZ:43:TYR:HD2	1.55	0.70
17:A2:5:VAL:CG2	17:A2:37:VAL:HG11	2.18	0.69
6:AG:128:ARG:HG3	6:AG:128:ARG:HH21	1.57	0.69
15:AR:27:THR:O	15:AR:89:VAL:HG22	1.92	0.69
20:AU:39:VAL:HG23	20:AU:41:GLY:H	1.55	0.69
21:AV:145:GLU:O	21:AV:145:GLU:CD	2.30	0.69
31:BA:1005:A:H5''	31:BA:1006:C:C5	2.27	0.69
31:BA:1060:C:H5''	40:BM:51:ARG:HG2	1.73	0.69
33:BF:18:TRP:NE1	44:BQ:54:PRO:HA	2.05	0.69
54:CA:792:A:H4'	54:CA:793:U:O5'	1.92	0.69
34:CG:105:VAL:HG13	34:CG:110:PHE:HB2	1.73	0.69
35:CH:42:GLY:CA	35:CH:66:MET:HG2	2.22	0.69
6:DG:112:PRO:CA	26:D4:37:SER:HB2	2.22	0.69
55:DA:2064:C:H2'	55:DA:2065:C:C6	2.26	0.69
55:DA:1786:A:C2	55:DA:2606:C:H1'	2.26	0.69
55:DA:273(F):C:H3'	55:DA:274:G:H5''	1.72	0.69
7:DH:168:PRO:O	7:DH:169:VAL:HG12	1.91	0.69
7:DH:30:LYS:HE3	7:DH:81:GLU:HA	1.73	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:DL:108:ALA:C	58:DL:111:LYS:CD	2.57	0.69
9:DM:45:ASN:HD22	9:DM:45:ASN:H	1.38	0.69
9:DM:7:LYS:O	9:DM:9:VAL:HG13	1.92	0.69
20:DU:86:ARG:HB2	20:DU:95:LYS:HD2	1.72	0.69
2:AB:40:U:C4	26:A4:1:MET:SD	2.85	0.69
1:AA:140:A:C8	1:AA:1408:C:O2'	2.45	0.69
11:AO:85:LEU:HA	11:AO:88:LEU:CB	2.22	0.69
31:BA:736:C:H2'	31:BA:737:A:H8	1.55	0.69
31:BA:792:A:H4'	31:BA:793:U:O5'	1.93	0.69
31:BA:954:G:H21	31:BA:1227:A:N6	1.89	0.69
33:BF:58:GLU:O	33:BF:64:VAL:HA	1.91	0.69
37:BJ:20:ASP:OD2	37:BJ:23:VAL:HG23	1.92	0.69
54:CA:1028(B):C:H3'	54:CA:1029:G:H5''	1.73	0.69
54:CA:418:C:H2'	54:CA:419:C:H6	1.57	0.69
54:CA:64:G:H4'	54:CA:65:U:H5'	1.73	0.69
54:CA:923:A:OP1	35:CH:21:ALA:HB2	1.92	0.69
52:CB:2:C:H2'	52:CB:3:C:C6	2.27	0.69
40:CM:38:ILE:HG12	40:CM:71:LEU:O	1.92	0.69
26:D4:38:LYS:C	26:D4:40:HIS:H	1.95	0.69
55:DA:2728:U:O2'	55:DA:2729:G:H5'	1.92	0.69
55:DA:974:G:O2'	55:DA:975:G:N7	2.24	0.69
56:DI:23:LEU:HD12	56:DI:23:LEU:H	1.57	0.69
11:DO:38:GLN:O	11:DO:41:ARG:N	2.23	0.69
20:DU:81:LYS:HZ3	20:DU:98:VAL:HG11	1.57	0.69
21:DV:48:PHE:HE2	21:DV:71:VAL:HG11	1.56	0.69
24:DW:40:SER:C	24:DW:42:GLY:H	1.94	0.69
57:DY:122:VAL:HA	57:DY:126:ALA:HB2	1.72	0.69
57:DY:129:PRO:HD2	57:DY:131:MET:N	2.07	0.69
57:DY:89:ALA:HB3	56:DJ:15:ALA:HB2	1.66	0.69
1:AA:1168:G:H2'	1:AA:1169:G:C8	2.28	0.69
1:AA:1657:C:H4'	4:AE:133:LYS:HB3	1.74	0.69
1:AA:2473:U:O2	1:AA:2473:U:H2'	1.92	0.69
1:AA:871:U:O2	1:AA:871:U:H2'	1.91	0.69
3:AD:35:LYS:HZ1	3:AD:65:ILE:HA	1.55	0.69
5:AF:22:ALA:C	5:AF:24:LEU:N	2.44	0.69
9:AM:67:LEU:O	9:AM:88:GLU:HB2	1.93	0.69
24:AW:51:ARG:HE	24:AW:55:ARG:HH12	1.38	0.69
31:BA:1095:U:P	31:BA:1108:G:H1	2.15	0.69
39:BL:17:VAL:HG11	39:BL:81:ILE:HD13	1.73	0.69
45:BR:82:ILE:HD11	45:BR:87:ILE:O	1.92	0.69
37:CJ:12:LEU:HD22	37:CJ:12:LEU:H	1.57	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:CM:40:LEU:HD12	40:CM:69:ASN:HB3	1.75	0.69
55:DA:1251:C:O2'	55:DA:1252:G:H3'	1.91	0.69
55:DA:1762:A:H5''	55:DA:1763:G:OP2	1.91	0.69
55:DA:1784:A:H4'	55:DA:1785:A:C5'	2.21	0.69
55:DA:205:G:O2'	55:DA:206:U:P	2.49	0.69
55:DA:468:G:N7	29:D7:39:ARG:NH2	2.34	0.69
55:DA:752:A:O2'	55:DA:753:C:OP2	2.10	0.69
3:DD:166:GLN:HE21	3:DD:166:GLN:CA	2.04	0.69
5:DF:136:THR:O	5:DF:140:LEU:HB2	1.92	0.69
5:DF:198:ALA:CA	5:DF:201:VAL:HG12	2.23	0.69
8:DK:76:THR:HG23	8:DK:139:GLN:HE22	1.56	0.69
58:DL:15:GLY:O	58:DL:16:LYS:HB2	1.90	0.69
58:DL:34:ILE:O	58:DL:34:ILE:HG23	1.92	0.69
58:DL:95:LYS:HB3	58:DL:136:VAL:CG2	2.20	0.69
17:A2:6:LYS:H	17:A2:37:VAL:CG1	2.06	0.69
1:AA:612:G:H5'	1:AA:612:G:C8	2.25	0.69
21:AV:106:GLY:C	21:AV:108:PRO:CD	2.49	0.69
24:AW:17:SER:HB2	24:AW:18:PRO:CA	2.21	0.69
31:BA:1306:A:N6	31:BA:1331:G:H1'	2.07	0.69
52:BD:21:A:C2'	52:BD:22:G:H5''	2.22	0.69
39:BL:28:VAL:HG13	39:BL:64:THR:HA	1.73	0.69
54:CA:1312:G:N7	49:CV:4:SER:OG	2.22	0.69
54:CA:1449:C:C3'	54:CA:1450:U:H5''	2.22	0.69
52:CD:2:C:H2'	52:CD:3:C:C6	2.27	0.69
43:CP:4:ILE:HG22	43:CP:5:ALA:N	2.08	0.69
55:DA:1086:A:C2	57:DY:41:ARG:NH2	2.60	0.69
55:DA:2298:A:N6	55:DA:2318:G:H8	1.90	0.69
55:DA:880:G:H2'	55:DA:880:G:N3	2.06	0.69
55:DA:889:C:H2'	55:DA:889:C:O2	1.92	0.69
3:DD:236:GLY:C	3:DD:237:GLU:OE2	2.30	0.69
3:DD:44:ASN:HD22	3:DD:44:ASN:C	1.92	0.69
6:DG:7:LEU:O	6:DG:7:LEU:HD23	1.92	0.69
55:DA:1113:U:OP1	7:DH:2:SER:N	2.25	0.69
56:DI:10:GLU:O	56:DI:14:GLN:HB3	1.92	0.69
56:DI:19:GLU:O	56:DI:20:LEU:C	2.29	0.69
57:DY:52:PHE:HD2	57:DY:52:PHE:N	1.91	0.69
23:DZ:41:ARG:HB2	23:DZ:43:TYR:CE1	2.28	0.69
16:A1:27:LEU:HB3	16:A1:31:SER:HB3	1.75	0.69
1:AA:1164:G:H2'	1:AA:1165:U:H6	1.56	0.69
1:AA:1803:A:H2	1:AA:1822:G:N3	1.89	0.69
1:AA:2051:A:N6	1:AA:2614:A:H2'	2.08	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AB:8:U:H6	2:AB:8:U:H5'	1.56	0.69
4:AE:175:VAL:O	4:AE:177:PRO:HD3	1.92	0.69
6:AG:101:ILE:HG13	6:AG:102:PHE:N	2.07	0.69
15:AR:90:GLN:HA	15:AR:90:GLN:HE21	1.57	0.69
24:AW:17:SER:CB	24:AW:18:PRO:CA	2.71	0.69
31:BA:1337:G:H5''	31:BA:1338:G:OP1	1.92	0.69
31:BA:737:A:H2'	31:BA:738:C:H6	1.57	0.69
31:BA:982:U:H4'	31:BA:983:A:O5'	1.93	0.69
52:BD:15:G:N1	52:BD:48:C:N4	2.40	0.69
52:BD:35:A:H61	52:BD:37:MIA:H153	1.57	0.69
26:A4:63:TYR:CE2	49:BV:41:VAL:HA	2.28	0.69
54:CA:1391:U:H2'	54:CA:1392:G:C8	2.28	0.69
54:CA:713:G:H21	54:CA:777:A:H1'	1.58	0.69
26:D4:32:TYR:C	26:D4:32:TYR:HD2	1.95	0.69
55:DA:1543:A:O2'	55:DA:1544:C:O5'	2.09	0.69
55:DA:2308:G:H5'	55:DA:2309:A:OP2	1.92	0.69
55:DA:386:G:H3'	55:DA:388:G:N2	2.08	0.69
55:DA:528:A:H2	55:DA:2043:C:C5'	2.05	0.69
8:DK:2:LYS:HA	8:DK:20:ASP:HA	1.72	0.69
58:DL:141:ALA:HB1	58:DL:142:PRO:C	2.13	0.69
57:DY:26:LEU:O	57:DY:111:LEU:CG	2.40	0.69
57:DY:92:THR:O	57:DY:96:PHE:O	2.09	0.69
1:AA:107:C:H2'	1:AA:108:U:H6	1.58	0.69
1:AA:1314:C:H5'	1:AA:1314:C:C6	2.27	0.69
1:AA:2282:G:O2'	1:AA:2283:C:OP2	2.10	0.69
1:AA:2864:G:OP1	15:AR:119:LYS:HD2	1.92	0.69
6:AG:10:LYS:HD3	6:AG:10:LYS:O	1.93	0.69
20:AU:20:TYR:OH	20:AU:42:VAL:HA	1.93	0.69
31:BA:652:U:O4	31:BA:752:G:H2'	1.93	0.69
32:BE:22:LYS:HA	32:BE:22:LYS:HE3	1.75	0.69
32:BE:69:LEU:HD21	32:BE:93:VAL:HG23	1.75	0.69
33:BF:148:GLY:HA3	33:BF:203:PHE:HB3	1.75	0.69
33:BF:83:ARG:O	33:BF:87:LEU:HG	1.93	0.69
38:BK:121:ASP:HB2	38:BK:125:ARG:HH22	1.58	0.69
39:BL:106:ALA:O	39:BL:108:VAL:HG13	1.93	0.69
50:BW:74:LYS:C	50:BW:76:ALA:H	1.94	0.69
54:CA:1032(A):G:H2'	54:CA:1032(B):G:C8	2.28	0.69
52:CB:19:G:O6	55:DA:881:G:H2'	1.93	0.69
32:CE:19:HIS:NE2	32:CE:206:ASP:HB2	2.07	0.69
43:CP:116:THR:O	43:CP:117:VAL:HG12	1.92	0.69
49:CV:5:LEU:HD22	49:CV:10:PHE:HE1	1.57	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:D4:23:GLU:H	26:D4:23:GLU:CD	1.95	0.69
55:DA:1332:G:H22	55:DA:1609:A:H2'	1.56	0.69
55:DA:654(Q):C:H2'	55:DA:654(R):C:C5	2.27	0.69
55:DA:848:G:H2'	55:DA:849:A:C8	2.26	0.69
55:DA:910:A:C5	12:DP:13:GLN:HG3	2.27	0.69
3:DD:17:THR:HG22	3:DD:205:VAL:N	2.08	0.69
4:DE:197:ILE:HD11	4:DE:199:ARG:HH22	1.56	0.69
6:DG:118:ARG:HA	6:DG:118:ARG:HE	1.57	0.69
57:DY:122:VAL:CB	57:DY:126:ALA:HB3	2.23	0.69
57:DY:43:ALA:HB3	57:DY:47:ASN:HD22	0.56	0.69
1:AA:1929:G:H2'	1:AA:1929:G:N3	2.07	0.69
1:AA:323:G:H5''	1:AA:324:A:H5'	1.74	0.69
1:AA:670:A:H4'	1:AA:671:C:OP1	1.92	0.69
3:AD:4:LYS:HZ1	3:AD:20:ASP:HA	1.57	0.69
4:AE:76:ARG:HD3	4:AE:195:LEU:HB2	1.75	0.69
7:AH:97:ARG:HG2	7:AH:98:LEU:H	1.58	0.69
9:AM:16:ILE:HD13	9:AM:137:LYS:HB2	1.75	0.69
21:AV:104:PHE:O	21:AV:105:VAL:HG12	1.91	0.69
31:BA:1374:A:C2'	31:BA:1375:A:H5'	2.23	0.69
1:AA:2602:A:H61	52:BB:76:A:H5'	1.56	0.69
32:BE:235:SER:O	32:BE:239:VAL:HG21	1.91	0.69
34:BG:21:LEU:N	34:BG:21:LEU:CD1	2.53	0.69
38:BK:21:LYS:O	38:BK:63:LEU:HD23	1.92	0.69
54:CA:713:G:H21	54:CA:777:A:C1'	2.05	0.69
54:CA:913:A:O2'	54:CA:914:A:OP2	2.10	0.69
33:CF:34:LEU:HD21	33:CF:38:ARG:HD2	1.73	0.69
16:D1:44:ASN:HD22	16:D1:44:ASN:N	1.89	0.69
12:DP:80:GLU:OE2	22:D3:4:LYS:HE3	1.92	0.69
26:D4:15:ILE:HD13	26:D4:15:ILE:N	2.06	0.69
26:D4:37:SER:HB3	26:D4:42:PHE:CD1	2.28	0.69
55:DA:34:C:O2'	55:DA:35:G:OP2	2.10	0.69
55:DA:443:A:H3'	5:DF:45:ARG:NH1	2.08	0.69
4:DE:57:LYS:CE	4:DE:59:VAL:HB	2.23	0.69
7:DH:46:GLU:OE2	7:DH:51:ARG:HD2	1.91	0.69
56:DJ:17:VAL:O	56:DJ:18:LEU:CB	2.41	0.69
58:DL:62:ASP:O	58:DL:63:ARG:HB2	1.91	0.69
57:DY:118:THR:CG2	57:DY:119:ALA:N	2.37	0.69
57:DY:24:PHE:C	57:DY:24:PHE:CD2	2.64	0.69
57:DY:74:LEU:CD1	57:DY:75:GLN:HG2	2.22	0.69
57:DY:75:GLN:OE1	57:DY:109:SER:OG	2.05	0.69
22:A3:2:ALA:O	22:A3:3:HIS:O	2.11	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1018:C:H2'	1:AA:1019:U:C6	2.28	0.69
1:AA:27:G:O2'	1:AA:28:A:P	2.51	0.69
1:AA:775:G:H4'	1:AA:776:G:O5'	1.92	0.69
1:AA:896:A:H1'	21:AV:176:PRO:HG2	1.74	0.69
2:AB:48:A:H2'	2:AB:49:C:C6	2.27	0.69
20:AU:63:LYS:HA	20:AU:63:LYS:NZ	2.07	0.69
31:BA:1127:G:H4'	31:BA:1148:U:O2	1.92	0.69
32:BE:22:LYS:HA	32:BE:22:LYS:CE	2.23	0.69
45:BR:25:THR:HG21	45:BR:70:LEU:HB2	1.75	0.69
51:BX:15:ARG:NH1	51:BX:15:ARG:HB2	2.08	0.69
54:CA:412:A:O2'	54:CA:413:G:OP2	2.10	0.69
54:CA:56:U:H2'	54:CA:57:G:C8	2.27	0.69
47:CT:74:LEU:HD12	47:CT:75:ARG:HG2	1.75	0.69
55:DA:1212:G:O2'	55:DA:1236:G:N2	2.23	0.69
55:DA:2133:G:H2'	55:DA:2157:G:H22	1.57	0.69
55:DA:2168:G:O4'	55:DA:2168:G:OP1	2.11	0.69
55:DA:1568:G:H5"	3:DD:61:LEU:HD22	1.73	0.69
7:DH:59:ARG:HG3	7:DH:59:ARG:HH11	1.58	0.69
58:DL:19:PRO:C	58:DL:25:PRO:HG2	2.13	0.69
58:DL:83:GLY:N	58:DL:99:ILE:HG23	2.07	0.69
20:DU:39:VAL:HG12	20:DU:40:GLU:N	2.06	0.69
57:DY:5:ARG:O	57:DY:7:VAL:CG1	2.41	0.69
30:A8:14:VAL:HG11	30:A8:22:VAL:CG1	2.22	0.69
1:AA:2866:U:H2'	1:AA:2866:U:O2	1.91	0.69
1:AA:654(O):G:H2'	1:AA:654(P):G:H8	1.58	0.69
7:AH:6:ARG:HB2	7:AH:66:GLY:HA2	1.75	0.69
21:AV:108:PRO:HB3	21:AV:142:SER:O	1.92	0.69
31:BA:1027:C:H2'	31:BA:1028:C:C6	2.28	0.69
31:BA:243:A:N6	31:BA:281:G:O2'	2.26	0.69
31:BA:887:G:C3'	31:BA:888:G:H5'	2.22	0.69
32:BE:47:THR:O	32:BE:51:LEU:HG	1.92	0.69
37:BJ:18:TYR:CD2	37:BJ:59:LEU:HB2	2.27	0.69
43:BP:10:PRO:HB2	43:BP:18:ALA:CB	2.13	0.69
54:CA:1022:G:H2'	54:CA:1023:G:O4'	1.93	0.69
54:CA:869:G:H4'	54:CA:872:A:C1'	2.23	0.69
43:CP:4:ILE:HG22	43:CP:5:ALA:H	1.56	0.69
16:D1:64:ARG:HH21	16:D1:64:ARG:CG	2.04	0.69
22:D3:68:GLU:OE1	22:D3:82:ARG:HB2	1.93	0.69
28:D6:36:LEU:HD12	28:D6:50:ARG:HG2	1.74	0.69
56:DI:25:ASP:HA	56:DI:28:LYS:HG2	1.75	0.69
58:DL:125:ARG:NE	58:DL:132:ARG:HH22	1.90	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:DV:117:LEU:N	21:DV:117:LEU:HD12	2.08	0.69
21:DV:127:LYS:O	21:DV:161:VAL:CB	2.41	0.69
24:DW:65:ASN:ND2	24:DW:69:ARG:HH21	1.90	0.69
25:DX:35:ARG:HH21	25:DX:37:LEU:HD21	1.58	0.69
57:DY:74:LEU:HB3	57:DY:120:LYS:N	2.08	0.69
57:DY:141:VAL:HG22	57:DY:142:LEU:H	1.57	0.69
57:DY:56:ASN:HA	57:DY:60:ARG:HG2	1.72	0.69
1:AA:2061:G:H5''	1:AA:2503:A:N1	2.08	0.69
1:AA:2523:G:H5'	1:AA:2523:G:H8	1.57	0.69
1:AA:270(K):C:HO2'	1:AA:270(L):U:H5	1.40	0.69
1:AA:387:U:H4'	1:AA:388:G:O5'	1.93	0.69
6:AG:131:TYR:HB3	6:AG:159:VAL:CG1	2.23	0.69
8:AK:56:LYS:HD2	8:AK:60:GLU:HB2	1.75	0.69
9:AM:68:GLU:HG2	9:AM:88:GLU:OE1	1.92	0.69
10:AN:71:ARG:HH21	10:AN:77:ILE:HG21	1.55	0.69
11:AO:63:PRO:O	11:AO:64:LYS:HB2	1.93	0.69
1:AA:480:A:H1'	20:AU:44:ILE:HD13	1.74	0.69
21:AV:76:LEU:H	21:AV:76:LEU:HD23	1.56	0.69
24:AW:17:SER:HB2	24:AW:18:PRO:C	2.12	0.69
53:B1:34:G:H2'	53:B1:35:A:C8	2.27	0.69
31:BA:1244:C:H2'	31:BA:1245:A:H8	1.58	0.69
31:BA:954:G:H4'	43:BP:120:LYS:HG2	1.75	0.69
36:BI:12:PRO:HG2	36:BI:13:ASN:H	1.57	0.69
41:BN:24:SER:HB3	41:BN:27:ASN:O	1.92	0.69
42:BO:25:PRO:O	42:BO:27:LEU:HD23	1.93	0.69
54:CA:160:A:H1'	54:CA:344:A:C8	2.27	0.69
33:CF:13:GLY:HA3	44:CQ:57:ARG:HE	1.58	0.69
46:CS:68:ASP:C	46:CS:70:ALA:H	1.96	0.69
55:DA:128:C:C4'	29:D7:49:ARG:HH21	2.05	0.69
55:DA:1064:C:O2'	58:DL:89:HIS:HB2	1.93	0.69
55:DA:1887:C:C3'	55:DA:1888:G:H5''	2.21	0.69
55:DA:1996:C:H4'	55:DA:1997:G:O5'	1.92	0.69
55:DA:221:A:H4'	55:DA:222:A:O5'	1.92	0.69
55:DA:2790:A:O2'	55:DA:2791:C:P	2.51	0.69
55:DA:405:U:O2	55:DA:405:U:H2'	1.93	0.69
55:DA:483:A:O2'	20:DU:59:GLY:HA2	1.92	0.69
55:DA:612:G:H2'	55:DA:613:U:O2	1.93	0.69
55:DA:1826:G:C4'	3:DD:242:ARG:HH21	2.05	0.69
3:DD:94:LEU:HD13	3:DD:94:LEU:C	2.12	0.69
55:DA:443:A:C5	5:DF:45:ARG:HD2	2.28	0.69
6:DG:126:ASP:CG	6:DG:130:ASN:HB2	2.13	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:DK:23:PRO:HB2	8:DK:27:ARG:HH12	1.56	0.69
58:DL:49:GLY:CA	58:DL:50:ASP:HB3	2.23	0.69
58:DL:56:GLU:HB3	58:DL:68:VAL:HG13	1.73	0.69
12:DP:27:VAL:HA	12:DP:105:GLU:OE1	1.92	0.69
21:DV:120:ILE:O	21:DV:171:ILE:HD13	1.93	0.69
25:DX:19:GLN:HE22	25:DX:52:HIS:CE1	2.10	0.69
57:DY:101:PRO:O	57:DY:102:LYS:HB3	1.92	0.69
57:DY:129:PRO:CD	57:DY:130:THR:H	2.06	0.69
57:DY:13:LEU:HD13	57:DY:13:LEU:O	1.93	0.69
57:DY:18:GLU:CG	57:DY:66:LEU:HD13	2.12	0.69
1:AA:2873:A:H8	13:A0:6:SER:N	1.88	0.69
1:AA:897:C:H2'	1:AA:898:C:H5'	1.73	0.69
11:AO:31:ALA:O	11:AO:32:THR:HG23	1.93	0.69
31:BA:977:A:H2'	31:BA:978:A:H5'	1.75	0.69
34:BG:34:GLU:C	34:BG:35:ARG:HG3	2.14	0.69
31:BA:718:G:H5'	41:BN:117:ASN:CG	2.14	0.69
47:BT:8:GLY:HA3	47:BT:23:VAL:HG22	1.74	0.69
50:BW:25:ARG:O	50:BW:29:LYS:HE3	1.93	0.69
31:BA:1286:A:H5''	51:BX:25:LYS:HD2	1.74	0.69
54:CA:703:G:H4'	54:CA:704:A:H5'	1.74	0.69
54:CA:789:U:H3'	54:CA:789:U:O2	1.93	0.69
34:CG:198:VAL:HG12	34:CG:199:ASN:N	2.08	0.69
49:CV:5:LEU:CD1	49:CV:5:LEU:C	2.61	0.69
26:D4:32:TYR:HD2	26:D4:33:VAL:N	1.91	0.69
55:DA:1076:C:H2'	55:DA:1077:A:C5'	2.02	0.69
55:DA:1270:C:H5''	55:DA:1271:G:O5'	1.93	0.69
55:DA:1431:U:H2'	55:DA:1432:C:C6	2.28	0.69
55:DA:1884:A:C2'	55:DA:1885:A:H5''	2.23	0.69
55:DA:2115:G:H21	55:DA:2172:U:H3	1.41	0.69
55:DA:2189:U:C3'	55:DA:2190:G:H5''	2.23	0.69
55:DA:405:U:H5''	55:DA:406:G:OP2	1.93	0.69
55:DA:648:G:O2'	55:DA:649:G:H5'	1.93	0.69
55:DA:774:A:C2	55:DA:787:U:O2'	2.46	0.69
6:DG:8:LYS:O	6:DG:11:TYR:HB3	1.93	0.69
56:DJ:19:GLU:OE1	56:DJ:19:GLU:HA	1.93	0.69
14:DQ:106:ARG:CA	14:DQ:110:LEU:HD11	2.22	0.69
24:DW:41:ILE:HD12	24:DW:41:ILE:C	2.13	0.69
57:DY:25:PHE:CD2	57:DY:82:PHE:CD1	2.80	0.69
57:DY:61:LEU:C	57:DY:63:LEU:H	1.95	0.69
23:DZ:76:ARG:HB2	23:DZ:94:LEU:HD11	1.75	0.69
28:A6:16:CYS:O	28:A6:17:LYS:HB2	1.92	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2393:A:OP1	30:A8:30:ARG:HB2	1.92	0.68
1:AA:1296:G:O2'	1:AA:1297:C:H5'	1.93	0.68
1:AA:2154:G:H2'	1:AA:2155:G:H8	1.56	0.68
1:AA:2753:A:H2'	1:AA:2754:U:C5'	2.14	0.68
1:AA:322:A:H3'	5:AF:169:ASN:HD21	1.57	0.68
1:AA:407:G:H2'	1:AA:408:G:H8	1.56	0.68
1:AA:497:A:H2'	1:AA:498:G:O4'	1.94	0.68
1:AA:583:G:H5''	16:A1:10:ARG:HH12	1.58	0.68
3:AD:267:SER:O	3:AD:269:PHE:N	2.26	0.68
10:AN:121:VAL:O	10:AN:122:LEU:HD23	1.92	0.68
21:AV:115:GLY:HA2	21:AV:177:PRO:HG2	0.74	0.68
25:AX:59:VAL:HG12	25:AX:60:GLU:N	2.07	0.68
31:BA:1007:C:C3'	31:BA:1008:C:H5''	2.24	0.68
31:BA:1410:G:H2'	31:BA:1411:C:C6	2.28	0.68
31:BA:250:A:H1'	31:BA:252:U:C5	2.28	0.68
32:BE:19:HIS:NE2	32:BE:206:ASP:HB2	2.08	0.68
34:BG:152:SER:O	34:BG:155:LEU:HB2	1.93	0.68
38:BK:12:ARG:HH11	38:BK:26:VAL:HA	1.59	0.68
40:BM:71:LEU:HD12	40:BM:72:VAL:H	1.57	0.68
31:BA:562:C:HO2'	42:BO:15:ARG:HB3	1.55	0.68
52:CD:18:G:H1'	52:CD:58:A:C2	2.29	0.68
33:CF:131:ARG:HG3	33:CF:131:ARG:HH11	1.58	0.68
33:CF:173:VAL:O	33:CF:175:LEU:HD12	1.92	0.68
37:CJ:155:ARG:N	37:CJ:155:ARG:HD3	2.07	0.68
38:CK:83:ILE:HG13	38:CK:137:VAL:HG22	1.73	0.68
43:CP:15:VAL:HG23	43:CP:43:THR:O	1.93	0.68
55:DA:2271:G:OP1	22:D3:18:ALA:HB1	1.94	0.68
26:D4:61:ARG:HD2	26:D4:61:ARG:N	2.08	0.68
55:DA:2657:A:C2	55:DA:2665:A:N7	2.61	0.68
55:DA:754:C:H2'	55:DA:755:C:H6	1.56	0.68
55:DA:780:G:H21	55:DA:783:A:H62	1.41	0.68
55:DA:943:U:OP2	11:DO:36:LYS:HG2	1.91	0.68
3:DD:35:LYS:HG2	3:DD:64:ILE:CG2	2.23	0.68
7:DH:150:ALA:C	7:DH:152:ARG:N	2.42	0.68
7:DH:35:VAL:HG13	7:DH:71:LEU:HG	1.75	0.68
8:DK:23:PRO:HB2	8:DK:27:ARG:NH1	2.08	0.68
11:DO:41:ARG:HH21	11:DO:41:ARG:HG3	1.57	0.68
12:DP:35:VAL:CG1	12:DP:130:LYS:HB3	2.22	0.68
18:DS:29:LEU:HD13	18:DS:69:LEU:HD13	1.75	0.68
21:DV:177:PRO:O	21:DV:177:PRO:HG2	1.91	0.68
57:DY:74:LEU:HD13	57:DY:75:GLN:N	2.08	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:7:VAL:O	57:DY:8:GLU:C	2.30	0.68
1:AA:508:G:O2'	1:AA:509:C:OP1	2.11	0.68
6:AG:118:ARG:HE	6:AG:118:ARG:HA	1.57	0.68
7:AH:20:ALA:HB1	7:AH:23:ARG:HE	1.58	0.68
8:AK:80:PRO:HA	8:AK:143:SER:HA	1.74	0.68
9:AM:18:ALA:O	9:AM:21:LYS:HB2	1.94	0.68
12:AP:42:ILE:N	12:AP:42:ILE:HD12	2.06	0.68
31:BA:801:U:H5'	31:BA:801:U:H6	1.57	0.68
42:BO:75:HIS:CD2	42:BO:77:LEU:HB2	2.27	0.68
31:BA:1226:C:O3'	43:BP:111:LYS:HE2	1.93	0.68
44:BQ:6:LEU:HD22	44:BQ:23:ARG:NH2	2.08	0.68
54:CA:630:G:C2'	54:CA:631:G:C5'	2.59	0.68
36:CI:97:PHE:O	48:CU:31:LEU:HD23	1.93	0.68
54:CA:1152:A:H5''	40:CM:13:HIS:CD2	2.28	0.68
45:CR:17:ARG:HG3	45:CR:17:ARG:HH11	1.57	0.68
54:CA:375:U:H4'	46:CS:17:TYR:CE2	2.29	0.68
46:CS:67:THR:O	46:CS:70:ALA:HB3	1.93	0.68
49:CV:80:TYR:CZ	49:CV:82:GLY:O	2.46	0.68
55:DA:2014:A:HO2'	27:D5:2:ALA:N	1.91	0.68
30:D8:44:LYS:N	30:D8:44:LYS:HD2	2.06	0.68
55:DA:2182:G:H2'	55:DA:2183:C:H6	1.57	0.68
55:DA:2458:G:O2'	55:DA:2460:U:O4	2.10	0.68
55:DA:614:U:H5'	55:DA:614:U:H6	1.57	0.68
55:DA:828:U:H3	55:DA:2247:A:H4'	1.58	0.68
55:DA:905:U:C3'	55:DA:906:G:H5''	2.23	0.68
8:DK:1:MET:HG3	8:DK:23:PRO:HB3	1.74	0.68
55:DA:1058:U:O2'	58:DL:115:LEU:HA	1.93	0.68
55:DA:1138:G:N2	9:DM:106:MET:HE3	2.07	0.68
14:DQ:107:GLU:N	14:DQ:110:LEU:HD11	2.08	0.68
1:AA:2122:U:H2'	1:AA:2123:G:O4'	1.93	0.68
1:AA:2712:U:OP1	1:AA:2714:G:H4'	1.93	0.68
1:AA:622:G:O2'	1:AA:623:G:H5'	1.93	0.68
1:AA:844:C:H2'	1:AA:845:G:O4'	1.93	0.68
3:AD:139:GLY:H	3:AD:165:ILE:HB	1.57	0.68
3:AD:43:ARG:HD2	3:AD:44:ASN:ND2	2.09	0.68
5:AF:140:LEU:HD13	5:AF:170:LEU:HD21	1.75	0.68
7:AH:101:ARG:HG3	7:AH:117:PRO:HG3	1.76	0.68
10:AN:2:ILE:HG23	10:AN:6:THR:HB	1.74	0.68
20:AU:97:ARG:HH21	20:AU:98:VAL:CB	2.03	0.68
21:AV:74:VAL:HG22	21:AV:86:VAL:HG13	1.75	0.68
39:BL:16:ARG:HH12	39:BL:64:THR:HB	1.58	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:1286:A:C5'	51:BX:25:LYS:HD2	2.22	0.68
54:CA:39:G:N7	54:CA:547:A:H8	1.91	0.68
34:CG:108:LEU:HB3	34:CG:110:PHE:CE1	2.28	0.68
38:CK:41:ARG:HH11	38:CK:41:ARG:CB	2.04	0.68
42:CO:83:VAL:HG21	42:CO:100:ILE:HG12	1.73	0.68
17:D2:25:LEU:HD12	17:D2:94:LEU:HD21	1.74	0.68
12:DP:2:LEU:HB3	12:DP:70:PRO:CG	2.24	0.68
57:DY:63:LEU:CD2	57:DY:65:GLU:OE1	2.36	0.68
30:A8:32:LEU:HG	30:A8:36:LYS:HG3	1.76	0.68
1:AA:1324:G:H1'	1:AA:1616:A:N6	2.09	0.68
1:AA:1444:G:H2'	1:AA:1445:C:C5	2.29	0.68
1:AA:1534:G:N2	1:AA:1538:G:C6	2.59	0.68
1:AA:1581:G:H2'	1:AA:1582:C:O4'	1.93	0.68
1:AA:1794:U:H2'	1:AA:1795:C:H6	1.58	0.68
1:AA:1948:G:H8	1:AA:1948:G:H5'	1.59	0.68
1:AA:2427:C:H5''	1:AA:2428:G:OP1	1.93	0.68
7:AH:35:VAL:CG1	7:AH:71:LEU:HG	2.23	0.68
12:AP:82:ARG:HG2	12:AP:82:ARG:NH1	2.02	0.68
21:AV:105:VAL:CG2	21:AV:106:GLY:N	2.55	0.68
21:AV:16:SER:O	21:AV:20:ARG:HG3	1.91	0.68
31:BA:1132:C:H2'	31:BA:1133:G:C8	2.29	0.68
31:BA:1262:C:H2'	31:BA:1263:C:C6	2.28	0.68
31:BA:1352:C:H2'	31:BA:1353:G:C8	2.29	0.68
31:BA:1365:G:O2'	31:BA:1366:C:H5'	1.92	0.68
31:BA:188:U:O2'	31:BA:189:U:H5'	1.93	0.68
52:BC:23:A:H2'	52:BC:24:G:H8	1.59	0.68
33:BF:54:ARG:HG2	33:BF:54:ARG:HH11	1.59	0.68
54:CA:1176:A:H2'	54:CA:1177:G:H5'	1.76	0.68
54:CA:47:C:C6	54:CA:365:U:H2'	2.29	0.68
32:CE:5:ILE:HG21	32:CE:221:LEU:HA	1.74	0.68
39:CL:106:ALA:O	39:CL:108:VAL:HG13	1.93	0.68
43:CP:116:THR:CG2	43:CP:117:VAL:N	2.56	0.68
55:DA:2331:G:H4'	22:D3:43:THR:H	1.58	0.68
28:D6:19:ARG:HE	28:D6:21:TYR:HE2	1.41	0.68
55:DA:2173:A:C5	55:DA:2174:C:H1'	2.27	0.68
55:DA:2273:A:O2'	55:DA:2274:A:H5'	1.93	0.68
55:DA:2636:U:P	4:DE:79:ARG:HA	2.32	0.68
55:DA:704:G:C2'	55:DA:726:G:H22	2.05	0.68
3:DD:17:THR:HG22	3:DD:205:VAL:H	1.58	0.68
55:DA:2086:U:OP1	3:DD:262:ARG:HD3	1.92	0.68
4:DE:24:THR:CG2	4:DE:188:VAL:HG11	2.19	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:DE:95:ILE:N	4:DE:95:ILE:HD12	2.07	0.68
56:DI:7:ARG:HE	56:DI:8:ILE:CD1	2.07	0.68
56:DJ:17:VAL:HG12	56:DJ:17:VAL:O	1.93	0.68
56:DJ:4:ASP:HA	56:DJ:7:ARG:HB3	1.76	0.68
9:DM:114:ARG:O	9:DM:116:LEU:N	2.25	0.68
21:DV:194:PRO:O	21:DV:196:VAL:HG12	1.92	0.68
57:DY:36:GLU:O	57:DY:37:THR:C	2.32	0.68
57:DY:43:ALA:CA	57:DY:47:ASN:HD22	2.07	0.68
57:DY:25:PHE:CA	57:DY:82:PHE:CZ	2.77	0.68
26:A4:9:LEU:CD2	26:A4:25:TYR:HB3	2.24	0.68
29:A7:30:VAL:HA	29:A7:33:ARG:HH12	1.58	0.68
1:AA:1209:G:N2	1:AA:1210:A:H62	1.92	0.68
1:AA:1406:U:H3'	1:AA:1407:C:H6	1.58	0.68
1:AA:1534:G:H3'	1:AA:1535:U:C5'	2.22	0.68
1:AA:2557:G:O2'	1:AA:2558:C:H5'	1.93	0.68
7:AH:10:PRO:HG2	7:AH:50:VAL:HG13	1.75	0.68
10:AN:2:ILE:HD11	10:AN:82:ASN:HD22	1.59	0.68
20:AU:72:VAL:HG23	20:AU:73:ARG:N	2.09	0.68
31:BA:186(D):C:H2'	31:BA:186(E):C:C6	2.28	0.68
31:BA:274:A:O2'	31:BA:275:G:O4'	2.10	0.68
31:BA:48:C:H5'	31:BA:49:U:OP2	1.93	0.68
31:BA:676:A:H2'	31:BA:677:U:C6	2.28	0.68
31:BA:991:U:H3	31:BA:1213:A:N6	1.91	0.68
36:BI:69:GLU:O	36:BI:72:VAL:HG12	1.94	0.68
54:CA:498:A:O2'	54:CA:500:G:C8	2.44	0.68
54:CA:818:G:H3'	54:CA:819:A:C5'	2.23	0.68
54:CA:957:U:H1'	54:CA:960:U:C5	2.28	0.68
48:CU:29:PHE:N	48:CU:29:PHE:CD2	2.60	0.68
48:CU:66:LEU:O	48:CU:70:ILE:HG13	1.92	0.68
50:CW:49:ALA:HB1	50:CW:99:LEU:HB2	1.76	0.68
27:D5:48:GLU:HG3	27:D5:59:GLU:HB2	1.75	0.68
55:DA:212:G:O2'	55:DA:213:A:H5'	1.93	0.68
55:DA:483:A:H3'	55:DA:484:C:H6	1.59	0.68
5:DF:59:TYR:CD1	5:DF:78:ILE:HB	2.28	0.68
58:DL:141:ALA:CB	58:DL:142:PRO:CA	2.71	0.68
58:DL:34:ILE:CD1	58:DL:38:VAL:HG22	2.23	0.68
18:DS:9:TYR:H	18:DS:102:HIS:CD2	2.12	0.68
21:DV:178:GLU:OE1	21:DV:180:VAL:C	2.32	0.68
57:DY:73:GLY:HA3	57:DY:112:LEU:CG	2.23	0.68
57:DY:51:LEU:CD2	57:DY:82:PHE:N	2.57	0.68
23:DZ:3:LYS:HD3	23:DZ:43:TYR:CD2	2.28	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2347:C:H4'	28:A6:39:TYR:CE2	2.29	0.68
1:AA:1171:G:OP2	1:AA:1171:G:H3'	1.93	0.68
1:AA:1458:C:H5''	1:AA:1459:G:H5'	1.75	0.68
1:AA:1558:A:H4'	1:AA:1559:G:O5'	1.94	0.68
1:AA:2552:U:H3'	1:AA:2554:U:OP2	1.93	0.68
1:AA:627:A:H4'	1:AA:628:G:OP1	1.92	0.68
1:AA:90:U:H2'	1:AA:91:A:H5''	1.74	0.68
3:AD:77:ALA:HB2	3:AD:97:TYR:CD2	2.28	0.68
4:AE:9:VAL:CG2	4:AE:10:GLY:H	2.06	0.68
6:AG:127:GLY:O	6:AG:128:ARG:HG2	1.94	0.68
21:AV:53:ILE:HG22	21:AV:71:VAL:O	1.93	0.68
31:BA:1218:C:H2'	31:BA:1219:U:C6	2.28	0.68
31:BA:1347:G:H21	31:BA:1373:G:H2'	1.59	0.68
31:BA:703:G:O2'	31:BA:704:A:H8	1.77	0.68
34:BG:108:LEU:HB3	34:BG:110:PHE:HE1	1.59	0.68
31:BA:1148:U:O2'	39:BL:14:VAL:HG11	1.93	0.68
50:BW:26:ASN:HD22	50:BW:27:LYS:N	1.91	0.68
54:CA:274:A:O2'	54:CA:275:G:H8	1.77	0.68
52:CC:7:A:H4'	52:CC:8:U:OP2	1.93	0.68
41:CN:21:ILE:HG13	41:CN:30:VAL:HG12	1.76	0.68
43:CP:84:ILE:HD11	49:CV:66:MET:HB3	1.74	0.68
46:CS:20:VAL:HG21	46:CS:32:TYR:CD1	2.28	0.68
55:DA:128:C:H4'	29:D7:49:ARG:HH21	1.57	0.68
55:DA:1075:C:H4'	21:DV:195:GLU:CD	2.14	0.68
55:DA:1203:G:H3'	55:DA:1204:A:H5''	1.74	0.68
55:DA:1291:C:C5'	55:DA:1536:A:H5'	2.23	0.68
55:DA:229:A:O2'	55:DA:230:U:OP2	2.12	0.68
55:DA:2620:C:OP1	4:DE:152:LYS:O	2.11	0.68
55:DA:483:A:H5'	20:DU:49:VAL:HG22	1.75	0.68
6:DG:101:ILE:HG13	6:DG:102:PHE:N	2.08	0.68
6:DG:88:ILE:HG23	6:DG:88:ILE:O	1.92	0.68
7:DH:152:ARG:HG3	7:DH:153:LYS:HE2	1.74	0.68
7:DH:20:ALA:CB	7:DH:21:PRO:CD	2.71	0.68
58:DL:119:ASP:O	58:DL:122:ALA:CB	2.40	0.68
20:DU:97:ARG:N	20:DU:97:ARG:HD3	2.08	0.68
21:DV:179:ASP:O	21:DV:180:VAL:HB	1.93	0.68
57:DY:112:LEU:CD1	57:DY:121:ASP:CB	2.69	0.68
57:DY:40:LEU:HD13	57:DY:41:ARG:HB2	1.76	0.68
57:DY:52:PHE:CD2	57:DY:52:PHE:N	2.61	0.68
57:DY:51:LEU:HD23	57:DY:82:PHE:O	1.88	0.68
1:AA:1151:G:H5''	16:A1:81:HIS:CE1	2.29	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2061:G:H5''	1:AA:2503:A:C2	2.28	0.68
1:AA:654(Q):C:H2'	1:AA:654(R):C:C5	2.29	0.68
4:AE:50:GLY:HA2	4:AE:78:LEU:HB3	1.74	0.68
5:AF:3:GLU:HA	5:AF:24:LEU:CG	2.22	0.68
7:AH:103:LEU:HD22	7:AH:123:PHE:CE1	2.29	0.68
9:AM:14:VAL:HG11	9:AM:137:LYS:HG3	1.75	0.68
12:AP:38:GLU:HB2	12:AP:127:ILE:CG2	2.23	0.68
31:BA:1032:A:H3'	31:BA:1032(A):G:H5''	1.75	0.68
31:BA:625:G:H2'	31:BA:626:U:C6	2.28	0.68
33:BF:63:ASN:N	33:BF:97:LYS:HD2	2.08	0.68
34:BG:21:LEU:HD12	34:BG:21:LEU:H	1.58	0.68
42:BO:23:LYS:HD3	42:BO:23:LYS:N	2.04	0.68
49:BV:49:ILE:CD1	49:BV:49:ILE:H	2.07	0.68
50:BW:50:GLU:HB2	50:BW:100:ILE:HG12	1.74	0.68
50:BW:33:ILE:CD1	50:BW:62:LEU:HB3	2.23	0.68
54:CA:160:A:H1'	54:CA:344:A:N7	2.09	0.68
54:CA:721:G:H4'	54:CA:722:A:O5'	1.93	0.68
34:CG:110:PHE:HD1	34:CG:110:PHE:H	1.42	0.68
37:CJ:78:ARG:HH11	37:CJ:79:ARG:H	1.41	0.68
42:CO:89:ARG:HE	42:CO:91:LYS:NZ	1.91	0.68
49:CV:53:ASN:HD21	49:CV:56:GLN:HG2	1.59	0.68
49:CV:80:TYR:CE1	49:CV:82:GLY:O	2.46	0.68
55:DA:345:A:H4'	55:DA:346:A:OP1	1.93	0.68
2:DB:81:G:N7	2:DB:96:G:C2	2.62	0.68
55:DA:1798:U:C5'	3:DD:259:THR:HG22	2.18	0.68
4:DE:35:GLN:HE21	4:DE:37:ARG:NE	1.92	0.68
9:DM:137:LYS:CG	9:DM:138:LEU:H	2.06	0.68
9:DM:43:THR:HB	9:DM:46:VAL:CG1	2.24	0.68
57:DY:142:LEU:CD2	57:DY:143:GLN:N	2.56	0.68
57:DY:97:ALA:O	57:DY:98:LYS:C	2.32	0.68
6:AG:67:LYS:HG3	26:A4:6:HIS:CB	2.24	0.68
30:A8:50:LEU:HD12	30:A8:53:PRO:C	2.14	0.68
1:AA:1204:A:H1'	1:AA:1206:G:C5	2.29	0.68
1:AA:140:A:H8	1:AA:1408:C:O2'	1.74	0.68
1:AA:782:A:O2'	3:AD:225:ALA:O	2.11	0.68
5:AF:132:VAL:HG13	5:AF:133:ASN:N	2.09	0.68
5:AF:83:PHE:O	5:AF:84:VAL:HB	1.93	0.68
1:AA:2745:C:H1'	7:AH:143:GLN:HG2	1.75	0.68
8:AK:68:LEU:HA	8:AK:71:ILE:HG22	1.76	0.68
23:AZ:87:PRO:O	23:AZ:91:LYS:N	2.25	0.68
31:BA:1022:G:H2'	31:BA:1023:G:O4'	1.92	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:577:G:H1'	31:BA:816:A:C4	2.28	0.68
31:BA:677:U:H1'	41:BN:119:CYS:SG	2.34	0.68
41:BN:34:ASP:HB3	41:BN:40:ILE:HD11	1.74	0.68
44:BQ:27:CYS:SG	44:BQ:29:ARG:HB2	2.34	0.68
46:BS:52:ASP:OD2	46:BS:54:GLU:HG2	1.94	0.68
54:CA:630:G:H3'	54:CA:630:G:H8	1.50	0.68
54:CA:678:U:H2'	54:CA:679:C:C6	2.29	0.68
54:CA:690:G:H2'	54:CA:691:G:O4'	1.94	0.68
34:CG:111:ALA:HB2	34:CG:120:LEU:HD11	1.76	0.68
35:CH:101:ILE:N	35:CH:101:ILE:HD13	2.09	0.68
38:CK:103:VAL:HG12	38:CK:108:GLY:HA3	1.75	0.68
39:CL:114:TYR:O	39:CL:114:TYR:HD2	1.77	0.68
28:D6:47:THR:HG22	28:D6:48:VAL:N	2.08	0.68
30:D8:34:TRP:CG	30:D8:35:GLN:N	2.59	0.68
55:DA:1510:A:H2	55:DA:1513:C:N4	1.91	0.68
55:DA:701:G:C2'	55:DA:702:G:H5''	2.23	0.68
20:DU:50:ARG:HB3	20:DU:53:PRO:HD2	1.75	0.68
57:DY:112:LEU:H	57:DY:112:LEU:HD23	1.54	0.68
1:AA:1116:C:H2'	1:AA:1117:G:H8	1.59	0.68
1:AA:70:G:H2'	1:AA:113:G:O2'	1.93	0.68
1:AA:387:U:O2'	1:AA:388:G:O5'	2.11	0.68
2:AB:39:A:H2'	26:A4:1:MET:HE2	1.75	0.68
3:AD:155:LEU:HD23	3:AD:177:LEU:HD22	1.76	0.68
3:AD:43:ARG:NH1	3:AD:44:ASN:HD21	1.89	0.68
4:AE:71:GLY:O	4:AE:73:GLU:N	2.27	0.68
14:AQ:66:ALA:O	14:AQ:69:VAL:HG12	1.94	0.68
20:AU:81:LYS:HD3	20:AU:97:ARG:HE	1.58	0.68
25:AX:24:LYS:HE3	25:AX:24:LYS:HA	1.74	0.68
31:BA:971:G:C5	31:BA:1365:G:H5'	2.29	0.68
31:BA:335:C:H2'	31:BA:336:C:H6	1.59	0.68
31:BA:1104:G:H4'	32:BE:111:ARG:NH2	2.08	0.68
43:BP:13:LYS:HA	43:BP:44:ARG:NH1	2.08	0.68
54:CA:382:A:H2'	54:CA:383:A:C8	2.28	0.68
54:CA:630:G:OP1	54:CA:630:G:H4'	1.94	0.68
32:CE:98:LEU:O	32:CE:101:MET:HG3	1.92	0.68
38:CK:129:VAL:HG23	38:CK:130:GLY:H	1.57	0.68
41:CN:22:HIS:HB3	41:CN:29:ILE:HG23	1.74	0.68
17:D2:89:GLN:HA	17:D2:89:GLN:NE2	2.09	0.68
55:DA:1742:C:H5'	55:DA:1743:G:OP2	1.93	0.68
55:DA:1945:G:H2'	55:DA:1946:U:C6	2.27	0.68
55:DA:2030:A:H4'	55:DA:2031:A:C8	2.28	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:796:C:H2'	55:DA:797:C:H6	1.57	0.68
3:DD:35:LYS:CG	3:DD:64:ILE:H	2.07	0.68
55:DA:2810:A:O3'	4:DE:61:ARG:CG	2.42	0.68
5:DF:178:PRO:HB2	5:DF:201:VAL:HG11	1.76	0.68
7:DH:37:VAL:HG12	7:DH:38:SER:N	2.07	0.68
55:DA:1244:G:OP1	11:DO:7:ARG:HD3	1.94	0.68
55:DA:483:A:H5''	20:DU:49:VAL:HG13	1.75	0.68
20:DU:76:CYS:HG	20:DU:77:PRO:HD2	1.55	0.68
57:DY:120:LYS:O	57:DY:121:ASP:CB	2.41	0.68
57:DY:51:LEU:CD1	57:DY:82:PHE:C	2.62	0.68
57:DY:58:LEU:H	57:DY:58:LEU:CD2	1.97	0.68
57:DY:8:GLU:OE2	57:DY:52:PHE:HB3	1.94	0.68
57:DY:91:LYS:HA	57:DY:94:VAL:HB	1.75	0.68
57:DY:8:GLU:O	57:DY:9:LEU:C	2.32	0.68
1:AA:1955:U:O2'	1:AA:1956:U:H5'	1.94	0.68
1:AA:2776:A:H4'	1:AA:2777:G:O5'	1.94	0.68
1:AA:2810:A:O3'	4:AE:61:ARG:HG3	1.92	0.68
1:AA:370:G:H4'	1:AA:371:A:OP2	1.94	0.68
2:AB:44:G:H5''	2:AB:45:A:OP1	1.94	0.68
4:AE:36:ARG:NH2	4:AE:88:GLY:CA	2.57	0.68
11:AO:88:LEU:HD11	11:AO:95:VAL:HG21	1.76	0.68
15:AR:90:GLN:HA	15:AR:90:GLN:NE2	2.08	0.68
21:AV:131:ARG:CG	21:AV:131:ARG:HH11	1.87	0.68
21:AV:163:LEU:HD23	21:AV:163:LEU:N	1.99	0.68
21:AV:145:GLU:OE1	21:AV:174:VAL:HG12	1.93	0.68
31:BA:121:C:H5'	31:BA:122:G:OP1	1.94	0.68
31:BA:946:A:H2'	31:BA:947:G:C8	2.29	0.68
31:BA:668:G:O4'	45:BR:49:ASP:HB2	1.93	0.68
54:CA:430:A:H2'	54:CA:431:A:H5'	1.76	0.68
54:CA:794:A:C2	54:CA:795:C:N3	2.62	0.68
38:CK:87:SER:HA	38:CK:93:VAL:HG23	1.76	0.68
47:CT:4:LYS:HE3	47:CT:6:LEU:CD2	2.24	0.68
13:D0:26:LYS:HE2	13:D0:70:LEU:O	1.94	0.68
55:DA:1057:A:C5	55:DA:1086:A:N3	2.62	0.68
55:DA:1645:G:H5''	55:DA:1646:C:H5'	1.75	0.68
55:DA:914:C:H2'	55:DA:915:C:H5'	1.75	0.68
7:DH:109:PHE:HZ	7:DH:152:ARG:HG2	1.58	0.68
7:DH:98:LEU:HB2	7:DH:125:VAL:CG1	2.24	0.68
56:DJ:1:MET:SD	56:DJ:2:ALA:N	2.63	0.68
9:DM:58:ASP:HB3	9:DM:95:PRO:HB3	1.75	0.68
9:DM:96:GLU:HG2	9:DM:97:ARG:N	2.08	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:DO:65:ARG:NH1	11:DO:65:ARG:HB2	2.09	0.68
57:DY:93:LEU:CD1	57:DY:97:ALA:O	2.42	0.68
1:AA:1267:U:C5	1:AA:2012:G:N2	2.62	0.67
1:AA:2150:U:H2'	1:AA:2151:G:C8	2.28	0.67
1:AA:2468:G:H22	1:AA:2481:G:H2'	1.59	0.67
1:AA:34:C:O2'	1:AA:35:G:H8	1.77	0.67
1:AA:404:C:O2'	1:AA:405:U:H5'	1.94	0.67
2:AB:81:G:N7	2:AB:96:G:N2	2.42	0.67
4:AE:105:THR:HG21	4:AE:164:ARG:NH1	2.09	0.67
4:AE:16:ARG:HD2	4:AE:16:ARG:O	1.93	0.67
10:AN:49:ARG:NH1	31:BA:1422:G:H4'	2.08	0.67
12:AP:130:LYS:NZ	21:AV:81:ARG:HG2	2.08	0.67
14:AQ:106:ARG:HA	14:AQ:110:LEU:HG	1.75	0.67
18:AS:59:VAL:HG23	18:AS:64:MET:H	1.59	0.67
24:AW:51:ARG:NE	24:AW:55:ARG:HH12	1.92	0.67
23:AZ:80:LEU:C	23:AZ:81:LYS:HZ3	1.98	0.67
31:BA:1322:C:HO2'	31:BA:1323:G:H5'	1.58	0.67
31:BA:89:U:H2'	31:BA:90:C:H6	1.59	0.67
52:BB:9:A:O2'	52:BB:10:G:OP1	2.12	0.67
52:BD:64:A:C2	52:BD:65:G:H1'	2.29	0.67
35:BH:53:LEU:O	35:BH:57:LYS:HG3	1.93	0.67
49:BV:46:GLY:N	49:BV:62:ILE:HG23	2.08	0.67
54:CA:1316:G:N2	54:CA:1318:A:H3'	2.09	0.67
54:CA:711:G:O2'	54:CA:712:A:H5'	1.94	0.67
33:CF:47:LEU:HD21	33:CF:68:VAL:HG11	1.76	0.67
39:CL:5:TYR:HA	39:CL:17:VAL:O	1.93	0.67
41:CN:124:LYS:HD2	41:CN:125:PHE:CE1	2.29	0.67
36:CI:97:PHE:CD2	48:CU:31:LEU:HD21	2.29	0.67
16:D1:28:ARG:HD3	16:D1:38:THR:OG1	1.94	0.67
28:D6:19:ARG:NE	28:D6:21:TYR:HE2	1.92	0.67
30:D8:17:THR:HG21	30:D8:21:LYS:HB2	1.76	0.67
30:D8:43:GLN:O	30:D8:44:LYS:HD2	1.93	0.67
30:D8:63:PRO:O	30:D8:64:TYR:HB2	1.93	0.67
55:DA:2345:G:O2'	55:DA:2381:C:H2'	1.93	0.67
55:DA:2392:A:C8	11:DO:60:MET:HG3	2.29	0.67
55:DA:5:A:P	55:DA:5:A:O4'	2.51	0.67
6:DG:26:GLN:NE2	6:DG:27:ASN:HB2	2.09	0.67
55:DA:1093:G:H4'	7:DH:170:ARG:NH2	2.08	0.67
55:DA:1111:A:H5'	7:DH:3:ARG:NH1	2.09	0.67
56:DI:30:ALA:CA	56:DJ:3:LEU:CD2	2.69	0.67
11:DO:126:VAL:HG12	11:DO:147:LEU:HD22	1.76	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:DO:147:LEU:O	11:DO:148:LEU:HB2	1.93	0.67
14:DQ:61:ASN:O	14:DQ:65:VAL:HG23	1.92	0.67
22:A3:14:ARG:O	22:A3:15:ASP:HB2	1.93	0.67
1:AA:1171:G:H4'	1:AA:1173:G:OP1	1.94	0.67
1:AA:1638:C:H4'	1:AA:2710:C:O2	1.93	0.67
3:AD:25:THR:HG21	3:AD:81:ALA:HB1	1.76	0.67
18:AS:39:THR:HG22	18:AS:44:ALA:HB2	1.74	0.67
1:AA:1341:U:O4	19:AT:16:LYS:HE2	1.94	0.67
21:AV:44:PHE:CZ	21:AV:86:VAL:HG21	2.29	0.67
25:AX:54:VAL:HG12	25:AX:55:ARG:N	2.08	0.67
53:B1:43:U:H5'	53:B1:43:U:H6	1.59	0.67
31:BA:477:G:H2'	31:BA:478:A:C8	2.29	0.67
34:BG:38:TYR:CD1	34:BG:45:GLN:HB3	2.29	0.67
35:BH:26:PHE:N	35:BH:26:PHE:CD1	2.63	0.67
38:BK:109:ILE:HG22	38:BK:137:VAL:O	1.95	0.67
48:BU:29:PHE:HE1	48:BU:31:LEU:HB3	1.59	0.67
48:BU:85:LEU:HD12	48:BU:85:LEU:O	1.93	0.67
54:CA:1047:G:H5'	44:CQ:4:LYS:HD2	1.77	0.67
54:CA:1240:U:O2'	37:CJ:38:LEU:HD23	1.95	0.67
54:CA:1363:A:H4'	54:CA:1364:U:OP1	1.94	0.67
52:CD:9:A:H4'	52:CD:46:G:H4'	1.76	0.67
48:CU:50:ILE:H	48:CU:50:ILE:HD12	1.59	0.67
55:DA:1011:G:H4'	55:DA:1012:U:OP1	1.94	0.67
55:DA:1211:U:H4'	55:DA:1212:G:OP2	1.94	0.67
55:DA:163:U:H2'	55:DA:164:U:H5'	1.75	0.67
55:DA:1937:A:O2'	55:DA:1938:A:OP1	2.09	0.67
55:DA:2148:G:O2'	55:DA:2149:G:H5'	1.93	0.67
55:DA:898:C:H3'	55:DA:899:A:C5'	2.23	0.67
55:DA:997:G:OP1	16:D1:93:LYS:HD2	1.93	0.67
4:DE:24:THR:HG21	4:DE:188:VAL:CG1	2.19	0.67
5:DF:157:VAL:HB	5:DF:194:MET:HB3	1.75	0.67
57:DY:130:THR:HG21	56:DJ:14:GLN:CD	2.13	0.67
58:DL:18:THR:CB	58:DL:19:PRO:CD	2.39	0.67
11:DO:35:HIS:O	11:DO:36:LYS:O	2.13	0.67
18:DS:15:ARG:HA	18:DS:18:ARG:HD2	1.75	0.67
55:DA:1056:G:P	57:DY:35:LYS:HD3	2.34	0.67
30:A8:30:ARG:O	30:A8:31:HIS:CG	2.47	0.67
1:AA:1249:U:O2	1:AA:1249:U:C2'	2.41	0.67
1:AA:1520:U:H2'	1:AA:1521:G:O4'	1.94	0.67
1:AA:2067:G:H4'	1:AA:2068:U:OP2	1.94	0.67
1:AA:2286:A:C5'	1:AA:2287:A:O4'	2.42	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:68:G:H3'	1:AA:69:C:H6	1.60	0.67
2:AB:39:A:C2	2:AB:44:G:C2	2.82	0.67
5:AF:37:VAL:O	5:AF:40:GLN:HB2	1.93	0.67
7:AH:137:ASP:HB3	7:AH:141:VAL:HG23	1.74	0.67
8:AK:8:PRO:HD3	8:AK:15:VAL:HG23	1.75	0.67
9:AM:30:ILE:HG22	9:AM:34:LEU:CD2	2.24	0.67
10:AN:43:VAL:HG23	10:AN:56:ASP:O	1.94	0.67
11:AO:97:PRO:O	11:AO:98:GLU:HB3	1.94	0.67
12:AP:28:ALA:HB2	12:AP:67:ARG:NH1	2.08	0.67
12:AP:75:THR:CA	12:AP:88:GLY:HA2	2.20	0.67
21:AV:75:ASN:O	21:AV:84:GLU:HG2	1.94	0.67
31:BA:1321:C:N4	31:BA:1322:C:H41	1.92	0.67
31:BA:328:C:H2'	31:BA:328:C:O2	1.94	0.67
31:BA:57:G:H2'	31:BA:58:C:H6	1.58	0.67
33:BF:129:ALA:HB3	33:BF:132:ARG:HD3	1.77	0.67
34:BG:155:LEU:O	34:BG:159:ARG:HG3	1.94	0.67
31:BA:1151:A:H1'	40:BM:39:PRO:HB2	1.77	0.67
51:BX:9:ARG:HH21	51:BX:10:ARG:NE	1.92	0.67
54:CA:160:A:H2'	54:CA:161:A:O4'	1.94	0.67
54:CA:659:U:OP1	45:CR:8:LYS:HE3	1.94	0.67
54:CA:65:U:H5'	54:CA:66:G:OP1	1.94	0.67
54:CA:1297:C:H2'	37:CJ:114:ARG:HH22	1.58	0.67
38:CK:116:LYS:HE2	38:CK:116:LYS:HA	1.76	0.67
26:D4:34:GLU:HG2	26:D4:35:VAL:N	2.08	0.67
55:DA:1288:U:O2'	55:DA:1647:G:N2	2.28	0.67
55:DA:372:G:O2'	55:DA:373:U:P	2.52	0.67
3:DD:10:THR:HG23	3:DD:13:ARG:CB	2.24	0.67
6:DG:77:ILE:HG22	6:DG:77:ILE:O	1.95	0.67
56:DI:13:SER:O	56:DI:16:THR:HG22	1.95	0.67
56:DJ:5:ILE:HG23	56:DJ:9:LYS:HG3	1.76	0.67
55:DA:662:G:OP1	11:DO:15:ARG:NE	2.28	0.67
14:DQ:106:ARG:HA	14:DQ:110:LEU:CD2	2.23	0.67
18:DS:70:TYR:HD2	18:DS:70:TYR:H	1.42	0.67
21:DV:9:TYR:CE2	21:DV:61:LEU:HD23	2.29	0.67
57:DY:56:ASN:HA	57:DY:60:ARG:CG	2.24	0.67
57:DY:98:LYS:HB3	57:DY:102:LYS:HE3	1.75	0.67
16:A1:25:TRP:O	16:A1:28:ARG:HB2	1.93	0.67
17:A2:61:VAL:HG13	17:A2:62:LEU:N	2.09	0.67
1:AA:1359:A:C8	1:AA:1359:A:C4'	2.77	0.67
1:AA:1431:U:H2'	1:AA:1432:C:C6	2.29	0.67
1:AA:1504:C:C2'	1:AA:1505:C:H5'	2.24	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1464:C:O2'	1:AA:1528:A:H8	1.71	0.67
1:AA:2645:G:H4'	1:AA:2732:G:HO2'	1.59	0.67
7:AH:102:ALA:CA	7:AH:117:PRO:HD3	2.23	0.67
10:AN:2:ILE:HG23	10:AN:6:THR:CB	2.24	0.67
11:AO:85:LEU:H	11:AO:85:LEU:HD23	1.59	0.67
23:AZ:82:LEU:CG	23:AZ:83:GLU:H	2.03	0.67
31:BA:1004:A:C2'	31:BA:1005:A:O5'	2.43	0.67
31:BA:843:U:H5'	31:BA:848:C:C6	2.29	0.67
54:CA:1508:G:H2'	54:CA:1509:C:H6	1.58	0.67
54:CA:192:U:H4'	50:CW:103:GLY:HA2	1.75	0.67
52:CB:7:A:H5'	52:CB:8:U:OP2	1.93	0.67
37:CJ:113:GLU:HB2	37:CJ:119:ARG:CG	2.22	0.67
38:CK:82:HIS:CD2	38:CK:82:HIS:C	2.67	0.67
42:CO:55:VAL:HG12	42:CO:56:ALA:H	1.58	0.67
49:CV:5:LEU:N	49:CV:5:LEU:HD12	2.09	0.67
7:DH:125:VAL:HA	7:DH:126:PRO:HB3	1.74	0.67
8:DK:88:ILE:HG12	8:DK:122:GLU:N	2.10	0.67
55:DA:1060:U:H5	58:DL:74:ALA:HB2	1.58	0.67
11:DO:84:ASN:HA	11:DO:115:LEU:O	1.92	0.67
57:DY:26:LEU:O	57:DY:111:LEU:HD22	1.94	0.67
57:DY:57:THR:HG22	57:DY:58:LEU:N	2.08	0.67
57:DY:74:LEU:CB	57:DY:120:LYS:HE2	2.25	0.67
57:DY:25:PHE:C	57:DY:82:PHE:HZ	1.96	0.67
26:A4:63:TYR:HE2	49:BV:41:VAL:HG22	1.58	0.67
1:AA:2286:A:OP2	28:A6:28:ARG:NH1	2.27	0.67
30:A8:49:VAL:HG13	30:A8:50:LEU:CD2	2.19	0.67
1:AA:1177:A:H5'	1:AA:1178:C:OP1	1.94	0.67
1:AA:1348:G:H2'	1:AA:1349:A:H5''	1.76	0.67
1:AA:2585:U:O2'	1:AA:2586:C:H5'	1.94	0.67
5:AF:24:LEU:HB3	5:AF:25:PRO:CD	2.19	0.67
6:AG:67:LYS:HZ3	26:A4:6:HIS:CE1	2.12	0.67
8:AK:62:LYS:HD2	8:AK:62:LYS:O	1.94	0.67
1:AA:558:G:OP2	9:AM:111:PRO:HD2	1.94	0.67
31:BA:457:C:H2'	31:BA:458:C:C6	2.30	0.67
31:BA:628:G:H2'	31:BA:629:G:C8	2.30	0.67
52:BB:10:G:H3'	52:BB:11:C:C5	2.27	0.67
32:BE:187:LEU:HD13	32:BE:187:LEU:O	1.94	0.67
42:BO:40:VAL:HG21	42:BO:78:GLN:CA	2.25	0.67
45:BR:39:LEU:HD12	45:BR:56:LEU:HB2	1.76	0.67
54:CA:130:A:C8	47:CT:63:ARG:HG3	2.30	0.67
54:CA:69:G:N2	54:CA:73:G:C8	2.63	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:CM:9:ARG:HG2	40:CM:69:ASN:OD1	1.94	0.67
40:CM:61:GLU:OE2	44:CQ:45:ARG:NH1	2.28	0.67
45:CR:87:ILE:CG2	45:CR:88:ARG:H	2.02	0.67
46:CS:22:THR:HA	46:CS:33:ILE:HG12	1.75	0.67
46:CS:4:ILE:HD12	46:CS:4:ILE:N	2.09	0.67
55:DA:1266:G:OP1	27:D5:19:ARG:HD2	1.93	0.67
27:D5:20:ARG:HA	27:D5:23:HIS:CE1	2.29	0.67
55:DA:1188:U:C5'	17:D2:79:VAL:HG22	2.25	0.67
55:DA:2347:C:H2'	55:DA:2348:U:C6	2.30	0.67
55:DA:754:C:H2'	55:DA:755:C:C6	2.29	0.67
4:DE:18:ASP:O	4:DE:19:ARG:O	2.12	0.67
5:DF:184:TYR:O	5:DF:188:ARG:HG3	1.94	0.67
6:DG:76:SER:O	6:DG:77:ILE:HD12	1.94	0.67
56:DI:5:ILE:O	56:DI:5:ILE:HG22	1.94	0.67
56:DJ:13:SER:CB	56:DJ:17:VAL:HG13	2.21	0.67
10:DN:93:PRO:HB3	10:DN:114:ILE:CD1	2.24	0.67
21:DV:186:GLU:HG3	21:DV:186:GLU:O	1.95	0.67
21:DV:61:LEU:HD11	21:DV:65:GLN:HG3	1.75	0.67
21:DV:76:LEU:HD23	21:DV:76:LEU:H	1.59	0.67
57:DY:142:LEU:CG	57:DY:143:GLN:N	2.56	0.67
1:AA:994:C:OP1	16:A1:53:ARG:NH2	2.27	0.67
1:AA:1610:A:H5''	1:AA:1611:C:OP2	1.94	0.67
1:AA:1954:G:O2'	1:AA:1956:U:C5	2.45	0.67
1:AA:484:C:H2'	1:AA:485:C:H6	1.57	0.67
2:AB:111:U:H2'	2:AB:112:G:C8	2.30	0.67
8:AK:10:GLU:OE1	8:AK:11:ASN:HB2	1.94	0.67
31:BA:535:A:H4'	31:BA:536:C:OP1	1.92	0.67
12:AP:56:ARG:HH21	52:BB:52:G:H4'	1.59	0.67
52:BB:9:A:H2	52:BB:11:C:H41	1.41	0.67
32:BE:7:VAL:HG22	32:BE:8:LYS:N	2.10	0.67
34:BG:31:CYS:C	34:BG:33:MET:N	2.48	0.67
34:BG:26:CYS:HA	34:BG:31:CYS:HB2	1.77	0.67
37:BJ:155:ARG:HG2	37:BJ:156:TRP:N	2.09	0.67
40:BM:30:SER:HB3	40:BM:84:GLN:HE21	1.58	0.67
54:CA:1024:G:C3'	54:CA:1025:U:H5''	2.25	0.67
54:CA:121:C:H5'	54:CA:122:G:OP1	1.95	0.67
54:CA:737:A:H2'	54:CA:738:C:H6	1.60	0.67
52:CB:19:G:N2	52:CB:56:C:N4	2.41	0.67
46:CS:50:LYS:HD3	46:CS:51:VAL:N	2.09	0.67
55:DA:2131:G:C5'	55:DA:2132:U:H5''	2.21	0.67
55:DA:2173:A:H3'	55:DA:2174:C:C6	2.30	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2401:U:H2'	55:DA:2402:C:H5''	1.75	0.67
55:DA:366:C:H5	55:DA:403:U:O2'	1.77	0.67
55:DA:1657:C:H4'	4:DE:133:LYS:HB3	1.76	0.67
4:DE:170:LEU:CD2	4:DE:185:LYS:HB2	2.22	0.67
5:DF:107:LYS:CD	5:DF:206:ILE:HD13	2.21	0.67
9:DM:6:PRO:HG3	9:DM:41:ASP:HB2	1.75	0.67
14:DQ:103:GLU:O	14:DQ:106:ARG:HG3	1.93	0.67
12:DP:140:ALA:HB2	21:DV:53:ILE:HD11	1.76	0.67
57:DY:111:LEU:C	57:DY:112:LEU:HD22	2.13	0.67
55:DA:1084:A:O2'	57:DY:53:VAL:HG21	1.94	0.67
23:DZ:80:LEU:O	23:DZ:81:LYS:HE2	1.95	0.67
16:A1:98:LEU:C	16:A1:100:VAL:N	2.44	0.67
1:AA:1054:A:H2'	1:AA:1055:G:C8	2.29	0.67
1:AA:1534:G:H3'	1:AA:1535:U:H5'	1.77	0.67
1:AA:1970:A:H5'	1:AA:1971:A:OP1	1.94	0.67
1:AA:2387:U:H5'	1:AA:2388:A:OP2	1.94	0.67
1:AA:2068:U:N3	1:AA:2430:A:H2	1.92	0.67
1:AA:2439:A:C8	1:AA:2439:A:H5'	2.29	0.67
1:AA:2795:G:H3'	1:AA:2797:U:C5'	2.25	0.67
20:AU:27:VAL:O	20:AU:27:VAL:HG23	1.94	0.67
20:AU:47:LYS:HG3	20:AU:60:PHE:HB3	1.76	0.67
23:AZ:76:ARG:HG3	23:AZ:94:LEU:HD13	1.76	0.67
31:BA:197:A:N6	31:BA:221:C:H5'	2.10	0.67
12:AP:56:ARG:NH2	52:BB:52:G:H4'	2.09	0.67
34:BG:153:ARG:NH1	34:BG:181:MET:HB2	2.09	0.67
43:BP:95:GLY:O	43:BP:110:ARG:HB3	1.95	0.67
49:BV:40:ILE:HG21	49:BV:66:MET:O	1.95	0.67
54:CA:1139:G:H1	54:CA:1144:G:H22	1.43	0.67
54:CA:501:C:H2'	54:CA:502:G:H8	1.59	0.67
52:CD:9:A:H4'	52:CD:46:G:C4'	2.24	0.67
37:CJ:108:ALA:HB2	37:CJ:123:GLU:HG2	1.75	0.67
41:CN:20:TYR:HB2	41:CN:31:THR:HG23	1.77	0.67
50:CW:82:SER:O	50:CW:86:ARG:CB	2.42	0.67
55:DA:1310:G:OP2	29:D7:9:ARG:NH1	2.28	0.67
30:D8:48:PHE:H	30:D8:48:PHE:HD1	1.41	0.67
55:DA:1024:G:C3'	55:DA:1025:G:H5''	2.25	0.67
6:DG:131:TYR:HB3	6:DG:159:VAL:HG13	1.76	0.67
6:DG:81:LYS:O	6:DG:82:LEU:HB2	1.92	0.67
12:DP:34:LEU:HD23	12:DP:104:PHE:CD1	2.30	0.67
21:DV:194:PRO:HG2	21:DV:196:VAL:HG13	1.70	0.67
16:A1:90:VAL:CG2	17:A2:39:LEU:HB3	2.24	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2391:G:O6	1:AA:2425:A:H8	1.78	0.67
1:AA:2716:U:O2'	1:AA:2717:G:H5'	1.95	0.67
1:AA:2815:C:H2'	1:AA:2816:C:C6	2.30	0.67
1:AA:704:G:C2'	1:AA:726:G:H22	2.07	0.67
3:AD:2:ALA:O	3:AD:3:VAL:HB	1.92	0.67
5:AF:124:LEU:HG	5:AF:124:LEU:O	1.93	0.67
20:AU:81:LYS:HB3	20:AU:97:ARG:CD	2.25	0.67
32:BE:97:TRP:CE2	32:BE:101:MET:HG3	2.30	0.67
36:BI:68:PRO:HG3	36:BI:71:ARG:HH21	1.60	0.67
40:BM:74:ILE:N	40:BM:74:ILE:HD13	2.10	0.67
42:BO:117:ARG:HB2	42:BO:122:THR:HB	1.77	0.67
54:CA:1152:A:H2'	54:CA:1153:C:H6	1.59	0.67
54:CA:1301:U:H2'	54:CA:1301:U:O2	1.95	0.67
37:CJ:13:GLN:O	37:CJ:24:THR:HG21	1.95	0.67
38:CK:33:GLU:HG2	38:CK:59:LEU:HD11	1.76	0.67
40:CM:78:ASN:O	40:CM:82:ILE:HG12	1.94	0.67
49:CV:29:ARG:HD3	49:CV:30:LEU:HD13	1.77	0.67
54:CA:1221:G:H4'	49:CV:77:THR:HG21	1.75	0.67
49:CV:9:VAL:O	49:CV:9:VAL:HG12	1.94	0.67
16:D1:52:ARG:HA	16:D1:55:ARG:HE	1.60	0.67
16:D1:81:HIS:NE2	16:D1:117:GLN:HG3	2.09	0.67
6:DG:112:PRO:HB3	26:D4:37:SER:CB	2.25	0.67
55:DA:1533:C:H3'	55:DA:1534:G:H5''	1.77	0.67
55:DA:2838:G:H1'	13:D0:45:ARG:HH21	1.60	0.67
55:DA:654(O):G:H2'	55:DA:654(P):G:C8	2.29	0.67
55:DA:886:C:O2	55:DA:887:A:N1	2.27	0.67
4:DE:26:ILE:HD13	4:DE:26:ILE:C	2.15	0.67
56:DI:1:MET:SD	56:DI:5:ILE:CG2	2.83	0.67
16:A1:34:LYS:HA	16:A1:34:LYS:HE2	1.75	0.67
16:A1:50:ARG:NH1	17:A2:72:VAL:HG11	2.10	0.67
28:A6:48:VAL:HG13	28:A6:49:HIS:N	2.10	0.67
1:AA:2094:G:OP1	8:AK:22:LYS:HD2	1.94	0.67
1:AA:900:A:H3'	1:AA:901:A:C8	2.24	0.67
2:AB:82:G:N2	2:AB:95:U:H1'	2.09	0.67
3:AD:28:GLU:HB2	3:AD:29:PRO:CD	2.25	0.67
9:AM:15:LEU:CG	9:AM:134:ARG:HE	2.05	0.67
14:AQ:25:ARG:HH11	14:AQ:25:ARG:CB	2.07	0.67
15:AR:6:LEU:HA	15:AR:9:LEU:HD12	1.77	0.67
19:AT:50:LYS:N	19:AT:87:GLN:HE22	1.91	0.67
31:BA:1152:A:H2'	31:BA:1153:C:C6	2.30	0.67
31:BA:949:A:H1'	31:BA:1364:U:H3	1.60	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:BL:96:LEU:HG	39:BL:102:LEU:HB2	1.75	0.67
40:BM:40:LEU:CG	40:BM:41:PRO:HD2	2.24	0.67
41:BN:27:ASN:OD1	41:BN:55:LYS:HB3	1.94	0.67
42:BO:86:ARG:HB2	42:BO:101:VAL:CG2	2.24	0.67
44:CQ:8:GLU:OE2	44:CQ:11:LYS:HD2	1.95	0.67
55:DA:1240:U:O2'	55:DA:1241:A:H5'	1.95	0.67
55:DA:27:G:O2'	55:DA:28:A:H8	1.78	0.67
55:DA:518:G:H4'	18:DS:18:ARG:HH12	1.58	0.67
55:DA:793:A:O2'	55:DA:794:G:OP2	2.13	0.67
3:DD:44:ASN:HB3	3:DD:49:ILE:HG22	1.76	0.67
4:DE:105:THR:OG1	4:DE:166:THR:HG23	1.95	0.67
56:DI:11:GLU:O	56:DI:14:GLN:HG2	1.95	0.67
58:DL:110:GLN:HG3	58:DL:111:LYS:CE	2.25	0.67
57:DY:94:VAL:O	57:DY:95:GLN:CB	2.43	0.67
1:AA:51:G:O2'	1:AA:119:A:N1	2.22	0.67
1:AA:2110:G:O2'	1:AA:2111:C:OP1	2.13	0.67
1:AA:832:G:C5'	11:AO:45:LEU:HD11	2.25	0.67
2:AB:15:A:C3'	2:AB:16:G:H5'	2.22	0.67
1:AA:2784:C:H1'	4:AE:37:ARG:NH2	2.09	0.67
5:AF:7:TYR:CE1	5:AF:10:PRO:HG3	2.30	0.67
6:AG:131:TYR:HB3	6:AG:159:VAL:HG13	1.75	0.67
1:AA:1138:G:H21	9:AM:106:MET:HE3	1.59	0.67
11:AO:124:LYS:HZ3	11:AO:143:GLY:HA3	1.57	0.67
20:AU:39:VAL:HG23	20:AU:40:GLU:N	2.10	0.67
21:AV:30:ASN:O	21:AV:32:HIS:N	2.28	0.67
31:BA:1125:U:OP2	31:BA:1145:C:N4	2.28	0.67
15:AR:107:ASP:HB2	31:BA:1432:G:OP1	1.95	0.67
31:BA:406:G:N2	34:BG:119:GLN:HE22	1.90	0.67
54:CA:1305:G:OP1	51:CX:2:GLY:HA3	1.95	0.67
54:CA:629:G:H5''	54:CA:630:G:P	2.35	0.67
54:CA:644:G:C2'	54:CA:645:C:H5'	2.25	0.67
32:CE:107:THR:HA	32:CE:110:GLN:HG3	1.77	0.67
35:CH:51:VAL:HB	35:CH:52:PRO:HD3	1.77	0.67
42:CO:19:ARG:HB3	42:CO:19:ARG:NH1	2.09	0.67
44:CQ:29:ARG:HD3	44:CQ:40:CYS:HB2	1.77	0.67
54:CA:192:U:C4'	50:CW:103:GLY:HA2	2.25	0.67
17:D2:35:LEU:HD23	17:D2:35:LEU:O	1.95	0.67
55:DA:1266:G:O2'	55:DA:1267:U:OP2	2.13	0.67
55:DA:2391:G:H1'	55:DA:2429:G:N2	2.09	0.67
55:DA:2512:C:H4'	4:DE:122:PHE:CE2	2.30	0.67
55:DA:2682:U:H5''	4:DE:11:MET:HB2	1.76	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:813:U:H2'	55:DA:814:C:C6	2.30	0.67
55:DA:893:C:H2'	55:DA:894:C:C6	2.30	0.67
3:DD:35:LYS:CD	3:DD:104:TYR:CD1	2.76	0.67
7:DH:80:SER:O	7:DH:81:GLU:HB2	1.95	0.67
58:DL:10:LEU:CD2	58:DL:55:VAL:HG11	2.25	0.67
58:DL:53:VAL:HG12	58:DL:72:PRO:HG2	1.75	0.67
12:DP:35:VAL:HG13	12:DP:130:LYS:HB3	1.76	0.67
19:DT:43:VAL:CG1	19:DT:51:VAL:HG21	2.25	0.67
57:DY:73:GLY:HA2	57:DY:119:ALA:O	1.95	0.67
57:DY:51:LEU:HD11	57:DY:82:PHE:C	2.16	0.67
1:AA:530:G:O6	1:AA:2023:G:OP1	2.12	0.66
1:AA:2776:A:O2'	1:AA:2777:G:OP2	2.12	0.66
1:AA:2853:C:H2'	1:AA:2854:G:H8	1.59	0.66
1:AA:956:G:H5'	1:AA:957:A:OP2	1.95	0.66
7:AH:9:ILE:HB	7:AH:49:VAL:HB	1.76	0.66
8:AK:5:LEU:HD12	8:AK:5:LEU:N	2.10	0.66
11:AO:18:ARG:O	11:AO:19:VAL:HB	1.95	0.66
12:AP:19:GLY:O	12:AP:98:LYS:HD3	1.95	0.66
18:AS:59:VAL:HG23	18:AS:65:LEU:N	2.08	0.66
18:AS:65:LEU:HD22	18:AS:67:ASP:H	1.60	0.66
19:AT:65:ARG:HG3	19:AT:65:ARG:HH11	1.60	0.66
20:AU:56:PRO:HB2	20:AU:57:GLN:NE2	2.10	0.66
21:AV:141:VAL:CG1	21:AV:141:VAL:O	2.42	0.66
24:AW:53:LEU:HD22	24:AW:57:ILE:HD11	1.77	0.66
31:BA:1053:G:O6	31:BA:1199:U:H2'	1.95	0.66
32:BE:20:GLU:HG2	32:BE:189:ASP:OD2	1.95	0.66
35:BH:135:THR:O	35:BH:138:ALA:HB3	1.95	0.66
35:BH:137:GLU:O	35:BH:141:GLN:HG3	1.95	0.66
50:BW:14:LYS:O	50:BW:18:GLN:HG3	1.94	0.66
54:CA:1014:A:H4'	49:CV:14:HIS:CE1	2.29	0.66
54:CA:629:G:H5"	54:CA:630:G:OP2	1.94	0.66
54:CA:644:G:H2'	54:CA:645:C:H5'	1.77	0.66
49:CV:63:THR:HG23	49:CV:65:ASN:HD21	1.60	0.66
13:D0:105:ARG:O	13:D0:105:ARG:HG3	1.93	0.66
16:D1:92:ARG:NH2	16:D1:94:ASN:HD22	1.92	0.66
55:DA:1484:G:H2'	55:DA:1485:G:C5'	2.12	0.66
55:DA:2101:G:H2'	55:DA:2102:U:C6	2.30	0.66
55:DA:2641:G:P	9:DM:74:ARG:HE	2.17	0.66
55:DA:2761:G:H8	55:DA:2761:G:H5'	1.59	0.66
55:DA:265:A:N6	55:DA:427:U:O2'	2.28	0.66
6:DG:77:ILE:HG22	6:DG:80:PHE:H	1.60	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:DL:77:LEU:HD22	58:DL:108:ALA:HB2	1.76	0.66
58:DL:18:THR:CG2	58:DL:19:PRO:HD3	2.24	0.66
11:DO:11:GLY:O	11:DO:12:ALA:CB	2.43	0.66
12:DP:81:VAL:O	12:DP:82:ARG:NE	2.27	0.66
19:DT:53:LYS:HB2	19:DT:82:GLN:HB3	1.76	0.66
21:DV:191:VAL:CG1	21:DV:197:ILE:HG13	2.13	0.66
16:A1:24:TYR:HB2	16:A1:29:SER:HB3	1.77	0.66
27:A5:47:PRO:HB3	27:A5:56:LYS:HZ1	1.60	0.66
30:A8:14:VAL:HG12	30:A8:15:LYS:N	2.10	0.66
1:AA:1444(A):A:H2'	1:AA:1444(A):A:N3	2.10	0.66
1:AA:1454:U:O2'	1:AA:1455:G:C8	2.47	0.66
1:AA:1803:A:C2	1:AA:1822:G:N3	2.64	0.66
1:AA:829:A:N7	1:AA:2248:C:H5'	2.11	0.66
1:AA:535:C:O2'	1:AA:536:A:H5'	1.95	0.66
4:AE:4:ILE:HD11	4:AE:28:ALA:HB3	1.76	0.66
5:AF:7:TYR:HE1	5:AF:10:PRO:HG3	1.60	0.66
6:AG:180:PHE:C	6:AG:182:LYS:H	1.99	0.66
7:AH:33:LEU:HD13	7:AH:75:ALA:HA	1.77	0.66
9:AM:55:VAL:HB	9:AM:126:PRO:CB	2.24	0.66
31:BA:173:U:O2'	31:BA:174:C:OP1	2.12	0.66
31:BA:85:U:O2	31:BA:85:U:H2'	1.94	0.66
33:BF:182:ILE:HG12	33:BF:203:PHE:HA	1.76	0.66
42:BO:32:PHE:HB3	42:BO:84:LEU:CD2	2.25	0.66
54:CA:738:C:H2'	54:CA:739:C:H6	1.61	0.66
54:CA:753:A:H4'	54:CA:754:C:O5'	1.96	0.66
34:CG:198:VAL:HG12	34:CG:199:ASN:H	1.61	0.66
34:CG:79:PHE:HD2	34:CG:79:PHE:C	1.98	0.66
40:CM:92:THR:HG23	40:CM:93:GLY:H	1.58	0.66
45:CR:3:ILE:HD13	45:CR:3:ILE:H	1.58	0.66
46:CS:6:LEU:HB3	46:CS:17:TYR:HD2	1.60	0.66
50:CW:96:GLY:O	50:CW:99:LEU:HD21	1.94	0.66
55:DA:1066:U:H3'	55:DA:1066:U:O2	1.95	0.66
55:DA:1077:A:OP1	55:DA:1077:A:H4'	1.94	0.66
55:DA:573:G:N1	55:DA:2031:A:OP2	2.20	0.66
55:DA:270(M):U:H1'	55:DA:270(N):G:C6	2.31	0.66
55:DA:2789:C:C2'	55:DA:2790:A:H5''	2.24	0.66
55:DA:2790:A:C2	55:DA:2894:G:H5''	2.30	0.66
7:DH:148:ILE:O	7:DH:151:ILE:HG12	1.95	0.66
58:DL:109:LYS:CA	58:DL:120:LEU:HD21	2.25	0.66
58:DL:133:SER:CA	58:DL:137:GLU:OE1	2.42	0.66
55:DA:2683:C:P	15:DR:53:ARG:HH22	2.17	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:107:VAL:CG1	57:DY:108:LYS:N	2.52	0.66
57:DY:137:GLU:HG3	57:DY:138:LEU:N	2.07	0.66
57:DY:6:ASN:O	57:DY:9:LEU:HB3	1.94	0.66
1:AA:1824:G:O2'	1:AA:1825:A:H5'	1.95	0.66
1:AA:2068:U:N3	1:AA:2430:A:C2	2.62	0.66
1:AA:270(E):G:H2'	1:AA:270(F):U:H6	1.61	0.66
1:AA:588:U:H2'	1:AA:589:C:C6	2.30	0.66
1:AA:603:A:H4'	1:AA:604:G:O5'	1.95	0.66
1:AA:644:A:H4'	1:AA:645:C:H5	1.60	0.66
3:AD:27:THR:O	3:AD:29:PRO:HD2	1.94	0.66
4:AE:60:ASN:O	4:AE:61:ARG:HB2	1.93	0.66
31:BA:1175:G:C6	31:BA:1176:A:N6	2.63	0.66
31:BA:511:C:C2	31:BA:512:U:C5	2.83	0.66
31:BA:524:G:H2'	31:BA:525:C:C6	2.30	0.66
31:BA:87:A:H2'	31:BA:87:A:N3	2.10	0.66
32:BE:137:ARG:HD3	32:BE:137:ARG:C	2.16	0.66
34:BG:19:LEU:O	34:BG:21:LEU:N	2.28	0.66
46:BS:8:ARG:HH11	46:BS:8:ARG:HG2	1.60	0.66
54:CA:1298:C:N4	37:CJ:114:ARG:HB3	2.10	0.66
54:CA:323:U:H5'	50:CW:23:ARG:HB2	1.76	0.66
46:CS:20:VAL:HG23	46:CS:35:LYS:HA	1.75	0.66
49:CV:63:THR:HG23	49:CV:66:MET:HG2	1.78	0.66
13:D0:10:LEU:O	13:D0:12:ARG:HG3	1.94	0.66
28:D6:25:LYS:HE2	28:D6:27:LYS:CE	2.24	0.66
55:DA:1060:U:H1'	55:DA:1061:U:H3'	1.77	0.66
55:DA:1062:G:O2'	55:DA:1077:A:N6	2.23	0.66
55:DA:1177:A:C4'	55:DA:1178:C:H5"	2.24	0.66
55:DA:2135:A:H3'	55:DA:2136:C:C5	2.30	0.66
55:DA:2591:C:OP1	3:DD:239:ARG:HG3	1.96	0.66
55:DA:467:G:OP1	29:D7:33:ARG:NH1	2.27	0.66
5:DF:180:GLY:O	5:DF:181:LEU:C	2.32	0.66
6:DG:173:LEU:HD22	6:DG:178:PHE:CZ	2.30	0.66
7:DH:150:ALA:O	7:DH:152:ARG:N	2.26	0.66
8:DK:114:LEU:CD1	8:DK:128:LEU:HD12	2.25	0.66
58:DL:59:ILE:HG22	58:DL:60:TYR:H	1.61	0.66
21:DV:178:GLU:C	21:DV:180:VAL:N	2.49	0.66
21:DV:187:ALA:HB2	21:DV:193:GLU:CG	2.21	0.66
57:DY:16:ASN:O	57:DY:19:ARG:O	2.13	0.66
57:DY:46:GLN:NE2	57:DY:46:GLN:HA	2.10	0.66
57:DY:99:SER:O	57:DY:100:ASN:C	2.34	0.66
28:A6:14:THR:HG22	28:A6:50:ARG:O	1.95	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1682:G:H2'	1:AA:1683:C:C6	2.30	0.66
3:AD:108:PRO:HB3	3:AD:143:HIS:CE1	2.31	0.66
1:AA:2820:A:H62	4:AE:192:ASN:H	0.68	0.66
21:AV:11:GLU:HG3	21:AV:12:GLY:H	1.60	0.66
21:AV:62:PRO:O	21:AV:63:ASP:C	2.34	0.66
31:BA:706:A:H1'	41:BN:29:ILE:HD11	1.77	0.66
35:CH:99:GLY:O	35:CH:117:ASP:HA	1.96	0.66
42:CO:7:ILE:O	42:CO:11:VAL:HG23	1.95	0.66
54:CA:1320:C:OP1	49:CV:70:LYS:HE3	1.95	0.66
26:D4:4:GLY:O	26:D4:5:ILE:HB	1.96	0.66
55:DA:1060:U:H5	58:DL:74:ALA:CB	2.09	0.66
55:DA:1084:A:N6	57:DY:31:GLY:HA3	2.10	0.66
55:DA:1771:C:H1'	55:DA:1786:A:C8	2.30	0.66
3:DD:44:ASN:HB2	3:DD:48:ARG:O	1.96	0.66
6:DG:113:ARG:NH1	6:DG:142:PRO:HA	2.10	0.66
8:DK:92:VAL:HG22	8:DK:92:VAL:O	1.96	0.66
58:DL:76:TYR:CG	58:DL:77:LEU:N	2.61	0.66
11:DO:111:ARG:HH12	11:DO:148:LEU:HD21	1.60	0.66
18:DS:29:LEU:HD21	18:DS:33:ARG:CZ	2.25	0.66
23:DZ:87:PRO:O	23:DZ:91:LYS:N	2.19	0.66
27:A5:47:PRO:HB3	27:A5:56:LYS:NZ	2.10	0.66
1:AA:1278:A:OP1	13:A0:36:THR:HG22	1.95	0.66
1:AA:1579:A:H2'	1:AA:1580:A:O4'	1.96	0.66
1:AA:270(Z):U:O2'	1:AA:271(A):C:H5	1.78	0.66
1:AA:443:A:H5''	1:AA:444:C:OP1	1.96	0.66
3:AD:206:LEU:HD22	3:AD:211:ARG:HG2	1.76	0.66
1:AA:779:U:OP1	3:AD:49:ILE:HG23	1.96	0.66
3:AD:27:THR:HG21	3:AD:83:GLU:HG2	1.78	0.66
6:AG:112:PRO:HB2	26:A4:37:SER:HA	1.77	0.66
15:AR:50:ILE:HD11	15:AR:102:ILE:CG1	2.25	0.66
19:AT:63:LYS:CE	19:AT:63:LYS:H	2.07	0.66
21:AV:23:LYS:HD3	21:AV:40:ASP:HA	1.76	0.66
25:AX:8:LEU:HD13	25:AX:31:LEU:HA	1.76	0.66
25:AX:8:LEU:CD1	25:AX:31:LEU:HD12	2.24	0.66
31:BA:179:A:H2'	31:BA:180:U:C6	2.31	0.66
31:BA:38:G:H4'	31:BA:547:A:N6	2.11	0.66
32:BE:82:ARG:HA	32:BE:92:TYR:HE1	1.61	0.66
40:BM:8:LEU:HG	40:BM:96:ILE:CG2	2.19	0.66
40:BM:99:LYS:CD	40:BM:100:THR:H	2.08	0.66
42:BO:18:VAL:HG23	42:BO:19:ARG:H	1.59	0.66
43:BP:19:LEU:O	43:BP:22:ILE:HG13	1.95	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
46:BS:82:GLN:O	46:BS:83:GLU:HB2	1.96	0.66
48:BU:41:LYS:HA	48:BU:44:LEU:HD13	1.76	0.66
49:BV:42:PRO:O	49:BV:43:GLU:C	2.33	0.66
50:BW:89:ARG:NH1	50:BW:104:LEU:HG	2.11	0.66
54:CA:976:G:N2	54:CA:1362(A):C:OP2	2.28	0.66
34:CG:33:MET:CE	34:CG:37:PRO:HA	2.24	0.66
13:D0:3:HIS:O	13:D0:5:LYS:N	2.22	0.66
12:DP:80:GLU:OE2	22:D3:4:LYS:NZ	2.28	0.66
55:DA:1541:U:H2'	55:DA:1542:G:O4'	1.94	0.66
55:DA:2287:A:N6	55:DA:2344:U:N3	2.40	0.66
55:DA:2750:A:O2'	55:DA:2751:G:OP1	2.13	0.66
55:DA:2756:U:O2'	55:DA:2757:A:H5''	1.95	0.66
55:DA:2836:U:H2'	55:DA:2837:G:C8	2.30	0.66
4:DE:52:LEU:HB2	4:DE:75:VAL:CG2	2.25	0.66
11:DO:6:LEU:O	11:DO:7:ARG:HG2	1.95	0.66
57:DY:7:VAL:HG13	57:DY:8:GLU:H	1.55	0.66
57:DY:8:GLU:OE1	57:DY:52:PHE:CD1	2.40	0.66
57:DY:93:LEU:CG	57:DY:126:ALA:HB1	2.25	0.66
23:DZ:60:PHE:HE2	23:DZ:91:LYS:HZ2	1.41	0.66
16:A1:76:TYR:O	16:A1:80:ILE:HG12	1.96	0.66
1:AA:1043:C:H2'	1:AA:1044:G:C5'	2.25	0.66
1:AA:1083:U:H1'	1:AA:1086:A:H61	1.59	0.66
1:AA:644:A:N6	1:AA:2349:G:H1'	2.11	0.66
1:AA:74:A:O2'	1:AA:75:G:OP2	2.11	0.66
3:AD:76:PRO:HG2	3:AD:98:VAL:HG21	1.76	0.66
4:AE:77:ILE:C	4:AE:78:LEU:HD23	2.16	0.66
5:AF:101:LEU:O	5:AF:106:ARG:NH1	2.29	0.66
24:AW:65:ASN:HD22	24:AW:69:ARG:HH21	1.43	0.66
31:BA:1144:G:H22	31:BA:1146:A:N6	1.93	0.66
50:BW:100:ILE:HD12	50:BW:100:ILE:H	1.60	0.66
50:BW:88:VAL:HA	50:BW:91:LEU:HD12	1.76	0.66
54:CA:1529:G:H5''	54:CA:1530:G:OP2	1.95	0.66
54:CA:22:G:H5'	54:CA:885:G:O4'	1.96	0.66
54:CA:50:A:O2'	54:CA:52:G:C8	2.49	0.66
33:CF:21:ARG:HD3	33:CF:21:ARG:H	1.61	0.66
34:CG:23:GLY:HA3	34:CG:112:VAL:HG21	1.76	0.66
39:CL:48:GLU:N	39:CL:49:PRO:HD2	2.11	0.66
55:DA:444:C:H4'	5:DF:49:ALA:HB2	1.77	0.66
55:DA:607:U:N3	55:DA:621:A:C2	2.63	0.66
3:DD:27:THR:O	3:DD:29:PRO:HD2	1.95	0.66
5:DF:46:ARG:HH11	5:DF:46:ARG:CG	2.08	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:DL:132:ARG:CD	58:DL:137:GLU:OE2	2.43	0.66
55:DA:1266:G:C8	18:DS:15:ARG:NH1	2.64	0.66
20:DU:52:SER:OG	20:DU:53:PRO:HD3	1.95	0.66
57:DY:14:LYS:HA	57:DY:14:LYS:CE	2.23	0.66
26:A4:56:VAL:HA	26:A4:60:GLN:HE22	1.61	0.66
1:AA:1945:G:H2'	1:AA:1946:U:C6	2.31	0.66
1:AA:2528:U:C2'	1:AA:2529:G:H5''	2.26	0.66
1:AA:2756:U:H5''	1:AA:2757:A:OP1	1.95	0.66
3:AD:35:LYS:NZ	3:AD:64:ILE:O	2.24	0.66
3:AD:65:ILE:HD11	3:AD:67:PHE:CE2	2.30	0.66
4:AE:60:ASN:O	4:AE:62:PRO:HD2	1.95	0.66
9:AM:35:ARG:HB3	9:AM:42:TRP:CZ3	2.30	0.66
9:AM:71:ILE:H	9:AM:71:ILE:HD12	1.60	0.66
11:AO:62:LEU:O	11:AO:62:LEU:HD13	1.96	0.66
19:AT:89:ILE:HG21	19:AT:92:LEU:HG	1.77	0.66
23:AZ:8:SER:HB3	23:AZ:66:HIS:ND1	2.11	0.66
31:BA:1347:G:N2	31:BA:1373:G:H2'	2.10	0.66
31:BA:17:U:H2'	31:BA:18:C:H6	1.59	0.66
31:BA:329:A:H4'	31:BA:330:C:OP1	1.95	0.66
32:BE:92:TYR:CD2	32:BE:151:GLY:HA3	2.31	0.66
38:BK:84:ARG:NH1	38:BK:86:ILE:HD13	2.10	0.66
39:BL:127:LYS:O	39:BL:128:ARG:HG2	1.95	0.66
40:BM:12:ASP:HB3	40:BM:15:THR:HG23	1.78	0.66
53:C1:53:U:O2'	53:C1:54:U:H5'	1.95	0.66
54:CA:953:G:H5'	54:CA:965:A:H61	1.60	0.66
26:D4:68:ARG:CA	26:D4:68:ARG:NH1	2.58	0.66
28:D6:37:ARG:HA	28:D6:37:ARG:NE	2.09	0.66
55:DA:2308:G:C2	55:DA:2311:A:H2	2.14	0.66
55:DA:2566:A:O2'	55:DA:2567:G:OP2	2.12	0.66
55:DA:654(L):G:N2	55:DA:654(M):C:H1'	2.09	0.66
55:DA:701:G:C3'	55:DA:702:G:H5''	2.26	0.66
8:DK:69:LYS:HE2	8:DK:73:GLU:OE2	1.96	0.66
58:DL:18:THR:CG2	58:DL:42:ASN:OD1	2.43	0.66
12:DP:81:VAL:C	12:DP:82:ARG:HG2	2.15	0.66
12:DP:86:GLY:C	12:DP:88:GLY:N	2.48	0.66
21:DV:48:PHE:CE2	21:DV:71:VAL:HG11	2.31	0.66
24:DW:28:LYS:HD2	24:DW:53:LEU:HD21	1.78	0.66
23:DZ:91:LYS:CE	23:DZ:91:LYS:HA	2.15	0.66
16:A1:66:ASN:ND2	16:A1:70:ARG:HE	1.94	0.66
17:A2:73:SER:HB3	17:A2:83:ARG:O	1.96	0.66
22:A3:36:ILE:O	22:A3:36:ILE:HD13	1.96	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1011:G:H2'	1:AA:1013:C:O4'	1.96	0.66
1:AA:1237:A:H4'	1:AA:1238:G:O5'	1.96	0.66
1:AA:1528:A:N1	1:AA:1543:A:C2	2.64	0.66
1:AA:1729:A:N1	1:AA:1731:G:N7	2.44	0.66
1:AA:2348:U:C2'	1:AA:2349:G:H5''	2.25	0.66
1:AA:2867:G:O2'	1:AA:2868:A:H8	1.79	0.66
5:AF:24:LEU:CB	5:AF:25:PRO:HD2	2.22	0.66
6:AG:115:ARG:HH12	43:BP:7:VAL:CG2	2.06	0.66
7:AH:87:LEU:HD22	7:AH:162:ILE:HG22	1.76	0.66
1:AA:1006:C:H1'	9:AM:106:MET:CE	2.26	0.66
11:AO:15:ARG:O	11:AO:16:ARG:C	2.34	0.66
20:AU:12:THR:HG23	20:AU:26:LYS:HE2	1.77	0.66
21:AV:181:GLU:HG2	21:AV:185:GLU:OE2	1.96	0.66
31:BA:1401:G:C2	31:BA:1402:C:H1'	2.31	0.66
32:BE:187:LEU:HA	32:BE:201:ILE:O	1.96	0.66
49:BV:53:ASN:HB2	49:BV:77:THR:HG22	1.77	0.66
54:CA:186(C):G:H2'	54:CA:186(D):C:C6	2.29	0.66
54:CA:328:C:O2'	54:CA:329:A:OP2	2.14	0.66
33:CF:40:ARG:HG2	33:CF:55:VAL:HG11	1.78	0.66
38:CK:87:SER:HB2	38:CK:93:VAL:HB	1.77	0.66
55:DA:2357:U:OP1	22:D3:20:ARG:NH1	2.29	0.66
30:D8:52:LYS:N	30:D8:53:PRO:CD	2.41	0.66
55:DA:1237:A:H4'	55:DA:1238:G:O5'	1.94	0.66
55:DA:1431:U:H2'	55:DA:1432:C:H6	1.61	0.66
55:DA:483:A:H3'	55:DA:484:C:C6	2.30	0.66
3:DD:132:PRO:HG3	3:DD:190:TYR:CE1	2.31	0.66
7:DH:19:VAL:HG12	7:DH:20:ALA:N	2.10	0.66
8:DK:74:ASN:N	8:DK:74:ASN:HD22	1.93	0.66
58:DL:93:ARG:HG2	58:DL:135:GLY:HA3	1.77	0.66
11:DO:26:GLY:O	11:DO:28:GLY:N	2.28	0.66
21:DV:174:VAL:O	21:DV:175:VAL:CG1	2.43	0.66
57:DY:101:PRO:HG2	57:DY:102:LYS:N	2.10	0.66
57:DY:134:LEU:HA	57:DY:137:GLU:CG	2.24	0.66
1:AA:1028:A:N6	1:AA:1125:G:H2'	2.11	0.66
1:AA:1188:U:O2'	1:AA:1189:A:H5'	1.95	0.66
1:AA:1914:C:H3'	1:AA:1914:C:O2	1.94	0.66
1:AA:1918:A:O2'	1:AA:1919:A:N7	2.28	0.66
1:AA:2091:U:C3'	1:AA:2092:U:H5'	2.20	0.66
1:AA:2183:C:H2'	1:AA:2184:G:C8	2.29	0.66
1:AA:877:U:O5'	1:AA:877:U:H6	1.78	0.66
3:AD:242:ARG:HD2	3:AD:242:ARG:N	2.09	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1049:C:H42	7:AH:2:SER:HB2	1.60	0.66
23:AZ:8:SER:HB3	23:AZ:66:HIS:CE1	2.30	0.66
31:BA:1176:A:H8	31:BA:1176:A:O5'	1.77	0.66
31:BA:280:C:O2'	31:BA:281:G:P	2.54	0.66
32:BE:172:ILE:HD12	32:BE:172:ILE:H	1.61	0.66
39:BL:111:ARG:HD2	44:BQ:61:TRP:OXT	1.96	0.66
39:BL:40:LEU:HD13	39:BL:74:ILE:HD11	1.76	0.66
40:BM:49:VAL:O	40:BM:60:ARG:HB2	1.96	0.66
45:BR:79:ARG:O	45:BR:83:GLU:HB2	1.95	0.66
54:CA:630:G:O4'	54:CA:630:G:P	2.53	0.66
35:CH:53:LEU:HD12	35:CH:53:LEU:H	1.61	0.66
54:CA:738:C:H5"	36:CI:69:GLU:HB2	1.77	0.66
50:CW:89:ARG:NH2	50:CW:104:LEU:HD21	2.10	0.66
55:DA:2882:A:OP1	13:D0:96:ARG:NH1	2.28	0.66
55:DA:2751:G:O6	7:DH:2:SER:HB3	1.95	0.66
55:DA:2752:C:H5'	55:DA:2753:A:OP2	1.96	0.66
55:DA:2810:A:O3'	4:DE:61:ARG:HG2	1.96	0.66
2:DB:15:A:H5'	2:DB:16:G:H8	1.59	0.66
4:DE:120:TRP:CD2	4:DE:155:LYS:HD3	2.30	0.66
56:DI:21:LYS:CA	56:DI:24:ILE:HD12	2.25	0.66
56:DJ:10:GLU:O	56:DJ:17:VAL:CG1	2.43	0.66
58:DL:95:LYS:N	58:DL:136:VAL:CG1	2.58	0.66
11:DO:14:LYS:O	11:DO:15:ARG:C	2.33	0.66
14:DQ:49:VAL:HG22	14:DQ:80:LEU:HD12	1.78	0.66
57:DY:74:LEU:HB3	57:DY:120:LYS:H	1.58	0.66
17:A2:38:LEU:HD12	17:A2:55:ALA:C	2.17	0.66
2:AB:40:U:N1	26:A4:1:MET:HE1	2.11	0.66
1:AA:2046:G:H5'	27:A5:19:ARG:HG3	1.77	0.66
1:AA:2688:U:H2'	1:AA:2719:G:N2	2.10	0.66
1:AA:49:A:H1'	1:AA:51:G:C4	2.31	0.66
1:AA:896:A:C5'	1:AA:897:C:OP2	2.42	0.66
3:AD:186:HIS:HB3	3:AD:189:CYS:SG	2.36	0.66
12:AP:33:GLY:HA2	12:AP:105:GLU:HA	1.75	0.66
18:AS:4:LYS:HG2	18:AS:106:ILE:HG22	1.77	0.66
18:AS:58:ALA:O	18:AS:62:HIS:HB2	1.95	0.66
21:AV:176:PRO:O	21:AV:178:GLU:N	2.28	0.66
31:BA:1347:G:O2'	31:BA:1348:U:OP2	2.13	0.66
33:BF:35:GLU:O	33:BF:39:ILE:HG13	1.96	0.66
38:CK:103:VAL:HG21	38:CK:110:ALA:HB2	1.78	0.66
54:CA:1346:A:C5'	39:CL:120:ARG:HH12	2.08	0.66
46:CS:6:LEU:HG	46:CS:17:TYR:HB3	1.76	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:D4:14:ILE:HG21	26:D4:21:VAL:HB	1.78	0.66
30:D8:41:ILE:HG13	30:D8:42:ARG:N	2.10	0.66
55:DA:142:G:H2'	55:DA:143:C:C6	2.31	0.66
55:DA:829:A:N7	55:DA:2248:C:H5'	2.11	0.66
55:DA:2347:C:H2'	55:DA:2348:U:H6	1.59	0.66
55:DA:2409:G:H2'	55:DA:2410:G:O4'	1.96	0.66
55:DA:583:G:H5''	16:D1:10:ARG:HH12	1.59	0.66
55:DA:609(A):G:H2'	55:DA:610:C:H6	1.61	0.66
55:DA:800:A:H4'	55:DA:801:G:O5'	1.95	0.66
4:DE:14:ILE:O	4:DE:15:PHE:CG	2.49	0.66
4:DE:53:PRO:O	4:DE:74:PRO:HA	1.96	0.66
6:DG:133:LEU:HD21	6:DG:157:ILE:HB	1.79	0.66
58:DL:25:PRO:HA	58:DL:27:LEU:CG	2.21	0.66
20:DU:35:TYR:CE1	20:DU:69:ALA:HB3	2.31	0.66
21:DV:174:VAL:O	21:DV:175:VAL:CG2	2.44	0.66
1:AA:2818:G:OP2	13:A0:42:LYS:NZ	2.28	0.65
22:A3:68:GLU:CG	22:A3:80:HIS:HB2	2.25	0.65
28:A6:41:PRO:HD2	28:A6:46:HIS:HA	1.76	0.65
28:A6:52:VAL:HG13	28:A6:53:LYS:N	2.11	0.65
1:AA:581:C:H2'	1:AA:582:G:H8	1.60	0.65
1:AA:877:U:O2'	1:AA:878:A:H5'	1.95	0.65
9:AM:17:ASP:O	9:AM:18:ALA:HB2	1.95	0.65
11:AO:83:VAL:HG23	11:AO:105:LEU:HD22	1.79	0.65
2:AB:113:C:O2'	14:AQ:46:VAL:HG13	1.96	0.65
15:AR:102:ILE:O	15:AR:106:SER:HB3	1.96	0.65
31:BA:1241:G:H2'	31:BA:1242:C:H6	1.61	0.65
31:BA:792:A:C2'	31:BA:794:A:N6	2.42	0.65
31:BA:942:G:N2	39:BL:124:GLN:HE22	1.93	0.65
33:BF:118:GLN:O	33:BF:122:GLU:HG3	1.94	0.65
33:BF:34:LEU:HG	33:BF:38:ARG:NH2	2.10	0.65
31:BA:1189:C:OP1	33:BF:5:ILE:HG21	1.95	0.65
34:BG:53:ASP:O	34:BG:57:ARG:HG2	1.94	0.65
37:BJ:113:GLU:CB	37:BJ:119:ARG:HG2	2.25	0.65
38:BK:20:TYR:HE2	38:BK:75:ARG:HD2	1.61	0.65
40:BM:4:ILE:HD11	40:BM:77:PRO:HB3	1.78	0.65
47:BT:87:LYS:O	47:BT:91:ARG:HG3	1.96	0.65
53:C1:52:U:C2'	53:C1:53:U:C5'	2.73	0.65
54:CA:1176:A:N6	54:CA:1177:G:C6	2.65	0.65
54:CA:129(A):G:C2	54:CA:188:U:O2'	2.48	0.65
17:D2:58:VAL:CB	17:D2:98:GLU:HB2	2.19	0.65
26:D4:32:TYR:C	26:D4:32:TYR:CD2	2.70	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:D6:15:GLU:CD	28:D6:44:ARG:HH22	2.00	0.65
55:DA:1102:C:O2'	55:DA:1103:A:H5''	1.96	0.65
55:DA:1112:G:O2'	7:DH:2:SER:HB2	1.96	0.65
55:DA:1359:A:C4'	55:DA:1359:A:H8	2.04	0.65
55:DA:1464:C:HO2'	55:DA:1528:A:H8	1.42	0.65
55:DA:1652:A:H3'	55:DA:1653:G:C8	2.31	0.65
55:DA:1933:G:H2'	55:DA:1934:C:H5''	1.78	0.65
55:DA:2219:G:H2'	55:DA:2224:G:H5'	1.77	0.65
55:DA:2591:C:H2'	55:DA:2592:G:C8	2.30	0.65
4:DE:117:MET:O	4:DE:117:MET:CG	2.44	0.65
4:DE:51:PHE:CD1	4:DE:52:LEU:HG	2.30	0.65
5:DF:178:PRO:HG2	5:DF:179:GLU:OE2	1.94	0.65
8:DK:32:PRO:C	8:DK:34:GLY:H	1.98	0.65
58:DL:100:THR:C	58:DL:102:GLU:N	2.50	0.65
58:DL:78:ILE:CA	58:DL:82:ALA:HB3	2.25	0.65
11:DO:47:ASP:OD1	11:DO:50:ARG:NH2	2.30	0.65
11:DO:97:PRO:O	11:DO:98:GLU:HB3	1.97	0.65
20:DU:76:CYS:HB3	20:DU:96:ILE:CD1	2.24	0.65
21:DV:48:PHE:CE2	21:DV:52:SER:HA	2.31	0.65
57:DY:75:GLN:CG	57:DY:110:GLY:O	2.44	0.65
17:A2:57:VAL:HG12	17:A2:99:ILE:HG13	1.77	0.65
1:AA:1859:A:N6	1:AA:1883:G:O2'	2.30	0.65
1:AA:2534:A:H5'	1:AA:2534:A:H8	1.61	0.65
1:AA:2820:A:O5'	13:A0:4:LEU:HD22	1.97	0.65
1:AA:405:U:O2	1:AA:405:U:H2'	1.95	0.65
1:AA:620:G:H4'	1:AA:621:A:H5'	1.78	0.65
8:AK:97:ILE:O	8:AK:100:ALA:HB3	1.96	0.65
9:AM:137:LYS:HZ3	9:AM:138:LEU:HD23	1.60	0.65
23:AZ:82:LEU:HG	23:AZ:83:GLU:N	2.09	0.65
23:AZ:87:PRO:O	23:AZ:88:LYS:C	2.34	0.65
31:BA:828:A:H5''	31:BA:859:A:C2	2.31	0.65
31:BA:954:G:H2'	31:BA:955:U:H6	1.61	0.65
33:BF:125:GLU:HG3	33:BF:189:ALA:HB1	1.78	0.65
33:BF:76:VAL:HG23	33:BF:77:ILE:N	2.11	0.65
34:BG:49:ARG:NH2	53:B1:57:U:H1'	2.12	0.65
40:BM:78:ASN:C	40:BM:80:LYS:H	1.98	0.65
51:BX:15:ARG:HH11	51:BX:15:ARG:HB2	1.59	0.65
54:CA:73:G:C2	54:CA:74:C:N4	2.64	0.65
52:CC:28:G:H2'	52:CC:29:G:H5'	1.78	0.65
39:CL:17:VAL:HG11	39:CL:81:ILE:HA	1.77	0.65
28:D6:20:ASN:ND2	28:D6:42:TRP:HH2	1.94	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1728:G:N1	55:DA:1730:U:OP2	2.29	0.65
55:DA:2211:G:H2'	55:DA:2211:G:N3	2.11	0.65
55:DA:2725:A:O2'	55:DA:2726:U:O5'	2.13	0.65
55:DA:445:C:O2'	55:DA:446:G:H5'	1.95	0.65
55:DA:686:G:N2	55:DA:788:A:H61	1.94	0.65
3:DD:241:PRO:O	3:DD:243:GLY:N	2.29	0.65
4:DE:78:LEU:CD2	4:DE:79:ARG:HE	2.10	0.65
5:DF:47:GLY:HA3	5:DF:95:ARG:O	1.95	0.65
56:DI:16:THR:O	56:DI:20:LEU:HD12	1.95	0.65
56:DJ:12:LEU:N	56:DJ:13:SER:HB3	2.02	0.65
56:DJ:11:GLU:HA	56:DJ:17:VAL:HG11	1.78	0.65
8:DK:77:LEU:HD11	8:DK:140:LEU:CA	2.27	0.65
19:DT:70:LEU:H	19:DT:70:LEU:HD23	1.61	0.65
20:DU:95:LYS:HB3	20:DU:100:ALA:CA	2.19	0.65
21:DV:119:GLU:CG	21:DV:119:GLU:O	2.42	0.65
55:DA:111:A:H4'	24:DW:69:ARG:HH22	1.60	0.65
57:DY:89:ALA:HB2	57:DY:125:LEU:CD1	2.26	0.65
57:DY:91:LYS:HZ3	57:DY:95:GLN:HE21	1.44	0.65
1:AA:1280:G:C2'	1:AA:1281:G:H5''	2.26	0.65
1:AA:1331:A:H2'	1:AA:1333:C:C5	2.32	0.65
1:AA:212:G:O2'	1:AA:213:A:H5'	1.96	0.65
1:AA:242:G:O2'	1:AA:243:U:OP2	2.13	0.65
1:AA:307:G:H21	1:AA:330:A:H62	1.43	0.65
1:AA:527:C:OP2	1:AA:2779:U:C5	2.50	0.65
1:AA:654(L):G:N2	1:AA:654(M):C:H1'	2.12	0.65
2:AB:42:C:C4'	6:AG:67:LYS:HD3	2.20	0.65
10:AN:69:ILE:HD12	10:AN:69:ILE:H	1.60	0.65
11:AO:83:VAL:CG1	11:AO:112:LEU:HD21	2.26	0.65
31:BA:13:U:H5'	31:BA:14:U:OP2	1.96	0.65
31:BA:977:A:O2'	31:BA:978:A:H5'	1.96	0.65
39:BL:96:LEU:HD23	39:BL:102:LEU:HD12	1.79	0.65
49:BV:48:THR:HG22	49:BV:61:TYR:HA	1.77	0.65
49:BV:63:THR:HG22	49:BV:66:MET:HE3	1.78	0.65
53:C1:52:U:O2'	53:C1:53:U:C5'	2.39	0.65
54:CA:244:U:O2'	54:CA:245:C:OP2	2.14	0.65
35:CH:41:VAL:CG1	35:CH:113:ALA:HB2	2.26	0.65
39:CL:9:ARG:HB2	39:CL:14:VAL:HG22	1.79	0.65
46:CS:45:THR:CG2	46:CS:47:ASP:H	2.03	0.65
50:CW:63:ILE:HG22	50:CW:77:ALA:HB1	1.79	0.65
13:D0:74:LYS:O	13:D0:75:LEU:HB3	1.96	0.65
55:DA:1578:U:H2'	55:DA:1579:A:H5'	1.79	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1796:U:H2'	55:DA:1797:C:C6	2.31	0.65
55:DA:1999:C:H4'	55:DA:2723:C:O2	1.96	0.65
55:DA:395:U:O2'	55:DA:396:G:C8	2.48	0.65
55:DA:883:G:O5'	55:DA:883:G:C8	2.49	0.65
3:DD:64:ILE:O	3:DD:64:ILE:HG12	1.96	0.65
55:DA:2667:C:H1'	7:DH:109:PHE:HD2	1.61	0.65
58:DL:41:PHE:C	58:DL:41:PHE:CD2	2.69	0.65
12:DP:110:THR:HB	12:DP:112:GLU:HG2	1.78	0.65
14:DQ:58:LEU:HD23	14:DQ:58:LEU:H	1.61	0.65
14:DQ:66:ALA:O	14:DQ:69:VAL:HG13	1.96	0.65
18:DS:88:ARG:HB3	18:DS:92:ARG:HB3	1.77	0.65
27:A5:3:LYS:HG3	27:A5:4:HIS:H	1.61	0.65
1:AA:954:G:O2'	1:AA:2274:A:N1	2.24	0.65
1:AA:521:G:H2'	1:AA:522:G:H8	1.62	0.65
1:AA:581:C:H2'	1:AA:582:G:C8	2.31	0.65
1:AA:637:A:H4'	1:AA:638:G:O5'	1.96	0.65
2:AB:52:A:O2'	2:AB:53:A:N7	2.29	0.65
6:AG:94:LEU:H	6:AG:94:LEU:HD23	1.62	0.65
20:AU:17:SER:CB	20:AU:71:LYS:HD2	2.26	0.65
31:BA:1057:G:O2'	31:BA:1058:G:H5'	1.97	0.65
31:BA:1126:U:O2'	31:BA:1127:G:OP2	2.12	0.65
31:BA:160:A:H1'	31:BA:344:A:N7	2.11	0.65
31:BA:412:A:O2'	31:BA:413:G:OP2	2.15	0.65
33:BF:182:ILE:HG23	33:BF:202:ILE:C	2.17	0.65
41:BN:59:TYR:O	41:BN:62:GLN:HB3	1.97	0.65
43:BP:91:ARG:HH22	43:BP:103:THR:HG21	1.59	0.65
54:CA:1053:G:H5'	54:CA:1054:C:C5'	2.24	0.65
54:CA:396:G:O2'	54:CA:398:C:OP1	2.09	0.65
32:CE:87:ARG:NE	32:CE:233:SER:HB2	2.11	0.65
42:CO:6:THR:H	42:CO:9:GLN:NE2	1.94	0.65
55:DA:1060:U:C5	58:DL:74:ALA:CB	2.79	0.65
55:DA:1204:A:H1'	55:DA:1206:G:C4	2.31	0.65
55:DA:1489:U:HO2'	55:DA:1490:A:H8	1.45	0.65
55:DA:1655:A:H3'	55:DA:1656:C:H6	1.60	0.65
55:DA:276:A:H2'	55:DA:277:C:C6	2.31	0.65
4:DE:186:GLY:O	4:DE:188:VAL:N	2.30	0.65
6:DG:55:LYS:NZ	6:DG:148:MET:HG3	2.12	0.65
58:DL:60:TYR:CD2	58:DL:63:ARG:HB3	2.31	0.65
58:DL:8:VAL:H	58:DL:57:ILE:CG1	2.09	0.65
12:DP:134:ARG:O	12:DP:135:ASP:O	2.13	0.65
15:DR:39:ARG:CG	15:DR:40:THR:H	2.09	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:DR:41:ARG:HG2	15:DR:41:ARG:HH11	1.61	0.65
57:DY:55:LYS:HG2	57:DY:56:ASN:N	2.10	0.65
28:A6:25:LYS:HA	30:A8:34:TRP:CH2	2.32	0.65
1:AA:1091:G:H2'	1:AA:1092:C:C6	2.32	0.65
1:AA:573:G:N1	1:AA:2031:A:OP2	2.25	0.65
1:AA:2602:A:O2'	1:AA:2603:G:OP2	2.13	0.65
1:AA:2758:A:H2'	1:AA:2759:G:C5'	2.23	0.65
1:AA:2789:C:C2'	1:AA:2790:A:H5''	2.26	0.65
1:AA:2892:A:H2'	1:AA:2893:G:O4'	1.95	0.65
1:AA:527:C:OP2	1:AA:2779:U:H5	1.79	0.65
1:AA:590:A:H2'	1:AA:591:C:C6	2.31	0.65
5:AF:53:THR:HG23	5:AF:55:GLY:H	1.61	0.65
7:AH:7:LEU:N	7:AH:8:PRO:CD	2.60	0.65
14:AQ:26:LEU:HD22	14:AQ:87:PHE:CD1	2.31	0.65
15:AR:98:LYS:HE3	15:AR:98:LYS:HA	1.76	0.65
21:AV:177:PRO:O	21:AV:180:VAL:N	2.30	0.65
31:BA:1293:G:O2'	31:BA:1294:G:H5'	1.97	0.65
31:BA:149:A:H2'	31:BA:150:C:C6	2.32	0.65
31:BA:411:A:N7	31:BA:413:G:N3	2.44	0.65
31:BA:947:G:H2'	31:BA:948:C:C6	2.31	0.65
31:BA:1190:G:OP1	33:BF:4:LYS:HA	1.95	0.65
36:BI:7:ASN:C	36:BI:8:ILE:HD12	2.17	0.65
54:CA:1364:U:H2'	54:CA:1364:U:O2	1.96	0.65
54:CA:186(A):C:H2'	54:CA:186(B):C:H6	1.62	0.65
54:CA:826:C:H2'	54:CA:827:U:O2	1.96	0.65
32:CE:35:GLU:O	32:CE:36:ARG:HD3	1.97	0.65
34:CG:112:VAL:HG13	34:CG:113:SER:H	1.61	0.65
39:CL:118:LYS:O	39:CL:119:ALA:HB3	1.97	0.65
43:CP:88:ARG:NH1	43:CP:88:ARG:HB3	2.04	0.65
48:CU:19:LYS:O	48:CU:20:ALA:HB2	1.95	0.65
54:CA:108:G:H1	50:CW:15:ARG:HH21	1.43	0.65
16:D1:87:GLY:O	17:D2:50:PRO:HD3	1.96	0.65
55:DA:140:A:H8	55:DA:1408:C:HO2'	1.44	0.65
55:DA:1479:G:H5'	55:DA:1558:A:H2	1.60	0.65
55:DA:1735:C:H6	55:DA:1735:C:H5'	1.62	0.65
55:DA:265:A:H2'	55:DA:266:G:O4'	1.96	0.65
2:DB:80:U:O2'	2:DB:81:G:H5''	1.97	0.65
56:DJ:14:GLN:N	56:DJ:15:ALA:O	2.30	0.65
8:DK:69:LYS:HG3	8:DK:136:VAL:HB	1.78	0.65
58:DL:101:TRP:NE1	58:DL:140:GLY:HA2	2.11	0.65
15:DR:23:ARG:HA	15:DR:52:ILE:HD11	1.79	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:DV:128:VAL:HG22	21:DV:129:SER:N	2.12	0.65
21:DV:16:SER:O	21:DV:20:ARG:HG3	1.97	0.65
57:DY:73:GLY:N	57:DY:112:LEU:HG	2.10	0.65
1:AA:2820:A:C1'	13:A0:3:HIS:HB3	2.27	0.65
17:A2:69:LYS:HG3	17:A2:86:GLY:HA3	1.78	0.65
1:AA:177:G:H5''	1:AA:178:G:OP2	1.97	0.65
1:AA:2348:U:H2'	1:AA:2349:G:C5'	2.27	0.65
1:AA:2657:A:C2	1:AA:2665:A:N7	2.64	0.65
21:AV:145:GLU:O	21:AV:146:ILE:CG1	2.44	0.65
31:BA:1296:C:H3'	31:BA:1297:C:C6	2.32	0.65
31:BA:1541:U:O2	31:BA:1541:U:H2'	1.96	0.65
32:BE:92:TYR:CE2	32:BE:151:GLY:HA3	2.31	0.65
33:BF:138:VAL:HG22	33:BF:151:VAL:HG23	1.78	0.65
42:BO:117:ARG:HH21	42:BO:124:LYS:HA	1.62	0.65
54:CA:959:A:H2	54:CA:1221:G:N3	1.95	0.65
54:CA:399:G:H2'	54:CA:400:C:C6	2.32	0.65
54:CA:890:G:O2'	54:CA:891:U:P	2.55	0.65
52:CD:9:A:H62	52:CD:23:A:N6	1.90	0.65
17:D2:41:GLY:HA3	17:D2:46:VAL:HG11	1.79	0.65
55:DA:1075:C:C5'	21:DV:195:GLU:CD	2.65	0.65
55:DA:1278:A:O3'	13:D0:34:ILE:HG23	1.97	0.65
55:DA:1478:G:O2'	55:DA:1479:G:H5'	1.96	0.65
55:DA:1791:A:N6	55:DA:1828:G:O2'	2.30	0.65
55:DA:2507:C:H5'	55:DA:2507:C:H6	1.62	0.65
7:DH:109:PHE:C	7:DH:111:HIS:H	1.99	0.65
56:DI:11:GLU:HA	56:DI:14:GLN:CD	2.17	0.65
9:DM:26:LEU:HG	9:DM:30:ILE:HD11	1.79	0.65
20:DU:95:LYS:HE3	20:DU:95:LYS:O	1.96	0.65
57:DY:101:PRO:O	57:DY:102:LYS:CB	2.43	0.65
57:DY:128:LEU:HD22	57:DY:129:PRO:HB3	1.77	0.65
16:A1:66:ASN:HD21	16:A1:70:ARG:HE	1.45	0.65
29:A7:30:VAL:HA	29:A7:33:ARG:NH1	2.12	0.65
1:AA:1022:G:N2	1:AA:1142(A):A:C2	2.63	0.65
1:AA:1086:A:C4'	1:AA:1103:A:H61	2.09	0.65
1:AA:1225:C:H5''	17:A2:85:LYS:HE2	1.77	0.65
1:AA:2150:U:H2'	1:AA:2151:G:H8	1.61	0.65
2:AB:55:U:H1'	6:AG:29:TRP:HE1	1.61	0.65
3:AD:108:PRO:HD2	3:AD:111:LEU:HG	1.78	0.65
4:AE:200:GLU:CG	4:AE:201:THR:H	2.09	0.65
4:AE:66:HIS:CE1	4:AE:73:GLU:HB2	2.28	0.65
7:AH:4:ILE:HB	7:AH:6:ARG:NH1	2.11	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:AH:97:ARG:HG2	7:AH:98:LEU:N	2.12	0.65
12:AP:140:ALA:O	12:AP:141:GLN:CB	2.45	0.65
19:AT:46:ALA:O	19:AT:48:LYS:HE2	1.97	0.65
20:AU:86:ARG:HB3	20:AU:88:LYS:NZ	2.12	0.65
21:AV:133:ILE:N	21:AV:133:ILE:HD12	2.11	0.65
31:BA:129(A):G:O2'	31:BA:189:U:H5''	1.96	0.65
32:BE:132:LYS:HA	32:BE:135:GLN:HE21	1.61	0.65
35:BH:20:GLN:NE2	35:BH:21:ALA:N	2.44	0.65
35:BH:40:ARG:NH2	35:BH:66:MET:HG2	2.11	0.65
49:BV:41:VAL:H	49:BV:44:MET:CE	2.07	0.65
50:BW:12:ALA:H	50:BW:13:LEU:HD12	1.60	0.65
54:CA:243:A:H4'	54:CA:244:U:C5'	2.26	0.65
54:CA:518:C:H2'	54:CA:518:C:O2	1.97	0.65
43:CP:88:ARG:CB	43:CP:88:ARG:HH11	2.01	0.65
46:CS:14:ASN:N	46:CS:15:PRO:HD3	2.12	0.65
46:CS:40:ASP:OD2	46:CS:42:ARG:HB2	1.97	0.65
50:CW:56:MET:HG3	50:CW:88:VAL:HG21	1.77	0.65
4:DE:111:ARG:HG2	13:D0:1:MET:SD	2.36	0.65
26:D4:67:TYR:O	26:D4:68:ARG:NH2	2.30	0.65
28:D6:17:LYS:HG3	28:D6:18:ARG:N	2.11	0.65
29:D7:9:ARG:HH21	29:D7:48:LYS:HB2	1.61	0.65
55:DA:1056:G:OP1	57:DY:35:LYS:CD	2.44	0.65
55:DA:1047:G:H2'	55:DA:1110:G:C2	2.31	0.65
55:DA:1652:A:H2'	55:DA:1653:G:O4'	1.97	0.65
55:DA:2275:C:O2'	12:DP:83:MET:HG3	1.97	0.65
4:DE:7:VAL:CG2	4:DE:8:LYS:H	1.99	0.65
5:DF:57:VAL:HG12	5:DF:59:TYR:H	1.61	0.65
8:DK:110:ASP:HB2	8:DK:112:LYS:N	2.12	0.65
15:DR:61:PHE:CE2	15:DR:76:PHE:HB2	2.32	0.65
20:DU:50:ARG:HB3	20:DU:53:PRO:CG	2.27	0.65
57:DY:104:ILE:HG13	57:DY:105:PRO:CD	2.09	0.65
1:AA:2168:G:H2'	1:AA:2168:G:N3	2.11	0.65
1:AA:2807:G:C3'	1:AA:2808:U:H5''	2.26	0.65
1:AA:946:G:HO2'	1:AA:947:G:H5'	1.54	0.65
3:AD:35:LYS:HD2	3:AD:104:TYR:CD1	2.31	0.65
3:AD:166:GLN:CA	3:AD:166:GLN:HE21	2.08	0.65
4:AE:24:THR:HG21	4:AE:188:VAL:HG12	1.79	0.65
5:AF:192:LEU:HD21	5:AF:194:MET:HE3	1.79	0.65
15:AR:45:PHE:CE2	15:AR:74:ARG:HB2	2.32	0.65
19:AT:26:TYR:OH	19:AT:88:LYS:HB2	1.97	0.65
20:AU:94:LYS:NZ	20:AU:101:LYS:HZ3	1.94	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:690:G:H2'	31:BA:691:G:O4'	1.97	0.65
34:BG:165:MET:HE2	34:BG:168:ARG:HB2	1.78	0.65
34:BG:12:CYS:SG	34:BG:21:LEU:HD22	2.36	0.65
37:BJ:18:TYR:HD2	37:BJ:59:LEU:HD22	1.61	0.65
45:BR:29:VAL:HG13	45:BR:63:ARG:HG3	1.78	0.65
47:BT:68:ARG:O	47:BT:69:LYS:HB2	1.97	0.65
54:CA:1032:A:H3'	54:CA:1032(A):G:C5'	2.26	0.65
54:CA:1442:G:H1	54:CA:1461:G:H21	1.43	0.65
54:CA:418:C:H2'	54:CA:419:C:C6	2.32	0.65
54:CA:126:G:H5'	54:CA:633:G:N2	2.11	0.65
54:CA:765:G:H1	54:CA:812:C:H2'	1.59	0.65
52:CD:64:A:C2	52:CD:65:G:H1'	2.31	0.65
32:CE:163:PHE:HA	32:CE:185:ILE:HG13	1.78	0.65
32:CE:51:LEU:HD23	32:CE:201:ILE:HD12	1.79	0.65
36:CI:46:ARG:HB3	36:CI:60:PHE:CE1	2.32	0.65
40:CM:39:PRO:HB3	40:CM:70:ARG:NH1	2.12	0.65
40:CM:54:PHE:CE2	40:CM:55:LYS:HD2	2.32	0.65
44:CQ:13:THR:N	44:CQ:14:PRO:CD	2.60	0.65
49:CV:87:ALA:O	49:CV:88:LYS:CD	2.40	0.65
50:CW:86:ARG:HG3	50:CW:86:ARG:HH11	1.61	0.65
55:DA:2723:C:H4'	13:D0:1:MET:HG2	1.79	0.65
55:DA:583:G:OP2	16:D1:10:ARG:NH1	2.27	0.65
29:D7:8:ASN:ND2	29:D7:8:ASN:C	2.46	0.65
55:DA:1844:C:O2'	55:DA:1845:G:H5'	1.96	0.65
55:DA:1918:A:O2'	55:DA:1919:A:N7	2.29	0.65
55:DA:1991:U:H2'	55:DA:1992:G:H5''	1.79	0.65
55:DA:2271:G:H5''	22:D3:20:ARG:NE	2.12	0.65
55:DA:2492:U:O2'	55:DA:2493:U:H5'	1.97	0.65
55:DA:72:U:C4	55:DA:112:U:H4'	2.32	0.65
3:DD:35:LYS:HD2	3:DD:104:TYR:CE1	2.31	0.65
55:DA:674:G:H1'	5:DF:74:ARG:HD3	1.78	0.65
58:DL:58:THR:OG1	58:DL:66:THR:HG23	1.96	0.65
55:DA:1070:A:N1	58:DL:9:LYS:HE3	2.12	0.65
21:DV:181:GLU:O	21:DV:181:GLU:CG	2.44	0.65
21:DV:51:ALA:HA	21:DV:55:HIS:CD2	2.31	0.65
24:DW:41:ILE:HD12	24:DW:41:ILE:O	1.97	0.65
57:DY:93:LEU:HD11	57:DY:97:ALA:O	1.97	0.65
23:DZ:78:LYS:HD2	23:DZ:80:LEU:HD12	1.79	0.65
28:A6:14:THR:O	28:A6:49:HIS:HA	1.97	0.65
28:A6:10:LEU:HA	30:A8:34:TRP:CZ3	2.32	0.65
1:AA:1140:C:C1'	1:AA:1143:A:N7	2.60	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:270(G):C:H2'	1:AA:270(H):C:C6	2.31	0.65
4:AE:171:GLU:HG2	4:AE:185:LYS:HG2	1.79	0.65
5:AF:153:SER:HB2	5:AF:190:GLU:H	1.62	0.65
5:AF:37:VAL:HA	5:AF:40:GLN:HG3	1.79	0.65
8:AK:88:ILE:CG2	8:AK:89:TYR:H	2.02	0.65
11:AO:3:LEU:HA	11:AO:6:LEU:HD23	1.78	0.65
15:AR:132:LYS:CG	15:AR:136:GLN:HE22	2.08	0.65
31:BA:1288:A:H1'	31:BA:1352:C:O2'	1.96	0.65
31:BA:388:G:O2'	31:BA:389:A:P	2.55	0.65
31:BA:545:C:H5''	34:BG:72:GLU:HG2	1.79	0.65
31:BA:908:A:H2'	31:BA:909:A:C8	2.31	0.65
52:BC:58:A:H1'	52:BC:60:U:H5	1.62	0.65
32:BE:78:GLN:CA	32:BE:94:ASN:HD21	2.10	0.65
34:BG:170:VAL:HG13	34:BG:174:LEU:O	1.97	0.65
36:BI:87:ARG:HG2	36:BI:87:ARG:HH11	1.62	0.65
38:BK:29:SER:HB3	38:BK:32:LYS:CG	2.27	0.65
42:BO:117:ARG:HG2	42:BO:117:ARG:HH11	1.61	0.65
54:CA:865:A:H5'	54:CA:1078:U:O4	1.97	0.65
54:CA:1116:C:C2'	54:CA:1117:G:H5''	2.27	0.65
54:CA:1225:A:H2'	54:CA:1225:A:N3	2.10	0.65
54:CA:1305:G:O2'	54:CA:1306:A:H8	1.78	0.65
33:CF:150:LYS:HG3	33:CF:169:ALA:HB2	1.79	0.65
38:CK:41:ARG:NH1	38:CK:41:ARG:HG3	2.08	0.65
42:CO:45:PRO:HB3	42:CO:92:ASP:HB3	1.77	0.65
54:CA:956:U:OP1	49:CV:87:ALA:HA	1.97	0.65
17:D2:1:MET:HG3	17:D2:43:GLU:HG2	1.77	0.65
26:D4:68:ARG:NH1	26:D4:68:ARG:CB	2.60	0.65
11:DO:61:ARG:HH12	30:D8:13:ARG:HG3	1.61	0.65
55:DA:1429:G:H2'	55:DA:1430:C:C6	2.32	0.65
55:DA:1747:G:O2'	55:DA:1748:G:H5'	1.97	0.65
55:DA:1799:G:N2	55:DA:1818:U:O2'	2.30	0.65
55:DA:33:U:H4'	55:DA:34:C:OP1	1.97	0.65
55:DA:443:A:H1'	55:DA:1201:C:O4'	1.96	0.65
55:DA:790:C:H4'	55:DA:791:C:OP1	1.97	0.65
6:DG:135:LEU:HD23	6:DG:140:ILE:HD11	1.79	0.65
6:DG:16:ARG:HB3	6:DG:17:PRO:CD	2.26	0.65
58:DL:108:ALA:HA	58:DL:111:LYS:NZ	2.12	0.65
58:DL:38:VAL:HG12	58:DL:42:ASN:OD1	1.97	0.65
58:DL:50:ASP:CG	58:DL:51:ALA:N	2.49	0.65
9:DM:35:ARG:HD3	9:DM:37:LYS:CD	2.27	0.65
11:DO:105:LEU:H	11:DO:105:LEU:HD12	1.62	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:DP:34:LEU:HD23	12:DP:104:PHE:HD1	1.62	0.65
57:DY:1:MET:CE	57:DY:3:ASN:ND2	2.60	0.65
57:DY:51:LEU:CD1	57:DY:82:PHE:H	1.88	0.65
1:AA:1210:A:H4'	1:AA:1211:U:O5'	1.96	0.65
1:AA:2258:C:H4'	1:AA:2259:G:OP2	1.94	0.65
1:AA:2304:G:H21	6:AG:156:ASP:CG	1.99	0.65
1:AA:278:A:H2'	1:AA:279:C:C6	2.32	0.65
7:AH:144:VAL:O	7:AH:148:ILE:HG12	1.96	0.65
8:AK:56:LYS:HE3	8:AK:60:GLU:HG2	1.79	0.65
14:AQ:42:ASP:O	14:AQ:43:GLU:HB2	1.97	0.65
31:BA:1152:A:H2'	31:BA:1153:C:H6	1.62	0.65
31:BA:1262:C:H2'	31:BA:1263:C:H6	1.62	0.65
31:BA:1326:C:OP2	51:BX:6:ARG:HD3	1.96	0.65
31:BA:1358:U:OP1	44:BQ:35:ARG:HG2	1.97	0.65
31:BA:1486:G:H2'	31:BA:1487:G:O4'	1.96	0.65
52:BD:46:G:N2	52:BD:48:C:O2	2.29	0.65
32:BE:7:VAL:HG22	32:BE:8:LYS:H	1.61	0.65
37:BJ:94:ARG:O	37:BJ:97:GLN:HB3	1.97	0.65
42:BO:27:LEU:HD23	42:BO:27:LEU:N	2.12	0.65
54:CA:1004:A:H5''	54:CA:1025:U:O4	1.97	0.65
54:CA:17:U:H2'	54:CA:18:C:H6	1.61	0.65
54:CA:198:G:H2'	54:CA:199:G:C8	2.32	0.65
54:CA:630:G:HO2'	54:CA:631:G:P	2.17	0.65
54:CA:784:C:H4'	55:DA:1837:C:OP1	1.96	0.65
54:CA:788:U:H2'	54:CA:789:U:H5'	1.79	0.65
54:CA:792:A:O2'	54:CA:793:U:OP2	2.15	0.65
54:CA:998:G:O2'	54:CA:998(A):C:H5'	1.97	0.65
34:CG:109:GLY:HA3	34:CG:165:MET:SD	2.37	0.65
38:CK:103:VAL:HG23	38:CK:110:ALA:HB2	1.78	0.65
41:CN:59:TYR:O	41:CN:62:GLN:HB3	1.97	0.65
47:CT:40:LYS:HG2	47:CT:41:LYS:N	2.12	0.65
17:D2:35:LEU:O	17:D2:37:VAL:N	2.30	0.65
26:D4:16:CYS:C	26:D4:18:CYS:N	2.49	0.65
55:DA:1444(A):A:H4'	55:DA:1460:A:O2'	1.95	0.65
55:DA:1496:A:H8	55:DA:1577:C:O2'	1.80	0.65
55:DA:1655:A:H3'	55:DA:1656:C:C6	2.32	0.65
55:DA:2317:C:H2'	55:DA:2318:G:C5'	2.26	0.65
3:DD:206:LEU:HB3	3:DD:211:ARG:HB3	1.79	0.65
3:DD:95:LEU:O	3:DD:95:LEU:HD12	1.97	0.65
6:DG:116:ASP:O	6:DG:117:PHE:HB3	1.96	0.65
8:DK:4:ILE:HG12	8:DK:18:VAL:CG2	2.25	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:DK:77:LEU:HD13	8:DK:78:THR:H	1.62	0.65
58:DL:52:ILE:CG1	58:DL:76:TYR:HB2	2.23	0.65
58:DL:78:ILE:O	58:DL:82:ALA:HB3	1.97	0.65
9:DM:45:ASN:HD22	9:DM:45:ASN:N	1.93	0.65
57:DY:73:GLY:O	57:DY:74:LEU:HB3	1.97	0.65
23:DZ:23:LYS:HE3	23:DZ:29:GLY:N	2.11	0.65
23:DZ:82:LEU:HD22	23:DZ:82:LEU:H	1.62	0.65
23:DZ:86:SER:N	23:DZ:87:PRO:CD	2.59	0.65
16:A1:8:VAL:HG12	16:A1:11:ARG:NH2	2.13	0.64
1:AA:1225:C:H5''	17:A2:85:LYS:CE	2.27	0.64
26:A4:61:ARG:HE	26:A4:61:ARG:HA	1.62	0.64
1:AA:1215:G:O2'	1:AA:1216:G:H5'	1.97	0.64
1:AA:1212:G:H2'	1:AA:1236:G:H22	1.61	0.64
1:AA:2425:A:H5'	1:AA:2427:C:O4'	1.98	0.64
1:AA:197:A:N6	1:AA:2430:A:H2'	2.12	0.64
1:AA:1050:A:H1'	1:AA:2751:G:N2	2.12	0.64
1:AA:757:U:H2'	1:AA:758:C:H6	1.62	0.64
1:AA:874:G:H2'	1:AA:875:G:H8	1.63	0.64
4:AE:8:LYS:CB	4:AE:192:ASN:HA	2.27	0.64
7:AH:152:ARG:C	7:AH:154:PRO:HD3	2.17	0.64
11:AO:146:VAL:HG13	11:AO:147:LEU:HD13	1.80	0.64
19:AT:64:LYS:HD3	19:AT:73:ARG:NE	2.12	0.64
31:BA:197:A:N1	31:BA:221:C:H4'	2.11	0.64
31:BA:366:C:O2'	31:BA:367:U:O5'	2.14	0.64
31:BA:448:A:C2	31:BA:449:C:N3	2.64	0.64
31:BA:532:A:O2'	31:BA:533:A:OP1	2.13	0.64
32:BE:179:LYS:NZ	32:BE:179:LYS:HB2	2.13	0.64
54:CA:651:C:H2'	54:CA:652:U:C6	2.32	0.64
52:CB:19:G:O2'	52:CB:20:U:P	2.55	0.64
33:CF:77:ILE:C	33:CF:83:ARG:HB3	2.17	0.64
38:CK:116:LYS:HE2	38:CK:116:LYS:N	2.11	0.64
39:CL:5:TYR:CD2	39:CL:6:GLY:N	2.65	0.64
54:CA:963:G:N2	40:CM:55:LYS:HD3	2.11	0.64
54:CA:690:G:N2	41:CN:55:LYS:NZ	2.40	0.64
48:CU:53:ARG:HG2	48:CU:58:LEU:O	1.96	0.64
49:CV:7:LYS:HB3	49:CV:7:LYS:HZ3	1.61	0.64
26:D4:49:PHE:O	26:D4:50:VAL:HB	1.97	0.64
6:DG:66:GLN:HA	26:D4:6:HIS:CE1	2.32	0.64
55:DA:1079:C:H1'	58:DL:129:GLY:O	1.96	0.64
55:DA:1165:U:H2'	55:DA:1166:C:H6	1.58	0.64
55:DA:2562:U:H4'	10:DN:25:LEU:HD21	1.79	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:443:A:N7	5:DF:45:ARG:HD2	2.13	0.64
55:DA:811:U:O2'	55:DA:1250:G:H2'	1.97	0.64
55:DA:897:C:C5	55:DA:897:C:OP2	2.49	0.64
4:DE:176:ILE:HD12	4:DE:176:ILE:N	2.12	0.64
4:DE:35:GLN:CG	4:DE:37:ARG:HG2	2.27	0.64
5:DF:185:ASP:HA	5:DF:188:ARG:HD3	1.78	0.64
56:DJ:10:GLU:C	56:DJ:17:VAL:CG1	2.65	0.64
8:DK:114:LEU:HD22	8:DK:130:TYR:CD1	2.31	0.64
8:DK:57:ARG:NH1	8:DK:57:ARG:HB2	2.12	0.64
58:DL:102:GLU:CG	58:DL:103:GLN:N	2.60	0.64
58:DL:112:MET:H	58:DL:113:PRO:HD2	1.58	0.64
58:DL:78:ILE:HG12	58:DL:131:ALA:HB2	1.79	0.64
5:DF:34:TRP:CE2	11:DO:8:PRO:HG3	2.32	0.64
55:DA:1083:U:O4'	57:DY:41:ARG:HD3	1.97	0.64
22:A3:49:LYS:HB2	22:A3:80:HIS:HB3	1.79	0.64
28:A6:17:LYS:HA	28:A6:17:LYS:HE3	1.79	0.64
1:AA:1266:G:O6	18:AS:13:SER:OG	2.06	0.64
1:AA:1967:C:C2'	1:AA:1968:G:H5'	2.27	0.64
1:AA:2335:A:O2'	1:AA:2336:A:H3'	1.97	0.64
1:AA:654(K):C:H2'	1:AA:654(L):G:C8	2.33	0.64
3:AD:34:VAL:O	3:AD:34:VAL:HG13	1.95	0.64
1:AA:1110:G:H4'	7:AH:3:ARG:HH22	1.61	0.64
11:AO:14:LYS:O	11:AO:15:ARG:C	2.35	0.64
20:AU:4:LYS:HE2	20:AU:4:LYS:HA	1.77	0.64
21:AV:125:LEU:HG	21:AV:164:ALA:HB1	1.77	0.64
21:AV:148:ASP:O	21:AV:149:SER:HB3	1.97	0.64
21:AV:67:LEU:CD2	21:AV:90:VAL:HG13	2.28	0.64
21:AV:95:PRO:O	21:AV:96:VAL:CB	2.45	0.64
25:AX:8:LEU:HB2	25:AX:28:LEU:HD13	1.79	0.64
31:BA:1000:A:H2'	31:BA:1001:G:H5'	1.79	0.64
31:BA:1132:C:H2'	31:BA:1133:G:H8	1.61	0.64
31:BA:1498:U:O2'	31:BA:1499:A:OP2	2.12	0.64
31:BA:466:C:H5"	31:BA:467:G:OP2	1.97	0.64
31:BA:676:A:H2'	31:BA:677:U:H6	1.62	0.64
39:BL:4:TYR:HA	39:BL:88:TYR:CE1	2.32	0.64
54:CA:658:G:H2'	54:CA:659:U:C6	2.31	0.64
32:CE:12:GLU:C	32:CE:14:GLY:H	2.00	0.64
33:CF:113:ALA:HB3	33:CF:114:PRO:HD3	1.78	0.64
26:D4:69:LYS:HD3	26:D4:70:GLY:CA	2.27	0.64
28:D6:33:LYS:O	28:D6:35:GLU:N	2.27	0.64
55:DA:2724:C:OP1	13:D0:1:MET:HE3	1.98	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2897:U:O2'	55:DA:2898:U:H5'	1.96	0.64
4:DE:116:VAL:HG22	4:DE:122:PHE:CG	2.32	0.64
5:DF:123:LEU:HD12	5:DF:192:LEU:O	1.97	0.64
5:DF:20:LEU:HD12	5:DF:21:ALA:N	2.12	0.64
11:DO:9:ASN:HB2	11:DO:10:PRO:HD2	1.77	0.64
2:DB:52:A:N6	14:DQ:33:LYS:HG3	2.13	0.64
21:DV:191:VAL:HG21	21:DV:197:ILE:CG1	2.26	0.64
57:DY:51:LEU:CG	57:DY:82:PHE:C	2.65	0.64
13:A0:63:ARG:HB2	13:A0:63:ARG:NH1	2.11	0.64
1:AA:1694:C:H1'	1:AA:1695:G:C2	2.32	0.64
1:AA:270(F):U:H2'	1:AA:270(G):C:C6	2.32	0.64
1:AA:2685:G:H1'	1:AA:2726:U:H5	1.62	0.64
1:AA:33:U:C4'	1:AA:34:C:OP1	2.43	0.64
1:AA:407:G:H2'	1:AA:408:G:C8	2.33	0.64
1:AA:662:G:OP1	11:AO:15:ARG:NE	2.31	0.64
21:AV:129:SER:O	21:AV:133:ILE:HD11	1.98	0.64
24:AW:17:SER:HA	24:AW:20:GLU:HG3	1.80	0.64
24:AW:30:ARG:HH11	24:AW:30:ARG:HG3	1.61	0.64
53:B1:41:U:H5'	53:B1:42:U:OP1	1.97	0.64
31:BA:1228:C:OP1	43:BP:115:LYS:HD2	1.96	0.64
31:BA:566:G:H4'	31:BA:567:G:OP1	1.97	0.64
52:BB:74:C:O2'	52:BB:75:C:P	2.56	0.64
32:BE:213:LEU:O	32:BE:216:SER:HB3	1.98	0.64
48:BU:31:LEU:H	48:BU:31:LEU:CD2	2.10	0.64
49:BV:40:ILE:HG22	49:BV:67:VAL:O	1.98	0.64
54:CA:1152:A:H2'	54:CA:1153:C:C6	2.32	0.64
54:CA:1305:G:O2'	54:CA:1306:A:C8	2.50	0.64
54:CA:653:A:H1'	38:CK:56:LYS:HD3	1.79	0.64
33:CF:15:THR:CG2	33:CF:181:ASN:HA	2.27	0.64
33:CF:64:VAL:HG12	33:CF:66:VAL:HG23	1.79	0.64
54:CA:27:G:H4'	34:CG:209:ARG:HG3	1.79	0.64
39:CL:59:PHE:CZ	39:CL:88:TYR:HE1	2.14	0.64
54:CA:1151:A:N3	40:CM:39:PRO:HG3	2.12	0.64
46:CS:76:GLN:O	46:CS:76:GLN:HG2	1.96	0.64
46:CS:74:LEU:O	46:CS:79:VAL:HG23	1.96	0.64
48:CU:70:ILE:O	48:CU:74:ARG:HG3	1.97	0.64
51:CX:12:LYS:HB3	51:CX:22:ARG:HD2	1.79	0.64
9:DM:42:TRP:O	16:D1:64:ARG:NH2	2.30	0.64
22:D3:56:ASP:O	22:D3:57:PHE:HB2	1.98	0.64
28:D6:41:PRO:HD2	28:D6:46:HIS:CA	2.27	0.64
55:DA:1880:C:H6	55:DA:1880:C:H5'	1.62	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2011:U:OP1	18:DS:42:ARG:NH1	2.29	0.64
55:DA:2238:G:H5'	55:DA:2239:G:OP1	1.98	0.64
55:DA:307:G:N2	55:DA:310:A:OP2	2.29	0.64
6:DG:109:VAL:CG1	26:D4:33:VAL:HG21	2.26	0.64
7:DH:30:LYS:HE3	7:DH:81:GLU:HG3	1.79	0.64
58:DL:86:LYS:CE	58:DL:86:LYS:CA	2.67	0.64
19:DT:41:ASN:O	19:DT:45:THR:HG23	1.98	0.64
25:DX:31:LEU:O	25:DX:32:GLN:HB2	1.95	0.64
16:A1:100:VAL:O	16:A1:101:ARG:HG2	1.97	0.64
28:A6:11:LEU:O	28:A6:24:GLU:O	2.15	0.64
1:AA:1180:C:H6	1:AA:1180:C:H5'	1.62	0.64
1:AA:242:G:O2'	1:AA:243:U:P	2.55	0.64
1:AA:299:A:N1	1:AA:322:A:O2'	2.20	0.64
1:AA:897:C:OP2	1:AA:897:C:C6	2.50	0.64
2:AB:0:A:H2'	2:AB:1:U:C6	2.32	0.64
4:AE:58:ARG:NE	4:AE:58:ARG:HA	2.09	0.64
7:AH:125:VAL:HG22	7:AH:126:PRO:CA	2.27	0.64
7:AH:92:ILE:HD12	7:AH:92:ILE:N	2.11	0.64
9:AM:45:ASN:HD22	9:AM:45:ASN:N	1.95	0.64
9:AM:55:VAL:HG23	9:AM:56:ASN:OD1	1.97	0.64
15:AR:27:THR:HG23	15:AR:90:GLN:HB3	1.79	0.64
21:AV:148:ASP:OD1	21:AV:174:VAL:N	2.30	0.64
31:BA:1226:C:C4'	31:BA:1227:A:OP1	2.40	0.64
31:BA:243:A:H4'	31:BA:244:U:O5'	1.97	0.64
52:BB:74:C:O2'	52:BB:75:C:OP2	2.16	0.64
32:BE:7:VAL:HG13	32:BE:8:LYS:H	1.63	0.64
26:A4:52:THR:HG21	43:BP:65:LYS:CD	2.27	0.64
31:BA:667:G:H4'	45:BR:51:HIS:CE1	2.32	0.64
49:BV:40:ILE:HD11	49:BV:62:ILE:HD13	1.79	0.64
54:CA:1285:A:O2'	54:CA:1286:A:OP2	2.14	0.64
54:CA:297:G:N2	54:CA:299:G:H3'	2.13	0.64
54:CA:686:U:H2'	54:CA:687:A:H8	1.59	0.64
54:CA:812:C:O2'	54:CA:813:U:P	2.56	0.64
32:CE:7:VAL:HG11	32:CE:217:ARG:NH2	2.13	0.64
36:CI:37:VAL:HG12	36:CI:38:GLU:N	2.12	0.64
39:CL:77:ILE:O	39:CL:81:ILE:HG12	1.98	0.64
46:CS:45:THR:HG23	46:CS:46:PRO:HD2	1.79	0.64
13:D0:44:LEU:HD22	13:D0:48:VAL:HG23	1.79	0.64
27:D5:16:ARG:NH1	27:D5:17:ASP:OD1	2.30	0.64
55:DA:372:G:C2'	55:DA:373:U:OP2	2.46	0.64
55:DA:654(R):C:O5'	55:DA:654(R):C:H6	1.78	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:DG:44:GLY:HA2	6:DG:88:ILE:HG12	1.79	0.64
56:DI:29:GLU:CG	56:DJ:6:GLU:OE1	2.40	0.64
58:DL:11:GLN:CG	58:DL:12:LEU:N	2.57	0.64
9:DM:58:ASP:N	9:DM:60:ILE:CD1	2.60	0.64
55:DA:2277:G:H5''	12:DP:85:LYS:HB2	1.80	0.64
28:A6:22:ALA:HB3	28:A6:42:TRP:CH2	2.32	0.64
1:AA:1281:G:H5'	1:AA:1281:G:C8	2.25	0.64
1:AA:1359:A:OP2	1:AA:1359:A:N7	2.30	0.64
1:AA:1829:A:H3'	1:AA:1830:C:C6	2.31	0.64
1:AA:2807:G:H2'	1:AA:2808:U:H5''	1.79	0.64
1:AA:959:A:N6	12:AP:82:ARG:NH2	2.43	0.64
3:AD:186:HIS:CD2	3:AD:188:GLU:H	2.15	0.64
4:AE:95:ILE:HD12	4:AE:95:ILE:N	2.12	0.64
8:AK:58:LEU:O	8:AK:62:LYS:HB3	1.97	0.64
15:AR:86:ILE:HG12	15:AR:86:ILE:O	1.98	0.64
18:AS:18:ARG:HG3	18:AS:76:VAL:HG12	1.80	0.64
21:AV:92:SER:O	21:AV:130:PRO:HG2	1.97	0.64
31:BA:1020:U:H2'	31:BA:1021:G:H5''	1.79	0.64
31:BA:1347:G:OP2	39:BL:107:ARG:HG2	1.98	0.64
32:BE:142:LEU:O	32:BE:142:LEU:HD23	1.97	0.64
34:BG:189:PRO:HB2	34:BG:194:LEU:CD2	2.27	0.64
34:BG:7:PRO:HB2	34:BG:10:ARG:HD2	1.78	0.64
39:BL:113:LYS:N	39:BL:113:LYS:HD2	2.11	0.64
40:BM:74:ILE:HD13	40:BM:74:ILE:H	1.62	0.64
42:BO:60:LEU:HD22	42:BO:60:LEU:N	2.13	0.64
54:CA:1129:C:H5'	54:CA:1130:A:OP1	1.97	0.64
54:CA:1151:A:H1'	40:CM:39:PRO:HB2	1.78	0.64
54:CA:737:A:H2'	54:CA:738:C:C6	2.33	0.64
52:CC:28:G:C2'	52:CC:29:G:H5'	2.27	0.64
54:CA:1108:G:H5'	33:CF:176:HIS:CD2	2.31	0.64
38:CK:60:ARG:HG3	38:CK:60:ARG:NH1	2.13	0.64
49:CV:85:LYS:CG	49:CV:86:GLU:N	2.57	0.64
55:DA:2336:A:H61	22:D3:43:THR:HG21	1.63	0.64
3:DD:122:ASP:CG	3:DD:123:ALA:H	1.98	0.64
3:DD:237:GLU:OE2	3:DD:237:GLU:N	2.30	0.64
57:DY:132:ASP:OD2	56:DJ:10:GLU:CD	2.35	0.64
11:DO:125:VAL:HG13	11:DO:125:VAL:O	1.98	0.64
12:DP:26:TYR:O	12:DP:27:VAL:HB	1.98	0.64
18:DS:29:LEU:HD21	18:DS:33:ARG:NH2	2.12	0.64
21:DV:186:GLU:O	21:DV:187:ALA:CB	2.45	0.64
25:DX:7:LYS:NZ	25:DX:32:GLN:HG3	2.13	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:56:ASN:O	57:DY:60:ARG:HB2	1.98	0.64
57:DY:6:ASN:C	57:DY:7:VAL:HG12	2.17	0.64
1:AA:2468:G:O2'	1:AA:2469:A:H5''	1.97	0.64
1:AA:2702:U:H2'	1:AA:2702:U:O2	1.96	0.64
1:AA:2748:A:N7	1:AA:2757:A:C6	2.65	0.64
1:AA:278:A:H4'	1:AA:279:C:OP1	1.97	0.64
1:AA:654(R):C:H6	1:AA:654(R):C:O5'	1.81	0.64
1:AA:5:A:H2'	1:AA:6:A:O4'	1.97	0.64
1:AA:881:G:H5'	1:AA:882:G:OP2	1.97	0.64
4:AE:117:MET:O	4:AE:117:MET:HG2	1.97	0.64
6:AG:37:VAL:O	6:AG:94:LEU:HD23	1.98	0.64
11:AO:64:LYS:C	11:AO:66:GLY:N	2.48	0.64
20:AU:5:MET:HE1	20:AU:32:PRO:HB3	1.80	0.64
21:AV:6:LYS:HB3	21:AV:8:TYR:CE2	2.32	0.64
23:AZ:51:VAL:HG11	23:AZ:74:VAL:CG2	2.25	0.64
31:BA:1203:C:O2'	31:BA:1204:A:H5'	1.97	0.64
31:BA:1442:G:N7	31:BA:1446:A:N1	2.46	0.64
32:BE:97:TRP:HZ2	32:BE:102:LEU:HD13	1.60	0.64
49:BV:23:ASN:HB2	49:BV:43:GLU:CD	2.17	0.64
54:CA:1032:A:H3'	54:CA:1032(A):G:H4'	1.80	0.64
54:CA:1346:A:H5''	39:CL:120:ARG:HH12	1.62	0.64
54:CA:179:A:O2'	54:CA:180:U:H5'	1.96	0.64
54:CA:918:A:H2'	54:CA:919:A:C8	2.32	0.64
32:CE:61:LEU:HG	32:CE:68:ILE:HD11	1.78	0.64
41:CN:108:ILE:H	48:CU:87:ARG:HE	1.44	0.64
29:D7:5:TRP:HE1	29:D7:7:PRO:HG3	1.61	0.64
55:DA:1340:U:O2'	55:DA:1602:U:H2'	1.98	0.64
55:DA:1348:G:H2'	55:DA:1349:A:C5'	2.26	0.64
55:DA:2127:G:C3'	55:DA:2128:C:H5''	2.27	0.64
55:DA:2656:U:C5	55:DA:2657:A:N7	2.66	0.64
55:DA:805:G:H4'	55:DA:806:C:OP2	1.97	0.64
3:DD:35:LYS:CD	3:DD:104:TYR:HD1	2.10	0.64
4:DE:65:GLY:HA2	4:DE:70:ALA:CB	2.27	0.64
56:DI:20:LEU:HA	56:DI:24:ILE:HG21	1.79	0.64
58:DL:93:ARG:NH1	58:DL:135:GLY:CA	2.58	0.64
24:DW:50:ILE:CD1	24:DW:51:ARG:N	2.59	0.64
57:DY:131:MET:O	57:DY:132:ASP:C	2.36	0.64
57:DY:59:ILE:HG13	57:DY:60:ARG:N	2.12	0.64
57:DY:74:LEU:HB3	57:DY:120:LYS:HE2	1.78	0.64
1:AA:1379:A:O4'	1:AA:1379:A:P	2.56	0.64
1:AA:118:A:N3	1:AA:178:G:H1'	2.13	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2602:A:H4'	1:AA:2603:G:C5'	2.27	0.64
1:AA:654(H):G:O5'	1:AA:654(H):G:H8	1.81	0.64
2:AB:15:A:H3'	2:AB:16:G:C5'	2.28	0.64
4:AE:55:ASN:HD21	4:AE:75:VAL:HA	1.63	0.64
7:AH:89:ILE:H	7:AH:89:ILE:HD13	1.62	0.64
31:BA:1348:U:H2'	31:BA:1349:A:H8	1.61	0.64
31:BA:598:U:H2'	31:BA:599:C:H6	1.62	0.64
31:BA:827:U:N3	31:BA:872:A:N6	2.43	0.64
39:BL:11:LYS:H	39:BL:104:ARG:NH2	1.95	0.64
39:BL:4:TYR:O	39:BL:18:PHE:HA	1.98	0.64
41:BN:13:GLN:HB2	41:BN:76:GLY:HA3	1.80	0.64
54:CA:163:C:O2'	54:CA:164:U:H5'	1.96	0.64
32:CE:215:LEU:O	32:CE:218:ALA:HB3	1.97	0.64
32:CE:69:LEU:HD13	32:CE:91:PRO:HB2	1.78	0.64
35:CH:7:GLU:HG2	35:CH:112:LEU:HD22	1.80	0.64
17:D2:44:LYS:O	17:D2:46:VAL:N	2.30	0.64
17:D2:76:LYS:O	17:D2:79:VAL:HG12	1.97	0.64
55:DA:1011:G:O2'	55:DA:1013:C:H5'	1.97	0.64
55:DA:1929:G:C4'	55:DA:1930:G:OP1	2.46	0.64
55:DA:2469:A:H61	55:DA:2481:G:H1'	1.62	0.64
4:DE:120:TRP:CE3	4:DE:155:LYS:HD3	2.33	0.64
6:DG:55:LYS:HZ1	6:DG:148:MET:HG3	1.62	0.64
55:DA:270(L):U:H3	8:DK:50:ARG:CZ	2.11	0.64
58:DL:115:LEU:HD11	58:DL:117:THR:HG1	1.63	0.64
58:DL:19:PRO:C	58:DL:25:PRO:CG	2.65	0.64
58:DL:72:PRO:N	58:DL:73:PRO:HD3	2.13	0.64
19:DT:41:ASN:N	19:DT:41:ASN:HD22	1.95	0.64
21:DV:61:LEU:CD1	21:DV:65:GLN:CB	2.66	0.64
57:DY:29:TYR:HD2	57:DY:30:GLN:H	1.45	0.64
57:DY:65:GLU:C	57:DY:66:LEU:HG	2.18	0.64
22:A3:32:ARG:N	22:A3:35:ASN:HD21	1.90	0.64
1:AA:1025:G:N3	1:AA:1025:G:H2'	2.13	0.64
1:AA:66:C:H5'	1:AA:456:C:O2	1.97	0.64
11:AO:71:VAL:HG13	11:AO:72:PRO:CD	2.27	0.64
21:AV:175:VAL:CA	21:AV:177:PRO:HD2	2.24	0.64
21:AV:184:ALA:C	21:AV:186:GLU:H	1.99	0.64
21:AV:94:GLU:O	21:AV:129:SER:HA	1.97	0.64
8:AK:27:ARG:HD2	23:AZ:71:TYR:CE1	2.33	0.64
32:BE:221:LEU:HD12	32:BE:221:LEU:C	2.18	0.64
32:BE:5:ILE:HD11	32:BE:221:LEU:HD22	1.80	0.64
35:BH:78:HIS:ND1	38:BK:107:LEU:HD12	2.11	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:BU:72:ARG:O	48:BU:76:LEU:HD12	1.97	0.64
54:CA:1371:G:O3'	39:CL:69:GLY:HA3	1.97	0.64
54:CA:1412:C:H2'	54:CA:1413:A:C8	2.33	0.64
54:CA:542:G:OP1	34:CG:10:ARG:NH2	2.31	0.64
54:CA:950:U:OP2	43:CP:102:ARG:HD2	1.96	0.64
52:CD:8:U:H2'	52:CD:13:C:N4	2.00	0.64
32:CE:55:PHE:HE1	32:CE:218:ALA:HA	1.62	0.64
35:CH:40:ARG:HH11	35:CH:40:ARG:CB	2.07	0.64
36:CI:62:TRP:CH2	36:CI:64:GLN:HB2	2.32	0.64
49:CV:63:THR:O	49:CV:66:MET:HG3	1.98	0.64
17:D2:46:VAL:HG13	17:D2:46:VAL:O	1.97	0.64
26:D4:70:GLY:O	26:D4:71:ARG:CB	2.44	0.64
30:D8:51:ALA:HA	30:D8:54:GLU:HG3	1.79	0.64
3:DD:70:TRP:HZ3	3:DD:146:GLU:OE2	1.79	0.64
57:DY:139:VAL:CG2	56:DJ:6:GLU:OE2	2.46	0.64
58:DL:146:ASP:OD1	58:DL:146:ASP:N	2.30	0.64
15:DR:129:ARG:O	15:DR:132:LYS:HB3	1.97	0.64
17:A2:5:VAL:HA	17:A2:37:VAL:HB	1.80	0.64
1:AA:2393:A:H5'	30:A8:30:ARG:HD3	1.80	0.64
30:A8:23:VAL:CG1	30:A8:47:LYS:HD3	2.28	0.64
30:A8:6:THR:HA	30:A8:61:LEU:HD11	1.79	0.64
1:AA:1045:A:C3'	1:AA:1046:A:H5''	2.28	0.64
1:AA:1085:A:H2'	1:AA:1086:A:C8	2.32	0.64
1:AA:1496:A:C8	1:AA:1577:C:O2'	2.46	0.64
1:AA:2154:G:H2'	1:AA:2155:G:C8	2.32	0.64
1:AA:2298:A:N6	1:AA:2318:G:H8	1.95	0.64
6:AG:82:LEU:HD23	6:AG:86:MET:SD	2.37	0.64
12:AP:43:THR:HA	12:AP:94:VAL:HG12	1.78	0.64
21:AV:142:SER:O	21:AV:144:LEU:N	2.30	0.64
21:AV:21:ALA:O	21:AV:23:LYS:HG2	1.97	0.64
21:AV:60:GLU:HA	21:AV:66:SER:HA	1.79	0.64
24:AW:13:ALA:O	24:AW:16:LEU:HG	1.97	0.64
24:AW:51:ARG:HH21	24:AW:55:ARG:HH12	1.46	0.64
31:BA:1130:A:N6	31:BA:1144:G:N2	2.45	0.64
31:BA:1176:A:C2'	31:BA:1177:G:H5'	2.28	0.64
31:BA:711:G:O2'	31:BA:712:A:H5'	1.98	0.64
52:BD:20:U:H2'	52:BD:21:A:C5'	2.27	0.64
32:BE:194:PRO:O	32:BE:196:LEU:N	2.30	0.64
34:BG:150:GLU:CD	34:BG:150:GLU:H	2.00	0.64
49:BV:31:ILE:HG23	49:BV:49:ILE:HG23	1.79	0.64
54:CA:199:G:O2'	54:CA:200:G:H5'	1.96	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:69:G:N1	54:CA:73:G:N7	2.46	0.64
52:CB:44:G:H2'	52:CB:45:U:C6	2.33	0.64
43:CP:123:ALA:HB1	43:CP:124:PRO:CD	2.27	0.64
43:CP:125:ARG:O	43:CP:126:LYS:O	2.16	0.64
48:CU:50:ILE:N	48:CU:50:ILE:HD12	2.13	0.64
55:DA:2182:G:H2'	55:DA:2183:C:C6	2.33	0.64
55:DA:2845:G:O2'	55:DA:2846:G:H5'	1.96	0.64
3:DD:25:THR:HG23	3:DD:27:THR:HB	1.79	0.64
7:DH:86:GLU:HG3	7:DH:165:ALA:CB	2.27	0.64
56:DJ:5:ILE:O	56:DJ:9:LYS:HB2	1.98	0.64
58:DL:50:ASP:N	58:DL:53:VAL:CG2	2.57	0.64
58:DL:7:VAL:HG12	58:DL:57:ILE:HD12	0.65	0.64
14:DQ:106:ARG:HA	14:DQ:110:LEU:CD1	2.26	0.64
15:DR:109:GLU:OE1	15:DR:112:ARG:HD3	1.98	0.64
10:DN:104:ARG:HH21	15:DR:43:GLN:NE2	1.96	0.64
20:DU:49:VAL:O	20:DU:51:VAL:N	2.31	0.64
20:DU:78:ALA:HB3	20:DU:81:LYS:HE3	1.79	0.64
21:DV:183:LEU:O	21:DV:184:ALA:CB	2.45	0.64
24:DW:50:ILE:HD12	24:DW:51:ARG:N	2.09	0.64
57:DY:134:LEU:O	57:DY:137:GLU:CG	2.46	0.64
57:DY:22:GLY:O	57:DY:23:SER:CB	2.45	0.64
57:DY:23:SER:OG	57:DY:114:GLY:CA	2.34	0.64
11:AO:64:LYS:NZ	30:A8:30:ARG:HA	2.12	0.64
1:AA:1410:G:H2'	1:AA:1411:C:C6	2.32	0.64
1:AA:1821:A:H2'	1:AA:1822:G:H5'	1.80	0.64
1:AA:999:U:H2'	1:AA:1000:A:C5'	2.27	0.64
3:AD:236:GLY:O	3:AD:237:GLU:CB	2.39	0.64
11:AO:9:ASN:HB2	11:AO:10:PRO:HD2	1.78	0.64
12:AP:12:GLN:HE21	12:AP:73:PRO:HD2	1.61	0.64
25:AX:44:ARG:O	25:AX:48:GLU:HG3	1.96	0.64
31:BA:1067:A:H4'	31:BA:1068:G:O5'	1.98	0.64
52:BB:75:C:H6	52:BB:75:C:H3'	1.63	0.64
48:CU:56:THR:HB	48:CU:58:LEU:CD1	2.28	0.64
30:D8:6:THR:O	30:D8:7:HIS:CB	2.46	0.64
55:DA:1608:A:H4'	55:DA:1609:A:OP1	1.97	0.64
4:DE:137:HIS:HB3	4:DE:138:PRO:HD2	1.79	0.64
58:DL:69:THR:C	58:DL:70:LYS:HG3	2.18	0.64
55:DA:958:U:OP2	12:DP:14:ARG:NH1	2.31	0.64
21:DV:6:LYS:O	21:DV:7:ALA:CB	2.45	0.64
24:DW:15:LYS:O	24:DW:16:LEU:HB3	1.98	0.64
57:DY:35:LYS:HE3	57:DY:35:LYS:O	1.98	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:38:HIS:O	57:DY:96:PHE:CZ	2.51	0.64
1:AA:1431:U:H2'	1:AA:1432:C:H6	1.63	0.63
1:AA:1813:G:H1'	3:AD:50:THR:OG1	1.98	0.63
1:AA:654(I):C:O2'	1:AA:654(J):A:C5'	2.46	0.63
3:AD:25:THR:CG2	3:AD:82:ILE:H	2.11	0.63
7:AH:16:SER:HB3	7:AH:26:VAL:O	1.97	0.63
8:AK:109:ILE:H	8:AK:109:ILE:CD1	2.04	0.63
8:AK:10:GLU:CD	8:AK:11:ASN:HB2	2.18	0.63
21:AV:178:GLU:O	21:AV:180:VAL:N	2.30	0.63
24:AW:4:SER:OG	24:AW:5:GLU:OE2	2.12	0.63
31:BA:1116:C:C3'	31:BA:1117:G:H5''	2.29	0.63
32:BE:56:ARG:CB	32:BE:56:ARG:HH11	2.08	0.63
37:BJ:60:LYS:O	37:BJ:63:LYS:HB3	1.98	0.63
38:BK:7:ALA:HB2	38:BK:85:ARG:CD	2.28	0.63
40:BM:47:PHE:HE1	40:BM:63:PHE:HB2	1.63	0.63
54:CA:1227:A:C2	49:CV:84:GLY:HA3	2.33	0.63
54:CA:641:U:C4'	54:CA:642:A:OP1	2.46	0.63
54:CA:687:A:O2'	54:CA:688:G:O4'	2.14	0.63
27:D5:40:LYS:HG2	27:D5:47:PRO:HD2	1.81	0.63
27:D5:56:LYS:N	27:D5:56:LYS:HD2	2.11	0.63
55:DA:2419:U:C5'	28:D6:23:THR:HG21	2.28	0.63
55:DA:1899:G:N2	55:DA:1902:C:C4	2.63	0.63
55:DA:2115:G:H2'	55:DA:2116:G:N7	2.13	0.63
55:DA:2448:A:C4'	55:DA:2449:U:OP2	2.42	0.63
55:DA:2472:G:H22	55:DA:2477:C:H5''	1.62	0.63
55:DA:2610:C:H4'	55:DA:2611:U:OP2	1.98	0.63
55:DA:2716:U:O2'	55:DA:2717:G:H5'	1.98	0.63
55:DA:654(C):G:H3'	55:DA:654(D):G:C8	2.31	0.63
55:DA:760:G:H2'	55:DA:761:A:O4'	1.98	0.63
10:DN:2:ILE:HD11	10:DN:82:ASN:ND2	2.11	0.63
15:DR:85:LYS:HE2	15:DR:87:ASP:OD2	1.97	0.63
21:DV:60:GLU:HG3	21:DV:61:LEU:N	2.13	0.63
25:DX:6:VAL:HB	25:DX:54:VAL:HG21	1.80	0.63
57:DY:122:VAL:CA	57:DY:126:ALA:CB	2.75	0.63
13:A0:104:ARG:NH1	13:A0:109:ALA:HB3	2.14	0.63
1:AA:1156:A:H5''	1:AA:1157:G:OP2	1.99	0.63
1:AA:2853:C:H2'	1:AA:2854:G:C8	2.33	0.63
1:AA:310:A:OP1	20:AU:17:SER:O	2.17	0.63
1:AA:654(B):C:H2'	1:AA:654(C):G:C1'	2.28	0.63
3:AD:28:GLU:HB2	3:AD:29:PRO:HD3	1.79	0.63
11:AO:48:PRO:O	11:AO:50:ARG:N	2.30	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:AQ:23:ARG:HG2	14:AQ:23:ARG:NH1	2.13	0.63
18:AS:39:THR:CG2	18:AS:44:ALA:HB2	2.27	0.63
24:AW:15:LYS:HD3	24:AW:67:LYS:HZ1	1.63	0.63
31:BA:1363:A:H4'	31:BA:1364:U:H5''	1.80	0.63
31:BA:468:A:H2'	31:BA:474:G:O4'	1.99	0.63
52:BB:9:A:H2	52:BB:11:C:N4	1.96	0.63
35:BH:144:THR:O	35:BH:148:VAL:HG23	1.98	0.63
42:BO:6:THR:H	42:BO:9:GLN:HE21	1.46	0.63
51:BX:9:ARG:O	51:BX:13:ILE:HG13	1.98	0.63
54:CA:1006:C:H2'	54:CA:1007:C:H6	1.64	0.63
54:CA:1007:C:H2'	54:CA:1008:C:C5'	2.15	0.63
54:CA:1072:G:H2'	54:CA:1073:U:C6	2.32	0.63
54:CA:511:C:H1'	34:CG:43:HIS:HE2	1.63	0.63
54:CA:530:G:O2'	54:CA:531:U:P	2.56	0.63
54:CA:714:G:H2'	54:CA:715:A:C8	2.33	0.63
54:CA:792:A:O2'	54:CA:793:U:P	2.57	0.63
52:CB:11:C:O2'	52:CB:12:U:H5'	1.97	0.63
52:CB:68:C:H2'	52:CB:69:G:C8	2.33	0.63
34:CG:144:ASP:HB2	34:CG:146:ILE:HD11	1.80	0.63
38:CK:12:ARG:HH11	38:CK:26:VAL:HA	1.63	0.63
13:D0:49:ASP:OD1	13:D0:95:THR:HG22	1.97	0.63
55:DA:1301:A:O2'	55:DA:1302:A:C3'	2.30	0.63
55:DA:242:G:O2'	55:DA:243:U:OP2	2.15	0.63
55:DA:2712:U:H2'	55:DA:2712(A):A:O5'	1.96	0.63
55:DA:312:G:H5'	55:DA:331:A:H2'	1.79	0.63
55:DA:654(M):C:C3'	55:DA:654(N):G:N7	2.52	0.63
7:DH:4:ILE:HD13	7:DH:4:ILE:N	2.12	0.63
56:DI:10:GLU:O	56:DI:14:GLN:CB	2.46	0.63
58:DL:41:PHE:CG	58:DL:42:ASN:N	2.65	0.63
11:DO:127:ALA:C	11:DO:147:LEU:HD23	2.18	0.63
12:DP:1:MET:HG2	12:DP:1:MET:O	1.96	0.63
14:DQ:52:SER:HB2	14:DQ:55:ALA:H	1.61	0.63
54:CA:1432:G:OP1	15:DR:107:ASP:HB2	1.99	0.63
15:DR:23:ARG:HB2	15:DR:24:PRO:HD2	1.78	0.63
57:DY:44:LEU:O	58:DL:119:ASP:HB3	1.98	0.63
23:DZ:23:LYS:HE3	23:DZ:29:GLY:HA2	1.79	0.63
22:A3:82:ARG:HG2	22:A3:84:LEU:HD22	1.80	0.63
1:AA:593:G:H1'	30:A8:4:MET:HE2	1.81	0.63
1:AA:1728:G:C6	1:AA:1730:U:OP2	2.50	0.63
1:AA:2165:G:N3	1:AA:2165:G:H2'	2.13	0.63
1:AA:528:A:C8	1:AA:528:A:H3'	2.34	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:616:A:O2'	1:AA:617:G:OP1	2.14	0.63
1:AA:869:G:O2'	1:AA:870:A:H5'	1.99	0.63
3:AD:228:PRO:HD3	3:AD:234:GLY:O	1.98	0.63
4:AE:6:GLY:HA2	4:AE:51:PHE:CZ	2.33	0.63
6:AG:101:ILE:HB	26:A4:25:TYR:CD2	2.33	0.63
15:AR:19:LEU:HD22	15:AR:86:ILE:HG21	1.80	0.63
21:AV:115:GLY:N	21:AV:177:PRO:HG2	2.10	0.63
23:AZ:92:LYS:O	23:AZ:94:LEU:N	2.31	0.63
31:BA:1055:A:N6	31:BA:1200:C:N3	2.47	0.63
31:BA:769:G:H4'	31:BA:1513:A:H4'	1.81	0.63
31:BA:971:G:N2	31:BA:1363:A:OP2	2.30	0.63
52:BB:2:C:H2'	52:BB:3:C:H6	1.63	0.63
34:BG:12:CYS:CA	34:BG:21:LEU:CD2	2.75	0.63
36:BI:45:LEU:HD23	36:BI:46:ARG:N	2.13	0.63
54:CA:1028(A):C:H2'	54:CA:1028(B):C:H6	1.62	0.63
54:CA:1322:C:HO2'	54:CA:1323:G:C5'	2.11	0.63
46:CS:72:ARG:HD3	46:CS:72:ARG:C	2.19	0.63
16:D1:83:LEU:HA	16:D1:88:ILE:CD1	2.27	0.63
55:DA:1085:A:O4'	55:DA:1105:U:H1'	1.97	0.63
55:DA:1203:G:H5'	11:DO:3:LEU:HD12	1.79	0.63
55:DA:1493:C:O2	55:DA:1493:C:H2'	1.97	0.63
55:DA:2503:A:O2'	55:DA:2505:G:OP2	2.16	0.63
55:DA:2:G:O2'	55:DA:3:U:H5'	1.98	0.63
4:DE:150:VAL:CG1	4:DE:154:LYS:HG3	2.29	0.63
5:DF:129:PHE:HA	5:DF:142:TRP:NE1	2.13	0.63
5:DF:32:LEU:HD21	5:DF:108:LYS:HB3	1.80	0.63
55:DA:468:G:H4'	5:DF:62:ARG:HH12	1.62	0.63
56:DJ:21:LYS:O	56:DJ:24:ILE:CB	2.45	0.63
11:DO:98:GLU:O	11:DO:101:VAL:HG12	1.98	0.63
19:DT:49:VAL:HG13	19:DT:87:GLN:NE2	2.13	0.63
20:DU:14:LEU:HD23	20:DU:14:LEU:C	2.18	0.63
57:DY:27:VAL:HG23	57:DY:80:VAL:HG21	1.79	0.63
23:DZ:60:PHE:HE2	23:DZ:91:LYS:NZ	1.96	0.63
1:AA:533:G:H5'	16:A1:24:TYR:CE2	2.34	0.63
16:A1:50:ARG:HH11	17:A2:72:VAL:CB	2.12	0.63
16:A1:50:ARG:NH2	16:A1:50:ARG:HB2	2.13	0.63
29:A7:24:THR:HG23	29:A7:27:GLY:N	2.10	0.63
1:AA:1022:G:O2'	1:AA:1023:U:P	2.56	0.63
1:AA:171:G:H2'	1:AA:172:C:H6	1.61	0.63
1:AA:1820:U:H4'	1:AA:1821:A:OP2	1.99	0.63
1:AA:1926:U:C6	1:AA:1928:A:OP2	2.52	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2879:C:C4'	1:AA:2880:C:OP1	2.45	0.63
5:AF:31:HIS:O	5:AF:34:TRP:HB3	1.98	0.63
6:AG:123:ASN:HD22	6:AG:123:ASN:N	1.96	0.63
11:AO:124:LYS:HG2	11:AO:145:PRO:HD3	1.80	0.63
4:AE:14:ILE:HD11	15:AR:14:TYR:OH	1.99	0.63
31:BA:1374:A:H2'	31:BA:1375:A:C5'	2.27	0.63
31:BA:404:U:H2'	31:BA:405:U:H6	1.63	0.63
31:BA:417:C:O2'	31:BA:418:C:H5'	1.99	0.63
31:BA:646:U:O2'	31:BA:647:C:H5'	1.98	0.63
31:BA:734:G:N2	48:BU:75:ILE:HD11	2.13	0.63
31:BA:745:C:H2'	31:BA:746:A:H8	1.64	0.63
31:BA:731:G:OP1	31:BA:766:A:H1'	1.99	0.63
31:BA:892:A:O2'	31:BA:1415:G:H4'	1.99	0.63
32:BE:63:MET:HG2	32:BE:225:ALA:HB1	1.79	0.63
32:BE:36:ARG:HG2	32:BE:37:ASN:ND2	2.13	0.63
34:BG:152:SER:HA	34:BG:155:LEU:HD12	1.80	0.63
34:BG:25:ARG:O	34:BG:27:TYR:N	2.31	0.63
36:BI:78:GLU:HA	36:BI:81:ILE:HD11	1.79	0.63
45:BR:64:ARG:NH1	45:BR:68:ARG:HH21	1.97	0.63
54:CA:474:G:O2'	54:CA:475:G:H5'	1.98	0.63
52:CD:20:U:O2'	52:CD:21:A:H5'	1.98	0.63
32:CE:52:GLU:HG2	32:CE:56:ARG:NH1	2.12	0.63
34:CG:148:VAL:HG12	34:CG:149:ALA:N	2.13	0.63
35:CH:110:LEU:CD2	35:CH:139:LEU:HD21	2.29	0.63
36:CI:82:ARG:HB2	36:CI:85:VAL:HG23	1.81	0.63
40:CM:40:LEU:HB2	40:CM:69:ASN:CB	2.27	0.63
44:CQ:40:CYS:N	44:CQ:43:CYS:HB2	2.13	0.63
48:CU:32:ARG:HA	48:CU:69:THR:HG21	1.80	0.63
26:D4:27:THR:O	26:D4:28:LYS:HB3	1.99	0.63
26:D4:65:ASP:O	26:D4:67:TYR:N	2.29	0.63
26:D4:68:ARG:HA	26:D4:68:ARG:NH2	2.11	0.63
55:DA:1082:U:OP1	55:DA:1082:U:H4'	1.98	0.63
55:DA:2506:U:O2'	55:DA:2507:C:C5'	2.46	0.63
55:DA:2895:U:H2'	55:DA:2896:C:H6	1.64	0.63
55:DA:303:U:H2'	55:DA:304:G:C8	2.33	0.63
55:DA:557:U:H2'	55:DA:558:G:H8	1.64	0.63
55:DA:605:C:O2'	55:DA:606:U:H5'	1.98	0.63
3:DD:27:THR:O	3:DD:28:GLU:HB2	1.99	0.63
4:DE:56:PRO:O	4:DE:57:LYS:HB2	1.98	0.63
8:DK:130:TYR:C	8:DK:131:LYS:HD2	2.18	0.63
58:DL:8:VAL:CA	58:DL:57:ILE:HG13	2.26	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:DM:32:THR:O	9:DM:35:ARG:O	2.16	0.63
11:DO:84:ASN:HB2	11:DO:87:ASP:OD2	1.98	0.63
12:DP:58:PHE:CD1	12:DP:61:GLY:HA3	2.32	0.63
10:DN:107:ARG:NH1	15:DR:36:GLU:HB3	2.13	0.63
15:DR:50:ILE:HA	15:DR:99:LEU:HD11	1.80	0.63
18:DS:41:LYS:HE3	27:D5:25:LEU:HD11	1.80	0.63
19:DT:64:LYS:C	19:DT:65:ARG:HD3	2.19	0.63
16:A1:92:ARG:O	16:A1:93:LYS:C	2.37	0.63
1:AA:1955:U:O2'	1:AA:1956:U:OP1	2.14	0.63
1:AA:2468:G:N2	1:AA:2481:G:H2'	2.14	0.63
1:AA:479:A:H4'	1:AA:480:A:OP1	1.98	0.63
3:AD:267:SER:C	3:AD:269:PHE:H	2.02	0.63
4:AE:11:MET:SD	4:AE:24:THR:HG22	2.39	0.63
9:AM:120:LEU:HD23	9:AM:120:LEU:O	1.98	0.63
20:AU:81:LYS:HZ3	20:AU:97:ARG:NH2	1.97	0.63
31:BA:1004:A:H2'	31:BA:1005:A:O5'	1.98	0.63
31:BA:50:A:O2'	31:BA:52:G:C8	2.52	0.63
31:BA:954:G:H2'	31:BA:955:U:C6	2.34	0.63
54:CA:371:G:H2'	54:CA:372:C:O4'	1.98	0.63
54:CA:630:G:O2'	54:CA:631:G:P	2.56	0.63
52:CD:2:C:O4'	52:CD:2:C:OP1	2.17	0.63
32:CE:140:HIS:O	32:CE:144:ARG:HG2	1.99	0.63
32:CE:172:ILE:O	32:CE:175:ARG:HB3	1.97	0.63
32:CE:235:SER:C	32:CE:237:ALA:H	2.02	0.63
32:CE:39:ILE:HG22	32:CE:40:HIS:O	1.97	0.63
39:CL:97:LYS:HB3	39:CL:98:PRO:HD3	1.81	0.63
40:CM:54:PHE:CD2	40:CM:55:LYS:HD2	2.33	0.63
47:CT:100:LYS:O	47:CT:101:ARG:HG3	1.98	0.63
50:CW:36:LEU:HD13	50:CW:39:LYS:HD3	1.79	0.63
16:D1:92:ARG:NH1	16:D1:95:LEU:HD11	2.13	0.63
17:D2:28:GLU:O	17:D2:61:VAL:HG11	1.99	0.63
11:DO:61:ARG:NH1	30:D8:13:ARG:HG3	2.14	0.63
55:DA:1062:G:C2'	55:DA:1077:A:H61	2.10	0.63
55:DA:1845:G:OP1	3:DD:258:LYS:NZ	2.30	0.63
55:DA:2158:A:H4'	55:DA:2159:G:O5'	1.99	0.63
55:DA:2839:G:C5'	13:D0:46:GLY:HA2	2.28	0.63
55:DA:288:C:O2'	55:DA:289:A:H5'	1.97	0.63
6:DG:65:GLY:HA3	26:D4:9:LEU:HD12	1.80	0.63
58:DL:82:ALA:O	58:DL:85:GLU:HB2	1.98	0.63
9:DM:61:ARG:HA	9:DM:61:ARG:HE	1.63	0.63
14:DQ:19:LYS:O	14:DQ:21:THR:N	2.31	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:DS:110:LYS:HG3	18:DS:111:HIS:ND1	2.13	0.63
21:DV:28:MET:SD	21:DV:37:VAL:HG11	2.39	0.63
21:DV:27:VAL:CG1	21:DV:87:ASP:HB3	2.28	0.63
57:DY:138:LEU:O	57:DY:140:GLY:N	2.31	0.63
57:DY:50:ARG:NH2	57:DY:83:TYR:HE1	1.96	0.63
16:A1:92:ARG:HH12	17:A2:11:GLN:HG3	1.63	0.63
28:A6:28:ARG:HB3	28:A6:30:THR:O	1.98	0.63
30:A8:9:GLY:O	30:A8:13:ARG:HG3	1.99	0.63
1:AA:1332:G:H5'	1:AA:1333:C:OP2	1.99	0.63
1:AA:1747:G:O2'	1:AA:1748:G:H5'	1.98	0.63
1:AA:2111:C:O2'	1:AA:2118:U:H4'	1.98	0.63
1:AA:2414:G:H21	11:AO:67:MET:CE	2.12	0.63
1:AA:2657:A:H2'	1:AA:2658:C:H5'	1.79	0.63
4:AE:201:THR:HG22	4:AE:202:LYS:N	2.11	0.63
4:AE:57:LYS:HZ3	4:AE:72:VAL:HG22	1.63	0.63
6:AG:41:GLN:O	6:AG:89:GLY:HA2	1.99	0.63
7:AH:149:ARG:HG3	7:AH:162:ILE:O	1.99	0.63
7:AH:41:MET:HG3	7:AH:54:ARG:CA	2.29	0.63
1:AA:1012:U:O4	9:AM:25:ARG:HA	1.99	0.63
21:AV:130:PRO:C	21:AV:133:ILE:HD11	2.19	0.63
21:AV:144:LEU:O	21:AV:146:ILE:N	2.31	0.63
21:AV:115:GLY:N	21:AV:177:PRO:CG	2.62	0.63
21:AV:37:VAL:O	21:AV:38:TYR:HB3	1.97	0.63
31:BA:1468:A:H2'	31:BA:1469:G:O4'	1.98	0.63
31:BA:328:C:O2'	31:BA:329:A:P	2.57	0.63
31:BA:508:C:H1'	31:BA:509:A:N7	2.14	0.63
34:BG:23:GLY:HA2	34:BG:27:TYR:CD1	2.34	0.63
36:BI:19:LEU:O	36:BI:23:LYS:HG3	1.98	0.63
40:BM:48:THR:CA	40:BM:62:HIS:HB3	2.25	0.63
42:BO:41:ARG:CB	42:BO:41:ARG:HH11	2.04	0.63
54:CA:1157:A:H1'	54:CA:1158:C:C4	2.32	0.63
54:CA:1293:G:H2'	54:CA:1294:G:H8	1.63	0.63
54:CA:411:A:N7	54:CA:413:G:N3	2.47	0.63
52:CB:74:C:O2'	52:CB:75:C:P	2.57	0.63
32:CE:122:PHE:HD1	32:CE:139:LYS:NZ	1.93	0.63
32:CE:141:GLU:O	32:CE:145:LEU:HD23	1.99	0.63
37:CJ:133:GLY:O	37:CJ:136:LYS:HB2	1.98	0.63
44:CQ:22:THR:O	44:CQ:23:ARG:HB2	1.97	0.63
49:CV:83:HIS:C	49:CV:85:LYS:N	2.52	0.63
55:DA:2173:A:C4	55:DA:2174:C:H1'	2.33	0.63
55:DA:601:C:O2'	55:DA:605:C:OP1	2.17	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:DL:104:VAL:O	58:DL:107:ILE:HB	1.98	0.63
58:DL:52:ILE:CG1	58:DL:53:VAL:H	2.06	0.63
58:DL:69:THR:HG22	58:DL:70:LYS:N	2.08	0.63
14:DQ:83:LYS:HE3	14:DQ:109:GLY:HA2	1.81	0.63
20:DU:79:CYS:O	20:DU:80:GLY:O	2.17	0.63
57:DY:19:ARG:CD	57:DY:20:ALA:N	2.62	0.63
57:DY:2:PRO:HG2	57:DY:3:ASN:N	2.02	0.63
57:DY:43:ALA:H	57:DY:47:ASN:ND2	1.95	0.63
30:A8:34:TRP:HD1	30:A8:35:GLN:H	1.45	0.63
30:A8:60:LEU:C	30:A8:61:LEU:HD12	2.18	0.63
1:AA:2189:U:C2'	1:AA:2190:G:H5''	2.29	0.63
1:AA:2189:U:H2'	1:AA:2190:G:H5''	1.80	0.63
1:AA:2311:A:H3'	1:AA:2312:U:C6	2.34	0.63
5:AF:117:ARG:NH2	5:AF:187:VAL:HA	2.13	0.63
8:AK:51:ILE:HG22	8:AK:52:ARG:N	2.14	0.63
10:AN:71:ARG:NH2	10:AN:77:ILE:HG21	2.13	0.63
20:AU:43:ASN:N	20:AU:43:ASN:ND2	2.46	0.63
31:BA:366:C:O2'	31:BA:394:G:N2	2.32	0.63
32:BE:22:LYS:HZ2	32:BE:22:LYS:N	1.95	0.63
39:BL:65:VAL:HG11	39:BL:73:GLN:HB3	1.81	0.63
43:BP:108:ARG:HD2	43:BP:108:ARG:N	2.14	0.63
8:AK:87:LYS:HD2	54:CA:359:U:P	2.38	0.63
34:CG:176:LEU:HD12	34:CG:177:ASP:H	1.64	0.63
35:CH:68:GLU:O	35:CH:68:GLU:HG3	1.98	0.63
54:CA:538:G:OP2	42:CO:115:LYS:HG3	1.99	0.63
46:CS:4:ILE:CD1	46:CS:64:ALA:HB1	2.26	0.63
41:CN:108:ILE:H	48:CU:87:ARG:NE	1.95	0.63
55:DA:1057:A:H4'	55:DA:1058:U:OP1	1.98	0.63
55:DA:1058:U:OP1	58:DL:5:VAL:CG2	2.46	0.63
55:DA:1668:A:N6	55:DA:1676:A:H61	1.95	0.63
55:DA:1879:C:H2'	55:DA:1880:C:C5'	2.29	0.63
55:DA:2291:U:H2'	55:DA:2292:C:C6	2.34	0.63
55:DA:2292:C:O2'	55:DA:2293:C:H5'	1.98	0.63
55:DA:642:G:N2	55:DA:644:A:H3'	2.14	0.63
55:DA:860:U:C5	55:DA:917:A:N1	2.59	0.63
7:DH:153:LYS:HG3	7:DH:161:GLY:HA2	1.79	0.63
7:DH:89:ILE:HD11	7:DH:129:THR:CB	2.18	0.63
56:DI:10:GLU:O	56:DI:14:GLN:N	2.31	0.63
56:DI:16:THR:O	56:DI:20:LEU:CD1	2.47	0.63
56:DJ:5:ILE:C	56:DJ:7:ARG:N	2.52	0.63
58:DL:95:LYS:CA	58:DL:136:VAL:HG11	2.29	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1077:A:C2'	58:DL:93:ARG:HH22	2.11	0.63
12:DP:33:GLY:HA2	12:DP:105:GLU:HA	1.81	0.63
55:DA:2318:G:H22	14:DQ:2:ALA:N	1.97	0.63
14:DQ:58:LEU:HD23	14:DQ:58:LEU:N	2.14	0.63
21:DV:185:GLU:OE1	21:DV:185:GLU:CA	2.45	0.63
57:DY:21:GLN:HE21	57:DY:21:GLN:C	1.96	0.63
57:DY:5:ARG:HD3	57:DY:7:VAL:HG11	1.79	0.63
16:A1:109:LEU:HD21	17:A2:40:LEU:HD21	1.80	0.63
17:A2:43:GLU:C	17:A2:44:LYS:HD3	2.20	0.63
27:A5:46:CYS:SG	27:A5:47:PRO:HD2	2.38	0.63
30:A8:41:ILE:O	30:A8:41:ILE:HD13	1.99	0.63
1:AA:1826:G:H2'	1:AA:1827:C:C6	2.34	0.63
1:AA:2341:G:H2'	1:AA:2342:C:C6	2.34	0.63
1:AA:1999:C:H4'	1:AA:2723:C:O2	1.99	0.63
1:AA:283:A:H4'	1:AA:284:U:OP2	1.98	0.63
1:AA:705:A:N6	1:AA:726:G:H1'	2.14	0.63
2:AB:17:C:H2'	2:AB:18:G:O4'	1.99	0.63
3:AD:92:ILE:HA	3:AD:107:ALA:H	1.62	0.63
5:AF:4:VAL:CG1	5:AF:17:ARG:HE	2.11	0.63
7:AH:86:GLU:O	7:AH:132:ARG:HA	1.99	0.63
9:AM:56:ASN:H	9:AM:126:PRO:HA	1.64	0.63
19:AT:8:ILE:HG23	19:AT:28:PHE:HD2	1.64	0.63
21:AV:62:PRO:O	21:AV:64:GLY:N	2.32	0.63
21:AV:6:LYS:O	21:AV:7:ALA:HB3	1.98	0.63
21:AV:94:GLU:HB3	21:AV:95:PRO:CD	2.29	0.63
31:BA:9:G:H2'	31:BA:10:A:H8	1.64	0.63
31:BA:1200:C:H1'	31:BA:1204:A:N6	2.14	0.63
31:BA:1472:U:O2'	31:BA:1473:A:H5'	1.99	0.63
52:BB:23:A:H2'	52:BB:24:G:O4'	1.99	0.63
33:BF:53:ALA:HB2	33:BF:115:LEU:HD21	1.81	0.63
34:BG:112:VAL:HG12	34:BG:116:GLN:OE1	1.98	0.63
34:BG:5:ILE:O	34:BG:5:ILE:HG22	1.99	0.63
34:BG:61:LYS:HZ1	34:BG:62:GLN:HE21	1.44	0.63
36:BI:14:LEU:HD21	36:BI:18:GLN:HB2	1.80	0.63
31:BA:932:C:H5''	37:BJ:3:ARG:HD2	1.80	0.63
40:BM:30:SER:HB3	40:BM:84:GLN:NE2	2.14	0.63
41:BN:30:VAL:HG21	41:BN:65:ALA:HA	1.81	0.63
31:BA:363:A:N7	42:BO:30:ALA:HB1	2.13	0.63
46:BS:53:VAL:O	46:BS:57:ARG:HG2	1.98	0.63
54:CA:1205:U:O2'	54:CA:1206:G:H5'	1.98	0.63
54:CA:502:G:OP1	42:CO:118:SER:HB2	1.99	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:517:G:O2'	54:CA:530:G:H4'	1.98	0.63
54:CA:789:U:C5	54:CA:792:A:OP2	2.52	0.63
34:CG:201:GLN:HA	34:CG:201:GLN:HE21	1.63	0.63
37:CJ:16:LEU:CD1	39:CL:42:ARG:HA	2.29	0.63
50:CW:13:LEU:HD12	50:CW:13:LEU:C	2.19	0.63
55:DA:2110:G:O2'	55:DA:2111:C:OP1	2.16	0.63
55:DA:2135:A:H3'	55:DA:2136:C:H5	1.64	0.63
55:DA:2167:U:H6	55:DA:2167:U:OP2	1.82	0.63
55:DA:2474:C:H3'	55:DA:2475:C:H6	1.63	0.63
55:DA:442:G:C4'	5:DF:46:ARG:HD3	2.29	0.63
2:DB:95:U:H3'	2:DB:95:U:C6	2.33	0.63
3:DD:65:ILE:CD1	3:DD:65:ILE:H	2.12	0.63
55:DA:444:C:C4'	5:DF:49:ALA:HB2	2.27	0.63
8:DK:82:ARG:HG3	8:DK:82:ARG:HH11	1.62	0.63
11:DO:57:THR:O	11:DO:60:MET:HB2	1.99	0.63
12:DP:90:VAL:HG13	12:DP:91:GLU:N	2.13	0.63
14:DQ:110:LEU:HA	14:DQ:112:PHE:CE1	2.33	0.63
15:DR:74:ARG:NH1	15:DR:74:ARG:HG2	2.13	0.63
57:DY:16:ASN:HA	57:DY:19:ARG:HD2	1.81	0.63
57:DY:49:ALA:HA	57:DY:84:GLU:O	1.99	0.63
57:DY:6:ASN:O	57:DY:7:VAL:C	2.36	0.63
1:AA:1341:U:H5''	19:AT:57:LEU:CG	2.28	0.63
1:AA:215:G:H4'	1:AA:216:A:O5'	1.99	0.63
1:AA:2228:G:H2'	1:AA:2229:C:C6	2.34	0.63
1:AA:2801:A:H2'	1:AA:2802:G:O4'	1.99	0.63
1:AA:557:U:H2'	1:AA:558:G:C8	2.32	0.63
2:AB:11:C:OP2	2:AB:12:C:N4	2.29	0.63
2:AB:34:U:O4	2:AB:44:G:H2'	1.99	0.63
3:AD:155:LEU:HD12	3:AD:155:LEU:N	2.13	0.63
8:AK:144:VAL:O	8:AK:145:VAL:HG22	1.98	0.63
14:AQ:106:ARG:CB	14:AQ:106:ARG:NH1	2.62	0.63
21:AV:115:GLY:O	21:AV:174:VAL:CG1	2.46	0.63
21:AV:144:LEU:HD11	21:AV:148:ASP:HA	1.81	0.63
21:AV:186:GLU:O	21:AV:187:ALA:HB2	1.99	0.63
24:AW:48:HIS:C	24:AW:48:HIS:CD2	2.72	0.63
31:BA:197:A:C6	31:BA:221:C:H4'	2.34	0.63
31:BA:457:C:H2'	31:BA:458:C:H6	1.64	0.63
33:BF:6:HIS:HD2	33:BF:7:PRO:HD2	1.62	0.63
43:BP:81:LEU:C	43:BP:83:ASP:N	2.51	0.63
54:CA:466:C:H5''	54:CA:467:G:OP2	1.99	0.63
52:CD:21:A:N3	52:CD:21:A:H3'	2.12	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:CH:102:ALA:HB1	35:CH:106:PRO:HG2	1.80	0.63
41:CN:124:LYS:HD2	41:CN:125:PHE:CZ	2.34	0.63
49:CV:65:ASN:HD22	49:CV:65:ASN:N	1.97	0.63
51:CX:10:ARG:HA	51:CX:13:ILE:HD12	1.80	0.63
16:D1:75:ASN:HB2	16:D1:78:THR:OG1	1.99	0.63
26:D4:10:VAL:CG2	26:D4:11:PRO:HD2	2.28	0.63
55:DA:1058:U:N3	55:DA:1059:G:O6	2.31	0.63
55:DA:1062:G:C2'	55:DA:1077:A:N6	2.61	0.63
55:DA:1512:G:H2'	55:DA:1513:C:O4'	1.99	0.63
55:DA:1688:U:H1'	55:DA:1701:A:C6	2.34	0.63
55:DA:242:G:H5''	30:D8:62:LEU:HD13	1.80	0.63
55:DA:2629:A:O2'	55:DA:2630:G:H5''	1.98	0.63
55:DA:2543:G:H21	55:DA:2646:C:H5''	1.64	0.63
55:DA:639:U:H2'	55:DA:640:C:H6	1.63	0.63
55:DA:649:G:H2'	55:DA:650:C:H6	1.59	0.63
3:DD:28:GLU:HB2	3:DD:29:PRO:CD	2.29	0.63
5:DF:9:ILE:HD11	5:DF:125:LEU:CG	2.28	0.63
7:DH:147:ASN:N	7:DH:147:ASN:HD22	1.96	0.63
58:DL:115:LEU:HD12	58:DL:116:ASN:N	2.14	0.63
58:DL:126:MET:H	58:DL:126:MET:CE	2.08	0.63
58:DL:143:GLU:HA	58:DL:143:GLU:OE1	1.97	0.63
58:DL:63:ARG:HE	58:DL:63:ARG:CA	2.12	0.63
12:DP:25:ASP:OD1	21:DV:78:LYS:HD3	1.99	0.63
12:DP:30:GLY:CA	12:DP:107:ALA:HB2	2.28	0.63
12:DP:87:LYS:C	12:DP:89:ASN:H	2.02	0.63
12:DP:39:PRO:HA	12:DP:97:VAL:O	1.98	0.63
14:DQ:20:ARG:HD3	14:DQ:21:THR:N	2.13	0.63
21:DV:169:GLU:OE1	21:DV:170:THR:N	2.31	0.63
1:AA:1069:A:H5'	1:AA:1070:A:C8	2.34	0.62
1:AA:329:G:O6	20:AU:19:LYS:HA	1.99	0.62
1:AA:919:G:C5'	2:AB:81:G:H1'	2.29	0.62
4:AE:71:GLY:C	4:AE:73:GLU:H	2.02	0.62
6:AG:135:LEU:N	6:AG:135:LEU:HD12	2.14	0.62
7:AH:153:LYS:HE3	7:AH:160:LYS:O	1.99	0.62
12:AP:7:MET:HB2	12:AP:10:ARG:HE	1.62	0.62
14:AQ:10:ARG:O	14:AQ:14:VAL:HG12	1.98	0.62
21:AV:106:GLY:CA	21:AV:108:PRO:HD2	2.29	0.62
24:AW:51:ARG:NH2	24:AW:55:ARG:HH12	1.97	0.62
31:BA:186:C:H1'	50:BW:81:LYS:HZ2	1.64	0.62
31:BA:920:U:H2'	31:BA:921:U:H6	1.64	0.62
32:BE:169:LYS:HD3	32:BE:169:LYS:O	1.99	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:BG:104:VAL:O	34:BG:108:LEU:HB2	1.99	0.62
54:CA:1234:C:H4'	54:CA:1364:U:C2'	2.29	0.62
54:CA:1443:G:C3'	54:CA:1446:A:H5''	2.29	0.62
54:CA:30:U:O2'	54:CA:31:G:OP1	2.16	0.62
54:CA:612:C:C2	54:CA:629:G:N2	2.67	0.62
54:CA:748:C:O2'	54:CA:749:C:P	2.57	0.62
32:CE:102:LEU:HB3	32:CE:180:LEU:CD1	2.29	0.62
32:CE:187:LEU:HD13	32:CE:187:LEU:O	1.99	0.62
34:CG:43:HIS:O	34:CG:46:LYS:HG2	1.99	0.62
35:CH:42:GLY:HA3	35:CH:66:MET:HG2	1.81	0.62
38:CK:115:SER:C	38:CK:116:LYS:HE2	2.19	0.62
39:CL:66:ARG:HH11	39:CL:66:ARG:HB3	1.64	0.62
43:CP:116:THR:HG22	43:CP:117:VAL:H	1.64	0.62
17:D2:38:LEU:HB3	17:D2:52:VAL:HG22	1.81	0.62
22:D3:51:VAL:HG23	22:D3:81:VAL:CG2	2.29	0.62
22:D3:5:LYS:O	22:D3:5:LYS:HD3	1.99	0.62
28:D6:44:ARG:O	28:D6:45:LYS:HB2	1.99	0.62
55:DA:1906:G:C8	55:DA:1929:G:N3	2.67	0.62
55:DA:704:G:HO2'	55:DA:705:A:P	2.22	0.62
3:DD:27:THR:CG2	3:DD:28:GLU:H	1.97	0.62
5:DF:128:ALA:O	5:DF:129:PHE:HB2	1.98	0.62
6:DG:16:ARG:HG2	6:DG:16:ARG:NH1	2.12	0.62
6:DG:181:ARG:HG2	6:DG:181:ARG:O	1.98	0.62
6:DG:67:LYS:CG	26:D4:5:ILE:HG22	2.26	0.62
6:DG:76:SER:OG	6:DG:83:ARG:HA	1.99	0.62
6:DG:81:LYS:HD3	6:DG:81:LYS:N	2.14	0.62
55:DA:2311:A:C8	6:DG:82:LEU:HD11	2.34	0.62
56:DI:29:GLU:N	56:DJ:2:ALA:HB1	2.13	0.62
58:DL:112:MET:SD	58:DL:120:LEU:CA	2.87	0.62
10:DN:2:ILE:HB	10:DN:33:ALA:HB3	1.80	0.62
15:DR:26:ASP:HB2	15:DR:90:GLN:O	1.98	0.62
15:DR:1:MET:O	15:DR:3:ARG:N	2.32	0.62
15:DR:51:ARG:CG	15:DR:98:LYS:HG3	2.25	0.62
21:DV:77:ASP:OD2	21:DV:80:ARG:HG3	1.98	0.62
57:DY:104:ILE:CG1	57:DY:105:PRO:CD	2.72	0.62
13:A0:62:ALA:O	13:A0:66:VAL:HG23	1.98	0.62
1:AA:2364:C:H4'	22:A3:56:ASP:OD2	1.99	0.62
1:AA:1096:A:C5	1:AA:1097:U:H1'	2.34	0.62
1:AA:1267:U:C4	1:AA:2012:G:N3	2.67	0.62
1:AA:1359:A:C5'	1:AA:1359:A:C8	2.81	0.62
1:AA:2732:G:H3'	1:AA:2733:A:C5'	2.27	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:78:A:H2'	1:AA:79:G:H8	1.64	0.62
3:AD:130:ALA:C	3:AD:131:LEU:HD12	2.19	0.62
3:AD:27:THR:O	3:AD:28:GLU:HB2	1.99	0.62
6:AG:109:VAL:O	6:AG:113:ARG:HG3	2.00	0.62
7:AH:106:THR:CG2	7:AH:112:PRO:HB3	2.27	0.62
12:AP:17:LEU:HD21	12:AP:41:TRP:HE1	1.64	0.62
14:AQ:11:LYS:HD2	14:AQ:15:ARG:NH2	2.14	0.62
21:AV:111:VAL:O	21:AV:111:VAL:HG13	1.99	0.62
21:AV:145:GLU:O	21:AV:145:GLU:CG	2.47	0.62
21:AV:175:VAL:CB	21:AV:176:PRO:CD	2.76	0.62
21:AV:179:ASP:O	21:AV:179:ASP:CG	2.37	0.62
31:BA:1077:G:N2	31:BA:1080:A:OP2	2.31	0.62
31:BA:1129:C:C5'	31:BA:1130:A:H5'	2.27	0.62
31:BA:1190:G:H3'	33:BF:3:ASN:ND2	2.14	0.62
31:BA:1316:G:H2'	31:BA:1317:C:H5''	1.81	0.62
31:BA:1470:G:O2'	31:BA:1471:G:H5'	1.99	0.62
31:BA:382:A:H2'	31:BA:383:A:H8	1.64	0.62
31:BA:440:A:H3'	31:BA:442:C:H6	1.64	0.62
52:BB:74:C:O2'	52:BB:75:C:C6	2.53	0.62
32:BE:131:PRO:O	32:BE:135:GLN:HG3	1.99	0.62
31:BA:1342:C:H1'	39:BL:124:GLN:HE22	1.64	0.62
54:CA:1002:G:C4	54:CA:1003:G:N7	2.66	0.62
54:CA:632:A:H3'	54:CA:633:G:H8	1.64	0.62
54:CA:689:C:H2'	54:CA:690:G:H5'	1.82	0.62
54:CA:1055:A:O3'	33:CF:161:GLU:OE2	2.17	0.62
34:CG:173:TRP:CZ3	34:CG:193:ASP:HB3	2.34	0.62
37:CJ:16:LEU:HD21	39:CL:45:ALA:HB2	1.80	0.62
40:CM:74:ILE:HD13	40:CM:74:ILE:H	1.63	0.62
43:CP:83:ASP:H	43:CP:93:ARG:NH2	1.96	0.62
28:D6:41:PRO:HB2	28:D6:44:ARG:NH1	2.13	0.62
55:DA:1872:A:H5'	55:DA:1878:G:OP2	1.99	0.62
55:DA:2789:C:O2'	55:DA:2790:A:C4'	2.48	0.62
55:DA:74:A:O2'	55:DA:75:G:OP2	2.18	0.62
3:DD:35:LYS:CE	3:DD:104:TYR:HB2	2.29	0.62
4:DE:51:PHE:HD1	4:DE:52:LEU:HG	1.62	0.62
7:DH:136:ILE:O	7:DH:136:ILE:HG22	1.97	0.62
58:DL:7:VAL:CG1	58:DL:58:THR:CA	2.75	0.62
11:DO:50:ARG:HH21	11:DO:50:ARG:HB3	1.64	0.62
57:DY:9:LEU:CD2	57:DY:10:LEU:N	2.62	0.62
57:DY:51:LEU:CD2	57:DY:82:PHE:CA	2.72	0.62
16:A1:61:TRP:O	16:A1:65:ILE:HD13	1.99	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:A2:35:LEU:O	17:A2:37:VAL:HG22	1.97	0.62
26:A4:60:GLN:NE2	26:A4:60:GLN:N	2.42	0.62
1:AA:1396:U:H2'	1:AA:1396:U:O2	1.99	0.62
1:AA:1267:U:O4	1:AA:2012:G:C4	2.52	0.62
1:AA:2275:C:O2	12:AP:83:MET:HG3	2.00	0.62
1:AA:2749:A:H5''	7:AH:6:ARG:NH1	2.14	0.62
1:AA:2857:G:N2	1:AA:2859:G:H3'	2.14	0.62
2:AB:104:A:H2'	2:AB:105:G:O4'	1.98	0.62
5:AF:179:GLU:N	5:AF:179:GLU:OE1	2.31	0.62
15:AR:94:ALA:C	15:AR:96:ARG:H	2.01	0.62
20:AU:8:LYS:O	20:AU:27:VAL:CG2	2.48	0.62
24:AW:24:LEU:HD22	24:AW:60:LEU:CD2	2.29	0.62
31:BA:1003:G:C3'	31:BA:1004:A:H5''	2.28	0.62
31:BA:1253:G:H2'	31:BA:1254:C:C6	2.34	0.62
31:BA:1327:C:O2'	31:BA:1328:C:H5'	2.00	0.62
31:BA:1348:U:N3	31:BA:1374:A:H2	1.97	0.62
31:BA:411:A:H62	31:BA:413:G:H21	1.47	0.62
32:BE:172:ILE:HD12	32:BE:172:ILE:N	2.13	0.62
33:BF:71:ALA:HA	33:BF:106:VAL:HB	1.80	0.62
35:BH:102:ALA:HB1	35:BH:106:PRO:HG2	1.81	0.62
39:BL:113:LYS:HD2	39:BL:113:LYS:H	1.64	0.62
53:C1:30:C:C4	53:C1:31:A:N7	2.66	0.62
54:CA:575:G:N2	54:CA:576:G:N7	2.46	0.62
39:CL:48:GLU:HB2	39:CL:78:LYS:HE3	1.80	0.62
43:CP:119:GLY:C	43:CP:120:LYS:HD3	2.19	0.62
54:CA:1245:A:P	51:CX:9:ARG:HH22	2.22	0.62
43:CP:65:LYS:HB3	26:D4:50:VAL:HG21	1.81	0.62
28:D6:20:ASN:ND2	28:D6:42:TRP:CH2	2.68	0.62
55:DA:1286:A:H2'	55:DA:1288:U:OP2	1.99	0.62
55:DA:1652:A:H4'	55:DA:1653:G:OP1	1.99	0.62
55:DA:1885:A:H3'	55:DA:1886:C:H6	1.63	0.62
55:DA:215:G:H4'	55:DA:216:A:O5'	1.99	0.62
55:DA:655:A:H2'	55:DA:656:G:H5'	1.81	0.62
55:DA:83:G:HO2'	55:DA:84:A:H8	1.46	0.62
3:DD:147:LEU:HD11	3:DD:183:ARG:NH1	2.14	0.62
4:DE:38:THR:OG1	4:DE:39:PRO:HD2	1.98	0.62
4:DE:39:PRO:HG2	4:DE:40:GLU:OE2	2.00	0.62
56:DJ:16:THR:OG1	56:DJ:17:VAL:HG23	1.99	0.62
58:DL:106:GLU:HA	58:DL:109:LYS:HB2	1.80	0.62
12:DP:109:VAL:CG1	12:DP:110:THR:N	2.62	0.62
18:DS:82:LEU:HB2	18:DS:98:LYS:HB2	1.81	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:71:A:H2	19:DT:31:HIS:CE1	2.17	0.62
20:DU:47:LYS:O	20:DU:49:VAL:HG23	1.99	0.62
57:DY:48:GLY:O	57:DY:50:ARG:HB3	2.00	0.62
28:A6:25:LYS:HB3	30:A8:34:TRP:CH2	2.34	0.62
1:AA:1043:C:H2'	1:AA:1044:G:H5''	1.80	0.62
1:AA:1406:U:H3'	1:AA:1407:C:C6	2.34	0.62
1:AA:2394:C:OP1	11:AO:63:PRO:CD	2.44	0.62
2:AB:44:G:C2	2:AB:48:A:C2	2.87	0.62
2:AB:50:G:P	14:AQ:62:LYS:HB2	2.39	0.62
6:AG:81:LYS:N	6:AG:81:LYS:HD3	2.13	0.62
31:BA:1510:U:H2'	31:BA:1511:G:C8	2.34	0.62
31:BA:243:A:H5'	31:BA:245:C:OP1	1.99	0.62
31:BA:961:U:O2	31:BA:1201:A:N1	2.32	0.62
33:BF:35:GLU:O	33:BF:38:ARG:HG2	2.00	0.62
34:BG:3:ARG:HE	34:BG:118:ARG:HD3	1.64	0.62
36:BI:6:VAL:HG12	36:BI:8:ILE:CD1	2.28	0.62
41:BN:99:GLN:OE1	41:BN:105:VAL:HG21	1.99	0.62
54:CA:1399:C:H4'	54:CA:1400:C:C5'	2.29	0.62
54:CA:1464:G:OP1	15:DR:108:ARG:HD2	1.99	0.62
54:CA:595:G:H5''	54:CA:596:C:OP1	1.99	0.62
32:CE:71:VAL:CG2	32:CE:164:VAL:HG22	2.29	0.62
32:CE:224:GLN:C	32:CE:226:ARG:H	2.03	0.62
42:CO:47:LYS:CB	42:CO:48:PRO:CD	2.76	0.62
48:CU:56:THR:HB	48:CU:58:LEU:HD13	1.81	0.62
13:D0:91:GLN:HE21	13:D0:91:GLN:N	1.96	0.62
26:D4:10:VAL:HG23	26:D4:11:PRO:HD2	1.80	0.62
26:D4:14:ILE:CG2	26:D4:21:VAL:HB	2.29	0.62
55:DA:322:A:H4'	55:DA:323:G:OP2	1.99	0.62
2:DB:81:G:C2	2:DB:82:G:C5	2.87	0.62
4:DE:119:ARG:HD3	4:DE:160:TYR:HB2	1.82	0.62
5:DF:65:TRP:HB2	5:DF:66:PRO:HD2	1.80	0.62
56:DJ:1:MET:CG	56:DJ:2:ALA:N	2.62	0.62
9:DM:30:ILE:HG22	9:DM:34:LEU:CD2	2.29	0.62
10:DN:7:TYR:C	10:DN:8:LEU:HD22	2.19	0.62
57:DY:27:VAL:CG2	57:DY:28:ASN:H	1.83	0.62
23:DZ:83:GLU:HG2	23:DZ:84:GLY:N	2.12	0.62
28:A6:9:LEU:HD22	28:A6:11:LEU:CD2	2.29	0.62
28:A6:31:PRO:C	28:A6:33:LYS:H	2.02	0.62
1:AA:2344:U:C2	28:A6:37:ARG:HD3	2.34	0.62
1:AA:1111:A:H5'	7:AH:3:ARG:CZ	2.29	0.62
1:AA:1252:G:O4'	16:A1:33:ARG:HD2	1.99	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:30:G:H2'	1:AA:31:C:C6	2.35	0.62
1:AA:320:A:H4'	1:AA:322:A:C8	2.35	0.62
1:AA:442:G:O4'	5:AF:46:ARG:HD3	2.00	0.62
5:AF:29:ASN:N	5:AF:112:MET:HE1	2.15	0.62
9:AM:126:PRO:O	9:AM:127:ASP:HB2	1.98	0.62
12:AP:54:MET:HE1	12:AP:118:LEU:HD22	1.80	0.62
21:AV:105:VAL:O	21:AV:140:ASP:HA	1.99	0.62
24:AW:54:LYS:O	24:AW:57:ILE:HG12	2.00	0.62
31:BA:1004:A:O4'	31:BA:1036:G:C6	2.52	0.62
31:BA:1529:G:H5'	31:BA:1530:G:P	2.40	0.62
31:BA:752:G:H1'	31:BA:754:C:H41	1.65	0.62
52:BD:35:A:H62	52:BD:37:MIA:H153	1.63	0.62
41:BN:41:THR:HG21	41:BN:71:LYS:HB2	1.81	0.62
50:BW:73:HIS:O	50:BW:76:ALA:HB3	2.00	0.62
54:CA:1095:U:P	54:CA:1108:G:H1	2.22	0.62
54:CA:148:G:H2'	54:CA:149:A:H8	1.64	0.62
54:CA:511:C:H1'	34:CG:43:HIS:NE2	2.13	0.62
54:CA:738:C:H2'	54:CA:739:C:C6	2.34	0.62
34:CG:119:GLN:HG3	34:CG:123:HIS:HD2	1.62	0.62
40:CM:64:GLU:HG2	44:CQ:59:ALA:HB2	1.82	0.62
40:CM:7:LYS:HG2	40:CM:71:LEU:HD13	1.81	0.62
47:CT:52:LYS:HD2	47:CT:55:ASP:OD1	1.99	0.62
16:D1:85:LYS:NZ	16:D1:117:GLN:HG2	2.14	0.62
55:DA:1249:U:O2	55:DA:1249:U:C2'	2.37	0.62
55:DA:1828:G:H8	55:DA:1828:G:OP2	1.82	0.62
55:DA:2051:A:N6	55:DA:2614:A:H2'	2.14	0.62
2:DB:21:G:H8	2:DB:21:G:H5'	1.64	0.62
3:DD:183:ARG:HG2	3:DD:183:ARG:NH1	2.12	0.62
4:DE:101:ARG:NH1	4:DE:171:GLU:HB2	2.15	0.62
58:DL:34:ILE:HD12	58:DL:37:PHE:O	1.98	0.62
9:DM:65:LYS:CB	9:DM:69:GLN:HE21	2.10	0.62
10:DN:53:LYS:HD2	10:DN:53:LYS:N	2.13	0.62
14:DQ:86:ALA:O	14:DQ:87:PHE:HB3	1.99	0.62
15:DR:95:ARG:HH11	15:DR:95:ARG:HG3	1.64	0.62
57:DY:130:THR:C	57:DY:132:ASP:H	2.03	0.62
57:DY:58:LEU:CA	57:DY:62:ALA:HB2	2.29	0.62
16:A1:90:VAL:HG22	17:A2:39:LEU:HD23	1.82	0.62
26:A4:58:ARG:HA	26:A4:61:ARG:HB2	1.81	0.62
1:AA:1669:A:C8	10:AN:5:GLN:HG2	2.33	0.62
1:AA:310:A:O2'	1:AA:311:A:H3'	1.99	0.62
1:AA:9:U:H3	1:AA:2629:A:N6	1.98	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:AD:10:THR:HG23	3:AD:13:ARG:CB	2.29	0.62
8:AK:5:LEU:HD11	8:AK:19:VAL:CG1	2.25	0.62
12:AP:4:PRO:HB2	12:AP:10:ARG:NH2	2.14	0.62
1:AA:851:U:H5'	25:AX:49:LYS:HD2	1.82	0.62
23:AZ:87:PRO:CA	23:AZ:90:ILE:HG22	2.22	0.62
23:AZ:91:LYS:O	23:AZ:92:LYS:C	2.38	0.62
31:BA:1053:G:C5'	31:BA:1054:C:H5'	2.26	0.62
31:BA:1216:G:H2'	31:BA:1217:C:H6	1.63	0.62
31:BA:1306:A:H2'	31:BA:1307:U:O4'	2.00	0.62
31:BA:1350:A:H2'	31:BA:1351:U:O4'	2.00	0.62
31:BA:421:U:H2'	31:BA:421:U:O2	1.99	0.62
32:BE:89:GLY:O	32:BE:154:LEU:HD13	1.99	0.62
34:BG:61:LYS:NZ	34:BG:62:GLN:NE2	2.47	0.62
37:BJ:69:VAL:HG12	37:BJ:69:VAL:O	1.99	0.62
42:BO:70:ILE:HD12	42:BO:77:LEU:HD12	1.80	0.62
46:BS:1:MET:HG3	46:BS:1:MET:O	1.99	0.62
47:BT:34:LYS:O	47:BT:36:ILE:HG23	1.99	0.62
49:BV:51:VAL:HB	49:BV:75:ALA:HB2	1.82	0.62
54:CA:628:G:C2	54:CA:629:G:C4	2.88	0.62
54:CA:693:G:H2'	54:CA:694:A:C8	2.34	0.62
54:CA:818:G:H3'	54:CA:819:A:H5'	1.81	0.62
52:CB:56:C:C6	55:DA:896:A:O2'	2.51	0.62
52:CC:46:G:O3'	52:CC:47:U:H4'	2.00	0.62
34:CG:79:PHE:CD2	34:CG:79:PHE:C	2.70	0.62
37:CJ:113:GLU:CG	37:CJ:119:ARG:HG2	2.29	0.62
55:DA:1027:A:N6	55:DA:1126:A:H1'	2.13	0.62
55:DA:1078:U:C1'	55:DA:1088:A:C2	2.82	0.62
55:DA:1099:G:H8	55:DA:1099:G:H5'	1.64	0.62
55:DA:1181:C:H5'	55:DA:1181:C:H6	1.63	0.62
55:DA:1427:A:O2'	55:DA:1428:C:OP2	2.14	0.62
55:DA:222:A:N6	55:DA:232:G:O2'	2.32	0.62
55:DA:2657:A:H1'	55:DA:2665:A:N6	2.14	0.62
55:DA:528:A:C2'	55:DA:529:A:H5''	2.29	0.62
55:DA:888:C:C2'	55:DA:889:C:H5'	2.30	0.62
58:DL:138:VAL:HG12	58:DL:139:VAL:H	1.62	0.62
58:DL:93:ARG:HG2	58:DL:135:GLY:HA2	1.81	0.62
55:DA:389:G:H22	11:DO:72:PRO:CD	2.13	0.62
14:DQ:70:GLY:O	14:DQ:73:LEU:HB3	1.99	0.62
20:DU:88:LYS:HB3	20:DU:90:LEU:CD2	2.29	0.62
21:DV:194:PRO:HB2	21:DV:196:VAL:HG11	1.80	0.62
1:AA:2002:G:OP1	13:A0:9:LYS:HE2	2.00	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:A1:88:ILE:C	16:A1:90:VAL:H	2.01	0.62
27:A5:40:LYS:HD2	27:A5:46:CYS:HB2	1.82	0.62
28:A6:10:LEU:C	28:A6:11:LEU:HD22	2.19	0.62
1:AA:1726:G:O2'	1:AA:1727:U:H5'	2.00	0.62
1:AA:171:G:H2'	1:AA:172:C:C6	2.34	0.62
1:AA:1784:A:H4'	1:AA:1785:A:C5'	2.29	0.62
1:AA:2065:C:H2'	1:AA:2066:C:C6	2.35	0.62
1:AA:266:G:C3'	1:AA:267:C:H5''	2.30	0.62
1:AA:2735:G:N2	1:AA:2770:G:H1'	2.13	0.62
1:AA:2820:A:N3	13:A0:4:LEU:CD2	2.63	0.62
4:AE:200:GLU:CG	4:AE:201:THR:N	2.62	0.62
6:AG:41:GLN:HE21	6:AG:60:LEU:HD12	1.64	0.62
6:AG:69:ALA:HB3	6:AG:91:ARG:HH21	1.65	0.62
7:AH:105:LEU:O	7:AH:107:VAL:HG13	1.99	0.62
7:AH:168:PRO:HG2	7:AH:169:VAL:H	1.65	0.62
7:AH:7:LEU:HD12	7:AH:8:PRO:N	2.15	0.62
7:AH:85:LYS:HA	7:AH:85:LYS:HE2	1.80	0.62
10:AN:87:ILE:HG22	10:AN:88:ASN:O	1.99	0.62
14:AQ:62:LYS:HB3	14:AQ:97:ARG:HD3	1.81	0.62
19:AT:50:LYS:H	19:AT:87:GLN:HE22	1.48	0.62
21:AV:57:ILE:N	21:AV:57:ILE:HD12	2.14	0.62
31:BA:1382:C:O2'	31:BA:1383:C:H5'	2.00	0.62
35:BH:80:ILE:HG22	38:BK:104:ARG:NE	2.15	0.62
36:BI:30:LEU:HD23	36:BI:75:LEU:HD11	1.80	0.62
39:BL:78:LYS:HB2	39:BL:78:LYS:NZ	2.14	0.62
42:BO:7:ILE:O	42:BO:10:LEU:N	2.33	0.62
54:CA:1443:G:H4'	54:CA:1446:A:OP2	1.99	0.62
54:CA:422:C:O2'	54:CA:423:G:H5''	2.00	0.62
54:CA:552:U:O2'	54:CA:553:A:H5'	2.00	0.62
54:CA:628:G:N2	54:CA:629:G:C2	2.67	0.62
32:CE:201:ILE:HG21	32:CE:214:ILE:HG21	1.81	0.62
33:CF:175:LEU:HD12	33:CF:175:LEU:N	2.14	0.62
36:CI:41:GLU:O	36:CI:43:LEU:HD12	1.99	0.62
36:CI:72:VAL:HG13	36:CI:73:ASN:N	2.14	0.62
43:CP:3:ARG:NH2	6:DG:139:LEU:HD13	2.15	0.62
48:CU:26:LEU:HD22	48:CU:42:ARG:CZ	2.30	0.62
50:CW:67:ALA:HA	50:CW:72:LEU:O	2.00	0.62
27:D5:51:TYR:CB	27:D5:56:LYS:HB3	2.30	0.62
55:DA:2158:A:H5''	55:DA:2159:G:OP1	2.00	0.62
55:DA:230:U:OP2	55:DA:230:U:H6	1.82	0.62
55:DA:2392:A:H2	55:DA:2424:C:N4	1.96	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:546:C:H3'	55:DA:547:A:C8	2.35	0.62
55:DA:718:A:H3'	55:DA:719:C:H6	1.64	0.62
5:DF:198:ALA:HA	5:DF:201:VAL:CG1	2.28	0.62
7:DH:86:GLU:O	7:DH:87:LEU:HB2	2.00	0.62
56:DJ:12:LEU:H	56:DJ:13:SER:CA	2.13	0.62
14:DQ:56:LEU:O	14:DQ:58:LEU:HD22	1.98	0.62
14:DQ:99:LYS:O	14:DQ:102:ALA:HB3	1.99	0.62
21:DV:30:ASN:OD1	21:DV:33:LEU:HB3	1.99	0.62
57:DY:104:ILE:HG23	57:DY:105:PRO:CD	2.29	0.62
1:AA:1278:A:H5''	13:A0:36:THR:HG22	1.81	0.62
26:A4:63:TYR:HE1	49:BV:39:THR:CG2	2.13	0.62
1:AA:208:C:H2'	1:AA:209:C:H6	1.65	0.62
1:AA:2340:G:O2'	1:AA:2341:G:H5'	2.00	0.62
1:AA:2478:A:H5'	1:AA:2479:G:OP2	1.99	0.62
1:AA:2698:U:H2'	1:AA:2699:C:C6	2.34	0.62
1:AA:5:A:O2'	1:AA:6:A:H5'	2.00	0.62
1:AA:768:G:O2'	1:AA:1379:A:N6	2.31	0.62
1:AA:884:C:O5'	1:AA:884:C:H6	1.83	0.62
4:AE:73:GLU:CG	4:AE:74:PRO:HD2	2.29	0.62
5:AF:136:THR:O	5:AF:140:LEU:HB2	1.99	0.62
6:AG:61:ALA:HB2	6:AG:68:PRO:CD	2.29	0.62
7:AH:106:THR:HG22	7:AH:112:PRO:CB	2.26	0.62
7:AH:18:GLU:HB2	7:AH:25:LYS:HB2	1.80	0.62
7:AH:18:GLU:CG	7:AH:25:LYS:HB2	2.29	0.62
11:AO:102:ARG:O	11:AO:102:ARG:HD2	2.00	0.62
12:AP:12:GLN:HG2	12:AP:73:PRO:HD2	1.81	0.62
14:AQ:70:GLY:HA3	14:AQ:104:GLY:HA3	1.80	0.62
52:BB:19:G:H1'	52:BB:57:G:C2	2.35	0.62
33:BF:156:ARG:HH21	33:BF:161:GLU:HA	1.64	0.62
39:BL:17:VAL:CG1	39:BL:81:ILE:HD13	2.29	0.62
41:BN:91:ARG:HG2	41:BN:91:ARG:HH11	1.64	0.62
43:BP:14:ARG:NH2	43:BP:16:ASP:OD1	2.33	0.62
49:BV:12:ASP:O	49:BV:16:LEU:HD13	1.99	0.62
50:BW:46:GLU:HG2	50:BW:46:GLU:O	2.00	0.62
54:CA:37:U:H2'	54:CA:38:G:H8	1.65	0.62
54:CA:69:G:N2	54:CA:101:A:C2	2.68	0.62
52:CB:76:A:H8	55:DA:2507:C:O4'	1.82	0.62
32:CE:69:LEU:HD12	32:CE:91:PRO:O	1.99	0.62
37:CJ:121:ALA:O	37:CJ:125:MET:HG3	1.99	0.62
47:CT:57:VAL:HG12	47:CT:76:LEU:HA	1.80	0.62
55:DA:2840:C:O3'	13:D0:53:HIS:CE1	2.53	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:D8:17:THR:HG23	30:D8:21:LYS:O	2.00	0.62
30:D8:6:THR:O	30:D8:7:HIS:HB2	2.00	0.62
55:DA:1057:A:O2'	55:DA:1058:U:O5'	2.17	0.62
55:DA:1332:G:H21	55:DA:1610:A:H8	1.47	0.62
55:DA:2733:A:H2'	55:DA:2734:A:O4'	1.99	0.62
6:DG:97:ASP:O	6:DG:100:TRP:N	2.32	0.62
56:DI:29:GLU:N	56:DJ:2:ALA:CB	2.62	0.62
8:DK:25:TYR:CE2	8:DK:29:TYR:HD2	2.18	0.62
58:DL:11:GLN:HA	58:DL:23:VAL:HG12	1.82	0.62
57:DY:57:THR:O	57:DY:58:LEU:O	2.17	0.62
1:AA:1924:C:H2'	1:AA:1925:C:O4'	1.99	0.62
1:AA:2162:G:O2'	1:AA:2163:C:H5'	2.00	0.62
1:AA:2789:C:O2'	1:AA:2790:A:C4'	2.48	0.62
1:AA:2820:A:H61	4:AE:192:ASN:N	1.93	0.62
4:AE:8:LYS:HE3	4:AE:188:VAL:CG1	2.27	0.62
8:AK:82:ARG:HD2	8:AK:146:ALA:HB2	1.81	0.62
11:AO:56:SER:O	11:AO:57:THR:HB	1.99	0.62
20:AU:5:MET:CE	20:AU:32:PRO:HB3	2.30	0.62
31:BA:1240:U:C4	37:BJ:32:ARG:HD2	2.35	0.62
31:BA:1347:G:O2'	31:BA:1348:U:P	2.57	0.62
31:BA:792:A:O2'	31:BA:793:U:OP2	2.18	0.62
31:BA:8:A:H1'	35:BH:102:ALA:N	2.15	0.62
36:BI:99:ALA:H	48:BU:31:LEU:HD22	1.65	0.62
39:BL:112:LYS:HD3	39:BL:112:LYS:C	2.20	0.62
46:BS:49:LEU:HD13	46:BS:73:LEU:HD22	1.81	0.62
47:BT:7:THR:O	47:BT:23:VAL:HG13	1.99	0.62
48:BU:78:LEU:O	48:BU:79:LEU:HD23	2.00	0.62
49:BV:80:TYR:CZ	49:BV:82:GLY:HA2	2.35	0.62
54:CA:1189:C:OP1	40:CM:51:ARG:NH2	2.26	0.62
54:CA:266:G:O2'	54:CA:267:C:OP2	2.13	0.62
34:CG:13:ARG:HB3	34:CG:33:MET:HG2	1.82	0.62
46:CS:20:VAL:HG23	46:CS:34:GLU:O	2.00	0.62
49:CV:49:ILE:O	49:CV:60:VAL:HG22	1.98	0.62
16:D1:74:LEU:HD13	16:D1:75:ASN:O	2.00	0.62
17:D2:55:ALA:HB1	17:D2:101:GLY:HA2	1.82	0.62
28:D6:33:LYS:HG3	28:D6:34:LEU:HD13	1.81	0.62
55:DA:1644:C:O2	55:DA:1644:C:H2'	1.99	0.62
55:DA:1728:G:C6	55:DA:1730:U:OP2	2.53	0.62
55:DA:2197:U:O2'	55:DA:2198:A:C8	2.52	0.62
55:DA:2391:G:H1'	55:DA:2429:G:H21	1.64	0.62
55:DA:2754:U:C5'	55:DA:2755:C:OP2	2.48	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2790:A:O2'	55:DA:2791:C:OP2	2.17	0.62
55:DA:27:G:N2	55:DA:512:G:O2'	2.33	0.62
55:DA:609(A):G:H2'	55:DA:610:C:C6	2.34	0.62
55:DA:654(C):G:C2	55:DA:654(S):G:C2	2.88	0.62
3:DD:36:PRO:O	3:DD:37:LEU:HD23	2.00	0.62
4:DE:95:ILE:CD1	4:DE:95:ILE:H	2.12	0.62
6:DG:101:ILE:CD1	26:D4:9:LEU:HD11	2.30	0.62
6:DG:41:GLN:HB3	6:DG:43:LEU:CD1	2.30	0.62
58:DL:36:GLU:O	58:DL:36:GLU:HG3	1.98	0.62
58:DL:18:THR:HG21	58:DL:38:VAL:HG11	1.80	0.62
58:DL:83:GLY:O	58:DL:84:LEU:C	2.37	0.62
55:DA:1012:U:C5	9:DM:28:THR:HG21	2.34	0.62
14:DQ:41:ASP:OD2	14:DQ:44:LYS:HB2	2.00	0.62
20:DU:57:GLN:HE21	20:DU:57:GLN:C	2.02	0.62
21:DV:152:ALA:HB1	21:DV:163:LEU:CD1	2.30	0.62
21:DV:190:GLU:C	21:DV:191:VAL:CG2	2.61	0.62
1:AA:1353:A:H4'	3:AD:38:LYS:HZ2	1.65	0.62
1:AA:1668:A:N6	1:AA:1676:A:H61	1.98	0.62
1:AA:1996:C:H5	10:AN:32:TYR:HH	1.48	0.62
1:AA:2009:G:O2'	1:AA:2010:G:H5'	2.00	0.62
1:AA:2273:A:O2'	1:AA:2274:A:H5'	2.00	0.62
1:AA:2328:A:H2'	1:AA:2329:G:C8	2.35	0.62
1:AA:2505:G:O2'	1:AA:2506:U:H6	1.83	0.62
1:AA:1966:A:H1'	1:AA:2593:U:H5'	1.82	0.62
1:AA:589:C:H2'	1:AA:590:A:H8	1.62	0.62
1:AA:877:U:H4'	1:AA:878:A:OP1	2.00	0.62
5:AF:63:LYS:CE	5:AF:67:GLN:HB2	2.29	0.62
8:AK:103:ARG:HH11	8:AK:103:ARG:HG2	1.65	0.62
9:AM:30:ILE:HG22	9:AM:34:LEU:HD21	1.82	0.62
9:AM:9:VAL:HG11	9:AM:39:ARG:HH12	1.65	0.62
11:AO:92:GLU:HA	11:AO:123:LEU:HD13	1.80	0.62
14:AQ:106:ARG:HH11	14:AQ:106:ARG:HB3	1.63	0.62
21:AV:114:GLY:C	21:AV:177:PRO:CB	2.68	0.62
1:AA:896:A:H1'	21:AV:176:PRO:CG	2.29	0.62
24:AW:70:GLN:HG2	24:AW:71:ASN:N	2.10	0.62
31:BA:1423:G:H2'	31:BA:1424:C:C6	2.35	0.62
31:BA:585:G:H4'	42:BO:8:ASN:ND2	2.12	0.62
31:BA:704:A:H5'	31:BA:705:U:OP2	2.00	0.62
31:BA:742:G:OP2	45:BR:35:ARG:NH2	2.32	0.62
31:BA:968:A:C4'	31:BA:969:A:OP2	2.48	0.62
52:BD:57:G:H2'	52:BD:58:A:H5''	1.82	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:8:A:H1'	35:BH:102:ALA:C	2.20	0.62
38:BK:116:LYS:HD2	38:BK:129:VAL:HG11	1.81	0.62
38:BK:77:GLU:HG2	38:BK:78:GLN:H	1.64	0.62
39:BL:127:LYS:HG3	39:BL:127:LYS:O	1.99	0.62
48:BU:22:VAL:HA	48:BU:25:THR:OG1	2.00	0.62
54:CA:1004:A:H2'	54:CA:1005:A:O4'	1.99	0.62
54:CA:1055:A:N6	54:CA:1200:C:N3	2.48	0.62
54:CA:1067:A:H1'	54:CA:1068:G:O4'	2.00	0.62
54:CA:853:G:O2'	54:CA:854:G:H5'	2.00	0.62
32:CE:97:TRP:CH2	32:CE:173:ALA:HA	2.35	0.62
33:CF:52:LEU:H	33:CF:52:LEU:HD23	1.63	0.62
34:CG:52:SER:H	34:CG:55:ALA:HB3	1.65	0.62
42:CO:55:VAL:HG12	42:CO:56:ALA:N	2.14	0.62
55:DA:1538:G:O5'	55:DA:1538:G:H8	1.83	0.62
55:DA:1820:U:C4'	55:DA:1821:A:OP2	2.48	0.62
55:DA:2031:A:C6	55:DA:2498:C:H1'	2.35	0.62
55:DA:386:G:H3'	55:DA:388:G:H22	1.64	0.62
55:DA:885:C:C2	55:DA:890:A:N6	2.68	0.62
55:DA:890:A:C8	55:DA:892:G:C8	2.87	0.62
8:DK:133:HIS:O	8:DK:134:PRO:C	2.38	0.62
12:DP:21:THR:O	12:DP:22:LYS:O	2.18	0.62
20:DU:50:ARG:HB3	20:DU:53:PRO:HG2	1.82	0.62
21:DV:9:TYR:CE2	21:DV:61:LEU:CD2	2.82	0.62
57:DY:26:LEU:O	57:DY:111:LEU:CD1	2.48	0.62
57:DY:25:PHE:CZ	57:DY:82:PHE:CG	2.82	0.62
1:AA:449:A:H4'	16:A1:3:ARG:HH11	1.65	0.61
1:AA:1020:A:N1	1:AA:1141:U:H2'	2.14	0.61
1:AA:1190:G:H5'	11:AO:32:THR:HA	1.82	0.61
1:AA:207:A:H2'	1:AA:208:C:O4'	2.00	0.61
1:AA:2425:A:H5''	1:AA:2426:A:H3'	1.82	0.61
1:AA:2746:U:O4	1:AA:2756:U:O4	2.18	0.61
4:AE:23:VAL:HG23	4:AE:24:THR:H	1.64	0.61
1:AA:2531:A:H4'	7:AH:157:TYR:HE2	1.62	0.61
7:AH:89:ILE:HD13	7:AH:90:LYS:H	1.63	0.61
14:AQ:14:VAL:HG21	14:AQ:89:ARG:NH1	2.14	0.61
20:AU:95:LYS:HB2	20:AU:95:LYS:NZ	2.14	0.61
21:AV:145:GLU:OE1	21:AV:145:GLU:C	2.38	0.61
24:AW:51:ARG:HB2	24:AW:55:ARG:HH11	1.62	0.61
23:AZ:91:LYS:HG3	23:AZ:92:LYS:N	2.15	0.61
31:BA:818:G:H3'	31:BA:819:A:C5'	2.29	0.61
52:BC:28:G:H2'	52:BC:29:G:H5'	1.82	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:BE:97:TRP:CZ2	32:BE:102:LEU:HD13	2.35	0.61
33:BF:64:VAL:HG23	33:BF:97:LYS:HE3	1.82	0.61
34:BG:141:ARG:N	34:BG:144:ASP:OD2	2.33	0.61
34:BG:187:ARG:HG2	34:BG:188:LEU:N	2.14	0.61
35:BH:127:ASN:O	35:BH:131:ILE:HG12	1.99	0.61
41:BN:54:ARG:HG2	41:BN:54:ARG:HH11	1.64	0.61
49:BV:64:GLU:O	49:BV:67:VAL:HG23	2.00	0.61
50:BW:33:ILE:HD13	50:BW:62:LEU:HB3	1.82	0.61
54:CA:161:A:H2'	54:CA:162:A:C8	2.35	0.61
54:CA:366:C:O2'	54:CA:394:G:N2	2.33	0.61
54:CA:678:U:H2'	54:CA:679:C:H6	1.64	0.61
54:CA:806:C:O2'	54:CA:807:A:H5'	1.99	0.61
34:CG:12:CYS:HA	34:CG:19:LEU:HD23	1.79	0.61
41:CN:79:SER:HB2	41:CN:106:LYS:CD	2.27	0.61
42:CO:25:PRO:C	42:CO:27:LEU:H	2.01	0.61
43:CP:3:ARG:HG2	43:CP:9:ILE:HG12	1.82	0.61
45:CR:39:LEU:HD13	45:CR:56:LEU:HB2	1.82	0.61
45:CR:8:LYS:NZ	45:CR:8:LYS:HB2	2.15	0.61
17:D2:66:ARG:NH1	17:D2:88:ARG:HD3	2.15	0.61
55:DA:1188:U:H5'	17:D2:79:VAL:HG22	1.82	0.61
55:DA:1059:G:OP1	58:DL:4:VAL:CG1	2.48	0.61
55:DA:1509:C:H3'	55:DA:1510:A:H4'	1.82	0.61
55:DA:2175:C:H2'	55:DA:2176:A:H5''	1.80	0.61
55:DA:2245:U:H5'	55:DA:2246:G:H5'	1.81	0.61
55:DA:2318:G:N2	14:DQ:2:ALA:N	2.48	0.61
55:DA:27:G:O2'	55:DA:28:A:C8	2.53	0.61
55:DA:556:G:H2'	55:DA:557:U:C6	2.35	0.61
55:DA:83:G:O2'	55:DA:84:A:H8	1.82	0.61
55:DA:860:U:O2	55:DA:860:U:O4'	2.17	0.61
2:DB:69:G:H2'	2:DB:70:C:H6	1.65	0.61
3:DD:35:LYS:HG2	3:DD:64:ILE:HG23	1.81	0.61
6:DG:98:ARG:O	6:DG:101:ILE:HG12	2.00	0.61
6:DG:94:LEU:HD23	6:DG:94:LEU:N	2.14	0.61
58:DL:108:ALA:HA	58:DL:111:LYS:HD3	1.82	0.61
58:DL:53:VAL:CG1	58:DL:72:PRO:HB2	2.30	0.61
9:DM:68:GLU:HG2	9:DM:88:GLU:CD	2.20	0.61
11:DO:61:ARG:O	11:DO:62:LEU:CD2	2.46	0.61
12:DP:90:VAL:O	12:DP:91:GLU:C	2.38	0.61
20:DU:81:LYS:HD3	20:DU:97:ARG:HE	1.63	0.61
57:DY:138:LEU:CG	57:DY:139:VAL:N	2.63	0.61
57:DY:49:ALA:HA	57:DY:84:GLU:H	1.65	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:686:G:H1'	29:A7:6:GLN:O	1.99	0.61
1:AA:1005:C:H2'	1:AA:1006:C:H6	1.63	0.61
1:AA:196:A:H5'	1:AA:197:A:OP2	1.99	0.61
1:AA:2102:U:H2'	1:AA:2103:C:C6	2.35	0.61
1:AA:2306:C:C3'	1:AA:2307:G:H5''	2.19	0.61
1:AA:2311:A:OP1	1:AA:2312:U:O4	2.19	0.61
1:AA:2439:A:O2'	1:AA:2440:C:OP2	2.19	0.61
1:AA:2712:U:HO2'	1:AA:2712(A):A:H8	1.42	0.61
1:AA:894:C:C5	1:AA:895:U:C5	2.88	0.61
1:AA:897:C:OP2	1:AA:897:C:O4'	2.18	0.61
1:AA:944:G:H2'	1:AA:944:G:N3	2.13	0.61
3:AD:165:ILE:HA	3:AD:175:LEU:HD23	1.81	0.61
4:AE:102:VAL:HB	4:AE:199:ARG:O	2.00	0.61
5:AF:170:LEU:HD23	5:AF:172:TRP:HE1	1.65	0.61
5:AF:89:VAL:HG12	5:AF:90:PHE:H	1.64	0.61
15:AR:3:ARG:HG2	15:AR:6:LEU:CB	2.29	0.61
21:AV:184:ALA:O	21:AV:186:GLU:N	2.33	0.61
34:BG:49:ARG:HH22	53:B1:57:U:C1'	2.10	0.61
31:BA:1469:G:H2'	31:BA:1470:G:H8	1.65	0.61
31:BA:86:U:H2'	31:BA:87:A:OP1	2.00	0.61
32:BE:165:VAL:HG23	32:BE:166:ASP:N	2.14	0.61
33:BF:7:PRO:O	33:BF:11:ARG:HG2	1.99	0.61
34:BG:146:ILE:H	34:BG:146:ILE:HD12	1.65	0.61
38:BK:14:ARG:O	38:BK:18:ARG:HD3	1.99	0.61
38:BK:20:TYR:HA	38:BK:65:TYR:CE2	2.35	0.61
40:BM:54:PHE:CE1	40:BM:55:LYS:HE3	2.35	0.61
43:BP:70:LEU:HD13	43:BP:71:ARG:N	2.15	0.61
53:C1:29:G:H2'	53:C1:30:C:C5	2.35	0.61
54:CA:69:G:C2	54:CA:73:G:N7	2.68	0.61
37:CJ:79:ARG:HH12	37:CJ:82:GLY:HA2	1.63	0.61
42:CO:117:ARG:O	42:CO:119:LYS:O	2.19	0.61
43:CP:125:ARG:O	43:CP:126:LYS:C	2.39	0.61
13:D0:33:ARG:NH2	27:D5:55:ARG:CG	2.55	0.61
9:DM:42:TRP:CD1	16:D1:63:VAL:HG11	2.35	0.61
22:D3:32:ARG:O	22:D3:35:ASN:ND2	2.33	0.61
55:DA:1077:A:C4'	58:DL:93:ARG:HH22	2.13	0.61
55:DA:1079:C:C3'	55:DA:1080:A:H8	2.11	0.61
55:DA:1288:U:H4'	55:DA:1289:C:OP2	1.99	0.61
55:DA:1301:A:H4'	55:DA:1302:A:OP1	2.00	0.61
55:DA:1406:U:H2'	55:DA:1407:C:C6	2.35	0.61
55:DA:2134:A:H2'	55:DA:2135:A:C8	2.35	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1049:C:C2	55:DA:2751:G:O6	2.53	0.61
55:DA:29:U:H2'	55:DA:30:G:C8	2.34	0.61
55:DA:481:G:HO2'	55:DA:507:A:H61	1.48	0.61
56:DI:21:LYS:C	56:DI:26:ALA:HB2	2.20	0.61
8:DK:74:ASN:ND2	8:DK:74:ASN:H	1.97	0.61
58:DL:125:ARG:O	58:DL:126:MET:C	2.38	0.61
58:DL:60:TYR:OH	58:DL:65:PHE:C	2.37	0.61
58:DL:79:ARG:C	58:DL:81:ALA:N	2.53	0.61
58:DL:86:LYS:N	58:DL:86:LYS:HE2	2.14	0.61
11:DO:136:GLU:HA	11:DO:139:LYS:HE3	1.81	0.61
21:DV:111:VAL:CG2	21:DV:146:ILE:N	2.63	0.61
21:DV:150:LEU:HD23	21:DV:151:HIS:N	2.08	0.61
55:DA:111:A:H4'	24:DW:69:ARG:NH2	2.15	0.61
17:A2:44:LYS:O	17:A2:46:VAL:N	2.33	0.61
1:AA:118:A:OP2	1:AA:119:A:H5''	1.99	0.61
1:AA:2228:G:H2'	1:AA:2229:C:H6	1.64	0.61
1:AA:2520:C:C6	1:AA:2567:G:H1'	2.35	0.61
1:AA:443:A:H1'	1:AA:1201:C:O4'	2.01	0.61
1:AA:528:A:H8	1:AA:528:A:H3'	1.64	0.61
1:AA:95:G:H1'	24:AW:47:ASN:OD1	2.00	0.61
6:AG:173:LEU:O	6:AG:178:PHE:HB2	2.00	0.61
7:AH:118:PRO:HG2	7:AH:121:ILE:HG13	1.83	0.61
7:AH:137:ASP:OD2	7:AH:140:LYS:HE2	2.00	0.61
1:AA:2562:U:H1'	10:AN:23:ARG:HH12	1.65	0.61
14:AQ:106:ARG:HA	14:AQ:110:LEU:CG	2.29	0.61
21:AV:6:LYS:HB3	21:AV:8:TYR:HE2	1.64	0.61
23:AZ:86:SER:H	23:AZ:87:PRO:HD3	1.65	0.61
31:BA:1181:G:C4	31:BA:1182:G:N2	2.68	0.61
31:BA:946:A:H2'	31:BA:947:G:H8	1.64	0.61
46:BS:43:LYS:HA	46:BS:48:TRP:CB	2.31	0.61
48:BU:53:ARG:HA	48:BU:56:THR:OG1	2.00	0.61
54:CA:511:C:O2'	54:CA:512:U:O5'	2.19	0.61
54:CA:690:G:H22	41:CN:55:LYS:HZ3	1.46	0.61
34:CG:199:ASN:OD1	34:CG:201:GLN:HB3	2.01	0.61
42:CO:28:LYS:HD2	42:CO:30:ALA:HB2	1.82	0.61
49:CV:41:VAL:HG12	49:CV:44:MET:H	1.65	0.61
49:CV:41:VAL:HG23	49:CV:67:VAL:HG13	1.81	0.61
30:D8:33:ASN:O	30:D8:34:TRP:C	2.38	0.61
55:DA:254:G:O6	30:D8:5:LYS:HG2	1.99	0.61
55:DA:1693:U:O2'	3:DD:14:ARG:NH2	2.27	0.61
55:DA:1718:G:C3'	55:DA:1725:G:H5''	2.30	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2723:C:OP1	13:D0:3:HIS:CD2	2.52	0.61
55:DA:2779:U:H1'	55:DA:2781:A:C5	2.35	0.61
55:DA:897:C:H6	55:DA:897:C:P	2.21	0.61
4:DE:14:ILE:O	4:DE:15:PHE:CB	2.47	0.61
4:DE:48:GLN:O	4:DE:49:LEU:HD12	1.99	0.61
5:DF:133:ASN:O	5:DF:135:LYS:N	2.32	0.61
7:DH:125:VAL:HA	7:DH:126:PRO:CB	2.30	0.61
7:DH:6:ARG:NE	7:DH:54:ARG:HH12	1.99	0.61
55:DA:943:U:OP2	11:DO:36:LYS:CE	2.48	0.61
23:DZ:56:GLN:N	23:DZ:56:GLN:NE2	2.45	0.61
30:A8:46:ARG:HH11	30:A8:46:ARG:HB2	1.65	0.61
1:AA:1494:A:H2'	1:AA:1495:A:C8	2.35	0.61
1:AA:2173:A:N3	1:AA:2173:A:H2'	2.15	0.61
1:AA:2331:G:O2'	22:A3:43:THR:HG22	2.00	0.61
1:AA:2611:U:O2	27:A5:3:LYS:HE2	2.00	0.61
1:AA:654(C):G:C3'	1:AA:654(D):G:H8	2.12	0.61
5:AF:27:GLU:O	5:AF:28:ILE:HG13	2.01	0.61
11:AO:11:GLY:O	11:AO:12:ALA:HB3	2.01	0.61
12:AP:42:ILE:HD11	12:AP:127:ILE:HD11	1.82	0.61
1:AA:2012:G:H5''	18:AS:96:ILE:HD11	1.82	0.61
21:AV:114:GLY:O	21:AV:116:VAL:N	2.33	0.61
31:BA:1160:G:C6	31:BA:1177:G:N2	2.67	0.61
31:BA:794:A:H2'	31:BA:795:C:H6	1.64	0.61
32:BE:166:ASP:OD1	32:BE:169:LYS:HB2	2.00	0.61
32:BE:59:GLU:HA	32:BE:221:LEU:HD13	1.82	0.61
38:BK:97:VAL:HG21	38:BK:128:GLY:HA2	1.83	0.61
40:BM:51:ARG:HB2	40:BM:60:ARG:HA	1.82	0.61
42:BO:33:ARG:O	42:BO:85:ILE:HG22	1.99	0.61
44:BQ:47:LEU:HA	44:BQ:50:LYS:HB2	1.81	0.61
54:CA:1541:U:O2	53:C1:32:A:C6	2.53	0.61
52:CD:15:G:N1	52:CD:48:C:N4	2.48	0.61
40:CM:63:PHE:HD1	44:CQ:58:LYS:HA	1.66	0.61
44:CQ:24:CYS:HB2	44:CQ:39:LEU:C	2.21	0.61
16:D1:91:ASP:O	16:D1:92:ARG:C	2.38	0.61
17:D2:41:GLY:HA3	17:D2:46:VAL:CG1	2.31	0.61
55:DA:1111:A:O2'	55:DA:1112:G:C4'	2.46	0.61
55:DA:1113:U:H5'	7:DH:2:SER:HB2	1.81	0.61
55:DA:2021:C:H5''	55:DA:2022:U:OP2	2.00	0.61
55:DA:2112:G:N1	55:DA:2169:A:N6	2.47	0.61
55:DA:2162:G:H4'	55:DA:2173:A:OP2	1.99	0.61
55:DA:2348:U:H4'	28:D6:42:TRP:CD1	2.35	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2258:C:O2'	55:DA:2427:C:OP2	2.17	0.61
55:DA:704:G:H2'	55:DA:726:G:N2	2.12	0.61
55:DA:70:G:H21	55:DA:71:A:H62	1.48	0.61
55:DA:890:A:H2'	55:DA:892:G:O4'	2.00	0.61
3:DD:131:LEU:HB2	3:DD:136:ILE:HD11	1.83	0.61
4:DE:1:MET:HG2	4:DE:83:ASP:O	1.99	0.61
56:DJ:16:THR:OG1	56:DJ:17:VAL:N	2.30	0.61
58:DL:53:VAL:CA	58:DL:72:PRO:HB2	2.30	0.61
9:DM:58:ASP:N	9:DM:60:ILE:HD11	2.15	0.61
21:DV:108:PRO:O	21:DV:109:ALA:HB3	1.99	0.61
27:A5:40:LYS:HZ2	27:A5:46:CYS:N	1.98	0.61
1:AA:1111:A:C4'	7:AH:3:ARG:HD3	2.29	0.61
1:AA:2129:C:H2'	1:AA:2130:U:C5'	2.30	0.61
1:AA:2315:G:H2'	1:AA:2316:C:C6	2.35	0.61
1:AA:2638:G:HO2'	1:AA:2639:A:H8	1.46	0.61
1:AA:531:C:H5''	1:AA:532:A:O4'	1.99	0.61
3:AD:186:HIS:HD2	3:AD:188:GLU:HB2	1.65	0.61
4:AE:119:ARG:HG2	4:AE:160:TYR:HB2	1.82	0.61
4:AE:8:LYS:HB3	4:AE:193:GLY:H	1.66	0.61
4:AE:57:LYS:NZ	4:AE:72:VAL:HG22	2.15	0.61
15:AR:8:LYS:HA	15:AR:11:GLU:OE1	2.01	0.61
15:AR:26:ASP:CB	15:AR:91:ARG:HA	2.31	0.61
18:AS:5:ALA:HB2	18:AS:54:ALA:HA	1.83	0.61
21:AV:63:ASP:O	21:AV:63:ASP:CG	2.38	0.61
21:AV:67:LEU:HD22	21:AV:90:VAL:HG13	1.83	0.61
24:AW:50:ILE:HD12	24:AW:51:ARG:N	2.10	0.61
31:BA:1365:G:H2'	31:BA:1366:C:H6	1.65	0.61
32:BE:178:ARG:HH22	32:BE:196:LEU:C	2.04	0.61
40:BM:63:PHE:HB3	44:BQ:57:ARG:O	2.00	0.61
45:BR:39:LEU:CD1	45:BR:56:LEU:HB2	2.29	0.61
26:A4:56:VAL:HG21	49:BV:64:GLU:OE2	1.99	0.61
54:CA:1007:C:C3'	54:CA:1008:C:H5''	2.29	0.61
34:CG:94:LEU:HA	34:CG:97:LEU:HD12	1.81	0.61
54:CA:939:G:H5''	37:CJ:102:ARG:NH2	2.14	0.61
13:D0:24:GLN:NE2	13:D0:36:THR:HG21	2.16	0.61
55:DA:1292:U:H2'	55:DA:1293:C:C6	2.36	0.61
55:DA:1906:G:C6	55:DA:1929:G:N2	2.67	0.61
55:DA:2760:C:O2'	55:DA:2761:G:H5''	1.99	0.61
55:DA:857:C:H1'	22:D3:26:TYR:CE2	2.36	0.61
2:DB:25:A:C2	2:DB:26:A:H1'	2.35	0.61
2:DB:31:C:O2	2:DB:31:C:H2'	2.00	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:DD:237:GLU:OE2	3:DD:237:GLU:CA	2.48	0.61
4:DE:21:VAL:HB	4:DE:22:PRO:CA	2.30	0.61
6:DG:21:ARG:HG2	6:DG:21:ARG:NH1	2.15	0.61
7:DH:132:ARG:HB2	7:DH:132:ARG:HH11	1.64	0.61
56:DJ:21:LYS:C	56:DJ:23:LEU:H	2.03	0.61
57:DY:28:ASN:HB2	57:DY:81:VAL:O	2.01	0.61
57:DY:8:GLU:OE1	57:DY:8:GLU:HA	2.00	0.61
1:AA:2329:G:H2'	1:AA:2330:G:C8	2.36	0.61
1:AA:2755:C:H4'	1:AA:2756:U:C5	2.25	0.61
1:AA:2631:G:N3	1:AA:2810:A:H2	1.98	0.61
1:AA:311:A:H1'	1:AA:332:A:C8	2.35	0.61
5:AF:192:LEU:HD21	5:AF:194:MET:CE	2.30	0.61
6:AG:126:ASP:OD1	6:AG:130:ASN:HB2	2.01	0.61
14:AQ:67:ARG:NH1	14:AQ:67:ARG:HB2	2.15	0.61
20:AU:72:VAL:O	20:AU:73:ARG:HB2	2.00	0.61
21:AV:11:GLU:CG	21:AV:12:GLY:H	2.13	0.61
21:AV:67:LEU:HD23	21:AV:68:PRO:CD	2.25	0.61
21:AV:70:LEU:HB2	21:AV:91:LEU:HD21	1.82	0.61
1:AA:95:G:H4'	24:AW:46:GLN:HB3	1.83	0.61
31:BA:115:G:O2'	31:BA:116:A:OP2	2.15	0.61
31:BA:1176:A:N6	31:BA:1177:G:C6	2.69	0.61
31:BA:1248:A:C2'	39:BL:70:LYS:HZ1	2.12	0.61
31:BA:66:G:OP2	31:BA:66:G:H8	1.83	0.61
31:BA:687:A:N6	31:BA:703:G:H1'	2.15	0.61
31:BA:723:U:H3	31:BA:1537:U:HO2'	1.47	0.61
52:BC:39:U:H2'	52:BC:40:C:C6	2.34	0.61
41:BN:23:ALA:HB1	41:BN:88:GLY:H	1.66	0.61
49:BV:29:ARG:HG3	49:BV:48:THR:HG1	1.64	0.61
54:CA:1238:A:H62	54:CA:1299:A:N6	1.99	0.61
54:CA:313:A:H2'	54:CA:314:C:C6	2.35	0.61
54:CA:531:U:H5''	54:CA:532:A:OP1	2.00	0.61
52:CD:19:G:N2	55:DA:2112:G:H21	1.99	0.61
40:CM:91:PRO:HB2	40:CM:94:VAL:HB	1.80	0.61
55:DA:1728:G:C2	55:DA:1730:U:OP2	2.54	0.61
55:DA:2531:A:H2	55:DA:2658:C:O2	1.84	0.61
55:DA:702:G:H5'	55:DA:702:G:H8	1.65	0.61
3:DD:263:ARG:CB	3:DD:263:ARG:HH11	2.14	0.61
3:DD:44:ASN:HB3	3:DD:49:ILE:HA	1.81	0.61
7:DH:3:ARG:HA	7:DH:3:ARG:HE	1.66	0.61
2:DB:116:G:H4'	14:DQ:54:LEU:HD12	1.82	0.61
20:DU:84:ARG:NH1	20:DU:97:ARG:HA	2.14	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:DV:102:LEU:O	21:DV:103:ARG:HD2	2.01	0.61
21:DV:194:PRO:CB	21:DV:196:VAL:HG11	2.31	0.61
57:DY:9:LEU:CG	57:DY:10:LEU:N	2.64	0.61
55:DA:1083:U:O5'	57:DY:47:ASN:OD1	2.17	0.61
16:A1:79:PHE:O	16:A1:79:PHE:HD2	1.83	0.61
16:A1:90:VAL:O	16:A1:92:ARG:N	2.33	0.61
1:AA:1359:A:H5'	1:AA:1359:A:C8	2.35	0.61
1:AA:805:G:H4'	1:AA:806:C:OP2	2.00	0.61
4:AE:61:ARG:C	4:AE:63:LEU:H	2.04	0.61
11:AO:82:GLY:HA2	11:AO:113:LYS:O	2.00	0.61
14:AQ:11:LYS:HD3	14:AQ:91:PRO:HD3	1.83	0.61
20:AU:52:SER:H	20:AU:53:PRO:HD3	1.64	0.61
31:BA:1109:C:H2'	31:BA:1110:A:O4'	2.00	0.61
31:BA:1319:A:H5'	31:BA:1320:C:OP1	2.01	0.61
31:BA:1391:U:H2'	31:BA:1392:G:C8	2.35	0.61
31:BA:281:G:H5''	31:BA:282:A:OP1	2.01	0.61
31:BA:625:G:H2'	31:BA:626:U:H6	1.66	0.61
1:AA:881:G:O3'	52:BB:19:G:N7	2.34	0.61
52:BD:59:U:H2'	52:BD:60:U:H5'	1.82	0.61
32:BE:168:THR:HA	32:BE:171:ALA:HB2	1.82	0.61
33:BF:182:ILE:HG23	33:BF:202:ILE:O	2.01	0.61
34:BG:25:ARG:CB	34:BG:25:ARG:HH11	1.99	0.61
39:BL:37:PHE:HB3	39:BL:43:ALA:CB	2.31	0.61
33:BF:20:SER:O	44:BQ:54:PRO:HG3	2.01	0.61
31:BA:668:G:O2'	45:BR:46:HIS:HD2	1.83	0.61
54:CA:1350:A:H2'	54:CA:1351:U:C6	2.36	0.61
54:CA:328:C:H2'	54:CA:328:C:O2	2.01	0.61
32:CE:75:LYS:HD3	32:CE:75:LYS:C	2.21	0.61
39:CL:62:TYR:C	39:CL:63:ILE:HD12	2.20	0.61
55:DA:1086:A:C4'	55:DA:1103:A:H61	2.12	0.61
55:DA:1085:A:C2'	55:DA:1086:A:N7	2.60	0.61
55:DA:1728:G:H5'	55:DA:1729:A:OP2	2.00	0.61
55:DA:2175:C:C3'	55:DA:2176:A:H5''	2.31	0.61
55:DA:270(O):U:H5''	55:DA:270(P):C:OP2	2.01	0.61
55:DA:304:G:H2'	55:DA:305:U:C6	2.36	0.61
3:DD:181:GLU:OE1	3:DD:270:ILE:HG23	2.00	0.61
7:DH:89:ILE:HD13	7:DH:89:ILE:C	2.19	0.61
58:DL:90:LYS:N	58:DL:91:PRO:CA	2.64	0.61
11:DO:140:ALA:O	11:DO:141:ALA:CB	2.49	0.61
19:DT:29:TRP:CZ3	19:DT:78:LYS:HG3	2.35	0.61
21:DV:191:VAL:HG12	21:DV:197:ILE:HG21	1.63	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:DV:63:ASP:O	21:DV:65:GLN:HG2	2.01	0.61
57:DY:101:PRO:CG	57:DY:102:LYS:N	2.63	0.61
57:DY:138:LEU:O	57:DY:139:VAL:CG1	2.44	0.61
28:A6:48:VAL:O	28:A6:49:HIS:HB2	1.99	0.61
1:AA:1338:G:N3	1:AA:1393:A:H2	1.99	0.61
1:AA:1810:A:H2'	1:AA:1811:G:H5'	1.81	0.61
1:AA:2571:C:C5'	1:AA:2572:A:H5''	2.30	0.61
1:AA:492:A:H2'	1:AA:493:G:O4'	2.00	0.61
1:AA:1826:G:C4'	3:AD:242:ARG:HH21	2.13	0.61
5:AF:4:VAL:HG11	5:AF:17:ARG:HE	1.66	0.61
6:AG:115:ARG:NH2	43:BP:7:VAL:HB	2.16	0.61
7:AH:122:THR:O	7:AH:133:VAL:HG13	2.01	0.61
7:AH:125:VAL:HG22	7:AH:126:PRO:CB	2.31	0.61
9:AM:13:TRP:O	9:AM:135:PRO:HD2	2.00	0.61
10:AN:23:ARG:HH11	10:AN:23:ARG:HG2	1.66	0.61
11:AO:107:LYS:O	11:AO:109:GLY:N	2.34	0.61
12:AP:133:ARG:O	12:AP:134:ARG:HB2	2.01	0.61
21:AV:4:ARG:NH1	21:AV:58:VAL:HG11	2.16	0.61
37:BJ:137:LYS:O	37:BJ:137:LYS:HE2	2.01	0.61
37:BJ:86:GLN:HB2	37:BJ:148:ASN:OD1	2.01	0.61
36:BI:97:PHE:O	48:BU:31:LEU:HD23	2.01	0.61
49:BV:62:ILE:HA	49:BV:66:MET:CE	2.31	0.61
54:CA:1160:G:C6	54:CA:1177:G:N2	2.68	0.61
54:CA:1190:G:OP1	33:CF:4:LYS:HA	2.00	0.61
54:CA:85:U:O2'	54:CA:86:U:O5'	2.16	0.61
54:CA:89:U:C2'	54:CA:90:C:O5'	2.49	0.61
52:CD:72:C:C3'	52:CD:73:A:H5''	2.30	0.61
32:CE:66:GLY:O	32:CE:67:THR:HG23	2.01	0.61
54:CA:1205:U:H1'	33:CF:195:VAL:CG2	2.31	0.61
37:CJ:21:VAL:HG23	37:CJ:22:LEU:N	2.14	0.61
38:CK:111:ILE:HG22	38:CK:112:LEU:N	2.16	0.61
39:CL:114:TYR:CD2	39:CL:114:TYR:O	2.53	0.61
43:CP:110:ARG:O	43:CP:110:ARG:HG2	2.01	0.61
45:CR:17:ARG:NH1	45:CR:77:ARG:NH1	2.48	0.61
45:CR:78:TYR:OH	45:CR:88:ARG:HG3	2.00	0.61
49:CV:41:VAL:CG1	49:CV:44:MET:HB2	2.28	0.61
50:CW:13:LEU:CD1	50:CW:17:ARG:HH12	2.13	0.61
16:D1:91:ASP:OD2	16:D1:96:ALA:HB2	2.00	0.61
27:D5:55:ARG:HG3	27:D5:57:VAL:N	2.10	0.61
55:DA:1085:A:H2'	55:DA:1086:A:H8	1.60	0.61
55:DA:1311:G:H21	55:DA:1603:A:H62	1.48	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2062:A:H2'	55:DA:2062:A:N3	2.15	0.61
55:DA:366:C:C5	55:DA:403:U:O2'	2.50	0.61
55:DA:654(B):C:H2'	55:DA:654(C):G:C1'	2.30	0.61
55:DA:1565:C:H5''	3:DD:18:VAL:HG11	1.83	0.61
3:DD:206:LEU:HA	3:DD:211:ARG:HH11	1.66	0.61
7:DH:109:PHE:CZ	7:DH:152:ARG:HG2	2.36	0.61
8:DK:41:GLU:H	8:DK:41:GLU:CD	2.04	0.61
11:DO:126:VAL:HG12	11:DO:147:LEU:HD21	1.82	0.61
14:DQ:13:ARG:O	14:DQ:13:ARG:HD2	2.01	0.61
55:DA:2378:A:H4'	14:DQ:23:ARG:NH1	2.15	0.61
14:DQ:49:VAL:HG21	14:DQ:77:ALA:HA	1.82	0.61
15:DR:94:ALA:O	15:DR:95:ARG:CB	2.49	0.61
57:DY:47:ASN:O	57:DY:48:GLY:O	2.18	0.61
28:A6:17:LYS:C	28:A6:19:ARG:H	2.04	0.61
1:AA:2030:A:H4'	1:AA:2031:A:C8	2.34	0.61
1:AA:2091:U:C3'	1:AA:2092:U:C5'	2.73	0.61
1:AA:2291:U:H2'	1:AA:2292:C:C6	2.35	0.61
1:AA:2404:C:H2'	1:AA:2405:G:O4'	2.01	0.61
1:AA:2648:C:H2'	1:AA:2649:U:C6	2.35	0.61
1:AA:2656:U:O4	1:AA:2657:A:C5	2.54	0.61
1:AA:971:C:H2'	1:AA:972:G:H5'	1.81	0.61
4:AE:197:ILE:O	4:AE:197:ILE:HG13	2.00	0.61
5:AF:132:VAL:HG22	5:AF:133:ASN:N	2.16	0.61
5:AF:20:LEU:HD23	5:AF:21:ALA:N	2.15	0.61
8:AK:76:THR:OG1	8:AK:77:LEU:N	2.33	0.61
9:AM:57:ALA:O	9:AM:60:ILE:HG13	2.01	0.61
10:AN:88:ASN:OD1	10:AN:92:GLU:HB2	2.00	0.61
15:AR:96:ARG:CZ	15:AR:96:ARG:HB2	2.31	0.61
1:AA:851:U:C5'	25:AX:49:LYS:HD2	2.30	0.61
23:AZ:44:PRO:HG2	23:AZ:46:LEU:CD1	2.29	0.61
31:BA:1067:A:O2'	31:BA:1068:G:C8	2.50	0.61
31:BA:1348:U:H3	31:BA:1374:A:H2	1.49	0.61
31:BA:250:A:O2'	31:BA:251:G:OP2	2.16	0.61
32:BE:111:ARG:HH11	32:BE:111:ARG:HA	1.66	0.61
32:BE:5:ILE:HD12	32:BE:59:GLU:HB2	1.83	0.61
32:BE:78:GLN:CB	32:BE:94:ASN:HD21	2.14	0.61
33:BF:35:GLU:HA	33:BF:38:ARG:CZ	2.30	0.61
34:BG:21:LEU:CD1	34:BG:21:LEU:H	1.80	0.61
31:BA:134:A:N6	46:BS:25:ARG:NH1	2.43	0.61
46:BS:70:ALA:O	46:BS:74:LEU:HD12	2.00	0.61
54:CA:1139:G:H5'	54:CA:1140:C:OP1	2.00	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1127:G:N2	54:CA:1145:C:C2	2.69	0.61
54:CA:781:A:OP1	54:CA:1523:G:H5'	2.00	0.61
32:CE:24:TRP:CZ3	32:CE:26:PRO:HA	2.34	0.61
54:CA:1205:U:H1'	33:CF:195:VAL:HG23	1.83	0.61
42:CO:28:LYS:CE	42:CO:33:ARG:HH22	2.13	0.61
55:DA:1082:U:O3'	58:DL:117:THR:HG23	2.01	0.61
55:DA:1332:G:N2	55:DA:1609:A:C2'	2.58	0.61
55:DA:1729:A:H2'	55:DA:1730:U:H5''	1.82	0.61
55:DA:192:C:H2'	55:DA:193:U:H5'	1.81	0.61
55:DA:2531:A:H61	55:DA:2662:A:H61	1.48	0.61
55:DA:1050:A:H8	55:DA:2751:G:H2'	1.66	0.61
3:DD:206:LEU:HA	3:DD:211:ARG:NH1	2.16	0.61
3:DD:25:THR:O	3:DD:27:THR:N	2.33	0.61
5:DF:46:ARG:HG2	5:DF:46:ARG:NH1	2.11	0.61
5:DF:57:VAL:CG1	5:DF:58:ALA:N	2.64	0.61
56:DI:3:LEU:HD21	56:DI:7:ARG:HD2	1.76	0.61
58:DL:53:VAL:HG12	58:DL:72:PRO:HB2	1.81	0.61
10:DN:86:ILE:CG2	10:DN:94:ARG:HD2	2.30	0.61
12:DP:37:LEU:HD21	12:DP:130:LYS:HE3	1.81	0.61
57:DY:71:LEU:CB	57:DY:112:LEU:C	2.61	0.61
57:DY:31:GLY:O	57:DY:32:LEU:HD13	2.01	0.61
57:DY:41:ARG:O	57:DY:41:ARG:HD2	2.01	0.61
6:AG:104:GLU:HG2	26:A4:23:GLU:HG2	1.82	0.61
29:A7:19:ARG:HG2	29:A7:19:ARG:NH1	2.10	0.61
1:AA:1099:G:H2'	1:AA:1100:C:C6	2.36	0.61
1:AA:1275:A:H4'	1:AA:1276:A:O5'	1.99	0.61
1:AA:1794:U:H2'	1:AA:1795:C:C6	2.35	0.61
1:AA:414:C:H1'	1:AA:1864:U:O2'	2.00	0.61
1:AA:1928:A:O2'	1:AA:1929:G:H5''	2.00	0.61
1:AA:2092:U:C6	1:AA:2092:U:H5''	2.36	0.61
1:AA:2439:A:H8	1:AA:2439:A:H5'	1.65	0.61
1:AA:387:U:H6	1:AA:387:U:O5'	1.84	0.61
1:AA:864:G:H2'	1:AA:865:C:C6	2.36	0.61
2:AB:15:A:H5'	2:AB:16:G:C8	2.35	0.61
7:AH:127:GLU:OE1	7:AH:130:ARG:HB3	1.99	0.61
8:AK:82:ARG:HG3	8:AK:146:ALA:H	1.65	0.61
18:AS:55:ALA:O	18:AS:58:ALA:HB3	2.01	0.61
20:AU:83:THR:HG21	20:AU:94:LYS:HG2	1.83	0.61
24:AW:13:ALA:HA	24:AW:16:LEU:HD21	1.83	0.61
1:AA:111:A:H4'	24:AW:69:ARG:NH2	2.16	0.61
52:BB:19:G:C1'	52:BB:57:G:N2	2.64	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:BE:231:GLU:HB3	32:BE:232:PRO:CD	2.29	0.61
34:BG:31:CYS:SG	34:BG:31:CYS:O	2.59	0.61
36:BI:8:ILE:HG21	36:BI:26:ILE:HD11	1.83	0.61
45:BR:2:PRO:HG2	45:BR:3:ILE:H	1.66	0.61
45:BR:82:ILE:O	45:BR:82:ILE:HD13	2.01	0.61
54:CA:157:G:O2'	54:CA:158:G:H5'	2.01	0.61
54:CA:399:G:H2'	54:CA:400:C:H6	1.66	0.61
52:CB:74:C:O2'	52:CB:75:C:H5'	2.01	0.61
33:CF:101:LEU:HD23	33:CF:102:ASN:N	2.16	0.61
33:CF:99:VAL:O	33:CF:99:VAL:HG23	2.01	0.61
36:CI:99:ALA:HB3	48:CU:29:PHE:CE2	2.36	0.61
44:CQ:40:CYS:SG	44:CQ:43:CYS:N	2.68	0.61
48:CU:43:PHE:CE2	48:CU:58:LEU:HD11	2.36	0.61
55:DA:1105:U:H2'	55:DA:1106:G:H8	1.64	0.61
55:DA:1794:U:H2'	55:DA:1795:C:H6	1.65	0.61
55:DA:859:G:O2'	55:DA:860:U:P	2.59	0.61
55:DA:890:A:H8	55:DA:892:G:C8	2.19	0.61
3:DD:62:TYR:HA	3:DD:87:ASN:HD21	1.64	0.61
4:DE:35:GLN:HG2	4:DE:37:ARG:HG2	1.83	0.61
9:DM:39:ARG:HB3	9:DM:39:ARG:NH1	2.16	0.61
19:DT:41:ASN:HD22	19:DT:41:ASN:H	1.47	0.61
21:DV:105:VAL:O	21:DV:140:ASP:HA	2.01	0.61
21:DV:193:GLU:N	21:DV:194:PRO:HD2	2.14	0.61
57:DY:35:LYS:HA	57:DY:35:LYS:CE	2.22	0.61
57:DY:87:VAL:O	57:DY:91:LYS:CB	2.46	0.61
23:DZ:80:LEU:O	23:DZ:80:LEU:HD22	2.00	0.61
28:A6:24:GLU:OE1	28:A6:24:GLU:HA	1.99	0.60
1:AA:1181:C:O2'	1:AA:1182:A:H5'	2.01	0.60
1:AA:2087:G:O2'	1:AA:2088:G:H5'	2.01	0.60
1:AA:847:U:H2'	1:AA:848:G:H5''	1.83	0.60
1:AA:878:A:N3	1:AA:878:A:H5'	2.16	0.60
1:AA:821:A:H2'	1:AA:946:G:H5''	1.82	0.60
5:AF:51:THR:HG23	5:AF:92:PRO:HG2	1.83	0.60
9:AM:128:HIS:NE2	9:AM:134:ARG:HD2	2.16	0.60
11:AO:101:VAL:HA	11:AO:105:LEU:O	2.01	0.60
1:AA:955:C:OP1	12:AP:13:GLN:HA	2.00	0.60
21:AV:175:VAL:HG22	21:AV:176:PRO:HD3	1.83	0.60
31:BA:119:A:O2'	31:BA:120:A:OP2	2.15	0.60
31:BA:1540:U:H2'	31:BA:1541:U:O4'	2.01	0.60
31:BA:748:C:O2'	31:BA:749:C:OP2	2.17	0.60
31:BA:973:G:H1'	40:BM:55:LYS:HG2	1.82	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:BB:74:C:C1'	52:BB:75:C:H5'	2.30	0.60
32:BE:19:HIS:CE1	32:BE:204:ASN:HB3	2.34	0.60
34:BG:34:GLU:O	34:BG:35:ARG:HG3	2.01	0.60
40:BM:33:GLN:HB2	40:BM:75:ILE:HG21	1.83	0.60
31:BA:667:G:H4'	45:BR:51:HIS:ND1	2.16	0.60
54:CA:1158:C:N3	54:CA:1160:G:N7	2.49	0.60
54:CA:1371:G:OP1	39:CL:12:GLU:HB2	2.01	0.60
54:CA:1400:C:H4'	54:CA:1401:G:OP2	2.00	0.60
54:CA:370:C:O2'	54:CA:371:G:H5'	2.01	0.60
54:CA:629:G:H5''	54:CA:630:G:O5'	2.01	0.60
54:CA:726:C:O2'	54:CA:727:G:H5'	2.01	0.60
52:CB:19:G:HO2'	52:CB:20:U:P	2.22	0.60
37:CJ:79:ARG:NH1	37:CJ:82:GLY:HA2	2.16	0.60
38:CK:16:ALA:HB2	38:CK:24:THR:HG21	1.83	0.60
39:CL:41:VAL:O	39:CL:41:VAL:HG12	2.01	0.60
42:CO:62:SER:O	42:CO:64:TYR:HD1	1.83	0.60
46:CS:4:ILE:HD11	46:CS:64:ALA:CB	2.26	0.60
49:CV:39:THR:HG22	49:CV:40:ILE:N	2.14	0.60
50:CW:101:GLY:O	50:CW:103:GLY:N	2.34	0.60
16:D1:65:ILE:HG12	16:D1:96:ALA:HB1	1.82	0.60
28:D6:28:ARG:HG3	28:D6:31:PRO:HD2	1.83	0.60
55:DA:1060:U:C1'	55:DA:1061:U:H3'	2.31	0.60
55:DA:1535:U:C5	55:DA:1537:C:O2	2.54	0.60
55:DA:1688:U:O2	55:DA:1700:A:H5''	2.01	0.60
55:DA:273(F):C:H3'	55:DA:274:G:C5'	2.31	0.60
58:DL:11:GLN:HA	58:DL:23:VAL:CG1	2.31	0.60
58:DL:59:ILE:CG2	58:DL:60:TYR:N	2.59	0.60
9:DM:63:THR:HG22	9:DM:66:LYS:HZ2	1.66	0.60
55:DA:636:G:OP1	11:DO:132:LYS:HB2	2.01	0.60
11:DO:71:VAL:HG13	11:DO:72:PRO:CD	2.29	0.60
18:DS:92:ARG:O	18:DS:93:ALA:HB3	2.01	0.60
57:DY:141:VAL:CG2	57:DY:142:LEU:N	2.53	0.60
17:A2:14:VAL:CB	17:A2:96:ILE:HG13	2.28	0.60
1:AA:1324:G:O2'	1:AA:1616:A:N7	2.28	0.60
1:AA:1688:U:H5'	1:AA:1689:A:OP1	2.01	0.60
1:AA:2531:A:H61	1:AA:2662:A:H61	1.47	0.60
1:AA:2584:U:C5	1:AA:2585:U:C5	2.89	0.60
1:AA:2657:A:O2'	7:AH:160:LYS:HE3	2.01	0.60
1:AA:414:C:H2'	1:AA:415:A:H8	1.66	0.60
1:AA:74:A:C5'	1:AA:75:G:O4'	2.49	0.60
1:AA:813:U:H2'	1:AA:814:C:H6	1.64	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1820:U:H2'	3:AD:158:ALA:O	2.00	0.60
12:AP:85:LYS:HG3	12:AP:86:GLY:N	2.15	0.60
14:AQ:59:LYS:HG2	14:AQ:60:GLY:N	2.15	0.60
15:AR:95:ARG:HH11	15:AR:95:ARG:HG3	1.65	0.60
1:AA:309:G:O3'	20:AU:18:GLY:HA3	2.01	0.60
31:BA:1024:G:N3	31:BA:1024:G:H3'	2.15	0.60
31:BA:1363:A:C4'	31:BA:1364:U:OP1	2.49	0.60
31:BA:1410:G:H2'	31:BA:1411:C:H6	1.65	0.60
31:BA:509:A:H5''	34:BG:55:ALA:HB2	1.82	0.60
32:BE:67:THR:C	32:BE:68:ILE:HD12	2.22	0.60
35:BH:51:VAL:O	35:BH:55:VAL:HG23	2.02	0.60
36:BI:77:ARG:HB3	36:BI:77:ARG:HH11	1.65	0.60
39:BL:46:ALA:HA	39:BL:78:LYS:HB2	1.83	0.60
39:BL:82:ALA:HB1	39:BL:96:LEU:HD11	1.82	0.60
43:BP:79:LYS:HE2	43:BP:82:MET:HE1	1.82	0.60
31:BA:376:G:OP2	46:BS:67:THR:HG21	2.00	0.60
47:BT:57:VAL:HG12	47:BT:76:LEU:HA	1.83	0.60
48:BU:50:ILE:CD1	48:BU:70:ILE:HG21	2.30	0.60
54:CA:186(A):C:H2'	54:CA:186(B):C:C6	2.35	0.60
52:CD:59:U:H3'	52:CD:60:U:H6	1.65	0.60
32:CE:204:ASN:HD22	32:CE:205:ASP:N	1.99	0.60
32:CE:29:ALA:HA	32:CE:32:ILE:HG22	1.83	0.60
36:CI:72:VAL:CG2	36:CI:90:VAL:HG11	2.30	0.60
38:CK:65:TYR:HA	38:CK:79:VAL:HG23	1.82	0.60
44:CQ:44:LEU:HD12	44:CQ:44:LEU:C	2.22	0.60
46:CS:75:ARG:C	46:CS:77:ALA:H	2.04	0.60
16:D1:96:ALA:C	16:D1:98:LEU:H	2.03	0.60
55:DA:2060:A:HO2'	55:DA:2061:G:P	2.24	0.60
55:DA:2060:A:O2'	55:DA:2061:G:P	2.59	0.60
55:DA:2688:U:C5	55:DA:2720:U:OP2	2.54	0.60
55:DA:775:G:O5'	55:DA:777:A:H1'	2.01	0.60
55:DA:784:A:C5	3:DD:229:VAL:HG21	2.36	0.60
55:DA:877:U:H2'	55:DA:878:A:O5'	2.01	0.60
55:DA:917:A:H2'	55:DA:918:A:H5'	1.83	0.60
3:DD:34:VAL:C	3:DD:35:LYS:HG3	2.21	0.60
4:DE:26:ILE:HD13	4:DE:27:LEU:N	2.15	0.60
5:DF:127:GLU:OE1	5:DF:127:GLU:HA	2.01	0.60
56:DI:20:LEU:O	56:DI:24:ILE:HG22	2.00	0.60
56:DI:24:ILE:HD13	56:DI:26:ALA:N	2.15	0.60
11:DO:113:LYS:HG2	11:DO:115:LEU:HD23	1.82	0.60
11:DO:122:PRO:CB	11:DO:141:ALA:HB1	2.25	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:DS:88:ARG:HH11	18:DS:88:ARG:HG2	1.66	0.60
57:DY:116:ILE:O	57:DY:116:ILE:CG1	2.49	0.60
57:DY:35:LYS:CA	57:DY:35:LYS:CE	2.79	0.60
57:DY:93:LEU:HD13	57:DY:97:ALA:C	2.22	0.60
1:AA:1252:G:N7	16:A1:36:ARG:NH1	2.49	0.60
1:AA:2219:G:H2'	1:AA:2224:G:H5'	1.83	0.60
1:AA:2734:A:H5'	1:AA:2735:G:OP2	2.01	0.60
1:AA:620:G:H4'	1:AA:621:A:C5'	2.31	0.60
1:AA:627:A:O2'	1:AA:628:G:C8	2.54	0.60
1:AA:85:G:OP1	20:AU:30:VAL:HG21	2.02	0.60
2:AB:39:A:N1	26:A4:1:MET:HB3	2.16	0.60
1:AA:2746:U:H5''	7:AH:138:LYS:HE3	1.83	0.60
1:AA:84:A:H5''	20:AU:8:LYS:HG2	1.83	0.60
31:BA:1126:U:H4'	31:BA:1127:G:C8	2.37	0.60
31:BA:347:G:O2'	31:BA:348:G:H5'	2.00	0.60
31:BA:857:C:H2'	31:BA:858:G:O4'	2.02	0.60
52:BB:5:G:H2'	52:BB:6:G:H8	1.65	0.60
52:BD:19:G:H4'	52:BD:57:G:H21	1.64	0.60
32:BE:54:THR:HB	32:BE:201:ILE:HD11	1.83	0.60
35:BH:76:ILE:CG2	35:BH:77:PRO:HD2	2.29	0.60
37:BJ:16:LEU:CD1	39:BL:42:ARG:HA	2.30	0.60
31:BA:716:A:N3	41:BN:117:ASN:O	2.35	0.60
42:BO:25:PRO:HD2	42:BO:98:TYR:OH	2.01	0.60
54:CA:1070:U:H2'	54:CA:1071:C:H6	1.65	0.60
54:CA:1330:U:H5'	54:CA:1331:G:OP2	2.01	0.60
54:CA:245:C:O2	54:CA:283:C:N3	2.34	0.60
54:CA:50:A:O2'	54:CA:51:A:P	2.59	0.60
54:CA:657:G:O2'	54:CA:658:G:H5'	2.01	0.60
34:CG:129:ASN:HA	34:CG:145:GLU:CB	2.31	0.60
34:CG:150:GLU:O	34:CG:152:SER:N	2.34	0.60
35:CH:32:VAL:HG23	35:CH:58:ALA:HB1	1.82	0.60
39:CL:53:VAL:O	39:CL:54:ASP:HB2	2.00	0.60
43:CP:23:TYR:HE1	43:CP:70:LEU:HD12	1.64	0.60
55:DA:1142(A):A:O2'	55:DA:1143:A:O5'	2.16	0.60
55:DA:1537:C:C5	55:DA:1538:G:C5	2.89	0.60
55:DA:2518:A:H5''	55:DA:2519:U:OP2	2.01	0.60
55:DA:2712:U:O2	55:DA:2712:U:H5''	2.01	0.60
55:DA:654(F):C:C2'	55:DA:654(G):C:OP1	2.48	0.60
55:DA:752:A:HO2'	55:DA:753:C:P	2.24	0.60
55:DA:686:G:H21	55:DA:788:A:H61	1.49	0.60
56:DJ:7:ARG:CG	56:DJ:7:ARG:NH1	2.62	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:DK:93:THR:O	8:DK:96:ASP:HB2	2.01	0.60
58:DL:102:GLU:CG	58:DL:103:GLN:H	2.14	0.60
9:DM:35:ARG:HD3	9:DM:37:LYS:HD2	1.84	0.60
12:DP:110:THR:HB	12:DP:112:GLU:OE2	2.02	0.60
20:DU:97:ARG:HH21	20:DU:98:VAL:CB	2.08	0.60
24:DW:15:LYS:H	24:DW:67:LYS:HE2	1.65	0.60
57:DY:131:MET:O	57:DY:133:GLU:HG3	2.02	0.60
57:DY:142:LEU:HD13	57:DY:143:GLN:CA	2.31	0.60
57:DY:87:VAL:HG13	57:DY:91:LYS:HG3	1.82	0.60
16:A1:16:LYS:O	16:A1:20:LEU:HD23	2.01	0.60
1:AA:2296:U:C4'	1:AA:2297:C:OP1	2.49	0.60
1:AA:2745:C:H4'	7:AH:142:GLY:O	2.02	0.60
1:AA:514:A:O2'	1:AA:515:A:H5'	2.01	0.60
2:AB:78:A:H2'	2:AB:79:C:O4'	2.02	0.60
4:AE:101:ARG:HD2	4:AE:169:ASN:O	2.01	0.60
5:AF:114:VAL:HG21	5:AF:202:PHE:CE1	2.36	0.60
7:AH:20:ALA:CB	7:AH:23:ARG:HE	2.13	0.60
10:AN:11:ALA:HB1	10:AN:99:PHE:O	2.01	0.60
14:AQ:106:ARG:HB2	14:AQ:106:ARG:CZ	2.32	0.60
14:AQ:49:VAL:HG21	14:AQ:77:ALA:HA	1.83	0.60
15:AR:95:ARG:NH1	15:AR:95:ARG:HG3	2.17	0.60
19:AT:30:VAL:HG12	19:AT:31:HIS:N	2.16	0.60
21:AV:107:THR:N	21:AV:108:PRO:CD	2.52	0.60
21:AV:122:ARG:HH11	21:AV:122:ARG:HG2	1.66	0.60
21:AV:170:THR:O	21:AV:171:ILE:HB	2.02	0.60
53:B1:36:G:C3'	53:B1:37:G:H5''	2.31	0.60
31:BA:1174:G:H2'	31:BA:1175:G:C8	2.36	0.60
31:BA:1321:C:C3'	31:BA:1322:C:H5''	2.32	0.60
31:BA:820:U:C4'	31:BA:821:G:OP2	2.49	0.60
32:BE:17:PHE:CZ	32:BE:44:LEU:HA	2.36	0.60
33:BF:129:ALA:CB	33:BF:132:ARG:HB3	2.30	0.60
35:BH:75:THR:HG23	35:BH:76:ILE:N	2.17	0.60
36:BI:82:ARG:HG2	36:BI:82:ARG:HH11	1.67	0.60
40:BM:6:ILE:HG22	40:BM:98:ILE:HA	1.84	0.60
42:BO:25:PRO:C	42:BO:27:LEU:H	2.03	0.60
44:BQ:15:LYS:NZ	44:BQ:16:PHE:H	1.98	0.60
51:BX:8:THR:O	51:BX:12:LYS:HB2	2.02	0.60
54:CA:1305:G:H22	54:CA:1331:G:C2'	2.02	0.60
54:CA:662:G:O2'	54:CA:836:G:H5''	2.01	0.60
52:CC:50:U:O2'	52:CC:51:U:H5'	2.01	0.60
32:CE:4:GLU:O	32:CE:5:ILE:HG23	2.00	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:CG:96:LEU:HD12	34:CG:139:ARG:CZ	2.32	0.60
48:CU:53:ARG:NH2	48:CU:60:ALA:N	2.49	0.60
50:CW:13:LEU:HD12	50:CW:17:ARG:HH12	1.66	0.60
26:D4:61:ARG:HA	26:D4:64:GLY:H	1.64	0.60
28:D6:11:LEU:HG	28:D6:51:GLU:OE2	2.01	0.60
30:D8:61:LEU:O	30:D8:62:LEU:HB2	2.01	0.60
55:DA:1059:G:OP2	55:DA:1060:U:H3'	2.01	0.60
55:DA:1071:G:H5'	55:DA:1088:A:O2'	2.00	0.60
55:DA:1092:C:H2'	55:DA:1093:G:O4'	2.01	0.60
55:DA:1538:G:O2'	55:DA:1539:G:H5'	2.01	0.60
55:DA:2677:G:H2'	55:DA:2678:C:C6	2.36	0.60
55:DA:548:A:H2'	55:DA:549:G:C5'	2.31	0.60
55:DA:968:G:H2'	55:DA:969:U:C6	2.36	0.60
56:DI:4:ASP:OD2	56:DI:5:ILE:HD12	2.02	0.60
58:DL:3:LYS:C	58:DL:4:VAL:HG23	2.22	0.60
24:DW:57:ILE:HG22	24:DW:61:LEU:HD12	1.83	0.60
57:DY:27:VAL:CG1	57:DY:110:GLY:HA3	2.19	0.60
57:DY:71:LEU:CA	57:DY:113:GLN:CB	2.69	0.60
6:AG:104:GLU:HG2	26:A4:23:GLU:HG3	1.83	0.60
1:AA:1171:G:P	1:AA:1171:G:H3'	2.42	0.60
1:AA:2707:G:H5''	13:A0:68:ARG:NH2	2.16	0.60
1:AA:518:G:H4'	18:AS:18:ARG:HH11	1.66	0.60
6:AG:171:ALA:O	6:AG:175:LEU:HG	2.01	0.60
11:AO:47:ASP:HB3	11:AO:48:PRO:CA	2.31	0.60
12:AP:52:VAL:O	12:AP:55:VAL:HG12	2.02	0.60
19:AT:34:ALA:CB	19:AT:39:ILE:HD11	2.28	0.60
21:AV:175:VAL:HG22	21:AV:176:PRO:CD	2.32	0.60
24:AW:24:LEU:HD21	24:AW:28:LYS:HE2	1.84	0.60
31:BA:1028(B):C:H3'	31:BA:1029:G:H5''	1.82	0.60
31:BA:1145:C:O2	31:BA:1145:C:H2'	2.02	0.60
31:BA:115:G:H1'	31:BA:116:A:N7	2.16	0.60
31:BA:531:U:C4'	31:BA:532:A:OP1	2.49	0.60
31:BA:965:A:C4'	31:BA:966:G:OP1	2.48	0.60
32:BE:101:MET:HE2	32:BE:108:ILE:HG21	1.83	0.60
36:BI:91:VAL:HG11	48:BU:72:ARG:HH12	1.66	0.60
46:BS:1:MET:HE1	46:BS:65:GLN:HB2	1.84	0.60
54:CA:1293:G:H2'	54:CA:1294:G:C8	2.36	0.60
54:CA:39:G:N7	54:CA:547:A:C8	2.69	0.60
54:CA:411:A:C5	54:CA:413:G:H1'	2.37	0.60
36:CI:38:GLU:HB2	36:CI:64:GLN:O	2.02	0.60
40:CM:48:THR:HA	40:CM:62:HIS:CB	2.20	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:D3:27:GLU:HA	22:D3:67:VAL:HG12	1.83	0.60
55:DA:630:G:OP2	30:D8:15:LYS:NZ	2.35	0.60
55:DA:118:A:H5'	55:DA:119:A:H8	1.67	0.60
55:DA:222:A:H3'	55:DA:421:U:H5''	1.83	0.60
55:DA:2347:C:P	28:D6:39:TYR:HH	2.24	0.60
55:DA:2875:C:H4'	15:DR:5:ALA:HB2	1.81	0.60
55:DA:878:A:N3	55:DA:878:A:H5''	2.16	0.60
3:DD:10:THR:HG23	3:DD:13:ARG:HB3	1.84	0.60
8:DK:60:GLU:HG3	8:DK:61:ARG:CZ	2.31	0.60
8:DK:74:ASN:ND2	8:DK:75:LEU:N	2.46	0.60
21:DV:23:LYS:HE2	21:DV:40:ASP:OD2	2.01	0.60
57:DY:73:GLY:HA2	57:DY:119:ALA:C	2.21	0.60
28:A6:9:LEU:HD22	28:A6:11:LEU:HD22	1.82	0.60
28:A6:25:LYS:NZ	28:A6:27:LYS:HD2	2.16	0.60
1:AA:1005:C:H2'	1:AA:1006:C:C6	2.37	0.60
1:AA:1497:U:H5''	1:AA:1498:C:OP2	2.02	0.60
1:AA:2820:A:N3	13:A0:4:LEU:HD23	2.16	0.60
2:AB:12:C:H4'	2:AB:13:A:H5''	1.83	0.60
3:AD:134:ARG:HG3	3:AD:135:PHE:CE2	2.37	0.60
4:AE:111:ARG:HD2	4:AE:118:LYS:HE3	1.83	0.60
7:AH:117:PRO:HB3	7:AH:123:PHE:HE1	1.67	0.60
8:AK:98:ALA:HA	8:AK:109:ILE:HD11	1.83	0.60
9:AM:99:LEU:O	9:AM:103:VAL:HG23	2.02	0.60
10:AN:69:ILE:HD12	10:AN:69:ILE:N	2.16	0.60
1:AA:2009:G:OP1	18:AS:41:LYS:HE3	2.02	0.60
18:AS:65:LEU:HD21	18:AS:67:ASP:OD2	2.01	0.60
18:AS:88:ARG:NH1	18:AS:94:ASP:OD1	2.34	0.60
20:AU:17:SER:HB2	20:AU:71:LYS:HD2	1.84	0.60
21:AV:178:GLU:C	21:AV:180:VAL:N	2.55	0.60
21:AV:80:ARG:O	21:AV:81:ARG:HG3	2.01	0.60
24:AW:26:ARG:HB3	24:AW:26:ARG:HH11	1.67	0.60
31:BA:1023:G:H2'	31:BA:1024:G:OP1	2.01	0.60
31:BA:37:U:O2'	31:BA:38:G:H5'	2.01	0.60
31:BA:951:G:H1'	31:BA:970:C:O2'	2.01	0.60
34:BG:24:GLU:H	34:BG:27:TYR:CB	2.14	0.60
34:BG:3:ARG:HB2	34:BG:3:ARG:NH2	2.16	0.60
39:BL:47:LEU:HD12	39:BL:47:LEU:N	2.17	0.60
43:BP:67:GLU:HG3	43:BP:68:GLY:H	1.66	0.60
33:BF:30:ARG:HD3	44:BQ:35:ARG:O	2.01	0.60
48:BU:23:LYS:HD2	48:BU:23:LYS:C	2.22	0.60
54:CA:1170:A:H2'	54:CA:1171:G:O4'	2.02	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1327:C:H2'	54:CA:1328:C:C6	2.36	0.60
54:CA:324:G:N2	54:CA:326:G:H3'	2.16	0.60
32:CE:122:PHE:HA	32:CE:139:LYS:NZ	2.17	0.60
34:CG:25:ARG:NH1	34:CG:30:LYS:HG3	2.16	0.60
35:CH:33:VAL:HG12	35:CH:112:LEU:HD12	1.83	0.60
37:CJ:26:PHE:O	37:CJ:30:ILE:HG12	2.01	0.60
45:CR:54:ARG:HH11	45:CR:54:ARG:HG2	1.66	0.60
50:CW:96:GLY:O	50:CW:97:ALA:HB3	2.01	0.60
13:D0:87:TYR:CE1	13:D0:118:GLU:HB3	2.36	0.60
13:D0:38:VAL:HB	13:D0:39:PRO:HD3	1.83	0.60
16:D1:58:ARG:HA	16:D1:61:TRP:CE3	2.37	0.60
27:D5:33:CYS:HB2	27:D5:40:LYS:CD	2.23	0.60
55:DA:1309:G:H4'	29:D7:7:PRO:HB2	1.84	0.60
55:DA:2165:G:N3	55:DA:2165:G:H2'	2.16	0.60
55:DA:386:G:H4'	55:DA:387:U:OP2	2.02	0.60
55:DA:894:C:H2'	55:DA:895:U:H6	1.65	0.60
55:DA:893:C:H3'	55:DA:894:C:H5	1.63	0.60
2:DB:104:A:H2'	2:DB:105:G:O4'	2.01	0.60
2:DB:77:U:P	21:DV:19:ARG:HH22	2.25	0.60
3:DD:130:ALA:C	3:DD:131:LEU:HD12	2.22	0.60
43:CP:7:VAL:CB	6:DG:115:ARG:HH22	2.14	0.60
7:DH:8:PRO:HG2	7:DH:69:ARG:HE	1.66	0.60
56:DI:4:ASP:HA	56:DI:8:ILE:CG1	2.32	0.60
58:DL:20:ALA:HB3	58:DL:21:PRO:CD	2.31	0.60
10:DN:104:ARG:HG2	10:DN:104:ARG:HH11	1.66	0.60
19:DT:8:ILE:N	19:DT:8:ILE:HD12	2.17	0.60
20:DU:74:PRO:O	20:DU:80:GLY:HA2	2.01	0.60
21:DV:178:GLU:C	21:DV:179:ASP:OD1	2.39	0.60
55:DA:1075:C:H5''	21:DV:195:GLU:CD	2.22	0.60
57:DY:25:PHE:CD1	57:DY:82:PHE:CD1	2.82	0.60
23:DZ:53:VAL:HB	23:DZ:58:ILE:HD12	1.82	0.60
1:AA:1454:U:H1'	13:A0:60:LEU:HD11	1.83	0.60
1:AA:2065:C:H2'	1:AA:2066:C:H6	1.66	0.60
1:AA:2581:G:H4'	1:AA:2582:G:N7	2.14	0.60
1:AA:2693:A:H2'	1:AA:2694:G:C8	2.37	0.60
1:AA:2842:G:O2'	1:AA:2843:G:H5'	2.01	0.60
1:AA:2867:G:HO2'	1:AA:2868:A:H8	1.46	0.60
1:AA:2:G:O2'	1:AA:3:U:H5'	2.01	0.60
1:AA:557:U:H2'	1:AA:558:G:H8	1.66	0.60
1:AA:654(F):C:C2'	1:AA:654(G):C:OP1	2.50	0.60
3:AD:10:THR:HG23	3:AD:13:ARG:HB2	1.82	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:AM:89:LYS:NZ	9:AM:89:LYS:HB3	2.17	0.60
11:AO:29:LYS:HD2	11:AO:30:THR:HG23	1.84	0.60
14:AQ:52:SER:O	14:AQ:56:LEU:CD2	2.49	0.60
15:AR:136:GLN:HG2	15:AR:136:GLN:O	2.01	0.60
18:AS:92:ARG:HH11	18:AS:92:ARG:HG2	1.66	0.60
20:AU:56:PRO:HB2	20:AU:57:GLN:HE22	1.66	0.60
20:AU:87:LYS:CB	20:AU:92:ASN:HB3	2.28	0.60
24:AW:10:LEU:O	24:AW:14:ARG:HB2	2.02	0.60
23:AZ:53:VAL:HG22	23:AZ:74:VAL:HG13	1.84	0.60
31:BA:1158:C:C2	31:BA:1160:G:N7	2.69	0.60
31:BA:1176:A:C6	31:BA:1177:G:C5	2.89	0.60
31:BA:191(F):U:H5'	31:BA:191(F):U:H6	1.66	0.60
31:BA:73:G:H2'	31:BA:74:C:C6	2.36	0.60
22:A3:5:LYS:HD2	52:BC:74:C:H5	1.65	0.60
32:BE:88:ALA:HB2	32:BE:219:VAL:HG23	1.84	0.60
42:BO:89:ARG:CG	42:BO:90:VAL:H	2.14	0.60
46:BS:5:ARG:NH2	46:BS:24:ALA:HA	2.16	0.60
47:BT:45:HIS:O	47:BT:73:VAL:HG12	2.02	0.60
54:CA:279:A:H5''	54:CA:280:C:H3'	1.84	0.60
54:CA:38:G:N3	54:CA:397:A:C2	2.70	0.60
34:CG:196:LEU:N	34:CG:196:LEU:HD12	2.12	0.60
55:DA:1077:A:N3	55:DA:1077:A:H2'	2.17	0.60
55:DA:2053:G:H5'	4:DE:144:ARG:O	2.00	0.60
55:DA:2426:A:O2'	55:DA:2427:C:OP1	2.19	0.60
55:DA:2665:A:O2'	55:DA:2666:C:H5'	2.01	0.60
55:DA:492:A:C2'	55:DA:493:G:H5'	2.32	0.60
7:DH:3:ARG:HA	7:DH:3:ARG:NE	2.17	0.60
56:DI:24:ILE:O	56:DI:26:ALA:N	2.35	0.60
56:DJ:20:LEU:HA	56:DJ:23:LEU:HB3	1.84	0.60
56:DJ:25:ASP:O	56:DJ:29:GLU:OE1	2.19	0.60
8:DK:133:HIS:CB	8:DK:134:PRO:HD2	2.31	0.60
9:DM:134:ARG:O	9:DM:136:GLU:N	2.35	0.60
10:DN:36:GLY:HA3	10:DN:109:LYS:HG3	1.83	0.60
19:DT:67:GLY:O	19:DT:69:TYR:N	2.34	0.60
21:DV:106:GLY:O	21:DV:107:THR:HG23	2.02	0.60
21:DV:120:ILE:O	21:DV:171:ILE:HA	2.00	0.60
21:DV:175:VAL:HB	21:DV:176:PRO:CA	2.32	0.60
57:DY:26:LEU:O	57:DY:111:LEU:N	2.30	0.60
17:A2:20:LEU:O	17:A2:93:GLU:HA	2.02	0.60
26:A4:10:VAL:HG13	26:A4:11:PRO:HD2	1.82	0.60
1:AA:2285:C:N4	28:A6:27:LYS:HE3	2.17	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:A8:32:LEU:CD2	30:A8:34:TRP:N	2.62	0.60
1:AA:1248:G:C4	16:A1:3:ARG:HG3	2.37	0.60
1:AA:2035:G:H4'	1:AA:2036:C:OP2	2.02	0.60
1:AA:2050:C:H1'	4:AE:156:MET:CE	2.32	0.60
1:AA:2505:G:O2'	1:AA:2506:U:C6	2.55	0.60
1:AA:752:A:C5	1:AA:1781:C:O4'	2.55	0.60
1:AA:932:G:H4'	1:AA:933:A:O5'	2.01	0.60
3:AD:91:ARG:O	3:AD:107:ALA:HB3	2.01	0.60
4:AE:11:MET:HA	4:AE:24:THR:HA	1.82	0.60
4:AE:27:LEU:HB2	4:AE:181:LEU:HD13	1.84	0.60
7:AH:152:ARG:O	7:AH:153:LYS:HB2	2.01	0.60
9:AM:38:HIS:ND1	9:AM:39:ARG:N	2.49	0.60
20:AU:86:ARG:HB3	20:AU:88:LYS:HZ1	1.67	0.60
21:AV:146:ILE:HG23	21:AV:147:GLY:N	2.00	0.60
21:AV:154:ASP:O	21:AV:155:LEU:O	2.19	0.60
21:AV:125:LEU:HG	21:AV:164:ALA:HB3	1.82	0.60
31:BA:926:G:N1	53:B1:45:U:H2'	2.16	0.60
31:BA:1085:U:H4'	31:BA:1086:U:OP1	2.01	0.60
31:BA:1380:U:H5''	31:BA:1381:U:OP1	2.01	0.60
31:BA:965:A:O2'	31:BA:966:G:H5'	2.01	0.60
34:BG:20:TYR:HB3	34:BG:27:TYR:CE1	2.36	0.60
31:BA:8:A:O2'	35:BH:103:GLY:N	2.35	0.60
40:BM:6:ILE:HA	40:BM:97:GLU:O	2.01	0.60
44:BQ:12:ARG:CZ	44:BQ:14:PRO:HG3	2.32	0.60
54:CA:1129:C:C5'	54:CA:1130:A:H5'	2.32	0.60
54:CA:74:C:H6	54:CA:74:C:O5'	1.85	0.60
33:CF:180:ALA:O	33:CF:181:ASN:HB3	2.02	0.60
34:CG:135:LEU:H	34:CG:135:LEU:HD22	1.66	0.60
54:CA:619:U:O2	34:CG:135:LEU:HD21	2.02	0.60
34:CG:11:LEU:C	34:CG:13:ARG:H	2.05	0.60
38:CK:6:ILE:N	38:CK:6:ILE:HD12	2.17	0.60
38:CK:91:ARG:HH11	38:CK:91:ARG:CG	2.15	0.60
40:CM:74:ILE:HD13	40:CM:74:ILE:N	2.17	0.60
41:CN:79:SER:CB	41:CN:106:LYS:HD2	2.28	0.60
55:DA:1299:G:H5''	55:DA:1300:U:OP1	2.01	0.60
55:DA:1332:G:N2	55:DA:1610:A:H8	2.00	0.60
55:DA:182:A:H2'	55:DA:183:C:O4'	2.01	0.60
55:DA:2848:G:O2'	55:DA:2849:U:O5'	2.19	0.60
55:DA:654(S):G:O3'	55:DA:654(T):A:H8	1.84	0.60
55:DA:774:A:H2	55:DA:787:U:O2'	1.84	0.60
5:DF:134:GLY:H	5:DF:162:LEU:HD22	1.65	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:DI:9:LYS:O	56:DI:11:GLU:C	2.39	0.60
10:DN:98:VAL:HG13	10:DN:117:LEU:HB2	1.82	0.60
10:DN:4:PRO:O	10:DN:5:GLN:CB	2.50	0.60
12:DP:108:GLY:O	12:DP:109:VAL:HG23	2.02	0.60
57:DY:115:GLN:CG	57:DY:115:GLN:O	2.50	0.60
57:DY:29:TYR:CZ	57:DY:32:LEU:HD11	2.33	0.60
1:AA:1008:C:N4	1:AA:1136:G:N1	2.50	0.60
1:AA:1111:A:O2'	1:AA:1112:G:H4'	2.02	0.60
1:AA:1161:C:O2'	1:AA:1162:G:H5'	2.02	0.60
1:AA:1286:A:C2'	1:AA:1288:U:OP2	2.49	0.60
1:AA:2494:G:OP1	22:A3:2:ALA:O	2.20	0.60
1:AA:2687:U:C4	1:AA:2688:U:C5	2.90	0.60
4:AE:134:ILE:O	4:AE:134:ILE:HG12	2.01	0.60
4:AE:31:CYS:HB3	4:AE:49:LEU:HB3	1.82	0.60
9:AM:120:LEU:HD23	9:AM:120:LEU:C	2.21	0.60
11:AO:42:SER:O	11:AO:43:GLY:C	2.40	0.60
20:AU:9:LYS:O	20:AU:27:VAL:HG22	2.02	0.60
20:AU:43:ASN:H	20:AU:43:ASN:HD22	1.49	0.60
20:AU:75:ILE:HB	20:AU:80:GLY:N	2.16	0.60
24:AW:23:LYS:O	24:AW:27:GLU:HG3	2.02	0.60
53:B1:31:A:H2'	53:B1:32:A:C8	2.37	0.60
31:BA:135:C:O2	46:BS:1:MET:HB3	2.01	0.60
31:BA:327:A:O2'	31:BA:329:A:O4'	2.13	0.60
31:BA:359:U:H2'	31:BA:360:A:C8	2.36	0.60
36:BI:3:ARG:HH22	36:BI:36:ARG:HH22	1.50	0.60
42:BO:18:VAL:O	42:BO:19:ARG:CB	2.50	0.60
43:BP:84:ILE:HG22	43:BP:85:GLY:H	1.66	0.60
54:CA:1053:G:O6	54:CA:1199:U:H2'	2.02	0.60
54:CA:243:A:H4'	54:CA:244:U:H5''	1.83	0.60
54:CA:948:C:C5	43:CP:106:ASN:ND2	2.69	0.60
32:CE:93:VAL:HG11	32:CE:97:TRP:HD1	1.66	0.60
33:CF:23:TYR:CG	33:CF:24:ALA:N	2.70	0.60
33:CF:92:ALA:HB2	33:CF:99:VAL:HG22	1.84	0.60
34:CG:63:LYS:HD3	34:CG:197:PRO:O	2.01	0.60
35:CH:11:ILE:HG13	35:CH:31:LEU:HD12	1.82	0.60
43:CP:116:THR:C	43:CP:117:VAL:CG1	2.70	0.60
49:CV:15:LEU:H	49:CV:15:LEU:CD2	2.13	0.60
49:CV:86:GLU:OE2	49:CV:86:GLU:CA	2.48	0.60
50:CW:98:PRO:O	50:CW:100:ILE:N	2.32	0.60
55:DA:1735:C:C5'	55:DA:1735:C:H6	2.14	0.60
55:DA:2790:A:H2	55:DA:2894:G:C5'	2.13	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:887:A:OP2	55:DA:887:A:O4'	2.19	0.60
2:DB:65:C:O2'	2:DB:66:A:H5'	2.02	0.60
4:DE:51:PHE:HD1	4:DE:52:LEU:CD1	2.14	0.60
5:DF:167:ALA:HB1	5:DF:173:VAL:HG11	1.83	0.60
6:DG:131:TYR:HB3	6:DG:159:VAL:CG1	2.32	0.60
6:DG:15:VAL:HG13	6:DG:175:LEU:HB2	1.84	0.60
12:DP:20:ALA:HB1	12:DP:99:PRO:HB2	1.84	0.60
15:DR:102:ILE:HA	15:DR:105:LEU:HD21	1.83	0.60
15:DR:16:ARG:HD3	15:DR:19:LEU:HG	1.82	0.60
15:DR:34:VAL:HG12	15:DR:35:LYS:N	2.16	0.60
15:DR:16:ARG:HH12	15:DR:81:PRO:HA	1.67	0.60
20:DU:47:LYS:C	20:DU:49:VAL:H	2.05	0.60
21:DV:52:SER:O	21:DV:54:HIS:N	2.35	0.60
25:DX:26:LEU:HD21	25:DX:46:ASN:HB2	1.84	0.60
57:DY:128:LEU:HA	57:DY:129:PRO:HB3	1.82	0.60
57:DY:96:PHE:O	57:DY:97:ALA:CB	2.50	0.60
26:A4:15:ILE:N	26:A4:15:ILE:HD12	2.16	0.60
28:A6:35:GLU:O	28:A6:36:LEU:HB2	2.02	0.60
29:A7:47:ARG:H	29:A7:47:ARG:HH11	1.49	0.60
1:AA:1082:U:H3'	1:AA:1082:U:C6	2.37	0.60
1:AA:1140:C:H5'	1:AA:1143:A:N6	2.17	0.60
1:AA:1579:A:H8	1:AA:1579:A:H5'	1.66	0.60
1:AA:2312:U:H6	1:AA:2312:U:O5'	1.85	0.60
1:AA:2341:G:H2'	1:AA:2342:C:H6	1.67	0.60
1:AA:363(C):G:H2'	1:AA:363(D):G:O4'	2.02	0.60
1:AA:846:C:H4'	1:AA:847:U:O5'	2.01	0.60
5:AF:150:GLY:HA2	5:AF:172:TRP:CE3	2.37	0.60
12:AP:42:ILE:HD13	12:AP:97:VAL:CG2	2.32	0.60
14:AQ:49:VAL:HG12	14:AQ:73:LEU:HD22	1.84	0.60
35:BH:110:LEU:O	35:BH:115:VAL:HG22	2.02	0.60
37:BJ:73:MET:HA	37:BJ:91:VAL:HG23	1.82	0.60
54:CA:1004:A:O4'	54:CA:1036:G:C6	2.55	0.60
54:CA:265:G:H2'	54:CA:267:C:H5	1.66	0.60
54:CA:703:G:O2'	54:CA:704:A:P	2.60	0.60
54:CA:789:U:C2	54:CA:791:G:OP2	2.55	0.60
52:CB:5:G:H2'	52:CB:6:G:H8	1.67	0.60
32:CE:17:PHE:CB	32:CE:44:LEU:HD11	2.32	0.60
32:CE:7:VAL:HG21	32:CE:217:ARG:HH11	1.65	0.60
34:CG:59:ARG:NH2	34:CG:62:GLN:HG3	2.17	0.60
34:CG:96:LEU:HD12	34:CG:139:ARG:NH1	2.17	0.60
35:CH:122:GLU:OE1	35:CH:131:ILE:HG13	2.01	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:CM:27:ALA:HB3	40:CM:34:VAL:HG21	1.83	0.60
43:CP:81:LEU:O	43:CP:84:ILE:HG22	2.02	0.60
43:CP:65:LYS:NZ	26:D4:52:THR:HB	2.17	0.60
28:D6:44:ARG:O	28:D6:45:LYS:CB	2.50	0.60
30:D8:54:GLU:O	30:D8:58:ILE:HG13	2.02	0.60
55:DA:1838:C:H4'	55:DA:1839:G:C8	2.36	0.60
55:DA:2023:G:H5'	55:DA:2617:C:H4'	1.84	0.60
55:DA:2312:U:O5'	55:DA:2312:U:H6	1.85	0.60
55:DA:2723:C:H5''	13:D0:1:MET:HG2	1.82	0.60
55:DA:304:G:H2'	55:DA:305:U:H6	1.67	0.60
6:DG:107:LEU:HD11	6:DG:178:PHE:CE1	2.37	0.60
56:DI:29:GLU:CA	56:DJ:2:ALA:CB	2.67	0.60
58:DL:57:ILE:HD12	58:DL:58:THR:O	2.02	0.60
58:DL:75:SER:O	58:DL:79:ARG:HD3	2.01	0.60
11:DO:119:GLU:HA	11:DO:119:GLU:OE1	2.01	0.60
14:DQ:42:ASP:O	14:DQ:43:GLU:HB2	1.99	0.60
21:DV:162:GLU:CG	21:DV:163:LEU:N	2.63	0.60
21:DV:82:ARG:HH11	21:DV:82:ARG:HG2	1.66	0.60
24:DW:18:PRO:O	24:DW:21:LEU:HB2	2.01	0.60
57:DY:23:SER:HB2	57:DY:67:GLY:O	2.02	0.60
16:A1:5:LYS:HB2	16:A1:5:LYS:HZ2	1.67	0.59
22:A3:43:THR:O	22:A3:43:THR:HG23	2.01	0.59
1:AA:1316:U:O2'	1:AA:1317:A:H5'	2.02	0.59
1:AA:1829:A:C8	1:AA:1830:C:C6	2.90	0.59
1:AA:2712:U:O2'	1:AA:2712(A):A:H3'	2.01	0.59
1:AA:278:A:H2'	1:AA:279:C:H6	1.67	0.59
1:AA:2807:G:H3'	1:AA:2808:U:H5''	1.83	0.59
4:AE:25:VAL:HA	4:AE:182:LEU:O	2.02	0.59
7:AH:117:PRO:HB3	7:AH:123:PHE:CE1	2.37	0.59
7:AH:136:ILE:N	7:AH:136:ILE:HD12	2.17	0.59
9:AM:15:LEU:HD13	9:AM:16:ILE:N	2.17	0.59
11:AO:62:LEU:HD12	30:A8:27:THR:HG22	1.82	0.59
20:AU:42:VAL:HG22	20:AU:65:ALA:HB3	1.83	0.59
20:AU:45:VAL:HA	20:AU:61:ILE:O	2.02	0.59
24:AW:65:ASN:ND2	24:AW:69:ARG:HH21	1.99	0.59
31:BA:411:A:H3'	31:BA:411:A:OP2	2.02	0.59
31:BA:535:A:H5''	31:BA:536:C:OP2	2.02	0.59
31:BA:628:G:O2'	31:BA:629:G:H5'	2.02	0.59
31:BA:939:G:H2'	31:BA:940:C:C6	2.37	0.59
52:BD:72:C:C2'	52:BD:73:A:H5''	2.30	0.59
33:BF:77:ILE:O	33:BF:83:ARG:HB3	2.02	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:BK:20:TYR:HD1	38:BK:65:TYR:CD2	2.19	0.59
31:BA:254:G:OP1	47:BT:67:LYS:O	2.19	0.59
49:BV:40:ILE:CG2	49:BV:67:VAL:HA	2.31	0.59
50:BW:23:ARG:O	50:BW:26:ASN:ND2	2.34	0.59
54:CA:1004:A:O5'	54:CA:1036:G:O6	2.20	0.59
54:CA:447:G:O6	54:CA:485:G:H2'	2.02	0.59
32:CE:194:PRO:O	32:CE:196:LEU:N	2.34	0.59
35:CH:41:VAL:HG13	35:CH:113:ALA:HB2	1.83	0.59
18:DS:15:ARG:HE	27:D5:20:ARG:CZ	2.15	0.59
28:D6:30:THR:N	28:D6:31:PRO:O	2.35	0.59
30:D8:29:LYS:HZ2	30:D8:44:LYS:HB2	1.65	0.59
55:DA:1803:A:H4'	3:DD:259:THR:CG2	2.31	0.59
55:DA:1805:U:O2	3:DD:50:THR:HB	2.02	0.59
55:DA:2698:U:H2'	55:DA:2699:C:C6	2.37	0.59
55:DA:2726:U:O2'	55:DA:2727:G:H5'	2.02	0.59
55:DA:2751:G:N3	7:DH:3:ARG:HB3	2.17	0.59
55:DA:481:G:O2'	55:DA:482:A:OP2	2.12	0.59
3:DD:17:THR:CG2	3:DD:204:ILE:HA	2.32	0.59
3:DD:94:LEU:HD11	3:DD:96:HIS:CE1	2.37	0.59
8:DK:95:LYS:CA	8:DK:111:PRO:HG3	2.27	0.59
8:DK:17:GLN:O	8:DK:18:VAL:HB	2.02	0.59
58:DL:63:ARG:HD3	58:DL:64:SER:O	2.02	0.59
58:DL:98:ARG:NH1	58:DL:98:ARG:N	2.50	0.59
10:DN:1:MET:HE3	10:DN:67:LYS:HG2	1.83	0.59
19:DT:54:VAL:HG22	19:DT:81:VAL:HB	1.84	0.59
21:DV:105:VAL:HG13	21:DV:140:ASP:HB3	1.82	0.59
21:DV:111:VAL:HG22	21:DV:145:GLU:HA	1.83	0.59
57:DY:128:LEU:HA	57:DY:129:PRO:CB	2.32	0.59
57:DY:141:VAL:CG2	57:DY:142:LEU:H	2.10	0.59
57:DY:71:LEU:C	57:DY:71:LEU:HD13	2.22	0.59
1:AA:559:G:H22	16:A1:49:HIS:CD2	2.20	0.59
1:AA:1341:U:H5''	19:AT:57:LEU:HD22	1.84	0.59
1:AA:1373:A:C6	1:AA:1374:G:C4	2.90	0.59
1:AA:2037:G:H2'	1:AA:2038:G:C8	2.37	0.59
1:AA:220:G:O2'	1:AA:233:A:N3	2.28	0.59
1:AA:391:G:H2'	1:AA:392:C:H6	1.67	0.59
5:AF:21:ALA:C	5:AF:23:ASP:H	2.05	0.59
31:BA:9:G:H2'	31:BA:10:A:C8	2.37	0.59
31:BA:164:U:H2'	31:BA:165:C:C6	2.37	0.59
31:BA:324:G:N2	31:BA:326:G:H3'	2.16	0.59
31:BA:757:U:H2'	31:BA:758:G:O4'	2.02	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:765:G:N2	31:BA:812:C:O2'	2.35	0.59
32:BE:12:GLU:HB3	32:BE:213:LEU:HD11	1.83	0.59
33:BF:90:GLU:O	33:BF:93:LYS:HB3	2.02	0.59
35:BH:78:HIS:HA	38:BK:105:ARG:HG3	1.84	0.59
38:BK:29:SER:HB3	38:BK:32:LYS:HD2	1.82	0.59
42:BO:84:LEU:HD22	42:BO:85:ILE:O	2.01	0.59
6:AG:147:ASP:HB3	43:BP:11:ARG:HH22	1.66	0.59
54:CA:1319:A:H5'	54:CA:1320:C:OP1	2.03	0.59
54:CA:265:G:H4'	47:CT:66:SER:HA	1.82	0.59
52:CB:57:G:H4'	21:DV:182:LYS:HZ3	1.67	0.59
52:CD:13:C:H2'	52:CD:14:A:H5'	1.83	0.59
52:CD:37:MIA:C11	52:CD:38:A:H1'	2.32	0.59
52:CD:56:C:N3	55:DA:2112:G:N2	2.49	0.59
32:CE:178:ARG:HH22	32:CE:196:LEU:C	2.05	0.59
32:CE:16:HIS:HD2	32:CE:210:SER:HA	1.68	0.59
32:CE:67:THR:C	32:CE:68:ILE:HD12	2.23	0.59
34:CG:188:LEU:HD23	34:CG:189:PRO:HD2	1.85	0.59
36:CI:99:ALA:HB1	48:CU:23:LYS:HZ2	1.64	0.59
48:CU:25:THR:O	48:CU:26:LEU:HD23	2.01	0.59
49:CV:24:ALA:C	49:CV:26:GLY:H	2.05	0.59
16:D1:108:GLU:CG	17:D2:44:LYS:HE3	2.33	0.59
17:D2:45:THR:HG22	17:D2:45:THR:O	2.02	0.59
17:D2:41:GLY:N	17:D2:46:VAL:HG13	2.17	0.59
22:D3:3:HIS:O	22:D3:4:LYS:CB	2.50	0.59
27:D5:40:LYS:NZ	27:D5:46:CYS:HB3	2.16	0.59
55:DA:1151:G:H5''	16:D1:81:HIS:CE1	2.37	0.59
55:DA:2250:G:C5	12:DP:82:ARG:HD3	2.36	0.59
55:DA:2377:A:H4'	14:DQ:111:GLU:O	2.02	0.59
55:DA:2823:A:OP1	4:DE:113:PHE:HB2	2.02	0.59
4:DE:131:ALA:HB1	4:DE:135:HIS:CE1	2.36	0.59
7:DH:151:ILE:O	7:DH:151:ILE:HG22	2.02	0.59
20:DU:47:LYS:HA	20:DU:60:PHE:HD1	1.67	0.59
21:DV:139:VAL:HG22	21:DV:155:LEU:HD22	1.85	0.59
21:DV:20:ARG:HG2	21:DV:20:ARG:NH1	2.10	0.59
57:DY:122:VAL:O	57:DY:125:LEU:N	2.36	0.59
57:DY:71:LEU:HA	57:DY:113:GLN:HA	1.83	0.59
1:AA:1106:G:H2'	1:AA:1107:G:H8	1.66	0.59
1:AA:1914:C:H2'	1:AA:1915:U:O4'	2.02	0.59
1:AA:671:C:H6	1:AA:671:C:H5'	1.67	0.59
5:AF:155:LEU:CD2	5:AF:186:ILE:HD13	2.30	0.59
5:AF:84:VAL:HG12	5:AF:85:GLY:N	2.16	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:AG:81:LYS:O	6:AG:82:LEU:HB2	2.02	0.59
15:AR:36:GLU:O	15:AR:36:GLU:HG3	2.02	0.59
21:AV:130:PRO:CA	21:AV:133:ILE:HD11	2.32	0.59
23:AZ:30:VAL:O	23:AZ:31:GLY:O	2.19	0.59
23:AZ:86:SER:N	23:AZ:87:PRO:HD2	2.17	0.59
31:BA:1020:U:O2'	31:BA:1021:G:H5''	2.02	0.59
31:BA:1067:A:H1'	31:BA:1068:G:C1'	2.31	0.59
31:BA:1148:U:C2'	31:BA:1149:C:H5'	2.33	0.59
31:BA:1492:A:H1'	53:B1:50:U:O2'	2.02	0.59
31:BA:408:A:O3'	34:BG:24:GLU:CD	2.40	0.59
31:BA:646:U:H2'	31:BA:647:C:H6	1.67	0.59
32:BE:142:LEU:HD23	32:BE:146:GLN:HB2	1.84	0.59
34:BG:112:VAL:HG22	34:BG:112:VAL:O	2.01	0.59
35:BH:12:LEU:HD23	35:BH:13:ILE:H	1.67	0.59
35:BH:32:VAL:HG12	35:BH:33:VAL:H	1.65	0.59
35:BH:69:VAL:O	35:BH:71:LEU:HG	2.03	0.59
37:BJ:71:PRO:HD3	37:BJ:103:TRP:HZ3	1.67	0.59
41:BN:21:ILE:HD13	41:BN:94:ALA:CB	2.32	0.59
46:BS:51:VAL:HG12	46:BS:52:ASP:N	2.17	0.59
26:A4:63:TYR:CE2	49:BV:41:VAL:CA	2.85	0.59
54:CA:1262:C:H2'	54:CA:1263:C:C6	2.37	0.59
54:CA:1299:A:C8	54:CA:1301:U:H1'	2.37	0.59
54:CA:1363:A:C4'	54:CA:1364:U:H5''	2.24	0.59
54:CA:194:C:C2'	54:CA:195:A:H5''	2.32	0.59
52:CB:68:C:H2'	52:CB:69:G:H8	1.67	0.59
52:CC:9:A:O2'	52:CC:10:G:N7	2.30	0.59
37:CJ:23:VAL:HG12	37:CJ:27:ILE:CD1	2.31	0.59
37:CJ:16:LEU:HD12	39:CL:42:ARG:HA	1.85	0.59
42:CO:51:ALA:O	42:CO:52:LEU:HG	2.02	0.59
43:CP:90:LEU:HD22	49:CV:78:ARG:NH2	2.12	0.59
44:CQ:15:LYS:HD2	44:CQ:16:PHE:CE2	2.37	0.59
49:CV:14:HIS:HD2	49:CV:35:SER:HB2	1.67	0.59
54:CA:261:U:C5	50:CW:79:ARG:NH1	2.69	0.59
49:CV:67:VAL:N	26:D4:59:PHE:HE1	1.97	0.59
27:D5:4:HIS:CB	27:D5:5:PRO:CD	2.77	0.59
55:DA:1904:G:O2'	55:DA:1905:C:H5'	2.02	0.59
55:DA:2197:U:O2'	55:DA:2198:A:H8	1.85	0.59
55:DA:222:A:O2'	55:DA:223:A:P	2.59	0.59
55:DA:278:A:O2'	55:DA:279:C:OP1	2.18	0.59
55:DA:55:G:N3	55:DA:127:A:H2	1.99	0.59
55:DA:811:U:OP2	11:DO:21:ARG:O	2.21	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:DD:155:LEU:HD23	3:DD:177:LEU:HD22	1.85	0.59
3:DD:206:LEU:HA	3:DD:211:ARG:HG2	1.83	0.59
5:DF:9:ILE:HD12	5:DF:123:LEU:CD2	2.32	0.59
6:DG:67:LYS:O	6:DG:67:LYS:HD2	2.02	0.59
7:DH:152:ARG:HE	7:DH:153:LYS:HE3	1.65	0.59
8:DK:2:LYS:HB3	8:DK:20:ASP:HB3	1.83	0.59
8:DK:38:LEU:N	8:DK:38:LEU:HD12	2.07	0.59
58:DL:123:ALA:HA	58:DL:126:MET:SD	2.42	0.59
58:DL:95:LYS:H	58:DL:136:VAL:CG1	2.15	0.59
12:DP:104:PHE:CE1	12:DP:125:LEU:HD11	2.35	0.59
12:DP:68:ILE:HD13	12:DP:103:MET:HE3	1.83	0.59
24:DW:43:GLN:O	24:DW:44:LEU:HG	2.02	0.59
17:A2:77:ALA:C	17:A2:78:LYS:HG2	2.22	0.59
26:A4:63:TYR:O	26:A4:63:TYR:CD2	2.56	0.59
29:A7:49:ARG:HD3	29:A7:49:ARG:OXT	2.02	0.59
1:AA:1771:C:H1'	1:AA:1786:A:C8	2.38	0.59
1:AA:426:C:O2'	1:AA:427:U:H5'	2.03	0.59
1:AA:458:G:C1'	1:AA:459:U:C5	2.80	0.59
1:AA:467:G:OP1	29:A7:33:ARG:NH1	2.36	0.59
1:AA:483:A:H5'	20:AU:49:VAL:HG22	1.84	0.59
1:AA:897:C:H2'	1:AA:898:C:C5'	2.32	0.59
5:AF:68:LYS:HB3	5:AF:69:HIS:CD2	2.37	0.59
8:AK:10:GLU:OE2	8:AK:11:ASN:HB2	2.01	0.59
8:AK:76:THR:C	8:AK:77:LEU:HD23	2.23	0.59
9:AM:46:VAL:O	9:AM:47:ALA:HB3	2.02	0.59
31:BA:1005:A:H5''	31:BA:1006:C:C6	2.37	0.59
31:BA:501:C:H2'	31:BA:502:G:H8	1.67	0.59
31:BA:947:G:H2'	31:BA:948:C:H6	1.67	0.59
31:BA:973:G:H1'	40:BM:55:LYS:CD	2.31	0.59
31:BA:975:A:H4'	31:BA:1358:U:H1'	1.84	0.59
52:BB:5:G:O2'	52:BB:6:G:H5'	2.02	0.59
33:BF:75:VAL:HG12	33:BF:75:VAL:O	2.01	0.59
37:BJ:21:VAL:HG23	37:BJ:22:LEU:H	1.68	0.59
40:BM:98:ILE:HD12	40:BM:98:ILE:N	2.16	0.59
48:BU:22:VAL:HG22	48:BU:23:LYS:H	1.67	0.59
49:BV:42:PRO:CA	49:BV:45:VAL:HG13	2.32	0.59
54:CA:1300:G:O2'	54:CA:1301:U:P	2.61	0.59
54:CA:1288:A:O4'	54:CA:1353:G:H4'	2.02	0.59
54:CA:1379:G:O6	37:CJ:2:ALA:HB3	2.01	0.59
54:CA:630:G:O2'	54:CA:631:G:H5''	2.00	0.59
52:CB:37:MIA:HN6	52:CB:37:MIA:C16	2.15	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:CD:2:C:C6	52:CD:2:C:H5''	2.32	0.59
32:CE:200:ILE:HD12	32:CE:200:ILE:N	2.13	0.59
34:CG:152:SER:HB3	34:CG:155:LEU:CD1	2.32	0.59
38:CK:112:LEU:HD12	38:CK:114:THR:HG23	1.83	0.59
40:CM:96:ILE:HD13	40:CM:96:ILE:H	1.67	0.59
48:CU:20:ALA:O	48:CU:22:VAL:N	2.35	0.59
48:CU:50:ILE:H	48:CU:50:ILE:CD1	2.13	0.59
22:D3:43:THR:O	22:D3:43:THR:HG23	2.00	0.59
55:DA:1021:A:C3'	55:DA:1021:A:C8	2.83	0.59
55:DA:1205:U:H4'	55:DA:1206:G:OP2	2.02	0.59
55:DA:1558:A:H4'	55:DA:1559:G:O5'	2.02	0.59
55:DA:2754:U:H5'	55:DA:2755:C:OP2	2.02	0.59
55:DA:2786:U:O2'	4:DE:62:PRO:HA	2.01	0.59
55:DA:498:G:H21	20:DU:47:LYS:NZ	1.99	0.59
55:DA:95:G:H4'	24:DW:46:GLN:HB3	1.85	0.59
5:DF:102:PRO:O	5:DF:106:ARG:HG2	2.01	0.59
6:DG:10:LYS:O	6:DG:15:VAL:HG23	2.02	0.59
6:DG:38:VAL:HG22	6:DG:93:THR:HG23	1.84	0.59
8:DK:57:ARG:HA	8:DK:60:GLU:HB3	1.83	0.59
58:DL:66:THR:O	58:DL:67:PHE:HB2	2.03	0.59
58:DL:93:ARG:C	58:DL:136:VAL:HG12	2.23	0.59
20:DU:76:CYS:SG	20:DU:77:PRO:CD	2.79	0.59
57:DY:75:GLN:HB3	57:DY:110:GLY:C	2.21	0.59
16:A1:108:GLU:OE1	17:A2:45:THR:HA	2.02	0.59
1:AA:1817:G:OP1	3:AD:88:ARG:NH2	2.30	0.59
1:AA:1:G:H2'	1:AA:2:G:C8	2.38	0.59
1:AA:2389:G:H5''	1:AA:2390:U:C5'	2.32	0.59
1:AA:2459:A:C2	1:AA:2460:U:H1'	2.37	0.59
1:AA:607:U:O4	1:AA:608:A:C5	2.54	0.59
1:AA:654(I):C:O2	1:AA:654(I):C:C2'	2.50	0.59
1:AA:919:G:H5'	2:AB:81:G:H1'	1.84	0.59
4:AE:13:ARG:HA	4:AE:21:VAL:O	2.02	0.59
7:AH:26:VAL:HG13	7:AH:27:LYS:N	2.18	0.59
8:AK:8:PRO:HD3	8:AK:15:VAL:CG2	2.32	0.59
1:AA:811:U:C4	11:AO:21:ARG:NH2	2.69	0.59
12:AP:127:ILE:HG22	12:AP:128:LYS:N	2.16	0.59
12:AP:14:ARG:HG2	12:AP:41:TRP:HH2	1.66	0.59
12:AP:89:ASN:O	12:AP:91:GLU:N	2.36	0.59
14:AQ:10:ARG:O	14:AQ:12:PHE:N	2.35	0.59
14:AQ:11:LYS:HD2	14:AQ:15:ARG:HH21	1.67	0.59
15:AR:74:ARG:CG	15:AR:74:ARG:HH11	2.16	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:AR:81:PRO:HD2	15:AR:82:LEU:HD12	1.84	0.59
1:AA:84:A:P	20:AU:8:LYS:HD3	2.42	0.59
31:BA:1285:A:O2'	31:BA:1286:A:OP2	2.18	0.59
31:BA:1390:U:H2'	31:BA:1391:U:C6	2.38	0.59
31:BA:209:U:O2'	31:BA:210:U:P	2.60	0.59
38:BK:88:LYS:HB3	38:BK:89:PRO:HD2	1.85	0.59
43:BP:90:LEU:CD1	49:BV:78:ARG:HH21	2.14	0.59
50:BW:70:SER:O	50:BW:73:HIS:HB2	2.03	0.59
51:BX:26:LYS:HZ3	51:BX:26:LYS:HA	1.65	0.59
54:CA:1032:A:H3'	54:CA:1032(A):G:C4'	2.32	0.59
54:CA:1106:G:H2'	54:CA:1107:C:H6	1.66	0.59
54:CA:135:C:H2'	54:CA:136:C:H5'	1.85	0.59
54:CA:245:C:O2'	54:CA:246:A:H5'	2.02	0.59
54:CA:346:G:H4'	15:DR:41:ARG:NH1	2.17	0.59
54:CA:882:C:O2'	54:CA:883:C:H5'	2.03	0.59
54:CA:913:A:H4'	54:CA:914:A:O5'	2.03	0.59
54:CA:968:A:H4'	54:CA:969:A:OP2	2.03	0.59
34:CG:119:GLN:NE2	34:CG:123:HIS:NE2	2.50	0.59
39:CL:22:GLY:N	39:CL:58:HIS:O	2.33	0.59
54:CA:267:C:OP1	47:CT:67:LYS:HB2	2.02	0.59
13:D0:9:LYS:O	13:D0:9:LYS:HG2	2.02	0.59
22:D3:36:ILE:H	22:D3:36:ILE:HD13	1.66	0.59
22:D3:50:ASN:HB2	22:D3:81:VAL:HB	1.83	0.59
55:DA:2285:C:N4	28:D6:27:LYS:HE2	2.17	0.59
55:DA:1030:G:OP2	12:DP:128:LYS:HE2	2.02	0.59
55:DA:1049:C:H1'	55:DA:1113:U:H4'	1.83	0.59
55:DA:1204:A:O2'	55:DA:1205:U:C5'	2.51	0.59
55:DA:284:U:H2'	55:DA:285:C:H6	1.65	0.59
55:DA:524:U:H2'	55:DA:525:U:C6	2.37	0.59
55:DA:673:C:OP1	5:DF:54:ARG:HD2	2.02	0.59
5:DF:155:LEU:CD1	5:DF:174:VAL:HG22	2.33	0.59
6:DG:161:THR:HG22	6:DG:162:THR:N	2.17	0.59
58:DL:52:ILE:HD12	58:DL:53:VAL:H	1.66	0.59
9:DM:30:ILE:HG22	9:DM:34:LEU:HD21	1.85	0.59
10:DN:104:ARG:NH1	15:DR:36:GLU:HG3	2.17	0.59
11:DO:96:THR:HG22	11:DO:126:VAL:HG23	1.83	0.59
20:DU:53:PRO:O	20:DU:54:LYS:C	2.39	0.59
21:DV:117:LEU:N	21:DV:117:LEU:CD1	2.66	0.59
21:DV:27:VAL:HG12	21:DV:87:ASP:HB3	1.84	0.59
21:DV:59:LEU:O	21:DV:61:LEU:N	2.35	0.59
57:DY:116:ILE:O	57:DY:116:ILE:HG13	2.02	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:138:LEU:C	57:DY:140:GLY:N	2.56	0.59
23:DZ:87:PRO:O	23:DZ:88:LYS:C	2.39	0.59
17:A2:64:HIS:HD2	17:A2:92:THR:HA	1.66	0.59
22:A3:27:GLU:OE1	22:A3:69:PHE:HB2	2.02	0.59
1:AA:1385:G:OP1	1:AA:1385:G:H4'	2.01	0.59
1:AA:1970:A:C5'	1:AA:1971:A:OP1	2.51	0.59
1:AA:2319:G:H1'	1:AA:2320:A:C4	2.38	0.59
1:AA:2376:A:H2	14:AQ:112:PHE:HB2	1.67	0.59
2:AB:81:G:C2	2:AB:82:G:C5	2.90	0.59
6:AG:77:ILE:O	6:AG:81:LYS:O	2.20	0.59
12:AP:25:ASP:OD1	12:AP:102:VAL:HB	2.02	0.59
12:AP:78:PRO:O	12:AP:79:LEU:CB	2.50	0.59
19:AT:63:LYS:HE3	19:AT:63:LYS:N	2.11	0.59
23:AZ:19:GLN:HB2	23:AZ:35:THR:O	2.02	0.59
31:BA:1158:C:O2'	32:BE:133:LYS:HE2	2.02	0.59
31:BA:1229:A:H2'	31:BA:1230:C:C6	2.38	0.59
31:BA:1319:A:H2'	31:BA:1323:G:N7	2.18	0.59
31:BA:559:A:H4'	31:BA:560:U:C5'	2.31	0.59
52:BC:44:G:H3'	52:BC:45:U:C6	2.38	0.59
32:BE:211:ILE:O	32:BE:215:LEU:HD23	2.02	0.59
33:BF:129:ALA:HB3	33:BF:132:ARG:HB3	1.85	0.59
31:BA:426:G:H4'	34:BG:41:GLY:O	2.02	0.59
37:BJ:21:VAL:HG23	37:BJ:22:LEU:N	2.17	0.59
31:BA:538:G:H3'	42:BO:115:LYS:NZ	2.17	0.59
6:AG:115:ARG:CB	43:BP:7:VAL:HG11	2.31	0.59
49:BV:33:THR:OG1	49:BV:34:TRP:N	2.36	0.59
50:BW:100:ILE:HD12	50:BW:100:ILE:N	2.18	0.59
50:BW:38:LYS:O	50:BW:41:ILE:HG12	2.02	0.59
54:CA:1323:G:H2'	54:CA:1324:A:H8	1.65	0.59
54:CA:971:G:N2	54:CA:1363:A:OP2	2.28	0.59
54:CA:992:U:O2'	54:CA:993:G:OP2	2.20	0.59
34:CG:117:ALA:O	34:CG:121:VAL:HG23	2.02	0.59
35:CH:10:MET:SD	35:CH:13:ILE:HD11	2.43	0.59
38:CK:64:LYS:HG2	38:CK:79:VAL:HG21	1.85	0.59
54:CA:881:G:P	42:CO:12:ARG:HH22	2.25	0.59
33:CF:13:GLY:HA3	44:CQ:57:ARG:NE	2.16	0.59
49:CV:81:ARG:HG2	49:CV:82:GLY:N	2.17	0.59
50:CW:17:ARG:HH11	50:CW:17:ARG:HG3	1.68	0.59
55:DA:1077:A:C1'	58:DL:93:ARG:HH22	2.15	0.59
55:DA:1062:G:H1'	55:DA:1088:A:C6	2.37	0.59
55:DA:228:A:C2'	55:DA:229:A:OP1	2.51	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2656:U:C5	55:DA:2664:G:N2	2.70	0.59
55:DA:2654:A:N9	55:DA:2656:U:O2	2.36	0.59
55:DA:458:G:O2'	55:DA:459:U:OP2	2.21	0.59
55:DA:894:C:O5'	55:DA:894:C:H6	1.86	0.59
7:DH:42:ARG:HG2	7:DH:42:ARG:HH11	1.67	0.59
7:DH:60:ARG:HG2	7:DH:60:ARG:HH11	1.68	0.59
56:DI:27:LEU:O	56:DI:29:GLU:HB2	2.02	0.59
58:DL:146:ASP:O	58:DL:147:ALA:CB	2.51	0.59
9:DM:115:ARG:O	9:DM:118:LYS:HB2	2.02	0.59
11:DO:19:VAL:HG22	11:DO:20:GLY:H	1.67	0.59
15:DR:27:THR:CG2	15:DR:90:GLN:HB3	2.27	0.59
18:DS:52:GLU:O	18:DS:55:ALA:HB3	2.03	0.59
19:DT:40:LYS:HG3	19:DT:51:VAL:HB	1.84	0.59
57:DY:104:ILE:CB	57:DY:105:PRO:HD2	2.31	0.59
57:DY:29:TYR:CD2	57:DY:32:LEU:HD21	2.37	0.59
1:AA:2882:A:C5'	13:A0:96:ARG:HG3	2.31	0.59
17:A2:87:HIS:HD1	17:A2:88:ARG:N	2.01	0.59
1:AA:1379:A:C2'	1:AA:1380:G:OP1	2.50	0.59
1:AA:2113:U:H3'	1:AA:2114:A:C5'	2.31	0.59
1:AA:2295:C:OP2	14:AQ:10:ARG:HD2	2.02	0.59
1:AA:2315:G:H2'	1:AA:2316:C:H6	1.66	0.59
1:AA:2748:A:H3'	1:AA:2748:A:N3	2.18	0.59
1:AA:2043:C:H1'	1:AA:2779:U:O4	2.02	0.59
1:AA:532:A:HO2'	1:AA:533:G:P	2.26	0.59
1:AA:654(C):G:H2'	1:AA:654(D):G:C1'	2.33	0.59
1:AA:78:A:H2'	1:AA:79:G:C8	2.38	0.59
1:AA:883:G:H2'	1:AA:884:C:C5	2.38	0.59
1:AA:914:C:C2'	1:AA:915:C:H5'	2.28	0.59
2:AB:81:G:H5'	2:AB:82:G:OP2	2.02	0.59
1:AA:2679:A:H5'	4:AE:165:VAL:HG11	1.84	0.59
11:AO:59:LEU:O	11:AO:61:ARG:HG2	2.03	0.59
1:AA:142:G:C1'	19:AT:37:THR:HG21	2.32	0.59
21:AV:20:ARG:HH11	21:AV:20:ARG:HG2	1.68	0.59
23:AZ:92:LYS:NZ	23:AZ:97:LEU:HG	2.16	0.59
33:BF:164:ARG:NE	53:B1:55:U:O4	2.36	0.59
31:BA:1112:C:C4	33:BF:178:LEU:HD23	2.38	0.59
31:BA:155:C:O2'	31:BA:156:G:H5'	2.02	0.59
31:BA:474:G:H5''	46:BS:81:ARG:HD3	1.84	0.59
31:BA:631:G:O2'	31:BA:632:A:OP1	2.19	0.59
31:BA:740:U:O2'	31:BA:741:G:H5'	2.02	0.59
32:BE:207:ALA:O	32:BE:211:ILE:HG13	2.02	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:BG:68:TYR:OH	34:BG:196:LEU:HD21	2.02	0.59
37:BJ:15:ASP:OD2	37:BJ:16:LEU:N	2.34	0.59
31:BA:599:C:H4'	38:BK:130:GLY:HA3	1.85	0.59
42:BO:90:VAL:O	42:BO:92:ASP:N	2.34	0.59
54:CA:1151:A:H2'	54:CA:1152:A:H8	1.64	0.59
54:CA:1254:C:H41	40:CM:43:ARG:HH12	1.50	0.59
54:CA:639:G:O2'	54:CA:640:A:H5'	2.03	0.59
54:CA:663:A:H2'	54:CA:664:G:O4'	2.01	0.59
32:CE:178:ARG:HD2	38:CK:71:GLY:C	2.23	0.59
32:CE:188:ALA:O	32:CE:202:PRO:HA	2.03	0.59
33:CF:79:ARG:HG2	33:CF:79:ARG:HH11	1.66	0.59
34:CG:156:GLU:HG2	34:CG:160:GLN:HE21	1.67	0.59
34:CG:98:GLU:OE1	34:CG:194:LEU:HD21	2.03	0.59
40:CM:6:ILE:CG2	40:CM:98:ILE:HG13	2.33	0.59
42:CO:28:LYS:O	42:CO:29:GLY:C	2.40	0.59
43:CP:116:THR:CG2	43:CP:117:VAL:H	2.15	0.59
43:CP:124:PRO:CB	43:CP:125:ARG:CA	2.81	0.59
33:CF:6:HIS:ND1	44:CQ:49:HIS:HB3	2.18	0.59
47:CT:59:ILE:HG21	47:CT:71:PHE:HB3	1.84	0.59
13:D0:67:LEU:HD12	13:D0:76:VAL:HG21	1.85	0.59
28:D6:17:LYS:O	28:D6:18:ARG:HB2	2.02	0.59
55:DA:54:G:O2'	29:D7:35:ARG:HD3	2.02	0.59
55:DA:1694:C:O2'	55:DA:1695:G:OP2	2.19	0.59
55:DA:2701:C:H3'	55:DA:2702:U:C5'	2.19	0.59
55:DA:868:U:C4	55:DA:869:G:N7	2.71	0.59
2:DB:95:U:C3'	2:DB:95:U:C6	2.86	0.59
4:DE:41:LYS:HE2	4:DE:41:LYS:HA	1.82	0.59
8:DK:29:TYR:C	8:DK:32:PRO:HD2	2.22	0.59
58:DL:101:TRP:C	58:DL:104:VAL:HB	2.23	0.59
58:DL:63:ARG:HE	58:DL:63:ARG:C	2.06	0.59
11:DO:115:LEU:CB	11:DO:131:SER:HB2	2.33	0.59
20:DU:81:LYS:NZ	20:DU:98:VAL:HG11	2.17	0.59
21:DV:176:PRO:O	21:DV:177:PRO:C	2.39	0.59
52:CB:57:G:H5''	21:DV:182:LYS:CE	2.32	0.59
21:DV:6:LYS:O	21:DV:7:ALA:HB2	2.02	0.59
57:DY:123:GLU:O	57:DY:127:GLU:HB2	2.01	0.59
9:AM:42:TRP:O	16:A1:64:ARG:NH2	2.36	0.59
26:A4:35:VAL:O	26:A4:37:SER:N	2.35	0.59
1:AA:1029:A:H5''	12:AP:128:LYS:HE2	1.84	0.59
1:AA:1810:A:H2'	1:AA:1811:G:C5'	2.33	0.59
1:AA:2147:G:H3'	1:AA:2147:G:H8	1.68	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2741:A:H2'	1:AA:2742:C:O4'	2.01	0.59
1:AA:2758:A:C3'	1:AA:2759:G:H5''	2.32	0.59
1:AA:331:A:O2'	1:AA:332:A:OP1	2.17	0.59
1:AA:633:A:H2'	1:AA:634:C:H5'	1.83	0.59
1:AA:864:G:H2'	1:AA:865:C:H6	1.67	0.59
1:AA:925:C:C3'	1:AA:926:A:H5''	2.32	0.59
2:AB:15:A:H5'	2:AB:16:G:H8	1.68	0.59
5:AF:184:TYR:O	5:AF:188:ARG:HB2	2.02	0.59
5:AF:25:PRO:O	5:AF:26:ALA:HB3	2.03	0.59
8:AK:124:GLY:O	8:AK:141:LYS:HA	2.02	0.59
11:AO:79:ARG:O	11:AO:111:ARG:HB2	2.03	0.59
15:AR:56:GLY:O	15:AR:59:THR:HG22	2.02	0.59
21:AV:183:LEU:O	21:AV:184:ALA:CB	2.51	0.59
31:BA:1027:C:H2'	31:BA:1028:C:C5	2.38	0.59
31:BA:1181:G:C5	31:BA:1182:G:N2	2.71	0.59
31:BA:1218:C:OP2	44:BQ:9:LYS:NZ	2.33	0.59
31:BA:137:C:H1'	46:BS:63:GLY:HA3	1.84	0.59
31:BA:1452:C:H1'	31:BA:1454:G:C4	2.38	0.59
31:BA:1535:C:C2'	31:BA:1536:C:H5'	2.33	0.59
31:BA:745:C:H2'	31:BA:746:A:C8	2.37	0.59
52:BB:69:G:C2'	52:BB:70:G:H5''	2.29	0.59
33:BF:116:VAL:HG11	33:BF:141:VAL:HG21	1.85	0.59
36:BI:33:TYR:OH	36:BI:78:GLU:HG3	2.03	0.59
31:BA:1216:G:OP1	44:BQ:2:ALA:HA	2.02	0.59
51:BX:9:ARG:NH1	51:BX:13:ILE:HD12	2.17	0.59
53:C1:31:A:O2'	53:C1:32:A:P	2.60	0.59
54:CA:951:G:O2'	54:CA:972:C:H5	1.85	0.59
52:CB:11:C:H2'	52:CB:12:U:C6	2.38	0.59
54:CA:921:U:O2	35:CH:19:MET:HB2	2.02	0.59
40:CM:54:PHE:CZ	40:CM:55:LYS:NZ	2.70	0.59
43:CP:12:ASN:HA	43:CP:46:LYS:HB2	1.85	0.59
50:CW:59:ALA:HA	50:CW:62:LEU:HD12	1.83	0.59
16:D1:5:LYS:HB2	16:D1:5:LYS:NZ	2.17	0.59
26:D4:13:ARG:O	26:D4:14:ILE:HB	2.03	0.59
49:CV:42:PRO:HD3	26:D4:63:TYR:CE2	2.37	0.59
30:D8:17:THR:CG2	30:D8:21:LYS:HB2	2.32	0.59
30:D8:22:VAL:HG21	30:D8:53:PRO:HB2	1.85	0.59
55:DA:1061:U:H5''	55:DA:1070:A:O2'	2.03	0.59
55:DA:1278:A:O2'	13:D0:34:ILE:HD12	2.02	0.59
55:DA:1379:A:HO2'	55:DA:1380:G:P	2.25	0.59
55:DA:1962:C:O2'	55:DA:1964:G:OP2	2.20	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:218:A:H2	55:DA:235:U:H4'	1.68	0.59
55:DA:590:A:H2'	55:DA:591:C:C6	2.37	0.59
3:DD:10:THR:HG23	3:DD:13:ARG:HB2	1.85	0.59
6:DG:34:LEU:HD13	6:DG:34:LEU:O	2.03	0.59
7:DH:24:VAL:O	7:DH:24:VAL:HG23	2.02	0.59
56:DI:24:ILE:C	56:DI:26:ALA:N	2.51	0.59
56:DJ:10:GLU:OE1	56:DJ:10:GLU:O	2.20	0.59
58:DL:106:GLU:OE1	58:DL:109:LYS:HD2	2.02	0.59
58:DL:7:VAL:HG12	58:DL:8:VAL:H	1.66	0.59
10:DN:87:ILE:HD12	10:DN:91:LEU:HD12	1.85	0.59
11:DO:112:LEU:HD22	11:DO:113:LYS:N	2.18	0.59
57:DY:23:SER:O	57:DY:25:PHE:N	2.36	0.59
57:DY:48:GLY:O	57:DY:49:ALA:C	2.40	0.59
57:DY:95:GLN:O	57:DY:96:PHE:CD2	2.56	0.59
17:A2:22:VAL:CG2	17:A2:23:GLU:H	2.15	0.59
28:A6:13:CYS:O	28:A6:21:TYR:HB3	2.03	0.59
1:AA:1010:A:H5'	16:A1:62:ILE:HG21	1.84	0.59
1:AA:1141:U:H5'	9:AM:25:ARG:HH21	1.68	0.59
1:AA:1963:U:O2	1:AA:1963:U:H2'	2.03	0.59
1:AA:2225:A:H4'	1:AA:2226:C:O5'	2.01	0.59
1:AA:419:C:H2'	1:AA:420:C:O4'	2.02	0.59
5:AF:128:ALA:O	5:AF:130:ALA:N	2.36	0.59
5:AF:18:ARG:O	5:AF:19:GLU:HB3	2.02	0.59
5:AF:63:LYS:NZ	5:AF:67:GLN:HB2	2.18	0.59
6:AG:114:ILE:HD13	6:AG:140:ILE:HG21	1.85	0.59
7:AH:87:LEU:HD21	7:AH:149:ARG:HB2	1.85	0.59
7:AH:20:ALA:O	7:AH:22:GLY:N	2.35	0.59
21:AV:182:LYS:HD3	21:AV:183:LEU:N	2.18	0.59
31:BA:1067:A:C1'	31:BA:1068:G:O4'	2.50	0.59
31:BA:1246:C:H2'	31:BA:1247:U:C6	2.38	0.59
31:BA:687:A:H61	31:BA:703:G:H1'	1.68	0.59
31:BA:877:C:O2'	31:BA:878:G:H5'	2.03	0.59
32:BE:7:VAL:HG13	32:BE:8:LYS:N	2.18	0.59
33:BF:53:ALA:HB2	33:BF:115:LEU:CD2	2.33	0.59
34:BG:14:ARG:HG3	34:BG:14:ARG:HH11	1.66	0.59
38:BK:23:SER:HA	38:BK:63:LEU:HD22	1.85	0.59
40:BM:54:PHE:CZ	40:BM:55:LYS:HD2	2.37	0.59
40:BM:82:ILE:HG22	40:BM:86:MET:SD	2.42	0.59
41:BN:110:ASP:HB3	48:BU:85:LEU:HD21	1.84	0.59
54:CA:1181:G:C4	54:CA:1182:G:N2	2.71	0.59
54:CA:765:G:N1	54:CA:812:C:H2'	2.18	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:911:U:H2'	54:CA:912:C:C6	2.38	0.59
34:CG:79:PHE:CE1	34:CG:207:TYR:HB2	2.37	0.59
35:CH:76:ILE:HB	35:CH:77:PRO:HD2	1.85	0.59
37:CJ:111:ARG:HH12	37:CJ:113:GLU:CD	2.07	0.59
42:CO:18:VAL:HG23	42:CO:19:ARG:N	2.10	0.59
43:CP:3:ARG:NE	43:CP:7:VAL:HG13	2.17	0.59
9:DM:42:TRP:C	16:D1:64:ARG:HH22	2.05	0.59
55:DA:1061:U:C5'	55:DA:1070:A:H1'	2.31	0.59
55:DA:1077:A:C4	55:DA:1078:U:H5''	2.36	0.59
55:DA:1313:U:H3'	55:DA:1314:C:H5'	1.84	0.59
55:DA:1360:A:H2'	55:DA:1361:G:H5'	1.85	0.59
55:DA:1412:A:H2'	55:DA:1413:G:C8	2.38	0.59
2:DB:50:G:OP2	14:DQ:62:LYS:HB2	2.02	0.59
4:DE:197:ILE:HD11	4:DE:199:ARG:CZ	2.32	0.59
5:DF:197:ASP:O	5:DF:198:ALA:HB3	2.03	0.59
43:CP:7:VAL:CG1	6:DG:115:ARG:HH22	2.16	0.59
7:DH:19:VAL:HG12	7:DH:20:ALA:H	1.64	0.59
56:DI:7:ARG:NE	56:DI:8:ILE:CD1	2.64	0.59
56:DJ:24:ILE:CG2	56:DJ:25:ASP:N	2.64	0.59
57:DY:18:GLU:CG	57:DY:66:LEU:HD11	2.32	0.59
13:A0:10:LEU:O	13:A0:12:ARG:HG3	2.03	0.59
30:A8:32:LEU:HB2	30:A8:36:LYS:NZ	2.18	0.59
1:AA:1348:G:C2'	1:AA:1349:A:H5''	2.33	0.59
1:AA:1716:U:O2'	1:AA:1717:G:H5'	2.03	0.59
1:AA:2113:U:H5'	1:AA:2114:A:C8	2.37	0.59
3:AD:25:THR:O	3:AD:27:THR:N	2.35	0.59
3:AD:44:ASN:HB2	3:AD:48:ARG:O	2.02	0.59
4:AE:8:LYS:CG	4:AE:192:ASN:HA	2.32	0.59
6:AG:111:LEU:HD11	6:AG:120:LEU:HD11	1.85	0.59
9:AM:73:THR:CG2	9:AM:84:LYS:HB3	2.30	0.59
15:AR:50:ILE:HD11	15:AR:102:ILE:HG12	1.85	0.59
21:AV:104:PHE:C	21:AV:105:VAL:HG12	2.22	0.59
24:AW:60:LEU:H	24:AW:60:LEU:HD12	1.67	0.59
31:BA:1216:G:H2'	31:BA:1217:C:C6	2.37	0.59
31:BA:189:U:N3	47:BT:72:ARG:NH1	2.50	0.59
31:BA:511:C:C2	31:BA:512:U:C4	2.91	0.59
31:BA:518:C:C4'	31:BA:519:C:O5'	2.49	0.59
31:BA:551:U:H2'	31:BA:552:U:C6	2.38	0.59
31:BA:633:G:H2'	31:BA:634:C:C6	2.38	0.59
34:BG:53:ASP:HB3	34:BG:57:ARG:HH12	1.66	0.59
37:BJ:76:ARG:HG2	37:BJ:76:ARG:HH11	1.67	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:BV:19:VAL:CG2	49:BV:44:MET:HB3	2.33	0.59
54:CA:1035:A:H3'	54:CA:1036:G:H5''	1.85	0.59
54:CA:1510:U:H2'	54:CA:1511:G:C8	2.38	0.59
54:CA:560:U:H4'	54:CA:561:U:H5''	1.84	0.59
54:CA:626:U:C2	54:CA:627:G:C8	2.91	0.59
33:CF:70:VAL:HG21	33:CF:76:VAL:HG11	1.84	0.59
54:CA:437:U:H5''	34:CG:155:LEU:HD22	1.85	0.59
36:CI:61:LEU:HB3	36:CI:63:TYR:HE2	1.68	0.59
43:CP:82:MET:HE1	43:CP:93:ARG:HA	1.83	0.59
48:CU:51:LEU:HB2	48:CU:56:THR:HG23	1.85	0.59
50:CW:83:ARG:HA	50:CW:86:ARG:HB3	1.85	0.59
13:D0:42:LYS:HA	13:D0:45:ARG:HH11	1.68	0.59
17:D2:44:LYS:C	17:D2:46:VAL:H	2.06	0.59
17:D2:49:THR:OG1	17:D2:50:PRO:CD	2.51	0.59
12:DP:80:GLU:HA	22:D3:4:LYS:HZ1	1.67	0.59
55:DA:1169:G:C3'	55:DA:1170:G:H5''	2.33	0.59
4:DE:111:ARG:HD2	4:DE:160:TYR:CD1	2.37	0.59
8:DK:139:GLN:C	8:DK:139:GLN:HE21	2.05	0.59
58:DL:125:ARG:NE	58:DL:132:ARG:NH2	2.51	0.59
58:DL:63:ARG:HE	58:DL:63:ARG:HA	1.67	0.59
12:DP:116:GLU:O	12:DP:120:ILE:HG12	2.03	0.59
19:DT:43:VAL:HG13	19:DT:51:VAL:HG21	1.85	0.59
21:DV:152:ALA:O	21:DV:153:SER:C	2.40	0.59
57:DY:96:PHE:O	57:DY:97:ALA:HB3	2.03	0.59
17:A2:84:LYS:O	17:A2:85:LYS:O	2.20	0.58
28:A6:23:THR:O	28:A6:24:GLU:HB2	2.02	0.58
1:AA:1041:C:H2'	1:AA:1042:G:H8	1.68	0.58
1:AA:1341:U:H4'	19:AT:56:THR:O	2.03	0.58
1:AA:2169:A:H2	1:AA:2170:A:N1	2.00	0.58
1:AA:2406:U:N3	11:AO:72:PRO:HB2	2.18	0.58
1:AA:2478:A:O2'	1:AA:2528:U:H1'	2.03	0.58
1:AA:85:G:P	20:AU:30:VAL:HG21	2.42	0.58
2:AB:40:U:O4	2:AB:43:C:OP1	2.21	0.58
3:AD:145:VAL:HG13	3:AD:191:ALA:HB2	1.84	0.58
4:AE:130:GLY:O	4:AE:131:ALA:HB3	2.03	0.58
4:AE:24:THR:HG21	4:AE:186:GLY:O	2.02	0.58
6:AG:106:LEU:O	6:AG:110:ALA:HB3	2.03	0.58
1:AA:2316:C:H1'	6:AG:128:ARG:HH22	1.68	0.58
9:AM:14:VAL:CG1	9:AM:137:LYS:HG3	2.33	0.58
12:AP:135:ASP:CG	21:AV:81:ARG:HH12	2.06	0.58
15:AR:26:ASP:O	15:AR:49:VAL:HG12	2.03	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:AV:125:LEU:C	21:AV:164:ALA:HB3	2.22	0.58
1:AA:896:A:H2	21:AV:178:GLU:HG2	1.67	0.58
21:AV:185:GLU:O	21:AV:186:GLU:CB	2.37	0.58
31:BA:1106:G:H2'	31:BA:1107:C:H6	1.68	0.58
31:BA:1298:C:H5''	37:BJ:114:ARG:HH22	1.68	0.58
31:BA:1328:C:C2'	31:BA:1329:A:H5'	2.33	0.58
31:BA:1498:U:H1'	31:BA:1499:A:N7	2.17	0.58
31:BA:50:A:N6	31:BA:361:G:H4'	2.19	0.58
52:BB:19:G:HO2'	52:BB:20:U:P	2.26	0.58
34:BG:22:LYS:HG3	34:BG:26:CYS:HB2	1.84	0.58
31:BA:1248:A:H2'	39:BL:70:LYS:NZ	2.16	0.58
42:BO:70:ILE:CD1	42:BO:100:ILE:HD12	2.33	0.58
43:BP:50:GLU:O	43:BP:53:VAL:HB	2.02	0.58
43:BP:5:ALA:HB2	43:BP:22:ILE:HD13	1.85	0.58
54:CA:1004:A:H2'	54:CA:1005:A:O5'	2.03	0.58
54:CA:1378:C:O2	37:CJ:156:TRP:HH2	1.86	0.58
54:CA:630:G:H8	54:CA:630:G:H5''	1.67	0.58
52:CB:37:MIA:H122	52:CB:38:A:N1	2.16	0.58
38:CK:80:ILE:HG23	38:CK:137:VAL:CG1	2.33	0.58
54:CA:973:G:C1'	40:CM:55:LYS:HE2	2.27	0.58
43:CP:97:PRO:HB2	43:CP:101:GLN:NE2	2.18	0.58
45:CR:56:LEU:HA	45:CR:59:MET:HE3	1.85	0.58
49:CV:81:ARG:O	49:CV:82:GLY:O	2.20	0.58
49:CV:83:HIS:CG	49:CV:84:GLY:H	2.18	0.58
55:DA:1454:U:OP1	13:D0:77:ARG:NH1	2.35	0.58
28:D6:30:THR:HG23	28:D6:30:THR:O	2.03	0.58
55:DA:1049:C:N3	55:DA:2751:G:C6	2.71	0.58
55:DA:1092:C:H2'	55:DA:1093:G:C5'	2.32	0.58
55:DA:1430:C:H2'	55:DA:1431:U:C6	2.38	0.58
55:DA:1735:C:C6	55:DA:1735:C:H5'	2.37	0.58
55:DA:2087:G:O2'	55:DA:2088:G:H5'	2.03	0.58
55:DA:271(B):G:H4'	55:DA:271(C):U:O5'	2.03	0.58
55:DA:2804:C:H2'	55:DA:2805:G:C8	2.38	0.58
55:DA:897:C:O5'	55:DA:897:C:H6	1.86	0.58
56:DI:15:ALA:O	56:DI:19:GLU:HG2	2.03	0.58
8:DK:142:VAL:HG23	8:DK:142:VAL:O	2.03	0.58
55:DA:1058:U:O3'	58:DL:4:VAL:HG11	2.03	0.58
58:DL:69:THR:CG2	58:DL:70:LYS:H	2.07	0.58
12:DP:120:ILE:O	12:DP:123:HIS:HB2	2.03	0.58
18:DS:80:PRO:O	18:DS:100:THR:CG2	2.51	0.58
55:DA:1082:U:OP2	57:DY:45:LYS:CG	2.51	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1020:A:H4'	1:AA:1021:A:O5'	2.03	0.58
1:AA:1024:G:H3'	1:AA:1025:G:C5'	2.26	0.58
1:AA:119:A:O2'	1:AA:120:U:OP2	2.20	0.58
1:AA:1945:G:H2'	1:AA:1946:U:H6	1.67	0.58
1:AA:2126:A:O2'	1:AA:2127:G:O5'	2.21	0.58
1:AA:2307:G:H1	6:AG:44:GLY:H	1.51	0.58
1:AA:2481:G:HO2'	1:AA:2482:G:P	2.26	0.58
1:AA:871:U:O2	1:AA:871:U:C2'	2.51	0.58
3:AD:267:SER:C	3:AD:269:PHE:N	2.57	0.58
6:AG:161:THR:HG22	6:AG:162:THR:N	2.18	0.58
6:AG:38:VAL:HG22	6:AG:93:THR:HG23	1.85	0.58
1:AA:385:C:O2	11:AO:71:VAL:HG21	2.03	0.58
14:AQ:58:LEU:N	14:AQ:58:LEU:HD23	2.17	0.58
15:AR:41:ARG:C	15:AR:42:ILE:HD12	2.23	0.58
20:AU:91:GLU:CG	20:AU:92:ASN:H	2.06	0.58
20:AU:98:VAL:HG13	20:AU:99:CYS:SG	2.42	0.58
21:AV:104:PHE:HA	21:AV:139:VAL:O	2.03	0.58
12:AP:134:ARG:HH22	21:AV:122:ARG:NH1	2.01	0.58
24:AW:24:LEU:HD22	24:AW:60:LEU:HD21	1.85	0.58
53:B1:53:U:C2'	53:B1:54:U:OP1	2.51	0.58
31:BA:973:G:H1'	40:BM:55:LYS:CE	2.33	0.58
52:BC:58:A:C1'	52:BC:60:U:H5	2.15	0.58
32:BE:42:ILE:HD13	32:BE:203:GLY:HA2	1.85	0.58
33:BF:36:ASP:OD1	33:BF:57:ILE:HG21	2.03	0.58
36:BI:100:ASN:O	48:BU:28:GLU:HB3	2.02	0.58
54:CA:1102:A:H2'	54:CA:1103:C:C6	2.38	0.58
54:CA:38:G:H4'	54:CA:547:A:N6	2.18	0.58
54:CA:957:U:H1'	54:CA:960:U:H5	1.66	0.58
32:CE:162:ILE:HG13	32:CE:162:ILE:O	2.02	0.58
32:CE:77:ALA:O	32:CE:81:VAL:HG23	2.03	0.58
42:CO:89:ARG:HE	42:CO:91:LYS:HZ3	1.50	0.58
42:CO:89:ARG:HH21	42:CO:91:LYS:HZ3	1.49	0.58
26:D4:58:ARG:CB	26:D4:62:ARG:HB3	2.32	0.58
27:D5:40:LYS:CG	27:D5:46:CYS:HB3	2.33	0.58
28:D6:17:LYS:C	28:D6:19:ARG:N	2.55	0.58
30:D8:29:LYS:HZ3	30:D8:44:LYS:HB2	1.69	0.58
55:DA:84:A:N6	55:DA:102:G:O2'	2.36	0.58
55:DA:1061:U:C2'	55:DA:1062:G:O5'	2.51	0.58
55:DA:1048:A:OP2	55:DA:1110:G:N2	2.35	0.58
55:DA:1926:U:H2'	55:DA:1928:A:OP2	2.03	0.58
55:DA:528:A:C2	55:DA:2043:C:C5'	2.85	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2113:U:H5'	55:DA:2114:A:H8	1.66	0.58
55:DA:2682:U:H6	55:DA:2682:U:H5'	1.68	0.58
55:DA:1050:A:O2'	55:DA:2752:C:H1'	2.03	0.58
55:DA:484:C:H2'	55:DA:485:C:C6	2.38	0.58
55:DA:654(A):A:N1	55:DA:654(T):A:N1	2.51	0.58
55:DA:969:U:H2'	55:DA:970:C:C6	2.38	0.58
2:DB:27:C:H5'	2:DB:28:C:OP2	2.02	0.58
55:DA:442:G:H4'	5:DF:46:ARG:HD3	1.85	0.58
58:DL:52:ILE:O	58:DL:53:VAL:O	2.20	0.58
58:DL:10:LEU:HD11	58:DL:55:VAL:HG21	1.85	0.58
58:DL:52:ILE:HG12	58:DL:76:TYR:CB	2.33	0.58
9:DM:134:ARG:N	9:DM:135:PRO:CD	2.63	0.58
14:DQ:83:LYS:HG2	14:DQ:109:GLY:N	2.17	0.58
15:DR:94:ALA:O	15:DR:95:ARG:HB2	2.02	0.58
18:DS:20:VAL:HG23	18:DS:47:VAL:HG21	1.85	0.58
20:DU:54:LYS:O	20:DU:55:TYR:HB2	2.02	0.58
24:DW:40:SER:C	24:DW:42:GLY:N	2.56	0.58
57:DY:50:ARG:HD3	57:DY:51:LEU:H	1.65	0.58
16:A1:95:LEU:HD12	17:A2:11:GLN:HB3	1.84	0.58
1:AA:1267:U:C5	1:AA:2012:G:N1	2.72	0.58
1:AA:2614:A:H4'	1:AA:2615:U:OP1	2.03	0.58
1:AA:2780:G:C2'	1:AA:2781:A:OP1	2.51	0.58
1:AA:27:G:O2'	1:AA:28:A:OP2	2.21	0.58
1:AA:654:A:N3	1:AA:654:A:H2'	2.17	0.58
1:AA:654(S):G:O3'	1:AA:654(T):A:H8	1.85	0.58
5:AF:28:ILE:HA	5:AF:112:MET:HE3	1.86	0.58
5:AF:60:SER:O	5:AF:61:GLY:C	2.41	0.58
6:AG:161:THR:HG22	6:AG:163:ALA:N	2.14	0.58
6:AG:34:LEU:HD13	6:AG:34:LEU:O	2.03	0.58
8:AK:120:ILE:HG22	8:AK:122:GLU:H	1.67	0.58
8:AK:52:ARG:C	8:AK:52:ARG:HD2	2.23	0.58
9:AM:111:PRO:HA	9:AM:114:ARG:CZ	2.33	0.58
9:AM:74:ARG:HH12	9:AM:85:ILE:HD11	1.67	0.58
11:AO:86:LYS:HG3	11:AO:87:ASP:N	2.17	0.58
23:AZ:7:ILE:HG12	23:AZ:91:LYS:HZ3	1.68	0.58
31:BA:321:A:N7	31:BA:328:C:O2	2.36	0.58
31:BA:872:A:C4'	31:BA:873:A:OP1	2.47	0.58
37:BJ:16:LEU:O	37:BJ:17:VAL:HG23	2.02	0.58
40:BM:33:GLN:N	40:BM:75:ILE:HG12	2.17	0.58
40:BM:8:LEU:CG	40:BM:96:ILE:HG22	2.21	0.58
48:BU:22:VAL:O	48:BU:23:LYS:CG	2.49	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:BV:42:PRO:C	49:BV:45:VAL:H	2.03	0.58
50:BW:87:LYS:O	50:BW:90:GLN:N	2.36	0.58
53:C1:30:C:H6	53:C1:30:C:H3'	1.67	0.58
54:CA:1117:G:H4'	39:CL:104:ARG:HH21	1.69	0.58
54:CA:243:A:H4'	54:CA:244:U:O5'	2.01	0.58
54:CA:518:C:C4	54:CA:530:G:N7	2.71	0.58
54:CA:869:G:H4'	54:CA:872:A:H1'	1.84	0.58
54:CA:963:G:H21	40:CM:55:LYS:CD	2.13	0.58
32:CE:4:GLU:C	32:CE:5:ILE:HD13	2.24	0.58
34:CG:19:LEU:HG	34:CG:21:LEU:HG	1.85	0.58
35:CH:55:VAL:O	35:CH:58:ALA:HB3	2.04	0.58
36:CI:60:PHE:C	36:CI:61:LEU:HD12	2.23	0.58
37:CJ:15:ASP:HB3	37:CJ:19:GLY:N	2.18	0.58
40:CM:62:HIS:CD2	40:CM:62:HIS:H	2.21	0.58
42:CO:26:ALA:O	42:CO:27:LEU:O	2.21	0.58
16:D1:108:GLU:HB2	17:D2:44:LYS:HE3	1.85	0.58
26:D4:56:VAL:HG13	26:D4:60:GLN:HG3	1.84	0.58
55:DA:1173:G:H4'	55:DA:1174:A:C2	2.39	0.58
55:DA:1771:C:C1'	55:DA:1786:A:H8	2.15	0.58
55:DA:270(K):C:O2'	55:DA:270(L):U:H5''	2.02	0.58
55:DA:858:U:O2'	55:DA:2268:A:H1'	2.02	0.58
5:DF:184:TYR:CE2	5:DF:188:ARG:HD2	2.38	0.58
7:DH:92:ILE:HD12	7:DH:92:ILE:N	2.13	0.58
56:DJ:3:LEU:O	56:DJ:6:GLU:N	2.36	0.58
8:DK:3:VAL:O	8:DK:18:VAL:HA	2.03	0.58
58:DL:95:LYS:CG	58:DL:136:VAL:HG21	2.32	0.58
58:DL:99:ILE:O	58:DL:138:VAL:HG13	2.03	0.58
58:DL:53:VAL:CG1	58:DL:76:TYR:CD2	2.86	0.58
15:DR:102:ILE:HA	15:DR:105:LEU:CD2	2.34	0.58
21:DV:122:ARG:HH11	21:DV:122:ARG:HG2	1.67	0.58
21:DV:155:LEU:O	21:DV:157:LEU:HD13	2.03	0.58
57:DY:126:ALA:O	57:DY:127:GLU:C	2.42	0.58
57:DY:50:ARG:H	57:DY:83:TYR:CA	2.16	0.58
26:A4:64:GLY:O	26:A4:70:GLY:HA2	2.02	0.58
30:A8:62:LEU:CB	30:A8:63:PRO:CD	2.81	0.58
1:AA:1332:G:N2	1:AA:1609:A:C2'	2.65	0.58
1:AA:2712:U:H1'	1:AA:2712(A):A:C8	2.38	0.58
1:AA:536:A:H2'	1:AA:537:C:C6	2.38	0.58
1:AA:558:G:P	9:AM:111:PRO:HD2	2.42	0.58
1:AA:654(C):G:C2	1:AA:654(S):G:C2	2.91	0.58
1:AA:669:G:H4'	1:AA:670:A:OP1	2.01	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:885:C:H2'	1:AA:886:C:O4'	2.04	0.58
20:AU:74:PRO:O	20:AU:80:GLY:HA2	2.03	0.58
21:AV:116:VAL:CG1	21:AV:117:LEU:H	2.13	0.58
32:BE:54:THR:HG22	32:BE:58:ILE:HD11	1.85	0.58
33:BF:83:ARG:NE	33:BF:87:LEU:HD11	2.18	0.58
37:BJ:48:LYS:O	37:BJ:52:GLU:HG2	2.02	0.58
42:BO:58:VAL:O	42:BO:65:GLU:HA	2.02	0.58
43:BP:49:THR:HG22	43:BP:51:ALA:H	1.69	0.58
54:CA:565:U:H5''	54:CA:566:G:H3'	1.85	0.58
54:CA:930:C:O2'	54:CA:931:C:H5'	2.03	0.58
54:CA:983:A:HO2'	54:CA:1049:U:HO2'	1.49	0.58
52:CD:22:G:O2'	52:CD:23:A:H5'	2.03	0.58
32:CE:229:VAL:HG12	32:CE:229:VAL:O	2.03	0.58
32:CE:71:VAL:HG12	32:CE:93:VAL:HB	1.83	0.58
36:CI:7:ASN:O	36:CI:88:VAL:HA	2.04	0.58
54:CA:600:C:OP1	38:CK:97:VAL:HG12	2.04	0.58
46:CS:28:ARG:NH1	46:CS:29:ASP:OD2	2.36	0.58
54:CA:376:G:OP1	46:CS:5:ARG:HB2	2.02	0.58
13:D0:2:ARG:HA	13:D0:5:LYS:HD2	1.84	0.58
30:D8:35:GLN:HA	30:D8:35:GLN:HE21	1.68	0.58
55:DA:1926:U:C1'	55:DA:1929:G:O6	2.51	0.58
55:DA:2126:A:O2'	55:DA:2127:G:O5'	2.22	0.58
55:DA:2308:G:H1	55:DA:2311:A:H2	1.37	0.58
7:DH:106:THR:HG22	7:DH:112:PRO:CB	2.30	0.58
8:DK:57:ARG:HH11	8:DK:57:ARG:HB2	1.68	0.58
58:DL:111:LYS:C	58:DL:113:PRO:CD	2.71	0.58
11:DO:143:GLY:C	11:DO:144:GLU:HG3	2.24	0.58
11:DO:61:ARG:HB2	11:DO:61:ARG:NH2	2.17	0.58
15:DR:88:ILE:HG13	15:DR:88:ILE:O	2.03	0.58
19:DT:70:LEU:HD23	19:DT:70:LEU:N	2.18	0.58
21:DV:196:VAL:C	21:DV:197:ILE:HD12	2.23	0.58
17:A2:82:ARG:NH1	17:A2:82:ARG:HG3	2.17	0.58
26:A4:2:LYS:HB3	26:A4:6:HIS:CE1	2.37	0.58
1:AA:1008:C:H5''	1:AA:1009:A:OP1	2.04	0.58
1:AA:1188:U:C2'	1:AA:1189:A:H5'	2.33	0.58
1:AA:2672:G:C3'	1:AA:2673:G:H5''	2.34	0.58
1:AA:270(B):A:H5''	1:AA:270(C):C:OP2	2.04	0.58
1:AA:690:G:H2'	1:AA:691:C:C6	2.38	0.58
1:AA:859:G:O2'	1:AA:860:U:P	2.61	0.58
1:AA:896:A:O5'	1:AA:897:C:C5	2.56	0.58
8:AK:10:GLU:CD	8:AK:11:ASN:N	2.56	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:AN:2:ILE:HD11	10:AN:82:ASN:ND2	2.19	0.58
11:AO:19:VAL:CG2	11:AO:20:GLY:H	1.95	0.58
2:AB:90:C:OP1	12:AP:16:ARG:HD2	2.02	0.58
53:B1:52:U:H4'	53:B1:52:U:OP1	2.03	0.58
31:BA:1139:G:H5'	31:BA:1140:C:OP1	2.04	0.58
31:BA:1277:C:O2'	31:BA:1279:A:H8	1.86	0.58
31:BA:1261:A:O4'	31:BA:1283:G:H5''	2.04	0.58
31:BA:192:U:H2'	31:BA:193:C:H6	1.68	0.58
31:BA:35:G:H2'	31:BA:36:C:C6	2.38	0.58
31:BA:409:G:H2'	31:BA:410:G:C8	2.37	0.58
31:BA:528:C:H5'	31:BA:535:A:N1	2.19	0.58
33:BF:34:LEU:HG	33:BF:38:ARG:HH21	1.68	0.58
37:BJ:23:VAL:HG13	37:BJ:43:PHE:HE2	1.67	0.58
41:BN:41:THR:OG1	41:BN:71:LYS:HD3	2.04	0.58
43:BP:70:LEU:HD22	43:BP:70:LEU:C	2.24	0.58
43:BP:23:TYR:CE1	43:BP:71:ARG:HD3	2.37	0.58
50:BW:18:GLN:HE21	50:BW:22:ARG:HH12	1.50	0.58
54:CA:1156:G:H5''	54:CA:1157:A:OP2	2.02	0.58
54:CA:946:A:H2'	54:CA:947:G:H8	1.66	0.58
37:CJ:94:ARG:HG3	37:CJ:94:ARG:NH1	2.14	0.58
54:CA:1313:U:OP1	49:CV:6:LYS:HB3	2.03	0.58
49:CV:8:GLY:O	49:CV:9:VAL:O	2.21	0.58
13:D0:1:MET:O	13:D0:2:ARG:HB2	2.02	0.58
13:D0:67:LEU:HD13	13:D0:76:VAL:HG21	1.85	0.58
16:D1:50:ARG:HG2	16:D1:53:ARG:NH2	2.18	0.58
17:D2:43:GLU:HA	17:D2:43:GLU:OE2	2.03	0.58
55:DA:1543:A:C8	55:DA:1545:A:H5''	2.38	0.58
55:DA:1847:A:H2'	55:DA:1847:A:N3	2.19	0.58
55:DA:2115:G:N2	55:DA:2172:U:H3	2.01	0.58
55:DA:2590:A:O2'	55:DA:2591:C:H5'	2.04	0.58
55:DA:654(J):A:N1	55:DA:654(L):G:O6	2.36	0.58
55:DA:880:G:C4'	55:DA:880:G:OP1	2.51	0.58
58:DL:67:PHE:C	58:DL:68:VAL:HG12	2.24	0.58
9:DM:113:GLY:C	9:DM:114:ARG:O	2.40	0.58
9:DM:18:ALA:HB3	9:DM:55:VAL:O	2.03	0.58
11:DO:38:GLN:HG2	11:DO:45:LEU:HD12	1.86	0.58
15:DR:125:ARG:HG2	15:DR:126:ALA:N	2.18	0.58
57:DY:58:LEU:C	57:DY:62:ALA:CB	2.72	0.58
57:DY:88:ALA:C	57:DY:92:THR:HB	2.24	0.58
17:A2:73:SER:HB2	17:A2:82:ARG:O	2.03	0.58
27:A5:3:LYS:HA	27:A5:3:LYS:CE	2.32	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:A7:8:ASN:ND2	29:A7:10:ARG:H	2.02	0.58
1:AA:1092:C:H2'	1:AA:1093:G:H5'	1.85	0.58
1:AA:1267:U:C4	1:AA:2012:G:N2	2.71	0.58
1:AA:1340:U:C2'	1:AA:1341:U:OP1	2.51	0.58
1:AA:2193:G:H8	1:AA:2193:G:H5'	1.69	0.58
1:AA:244:A:H2'	1:AA:245:G:O4'	2.03	0.58
1:AA:2839:G:H5'	13:A0:46:GLY:HA2	1.86	0.58
2:AB:40:U:C2	26:A4:1:MET:SD	2.96	0.58
4:AE:11:MET:O	4:AE:12:THR:HG23	2.03	0.58
5:AF:155:LEU:HD22	5:AF:185:ASP:O	2.02	0.58
6:AG:43:LEU:HD22	6:AG:90:LEU:HD23	1.86	0.58
10:AN:10:VAL:HG22	10:AN:17:ARG:O	2.03	0.58
14:AQ:83:LYS:HG2	14:AQ:109:GLY:H	1.69	0.58
14:AQ:3:ARG:HG2	14:AQ:4:LEU:N	2.17	0.58
18:AS:22:ASP:HA	18:AS:25:ARG:HH12	1.67	0.58
53:B1:57:U:H2'	53:B1:57:U:O2	2.01	0.58
31:BA:1326:C:OP1	51:BX:12:LYS:HE2	2.03	0.58
31:BA:1423:G:H2'	31:BA:1424:C:H6	1.67	0.58
31:BA:1513:A:H2'	31:BA:1514:C:H6	1.69	0.58
31:BA:279:A:O2'	31:BA:280:C:P	2.62	0.58
31:BA:423:G:N2	31:BA:424:G:C8	2.72	0.58
31:BA:434:U:H2'	31:BA:435:C:C6	2.39	0.58
31:BA:609:A:H2'	31:BA:610:G:O4'	2.04	0.58
33:BF:119:ARG:HH21	33:BF:137:ALA:HA	1.67	0.58
38:BK:84:ARG:HH12	38:BK:86:ILE:CD1	2.16	0.58
40:BM:79:ARG:O	40:BM:83:GLU:HB2	2.03	0.58
42:BO:86:ARG:HG2	42:BO:86:ARG:O	2.04	0.58
53:C1:30:C:N4	53:C1:31:A:N7	2.51	0.58
54:CA:1221:G:H4'	49:CV:77:THR:CG2	2.33	0.58
54:CA:1347:G:O2'	54:CA:1348:U:P	2.62	0.58
54:CA:444:C:H2'	54:CA:445:G:H8	1.69	0.58
54:CA:474:G:OP1	46:CS:81:ARG:HG3	2.03	0.58
54:CA:531:U:H4'	54:CA:532:A:OP1	2.04	0.58
54:CA:923:A:H2'	54:CA:924:C:C6	2.38	0.58
54:CA:91:C:H2'	54:CA:92:G:C5'	2.32	0.58
54:CA:991:U:O2	54:CA:993:G:H8	1.85	0.58
34:CG:106:TYR:HE1	34:CG:112:VAL:O	1.87	0.58
39:CL:96:LEU:HD23	39:CL:102:LEU:HD12	1.86	0.58
40:CM:4:ILE:HB	40:CM:74:ILE:CG1	2.32	0.58
42:CO:126:LYS:HE2	42:CO:128:ALA:HB3	1.85	0.58
42:CO:6:THR:OG1	42:CO:9:GLN:HG3	2.04	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
46:CS:20:VAL:HG22	46:CS:21:VAL:N	2.18	0.58
47:CT:45:HIS:HE2	47:CT:47:PRO:HG3	1.68	0.58
49:CV:41:VAL:N	49:CV:44:MET:SD	2.73	0.58
49:CV:42:PRO:O	49:CV:45:VAL:HG13	2.04	0.58
27:D5:42:PRO:O	27:D5:44:THR:HG23	2.04	0.58
55:DA:1082:U:OP1	55:DA:1082:U:C4'	2.52	0.58
55:DA:1071:G:H1'	55:DA:1089:G:H3'	1.86	0.58
55:DA:1170:G:H8	55:DA:1170:G:H5'	1.69	0.58
55:DA:1238:G:O2'	55:DA:1239:G:H5'	2.03	0.58
55:DA:1820:U:H2'	3:DD:159:ALA:O	2.03	0.58
55:DA:414:C:O2	55:DA:1864:U:O2'	2.21	0.58
55:DA:1934:C:C5'	55:DA:1934:C:H6	2.08	0.58
55:DA:2315:G:H2'	55:DA:2316:C:C6	2.38	0.58
55:DA:616:A:O2'	55:DA:617:G:O4'	2.20	0.58
55:DA:654(C):G:C3'	55:DA:654(D):G:H8	2.16	0.58
4:DE:116:VAL:HG21	4:DE:122:PHE:CD2	2.39	0.58
5:DF:195:ASP:O	5:DF:197:ASP:O	2.22	0.58
55:DA:1058:U:C5'	58:DL:4:VAL:HB	2.33	0.58
58:DL:77:LEU:HD21	58:DL:111:LYS:HZ1	1.68	0.58
11:DO:79:ARG:HD3	11:DO:110:TYR:CE1	2.38	0.58
14:DQ:109:GLY:O	14:DQ:110:LEU:HB2	2.03	0.58
20:DU:50:ARG:HB3	20:DU:53:PRO:CD	2.32	0.58
57:DY:73:GLY:C	57:DY:119:ALA:C	2.60	0.58
57:DY:74:LEU:HG	57:DY:120:LYS:HD3	1.85	0.58
57:DY:121:ASP:OD1	57:DY:122:VAL:N	2.36	0.58
57:DY:26:LEU:HD23	57:DY:112:LEU:HB3	1.86	0.58
16:A1:90:VAL:HA	17:A2:39:LEU:CD2	2.34	0.58
1:AA:1240:U:O2'	1:AA:1241:A:H5'	2.03	0.58
1:AA:769:G:H5'	1:AA:1379:A:N6	2.18	0.58
1:AA:13:A:O2'	1:AA:15:G:N7	2.33	0.58
1:AA:1954:G:O2'	1:AA:1955:U:P	2.60	0.58
1:AA:2148:G:O2'	1:AA:2149:G:H5'	2.03	0.58
1:AA:2174:C:O2'	1:AA:2175:C:H5'	2.03	0.58
1:AA:242:G:C2'	1:AA:243:U:OP2	2.51	0.58
1:AA:307:G:N2	1:AA:309:G:H3'	2.18	0.58
1:AA:671:C:O2'	1:AA:672:C:H5'	2.04	0.58
2:AB:46:A:H2'	2:AB:47:C:H6	1.65	0.58
4:AE:3:GLY:C	4:AE:4:ILE:HG23	2.23	0.58
5:AF:152:GLU:HA	5:AF:190:GLU:OE2	2.03	0.58
8:AK:76:THR:HG21	8:AK:138:ILE:HG12	1.85	0.58
9:AM:36:GLY:O	9:AM:42:TRP:HB2	2.03	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AO:85:LEU:CA	11:AO:88:LEU:HB3	2.32	0.58
20:AU:95:LYS:HA	20:AU:101:LYS:HB2	1.85	0.58
21:AV:69:THR:HA	21:AV:91:LEU:HD23	1.86	0.58
21:AV:77:ASP:OD2	21:AV:80:ARG:HB2	2.03	0.58
31:BA:1090:U:O2'	31:BA:1091:U:H5'	2.03	0.58
31:BA:530:G:N2	31:BA:1492:A:N1	2.51	0.58
31:BA:38:G:H22	31:BA:397:A:H5''	1.69	0.58
31:BA:710:G:OP1	36:BI:54:LYS:HE3	2.03	0.58
31:BA:812:C:O2'	31:BA:813:U:P	2.61	0.58
32:BE:212:GLN:NE2	32:BE:216:SER:HB2	2.17	0.58
35:BH:41:VAL:HG23	35:BH:69:VAL:HG21	1.86	0.58
42:BO:60:LEU:HB2	42:BO:64:TYR:CB	2.32	0.58
44:BQ:48:ALA:N	44:BQ:53:LEU:HD12	2.18	0.58
46:BS:74:LEU:O	46:BS:79:VAL:HG23	2.03	0.58
47:BT:86:GLU:O	47:BT:90:ILE:HG12	2.03	0.58
54:CA:1002:G:C2'	54:CA:1003:G:H5'	2.34	0.58
54:CA:1300:G:O2'	54:CA:1301:U:O5'	2.19	0.58
54:CA:197:A:N6	54:CA:221:C:H5''	2.18	0.58
54:CA:498:A:O2'	54:CA:500:G:O5'	2.22	0.58
52:CB:6:G:H2'	52:CB:7:A:O4'	2.04	0.58
52:CD:20:U:H2'	52:CD:20:U:O2	2.03	0.58
42:CO:119:LYS:HB2	42:CO:120:TYR:CD1	2.39	0.58
43:CP:14:ARG:N	43:CP:44:ARG:HD2	2.19	0.58
13:D0:85:PRO:O	13:D0:87:TYR:N	2.37	0.58
55:DA:18:C:H4'	16:D1:23:GLY:O	2.04	0.58
22:D3:33:ALA:HB2	22:D3:63:VAL:HA	1.85	0.58
22:D3:53:MET:HA	22:D3:58:THR:O	2.04	0.58
55:DA:2285:C:H41	28:D6:27:LYS:HE2	1.68	0.58
55:DA:1061:U:H2'	55:DA:1062:G:O5'	2.04	0.58
55:DA:1166:C:O2'	55:DA:1167:U:H5'	2.04	0.58
55:DA:1510:A:OP1	55:DA:1510:A:H4'	2.03	0.58
55:DA:1694:C:H1'	55:DA:1695:G:N2	2.18	0.58
55:DA:1930:G:O2'	55:DA:1931:U:OP2	2.19	0.58
55:DA:2712:U:O2'	55:DA:2712(A):A:O5'	2.21	0.58
55:DA:373:U:O2	55:DA:423:A:H2	1.85	0.58
55:DA:492:A:H2'	55:DA:493:G:H5'	1.85	0.58
56:DJ:12:LEU:N	56:DJ:13:SER:CA	2.67	0.58
55:DA:969:U:OP1	25:DX:17:LYS:HG3	2.03	0.58
16:A1:91:ASP:O	16:A1:92:ARG:HB3	2.04	0.58
1:AA:1060:U:H5''	1:AA:1061:U:OP1	2.03	0.58
1:AA:1250:G:O2'	1:AA:1251:C:OP1	2.20	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1272:A:H3'	1:AA:1273:U:C5'	2.34	0.58
1:AA:1364:G:OP2	23:AZ:2:SER:HA	2.04	0.58
1:AA:1427:A:H4'	1:AA:1428:C:O4'	2.04	0.58
1:AA:1705:G:H2'	1:AA:1706:U:O4'	2.04	0.58
1:AA:1731:G:C2'	1:AA:1732:A:H5'	2.32	0.58
1:AA:205:G:HO2'	1:AA:206:U:P	2.26	0.58
1:AA:2060:A:HO2'	1:AA:2061:G:P	2.26	0.58
1:AA:2763:G:H5'	1:AA:2764:A:OP2	2.04	0.58
1:AA:2720:U:N3	1:AA:2873:A:C2	2.66	0.58
1:AA:908:C:O2'	1:AA:909:A:H5'	2.04	0.58
8:AK:117:GLU:CD	8:AK:117:GLU:N	2.50	0.58
8:AK:114:LEU:HD23	8:AK:130:TYR:HB2	1.86	0.58
8:AK:68:LEU:O	8:AK:71:ILE:HG22	2.03	0.58
10:AN:10:VAL:HG23	10:AN:10:VAL:O	2.03	0.58
11:AO:124:LYS:HZ2	11:AO:143:GLY:HA3	1.67	0.58
12:AP:109:VAL:HG12	12:AP:110:THR:N	2.18	0.58
12:AP:140:ALA:O	12:AP:141:GLN:HB2	2.02	0.58
12:AP:70:PRO:HA	12:AP:94:VAL:O	2.04	0.58
25:AX:23:LEU:HD11	25:AX:53:LEU:CD1	2.33	0.58
31:BA:1328:C:O2'	31:BA:1329:A:H5'	2.03	0.58
31:BA:1532:U:H2'	31:BA:1533:C:C6	2.39	0.58
31:BA:335:C:H2'	31:BA:336:C:C6	2.39	0.58
22:A3:5:LYS:CE	52:BC:73:A:O2'	2.42	0.58
32:BE:206:ASP:O	32:BE:211:ILE:HD11	2.04	0.58
32:BE:22:LYS:H	32:BE:22:LYS:NZ	2.01	0.58
34:BG:149:ALA:O	34:BG:150:GLU:O	2.21	0.58
39:BL:82:ALA:HB1	39:BL:96:LEU:HD21	1.85	0.58
40:BM:6:ILE:O	40:BM:6:ILE:HG13	2.03	0.58
43:BP:56:LEU:O	43:BP:60:VAL:HG23	2.03	0.58
43:BP:85:GLY:O	43:BP:86:CYS:C	2.42	0.58
44:BQ:26:ARG:HD3	44:BQ:43:CYS:CB	2.34	0.58
45:BR:54:ARG:HG2	45:BR:58:MET:HE2	1.85	0.58
53:C1:31:A:O2'	53:C1:32:A:O5'	2.21	0.58
54:CA:1006:C:H2'	54:CA:1007:C:C6	2.39	0.58
54:CA:1498:U:O2'	54:CA:1499:A:OP2	2.20	0.58
54:CA:32:A:H2'	54:CA:33:A:C8	2.39	0.58
54:CA:564:C:H5'	47:CT:32:TYR:HE2	1.67	0.58
54:CA:22:G:H4'	54:CA:885:G:C8	2.38	0.58
52:CB:46:G:H5''	52:CB:47:U:OP2	2.04	0.58
32:CE:4:GLU:HG2	32:CE:5:ILE:N	2.18	0.58
43:CP:50:GLU:O	43:CP:54:VAL:HG23	2.02	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
45:CR:82:ILE:CG2	45:CR:83:GLU:N	2.66	0.58
48:CU:45:SER:HG	48:CU:47:THR:HG1	1.52	0.58
16:D1:11:ARG:O	16:D1:15:LYS:HG3	2.04	0.58
16:D1:52:ARG:HG2	16:D1:52:ARG:HH11	1.69	0.58
28:D6:9:LEU:HD13	28:D6:11:LEU:CD2	2.33	0.58
55:DA:1945:G:H2'	55:DA:1946:U:H6	1.69	0.58
55:DA:2443:C:O2'	55:DA:2444:G:H5'	2.03	0.58
55:DA:2702:U:OP1	55:DA:2702:U:O4'	2.22	0.58
55:DA:2771:C:H2'	55:DA:2772:C:C6	2.38	0.58
55:DA:328:U:O2'	20:DU:71:LYS:HD3	2.04	0.58
2:DB:113:C:O2'	14:DQ:46:VAL:HG13	2.04	0.58
3:DD:134:ARG:HB2	3:DD:135:PHE:HD2	1.68	0.58
3:DD:35:LYS:HG2	3:DD:64:ILE:CA	2.34	0.58
4:DE:174:ASP:HB3	4:DE:183:LEU:HD22	1.85	0.58
5:DF:31:HIS:HB2	11:DO:9:ASN:ND2	2.16	0.58
55:DA:1079:C:H1'	58:DL:129:GLY:HA3	1.84	0.58
58:DL:86:LYS:C	58:DL:88:ALA:H	2.05	0.58
58:DL:99:ILE:HG13	58:DL:138:VAL:CG2	2.25	0.58
19:DT:15:GLU:N	19:DT:15:GLU:OE1	2.26	0.58
21:DV:189:ALA:HB2	21:DV:190:GLU:HG2	1.71	0.58
57:DY:12:THR:HB	57:DY:52:PHE:CD2	2.38	0.58
57:DY:75:GLN:HG3	57:DY:110:GLY:H	1.69	0.58
1:AA:1060:U:O2	1:AA:1088:A:H8	1.85	0.58
1:AA:1106:G:H2'	1:AA:1107:G:C8	2.39	0.58
1:AA:1932:A:H2'	1:AA:1933:G:O4'	2.03	0.58
1:AA:2091:U:H3'	1:AA:2092:U:H5''	1.84	0.58
1:AA:2126:A:HO2'	1:AA:2127:G:C5'	2.16	0.58
1:AA:2656:U:C6	1:AA:2656:U:H3'	2.39	0.58
1:AA:897:C:OP2	1:AA:897:C:H6	1.87	0.58
3:AD:70:TRP:CH2	3:AD:150:LYS:HA	2.39	0.58
4:AE:203:LYS:HD3	4:AE:203:LYS:C	2.24	0.58
5:AF:7:TYR:CD2	5:AF:18:ARG:HB2	2.39	0.58
6:AG:53:LEU:C	6:AG:53:LEU:HD23	2.24	0.58
7:AH:143:GLN:C	7:AH:143:GLN:HE21	2.07	0.58
8:AK:130:TYR:O	8:AK:136:VAL:HG13	2.04	0.58
9:AM:45:ASN:ND2	9:AM:45:ASN:H	2.02	0.58
14:AQ:73:LEU:O	14:AQ:73:LEU:HD13	2.04	0.58
24:AW:31:GLU:O	24:AW:34:GLU:N	2.36	0.58
31:BA:412:A:O2'	31:BA:413:G:P	2.61	0.58
31:BA:687:A:O2'	31:BA:688:G:OP2	2.21	0.58
31:BA:77:C:H2'	31:BA:78:G:C5'	2.25	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:95:G:H2'	31:BA:96:G:H5'	1.85	0.58
31:BA:957:U:H1'	31:BA:960:U:C6	2.39	0.58
31:BA:998(A):C:O2'	31:BA:999:U:H5'	2.04	0.58
39:BL:95:LYS:HA	39:BL:99:LEU:HD23	1.86	0.58
31:BA:972:C:H4'	40:BM:57:LYS:HG3	1.85	0.58
42:BO:60:LEU:HD21	42:BO:66:VAL:CG2	2.33	0.58
6:AG:115:ARG:HH22	43:BP:7:VAL:HB	1.69	0.58
48:BU:22:VAL:C	48:BU:24:ALA:N	2.57	0.58
50:BW:10:LEU:HD13	50:BW:10:LEU:O	2.04	0.58
54:CA:1151:A:H2'	54:CA:1152:A:C8	2.38	0.58
54:CA:1196:U:O2	33:CF:162:GLN:NE2	2.37	0.58
54:CA:1521:G:H2'	54:CA:1522:U:C6	2.39	0.58
54:CA:31:G:O2'	54:CA:32:A:P	2.61	0.58
54:CA:88:C:H3'	54:CA:89:U:C6	2.39	0.58
52:CD:56:C:C6	55:DA:2169:A:N7	2.72	0.58
32:CE:71:VAL:HG21	32:CE:164:VAL:HG22	1.85	0.58
32:CE:221:LEU:O	32:CE:221:LEU:HD13	2.03	0.58
32:CE:7:VAL:HG22	32:CE:8:LYS:N	2.18	0.58
40:CM:38:ILE:HG13	40:CM:38:ILE:O	2.04	0.58
42:CO:48:PRO:O	42:CO:49:ASN:ND2	2.37	0.58
43:CP:97:PRO:HA	43:CP:110:ARG:HD3	1.84	0.58
43:CP:11:ARG:CB	43:CP:11:ARG:HH11	2.16	0.58
43:CP:3:ARG:HH21	6:DG:139:LEU:HD13	1.67	0.58
9:DM:4:TYR:CD2	16:D1:100:VAL:HG11	2.38	0.58
16:D1:95:LEU:HD12	17:D2:11:GLN:NE2	2.18	0.58
26:D4:13:ARG:HB2	26:D4:30:GLU:HA	1.85	0.58
6:DG:67:LYS:HG3	26:D4:6:HIS:CE1	2.38	0.58
55:DA:594:U:OP1	30:D8:61:LEU:HD22	2.04	0.58
55:DA:1056:G:H2'	55:DA:1057:A:OP2	2.04	0.58
55:DA:1973:G:H2'	55:DA:1974:C:C6	2.38	0.58
55:DA:2012:G:H4'	18:DS:96:ILE:HD11	1.86	0.58
55:DA:2038:G:H2'	55:DA:2039:C:C6	2.39	0.58
55:DA:2150:U:H2'	55:DA:2151:G:H8	1.65	0.58
55:DA:2197:U:H1'	55:DA:2198:A:C8	2.38	0.58
55:DA:2662:A:H2'	55:DA:2663:G:O4'	2.04	0.58
55:DA:2688:U:H5	55:DA:2720:U:OP2	1.86	0.58
56:DJ:5:ILE:O	56:DJ:9:LYS:N	2.33	0.58
8:DK:144:VAL:HG22	8:DK:145:VAL:N	2.18	0.58
8:DK:21:VAL:HG21	8:DK:25:TYR:CD1	2.39	0.58
58:DL:14:ALA:CB	58:DL:50:ASP:HB3	2.29	0.58
58:DL:28:GLY:C	58:DL:30:HIS:H	2.07	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:CB:57:G:C5'	21:DV:182:LYS:HZ1	2.01	0.58
21:DV:60:GLU:CG	21:DV:61:LEU:H	2.14	0.58
24:DW:14:ARG:NH1	24:DW:66:GLU:OE1	2.37	0.58
57:DY:100:ASN:ND2	57:DY:100:ASN:O	2.36	0.58
57:DY:89:ALA:HB2	57:DY:125:LEU:HD11	1.85	0.58
13:A0:44:LEU:C	13:A0:44:LEU:HD13	2.24	0.58
13:A0:78:LYS:O	13:A0:82:GLU:HB3	2.04	0.58
1:AA:1826:G:H2'	1:AA:1827:C:H6	1.68	0.58
1:AA:208:C:H2'	1:AA:209:C:C6	2.39	0.58
1:AA:2469:A:H5'	1:AA:2470:G:C8	2.39	0.58
1:AA:273(E):U:C2'	1:AA:273(F):C:H5'	2.33	0.58
1:AA:2850:A:H5'	1:AA:2868:A:H2	1.69	0.58
1:AA:372:G:O2'	1:AA:373:U:OP2	2.22	0.58
1:AA:387:U:OP2	1:AA:387:U:C6	2.56	0.58
3:AD:165:ILE:HD13	3:AD:175:LEU:HD21	1.85	0.58
3:AD:25:THR:HG21	3:AD:82:ILE:N	2.19	0.58
7:AH:54:ARG:HB2	7:AH:55:PRO:HD2	1.84	0.58
7:AH:83:TYR:HA	7:AH:134:SER:CB	2.33	0.58
10:AN:69:ILE:HD12	10:AN:77:ILE:O	2.02	0.58
14:AQ:108:GLY:O	14:AQ:110:LEU:N	2.36	0.58
15:AR:82:LEU:N	15:AR:82:LEU:HD12	2.18	0.58
21:AV:92:SER:O	21:AV:94:GLU:N	2.37	0.58
31:BA:1101:A:H4'	31:BA:1102:A:C4'	2.34	0.58
31:BA:1187:G:H21	44:BQ:60:SER:HB3	1.69	0.58
31:BA:678:U:H2'	31:BA:679:C:C6	2.39	0.58
32:BE:86:GLU:C	32:BE:88:ALA:H	2.07	0.58
34:BG:128:VAL:HG12	34:BG:129:ASN:ND2	2.19	0.58
42:BO:117:ARG:HH21	42:BO:124:LYS:CA	2.17	0.58
54:CA:1181:G:C2	54:CA:1182:G:N2	2.71	0.58
54:CA:1321:C:C5	54:CA:1322:C:N3	2.71	0.58
54:CA:838:G:C6	54:CA:842:C:H1'	2.38	0.58
32:CE:200:ILE:O	32:CE:201:ILE:HD13	2.04	0.58
35:CH:100:VAL:HG23	35:CH:116:THR:O	2.04	0.58
39:CL:50:LEU:HB3	39:CL:55:ALA:O	2.03	0.58
42:CO:38:THR:HG21	42:CO:65:GLU:OE2	2.03	0.58
43:CP:11:ARG:HH11	43:CP:11:ARG:HB3	1.69	0.58
43:CP:7:VAL:HB	6:DG:115:ARG:HH12	1.69	0.58
22:D3:24:LYS:O	22:D3:25:ARG:HG2	2.04	0.58
43:CP:62:ASN:OD1	26:D4:49:PHE:HD2	1.87	0.58
28:D6:13:CYS:HB2	28:D6:22:ALA:O	2.04	0.58
55:DA:1970:A:H5'	55:DA:1971:A:OP1	2.03	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2135:A:H2'	55:DA:2136:C:OP1	2.03	0.58
55:DA:2186:G:H2'	55:DA:2187:G:H8	1.69	0.58
55:DA:223:A:O2'	55:DA:420:C:O2	2.21	0.58
55:DA:228:A:O2'	55:DA:229:A:P	2.62	0.58
55:DA:2068:U:N3	55:DA:2430:A:H2	1.99	0.58
55:DA:270(P):C:H2'	55:DA:270(Q):C:C6	2.39	0.58
55:DA:2776:A:HO2'	55:DA:2781:A:HO2'	1.52	0.58
55:DA:320:A:H2'	5:DF:136:THR:HG21	1.85	0.58
56:DJ:13:SER:C	56:DJ:17:VAL:CG2	2.69	0.58
8:DK:47:LEU:HA	8:DK:50:ARG:HD3	1.86	0.58
58:DL:76:TYR:O	58:DL:77:LEU:C	2.43	0.58
10:DN:96:THR:O	10:DN:97:ARG:O	2.21	0.58
15:DR:105:LEU:HG	15:DR:105:LEU:O	2.03	0.58
21:DV:53:ILE:H	21:DV:71:VAL:CG1	2.16	0.58
57:DY:28:ASN:CB	57:DY:81:VAL:O	2.52	0.58
16:A1:79:PHE:CE2	16:A1:83:LEU:HD13	2.34	0.57
22:A3:6:GLY:O	22:A3:7:LEU:O	2.22	0.57
26:A4:58:ARG:HA	26:A4:61:ARG:CB	2.34	0.57
1:AA:1098:A:H2'	1:AA:1099:G:H5''	1.86	0.57
1:AA:1771:C:C1'	1:AA:1786:A:H8	2.16	0.57
1:AA:1913:A:C8	31:BA:1494:G:H4'	2.39	0.57
1:AA:19:C:H2'	1:AA:20:C:H6	1.69	0.57
1:AA:2181:G:O2'	1:AA:2182:G:H5'	2.05	0.57
1:AA:302:C:O2'	1:AA:303:U:H5'	2.03	0.57
1:AA:638:G:H2'	1:AA:639:U:C6	2.39	0.57
2:AB:5:C:O2'	2:AB:6:C:H5'	2.04	0.57
4:AE:132:HIS:O	4:AE:134:ILE:N	2.37	0.57
4:AE:137:HIS:HB3	4:AE:138:PRO:CD	2.34	0.57
6:AG:127:GLY:HA2	6:AG:166:ASP:CG	2.24	0.57
7:AH:86:GLU:O	7:AH:87:LEU:HG	2.04	0.57
8:AK:128:LEU:O	8:AK:138:ILE:HG22	2.03	0.57
11:AO:47:ASP:HB3	11:AO:48:PRO:O	2.04	0.57
11:AO:63:PRO:O	11:AO:64:LYS:CB	2.52	0.57
15:AR:8:LYS:HB3	15:AR:8:LYS:NZ	2.18	0.57
19:AT:21:PHE:C	19:AT:23:GLU:H	2.07	0.57
21:AV:57:ILE:HG22	21:AV:58:VAL:N	2.17	0.57
23:AZ:80:LEU:N	23:AZ:80:LEU:HD22	2.18	0.57
31:BA:1380:U:O2'	31:BA:1381:U:H5''	2.04	0.57
31:BA:429:U:H1'	31:BA:430:A:H5''	1.86	0.57
31:BA:84:U:H3'	31:BA:85:U:C5	2.39	0.57
33:BF:73:PRO:HA	33:BF:76:VAL:HG13	1.85	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:BG:117:ALA:O	34:BG:121:VAL:HG23	2.04	0.57
31:BA:878:G:H5'	38:BK:89:PRO:HG2	1.86	0.57
40:BM:83:GLU:O	40:BM:86:MET:HB2	2.03	0.57
43:BP:67:GLU:O	43:BP:70:LEU:HD13	2.04	0.57
43:BP:8:GLU:CD	43:BP:22:ILE:HA	2.24	0.57
46:BS:20:VAL:HG23	46:BS:32:TYR:HB2	1.86	0.57
47:BT:59:ILE:CD1	47:BT:73:VAL:HA	2.33	0.57
49:BV:11:VAL:HG23	49:BV:38:SER:HB2	1.86	0.57
50:BW:41:ILE:O	50:BW:44:ALA:HB3	2.04	0.57
51:BX:26:LYS:HZ2	51:BX:26:LYS:HA	1.67	0.57
51:BX:6:ARG:O	51:BX:12:LYS:HE3	2.04	0.57
54:CA:1072:G:H2'	54:CA:1073:U:H6	1.69	0.57
54:CA:1190:G:O2'	54:CA:1191:A:OP2	2.19	0.57
54:CA:940:C:O2'	54:CA:941:G:H5'	2.04	0.57
32:CE:42:ILE:CD1	32:CE:202:PRO:HB2	2.34	0.57
38:CK:39:LEU:HB3	38:CK:45:ILE:HG12	1.85	0.57
17:D2:37:VAL:O	17:D2:37:VAL:HG23	2.04	0.57
55:DA:2349:G:OP2	30:D8:42:ARG:HD3	2.04	0.57
55:DA:1178:C:C2'	55:DA:1179:C:C6	2.75	0.57
55:DA:1543:A:O2'	55:DA:1544:C:P	2.61	0.57
55:DA:1652:A:C2'	55:DA:1653:G:H5'	2.33	0.57
55:DA:2291:U:O2'	55:DA:2374:C:H1'	2.04	0.57
55:DA:654(J):A:HO2'	55:DA:654(K):C:P	2.27	0.57
6:DG:106:LEU:HA	6:DG:110:ALA:HB3	1.85	0.57
55:DA:1059:G:H21	58:DL:126:MET:C	2.07	0.57
10:DN:63:VAL:HG12	10:DN:106:LEU:HD11	1.86	0.57
12:DP:109:VAL:CG1	12:DP:113:GLN:HB3	2.34	0.57
14:DQ:101:LEU:HD13	14:DQ:101:LEU:O	2.04	0.57
14:DQ:89:ARG:HG3	14:DQ:89:ARG:O	2.04	0.57
15:DR:125:ARG:HA	15:DR:128:GLU:HB3	1.84	0.57
15:DR:3:ARG:O	15:DR:4:GLY:C	2.43	0.57
18:DS:66:GLU:O	18:DS:68:ARG:N	2.37	0.57
21:DV:116:VAL:CG1	21:DV:117:LEU:N	2.66	0.57
21:DV:117:LEU:CD1	21:DV:117:LEU:H	2.13	0.57
57:DY:27:VAL:HG22	57:DY:80:VAL:HG11	1.84	0.57
55:DA:1083:U:P	57:DY:47:ASN:OD1	2.62	0.57
57:DY:88:ALA:O	57:DY:91:LYS:N	2.36	0.57
57:DY:90:ALA:O	57:DY:94:VAL:CG2	2.51	0.57
55:DA:2199:A:H5'	23:DZ:50:ARG:NH2	2.19	0.57
23:DZ:49:VAL:HG11	23:DZ:70:VAL:HG11	1.86	0.57
26:A4:57:GLU:O	26:A4:61:ARG:N	2.36	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:A8:62:LEU:HB2	30:A8:63:PRO:HD3	1.86	0.57
1:AA:1096:A:N7	1:AA:1097:U:H1'	2.20	0.57
1:AA:1099:G:H2'	1:AA:1100:C:H6	1.69	0.57
1:AA:1360:A:C5'	1:AA:1361:G:OP2	2.51	0.57
1:AA:1645:G:C5'	1:AA:1646:C:H5'	2.33	0.57
1:AA:2128:C:O2'	1:AA:2173:A:N3	2.36	0.57
1:AA:2298:A:H62	1:AA:2318:G:H8	1.52	0.57
1:AA:2496:C:P	12:AP:81:VAL:HG13	2.44	0.57
1:AA:274:G:OP1	1:AA:274:G:O4'	2.22	0.57
1:AA:448:U:O2'	5:AF:84:VAL:HG13	2.04	0.57
1:AA:991:C:H5'	1:AA:991:C:H6	1.69	0.57
2:AB:8:U:H6	2:AB:8:U:C5'	2.16	0.57
3:AD:85:ASP:HB2	3:AD:92:ILE:HD13	1.85	0.57
4:AE:27:LEU:HA	4:AE:180:ASN:O	2.04	0.57
4:AE:21:VAL:HB	4:AE:22:PRO:HD2	1.86	0.57
1:AA:618:G:H5'	5:AF:107:LYS:HE2	1.85	0.57
9:AM:74:ARG:HH12	9:AM:85:ILE:CD1	2.17	0.57
10:AN:24:VAL:HA	10:AN:39:ILE:HG22	1.87	0.57
10:AN:79:PHE:HE2	10:AN:101:PRO:HB2	1.69	0.57
14:AQ:106:ARG:HB2	14:AQ:106:ARG:NH1	2.18	0.57
20:AU:20:TYR:HE2	20:AU:42:VAL:N	2.03	0.57
21:AV:140:ASP:O	21:AV:141:VAL:HG12	2.03	0.57
23:AZ:51:VAL:HG12	23:AZ:53:VAL:HG23	1.85	0.57
31:BA:723:U:N3	31:BA:1537:U:O2'	2.37	0.57
31:BA:216:G:O2'	31:BA:217:C:C5'	2.52	0.57
31:BA:91:C:H2'	31:BA:92:G:H5''	1.86	0.57
52:BD:48:C:N4	52:BD:59:U:C2	2.72	0.57
35:BH:101:ILE:HD13	35:BH:101:ILE:N	2.19	0.57
38:BK:23:SER:HA	38:BK:63:LEU:CD2	2.34	0.57
38:BK:7:ALA:HB2	38:BK:85:ARG:HD2	1.85	0.57
31:BA:797:C:OP1	41:BN:124:LYS:HE2	2.04	0.57
46:BS:72:ARG:HH11	46:BS:73:LEU:HG	1.68	0.57
47:BT:10:VAL:HG23	47:BT:55:ASP:O	2.04	0.57
50:BW:97:ALA:O	50:BW:99:LEU:HG	2.04	0.57
54:CA:1001:G:H5'	54:CA:1001:G:H8	1.67	0.57
54:CA:1004:A:C2'	54:CA:1005:A:O5'	2.52	0.57
54:CA:256:U:H2'	54:CA:257:G:C8	2.39	0.57
54:CA:794:A:C5'	54:CA:794:A:H8	2.17	0.57
32:CE:163:PHE:CE1	32:CE:215:LEU:HD22	2.39	0.57
32:CE:239:VAL:HG12	32:CE:240:GLN:NE2	2.19	0.57
33:CF:116:VAL:O	33:CF:119:ARG:HB3	2.03	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:CH:9:LYS:HB3	35:CH:112:LEU:HD11	1.85	0.57
41:CN:20:TYR:HB2	41:CN:31:THR:CG2	2.34	0.57
55:DA:1142(A):A:HO2'	55:DA:1143:A:C5'	2.16	0.57
55:DA:1933:G:C2'	55:DA:1934:C:H5''	2.33	0.57
55:DA:213:A:H2'	55:DA:214:G:O4'	2.04	0.57
55:DA:2463:C:O2'	55:DA:2464:C:H5'	2.04	0.57
55:DA:2712:U:H1'	55:DA:2712(A):A:N7	2.18	0.57
55:DA:298:G:P	20:DU:85:VAL:HG22	2.45	0.57
55:DA:222:A:H3'	55:DA:421:U:C5'	2.33	0.57
55:DA:862:G:H2'	55:DA:863:A:O4'	2.04	0.57
3:DD:69:ARG:C	3:DD:71:ASP:H	2.06	0.57
4:DE:36:ARG:HH21	4:DE:86:PRO:HD2	1.70	0.57
4:DE:37:ARG:HD3	4:DE:42:ASP:OD2	2.04	0.57
8:DK:77:LEU:HD13	8:DK:78:THR:N	2.18	0.57
58:DL:18:THR:HG22	58:DL:38:VAL:HG12	1.70	0.57
58:DL:41:PHE:CD2	58:DL:45:THR:OG1	2.57	0.57
12:DP:75:THR:HA	12:DP:88:GLY:C	2.24	0.57
57:DY:29:TYR:CE2	57:DY:32:LEU:HD21	2.39	0.57
23:DZ:41:ARG:HG3	23:DZ:41:ARG:HH11	1.69	0.57
1:AA:1484:G:H2'	1:AA:1485:G:C5'	2.13	0.57
1:AA:1517:G:O2'	1:AA:1518:C:H5'	2.04	0.57
1:AA:2300:G:H2'	1:AA:2301:C:C6	2.39	0.57
1:AA:2317:C:O2'	1:AA:2318:G:H5'	2.04	0.57
1:AA:2319:G:H5''	1:AA:2320:A:OP1	2.04	0.57
1:AA:2554:U:O2	52:BB:74:C:C5	2.57	0.57
1:AA:329:G:C6	20:AU:19:LYS:HG2	2.39	0.57
4:AE:34:VAL:CG1	4:AE:64:LYS:HD3	2.34	0.57
6:AG:81:LYS:H	6:AG:81:LYS:HD3	1.69	0.57
11:AO:62:LEU:HD23	11:AO:64:LYS:HD2	1.86	0.57
12:AP:39:PRO:HA	12:AP:97:VAL:O	2.05	0.57
20:AU:95:LYS:NZ	20:AU:96:ILE:O	2.37	0.57
21:AV:114:GLY:O	21:AV:115:GLY:C	2.42	0.57
21:AV:175:VAL:CB	21:AV:176:PRO:HD2	2.31	0.57
31:BA:328:C:O2'	31:BA:329:A:OP2	2.18	0.57
31:BA:598:U:H2'	31:BA:599:C:C6	2.38	0.57
52:BC:38:A:C2'	52:BC:39:U:H5'	2.34	0.57
33:BF:59:ARG:HG2	33:BF:64:VAL:HG22	1.86	0.57
35:BH:101:ILE:H	35:BH:101:ILE:HD13	1.68	0.57
39:BL:66:ARG:HB3	39:BL:66:ARG:CZ	2.34	0.57
31:BA:1059:C:O2	40:BM:53:PRO:HG3	2.05	0.57
42:BO:89:ARG:HA	42:BO:97:ARG:HA	1.86	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:BU:31:LEU:HD23	48:BU:31:LEU:H	1.68	0.57
50:BW:70:SER:O	50:BW:71:THR:C	2.41	0.57
50:BW:74:LYS:C	50:BW:76:ALA:N	2.58	0.57
54:CA:1004:A:C5'	54:CA:1025:U:O4	2.52	0.57
54:CA:1347:G:C2'	54:CA:1348:U:OP2	2.51	0.57
54:CA:136:C:H2'	54:CA:137:C:C6	2.39	0.57
54:CA:328:C:H4'	54:CA:329:A:C5'	2.34	0.57
54:CA:428:G:C2	54:CA:430:A:N6	2.72	0.57
52:CC:72:C:C2'	52:CC:73:A:H5'	2.34	0.57
35:CH:94:ALA:HB2	35:CH:119:LEU:HG	1.85	0.57
37:CJ:78:ARG:HG3	37:CJ:79:ARG:N	2.19	0.57
46:CS:8:ARG:O	46:CS:9:PHE:HD2	1.86	0.57
49:CV:69:HIS:HB3	49:CV:73:GLU:OE2	2.04	0.57
26:D4:39:CYS:C	26:D4:41:PRO:HD3	2.23	0.57
55:DA:1057:A:H3'	55:DA:1058:U:C5	2.38	0.57
55:DA:1059:G:O2'	58:DL:73:PRO:HG2	2.04	0.57
55:DA:565:C:H4'	55:DA:1253:A:C6	2.38	0.57
55:DA:2614:A:H4'	55:DA:2615:U:OP1	2.04	0.57
55:DA:2815:C:H2'	55:DA:2816:C:H6	1.69	0.57
55:DA:2895:U:H2'	55:DA:2896:C:C6	2.39	0.57
55:DA:943:U:OP2	11:DO:36:LYS:CG	2.52	0.57
4:DE:75:VAL:O	4:DE:76:ARG:HG3	2.05	0.57
6:DG:137:GLU:HB3	6:DG:152:LEU:HD13	1.86	0.57
7:DH:169:VAL:HG22	7:DH:170:ARG:H	1.69	0.57
8:DK:78:THR:O	8:DK:79:ILE:HG12	2.04	0.57
58:DL:8:VAL:O	58:DL:57:ILE:HB	2.04	0.57
58:DL:62:ASP:O	58:DL:63:ARG:CB	2.52	0.57
58:DL:76:TYR:C	58:DL:78:ILE:H	2.07	0.57
9:DM:43:THR:HB	9:DM:46:VAL:HG11	1.84	0.57
57:DY:25:PHE:CD1	57:DY:82:PHE:CZ	2.93	0.57
26:A4:7:PRO:O	26:A4:8:LYS:O	2.22	0.57
28:A6:29:ASN:N	28:A6:29:ASN:ND2	2.52	0.57
30:A8:50:LEU:O	30:A8:51:ALA:CB	2.52	0.57
1:AA:1371:G:O2'	1:AA:1372:U:H5	1.87	0.57
1:AA:1925:C:C2'	1:AA:1925:C:O2	2.48	0.57
1:AA:2472:G:N1	1:AA:2477:C:OP1	2.36	0.57
1:AA:315:G:H2'	1:AA:316:C:C6	2.40	0.57
1:AA:387:U:H6	1:AA:387:U:P	2.27	0.57
1:AA:49:A:H1'	1:AA:51:G:C5	2.38	0.57
1:AA:654(J):A:N1	1:AA:654(L):G:O6	2.38	0.57
1:AA:755:C:H2'	1:AA:756:C:C6	2.39	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AB:40:U:O4	26:A4:1:MET:HG2	2.03	0.57
1:AA:2572:A:N7	4:AE:144:ARG:HD2	2.18	0.57
8:AK:41:GLU:HA	8:AK:44:LEU:HB2	1.87	0.57
12:AP:57:HIS:CE1	12:AP:113:GLN:HE21	2.16	0.57
12:AP:7:MET:CB	12:AP:10:ARG:HE	2.17	0.57
15:AR:36:GLU:OE1	15:AR:41:ARG:HD2	2.04	0.57
18:AS:1:MET:HE2	18:AS:2:GLU:O	2.04	0.57
21:AV:175:VAL:HG12	21:AV:177:PRO:HD3	0.61	0.57
24:AW:16:LEU:HD12	24:AW:16:LEU:O	2.03	0.57
31:BA:1024:G:C2'	31:BA:1025:U:H5''	2.33	0.57
31:BA:1041:A:H2'	31:BA:1042:G:O4'	2.05	0.57
31:BA:1174:G:H2'	31:BA:1175:G:H8	1.69	0.57
31:BA:1251:A:H2'	31:BA:1252:A:C8	2.39	0.57
31:BA:47:C:H5''	31:BA:48:C:OP1	2.05	0.57
31:BA:89:U:H2'	31:BA:90:C:C6	2.39	0.57
52:BB:11:C:H2'	52:BB:12:U:C6	2.39	0.57
34:BG:24:GLU:N	34:BG:27:TYR:CB	2.67	0.57
35:BH:102:ALA:HB1	35:BH:106:PRO:CG	2.34	0.57
36:BI:50:TYR:CE2	36:BI:52:ILE:HD11	2.40	0.57
37:BJ:20:ASP:OD2	37:BJ:22:LEU:HB3	2.04	0.57
42:BO:17:LYS:C	42:BO:17:LYS:HD3	2.25	0.57
47:BT:67:LYS:O	47:BT:68:ARG:C	2.42	0.57
54:CA:1036:G:H3'	54:CA:1037:C:C5	2.38	0.57
54:CA:1106:G:H2'	54:CA:1107:C:C6	2.39	0.57
54:CA:1322:C:H5'	43:CP:100:GLY:HA2	1.85	0.57
54:CA:528:C:H41	42:CO:49:ASN:ND2	2.03	0.57
54:CA:689:C:C2'	54:CA:690:G:H5'	2.34	0.57
32:CE:207:ALA:O	32:CE:209:ARG:N	2.37	0.57
32:CE:96:ARG:N	32:CE:96:ARG:HD2	2.19	0.57
46:CS:53:VAL:O	46:CS:57:ARG:HG2	2.05	0.57
55:DA:2656:U:O4	55:DA:2657:A:C5	2.57	0.57
55:DA:27:G:N2	55:DA:512:G:H2'	2.19	0.57
55:DA:654(G):C:H2'	55:DA:654(H):G:N7	2.19	0.57
3:DD:148:GLU:HB2	3:DD:151:LYS:HD2	1.86	0.57
4:DE:61:ARG:O	4:DE:63:LEU:N	2.38	0.57
7:DH:143:GLN:HE21	7:DH:143:GLN:C	2.08	0.57
7:DH:169:VAL:HG22	7:DH:170:ARG:N	2.18	0.57
58:DL:92:GLY:O	58:DL:135:GLY:O	2.22	0.57
9:DM:46:VAL:HG13	9:DM:48:MET:HG3	1.85	0.57
12:DP:32:TYR:CZ	12:DP:111:GLU:HB3	2.39	0.57
14:DQ:106:ARG:N	14:DQ:110:LEU:HD21	2.20	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:DR:33:LYS:HE2	15:DR:84:GLN:HB3	1.86	0.57
19:DT:49:VAL:HG11	19:DT:83:VAL:HG22	1.86	0.57
21:DV:72:ARG:NH1	21:DV:72:ARG:HG3	2.19	0.57
57:DY:46:GLN:HE21	57:DY:46:GLN:HA	1.70	0.57
1:AA:1088:A:H4'	1:AA:1089:G:C8	2.39	0.57
1:AA:387:U:O2'	1:AA:388:G:OP2	2.19	0.57
1:AA:718:A:H2'	1:AA:719:C:O4'	2.04	0.57
4:AE:132:HIS:O	4:AE:134:ILE:HG23	2.05	0.57
4:AE:87:GLU:O	4:AE:89:ASP:N	2.37	0.57
14:AQ:9:ARG:O	14:AQ:12:PHE:HB2	2.05	0.57
15:AR:98:LYS:HE3	15:AR:98:LYS:CA	2.35	0.57
21:AV:150:LEU:HD23	21:AV:150:LEU:C	2.25	0.57
25:AX:23:LEU:HG	25:AX:50:VAL:HG11	1.85	0.57
31:BA:345:C:O2'	31:BA:346:G:N3	2.38	0.57
31:BA:404:U:H2'	31:BA:405:U:C6	2.39	0.57
52:BC:60:U:H5''	52:BC:61:C:H5	1.70	0.57
41:BN:116:HIS:O	41:BN:117:ASN:HB2	2.04	0.57
48:BU:30:ASP:OD2	48:BU:32:ARG:HB3	2.03	0.57
50:BW:13:LEU:HD12	50:BW:13:LEU:N	2.19	0.57
54:CA:1054:C:O2'	54:CA:1055:A:H5''	2.04	0.57
54:CA:1347:G:H22	54:CA:1373:G:H2'	1.70	0.57
54:CA:437:U:C2'	54:CA:438:G:H5'	2.34	0.57
54:CA:825:G:O2'	54:CA:826:C:H5'	2.05	0.57
54:CA:91:C:C2'	54:CA:92:G:H5''	2.34	0.57
54:CA:95:G:H2'	54:CA:96:G:H5''	1.87	0.57
32:CE:21:ARG:HG3	32:CE:38:GLY:C	2.25	0.57
39:CL:112:LYS:HD3	39:CL:113:LYS:N	2.19	0.57
40:CM:13:HIS:HB3	40:CM:68:HIS:CE1	2.40	0.57
43:CP:87:TYR:HA	43:CP:90:LEU:HG	1.86	0.57
17:D2:41:GLY:H	17:D2:46:VAL:HG13	1.69	0.57
17:D2:49:THR:HB	17:D2:50:PRO:HD2	1.84	0.57
55:DA:2462:U:H1'	55:DA:2491:U:O4	2.04	0.57
55:DA:271(B):G:O2'	55:DA:271(C):U:OP2	2.22	0.57
55:DA:871:U:C2'	55:DA:871:U:O2	2.47	0.57
55:DA:874:G:O2'	55:DA:875:G:H5'	2.05	0.57
3:DD:166:GLN:HE21	3:DD:166:GLN:N	2.02	0.57
5:DF:110:LEU:HD11	5:DF:181:LEU:CD1	2.35	0.57
7:DH:50:VAL:HG22	7:DH:50:VAL:O	2.05	0.57
56:DI:3:LEU:CD2	56:DI:4:ASP:N	2.67	0.57
56:DJ:11:GLU:C	56:DJ:17:VAL:HG11	2.25	0.57
58:DL:101:TRP:HD1	58:DL:101:TRP:N	1.95	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:DL:90:LYS:HE3	58:DL:92:GLY:HA2	1.85	0.57
58:DL:98:ARG:NH1	58:DL:98:ARG:HB3	2.19	0.57
55:DA:195:A:OP1	11:DO:46:LYS:HE2	2.03	0.57
11:DO:96:THR:HG22	11:DO:126:VAL:CG2	2.34	0.57
15:DR:41:ARG:NH2	15:DR:43:GLN:HB3	2.19	0.57
20:DU:75:ILE:C	20:DU:75:ILE:HD13	2.24	0.57
20:DU:88:LYS:C	20:DU:90:LEU:H	2.07	0.57
21:DV:111:VAL:O	21:DV:112:ARG:O	2.22	0.57
57:DY:32:LEU:N	57:DY:32:LEU:HD22	2.19	0.57
16:A1:88:ILE:C	16:A1:90:VAL:N	2.57	0.57
17:A2:49:THR:CB	17:A2:50:PRO:CD	2.81	0.57
22:A3:83:PRO:O	22:A3:84:LEU:O	2.23	0.57
1:AA:1312:U:H4'	1:AA:1313:U:O5'	2.04	0.57
1:AA:1320:C:H5	1:AA:1329:U:H5'	1.69	0.57
1:AA:1379:A:O2'	1:AA:1380:G:P	2.61	0.57
1:AA:1796:U:H2'	1:AA:1797:C:H6	1.67	0.57
1:AA:2426:A:C4'	1:AA:2427:C:OP2	2.51	0.57
1:AA:2531:A:H61	1:AA:2662:A:N6	2.02	0.57
1:AA:27:G:N2	1:AA:512:G:H2'	2.18	0.57
1:AA:30:G:H2'	1:AA:31:C:H6	1.68	0.57
1:AA:363(B):G:O2'	1:AA:363(C):G:H5'	2.04	0.57
1:AA:412:A:H2'	1:AA:413:C:H5'	1.87	0.57
1:AA:68:G:H3'	1:AA:69:C:C6	2.39	0.57
5:AF:107:LYS:HB3	5:AF:206:ILE:CG2	2.34	0.57
5:AF:113:ALA:HB1	5:AF:186:ILE:HG21	1.86	0.57
5:AF:122:LYS:HD2	5:AF:191:ARG:HG2	1.85	0.57
5:AF:124:LEU:O	5:AF:126:VAL:N	2.37	0.57
6:AG:99:MET:HG3	6:AG:100:TRP:N	2.18	0.57
6:AG:72:ARG:HH11	6:AG:72:ARG:HG3	1.68	0.57
7:AH:89:ILE:HD11	7:AH:129:THR:CB	2.34	0.57
11:AO:106:LEU:O	11:AO:107:LYS:HB2	2.05	0.57
11:AO:94:GLU:O	11:AO:96:THR:HG23	2.05	0.57
12:AP:20:ALA:HA	12:AP:99:PRO:HG2	1.87	0.57
15:AR:16:ARG:HE	15:AR:19:LEU:HD11	1.68	0.57
18:AS:59:VAL:CG2	18:AS:65:LEU:H	2.15	0.57
20:AU:94:LYS:NZ	20:AU:101:LYS:NZ	2.52	0.57
20:AU:62:GLU:CD	20:AU:63:LYS:H	2.08	0.57
20:AU:8:LYS:O	20:AU:27:VAL:HG21	2.04	0.57
21:AV:135:GLU:HG3	21:AV:136:PHE:CD2	2.39	0.57
31:BA:1177:G:H2'	31:BA:1178:G:N3	2.19	0.57
31:BA:1316:G:C2'	31:BA:1317:C:H5''	2.35	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:1320:C:C4	49:BV:36:ARG:HG3	2.39	0.57
31:BA:922:G:N3	31:BA:1398:A:H2	2.02	0.57
52:BC:57:G:H2'	52:BC:58:A:H5''	1.86	0.57
32:BE:141:GLU:O	32:BE:145:LEU:HB2	2.04	0.57
32:BE:82:ARG:HA	32:BE:92:TYR:CE1	2.40	0.57
34:BG:132:ARG:HG3	34:BG:132:ARG:O	2.05	0.57
42:BO:62:SER:O	42:BO:64:TYR:N	2.38	0.57
43:BP:81:LEU:HD21	43:BP:88:ARG:NH1	2.19	0.57
49:BV:36:ARG:HH21	49:BV:75:ALA:HB3	1.68	0.57
54:CA:939:G:H2'	54:CA:940:C:C6	2.38	0.57
52:CD:37:MIA:H131	52:CD:37:MIA:N1	2.20	0.57
34:CG:111:ALA:HB2	34:CG:120:LEU:CD1	2.34	0.57
36:CI:22:GLU:O	36:CI:26:ILE:HG13	2.05	0.57
38:CK:100:ILE:HB	38:CK:125:ARG:NH1	2.20	0.57
51:CX:10:ARG:HG2	51:CX:13:ILE:HD12	1.86	0.57
26:D4:42:PHE:C	26:D4:42:PHE:CD1	2.77	0.57
55:DA:1204:A:O2'	55:DA:1205:U:H5''	2.04	0.57
55:DA:1472:A:H2'	55:DA:1473:G:O4'	2.04	0.57
55:DA:1480:G:O6	55:DA:1510:A:C2	2.58	0.57
55:DA:1885:A:H3'	55:DA:1886:C:C6	2.39	0.57
55:DA:2427:C:H5''	55:DA:2428:G:OP1	2.05	0.57
55:DA:2602:A:OP2	55:DA:2603:G:H5''	2.04	0.57
4:DE:37:ARG:HB3	4:DE:42:ASP:CG	2.25	0.57
7:DH:13:LYS:HE2	7:DH:13:LYS:CA	2.30	0.57
7:DH:12:PRO:HG3	7:DH:48:GLY:O	2.05	0.57
58:DL:140:GLY:O	58:DL:141:ALA:CB	2.47	0.57
55:DA:1012:U:O4	9:DM:25:ARG:HA	2.04	0.57
9:DM:68:GLU:HG2	9:DM:88:GLU:OE2	2.04	0.57
15:DR:66:VAL:HG12	15:DR:67:SER:N	2.19	0.57
19:DT:49:VAL:HG11	19:DT:83:VAL:CG2	2.35	0.57
57:DY:122:VAL:O	57:DY:123:GLU:C	2.40	0.57
57:DY:135:ARG:HH11	57:DY:138:LEU:CG	2.06	0.57
57:DY:24:PHE:O	57:DY:25:PHE:C	2.43	0.57
57:DY:50:ARG:N	57:DY:83:TYR:CB	2.68	0.57
13:A0:37:THR:HG22	13:A0:39:PRO:CD	2.26	0.57
1:AA:1299:G:H5''	1:AA:1300:U:H5''	1.86	0.57
1:AA:1371:G:HO2'	1:AA:1372:U:H5	1.51	0.57
1:AA:1777:U:O2'	1:AA:1778:U:H5'	2.05	0.57
1:AA:752:A:C6	1:AA:1781:C:O4'	2.58	0.57
1:AA:2820:A:N6	4:AE:192:ASN:CA	2.66	0.57
1:AA:322:A:OP1	5:AF:169:ASN:HB2	2.04	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:401:A:H2'	1:AA:402:A:O4'	2.04	0.57
1:AA:654(D):G:N3	1:AA:654(D):G:H2'	2.18	0.57
1:AA:75:G:H4'	24:AW:55:ARG:HH21	1.69	0.57
1:AA:946:G:H2'	1:AA:947:G:O5'	2.02	0.57
2:AB:86:G:O2'	2:AB:87:G:H5'	2.05	0.57
4:AE:199:ARG:NH1	4:AE:199:ARG:HB2	2.19	0.57
5:AF:178:PRO:HB3	5:AF:198:ALA:HB1	1.87	0.57
6:AG:67:LYS:NZ	26:A4:6:HIS:CE1	2.71	0.57
7:AH:26:VAL:HG13	7:AH:27:LYS:H	1.70	0.57
14:AQ:36:TYR:CD2	14:AQ:52:SER:HB3	2.40	0.57
21:AV:11:GLU:CG	21:AV:12:GLY:N	2.66	0.57
24:AW:48:HIS:O	24:AW:52:ASP:HB2	2.04	0.57
24:AW:68:ARG:HG3	24:AW:68:ARG:NH1	2.20	0.57
31:BA:1449:C:H3'	31:BA:1450:U:H4'	1.85	0.57
31:BA:413:G:O2'	31:BA:428:G:N2	2.38	0.57
31:BA:957:U:H1'	31:BA:960:U:C5	2.39	0.57
32:BE:9:GLU:O	32:BE:12:GLU:HG3	2.03	0.57
36:BI:10:LEU:HD12	36:BI:10:LEU:N	2.20	0.57
44:BQ:12:ARG:HG2	44:BQ:14:PRO:CD	2.28	0.57
48:BU:23:LYS:HE2	48:BU:57:GLY:O	2.04	0.57
49:BV:6:LYS:H	49:BV:6:LYS:HD2	1.69	0.57
54:CA:1004:A:O4'	54:CA:1036:G:O6	2.22	0.57
54:CA:1376:U:H2'	54:CA:1377:A:H8	1.68	0.57
54:CA:50:A:HO2'	54:CA:52:G:H8	1.46	0.57
52:CB:10:G:H3'	52:CB:11:C:H5	1.69	0.57
52:CB:37:MIA:C16	52:CB:37:MIA:N6	2.67	0.57
32:CE:17:PHE:HB3	32:CE:44:LEU:HD11	1.87	0.57
32:CE:209:ARG:HD3	32:CE:240:GLN:OE1	2.04	0.57
38:CK:1:MET:CE	38:CK:1:MET:H3	2.16	0.57
39:CL:112:LYS:HD3	39:CL:113:LYS:O	2.05	0.57
39:CL:23:ASN:HB2	39:CL:25:LYS:HG2	1.87	0.57
49:CV:7:LYS:HB2	49:CV:7:LYS:NZ	2.20	0.57
55:DA:1138:G:H21	9:DM:106:MET:CE	2.13	0.57
55:DA:1678:G:N2	55:DA:1989:G:H1	2.02	0.57
55:DA:2741:A:H2'	55:DA:2742:C:O4'	2.05	0.57
3:DD:43:ARG:HD2	3:DD:49:ILE:HG22	1.87	0.57
4:DE:34:VAL:HG21	4:DE:77:ILE:CG2	2.34	0.57
7:DH:37:VAL:CG1	7:DH:38:SER:H	2.16	0.57
56:DJ:13:SER:HG	56:DJ:17:VAL:HG22	1.64	0.57
56:DJ:9:LYS:NZ	56:DJ:9:LYS:O	2.37	0.57
11:DO:50:ARG:HB2	11:DO:50:ARG:HH21	1.67	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:DO:9:ASN:CB	11:DO:10:PRO:HD2	2.35	0.57
15:DR:56:GLY:C	15:DR:57:PHE:O	2.41	0.57
20:DU:77:PRO:O	20:DU:78:ALA:HB2	2.05	0.57
21:DV:120:ILE:HB	21:DV:171:ILE:N	2.18	0.57
21:DV:140:ASP:C	21:DV:141:VAL:HG23	2.25	0.57
21:DV:192:ALA:C	21:DV:194:PRO:HD2	2.24	0.57
57:DY:75:GLN:HE21	57:DY:75:GLN:CA	2.16	0.57
57:DY:49:ALA:C	57:DY:83:TYR:CD1	2.73	0.57
16:A1:83:LEU:HG	16:A1:88:ILE:HG13	1.86	0.57
16:A1:92:ARG:O	16:A1:94:ASN:N	2.37	0.57
1:AA:1359:A:OP2	1:AA:1359:A:C8	2.57	0.57
1:AA:1291:C:H5''	1:AA:1536:A:H5''	1.86	0.57
1:AA:2095:C:H2'	1:AA:2096:U:O4'	2.05	0.57
1:AA:234:C:H2'	1:AA:235:U:H6	1.70	0.57
1:AA:521:G:H2'	1:AA:522:G:C8	2.38	0.57
2:AB:31:C:H2'	2:AB:31:C:O2	2.04	0.57
2:AB:74:U:C3'	2:AB:75:G:H5''	2.35	0.57
4:AE:197:ILE:HD11	4:AE:199:ARG:CZ	2.35	0.57
5:AF:123:LEU:CD1	5:AF:192:LEU:HD22	2.34	0.57
6:AG:23:PHE:HB2	6:AG:25:TYR:CE2	2.40	0.57
6:AG:47:LYS:HD3	6:AG:81:LYS:HB2	1.85	0.57
9:AM:137:LYS:HZ3	9:AM:137:LYS:HA	1.70	0.57
12:AP:134:ARG:NH2	21:AV:122:ARG:NH1	2.52	0.57
12:AP:42:ILE:HG22	12:AP:47:ILE:HG13	1.86	0.57
21:AV:122:ARG:HD3	21:AV:123:ASP:OD2	2.04	0.57
21:AV:144:LEU:HD13	21:AV:146:ILE:O	2.05	0.57
31:BA:751:U:H2'	31:BA:752:G:O4'	2.04	0.57
31:BA:85:U:C2'	31:BA:86:U:OP1	2.53	0.57
31:BA:1055:A:H4'	33:BF:161:GLU:OE2	2.04	0.57
34:BG:11:LEU:C	34:BG:13:ARG:N	2.55	0.57
36:BI:68:PRO:HG3	36:BI:71:ARG:NH2	2.20	0.57
31:BA:1199:U:H4'	40:BM:54:PHE:CE1	2.39	0.57
40:BM:7:LYS:HG3	40:BM:71:LEU:HD13	1.85	0.57
43:BP:4:ILE:HG12	43:BP:5:ALA:N	2.19	0.57
46:BS:18:ARG:O	46:BS:20:VAL:HG12	2.04	0.57
46:BS:34:GLU:OE2	46:BS:55:ARG:HD3	2.05	0.57
54:CA:1053:G:C4'	54:CA:1054:C:H5'	2.35	0.57
54:CA:1292:U:O2'	54:CA:1293:G:H5'	2.05	0.57
54:CA:164:U:H2'	54:CA:165:C:H6	1.68	0.57
35:CH:142:LEU:O	35:CH:143:ARG:NH1	2.29	0.57
39:CL:86:VAL:O	39:CL:90:PRO:HA	2.04	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:CM:46:ARG:HA	40:CM:64:GLU:HA	1.86	0.57
49:CV:11:VAL:HG23	49:CV:38:SER:HB2	1.86	0.57
54:CA:263:A:OP2	50:CW:79:ARG:NH1	2.34	0.57
55:DA:1043:C:H2'	55:DA:1044:G:C5'	2.30	0.57
55:DA:1079:C:C3'	55:DA:1080:A:C8	2.86	0.57
55:DA:1926:U:C6	55:DA:1928:A:OP2	2.57	0.57
55:DA:2334:G:H4'	55:DA:2335:A:OP2	2.04	0.57
55:DA:270(P):C:O2'	55:DA:270(Q):C:H5'	2.04	0.57
55:DA:779:U:OP1	3:DD:49:ILE:HG12	2.04	0.57
55:DA:865:C:H5'	55:DA:866:A:OP1	2.04	0.57
55:DA:729:G:C5	3:DD:208:LYS:HB2	2.40	0.57
3:DD:181:GLU:CA	3:DD:272:ALA:HB3	2.35	0.57
3:DD:65:ILE:CD1	3:DD:65:ILE:N	2.67	0.57
55:DA:2683:C:H4'	4:DE:13:ARG:NH2	2.19	0.57
7:DH:40:GLU:O	7:DH:41:MET:HB2	2.05	0.57
7:DH:28:GLY:HA3	7:DH:79:VAL:HB	1.86	0.57
58:DL:106:GLU:HA	58:DL:109:LYS:CB	2.35	0.57
58:DL:7:VAL:CG1	58:DL:58:THR:H	2.03	0.57
9:DM:36:GLY:O	9:DM:42:TRP:HB2	2.05	0.57
10:DN:75:SER:CB	15:DR:74:ARG:HH12	2.18	0.57
57:DY:122:VAL:CG1	57:DY:126:ALA:CB	2.78	0.57
57:DY:30:GLN:OE1	57:DY:79:ALA:HB1	2.05	0.57
1:AA:1041:C:H2'	1:AA:1042:G:C8	2.39	0.57
1:AA:1168:G:H2'	1:AA:1169:G:H8	1.68	0.57
1:AA:1281:G:H2'	1:AA:1282:U:O4'	2.04	0.57
1:AA:1290:C:H2'	1:AA:1291:C:C6	2.40	0.57
1:AA:1954:G:O2'	1:AA:1956:U:H5	1.86	0.57
1:AA:458:G:O2'	1:AA:459:U:O5'	2.22	0.57
1:AA:888:C:HO2'	1:AA:889:C:P	2.24	0.57
1:AA:893:C:C2	1:AA:894:C:C5	2.92	0.57
4:AE:179:GLU:HB3	4:AE:181:LEU:HD23	1.86	0.57
5:AF:187:VAL:HG11	11:AO:6:LEU:HD21	1.86	0.57
10:AN:99:PHE:CD2	10:AN:99:PHE:N	2.73	0.57
11:AO:80:TYR:CE1	11:AO:111:ARG:HB3	2.40	0.57
12:AP:12:GLN:HE21	12:AP:73:PRO:HD3	1.69	0.57
14:AQ:14:VAL:HG21	14:AQ:89:ARG:HH11	1.70	0.57
19:AT:41:ASN:O	19:AT:45:THR:HG23	2.04	0.57
20:AU:13:VAL:HG22	20:AU:14:LEU:O	2.04	0.57
31:BA:1273:G:H3'	31:BA:1274:G:H8	1.70	0.57
31:BA:1469:G:H2'	31:BA:1470:G:C8	2.40	0.57
31:BA:235:C:H1'	47:BT:61:GLU:OE1	2.05	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:501:C:O2'	31:BA:502:G:H5'	2.05	0.57
31:BA:553:A:H2'	31:BA:554:C:C6	2.39	0.57
32:BE:45:GLN:O	32:BE:49:GLU:HG2	2.03	0.57
33:BF:6:HIS:CD2	33:BF:7:PRO:HD2	2.39	0.57
34:BG:11:LEU:O	34:BG:12:CYS:C	2.43	0.57
46:BS:38:TYR:CZ	46:BS:50:LYS:HB3	2.40	0.57
49:BV:20:LEU:HD22	49:BV:43:GLU:HG2	1.85	0.57
53:C1:56:U:H3'	53:C1:56:U:O2	2.03	0.57
54:CA:1019:C:O2'	54:CA:1020:U:H5'	2.05	0.57
54:CA:1032:A:N7	54:CA:1032(A):G:H1'	2.19	0.57
54:CA:188:U:C2'	54:CA:189:U:H5''	2.27	0.57
54:CA:190:G:N3	54:CA:190:G:H2'	2.20	0.57
54:CA:197:A:N6	54:CA:221:C:C5'	2.68	0.57
54:CA:337:C:H2'	54:CA:338:A:C8	2.40	0.57
54:CA:965:A:H4'	54:CA:966:G:O5'	2.03	0.57
52:CB:62:C:O2'	52:CB:63:G:H5'	2.05	0.57
35:CH:147:ASP:O	35:CH:151:LEU:HG	2.04	0.57
49:CV:11:VAL:HG13	49:CV:16:LEU:HD22	1.85	0.57
16:D1:62:ILE:HG23	16:D1:76:TYR:CE1	2.40	0.57
17:D2:76:LYS:HB2	17:D2:81:TYR:HB3	1.84	0.57
26:D4:55:ARG:HD2	26:D4:56:VAL:HG23	1.85	0.57
27:D5:56:LYS:HG2	27:D5:58:LEU:HG	1.87	0.57
55:DA:1165:U:C2	55:DA:1166:C:C5	2.92	0.57
55:DA:1485:G:O2'	55:DA:1486:A:H5'	2.05	0.57
55:DA:1731:G:H8	55:DA:1731:G:OP2	1.88	0.57
55:DA:2723:C:C4'	13:D0:1:MET:HG2	2.34	0.57
55:DA:481:G:O2'	55:DA:507:A:N6	2.37	0.57
55:DA:895:U:H5''	55:DA:896:A:OP2	2.05	0.57
3:DD:101:GLU:OE1	3:DD:103:ARG:HD3	2.05	0.57
56:DI:28:LYS:C	56:DJ:2:ALA:HB1	2.24	0.57
58:DL:111:LYS:CD	58:DL:111:LYS:N	2.67	0.57
58:DL:95:LYS:C	58:DL:97:GLY:N	2.56	0.57
12:DP:20:ALA:O	12:DP:21:THR:HG23	2.04	0.57
21:DV:110:GLY:HA3	21:DV:144:LEU:O	2.05	0.57
21:DV:110:GLY:O	21:DV:111:VAL:C	2.43	0.57
21:DV:179:ASP:O	21:DV:180:VAL:CB	2.52	0.57
57:DY:61:LEU:C	57:DY:63:LEU:N	2.58	0.57
16:A1:72:HIS:CE1	16:A1:107:ALA:HA	2.37	0.57
28:A6:9:LEU:HD23	28:A6:10:LEU:H	1.70	0.57
1:AA:99:U:H4'	1:AA:102:G:H1'	1.85	0.57
1:AA:1614:A:H61	18:AS:88:ARG:H	1.53	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2320:A:H1'	1:AA:2321:G:C6	2.40	0.57
1:AA:2377:A:H2'	1:AA:2378:A:C8	2.40	0.57
1:AA:2554:U:C2	52:BB:74:C:H5	2.23	0.57
1:AA:2581:G:H4'	1:AA:2582:G:C8	2.40	0.57
1:AA:971:C:H2'	1:AA:972:G:C5'	2.35	0.57
3:AD:109:ASP:HB2	3:AD:197:GLY:HA2	1.86	0.57
4:AE:4:ILE:HA	4:AE:49:LEU:CD1	2.35	0.57
4:AE:77:ILE:O	4:AE:78:LEU:O	2.23	0.57
14:AQ:24:LEU:HD22	14:AQ:24:LEU:N	2.20	0.57
15:AR:26:ASP:CB	15:AR:91:ARG:HG2	2.35	0.57
20:AU:87:LYS:HA	20:AU:92:ASN:HA	1.86	0.57
20:AU:91:GLU:HG3	20:AU:92:ASN:N	2.09	0.57
53:B1:53:U:H1'	53:B1:54:U:OP1	2.05	0.57
31:BA:1027:C:O2'	31:BA:1028:C:OP1	2.20	0.57
31:BA:1035:A:H3'	31:BA:1036:G:H5''	1.87	0.57
32:BE:187:LEU:HD23	32:BE:201:ILE:O	2.05	0.57
32:BE:55:PHE:HA	32:BE:58:ILE:HG12	1.86	0.57
33:BF:119:ARG:O	33:BF:123:GLN:HG3	2.05	0.57
33:BF:22:TRP:CB	33:BF:59:ARG:HB2	2.35	0.57
43:BP:48:LEU:HD23	43:BP:48:LEU:H	1.70	0.57
54:CA:792:A:H2'	54:CA:794:A:C6	2.37	0.57
54:CA:814:A:N7	54:CA:816:A:C4	2.73	0.57
54:CA:865:A:H2'	54:CA:866:C:C6	2.40	0.57
52:CB:60:U:H5'	52:CB:61:C:OP2	2.04	0.57
52:CC:50:U:H2'	52:CC:51:U:O4'	2.05	0.57
32:CE:168:THR:HG23	32:CE:169:LYS:N	2.20	0.57
32:CE:82:ARG:O	32:CE:86:GLU:HG3	2.05	0.57
54:CA:1374:A:O2'	37:CJ:28:ASN:HB3	2.05	0.57
38:CK:111:ILE:O	38:CK:112:LEU:HB3	2.05	0.57
39:CL:118:LYS:HB3	39:CL:118:LYS:NZ	2.20	0.57
50:CW:100:ILE:HG13	50:CW:102:GLY:N	2.08	0.57
26:D4:12:ALA:CB	26:D4:29:PRO:HA	2.34	0.57
55:DA:1079:C:O2	58:DL:129:GLY:HA3	2.04	0.57
55:DA:1799:G:H4'	55:DA:1800:C:O5'	2.05	0.57
55:DA:2776:A:O2'	55:DA:2781:A:H4'	2.04	0.57
55:DA:2787:C:O2'	4:DE:61:ARG:CD	2.52	0.57
55:DA:446:G:H4'	55:DA:449:A:N3	2.20	0.57
55:DA:803:U:C2'	55:DA:804:A:H5'	2.35	0.57
55:DA:860:U:C5	55:DA:917:A:H2	2.20	0.57
3:DD:242:ARG:N	3:DD:242:ARG:HD2	2.20	0.57
3:DD:80:ALA:HB3	3:DD:94:LEU:CD1	2.34	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:DE:30:PRO:O	4:DE:32:PRO:HD3	2.04	0.57
4:DE:54:GLN:O	4:DE:55:ASN:HB2	2.04	0.57
4:DE:62:PRO:O	4:DE:63:LEU:HG	2.05	0.57
56:DI:21:LYS:HA	56:DI:24:ILE:CD1	2.33	0.57
56:DI:28:LYS:C	56:DJ:2:ALA:CB	2.74	0.57
12:DP:92:GLY:C	12:DP:93:TYR:CD1	2.78	0.57
15:DR:54:ARG:HH11	15:DR:54:ARG:HG2	1.69	0.57
21:DV:178:GLU:OE1	21:DV:180:VAL:CA	2.53	0.57
25:DX:7:LYS:HZ1	25:DX:32:GLN:HG3	1.70	0.57
57:DY:71:LEU:HB3	57:DY:113:GLN:HB3	0.60	0.57
57:DY:122:VAL:O	57:DY:126:ALA:N	2.31	0.57
57:DY:130:THR:HG21	56:DJ:14:GLN:OE1	2.05	0.57
57:DY:2:PRO:CG	57:DY:3:ASN:N	2.52	0.57
23:DZ:85:LEU:N	23:DZ:85:LEU:HD22	2.20	0.57
1:AA:1316:U:H2'	1:AA:1317:A:H8	1.70	0.56
1:AA:2893:G:H8	1:AA:2893:G:OP2	1.88	0.56
1:AA:220:G:H2'	1:AA:427:U:O4	2.05	0.56
1:AA:639:U:H2'	1:AA:640:C:H6	1.69	0.56
1:AA:788:A:O2'	1:AA:789:A:OP2	2.23	0.56
2:AB:14:U:H4'	2:AB:106:G:N2	2.20	0.56
5:AF:198:ALA:O	5:AF:201:VAL:HG12	2.05	0.56
6:AG:104:GLU:O	6:AG:108:ASN:HB2	2.05	0.56
6:AG:118:ARG:NH2	26:A4:42:PHE:CZ	2.73	0.56
6:AG:118:ARG:NH2	26:A4:42:PHE:HZ	2.03	0.56
6:AG:124:SER:HB2	6:AG:131:TYR:CE1	2.40	0.56
6:AG:145:THR:O	6:AG:146:TYR:HB3	2.05	0.56
6:AG:16:ARG:HG2	6:AG:16:ARG:NH1	2.20	0.56
14:AQ:34:HIS:HB3	14:AQ:53:SER:HB3	1.86	0.56
21:AV:140:ASP:O	21:AV:141:VAL:CG1	2.53	0.56
24:AW:33:MET:HG2	24:AW:37:PHE:CE1	2.39	0.56
31:BA:1065:U:O2'	31:BA:1066:C:P	2.62	0.56
31:BA:1106:G:H5''	33:BF:172:ARG:CG	2.32	0.56
31:BA:1239:A:H2'	31:BA:1298:C:H42	1.70	0.56
31:BA:913:A:O2'	31:BA:914:A:OP2	2.21	0.56
32:BE:74:LYS:O	32:BE:78:GLN:HG3	2.05	0.56
33:BF:8:ILE:HG23	33:BF:16:ARG:HG2	1.86	0.56
34:BG:19:LEU:CD1	34:BG:21:LEU:HD23	2.35	0.56
35:BH:141:GLN:HA	35:BH:143:ARG:HH21	1.70	0.56
31:BA:878:G:C5'	38:BK:89:PRO:HG2	2.35	0.56
49:BV:76:PRO:HB2	49:BV:78:ARG:CD	2.35	0.56
50:BW:48:LYS:HB3	50:BW:51:GLU:CG	2.35	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:BX:3:LYS:HA	51:BX:11:GLY:HA2	1.87	0.56
54:CA:1015:A:H2'	54:CA:1016:A:C8	2.39	0.56
54:CA:328:C:H4'	54:CA:329:A:H5'	1.86	0.56
54:CA:990:C:H2'	54:CA:991:U:C6	2.40	0.56
52:CB:58:A:H4'	52:CB:59:U:OP1	2.05	0.56
32:CE:80:ILE:CD1	32:CE:208:ILE:HG23	2.32	0.56
32:CE:86:GLU:C	32:CE:88:ALA:H	2.07	0.56
34:CG:131:ARG:H	34:CG:131:ARG:HD3	1.69	0.56
34:CG:173:TRP:NE1	34:CG:174:LEU:HG	2.19	0.56
35:CH:90:VAL:O	35:CH:120:THR:HA	2.05	0.56
37:CJ:111:ARG:HB3	37:CJ:111:ARG:NH1	2.20	0.56
54:CA:1297:C:C2'	37:CJ:114:ARG:HH22	2.17	0.56
38:CK:49:GLU:HG3	38:CK:49:GLU:O	2.04	0.56
39:CL:47:LEU:HD22	39:CL:47:LEU:N	2.16	0.56
43:CP:82:MET:O	43:CP:84:ILE:N	2.37	0.56
50:CW:14:LYS:HG3	50:CW:17:ARG:NH2	2.19	0.56
16:D1:83:LEU:CA	16:D1:88:ILE:HD11	2.30	0.56
55:DA:1048:A:C8	55:DA:1049:C:H5	2.23	0.56
55:DA:196:A:H5'	55:DA:197:A:OP2	2.04	0.56
55:DA:803:U:O2'	55:DA:804:A:H5'	2.05	0.56
55:DA:1797:C:H4'	3:DD:257:LEU:O	2.05	0.56
4:DE:78:LEU:HD21	4:DE:79:ARG:NE	2.20	0.56
5:DF:132:VAL:O	5:DF:133:ASN:C	2.44	0.56
5:DF:174:VAL:O	5:DF:174:VAL:HG22	2.03	0.56
7:DH:152:ARG:O	7:DH:153:LYS:CD	2.51	0.56
9:DM:120:LEU:CD1	9:DM:122:VAL:HG23	2.35	0.56
11:DO:91:PHE:CE2	11:DO:95:VAL:HG22	2.39	0.56
21:DV:165:VAL:O	21:DV:167:PRO:HD3	2.04	0.56
57:DY:71:LEU:HD22	57:DY:72:ASP:HA	1.87	0.56
57:DY:50:ARG:NH2	57:DY:83:TYR:CE1	2.73	0.56
57:DY:93:LEU:CD2	57:DY:97:ALA:CB	2.71	0.56
22:A3:53:MET:HA	22:A3:58:THR:O	2.04	0.56
1:AA:1698:A:H1'	1:AA:1699:G:O3'	2.05	0.56
1:AA:1688:U:O2	1:AA:1700:A:H8	1.89	0.56
1:AA:1930:G:C2'	1:AA:1931:U:OP2	2.52	0.56
1:AA:2286:A:O5'	28:A6:28:ARG:NE	2.38	0.56
1:AA:2533:A:H2'	1:AA:2534:A:C5'	2.24	0.56
1:AA:270(F):U:H2'	1:AA:270(G):C:H6	1.69	0.56
1:AA:811:U:OP2	11:AO:21:ARG:O	2.22	0.56
1:AA:856:C:H4'	1:AA:857:C:OP1	2.06	0.56
2:AB:31:C:H4'	6:AG:29:TRP:CH2	2.41	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:AH:109:PHE:CE1	7:AH:152:ARG:HD3	2.39	0.56
7:AH:75:ALA:O	7:AH:79:VAL:HG22	2.05	0.56
10:AN:11:ALA:O	10:AN:12:ASP:HB3	2.05	0.56
14:AQ:98:VAL:O	14:AQ:101:LEU:HB3	2.06	0.56
15:AR:26:ASP:HB2	15:AR:91:ARG:HG2	1.87	0.56
1:AA:1341:U:O4'	19:AT:57:LEU:HD22	2.04	0.56
20:AU:75:ILE:HB	20:AU:80:GLY:H	1.70	0.56
25:AX:8:LEU:CD1	25:AX:31:LEU:HA	2.34	0.56
31:BA:1026:G:C6	31:BA:1036:G:C2	2.93	0.56
31:BA:1503:A:O2'	31:BA:1504:G:OP1	2.22	0.56
31:BA:280:C:H5'	31:BA:281:G:C8	2.39	0.56
33:BF:116:VAL:O	33:BF:119:ARG:HB3	2.04	0.56
54:CA:1191:A:OP1	33:CF:3:ASN:ND2	2.38	0.56
54:CA:1446:A:H1'	15:DR:125:ARG:NH2	2.12	0.56
54:CA:328:C:O2'	54:CA:329:A:P	2.62	0.56
54:CA:574:A:H5''	54:CA:575:G:OP2	2.05	0.56
54:CA:872:A:C2'	54:CA:873:A:H3'	2.35	0.56
54:CA:973:G:C4	40:CM:55:LYS:HE2	2.40	0.56
52:CD:71:G:H2'	52:CD:72:C:H6	1.70	0.56
32:CE:4:GLU:HG2	32:CE:5:ILE:H	1.69	0.56
33:CF:175:LEU:H	33:CF:175:LEU:CD1	2.14	0.56
33:CF:188:LEU:O	33:CF:189:ALA:HB2	2.05	0.56
33:CF:70:VAL:HG12	33:CF:71:ALA:H	1.69	0.56
34:CG:148:VAL:HG12	34:CG:149:ALA:H	1.69	0.56
37:CJ:102:ARG:O	37:CJ:106:GLN:HG3	2.04	0.56
37:CJ:15:ASP:O	37:CJ:19:GLY:HA2	2.05	0.56
40:CM:6:ILE:CD1	40:CM:72:VAL:HB	2.36	0.56
49:CV:24:ALA:C	49:CV:26:GLY:N	2.58	0.56
30:D8:22:VAL:CG2	30:D8:53:PRO:HB2	2.36	0.56
55:DA:105:C:H2'	55:DA:106:C:H6	1.70	0.56
55:DA:2564:A:OP1	55:DA:2648:C:H4'	2.05	0.56
55:DA:2654:A:C8	55:DA:2656:U:O2	2.57	0.56
55:DA:270(O):U:H2'	55:DA:270(O):U:O2	2.03	0.56
55:DA:39:C:H2'	55:DA:40:C:C6	2.41	0.56
55:DA:877:U:O4	55:DA:899:A:C6	2.58	0.56
3:DD:70:TRP:CZ3	3:DD:146:GLU:OE2	2.57	0.56
6:DG:22:ARG:HH12	6:DG:175:LEU:HD21	1.70	0.56
6:DG:88:ILE:C	6:DG:88:ILE:HD13	2.25	0.56
9:DM:29:LYS:O	9:DM:33:LEU:HD13	2.05	0.56
9:DM:63:THR:HG22	9:DM:66:LYS:NZ	2.20	0.56
14:DQ:106:ARG:CZ	14:DQ:106:ARG:HB2	2.28	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:DR:16:ARG:NH2	15:DR:83:ILE:O	2.37	0.56
18:DS:1:MET:HG3	18:DS:2:GLU:N	2.20	0.56
19:DT:35:THR:O	19:DT:39:ILE:HG13	2.04	0.56
21:DV:67:LEU:HD22	21:DV:90:VAL:HG13	1.87	0.56
57:DY:72:ASP:OD1	57:DY:74:LEU:C	2.44	0.56
17:A2:3:ALA:HB1	17:A2:38:LEU:CD2	2.36	0.56
1:AA:2014:A:HO2'	27:A5:2:ALA:N	2.02	0.56
1:AA:2243:U:O2'	1:AA:2244:U:H5'	2.05	0.56
1:AA:2720:U:H3	1:AA:2873:A:H2	1.48	0.56
1:AA:2749:A:OP1	1:AA:2750:A:H5''	2.06	0.56
1:AA:527:C:P	1:AA:2779:U:H5	2.28	0.56
1:AA:874:G:H2'	1:AA:875:G:C8	2.40	0.56
3:AD:186:HIS:CD2	3:AD:188:GLU:HB2	2.39	0.56
1:AA:764:A:OP1	3:AD:208:LYS:HE3	2.04	0.56
8:AK:79:ILE:O	8:AK:81:VAL:HG23	2.04	0.56
9:AM:40:PRO:O	16:A1:64:ARG:HG3	2.04	0.56
4:AE:18:ASP:OD1	15:AR:82:LEU:HG	2.05	0.56
25:AX:12:PRO:HB2	25:AX:20:LYS:HG2	1.87	0.56
31:BA:632:A:H4'	31:BA:633:G:O5'	2.03	0.56
31:BA:634:C:O2'	31:BA:635:G:H5'	2.04	0.56
32:BE:217:ARG:HH11	32:BE:217:ARG:HG3	1.70	0.56
33:BF:6:HIS:HE1	44:BQ:50:LYS:HE2	1.70	0.56
34:BG:176:LEU:HD12	34:BG:177:ASP:N	2.16	0.56
34:BG:24:GLU:N	34:BG:27:TYR:HB2	2.21	0.56
34:BG:31:CYS:O	34:BG:32:ALA:HB3	2.05	0.56
35:BH:51:VAL:CB	35:BH:52:PRO:HD3	2.27	0.56
40:BM:75:ILE:HG13	40:BM:76:ASN:N	2.20	0.56
42:BO:27:LEU:HB2	42:BO:33:ARG:HD2	1.86	0.56
45:BR:10:LYS:HA	45:BR:10:LYS:HE3	1.87	0.56
50:BW:57:ARG:HG2	50:BW:102:GLY:O	2.05	0.56
33:CF:164:ARG:NH1	53:C1:55:U:O2	2.39	0.56
54:CA:117:G:O5'	54:CA:117:G:H8	1.89	0.56
54:CA:881:G:OP2	42:CO:12:ARG:NH2	2.38	0.56
54:CA:986:A:H2'	54:CA:987:G:C8	2.40	0.56
32:CE:233:SER:OG	32:CE:234:PRO:HD2	2.05	0.56
34:CG:11:LEU:O	34:CG:13:ARG:N	2.38	0.56
36:CI:42:GLU:C	36:CI:44:GLY:H	2.08	0.56
39:CL:26:VAL:HG13	39:CL:61:ALA:O	2.05	0.56
43:CP:116:THR:O	43:CP:117:VAL:CG1	2.54	0.56
50:CW:84:LEU:HD13	50:CW:85:MET:N	2.20	0.56
13:D0:117:VAL:CG2	13:D0:118:GLU:H	1.95	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
27:D5:51:TYR:HB2	27:D5:56:LYS:HB3	1.86	0.56
55:DA:1164:G:H2'	55:DA:1165:U:C6	2.40	0.56
55:DA:1458:C:H5''	55:DA:1459:G:C5'	2.36	0.56
55:DA:1349:A:N6	55:DA:1598:C:N4	2.53	0.56
55:DA:1794:U:H1'	55:DA:1900:A:N3	2.19	0.56
55:DA:2093:G:OP1	8:DK:23:PRO:HG2	2.04	0.56
55:DA:719:C:H2'	55:DA:720:C:H6	1.69	0.56
5:DF:206:ILE:HD12	5:DF:207:GLY:N	2.21	0.56
7:DH:152:ARG:HG3	7:DH:153:LYS:HE3	1.84	0.56
8:DK:76:THR:HG22	8:DK:77:LEU:N	2.19	0.56
58:DL:74:ALA:O	58:DL:75:SER:OG	2.18	0.56
58:DL:82:ALA:HA	58:DL:99:ILE:HG21	1.87	0.56
20:DU:94:LYS:HD2	20:DU:101:LYS:HZ3	1.69	0.56
25:DX:7:LYS:O	25:DX:54:VAL:HG23	2.05	0.56
57:DY:58:LEU:C	57:DY:62:ALA:HB3	2.26	0.56
57:DY:74:LEU:CD1	57:DY:74:LEU:C	2.69	0.56
30:A8:39:LYS:O	30:A8:40:GLU:HB2	2.06	0.56
1:AA:121:G:H4'	1:AA:149:A:H5'	1.86	0.56
1:AA:1678:G:N2	1:AA:1989:G:H22	2.03	0.56
1:AA:2807:G:C2'	1:AA:2808:U:H5''	2.34	0.56
1:AA:414:C:H2'	1:AA:415:A:C8	2.40	0.56
1:AA:961:C:H5''	1:AA:962:G:OP2	2.06	0.56
2:AB:40:U:C2	26:A4:1:MET:HE1	2.41	0.56
4:AE:108:SER:HB3	4:AE:165:VAL:CG2	2.35	0.56
5:AF:18:ARG:HG2	5:AF:19:GLU:H	1.71	0.56
5:AF:20:LEU:O	5:AF:21:ALA:O	2.23	0.56
6:AG:115:ARG:HH11	6:AG:115:ARG:HB3	1.69	0.56
8:AK:129:THR:HA	8:AK:137:PRO:HA	1.85	0.56
1:AA:2093:G:O5'	8:AK:24:GLY:HA3	2.06	0.56
18:AS:68:ARG:HH22	18:AS:112:GLY:HA2	1.70	0.56
21:AV:103:ARG:HG3	21:AV:105:VAL:HG12	1.88	0.56
31:BA:1036:G:H3'	31:BA:1037:C:C5	2.41	0.56
31:BA:1037:C:H2'	31:BA:1038:C:C6	2.40	0.56
31:BA:1129:C:N4	31:BA:1141:C:H41	2.03	0.56
31:BA:186(F):C:H2'	31:BA:187:C:O4'	2.05	0.56
34:BG:112:VAL:HG12	34:BG:116:GLN:CD	2.26	0.56
34:BG:30:LYS:HB2	34:BG:35:ARG:HH11	1.70	0.56
37:BJ:138:LYS:HG2	37:BJ:142:GLU:OE1	2.05	0.56
39:BL:63:ILE:HD12	39:BL:63:ILE:N	2.20	0.56
43:BP:96:LEU:HB3	43:BP:97:PRO:HD2	1.87	0.56
48:BU:18:ARG:O	48:BU:19:LYS:CB	2.41	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1095:U:H2'	54:CA:1096:C:H6	1.70	0.56
54:CA:977:A:C2'	54:CA:978:A:H5'	2.36	0.56
43:CP:126:LYS:HG3	52:CC:26:A:OP2	2.05	0.56
52:CD:61:C:H2'	52:CD:62:C:H6	1.70	0.56
32:CE:75:LYS:HA	32:CE:78:GLN:HG3	1.87	0.56
33:CF:134:ILE:CG2	33:CF:168:ALA:HB3	2.36	0.56
34:CG:11:LEU:O	34:CG:12:CYS:C	2.44	0.56
54:CA:878:G:H5'	38:CK:89:PRO:HG2	1.86	0.56
43:CP:23:TYR:CE1	43:CP:70:LEU:HD12	2.39	0.56
46:CS:51:VAL:HG11	46:CS:74:LEU:HD23	1.88	0.56
50:CW:72:LEU:HD23	50:CW:73:HIS:N	2.19	0.56
13:D0:52:ILE:O	13:D0:55:ALA:N	2.38	0.56
13:D0:8:ARG:HG3	13:D0:43:GLU:OE2	2.04	0.56
9:DM:41:ASP:C	16:D1:64:ARG:NH2	2.57	0.56
30:D8:41:ILE:HG13	30:D8:42:ARG:H	1.70	0.56
55:DA:1144:G:H2'	55:DA:1145:C:C6	2.40	0.56
55:DA:1177:A:C5'	55:DA:1178:C:H5''	2.36	0.56
55:DA:1496:A:C8	55:DA:1577:C:O2'	2.57	0.56
55:DA:1899:G:O2'	55:DA:1900:A:P	2.63	0.56
55:DA:2127:G:H2'	55:DA:2128:C:C4'	2.36	0.56
55:DA:2533:A:C2'	55:DA:2534:A:H5'	2.36	0.56
55:DA:2879:C:H4'	55:DA:2880:C:OP1	2.04	0.56
55:DA:620:G:H4'	55:DA:621:A:O5'	2.05	0.56
55:DA:654(I):C:O2	55:DA:654(I):C:H2'	2.05	0.56
55:DA:654(J):A:C2	55:DA:654(L):G:N7	2.73	0.56
55:DA:83:G:N2	55:DA:102:G:O2'	2.39	0.56
3:DD:124:PRO:HG2	3:DD:129:ASN:ND2	2.21	0.56
3:DD:129:ASN:O	3:DD:193:VAL:HG12	2.04	0.56
5:DF:9:ILE:HG23	5:DF:20:LEU:O	2.05	0.56
56:DI:8:ILE:O	56:DI:9:LYS:C	2.44	0.56
58:DL:10:LEU:O	58:DL:23:VAL:CG1	2.54	0.56
55:DA:1082:U:C3'	58:DL:117:THR:HG21	2.35	0.56
9:DM:17:ASP:O	9:DM:18:ALA:HB3	2.05	0.56
55:DA:196:A:OP2	11:DO:46:LYS:NZ	2.38	0.56
15:DR:23:ARG:HG2	15:DR:120:ARG:HH12	1.70	0.56
21:DV:196:VAL:O	21:DV:196:VAL:HG22	2.05	0.56
30:A8:16:ILE:HD13	30:A8:57:ARG:HD2	1.87	0.56
1:AA:1095:A:H2'	1:AA:1095:A:N3	2.20	0.56
1:AA:1203:G:H3'	1:AA:1204:A:H5''	1.87	0.56
1:AA:1387:C:C2	1:AA:1388:G:C8	2.93	0.56
1:AA:1937:A:O2'	1:AA:1938:A:P	2.62	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:AD:27:THR:HG22	3:AD:28:GLU:N	2.19	0.56
10:AN:102:VAL:HB	10:AN:106:LEU:CD1	2.34	0.56
12:AP:57:HIS:NE2	12:AP:116:GLU:HG3	2.20	0.56
15:AR:24:PRO:HA	15:AR:49:VAL:CG1	2.30	0.56
21:AV:105:VAL:HG23	21:AV:106:GLY:H	1.71	0.56
31:BA:1177:G:H2'	31:BA:1178:G:C2	2.40	0.56
31:BA:1200:C:H1'	31:BA:1204:A:H62	1.71	0.56
31:BA:1307:U:H2'	31:BA:1308:U:C6	2.40	0.56
31:BA:1380:U:H4'	31:BA:1381:U:O5'	2.03	0.56
31:BA:1393:U:O4'	31:BA:1502:A:H5''	2.06	0.56
31:BA:1527:C:H2'	31:BA:1528:U:O4'	2.05	0.56
32:BE:41:ILE:N	32:BE:41:ILE:HD12	2.21	0.56
32:BE:67:THR:HG22	32:BE:90:MET:SD	2.45	0.56
33:BF:113:ALA:HB3	33:BF:114:PRO:HD3	1.88	0.56
33:BF:178:LEU:C	33:BF:180:ALA:H	2.08	0.56
34:BG:12:CYS:CB	34:BG:21:LEU:CD2	2.79	0.56
36:BI:42:GLU:C	36:BI:44:GLY:H	2.09	0.56
31:BA:1382:C:H1'	37:BJ:79:ARG:NH1	2.21	0.56
40:BM:46:ARG:HG2	40:BM:64:GLU:HB3	1.88	0.56
43:BP:96:LEU:C	43:BP:110:ARG:HE	2.09	0.56
45:BR:78:TYR:O	45:BR:82:ILE:HG22	2.05	0.56
50:BW:50:GLU:HG3	50:BW:51:GLU:N	2.20	0.56
54:CA:1002:G:H2'	54:CA:1003:G:H5'	1.87	0.56
54:CA:1133:G:H2'	54:CA:1134:G:H8	1.71	0.56
54:CA:1322:C:O2	54:CA:1322:C:C2'	2.54	0.56
54:CA:191(C):G:H2'	54:CA:191(D):U:C6	2.41	0.56
54:CA:625:G:H2'	54:CA:626:U:C6	2.40	0.56
54:CA:69:G:C2	54:CA:73:G:C8	2.93	0.56
54:CA:753:A:O2'	54:CA:754:C:OP2	2.19	0.56
32:CE:45:GLN:O	32:CE:49:GLU:HG3	2.05	0.56
39:CL:79:LEU:O	39:CL:79:LEU:HD13	2.06	0.56
40:CM:38:ILE:HD11	40:CM:71:LEU:HB3	1.87	0.56
50:CW:69:GLY:O	50:CW:73:HIS:CD2	2.59	0.56
6:DG:67:LYS:HG3	26:D4:6:HIS:ND1	2.21	0.56
26:D4:9:LEU:H	26:D4:27:THR:HG22	1.71	0.56
28:D6:34:LEU:H	28:D6:34:LEU:CD1	2.17	0.56
55:DA:1026:U:O2	55:DA:1126:A:N1	2.39	0.56
55:DA:1130:U:O2'	55:DA:1131:G:O5'	2.22	0.56
55:DA:1167:U:H2'	55:DA:1168:G:C8	2.41	0.56
55:DA:1188:U:C2'	55:DA:1189:A:H5'	2.35	0.56
55:DA:1528:A:N1	55:DA:1543:A:C2	2.74	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2101:G:H2'	55:DA:2102:U:H6	1.70	0.56
55:DA:2346:A:H5'	55:DA:2383:G:O4'	2.06	0.56
55:DA:370:G:H4'	55:DA:371:A:OP2	2.04	0.56
55:DA:481:G:O2'	55:DA:482:A:P	2.61	0.56
55:DA:690:G:H2'	55:DA:691:C:C6	2.40	0.56
55:DA:848:G:O6	55:DA:929:G:H2'	2.05	0.56
55:DA:85:G:OP1	20:DU:30:VAL:HG21	2.06	0.56
4:DE:64:LYS:O	4:DE:70:ALA:HB2	2.04	0.56
7:DH:151:ILE:C	7:DH:152:ARG:O	2.42	0.56
7:DH:154:PRO:HD3	7:DH:161:GLY:CA	2.35	0.56
56:DI:7:ARG:HH21	56:DI:8:ILE:HG23	1.71	0.56
58:DL:124:ALA:O	58:DL:126:MET:N	2.38	0.56
58:DL:52:ILE:HG21	58:DL:75:SER:CA	2.34	0.56
9:DM:45:ASN:H	9:DM:45:ASN:ND2	2.04	0.56
10:DN:86:ILE:HG22	10:DN:94:ARG:HB2	1.87	0.56
11:DO:84:ASN:ND2	11:DO:116:GLY:HA3	2.21	0.56
19:DT:12:VAL:CG1	19:DT:27:THR:HG23	2.35	0.56
57:DY:88:ALA:O	57:DY:89:ALA:C	2.43	0.56
16:A1:101:ARG:O	16:A1:102:GLU:HG2	2.06	0.56
16:A1:102:GLU:N	16:A1:103:PRO:CD	2.69	0.56
17:A2:29:PRO:C	17:A2:61:VAL:HG11	2.26	0.56
26:A4:12:ALA:CB	26:A4:24:THR:HG21	2.35	0.56
30:A8:46:ARG:HH11	30:A8:46:ARG:CB	2.19	0.56
1:AA:1086:A:H5''	1:AA:1103:A:N1	2.21	0.56
1:AA:1342:A:C8	1:AA:1345:C:C5	2.94	0.56
1:AA:1996:C:H4'	1:AA:1997:G:O5'	2.05	0.56
1:AA:2065:C:H1'	1:AA:2449:U:O2	2.06	0.56
1:AA:320:A:H4'	1:AA:322:A:H8	1.70	0.56
1:AA:39:C:H2'	1:AA:40:C:C6	2.41	0.56
1:AA:460:A:H2'	1:AA:461:C:O4'	2.06	0.56
1:AA:518:G:H4'	18:AS:18:ARG:NH1	2.21	0.56
2:AB:12:C:H4'	2:AB:13:A:OP1	2.05	0.56
2:AB:15:A:H1'	2:AB:109:G:C4	2.40	0.56
4:AE:105:THR:HG21	4:AE:164:ARG:CZ	2.36	0.56
4:AE:199:ARG:HG2	4:AE:200:GLU:H	1.71	0.56
4:AE:78:LEU:N	4:AE:78:LEU:CD2	2.69	0.56
5:AF:5:ALA:HB1	5:AF:125:LEU:HD21	1.86	0.56
11:AO:111:ARG:HG3	11:AO:128:HIS:ND1	2.20	0.56
12:AP:6:ARG:O	12:AP:7:MET:HB2	2.06	0.56
12:AP:81:VAL:O	12:AP:82:ARG:HG2	2.05	0.56
14:AQ:106:ARG:O	14:AQ:107:GLU:CB	2.53	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:AR:80:SER:HB3	15:AR:83:ILE:CD1	2.36	0.56
1:AA:138:G:N2	19:AT:44:GLU:OE2	2.39	0.56
21:AV:130:PRO:HA	21:AV:133:ILE:HD11	1.88	0.56
21:AV:158:PRO:O	21:AV:161:VAL:HG22	2.05	0.56
24:AW:9:GLN:HA	24:AW:12:GLU:HB3	1.86	0.56
31:BA:1041:A:C3'	31:BA:1042:G:H5''	2.36	0.56
31:BA:1372:U:H2'	31:BA:1373:G:H5'	1.87	0.56
31:BA:474:G:OP1	46:BS:81:ARG:HB2	2.05	0.56
31:BA:532:A:O2'	31:BA:533:A:P	2.63	0.56
31:BA:789:U:H2'	31:BA:791:G:OP2	2.06	0.56
31:BA:85:U:H2'	31:BA:86:U:OP1	2.06	0.56
31:BA:986:A:H1'	49:BV:54:GLY:O	2.05	0.56
38:BK:12:ARG:NH1	38:BK:26:VAL:HA	2.20	0.56
39:BL:53:VAL:HG23	39:BL:55:ALA:HB3	1.88	0.56
45:BR:54:ARG:HG2	45:BR:58:MET:CE	2.35	0.56
46:BS:66:PRO:HG2	46:BS:71:ARG:HG3	1.87	0.56
54:CA:104:G:O2'	54:CA:105:G:H5'	2.05	0.56
54:CA:372:C:H42	54:CA:389:A:H62	1.54	0.56
54:CA:468:A:H4'	46:CS:80:PHE:O	2.06	0.56
54:CA:933:G:OP2	37:CJ:3:ARG:HB2	2.05	0.56
43:CP:121:LYS:O	43:CP:122:LYS:HB2	2.06	0.56
44:CQ:51:GLY:C	44:CQ:53:LEU:H	2.09	0.56
50:CW:63:ILE:CG2	50:CW:77:ALA:HB1	2.35	0.56
22:D3:68:GLU:CG	22:D3:80:HIS:HB2	2.34	0.56
26:D4:11:PRO:HB3	26:D4:25:TYR:CE1	2.41	0.56
28:D6:22:ALA:CB	28:D6:42:TRP:HZ2	2.13	0.56
11:DO:64:LYS:HB3	30:D8:25:MET:HG3	1.88	0.56
55:DA:1020:A:N1	55:DA:1141:U:C2'	2.68	0.56
55:DA:1324:G:C3'	55:DA:1325:G:H5'	2.36	0.56
55:DA:1937:A:O2'	55:DA:1938:A:P	2.63	0.56
55:DA:1999:C:H5''	55:DA:2723:C:O2'	2.05	0.56
55:DA:27:G:H1'	55:DA:513:A:H62	1.70	0.56
55:DA:702:G:H5'	55:DA:702:G:C8	2.41	0.56
55:DA:718:A:H3'	55:DA:719:C:C6	2.41	0.56
2:DB:74:U:C3'	2:DB:75:G:H5''	2.35	0.56
3:DD:92:ILE:HD13	3:DD:104:TYR:CE2	2.40	0.56
5:DF:153:SER:OG	5:DF:189:THR:HA	2.04	0.56
7:DH:94:TYR:CD2	7:DH:107:VAL:HG12	2.41	0.56
10:DN:86:ILE:N	10:DN:86:ILE:HD12	2.21	0.56
52:CB:57:G:C5'	21:DV:182:LYS:HZ3	2.19	0.56
21:DV:198:LYS:O	21:DV:199:LYS:CB	2.53	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:DV:75:ASN:O	21:DV:84:GLU:HG2	2.05	0.56
25:DX:5:LYS:HE2	25:DX:34:GLU:OE1	2.05	0.56
25:DX:7:LYS:HE2	25:DX:32:GLN:HA	1.87	0.56
57:DY:104:ILE:CG2	57:DY:105:PRO:N	2.67	0.56
57:DY:27:VAL:HG23	57:DY:110:GLY:HA2	1.76	0.56
23:DZ:23:LYS:HB3	23:DZ:29:GLY:HA3	1.88	0.56
23:DZ:78:LYS:HD2	23:DZ:78:LYS:O	2.06	0.56
16:A1:74:LEU:H	16:A1:74:LEU:HD12	1.69	0.56
26:A4:48:ARG:HG2	26:A4:51:ASP:HB3	1.87	0.56
1:AA:2331:G:H4'	22:A3:43:THR:N	2.21	0.56
1:AA:2722:G:H2'	1:AA:2723:C:C6	2.41	0.56
1:AA:2819:G:H2'	1:AA:2820:A:OP2	2.05	0.56
1:AA:654(C):G:C2	1:AA:654(D):G:C4	2.94	0.56
1:AA:664:C:H4'	1:AA:941:A:OP1	2.05	0.56
4:AE:176:ILE:HD12	4:AE:176:ILE:N	2.21	0.56
6:AG:13:GLU:O	6:AG:14:GLU:HB2	2.05	0.56
7:AH:143:GLN:HE22	7:AH:147:ASN:HD21	1.51	0.56
7:AH:10:PRO:CG	7:AH:50:VAL:HG13	2.35	0.56
7:AH:94:TYR:HE2	7:AH:153:LYS:HE2	1.71	0.56
8:AK:62:LYS:HD2	8:AK:62:LYS:C	2.25	0.56
11:AO:84:ASN:CG	11:AO:116:GLY:HA3	2.25	0.56
12:AP:80:GLU:HA	12:AP:80:GLU:OE2	2.06	0.56
12:AP:82:ARG:HH11	12:AP:82:ARG:CG	2.07	0.56
12:AP:75:THR:CG2	12:AP:88:GLY:HA3	2.29	0.56
15:AR:115:ARG:CD	15:AR:115:ARG:H	2.10	0.56
20:AU:35:TYR:CE1	20:AU:69:ALA:HB3	2.40	0.56
31:BA:255:G:H1'	47:BT:16:GLN:HE21	1.69	0.56
31:BA:160:A:H1'	31:BA:344:A:C5	2.40	0.56
31:BA:534:U:H5'	31:BA:535:A:OP2	2.06	0.56
31:BA:639:G:O2'	31:BA:640:A:H5'	2.06	0.56
31:BA:658:G:O2'	31:BA:659:U:H5'	2.05	0.56
31:BA:957:U:H2'	31:BA:959:A:OP2	2.05	0.56
52:BB:42:C:O2'	52:BB:43:C:H5'	2.06	0.56
52:BD:58:A:H1'	52:BD:60:U:C5	2.41	0.56
37:BJ:137:LYS:O	37:BJ:141:VAL:HG23	2.05	0.56
44:BQ:47:LEU:HB2	44:BQ:53:LEU:HD11	1.87	0.56
42:BO:8:ASN:ND2	47:BT:34:LYS:HE2	2.13	0.56
50:BW:84:LEU:O	50:BW:88:VAL:HG23	2.06	0.56
54:CA:1205:U:H5'	33:CF:190:ARG:NH2	2.20	0.56
54:CA:1348:U:H2'	54:CA:1349:A:H8	1.71	0.56
32:CE:70:PHE:CE2	32:CE:163:PHE:HD1	2.24	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1057:G:H5''	33:CF:154:SER:O	2.05	0.56
34:CG:156:GLU:HG2	34:CG:160:GLN:NE2	2.20	0.56
38:CK:29:SER:HB3	38:CK:32:LYS:HB2	1.88	0.56
39:CL:21:PRO:HA	39:CL:58:HIS:O	2.05	0.56
45:CR:56:LEU:O	45:CR:60:VAL:HG23	2.05	0.56
50:CW:94:ALA:O	50:CW:95:ALA:HB3	2.04	0.56
16:D1:34:LYS:HE2	16:D1:34:LYS:CA	2.28	0.56
11:DO:23:PRO:HB3	17:D2:80:GLN:HG3	1.87	0.56
55:DA:1523:U:H2'	55:DA:1524:G:O4'	2.06	0.56
55:DA:2506:U:O2'	55:DA:2507:C:H5''	2.05	0.56
55:DA:2739:U:O2'	55:DA:2740:A:H5'	2.06	0.56
55:DA:1:G:H2'	55:DA:2:G:C8	2.41	0.56
3:DD:149:PRO:O	3:DD:150:LYS:HB2	2.06	0.56
3:DD:223:GLY:O	3:DD:224:ALA:C	2.44	0.56
4:DE:46:ALA:O	4:DE:47:VAL:HG13	2.06	0.56
5:DF:133:ASN:O	5:DF:134:GLY:C	2.44	0.56
6:DG:96:ARG:HH11	6:DG:96:ARG:HG3	1.70	0.56
57:DY:132:ASP:OD1	56:DJ:7:ARG:NH1	2.38	0.56
8:DK:40:THR:O	8:DK:44:LEU:HB2	2.04	0.56
58:DL:106:GLU:CD	58:DL:109:LYS:HD2	2.26	0.56
55:DA:661:C:H1'	11:DO:12:ALA:HA	1.88	0.56
11:DO:88:LEU:HD23	11:DO:89:ALA:N	2.21	0.56
15:DR:19:LEU:HD22	15:DR:86:ILE:CG2	2.36	0.56
19:DT:50:LYS:N	19:DT:87:GLN:HE22	2.00	0.56
57:DY:26:LEU:O	57:DY:111:LEU:CD2	2.54	0.56
57:DY:29:TYR:CA	57:DY:81:VAL:CG1	2.45	0.56
26:A4:18:CYS:SG	26:A4:19:GLY:N	2.77	0.56
26:A4:63:TYR:CE2	49:BV:41:VAL:HG22	2.38	0.56
1:AA:749:C:C4	1:AA:1618:A:C2	2.94	0.56
1:AA:1772:G:N2	1:AA:1774:C:H5''	2.21	0.56
1:AA:1786:A:N1	1:AA:2606:C:H1'	2.20	0.56
1:AA:2211:G:H3'	1:AA:2211:G:OP2	2.05	0.56
1:AA:2433:A:H5''	1:AA:2434:A:OP1	2.05	0.56
1:AA:2522:U:O2'	1:AA:2647:U:H5''	2.06	0.56
1:AA:34:C:C2'	1:AA:35:G:OP2	2.54	0.56
1:AA:587:C:O2	11:AO:33:ARG:NH1	2.39	0.56
1:AA:774:A:H2	1:AA:787:U:HO2'	1.51	0.56
1:AA:90:U:O2'	1:AA:91:A:H8	1.78	0.56
2:AB:110:G:H2'	2:AB:111:U:O4'	2.06	0.56
2:AB:81:G:N2	2:AB:82:G:C5	2.74	0.56
2:AB:96:G:N2	2:AB:97:G:H1'	2.21	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:AD:58:HIS:HD2	3:AD:59:LYS:O	1.89	0.56
3:AD:62:TYR:HA	3:AD:87:ASN:HD21	1.71	0.56
5:AF:124:LEU:O	5:AF:125:LEU:C	2.43	0.56
6:AG:123:ASN:ND2	6:AG:123:ASN:N	2.54	0.56
8:AK:94:ALA:HB1	8:AK:112:LYS:H	1.70	0.56
21:AV:9:TYR:CE2	21:AV:61:LEU:HD13	2.41	0.56
31:BA:1229:A:H2'	31:BA:1230:C:H6	1.69	0.56
31:BA:1324:A:C4'	31:BA:1362:C:H4'	2.36	0.56
31:BA:881:G:P	42:BO:12:ARG:HH22	2.28	0.56
52:BB:51:U:H2'	52:BB:52:G:C8	2.40	0.56
38:BK:84:ARG:O	38:BK:135:CYS:HB2	2.06	0.56
54:CA:1025:U:O2'	54:CA:1026:G:H8	1.88	0.56
54:CA:1067:A:O2'	54:CA:1068:G:C8	2.41	0.56
54:CA:500:G:H2'	54:CA:501:C:C6	2.41	0.56
54:CA:50:A:O2'	54:CA:52:G:H8	1.86	0.56
54:CA:977:A:H2'	54:CA:978:A:H5'	1.87	0.56
52:CB:76:A:C8	55:DA:2507:C:O4'	2.58	0.56
32:CE:101:MET:HA	32:CE:108:ILE:HG13	1.88	0.56
32:CE:132:LYS:HA	32:CE:135:GLN:CD	2.26	0.56
32:CE:58:ILE:CG2	32:CE:221:LEU:HD12	2.36	0.56
34:CG:98:GLU:CD	34:CG:107:ARG:HE	2.07	0.56
42:CO:38:THR:CG2	42:CO:57:LYS:HB3	2.36	0.56
49:CV:83:HIS:C	49:CV:85:LYS:H	2.08	0.56
17:D2:55:ALA:HB2	17:D2:101:GLY:OXT	2.06	0.56
26:D4:68:ARG:CA	26:D4:68:ARG:NE	2.68	0.56
55:DA:1484:G:C3'	55:DA:1485:G:H5''	2.35	0.56
55:DA:961:C:H42	55:DA:2031:A:H1'	1.71	0.56
55:DA:211:A:H2'	55:DA:212:G:O4'	2.05	0.56
55:DA:2715:C:O2'	55:DA:2716:U:H5'	2.06	0.56
3:DD:131:LEU:CD1	3:DD:131:LEU:N	2.69	0.56
4:DE:52:LEU:N	4:DE:52:LEU:HD12	2.14	0.56
4:DE:92:THR:N	4:DE:95:ILE:HD11	2.20	0.56
5:DF:168:ARG:HG3	5:DF:175:THR:HG21	1.87	0.56
58:DL:11:GLN:CG	58:DL:41:PHE:HZ	2.10	0.56
9:DM:131:GLN:HE21	9:DM:131:GLN:H	1.53	0.56
11:DO:38:GLN:O	11:DO:39:LYS:C	2.40	0.56
20:DU:73:ARG:NH2	20:DU:82:PRO:HD3	2.20	0.56
16:A1:81:HIS:NE2	16:A1:85:LYS:HD2	2.21	0.56
17:A2:84:LYS:HB2	17:A2:84:LYS:NZ	2.21	0.56
22:A3:74:ARG:HG2	22:A3:75:LEU:HD23	1.87	0.56
26:A4:69:LYS:HD3	26:A4:69:LYS:C	2.26	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1372:U:C5'	1:AA:1372:U:C5	2.87	0.56
1:AA:2798:C:OP2	1:AA:2799:A:N7	2.39	0.56
1:AA:706:A:H2'	1:AA:707:G:O4'	2.06	0.56
1:AA:800:A:C4'	1:AA:801:G:O5'	2.52	0.56
1:AA:817:C:O2'	1:AA:839:U:OP1	2.23	0.56
2:AB:44:G:H1'	2:AB:47:C:N4	2.21	0.56
3:AD:3:VAL:HG13	3:AD:3:VAL:O	2.03	0.56
14:AQ:69:VAL:HG13	14:AQ:101:LEU:HD23	1.88	0.56
21:AV:30:ASN:OD1	21:AV:33:LEU:HB3	2.05	0.56
25:AX:18:ASP:O	25:AX:21:ALA:HB3	2.06	0.56
31:BA:1002:G:H2'	31:BA:1003:G:H8	1.68	0.56
31:BA:1217:C:O2'	31:BA:1218:C:H5'	2.04	0.56
31:BA:1234:C:C4'	31:BA:1364:U:O2'	2.54	0.56
31:BA:163:C:H2'	31:BA:164:U:O4'	2.06	0.56
34:BG:159:ARG:O	34:BG:163:GLU:HG3	2.06	0.56
34:BG:189:PRO:O	34:BG:190:ASP:HB2	2.06	0.56
31:BA:15:G:H1'	35:BH:19:MET:CE	2.35	0.56
37:BJ:155:ARG:HG2	37:BJ:156:TRP:H	1.69	0.56
37:BJ:76:ARG:HG3	37:BJ:89:MET:HB2	1.88	0.56
39:BL:66:ARG:HB3	39:BL:66:ARG:HH11	1.70	0.56
40:BM:6:ILE:CD1	40:BM:23:ILE:HG21	2.36	0.56
40:BM:89:ASP:HB3	40:BM:91:PRO:HD3	1.87	0.56
41:BN:55:LYS:C	41:BN:57:THR:H	2.09	0.56
45:BR:82:ILE:C	45:BR:82:ILE:HD13	2.26	0.56
50:BW:10:LEU:HD22	50:BW:10:LEU:C	2.25	0.56
54:CA:1098:C:O2'	54:CA:1099:G:H5'	2.05	0.56
54:CA:264:U:O2'	47:CT:64:PRO:HD2	2.05	0.56
54:CA:788:U:N3	54:CA:795:C:N4	2.53	0.56
34:CG:80:GLU:OE2	34:CG:84:LYS:HE2	2.06	0.56
35:CH:72:GLN:C	35:CH:74:GLY:H	2.08	0.56
54:CA:1298:C:H41	37:CJ:114:ARG:HB3	1.71	0.56
37:CJ:15:ASP:H	37:CJ:20:ASP:N	1.95	0.56
47:CT:40:LYS:HG2	47:CT:41:LYS:H	1.69	0.56
49:CV:23:ASN:OD1	49:CV:43:GLU:HB2	2.05	0.56
50:CW:93:GLU:OE1	50:CW:94:ALA:N	2.39	0.56
17:D2:38:LEU:O	17:D2:51:VAL:HA	2.05	0.56
22:D3:50:ASN:CB	22:D3:81:VAL:HB	2.36	0.56
28:D6:31:PRO:O	28:D6:32:ASN:HB3	2.06	0.56
55:DA:1019:U:H2'	55:DA:1020:A:H8	1.71	0.56
55:DA:1175:U:O2'	55:DA:1176:G:C4'	2.48	0.56
55:DA:1280:G:C2'	55:DA:1281:G:H5''	2.36	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1510:A:OP1	55:DA:1510:A:O3'	2.24	0.56
55:DA:1567:A:H3'	3:DD:86:PRO:HG3	1.88	0.56
55:DA:2341:G:H2'	55:DA:2342:C:C6	2.41	0.56
55:DA:2447:G:O2'	55:DA:2448:A:OP2	2.23	0.56
55:DA:302:C:O2'	55:DA:303:U:H5'	2.05	0.56
55:DA:482:A:H4'	20:DU:47:LYS:CD	2.36	0.56
55:DA:654(C):G:H2'	55:DA:654(D):G:C1'	2.36	0.56
7:DH:136:ILE:HD12	7:DH:136:ILE:N	2.21	0.56
58:DL:110:GLN:O	58:DL:111:LYS:HE2	2.06	0.56
58:DL:111:LYS:CA	58:DL:113:PRO:HD2	2.35	0.56
58:DL:112:MET:CG	58:DL:118:THR:O	2.48	0.56
10:DN:77:ILE:HD13	10:DN:78:ARG:N	2.21	0.56
23:DZ:44:PRO:HG2	23:DZ:46:LEU:HD13	1.88	0.56
17:A2:70:ILE:O	17:A2:71:LEU:HB3	2.06	0.56
17:A2:79:VAL:HG23	17:A2:80:GLN:H	1.71	0.56
22:A3:81:VAL:O	22:A3:83:PRO:HD3	2.04	0.56
30:A8:47:LYS:O	30:A8:48:PHE:HB3	2.06	0.56
30:A8:60:LEU:O	30:A8:61:LEU:HD12	2.05	0.56
1:AA:1316:U:H2'	1:AA:1317:A:C8	2.41	0.56
1:AA:2130:U:H4'	1:AA:2134:A:H5'	1.87	0.56
1:AA:2159:G:H2'	1:AA:2160:G:C8	2.41	0.56
1:AA:2311:A:H3'	1:AA:2312:U:H5	1.67	0.56
1:AA:2475:C:H5'	1:AA:2476:A:OP2	2.06	0.56
1:AA:2571:C:H5'	1:AA:2572:A:H5'	1.87	0.56
1:AA:2602:A:H4'	1:AA:2603:G:O5'	2.05	0.56
2:AB:111:U:H2'	2:AB:112:G:H8	1.67	0.56
2:AB:11:C:H3'	2:AB:12:C:C6	2.41	0.56
3:AD:27:THR:O	3:AD:28:GLU:CB	2.53	0.56
4:AE:107:THR:O	4:AE:190:GLY:HA2	2.06	0.56
5:AF:61:GLY:O	5:AF:77:ASP:HB2	2.06	0.56
6:AG:88:ILE:C	6:AG:88:ILE:HD13	2.27	0.56
9:AM:128:HIS:HE2	9:AM:134:ARG:HD2	1.69	0.56
14:AQ:27:SER:HA	14:AQ:88:ASP:CB	2.35	0.56
21:AV:52:SER:O	21:AV:54:HIS:N	2.39	0.56
24:AW:69:ARG:HH11	24:AW:69:ARG:HG2	1.70	0.56
31:BA:495:A:H5'	31:BA:496:A:OP1	2.06	0.56
31:BA:64:G:H5'	31:BA:65:U:OP1	2.04	0.56
31:BA:939:G:C6	31:BA:940:C:N4	2.74	0.56
33:BF:112:SER:HB3	33:BF:115:LEU:HD12	1.86	0.56
38:BK:31:PHE:O	38:BK:35:ILE:HG13	2.05	0.56
47:BT:59:ILE:CG2	47:BT:71:PHE:HB3	2.36	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1070:U:H2'	54:CA:1071:C:C6	2.40	0.56
54:CA:67:C:O2'	54:CA:171:A:H1'	2.05	0.56
54:CA:86:U:O3'	54:CA:87:A:O4'	2.24	0.56
32:CE:4:GLU:HG2	32:CE:5:ILE:HD13	1.87	0.56
36:CI:27:GLN:HA	36:CI:27:GLN:NE2	2.17	0.56
38:CK:100:ILE:HB	38:CK:125:ARG:HH12	1.70	0.56
38:CK:13:ILE:O	38:CK:17:THR:HG23	2.06	0.56
46:CS:48:TRP:O	46:CS:49:LEU:HB2	2.06	0.56
13:D0:18:LEU:HD22	13:D0:18:LEU:O	2.05	0.56
16:D1:110:VAL:O	16:D1:114:LYS:HG2	2.05	0.56
30:D8:30:ARG:O	30:D8:31:HIS:CB	2.54	0.56
55:DA:1009:A:OP2	9:DM:37:LYS:NZ	2.39	0.56
55:DA:1934:C:O2'	55:DA:1935:G:H5'	2.06	0.56
55:DA:2119:A:H8	55:DA:2119:A:O5'	1.89	0.56
55:DA:2135:A:C2'	55:DA:2136:C:OP1	2.54	0.56
55:DA:90:U:H1'	55:DA:91:A:N7	2.21	0.56
55:DA:99:U:C6	55:DA:102:G:C2	2.94	0.56
2:DB:28:C:H2'	2:DB:29:A:O4'	2.05	0.56
4:DE:159:HIS:HE1	4:DE:162:ALA:HB3	1.71	0.56
58:DL:137:GLU:HA	58:DL:137:GLU:OE2	2.06	0.56
58:DL:4:VAL:HG12	58:DL:4:VAL:O	2.05	0.56
58:DL:78:ILE:C	58:DL:82:ALA:HB3	2.26	0.56
11:DO:105:LEU:N	11:DO:105:LEU:HD12	2.20	0.56
14:DQ:3:ARG:HG2	14:DQ:4:LEU:H	1.70	0.56
20:DU:43:ASN:HB3	20:DU:64:GLU:HA	1.88	0.56
57:DY:71:LEU:HB3	57:DY:112:LEU:O	1.97	0.56
13:A0:74:LYS:HE2	13:A0:77:ARG:NH2	2.19	0.56
26:A4:24:THR:HG22	26:A4:25:TYR:N	2.21	0.56
26:A4:39:CYS:O	26:A4:40:HIS:CB	2.53	0.56
26:A4:67:TYR:O	26:A4:69:LYS:N	2.39	0.56
28:A6:41:PRO:HD2	28:A6:46:HIS:CA	2.36	0.56
1:AA:1098:A:H3'	1:AA:1099:G:H5'	1.86	0.56
1:AA:235:U:H2'	1:AA:236:C:C6	2.41	0.56
1:AA:616:A:N3	1:AA:616:A:H2'	2.20	0.56
1:AA:654(G):C:H2'	1:AA:654(H):G:N7	2.20	0.56
1:AA:85:G:N3	1:AA:103:A:C2	2.74	0.56
4:AE:6:GLY:HA3	4:AE:28:ALA:HA	1.88	0.56
4:AE:55:ASN:HD21	4:AE:75:VAL:CA	2.19	0.56
5:AF:9:ILE:HG12	5:AF:15:SER:HA	1.88	0.56
11:AO:85:LEU:H	11:AO:85:LEU:CD2	2.19	0.56
12:AP:22:LYS:O	12:AP:24:GLY:N	2.39	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:AR:97:ALA:HB1	15:AR:98:LYS:HZ3	1.71	0.56
18:AS:69:LEU:HD12	18:AS:69:LEU:O	2.06	0.56
20:AU:72:VAL:CG2	20:AU:73:ARG:H	2.07	0.56
25:AX:6:VAL:HG13	25:AX:56:VAL:CG2	2.35	0.56
31:BA:1306:A:H1'	31:BA:1332:A:C2	2.41	0.56
31:BA:186(D):C:H2'	31:BA:186(E):C:H6	1.69	0.56
31:BA:194:C:H2'	31:BA:195:A:H5''	1.87	0.56
31:BA:539:A:H2'	31:BA:540:G:H8	1.71	0.56
31:BA:692:U:H2'	31:BA:694:A:OP2	2.06	0.56
33:BF:150:LYS:HB3	33:BF:201:TYR:HB2	1.88	0.56
34:BG:21:LEU:HD12	34:BG:26:CYS:O	2.05	0.56
39:BL:65:VAL:O	39:BL:66:ARG:HB2	2.05	0.56
31:BA:973:G:N3	40:BM:55:LYS:HE2	2.20	0.56
40:BM:7:LYS:CG	40:BM:71:LEU:HD13	2.36	0.56
44:BQ:4:LYS:O	44:BQ:7:ILE:HG12	2.06	0.56
54:CA:1158:C:O2	54:CA:1158:C:C2'	2.54	0.56
54:CA:1238:A:N6	54:CA:1299:A:N6	2.54	0.56
54:CA:191(C):G:H2'	54:CA:191(D):U:O4'	2.06	0.56
54:CA:481:G:H1'	54:CA:482:A:N7	2.20	0.56
54:CA:807:A:H2'	54:CA:808:C:H6	1.68	0.56
54:CA:84:U:C5'	54:CA:84:U:C6	2.87	0.56
54:CA:960:U:O2	54:CA:960:U:C2'	2.53	0.56
35:CH:72:GLN:NE2	35:CH:144:THR:HG22	2.21	0.56
54:CA:973:G:N3	40:CM:55:LYS:HE2	2.21	0.56
46:CS:7:ALA:O	46:CS:9:PHE:CD2	2.58	0.56
17:D2:66:ARG:CB	17:D2:66:ARG:HH11	2.19	0.56
55:DA:1084:A:C6	55:DA:1085:A:N6	2.73	0.56
55:DA:139:G:N3	55:DA:141:A:N1	2.54	0.56
55:DA:1789:A:OP1	3:DD:221:VAL:HA	2.06	0.56
55:DA:2082:A:H2'	55:DA:2083:G:O4'	2.06	0.56
55:DA:2346:A:O2'	55:DA:2347:C:P	2.64	0.56
55:DA:2068:U:N3	55:DA:2430:A:C2	2.63	0.56
55:DA:2791:C:C5	55:DA:2893:G:C5	2.94	0.56
55:DA:259:G:N2	55:DA:621:A:H8	1.96	0.56
55:DA:889:C:O2	55:DA:889:C:C2'	2.53	0.56
58:DL:128:ALA:O	58:DL:132:ARG:HB3	2.06	0.56
58:DL:21:PRO:HB2	58:DL:22:PRO:HD2	1.88	0.56
58:DL:51:ALA:HB1	58:DL:79:ARG:HE	1.71	0.56
58:DL:53:VAL:HA	58:DL:72:PRO:HB2	1.88	0.56
9:DM:18:ALA:HB2	9:DM:54:VAL:HG13	1.87	0.56
10:DN:1:MET:HA	10:DN:33:ALA:O	2.06	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:389:G:H22	11:DO:72:PRO:HD3	1.70	0.56
18:DS:59:VAL:HG12	18:DS:60:ASN:HD22	1.71	0.56
21:DV:105:VAL:HG12	21:DV:140:ASP:CB	2.33	0.56
17:A2:22:VAL:CG2	17:A2:23:GLU:N	2.68	0.55
17:A2:41:GLY:HA3	17:A2:46:VAL:CG1	2.36	0.55
17:A2:58:VAL:HB	17:A2:98:GLU:HG3	1.87	0.55
30:A8:47:LYS:C	30:A8:48:PHE:HD1	2.09	0.55
1:AA:1044:G:O3'	1:AA:1045:A:H4'	2.06	0.55
1:AA:1400:G:O2'	1:AA:1401:G:H5'	2.06	0.55
1:AA:2102:U:H2'	1:AA:2103:C:H6	1.72	0.55
1:AA:2271:G:H5''	22:A3:20:ARG:HE	1.70	0.55
1:AA:2741:A:H61	1:AA:2763:G:C2'	2.19	0.55
1:AA:384:U:C3'	1:AA:385:C:H5'	2.34	0.55
1:AA:559:G:O2'	1:AA:560:C:H5'	2.06	0.55
1:AA:654(O):G:H2'	1:AA:654(P):G:C8	2.41	0.55
3:AD:27:THR:HG21	3:AD:83:GLU:OE2	2.05	0.55
4:AE:4:ILE:C	4:AE:5:LEU:HD23	2.26	0.55
6:AG:97:ASP:N	6:AG:100:TRP:HD1	2.02	0.55
7:AH:136:ILE:H	7:AH:136:ILE:HD12	1.69	0.55
8:AK:1:MET:HB2	8:AK:21:VAL:O	2.06	0.55
11:AO:105:LEU:O	11:AO:106:LEU:HB2	2.05	0.55
11:AO:128:HIS:O	11:AO:147:LEU:HB3	2.06	0.55
14:AQ:110:LEU:HD22	14:AQ:111:GLU:N	2.20	0.55
18:AS:70:TYR:H	18:AS:70:TYR:HD2	1.54	0.55
20:AU:20:TYR:C	20:AU:22:GLY:N	2.45	0.55
21:AV:183:LEU:O	21:AV:184:ALA:HB2	2.05	0.55
23:AZ:79:GLY:C	23:AZ:80:LEU:HD13	2.26	0.55
31:BA:1091:U:H2'	31:BA:1093:A:OP2	2.06	0.55
31:BA:1240:U:OP2	37:BJ:116:ALA:HB2	2.07	0.55
31:BA:1297:C:O2'	31:BA:1298:C:P	2.64	0.55
31:BA:1321:C:N4	31:BA:1322:C:N4	2.53	0.55
31:BA:1327:C:H2'	31:BA:1328:C:H6	1.72	0.55
31:BA:1341:U:O2'	31:BA:1342:C:H5'	2.06	0.55
31:BA:1381:U:H3	37:BJ:79:ARG:NH2	2.04	0.55
31:BA:80:G:N2	31:BA:90:C:H1'	2.21	0.55
31:BA:882:C:O2'	31:BA:883:C:H5'	2.06	0.55
31:BA:973:G:C1'	40:BM:55:LYS:HG2	2.36	0.55
52:BB:61:C:O2'	52:BB:62:C:H5'	2.06	0.55
31:BA:438:G:H4'	34:BG:123:HIS:ND1	2.21	0.55
34:BG:59:ARG:HH21	34:BG:66:ARG:NH1	2.04	0.55
34:BG:9:CYS:HA	34:BG:12:CYS:CB	2.32	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:BL:96:LEU:HD12	39:BL:101:PHE:HB2	1.87	0.55
39:BL:95:LYS:HZ2	39:BL:96:LEU:HD13	1.71	0.55
46:BS:21:VAL:CG1	46:BS:34:GLU:HB3	2.35	0.55
49:BV:23:ASN:O	49:BV:25:LYS:N	2.39	0.55
50:BW:72:LEU:HD11	50:BW:80:ARG:HD3	1.89	0.55
54:CA:1260:C:OP1	54:CA:1284:C:H4'	2.06	0.55
32:CE:68:ILE:N	32:CE:68:ILE:HD12	2.21	0.55
34:CG:150:GLU:N	34:CG:150:GLU:OE1	2.37	0.55
36:CI:19:LEU:HD11	36:CI:59:TYR:CZ	2.41	0.55
36:CI:30:LEU:C	36:CI:35:ALA:HB3	2.26	0.55
54:CA:1251:A:H4'	39:CL:12:GLU:OE1	2.06	0.55
54:CA:1189:C:P	40:CM:51:ARG:HH22	2.29	0.55
41:CN:126:ARG:HG2	41:CN:126:ARG:HH11	1.72	0.55
47:CT:4:LYS:CE	47:CT:6:LEU:HD21	2.31	0.55
17:D2:35:LEU:N	17:D2:35:LEU:HD22	2.12	0.55
26:D4:56:VAL:O	26:D4:60:GLN:N	2.39	0.55
29:D7:10:ARG:O	29:D7:14:LYS:HB2	2.05	0.55
55:DA:1151:G:H4'	16:D1:81:HIS:CD2	2.40	0.55
55:DA:1385:G:C4'	55:DA:1386:C:OP1	2.54	0.55
55:DA:1933:G:H2'	55:DA:1934:C:C5'	2.35	0.55
55:DA:2028:U:H2'	55:DA:2029:G:O4'	2.05	0.55
55:DA:2112:G:O2'	55:DA:2113:U:H5''	2.06	0.55
55:DA:2134:A:OP2	55:DA:2157:G:N2	2.38	0.55
55:DA:2141:G:O2'	55:DA:2142:C:H5'	2.05	0.55
55:DA:2344:U:H4'	55:DA:2345:G:OP1	2.04	0.55
55:DA:2591:C:H2'	55:DA:2592:G:H8	1.69	0.55
55:DA:2670:A:O2'	55:DA:2671:A:H5'	2.06	0.55
55:DA:654(C):G:H2'	55:DA:654(D):G:C8	2.41	0.55
55:DA:856:C:H4'	55:DA:857:C:OP1	2.06	0.55
55:DA:897:C:H5	55:DA:897:C:OP2	1.87	0.55
55:DA:999:U:H5''	55:DA:1154:G:O6	2.06	0.55
3:DD:263:ARG:HB2	3:DD:263:ARG:NH1	2.21	0.55
4:DE:167:VAL:HG21	4:DE:187:ALA:O	2.06	0.55
4:DE:51:PHE:HD1	4:DE:52:LEU:CG	2.19	0.55
6:DG:41:GLN:HE21	6:DG:60:LEU:CD1	2.19	0.55
7:DH:17:VAL:HG12	7:DH:17:VAL:O	2.06	0.55
8:DK:11:ASN:O	8:DK:12:LEU:CB	2.51	0.55
8:DK:76:THR:HG22	8:DK:77:LEU:HD12	1.87	0.55
8:DK:76:THR:HG22	8:DK:77:LEU:H	1.71	0.55
8:DK:77:LEU:CD1	8:DK:78:THR:N	2.69	0.55
58:DL:112:MET:HE3	58:DL:118:THR:C	2.26	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:DL:93:ARG:CA	58:DL:136:VAL:HG12	2.36	0.55
9:DM:75:TYR:HA	9:DM:81:GLY:O	2.06	0.55
10:DN:104:ARG:HD3	15:DR:36:GLU:CD	2.27	0.55
55:DA:910:A:H62	12:DP:12:GLN:HA	1.71	0.55
12:DP:23:GLY:HA3	12:DP:101:ARG:HD2	1.88	0.55
20:DU:33:LYS:HG3	20:DU:34:LYS:H	1.71	0.55
57:DY:138:LEU:C	57:DY:138:LEU:CD1	2.72	0.55
57:DY:38:HIS:O	57:DY:96:PHE:CE2	2.59	0.55
57:DY:38:HIS:CE1	57:DY:40:LEU:HD12	2.41	0.55
57:DY:91:LYS:HD3	57:DY:94:VAL:HG12	1.87	0.55
16:A1:91:ASP:OD2	16:A1:96:ALA:HB2	2.07	0.55
17:A2:48:GLY:HA3	17:A2:52:VAL:CG2	2.31	0.55
17:A2:70:ILE:HB	17:A2:86:GLY:O	2.05	0.55
1:AA:1098:A:C3'	1:AA:1099:G:C5'	2.85	0.55
1:AA:2158:A:H4'	1:AA:2159:G:O5'	2.07	0.55
1:AA:2298:A:N6	1:AA:2318:G:C8	2.74	0.55
1:AA:643:A:N1	1:AA:2369:A:O2'	2.39	0.55
1:AA:897:C:N3	1:AA:898:C:C5	2.74	0.55
2:AB:43:C:H1'	6:AG:93:THR:O	2.06	0.55
2:AB:94:C:O2'	2:AB:95:U:H5'	2.06	0.55
1:AA:1798:U:H5'	3:AD:259:THR:OG1	2.06	0.55
4:AE:37:ARG:HA	4:AE:42:ASP:OD2	2.06	0.55
5:AF:164:ARG:HG3	5:AF:175:THR:HG21	1.88	0.55
7:AH:105:LEU:HD13	7:AH:105:LEU:N	2.21	0.55
7:AH:42:ARG:HG2	7:AH:42:ARG:HH11	1.70	0.55
8:AK:47:LEU:O	8:AK:51:ILE:HG13	2.06	0.55
1:AA:195:A:OP1	11:AO:46:LYS:HE2	2.06	0.55
11:AO:55:ARG:O	11:AO:56:SER:C	2.44	0.55
15:AR:62:THR:HG22	15:AR:75:ILE:HG12	1.88	0.55
21:AV:63:ASP:O	21:AV:65:GLN:N	2.39	0.55
31:BA:135:C:H2'	31:BA:136:C:H5'	1.88	0.55
31:BA:157:G:H2'	31:BA:158:G:H8	1.71	0.55
31:BA:5:U:C2'	31:BA:6:G:OP2	2.53	0.55
52:BB:58:A:H4'	52:BB:59:U:OP1	2.07	0.55
31:BA:1190:G:H5'	33:BF:176:HIS:NE2	2.21	0.55
40:BM:78:ASN:C	40:BM:80:LYS:N	2.59	0.55
31:BA:951:G:OP2	43:BP:102:ARG:NH2	2.37	0.55
45:BR:17:ARG:NH1	45:BR:77:ARG:NH1	2.54	0.55
54:CA:130:A:H1'	54:CA:264:U:C4'	2.36	0.55
54:CA:1358:U:H5''	44:CQ:33:VAL:O	2.06	0.55
54:CA:673:G:H2'	54:CA:674:G:C8	2.41	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:85:U:O2'	54:CA:86:U:P	2.64	0.55
54:CA:963:G:N3	40:CM:55:LYS:NZ	2.54	0.55
55:DA:857:C:H5'	22:D3:77:ARG:NH2	2.21	0.55
27:D5:4:HIS:HB3	27:D5:5:PRO:HD2	1.84	0.55
27:D5:56:LYS:O	27:D5:57:VAL:HG12	2.06	0.55
30:D8:56:GLU:O	30:D8:59:LYS:N	2.37	0.55
55:DA:1286:A:C2'	55:DA:1288:U:OP2	2.54	0.55
55:DA:1558:A:O2'	55:DA:1559:G:OP2	2.23	0.55
55:DA:1729:A:C2'	55:DA:1730:U:H5''	2.37	0.55
55:DA:2033:A:H4'	55:DA:2034:U:OP1	2.07	0.55
55:DA:2475:C:O2	55:DA:2475:C:C2'	2.52	0.55
55:DA:2754:U:H2'	55:DA:2756:U:OP1	2.06	0.55
55:DA:2815:C:H2'	55:DA:2816:C:C6	2.41	0.55
55:DA:881:G:H3'	55:DA:882:G:O4'	2.05	0.55
3:DD:124:PRO:HB2	3:DD:126:GLN:HE22	1.70	0.55
5:DF:175:THR:O	5:DF:176:LEU:HB2	2.05	0.55
7:DH:118:PRO:HG2	7:DH:121:ILE:HD12	1.88	0.55
7:DH:153:LYS:HG3	7:DH:161:GLY:HA3	1.88	0.55
56:DI:29:GLU:O	56:DI:30:ALA:O	2.25	0.55
58:DL:100:THR:O	58:DL:102:GLU:N	2.39	0.55
55:DA:1079:C:H1'	58:DL:129:GLY:C	2.27	0.55
58:DL:93:ARG:HA	58:DL:136:VAL:HG12	1.87	0.55
55:DA:2563:U:O2'	10:DN:28:SER:HB3	2.07	0.55
11:DO:15:ARG:O	11:DO:17:LYS:N	2.38	0.55
12:DP:24:GLY:HA2	12:DP:67:ARG:NH2	2.21	0.55
55:DA:955:C:OP1	12:DP:85:LYS:HE2	2.07	0.55
18:DS:29:LEU:HD13	18:DS:69:LEU:CD1	2.36	0.55
57:DY:35:LYS:HA	57:DY:35:LYS:NZ	2.22	0.55
57:DY:43:ALA:O	57:DY:44:LEU:CB	2.54	0.55
57:DY:50:ARG:HB2	57:DY:50:ARG:CZ	2.36	0.55
57:DY:98:LYS:HG3	57:DY:99:SER:N	2.21	0.55
17:A2:46:VAL:HG22	17:A2:46:VAL:O	2.06	0.55
1:AA:1465:G:H5'	1:AA:1528:A:H1'	1.89	0.55
1:AA:1652:A:OP1	13:A0:8:ARG:NH1	2.39	0.55
1:AA:1929:G:N3	1:AA:1929:G:C2'	2.69	0.55
1:AA:2449:U:O2'	1:AA:2450:A:H8	1.87	0.55
1:AA:2611:U:H6	1:AA:2611:U:H5'	1.72	0.55
1:AA:500:G:N2	1:AA:502:A:H3'	2.21	0.55
4:AE:39:PRO:HA	4:AE:43:GLY:H	1.70	0.55
5:AF:136:THR:HG23	5:AF:170:LEU:HD11	1.89	0.55
8:AK:142:VAL:O	8:AK:143:SER:HB2	2.06	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:AM:49:GLY:O	9:AM:119:ARG:NH1	2.39	0.55
9:AM:94:HIS:HB2	9:AM:96:GLU:OE2	2.05	0.55
11:AO:57:THR:CG2	11:AO:60:MET:HB2	2.36	0.55
12:AP:130:LYS:HZ3	21:AV:81:ARG:HG2	1.71	0.55
14:AQ:100:ALA:O	14:AQ:103:GLU:HG2	2.06	0.55
14:AQ:83:LYS:O	14:AQ:109:GLY:HA3	2.06	0.55
15:AR:88:ILE:HD12	15:AR:88:ILE:C	2.26	0.55
20:AU:30:VAL:O	20:AU:36:ALA:O	2.23	0.55
21:AV:103:ARG:HA	21:AV:103:ARG:NE	2.21	0.55
23:AZ:90:ILE:CG2	23:AZ:91:LYS:N	2.68	0.55
31:BA:1054:C:O2'	31:BA:1055:A:H5''	2.07	0.55
31:BA:1246:C:H2'	31:BA:1247:U:H6	1.72	0.55
31:BA:405:U:H3'	31:BA:406:G:H5'	1.89	0.55
31:BA:67:C:O2'	31:BA:171:A:H1'	2.05	0.55
31:BA:913:A:H4'	31:BA:914:A:O5'	2.06	0.55
32:BE:22:LYS:HA	32:BE:22:LYS:NZ	2.21	0.55
31:BA:1125:U:O4	40:BM:5:ARG:NH1	2.39	0.55
41:BN:69:ALA:O	41:BN:73:MET:HG2	2.06	0.55
47:BT:22:LEU:HD11	47:BT:39:SER:HB2	1.88	0.55
49:BV:6:LYS:HZ2	49:BV:10:PHE:HZ	1.53	0.55
49:BV:76:PRO:CB	49:BV:78:ARG:HD3	2.36	0.55
50:BW:8:ARG:NH1	50:BW:8:ARG:HG3	2.19	0.55
54:CA:1401:G:C2	54:CA:1402:C:H1'	2.42	0.55
54:CA:35:G:H2'	54:CA:36:C:C6	2.41	0.55
54:CA:8:A:O2'	35:CH:103:GLY:N	2.38	0.55
52:CB:7:A:H4'	52:CB:8:U:OP1	2.04	0.55
33:CF:134:ILE:HG22	33:CF:168:ALA:HB3	1.88	0.55
33:CF:34:LEU:CD2	33:CF:38:ARG:HD2	2.37	0.55
34:CG:129:ASN:HA	34:CG:145:GLU:CG	2.35	0.55
34:CG:29:PRO:O	34:CG:30:LYS:HB3	2.06	0.55
37:CJ:12:LEU:HD22	37:CJ:12:LEU:N	2.21	0.55
54:CA:177:C:OP2	50:CW:65:LYS:HE2	2.06	0.55
13:D0:74:LYS:O	13:D0:75:LEU:CB	2.54	0.55
16:D1:112:ARG:HH11	16:D1:112:ARG:HG2	1.71	0.55
55:DA:1162:G:O3'	17:D2:24:LYS:NZ	2.40	0.55
6:DG:145:THR:HG22	26:D4:28:LYS:HZ1	1.71	0.55
55:DA:1057:A:C6	55:DA:1086:A:C2	2.94	0.55
55:DA:1332:G:H5''	55:DA:1333:C:OP2	2.07	0.55
55:DA:1359:A:N7	55:DA:1359:A:OP2	2.38	0.55
55:DA:2054:A:H5''	55:DA:2055:C:O5'	2.05	0.55
55:DA:2700:C:O2'	55:DA:2701:C:H5'	2.05	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:DD:134:ARG:HG3	3:DD:187:GLY:C	2.26	0.55
3:DD:223:GLY:O	3:DD:225:ALA:N	2.38	0.55
55:DA:2304:G:N2	6:DG:156:ASP:OD2	2.31	0.55
6:DG:180:PHE:C	6:DG:182:LYS:N	2.60	0.55
55:DA:1111:A:C5'	7:DH:3:ARG:NH1	2.69	0.55
56:DI:16:THR:CG2	56:DI:17:VAL:N	2.56	0.55
58:DL:144:VAL:HG13	58:DL:145:LYS:N	2.13	0.55
58:DL:7:VAL:CG1	58:DL:58:THR:C	2.67	0.55
55:DA:637:A:P	11:DO:116:GLY:HA2	2.46	0.55
20:DU:81:LYS:HB3	20:DU:97:ARG:CD	2.36	0.55
21:DV:20:ARG:C	21:DV:22:GLY:H	2.09	0.55
57:DY:9:LEU:CD1	57:DY:10:LEU:N	2.51	0.55
57:DY:9:LEU:CD2	57:DY:9:LEU:C	2.71	0.55
23:DZ:82:LEU:HD22	23:DZ:82:LEU:N	2.22	0.55
17:A2:2:PHE:CD2	17:A2:42:GLY:HA2	2.42	0.55
26:A4:12:ALA:C	26:A4:24:THR:OG1	2.44	0.55
1:AA:1116:C:H2'	1:AA:1117:G:C8	2.42	0.55
1:AA:1340:U:HO2'	1:AA:1341:U:P	2.29	0.55
1:AA:1469:A:H2'	1:AA:1470:G:C8	2.41	0.55
1:AA:1779:U:OP2	1:AA:1784:A:N6	2.38	0.55
1:AA:1798:U:OP1	3:AD:260:ARG:HB2	2.07	0.55
1:AA:191:A:O2'	1:AA:192:C:H5'	2.06	0.55
1:AA:1945:G:C4	1:AA:1946:U:C5	2.95	0.55
1:AA:2425:A:H5''	1:AA:2426:A:O5'	2.07	0.55
1:AA:2531:A:H3'	1:AA:2532:G:H8	1.71	0.55
1:AA:2776:A:O2'	1:AA:2781:A:H4'	2.06	0.55
1:AA:2851:A:O2'	1:AA:2852:G:H5'	2.05	0.55
1:AA:2791:C:C5	1:AA:2893:G:C5	2.94	0.55
1:AA:372:G:O2'	1:AA:400:G:N1	2.40	0.55
3:AD:35:LYS:HG2	3:AD:64:ILE:CA	2.36	0.55
3:AD:27:THR:HG21	3:AD:83:GLU:CG	2.35	0.55
5:AF:17:ARG:O	5:AF:17:ARG:HD3	2.06	0.55
5:AF:4:VAL:HG22	5:AF:19:GLU:OE1	2.07	0.55
8:AK:13:GLY:HA3	8:AK:17:GLN:OE1	2.07	0.55
1:AA:1142(A):A:H4'	9:AM:25:ARG:HH22	1.71	0.55
9:AM:99:LEU:HD13	9:AM:99:LEU:O	2.06	0.55
2:AB:8:U:O2'	14:AQ:40:ILE:HD13	2.07	0.55
15:AR:2:ASN:O	15:AR:3:ARG:HB2	2.05	0.55
31:BA:1346:A:O2'	31:BA:1347:G:P	2.64	0.55
31:BA:983:A:H2	31:BA:984:C:C6	2.24	0.55
33:BF:42:LEU:HD11	33:BF:46:GLU:OE2	2.06	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:BF:70:VAL:HG12	33:BF:72:LYS:N	2.20	0.55
34:BG:189:PRO:CB	34:BG:194:LEU:HD21	2.36	0.55
36:BI:28:ARG:HA	36:BI:28:ARG:HH11	1.70	0.55
36:BI:38:GLU:HB3	36:BI:64:GLN:O	2.07	0.55
38:BK:29:SER:O	38:BK:32:LYS:HB2	2.05	0.55
39:BL:99:LEU:HB3	39:BL:101:PHE:HE1	1.71	0.55
39:BL:105:ASP:OD2	39:BL:107:ARG:HD3	2.07	0.55
40:BM:71:LEU:HD12	40:BM:72:VAL:N	2.21	0.55
43:BP:78:ILE:O	43:BP:81:LEU:N	2.38	0.55
43:BP:97:PRO:CA	43:BP:110:ARG:HD3	2.36	0.55
31:BA:237:C:O3'	47:BT:25:ARG:NH2	2.40	0.55
54:CA:1149:C:H2'	54:CA:1150:U:O4'	2.06	0.55
54:CA:1152:A:O2'	54:CA:1153:C:H5'	2.06	0.55
54:CA:1450:U:O2	54:CA:1452:C:H5"	2.06	0.55
54:CA:432:A:H2'	54:CA:433:C:O4'	2.06	0.55
52:CD:2:C:H2'	52:CD:3:C:H6	1.70	0.55
32:CE:144:ARG:O	32:CE:147:LYS:HB3	2.06	0.55
32:CE:163:PHE:HA	32:CE:185:ILE:CG1	2.35	0.55
33:CF:143:GLU:C	33:CF:145:GLY:H	2.09	0.55
54:CA:1190:G:OP2	33:CF:5:ILE:HG23	2.07	0.55
40:CM:38:ILE:HD11	40:CM:71:LEU:CD2	2.32	0.55
43:CP:40:ASN:HD21	43:CP:42:ALA:HB3	1.72	0.55
16:D1:17:ILE:HG23	16:D1:39:LEU:HD12	1.88	0.55
17:D2:16:PRO:HA	17:D2:96:ILE:HG22	1.89	0.55
55:DA:1259:G:O2'	55:DA:1260:G:H5'	2.07	0.55
55:DA:1709:U:H1'	55:DA:2860:A:N3	2.22	0.55
55:DA:1963:U:C2'	55:DA:1963:U:O2	2.53	0.55
55:DA:2631:G:N3	55:DA:2810:A:H2	2.05	0.55
55:DA:2872:G:H2'	55:DA:2873:A:C8	2.41	0.55
55:DA:2876:G:OP1	15:DR:4:GLY:N	2.38	0.55
55:DA:660:G:H5'	5:DF:99:TYR:CD2	2.42	0.55
7:DH:123:PHE:O	7:DH:125:VAL:HG23	2.06	0.55
8:DK:77:LEU:CD1	8:DK:140:LEU:HB2	2.23	0.55
8:DK:86:THR:HG22	8:DK:86:THR:O	2.07	0.55
15:DR:3:ARG:HG3	15:DR:7:ILE:CG1	2.36	0.55
18:DS:82:LEU:HD12	18:DS:82:LEU:N	2.21	0.55
21:DV:175:VAL:CB	21:DV:176:PRO:HA	2.36	0.55
16:A1:105:VAL:HG23	16:A1:106:PHE:N	2.21	0.55
1:AA:1329:U:H5"	1:AA:1330:C:H5	1.71	0.55
1:AA:1419:A:H2'	1:AA:1421:G:N7	2.21	0.55
1:AA:1538:G:O2'	1:AA:1539:G:H5'	2.07	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1558:A:HO2'	1:AA:1559:G:P	2.28	0.55
1:AA:2285:C:C5'	1:AA:2286:A:OP2	2.55	0.55
1:AA:27:G:C2'	1:AA:28:A:OP2	2.55	0.55
1:AA:2825:C:H2'	1:AA:2826:A:O4'	2.07	0.55
1:AA:289:A:N3	1:AA:289:A:H2'	2.21	0.55
1:AA:870:A:H2'	1:AA:871:U:O4'	2.05	0.55
2:AB:29:A:H2'	2:AB:30:C:O4'	2.07	0.55
3:AD:239:ARG:O	3:AD:240:ALA:CB	2.51	0.55
4:AE:8:LYS:O	4:AE:9:VAL:HG13	2.06	0.55
5:AF:38:ARG:HD3	5:AF:99:TYR:OH	2.06	0.55
7:AH:33:LEU:CD1	7:AH:79:VAL:HG13	2.36	0.55
15:AR:58:ASN:N	15:AR:58:ASN:HD22	2.03	0.55
20:AU:84:ARG:HB3	20:AU:95:LYS:HE3	1.89	0.55
31:BA:1285:A:O2'	31:BA:1286:A:P	2.64	0.55
31:BA:191(C):G:H5'	31:BA:191(D):U:OP2	2.07	0.55
31:BA:192:U:H2'	31:BA:193:C:C6	2.41	0.55
31:BA:24:U:H2'	31:BA:25:C:C6	2.42	0.55
31:BA:498:A:O2'	31:BA:500:G:H8	1.83	0.55
37:BJ:84:ASN:OD1	52:BD:37:MIA:H131	2.06	0.55
32:BE:109:SER:HA	32:BE:112:VAL:HG23	1.87	0.55
34:BG:199:ASN:O	34:BG:201:GLN:N	2.38	0.55
34:BG:36:ARG:HB2	34:BG:38:TYR:CE2	2.41	0.55
35:BH:48:ALA:HB2	35:BH:57:LYS:HD3	1.88	0.55
39:BL:3:GLN:NE2	39:BL:20:ARG:HH22	2.05	0.55
54:CA:366:C:O2'	54:CA:367:U:P	2.65	0.55
52:CD:1:G:C2'	52:CD:2:C:OP1	2.55	0.55
32:CE:102:LEU:N	32:CE:102:LEU:HD12	2.20	0.55
32:CE:163:PHE:CD2	32:CE:185:ILE:HG13	2.41	0.55
35:CH:64:ARG:HG3	35:CH:64:ARG:HH11	1.72	0.55
38:CK:51:VAL:HG11	38:CK:60:ARG:CG	2.37	0.55
42:CO:27:LEU:HD23	42:CO:64:TYR:OH	2.06	0.55
43:CP:65:LYS:HB2	43:CP:69:GLU:CB	2.36	0.55
44:CQ:33:VAL:HG13	44:CQ:39:LEU:O	2.06	0.55
48:CU:30:ASP:C	48:CU:32:ARG:H	2.10	0.55
16:D1:65:ILE:HG12	16:D1:96:ALA:CB	2.36	0.55
16:D1:95:LEU:HD13	17:D2:4:ILE:HG21	1.88	0.55
55:DA:1931:U:H5	55:DA:1969:A:N7	2.04	0.55
55:DA:2162:G:O2'	55:DA:2163:C:H5'	2.07	0.55
55:DA:2311:A:C8	6:DG:88:ILE:HG13	2.41	0.55
55:DA:2529:G:OP2	55:DA:2530:A:H5''	2.07	0.55
55:DA:270(P):C:H2'	55:DA:270(Q):C:H6	1.72	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:593:G:H2'	55:DA:594:U:C6	2.41	0.55
55:DA:654(R):C:C2	55:DA:654(S):G:C8	2.94	0.55
55:DA:811:U:H2'	11:DO:21:ARG:HG3	1.88	0.55
55:DA:833:U:H1'	11:DO:55:ARG:NH1	2.21	0.55
55:DA:887:A:O2'	55:DA:888:C:C5'	2.54	0.55
55:DA:990:A:OP2	55:DA:991:C:OP2	2.24	0.55
5:DF:114:VAL:HG21	5:DF:202:PHE:CZ	2.41	0.55
8:DK:125:GLU:OE1	8:DK:141:LYS:HA	2.05	0.55
58:DL:55:VAL:HA	58:DL:69:THR:OG1	2.06	0.55
10:DN:1:MET:HE3	10:DN:67:LYS:HE2	1.87	0.55
11:DO:47:ASP:C	11:DO:47:ASP:OD2	2.42	0.55
10:DN:104:ARG:HD3	15:DR:36:GLU:OE2	2.06	0.55
20:DU:19:LYS:O	20:DU:20:TYR:CG	2.60	0.55
20:DU:42:VAL:HG12	20:DU:65:ALA:HB3	1.88	0.55
20:DU:90:LEU:HD22	20:DU:90:LEU:N	2.22	0.55
21:DV:183:LEU:N	21:DV:183:LEU:HD23	2.22	0.55
57:DY:26:LEU:C	57:DY:111:LEU:H	2.09	0.55
57:DY:23:SER:OG	57:DY:68:LEU:CB	2.54	0.55
57:DY:75:GLN:HB3	57:DY:111:LEU:CA	2.33	0.55
13:A0:17:ARG:O	13:A0:20:LEU:HB3	2.06	0.55
26:A4:9:LEU:O	26:A4:9:LEU:HD23	2.07	0.55
1:AA:2372:G:H1'	28:A6:46:HIS:CE1	2.42	0.55
1:AA:1694:C:O2'	1:AA:1695:G:OP2	2.23	0.55
1:AA:2532:G:H4'	1:AA:2657:A:C2	2.42	0.55
1:AA:2648:C:H2'	1:AA:2649:U:H6	1.72	0.55
1:AA:2751:G:O6	7:AH:2:SER:HB3	2.06	0.55
1:AA:38:A:H2'	1:AA:39:C:C6	2.42	0.55
1:AA:87:C:OP2	1:AA:90:U:O4	2.24	0.55
1:AA:2823:A:OP1	4:AE:113:PHE:HB2	2.06	0.55
4:AE:8:LYS:O	4:AE:9:VAL:CG2	2.52	0.55
6:AG:67:LYS:HG3	26:A4:6:HIS:CG	2.42	0.55
7:AH:118:PRO:HG2	7:AH:121:ILE:CG1	2.37	0.55
11:AO:124:LYS:HA	11:AO:143:GLY:O	2.06	0.55
12:AP:38:GLU:O	12:AP:127:ILE:HD13	2.06	0.55
21:AV:145:GLU:HA	21:AV:174:VAL:CG1	2.25	0.55
31:BA:1058:G:H2'	31:BA:1059:C:C6	2.42	0.55
31:BA:1321:C:C5'	31:BA:1322:C:H5''	2.36	0.55
31:BA:173:U:H5''	31:BA:197:A:O4'	2.07	0.55
31:BA:432:A:H3'	31:BA:433:C:C6	2.40	0.55
31:BA:439:A:OP2	31:BA:493:G:N1	2.35	0.55
31:BA:551:U:H2'	31:BA:552:U:H6	1.72	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:703:G:O2'	31:BA:704:A:OP2	2.19	0.55
31:BA:75:C:H2'	31:BA:76:G:O4'	2.06	0.55
52:BC:19:G:H21	52:BC:57:G:H1'	1.72	0.55
52:BC:43:C:O2'	52:BC:44:G:OP1	2.21	0.55
52:BC:58:A:H2'	52:BC:60:U:OP2	2.05	0.55
52:BD:26:A:H61	52:BD:44:G:H1	1.55	0.55
32:BE:17:PHE:HZ	32:BE:44:LEU:HA	1.71	0.55
34:BG:19:LEU:HD12	34:BG:21:LEU:CD2	2.36	0.55
35:BH:87:SER:HB3	35:BH:131:ILE:HD13	1.88	0.55
36:BI:14:LEU:HD21	36:BI:18:GLN:CB	2.37	0.55
42:BO:32:PHE:CB	42:BO:84:LEU:HD21	2.36	0.55
44:BQ:18:VAL:C	44:BQ:20:ALA:H	2.09	0.55
50:BW:26:ASN:CB	50:BW:71:THR:HG23	2.34	0.55
54:CA:1283:G:O2'	54:CA:1284:C:H5'	2.06	0.55
54:CA:1310:G:O2'	54:CA:1311:G:H5'	2.07	0.55
54:CA:451:A:H4'	54:CA:452:A:O4'	2.06	0.55
54:CA:624:C:O3'	46:CS:10:GLY:HA2	2.06	0.55
52:CD:37:MIA:H3'	52:CD:38:A:H8	1.71	0.55
42:CO:64:TYR:O	42:CO:65:GLU:HB2	2.07	0.55
47:CT:78:GLU:O	47:CT:78:GLU:HG2	2.06	0.55
54:CA:1305:G:C5'	51:CX:4:GLY:HA3	2.34	0.55
28:D6:27:LYS:CB	28:D6:27:LYS:HZ2	2.13	0.55
55:DA:1020:A:N6	55:DA:1141:U:HO2'	2.04	0.55
55:DA:1026:U:H1'	55:DA:1027:A:O5'	2.07	0.55
55:DA:1180:C:C2'	55:DA:1181:C:H5'	2.35	0.55
55:DA:1509:C:N4	55:DA:1511:A:N6	2.54	0.55
55:DA:1448:G:H1'	55:DA:1528:A:H62	1.71	0.55
55:DA:1530:G:H2'	55:DA:1531:C:H6	1.72	0.55
55:DA:1771:C:C1'	55:DA:1786:A:C8	2.89	0.55
55:DA:877:U:C2'	55:DA:878:A:O5'	2.53	0.55
55:DA:894:C:C2'	55:DA:895:U:C6	2.85	0.55
4:DE:59:VAL:CG1	4:DE:63:LEU:HB3	2.36	0.55
5:DF:11:VAL:HB	5:DF:18:ARG:HG3	1.88	0.55
56:DJ:3:LEU:O	56:DJ:5:ILE:N	2.40	0.55
58:DL:93:ARG:HA	58:DL:135:GLY:C	2.26	0.55
58:DL:21:PRO:CB	58:DL:22:PRO:CD	2.82	0.55
11:DO:62:LEU:O	11:DO:62:LEU:CD2	2.41	0.55
55:DA:1341:U:O2'	19:DT:55:ASN:HB3	2.05	0.55
19:DT:63:LYS:O	19:DT:64:LYS:HD2	2.05	0.55
21:DV:152:ALA:C	21:DV:154:ASP:N	2.59	0.55
57:DY:33:PRO:O	57:DY:34:ALA:C	2.44	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:5:ARG:N	57:DY:5:ARG:HD2	2.21	0.55
57:DY:90:ALA:H	56:DJ:15:ALA:CA	2.18	0.55
1:AA:1278:A:O3'	13:A0:34:ILE:CD1	2.54	0.55
17:A2:43:GLU:O	17:A2:44:LYS:HD3	2.07	0.55
17:A2:58:VAL:HB	17:A2:98:GLU:CG	2.37	0.55
1:AA:1283:G:N2	1:AA:1286:A:OP2	2.37	0.55
1:AA:1667:G:O2'	1:AA:1669:A:N6	2.39	0.55
1:AA:1683:C:H2'	1:AA:1684:C:H6	1.71	0.55
1:AA:2278:A:H2'	1:AA:2279:G:O5'	2.07	0.55
1:AA:960:A:C4'	1:AA:2457:U:H4'	2.36	0.55
1:AA:957:A:C2	1:AA:2458:G:H4'	2.41	0.55
1:AA:2553:G:H2'	1:AA:2554:U:O4'	2.07	0.55
1:AA:270(R):G:H2'	1:AA:270(S):G:C8	2.41	0.55
1:AA:2777:G:H5''	1:AA:2778:A:C5'	2.34	0.55
1:AA:2848:G:O2'	1:AA:2849:U:P	2.63	0.55
1:AA:440:G:H2'	1:AA:441:U:H6	1.72	0.55
1:AA:752:A:HO2'	1:AA:753:C:P	2.27	0.55
3:AD:7:LYS:NZ	3:AD:7:LYS:HB3	2.21	0.55
5:AF:9:ILE:HG12	5:AF:15:SER:CA	2.36	0.55
6:AG:101:ILE:HD13	26:A4:9:LEU:HD11	1.88	0.55
11:AO:37:GLY:O	11:AO:39:LYS:N	2.39	0.55
11:AO:6:LEU:O	11:AO:7:ARG:HG2	2.07	0.55
18:AS:20:VAL:O	18:AS:23:LEU:HB2	2.06	0.55
21:AV:98:MET:O	21:AV:125:LEU:HA	2.07	0.55
21:AV:6:LYS:O	21:AV:7:ALA:CB	2.55	0.55
25:AX:19:GLN:HE22	25:AX:52:HIS:HE1	1.53	0.55
25:AX:19:GLN:O	25:AX:22:ALA:HB3	2.06	0.55
23:AZ:23:LYS:HD3	23:AZ:28:GLY:HA3	1.89	0.55
31:BA:1056:U:H5'	33:BF:163:ALA:CB	2.37	0.55
31:BA:430:A:OP2	34:BG:8:VAL:HG22	2.07	0.55
31:BA:744:C:O2'	31:BA:745:C:H5'	2.07	0.55
52:BB:8:U:OP2	52:BB:8:U:H6	1.89	0.55
52:BD:59:U:H2'	52:BD:60:U:C5'	2.37	0.55
33:BF:63:ASN:O	33:BF:64:VAL:HB	2.06	0.55
34:BG:23:GLY:C	34:BG:24:GLU:HG2	2.26	0.55
35:BH:35:GLY:HA3	35:BH:112:LEU:O	2.07	0.55
37:BJ:20:ASP:HB3	37:BJ:23:VAL:HG23	1.87	0.55
39:BL:14:VAL:O	39:BL:65:VAL:HG23	2.06	0.55
40:BM:5:ARG:HA	40:BM:73:ASP:OD2	2.07	0.55
43:BP:97:PRO:HA	43:BP:110:ARG:HD3	1.89	0.55
45:BR:3:ILE:HD13	45:BR:3:ILE:H	1.72	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:BT:32:TYR:O	47:BT:34:LYS:N	2.40	0.55
36:BI:91:VAL:HG11	48:BU:72:ARG:CZ	2.35	0.55
54:CA:1120:G:H2'	54:CA:1121:U:C6	2.42	0.55
54:CA:1435:G:H2'	54:CA:1436:U:C6	2.42	0.55
54:CA:34:C:O2'	54:CA:35:G:H5'	2.06	0.55
54:CA:353:A:H5'	54:CA:353:A:C8	2.36	0.55
54:CA:502:G:H2'	54:CA:503:C:C6	2.42	0.55
54:CA:49:U:H5''	54:CA:50:A:OP2	2.07	0.55
33:CF:78:GLY:HA3	33:CF:83:ARG:H	1.72	0.55
37:CJ:57:GLU:O	37:CJ:61:VAL:HG23	2.06	0.55
38:CK:5:PRO:O	38:CK:8:ASP:HB3	2.07	0.55
40:CM:48:THR:HG23	40:CM:62:HIS:HB3	1.89	0.55
36:CI:46:ARG:NH2	48:CU:37:VAL:HG23	2.22	0.55
55:DA:2870:C:H5'	13:D0:61:HIS:HE1	1.70	0.55
17:D2:3:ALA:O	17:D2:4:ILE:HD13	2.06	0.55
55:DA:1331:A:H2'	55:DA:1333:C:H5	1.72	0.55
55:DA:174:C:H2'	55:DA:175:G:O4'	2.07	0.55
55:DA:1904:G:C2'	55:DA:1905:C:H5'	2.37	0.55
55:DA:2167:U:H2'	55:DA:2167:U:O2	2.05	0.55
55:DA:2061:G:OP2	55:DA:2502:G:H5'	2.07	0.55
55:DA:2650:U:H2'	55:DA:2651:C:C6	2.41	0.55
55:DA:524:U:H4'	55:DA:554:U:H4'	1.87	0.55
2:DB:105:G:O2'	2:DB:106:G:H5'	2.07	0.55
4:DE:2:LYS:NZ	4:DE:95:ILE:O	2.40	0.55
7:DH:20:ALA:CB	7:DH:21:PRO:HD2	2.34	0.55
57:DY:89:ALA:C	56:DJ:15:ALA:HB2	2.26	0.55
56:DJ:18:LEU:O	56:DJ:21:LYS:HB2	2.07	0.55
56:DI:29:GLU:CD	56:DJ:6:GLU:OE2	2.45	0.55
58:DL:78:ILE:HG12	58:DL:131:ALA:CB	2.36	0.55
58:DL:84:LEU:N	58:DL:84:LEU:HD12	2.22	0.55
14:DQ:18:ILE:C	14:DQ:19:LYS:O	2.43	0.55
14:DQ:46:VAL:HG12	14:DQ:47:THR:N	2.22	0.55
55:DA:2849:U:OP1	15:DR:95:ARG:NH1	2.40	0.55
57:DY:73:GLY:CA	57:DY:119:ALA:C	2.74	0.55
57:DY:135:ARG:HD2	57:DY:138:LEU:HG	1.88	0.55
57:DY:15:GLU:O	57:DY:15:GLU:CG	2.46	0.55
57:DY:50:ARG:O	57:DY:51:LEU:CD2	2.55	0.55
57:DY:59:ILE:HD11	57:DY:60:ARG:CZ	2.36	0.55
16:A1:79:PHE:CD2	16:A1:79:PHE:C	2.80	0.55
22:A3:72:ARG:NH2	22:A3:75:LEU:HD12	2.22	0.55
26:A4:9:LEU:CG	26:A4:25:TYR:HB3	2.37	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:A8:14:VAL:HG12	30:A8:15:LYS:H	1.72	0.55
1:AA:1162:G:H2'	1:AA:1163:G:H8	1.71	0.55
1:AA:2211:G:O2'	1:AA:2212:A:OP2	2.21	0.55
1:AA:2505:G:HO2'	1:AA:2506:U:H6	1.52	0.55
1:AA:262:A:H2'	1:AA:263:C:O4'	2.06	0.55
1:AA:2750:A:H5''	1:AA:2751:G:OP2	2.07	0.55
1:AA:2050:C:H1'	4:AE:156:MET:HE2	1.88	0.55
4:AE:203:LYS:O	4:AE:204:ALA:HB2	2.06	0.55
4:AE:24:THR:O	4:AE:25:VAL:HB	2.07	0.55
4:AE:37:ARG:HD3	4:AE:44:TYR:OH	2.07	0.55
6:AG:114:ILE:HB	6:AG:117:PHE:HB2	1.88	0.55
7:AH:10:PRO:HG3	7:AH:69:ARG:CZ	2.36	0.55
7:AH:72:ILE:O	7:AH:75:ALA:HB3	2.07	0.55
8:AK:125:GLU:HA	8:AK:141:LYS:CB	2.35	0.55
1:AA:2377:A:H4'	14:AQ:111:GLU:O	2.07	0.55
21:AV:177:PRO:O	21:AV:180:VAL:HA	2.07	0.55
24:AW:51:ARG:CZ	24:AW:55:ARG:HH12	2.20	0.55
2:AB:83:G:H5''	25:AX:52:HIS:ND1	2.22	0.55
31:BA:1037:C:H2'	31:BA:1038:C:H6	1.72	0.55
31:BA:1225:A:N3	31:BA:1225:A:H2'	2.22	0.55
31:BA:1253:G:H2'	31:BA:1254:C:H6	1.72	0.55
31:BA:1275:A:H2'	31:BA:1276:G:O4'	2.07	0.55
52:BC:38:A:H2'	52:BC:39:U:H5'	1.88	0.55
32:BE:59:GLU:O	32:BE:62:ALA:HB3	2.07	0.55
34:BG:98:GLU:OE2	34:BG:103:ASN:ND2	2.40	0.55
34:BG:209:ARG:O	34:BG:209:ARG:HG2	2.06	0.55
35:BH:11:ILE:HD13	35:BH:105:VAL:HA	1.88	0.55
37:BJ:27:ILE:H	37:BJ:27:ILE:HD12	1.71	0.55
37:BJ:77:SER:HB2	52:BD:32:U:O3'	2.06	0.55
31:BA:1368:G:OP2	39:BL:112:LYS:HD2	2.06	0.55
43:BP:115:LYS:H	43:BP:115:LYS:HD2	1.71	0.55
48:BU:31:LEU:HD23	48:BU:31:LEU:N	2.22	0.55
49:BV:36:ARG:NH2	49:BV:75:ALA:HB3	2.22	0.55
49:BV:40:ILE:O	49:BV:40:ILE:HG23	2.06	0.55
51:BX:8:THR:HG22	51:BX:9:ARG:N	2.21	0.55
54:CA:1189:C:P	40:CM:51:ARG:NH2	2.80	0.55
54:CA:439:A:OP2	54:CA:493:G:N1	2.39	0.55
52:CD:8:U:O4'	52:CD:48:C:O2'	2.25	0.55
32:CE:61:LEU:HD21	32:CE:161:ALA:HB3	1.87	0.55
32:CE:75:LYS:HA	32:CE:78:GLN:HE21	1.71	0.55
54:CA:1189:C:O3'	33:CF:5:ILE:HD12	2.07	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:CL:113:LYS:N	39:CL:113:LYS:HD2	2.22	0.55
42:CO:43:VAL:O	42:CO:44:THR:HG23	2.05	0.55
43:CP:108:ARG:HH11	43:CP:108:ARG:HA	1.69	0.55
44:CQ:24:CYS:HA	44:CQ:38:GLY:O	2.07	0.55
45:CR:3:ILE:HD13	45:CR:3:ILE:N	2.21	0.55
46:CS:6:LEU:CD1	46:CS:19:ILE:HD13	2.37	0.55
49:CV:44:MET:O	49:CV:46:GLY:N	2.40	0.55
17:D2:91:TYR:C	17:D2:91:TYR:CD1	2.80	0.55
55:DA:1265:A:H1'	55:DA:1267:U:C6	2.42	0.55
55:DA:1905:C:H2'	55:DA:1930:G:C8	2.42	0.55
55:DA:2332:U:H5'	55:DA:2336:A:N6	2.22	0.55
55:DA:412:A:N7	55:DA:2411:A:H2	2.04	0.55
55:DA:1799:G:H2'	3:DD:181:GLU:OE2	2.06	0.55
6:DG:126:ASP:OD1	6:DG:130:ASN:HB2	2.06	0.55
56:DI:14:GLN:O	56:DI:15:ALA:HB3	2.06	0.55
56:DI:24:ILE:CD1	56:DI:25:ASP:HB2	2.37	0.55
11:DO:30:THR:O	11:DO:31:ALA:CB	2.54	0.55
15:DR:50:ILE:HA	15:DR:99:LEU:CD1	2.36	0.55
18:DS:95:ILE:O	18:DS:95:ILE:HD12	2.06	0.55
21:DV:196:VAL:C	21:DV:197:ILE:CD1	2.75	0.55
13:A0:10:LEU:O	13:A0:11:ASN:C	2.45	0.55
17:A2:35:LEU:HD23	17:A2:37:VAL:CG2	2.37	0.55
17:A2:97:LYS:O	17:A2:98:GLU:HG2	2.07	0.55
1:AA:517:C:OP1	27:A5:16:ARG:NH2	2.40	0.55
1:AA:2815:C:O2'	27:A5:43:HIS:HD2	1.90	0.55
1:AA:593:G:C1'	30:A8:4:MET:HE2	2.36	0.55
1:AA:1055:G:H2'	1:AA:1056:G:H5'	1.89	0.55
1:AA:1097:U:H2'	1:AA:1098:A:O4'	2.07	0.55
1:AA:347:A:O2'	1:AA:348:G:H5'	2.06	0.55
1:AA:654(C):G:C3'	1:AA:654(D):G:C8	2.90	0.55
2:AB:88:C:H3'	2:AB:89:G:C8	2.37	0.55
5:AF:20:LEU:HD13	5:AF:199:TRP:CH2	2.42	0.55
5:AF:25:PRO:C	5:AF:27:GLU:H	2.09	0.55
5:AF:64:ILE:HG13	5:AF:65:TRP:N	2.22	0.55
6:AG:113:ARG:HD3	6:AG:140:ILE:O	2.06	0.55
6:AG:34:LEU:HD21	6:AG:159:VAL:HG23	1.89	0.55
8:AK:76:THR:HG21	8:AK:138:ILE:CG1	2.36	0.55
11:AO:147:LEU:HD23	11:AO:148:LEU:O	2.05	0.55
18:AS:38:TYR:O	27:A5:28:PRO:HB3	2.07	0.55
19:AT:14:SER:H	19:AT:17:ALA:HB3	1.71	0.55
31:BA:1052:U:H2'	31:BA:1055:A:OP1	2.07	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:1144:G:N2	31:BA:1146:A:N6	2.54	0.55
31:BA:1357:A:N7	31:BA:1358:U:C5	2.75	0.55
31:BA:1439:C:OP1	50:BW:38:LYS:HD2	2.05	0.55
31:BA:382:A:H2'	31:BA:383:A:C8	2.41	0.55
31:BA:429:U:O2'	31:BA:430:A:H5''	2.07	0.55
31:BA:715:A:H2'	31:BA:716:A:C8	2.42	0.55
33:BF:164:ARG:HG2	33:BF:165:THR:H	1.71	0.55
34:BG:49:ARG:HA	34:BG:49:ARG:HE	1.70	0.55
40:BM:32:ALA:HA	40:BM:75:ILE:HD11	1.89	0.55
54:CA:1348:U:N3	54:CA:1374:A:C2	2.68	0.55
54:CA:1452:C:C2'	54:CA:1453:G:OP2	2.54	0.55
52:CB:37:MIA:H122	52:CB:38:A:C2	2.42	0.55
52:CC:19:G:O2'	52:CC:20:U:P	2.65	0.55
33:CF:5:ILE:HD13	33:CF:5:ILE:H	1.72	0.55
54:CA:542:G:H5'	34:CG:41:GLY:HA3	1.88	0.55
37:CJ:111:ARG:NH1	37:CJ:113:GLU:CD	2.60	0.55
37:CJ:115:ARG:HB2	37:CJ:118:VAL:CG1	2.37	0.55
38:CK:51:VAL:HG11	38:CK:60:ARG:HG3	1.88	0.55
38:CK:88:LYS:HB3	38:CK:89:PRO:HD2	1.89	0.55
42:CO:90:VAL:O	42:CO:92:ASP:N	2.40	0.55
43:CP:2:ALA:O	43:CP:10:PRO:HD2	2.06	0.55
43:CP:5:ALA:O	43:CP:7:VAL:N	2.39	0.55
16:D1:92:ARG:O	16:D1:92:ARG:HG2	2.06	0.55
55:DA:1162:G:H21	17:D2:89:GLN:HE22	1.54	0.55
26:D4:68:ARG:C	26:D4:70:GLY:N	2.58	0.55
55:DA:1171:G:C6	55:DA:1174:A:N6	2.74	0.55
55:DA:2355:C:O4'	22:D3:36:ILE:HD11	2.06	0.55
55:DA:2496:C:OP1	12:DP:81:VAL:HG12	2.07	0.55
55:DA:991:C:H2'	55:DA:992:C:H6	1.72	0.55
2:DB:11:C:OP2	2:DB:12:C:N4	2.32	0.55
2:DB:12:C:C4'	2:DB:13:A:OP1	2.52	0.55
3:DD:96:HIS:CE1	3:DD:102:LYS:NZ	2.75	0.55
3:DD:136:ILE:HG22	3:DD:165:ILE:HD12	1.89	0.55
56:DI:17:VAL:HB	56:DI:21:LYS:HE3	1.89	0.55
58:DL:70:LYS:O	58:DL:72:PRO:HD3	2.07	0.55
58:DL:78:ILE:HA	58:DL:82:ALA:CB	2.34	0.55
10:DN:104:ARG:HG2	10:DN:104:ARG:NH1	2.19	0.55
18:DS:70:TYR:N	18:DS:70:TYR:CD2	2.75	0.55
21:DV:150:LEU:O	21:DV:151:HIS:ND1	2.40	0.55
57:DY:1:MET:HE3	57:DY:3:ASN:ND2	2.22	0.55
13:A0:84:ALA:HB3	13:A0:85:PRO:HD3	1.89	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:A0:87:TYR:HE1	13:A0:117:VAL:HG13	1.72	0.55
17:A2:5:VAL:HG22	17:A2:6:LYS:N	2.22	0.55
26:A4:34:GLU:O	26:A4:35:VAL:C	2.46	0.55
1:AA:1259:G:O2'	1:AA:1260:G:H5'	2.06	0.55
1:AA:1403:C:H5''	1:AA:1471:A:C1'	2.37	0.55
1:AA:1458:C:H5''	1:AA:1459:G:C5'	2.36	0.55
1:AA:1537:C:H6	1:AA:1537:C:O5'	1.89	0.55
1:AA:1754:C:OP1	15:AR:96:ARG:NH1	2.39	0.55
1:AA:2147:G:C8	1:AA:2147:G:H3'	2.42	0.55
1:AA:2694:G:O2'	1:AA:2695:C:H5'	2.07	0.55
1:AA:2791:C:H2'	1:AA:2792:G:O4'	2.07	0.55
4:AE:27:LEU:HD13	4:AE:181:LEU:HD22	1.87	0.55
4:AE:49:LEU:HD23	4:AE:49:LEU:N	2.22	0.55
8:AK:61:ARG:HA	8:AK:64:GLU:HG2	1.88	0.55
9:AM:7:LYS:HZ3	9:AM:7:LYS:N	2.05	0.55
1:AA:627:A:H62	11:AO:116:GLY:HA2	1.71	0.55
18:AS:106:ILE:HG13	18:AS:106:ILE:O	2.05	0.55
21:AV:108:PRO:HG3	21:AV:142:SER:HA	1.89	0.55
21:AV:5:LEU:HD13	21:AV:5:LEU:O	2.06	0.55
24:AW:46:GLN:O	24:AW:49:LYS:HE3	2.07	0.55
25:AX:29:ARG:HG3	25:AX:29:ARG:HH11	1.72	0.55
31:BA:1129:C:N4	31:BA:1141:C:N4	2.55	0.55
31:BA:1291:G:H4'	39:BL:39:GLY:HA3	1.87	0.55
31:BA:151:A:O2'	31:BA:152:A:H5'	2.07	0.55
52:BB:56:C:H2'	52:BB:57:G:H8	1.71	0.55
32:BE:21:ARG:NH2	32:BE:38:GLY:HA3	2.19	0.55
34:BG:14:ARG:O	34:BG:14:ARG:CG	2.54	0.55
34:BG:201:GLN:O	34:BG:205:GLU:HG3	2.07	0.55
37:BJ:109:ASN:HA	37:BJ:119:ARG:HE	1.71	0.55
39:BL:20:ARG:O	39:BL:60:ASP:HB3	2.07	0.55
31:BA:1250:A:OP1	39:BL:66:ARG:HG2	2.06	0.55
42:BO:117:ARG:NH2	42:BO:124:LYS:HA	2.21	0.55
44:BQ:9:LYS:HA	44:BQ:12:ARG:HB3	1.89	0.55
50:BW:67:ALA:HA	50:BW:73:HIS:H	1.71	0.55
54:CA:1019:C:C2'	54:CA:1020:U:H5'	2.36	0.55
54:CA:1028(B):C:H3'	54:CA:1029:G:C5'	2.36	0.55
54:CA:174:C:H2'	54:CA:175:C:H6	1.71	0.55
54:CA:309:G:H1'	54:CA:608:A:C2	2.41	0.55
54:CA:746:A:H2'	54:CA:747:C:O4'	2.07	0.55
54:CA:849:C:O2'	54:CA:850:U:H5'	2.06	0.55
32:CE:175:ARG:HG2	32:CE:175:ARG:NH1	2.22	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:CE:56:ARG:HH11	32:CE:56:ARG:HG2	1.73	0.55
35:CH:32:VAL:HG12	35:CH:33:VAL:N	2.21	0.55
37:CJ:76:ARG:HD3	37:CJ:156:TRP:HZ2	1.72	0.55
38:CK:121:ASP:HB2	38:CK:125:ARG:HH22	1.72	0.55
46:CS:40:ASP:C	46:CS:42:ARG:H	2.10	0.55
51:CX:15:ARG:HG2	51:CX:15:ARG:NH1	2.21	0.55
13:D0:57:ARG:HD3	13:D0:59:ASP:CG	2.28	0.55
17:D2:66:ARG:HB3	17:D2:66:ARG:HH11	1.72	0.55
17:D2:76:LYS:HB3	17:D2:79:VAL:HG11	1.88	0.55
22:D3:31:VAL:HB	22:D3:35:ASN:ND2	2.21	0.55
28:D6:25:LYS:HE2	28:D6:27:LYS:HE2	1.89	0.55
28:D6:30:THR:CA	28:D6:31:PRO:O	2.55	0.55
55:DA:1056:G:N2	55:DA:1087:G:H1	2.03	0.55
55:DA:1144:G:H2'	55:DA:1145:C:H6	1.72	0.55
55:DA:1180:C:H2'	55:DA:1181:C:C5'	2.36	0.55
55:DA:1925:C:N4	55:DA:1926:U:C5	2.75	0.55
55:DA:570:G:H2'	55:DA:2030:A:N7	2.22	0.55
55:DA:2038:G:H2'	55:DA:2039:C:H6	1.72	0.55
55:DA:2649:U:H2'	55:DA:2650:U:C6	2.42	0.55
55:DA:2761:G:H5'	55:DA:2761:G:C8	2.42	0.55
55:DA:884:C:C2'	55:DA:885:C:OP1	2.55	0.55
4:DE:179:GLU:O	4:DE:180:ASN:HB2	2.06	0.55
56:DJ:12:LEU:H	56:DJ:13:SER:C	2.10	0.55
8:DK:132:PRO:O	8:DK:133:HIS:ND1	2.40	0.55
8:DK:71:ILE:O	8:DK:71:ILE:HD13	2.07	0.55
58:DL:86:LYS:NZ	58:DL:86:LYS:HA	2.22	0.55
12:DP:138:ASP:O	12:DP:139:GLU:O	2.25	0.55
14:DQ:27:SER:HA	14:DQ:88:ASP:CB	2.37	0.55
57:DY:13:LEU:HD13	57:DY:13:LEU:C	2.28	0.55
57:DY:52:PHE:O	57:DY:53:VAL:HG22	2.07	0.55
1:AA:1085:A:H2'	1:AA:1086:A:H8	1.72	0.54
1:AA:1154:G:OP1	16:A1:58:ARG:HD2	2.06	0.54
1:AA:1220:A:H5'	1:AA:1221:C:OP2	2.07	0.54
1:AA:1280:G:C3'	1:AA:1281:G:H5''	2.38	0.54
1:AA:1300:U:O2'	1:AA:1301:A:OP2	2.21	0.54
1:AA:1694:C:O2'	1:AA:1695:G:P	2.66	0.54
1:AA:1887:C:C3'	1:AA:1888:G:H5''	2.36	0.54
1:AA:2285:C:H41	28:A6:27:LYS:HE3	1.72	0.54
1:AA:2642:G:H5''	9:AM:78:TYR:CD2	2.42	0.54
1:AA:2765:A:H2	1:AA:2766:G:O4'	1.90	0.54
1:AA:403:U:HO2'	1:AA:404:C:P	2.30	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:403:U:O2'	1:AA:404:C:P	2.66	0.54
1:AA:265:A:N6	1:AA:428:A:C8	2.75	0.54
1:AA:467:G:O2'	1:AA:468:G:H5'	2.07	0.54
3:AD:32:SER:O	3:AD:33:LEU:HB2	2.06	0.54
7:AH:151:ILE:HG22	7:AH:151:ILE:O	2.08	0.54
15:AR:48:ILE:HD12	15:AR:48:ILE:N	2.22	0.54
20:AU:48:ALA:O	20:AU:50:ARG:N	2.40	0.54
21:AV:63:ASP:C	21:AV:65:GLN:H	2.10	0.54
21:AV:8:TYR:N	21:AV:8:TYR:CD2	2.73	0.54
23:AZ:83:GLU:OE2	23:AZ:85:LEU:HD23	2.07	0.54
31:BA:1041:A:H3'	31:BA:1042:G:H5''	1.89	0.54
31:BA:978:A:H1'	31:BA:1322:C:O2	2.07	0.54
31:BA:828:A:H2'	31:BA:829:G:O4'	2.07	0.54
32:BE:103:THR:HA	32:BE:180:LEU:HD11	1.89	0.54
33:BF:155:GLY:O	33:BF:196:LEU:HD13	2.07	0.54
34:BG:14:ARG:HG3	34:BG:14:ARG:NH1	2.21	0.54
35:BH:20:GLN:HE22	35:BH:21:ALA:HB3	1.72	0.54
36:BI:12:PRO:O	36:BI:14:LEU:N	2.38	0.54
39:BL:26:VAL:CG2	39:BL:61:ALA:HB3	2.29	0.54
39:BL:15:ALA:HA	39:BL:65:VAL:HA	1.89	0.54
43:BP:15:VAL:C	43:BP:17:VAL:H	2.09	0.54
47:BT:59:ILE:HG23	47:BT:71:PHE:HB3	1.88	0.54
49:BV:53:ASN:CB	49:BV:77:THR:HG22	2.37	0.54
50:BW:67:ALA:HA	50:BW:73:HIS:HA	1.89	0.54
51:BX:15:ARG:HH11	51:BX:15:ARG:CB	2.20	0.54
54:CA:222:U:H2'	54:CA:223:U:C6	2.42	0.54
54:CA:748:C:H1'	54:CA:749:C:H5	1.72	0.54
54:CA:792:A:C2'	54:CA:794:A:N7	2.71	0.54
52:CC:36:A:H5'	52:CC:36:A:C8	2.36	0.54
32:CE:141:GLU:HA	32:CE:144:ARG:HD3	1.89	0.54
32:CE:92:TYR:CZ	32:CE:151:GLY:HA3	2.41	0.54
54:CA:970:C:N4	39:CL:128:ARG:OXT	2.39	0.54
39:CL:17:VAL:HG21	39:CL:81:ILE:N	2.22	0.54
41:CN:21:ILE:HD12	41:CN:21:ILE:N	2.22	0.54
43:CP:92:HIS:HA	43:CP:110:ARG:NH2	2.22	0.54
33:CF:33:LEU:HD11	44:CQ:53:LEU:HD23	1.89	0.54
49:CV:64:GLU:HG3	49:CV:65:ASN:H	1.69	0.54
28:D6:43:CYS:SG	28:D6:44:ARG:NH1	2.80	0.54
55:DA:1174:A:N7	55:DA:1175:U:C1'	2.68	0.54
55:DA:1803:A:C8	55:DA:1804:C:C5	2.95	0.54
55:DA:2657:A:C1'	55:DA:2665:A:N6	2.69	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2752:C:OP2	55:DA:2752:C:C6	2.55	0.54
55:DA:2884:U:C2'	55:DA:2885:C:H5'	2.37	0.54
55:DA:557:U:H2'	55:DA:558:G:C8	2.41	0.54
55:DA:814:C:O2'	55:DA:815:C:H5'	2.06	0.54
8:DK:10:GLU:O	8:DK:11:ASN:HB3	2.06	0.54
58:DL:108:ALA:HA	58:DL:111:LYS:CD	2.37	0.54
11:DO:120:ALA:HB1	11:DO:138:LEU:CA	2.37	0.54
15:DR:41:ARG:C	15:DR:42:ILE:HG13	2.26	0.54
19:DT:83:VAL:HG13	19:DT:87:GLN:HE21	1.71	0.54
21:DV:116:VAL:HG13	21:DV:117:LEU:N	2.22	0.54
24:DW:28:LYS:NZ	24:DW:56:GLN:HE22	2.04	0.54
57:DY:5:ARG:O	57:DY:7:VAL:CB	2.55	0.54
57:DY:51:LEU:HD11	57:DY:82:PHE:CA	2.37	0.54
23:DZ:92:LYS:O	23:DZ:95:LEU:N	2.39	0.54
17:A2:48:GLY:O	17:A2:49:THR:O	2.24	0.54
17:A2:4:ILE:HD12	17:A2:4:ILE:N	2.22	0.54
29:A7:8:ASN:ND2	29:A7:8:ASN:C	2.59	0.54
29:A7:8:ASN:HD22	29:A7:9:ARG:N	2.05	0.54
30:A8:50:LEU:O	30:A8:51:ALA:HB2	2.06	0.54
1:AA:99:U:H1'	1:AA:102:G:C2	2.43	0.54
1:AA:1998:G:H2'	1:AA:1999:C:O4'	2.07	0.54
1:AA:2712:U:O2'	1:AA:2712(A):A:H8	1.90	0.54
1:AA:28:A:N6	1:AA:512:G:H1'	2.22	0.54
2:AB:12:C:C4'	2:AB:13:A:OP1	2.55	0.54
2:AB:95:U:O4	2:AB:96:G:O6	2.25	0.54
5:AF:26:ALA:O	5:AF:27:GLU:HG2	2.07	0.54
9:AM:36:GLY:C	9:AM:42:TRP:HB2	2.27	0.54
11:AO:127:ALA:O	11:AO:147:LEU:HA	2.06	0.54
11:AO:98:GLU:HG3	11:AO:99:LEU:N	2.21	0.54
18:AS:68:ARG:HH22	18:AS:112:GLY:CA	2.21	0.54
24:AW:17:SER:HB2	24:AW:18:PRO:HA	1.86	0.54
25:AX:6:VAL:CG1	25:AX:54:VAL:HG11	2.35	0.54
53:B1:35:A:O2'	53:B1:36:G:H5'	2.08	0.54
31:BA:1190:G:OP1	33:BF:5:ILE:N	2.37	0.54
31:BA:4:U:O4	38:BK:105:ARG:HD3	2.07	0.54
31:BA:518:C:H5'	31:BA:519:C:C6	2.42	0.54
31:BA:674:G:H2'	31:BA:675:A:H8	1.72	0.54
31:BA:792:A:C4	31:BA:794:A:N6	2.75	0.54
31:BA:859:A:H2'	31:BA:860:A:O4'	2.07	0.54
31:BA:948:C:H2'	31:BA:949:A:H8	1.71	0.54
32:BE:88:ALA:CA	32:BE:226:ARG:HH12	2.18	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:BJ:76:ARG:HD3	37:BJ:89:MET:HG3	1.89	0.54
38:BK:109:ILE:HG12	38:BK:110:ALA:N	2.22	0.54
38:BK:1:MET:H3	38:BK:1:MET:HE2	1.72	0.54
39:BL:95:LYS:NZ	39:BL:96:LEU:HD13	2.22	0.54
41:BN:121:PRO:HB2	41:BN:126:ARG:HG3	1.89	0.54
50:BW:56:MET:HG3	50:BW:84:LEU:HD12	1.88	0.54
54:CA:1443:G:H2'	15:DR:122:ASP:OD2	2.07	0.54
54:CA:703:G:O2'	54:CA:704:A:OP2	2.24	0.54
52:CB:36:A:H62	52:CB:37:MIA:H163	1.72	0.54
52:CC:20:U:H3'	52:CC:21:A:H5''	1.89	0.54
52:CC:19:G:C4	52:CC:57:G:N2	2.75	0.54
52:CD:43:C:H5'	52:CD:44:G:OP2	2.06	0.54
32:CE:219:VAL:O	32:CE:222:ILE:HB	2.06	0.54
33:CF:23:TYR:CD2	33:CF:24:ALA:N	2.76	0.54
33:CF:94:LEU:HD12	33:CF:95:THR:N	2.21	0.54
38:CK:64:LYS:C	38:CK:65:TYR:CD1	2.80	0.54
39:CL:25:LYS:HG3	39:CL:60:ASP:OD1	2.06	0.54
40:CM:56:HIS:O	40:CM:58:ASP:N	2.41	0.54
46:CS:40:ASP:O	46:CS:42:ARG:N	2.40	0.54
47:CT:44:ALA:HA	47:CT:71:PHE:O	2.07	0.54
47:CT:65:ILE:HD12	47:CT:65:ILE:N	2.22	0.54
50:CW:22:ARG:O	50:CW:26:ASN:ND2	2.40	0.54
17:D2:49:THR:CB	17:D2:50:PRO:CD	2.86	0.54
55:DA:1313:U:H5''	55:DA:1314:C:OP2	2.07	0.54
55:DA:2168:G:OP1	55:DA:2168:G:C8	2.61	0.54
55:DA:2259:G:H1'	55:DA:2427:C:C2	2.42	0.54
55:DA:2430:A:H8	55:DA:2431:U:H5	1.55	0.54
55:DA:2842:G:O2'	55:DA:2843:G:H5'	2.07	0.54
55:DA:452:G:N3	55:DA:457:A:H2	2.04	0.54
55:DA:546:C:H3'	55:DA:547:A:H8	1.72	0.54
55:DA:55:G:N3	55:DA:127:A:C2	2.75	0.54
55:DA:4:C:C2'	55:DA:5:A:OP2	2.54	0.54
55:DA:810:U:O5'	55:DA:810:U:H6	1.89	0.54
55:DA:7:G:O2'	55:DA:8:A:H5'	2.07	0.54
3:DD:35:LYS:NZ	3:DD:64:ILE:O	2.39	0.54
3:DD:35:LYS:CE	3:DD:64:ILE:C	2.73	0.54
6:DG:77:ILE:CG2	6:DG:80:PHE:H	2.20	0.54
7:DH:154:PRO:HG3	7:DH:163:TYR:CD1	2.42	0.54
56:DJ:14:GLN:N	56:DJ:17:VAL:CG2	2.69	0.54
8:DK:61:ARG:HA	8:DK:61:ARG:CZ	2.37	0.54
58:DL:115:LEU:C	58:DL:115:LEU:HD12	2.27	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:DL:19:PRO:CB	58:DL:25:PRO:HG2	2.36	0.54
9:DM:28:THR:HG22	9:DM:29:LYS:N	2.22	0.54
10:DN:2:ILE:CD1	10:DN:82:ASN:HD22	2.18	0.54
21:DV:72:ARG:HH11	21:DV:72:ARG:HG3	1.71	0.54
24:DW:28:LYS:HD2	24:DW:53:LEU:CD2	2.37	0.54
57:DY:138:LEU:CD1	57:DY:139:VAL:N	2.70	0.54
57:DY:50:ARG:N	57:DY:83:TYR:HB3	2.22	0.54
57:DY:70:GLU:O	57:DY:71:LEU:HG	2.07	0.54
57:DY:7:VAL:CG2	57:DY:8:GLU:N	2.42	0.54
17:A2:66:ARG:HE	17:A2:88:ARG:HD2	1.72	0.54
29:A7:45:ALA:O	29:A7:46:VAL:HG23	2.07	0.54
1:AA:1171:G:HO2'	1:AA:1173:G:C1'	2.20	0.54
1:AA:1345:C:OP2	1:AA:1346:G:OP2	2.25	0.54
1:AA:1786:A:O2'	1:AA:1938:A:N6	2.34	0.54
1:AA:2655:G:N2	1:AA:2665:A:OP2	2.41	0.54
1:AA:2789:C:O2'	1:AA:2790:A:H4'	2.06	0.54
1:AA:323:G:O2'	1:AA:1205:U:N3	2.38	0.54
1:AA:944:G:C2'	1:AA:944:G:N3	2.69	0.54
6:AG:106:LEU:HD12	6:AG:110:ALA:HB3	1.88	0.54
8:AK:54:GLN:HG3	8:AK:55:ALA:N	2.23	0.54
11:AO:57:THR:O	11:AO:59:LEU:N	2.40	0.54
14:AQ:26:LEU:HD22	14:AQ:87:PHE:HD1	1.70	0.54
20:AU:28:LYS:HA	20:AU:28:LYS:HE2	1.88	0.54
20:AU:63:LYS:HA	20:AU:63:LYS:HZ2	1.71	0.54
21:AV:115:GLY:CA	21:AV:175:VAL:O	2.54	0.54
24:AW:46:GLN:HB2	24:AW:49:LYS:CE	2.36	0.54
53:B1:30:C:O2'	53:B1:31:A:H5'	2.08	0.54
31:BA:1084:G:OP1	31:BA:1086:U:C2	2.61	0.54
31:BA:967:C:H2'	31:BA:968:A:N7	2.22	0.54
52:BB:19:G:N2	52:BB:56:C:N3	2.55	0.54
36:BI:19:LEU:HD23	36:BI:19:LEU:O	2.08	0.54
38:BK:104:ARG:O	38:BK:105:ARG:C	2.46	0.54
42:BO:5:PRO:HG2	42:BO:10:LEU:HD21	1.89	0.54
26:A4:52:THR:CG2	43:BP:65:LYS:HD3	2.32	0.54
54:CA:1200:C:O2'	54:CA:1201:A:OP2	2.25	0.54
54:CA:1504:G:O2'	54:CA:1505:G:P	2.65	0.54
54:CA:341:C:H2'	54:CA:342:C:C6	2.43	0.54
54:CA:476:G:O2'	54:CA:477:G:H5'	2.07	0.54
54:CA:50:A:H4'	54:CA:51:A:H5'	1.88	0.54
54:CA:722:A:H3'	54:CA:722:A:N3	2.22	0.54
52:CD:58:A:N6	52:CD:61:C:C6	2.75	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:CH:83:GLU:HG2	35:CH:88:LYS:HB2	1.88	0.54
36:CI:24:GLU:HA	36:CI:27:GLN:HG2	1.89	0.54
37:CJ:32:ARG:O	37:CJ:33:ASP:HB2	2.07	0.54
54:CA:689:C:P	41:CN:46:GLY:HA3	2.48	0.54
43:CP:7:VAL:HB	6:DG:115:ARG:NH2	2.23	0.54
46:CS:81:ARG:HH11	46:CS:81:ARG:HB3	1.70	0.54
49:CV:5:LEU:H	49:CV:5:LEU:HD12	1.71	0.54
4:DE:111:ARG:HA	13:D0:1:MET:SD	2.48	0.54
28:D6:25:LYS:HD2	30:D8:34:TRP:CZ2	2.43	0.54
28:D6:40:CYS:SG	28:D6:45:LYS:CD	2.96	0.54
55:DA:1155:A:H4'	16:D1:55:ARG:NH1	2.22	0.54
55:DA:1181:C:O2'	55:DA:1182:A:H5'	2.06	0.54
55:DA:699:A:H4'	55:DA:1634:A:N7	2.22	0.54
55:DA:163:U:C2'	55:DA:164:U:H5'	2.37	0.54
55:DA:2219:G:C2'	55:DA:2224:G:H5'	2.37	0.54
55:DA:2749:A:N1	55:DA:2750:A:N6	2.54	0.54
55:DA:2832:U:C4'	55:DA:2833:G:H5''	2.32	0.54
55:DA:558:G:P	9:DM:111:PRO:HD2	2.47	0.54
55:DA:654(N):G:H8	55:DA:654(N):G:OP1	1.91	0.54
55:DA:780:G:N2	55:DA:783:A:H62	2.06	0.54
3:DD:168:ARG:HG3	3:DD:168:ARG:HH11	1.72	0.54
3:DD:182:LEU:N	3:DD:272:ALA:HB3	2.21	0.54
4:DE:128:SER:OG	4:DE:129:HIS:N	2.38	0.54
5:DF:132:VAL:HG23	5:DF:133:ASN:ND2	2.22	0.54
8:DK:88:ILE:HB	8:DK:90:GLY:O	2.08	0.54
58:DL:52:ILE:HD11	58:DL:76:TYR:CA	2.37	0.54
11:DO:14:LYS:O	11:DO:16:ARG:N	2.40	0.54
11:DO:34:GLY:O	11:DO:36:LYS:N	2.40	0.54
11:DO:70:GLN:N	11:DO:70:GLN:CD	2.61	0.54
12:DP:47:ILE:HD12	12:DP:70:PRO:HD3	1.90	0.54
55:DA:2875:C:C4'	15:DR:5:ALA:HB2	2.37	0.54
57:DY:26:LEU:HB3	57:DY:112:LEU:HD13	1.89	0.54
13:A0:28:LEU:HD13	13:A0:28:LEU:O	2.07	0.54
28:A6:34:LEU:CD2	28:A6:34:LEU:H	2.20	0.54
1:AA:1856:G:C2'	1:AA:1857:G:H5'	2.37	0.54
1:AA:2389:G:H5''	1:AA:2390:U:H5'	1.90	0.54
1:AA:2531:A:H3'	1:AA:2532:G:C8	2.43	0.54
1:AA:2623:G:H4'	1:AA:2825:C:O2	2.08	0.54
1:AA:67:U:H2'	1:AA:68:G:H8	1.72	0.54
2:AB:55:U:O2'	2:AB:56:G:H5'	2.07	0.54
5:AF:4:VAL:HG11	5:AF:17:ARG:HH21	1.73	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:AF:9:ILE:CG1	5:AF:15:SER:HA	2.37	0.54
8:AK:56:LYS:HD2	8:AK:56:LYS:C	2.27	0.54
8:AK:75:LEU:HD22	8:AK:77:LEU:CD2	2.35	0.54
9:AM:98:VAL:HG23	9:AM:99:LEU:N	2.23	0.54
1:AA:2415:G:H4'	11:AO:67:MET:N	2.21	0.54
11:AO:88:LEU:HD11	11:AO:95:VAL:CG2	2.36	0.54
12:AP:99:PRO:HG3	21:AV:79:ARG:HH12	1.70	0.54
14:AQ:106:ARG:HH11	14:AQ:106:ARG:CB	2.20	0.54
14:AQ:18:ILE:CD1	14:AQ:88:ASP:HA	2.38	0.54
15:AR:29:ARG:HH11	15:AR:29:ARG:CG	2.16	0.54
15:AR:45:PHE:CD1	15:AR:65:LYS:HE2	2.42	0.54
20:AU:35:TYR:CD1	20:AU:69:ALA:HB3	2.43	0.54
21:AV:102:LEU:HD21	21:AV:124:ILE:CG2	2.37	0.54
21:AV:12:GLY:C	21:AV:13:GLU:HG3	2.27	0.54
31:BA:1137:C:H4'	31:BA:1138:G:C2	2.43	0.54
31:BA:1160:G:H2'	31:BA:1160:G:N3	2.21	0.54
31:BA:1319:A:H2'	31:BA:1323:G:C8	2.43	0.54
32:BE:91:PRO:HA	32:BE:154:LEU:HD12	1.88	0.54
32:BE:236:TYR:HA	32:BE:239:VAL:HG23	1.89	0.54
33:BF:119:ARG:HA	33:BF:122:GLU:OE2	2.07	0.54
35:BH:111:GLU:C	35:BH:113:ALA:H	2.10	0.54
36:BI:8:ILE:HG22	36:BI:10:LEU:CD1	2.38	0.54
31:BA:1151:A:H1'	40:BM:39:PRO:CB	2.37	0.54
43:BP:27:LYS:CE	43:BP:31:LYS:HE3	2.34	0.54
54:CA:1032(A):G:H2'	54:CA:1032(B):G:H8	1.71	0.54
54:CA:1074:G:C4'	32:CE:104:ASN:HB2	2.38	0.54
54:CA:949:A:O4'	54:CA:1364:U:O4	2.25	0.54
54:CA:1376:U:H2'	54:CA:1377:A:C8	2.42	0.54
54:CA:430:A:C2'	54:CA:431:A:H5'	2.37	0.54
54:CA:437:U:H2'	54:CA:438:G:O4'	2.07	0.54
54:CA:579:G:H5'	54:CA:728:A:C1'	2.29	0.54
54:CA:779:C:H2'	54:CA:780:A:O4'	2.08	0.54
54:CA:986:A:H2'	54:CA:987:G:O4'	2.07	0.54
54:CA:991:U:O4	54:CA:1212:U:O2'	2.26	0.54
32:CE:97:TRP:HZ3	32:CE:172:ILE:HG22	1.72	0.54
34:CG:116:GLN:NE2	34:CG:157:LEU:HD21	2.22	0.54
34:CG:9:CYS:C	34:CG:11:LEU:H	2.09	0.54
42:CO:53:ARG:HH12	42:CO:92:ASP:CB	2.21	0.54
43:CP:90:LEU:C	43:CP:90:LEU:HD12	2.26	0.54
46:CS:68:ASP:O	46:CS:70:ALA:N	2.40	0.54
50:CW:57:ARG:HD3	50:CW:102:GLY:O	2.07	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:D1:96:ALA:C	16:D1:98:LEU:N	2.57	0.54
26:D4:49:PHE:O	26:D4:50:VAL:CB	2.55	0.54
55:DA:2016:U:H1'	27:D5:6:VAL:HG13	1.90	0.54
28:D6:15:GLU:CD	28:D6:44:ARG:NH2	2.61	0.54
55:DA:1444:G:H2'	55:DA:1445:C:C5	2.43	0.54
55:DA:1535:U:H3'	55:DA:1536:A:H5''	1.90	0.54
55:DA:1291:C:O4'	55:DA:1536:A:OP2	2.25	0.54
55:DA:1537:C:H2'	55:DA:1538:G:O4'	2.07	0.54
55:DA:1547:C:H2'	55:DA:1548:C:H6	1.72	0.54
55:DA:200:U:H2'	55:DA:201:C:H5'	1.88	0.54
55:DA:2051:A:OP2	55:DA:2051:A:H8	1.90	0.54
55:DA:2691:C:H6	55:DA:2691:C:H5'	1.73	0.54
55:DA:2849:U:C2'	55:DA:2866:U:O2	2.56	0.54
55:DA:372:G:O2'	55:DA:373:U:OP2	2.25	0.54
55:DA:654(D):G:H2'	55:DA:654(D):G:N3	2.22	0.54
55:DA:662:G:H5''	11:DO:15:ARG:O	2.07	0.54
3:DD:123:ALA:HB3	3:DD:131:LEU:HG	1.89	0.54
3:DD:27:THR:CG2	3:DD:83:GLU:HB3	2.33	0.54
3:DD:91:ARG:O	3:DD:107:ALA:HB3	2.07	0.54
4:DE:70:ALA:O	4:DE:71:GLY:C	2.45	0.54
8:DK:116:LEU:O	8:DK:116:LEU:HG	2.08	0.54
58:DL:65:PHE:O	58:DL:65:PHE:CD2	2.60	0.54
11:DO:138:LEU:C	11:DO:140:ALA:N	2.60	0.54
12:DP:87:LYS:C	12:DP:89:ASN:N	2.57	0.54
26:A4:25:TYR:O	26:A4:26:SER:C	2.45	0.54
26:A4:1:MET:O	26:A4:2:LYS:HD3	2.07	0.54
1:AA:2393:A:C5'	30:A8:30:ARG:HD3	2.37	0.54
30:A8:50:LEU:CD1	30:A8:54:GLU:HA	2.37	0.54
1:AA:1231:G:O2'	1:AA:1232:G:H5'	2.07	0.54
1:AA:1278:A:O2'	13:A0:34:ILE:HD11	2.07	0.54
1:AA:1381:G:H2'	1:AA:1382:G:H5'	1.90	0.54
1:AA:1827:C:C3'	1:AA:1828:G:H5'	2.38	0.54
1:AA:9:U:H3	1:AA:2629:A:H61	1.53	0.54
1:AA:709:U:H2'	1:AA:710:G:C8	2.43	0.54
1:AA:863:A:O2'	1:AA:864:G:H5'	2.07	0.54
1:AA:863:A:H2'	1:AA:864:G:C8	2.41	0.54
3:AD:69:ARG:NH1	3:AD:128:GLY:O	2.40	0.54
4:AE:27:LEU:HD12	4:AE:180:ASN:O	2.07	0.54
5:AF:192:LEU:HD23	5:AF:192:LEU:C	2.28	0.54
7:AH:80:SER:O	7:AH:81:GLU:HB2	2.08	0.54
8:AK:56:LYS:HG3	8:AK:57:ARG:N	2.23	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:AM:60:ILE:HD13	9:AM:99:LEU:HD23	1.89	0.54
11:AO:114:ILE:O	11:AO:114:ILE:HG13	2.07	0.54
12:AP:133:ARG:HD3	12:AP:133:ARG:C	2.27	0.54
15:AR:106:SER:O	15:AR:107:ASP:CB	2.55	0.54
15:AR:106:SER:O	15:AR:107:ASP:HB3	2.08	0.54
15:AR:74:ARG:HG2	15:AR:74:ARG:NH1	2.22	0.54
19:AT:36:LYS:HG2	19:AT:56:THR:HG23	1.88	0.54
21:AV:106:GLY:O	21:AV:108:PRO:HD2	2.04	0.54
21:AV:147:GLY:O	21:AV:148:ASP:C	2.44	0.54
21:AV:184:ALA:C	21:AV:186:GLU:N	2.60	0.54
31:BA:1238:A:N7	31:BA:1301:U:O4	2.41	0.54
31:BA:1221:G:OP1	31:BA:1321:C:N4	2.41	0.54
31:BA:1362(A):C:H5'	31:BA:1363:A:O5'	2.08	0.54
31:BA:201:C:C4	31:BA:209:U:H6	2.25	0.54
31:BA:986:A:H2'	31:BA:987:G:H8	1.69	0.54
31:BA:1104:G:H4'	32:BE:111:ARG:CZ	2.38	0.54
32:BE:134:GLU:O	32:BE:138:LEU:HG	2.07	0.54
32:BE:22:LYS:NZ	32:BE:40:HIS:CE1	2.75	0.54
33:BF:59:ARG:NE	33:BF:64:VAL:HG22	2.21	0.54
33:BF:76:VAL:CG2	33:BF:77:ILE:N	2.70	0.54
50:BW:57:ARG:HD3	50:BW:102:GLY:O	2.06	0.54
54:CA:1363:A:C4'	54:CA:1364:U:OP1	2.55	0.54
54:CA:1410:G:H2'	54:CA:1411:C:C6	2.43	0.54
54:CA:1424:C:H2'	54:CA:1425:U:C6	2.43	0.54
54:CA:776:G:N2	54:CA:802:A:OP2	2.39	0.54
54:CA:91:C:H2'	54:CA:92:G:H5''	1.89	0.54
52:CB:58:A:O2'	52:CB:59:U:O5'	2.20	0.54
32:CE:167:PRO:HG3	32:CE:188:ALA:HB2	1.88	0.54
35:CH:72:GLN:HE21	35:CH:144:THR:HG22	1.71	0.54
38:CK:82:HIS:HD2	38:CK:83:ILE:N	2.06	0.54
41:CN:85:ARG:HG2	41:CN:111:ASP:O	2.07	0.54
43:CP:25:ILE:HD11	43:CP:66:LEU:CD2	2.38	0.54
46:CS:28:ARG:NH1	46:CS:28:ARG:HG2	2.20	0.54
55:DA:1049:C:C6	55:DA:1049:C:H5'	2.43	0.54
55:DA:1142(A):A:HO2'	55:DA:1143:A:P	2.31	0.54
55:DA:49:A:C8	55:DA:120:U:H5	2.21	0.54
55:DA:811:U:HO2'	55:DA:1250:G:H2'	1.73	0.54
55:DA:1803:A:H4'	3:DD:259:THR:HG23	1.89	0.54
55:DA:2238:G:H5'	55:DA:2239:G:N7	2.23	0.54
55:DA:2341:G:H2'	55:DA:2342:C:H6	1.71	0.54
55:DA:2591:C:P	3:DD:239:ARG:HG3	2.48	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2864:G:OP1	15:DR:119:LYS:HD2	2.07	0.54
55:DA:613:U:O4'	55:DA:613:U:O2	2.21	0.54
55:DA:84:A:C4'	55:DA:85:G:O5'	2.50	0.54
55:DA:950:G:H2'	55:DA:951:C:C6	2.42	0.54
3:DD:110:GLY:O	3:DD:111:LEU:C	2.45	0.54
3:DD:11:PRO:O	3:DD:12:SER:OG	2.18	0.54
7:DH:30:LYS:HE3	7:DH:81:GLU:CA	2.37	0.54
8:DK:52:ARG:HG3	8:DK:53:ALA:N	2.22	0.54
55:DA:1081:U:O2'	58:DL:126:MET:SD	2.65	0.54
58:DL:25:PRO:C	58:DL:27:LEU:H	2.11	0.54
11:DO:125:VAL:O	11:DO:145:PRO:HD2	2.07	0.54
11:DO:31:ALA:O	11:DO:32:THR:CG2	2.55	0.54
15:DR:50:ILE:HD11	15:DR:102:ILE:CD1	2.31	0.54
21:DV:162:GLU:HG2	21:DV:163:LEU:N	2.20	0.54
57:DY:69:PRO:O	57:DY:113:GLN:HB2	2.07	0.54
23:DZ:80:LEU:C	23:DZ:81:LYS:HE2	2.27	0.54
13:A0:41:ALA:O	13:A0:43:GLU:N	2.40	0.54
1:AA:142:G:H4'	19:AT:35:THR:HG21	1.87	0.54
1:AA:1885:A:C5'	1:AA:1886:C:OP2	2.55	0.54
1:AA:2275:C:O2'	1:AA:2276:G:OP2	2.24	0.54
1:AA:270:A:O2'	1:AA:270(A):A:H5'	2.08	0.54
1:AA:858:U:C2	1:AA:2268:A:C2	2.95	0.54
1:AA:893:C:N3	1:AA:894:C:N4	2.56	0.54
2:AB:31:C:H4'	6:AG:29:TRP:HH2	1.73	0.54
3:AD:24:ILE:O	3:AD:24:ILE:HG23	2.07	0.54
1:AA:1568:G:P	3:AD:63:ARG:HH12	2.30	0.54
4:AE:154:LYS:HA	4:AE:154:LYS:CE	2.34	0.54
5:AF:29:ASN:N	5:AF:112:MET:CE	2.71	0.54
8:AK:56:LYS:HE2	8:AK:57:ARG:HG2	1.89	0.54
10:AN:71:ARG:NE	10:AN:105:GLU:OE2	2.41	0.54
10:AN:91:LEU:HD22	10:AN:91:LEU:N	2.23	0.54
11:AO:117:GLU:N	11:AO:117:GLU:OE1	2.38	0.54
11:AO:62:LEU:C	11:AO:62:LEU:HD22	2.27	0.54
12:AP:137:TYR:O	12:AP:138:ASP:O	2.25	0.54
18:AS:71:VAL:HA	18:AS:107:LEU:HD12	1.90	0.54
20:AU:27:VAL:HG12	20:AU:39:VAL:HG12	1.89	0.54
21:AV:115:GLY:O	21:AV:174:VAL:HG13	2.08	0.54
53:B1:28:G:H2'	53:B1:29:G:C8	2.42	0.54
31:BA:1216:G:H5''	44:BQ:5:ALA:CB	2.37	0.54
31:BA:132:C:O3'	50:BW:74:LYS:HE2	2.06	0.54
31:BA:1363:A:H1'	31:BA:1365:G:C8	2.41	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:345:C:H1'	31:BA:346:G:C2	2.41	0.54
31:BA:464:G:N1	31:BA:467:G:OP2	2.36	0.54
31:BA:946:A:O2'	31:BA:1333:A:H1'	2.07	0.54
33:BF:15:THR:HG22	33:BF:16:ARG:N	2.23	0.54
35:BH:70:PRO:O	35:BH:71:LEU:HD23	2.08	0.54
42:BO:28:LYS:O	42:BO:30:ALA:N	2.40	0.54
48:BU:36:ASN:HD22	48:BU:39:VAL:HG21	1.72	0.54
54:CA:1067:A:H4'	54:CA:1068:G:O5'	2.06	0.54
54:CA:1499:A:C1'	54:CA:1520:G:H5'	2.34	0.54
54:CA:645:C:H2'	54:CA:646:U:C6	2.43	0.54
54:CA:655:A:C2	54:CA:754:C:N4	2.76	0.54
54:CA:837:G:O2'	54:CA:838:G:H5'	2.08	0.54
54:CA:869:G:C4'	54:CA:872:A:H1'	2.37	0.54
33:CF:124:ILE:C	33:CF:126:ARG:H	2.09	0.54
34:CG:25:ARG:O	34:CG:27:TYR:N	2.29	0.54
36:CI:75:LEU:C	36:CI:75:LEU:HD23	2.28	0.54
38:CK:104:ARG:C	38:CK:106:GLY:H	2.11	0.54
39:CL:17:VAL:HG13	39:CL:81:ILE:HD13	1.90	0.54
33:CF:23:TYR:HA	40:CM:11:PHE:CE1	2.42	0.54
40:CM:29:ARG:HG2	40:CM:29:ARG:O	2.07	0.54
54:CA:972:C:OP2	40:CM:57:LYS:HE2	2.08	0.54
40:CM:51:ARG:NE	40:CM:60:ARG:O	2.40	0.54
50:CW:33:ILE:HD12	50:CW:63:ILE:HG12	1.89	0.54
16:D1:92:ARG:O	16:D1:92:ARG:CG	2.55	0.54
22:D3:23:VAL:HG12	22:D3:25:ARG:H	1.71	0.54
12:DP:80:GLU:HA	22:D3:4:LYS:HE3	1.88	0.54
55:DA:1329:U:H5''	55:DA:1330:C:H5	1.73	0.54
55:DA:2131:G:OP1	55:DA:2132:U:H3'	2.08	0.54
55:DA:2290:G:H8	55:DA:2290:G:H5'	1.71	0.54
55:DA:2481:G:HO2'	55:DA:2482:G:P	2.30	0.54
55:DA:2675:A:H5'	55:DA:2676:C:OP2	2.08	0.54
55:DA:439:G:O2'	55:DA:440:G:H5'	2.07	0.54
55:DA:803:U:H2'	55:DA:804:A:C5'	2.38	0.54
55:DA:805:G:C4'	55:DA:806:C:OP2	2.55	0.54
55:DA:897:C:H6	55:DA:897:C:C5'	2.20	0.54
58:DL:46:ALA:O	58:DL:47:ASN:C	2.45	0.54
55:DA:637:A:O5'	11:DO:116:GLY:HA2	2.07	0.54
12:DP:23:GLY:O	12:DP:24:GLY:C	2.46	0.54
14:DQ:18:ILE:O	14:DQ:19:LYS:O	2.24	0.54
19:DT:31:HIS:NE2	19:DT:33:LYS:HB2	2.22	0.54
20:DU:49:VAL:HG12	20:DU:50:ARG:N	2.22	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:DU:57:GLN:HE21	20:DU:58:GLY:N	2.06	0.54
20:DU:75:ILE:CG1	20:DU:80:GLY:H	2.21	0.54
21:DV:177:PRO:CG	21:DV:177:PRO:O	2.55	0.54
13:A0:97:VAL:HG13	13:A0:114:VAL:HG22	1.88	0.54
26:A4:59:PHE:HB3	26:A4:60:GLN:HE22	1.73	0.54
29:A7:12:ARG:CD	29:A7:46:VAL:HG21	2.31	0.54
30:A8:6:THR:O	30:A8:7:HIS:HB3	2.07	0.54
1:AA:1098:A:C3'	1:AA:1099:G:H5''	2.36	0.54
1:AA:1448:G:H2'	1:AA:1449:A:C8	2.42	0.54
1:AA:1751:C:O2'	1:AA:1752:C:H5'	2.07	0.54
1:AA:236:C:H2'	1:AA:237:C:H6	1.73	0.54
1:AA:556:G:H8	1:AA:556:G:O5'	1.91	0.54
1:AA:654(I):C:C2	1:AA:654(J):A:C8	2.95	0.54
3:AD:94:LEU:HD23	3:AD:95:LEU:N	2.23	0.54
4:AE:3:GLY:HA3	4:AE:81:ILE:HD12	1.90	0.54
7:AH:121:ILE:HG22	7:AH:122:THR:N	2.23	0.54
11:AO:146:VAL:HG22	11:AO:147:LEU:N	2.15	0.54
12:AP:78:PRO:O	12:AP:79:LEU:HB2	2.07	0.54
31:BA:1126:U:O2'	31:BA:1127:G:P	2.65	0.54
31:BA:1534:A:H2'	31:BA:1535:C:C5	2.43	0.54
31:BA:508:C:H5''	31:BA:509:A:OP1	2.07	0.54
52:BB:11:C:H2'	52:BB:12:U:H6	1.72	0.54
32:BE:121:LEU:HD22	32:BE:127:ILE:HD13	1.90	0.54
32:BE:178:ARG:HB2	32:BE:178:ARG:HH11	1.73	0.54
32:BE:36:ARG:H	32:BE:41:ILE:HD13	1.72	0.54
33:BF:141:VAL:HG11	33:BF:202:ILE:HD12	1.89	0.54
36:BI:14:LEU:CD2	36:BI:18:GLN:HB2	2.37	0.54
37:BJ:113:GLU:HB2	37:BJ:119:ARG:CG	2.33	0.54
31:BA:1347:G:C8	39:BL:107:ARG:HB3	2.42	0.54
40:BM:65:LEU:HA	44:BQ:55:GLY:O	2.07	0.54
42:BO:6:THR:N	42:BO:9:GLN:HE21	2.05	0.54
43:BP:97:PRO:N	43:BP:110:ARG:HD3	2.23	0.54
44:BQ:14:PRO:CG	44:BQ:15:LYS:H	2.18	0.54
54:CA:1052:U:H2'	54:CA:1055:A:OP1	2.07	0.54
54:CA:1322:C:O2'	54:CA:1323:G:C5'	2.53	0.54
54:CA:181:G:O2'	54:CA:182:U:C6	2.60	0.54
54:CA:448:A:C2	54:CA:487:A:C2	2.95	0.54
54:CA:95:G:C2'	54:CA:96:G:H5''	2.38	0.54
32:CE:92:TYR:CE2	32:CE:151:GLY:HA3	2.41	0.54
32:CE:54:THR:CG2	32:CE:201:ILE:HD11	2.29	0.54
37:CJ:54:THR:O	37:CJ:56:GLN:N	2.41	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:CH:78:HIS:HB3	38:CK:107:LEU:HD12	1.89	0.54
40:CM:80:LYS:HB2	40:CM:80:LYS:NZ	2.23	0.54
41:CN:80:VAL:HG13	41:CN:103:LEU:HD12	1.90	0.54
42:CO:62:SER:HB2	42:CO:64:TYR:CD1	2.42	0.54
54:CA:377:G:OP1	46:CS:3:LYS:HD2	2.06	0.54
47:CT:24:GLU:HG2	47:CT:39:SER:HB3	1.89	0.54
47:CT:46:ASP:OD2	47:CT:51:TYR:HD1	1.91	0.54
49:CV:80:TYR:CE2	49:CV:81:ARG:O	2.61	0.54
49:CV:80:TYR:CD2	49:CV:81:ARG:O	2.60	0.54
54:CA:1305:G:H5''	51:CX:4:GLY:C	2.28	0.54
55:DA:2839:G:H5''	13:D0:46:GLY:HA2	1.90	0.54
17:D2:25:LEU:CD1	17:D2:94:LEU:HD21	2.38	0.54
55:DA:1155:A:H4'	16:D1:55:ARG:HH12	1.73	0.54
55:DA:1547:C:H2'	55:DA:1548:C:C6	2.43	0.54
55:DA:2111:C:C2	55:DA:2118:U:O2'	2.61	0.54
55:DA:2439:A:H5'	55:DA:2439:A:C8	2.43	0.54
55:DA:2524:G:H2'	55:DA:2741:A:H2	1.72	0.54
55:DA:2531:A:H2'	55:DA:2531:A:N3	2.22	0.54
55:DA:2884:U:H2'	55:DA:2885:C:H5'	1.88	0.54
55:DA:556:G:H2'	55:DA:557:U:H6	1.73	0.54
3:DD:134:ARG:HB2	3:DD:135:PHE:CE2	2.43	0.54
55:DA:323:G:H2'	5:DF:169:ASN:ND2	2.23	0.54
6:DG:13:GLU:HG3	6:DG:13:GLU:O	2.08	0.54
6:DG:77:ILE:O	6:DG:81:LYS:O	2.25	0.54
58:DL:16:LYS:O	58:DL:17:ALA:HB2	2.08	0.54
9:DM:62:VAL:CG1	9:DM:66:LYS:HD2	2.37	0.54
11:DO:95:VAL:HG13	11:DO:100:LEU:HD21	1.89	0.54
15:DR:96:ARG:CZ	15:DR:96:ARG:HB2	2.38	0.54
21:DV:128:VAL:HG23	21:DV:161:VAL:HG23	1.90	0.54
55:DA:1086:A:H2	57:DY:41:ARG:NH2	2.03	0.54
57:DY:7:VAL:CG1	57:DY:8:GLU:H	2.05	0.54
13:A0:33:ARG:HG2	13:A0:115:GLU:CG	2.38	0.54
6:AG:101:ILE:HB	26:A4:25:TYR:HD2	1.73	0.54
30:A8:33:ASN:O	30:A8:34:TRP:HB3	2.06	0.54
30:A8:50:LEU:CG	30:A8:51:ALA:N	2.59	0.54
1:AA:1406:U:H2'	1:AA:1406:U:O2	2.07	0.54
1:AA:1543:A:O2'	1:AA:1544:C:H3'	2.06	0.54
1:AA:1657:C:H2'	1:AA:1658:C:H6	1.71	0.54
1:AA:1928:A:C3'	1:AA:1929:G:C5'	2.85	0.54
1:AA:2245:U:C5'	1:AA:2246:G:H5'	2.36	0.54
1:AA:2537:U:H2'	1:AA:2538:C:H6	1.73	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2702:U:HO2'	1:AA:2703:C:H6	1.50	0.54
1:AA:2790:A:O2'	1:AA:2791:C:P	2.66	0.54
1:AA:2852:G:H2'	1:AA:2853:C:C6	2.43	0.54
1:AA:313:C:O2'	1:AA:314:A:H5'	2.08	0.54
1:AA:947:G:H2'	1:AA:948:G:H8	1.72	0.54
2:AB:24:G:H5''	2:AB:25:A:OP1	2.07	0.54
3:AD:117:VAL:HG22	3:AD:118:VAL:H	1.72	0.54
1:AA:1693:U:O2	3:AD:14:ARG:NH1	2.41	0.54
5:AF:132:VAL:O	5:AF:134:GLY:N	2.31	0.54
7:AH:94:TYR:CE2	7:AH:153:LYS:HE2	2.43	0.54
8:AK:133:HIS:ND1	8:AK:134:PRO:HD3	2.22	0.54
8:AK:88:ILE:CG2	8:AK:89:TYR:N	2.64	0.54
11:AO:105:LEU:HD12	11:AO:105:LEU:N	2.23	0.54
12:AP:34:LEU:HD13	12:AP:118:LEU:HB3	1.90	0.54
14:AQ:78:LEU:HD11	14:AQ:107:GLU:O	2.07	0.54
15:AR:98:LYS:HB3	15:AR:100:TYR:CE1	2.43	0.54
20:AU:14:LEU:H	20:AU:14:LEU:HD23	1.72	0.54
20:AU:81:LYS:HB2	20:AU:96:ILE:CG2	2.38	0.54
24:AW:16:LEU:CD1	24:AW:16:LEU:O	2.55	0.54
23:AZ:91:LYS:CA	23:AZ:91:LYS:HE3	2.30	0.54
31:BA:1169:A:H2'	31:BA:1170:A:C8	2.43	0.54
31:BA:1326:C:O2'	31:BA:1327:C:H5'	2.08	0.54
31:BA:1346:A:O2'	31:BA:1347:G:OP2	2.26	0.54
31:BA:246:A:O2'	31:BA:247:G:O5'	2.26	0.54
31:BA:30:U:O2'	31:BA:31:G:OP1	2.21	0.54
31:BA:31:G:O2'	31:BA:32:A:P	2.65	0.54
31:BA:707:C:H2'	31:BA:708:C:C6	2.41	0.54
31:BA:930:C:O2'	31:BA:931:C:H5'	2.08	0.54
32:BE:87:ARG:NH1	32:BE:219:VAL:HG13	2.22	0.54
33:BF:149:ALA:O	33:BF:169:ALA:HB1	2.08	0.54
33:BF:60:ALA:O	33:BF:61:ALA:HB2	2.08	0.54
34:BG:49:ARG:NE	34:BG:49:ARG:HA	2.23	0.54
35:BH:63:ARG:HA	35:BH:66:MET:HE1	1.90	0.54
34:BG:88:VAL:HG13	35:BH:97:GLY:HA2	1.90	0.54
36:BI:60:PHE:C	36:BI:61:LEU:HD12	2.28	0.54
37:BJ:131:LYS:NZ	37:BJ:131:LYS:HB2	2.22	0.54
47:BT:80:GLY:O	47:BT:81:ARG:HB2	2.08	0.54
54:CA:1118:C:H1'	54:CA:1179:A:C4	2.43	0.54
54:CA:1176:A:H2'	54:CA:1177:G:C5'	2.37	0.54
54:CA:161:A:H2'	54:CA:162:A:H8	1.72	0.54
54:CA:46:G:O2'	54:CA:365:U:H1'	2.08	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:CK:11:THR:HG23	38:CK:14:ARG:HH12	1.71	0.54
37:CJ:16:LEU:HD11	39:CL:45:ALA:HB2	1.90	0.54
54:CA:625:G:H5'	46:CS:10:GLY:HA2	1.89	0.54
54:CA:1453:G:H2'	50:CW:39:LYS:HZ1	1.73	0.54
26:D4:1:MET:HB3	26:D4:6:HIS:CD2	2.43	0.54
29:D7:30:VAL:HG12	29:D7:31:LEU:N	2.22	0.54
55:DA:1204:A:H1'	55:DA:1206:G:C5	2.42	0.54
55:DA:1716:U:O2'	55:DA:1717:G:H5'	2.07	0.54
55:DA:1550:C:H4'	55:DA:1734:C:O2	2.08	0.54
55:DA:2064:C:H2'	55:DA:2065:C:H6	1.73	0.54
55:DA:2405:G:O2'	55:DA:2406:U:OP2	2.24	0.54
55:DA:439:G:H2'	55:DA:440:G:C8	2.43	0.54
55:DA:587:C:O2'	55:DA:588:U:OP2	2.23	0.54
3:DD:158:ALA:O	3:DD:159:ALA:C	2.46	0.54
3:DD:166:GLN:CA	3:DD:166:GLN:NE2	2.71	0.54
6:DG:68:PRO:CG	6:DG:90:LEU:HD12	2.37	0.54
7:DH:35:VAL:CG1	7:DH:71:LEU:HG	2.36	0.54
56:DI:24:ILE:HD11	56:DI:25:ASP:HB2	1.90	0.54
58:DL:34:ILE:HG13	58:DL:38:VAL:HG22	1.90	0.54
58:DL:80:LYS:C	58:DL:80:LYS:HD2	2.27	0.54
11:DO:38:GLN:N	11:DO:41:ARG:HG2	2.21	0.54
12:DP:35:VAL:HG11	12:DP:130:LYS:HD2	1.90	0.54
19:DT:26:TYR:OH	19:DT:88:LYS:HB2	2.07	0.54
21:DV:76:LEU:HD23	21:DV:76:LEU:N	2.23	0.54
57:DY:93:LEU:HG	57:DY:126:ALA:O	2.06	0.54
1:AA:1099:G:H2'	1:AA:1100:C:O4'	2.08	0.54
1:AA:1128:A:O2'	1:AA:1129:A:O4'	2.23	0.54
1:AA:1278:A:O2'	1:AA:1279:G:H5'	2.08	0.54
1:AA:1567:A:OP2	3:AD:86:PRO:HB3	2.07	0.54
1:AA:2018:G:H2'	1:AA:2019:A:C8	2.43	0.54
1:AA:2311:A:O2'	1:AA:2312:U:H5'	2.08	0.54
1:AA:2656:U:C5	1:AA:2657:A:N7	2.76	0.54
1:AA:2807:G:O6	1:AA:2893:G:O6	2.25	0.54
1:AA:566:U:H2'	1:AA:567:A:O4'	2.08	0.54
1:AA:654(M):C:C2'	1:AA:654(N):G:OP1	2.55	0.54
1:AA:654(N):G:H8	1:AA:654(N):G:OP1	1.91	0.54
1:AA:89:G:OP2	1:AA:90:U:C6	2.60	0.54
4:AE:55:ASN:C	4:AE:57:LYS:H	2.11	0.54
6:AG:144:ILE:HG22	6:AG:146:TYR:H	1.73	0.54
2:AB:54:G:H21	6:AG:29:TRP:HZ2	1.56	0.54
1:AA:270(L):U:O4	8:AK:50:ARG:NH1	2.41	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:AQ:86:ALA:O	14:AQ:87:PHE:HB2	2.08	0.54
21:AV:58:VAL:C	21:AV:60:GLU:H	2.11	0.54
53:B1:35:A:H2'	53:B1:36:G:H8	1.72	0.54
31:BA:1004:A:C4	31:BA:1025:U:C2	2.96	0.54
31:BA:1062:U:H2'	31:BA:1063:C:C5	2.42	0.54
31:BA:1182:G:H4'	31:BA:1183:A:H5''	1.90	0.54
31:BA:186(C):G:H2'	31:BA:186(D):C:H6	1.72	0.54
31:BA:328:C:C2'	31:BA:328:C:O2	2.56	0.54
31:BA:794:A:O2'	31:BA:795:C:H5'	2.08	0.54
31:BA:841:U:H4'	31:BA:842:C:C5	2.43	0.54
52:BC:18:G:H22	52:BC:57:G:H2'	1.73	0.54
32:BE:42:ILE:HD11	32:BE:202:PRO:HB2	1.89	0.54
33:BF:92:ALA:HB2	33:BF:99:VAL:CG2	2.38	0.54
31:BA:1372:U:OP2	39:BL:11:LYS:NZ	2.39	0.54
40:BM:27:ALA:HB2	40:BM:85:LEU:HD11	1.90	0.54
40:BM:49:VAL:HG12	40:BM:50:ILE:N	2.23	0.54
42:BO:68:ALA:HB1	42:BO:100:ILE:HG13	1.89	0.54
42:BO:68:ALA:HA	42:BO:98:TYR:O	2.07	0.54
54:CA:1306:A:N6	54:CA:1331:G:H1'	2.22	0.54
33:CF:15:THR:HG21	33:CF:181:ASN:HA	1.88	0.54
34:CG:146:ILE:N	34:CG:146:ILE:HD12	2.23	0.54
36:CI:63:TYR:CD2	36:CI:63:TYR:N	2.76	0.54
38:CK:20:TYR:CE1	38:CK:78:GLN:NE2	2.76	0.54
40:CM:4:ILE:CG2	40:CM:74:ILE:HD11	2.38	0.54
40:CM:50:ILE:HD13	40:CM:60:ARG:HD3	1.89	0.54
45:CR:26:GLU:HA	45:CR:81:LEU:HD22	1.90	0.54
55:DA:2840:C:O3'	13:D0:53:HIS:HE1	1.90	0.54
55:DA:105:C:H2'	55:DA:106:C:C6	2.43	0.54
55:DA:1209:G:N2	55:DA:1210:A:H62	2.03	0.54
55:DA:1264:G:O5'	55:DA:1264:G:H8	1.91	0.54
55:DA:729:G:H2'	55:DA:1775:U:O2	2.08	0.54
55:DA:1926:U:H1'	55:DA:1929:G:O6	2.08	0.54
55:DA:2600:A:C6	55:DA:2601:C:N4	2.76	0.54
2:DB:83:G:H4'	25:DX:52:HIS:CG	2.42	0.54
3:DD:142:VAL:HG23	3:DD:193:VAL:CA	2.37	0.54
3:DD:65:ILE:O	3:DD:65:ILE:HD13	2.08	0.54
4:DE:13:ARG:CB	4:DE:21:VAL:HA	2.37	0.54
5:DF:65:TRP:HZ3	5:DF:73:ALA:O	1.91	0.54
58:DL:14:ALA:HB2	58:DL:50:ASP:CA	2.38	0.54
55:DA:1059:G:O2'	58:DL:73:PRO:HD2	2.08	0.54
58:DL:77:LEU:HB3	58:DL:107:ILE:HG12	1.88	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:DM:78:TYR:CD1	9:DM:78:TYR:N	2.75	0.54
11:DO:18:ARG:O	11:DO:19:VAL:HB	2.08	0.54
11:DO:61:ARG:O	11:DO:62:LEU:CB	2.55	0.54
21:DV:180:VAL:CG1	21:DV:181:GLU:N	2.71	0.54
21:DV:43:GLU:O	21:DV:47:VAL:HG23	2.08	0.54
24:DW:15:LYS:HE3	24:DW:67:LYS:HE2	1.89	0.54
23:DZ:13:ILE:HD11	23:DZ:42:GLN:OE1	2.07	0.54
6:AG:109:VAL:HA	26:A4:37:SER:OG	2.08	0.54
28:A6:17:LYS:H	28:A6:19:ARG:HG2	1.73	0.54
30:A8:48:PHE:C	30:A8:49:VAL:CG2	2.76	0.54
1:AA:1103:A:H2'	1:AA:1104:C:H5'	1.90	0.54
1:AA:1139:G:O2'	1:AA:1143:A:N6	2.41	0.54
1:AA:1236:G:C4'	1:AA:1237:A:OP1	2.50	0.54
1:AA:2092:U:C4	1:AA:2225:A:O2'	2.61	0.54
1:AA:2322:A:H3'	1:AA:2323:G:C8	2.39	0.54
1:AA:2526:G:H5'	1:AA:2742:C:O2'	2.08	0.54
1:AA:273(F):C:H3'	1:AA:274:G:C5'	2.38	0.54
1:AA:567:A:OP2	11:AO:29:LYS:NZ	2.41	0.54
1:AA:1993:U:H4'	4:AE:128:SER:CB	2.38	0.54
1:AA:2572:A:C5	4:AE:144:ARG:NH1	2.77	0.54
5:AF:123:LEU:HD13	5:AF:192:LEU:HD22	1.89	0.54
6:AG:38:VAL:CG2	6:AG:93:THR:HG23	2.37	0.54
6:AG:58:GLN:O	6:AG:62:LEU:HB2	2.08	0.54
7:AH:154:PRO:O	7:AH:155:SER:HB2	2.08	0.54
7:AH:18:GLU:HG3	7:AH:25:LYS:HB2	1.90	0.54
15:AR:3:ARG:O	15:AR:7:ILE:HG12	2.08	0.54
1:AA:1312:U:OP2	19:AT:63:LYS:HD3	2.08	0.54
21:AV:105:VAL:HG23	21:AV:106:GLY:N	2.23	0.54
21:AV:111:VAL:O	21:AV:113:ALA:N	2.41	0.54
21:AV:71:VAL:HA	21:AV:88:PHE:HA	1.89	0.54
24:AW:50:ILE:CD1	24:AW:51:ARG:N	2.67	0.54
24:AW:53:LEU:O	24:AW:56:GLN:HB2	2.07	0.54
31:BA:1036:G:H3'	31:BA:1037:C:C6	2.43	0.54
31:BA:1130:A:N6	31:BA:1144:G:H21	2.05	0.54
31:BA:1176:A:C8	31:BA:1176:A:O5'	2.61	0.54
31:BA:1187:G:H21	44:BQ:60:SER:CB	2.21	0.54
31:BA:1194:U:H2'	31:BA:1195:C:C6	2.43	0.54
31:BA:1268:A:C6	31:BA:1269:A:C6	2.96	0.54
31:BA:1292:U:H2'	31:BA:1293:G:C8	2.43	0.54
31:BA:1537:U:H2'	31:BA:1538:C:H6	1.68	0.54
31:BA:452:A:O2'	31:BA:453:A:O4'	2.25	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:BH:36:ASP:O	35:BH:38:GLN:HG2	2.07	0.54
36:BI:62:TRP:CH2	36:BI:64:GLN:HB2	2.42	0.54
37:BJ:20:ASP:HB3	37:BJ:23:VAL:CG2	2.38	0.54
38:BK:29:SER:HB3	38:BK:32:LYS:HB2	1.89	0.54
42:BO:60:LEU:HD23	42:BO:65:GLU:H	1.73	0.54
46:BS:63:GLY:O	46:BS:64:ALA:HB2	2.08	0.54
49:BV:42:PRO:CA	49:BV:45:VAL:HG22	2.38	0.54
49:BV:63:THR:HG22	49:BV:66:MET:HE2	1.88	0.54
54:CA:1026:G:C6	54:CA:1036:G:N2	2.76	0.54
54:CA:1125:U:OP2	54:CA:1145:C:N4	2.41	0.54
54:CA:115:G:O2'	54:CA:116:A:OP2	2.23	0.54
54:CA:38:G:N2	54:CA:397:A:H2	2.03	0.54
54:CA:629:G:C5'	54:CA:630:G:P	2.95	0.54
54:CA:961:U:H2'	54:CA:962:C:O4'	2.08	0.54
32:CE:4:GLU:CG	32:CE:5:ILE:H	2.21	0.54
49:CV:64:GLU:HA	49:CV:67:VAL:HG23	1.89	0.54
16:D1:89:GLU:O	16:D1:90:VAL:O	2.25	0.54
26:D4:17:GLY:H	26:D4:35:VAL:HA	1.73	0.54
55:DA:1063:G:H1'	55:DA:1077:A:N7	2.23	0.54
55:DA:1854:A:H62	55:DA:1888:G:H8	1.56	0.54
55:DA:2175:C:C2'	55:DA:2176:A:H5''	2.38	0.54
2:DB:1(M):A:H2'	2:DB:1(M):A:N3	2.22	0.54
3:DD:238:GLY:C	3:DD:239:ARG:O	2.46	0.54
7:DH:130:ARG:O	7:DH:130:ARG:HD2	2.08	0.54
8:DK:37:VAL:HG13	8:DK:38:LEU:HD12	1.89	0.54
58:DL:108:ALA:CA	58:DL:111:LYS:CD	2.80	0.54
58:DL:18:THR:HG21	58:DL:42:ASN:OD1	2.07	0.54
9:DM:137:LYS:HG3	9:DM:138:LEU:N	2.13	0.54
9:DM:85:ILE:HG23	9:DM:89:LYS:HG2	1.88	0.54
11:DO:77:ARG:HB2	11:DO:78:PRO:HD2	1.90	0.54
14:DQ:106:ARG:HA	14:DQ:110:LEU:CG	2.38	0.54
55:DA:2875:C:O2'	15:DR:5:ALA:HB3	2.08	0.54
21:DV:125:LEU:HG	21:DV:164:ALA:HB3	1.90	0.54
57:DY:70:GLU:O	57:DY:71:LEU:HB3	2.08	0.54
23:DZ:53:VAL:HG22	23:DZ:74:VAL:HG13	1.90	0.54
1:AA:449:A:C4'	16:A1:3:ARG:NH1	2.68	0.53
1:AA:996:A:O3'	16:A1:92:ARG:HD2	2.08	0.53
16:A1:92:ARG:HH12	17:A2:11:GLN:CG	2.21	0.53
1:AA:1373:A:N6	1:AA:1374:G:C2	2.77	0.53
1:AA:1416:G:H2'	1:AA:1417:C:C6	2.43	0.53
1:AA:1773:A:H2'	1:AA:1774:C:H5'	1.90	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1930:G:O2'	1:AA:1931:U:P	2.66	0.53
1:AA:530:G:N3	1:AA:2021:C:O2'	2.41	0.53
1:AA:528:A:H2	1:AA:2043:C:H5'	1.73	0.53
1:AA:2210:G:H3'	1:AA:2210:G:N3	2.22	0.53
1:AA:2303:G:O2'	1:AA:2304:G:H5'	2.08	0.53
1:AA:2327:A:H2'	1:AA:2328:A:C8	2.43	0.53
1:AA:2335:A:HO2'	1:AA:2336:A:H3'	1.72	0.53
1:AA:2702:U:OP1	1:AA:2702:U:O4'	2.27	0.53
1:AA:2888:C:H2'	1:AA:2889:C:H6	1.73	0.53
1:AA:34:C:O2'	1:AA:35:G:P	2.65	0.53
1:AA:881:G:H2'	52:BB:19:G:O6	2.08	0.53
3:AD:106:ILE:O	3:AD:106:ILE:HD13	2.08	0.53
3:AD:35:LYS:CA	3:AD:64:ILE:HG23	2.38	0.53
4:AE:9:VAL:HG21	4:AE:25:VAL:CG1	2.38	0.53
5:AF:29:ASN:H	5:AF:112:MET:CE	2.21	0.53
1:AA:2443:C:OP1	5:AF:68:LYS:HG2	2.08	0.53
7:AH:84:SER:O	7:AH:85:LYS:HB2	2.08	0.53
31:BA:1144:G:H22	31:BA:1146:A:H62	1.53	0.53
31:BA:1176:A:H2'	31:BA:1177:G:C5'	2.38	0.53
31:BA:191(A):G:O2'	31:BA:191(B):G:H5'	2.08	0.53
31:BA:788:U:N3	31:BA:795:C:N4	2.55	0.53
31:BA:926:G:H22	53:B1:45:U:H3'	1.73	0.53
52:BD:15:G:C2	52:BD:48:C:N4	2.76	0.53
52:BD:20:U:H2'	52:BD:21:A:C4'	2.37	0.53
35:BH:32:VAL:CG1	35:BH:33:VAL:N	2.71	0.53
37:BJ:84:ASN:HB2	52:BD:37:MIA:C16	2.36	0.53
38:BK:11:THR:HG23	38:BK:14:ARG:NH1	2.24	0.53
40:BM:7:LYS:CE	40:BM:71:LEU:HD22	2.39	0.53
43:BP:10:PRO:CG	43:BP:18:ALA:HB1	2.38	0.53
45:BR:8:LYS:O	45:BR:12:ILE:HG13	2.08	0.53
31:BA:1288:A:O3'	51:BX:9:ARG:NH1	2.42	0.53
54:CA:1297:C:O2'	54:CA:1298:C:P	2.65	0.53
54:CA:1433:A:C4	54:CA:1468:A:C2	2.96	0.53
54:CA:200:G:H2'	54:CA:201:C:O4'	2.08	0.53
54:CA:405:U:H5''	54:CA:495:A:H2	1.73	0.53
54:CA:595:G:C6	54:CA:641:U:H2'	2.43	0.53
54:CA:605:U:O2'	54:CA:606:G:H5'	2.09	0.53
54:CA:73:G:H2'	54:CA:74:C:C6	2.43	0.53
54:CA:832:C:O4'	54:CA:1539:C:OP1	2.26	0.53
52:CD:55:U:H2'	52:CD:56:C:H5'	1.89	0.53
34:CG:173:TRP:O	34:CG:186:LEU:HB2	2.08	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:CI:8:ILE:HG22	36:CI:10:LEU:HD12	1.90	0.53
38:CK:97:VAL:HG13	38:CK:98:LYS:N	2.24	0.53
40:CM:100:THR:O	40:CM:101:VAL:HB	2.08	0.53
13:D0:84:ALA:HB3	13:D0:85:PRO:HD3	1.90	0.53
27:D5:49:CYS:SG	27:D5:58:LEU:HB2	2.47	0.53
55:DA:1057:A:O2'	55:DA:1058:U:P	2.66	0.53
55:DA:1535:U:C2	55:DA:1536:A:H5''	2.43	0.53
55:DA:1311:G:N2	55:DA:1603:A:H62	2.05	0.53
55:DA:1641:A:H2'	55:DA:1642:G:O4'	2.08	0.53
55:DA:2626:C:H2'	55:DA:2627:G:O4'	2.08	0.53
55:DA:622:G:O2'	55:DA:623:G:H5'	2.08	0.53
3:DD:25:THR:HG22	3:DD:82:ILE:H	1.71	0.53
3:DD:94:LEU:HD13	3:DD:95:LEU:N	2.23	0.53
6:DG:109:VAL:C	6:DG:112:PRO:HD2	2.28	0.53
7:DH:103:LEU:HD21	7:DH:115:VAL:HB	1.90	0.53
7:DH:127:GLU:HG2	7:DH:128:PRO:CD	2.38	0.53
8:DK:35:LEU:O	8:DK:36:ALA:HB2	2.08	0.53
58:DL:136:VAL:O	58:DL:137:GLU:CB	2.50	0.53
58:DL:76:TYR:CD2	58:DL:77:LEU:HD12	2.43	0.53
58:DL:95:LYS:HE2	58:DL:136:VAL:HB	1.90	0.53
12:DP:14:ARG:O	12:DP:72:LYS:NZ	2.41	0.53
21:DV:30:ASN:O	21:DV:32:HIS:N	2.41	0.53
57:DY:122:VAL:O	57:DY:124:ALA:N	2.41	0.53
57:DY:13:LEU:HD23	57:DY:62:ALA:CB	2.06	0.53
13:A0:94:TYR:C	13:A0:117:VAL:HG12	2.29	0.53
27:A5:4:HIS:CB	27:A5:5:PRO:CD	2.80	0.53
30:A8:56:GLU:HA	30:A8:59:LYS:HE3	1.89	0.53
1:AA:1044:G:H2'	1:AA:1045:A:H5''	1.90	0.53
1:AA:1171:G:H1'	1:AA:1173:G:H5'	1.90	0.53
1:AA:456:C:HO2'	1:AA:457:A:P	2.31	0.53
1:AA:523:C:H2'	1:AA:524:U:O4'	2.07	0.53
1:AA:603:A:O2'	1:AA:604:G:OP2	2.26	0.53
1:AA:654(S):G:C2'	1:AA:654(T):A:C8	2.91	0.53
1:AA:684:G:C2	1:AA:774:A:C2	2.96	0.53
3:AD:105:ILE:HD13	3:AD:106:ILE:N	2.23	0.53
3:AD:24:ILE:O	3:AD:25:THR:O	2.26	0.53
3:AD:72:LYS:HE3	3:AD:101:GLU:HB3	1.89	0.53
5:AF:150:GLY:HA2	5:AF:172:TRP:CD2	2.44	0.53
8:AK:114:LEU:O	8:AK:115:ALA:HB3	2.07	0.53
11:AO:146:VAL:HG13	11:AO:147:LEU:N	2.23	0.53
12:AP:29:PHE:CD1	12:AP:29:PHE:N	2.76	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AP:38:GLU:O	12:AP:127:ILE:HG21	2.08	0.53
14:AQ:67:ARG:O	14:AQ:71:ARG:HG3	2.08	0.53
15:AR:125:ARG:O	15:AR:129:ARG:HG3	2.08	0.53
18:AS:18:ARG:HE	18:AS:76:VAL:HG13	1.73	0.53
21:AV:28:MET:O	21:AV:34:ASN:HA	2.07	0.53
24:AW:15:LYS:H	24:AW:67:LYS:HE2	1.73	0.53
31:BA:1274:G:N2	31:BA:1275:A:H62	2.06	0.53
31:BA:591:U:H2'	31:BA:592:G:C8	2.42	0.53
31:BA:678:U:H2'	31:BA:679:C:H6	1.73	0.53
31:BA:959:A:O3'	31:BA:960:U:H4'	2.08	0.53
32:BE:144:ARG:HG3	32:BE:145:LEU:N	2.23	0.53
35:BH:80:ILE:HG22	38:BK:104:ARG:CZ	2.39	0.53
40:BM:47:PHE:O	40:BM:47:PHE:HD1	1.90	0.53
40:BM:99:LYS:HD3	40:BM:100:THR:H	1.72	0.53
41:BN:29:ILE:CB	41:BN:44:SER:HB3	2.33	0.53
54:CA:1004:A:H2'	54:CA:1005:A:N3	2.22	0.53
54:CA:1276:G:H2'	54:CA:1277:C:H6	1.73	0.53
54:CA:1315:U:H2'	54:CA:1316:G:O4'	2.08	0.53
54:CA:974:A:H1'	44:CQ:31:ARG:NE	2.23	0.53
32:CE:112:VAL:O	32:CE:116:GLU:HG3	2.07	0.53
32:CE:14:GLY:C	32:CE:15:VAL:HG22	2.29	0.53
32:CE:175:ARG:HG2	32:CE:175:ARG:HH11	1.72	0.53
32:CE:29:ALA:O	32:CE:32:ILE:HG22	2.08	0.53
32:CE:77:ALA:HB2	32:CE:211:ILE:HD13	1.90	0.53
34:CG:108:LEU:HB3	34:CG:110:PHE:HE1	1.71	0.53
36:CI:14:LEU:HA	36:CI:18:GLN:NE2	2.24	0.53
17:D2:55:ALA:CB	17:D2:101:GLY:HA2	2.37	0.53
55:DA:1021:A:H8	55:DA:1022:G:H5''	1.74	0.53
55:DA:1079:C:C5'	55:DA:1079:C:H6	2.20	0.53
55:DA:1562:A:H2'	55:DA:1563:G:C8	2.43	0.53
55:DA:2391:G:HO2'	55:DA:2392:A:P	2.31	0.53
55:DA:2751:G:C2	7:DH:3:ARG:HB3	2.42	0.53
55:DA:29:U:H2'	55:DA:30:G:H8	1.73	0.53
55:DA:390:A:H4'	55:DA:391:G:O5'	2.07	0.53
55:DA:803:U:C5'	55:DA:803:U:H6	2.10	0.53
55:DA:94:G:N3	24:DW:47:ASN:OD1	2.42	0.53
2:DB:24:G:N7	2:DB:56:G:H2'	2.24	0.53
4:DE:6:GLY:HA2	4:DE:51:PHE:CE2	2.43	0.53
6:DG:55:LYS:HZ1	6:DG:148:MET:CG	2.21	0.53
6:DG:107:LEU:HD11	6:DG:178:PHE:CD1	2.43	0.53
7:DH:98:LEU:HA	7:DH:103:LEU:HA	1.89	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:DI:9:LYS:HG2	56:DI:10:GLU:N	2.23	0.53
56:DI:17:VAL:O	56:DI:18:LEU:C	2.46	0.53
56:DI:26:ALA:O	56:DI:27:LEU:O	2.26	0.53
58:DL:41:PHE:CD2	58:DL:42:ASN:N	2.77	0.53
58:DL:89:HIS:O	58:DL:90:LYS:CB	2.51	0.53
58:DL:95:LYS:HD3	58:DL:136:VAL:CG2	2.32	0.53
11:DO:105:LEU:O	11:DO:106:LEU:CB	2.48	0.53
11:DO:61:ARG:HB2	11:DO:61:ARG:HH21	1.72	0.53
11:DO:86:LYS:HB3	11:DO:118:GLY:HA3	1.89	0.53
15:DR:66:VAL:HG12	15:DR:67:SER:H	1.73	0.53
20:DU:94:LYS:CE	20:DU:101:LYS:HZ3	2.21	0.53
21:DV:152:ALA:HB1	21:DV:163:LEU:HD13	1.90	0.53
21:DV:180:VAL:O	21:DV:181:GLU:C	2.47	0.53
12:DP:20:ALA:H	21:DV:79:ARG:NH2	2.02	0.53
24:DW:52:ASP:O	24:DW:56:GLN:HB2	2.08	0.53
25:DX:28:LEU:HA	25:DX:33:GLN:OE1	2.08	0.53
57:DY:142:LEU:HD13	57:DY:143:GLN:HG2	1.89	0.53
57:DY:4:LYS:HB3	57:DY:5:ARG:HD2	1.90	0.53
23:DZ:67:ILE:N	23:DZ:68:PRO:CD	2.70	0.53
23:DZ:87:PRO:O	23:DZ:91:LYS:HB2	2.08	0.53
13:A0:18:LEU:HD11	13:A0:22:ARG:NE	2.24	0.53
16:A1:8:VAL:HG12	16:A1:11:ARG:HH21	1.73	0.53
17:A2:38:LEU:CD1	17:A2:55:ALA:HB1	2.39	0.53
1:AA:1236:G:O2'	1:AA:1237:A:C8	2.62	0.53
1:AA:1368:G:O2'	1:AA:1369:G:H5'	2.08	0.53
1:AA:1332:G:N2	1:AA:1610:A:H8	2.06	0.53
1:AA:2012:G:C5'	18:AS:96:ILE:HD11	2.38	0.53
1:AA:2238:G:H5'	1:AA:2239:G:N7	2.24	0.53
1:AA:352:G:O2'	1:AA:353:G:OP1	2.27	0.53
1:AA:894:C:C6	1:AA:895:U:C5	2.96	0.53
2:AB:50:G:OP1	14:AQ:62:LYS:HB2	2.08	0.53
1:AA:729:G:C5	3:AD:208:LYS:HB2	2.43	0.53
4:AE:128:SER:OG	4:AE:129:HIS:N	2.39	0.53
4:AE:76:ARG:HD3	4:AE:195:LEU:HD13	1.90	0.53
9:AM:115:ARG:HH11	9:AM:115:ARG:HG2	1.73	0.53
11:AO:3:LEU:O	11:AO:5:ASP:N	2.39	0.53
12:AP:21:THR:HG22	12:AP:99:PRO:O	2.09	0.53
14:AQ:35:ILE:O	14:AQ:35:ILE:HG23	2.09	0.53
21:AV:135:GLU:O	21:AV:136:PHE:HB3	2.09	0.53
21:AV:39:VAL:HG21	21:AV:44:PHE:HB2	1.89	0.53
53:B1:33:G:H2'	53:B1:34:G:C8	2.38	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:1025:U:HO2'	31:BA:1026:G:H8	1.55	0.53
31:BA:1026:G:C5	31:BA:1036:G:N2	2.75	0.53
31:BA:1096:C:O2'	31:BA:1097:C:H5'	2.08	0.53
31:BA:1325:C:H4'	51:BX:17:THR:HG21	1.90	0.53
31:BA:142:G:H2'	31:BA:143:A:C8	2.44	0.53
31:BA:210:U:O4'	31:BA:210:U:OP2	2.26	0.53
32:BE:12:GLU:HB3	32:BE:213:LEU:CD1	2.38	0.53
42:BO:82:VAL:O	42:BO:106:ASP:HB2	2.08	0.53
42:BO:27:LEU:HG	42:BO:33:ARG:HG2	1.88	0.53
54:CA:52:G:O2'	54:CA:53:A:H5'	2.08	0.53
54:CA:77:C:C3'	54:CA:78:G:H5''	2.38	0.53
54:CA:800:G:O2'	54:CA:801:U:H5'	2.08	0.53
54:CA:812:C:OP1	54:CA:903:G:H1'	2.08	0.53
52:CB:5:G:H2'	52:CB:6:G:C8	2.43	0.53
33:CF:14:ILE:HG12	33:CF:15:THR:N	2.23	0.53
33:CF:95:THR:C	33:CF:97:LYS:H	2.11	0.53
36:CI:63:TYR:N	36:CI:63:TYR:HD2	2.06	0.53
54:CA:523:A:N6	42:CO:92:ASP:HB2	2.22	0.53
54:CA:982:U:H5''	44:CQ:6:LEU:CD1	2.38	0.53
46:CS:6:LEU:HB3	46:CS:17:TYR:CD2	2.41	0.53
50:CW:10:LEU:O	50:CW:12:ALA:N	2.42	0.53
55:DA:1021:A:C8	55:DA:1022:G:H5''	2.43	0.53
55:DA:1056:G:C2'	55:DA:1057:A:OP2	2.56	0.53
55:DA:1199:U:H2'	55:DA:1200:C:O4'	2.07	0.53
55:DA:1278:A:H5''	13:D0:36:THR:HG22	1.90	0.53
55:DA:1314:C:OP1	55:DA:1332:G:H5'	2.07	0.53
55:DA:1431:U:O2'	55:DA:1432:C:H5'	2.09	0.53
55:DA:1537:C:H2'	55:DA:1538:G:C8	2.44	0.53
55:DA:1673:U:O2'	55:DA:1674:G:H5'	2.07	0.53
55:DA:1863:G:H2'	55:DA:1864:U:O4'	2.08	0.53
55:DA:1925:C:H3'	55:DA:1925:C:H6	1.71	0.53
55:DA:788:A:O2'	55:DA:789:A:OP2	2.23	0.53
4:DE:5:LEU:O	4:DE:51:PHE:CE2	2.61	0.53
5:DF:7:TYR:CD1	5:DF:7:TYR:N	2.76	0.53
6:DG:113:ARG:HD3	6:DG:140:ILE:O	2.07	0.53
56:DJ:12:LEU:CB	56:DJ:13:SER:OG	2.39	0.53
8:DK:125:GLU:OE2	8:DK:141:LYS:HG2	2.08	0.53
58:DL:122:ALA:C	58:DL:124:ALA:H	2.12	0.53
58:DL:53:VAL:HG12	58:DL:72:PRO:CB	2.38	0.53
55:DA:2414:G:N2	11:DO:67:MET:HE1	2.18	0.53
15:DR:39:ARG:HG2	15:DR:39:ARG:HH11	1.72	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:484:C:OP1	20:DU:51:VAL:HB	2.08	0.53
55:DA:85:G:OP2	20:DU:9:LYS:HG3	2.09	0.53
57:DY:13:LEU:CD2	57:DY:13:LEU:O	2.51	0.53
57:DY:89:ALA:N	57:DY:92:THR:CB	2.71	0.53
26:A4:65:ASP:O	26:A4:67:TYR:N	2.40	0.53
1:AA:1543:A:H1'	1:AA:1545:A:C4'	2.38	0.53
1:AA:1805:U:O2	3:AD:50:THR:HB	2.08	0.53
1:AA:2092:U:H6	1:AA:2092:U:C5'	2.21	0.53
1:AA:192:C:O2'	1:AA:802:A:N3	2.42	0.53
3:AD:132:PRO:HG3	3:AD:190:TYR:CE1	2.43	0.53
5:AF:29:ASN:O	5:AF:112:MET:HE1	2.08	0.53
8:AK:51:ILE:C	8:AK:53:ALA:H	2.12	0.53
10:AN:10:VAL:CG2	10:AN:17:ARG:HA	2.39	0.53
11:AO:112:LEU:H	11:AO:128:HIS:CD2	2.26	0.53
12:AP:32:TYR:O	12:AP:105:GLU:HA	2.07	0.53
12:AP:16:ARG:CB	12:AP:16:ARG:HH11	2.22	0.53
18:AS:92:ARG:O	18:AS:93:ALA:HB3	2.08	0.53
1:AA:1312:U:H3'	19:AT:63:LYS:NZ	2.23	0.53
21:AV:114:GLY:C	21:AV:177:PRO:HB2	2.29	0.53
21:AV:141:VAL:HG22	21:AV:144:LEU:HB3	1.91	0.53
31:BA:1078:U:H2'	31:BA:1079:G:O4'	2.08	0.53
31:BA:1084:G:C5	31:BA:1085:U:C4	2.96	0.53
31:BA:1314:C:OP2	49:BV:6:LYS:HG2	2.09	0.53
31:BA:1523:G:H2'	31:BA:1524:C:C6	2.44	0.53
31:BA:451:A:H4'	31:BA:452:A:O4'	2.08	0.53
31:BA:511:C:C4'	34:BG:43:HIS:CD2	2.92	0.53
31:BA:545:C:O2'	31:BA:549:C:OP1	2.24	0.53
52:BB:51:U:H2'	52:BB:52:G:H8	1.73	0.53
52:BC:44:G:H2'	52:BC:45:U:O4'	2.08	0.53
32:BE:200:ILE:N	32:BE:200:ILE:HD12	2.23	0.53
32:BE:24:TRP:CZ3	32:BE:26:PRO:HA	2.44	0.53
35:BH:105:VAL:HB	35:BH:106:PRO:CD	2.38	0.53
35:BH:153:LYS:NZ	35:BH:153:LYS:HB2	2.24	0.53
39:BL:17:VAL:HA	39:BL:63:ILE:HG13	1.91	0.53
39:BL:26:VAL:HA	39:BL:61:ALA:O	2.08	0.53
42:BO:70:ILE:HD11	42:BO:100:ILE:HD12	1.91	0.53
44:BQ:41:ARG:HG3	44:BQ:42:ILE:N	2.24	0.53
49:BV:47:HIS:H	49:BV:62:ILE:CG2	2.21	0.53
31:BA:1327:C:OP1	51:BX:21:TYR:HD1	1.92	0.53
54:CA:183:G:H2'	54:CA:184:G:C8	2.44	0.53
54:CA:401:C:O2'	54:CA:402:G:H5'	2.08	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:914:A:O2'	54:CA:915:A:H5'	2.09	0.53
32:CE:7:VAL:HG11	32:CE:217:ARG:CZ	2.39	0.53
33:CF:116:VAL:CG2	33:CF:202:ILE:HD11	2.35	0.53
35:CH:60:TYR:CE1	35:CH:64:ARG:NH2	2.75	0.53
37:CJ:126:ASP:HB3	37:CJ:131:LYS:HG3	1.89	0.53
38:CK:112:LEU:HA	38:CK:134:ILE:H	1.73	0.53
42:CO:127:GLU:OE1	42:CO:127:GLU:N	2.42	0.53
54:CA:1492:A:OP1	42:CO:47:LYS:N	2.41	0.53
43:CP:117:VAL:O	43:CP:118:ALA:O	2.25	0.53
45:CR:64:ARG:HH11	45:CR:64:ARG:HG3	1.73	0.53
54:CA:255:G:H1'	47:CT:16:GLN:HE21	1.74	0.53
13:D0:61:HIS:O	13:D0:65:LEU:HD13	2.08	0.53
13:D0:92:GLY:N	13:D0:94:TYR:CE2	2.76	0.53
22:D3:8:GLY:O	22:D3:9:SER:O	2.26	0.53
55:DA:1082:U:O4	55:DA:1083:U:C2	2.62	0.53
55:DA:1084:A:C5'	55:DA:1085:A:OP2	2.55	0.53
55:DA:1368:G:O2'	55:DA:1369:G:H5'	2.09	0.53
55:DA:1916:A:H2'	55:DA:1917:U:O4'	2.07	0.53
55:DA:201:C:H2'	55:DA:202:U:H5'	1.90	0.53
55:DA:2168:G:H2'	55:DA:2168:G:N3	2.23	0.53
55:DA:185:U:H4'	55:DA:218:A:H4'	1.89	0.53
55:DA:270(J):G:H2'	55:DA:270(K):C:C6	2.44	0.53
55:DA:704:G:O2'	55:DA:705:A:P	2.65	0.53
55:DA:775:G:H4'	55:DA:776:G:O5'	2.08	0.53
3:DD:145:VAL:HG12	3:DD:146:GLU:O	2.08	0.53
3:DD:34:VAL:HG21	3:DD:103:ARG:HA	1.90	0.53
4:DE:61:ARG:CB	4:DE:62:PRO:CD	2.58	0.53
7:DH:124:GLU:HB2	7:DH:132:ARG:CD	2.36	0.53
7:DH:123:PHE:CE2	7:DH:133:VAL:HG22	2.43	0.53
58:DL:125:ARG:HD2	58:DL:132:ARG:HH21	1.73	0.53
11:DO:83:VAL:HG13	11:DO:114:ILE:HA	1.89	0.53
15:DR:133:GLU:C	15:DR:135:ALA:H	2.12	0.53
15:DR:95:ARG:NH1	15:DR:95:ARG:HG3	2.20	0.53
21:DV:116:VAL:CG1	21:DV:117:LEU:CD1	2.75	0.53
57:DY:71:LEU:HA	57:DY:113:GLN:CA	2.39	0.53
22:A3:31:VAL:CB	22:A3:35:ASN:HD22	2.18	0.53
22:A3:74:ARG:HH11	22:A3:74:ARG:CG	2.20	0.53
1:AA:1018:C:H2'	1:AA:1019:U:H6	1.71	0.53
1:AA:1103:A:H8	1:AA:1103:A:H5'	1.73	0.53
1:AA:1668:A:H61	1:AA:1676:A:H61	1.56	0.53
1:AA:1818:U:HO2'	1:AA:1819:A:P	2.31	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1856:G:H2'	1:AA:1857:G:H5'	1.90	0.53
1:AA:2311:A:C2'	1:AA:2312:U:C6	2.91	0.53
1:AA:2376:A:C2	14:AQ:112:PHE:HB2	2.43	0.53
1:AA:479:A:HO2'	1:AA:480:A:P	2.31	0.53
1:AA:49:A:O2'	1:AA:50:U:OP2	2.26	0.53
1:AA:511:U:H5''	1:AA:512:G:OP2	2.07	0.53
1:AA:654(C):G:H2'	1:AA:654(D):G:C8	2.44	0.53
1:AA:654(M):C:C3'	1:AA:654(N):G:N7	2.63	0.53
1:AA:864:G:H1'	1:AA:914:C:H42	1.72	0.53
2:AB:14:U:OP2	2:AB:70:C:O2'	2.19	0.53
3:AD:117:VAL:HG22	3:AD:118:VAL:N	2.24	0.53
4:AE:22:PRO:HB2	4:AE:186:GLY:CA	2.38	0.53
4:AE:53:PRO:CG	4:AE:54:GLN:H	2.21	0.53
10:AN:7:TYR:HE1	10:AN:20:MET:HE3	1.73	0.53
11:AO:112:LEU:HD22	11:AO:113:LYS:H	1.72	0.53
11:AO:138:LEU:CD1	11:AO:144:GLU:HG2	2.39	0.53
12:AP:2:LEU:H	12:AP:2:LEU:HD12	1.73	0.53
19:AT:88:LYS:HD2	19:AT:93:GLU:OE2	2.09	0.53
20:AU:86:ARG:HG3	20:AU:86:ARG:HH11	1.74	0.53
21:AV:103:ARG:CB	21:AV:137:ILE:O	2.56	0.53
21:AV:108:PRO:O	21:AV:109:ALA:HB3	2.09	0.53
31:BA:926:G:N2	53:B1:45:U:H3'	2.24	0.53
31:BA:255:G:H2'	31:BA:256:U:C6	2.43	0.53
31:BA:511:C:H4'	34:BG:43:HIS:CD2	2.43	0.53
31:BA:737:A:O2'	36:BI:72:VAL:HG13	2.08	0.53
52:BC:46:G:H5''	52:BC:47:U:OP2	2.08	0.53
32:BE:46:LYS:HA	32:BE:49:GLU:CG	2.38	0.53
37:BJ:62:PHE:O	37:BJ:66:VAL:HG23	2.09	0.53
41:BN:16:SER:O	41:BN:35:PRO:HG3	2.09	0.53
44:BQ:36:PHE:C	44:BQ:36:PHE:CD1	2.82	0.53
54:CA:363:A:H62	42:CO:28:LYS:HD3	1.74	0.53
52:CB:9:A:H2	52:CB:11:C:H41	1.55	0.53
52:CD:56:C:H2'	52:CD:57:G:C8	2.43	0.53
33:CF:16:ARG:HH11	33:CF:16:ARG:CB	2.20	0.53
34:CG:30:LYS:C	34:CG:32:ALA:N	2.55	0.53
35:CH:10:MET:CB	35:CH:32:VAL:HG22	2.38	0.53
54:CA:1297:C:H2'	37:CJ:114:ARG:NH2	2.23	0.53
38:CK:82:HIS:HD2	38:CK:82:HIS:C	2.10	0.53
42:CO:24:VAL:HG12	42:CO:24:VAL:O	2.08	0.53
42:CO:71:PRO:HG3	42:CO:99:HIS:CD2	2.40	0.53
46:CS:71:ARG:O	46:CS:74:LEU:N	2.42	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:D2:31:ALA:O	17:D2:61:VAL:HG12	2.09	0.53
17:D2:81:TYR:C	17:D2:82:ARG:HG3	2.28	0.53
26:D4:69:LYS:HD3	26:D4:70:GLY:HA3	1.91	0.53
55:DA:1059:G:N1	55:DA:1080:A:C2	2.76	0.53
55:DA:1530:G:H2'	55:DA:1531:C:C6	2.43	0.53
55:DA:1688:U:O2	55:DA:1700:A:H8	1.91	0.53
55:DA:2654:A:N1	55:DA:2665:A:H5'	2.24	0.53
5:DF:125:LEU:HD21	5:DF:199:TRP:CE3	2.43	0.53
5:DF:88:VAL:HG11	5:DF:91:GLY:HA3	1.90	0.53
6:DG:99:MET:HG3	6:DG:100:TRP:N	2.24	0.53
6:DG:142:PRO:HG2	6:DG:143:GLU:H	1.74	0.53
7:DH:109:PHE:C	7:DH:111:HIS:N	2.62	0.53
7:DH:153:LYS:HA	7:DH:153:LYS:HE2	1.91	0.53
56:DJ:15:ALA:O	56:DJ:16:THR:CB	2.57	0.53
58:DL:105:LEU:C	58:DL:107:ILE:H	2.12	0.53
19:DT:49:VAL:HG12	19:DT:50:LYS:N	2.23	0.53
20:DU:81:LYS:HB3	20:DU:97:ARG:HD3	1.89	0.53
21:DV:108:PRO:O	21:DV:109:ALA:CB	2.57	0.53
21:DV:179:ASP:OD1	21:DV:179:ASP:N	2.41	0.53
21:DV:53:ILE:HG22	21:DV:71:VAL:HG13	1.91	0.53
57:DY:112:LEU:O	57:DY:113:GLN:HG2	2.08	0.53
57:DY:144:ALA:CB	57:DY:145:PRO:CD	2.81	0.53
29:A7:33:ARG:HH11	29:A7:33:ARG:HB2	1.73	0.53
1:AA:1043:C:H2'	1:AA:1044:G:H5'	1.91	0.53
1:AA:1290:C:H2'	1:AA:1291:C:H6	1.73	0.53
1:AA:2638:G:O2'	1:AA:2639:A:H8	1.90	0.53
1:AA:2729:G:H1'	4:AE:187:ALA:HB3	1.90	0.53
1:AA:681:G:H2'	1:AA:682:G:O4'	2.08	0.53
1:AA:992:C:H2'	1:AA:993:G:H8	1.72	0.53
2:AB:39:A:C2	2:AB:44:G:C4	2.97	0.53
3:AD:30:GLU:CG	3:AD:63:ARG:CZ	2.85	0.53
3:AD:35:LYS:HE3	3:AD:63:ARG:O	2.08	0.53
4:AE:3:GLY:O	4:AE:4:ILE:HG23	2.08	0.53
4:AE:60:ASN:O	4:AE:61:ARG:CB	2.57	0.53
5:AF:20:LEU:HD23	5:AF:21:ALA:CB	2.38	0.53
9:AM:30:ILE:CD1	9:AM:99:LEU:HD11	2.38	0.53
12:AP:23:GLY:HA2	21:AV:78:LYS:CE	2.35	0.53
20:AU:50:ARG:HB3	20:AU:53:PRO:CG	2.35	0.53
20:AU:42:VAL:HG12	20:AU:67:LEU:HD13	1.90	0.53
21:AV:162:GLU:O	21:AV:163:LEU:C	2.47	0.53
31:BA:1045:C:O2	31:BA:1045:C:H2'	2.09	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:1321:C:H3'	31:BA:1322:C:H5''	1.91	0.53
31:BA:251:G:N1	31:BA:266:G:C6	2.76	0.53
31:BA:498:A:H4'	31:BA:500:G:OP1	2.08	0.53
31:BA:566:G:C4'	31:BA:567:G:OP1	2.56	0.53
31:BA:620:C:H2'	31:BA:621:A:O4'	2.09	0.53
32:BE:19:HIS:CE1	32:BE:204:ASN:HD22	2.26	0.53
33:BF:150:LYS:HG3	33:BF:169:ALA:HB2	1.90	0.53
39:BL:55:ALA:HA	39:BL:58:HIS:HD2	1.74	0.53
40:BM:48:THR:HA	40:BM:62:HIS:CB	2.34	0.53
42:BO:68:ALA:CB	42:BO:100:ILE:HG13	2.39	0.53
46:BS:18:ARG:HA	46:BS:38:TYR:HA	1.91	0.53
48:BU:70:ILE:HG23	48:BU:79:LEU:HD12	1.89	0.53
50:BW:48:LYS:HB3	50:BW:51:GLU:HG3	1.90	0.53
53:C1:57:U:C3'	53:C1:57:U:O2	2.56	0.53
54:CA:1004:A:OP1	54:CA:1025:U:O4	2.26	0.53
54:CA:1081:G:H2'	54:CA:1082:G:H8	1.74	0.53
54:CA:1068:G:OP2	54:CA:1094:G:H5''	2.09	0.53
54:CA:197:A:C5	54:CA:221:C:H4'	2.43	0.53
54:CA:341:C:H2'	54:CA:342:C:H6	1.74	0.53
54:CA:371:G:N2	54:CA:374:A:N6	2.55	0.53
54:CA:562:C:O2'	42:CO:15:ARG:HD2	2.09	0.53
54:CA:624:C:H2'	54:CA:625:G:H8	1.73	0.53
54:CA:5:U:O2'	54:CA:6:G:C4	2.59	0.53
52:CB:37:MIA:HN6	52:CB:37:MIA:H162	1.73	0.53
52:CD:64:A:N3	52:CD:65:G:H1'	2.23	0.53
32:CE:17:PHE:CD1	32:CE:17:PHE:O	2.62	0.53
35:CH:137:GLU:O	35:CH:141:GLN:HG3	2.09	0.53
46:CS:8:ARG:HG2	46:CS:8:ARG:HH11	1.73	0.53
48:CU:18:ARG:N	48:CU:18:ARG:HD3	2.16	0.53
16:D1:24:TYR:HE1	16:D1:39:LEU:HD23	1.73	0.53
18:DS:15:ARG:HE	27:D5:20:ARG:NH1	2.07	0.53
27:D5:57:VAL:HG13	27:D5:57:VAL:O	2.07	0.53
55:DA:1125:G:C6	55:DA:1126:A:N6	2.77	0.53
55:DA:1019:U:N3	55:DA:1142(A):A:N6	2.47	0.53
55:DA:2212:A:H1'	55:DA:2215:G:C5	2.44	0.53
55:DA:2211:G:O2'	55:DA:2212:A:P	2.67	0.53
55:DA:2336:A:H61	22:D3:43:THR:CG2	2.20	0.53
55:DA:27:G:N2	55:DA:512:G:C2'	2.72	0.53
55:DA:897:C:C6	55:DA:897:C:OP1	2.62	0.53
55:DA:2619:C:H5''	4:DE:152:LYS:HA	1.90	0.53
6:DG:161:THR:HG22	6:DG:162:THR:H	1.74	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:DH:20:ALA:HB1	7:DH:21:PRO:CD	2.38	0.53
7:DH:86:GLU:O	7:DH:87:LEU:CB	2.56	0.53
8:DK:17:GLN:HE21	8:DK:19:VAL:HB	1.74	0.53
8:DK:68:LEU:HA	8:DK:71:ILE:CG2	2.38	0.53
58:DL:83:GLY:C	58:DL:85:GLU:N	2.58	0.53
12:DP:51:ARG:HH11	12:DP:51:ARG:HG2	1.73	0.53
12:DP:66:ILE:HA	12:DP:104:PHE:CA	2.31	0.53
20:DU:47:LYS:HA	20:DU:60:PHE:CD1	2.43	0.53
21:DV:6:LYS:NZ	21:DV:43:GLU:HG3	2.23	0.53
21:DV:58:VAL:O	21:DV:67:LEU:O	2.26	0.53
24:DW:31:GLU:O	24:DW:35:LEU:HD23	2.08	0.53
57:DY:15:GLU:HG3	57:DY:19:ARG:NH2	2.24	0.53
57:DY:51:LEU:HD11	57:DY:83:TYR:N	2.23	0.53
17:A2:55:ALA:HA	17:A2:101:GLY:HA2	1.91	0.53
1:AA:2420:C:N4	30:A8:31:HIS:HB3	2.13	0.53
1:AA:1358:G:C2'	1:AA:1359:A:OP2	2.57	0.53
1:AA:2134:A:N6	1:AA:2157:G:O2'	2.41	0.53
1:AA:2278:A:C2'	1:AA:2279:G:O5'	2.57	0.53
1:AA:2619:C:OP1	4:AE:152:LYS:HE3	2.08	0.53
1:AA:34:C:H2'	1:AA:35:G:OP2	2.09	0.53
1:AA:986:C:C2'	1:AA:987:G:H5'	2.39	0.53
2:AB:12:C:H5''	2:AB:13:A:OP1	2.09	0.53
3:AD:94:LEU:CD2	3:AD:94:LEU:C	2.77	0.53
5:AF:5:ALA:H	5:AF:19:GLU:HA	1.74	0.53
5:AF:32:LEU:O	5:AF:36:VAL:HG23	2.08	0.53
6:AG:139:LEU:HD12	6:AG:140:ILE:N	2.22	0.53
7:AH:4:ILE:HD11	7:AH:7:LEU:HB3	1.90	0.53
11:AO:107:LYS:C	11:AO:109:GLY:H	2.12	0.53
20:AU:43:ASN:HA	20:AU:63:LYS:O	2.09	0.53
23:AZ:7:ILE:HG12	23:AZ:91:LYS:HZ1	1.73	0.53
31:BA:1004:A:O2'	31:BA:1005:A:O4'	2.26	0.53
31:BA:1234:C:O2'	31:BA:1235:U:H5'	2.09	0.53
31:BA:1402:C:H2'	31:BA:1403:C:O4'	2.09	0.53
31:BA:890:G:O2'	31:BA:891:U:OP2	2.27	0.53
31:BA:99:C:H2'	31:BA:101:A:C8	2.43	0.53
52:BB:25:C:H5'	52:BB:26:A:OP2	2.08	0.53
52:BD:21:A:N3	52:BD:21:A:C3'	2.71	0.53
52:BD:7:A:H5'	52:BD:8:U:OP2	2.09	0.53
34:BG:16:GLY:O	34:BG:17:VAL:C	2.47	0.53
37:BJ:97:GLN:NE2	37:BJ:101:LEU:HD11	2.16	0.53
37:BJ:111:ARG:NH1	37:BJ:123:GLU:N	2.56	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:BL:47:LEU:CD1	39:BL:47:LEU:H	2.21	0.53
39:BL:47:LEU:HD12	39:BL:47:LEU:H	1.74	0.53
40:BM:49:VAL:HG11	44:BQ:41:ARG:O	2.09	0.53
42:BO:40:VAL:HG12	42:BO:40:VAL:O	2.08	0.53
42:BO:74:GLY:O	42:BO:75:HIS:HB3	2.08	0.53
44:BQ:14:PRO:HG2	44:BQ:15:LYS:N	2.21	0.53
54:CA:1058:G:H2'	54:CA:1059:C:O4'	2.08	0.53
54:CA:115:G:H1'	54:CA:116:A:N7	2.24	0.53
54:CA:1235:U:H2'	54:CA:1236:A:O4'	2.09	0.53
54:CA:1237:C:O4'	54:CA:1334:G:N2	2.42	0.53
54:CA:275:G:OP1	47:CT:14:LYS:HG2	2.08	0.53
52:CC:18:G:H1'	52:CC:58:A:H2	1.74	0.53
33:CF:19:GLU:HA	33:CF:54:ARG:NH2	2.24	0.53
35:CH:41:VAL:HG22	35:CH:69:VAL:HG21	1.90	0.53
38:CK:132:GLU:O	38:CK:134:ILE:N	2.42	0.53
39:CL:113:LYS:HD3	39:CL:119:ALA:HA	1.91	0.53
16:D1:95:LEU:O	16:D1:98:LEU:HB3	2.09	0.53
17:D2:47:VAL:HG13	17:D2:48:GLY:N	2.24	0.53
55:DA:1140:C:OP1	9:DM:23:LEU:HB3	2.08	0.53
55:DA:1266:G:C6	18:DS:16:LYS:HE2	2.44	0.53
55:DA:1385:G:H5'	55:DA:1386:C:OP1	2.09	0.53
55:DA:1567:A:H5'	3:DD:58:HIS:CD2	2.44	0.53
55:DA:1709:U:H2'	55:DA:1710:C:H6	1.72	0.53
55:DA:1778:U:H2'	55:DA:1784:A:H62	1.72	0.53
55:DA:1931:U:O2	55:DA:1931:U:O4'	2.25	0.53
55:DA:857:C:H5'	22:D3:77:ARG:HH21	1.74	0.53
3:DD:72:LYS:CG	3:DD:103:ARG:NH2	2.71	0.53
8:DK:19:VAL:HG22	8:DK:20:ASP:N	2.24	0.53
58:DL:11:GLN:CG	58:DL:41:PHE:CZ	2.88	0.53
9:DM:43:THR:CG2	9:DM:45:ASN:ND2	2.72	0.53
55:DA:943:U:OP2	11:DO:36:LYS:HE3	2.08	0.53
12:DP:66:ILE:CA	12:DP:104:PHE:HA	2.31	0.53
15:DR:1:MET:O	15:DR:3:ARG:HG2	2.08	0.53
21:DV:113:ALA:O	21:DV:114:GLY:C	2.45	0.53
57:DY:131:MET:O	57:DY:133:GLU:CG	2.56	0.53
57:DY:138:LEU:HG	57:DY:139:VAL:H	1.74	0.53
57:DY:2:PRO:O	57:DY:3:ASN:CB	2.55	0.53
22:A3:32:ARG:H	22:A3:35:ASN:HD22	1.42	0.53
22:A3:36:ILE:HD13	22:A3:36:ILE:N	2.24	0.53
26:A4:15:ILE:HD12	26:A4:15:ILE:H	1.74	0.53
1:AA:1098:A:H3'	1:AA:1099:G:H5''	1.91	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1341:U:H5''	19:AT:57:LEU:CD2	2.38	0.53
1:AA:1429:G:H2'	1:AA:1430:C:H6	1.69	0.53
1:AA:1646:C:C2'	1:AA:1647:G:OP1	2.56	0.53
1:AA:2287:A:O2'	1:AA:2288:A:H3'	2.08	0.53
1:AA:270(H):C:O2'	1:AA:270(I):G:H5'	2.09	0.53
1:AA:2832:U:O4	1:AA:2883:A:H5''	2.09	0.53
1:AA:790:C:O2'	1:AA:791:C:OP1	2.22	0.53
3:AD:227:ASN:HB3	3:AD:228:PRO:HD2	1.90	0.53
3:AD:27:THR:HG21	3:AD:83:GLU:CD	2.30	0.53
4:AE:23:VAL:HG23	4:AE:24:THR:N	2.24	0.53
4:AE:47:VAL:O	4:AE:80:GLU:HG2	2.09	0.53
6:AG:60:LEU:O	6:AG:64:THR:HG22	2.08	0.53
7:AH:20:ALA:HB3	7:AH:23:ARG:HG3	1.91	0.53
1:AA:943:U:OP2	11:AO:36:LYS:CE	2.56	0.53
11:AO:62:LEU:HD22	11:AO:63:PRO:O	2.08	0.53
14:AQ:11:LYS:HB2	14:AQ:91:PRO:HD3	1.91	0.53
20:AU:51:VAL:HG22	20:AU:51:VAL:O	2.08	0.53
21:AV:106:GLY:O	21:AV:107:THR:CB	2.56	0.53
31:BA:1375:A:H4'	37:BJ:29:LYS:NZ	2.24	0.53
31:BA:197:A:O2'	31:BA:198:G:P	2.67	0.53
31:BA:279:A:HO2'	31:BA:280:C:P	2.30	0.53
31:BA:421:U:C2'	31:BA:421:U:O2	2.57	0.53
31:BA:689:C:H2'	31:BA:690:G:C5'	2.37	0.53
52:BC:18:G:N3	52:BC:18:G:H2'	2.23	0.53
52:BC:65:G:O2'	52:BC:66:U:H5'	2.08	0.53
52:BC:65:G:H2'	52:BC:66:U:C6	2.43	0.53
52:BD:14:A:H2'	52:BD:15:G:C8	2.43	0.53
32:BE:108:ILE:HG22	32:BE:108:ILE:O	2.08	0.53
32:BE:97:TRP:CZ3	32:BE:173:ALA:HA	2.44	0.53
32:BE:8:LYS:O	32:BE:9:GLU:HB3	2.09	0.53
33:BF:127:ARG:HH11	33:BF:127:ARG:HG2	1.73	0.53
33:BF:58:GLU:HB2	33:BF:65:ALA:HB2	1.89	0.53
34:BG:53:ASP:HB3	34:BG:57:ARG:NH1	2.24	0.53
41:BN:127:LYS:O	41:BN:128:ALA:HB3	2.08	0.53
42:BO:109:GLY:HA3	42:BO:121:GLY:O	2.09	0.53
50:BW:65:LYS:O	50:BW:68:LYS:HG3	2.08	0.53
54:CA:1139:G:N2	54:CA:1143:G:C6	2.77	0.53
54:CA:1158:C:C2	54:CA:1160:G:N7	2.77	0.53
54:CA:1179:A:O3'	39:CL:103:THR:HG23	2.09	0.53
54:CA:1306:A:H61	54:CA:1331:G:H1'	1.74	0.53
54:CA:1336:C:O2'	54:CA:1337:G:C4	2.61	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1390:U:H2'	54:CA:1391:U:C6	2.43	0.53
54:CA:1530:G:H2'	54:CA:1531:A:C8	2.43	0.53
54:CA:344:A:H5''	54:CA:345:C:OP2	2.09	0.53
54:CA:691:G:H1'	54:CA:696:A:N6	2.24	0.53
54:CA:794:A:C2	54:CA:795:C:C4	2.97	0.53
32:CE:33:TYR:HD2	32:CE:43:ASP:HB2	1.73	0.53
34:CG:198:VAL:CG1	34:CG:199:ASN:H	2.21	0.53
34:CG:199:ASN:O	34:CG:201:GLN:N	2.41	0.53
38:CK:97:VAL:HG13	38:CK:98:LYS:H	1.74	0.53
43:CP:66:LEU:O	43:CP:67:GLU:C	2.47	0.53
50:CW:18:GLN:O	50:CW:22:ARG:HG3	2.07	0.53
55:DA:1317:A:H2'	55:DA:1318:C:C6	2.44	0.53
55:DA:1497:U:C5'	55:DA:1498:C:OP2	2.55	0.53
55:DA:1503:U:H2'	55:DA:1504:C:C6	2.44	0.53
55:DA:1888:G:N3	55:DA:1888:G:H5'	2.24	0.53
55:DA:363(A):A:O2'	55:DA:363(B):G:H5'	2.08	0.53
55:DA:583:G:H5''	16:D1:10:ARG:NH1	2.23	0.53
55:DA:80:G:O2'	55:DA:81:G:H5'	2.09	0.53
2:DB:40:U:HO2'	2:DB:43:C:H5	1.57	0.53
4:DE:54:GLN:HG2	4:DE:75:VAL:HG22	1.90	0.53
56:DJ:7:ARG:O	56:DJ:8:ILE:HD13	2.09	0.53
58:DL:95:LYS:CG	58:DL:136:VAL:HG11	2.38	0.53
58:DL:21:PRO:O	58:DL:24:GLY:O	2.27	0.53
58:DL:63:ARG:O	58:DL:64:SER:CB	2.55	0.53
55:DA:1059:G:H4'	58:DL:71:THR:HB	1.90	0.53
58:DL:77:LEU:C	58:DL:107:ILE:HD11	2.28	0.53
11:DO:61:ARG:O	11:DO:62:LEU:HB3	2.08	0.53
20:DU:42:VAL:HG11	20:DU:65:ALA:HB3	1.88	0.53
57:DY:138:LEU:CG	57:DY:139:VAL:H	2.21	0.53
57:DY:27:VAL:N	57:DY:111:LEU:H	2.06	0.53
57:DY:28:ASN:ND2	57:DY:83:TYR:CE2	2.75	0.53
57:DY:52:PHE:O	57:DY:53:VAL:HG13	2.09	0.53
26:A4:63:TYR:CE2	49:BV:41:VAL:HG13	2.44	0.53
6:AG:67:LYS:HE2	26:A4:6:HIS:CD2	2.44	0.53
27:A5:41:PRO:HG2	27:A5:44:THR:OG1	2.09	0.53
30:A8:29:LYS:O	30:A8:30:ARG:C	2.47	0.53
1:AA:1022:G:O2'	1:AA:1023:U:O5'	2.27	0.53
1:AA:1062:G:H2'	1:AA:1063:G:H8	1.74	0.53
1:AA:1814:G:H2'	1:AA:1815:A:C8	2.43	0.53
1:AA:1924:C:N4	1:AA:1925:C:C5	2.76	0.53
1:AA:2443:C:O2'	1:AA:2444:G:H5'	2.09	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:428:A:H8	1:AA:428:A:OP2	1.92	0.53
1:AA:570:G:H2'	1:AA:2030:A:C5	2.44	0.53
1:AA:5:A:H2'	1:AA:6:A:C8	2.44	0.53
1:AA:986:C:O2'	1:AA:987:G:H5'	2.08	0.53
2:AB:90:C:P	12:AP:16:ARG:HD2	2.49	0.53
3:AD:102:LYS:C	3:AD:103:ARG:HG2	2.29	0.53
4:AE:203:LYS:HD3	4:AE:203:LYS:O	2.09	0.53
1:AA:2786:U:H4'	4:AE:64:LYS:C	2.29	0.53
7:AH:18:GLU:CB	7:AH:25:LYS:HB2	2.39	0.53
8:AK:51:ILE:C	8:AK:53:ALA:N	2.61	0.53
10:AN:86:ILE:H	10:AN:86:ILE:HD12	1.73	0.53
11:AO:78:PRO:HB3	11:AO:111:ARG:NH2	2.24	0.53
15:AR:102:ILE:HD12	15:AR:103:ARG:N	2.24	0.53
10:AN:73:ASP:OD1	15:AR:32:TYR:OH	2.26	0.53
19:AT:23:GLU:O	19:AT:25:LYS:N	2.39	0.53
21:AV:10:ARG:NH1	21:AV:36:LYS:HD3	2.24	0.53
24:AW:48:HIS:CD2	24:AW:49:LYS:N	2.77	0.53
31:BA:1260:C:H4'	31:BA:1284:C:H5'	1.89	0.53
31:BA:353:A:H2'	31:BA:354:G:OP2	2.08	0.53
31:BA:389:A:H2'	31:BA:389:A:N3	2.23	0.53
31:BA:425:G:O2'	31:BA:426:G:H5'	2.08	0.53
31:BA:500:G:H2'	31:BA:501:C:C6	2.44	0.53
31:BA:92:G:O2'	31:BA:93:U:H5'	2.09	0.53
31:BA:948:C:C5	43:BP:106:ASN:ND2	2.77	0.53
52:BD:24:G:O2'	52:BD:25:C:H5'	2.09	0.53
34:BG:29:PRO:HD2	34:BG:30:LYS:HD3	1.91	0.53
34:BG:30:LYS:C	34:BG:32:ALA:H	2.12	0.53
31:BA:1240:U:O2'	37:BJ:38:LEU:HD23	2.09	0.53
40:BM:30:SER:OG	40:BM:81:THR:HA	2.09	0.53
40:BM:90:LEU:HD12	40:BM:90:LEU:N	2.24	0.53
45:BR:11:VAL:HG21	45:BR:34:LEU:HD22	1.89	0.53
49:BV:24:ALA:O	49:BV:27:GLU:OE1	2.27	0.53
49:BV:36:ARG:HB2	49:BV:72:GLY:H	1.74	0.53
54:CA:1158:C:H2'	54:CA:1158:C:O2	2.09	0.53
54:CA:1161:C:H2'	54:CA:1162:C:H6	1.74	0.53
54:CA:1285:A:H4'	54:CA:1286:A:O5'	2.08	0.53
54:CA:389:A:H2'	54:CA:390:C:O4'	2.09	0.53
54:CA:406:G:C5'	34:CG:5:ILE:HD13	2.36	0.53
54:CA:872:A:O2'	54:CA:873:A:C3'	2.55	0.53
54:CA:95:G:H2'	54:CA:96:G:C5'	2.38	0.53
52:CB:57:G:H4'	21:DV:182:LYS:NZ	2.23	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:CC:51:U:H2'	52:CC:52:G:C8	2.44	0.53
52:CC:53:G:O2'	52:CC:54:U:H5'	2.08	0.53
34:CG:131:ARG:HH11	34:CG:131:ARG:HG3	1.73	0.53
35:CH:82:VAL:CG1	35:CH:83:GLU:N	2.71	0.53
13:D0:12:ARG:HG3	13:D0:12:ARG:NH1	2.22	0.53
13:D0:78:LYS:HE2	13:D0:83:ILE:HD11	1.90	0.53
17:D2:65:GLY:HA3	17:D2:91:TYR:CE1	2.44	0.53
17:D2:64:HIS:CG	17:D2:92:THR:HG22	2.44	0.53
55:DA:1084:A:H3'	55:DA:1085:A:C8	2.44	0.53
55:DA:1142(A):A:C5	55:DA:1144:G:N7	2.76	0.53
55:DA:1175:U:H4'	55:DA:1176:G:OP1	2.07	0.53
55:DA:1577:C:H2'	55:DA:1578:U:C6	2.44	0.53
55:DA:2639:A:H2'	55:DA:2640:G:H5'	1.89	0.53
55:DA:2780:G:H22	9:DM:100:GLU:CD	2.11	0.53
55:DA:27:G:H22	55:DA:512:G:C2'	2.22	0.53
55:DA:2862:G:H2'	55:DA:2863:C:H6	1.74	0.53
55:DA:654(F):C:H2'	55:DA:654(G):C:OP1	2.08	0.53
55:DA:900:A:H3'	55:DA:901:A:H8	1.74	0.53
3:DD:35:LYS:CG	3:DD:64:ILE:HG23	2.39	0.53
4:DE:116:VAL:CG2	4:DE:122:PHE:CD2	2.92	0.53
6:DG:121:ASN:HD22	6:DG:121:ASN:C	2.12	0.53
6:DG:34:LEU:HD21	6:DG:99:MET:HE1	1.91	0.53
7:DH:9:ILE:O	7:DH:9:ILE:HG13	2.09	0.53
56:DJ:8:ILE:HA	56:DJ:11:GLU:HB2	1.90	0.53
56:DJ:4:ASP:O	56:DJ:7:ARG:C	2.47	0.53
9:DM:112:LEU:C	9:DM:112:LEU:HD23	2.28	0.53
12:DP:51:ARG:O	12:DP:55:VAL:CG1	2.57	0.53
15:DR:49:VAL:HG13	15:DR:49:VAL:O	2.08	0.53
20:DU:9:LYS:O	20:DU:27:VAL:HG22	2.09	0.53
24:DW:69:ARG:HH11	24:DW:69:ARG:HG2	1.74	0.53
57:DY:89:ALA:HB2	57:DY:125:LEU:HD12	1.89	0.53
17:A2:49:THR:CG2	17:A2:50:PRO:HD3	2.38	0.53
26:A4:11:PRO:HA	26:A4:24:THR:HB	1.90	0.53
26:A4:55:ARG:HE	26:A4:56:VAL:HB	1.74	0.53
1:AA:1053:C:C2'	1:AA:1054:A:H5''	2.39	0.53
1:AA:996:A:N6	1:AA:1160:G:C6	2.77	0.53
1:AA:1466:G:H5'	1:AA:1467:C:OP1	2.08	0.53
1:AA:1490:A:H5'	1:AA:1491:G:OP2	2.07	0.53
1:AA:1771:C:C1'	1:AA:1786:A:C8	2.92	0.53
1:AA:2119:A:N6	1:AA:2170:A:C5	2.77	0.53
1:AA:2185:C:H2'	1:AA:2186:G:C8	2.44	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2614:A:C4'	1:AA:2615:U:OP1	2.56	0.53
1:AA:2672:G:H3'	1:AA:2673:G:H5''	1.91	0.53
1:AA:2820:A:H61	4:AE:192:ASN:CA	2.22	0.53
1:AA:2824:C:C2'	1:AA:2825:C:H5'	2.39	0.53
1:AA:877:U:O2'	1:AA:878:A:C5'	2.56	0.53
2:AB:44:G:OP2	26:A4:1:MET:N	2.42	0.53
2:AB:96:G:N2	2:AB:97:G:C1'	2.72	0.53
4:AE:101:ARG:HH11	4:AE:171:GLU:N	2.06	0.53
4:AE:71:GLY:C	4:AE:73:GLU:N	2.61	0.53
6:AG:128:ARG:HG3	6:AG:128:ARG:NH2	2.23	0.53
6:AG:51:ARG:HB2	6:AG:51:ARG:NH1	2.23	0.53
7:AH:111:HIS:ND1	7:AH:112:PRO:HD2	2.24	0.53
7:AH:37:VAL:HG22	7:AH:38:SER:N	2.24	0.53
9:AM:126:PRO:O	9:AM:127:ASP:CB	2.56	0.53
9:AM:30:ILE:O	9:AM:34:LEU:HD23	2.09	0.53
11:AO:52:GLU:OE1	11:AO:52:GLU:C	2.47	0.53
12:AP:69:PHE:CD1	12:AP:70:PRO:HD2	2.44	0.53
14:AQ:100:ALA:HA	14:AQ:103:GLU:HG2	1.90	0.53
31:BA:1277:C:O2'	31:BA:1279:A:H1'	2.09	0.53
31:BA:166:G:O2'	31:BA:167:G:H5'	2.09	0.53
31:BA:350:G:H5'	31:BA:351:G:OP2	2.09	0.53
31:BA:686:U:H2'	31:BA:687:A:H8	1.73	0.53
52:BB:22:G:N7	52:BB:46:G:N2	2.57	0.53
32:BE:98:LEU:HB2	32:BE:101:MET:CE	2.39	0.53
37:BJ:129:GLU:O	37:BJ:129:GLU:HG3	2.09	0.53
40:BM:30:SER:C	40:BM:81:THR:HG22	2.30	0.53
31:BA:568:G:O6	42:BO:5:PRO:HD3	2.08	0.53
43:BP:13:LYS:CA	43:BP:44:ARG:HH11	2.19	0.53
54:CA:1160:G:N1	54:CA:1177:G:N2	2.57	0.53
54:CA:1234:C:H5'	54:CA:1365:G:OP1	2.09	0.53
54:CA:1286:A:C8	54:CA:1287:A:H4'	2.44	0.53
54:CA:1380:U:H4'	54:CA:1381:U:O5'	2.09	0.53
54:CA:556:C:C2'	54:CA:557:G:H5'	2.38	0.53
32:CE:88:ALA:HB2	32:CE:219:VAL:HG13	1.89	0.53
35:CH:8:GLU:OE2	35:CH:63:ARG:NH2	2.41	0.53
39:CL:9:ARG:HA	39:CL:76:ALA:HB1	1.91	0.53
42:CO:89:ARG:NH2	42:CO:91:LYS:HZ3	2.07	0.53
43:CP:4:ILE:CG2	43:CP:5:ALA:H	2.18	0.53
44:CQ:39:LEU:HB3	44:CQ:43:CYS:CB	2.37	0.53
49:CV:41:VAL:HG12	49:CV:44:MET:N	2.24	0.53
55:DA:1493:C:O2	55:DA:1493:C:C2'	2.57	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1992:G:O2'	55:DA:1993:U:OP2	2.18	0.53
52:CD:19:G:O6	55:DA:2112:G:O4'	2.26	0.53
55:DA:2128:C:O2'	55:DA:2173:A:N3	2.42	0.53
55:DA:2318:G:C2'	55:DA:2319:G:OP1	2.57	0.53
55:DA:2688:U:H1'	55:DA:2721:A:N6	2.24	0.53
55:DA:616:A:O2'	55:DA:617:G:P	2.66	0.53
55:DA:686:G:H4'	55:DA:687:C:OP1	2.05	0.53
55:DA:747:U:C4	55:DA:2613:U:C4	2.96	0.53
55:DA:90:U:H5''	55:DA:91:A:OP1	2.09	0.53
5:DF:110:LEU:HD11	5:DF:181:LEU:HD12	1.91	0.53
7:DH:6:ARG:C	7:DH:8:PRO:HD2	2.30	0.53
8:DK:14:ASP:O	8:DK:16:GLY:N	2.39	0.53
55:DA:1081:U:O2	58:DL:115:LEU:HD22	2.09	0.53
58:DL:125:ARG:HD2	58:DL:132:ARG:NH2	2.24	0.53
58:DL:58:THR:CB	58:DL:66:THR:HG23	2.38	0.53
15:DR:57:PHE:O	15:DR:58:ASN:ND2	2.37	0.53
20:DU:95:LYS:HA	20:DU:101:LYS:HB2	1.91	0.53
21:DV:154:ASP:N	21:DV:154:ASP:OD2	2.40	0.53
24:DW:50:ILE:HD13	24:DW:51:ARG:H	1.69	0.53
24:DW:65:ASN:ND2	24:DW:69:ARG:NH2	2.46	0.53
57:DY:135:ARG:HG2	56:DJ:10:GLU:CB	2.39	0.53
17:A2:37:VAL:HG23	17:A2:38:LEU:N	2.23	0.52
16:A1:88:ILE:HG22	17:A2:49:THR:O	2.09	0.52
30:A8:50:LEU:HB2	30:A8:53:PRO:HG2	1.90	0.52
1:AA:727:A:OP1	1:AA:1431:U:O2'	2.27	0.52
1:AA:1925:C:H42	1:AA:1929:G:N2	2.08	0.52
1:AA:2498:C:O2'	1:AA:2499:C:H5'	2.07	0.52
1:AA:2556:C:O2'	1:AA:2557:G:H5'	2.09	0.52
1:AA:606:U:H4'	1:AA:658:C:H4'	1.90	0.52
1:AA:84:A:C8	1:AA:99:U:H5	2.26	0.52
4:AE:64:LYS:HG2	4:AE:65:GLY:H	1.73	0.52
5:AF:117:ARG:HD3	5:AF:120:GLU:OE1	2.08	0.52
5:AF:29:ASN:H	5:AF:112:MET:HE1	1.74	0.52
6:AG:60:LEU:C	6:AG:62:LEU:H	2.12	0.52
11:AO:66:GLY:O	11:AO:67:MET:CB	2.57	0.52
19:AT:64:LYS:NZ	19:AT:73:ARG:NH2	2.57	0.52
23:AZ:41:ARG:HH11	23:AZ:41:ARG:HG3	1.74	0.52
31:BA:116:A:O5'	31:BA:116:A:H8	1.92	0.52
31:BA:188:U:O2'	31:BA:189:U:C5'	2.57	0.52
31:BA:765:G:N1	31:BA:812:C:H2'	2.24	0.52
32:BE:46:LYS:HA	32:BE:49:GLU:HG2	1.90	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:BF:181:ASN:ND2	33:BF:204:LEU:HD12	2.24	0.52
38:BK:20:TYR:CE2	38:BK:75:ARG:HB3	2.44	0.52
39:BL:125:TYR:HD2	39:BL:126:SER:N	2.07	0.52
40:BM:58:ASP:O	40:BM:59:SER:C	2.47	0.52
40:BM:84:GLN:HB3	40:BM:88:LEU:HD23	1.90	0.52
45:BR:17:ARG:HG3	45:BR:17:ARG:NH1	2.23	0.52
45:BR:3:ILE:HG22	45:BR:38:ARG:NH2	2.24	0.52
50:BW:33:ILE:HD11	50:BW:62:LEU:HB3	1.90	0.52
31:BA:186:C:H1'	50:BW:81:LYS:HZ1	1.73	0.52
54:CA:1024:G:H3'	54:CA:1024:G:N3	2.24	0.52
54:CA:1074:G:H2'	54:CA:1075:C:H6	1.74	0.52
54:CA:1101:A:C4'	54:CA:1102:A:O5'	2.30	0.52
54:CA:346:G:H2'	54:CA:346:G:N3	2.23	0.52
52:CB:25:C:H5'	52:CB:26:A:OP2	2.10	0.52
32:CE:200:ILE:HG22	32:CE:201:ILE:N	2.24	0.52
33:CF:11:ARG:O	33:CF:13:GLY:N	2.42	0.52
39:CL:54:ASP:O	39:CL:56:LEU:N	2.42	0.52
42:CO:19:ARG:HH11	42:CO:19:ARG:HB3	1.75	0.52
47:CT:76:LEU:HD11	47:CT:79:SER:HB2	1.91	0.52
48:CU:39:VAL:HA	48:CU:42:ARG:NH1	2.24	0.52
51:CX:25:LYS:HE2	51:CX:26:LYS:O	2.09	0.52
17:D2:35:LEU:C	17:D2:37:VAL:N	2.59	0.52
22:D3:3:HIS:CG	22:D3:4:LYS:H	2.27	0.52
27:D5:55:ARG:C	27:D5:57:VAL:H	2.11	0.52
28:D6:9:LEU:HD22	28:D6:11:LEU:CD2	2.36	0.52
55:DA:1049:C:C2'	55:DA:1050:A:H5''	2.36	0.52
55:DA:1092:C:H2'	55:DA:1093:G:H4'	1.91	0.52
55:DA:1003:G:N2	55:DA:1153:C:C2	2.77	0.52
55:DA:1291:C:H2'	55:DA:1292:U:H6	1.74	0.52
55:DA:2335:A:C8	55:DA:2337:G:C5	2.97	0.52
55:DA:249:C:O2	30:D8:12:LYS:NZ	2.38	0.52
55:DA:2540:C:O2	55:DA:2740:A:H2	1.92	0.52
55:DA:2762:G:C2'	55:DA:2763:G:H5'	2.40	0.52
55:DA:991:C:H5'	55:DA:991:C:H6	1.74	0.52
3:DD:176:ARG:HG2	3:DD:176:ARG:NH1	2.21	0.52
4:DE:34:VAL:HG23	4:DE:48:GLN:CB	2.40	0.52
55:DA:2810:A:HO2'	4:DE:61:ARG:HG3	1.71	0.52
7:DH:147:ASN:N	7:DH:147:ASN:ND2	2.57	0.52
7:DH:87:LEU:HD13	7:DH:148:ILE:CG2	2.38	0.52
56:DI:12:LEU:N	56:DI:12:LEU:HD12	2.23	0.52
56:DI:5:ILE:O	56:DI:6:GLU:CB	2.58	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2780:G:OP2	9:DM:118:LYS:HD3	2.09	0.52
57:DY:50:ARG:HA	57:DY:83:TYR:CE1	2.42	0.52
57:DY:51:LEU:CD1	57:DY:81:VAL:C	2.57	0.52
57:DY:91:LYS:HA	57:DY:94:VAL:CG1	2.39	0.52
13:A0:54:LEU:HD23	13:A0:66:VAL:HG22	1.91	0.52
1:AA:1010:A:H5'	16:A1:62:ILE:CG2	2.39	0.52
1:AA:1174:A:C5'	1:AA:1175:U:H5''	2.39	0.52
1:AA:1654:A:OP1	13:A0:2:ARG:HD3	2.09	0.52
1:AA:1952:A:C6	1:AA:1953:A:N1	2.78	0.52
1:AA:532:A:N7	1:AA:2021:C:H2'	2.24	0.52
1:AA:2061:G:HO2'	1:AA:2062:A:P	2.32	0.52
1:AA:2512:C:H4'	4:AE:122:PHE:CE2	2.44	0.52
1:AA:2531:A:N6	1:AA:2662:A:H61	2.06	0.52
1:AA:2656:U:C6	1:AA:2656:U:C3'	2.93	0.52
1:AA:2788:C:H2'	1:AA:2789:C:O4'	2.08	0.52
1:AA:2862:G:H2'	1:AA:2863:C:H6	1.73	0.52
1:AA:361:G:H2'	1:AA:362:U:O4'	2.09	0.52
1:AA:84:A:C4'	1:AA:85:G:O5'	2.52	0.52
1:AA:865:C:C4'	1:AA:866:A:OP1	2.56	0.52
1:AA:942:G:O2'	1:AA:943:U:H5'	2.08	0.52
1:AA:946:G:O2'	1:AA:947:G:C4'	2.55	0.52
1:AA:973:A:O2'	1:AA:1186:G:N2	2.41	0.52
3:AD:64:ILE:O	3:AD:64:ILE:CG1	2.55	0.52
5:AF:123:LEU:HG	5:AF:125:LEU:HD22	1.91	0.52
9:AM:46:VAL:HG13	9:AM:47:ALA:N	2.23	0.52
1:AA:626:U:O2	11:AO:105:LEU:HG	2.09	0.52
1:AA:627:A:N7	11:AO:84:ASN:ND2	2.54	0.52
1:AA:911:A:H2'	12:AP:9:TYR:OH	2.08	0.52
14:AQ:58:LEU:H	14:AQ:58:LEU:HD23	1.74	0.52
15:AR:107:ASP:O	15:AR:110:ILE:HG22	2.08	0.52
15:AR:96:ARG:HH11	15:AR:96:ARG:HB2	1.72	0.52
21:AV:33:LEU:HG	21:AV:34:ASN:N	2.24	0.52
31:BA:180:U:H2'	31:BA:181:G:H5'	1.90	0.52
31:BA:366:C:O2'	31:BA:367:U:OP1	2.27	0.52
31:BA:412:A:H61	34:BG:35:ARG:HA	1.74	0.52
31:BA:511:C:O4'	34:BG:43:HIS:CD2	2.61	0.52
31:BA:64:G:C4'	31:BA:65:U:H5''	2.18	0.52
31:BA:992:U:O2'	31:BA:993:G:OP2	2.21	0.52
1:AA:882:G:O5'	52:BB:19:G:N7	2.41	0.52
52:BC:51:U:H2'	52:BC:52:G:H8	1.73	0.52
37:BJ:15:ASP:O	37:BJ:19:GLY:HA2	2.09	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:BO:57:LYS:CG	42:BO:67:THR:HG22	2.38	0.52
48:BU:47:THR:HB	48:BU:49:LYS:HE3	1.90	0.52
50:BW:100:ILE:H	50:BW:100:ILE:CD1	2.21	0.52
53:C1:30:C:C6	53:C1:30:C:H3'	2.44	0.52
54:CA:1157:A:N3	54:CA:1157:A:H2'	2.24	0.52
54:CA:1223:C:OP1	54:CA:1224:G:H3'	2.09	0.52
54:CA:412:A:H4'	54:CA:413:G:O5'	2.09	0.52
54:CA:653:A:C1'	38:CK:56:LYS:HD3	2.38	0.52
54:CA:815:A:O2'	54:CA:816:A:P	2.67	0.52
52:CB:42:C:O2'	52:CB:43:C:H5'	2.09	0.52
32:CE:180:LEU:O	32:CE:181:PHE:HB2	2.08	0.52
36:CI:48:LEU:HG	36:CI:57:GLN:HA	1.91	0.52
38:CK:110:ALA:HB3	38:CK:121:ASP:HB3	1.90	0.52
39:CL:118:LYS:NZ	39:CL:118:LYS:CB	2.71	0.52
16:D1:49:HIS:O	16:D1:50:ARG:C	2.45	0.52
16:D1:108:GLU:CB	17:D2:44:LYS:HE3	2.39	0.52
43:CP:3:ARG:HD2	26:D4:34:GLU:OE2	2.08	0.52
55:DA:1099:G:H2'	55:DA:1100:C:O4'	2.09	0.52
55:DA:1288:U:C4'	55:DA:1289:C:OP2	2.57	0.52
55:DA:1289:C:H2'	55:DA:1290:C:C6	2.44	0.52
55:DA:1336:A:H2'	55:DA:1337:G:C8	2.45	0.52
55:DA:1778:U:C2'	55:DA:1784:A:N6	2.69	0.52
52:CD:56:C:H5''	55:DA:2169:A:H8	1.75	0.52
55:DA:1782:C:H1'	55:DA:2609:U:O4'	2.08	0.52
55:DA:2841:C:O2'	55:DA:2842:G:H5'	2.08	0.52
55:DA:484:C:H2'	55:DA:485:C:H6	1.74	0.52
55:DA:614:U:O4	5:DF:175:THR:HG23	2.08	0.52
55:DA:662:G:H5'	11:DO:15:ARG:HA	1.91	0.52
6:DG:101:ILE:HD13	26:D4:9:LEU:HD11	1.91	0.52
21:DV:135:GLU:O	21:DV:136:PHE:HB3	2.09	0.52
57:DY:113:GLN:HG3	57:DY:113:GLN:O	2.09	0.52
57:DY:120:LYS:O	57:DY:121:ASP:HB2	2.09	0.52
26:A4:56:VAL:N	26:A4:59:PHE:HB3	2.24	0.52
28:A6:18:ARG:HE	28:A6:44:ARG:NH1	2.07	0.52
1:AA:1082:U:C3'	1:AA:1082:U:C6	2.92	0.52
1:AA:1342:A:N7	1:AA:1345:C:C4	2.78	0.52
1:AA:1495:A:O2'	1:AA:1496:A:H5'	2.09	0.52
1:AA:1512:G:O2'	1:AA:1513:C:H5'	2.09	0.52
1:AA:1556:C:H2'	1:AA:1557:C:C6	2.44	0.52
1:AA:1838:C:N4	1:AA:1898:U:H2'	2.23	0.52
1:AA:2590:A:H2'	1:AA:2591:C:C6	2.45	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2611:U:O2'	27:A5:3:LYS:HD3	2.09	0.52
1:AA:2768:C:C4	1:AA:2769:C:C5	2.97	0.52
1:AA:299:A:N6	1:AA:322:A:HO2'	2.08	0.52
1:AA:654(S):G:C3'	1:AA:654(T):A:C8	2.92	0.52
1:AA:692:C:O2'	1:AA:693:C:H5'	2.10	0.52
1:AA:796:C:H2'	1:AA:797:C:C6	2.43	0.52
1:AA:80:G:C2'	1:AA:81:G:H5'	2.40	0.52
1:AA:888:C:H1'	1:AA:889:C:OP2	2.10	0.52
3:AD:77:ALA:HB2	3:AD:97:TYR:CG	2.45	0.52
5:AF:53:THR:HG23	5:AF:55:GLY:N	2.24	0.52
7:AH:24:VAL:O	7:AH:24:VAL:HG23	2.10	0.52
7:AH:4:ILE:HD12	7:AH:6:ARG:NE	2.25	0.52
8:AK:37:VAL:HG12	8:AK:38:LEU:N	2.25	0.52
9:AM:91:LEU:CA	9:AM:95:PRO:HB3	2.30	0.52
15:AR:29:ARG:HE	15:AR:44:ASP:HB3	1.73	0.52
20:AU:95:LYS:HB2	20:AU:95:LYS:HZ3	1.74	0.52
21:AV:11:GLU:O	21:AV:12:GLY:O	2.27	0.52
21:AV:145:GLU:OE1	21:AV:174:VAL:CB	2.58	0.52
21:AV:53:ILE:HA	21:AV:70:LEU:CD2	2.39	0.52
53:B1:37:G:H5'	53:B1:37:G:H8	1.74	0.52
31:BA:10:A:O2'	31:BA:11:G:H5'	2.09	0.52
31:BA:128:G:O2'	31:BA:129:U:H5'	2.09	0.52
31:BA:1320:C:H2'	31:BA:1321:C:C6	2.44	0.52
31:BA:337:C:H2'	31:BA:338:A:C8	2.44	0.52
31:BA:688:G:O2'	31:BA:689:C:H5'	2.09	0.52
52:BC:57:G:C2'	52:BC:58:A:H5''	2.39	0.52
34:BG:21:LEU:CD1	34:BG:26:CYS:O	2.57	0.52
36:BI:75:LEU:HD21	36:BI:79:LEU:HD11	1.90	0.52
39:BL:95:LYS:O	39:BL:99:LEU:HB2	2.08	0.52
42:BO:47:LYS:CB	42:BO:48:PRO:CD	2.74	0.52
43:BP:81:LEU:HD13	43:BP:88:ARG:HG2	1.90	0.52
48:BU:19:LYS:O	48:BU:20:ALA:O	2.28	0.52
54:CA:505:G:H5'	54:CA:534:U:H2'	1.91	0.52
54:CA:681:C:O2'	54:CA:682:G:H5'	2.09	0.52
33:CF:182:ILE:HA	33:CF:202:ILE:O	2.09	0.52
40:CM:8:LEU:CD2	40:CM:96:ILE:HG22	2.40	0.52
41:CN:105:VAL:O	41:CN:105:VAL:HG23	2.09	0.52
42:CO:43:VAL:HG23	42:CO:44:THR:N	2.24	0.52
45:CR:82:ILE:HD11	45:CR:88:ARG:CG	2.39	0.52
46:CS:6:LEU:HD11	46:CS:19:ILE:HD13	1.91	0.52
55:DA:1021:A:H8	55:DA:1022:G:C5'	2.21	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1360:A:N1	55:DA:1372:U:C4	2.78	0.52
55:DA:1570:A:H2'	55:DA:1571:A:C8	2.43	0.52
55:DA:1698:A:O2'	55:DA:1699:G:OP2	2.24	0.52
55:DA:1681:G:O2'	55:DA:1762:A:O2'	2.25	0.52
55:DA:1824:G:OP1	3:DD:52:ARG:HD3	2.09	0.52
55:DA:2174:C:H2'	55:DA:2175:C:C6	2.44	0.52
55:DA:2318:G:H2'	55:DA:2319:G:OP1	2.10	0.52
55:DA:2638:G:O2'	55:DA:2639:A:H8	1.92	0.52
55:DA:440:G:H2'	55:DA:441:U:C6	2.44	0.52
55:DA:603:A:C4'	55:DA:604:G:O5'	2.43	0.52
3:DD:35:LYS:HZ1	3:DD:65:ILE:HA	1.73	0.52
4:DE:50:GLY:N	4:DE:77:ILE:HA	2.24	0.52
5:DF:29:ASN:HB3	5:DF:112:MET:HE1	1.91	0.52
5:DF:124:LEU:HD12	5:DF:125:LEU:N	2.24	0.52
6:DG:106:LEU:HD12	6:DG:110:ALA:HB3	1.92	0.52
7:DH:10:PRO:HD2	7:DH:50:VAL:O	2.08	0.52
56:DJ:24:ILE:O	56:DJ:27:LEU:HB2	2.09	0.52
58:DL:115:LEU:HD11	58:DL:117:THR:OG1	2.09	0.52
10:DN:23:ARG:HG3	10:DN:24:VAL:N	2.21	0.52
11:DO:75:ILE:HG12	11:DO:77:ARG:NH1	2.24	0.52
12:DP:71:ASP:C	12:DP:71:ASP:OD2	2.48	0.52
21:DV:152:ALA:CB	21:DV:163:LEU:HD13	2.39	0.52
21:DV:182:LYS:C	21:DV:183:LEU:HD23	2.30	0.52
21:DV:197:ILE:N	21:DV:197:ILE:CD1	2.62	0.52
21:DV:24:LEU:HD21	21:DV:86:VAL:HG23	1.90	0.52
25:DX:21:ALA:O	25:DX:24:LYS:HB3	2.10	0.52
57:DY:119:ALA:O	57:DY:121:ASP:N	2.43	0.52
57:DY:5:ARG:C	57:DY:7:VAL:HG12	2.30	0.52
17:A2:76:LYS:NZ	17:A2:82:ARG:HD3	2.24	0.52
28:A6:31:PRO:C	28:A6:33:LYS:N	2.62	0.52
1:AA:2086:U:H2'	1:AA:2087:G:C8	2.44	0.52
1:AA:2748:A:O2'	1:AA:2749:A:H5'	2.09	0.52
1:AA:84:A:H1'	1:AA:85:G:O4'	2.09	0.52
1:AA:943:U:OP2	11:AO:36:LYS:NZ	2.41	0.52
3:AD:32:SER:HA	3:AD:36:PRO:CD	2.30	0.52
4:AE:41:LYS:HG3	4:AE:42:ASP:OD2	2.09	0.52
4:AE:64:LYS:C	4:AE:66:HIS:H	2.11	0.52
1:AA:2634:G:O3'	4:AE:77:ILE:HG21	2.09	0.52
5:AF:122:LYS:O	5:AF:123:LEU:CB	2.58	0.52
11:AO:83:VAL:O	11:AO:83:VAL:HG13	2.09	0.52
15:AR:50:ILE:HG22	15:AR:62:THR:OG1	2.08	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:AU:61:ILE:N	20:AU:61:ILE:HD12	2.24	0.52
23:AZ:67:ILE:N	23:AZ:68:PRO:CD	2.73	0.52
31:BA:1092:A:C2	31:BA:1183:A:C2	2.97	0.52
31:BA:1314:C:H2'	31:BA:1315:U:C6	2.44	0.52
31:BA:132:C:O2'	31:BA:133:U:H5'	2.10	0.52
31:BA:376:G:H4'	46:BS:5:ARG:HD3	1.91	0.52
31:BA:376:G:O2'	31:BA:377:G:H5'	2.08	0.52
31:BA:9:G:H5'	35:BH:122:GLU:OE2	2.08	0.52
52:BB:19:G:O2'	52:BB:20:U:P	2.68	0.52
33:BF:186:PHE:HD1	33:BF:198:VAL:O	1.91	0.52
34:BG:31:CYS:O	34:BG:33:MET:N	2.37	0.52
34:BG:3:ARG:NH2	34:BG:5:ILE:HD13	2.24	0.52
39:BL:28:VAL:HG22	39:BL:63:ILE:HB	1.92	0.52
39:BL:63:ILE:HG22	39:BL:64:THR:N	2.25	0.52
40:BM:7:LYS:C	40:BM:8:LEU:HD12	2.30	0.52
42:BO:79:GLU:HG3	42:BO:80:HIS:N	2.23	0.52
44:BQ:47:LEU:O	44:BQ:50:LYS:N	2.42	0.52
49:BV:76:PRO:HB2	49:BV:78:ARG:CZ	2.40	0.52
50:BW:32:ALA:O	50:BW:36:LEU:HB2	2.09	0.52
50:BW:71:THR:HG22	50:BW:72:LEU:N	2.24	0.52
54:CA:1347:G:H2'	54:CA:1348:U:OP2	2.10	0.52
54:CA:176:C:O2'	54:CA:177:C:H5'	2.09	0.52
54:CA:115:G:H1	54:CA:312:C:H42	1.55	0.52
54:CA:551:U:H2'	54:CA:552:U:C6	2.44	0.52
54:CA:566:G:C4'	54:CA:567:G:OP1	2.56	0.52
54:CA:748:C:O2'	54:CA:749:C:OP2	2.23	0.52
54:CA:817:C:H4'	54:CA:818:G:OP1	2.07	0.52
54:CA:948:C:O2'	54:CA:949:A:H5'	2.09	0.52
35:CH:78:HIS:HA	38:CK:105:ARG:HB2	1.91	0.52
38:CK:25:ASP:N	38:CK:25:ASP:OD1	2.41	0.52
39:CL:9:ARG:HG2	39:CL:104:ARG:HD2	1.92	0.52
39:CL:53:VAL:HG21	39:CL:92:TYR:CD1	2.45	0.52
41:CN:17:GLY:HA3	41:CN:77:MET:HE3	1.91	0.52
44:CQ:23:ARG:HD2	44:CQ:28:GLY:O	2.10	0.52
55:DA:1653:G:C6	13:D0:9:LYS:HG2	2.44	0.52
27:D5:58:LEU:HD13	27:D5:60:VAL:CB	2.34	0.52
55:DA:1088:A:H3'	55:DA:1088:A:N3	2.24	0.52
55:DA:1266:G:O2'	55:DA:2012:G:N1	2.41	0.52
55:DA:142:G:H2'	55:DA:143:C:H6	1.74	0.52
55:DA:2096:U:H2'	55:DA:2097:C:C6	2.44	0.52
55:DA:2112:G:C2'	55:DA:2113:U:H5''	2.40	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2656:U:N3	55:DA:2665:A:C2	2.76	0.52
3:DD:35:LYS:HZ3	3:DD:104:TYR:HD1	1.57	0.52
3:DD:75:ILE:HD13	3:DD:99:ASP:OD1	2.10	0.52
55:DA:2729:G:C1'	4:DE:187:ALA:HB2	2.35	0.52
4:DE:197:ILE:HD11	4:DE:199:ARG:NH1	2.23	0.52
4:DE:63:LEU:O	4:DE:64:LYS:HB2	2.10	0.52
43:CP:68:GLY:HA3	6:DG:116:ASP:OD2	2.09	0.52
56:DI:18:LEU:O	56:DI:19:GLU:C	2.48	0.52
56:DJ:13:SER:OG	56:DJ:17:VAL:HG13	2.10	0.52
8:DK:3:VAL:HB	8:DK:37:VAL:O	2.09	0.52
8:DK:77:LEU:HD11	8:DK:140:LEU:HA	1.89	0.52
58:DL:109:LYS:CB	58:DL:120:LEU:HD21	2.40	0.52
9:DM:43:THR:HB	9:DM:46:VAL:HG12	1.89	0.52
11:DO:147:LEU:O	11:DO:148:LEU:CB	2.57	0.52
55:DA:389:G:H22	11:DO:72:PRO:CG	2.22	0.52
19:DT:14:SER:H	19:DT:17:ALA:HB3	1.74	0.52
20:DU:44:ILE:CD1	20:DU:45:VAL:HG23	2.39	0.52
21:DV:118:GLN:CA	21:DV:118:GLN:HE21	1.97	0.52
21:DV:174:VAL:O	21:DV:175:VAL:CB	2.57	0.52
55:DA:896:A:H2	21:DV:178:GLU:OE2	1.84	0.52
57:DY:132:ASP:C	57:DY:134:LEU:HD22	2.29	0.52
16:A1:90:VAL:HG13	17:A2:39:LEU:CB	2.40	0.52
28:A6:10:LEU:HA	30:A8:34:TRP:CH2	2.44	0.52
28:A6:15:GLU:CD	28:A6:44:ARG:NH2	2.63	0.52
1:AA:1059:G:H2'	1:AA:1060:U:C6	2.44	0.52
1:AA:1131:G:H4'	9:AM:82:LEU:HB2	1.91	0.52
1:AA:1174:A:H3'	1:AA:1175:U:H4'	1.90	0.52
1:AA:1365:A:OP2	23:AZ:3:LYS:HB2	2.09	0.52
1:AA:1666:G:H2'	1:AA:1667:G:O4'	2.09	0.52
1:AA:1991:U:H2'	1:AA:1992:G:H5'	1.91	0.52
1:AA:2290:G:H4'	1:AA:2381:C:O2'	2.09	0.52
1:AA:2861:G:H2'	1:AA:2862:G:H8	1.74	0.52
1:AA:387:U:P	1:AA:387:U:C6	3.03	0.52
1:AA:816:C:O2'	1:AA:817:C:H5'	2.10	0.52
2:AB:24:G:H1'	2:AB:27:C:N4	2.24	0.52
3:AD:48:ARG:NH1	3:AD:48:ARG:HG3	2.23	0.52
4:AE:95:ILE:H	4:AE:95:ILE:HD12	1.73	0.52
5:AF:121:GLY:O	5:AF:122:LYS:HD3	2.09	0.52
6:AG:130:ASN:OD1	6:AG:160:VAL:HA	2.09	0.52
11:AO:9:ASN:CB	11:AO:10:PRO:HD2	2.40	0.52
14:AQ:69:VAL:HG13	14:AQ:101:LEU:CD2	2.39	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:AU:15:VAL:CB	20:AU:22:GLY:HA3	2.23	0.52
20:AU:43:ASN:HB3	20:AU:64:GLU:HA	1.89	0.52
31:BA:1058:G:H2'	31:BA:1059:C:O4'	2.09	0.52
31:BA:1260:C:O5'	31:BA:1284:C:H4'	2.09	0.52
37:BJ:63:LYS:HG3	37:BJ:64:GLN:OE1	2.09	0.52
35:BH:78:HIS:CB	38:BK:104:ARG:HD2	2.34	0.52
38:BK:6:ILE:HD12	38:BK:6:ILE:N	2.25	0.52
39:BL:3:GLN:CD	39:BL:20:ARG:HH12	2.12	0.52
41:BN:105:VAL:O	41:BN:105:VAL:HG23	2.09	0.52
42:BO:47:LYS:HB3	42:BO:48:PRO:HD3	1.84	0.52
42:BO:46:LYS:CG	42:BO:47:LYS:N	2.71	0.52
31:BA:1316:G:H4'	44:BQ:18:VAL:CG1	2.39	0.52
54:CA:1452:C:H2'	54:CA:1453:G:OP2	2.10	0.52
54:CA:37:U:O2'	54:CA:38:G:H5'	2.10	0.52
54:CA:522:C:O2'	54:CA:523:A:H5'	2.09	0.52
8:AK:82:ARG:NE	54:CA:56:U:H4'	2.20	0.52
34:CG:141:ARG:HB3	34:CG:142:PRO:HD2	1.92	0.52
35:CH:107:ARG:HG2	35:CH:108:ALA:N	2.24	0.52
37:CJ:115:ARG:O	37:CJ:118:VAL:HG13	2.09	0.52
39:CL:48:GLU:N	39:CL:49:PRO:CD	2.71	0.52
44:CQ:24:CYS:HB3	44:CQ:40:CYS:HB3	1.92	0.52
30:D8:51:ALA:N	30:D8:53:PRO:HD2	2.24	0.52
55:DA:1272:A:H5'	55:DA:1273:U:OP2	2.09	0.52
55:DA:1280:G:H2'	55:DA:1281:G:H5''	1.91	0.52
55:DA:1445:C:H2'	55:DA:1446:C:H6	1.74	0.52
55:DA:2574:G:O2'	4:DE:143:ASN:HB3	2.10	0.52
55:DA:502:A:H2'	55:DA:503:A:H5'	1.90	0.52
55:DA:506:G:O3'	55:DA:507:A:H8	1.92	0.52
2:DB:21:G:H2'	2:DB:22:U:O4'	2.10	0.52
3:DD:35:LYS:CE	3:DD:104:TYR:CD1	2.93	0.52
4:DE:174:ASP:OD2	4:DE:175:VAL:N	2.38	0.52
7:DH:18:GLU:O	7:DH:24:VAL:HA	2.10	0.52
8:DK:101:LEU:C	8:DK:101:LEU:HD23	2.30	0.52
58:DL:14:ALA:CA	58:DL:49:GLY:HA3	2.15	0.52
58:DL:53:VAL:HG12	58:DL:72:PRO:CG	2.38	0.52
9:DM:131:GLN:NE2	9:DM:132:ALA:H	2.07	0.52
9:DM:39:ARG:NH1	9:DM:41:ASP:OD1	2.42	0.52
12:DP:31:ASP:N	12:DP:106:VAL:O	2.42	0.52
12:DP:35:VAL:HG22	12:DP:36:ALA:N	2.25	0.52
55:DA:957:A:OP1	12:DP:76:LYS:HD2	2.09	0.52
14:DQ:34:HIS:HB3	14:DQ:53:SER:HB3	1.92	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:15:GLU:O	57:DY:16:ASN:HB2	2.09	0.52
57:DY:23:SER:OG	57:DY:68:LEU:HB3	2.09	0.52
17:A2:81:TYR:HB2	17:A2:83:ARG:NH1	2.25	0.52
1:AA:2815:C:O2'	27:A5:43:HIS:CD2	2.62	0.52
1:AA:2016:U:H1'	27:A5:6:VAL:HG22	1.91	0.52
1:AA:1024:G:O6	1:AA:1025:G:N1	2.42	0.52
1:AA:1652:A:H3'	1:AA:1653:G:C8	2.45	0.52
1:AA:2081:C:O2'	1:AA:2082:A:H5'	2.10	0.52
1:AA:2250:G:H2'	1:AA:2496:C:OP1	2.09	0.52
1:AA:2637:U:H5''	1:AA:2638:G:OP2	2.10	0.52
1:AA:607:U:O2	1:AA:620:G:C8	2.63	0.52
1:AA:788:A:O2'	1:AA:789:A:P	2.68	0.52
1:AA:80:G:O2'	1:AA:81:G:H5'	2.09	0.52
1:AA:954:G:N3	1:AA:954:G:H2'	2.24	0.52
2:AB:39:A:N1	2:AB:44:G:C6	2.78	0.52
2:AB:8:U:H2'	2:AB:9:G:C8	2.45	0.52
3:AD:176:ARG:NH1	3:AD:176:ARG:HG2	2.20	0.52
3:AD:34:VAL:C	3:AD:35:LYS:HG3	2.29	0.52
1:AA:2723:C:OP1	4:AE:109:LYS:HE2	2.09	0.52
4:AE:9:VAL:CG2	4:AE:10:GLY:N	2.61	0.52
6:AG:111:LEU:CD1	6:AG:120:LEU:HD11	2.39	0.52
6:AG:142:PRO:HG2	6:AG:143:GLU:OE2	2.10	0.52
1:AA:2757:A:H2	7:AH:64:LEU:HD23	1.75	0.52
7:AH:4:ILE:HD11	7:AH:7:LEU:CG	2.40	0.52
1:AA:2467:C:H4'	12:AP:123:HIS:CD2	2.45	0.52
14:AQ:110:LEU:HD22	14:AQ:111:GLU:H	1.74	0.52
19:AT:65:ARG:HG3	19:AT:65:ARG:NH1	2.21	0.52
53:B1:43:U:H2'	53:B1:44:U:H5'	1.91	0.52
31:BA:1026:G:C6	31:BA:1036:G:N2	2.77	0.52
31:BA:1027:C:O2'	31:BA:1028:C:P	2.67	0.52
31:BA:105:G:H2'	31:BA:106:C:C6	2.44	0.52
31:BA:1098:C:H2'	31:BA:1099:G:O4'	2.08	0.52
31:BA:1343:G:H2'	31:BA:1344:C:H6	1.68	0.52
31:BA:1347:G:HO2'	31:BA:1373:G:H1	1.57	0.52
31:BA:533:A:O2'	31:BA:534:U:OP1	2.24	0.52
32:BE:78:GLN:HA	32:BE:94:ASN:OD1	2.09	0.52
33:BF:16:ARG:HH11	33:BF:16:ARG:CA	2.22	0.52
35:BH:9:LYS:O	35:BH:33:VAL:HG23	2.08	0.52
37:BJ:20:ASP:O	37:BJ:23:VAL:HB	2.10	0.52
40:BM:23:ILE:HA	40:BM:26:ALA:HB3	1.92	0.52
42:BO:37:CYS:HA	42:BO:57:LYS:O	2.09	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:1329:A:H4'	43:BP:24:GLY:O	2.10	0.52
43:BP:22:ILE:CB	43:BP:25:ILE:HG12	2.15	0.52
43:BP:45:VAL:HA	43:BP:48:LEU:HD22	1.91	0.52
43:BP:8:GLU:OE1	43:BP:22:ILE:HG12	2.10	0.52
50:BW:67:ALA:HA	50:BW:73:HIS:CA	2.40	0.52
53:C1:37:G:H3'	53:C1:38:U:C6	2.44	0.52
54:CA:134:A:N6	46:CS:25:ARG:NH1	2.58	0.52
54:CA:152:A:N6	54:CA:169:C:N4	2.56	0.52
54:CA:153:C:H2'	54:CA:154:C:C6	2.44	0.52
54:CA:501:C:H2'	54:CA:502:G:C8	2.41	0.52
54:CA:671:G:H2'	54:CA:672:U:H6	1.74	0.52
54:CA:766:A:H61	54:CA:1511:G:H1'	1.75	0.52
54:CA:965:A:H4'	54:CA:966:G:C5'	2.40	0.52
32:CE:55:PHE:HA	32:CE:58:ILE:CG1	2.39	0.52
32:CE:82:ARG:NH1	32:CE:86:GLU:OE2	2.43	0.52
54:CA:1055:A:H4'	33:CF:161:GLU:OE1	2.10	0.52
33:CF:62:ASP:HA	33:CF:97:LYS:HD2	1.91	0.52
40:CM:3:LYS:O	40:CM:100:THR:HA	2.10	0.52
50:CW:35:THR:O	50:CW:39:LYS:HG3	2.10	0.52
50:CW:89:ARG:HH21	50:CW:89:ARG:HG3	1.74	0.52
16:D1:34:LYS:HA	16:D1:34:LYS:CE	2.31	0.52
27:D5:13:LYS:HG2	27:D5:16:ARG:HH21	1.74	0.52
55:DA:2346:A:O3'	28:D6:39:TYR:OH	2.27	0.52
30:D8:37:SER:O	30:D8:40:GLU:HB3	2.09	0.52
55:DA:119:A:H4'	55:DA:120:U:OP1	2.09	0.52
55:DA:1292:U:H2'	55:DA:1293:C:H6	1.73	0.52
55:DA:1329:U:H5''	55:DA:1330:C:C5	2.45	0.52
55:DA:1871:A:H2'	55:DA:1872:A:C8	2.44	0.52
55:DA:1964:G:C8	55:DA:1964:G:OP1	2.62	0.52
55:DA:2552:U:H2'	55:DA:2554:U:H5''	1.91	0.52
55:DA:729:G:OP2	3:DD:13:ARG:NH1	2.41	0.52
3:DD:65:ILE:H	3:DD:65:ILE:HD12	1.74	0.52
6:DG:104:GLU:O	6:DG:108:ASN:HB2	2.09	0.52
6:DG:41:GLN:HE21	6:DG:60:LEU:HD12	1.75	0.52
57:DY:135:ARG:HB2	56:DJ:19:GLU:CD	2.30	0.52
58:DL:102:GLU:HG2	58:DL:103:GLN:H	1.75	0.52
58:DL:105:LEU:C	58:DL:107:ILE:N	2.62	0.52
58:DL:11:GLN:CB	58:DL:41:PHE:CZ	2.91	0.52
9:DM:131:GLN:N	9:DM:131:GLN:HE21	2.07	0.52
9:DM:35:ARG:HB2	9:DM:42:TRP:CZ3	2.45	0.52
9:DM:98:VAL:HG13	9:DM:99:LEU:N	2.25	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:DN:107:ARG:CZ	15:DR:36:GLU:HB3	2.40	0.52
10:DN:64:ARG:O	10:DN:82:ASN:HA	2.09	0.52
14:DQ:71:ARG:CG	14:DQ:104:GLY:HA2	2.32	0.52
57:DY:24:PHE:CZ	57:DY:88:ALA:CB	2.84	0.52
57:DY:27:VAL:CA	57:DY:111:LEU:N	2.68	0.52
22:A3:72:ARG:NH1	22:A3:72:ARG:HG3	2.22	0.52
1:AA:1114:G:H2'	1:AA:1115:G:C8	2.44	0.52
1:AA:1254:A:H5'	1:AA:1255:U:H5'	1.92	0.52
1:AA:1838:C:H5'	1:AA:1839:G:OP1	2.09	0.52
1:AA:2461:C:H2'	1:AA:2462:U:H6	1.74	0.52
1:AA:2646:C:H6	1:AA:2646:C:O5'	1.93	0.52
1:AA:2688:U:H3'	1:AA:2688:U:O2	2.10	0.52
1:AA:483:A:H3'	1:AA:484:C:H6	1.74	0.52
1:AA:527:C:O2	1:AA:2779:U:C5	2.63	0.52
1:AA:58:G:H2'	1:AA:59:U:C6	2.45	0.52
1:AA:691:C:H2'	1:AA:692:C:H6	1.74	0.52
1:AA:946:G:O6	1:AA:972:G:C2	2.62	0.52
2:AB:27:C:C4	2:AB:28:C:C4	2.98	0.52
3:AD:122:ASP:OD1	3:AD:123:ALA:N	2.42	0.52
4:AE:48:GLN:HE22	4:AE:64:LYS:HE3	1.75	0.52
4:AE:60:ASN:C	4:AE:62:PRO:CD	2.67	0.52
5:AF:46:ARG:HG2	5:AF:46:ARG:NH1	2.18	0.52
11:AO:101:VAL:HG13	11:AO:102:ARG:N	2.25	0.52
11:AO:148:LEU:N	11:AO:148:LEU:HD23	2.24	0.52
12:AP:42:ILE:HA	12:AP:46:GLN:OE1	2.09	0.52
12:AP:42:ILE:HD13	12:AP:97:VAL:HB	1.91	0.52
12:AP:82:ARG:NH1	12:AP:82:ARG:CG	2.70	0.52
14:AQ:11:LYS:CD	14:AQ:15:ARG:HH21	2.23	0.52
15:AR:53:ARG:HD3	15:AR:60:THR:OG1	2.09	0.52
19:AT:64:LYS:HZ3	19:AT:73:ARG:NH2	2.08	0.52
21:AV:37:VAL:HG23	21:AV:38:TYR:N	2.25	0.52
25:AX:4:LEU:CD2	25:AX:56:VAL:HG13	2.39	0.52
31:BA:1144:G:N2	31:BA:1146:A:H62	2.08	0.52
31:BA:1064:G:O2'	31:BA:1190:G:N2	2.42	0.52
31:BA:338:A:H61	31:BA:351:G:H1	1.58	0.52
31:BA:389:A:H3'	31:BA:390:C:H6	1.74	0.52
31:BA:641:U:H4'	31:BA:642:A:OP1	2.10	0.52
31:BA:794:A:C2	31:BA:795:C:C2	2.98	0.52
31:BA:960:U:O4	31:BA:1225:A:H1'	2.10	0.52
52:BB:46:G:H5''	52:BB:47:U:OP2	2.10	0.52
33:BF:16:ARG:HH11	33:BF:16:ARG:HA	1.74	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:BF:57:ILE:HG23	33:BF:64:VAL:HG13	1.92	0.52
37:BJ:18:TYR:CE2	37:BJ:59:LEU:HB2	2.43	0.52
38:BK:38:ILE:HD11	38:BK:118:VAL:O	2.10	0.52
43:BP:70:LEU:HD13	43:BP:71:ARG:H	1.73	0.52
48:BU:30:ASP:C	48:BU:32:ARG:H	2.13	0.52
49:BV:5:LEU:HD13	49:BV:6:LYS:N	2.25	0.52
31:BA:1453:G:H3'	50:BW:39:LYS:NZ	2.23	0.52
54:CA:1151:A:H1'	40:CM:39:PRO:CB	2.39	0.52
54:CA:1199:U:H4'	40:CM:54:PHE:CZ	2.44	0.52
54:CA:723:U:OP1	53:C1:37:G:O2'	2.28	0.52
52:CB:10:G:O2'	52:CB:11:C:OP1	2.28	0.52
52:CD:18:G:HO2'	52:CD:19:G:P	2.33	0.52
36:CI:42:GLU:O	36:CI:44:GLY:N	2.43	0.52
36:CI:79:LEU:O	36:CI:85:VAL:HG11	2.10	0.52
54:CA:640:A:O2'	38:CK:115:SER:HB3	2.09	0.52
42:CO:28:LYS:C	42:CO:30:ALA:N	2.61	0.52
42:CO:33:ARG:HE	42:CO:33:ARG:HA	1.74	0.52
43:CP:15:VAL:HG12	43:CP:19:LEU:HD21	1.92	0.52
6:DG:6:ALA:CB	26:D4:23:GLU:HG3	2.39	0.52
30:D8:56:GLU:N	30:D8:56:GLU:OE1	2.43	0.52
55:DA:1048:A:C5	55:DA:1049:C:C5	2.98	0.52
55:DA:1084:A:H1'	57:DY:53:VAL:CG2	2.40	0.52
55:DA:1079:C:C4	55:DA:1088:A:C8	2.97	0.52
55:DA:1936:A:H4'	55:DA:1937:A:OP2	2.08	0.52
55:DA:2723:C:H5''	13:D0:1:MET:CG	2.40	0.52
55:DA:2891:G:H5'	55:DA:2892:A:OP1	2.09	0.52
55:DA:440:G:H2'	55:DA:441:U:H6	1.75	0.52
55:DA:654(M):C:C2'	55:DA:654(N):G:OP1	2.58	0.52
55:DA:895:U:C5'	55:DA:896:A:OP2	2.58	0.52
4:DE:63:LEU:O	4:DE:64:LYS:CB	2.58	0.52
5:DF:129:PHE:O	5:DF:142:TRP:CD1	2.63	0.52
5:DF:155:LEU:HD13	5:DF:174:VAL:HG22	1.92	0.52
6:DG:16:ARG:CG	6:DG:16:ARG:HH11	2.15	0.52
7:DH:161:GLY:O	7:DH:163:TYR:HD1	1.93	0.52
8:DK:112:LYS:O	8:DK:113:ARG:HB2	2.09	0.52
58:DL:105:LEU:HD12	58:DL:106:GLU:HB2	1.91	0.52
58:DL:112:MET:CE	58:DL:118:THR:O	2.57	0.52
11:DO:29:LYS:HD2	11:DO:30:THR:HG22	1.91	0.52
11:DO:90:ARG:NH2	11:DO:91:PHE:HB3	2.25	0.52
15:DR:110:ILE:HG22	15:DR:111:ARG:NH1	2.24	0.52
57:DY:118:THR:OG1	57:DY:119:ALA:N	2.39	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:129:PRO:C	57:DY:130:THR:OG1	2.48	0.52
57:DY:142:LEU:C	57:DY:142:LEU:CD2	2.78	0.52
57:DY:98:LYS:CB	57:DY:102:LYS:HE3	2.40	0.52
23:DZ:89:GLU:OE2	23:DZ:93:GLU:OE1	2.28	0.52
13:A0:104:ARG:HH11	13:A0:104:ARG:CB	2.23	0.52
26:A4:35:VAL:C	26:A4:37:SER:N	2.59	0.52
1:AA:1024:G:C6	1:AA:1025:G:C6	2.98	0.52
1:AA:1427:A:OP1	1:AA:1427:A:H8	1.93	0.52
1:AA:1488:G:H5'	1:AA:1489:U:OP2	2.10	0.52
1:AA:1654:A:H2	4:AE:113:PHE:CD2	2.28	0.52
1:AA:1856:G:H2'	1:AA:1857:G:C5'	2.40	0.52
1:AA:1833:U:O2'	1:AA:1969:A:N1	2.39	0.52
1:AA:2112:G:O2'	1:AA:2113:U:H5''	2.10	0.52
1:AA:2238:G:H5''	1:AA:2239:G:OP1	2.10	0.52
1:AA:2468:G:C2'	1:AA:2469:A:H5''	2.40	0.52
1:AA:2612:C:C5	1:AA:2613:U:C5	2.97	0.52
1:AA:332:A:H4'	1:AA:333:G:OP1	2.10	0.52
1:AA:384:U:O2'	1:AA:385:C:H5'	2.06	0.52
1:AA:753:C:H2'	1:AA:754:C:H6	1.75	0.52
1:AA:839:U:H2'	1:AA:840:C:C6	2.45	0.52
4:AE:47:VAL:HG12	4:AE:48:GLN:N	2.22	0.52
5:AF:164:ARG:HG2	5:AF:164:ARG:HH11	1.74	0.52
8:AK:101:LEU:HG	8:AK:109:ILE:CD1	2.35	0.52
9:AM:30:ILE:HG22	9:AM:34:LEU:HD23	1.90	0.52
12:AP:32:TYR:CZ	12:AP:111:GLU:HB2	2.45	0.52
12:AP:89:ASN:ND2	12:AP:89:ASN:N	2.54	0.52
14:AQ:78:LEU:HD11	14:AQ:107:GLU:HG3	1.92	0.52
15:AR:12:SER:C	15:AR:14:TYR:H	2.13	0.52
18:AS:47:VAL:O	18:AS:50:VAL:HG12	2.10	0.52
31:BA:436:C:H2'	31:BA:437:U:C6	2.45	0.52
52:BB:11:C:O5'	52:BB:11:C:H6	1.93	0.52
31:BA:966:G:C2	52:BC:34:G:H5'	2.45	0.52
32:BE:216:SER:C	32:BE:218:ALA:H	2.12	0.52
34:BG:128:VAL:HG12	34:BG:129:ASN:HD22	1.74	0.52
35:BH:9:LYS:CB	35:BH:112:LEU:HD11	2.40	0.52
42:BO:18:VAL:O	42:BO:19:ARG:HB3	2.10	0.52
42:BO:60:LEU:HD21	42:BO:66:VAL:HG23	1.92	0.52
42:BO:55:VAL:HG13	42:BO:68:ALA:O	2.08	0.52
45:BR:64:ARG:HH12	45:BR:68:ARG:HH21	1.57	0.52
51:BX:9:ARG:NH2	51:BX:10:ARG:HE	2.07	0.52
54:CA:288:A:H2'	54:CA:289:G:H4'	1.91	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:297:G:H4'	54:CA:557:G:H4'	1.91	0.52
54:CA:67:C:H2'	54:CA:68:G:H8	1.71	0.52
54:CA:74:C:C4	54:CA:75:C:C5	2.98	0.52
54:CA:817:C:H1'	54:CA:819:A:H5'	1.91	0.52
52:CD:1:G:H2'	52:CD:2:C:OP1	2.10	0.52
32:CE:67:THR:HG21	32:CE:155:LEU:CD1	2.39	0.52
32:CE:163:PHE:HD2	32:CE:185:ILE:HG13	1.75	0.52
32:CE:44:LEU:N	32:CE:44:LEU:HD12	2.24	0.52
35:CH:10:MET:HB2	35:CH:32:VAL:HG22	1.91	0.52
37:CJ:115:ARG:HB2	37:CJ:118:VAL:HG13	1.91	0.52
37:CJ:38:LEU:O	37:CJ:42:ILE:HG13	2.09	0.52
37:CJ:87:VAL:CG2	37:CJ:154:TYR:HB2	2.40	0.52
38:CK:85:ARG:HD3	38:CK:88:LYS:HG2	1.92	0.52
38:CK:85:ARG:NE	38:CK:87:SER:O	2.43	0.52
54:CA:644:G:C4'	38:CK:92:ARG:HH21	2.21	0.52
41:CN:48:ILE:HD11	41:CN:64:ALA:HA	1.92	0.52
42:CO:38:THR:HG23	42:CO:57:LYS:HB3	1.92	0.52
49:CV:83:HIS:O	49:CV:85:LYS:N	2.43	0.52
50:CW:35:THR:O	50:CW:38:LYS:HB2	2.09	0.52
13:D0:1:MET:O	13:D0:2:ARG:CB	2.57	0.52
27:D5:51:TYR:O	27:D5:56:LYS:HE2	2.09	0.52
28:D6:41:PRO:HG3	28:D6:47:THR:OG1	2.10	0.52
55:DA:1103:A:H5'	55:DA:1103:A:C8	2.44	0.52
55:DA:1936:A:C2	55:DA:1945:G:C6	2.98	0.52
55:DA:286:C:O2'	55:DA:287:C:H5'	2.10	0.52
55:DA:606:U:H4'	55:DA:658:C:H4'	1.92	0.52
55:DA:782:A:H4'	55:DA:783:A:O5'	2.10	0.52
55:DA:90:U:H1'	55:DA:91:A:C8	2.44	0.52
4:DE:14:ILE:O	4:DE:15:PHE:HB2	2.08	0.52
5:DF:89:VAL:HG12	5:DF:90:PHE:N	2.25	0.52
6:DG:33:ARG:O	6:DG:162:THR:HG23	2.10	0.52
7:DH:143:GLN:HE22	7:DH:147:ASN:ND2	2.04	0.52
7:DH:26:VAL:O	7:DH:27:LYS:O	2.28	0.52
58:DL:144:VAL:O	58:DL:145:LYS:HB2	2.09	0.52
9:DM:6:PRO:CG	9:DM:41:ASP:HB2	2.40	0.52
10:DN:5:GLN:O	10:DN:20:MET:HE2	2.08	0.52
11:DO:39:LYS:HA	11:DO:45:LEU:HD13	1.91	0.52
55:DA:911:A:H2'	12:DP:9:TYR:OH	2.10	0.52
19:DT:44:GLU:HB3	19:DT:49:VAL:O	2.09	0.52
19:DT:3:THR:HG22	19:DT:6:ASP:OD2	2.09	0.52
24:DW:28:LYS:NZ	24:DW:56:GLN:NE2	2.58	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:DZ:70:VAL:O	23:DZ:74:VAL:HG23	2.10	0.52
23:DZ:80:LEU:HD13	23:DZ:80:LEU:N	2.25	0.52
1:AA:1248:G:H2'	16:A1:3:ARG:HA	1.90	0.52
17:A2:47:VAL:HG22	17:A2:47:VAL:O	2.10	0.52
17:A2:70:ILE:O	17:A2:71:LEU:CB	2.57	0.52
28:A6:31:PRO:HB2	28:A6:33:LYS:CG	2.38	0.52
1:AA:1055:G:H2'	1:AA:1056:G:C5'	2.39	0.52
1:AA:1209:G:O3'	1:AA:1212:G:H5'	2.09	0.52
1:AA:1278:A:H4'	13:A0:34:ILE:CD1	2.40	0.52
1:AA:1494:A:H2'	1:AA:1495:A:H8	1.75	0.52
1:AA:1668:A:H2'	1:AA:1674:G:N7	2.25	0.52
1:AA:2320:A:H61	1:AA:2333:A:H2'	1.74	0.52
1:AA:2393:A:OP2	30:A8:30:ARG:HB2	2.10	0.52
1:AA:2786:U:O4'	4:AE:64:LYS:HA	2.09	0.52
1:AA:389:G:O6	11:AO:70:GLN:HB3	2.10	0.52
1:AA:654(R):C:C2	1:AA:654(S):G:C8	2.98	0.52
1:AA:674:G:H1'	5:AF:74:ARG:CD	2.28	0.52
1:AA:956:G:C5'	1:AA:957:A:OP2	2.58	0.52
3:AD:171:ASP:O	3:AD:187:GLY:N	2.41	0.52
3:AD:130:ALA:HB2	3:AD:192:THR:HB	1.92	0.52
3:AD:35:LYS:HA	3:AD:64:ILE:CG2	2.40	0.52
3:AD:35:LYS:CG	3:AD:64:ILE:N	2.63	0.52
4:AE:129:HIS:O	4:AE:130:GLY:C	2.48	0.52
7:AH:86:GLU:HA	7:AH:132:ARG:CB	2.37	0.52
8:AK:82:ARG:CD	8:AK:146:ALA:HB2	2.40	0.52
9:AM:103:VAL:O	9:AM:106:MET:N	2.37	0.52
10:AN:104:ARG:CB	10:AN:104:ARG:HH11	2.17	0.52
18:AS:47:VAL:HA	18:AS:50:VAL:HG12	1.92	0.52
18:AS:73:ALA:HB3	18:AS:106:ILE:CG1	2.36	0.52
24:AW:31:GLU:O	24:AW:34:GLU:HB2	2.10	0.52
53:B1:36:G:H2'	53:B1:37:G:C5'	2.31	0.52
31:BA:1127:G:N2	31:BA:1145:C:C2	2.78	0.52
31:BA:1256:A:H62	31:BA:1278:U:H6	1.58	0.52
31:BA:812:C:O2'	31:BA:813:U:OP2	2.27	0.52
31:BA:92:G:H2'	31:BA:93:U:O4'	2.10	0.52
52:BB:24:G:H2'	52:BB:24:G:N3	2.25	0.52
32:BE:162:ILE:O	32:BE:185:ILE:HG12	2.10	0.52
32:BE:5:ILE:HD13	32:BE:55:PHE:HB3	1.92	0.52
32:BE:5:ILE:O	32:BE:6:THR:HG22	2.09	0.52
34:BG:173:TRP:O	34:BG:186:LEU:HB2	2.09	0.52
34:BG:190:ASP:C	34:BG:192:GLU:N	2.63	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:BI:2:ARG:O	36:BI:66:GLU:HA	2.10	0.52
38:BK:127:LEU:H	38:BK:127:LEU:HD22	1.75	0.52
43:BP:56:LEU:HD13	43:BP:56:LEU:C	2.29	0.52
44:BQ:44:LEU:HD12	44:BQ:44:LEU:C	2.29	0.52
50:BW:29:LYS:O	50:BW:33:ILE:HG12	2.09	0.52
50:BW:50:GLU:HG3	50:BW:51:GLU:H	1.75	0.52
54:CA:1128:C:H1'	54:CA:1146:A:H61	1.75	0.52
54:CA:1161:C:H2'	54:CA:1162:C:C6	2.45	0.52
54:CA:191(B):G:O2'	54:CA:191(C):G:H5'	2.09	0.52
54:CA:262:A:H2'	54:CA:263:A:C8	2.45	0.52
54:CA:563:A:H1'	54:CA:566:G:O2'	2.10	0.52
54:CA:595:G:N1	54:CA:641:U:H2'	2.24	0.52
54:CA:629:G:H5'	54:CA:630:G:OP2	2.07	0.52
54:CA:707:C:O2'	54:CA:708:C:H5'	2.10	0.52
52:CC:72:C:H2'	52:CC:73:A:H5'	1.92	0.52
52:CD:61:C:H2'	52:CD:62:C:C6	2.45	0.52
34:CG:49:ARG:O	34:CG:50:ARG:C	2.48	0.52
35:CH:12:LEU:HD23	35:CH:13:ILE:H	1.74	0.52
35:CH:153:LYS:NZ	35:CH:153:LYS:HB2	2.25	0.52
35:CH:78:HIS:HE1	35:CH:143:ARG:H	1.56	0.52
38:CK:6:ILE:CD1	38:CK:6:ILE:H	2.20	0.52
46:CS:8:ARG:NH2	46:CS:15:PRO:HG3	2.24	0.52
26:D4:42:PHE:O	26:D4:44:THR:N	2.38	0.52
55:DA:84:A:N6	55:DA:102:G:H1'	2.24	0.52
55:DA:1800:C:OP1	3:DD:266:SER:OG	2.27	0.52
55:DA:2155:G:H2'	55:DA:2156:G:O4'	2.10	0.52
55:DA:2243:U:H2'	55:DA:2244:U:C6	2.45	0.52
55:DA:2331:G:O3'	22:D3:43:THR:HG22	2.09	0.52
55:DA:279:C:H2'	55:DA:280:C:H6	1.74	0.52
55:DA:30:G:H2'	55:DA:31:C:C6	2.44	0.52
55:DA:457:A:H1'	55:DA:459:U:C6	2.45	0.52
55:DA:686:G:O6	29:D7:12:ARG:NH1	2.40	0.52
55:DA:859:G:C2'	55:DA:860:U:OP2	2.57	0.52
2:DB:14:U:H4'	2:DB:106:G:N2	2.25	0.52
2:DB:31:C:C2	2:DB:32:C:C5	2.98	0.52
2:DB:30:C:N3	2:DB:31:C:H1'	2.25	0.52
3:DD:69:ARG:HD3	3:DD:105:ILE:CD1	2.37	0.52
6:DG:136:ARG:O	6:DG:137:GLU:C	2.47	0.52
56:DI:29:GLU:CD	56:DJ:6:GLU:CD	2.68	0.52
8:DK:110:ASP:HB3	8:DK:112:LYS:HG3	1.92	0.52
11:DO:103:ALA:HB3	11:DO:105:LEU:CD1	2.39	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:DO:85:LEU:HA	11:DO:88:LEU:CD2	2.34	0.52
12:DP:2:LEU:O	12:DP:3:MET:C	2.49	0.52
20:DU:47:LYS:HG2	20:DU:60:PHE:CD1	2.45	0.52
21:DV:185:GLU:OE1	21:DV:185:GLU:HA	2.09	0.52
27:A5:42:PRO:O	27:A5:44:THR:HG23	2.10	0.52
11:AO:49:ARG:HD2	30:A8:58:ILE:HG22	1.91	0.52
1:AA:999:U:O2'	1:AA:1000:A:H5''	2.10	0.52
1:AA:84:A:N6	1:AA:102:G:C2'	2.71	0.52
1:AA:1328:G:H2'	1:AA:1330:C:C5	2.45	0.52
1:AA:1341:U:O2	19:AT:80:ILE:HD13	2.10	0.52
1:AA:1536:A:OP1	1:AA:1537:C:N4	2.43	0.52
1:AA:2348:U:O4	1:AA:2382:G:C2	2.63	0.52
1:AA:239:U:H2'	1:AA:240:G:O4'	2.10	0.52
1:AA:1462:C:H4'	1:AA:2703:C:H5'	1.91	0.52
1:AA:2839:G:C5'	13:A0:46:GLY:HA2	2.40	0.52
2:AB:113:C:H2'	2:AB:114:G:C8	2.45	0.52
4:AE:171:GLU:HG2	4:AE:185:LYS:CG	2.41	0.52
6:AG:64:THR:OG1	6:AG:94:LEU:HD13	2.10	0.52
6:AG:74:LYS:O	6:AG:84:LYS:HD3	2.09	0.52
7:AH:89:ILE:CD1	7:AH:129:THR:HB	2.37	0.52
7:AH:86:GLU:O	7:AH:87:LEU:CB	2.57	0.52
8:AK:82:ARG:NH1	8:AK:146:ALA:HB2	2.25	0.52
1:AA:138:G:H22	19:AT:44:GLU:CD	2.13	0.52
25:AX:48:GLU:HA	25:AX:51:ALA:HB2	1.92	0.52
31:BA:1067:A:O2'	31:BA:1068:G:P	2.68	0.52
31:BA:1151:A:H2'	31:BA:1152:A:H8	1.75	0.52
31:BA:197:A:C6	31:BA:221:C:C4'	2.93	0.52
31:BA:261:U:OP2	50:BW:79:ARG:NH2	2.43	0.52
31:BA:433:C:O2'	31:BA:434:U:H5'	2.09	0.52
31:BA:452:A:O2'	46:BS:72:ARG:HG3	2.09	0.52
31:BA:518:C:H5''	31:BA:519:C:O5'	2.10	0.52
31:BA:680:C:H2'	31:BA:681:C:H6	1.76	0.52
31:BA:737:A:O2'	36:BI:72:VAL:CG1	2.58	0.52
52:BB:55:U:H2'	52:BB:57:G:OP2	2.10	0.52
34:BG:173:TRP:NE1	34:BG:174:LEU:HG	2.25	0.52
38:BK:100:ILE:HD12	38:BK:125:ARG:HG3	1.91	0.52
40:BM:12:ASP:HB3	40:BM:15:THR:CG2	2.39	0.52
40:BM:30:SER:O	40:BM:81:THR:HG22	2.10	0.52
43:BP:34:LEU:O	43:BP:38:GLY:N	2.43	0.52
48:BU:44:LEU:N	48:BU:44:LEU:HD12	2.25	0.52
50:BW:48:LYS:HD2	50:BW:51:GLU:OE2	2.09	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
50:BW:51:GLU:O	50:BW:55:ILE:HG12	2.10	0.52
34:CG:47:ARG:HH21	53:C1:57:U:H5	1.52	0.52
54:CA:1123:A:H4'	40:CM:36:GLY:HA3	1.91	0.52
54:CA:1157:A:H62	54:CA:1178:G:N2	2.08	0.52
54:CA:313:A:H2'	54:CA:314:C:H6	1.73	0.52
54:CA:428:G:O2'	54:CA:429:U:OP2	2.26	0.52
54:CA:529:G:O6	42:CO:49:ASN:HA	2.10	0.52
54:CA:547:A:C4'	54:CA:548:G:O5'	2.34	0.52
54:CA:947:G:H2'	54:CA:948:C:C6	2.45	0.52
32:CE:194:PRO:HG2	32:CE:195:ASP:H	1.75	0.52
32:CE:239:VAL:HG12	32:CE:240:GLN:CD	2.30	0.52
34:CG:9:CYS:O	34:CG:13:ARG:HG2	2.10	0.52
37:CJ:26:PHE:CE2	37:CJ:30:ILE:HD11	2.45	0.52
37:CJ:91:VAL:HB	37:CJ:96:GLN:HG2	1.91	0.52
39:CL:95:LYS:HD3	39:CL:96:LEU:N	2.24	0.52
40:CM:70:ARG:HG3	40:CM:70:ARG:HH11	1.75	0.52
44:CQ:40:CYS:N	44:CQ:43:CYS:SG	2.83	0.52
46:CS:68:ASP:C	46:CS:70:ALA:N	2.63	0.52
50:CW:23:ARG:CA	50:CW:26:ASN:HD21	2.19	0.52
13:D0:81:ASP:O	13:D0:82:GLU:HG2	2.10	0.52
55:DA:489:G:N2	55:DA:1321:A:OP1	2.43	0.52
55:DA:2653:U:H2'	55:DA:2654:A:C8	2.44	0.52
55:DA:2787:C:C1'	4:DE:62:PRO:HD3	2.34	0.52
55:DA:917:A:O4'	55:DA:917:A:N3	2.43	0.52
2:DB:25:A:H2'	2:DB:26:A:O4'	2.10	0.52
4:DE:34:VAL:O	4:DE:35:GLN:HB2	2.09	0.52
6:DG:166:ASP:HA	6:DG:169:ALA:HB3	1.92	0.52
7:DH:10:PRO:O	7:DH:11:VAL:CG1	2.56	0.52
56:DJ:19:GLU:O	56:DJ:23:LEU:HB2	2.09	0.52
56:DJ:18:LEU:O	56:DJ:20:LEU:N	2.43	0.52
8:DK:79:ILE:HB	8:DK:141:LYS:O	2.09	0.52
12:DP:109:VAL:HG13	12:DP:110:THR:N	2.23	0.52
14:DQ:60:GLY:O	14:DQ:61:ASN:CB	2.58	0.52
14:DQ:7:TYR:CE2	14:DQ:91:PRO:HG3	2.44	0.52
18:DS:82:LEU:H	18:DS:82:LEU:HD12	1.74	0.52
21:DV:35:ARG:NH1	21:DV:35:ARG:HB3	2.25	0.52
57:DY:144:ALA:HB1	57:DY:145:PRO:CD	2.10	0.52
12:AP:83:MET:HB3	22:A3:8:GLY:HA2	1.91	0.51
27:A5:40:LYS:HZ2	27:A5:45:VAL:CA	2.22	0.51
30:A8:63:PRO:O	30:A8:64:TYR:O	2.27	0.51
1:AA:1033:U:O2	1:AA:2750:A:H2	1.93	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1585:C:O2	1:AA:1585:C:H2'	2.08	0.51
1:AA:1729:A:C6	1:AA:1731:G:N7	2.78	0.51
1:AA:1837:C:OP1	31:BA:784:C:H4'	2.10	0.51
1:AA:1856:G:O2'	1:AA:1857:G:H5'	2.10	0.51
1:AA:1924:C:H2'	1:AA:1925:C:C1'	2.40	0.51
1:AA:2119:A:N1	1:AA:2171:A:H1'	2.24	0.51
1:AA:2162:G:H2'	1:AA:2163:C:C6	2.44	0.51
1:AA:2552:U:C2	1:AA:2554:U:H5'	2.46	0.51
1:AA:404:C:H1'	1:AA:406:G:N7	2.24	0.51
1:AA:440:G:H2'	1:AA:441:U:C6	2.45	0.51
1:AA:524:U:H2'	1:AA:525:U:C6	2.46	0.51
1:AA:547:A:H2'	1:AA:548:A:C8	2.45	0.51
1:AA:99:U:H1'	1:AA:102:G:N3	2.25	0.51
3:AD:111:LEU:HD13	3:AD:112:GLN:N	2.25	0.51
4:AE:110:GLY:HA3	4:AE:162:ALA:HB2	1.91	0.51
6:AG:101:ILE:O	6:AG:105:LYS:HG3	2.10	0.51
6:AG:16:ARG:NE	6:AG:31:VAL:HG11	2.25	0.51
8:AK:53:ALA:O	8:AK:56:LYS:HG3	2.11	0.51
8:AK:72:LEU:O	8:AK:74:ASN:N	2.37	0.51
9:AM:11:PRO:HB2	9:AM:51:PHE:HE1	1.74	0.51
9:AM:137:LYS:NZ	9:AM:138:LEU:HD23	2.24	0.51
10:AN:47:ILE:O	10:AN:48:PRO:C	2.48	0.51
11:AO:59:LEU:CD2	11:AO:59:LEU:C	2.79	0.51
1:AA:910:A:H62	12:AP:12:GLN:HA	1.75	0.51
12:AP:2:LEU:O	12:AP:3:MET:HG2	2.10	0.51
12:AP:68:ILE:HD13	12:AP:103:MET:CG	2.40	0.51
12:AP:78:PRO:O	12:AP:79:LEU:CG	2.58	0.51
21:AV:10:ARG:O	21:AV:36:LYS:HB2	2.10	0.51
31:BA:1067:A:N3	31:BA:1068:G:H1'	2.25	0.51
31:BA:1104:G:O2'	31:BA:1105:A:H5'	2.10	0.51
31:BA:1326:C:H2'	31:BA:1327:C:H6	1.74	0.51
1:AA:1915:U:O4	31:BA:1409:C:H4'	2.10	0.51
31:BA:209:U:O2'	31:BA:210:U:OP2	2.25	0.51
31:BA:123:C:OP1	31:BA:312:C:H5'	2.10	0.51
31:BA:373:A:O2'	31:BA:374:A:H5'	2.10	0.51
31:BA:438:G:H4'	34:BG:123:HIS:CE1	2.45	0.51
31:BA:438:G:OP1	34:BG:151:LYS:HE2	2.10	0.51
31:BA:451:A:H1'	31:BA:452:A:N7	2.24	0.51
31:BA:817:C:H4'	31:BA:818:G:OP1	2.09	0.51
31:BA:869:G:H4'	31:BA:872:A:H1'	1.92	0.51
32:BE:12:GLU:O	32:BE:15:VAL:N	2.42	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:BE:195:ASP:C	32:BE:197:VAL:H	2.13	0.51
32:BE:187:LEU:HD11	32:BE:204:ASN:O	2.09	0.51
32:BE:30:ARG:HB2	32:BE:46:LYS:NZ	2.25	0.51
32:BE:98:LEU:HB2	32:BE:101:MET:HE1	1.92	0.51
33:BF:62:ASP:C	33:BF:97:LYS:HB2	2.30	0.51
34:BG:150:GLU:N	34:BG:150:GLU:OE2	2.42	0.51
36:BI:6:VAL:CG1	36:BI:8:ILE:HD11	2.34	0.51
37:BJ:22:LEU:HG	37:BJ:62:PHE:CE2	2.41	0.51
31:BA:4:U:H3	38:BK:102:ARG:NH1	2.07	0.51
40:BM:65:LEU:HD23	40:BM:65:LEU:O	2.11	0.51
41:BN:98:LEU:HA	41:BN:101:SER:HB3	1.91	0.51
47:BT:65:ILE:O	47:BT:66:SER:HB3	2.11	0.51
49:BV:11:VAL:HG21	49:BV:16:LEU:HD11	1.90	0.51
49:BV:29:ARG:O	49:BV:29:ARG:HG2	2.10	0.51
53:C1:52:U:H2'	53:C1:53:U:H5'	1.92	0.51
54:CA:1005:A:H5''	54:CA:1006:C:C6	2.44	0.51
54:CA:1128:C:O2'	54:CA:1130:A:C8	2.63	0.51
54:CA:1262:C:H2'	54:CA:1263:C:H6	1.74	0.51
54:CA:1342:C:H2'	54:CA:1343:G:H8	1.74	0.51
54:CA:192:U:O2'	54:CA:193:C:H5'	2.10	0.51
54:CA:632:A:H8	54:CA:633:G:C8	2.28	0.51
54:CA:685:G:N2	54:CA:686:U:N3	2.58	0.51
54:CA:652:U:O4	54:CA:752:G:H2'	2.09	0.51
52:CB:74:C:O2'	52:CB:75:C:C6	2.64	0.51
52:CC:54:U:H2'	52:CC:55:U:O4'	2.09	0.51
33:CF:21:ARG:CD	33:CF:21:ARG:N	2.72	0.51
33:CF:45:LYS:HB2	33:CF:45:LYS:NZ	2.25	0.51
33:CF:82:GLU:O	33:CF:86:VAL:HG13	2.09	0.51
36:CI:30:LEU:HB3	36:CI:35:ALA:HB3	1.92	0.51
37:CJ:151:TYR:HA	37:CJ:153:HIS:CE1	2.45	0.51
38:CK:12:ARG:NH1	38:CK:27:PRO:CD	2.71	0.51
39:CL:79:LEU:CD1	39:CL:83:ARG:HD2	2.39	0.51
43:CP:115:LYS:O	43:CP:117:VAL:CG1	2.58	0.51
46:CS:8:ARG:C	46:CS:9:PHE:HD2	2.12	0.51
27:D5:40:LYS:CB	27:D5:46:CYS:SG	2.97	0.51
55:DA:1079:C:H2'	55:DA:1080:A:C8	2.45	0.51
55:DA:1331:A:H2'	55:DA:1333:C:C5	2.45	0.51
55:DA:1418:G:H8	55:DA:1418:G:O5'	1.92	0.51
55:DA:2019:A:H2'	55:DA:2020:A:O5'	2.11	0.51
55:DA:2423:U:O2'	55:DA:2425:A:H2'	2.09	0.51
55:DA:287:C:O2'	55:DA:288:C:H5'	2.09	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DB:40:U:C2	2:DB:43:C:OP2	2.63	0.51
5:DF:118:ALA:HA	5:DF:123:LEU:HB3	1.92	0.51
7:DH:153:LYS:CB	7:DH:154:PRO:CD	2.72	0.51
7:DH:42:ARG:HG3	7:DH:42:ARG:O	2.09	0.51
56:DJ:24:ILE:HG22	56:DJ:25:ASP:H	1.75	0.51
8:DK:110:ASP:HB2	8:DK:111:PRO:CA	2.40	0.51
8:DK:127:VAL:HG13	8:DK:139:GLN:HB3	1.92	0.51
8:DK:82:ARG:NH1	8:DK:82:ARG:HG3	2.24	0.51
58:DL:47:ASN:O	58:DL:48:MET:CB	2.58	0.51
58:DL:95:LYS:O	58:DL:96:VAL:C	2.49	0.51
14:DQ:26:LEU:HG	14:DQ:39:ILE:HD13	1.92	0.51
19:DT:12:VAL:O	19:DT:12:VAL:HG13	2.10	0.51
55:DA:1599:C:OP1	19:DT:36:LYS:HG3	2.10	0.51
20:DU:48:ALA:H	20:DU:60:PHE:HA	1.75	0.51
57:DY:9:LEU:HD13	57:DY:10:LEU:CB	2.40	0.51
57:DY:27:VAL:CG1	57:DY:110:GLY:CA	2.81	0.51
57:DY:130:THR:HG22	56:DJ:14:GLN:NE2	2.12	0.51
57:DY:26:LEU:HB3	57:DY:112:LEU:HD22	1.91	0.51
57:DY:71:LEU:HD22	57:DY:72:ASP:CA	2.40	0.51
30:A8:50:LEU:CD1	30:A8:54:GLU:CA	2.88	0.51
1:AA:1128:A:O2'	1:AA:1129:A:O5'	2.27	0.51
1:AA:1271:G:C2	1:AA:1617:C:H4'	2.44	0.51
1:AA:1773:A:C2'	1:AA:1774:C:H5'	2.39	0.51
1:AA:1857:G:O2'	1:AA:1885:A:N6	2.41	0.51
1:AA:197:A:H62	1:AA:2430:A:H2'	1.75	0.51
1:AA:19:C:H2'	1:AA:20:C:C6	2.46	0.51
1:AA:2211:G:O2'	1:AA:2212:A:P	2.68	0.51
1:AA:921:G:H4'	1:AA:2269:A:C5	2.45	0.51
1:AA:2282:G:H5''	1:AA:2283:C:O4'	2.09	0.51
1:AA:2785:C:H2'	1:AA:2786:U:C6	2.44	0.51
1:AA:662:G:H5'	11:AO:15:ARG:CA	2.37	0.51
1:AA:729:G:C6	3:AD:208:LYS:HB2	2.44	0.51
1:AA:858:U:O2'	1:AA:2268:A:C2'	2.58	0.51
2:AB:38:C:O2	2:AB:48:A:H1'	2.10	0.51
4:AE:73:GLU:HG2	4:AE:74:PRO:HD2	1.92	0.51
5:AF:18:ARG:CG	5:AF:19:GLU:H	2.22	0.51
5:AF:59:TYR:CD2	5:AF:59:TYR:N	2.74	0.51
6:AG:180:PHE:C	6:AG:182:LYS:N	2.63	0.51
6:AG:41:GLN:NE2	6:AG:60:LEU:HD12	2.25	0.51
8:AK:82:ARG:CG	8:AK:82:ARG:HH11	2.23	0.51
11:AO:47:ASP:HB3	11:AO:48:PRO:C	2.31	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AO:56:SER:O	11:AO:57:THR:CB	2.58	0.51
11:AO:59:LEU:C	11:AO:59:LEU:HD23	2.30	0.51
12:AP:31:ASP:HA	12:AP:134:ARG:HD2	1.92	0.51
12:AP:17:LEU:HD21	12:AP:41:TRP:NE1	2.25	0.51
19:AT:21:PHE:O	19:AT:23:GLU:N	2.43	0.51
19:AT:30:VAL:HG12	19:AT:31:HIS:H	1.73	0.51
20:AU:27:VAL:HA	20:AU:39:VAL:HG12	1.91	0.51
21:AV:30:ASN:HB3	21:AV:90:VAL:HB	1.91	0.51
53:B1:34:G:H2'	53:B1:35:A:H8	1.75	0.51
31:BA:101:A:O2'	31:BA:102:G:H5'	2.10	0.51
31:BA:1104:G:H4'	32:BE:111:ARG:HH21	1.74	0.51
31:BA:1236:A:H2'	31:BA:1237:C:C6	2.44	0.51
31:BA:1301:U:C2'	31:BA:1301:U:O2	2.55	0.51
31:BA:1491:G:O3'	42:BO:46:LYS:HB2	2.10	0.51
31:BA:388:G:HO2'	31:BA:389:A:P	2.33	0.51
31:BA:555:C:H2'	31:BA:556:C:C6	2.44	0.51
31:BA:652:U:O2'	31:BA:653:A:H5''	2.10	0.51
31:BA:86:U:C2'	31:BA:87:A:OP1	2.58	0.51
52:BB:33:U:O4'	52:BB:37:MIA:H161	2.10	0.51
52:BC:19:G:C4	52:BC:57:G:N2	2.78	0.51
33:BF:126:ARG:O	33:BF:128:PHE:N	2.44	0.51
33:BF:25:GLY:C	33:BF:27:LYS:N	2.59	0.51
33:BF:53:ALA:HB1	33:BF:114:PRO:HB2	1.91	0.51
34:BG:110:PHE:CD1	34:BG:110:PHE:N	2.70	0.51
38:BK:29:SER:HB3	38:BK:32:LYS:CD	2.40	0.51
39:BL:3:GLN:NE2	39:BL:20:ARG:NH2	2.57	0.51
40:BM:33:GLN:O	40:BM:75:ILE:HG23	2.09	0.51
47:BT:11:VAL:O	47:BT:12:SER:HB2	2.10	0.51
54:CA:1276:G:H2'	54:CA:1277:C:C6	2.45	0.51
54:CA:1305:G:N2	54:CA:1331:G:C2'	2.69	0.51
54:CA:546:G:P	34:CG:72:GLU:HB3	2.49	0.51
54:CA:628:G:N2	54:CA:629:G:C4	2.79	0.51
54:CA:687:A:C6	54:CA:704:A:N7	2.79	0.51
32:CE:139:LYS:N	32:CE:139:LYS:HD2	2.25	0.51
32:CE:160:ASP:O	32:CE:183:PRO:HD2	2.10	0.51
32:CE:216:SER:O	32:CE:218:ALA:N	2.44	0.51
33:CF:78:GLY:HA3	33:CF:83:ARG:HB2	1.90	0.51
34:CG:116:GLN:HE22	34:CG:157:LEU:HD11	1.74	0.51
35:CH:12:LEU:O	35:CH:13:ILE:HD12	2.09	0.51
39:CL:17:VAL:CG1	39:CL:81:ILE:HD13	2.40	0.51
41:CN:27:ASN:OD1	41:CN:55:LYS:HB3	2.10	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:CT:11:VAL:HG23	47:CT:20:THR:HB	1.91	0.51
22:D3:63:VAL:HG23	22:D3:64:ASP:O	2.10	0.51
26:D4:47:GLN:O	26:D4:48:ARG:HB2	2.11	0.51
30:D8:35:GLN:HE21	30:D8:35:GLN:CA	2.22	0.51
55:DA:1086:A:H5'	55:DA:1103:A:N6	2.18	0.51
55:DA:1528:A:O2'	55:DA:1529:A:H5'	2.11	0.51
55:DA:1899:G:H22	55:DA:1902:C:H41	0.62	0.51
55:DA:2060:A:H4'	55:DA:2061:G:OP2	2.10	0.51
55:DA:2062:A:C2'	55:DA:2062:A:N3	2.73	0.51
55:DA:2250:G:C8	55:DA:2496:C:H5''	2.46	0.51
55:DA:2282:G:OP1	55:DA:2283:C:H1'	2.10	0.51
55:DA:2310:A:N6	6:DG:79:ASN:HD22	2.07	0.51
55:DA:2439:A:O2'	55:DA:2440:C:OP2	2.28	0.51
55:DA:2747:G:O3'	7:DH:70:THR:HG21	2.10	0.51
55:DA:725:G:C6	55:DA:726:G:N1	2.78	0.51
2:DB:48:A:H2'	2:DB:49:C:H6	1.70	0.51
55:DA:2224:G:OP1	3:DD:268:ARG:HD3	2.10	0.51
3:DD:35:LYS:HA	3:DD:64:ILE:HG22	1.90	0.51
3:DD:65:ILE:N	3:DD:65:ILE:HD13	2.25	0.51
5:DF:155:LEU:HD12	5:DF:174:VAL:O	2.10	0.51
6:DG:41:GLN:HB3	6:DG:43:LEU:HD13	1.93	0.51
56:DI:23:LEU:C	56:DI:27:LEU:HD12	2.28	0.51
57:DY:90:ALA:HB3	56:DJ:15:ALA:N	2.25	0.51
10:DN:7:TYR:CE1	10:DN:20:MET:HE3	2.38	0.51
11:DO:30:THR:O	11:DO:31:ALA:HB3	2.10	0.51
12:DP:20:ALA:HB2	12:DP:99:PRO:HD2	1.93	0.51
21:DV:105:VAL:CG1	21:DV:140:ASP:CB	2.83	0.51
25:DX:59:VAL:CG1	25:DX:60:GLU:N	2.73	0.51
57:DY:142:LEU:CD1	57:DY:143:GLN:HG2	2.40	0.51
57:DY:143:GLN:O	57:DY:144:ALA:O	2.28	0.51
57:DY:19:ARG:HH22	57:DY:84:GLU:CD	2.14	0.51
57:DY:93:LEU:HD13	57:DY:97:ALA:O	2.08	0.51
57:DY:93:LEU:HG	57:DY:126:ALA:CA	2.40	0.51
17:A2:1:MET:SD	17:A2:42:GLY:HA3	2.50	0.51
28:A6:47:THR:HG23	28:A6:48:VAL:H	1.76	0.51
11:AO:64:LYS:CE	30:A8:30:ARG:CZ	2.81	0.51
1:AA:127:A:H5''	1:AA:128:C:C6	2.45	0.51
1:AA:2051:A:H8	1:AA:2051:A:OP2	1.94	0.51
1:AA:2287:A:H2	1:AA:2346:A:C2	2.28	0.51
1:AA:2657:A:N9	1:AA:2665:A:N6	2.59	0.51
1:AA:302:C:H2'	1:AA:303:U:H6	1.74	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:384:U:O2'	1:AA:385:C:C5'	2.58	0.51
1:AA:608:A:N9	1:AA:621:A:N6	2.58	0.51
4:AE:107:THR:HG22	4:AE:107:THR:O	2.11	0.51
5:AF:110:LEU:HD22	5:AF:202:PHE:HE1	1.75	0.51
5:AF:37:VAL:HA	5:AF:40:GLN:CG	2.40	0.51
6:AG:121:ASN:HD22	6:AG:122:PRO:N	2.08	0.51
6:AG:133:LEU:HD23	6:AG:133:LEU:N	2.25	0.51
6:AG:53:LEU:HD23	6:AG:54:GLU:N	2.24	0.51
7:AH:26:VAL:HG11	7:AH:33:LEU:HB2	1.92	0.51
7:AH:88:LEU:HD13	7:AH:164:TYR:O	2.10	0.51
7:AH:89:ILE:HG12	7:AH:90:LYS:N	2.26	0.51
9:AM:26:LEU:HD23	9:AM:99:LEU:HD21	1.92	0.51
10:AN:8:LEU:HD13	10:AN:82:ASN:O	2.10	0.51
12:AP:9:TYR:O	12:AP:10:ARG:HD3	2.11	0.51
14:AQ:106:ARG:HG2	14:AQ:110:LEU:HD11	1.92	0.51
15:AR:132:LYS:HG2	15:AR:136:GLN:NE2	2.16	0.51
19:AT:18:TYR:CA	19:AT:21:PHE:HD2	2.14	0.51
19:AT:17:ALA:HA	19:AT:27:THR:HG21	1.93	0.51
21:AV:8:TYR:HD1	21:AV:38:TYR:HH	1.59	0.51
31:BA:1023:G:C2'	31:BA:1024:G:OP1	2.59	0.51
31:BA:1095:U:H5''	31:BA:1109:C:O2	2.11	0.51
31:BA:1318:A:O2'	49:BV:37:ARG:HB3	2.10	0.51
31:BA:1350:A:N6	31:BA:1373:G:N2	2.58	0.51
31:BA:153:C:H2'	31:BA:154:C:H6	1.75	0.51
31:BA:177:C:H2'	31:BA:178:C:H6	1.75	0.51
31:BA:366:C:HO2'	31:BA:367:U:P	2.29	0.51
31:BA:39:G:O2'	31:BA:40:C:H5'	2.11	0.51
31:BA:543:C:OP1	34:BG:14:ARG:NE	2.43	0.51
31:BA:766:A:H2'	31:BA:767:A:O4'	2.10	0.51
31:BA:827:U:O5'	31:BA:827:U:H6	1.93	0.51
32:BE:213:LEU:HD23	32:BE:213:LEU:C	2.31	0.51
32:BE:5:ILE:HA	32:BE:224:GLN:HE21	1.74	0.51
32:BE:73:THR:HG22	32:BE:94:ASN:C	2.31	0.51
33:BF:11:ARG:HH11	33:BF:11:ARG:HG2	1.75	0.51
33:BF:32:LEU:O	33:BF:35:GLU:HB3	2.10	0.51
34:BG:153:ARG:HH12	34:BG:181:MET:HB2	1.75	0.51
35:BH:80:ILE:HD12	35:BH:82:VAL:HG23	1.92	0.51
31:BA:737:A:H5'	36:BI:90:VAL:O	2.09	0.51
39:BL:10:ARG:NH2	39:BL:11:LYS:HB2	2.21	0.51
49:BV:80:TYR:O	49:BV:81:ARG:CB	2.59	0.51
31:BA:1453:G:C3'	50:BW:39:LYS:HZ2	2.22	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1296:C:C5'	54:CA:1297:C:OP2	2.54	0.51
54:CA:1391:U:H2'	54:CA:1392:G:H8	1.75	0.51
54:CA:1402:C:H2'	54:CA:1403:C:O4'	2.10	0.51
54:CA:1535:C:H2'	54:CA:1536:C:H5'	1.91	0.51
54:CA:181:G:N2	54:CA:183:G:N2	2.58	0.51
54:CA:614:A:H2'	54:CA:615:C:C6	2.45	0.51
32:CE:55:PHE:CE1	32:CE:218:ALA:HA	2.44	0.51
33:CF:86:VAL:O	33:CF:90:GLU:HG2	2.11	0.51
36:CI:61:LEU:N	36:CI:61:LEU:HD12	2.24	0.51
37:CJ:76:ARG:HD3	37:CJ:156:TRP:CZ2	2.45	0.51
37:CJ:23:VAL:HG12	37:CJ:27:ILE:HD13	1.92	0.51
40:CM:19:SER:O	40:CM:23:ILE:HG13	2.11	0.51
40:CM:29:ARG:HH11	40:CM:29:ARG:HG2	1.75	0.51
48:CU:66:LEU:HD11	48:CU:70:ILE:HD11	1.92	0.51
16:D1:60:LEU:CD2	16:D1:64:ARG:HG3	2.37	0.51
29:D7:24:THR:O	29:D7:28:ARG:HG3	2.10	0.51
55:DA:1045:A:O2'	55:DA:1047:G:C5	2.61	0.51
55:DA:1252:G:C2	55:DA:1253:A:C2	2.98	0.51
55:DA:12:U:O5'	55:DA:12:U:H6	1.93	0.51
55:DA:1535:U:N3	55:DA:1536:A:H3'	2.26	0.51
55:DA:1851:U:H2'	55:DA:1852:C:O4'	2.10	0.51
55:DA:215:G:H4'	55:DA:216:A:H4'	1.91	0.51
55:DA:2637:U:H5"	4:DE:82:ARG:HH21	1.76	0.51
55:DA:2712:U:H3'	55:DA:2712:U:O2	2.10	0.51
55:DA:2737:G:H2'	55:DA:2738:A:H8	1.75	0.51
55:DA:2756:U:H4'	55:DA:2757:A:O5'	2.10	0.51
55:DA:355:G:O2'	55:DA:356:G:H5'	2.10	0.51
55:DA:817:C:O2'	55:DA:839:U:H5"	2.10	0.51
55:DA:921:G:H4'	55:DA:2269:A:C5	2.45	0.51
55:DA:847:U:H5	55:DA:933:A:C2	2.28	0.51
2:DB:73:A:H2'	2:DB:73:A:N3	2.25	0.51
7:DH:123:PHE:HE2	7:DH:133:VAL:HG22	1.75	0.51
55:DA:622:G:OP2	11:DO:108:LYS:HE3	2.10	0.51
11:DO:126:VAL:HG13	11:DO:145:PRO:HB2	1.92	0.51
15:DR:26:ASP:CB	15:DR:91:ARG:HA	2.30	0.51
21:DV:5:LEU:O	21:DV:5:LEU:HD13	2.10	0.51
57:DY:43:ALA:HB3	57:DY:47:ASN:CA	2.39	0.51
13:A0:117:VAL:O	13:A0:118:GLU:HB2	2.10	0.51
17:A2:4:ILE:HD13	17:A2:40:LEU:HB2	1.93	0.51
22:A3:65:GLY:HA3	22:A3:81:VAL:HG12	1.92	0.51
27:A5:52:TYR:O	27:A5:53:ALA:HB2	2.10	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1210:A:H5'	1:AA:1212:G:O4'	2.11	0.51
1:AA:1525:G:H2'	1:AA:1526:G:C8	2.45	0.51
1:AA:1528:A:C2	1:AA:1543:A:N1	2.78	0.51
1:AA:1657:C:H2'	1:AA:1658:C:C6	2.45	0.51
1:AA:1799:G:H2'	3:AD:181:GLU:OE2	2.11	0.51
1:AA:1944:U:O2	1:AA:1955:U:H5''	2.10	0.51
1:AA:2064:C:H2'	1:AA:2065:C:C6	2.45	0.51
1:AA:2197:U:O2'	1:AA:2198:A:C8	2.58	0.51
1:AA:2655:G:H2'	1:AA:2656:U:OP2	2.10	0.51
1:AA:271(C):U:O2	1:AA:271(C):U:H2'	2.10	0.51
1:AA:654(J):A:C2	1:AA:654(L):G:N7	2.79	0.51
1:AA:814:C:O2'	1:AA:815:C:H5'	2.10	0.51
2:AB:58:A:H3'	2:AB:59:A:H8	1.74	0.51
3:AD:9:TYR:CD2	3:AD:10:THR:HG22	2.45	0.51
4:AE:130:GLY:O	4:AE:131:ALA:HB2	2.10	0.51
6:AG:112:PRO:CB	26:A4:37:SER:HA	2.40	0.51
1:AA:2667:C:H1'	7:AH:109:PHE:CD2	2.45	0.51
7:AH:149:ARG:HA	7:AH:162:ILE:HG21	1.93	0.51
8:AK:88:ILE:HG22	8:AK:90:GLY:H	1.76	0.51
9:AM:17:ASP:O	9:AM:18:ALA:CB	2.57	0.51
10:AN:12:ASP:C	10:AN:12:ASP:OD2	2.48	0.51
11:AO:115:LEU:CD2	11:AO:131:SER:HB2	2.40	0.51
15:AR:29:ARG:HG2	15:AR:46:GLU:HB2	1.92	0.51
15:AR:58:ASN:ND2	15:AR:58:ASN:N	2.59	0.51
21:AV:121:HIS:HB2	21:AV:171:ILE:HD13	1.92	0.51
21:AV:108:PRO:HG2	21:AV:141:VAL:O	2.03	0.51
21:AV:6:LYS:O	21:AV:59:LEU:O	2.29	0.51
25:AX:47:VAL:HG11	25:AX:56:VAL:HG21	1.93	0.51
31:BA:1267:C:O2	31:BA:1267:C:H2'	2.10	0.51
31:BA:1402:C:O2	31:BA:1500:A:N1	2.42	0.51
31:BA:406:G:H2'	31:BA:407:G:H8	1.75	0.51
31:BA:409:G:H2'	31:BA:410:G:H8	1.75	0.51
31:BA:85:U:O2	31:BA:85:U:C2'	2.58	0.51
31:BA:960:U:O2	31:BA:960:U:H2'	2.11	0.51
52:BB:11:C:O2'	52:BB:12:U:H5'	2.10	0.51
52:BB:23:A:H2'	52:BB:24:G:C1'	2.40	0.51
1:AA:1924:C:O2'	52:BC:13:C:H4'	2.10	0.51
32:BE:185:ILE:HG22	32:BE:199:TYR:HB2	1.91	0.51
33:BF:12:LEU:HA	33:BF:16:ARG:O	2.09	0.51
33:BF:16:ARG:HH21	33:BF:183:ASP:HA	1.74	0.51
33:BF:70:VAL:HG12	33:BF:71:ALA:N	2.26	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:BG:19:LEU:O	34:BG:20:TYR:C	2.48	0.51
36:BI:2:ARG:O	36:BI:66:GLU:HG3	2.10	0.51
40:BM:18:ALA:O	40:BM:21:GLN:HB3	2.11	0.51
54:CA:1424:C:H2'	54:CA:1425:U:H6	1.74	0.51
54:CA:167:G:O2'	54:CA:168:G:H5'	2.11	0.51
54:CA:434:U:H2'	54:CA:435:C:C6	2.46	0.51
54:CA:895:G:H2'	54:CA:896:C:H6	1.75	0.51
54:CA:951:G:O2'	54:CA:972:C:C5	2.62	0.51
52:CB:9:A:C2	52:CB:11:C:N4	2.79	0.51
33:CF:126:ARG:HH11	33:CF:126:ARG:HG2	1.76	0.51
37:CJ:30:ILE:HD12	37:CJ:120:ILE:HD12	1.92	0.51
37:CJ:69:VAL:O	37:CJ:69:VAL:HG12	2.11	0.51
38:CK:63:LEU:HB2	38:CK:65:TYR:HE1	1.75	0.51
39:CL:20:ARG:O	39:CL:60:ASP:HB2	2.09	0.51
43:CP:14:ARG:CA	43:CP:44:ARG:HA	2.37	0.51
46:CS:13:HIS:C	46:CS:15:PRO:HD3	2.30	0.51
46:CS:50:LYS:HD3	46:CS:50:LYS:C	2.30	0.51
46:CS:4:ILE:HD13	46:CS:66:PRO:N	2.24	0.51
30:D8:62:LEU:O	30:D8:63:PRO:C	2.48	0.51
30:D8:62:LEU:HB2	30:D8:63:PRO:HD3	1.91	0.51
55:DA:127:A:H5''	55:DA:128:C:C6	2.46	0.51
55:DA:1360:A:C2'	55:DA:1361:G:H5'	2.40	0.51
55:DA:1558:A:HO2'	55:DA:1559:G:P	2.33	0.51
55:DA:1820:U:O2'	3:DD:201:HIS:HD2	1.93	0.51
55:DA:189:G:O6	55:DA:205:G:H2'	2.10	0.51
55:DA:2520:C:C6	55:DA:2567:G:H1'	2.44	0.51
55:DA:2657:A:N9	55:DA:2665:A:N6	2.58	0.51
55:DA:343:C:H5'	55:DA:344:G:OP2	2.10	0.51
55:DA:403:U:H4'	55:DA:404:C:O5'	2.10	0.51
55:DA:693:C:O2'	55:DA:694:U:H5'	2.11	0.51
55:DA:885:C:OP1	55:DA:885:C:C4'	2.58	0.51
3:DD:182:LEU:H	3:DD:272:ALA:CB	2.23	0.51
6:DG:13:GLU:O	6:DG:14:GLU:CB	2.58	0.51
6:DG:146:TYR:C	6:DG:148:MET:H	2.13	0.51
7:DH:25:LYS:HG2	7:DH:34:GLU:HG2	1.92	0.51
8:DK:127:VAL:HG22	8:DK:139:GLN:HB3	1.93	0.51
8:DK:1:MET:HG3	8:DK:23:PRO:CB	2.41	0.51
58:DL:51:ALA:HB1	58:DL:79:ARG:NE	2.26	0.51
58:DL:59:ILE:O	58:DL:60:TYR:CD1	2.63	0.51
58:DL:72:PRO:CD	58:DL:73:PRO:HD3	2.40	0.51
58:DL:77:LEU:HB3	58:DL:107:ILE:CG1	2.39	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:DQ:108:GLY:O	14:DQ:110:LEU:N	2.43	0.51
55:DA:2683:C:P	15:DR:53:ARG:NH2	2.82	0.51
19:DT:31:HIS:CD2	19:DT:33:LYS:H	2.28	0.51
21:DV:69:THR:HB	21:DV:88:PHE:HB3	1.92	0.51
17:A2:35:LEU:H	17:A2:35:LEU:HD22	1.74	0.51
26:A4:55:ARG:O	26:A4:55:ARG:HG3	2.10	0.51
28:A6:48:VAL:O	28:A6:49:HIS:CB	2.59	0.51
1:AA:1198:U:H2'	1:AA:1199:U:C6	2.45	0.51
1:AA:1229:G:N2	1:AA:1229(A):G:H1'	2.25	0.51
1:AA:1288:U:H5''	1:AA:1289:C:OP2	2.09	0.51
1:AA:2404:C:O2'	1:AA:2405:G:H5'	2.10	0.51
1:AA:2420:C:OP1	30:A8:33:ASN:O	2.28	0.51
1:AA:2461:C:C2	1:AA:2462:U:C5	2.98	0.51
1:AA:2808:U:O2'	1:AA:2809:A:H5'	2.10	0.51
1:AA:300:A:P	20:AU:84:ARG:HH12	2.34	0.51
1:AA:887:A:O2'	43:BP:93:ARG:HG2	2.10	0.51
3:AD:106:ILE:HD13	3:AD:106:ILE:C	2.31	0.51
5:AF:164:ARG:HG3	5:AF:175:THR:OG1	2.10	0.51
5:AF:3:GLU:CA	5:AF:24:LEU:HG	2.31	0.51
5:AF:36:VAL:HB	5:AF:183:VAL:HG21	1.92	0.51
6:AG:88:ILE:HD13	6:AG:89:GLY:N	2.25	0.51
11:AO:86:LYS:CG	11:AO:87:ASP:N	2.73	0.51
15:AR:51:ARG:HH11	15:AR:51:ARG:CG	2.24	0.51
19:AT:21:PHE:C	19:AT:23:GLU:N	2.63	0.51
20:AU:39:VAL:CG2	20:AU:40:GLU:H	2.19	0.51
23:AZ:66:HIS:O	23:AZ:67:ILE:C	2.49	0.51
23:AZ:78:LYS:HD2	23:AZ:80:LEU:CD2	2.26	0.51
31:BA:1151:A:C1'	40:BM:39:PRO:HB2	2.39	0.51
31:BA:1238:A:H2	31:BA:1241:G:N3	2.09	0.51
31:BA:949:A:H1'	31:BA:1364:U:N3	2.24	0.51
31:BA:1501:C:OP2	31:BA:1504:G:H2'	2.10	0.51
31:BA:713:G:N2	31:BA:777:A:H1'	2.25	0.51
31:BA:962:C:H2'	31:BA:963:G:O4'	2.10	0.51
1:AA:2394:C:N4	52:BD:76:A:H8	2.03	0.51
32:BE:47:THR:HA	32:BE:202:PRO:HG2	1.92	0.51
32:BE:76:GLN:O	32:BE:208:ILE:HG12	2.10	0.51
33:BF:23:TYR:OH	33:BF:25:GLY:HA2	2.11	0.51
38:BK:96:GLY:H	38:BK:99:GLU:CD	2.13	0.51
39:BL:117:HIS:O	39:BL:118:LYS:HG3	2.10	0.51
39:BL:3:GLN:CG	39:BL:20:ARG:HH12	2.23	0.51
39:BL:7:THR:HG21	39:BL:9:ARG:HH21	1.75	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:BM:40:LEU:HB3	40:BM:69:ASN:CB	2.27	0.51
40:BM:99:LYS:HD2	40:BM:100:THR:H	1.74	0.51
42:BO:59:ARG:HA	42:BO:65:GLU:HG2	1.93	0.51
46:BS:20:VAL:HG23	46:BS:32:TYR:CB	2.41	0.51
46:BS:75:ARG:HG3	46:BS:80:PHE:HD1	1.76	0.51
48:BU:37:VAL:O	48:BU:40:LEU:N	2.43	0.51
49:BV:46:GLY:H	49:BV:62:ILE:HG23	1.76	0.51
54:CA:1202:G:C2	44:CQ:42:ILE:HG21	2.46	0.51
54:CA:1269:A:C2	54:CA:1313:U:O4'	2.63	0.51
54:CA:448:A:N3	54:CA:449:C:O2	2.43	0.51
54:CA:683:G:H2'	54:CA:684:A:C8	2.46	0.51
54:CA:688:G:H2'	54:CA:689:C:H6	1.76	0.51
54:CA:718:G:C8	41:CN:116:HIS:HB3	2.46	0.51
54:CA:777:A:H2'	54:CA:778:G:C8	2.45	0.51
52:CB:58:A:H1'	52:CB:60:U:C5	2.45	0.51
52:CB:74:C:H1'	52:CB:75:C:H5'	1.91	0.51
32:CE:42:ILE:O	32:CE:44:LEU:HD12	2.10	0.51
32:CE:9:GLU:HB3	32:CE:48:MET:SD	2.51	0.51
33:CF:150:LYS:HE2	33:CF:152:ILE:HD11	1.93	0.51
54:CA:8:A:N7	34:CG:208:SER:O	2.42	0.51
39:CL:53:VAL:HB	39:CL:95:LYS:HE3	1.93	0.51
42:CO:83:VAL:HG23	42:CO:100:ILE:HG23	1.93	0.51
48:CU:36:ASN:HB2	48:CU:39:VAL:HG23	1.91	0.51
13:D0:75:LEU:O	13:D0:79:LEU:HB2	2.11	0.51
26:D4:48:ARG:NH2	26:D4:51:ASP:HA	2.25	0.51
28:D6:15:GLU:OE1	28:D6:44:ARG:NH2	2.41	0.51
55:DA:1055:G:C6	55:DA:1056:G:N7	2.79	0.51
55:DA:1360:A:H2'	55:DA:1361:G:C5'	2.40	0.51
55:DA:1629:U:H2'	55:DA:1630:G:C8	2.44	0.51
55:DA:2126:A:HO2'	55:DA:2127:G:C5'	2.24	0.51
55:DA:2175:C:H3'	55:DA:2176:A:H5''	1.92	0.51
55:DA:2319:G:H4'	55:DA:2320:A:OP1	2.10	0.51
55:DA:2615:U:H2'	55:DA:2616:C:H6	1.75	0.51
55:DA:2778:A:H4'	55:DA:2779:U:OP1	2.09	0.51
55:DA:803:U:H2'	55:DA:804:A:H5'	1.91	0.51
55:DA:941:A:H2'	55:DA:942:G:C8	2.46	0.51
55:DA:961:C:N4	55:DA:2031:A:H1'	2.25	0.51
2:DB:37:C:O2'	2:DB:38:C:H5'	2.10	0.51
3:DD:102:LYS:O	3:DD:103:ARG:HG2	2.10	0.51
3:DD:149:PRO:HD3	3:DD:186:HIS:HB3	1.93	0.51
3:DD:69:ARG:C	3:DD:71:ASP:N	2.63	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:DE:111:ARG:HD2	4:DE:160:TYR:CE1	2.45	0.51
6:DG:28:VAL:O	6:DG:31:VAL:HG12	2.11	0.51
8:DK:114:LEU:O	8:DK:115:ALA:HB2	2.11	0.51
8:DK:31:LEU:HB2	8:DK:32:PRO:HD3	1.93	0.51
58:DL:83:GLY:N	58:DL:99:ILE:CG2	2.56	0.51
10:DN:64:ARG:NH1	10:DN:81:ASP:OD1	2.43	0.51
19:DT:13:LEU:HB3	19:DT:18:TYR:OH	2.10	0.51
21:DV:151:HIS:ND1	21:DV:151:HIS:N	2.56	0.51
57:DY:99:SER:C	57:DY:100:ASN:O	2.48	0.51
57:DY:25:PHE:CE1	57:DY:82:PHE:CA	2.93	0.51
57:DY:89:ALA:HB3	56:DJ:15:ALA:CA	2.32	0.51
57:DY:92:THR:CG2	57:DY:93:LEU:N	2.74	0.51
57:DY:98:LYS:HG2	57:DY:102:LYS:CA	2.38	0.51
22:A3:55:ARG:HB3	22:A3:55:ARG:CZ	2.41	0.51
1:AA:1384:A:N3	1:AA:1405:U:H1'	2.25	0.51
1:AA:1729:A:H2'	1:AA:1730:U:H5''	1.92	0.51
1:AA:1761:C:C4	1:AA:1762:A:C2	2.98	0.51
1:AA:1819:A:H1'	1:AA:1821:A:C6	2.46	0.51
1:AA:2184:G:O2'	1:AA:2185:C:H5'	2.11	0.51
1:AA:2790:A:O2'	1:AA:2791:C:OP2	2.23	0.51
1:AA:562:U:O2'	1:AA:572:A:O4'	2.17	0.51
1:AA:654(F):C:H2'	1:AA:654(G):C:OP1	2.09	0.51
1:AA:68:G:N3	1:AA:68:G:H2'	2.26	0.51
3:AD:102:LYS:O	3:AD:103:ARG:HG2	2.10	0.51
3:AD:224:ALA:O	3:AD:225:ALA:HB2	2.11	0.51
5:AF:132:VAL:HG13	5:AF:133:ASN:H	1.75	0.51
5:AF:185:ASP:CG	5:AF:188:ARG:HH21	2.14	0.51
6:AG:37:VAL:HG23	6:AG:99:MET:HE3	1.93	0.51
6:AG:54:GLU:O	6:AG:57:ALA:HB3	2.11	0.51
7:AH:97:ARG:CG	7:AH:98:LEU:H	2.19	0.51
9:AM:129:PRO:O	9:AM:131:GLN:N	2.43	0.51
9:AM:138:LEU:HD23	9:AM:138:LEU:N	2.26	0.51
10:AN:107:ARG:HD3	15:AR:37:GLY:H	1.76	0.51
11:AO:107:LYS:C	11:AO:109:GLY:N	2.63	0.51
11:AO:19:VAL:HG21	11:AO:21:ARG:HD2	1.91	0.51
21:AV:103:ARG:N	21:AV:137:ILE:O	2.35	0.51
21:AV:106:GLY:HA2	21:AV:140:ASP:OD1	2.10	0.51
53:B1:30:C:H6	53:B1:30:C:O5'	1.93	0.51
53:B1:55:U:H6	53:B1:55:U:OP2	1.94	0.51
31:BA:865:A:H2'	31:BA:866:C:C6	2.46	0.51
1:AA:2602:A:H5''	52:BC:74:C:H5'	1.92	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:BD:12:U:H2'	52:BD:13:C:O4'	2.09	0.51
52:BD:44:G:O2'	52:BD:45:U:H5'	2.11	0.51
40:BM:6:ILE:HD11	40:BM:23:ILE:HG21	1.92	0.51
41:BN:115:PRO:C	41:BN:117:ASN:H	2.14	0.51
43:BP:67:GLU:N	43:BP:70:LEU:HD12	2.26	0.51
47:BT:60:ILE:HB	47:BT:74:LEU:HD23	1.93	0.51
48:BU:29:PHE:CE1	48:BU:31:LEU:HB3	2.42	0.51
50:BW:56:MET:O	50:BW:59:ALA:HB3	2.11	0.51
54:CA:1067:A:O2'	54:CA:1068:G:P	2.69	0.51
54:CA:1145:C:H5'	54:CA:1146:A:OP1	2.10	0.51
54:CA:487:A:C2	54:CA:488:C:H1'	2.46	0.51
54:CA:560:U:H4'	54:CA:561:U:C5'	2.41	0.51
54:CA:753:A:HO2'	54:CA:754:C:P	2.34	0.51
54:CA:963:G:N2	40:CM:55:LYS:HZ3	2.08	0.51
52:CB:21:A:N6	52:CB:46:G:C4	2.79	0.51
52:CD:18:G:H2'	52:CD:57:G:N2	2.26	0.51
52:CD:76:A:H8	55:DA:2394:C:N4	2.03	0.51
32:CE:238:LEU:HG	32:CE:238:LEU:O	2.11	0.51
33:CF:15:THR:HG23	33:CF:181:ASN:HA	1.92	0.51
33:CF:29:TYR:O	33:CF:29:TYR:HD2	1.93	0.51
54:CA:8:A:O2'	35:CH:102:ALA:C	2.49	0.51
44:CQ:3:ARG:O	44:CQ:6:LEU:N	2.35	0.51
36:CI:99:ALA:HB1	48:CU:23:LYS:HZ1	1.73	0.51
36:CI:60:PHE:CE2	48:CU:78:LEU:HD21	2.46	0.51
49:CV:42:PRO:HD3	26:D4:63:TYR:HE2	1.74	0.51
49:CV:44:MET:HA	49:CV:47:HIS:CD2	2.36	0.51
50:CW:26:ASN:O	50:CW:30:LYS:HB2	2.10	0.51
26:D4:55:ARG:C	26:D4:59:PHE:HB2	2.31	0.51
55:DA:1111:A:HO2'	55:DA:1112:G:H4'	1.76	0.51
55:DA:1204:A:O2'	55:DA:1205:U:O5'	2.29	0.51
55:DA:1317:A:H2'	55:DA:1318:C:H6	1.76	0.51
55:DA:1528:A:N1	55:DA:1543:A:H2	2.08	0.51
55:DA:155:C:H2'	55:DA:161:U:H5'	1.93	0.51
55:DA:1498:C:O4'	55:DA:1577:C:H4'	2.11	0.51
55:DA:1826:G:O2'	3:DD:242:ARG:NH2	2.44	0.51
55:DA:1946:U:H2'	55:DA:1947:C:H6	1.76	0.51
55:DA:1954:G:O2'	55:DA:1956:U:O4	2.25	0.51
55:DA:2141:G:H2'	55:DA:2142:C:O4'	2.09	0.51
55:DA:2199:A:H3'	55:DA:2205:C:C6	2.46	0.51
55:DA:242:G:N2	55:DA:254:G:H2'	2.26	0.51
55:DA:2490:G:H4'	55:DA:2491:U:O5'	2.10	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2664:G:O5'	55:DA:2664:G:H8	1.93	0.51
55:DA:2524:G:H2'	55:DA:2741:A:C2	2.46	0.51
55:DA:332:A:O2'	55:DA:333:G:OP1	2.28	0.51
55:DA:532:A:O2'	55:DA:2021:C:N4	2.44	0.51
55:DA:616:A:O2'	55:DA:617:G:OP1	2.23	0.51
3:DD:143:HIS:O	3:DD:144:ALA:C	2.49	0.51
3:DD:27:THR:CG2	3:DD:28:GLU:N	2.65	0.51
6:DG:51:ARG:HB3	6:DG:51:ARG:HH11	1.74	0.51
58:DL:112:MET:O	58:DL:113:PRO:C	2.48	0.51
58:DL:41:PHE:O	58:DL:42:ASN:C	2.49	0.51
58:DL:56:GLU:HB3	58:DL:68:VAL:O	2.11	0.51
58:DL:80:LYS:NZ	58:DL:107:ILE:HG13	2.25	0.51
9:DM:63:THR:HG23	9:DM:66:LYS:HE3	1.92	0.51
11:DO:86:LYS:HB3	11:DO:118:GLY:CA	2.41	0.51
12:DP:20:ALA:CB	12:DP:99:PRO:HB2	2.41	0.51
12:DP:52:VAL:HA	12:DP:55:VAL:HG13	1.91	0.51
18:DS:12:ILE:HD13	18:DS:17:VAL:HG22	1.93	0.51
25:DX:11:SER:OG	25:DX:13:ILE:HG12	2.11	0.51
57:DY:129:PRO:CD	57:DY:130:THR:N	2.73	0.51
13:A0:66:VAL:HG12	13:A0:70:LEU:HD12	1.92	0.51
22:A3:82:ARG:CG	22:A3:84:LEU:HD22	2.41	0.51
28:A6:41:PRO:HD2	28:A6:45:LYS:C	2.29	0.51
28:A6:52:VAL:HG13	28:A6:53:LYS:HG2	1.93	0.51
1:AA:1141:U:O5'	9:AM:63:THR:HG21	2.10	0.51
1:AA:1270:C:H5"	1:AA:1271:G:O5'	2.10	0.51
1:AA:1416:G:H2'	1:AA:1417:C:C5	2.45	0.51
1:AA:1493:C:H4'	1:AA:1494:A:OP2	2.11	0.51
1:AA:2376:A:H2'	1:AA:2377:A:O4'	2.11	0.51
1:AA:2756:U:H4'	1:AA:2757:A:O5'	2.11	0.51
1:AA:616:A:O2'	1:AA:617:G:P	2.68	0.51
1:AA:768:G:H2'	1:AA:769:G:H8	1.74	0.51
3:AD:130:ALA:HA	3:AD:192:THR:HA	1.93	0.51
1:AA:2591:C:OP1	3:AD:239:ARG:HG3	2.09	0.51
3:AD:70:TRP:CD1	3:AD:71:ASP:N	2.78	0.51
5:AF:181:LEU:HD21	5:AF:194:MET:HE1	1.92	0.51
7:AH:108:GLY:HA3	7:AH:152:ARG:NH2	2.25	0.51
8:AK:97:ILE:O	8:AK:101:LEU:CD2	2.57	0.51
11:AO:48:PRO:HG2	11:AO:49:ARG:N	2.23	0.51
12:AP:133:ARG:HH11	12:AP:133:ARG:HG3	1.74	0.51
18:AS:21:VAL:HG13	18:AS:74:ALA:HB1	1.93	0.51
21:AV:130:PRO:C	21:AV:133:ILE:CD1	2.79	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:107:G:C2'	31:BA:108:G:H5'	2.41	0.51
31:BA:562:C:O2'	42:BO:15:ARG:HD2	2.11	0.51
31:BA:908:A:H2'	31:BA:909:A:H8	1.74	0.51
32:BE:210:SER:O	32:BE:214:ILE:HG12	2.11	0.51
32:BE:76:GLN:OE1	32:BE:206:ASP:HB3	2.11	0.51
33:BF:155:GLY:O	33:BF:156:ARG:CB	2.56	0.51
34:BG:203:VAL:O	34:BG:206:PHE:HB3	2.10	0.51
37:BJ:22:LEU:CD1	37:BJ:97:GLN:HE22	2.23	0.51
38:BK:11:THR:HG23	38:BK:14:ARG:HH11	1.75	0.51
31:BA:1342:C:H4'	39:BL:125:TYR:HB3	1.92	0.51
39:BL:93:ARG:C	39:BL:95:LYS:H	2.14	0.51
40:BM:47:PHE:CE1	40:BM:63:PHE:HB2	2.45	0.51
40:BM:30:SER:CB	40:BM:81:THR:HA	2.41	0.51
41:BN:74:ALA:C	41:BN:76:GLY:H	2.13	0.51
31:BA:538:G:H3'	42:BO:115:LYS:HZ2	1.73	0.51
42:BO:22:SER:C	42:BO:24:VAL:H	2.14	0.51
42:BO:55:VAL:HG13	42:BO:68:ALA:C	2.31	0.51
42:BO:27:LEU:HD12	42:BO:61:THR:OG1	2.11	0.51
48:BU:85:LEU:HD12	48:BU:85:LEU:C	2.30	0.51
31:BA:1313:U:P	49:BV:6:LYS:HB3	2.51	0.51
54:CA:1329:A:P	43:CP:28:ALA:HB3	2.50	0.51
54:CA:556:C:O2'	54:CA:557:G:H5'	2.11	0.51
54:CA:792:A:C4	54:CA:794:A:C6	2.98	0.51
54:CA:830:G:H2'	54:CA:831:U:C6	2.46	0.51
54:CA:848:C:O5'	54:CA:848:C:H6	1.94	0.51
52:CB:51:U:H2'	52:CB:52:G:H8	1.76	0.51
52:CB:8:U:OP2	52:CB:8:U:C6	2.63	0.51
33:CF:70:VAL:CG1	33:CF:71:ALA:N	2.74	0.51
54:CA:921:U:O2'	35:CH:19:MET:O	2.18	0.51
37:CJ:88:PRO:O	37:CJ:89:MET:HB3	2.10	0.51
40:CM:58:ASP:O	40:CM:59:SER:C	2.48	0.51
40:CM:38:ILE:CG1	40:CM:71:LEU:HB3	2.41	0.51
42:CO:22:SER:C	42:CO:24:VAL:H	2.13	0.51
42:CO:27:LEU:C	42:CO:29:GLY:H	2.14	0.51
54:CA:1358:U:P	44:CQ:35:ARG:HG3	2.51	0.51
22:D3:10:THR:O	22:D3:10:THR:CG2	2.58	0.51
26:D4:68:ARG:HB3	26:D4:71:ARG:C	2.31	0.51
55:DA:2815:C:O2'	27:D5:43:HIS:CD2	2.64	0.51
28:D6:25:LYS:HD2	30:D8:34:TRP:HZ2	1.75	0.51
28:D6:15:GLU:OE2	28:D6:44:ARG:CZ	2.59	0.51
55:DA:254:G:N7	30:D8:5:LYS:HE2	2.25	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:550:G:N3	55:DA:1220:A:H2	2.09	0.51
55:DA:1419:A:C3'	55:DA:1420:U:H5''	2.41	0.51
55:DA:141:A:C8	55:DA:1408:C:H1'	2.45	0.51
55:DA:1458:C:H5''	55:DA:1459:G:H5'	1.93	0.51
55:DA:1533:C:O2	55:DA:1534:G:O6	2.29	0.51
55:DA:1706:U:O2	55:DA:1757:U:H5'	2.10	0.51
55:DA:2650:U:H2'	55:DA:2651:C:H6	1.75	0.51
55:DA:27:G:O2'	55:DA:28:A:P	2.69	0.51
55:DA:654(C):G:C3'	55:DA:654(D):G:C8	2.93	0.51
55:DA:660:G:O3'	5:DF:38:ARG:NH2	2.44	0.51
55:DA:903:C:H2'	55:DA:904:C:H6	1.75	0.51
55:DA:952:G:P	12:DP:16:ARG:HH12	2.33	0.51
5:DF:196:LEU:O	5:DF:200:GLU:HG2	2.10	0.51
43:CP:7:VAL:HG11	6:DG:115:ARG:NH2	2.26	0.51
7:DH:109:PHE:HB2	7:DH:111:HIS:O	2.10	0.51
7:DH:16:SER:O	7:DH:17:VAL:CB	2.59	0.51
58:DL:101:TRP:CE2	58:DL:140:GLY:CA	2.94	0.51
58:DL:10:LEU:CG	58:DL:55:VAL:HG11	2.40	0.51
9:DM:120:LEU:HD11	9:DM:122:VAL:HG23	1.92	0.51
9:DM:15:LEU:O	9:DM:136:GLU:HA	2.11	0.51
10:DN:8:LEU:HB2	10:DN:19:ILE:CD1	2.41	0.51
11:DO:38:GLN:C	11:DO:40:SER:N	2.60	0.51
15:DR:88:ILE:HD12	15:DR:90:GLN:H	1.75	0.51
18:DS:71:VAL:HA	18:DS:107:LEU:HD12	1.93	0.51
18:DS:92:ARG:NH1	18:DS:94:ASP:OD2	2.44	0.51
19:DT:12:VAL:HG13	19:DT:17:ALA:HB1	1.92	0.51
21:DV:111:VAL:HG23	21:DV:146:ILE:HG13	1.91	0.51
21:DV:125:LEU:HG	21:DV:164:ALA:CB	2.41	0.51
21:DV:114:GLY:CA	21:DV:179:ASP:OD1	2.57	0.51
24:DW:41:ILE:HD11	24:DW:44:LEU:CG	2.41	0.51
25:DX:19:GLN:NE2	25:DX:52:HIS:CE1	2.77	0.51
57:DY:27:VAL:CA	57:DY:111:LEU:HD13	2.29	0.51
55:DA:1084:A:C8	57:DY:53:VAL:HG11	2.46	0.51
1:AA:2840:C:H5''	13:A0:53:HIS:CD2	2.46	0.51
17:A2:44:LYS:HG2	17:A2:45:THR:N	2.25	0.51
17:A2:28:GLU:O	17:A2:61:VAL:HG21	2.11	0.51
13:A0:33:ARG:NH1	27:A5:55:ARG:HH21	2.09	0.51
30:A8:29:LYS:O	30:A8:31:HIS:N	2.44	0.51
1:AA:1114:G:H2'	1:AA:1115:G:H8	1.75	0.51
1:AA:1142(A):A:C5	1:AA:1144:G:C5	2.99	0.51
1:AA:1208:C:C4	1:AA:1209:G:N7	2.79	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1559:G:O2'	1:AA:1560:G:OP1	2.24	0.51
1:AA:1683:C:H2'	1:AA:1684:C:C6	2.46	0.51
1:AA:1747:G:H2'	1:AA:1748:G:H8	1.75	0.51
1:AA:1773:A:N7	1:AA:1829:A:H1'	2.25	0.51
1:AA:1:G:H2'	1:AA:2:G:H8	1.74	0.51
1:AA:2191:G:N2	1:AA:2192:G:H1'	2.26	0.51
1:AA:2347:C:H2'	1:AA:2348:U:C6	2.46	0.51
1:AA:2348:U:H4'	28:A6:42:TRP:HD1	1.75	0.51
1:AA:646:A:C8	1:AA:2349:G:N2	2.78	0.51
1:AA:2420:C:H41	30:A8:31:HIS:CB	2.16	0.51
1:AA:2391:G:H1'	1:AA:2429:G:N2	2.25	0.51
1:AA:2469:A:H1'	1:AA:2482:G:C6	2.46	0.51
1:AA:2490:G:C5'	1:AA:2491:U:OP1	2.58	0.51
1:AA:2522:U:H2'	1:AA:2523:G:H5''	1.92	0.51
1:AA:27:G:H1'	1:AA:513:A:H62	1.76	0.51
1:AA:548:A:H2'	1:AA:549:G:H5'	1.92	0.51
3:AD:92:ILE:HD12	3:AD:104:TYR:CD2	2.46	0.51
3:AD:215:LEU:HB2	3:AD:217:ARG:HG3	1.93	0.51
3:AD:35:LYS:HE3	3:AD:65:ILE:HG22	1.92	0.51
5:AF:18:ARG:HG2	5:AF:19:GLU:N	2.25	0.51
6:AG:75:LYS:HG3	6:AG:76:SER:H	1.76	0.51
8:AK:82:ARG:NH1	8:AK:146:ALA:CB	2.74	0.51
9:AM:21:LYS:O	9:AM:61:ARG:N	2.42	0.51
9:AM:11:PRO:HB2	9:AM:51:PHE:CE1	2.45	0.51
12:AP:88:GLY:O	12:AP:89:ASN:CB	2.59	0.51
14:AQ:48:LEU:N	14:AQ:48:LEU:HD12	2.26	0.51
14:AQ:52:SER:O	14:AQ:56:LEU:HD22	2.11	0.51
21:AV:116:VAL:N	21:AV:177:PRO:HG3	2.25	0.51
21:AV:30:ASN:HA	21:AV:89:PHE:HE2	1.76	0.51
21:AV:5:LEU:CD2	21:AV:47:VAL:HG21	2.35	0.51
21:AV:53:ILE:HD12	21:AV:53:ILE:C	2.31	0.51
25:AX:54:VAL:CG1	25:AX:55:ARG:N	2.73	0.51
23:AZ:81:LYS:HB3	23:AZ:81:LYS:NZ	2.26	0.51
31:BA:1073:U:H2'	31:BA:1074:G:H8	1.76	0.51
31:BA:1053:G:C6	31:BA:1199:U:H2'	2.45	0.51
31:BA:1298:C:C6	37:BJ:114:ARG:NH1	2.79	0.51
31:BA:1388:C:H2'	31:BA:1389:C:C6	2.46	0.51
31:BA:151:A:H2'	31:BA:152:A:O4'	2.10	0.51
31:BA:199:G:O2'	31:BA:200:G:H5'	2.10	0.51
31:BA:411:A:C4	31:BA:413:G:H1'	2.46	0.51
31:BA:426:G:O2'	31:BA:427:U:H5'	2.10	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:428:G:H4'	31:BA:429:U:O5'	2.10	0.51
31:BA:65:U:H4'	31:BA:66:G:O5'	2.11	0.51
31:BA:696:A:H2'	31:BA:697:U:C5'	2.26	0.51
31:BA:868:C:H2'	31:BA:869:G:O4'	2.11	0.51
1:AA:2554:U:C2	52:BB:74:C:C5	2.99	0.51
32:BE:26:PRO:C	32:BE:28:PHE:H	2.13	0.51
34:BG:61:LYS:HZ1	34:BG:62:GLN:NE2	2.05	0.51
34:BG:70:ILE:HD12	34:BG:100:ARG:CZ	2.41	0.51
35:BH:76:ILE:HG22	35:BH:78:HIS:H	1.75	0.51
36:BI:100:ASN:HB3	48:BU:28:GLU:HA	1.92	0.51
39:BL:5:TYR:OH	39:BL:16:ARG:HG2	2.11	0.51
42:BO:86:ARG:HB2	42:BO:101:VAL:HG22	1.91	0.51
43:BP:81:LEU:HD22	43:BP:88:ARG:HG2	1.92	0.51
48:BU:41:LYS:HD3	48:BU:41:LYS:C	2.29	0.51
54:CA:1004:A:H5''	54:CA:1025:U:C4	2.45	0.51
54:CA:1049:U:H4'	54:CA:1050:G:OP2	2.10	0.51
54:CA:1064:G:O2'	54:CA:1065:U:O5'	2.28	0.51
54:CA:1370:G:C2	54:CA:1371:G:C8	2.98	0.51
54:CA:22:G:O2'	54:CA:23:C:H5'	2.11	0.51
54:CA:864:A:H2'	54:CA:865:A:C8	2.45	0.51
54:CA:1104:G:O5'	32:CE:111:ARG:HD2	2.11	0.51
32:CE:240:GLN:O	32:CE:240:GLN:HG2	2.11	0.51
34:CG:114:ARG:O	34:CG:117:ALA:HB3	2.11	0.51
34:CG:198:VAL:CG1	34:CG:199:ASN:N	2.73	0.51
35:CH:135:THR:O	35:CH:138:ALA:HB3	2.11	0.51
35:CH:140:ARG:CB	35:CH:140:ARG:HH11	2.23	0.51
35:CH:78:HIS:CD2	38:CK:104:ARG:HE	2.28	0.51
36:CI:59:TYR:HD2	36:CI:61:LEU:HD11	1.76	0.51
40:CM:13:HIS:CG	40:CM:14:LYS:N	2.78	0.51
40:CM:51:ARG:HG3	40:CM:60:ARG:C	2.31	0.51
40:CM:48:THR:HG23	40:CM:62:HIS:CG	2.46	0.51
43:CP:92:HIS:HD2	43:CP:110:ARG:NH2	2.09	0.51
43:CP:44:ARG:O	43:CP:46:LYS:N	2.42	0.51
49:CV:5:LEU:HD13	49:CV:5:LEU:C	2.31	0.51
13:D0:85:PRO:C	13:D0:87:TYR:H	2.13	0.51
16:D1:108:GLU:O	16:D1:111:GLU:HB2	2.10	0.51
22:D3:36:ILE:N	22:D3:36:ILE:HD13	2.25	0.51
55:DA:1538:G:N2	55:DA:1539:G:C4	2.79	0.51
55:DA:1340:U:H1'	55:DA:1603:A:H5'	1.92	0.51
55:DA:2360:A:H8	55:DA:2360:A:O5'	1.92	0.51
55:DA:957:A:N6	55:DA:2459:A:C8	2.79	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:270(L):U:H3	8:DK:50:ARG:NE	2.09	0.51
55:DA:307:G:H21	55:DA:330:A:H62	1.59	0.51
55:DA:774:A:H2	55:DA:787:U:HO2'	1.52	0.51
55:DA:802:A:H2'	55:DA:803:U:C5'	2.41	0.51
3:DD:168:ARG:NH1	3:DD:168:ARG:HG3	2.26	0.51
6:DG:146:TYR:C	6:DG:148:MET:N	2.64	0.51
7:DH:12:PRO:O	7:DH:13:LYS:HB2	2.10	0.51
8:DK:49:ALA:O	8:DK:52:ARG:HG2	2.10	0.51
8:DK:61:ARG:HH21	8:DK:64:GLU:CD	2.13	0.51
58:DL:19:PRO:O	58:DL:20:ALA:CB	2.59	0.51
9:DM:5:VAL:O	9:DM:5:VAL:HG13	2.11	0.51
9:DM:62:VAL:HG21	9:DM:87:LEU:HD11	1.92	0.51
12:DP:34:LEU:HD11	12:DP:129:THR:HB	1.92	0.51
14:DQ:48:LEU:HD23	14:DQ:82:ILE:HD11	1.91	0.51
18:DS:42:ARG:HG2	18:DS:42:ARG:NH1	2.26	0.51
18:DS:60:ASN:N	18:DS:60:ASN:HD22	2.07	0.51
19:DT:64:LYS:NZ	19:DT:73:ARG:NH2	2.59	0.51
20:DU:94:LYS:NZ	20:DU:101:LYS:NZ	2.59	0.51
57:DY:120:LYS:O	57:DY:121:ASP:HB3	2.11	0.51
57:DY:40:LEU:HD22	57:DY:41:ARG:N	2.25	0.51
16:A1:47:TYR:HA	16:A1:50:ARG:NH1	2.25	0.51
17:A2:1:MET:HG2	17:A2:42:GLY:HA3	1.92	0.51
17:A2:59:ALA:HB2	17:A2:96:ILE:HD13	1.93	0.51
22:A3:53:MET:CB	22:A3:59:LEU:HD23	2.40	0.51
26:A4:34:GLU:OE1	43:BP:3:ARG:HB2	2.11	0.51
30:A8:33:ASN:O	30:A8:34:TRP:CB	2.58	0.51
1:AA:1053:C:H2'	1:AA:1054:A:H5''	1.92	0.51
1:AA:1111:A:H4'	7:AH:3:ARG:CD	2.35	0.51
1:AA:1144:G:H2'	1:AA:1145:C:H6	1.76	0.51
1:AA:1394:U:H3'	1:AA:1394:U:H6	1.76	0.51
1:AA:1525:G:H2'	1:AA:1526:G:H8	1.76	0.51
1:AA:2182:G:H2'	1:AA:2183:C:H6	1.76	0.51
1:AA:274:G:H2'	1:AA:275:G:O4'	2.10	0.51
1:AA:319:C:H2'	1:AA:320:A:O4'	2.11	0.51
1:AA:363(B):G:H8	1:AA:363(B):G:H5'	1.76	0.51
1:AA:376:C:H2'	1:AA:377:C:C6	2.46	0.51
1:AA:503:A:H5''	1:AA:504:U:OP1	2.11	0.51
1:AA:833:U:H4'	11:AO:51:PHE:O	2.11	0.51
1:AA:83:G:C2	1:AA:102:G:H2'	2.46	0.51
1:AA:956:G:N2	1:AA:959:A:H3'	2.26	0.51
2:AB:79:C:C2'	2:AB:80:U:H5'	2.41	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:AD:34:VAL:HG13	3:AD:104:TYR:HE1	1.73	0.51
4:AE:6:GLY:CA	4:AE:28:ALA:HA	2.41	0.51
4:AE:76:ARG:O	4:AE:78:LEU:N	2.44	0.51
6:AG:38:VAL:HG22	6:AG:93:THR:HA	1.92	0.51
11:AO:59:LEU:HD22	11:AO:60:MET:N	2.26	0.51
14:AQ:83:LYS:HG2	14:AQ:109:GLY:N	2.25	0.51
14:AQ:26:LEU:HB3	14:AQ:87:PHE:HA	1.93	0.51
24:AW:14:ARG:HA	24:AW:63:VAL:HG11	1.93	0.51
24:AW:15:LYS:HD3	24:AW:67:LYS:NZ	2.26	0.51
25:AX:19:GLN:HE22	25:AX:52:HIS:CE1	2.28	0.51
31:BA:1124:G:O2'	31:BA:1125:U:P	2.69	0.51
31:BA:1162:C:O2'	31:BA:1163:C:H5'	2.11	0.51
31:BA:1167:A:OP1	31:BA:1167:A:H8	1.94	0.51
31:BA:1210:C:H4'	31:BA:1214:C:N4	2.25	0.51
31:BA:181:G:O2'	31:BA:182:U:O5'	2.29	0.51
31:BA:977:A:O2'	31:BA:979:C:OP2	2.24	0.51
32:BE:96:ARG:O	32:BE:98:LEU:N	2.43	0.51
31:BA:1190:G:OP2	33:BF:5:ILE:HG23	2.10	0.51
34:BG:108:LEU:HB3	34:BG:110:PHE:CD1	2.45	0.51
36:BI:42:GLU:C	36:BI:44:GLY:N	2.64	0.51
31:BA:1346:A:N7	37:BJ:10:ARG:NH2	2.59	0.51
38:BK:82:HIS:CD2	38:BK:138:TRP:NE1	2.68	0.51
39:BL:20:ARG:O	39:BL:22:GLY:N	2.44	0.51
40:BM:89:ASP:C	40:BM:90:LEU:HD12	2.31	0.51
41:BN:38:ASN:N	41:BN:38:ASN:HD22	2.09	0.51
43:BP:4:ILE:HG23	43:BP:5:ALA:N	2.19	0.51
31:BA:956:U:H4'	49:BV:83:HIS:HB3	1.92	0.51
31:BA:1286:A:H4'	51:BX:25:LYS:HD2	1.92	0.51
54:CA:134:A:H1'	54:CA:325:A:C5	2.46	0.51
54:CA:1534:A:N3	54:CA:1535:C:N4	2.42	0.51
54:CA:6:G:H4'	54:CA:298:A:H4'	1.92	0.51
54:CA:309:G:O2'	54:CA:607:A:N1	2.43	0.51
54:CA:96:G:H2'	54:CA:97:U:C4'	2.40	0.51
32:CE:104:ASN:O	32:CE:108:ILE:HG12	2.11	0.51
32:CE:124:SER:HB2	32:CE:125:PRO:HD2	1.92	0.51
33:CF:127:ARG:HG2	33:CF:127:ARG:HH11	1.76	0.51
36:CI:10:LEU:HA	36:CI:84:ASN:O	2.10	0.51
39:CL:9:ARG:O	39:CL:104:ARG:HD2	2.11	0.51
40:CM:92:THR:HG23	40:CM:93:GLY:N	2.26	0.51
43:CP:65:LYS:HB2	43:CP:69:GLU:HB3	1.93	0.51
47:CT:77:VAL:HG12	47:CT:77:VAL:O	2.11	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1509:C:H3'	55:DA:1510:A:C4'	2.40	0.51
55:DA:1511:A:C2'	55:DA:1512:G:H5'	2.40	0.51
55:DA:1531:C:O2'	55:DA:1532:C:H5'	2.10	0.51
55:DA:1652:A:O3'	55:DA:1653:G:H8	1.94	0.51
55:DA:2427:C:C5'	55:DA:2428:G:OP1	2.59	0.51
55:DA:2563:U:H4'	10:DN:28:SER:HA	1.92	0.51
55:DA:2869:G:H2'	55:DA:2870:C:C6	2.46	0.51
55:DA:322:A:H1'	55:DA:339:U:O2	2.10	0.51
55:DA:704:G:C2'	55:DA:726:G:N2	2.70	0.51
55:DA:704:G:H1'	55:DA:727:A:N6	2.25	0.51
3:DD:98:VAL:C	3:DD:100:GLY:N	2.62	0.51
55:DA:1354:A:OP1	3:DD:38:LYS:HE2	2.10	0.51
4:DE:107:THR:O	4:DE:190:GLY:HA2	2.10	0.51
6:DG:143:GLU:HG3	26:D4:31:ILE:CD1	2.41	0.51
7:DH:109:PHE:O	7:DH:111:HIS:N	2.37	0.51
8:DK:132:PRO:O	8:DK:133:HIS:O	2.28	0.51
58:DL:132:ARG:O	58:DL:137:GLU:OE1	2.29	0.51
58:DL:13:PRO:O	58:DL:14:ALA:HB2	2.11	0.51
15:DR:105:LEU:HD23	15:DR:105:LEU:N	2.26	0.51
15:DR:54:ARG:NH1	15:DR:54:ARG:HG2	2.25	0.51
18:DS:1:MET:HG3	18:DS:64:MET:HE3	1.92	0.51
55:DA:71:A:C2	19:DT:31:HIS:HE1	2.28	0.51
19:DT:90:GLU:O	19:DT:91:ALA:C	2.50	0.51
20:DU:86:ARG:O	20:DU:92:ASN:HA	2.11	0.51
21:DV:107:THR:OG1	21:DV:108:PRO:CD	2.59	0.51
21:DV:142:SER:O	21:DV:143:GLY:O	2.29	0.51
21:DV:26:GLY:HA2	21:DV:86:VAL:O	2.11	0.51
25:DX:59:VAL:HG12	25:DX:60:GLU:N	2.26	0.51
57:DY:134:LEU:HA	57:DY:137:GLU:HB3	1.93	0.51
55:DA:1083:U:OP2	57:DY:45:LYS:O	2.29	0.51
57:DY:47:ASN:O	57:DY:48:GLY:C	2.48	0.51
57:DY:88:ALA:CB	57:DY:92:THR:OG1	2.54	0.51
16:A1:104:GLN:O	16:A1:107:ALA:HB3	2.11	0.51
16:A1:98:LEU:CB	16:A1:102:GLU:HB2	2.41	0.51
27:A5:20:ARG:HA	27:A5:23:HIS:CD2	2.46	0.51
1:AA:1043:C:C4	1:AA:1044:G:N7	2.79	0.51
1:AA:1115:G:O2'	1:AA:1116:C:H5'	2.10	0.51
1:AA:1616:A:C2	1:AA:1616:A:OP1	2.64	0.51
1:AA:1789:A:H2'	1:AA:1790:C:C6	2.45	0.51
1:AA:1913:A:O2'	1:AA:1914:C:OP2	2.27	0.51
1:AA:387:U:HO2'	1:AA:388:G:P	2.27	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:498:G:H21	20:AU:47:LYS:NZ	2.09	0.51
1:AA:88:G:H5'	1:AA:90:U:H5	1.75	0.51
1:AA:946:G:HO2'	1:AA:947:G:C4'	2.24	0.51
4:AE:25:VAL:HG13	4:AE:181:LEU:HD12	1.92	0.51
4:AE:70:ALA:O	4:AE:72:VAL:N	2.44	0.51
15:AR:8:LYS:C	15:AR:10:VAL:N	2.60	0.51
20:AU:94:LYS:CE	20:AU:101:LYS:HZ3	2.23	0.51
25:AX:54:VAL:HG12	25:AX:55:ARG:H	1.74	0.51
31:BA:1075:C:H4'	31:BA:1101:A:C6	2.46	0.51
31:BA:1320:C:N3	31:BA:1321:C:N4	2.59	0.51
31:BA:250:A:O2'	31:BA:251:G:P	2.69	0.51
31:BA:812:C:O2'	31:BA:813:U:H6	1.94	0.51
31:BA:89:U:H4'	31:BA:90:C:OP1	2.10	0.51
31:BA:991:U:O2	31:BA:993:G:H8	1.95	0.51
32:BE:223:ILE:HA	32:BE:226:ARG:HB3	1.92	0.51
33:BF:33:LEU:O	33:BF:35:GLU:N	2.44	0.51
39:BL:53:VAL:O	39:BL:55:ALA:N	2.44	0.51
47:BT:56:VAL:O	47:BT:77:VAL:HB	2.11	0.51
47:BT:84:LEU:O	47:BT:87:LYS:HB2	2.11	0.51
54:CA:1054:C:N3	52:CB:34:G:H1'	2.26	0.51
54:CA:1157:A:N6	54:CA:1178:G:N2	2.59	0.51
54:CA:1216:G:H5''	44:CQ:5:ALA:CB	2.41	0.51
54:CA:1298:C:H4'	54:CA:1299:A:C8	2.46	0.51
54:CA:340:U:H2'	54:CA:341:C:H6	1.76	0.51
52:CD:19:G:O4'	52:CD:57:G:N2	2.44	0.51
54:CA:1190:G:P	33:CF:5:ILE:HD12	2.50	0.51
33:CF:6:HIS:CB	44:CQ:49:HIS:HD2	2.24	0.51
34:CG:106:TYR:CE1	34:CG:112:VAL:O	2.64	0.51
38:CK:38:ILE:O	38:CK:42:GLU:HG2	2.11	0.51
49:CV:30:LEU:C	49:CV:30:LEU:HD22	2.31	0.51
49:CV:7:LYS:HB2	49:CV:7:LYS:HZ3	1.75	0.51
13:D0:33:ARG:HG2	13:D0:34:ILE:N	2.24	0.51
13:D0:78:LYS:O	13:D0:82:GLU:HB2	2.10	0.51
22:D3:25:ARG:HA	22:D3:29:GLN:NE2	2.26	0.51
26:D4:56:VAL:O	26:D4:60:GLN:HG2	2.11	0.51
30:D8:14:VAL:HG13	30:D8:15:LYS:N	2.25	0.51
55:DA:1149:G:H2'	55:DA:1150:C:C6	2.46	0.51
55:DA:1204:A:H1'	55:DA:1206:G:C8	2.46	0.51
55:DA:1251:C:O2'	55:DA:1252:G:H5''	2.11	0.51
55:DA:1489:U:O2'	55:DA:1490:A:H8	1.94	0.51
55:DA:1528:A:C2	55:DA:1543:A:N1	2.79	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1794:U:H2'	55:DA:1795:C:C6	2.45	0.51
55:DA:1934:C:H2'	55:DA:1935:G:O5'	2.11	0.51
55:DA:2159:G:H2'	55:DA:2160:G:O4'	2.11	0.51
55:DA:2286:A:C8	55:DA:2287:A:C6	2.99	0.51
55:DA:2656:U:O4	55:DA:2665:A:C6	2.64	0.51
55:DA:2:G:H2'	55:DA:3:U:O4'	2.11	0.51
55:DA:847:U:C5	55:DA:933:A:N1	2.79	0.51
2:DB:44:G:H1'	2:DB:47:C:N4	2.25	0.51
5:DF:107:LYS:HD2	5:DF:206:ILE:CD1	2.29	0.51
55:DA:1248:G:N2	5:DF:88:VAL:HG22	2.26	0.51
7:DH:132:ARG:CB	7:DH:132:ARG:HH11	2.24	0.51
56:DI:14:GLN:C	56:DI:16:THR:H	2.14	0.51
58:DL:144:VAL:CG1	58:DL:145:LYS:H	2.14	0.51
58:DL:60:TYR:N	58:DL:60:TYR:CD1	2.79	0.51
9:DM:133:GLN:O	9:DM:134:ARG:HB3	2.10	0.51
14:DQ:65:VAL:O	14:DQ:69:VAL:HG12	2.11	0.51
14:DQ:5:THR:C	14:DQ:7:TYR:N	2.64	0.51
18:DS:11:ARG:NH2	18:DS:99:ARG:O	2.42	0.51
21:DV:109:ALA:O	21:DV:110:GLY:O	2.29	0.51
57:DY:18:GLU:HG3	57:DY:66:LEU:HD11	1.93	0.51
57:DY:70:GLU:C	57:DY:71:LEU:CD1	2.70	0.51
57:DY:7:VAL:O	57:DY:9:LEU:N	2.44	0.51
13:A0:23:ASN:N	13:A0:23:ASN:HD22	2.09	0.50
17:A2:74:LYS:NZ	17:A2:74:LYS:HB3	2.26	0.50
26:A4:14:ILE:CG2	26:A4:21:VAL:HG23	2.40	0.50
1:AA:2394:C:OP1	30:A8:30:ARG:NH1	2.44	0.50
1:AA:1028:A:H61	1:AA:1125:G:H2'	1.76	0.50
1:AA:1204:A:C2	1:AA:1241:A:C2	2.99	0.50
1:AA:1311:G:H21	1:AA:1603:A:H62	1.59	0.50
1:AA:1761:C:N4	1:AA:1762:A:C2	2.79	0.50
1:AA:1885:A:H3'	1:AA:1886:C:C6	2.37	0.50
1:AA:2012:G:H4'	18:AS:96:ILE:CD1	2.42	0.50
1:AA:2165:G:N2	1:AA:2166:G:H1'	2.25	0.50
1:AA:2331:G:O3'	22:A3:43:THR:HG22	2.11	0.50
1:AA:2464:C:H2'	1:AA:2465:C:H6	1.76	0.50
1:AA:760:G:H2'	1:AA:761:A:O4'	2.11	0.50
5:AF:37:VAL:O	5:AF:38:ARG:C	2.48	0.50
6:AG:106:LEU:HD12	6:AG:110:ALA:CB	2.41	0.50
6:AG:7:LEU:HD21	6:AG:176:LEU:HD22	1.93	0.50
7:AH:107:VAL:O	7:AH:152:ARG:NH2	2.44	0.50
11:AO:93:GLY:O	11:AO:94:GLU:HB2	2.11	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AP:56:ARG:CB	12:AP:56:ARG:HH11	2.21	0.50
12:AP:68:ILE:HD13	12:AP:103:MET:HG2	1.92	0.50
15:AR:106:SER:C	15:AR:107:ASP:OD1	2.49	0.50
18:AS:20:VAL:CG2	18:AS:47:VAL:HG21	2.40	0.50
18:AS:82:LEU:HB2	18:AS:98:LYS:HB2	1.93	0.50
20:AU:13:VAL:HG22	20:AU:14:LEU:N	2.26	0.50
21:AV:177:PRO:O	21:AV:180:VAL:CA	2.59	0.50
24:AW:17:SER:HA	24:AW:20:GLU:CG	2.41	0.50
24:AW:51:ARG:HB2	24:AW:55:ARG:NH1	2.26	0.50
31:BA:1048:G:H2'	31:BA:1050:G:H8	1.75	0.50
31:BA:1054:C:O2'	31:BA:1055:A:C5'	2.59	0.50
31:BA:1347:G:H2'	31:BA:1373:G:O6	2.10	0.50
31:BA:481:G:H1'	31:BA:482:A:N7	2.26	0.50
31:BA:812:C:O2'	31:BA:813:U:C6	2.61	0.50
52:BC:58:A:O2'	52:BC:59:U:P	2.69	0.50
32:BE:200:ILE:H	32:BE:200:ILE:HD12	1.77	0.50
33:BF:34:LEU:C	33:BF:38:ARG:HH21	2.14	0.50
33:BF:8:ILE:O	33:BF:11:ARG:N	2.36	0.50
35:BH:12:LEU:HD23	35:BH:13:ILE:N	2.26	0.50
37:BJ:146:GLU:OE1	37:BJ:149:ARG:HD2	2.09	0.50
39:BL:42:ARG:NH1	39:BL:71:SER:HA	2.26	0.50
42:BO:46:LYS:HG2	42:BO:47:LYS:H	1.75	0.50
53:C1:39:U:H2'	53:C1:40:U:C6	2.45	0.50
54:CA:1054:C:N4	52:CB:34:G:N9	2.59	0.50
54:CA:1500:A:OP2	54:CA:1505:G:OP1	2.28	0.50
54:CA:262:A:N6	54:CA:263:A:N6	2.59	0.50
54:CA:339:C:H2'	54:CA:340:U:C6	2.45	0.50
54:CA:518:C:H1'	54:CA:529:G:C2	2.46	0.50
54:CA:562:C:N3	42:CO:16:GLU:HB3	2.26	0.50
54:CA:665:A:N3	54:CA:732:C:H2'	2.25	0.50
54:CA:758:G:H8	54:CA:758:G:O5'	1.94	0.50
52:CD:47:U:OP2	52:CD:47:U:O4'	2.29	0.50
52:CD:70:G:H2'	52:CD:71:G:C8	2.47	0.50
32:CE:91:PRO:HG3	32:CE:155:LEU:HD23	1.93	0.50
54:CA:1055:A:H4'	33:CF:161:GLU:CD	2.31	0.50
33:CF:181:ASN:HD22	33:CF:204:LEU:HB2	1.76	0.50
33:CF:60:ALA:O	33:CF:61:ALA:HB2	2.11	0.50
33:CF:83:ARG:O	33:CF:85:ARG:N	2.43	0.50
34:CG:144:ASP:O	34:CG:146:ILE:HD12	2.11	0.50
34:CG:199:ASN:O	34:CG:200:GLU:C	2.50	0.50
34:CG:84:LYS:N	34:CG:84:LYS:HD2	2.25	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:CM:40:LEU:HB3	40:CM:41:PRO:HD2	1.94	0.50
42:CO:83:VAL:HG21	42:CO:100:ILE:CG1	2.39	0.50
42:CO:28:LYS:HZ1	42:CO:33:ARG:HH22	1.56	0.50
55:DA:534:U:O2'	16:D1:49:HIS:CD2	2.64	0.50
55:DA:1103:A:H2'	55:DA:1104:C:C5'	2.33	0.50
55:DA:1231:G:H2'	55:DA:1232:G:C8	2.46	0.50
55:DA:1268:A:H2'	55:DA:1269:A:O4'	2.10	0.50
55:DA:1372:U:C5	55:DA:1372:U:C5'	2.81	0.50
55:DA:1486:A:H2'	55:DA:1487:G:H8	1.75	0.50
55:DA:1525:G:H2'	55:DA:1526:G:C8	2.46	0.50
55:DA:1655:A:H2'	55:DA:1656:C:O4'	2.12	0.50
55:DA:201:C:C2'	55:DA:202:U:H5'	2.41	0.50
55:DA:2092:U:C2	55:DA:2225:A:O2'	2.64	0.50
55:DA:2677:G:H2'	55:DA:2678:C:H6	1.75	0.50
4:DE:13:ARG:HB3	4:DE:21:VAL:CG1	2.39	0.50
4:DE:154:LYS:HD3	4:DE:155:LYS:N	2.26	0.50
4:DE:104:VAL:HG22	4:DE:198:VAL:HG22	1.93	0.50
6:DG:7:LEU:N	6:DG:104:GLU:OE2	2.33	0.50
6:DG:121:ASN:HD22	6:DG:122:PRO:N	2.10	0.50
6:DG:129:GLY:HA2	6:DG:169:ALA:CB	2.41	0.50
7:DH:22:GLY:O	7:DH:37:VAL:HB	2.10	0.50
8:DK:8:PRO:HG3	8:DK:14:ASP:HB2	1.92	0.50
12:DP:5:ARG:O	12:DP:6:ARG:HB3	2.09	0.50
14:DQ:32:LEU:O	14:DQ:62:LYS:HE2	2.11	0.50
17:A2:39:LEU:N	17:A2:39:LEU:HD12	2.25	0.50
22:A3:25:ARG:HG3	22:A3:29:GLN:NE2	2.26	0.50
22:A3:50:ASN:HA	22:A3:62:LEU:HD12	1.93	0.50
1:AA:1072:C:H2'	1:AA:1093:G:O6	2.11	0.50
1:AA:1140:C:C4'	1:AA:1143:A:N6	2.74	0.50
1:AA:1192:G:O2'	1:AA:1193:G:H5'	2.10	0.50
1:AA:1205:U:H4'	1:AA:1206:G:OP2	2.11	0.50
1:AA:1844:C:O2'	1:AA:1845:G:H5'	2.12	0.50
1:AA:1914:C:C3'	1:AA:1914:C:O2	2.58	0.50
1:AA:191:A:H2'	1:AA:192:C:C6	2.46	0.50
1:AA:2655:G:C2'	1:AA:2656:U:OP2	2.59	0.50
1:AA:1758:G:C2	1:AA:2696:U:H5'	2.46	0.50
1:AA:600:G:H5'	5:AF:32:LEU:HD12	1.93	0.50
1:AA:620:G:C5'	1:AA:621:A:OP1	2.53	0.50
1:AA:757:U:H2'	1:AA:758:C:C6	2.44	0.50
1:AA:920:G:O2'	1:AA:921:G:H5'	2.11	0.50
3:AD:166:GLN:CA	3:AD:166:GLN:NE2	2.74	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:AD:65:ILE:HD12	3:AD:65:ILE:C	2.31	0.50
1:AA:442:G:C4'	5:AF:46:ARG:HD3	2.40	0.50
7:AH:22:GLY:HA3	7:AH:37:VAL:H	1.76	0.50
7:AH:89:ILE:H	7:AH:89:ILE:CD1	2.22	0.50
8:AK:84:GLY:O	8:AK:85:GLU:HB2	2.09	0.50
18:AS:27:LYS:O	18:AS:70:TYR:HB2	2.11	0.50
20:AU:95:LYS:CB	20:AU:100:ALA:HA	2.39	0.50
20:AU:63:LYS:NZ	20:AU:64:GLU:H	1.94	0.50
12:AP:134:ARG:NH2	21:AV:122:ARG:HH12	2.08	0.50
31:BA:1129:C:C2	31:BA:1132:C:N4	2.76	0.50
31:BA:32:A:H4'	31:BA:48:C:H41	1.76	0.50
31:BA:394:G:H2'	31:BA:395:C:C6	2.45	0.50
31:BA:542:G:OP1	34:BG:10:ARG:NH2	2.39	0.50
31:BA:579:G:H2'	31:BA:580:U:C6	2.46	0.50
31:BA:676:A:O2'	31:BA:677:U:H5'	2.11	0.50
31:BA:722:A:HO2'	31:BA:723:U:H5	1.58	0.50
31:BA:890:G:O2'	31:BA:906:G:N1	2.44	0.50
31:BA:938:A:H2'	31:BA:939:G:O4'	2.11	0.50
32:BE:172:ILE:HG22	32:BE:172:ILE:O	2.11	0.50
33:BF:11:ARG:O	33:BF:12:LEU:C	2.50	0.50
33:BF:59:ARG:HE	33:BF:64:VAL:CG2	2.24	0.50
36:BI:48:LEU:HD13	36:BI:52:ILE:CG1	2.40	0.50
37:BJ:103:TRP:NE1	37:BJ:137:LYS:HD3	2.26	0.50
37:BJ:27:ILE:HD11	37:BJ:43:PHE:CD2	2.47	0.50
39:BL:27:THR:HG23	39:BL:31:GLN:N	2.25	0.50
47:BT:13:ASP:C	47:BT:15:MET:H	2.15	0.50
48:BU:75:ILE:C	48:BU:77:GLY:H	2.14	0.50
49:BV:41:VAL:CG1	49:BV:42:PRO:CD	2.40	0.50
54:CA:1131:G:H1	54:CA:1144:G:H21	1.57	0.50
54:CA:1414:U:H2'	54:CA:1415:G:H8	1.76	0.50
52:CC:72:C:O2'	52:CC:73:A:H5'	2.11	0.50
54:CA:1112:C:H1'	33:CF:179:ARG:HD3	1.93	0.50
34:CG:88:VAL:O	34:CG:88:VAL:HG12	2.11	0.50
35:CH:101:ILE:HD13	35:CH:101:ILE:H	1.76	0.50
38:CK:101:PRO:HG2	38:CK:133:LEU:HD11	1.92	0.50
39:CL:80:GLY:O	39:CL:81:ILE:C	2.49	0.50
43:CP:123:ALA:CB	43:CP:124:PRO:CD	2.86	0.50
55:DA:2723:C:C5'	13:D0:1:MET:HG2	2.40	0.50
17:D2:38:LEU:H	17:D2:51:VAL:HG13	1.76	0.50
55:DA:1104:C:H2'	55:DA:1105:U:C6	2.46	0.50
55:DA:1188:U:H2'	55:DA:1189:A:C5'	2.41	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1535:U:O2	55:DA:1535:U:C3'	2.59	0.50
55:DA:228:A:N3	55:DA:228:A:H2'	2.26	0.50
55:DA:372:G:H2'	55:DA:373:U:OP2	2.10	0.50
55:DA:531:C:H5''	55:DA:532:A:O4'	2.10	0.50
55:DA:949:C:O2'	55:DA:950:G:H5'	2.11	0.50
2:DB:78:A:C2	2:DB:99:A:C4	3.00	0.50
3:DD:108:PRO:HB3	3:DD:143:HIS:HE1	1.73	0.50
55:DA:1800:C:OP1	3:DD:264:LYS:HE2	2.11	0.50
4:DE:167:VAL:HG11	4:DE:187:ALA:O	2.11	0.50
4:DE:29:GLY:HA2	4:DE:180:ASN:HB3	1.92	0.50
5:DF:46:ARG:O	5:DF:47:GLY:C	2.49	0.50
5:DF:75:HIS:CE1	5:DF:82:ILE:HD12	2.46	0.50
6:DG:111:LEU:HB2	6:DG:112:PRO:HD3	1.93	0.50
6:DG:16:ARG:HB3	6:DG:17:PRO:HD3	1.92	0.50
8:DK:120:ILE:HD12	8:DK:121:LYS:N	2.26	0.50
58:DL:104:VAL:CG1	58:DL:105:LEU:N	2.61	0.50
9:DM:90:MET:CE	9:DM:90:MET:HA	2.41	0.50
11:DO:90:ARG:CZ	11:DO:91:PHE:HB3	2.41	0.50
12:DP:32:TYR:CE1	12:DP:133:ARG:HG3	2.45	0.50
12:DP:43:THR:HG1	12:DP:46:GLN:HG3	1.76	0.50
12:DP:51:ARG:O	12:DP:55:VAL:HG12	2.10	0.50
20:DU:39:VAL:CG1	20:DU:40:GLU:H	2.12	0.50
57:DY:38:HIS:CD2	57:DY:40:LEU:O	2.65	0.50
23:DZ:52:ARG:HA	23:DZ:57:GLU:HA	1.93	0.50
23:DZ:97:LEU:HD23	23:DZ:98:LEU:N	2.26	0.50
17:A2:32:THR:HG22	17:A2:58:VAL:HG12	1.93	0.50
17:A2:77:ALA:O	17:A2:78:LYS:HG2	2.11	0.50
17:A2:79:VAL:O	17:A2:80:GLN:CG	2.59	0.50
17:A2:7:THR:C	17:A2:9:GLY:H	2.15	0.50
1:AA:1062:G:H2'	1:AA:1063:G:C8	2.46	0.50
1:AA:1113:U:H2'	1:AA:1114:G:H8	1.75	0.50
1:AA:1275:A:O2'	1:AA:1276:A:C8	2.65	0.50
1:AA:1430:C:H2'	1:AA:1431:U:C6	2.46	0.50
1:AA:758:C:O2	1:AA:1981:A:H2	1.94	0.50
1:AA:2092:U:H6	1:AA:2092:U:H5''	1.77	0.50
1:AA:2159:G:H2'	1:AA:2160:G:H8	1.76	0.50
1:AA:2303:G:C2'	1:AA:2304:G:H5'	2.42	0.50
1:AA:2297:C:N4	1:AA:2320:A:H8	2.09	0.50
1:AA:2346:A:C2	1:AA:2383:G:C2	3.00	0.50
1:AA:2348:U:C4'	28:A6:42:TRP:HD1	2.24	0.50
1:AA:2467:C:H2'	1:AA:2468:G:O4'	2.11	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2602:A:N6	52:BB:76:A:C5'	2.68	0.50
1:AA:1786:A:H2	1:AA:2606:C:H1'	1.73	0.50
1:AA:372:G:O2'	1:AA:373:U:P	2.69	0.50
1:AA:528:A:C3'	1:AA:528:A:C8	2.94	0.50
1:AA:649:G:H2'	1:AA:650:C:C6	2.46	0.50
1:AA:83:G:O2'	1:AA:84:A:C8	2.65	0.50
1:AA:889:C:C4	1:AA:890:A:H1'	2.46	0.50
1:AA:897:C:N3	1:AA:898:C:H5	2.08	0.50
4:AE:202:LYS:N	4:AE:202:LYS:HE3	2.27	0.50
6:AG:61:ALA:HA	6:AG:64:THR:CG2	2.42	0.50
6:AG:7:LEU:HB2	6:AG:104:GLU:CD	2.32	0.50
8:AK:1:MET:HG3	8:AK:23:PRO:HA	1.94	0.50
1:AA:2416:C:OP1	11:AO:64:LYS:O	2.29	0.50
11:AO:91:PHE:N	11:AO:91:PHE:CD1	2.79	0.50
12:AP:58:PHE:CD1	12:AP:61:GLY:HA2	2.46	0.50
14:AQ:89:ARG:O	14:AQ:90:GLY:C	2.49	0.50
18:AS:9:TYR:H	18:AS:102:HIS:CD2	2.25	0.50
19:AT:55:ASN:O	19:AT:79:ALA:HA	2.11	0.50
20:AU:27:VAL:HG12	20:AU:39:VAL:CG1	2.40	0.50
20:AU:52:SER:N	20:AU:53:PRO:CD	2.74	0.50
53:B1:52:U:OP1	53:B1:52:U:C4'	2.59	0.50
31:BA:1160:G:H1	31:BA:1177:G:N2	1.97	0.50
31:BA:1176:A:N6	31:BA:1177:G:O6	2.45	0.50
31:BA:434:U:H2'	31:BA:435:C:H6	1.76	0.50
31:BA:690:G:O2'	31:BA:691:G:H5'	2.10	0.50
31:BA:780:A:C2	31:BA:803:G:C6	3.00	0.50
31:BA:765:G:N2	31:BA:813:U:H5	2.09	0.50
31:BA:980:C:H5''	31:BA:981:U:C5	2.47	0.50
52:BD:59:U:C2'	52:BD:60:U:H5'	2.42	0.50
32:BE:52:GLU:O	32:BE:56:ARG:HG3	2.11	0.50
36:BI:14:LEU:HD22	36:BI:19:LEU:HB2	1.93	0.50
39:BL:112:LYS:HD3	39:BL:113:LYS:O	2.12	0.50
40:BM:54:PHE:C	40:BM:55:LYS:HG3	2.31	0.50
40:BM:54:PHE:CG	40:BM:55:LYS:N	2.79	0.50
40:BM:7:LYS:HE2	40:BM:71:LEU:HD22	1.92	0.50
42:BO:102:ARG:O	42:BO:104:VAL:HG23	2.11	0.50
43:BP:108:ARG:NH2	43:BP:114:ARG:HG2	2.25	0.50
43:BP:81:LEU:C	43:BP:83:ASP:H	2.14	0.50
54:CA:107:G:H2'	54:CA:108:G:C5'	2.39	0.50
54:CA:1094:G:O2'	54:CA:1095:U:OP2	2.27	0.50
54:CA:1261:A:H2'	54:CA:1262:C:H5'	1.93	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1380:U:H5''	54:CA:1381:U:OP1	2.11	0.50
54:CA:328:C:H4'	54:CA:329:A:O5'	2.12	0.50
54:CA:721:G:H4'	54:CA:722:A:C5'	2.42	0.50
54:CA:728:A:C6	45:CR:54:ARG:HD2	2.47	0.50
54:CA:942:G:O2'	54:CA:943:U:H5'	2.11	0.50
32:CE:95:GLN:O	32:CE:96:ARG:C	2.50	0.50
34:CG:88:VAL:HA	35:CH:97:GLY:HA2	1.92	0.50
35:CH:137:GLU:CA	35:CH:140:ARG:NH1	2.73	0.50
35:CH:62:ALA:O	35:CH:65:ASN:N	2.45	0.50
38:CK:48:TYR:HA	38:CK:60:ARG:O	2.11	0.50
38:CK:49:GLU:O	38:CK:51:VAL:HG13	2.10	0.50
38:CK:97:VAL:HG13	38:CK:98:LYS:HG3	1.93	0.50
39:CL:95:LYS:C	39:CL:95:LYS:HD3	2.31	0.50
42:CO:47:LYS:CB	42:CO:48:PRO:HD2	2.40	0.50
43:CP:44:ARG:C	43:CP:46:LYS:N	2.63	0.50
43:CP:7:VAL:CG1	6:DG:115:ARG:NH2	2.75	0.50
54:CA:280:C:C2	47:CT:38:ARG:HG3	2.46	0.50
54:CA:263:A:P	50:CW:79:ARG:NH1	2.85	0.50
13:D0:17:ARG:O	13:D0:20:LEU:HB3	2.12	0.50
17:D2:81:TYR:C	17:D2:82:ARG:CG	2.80	0.50
30:D8:39:LYS:O	30:D8:43:GLN:HB2	2.11	0.50
55:DA:1083:U:O5'	57:DY:41:ARG:HD3	2.10	0.50
55:DA:1056:G:H1'	55:DA:1086:A:C1'	2.41	0.50
55:DA:1278:A:H2'	55:DA:1279:G:C8	2.46	0.50
55:DA:1750:G:O2'	55:DA:1751:C:H5'	2.11	0.50
55:DA:1930:G:H2'	55:DA:1968:G:O6	2.10	0.50
55:DA:1991:U:H2'	55:DA:1992:G:C5'	2.41	0.50
55:DA:265:A:HO2'	55:DA:266:G:P	2.35	0.50
55:DA:92:G:H2'	55:DA:93:C:C6	2.47	0.50
3:DD:62:TYR:CG	3:DD:63:ARG:N	2.79	0.50
55:DA:2311:A:H8	6:DG:82:LEU:HD11	1.76	0.50
7:DH:16:SER:O	7:DH:17:VAL:HB	2.11	0.50
55:DA:1064:C:O3'	58:DL:89:HIS:CB	2.60	0.50
10:DN:26:LYS:HB2	10:DN:30:ALA:HB2	1.94	0.50
11:DO:140:ALA:O	11:DO:141:ALA:HB2	2.10	0.50
14:DQ:20:ARG:C	14:DQ:22:GLY:H	2.14	0.50
20:DU:43:ASN:HA	20:DU:64:GLU:HA	1.93	0.50
21:DV:115:GLY:HA3	21:DV:146:ILE:HD11	1.93	0.50
21:DV:170:THR:O	21:DV:171:ILE:CB	2.58	0.50
24:DW:8:LYS:NZ	24:DW:8:LYS:HB2	2.26	0.50
25:DX:6:VAL:HG11	25:DX:47:VAL:HG13	1.92	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:A1:90:VAL:HG12	16:A1:91:ASP:N	2.26	0.50
16:A1:40:PHE:HE2	17:A2:83:ARG:HH22	1.59	0.50
22:A3:43:THR:C	22:A3:45:PHE:H	2.14	0.50
1:AA:1001:A:H2'	1:AA:1002:G:H5'	1.94	0.50
1:AA:1340:U:O2'	1:AA:1341:U:OP2	2.30	0.50
1:AA:1347:G:N3	29:A7:49:ARG:NH2	2.58	0.50
1:AA:1530:G:H2'	1:AA:1531:C:H6	1.76	0.50
1:AA:1728:G:N1	1:AA:1730:U:OP2	2.44	0.50
1:AA:1814:G:H2'	1:AA:1815:A:N7	2.25	0.50
1:AA:1967:C:H2'	1:AA:1968:G:C5'	2.39	0.50
1:AA:2129:C:N4	1:AA:2130:U:O4	2.44	0.50
1:AA:2178:C:O2'	1:AA:2179:C:H5'	2.11	0.50
1:AA:234:C:O2'	1:AA:235:U:H5'	2.12	0.50
1:AA:2528:U:O2'	1:AA:2529:G:H5''	2.11	0.50
1:AA:2563:U:H2'	1:AA:2565:A:OP2	2.10	0.50
1:AA:2848:G:HO2'	1:AA:2849:U:P	2.34	0.50
1:AA:452:G:N3	1:AA:457:A:H2	2.08	0.50
1:AA:455:C:H5''	1:AA:456:C:OP2	2.11	0.50
1:AA:529:A:C5'	1:AA:530:G:OP1	2.57	0.50
1:AA:580:C:H2'	1:AA:581:C:H6	1.75	0.50
1:AA:729:G:H5'	1:AA:730:C:H5''	1.93	0.50
1:AA:805:G:C4'	1:AA:806:C:OP2	2.59	0.50
1:AA:89:G:OP2	1:AA:90:U:H6	1.94	0.50
1:AA:950:G:H2'	1:AA:951:C:C6	2.47	0.50
2:AB:11:C:H3'	2:AB:12:C:C5	2.46	0.50
4:AE:91:VAL:O	4:AE:91:VAL:HG13	2.11	0.50
6:AG:108:ASN:O	26:A4:37:SER:HB2	2.11	0.50
6:AG:136:ARG:O	6:AG:154:GLY:CA	2.56	0.50
6:AG:51:ARG:CB	6:AG:51:ARG:NH1	2.75	0.50
7:AH:33:LEU:HD11	7:AH:79:VAL:HG13	1.93	0.50
1:AA:598:G:H5'	11:AO:11:GLY:HA3	1.94	0.50
12:AP:133:ARG:O	12:AP:134:ARG:CB	2.59	0.50
14:AQ:61:ASN:O	14:AQ:65:VAL:HG23	2.11	0.50
18:AS:50:VAL:HG11	18:AS:103:ILE:HG21	1.92	0.50
20:AU:46:LYS:O	20:AU:48:ALA:N	2.42	0.50
20:AU:42:VAL:CG2	20:AU:65:ALA:HB3	2.41	0.50
20:AU:83:THR:HG22	20:AU:85:VAL:H	1.77	0.50
21:AV:178:GLU:O	21:AV:179:ASP:C	2.48	0.50
31:BA:1020:U:H2'	31:BA:1021:G:C5'	2.40	0.50
31:BA:1056:U:C5'	33:BF:163:ALA:HB2	2.42	0.50
31:BA:1075:C:H4'	31:BA:1101:A:N6	2.26	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:1124:G:O2'	31:BA:1125:U:O5'	2.30	0.50
31:BA:1220:G:H4'	49:BV:34:TRP:O	2.11	0.50
31:BA:1236:A:H4'	31:BA:1304:G:H4'	1.92	0.50
32:BE:92:TYR:HE2	32:BE:151:GLY:N	2.09	0.50
33:BF:154:SER:O	33:BF:155:GLY:O	2.30	0.50
31:BA:1106:G:H4'	33:BF:171:GLY:O	2.11	0.50
36:BI:44:GLY:O	36:BI:46:ARG:HG3	2.12	0.50
37:BJ:44:TYR:HA	37:BJ:47:CYS:SG	2.51	0.50
43:BP:110:ARG:HG2	43:BP:110:ARG:HH11	1.76	0.50
48:BU:22:VAL:O	48:BU:24:ALA:N	2.45	0.50
49:BV:47:HIS:O	49:BV:48:THR:HG23	2.11	0.50
54:CA:1054:C:O2	54:CA:1054:C:H2'	2.10	0.50
54:CA:1116:C:H2'	54:CA:1117:G:H5'	1.92	0.50
32:CE:83:MET:O	32:CE:86:GLU:N	2.44	0.50
33:CF:52:LEU:CD2	33:CF:52:LEU:H	2.24	0.50
34:CG:9:CYS:C	34:CG:11:LEU:N	2.65	0.50
38:CK:42:GLU:HG3	38:CK:109:ILE:CD1	2.37	0.50
43:CP:70:LEU:O	43:CP:74:VAL:HG23	2.11	0.50
47:CT:13:ASP:C	47:CT:15:MET:H	2.14	0.50
49:CV:25:LYS:HA	49:CV:25:LYS:HZ1	1.76	0.50
50:CW:96:GLY:O	50:CW:97:ALA:CB	2.58	0.50
16:D1:24:TYR:HE1	16:D1:39:LEU:CD2	2.25	0.50
55:DA:1019:U:H2'	55:DA:1020:A:C8	2.46	0.50
55:DA:1280:G:C2'	55:DA:1281:G:C5'	2.87	0.50
55:DA:1281:G:O2'	55:DA:1282:U:H5'	2.11	0.50
55:DA:1614:A:N1	18:DS:91:GLY:HA2	2.26	0.50
55:DA:1785:A:H2'	55:DA:1787:A:N7	2.26	0.50
55:DA:1831:G:H2'	55:DA:1832:C:C6	2.47	0.50
55:DA:2297:C:O2'	55:DA:2298:A:H5'	2.11	0.50
55:DA:2453:A:O2'	55:DA:2572:A:H1'	2.11	0.50
55:DA:2646:C:OP2	55:DA:2732:G:O2'	2.26	0.50
55:DA:2785:C:H2'	55:DA:2786:U:O4'	2.10	0.50
55:DA:654(T):A:H2'	55:DA:654(U):A:O4'	2.11	0.50
55:DA:6:A:H2'	55:DA:7:G:O4'	2.12	0.50
4:DE:10:GLY:O	4:DE:25:VAL:HG22	2.12	0.50
4:DE:131:ALA:O	4:DE:132:HIS:CB	2.60	0.50
7:DH:16:SER:HB3	7:DH:26:VAL:O	2.12	0.50
8:DK:52:ARG:HH11	8:DK:52:ARG:CB	2.25	0.50
11:DO:35:HIS:O	11:DO:36:LYS:C	2.48	0.50
12:DP:118:LEU:HD13	12:DP:131:ILE:HG23	1.93	0.50
12:DP:75:THR:HA	12:DP:88:GLY:O	2.11	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:DS:14:PRO:HG2	18:DS:78:GLU:CG	2.41	0.50
57:DY:127:GLU:O	57:DY:128:LEU:CD2	2.51	0.50
57:DY:20:ALA:C	57:DY:21:GLN:O	2.45	0.50
57:DY:12:THR:HB	57:DY:52:PHE:CE2	2.46	0.50
16:A1:5:LYS:O	16:A1:6:THR:C	2.50	0.50
22:A3:47:PRO:HB2	22:A3:51:VAL:O	2.12	0.50
1:AA:141:A:C8	1:AA:1408:C:H1'	2.45	0.50
1:AA:1678:G:N2	1:AA:1989:G:H1	2.09	0.50
1:AA:1802:A:N1	1:AA:1822:G:H1'	2.27	0.50
1:AA:1879:C:H2'	1:AA:1880:C:H5'	1.93	0.50
1:AA:2173:A:C6	1:AA:2174:C:H1'	2.46	0.50
1:AA:2271:G:H2'	1:AA:2272:U:C6	2.46	0.50
1:AA:2495:G:OP1	22:A3:4:LYS:HE3	2.12	0.50
1:AA:2787:C:O2	1:AA:2787:C:H2'	2.11	0.50
1:AA:2808:U:H2'	1:AA:2809:A:H8	1.75	0.50
1:AA:361:G:N3	1:AA:362:U:H1'	2.26	0.50
1:AA:552:G:H2'	1:AA:553:U:O4'	2.11	0.50
1:AA:887:A:N6	1:AA:889:C:C6	2.73	0.50
1:AA:896:A:C2	21:AV:178:GLU:CG	2.90	0.50
1:AA:922:U:H2'	1:AA:923:C:H6	1.71	0.50
3:AD:182:LEU:O	3:AD:271:ILE:HG13	2.12	0.50
4:AE:16:ARG:O	4:AE:17:ASP:HB3	2.12	0.50
4:AE:199:ARG:CB	4:AE:199:ARG:NH1	2.75	0.50
4:AE:52:LEU:O	4:AE:74:PRO:CA	2.57	0.50
5:AF:185:ASP:HA	5:AF:188:ARG:HE	1.77	0.50
6:AG:61:ALA:HA	6:AG:64:THR:HG22	1.93	0.50
7:AH:17:VAL:HA	7:AH:26:VAL:HA	1.91	0.50
11:AO:52:GLU:CB	11:AO:55:ARG:HD2	2.41	0.50
12:AP:75:THR:HG22	12:AP:88:GLY:CA	2.28	0.50
14:AQ:14:VAL:O	14:AQ:18:ILE:HG13	2.10	0.50
15:AR:27:THR:CG2	15:AR:90:GLN:HB3	2.40	0.50
18:AS:55:ALA:O	18:AS:56:ALA:C	2.49	0.50
31:BA:1532:U:O5'	31:BA:1532:U:H6	1.94	0.50
31:BA:277:C:H5'	47:BT:68:ARG:NH1	2.27	0.50
31:BA:373:A:C2	31:BA:374:A:C8	2.99	0.50
31:BA:394:G:H2'	31:BA:395:C:H6	1.76	0.50
31:BA:518:C:H5'	31:BA:519:C:H6	1.77	0.50
31:BA:657:G:O2'	31:BA:658:G:H5'	2.12	0.50
52:BC:18:G:O6	52:BC:55:U:H1'	2.11	0.50
32:BE:114:ARG:HA	32:BE:117:GLU:CG	2.42	0.50
32:BE:12:GLU:O	32:BE:14:GLY:N	2.44	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:BE:61:LEU:HG	32:BE:68:ILE:HD11	1.94	0.50
34:BG:100:ARG:NH1	34:BG:137:SER:CB	2.74	0.50
34:BG:96:LEU:HD12	34:BG:139:ARG:CZ	2.41	0.50
35:BH:72:GLN:O	35:BH:73:ASN:HB2	2.12	0.50
36:BI:1:MET:HB2	36:BI:66:GLU:HG2	1.94	0.50
37:BJ:153:HIS:C	37:BJ:155:ARG:H	2.15	0.50
38:BK:85:ARG:HH11	38:BK:85:ARG:HG3	1.76	0.50
40:BM:30:SER:HB2	40:BM:81:THR:HA	1.94	0.50
48:BU:52:PRO:O	48:BU:56:THR:HG23	2.11	0.50
31:BA:191(F):U:O2	50:BW:105:SER:HA	2.11	0.50
54:CA:1069:C:H2'	54:CA:1070:U:O5'	2.11	0.50
54:CA:1400:C:C4'	54:CA:1401:G:OP2	2.60	0.50
54:CA:243:A:C2	54:CA:245:C:H2'	2.47	0.50
54:CA:344:A:H3'	54:CA:346:G:O6	2.11	0.50
54:CA:381:C:H2'	54:CA:382:A:O4'	2.12	0.50
54:CA:522:C:H41	42:CO:53:ARG:NH2	2.10	0.50
54:CA:780:A:C2	54:CA:803:G:C6	3.00	0.50
54:CA:922:G:C2	54:CA:923:A:C4	3.00	0.50
52:CB:21:A:N7	52:CB:46:G:C6	2.80	0.50
32:CE:168:THR:HG23	32:CE:169:LYS:H	1.77	0.50
32:CE:47:THR:O	32:CE:50:GLU:HB2	2.11	0.50
33:CF:131:ARG:HG3	33:CF:131:ARG:NH1	2.26	0.50
34:CG:112:VAL:HG12	34:CG:116:GLN:CD	2.31	0.50
34:CG:58:LEU:O	34:CG:58:LEU:HD22	2.12	0.50
39:CL:17:VAL:HG21	39:CL:80:GLY:HA3	1.92	0.50
40:CM:31:GLY:HA2	40:CM:78:ASN:ND2	2.26	0.50
40:CM:45:ARG:HG3	40:CM:45:ARG:HH11	1.76	0.50
44:CQ:24:CYS:SG	44:CQ:27:CYS:N	2.84	0.50
45:CR:17:ARG:NH1	45:CR:77:ARG:CZ	2.75	0.50
46:CS:53:VAL:HG23	46:CS:54:GLU:N	2.26	0.50
48:CU:50:ILE:HD11	48:CU:74:ARG:NH1	2.26	0.50
16:D1:88:ILE:N	16:D1:88:ILE:HD13	2.23	0.50
55:DA:1204:A:O2'	55:DA:1205:U:P	2.69	0.50
55:DA:1262:A:N3	27:D5:10:LYS:HE3	2.27	0.50
55:DA:1347:G:O2'	55:DA:1348:G:H5'	2.11	0.50
55:DA:1798:U:H5'	3:DD:259:THR:CG2	2.29	0.50
55:DA:1936:A:N3	55:DA:1945:G:C6	2.80	0.50
55:DA:1992:G:N2	55:DA:1996:C:O2'	2.44	0.50
55:DA:445:C:C2'	55:DA:446:G:H5'	2.41	0.50
55:DA:790:C:O2'	55:DA:791:C:H5'	2.11	0.50
55:DA:833:U:H2'	55:DA:834:C:C6	2.47	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:DD:27:THR:O	3:DD:28:GLU:CB	2.59	0.50
3:DD:95:LEU:HD11	3:DD:97:TYR:HE1	1.76	0.50
58:DL:52:ILE:HG12	58:DL:76:TYR:CA	2.41	0.50
58:DL:87:GLY:C	58:DL:96:VAL:HG11	2.32	0.50
9:DM:96:GLU:CG	9:DM:97:ARG:N	2.73	0.50
9:DM:94:HIS:CD2	9:DM:97:ARG:HH21	2.30	0.50
10:DN:10:VAL:HG23	10:DN:17:ARG:O	2.11	0.50
54:CA:346:G:C5'	15:DR:41:ARG:HD2	2.41	0.50
19:DT:28:PHE:CD1	19:DT:28:PHE:N	2.79	0.50
20:DU:43:ASN:CB	20:DU:64:GLU:HA	2.41	0.50
55:DA:1075:C:C4'	21:DV:195:GLU:CG	2.58	0.50
57:DY:25:PHE:O	57:DY:112:LEU:HA	2.12	0.50
17:A2:3:ALA:HB1	17:A2:38:LEU:HD22	1.94	0.50
26:A4:38:LYS:C	26:A4:38:LYS:HD2	2.30	0.50
1:AA:1022:G:O2'	1:AA:1023:U:OP2	2.30	0.50
1:AA:1130:U:O2	1:AA:2025:C:H5''	2.11	0.50
1:AA:976:C:C5'	1:AA:1156:A:N6	2.72	0.50
1:AA:1205:U:O2'	1:AA:1206:G:OP1	2.30	0.50
1:AA:1373:A:H2'	1:AA:1374:G:O4'	2.12	0.50
1:AA:13:A:H61	1:AA:525:U:H3'	1.77	0.50
1:AA:1484:G:C3'	1:AA:1485:G:H5''	2.41	0.50
1:AA:1490:A:H4'	1:AA:1491:G:OP2	2.12	0.50
1:AA:1930:G:H2'	1:AA:1931:U:OP2	2.12	0.50
1:AA:1940:U:H5'	1:AA:1965:C:C5	2.47	0.50
1:AA:2173:A:H5''	1:AA:2174:C:C6	2.47	0.50
1:AA:644:A:H61	1:AA:2349:G:H1'	1.76	0.50
1:AA:2355:C:H5'	22:A3:36:ILE:CD1	2.33	0.50
1:AA:2898:U:C2	1:AA:2899:G:C8	3.00	0.50
1:AA:530:G:H2'	1:AA:531:C:OP2	2.11	0.50
1:AA:65:C:O2'	1:AA:66:C:H5'	2.12	0.50
1:AA:780:G:OP1	3:AD:218:ARG:NH2	2.44	0.50
1:AA:894:C:C3'	1:AA:895:U:H6	2.08	0.50
1:AA:864:G:H1'	1:AA:914:C:N4	2.27	0.50
2:AB:30:C:H4'	2:AB:58:A:H2	1.77	0.50
4:AE:10:GLY:O	4:AE:11:MET:CB	2.60	0.50
5:AF:117:ARG:HH22	5:AF:187:VAL:HA	1.76	0.50
6:AG:2:PRO:C	6:AG:4:ASP:H	2.15	0.50
11:AO:27:HIS:ND1	11:AO:27:HIS:N	2.59	0.50
21:AV:128:VAL:HG22	21:AV:129:SER:N	2.26	0.50
21:AV:180:VAL:O	21:AV:181:GLU:O	2.30	0.50
31:BA:60:A:N6	31:BA:110:C:N3	2.58	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:1114:C:H2'	31:BA:1115:C:H6	1.76	0.50
31:BA:1356:G:H2'	31:BA:1357:A:C8	2.46	0.50
31:BA:429:U:H5'	34:BG:9:CYS:HB2	1.93	0.50
31:BA:593:G:O2'	31:BA:594:G:H5'	2.11	0.50
32:BE:36:ARG:HH11	32:BE:36:ARG:HG3	1.77	0.50
33:BF:117:ALA:HB2	33:BF:200:ALA:HB2	1.93	0.50
33:BF:58:GLU:HB2	33:BF:65:ALA:CB	2.41	0.50
35:BH:140:ARG:HG2	35:BH:140:ARG:O	2.12	0.50
31:BA:1080:A:OP1	35:BH:47:LYS:HE3	2.11	0.50
38:BK:63:LEU:H	38:BK:63:LEU:HD22	1.76	0.50
39:BL:4:TYR:CE2	39:BL:88:TYR:HB2	2.46	0.50
41:BN:16:SER:HA	41:BN:79:SER:O	2.12	0.50
43:BP:15:VAL:C	43:BP:17:VAL:N	2.65	0.50
43:BP:66:LEU:HA	43:BP:70:LEU:HD12	1.94	0.50
46:BS:21:VAL:HG13	46:BS:21:VAL:O	2.11	0.50
54:CA:1178:G:C8	54:CA:1180:A:OP2	2.65	0.50
54:CA:1250:A:H2'	54:CA:1251:A:C8	2.47	0.50
54:CA:31:G:O2'	54:CA:32:A:O5'	2.27	0.50
54:CA:41:G:H2'	54:CA:42:G:C8	2.46	0.50
32:CE:12:GLU:C	32:CE:14:GLY:N	2.65	0.50
32:CE:122:PHE:HA	32:CE:139:LYS:HZ3	1.75	0.50
32:CE:157:ARG:HG2	32:CE:158:LEU:HD12	1.94	0.50
32:CE:168:THR:HB	32:CE:192:SER:OG	2.11	0.50
33:CF:8:ILE:HG23	33:CF:16:ARG:HG2	1.94	0.50
33:CF:62:ASP:HA	33:CF:97:LYS:CD	2.42	0.50
34:CG:52:SER:HB3	34:CG:55:ALA:HB2	1.92	0.50
54:CA:430:A:OP1	34:CG:9:CYS:HB2	2.11	0.50
37:CJ:95:ARG:HH11	37:CJ:95:ARG:HG3	1.75	0.50
54:CA:562:C:H2'	42:CO:16:GLU:O	2.10	0.50
42:CO:60:LEU:HD22	42:CO:60:LEU:N	2.26	0.50
26:D4:39:CYS:C	26:D4:41:PRO:CD	2.80	0.50
26:D4:50:VAL:O	26:D4:52:THR:N	2.45	0.50
27:D5:40:LYS:HB2	27:D5:46:CYS:SG	2.51	0.50
27:D5:36:CYS:SG	27:D5:48:GLU:O	2.60	0.50
27:D5:54:GLY:O	27:D5:55:ARG:C	2.49	0.50
55:DA:1177:A:H5''	55:DA:1178:C:C5'	2.42	0.50
55:DA:1751:C:O2'	55:DA:1752:C:H5'	2.11	0.50
55:DA:1887:C:H2'	55:DA:1888:G:H5''	1.93	0.50
55:DA:2188:C:H2'	55:DA:2189:U:O4'	2.12	0.50
55:DA:2418:A:H2'	55:DA:2419:U:C6	2.47	0.50
55:DA:270(Q):C:H2'	55:DA:270(R):G:O4'	2.12	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2808:U:O2'	55:DA:2809:A:H5'	2.12	0.50
55:DA:721:C:O2	55:DA:721:C:H2'	2.11	0.50
55:DA:664:C:H4'	55:DA:941:A:OP1	2.12	0.50
6:DG:129:GLY:HA2	6:DG:169:ALA:HB2	1.93	0.50
7:DH:103:LEU:CD2	7:DH:115:VAL:HB	2.41	0.50
7:DH:125:VAL:HG12	7:DH:125:VAL:O	2.12	0.50
58:DL:12:LEU:HD12	58:DL:13:PRO:C	2.31	0.50
58:DL:58:THR:CB	58:DL:66:THR:CG2	2.90	0.50
2:DB:116:G:H4'	14:DQ:54:LEU:CD1	2.41	0.50
14:DQ:88:ASP:CG	14:DQ:90:GLY:H	2.14	0.50
15:DR:86:ILE:O	15:DR:86:ILE:HG12	2.12	0.50
21:DV:180:VAL:HG13	21:DV:181:GLU:H	1.72	0.50
57:DY:75:GLN:OE1	57:DY:109:SER:CB	2.59	0.50
16:A1:17:ILE:HG23	16:A1:39:LEU:HD12	1.94	0.50
16:A1:5:LYS:NZ	16:A1:5:LYS:CB	2.74	0.50
1:AA:1161:C:C1'	17:A2:8:GLY:O	2.56	0.50
29:A7:10:ARG:O	29:A7:14:LYS:HB2	2.12	0.50
1:AA:1992:G:O2'	1:AA:1993:U:OP2	2.22	0.50
1:AA:2300:G:H2'	1:AA:2301:C:H6	1.77	0.50
1:AA:270:A:C2'	1:AA:270(A):A:H5'	2.42	0.50
1:AA:26:G:C2	1:AA:27:G:N2	2.80	0.50
1:AA:2886:G:H2'	1:AA:2887:U:H6	1.77	0.50
1:AA:322:A:O4'	1:AA:322:A:OP2	2.30	0.50
1:AA:35:G:H2'	1:AA:36:G:O4'	2.11	0.50
1:AA:945:A:O4'	1:AA:946:G:OP1	2.30	0.50
2:AB:56:G:H4'	2:AB:57:A:C8	2.46	0.50
2:AB:7:G:H3'	2:AB:8:U:C5'	2.36	0.50
1:AA:1695:G:N7	3:AD:14:ARG:NH2	2.59	0.50
3:AD:25:THR:C	3:AD:27:THR:H	2.14	0.50
5:AF:3:GLU:HA	5:AF:24:LEU:CD2	2.40	0.50
5:AF:65:TRP:HZ3	5:AF:73:ALA:O	1.94	0.50
6:AG:111:LEU:HB2	6:AG:112:PRO:HD3	1.92	0.50
7:AH:13:LYS:O	7:AH:15:VAL:HG13	2.11	0.50
8:AK:82:ARG:HH11	8:AK:146:ALA:H	1.60	0.50
1:AA:952:G:P	12:AP:16:ARG:HH22	2.35	0.50
20:AU:47:LYS:C	20:AU:49:VAL:H	2.15	0.50
20:AU:47:LYS:HG3	20:AU:60:PHE:CD1	2.45	0.50
21:AV:158:PRO:O	21:AV:161:VAL:CG2	2.60	0.50
21:AV:179:ASP:O	21:AV:179:ASP:OD2	2.30	0.50
31:BA:1027:C:C2	31:BA:1028:C:C5	2.99	0.50
31:BA:1353:G:O2'	31:BA:1354:C:H5'	2.12	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:152:A:C8	31:BA:153:C:C5	3.00	0.50
31:BA:792:A:O2'	31:BA:793:U:P	2.69	0.50
52:BD:58:A:O2'	52:BD:59:U:P	2.70	0.50
32:BE:114:ARG:HA	32:BE:117:GLU:HG3	1.93	0.50
33:BF:173:VAL:N	33:BF:174:PRO:HD3	2.27	0.50
42:BO:79:GLU:HG3	42:BO:80:HIS:CG	2.47	0.50
42:BO:91:LYS:HG2	42:BO:91:LYS:O	2.11	0.50
43:BP:94:ARG:HG3	43:BP:96:LEU:HG	1.94	0.50
44:BQ:26:ARG:HD3	44:BQ:43:CYS:SG	2.52	0.50
49:BV:63:THR:H	49:BV:66:MET:CE	2.01	0.50
50:BW:93:GLU:OE1	50:BW:94:ALA:N	2.45	0.50
54:CA:1179:A:H5''	39:CL:102:LEU:O	2.11	0.50
54:CA:1189:C:H5''	33:CF:5:ILE:CG2	2.34	0.50
54:CA:1281:U:H5'	54:CA:1282:C:OP2	2.12	0.50
54:CA:1347:G:C8	39:CL:107:ARG:HB3	2.47	0.50
54:CA:152:A:H62	54:CA:169:C:H42	1.58	0.50
54:CA:152:A:N6	54:CA:169:C:H42	2.10	0.50
54:CA:1538:C:H2'	54:CA:1539:C:C6	2.46	0.50
54:CA:664:G:N2	54:CA:741:G:H1	2.07	0.50
54:CA:890:G:O2'	54:CA:906:G:N1	2.44	0.50
52:CB:11:C:H2'	52:CB:12:U:H6	1.76	0.50
52:CD:37:MIA:H3'	52:CD:38:A:C8	2.47	0.50
33:CF:6:HIS:HD2	33:CF:8:ILE:H	1.60	0.50
34:CG:61:LYS:HE2	34:CG:65:ARG:HD2	1.93	0.50
35:CH:140:ARG:HH11	35:CH:140:ARG:HB3	1.77	0.50
39:CL:47:LEU:N	39:CL:47:LEU:HD13	2.27	0.50
42:CO:41:ARG:HH11	42:CO:41:ARG:HG3	1.77	0.50
47:CT:67:LYS:CA	47:CT:70:ARG:HH12	2.08	0.50
13:D0:53:HIS:HA	13:D0:56:LYS:HB2	1.93	0.50
55:DA:1047:G:C2'	55:DA:1110:G:H22	2.23	0.50
55:DA:1289:C:H2'	55:DA:1290:C:H6	1.76	0.50
55:DA:1379:A:O2'	55:DA:1380:G:OP1	2.30	0.50
55:DA:1507:A:C3'	55:DA:1508:A:H5''	2.34	0.50
55:DA:1826:G:H2'	55:DA:1827:C:H6	1.76	0.50
55:DA:1887:C:C2'	55:DA:1888:G:H5''	2.41	0.50
55:DA:2035:G:H4'	55:DA:2036:C:OP2	2.12	0.50
55:DA:2119:A:N6	55:DA:2170:A:N7	2.58	0.50
55:DA:2317:C:O2'	55:DA:2318:G:H5'	2.11	0.50
55:DA:2468:G:O2'	55:DA:2469:A:O5'	2.30	0.50
55:DA:654(C):G:C2	55:DA:654(D):G:C4	2.99	0.50
2:DB:65:C:C2'	2:DB:66:A:H5'	2.41	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:DD:131:LEU:HB2	3:DD:136:ILE:CD1	2.42	0.50
3:DD:35:LYS:HD3	3:DD:63:ARG:C	2.32	0.50
7:DH:30:LYS:HB2	7:DH:79:VAL:O	2.12	0.50
56:DI:3:LEU:CD2	56:DI:4:ASP:H	2.23	0.50
57:DY:89:ALA:CA	56:DJ:15:ALA:CB	2.90	0.50
8:DK:18:VAL:O	8:DK:18:VAL:HG12	2.10	0.50
12:DP:20:ALA:CB	12:DP:99:PRO:HD2	2.42	0.50
18:DS:70:TYR:N	18:DS:70:TYR:HD2	2.07	0.50
21:DV:130:PRO:HA	21:DV:133:ILE:CD1	2.37	0.50
21:DV:178:GLU:HG3	21:DV:180:VAL:CA	2.41	0.50
57:DY:122:VAL:C	57:DY:126:ALA:H	2.14	0.50
57:DY:16:ASN:C	57:DY:16:ASN:OD1	2.49	0.50
57:DY:52:PHE:HD2	57:DY:52:PHE:H	1.56	0.50
1:AA:17:G:H4'	16:A1:25:TRP:CZ3	2.47	0.50
16:A1:90:VAL:HG13	17:A2:39:LEU:HB2	1.94	0.50
17:A2:32:THR:HG22	17:A2:58:VAL:CG1	2.42	0.50
26:A4:58:ARG:HD2	26:A4:58:ARG:C	2.32	0.50
30:A8:32:LEU:CD2	30:A8:33:ASN:N	2.75	0.50
1:AA:1050:A:H1'	1:AA:2751:G:H22	1.77	0.50
1:AA:1272:A:C5	1:AA:1618:A:H1'	2.47	0.50
1:AA:2182:G:H2'	1:AA:2183:C:C6	2.47	0.50
1:AA:2653:U:C2	1:AA:2654:A:N7	2.80	0.50
1:AA:2820:A:N3	13:A0:4:LEU:HD21	2.27	0.50
1:AA:532:A:O2'	1:AA:533:G:OP2	2.30	0.50
2:AB:44:G:H1'	2:AB:47:C:H42	1.76	0.50
2:AB:81:G:N2	2:AB:82:G:O6	2.44	0.50
3:AD:166:GLN:NE2	3:AD:166:GLN:HA	2.26	0.50
4:AE:24:THR:HG23	4:AE:186:GLY:HA2	1.94	0.50
4:AE:63:LEU:O	4:AE:63:LEU:HD23	2.12	0.50
5:AF:117:ARG:HG3	5:AF:122:LYS:HB2	1.92	0.50
6:AG:51:ARG:HB2	6:AG:51:ARG:CZ	2.41	0.50
15:AR:97:ALA:HB1	15:AR:98:LYS:NZ	2.26	0.50
24:AW:70:GLN:CG	24:AW:71:ASN:H	2.09	0.50
23:AZ:86:SER:O	23:AZ:87:PRO:C	2.49	0.50
53:B1:42:U:O2'	53:B1:44:U:OP2	2.27	0.50
31:BA:1118:C:O2'	31:BA:1119:C:H5'	2.12	0.50
31:BA:1178:G:N2	31:BA:1181:G:O6	2.44	0.50
31:BA:1541:U:O2	31:BA:1541:U:C2'	2.60	0.50
31:BA:713:G:H21	31:BA:777:A:C4'	2.25	0.50
31:BA:889:A:O3'	31:BA:890:G:C4'	2.56	0.50
52:BB:75:C:H3'	52:BB:75:C:C6	2.46	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:BC:18:G:N2	52:BC:58:A:O4'	2.45	0.50
32:BE:163:PHE:HA	32:BE:185:ILE:HG13	1.94	0.50
33:BF:41:GLY:C	33:BF:45:LYS:HE3	2.32	0.50
36:BI:36:ARG:HG2	36:BI:36:ARG:O	2.12	0.50
38:BK:89:PRO:HA	38:BK:92:ARG:NH1	2.24	0.50
40:BM:5:ARG:HG3	40:BM:71:LEU:HD11	1.94	0.50
41:BN:84:VAL:HG11	41:BN:95:ILE:HD11	1.93	0.50
45:BR:30:ALA:HA	45:BR:85:LEU:HD11	1.93	0.50
45:BR:32:LEU:O	45:BR:36:ILE:HG13	2.11	0.50
46:BS:54:GLU:O	46:BS:57:ARG:HB2	2.12	0.50
54:CA:1124:G:H8	54:CA:1124:G:OP2	1.95	0.50
54:CA:1128:C:N3	54:CA:1139:G:N1	2.60	0.50
54:CA:1240:U:OP2	37:CJ:116:ALA:HB2	2.11	0.50
54:CA:179:A:C2'	54:CA:180:U:H5'	2.42	0.50
54:CA:112:G:C4'	54:CA:389:A:H5''	2.41	0.50
54:CA:562:C:C2	42:CO:16:GLU:HB3	2.47	0.50
54:CA:60:A:O2'	54:CA:61:G:OP2	2.30	0.50
54:CA:127:G:OP1	54:CA:635:G:H1'	2.11	0.50
52:CB:18:G:O2'	52:CB:60:U:N3	2.45	0.50
52:CD:18:G:O2'	52:CD:19:G:P	2.70	0.50
33:CF:119:ARG:HG3	33:CF:119:ARG:HH11	1.77	0.50
36:CI:49:ALA:HB1	48:CU:80:PRO:HG3	1.93	0.50
37:CJ:54:THR:HG23	37:CJ:54:THR:O	2.12	0.50
39:CL:117:HIS:CD2	39:CL:123:PRO:HA	2.47	0.50
39:CL:47:LEU:C	39:CL:49:PRO:HD2	2.32	0.50
43:CP:115:LYS:O	43:CP:117:VAL:HG13	2.11	0.50
44:CQ:47:LEU:O	44:CQ:50:LYS:HG3	2.12	0.50
16:D1:92:ARG:CZ	16:D1:94:ASN:HD22	2.25	0.50
17:D2:99:ILE:HD13	17:D2:99:ILE:N	2.25	0.50
28:D6:20:ASN:CG	28:D6:21:TYR:N	2.61	0.50
28:D6:34:LEU:O	28:D6:36:LEU:HB3	2.12	0.50
28:D6:14:THR:HG22	28:D6:50:ARG:O	2.12	0.50
29:D7:20:ALA:HA	29:D7:23:ARG:CZ	2.42	0.50
55:DA:1062:G:N2	55:DA:1077:A:N7	2.56	0.50
55:DA:1155:A:O3'	16:D1:55:ARG:NH1	2.45	0.50
55:DA:990:A:N6	55:DA:1186:G:H1'	2.27	0.50
55:DA:695:G:OP1	55:DA:1380:G:O2'	2.29	0.50
55:DA:1799:G:N3	55:DA:1800:C:H5	2.10	0.50
55:DA:2059:A:O3'	5:DF:69:HIS:HA	2.12	0.50
55:DA:2302:G:C6	55:DA:2303:G:N7	2.79	0.50
55:DA:2377:A:O2'	55:DA:2378:A:H5'	2.11	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2657:A:C4	55:DA:2665:A:N6	2.79	0.50
55:DA:805:G:N2	55:DA:829:A:OP1	2.44	0.50
4:DE:35:GLN:CD	4:DE:37:ARG:HG2	2.33	0.50
4:DE:64:LYS:C	4:DE:66:HIS:N	2.64	0.50
4:DE:78:LEU:CD2	4:DE:79:ARG:HB2	2.32	0.50
5:DF:174:VAL:O	5:DF:174:VAL:CG2	2.60	0.50
5:DF:36:VAL:HG11	5:DF:183:VAL:CG1	2.41	0.50
6:DG:82:LEU:HA	6:DG:86:MET:SD	2.52	0.50
6:DG:95:ARG:O	6:DG:96:ARG:O	2.30	0.50
7:DH:7:LEU:N	7:DH:8:PRO:CD	2.74	0.50
58:DL:109:LYS:C	58:DL:111:LYS:H	2.13	0.50
10:DN:104:ARG:CZ	15:DR:34:VAL:HG11	2.42	0.50
20:DU:97:ARG:CD	20:DU:97:ARG:H	2.18	0.50
21:DV:174:VAL:C	21:DV:175:VAL:HG22	2.31	0.50
21:DV:194:PRO:HG2	21:DV:196:VAL:HG12	1.84	0.50
25:DX:38:GLU:C	25:DX:40:THR:H	2.15	0.50
57:DY:122:VAL:CB	57:DY:126:ALA:CB	2.90	0.50
57:DY:122:VAL:HB	57:DY:126:ALA:HB3	1.93	0.50
57:DY:73:GLY:O	57:DY:119:ALA:CB	2.59	0.50
16:A1:65:ILE:HD11	16:A1:96:ALA:HB3	1.92	0.50
16:A1:92:ARG:C	16:A1:94:ASN:N	2.64	0.50
26:A4:47:GLN:O	26:A4:48:ARG:C	2.51	0.50
30:A8:32:LEU:CB	30:A8:36:LYS:NZ	2.75	0.50
1:AA:200:U:H4'	23:AZ:34:THR:CG2	2.42	0.50
1:AA:2029:G:N1	1:AA:2033:A:OP1	2.45	0.50
1:AA:2111:C:O2	1:AA:2118:U:O2'	2.30	0.50
1:AA:954:G:H2'	1:AA:2274:A:C2	2.46	0.50
1:AA:2489:G:O2'	1:AA:2518:A:N6	2.40	0.50
1:AA:705:A:H62	1:AA:726:G:H1'	1.75	0.50
1:AA:704:G:HO2'	1:AA:705:A:P	2.35	0.50
1:AA:704:G:O2'	1:AA:705:A:P	2.70	0.50
4:AE:61:ARG:N	4:AE:62:PRO:HD2	2.27	0.50
7:AH:89:ILE:CD1	7:AH:90:LYS:H	2.25	0.50
8:AK:78:THR:HG23	8:AK:80:PRO:HD3	1.93	0.50
9:AM:120:LEU:HD21	9:AM:122:VAL:CG2	2.39	0.50
12:AP:33:GLY:HA2	12:AP:105:GLU:CA	2.41	0.50
14:AQ:42:ASP:C	14:AQ:44:LYS:H	2.14	0.50
20:AU:81:LYS:NZ	20:AU:97:ARG:NH2	2.59	0.50
21:AV:145:GLU:O	21:AV:145:GLU:OE1	2.30	0.50
23:AZ:96:LYS:HG3	23:AZ:97:LEU:H	1.77	0.50
31:BA:1001:G:H5'	31:BA:1001:G:H8	1.77	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:1037:C:H6	31:BA:1037:C:O5'	1.95	0.50
31:BA:1050:G:H2'	31:BA:1050:G:N3	2.27	0.50
31:BA:1270:C:O2'	31:BA:1271:G:H5'	2.11	0.50
31:BA:1305:G:O2'	31:BA:1306:A:O5'	2.29	0.50
31:BA:517:G:C6	31:BA:531:U:H1'	2.47	0.50
31:BA:894:G:O2'	31:BA:895:G:H5'	2.12	0.50
31:BA:992:U:H4'	31:BA:993:G:O5'	2.12	0.50
52:BB:19:G:O2'	52:BB:20:U:OP2	2.29	0.50
52:BC:65:G:H2'	52:BC:66:U:H6	1.76	0.50
31:BA:1206:G:O2'	33:BF:193:TYR:HA	2.12	0.50
34:BG:9:CYS:HA	34:BG:12:CYS:H	1.77	0.50
39:BL:17:VAL:HG22	39:BL:63:ILE:HG12	1.94	0.50
42:BO:127:GLU:O	42:BO:128:ALA:HB3	2.12	0.50
43:BP:84:ILE:HG22	43:BP:85:GLY:N	2.27	0.50
44:BQ:17:LYS:O	44:BQ:20:ALA:HB3	2.12	0.50
50:BW:87:LYS:O	50:BW:88:VAL:C	2.49	0.50
54:CA:1003:G:N2	54:CA:1004:A:O3'	2.40	0.50
54:CA:1326:C:O2'	54:CA:1327:C:H5'	2.12	0.50
54:CA:191(D):U:H2'	54:CA:191(E):G:C8	2.47	0.50
54:CA:243:A:O2'	54:CA:244:U:OP2	2.26	0.50
32:CE:93:VAL:HG11	32:CE:97:TRP:CD1	2.47	0.50
34:CG:150:GLU:H	34:CG:150:GLU:CD	2.15	0.50
35:CH:126:ARG:HG3	35:CH:126:ARG:NH1	2.26	0.50
35:CH:7:GLU:OE2	35:CH:7:GLU:HA	2.12	0.50
39:CL:5:TYR:HD2	39:CL:6:GLY:H	1.57	0.50
40:CM:22:LYS:HD2	40:CM:22:LYS:C	2.32	0.50
40:CM:39:PRO:CB	40:CM:70:ARG:HH12	2.20	0.50
40:CM:75:ILE:HG13	40:CM:76:ASN:N	2.23	0.50
42:CO:45:PRO:C	42:CO:46:LYS:O	2.48	0.50
45:CR:56:LEU:HA	45:CR:59:MET:CE	2.41	0.50
45:CR:76:GLU:C	45:CR:78:TYR:H	2.14	0.50
49:CV:41:VAL:HA	49:CV:44:MET:CG	2.41	0.50
13:D0:42:LYS:CA	13:D0:45:ARG:HH11	2.25	0.50
16:D1:106:PHE:O	16:D1:109:LEU:HB2	2.11	0.50
17:D2:39:LEU:HD13	17:D2:39:LEU:N	2.27	0.50
26:D4:56:VAL:HA	26:D4:60:GLN:HG2	1.94	0.50
26:D4:56:VAL:O	26:D4:57:GLU:C	2.50	0.50
27:D5:42:PRO:HB2	27:D5:43:HIS:HD2	1.77	0.50
28:D6:36:LEU:HD23	28:D6:36:LEU:C	2.32	0.50
55:DA:1680:U:O2	55:DA:1763:G:H3'	2.11	0.50
55:DA:2123:G:O2'	55:DA:2124:G:H5'	2.11	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2392:A:C2	55:DA:2424:C:N4	2.76	0.50
55:DA:2468:G:O2'	55:DA:2469:A:OP2	2.30	0.50
55:DA:2481:G:O2'	55:DA:2482:G:O5'	2.30	0.50
55:DA:2655:G:C2'	55:DA:2656:U:OP2	2.60	0.50
55:DA:579:G:H2'	55:DA:580:C:C6	2.46	0.50
55:DA:5:A:C2'	55:DA:6:A:H5'	2.42	0.50
55:DA:654(H):G:O5'	55:DA:654(H):G:H8	1.94	0.50
55:DA:654(L):G:C4	55:DA:654(M):C:C6	2.99	0.50
55:DA:814:C:H4'	55:DA:1225:C:O2	2.11	0.50
4:DE:34:VAL:HG23	4:DE:48:GLN:HB2	1.94	0.50
6:DG:115:ARG:CG	6:DG:115:ARG:HH11	2.25	0.50
6:DG:49:ASP:HB3	6:DG:52:ILE:HG12	1.94	0.50
7:DH:26:VAL:HG13	7:DH:27:LYS:N	2.26	0.50
7:DH:34:GLU:O	7:DH:36:PRO:HD3	2.12	0.50
56:DI:4:ASP:HA	56:DI:7:ARG:CD	2.37	0.50
8:DK:129:THR:HA	8:DK:137:PRO:HA	1.94	0.50
8:DK:33:ARG:HB3	8:DK:35:LEU:HG	1.94	0.50
8:DK:30:LEU:HB3	8:DK:36:ALA:HB3	1.92	0.50
10:DN:34:THR:O	10:DN:62:VAL:HB	2.11	0.50
12:DP:18:LYS:O	12:DP:19:GLY:O	2.30	0.50
15:DR:29:ARG:HH11	15:DR:29:ARG:HB2	1.77	0.50
20:DU:46:LYS:O	20:DU:48:ALA:N	2.45	0.50
21:DV:116:VAL:HG12	21:DV:118:GLN:CG	2.39	0.50
57:DY:64:LYS:O	57:DY:65:GLU:HB2	2.11	0.50
22:A3:45:PHE:CE2	22:A3:69:PHE:HE2	2.31	0.49
26:A4:50:VAL:O	26:A4:50:VAL:HG22	2.11	0.49
26:A4:51:ASP:O	26:A4:52:THR:C	2.50	0.49
26:A4:56:VAL:HG12	26:A4:57:GLU:N	2.27	0.49
1:AA:1110:G:O2'	1:AA:1111:A:O4'	2.30	0.49
1:AA:818:G:N1	1:AA:1188:U:OP2	2.32	0.49
1:AA:141(A):C:H6	1:AA:141(A):C:O5'	1.95	0.49
1:AA:1729:A:N6	1:AA:1731:G:N7	2.60	0.49
1:AA:2012:G:H8	1:AA:2012:G:O5'	1.95	0.49
1:AA:2210:G:O2'	1:AA:2211:G:OP1	2.24	0.49
1:AA:234:C:H2'	1:AA:235:U:C6	2.47	0.49
1:AA:2346:A:H5'	1:AA:2383:G:H1'	1.93	0.49
1:AA:2654:A:C4	1:AA:2656:U:O2	2.65	0.49
1:AA:288:C:O2	1:AA:288:C:H2'	2.13	0.49
1:AA:580:C:H2'	1:AA:581:C:C6	2.47	0.49
2:AB:66:A:C2'	2:AB:67:G:OP2	2.60	0.49
1:AA:919:G:H5''	2:AB:81:G:H1'	1.93	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:AD:24:ILE:HD11	3:AD:84:TYR:N	2.27	0.49
5:AF:6:VAL:O	5:AF:6:VAL:HG12	2.12	0.49
6:AG:8:LYS:O	6:AG:11:TYR:HB3	2.12	0.49
7:AH:94:TYR:N	7:AH:94:TYR:CD1	2.80	0.49
8:AK:5:LEU:CD1	8:AK:19:VAL:HG12	2.30	0.49
10:AN:8:LEU:N	10:AN:8:LEU:HD23	2.26	0.49
15:AR:16:ARG:HG3	15:AR:79:HIS:HA	1.94	0.49
19:AT:89:ILE:CG2	19:AT:92:LEU:HG	2.42	0.49
20:AU:17:SER:HB3	20:AU:71:LYS:HD2	1.93	0.49
21:AV:56:VAL:HG12	21:AV:57:ILE:N	2.26	0.49
31:BA:1064:G:HO2'	31:BA:1065:U:P	2.34	0.49
31:BA:1182:G:H4'	31:BA:1183:A:C5'	2.42	0.49
31:BA:321:A:H2'	31:BA:322:C:C6	2.47	0.49
31:BA:339:C:O2'	31:BA:340:U:H5'	2.12	0.49
31:BA:160:A:O2'	31:BA:344:A:N6	2.45	0.49
31:BA:450:G:H5'	46:BS:41:PRO:O	2.12	0.49
31:BA:490:G:OP2	34:BG:132:ARG:NH2	2.39	0.49
31:BA:704:A:N3	31:BA:704:A:H2'	2.27	0.49
31:BA:737:A:H1'	36:BI:73:ASN:ND2	2.27	0.49
32:BE:5:ILE:HD12	32:BE:59:GLU:CB	2.42	0.49
33:BF:21:ARG:O	33:BF:22:TRP:HB3	2.12	0.49
36:BI:8:ILE:HG21	36:BI:26:ILE:CD1	2.42	0.49
37:BJ:113:GLU:HB3	37:BJ:118:VAL:CG2	2.42	0.49
37:BJ:95:ARG:HG3	37:BJ:95:ARG:HH11	1.76	0.49
39:BL:10:ARG:HD3	39:BL:75:ASP:OD1	2.12	0.49
31:BA:942:G:N2	39:BL:124:GLN:NE2	2.58	0.49
41:BN:96:ARG:HA	41:BN:99:GLN:HG2	1.94	0.49
42:BO:33:ARG:HE	42:BO:33:ARG:HA	1.77	0.49
42:BO:55:VAL:HG12	42:BO:56:ALA:N	2.26	0.49
44:BQ:15:LYS:HA	44:BQ:15:LYS:HE2	1.94	0.49
44:BQ:39:LEU:HD13	44:BQ:47:LEU:HD12	1.94	0.49
54:CA:1003:G:C2	54:CA:1004:A:H4'	2.47	0.49
54:CA:1015:A:H1'	54:CA:1218:C:O2'	2.12	0.49
54:CA:1094:G:C2'	54:CA:1095:U:OP2	2.59	0.49
54:CA:136:C:H2'	54:CA:137:C:H6	1.77	0.49
54:CA:1540:U:N3	54:CA:1541:U:H1'	2.27	0.49
54:CA:464:G:O6	54:CA:466:C:H5''	2.12	0.49
54:CA:552:U:C2'	54:CA:553:A:H5'	2.41	0.49
54:CA:662:G:O2'	54:CA:836:G:C5'	2.59	0.49
54:CA:833:U:H2'	54:CA:834:C:H6	1.75	0.49
33:CF:70:VAL:HG21	33:CF:76:VAL:CG1	2.42	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:CG:108:LEU:HB3	34:CG:110:PHE:CD1	2.47	0.49
34:CG:47:ARG:NH2	53:C1:57:U:C5	2.63	0.49
35:CH:74:GLY:O	35:CH:115:VAL:HA	2.12	0.49
36:CI:19:LEU:HD23	36:CI:19:LEU:C	2.32	0.49
39:CL:79:LEU:HD13	39:CL:83:ARG:HD2	1.93	0.49
48:CU:33:ASP:O	48:CU:40:LEU:HD11	2.12	0.49
55:DA:2690:C:OP2	13:D0:14:SER:HB3	2.12	0.49
26:D4:68:ARG:CB	26:D4:68:ARG:CZ	2.90	0.49
27:D5:52:TYR:O	27:D5:53:ALA:HB2	2.12	0.49
55:DA:1062:G:H2'	55:DA:1077:A:N6	2.26	0.49
55:DA:1429:G:H2'	55:DA:1430:C:H6	1.73	0.49
55:DA:1493:C:H4'	55:DA:1494:A:OP1	2.12	0.49
55:DA:1608:A:C4'	55:DA:1609:A:OP1	2.60	0.49
55:DA:1630:G:H2'	55:DA:1630(A):C:C6	2.47	0.49
55:DA:1657:C:H2'	55:DA:1658:C:C6	2.47	0.49
55:DA:1694:C:H1'	55:DA:1695:G:C2	2.46	0.49
55:DA:189:G:N1	55:DA:205:G:O2'	2.38	0.49
55:DA:1926:U:O4'	55:DA:1929:G:O6	2.30	0.49
55:DA:191:A:H2'	55:DA:192:C:H6	1.77	0.49
55:DA:2227:A:H5'	3:DD:263:ARG:NH1	2.26	0.49
55:DA:2432:A:H2'	55:DA:2433:A:C8	2.46	0.49
55:DA:2439:A:H5'	55:DA:2439:A:H8	1.77	0.49
55:DA:2614:A:C4'	55:DA:2615:U:OP1	2.60	0.49
55:DA:2638:G:O2'	55:DA:2639:A:C8	2.65	0.49
55:DA:2712:U:O2'	55:DA:2712(A):A:OP1	2.30	0.49
55:DA:2526:G:H5'	55:DA:2742:C:O2'	2.12	0.49
55:DA:2740:A:C6	55:DA:2764:A:C8	3.00	0.49
55:DA:1:G:H2'	55:DA:2:G:H8	1.76	0.49
55:DA:528:A:C3'	55:DA:529:A:C5'	2.86	0.49
55:DA:528:A:H3'	55:DA:529:A:H5''	1.90	0.49
55:DA:536:A:P	16:D1:53:ARG:NH1	2.85	0.49
3:DD:44:ASN:C	3:DD:44:ASN:ND2	2.64	0.49
55:DA:1695:G:H1'	3:DD:8:PRO:O	2.12	0.49
4:DE:176:ILE:HG22	4:DE:179:GLU:H	1.76	0.49
4:DE:67:PHE:C	4:DE:69:LYS:H	2.09	0.49
6:DG:77:ILE:CG2	6:DG:77:ILE:O	2.60	0.49
9:DM:60:ILE:HD13	9:DM:60:ILE:H	1.77	0.49
12:DP:109:VAL:HG13	12:DP:110:THR:H	1.76	0.49
14:DQ:66:ALA:O	14:DQ:69:VAL:CG1	2.60	0.49
57:DY:40:LEU:O	57:DY:41:ARG:CB	2.59	0.49
57:DY:43:ALA:O	57:DY:44:LEU:HB2	2.12	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1046:A:O2'	57:DY:57:THR:HA	2.11	0.49
13:A0:101:ALA:O	13:A0:102:GLU:HB3	2.12	0.49
13:A0:13:HIS:O	13:A0:14:SER:C	2.51	0.49
16:A1:6:THR:O	16:A1:9:VAL:HG23	2.12	0.49
1:AA:996:A:O3'	16:A1:92:ARG:CD	2.59	0.49
22:A3:5:LYS:HE2	52:BC:73:A:C2'	2.42	0.49
1:AA:1379:A:O2'	1:AA:1380:G:OP1	2.29	0.49
1:AA:1810:A:C2'	1:AA:1811:G:H5'	2.41	0.49
1:AA:1893:C:C5	1:AA:1894:C:C5	3.00	0.49
1:AA:1935:G:H3'	1:AA:1962:C:N4	2.17	0.49
1:AA:1952:A:C5	10:AN:22:ILE:HD12	2.46	0.49
1:AA:1973:G:H2'	1:AA:1974:C:C6	2.47	0.49
1:AA:2249:U:H4'	1:AA:2275:C:C5	2.47	0.49
1:AA:523:C:C2'	1:AA:524:U:H5'	2.41	0.49
1:AA:655:A:C2'	1:AA:656:G:H5'	2.42	0.49
1:AA:666:G:H5''	1:AA:667:U:OP2	2.12	0.49
1:AA:859:G:HO2'	1:AA:860:U:H6	1.53	0.49
2:AB:110:G:C2'	2:AB:111:U:H5'	2.42	0.49
3:AD:262:ARG:NH1	3:AD:262:ARG:HG3	2.27	0.49
1:AA:779:U:OP1	3:AD:49:ILE:CG2	2.60	0.49
4:AE:72:VAL:O	4:AE:73:GLU:O	2.30	0.49
6:AG:88:ILE:O	6:AG:88:ILE:HG23	2.12	0.49
7:AH:130:ARG:NH1	7:AH:132:ARG:NH1	2.60	0.49
7:AH:19:VAL:HG12	7:AH:20:ALA:N	2.27	0.49
11:AO:101:VAL:HG23	11:AO:107:LYS:H	1.76	0.49
11:AO:92:GLU:HA	11:AO:123:LEU:CD1	2.42	0.49
12:AP:130:LYS:O	12:AP:130:LYS:HG2	2.11	0.49
14:AQ:3:ARG:CG	14:AQ:4:LEU:N	2.75	0.49
18:AS:79:GLY:O	18:AS:80:PRO:O	2.30	0.49
21:AV:74:VAL:HG13	21:AV:86:VAL:HG22	1.93	0.49
31:BA:1117:G:H5'	31:BA:1117:G:H8	1.77	0.49
31:BA:622:A:C8	31:BA:623:C:C6	3.00	0.49
31:BA:777:A:H2'	31:BA:778:G:C8	2.47	0.49
34:BG:126:ILE:HG22	34:BG:127:THR:N	2.27	0.49
34:BG:199:ASN:O	34:BG:200:GLU:CG	2.59	0.49
34:BG:200:GLU:HG3	34:BG:201:GLN:N	2.27	0.49
37:BJ:66:VAL:HG12	37:BJ:70:LYS:HE3	1.94	0.49
38:BK:6:ILE:H	38:BK:6:ILE:HD12	1.76	0.49
31:BA:1147:C:H2'	39:BL:16:ARG:HD3	1.94	0.49
39:BL:47:LEU:O	39:BL:50:LEU:HB2	2.12	0.49
43:BP:48:LEU:H	43:BP:48:LEU:CD2	2.24	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1069:C:C2'	54:CA:1070:U:O5'	2.61	0.49
54:CA:1128:C:C2	54:CA:1139:G:C6	3.00	0.49
54:CA:1399:C:C2	54:CA:1502:A:N6	2.80	0.49
54:CA:397:A:N7	54:CA:548:G:C8	2.80	0.49
54:CA:531:U:C4'	54:CA:532:A:OP1	2.60	0.49
54:CA:827:U:C5'	54:CA:828:A:OP2	2.59	0.49
54:CA:890:G:C2'	54:CA:891:U:OP2	2.60	0.49
54:CA:947:G:H2'	54:CA:948:C:H6	1.77	0.49
52:CD:12:U:H2'	52:CD:13:C:O4'	2.12	0.49
52:CD:15:G:H2'	52:CD:16:U:H5'	1.94	0.49
32:CE:20:GLU:HG2	32:CE:189:ASP:OD2	2.12	0.49
35:CH:32:VAL:HG12	35:CH:33:VAL:O	2.11	0.49
39:CL:4:TYR:CD1	39:CL:88:TYR:HB2	2.47	0.49
43:CP:78:ILE:O	43:CP:81:LEU:HB2	2.11	0.49
43:CP:7:VAL:CG2	6:DG:115:ARG:HH12	2.25	0.49
45:CR:65:ARG:NH1	45:CR:65:ARG:HB2	2.27	0.49
46:CS:43:LYS:HE2	46:CS:48:TRP:CZ3	2.47	0.49
49:CV:67:VAL:HG12	49:CV:67:VAL:O	2.12	0.49
13:D0:33:ARG:NH1	27:D5:57:VAL:HG23	2.27	0.49
17:D2:40:LEU:HD23	17:D2:47:VAL:HA	1.93	0.49
26:D4:56:VAL:HG13	26:D4:60:GLN:CG	2.41	0.49
55:DA:1045:A:O2'	55:DA:1047:G:C4	2.62	0.49
55:DA:191:A:H2'	55:DA:192:C:C6	2.46	0.49
55:DA:2821:A:H2'	55:DA:2822:G:O4'	2.12	0.49
3:DD:61:LEU:O	3:DD:63:ARG:NH1	2.45	0.49
4:DE:68:ALA:C	4:DE:69:LYS:HG3	2.32	0.49
6:DG:4:ASP:O	6:DG:5:VAL:HB	2.12	0.49
6:DG:73:ALA:O	6:DG:84:LYS:O	2.30	0.49
56:DI:24:ILE:O	56:DI:25:ASP:C	2.49	0.49
57:DY:139:VAL:CG2	56:DJ:6:GLU:CD	2.77	0.49
8:DK:92:VAL:O	8:DK:120:ILE:HG23	2.12	0.49
58:DL:135:GLY:C	58:DL:136:VAL:CG1	2.80	0.49
58:DL:132:ARG:O	58:DL:137:GLU:OE2	2.30	0.49
58:DL:63:ARG:CA	58:DL:63:ARG:NE	2.73	0.49
58:DL:63:ARG:HA	58:DL:63:ARG:NE	2.26	0.49
55:DA:2469:A:O2'	12:DP:56:ARG:HG2	2.11	0.49
14:DQ:5:THR:HG1	14:DQ:8:GLU:HG3	1.74	0.49
18:DS:4:LYS:CB	18:DS:106:ILE:HG22	2.42	0.49
21:DV:154:ASP:O	21:DV:155:LEU:C	2.50	0.49
57:DY:16:ASN:OD1	57:DY:25:PHE:CE2	2.65	0.49
57:DY:70:GLU:C	57:DY:113:GLN:CB	2.81	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:A0:81:ASP:O	13:A0:82:GLU:HB2	2.12	0.49
26:A4:9:LEU:HG	26:A4:25:TYR:C	2.32	0.49
28:A6:41:PRO:CD	28:A6:45:LYS:C	2.75	0.49
30:A8:46:ARG:NH1	30:A8:46:ARG:HB2	2.26	0.49
30:A8:48:PHE:O	30:A8:49:VAL:HG22	2.13	0.49
1:AA:1131:G:O2'	1:AA:1132:A:O5'	2.30	0.49
1:AA:1530:G:H2'	1:AA:1531:C:C6	2.48	0.49
1:AA:1656:C:H2'	1:AA:1657:C:H6	1.77	0.49
1:AA:1912:A:N6	1:AA:1918:A:H1'	2.27	0.49
1:AA:807:U:O2'	1:AA:2060:A:N1	2.39	0.49
1:AA:2656:U:O4	1:AA:2657:A:N7	2.46	0.49
1:AA:2660:A:H2'	1:AA:2661:G:O4'	2.12	0.49
1:AA:2715:C:O2'	1:AA:2716:U:H5'	2.12	0.49
1:AA:2737:G:O2'	1:AA:2738:A:H5'	2.11	0.49
1:AA:2897:U:H2'	1:AA:2898:U:C6	2.48	0.49
1:AA:566:U:H4'	1:AA:809:G:OP2	2.12	0.49
1:AA:654(S):G:O2'	1:AA:654(T):A:O4'	2.30	0.49
3:AD:174:ILE:N	3:AD:174:ILE:HD12	2.27	0.49
6:AG:154:GLY:O	6:AG:155:MET:HB3	2.13	0.49
6:AG:56:ALA:HB2	6:AG:153:ARG:NE	2.25	0.49
6:AG:83:ARG:O	6:AG:84:LYS:C	2.51	0.49
7:AH:39:PRO:O	7:AH:40:GLU:HG3	2.11	0.49
8:AK:2:LYS:HB2	8:AK:39:ALA:CB	2.42	0.49
11:AO:19:VAL:CG2	11:AO:21:ARG:HD2	2.43	0.49
14:AQ:74:ALA:O	14:AQ:75:GLU:C	2.51	0.49
15:AR:124:ASP:HB3	15:AR:128:GLU:OE2	2.13	0.49
15:AR:61:PHE:CE2	15:AR:76:PHE:HB2	2.47	0.49
21:AV:76:LEU:HA	21:AV:83:PRO:HA	1.95	0.49
24:AW:19:VAL:O	24:AW:23:LYS:HG3	2.12	0.49
31:BA:1399:C:C4'	31:BA:1400:C:O5'	2.54	0.49
31:BA:251:G:C4	31:BA:266:G:N7	2.80	0.49
31:BA:297:G:N2	31:BA:299:G:H3'	2.27	0.49
31:BA:58:C:O2'	31:BA:59:A:H5'	2.13	0.49
52:BB:37:MIA:H3'	52:BB:38:A:H8	1.76	0.49
52:BC:53:G:O2'	52:BC:54:U:H5'	2.12	0.49
52:BC:58:A:O2'	52:BC:59:U:O5'	2.24	0.49
32:BE:97:TRP:CD2	32:BE:101:MET:HG3	2.47	0.49
32:BE:228:GLY:O	32:BE:230:VAL:HG13	2.12	0.49
32:BE:32:ILE:HD12	32:BE:33:TYR:H	1.77	0.49
31:BA:1206:G:O4'	33:BF:194:GLY:N	2.45	0.49
34:BG:107:ARG:HH12	34:BG:114:ARG:NH2	2.10	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:BI:10:LEU:HD13	36:BI:61:LEU:HD13	1.93	0.49
39:BL:114:TYR:CD1	40:BM:60:ARG:HG2	2.47	0.49
39:BL:43:ALA:C	39:BL:45:ALA:H	2.15	0.49
40:BM:96:ILE:N	40:BM:96:ILE:HD13	2.16	0.49
44:BQ:21:TYR:CD2	44:BQ:22:THR:O	2.65	0.49
50:BW:67:ALA:HA	50:BW:73:HIS:N	2.27	0.49
53:C1:32:A:H2'	53:C1:33:G:C4'	2.41	0.49
54:CA:1002:G:N3	54:CA:1003:G:C8	2.80	0.49
54:CA:1064:G:H1'	54:CA:1066:C:C6	2.47	0.49
54:CA:1074:G:H4'	32:CE:104:ASN:HB2	1.93	0.49
54:CA:376:G:C5'	46:CS:5:ARG:HD2	2.41	0.49
54:CA:426:G:H2'	54:CA:427:U:C6	2.47	0.49
54:CA:484:G:O2'	54:CA:485:G:OP2	2.28	0.49
54:CA:848:C:O2'	54:CA:849:C:H5'	2.12	0.49
52:CC:51:U:H2'	52:CC:52:G:H8	1.76	0.49
32:CE:185:ILE:HA	32:CE:199:TYR:O	2.13	0.49
34:CG:196:LEU:O	34:CG:198:VAL:N	2.43	0.49
37:CJ:108:ALA:O	37:CJ:111:ARG:HG3	2.12	0.49
37:CJ:44:TYR:C	37:CJ:46:ALA:N	2.64	0.49
40:CM:27:ALA:CB	40:CM:34:VAL:HG21	2.42	0.49
54:CA:685:G:H5'	41:CN:39:PRO:O	2.11	0.49
43:CP:116:THR:C	43:CP:117:VAL:HG13	2.33	0.49
13:D0:41:ALA:O	13:D0:43:GLU:N	2.46	0.49
26:D4:14:ILE:HA	26:D4:31:ILE:O	2.12	0.49
30:D8:4:MET:O	30:D8:62:LEU:HD12	2.12	0.49
55:DA:1080:A:H2'	58:DL:126:MET:HE2	1.94	0.49
55:DA:1081:U:O2	58:DL:115:LEU:CD2	2.60	0.49
55:DA:1177:A:H5''	55:DA:1178:C:O5'	2.13	0.49
55:DA:1386:C:H2'	55:DA:1387:C:C6	2.47	0.49
55:DA:1854:A:H3'	55:DA:1855:G:H8	1.77	0.49
55:DA:2505:G:O6	55:DA:2576:G:H2'	2.12	0.49
55:DA:265:A:O2'	55:DA:266:G:O5'	2.30	0.49
55:DA:371:A:H1'	55:DA:373:U:C6	2.47	0.49
55:DA:699:A:H2'	55:DA:700:G:O4'	2.12	0.49
3:DD:35:LYS:HZ3	3:DD:104:TYR:HB2	1.75	0.49
3:DD:125:ILE:HD13	3:DD:131:LEU:HD21	1.95	0.49
3:DD:263:ARG:CB	3:DD:263:ARG:NH1	2.73	0.49
6:DG:61:ALA:HB2	6:DG:68:PRO:HD3	1.95	0.49
56:DI:27:LEU:O	56:DI:28:LYS:C	2.49	0.49
9:DM:60:ILE:HD13	9:DM:60:ILE:N	2.27	0.49
15:DR:39:ARG:HG2	15:DR:40:THR:N	2.14	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:DU:12:THR:OG1	20:DU:26:LYS:HE2	2.12	0.49
20:DU:14:LEU:HD23	20:DU:14:LEU:O	2.12	0.49
20:DU:38:ILE:CG2	20:DU:66:PRO:HG3	2.43	0.49
20:DU:38:ILE:HG22	20:DU:66:PRO:HA	1.94	0.49
21:DV:86:VAL:HG12	21:DV:87:ASP:N	2.27	0.49
57:DY:62:ALA:O	57:DY:63:LEU:O	2.29	0.49
57:DY:51:LEU:CG	57:DY:82:PHE:H	2.23	0.49
17:A2:2:PHE:HD2	17:A2:42:GLY:HA2	1.76	0.49
1:AA:1225:C:C3'	17:A2:85:LYS:HB2	2.42	0.49
28:A6:17:LYS:O	28:A6:18:ARG:HB3	2.12	0.49
1:AA:1113:U:H2'	1:AA:1114:G:C8	2.47	0.49
1:AA:1005:C:N1	1:AA:1143:A:C2	2.80	0.49
1:AA:2211:G:H2'	1:AA:2211:G:N3	2.28	0.49
1:AA:2645:G:H3'	1:AA:2646:C:H5'	1.93	0.49
1:AA:391:G:H2'	1:AA:392:C:C6	2.47	0.49
1:AA:479:A:H4'	1:AA:480:A:H5'	1.94	0.49
1:AA:507:A:O4'	1:AA:509:C:C2	2.66	0.49
1:AA:83:G:N2	1:AA:102:G:H2'	2.27	0.49
1:AA:953:A:C4	1:AA:954:G:C8	3.00	0.49
3:AD:211:ARG:O	3:AD:215:LEU:HG	2.12	0.49
1:AA:1798:U:OP2	3:AD:273:ARG:NH1	2.45	0.49
3:AD:70:TRP:HD1	3:AD:70:TRP:C	2.13	0.49
4:AE:73:GLU:HG3	4:AE:74:PRO:HD2	1.93	0.49
5:AF:34:TRP:CZ3	11:AO:8:PRO:HB3	2.47	0.49
5:AF:65:TRP:O	5:AF:67:GLN:N	2.45	0.49
10:AN:88:ASN:HD21	10:AN:90:GLN:CB	2.20	0.49
11:AO:47:ASP:HB3	11:AO:48:PRO:HA	1.93	0.49
12:AP:86:GLY:C	12:AP:88:GLY:N	2.61	0.49
14:AQ:29:PHE:CD2	14:AQ:30:ARG:N	2.81	0.49
21:AV:120:ILE:O	21:AV:121:HIS:HB2	2.11	0.49
21:AV:177:PRO:O	21:AV:178:GLU:OE1	2.30	0.49
31:BA:280:C:O2	31:BA:280:C:H2'	2.12	0.49
31:BA:328:C:HO2'	31:BA:329:A:P	2.35	0.49
52:BC:8:U:O2'	52:BC:48:C:H1'	2.12	0.49
52:BC:8:U:O4'	52:BC:48:C:O2'	2.30	0.49
35:BH:31:LEU:HD23	35:BH:32:VAL:N	2.27	0.49
31:BA:1298:C:H5	37:BJ:114:ARG:HD2	1.70	0.49
40:BM:31:GLY:O	40:BM:81:THR:HG21	2.13	0.49
41:BN:114:VAL:O	41:BN:114:VAL:HG13	2.11	0.49
42:BO:78:GLN:HG3	42:BO:79:GLU:N	2.28	0.49
43:BP:3:ARG:HA	43:BP:9:ILE:HG12	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:BQ:6:LEU:C	44:BQ:8:GLU:H	2.15	0.49
49:BV:51:VAL:HG21	49:BV:71:LEU:HB3	1.93	0.49
53:C1:42:U:O2'	53:C1:43:U:OP1	2.28	0.49
54:CA:1402:C:O2	54:CA:1500:A:N1	2.46	0.49
54:CA:450:G:N7	54:CA:481:G:C6	2.81	0.49
54:CA:801:U:H5'	54:CA:801:U:H6	1.77	0.49
52:CB:67:C:H2'	52:CB:68:C:C6	2.47	0.49
52:CB:9:A:H2	52:CB:11:C:N4	2.10	0.49
52:CC:40:C:O2	52:CC:40:C:H2'	2.13	0.49
34:CG:146:ILE:HD12	34:CG:146:ILE:H	1.76	0.49
37:CJ:16:LEU:CD2	39:CL:45:ALA:HB2	2.42	0.49
42:CO:83:VAL:CG2	42:CO:100:ILE:HG23	2.42	0.49
47:CT:63:ARG:HG2	47:CT:64:PRO:N	2.27	0.49
50:CW:39:LYS:O	50:CW:43:LEU:HG	2.13	0.49
17:D2:35:LEU:HD22	17:D2:57:VAL:O	2.12	0.49
55:DA:2108:C:O2'	55:DA:2109:U:H5'	2.12	0.49
55:DA:2163:C:OP1	55:DA:2172:U:H5	1.96	0.49
55:DA:2444:G:OP2	5:DF:68:LYS:HE3	2.13	0.49
55:DA:2618:G:H2'	55:DA:2619:C:H6	1.77	0.49
55:DA:2771:C:H5''	4:DE:202:LYS:HG2	1.95	0.49
55:DA:2817:G:N2	55:DA:2830:G:H1'	2.28	0.49
3:DD:58:HIS:CD2	3:DD:59:LYS:O	2.48	0.49
5:DF:39:TRP:CB	5:DF:101:LEU:HD22	2.43	0.49
5:DF:126:VAL:HG23	5:DF:127:GLU:N	2.27	0.49
6:DG:97:ASP:H	6:DG:100:TRP:HD1	1.59	0.49
7:DH:98:LEU:HB2	7:DH:125:VAL:CB	2.43	0.49
7:DH:126:PRO:HB2	7:DH:130:ARG:O	2.12	0.49
57:DY:135:ARG:NH2	56:DJ:18:LEU:HD13	2.24	0.49
56:DJ:2:ALA:O	56:DJ:6:GLU:HG3	2.12	0.49
8:DK:52:ARG:NH1	8:DK:52:ARG:HB3	2.27	0.49
58:DL:111:LYS:HG2	58:DL:111:LYS:O	2.11	0.49
58:DL:25:PRO:HA	58:DL:27:LEU:CD2	2.42	0.49
11:DO:148:LEU:O	11:DO:149:GLU:HG3	2.13	0.49
55:DA:806:C:OP2	11:DO:41:ARG:NH2	2.45	0.49
14:DQ:52:SER:O	14:DQ:56:LEU:CD2	2.60	0.49
18:DS:24:ILE:C	18:DS:24:ILE:HD12	2.32	0.49
21:DV:140:ASP:CG	21:DV:141:VAL:N	2.63	0.49
21:DV:146:ILE:HA	21:DV:174:VAL:CG1	2.42	0.49
21:DV:152:ALA:O	21:DV:154:ASP:C	2.51	0.49
25:DX:46:ASN:O	25:DX:50:VAL:HG22	2.12	0.49
23:DZ:63:ALA:O	23:DZ:65:SER:N	2.46	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:DZ:51:VAL:HG11	23:DZ:74:VAL:CG2	2.42	0.49
16:A1:102:GLU:O	16:A1:105:VAL:HG22	2.12	0.49
17:A2:40:LEU:HD23	17:A2:47:VAL:HA	1.95	0.49
17:A2:98:GLU:C	17:A2:99:ILE:HD12	2.33	0.49
29:A7:12:ARG:HD3	29:A7:46:VAL:CG2	2.31	0.49
1:AA:1131:G:O2'	1:AA:1132:A:O4'	2.28	0.49
1:AA:1157:G:H2'	1:AA:1158:C:H6	1.77	0.49
1:AA:1169:G:H2'	1:AA:1170:G:O4'	2.13	0.49
1:AA:1688:U:O2	1:AA:1700:A:H5'	2.12	0.49
1:AA:1995:U:H2'	1:AA:1996:C:C5	2.48	0.49
1:AA:2590:A:H2'	1:AA:2591:C:H6	1.77	0.49
1:AA:59:U:H6	1:AA:59:U:O5'	1.94	0.49
1:AA:654(T):A:H2'	1:AA:654(U):A:O4'	2.11	0.49
1:AA:829:A:C5	1:AA:2248:C:H5'	2.48	0.49
4:AE:36:ARG:HG2	4:AE:36:ARG:NH1	2.27	0.49
5:AF:84:VAL:C	5:AF:86:GLY:N	2.63	0.49
10:AN:54:GLU:C	10:AN:56:ASP:H	2.15	0.49
5:AF:34:TRP:HA	11:AO:6:LEU:HD12	1.95	0.49
15:AR:50:ILE:O	15:AR:99:LEU:HD12	2.13	0.49
21:AV:178:GLU:OE1	21:AV:178:GLU:O	2.30	0.49
24:AW:46:GLN:HA	24:AW:46:GLN:OE1	2.12	0.49
1:AA:849:A:N1	25:AX:25:ALA:HA	2.26	0.49
31:BA:1127:G:C2'	31:BA:1147:C:H42	2.26	0.49
31:BA:1201:A:O2'	31:BA:1202:G:OP2	2.26	0.49
31:BA:1399:C:H4'	31:BA:1400:C:C5'	2.41	0.49
31:BA:411:A:H62	31:BA:413:G:N2	2.10	0.49
31:BA:468:A:C2'	31:BA:474:G:H5'	2.42	0.49
31:BA:405:U:H5''	31:BA:495:A:H2	1.76	0.49
31:BA:652:U:C1'	31:BA:653:A:H2	2.17	0.49
31:BA:818:G:C3'	31:BA:819:A:C5'	2.90	0.49
52:BC:76:A:H8	52:BC:76:A:H5'	1.77	0.49
32:BE:54:THR:O	32:BE:57:PHE:HB3	2.12	0.49
34:BG:8:VAL:O	34:BG:11:LEU:HB2	2.12	0.49
35:BH:68:GLU:O	35:BH:68:GLU:HG3	2.12	0.49
36:BI:87:ARG:HG2	36:BI:87:ARG:NH1	2.27	0.49
37:BJ:105:VAL:O	37:BJ:108:ALA:HB3	2.13	0.49
37:BJ:50:ILE:O	37:BJ:54:THR:HG23	2.11	0.49
38:BK:91:ARG:HH11	38:BK:91:ARG:HG2	1.77	0.49
31:BA:973:G:H1'	40:BM:55:LYS:CG	2.42	0.49
42:BO:83:VAL:CG2	42:BO:100:ILE:HG23	2.43	0.49
31:BA:1329:A:P	43:BP:28:ALA:HB3	2.53	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
43:BP:34:LEU:HA	43:BP:37:THR:OG1	2.12	0.49
46:BS:52:ASP:OD2	46:BS:54:GLU:CG	2.61	0.49
47:BT:89:LEU:O	47:BT:92:ARG:HB3	2.13	0.49
26:A4:63:TYR:CD2	49:BV:41:VAL:HA	2.48	0.49
54:CA:1288:A:H2'	54:CA:1289:A:C8	2.48	0.49
54:CA:1453:G:O6	50:CW:51:GLU:HB3	2.13	0.49
54:CA:1535:C:H5'	54:CA:1535:C:C6	2.48	0.49
54:CA:397:A:N6	54:CA:548:G:C5	2.80	0.49
54:CA:865:A:C2	54:CA:918:A:H4'	2.47	0.49
54:CA:942:G:C2	54:CA:943:U:C6	3.00	0.49
52:CD:56:C:H2'	52:CD:57:G:H8	1.76	0.49
32:CE:216:SER:C	32:CE:218:ALA:H	2.16	0.49
33:CF:107:GLN:H	33:CF:107:GLN:CD	2.16	0.49
34:CG:147:ALA:HB1	34:CG:181:MET:O	2.12	0.49
35:CH:79:GLU:OE1	35:CH:79:GLU:O	2.30	0.49
39:CL:5:TYR:HD2	39:CL:17:VAL:O	1.95	0.49
41:CN:69:ALA:O	41:CN:70:LYS:C	2.50	0.49
43:CP:8:GLU:OE1	43:CP:67:GLU:HB2	2.12	0.49
54:CA:564:C:H5'	47:CT:32:TYR:CD2	2.47	0.49
48:CU:23:LYS:HG3	48:CU:24:ALA:N	2.28	0.49
49:CV:88:LYS:O	49:CV:89:ALA:O	2.30	0.49
50:CW:15:ARG:O	50:CW:19:SER:HB2	2.12	0.49
13:D0:57:ARG:HD2	13:D0:62:ALA:HB2	1.94	0.49
55:DA:1056:G:OP1	57:DY:35:LYS:CG	2.60	0.49
55:DA:1077:A:N1	55:DA:1088:A:N6	2.60	0.49
55:DA:1108:U:H2'	55:DA:1109:C:O4'	2.13	0.49
55:DA:1358:G:H2'	55:DA:1359:A:OP2	2.12	0.49
55:DA:1396:U:H2'	55:DA:1396:U:O2	2.12	0.49
55:DA:152:G:H2'	55:DA:153:C:C6	2.47	0.49
55:DA:1639:U:O2'	55:DA:1640:C:H5''	2.12	0.49
55:DA:1678:G:N2	55:DA:1989:G:N2	2.58	0.49
55:DA:1869:G:H5'	55:DA:1870:C:P	2.52	0.49
55:DA:2291:U:O2'	55:DA:2374:C:O2	2.25	0.49
55:DA:2472:G:N2	55:DA:2477:C:H5''	2.26	0.49
55:DA:312:G:H5'	55:DA:331:A:C2'	2.40	0.49
55:DA:602:G:O2'	55:DA:655:A:N6	2.45	0.49
55:DA:654(S):G:O2'	55:DA:654(T):A:O4'	2.30	0.49
55:DA:71:A:C2	19:DT:31:HIS:CE1	2.99	0.49
55:DA:828:U:C2'	55:DA:828:U:O2	2.60	0.49
4:DE:27:LEU:HD21	15:DR:1:MET:CE	2.42	0.49
5:DF:123:LEU:HD13	5:DF:192:LEU:HB3	1.92	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:DJ:7:ARG:HA	56:DJ:11:GLU:OE2	2.13	0.49
8:DK:29:TYR:HD1	8:DK:33:ARG:HE	1.60	0.49
58:DL:108:ALA:HA	58:DL:111:LYS:HZ2	1.76	0.49
58:DL:20:ALA:HB3	58:DL:21:PRO:HD3	1.95	0.49
58:DL:51:ALA:C	58:DL:52:ILE:CG1	2.78	0.49
58:DL:55:VAL:HG23	58:DL:69:THR:OG1	2.13	0.49
58:DL:72:PRO:HG2	58:DL:76:TYR:CE2	2.47	0.49
5:DF:34:TRP:HB2	11:DO:6:LEU:HD12	1.95	0.49
15:DR:136:GLN:HG3	15:DR:137:LYS:N	2.28	0.49
15:DR:32:TYR:O	15:DR:34:VAL:HG23	2.12	0.49
18:DS:42:ARG:HG2	18:DS:42:ARG:HH11	1.78	0.49
21:DV:116:VAL:O	21:DV:174:VAL:CA	2.48	0.49
24:DW:32:LEU:HD12	24:DW:57:ILE:HD12	1.93	0.49
16:A1:50:ARG:HH11	17:A2:72:VAL:HG11	1.76	0.49
16:A1:59:ARG:O	16:A1:63:VAL:HG23	2.13	0.49
17:A2:71:LEU:CA	17:A2:86:GLY:HA3	2.43	0.49
26:A4:14:ILE:HG13	26:A4:31:ILE:HB	1.94	0.49
11:AO:62:LEU:HD21	30:A8:25:MET:HB3	1.95	0.49
1:AA:1005:C:C6	1:AA:1143:A:N3	2.80	0.49
1:AA:1379:A:OP1	1:AA:1379:A:O4'	2.30	0.49
1:AA:142:G:H5''	1:AA:1598:C:O2'	2.12	0.49
1:AA:2639:A:H2'	1:AA:2640:G:H5'	1.95	0.49
1:AA:2713:A:C3'	1:AA:2714:G:H5''	2.42	0.49
1:AA:506:G:H4'	1:AA:507:A:O5'	2.12	0.49
1:AA:621:A:C2	1:AA:622:G:C4	3.00	0.49
1:AA:96:G:H4'	24:AW:48:HIS:ND1	2.28	0.49
1:AA:99:U:O2	1:AA:99:U:C2'	2.56	0.49
2:AB:82:G:N1	2:AB:95:U:O2	2.46	0.49
4:AE:14:ILE:O	4:AE:21:VAL:HG22	2.12	0.49
4:AE:65:GLY:O	4:AE:66:HIS:C	2.50	0.49
4:AE:72:VAL:O	4:AE:72:VAL:HG12	2.13	0.49
8:AK:78:THR:CG2	8:AK:104:GLN:HE22	2.22	0.49
8:AK:6:LEU:HD13	8:AK:36:ALA:CA	2.38	0.49
10:AN:4:PRO:O	10:AN:5:GLN:CB	2.43	0.49
1:AA:671:C:OP1	11:AO:42:SER:O	2.29	0.49
18:AS:59:VAL:CG1	18:AS:60:ASN:N	2.75	0.49
21:AV:9:TYR:HE2	21:AV:61:LEU:HD13	1.76	0.49
24:AW:38:GLN:O	24:AW:41:ILE:HG12	2.11	0.49
23:AZ:5:CYS:CB	23:AZ:8:SER:HG	2.25	0.49
23:AZ:70:VAL:O	23:AZ:74:VAL:HG23	2.12	0.49
53:B1:29:G:H2'	53:B1:30:C:O4'	2.12	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:1048:G:H2'	31:BA:1050:G:C8	2.48	0.49
31:BA:1305:G:O2'	31:BA:1306:A:H8	1.83	0.49
31:BA:48:C:H6	31:BA:365:U:O4	1.95	0.49
52:BC:66:U:H2'	52:BC:67:C:C6	2.48	0.49
32:BE:224:GLN:CA	32:BE:229:VAL:HG22	2.30	0.49
33:BF:117:ALA:HB2	33:BF:200:ALA:CB	2.42	0.49
34:BG:161:ASN:O	34:BG:164:ALA:N	2.45	0.49
34:BG:27:TYR:O	34:BG:28:SER:CB	2.61	0.49
47:BT:12:SER:HB3	47:BT:20:THR:HB	1.93	0.49
48:BU:62:GLU:HA	48:BU:65:ILE:HD11	1.95	0.49
48:BU:32:ARG:C	48:BU:69:THR:HG21	2.33	0.49
49:BV:58:VAL:O	49:BV:58:VAL:HG23	2.12	0.49
33:CF:162:GLN:CG	53:C1:54:U:O2	2.60	0.49
54:CA:1120:G:H2'	54:CA:1121:U:H6	1.77	0.49
54:CA:1157:A:N6	54:CA:1180:A:C5	2.81	0.49
54:CA:1206:G:C6	54:CA:1207:G:C5	3.01	0.49
54:CA:1210:C:H5'	54:CA:1214:C:N4	2.28	0.49
54:CA:1237:C:O2'	54:CA:1300:G:N2	2.44	0.49
54:CA:1358:U:OP2	54:CA:1359:C:N4	2.45	0.49
54:CA:485:G:O2'	54:CA:486:U:P	2.71	0.49
54:CA:518:C:H1'	54:CA:529:G:N1	2.27	0.49
54:CA:826:C:C2	54:CA:827:U:O2	2.66	0.49
54:CA:957:U:O2	54:CA:960:U:H6	1.94	0.49
52:CC:22:G:O2'	52:CC:23:A:H5'	2.13	0.49
32:CE:19:HIS:CD2	32:CE:206:ASP:HB2	2.48	0.49
35:CH:12:LEU:HB3	35:CH:31:LEU:HB3	1.94	0.49
37:CJ:15:ASP:HB3	37:CJ:19:GLY:H	1.75	0.49
40:CM:96:ILE:N	40:CM:96:ILE:HD13	2.27	0.49
42:CO:27:LEU:C	42:CO:29:GLY:N	2.66	0.49
42:CO:28:LYS:HE3	42:CO:33:ARG:HH12	1.77	0.49
42:CO:28:LYS:NZ	42:CO:33:ARG:NH2	2.55	0.49
43:CP:66:LEU:HD12	43:CP:66:LEU:N	2.26	0.49
43:CP:84:ILE:HD11	49:CV:66:MET:CB	2.43	0.49
45:CR:8:LYS:HB2	45:CR:8:LYS:HZ2	1.76	0.49
48:CU:31:LEU:CD2	48:CU:31:LEU:H	2.26	0.49
16:D1:25:TRP:O	16:D1:28:ARG:HB2	2.13	0.49
26:D4:14:ILE:O	26:D4:14:ILE:HG23	2.13	0.49
26:D4:22:ILE:O	26:D4:23:GLU:C	2.51	0.49
55:DA:1612:C:H4'	29:D7:5:TRP:O	2.13	0.49
55:DA:1092:C:H3'	55:DA:1092:C:C6	2.47	0.49
55:DA:1163:G:O2'	55:DA:1164:G:H5'	2.13	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1179:C:H2'	55:DA:1180:C:O4'	2.12	0.49
55:DA:1358:G:C2'	55:DA:1359:A:OP2	2.61	0.49
55:DA:1538:G:C2	55:DA:1539:G:C5	3.01	0.49
55:DA:171:G:O2'	55:DA:172:C:H5'	2.11	0.49
55:DA:2232:U:P	23:DZ:40:ARG:HH12	2.35	0.49
55:DA:2307:G:O4'	55:DA:2307:G:OP1	2.30	0.49
55:DA:310:A:O2'	55:DA:311:A:H2'	2.13	0.49
55:DA:479:A:C4'	55:DA:480:A:OP1	2.60	0.49
55:DA:654(S):G:C2'	55:DA:654(T):A:C8	2.95	0.49
3:DD:236:GLY:O	3:DD:237:GLU:OE2	2.30	0.49
3:DD:70:TRP:CD1	3:DD:70:TRP:C	2.86	0.49
7:DH:123:PHE:HB3	7:DH:125:VAL:HG22	1.95	0.49
8:DK:76:THR:CG2	8:DK:139:GLN:HE22	2.24	0.49
8:DK:57:ARG:O	8:DK:61:ARG:HG2	2.11	0.49
58:DL:132:ARG:O	58:DL:137:GLU:CD	2.51	0.49
58:DL:44:ALA:C	58:DL:46:ALA:N	2.59	0.49
58:DL:52:ILE:CD1	58:DL:53:VAL:H	2.21	0.49
58:DL:60:TYR:O	58:DL:61:ALA:HB2	2.11	0.49
9:DM:26:LEU:HG	9:DM:30:ILE:CD1	2.41	0.49
10:DN:71:ARG:NH2	10:DN:77:ILE:HG21	2.28	0.49
12:DP:110:THR:O	12:DP:113:GLN:N	2.46	0.49
19:DT:28:PHE:CE2	19:DT:92:LEU:HD11	2.48	0.49
57:DY:142:LEU:O	57:DY:143:GLN:O	2.31	0.49
57:DY:50:ARG:O	57:DY:83:TYR:N	2.45	0.49
57:DY:73:GLY:CA	57:DY:112:LEU:HG	2.42	0.49
57:DY:73:GLY:N	57:DY:118:THR:O	2.45	0.49
23:DZ:60:PHE:CE2	23:DZ:91:LYS:NZ	2.78	0.49
16:A1:98:LEU:O	16:A1:100:VAL:N	2.45	0.49
1:AA:1080:A:H2'	1:AA:1081:U:H6	1.77	0.49
1:AA:1493:C:H5'	1:AA:1494:A:OP2	2.12	0.49
1:AA:1790:C:H5''	1:AA:1791:A:OP1	2.12	0.49
1:AA:182:A:H2'	1:AA:183:C:O4'	2.12	0.49
1:AA:2130:U:C4'	1:AA:2134:A:H5'	2.42	0.49
1:AA:2522:U:H2'	1:AA:2523:G:C5'	2.43	0.49
3:AD:168:ARG:HA	3:AD:173:VAL:HA	1.95	0.49
1:AA:607:U:OP1	5:AF:103:LYS:HG3	2.11	0.49
5:AF:128:ALA:C	5:AF:142:TRP:HE1	2.16	0.49
1:AA:811:U:O4	11:AO:21:ARG:NH2	2.45	0.49
15:AR:12:SER:HB3	15:AR:15:VAL:CG1	2.41	0.49
1:AA:1615:C:H1'	18:AS:87:PRO:HG2	1.95	0.49
25:AX:4:LEU:HD23	25:AX:57:GLU:O	2.13	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:1049:U:O2'	31:BA:1050:G:OP2	2.28	0.49
31:BA:1213:A:N6	31:BA:1215:G:C2	2.81	0.49
31:BA:1275:A:O2'	31:BA:1276:G:H5'	2.12	0.49
31:BA:1298:C:O2'	31:BA:1299:A:C4	2.66	0.49
31:BA:397:A:N7	31:BA:547:A:O2'	2.45	0.49
31:BA:410:G:H4'	31:BA:411:A:OP1	2.12	0.49
32:BE:67:THR:HG21	32:BE:155:LEU:HG	1.93	0.49
35:BH:34:VAL:O	35:BH:42:GLY:N	2.44	0.49
37:BJ:131:LYS:O	37:BJ:131:LYS:HG2	2.13	0.49
31:BA:1118:C:P	39:BL:104:ARG:HD3	2.53	0.49
39:BL:3:GLN:NE2	39:BL:20:ARG:HH12	2.10	0.49
39:BL:7:THR:HG21	39:BL:9:ARG:NH2	2.27	0.49
43:BP:108:ARG:CZ	43:BP:114:ARG:HG2	2.43	0.49
49:BV:40:ILE:HD12	49:BV:71:LEU:HD23	1.94	0.49
50:BW:84:LEU:HD13	50:BW:84:LEU:C	2.32	0.49
54:CA:119:A:H5'	54:CA:120:A:C4	2.48	0.49
54:CA:1300:G:HO2'	54:CA:1301:U:P	2.34	0.49
54:CA:47:C:H6	54:CA:365:U:H2'	1.75	0.49
54:CA:424:G:H2'	54:CA:425:G:H8	1.77	0.49
54:CA:429:U:O2'	54:CA:430:A:H5''	2.12	0.49
54:CA:637:G:O2'	54:CA:638:G:H5'	2.13	0.49
54:CA:777:A:H2'	54:CA:778:G:H8	1.78	0.49
52:CC:18:G:C4	52:CC:58:A:C2	3.00	0.49
33:CF:79:ARG:CG	33:CF:79:ARG:HH11	2.25	0.49
35:CH:144:THR:O	35:CH:148:VAL:HG23	2.12	0.49
37:CJ:43:PHE:O	37:CJ:46:ALA:HB3	2.12	0.49
39:CL:53:VAL:O	39:CL:54:ASP:CB	2.60	0.49
40:CM:16:LEU:C	40:CM:16:LEU:HD13	2.33	0.49
40:CM:54:PHE:C	40:CM:55:LYS:HG3	2.33	0.49
54:CA:538:G:OP2	42:CO:115:LYS:CG	2.60	0.49
47:CT:45:HIS:O	47:CT:73:VAL:HG23	2.13	0.49
13:D0:96:ARG:HG2	13:D0:97:VAL:N	2.28	0.49
28:D6:20:ASN:ND2	28:D6:21:TYR:N	2.54	0.49
55:DA:2347:C:H4'	28:D6:39:TYR:HE2	1.78	0.49
55:DA:1058:U:H2'	55:DA:1059:G:H8	1.64	0.49
55:DA:1213:A:N3	55:DA:1238:G:H1'	2.27	0.49
55:DA:1946:U:H2'	55:DA:1947:C:C6	2.48	0.49
55:DA:1954:G:O2'	55:DA:1955:U:OP2	2.31	0.49
55:DA:2228:G:C5	55:DA:2229:C:C5	3.00	0.49
55:DA:921:G:H4'	55:DA:2269:A:C6	2.48	0.49
55:DA:2286:A:OP2	28:D6:28:ARG:CD	2.60	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2478:A:H3'	55:DA:2479:G:H8	1.78	0.49
55:DA:2638:G:HO2'	55:DA:2639:A:H8	1.57	0.49
55:DA:270(T):G:O2'	55:DA:270(U):C:H5'	2.11	0.49
55:DA:755:C:H2'	55:DA:756:C:C6	2.47	0.49
55:DA:852:G:O2'	55:DA:853:G:H5'	2.12	0.49
55:DA:855:G:H2'	55:DA:856:C:C6	2.47	0.49
2:DB:40:U:O2'	2:DB:41:U:OP1	2.25	0.49
4:DE:15:PHE:HA	4:DE:19:ARG:O	2.11	0.49
5:DF:183:VAL:O	5:DF:187:VAL:HG23	2.13	0.49
5:DF:42:ALA:O	5:DF:45:ARG:HB2	2.13	0.49
6:DG:120:LEU:O	6:DG:122:PRO:HD3	2.12	0.49
8:DK:111:PRO:O	8:DK:114:LEU:HB2	2.13	0.49
58:DL:135:GLY:C	58:DL:136:VAL:HG13	2.24	0.49
58:DL:60:TYR:HD2	58:DL:63:ARG:HB3	1.74	0.49
58:DL:69:THR:C	58:DL:70:LYS:CG	2.81	0.49
15:DR:24:PRO:HA	15:DR:49:VAL:CG1	2.33	0.49
19:DT:65:ARG:CD	19:DT:65:ARG:N	2.66	0.49
55:DA:498:G:N2	20:DU:47:LYS:NZ	2.61	0.49
20:DU:97:ARG:CD	20:DU:97:ARG:N	2.75	0.49
21:DV:33:LEU:HG	21:DV:34:ASN:N	2.28	0.49
57:DY:13:LEU:HD21	57:DY:62:ALA:O	2.11	0.49
57:DY:28:ASN:CG	57:DY:83:TYR:CD2	2.85	0.49
17:A2:1:MET:O	17:A2:2:PHE:C	2.51	0.49
17:A2:35:LEU:HD23	17:A2:37:VAL:CG1	2.43	0.49
16:A1:50:ARG:NH1	17:A2:72:VAL:CG1	2.74	0.49
30:A8:48:PHE:CD1	30:A8:48:PHE:N	2.80	0.49
1:AA:1069:A:H5'	1:AA:1070:A:H8	1.73	0.49
1:AA:323:G:O2'	1:AA:1205:U:O2	2.30	0.49
1:AA:1210:A:C4'	1:AA:1211:U:O5'	2.59	0.49
1:AA:1344:G:H4'	1:AA:1384:A:N7	2.28	0.49
1:AA:1654:A:C2	4:AE:113:PHE:CD2	3.01	0.49
1:AA:1923:U:H2'	1:AA:1924:C:H6	1.77	0.49
1:AA:2129:C:C2'	1:AA:2130:U:H5'	2.38	0.49
1:AA:908:C:OP1	12:AP:22:LYS:CG	2.61	0.49
1:AA:91:A:C2'	1:AA:92:G:H5'	2.42	0.49
2:AB:15:A:H1'	2:AB:109:G:C8	2.48	0.49
4:AE:22:PRO:O	4:AE:23:VAL:CG1	2.59	0.49
4:AE:54:GLN:O	4:AE:75:VAL:HG22	2.13	0.49
6:AG:96:ARG:O	6:AG:98:ARG:N	2.44	0.49
1:AA:1111:A:H4'	7:AH:3:ARG:HH11	1.78	0.49
10:AN:2:ILE:HD11	10:AN:82:ASN:HB3	1.93	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AO:115:LEU:HD23	11:AO:131:SER:HB2	1.94	0.49
11:AO:15:ARG:HG3	11:AO:16:ARG:N	2.28	0.49
20:AU:89:PHE:HD1	20:AU:90:LEU:CD2	2.16	0.49
21:AV:105:VAL:HG22	21:AV:106:GLY:N	2.27	0.49
21:AV:170:THR:O	21:AV:171:ILE:CB	2.60	0.49
31:BA:1065:U:O2'	31:BA:1066:C:OP2	2.30	0.49
31:BA:1136:U:H5''	31:BA:1137:C:C5	2.48	0.49
31:BA:132:C:H5'	31:BA:262:A:O2'	2.13	0.49
31:BA:1490:C:O2'	31:BA:1491:G:H5'	2.13	0.49
31:BA:831:U:O2'	31:BA:1539:C:OP1	2.31	0.49
31:BA:191(F):U:O2'	31:BA:191:G:H5'	2.12	0.49
31:BA:277:C:H5''	47:BT:68:ARG:HH22	1.78	0.49
31:BA:336:C:O2'	31:BA:337:C:H5'	2.13	0.49
31:BA:428:G:O2'	31:BA:429:U:P	2.70	0.49
31:BA:42:G:H2'	31:BA:43:C:O4'	2.13	0.49
52:BD:75:C:O2'	52:BD:76:A:H2	1.95	0.49
32:BE:102:LEU:HD23	32:BE:182:ILE:HD12	1.95	0.49
32:BE:19:HIS:O	32:BE:39:ILE:HG23	2.12	0.49
32:BE:68:ILE:N	32:BE:68:ILE:HD12	2.28	0.49
37:BJ:29:LYS:O	37:BJ:105:VAL:HG11	2.13	0.49
39:BL:26:VAL:HG13	39:BL:61:ALA:HB3	1.95	0.49
44:BQ:60:SER:O	44:BQ:61:TRP:HB3	2.13	0.49
47:BT:52:LYS:H	47:BT:52:LYS:HD2	1.78	0.49
47:BT:62:SER:HB2	47:BT:72:ARG:NH1	2.28	0.49
31:BA:1353:G:H5''	51:BX:13:ILE:HG21	1.95	0.49
54:CA:1097:C:H2'	54:CA:1098:C:C6	2.48	0.49
54:CA:1164:G:C6	54:CA:1173:G:C6	3.01	0.49
54:CA:1353:G:O2'	54:CA:1354:C:H5'	2.13	0.49
54:CA:265:G:H2'	54:CA:267:C:C5	2.48	0.49
54:CA:429:U:C1'	54:CA:430:A:H5''	2.37	0.49
54:CA:500:G:H2'	54:CA:501:C:H6	1.78	0.49
54:CA:625:G:H2'	54:CA:626:U:H6	1.77	0.49
54:CA:627:G:O2'	54:CA:628:G:H5'	2.13	0.49
54:CA:91:C:H2'	54:CA:92:G:H5'	1.94	0.49
32:CE:127:ILE:HD11	32:CE:139:LYS:HE2	1.95	0.49
33:CF:6:HIS:HB3	44:CQ:49:HIS:HD2	1.77	0.49
33:CF:95:THR:O	33:CF:97:LYS:N	2.41	0.49
34:CG:114:ARG:CG	34:CG:114:ARG:NH1	2.72	0.49
36:CI:21:LEU:O	36:CI:25:ILE:HG12	2.13	0.49
36:CI:74:ASP:O	36:CI:77:ARG:N	2.46	0.49
36:CI:9:VAL:HA	36:CI:59:TYR:O	2.12	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:CL:65:VAL:HG21	39:CL:73:GLN:NE2	2.27	0.49
39:CL:99:LEU:HD22	39:CL:99:LEU:N	2.28	0.49
43:CP:57:ARG:HH21	26:D4:34:GLU:CB	2.24	0.49
44:CQ:12:ARG:C	44:CQ:14:PRO:CD	2.75	0.49
44:CQ:24:CYS:CB	44:CQ:40:CYS:HB3	2.42	0.49
49:CV:47:HIS:O	49:CV:62:ILE:HB	2.13	0.49
49:CV:67:VAL:HB	26:D4:59:PHE:CD1	2.48	0.49
55:DA:1654:A:OP1	13:D0:1:MET:O	2.31	0.49
22:D3:32:ARG:O	22:D3:33:ALA:C	2.51	0.49
28:D6:15:GLU:CG	28:D6:16:CYS:H	1.94	0.49
29:D7:46:VAL:HG12	29:D7:47:ARG:H	1.78	0.49
55:DA:1083:U:C2'	55:DA:1085:A:OP2	2.49	0.49
55:DA:1178:C:H2'	55:DA:1179:C:C5	2.41	0.49
55:DA:1283:G:N2	55:DA:1286:A:OP2	2.45	0.49
55:DA:1657:C:H2'	55:DA:1658:C:H6	1.78	0.49
55:DA:1668:A:H61	55:DA:1676:A:H61	1.61	0.49
55:DA:2098:U:H2'	55:DA:2099:U:C6	2.48	0.49
55:DA:2173:A:C6	55:DA:2174:C:H1'	2.48	0.49
55:DA:2173:A:H5''	55:DA:2174:C:C5	2.48	0.49
55:DA:2305:A:H2'	55:DA:2306:C:O4'	2.12	0.49
55:DA:2355:C:O4'	22:D3:36:ILE:CD1	2.61	0.49
55:DA:2461:C:H2'	55:DA:2462:U:C6	2.47	0.49
55:DA:2475:C:H41	55:DA:2529:G:H22	1.61	0.49
55:DA:2593:U:C2	55:DA:2594:C:C5	3.01	0.49
55:DA:2835:A:H5'	55:DA:2836:U:OP1	2.13	0.49
55:DA:2846:G:P	15:DR:54:ARG:HB2	2.53	0.49
55:DA:639:U:C2	55:DA:640:C:C5	3.00	0.49
55:DA:860:U:C2	55:DA:2268:A:O4'	2.66	0.49
3:DD:134:ARG:HG2	3:DD:187:GLY:HA3	1.95	0.49
3:DD:43:ARG:HB2	3:DD:54:ARG:HB2	1.95	0.49
6:DG:137:GLU:CB	6:DG:152:LEU:HD22	2.43	0.49
7:DH:153:LYS:HG2	7:DH:162:ILE:CG1	2.41	0.49
56:DI:4:ASP:HA	56:DI:8:ILE:HG12	1.95	0.49
8:DK:60:GLU:CG	8:DK:61:ARG:HH12	2.22	0.49
58:DL:11:GLN:HB3	58:DL:41:PHE:CE1	2.46	0.49
58:DL:41:PHE:HE2	58:DL:45:THR:HG1	1.52	0.49
9:DM:38:HIS:N	9:DM:38:HIS:ND1	2.60	0.49
10:DN:3:GLN:O	10:DN:4:PRO:C	2.49	0.49
10:DN:86:ILE:HD12	10:DN:86:ILE:H	1.78	0.49
19:DT:51:VAL:HG13	19:DT:81:VAL:HG23	1.94	0.49
24:DW:11:GLU:HA	24:DW:14:ARG:HD2	1.95	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:DX:6:VAL:HG12	25:DX:56:VAL:HG13	1.93	0.49
57:DY:23:SER:CB	57:DY:68:LEU:O	2.58	0.49
1:AA:996:A:H4'	16:A1:92:ARG:HD2	1.95	0.49
17:A2:14:VAL:HA	17:A2:18:LEU:HD13	1.95	0.49
22:A3:36:ILE:CD1	22:A3:36:ILE:N	2.74	0.49
26:A4:63:TYR:HE2	49:BV:41:VAL:CG2	2.24	0.49
28:A6:18:ARG:NE	28:A6:43:CYS:HB3	2.27	0.49
28:A6:48:VAL:HG22	28:A6:49:HIS:N	2.27	0.49
1:AA:1009:A:OP2	1:AA:1010:A:OP2	2.30	0.49
1:AA:1049:C:C4	7:AH:2:SER:HB2	2.48	0.49
1:AA:1213:A:N3	1:AA:1238:G:H1'	2.27	0.49
1:AA:2033:A:O2'	1:AA:2034:U:P	2.71	0.49
1:AA:227:A:O2'	1:AA:228:A:OP2	2.29	0.49
1:AA:2391:G:C2'	1:AA:2424:C:H41	2.26	0.49
1:AA:2258:C:H2'	1:AA:2427:C:OP2	2.12	0.49
1:AA:634:C:H2'	1:AA:635:C:C6	2.48	0.49
1:AA:888:C:C2'	1:AA:889:C:OP2	2.60	0.49
1:AA:903:C:H2'	1:AA:904:C:H6	1.78	0.49
1:AA:915:C:O2'	1:AA:916:G:H5'	2.13	0.49
2:AB:12:C:C5'	2:AB:13:A:OP1	2.61	0.49
3:AD:201:HIS:O	3:AD:204:ILE:HG12	2.13	0.49
3:AD:2:ALA:O	3:AD:3:VAL:CB	2.59	0.49
8:AK:111:PRO:O	8:AK:113:ARG:HG2	2.12	0.49
8:AK:91:SER:OG	8:AK:119:PRO:HB2	2.13	0.49
10:AN:63:VAL:HG23	10:AN:64:ARG:HG3	1.93	0.49
12:AP:99:PRO:HG3	21:AV:79:ARG:NH1	2.27	0.49
15:AR:28:VAL:HG22	15:AR:29:ARG:N	2.27	0.49
15:AR:77:PRO:O	15:AR:79:HIS:N	2.46	0.49
21:AV:4:ARG:HH11	21:AV:58:VAL:HG11	1.77	0.49
25:AX:43:ILE:O	25:AX:47:VAL:HG23	2.13	0.49
31:BA:518:C:H2'	31:BA:530:G:N3	2.28	0.49
31:BA:517:G:H21	31:BA:530:G:P	2.36	0.49
31:BA:737:A:H1'	36:BI:73:ASN:HD21	1.78	0.49
31:BA:991:U:O2	31:BA:993:G:H1'	2.13	0.49
52:BD:44:G:H2'	52:BD:45:U:O4'	2.12	0.49
32:BE:17:PHE:CD1	32:BE:42:ILE:HG23	2.48	0.49
33:BF:156:ARG:NE	33:BF:160:ALA:O	2.42	0.49
35:BH:105:VAL:HB	35:BH:106:PRO:HD3	1.95	0.49
35:BH:149:GLU:O	35:BH:153:LYS:N	2.46	0.49
40:BM:35:SER:O	40:BM:36:GLY:O	2.31	0.49
41:BN:87:THR:O	41:BN:87:THR:HG22	2.13	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:BO:22:SER:O	42:BO:24:VAL:N	2.46	0.49
54:CA:1276:G:H2'	54:CA:1277:C:O4'	2.12	0.49
54:CA:1327:C:H2'	54:CA:1328:C:H6	1.74	0.49
54:CA:160:A:N6	54:CA:347:G:H1'	2.24	0.49
54:CA:88:C:H5''	54:CA:89:U:OP2	2.13	0.49
52:CB:29:G:O2'	52:CB:30:G:H5'	2.13	0.49
52:CB:66:U:H3'	52:CB:67:C:C6	2.48	0.49
52:CD:14:A:N6	52:CD:22:G:C5	2.80	0.49
33:CF:106:VAL:O	33:CF:107:GLN:C	2.51	0.49
17:D2:18:LEU:HD23	17:D2:19:LYS:N	2.28	0.49
55:DA:2355:C:H5'	22:D3:36:ILE:HD11	1.94	0.49
26:D4:69:LYS:CG	26:D4:70:GLY:N	2.76	0.49
55:DA:1092:C:H6	55:DA:1092:C:H5'	1.78	0.49
55:DA:1095:A:N3	55:DA:1095:A:C2'	2.75	0.49
55:DA:1098:A:H2'	55:DA:1099:G:H5''	1.95	0.49
55:DA:1357:U:H2'	55:DA:1358:G:O4'	2.13	0.49
55:DA:1464:C:O2'	55:DA:1528:A:H8	1.93	0.49
55:DA:1734:C:H3'	55:DA:1735:C:H5''	1.93	0.49
55:DA:1786:A:N1	55:DA:2606:C:H1'	2.27	0.49
55:DA:2531:A:N3	55:DA:2658:C:O2'	2.34	0.49
55:DA:508:G:N7	18:DS:80:PRO:HG3	2.28	0.49
55:DA:828:U:H3	55:DA:2247:A:C4'	2.25	0.49
3:DD:92:ILE:HD13	3:DD:104:TYR:CD2	2.48	0.49
6:DG:107:LEU:O	26:D4:38:LYS:HE2	2.13	0.49
6:DG:110:ALA:C	6:DG:112:PRO:HD2	2.32	0.49
7:DH:89:ILE:HA	7:DH:162:ILE:HA	1.94	0.49
8:DK:127:VAL:HA	8:DK:139:GLN:HA	1.95	0.49
8:DK:77:LEU:CD1	8:DK:78:THR:H	2.25	0.49
58:DL:98:ARG:H	58:DL:98:ARG:HH11	1.57	0.49
9:DM:34:LEU:O	9:DM:49:GLY:HA3	2.13	0.49
55:DA:2562:U:O2'	10:DN:23:ARG:NH1	2.46	0.49
12:DP:30:GLY:HA2	12:DP:107:ALA:CB	2.41	0.49
12:DP:52:VAL:O	12:DP:54:MET:N	2.46	0.49
18:DS:14:PRO:HG2	18:DS:78:GLU:HB2	1.95	0.49
18:DS:38:TYR:OH	27:D5:47:PRO:HG3	2.13	0.49
19:DT:8:ILE:N	19:DT:8:ILE:CD1	2.76	0.49
21:DV:192:ALA:HA	21:DV:193:GLU:OE1	2.12	0.49
21:DV:5:LEU:O	21:DV:5:LEU:HD22	2.13	0.49
22:A3:4:LYS:O	22:A3:5:LYS:O	2.30	0.49
26:A4:34:GLU:CB	43:BP:57:ARG:NH1	2.75	0.49
27:A5:16:ARG:HH11	27:A5:16:ARG:CG	2.18	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:747:U:C2	27:A5:2:ALA:N	2.80	0.49
29:A7:34:ARG:HB2	29:A7:42:LEU:HD22	1.94	0.49
1:AA:1140:C:C4'	1:AA:1143:A:H62	2.26	0.49
1:AA:116:C:C2'	1:AA:117:G:H5'	2.43	0.49
1:AA:116:C:O2'	1:AA:117:G:H5'	2.12	0.49
1:AA:1238:G:O2'	1:AA:1239:G:H5'	2.13	0.49
1:AA:1252:G:O4'	16:A1:33:ARG:CD	2.60	0.49
1:AA:1785:A:H2'	1:AA:1787:A:N7	2.28	0.49
1:AA:1853:A:H2'	1:AA:1854:A:C8	2.48	0.49
1:AA:195:A:C8	1:AA:197:A:OP1	2.66	0.49
1:AA:2126:A:H4'	1:AA:2127:G:O5'	2.13	0.49
1:AA:226:G:H1'	1:AA:228:A:N6	2.25	0.49
1:AA:2396:G:O2'	23:AZ:29:GLY:HA3	2.13	0.49
1:AA:2:G:H2'	1:AA:3:U:O4'	2.12	0.49
1:AA:374:A:H1'	1:AA:401:A:N6	2.27	0.49
1:AA:479:A:O2'	1:AA:480:A:O5'	2.29	0.49
2:AB:110:G:C2	2:AB:111:U:H1'	2.47	0.49
2:AB:28:C:O2'	2:AB:29:A:H5'	2.13	0.49
7:AH:125:VAL:HG22	7:AH:126:PRO:HA	1.94	0.49
8:AK:79:ILE:CA	8:AK:142:VAL:HG11	2.42	0.49
10:AN:34:THR:O	10:AN:37:ASP:HB2	2.12	0.49
19:AT:23:GLU:C	19:AT:25:LYS:H	2.16	0.49
19:AT:39:ILE:O	19:AT:40:LYS:C	2.51	0.49
19:AT:25:LYS:NZ	19:AT:82:GLN:OE1	2.39	0.49
20:AU:75:ILE:C	20:AU:75:ILE:HD13	2.33	0.49
21:AV:122:ARG:NH1	21:AV:122:ARG:HG2	2.28	0.49
25:AX:4:LEU:O	25:AX:36:VAL:HA	2.13	0.49
31:BA:1004:A:C2	31:BA:1005:A:C2	3.01	0.49
31:BA:1053:G:N7	31:BA:1200:C:H5"	2.27	0.49
31:BA:1288:A:H2'	31:BA:1289:A:C8	2.48	0.49
31:BA:1502:A:C2	31:BA:1505:G:N1	2.62	0.49
31:BA:437:U:H4'	34:BG:125:HIS:HE2	1.77	0.49
31:BA:547:A:H1'	31:BA:548:G:O4'	2.13	0.49
31:BA:658:G:H2'	31:BA:659:U:H6	1.78	0.49
31:BA:975:A:H5"	31:BA:976:G:O5'	2.13	0.49
52:BC:58:A:H1'	52:BC:60:U:C6	2.47	0.49
32:BE:97:TRP:CZ2	32:BE:101:MET:HB2	2.47	0.49
32:BE:24:TRP:H	32:BE:24:TRP:HD1	1.59	0.49
33:BF:22:TRP:HB2	33:BF:59:ARG:HB2	1.95	0.49
39:BL:82:ALA:CB	39:BL:96:LEU:HD11	2.42	0.49
40:BM:51:ARG:CB	40:BM:60:ARG:HA	2.42	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:BN:27:ASN:ND2	41:BN:29:ILE:HG22	2.28	0.49
42:BO:19:ARG:HG2	42:BO:20:LYS:N	2.28	0.49
43:BP:36:LYS:HB2	43:BP:59:TYR:CE2	2.48	0.49
43:BP:94:ARG:O	43:BP:94:ARG:HG3	2.13	0.49
49:BV:31:ILE:CG2	49:BV:49:ILE:HA	2.43	0.49
49:BV:49:ILE:HG22	49:BV:50:ALA:H	1.77	0.49
50:BW:50:GLU:CG	50:BW:51:GLU:N	2.76	0.49
53:C1:42:U:O2'	53:C1:43:U:P	2.71	0.49
54:CA:1003:G:H2'	54:CA:1004:A:C5'	2.16	0.49
54:CA:1139:G:C2	54:CA:1143:G:O6	2.66	0.49
54:CA:153:C:H2'	54:CA:154:C:H6	1.77	0.49
54:CA:258:G:C2	54:CA:259:G:C8	3.00	0.49
54:CA:255:G:O6	54:CA:266:G:O6	2.31	0.49
54:CA:369:C:OP2	54:CA:388:G:N2	2.46	0.49
54:CA:681:C:H2'	54:CA:682:G:H8	1.78	0.49
54:CA:992:U:H4'	54:CA:993:G:O5'	2.13	0.49
52:CC:35:A:C3'	52:CC:36:A:H5''	2.43	0.49
34:CG:24:GLU:O	34:CG:27:TYR:HB2	2.13	0.49
34:CG:3:ARG:O	34:CG:5:ILE:HG13	2.13	0.49
39:CL:118:LYS:HB3	39:CL:118:LYS:HZ3	1.77	0.49
54:CA:948:C:C6	43:CP:106:ASN:ND2	2.81	0.49
43:CP:120:LYS:O	43:CP:121:LYS:HB2	2.13	0.49
45:CR:3:ILE:O	45:CR:3:ILE:HG12	2.13	0.49
45:CR:26:GLU:CD	45:CR:77:ARG:HH12	2.16	0.49
46:CS:39:TYR:CZ	46:CS:41:PRO:HB3	2.48	0.49
17:D2:62:LEU:CD1	17:D2:95:LEU:HB2	2.42	0.49
26:D4:23:GLU:CD	26:D4:23:GLU:N	2.65	0.49
28:D6:40:CYS:HA	28:D6:46:HIS:HA	1.95	0.49
55:DA:686:G:N7	29:D7:5:TRP:CH2	2.81	0.49
55:DA:1059:G:C2	55:DA:1060:U:O4	2.65	0.49
55:DA:1639:U:C2'	55:DA:1640:C:H5''	2.43	0.49
55:DA:1681:G:O2'	55:DA:1762:A:C2'	2.61	0.49
55:DA:1853:A:N1	55:DA:2087:G:H1'	2.27	0.49
55:DA:2146:C:H4'	55:DA:2147:G:C8	2.48	0.49
55:DA:2430:A:H8	55:DA:2431:U:C5	2.30	0.49
55:DA:2661:G:H2'	55:DA:2662:A:O4'	2.13	0.49
55:DA:345:A:C4'	55:DA:346:A:OP1	2.59	0.49
55:DA:654(S):G:C3'	55:DA:654(T):A:C8	2.95	0.49
55:DA:898:C:C3'	55:DA:899:A:C5'	2.90	0.49
4:DE:49:LEU:O	4:DE:50:GLY:O	2.31	0.49
5:DF:125:LEU:HA	5:DF:194:MET:O	2.12	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:660:G:H5'	5:DF:99:TYR:CE2	2.48	0.49
7:DH:152:ARG:HE	7:DH:153:LYS:CE	2.26	0.49
8:DK:62:LYS:O	8:DK:66:GLU:HG2	2.13	0.49
58:DL:14:ALA:HB1	58:DL:50:ASP:HB3	1.92	0.49
58:DL:63:ARG:O	58:DL:64:SER:HB3	2.13	0.49
14:DQ:94:TYR:CE2	14:DQ:99:LYS:HG3	2.48	0.49
20:DU:54:LYS:O	20:DU:55:TYR:CB	2.60	0.49
20:DU:84:ARG:O	20:DU:95:LYS:HD3	2.13	0.49
24:DW:4:SER:OG	24:DW:5:GLU:OE2	2.26	0.49
23:DZ:78:LYS:CD	23:DZ:78:LYS:O	2.61	0.49
26:A4:71:ARG:NH1	26:A4:71:ARG:HB3	2.28	0.48
1:AA:686:G:O6	29:A7:12:ARG:HG3	2.13	0.48
1:AA:1173:G:H2'	1:AA:1174:A:OP2	2.13	0.48
1:AA:330:A:H2	1:AA:1210:A:H2'	1.77	0.48
1:AA:1372:U:H2'	1:AA:1373:A:H5'	1.95	0.48
1:AA:1971:A:C5	3:AD:241:PRO:HD3	2.48	0.48
1:AA:2115:G:H2'	1:AA:2116:G:C8	2.47	0.48
1:AA:2115:G:H2'	1:AA:2116:G:N7	2.28	0.48
1:AA:2521:C:H2'	1:AA:2522:U:O4'	2.12	0.48
1:AA:421:U:O2'	1:AA:422:A:P	2.71	0.48
1:AA:487:C:N4	1:AA:488:G:C6	2.81	0.48
1:AA:476:G:H4'	1:AA:502:A:N1	2.28	0.48
1:AA:610:C:H2'	1:AA:611:C:C6	2.47	0.48
1:AA:633:A:C2'	1:AA:634:C:H5'	2.42	0.48
1:AA:946:G:O6	1:AA:972:G:N2	2.46	0.48
1:AA:974:G:O2'	1:AA:975:G:N7	2.33	0.48
3:AD:268:ARG:HD3	3:AD:269:PHE:CE1	2.48	0.48
1:AA:1993:U:H4'	4:AE:128:SER:HB2	1.94	0.48
5:AF:60:SER:O	5:AF:61:GLY:O	2.31	0.48
8:AK:58:LEU:C	8:AK:60:GLU:H	2.15	0.48
9:AM:51:PHE:CE2	9:AM:119:ARG:HD2	2.48	0.48
10:AN:35:VAL:HG11	10:AN:103:ALA:CB	2.33	0.48
15:AR:31:SER:HB3	15:AR:42:ILE:CG2	2.42	0.48
15:AR:90:GLN:CD	15:AR:91:ARG:H	2.17	0.48
20:AU:21:LYS:O	20:AU:21:LYS:HG3	2.13	0.48
21:AV:178:GLU:OE1	21:AV:181:GLU:O	2.30	0.48
21:AV:39:VAL:CG2	21:AV:44:PHE:HB2	2.42	0.48
21:AV:7:ALA:C	21:AV:8:TYR:CD2	2.87	0.48
31:BA:526:C:OP2	42:BO:91:LYS:HE2	2.13	0.48
31:BA:556:C:O2'	31:BA:557:G:H5'	2.13	0.48
31:BA:748:C:O2'	31:BA:749:C:P	2.71	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:777:A:H2'	31:BA:778:G:H8	1.77	0.48
31:BA:577:G:H1'	31:BA:816:A:N3	2.27	0.48
33:BF:181:ASN:HD21	33:BF:204:LEU:HD12	1.78	0.48
33:BF:5:ILE:HD13	33:BF:5:ILE:O	2.13	0.48
35:BH:59:GLY:O	35:BH:60:TYR:C	2.51	0.48
31:BA:1241:G:OP2	37:BJ:38:LEU:HD21	2.13	0.48
40:BM:90:LEU:N	40:BM:91:PRO:CD	2.76	0.48
41:BN:95:ILE:CG2	41:BN:108:ILE:HD13	2.43	0.48
45:BR:17:ARG:HD3	45:BR:26:GLU:CD	2.34	0.48
45:BR:74:ASP:OD2	45:BR:76:GLU:HB3	2.13	0.48
54:CA:195:A:H1'	54:CA:222:U:O2'	2.13	0.48
54:CA:952:U:H4'	54:CA:964:A:N1	2.28	0.48
54:CA:973:G:H3'	54:CA:974:A:H5''	1.95	0.48
34:CG:103:ASN:O	34:CG:106:TYR:HB3	2.11	0.48
34:CG:29:PRO:O	34:CG:30:LYS:CB	2.61	0.48
35:CH:11:ILE:O	35:CH:12:LEU:CB	2.54	0.48
35:CH:78:HIS:CE1	35:CH:143:ARG:H	2.30	0.48
39:CL:70:LYS:O	39:CL:74:ILE:HG13	2.12	0.48
39:CL:79:LEU:HD13	39:CL:79:LEU:C	2.34	0.48
39:CL:83:ARG:C	39:CL:86:VAL:HG12	2.33	0.48
40:CM:101:VAL:HG22	40:CM:101:VAL:O	2.12	0.48
40:CM:5:ARG:NH2	40:CM:99:LYS:HD2	2.27	0.48
41:CN:81:ASP:O	41:CN:82:VAL:O	2.31	0.48
46:CS:1:MET:HG2	46:CS:2:VAL:N	2.28	0.48
47:CT:14:LYS:HZ2	47:CT:14:LYS:H	1.61	0.48
47:CT:27:PHE:CE1	47:CT:36:ILE:HD11	2.48	0.48
48:CU:31:LEU:H	48:CU:31:LEU:HD23	1.77	0.48
55:DA:2816:C:H5''	13:D0:99:LYS:NZ	2.27	0.48
28:D6:30:THR:HA	28:D6:31:PRO:O	2.12	0.48
30:D8:56:GLU:O	30:D8:57:ARG:C	2.51	0.48
55:DA:1056:G:O2'	55:DA:1086:A:H1'	2.13	0.48
55:DA:118:A:OP2	55:DA:119:A:H5''	2.13	0.48
55:DA:1537:C:C5	55:DA:1538:G:C6	3.01	0.48
55:DA:1844:C:H5''	3:DD:258:LYS:HG3	1.95	0.48
55:DA:1929:G:C5'	55:DA:1930:G:OP1	2.61	0.48
55:DA:1991:U:C2'	55:DA:1992:G:H5''	2.43	0.48
55:DA:2050:C:H2'	55:DA:2051:A:O4'	2.13	0.48
55:DA:2391:G:H2'	55:DA:2424:C:H41	1.77	0.48
55:DA:2543:G:H5'	55:DA:2543:G:H8	1.78	0.48
55:DA:2694:G:O2'	55:DA:2695:C:H5'	2.13	0.48
55:DA:2879:C:C4'	55:DA:2880:C:OP1	2.60	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:888:C:O5'	55:DA:889:C:H5	1.95	0.48
3:DD:13:ARG:CZ	3:DD:16:MET:CE	2.91	0.48
3:DD:30:GLU:HG3	3:DD:63:ARG:NH2	2.27	0.48
4:DE:39:PRO:HA	4:DE:44:TYR:N	2.28	0.48
4:DE:51:PHE:CD1	4:DE:52:LEU:N	2.81	0.48
5:DF:123:LEU:HD13	5:DF:192:LEU:HD13	1.95	0.48
5:DF:206:ILE:C	5:DF:206:ILE:HD12	2.34	0.48
6:DG:106:LEU:HA	6:DG:110:ALA:CB	2.43	0.48
6:DG:133:LEU:CD2	6:DG:157:ILE:HB	2.42	0.48
7:DH:149:ARG:HG3	7:DH:162:ILE:O	2.12	0.48
55:DA:1080:A:O2'	58:DL:126:MET:HE2	2.13	0.48
9:DM:120:LEU:HD11	9:DM:122:VAL:CG2	2.43	0.48
9:DM:71:ILE:H	9:DM:71:ILE:HD13	1.74	0.48
10:DN:77:ILE:HD12	10:DN:79:PHE:CE1	2.48	0.48
20:DU:44:ILE:HD12	20:DU:45:VAL:H	1.78	0.48
21:DV:114:GLY:O	21:DV:116:VAL:HG23	2.12	0.48
21:DV:28:MET:O	21:DV:34:ASN:HA	2.12	0.48
57:DY:15:GLU:HG3	57:DY:19:ARG:HH22	1.78	0.48
57:DY:97:ALA:O	57:DY:98:LYS:O	2.30	0.48
13:A0:21:TYR:OH	13:A0:43:GLU:HG2	2.13	0.48
16:A1:76:TYR:CZ	16:A1:80:ILE:HG13	2.48	0.48
22:A3:7:LEU:N	22:A3:7:LEU:HD13	2.28	0.48
26:A4:9:LEU:HA	26:A4:26:SER:O	2.13	0.48
29:A7:23:ARG:HH11	29:A7:23:ARG:HG3	1.78	0.48
1:AA:1022:G:O2'	1:AA:1024:G:N7	2.37	0.48
1:AA:1212:G:HO2'	1:AA:1213:A:P	2.36	0.48
1:AA:1340:U:O2'	1:AA:1341:U:OP1	2.30	0.48
1:AA:1362:C:O2'	1:AA:1363:C:H5'	2.13	0.48
1:AA:136:G:H2'	1:AA:137:C:H6	1.78	0.48
1:AA:142:G:H1'	19:AT:37:THR:CG2	2.35	0.48
1:AA:1858:G:C6	1:AA:1883:G:C6	3.01	0.48
1:AA:1991:U:H2'	1:AA:1992:G:C5'	2.43	0.48
1:AA:2056:G:C2	1:AA:2057:A:C8	3.01	0.48
1:AA:2162:G:H2'	1:AA:2163:C:H6	1.78	0.48
1:AA:2695:C:H2'	1:AA:2696:U:C6	2.49	0.48
1:AA:273(C):C:H3'	1:AA:273(D):C:H6	1.77	0.48
1:AA:2884:U:H2'	1:AA:2885:C:O4'	2.13	0.48
1:AA:311:A:O4'	1:AA:332:A:C4	2.67	0.48
1:AA:621:A:C2	1:AA:622:G:C8	3.01	0.48
1:AA:832:G:H2'	1:AA:833:U:C6	2.48	0.48
1:AA:886:C:C2	1:AA:890:A:N1	2.82	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:AD:35:LYS:NZ	3:AD:104:TYR:CB	2.65	0.48
3:AD:85:ASP:HB2	3:AD:92:ILE:CD1	2.44	0.48
4:AE:117:MET:CE	4:AE:124:GLY:HA3	2.43	0.48
4:AE:201:THR:CG2	4:AE:202:LYS:H	2.19	0.48
8:AK:47:LEU:HG	8:AK:51:ILE:CD1	2.43	0.48
8:AK:56:LYS:CG	8:AK:57:ARG:N	2.76	0.48
10:AN:23:ARG:HG3	10:AN:24:VAL:H	1.78	0.48
11:AO:110:TYR:O	11:AO:111:ARG:C	2.52	0.48
15:AR:42:ILE:HD12	15:AR:42:ILE:N	2.28	0.48
21:AV:11:GLU:OE2	21:AV:12:GLY:N	2.46	0.48
23:AZ:30:VAL:HG23	23:AZ:30:VAL:O	2.14	0.48
31:BA:1025:U:H2'	31:BA:1026:G:H8	1.78	0.48
15:AR:118:ARG:CZ	31:BA:1446:A:C6	2.96	0.48
31:BA:903:G:H2'	31:BA:904:C:C6	2.48	0.48
35:BH:9:LYS:HB3	35:BH:112:LEU:HD11	1.94	0.48
40:BM:4:ILE:HD12	40:BM:74:ILE:HG13	1.94	0.48
40:BM:5:ARG:CG	40:BM:71:LEU:HD11	2.43	0.48
41:BN:57:THR:HG22	41:BN:59:TYR:H	1.78	0.48
43:BP:15:VAL:O	43:BP:19:LEU:HD23	2.14	0.48
47:BT:8:GLY:CA	47:BT:23:VAL:HG22	2.43	0.48
47:BT:65:ILE:HG22	47:BT:65:ILE:O	2.12	0.48
54:CA:1132:C:H2'	54:CA:1133:G:C8	2.48	0.48
54:CA:1137:C:O2'	54:CA:1138:G:H5''	2.12	0.48
54:CA:11:G:H2'	54:CA:12:U:C6	2.48	0.48
54:CA:398:C:O2'	54:CA:399:G:H5'	2.13	0.48
54:CA:691:G:H2'	54:CA:692:U:C6	2.48	0.48
54:CA:84:U:O2	54:CA:84:U:H2'	2.12	0.48
54:CA:892:A:O2'	54:CA:1415:G:H4'	2.13	0.48
32:CE:77:ALA:HB2	32:CE:211:ILE:CD1	2.42	0.48
33:CF:61:ALA:O	33:CF:62:ASP:C	2.51	0.48
34:CG:144:ASP:C	34:CG:146:ILE:HD12	2.34	0.48
54:CA:543:C:P	34:CG:14:ARG:HH21	2.37	0.48
34:CG:7:PRO:HB2	34:CG:10:ARG:HD2	1.94	0.48
35:CH:42:GLY:HA2	35:CH:65:ASN:O	2.12	0.48
38:CK:44:PHE:HA	38:CK:79:VAL:CG1	2.43	0.48
54:CA:718:G:H1'	41:CN:116:HIS:HA	1.95	0.48
42:CO:89:ARG:NE	42:CO:91:LYS:HZ3	2.10	0.48
47:CT:52:LYS:HD2	47:CT:52:LYS:H	1.78	0.48
54:CA:1014:A:H4'	49:CV:14:HIS:ND1	2.28	0.48
17:D2:1:MET:HG2	17:D2:42:GLY:H	1.78	0.48
22:D3:25:ARG:HD3	22:D3:29:GLN:HE22	1.78	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:D6:34:LEU:O	28:D6:35:GLU:C	2.51	0.48
55:DA:1020:A:N6	55:DA:1141:U:O2'	2.46	0.48
55:DA:1142(A):A:C5	55:DA:1144:G:C5	3.01	0.48
55:DA:1322:A:OP1	18:DS:11:ARG:HG3	2.13	0.48
55:DA:1336:A:H2'	55:DA:1337:G:H8	1.76	0.48
55:DA:1827:C:O2	55:DA:1827:C:H2'	2.13	0.48
55:DA:1885:A:H2'	55:DA:1886:C:O4'	2.12	0.48
55:DA:1889:A:O2'	55:DA:2087:G:H5'	2.14	0.48
55:DA:1899:G:HO2'	55:DA:1900:A:P	2.35	0.48
55:DA:1946:U:C2	55:DA:1947:C:C5	3.01	0.48
55:DA:532:A:N7	55:DA:2021:C:H2'	2.28	0.48
55:DA:2127:G:H21	55:DA:2173:A:C1'	2.27	0.48
55:DA:2210:G:H4'	55:DA:2211:G:OP1	2.13	0.48
55:DA:526:A:H2	55:DA:2625:G:N3	2.11	0.48
55:DA:2839:G:H5'	13:D0:46:GLY:HA2	1.95	0.48
55:DA:301:G:HO2'	55:DA:302:C:H6	1.57	0.48
55:DA:554:U:O2'	55:DA:556:G:C8	2.65	0.48
55:DA:821:A:H5''	55:DA:822:U:H6	1.78	0.48
55:DA:950:G:H2'	55:DA:951:C:H6	1.77	0.48
8:DK:79:ILE:O	8:DK:79:ILE:HG22	2.12	0.48
10:DN:71:ARG:HG3	10:DN:71:ARG:NH1	2.25	0.48
21:DV:158:PRO:C	21:DV:160:GLY:H	2.15	0.48
21:DV:174:VAL:C	21:DV:175:VAL:HG13	2.34	0.48
21:DV:190:GLU:O	21:DV:191:VAL:HB	2.13	0.48
25:DX:18:ASP:N	25:DX:18:ASP:OD1	2.46	0.48
55:DA:1082:U:O2	57:DY:41:ARG:NH2	2.46	0.48
23:DZ:63:ALA:O	23:DZ:64:ALA:C	2.51	0.48
16:A1:90:VAL:HA	17:A2:39:LEU:HD23	1.93	0.48
17:A2:57:VAL:CG1	17:A2:99:ILE:HG13	2.43	0.48
1:AA:2346:A:H61	28:A6:28:ARG:NH2	2.11	0.48
28:A6:35:GLU:O	28:A6:36:LEU:CB	2.61	0.48
1:AA:1175:U:H2'	1:AA:1176:G:C4'	2.41	0.48
1:AA:1301:A:O2'	1:AA:1302:A:H3'	2.13	0.48
1:AA:1541:U:H2'	1:AA:1542:G:O4'	2.13	0.48
1:AA:1667:G:OP2	1:AA:1667:G:O4'	2.30	0.48
1:AA:1815:A:P	3:AD:54:ARG:HH22	2.36	0.48
1:AA:1842:G:H2'	1:AA:1843:C:C6	2.47	0.48
1:AA:1888:G:H5'	1:AA:1888:G:N3	2.28	0.48
1:AA:1930:G:H2'	1:AA:1968:G:H1	1.78	0.48
1:AA:2119:A:N6	1:AA:2171:A:N3	2.62	0.48
1:AA:2366:A:H2'	1:AA:2367:G:O4'	2.12	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2387:U:C5'	1:AA:2388:A:OP2	2.62	0.48
1:AA:270(N):G:H1'	1:AA:270(P):C:H1'	1.96	0.48
1:AA:270(V):G:O2'	1:AA:270(W):G:H5'	2.13	0.48
1:AA:2713:A:H3'	1:AA:2714:G:C5'	2.43	0.48
1:AA:2864:G:O2'	1:AA:2865:U:H5'	2.12	0.48
1:AA:616:A:H2'	1:AA:617:G:O4'	2.14	0.48
2:AB:12:C:H4'	2:AB:13:A:C5'	2.43	0.48
2:AB:55:U:HO2'	6:AG:29:TRP:HD1	1.60	0.48
3:AD:109:ASP:HB2	3:AD:197:GLY:CA	2.43	0.48
4:AE:37:ARG:CG	4:AE:46:ALA:HB3	2.32	0.48
5:AF:141:ALA:O	5:AF:144:LYS:HB3	2.13	0.48
5:AF:63:LYS:HZ3	5:AF:67:GLN:HB2	1.78	0.48
5:AF:8:GLN:HG2	5:AF:126:VAL:HG12	1.95	0.48
12:AP:30:GLY:N	12:AP:65:PHE:HE2	2.11	0.48
12:AP:35:VAL:HG22	12:AP:36:ALA:N	2.28	0.48
15:AR:23:ARG:HG2	15:AR:120:ARG:NH1	2.29	0.48
15:AR:29:ARG:NE	15:AR:44:ASP:HB3	2.28	0.48
21:AV:144:LEU:HD12	21:AV:144:LEU:C	2.33	0.48
31:BA:1027:C:HO2'	31:BA:1028:C:P	2.36	0.48
31:BA:1054:C:H2'	31:BA:1055:A:H5''	1.95	0.48
31:BA:1213:A:C6	31:BA:1215:G:C4	3.01	0.48
31:BA:1301:U:C4	31:BA:1303:C:C6	3.01	0.48
31:BA:1372:U:H2'	31:BA:1373:G:C5'	2.42	0.48
31:BA:1448:C:H2'	31:BA:1449:C:O4'	2.12	0.48
31:BA:176:C:H2'	31:BA:177:C:H6	1.78	0.48
31:BA:357:G:C2	31:BA:358:U:C5	3.02	0.48
31:BA:575:G:O2'	31:BA:576:G:P	2.71	0.48
31:BA:834:C:H2'	31:BA:835:U:H6	1.78	0.48
31:BA:977:A:N3	31:BA:977:A:H3'	2.27	0.48
52:BB:57:G:H2'	52:BB:58:A:H5''	1.94	0.48
32:BE:161:ALA:HA	32:BE:183:PRO:O	2.13	0.48
32:BE:212:GLN:HE22	32:BE:216:SER:HB2	1.78	0.48
33:BF:64:VAL:CG2	33:BF:97:LYS:HE3	2.43	0.48
35:BH:146:ALA:O	35:BH:148:VAL:N	2.46	0.48
35:BH:83:GLU:HG2	35:BH:88:LYS:HD2	1.94	0.48
42:BO:60:LEU:CD2	42:BO:60:LEU:N	2.76	0.48
54:CA:1004:A:C2'	54:CA:1005:A:O4'	2.61	0.48
54:CA:1239:A:H1'	54:CA:1241:G:C5	2.48	0.48
54:CA:1336:C:H1'	54:CA:1337:G:C2	2.47	0.48
54:CA:1542:U:OP1	54:CA:1542:U:H4'	2.11	0.48
54:CA:481:G:O2'	54:CA:482:A:P	2.70	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:485:G:C2'	54:CA:486:U:OP2	2.61	0.48
54:CA:502:G:H2'	54:CA:503:C:H6	1.79	0.48
54:CA:509:A:C6	54:CA:510:A:N1	2.81	0.48
54:CA:50:A:O2'	54:CA:51:A:OP2	2.31	0.48
54:CA:64:G:N2	54:CA:67:C:C4	2.81	0.48
54:CA:676:A:O2'	54:CA:677:U:H5'	2.13	0.48
54:CA:817:C:H4'	54:CA:818:G:O5'	2.12	0.48
54:CA:982:U:C4'	54:CA:983:A:O5'	2.50	0.48
32:CE:101:MET:C	32:CE:102:LEU:HD12	2.33	0.48
32:CE:11:LEU:HG	32:CE:213:LEU:HD11	1.94	0.48
33:CF:33:LEU:O	33:CF:37:GLN:HG2	2.13	0.48
37:CJ:33:ASP:C	37:CJ:35:LYS:H	2.16	0.48
37:CJ:78:ARG:HG3	37:CJ:79:ARG:H	1.77	0.48
39:CL:111:ARG:O	39:CL:113:LYS:HE3	2.14	0.48
39:CL:127:LYS:CE	52:CC:34:G:OP2	2.61	0.48
43:CP:20:THR:C	43:CP:22:ILE:H	2.17	0.48
43:CP:19:LEU:HB3	43:CP:25:ILE:HG21	1.94	0.48
54:CA:254:G:H21	47:CT:16:GLN:NE2	2.12	0.48
49:CV:44:MET:C	49:CV:62:ILE:HG21	2.34	0.48
49:CV:64:GLU:HG3	49:CV:65:ASN:N	2.27	0.48
50:CW:71:THR:CG2	50:CW:72:LEU:H	1.93	0.48
13:D0:53:HIS:CD2	13:D0:56:LYS:HE3	2.48	0.48
17:D2:35:LEU:CD2	17:D2:57:VAL:HG22	2.28	0.48
6:DG:108:ASN:O	26:D4:38:LYS:HB2	2.13	0.48
27:D5:49:CYS:HA	27:D5:58:LEU:CB	2.36	0.48
29:D7:8:ASN:ND2	29:D7:10:ARG:N	2.61	0.48
55:DA:1682:G:H5'	55:DA:1762:A:O2'	2.13	0.48
55:DA:1954:G:O2'	55:DA:1956:U:C5	2.66	0.48
55:DA:1970:A:C5'	55:DA:1971:A:OP1	2.61	0.48
55:DA:222:A:O2'	55:DA:223:A:O5'	2.25	0.48
55:DA:2455:G:H2'	55:DA:2456:C:C6	2.48	0.48
55:DA:2537:U:H2'	55:DA:2538:C:C6	2.48	0.48
55:DA:12:U:O2	55:DA:2627:G:OP1	2.31	0.48
55:DA:2848:G:O2'	55:DA:2849:U:P	2.71	0.48
55:DA:287:C:H2'	55:DA:288:C:H6	1.78	0.48
55:DA:447:A:H2'	55:DA:473:G:N7	2.29	0.48
55:DA:465:G:N2	55:DA:466:A:C2	2.81	0.48
55:DA:646:A:H5'	55:DA:646:A:N3	2.27	0.48
55:DA:792:G:C5'	55:DA:793:A:H5'	2.30	0.48
4:DE:196:VAL:O	4:DE:197:ILE:HG22	2.12	0.48
56:DJ:18:LEU:C	56:DJ:21:LYS:HB2	2.33	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:DK:124:GLY:H	8:DK:142:VAL:CG2	2.27	0.48
8:DK:32:PRO:C	8:DK:34:GLY:N	2.65	0.48
8:DK:61:ARG:NH2	8:DK:64:GLU:OE2	2.46	0.48
58:DL:101:TRP:CE2	58:DL:140:GLY:HA3	2.48	0.48
58:DL:18:THR:HG22	58:DL:38:VAL:HG13	1.86	0.48
58:DL:25:PRO:CB	58:DL:27:LEU:HG	2.43	0.48
9:DM:43:THR:CG2	9:DM:45:ASN:HD21	2.26	0.48
9:DM:43:THR:HG22	9:DM:45:ASN:HD22	1.75	0.48
9:DM:75:TYR:C	9:DM:76:SER:O	2.48	0.48
9:DM:78:TYR:H	9:DM:78:TYR:HD1	1.60	0.48
11:DO:41:ARG:HH21	11:DO:41:ARG:CG	2.18	0.48
21:DV:191:VAL:CB	21:DV:197:ILE:HG12	2.43	0.48
21:DV:51:ALA:CB	21:DV:57:ILE:HD11	2.43	0.48
57:DY:117:LEU:HD13	57:DY:117:LEU:N	2.27	0.48
57:DY:90:ALA:HB3	56:DJ:15:ALA:H	1.78	0.48
16:A1:50:ARG:HH21	16:A1:50:ARG:HB2	1.78	0.48
26:A4:43:TYR:O	26:A4:43:TYR:CG	2.66	0.48
1:AA:119:A:O2'	1:AA:120:U:P	2.72	0.48
1:AA:1292:U:H2'	1:AA:1293:C:C6	2.48	0.48
1:AA:1612:C:H4'	29:A7:5:TRP:O	2.14	0.48
1:AA:2307:G:O2'	1:AA:2308:G:C5	2.64	0.48
1:AA:2309:A:H2'	1:AA:2310:A:O4'	2.13	0.48
1:AA:2317:C:C2'	1:AA:2318:G:H5'	2.43	0.48
1:AA:235:U:H2'	1:AA:236:C:H6	1.77	0.48
1:AA:2414:G:H21	11:AO:67:MET:HE1	1.77	0.48
1:AA:2682:U:H6	1:AA:2682:U:H5'	1.77	0.48
1:AA:2681:C:O2'	1:AA:2682:U:OP2	2.28	0.48
1:AA:2867:G:C2'	1:AA:2868:A:OP2	2.61	0.48
1:AA:2867:G:O2'	1:AA:2868:A:C8	2.57	0.48
1:AA:2867:G:O2'	1:AA:2868:A:P	2.71	0.48
1:AA:456:C:O2'	1:AA:457:A:O5'	2.30	0.48
1:AA:654(L):G:C4	1:AA:654(M):C:C6	3.01	0.48
1:AA:723:G:H2'	1:AA:724:U:C6	2.48	0.48
1:AA:857:C:H1'	22:A3:26:TYR:CE2	2.49	0.48
5:AF:205:ARG:O	5:AF:206:ILE:HG23	2.13	0.48
9:AM:134:ARG:HG2	9:AM:134:ARG:O	2.13	0.48
9:AM:15:LEU:HD21	9:AM:55:VAL:HG13	1.95	0.48
10:AN:86:ILE:N	10:AN:86:ILE:HD12	2.27	0.48
11:AO:62:LEU:CD2	11:AO:63:PRO:O	2.61	0.48
18:AS:29:LEU:HD21	18:AS:33:ARG:NH2	2.28	0.48
19:AT:14:SER:O	19:AT:17:ALA:HB3	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:AU:17:SER:OG	20:AU:18:GLY:N	2.46	0.48
21:AV:56:VAL:CG1	21:AV:57:ILE:N	2.76	0.48
31:BA:1059:C:O2'	40:BM:53:PRO:HD3	2.14	0.48
31:BA:1073:U:H2'	31:BA:1074:G:C8	2.47	0.48
31:BA:865:A:H5'	31:BA:1078:U:O4	2.14	0.48
31:BA:1111:A:H2'	31:BA:1112:C:C6	2.49	0.48
31:BA:746:A:O2'	31:BA:747:C:H5'	2.14	0.48
31:BA:787:A:C2	31:BA:796:C:N3	2.82	0.48
32:BE:215:LEU:O	32:BE:218:ALA:HB3	2.13	0.48
33:BF:113:ALA:HB2	33:BF:202:ILE:HG12	1.94	0.48
33:BF:71:ALA:CA	33:BF:106:VAL:HB	2.43	0.48
34:BG:33:MET:O	34:BG:34:GLU:HB2	2.13	0.48
37:BJ:75:VAL:O	37:BJ:75:VAL:HG23	2.14	0.48
38:BK:20:TYR:CE2	38:BK:75:ARG:HD2	2.45	0.48
49:BV:31:ILE:CG2	49:BV:49:ILE:HG23	2.43	0.48
54:CA:1027:C:O2'	54:CA:1028:C:P	2.70	0.48
54:CA:1054:C:O2'	54:CA:1055:A:C5'	2.60	0.48
54:CA:1238:A:N6	54:CA:1299:A:H61	2.11	0.48
54:CA:1486:G:H2'	54:CA:1487:G:O4'	2.13	0.48
54:CA:353:A:H2'	54:CA:354:G:OP2	2.13	0.48
54:CA:392:G:H2'	54:CA:393:A:H8	1.78	0.48
54:CA:533:A:O2'	54:CA:535:A:OP2	2.19	0.48
54:CA:64:G:H5"	54:CA:65:U:OP1	2.14	0.48
54:CA:831:U:H2'	54:CA:832:C:H6	1.79	0.48
54:CA:895:G:H2'	54:CA:896:C:C6	2.49	0.48
54:CA:93:U:C3'	54:CA:95:G:H5"	2.44	0.48
52:CD:37:MIA:H111	52:CD:38:A:H1'	1.96	0.48
32:CE:117:GLU:O	32:CE:121:LEU:HB2	2.13	0.48
34:CG:108:LEU:HD23	34:CG:110:PHE:CE1	2.47	0.48
54:CA:559:A:OP1	35:CH:126:ARG:NH2	2.45	0.48
37:CJ:41:ARG:HH11	37:CJ:41:ARG:HG2	1.78	0.48
54:CA:826:C:C5'	38:CK:12:ARG:HH21	2.24	0.48
38:CK:83:ILE:HA	38:CK:136:GLU:O	2.13	0.48
38:CK:31:PHE:O	38:CK:34:GLU:HB2	2.13	0.48
38:CK:84:ARG:NH1	38:CK:86:ILE:HD13	2.28	0.48
54:CA:1149:C:P	39:CL:9:ARG:HH21	2.37	0.48
41:CN:127:LYS:HE2	41:CN:127:LYS:CA	2.32	0.48
47:CT:63:ARG:O	47:CT:65:ILE:HD12	2.14	0.48
48:CU:22:VAL:O	48:CU:25:THR:HB	2.14	0.48
26:D4:13:ARG:O	26:D4:24:THR:HG21	2.14	0.48
26:D4:55:ARG:HD2	26:D4:56:VAL:H	1.79	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1188:U:H4'	17:D2:79:VAL:HG22	1.94	0.48
55:DA:1301:A:C8	55:DA:1303:G:C8	3.01	0.48
55:DA:1485:G:H8	55:DA:1485:G:H5'	1.78	0.48
55:DA:2212:A:H1'	55:DA:2215:G:C4	2.48	0.48
55:DA:226:G:OP1	55:DA:256:A:O2'	2.31	0.48
55:DA:236:C:H2'	55:DA:237:C:H6	1.79	0.48
55:DA:2600:A:H2'	55:DA:2601:C:C6	2.48	0.48
55:DA:2771:C:H2'	55:DA:2772:C:H6	1.78	0.48
2:DB:79:C:H2'	2:DB:80:U:O4'	2.13	0.48
4:DE:203:LYS:HD2	4:DE:203:LYS:O	2.13	0.48
7:DH:98:LEU:HD12	7:DH:102:ALA:C	2.34	0.48
56:DI:1:MET:HB3	56:DI:5:ILE:CG2	2.43	0.48
58:DL:46:ALA:C	58:DL:48:MET:N	2.62	0.48
55:DA:1059:G:H4'	58:DL:71:THR:CB	2.43	0.48
55:DA:1203:G:C5'	11:DO:3:LEU:HD12	2.43	0.48
12:DP:118:LEU:CD1	12:DP:131:ILE:HG23	2.42	0.48
20:DU:61:ILE:HG23	20:DU:62:GLU:N	2.28	0.48
21:DV:8:TYR:HD1	21:DV:38:TYR:CZ	2.31	0.48
55:DA:1082:U:OP2	57:DY:45:LYS:HG3	2.13	0.48
57:DY:43:ALA:CB	57:DY:47:ASN:HA	2.41	0.48
13:A0:9:LYS:O	13:A0:11:ASN:N	2.46	0.48
1:AA:2821:A:OP2	13:A0:3:HIS:CE1	2.67	0.48
13:A0:44:LEU:O	13:A0:44:LEU:HD13	2.13	0.48
13:A0:80:PHE:O	13:A0:85:PRO:HD3	2.13	0.48
11:AO:64:LYS:HZ1	30:A8:30:ARG:HA	1.77	0.48
1:AA:1011:G:H4'	1:AA:1011:G:OP1	2.13	0.48
1:AA:1196:C:O4'	1:AA:1227:A:C2	2.67	0.48
1:AA:1204:A:C2'	1:AA:1205:U:OP2	2.61	0.48
1:AA:1667:G:OP2	1:AA:1667:G:H8	1.96	0.48
1:AA:1946:U:H2'	1:AA:1947:C:C6	2.49	0.48
1:AA:2146:C:C5'	1:AA:2147:G:OP1	2.59	0.48
1:AA:2190:G:H5'	1:AA:2190:G:H8	1.78	0.48
1:AA:2285:C:C5	28:A6:27:LYS:HE3	2.49	0.48
1:AA:2378:A:O2'	14:AQ:23:ARG:HD3	2.12	0.48
1:AA:2390:U:O5'	1:AA:2390:U:H6	1.96	0.48
1:AA:363(A):A:C3'	1:AA:363(B):G:H5''	2.43	0.48
1:AA:433:C:H2'	1:AA:434:U:C6	2.48	0.48
1:AA:733:G:O6	1:AA:761:A:C8	2.67	0.48
1:AA:940:G:H2'	1:AA:941:A:O4'	2.14	0.48
1:AA:971:C:C2'	1:AA:972:G:H5'	2.43	0.48
2:AB:10:C:O2'	2:AB:11:C:H5'	2.14	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AB:66:A:O2'	2:AB:67:G:P	2.71	0.48
3:AD:69:ARG:NH2	3:AD:192:THR:HB	2.28	0.48
3:AD:43:ARG:NH1	3:AD:44:ASN:HD22	1.95	0.48
4:AE:25:VAL:CG1	4:AE:181:LEU:HD12	2.43	0.48
6:AG:15:VAL:HG13	6:AG:175:LEU:HB2	1.95	0.48
6:AG:38:VAL:HG13	6:AG:92:VAL:O	2.12	0.48
6:AG:94:LEU:N	6:AG:94:LEU:HD23	2.27	0.48
8:AK:72:LEU:C	8:AK:74:ASN:H	2.17	0.48
11:AO:108:LYS:C	11:AO:110:TYR:H	2.17	0.48
11:AO:48:PRO:CG	11:AO:49:ARG:H	2.24	0.48
11:AO:83:VAL:HG12	11:AO:112:LEU:CD2	2.41	0.48
15:AR:58:ASN:ND2	15:AR:58:ASN:H	2.11	0.48
20:AU:83:THR:HG22	20:AU:85:VAL:HG22	1.95	0.48
24:AW:47:ASN:O	24:AW:48:HIS:C	2.52	0.48
31:BA:1054:C:C2'	31:BA:1055:A:H5''	2.44	0.48
31:BA:1206:G:C6	31:BA:1207:G:C5	3.01	0.48
31:BA:1223:C:OP1	31:BA:1224:G:H3'	2.14	0.48
31:BA:978:A:C2	31:BA:1319:A:H1'	2.48	0.48
31:BA:1363:A:N3	31:BA:1363:A:H2'	2.28	0.48
31:BA:179:A:H2'	31:BA:180:U:H6	1.77	0.48
31:BA:191(F):U:C2'	31:BA:191:G:H5'	2.43	0.48
31:BA:48:C:C5	31:BA:364:A:H2	2.32	0.48
31:BA:389:A:H3'	31:BA:390:C:C6	2.47	0.48
31:BA:397:A:H5'	31:BA:398:C:OP1	2.14	0.48
31:BA:631:G:O3'	31:BA:632:A:C8	2.56	0.48
31:BA:668:G:O2'	31:BA:669:U:H5'	2.14	0.48
31:BA:758:G:H5''	31:BA:880:C:H1'	1.96	0.48
52:BD:8:U:C2'	52:BD:13:C:H41	2.18	0.48
32:BE:17:PHE:CE2	32:BE:44:LEU:HB3	2.48	0.48
34:BG:29:PRO:CD	34:BG:30:LYS:HD3	2.44	0.48
37:BJ:148:ASN:HD22	37:BJ:148:ASN:N	2.11	0.48
38:BK:11:THR:HA	38:BK:14:ARG:NH1	2.29	0.48
39:BL:113:LYS:CD	39:BL:113:LYS:H	2.25	0.48
43:BP:90:LEU:HD13	49:BV:78:ARG:NH2	2.22	0.48
26:A4:63:TYR:CZ	49:BV:39:THR:HB	2.48	0.48
49:BV:8:GLY:O	49:BV:9:VAL:O	2.31	0.48
53:C1:38:U:H2'	53:C1:39:U:H5'	1.95	0.48
54:CA:1028(B):C:N4	54:CA:1029:G:H1'	2.28	0.48
54:CA:1181:G:N3	54:CA:1182:G:N2	2.61	0.48
54:CA:1280:A:O2'	54:CA:1281:U:OP1	2.27	0.48
54:CA:1298:C:O2'	54:CA:1299:A:C2	2.62	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1314:C:H2'	54:CA:1315:U:C6	2.48	0.48
54:CA:1221:G:OP1	54:CA:1321:C:N3	2.46	0.48
54:CA:1355:G:H2'	54:CA:1356:G:C8	2.47	0.48
54:CA:485:G:HO2'	54:CA:486:U:P	2.37	0.48
54:CA:502:G:OP1	42:CO:118:SER:CB	2.61	0.48
54:CA:788:U:C2'	54:CA:789:U:H5'	2.43	0.48
54:CA:945:G:H2'	54:CA:945:G:N3	2.28	0.48
52:CC:18:G:H4'	52:CC:18:G:OP2	2.13	0.48
52:CC:2:C:H2'	52:CC:3:C:H6	1.77	0.48
52:CD:71:G:H2'	52:CD:72:C:C6	2.48	0.48
32:CE:102:LEU:HB3	32:CE:180:LEU:HD12	1.94	0.48
32:CE:167:PRO:HG2	32:CE:168:THR:H	1.78	0.48
32:CE:216:SER:C	32:CE:218:ALA:N	2.67	0.48
33:CF:123:GLN:O	33:CF:128:PHE:HB2	2.13	0.48
37:CJ:85:TYR:HD1	37:CJ:154:TYR:CD1	2.31	0.48
37:CJ:77:SER:HB2	37:CJ:84:ASN:OD1	2.14	0.48
39:CL:81:ILE:O	39:CL:85:LEU:HG	2.14	0.48
43:CP:3:ARG:NH2	43:CP:7:VAL:HG13	2.28	0.48
46:CS:14:ASN:N	46:CS:15:PRO:CD	2.75	0.48
47:CT:22:LEU:HD13	47:CT:41:LYS:HG2	1.95	0.48
49:CV:45:VAL:HA	49:CV:62:ILE:CG2	2.43	0.48
49:CV:36:ARG:HB3	49:CV:51:VAL:CG1	2.43	0.48
13:D0:94:TYR:O	13:D0:117:VAL:HG12	2.14	0.48
17:D2:22:VAL:CG1	17:D2:23:GLU:N	2.76	0.48
55:DA:1120:G:H2'	55:DA:1121:C:C6	2.49	0.48
55:DA:140:A:H8	55:DA:1408:C:O2'	1.96	0.48
55:DA:1421:G:C2	55:DA:1422:G:C8	3.02	0.48
55:DA:2296:U:C4'	55:DA:2297:C:OP1	2.56	0.48
55:DA:2517:C:C2	55:DA:2542:A:N6	2.81	0.48
55:DA:2702:U:O2'	55:DA:2703:C:P	2.71	0.48
55:DA:247:G:H4'	55:DA:386:G:C5	2.48	0.48
55:DA:468:G:H4'	5:DF:62:ARG:NH1	2.28	0.48
55:DA:57:C:H2'	55:DA:58:G:O5'	2.14	0.48
2:DB:15:A:H1'	2:DB:109:G:N9	2.28	0.48
3:DD:35:LYS:CE	3:DD:104:TYR:HD1	2.24	0.48
6:DG:139:LEU:HA	6:DG:144:ILE:HG21	1.96	0.48
6:DG:20:ILE:O	6:DG:24:GLY:HA2	2.13	0.48
7:DH:96:ALA:HB2	7:DH:105:LEU:HB3	1.95	0.48
56:DI:7:ARG:NE	56:DI:8:ILE:HD13	2.28	0.48
57:DY:88:ALA:O	56:DJ:15:ALA:CB	2.59	0.48
8:DK:114:LEU:O	8:DK:115:ALA:CB	2.61	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:DK:69:LYS:HG3	8:DK:136:VAL:CG2	2.43	0.48
8:DK:76:THR:CG2	8:DK:77:LEU:H	2.25	0.48
58:DL:106:GLU:HG2	58:DL:109:LYS:HD2	1.95	0.48
58:DL:28:GLY:C	58:DL:30:HIS:N	2.66	0.48
55:DA:1064:C:H5'	58:DL:88:ALA:O	2.12	0.48
11:DO:144:GLU:N	11:DO:145:PRO:CD	2.74	0.48
14:DQ:20:ARG:HE	14:DQ:21:THR:HA	1.78	0.48
18:DS:12:ILE:CD1	18:DS:17:VAL:HG22	2.44	0.48
20:DU:50:ARG:CB	20:DU:53:PRO:HD2	2.44	0.48
20:DU:56:PRO:O	20:DU:57:GLN:NE2	2.47	0.48
57:DY:30:GLN:CD	57:DY:79:ALA:O	2.47	0.48
17:A2:81:TYR:N	17:A2:81:TYR:CD1	2.79	0.48
30:A8:54:GLU:O	30:A8:57:ARG:HB2	2.13	0.48
1:AA:1069:A:C4'	1:AA:1070:A:H5''	2.35	0.48
1:AA:1378:A:O2'	1:AA:1379:A:P	2.72	0.48
1:AA:1387:C:C4	1:AA:1388:G:N7	2.82	0.48
1:AA:1386:C:OP2	1:AA:1396:U:C5	2.66	0.48
1:AA:1473:G:H2'	1:AA:1474:C:O4'	2.14	0.48
1:AA:1693:U:O2'	1:AA:1694:C:OP1	2.26	0.48
1:AA:2130:U:H6	1:AA:2130:U:H3'	1.79	0.48
1:AA:2158:A:O5'	1:AA:2158:A:H8	1.96	0.48
1:AA:2206:C:H2'	1:AA:2207:C:C6	2.49	0.48
1:AA:248:G:C2	1:AA:2431:U:H4'	2.49	0.48
1:AA:2031:A:C6	1:AA:2498:C:H1'	2.49	0.48
1:AA:2748:A:N7	1:AA:2757:A:N6	2.62	0.48
1:AA:523:C:O2'	1:AA:524:U:H5'	2.13	0.48
2:AB:87:G:N2	2:AB:89(A):A:OP1	2.47	0.48
4:AE:76:ARG:CG	4:AE:195:LEU:HD22	2.43	0.48
5:AF:107:LYS:O	5:AF:206:ILE:HG21	2.14	0.48
5:AF:165:ARG:CB	5:AF:165:ARG:HH11	2.26	0.48
6:AG:41:GLN:O	6:AG:43:LEU:HD13	2.14	0.48
6:AG:97:ASP:O	6:AG:100:TRP:N	2.47	0.48
7:AH:125:VAL:CG2	7:AH:126:PRO:HA	2.43	0.48
7:AH:140:LYS:O	7:AH:144:VAL:HG23	2.13	0.48
7:AH:94:TYR:N	7:AH:94:TYR:HD1	2.12	0.48
9:AM:112:LEU:HA	9:AM:115:ARG:CB	2.42	0.48
9:AM:56:ASN:HB3	9:AM:125:GLY:C	2.34	0.48
9:AM:76:SER:O	9:AM:78:TYR:N	2.46	0.48
10:AN:19:ILE:HG22	10:AN:43:VAL:HA	1.95	0.48
1:AA:832:G:H5'	11:AO:38:GLN:OE1	2.13	0.48
12:AP:21:THR:CG2	12:AP:21:THR:O	2.59	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AP:42:ILE:CG2	12:AP:47:ILE:HG13	2.43	0.48
14:AQ:7:TYR:CZ	14:AQ:91:PRO:HG3	2.49	0.48
21:AV:94:GLU:C	21:AV:95:PRO:O	2.50	0.48
24:AW:15:LYS:HA	24:AW:67:LYS:NZ	2.25	0.48
23:AZ:64:ALA:HA	23:AZ:67:ILE:HG13	1.95	0.48
23:AZ:82:LEU:HD23	23:AZ:82:LEU:N	2.22	0.48
31:BA:1101:A:C4'	31:BA:1102:A:O5'	2.59	0.48
31:BA:1338:G:C6	31:BA:1339:A:C6	3.02	0.48
31:BA:1348:U:H4'	39:BL:120:ARG:HD2	1.96	0.48
31:BA:355:C:H4'	31:BA:388:G:O2'	2.13	0.48
31:BA:714:G:H2'	31:BA:715:A:C8	2.49	0.48
31:BA:91:C:C2'	31:BA:92:G:H5''	2.44	0.48
52:BC:53:G:H2'	52:BC:54:U:C6	2.47	0.48
31:BA:936:C:O2'	52:BD:34:G:H5'	2.13	0.48
52:BD:75:C:HO2'	52:BD:76:A:H2	1.61	0.48
34:BG:29:PRO:HD2	34:BG:30:LYS:HE2	1.95	0.48
35:BH:48:ALA:HB1	35:BH:49:PRO:CD	2.35	0.48
40:BM:22:LYS:O	40:BM:26:ALA:N	2.47	0.48
43:BP:90:LEU:CD1	49:BV:78:ARG:NH2	2.77	0.48
46:BS:14:ASN:N	46:BS:15:PRO:CD	2.75	0.48
54:CA:1128:C:H2'	54:CA:1139:G:O6	2.13	0.48
54:CA:729:A:H2	54:CA:764:C:O2	1.96	0.48
32:CE:200:ILE:H	32:CE:200:ILE:CD1	2.18	0.48
32:CE:204:ASN:HD21	32:CE:206:ASP:H	1.51	0.48
32:CE:29:ALA:HA	32:CE:32:ILE:CG2	2.43	0.48
32:CE:7:VAL:CG2	32:CE:8:LYS:N	2.77	0.48
33:CF:189:ALA:HB3	33:CF:196:LEU:CB	2.44	0.48
34:CG:122:ARG:O	34:CG:122:ARG:HD3	2.14	0.48
34:CG:68:TYR:OH	34:CG:196:LEU:HD21	2.14	0.48
36:CI:69:GLU:HG2	36:CI:70:ASP:N	2.28	0.48
37:CJ:41:ARG:NH1	37:CJ:41:ARG:HG2	2.29	0.48
35:CH:152:ARG:HD3	38:CK:44:PHE:CE1	2.49	0.48
41:CN:67:ASP:OD2	41:CN:71:LYS:HE3	2.13	0.48
43:CP:126:LYS:OXT	52:CC:27:G:C5	2.63	0.48
54:CA:982:U:H5''	44:CQ:6:LEU:HD13	1.94	0.48
45:CR:79:ARG:O	45:CR:82:ILE:HG22	2.12	0.48
46:CS:19:ILE:H	46:CS:38:TYR:HA	1.79	0.48
49:CV:7:LYS:HB3	49:CV:7:LYS:NZ	2.22	0.48
9:DM:42:TRP:N	16:D1:64:ARG:NH2	2.62	0.48
55:DA:1188:U:H4'	17:D2:79:VAL:CG2	2.43	0.48
22:D3:25:ARG:HD3	22:D3:29:GLN:NE2	2.29	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:D7:25:PRO:HA	29:D7:28:ARG:NH1	2.29	0.48
55:DA:120:U:O2	55:DA:120:U:O4'	2.29	0.48
55:DA:1354:A:P	3:DD:38:LYS:HE2	2.53	0.48
55:DA:1557:C:OP2	55:DA:1558:A:H2'	2.13	0.48
55:DA:1581:G:H2'	55:DA:1582:C:O4'	2.14	0.48
55:DA:1698:A:H4'	55:DA:1699:G:H3'	1.95	0.48
55:DA:1893:C:C5	55:DA:1894:C:C5	3.01	0.48
55:DA:270(U):C:C2	55:DA:270(V):G:C8	3.01	0.48
55:DA:2836:U:C4	55:DA:2883:A:N6	2.81	0.48
55:DA:345:A:H2'	55:DA:347:A:H62	1.77	0.48
55:DA:968:G:H2'	55:DA:969:U:H6	1.79	0.48
2:DB:91:C:OP1	21:DV:79:ARG:NH2	2.46	0.48
6:DG:67:LYS:HG2	26:D4:5:ILE:HG21	1.92	0.48
8:DK:37:VAL:CG1	8:DK:38:LEU:HD12	2.43	0.48
11:DO:127:ALA:N	11:DO:147:LEU:HD23	2.29	0.48
11:DO:146:VAL:HG13	11:DO:146:VAL:O	2.13	0.48
12:DP:90:VAL:HG13	12:DP:91:GLU:H	1.77	0.48
14:DQ:26:LEU:HD22	14:DQ:87:PHE:HD1	1.78	0.48
21:DV:105:VAL:C	21:DV:140:ASP:HA	2.32	0.48
57:DY:71:LEU:N	57:DY:113:GLN:HB3	2.29	0.48
57:DY:127:GLU:CA	57:DY:127:GLU:OE2	2.61	0.48
16:A1:95:LEU:CD1	17:A2:11:GLN:HB3	2.43	0.48
17:A2:88:ARG:O	17:A2:90:PRO:HD3	2.14	0.48
22:A3:48:GLY:HA3	22:A3:80:HIS:ND1	2.28	0.48
29:A7:48:LYS:CD	29:A7:49:ARG:H	2.26	0.48
30:A8:62:LEU:HB2	30:A8:63:PRO:CD	2.43	0.48
1:AA:1288:U:C4'	1:AA:1289:C:OP2	2.62	0.48
1:AA:1359:A:H2'	1:AA:1360:A:O5'	2.14	0.48
1:AA:1932:A:H61	1:AA:1968:G:H1'	1.79	0.48
1:AA:2009:G:C2'	1:AA:2010:G:H5'	2.44	0.48
1:AA:2057:A:H2'	1:AA:2058:A:O4'	2.14	0.48
1:AA:2291:U:H2'	1:AA:2292:C:H6	1.79	0.48
1:AA:270(K):C:O2'	1:AA:270(L):U:H5	1.96	0.48
1:AA:2748:A:C2'	1:AA:2749:A:H5'	2.42	0.48
1:AA:2763:G:H5'	1:AA:2764:A:P	2.53	0.48
1:AA:2785:C:H2'	1:AA:2786:U:H6	1.77	0.48
1:AA:309:G:H4'	20:AU:19:LYS:H	1.78	0.48
1:AA:479:A:H4'	1:AA:480:A:C5'	2.44	0.48
1:AA:670:A:C4'	1:AA:671:C:OP1	2.60	0.48
1:AA:792:G:H5''	1:AA:793:A:O5'	2.13	0.48
1:AA:994:C:OP2	16:A1:54:LYS:NZ	2.44	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:AD:25:THR:HG22	3:AD:82:ILE:O	2.13	0.48
4:AE:199:ARG:CB	4:AE:199:ARG:HH11	2.27	0.48
5:AF:124:LEU:HG	5:AF:126:VAL:HG12	1.94	0.48
5:AF:195:ASP:OD1	5:AF:197:ASP:CB	2.61	0.48
5:AF:50:SER:HB2	5:AF:94:PRO:HD3	1.95	0.48
5:AF:8:GLN:C	5:AF:8:GLN:CD	2.72	0.48
8:AK:109:ILE:N	8:AK:109:ILE:HD13	2.09	0.48
8:AK:120:ILE:O	8:AK:121:LYS:HB2	2.14	0.48
1:AA:270(Q):C:H5'	8:AK:45:LYS:NZ	2.28	0.48
9:AM:15:LEU:HD22	9:AM:53:VAL:O	2.14	0.48
12:AP:58:PHE:HD1	12:AP:61:GLY:HA2	1.78	0.48
19:AT:83:VAL:CG2	19:AT:87:GLN:HE21	2.26	0.48
20:AU:39:VAL:C	20:AU:40:GLU:OE2	2.51	0.48
20:AU:54:LYS:C	20:AU:55:TYR:CG	2.86	0.48
21:AV:94:GLU:H	21:AV:130:PRO:HD2	1.77	0.48
12:AP:60:ARG:HD3	21:AV:185:GLU:HG3	1.96	0.48
21:AV:28:MET:CG	21:AV:37:VAL:HG11	2.41	0.48
19:AT:11:PRO:HG3	24:AW:37:PHE:CD2	2.49	0.48
23:AZ:83:GLU:O	23:AZ:85:LEU:N	2.47	0.48
31:BA:1009:G:O2'	31:BA:1010:G:H5'	2.14	0.48
31:BA:1218:C:H2'	31:BA:1219:U:H6	1.75	0.48
31:BA:1305:G:N2	31:BA:1331:G:C2'	2.75	0.48
31:BA:1378:C:N3	31:BA:1379:G:H1'	2.29	0.48
31:BA:197:A:H61	31:BA:221:C:H5'	1.78	0.48
31:BA:687:A:HO2'	31:BA:688:G:P	2.37	0.48
52:BC:42:C:C3'	52:BC:43:C:C5'	2.90	0.48
36:BI:45:LEU:N	36:BI:59:TYR:CD1	2.82	0.48
40:BM:54:PHE:CG	40:BM:55:LYS:HG3	2.49	0.48
40:BM:56:HIS:O	40:BM:58:ASP:N	2.45	0.48
41:BN:91:ARG:O	41:BN:94:ALA:HB3	2.13	0.48
43:BP:15:VAL:HG22	43:BP:41:PRO:O	2.14	0.48
43:BP:81:LEU:O	43:BP:83:ASP:N	2.46	0.48
43:BP:86:CYS:O	43:BP:86:CYS:SG	2.72	0.48
46:BS:15:PRO:O	46:BS:41:PRO:HD2	2.14	0.48
48:BU:61:LYS:O	48:BU:65:ILE:HG13	2.13	0.48
49:BV:20:LEU:CD2	49:BV:43:GLU:HG2	2.43	0.48
50:BW:49:ALA:O	50:BW:53:LEU:HG	2.14	0.48
54:CA:1004:A:C4	54:CA:1025:U:C2	3.02	0.48
54:CA:1126:U:C1'	54:CA:1280:A:N7	2.76	0.48
54:CA:555:C:H2'	54:CA:556:C:C6	2.49	0.48
54:CA:913:A:HO2'	54:CA:914:A:P	2.35	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:948:C:H2'	54:CA:949:A:H8	1.79	0.48
32:CE:5:ILE:N	32:CE:5:ILE:HD13	2.29	0.48
33:CF:117:ALA:O	33:CF:118:GLN:C	2.52	0.48
54:CA:402:G:OP1	34:CG:74:GLN:HG2	2.14	0.48
39:CL:80:GLY:O	39:CL:82:ALA:N	2.46	0.48
54:CA:1178:G:C5'	39:CL:93:ARG:HH21	2.08	0.48
42:CO:37:CYS:SG	42:CO:83:VAL:HG11	2.53	0.48
43:CP:90:LEU:HD12	43:CP:91:ARG:N	2.28	0.48
45:CR:64:ARG:CZ	45:CR:64:ARG:HB2	2.44	0.48
46:CS:71:ARG:HB2	46:CS:71:ARG:NH1	2.28	0.48
54:CA:267:C:OP2	47:CT:67:LYS:HD2	2.14	0.48
48:CU:56:THR:O	48:CU:58:LEU:HD12	2.14	0.48
49:CV:5:LEU:O	49:CV:6:LYS:O	2.30	0.48
27:D5:44:THR:O	27:D5:46:CYS:N	2.47	0.48
55:DA:990:A:H5'	55:DA:1157:G:OP1	2.14	0.48
55:DA:1449:A:N3	55:DA:1530:G:H1'	2.29	0.48
55:DA:1547:C:O2'	55:DA:1548:C:H5'	2.13	0.48
55:DA:163:U:OP2	55:DA:164:U:H5	1.96	0.48
55:DA:1784:A:H4'	55:DA:1785:A:H5''	1.94	0.48
55:DA:1884:A:C3'	55:DA:1885:A:H5''	2.44	0.48
55:DA:2009:G:N3	13:D0:107:ASP:HA	2.28	0.48
55:DA:2168:G:OP1	55:DA:2168:G:H8	1.97	0.48
55:DA:270(U):C:H2'	55:DA:270(V):G:H8	1.79	0.48
55:DA:57:C:C2'	55:DA:58:G:O5'	2.62	0.48
55:DA:623:G:H2'	55:DA:624:C:C6	2.49	0.48
55:DA:747:U:C2	27:D5:2:ALA:N	2.82	0.48
55:DA:865:C:H4'	55:DA:866:A:OP1	2.13	0.48
56:DI:20:LEU:C	56:DI:24:ILE:HG23	2.28	0.48
55:DA:1080:A:C2'	58:DL:126:MET:HE2	2.43	0.48
58:DL:34:ILE:HD11	58:DL:38:VAL:HG22	1.95	0.48
55:DA:1064:C:O3'	58:DL:89:HIS:HB3	2.13	0.48
11:DO:45:LEU:HD12	11:DO:45:LEU:HA	1.50	0.48
14:DQ:48:LEU:CD2	14:DQ:82:ILE:HD11	2.44	0.48
20:DU:19:LYS:O	20:DU:19:LYS:HD2	2.13	0.48
20:DU:42:VAL:HB	20:DU:67:LEU:CD1	2.41	0.48
21:DV:191:VAL:CB	21:DV:197:ILE:CG1	2.91	0.48
21:DV:30:ASN:HD21	21:DV:32:HIS:CD2	2.32	0.48
57:DY:16:ASN:O	57:DY:16:ASN:OD1	2.30	0.48
57:DY:30:GLN:C	57:DY:31:GLY:O	2.51	0.48
57:DY:52:PHE:C	57:DY:53:VAL:CG2	2.80	0.48
57:DY:80:VAL:CG1	57:DY:81:VAL:N	2.54	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:A2:61:VAL:O	17:A2:93:GLU:O	2.32	0.48
1:AA:1098:A:C2'	1:AA:1099:G:H5''	2.43	0.48
1:AA:1558:A:O2'	1:AA:1559:G:P	2.72	0.48
1:AA:1639:U:O2'	1:AA:1640:C:H5''	2.14	0.48
1:AA:1669:A:N3	1:AA:1669:A:H2'	2.27	0.48
1:AA:1937:A:O2'	1:AA:1938:A:OP1	2.30	0.48
1:AA:2145:C:H5''	1:AA:2146:C:OP2	2.14	0.48
1:AA:2369:A:H2'	1:AA:2370:G:C8	2.49	0.48
1:AA:2519:U:C4'	1:AA:2520:C:OP1	2.43	0.48
1:AA:2572:A:OP1	4:AE:144:ARG:HB2	2.13	0.48
1:AA:2633:G:H5'	1:AA:2811:G:O2'	2.13	0.48
1:AA:2819:G:C2'	1:AA:2820:A:OP2	2.61	0.48
1:AA:510:C:H3'	1:AA:510:C:OP1	2.13	0.48
1:AA:85:G:N3	1:AA:103:A:H2	2.10	0.48
1:AA:947:G:H2'	1:AA:948:G:C8	2.48	0.48
1:AA:980:A:N7	1:AA:1136:G:H5''	2.29	0.48
3:AD:268:ARG:C	3:AD:269:PHE:HD1	2.17	0.48
4:AE:101:ARG:NH1	4:AE:171:GLU:HB3	2.28	0.48
4:AE:117:MET:HE1	4:AE:124:GLY:HA3	1.96	0.48
4:AE:144:ARG:HB3	4:AE:145:LYS:H	1.51	0.48
6:AG:2:PRO:O	6:AG:4:ASP:N	2.47	0.48
7:AH:44:VAL:HG22	7:AH:46:GLU:HG3	1.96	0.48
12:AP:43:THR:OG1	12:AP:45:GLN:HG2	2.13	0.48
12:AP:58:PHE:HD1	12:AP:58:PHE:O	1.96	0.48
12:AP:84:GLY:CA	22:A3:10:THR:HG21	2.44	0.48
1:AA:2318:G:N2	14:AQ:2:ALA:N	2.57	0.48
21:AV:111:VAL:O	21:AV:112:ARG:C	2.51	0.48
25:AX:23:LEU:HD11	25:AX:53:LEU:HD13	1.95	0.48
25:AX:57:GLU:OE2	25:AX:59:VAL:HG22	2.14	0.48
23:AZ:83:GLU:HG2	23:AZ:84:GLY:N	2.28	0.48
31:BA:1024:G:O4'	31:BA:1024:G:OP1	2.32	0.48
31:BA:1145:C:O2	31:BA:1145:C:C2'	2.62	0.48
31:BA:1161:C:H2'	31:BA:1162:C:C6	2.48	0.48
31:BA:1428:A:H2'	31:BA:1429:C:C6	2.49	0.48
31:BA:650:G:O2'	31:BA:651:C:H5'	2.12	0.48
52:BC:53:G:H2'	52:BC:54:U:H6	1.79	0.48
32:BE:9:GLU:C	32:BE:11:LEU:N	2.65	0.48
32:BE:137:ARG:C	32:BE:139:LYS:H	2.17	0.48
32:BE:5:ILE:CD1	32:BE:55:PHE:HB3	2.43	0.48
34:BG:9:CYS:C	34:BG:11:LEU:N	2.65	0.48
34:BG:4:TYR:CE2	34:BG:11:LEU:HD11	2.49	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:BH:101:ILE:HD11	35:BH:119:LEU:HD23	1.96	0.48
38:BK:64:LYS:HG2	38:BK:79:VAL:HG21	1.94	0.48
39:BL:4:TYR:HB2	39:BL:19:LEU:CB	2.40	0.48
40:BM:31:GLY:O	40:BM:32:ALA:HB2	2.14	0.48
50:BW:10:LEU:O	50:BW:12:ALA:N	2.47	0.48
54:CA:1172:C:H2'	54:CA:1173:G:H8	1.79	0.48
54:CA:1322:C:O4'	54:CA:1322:C:OP1	2.32	0.48
54:CA:412:A:HO2'	54:CA:413:G:P	2.36	0.48
54:CA:622:A:C8	54:CA:623:C:C6	3.02	0.48
52:CC:46:G:H5''	52:CC:47:U:OP2	2.14	0.48
52:CD:58:A:C6	52:CD:61:C:C6	3.02	0.48
52:CD:70:G:H2'	52:CD:71:G:H8	1.79	0.48
35:CH:9:LYS:CB	35:CH:112:LEU:HD11	2.43	0.48
36:CI:24:GLU:HA	36:CI:27:GLN:CG	2.43	0.48
36:CI:2:ARG:HB2	36:CI:4:TYR:CE2	2.49	0.48
36:CI:75:LEU:HD23	36:CI:79:LEU:HG	1.95	0.48
37:CJ:90:GLU:H	37:CJ:90:GLU:CD	2.16	0.48
45:CR:79:ARG:C	45:CR:82:ILE:HG22	2.34	0.48
49:CV:41:VAL:CB	49:CV:42:PRO:CA	2.74	0.48
16:D1:52:ARG:CG	16:D1:52:ARG:HH11	2.26	0.48
16:D1:92:ARG:HD3	16:D1:92:ARG:O	2.13	0.48
6:DG:5:VAL:HG22	26:D4:25:TYR:CE2	2.48	0.48
13:D0:33:ARG:HH12	27:D5:57:VAL:HG23	1.78	0.48
28:D6:15:GLU:CG	28:D6:16:CYS:N	2.60	0.48
30:D8:17:THR:HG23	30:D8:21:LYS:C	2.34	0.48
55:DA:1101:U:H2'	55:DA:1102:C:H6	1.79	0.48
55:DA:2645:G:H3'	55:DA:2646:C:H5'	1.94	0.48
55:DA:557:U:O2	9:DM:45:ASN:HB2	2.13	0.48
4:DE:13:ARG:HA	4:DE:21:VAL:HA	1.95	0.48
4:DE:21:VAL:CB	4:DE:22:PRO:HB3	2.16	0.48
7:DH:153:LYS:HE2	7:DH:153:LYS:CA	2.43	0.48
56:DJ:14:GLN:CA	56:DJ:15:ALA:O	2.61	0.48
58:DL:13:PRO:O	58:DL:50:ASP:HA	2.14	0.48
11:DO:108:LYS:O	11:DO:110:TYR:N	2.44	0.48
14:DQ:5:THR:HG1	14:DQ:7:TYR:HB3	1.76	0.48
20:DU:44:ILE:HD12	20:DU:45:VAL:HG23	1.95	0.48
21:DV:184:ALA:O	21:DV:185:GLU:CB	2.51	0.48
21:DV:187:ALA:HB1	21:DV:193:GLU:HG2	1.90	0.48
16:A1:58:ARG:HA	16:A1:61:TRP:HE3	1.74	0.48
26:A4:67:TYR:C	26:A4:67:TYR:CD2	2.87	0.48
1:AA:1015:G:C6	1:AA:1148:A:C6	3.01	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1254:A:H5'	1:AA:1255:U:C5'	2.43	0.48
1:AA:1272:A:C3'	1:AA:1273:U:C5'	2.92	0.48
1:AA:2114:A:H1'	1:AA:2168:G:H1'	1.95	0.48
1:AA:365:C:O2'	1:AA:366:C:H5'	2.14	0.48
1:AA:404:C:O2'	1:AA:405:U:P	2.72	0.48
1:AA:532:A:O2'	1:AA:533:G:P	2.69	0.48
1:AA:649:G:H2'	1:AA:650:C:H6	1.79	0.48
1:AA:916:G:C2'	1:AA:917:A:H5''	2.43	0.48
3:AD:131:LEU:CD1	3:AD:131:LEU:N	2.77	0.48
1:AA:2052:G:O4'	4:AE:142:GLY:HA3	2.14	0.48
5:AF:170:LEU:HD23	5:AF:172:TRP:NE1	2.28	0.48
10:AN:26:LYS:HB2	10:AN:30:ALA:HB2	1.95	0.48
10:AN:88:ASN:ND2	10:AN:90:GLN:H	2.12	0.48
11:AO:109:GLY:O	11:AO:110:TYR:O	2.31	0.48
11:AO:66:GLY:O	11:AO:67:MET:HB3	2.14	0.48
14:AQ:36:TYR:CD1	14:AQ:36:TYR:N	2.82	0.48
15:AR:92:GLY:HA2	15:AR:117:ASP:H	1.78	0.48
20:AU:42:VAL:CG1	20:AU:67:LEU:HD13	2.44	0.48
20:AU:75:ILE:HD13	20:AU:76:CYS:N	2.29	0.48
1:AA:896:A:C2	21:AV:178:GLU:OE2	2.57	0.48
25:AX:6:VAL:HG13	25:AX:56:VAL:HG23	1.95	0.48
31:BA:1145:C:H5'	31:BA:1146:A:OP1	2.13	0.48
31:BA:1253:G:N1	31:BA:1285:A:N6	2.62	0.48
31:BA:1364:U:O2'	31:BA:1365:G:OP1	2.29	0.48
31:BA:923:A:O2'	31:BA:1399:C:OP2	2.31	0.48
31:BA:1428:A:H2'	31:BA:1429:C:O4'	2.14	0.48
31:BA:115:G:C2	31:BA:289:G:N7	2.82	0.48
31:BA:44:G:C2	31:BA:45:U:H1'	2.48	0.48
31:BA:726:C:O2'	31:BA:727:G:H5'	2.14	0.48
31:BA:738:C:H2'	31:BA:739:C:H6	1.79	0.48
31:BA:819:A:H4'	31:BA:820:U:OP2	2.14	0.48
52:BB:42:C:H2'	52:BB:43:C:C6	2.48	0.48
52:BB:21:A:N6	52:BB:46:G:C4	2.82	0.48
1:AA:2554:U:N3	52:BB:74:C:H5	2.12	0.48
52:BB:9:A:HO2'	52:BB:10:G:P	2.33	0.48
33:BF:39:ILE:O	33:BF:43:LEU:HB2	2.14	0.48
34:BG:190:ASP:O	34:BG:191:ARG:C	2.52	0.48
34:BG:24:GLU:N	34:BG:27:TYR:HB3	2.29	0.48
37:BJ:111:ARG:HH12	37:BJ:122:HIS:CB	2.27	0.48
39:BL:3:GLN:HG2	39:BL:20:ARG:NH1	2.29	0.48
40:BM:24:VAL:O	40:BM:28:ARG:HB2	2.14	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:BN:103:LEU:N	41:BN:103:LEU:HD22	2.29	0.48
41:BN:48:ILE:HD11	41:BN:64:ALA:HA	1.95	0.48
46:BS:18:ARG:HD3	46:BS:35:LYS:CD	2.43	0.48
26:A4:63:TYR:HE1	49:BV:39:THR:HG21	1.77	0.48
49:BV:76:PRO:HB2	49:BV:78:ARG:HH11	1.75	0.48
54:CA:1023:G:H3'	54:CA:1024:G:H5''	1.96	0.48
54:CA:113:G:H2'	54:CA:114:U:H6	1.78	0.48
54:CA:1241:G:H2'	54:CA:1242:C:H6	1.77	0.48
54:CA:1256:A:H2	54:CA:1277:C:C2'	2.26	0.48
54:CA:392:G:H5'	46:CS:12:LYS:HG3	1.94	0.48
54:CA:812:C:HO2'	54:CA:813:U:H6	1.57	0.48
54:CA:890:G:H2'	54:CA:906:G:O6	2.13	0.48
52:CD:40:C:H2'	52:CD:41:C:C6	2.48	0.48
32:CE:161:ALA:HA	32:CE:183:PRO:O	2.14	0.48
34:CG:142:PRO:HA	34:CG:185:PHE:O	2.14	0.48
34:CG:61:LYS:HE2	34:CG:65:ARG:CD	2.44	0.48
38:CK:1:MET:O	38:CK:2:LEU:HB2	2.14	0.48
40:CM:17:ASP:C	40:CM:17:ASP:OD2	2.52	0.48
40:CM:98:ILE:HD12	40:CM:98:ILE:N	2.28	0.48
41:CN:30:VAL:HG21	41:CN:65:ALA:HA	1.95	0.48
46:CS:20:VAL:HG22	46:CS:21:VAL:H	1.79	0.48
46:CS:43:LYS:HA	46:CS:48:TRP:HB2	1.96	0.48
55:DA:1151:G:H5''	16:D1:81:HIS:NE2	2.28	0.48
28:D6:52:VAL:O	28:D6:53:LYS:C	2.52	0.48
55:DA:101:G:HO2'	55:DA:102:G:P	2.37	0.48
55:DA:1077:A:C2'	55:DA:1078:U:C5'	2.92	0.48
55:DA:141(A):C:O5'	55:DA:141(A):C:H6	1.97	0.48
55:DA:184:C:H2'	55:DA:185:U:C6	2.49	0.48
55:DA:1899:G:O2'	55:DA:1900:A:C5'	2.62	0.48
55:DA:1950:G:O6	55:DA:1954:G:H2'	2.14	0.48
55:DA:2210:G:H5'	55:DA:2211:G:OP2	2.14	0.48
55:DA:26:G:C6	55:DA:27:G:N1	2.82	0.48
55:DA:2732:G:H3'	55:DA:2733:A:C5'	2.40	0.48
55:DA:299:A:OP2	55:DA:299:A:H8	1.95	0.48
55:DA:634:C:H2'	55:DA:635:C:C6	2.47	0.48
2:DB:37:C:C2'	2:DB:38:C:H5'	2.44	0.48
4:DE:103:ASP:CG	4:DE:201:THR:HA	2.33	0.48
4:DE:1:MET:O	4:DE:2:LYS:C	2.52	0.48
7:DH:92:ILE:CD1	7:DH:160:LYS:HD3	2.44	0.48
58:DL:95:LYS:CB	58:DL:136:VAL:HG11	2.44	0.48
10:DN:1:MET:CE	10:DN:67:LYS:HG2	2.43	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:DN:36:GLY:HA2	10:DN:106:LEU:HD23	1.94	0.48
10:DN:59:LYS:O	10:DN:87:ILE:HG12	2.14	0.48
15:DR:28:VAL:CG2	15:DR:86:ILE:HG13	2.44	0.48
19:DT:88:LYS:NZ	19:DT:88:LYS:HB3	2.28	0.48
55:DA:1107:G:OP1	57:DY:56:ASN:OD1	2.32	0.48
23:DZ:76:ARG:N	23:DZ:76:ARG:HD2	2.29	0.48
16:A1:31:SER:O	16:A1:32:PHE:C	2.52	0.48
17:A2:18:LEU:HD23	17:A2:19:LYS:N	2.29	0.48
17:A2:65:GLY:O	17:A2:91:TYR:HB2	2.14	0.48
17:A2:85:LYS:HD3	17:A2:86:GLY:H	1.78	0.48
11:AO:64:LYS:HG3	30:A8:30:ARG:HH12	1.78	0.48
30:A8:61:LEU:CD1	30:A8:62:LEU:N	2.77	0.48
1:AA:1100:C:H2'	1:AA:1101:U:C6	2.49	0.48
1:AA:1225:C:O2'	17:A2:84:LYS:HA	2.13	0.48
1:AA:1301:A:C2'	1:AA:1302:A:H5''	2.43	0.48
1:AA:139:G:O2'	1:AA:140:A:C2	2.66	0.48
1:AA:1790:C:H4'	3:AD:209:ALA:HB2	1.94	0.48
1:AA:2206:C:H2'	1:AA:2207:C:H6	1.79	0.48
1:AA:221:A:H1'	1:AA:233:A:H1'	1.96	0.48
1:AA:2345:G:O2'	1:AA:2382:G:H5'	2.14	0.48
1:AA:2432:A:C6	23:AZ:33:LYS:HB3	2.49	0.48
1:AA:2458:G:O6	1:AA:2490:G:O2'	2.29	0.48
1:AA:2543:G:H2'	1:AA:2544:G:C8	2.48	0.48
1:AA:2729:G:C6	1:AA:2730:C:C4	3.02	0.48
1:AA:2870:C:H2'	1:AA:2871:C:H5'	1.95	0.48
1:AA:414:C:O2	1:AA:1864:U:O2'	2.27	0.48
1:AA:483:A:H3'	1:AA:484:C:C6	2.48	0.48
1:AA:937:U:H2'	1:AA:938:G:O4'	2.14	0.48
3:AD:132:PRO:O	3:AD:136:ILE:HD13	2.14	0.48
4:AE:39:PRO:HA	4:AE:43:GLY:CA	2.44	0.48
4:AE:61:ARG:C	4:AE:63:LEU:N	2.66	0.48
6:AG:75:LYS:HG3	6:AG:76:SER:N	2.29	0.48
1:AA:1012:U:H3	9:AM:25:ARG:HD3	1.78	0.48
20:AU:49:VAL:O	20:AU:50:ARG:C	2.52	0.48
21:AV:124:ILE:HG23	21:AV:124:ILE:O	2.13	0.48
23:AZ:87:PRO:O	23:AZ:89:GLU:N	2.45	0.48
53:B1:35:A:H2'	53:B1:36:G:C8	2.48	0.48
31:BA:1043:C:O2'	31:BA:1044:A:H5'	2.14	0.48
31:BA:1171:G:O2'	31:BA:1172:C:H5'	2.14	0.48
31:BA:426:G:OP1	34:BG:38:TYR:OH	2.23	0.48
31:BA:4:U:O2'	31:BA:5:U:P	2.72	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:509:A:H2'	31:BA:510:A:C8	2.49	0.48
31:BA:553:A:H2'	31:BA:554:C:H6	1.78	0.48
31:BA:5:U:H2'	31:BA:5:U:O2	2.13	0.48
31:BA:713:G:H21	31:BA:777:A:C1'	2.25	0.48
52:BB:23:A:H2'	52:BB:24:G:C8	2.48	0.48
1:AA:2395:C:C2	52:BD:76:A:H1'	2.49	0.48
32:BE:80:ILE:HD12	32:BE:212:GLN:HA	1.96	0.48
33:BF:76:VAL:HG23	33:BF:77:ILE:HG13	1.95	0.48
34:BG:18:LYS:HG2	34:BG:21:LEU:HD11	1.96	0.48
34:BG:34:GLU:O	34:BG:35:ARG:CB	2.61	0.48
34:BG:3:ARG:CB	34:BG:3:ARG:NH2	2.77	0.48
37:BJ:121:ALA:O	37:BJ:125:MET:HG3	2.14	0.48
31:BA:186:C:O2'	50:BW:82:SER:HA	2.14	0.48
54:CA:371:G:H21	54:CA:374:A:N6	2.12	0.48
54:CA:646:U:H2'	54:CA:647:C:C6	2.49	0.48
54:CA:659:U:C2	54:CA:660:G:C8	3.02	0.48
54:CA:872:A:C6	54:CA:874:G:C5	3.02	0.48
54:CA:96:G:H2'	54:CA:97:U:O4'	2.14	0.48
54:CA:9:G:OP1	35:CH:121:LYS:HD2	2.14	0.48
36:CI:98:LEU:HA	48:CU:31:LEU:HD22	1.95	0.48
37:CJ:18:TYR:OH	37:CJ:58:PRO:HG2	2.14	0.48
45:CR:39:LEU:O	45:CR:42:HIS:HB3	2.13	0.48
45:CR:76:GLU:O	45:CR:78:TYR:N	2.47	0.48
46:CS:1:MET:HG2	46:CS:2:VAL:O	2.13	0.48
36:CI:43:LEU:HD11	48:CU:35:ARG:NH1	2.29	0.48
49:CV:44:MET:O	49:CV:47:HIS:N	2.44	0.48
50:CW:89:ARG:O	50:CW:93:GLU:HB3	2.14	0.48
16:D1:68:ALA:HB1	16:D1:106:PHE:CE2	2.48	0.48
16:D1:83:LEU:CD1	16:D1:113:ALA:HB2	2.44	0.48
16:D1:88:ILE:HB	16:D1:90:VAL:HG23	1.95	0.48
55:DA:1021:A:H3'	55:DA:1021:A:H8	1.77	0.48
55:DA:1057:A:N1	55:DA:1082:U:O2	2.47	0.48
55:DA:1729:A:O2'	55:DA:1730:U:H5''	2.14	0.48
55:DA:2010:G:H5''	18:DS:42:ARG:HB2	1.95	0.48
55:DA:2345:G:O2'	55:DA:2381:C:C2'	2.61	0.48
55:DA:2567:G:H2'	55:DA:2568:C:C6	2.49	0.48
55:DA:2602:A:H5'	55:DA:2603:G:C5'	2.44	0.48
55:DA:2783:G:H2'	55:DA:2784:C:C6	2.49	0.48
55:DA:321:G:O2'	55:DA:340:A:H1'	2.13	0.48
55:DA:363(B):G:H2'	55:DA:363(C):G:H8	1.78	0.48
55:DA:363(F):A:H1'	55:DA:364:C:C5	2.49	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:447:A:N1	55:DA:454:A:H2'	2.29	0.48
55:DA:492:A:H2'	55:DA:493:G:O4'	2.14	0.48
55:DA:859:G:O2'	55:DA:860:U:OP2	2.31	0.48
3:DD:130:ALA:HA	3:DD:192:THR:HA	1.96	0.48
3:DD:253:GLN:HB2	3:DD:257:LEU:HG	1.95	0.48
5:DF:178:PRO:HB3	5:DF:198:ALA:CB	2.44	0.48
6:DG:151:ALA:HB3	6:DG:153:ARG:HH12	1.79	0.48
6:DG:88:ILE:C	6:DG:88:ILE:CD1	2.83	0.48
58:DL:110:GLN:C	58:DL:111:LYS:CE	2.82	0.48
58:DL:67:PHE:O	58:DL:68:VAL:HG12	2.14	0.48
58:DL:53:VAL:HG12	58:DL:76:TYR:CD2	2.48	0.48
9:DM:63:THR:O	9:DM:66:LYS:HG3	2.14	0.48
15:DR:102:ILE:HB	15:DR:110:ILE:HD11	1.92	0.48
15:DR:136:GLN:HG3	15:DR:137:LYS:H	1.78	0.48
15:DR:16:ARG:HG2	15:DR:18:ASP:OD1	2.14	0.48
20:DU:45:VAL:HG12	20:DU:60:PHE:HB3	1.95	0.48
20:DU:75:ILE:HD13	20:DU:76:CYS:N	2.28	0.48
21:DV:127:LYS:O	21:DV:161:VAL:CG2	2.62	0.48
21:DV:150:LEU:CD2	21:DV:154:ASP:OD1	2.62	0.48
21:DV:38:TYR:CG	21:DV:38:TYR:O	2.66	0.48
24:DW:69:ARG:HB2	24:DW:69:ARG:CZ	2.44	0.48
57:DY:121:ASP:O	57:DY:122:VAL:C	2.52	0.48
57:DY:73:GLY:CA	57:DY:112:LEU:CG	2.92	0.48
17:A2:43:GLU:HA	17:A2:43:GLU:OE2	2.14	0.47
17:A2:5:VAL:HG23	17:A2:37:VAL:HG21	1.95	0.47
28:A6:28:ARG:C	28:A6:29:ASN:HD22	2.17	0.47
30:A8:61:LEU:HD13	30:A8:62:LEU:N	2.28	0.47
1:AA:1198:U:C2	1:AA:1199:U:C5	3.02	0.47
1:AA:323:G:O2'	1:AA:1205:U:C2	2.67	0.47
1:AA:1515:C:H2'	1:AA:1516:U:H6	1.78	0.47
1:AA:1614:A:H62	18:AS:93:ALA:CB	2.22	0.47
1:AA:1688:U:O2	1:AA:1700:A:C8	2.67	0.47
1:AA:1829:A:O5'	1:AA:1830:C:H5	1.97	0.47
1:AA:184:C:O4'	1:AA:216:A:H2	1.97	0.47
1:AA:2180:U:H2'	1:AA:2181:G:C8	2.49	0.47
1:AA:2331:G:O2'	1:AA:2336:A:N1	2.36	0.47
1:AA:2439:A:H4'	1:AA:2440:C:O5'	2.14	0.47
1:AA:2554:U:O2	52:BB:74:C:H5	1.96	0.47
1:AA:2749:A:H62	1:AA:2750:A:H61	1.56	0.47
1:AA:2795:G:H3'	1:AA:2797:U:H5'	1.94	0.47
1:AA:374:A:H3'	1:AA:375:C:H6	1.79	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:654(F):C:H6	1:AA:654(F):C:O5'	1.96	0.47
1:AA:753:C:H2'	1:AA:754:C:C6	2.49	0.47
1:AA:1828:G:O6	3:AD:222:ARG:HD3	2.14	0.47
5:AF:125:LEU:HD12	5:AF:196:LEU:HD23	1.96	0.47
12:AP:59:ARG:O	12:AP:60:ARG:HB2	2.14	0.47
18:AS:75:TYR:OH	18:AS:104:THR:HG21	2.13	0.47
20:AU:33:LYS:CE	20:AU:34:LYS:HG2	2.44	0.47
53:B1:51:U:H3'	53:B1:51:U:C6	2.49	0.47
31:BA:1190:G:OP1	33:BF:5:ILE:HD12	2.14	0.47
31:BA:1320:C:O2	49:BV:72:GLY:HA3	2.14	0.47
31:BA:353:A:C8	31:BA:353:A:H5'	2.43	0.47
31:BA:477:G:H2'	31:BA:478:A:H8	1.78	0.47
31:BA:649:G:C6	31:BA:650:G:N7	2.82	0.47
31:BA:680:C:H2'	31:BA:681:C:C6	2.49	0.47
31:BA:5:U:H2'	31:BA:6:G:OP2	2.13	0.47
31:BA:700:G:H4'	31:BA:704:A:H1'	1.96	0.47
31:BA:877:C:H1'	38:BK:3:THR:OG1	2.12	0.47
34:BG:173:TRP:CD1	34:BG:174:LEU:HG	2.49	0.47
34:BG:19:LEU:O	34:BG:21:LEU:HG	2.14	0.47
34:BG:59:ARG:NH2	34:BG:66:ARG:HH12	2.12	0.47
36:BI:1:MET:SD	36:BI:68:PRO:HD3	2.54	0.47
40:BM:34:VAL:CG2	40:BM:74:ILE:HG22	2.35	0.47
43:BP:15:VAL:HG23	43:BP:41:PRO:HA	1.96	0.47
46:BS:21:VAL:HG11	46:BS:34:GLU:HB3	1.95	0.47
49:BV:47:HIS:O	49:BV:48:THR:OG1	2.26	0.47
54:CA:1234:C:O2'	54:CA:1235:U:H5'	2.14	0.47
54:CA:1256:A:H2	54:CA:1277:C:H2'	1.79	0.47
54:CA:250:A:O2'	54:CA:251:G:OP2	2.29	0.47
54:CA:340:U:H2'	54:CA:341:C:C6	2.49	0.47
54:CA:405:U:C3'	54:CA:406:G:H5'	2.40	0.47
54:CA:789:U:C3'	54:CA:789:U:O2	2.61	0.47
54:CA:7:G:H5'	54:CA:298:A:O4'	2.13	0.47
54:CA:815:A:O2'	54:CA:1527:C:O4'	2.32	0.47
52:CC:56:C:H42	6:DG:83:ARG:HH22	1.61	0.47
32:CE:221:LEU:C	32:CE:221:LEU:HD13	2.34	0.47
32:CE:5:ILE:O	32:CE:6:THR:O	2.32	0.47
32:CE:83:MET:O	32:CE:85:ALA:N	2.47	0.47
33:CF:52:LEU:N	33:CF:52:LEU:HD23	2.29	0.47
43:CP:74:VAL:O	43:CP:78:ILE:HG13	2.14	0.47
46:CS:75:ARG:O	46:CS:77:ALA:N	2.41	0.47
50:CW:86:ARG:HG3	50:CW:86:ARG:NH1	2.28	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:D1:78:THR:O	16:D1:79:PHE:C	2.52	0.47
6:DG:67:LYS:CE	26:D4:6:HIS:NE2	2.67	0.47
27:D5:31:VAL:HG13	27:D5:42:PRO:HG3	1.96	0.47
28:D6:20:ASN:O	28:D6:21:TYR:HB2	2.14	0.47
55:DA:2656:U:O4	55:DA:2657:A:N7	2.46	0.47
55:DA:2712:U:OP1	55:DA:2714:G:H4'	2.14	0.47
55:DA:26:G:N1	55:DA:27:G:N2	2.62	0.47
55:DA:338:G:N2	55:DA:339:U:H1'	2.29	0.47
55:DA:654(S):G:O2'	55:DA:654(T):A:O5'	2.17	0.47
55:DA:72:U:H3	24:DW:62:THR:HG22	1.79	0.47
55:DA:698:C:O2'	55:DA:734:A:N6	2.47	0.47
55:DA:756:C:N4	55:DA:757:U:C4	2.82	0.47
55:DA:881:G:C4	55:DA:882:G:H1'	2.49	0.47
3:DD:134:ARG:CB	3:DD:135:PHE:HD2	2.26	0.47
3:DD:28:GLU:HB2	3:DD:29:PRO:HD2	1.96	0.47
4:DE:5:LEU:O	4:DE:51:PHE:HE2	1.97	0.47
7:DH:154:PRO:HD3	7:DH:161:GLY:C	2.34	0.47
7:DH:56:SER:OG	7:DH:58:GLU:HG3	2.13	0.47
56:DJ:1:MET:O	56:DJ:5:ILE:HG13	2.14	0.47
8:DK:37:VAL:HG12	8:DK:38:LEU:N	2.28	0.47
58:DL:21:PRO:HG2	58:DL:24:GLY:CA	2.26	0.47
9:DM:61:ARG:HE	9:DM:61:ARG:CA	2.26	0.47
9:DM:99:LEU:O	9:DM:99:LEU:HD22	2.13	0.47
11:DO:112:LEU:HD11	11:DO:114:ILE:HG22	1.94	0.47
11:DO:91:PHE:HE2	11:DO:95:VAL:HG22	1.78	0.47
15:DR:2:ASN:O	15:DR:3:ARG:HB3	2.14	0.47
15:DR:39:ARG:CG	15:DR:40:THR:N	2.76	0.47
21:DV:185:GLU:OE1	21:DV:185:GLU:C	2.52	0.47
21:DV:191:VAL:CG1	21:DV:197:ILE:CB	2.49	0.47
21:DV:35:ARG:NH1	21:DV:35:ARG:CB	2.77	0.47
24:DW:41:ILE:O	24:DW:43:GLN:N	2.47	0.47
57:DY:29:TYR:HA	57:DY:80:VAL:HG13	1.96	0.47
23:DZ:85:LEU:N	23:DZ:85:LEU:CD2	2.77	0.47
23:DZ:58:ILE:HG21	23:DZ:87:PRO:HG3	1.96	0.47
13:A0:25:ALA:O	13:A0:29:LEU:HB2	2.14	0.47
16:A1:60:LEU:HD11	16:A1:64:ARG:HH21	1.80	0.47
17:A2:5:VAL:CG2	17:A2:6:LYS:N	2.77	0.47
22:A3:55:ARG:NH1	22:A3:55:ARG:CB	2.75	0.47
26:A4:53:GLU:CG	26:A4:54:GLY:N	2.66	0.47
30:A8:60:LEU:O	30:A8:61:LEU:C	2.51	0.47
1:AA:1061:U:H1'	1:AA:1070:A:H1'	1.96	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1204:A:N3	1:AA:1206:G:C2	2.82	0.47
1:AA:141:A:H8	1:AA:1595:G:N2	2.04	0.47
1:AA:1485:G:O2'	1:AA:1486:A:H5'	2.14	0.47
1:AA:1543:A:C1'	1:AA:1545:A:O4'	2.59	0.47
1:AA:1682:G:H2'	1:AA:1683:C:H6	1.75	0.47
1:AA:1753:G:N1	1:AA:1756:G:C2	2.82	0.47
1:AA:189:G:H1	1:AA:205:G:HO2'	1.61	0.47
1:AA:2001:A:H2'	1:AA:2002:G:C8	2.49	0.47
1:AA:2187:G:O2'	1:AA:2188:C:H5'	2.14	0.47
1:AA:2378:A:H4'	14:AQ:23:ARG:CZ	2.44	0.47
1:AA:270(Q):C:H4'	8:AK:45:LYS:HE3	1.95	0.47
1:AA:2851:A:H2'	1:AA:2852:G:C8	2.49	0.47
1:AA:446:G:H4'	1:AA:449:A:N3	2.30	0.47
1:AA:859:G:C2'	1:AA:860:U:OP2	2.63	0.47
1:AA:923:C:H2'	1:AA:924:C:C6	2.49	0.47
1:AA:955:C:H5'	1:AA:956:G:P	2.54	0.47
3:AD:93:ALA:N	3:AD:107:ALA:HB2	2.29	0.47
4:AE:76:ARG:HG3	4:AE:195:LEU:HD22	1.95	0.47
6:AG:97:ASP:O	6:AG:101:ILE:HG23	2.14	0.47
6:AG:139:LEU:HD12	6:AG:139:LEU:C	2.35	0.47
7:AH:23:ARG:N	7:AH:36:PRO:HA	2.29	0.47
7:AH:43:VAL:O	7:AH:43:VAL:HG23	2.14	0.47
7:AH:92:ILE:HG22	7:AH:93:GLY:N	2.28	0.47
8:AK:104:GLN:O	8:AK:105:HIS:HD2	1.97	0.47
9:AM:39:ARG:HG2	9:AM:40:PRO:HD2	1.97	0.47
15:AR:24:PRO:O	15:AR:94:ALA:HB2	2.13	0.47
18:AS:4:LYS:HG2	18:AS:106:ILE:CG2	2.42	0.47
20:AU:63:LYS:HA	20:AU:63:LYS:HZ3	1.77	0.47
21:AV:145:GLU:OE1	21:AV:174:VAL:HB	2.13	0.47
21:AV:53:ILE:C	21:AV:70:LEU:HD21	2.33	0.47
31:BA:1279:A:O2'	31:BA:1282:C:N4	2.48	0.47
31:BA:1336:C:O2'	31:BA:1337:G:OP2	2.25	0.47
31:BA:384:G:H2'	31:BA:385:C:C6	2.49	0.47
31:BA:658:G:H2'	31:BA:659:U:C6	2.49	0.47
31:BA:674:G:H2'	31:BA:675:A:C8	2.49	0.47
31:BA:713:G:N2	31:BA:777:A:C1'	2.76	0.47
32:BE:51:LEU:HD23	32:BE:201:ILE:HG23	1.97	0.47
33:BF:25:GLY:O	33:BF:27:LYS:N	2.47	0.47
34:BG:33:MET:O	34:BG:34:GLU:CB	2.61	0.47
37:BJ:18:TYR:CD2	37:BJ:59:LEU:HD22	2.47	0.47
39:BL:16:ARG:O	39:BL:63:ILE:HG23	2.14	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:BM:34:VAL:HA	40:BM:74:ILE:HA	1.95	0.47
42:BO:36:VAL:O	42:BO:58:VAL:HA	2.13	0.47
49:BV:40:ILE:CD1	49:BV:62:ILE:HD13	2.44	0.47
54:CA:1028:C:C4	54:CA:1028(A):C:C5	3.03	0.47
54:CA:128:G:C5	54:CA:129:U:C5	3.02	0.47
54:CA:1336:C:H2'	54:CA:1336:C:O2	2.13	0.47
54:CA:411:A:C6	54:CA:429:U:C5	3.02	0.47
54:CA:498:A:O2'	54:CA:500:G:C5'	2.62	0.47
54:CA:647:C:O2'	54:CA:648:A:H5'	2.14	0.47
54:CA:833:U:H2'	54:CA:834:C:C6	2.49	0.47
52:CD:33:U:O5'	52:CD:33:U:H6	1.98	0.47
33:CF:114:PRO:HD3	33:CF:183:ASP:OD1	2.15	0.47
33:CF:28:GLN:HA	33:CF:31:HIS:HD2	1.78	0.47
33:CF:52:LEU:O	33:CF:115:LEU:HD21	2.14	0.47
35:CH:153:LYS:HD3	35:CH:153:LYS:C	2.35	0.47
35:CH:10:MET:HA	35:CH:32:VAL:HG13	1.97	0.47
35:CH:50:GLU:CD	35:CH:51:VAL:H	2.17	0.47
44:CQ:40:CYS:SG	44:CQ:42:ILE:HB	2.54	0.47
49:CV:63:THR:HG23	49:CV:65:ASN:ND2	2.26	0.47
54:CA:191:G:N3	50:CW:105:SER:HB3	2.29	0.47
28:D6:52:VAL:HG22	28:D6:53:LYS:N	2.25	0.47
55:DA:1048:A:H2	55:DA:1112:G:N3	2.12	0.47
55:DA:1044:G:N3	55:DA:1111:A:N1	2.63	0.47
55:DA:1839:G:N3	55:DA:1839:G:H2'	2.29	0.47
55:DA:1858:G:O2'	55:DA:1884:A:N6	2.46	0.47
55:DA:1879:C:C2'	55:DA:1880:C:C5'	2.91	0.47
55:DA:2173:A:H3'	55:DA:2174:C:O4'	2.14	0.47
55:DA:2174:C:H2'	55:DA:2175:C:H6	1.79	0.47
55:DA:2286:A:OP2	28:D6:28:ARG:CG	2.62	0.47
55:DA:2290:G:H2'	55:DA:2291:U:O4'	2.15	0.47
55:DA:2698:U:H2'	55:DA:2699:C:H6	1.77	0.47
55:DA:331:A:H4'	55:DA:332:A:OP1	2.14	0.47
55:DA:654(J):A:C2	55:DA:654(K):C:C4	3.02	0.47
3:DD:35:LYS:HB3	3:DD:36:PRO:HA	1.96	0.47
5:DF:128:ALA:O	5:DF:129:PHE:CB	2.61	0.47
5:DF:170:LEU:HD23	5:DF:172:TRP:CZ2	2.49	0.47
5:DF:57:VAL:HG13	5:DF:58:ALA:H	1.78	0.47
6:DG:173:LEU:O	6:DG:178:PHE:HB2	2.14	0.47
7:DH:126:PRO:HG2	7:DH:130:ARG:O	2.14	0.47
7:DH:151:ILE:O	7:DH:152:ARG:HG2	2.15	0.47
8:DK:92:VAL:O	8:DK:120:ILE:CG2	2.62	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:DN:93:PRO:CG	10:DN:113:LYS:HD3	2.44	0.47
21:DV:5:LEU:O	21:DV:6:LYS:C	2.52	0.47
21:DV:5:LEU:C	21:DV:6:LYS:HG3	2.34	0.47
24:DW:13:ALA:HA	24:DW:16:LEU:CD2	2.43	0.47
57:DY:132:ASP:HA	57:DY:134:LEU:HD22	1.97	0.47
57:DY:5:ARG:O	57:DY:7:VAL:HB	2.14	0.47
13:A0:104:ARG:HH11	13:A0:104:ARG:HB2	1.80	0.47
1:AA:1225:C:H5"	17:A2:85:LYS:HE3	1.96	0.47
22:A3:25:ARG:HG2	22:A3:25:ARG:HH11	1.78	0.47
2:AB:39:A:C2'	26:A4:1:MET:CE	2.76	0.47
28:A6:53:LYS:NZ	28:A6:53:LYS:HB3	2.29	0.47
29:A7:47:ARG:NH1	29:A7:47:ARG:HB2	2.29	0.47
29:A7:47:ARG:HD3	29:A7:47:ARG:N	2.29	0.47
1:AA:1360:A:N6	1:AA:1372:U:O4	2.47	0.47
1:AA:1537:C:O2'	1:AA:1538:G:O4'	2.28	0.47
1:AA:1694:C:H1'	1:AA:1695:G:N2	2.30	0.47
1:AA:729:G:H2'	1:AA:1775:U:O2	2.13	0.47
1:AA:1821:A:C2'	1:AA:1822:G:H5'	2.43	0.47
1:AA:1956:U:C4	1:AA:1957:C:C5	3.02	0.47
1:AA:198:C:H5'	1:AA:2244:U:OP1	2.14	0.47
1:AA:2146:C:OP2	1:AA:2146:C:H6	1.97	0.47
1:AA:2311:A:C3'	1:AA:2312:U:C6	2.97	0.47
1:AA:2646:C:H2'	1:AA:2647:U:O4'	2.14	0.47
1:AA:374:A:H3'	1:AA:375:C:C6	2.49	0.47
1:AA:372:G:N2	1:AA:401:A:OP2	2.29	0.47
1:AA:541:C:C2	1:AA:542:C:C5	3.02	0.47
1:AA:642:G:H21	1:AA:646:A:H2	1.56	0.47
1:AA:911:A:H2'	12:AP:9:TYR:CZ	2.48	0.47
2:AB:15:A:H1'	2:AB:109:G:N9	2.28	0.47
2:AB:42:C:O3'	6:AG:67:LYS:NZ	2.44	0.47
1:AA:1971:A:C4	3:AD:241:PRO:HD3	2.50	0.47
4:AE:101:ARG:CZ	4:AE:171:GLU:HB3	2.44	0.47
4:AE:29:GLY:H	4:AE:51:PHE:HE1	1.62	0.47
4:AE:35:GLN:CG	4:AE:64:LYS:NZ	2.76	0.47
7:AH:166:GLY:O	7:AH:167:GLU:O	2.32	0.47
7:AH:54:ARG:HG3	7:AH:56:SER:O	2.14	0.47
10:AN:68:GLU:HB3	10:AN:78:ARG:HH11	1.78	0.47
11:AO:79:ARG:HD3	11:AO:109:GLY:HA2	1.96	0.47
12:AP:96:VAL:HG12	12:AP:96:VAL:O	2.14	0.47
1:AA:851:U:H5"	25:AX:49:LYS:HD2	1.96	0.47
31:BA:1073:U:C2	31:BA:1074:G:C8	3.02	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:1136:U:H6	31:BA:1136:U:O5'	1.96	0.47
31:BA:168:G:C3'	31:BA:169:C:H5''	2.44	0.47
31:BA:181:G:O2'	31:BA:182:U:P	2.71	0.47
31:BA:942:G:H2'	31:BA:943:U:H6	1.79	0.47
34:BG:112:VAL:HG12	34:BG:116:GLN:NE2	2.29	0.47
35:BH:107:ARG:O	35:BH:110:LEU:N	2.30	0.47
35:BH:109:ILE:O	35:BH:109:ILE:HG22	2.12	0.47
35:BH:32:VAL:CG1	35:BH:33:VAL:H	2.26	0.47
36:BI:42:GLU:O	36:BI:44:GLY:N	2.47	0.47
37:BJ:113:GLU:CG	37:BJ:119:ARG:HG2	2.44	0.47
41:BN:34:ASP:HB2	41:BN:35:PRO:CD	2.44	0.47
47:BT:58:GLU:CB	47:BT:74:LEU:HB3	2.44	0.47
54:CA:425:G:O2'	54:CA:426:G:H5'	2.15	0.47
54:CA:976:G:H5'	54:CA:977:A:OP1	2.14	0.47
33:CF:77:ILE:HA	33:CF:84:ILE:HB	1.95	0.47
34:CG:25:ARG:HH12	34:CG:30:LYS:HB2	1.78	0.47
38:CK:116:LYS:HB2	38:CK:119:LEU:HD11	1.96	0.47
38:CK:63:LEU:HB2	38:CK:65:TYR:CE1	2.49	0.47
39:CL:88:TYR:CZ	39:CL:89:ASN:ND2	2.82	0.47
40:CM:23:ILE:HG23	40:CM:85:LEU:HD13	1.97	0.47
40:CM:6:ILE:O	40:CM:71:LEU:HD12	2.13	0.47
54:CA:1228:C:P	43:CP:108:ARG:NH2	2.87	0.47
44:CQ:12:ARG:CA	44:CQ:14:PRO:HD2	2.44	0.47
45:CR:12:ILE:C	45:CR:14:GLU:H	2.16	0.47
46:CS:26:ARG:NH2	46:CS:31:LYS:HE2	2.29	0.47
47:CT:58:GLU:O	47:CT:74:LEU:N	2.46	0.47
17:D2:59:ALA:HA	17:D2:95:LEU:O	2.15	0.47
27:D5:48:GLU:CG	27:D5:59:GLU:HG3	2.45	0.47
55:DA:1156:A:H4'	55:DA:1157:G:OP2	2.14	0.47
55:DA:1271:G:C2	55:DA:1617:C:H4'	2.49	0.47
55:DA:1791:A:H5'	55:DA:1792:G:OP2	2.14	0.47
55:DA:1930:G:H2'	55:DA:1968:G:C6	2.50	0.47
55:DA:2039:C:O2'	55:DA:2040:C:H5'	2.14	0.47
55:DA:2087:G:C2'	55:DA:2088:G:H5'	2.44	0.47
55:DA:20:C:O2'	55:DA:21:A:H5'	2.15	0.47
55:DA:2121:G:H2'	55:DA:2122:U:C6	2.49	0.47
55:DA:2205:C:O2	55:DA:2226:C:N4	2.46	0.47
55:DA:221:A:H1'	55:DA:233:A:H1'	1.97	0.47
55:DA:2292:C:H2'	55:DA:2293:C:C6	2.49	0.47
55:DA:218:A:C2	55:DA:235:U:H4'	2.48	0.47
55:DA:2414:G:H21	11:DO:67:MET:HE3	1.78	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2540:C:C2'	55:DA:2541:A:H5'	2.44	0.47
55:DA:2784:C:O2'	4:DE:37:ARG:HD2	2.14	0.47
55:DA:74:A:H5'	55:DA:75:G:O4'	2.13	0.47
55:DA:887:A:H1'	55:DA:889:C:C5	2.49	0.47
2:DB:42:C:H4'	6:DG:67:LYS:HD2	1.91	0.47
2:DB:55:U:H2'	2:DB:56:G:O4'	2.14	0.47
3:DD:263:ARG:HH11	3:DD:263:ARG:HB3	1.78	0.47
5:DF:204:ASN:C	5:DF:206:ILE:N	2.67	0.47
5:DF:54:ARG:HG2	5:DF:81:PRO:HD3	1.96	0.47
7:DH:126:PRO:CD	7:DH:127:GLU:N	2.47	0.47
56:DI:20:LEU:O	56:DI:21:LYS:C	2.51	0.47
8:DK:12:LEU:O	8:DK:13:GLY:O	2.33	0.47
58:DL:21:PRO:CB	58:DL:22:PRO:HD3	2.43	0.47
9:DM:95:PRO:O	9:DM:96:GLU:OE2	2.31	0.47
15:DR:3:ARG:HD3	15:DR:7:ILE:HD11	1.97	0.47
25:DX:7:LYS:C	25:DX:54:VAL:HG23	2.34	0.47
57:DY:70:GLU:C	57:DY:113:GLN:HB3	2.34	0.47
57:DY:132:ASP:CA	57:DY:134:LEU:HD22	2.45	0.47
57:DY:21:GLN:HE21	57:DY:22:GLY:H	0.54	0.47
57:DY:91:LYS:HA	57:DY:94:VAL:CB	2.43	0.47
16:A1:17:ILE:O	16:A1:20:LEU:HB2	2.14	0.47
30:A8:50:LEU:HD12	30:A8:54:GLU:N	2.28	0.47
1:AA:973:A:H1'	1:AA:1188:U:C5	2.49	0.47
1:AA:1278:A:C5'	13:A0:36:THR:HG22	2.44	0.47
1:AA:1344:G:C2	1:AA:1385:G:C8	3.03	0.47
1:AA:1761:C:N4	1:AA:1762:A:N1	2.62	0.47
1:AA:1897:G:H2'	1:AA:1898:U:C6	2.49	0.47
1:AA:2171:A:H2'	1:AA:2172:U:C6	2.48	0.47
1:AA:229:A:OP2	1:AA:229:A:H4'	2.14	0.47
1:AA:2329:G:H2'	1:AA:2330:G:H8	1.78	0.47
1:AA:2334:G:H4'	1:AA:2335:A:OP2	2.11	0.47
1:AA:2552:U:O2	1:AA:2554:U:H5'	2.14	0.47
1:AA:2453:A:HO2'	1:AA:2572:A:H1'	1.80	0.47
1:AA:2817:G:OP1	13:A0:42:LYS:NZ	2.44	0.47
1:AA:386:G:H3'	1:AA:388:G:N2	2.29	0.47
1:AA:390:A:H4'	1:AA:391:G:O5'	2.14	0.47
1:AA:529:A:C4'	1:AA:530:G:OP1	2.62	0.47
1:AA:607:U:OP1	5:AF:102:PRO:HA	2.14	0.47
1:AA:648:G:O2'	1:AA:649:G:H5'	2.14	0.47
1:AA:830:G:H22	1:AA:2446:G:H5'	1.79	0.47
1:AA:889:C:C5	1:AA:890:A:H1'	2.50	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:897:C:H6	1:AA:897:C:P	2.37	0.47
4:AE:116:VAL:HG23	4:AE:120:TRP:HD1	1.79	0.47
4:AE:61:ARG:N	4:AE:62:PRO:CD	2.78	0.47
6:AG:64:THR:HG23	6:AG:65:GLY:N	2.29	0.47
7:AH:45:VAL:O	7:AH:45:VAL:HG13	2.14	0.47
10:AN:87:ILE:HD13	10:AN:93:PRO:HA	1.95	0.47
1:AA:666:G:OP1	11:AO:47:ASP:O	2.32	0.47
11:AO:84:ASN:HD22	11:AO:84:ASN:N	2.12	0.47
12:AP:11:LYS:CE	12:AP:86:GLY:O	2.62	0.47
14:AQ:25:ARG:HH11	14:AQ:25:ARG:HB2	1.77	0.47
15:AR:50:ILE:HA	15:AR:99:LEU:CD1	2.44	0.47
21:AV:175:VAL:CG2	21:AV:176:PRO:HD2	2.44	0.47
24:AW:30:ARG:NH1	24:AW:30:ARG:HG3	2.29	0.47
53:B1:42:U:H5'	53:B1:42:U:H6	1.79	0.47
53:B1:53:U:HO2'	53:B1:54:U:P	2.31	0.47
31:BA:182:U:OP2	31:BA:183:G:N7	2.47	0.47
31:BA:371:G:O2'	31:BA:372:C:H5'	2.14	0.47
31:BA:501:C:H2'	31:BA:502:G:C8	2.47	0.47
31:BA:869:G:H4'	31:BA:872:A:C1'	2.45	0.47
34:BG:24:GLU:HG3	34:BG:25:ARG:H	1.78	0.47
35:BH:31:LEU:HD21	35:BH:43:LEU:CD1	2.26	0.47
39:BL:3:GLN:NE2	39:BL:20:ARG:NH1	2.62	0.47
39:BL:53:VAL:C	39:BL:55:ALA:H	2.18	0.47
43:BP:115:LYS:O	43:BP:116:THR:C	2.53	0.47
43:BP:49:THR:N	43:BP:52:GLU:OE1	2.47	0.47
46:BS:59:TRP:O	46:BS:64:ALA:HB3	2.14	0.47
47:BT:58:GLU:HB2	47:BT:74:LEU:HB3	1.97	0.47
48:BU:34:TYR:HA	48:BU:40:LEU:HD11	1.97	0.47
54:CA:1046:A:H3'	54:CA:1047:G:H8	1.79	0.47
54:CA:1160:G:H2'	54:CA:1161:C:O5'	2.14	0.47
54:CA:1366:C:O2'	40:CM:60:ARG:NH2	2.45	0.47
54:CA:1447:G:OP2	54:CA:1447:G:C8	2.68	0.47
54:CA:1449:C:H2'	54:CA:1450:U:C5'	2.25	0.47
54:CA:533:A:O2'	54:CA:534:U:P	2.72	0.47
54:CA:723:U:H5''	54:CA:724:G:OP2	2.15	0.47
54:CA:789:U:O4	54:CA:792:A:OP2	2.30	0.47
54:CA:794:A:H2'	54:CA:795:C:O4'	2.13	0.47
52:CD:24:G:O2'	52:CD:25:C:H5'	2.13	0.47
32:CE:70:PHE:HB3	32:CE:81:VAL:HG13	1.96	0.47
33:CF:6:HIS:CD2	33:CF:8:ILE:H	2.31	0.47
34:CG:163:GLU:C	34:CG:165:MET:H	2.17	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:CH:33:VAL:CG1	35:CH:112:LEU:HD12	2.43	0.47
36:CI:42:GLU:C	36:CI:44:GLY:N	2.66	0.47
37:CJ:95:ARG:HG2	37:CJ:99:LEU:CD1	2.44	0.47
38:CK:1:MET:N	38:CK:1:MET:HE2	2.29	0.47
39:CL:113:LYS:N	39:CL:113:LYS:CD	2.78	0.47
39:CL:21:PRO:HA	39:CL:59:PHE:HA	1.95	0.47
40:CM:4:ILE:HG22	40:CM:74:ILE:HD11	1.95	0.47
33:CF:30:ARG:HD2	44:CQ:38:GLY:HA3	1.96	0.47
44:CQ:3:ARG:O	44:CQ:4:LYS:C	2.52	0.47
46:CS:4:ILE:HG13	46:CS:21:VAL:HG12	1.97	0.47
51:CX:3:LYS:HB3	51:CX:14:TRP:CD1	2.49	0.47
6:DG:112:PRO:CB	26:D4:37:SER:H	2.12	0.47
28:D6:20:ASN:O	28:D6:21:TYR:CB	2.62	0.47
55:DA:1045:A:H1'	55:DA:1047:G:N3	2.29	0.47
55:DA:1838:C:H4'	55:DA:1839:G:H8	1.79	0.47
55:DA:1952:A:C6	55:DA:1953:A:C6	3.02	0.47
55:DA:2540:C:O2'	55:DA:2541:A:H5'	2.15	0.47
55:DA:2831:G:OP1	55:DA:2834:G:H4'	2.14	0.47
55:DA:2849:U:H5	15:DR:93:ARG:HH12	1.62	0.47
55:DA:2891:G:H5'	55:DA:2892:A:P	2.55	0.47
55:DA:774:A:C2'	55:DA:775:G:OP2	2.62	0.47
55:DA:802:A:H2'	55:DA:803:U:H5'	1.97	0.47
2:DB:14:U:O2'	2:DB:107:U:H4'	2.15	0.47
2:DB:82:G:N1	2:DB:95:U:O2	2.46	0.47
3:DD:35:LYS:HE2	3:DD:104:TYR:HB2	1.95	0.47
5:DF:129:PHE:C	5:DF:131:GLY:H	2.17	0.47
7:DH:19:VAL:O	7:DH:20:ALA:HB2	2.14	0.47
8:DK:78:THR:CG2	8:DK:141:LYS:HD2	2.45	0.47
9:DM:137:LYS:CG	9:DM:138:LEU:N	2.75	0.47
12:DP:51:ARG:NH1	12:DP:51:ARG:HG2	2.29	0.47
14:DQ:108:GLY:O	14:DQ:110:LEU:HG	2.13	0.47
14:DQ:18:ILE:HD13	14:DQ:87:PHE:O	2.15	0.47
15:DR:23:ARG:HA	15:DR:52:ILE:CD1	2.44	0.47
18:DS:29:LEU:HD11	18:DS:55:ALA:HB2	1.96	0.47
19:DT:12:VAL:HG11	19:DT:27:THR:HG23	1.96	0.47
19:DT:57:LEU:CD1	19:DT:57:LEU:N	2.77	0.47
21:DV:111:VAL:CG2	21:DV:145:GLU:HA	2.45	0.47
21:DV:178:GLU:HG3	21:DV:180:VAL:HA	1.96	0.47
12:DP:136:ALA:HB1	21:DV:52:SER:HB2	1.96	0.47
25:DX:36:VAL:O	25:DX:37:LEU:HD23	2.15	0.47
25:DX:7:LYS:HE2	25:DX:32:GLN:CA	2.44	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:74:LEU:HD13	57:DY:75:GLN:CG	2.43	0.47
23:DZ:21:ARG:HG3	23:DZ:35:THR:HG23	1.96	0.47
16:A1:50:ARG:HH11	17:A2:72:VAL:CG1	2.28	0.47
22:A3:74:ARG:NH1	22:A3:74:ARG:HG3	2.26	0.47
26:A4:65:ASP:O	26:A4:66:SER:C	2.53	0.47
30:A8:6:THR:O	30:A8:7:HIS:CB	2.63	0.47
1:AA:1053:C:H3'	1:AA:1054:A:C5'	2.43	0.47
1:AA:1085:A:O2'	1:AA:1086:A:P	2.72	0.47
1:AA:330:A:C2	1:AA:1210:A:H2'	2.49	0.47
1:AA:1406:U:C3'	1:AA:1407:C:H6	2.25	0.47
1:AA:1925:C:H42	1:AA:1929:G:H21	1.62	0.47
1:AA:223:A:O2'	1:AA:420:C:O2	2.31	0.47
1:AA:2540:C:O2	1:AA:2540:C:H2'	2.13	0.47
1:AA:2540:C:O2'	1:AA:2740:A:N3	2.32	0.47
1:AA:613:U:O4'	1:AA:613:U:O2	2.32	0.47
1:AA:789:A:H4'	1:AA:790:C:OP1	2.14	0.47
1:AA:954:G:N2	1:AA:955:C:H1'	2.30	0.47
2:AB:40:U:H2'	2:AB:41:U:OP1	2.14	0.47
4:AE:58:ARG:NH2	4:AE:58:ARG:HA	2.27	0.47
1:AA:2749:A:H1'	7:AH:63:SER:OG	2.14	0.47
9:AM:28:THR:HA	9:AM:106:MET:CE	2.44	0.47
9:AM:120:LEU:CD2	9:AM:122:VAL:HG23	2.38	0.47
10:AN:86:ILE:O	10:AN:87:ILE:HD13	2.15	0.47
11:AO:114:ILE:HG21	11:AO:125:VAL:HG21	1.95	0.47
12:AP:104:PHE:O	12:AP:105:GLU:O	2.32	0.47
14:AQ:73:LEU:C	14:AQ:73:LEU:HD13	2.34	0.47
1:AA:1754:C:P	15:AR:96:ARG:HH12	2.38	0.47
19:AT:18:TYR:O	19:AT:21:PHE:HB2	2.14	0.47
20:AU:6:HIS:HE2	20:AU:72:VAL:CG2	2.27	0.47
23:AZ:96:LYS:O	23:AZ:97:LEU:C	2.53	0.47
31:BA:1024:G:C3'	31:BA:1025:U:H5''	2.44	0.47
31:BA:109:A:C6	31:BA:326:G:C6	3.02	0.47
31:BA:1126:U:N3	31:BA:1281:U:O4'	2.47	0.47
31:BA:1320:C:C2	49:BV:72:GLY:HA3	2.48	0.47
31:BA:178:C:O2'	31:BA:179:A:H5'	2.14	0.47
31:BA:359:U:H2'	31:BA:360:A:H8	1.77	0.47
31:BA:46:G:O2'	31:BA:365:U:H1'	2.15	0.47
31:BA:583:A:H2'	31:BA:584:G:O4'	2.15	0.47
31:BA:663:A:H5'	31:BA:836:G:OP1	2.15	0.47
31:BA:959:A:C3'	31:BA:960:U:H4'	2.44	0.47
31:BA:965:A:C5'	31:BA:966:G:OP1	2.61	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:BB:9:A:C2	52:BB:11:C:N4	2.80	0.47
34:BG:190:ASP:O	34:BG:192:GLU:N	2.48	0.47
38:BK:29:SER:HB3	38:BK:32:LYS:HG3	1.97	0.47
39:BL:14:VAL:C	39:BL:65:VAL:HG23	2.35	0.47
40:BM:33:GLN:H	40:BM:75:ILE:HG12	1.79	0.47
40:BM:75:ILE:CG1	40:BM:76:ASN:N	2.77	0.47
43:BP:15:VAL:CG2	43:BP:41:PRO:HA	2.44	0.47
44:BQ:22:THR:O	44:BQ:23:ARG:HB2	2.13	0.47
48:BU:21:LYS:HB3	48:BU:22:VAL:H	1.45	0.47
49:BV:5:LEU:HG	49:BV:9:VAL:CA	2.36	0.47
54:CA:1273:G:H3'	54:CA:1274:G:H8	1.79	0.47
54:CA:1298:C:H1'	54:CA:1299:A:C6	2.49	0.47
54:CA:1316:G:H4'	44:CQ:18:VAL:CG1	2.44	0.47
54:CA:1336:C:C2'	54:CA:1337:G:OP2	2.63	0.47
54:CA:1346:A:N1	54:CA:1374:A:H5''	2.30	0.47
54:CA:384:G:H2'	54:CA:385:C:C6	2.49	0.47
54:CA:489:C:H2'	54:CA:490:G:H8	1.80	0.47
33:CF:34:LEU:O	33:CF:38:ARG:HG3	2.15	0.47
33:CF:35:GLU:OE2	33:CF:95:THR:HG23	2.15	0.47
34:CG:160:GLN:O	34:CG:163:GLU:HB3	2.14	0.47
34:CG:25:ARG:C	34:CG:27:TYR:H	2.11	0.47
36:CI:21:LEU:O	36:CI:24:GLU:N	2.47	0.47
36:CI:41:GLU:O	36:CI:41:GLU:HG2	2.14	0.47
40:CM:32:ALA:CB	40:CM:76:ASN:HB2	2.45	0.47
40:CM:94:VAL:HG12	40:CM:95:GLU:N	2.30	0.47
42:CO:78:GLN:O	42:CO:80:HIS:N	2.47	0.47
43:CP:16:ASP:N	43:CP:16:ASP:OD2	2.46	0.47
46:CS:51:VAL:HG12	46:CS:52:ASP:N	2.30	0.47
48:CU:26:LEU:HD22	48:CU:42:ARG:NE	2.29	0.47
48:CU:58:LEU:N	48:CU:58:LEU:HD12	2.29	0.47
27:D5:2:ALA:O	27:D5:3:LYS:HD2	2.13	0.47
28:D6:15:GLU:OE2	28:D6:44:ARG:NH1	2.47	0.47
55:DA:1044:G:O2'	55:DA:1045:A:H5''	2.14	0.47
55:DA:1056:G:C2	55:DA:1103:A:N7	2.81	0.47
55:DA:1242:A:C5'	55:DA:1243:G:OP2	2.63	0.47
55:DA:1360:A:C8	55:DA:1361:G:C8	3.02	0.47
55:DA:1416:G:C4	55:DA:1417:C:C5	3.03	0.47
55:DA:1427:A:H4'	55:DA:1428:C:O4'	2.14	0.47
55:DA:1503:U:H2'	55:DA:1504:C:H6	1.77	0.47
55:DA:1964:G:O2'	55:DA:1967:C:OP2	2.29	0.47
55:DA:2126:A:H4'	55:DA:2127:G:O5'	2.14	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2311:A:H3'	55:DA:2312:U:C6	2.49	0.47
55:DA:2408:U:H2'	55:DA:2409:G:C8	2.49	0.47
55:DA:2728:U:H2'	55:DA:2729:G:C8	2.50	0.47
55:DA:2762:G:H2'	55:DA:2763:G:H5'	1.97	0.47
55:DA:565:C:H4'	55:DA:1253:A:N6	2.30	0.47
55:DA:719:C:H2'	55:DA:720:C:C6	2.48	0.47
55:DA:851:U:C4'	25:DX:46:ASN:ND2	2.77	0.47
55:DA:888:C:O5'	55:DA:889:C:C5	2.67	0.47
3:DD:166:GLN:HE21	3:DD:166:GLN:HA	1.80	0.47
5:DF:107:LYS:O	5:DF:108:LYS:C	2.53	0.47
5:DF:178:PRO:HB2	5:DF:201:VAL:CG1	2.42	0.47
6:DG:122:PRO:C	6:DG:123:ASN:HD22	2.18	0.47
6:DG:88:ILE:O	6:DG:88:ILE:CG2	2.62	0.47
7:DH:164:TYR:O	7:DH:166:GLY:N	2.47	0.47
56:DI:26:ALA:O	56:DI:28:LYS:O	2.32	0.47
58:DL:19:PRO:O	58:DL:20:ALA:HB3	2.13	0.47
58:DL:34:ILE:CG1	58:DL:38:VAL:HG22	2.45	0.47
9:DM:35:ARG:O	9:DM:37:LYS:N	2.48	0.47
11:DO:36:LYS:HG3	11:DO:36:LYS:HZ3	1.51	0.47
55:DA:2393:A:H5'	11:DO:62:LEU:HB2	1.97	0.47
4:DE:9:VAL:HG11	15:DR:7:ILE:HB	1.95	0.47
21:DV:62:PRO:O	21:DV:63:ASP:HB3	2.14	0.47
57:DY:50:ARG:CD	57:DY:51:LEU:N	2.70	0.47
57:DY:6:ASN:C	57:DY:7:VAL:CG1	2.82	0.47
57:DY:97:ALA:C	57:DY:98:LYS:O	2.53	0.47
26:A4:56:VAL:O	26:A4:57:GLU:HB2	2.15	0.47
1:AA:1141:U:H5'	9:AM:25:ARG:NH2	2.30	0.47
1:AA:1165:U:H2'	1:AA:1166:C:H6	1.72	0.47
1:AA:51:G:N2	1:AA:119:A:N3	2.63	0.47
1:AA:27:G:H22	1:AA:512:G:H2'	1.79	0.47
1:AA:727:A:C2	3:AD:9:TYR:CD2	3.03	0.47
2:AB:14:U:O2'	2:AB:107:U:H1'	2.14	0.47
4:AE:111:ARG:HG2	13:A0:2:ARG:HH22	1.79	0.47
6:AG:151:ALA:HB3	6:AG:153:ARG:HH11	1.79	0.47
7:AH:13:LYS:HA	7:AH:13:LYS:HZ3	1.79	0.47
7:AH:4:ILE:HG13	7:AH:5:GLY:N	2.28	0.47
10:AN:16:ALA:HA	10:AN:46:ALA:HA	1.96	0.47
11:AO:114:ILE:HG12	11:AO:130:PHE:CD1	2.49	0.47
15:AR:98:LYS:N	15:AR:98:LYS:HD2	2.30	0.47
18:AS:70:TYR:CD2	18:AS:70:TYR:N	2.83	0.47
18:AS:13:SER:HA	18:AS:99:ARG:HB2	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:AU:20:TYR:N	20:AU:20:TYR:CD1	2.83	0.47
21:AV:44:PHE:CE2	21:AV:86:VAL:HG11	2.49	0.47
23:AZ:92:LYS:HZ1	23:AZ:97:LEU:CG	2.20	0.47
53:B1:51:U:C3'	53:B1:51:U:C6	2.98	0.47
31:BA:1342:C:H1'	39:BL:124:GLN:HE21	1.76	0.47
31:BA:1342:C:O2'	31:BA:1343:G:H5'	2.15	0.47
31:BA:1365:G:H2'	31:BA:1366:C:C6	2.48	0.47
31:BA:1535:C:O2'	31:BA:1536:C:H5'	2.14	0.47
31:BA:247:G:O2'	31:BA:248:C:H5'	2.14	0.47
31:BA:508:C:H4'	31:BA:509:A:O5'	2.14	0.47
31:BA:707:C:O2'	31:BA:708:C:H5'	2.14	0.47
52:BC:42:C:C2'	52:BC:43:C:H5''	2.44	0.47
33:BF:64:VAL:HG12	33:BF:66:VAL:HG23	1.96	0.47
34:BG:139:ARG:NH1	34:BG:139:ARG:HG3	2.22	0.47
34:BG:19:LEU:C	34:BG:21:LEU:HG	2.35	0.47
35:BH:78:HIS:CG	38:BK:104:ARG:HD2	2.50	0.47
31:BA:718:G:C5'	41:BN:117:ASN:OD1	2.58	0.47
42:BO:70:ILE:CD1	42:BO:77:LEU:HD12	2.45	0.47
43:BP:49:THR:HB	43:BP:52:GLU:CG	2.43	0.47
43:BP:97:PRO:HA	43:BP:110:ARG:CD	2.44	0.47
45:BR:66:LEU:O	45:BR:69:TYR:HB3	2.14	0.47
48:BU:45:SER:C	48:BU:47:THR:H	2.18	0.47
50:BW:26:ASN:HD22	50:BW:27:LYS:H	1.62	0.47
54:CA:1003:G:H5'	54:CA:1003:G:H8	1.80	0.47
54:CA:1102:A:C6	54:CA:1103:C:N4	2.83	0.47
54:CA:1175:G:H2'	54:CA:1176:A:H8	1.73	0.47
54:CA:129(A):G:H1'	54:CA:190:G:C5'	2.45	0.47
54:CA:1313:U:OP2	49:CV:6:LYS:N	2.47	0.47
54:CA:143:A:H2	54:CA:220:G:H1	1.62	0.47
54:CA:478:A:O2'	54:CA:479:C:H5'	2.15	0.47
54:CA:715:A:H2'	54:CA:716:A:C8	2.50	0.47
54:CA:736:C:H2'	54:CA:737:A:C8	2.50	0.47
54:CA:788:U:H3	54:CA:795:C:N4	2.11	0.47
54:CA:953:G:H2'	54:CA:954:G:O4'	2.14	0.47
52:CB:22:G:O2'	52:CB:23:A:H5'	2.14	0.47
52:CB:25:C:C4	52:CB:26:A:C2	3.02	0.47
52:CB:2:C:H2'	52:CB:3:C:H6	1.78	0.47
52:CC:21:A:N6	52:CC:46:G:H2'	2.30	0.47
33:CF:5:ILE:CD1	33:CF:5:ILE:H	2.27	0.47
35:CH:79:GLU:HB3	35:CH:92:LYS:HG3	1.95	0.47
37:CJ:87:VAL:HG21	37:CJ:154:TYR:HB2	1.96	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:CK:53:VAL:O	38:CK:54:ASP:HB2	2.14	0.47
54:CA:1346:A:H5'	39:CL:120:ARG:HH12	1.80	0.47
54:CA:1128:C:C5'	39:CL:16:ARG:HH22	2.16	0.47
54:CA:1151:A:H5'	40:CM:40:LEU:O	2.15	0.47
40:CM:11:PHE:HA	40:CM:66:ARG:O	2.15	0.47
43:CP:48:LEU:N	43:CP:48:LEU:HD23	2.28	0.47
48:CU:50:ILE:N	48:CU:50:ILE:CD1	2.76	0.47
50:CW:94:ALA:O	50:CW:95:ALA:CB	2.63	0.47
54:CA:1244:C:OP2	51:CX:9:ARG:HB2	2.15	0.47
13:D0:94:TYR:C	13:D0:117:VAL:HG12	2.35	0.47
16:D1:92:ARG:HD2	16:D1:95:LEU:HD12	1.96	0.47
17:D2:48:GLY:O	17:D2:49:THR:O	2.32	0.47
55:DA:242:G:OP2	30:D8:3:LYS:HE3	2.14	0.47
55:DA:1082:U:OP2	57:DY:45:LYS:HG2	2.15	0.47
55:DA:1278:A:H2'	55:DA:1279:G:H8	1.79	0.47
55:DA:1533:C:C2'	55:DA:1534:G:N7	2.65	0.47
55:DA:1923:U:H2'	55:DA:1924:C:C6	2.49	0.47
55:DA:212:G:C2'	55:DA:213:A:H5'	2.45	0.47
55:DA:21:A:O2'	55:DA:22:C:H5'	2.15	0.47
55:DA:235:U:H2'	55:DA:236:C:C6	2.49	0.47
55:DA:234:C:H2'	55:DA:235:U:O4'	2.15	0.47
55:DA:1629:U:O2	55:DA:2698:U:H5''	2.14	0.47
55:DA:469:G:H2'	55:DA:470:A:H5''	1.97	0.47
55:DA:74:A:C5'	55:DA:75:G:O4'	2.62	0.47
55:DA:916:G:C2'	55:DA:917:A:H5''	2.44	0.47
4:DE:109:LYS:HE2	4:DE:191:PRO:HA	1.97	0.47
5:DF:164:ARG:HG3	5:DF:175:THR:OG1	2.15	0.47
6:DG:122:PRO:HB3	6:DG:180:PHE:HD2	1.79	0.47
6:DG:96:ARG:O	6:DG:98:ARG:N	2.48	0.47
7:DH:127:GLU:CG	7:DH:128:PRO:HD2	2.45	0.47
7:DH:22:GLY:O	7:DH:37:VAL:N	2.48	0.47
8:DK:27:ARG:HB2	23:DZ:71:TYR:CZ	2.49	0.47
58:DL:58:THR:HB	58:DL:66:THR:HG21	1.95	0.47
9:DM:131:GLN:NE2	9:DM:132:ALA:N	2.62	0.47
10:DN:8:LEU:HB2	10:DN:19:ILE:HD11	1.96	0.47
14:DQ:58:LEU:N	14:DQ:58:LEU:CD2	2.78	0.47
15:DR:29:ARG:HH12	15:DR:89:VAL:HG11	1.79	0.47
18:DS:9:TYR:N	18:DS:102:HIS:HD2	2.03	0.47
57:DY:4:LYS:C	57:DY:5:ARG:HG3	2.35	0.47
57:DY:88:ALA:C	57:DY:92:THR:CB	2.83	0.47
13:A0:103:ARG:HD3	13:A0:110:PRO:HA	1.95	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:A1:102:GLU:H	16:A1:103:PRO:CD	2.28	0.47
1:AA:1151:G:H5''	16:A1:81:HIS:NE2	2.30	0.47
26:A4:16:CYS:C	26:A4:18:CYS:H	2.18	0.47
30:A8:37:SER:O	30:A8:38:GLY:C	2.53	0.47
30:A8:64:TYR:HB2	30:A8:65:GLU:H	1.36	0.47
1:AA:1011:G:OP1	16:A1:75:ASN:HB3	2.15	0.47
1:AA:1202:C:N4	1:AA:1203:G:C6	2.83	0.47
1:AA:1219:G:OP2	16:A1:19:LYS:NZ	2.48	0.47
1:AA:1462:C:H2'	1:AA:1463:C:H6	1.79	0.47
1:AA:1937:A:HO2'	1:AA:1938:A:P	2.38	0.47
1:AA:2037:G:H2'	1:AA:2038:G:H8	1.79	0.47
1:AA:2169:A:N3	1:AA:2169:A:H2'	2.30	0.47
1:AA:2453:A:O2'	1:AA:2572:A:H1'	2.15	0.47
1:AA:2672:G:H2'	1:AA:2673:G:H5''	1.97	0.47
1:AA:2850:A:H5'	1:AA:2868:A:C2	2.49	0.47
1:AA:345:A:O2'	1:AA:346:A:P	2.72	0.47
1:AA:395:U:H2'	1:AA:396:G:N7	2.30	0.47
1:AA:458:G:H2'	1:AA:459:U:OP2	2.11	0.47
5:AF:110:LEU:O	5:AF:113:ALA:HB3	2.14	0.47
5:AF:24:LEU:O	5:AF:25:PRO:O	2.33	0.47
6:AG:7:LEU:HD22	6:AG:100:TRP:CZ3	2.49	0.47
6:AG:47:LYS:HG3	6:AG:82:LEU:HD23	1.96	0.47
1:AA:270(Q):C:H5'	8:AK:45:LYS:CE	2.44	0.47
9:AM:56:ASN:HB3	9:AM:126:PRO:N	2.30	0.47
12:AP:105:GLU:HG3	12:AP:105:GLU:O	2.14	0.47
12:AP:37:LEU:HD21	12:AP:130:LYS:HD2	1.96	0.47
14:AQ:103:GLU:O	14:AQ:106:ARG:HG3	2.15	0.47
14:AQ:83:LYS:HE2	14:AQ:84:GLN:HG3	1.96	0.47
15:AR:108:ARG:O	15:AR:108:ARG:HG2	2.15	0.47
21:AV:67:LEU:HD21	21:AV:90:VAL:HG13	1.96	0.47
21:AV:73:GLN:HB3	21:AV:87:ASP:OD2	2.15	0.47
23:AZ:15:ALA:HB2	23:AZ:42:GLN:OE1	2.14	0.47
23:AZ:82:LEU:H	23:AZ:82:LEU:CD2	2.21	0.47
31:BA:1004:A:O5'	31:BA:1025:U:C4	2.66	0.47
31:BA:1053:G:N7	31:BA:1199:U:H3'	2.29	0.47
31:BA:126:G:H4'	31:BA:634:C:O2	2.15	0.47
31:BA:1297:C:H6	31:BA:1297:C:OP2	1.97	0.47
31:BA:791:G:C5	31:BA:792:A:N7	2.82	0.47
52:BB:5:G:H2'	52:BB:6:G:C8	2.46	0.47
33:BF:76:VAL:CG2	33:BF:77:ILE:H	2.28	0.47
35:BH:129:ILE:O	35:BH:132:ALA:HB3	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:BH:144:THR:O	35:BH:147:ASP:OD2	2.33	0.47
35:BH:80:ILE:CD1	35:BH:82:VAL:HG23	2.44	0.47
42:BO:35:GLY:HA3	42:BO:58:VAL:CG1	2.45	0.47
43:BP:108:ARG:NH1	43:BP:108:ARG:HG3	2.30	0.47
44:BQ:48:ALA:CA	44:BQ:53:LEU:HD12	2.44	0.47
46:BS:75:ARG:HG3	46:BS:80:PHE:CD1	2.50	0.47
53:C1:56:U:O2'	53:C1:57:U:O5'	2.33	0.47
54:CA:999:U:H2'	54:CA:1000:A:O4'	2.15	0.47
54:CA:184:G:O4'	54:CA:224:C:H4'	2.15	0.47
54:CA:41:G:H2'	54:CA:42:G:H8	1.80	0.47
54:CA:630:G:H8	54:CA:630:G:C3'	2.16	0.47
54:CA:872:A:N1	54:CA:874:G:C6	2.82	0.47
54:CA:9:G:OP2	54:CA:9:G:O4'	2.33	0.47
32:CE:41:ILE:HD12	32:CE:41:ILE:N	2.29	0.47
32:CE:33:TYR:CD2	32:CE:43:ASP:HB2	2.48	0.47
32:CE:48:MET:C	32:CE:50:GLU:N	2.66	0.47
32:CE:97:TRP:CH2	32:CE:176:GLU:OE2	2.68	0.47
33:CF:153:VAL:HG13	33:CF:196:LEU:HD12	1.95	0.47
54:CA:1055:A:O2'	33:CF:161:GLU:OE1	2.33	0.47
34:CG:8:VAL:O	34:CG:11:LEU:N	2.45	0.47
35:CH:78:HIS:CB	38:CK:104:ARG:HG2	2.45	0.47
36:CI:99:ALA:O	48:CU:28:GLU:HA	2.14	0.47
37:CJ:122:HIS:O	37:CJ:125:MET:HB2	2.14	0.47
37:CJ:45:ASP:O	37:CJ:49:ILE:HG12	2.15	0.47
46:CS:21:VAL:O	46:CS:21:VAL:HG23	2.15	0.47
54:CA:275:G:H5'	47:CT:14:LYS:HB3	1.95	0.47
50:CW:45:GLN:HB2	50:CW:91:LEU:HD22	1.96	0.47
22:D3:14:ARG:O	22:D3:15:ASP:HB2	2.13	0.47
55:DA:1011:G:P	16:D1:77:SER:HG	2.38	0.47
55:DA:1458:C:H5"	55:DA:1459:G:O5'	2.15	0.47
55:DA:1978:A:H2'	55:DA:1979:C:C6	2.50	0.47
55:DA:2219:G:H2'	55:DA:2224:G:C5'	2.45	0.47
55:DA:2241:A:H2'	55:DA:2242:G:C8	2.49	0.47
55:DA:2408:U:O5'	55:DA:2408:U:H6	1.98	0.47
55:DA:2618:G:H2'	55:DA:2619:C:C6	2.50	0.47
55:DA:2779:U:O2'	55:DA:2781:A:N7	2.42	0.47
55:DA:33:U:O2'	55:DA:34:C:P	2.73	0.47
55:DA:528:A:C2	55:DA:2043:C:H4'	2.49	0.47
55:DA:77:C:H5"	24:DW:10:LEU:HD11	1.97	0.47
55:DA:847:U:H5	55:DA:933:A:N1	2.12	0.47
2:DB:35:U:H2'	2:DB:36:C:C6	2.50	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:DD:66:ASP:OD1	3:DD:68:LYS:O	2.33	0.47
7:DH:137:ASP:O	7:DH:138:LYS:HB2	2.15	0.47
56:DI:10:GLU:O	56:DI:14:GLN:CA	2.63	0.47
58:DL:69:THR:O	58:DL:70:LYS:CG	2.63	0.47
10:DN:31:LYS:HB3	10:DN:32:TYR:CD1	2.50	0.47
10:DN:79:PHE:HE2	10:DN:101:PRO:HB2	1.79	0.47
11:DO:79:ARG:HD3	11:DO:110:TYR:HE1	1.77	0.47
11:DO:88:LEU:C	11:DO:88:LEU:HD23	2.35	0.47
14:DQ:3:ARG:CG	14:DQ:4:LEU:N	2.75	0.47
20:DU:91:GLU:HG3	20:DU:92:ASN:N	2.30	0.47
21:DV:178:GLU:O	21:DV:179:ASP:CB	2.61	0.47
21:DV:192:ALA:CA	21:DV:193:GLU:OE1	2.62	0.47
57:DY:116:ILE:O	57:DY:117:LEU:CB	2.59	0.47
57:DY:59:ILE:C	57:DY:61:LEU:N	2.68	0.47
16:A1:28:ARG:NH1	16:A1:38:THR:OG1	2.47	0.47
1:AA:2330:G:H1'	22:A3:41:ARG:O	2.15	0.47
1:AA:1061:U:O2'	1:AA:1070:A:O4'	2.33	0.47
1:AA:1498:C:O4'	1:AA:1577:C:H4'	2.15	0.47
1:AA:1652:A:H62	13:A0:11:ASN:ND2	2.03	0.47
1:AA:1827:C:H1'	1:AA:1970:A:O2'	2.15	0.47
1:AA:2147:G:C8	1:AA:2147:G:C3'	2.97	0.47
1:AA:233:A:O2'	1:AA:234:C:H5'	2.15	0.47
1:AA:2387:U:H1'	22:A3:41:ARG:HD2	1.96	0.47
1:AA:2407:G:O5'	1:AA:2407:G:C8	2.68	0.47
1:AA:2418:A:OP2	30:A8:29:LYS:CE	2.63	0.47
1:AA:2468:G:OP1	12:AP:119:ARG:NH2	2.45	0.47
1:AA:2517:C:C2	1:AA:2542:A:N6	2.83	0.47
1:AA:2758:A:C2	7:AH:67:LEU:HD21	2.49	0.47
1:AA:2861:G:O2'	1:AA:2862:G:H5'	2.15	0.47
1:AA:322:A:H3'	5:AF:169:ASN:ND2	2.28	0.47
1:AA:332:A:O2'	1:AA:334:C:OP2	2.26	0.47
1:AA:370:G:O2'	1:AA:371:A:OP1	2.27	0.47
1:AA:528:A:N1	1:AA:2042:A:H2'	2.28	0.47
1:AA:588:U:H2'	1:AA:589:C:H6	1.79	0.47
1:AA:612:G:C5'	1:AA:612:G:H8	2.19	0.47
1:AA:758:C:H2'	1:AA:758:C:O2	2.15	0.47
1:AA:863:A:H2'	1:AA:864:G:H8	1.80	0.47
2:AB:14:U:O2'	2:AB:107:U:H4'	2.15	0.47
2:AB:2:C:H2'	2:AB:3:C:H6	1.80	0.47
2:AB:58:A:H5'	2:AB:59:A:OP2	2.15	0.47
2:AB:93:C:O2'	2:AB:94:C:H5'	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:AH:67:LEU:O	7:AH:71:LEU:HB2	2.15	0.47
12:AP:89:ASN:HD22	12:AP:89:ASN:N	2.12	0.47
14:AQ:62:LYS:O	14:AQ:65:VAL:HB	2.15	0.47
20:AU:75:ILE:CB	20:AU:80:GLY:H	2.28	0.47
20:AU:75:ILE:CG1	20:AU:80:GLY:H	2.28	0.47
21:AV:178:GLU:C	21:AV:180:VAL:H	2.17	0.47
53:B1:42:U:H4'	53:B1:43:U:OP1	2.15	0.47
31:BA:1143:G:H2'	31:BA:1144:G:C8	2.50	0.47
31:BA:1157:A:N6	31:BA:1180:A:N7	2.62	0.47
31:BA:1370:G:O2'	31:BA:1371:G:H5'	2.14	0.47
31:BA:1381:U:H2'	31:BA:1381:U:O2	2.15	0.47
31:BA:1502:A:O2'	31:BA:1503:A:OP1	2.23	0.47
31:BA:209:U:HO2'	31:BA:210:U:P	2.35	0.47
31:BA:208:U:O2'	31:BA:209:U:OP1	2.28	0.47
31:BA:542:G:H5'	34:BG:41:GLY:CA	2.45	0.47
31:BA:913:A:H1'	31:BA:914:A:O4'	2.15	0.47
52:BC:39:U:C2	52:BC:40:C:C5	3.03	0.47
33:BF:148:GLY:CA	33:BF:203:PHE:HB3	2.43	0.47
34:BG:119:GLN:OE1	34:BG:123:HIS:CD2	2.68	0.47
34:BG:200:GLU:O	34:BG:203:VAL:N	2.48	0.47
34:BG:9:CYS:C	34:BG:11:LEU:H	2.17	0.47
38:BK:31:PHE:CE2	38:BK:35:ILE:HD11	2.49	0.47
39:BL:28:VAL:HG22	39:BL:63:ILE:O	2.15	0.47
40:BM:97:GLU:C	40:BM:98:ILE:HD12	2.34	0.47
31:BA:1455:G:OP1	50:BW:35:THR:HG21	2.15	0.47
53:C1:30:C:C6	53:C1:30:C:C3'	2.97	0.47
54:CA:1139:G:N2	54:CA:1143:G:O6	2.47	0.47
54:CA:328:C:C2'	54:CA:328:C:O2	2.63	0.47
54:CA:392:G:H2'	54:CA:393:A:C8	2.50	0.47
54:CA:828:A:H5''	54:CA:859:A:C2	2.50	0.47
54:CA:957:U:H1'	54:CA:960:U:C6	2.49	0.47
52:CD:9:A:H5''	52:CD:10:G:OP2	2.15	0.47
32:CE:212:GLN:CG	32:CE:235:SER:HB2	2.40	0.47
32:CE:60:ASP:HB3	32:CE:64:ARG:CZ	2.44	0.47
33:CF:91:LEU:CD1	33:CF:101:LEU:HD12	2.43	0.47
34:CG:131:ARG:NH1	34:CG:131:ARG:HG3	2.28	0.47
40:CM:49:VAL:CG1	40:CM:50:ILE:N	2.78	0.47
41:CN:78:GLN:O	41:CN:103:LEU:HA	2.15	0.47
42:CO:82:VAL:HG23	42:CO:106:ASP:OD2	2.14	0.47
42:CO:18:VAL:O	42:CO:19:ARG:HB2	2.14	0.47
42:CO:90:VAL:O	42:CO:91:LYS:C	2.53	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:CT:80:GLY:O	47:CT:82:MET:HG2	2.15	0.47
16:D1:44:ASN:ND2	16:D1:44:ASN:N	2.59	0.47
16:D1:92:ARG:HD2	17:D2:11:GLN:HE21	1.70	0.47
55:DA:857:C:H1'	22:D3:26:TYR:HE2	1.79	0.47
55:DA:2420:C:OP1	30:D8:34:TRP:N	2.46	0.47
55:DA:1074:G:H2'	55:DA:1075:C:C6	2.49	0.47
55:DA:1710:C:H2'	55:DA:1711:C:C6	2.50	0.47
55:DA:1906:G:C4	55:DA:1929:G:N2	2.83	0.47
55:DA:2273:A:C2'	55:DA:2274:A:H5'	2.44	0.47
55:DA:2477:C:O5'	55:DA:2477:C:H6	1.97	0.47
55:DA:2737:G:H2'	55:DA:2738:A:C8	2.50	0.47
55:DA:321:G:O2'	55:DA:340:A:O2'	2.27	0.47
55:DA:448:U:C4	55:DA:583:G:H1'	2.50	0.47
55:DA:471:A:O5'	55:DA:471:A:H8	1.98	0.47
55:DA:510:C:O2'	55:DA:511:U:H5'	2.15	0.47
55:DA:633:A:H2'	55:DA:634:C:H5'	1.95	0.47
55:DA:684:G:C2	55:DA:774:A:C2	3.03	0.47
3:DD:94:LEU:C	3:DD:94:LEU:CD1	2.81	0.47
4:DE:179:GLU:HB3	4:DE:181:LEU:HD23	1.95	0.47
55:DA:2811:G:OP1	4:DE:61:ARG:HG2	2.14	0.47
6:DG:108:ASN:HA	26:D4:38:LYS:CG	2.44	0.47
6:DG:75:LYS:HD2	6:DG:77:ILE:HD11	1.96	0.47
7:DH:16:SER:O	7:DH:17:VAL:HG23	2.15	0.47
7:DH:2:SER:O	7:DH:3:ARG:C	2.53	0.47
7:DH:89:ILE:HD13	7:DH:90:LYS:H	1.78	0.47
58:DL:78:ILE:HG21	58:DL:131:ALA:HB2	1.97	0.47
9:DM:7:LYS:CD	9:DM:7:LYS:H	2.23	0.47
10:DN:12:ASP:C	10:DN:12:ASP:OD2	2.53	0.47
10:DN:5:GLN:O	10:DN:20:MET:CE	2.63	0.47
12:DP:6:ARG:O	12:DP:7:MET:HB2	2.15	0.47
19:DT:47:PHE:O	19:DT:48:LYS:C	2.53	0.47
12:DP:140:ALA:CB	21:DV:53:ILE:HD11	2.43	0.47
57:DY:38:HIS:C	57:DY:40:LEU:N	2.66	0.47
57:DY:41:ARG:O	58:DL:116:ASN:O	2.33	0.47
57:DY:89:ALA:CA	56:DJ:15:ALA:HB2	2.44	0.47
23:DZ:15:ALA:O	23:DZ:40:ARG:HG3	2.14	0.47
13:A0:63:ARG:HH11	13:A0:63:ARG:CB	2.24	0.47
16:A1:75:ASN:HB2	16:A1:78:THR:OG1	2.15	0.47
17:A2:38:LEU:HD21	17:A2:57:VAL:HG13	1.97	0.47
17:A2:44:LYS:HG2	17:A2:45:THR:HG23	1.97	0.47
30:A8:36:LYS:HB2	30:A8:41:ILE:HG13	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:A8:61:LEU:O	30:A8:62:LEU:C	2.53	0.47
1:AA:1171:G:H1'	1:AA:1173:G:O4'	2.14	0.47
1:AA:1291:C:C5'	1:AA:1536:A:H5''	2.44	0.47
1:AA:1297:C:H2'	1:AA:1298:C:H6	1.80	0.47
1:AA:1517:G:H2'	1:AA:1518:C:C6	2.50	0.47
1:AA:1819:A:OP1	3:AD:161:THR:HG21	2.13	0.47
1:AA:1930:G:H2'	1:AA:1968:G:N1	2.30	0.47
1:AA:2113:U:H3'	1:AA:2114:A:H5'	1.96	0.47
1:AA:2133:G:O2'	1:AA:2158:A:N1	2.45	0.47
1:AA:2753:A:C3'	1:AA:2754:U:H5''	2.44	0.47
1:AA:2851:A:H2'	1:AA:2852:G:H8	1.80	0.47
1:AA:2852:G:O2'	1:AA:2853:C:H5'	2.14	0.47
1:AA:457:A:O2'	1:AA:458:G:OP2	2.30	0.47
1:AA:640:C:O2'	1:AA:641:C:H5'	2.15	0.47
1:AA:920:G:H2'	1:AA:921:G:H8	1.80	0.47
1:AA:957:A:N6	1:AA:959:A:C2	2.83	0.47
1:AA:96:G:O5'	24:AW:48:HIS:ND1	2.45	0.47
1:AA:988:A:H3'	25:AX:11:SER:OG	2.14	0.47
2:AB:14:U:O2'	2:AB:107:U:C1'	2.62	0.47
4:AE:93:VAL:HG21	4:AE:180:ASN:HA	1.97	0.47
4:AE:93:VAL:C	4:AE:95:ILE:H	2.17	0.47
8:AK:100:ALA:O	8:AK:102:SER:N	2.48	0.47
8:AK:52:ARG:HG2	8:AK:52:ARG:HH11	1.80	0.47
12:AP:21:THR:CG2	12:AP:100:GLY:HA3	2.41	0.47
12:AP:22:LYS:HD3	12:AP:101:ARG:NH1	2.30	0.47
15:AR:80:SER:HB3	15:AR:83:ILE:HG13	1.96	0.47
1:AA:310:A:P	20:AU:18:GLY:HA2	2.54	0.47
21:AV:33:LEU:O	21:AV:34:ASN:HB2	2.15	0.47
24:AW:47:ASN:O	24:AW:50:ILE:HG13	2.14	0.47
31:BA:1128:C:C4	31:BA:1139:G:C2	3.03	0.47
31:BA:1161:C:H2'	31:BA:1162:C:H6	1.80	0.47
31:BA:198:G:O2'	31:BA:199:G:H5'	2.14	0.47
31:BA:302:G:N3	31:BA:556:C:H4'	2.29	0.47
52:BB:7:A:C5'	52:BB:8:U:OP2	2.53	0.47
32:BE:158:LEU:HD12	32:BE:158:LEU:H	1.78	0.47
32:BE:172:ILE:H	32:BE:172:ILE:CD1	2.25	0.47
34:BG:200:GLU:HG3	34:BG:201:GLN:H	1.80	0.47
37:BJ:38:LEU:O	37:BJ:42:ILE:HG13	2.14	0.47
39:BL:3:GLN:CG	39:BL:20:ARG:NH1	2.77	0.47
39:BL:37:PHE:CE1	39:BL:74:ILE:HG12	2.49	0.47
40:BM:22:LYS:HD2	40:BM:22:LYS:C	2.35	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:BM:48:THR:CB	40:BM:62:HIS:HB3	2.43	0.47
42:BO:57:LYS:HG3	42:BO:67:THR:HG22	1.96	0.47
45:BR:33:THR:HG23	45:BR:63:ARG:NH1	2.29	0.47
47:BT:67:LYS:HG2	47:BT:68:ARG:N	2.30	0.47
47:BT:95:TYR:HD1	47:BT:98:LEU:HD12	1.79	0.47
49:BV:67:VAL:C	49:BV:69:HIS:H	2.17	0.47
50:BW:18:GLN:NE2	50:BW:22:ARG:HH12	2.13	0.47
51:BX:2:GLY:C	51:BX:4:GLY:H	2.18	0.47
54:CA:1037:C:H2'	54:CA:1038:C:C6	2.50	0.47
54:CA:1100:C:O2	54:CA:1102:A:H5'	2.14	0.47
54:CA:1095:U:OP1	54:CA:1108:G:N1	2.47	0.47
54:CA:1176:A:N6	54:CA:1177:G:N1	2.63	0.47
54:CA:170:U:O2'	54:CA:171:A:H5'	2.15	0.47
54:CA:818:G:C3'	54:CA:819:A:C5'	2.93	0.47
54:CA:973:G:O4'	40:CM:55:LYS:HG2	2.15	0.47
33:CF:162:GLN:HG2	53:C1:54:U:O2	2.14	0.47
33:CF:23:TYR:O	33:CF:24:ALA:HB2	2.14	0.47
33:CF:70:VAL:O	33:CF:106:VAL:HG23	2.15	0.47
34:CG:151:LYS:HD3	34:CG:151:LYS:O	2.15	0.47
34:CG:60:GLU:OE2	34:CG:198:VAL:HA	2.15	0.47
34:CG:23:GLY:O	34:CG:27:TYR:HD1	1.97	0.47
39:CL:50:LEU:HD23	39:CL:85:LEU:CD2	2.44	0.47
41:CN:21:ILE:HG13	41:CN:30:VAL:CG1	2.45	0.47
45:CR:74:ASP:OD2	45:CR:77:ARG:HD3	2.14	0.47
22:D3:14:ARG:HB2	22:D3:14:ARG:HE	1.23	0.47
22:D3:3:HIS:CG	22:D3:4:LYS:N	2.83	0.47
22:D3:50:ASN:O	22:D3:81:VAL:HG21	2.15	0.47
30:D8:15:LYS:HD3	30:D8:16:ILE:N	2.30	0.47
55:DA:1071:G:N1	55:DA:1091:G:N7	2.63	0.47
55:DA:1082:U:H2'	57:DY:41:ARG:CZ	2.45	0.47
55:DA:138:G:H5'	55:DA:138:G:N3	2.30	0.47
55:DA:1579:A:H2'	55:DA:1580:A:O4'	2.15	0.47
55:DA:1952:A:C6	55:DA:1953:A:N1	2.82	0.47
55:DA:2112:G:H1	55:DA:2169:A:N6	2.13	0.47
55:DA:2243:U:O2	55:DA:2434:A:C2	2.68	0.47
55:DA:2330:G:H2'	55:DA:2331:G:O4'	2.15	0.47
55:DA:2481:G:O2'	55:DA:2482:G:P	2.73	0.47
55:DA:2702:U:H2'	55:DA:2702:U:O2	2.15	0.47
55:DA:2717:G:O2'	15:DR:96:ARG:HD3	2.15	0.47
55:DA:2760:C:C2'	55:DA:2761:G:C5'	2.76	0.47
55:DA:2740:A:N6	55:DA:2764:A:C8	2.82	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2779:U:H5'	55:DA:2780:G:OP1	2.14	0.47
55:DA:3:U:C2	55:DA:4:C:C6	3.03	0.47
55:DA:510:C:C2'	55:DA:511:U:H5'	2.44	0.47
55:DA:572:A:OP2	17:D2:78:LYS:NZ	2.43	0.47
3:DD:133:LEU:HA	3:DD:136:ILE:HD13	1.97	0.47
3:DD:2:ALA:O	3:DD:3:VAL:HB	2.15	0.47
4:DE:134:ILE:HD12	4:DE:134:ILE:C	2.36	0.47
5:DF:63:LYS:HE2	5:DF:67:GLN:HB2	1.96	0.47
7:DH:59:ARG:HG3	7:DH:59:ARG:NH1	2.27	0.47
56:DI:24:ILE:O	56:DI:27:LEU:CB	2.61	0.47
56:DJ:14:GLN:H	56:DJ:17:VAL:HG23	1.79	0.47
8:DK:29:TYR:O	8:DK:32:PRO:HD2	2.14	0.47
8:DK:41:GLU:O	8:DK:45:LYS:HB2	2.15	0.47
8:DK:57:ARG:CA	8:DK:60:GLU:HB3	2.45	0.47
8:DK:64:GLU:HA	8:DK:64:GLU:OE1	2.14	0.47
11:DO:31:ALA:O	11:DO:32:THR:HG22	2.15	0.47
12:DP:18:LYS:O	12:DP:19:GLY:C	2.53	0.47
21:DV:192:ALA:O	21:DV:193:GLU:OE1	2.32	0.47
24:DW:53:LEU:O	24:DW:57:ILE:HG13	2.14	0.47
57:DY:102:LYS:CD	57:DY:103:GLY:N	2.78	0.47
57:DY:75:GLN:NE2	57:DY:75:GLN:CA	2.75	0.47
13:A0:41:ALA:HB1	13:A0:97:VAL:CG1	2.45	0.47
6:AG:5:VAL:HG22	26:A4:25:TYR:CZ	2.50	0.47
1:AA:1043:C:H42	1:AA:1112:G:H1	1.63	0.47
1:AA:1093:G:O2'	1:AA:1094:U:H5'	2.15	0.47
1:AA:1287:A:OP1	13:A0:105:ARG:O	2.33	0.47
1:AA:1665:A:H2'	1:AA:1666:G:O4'	2.14	0.47
1:AA:1965:C:H2'	1:AA:1966:A:C8	2.48	0.47
1:AA:2102:U:O2'	1:AA:2103:C:H5'	2.15	0.47
1:AA:2279:G:N2	1:AA:2280:G:H1'	2.30	0.47
1:AA:227:A:O2'	1:AA:228:A:P	2.73	0.47
1:AA:21:A:O2'	1:AA:22:C:H5'	2.14	0.47
1:AA:2389:G:H5''	1:AA:2390:U:O4'	2.15	0.47
1:AA:2756:U:H1'	1:AA:2757:A:C8	2.50	0.47
1:AA:2780:G:OP2	9:AM:118:LYS:HE2	2.15	0.47
1:AA:372:G:C2'	1:AA:373:U:OP2	2.62	0.47
1:AA:471:A:H8	1:AA:471:A:O5'	1.97	0.47
1:AA:726:G:HO2'	1:AA:727:A:H8	1.62	0.47
1:AA:807:U:O2'	1:AA:808:G:H5'	2.15	0.47
3:AD:166:GLN:HE21	3:AD:166:GLN:HA	1.79	0.47
4:AE:39:PRO:HA	4:AE:43:GLY:N	2.29	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:AF:116:ASP:OD2	11:AO:1:MET:N	2.47	0.47
5:AF:17:ARG:HH11	5:AF:17:ARG:HG3	1.80	0.47
9:AM:30:ILE:CG2	9:AM:34:LEU:HD21	2.44	0.47
9:AM:62:VAL:CG2	9:AM:66:LYS:HG3	2.39	0.47
10:AN:98:VAL:HG12	10:AN:117:LEU:HB3	1.97	0.47
11:AO:138:LEU:HD12	11:AO:139:LYS:N	2.30	0.47
11:AO:48:PRO:O	11:AO:51:PHE:N	2.47	0.47
11:AO:52:GLU:HB2	11:AO:53:GLY:H	1.60	0.47
12:AP:26:TYR:O	12:AP:27:VAL:C	2.52	0.47
12:AP:78:PRO:O	12:AP:79:LEU:HG	2.15	0.47
18:AS:75:TYR:CE2	18:AS:104:THR:HB	2.50	0.47
18:AS:20:VAL:O	18:AS:21:VAL:C	2.53	0.47
19:AT:12:VAL:HG22	19:AT:17:ALA:HB2	1.97	0.47
21:AV:81:ARG:HD3	21:AV:81:ARG:O	2.15	0.47
24:AW:69:ARG:HB2	24:AW:69:ARG:CZ	2.45	0.47
53:B1:43:U:C2'	53:B1:44:U:H5'	2.45	0.47
31:BA:1296:C:H3'	31:BA:1297:C:H6	1.78	0.47
31:BA:1506:U:O2'	31:BA:1507:A:P	2.73	0.47
31:BA:22:G:H2'	31:BA:23:C:H6	1.80	0.47
31:BA:255:G:H2'	31:BA:256:U:H6	1.80	0.47
31:BA:560:U:O2'	31:BA:561:U:OP2	2.22	0.47
31:BA:641:U:C4'	31:BA:642:A:OP1	2.62	0.47
31:BA:691:G:O2'	31:BA:797:C:H4'	2.15	0.47
31:BA:916:G:H2'	31:BA:916:G:N3	2.30	0.47
52:BD:70:G:H2'	52:BD:71:G:C8	2.50	0.47
33:BF:78:GLY:HA3	33:BF:83:ARG:HB3	1.97	0.47
34:BG:38:TYR:O	34:BG:38:TYR:HD2	1.98	0.47
37:BJ:118:VAL:HG23	37:BJ:119:ARG:N	2.30	0.47
39:BL:4:TYR:HA	39:BL:88:TYR:HE1	1.80	0.47
43:BP:70:LEU:O	43:BP:74:VAL:HG23	2.15	0.47
49:BV:29:ARG:O	49:BV:30:LEU:CB	2.57	0.47
50:BW:26:ASN:HA	50:BW:29:LYS:HG2	1.96	0.47
54:CA:1137:C:O2'	54:CA:1138:G:N3	2.47	0.47
54:CA:1473:A:O2'	54:CA:1474:G:H5'	2.14	0.47
54:CA:419:C:H2'	54:CA:420:U:H5'	1.96	0.47
54:CA:602:A:H2'	54:CA:603:U:C6	2.50	0.47
54:CA:953:G:C5'	54:CA:965:A:H61	2.25	0.47
52:CC:2:C:H2'	52:CC:3:C:C6	2.50	0.47
52:CC:68:C:H2'	52:CC:69:G:C8	2.50	0.47
52:CD:58:A:O2'	52:CD:59:U:OP1	2.32	0.47
32:CE:213:LEU:HD21	32:CE:217:ARG:HH11	1.78	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:CE:69:LEU:CD1	32:CE:91:PRO:HB2	2.43	0.47
37:CJ:73:MET:HG2	37:CJ:90:GLU:CA	2.42	0.47
37:CJ:78:ARG:HH11	37:CJ:78:ARG:HG3	1.80	0.47
41:CN:81:ASP:O	41:CN:82:VAL:C	2.53	0.47
43:CP:57:ARG:HH11	43:CP:57:ARG:CB	2.20	0.47
48:CU:53:ARG:NH2	48:CU:59:SER:C	2.68	0.47
13:D0:63:ARG:HA	13:D0:80:PHE:CZ	2.50	0.47
16:D1:107:ALA:O	16:D1:110:VAL:HB	2.14	0.47
55:DA:1225:C:O2'	17:D2:85:LYS:HA	2.15	0.47
55:DA:1059:G:H3'	55:DA:1059:G:C8	2.50	0.47
55:DA:1173:G:H4'	55:DA:1174:A:N1	2.30	0.47
55:DA:1188:U:H5'	17:D2:79:VAL:CG2	2.43	0.47
55:DA:1188:U:H2'	55:DA:1189:A:O5'	2.15	0.47
55:DA:1511:A:H2'	55:DA:1512:G:H5'	1.97	0.47
55:DA:2168:G:N2	55:DA:2170:A:OP2	2.47	0.47
55:DA:2444:G:OP2	5:DF:68:LYS:CE	2.63	0.47
55:DA:2808:U:C2'	55:DA:2809:A:H5'	2.44	0.47
55:DA:2813:A:H2'	55:DA:2814:C:O4'	2.15	0.47
55:DA:28:A:N6	55:DA:512:G:H1'	2.28	0.47
55:DA:65:C:H2'	55:DA:66:C:H6	1.78	0.47
55:DA:705:A:C2	55:DA:727:A:H1'	2.50	0.47
55:DA:95:G:H5'	24:DW:46:GLN:OE1	2.15	0.47
2:DB:29:A:H2'	2:DB:30:C:C6	2.50	0.47
3:DD:177:LEU:HD12	3:DD:181:GLU:HG2	1.97	0.47
3:DD:72:LYS:HB3	3:DD:75:ILE:HD12	1.96	0.47
6:DG:68:PRO:HG2	6:DG:90:LEU:HD12	1.97	0.47
56:DI:4:ASP:OD2	56:DI:5:ILE:CD1	2.62	0.47
8:DK:82:ARG:O	8:DK:89:TYR:HD1	1.98	0.47
11:DO:66:GLY:O	11:DO:67:MET:CB	2.63	0.47
11:DO:81:GLN:NE2	11:DO:106:LEU:O	2.48	0.47
12:DP:103:MET:HE1	12:DP:125:LEU:HD13	1.96	0.47
14:DQ:25:ARG:CB	14:DQ:25:ARG:HH11	2.27	0.47
14:DQ:25:ARG:HB3	14:DQ:25:ARG:HH11	1.80	0.47
14:DQ:62:LYS:HB3	14:DQ:97:ARG:CD	2.44	0.47
55:DA:138:G:N2	19:DT:44:GLU:OE2	2.35	0.47
21:DV:10:ARG:NH2	21:DV:26:GLY:O	2.48	0.47
13:A0:94:TYR:O	13:A0:117:VAL:HG12	2.15	0.47
13:A0:28:LEU:HD22	13:A0:28:LEU:O	2.14	0.47
13:A0:31:HIS:C	13:A0:33:ARG:H	2.18	0.47
16:A1:33:ARG:O	16:A1:37:GLU:HG3	2.14	0.47
26:A4:24:THR:O	26:A4:25:TYR:HB2	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
27:A5:6:VAL:HG22	27:A5:7:PRO:HD2	1.97	0.47
1:AA:1204:A:C2	1:AA:1206:G:N2	2.83	0.47
1:AA:55:G:O2'	1:AA:127:A:N1	2.35	0.47
1:AA:1412:A:H2'	1:AA:1413:G:C8	2.49	0.47
1:AA:1455:G:C2	1:AA:1456:G:C8	3.03	0.47
1:AA:1517:G:H2'	1:AA:1518:C:H6	1.80	0.47
1:AA:2212:A:H1'	1:AA:2215:G:C5	2.50	0.47
1:AA:2262:U:H4'	1:AA:2328:A:C2	2.50	0.47
1:AA:2348:U:H4'	28:A6:42:TRP:CD1	2.49	0.47
1:AA:322:A:H2	1:AA:339:U:O4	1.98	0.47
1:AA:3:U:C2	1:AA:4:C:C6	3.03	0.47
1:AA:610:C:H2'	1:AA:611:C:H6	1.80	0.47
1:AA:608:A:N1	1:AA:621:A:N7	2.63	0.47
1:AA:824:A:H1'	1:AA:2358:G:N7	2.30	0.47
1:AA:865:C:H2'	1:AA:865:C:O2	2.14	0.47
1:AA:923:C:H2'	1:AA:924:C:H6	1.80	0.47
1:AA:999:U:C2'	1:AA:1000:A:C5'	2.88	0.47
2:AB:78:A:H3'	2:AB:79:C:C6	2.50	0.47
3:AD:263:ARG:HB2	3:AD:263:ARG:CZ	2.43	0.47
4:AE:199:ARG:CZ	4:AE:199:ARG:HB2	2.45	0.47
8:AK:144:VAL:O	8:AK:145:VAL:CG2	2.62	0.47
9:AM:132:ALA:O	9:AM:133:GLN:C	2.53	0.47
11:AO:31:ALA:C	11:AO:32:THR:HG23	2.35	0.47
11:AO:95:VAL:HA	11:AO:99:LEU:HD23	1.97	0.47
14:AQ:48:LEU:CD2	14:AQ:82:ILE:HD11	2.45	0.47
14:AQ:52:SER:O	14:AQ:56:LEU:HD21	2.15	0.47
20:AU:81:LYS:O	20:AU:96:ILE:HG22	2.15	0.47
21:AV:108:PRO:HB2	21:AV:141:VAL:O	2.02	0.47
24:AW:32:LEU:HB2	24:AW:53:LEU:HD13	1.96	0.47
53:B1:52:U:C2'	53:B1:53:U:O5'	2.63	0.47
31:BA:1387:G:C4	31:BA:1388:C:C5	3.03	0.47
15:AR:118:ARG:NH1	31:BA:1446:A:C6	2.83	0.47
31:BA:209:U:O2	31:BA:209:U:H2'	2.13	0.47
31:BA:570:G:C6	31:BA:873:A:C2	3.03	0.47
31:BA:595:G:N1	31:BA:641:U:H2'	2.30	0.47
31:BA:686:U:O2'	31:BA:687:A:O5'	2.33	0.47
31:BA:802:A:H2'	31:BA:803:G:O4'	2.15	0.47
31:BA:953:G:H2'	31:BA:954:G:O4'	2.14	0.47
52:BC:37:MIA:N3	52:BC:37:MIA:H2'	2.30	0.47
52:BD:24:G:C2'	52:BD:25:C:H5'	2.45	0.47
34:BG:209:ARG:HG3	34:BG:209:ARG:HH11	1.80	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:BG:59:ARG:NH2	34:BG:66:ARG:NH1	2.63	0.47
37:BJ:131:LYS:HZ3	37:BJ:131:LYS:HB2	1.79	0.47
37:BJ:78:ARG:HD2	37:BJ:79:ARG:H	1.80	0.47
38:BK:63:LEU:HB2	38:BK:65:TYR:CE1	2.50	0.47
40:BM:50:ILE:HA	40:BM:60:ARG:HB2	1.97	0.47
54:CA:1157:A:C2	54:CA:1181:G:H1'	2.50	0.47
54:CA:1178:G:N7	54:CA:1180:A:OP2	2.48	0.47
54:CA:140:A:H2'	54:CA:141:A:O4'	2.14	0.47
54:CA:1528:U:O2'	54:CA:1529:G:P	2.73	0.47
54:CA:129(A):G:N3	54:CA:188:U:O2'	2.47	0.47
54:CA:262:A:C6	54:CA:263:A:N6	2.82	0.47
54:CA:430:A:H2'	54:CA:431:A:C5'	2.44	0.47
54:CA:644:G:H2'	54:CA:645:C:C5'	2.44	0.47
32:CE:55:PHE:HA	32:CE:58:ILE:HG12	1.97	0.47
33:CF:6:HIS:HB3	44:CQ:49:HIS:CD2	2.50	0.47
34:CG:172:PRO:HD2	34:CG:173:TRP:CZ3	2.50	0.47
34:CG:43:HIS:HA	34:CG:46:LYS:HG2	1.97	0.47
54:CA:15:G:H4'	35:CH:24:ARG:NH1	2.29	0.47
37:CJ:17:VAL:HG12	37:CJ:18:TYR:CD1	2.50	0.47
39:CL:65:VAL:C	39:CL:66:ARG:HG3	2.36	0.47
54:CA:255:G:O4'	47:CT:16:GLN:HB2	2.15	0.47
50:CW:87:LYS:O	50:CW:88:VAL:C	2.52	0.47
28:D6:47:THR:HG22	28:D6:48:VAL:H	1.78	0.47
55:DA:1110:G:H2'	55:DA:1111:A:H8	1.80	0.47
55:DA:1173:G:N3	55:DA:1175:U:C5	2.83	0.47
55:DA:1177:A:H5''	55:DA:1178:C:H5''	1.97	0.47
55:DA:1485:G:H2'	55:DA:1486:A:O5'	2.15	0.47
55:DA:2097:C:C2'	55:DA:2098:U:H5'	2.45	0.47
55:DA:2186:G:H2'	55:DA:2187:G:C8	2.50	0.47
55:DA:2426:A:HO2'	55:DA:2427:C:P	2.38	0.47
55:DA:991:C:H2'	55:DA:992:C:C6	2.49	0.47
3:DD:89:SER:C	3:DD:159:ALA:HB2	2.35	0.47
3:DD:134:ARG:HD2	3:DD:188:GLU:OE2	2.15	0.47
3:DD:25:THR:O	3:DD:26:LYS:C	2.53	0.47
5:DF:23:ASP:O	5:DF:24:LEU:O	2.32	0.47
6:DG:177:GLY:O	6:DG:179:PRO:HD3	2.15	0.47
58:DL:10:LEU:HG	58:DL:55:VAL:HG11	1.97	0.47
58:DL:20:ALA:CB	58:DL:21:PRO:CD	2.91	0.47
58:DL:21:PRO:O	58:DL:24:GLY:C	2.54	0.47
58:DL:52:ILE:CG1	58:DL:76:TYR:CA	2.93	0.47
58:DL:77:LEU:HB3	58:DL:107:ILE:HD13	1.96	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:DM:15:LEU:HB2	9:DM:134:ARG:HB2	1.95	0.47
12:DP:42:ILE:HD13	12:DP:97:VAL:HG21	1.96	0.47
14:DQ:100:ALA:O	14:DQ:103:GLU:HG2	2.15	0.47
15:DR:23:ARG:HG2	15:DR:120:ARG:NH1	2.30	0.47
55:DA:2012:G:H4'	18:DS:96:ILE:CD1	2.45	0.47
21:DV:33:LEU:HD23	21:DV:90:VAL:HG21	1.97	0.47
57:DY:38:HIS:O	57:DY:96:PHE:HZ	1.97	0.47
16:A1:98:LEU:O	16:A1:99:ALA:HB3	2.15	0.46
17:A2:85:LYS:CG	17:A2:86:GLY:N	2.78	0.46
22:A3:43:THR:C	22:A3:45:PHE:N	2.68	0.46
22:A3:74:ARG:HG2	22:A3:75:LEU:CD2	2.44	0.46
26:A4:58:ARG:HG3	26:A4:59:PHE:N	2.29	0.46
30:A8:50:LEU:CD1	30:A8:53:PRO:O	2.63	0.46
1:AA:1019:U:O2'	1:AA:1021:A:C2	2.63	0.46
1:AA:1288:U:H4'	1:AA:1289:C:OP2	2.15	0.46
1:AA:1359:A:C5'	1:AA:1359:A:H8	2.25	0.46
1:AA:1640:C:H2'	1:AA:1641:A:O4'	2.13	0.46
1:AA:2038:G:H2'	1:AA:2039:C:O4'	2.15	0.46
1:AA:910:A:H2	1:AA:2264:C:O2	1.97	0.46
1:AA:2314:C:C2'	1:AA:2315:G:H5'	2.45	0.46
1:AA:2419:U:O4	30:A8:31:HIS:CE1	2.67	0.46
1:AA:2553:G:H2'	1:AA:2554:U:C4'	2.45	0.46
1:AA:2759:G:H8	1:AA:2759:G:H5'	1.80	0.46
1:AA:2780:G:O2'	1:AA:2781:A:P	2.73	0.46
1:AA:454:A:HO2'	1:AA:455:C:P	2.39	0.46
1:AA:448:U:C4	1:AA:583:G:H1'	2.50	0.46
1:AA:742:G:O2'	1:AA:743:G:H5'	2.15	0.46
1:AA:917:A:C2	1:AA:918:A:H1'	2.50	0.46
2:AB:94:C:C2'	2:AB:95:U:H5'	2.46	0.46
4:AE:203:LYS:O	4:AE:204:ALA:CB	2.63	0.46
9:AM:63:THR:O	9:AM:64:GLY:O	2.32	0.46
11:AO:23:PRO:O	11:AO:24:GLY:C	2.52	0.46
11:AO:38:GLN:HG2	11:AO:45:LEU:HD13	1.97	0.46
15:AR:74:ARG:HG2	15:AR:74:ARG:HH11	1.78	0.46
18:AS:69:LEU:HA	18:AS:108:GLY:O	2.15	0.46
1:AA:1342:A:OP2	19:AT:56:THR:O	2.33	0.46
20:AU:21:LYS:O	20:AU:21:LYS:CG	2.62	0.46
20:AU:84:ARG:HB3	20:AU:95:LYS:CE	2.45	0.46
20:AU:97:ARG:NH1	20:AU:97:ARG:HG2	2.30	0.46
24:AW:46:GLN:C	24:AW:49:LYS:HE3	2.35	0.46
24:AW:56:GLN:O	24:AW:60:LEU:CD1	2.64	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:B1:43:U:H2'	53:B1:44:U:C5'	2.45	0.46
31:BA:1006:C:O2'	31:BA:1007:C:H5'	2.14	0.46
31:BA:1014:A:H4'	49:BV:14:HIS:CE1	2.50	0.46
31:BA:1032(A):G:H2'	31:BA:1032(B):G:C8	2.50	0.46
31:BA:1327:C:H2'	31:BA:1328:C:C6	2.49	0.46
31:BA:191(F):U:H2'	31:BA:191:G:C5'	2.44	0.46
31:BA:24:U:H2'	31:BA:25:C:H6	1.80	0.46
31:BA:641:U:H5''	31:BA:642:A:OP1	2.15	0.46
31:BA:923:A:H2'	31:BA:924:C:C6	2.49	0.46
33:BF:145:GLY:O	33:BF:146:ALA:O	2.33	0.46
33:BF:3:ASN:N	33:BF:3:ASN:OD1	2.48	0.46
33:BF:79:ARG:C	33:BF:81:GLY:H	2.18	0.46
34:BG:26:CYS:SG	34:BG:31:CYS:O	2.73	0.46
34:BG:4:TYR:HE2	34:BG:7:PRO:O	1.98	0.46
37:BJ:57:GLU:OE2	37:BJ:60:LYS:HD3	2.15	0.46
39:BL:47:LEU:CD1	39:BL:47:LEU:N	2.78	0.46
41:BN:95:ILE:HG21	41:BN:108:ILE:HD13	1.97	0.46
43:BP:68:GLY:HA2	43:BP:71:ARG:HB3	1.96	0.46
33:BF:29:TYR:OH	44:BQ:54:PRO:HD2	2.14	0.46
33:BF:13:GLY:HA2	44:BQ:57:ARG:CZ	2.45	0.46
48:BU:25:THR:O	48:BU:25:THR:HG22	2.15	0.46
48:BU:62:GLU:HA	48:BU:65:ILE:CG1	2.45	0.46
50:BW:53:LEU:HD12	50:BW:100:ILE:O	2.15	0.46
50:BW:58:LYS:O	50:BW:58:LYS:HD3	2.15	0.46
53:C1:31:A:O2'	53:C1:32:A:OP1	2.33	0.46
53:C1:52:U:H2'	53:C1:53:U:C5'	2.45	0.46
53:C1:55:U:O2'	53:C1:56:U:C2	2.69	0.46
54:CA:1091:U:H2'	54:CA:1093:A:OP2	2.16	0.46
54:CA:1148:U:C2'	54:CA:1149:C:H5'	2.45	0.46
54:CA:1270:C:O2'	54:CA:1314:C:H5'	2.14	0.46
54:CA:1404:C:H6	54:CA:1404:C:O5'	1.97	0.46
54:CA:1453:G:H2'	50:CW:39:LYS:NZ	2.28	0.46
54:CA:1534:A:C2	53:C1:39:U:C2	3.03	0.46
52:CC:36:A:H2'	52:CC:37:MIA:O4'	2.15	0.46
32:CE:165:VAL:HG23	32:CE:166:ASP:N	2.25	0.46
32:CE:25:ASN:ND2	32:CE:193:ASP:HB3	2.31	0.46
33:CF:120:VAL:HB	33:CF:198:VAL:HG11	1.95	0.46
37:CJ:23:VAL:HG13	37:CJ:43:PHE:CE2	2.50	0.46
43:CP:119:GLY:CA	43:CP:120:LYS:HD3	2.44	0.46
43:CP:19:LEU:N	43:CP:19:LEU:HD22	2.30	0.46
45:CR:37:ASN:HD22	45:CR:37:ASN:N	2.13	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:CV:5:LEU:C	49:CV:5:LEU:HD12	2.34	0.46
27:D5:50:GLY:O	27:D5:51:TYR:HB2	2.14	0.46
27:D5:58:LEU:C	27:D5:60:VAL:H	2.19	0.46
29:D7:36:GLN:HG2	29:D7:36:GLN:O	2.14	0.46
55:DA:1079:C:C2'	55:DA:1080:A:C8	2.98	0.46
55:DA:1091:G:H2'	55:DA:1092:C:H5'	1.97	0.46
55:DA:1092:C:C2'	55:DA:1093:G:H5'	2.45	0.46
55:DA:1730:U:H3'	55:DA:1730:U:H6	1.79	0.46
55:DA:858:U:O2'	55:DA:2268:A:C2'	2.64	0.46
55:DA:2543:G:H2'	55:DA:2544:G:O4'	2.16	0.46
55:DA:2688:U:O2	55:DA:2688:U:H3'	2.15	0.46
55:DA:1637:A:H4'	55:DA:2711:A:O2'	2.15	0.46
55:DA:2803:C:O2'	55:DA:2804:C:H5'	2.15	0.46
55:DA:2850:A:H2	13:D0:61:HIS:CG	2.33	0.46
55:DA:330:A:H2	55:DA:1210:A:H2'	1.79	0.46
55:DA:802:A:C2'	55:DA:803:U:H5''	2.45	0.46
3:DD:96:HIS:ND1	3:DD:102:LYS:HG2	2.29	0.46
3:DD:83:GLU:HG3	3:DD:92:ILE:CD1	2.44	0.46
4:DE:26:ILE:HG13	4:DE:196:VAL:HG21	1.96	0.46
6:DG:118:ARG:HA	6:DG:118:ARG:NE	2.28	0.46
6:DG:91:ARG:HG2	6:DG:92:VAL:N	2.30	0.46
7:DH:124:GLU:CB	7:DH:132:ARG:HG3	2.45	0.46
56:DI:8:ILE:O	56:DI:11:GLU:N	2.49	0.46
8:DK:69:LYS:HG3	8:DK:136:VAL:CB	2.43	0.46
58:DL:56:GLU:HG2	58:DL:57:ILE:O	2.15	0.46
58:DL:98:ARG:HB3	58:DL:98:ARG:CZ	2.45	0.46
9:DM:114:ARG:O	9:DM:115:ARG:HB3	2.14	0.46
11:DO:85:LEU:CD2	11:DO:114:ILE:HD12	2.45	0.46
12:DP:90:VAL:CG1	12:DP:91:GLU:N	2.78	0.46
15:DR:18:ASP:N	15:DR:18:ASP:OD1	2.48	0.46
18:DS:22:ASP:HA	18:DS:25:ARG:HH12	1.79	0.46
20:DU:49:VAL:HG11	20:DU:50:ARG:NH2	2.30	0.46
57:DY:75:GLN:HB2	57:DY:111:LEU:HA	1.90	0.46
16:A1:46:ALA:O	16:A1:47:TYR:C	2.53	0.46
17:A2:61:VAL:CG1	17:A2:62:LEU:H	2.26	0.46
17:A2:85:LYS:HG3	17:A2:86:GLY:N	2.26	0.46
29:A7:17:GLY:O	29:A7:21:ARG:HG2	2.16	0.46
1:AA:1311:G:N2	1:AA:1603:A:H62	2.12	0.46
1:AA:1357:U:H2'	1:AA:1358:G:O4'	2.15	0.46
1:AA:2033:A:H2'	1:AA:2035:G:OP2	2.15	0.46
1:AA:2050:C:H2'	1:AA:2051:A:O4'	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:224:G:N7	1:AA:420:C:H4'	2.30	0.46
1:AA:2281:C:O2'	1:AA:2282:G:H5'	2.15	0.46
1:AA:2288:A:H5''	1:AA:2289:G:OP2	2.16	0.46
1:AA:2311:A:H2'	1:AA:2312:U:C6	2.51	0.46
1:AA:2612:C:C5	1:AA:2613:U:H5	2.32	0.46
1:AA:2695:C:H2'	1:AA:2696:U:H6	1.80	0.46
1:AA:2813:A:H2'	1:AA:2814:C:O4'	2.15	0.46
1:AA:2807:G:C6	1:AA:2893:G:O6	2.68	0.46
1:AA:554:U:HO2'	1:AA:556:G:H8	1.57	0.46
1:AA:72:U:C4	1:AA:112:U:H4'	2.50	0.46
2:AB:105:G:O2'	2:AB:106:G:H5'	2.15	0.46
2:AB:40:U:N3	2:AB:43:C:OP2	2.47	0.46
1:AA:1825:A:C4'	3:AD:254:THR:HG21	2.44	0.46
3:AD:35:LYS:HD3	3:AD:63:ARG:CG	2.44	0.46
4:AE:15:PHE:CB	15:AR:81:PRO:HG2	2.44	0.46
4:AE:47:VAL:HG23	4:AE:84:PHE:HB3	1.96	0.46
4:AE:35:GLN:CG	4:AE:64:LYS:HZ2	2.20	0.46
5:AF:156:LEU:HD21	5:AF:163:VAL:HG12	1.97	0.46
5:AF:80:ALA:O	5:AF:83:PHE:HB2	2.14	0.46
8:AK:123:LEU:HD22	8:AK:143:SER:CB	2.43	0.46
11:AO:33:ARG:O	11:AO:34:GLY:O	2.33	0.46
12:AP:21:THR:O	12:AP:22:LYS:C	2.53	0.46
14:AQ:56:LEU:HG	14:AQ:58:LEU:HD22	1.98	0.46
20:AU:28:LYS:O	20:AU:29:GLU:O	2.32	0.46
20:AU:97:ARG:NH2	20:AU:98:VAL:HB	2.10	0.46
21:AV:175:VAL:CG2	21:AV:176:PRO:CD	2.93	0.46
23:AZ:76:ARG:CB	23:AZ:94:LEU:HD13	2.45	0.46
53:B1:57:U:C2'	53:B1:57:U:O2	2.63	0.46
31:BA:1021:G:H5'	31:BA:1021:G:H8	1.79	0.46
31:BA:1158:C:C2'	31:BA:1158:C:O2	2.64	0.46
31:BA:977:A:N6	31:BA:1224:G:OP1	2.48	0.46
10:AN:49:ARG:NH1	31:BA:1422:G:O3'	2.49	0.46
31:BA:1530:G:H2'	31:BA:1531:A:N7	2.30	0.46
31:BA:167:G:O2'	31:BA:168:G:H5'	2.15	0.46
31:BA:513:C:H2'	31:BA:514:C:H6	1.80	0.46
31:BA:948:C:O2'	31:BA:949:A:H5'	2.16	0.46
52:BB:8:U:O2'	52:BB:9:A:C5'	2.63	0.46
32:BE:126:GLU:C	32:BE:128:GLU:N	2.69	0.46
32:BE:194:PRO:HG2	32:BE:195:ASP:OD1	2.15	0.46
32:BE:196:LEU:HD12	32:BE:197:VAL:HG23	1.97	0.46
32:BE:83:MET:SD	32:BE:234:PRO:HG2	2.55	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:BF:11:ARG:O	33:BF:13:GLY:N	2.47	0.46
33:BF:134:ILE:HG21	33:BF:168:ALA:HB3	1.97	0.46
34:BG:10:ARG:HH11	34:BG:10:ARG:HG3	1.80	0.46
34:BG:18:LYS:HG2	34:BG:21:LEU:HD21	1.96	0.46
35:BH:82:VAL:HG21	35:BH:141:GLN:NE2	2.30	0.46
39:BL:40:LEU:HD11	39:BL:70:LYS:CG	2.41	0.46
42:BO:41:ARG:HB3	42:BO:42:THR:H	1.63	0.46
46:BS:53:VAL:HG23	46:BS:54:GLU:N	2.30	0.46
47:BT:8:GLY:HA3	47:BT:21:VAL:HG12	1.97	0.46
26:A4:63:TYR:HH	49:BV:39:THR:HB	1.74	0.46
49:BV:42:PRO:HA	49:BV:45:VAL:CG2	2.44	0.46
51:BX:9:ARG:CZ	51:BX:10:ARG:HA	2.46	0.46
53:C1:30:C:C4	53:C1:31:A:C8	3.03	0.46
53:C1:33:G:H2'	53:C1:34:G:N7	2.30	0.46
54:CA:1004:A:O5'	54:CA:1025:U:O4	2.32	0.46
54:CA:1324:A:H2'	54:CA:1325:C:H6	1.79	0.46
54:CA:687:A:H4'	54:CA:688:G:O5'	2.15	0.46
54:CA:687:A:O2'	54:CA:688:G:P	2.73	0.46
54:CA:719:C:O2	48:CU:50:ILE:HD13	2.15	0.46
52:CB:51:U:H2'	52:CB:52:G:C8	2.49	0.46
52:CD:2:C:OP1	52:CD:2:C:C4'	2.63	0.46
32:CE:16:HIS:CD2	32:CE:210:SER:HA	2.47	0.46
33:CF:11:ARG:HB3	33:CF:14:ILE:O	2.15	0.46
34:CG:9:CYS:SG	34:CG:26:CYS:SG	3.12	0.46
38:CK:111:ILE:HG22	38:CK:112:LEU:H	1.80	0.46
38:CK:36:LEU:O	38:CK:39:LEU:HB2	2.14	0.46
39:CL:9:ARG:HB3	39:CL:14:VAL:HG13	1.97	0.46
39:CL:46:ALA:HA	39:CL:78:LYS:HB2	1.97	0.46
39:CL:45:ALA:HA	39:CL:48:GLU:CD	2.35	0.46
42:CO:20:LYS:CD	42:CO:20:LYS:H	2.18	0.46
43:CP:11:ARG:CB	43:CP:11:ARG:NH1	2.77	0.46
48:CU:82:THR:HG22	48:CU:83:GLU:N	2.30	0.46
50:CW:49:ALA:CB	50:CW:99:LEU:HB2	2.43	0.46
22:D3:53:MET:HB3	22:D3:59:LEU:CD2	2.46	0.46
55:DA:1103:A:H5'	55:DA:1103:A:H8	1.80	0.46
55:DA:1107:G:OP1	57:DY:54:ALA:HA	2.15	0.46
55:DA:1535:U:H3'	55:DA:1536:A:C5'	2.45	0.46
55:DA:1535:U:C4	55:DA:1537:C:O2	2.68	0.46
55:DA:1578:U:H2'	55:DA:1579:A:C5'	2.45	0.46
55:DA:1652:A:O3'	55:DA:1653:G:C8	2.68	0.46
55:DA:1826:G:H2'	55:DA:1827:C:C6	2.50	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1906:G:N1	55:DA:1907:G:C5	2.83	0.46
55:DA:2198:A:HO2'	55:DA:2199:A:P	2.38	0.46
55:DA:2284:C:H41	28:D6:25:LYS:HZ1	1.63	0.46
55:DA:2346:A:HO2'	55:DA:2347:C:P	2.38	0.46
55:DA:2675:A:H61	55:DA:2732:G:H1	1.62	0.46
55:DA:527:C:OP2	55:DA:2779:U:C4	2.68	0.46
55:DA:297:C:H2'	55:DA:298:G:O4'	2.15	0.46
55:DA:723:G:H2'	55:DA:724:U:O4'	2.15	0.46
3:DD:136:ILE:CG2	3:DD:165:ILE:HD12	2.46	0.46
7:DH:62:LYS:O	7:DH:65:HIS:HB3	2.15	0.46
58:DL:105:LEU:CD1	58:DL:106:GLU:N	2.53	0.46
58:DL:52:ILE:CG1	58:DL:53:VAL:N	2.74	0.46
12:DP:29:PHE:N	12:DP:105:GLU:OE2	2.45	0.46
12:DP:26:TYR:O	12:DP:27:VAL:CB	2.61	0.46
14:DQ:5:THR:HG23	14:DQ:8:GLU:OE2	2.14	0.46
15:DR:29:ARG:NH1	15:DR:29:ARG:HB2	2.30	0.46
18:DS:60:ASN:ND2	18:DS:60:ASN:N	2.64	0.46
18:DS:73:ALA:HB3	18:DS:106:ILE:HD11	1.98	0.46
19:DT:44:GLU:CB	19:DT:49:VAL:O	2.63	0.46
20:DU:91:GLU:HG3	20:DU:92:ASN:OD1	2.15	0.46
55:DA:76:C:OP1	24:DW:55:ARG:HD3	2.15	0.46
16:A1:108:GLU:C	16:A1:110:VAL:N	2.68	0.46
16:A1:113:ALA:C	16:A1:115:ALA:H	2.18	0.46
16:A1:50:ARG:HH11	17:A2:72:VAL:HG21	1.80	0.46
17:A2:96:ILE:HG22	17:A2:99:ILE:HD11	1.97	0.46
26:A4:35:VAL:HG23	26:A4:37:SER:H	1.80	0.46
1:AA:686:G:C5'	29:A7:11:LYS:HE2	2.43	0.46
30:A8:52:LYS:H	30:A8:53:PRO:HD2	1.81	0.46
1:AA:1033:U:O2	1:AA:2750:A:C2	2.67	0.46
1:AA:1174:A:H5'	1:AA:1175:U:H5''	1.96	0.46
1:AA:1511:A:H2'	1:AA:1512:G:C8	2.50	0.46
1:AA:1872:A:H5'	1:AA:1878:G:OP2	2.16	0.46
1:AA:1948:G:C5'	1:AA:1948:G:H8	2.27	0.46
1:AA:2111:C:OP2	1:AA:2145:C:N4	2.49	0.46
1:AA:2517:C:HO2'	1:AA:2518:A:H3'	1.75	0.46
1:AA:2729:G:C5	1:AA:2730:C:C5	3.03	0.46
1:AA:2720:U:N3	1:AA:2873:A:H2	2.10	0.46
1:AA:2876:G:H4'	15:AR:2:ASN:O	2.16	0.46
1:AA:302:C:H2'	1:AA:303:U:C6	2.50	0.46
1:AA:411:G:H5''	1:AA:412:A:OP1	2.15	0.46
1:AA:70:G:H4'	1:AA:73:A:O4'	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:901:A:H2'	1:AA:901:A:N3	2.30	0.46
2:AB:16:G:O2'	2:AB:17:C:H5'	2.16	0.46
2:AB:94:C:H2'	2:AB:95:U:O4'	2.16	0.46
4:AE:16:ARG:O	4:AE:17:ASP:CB	2.62	0.46
4:AE:102:VAL:HA	4:AE:200:GLU:O	2.15	0.46
4:AE:102:VAL:HA	4:AE:201:THR:OG1	2.15	0.46
1:AA:1257:C:H4'	5:AF:83:PHE:CE2	2.51	0.46
6:AG:135:LEU:CD1	6:AG:135:LEU:N	2.78	0.46
6:AG:16:ARG:HB3	6:AG:17:PRO:CD	2.45	0.46
7:AH:102:ALA:CB	7:AH:117:PRO:HD3	2.45	0.46
7:AH:89:ILE:CG1	7:AH:90:LYS:N	2.79	0.46
18:AS:6:ILE:HG23	18:AS:104:THR:OG1	2.15	0.46
21:AV:2:GLU:OE2	21:AV:4:ARG:NH2	2.47	0.46
1:AA:850:C:O3'	25:AX:49:LYS:HE2	2.14	0.46
2:AB:83:G:H5''	25:AX:52:HIS:CE1	2.49	0.46
31:BA:1091:U:O2	31:BA:1093:A:C8	2.68	0.46
31:BA:1128:C:C2	31:BA:1139:G:C6	3.03	0.46
31:BA:1190:G:H8	31:BA:1190:G:O5'	1.99	0.46
31:BA:1273:G:H3'	31:BA:1274:G:C8	2.49	0.46
31:BA:1277:C:HO2'	31:BA:1279:A:H1'	1.80	0.46
31:BA:1499:A:C2	31:BA:1500:A:C8	3.03	0.46
31:BA:652:U:O4	31:BA:752:G:C2'	2.62	0.46
31:BA:818:G:C2	31:BA:820:U:H2'	2.51	0.46
31:BA:836:G:C6	31:BA:851:G:C6	3.04	0.46
52:BC:19:G:C2	52:BC:57:G:N3	2.83	0.46
32:BE:12:GLU:OE1	32:BE:16:HIS:HB2	2.15	0.46
32:BE:40:HIS:CB	32:BE:190:THR:HG21	2.40	0.46
32:BE:79:ASP:O	32:BE:82:ARG:N	2.43	0.46
35:BH:48:ALA:CB	35:BH:49:PRO:HD2	2.32	0.46
38:BK:63:LEU:N	38:BK:63:LEU:HD22	2.30	0.46
40:BM:33:GLN:H	40:BM:75:ILE:CG1	2.28	0.46
40:BM:54:PHE:CE1	40:BM:55:LYS:CE	2.98	0.46
41:BN:21:ILE:HG12	41:BN:30:VAL:HG12	1.98	0.46
42:BO:28:LYS:O	42:BO:29:GLY:C	2.54	0.46
42:BO:34:ARG:HG2	42:BO:35:GLY:N	2.30	0.46
43:BP:8:GLU:OE2	43:BP:22:ILE:HA	2.14	0.46
45:BR:62:GLN:O	45:BR:66:LEU:HD13	2.15	0.46
47:BT:80:GLY:O	47:BT:81:ARG:CB	2.64	0.46
49:BV:6:LYS:NZ	49:BV:10:PHE:HZ	2.12	0.46
54:CA:1003:G:C8	54:CA:1003:G:H5'	2.51	0.46
54:CA:1003:G:N2	54:CA:1004:A:O2'	2.48	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1171:G:H2'	54:CA:1172:C:C6	2.51	0.46
54:CA:1469:G:H2'	54:CA:1470:G:C8	2.50	0.46
54:CA:254:G:H21	47:CT:16:GLN:HE21	1.62	0.46
54:CA:843:U:H2'	54:CA:848:C:OP1	2.15	0.46
54:CA:927:G:N2	54:CA:1391:U:H1'	2.30	0.46
33:CF:113:ALA:HB2	33:CF:183:ASP:HB3	1.97	0.46
33:CF:179:ARG:O	33:CF:206:GLU:HA	2.14	0.46
36:CI:78:GLU:OE2	36:CI:81:ILE:HD12	2.15	0.46
37:CJ:144:MET:HE3	52:CD:31:A:H1'	1.98	0.46
38:CK:9:MET:O	38:CK:12:ARG:N	2.49	0.46
40:CM:54:PHE:CE1	40:CM:55:LYS:NZ	2.79	0.46
40:CM:34:VAL:CG2	40:CM:74:ILE:HG22	2.40	0.46
41:CN:59:TYR:CZ	41:CN:63:LEU:HD11	2.51	0.46
43:CP:125:ARG:HD3	43:CP:126:LYS:N	2.23	0.46
54:CA:1308:U:OP1	43:CP:98:VAL:N	2.48	0.46
45:CR:82:ILE:CG2	45:CR:83:GLU:H	2.27	0.46
46:CS:2:VAL:HG22	46:CS:3:LYS:N	2.31	0.46
48:CU:20:ALA:C	48:CU:22:VAL:H	2.19	0.46
50:CW:89:ARG:O	50:CW:93:GLU:N	2.48	0.46
13:D0:8:ARG:HH11	13:D0:39:PRO:HB3	1.80	0.46
17:D2:1:MET:HE2	17:D2:43:GLU:HB2	1.96	0.46
17:D2:38:LEU:O	17:D2:51:VAL:HG13	2.15	0.46
22:D3:6:GLY:O	22:D3:7:LEU:HB2	2.14	0.46
55:DA:1059:G:C3'	55:DA:1059:G:C8	2.99	0.46
55:DA:1360:A:H2'	55:DA:1361:G:O4'	2.14	0.46
55:DA:1419:A:H5'	55:DA:1420:U:OP2	2.16	0.46
55:DA:1782:C:H2'	55:DA:2608:G:O2'	2.15	0.46
55:DA:1932:A:H2'	55:DA:1933:G:O4'	2.15	0.46
55:DA:2469:A:O4'	55:DA:2469:A:N3	2.47	0.46
55:DA:2536:G:C6	55:DA:2537:U:C4	3.03	0.46
55:DA:2554:U:H2'	55:DA:2555:U:C6	2.50	0.46
55:DA:2872:G:C2	55:DA:2873:A:N6	2.84	0.46
55:DA:320:A:H2'	5:DF:136:THR:CG2	2.45	0.46
55:DA:530:G:C5	55:DA:2022:U:H5''	2.50	0.46
55:DA:760:G:C2'	55:DA:761:A:H5'	2.45	0.46
3:DD:72:LYS:HG2	3:DD:103:ARG:HH22	1.80	0.46
3:DD:18:VAL:HG22	3:DD:211:ARG:NH2	2.30	0.46
3:DD:35:LYS:NZ	3:DD:65:ILE:HA	2.31	0.46
4:DE:15:PHE:CE1	4:DE:20:ALA:HB2	2.50	0.46
6:DG:125:PHE:HE1	6:DG:180:PHE:HE2	1.63	0.46
56:DJ:21:LYS:O	56:DJ:24:ILE:N	2.49	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:DM:97:ARG:HA	9:DM:100:GLU:HB3	1.96	0.46
10:DN:8:LEU:HD13	10:DN:82:ASN:CB	2.45	0.46
11:DO:135:LEU:O	11:DO:136:GLU:C	2.53	0.46
54:CA:346:G:H5'	15:DR:41:ARG:HD2	1.97	0.46
15:DR:19:LEU:HD22	15:DR:86:ILE:HG22	1.97	0.46
15:DR:91:ARG:HB2	15:DR:121:ILE:CG1	2.41	0.46
55:DA:498:G:N2	20:DU:47:LYS:HZ1	2.13	0.46
21:DV:162:GLU:CG	21:DV:163:LEU:H	2.28	0.46
21:DV:9:TYR:CZ	21:DV:61:LEU:HD21	2.50	0.46
57:DY:89:ALA:CA	57:DY:92:THR:HB	2.45	0.46
23:DZ:90:ILE:O	23:DZ:94:LEU:HB3	2.15	0.46
13:A0:103:ARG:NH1	13:A0:108:GLY:O	2.49	0.46
16:A1:108:GLU:C	16:A1:110:VAL:H	2.19	0.46
27:A5:40:LYS:HD2	27:A5:46:CYS:CB	2.45	0.46
28:A6:9:LEU:HD23	28:A6:10:LEU:N	2.29	0.46
1:AA:1007:C:H5''	9:AM:35:ARG:NH1	2.30	0.46
1:AA:1059:G:H2'	1:AA:1060:U:C5	2.50	0.46
1:AA:1335:U:O2'	1:AA:1336:A:H5'	2.16	0.46
1:AA:1788:C:O2'	1:AA:1789:A:H5'	2.15	0.46
1:AA:1906:G:C5	1:AA:1929:G:N2	2.83	0.46
1:AA:2031:A:O4'	1:AA:2031:A:OP1	2.33	0.46
1:AA:2131:G:H5'	1:AA:2132:U:C5'	2.33	0.46
1:AA:2137:C:OP1	54:CA:1000:A:H4'	2.16	0.46
1:AA:2368:C:O2'	1:AA:2369:A:H5'	2.15	0.46
1:AA:263:C:O2'	1:AA:264:C:H5'	2.15	0.46
1:AA:2688:U:H1'	1:AA:2721:A:N6	2.31	0.46
1:AA:2791:C:C2	1:AA:2792:G:C8	3.04	0.46
1:AA:2852:G:C2	1:AA:2853:C:C2	3.03	0.46
1:AA:441:U:H2'	1:AA:442:G:C8	2.50	0.46
1:AA:607:U:C4	1:AA:608:A:N7	2.84	0.46
1:AA:691:C:H2'	1:AA:692:C:C6	2.49	0.46
1:AA:718:A:H3'	1:AA:719:C:H6	1.80	0.46
1:AA:810:U:O5'	1:AA:810:U:H6	1.99	0.46
2:AB:43:C:OP1	6:AG:67:LYS:NZ	2.47	0.46
1:AA:2820:A:N6	4:AE:192:ASN:HB2	2.21	0.46
4:AE:58:ARG:O	4:AE:60:ASN:N	2.49	0.46
5:AF:119:ARG:HG2	5:AF:119:ARG:HH11	1.80	0.46
7:AH:86:GLU:O	7:AH:87:LEU:HB2	2.16	0.46
9:AM:34:LEU:HD12	9:AM:116:LEU:O	2.15	0.46
9:AM:41:ASP:OD1	16:A1:100:VAL:HG22	2.14	0.46
10:AN:64:ARG:NH1	10:AN:83:ALA:HB2	2.30	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:AO:86:LYS:HB3	11:AO:117:GLU:O	2.16	0.46
12:AP:31:ASP:OD2	12:AP:107:ALA:HA	2.14	0.46
12:AP:16:ARG:HB2	12:AP:16:ARG:HH11	1.81	0.46
14:AQ:25:ARG:HB3	14:AQ:25:ARG:NH1	2.30	0.46
19:AT:49:VAL:HB	19:AT:83:VAL:HG23	1.96	0.46
31:BA:1005:A:C2	31:BA:1006:C:C2	3.04	0.46
31:BA:1126:U:H3	31:BA:1281:U:H1'	1.79	0.46
31:BA:1324:A:H4'	31:BA:1362:C:H4'	1.95	0.46
31:BA:924:C:H5'	31:BA:1399:C:OP2	2.16	0.46
31:BA:678:U:H4'	31:BA:778:G:OP1	2.16	0.46
31:BA:728:A:N1	31:BA:729:A:C6	2.83	0.46
33:BF:148:GLY:HA3	33:BF:172:ARG:O	2.15	0.46
33:BF:207:VAL:O	33:BF:207:VAL:HG12	2.15	0.46
33:BF:22:TRP:HB3	33:BF:59:ARG:HB2	1.97	0.46
33:BF:29:TYR:HE2	33:BF:33:LEU:HD13	1.79	0.46
39:BL:69:GLY:O	39:BL:70:LYS:C	2.53	0.46
48:BU:21:LYS:HA	48:BU:21:LYS:HD2	1.48	0.46
48:BU:53:ARG:C	48:BU:55:ARG:N	2.69	0.46
54:CA:1002:G:C2	54:CA:1003:G:C5	3.04	0.46
54:CA:1008:C:C5'	54:CA:1008:C:H6	2.27	0.46
54:CA:1015:A:H2'	54:CA:1016:A:O4'	2.15	0.46
54:CA:1074:G:H2'	54:CA:1075:C:C6	2.50	0.46
54:CA:1175:G:N1	54:CA:1176:A:N6	2.64	0.46
54:CA:530:G:O6	53:C1:51:U:H1'	2.15	0.46
54:CA:701:C:O2	54:CA:703:G:N2	2.48	0.46
54:CA:74:C:N4	54:CA:75:C:N4	2.63	0.46
54:CA:794:A:C8	54:CA:794:A:C4'	2.99	0.46
54:CA:865:A:H2'	54:CA:866:C:H6	1.80	0.46
52:CC:13:C:O2'	55:DA:1924:C:H4'	2.16	0.46
52:CD:9:A:H4'	52:CD:46:G:O4'	2.16	0.46
32:CE:67:THR:HG21	32:CE:155:LEU:HD11	1.97	0.46
34:CG:98:GLU:OE2	34:CG:103:ASN:ND2	2.48	0.46
36:CI:41:GLU:O	36:CI:43:LEU:N	2.47	0.46
37:CJ:95:ARG:NH1	37:CJ:95:ARG:HG3	2.31	0.46
43:CP:15:VAL:O	43:CP:19:LEU:HD23	2.15	0.46
43:CP:25:ILE:HD11	43:CP:66:LEU:HD21	1.98	0.46
49:CV:25:LYS:HA	49:CV:25:LYS:NZ	2.31	0.46
27:D5:20:ARG:C	27:D5:22:HIS:H	2.19	0.46
29:D7:38:GLY:O	29:D7:39:ARG:C	2.54	0.46
11:DO:64:LYS:CB	30:D8:25:MET:HG3	2.46	0.46
55:DA:1022:G:N2	55:DA:1142(A):A:C2	2.73	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1339:G:N2	55:DA:1603:A:H1'	2.30	0.46
55:DA:1511:A:O2'	55:DA:1512:G:H5'	2.14	0.46
55:DA:1545(A):A:H2'	55:DA:1546:C:O4'	2.15	0.46
55:DA:17:G:H2'	55:DA:18:C:C6	2.49	0.46
55:DA:1978:A:H2'	55:DA:1979:C:H6	1.79	0.46
55:DA:2044:C:H6	55:DA:2044:C:C5'	2.22	0.46
55:DA:2086:U:H2'	55:DA:2087:G:C8	2.50	0.46
55:DA:2178:C:O2'	55:DA:2179:C:H5'	2.16	0.46
55:DA:2298:A:N6	55:DA:2318:G:C8	2.73	0.46
55:DA:2439:A:H4'	55:DA:2440:C:O5'	2.16	0.46
55:DA:2469:A:N1	55:DA:2481:G:C2	2.84	0.46
55:DA:2592:G:C5	55:DA:2593:U:C5	3.04	0.46
55:DA:2655:G:O2'	55:DA:2656:U:OP2	2.33	0.46
55:DA:738:G:C6	55:DA:739:G:C2	3.04	0.46
3:DD:36:PRO:HB3	3:DD:62:TYR:O	2.16	0.46
4:DE:14:ILE:O	4:DE:15:PHE:HD2	1.89	0.46
8:DK:52:ARG:NH1	8:DK:52:ARG:CB	2.79	0.46
58:DL:103:GLN:O	58:DL:107:ILE:CG1	2.64	0.46
58:DL:112:MET:HE1	58:DL:123:ALA:HB2	1.94	0.46
58:DL:62:ASP:OD2	58:DL:63:ARG:N	2.48	0.46
55:DA:1666:G:O3'	10:DN:6:THR:HG23	2.15	0.46
10:DN:86:ILE:HG21	10:DN:94:ARG:HD2	1.96	0.46
11:DO:92:GLU:HG3	11:DO:121:LYS:HD2	1.97	0.46
12:DP:38:GLU:OE2	12:DP:128:LYS:HG3	2.15	0.46
14:DQ:7:TYR:HA	14:DQ:10:ARG:HH11	1.81	0.46
55:DA:1335:U:OP1	19:DT:65:ARG:NE	2.48	0.46
20:DU:51:VAL:CG2	20:DU:57:GLN:HA	2.46	0.46
24:DW:17:SER:HB2	24:DW:18:PRO:CA	2.46	0.46
57:DY:120:LYS:HD3	57:DY:120:LYS:HA	1.63	0.46
57:DY:12:THR:C	57:DY:14:LYS:H	2.19	0.46
13:A0:30:THR:HA	13:A0:78:LYS:NZ	2.31	0.46
13:A0:92:GLY:O	13:A0:94:TYR:CD1	2.69	0.46
2:AB:39:A:C2'	26:A4:1:MET:HE2	2.45	0.46
30:A8:47:LYS:C	30:A8:48:PHE:CD1	2.89	0.46
1:AA:1300:U:H2'	1:AA:1635:G:OP1	2.16	0.46
1:AA:1372:U:C3'	1:AA:1372:U:C6	2.95	0.46
1:AA:1668:A:H4'	1:AA:1669:A:O5'	2.15	0.46
1:AA:195:A:H3'	1:AA:196:A:H5'	1.98	0.46
1:AA:1992:G:N2	1:AA:1996:C:O2'	2.48	0.46
1:AA:2009:G:C6	1:AA:2010:G:N7	2.84	0.46
1:AA:212:G:C2'	1:AA:213:A:H5'	2.45	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:579:G:H2'	1:AA:580:C:C6	2.50	0.46
1:AA:605:C:O2	1:AA:657:U:O2'	2.27	0.46
1:AA:704:G:C2'	1:AA:705:A:OP2	2.63	0.46
1:AA:751:A:O5'	1:AA:751:A:H8	1.98	0.46
1:AA:957:A:N1	1:AA:2459:A:C8	2.84	0.46
3:AD:8:PRO:HB3	3:AD:14:ARG:HE	1.80	0.46
3:AD:35:LYS:HD3	3:AD:63:ARG:CA	2.45	0.46
6:AG:101:ILE:HD13	26:A4:25:TYR:HB2	1.98	0.46
6:AG:51:ARG:HH11	6:AG:51:ARG:HB3	1.80	0.46
7:AH:156:ALA:HB3	7:AH:159:GLU:O	2.15	0.46
8:AK:110:ASP:OD2	8:AK:130:TYR:CE1	2.68	0.46
1:AA:953:A:OP2	12:AP:16:ARG:NH1	2.45	0.46
18:AS:95:ILE:HG13	18:AS:95:ILE:O	2.15	0.46
20:AU:81:LYS:HB3	20:AU:97:ARG:HD3	1.97	0.46
23:AZ:89:GLU:O	23:AZ:93:GLU:HB2	2.15	0.46
23:AZ:91:LYS:HA	23:AZ:91:LYS:CE	2.31	0.46
31:BA:1095:U:OP1	31:BA:1108:G:N2	2.48	0.46
31:BA:1106:G:O2'	31:BA:1107:C:H5'	2.16	0.46
31:BA:274:A:H4'	31:BA:275:G:OP1	2.16	0.46
31:BA:280:C:H1'	47:BT:38:ARG:NH1	2.29	0.46
31:BA:310:G:OP1	46:BS:27:LYS:HD3	2.16	0.46
31:BA:353:A:C2'	31:BA:354:G:OP2	2.64	0.46
31:BA:528:C:H4'	31:BA:535:A:C6	2.50	0.46
31:BA:560:U:H4'	31:BA:561:U:O5'	2.16	0.46
31:BA:685:G:H5'	41:BN:39:PRO:O	2.16	0.46
31:BA:784:C:H2'	31:BA:785:G:H8	1.81	0.46
31:BA:807:A:H2'	31:BA:808:C:H6	1.75	0.46
52:BD:18:G:C2	52:BD:57:G:O6	2.69	0.46
33:BF:167:TRP:O	33:BF:168:ALA:HB2	2.15	0.46
33:BF:95:THR:HG22	33:BF:97:LYS:CG	2.42	0.46
34:BG:64:LEU:CD1	34:BG:196:LEU:HD23	2.45	0.46
38:BK:12:ARG:HH12	38:BK:27:PRO:HD3	1.78	0.46
39:BL:114:TYR:HD1	40:BM:60:ARG:HG2	1.81	0.46
42:BO:90:VAL:O	42:BO:90:VAL:HG12	2.14	0.46
44:BQ:12:ARG:C	44:BQ:14:PRO:HD3	2.36	0.46
50:BW:20:LEU:O	50:BW:21:LYS:C	2.52	0.46
54:CA:1054:C:N4	52:CB:34:G:C8	2.84	0.46
54:CA:1228:C:OP1	43:CP:108:ARG:NH2	2.47	0.46
54:CA:748:C:OP2	54:CA:748:C:H6	1.97	0.46
54:CA:75:C:H2'	54:CA:76:G:O4'	2.16	0.46
54:CA:812:C:O2'	54:CA:813:U:OP2	2.33	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:CB:8:U:H6	52:CB:8:U:OP2	1.98	0.46
52:CC:18:G:N2	52:CC:57:G:H2'	2.31	0.46
32:CE:134:GLU:O	32:CE:137:ARG:HB3	2.15	0.46
35:CH:18:ARG:HG2	35:CH:19:MET:N	2.29	0.46
39:CL:91:ASP:C	39:CL:93:ARG:H	2.19	0.46
41:CN:43:SER:HB3	41:CN:68:ALA:HB2	1.98	0.46
43:CP:120:LYS:HD3	43:CP:120:LYS:H	1.76	0.46
43:CP:48:LEU:O	43:CP:49:THR:C	2.54	0.46
45:CR:21:ASP:OD1	45:CR:24:SER:HB2	2.16	0.46
16:D1:86:ALA:HB3	16:D1:88:ILE:CD1	2.45	0.46
16:D1:98:LEU:C	16:D1:98:LEU:CD2	2.81	0.46
55:DA:1092:C:C3'	55:DA:1092:C:C6	2.98	0.46
55:DA:1104:C:H6	55:DA:1104:C:O5'	1.98	0.46
55:DA:1188:U:H2'	55:DA:1189:A:H5'	1.97	0.46
55:DA:1509:C:C3'	55:DA:1510:A:H4'	2.44	0.46
55:DA:1938:A:H5'	55:DA:1939:U:OP2	2.15	0.46
55:DA:2631:G:H2'	55:DA:2632:A:O5'	2.15	0.46
55:DA:2655:G:N2	55:DA:2665:A:OP2	2.49	0.46
55:DA:2531:A:N6	55:DA:2662:A:H61	2.14	0.46
55:DA:2712:U:O2	55:DA:2712:U:C3'	2.63	0.46
55:DA:315:G:H2'	55:DA:316:C:C6	2.50	0.46
55:DA:405:U:C2'	55:DA:405:U:O2	2.59	0.46
55:DA:531:C:H5"	55:DA:532:A:C4	2.50	0.46
55:DA:654(J):A:H2	55:DA:654(L):G:N7	2.14	0.46
3:DD:142:VAL:HG23	3:DD:192:THR:C	2.36	0.46
3:DD:166:GLN:HA	3:DD:166:GLN:NE2	2.30	0.46
3:DD:142:VAL:HG23	3:DD:193:VAL:N	2.30	0.46
4:DE:1:MET:HB3	4:DE:200:GLU:OE1	2.15	0.46
5:DF:161:GLU:O	5:DF:164:ARG:HB3	2.15	0.46
5:DF:63:LYS:HE3	5:DF:65:TRP:O	2.16	0.46
6:DG:125:PHE:HB3	6:DG:166:ASP:HB2	1.98	0.46
7:DH:9:ILE:HG22	7:DH:51:ARG:CG	2.42	0.46
56:DI:17:VAL:CG1	56:DI:21:LYS:HE3	2.45	0.46
58:DL:68:VAL:CG2	58:DL:69:THR:N	2.78	0.46
58:DL:82:ALA:C	58:DL:84:LEU:N	2.69	0.46
9:DM:120:LEU:HD13	9:DM:120:LEU:C	2.36	0.46
10:DN:2:ILE:HG13	10:DN:8:LEU:HD11	1.97	0.46
12:DP:111:GLU:O	12:DP:115:MET:HG3	2.15	0.46
21:DV:106:GLY:O	21:DV:107:THR:OG1	2.30	0.46
21:DV:155:LEU:O	21:DV:156:LYS:C	2.54	0.46
57:DY:25:PHE:CE1	57:DY:82:PHE:C	2.89	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:54:ALA:HB3	57:DY:58:LEU:CD2	2.45	0.46
13:A0:66:VAL:CG1	13:A0:70:LEU:HD12	2.45	0.46
13:A0:55:ALA:CB	13:A0:79:LEU:HD22	2.45	0.46
16:A1:50:ARG:HH21	16:A1:50:ARG:CB	2.29	0.46
16:A1:58:ARG:O	16:A1:62:ILE:HD13	2.14	0.46
16:A1:60:LEU:HD21	16:A1:64:ARG:NH2	2.31	0.46
22:A3:55:ARG:HH11	22:A3:55:ARG:CB	2.28	0.46
26:A4:66:SER:O	26:A4:67:TYR:O	2.33	0.46
27:A5:59:GLU:O	27:A5:60:VAL:C	2.53	0.46
1:AA:2233:U:H2'	1:AA:2234:G:C8	2.50	0.46
1:AA:2282:G:H4'	1:AA:2389:G:O2'	2.15	0.46
1:AA:2391:G:O2'	1:AA:2392:A:P	2.72	0.46
1:AA:2552:U:C5'	1:AA:2553:G:OP2	2.64	0.46
1:AA:2627:G:N3	1:AA:2781:A:H2	2.14	0.46
1:AA:2883:A:H3'	1:AA:2884:U:H5'	1.96	0.46
1:AA:699:A:H2'	1:AA:700:G:O4'	2.15	0.46
1:AA:895:U:O2	1:AA:895:U:C3'	2.63	0.46
1:AA:948:G:H2'	1:AA:949:C:C6	2.50	0.46
2:AB:2:C:H2'	2:AB:3:C:C6	2.50	0.46
2:AB:87:G:H3'	2:AB:88:C:H5'	1.92	0.46
3:AD:35:LYS:HE3	3:AD:64:ILE:C	2.36	0.46
4:AE:16:ARG:HH11	4:AE:16:ARG:HG3	1.81	0.46
4:AE:7:VAL:O	4:AE:26:ILE:O	2.34	0.46
5:AF:125:LEU:HD12	5:AF:196:LEU:CD2	2.45	0.46
5:AF:84:VAL:O	5:AF:86:GLY:N	2.49	0.46
7:AH:126:PRO:O	7:AH:127:GLU:O	2.32	0.46
7:AH:12:PRO:C	7:AH:15:VAL:HG22	2.36	0.46
9:AM:10:GLU:OE2	9:AM:11:PRO:HD2	2.15	0.46
12:AP:133:ARG:HH11	12:AP:133:ARG:CG	2.29	0.46
12:AP:98:LYS:HB3	12:AP:99:PRO:CD	2.40	0.46
15:AR:126:ALA:C	15:AR:128:GLU:H	2.18	0.46
21:AV:178:GLU:OE1	21:AV:181:GLU:N	2.49	0.46
21:AV:182:LYS:NZ	52:BB:57:G:H5'	2.30	0.46
24:AW:56:GLN:O	24:AW:60:LEU:HD12	2.15	0.46
24:AW:71:ASN:O	24:AW:72:ALA:C	2.52	0.46
31:BA:1198:G:H2'	31:BA:1199:U:C6	2.51	0.46
31:BA:1254:C:OP1	40:BM:45:ARG:HA	2.16	0.46
31:BA:1270:C:OP2	51:BX:24:ARG:NH2	2.49	0.46
31:BA:1347:G:N2	31:BA:1374:A:OP2	2.44	0.46
31:BA:1442:G:N7	31:BA:1446:A:C2	2.84	0.46
31:BA:147:G:O2'	31:BA:148:G:H5'	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:228:A:H2'	31:BA:229:U:H6	1.79	0.46
31:BA:339:C:H2'	31:BA:340:U:C6	2.50	0.46
31:BA:36:C:H2'	31:BA:37:U:H5'	1.98	0.46
31:BA:4:U:C4	38:BK:105:ARG:NH1	2.83	0.46
31:BA:537:G:H5''	42:BO:113:ARG:HH12	1.80	0.46
31:BA:621:A:H2'	31:BA:622:A:C8	2.50	0.46
31:BA:623:C:C4	31:BA:624:C:C5	3.03	0.46
31:BA:577:G:C1'	31:BA:816:A:C4	2.98	0.46
31:BA:889:A:O3'	31:BA:890:G:H4'	2.15	0.46
31:BA:982:U:H6	31:BA:982:U:OP1	1.99	0.46
32:BE:86:GLU:O	32:BE:88:ALA:N	2.45	0.46
32:BE:85:ALA:HB1	32:BE:92:TYR:HB3	1.98	0.46
33:BF:136:GLN:O	33:BF:140:ARG:N	2.48	0.46
35:BH:13:ILE:O	35:BH:13:ILE:HG12	2.15	0.46
37:BJ:44:TYR:O	37:BJ:47:CYS:N	2.48	0.46
38:BK:39:LEU:O	38:BK:44:PHE:HB2	2.16	0.46
33:BF:18:TRP:CD1	44:BQ:54:PRO:HA	2.50	0.46
47:BT:59:ILE:HG22	47:BT:71:PHE:CD1	2.50	0.46
48:BU:62:GLU:O	48:BU:65:ILE:CD1	2.64	0.46
54:CA:1333:A:C2	54:CA:1334:G:H1'	2.50	0.46
54:CA:266:G:H5''	54:CA:268:C:N4	2.14	0.46
54:CA:424:G:O2'	54:CA:425:G:H5'	2.16	0.46
54:CA:481:G:O2'	54:CA:482:A:C8	2.64	0.46
54:CA:586:C:H1'	54:CA:878:G:O2'	2.16	0.46
54:CA:96:G:C8	54:CA:96:G:H5'	2.43	0.46
32:CE:213:LEU:HD23	32:CE:213:LEU:C	2.36	0.46
34:CG:152:SER:HB3	34:CG:155:LEU:HD12	1.96	0.46
35:CH:102:ALA:HB1	35:CH:106:PRO:CG	2.43	0.46
35:CH:10:MET:SD	35:CH:13:ILE:CD1	3.03	0.46
36:CI:19:LEU:HD11	36:CI:59:TYR:CE2	2.50	0.46
38:CK:115:SER:HB2	38:CK:116:LYS:HE3	1.98	0.46
42:CO:45:PRO:O	42:CO:46:LYS:O	2.34	0.46
46:CS:82:GLN:O	46:CS:83:GLU:HB3	2.16	0.46
48:CU:53:ARG:HH21	48:CU:60:ALA:N	2.12	0.46
13:D0:91:GLN:NE2	13:D0:91:GLN:H	2.13	0.46
16:D1:83:LEU:HD12	16:D1:88:ILE:HD11	1.97	0.46
29:D7:25:PRO:HA	29:D7:28:ARG:CZ	2.46	0.46
55:DA:2420:C:OP1	30:D8:34:TRP:CA	2.63	0.46
55:DA:1062:G:N2	55:DA:1077:A:C8	2.83	0.46
55:DA:1283:G:H2'	55:DA:1285:G:OP2	2.15	0.46
55:DA:1311:G:H1'	55:DA:1313:U:O4	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1480:G:O6	55:DA:1510:A:N1	2.49	0.46
55:DA:1999:C:H2'	55:DA:2000:G:O4'	2.16	0.46
55:DA:265:A:C2'	55:DA:266:G:O4'	2.64	0.46
55:DA:2680:C:H5'	4:DE:189:PRO:HA	1.98	0.46
55:DA:2836:U:H2'	55:DA:2837:G:H8	1.79	0.46
55:DA:381:G:O2'	55:DA:382:G:H5'	2.16	0.46
55:DA:492:A:H2'	55:DA:493:G:C5'	2.46	0.46
55:DA:721:C:H3'	55:DA:722:A:H8	1.81	0.46
55:DA:897:C:C6	55:DA:897:C:C4'	2.98	0.46
2:DB:41:U:O4	6:DG:70:VAL:O	2.34	0.46
2:DB:52:A:H2'	2:DB:53:A:N7	2.30	0.46
3:DD:30:GLU:CG	3:DD:63:ARG:NH2	2.78	0.46
4:DE:62:PRO:O	4:DE:63:LEU:C	2.52	0.46
4:DE:50:GLY:CA	4:DE:74:PRO:HG3	2.44	0.46
5:DF:34:TRP:CH2	11:DO:8:PRO:HG3	2.48	0.46
6:DG:26:GLN:HE21	6:DG:27:ASN:HB2	1.80	0.46
6:DG:44:GLY:O	6:DG:47:LYS:HB2	2.16	0.46
6:DG:75:LYS:HD2	6:DG:77:ILE:CD1	2.45	0.46
56:DI:10:GLU:O	56:DI:11:GLU:C	2.53	0.46
8:DK:139:GLN:NE2	8:DK:139:GLN:O	2.49	0.46
8:DK:71:ILE:C	8:DK:74:ASN:HD21	2.19	0.46
58:DL:12:LEU:HA	58:DL:12:LEU:HD13	1.42	0.46
58:DL:138:VAL:CG1	58:DL:139:VAL:H	2.25	0.46
9:DM:47:ALA:HB1	9:DM:116:LEU:HD21	1.97	0.46
9:DM:131:GLN:OE1	9:DM:132:ALA:HB2	2.15	0.46
15:DR:78:LEU:O	15:DR:78:LEU:HD13	2.16	0.46
19:DT:49:VAL:CG1	19:DT:50:LYS:N	2.78	0.46
20:DU:91:GLU:O	20:DU:92:ASN:HB3	2.16	0.46
20:DU:96:ILE:HD11	20:DU:99:CYS:SG	2.55	0.46
21:DV:194:PRO:CB	21:DV:196:VAL:CG1	2.92	0.46
21:DV:61:LEU:HD22	21:DV:62:PRO:O	2.15	0.46
57:DY:25:PHE:CA	57:DY:82:PHE:CE2	2.99	0.46
28:A6:25:LYS:HB2	28:A6:27:LYS:CD	2.46	0.46
30:A8:8:LYS:O	30:A8:12:LYS:HG3	2.15	0.46
1:AA:1235:G:N1	1:AA:1236:G:N2	2.64	0.46
1:AA:1665:A:C2'	1:AA:1666:G:H5'	2.46	0.46
1:AA:1798:U:O2	1:AA:1802:A:H2	1.98	0.46
1:AA:2168:G:C2'	1:AA:2168:G:N3	2.78	0.46
1:AA:2219:G:C2'	1:AA:2224:G:H5'	2.45	0.46
1:AA:2580:U:H4'	4:AE:130:GLY:HA3	1.95	0.46
1:AA:270(B):A:C5'	1:AA:270(C):C:OP2	2.64	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:273(B):C:H2'	1:AA:273(C):C:C6	2.51	0.46
1:AA:2867:G:H2'	1:AA:2868:A:OP2	2.16	0.46
1:AA:434:U:C4'	1:AA:435:C:OP1	2.50	0.46
1:AA:529:A:H2'	1:AA:529:A:N3	2.30	0.46
1:AA:621:A:H2'	1:AA:622:G:C5'	2.45	0.46
1:AA:654(S):G:H2'	1:AA:654(T):A:N9	2.30	0.46
1:AA:669:G:N3	1:AA:669:G:C2'	2.76	0.46
1:AA:806:C:OP2	11:AO:41:ARG:NH2	2.48	0.46
3:AD:16:MET:HG3	3:AD:206:LEU:O	2.15	0.46
5:AF:84:VAL:C	5:AF:86:GLY:H	2.19	0.46
6:AG:115:ARG:O	6:AG:116:ASP:HB2	2.15	0.46
7:AH:149:ARG:C	7:AH:151:ILE:H	2.19	0.46
7:AH:20:ALA:HB3	7:AH:23:ARG:CG	2.46	0.46
8:AK:122:GLU:O	8:AK:126:TYR:OH	2.29	0.46
8:AK:68:LEU:HA	8:AK:71:ILE:CG2	2.45	0.46
8:AK:88:ILE:HG12	8:AK:121:LYS:C	2.36	0.46
9:AM:28:THR:HA	9:AM:106:MET:HE1	1.98	0.46
9:AM:9:VAL:HG11	9:AM:39:ARG:NH1	2.30	0.46
10:AN:8:LEU:HD12	10:AN:84:ALA:HB2	1.97	0.46
12:AP:35:VAL:HG11	12:AP:130:LYS:HD3	1.97	0.46
14:AQ:88:ASP:CG	14:AQ:89:ARG:H	2.19	0.46
15:AR:70:VAL:HG12	15:AR:71:GLY:N	2.31	0.46
21:AV:148:ASP:OD1	21:AV:174:VAL:HG23	2.16	0.46
21:AV:7:ALA:HB2	21:AV:39:VAL:HG12	1.98	0.46
25:AX:42:ALA:O	25:AX:43:ILE:C	2.52	0.46
53:B1:53:U:O2'	53:B1:54:U:P	2.73	0.46
31:BA:1028:C:C4	31:BA:1028(A):C:C5	3.03	0.46
31:BA:1060:C:O2'	31:BA:1061:G:H5'	2.16	0.46
31:BA:1298:C:H5''	37:BJ:114:ARG:HH12	1.81	0.46
31:BA:1323:G:H2'	31:BA:1324:A:C8	2.51	0.46
31:BA:1234:C:H5'	31:BA:1364:U:O2'	2.16	0.46
31:BA:979:C:O2	44:BQ:19:ARG:HG2	2.15	0.46
52:BB:34:G:O2'	52:BB:35:A:H5'	2.15	0.46
52:BC:75:C:H5''	52:BC:76:A:OP2	2.16	0.46
32:BE:16:HIS:NE2	32:BE:209:ARG:HD2	2.30	0.46
32:BE:219:VAL:HA	32:BE:222:ILE:CD1	2.42	0.46
32:BE:67:THR:HA	32:BE:90:MET:SD	2.55	0.46
32:BE:95:GLN:O	32:BE:96:ARG:C	2.54	0.46
34:BG:33:MET:C	34:BG:34:GLU:O	2.48	0.46
38:BK:36:LEU:O	38:BK:39:LEU:N	2.49	0.46
39:BL:118:LYS:O	39:BL:119:ALA:HB3	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:BO:117:ARG:CG	42:BO:117:ARG:HH11	2.29	0.46
43:BP:11:ARG:O	43:BP:13:LYS:N	2.47	0.46
43:BP:34:LEU:CD1	43:BP:41:PRO:HB3	2.46	0.46
44:BQ:61:TRP:CG	44:BQ:61:TRP:OXT	2.68	0.46
46:BS:19:ILE:N	46:BS:37:GLY:O	2.44	0.46
31:BA:191(F):U:H3	50:BW:105:SER:HB3	1.81	0.46
50:BW:26:ASN:HA	50:BW:29:LYS:CG	2.46	0.46
35:CH:15:ARG:HD3	53:C1:56:U:OP2	2.16	0.46
54:CA:1155:G:O2'	54:CA:1156:G:H5'	2.16	0.46
54:CA:1324:A:H2'	54:CA:1325:C:C6	2.51	0.46
54:CA:336:C:O2'	54:CA:337:C:H5'	2.15	0.46
54:CA:529:G:O6	42:CO:49:ASN:ND2	2.49	0.46
54:CA:556:C:H2'	54:CA:557:G:H5'	1.96	0.46
54:CA:723:U:O2	54:CA:723:U:H2'	2.16	0.46
54:CA:956:U:H5'	49:CV:87:ALA:HB2	1.98	0.46
32:CE:212:GLN:C	32:CE:212:GLN:CD	2.75	0.46
32:CE:239:VAL:HG12	32:CE:240:GLN:OE1	2.16	0.46
32:CE:55:PHE:CD1	32:CE:221:LEU:HG	2.51	0.46
33:CF:45:LYS:O	33:CF:48:TYR:HB3	2.16	0.46
34:CG:188:LEU:CD2	34:CG:189:PRO:HD2	2.44	0.46
38:CK:97:VAL:O	38:CK:100:ILE:HG13	2.15	0.46
38:CK:34:GLU:CB	38:CK:118:VAL:HG21	2.38	0.46
38:CK:83:ILE:HG23	38:CK:83:ILE:O	2.16	0.46
40:CM:48:THR:HG23	40:CM:62:HIS:ND1	2.31	0.46
54:CA:502:G:P	42:CO:118:SER:HB2	2.56	0.46
42:CO:109:GLY:CA	42:CO:121:GLY:O	2.56	0.46
42:CO:19:ARG:HH11	42:CO:19:ARG:CB	2.28	0.46
47:CT:13:ASP:O	47:CT:15:MET:N	2.48	0.46
17:D2:48:GLY:O	17:D2:49:THR:C	2.54	0.46
55:DA:2355:C:C5'	22:D3:36:ILE:HD11	2.45	0.46
26:D4:37:SER:O	26:D4:38:LYS:O	2.34	0.46
28:D6:38:LYS:HA	28:D6:48:VAL:O	2.15	0.46
11:DO:62:LEU:CD2	30:D8:25:MET:HB2	2.36	0.46
55:DA:1287:A:OP1	13:D0:105:ARG:O	2.34	0.46
55:DA:1533:C:C5'	55:DA:1534:G:OP2	2.58	0.46
55:DA:165:U:O2	55:DA:165:U:C2'	2.64	0.46
55:DA:1810:A:H2'	55:DA:1811:G:O4'	2.16	0.46
55:DA:1944:U:H5''	55:DA:1945:G:OP2	2.16	0.46
55:DA:2152:G:H2'	55:DA:2153:G:C8	2.50	0.46
55:DA:2469:A:N6	55:DA:2481:G:H1'	2.30	0.46
55:DA:2506:U:O2'	55:DA:2507:C:H5'	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:554:U:O2'	55:DA:556:G:H8	1.97	0.46
55:DA:638:G:H2'	55:DA:639:U:H6	1.72	0.46
55:DA:879:G:C4	55:DA:880:G:H1'	2.51	0.46
6:DG:77:ILE:HG22	6:DG:80:PHE:N	2.27	0.46
7:DH:89:ILE:HD12	7:DH:129:THR:O	2.16	0.46
7:DH:26:VAL:CG1	7:DH:33:LEU:HB2	2.45	0.46
8:DK:78:THR:O	8:DK:79:ILE:CB	2.63	0.46
55:DA:1058:U:O3'	58:DL:4:VAL:CG1	2.64	0.46
9:DM:46:VAL:O	9:DM:47:ALA:HB3	2.15	0.46
11:DO:30:THR:OG1	11:DO:31:ALA:N	2.47	0.46
11:DO:41:ARG:HD3	11:DO:41:ARG:HA	1.65	0.46
14:DQ:15:ARG:NE	14:DQ:88:ASP:OD1	2.48	0.46
20:DU:48:ALA:HB2	20:DU:61:ILE:HD13	1.98	0.46
24:DW:42:GLY:C	24:DW:44:LEU:H	2.07	0.46
16:A1:47:TYR:HA	16:A1:50:ARG:HH12	1.81	0.46
28:A6:29:ASN:HD22	28:A6:29:ASN:N	2.12	0.46
1:AA:1019:U:H2'	1:AA:1020:A:C8	2.50	0.46
1:AA:1301:A:O2'	1:AA:1302:A:C3'	2.63	0.46
1:AA:1428:C:C5	1:AA:1569:A:H5"	2.51	0.46
1:AA:1432:C:H2'	1:AA:1433:U:O4'	2.15	0.46
1:AA:1814:G:C2	1:AA:1815:A:N6	2.84	0.46
1:AA:1820:U:C2	3:AD:202:LYS:HB3	2.50	0.46
1:AA:981:A:H2	1:AA:2027:G:N3	2.13	0.46
1:AA:2391:G:HO2'	1:AA:2392:A:P	2.38	0.46
1:AA:2556:C:C2'	1:AA:2557:G:H5'	2.46	0.46
1:AA:2661:G:H2'	1:AA:2662:A:C8	2.51	0.46
1:AA:2777:G:H4'	1:AA:2778:A:H5'	1.98	0.46
1:AA:818:G:O2'	1:AA:838:C:O2'	2.25	0.46
1:AA:907:U:OP1	12:AP:24:GLY:C	2.53	0.46
1:AA:913:U:O2'	1:AA:914:C:O5'	2.34	0.46
1:AA:2239:G:P	3:AD:244:ARG:HH22	2.38	0.46
3:AD:63:ARG:O	3:AD:65:ILE:HG22	2.15	0.46
1:AA:2822:G:OP1	4:AE:112:GLY:HA2	2.16	0.46
5:AF:20:LEU:HD22	5:AF:23:ASP:OD2	2.15	0.46
7:AH:91:GLY:O	7:AH:94:TYR:HB2	2.15	0.46
8:AK:129:THR:HG22	8:AK:137:PRO:CB	2.34	0.46
9:AM:98:VAL:CG2	9:AM:99:LEU:N	2.78	0.46
1:AA:955:C:H5'	12:AP:14:ARG:HH21	1.81	0.46
1:AA:2485:G:H5"	12:AP:46:GLN:NE2	2.31	0.46
12:AP:93:TYR:CD1	12:AP:93:TYR:N	2.83	0.46
14:AQ:88:ASP:OD2	14:AQ:89:ARG:N	2.49	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:AR:90:GLN:CG	15:AR:91:ARG:N	2.79	0.46
18:AS:5:ALA:HB3	18:AS:54:ALA:HB2	1.97	0.46
20:AU:68:HIS:O	20:AU:71:LYS:CG	2.64	0.46
21:AV:26:GLY:C	21:AV:37:VAL:HG22	2.36	0.46
24:AW:59:ARG:O	24:AW:60:LEU:C	2.55	0.46
23:AZ:73:LEU:HB3	23:AZ:90:ILE:CD1	2.46	0.46
31:BA:1162:C:C2	31:BA:1175:G:N2	2.83	0.46
31:BA:1492:A:H1'	53:B1:50:U:HO2'	1.78	0.46
31:BA:1512:U:H2'	31:BA:1513:A:H8	1.80	0.46
31:BA:663:A:H2'	31:BA:664:G:O4'	2.15	0.46
31:BA:901:A:C5	31:BA:902:G:H1'	2.50	0.46
31:BA:956:U:H2'	31:BA:957:U:O4'	2.15	0.46
52:BB:16:U:H2'	52:BB:18:G:OP2	2.15	0.46
52:BC:1:G:C4	52:BC:2:C:C5	3.04	0.46
32:BE:160:ASP:O	32:BE:161:ALA:HB2	2.15	0.46
32:BE:218:ALA:O	32:BE:221:LEU:HG	2.15	0.46
32:BE:68:ILE:O	32:BE:90:MET:HB3	2.16	0.46
32:BE:72:GLY:C	32:BE:74:LYS:H	2.19	0.46
32:BE:87:ARG:HD2	32:BE:87:ARG:O	2.15	0.46
33:BF:14:ILE:O	33:BF:15:THR:HB	2.15	0.46
33:BF:22:TRP:CZ3	33:BF:24:ALA:HB2	2.51	0.46
33:BF:33:LEU:C	33:BF:35:GLU:N	2.69	0.46
33:BF:59:ARG:CG	33:BF:64:VAL:HG22	2.44	0.46
31:BA:426:G:P	34:BG:36:ARG:HH21	2.38	0.46
34:BG:38:TYR:CE1	34:BG:45:GLN:HB3	2.50	0.46
35:BH:19:MET:CE	35:BH:24:ARG:HB3	2.45	0.46
35:BH:64:ARG:HH11	35:BH:64:ARG:HG3	1.81	0.46
37:BJ:95:ARG:HG3	37:BJ:95:ARG:NH1	2.31	0.46
38:BK:60:ARG:HB2	38:BK:60:ARG:NH1	2.31	0.46
43:BP:91:ARG:NH2	43:BP:103:THR:HG21	2.28	0.46
43:BP:23:TYR:HD1	43:BP:71:ARG:CZ	2.28	0.46
45:BR:17:ARG:NH1	45:BR:77:ARG:HH12	2.12	0.46
49:BV:8:GLY:O	49:BV:9:VAL:C	2.54	0.46
53:C1:57:U:C2'	53:C1:57:U:O2	2.64	0.46
54:CA:1027:C:HO2'	54:CA:1028:C:P	2.38	0.46
54:CA:960:U:O4	54:CA:1225:A:H1'	2.16	0.46
54:CA:321:A:C2	54:CA:333:G:C2	3.04	0.46
54:CA:344:A:C5'	54:CA:345:C:OP2	2.63	0.46
52:CC:35:A:H2'	52:CC:36:A:C5'	2.37	0.46
52:CD:8:U:C4'	52:CD:9:A:OP1	2.59	0.46
32:CE:224:GLN:C	32:CE:226:ARG:N	2.67	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:CF:101:LEU:HD23	33:CF:102:ASN:O	2.16	0.46
33:CF:153:VAL:HG22	33:CF:198:VAL:HG22	1.97	0.46
33:CF:83:ARG:C	33:CF:85:ARG:N	2.69	0.46
54:CA:542:G:H5'	34:CG:41:GLY:CA	2.45	0.46
35:CH:76:ILE:HD11	35:CH:118:ILE:CD1	2.46	0.46
35:CH:32:VAL:CG2	35:CH:58:ALA:HB1	2.45	0.46
36:CI:18:GLN:O	36:CI:21:LEU:HB2	2.16	0.46
38:CK:94:TYR:HE1	38:CK:132:GLU:HB2	1.79	0.46
39:CL:19:LEU:CD1	39:CL:85:LEU:HB3	2.45	0.46
40:CM:5:ARG:O	40:CM:98:ILE:HA	2.16	0.46
41:CN:126:ARG:HG2	41:CN:126:ARG:NH1	2.30	0.46
43:CP:108:ARG:NH1	43:CP:108:ARG:HG3	2.30	0.46
43:CP:32:GLU:O	43:CP:35:GLU:HG2	2.16	0.46
43:CP:7:VAL:CG2	6:DG:115:ARG:NH1	2.78	0.46
49:CV:3:ARG:O	49:CV:4:SER:CB	2.61	0.46
49:CV:58:VAL:O	49:CV:58:VAL:HG23	2.16	0.46
49:CV:63:THR:CG2	49:CV:65:ASN:HD21	2.28	0.46
50:CW:17:ARG:NH1	50:CW:17:ARG:HG3	2.30	0.46
50:CW:9:ASN:HD22	50:CW:9:ASN:C	2.18	0.46
13:D0:41:ALA:C	13:D0:43:GLU:H	2.19	0.46
16:D1:108:GLU:HG3	17:D2:44:LYS:CE	2.41	0.46
17:D2:30:GLY:N	17:D2:61:VAL:HG13	2.31	0.46
28:D6:19:ARG:HA	28:D6:19:ARG:HD2	1.73	0.46
55:DA:1094:U:O2'	55:DA:1096:A:OP1	2.33	0.46
55:DA:72:U:C5	55:DA:112:U:H4'	2.51	0.46
55:DA:118:A:H5'	55:DA:119:A:C8	2.49	0.46
55:DA:1231:G:H2'	55:DA:1232:G:H8	1.80	0.46
55:DA:2063:C:C4	55:DA:2064:C:C4	3.04	0.46
55:DA:2134:A:N6	55:DA:2157:G:C1'	2.76	0.46
55:DA:2199:A:H8	55:DA:2199:A:OP2	1.99	0.46
55:DA:2562:U:H4'	10:DN:25:LEU:CD2	2.46	0.46
55:DA:2629:A:O2'	55:DA:2630:G:C5'	2.64	0.46
55:DA:343:C:O2	55:DA:343:C:H2'	2.15	0.46
55:DA:531:C:C5	55:DA:2035:G:C2	3.04	0.46
55:DA:884:C:N3	55:DA:885:C:C4	2.84	0.46
3:DD:12:SER:O	3:DD:16:MET:HB2	2.16	0.46
3:DD:46:GLN:H	3:DD:46:GLN:HG3	1.30	0.46
4:DE:104:VAL:HG11	4:DE:188:VAL:HG23	1.97	0.46
5:DF:103:LYS:HA	5:DF:106:ARG:CG	2.32	0.46
6:DG:34:LEU:HD22	6:DG:34:LEU:C	2.36	0.46
7:DH:151:ILE:O	7:DH:152:ARG:O	2.34	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:135:ARG:HB3	56:DJ:10:GLU:OE2	2.15	0.46
56:DJ:14:GLN:N	56:DJ:17:VAL:HG23	2.31	0.46
58:DL:101:TRP:O	58:DL:105:LEU:HG	2.16	0.46
9:DM:74:ARG:NH1	9:DM:85:ILE:HD11	2.31	0.46
9:DM:91:LEU:O	9:DM:95:PRO:HG3	2.16	0.46
11:DO:12:ALA:C	11:DO:14:LYS:H	2.19	0.46
11:DO:138:LEU:O	11:DO:140:ALA:N	2.43	0.46
11:DO:75:ILE:N	11:DO:75:ILE:CD1	2.62	0.46
12:DP:20:ALA:HB3	21:DV:79:ARG:CZ	2.46	0.46
12:DP:21:THR:HB	12:DP:22:LYS:H	1.38	0.46
15:DR:104:ASN:O	15:DR:105:LEU:HB3	2.16	0.46
20:DU:94:LYS:CD	20:DU:101:LYS:HZ3	2.28	0.46
20:DU:19:LYS:O	20:DU:19:LYS:CG	2.64	0.46
12:DP:140:ALA:HB1	21:DV:123:ASP:OD1	2.16	0.46
21:DV:63:ASP:O	21:DV:65:GLN:N	2.48	0.46
21:DV:82:ARG:HG3	21:DV:83:PRO:CD	2.46	0.46
57:DY:104:ILE:CB	57:DY:105:PRO:CD	2.94	0.46
57:DY:104:ILE:HD12	57:DY:104:ILE:HA	1.77	0.46
57:DY:5:ARG:O	57:DY:7:VAL:N	2.49	0.46
16:A1:105:VAL:O	16:A1:106:PHE:C	2.54	0.46
17:A2:79:VAL:O	17:A2:80:GLN:CB	2.63	0.46
27:A5:46:CYS:SG	27:A5:48:GLU:HG2	2.56	0.46
29:A7:24:THR:C	29:A7:26:GLY:N	2.68	0.46
1:AA:1171:G:H1'	1:AA:1173:G:C5'	2.46	0.46
1:AA:1880:C:H2'	1:AA:1881:C:O4'	2.15	0.46
1:AA:1924:C:C2	1:AA:1925:C:H6	2.25	0.46
1:AA:2328:A:H2'	1:AA:2329:G:H8	1.80	0.46
1:AA:2552:U:H5'	1:AA:2553:G:OP2	2.16	0.46
1:AA:2631:G:N3	1:AA:2810:A:C2	2.81	0.46
1:AA:2707:G:H5''	13:A0:68:ARG:CZ	2.46	0.46
1:AA:396:G:O3'	23:AZ:44:PRO:HA	2.16	0.46
1:AA:644:A:O3'	1:AA:645:C:H6	1.99	0.46
1:AA:654(H):G:H3'	1:AA:654(I):C:C5'	2.46	0.46
1:AA:5:A:H2'	1:AA:6:A:H8	1.81	0.46
1:AA:196:A:C2'	1:AA:805:G:O6	2.61	0.46
5:AF:22:ALA:O	5:AF:24:LEU:N	2.48	0.46
6:AG:95:ARG:O	6:AG:96:ARG:O	2.34	0.46
10:AN:13:ASN:ND2	10:AN:97:ARG:HB2	2.31	0.46
11:AO:85:LEU:HD22	11:AO:115:LEU:O	2.16	0.46
11:AO:12:ALA:C	11:AO:14:LYS:H	2.18	0.46
12:AP:139:GLU:N	12:AP:139:GLU:OE1	2.48	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:AB:37:C:O2	14:AQ:95:HIS:NE2	2.49	0.46
18:AS:92:ARG:HG2	18:AS:92:ARG:NH1	2.31	0.46
21:AV:157:LEU:C	21:AV:161:VAL:HG21	2.37	0.46
21:AV:82:ARG:HH11	21:AV:82:ARG:HG2	1.81	0.46
24:AW:26:ARG:HB3	24:AW:26:ARG:NH1	2.29	0.46
25:AX:31:LEU:C	25:AX:33:GLN:H	2.19	0.46
23:AZ:50:ARG:HA	23:AZ:58:ILE:O	2.16	0.46
23:AZ:76:ARG:CG	23:AZ:94:LEU:HD13	2.43	0.46
31:BA:1179:A:OP2	39:BL:93:ARG:NH2	2.47	0.46
31:BA:1317:C:N4	44:BQ:19:ARG:HH21	2.13	0.46
31:BA:181:G:HO2'	31:BA:182:U:H6	1.62	0.46
31:BA:191(F):U:H2'	31:BA:191:G:H5'	1.98	0.46
31:BA:392:G:H2'	31:BA:393:A:H8	1.81	0.46
31:BA:481:G:H2'	31:BA:483:C:H41	1.81	0.46
31:BA:793:U:O4	31:BA:1517:G:H5'	2.16	0.46
31:BA:965:A:H5''	31:BA:966:G:OP1	2.15	0.46
52:BC:44:G:H3'	52:BC:45:U:H6	1.78	0.46
32:BE:19:HIS:HD1	32:BE:204:ASN:ND2	2.14	0.46
32:BE:19:HIS:O	32:BE:20:GLU:C	2.53	0.46
32:BE:4:GLU:O	32:BE:6:THR:N	2.48	0.46
33:BF:15:THR:HG22	33:BF:16:ARG:HH12	1.80	0.46
34:BG:181:MET:O	34:BG:182:LYS:HB3	2.15	0.46
42:BO:86:ARG:HB2	42:BO:101:VAL:HG23	1.97	0.46
31:BA:1368:G:H4'	44:BQ:61:TRP:HZ2	1.80	0.46
48:BU:36:ASN:ND2	48:BU:39:VAL:HG21	2.29	0.46
50:BW:36:LEU:CD1	50:BW:55:ILE:HG23	2.46	0.46
53:C1:39:U:H2'	53:C1:40:U:H6	1.81	0.46
53:C1:44:U:O2'	53:C1:45:U:P	2.74	0.46
54:CA:1009:G:O2'	54:CA:1010:G:H5'	2.16	0.46
54:CA:1081:G:OP1	35:CH:16:THR:OG1	2.34	0.46
54:CA:1297:C:O2'	54:CA:1298:C:H6	1.98	0.46
54:CA:411:A:C8	54:CA:413:G:H1'	2.50	0.46
54:CA:45:U:H2'	54:CA:46:G:C8	2.50	0.46
54:CA:522:C:H41	42:CO:53:ARG:HH22	1.64	0.46
54:CA:767:A:H2'	54:CA:768:A:O4'	2.16	0.46
52:CC:68:C:H2'	52:CC:69:G:H8	1.79	0.46
33:CF:189:ALA:HB3	33:CF:196:LEU:HB3	1.96	0.46
34:CG:94:LEU:HD13	34:CG:191:ARG:HH11	1.81	0.46
36:CI:89:MET:O	36:CI:90:VAL:C	2.53	0.46
38:CK:39:LEU:CD1	38:CK:111:ILE:HD11	2.46	0.46
38:CK:82:HIS:CD2	38:CK:83:ILE:N	2.83	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:CL:10:ARG:HG3	39:CL:105:ASP:HB2	1.98	0.46
40:CM:64:GLU:OE2	40:CM:66:ARG:HD2	2.15	0.46
44:CQ:4:LYS:O	44:CQ:7:ILE:HG12	2.16	0.46
45:CR:39:LEU:CD1	45:CR:56:LEU:HB2	2.46	0.46
54:CA:375:U:H4'	46:CS:17:TYR:HE2	1.76	0.46
48:CU:66:LEU:CG	48:CU:70:ILE:HD11	2.46	0.46
50:CW:16:HIS:O	50:CW:19:SER:HB3	2.16	0.46
55:DA:1649:G:N3	13:D0:107:ASP:HB2	2.30	0.46
13:D0:66:VAL:HG12	13:D0:70:LEU:HD12	1.97	0.46
16:D1:92:ARG:NH1	17:D2:11:GLN:HB2	2.31	0.46
17:D2:44:LYS:C	17:D2:46:VAL:N	2.68	0.46
55:DA:517:C:OP1	27:D5:16:ARG:NH2	2.49	0.46
28:D6:16:CYS:O	28:D6:17:LYS:HB3	2.15	0.46
28:D6:18:ARG:O	28:D6:19:ARG:O	2.32	0.46
55:DA:1126:A:H4'	55:DA:1127:A:O5'	2.16	0.46
55:DA:1639:U:O2'	55:DA:2699:C:H4'	2.16	0.46
55:DA:2167:U:C6	55:DA:2167:U:OP2	2.65	0.46
55:DA:2197:U:H2'	55:DA:2224:G:H1	1.80	0.46
55:DA:2443:C:OP1	5:DF:68:LYS:HD2	2.15	0.46
55:DA:2529:G:H5''	55:DA:2530:A:C5'	2.46	0.46
55:DA:338:G:H2'	55:DA:339:U:H6	1.81	0.46
55:DA:632:A:H2'	55:DA:633:A:C8	2.51	0.46
55:DA:865:C:C4'	55:DA:866:A:OP1	2.63	0.46
55:DA:928:G:H3'	55:DA:929:G:H8	1.81	0.46
3:DD:177:LEU:HD12	3:DD:181:GLU:CG	2.46	0.46
4:DE:34:VAL:HG21	4:DE:77:ILE:HG21	1.98	0.46
4:DE:61:ARG:O	4:DE:63:LEU:HD23	2.16	0.46
55:DA:614:U:O4	5:DF:175:THR:O	2.34	0.46
7:DH:41:MET:HE1	7:DH:64:LEU:HB2	1.98	0.46
56:DJ:17:VAL:O	56:DJ:18:LEU:HG	2.16	0.46
56:DJ:21:LYS:HD3	56:DJ:21:LYS:HA	1.49	0.46
8:DK:9:LEU:O	8:DK:10:GLU:O	2.34	0.46
58:DL:48:MET:CG	58:DL:48:MET:O	2.63	0.46
58:DL:8:VAL:N	58:DL:57:ILE:HG21	2.30	0.46
11:DO:36:LYS:CB	11:DO:40:SER:HB3	2.40	0.46
12:DP:70:PRO:HA	12:DP:94:VAL:O	2.16	0.46
14:DQ:38:GLN:HG3	14:DQ:47:THR:CG2	2.46	0.46
14:DQ:56:LEU:HG	14:DQ:58:LEU:HD22	1.98	0.46
14:DQ:5:THR:C	14:DQ:7:TYR:H	2.19	0.46
14:DQ:62:LYS:HB3	14:DQ:97:ARG:HD3	1.96	0.46
15:DR:28:VAL:HG21	15:DR:86:ILE:CD1	2.46	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:DR:84:GLN:HG3	15:DR:85:LYS:N	2.30	0.46
19:DT:35:THR:HG22	19:DT:38:GLU:OE1	2.16	0.46
20:DU:84:ARG:HH22	20:DU:97:ARG:CB	2.28	0.46
20:DU:8:LYS:O	20:DU:27:VAL:HG21	2.15	0.46
21:DV:102:LEU:C	21:DV:103:ARG:HD2	2.36	0.46
21:DV:191:VAL:O	21:DV:192:ALA:HB3	2.10	0.46
21:DV:81:ARG:HG3	21:DV:81:ARG:O	2.15	0.46
57:DY:73:GLY:CA	57:DY:112:LEU:CD1	2.73	0.46
57:DY:71:LEU:HB3	57:DY:113:GLN:HG2	1.91	0.46
13:A0:16:HIS:O	13:A0:19:ALA:HB3	2.16	0.46
13:A0:84:ALA:N	13:A0:85:PRO:CD	2.79	0.46
17:A2:1:MET:N	17:A2:16:PRO:HD3	2.31	0.46
17:A2:75:PHE:CD2	17:A2:81:TYR:CD1	3.03	0.46
17:A2:96:ILE:CG2	17:A2:99:ILE:HD11	2.46	0.46
22:A3:27:GLU:HA	22:A3:67:VAL:O	2.15	0.46
22:A3:2:ALA:C	22:A3:3:HIS:O	2.53	0.46
28:A6:17:LYS:O	28:A6:18:ARG:CB	2.64	0.46
30:A8:39:LYS:O	30:A8:40:GLU:CB	2.64	0.46
1:AA:1060:U:H1'	1:AA:1062:G:H5'	1.98	0.46
1:AA:1204:A:O2'	1:AA:1205:U:C5'	2.60	0.46
1:AA:699:A:H4'	1:AA:1634:A:N7	2.31	0.46
1:AA:1733:G:C2'	1:AA:1734:C:H5'	2.46	0.46
1:AA:1779:U:C6	1:AA:1783:A:N7	2.84	0.46
1:AA:2061:G:O2'	1:AA:2062:A:P	2.74	0.46
1:AA:2188:C:H2'	1:AA:2189:U:O4'	2.16	0.46
1:AA:2567:G:H2'	1:AA:2568:C:C6	2.51	0.46
1:AA:2654:A:O2'	1:AA:2655:G:H4'	2.16	0.46
1:AA:276:A:C4	1:AA:277:C:H5	2.33	0.46
1:AA:460:A:H3'	1:AA:461:C:H6	1.80	0.46
1:AA:49:A:C4'	1:AA:50:U:H5'	2.42	0.46
1:AA:639:U:C2	1:AA:640:C:C5	3.03	0.46
1:AA:886:C:N3	1:AA:890:A:N1	2.64	0.46
3:AD:152:GLY:O	3:AD:154:LYS:HG3	2.16	0.46
3:AD:237:GLU:O	3:AD:238:GLY:C	2.54	0.46
4:AE:58:ARG:O	4:AE:59:VAL:C	2.54	0.46
7:AH:10:PRO:HG3	7:AH:69:ARG:NE	2.31	0.46
1:AA:943:U:OP2	11:AO:36:LYS:HE3	2.16	0.46
11:AO:62:LEU:CD1	30:A8:27:THR:HG22	2.46	0.46
12:AP:137:TYR:O	12:AP:138:ASP:C	2.54	0.46
15:AR:60:THR:HG22	15:AR:77:PRO:HA	1.98	0.46
21:AV:144:LEU:HG	21:AV:174:VAL:CG2	2.45	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:AV:157:LEU:N	21:AV:158:PRO:CD	2.78	0.46
21:AV:57:ILE:CG2	21:AV:58:VAL:N	2.79	0.46
24:AW:12:GLU:HG3	24:AW:13:ALA:N	2.30	0.46
24:AW:51:ARG:HH21	24:AW:55:ARG:HH22	1.63	0.46
31:BA:1116:C:H2'	31:BA:1117:G:C5'	2.25	0.46
31:BA:1346:A:O2'	31:BA:1347:G:O4'	2.34	0.46
31:BA:388:G:H4'	31:BA:389:A:OP2	2.16	0.46
31:BA:4:U:N3	38:BK:102:ARG:NH1	2.63	0.46
31:BA:758:G:C5'	31:BA:880:C:H1'	2.46	0.46
52:BB:42:C:H2'	52:BB:43:C:H6	1.81	0.46
33:BF:18:TRP:HE1	44:BQ:53:LEU:C	2.20	0.46
33:BF:34:LEU:CB	33:BF:38:ARG:HH21	2.28	0.46
37:BJ:53:LYS:HE2	37:BJ:53:LYS:HA	1.98	0.46
38:BK:97:VAL:HA	38:BK:100:ILE:CD1	2.40	0.46
39:BL:103:THR:HG22	39:BL:105:ASP:H	1.81	0.46
47:BT:69:LYS:C	47:BT:70:ARG:HD2	2.36	0.46
48:BU:62:GLU:O	48:BU:65:ILE:HD12	2.16	0.46
49:BV:9:VAL:CG1	49:BV:10:PHE:N	2.74	0.46
54:CA:108:G:N1	50:CW:15:ARG:NH2	2.56	0.46
54:CA:1352:C:OP1	51:CX:3:LYS:NZ	2.47	0.46
54:CA:138:G:O2'	54:CA:139:G:H5'	2.15	0.46
54:CA:1454:G:OP1	50:CW:39:LYS:NZ	2.40	0.46
54:CA:28:G:H2'	54:CA:29:G:O4'	2.15	0.46
54:CA:708:C:H2'	54:CA:709:G:H8	1.81	0.46
54:CA:865:A:H5'	54:CA:1078:U:C4	2.51	0.46
52:CB:69:G:H2'	52:CB:70:G:H5''	1.98	0.46
34:CG:18:LYS:NZ	34:CG:34:GLU:HG2	2.31	0.46
46:CS:20:VAL:CG2	46:CS:34:GLU:O	2.63	0.46
46:CS:55:ARG:C	46:CS:57:ARG:N	2.68	0.46
47:CT:5:VAL:HA	47:CT:59:ILE:O	2.16	0.46
49:CV:3:ARG:HH11	49:CV:3:ARG:CG	2.29	0.46
50:CW:39:LYS:HB2	50:CW:55:ILE:HG21	1.97	0.46
13:D0:51:LEU:HD22	13:D0:66:VAL:HG13	1.98	0.46
13:D0:45:ARG:HA	13:D0:95:THR:HG21	1.98	0.46
16:D1:85:LYS:HZ2	16:D1:117:GLN:HG2	1.79	0.46
30:D8:50:LEU:O	30:D8:51:ALA:C	2.53	0.46
55:DA:1080:A:O2'	58:DL:126:MET:N	2.49	0.46
55:DA:1171:G:C6	55:DA:1174:A:C6	3.04	0.46
55:DA:1509:C:H3'	55:DA:1510:A:C5'	2.46	0.46
55:DA:1557:C:H5''	55:DA:1558:A:OP2	2.15	0.46
55:DA:163:U:OP2	55:DA:164:U:C5	2.69	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1813:G:H2'	55:DA:1814:G:H5'	1.98	0.46
55:DA:2134:A:C2	55:DA:2159:G:H1'	2.51	0.46
55:DA:2146:C:H4'	55:DA:2147:G:N7	2.31	0.46
55:DA:2228:G:C6	55:DA:2229:C:C4	3.03	0.46
55:DA:2361:A:OP1	30:D8:27:THR:OG1	2.33	0.46
55:DA:2751:G:C2	7:DH:3:ARG:CB	2.99	0.46
55:DA:2747:G:O6	55:DA:2754:U:H3'	2.16	0.46
55:DA:483:A:O5'	55:DA:484:C:H5	1.99	0.46
55:DA:536:A:OP1	16:D1:53:ARG:NH1	2.49	0.46
55:DA:687:C:H2'	55:DA:687:C:O2	2.16	0.46
55:DA:811:U:C4	11:DO:21:ARG:NH2	2.84	0.46
2:DB:106:G:O2'	2:DB:107:U:H5'	2.17	0.46
3:DD:79:VAL:HG21	3:DD:111:LEU:HD21	1.97	0.46
6:DG:98:ARG:HA	6:DG:101:ILE:HG12	1.97	0.46
6:DG:7:LEU:HD23	6:DG:7:LEU:C	2.37	0.46
7:DH:127:GLU:HG2	7:DH:128:PRO:HD3	1.98	0.46
7:DH:54:ARG:HB2	7:DH:55:PRO:HD2	1.97	0.46
58:DL:80:LYS:O	58:DL:81:ALA:HB2	2.15	0.46
55:DA:558:G:OP1	9:DM:111:PRO:HD2	2.16	0.46
10:DN:65:THR:HA	10:DN:82:ASN:HA	1.97	0.46
55:DA:1191:G:OP1	11:DO:32:THR:HB	2.16	0.46
11:DO:42:SER:O	11:DO:43:GLY:C	2.51	0.46
11:DO:38:GLN:HG2	11:DO:45:LEU:CD1	2.45	0.46
15:DR:111:ARG:O	15:DR:112:ARG:CG	2.59	0.46
15:DR:28:VAL:HG21	15:DR:86:ILE:HD11	1.98	0.46
19:DT:12:VAL:HG13	19:DT:17:ALA:CB	2.45	0.46
19:DT:29:TRP:CZ2	19:DT:76:ARG:NH2	2.84	0.46
57:DY:24:PHE:CG	57:DY:25:PHE:N	2.81	0.46
13:A0:54:LEU:HD23	13:A0:66:VAL:CG2	2.46	0.45
13:A0:92:GLY:O	13:A0:94:TYR:CE1	2.68	0.45
1:AA:449:A:C4'	16:A1:3:ARG:HH12	2.29	0.45
17:A2:16:PRO:HA	17:A2:96:ILE:O	2.15	0.45
17:A2:95:LEU:HD22	17:A2:97:LYS:HB2	1.98	0.45
26:A4:59:PHE:HB3	26:A4:60:GLN:NE2	2.31	0.45
30:A8:16:ILE:HG21	30:A8:57:ARG:HH11	1.81	0.45
1:AA:1084:A:H2'	1:AA:1085:A:C8	2.51	0.45
1:AA:1206:G:H8	1:AA:1206:G:OP2	1.99	0.45
1:AA:1274:A:N3	1:AA:1297:C:H1'	2.30	0.45
1:AA:1367:A:C5'	1:AA:1368:G:OP2	2.64	0.45
1:AA:1810:A:H2'	1:AA:1811:G:O4'	2.17	0.45
1:AA:1824:G:H1'	3:AD:254:THR:OG1	2.15	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2098:U:H2'	1:AA:2099:U:C6	2.52	0.45
1:AA:2173:A:C5	1:AA:2174:C:H1'	2.52	0.45
1:AA:2191:G:O2'	1:AA:2192:G:OP1	2.29	0.45
1:AA:2493:U:H2'	1:AA:2494:G:O4'	2.16	0.45
1:AA:270(B):A:N1	1:AA:273:G:O2'	2.35	0.45
1:AA:332:A:O2'	1:AA:333:G:P	2.73	0.45
1:AA:37:C:H4'	1:AA:451:C:OP1	2.16	0.45
1:AA:383:U:O2	1:AA:385:C:N4	2.49	0.45
1:AA:536:A:H2'	1:AA:537:C:H6	1.81	0.45
1:AA:607:U:P	5:AF:103:LYS:HG3	2.56	0.45
1:AA:839:U:H2'	1:AA:840:C:H6	1.81	0.45
1:AA:849:A:H3'	1:AA:850:C:C6	2.52	0.45
1:AA:896:A:H4'	1:AA:897:C:OP1	2.15	0.45
1:AA:952:G:C6	1:AA:966:G:C6	3.04	0.45
4:AE:51:PHE:CE2	4:AE:52:LEU:HG	2.51	0.45
4:AE:63:LEU:O	4:AE:66:HIS:CG	2.69	0.45
4:AE:89:ASP:O	4:AE:90:THR:HB	2.15	0.45
6:AG:115:ARG:HA	43:BP:7:VAL:HG11	1.98	0.45
6:AG:115:ARG:CB	6:AG:115:ARG:HH11	2.29	0.45
6:AG:61:ALA:HB2	6:AG:67:LYS:HA	1.99	0.45
7:AH:29:PRO:HG2	7:AH:30:LYS:CD	2.47	0.45
7:AH:9:ILE:HD12	7:AH:49:VAL:CG1	2.43	0.45
9:AM:15:LEU:HD13	9:AM:15:LEU:C	2.37	0.45
10:AN:13:ASN:HD21	10:AN:97:ARG:HB2	1.82	0.45
11:AO:123:LEU:HD23	11:AO:123:LEU:N	2.31	0.45
11:AO:55:ARG:HG2	11:AO:56:SER:N	2.25	0.45
1:AA:2845:G:H5''	15:AR:54:ARG:O	2.15	0.45
15:AR:85:LYS:HG3	15:AR:86:ILE:N	2.31	0.45
19:AT:12:VAL:CG1	19:AT:27:THR:O	2.64	0.45
21:AV:103:ARG:HD3	21:AV:104:PHE:O	2.16	0.45
25:AX:46:ASN:O	25:AX:49:LYS:HB3	2.16	0.45
31:BA:1378:C:O2	31:BA:1378:C:H2'	2.15	0.45
31:BA:277:C:H5''	47:BT:68:ARG:NH2	2.31	0.45
31:BA:579:G:H2'	31:BA:580:U:H6	1.80	0.45
31:BA:601:C:O2'	31:BA:602:A:H5'	2.16	0.45
31:BA:755:G:OP2	45:BR:65:ARG:HD3	2.15	0.45
31:BA:819:A:C4'	31:BA:820:U:OP2	2.63	0.45
31:BA:890:G:H2'	31:BA:906:G:O6	2.16	0.45
52:BB:22:G:O2'	52:BB:23:A:H5'	2.15	0.45
32:BE:179:LYS:HZ3	32:BE:179:LYS:HB2	1.81	0.45
32:BE:55:PHE:HD1	32:BE:221:LEU:CD2	2.30	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:BF:99:VAL:HG23	33:BF:99:VAL:O	2.15	0.45
35:BH:20:GLN:O	35:BH:21:ALA:C	2.54	0.45
37:BJ:113:GLU:HB3	37:BJ:118:VAL:HG23	1.97	0.45
38:BK:28:ALA:HA	38:BK:59:LEU:HG	1.97	0.45
39:BL:46:ALA:N	39:BL:78:LYS:HZ2	2.14	0.45
42:BO:28:LYS:C	42:BO:30:ALA:N	2.69	0.45
42:BO:27:LEU:HB2	42:BO:33:ARG:CD	2.46	0.45
43:BP:79:LYS:HE2	43:BP:82:MET:HE3	1.97	0.45
43:BP:7:VAL:HG22	43:BP:67:GLU:HG2	1.99	0.45
45:BR:87:ILE:HG22	45:BR:88:ARG:N	2.22	0.45
50:BW:50:GLU:CB	50:BW:100:ILE:HG12	2.43	0.45
54:CA:1008:C:H5'	54:CA:1008:C:H6	1.81	0.45
54:CA:1177:G:H2'	54:CA:1178:G:C4	2.50	0.45
54:CA:1343:G:H2'	54:CA:1344:C:C6	2.51	0.45
54:CA:176:C:H2'	54:CA:177:C:H6	1.81	0.45
54:CA:433:C:O2'	54:CA:434:U:H5'	2.15	0.45
54:CA:44:G:H2'	54:CA:45:U:O4'	2.15	0.45
54:CA:511:C:O2'	54:CA:512:U:P	2.73	0.45
54:CA:61:G:P	50:CW:10:LEU:HD11	2.55	0.45
54:CA:780:A:H2	54:CA:803:G:C6	2.33	0.45
54:CA:973:G:C6	54:CA:974:A:C6	3.04	0.45
52:CC:53:G:H2'	52:CC:54:U:H6	1.81	0.45
32:CE:5:ILE:CG2	32:CE:221:LEU:HA	2.46	0.45
32:CE:223:ILE:O	32:CE:226:ARG:HB3	2.16	0.45
33:CF:12:LEU:O	33:CF:16:ARG:O	2.34	0.45
33:CF:30:ARG:HB2	44:CQ:36:PHE:O	2.17	0.45
33:CF:35:GLU:O	33:CF:39:ILE:HG13	2.16	0.45
33:CF:44:GLU:O	33:CF:48:TYR:HB2	2.16	0.45
34:CG:8:VAL:C	34:CG:10:ARG:N	2.68	0.45
40:CM:4:ILE:HB	40:CM:74:ILE:CD1	2.45	0.45
42:CO:92:ASP:O	42:CO:94:PRO:HD3	2.16	0.45
43:CP:88:ARG:O	43:CP:88:ARG:HD2	2.15	0.45
44:CQ:42:ILE:O	44:CQ:43:CYS:C	2.54	0.45
46:CS:40:ASP:C	46:CS:42:ARG:N	2.69	0.45
46:CS:49:LEU:HD22	46:CS:73:LEU:HD22	1.97	0.45
47:CT:83:ASP:O	47:CT:86:GLU:HB2	2.16	0.45
49:CV:88:LYS:O	49:CV:89:ALA:C	2.54	0.45
13:D0:13:HIS:CE1	13:D0:15:SER:OG	2.68	0.45
30:D8:33:ASN:O	30:D8:35:GLN:N	2.49	0.45
55:DA:1204:A:N1	55:DA:1241:A:C2	2.84	0.45
55:DA:1444(A):A:H5'	55:DA:1445:C:OP2	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2285:C:C5	28:D6:27:LYS:CE	2.99	0.45
55:DA:2614:A:H5''	55:DA:2615:U:OP1	2.16	0.45
55:DA:2789:C:O2'	55:DA:2790:A:H4'	2.15	0.45
55:DA:2844:G:H3'	55:DA:2845:G:H8	1.80	0.45
55:DA:2862:G:H2'	55:DA:2863:C:C6	2.51	0.45
55:DA:314:A:O2'	55:DA:315:G:H5'	2.16	0.45
55:DA:818:G:N7	55:DA:1187:G:C6	2.84	0.45
2:DB:90:C:OP1	12:DP:16:ARG:CG	2.64	0.45
55:DA:1693:U:H1'	3:DD:14:ARG:NH2	2.31	0.45
3:DD:106:ILE:HD11	3:DD:196:VAL:HG13	1.98	0.45
3:DD:218:ARG:HG3	3:DD:218:ARG:HH11	1.81	0.45
3:DD:238:GLY:O	3:DD:239:ARG:C	2.51	0.45
3:DD:8:PRO:HB3	3:DD:14:ARG:HB2	1.97	0.45
4:DE:62:PRO:C	4:DE:63:LEU:HD23	2.36	0.45
5:DF:65:TRP:CB	5:DF:66:PRO:HD2	2.45	0.45
6:DG:51:ARG:HB3	6:DG:51:ARG:NH1	2.31	0.45
8:DK:74:ASN:CG	8:DK:75:LEU:H	2.18	0.45
58:DL:40:ALA:HA	58:DL:43:ALA:HB3	1.98	0.45
55:DA:1059:G:O2'	58:DL:73:PRO:CG	2.64	0.45
9:DM:94:HIS:C	9:DM:95:PRO:O	2.51	0.45
11:DO:67:MET:H	11:DO:68:GLN:NE2	2.14	0.45
55:DA:2875:C:O2'	15:DR:5:ALA:CB	2.64	0.45
21:DV:113:ALA:O	21:DV:115:GLY:N	2.49	0.45
21:DV:5:LEU:O	21:DV:6:LYS:HG3	2.15	0.45
57:DY:105:PRO:O	57:DY:106:GLN:O	2.34	0.45
57:DY:12:THR:CG2	57:DY:58:LEU:HD13	2.46	0.45
23:DZ:21:ARG:HG3	23:DZ:35:THR:CG2	2.47	0.45
16:A1:31:SER:O	16:A1:34:LYS:N	2.50	0.45
16:A1:79:PHE:C	16:A1:79:PHE:HD2	2.18	0.45
17:A2:38:LEU:HD13	17:A2:55:ALA:HB1	1.97	0.45
27:A5:16:ARG:HD2	27:A5:20:ARG:NH2	2.31	0.45
1:AA:1255:U:H5'	1:AA:1256:G:H5''	1.99	0.45
1:AA:1555:G:O2'	1:AA:1556:C:H5'	2.17	0.45
1:AA:1608:A:HO2'	1:AA:1610:A:P	2.40	0.45
1:AA:1728:G:H8	1:AA:1732:A:N6	2.15	0.45
1:AA:1746:G:O2'	1:AA:1747:G:H5'	2.16	0.45
1:AA:270(N):G:H1'	1:AA:270(P):C:C1'	2.46	0.45
1:AA:2820:A:C2	13:A0:4:LEU:HD21	2.52	0.45
1:AA:304:G:H2'	1:AA:305:U:C6	2.51	0.45
1:AA:357:A:H2'	1:AA:358:U:C6	2.51	0.45
1:AA:594:U:H2'	1:AA:595:C:C6	2.51	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:839:U:H1'	1:AA:1191:G:H1'	1.98	0.45
1:AA:918:A:C6	1:AA:919:G:H1'	2.51	0.45
2:AB:41:U:C4	6:AG:70:VAL:O	2.69	0.45
1:AA:2591:C:P	3:AD:239:ARG:HG3	2.57	0.45
5:AF:161:GLU:O	5:AF:165:ARG:N	2.42	0.45
5:AF:75:HIS:CE1	5:AF:82:ILE:HD11	2.51	0.45
6:AG:125:PHE:C	6:AG:127:GLY:H	2.19	0.45
7:AH:153:LYS:O	7:AH:155:SER:N	2.49	0.45
8:AK:79:ILE:O	8:AK:79:ILE:HG22	2.16	0.45
11:AO:34:GLY:O	11:AO:35:HIS:C	2.54	0.45
12:AP:2:LEU:N	12:AP:2:LEU:HD12	2.30	0.45
12:AP:11:LYS:HE3	12:AP:86:GLY:O	2.17	0.45
14:AQ:19:LYS:O	14:AQ:20:ARG:CB	2.56	0.45
20:AU:75:ILE:HG13	20:AU:79:CYS:CA	2.32	0.45
21:AV:106:GLY:O	21:AV:107:THR:OG1	2.31	0.45
21:AV:44:PHE:CZ	21:AV:86:VAL:HG11	2.50	0.45
24:AW:51:ARG:HH21	24:AW:55:ARG:NH1	2.11	0.45
31:BA:1067:A:C4'	31:BA:1068:G:O5'	2.65	0.45
31:BA:1286:A:H5''	51:BX:25:LYS:CD	2.45	0.45
31:BA:1502:A:C2	31:BA:1505:G:N2	2.82	0.45
31:BA:553:A:H2'	31:BA:554:C:O4'	2.17	0.45
32:BE:221:LEU:HA	32:BE:224:GLN:HG2	1.97	0.45
33:BF:16:ARG:HB2	33:BF:16:ARG:NH1	2.31	0.45
31:BA:972:C:O3'	40:BM:57:LYS:HG2	2.17	0.45
41:BN:85:ARG:HE	41:BN:111:ASP:HB3	1.80	0.45
31:BA:376:G:OP1	46:BS:67:THR:HG21	2.15	0.45
48:BU:25:THR:HG22	48:BU:42:ARG:HH12	1.81	0.45
54:CA:1171:G:H2'	54:CA:1172:C:H6	1.79	0.45
54:CA:233:C:H2'	54:CA:234:C:H6	1.80	0.45
54:CA:300:A:H2'	54:CA:301:G:O5'	2.17	0.45
54:CA:335:C:H2'	54:CA:336:C:C6	2.51	0.45
54:CA:44:G:C2	54:CA:45:U:H1'	2.52	0.45
54:CA:974:A:OP2	44:CQ:29:ARG:NH2	2.39	0.45
52:CB:66:U:H2'	52:CB:67:C:C6	2.51	0.45
32:CE:115:LEU:O	32:CE:119:GLU:HB2	2.16	0.45
32:CE:178:ARG:NH2	38:CK:68:ARG:HH22	2.14	0.45
32:CE:20:GLU:HG3	32:CE:191:ASP:HB2	1.98	0.45
32:CE:23:ARG:HG2	32:CE:23:ARG:HH11	1.81	0.45
33:CF:188:LEU:HB3	33:CF:189:ALA:H	1.49	0.45
33:CF:5:ILE:HD13	33:CF:5:ILE:N	2.32	0.45
34:CG:10:ARG:HG2	34:CG:11:LEU:HG	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:CG:4:TYR:HE2	34:CG:7:PRO:O	1.99	0.45
36:CI:67:MET:SD	36:CI:75:LEU:HD13	2.57	0.45
37:CJ:12:LEU:H	37:CJ:12:LEU:CD2	2.26	0.45
37:CJ:85:TYR:CE1	37:CJ:154:TYR:HE1	2.35	0.45
38:CK:65:TYR:N	38:CK:65:TYR:CD1	2.84	0.45
32:CE:178:ARG:HH21	38:CK:68:ARG:HH22	1.64	0.45
39:CL:43:ALA:C	39:CL:45:ALA:H	2.19	0.45
40:CM:13:HIS:ND1	40:CM:14:LYS:N	2.64	0.45
41:CN:59:TYR:CE2	41:CN:63:LEU:HD11	2.51	0.45
43:CP:49:THR:C	43:CP:51:ALA:N	2.67	0.45
44:CQ:13:THR:O	44:CQ:14:PRO:O	2.35	0.45
45:CR:69:TYR:CE1	45:CR:73:GLU:HG3	2.51	0.45
48:CU:22:VAL:O	48:CU:23:LYS:C	2.53	0.45
16:D1:105:VAL:HG22	17:D2:44:LYS:HG2	1.97	0.45
30:D8:48:PHE:N	30:D8:48:PHE:CD1	2.82	0.45
55:DA:1059:G:O5'	55:DA:1060:U:OP2	2.34	0.45
55:DA:1710:C:H2'	55:DA:1711:C:H6	1.81	0.45
55:DA:1725:G:O2'	55:DA:1726:G:H5'	2.16	0.45
55:DA:2023:G:H4'	55:DA:2617:C:O3'	2.16	0.45
55:DA:2134:A:H62	55:DA:2157:G:C1'	2.14	0.45
55:DA:2173:A:C4	55:DA:2174:C:C1'	2.99	0.45
55:DA:2552:U:C2	55:DA:2554:U:H5'	2.51	0.45
55:DA:2729:G:N3	4:DE:187:ALA:HB2	2.32	0.45
55:DA:92:G:H2'	55:DA:93:C:H6	1.81	0.45
2:DB:69:G:H2'	2:DB:70:C:C6	2.47	0.45
58:DL:11:GLN:NE2	58:DL:18:THR:HA	2.31	0.45
9:DM:26:LEU:CG	9:DM:30:ILE:HD11	2.46	0.45
11:DO:127:ALA:CA	11:DO:147:LEU:HD23	2.45	0.45
14:DQ:85:VAL:HG23	14:DQ:112:PHE:CE1	2.51	0.45
10:DN:80:ASP:OD2	15:DR:71:GLY:HA3	2.16	0.45
20:DU:9:LYS:HE3	20:DU:28:LYS:O	2.16	0.45
20:DU:5:MET:HG2	20:DU:35:TYR:CE2	2.51	0.45
24:DW:31:GLU:HB2	24:DW:53:LEU:HD11	1.97	0.45
57:DY:23:SER:O	57:DY:24:PHE:CB	2.56	0.45
57:DY:74:LEU:N	57:DY:74:LEU:CD1	2.79	0.45
57:DY:49:ALA:CA	57:DY:84:GLU:HB2	2.45	0.45
16:A1:110:VAL:O	16:A1:113:ALA:HB3	2.16	0.45
26:A4:24:THR:HG22	26:A4:25:TYR:H	1.80	0.45
28:A6:47:THR:OG1	28:A6:48:VAL:N	2.49	0.45
1:AA:107:C:C2	1:AA:108:U:C5	3.03	0.45
1:AA:1195:G:O2'	1:AA:1196:C:H5'	2.15	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1288:U:H2'	1:AA:1327:C:O2	2.16	0.45
1:AA:1358:G:H2'	1:AA:1359:A:OP2	2.16	0.45
1:AA:1528:A:C6	1:AA:1545:A:N1	2.84	0.45
1:AA:1946:U:H2'	1:AA:1947:C:H6	1.79	0.45
1:AA:1991:U:O2'	1:AA:1992:G:H5''	2.17	0.45
1:AA:2112:G:C2'	1:AA:2113:U:H5''	2.46	0.45
1:AA:2879:C:H5'	1:AA:2880:C:OP1	2.15	0.45
1:AA:530:G:C2'	1:AA:531:C:OP2	2.64	0.45
1:AA:69:C:O2	1:AA:69:C:H2'	2.16	0.45
2:AB:43:C:P	6:AG:67:LYS:NZ	2.90	0.45
2:AB:96:G:C6	2:AB:97:G:N7	2.85	0.45
4:AE:81:ILE:O	4:AE:82:ARG:HB3	2.17	0.45
5:AF:153:SER:HB2	5:AF:190:GLU:N	2.30	0.45
5:AF:17:ARG:HD3	5:AF:17:ARG:C	2.36	0.45
6:AG:60:LEU:C	6:AG:62:LEU:N	2.70	0.45
6:AG:63:ILE:HD11	6:AG:155:MET:HE1	1.99	0.45
7:AH:50:VAL:O	7:AH:50:VAL:HG22	2.16	0.45
8:AK:1:MET:O	8:AK:3:VAL:HG13	2.16	0.45
8:AK:66:GLU:HA	8:AK:66:GLU:OE1	2.16	0.45
11:AO:11:GLY:O	11:AO:12:ALA:CB	2.64	0.45
12:AP:85:LYS:HG3	12:AP:86:GLY:H	1.79	0.45
18:AS:14:PRO:HB3	18:AS:18:ARG:NH2	2.29	0.45
18:AS:20:VAL:HG23	18:AS:47:VAL:HG21	1.98	0.45
25:AX:4:LEU:HD21	25:AX:56:VAL:CG1	2.43	0.45
35:BH:15:ARG:HH22	53:B1:55:U:P	2.40	0.45
31:BA:1003:G:N3	31:BA:1004:A:O3'	2.49	0.45
31:BA:1238:A:C8	31:BA:1303:C:H1'	2.51	0.45
31:BA:1388:C:H2'	31:BA:1389:C:H6	1.81	0.45
31:BA:1452:C:C2'	31:BA:1453:G:OP2	2.64	0.45
31:BA:228:A:H2'	31:BA:229:U:C6	2.51	0.45
31:BA:339:C:H2'	31:BA:340:U:H6	1.80	0.45
31:BA:36:C:C2'	31:BA:37:U:H5'	2.46	0.45
31:BA:393:A:H5'	31:BA:483:C:O2'	2.17	0.45
31:BA:485:G:C2'	31:BA:486:U:OP2	2.64	0.45
31:BA:513:C:H2'	31:BA:514:C:C6	2.52	0.45
34:BG:76:ARG:O	34:BG:79:PHE:HB3	2.17	0.45
37:BJ:76:ARG:HG2	37:BJ:76:ARG:NH1	2.31	0.45
39:BL:113:LYS:CD	39:BL:113:LYS:N	2.80	0.45
6:AG:115:ARG:HA	43:BP:7:VAL:CG1	2.46	0.45
50:BW:26:ASN:HB3	50:BW:71:THR:OG1	2.16	0.45
54:CA:1006:C:O2'	54:CA:1007:C:H5'	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1037:C:H2'	54:CA:1038:C:H6	1.80	0.45
54:CA:1145:C:C5'	54:CA:1146:A:OP1	2.64	0.45
54:CA:277:C:H5''	47:CT:68:ARG:NH2	2.31	0.45
54:CA:533:A:H4'	54:CA:534:U:OP1	2.16	0.45
54:CA:591:U:OP2	38:CK:30:ARG:NH1	2.48	0.45
54:CA:626:U:H2'	54:CA:627:G:H8	1.81	0.45
54:CA:657:G:C2	54:CA:658:G:C8	3.04	0.45
54:CA:674:G:H2'	54:CA:675:A:H8	1.80	0.45
52:CC:65:G:H2'	52:CC:66:U:C6	2.51	0.45
32:CE:78:GLN:HB3	32:CE:94:ASN:HD22	1.81	0.45
36:CI:14:LEU:HB3	36:CI:18:GLN:HE21	1.81	0.45
37:CJ:155:ARG:N	37:CJ:155:ARG:CD	2.78	0.45
49:CV:85:LYS:HG2	49:CV:86:GLU:N	2.30	0.45
13:D0:91:GLN:NE2	13:D0:91:GLN:N	2.62	0.45
26:D4:62:ARG:HG3	26:D4:62:ARG:O	2.16	0.45
55:DA:1066:U:C3'	55:DA:1066:U:O2	2.63	0.45
55:DA:550:G:N3	55:DA:1220:A:C2	2.84	0.45
55:DA:1266:G:O2'	55:DA:1267:U:P	2.74	0.45
55:DA:1416:G:H2'	55:DA:1417:C:C6	2.51	0.45
55:DA:1445:C:H2'	55:DA:1446:C:C6	2.50	0.45
55:DA:1449:A:H5'	55:DA:1449(A):G:OP2	2.16	0.45
55:DA:1773:A:N7	55:DA:1829:A:C1'	2.80	0.45
55:DA:1956:U:C4	55:DA:1957:C:C5	3.05	0.45
55:DA:2001:A:H2'	55:DA:2002:G:C8	2.52	0.45
55:DA:2162:G:H2'	55:DA:2163:C:C6	2.52	0.45
55:DA:21:A:H2'	55:DA:22:C:O4'	2.15	0.45
55:DA:2533:A:H2'	55:DA:2534:A:H5'	1.98	0.45
55:DA:26:G:H1'	55:DA:514:A:H61	1.77	0.45
55:DA:2887:U:O2'	55:DA:2888:C:H5'	2.16	0.45
55:DA:626:U:H5'	55:DA:627:A:C5'	2.47	0.45
55:DA:752:A:H4'	55:DA:753:C:O5'	2.17	0.45
55:DA:846:C:HO2'	55:DA:847:U:P	2.39	0.45
55:DA:903:C:H2'	55:DA:904:C:C6	2.51	0.45
55:DA:979:G:H3'	55:DA:980:A:H5''	1.98	0.45
3:DD:98:VAL:C	3:DD:100:GLY:H	2.19	0.45
3:DD:115:GLN:HG2	3:DD:116:GLN:O	2.16	0.45
3:DD:72:LYS:HG2	3:DD:103:ARG:NH2	2.30	0.45
4:DE:115:GLY:O	4:DE:119:ARG:HB2	2.16	0.45
4:DE:92:THR:HG22	4:DE:93:VAL:N	2.31	0.45
5:DF:132:VAL:HG23	5:DF:133:ASN:N	2.31	0.45
7:DH:19:VAL:CG1	7:DH:20:ALA:N	2.79	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:DI:24:ILE:C	56:DI:26:ALA:H	2.20	0.45
58:DL:112:MET:HE2	58:DL:123:ALA:H	1.81	0.45
9:DM:30:ILE:O	9:DM:34:LEU:CD2	2.64	0.45
55:DA:1140:C:H5'	9:DM:66:LYS:NZ	2.32	0.45
9:DM:96:GLU:O	9:DM:98:VAL:HG12	2.16	0.45
12:DP:31:ASP:OD1	12:DP:134:ARG:HD2	2.17	0.45
20:DU:9:LYS:HA	20:DU:27:VAL:CG2	2.46	0.45
21:DV:53:ILE:HG22	21:DV:71:VAL:O	2.16	0.45
57:DY:17:LEU:O	57:DY:22:GLY:HA3	2.16	0.45
55:DA:396:G:O3'	23:DZ:44:PRO:HA	2.16	0.45
17:A2:69:LYS:HA	17:A2:87:HIS:O	2.15	0.45
27:A5:30:LEU:HA	27:A5:42:PRO:HD3	1.98	0.45
30:A8:14:VAL:CG1	30:A8:15:LYS:N	2.77	0.45
1:AA:1039:G:O2'	1:AA:1040:C:H5'	2.16	0.45
1:AA:1117:G:O2'	1:AA:1118:C:H5'	2.16	0.45
1:AA:1322:A:O2'	1:AA:1323:U:H5'	2.17	0.45
1:AA:1449:A:O2'	1:AA:1530:G:N2	2.35	0.45
1:AA:1825:A:O4'	3:AD:254:THR:HG21	2.16	0.45
1:AA:2000:G:HO2'	1:AA:2689:U:H5	1.63	0.45
1:AA:2092:U:C6	1:AA:2092:U:C5'	2.99	0.45
1:AA:2335:A:HO2'	1:AA:2336:A:P	2.36	0.45
1:AA:2590:A:O2'	1:AA:2591:C:H5'	2.17	0.45
1:AA:278:A:N3	1:AA:279:C:C6	2.85	0.45
1:AA:2815:C:C2	1:AA:2816:C:C6	3.04	0.45
1:AA:2851:A:C2'	1:AA:2852:G:O5'	2.65	0.45
1:AA:2851:A:H2'	1:AA:2852:G:O5'	2.16	0.45
1:AA:310:A:O2'	1:AA:311:A:O5'	2.28	0.45
1:AA:621:A:H2'	1:AA:622:G:O5'	2.16	0.45
1:AA:747:U:O4'	27:A5:2:ALA:HB3	2.17	0.45
1:AA:859:G:O2'	1:AA:860:U:C6	2.69	0.45
1:AA:896:A:H2	21:AV:178:GLU:CG	2.27	0.45
3:AD:165:ILE:HD13	3:AD:175:LEU:CD2	2.46	0.45
3:AD:53:PHE:HA	3:AD:218:ARG:HB2	1.98	0.45
3:AD:43:ARG:CB	3:AD:54:ARG:HB2	2.47	0.45
5:AF:21:ALA:HB3	5:AF:23:ASP:CG	2.37	0.45
8:AK:125:GLU:CA	8:AK:141:LYS:HB3	2.40	0.45
8:AK:56:LYS:HG3	8:AK:57:ARG:H	1.81	0.45
11:AO:97:PRO:O	11:AO:98:GLU:CB	2.62	0.45
14:AQ:62:LYS:HB3	14:AQ:97:ARG:CD	2.47	0.45
15:AR:96:ARG:HH11	15:AR:96:ARG:CB	2.30	0.45
18:AS:75:TYR:CZ	18:AS:104:THR:HG21	2.51	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:AU:80:GLY:O	20:AU:81:LYS:HG3	2.16	0.45
21:AV:59:LEU:O	21:AV:60:GLU:HG2	2.17	0.45
31:BA:1106:G:H2'	31:BA:1107:C:C6	2.49	0.45
31:BA:330:C:H5''	31:BA:330:C:H6	1.80	0.45
31:BA:491:G:O2'	31:BA:492:G:H5'	2.17	0.45
31:BA:645:C:H2'	31:BA:646:U:C6	2.52	0.45
31:BA:696:A:H1'	31:BA:786:G:O2'	2.17	0.45
52:BC:15:G:H2'	52:BC:16:U:C6	2.51	0.45
52:BD:16:U:H2'	52:BD:16:U:O2	2.16	0.45
52:BD:46:G:N2	52:BD:47:U:O2'	2.50	0.45
32:BE:124:SER:C	32:BE:126:GLU:H	2.19	0.45
32:BE:212:GLN:O	32:BE:216:SER:N	2.45	0.45
32:BE:60:ASP:HA	32:BE:63:MET:HE2	1.97	0.45
33:BF:125:GLU:HG2	33:BF:190:ARG:O	2.17	0.45
33:BF:34:LEU:CG	33:BF:38:ARG:HH21	2.30	0.45
33:BF:39:ILE:C	33:BF:41:GLY:N	2.69	0.45
34:BG:61:LYS:HZ2	34:BG:62:GLN:NE2	2.14	0.45
35:BH:91:LEU:HA	35:BH:120:THR:HG22	1.97	0.45
34:BG:88:VAL:HG13	35:BH:97:GLY:CA	2.47	0.45
36:BI:8:ILE:HD12	36:BI:8:ILE:N	2.31	0.45
39:BL:3:GLN:HA	39:BL:19:LEU:O	2.17	0.45
43:BP:108:ARG:HH11	43:BP:108:ARG:HG3	1.82	0.45
43:BP:15:VAL:O	43:BP:17:VAL:N	2.50	0.45
43:BP:81:LEU:O	43:BP:82:MET:C	2.54	0.45
45:BR:3:ILE:HG22	45:BR:38:ARG:NE	2.32	0.45
46:BS:7:ALA:CB	46:BS:20:VAL:HG11	2.39	0.45
47:BT:10:VAL:HA	47:BT:20:THR:O	2.16	0.45
48:BU:50:ILE:HD12	48:BU:70:ILE:HD12	1.98	0.45
54:CA:1337:G:H5''	54:CA:1338:G:OP1	2.15	0.45
54:CA:1346:A:H5'	39:CL:120:ARG:NH1	2.31	0.45
54:CA:16:A:O2'	54:CA:17:U:H5'	2.16	0.45
54:CA:184:G:H2'	54:CA:185:A:H8	1.82	0.45
54:CA:265:G:H4'	47:CT:66:SER:CA	2.46	0.45
54:CA:382:A:H2'	54:CA:383:A:H8	1.77	0.45
54:CA:859:A:H2'	54:CA:860:A:O4'	2.17	0.45
54:CA:91:C:O2'	54:CA:92:G:H5''	2.17	0.45
54:CA:956:U:O2	54:CA:960:U:C5	2.69	0.45
52:CD:58:A:H2'	52:CD:60:U:OP2	2.17	0.45
52:CD:65:G:N3	52:CD:65:G:H2'	2.31	0.45
54:CA:1060:C:C5	33:CF:2:GLY:HA3	2.50	0.45
34:CG:147:ALA:HB2	34:CG:182:LYS:CB	2.46	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:CJ:63:LYS:HG3	37:CJ:64:GLN:N	2.31	0.45
38:CK:80:ILE:HG23	38:CK:137:VAL:HG13	1.98	0.45
38:CK:63:LEU:N	38:CK:63:LEU:HD22	2.30	0.45
33:CF:29:TYR:CD1	44:CQ:36:PHE:CE1	3.05	0.45
46:CS:81:ARG:CB	46:CS:81:ARG:HH11	2.29	0.45
54:CA:255:G:H1'	47:CT:16:GLN:NE2	2.31	0.45
47:CT:62:SER:CB	47:CT:72:ARG:NE	2.79	0.45
13:D0:53:HIS:HB2	13:D0:94:TYR:HE1	1.81	0.45
17:D2:39:LEU:CD1	17:D2:39:LEU:N	2.79	0.45
28:D6:44:ARG:O	28:D6:45:LYS:HG2	2.17	0.45
30:D8:48:PHE:N	30:D8:48:PHE:HD1	2.12	0.45
55:DA:1374:G:C5	55:DA:1375:C:C5	3.04	0.45
55:DA:1385:G:HO2'	55:DA:1386:C:H6	1.61	0.45
55:DA:1638:C:H4'	55:DA:2710:C:O2	2.17	0.45
55:DA:1819:A:H5''	3:DD:158:ALA:HB3	1.97	0.45
55:DA:226:G:O2'	55:DA:227:A:P	2.74	0.45
55:DA:2399:G:H2'	55:DA:2400:G:O4'	2.17	0.45
55:DA:2638:G:P	4:DE:82:ARG:NH2	2.89	0.45
55:DA:2820:A:O4'	13:D0:4:LEU:HD23	2.17	0.45
55:DA:767:U:O2'	55:DA:768:G:H5'	2.17	0.45
2:DB:15:A:H1'	2:DB:109:G:C4	2.51	0.45
3:DD:12:SER:HB2	3:DD:208:LYS:HB3	1.97	0.45
56:DI:16:THR:CG2	56:DI:17:VAL:HG22	2.44	0.45
9:DM:112:LEU:O	9:DM:112:LEU:HD23	2.17	0.45
11:DO:11:GLY:O	11:DO:12:ALA:HB3	2.13	0.45
55:DA:2415:G:H4'	11:DO:66:GLY:C	2.36	0.45
12:DP:118:LEU:O	12:DP:119:ARG:C	2.55	0.45
12:DP:25:ASP:HA	12:DP:100:GLY:O	2.16	0.45
14:DQ:106:ARG:O	14:DQ:107:GLU:CB	2.64	0.45
15:DR:106:SER:HA	15:DR:110:ILE:HB	1.98	0.45
15:DR:133:GLU:HA	15:DR:136:GLN:HG2	1.98	0.45
57:DY:38:HIS:HB3	57:DY:40:LEU:H	1.81	0.45
23:DZ:80:LEU:CD2	23:DZ:80:LEU:O	2.64	0.45
16:A1:83:LEU:HD12	16:A1:88:ILE:HD11	1.97	0.45
16:A1:92:ARG:NH1	17:A2:11:GLN:CD	2.69	0.45
1:AA:2271:G:H5''	22:A3:20:ARG:NE	2.31	0.45
1:AA:1048:A:N6	7:AH:2:SER:OG	2.49	0.45
1:AA:1343:G:H2'	1:AA:1343:G:N3	2.31	0.45
1:AA:1575:C:H2'	1:AA:1575:C:O2	2.16	0.45
1:AA:1822:G:H8	1:AA:1822:G:H5'	1.81	0.45
1:AA:2059:A:H5'	1:AA:2060:A:OP2	2.17	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2154:G:O2'	1:AA:2155:G:H5'	2.17	0.45
1:AA:2393:A:H2'	1:AA:2394:C:O4'	2.17	0.45
1:AA:2451:A:H5'	52:BC:76:A:C2	2.51	0.45
1:AA:2791:C:N3	1:AA:2792:G:C5	2.85	0.45
1:AA:315:G:H2'	1:AA:316:C:O4'	2.16	0.45
1:AA:603:A:HO2'	1:AA:604:G:P	2.40	0.45
4:AE:36:ARG:HG2	4:AE:36:ARG:HH11	1.81	0.45
4:AE:95:ILE:CD1	4:AE:95:ILE:N	2.79	0.45
5:AF:124:LEU:HG	5:AF:126:VAL:CG1	2.47	0.45
6:AG:72:ARG:NH1	6:AG:72:ARG:HG3	2.31	0.45
7:AH:152:ARG:O	7:AH:153:LYS:CB	2.65	0.45
7:AH:29:PRO:HG2	7:AH:30:LYS:HD2	1.98	0.45
11:AO:124:LYS:HG3	11:AO:143:GLY:C	2.37	0.45
11:AO:61:ARG:O	11:AO:62:LEU:CB	2.55	0.45
15:AR:137:LYS:NZ	15:AR:137:LYS:HB3	2.32	0.45
15:AR:16:ARG:NE	15:AR:19:LEU:HD11	2.31	0.45
15:AR:94:ALA:C	15:AR:96:ARG:N	2.68	0.45
15:AR:98:LYS:N	15:AR:98:LYS:CD	2.79	0.45
1:AA:483:A:C5'	20:AU:49:VAL:HG13	2.43	0.45
21:AV:131:ARG:CG	21:AV:131:ARG:NH1	2.58	0.45
21:AV:58:VAL:O	21:AV:59:LEU:HB2	2.15	0.45
31:BA:1058:G:H2'	31:BA:1059:C:H6	1.81	0.45
31:BA:1320:C:N3	49:BV:36:ARG:HG3	2.32	0.45
31:BA:1315:U:O2	31:BA:1360:A:H2	2.00	0.45
31:BA:1498:U:O2'	31:BA:1499:A:P	2.74	0.45
31:BA:414:A:OP2	31:BA:428:G:N2	2.46	0.45
31:BA:783:C:N4	31:BA:800:G:N2	2.65	0.45
31:BA:577:G:C8	31:BA:816:A:C6	3.04	0.45
52:BD:44:G:H2'	52:BD:45:U:C6	2.52	0.45
1:AA:1851:U:H4'	52:BD:70:G:N2	2.32	0.45
32:BE:100:GLY:O	32:BE:101:MET:C	2.55	0.45
32:BE:119:GLU:C	32:BE:121:LEU:H	2.20	0.45
33:BF:81:GLY:O	33:BF:82:GLU:CB	2.64	0.45
35:BH:103:GLY:O	35:BH:104:ALA:C	2.53	0.45
39:BL:53:VAL:C	39:BL:55:ALA:N	2.69	0.45
42:BO:89:ARG:HH12	42:BO:91:LYS:HA	1.82	0.45
49:BV:86:GLU:HA	49:BV:86:GLU:OE1	2.16	0.45
50:BW:23:ARG:HA	50:BW:26:ASN:HD21	1.81	0.45
50:BW:25:ARG:HG2	50:BW:29:LYS:NZ	2.32	0.45
50:BW:56:MET:HG3	50:BW:84:LEU:HD11	1.95	0.45
54:CA:1032:A:O5'	54:CA:1032(A):G:H5''	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1091:U:O2	54:CA:1093:A:C8	2.70	0.45
54:CA:1295:G:O2'	43:CP:14:ARG:NH1	2.45	0.45
54:CA:1299:A:C6	54:CA:1301:U:C2	3.04	0.45
54:CA:1301:U:C2'	54:CA:1301:U:O2	2.64	0.45
54:CA:412:A:O2'	54:CA:413:G:P	2.75	0.45
54:CA:624:C:H2'	54:CA:625:G:C8	2.51	0.45
54:CA:975:A:H4'	54:CA:1358:U:H1'	1.98	0.45
52:CD:73:A:H8	52:CD:73:A:H5'	1.80	0.45
32:CE:138:LEU:C	32:CE:140:HIS:H	2.18	0.45
33:CF:87:LEU:C	33:CF:89:GLU:N	2.67	0.45
34:CG:92:VAL:O	34:CG:96:LEU:HD22	2.16	0.45
54:CA:9:G:H5'	35:CH:122:GLU:OE2	2.17	0.45
35:CH:12:LEU:CD2	35:CH:13:ILE:N	2.77	0.45
38:CK:41:ARG:HH11	38:CK:41:ARG:HB3	1.79	0.45
38:CK:91:ARG:NH1	38:CK:91:ARG:CG	2.77	0.45
43:CP:84:ILE:HD13	49:CV:65:ASN:CG	2.36	0.45
44:CQ:8:GLU:C	44:CQ:10:ALA:H	2.20	0.45
45:CR:54:ARG:NH1	45:CR:54:ARG:HG2	2.31	0.45
46:CS:76:GLN:O	46:CS:76:GLN:CG	2.64	0.45
50:CW:13:LEU:CD1	50:CW:17:ARG:NH1	2.78	0.45
17:D2:35:LEU:C	17:D2:37:VAL:H	2.20	0.45
28:D6:14:THR:O	28:D6:49:HIS:HA	2.16	0.45
30:D8:61:LEU:C	30:D8:63:PRO:HD2	2.37	0.45
55:DA:1048:A:H3'	55:DA:1049:C:H5'	1.97	0.45
55:DA:1055:G:H2'	55:DA:1056:G:C5'	2.44	0.45
55:DA:1060:U:C4'	55:DA:1061:U:O5'	2.53	0.45
55:DA:1398:C:O2'	55:DA:1399:C:H5'	2.15	0.45
55:DA:1416:G:O2'	55:DA:1417:C:O5'	2.34	0.45
55:DA:1853:A:H2'	55:DA:1854:A:C8	2.52	0.45
55:DA:1902:C:H5'	3:DD:246:PRO:HD3	1.98	0.45
55:DA:2143:C:H2'	55:DA:2144:U:O4'	2.17	0.45
55:DA:2646:C:H2'	55:DA:2647:U:O4'	2.16	0.45
55:DA:2773:C:OP1	4:DE:166:THR:OG1	2.35	0.45
55:DA:301:G:O2'	55:DA:302:C:P	2.74	0.45
55:DA:616:A:H4'	55:DA:617:G:OP1	2.16	0.45
55:DA:843:G:O2'	55:DA:844:C:H5'	2.16	0.45
55:DA:897:C:OP1	55:DA:897:C:H5	1.84	0.45
3:DD:134:ARG:CB	3:DD:135:PHE:CD2	2.95	0.45
4:DE:3:GLY:O	4:DE:4:ILE:HB	2.16	0.45
5:DF:117:ARG:HH21	5:DF:187:VAL:HA	1.82	0.45
6:DG:39:ILE:HB	6:DG:92:VAL:CG1	2.47	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:DI:1:MET:HB3	56:DI:5:ILE:HG22	1.98	0.45
57:DY:138:LEU:HD22	56:DJ:22:GLN:OE1	2.16	0.45
8:DK:3:VAL:CB	8:DK:37:VAL:O	2.65	0.45
55:DA:1006:C:O2'	9:DM:106:MET:HB3	2.17	0.45
12:DP:57:HIS:NE2	12:DP:116:GLU:HG2	2.31	0.45
15:DR:14:TYR:CD1	15:DR:14:TYR:N	2.83	0.45
21:DV:108:PRO:HG3	21:DV:142:SER:O	2.16	0.45
21:DV:72:ARG:O	21:DV:73:GLN:HB2	2.17	0.45
57:DY:73:GLY:CA	57:DY:119:ALA:HA	2.47	0.45
57:DY:131:MET:C	57:DY:133:GLU:N	2.70	0.45
57:DY:141:VAL:O	57:DY:142:LEU:C	2.55	0.45
57:DY:49:ALA:HA	57:DY:83:TYR:HB3	1.98	0.45
17:A2:10:LYS:N	17:A2:10:LYS:HD2	2.31	0.45
17:A2:98:GLU:HA	17:A2:98:GLU:OE1	2.16	0.45
1:AA:2884:U:OP2	27:A5:43:HIS:HE1	1.98	0.45
1:AA:2372:G:H1'	28:A6:46:HIS:HE1	1.79	0.45
29:A7:13:ALA:O	29:A7:17:GLY:HA3	2.16	0.45
1:AA:1029:A:H2'	1:AA:1030:G:O4'	2.16	0.45
1:AA:1102:C:C2'	1:AA:1103:A:H5''	2.46	0.45
1:AA:1278:A:H2'	1:AA:1279:G:C8	2.52	0.45
1:AA:1507:A:O2'	1:AA:1510:A:N1	2.49	0.45
1:AA:1639:U:C2'	1:AA:1640:C:H5''	2.46	0.45
1:AA:1930:G:N2	1:AA:1968:G:H2'	2.32	0.45
1:AA:1131:G:C8	1:AA:2025:C:H4'	2.51	0.45
1:AA:214:G:H21	1:AA:216:A:H1'	1.82	0.45
1:AA:2093:G:H21	1:AA:2198:A:N6	2.15	0.45
1:AA:2215:G:O2'	1:AA:2216:G:H5'	2.17	0.45
1:AA:2522:U:C2'	1:AA:2523:G:H5''	2.47	0.45
1:AA:2577:A:H5''	1:AA:2578:G:H5'	1.99	0.45
1:AA:2654:A:O2'	1:AA:2655:G:C4'	2.65	0.45
1:AA:2665:A:H2'	1:AA:2666:C:H6	1.81	0.45
1:AA:2820:A:O5'	13:A0:4:LEU:CD2	2.62	0.45
1:AA:284:U:O2'	1:AA:285:C:H5'	2.16	0.45
1:AA:608:A:C2	1:AA:621:A:N7	2.85	0.45
1:AA:654(N):G:C8	1:AA:654(N):G:OP1	2.70	0.45
1:AA:685:A:OP1	1:AA:686:G:N2	2.50	0.45
1:AA:894:C:H5'	1:AA:895:U:OP2	2.17	0.45
1:AA:93:C:H2'	1:AA:93:C:O2	2.17	0.45
2:AB:44:G:N2	2:AB:48:A:N3	2.65	0.45
2:AB:81:G:N2	2:AB:82:G:C6	2.84	0.45
4:AE:68:ALA:HA	4:AE:71:GLY:HA2	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:AF:24:LEU:HD12	5:AF:25:PRO:HD2	1.99	0.45
6:AG:101:ILE:CD1	26:A4:25:TYR:HB2	2.46	0.45
6:AG:47:LYS:NZ	6:AG:81:LYS:HG2	2.32	0.45
7:AH:13:LYS:HA	7:AH:13:LYS:NZ	2.31	0.45
9:AM:27:ALA:HB1	9:AM:103:VAL:HG22	1.97	0.45
9:AM:129:PRO:O	9:AM:130:HIS:C	2.54	0.45
11:AO:118:GLY:O	11:AO:119:GLU:C	2.55	0.45
14:AQ:49:VAL:HG22	14:AQ:80:LEU:HD12	1.98	0.45
15:AR:102:ILE:C	15:AR:102:ILE:HD12	2.36	0.45
18:AS:1:MET:CE	18:AS:2:GLU:H	2.30	0.45
20:AU:17:SER:HB3	20:AU:71:LYS:HB3	1.98	0.45
20:AU:40:GLU:N	20:AU:40:GLU:OE2	2.49	0.45
20:AU:89:PHE:O	20:AU:90:LEU:O	2.34	0.45
21:AV:133:ILE:N	21:AV:133:ILE:CD1	2.79	0.45
21:AV:149:SER:HB2	21:AV:151:HIS:CE1	2.52	0.45
24:AW:26:ARG:NH1	24:AW:26:ARG:CB	2.80	0.45
25:AX:19:GLN:NE2	25:AX:52:HIS:CE1	2.85	0.45
1:AA:380:U:O3'	23:AZ:16:ASN:HB2	2.17	0.45
31:BA:1463:C:O2'	31:BA:1464:G:H5'	2.17	0.45
31:BA:148:G:H2'	31:BA:149:A:H8	1.82	0.45
31:BA:270:A:H2'	31:BA:271:C:O4'	2.17	0.45
31:BA:626:U:C2	31:BA:627:G:C8	3.05	0.45
31:BA:794:A:N3	31:BA:795:C:C2	2.85	0.45
31:BA:848:C:O2'	31:BA:849:C:H5'	2.16	0.45
31:BA:890:G:HO2'	31:BA:891:U:H5	1.61	0.45
32:BE:102:LEU:HD12	32:BE:102:LEU:N	2.32	0.45
33:BF:148:GLY:O	33:BF:203:PHE:N	2.43	0.45
33:BF:94:LEU:HD12	33:BF:95:THR:N	2.31	0.45
34:BG:2:GLY:O	34:BG:3:ARG:HB2	2.16	0.45
36:BI:15:ASP:OD1	36:BI:16:GLN:N	2.49	0.45
37:BJ:47:CYS:O	37:BJ:50:ILE:HB	2.17	0.45
39:BL:78:LYS:HZ2	39:BL:78:LYS:HB2	1.80	0.45
41:BN:120:ARG:HA	41:BN:121:PRO:HD3	1.79	0.45
43:BP:109:THR:O	43:BP:109:THR:HG22	2.16	0.45
26:A4:49:PHE:CZ	43:BP:61:GLU:O	2.70	0.45
43:BP:79:LYS:C	43:BP:79:LYS:HD3	2.36	0.45
1:AA:887:A:H2	43:BP:79:LYS:HE3	1.81	0.45
54:CA:1026:G:C2	54:CA:1027:C:H1'	2.51	0.45
54:CA:101:A:C2	54:CA:102:G:C8	3.05	0.45
54:CA:1126:U:OP2	54:CA:1281:U:O2	2.35	0.45
54:CA:1148:U:H2'	54:CA:1149:C:H5'	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:191(F):U:O2'	54:CA:191:G:H5'	2.17	0.45
54:CA:748:C:HO2'	54:CA:749:C:P	2.34	0.45
52:CB:9:A:HO2'	52:CB:10:G:P	2.38	0.45
33:CF:14:ILE:CG1	33:CF:15:THR:N	2.79	0.45
33:CF:174:PRO:C	33:CF:176:HIS:H	2.20	0.45
34:CG:104:VAL:O	34:CG:105:VAL:C	2.55	0.45
34:CG:52:SER:HB3	34:CG:55:ALA:CB	2.47	0.45
34:CG:74:GLN:HA	34:CG:77:ASN:HD22	1.80	0.45
35:CH:140:ARG:CB	35:CH:140:ARG:NH1	2.79	0.45
35:CH:67:VAL:HG22	35:CH:68:GLU:N	2.31	0.45
37:CJ:15:ASP:OD2	37:CJ:44:TYR:OH	2.35	0.45
37:CJ:31:MET:SD	37:CJ:36:LYS:HB2	2.57	0.45
35:CH:148:VAL:CG2	38:CK:107:LEU:HD22	2.37	0.45
54:CA:1367:C:OP1	39:CL:114:TYR:HA	2.16	0.45
39:CL:40:LEU:C	39:CL:42:ARG:H	2.20	0.45
54:CA:539:A:OP2	42:CO:115:LYS:HE3	2.16	0.45
42:CO:6:THR:HG23	42:CO:9:GLN:HE21	1.81	0.45
54:CA:1331:G:OP2	43:CP:23:TYR:CD2	2.69	0.45
44:CQ:8:GLU:C	44:CQ:10:ALA:N	2.68	0.45
44:CQ:14:PRO:O	44:CQ:15:LYS:HB2	2.16	0.45
51:CX:12:LYS:HE2	51:CX:19:GLY:N	2.31	0.45
17:D2:89:GLN:HE21	17:D2:89:GLN:CA	2.07	0.45
17:D2:25:LEU:H	17:D2:92:THR:CG2	2.30	0.45
26:D4:21:VAL:O	26:D4:24:THR:HG23	2.17	0.45
55:DA:1167:U:H2'	55:DA:1168:G:H8	1.81	0.45
55:DA:1341:U:O4	19:DT:16:LYS:HE2	2.16	0.45
55:DA:1525:G:H2'	55:DA:1526:G:H8	1.82	0.45
55:DA:1534:G:H5''	55:DA:1534:G:C8	2.51	0.45
54:CA:1418:A:H2	55:DA:1948:G:N3	2.14	0.45
55:DA:1970:A:H1'	55:DA:1972:A:C8	2.51	0.45
55:DA:2197:U:O2'	55:DA:2198:A:P	2.75	0.45
55:DA:228:A:O2'	55:DA:229:A:OP1	2.35	0.45
55:DA:2615:U:H2'	55:DA:2616:C:C6	2.51	0.45
55:DA:321:G:HO2'	55:DA:340:A:C2'	2.30	0.45
55:DA:2:G:H2'	55:DA:3:U:H6	1.82	0.45
55:DA:528:A:H3'	55:DA:529:A:C5'	2.47	0.45
55:DA:535:C:O3'	16:D1:53:ARG:NH1	2.49	0.45
55:DA:608:A:C2	55:DA:621:A:N7	2.85	0.45
55:DA:808:G:O2'	55:DA:809:G:H5'	2.16	0.45
55:DA:885:C:N3	55:DA:892:G:C2	2.85	0.45
55:DA:940:G:H3'	55:DA:941:A:H5''	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:971:C:OP1	55:DA:974:G:C8	2.70	0.45
3:DD:118:VAL:HG22	3:DD:119:ALA:N	2.32	0.45
4:DE:116:VAL:O	4:DE:117:MET:HB3	2.16	0.45
5:DF:197:ASP:C	5:DF:199:TRP:H	2.20	0.45
5:DF:65:TRP:CZ3	5:DF:73:ALA:O	2.69	0.45
7:DH:97:ARG:O	7:DH:125:VAL:HG11	2.17	0.45
58:DL:60:TYR:OH	58:DL:66:THR:CG2	2.64	0.45
58:DL:76:TYR:O	58:DL:79:ARG:N	2.50	0.45
9:DM:71:ILE:HG22	9:DM:86:PRO:HA	1.99	0.45
10:DN:104:ARG:NH1	15:DR:36:GLU:CD	2.70	0.45
11:DO:41:ARG:NH2	11:DO:41:ARG:CG	2.79	0.45
14:DQ:24:LEU:N	14:DQ:24:LEU:HD22	2.32	0.45
14:DQ:25:ARG:NH1	14:DQ:25:ARG:HB3	2.32	0.45
14:DQ:38:GLN:HG3	14:DQ:47:THR:HG21	1.98	0.45
19:DT:3:THR:O	19:DT:6:ASP:HB2	2.16	0.45
20:DU:4:LYS:O	20:DU:5:MET:O	2.35	0.45
21:DV:122:ARG:NH1	21:DV:122:ARG:HG2	2.32	0.45
57:DY:74:LEU:HB2	57:DY:120:LYS:HE2	1.99	0.45
57:DY:59:ILE:O	57:DY:61:LEU:N	2.49	0.45
13:A0:41:ALA:HB1	13:A0:97:VAL:HG11	1.99	0.45
17:A2:12:TYR:CG	17:A2:20:LEU:HD21	2.52	0.45
17:A2:35:LEU:HD23	17:A2:37:VAL:HG21	1.98	0.45
17:A2:41:GLY:N	17:A2:46:VAL:CG1	2.79	0.45
17:A2:80:GLN:N	17:A2:80:GLN:NE2	2.64	0.45
26:A4:67:TYR:HD2	26:A4:67:TYR:C	2.20	0.45
28:A6:26:ASN:C	28:A6:27:LYS:HD3	2.36	0.45
28:A6:52:VAL:CG1	28:A6:53:LYS:N	2.80	0.45
1:AA:1060:U:H1'	1:AA:1062:G:C4'	2.47	0.45
1:AA:1195:G:N3	1:AA:1227:A:H2	2.14	0.45
1:AA:1812:A:H2'	1:AA:1813:G:C8	2.52	0.45
1:AA:222:A:O2'	1:AA:223:A:P	2.75	0.45
1:AA:2365:G:H4'	22:A3:60:PHE:CZ	2.52	0.45
1:AA:2406:U:C4	11:AO:72:PRO:HB2	2.52	0.45
1:AA:2791:C:N4	1:AA:2792:G:O6	2.49	0.45
1:AA:301:G:O2'	1:AA:302:C:O4'	2.34	0.45
1:AA:303:U:C2	1:AA:304:G:C8	3.04	0.45
2:AB:17:C:O2'	2:AB:18:G:H5'	2.17	0.45
1:AA:1820:U:C4	3:AD:160:GLY:HA3	2.51	0.45
3:AD:30:GLU:CG	3:AD:63:ARG:NH2	2.79	0.45
3:AD:70:TRP:O	3:AD:73:VAL:HG22	2.16	0.45
4:AE:124:GLY:HA2	4:AE:137:HIS:O	2.15	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:AF:103:LYS:HA	5:AF:106:ARG:HG3	1.98	0.45
5:AF:63:LYS:CE	5:AF:67:GLN:CB	2.95	0.45
5:AF:7:TYR:O	5:AF:17:ARG:N	2.49	0.45
1:AA:1111:A:H5'	7:AH:3:ARG:NH1	2.32	0.45
8:AK:35:LEU:O	8:AK:36:ALA:HB2	2.17	0.45
12:AP:46:GLN:O	12:AP:47:ILE:C	2.54	0.45
14:AQ:110:LEU:HA	14:AQ:112:PHE:CE1	2.51	0.45
15:AR:1:MET:O	15:AR:3:ARG:N	2.46	0.45
15:AR:90:GLN:CA	15:AR:90:GLN:HE21	2.23	0.45
21:AV:35:ARG:NH1	21:AV:35:ARG:HB3	2.32	0.45
21:AV:96:VAL:O	21:AV:96:VAL:HG12	2.15	0.45
1:AA:61:G:OP1	24:AW:50:ILE:HD13	2.17	0.45
31:BA:1025:U:O2'	31:BA:1026:G:O4'	2.31	0.45
31:BA:1126:U:H5''	31:BA:1280:A:N7	2.32	0.45
31:BA:1372:U:OP1	39:BL:71:SER:HB3	2.16	0.45
31:BA:1411:C:H2'	31:BA:1412:C:C6	2.51	0.45
31:BA:149:A:O2'	31:BA:150:C:H5'	2.17	0.45
31:BA:197:A:H4'	31:BA:198:G:O5'	2.16	0.45
31:BA:197:A:N6	31:BA:221:C:C5'	2.79	0.45
31:BA:210:U:O2'	31:BA:216:G:P	2.75	0.45
31:BA:278:G:O4'	31:BA:282:A:H1'	2.16	0.45
31:BA:533:A:O2'	31:BA:535:A:OP2	2.34	0.45
31:BA:560:U:H4'	31:BA:561:U:C5'	2.47	0.45
31:BA:989:C:H1'	31:BA:1016:A:H2	1.81	0.45
32:BE:10:LEU:O	32:BE:13:ALA:CB	2.65	0.45
32:BE:67:THR:HG22	32:BE:90:MET:CE	2.46	0.45
32:BE:67:THR:HG22	32:BE:90:MET:HE1	1.98	0.45
33:BF:115:LEU:O	33:BF:116:VAL:C	2.55	0.45
34:BG:159:ARG:O	34:BG:160:GLN:C	2.55	0.45
36:BI:53:ALA:O	36:BI:54:LYS:CB	2.64	0.45
39:BL:48:GLU:N	39:BL:49:PRO:HD2	2.32	0.45
43:BP:15:VAL:HG12	43:BP:45:VAL:HG22	1.99	0.45
48:BU:22:VAL:HG22	48:BU:23:LYS:N	2.30	0.45
49:BV:23:ASN:O	49:BV:26:GLY:N	2.48	0.45
49:BV:31:ILE:HG23	49:BV:49:ILE:HA	1.99	0.45
54:CA:1003:G:H21	54:CA:1005:A:P	2.39	0.45
54:CA:1176:A:N6	54:CA:1177:G:C2	2.85	0.45
54:CA:1351:U:O2'	54:CA:1352:C:H5'	2.17	0.45
54:CA:328:C:HO2'	54:CA:329:A:P	2.35	0.45
52:CB:4:C:H2'	52:CB:5:G:H8	1.82	0.45
52:CD:59:U:H2'	52:CD:60:U:O4'	2.17	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:CE:70:PHE:O	32:CE:93:VAL:N	2.49	0.45
35:CH:108:ALA:O	35:CH:112:LEU:HG	2.17	0.45
35:CH:140:ARG:HB2	35:CH:140:ARG:NH1	2.32	0.45
35:CH:72:GLN:C	35:CH:74:GLY:N	2.69	0.45
36:CI:72:VAL:CG1	36:CI:73:ASN:N	2.78	0.45
39:CL:111:ARG:O	39:CL:113:LYS:HD2	2.17	0.45
39:CL:11:LYS:O	39:CL:11:LYS:HD2	2.17	0.45
41:CN:34:ASP:HB2	41:CN:35:PRO:CD	2.47	0.45
46:CS:19:ILE:CG2	46:CS:36:ILE:HG13	2.43	0.45
50:CW:87:LYS:O	50:CW:90:GLN:N	2.49	0.45
13:D0:96:ARG:HH11	13:D0:96:ARG:HG3	1.82	0.45
26:D4:12:ALA:HB1	26:D4:30:GLU:N	2.22	0.45
26:D4:55:ARG:O	26:D4:57:GLU:N	2.49	0.45
27:D5:2:ALA:O	27:D5:3:LYS:HB2	2.16	0.45
55:DA:1180:C:C2'	55:DA:1181:C:C5'	2.94	0.45
55:DA:1340:U:HO2'	55:DA:1602:U:H2'	1.80	0.45
55:DA:1893:C:O2'	55:DA:1894:C:H5'	2.17	0.45
55:DA:1914:C:H2'	55:DA:1915:U:O4'	2.17	0.45
55:DA:2165:G:C2'	55:DA:2166:G:H5'	2.46	0.45
55:DA:2115:G:C1'	55:DA:2171:A:H61	2.30	0.45
55:DA:2353:G:H5"	22:D3:32:ARG:NH1	2.32	0.45
55:DA:2428:G:H5"	55:DA:2429:G:OP1	2.16	0.45
55:DA:2657:A:H2'	55:DA:2658:C:H5'	1.98	0.45
55:DA:372:G:HO2'	55:DA:373:U:H5	1.61	0.45
55:DA:442:G:O4'	5:DF:46:ARG:HD3	2.17	0.45
55:DA:469:G:C2'	55:DA:470:A:H5"	2.46	0.45
55:DA:621:A:H2'	55:DA:622:G:H5'	1.98	0.45
55:DA:774:A:H2	55:DA:787:U:C2'	2.30	0.45
2:DB:40:U:C2'	2:DB:41:U:OP1	2.65	0.45
55:DA:782:A:N1	3:DD:226:MET:HE1	2.31	0.45
3:DD:34:VAL:CG1	3:DD:34:VAL:O	2.59	0.45
3:DD:85:ASP:HB2	3:DD:92:ILE:HG13	1.99	0.45
4:DE:119:ARG:HB3	4:DE:120:TRP:CD1	2.52	0.45
4:DE:13:ARG:CA	4:DE:21:VAL:HA	2.47	0.45
6:DG:145:THR:HG22	26:D4:28:LYS:NZ	2.30	0.45
6:DG:58:GLN:HE22	6:DG:148:MET:CE	2.30	0.45
58:DL:10:LEU:HB2	58:DL:11:GLN:H	1.54	0.45
58:DL:19:PRO:HD3	58:DL:38:VAL:CG1	2.43	0.45
58:DL:52:ILE:O	58:DL:53:VAL:C	2.54	0.45
9:DM:133:GLN:CB	9:DM:135:PRO:HD3	2.42	0.45
10:DN:104:ARG:HG3	10:DN:122:LEU:O	2.17	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:DO:19:VAL:HG13	11:DO:20:GLY:N	2.32	0.45
14:DQ:30:ARG:HG2	14:DQ:30:ARG:HH11	1.81	0.45
21:DV:111:VAL:HG21	21:DV:146:ILE:N	2.31	0.45
21:DV:194:PRO:O	21:DV:196:VAL:CG1	2.62	0.45
57:DY:10:LEU:O	57:DY:11:ALA:O	2.35	0.45
57:DY:112:LEU:HD13	57:DY:121:ASP:CG	2.34	0.45
57:DY:122:VAL:CG1	57:DY:126:ALA:HB2	2.42	0.45
57:DY:23:SER:HB3	57:DY:68:LEU:CB	2.31	0.45
16:A1:56:ASP:O	16:A1:59:ARG:HB2	2.16	0.45
16:A1:88:ILE:O	16:A1:90:VAL:N	2.49	0.45
17:A2:97:LYS:HG2	17:A2:97:LYS:O	2.17	0.45
26:A4:61:ARG:O	26:A4:62:ARG:NH1	2.49	0.45
30:A8:38:GLY:C	30:A8:41:ILE:HG22	2.36	0.45
1:AA:1005:C:C5	1:AA:1143:A:H1'	2.51	0.45
1:AA:1699:G:O2'	1:AA:1700:A:P	2.75	0.45
1:AA:2031:A:N3	1:AA:2455:G:O2'	2.37	0.45
1:AA:2130:U:C6	1:AA:2130:U:H3'	2.52	0.45
1:AA:2239:G:OP2	3:AD:244:ARG:NH2	2.40	0.45
1:AA:2346:A:N6	28:A6:28:ARG:NH2	2.64	0.45
1:AA:2362:G:H2'	1:AA:2363:C:H5'	1.98	0.45
1:AA:2656:U:N3	1:AA:2665:A:C2	2.84	0.45
1:AA:2836:U:C4	1:AA:2883:A:N6	2.84	0.45
1:AA:2888:C:H2'	1:AA:2889:C:C6	2.52	0.45
1:AA:2898:U:C2	1:AA:2899:G:N7	2.85	0.45
1:AA:908:C:OP1	12:AP:22:LYS:HG3	2.17	0.45
1:AA:918:A:H2'	1:AA:919:G:H5'	1.99	0.45
1:AA:928:G:H3'	1:AA:929:G:C8	2.52	0.45
1:AA:957:A:N6	1:AA:2494:G:N2	2.64	0.45
3:AD:144:ALA:HB3	3:AD:192:THR:CG2	2.46	0.45
4:AE:102:VAL:HA	4:AE:201:THR:HG1	1.82	0.45
4:AE:95:ILE:CD1	4:AE:95:ILE:H	2.30	0.45
5:AF:9:ILE:HG12	5:AF:15:SER:N	2.32	0.45
5:AF:172:TRP:CD2	5:AF:173:VAL:HG23	2.52	0.45
5:AF:113:ALA:HB2	5:AF:183:VAL:HG12	1.98	0.45
7:AH:115:VAL:HG11	7:AH:148:ILE:HD12	1.98	0.45
14:AQ:66:ALA:HA	14:AQ:69:VAL:HG12	1.98	0.45
15:AR:117:ASP:OD1	15:AR:120:ARG:NE	2.43	0.45
19:AT:36:LYS:HE3	19:AT:54:VAL:O	2.17	0.45
20:AU:13:VAL:CG2	20:AU:14:LEU:N	2.79	0.45
21:AV:182:LYS:HD3	21:AV:183:LEU:H	1.82	0.45
21:AV:5:LEU:O	21:AV:6:LYS:C	2.55	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:1007:C:H2'	31:BA:1008:C:C5'	2.33	0.45
31:BA:1028(B):C:H3'	31:BA:1029:G:C5'	2.47	0.45
31:BA:1190:G:C5'	33:BF:176:HIS:NE2	2.79	0.45
31:BA:1238:A:C2	31:BA:1241:G:N3	2.85	0.45
31:BA:1261:A:H2'	31:BA:1262:C:H5'	1.97	0.45
31:BA:201:C:O2'	31:BA:208:U:P	2.75	0.45
31:BA:397:A:C3'	31:BA:397:A:N3	2.71	0.45
31:BA:560:U:H4'	31:BA:561:U:H5''	1.99	0.45
31:BA:701:C:C2'	31:BA:702:A:OP2	2.65	0.45
31:BA:984:C:O2'	31:BA:985:C:H5'	2.17	0.45
52:BB:18:G:HO2'	52:BB:19:G:P	2.40	0.45
52:BC:18:G:C5'	52:BC:19:G:OP2	2.55	0.45
52:BC:51:U:H2'	52:BC:52:G:C8	2.52	0.45
37:BJ:84:ASN:HB2	52:BD:37:MIA:C14	2.47	0.45
32:BE:15:VAL:C	32:BE:16:HIS:ND1	2.70	0.45
32:BE:60:ASP:HA	32:BE:63:MET:CE	2.47	0.45
35:BH:31:LEU:CD2	35:BH:43:LEU:HD11	2.26	0.45
37:BJ:135:VAL:O	37:BJ:138:LYS:N	2.50	0.45
37:BJ:147:ALA:C	37:BJ:149:ARG:H	2.19	0.45
38:BK:114:THR:HG22	38:BK:130:GLY:O	2.17	0.45
39:BL:55:ALA:HA	39:BL:58:HIS:CD2	2.52	0.45
39:BL:46:ALA:CA	39:BL:78:LYS:HZ2	2.29	0.45
43:BP:5:ALA:HB2	43:BP:22:ILE:CD1	2.45	0.45
49:BV:40:ILE:HA	49:BV:44:MET:HE3	1.99	0.45
51:BX:9:ARG:NH2	51:BX:10:ARG:NE	2.62	0.45
53:C1:30:C:N3	53:C1:31:A:C8	2.84	0.45
54:CA:101:A:O2'	54:CA:102:G:H5'	2.16	0.45
54:CA:1132:C:H2'	54:CA:1133:G:H8	1.81	0.45
54:CA:1134:G:H2'	54:CA:1135:U:H5'	1.99	0.45
54:CA:1245:A:OP2	51:CX:9:ARG:NH1	2.50	0.45
54:CA:1333:A:H2'	54:CA:1334:G:O4'	2.17	0.45
54:CA:1343:G:C1'	39:CL:121:ARG:HH12	2.29	0.45
54:CA:1350:A:C5	54:CA:1351:U:C4	3.04	0.45
54:CA:177:C:H2'	54:CA:178:C:H6	1.82	0.45
54:CA:198:G:N2	54:CA:220:G:H1'	2.31	0.45
54:CA:451:A:N7	54:CA:481:G:C2	2.85	0.45
54:CA:61:G:H2'	54:CA:62:U:O4'	2.17	0.45
54:CA:91:C:C2'	54:CA:92:G:C5'	2.94	0.45
52:CD:53:G:O2'	52:CD:54:U:H5'	2.17	0.45
32:CE:167:PRO:HG3	32:CE:188:ALA:CB	2.47	0.45
33:CF:34:LEU:HD23	33:CF:35:GLU:N	2.32	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:CG:157:LEU:O	34:CG:161:ASN:ND2	2.50	0.45
34:CG:173:TRP:HZ3	34:CG:193:ASP:HB3	1.78	0.45
41:CN:12:ARG:CG	41:CN:13:GLN:N	2.80	0.45
43:CP:14:ARG:H	43:CP:44:ARG:HD2	1.81	0.45
54:CA:974:A:H1'	44:CQ:31:ARG:HE	1.82	0.45
45:CR:51:HIS:O	45:CR:54:ARG:HB3	2.17	0.45
46:CS:21:VAL:O	46:CS:33:ILE:HG12	2.17	0.45
17:D2:75:PHE:C	17:D2:75:PHE:CD1	2.90	0.45
26:D4:30:GLU:O	26:D4:31:ILE:O	2.35	0.45
55:DA:2419:U:H5'	28:D6:23:THR:HG21	1.98	0.45
55:DA:1032:A:H2	55:DA:1122:G:H22	1.65	0.45
55:DA:1048:A:N7	55:DA:1049:C:C5	2.85	0.45
55:DA:1086:A:H4'	55:DA:1103:A:N6	2.32	0.45
55:DA:111:A:C2'	55:DA:112:U:H5'	2.47	0.45
55:DA:1312:U:O2'	55:DA:1313:U:P	2.75	0.45
55:DA:1825:A:OP1	3:DD:249:PRO:HD3	2.17	0.45
55:DA:1980:G:H4'	55:DA:1981:A:OP2	2.17	0.45
55:DA:2804:C:O2'	55:DA:2805:G:H5'	2.17	0.45
55:DA:414:C:H2'	55:DA:415:A:C8	2.52	0.45
55:DA:576:U:H2'	55:DA:577:G:C8	2.52	0.45
55:DA:586:A:H5'	5:DF:89:VAL:HG21	1.98	0.45
55:DA:858:U:OP2	22:D3:77:ARG:NH2	2.33	0.45
4:DE:131:ALA:CB	4:DE:135:HIS:HE1	2.24	0.45
8:DK:23:PRO:O	8:DK:27:ARG:HG2	2.15	0.45
58:DL:100:THR:O	58:DL:104:VAL:HG23	2.17	0.45
58:DL:122:ALA:C	58:DL:124:ALA:N	2.69	0.45
55:DA:1059:G:N2	58:DL:126:MET:HB3	2.31	0.45
58:DL:145:LYS:HA	58:DL:145:LYS:HD2	1.74	0.45
58:DL:8:VAL:H	58:DL:57:ILE:CD1	2.30	0.45
9:DM:15:LEU:HD13	9:DM:15:LEU:C	2.37	0.45
9:DM:63:THR:CG2	9:DM:66:LYS:NZ	2.80	0.45
9:DM:74:ARG:O	9:DM:83:LYS:N	2.40	0.45
11:DO:75:ILE:CG1	11:DO:77:ARG:HH12	2.29	0.45
55:DA:956:G:OP2	12:DP:14:ARG:NH2	2.50	0.45
15:DR:27:THR:HA	15:DR:48:ILE:HA	1.98	0.45
15:DR:33:LYS:HE2	15:DR:84:GLN:CB	2.47	0.45
18:DS:59:VAL:HG12	18:DS:60:ASN:ND2	2.30	0.45
21:DV:197:ILE:O	21:DV:198:LYS:O	2.35	0.45
21:DV:92:SER:O	21:DV:94:GLU:HG2	2.16	0.45
21:DV:95:PRO:HA	21:DV:128:VAL:O	2.17	0.45
57:DY:134:LEU:HA	57:DY:137:GLU:CB	2.46	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1083:U:C5'	57:DY:41:ARG:HD3	2.47	0.45
57:DY:51:LEU:HD23	57:DY:51:LEU:HA	1.57	0.45
16:A1:91:ASP:O	16:A1:92:ARG:O	2.35	0.45
26:A4:12:ALA:HB3	26:A4:24:THR:CG2	2.43	0.45
28:A6:47:THR:HG23	28:A6:49:HIS:CE1	2.51	0.45
11:AO:64:LYS:HZ2	30:A8:30:ARG:HA	1.80	0.45
30:A8:41:ILE:H	30:A8:43:GLN:H	1.64	0.45
1:AA:1080:A:O2'	1:AA:1081:U:H5'	2.17	0.45
1:AA:1142(A):A:N7	1:AA:1144:G:C6	2.84	0.45
1:AA:1170:G:H2'	1:AA:1170:G:N3	2.31	0.45
1:AA:1613:G:C2	1:AA:1617:C:C2	3.05	0.45
1:AA:1652:A:O3'	1:AA:1653:G:C8	2.69	0.45
1:AA:1829:A:N7	1:AA:1830:C:C4	2.85	0.45
1:AA:1879:C:H2'	1:AA:1880:C:C5'	2.47	0.45
1:AA:2344:U:H4'	1:AA:2345:G:OP1	2.16	0.45
1:AA:2777:G:OP2	1:AA:2781:A:O2'	2.32	0.45
1:AA:2791:C:C4	1:AA:2792:G:C5	3.04	0.45
1:AA:310:A:HO2'	1:AA:311:A:H3'	1.82	0.45
1:AA:48:G:H2'	1:AA:49:A:H2	1.82	0.45
1:AA:654(J):A:H2	1:AA:654(L):G:N7	2.15	0.45
1:AA:654(M):C:H2'	1:AA:654(N):G:OP1	2.16	0.45
3:AD:112:GLN:O	3:AD:115:GLN:CB	2.65	0.45
6:AG:16:ARG:HE	6:AG:31:VAL:CG1	2.30	0.45
1:AA:2746:U:C4'	7:AH:138:LYS:HG3	2.27	0.45
8:AK:144:VAL:HG12	8:AK:145:VAL:HG22	1.99	0.45
9:AM:65:LYS:O	9:AM:66:LYS:C	2.54	0.45
11:AO:83:VAL:CG2	11:AO:105:LEU:HD22	2.45	0.45
12:AP:134:ARG:NH1	12:AP:134:ARG:HG2	2.31	0.45
15:AR:9:LEU:O	15:AR:12:SER:HB2	2.16	0.45
1:AA:2683:C:P	15:AR:53:ARG:HH22	2.38	0.45
18:AS:62:HIS:O	18:AS:63:ASP:O	2.35	0.45
20:AU:89:PHE:HD1	20:AU:90:LEU:H	1.62	0.45
20:AU:81:LYS:CE	20:AU:97:ARG:NH2	2.79	0.45
23:AZ:50:ARG:HD2	23:AZ:57:GLU:OE1	2.17	0.45
31:BA:1148:U:H2'	31:BA:1149:C:H5'	1.99	0.45
31:BA:1240:U:C5	37:BJ:32:ARG:HD2	2.52	0.45
31:BA:1512:U:N3	31:BA:1513:A:N7	2.64	0.45
31:BA:158:G:C2'	31:BA:159:G:H5'	2.47	0.45
31:BA:210:U:C2'	31:BA:216:G:OP2	2.64	0.45
31:BA:464:G:H1'	31:BA:468:A:H61	1.82	0.45
31:BA:45:U:H2'	31:BA:46:G:C8	2.52	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:511:C:H1'	31:BA:512:U:C6	2.52	0.45
31:BA:828:A:H5''	31:BA:859:A:N1	2.32	0.45
52:BB:66:U:H2'	52:BB:67:C:C6	2.51	0.45
52:BD:19:G:N3	52:BD:19:G:H2'	2.32	0.45
32:BE:22:LYS:CA	32:BE:22:LYS:NZ	2.79	0.45
32:BE:82:ARG:HG2	32:BE:82:ARG:HH11	1.82	0.45
33:BF:180:ALA:O	33:BF:181:ASN:HB3	2.17	0.45
33:BF:86:VAL:HG23	33:BF:87:LEU:N	2.32	0.45
34:BG:199:ASN:O	34:BG:200:GLU:HG3	2.16	0.45
36:BI:14:LEU:HD23	36:BI:15:ASP:N	2.32	0.45
42:BO:117:ARG:NH1	42:BO:117:ARG:HG2	2.31	0.45
43:BP:91:ARG:NH2	43:BP:96:LEU:HB3	2.32	0.45
50:BW:13:LEU:O	50:BW:17:ARG:HG3	2.16	0.45
33:CF:164:ARG:NH2	53:C1:56:U:O4	2.50	0.45
54:CA:1026:G:O6	54:CA:1036:G:C2	2.70	0.45
54:CA:1156:G:H3'	54:CA:1157:A:H5''	1.99	0.45
54:CA:1200:C:C4'	54:CA:1201:A:H5''	2.40	0.45
54:CA:815:A:HO2'	54:CA:1527:C:C1'	2.30	0.45
54:CA:51:A:N7	54:CA:114:U:O2'	2.47	0.45
54:CA:563:A:O2'	54:CA:566:G:O2'	2.21	0.45
52:CB:18:G:O2'	52:CB:19:G:P	2.74	0.45
52:CB:30:G:O2'	52:CB:31:A:H5'	2.17	0.45
32:CE:219:VAL:HA	32:CE:222:ILE:HD12	1.99	0.45
35:CH:10:MET:CA	35:CH:32:VAL:HG13	2.47	0.45
38:CK:102:ARG:O	38:CK:103:VAL:C	2.54	0.45
38:CK:102:ARG:NH1	38:CK:105:ARG:NH2	2.52	0.45
40:CM:24:VAL:HG21	40:CM:37:PRO:CG	2.42	0.45
40:CM:8:LEU:HB3	40:CM:16:LEU:HD21	1.98	0.45
41:CN:71:LYS:O	41:CN:74:ALA:N	2.48	0.45
46:CS:72:ARG:CD	46:CS:72:ARG:C	2.85	0.45
48:CU:31:LEU:N	48:CU:31:LEU:HD23	2.30	0.45
55:DA:2009:G:C6	55:DA:2010:G:N7	2.85	0.45
55:DA:222:A:N6	55:DA:224:G:C2	2.85	0.45
55:DA:2642:G:O2'	55:DA:2643:G:H5'	2.17	0.45
55:DA:2654:A:C4	55:DA:2656:U:O2	2.70	0.45
55:DA:2692:C:O2'	55:DA:2693:A:H5'	2.17	0.45
55:DA:2776:A:O2'	55:DA:2781:A:O2'	2.28	0.45
55:DA:2804:C:H2'	55:DA:2805:G:H8	1.78	0.45
55:DA:340:A:H2'	55:DA:341:G:H5'	1.99	0.45
55:DA:676:A:C8	55:DA:2069:G:N2	2.55	0.45
55:DA:740:U:H2'	55:DA:741:G:C8	2.52	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:889:C:C2'	55:DA:890:A:O4'	2.48	0.45
55:DA:997:G:O2'	55:DA:998:C:H5'	2.17	0.45
2:DB:43:C:H1'	6:DG:93:THR:O	2.17	0.45
3:DD:79:VAL:HG11	3:DD:111:LEU:CD1	2.47	0.45
3:DD:13:ARG:NH1	3:DD:16:MET:SD	2.89	0.45
3:DD:176:ARG:CG	3:DD:176:ARG:HH11	2.23	0.45
4:DE:48:GLN:HE21	4:DE:48:GLN:HB3	1.56	0.45
4:DE:55:ASN:C	4:DE:57:LYS:N	2.61	0.45
4:DE:1:MET:H2	4:DE:84:PHE:HD2	1.65	0.45
58:DL:65:PHE:HD2	58:DL:66:THR:N	2.14	0.45
10:DN:2:ILE:CD1	10:DN:82:ASN:ND2	2.78	0.45
10:DN:88:ASN:HD21	10:DN:92:GLU:HB2	1.82	0.45
11:DO:18:ARG:HB3	11:DO:19:VAL:H	1.64	0.45
11:DO:95:VAL:O	11:DO:96:THR:HG23	2.17	0.45
18:DS:1:MET:HE2	18:DS:2:GLU:H	1.82	0.45
24:DW:17:SER:CB	24:DW:18:PRO:HA	2.47	0.45
57:DY:112:LEU:O	57:DY:113:GLN:CB	2.65	0.45
57:DY:16:ASN:N	57:DY:19:ARG:NH1	2.62	0.45
57:DY:55:LYS:HD2	57:DY:78:SER:O	2.17	0.45
26:A4:56:VAL:H	26:A4:59:PHE:HB3	1.82	0.45
1:AA:1055:G:O5'	1:AA:1055:G:H8	2.00	0.45
1:AA:141:A:H5''	1:AA:141(A):C:H5	1.82	0.45
1:AA:2439:A:HO2'	1:AA:2440:C:P	2.40	0.45
1:AA:2770:G:H5'	1:AA:2771:C:OP2	2.17	0.45
1:AA:2849:U:O2'	1:AA:2850:A:P	2.75	0.45
1:AA:288:C:H2'	1:AA:289:A:C1'	2.47	0.45
1:AA:621:A:C2	1:AA:622:G:N9	2.85	0.45
1:AA:768:G:H2'	1:AA:769:G:C8	2.51	0.45
1:AA:768:G:O2'	1:AA:769:G:H5'	2.17	0.45
1:AA:945:A:N3	1:AA:945:A:H2'	2.31	0.45
1:AA:977:G:O2'	1:AA:978:G:H5'	2.17	0.45
3:AD:121:PRO:HB3	3:AD:135:PHE:CE1	2.52	0.45
1:AA:781:A:C8	3:AD:219:PRO:HG3	2.52	0.45
1:AA:2820:A:N6	4:AE:192:ASN:CB	2.75	0.45
5:AF:32:LEU:HD23	5:AF:32:LEU:C	2.38	0.45
6:AG:48:GLU:O	6:AG:48:GLU:HG3	2.17	0.45
11:AO:110:TYR:HD2	11:AO:111:ARG:HH21	1.64	0.45
12:AP:10:ARG:HG3	12:AP:10:ARG:HH11	1.82	0.45
12:AP:120:ILE:HA	12:AP:120:ILE:HD13	1.82	0.45
1:AA:910:A:N7	12:AP:13:GLN:HG3	2.32	0.45
14:AQ:54:LEU:O	14:AQ:56:LEU:N	2.46	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:AR:115:ARG:N	15:AR:115:ARG:HD3	2.18	0.45
15:AR:124:ASP:O	15:AR:128:GLU:HG3	2.16	0.45
15:AR:48:ILE:H	15:AR:48:ILE:HD12	1.82	0.45
19:AT:63:LYS:CE	19:AT:63:LYS:N	2.74	0.45
20:AU:20:TYR:CE2	20:AU:42:VAL:N	2.85	0.45
20:AU:81:LYS:NZ	20:AU:98:VAL:HG21	2.31	0.45
21:AV:12:GLY:O	21:AV:13:GLU:O	2.35	0.45
24:AW:48:HIS:HD2	24:AW:48:HIS:C	2.20	0.45
31:BA:1015:A:N6	31:BA:1016:A:N6	2.64	0.45
31:BA:1268:A:N3	31:BA:1326:C:O2'	2.46	0.45
31:BA:1321:C:H5'	31:BA:1322:C:H5''	1.99	0.45
31:BA:511:C:N3	31:BA:512:U:C4	2.85	0.45
31:BA:983:A:N1	31:BA:1222:G:N2	2.64	0.45
52:BB:17:C:H5'	52:BB:18:G:OP2	2.17	0.45
52:BD:48:C:H6	52:BD:48:C:H3'	1.81	0.45
33:BF:63:ASN:O	33:BF:64:VAL:CB	2.65	0.45
31:BA:619:U:N3	34:BG:134:ASP:OD2	2.34	0.45
34:BG:3:ARG:NH2	34:BG:5:ILE:CD1	2.79	0.45
35:BH:147:ASP:OD2	35:BH:147:ASP:N	2.49	0.45
35:BH:147:ASP:O	35:BH:151:LEU:HD23	2.17	0.45
38:BK:29:SER:HB3	38:BK:32:LYS:CB	2.46	0.45
39:BL:117:HIS:HD2	39:BL:123:PRO:HA	1.82	0.45
39:BL:27:THR:HA	39:BL:31:GLN:O	2.17	0.45
39:BL:78:LYS:CB	39:BL:78:LYS:NZ	2.80	0.45
42:BO:57:LYS:HG2	42:BO:67:THR:HG22	1.99	0.45
43:BP:3:ARG:HA	43:BP:8:GLU:O	2.17	0.45
44:BQ:39:LEU:CD1	44:BQ:47:LEU:HD12	2.47	0.45
49:BV:15:LEU:HD22	49:BV:31:ILE:HD11	1.99	0.45
49:BV:19:VAL:HG22	49:BV:44:MET:HB3	1.98	0.45
53:C1:38:U:C2'	53:C1:39:U:H5'	2.47	0.45
54:CA:1004:A:C6	54:CA:1025:U:H1'	2.51	0.45
54:CA:291:C:O2'	54:CA:292:G:H5'	2.17	0.45
54:CA:544:G:C6	54:CA:545:C:C4	3.05	0.45
54:CA:627:G:H2'	54:CA:628:G:H8	1.82	0.45
54:CA:782:A:C8	54:CA:783:C:C5	3.05	0.45
54:CA:819:A:C5'	54:CA:820:U:OP2	2.58	0.45
52:CC:14:A:C2'	52:CC:15:G:H5'	2.47	0.45
32:CE:21:ARG:HG2	32:CE:21:ARG:O	2.17	0.45
33:CF:76:VAL:HG21	33:CF:103:VAL:CG1	2.47	0.45
34:CG:50:ARG:NH1	53:C1:57:U:O2'	2.50	0.45
35:CH:79:GLU:HB3	35:CH:93:PRO:HD2	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:CJ:50:ILE:O	37:CJ:50:ILE:HG22	2.16	0.45
38:CK:39:LEU:HD11	38:CK:111:ILE:HD11	1.99	0.45
42:CO:28:LYS:HE3	42:CO:33:ARG:HH22	1.82	0.45
42:CO:71:PRO:CG	42:CO:99:HIS:HD2	2.26	0.45
43:CP:120:LYS:CD	43:CP:120:LYS:N	2.60	0.45
43:CP:90:LEU:HD13	49:CV:78:ARG:NH2	2.31	0.45
48:CU:51:LEU:HB2	48:CU:56:THR:CG2	2.48	0.45
26:D4:69:LYS:CD	26:D4:70:GLY:H	2.25	0.45
27:D5:40:LYS:HG2	27:D5:46:CYS:CB	2.45	0.45
28:D6:33:LYS:C	28:D6:35:GLU:H	2.18	0.45
55:DA:1011:G:O2'	55:DA:1013:C:C5'	2.65	0.45
55:DA:1046:A:H5''	55:DA:1046:A:N3	2.31	0.45
55:DA:1047:G:H2'	55:DA:1110:G:N1	2.31	0.45
55:DA:1049:C:H6	55:DA:1049:C:H5'	1.80	0.45
55:DA:1106:G:H4'	57:DY:53:VAL:HG11	1.97	0.45
55:DA:1192:G:O2'	55:DA:1193:G:H5'	2.17	0.45
55:DA:1479:G:C5'	55:DA:1558:A:H2	2.29	0.45
55:DA:1498:C:O4'	55:DA:1577:C:C4'	2.65	0.45
55:DA:2079:U:H2'	55:DA:2080:G:O4'	2.16	0.45
55:DA:2168:G:O4'	55:DA:2168:G:P	2.75	0.45
55:DA:2405:G:HO2'	55:DA:2406:U:P	2.39	0.45
55:DA:2680:C:N3	55:DA:2681:C:N4	2.65	0.45
55:DA:2695:C:H2'	55:DA:2696:U:H6	1.81	0.45
55:DA:347:A:H2'	55:DA:348:G:C8	2.52	0.45
55:DA:469:G:O6	29:D7:37:LYS:CE	2.64	0.45
55:DA:503:A:O2'	55:DA:504:U:OP2	2.33	0.45
2:DB:81:G:N2	2:DB:82:G:C5	2.85	0.45
3:DD:69:ARG:CD	3:DD:105:ILE:HD11	2.39	0.45
3:DD:227:ASN:O	3:DD:230:ASP:OD2	2.34	0.45
3:DD:182:LEU:HB2	3:DD:271:ILE:HB	1.99	0.45
4:DE:55:ASN:O	4:DE:57:LYS:N	2.47	0.45
5:DF:155:LEU:HD12	5:DF:174:VAL:HG22	1.99	0.45
6:DG:151:ALA:HB3	6:DG:153:ARG:HH11	1.80	0.45
7:DH:35:VAL:CG2	7:DH:75:ALA:HB2	2.47	0.45
56:DI:7:ARG:CG	56:DI:7:ARG:HH11	2.30	0.45
8:DK:112:LYS:O	8:DK:113:ARG:CB	2.65	0.45
8:DK:10:GLU:OE2	8:DK:11:ASN:N	2.50	0.45
8:DK:25:TYR:CE2	8:DK:29:TYR:CD2	2.95	0.45
58:DL:63:ARG:HB3	58:DL:64:SER:H	1.58	0.45
58:DL:58:THR:HG21	58:DL:66:THR:CG2	2.47	0.45
58:DL:72:PRO:HD2	58:DL:73:PRO:HD3	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:DO:106:LEU:HD23	11:DO:106:LEU:HA	1.69	0.45
14:DQ:15:ARG:HG3	14:DQ:19:LYS:HE2	1.99	0.45
18:DS:1:MET:C	18:DS:64:MET:HE3	2.36	0.45
20:DU:51:VAL:HG22	20:DU:57:GLN:HA	1.99	0.45
24:DW:47:ASN:N	24:DW:47:ASN:ND2	2.47	0.45
57:DY:8:GLU:CD	57:DY:52:PHE:HD1	2.18	0.45
23:DZ:19:GLN:O	23:DZ:35:THR:N	2.50	0.45
23:DZ:7:ILE:HD13	23:DZ:69:LYS:HB3	1.99	0.45
13:A0:55:ALA:HA	13:A0:80:PHE:CE2	2.51	0.44
17:A2:52:VAL:O	17:A2:55:ALA:HB3	2.17	0.44
17:A2:57:VAL:HG12	17:A2:99:ILE:HA	1.99	0.44
28:A6:17:LYS:HA	28:A6:17:LYS:CE	2.44	0.44
30:A8:32:LEU:HB2	30:A8:36:LYS:HZ1	1.82	0.44
1:AA:1141:U:H1'	1:AA:1142(A):A:C2	2.52	0.44
1:AA:1354:A:C8	1:AA:1355:G:C8	3.05	0.44
1:AA:1861:G:O2'	1:AA:1862:G:H5'	2.17	0.44
1:AA:1946:U:C2	1:AA:1947:C:C5	3.05	0.44
1:AA:2271:G:H2'	1:AA:2272:U:H6	1.82	0.44
1:AA:2478:A:H3'	1:AA:2479:G:H8	1.82	0.44
1:AA:2614:A:H5''	1:AA:2615:U:OP1	2.17	0.44
1:AA:270(C):C:O2'	1:AA:270(D):C:H5'	2.17	0.44
1:AA:2791:C:C4	1:AA:2792:G:N7	2.85	0.44
1:AA:2898:U:O2'	1:AA:2899:G:H5'	2.17	0.44
1:AA:736:C:H2'	1:AA:737:C:H6	1.82	0.44
1:AA:858:U:O2'	1:AA:2268:A:H1'	2.17	0.44
1:AA:99:U:C4'	1:AA:102:G:H1'	2.47	0.44
4:AE:11:MET:CA	4:AE:24:THR:HA	2.47	0.44
4:AE:51:PHE:O	4:AE:52:LEU:HB2	2.17	0.44
5:AF:164:ARG:HG3	5:AF:175:THR:CG2	2.47	0.44
5:AF:155:LEU:CD2	5:AF:186:ILE:HA	2.47	0.44
5:AF:20:LEU:HD13	5:AF:199:TRP:CZ3	2.52	0.44
6:AG:16:ARG:NH1	6:AG:16:ARG:CG	2.78	0.44
7:AH:16:SER:O	7:AH:17:VAL:HB	2.18	0.44
8:AK:5:LEU:CD1	8:AK:5:LEU:H	2.24	0.44
10:AN:12:ASP:HA	10:AN:98:VAL:HA	1.99	0.44
10:AN:32:TYR:CD1	10:AN:32:TYR:N	2.84	0.44
11:AO:138:LEU:HD13	11:AO:144:GLU:HG2	1.98	0.44
11:AO:13:ASN:C	11:AO:15:ARG:N	2.70	0.44
11:AO:57:THR:O	11:AO:57:THR:CG2	2.59	0.44
14:AQ:60:GLY:O	14:AQ:61:ASN:HB2	2.17	0.44
20:AU:86:ARG:O	20:AU:93:GLY:N	2.49	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:AZ:92:LYS:O	23:AZ:93:GLU:C	2.54	0.44
31:BA:1203:C:H2'	31:BA:1204:A:O4'	2.17	0.44
31:BA:1535:C:H2'	31:BA:1536:C:H5'	1.98	0.44
31:BA:190:G:O6	31:BA:264:U:H5''	2.16	0.44
31:BA:22:G:H2'	31:BA:23:C:C6	2.52	0.44
31:BA:265:G:N2	31:BA:267:C:H5''	2.31	0.44
31:BA:280:C:C2'	31:BA:280:C:O2	2.65	0.44
31:BA:29:G:O2'	31:BA:30:U:H5'	2.16	0.44
31:BA:625:G:C4	31:BA:626:U:C5	3.05	0.44
31:BA:721:G:H1'	31:BA:722:A:N1	2.32	0.44
32:BE:122:PHE:C	32:BE:124:SER:H	2.21	0.44
32:BE:237:ALA:O	32:BE:238:LEU:HB3	2.17	0.44
33:BF:73:PRO:HA	33:BF:76:VAL:CG1	2.47	0.44
33:BF:90:GLU:OE2	33:BF:93:LYS:HD2	2.17	0.44
35:BH:127:ASN:O	35:BH:128:PRO:C	2.55	0.44
35:BH:55:VAL:O	35:BH:58:ALA:HB3	2.17	0.44
36:BI:8:ILE:HG22	36:BI:10:LEU:HD11	1.98	0.44
36:BI:48:LEU:HD13	36:BI:52:ILE:HG13	1.98	0.44
31:BA:1375:A:H4'	37:BJ:29:LYS:HZ2	1.82	0.44
38:BK:132:GLU:O	38:BK:134:ILE:N	2.51	0.44
39:BL:62:TYR:C	39:BL:63:ILE:HD12	2.37	0.44
39:BL:66:ARG:CZ	39:BL:66:ARG:CB	2.94	0.44
45:BR:61:GLY:O	45:BR:65:ARG:HG3	2.16	0.44
45:BR:7:GLU:O	45:BR:11:VAL:HG23	2.17	0.44
46:BS:48:TRP:CE3	46:BS:49:LEU:HB2	2.51	0.44
48:BU:22:VAL:HG11	48:BU:56:THR:HA	1.98	0.44
26:A4:63:TYR:HE2	49:BV:41:VAL:CB	2.30	0.44
50:BW:8:ARG:HG2	50:BW:8:ARG:O	2.17	0.44
33:CF:131:ARG:HH21	53:C1:56:U:H3	1.65	0.44
54:CA:1228:C:P	43:CP:108:ARG:HH22	2.40	0.44
54:CA:1306:A:O2'	54:CA:1307:U:H5'	2.17	0.44
54:CA:1347:G:H5''	39:CL:107:ARG:HA	2.00	0.44
54:CA:1347:G:H22	54:CA:1374:A:P	2.39	0.44
54:CA:1536:C:H1'	53:C1:37:G:H22	1.78	0.44
54:CA:1542:U:HO3'	54:CA:1542:U:P	2.29	0.44
54:CA:191(E):G:H2'	54:CA:191(F):U:C6	2.52	0.44
54:CA:352:C:H4'	54:CA:354:G:OP1	2.17	0.44
54:CA:531:U:C5'	54:CA:532:A:OP1	2.65	0.44
54:CA:585:G:OP1	47:CT:37:LYS:HE2	2.17	0.44
54:CA:609:A:C2'	54:CA:610:G:H5'	2.46	0.44
54:CA:630:G:C8	54:CA:630:G:C4'	3.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1191:A:P	33:CF:3:ASN:HD21	2.40	0.44
33:CF:79:ARG:NH1	33:CF:82:GLU:HG3	2.31	0.44
54:CA:1346:A:C4	37:CJ:10:ARG:NH1	2.84	0.44
43:CP:123:ALA:HB1	43:CP:124:PRO:HD2	1.97	0.44
43:CP:88:ARG:HD3	43:CP:98:VAL:CG1	2.48	0.44
46:CS:83:GLU:HA	46:CS:83:GLU:OE2	2.17	0.44
50:CW:26:ASN:HB2	50:CW:71:THR:CG2	2.33	0.44
55:DA:1278:A:OP1	13:D0:36:THR:HG22	2.17	0.44
16:D1:88:ILE:CD1	16:D1:88:ILE:H	2.25	0.44
22:D3:11:ARG:CZ	22:D3:11:ARG:HB2	2.45	0.44
55:DA:2331:G:C4'	22:D3:42:GLY:HA3	2.47	0.44
26:D4:67:TYR:HB3	26:D4:68:ARG:H	1.38	0.44
55:DA:1010:A:H1'	55:DA:1153:C:H1'	1.98	0.44
55:DA:1487:G:C2	55:DA:1488:G:C8	3.05	0.44
55:DA:1689:A:N6	55:DA:1698:A:H2	1.87	0.44
55:DA:2465:C:O2'	55:DA:2466:C:H5'	2.17	0.44
55:DA:270(J):G:H2'	55:DA:270(K):C:O4'	2.17	0.44
55:DA:2808:U:H2'	55:DA:2809:A:H5'	1.98	0.44
55:DA:2866:U:O2'	55:DA:2867:G:P	2.75	0.44
55:DA:439:G:H2'	55:DA:440:G:H8	1.81	0.44
2:DB:60:C:O2'	2:DB:61:G:H5'	2.16	0.44
3:DD:13:ARG:NH2	3:DD:16:MET:HE3	2.32	0.44
3:DD:223:GLY:C	3:DD:225:ALA:N	2.71	0.44
3:DD:35:LYS:HD3	3:DD:63:ARG:HA	1.96	0.44
3:DD:95:LEU:C	3:DD:95:LEU:HD12	2.37	0.44
4:DE:72:VAL:O	4:DE:73:GLU:C	2.55	0.44
6:DG:110:ALA:O	6:DG:111:LEU:C	2.56	0.44
6:DG:41:GLN:HB3	6:DG:43:LEU:HD11	1.99	0.44
56:DI:29:GLU:CG	56:DJ:6:GLU:CD	2.85	0.44
56:DJ:13:SER:CB	56:DJ:17:VAL:HG11	2.15	0.44
8:DK:52:ARG:HH11	8:DK:52:ARG:HB2	1.81	0.44
58:DL:126:MET:O	58:DL:129:GLY:N	2.37	0.44
58:DL:125:ARG:CD	58:DL:132:ARG:NH2	2.79	0.44
5:DF:34:TRP:CA	11:DO:6:LEU:HD12	2.47	0.44
11:DO:7:ARG:HA	11:DO:8:PRO:HD3	1.74	0.44
11:DO:88:LEU:HD12	11:DO:95:VAL:HG11	1.98	0.44
15:DR:45:PHE:HE2	15:DR:63:VAL:HB	1.82	0.44
24:DW:47:ASN:HD22	24:DW:47:ASN:N	1.91	0.44
55:DA:102:G:OP1	24:DW:7:ARG:NH2	2.50	0.44
57:DY:50:ARG:H	57:DY:83:TYR:CB	2.29	0.44
23:DZ:11:ARG:HB2	23:DZ:12:PRO:HD2	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:DZ:8:SER:OG	23:DZ:10:LYS:HG3	2.17	0.44
26:A4:38:LYS:C	26:A4:40:HIS:H	2.09	0.44
1:AA:1001:A:C2'	1:AA:1002:G:H5'	2.48	0.44
1:AA:1028:A:H2'	1:AA:1029:A:C8	2.52	0.44
1:AA:1204:A:H2	1:AA:1241:A:C2	2.35	0.44
1:AA:1381:G:C2'	1:AA:1382:G:H5'	2.46	0.44
1:AA:1421:G:C2	1:AA:1422:G:N7	2.85	0.44
1:AA:1512:G:H2'	1:AA:1513:C:H6	1.81	0.44
1:AA:1332:G:H21	1:AA:1610:A:H8	1.63	0.44
1:AA:2091:U:P	1:AA:2092:U:H3'	2.57	0.44
1:AA:2259:G:H1'	1:AA:2427:C:C2	2.52	0.44
1:AA:2555:U:H2'	1:AA:2556:C:H5'	1.98	0.44
1:AA:2712:U:H2'	1:AA:2713:A:H5''	1.99	0.44
1:AA:35:G:H1'	1:AA:454:A:C4	2.52	0.44
1:AA:503:A:H4'	1:AA:504:U:H5''	1.99	0.44
1:AA:527:C:O5'	1:AA:2779:U:H5	2.01	0.44
1:AA:607:U:C5	1:AA:608:A:N7	2.85	0.44
1:AA:70:G:H5''	1:AA:75:G:N2	2.33	0.44
2:AB:39:A:C2	2:AB:44:G:N3	2.86	0.44
3:AD:35:LYS:CE	3:AD:104:TYR:HB2	2.46	0.44
3:AD:257:LEU:C	3:AD:257:LEU:HD23	2.37	0.44
4:AE:23:VAL:O	4:AE:24:THR:OG1	2.28	0.44
5:AF:162:LEU:HD12	5:AF:162:LEU:H	1.82	0.44
1:AA:323:G:H2'	5:AF:169:ASN:OD1	2.17	0.44
5:AF:61:GLY:O	5:AF:77:ASP:CB	2.65	0.44
7:AH:86:GLU:O	7:AH:87:LEU:CG	2.65	0.44
8:AK:81:VAL:H	8:AK:143:SER:HA	1.82	0.44
9:AM:53:VAL:HG11	9:AM:128:HIS:CB	2.47	0.44
11:AO:101:VAL:HG21	11:AO:108:LYS:H	1.81	0.44
12:AP:35:VAL:HG23	12:AP:100:GLY:O	2.17	0.44
12:AP:58:PHE:O	12:AP:59:ARG:C	2.56	0.44
18:AS:20:VAL:O	18:AS:23:LEU:N	2.41	0.44
1:AA:1599:C:OP2	19:AT:36:LYS:HD3	2.17	0.44
20:AU:75:ILE:HG12	20:AU:76:CYS:N	2.32	0.44
21:AV:148:ASP:O	21:AV:149:SER:OG	2.32	0.44
21:AV:157:LEU:HD22	21:AV:163:LEU:HD22	1.98	0.44
21:AV:157:LEU:O	21:AV:161:VAL:HG21	2.17	0.44
21:AV:178:GLU:OE1	21:AV:180:VAL:C	2.55	0.44
24:AW:24:LEU:O	24:AW:24:LEU:HD23	2.17	0.44
23:AZ:82:LEU:CG	23:AZ:83:GLU:N	2.73	0.44
31:BA:1006:C:C2	31:BA:1007:C:C5	3.05	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:285:G:O2'	31:BA:286:G:H5'	2.17	0.44
31:BA:464:G:O5'	31:BA:464:G:H8	2.01	0.44
31:BA:624:C:H2'	31:BA:625:G:C8	2.52	0.44
31:BA:689:C:C2'	31:BA:690:G:C5'	2.91	0.44
31:BA:693:G:H2'	31:BA:694:A:O4'	2.17	0.44
31:BA:6:G:C4	35:BH:119:LEU:HD11	2.52	0.44
31:BA:945:G:C2	31:BA:946:A:C8	3.05	0.44
52:BB:75:C:C3'	52:BB:75:C:C6	2.99	0.44
32:BE:137:ARG:HD3	32:BE:137:ARG:O	2.17	0.44
33:BF:54:ARG:HG2	33:BF:54:ARG:NH1	2.27	0.44
34:BG:61:LYS:HZ2	34:BG:62:GLN:HE21	1.64	0.44
38:BK:39:LEU:HD12	38:BK:39:LEU:HA	1.81	0.44
39:BL:15:ALA:HB2	39:BL:65:VAL:CG2	2.47	0.44
39:BL:46:ALA:HA	39:BL:78:LYS:HZ2	1.81	0.44
39:BL:7:THR:HB	39:BL:83:ARG:NH1	2.32	0.44
31:BA:1228:C:H4'	43:BP:116:THR:O	2.18	0.44
45:BR:2:PRO:HG2	45:BR:3:ILE:HD13	1.99	0.44
36:BI:99:ALA:N	48:BU:31:LEU:HD22	2.31	0.44
26:A4:59:PHE:CE1	49:BV:67:VAL:HB	2.52	0.44
54:CA:1065:U:O2'	54:CA:1066:C:OP2	2.26	0.44
54:CA:1129:C:H5'	54:CA:1130:A:C5'	2.48	0.44
54:CA:157:G:C2'	54:CA:158:G:H5'	2.47	0.44
54:CA:302:G:N3	54:CA:556:C:H4'	2.33	0.44
54:CA:720:C:C5	54:CA:721:G:H2'	2.53	0.44
54:CA:750:G:N3	45:CR:23:GLY:HA3	2.32	0.44
54:CA:836:G:C6	54:CA:851:G:C6	3.06	0.44
54:CA:872:A:C4	54:CA:874:G:N7	2.84	0.44
52:CB:8:U:H2'	52:CB:9:A:OP2	2.18	0.44
52:CC:44:G:O4'	52:CC:44:G:N3	2.50	0.44
52:CD:56:C:H6	55:DA:2169:A:N7	2.15	0.44
32:CE:9:GLU:HA	32:CE:12:GLU:OE1	2.17	0.44
34:CG:140:VAL:HG12	34:CG:144:ASP:OD2	2.17	0.44
34:CG:76:ARG:HH11	34:CG:76:ARG:HG2	1.82	0.44
35:CH:6:PHE:HA	35:CH:6:PHE:HD2	1.67	0.44
37:CJ:146:GLU:O	37:CJ:149:ARG:HB2	2.18	0.44
40:CM:21:GLN:HG2	40:CM:21:GLN:O	2.16	0.44
40:CM:38:ILE:CD1	40:CM:71:LEU:HB3	2.47	0.44
41:CN:121:PRO:HD2	41:CN:126:ARG:HD3	1.99	0.44
42:CO:27:LEU:O	42:CO:29:GLY:N	2.49	0.44
54:CA:1358:U:OP1	44:CQ:35:ARG:HG3	2.17	0.44
46:CS:48:TRP:CZ2	46:CS:76:GLN:OE1	2.70	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
46:CS:82:GLN:O	46:CS:83:GLU:CB	2.65	0.44
48:CU:73:ALA:O	48:CU:76:LEU:HB2	2.17	0.44
16:D1:35:ALA:O	16:D1:39:LEU:HG	2.16	0.44
55:DA:446:G:OP1	16:D1:3:ARG:HD3	2.16	0.44
26:D4:12:ALA:HB1	26:D4:29:PRO:HA	1.99	0.44
6:DG:118:ARG:O	26:D4:43:TYR:CZ	2.70	0.44
26:D4:3:GLU:HG3	26:D4:4:GLY:N	2.31	0.44
30:D8:16:ILE:HG23	30:D8:16:ILE:O	2.16	0.44
55:DA:1022:G:C5	55:DA:1140:C:C4	3.05	0.44
55:DA:1077:A:O4'	58:DL:93:ARG:NH2	2.51	0.44
55:DA:1131:G:HO2'	55:DA:1132:A:H8	1.65	0.44
55:DA:1284:A:H2'	55:DA:1285:G:O4'	2.16	0.44
55:DA:1299:G:C5'	55:DA:1300:U:OP1	2.65	0.44
55:DA:1467:C:C5	55:DA:1546:C:H2'	2.53	0.44
55:DA:1629:U:H2'	55:DA:1630:G:O4'	2.17	0.44
55:DA:1820:U:H1'	3:DD:202:LYS:HB3	1.98	0.44
55:DA:2019:A:C2'	55:DA:2020:A:O5'	2.64	0.44
55:DA:205:G:C2'	55:DA:206:U:OP2	2.65	0.44
55:DA:2131:G:O5'	55:DA:2133:G:OP1	2.35	0.44
55:DA:2182:G:O2'	55:DA:2183:C:H5'	2.16	0.44
55:DA:226:G:H1'	55:DA:228:A:H61	1.80	0.44
55:DA:2292:C:H2'	55:DA:2293:C:H6	1.80	0.44
55:DA:228:A:H2'	55:DA:229:A:OP1	2.17	0.44
55:DA:388:G:H5'	23:DZ:25:LYS:HB3	1.98	0.44
55:DA:534:U:O2'	16:D1:49:HIS:HD2	2.00	0.44
55:DA:614:U:H2'	55:DA:615:G:OP1	2.16	0.44
2:DB:31:C:O2	2:DB:31:C:C2'	2.64	0.44
4:DE:116:VAL:CG2	4:DE:122:PHE:CG	3.00	0.44
4:DE:7:VAL:CG2	4:DE:27:LEU:HD23	2.42	0.44
4:DE:69:LYS:CD	4:DE:89:ASP:OD1	2.65	0.44
5:DF:46:ARG:CG	5:DF:46:ARG:NH1	2.72	0.44
5:DF:74:ARG:HG2	5:DF:74:ARG:O	2.16	0.44
7:DH:146:ALA:O	7:DH:150:ALA:N	2.43	0.44
7:DH:153:LYS:HA	7:DH:153:LYS:CE	2.47	0.44
8:DK:60:GLU:CG	8:DK:61:ARG:HH22	2.27	0.44
11:DO:107:LYS:HB3	11:DO:110:TYR:HD2	1.82	0.44
15:DR:57:PHE:O	15:DR:58:ASN:C	2.56	0.44
21:DV:106:GLY:O	21:DV:107:THR:CB	2.65	0.44
21:DV:35:ARG:HH11	21:DV:35:ARG:CB	2.30	0.44
57:DY:64:LYS:HD2	57:DY:64:LYS:N	2.32	0.44
16:A1:50:ARG:HH11	17:A2:72:VAL:HB	1.82	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1086:A:H4'	1:AA:1103:A:N6	2.17	0.44
1:AA:1087:G:C4	1:AA:1089:G:O2'	2.70	0.44
1:AA:1533:C:H3'	1:AA:1534:G:O4'	2.16	0.44
1:AA:141:A:N6	1:AA:1596:A:H5'	2.32	0.44
1:AA:2035:G:C4'	1:AA:2036:C:OP2	2.65	0.44
1:AA:2170:A:C2'	1:AA:2171:A:H5'	2.48	0.44
1:AA:2232:U:P	23:AZ:40:ARG:HH12	2.41	0.44
1:AA:2250:G:C5	12:AP:82:ARG:HG3	2.53	0.44
1:AA:2358:G:N2	1:AA:2359:C:H1'	2.32	0.44
1:AA:236:C:H2'	1:AA:237:C:C6	2.53	0.44
1:AA:2790:A:O2'	1:AA:2893:G:O2'	2.33	0.44
1:AA:90:U:H2'	1:AA:91:A:C5'	2.47	0.44
3:AD:124:PRO:HG2	3:AD:129:ASN:ND2	2.33	0.44
1:AA:1790:C:H4'	3:AD:209:ALA:CB	2.47	0.44
3:AD:224:ALA:N	3:AD:233:HIS:HB2	2.32	0.44
3:AD:34:VAL:O	3:AD:35:LYS:CB	2.64	0.44
4:AE:118:LYS:H	4:AE:121:ASN:H	1.65	0.44
4:AE:47:VAL:HG12	4:AE:49:LEU:HD23	2.00	0.44
4:AE:7:VAL:HG13	4:AE:51:PHE:CE2	2.53	0.44
4:AE:79:ARG:HB3	4:AE:80:GLU:H	1.60	0.44
6:AG:32:PRO:HB3	6:AG:163:ALA:HB2	1.99	0.44
6:AG:78:SER:OG	52:BC:19:G:N2	2.50	0.44
7:AH:136:ILE:O	7:AH:137:ASP:HB2	2.17	0.44
7:AH:40:GLU:HB3	7:AH:41:MET:SD	2.58	0.44
7:AH:70:THR:HA	7:AH:73:ALA:HB3	1.99	0.44
8:AK:118:LYS:O	8:AK:119:PRO:O	2.36	0.44
9:AM:128:HIS:CE1	9:AM:134:ARG:CZ	3.00	0.44
9:AM:43:THR:HB	9:AM:46:VAL:HG11	1.99	0.44
11:AO:106:LEU:HD11	11:AO:112:LEU:CD2	2.40	0.44
12:AP:35:VAL:HG23	12:AP:101:ARG:O	2.17	0.44
12:AP:41:TRP:O	12:AP:42:ILE:C	2.55	0.44
14:AQ:52:SER:HB2	14:AQ:55:ALA:HB3	2.00	0.44
18:AS:29:LEU:HD11	18:AS:51:LEU:HD11	2.00	0.44
18:AS:51:LEU:HD13	18:AS:51:LEU:C	2.36	0.44
20:AU:9:LYS:O	20:AU:27:VAL:CG2	2.66	0.44
20:AU:27:VAL:O	20:AU:27:VAL:CG2	2.65	0.44
21:AV:145:GLU:O	21:AV:146:ILE:HG12	2.16	0.44
53:B1:30:C:H2'	53:B1:31:A:C8	2.53	0.44
31:BA:1176:A:O2'	31:BA:1177:G:H5'	2.17	0.44
31:BA:1328:C:H2'	31:BA:1329:A:H5'	1.99	0.44
31:BA:1466:C:H2'	31:BA:1467:G:O4'	2.16	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:153:C:H2'	31:BA:154:C:C6	2.51	0.44
31:BA:198:G:H2'	31:BA:199:G:H8	1.82	0.44
31:BA:312:C:H2'	31:BA:313:A:C8	2.52	0.44
31:BA:428:G:HO2'	31:BA:429:U:P	2.40	0.44
31:BA:633:G:H2'	31:BA:634:C:H6	1.81	0.44
31:BA:713:G:H2'	31:BA:714:G:C8	2.53	0.44
31:BA:955:U:O2'	49:BV:83:HIS:HB2	2.18	0.44
52:BB:21:A:N7	52:BB:46:G:C6	2.86	0.44
32:BE:22:LYS:CA	32:BE:22:LYS:CE	2.93	0.44
32:BE:38:GLY:O	32:BE:39:ILE:O	2.35	0.44
33:BF:58:GLU:O	33:BF:59:ARG:HG3	2.18	0.44
33:BF:69:HIS:CD2	33:BF:104:GLN:HB2	2.53	0.44
34:BG:88:VAL:O	34:BG:92:VAL:HG23	2.17	0.44
36:BI:82:ARG:HG2	36:BI:82:ARG:NH1	2.32	0.44
40:BM:35:SER:OG	40:BM:73:ASP:HB2	2.17	0.44
40:BM:70:ARG:HG3	40:BM:70:ARG:HH11	1.81	0.44
42:BO:60:LEU:O	42:BO:61:THR:C	2.55	0.44
44:BQ:10:ALA:HB1	44:BQ:23:ARG:HB3	1.98	0.44
31:BA:1217:C:OP1	44:BQ:9:LYS:HE3	2.17	0.44
31:BA:657:G:H4'	45:BR:28:GLN:HG2	1.99	0.44
45:BR:33:THR:HG23	45:BR:63:ARG:HH12	1.82	0.44
48:BU:32:ARG:HG3	48:BU:32:ARG:O	2.17	0.44
48:BU:72:ARG:O	48:BU:75:ILE:N	2.50	0.44
54:CA:1064:G:HO2'	54:CA:1065:U:P	2.39	0.44
54:CA:113:G:H2'	54:CA:114:U:C6	2.53	0.44
54:CA:1540:U:C2	54:CA:1541:U:H1'	2.52	0.44
54:CA:579:G:H2'	54:CA:580:U:C6	2.53	0.44
54:CA:703:G:H4'	54:CA:704:A:C5'	2.46	0.44
54:CA:857:C:H2'	54:CA:858:G:O4'	2.17	0.44
54:CA:955:U:O2'	54:CA:956:U:H5'	2.17	0.44
32:CE:167:PRO:HG2	32:CE:192:SER:OG	2.18	0.44
32:CE:84:GLU:HG3	32:CE:215:LEU:HD12	1.99	0.44
34:CG:148:VAL:CG1	34:CG:149:ALA:N	2.80	0.44
34:CG:8:VAL:C	34:CG:10:ARG:H	2.21	0.44
36:CI:28:ARG:HG3	36:CI:28:ARG:HH11	1.82	0.44
37:CJ:23:VAL:O	37:CJ:27:ILE:CD1	2.66	0.44
37:CJ:27:ILE:HD11	37:CJ:43:PHE:CD2	2.53	0.44
37:CJ:79:ARG:HH11	37:CJ:79:ARG:HG2	1.81	0.44
39:CL:114:TYR:CE1	40:CM:60:ARG:N	2.84	0.44
42:CO:27:LEU:HD13	42:CO:28:LYS:N	2.28	0.44
44:CQ:34:TYR:O	44:CQ:35:ARG:C	2.54	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:D0:37:THR:OG1	13:D0:40:LYS:HG3	2.17	0.44
13:D0:41:ALA:C	13:D0:43:GLU:N	2.70	0.44
13:D0:85:PRO:C	13:D0:87:TYR:N	2.70	0.44
16:D1:83:LEU:HD13	16:D1:113:ALA:HB2	1.98	0.44
55:DA:1188:U:C4'	17:D2:79:VAL:HG22	2.47	0.44
26:D4:32:TYR:CD2	26:D4:33:VAL:N	2.79	0.44
29:D7:12:ARG:NH1	29:D7:12:ARG:HG3	2.33	0.44
30:D8:17:THR:HG22	30:D8:23:VAL:HG23	1.98	0.44
55:DA:1027:A:N6	55:DA:1126:A:C1'	2.79	0.44
55:DA:1057:A:N7	55:DA:1086:A:N3	2.65	0.44
55:DA:1133:U:O4	55:DA:2026:C:H1'	2.17	0.44
55:DA:1181:C:C2'	55:DA:1182:A:H5'	2.46	0.44
55:DA:197:A:N6	55:DA:2430:A:H2'	2.32	0.44
55:DA:2020:A:O2'	55:DA:2021:C:H3'	2.16	0.44
55:DA:2198:A:O2'	55:DA:2199:A:P	2.75	0.44
55:DA:1889:A:N1	55:DA:2234:G:H1'	2.32	0.44
55:DA:2302:G:H21	6:DG:126:ASP:HB2	1.82	0.44
55:DA:2512:C:H2'	55:DA:2513:G:O4'	2.17	0.44
55:DA:2637:U:H2'	55:DA:2638:G:O4'	2.17	0.44
55:DA:2893:G:H5''	55:DA:2894:G:C4'	2.47	0.44
55:DA:548:A:C2'	55:DA:549:G:H5'	2.46	0.44
55:DA:70:G:O2'	55:DA:71:A:OP2	2.25	0.44
55:DA:828:U:H2'	55:DA:828:U:O2	2.18	0.44
55:DA:884:C:O2'	55:DA:885:C:OP1	2.35	0.44
2:DB:24:G:H1'	2:DB:27:C:N4	2.32	0.44
2:DB:44:G:C2	2:DB:48:A:C2	3.06	0.44
2:DB:90:C:OP1	12:DP:16:ARG:HG2	2.17	0.44
3:DD:134:ARG:HB3	3:DD:134:ARG:HE	1.59	0.44
4:DE:87:GLU:O	4:DE:89:ASP:N	2.50	0.44
6:DG:179:PRO:HG3	26:D4:38:LYS:NZ	2.32	0.44
7:DH:4:ILE:O	7:DH:6:ARG:N	2.42	0.44
8:DK:114:LEU:HD22	8:DK:130:TYR:CE1	2.53	0.44
58:DL:102:GLU:HG3	58:DL:103:GLN:N	2.32	0.44
58:DL:111:LYS:HA	58:DL:113:PRO:CG	2.41	0.44
58:DL:54:PRO:HG2	58:DL:55:VAL:N	2.33	0.44
11:DO:113:LYS:HE2	11:DO:115:LEU:HD23	1.98	0.44
11:DO:65:ARG:O	11:DO:66:GLY:C	2.55	0.44
12:DP:58:PHE:O	12:DP:61:GLY:N	2.50	0.44
15:DR:105:LEU:CD2	15:DR:105:LEU:N	2.80	0.44
4:DE:27:LEU:CD2	15:DR:1:MET:HE1	2.47	0.44
15:DR:3:ARG:HA	59:DR:204:MG:MG	1.42	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:DU:56:PRO:O	20:DU:57:GLN:O	2.36	0.44
21:DV:127:LYS:O	21:DV:161:VAL:HG21	2.17	0.44
21:DV:9:TYR:CE2	21:DV:61:LEU:HD21	2.52	0.44
57:DY:98:LYS:HG3	57:DY:99:SER:O	2.18	0.44
13:A0:101:ALA:HB2	27:A5:44:THR:CB	2.47	0.44
13:A0:2:ARG:CG	13:A0:2:ARG:HH11	2.29	0.44
26:A4:34:GLU:HB2	26:A4:35:VAL:H	1.49	0.44
1:AA:2348:U:H5'	28:A6:42:TRP:CD1	2.52	0.44
30:A8:23:VAL:HG13	30:A8:47:LYS:HB3	1.98	0.44
30:A8:56:GLU:O	30:A8:57:ARG:C	2.55	0.44
1:AA:1102:C:H2'	1:AA:1103:A:H5''	2.00	0.44
1:AA:1102:C:H2'	1:AA:1103:A:C5'	2.47	0.44
1:AA:1104:C:O5'	1:AA:1104:C:H6	2.00	0.44
1:AA:111:A:H4'	24:AW:69:ARG:HH22	1.81	0.44
1:AA:532:A:O2'	1:AA:2021:C:N4	2.50	0.44
1:AA:2173:A:H3'	1:AA:2174:C:H6	1.81	0.44
1:AA:2298:A:H2'	1:AA:2299:G:O4'	2.17	0.44
1:AA:2556:C:H2'	1:AA:2557:G:O4'	2.17	0.44
1:AA:2698:U:H2'	1:AA:2699:C:H6	1.80	0.44
1:AA:2727:G:H4'	10:AN:70:LYS:HD2	1.98	0.44
1:AA:414:C:O2'	1:AA:415:A:H5'	2.17	0.44
1:AA:654(H):G:H3'	1:AA:654(I):C:H5''	1.99	0.44
1:AA:704:G:O2'	1:AA:726:G:N2	2.51	0.44
1:AA:846:C:HO2'	1:AA:847:U:C5'	2.30	0.44
1:AA:85:G:H2'	1:AA:86:C:O4'	2.18	0.44
1:AA:912:C:H2'	1:AA:912:C:O2	2.17	0.44
2:AB:40:U:H6	2:AB:40:U:P	2.40	0.44
2:AB:78:A:N6	2:AB:98:G:H1'	2.33	0.44
3:AD:236:GLY:O	3:AD:237:GLU:OE1	2.34	0.44
3:AD:83:GLU:OE1	3:AD:104:TYR:CE2	2.71	0.44
4:AE:51:PHE:CD2	4:AE:52:LEU:HG	2.52	0.44
4:AE:51:PHE:O	4:AE:74:PRO:CB	2.66	0.44
4:AE:7:VAL:HG13	4:AE:51:PHE:HE2	1.82	0.44
5:AF:30:PRO:O	5:AF:33:LEU:N	2.51	0.44
6:AG:7:LEU:H	6:AG:104:GLU:CD	2.18	0.44
8:AK:130:TYR:O	8:AK:136:VAL:CG1	2.65	0.44
9:AM:25:ARG:O	9:AM:28:THR:HB	2.17	0.44
9:AM:46:VAL:HG13	9:AM:47:ALA:H	1.83	0.44
1:AA:662:G:C5'	11:AO:15:ARG:HA	2.36	0.44
11:AO:59:LEU:CD2	11:AO:60:MET:N	2.80	0.44
12:AP:33:GLY:HA2	12:AP:105:GLU:CB	2.48	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:AQ:12:PHE:HA	14:AQ:12:PHE:HD2	1.75	0.44
14:AQ:34:HIS:HB2	14:AQ:36:TYR:CE1	2.52	0.44
18:AS:25:ARG:NH2	18:AS:74:ALA:O	2.48	0.44
18:AS:88:ARG:HB3	18:AS:92:ARG:HB3	2.00	0.44
21:AV:140:ASP:HB3	21:AV:141:VAL:H	1.46	0.44
21:AV:15:PRO:HB2	21:AV:19:ARG:NH2	2.33	0.44
21:AV:30:ASN:HA	21:AV:89:PHE:CE2	2.52	0.44
31:BA:1053:G:H5'	31:BA:1054:C:C5'	2.37	0.44
31:BA:107:G:H2'	31:BA:108:G:H5'	1.98	0.44
31:BA:1297:C:O2'	37:BJ:114:ARG:NH2	2.51	0.44
31:BA:1296:C:H4'	31:BA:1302:U:O4	2.17	0.44
31:BA:1394:A:C5'	31:BA:1395:C:OP2	2.64	0.44
10:AN:49:ARG:HH22	31:BA:1423:G:C5'	2.29	0.44
31:BA:259:G:O2'	31:BA:260:G:H5'	2.17	0.44
31:BA:377:G:OP1	46:BS:3:LYS:HD3	2.17	0.44
31:BA:622:A:C8	31:BA:623:C:C5	3.05	0.44
31:BA:794:A:H2'	31:BA:795:C:O4'	2.17	0.44
31:BA:873:A:H4'	31:BA:874:G:OP1	2.13	0.44
31:BA:922:G:H2'	31:BA:923:A:C8	2.52	0.44
32:BE:17:PHE:HD1	32:BE:42:ILE:HG23	1.82	0.44
32:BE:74:LYS:HE2	32:BE:74:LYS:HB3	1.75	0.44
32:BE:92:TYR:C	32:BE:92:TYR:CD2	2.91	0.44
33:BF:129:ALA:HB1	33:BF:132:ARG:HB3	2.00	0.44
33:BF:91:LEU:C	33:BF:93:LYS:N	2.71	0.44
37:BJ:143:ARG:O	37:BJ:146:GLU:N	2.49	0.44
38:BK:11:THR:O	38:BK:12:ARG:C	2.56	0.44
38:BK:14:ARG:HG2	38:BK:14:ARG:O	2.17	0.44
31:BA:942:G:H21	39:BL:124:GLN:NE2	2.15	0.44
31:BA:963:G:N3	40:BM:55:LYS:NZ	2.65	0.44
40:BM:3:LYS:HB2	40:BM:77:PRO:HD3	2.00	0.44
31:BA:362:G:O3'	42:BO:33:ARG:NH2	2.51	0.44
42:BO:6:THR:OG1	42:BO:9:GLN:HG3	2.18	0.44
45:BR:26:GLU:HG2	45:BR:26:GLU:H	1.54	0.44
45:BR:3:ILE:HG22	45:BR:38:ARG:HH21	1.82	0.44
47:BT:25:ARG:O	47:BT:25:ARG:HG3	2.16	0.44
48:BU:43:PHE:O	48:BU:51:LEU:HG	2.18	0.44
49:BV:52:TYR:HB2	49:BV:57:HIS:CE1	2.53	0.44
49:BV:58:VAL:HG21	49:BV:75:ALA:HA	2.00	0.44
54:CA:1158:C:C2	54:CA:1160:G:C8	3.05	0.44
54:CA:1535:C:H2'	54:CA:1536:C:C5'	2.47	0.44
54:CA:174:C:H2'	54:CA:175:C:C6	2.50	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:192:U:O4'	50:CW:103:GLY:HA2	2.17	0.44
54:CA:445:G:O2'	54:CA:446:G:H5'	2.17	0.44
54:CA:642:A:H2'	54:CA:643:C:H6	1.82	0.44
54:CA:659:U:H2'	54:CA:660:G:H8	1.81	0.44
54:CA:815:A:H4'	54:CA:816:A:OP2	2.17	0.44
33:CF:156:ARG:HB3	33:CF:160:ALA:O	2.17	0.44
33:CF:54:ARG:HH12	33:CF:56:ASP:CG	2.21	0.44
33:CF:78:GLY:O	33:CF:79:ARG:C	2.56	0.44
37:CJ:62:PHE:HA	37:CJ:124:LEU:HD22	1.98	0.44
37:CJ:87:VAL:HG21	37:CJ:154:TYR:CB	2.48	0.44
37:CJ:46:ALA:O	37:CJ:49:ILE:N	2.49	0.44
40:CM:34:VAL:HG13	40:CM:73:ASP:O	2.18	0.44
41:CN:21:ILE:CD1	41:CN:82:VAL:HG13	2.47	0.44
43:CP:122:LYS:HG3	43:CP:123:ALA:N	2.33	0.44
46:CS:45:THR:HG23	46:CS:46:PRO:CD	2.45	0.44
48:CU:25:THR:O	48:CU:25:THR:HG22	2.17	0.44
49:CV:32:LYS:HA	49:CV:50:ALA:HB3	1.99	0.44
49:CV:41:VAL:CG2	49:CV:67:VAL:HG22	2.41	0.44
54:CA:1455:G:OP1	50:CW:35:THR:HG21	2.17	0.44
55:DA:2880:C:H1'	13:D0:92:GLY:O	2.17	0.44
9:DM:41:ASP:C	16:D1:64:ARG:HH22	2.19	0.44
16:D1:8:VAL:HG23	16:D1:11:ARG:NH2	2.32	0.44
17:D2:98:GLU:OE1	17:D2:98:GLU:HA	2.18	0.44
6:DG:67:LYS:CE	26:D4:6:HIS:CE1	2.88	0.44
27:D5:56:LYS:C	27:D5:58:LEU:N	2.71	0.44
30:D8:36:LYS:CB	30:D8:40:GLU:HG2	2.30	0.44
55:DA:1206:G:C6	55:DA:1207:C:C4	3.06	0.44
55:DA:1936:A:H2'	55:DA:1945:G:O6	2.17	0.44
55:DA:2199:A:H3'	55:DA:2205:C:H6	1.82	0.44
55:DA:2213:U:H5'	23:DZ:52:ARG:HH12	1.82	0.44
55:DA:2702:U:O2'	55:DA:2703:C:C6	2.70	0.44
55:DA:581:C:H2'	55:DA:582:G:C8	2.53	0.44
55:DA:784:A:N7	3:DD:229:VAL:CG2	2.81	0.44
2:DB:17:C:O2'	2:DB:18:G:H5'	2.16	0.44
3:DD:45:ASN:OD1	3:DD:45:ASN:C	2.55	0.44
5:DF:165:ARG:HB3	5:DF:165:ARG:HH11	1.83	0.44
7:DH:9:ILE:O	7:DH:10:PRO:O	2.36	0.44
7:DH:103:LEU:CD1	7:DH:131:VAL:HG11	2.48	0.44
56:DI:29:GLU:O	56:DI:30:ALA:HB3	2.17	0.44
56:DJ:8:ILE:CA	56:DJ:11:GLU:HB2	2.48	0.44
58:DL:39:LYS:O	58:DL:41:PHE:N	2.51	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:DL:55:VAL:HG13	58:DL:56:GLU:N	2.33	0.44
11:DO:16:ARG:HE	11:DO:16:ARG:CA	2.30	0.44
14:DQ:35:ILE:CG2	14:DQ:69:VAL:HG11	2.48	0.44
14:DQ:59:LYS:CG	14:DQ:60:GLY:N	2.71	0.44
18:DS:17:VAL:O	18:DS:20:VAL:HG22	2.18	0.44
20:DU:47:LYS:C	20:DU:49:VAL:N	2.69	0.44
21:DV:6:LYS:CE	21:DV:43:GLU:HG3	2.48	0.44
57:DY:102:LYS:HD2	57:DY:103:GLY:N	2.32	0.44
57:DY:134:LEU:HD21	56:DJ:19:GLU:CG	2.46	0.44
57:DY:6:ASN:O	57:DY:7:VAL:HG12	2.18	0.44
17:A2:18:LEU:HD23	17:A2:19:LYS:C	2.38	0.44
17:A2:59:ALA:HB1	17:A2:94:LEU:HB3	2.00	0.44
22:A3:11:ARG:O	22:A3:14:ARG:NH2	2.50	0.44
30:A8:61:LEU:HD13	30:A8:62:LEU:H	1.82	0.44
1:AA:1157:G:H2'	1:AA:1158:C:C6	2.53	0.44
1:AA:1278:A:H5''	13:A0:36:THR:CG2	2.46	0.44
1:AA:1285:G:N2	1:AA:1328:G:H5''	2.33	0.44
1:AA:1781:C:H2'	1:AA:1781:C:O2	2.16	0.44
1:AA:526:A:O2'	1:AA:2043:C:O2	2.33	0.44
1:AA:2131:G:C5'	1:AA:2132:U:H5''	2.33	0.44
1:AA:2159:G:H2'	1:AA:2160:G:O4'	2.17	0.44
1:AA:2302:G:C6	1:AA:2315:G:C6	3.06	0.44
1:AA:2405:G:HO2'	1:AA:2406:U:P	2.39	0.44
1:AA:957:A:H2	1:AA:2458:G:H4'	1.82	0.44
1:AA:2643:G:H2'	1:AA:2644:G:O4'	2.18	0.44
1:AA:2656:U:O4	1:AA:2665:A:C6	2.70	0.44
1:AA:2822:G:OP2	4:AE:110:GLY:O	2.36	0.44
1:AA:2840:C:H4'	13:A0:53:HIS:HD2	1.82	0.44
1:AA:2854:G:N1	1:AA:2864:G:C6	2.86	0.44
1:AA:558:G:C5'	9:AM:112:LEU:HD22	2.48	0.44
1:AA:582:G:H2'	1:AA:583:G:C8	2.52	0.44
1:AA:654(C):G:C2'	1:AA:654(D):G:C8	3.00	0.44
1:AA:72:U:H1'	24:AW:58:ALA:HA	1.99	0.44
1:AA:741:G:O2'	1:AA:742:G:H5'	2.18	0.44
1:AA:7:G:O2'	1:AA:8:A:H5'	2.18	0.44
1:AA:912:C:O2'	1:AA:913:U:H5'	2.18	0.44
1:AA:991:C:H2'	1:AA:992:C:H6	1.82	0.44
2:AB:15:A:O2'	2:AB:109:G:C8	2.58	0.44
2:AB:20:C:O2'	2:AB:21:G:H5'	2.18	0.44
3:AD:127:VAL:HG12	3:AD:127:VAL:O	2.17	0.44
3:AD:48:ARG:HH11	3:AD:48:ARG:HG3	1.82	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:AE:175:VAL:O	4:AE:177:PRO:CD	2.64	0.44
4:AE:42:ASP:HB3	4:AE:44:TYR:CE1	2.52	0.44
5:AF:114:VAL:HG21	5:AF:202:PHE:CZ	2.53	0.44
6:AG:109:VAL:HG11	6:AG:142:PRO:HB3	1.98	0.44
6:AG:41:GLN:HG2	6:AG:155:MET:HB3	1.98	0.44
8:AK:9:LEU:HD23	8:AK:9:LEU:H	1.82	0.44
1:AA:636:G:C4	11:AO:115:LEU:HD11	2.53	0.44
12:AP:35:VAL:HG12	12:AP:130:LYS:O	2.17	0.44
15:AR:26:ASP:HB3	15:AR:91:ARG:HA	1.98	0.44
15:AR:74:ARG:CB	15:AR:74:ARG:HH11	2.30	0.44
15:AR:82:LEU:H	15:AR:82:LEU:HD12	1.80	0.44
15:AR:90:GLN:HG3	15:AR:91:ARG:N	2.33	0.44
15:AR:94:ALA:O	15:AR:96:ARG:N	2.51	0.44
20:AU:97:ARG:CD	20:AU:97:ARG:H	2.22	0.44
23:AZ:65:SER:O	23:AZ:66:HIS:HD2	1.99	0.44
53:B1:54:U:H2'	53:B1:55:U:OP2	2.18	0.44
31:BA:1041:A:H2'	31:BA:1042:G:H5''	1.98	0.44
31:BA:1207:G:C5	31:BA:1208:C:C5	3.06	0.44
31:BA:1219:U:OP1	44:BQ:19:ARG:NH1	2.46	0.44
1:AA:1916:A:N6	31:BA:1408:A:O2'	2.49	0.44
31:BA:399:G:H2'	31:BA:400:C:H6	1.77	0.44
31:BA:48:C:C5'	31:BA:49:U:OP2	2.64	0.44
31:BA:502:G:H2'	31:BA:503:C:O4'	2.17	0.44
31:BA:646:U:H2'	31:BA:647:C:C6	2.49	0.44
31:BA:738:C:H2'	31:BA:739:C:C6	2.52	0.44
31:BA:794:A:H2	31:BA:795:C:N3	2.09	0.44
1:AA:2554:U:H3	52:BB:74:C:H5	1.65	0.44
33:BF:137:ALA:O	33:BF:141:VAL:HG23	2.17	0.44
33:BF:41:GLY:O	33:BF:45:LYS:HG3	2.17	0.44
34:BG:73:ARG:O	34:BG:77:ASN:ND2	2.50	0.44
36:BI:7:ASN:O	36:BI:88:VAL:HA	2.17	0.44
37:BJ:113:GLU:HG3	37:BJ:119:ARG:HA	1.98	0.44
39:BL:41:VAL:O	39:BL:41:VAL:HG12	2.18	0.44
41:BN:59:TYR:CE1	41:BN:63:LEU:HD21	2.52	0.44
43:BP:66:LEU:O	43:BP:69:GLU:HB3	2.18	0.44
46:BS:45:THR:HB	46:BS:46:PRO:HD2	1.98	0.44
48:BU:70:ILE:HG23	48:BU:79:LEU:CD1	2.48	0.44
49:BV:47:HIS:O	49:BV:48:THR:CB	2.66	0.44
54:CA:1005:A:H5''	54:CA:1006:C:C5	2.52	0.44
54:CA:1006:C:C2	54:CA:1007:C:C5	3.06	0.44
54:CA:1124:G:N2	54:CA:1280:A:N6	2.66	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1225:A:N3	54:CA:1225:A:C2'	2.77	0.44
54:CA:1258:G:O2'	54:CA:1259:C:H5'	2.17	0.44
54:CA:1362(A):C:H5'	54:CA:1363:A:O5'	2.17	0.44
54:CA:177:C:OP1	50:CW:65:LYS:NZ	2.47	0.44
54:CA:751:U:H2'	54:CA:752:G:O4'	2.17	0.44
54:CA:954:G:H2'	54:CA:955:U:C6	2.53	0.44
32:CE:200:ILE:HG22	32:CE:202:PRO:HD3	1.99	0.44
32:CE:237:ALA:C	32:CE:239:VAL:H	2.20	0.44
33:CF:108:ASN:OD1	33:CF:110:ASN:HB2	2.18	0.44
33:CF:50:ALA:HB2	33:CF:75:VAL:HB	2.00	0.44
39:CL:122:ALA:HB1	39:CL:123:PRO:HD2	1.98	0.44
39:CL:66:ARG:HH11	39:CL:66:ARG:CB	2.30	0.44
39:CL:10:ARG:HD2	39:CL:75:ASP:HB3	1.99	0.44
41:CN:120:ARG:NH1	41:CN:126:ARG:NH2	2.65	0.44
44:CQ:13:THR:HG23	44:CQ:20:ALA:HB2	2.00	0.44
47:CT:3:LYS:HD3	47:CT:61:GLU:O	2.18	0.44
55:DA:2723:C:O3'	13:D0:1:MET:HE2	2.16	0.44
13:D0:2:ARG:HG3	13:D0:2:ARG:HH11	1.83	0.44
55:DA:2355:C:O3'	22:D3:24:LYS:HD2	2.17	0.44
28:D6:43:CYS:O	28:D6:44:ARG:HD3	2.17	0.44
55:DA:1062:G:H1'	55:DA:1088:A:C5	2.51	0.44
55:DA:1116:C:H2'	55:DA:1117:G:H8	1.81	0.44
55:DA:1173:G:C5'	55:DA:1174:A:OP1	2.56	0.44
55:DA:1203:G:H5'	11:DO:3:LEU:CD1	2.46	0.44
55:DA:1203:G:O6	55:DA:1204:A:N6	2.50	0.44
55:DA:1310:G:H2'	55:DA:1311:G:O4'	2.18	0.44
55:DA:1538:G:C2	55:DA:1539:G:C8	3.05	0.44
55:DA:1130:U:C2	55:DA:2025:C:H5''	2.52	0.44
55:DA:2158:A:O2'	55:DA:2159:G:C8	2.70	0.44
55:DA:221:A:O2'	55:DA:222:A:OP2	2.34	0.44
55:DA:227:A:OP2	55:DA:227:A:H8	2.01	0.44
55:DA:2581:G:H4'	55:DA:2582:G:C8	2.53	0.44
55:DA:1786:A:H2	55:DA:2606:C:H1'	1.79	0.44
55:DA:2865:U:C4	55:DA:2866:U:C4	3.04	0.44
55:DA:289:A:H2'	55:DA:290:G:O4'	2.18	0.44
55:DA:347:A:H2'	55:DA:348:G:H8	1.83	0.44
55:DA:84:A:H5''	20:DU:8:LYS:HG2	2.00	0.44
2:DB:11:C:H3'	2:DB:12:C:C6	2.53	0.44
2:DB:74:U:C3'	2:DB:75:G:C5'	2.95	0.44
2:DB:96:G:N2	2:DB:97:G:H1'	2.33	0.44
5:DF:88:VAL:CG1	5:DF:91:GLY:HA3	2.47	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:DH:23:ARG:HD2	7:DH:34:GLU:OE2	2.17	0.44
58:DL:146:ASP:HB2	58:DL:147:ALA:H	1.58	0.44
9:DM:65:LYS:HB2	9:DM:69:GLN:NE2	2.19	0.44
10:DN:32:TYR:CD1	10:DN:32:TYR:N	2.85	0.44
11:DO:16:ARG:H	11:DO:16:ARG:HE	1.65	0.44
20:DU:46:LYS:NZ	20:DU:63:LYS:HG2	2.33	0.44
20:DU:84:ARG:HH12	20:DU:97:ARG:HB3	1.80	0.44
57:DY:112:LEU:O	57:DY:113:GLN:CG	2.65	0.44
57:DY:143:GLN:HB3	57:DY:144:ALA:H	1.34	0.44
57:DY:93:LEU:HA	57:DY:93:LEU:HD22	1.79	0.44
13:A0:45:ARG:HH11	13:A0:45:ARG:HG3	1.83	0.44
13:A0:54:LEU:HD12	13:A0:54:LEU:HA	1.82	0.44
1:AA:1242:A:H3'	1:AA:1243:G:H8	1.82	0.44
1:AA:1536:A:P	1:AA:1537:C:N4	2.91	0.44
1:AA:1827:C:H3'	1:AA:1828:G:H8	1.82	0.44
1:AA:570:G:H2'	1:AA:2030:A:N7	2.33	0.44
1:AA:2362:G:C2'	1:AA:2363:C:H5'	2.47	0.44
1:AA:2364:C:H2'	1:AA:2365:G:O4'	2.17	0.44
1:AA:2449:U:O2'	1:AA:2450:A:C8	2.68	0.44
1:AA:2491:U:H5'	1:AA:2570:G:H5''	1.98	0.44
1:AA:1966:A:C2	1:AA:2593:U:O4'	2.70	0.44
1:AA:404:C:C1'	1:AA:406:G:C8	2.99	0.44
1:AA:604:G:O2'	1:AA:605:C:H5'	2.17	0.44
1:AA:747:U:C5	1:AA:2613:U:C5	3.06	0.44
2:AB:104:A:H5'	21:AV:72:ARG:HD3	2.00	0.44
2:AB:66:A:O2'	2:AB:67:G:OP2	2.32	0.44
4:AE:137:HIS:CB	4:AE:138:PRO:HD2	2.43	0.44
1:AA:2050:C:H1'	4:AE:156:MET:HE1	1.99	0.44
4:AE:62:PRO:C	4:AE:64:LYS:H	2.20	0.44
6:AG:51:ARG:HH11	6:AG:51:ARG:CB	2.30	0.44
6:AG:70:VAL:HG12	6:AG:90:LEU:HD11	1.99	0.44
6:AG:9:ARG:HG2	6:AG:13:GLU:OE1	2.18	0.44
7:AH:4:ILE:HD11	7:AH:7:LEU:HD23	1.99	0.44
8:AK:102:SER:OG	8:AK:103:ARG:N	2.51	0.44
8:AK:2:LYS:HB2	8:AK:39:ALA:HB3	1.99	0.44
12:AP:111:GLU:OE1	12:AP:133:ARG:NH2	2.51	0.44
19:AT:34:ALA:HA	19:AT:38:GLU:OE2	2.18	0.44
20:AU:81:LYS:HB2	20:AU:96:ILE:HG22	1.99	0.44
21:AV:175:VAL:HG22	21:AV:176:PRO:HD2	2.00	0.44
21:AV:41:LEU:O	21:AV:42:VAL:C	2.55	0.44
53:B1:29:G:H2'	53:B1:30:C:C6	2.53	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:1054:C:OP2	31:BA:1197:G:OP2	2.36	0.44
31:BA:1326:C:C2	31:BA:1327:C:C5	3.06	0.44
31:BA:1347:G:C2'	31:BA:1348:U:OP2	2.65	0.44
1:AA:1915:U:O4	31:BA:1409:C:C5'	2.66	0.44
31:BA:614:A:H2'	31:BA:615:C:O4'	2.17	0.44
31:BA:790:A:H5'	52:BC:38:A:O3'	2.17	0.44
31:BA:6:G:O2'	31:BA:7:G:O5'	2.31	0.44
31:BA:976:G:P	44:BQ:32:SER:H	2.40	0.44
33:BF:110:ASN:HD22	33:BF:144:SER:HB3	1.82	0.44
33:BF:113:ALA:HB2	33:BF:202:ILE:CG1	2.48	0.44
33:BF:114:PRO:O	33:BF:118:GLN:HG3	2.17	0.44
33:BF:73:PRO:HB2	33:BF:76:VAL:CG2	2.47	0.44
36:BI:15:ASP:C	36:BI:15:ASP:OD1	2.55	0.44
36:BI:1:MET:HE1	36:BI:68:PRO:HD3	2.00	0.44
38:BK:124:ALA:O	38:BK:128:GLY:N	2.50	0.44
38:BK:95:VAL:HG11	38:BK:133:LEU:HD12	2.00	0.44
39:BL:114:TYR:N	39:BL:114:TYR:CD2	2.85	0.44
39:BL:11:LYS:H	39:BL:104:ARG:HH21	1.63	0.44
40:BM:22:LYS:NZ	40:BM:23:ILE:HG12	2.33	0.44
33:BF:18:TRP:CZ2	44:BQ:56:VAL:O	2.70	0.44
46:BS:51:VAL:HG11	46:BS:74:LEU:HD23	1.98	0.44
47:BT:52:LYS:N	47:BT:52:LYS:HD2	2.33	0.44
48:BU:36:ASN:HD22	48:BU:39:VAL:CG2	2.30	0.44
50:BW:43:LEU:HB3	50:BW:48:LYS:HG3	1.99	0.44
54:CA:1200:C:C2'	54:CA:1201:A:OP2	2.65	0.44
54:CA:960:U:N3	54:CA:1225:A:C5	2.85	0.44
54:CA:1256:A:C2	54:CA:1277:C:H3'	2.53	0.44
54:CA:335:C:H2'	54:CA:336:C:H6	1.83	0.44
54:CA:509:A:H1'	54:CA:543:C:O2'	2.18	0.44
54:CA:700:G:C2'	54:CA:701:C:H5''	2.47	0.44
54:CA:89:U:O2'	54:CA:90:C:O5'	2.33	0.44
52:CB:10:G:H3'	52:CB:11:C:C5	2.51	0.44
52:CC:43:C:H2'	52:CC:44:G:O4'	2.17	0.44
52:CD:18:G:O2'	52:CD:19:G:OP1	2.31	0.44
52:CD:20:U:OP1	52:CD:20:U:H6	2.00	0.44
52:CD:37:MIA:H162	53:C1:43:U:O4	2.17	0.44
32:CE:14:GLY:O	32:CE:15:VAL:HG22	2.18	0.44
32:CE:207:ALA:O	32:CE:208:ILE:C	2.55	0.44
33:CF:12:LEU:C	33:CF:14:ILE:H	2.20	0.44
40:CM:5:ARG:O	40:CM:6:ILE:HG23	2.18	0.44
40:CM:90:LEU:N	40:CM:90:LEU:HD12	2.32	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
43:CP:45:VAL:O	43:CP:48:LEU:HD22	2.17	0.44
43:CP:82:MET:HE2	43:CP:93:ARG:N	2.32	0.44
46:CS:6:LEU:HD11	46:CS:19:ILE:CD1	2.47	0.44
47:CT:76:LEU:HD11	47:CT:79:SER:H	1.83	0.44
48:CU:26:LEU:HD21	48:CU:42:ARG:HD2	1.99	0.44
49:CV:11:VAL:CG1	49:CV:16:LEU:HD22	2.48	0.44
26:D4:54:GLY:O	26:D4:55:ARG:O	2.35	0.44
28:D6:31:PRO:O	28:D6:32:ASN:CB	2.65	0.44
28:D6:41:PRO:HG2	28:D6:45:LYS:H	1.82	0.44
55:DA:1059:G:OP1	58:DL:4:VAL:HG13	2.16	0.44
55:DA:1142(A):A:C4	55:DA:1144:G:N7	2.86	0.44
55:DA:1188:U:C2'	55:DA:1189:A:C5'	2.96	0.44
55:DA:1204:A:C2'	55:DA:1205:U:OP2	2.66	0.44
55:DA:1204:A:H1'	55:DA:1206:G:N9	2.33	0.44
55:DA:1523:U:H2'	55:DA:1524:G:C8	2.53	0.44
55:DA:1556:C:H2'	55:DA:1557:C:C6	2.52	0.44
55:DA:1694:C:O2'	55:DA:1695:G:P	2.76	0.44
55:DA:1811:G:H2'	55:DA:1812:A:O4'	2.17	0.44
55:DA:1800:C:N4	55:DA:1819:A:N7	2.51	0.44
55:DA:1850:G:H2'	55:DA:1851:U:O4'	2.16	0.44
55:DA:2205:C:H5''	55:DA:2205:C:C6	2.52	0.44
55:DA:2065:C:H1'	55:DA:2449:U:O2	2.17	0.44
55:DA:2060:A:H1'	55:DA:2502:G:O4'	2.18	0.44
55:DA:270(Z):U:O2'	55:DA:271(A):C:OP2	2.26	0.44
55:DA:2810:A:H61	55:DA:2891:G:C2'	2.30	0.44
55:DA:303:U:O2'	55:DA:304:G:H5'	2.17	0.44
55:DA:578:A:H5'	55:DA:1254:A:OP1	2.17	0.44
55:DA:894:C:C6	55:DA:894:C:O5'	2.68	0.44
55:DA:860:U:C4	55:DA:917:A:H2	2.35	0.44
6:DG:25:TYR:CE2	6:DG:31:VAL:HA	2.53	0.44
8:DK:114:LEU:HD22	8:DK:130:TYR:HD1	1.79	0.44
58:DL:48:MET:O	58:DL:49:GLY:O	2.35	0.44
58:DL:14:ALA:HA	58:DL:50:ASP:HB3	2.00	0.44
11:DO:16:ARG:HE	11:DO:16:ARG:N	2.16	0.44
11:DO:83:VAL:CG1	11:DO:112:LEU:CD2	2.93	0.44
12:DP:54:MET:HE1	12:DP:104:PHE:HB3	1.99	0.44
15:DR:41:ARG:NH2	15:DR:43:GLN:CB	2.81	0.44
21:DV:178:GLU:O	21:DV:180:VAL:N	2.39	0.44
21:DV:178:GLU:OE1	21:DV:181:GLU:N	2.50	0.44
21:DV:196:VAL:CA	21:DV:197:ILE:HD12	2.48	0.44
21:DV:69:THR:HG22	21:DV:90:VAL:CA	2.43	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:DX:38:GLU:O	25:DX:40:THR:N	2.51	0.44
57:DY:1:MET:O	57:DY:1:MET:CG	2.64	0.44
57:DY:49:ALA:HA	57:DY:84:GLU:N	2.32	0.44
13:A0:52:ILE:O	13:A0:55:ALA:N	2.51	0.44
1:AA:1161:C:O2'	17:A2:23:GLU:HG2	2.17	0.44
22:A3:43:THR:CG2	22:A3:46:LYS:HE2	2.48	0.44
1:AA:1042:G:H2'	1:AA:1043:C:C6	2.53	0.44
1:AA:1125:G:C6	1:AA:1126:A:N6	2.85	0.44
1:AA:12:U:O2	1:AA:12:U:H2'	2.16	0.44
1:AA:1378:A:O2'	1:AA:1379:A:OP1	2.29	0.44
1:AA:1428:C:N4	1:AA:1569:A:H3'	2.31	0.44
1:AA:1512:G:H2'	1:AA:1513:C:C6	2.52	0.44
1:AA:1467:C:C5	1:AA:1546:C:H2'	2.52	0.44
1:AA:1683:C:C2	1:AA:1684:C:C5	3.06	0.44
1:AA:1762:A:C4'	1:AA:1763:G:OP2	2.65	0.44
1:AA:184:C:H2'	1:AA:185:U:C6	2.53	0.44
1:AA:204:A:O2'	1:AA:205:G:P	2.76	0.44
1:AA:412:A:N7	1:AA:2411:A:H2	2.16	0.44
1:AA:2447:G:H3'	1:AA:2500:U:OP2	2.18	0.44
1:AA:2483:C:N3	12:AP:124:LYS:NZ	2.65	0.44
1:AA:2491:U:O2'	1:AA:2492:U:H5'	2.18	0.44
1:AA:2496:C:OP1	12:AP:81:VAL:CG1	2.63	0.44
1:AA:2720:U:C2	1:AA:2873:A:H2	2.36	0.44
1:AA:2866:U:O2	1:AA:2866:U:C2'	2.61	0.44
1:AA:751:A:H5'	18:AS:90:ARG:HA	1.99	0.44
1:AA:917:A:H2'	1:AA:918:A:O5'	2.17	0.44
1:AA:91:A:O2'	1:AA:92:G:H5'	2.18	0.44
3:AD:159:ALA:N	3:AD:196:VAL:HG11	2.26	0.44
4:AE:103:ASP:N	4:AE:200:GLU:O	2.50	0.44
6:AG:9:ARG:C	6:AG:11:TYR:H	2.21	0.44
8:AK:37:VAL:CG1	8:AK:38:LEU:N	2.81	0.44
9:AM:19:GLU:HB3	9:AM:59:LYS:HE3	1.99	0.44
11:AO:79:ARG:HG3	11:AO:109:GLY:O	2.17	0.44
11:AO:21:ARG:O	11:AO:28:GLY:HA2	2.18	0.44
12:AP:22:LYS:O	12:AP:23:GLY:C	2.56	0.44
14:AQ:48:LEU:HD22	14:AQ:82:ILE:HD11	2.00	0.44
20:AU:13:VAL:HG21	20:AU:72:VAL:HB	2.00	0.44
20:AU:81:LYS:HB3	20:AU:97:ARG:HD2	1.97	0.44
31:BA:1053:G:C4'	31:BA:1054:C:H5'	2.48	0.44
31:BA:1128:C:H2'	31:BA:1130:A:C8	2.53	0.44
31:BA:1235:U:O2'	31:BA:1305:G:O5'	2.35	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:1346:A:H5'	39:BL:120:ARG:HH12	1.82	0.44
31:BA:1535:C:H2'	31:BA:1536:C:O4'	2.17	0.44
52:BD:57:G:C2'	52:BD:58:A:H5''	2.47	0.44
34:BG:190:ASP:O	34:BG:194:LEU:HD23	2.18	0.44
35:BH:90:VAL:O	35:BH:120:THR:HA	2.18	0.44
36:BI:45:LEU:HD21	36:BI:57:GLN:CD	2.38	0.44
31:BA:1346:A:N6	37:BJ:10:ARG:NE	2.65	0.44
37:BJ:111:ARG:HH12	37:BJ:122:HIS:HB3	1.83	0.44
40:BM:47:PHE:CD1	40:BM:47:PHE:O	2.71	0.44
40:BM:69:ASN:O	40:BM:70:ARG:HB2	2.18	0.44
46:BS:48:TRP:CZ3	46:BS:49:LEU:HB2	2.52	0.44
50:BW:67:ALA:O	50:BW:73:HIS:CD2	2.70	0.44
50:BW:82:SER:O	50:BW:83:ARG:C	2.56	0.44
54:CA:1016:A:H2'	54:CA:1017:G:O4'	2.18	0.44
54:CA:1028:C:C3'	54:CA:1028(A):C:H5''	2.47	0.44
54:CA:1116:C:C2'	54:CA:1117:G:C5'	2.90	0.44
54:CA:1280:A:HO2'	54:CA:1281:U:P	2.41	0.44
54:CA:1363:A:N3	54:CA:1365:G:C6	2.86	0.44
54:CA:229:U:O2'	54:CA:230:G:H5'	2.18	0.44
54:CA:33:A:C4	54:CA:34:C:C5	3.06	0.44
54:CA:428:G:H4'	54:CA:429:U:O5'	2.18	0.44
54:CA:509:A:C8	54:CA:509:A:H3'	2.52	0.44
54:CA:601:C:O2'	54:CA:602:A:H5'	2.18	0.44
54:CA:645:C:H2'	54:CA:646:U:H6	1.81	0.44
54:CA:738:C:C2	54:CA:739:C:C5	3.05	0.44
54:CA:873:A:H4'	54:CA:874:G:OP1	2.16	0.44
32:CE:126:GLU:O	32:CE:126:GLU:HG2	2.17	0.44
32:CE:118:LEU:HB3	32:CE:142:LEU:HD12	1.98	0.44
34:CG:83:SER:C	34:CG:85:LYS:H	2.21	0.44
38:CK:10:LEU:HD23	38:CK:10:LEU:N	2.33	0.44
38:CK:129:VAL:HG23	38:CK:130:GLY:N	2.29	0.44
40:CM:95:GLU:HG3	40:CM:96:ILE:N	2.33	0.44
41:CN:54:ARG:O	41:CN:57:THR:HB	2.17	0.44
43:CP:44:ARG:C	43:CP:46:LYS:H	2.21	0.44
43:CP:25:ILE:HG13	43:CP:66:LEU:HD23	2.00	0.44
43:CP:82:MET:CE	43:CP:93:ARG:HA	2.45	0.44
46:CS:80:PHE:CD1	46:CS:80:PHE:N	2.86	0.44
47:CT:29:HIS:CE1	47:CT:32:TYR:HD1	2.36	0.44
13:D0:20:LEU:O	13:D0:21:TYR:C	2.55	0.44
13:D0:38:VAL:HG12	13:D0:42:LYS:HE3	1.99	0.44
55:DA:1653:G:C6	13:D0:9:LYS:O	2.66	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:D1:90:VAL:C	16:D1:92:ARG:N	2.68	0.44
17:D2:41:GLY:CA	17:D2:46:VAL:CG1	2.96	0.44
17:D2:62:LEU:HD12	17:D2:62:LEU:N	2.32	0.44
17:D2:66:ARG:CG	17:D2:66:ARG:HH11	2.30	0.44
17:D2:69:LYS:HA	17:D2:87:HIS:O	2.17	0.44
29:D7:47:ARG:HB2	29:D7:48:LYS:H	1.55	0.44
30:D8:22:VAL:HB	30:D8:53:PRO:HB2	1.99	0.44
55:DA:1083:U:C1'	55:DA:1086:A:H61	2.03	0.44
55:DA:1175:U:HO2'	55:DA:1176:G:H4'	1.79	0.44
55:DA:1892:C:O2'	55:DA:1893:C:H5'	2.18	0.44
55:DA:2467:C:H2'	55:DA:2468:G:O4'	2.17	0.44
55:DA:341:G:H2'	55:DA:342:G:O4'	2.17	0.44
55:DA:540:G:H3'	55:DA:541:C:H6	1.83	0.44
55:DA:552:G:H2'	55:DA:553:U:O4'	2.18	0.44
55:DA:627:A:H4'	55:DA:628:G:OP1	2.16	0.44
55:DA:637:A:OP1	11:DO:133:SER:HB3	2.18	0.44
55:DA:70:G:N2	55:DA:71:A:H62	2.13	0.44
55:DA:751:A:H5'	18:DS:90:ARG:HA	1.99	0.44
55:DA:916:G:H2'	55:DA:917:A:H5''	1.99	0.44
3:DD:52:ARG:H	3:DD:52:ARG:HG3	1.61	0.44
4:DE:7:VAL:HG22	4:DE:27:LEU:CD2	2.42	0.44
4:DE:91:VAL:HG13	4:DE:95:ILE:HG12	1.99	0.44
7:DH:16:SER:OG	7:DH:17:VAL:N	2.51	0.44
8:DK:101:LEU:CD2	8:DK:107:VAL:HB	2.42	0.44
55:DA:1080:A:O2'	58:DL:126:MET:CE	2.66	0.44
58:DL:32:ALA:O	58:DL:64:SER:OG	2.34	0.44
9:DM:19:GLU:HG3	9:DM:59:LYS:HB3	1.99	0.44
12:DP:41:TRP:HB3	12:DP:94:VAL:HB	2.00	0.44
12:DP:78:PRO:C	12:DP:79:LEU:HG	2.38	0.44
2:DB:50:G:P	14:DQ:63:THR:HG23	2.56	0.44
14:DQ:78:LEU:HD11	14:DQ:107:GLU:O	2.18	0.44
18:DS:34:ASN:O	18:DS:37:ARG:HB3	2.17	0.44
21:DV:112:ARG:CD	21:DV:112:ARG:N	2.70	0.44
21:DV:105:VAL:HG12	21:DV:140:ASP:CA	2.48	0.44
57:DY:57:THR:O	57:DY:58:LEU:C	2.56	0.44
57:DY:6:ASN:O	57:DY:7:VAL:CG1	2.65	0.44
13:A0:97:VAL:HG13	13:A0:114:VAL:CG2	2.47	0.44
1:AA:1058:U:H2'	1:AA:1059:G:H8	1.83	0.44
1:AA:1060:U:C1'	1:AA:1062:G:H5'	2.47	0.44
1:AA:1005:C:C1'	1:AA:1143:A:C2	3.01	0.44
1:AA:1342:A:O2'	1:AA:1343:G:O5'	2.36	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1434:A:O2'	1:AA:1435:G:H5'	2.18	0.44
1:AA:1665:A:O2'	1:AA:1666:G:H5'	2.18	0.44
1:AA:1695:G:H2'	1:AA:1696:G:C5'	2.48	0.44
1:AA:1762:A:C5'	1:AA:1763:G:OP2	2.65	0.44
1:AA:2353:G:H2'	1:AA:2354:G:O4'	2.17	0.44
1:AA:2713:A:H3'	1:AA:2714:G:H5''	1.99	0.44
1:AA:305:U:H2'	1:AA:306:U:C6	2.53	0.44
1:AA:312:G:P	1:AA:312:G:C8	3.11	0.44
1:AA:469:G:H2'	1:AA:470:A:H5''	1.98	0.44
1:AA:481:G:H2'	1:AA:507:A:N1	2.33	0.44
1:AA:548:A:C2'	1:AA:549:G:H5'	2.47	0.44
1:AA:616:A:O2'	1:AA:617:G:C5'	2.66	0.44
1:AA:688:U:H5'	1:AA:1780:A:C2	2.53	0.44
1:AA:795:C:H2'	1:AA:796:C:C6	2.52	0.44
1:AA:968:G:H2'	1:AA:969:U:C6	2.52	0.44
4:AE:12:THR:H	4:AE:23:VAL:HG23	1.83	0.44
12:AP:66:ILE:HG13	12:AP:67:ARG:N	2.31	0.44
15:AR:98:LYS:HD2	15:AR:98:LYS:H	1.82	0.44
21:AV:141:VAL:CG2	21:AV:144:LEU:CD2	2.90	0.44
21:AV:52:SER:O	21:AV:53:ILE:HG13	2.18	0.44
24:AW:32:LEU:HA	24:AW:35:LEU:HD23	1.98	0.44
25:AX:36:VAL:HG23	25:AX:36:VAL:O	2.17	0.44
31:BA:1211:U:O2'	31:BA:1213:A:N3	2.51	0.44
31:BA:1300:G:O2'	31:BA:1301:U:O5'	2.25	0.44
31:BA:130:A:H1'	31:BA:264:U:H5'	1.99	0.44
31:BA:251:G:C2	31:BA:266:G:C6	3.06	0.44
31:BA:464:G:O6	31:BA:466:C:H5'	2.18	0.44
31:BA:568:G:N3	31:BA:574:A:H2	2.16	0.44
31:BA:666:G:C2	31:BA:741:G:C4	3.06	0.44
31:BA:748:C:H1'	31:BA:749:C:H5	1.83	0.44
31:BA:750:G:H1'	45:BR:23:GLY:H	1.81	0.44
31:BA:752:G:H1'	31:BA:754:C:N4	2.29	0.44
31:BA:89:U:O2'	31:BA:90:C:P	2.76	0.44
31:BA:950:U:H3'	43:BP:102:ARG:HH22	1.83	0.44
33:BF:136:GLN:O	33:BF:137:ALA:C	2.56	0.44
33:BF:141:VAL:HG11	33:BF:202:ILE:CD1	2.47	0.44
34:BG:146:ILE:H	34:BG:146:ILE:CD1	2.28	0.44
35:BH:147:ASP:HA	35:BH:150:ARG:CB	2.47	0.44
44:BQ:15:LYS:HA	44:BQ:15:LYS:CE	2.48	0.44
48:BU:43:PHE:CA	48:BU:51:LEU:HD12	2.44	0.44
49:BV:49:ILE:HG22	49:BV:50:ALA:N	2.32	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1002:G:N3	54:CA:1003:G:N7	2.66	0.44
54:CA:107:G:O2'	54:CA:108:G:H5'	2.18	0.44
54:CA:1174:G:H2'	54:CA:1175:G:C8	2.52	0.44
54:CA:960:U:N3	54:CA:1225:A:C8	2.76	0.44
54:CA:1239:A:H4'	54:CA:1240:U:O5'	2.18	0.44
54:CA:945:G:C6	54:CA:1337:G:C5	3.05	0.44
54:CA:1356:G:H2'	54:CA:1357:A:C8	2.52	0.44
54:CA:109:A:C6	54:CA:326:G:C6	3.06	0.44
54:CA:372:C:HO2'	54:CA:373:A:P	2.41	0.44
54:CA:487:A:H3'	54:CA:488:C:H6	1.83	0.44
54:CA:545:C:OP1	34:CG:61:LYS:NZ	2.50	0.44
54:CA:66:G:H2'	54:CA:66:G:N3	2.33	0.44
54:CA:701:C:H1'	54:CA:703:G:N3	2.31	0.44
54:CA:954:G:H2'	54:CA:955:U:H6	1.83	0.44
52:CD:27:G:H2'	52:CD:28:G:H8	1.83	0.44
37:CJ:26:PHE:HD1	37:CJ:101:LEU:HD22	1.83	0.44
42:CO:26:ALA:C	42:CO:27:LEU:O	2.56	0.44
54:CA:659:U:OP1	45:CR:9:GLN:NE2	2.51	0.44
46:CS:83:GLU:HG3	46:CS:84:ALA:N	2.32	0.44
47:CT:46:ASP:OD2	47:CT:51:TYR:CD1	2.71	0.44
48:CU:22:VAL:O	48:CU:25:THR:CB	2.66	0.44
48:CU:66:LEU:HG	48:CU:70:ILE:HD11	1.99	0.44
49:CV:53:ASN:HD22	49:CV:58:VAL:HG13	1.83	0.44
49:CV:63:THR:CG2	49:CV:66:MET:HE3	2.47	0.44
50:CW:14:LYS:HA	50:CW:17:ARG:NH1	2.32	0.44
27:D5:31:VAL:CG1	27:D5:42:PRO:HG3	2.47	0.44
55:DA:1014:U:H2'	55:DA:1015:G:H8	1.83	0.44
55:DA:1059:G:H21	58:DL:126:MET:HB3	1.82	0.44
55:DA:1077:A:H4'	58:DL:93:ARG:HH21	1.80	0.44
55:DA:1057:A:C8	55:DA:1086:A:H2'	2.38	0.44
55:DA:1204:A:H2'	55:DA:1205:U:OP2	2.18	0.44
55:DA:1230:C:H2'	55:DA:1231:G:H8	1.83	0.44
55:DA:1578:U:H2'	55:DA:1578:U:O2	2.18	0.44
55:DA:1668:A:H4'	55:DA:1669:A:O5'	2.18	0.44
55:DA:2321:G:N3	55:DA:2321:G:H2'	2.32	0.44
55:DA:248:G:H5'	55:DA:250:G:N7	2.32	0.44
55:DA:270(G):C:H2'	55:DA:270(H):C:H6	1.82	0.44
55:DA:2809:A:C2	55:DA:2892:A:C4	3.05	0.44
55:DA:616:A:H2'	55:DA:616:A:N3	2.33	0.44
55:DA:657:U:O2'	55:DA:658:C:H5'	2.18	0.44
55:DA:844:C:H2'	55:DA:845:G:O4'	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:934:G:H2'	55:DA:935:C:H6	1.82	0.44
4:DE:108:SER:O	4:DE:162:ALA:HA	2.18	0.44
5:DF:115:ALA:O	5:DF:116:ASP:C	2.55	0.44
6:DG:111:LEU:N	6:DG:112:PRO:CD	2.80	0.44
6:DG:181:ARG:CG	6:DG:181:ARG:O	2.64	0.44
6:DG:32:PRO:HA	6:DG:162:THR:OG1	2.18	0.44
7:DH:41:MET:HA	7:DH:53:GLU:O	2.18	0.44
7:DH:83:TYR:HB2	7:DH:84:SER:H	1.46	0.44
57:DY:135:ARG:HG2	56:DJ:10:GLU:HB3	1.99	0.44
8:DK:124:GLY:N	8:DK:142:VAL:HG21	2.33	0.44
8:DK:9:LEU:O	8:DK:10:GLU:HG3	2.17	0.44
58:DL:10:LEU:CD1	58:DL:55:VAL:HG21	2.47	0.44
11:DO:114:ILE:O	11:DO:115:LEU:HB3	2.18	0.44
19:DT:49:VAL:CG1	19:DT:83:VAL:HG13	2.48	0.44
21:DV:100:VAL:O	21:DV:124:ILE:HG22	2.18	0.44
25:DX:50:VAL:O	25:DX:51:ALA:C	2.55	0.44
23:DZ:80:LEU:HB2	23:DZ:82:LEU:CD2	2.47	0.44
23:DZ:83:GLU:HG2	23:DZ:84:GLY:H	1.80	0.44
26:A4:59:PHE:HE1	49:BV:67:VAL:HB	1.83	0.44
28:A6:25:LYS:CB	28:A6:25:LYS:HZ2	2.31	0.44
28:A6:27:LYS:HB2	28:A6:27:LYS:HE2	1.72	0.44
29:A7:24:THR:O	29:A7:25:PRO:C	2.56	0.44
30:A8:26:LYS:HB2	30:A8:44:LYS:O	2.17	0.44
1:AA:1180:C:H2'	1:AA:1181:C:O4'	2.17	0.44
1:AA:1670:C:O2	4:AE:129:HIS:NE2	2.42	0.44
1:AA:191:A:H2'	1:AA:192:C:H6	1.83	0.44
1:AA:2115:G:N7	1:AA:2117:A:OP2	2.51	0.44
1:AA:2274:A:C5	1:AA:2276:G:C8	3.06	0.44
1:AA:2418:A:OP2	30:A8:29:LYS:HE3	2.18	0.44
1:AA:2464:C:H2'	1:AA:2465:C:C6	2.53	0.44
1:AA:2649:U:H2'	1:AA:2650:U:C6	2.52	0.44
1:AA:2734:A:C8	1:AA:2735:G:C8	3.05	0.44
1:AA:569:U:C4	1:AA:570:G:C6	3.06	0.44
1:AA:654(S):G:O2'	1:AA:654(T):A:O5'	2.19	0.44
1:AA:733:G:H8	1:AA:733:G:O5'	2.00	0.44
1:AA:972:G:N1	1:AA:973:A:N6	2.66	0.44
3:AD:118:VAL:HG22	3:AD:119:ALA:N	2.33	0.44
4:AE:66:HIS:C	4:AE:68:ALA:N	2.63	0.44
5:AF:135:LYS:HB3	5:AF:138:GLU:HG3	1.99	0.44
5:AF:30:PRO:O	5:AF:31:HIS:C	2.55	0.44
6:AG:70:VAL:HA	6:AG:90:LEU:HD13	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:AN:64:ARG:NH1	10:AN:83:ALA:CB	2.81	0.44
11:AO:12:ALA:C	11:AO:14:LYS:N	2.70	0.44
11:AO:71:VAL:HG12	11:AO:72:PRO:HD3	1.98	0.44
14:AQ:110:LEU:CD2	14:AQ:111:GLU:N	2.80	0.44
20:AU:42:VAL:HG13	20:AU:65:ALA:O	2.18	0.44
25:AX:46:ASN:HA	25:AX:46:ASN:HD22	1.53	0.44
31:BA:1114:C:H2'	31:BA:1115:C:C6	2.52	0.44
1:AA:1948:G:N3	31:BA:1418:A:H2	2.15	0.44
31:BA:1512:U:H2'	31:BA:1513:A:C8	2.53	0.44
31:BA:1523:G:H2'	31:BA:1524:C:H6	1.83	0.44
31:BA:347:G:C2'	31:BA:348:G:H5'	2.47	0.44
31:BA:355:C:C4'	31:BA:388:G:O2'	2.66	0.44
31:BA:496:A:H2'	31:BA:496:A:N3	2.32	0.44
31:BA:558:G:H2'	31:BA:559:A:H2	1.83	0.44
31:BA:582:U:C2	31:BA:760:G:C6	3.06	0.44
31:BA:635:G:C6	31:BA:636:U:C4	3.06	0.44
31:BA:668:G:O2'	45:BR:46:HIS:CD2	2.68	0.44
31:BA:819:A:C5'	31:BA:820:U:OP2	2.66	0.44
31:BA:834:C:H2'	31:BA:835:U:C6	2.53	0.44
31:BA:849:C:O2'	31:BA:850:U:H5'	2.18	0.44
31:BA:860:A:H2'	31:BA:861:G:O4'	2.17	0.44
52:BD:18:G:H1'	52:BD:58:A:H2	1.82	0.44
52:BD:72:C:O2	52:BD:72:C:H2'	2.17	0.44
38:BK:36:LEU:HA	38:BK:39:LEU:HB2	2.00	0.44
38:BK:85:ARG:NH1	38:BK:85:ARG:HG3	2.33	0.44
40:BM:22:LYS:CD	40:BM:26:ALA:HB2	2.43	0.44
31:BA:1060:C:C5'	40:BM:51:ARG:HG2	2.44	0.44
40:BM:8:LEU:N	40:BM:8:LEU:HD12	2.33	0.44
41:BN:31:THR:O	41:BN:31:THR:HG23	2.18	0.44
41:BN:38:ASN:ND2	41:BN:38:ASN:N	2.66	0.44
42:BO:36:VAL:HG12	42:BO:37:CYS:N	2.32	0.44
43:BP:6:GLY:O	43:BP:7:VAL:HG13	2.17	0.44
54:CA:1088:G:O2'	54:CA:1089:G:H5'	2.18	0.44
54:CA:1125:U:O4	40:CM:5:ARG:HD3	2.18	0.44
54:CA:1158:C:H4'	32:CE:133:LYS:NZ	2.33	0.44
54:CA:1347:G:N2	54:CA:1374:A:OP2	2.51	0.44
54:CA:1521:G:H2'	54:CA:1522:U:H6	1.83	0.44
54:CA:243:A:H2	54:CA:245:C:H2'	1.82	0.44
54:CA:468:A:H2'	54:CA:474:G:H5'	2.00	0.44
54:CA:889:A:H1'	54:CA:891:U:C6	2.52	0.44
54:CA:991:U:O2	54:CA:993:G:C8	2.69	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:CF:118:GLN:O	33:CF:119:ARG:C	2.55	0.44
33:CF:174:PRO:C	33:CF:176:HIS:N	2.71	0.44
36:CI:99:ALA:O	36:CI:100:ASN:HB3	2.18	0.44
39:CL:80:GLY:C	39:CL:82:ALA:N	2.71	0.44
40:CM:54:PHE:CE1	40:CM:55:LYS:CE	3.01	0.44
42:CO:104:VAL:HG12	42:CO:105:TYR:CE1	2.53	0.44
43:CP:11:ARG:HB2	43:CP:11:ARG:NH1	2.32	0.44
51:CX:12:LYS:O	51:CX:16:GLY:N	2.50	0.44
51:CX:17:THR:O	51:CX:22:ARG:HD3	2.18	0.44
16:D1:39:LEU:O	16:D1:40:PHE:C	2.54	0.44
16:D1:74:LEU:C	16:D1:74:LEU:HD12	2.38	0.44
16:D1:69:CYS:O	16:D1:74:LEU:HD12	2.18	0.44
16:D1:81:HIS:CD2	16:D1:117:GLN:HG3	2.52	0.44
17:D2:64:HIS:N	17:D2:64:HIS:ND1	2.63	0.44
26:D4:23:GLU:OE1	26:D4:23:GLU:N	2.50	0.44
26:D4:38:LYS:HD3	26:D4:38:LYS:HA	1.47	0.44
26:D4:68:ARG:CB	26:D4:71:ARG:C	2.86	0.44
55:DA:1098:A:H2'	55:DA:1099:G:C5'	2.48	0.44
55:DA:1590:U:H2'	55:DA:1591:G:C8	2.53	0.44
55:DA:1694:C:HO2'	55:DA:1695:G:P	2.41	0.44
55:DA:1827:C:C2'	55:DA:1827:C:O2	2.64	0.44
55:DA:185:U:H2'	55:DA:186:G:C8	2.53	0.44
55:DA:2025:C:H2'	55:DA:2026:C:C6	2.53	0.44
55:DA:2311:A:O2'	55:DA:2312:U:O4'	2.29	0.44
55:DA:2529:G:H5''	55:DA:2530:A:H5''	1.99	0.44
55:DA:2555:U:H2'	55:DA:2556:C:H5'	2.00	0.44
55:DA:2850:A:H2	13:D0:61:HIS:CD2	2.36	0.44
55:DA:28:A:C2	55:DA:513:A:C8	3.05	0.44
55:DA:49:A:H5''	55:DA:50:U:C3'	2.33	0.44
55:DA:772:C:H5'	55:DA:1355:G:O2'	2.17	0.44
55:DA:794:G:H2'	55:DA:795:C:C6	2.53	0.44
55:DA:888:C:C3'	55:DA:889:C:H5'	2.48	0.44
2:DB:95:U:O5'	2:DB:95:U:H6	2.01	0.44
3:DD:70:TRP:HZ3	3:DD:146:GLU:CD	2.20	0.44
55:DA:727:A:H2	3:DD:9:TYR:CD2	2.36	0.44
4:DE:188:VAL:HG13	4:DE:188:VAL:O	2.18	0.44
5:DF:110:LEU:HD11	5:DF:181:LEU:HD13	2.00	0.44
5:DF:135:LYS:O	5:DF:138:GLU:N	2.49	0.44
5:DF:179:GLU:CD	5:DF:179:GLU:N	2.72	0.44
7:DH:54:ARG:NH2	7:DH:57:ASP:OD1	2.47	0.44
55:DA:2749:A:H4'	7:DH:62:LYS:HB3	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:DL:67:PHE:O	58:DL:68:VAL:CB	2.66	0.44
9:DM:128:HIS:O	9:DM:130:HIS:CD2	2.71	0.44
10:DN:112:MET:O	10:DN:113:LYS:C	2.57	0.44
11:DO:19:VAL:HG12	11:DO:27:HIS:HD2	1.83	0.44
15:DR:110:ILE:CG2	15:DR:111:ARG:HH11	2.31	0.44
20:DU:30:VAL:HG13	20:DU:37:VAL:HG12	1.99	0.44
24:DW:42:GLY:C	24:DW:44:LEU:N	2.69	0.44
57:DY:54:ALA:O	57:DY:55:LYS:C	2.56	0.44
55:DA:2199:A:H5'	23:DZ:50:ARG:HH21	1.81	0.44
23:DZ:56:GLN:NE2	23:DZ:56:GLN:H	2.14	0.44
13:A0:103:ARG:HD2	13:A0:108:GLY:O	2.18	0.43
28:A6:25:LYS:HB3	28:A6:25:LYS:NZ	2.33	0.43
28:A6:9:LEU:HD22	28:A6:11:LEU:HD21	1.99	0.43
30:A8:4:MET:HB3	30:A8:61:LEU:CD2	2.48	0.43
1:AA:1024:G:OP2	1:AA:1026:U:OP1	2.36	0.43
1:AA:1191:G:O2'	1:AA:1192:G:H5'	2.18	0.43
1:AA:123:G:O3'	1:AA:1376:C:H4'	2.18	0.43
1:AA:1417:C:H2'	1:AA:1418:G:C5'	2.48	0.43
1:AA:1454:U:C4'	1:AA:1455:G:OP1	2.55	0.43
1:AA:1490:A:C4'	1:AA:1491:G:OP2	2.66	0.43
1:AA:1936:A:H2'	1:AA:1945:G:N7	2.33	0.43
1:AA:2055:C:H4'	1:AA:2056:G:H5''	2.00	0.43
1:AA:2078:C:O2'	1:AA:2079:U:H5'	2.18	0.43
1:AA:2168:G:C2	1:AA:2170:A:OP2	2.71	0.43
1:AA:2311:A:O2'	1:AA:2312:U:C5'	2.66	0.43
1:AA:2592:G:C5	1:AA:2593:U:C5	3.06	0.43
1:AA:2702:U:C2'	1:AA:2702:U:O2	2.66	0.43
1:AA:281:G:O2'	1:AA:282:A:O4'	2.35	0.43
1:AA:2886:G:H2'	1:AA:2887:U:C6	2.53	0.43
1:AA:182:A:H2	1:AA:433:C:O2	2.01	0.43
1:AA:60:G:H5''	24:AW:54:LYS:NZ	2.32	0.43
1:AA:669:G:H3'	1:AA:670:A:C8	2.52	0.43
1:AA:727:A:H2	3:AD:9:TYR:CD2	2.36	0.43
1:AA:774:A:C2	1:AA:787:U:O2'	2.70	0.43
1:AA:969:U:H6	1:AA:969:U:O5'	2.01	0.43
1:AA:981:A:C2	1:AA:2027:G:N3	2.86	0.43
1:AA:992:C:H2'	1:AA:993:G:C8	2.53	0.43
3:AD:45:ASN:C	3:AD:45:ASN:OD1	2.56	0.43
4:AE:188:VAL:O	4:AE:188:VAL:HG13	2.18	0.43
4:AE:77:ILE:HA	4:AE:78:LEU:HD23	2.00	0.43
5:AF:1:MET:O	5:AF:2:LYS:C	2.56	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:AF:53:THR:O	5:AF:57:VAL:HG23	2.17	0.43
6:AG:129:GLY:O	6:AG:161:THR:HB	2.17	0.43
7:AH:44:VAL:CG1	7:AH:51:ARG:HB2	2.48	0.43
9:AM:66:LYS:O	9:AM:67:LEU:C	2.56	0.43
10:AN:25:LEU:HB2	10:AN:38:VAL:HG23	1.99	0.43
12:AP:10:ARG:HG3	12:AP:10:ARG:NH1	2.33	0.43
4:AE:14:ILE:CG1	15:AR:14:TYR:CZ	2.99	0.43
18:AS:43:GLY:O	18:AS:44:ALA:C	2.56	0.43
1:AA:138:G:N2	19:AT:50:LYS:HZ1	2.16	0.43
21:AV:91:LEU:HD12	21:AV:130:PRO:HG3	1.99	0.43
21:AV:150:LEU:CD2	21:AV:150:LEU:C	2.86	0.43
24:AW:40:SER:C	24:AW:42:GLY:H	2.20	0.43
25:AX:47:VAL:CG1	25:AX:56:VAL:HG21	2.47	0.43
23:AZ:78:LYS:CD	23:AZ:80:LEU:HD21	2.28	0.43
31:BA:1049:U:OP1	44:BQ:3:ARG:NH1	2.51	0.43
31:BA:1070:U:O2'	31:BA:1071:C:H5'	2.18	0.43
31:BA:1095:U:H2'	31:BA:1096:C:O4'	2.18	0.43
31:BA:1162:C:C2	31:BA:1175:G:C2	3.05	0.43
31:BA:1350:A:H2'	31:BA:1351:U:C6	2.52	0.43
31:BA:1508:G:O2'	31:BA:1509:C:H5'	2.18	0.43
31:BA:511:C:N1	31:BA:512:U:C5	2.85	0.43
52:BC:66:U:C4	52:BC:67:C:N4	2.86	0.43
34:BG:121:VAL:HG22	34:BG:126:ILE:HG13	2.00	0.43
35:BH:79:GLU:HB3	35:BH:93:PRO:HD2	2.00	0.43
37:BJ:140:ASP:O	37:BJ:143:ARG:HB2	2.18	0.43
37:BJ:74:GLU:HG2	37:BJ:91:VAL:HG22	2.00	0.43
38:BK:25:ASP:OD1	38:BK:25:ASP:N	2.51	0.43
40:BM:54:PHE:C	40:BM:55:LYS:CG	2.86	0.43
42:BO:27:LEU:HD11	42:BO:60:LEU:CB	2.45	0.43
43:BP:26:GLY:O	43:BP:28:ALA:N	2.50	0.43
26:A4:49:PHE:HZ	43:BP:61:GLU:O	2.01	0.43
49:BV:78:ARG:HH11	49:BV:78:ARG:HG2	1.83	0.43
53:C1:38:U:H2'	53:C1:39:U:C5'	2.48	0.43
54:CA:1003:G:C8	54:CA:1003:G:H3'	2.52	0.43
54:CA:186(C):G:C6	54:CA:191(E):G:C6	3.06	0.43
54:CA:197:A:N7	54:CA:221:C:H4'	2.33	0.43
54:CA:25:C:H2'	54:CA:26:A:C8	2.53	0.43
54:CA:404:U:H2'	54:CA:405:U:C6	2.53	0.43
54:CA:419:C:C4	54:CA:420:U:C5	3.06	0.43
54:CA:535:A:H4'	54:CA:536:C:OP1	2.13	0.43
54:CA:863:U:H2'	54:CA:865:A:OP2	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:96:G:C2	54:CA:97:U:H1'	2.53	0.43
32:CE:164:VAL:O	32:CE:186:ALA:HA	2.17	0.43
32:CE:193:ASP:OD2	32:CE:193:ASP:O	2.36	0.43
32:CE:33:TYR:HB2	32:CE:43:ASP:CB	2.42	0.43
32:CE:56:ARG:HH11	32:CE:56:ARG:CG	2.30	0.43
34:CG:159:ARG:O	34:CG:163:GLU:HB2	2.18	0.43
34:CG:31:CYS:O	34:CG:32:ALA:HB3	2.18	0.43
35:CH:114:GLY:O	35:CH:115:VAL:C	2.57	0.43
35:CH:11:ILE:HD12	35:CH:105:VAL:HG13	1.99	0.43
35:CH:50:GLU:HG3	35:CH:52:PRO:CD	2.37	0.43
35:CH:78:HIS:HB2	38:CK:104:ARG:HG2	1.99	0.43
38:CK:134:ILE:O	38:CK:135:CYS:HB3	2.17	0.43
38:CK:94:TYR:CE1	38:CK:132:GLU:HB2	2.53	0.43
54:CA:963:G:H21	40:CM:55:LYS:HZ3	1.65	0.43
43:CP:67:GLU:CD	43:CP:68:GLY:N	2.71	0.43
45:CR:76:GLU:HA	45:CR:79:ARG:HH12	1.83	0.43
48:CU:36:ASN:O	48:CU:37:VAL:C	2.56	0.43
13:D0:21:TYR:OH	13:D0:43:GLU:HG2	2.18	0.43
17:D2:38:LEU:N	17:D2:51:VAL:HG13	2.33	0.43
55:DA:1122:G:C2	55:DA:1123:C:C6	3.07	0.43
55:DA:1494:A:N3	55:DA:1494:A:H2'	2.33	0.43
55:DA:1726:G:C2'	55:DA:1727:U:H5'	2.47	0.43
55:DA:1931:U:C5	55:DA:1969:A:N7	2.86	0.43
55:DA:204:A:O2'	55:DA:205:G:P	2.76	0.43
55:DA:226:G:H2'	55:DA:227:A:C8	2.53	0.43
55:DA:2639:A:C2'	55:DA:2640:G:H5'	2.47	0.43
55:DA:270(F):U:H2'	55:DA:270(G):C:H6	1.79	0.43
55:DA:2712:U:H2'	55:DA:2712(A):A:H3'	1.98	0.43
55:DA:498:G:H21	20:DU:47:LYS:HZ2	1.66	0.43
55:DA:587:C:OP2	11:DO:21:ARG:NH2	2.50	0.43
55:DA:805:G:H5'	55:DA:806:C:OP2	2.18	0.43
55:DA:826:U:H2'	55:DA:828:U:O4'	2.18	0.43
55:DA:858:U:O2'	55:DA:2268:A:C1'	2.66	0.43
55:DA:880:G:N3	55:DA:880:G:C2'	2.79	0.43
55:DA:887:A:H1'	55:DA:889:C:N4	2.33	0.43
3:DD:35:LYS:CA	3:DD:64:ILE:HG22	2.48	0.43
7:DH:128:PRO:HG2	7:DH:129:THR:H	1.83	0.43
56:DI:5:ILE:O	56:DI:6:GLU:CG	2.66	0.43
8:DK:86:THR:O	8:DK:87:LYS:HB2	2.18	0.43
10:DN:77:ILE:HG23	10:DN:77:ILE:O	2.18	0.43
14:DQ:83:LYS:HE2	14:DQ:84:GLN:HG3	1.99	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:DW:36:ARG:O	24:DW:40:SER:HB2	2.18	0.43
57:DY:50:ARG:H	57:DY:84:GLU:N	2.15	0.43
13:A0:34:ILE:HG22	13:A0:114:VAL:HB	2.00	0.43
1:AA:2820:A:H1'	13:A0:3:HIS:CB	2.38	0.43
16:A1:113:ALA:C	16:A1:115:ALA:N	2.70	0.43
16:A1:90:VAL:HA	17:A2:39:LEU:HD22	1.99	0.43
16:A1:93:LYS:H	16:A1:93:LYS:CD	2.31	0.43
17:A2:29:PRO:HA	17:A2:61:VAL:HG21	1.99	0.43
26:A4:55:ARG:HD2	26:A4:56:VAL:HG23	2.00	0.43
29:A7:38:GLY:O	29:A7:39:ARG:C	2.55	0.43
30:A8:55:ALA:O	30:A8:56:GLU:C	2.57	0.43
1:AA:1021:A:H3'	1:AA:1021:A:H8	1.82	0.43
1:AA:1254:A:H5'	1:AA:1255:U:O5'	2.17	0.43
1:AA:141(A):C:H2'	1:AA:142:G:O4'	2.18	0.43
1:AA:2067:G:C4'	1:AA:2068:U:OP2	2.63	0.43
1:AA:2247:A:H2'	1:AA:2248:C:H6	1.83	0.43
1:AA:2473:U:C2'	1:AA:2473:U:O2	2.62	0.43
1:AA:2650:U:H2'	1:AA:2651:C:C6	2.53	0.43
1:AA:325:G:H2'	1:AA:326:G:O4'	2.18	0.43
1:AA:686:G:N2	1:AA:788:A:N6	2.60	0.43
1:AA:774:A:H2	1:AA:787:U:O2'	2.01	0.43
1:AA:918:A:H5''	2:AB:97:G:O2'	2.18	0.43
1:AA:846:C:H2'	1:AA:930:U:O4	2.17	0.43
1:AA:840:C:P	1:AA:932:G:H22	2.41	0.43
1:AA:955:C:H2'	1:AA:955:C:O2	2.18	0.43
1:AA:959:A:C6	1:AA:960:A:N1	2.87	0.43
1:AA:966:G:H2'	1:AA:967:C:C6	2.53	0.43
2:AB:44:G:C2	2:AB:48:A:N3	2.86	0.43
2:AB:30:C:C4'	2:AB:58:A:H2	2.31	0.43
3:AD:132:PRO:HD2	3:AD:135:PHE:CD1	2.53	0.43
3:AD:109:ASP:N	3:AD:196:VAL:O	2.51	0.43
3:AD:224:ALA:HA	3:AD:233:HIS:O	2.18	0.43
4:AE:4:ILE:CD1	4:AE:28:ALA:HB3	2.44	0.43
5:AF:132:VAL:C	5:AF:134:GLY:H	2.18	0.43
5:AF:39:TRP:O	5:AF:40:GLN:C	2.57	0.43
6:AG:101:ILE:CG1	6:AG:102:PHE:N	2.79	0.43
7:AH:4:ILE:HB	7:AH:6:ARG:NE	2.33	0.43
7:AH:44:VAL:O	7:AH:50:VAL:HG23	2.17	0.43
8:AK:18:VAL:O	8:AK:18:VAL:HG12	2.17	0.43
8:AK:81:VAL:HG12	8:AK:82:ARG:N	2.33	0.43
10:AN:24:VAL:HB	10:AN:33:ALA:HB2	1.99	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AP:110:THR:OG1	12:AP:112:GLU:HB2	2.18	0.43
14:AQ:25:ARG:HB3	14:AQ:25:ARG:HH11	1.79	0.43
15:AR:45:PHE:CE1	15:AR:65:LYS:HE2	2.53	0.43
15:AR:76:PHE:HA	15:AR:77:PRO:HD3	1.75	0.43
20:AU:76:CYS:O	20:AU:78:ALA:N	2.52	0.43
23:AZ:5:CYS:HB3	23:AZ:8:SER:OG	2.18	0.43
31:BA:1425:U:H2'	31:BA:1426:C:C6	2.53	0.43
31:BA:1460:A:H2'	31:BA:1461:G:O4'	2.18	0.43
31:BA:557:G:H2'	31:BA:558:G:C8	2.52	0.43
31:BA:65:U:H4'	31:BA:66:G:C5'	2.48	0.43
31:BA:697:U:H3'	31:BA:698:G:H8	1.83	0.43
31:BA:748:C:HO2'	31:BA:749:C:P	2.39	0.43
31:BA:995:C:O2'	31:BA:996:A:H5'	2.18	0.43
32:BE:51:LEU:HD23	32:BE:201:ILE:HD12	1.99	0.43
34:BG:138:TYR:C	34:BG:138:TYR:CD2	2.92	0.43
31:BA:412:A:N6	34:BG:35:ARG:HA	2.33	0.43
31:BA:921:U:O2	35:BH:19:MET:HB3	2.17	0.43
35:BH:63:ARG:O	35:BH:66:MET:HE1	2.18	0.43
36:BI:4:TYR:CE1	36:BI:92:LYS:HG2	2.53	0.43
36:BI:98:LEU:HD12	36:BI:98:LEU:O	2.18	0.43
38:BK:60:ARG:HH11	38:BK:60:ARG:HB2	1.82	0.43
42:BO:71:PRO:O	42:BO:102:ARG:NH1	2.50	0.43
43:BP:106:ASN:O	43:BP:107:ALA:HB3	2.18	0.43
44:BQ:23:ARG:O	44:BQ:24:CYS:C	2.56	0.43
45:BR:48:LYS:HD3	45:BR:48:LYS:HA	1.77	0.43
50:BW:48:LYS:O	50:BW:50:GLU:N	2.51	0.43
53:C1:56:U:C2'	53:C1:56:U:O2	2.66	0.43
54:CA:1000:A:H2'	54:CA:1001:G:H5'	2.00	0.43
54:CA:10:A:H2'	54:CA:11:G:H8	1.83	0.43
54:CA:1266:G:N2	54:CA:1270:C:C4	2.86	0.43
54:CA:1338:G:C6	54:CA:1339:A:C6	3.06	0.43
54:CA:188:U:C2'	54:CA:189:U:C5'	2.94	0.43
54:CA:38:G:N2	54:CA:397:A:C2	2.82	0.43
54:CA:408:A:C4	54:CA:409:G:C8	3.06	0.43
54:CA:40:C:H2'	54:CA:41:G:O4'	2.19	0.43
54:CA:411:A:C5	54:CA:429:U:C5	3.06	0.43
54:CA:518:C:O2	54:CA:518:C:C2'	2.64	0.43
54:CA:721:G:H1'	54:CA:722:A:C2	2.54	0.43
54:CA:939:G:C6	54:CA:940:C:N4	2.85	0.43
52:CB:16:U:H2'	52:CB:17:C:H5'	2.00	0.43
52:CB:19:G:H22	52:CB:56:C:N4	2.07	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:CE:71:VAL:O	32:CE:165:VAL:HG22	2.18	0.43
32:CE:7:VAL:HG23	32:CE:8:LYS:HD3	2.00	0.43
33:CF:58:GLU:HB2	33:CF:65:ALA:CB	2.40	0.43
34:CG:116:GLN:O	34:CG:119:GLN:HB3	2.18	0.43
34:CG:107:ARG:NH2	34:CG:194:LEU:HD11	2.33	0.43
34:CG:88:VAL:O	34:CG:90:GLY:N	2.44	0.43
36:CI:78:GLU:HA	36:CI:81:ILE:CD1	2.47	0.43
38:CK:103:VAL:HG21	38:CK:109:ILE:C	2.38	0.43
41:CN:57:THR:HG23	41:CN:58:PRO:HD2	1.99	0.43
42:CO:101:VAL:HG11	42:CO:104:VAL:HG21	2.01	0.43
43:CP:108:ARG:NH1	43:CP:112:GLY:O	2.50	0.43
46:CS:75:ARG:C	46:CS:77:ALA:N	2.69	0.43
48:CU:76:LEU:HB3	48:CU:78:LEU:HG	2.00	0.43
50:CW:101:GLY:O	50:CW:102:GLY:C	2.56	0.43
16:D1:52:ARG:CG	16:D1:52:ARG:NH1	2.80	0.43
6:DG:67:LYS:H	26:D4:6:HIS:CE1	2.36	0.43
27:D5:20:ARG:C	27:D5:22:HIS:N	2.71	0.43
55:DA:1042:G:H2'	55:DA:1043:C:C6	2.53	0.43
55:DA:1070:A:N6	55:DA:1096:A:C6	2.86	0.43
55:DA:1079:C:N4	55:DA:1080:A:N6	2.66	0.43
55:DA:1242:A:H5''	55:DA:1243:G:OP2	2.18	0.43
55:DA:1312:U:O2'	55:DA:1313:U:OP2	2.35	0.43
55:DA:1330:C:O2'	55:DA:1331:A:H5'	2.18	0.43
55:DA:1827:C:C2'	55:DA:1828:G:H5'	2.48	0.43
55:DA:1902:C:H2'	55:DA:1903:G:O5'	2.18	0.43
55:DA:2063:C:C5	55:DA:2064:C:C5	3.06	0.43
55:DA:2517:C:HO2'	55:DA:2518:A:P	2.40	0.43
55:DA:2735:G:H2'	55:DA:2736:G:H8	1.84	0.43
55:DA:2884:U:H2'	55:DA:2885:C:C5'	2.48	0.43
55:DA:405:U:C5'	55:DA:406:G:OP2	2.63	0.43
55:DA:481:G:C1'	55:DA:506:G:N2	2.79	0.43
55:DA:918:A:H2'	55:DA:919:G:O5'	2.18	0.43
3:DD:32:SER:O	3:DD:33:LEU:CB	2.47	0.43
6:DG:121:ASN:ND2	6:DG:121:ASN:C	2.71	0.43
6:DG:146:TYR:O	6:DG:149:VAL:N	2.50	0.43
6:DG:125:PHE:CD2	6:DG:166:ASP:HB2	2.53	0.43
6:DG:2:PRO:O	6:DG:4:ASP:N	2.50	0.43
6:DG:34:LEU:HD12	6:DG:100:TRP:CH2	2.53	0.43
6:DG:51:ARG:CB	6:DG:51:ARG:NH1	2.81	0.43
7:DH:155:SER:O	7:DH:156:ALA:O	2.36	0.43
7:DH:9:ILE:HD12	7:DH:49:VAL:HG11	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:DK:94:ALA:H	8:DK:116:LEU:HD13	1.83	0.43
8:DK:73:GLU:HB2	8:DK:136:VAL:CG2	2.48	0.43
8:DK:27:ARG:C	8:DK:28:ASN:HD22	2.21	0.43
58:DL:134:MET:O	58:DL:135:GLY:C	2.55	0.43
58:DL:143:GLU:O	58:DL:145:LYS:HG2	2.18	0.43
58:DL:44:ALA:C	58:DL:46:ALA:H	2.20	0.43
58:DL:58:THR:O	58:DL:59:ILE:HG12	2.17	0.43
58:DL:78:ILE:HD11	58:DL:127:ILE:CG2	2.48	0.43
10:DN:86:ILE:CD1	10:DN:86:ILE:H	2.31	0.43
12:DP:140:ALA:O	12:DP:141:GLN:C	2.56	0.43
55:DA:2496:C:OP1	12:DP:81:VAL:CG1	2.66	0.43
15:DR:122:ASP:O	15:DR:123:GLN:O	2.36	0.43
15:DR:67:SER:N	15:DR:70:VAL:O	2.51	0.43
18:DS:58:ALA:HB1	18:DS:64:MET:HB2	1.99	0.43
20:DU:50:ARG:C	20:DU:53:PRO:HD2	2.38	0.43
21:DV:182:LYS:HB3	21:DV:183:LEU:HD23	1.99	0.43
21:DV:1:MET:HB3	21:DV:2:GLU:H	1.56	0.43
57:DY:10:LEU:O	57:DY:11:ALA:C	2.57	0.43
57:DY:142:LEU:HD13	57:DY:143:GLN:CB	2.47	0.43
57:DY:13:LEU:CD2	57:DY:62:ALA:O	2.66	0.43
16:A1:50:ARG:HH11	17:A2:72:VAL:CG2	2.31	0.43
1:AA:995:C:H1'	16:A1:61:TRP:HZ2	1.82	0.43
17:A2:46:VAL:O	17:A2:48:GLY:N	2.51	0.43
29:A7:47:ARG:HB2	29:A7:47:ARG:HH11	1.83	0.43
1:AA:1032:A:H2	1:AA:1122:G:H22	1.65	0.43
1:AA:1187:G:H8	1:AA:1187:G:O5'	2.02	0.43
1:AA:1363:C:H2'	1:AA:1364:G:H8	1.82	0.43
1:AA:1407:C:H2'	1:AA:1407:C:O2	2.17	0.43
1:AA:1697:G:H3'	1:AA:1698:A:H5''	2.01	0.43
1:AA:2129:C:C2'	1:AA:2130:U:C5'	2.96	0.43
1:AA:2227:A:N6	1:AA:2228:G:C6	2.86	0.43
1:AA:2283:C:H2'	1:AA:2284:C:H5'	2.00	0.43
1:AA:2513:G:H2'	1:AA:2514:U:C6	2.53	0.43
1:AA:2679:A:O2'	1:AA:2680:C:H5'	2.19	0.43
1:AA:312:G:OP2	1:AA:312:G:C8	2.60	0.43
1:AA:337:C:H2'	1:AA:338:G:O4'	2.19	0.43
1:AA:26:G:H1'	1:AA:514:A:N6	2.33	0.43
1:AA:626:U:H5'	1:AA:627:A:C5'	2.47	0.43
1:AA:726:G:O2'	1:AA:727:A:P	2.76	0.43
1:AA:818:G:C2	1:AA:1190:G:O6	2.71	0.43
1:AA:830:G:H1'	1:AA:2448:A:C6	2.53	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:90:U:O2	1:AA:90:U:C2'	2.42	0.43
1:AA:91:A:H2'	1:AA:92:G:O4'	2.18	0.43
4:AE:5:LEU:N	4:AE:5:LEU:HD23	2.33	0.43
5:AF:154:VAL:HA	5:AF:191:ARG:O	2.18	0.43
6:AG:111:LEU:HD13	6:AG:120:LEU:HD21	1.99	0.43
6:AG:44:GLY:O	6:AG:47:LYS:HB2	2.18	0.43
6:AG:67:LYS:O	6:AG:67:LYS:HD2	2.19	0.43
8:AK:101:LEU:CG	8:AK:109:ILE:HD12	2.42	0.43
8:AK:114:LEU:HA	8:AK:130:TYR:HB2	2.00	0.43
8:AK:117:GLU:HB2	8:AK:118:LYS:H	1.50	0.43
8:AK:51:ILE:O	8:AK:53:ALA:N	2.51	0.43
8:AK:69:LYS:HD2	8:AK:69:LYS:C	2.38	0.43
10:AN:111:PHE:HB3	10:AN:114:ILE:CD1	2.45	0.43
11:AO:31:ALA:C	11:AO:32:THR:CG2	2.86	0.43
5:AF:31:HIS:CB	11:AO:9:ASN:ND2	2.78	0.43
12:AP:28:ALA:CB	12:AP:67:ARG:NH1	2.81	0.43
14:AQ:47:THR:C	14:AQ:48:LEU:HD12	2.39	0.43
20:AU:47:LYS:HG3	20:AU:60:PHE:HD1	1.84	0.43
20:AU:75:ILE:CG1	20:AU:76:CYS:N	2.81	0.43
21:AV:93:ASP:OD1	21:AV:93:ASP:O	2.37	0.43
53:B1:35:A:H2'	53:B1:36:G:O4'	2.18	0.43
31:BA:1069:C:O2'	31:BA:1192:C:H1'	2.18	0.43
31:BA:1187:G:H5'	31:BA:1188:A:OP2	2.17	0.43
31:BA:1190:G:P	33:BF:5:ILE:HG23	2.58	0.43
31:BA:1281:U:C2'	31:BA:1282:C:OP1	2.66	0.43
31:BA:1354:C:H2'	31:BA:1355:G:H8	1.83	0.43
31:BA:187:C:O2	31:BA:191(A):G:C6	2.72	0.43
31:BA:452:A:HO2'	31:BA:453:A:C4'	2.29	0.43
31:BA:511:C:O2'	31:BA:512:U:H5''	2.18	0.43
31:BA:563:A:H1'	31:BA:566:G:HO2'	1.83	0.43
31:BA:733:A:O2'	31:BA:734:G:C5'	2.66	0.43
31:BA:806:C:H2'	31:BA:807:A:H8	1.82	0.43
32:BE:9:GLU:O	32:BE:11:LEU:N	2.51	0.43
32:BE:137:ARG:C	32:BE:139:LYS:N	2.71	0.43
32:BE:221:LEU:HA	32:BE:224:GLN:CG	2.48	0.43
33:BF:35:GLU:N	33:BF:38:ARG:NH2	2.67	0.43
35:BH:36:ASP:C	35:BH:38:GLN:H	2.22	0.43
37:BJ:50:ILE:C	37:BJ:52:GLU:H	2.21	0.43
47:BT:66:SER:C	47:BT:67:LYS:O	2.55	0.43
48:BU:50:ILE:HD11	48:BU:70:ILE:CG2	2.42	0.43
49:BV:11:VAL:HG22	49:BV:12:ASP:N	2.33	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
50:BW:73:HIS:O	50:BW:74:LYS:HB2	2.18	0.43
31:BA:186:C:C1'	50:BW:81:LYS:NZ	2.80	0.43
54:CA:1004:A:C8	54:CA:1036:G:N2	2.86	0.43
54:CA:107:G:O6	50:CW:15:ARG:HD3	2.18	0.43
54:CA:152:A:N6	54:CA:170:U:C2	2.86	0.43
54:CA:420:U:O2	54:CA:420:U:H2'	2.18	0.43
54:CA:586:C:O2'	54:CA:878:G:H4'	2.18	0.43
54:CA:616:G:H2'	54:CA:616:G:N3	2.32	0.43
54:CA:628:G:C2	54:CA:629:G:N3	2.85	0.43
54:CA:967:C:H2'	54:CA:968:A:C8	2.53	0.43
54:CA:96:G:O2'	54:CA:97:U:H5'	2.18	0.43
52:CB:64:A:C2'	52:CB:65:G:H5'	2.48	0.43
32:CE:8:LYS:CE	32:CE:11:LEU:HD22	2.48	0.43
33:CF:109:PRO:C	33:CF:111:LEU:H	2.22	0.43
33:CF:126:ARG:NH1	33:CF:126:ARG:HG2	2.32	0.43
33:CF:70:VAL:CG1	33:CF:71:ALA:H	2.30	0.43
37:CJ:111:ARG:NH1	37:CJ:113:GLU:OE2	2.51	0.43
39:CL:8:GLY:CA	39:CL:79:LEU:HD12	2.30	0.43
39:CL:7:THR:O	39:CL:83:ARG:HD3	2.18	0.43
39:CL:92:TYR:O	39:CL:96:LEU:HB2	2.18	0.43
54:CA:1359:C:C5	44:CQ:35:ARG:HD2	2.53	0.43
45:CR:17:ARG:NH1	45:CR:17:ARG:HG3	2.27	0.43
45:CR:61:GLY:C	45:CR:65:ARG:HH12	2.20	0.43
46:CS:21:VAL:HG22	46:CS:34:GLU:O	2.19	0.43
13:D0:12:ARG:CG	13:D0:12:ARG:NH1	2.80	0.43
13:D0:70:LEU:HD12	13:D0:76:VAL:HG22	2.00	0.43
13:D0:75:LEU:HD13	13:D0:75:LEU:C	2.39	0.43
17:D2:61:VAL:C	17:D2:62:LEU:HD12	2.37	0.43
28:D6:37:ARG:O	28:D6:48:VAL:O	2.36	0.43
55:DA:1794:U:O4'	55:DA:1900:A:C2	2.71	0.43
55:DA:287:C:H2'	55:DA:288:C:C6	2.53	0.43
55:DA:2809:A:H62	55:DA:2891:G:H2'	1.83	0.43
55:DA:30:G:H2'	55:DA:31:C:H6	1.84	0.43
55:DA:270:A:H1'	55:DA:370:G:C2	2.53	0.43
55:DA:373:U:O2	55:DA:423:A:C2	2.69	0.43
55:DA:73:A:O5'	55:DA:73:A:H8	2.01	0.43
55:DA:74:A:O5'	55:DA:74:A:C8	2.72	0.43
55:DA:824:A:H1'	55:DA:2358:G:N7	2.33	0.43
55:DA:883:G:P	55:DA:883:G:H8	2.42	0.43
55:DA:97:C:O2	55:DA:97:C:H2'	2.17	0.43
2:DB:41:U:C4	6:DG:70:VAL:O	2.70	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:DB:58:A:H5'	2:DB:59:A:OP2	2.17	0.43
3:DD:89:SER:HB2	3:DD:159:ALA:CB	2.48	0.43
3:DD:25:THR:HG21	3:DD:82:ILE:N	2.28	0.43
3:DD:85:ASP:OD1	3:DD:85:ASP:C	2.54	0.43
3:DD:95:LEU:CD1	3:DD:97:TYR:CE1	3.01	0.43
4:DE:16:ARG:O	4:DE:17:ASP:HB2	2.18	0.43
6:DG:47:LYS:HD3	6:DG:81:LYS:CB	2.35	0.43
56:DI:27:LEU:O	56:DI:29:GLU:N	2.51	0.43
8:DK:110:ASP:OD2	8:DK:130:TYR:CE1	2.71	0.43
58:DL:102:GLU:C	58:DL:104:VAL:N	2.72	0.43
58:DL:54:PRO:HD2	58:DL:72:PRO:CB	2.47	0.43
58:DL:85:GLU:C	58:DL:86:LYS:HE2	2.38	0.43
55:DA:1666:G:C4'	10:DN:6:THR:HG23	2.46	0.43
10:DN:7:TYR:CE1	10:DN:20:MET:HB2	2.53	0.43
12:DP:2:LEU:HA	12:DP:2:LEU:HD13	1.74	0.43
14:DQ:111:GLU:O	14:DQ:112:PHE:HD1	2.01	0.43
15:DR:95:ARG:HA	15:DR:95:ARG:HD2	1.81	0.43
18:DS:28:SER:HB3	18:DS:31:GLU:HB2	2.01	0.43
20:DU:43:ASN:CA	20:DU:64:GLU:HA	2.48	0.43
20:DU:77:PRO:O	20:DU:78:ALA:CB	2.66	0.43
21:DV:124:ILE:HG23	21:DV:124:ILE:O	2.19	0.43
21:DV:145:GLU:O	21:DV:146:ILE:C	2.57	0.43
24:DW:17:SER:OG	24:DW:67:LYS:HE3	2.17	0.43
24:DW:18:PRO:CD	24:DW:19:VAL:H	2.28	0.43
24:DW:4:SER:OG	24:DW:5:GLU:N	2.51	0.43
57:DY:89:ALA:CB	57:DY:125:LEU:HD12	2.47	0.43
57:DY:133:GLU:HG2	57:DY:133:GLU:H	1.29	0.43
57:DY:28:ASN:N	57:DY:80:VAL:HG11	2.34	0.43
57:DY:50:ARG:N	57:DY:83:TYR:CA	2.80	0.43
13:A0:103:ARG:HD2	13:A0:108:GLY:C	2.38	0.43
13:A0:31:HIS:O	13:A0:33:ARG:N	2.52	0.43
16:A1:50:ARG:NH1	17:A2:72:VAL:HB	2.34	0.43
16:A1:50:ARG:NH1	17:A2:72:VAL:CB	2.80	0.43
30:A8:50:LEU:HD12	30:A8:53:PRO:O	2.17	0.43
1:AA:1008:C:N4	1:AA:1136:G:C6	2.87	0.43
1:AA:1162:G:H2'	1:AA:1163:G:C8	2.53	0.43
1:AA:1283:G:N2	1:AA:1285:G:H3'	2.32	0.43
1:AA:141:A:H1'	1:AA:1408:C:C1'	2.47	0.43
1:AA:1565:C:O2'	1:AA:1566:A:H8	2.02	0.43
1:AA:1603:A:OP1	1:AA:1604:C:OP2	2.37	0.43
1:AA:2438:U:O3'	1:AA:2439:A:H3'	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2523:G:H5'	1:AA:2523:G:C8	2.45	0.43
1:AA:2649:U:H2'	1:AA:2650:U:H6	1.83	0.43
1:AA:2653:U:O5'	1:AA:2653:U:H6	2.01	0.43
1:AA:2707:G:H2'	1:AA:2708:G:H8	1.83	0.43
1:AA:2791:C:C4	1:AA:2893:G:C6	3.06	0.43
1:AA:2824:C:H2'	1:AA:2825:C:C5'	2.43	0.43
1:AA:300:A:H2'	1:AA:334:C:H1'	1.99	0.43
1:AA:246:C:H4'	1:AA:385:C:O4'	2.18	0.43
1:AA:726:G:O2'	1:AA:727:A:H8	2.01	0.43
1:AA:870:A:C2	1:AA:871:U:H1'	2.53	0.43
2:AB:95:U:C6	2:AB:95:U:H3'	2.53	0.43
4:AE:199:ARG:HH11	4:AE:199:ARG:HB3	1.82	0.43
4:AE:33:VAL:O	4:AE:33:VAL:HG13	2.18	0.43
6:AG:75:LYS:HD2	6:AG:77:ILE:HD11	1.99	0.43
7:AH:115:VAL:HG11	7:AH:148:ILE:CD1	2.48	0.43
8:AK:133:HIS:N	8:AK:134:PRO:HD2	2.32	0.43
8:AK:56:LYS:C	8:AK:58:LEU:N	2.71	0.43
8:AK:98:ALA:O	8:AK:99:GLU:C	2.57	0.43
9:AM:36:GLY:H	9:AM:42:TRP:HE3	1.67	0.43
11:AO:147:LEU:C	11:AO:148:LEU:HD23	2.39	0.43
21:AV:121:HIS:HB2	21:AV:171:ILE:CD1	2.48	0.43
24:AW:33:MET:O	24:AW:36:ARG:HB2	2.16	0.43
24:AW:46:GLN:H	24:AW:49:LYS:HE3	1.83	0.43
53:B1:28:G:O2'	53:B1:29:G:O4'	2.28	0.43
53:B1:53:U:O2'	53:B1:54:U:H5''	2.18	0.43
31:BA:1365:G:C5	31:BA:1366:C:C5	3.07	0.43
31:BA:518:C:H4'	31:BA:519:C:C5'	2.49	0.43
31:BA:627:G:O2'	31:BA:628:G:H5'	2.18	0.43
31:BA:765:G:H22	31:BA:812:C:C2'	2.32	0.43
31:BA:881:G:C6	31:BA:882:C:C4	3.06	0.43
31:BA:960:U:C2'	31:BA:960:U:O2	2.64	0.43
52:BC:50:U:O2'	52:BC:51:U:H5'	2.19	0.43
52:BD:71:G:H2'	52:BD:72:C:O4'	2.19	0.43
32:BE:101:MET:CE	32:BE:108:ILE:HG21	2.48	0.43
32:BE:9:GLU:C	32:BE:11:LEU:H	2.21	0.43
33:BF:88:ARG:HG2	33:BF:101:LEU:HB3	2.00	0.43
34:BG:3:ARG:HD2	34:BG:118:ARG:HE	1.83	0.43
37:BJ:78:ARG:HD2	37:BJ:79:ARG:N	2.33	0.43
38:BK:5:PRO:O	38:BK:8:ASP:HB3	2.18	0.43
40:BM:74:ILE:CD1	40:BM:74:ILE:N	2.80	0.43
42:BO:60:LEU:HD21	42:BO:66:VAL:HG22	2.01	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:BU:31:LEU:HG	48:BU:65:ILE:CD1	2.44	0.43
50:BW:14:LYS:HA	50:BW:17:ARG:NE	2.34	0.43
54:CA:1060:C:N4	33:CF:2:GLY:HA3	2.31	0.43
54:CA:1152:A:OP1	40:CM:68:HIS:CD2	2.71	0.43
54:CA:1403:C:H1'	54:CA:1500:A:N1	2.34	0.43
54:CA:250:A:O2'	54:CA:251:G:P	2.76	0.43
54:CA:258:G:N3	54:CA:259:G:C8	2.86	0.43
54:CA:274:A:H1'	54:CA:275:G:O4'	2.18	0.43
54:CA:397:A:N7	54:CA:547:A:O2'	2.51	0.43
54:CA:412:A:H4'	54:CA:413:G:C5'	2.48	0.43
54:CA:701:C:C2'	54:CA:702:A:OP2	2.66	0.43
54:CA:718:G:C1'	41:CN:116:HIS:HA	2.48	0.43
54:CA:74:C:H2'	54:CA:75:C:H5'	2.00	0.43
54:CA:794:A:C5'	54:CA:794:A:C8	2.99	0.43
52:CD:59:U:H3'	52:CD:60:U:C6	2.50	0.43
35:CH:20:GLN:O	35:CH:21:ALA:O	2.35	0.43
37:CJ:113:GLU:HB3	37:CJ:118:VAL:HG22	2.01	0.43
37:CJ:6:ARG:HG2	37:CJ:6:ARG:O	2.18	0.43
42:CO:71:PRO:O	42:CO:102:ARG:HD2	2.19	0.43
45:CR:26:GLU:OE2	45:CR:77:ARG:NH1	2.49	0.43
48:CU:66:LEU:CD1	48:CU:70:ILE:HD11	2.49	0.43
30:D8:22:VAL:CB	30:D8:53:PRO:HB2	2.49	0.43
55:DA:105:C:C2	55:DA:106:C:C5	3.07	0.43
55:DA:1279:G:H4'	13:D0:31:HIS:CD2	2.53	0.43
55:DA:1562:A:H2'	55:DA:1563:G:H8	1.82	0.43
55:DA:195:A:H5''	11:DO:46:LYS:HZ1	1.83	0.43
55:DA:2097:C:O2'	55:DA:2098:U:H5'	2.18	0.43
55:DA:2262:U:H4'	55:DA:2328:A:C2	2.53	0.43
55:DA:2331:G:O2'	55:DA:2336:A:N1	2.36	0.43
55:DA:2880:C:O2'	13:D0:90:ARG:NH1	2.50	0.43
55:DA:627:A:H4'	55:DA:628:G:H5'	1.99	0.43
55:DA:846:C:C4	55:DA:930:U:C5	3.06	0.43
3:DD:92:ILE:HG22	3:DD:106:ILE:HA	2.01	0.43
3:DD:73:VAL:HA	3:DD:119:ALA:O	2.18	0.43
4:DE:103:ASP:OD2	4:DE:168:MET:CE	2.66	0.43
5:DF:57:VAL:HG13	5:DF:58:ALA:N	2.33	0.43
7:DH:6:ARG:HA	7:DH:66:GLY:HA2	2.00	0.43
58:DL:133:SER:C	58:DL:137:GLU:OE1	2.57	0.43
14:DQ:89:ARG:O	14:DQ:90:GLY:C	2.57	0.43
18:DS:66:GLU:C	18:DS:68:ARG:H	2.22	0.43
19:DT:39:ILE:O	19:DT:40:LYS:C	2.57	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:DU:63:LYS:HZ2	20:DU:64:GLU:HG3	1.83	0.43
21:DV:100:VAL:HG11	21:DV:137:ILE:HG13	1.99	0.43
25:DX:4:LEU:O	25:DX:36:VAL:HA	2.18	0.43
57:DY:135:ARG:HB2	56:DJ:19:GLU:OE2	2.19	0.43
57:DY:15:GLU:CD	57:DY:19:ARG:NH2	2.72	0.43
57:DY:54:ALA:O	57:DY:57:THR:HB	2.17	0.43
57:DY:74:LEU:N	57:DY:74:LEU:HD12	2.33	0.43
13:A0:13:HIS:O	13:A0:16:HIS:N	2.52	0.43
17:A2:44:LYS:C	17:A2:46:VAL:H	2.21	0.43
17:A2:71:LEU:O	17:A2:72:VAL:O	2.37	0.43
22:A3:74:ARG:NH1	22:A3:74:ARG:CG	2.78	0.43
26:A4:46:GLN:HE21	26:A4:46:GLN:HA	1.83	0.43
26:A4:51:ASP:O	26:A4:51:ASP:OD1	2.36	0.43
27:A5:33:CYS:O	27:A5:36:CYS:O	2.36	0.43
30:A8:32:LEU:CD2	30:A8:33:ASN:H	2.30	0.43
1:AA:1043:C:N4	1:AA:1044:G:N7	2.67	0.43
1:AA:1144:G:H3'	1:AA:1144:G:P	2.57	0.43
1:AA:1270:C:H5''	1:AA:1271:G:C5'	2.49	0.43
1:AA:1466:G:H2'	1:AA:1466:G:N3	2.33	0.43
1:AA:1509:C:OP1	1:AA:1509:C:H4'	2.18	0.43
1:AA:1528:A:N1	1:AA:1543:A:N1	2.66	0.43
1:AA:749:C:C2	1:AA:1618:A:H2'	2.53	0.43
1:AA:1881:C:H2'	1:AA:1881:C:O2	2.18	0.43
1:AA:1944:U:C1'	1:AA:1955:U:O4'	2.61	0.43
1:AA:1979:C:O2	1:AA:1979:C:H2'	2.18	0.43
1:AA:241:A:N1	1:AA:255:A:H5''	2.33	0.43
1:AA:1786:A:N1	1:AA:2606:C:C1'	2.81	0.43
1:AA:2652:C:O2'	1:AA:2653:U:H5'	2.18	0.43
1:AA:361:G:N2	1:AA:362:U:H1'	2.33	0.43
1:AA:371:A:H1'	1:AA:373:U:C6	2.54	0.43
1:AA:442:G:C2	1:AA:444:C:C5	3.05	0.43
1:AA:598:G:H2'	1:AA:599:G:O4'	2.17	0.43
1:AA:63:U:O2'	1:AA:64:A:C8	2.70	0.43
1:AA:73:A:O5'	1:AA:73:A:H8	2.00	0.43
1:AA:974:G:C2	1:AA:1186:G:C4	3.06	0.43
2:AB:27:C:N4	2:AB:28:C:N4	2.66	0.43
2:AB:56:G:H4'	2:AB:57:A:H8	1.82	0.43
3:AD:268:ARG:HD3	3:AD:269:PHE:CD1	2.54	0.43
4:AE:108:SER:O	4:AE:162:ALA:HA	2.19	0.43
4:AE:110:GLY:HA2	4:AE:162:ALA:N	2.34	0.43
4:AE:32:PRO:O	4:AE:33:VAL:HB	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:AE:34:VAL:HG21	4:AE:78:LEU:HD13	2.00	0.43
4:AE:69:LYS:HD2	4:AE:69:LYS:HA	1.79	0.43
6:AG:103:LEU:HD22	6:AG:178:PHE:HZ	1.83	0.43
6:AG:125:PHE:O	6:AG:127:GLY:N	2.52	0.43
6:AG:175:LEU:O	6:AG:176:LEU:C	2.54	0.43
7:AH:105:LEU:O	7:AH:105:LEU:CD2	2.66	0.43
8:AK:91:SER:OG	8:AK:119:PRO:CB	2.67	0.43
12:AP:134:ARG:O	12:AP:134:ARG:HG2	2.18	0.43
12:AP:54:MET:O	12:AP:55:VAL:C	2.56	0.43
12:AP:55:VAL:HG13	12:AP:56:ARG:N	2.34	0.43
12:AP:37:LEU:O	12:AP:99:PRO:HB3	2.19	0.43
14:AQ:106:ARG:CB	14:AQ:106:ARG:CZ	2.96	0.43
15:AR:111:ARG:HG3	15:AR:111:ARG:HH11	1.84	0.43
18:AS:70:TYR:O	18:AS:107:LEU:HA	2.19	0.43
18:AS:68:ARG:HB3	18:AS:110:LYS:H	1.84	0.43
19:AT:80:ILE:O	19:AT:80:ILE:HG12	2.18	0.43
1:AA:328:U:H4'	20:AU:68:HIS:CD2	2.53	0.43
20:AU:96:ILE:HD12	20:AU:98:VAL:HG12	2.01	0.43
21:AV:57:ILE:O	21:AV:68:PRO:HA	2.18	0.43
21:AV:81:ARG:CG	21:AV:81:ARG:O	2.66	0.43
25:AX:7:LYS:O	25:AX:54:VAL:HG13	2.18	0.43
31:BA:1028(A):C:H5'	31:BA:1028(A):C:H6	1.83	0.43
31:BA:1028:C:N4	31:BA:1028(A):C:N4	2.67	0.43
31:BA:1049:U:O2'	31:BA:1050:G:P	2.77	0.43
31:BA:1052:U:C2'	31:BA:1055:A:OP1	2.67	0.43
31:BA:1207:G:C6	31:BA:1208:C:C4	3.06	0.43
31:BA:1314:C:H2'	31:BA:1315:U:H6	1.83	0.43
31:BA:194:C:C2'	31:BA:195:A:H5''	2.48	0.43
31:BA:61:G:H2'	31:BA:62:U:O4'	2.18	0.43
31:BA:595:G:C6	31:BA:641:U:H2'	2.54	0.43
31:BA:862:C:O2'	31:BA:874:G:H5''	2.19	0.43
52:BB:39:U:H2'	52:BB:40:C:H6	1.82	0.43
52:BC:9:A:H5'	52:BC:46:G:O2'	2.19	0.43
52:BD:51:U:H2'	52:BD:52:G:C8	2.54	0.43
52:BD:67:C:H2'	52:BD:68:C:C6	2.53	0.43
32:BE:10:LEU:O	32:BE:13:ALA:HB3	2.18	0.43
32:BE:58:ILE:O	32:BE:62:ALA:N	2.52	0.43
31:BA:1190:G:P	33:BF:5:ILE:HD12	2.59	0.43
33:BF:63:ASN:HB3	33:BF:98:ASN:HB3	2.01	0.43
34:BG:105:VAL:HG13	34:BG:110:PHE:HB2	2.01	0.43
35:BH:40:ARG:HH11	35:BH:40:ARG:HG2	1.84	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:BI:67:MET:HB2	36:BI:68:PRO:HD2	2.01	0.43
39:BL:117:HIS:C	39:BL:118:LYS:HG3	2.39	0.43
40:BM:82:ILE:HG22	40:BM:82:ILE:O	2.19	0.43
49:BV:41:VAL:O	49:BV:44:MET:SD	2.76	0.43
31:BA:1313:U:C5	49:BV:4:SER:HB2	2.54	0.43
26:A4:59:PHE:HE1	49:BV:68:GLY:H	1.61	0.43
54:CA:920:U:O4'	54:CA:1080:A:C2	2.72	0.43
54:CA:1270:C:O2'	54:CA:1271:G:H5'	2.17	0.43
54:CA:452:A:H2'	54:CA:453:A:C8	2.53	0.43
54:CA:455:C:H6	54:CA:455:C:O5'	2.01	0.43
54:CA:565:U:C5'	54:CA:566:G:H3'	2.48	0.43
54:CA:724:G:O2'	54:CA:725:G:H5'	2.19	0.43
54:CA:827:U:C5	54:CA:870:U:C4	3.07	0.43
54:CA:940:C:H2'	54:CA:941:G:H8	1.83	0.43
32:CE:169:LYS:O	32:CE:169:LYS:HD3	2.18	0.43
32:CE:17:PHE:HB2	32:CE:44:LEU:HD11	2.01	0.43
33:CF:27:LYS:HZ3	33:CF:27:LYS:HA	1.83	0.43
35:CH:34:VAL:HG11	35:CH:63:ARG:HG2	2.00	0.43
36:CI:16:GLN:CD	36:CI:16:GLN:H	2.21	0.43
36:CI:67:MET:HB2	36:CI:68:PRO:HD2	1.99	0.43
37:CJ:140:ASP:HA	37:CJ:143:ARG:CZ	2.45	0.43
37:CJ:75:VAL:O	37:CJ:75:VAL:HG23	2.18	0.43
38:CK:6:ILE:CG2	38:CK:85:ARG:HH12	2.30	0.43
41:CN:58:PRO:HD3	41:CN:89:ALA:HB1	2.00	0.43
43:CP:10:PRO:O	43:CP:11:ARG:HB3	2.18	0.43
43:CP:54:VAL:O	43:CP:58:GLU:HG3	2.18	0.43
45:CR:58:MET:O	45:CR:61:GLY:N	2.51	0.43
47:CT:6:LEU:O	47:CT:58:GLU:HA	2.19	0.43
36:CI:98:LEU:HA	48:CU:31:LEU:CD2	2.48	0.43
48:CU:53:ARG:O	48:CU:55:ARG:N	2.52	0.43
16:D1:72:HIS:CE1	16:D1:107:ALA:HA	2.53	0.43
17:D2:37:VAL:O	17:D2:37:VAL:CG2	2.66	0.43
17:D2:72:VAL:HG13	17:D2:72:VAL:O	2.19	0.43
27:D5:58:LEU:O	27:D5:60:VAL:N	2.43	0.43
55:DA:1078:U:H3'	55:DA:1078:U:C6	2.54	0.43
55:DA:1265:A:N6	55:DA:2014:A:OP2	2.52	0.43
55:DA:1324:G:C4	55:DA:1328:G:O6	2.72	0.43
55:DA:1388:G:O2'	55:DA:1389:G:H5'	2.18	0.43
55:DA:1411:C:H5'	55:DA:1412:A:OP2	2.18	0.43
55:DA:1725:G:C5'	55:DA:1725:G:C8	2.93	0.43
55:DA:860:U:O2	55:DA:2268:A:O4'	2.36	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2307:G:H1'	55:DA:2308:G:C2	2.53	0.43
55:DA:2786:U:O2	4:DE:62:PRO:HG3	2.19	0.43
55:DA:372:G:O2'	55:DA:373:U:C6	2.70	0.43
55:DA:479:A:H4'	55:DA:480:A:C5'	2.49	0.43
55:DA:541:C:C2	55:DA:542:C:C5	3.07	0.43
55:DA:588:U:H1'	5:DF:90:PHE:HB3	2.01	0.43
3:DD:167:GLY:O	3:DD:173:VAL:HG23	2.18	0.43
3:DD:264:LYS:HG2	3:DD:266:SER:HB3	2.00	0.43
4:DE:144:ARG:HB3	4:DE:145:LYS:H	1.53	0.43
5:DF:37:VAL:HG12	5:DF:41:LEU:HD12	2.00	0.43
6:DG:115:ARG:O	6:DG:116:ASP:HB2	2.18	0.43
6:DG:146:TYR:O	6:DG:148:MET:N	2.52	0.43
55:DA:2313:C:H5''	6:DG:91:ARG:HD3	1.99	0.43
8:DK:31:LEU:N	8:DK:32:PRO:CD	2.82	0.43
58:DL:112:MET:SD	58:DL:120:LEU:CD1	2.99	0.43
58:DL:109:LYS:HB2	58:DL:120:LEU:HD21	2.01	0.43
58:DL:51:ALA:C	58:DL:52:ILE:HG13	2.38	0.43
58:DL:59:ILE:O	58:DL:60:TYR:HB2	2.18	0.43
58:DL:56:GLU:CB	58:DL:68:VAL:HG13	2.44	0.43
58:DL:52:ILE:CG1	58:DL:76:TYR:N	2.82	0.43
11:DO:19:VAL:CG2	11:DO:21:ARG:HD2	2.49	0.43
57:DY:7:VAL:O	57:DY:10:LEU:N	2.51	0.43
57:DY:117:LEU:HD21	56:DJ:24:ILE:HD11	2.00	0.43
57:DY:91:LYS:HZ2	57:DY:95:GLN:NE2	2.12	0.43
13:A0:72:ASP:HB3	13:A0:75:LEU:HB3	2.01	0.43
17:A2:58:VAL:HB	17:A2:98:GLU:HB2	2.00	0.43
26:A4:14:ILE:HG23	26:A4:14:ILE:O	2.19	0.43
28:A6:45:LYS:HB2	28:A6:46:HIS:H	1.36	0.43
30:A8:56:GLU:O	30:A8:59:LYS:N	2.44	0.43
1:AA:1091:G:H2'	1:AA:1092:C:H6	1.82	0.43
1:AA:1153:C:H5'	16:A1:76:TYR:HE2	1.84	0.43
1:AA:1482:U:H5'	1:AA:1483:G:OP2	2.18	0.43
1:AA:1652:A:H4'	1:AA:1653:G:OP1	2.18	0.43
1:AA:1812:A:H2'	1:AA:1813:G:H8	1.83	0.43
1:AA:2014:A:H2'	1:AA:2015:A:C8	2.52	0.43
1:AA:2287:A:O2'	1:AA:2288:A:O5'	2.27	0.43
1:AA:2392:A:N1	1:AA:2424:C:N3	2.67	0.43
1:AA:2063:C:O2	1:AA:2450:A:N1	2.52	0.43
1:AA:2457:U:H2'	1:AA:2458:G:O4'	2.18	0.43
1:AA:2552:U:OP2	1:AA:2552:U:H6	2.02	0.43
1:AA:2720:U:H2'	1:AA:2721:A:H8	1.82	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2852:G:C6	1:AA:2853:C:C4	3.07	0.43
1:AA:291:C:O2'	1:AA:292:C:H5'	2.19	0.43
1:AA:299:A:N6	1:AA:322:A:O2'	2.51	0.43
1:AA:654(C):G:C6	1:AA:654(D):G:C6	3.06	0.43
1:AA:840:C:OP2	1:AA:932:G:N2	2.48	0.43
1:AA:855:G:H2'	1:AA:856:C:C6	2.53	0.43
1:AA:943:U:O2'	1:AA:944:G:H5'	2.17	0.43
1:AA:9:U:H5'	9:AM:115:ARG:HH22	1.84	0.43
2:AB:45:A:OP1	2:AB:45:A:H8	2.01	0.43
2:AB:79:C:H2'	2:AB:80:U:H5'	1.99	0.43
3:AD:154:LYS:C	3:AD:155:LEU:HD12	2.38	0.43
3:AD:176:ARG:HH11	3:AD:176:ARG:CG	2.22	0.43
3:AD:176:ARG:NH1	3:AD:176:ARG:CG	2.80	0.43
3:AD:223:GLY:HA3	3:AD:231:HIS:CE1	2.53	0.43
1:AA:1816:G:O6	3:AD:37:LEU:HD11	2.19	0.43
3:AD:68:LYS:O	3:AD:70:TRP:N	2.51	0.43
5:AF:195:ASP:OD1	5:AF:197:ASP:HB3	2.18	0.43
7:AH:87:LEU:CD2	7:AH:149:ARG:HB2	2.49	0.43
7:AH:37:VAL:HG22	7:AH:38:SER:H	1.83	0.43
7:AH:51:ARG:HH11	7:AH:51:ARG:HG3	1.84	0.43
8:AK:72:LEU:CD2	8:AK:107:VAL:HG21	2.45	0.43
9:AM:46:VAL:O	9:AM:47:ALA:CB	2.67	0.43
11:AO:64:LYS:O	11:AO:66:GLY:N	2.47	0.43
12:AP:24:GLY:O	12:AP:25:ASP:CB	2.56	0.43
12:AP:50:ALA:O	12:AP:53:ALA:HB3	2.18	0.43
20:AU:20:TYR:CE2	20:AU:42:VAL:HA	2.53	0.43
20:AU:47:LYS:CA	20:AU:60:PHE:HB3	2.41	0.43
21:AV:94:GLU:O	21:AV:130:PRO:HD3	2.19	0.43
31:BA:1024:G:H2'	31:BA:1025:U:O4'	2.18	0.43
31:BA:1026:G:C2	31:BA:1027:C:H1'	2.54	0.43
31:BA:1139:G:N2	31:BA:1143:G:C6	2.86	0.43
31:BA:31:G:O2'	31:BA:32:A:OP1	2.36	0.43
31:BA:495:A:O2'	31:BA:496:A:H5''	2.18	0.43
31:BA:701:C:H1'	31:BA:703:G:N3	2.33	0.43
31:BA:788:U:H3	31:BA:795:C:N4	2.15	0.43
31:BA:811:C:O2'	31:BA:901:A:N1	2.46	0.43
31:BA:991:U:O2	31:BA:993:G:C8	2.72	0.43
52:BC:14:A:C5	52:BC:22:G:C2	3.06	0.43
52:BC:17:C:C6	52:BC:17:C:H3'	2.53	0.43
52:BD:9:A:H4'	52:BD:46:G:C4'	2.48	0.43
52:BD:70:G:O2'	52:BD:71:G:H5'	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:BG:24:GLU:CG	34:BG:25:ARG:H	2.32	0.43
34:BG:34:GLU:O	34:BG:35:ARG:CG	2.66	0.43
40:BM:4:ILE:HD12	40:BM:74:ILE:CG1	2.48	0.43
40:BM:67:THR:HG22	40:BM:67:THR:O	2.18	0.43
40:BM:92:THR:HG23	40:BM:93:GLY:N	2.34	0.43
44:BQ:6:LEU:C	44:BQ:8:GLU:N	2.72	0.43
46:BS:1:MET:HE1	46:BS:65:GLN:CB	2.48	0.43
49:BV:53:ASN:HB2	49:BV:77:THR:CG2	2.46	0.43
50:BW:68:LYS:C	50:BW:68:LYS:HD2	2.38	0.43
54:CA:1126:U:O4	54:CA:1127:G:C2	2.71	0.43
54:CA:815:A:N6	54:CA:1509:C:H1'	2.33	0.43
54:CA:262:A:C6	54:CA:263:A:C6	3.06	0.43
54:CA:447:G:H2'	54:CA:485:G:N2	2.33	0.43
54:CA:577:G:H1'	54:CA:816:A:C4	2.54	0.43
54:CA:629:G:O3'	54:CA:630:G:O4'	2.37	0.43
54:CA:748:C:H1'	54:CA:749:C:C5	2.52	0.43
54:CA:976:G:C8	54:CA:1358:U:C2	3.07	0.43
52:CB:4:C:O2'	52:CB:5:G:H5'	2.19	0.43
32:CE:125:PRO:O	32:CE:126:GLU:HB2	2.17	0.43
32:CE:57:PHE:CE2	32:CE:61:LEU:HD22	2.54	0.43
33:CF:130:VAL:HG21	33:CF:157:ILE:HG23	2.01	0.43
35:CH:110:LEU:O	35:CH:115:VAL:HB	2.18	0.43
35:CH:73:ASN:O	35:CH:73:ASN:ND2	2.52	0.43
36:CI:83:ASP:N	36:CI:83:ASP:OD2	2.51	0.43
37:CJ:91:VAL:HG12	37:CJ:95:ARG:HB3	2.00	0.43
43:CP:65:LYS:HB2	43:CP:69:GLU:HB2	2.01	0.43
49:CV:33:THR:OG1	49:CV:34:TRP:N	2.52	0.43
13:D0:18:LEU:O	13:D0:22:ARG:HG3	2.18	0.43
17:D2:52:VAL:O	17:D2:52:VAL:HG23	2.19	0.43
29:D7:46:VAL:O	29:D7:47:ARG:HB3	2.18	0.43
55:DA:1025:G:O2'	55:DA:1026:U:P	2.76	0.43
55:DA:1034:G:H2'	55:DA:1035:U:O4'	2.18	0.43
55:DA:1187:G:H8	55:DA:1187:G:O5'	2.02	0.43
55:DA:1235:G:C6	55:DA:1236:G:N1	2.87	0.43
55:DA:1252:G:O2'	55:DA:1253:A:O4'	2.36	0.43
55:DA:1359:A:N3	55:DA:1373:A:C2	2.86	0.43
55:DA:1879:C:C2'	55:DA:1880:C:H5'	2.46	0.43
55:DA:2340:G:HO2'	55:DA:2341:G:H5'	1.81	0.43
55:DA:2583:G:H2'	55:DA:2584:U:O2	2.17	0.43
55:DA:2657:A:H2'	55:DA:2658:C:C5'	2.48	0.43
55:DA:2884:U:H2'	55:DA:2885:C:O4'	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:299:A:C5'	55:DA:300:A:OP2	2.66	0.43
55:DA:301:G:O2'	55:DA:302:C:O4'	2.36	0.43
55:DA:310:A:O2'	55:DA:311:A:C2'	2.67	0.43
55:DA:340:A:H2'	55:DA:341:G:C5'	2.49	0.43
55:DA:885:C:C5	55:DA:886:C:C2	3.07	0.43
55:DA:897:C:H6	55:DA:897:C:C4'	2.30	0.43
3:DD:96:HIS:CE1	3:DD:102:LYS:HZ3	2.35	0.43
3:DD:143:HIS:HB2	3:DD:156:ALA:O	2.18	0.43
3:DD:96:HIS:CE1	3:DD:102:LYS:HZ2	2.36	0.43
5:DF:119:ARG:HG2	5:DF:119:ARG:HH11	1.83	0.43
6:DG:137:GLU:HB2	6:DG:152:LEU:HD22	2.00	0.43
6:DG:34:LEU:HD22	6:DG:35:GLU:N	2.33	0.43
6:DG:83:ARG:HG2	6:DG:83:ARG:NH1	2.34	0.43
7:DH:124:GLU:HB2	7:DH:132:ARG:HG3	1.99	0.43
7:DH:147:ASN:ND2	7:DH:147:ASN:H	2.15	0.43
8:DK:38:LEU:H	8:DK:38:LEU:CD1	2.05	0.43
9:DM:67:LEU:HA	9:DM:87:LEU:HD13	2.00	0.43
11:DO:75:ILE:HG12	11:DO:77:ARG:HH12	1.82	0.43
55:DA:2846:G:OP2	15:DR:54:ARG:HB2	2.19	0.43
21:DV:108:PRO:HG2	21:DV:109:ALA:H	1.83	0.43
21:DV:98:MET:O	21:DV:125:LEU:HA	2.18	0.43
21:DV:140:ASP:OD2	21:DV:141:VAL:N	2.52	0.43
24:DW:14:ARG:HG2	24:DW:14:ARG:HH11	1.84	0.43
57:DY:104:ILE:CG2	57:DY:105:PRO:HD2	2.48	0.43
57:DY:49:ALA:O	57:DY:50:ARG:CB	2.64	0.43
23:DZ:92:LYS:HE3	23:DZ:92:LYS:HB2	1.73	0.43
13:A0:37:THR:HB	13:A0:40:LYS:CG	2.49	0.43
26:A4:2:LYS:CB	26:A4:6:HIS:NE2	2.75	0.43
27:A5:46:CYS:HB3	27:A5:49:CYS:SG	2.59	0.43
28:A6:20:ASN:ND2	28:A6:42:TRP:CE3	2.84	0.43
1:AA:1024:G:C6	1:AA:1025:G:N1	2.86	0.43
1:AA:1152:C:H2'	1:AA:1153:C:H6	1.83	0.43
1:AA:1188:U:H2'	1:AA:1189:A:H5'	1.98	0.43
1:AA:1417:C:C2'	1:AA:1418:G:H5'	2.48	0.43
1:AA:1441:G:H2'	1:AA:1442:G:H8	1.83	0.43
1:AA:1773:A:C5	1:AA:1829:A:H1'	2.53	0.43
1:AA:21:A:H2'	1:AA:22:C:C6	2.54	0.43
1:AA:2317:C:H3'	1:AA:2318:G:H21	1.84	0.43
1:AA:2724:C:H2'	1:AA:2725:A:C8	2.53	0.43
1:AA:2741:A:N6	1:AA:2763:G:H2'	2.30	0.43
1:AA:2795:G:C3'	1:AA:2797:U:C5'	2.96	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2889:C:H3'	1:AA:2891:G:H8	1.84	0.43
1:AA:65:C:O2'	1:AA:456:C:O2	2.34	0.43
1:AA:554:U:H2'	1:AA:556:G:N7	2.34	0.43
1:AA:803:U:O2'	1:AA:804:A:H5'	2.19	0.43
1:AA:958:U:O2	2:AB:89(A):A:O2'	2.36	0.43
2:AB:33:G:H2'	2:AB:34:U:O4'	2.18	0.43
3:AD:124:PRO:O	3:AD:129:ASN:ND2	2.48	0.43
3:AD:131:LEU:HB2	3:AD:136:ILE:CD1	2.48	0.43
3:AD:161:THR:O	3:AD:196:VAL:HG23	2.19	0.43
1:AA:1820:U:O2'	3:AD:201:HIS:CD2	2.72	0.43
3:AD:218:ARG:HB3	3:AD:219:PRO:HD2	1.99	0.43
3:AD:31:LYS:NZ	3:AD:102:LYS:NZ	2.67	0.43
3:AD:31:LYS:O	3:AD:35:LYS:O	2.36	0.43
5:AF:17:ARG:NH1	5:AF:17:ARG:HG3	2.33	0.43
5:AF:68:LYS:HD2	5:AF:68:LYS:HA	1.37	0.43
6:AG:107:LEU:HD11	6:AG:178:PHE:CE1	2.54	0.43
2:AB:56:G:H5'	6:AG:27:ASN:HD21	1.83	0.43
6:AG:76:SER:C	6:AG:77:ILE:HD12	2.38	0.43
9:AM:95:PRO:C	9:AM:97:ARG:N	2.70	0.43
10:AN:8:LEU:N	10:AN:8:LEU:CD2	2.81	0.43
12:AP:101:ARG:HG3	12:AP:102:VAL:H	1.83	0.43
12:AP:139:GLU:HB2	12:AP:140:ALA:H	1.54	0.43
14:AQ:106:ARG:NH1	14:AQ:106:ARG:HB3	2.27	0.43
15:AR:19:LEU:HA	15:AR:20:PRO:HD3	1.61	0.43
18:AS:66:GLU:O	18:AS:68:ARG:N	2.52	0.43
19:AT:35:THR:C	19:AT:37:THR:N	2.72	0.43
21:AV:120:ILE:HG22	21:AV:121:HIS:CD2	2.54	0.43
21:AV:147:GLY:O	21:AV:148:ASP:O	2.36	0.43
24:AW:33:MET:CG	24:AW:37:PHE:HE1	2.32	0.43
1:AA:988:A:OP2	25:AX:11:SER:HB2	2.19	0.43
25:AX:8:LEU:HD22	25:AX:9:VAL:N	2.34	0.43
31:BA:1004:A:HO2'	31:BA:1005:A:C4'	2.31	0.43
31:BA:1067:A:O2'	31:BA:1068:G:O4'	2.34	0.43
31:BA:1095:U:C5'	31:BA:1109:C:O2	2.66	0.43
31:BA:1152:A:O2'	31:BA:1153:C:H5'	2.18	0.43
31:BA:1382:C:H2'	31:BA:1383:C:H6	1.83	0.43
31:BA:1496:C:H2'	31:BA:1497:G:O4'	2.18	0.43
31:BA:19:C:O2'	31:BA:20:U:H5'	2.19	0.43
31:BA:360:A:H2'	31:BA:361:G:O4'	2.18	0.43
31:BA:60:A:O2'	31:BA:61:G:OP2	2.36	0.43
31:BA:823:G:H21	38:BK:1:MET:HE1	1.84	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:BB:44:G:H2'	52:BB:45:U:C6	2.54	0.43
52:BC:33:U:C2	52:BC:35:A:OP2	2.72	0.43
52:BC:68:C:H2'	52:BC:69:G:C8	2.53	0.43
32:BE:213:LEU:HD23	32:BE:213:LEU:O	2.18	0.43
32:BE:5:ILE:HB	32:BE:59:GLU:CD	2.39	0.43
31:BA:1055:A:O2'	33:BF:156:ARG:NH2	2.52	0.43
33:BF:196:LEU:N	33:BF:196:LEU:HD22	2.34	0.43
33:BF:21:ARG:O	33:BF:58:GLU:HA	2.18	0.43
33:BF:6:HIS:NE2	33:BF:184:TYR:CE2	2.87	0.43
33:BF:76:VAL:HG23	33:BF:77:ILE:H	1.84	0.43
33:BF:90:GLU:O	33:BF:93:LYS:CB	2.64	0.43
39:BL:117:HIS:CD2	39:BL:123:PRO:HA	2.54	0.43
39:BL:14:VAL:O	39:BL:65:VAL:HA	2.18	0.43
42:BO:89:ARG:CG	42:BO:90:VAL:N	2.81	0.43
43:BP:17:VAL:O	43:BP:19:LEU:N	2.51	0.43
44:BQ:29:ARG:HG2	44:BQ:40:CYS:HB3	1.97	0.43
31:BA:134:A:N6	46:BS:25:ARG:HH12	2.06	0.43
48:BU:18:ARG:HA	48:BU:18:ARG:HD2	1.73	0.43
49:BV:26:GLY:C	49:BV:27:GLU:OE1	2.57	0.43
49:BV:22:LEU:HD11	49:BV:28:LYS:O	2.18	0.43
49:BV:36:ARG:HE	49:BV:72:GLY:CA	2.32	0.43
49:BV:71:LEU:O	49:BV:73:GLU:N	2.51	0.43
53:C1:47:U:H2'	53:C1:48:U:O4'	2.19	0.43
54:CA:1149:C:H2'	54:CA:1150:U:C6	2.54	0.43
54:CA:210:U:O2'	54:CA:216:G:C8	2.66	0.43
54:CA:503:C:H2'	54:CA:504:C:H6	1.83	0.43
52:CB:18:G:O2'	52:CB:19:G:O5'	2.37	0.43
32:CE:166:ASP:OD2	32:CE:168:THR:HG22	2.19	0.43
32:CE:30:ARG:HG3	32:CE:31:TYR:CD1	2.52	0.43
33:CF:29:TYR:CD2	33:CF:29:TYR:C	2.92	0.43
35:CH:12:LEU:O	35:CH:30:ALA:HA	2.19	0.43
36:CI:2:ARG:CZ	36:CI:69:GLU:HG3	2.48	0.43
39:CL:43:ALA:O	39:CL:45:ALA:N	2.52	0.43
54:CA:1151:A:C1'	40:CM:39:PRO:HB2	2.48	0.43
54:CA:1525:G:P	41:CN:120:ARG:HH22	2.42	0.43
41:CN:77:MET:HG3	41:CN:78:GLN:N	2.34	0.43
44:CQ:51:GLY:O	44:CQ:53:LEU:N	2.51	0.43
45:CR:50:HIS:O	45:CR:53:HIS:HB3	2.19	0.43
45:CR:64:ARG:HH11	45:CR:64:ARG:CG	2.31	0.43
49:CV:5:LEU:HD13	49:CV:5:LEU:O	2.18	0.43
26:D4:27:THR:O	26:D4:28:LYS:CB	2.67	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2344:U:OP1	28:D6:38:LYS:HE2	2.19	0.43
29:D7:25:PRO:CA	29:D7:28:ARG:NH1	2.81	0.43
55:DA:1040:C:H2'	55:DA:1041:C:H6	1.83	0.43
55:DA:1078:U:C1'	55:DA:1088:A:H2	2.13	0.43
55:DA:1509:C:H3'	55:DA:1510:A:H5''	2.01	0.43
55:DA:1434:A:N6	55:DA:1558:A:N6	2.50	0.43
55:DA:1643:G:N2	55:DA:1644:C:H1'	2.34	0.43
55:DA:190:A:P	55:DA:205:G:H22	2.42	0.43
55:DA:2037:G:H2'	55:DA:2038:G:C8	2.54	0.43
55:DA:2134:A:H8	55:DA:2157:G:H21	1.63	0.43
55:DA:2115:G:N3	55:DA:2171:A:N1	2.67	0.43
55:DA:2325:G:H8	55:DA:2325:G:O5'	2.01	0.43
55:DA:2415:G:C4	55:DA:2416:C:C5	3.07	0.43
55:DA:2486:G:H2'	55:DA:2487:G:O4'	2.18	0.43
55:DA:2728:U:H2'	55:DA:2729:G:H8	1.83	0.43
55:DA:2772:C:C2	55:DA:2773:C:C5	3.06	0.43
55:DA:27:G:C2'	55:DA:28:A:OP2	2.66	0.43
55:DA:2816:C:H4'	13:D0:99:LYS:HZ1	1.84	0.43
55:DA:307:G:N2	55:DA:310:A:C8	2.87	0.43
55:DA:366:C:H5	55:DA:403:U:HO2'	1.47	0.43
55:DA:442:G:N3	5:DF:48:THR:CG2	2.73	0.43
55:DA:587:C:N3	11:DO:33:ARG:NH1	2.67	0.43
55:DA:69:C:C2'	55:DA:70:G:H5'	2.47	0.43
55:DA:753:C:O2'	55:DA:754:C:H5'	2.18	0.43
55:DA:784:A:H5''	3:DD:227:ASN:ND2	2.34	0.43
55:DA:687:C:H42	55:DA:787:U:H4'	1.84	0.43
55:DA:994:C:OP2	16:D1:54:LYS:NZ	2.37	0.43
3:DD:168:ARG:HA	3:DD:173:VAL:HA	2.01	0.43
3:DD:220:HIS:C	3:DD:220:HIS:CD2	2.90	0.43
3:DD:273:ARG:CG	3:DD:273:ARG:O	2.53	0.43
5:DF:9:ILE:CG2	5:DF:20:LEU:O	2.66	0.43
7:DH:43:VAL:HG23	7:DH:43:VAL:O	2.19	0.43
7:DH:12:PRO:HD3	7:DH:48:GLY:O	2.19	0.43
58:DL:99:ILE:H	58:DL:138:VAL:HG22	1.83	0.43
58:DL:52:ILE:HG22	58:DL:75:SER:HB2	1.81	0.43
9:DM:95:PRO:O	9:DM:96:GLU:CD	2.57	0.43
11:DO:66:GLY:O	11:DO:67:MET:HB2	2.19	0.43
21:DV:117:LEU:CD1	21:DV:118:GLN:N	2.59	0.43
55:DA:1075:C:C4'	21:DV:195:GLU:CD	2.87	0.43
13:A0:76:VAL:CG1	13:A0:77:ARG:N	2.82	0.43
17:A2:87:HIS:ND1	17:A2:88:ARG:N	2.65	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:A3:84:LEU:CD1	22:A3:84:LEU:H	2.32	0.43
27:A5:16:ARG:NH1	27:A5:17:ASP:OD1	2.51	0.43
28:A6:34:LEU:HG	28:A6:36:LEU:HG	1.99	0.43
29:A7:25:PRO:HD3	29:A7:28:ARG:HH22	1.83	0.43
29:A7:48:LYS:HD3	29:A7:49:ARG:H	1.83	0.43
1:AA:51:G:N3	1:AA:119:A:C2	2.87	0.43
1:AA:1342:A:C2	1:AA:1397:U:C2	3.07	0.43
1:AA:1388:G:O2'	1:AA:1389:G:H5'	2.19	0.43
1:AA:1427:A:O2'	1:AA:1428:C:OP2	2.29	0.43
1:AA:1542:G:C5	1:AA:1543:A:N1	2.87	0.43
1:AA:1547:C:H2'	1:AA:1548:C:C6	2.54	0.43
1:AA:1577:C:H2'	1:AA:1578:U:C1'	2.48	0.43
1:AA:1585:C:O2	1:AA:1585:C:C2'	2.66	0.43
1:AA:1838:C:H2'	1:AA:1898:U:O4	2.18	0.43
1:AA:2080:G:H2'	1:AA:2081:C:C6	2.54	0.43
1:AA:2293:C:C4	1:AA:2294:C:C5	3.07	0.43
1:AA:2402:C:H41	1:AA:2416:C:H1'	1.83	0.43
1:AA:2576:G:O2'	1:AA:2579:C:OP2	2.25	0.43
1:AA:2668:G:O2'	1:AA:2669:G:H5'	2.19	0.43
1:AA:2735:G:H2'	1:AA:2736:G:C8	2.45	0.43
1:AA:2780:G:H2'	1:AA:2781:A:OP1	2.18	0.43
1:AA:560:C:H2'	1:AA:561:G:O4'	2.18	0.43
1:AA:646:A:N3	1:AA:646:A:H5'	2.32	0.43
1:AA:805:G:N2	1:AA:829:A:OP1	2.51	0.43
1:AA:836:G:H2'	1:AA:837:C:C6	2.53	0.43
1:AA:917:A:H2'	1:AA:918:A:C5'	2.49	0.43
3:AD:35:LYS:CD	3:AD:63:ARG:HB3	2.41	0.43
4:AE:37:ARG:HA	4:AE:42:ASP:CG	2.39	0.43
4:AE:52:LEU:HD12	4:AE:76:ARG:HD2	2.01	0.43
5:AF:21:ALA:C	5:AF:23:ASP:N	2.72	0.43
6:AG:151:ALA:O	6:AG:153:ARG:NH1	2.52	0.43
7:AH:35:VAL:O	7:AH:35:VAL:HG12	2.19	0.43
7:AH:45:VAL:O	7:AH:45:VAL:HG22	2.18	0.43
7:AH:7:LEU:HD12	7:AH:7:LEU:C	2.37	0.43
8:AK:8:PRO:CD	8:AK:15:VAL:HG23	2.45	0.43
11:AO:96:THR:O	11:AO:100:LEU:HD23	2.19	0.43
12:AP:52:VAL:C	12:AP:55:VAL:HG12	2.38	0.43
12:AP:56:ARG:O	12:AP:57:HIS:C	2.57	0.43
1:AA:912:C:OP1	12:AP:9:TYR:HE2	2.02	0.43
21:AV:177:PRO:C	21:AV:178:GLU:CG	2.85	0.43
21:AV:52:SER:C	21:AV:54:HIS:H	2.22	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:AW:33:MET:HG2	24:AW:37:PHE:HE1	1.82	0.43
53:B1:56:U:HO2'	53:B1:57:U:P	2.41	0.43
31:BA:119:A:C2	31:BA:240:C:C5	3.07	0.43
31:BA:1225:A:H5''	31:BA:1226:C:OP2	2.18	0.43
31:BA:1233:G:H2'	31:BA:1234:C:C6	2.53	0.43
31:BA:1349:A:O2'	31:BA:1350:A:H5'	2.19	0.43
31:BA:1512:U:C2	31:BA:1513:A:C8	3.07	0.43
31:BA:191(B):G:H2'	31:BA:191(C):G:O4'	2.19	0.43
31:BA:243:A:H62	31:BA:281:G:C2'	2.32	0.43
31:BA:437:U:O5'	31:BA:437:U:H6	2.02	0.43
31:BA:437:U:H2'	31:BA:438:G:O4'	2.19	0.43
31:BA:501:C:H1'	31:BA:549:C:H1'	2.00	0.43
31:BA:572:A:H5''	31:BA:917:G:H4'	2.01	0.43
31:BA:605:U:O2'	31:BA:606:G:H5'	2.18	0.43
32:BE:171:ALA:C	32:BE:173:ALA:H	2.21	0.43
32:BE:178:ARG:CG	32:BE:178:ARG:HH11	2.31	0.43
32:BE:214:ILE:C	32:BE:216:SER:N	2.72	0.43
32:BE:30:ARG:HB2	32:BE:46:LYS:HZ3	1.83	0.43
32:BE:97:TRP:CZ3	32:BE:176:GLU:OE2	2.72	0.43
33:BF:151:VAL:HG12	33:BF:152:ILE:N	2.33	0.43
35:BH:137:GLU:OE1	35:BH:141:GLN:NE2	2.52	0.43
35:BH:110:LEU:HD21	35:BH:139:LEU:HD21	2.01	0.43
39:BL:15:ALA:HB2	39:BL:65:VAL:HB	1.99	0.43
39:BL:46:ALA:HA	39:BL:78:LYS:CB	2.48	0.43
39:BL:40:LEU:CD1	39:BL:74:ILE:HD11	2.45	0.43
43:BP:64:TRP:HD1	43:BP:64:TRP:O	2.02	0.43
31:BA:994:A:O2'	44:BQ:8:GLU:HG3	2.19	0.43
49:BV:36:ARG:HB2	49:BV:72:GLY:N	2.34	0.43
50:BW:67:ALA:O	50:BW:68:LYS:C	2.57	0.43
53:C1:51:U:O5'	53:C1:51:U:H6	2.02	0.43
54:CA:1133:G:N1	54:CA:1142:G:C6	2.87	0.43
54:CA:1171:G:O2'	54:CA:1172:C:H5'	2.19	0.43
54:CA:1200:C:O2'	54:CA:1201:A:P	2.77	0.43
54:CA:1297:C:O2'	37:CJ:114:ARG:NH2	2.51	0.43
54:CA:280:C:N3	47:CT:39:SER:OG	2.50	0.43
54:CA:310:G:OP2	46:CS:27:LYS:HE3	2.19	0.43
54:CA:464:G:O6	54:CA:466:C:C5'	2.67	0.43
54:CA:518:C:C5	54:CA:530:G:N7	2.86	0.43
54:CA:644:G:O2'	54:CA:645:C:H5'	2.18	0.43
54:CA:801:U:H2'	54:CA:802:A:O5'	2.19	0.43
54:CA:953:G:O2'	54:CA:954:G:H5'	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:CD:44:G:O2'	52:CD:45:U:OP1	2.29	0.43
33:CF:28:GLN:O	33:CF:29:TYR:C	2.58	0.43
34:CG:100:ARG:CZ	34:CG:137:SER:HA	2.48	0.43
35:CH:103:GLY:O	35:CH:107:ARG:HB3	2.19	0.43
36:CI:2:ARG:HD2	36:CI:4:TYR:OH	2.19	0.43
36:CI:89:MET:HG2	36:CI:91:VAL:HG23	2.01	0.43
54:CA:878:G:C5'	38:CK:89:PRO:HG2	2.48	0.43
42:CO:28:LYS:HB3	42:CO:30:ALA:HB2	2.01	0.43
43:CP:88:ARG:HD3	43:CP:98:VAL:HG11	1.99	0.43
45:CR:66:LEU:O	45:CR:69:TYR:HB3	2.18	0.43
54:CA:265:G:H5'	47:CT:64:PRO:O	2.19	0.43
47:CT:62:SER:HB3	47:CT:72:ARG:HH21	1.84	0.43
48:CU:26:LEU:CD2	48:CU:42:ARG:HD2	2.49	0.43
55:DA:1083:U:O5'	57:DY:41:ARG:CD	2.66	0.43
55:DA:1495:A:H1'	55:DA:1579:A:H5''	2.00	0.43
55:DA:1914:C:H3'	55:DA:1914:C:O2	2.19	0.43
55:DA:2285:C:C5	28:D6:27:LYS:NZ	2.86	0.43
55:DA:271(C):U:H2'	55:DA:271:G:OP1	2.18	0.43
55:DA:33:U:HO2'	55:DA:34:C:P	2.42	0.43
55:DA:464:U:O2'	55:DA:465:G:H5'	2.19	0.43
55:DA:521:G:H2'	55:DA:522:G:C8	2.54	0.43
55:DA:554:U:HO2'	55:DA:556:G:H8	1.61	0.43
55:DA:779:U:P	3:DD:49:ILE:HG12	2.59	0.43
5:DF:32:LEU:CD2	5:DF:108:LYS:HB3	2.48	0.43
6:DG:106:LEU:O	6:DG:110:ALA:HB3	2.19	0.43
6:DG:68:PRO:HG2	6:DG:90:LEU:CD1	2.49	0.43
7:DH:144:VAL:O	7:DH:148:ILE:HG12	2.19	0.43
8:DK:79:ILE:HD13	8:DK:79:ILE:HA	1.87	0.43
58:DL:106:GLU:CG	58:DL:109:LYS:HD2	2.49	0.43
58:DL:82:ALA:HB1	58:DL:99:ILE:HD13	2.01	0.43
58:DL:9:LYS:N	58:DL:9:LYS:HZ3	2.16	0.43
9:DM:15:LEU:HB3	9:DM:136:GLU:HB3	2.01	0.43
12:DP:32:TYR:OH	12:DP:111:GLU:HB3	2.19	0.43
12:DP:74:TYR:CE2	12:DP:91:GLU:HG3	2.53	0.43
14:DQ:107:GLU:H	14:DQ:110:LEU:CD1	2.19	0.43
14:DQ:95:HIS:O	14:DQ:96:GLY:C	2.56	0.43
15:DR:34:VAL:HG12	15:DR:35:LYS:H	1.81	0.43
10:DN:104:ARG:NH1	15:DR:36:GLU:OE1	2.52	0.43
18:DS:14:PRO:HG2	18:DS:78:GLU:CB	2.47	0.43
21:DV:120:ILE:HB	21:DV:171:ILE:CA	2.48	0.43
21:DV:30:ASN:ND2	21:DV:32:HIS:CD2	2.87	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:DW:12:GLU:O	24:DW:16:LEU:HD23	2.19	0.43
57:DY:70:GLU:C	57:DY:113:GLN:HB2	2.39	0.43
57:DY:135:ARG:HG2	56:DJ:10:GLU:HB2	2.00	0.43
57:DY:13:LEU:CD1	57:DY:13:LEU:O	2.65	0.43
55:DA:1082:U:C5'	57:DY:45:LYS:O	2.59	0.43
13:A0:41:ALA:O	13:A0:44:LEU:N	2.51	0.43
13:A0:96:ARG:NH1	13:A0:96:ARG:HB3	2.33	0.43
22:A3:48:GLY:H	22:A3:51:VAL:HB	1.82	0.43
26:A4:17:GLY:HA3	26:A4:34:GLU:O	2.18	0.43
26:A4:69:LYS:C	26:A4:69:LYS:CD	2.86	0.43
1:AA:2347:C:H4'	28:A6:39:TYR:CZ	2.53	0.43
29:A7:19:ARG:CG	29:A7:19:ARG:NH1	2.77	0.43
28:A6:25:LYS:CA	30:A8:34:TRP:CH2	3.02	0.43
1:AA:1055:G:N2	1:AA:1085:A:O2'	2.51	0.43
1:AA:1085:A:O2'	1:AA:1104:C:O2	2.36	0.43
1:AA:1179:C:H2'	1:AA:1180:C:C5'	2.49	0.43
1:AA:1417:C:O2'	1:AA:1418:G:H5'	2.19	0.43
1:AA:1475:G:H5'	1:AA:1476:C:OP2	2.18	0.43
1:AA:2517:C:O2'	1:AA:2518:A:P	2.76	0.43
1:AA:2748:A:N1	1:AA:2749:A:C6	2.87	0.43
1:AA:2776:A:O2'	1:AA:2781:A:C4'	2.66	0.43
1:AA:355:G:H2'	1:AA:356:G:O4'	2.19	0.43
1:AA:603:A:O2'	1:AA:604:G:P	2.77	0.43
1:AA:663:G:C6	1:AA:664:C:C4	3.07	0.43
1:AA:959:A:C6	1:AA:960:A:C2	3.07	0.43
2:AB:34:U:C4	2:AB:44:G:H2'	2.54	0.43
1:AA:1902:C:H5'	3:AD:246:PRO:HD3	2.01	0.43
4:AE:64:LYS:C	4:AE:66:HIS:N	2.72	0.43
4:AE:63:LEU:CD2	4:AE:66:HIS:HB2	2.49	0.43
4:AE:73:GLU:HG2	4:AE:74:PRO:CD	2.48	0.43
5:AF:40:GLN:HE21	5:AF:40:GLN:HA	1.84	0.43
6:AG:5:VAL:HA	26:A4:25:TYR:OH	2.19	0.43
7:AH:4:ILE:HD12	7:AH:6:ARG:CD	2.49	0.43
8:AK:89:TYR:HA	54:CA:357:G:O2'	2.18	0.43
1:AA:1012:U:O2	9:AM:25:ARG:NH1	2.52	0.43
9:AM:1:MET:O	9:AM:2:LYS:HB2	2.18	0.43
1:AA:1242:A:C6	11:AO:4:SER:HB2	2.53	0.43
19:AT:35:THR:H	19:AT:38:GLU:HG2	1.84	0.43
21:AV:63:ASP:OD2	21:AV:65:GLN:HG2	2.19	0.43
31:BA:1238:A:H62	31:BA:1301:U:H3	1.66	0.43
31:BA:1517:G:H2'	31:BA:1518:A:C8	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:164:U:H2'	31:BA:165:C:C5	2.53	0.43
31:BA:210:U:O2	31:BA:210:U:C2'	2.59	0.43
31:BA:291:C:O2'	31:BA:292:G:H5'	2.19	0.43
31:BA:309:G:H1'	31:BA:608:A:C2	2.54	0.43
31:BA:452:A:H2'	31:BA:453:A:C8	2.54	0.43
52:BD:29:G:O2'	52:BD:30:G:H5'	2.18	0.43
32:BE:216:SER:O	32:BE:218:ALA:N	2.51	0.43
33:BF:63:ASN:CB	33:BF:98:ASN:HB3	2.48	0.43
34:BG:20:TYR:N	34:BG:20:TYR:CD1	2.86	0.43
38:BK:26:VAL:HG23	38:BK:27:PRO:HD2	2.01	0.43
26:A4:50:VAL:HG11	43:BP:65:LYS:HB3	2.01	0.43
46:BS:55:ARG:O	46:BS:56:ALA:C	2.57	0.43
48:BU:23:LYS:CA	48:BU:26:LEU:HD11	2.44	0.43
49:BV:63:THR:CG2	49:BV:66:MET:HG3	2.49	0.43
50:BW:88:VAL:O	50:BW:91:LEU:HB2	2.19	0.43
54:CA:1355:G:H2'	54:CA:1356:G:H8	1.84	0.43
54:CA:1366:C:H2'	54:CA:1367:C:C6	2.54	0.43
54:CA:366:C:HO2'	54:CA:367:U:P	2.40	0.43
54:CA:397:A:H5'	54:CA:398:C:OP1	2.19	0.43
54:CA:427:U:H1'	54:CA:541:G:OP1	2.18	0.43
54:CA:452:A:C6	54:CA:453:A:C6	3.06	0.43
54:CA:555:C:OP1	42:CO:20:LYS:HE2	2.18	0.43
54:CA:575:G:H4'	54:CA:576:G:O5'	2.18	0.43
54:CA:736:C:H2'	54:CA:737:A:H8	1.84	0.43
52:CB:9:A:O4'	52:CB:46:G:O4'	2.37	0.43
52:CC:47:U:C2'	52:CC:47:U:O2	2.59	0.43
33:CF:64:VAL:CG1	33:CF:66:VAL:HG23	2.47	0.43
34:CG:124:GLY:C	34:CG:126:ILE:H	2.22	0.43
34:CG:19:LEU:CD2	34:CG:19:LEU:N	2.82	0.43
34:CG:20:TYR:CD1	34:CG:20:TYR:N	2.87	0.43
34:CG:39:PRO:O	34:CG:44:GLY:HA3	2.19	0.43
36:CI:62:TRP:CD1	48:CU:35:ARG:CZ	3.02	0.43
36:CI:83:ASP:C	36:CI:85:VAL:H	2.21	0.43
36:CI:4:TYR:CD1	36:CI:92:LYS:HA	2.54	0.43
39:CL:112:LYS:HD3	39:CL:112:LYS:C	2.39	0.43
39:CL:9:ARG:O	39:CL:9:ARG:HG2	2.18	0.43
41:CN:120:ARG:HH12	41:CN:126:ARG:NH2	2.17	0.43
44:CQ:13:THR:O	44:CQ:14:PRO:C	2.56	0.43
54:CA:452:A:C4'	46:CS:72:ARG:NH2	2.76	0.43
47:CT:68:ARG:N	47:CT:70:ARG:NH1	2.66	0.43
49:CV:83:HIS:O	49:CV:87:ALA:N	2.41	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
50:CW:75:ASN:N	50:CW:75:ASN:OD1	2.51	0.43
22:D3:31:VAL:HB	22:D3:35:ASN:HD22	1.83	0.43
12:DP:80:GLU:CD	22:D3:4:LYS:HZ2	2.22	0.43
27:D5:48:GLU:HG3	27:D5:59:GLU:HG3	2.01	0.43
28:D6:35:GLU:OE2	28:D6:51:GLU:OE2	2.37	0.43
28:D6:36:LEU:HD23	28:D6:36:LEU:O	2.19	0.43
55:DA:2347:C:H4'	28:D6:39:TYR:CE2	2.54	0.43
55:DA:1081:U:H3'	55:DA:1082:U:O4'	2.19	0.43
55:DA:1191:G:H2'	55:DA:1192:G:O4'	2.19	0.43
55:DA:1212:G:N2	55:DA:1236:G:O2'	2.51	0.43
55:DA:1397:U:O2'	55:DA:1398:C:OP1	2.33	0.43
55:DA:142:G:H1'	19:DT:37:THR:CG2	2.48	0.43
55:DA:1380:G:N2	55:DA:1570:A:C2	2.87	0.43
55:DA:1773:A:N7	55:DA:1829:A:H1'	2.34	0.43
55:DA:1893:C:C2'	55:DA:1894:C:H5'	2.49	0.43
55:DA:1906:G:C4	55:DA:1929:G:C2	3.07	0.43
55:DA:1678:G:H22	55:DA:1989:G:H1	1.65	0.43
55:DA:2163:C:OP1	55:DA:2172:U:C5	2.72	0.43
55:DA:226:G:C2'	55:DA:227:A:OP2	2.67	0.43
55:DA:2667:C:H2'	55:DA:2668:G:O4'	2.18	0.43
55:DA:447:A:H5''	55:DA:448:U:OP1	2.19	0.43
55:DA:500:G:N2	55:DA:502:A:H2'	2.33	0.43
55:DA:654(C):G:C2'	55:DA:654(D):G:C8	3.01	0.43
55:DA:856:C:H5''	55:DA:856:C:C6	2.54	0.43
3:DD:13:ARG:CZ	3:DD:16:MET:HE1	2.49	0.43
3:DD:228:PRO:HD3	3:DD:234:GLY:O	2.18	0.43
4:DE:63:LEU:HG	4:DE:64:LYS:N	2.33	0.43
43:CP:7:VAL:CB	6:DG:115:ARG:HH12	2.31	0.43
2:DB:45:A:H1'	6:DG:95:ARG:NH2	2.33	0.43
7:DH:127:GLU:CG	7:DH:128:PRO:CD	2.96	0.43
57:DY:135:ARG:HE	56:DJ:18:LEU:HD13	1.84	0.43
9:DM:30:ILE:O	9:DM:34:LEU:HD22	2.18	0.43
10:DN:35:VAL:HA	10:DN:62:VAL:HG12	2.01	0.43
11:DO:127:ALA:O	11:DO:147:LEU:HD23	2.19	0.43
11:DO:75:ILE:O	11:DO:75:ILE:HG12	2.19	0.43
14:DQ:88:ASP:CG	14:DQ:89:ARG:N	2.71	0.43
15:DR:124:ASP:O	15:DR:128:GLU:N	2.51	0.43
15:DR:40:THR:HG22	15:DR:40:THR:O	2.17	0.43
15:DR:62:THR:HB	15:DR:75:ILE:HG12	2.00	0.43
55:DA:2847:U:OP1	15:DR:98:LYS:HD3	2.19	0.43
20:DU:75:ILE:HB	20:DU:80:GLY:HA2	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:DU:81:LYS:HZ3	20:DU:98:VAL:CG1	2.29	0.43
57:DY:136:ALA:O	57:DY:139:VAL:CB	2.62	0.43
57:DY:29:TYR:CD2	57:DY:32:LEU:CD2	3.01	0.43
57:DY:58:LEU:CA	57:DY:62:ALA:CB	2.97	0.43
13:A0:44:LEU:HD22	13:A0:44:LEU:O	2.18	0.43
17:A2:95:LEU:HD22	17:A2:97:LYS:HE3	2.01	0.43
1:AA:2356:C:H4'	22:A3:20:ARG:HG3	2.00	0.43
1:AA:1073:A:OP2	1:AA:1094:U:O4	2.37	0.43
1:AA:1543:A:H1'	1:AA:1545:A:C1'	2.47	0.43
1:AA:1554:A:C2	1:AA:1634:A:N6	2.85	0.43
1:AA:1428:C:N4	1:AA:1570:A:OP2	2.36	0.43
1:AA:1332:G:N2	1:AA:1610:A:C8	2.87	0.43
1:AA:1625:C:H2'	1:AA:1626:G:O4'	2.19	0.43
1:AA:1936:A:H5''	1:AA:1937:A:O5'	2.19	0.43
1:AA:1944:U:H1'	1:AA:1955:U:C4'	2.49	0.43
1:AA:2032:G:OP2	1:AA:2454:G:O2'	2.34	0.43
1:AA:2462:U:H2'	1:AA:2463:C:O4'	2.19	0.43
1:AA:2602:A:H4'	1:AA:2603:G:H5'	2.00	0.43
1:AA:2642:G:C5'	9:AM:78:TYR:CD2	3.02	0.43
1:AA:588:U:C2	1:AA:589:C:C5	3.06	0.43
1:AA:654(I):C:O2	1:AA:654(I):C:O2'	2.34	0.43
1:AA:605:C:H1'	1:AA:657:U:O2'	2.19	0.43
3:AD:268:ARG:HD3	3:AD:268:ARG:O	2.19	0.43
3:AD:27:THR:C	3:AD:29:PRO:HD2	2.38	0.43
4:AE:146:THR:HA	4:AE:147:PRO:C	2.38	0.43
4:AE:16:ARG:NH1	4:AE:16:ARG:HG3	2.34	0.43
4:AE:179:GLU:HG3	4:AE:181:LEU:CD2	2.48	0.43
4:AE:25:VAL:HG22	4:AE:183:LEU:CD1	2.49	0.43
4:AE:29:GLY:N	4:AE:51:PHE:HE1	2.16	0.43
4:AE:3:GLY:O	4:AE:198:VAL:O	2.36	0.43
9:AM:97:ARG:HH11	9:AM:97:ARG:HG2	1.84	0.43
12:AP:60:ARG:O	12:AP:61:GLY:O	2.37	0.43
14:AQ:25:ARG:NH1	14:AQ:25:ARG:CB	2.78	0.43
14:AQ:61:ASN:O	14:AQ:62:LYS:C	2.57	0.43
20:AU:62:GLU:CG	20:AU:63:LYS:H	2.31	0.43
21:AV:150:LEU:HD23	21:AV:154:ASP:HB2	2.00	0.43
24:AW:39:ALA:O	24:AW:42:GLY:HA2	2.19	0.43
24:AW:64:LEU:O	24:AW:64:LEU:HD23	2.19	0.43
25:AX:54:VAL:CG1	25:AX:55:ARG:H	2.31	0.43
23:AZ:66:HIS:C	23:AZ:68:PRO:HD2	2.38	0.43
33:BF:164:ARG:NH2	53:B1:55:U:O4	2.50	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:1053:G:N7	31:BA:1200:C:C5'	2.82	0.43
31:BA:1095:U:H2'	31:BA:1096:C:C6	2.54	0.43
31:BA:1122:U:C2	31:BA:1123:A:C8	3.06	0.43
31:BA:1133:G:N1	31:BA:1142:G:C6	2.87	0.43
31:BA:115:G:H4'	31:BA:116:A:O5'	2.18	0.43
31:BA:1294:G:O2'	31:BA:1295:G:H5'	2.19	0.43
31:BA:1303:C:H2'	31:BA:1304:G:H5'	2.00	0.43
31:BA:216:G:O2'	31:BA:217:C:O4'	2.36	0.43
31:BA:39:G:N7	31:BA:547:A:C8	2.87	0.43
31:BA:642:A:N3	38:BK:113:SER:OG	2.38	0.43
31:BA:669:U:H2'	31:BA:670:G:C8	2.54	0.43
52:BC:40:C:O2	52:BC:40:C:H2'	2.18	0.43
32:BE:121:LEU:O	32:BE:127:ILE:HG12	2.19	0.43
32:BE:158:LEU:N	32:BE:158:LEU:HD12	2.34	0.43
32:BE:71:VAL:HB	32:BE:164:VAL:HG22	2.01	0.43
32:BE:25:ASN:HB2	32:BE:191:ASP:O	2.19	0.43
32:BE:5:ILE:O	32:BE:5:ILE:HG23	2.19	0.43
33:BF:126:ARG:HB2	33:BF:128:PHE:HD1	1.84	0.43
33:BF:157:ILE:C	33:BF:159:GLY:H	2.21	0.43
34:BG:182:LYS:HG3	34:BG:182:LYS:O	2.19	0.43
34:BG:20:TYR:HB3	34:BG:27:TYR:CD1	2.54	0.43
39:BL:45:ALA:O	39:BL:48:GLU:HB2	2.18	0.43
40:BM:12:ASP:C	40:BM:14:LYS:H	2.22	0.43
31:BA:972:C:H4'	40:BM:57:LYS:CG	2.48	0.43
44:BQ:46:GLU:O	44:BQ:47:LEU:C	2.57	0.43
47:BT:99:SER:O	47:BT:101:ARG:N	2.49	0.43
48:BU:56:THR:HG21	48:BU:63:GLN:OE1	2.19	0.43
49:BV:44:MET:HA	49:BV:47:HIS:HD2	1.80	0.43
54:CA:1174:G:H2'	54:CA:1175:G:H8	1.84	0.43
54:CA:1360:A:H2'	54:CA:1361:G:C8	2.53	0.43
54:CA:1363:A:C1'	54:CA:1365:G:N7	2.65	0.43
54:CA:428:G:O2'	54:CA:429:U:P	2.77	0.43
54:CA:442:C:H2'	54:CA:442:C:O2	2.18	0.43
54:CA:675:A:H1'	41:CN:116:HIS:CD2	2.54	0.43
54:CA:89:U:H2'	54:CA:90:C:O5'	2.17	0.43
52:CD:13:C:C2'	52:CD:14:A:H5'	2.49	0.43
32:CE:158:LEU:O	32:CE:158:LEU:HD12	2.19	0.43
34:CG:173:TRP:CE3	34:CG:193:ASP:HB3	2.54	0.43
35:CH:32:VAL:CG1	35:CH:33:VAL:N	2.82	0.43
35:CH:48:ALA:HB2	35:CH:57:LYS:HD3	2.00	0.43
39:CL:57:GLY:O	39:CL:58:HIS:C	2.56	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1149:C:P	39:CL:9:ARG:NH2	2.91	0.43
41:CN:64:ALA:O	41:CN:65:ALA:C	2.56	0.43
42:CO:89:ARG:HH11	42:CO:89:ARG:HB2	1.84	0.43
42:CO:43:VAL:HG21	42:CO:93:LEU:HD22	2.00	0.43
44:CQ:19:ARG:O	44:CQ:21:TYR:HD1	2.01	0.43
44:CQ:42:ILE:HG22	44:CQ:46:GLU:CG	2.49	0.43
46:CS:71:ARG:CB	46:CS:71:ARG:HH11	2.32	0.43
47:CT:84:LEU:C	47:CT:86:GLU:N	2.72	0.43
49:CV:83:HIS:O	49:CV:86:GLU:CA	2.67	0.43
55:DA:1162:G:H1'	17:D2:23:GLU:OE2	2.19	0.43
22:D3:53:MET:HB3	22:D3:59:LEU:HD23	2.01	0.43
22:D3:7:LEU:HD22	22:D3:7:LEU:N	2.34	0.43
6:DG:179:PRO:HG3	26:D4:38:LYS:HZ1	1.84	0.43
27:D5:58:LEU:HD13	27:D5:60:VAL:CG1	2.49	0.43
55:DA:1001:A:H2'	55:DA:1002:G:O4'	2.18	0.43
55:DA:1324:G:C2'	55:DA:1325:G:H5'	2.49	0.43
55:DA:1359:A:C2	55:DA:1373:A:C4	3.07	0.43
55:DA:1879:C:C2'	55:DA:1880:C:H5''	2.48	0.43
55:DA:196:A:P	11:DO:46:LYS:HZ1	2.41	0.43
55:DA:530:G:C6	55:DA:2022:U:H5''	2.53	0.43
55:DA:2146:C:C5'	55:DA:2147:G:OP1	2.62	0.43
55:DA:2166:G:C2'	55:DA:2167:U:OP1	2.67	0.43
55:DA:2234:G:H2'	55:DA:2235:G:O4'	2.18	0.43
55:DA:2286:A:OP2	28:D6:28:ARG:HD3	2.19	0.43
55:DA:2572:A:O2'	55:DA:2573:C:OP2	2.36	0.43
55:DA:2707:G:H2'	55:DA:2708:G:H8	1.84	0.43
55:DA:454:A:H4'	55:DA:455:C:OP2	2.16	0.43
55:DA:2600:A:N7	3:DD:237:GLU:HG2	2.34	0.43
4:DE:24:THR:CB	4:DE:188:VAL:HG11	2.49	0.43
55:DA:2635:C:H5'	4:DE:77:ILE:CD1	2.49	0.43
4:DE:7:VAL:CG2	4:DE:8:LYS:N	2.57	0.43
4:DE:92:THR:C	4:DE:95:ILE:HD13	2.39	0.43
4:DE:92:THR:H	4:DE:95:ILE:HD11	1.83	0.43
7:DH:41:MET:CE	7:DH:64:LEU:HB2	2.49	0.43
58:DL:109:LYS:HA	58:DL:120:LEU:CD2	2.43	0.43
10:DN:106:LEU:HD23	10:DN:106:LEU:HA	1.73	0.43
15:DR:96:ARG:HB2	15:DR:96:ARG:HH11	1.81	0.43
20:DU:19:LYS:O	20:DU:20:TYR:CD1	2.72	0.43
20:DU:50:ARG:C	20:DU:52:SER:N	2.73	0.43
21:DV:151:HIS:CG	21:DV:169:GLU:O	2.72	0.43
57:DY:101:PRO:CD	57:DY:102:LYS:H	2.32	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:DY:138:LEU:HD21	56:DJ:22:GLN:CD	2.34	0.43
17:A2:71:LEU:CA	17:A2:86:GLY:CA	2.97	0.42
30:A8:32:LEU:HD23	30:A8:33:ASN:H	1.83	0.42
1:AA:1121:C:H2'	1:AA:1122:G:O4'	2.19	0.42
1:AA:1264:G:C2'	1:AA:1265:A:OP1	2.67	0.42
1:AA:1298:C:H2'	1:AA:1298:C:O2	2.20	0.42
1:AA:1355:G:C4	1:AA:1356:G:C8	3.06	0.42
1:AA:1544:C:O2'	1:AA:1545:A:OP1	2.28	0.42
1:AA:14:A:N6	1:AA:15:G:C2	2.87	0.42
1:AA:528:A:H2	1:AA:2042:A:H2'	1.78	0.42
1:AA:2107:C:H2'	1:AA:2108:C:O4'	2.19	0.42
1:AA:2113:U:C6	1:AA:2114:A:O4'	2.71	0.42
1:AA:214:G:H1'	1:AA:216:A:O2'	2.19	0.42
1:AA:858:U:HO2'	1:AA:2268:A:H1'	1.84	0.42
1:AA:2393:A:H3'	1:AA:2394:C:H6	1.83	0.42
1:AA:247:G:H4'	1:AA:386:G:C6	2.53	0.42
1:AA:2534:A:C2	1:AA:2535:G:H1'	2.54	0.42
1:AA:2720:U:H2'	1:AA:2721:A:C8	2.54	0.42
1:AA:2728:U:O2'	1:AA:2729:G:H5'	2.19	0.42
1:AA:2748:A:N7	1:AA:2757:A:N1	2.67	0.42
1:AA:2780:G:O2'	1:AA:2781:A:OP1	2.36	0.42
1:AA:2810:A:H61	1:AA:2891:G:C2'	2.32	0.42
1:AA:312:G:P	1:AA:312:G:H8	2.42	0.42
1:AA:39:C:H2'	1:AA:40:C:H6	1.83	0.42
1:AA:404:C:O2'	1:AA:405:U:C5'	2.65	0.42
1:AA:630:G:N2	1:AA:633:A:OP2	2.45	0.42
1:AA:74:A:C8	1:AA:74:A:OP2	2.72	0.42
1:AA:852:G:O2'	1:AA:853:G:H5'	2.19	0.42
1:AA:916:G:O2'	1:AA:917:A:H5''	2.19	0.42
2:AB:78:A:H61	2:AB:98:G:H1'	1.84	0.42
3:AD:169:GLU:HG2	3:AD:174:ILE:CD1	2.44	0.42
3:AD:222:ARG:O	3:AD:225:ALA:HB3	2.18	0.42
3:AD:270:ILE:HG22	3:AD:271:ILE:N	2.34	0.42
4:AE:179:GLU:O	4:AE:180:ASN:HB2	2.18	0.42
5:AF:79:GLY:CA	5:AF:86:GLY:HA2	2.42	0.42
5:AF:8:GLN:NE2	5:AF:127:GLU:HB3	2.34	0.42
6:AG:129:GLY:O	6:AG:130:ASN:CG	2.58	0.42
9:AM:35:ARG:NH2	9:AM:42:TRP:HH2	2.16	0.42
9:AM:89:LYS:HZ2	9:AM:89:LYS:HB3	1.83	0.42
11:AO:112:LEU:HD13	11:AO:112:LEU:C	2.39	0.42
12:AP:109:VAL:CG1	12:AP:110:THR:N	2.82	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AP:87:LYS:O	12:AP:88:GLY:O	2.37	0.42
14:AQ:20:ARG:HE	14:AQ:21:THR:HA	1.83	0.42
14:AQ:39:ILE:O	14:AQ:47:THR:HG23	2.19	0.42
18:AS:40:ASN:O	18:AS:41:LYS:HG2	2.18	0.42
20:AU:12:THR:HG22	20:AU:26:LYS:HE2	1.97	0.42
21:AV:41:LEU:HD23	21:AV:41:LEU:C	2.38	0.42
24:AW:53:LEU:O	24:AW:56:GLN:N	2.51	0.42
23:AZ:6:GLU:OE1	23:AZ:60:PHE:HA	2.19	0.42
53:B1:51:U:H3'	53:B1:51:U:H6	1.83	0.42
31:BA:1269:A:C2	31:BA:1313:U:O4'	2.72	0.42
31:BA:1306:A:H2'	31:BA:1307:U:C6	2.54	0.42
31:BA:1478:C:H2'	31:BA:1479:C:H6	1.84	0.42
31:BA:197:A:O2'	31:BA:198:G:OP2	2.36	0.42
31:BA:624:C:H2'	31:BA:625:G:H8	1.83	0.42
32:BE:55:PHE:CD1	32:BE:221:LEU:CD2	3.02	0.42
33:BF:70:VAL:O	33:BF:106:VAL:HG23	2.17	0.42
34:BG:108:LEU:CB	34:BG:110:PHE:HE1	2.28	0.42
34:BG:20:TYR:HD2	34:BG:27:TYR:CD1	2.37	0.42
34:BG:90:GLY:O	34:BG:94:LEU:HG	2.18	0.42
40:BM:6:ILE:CA	40:BM:97:GLU:O	2.67	0.42
41:BN:21:ILE:HB	41:BN:84:VAL:HG12	2.01	0.42
41:BN:93:GLN:O	41:BN:97:ALA:HB2	2.18	0.42
43:BP:88:ARG:HH11	43:BP:88:ARG:HG2	1.83	0.42
49:BV:19:VAL:HG12	49:BV:20:LEU:N	2.34	0.42
49:BV:48:THR:HG22	49:BV:61:TYR:HD1	1.84	0.42
54:CA:1107:C:C4	54:CA:1108:G:C8	3.07	0.42
54:CA:1345:U:H5''	54:CA:1346:A:OP1	2.19	0.42
54:CA:1410:G:H2'	54:CA:1411:C:H6	1.84	0.42
54:CA:781:A:O2'	54:CA:1522:U:O2	2.35	0.42
54:CA:376:G:C2'	54:CA:377:G:H5'	2.49	0.42
54:CA:377:G:OP1	46:CS:3:LYS:NZ	2.49	0.42
54:CA:373:A:H61	54:CA:391:G:H1'	1.83	0.42
54:CA:624:C:H4'	46:CS:10:GLY:O	2.18	0.42
54:CA:719:C:O2'	48:CU:49:LYS:HB3	2.18	0.42
54:CA:878:G:H2'	54:CA:879:C:C6	2.54	0.42
54:CA:922:G:H4'	35:CH:20:GLN:HA	2.01	0.42
52:CD:2:C:C5'	52:CD:2:C:H6	2.24	0.42
52:CD:37:MIA:H112	52:CD:38:A:H1'	1.99	0.42
52:CD:72:C:C3'	52:CD:73:A:C5'	2.97	0.42
32:CE:28:PHE:CZ	32:CE:189:ASP:HA	2.54	0.42
33:CF:14:ILE:O	33:CF:15:THR:HB	2.18	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:CF:22:TRP:CZ2	44:CQ:54:PRO:HG2	2.53	0.42
33:CF:3:ASN:OD1	33:CF:3:ASN:N	2.52	0.42
33:CF:79:ARG:NH1	33:CF:79:ARG:CG	2.82	0.42
34:CG:132:ARG:NH1	34:CG:132:ARG:HG2	2.34	0.42
34:CG:163:GLU:O	34:CG:165:MET:N	2.52	0.42
34:CG:92:VAL:O	34:CG:96:LEU:CD2	2.67	0.42
35:CH:20:GLN:O	35:CH:21:ALA:C	2.57	0.42
35:CH:89:ILE:HG12	35:CH:91:LEU:CD1	2.49	0.42
38:CK:82:HIS:O	38:CK:137:VAL:HA	2.19	0.42
41:CN:30:VAL:O	41:CN:30:VAL:HG23	2.19	0.42
42:CO:83:VAL:CG2	42:CO:100:ILE:HG12	2.44	0.42
42:CO:61:THR:OG1	42:CO:62:SER:N	2.52	0.42
43:CP:29:ARG:HD3	43:CP:64:TRP:CE2	2.54	0.42
48:CU:66:LEU:HG	48:CU:70:ILE:CD1	2.49	0.42
13:D0:13:HIS:HE1	13:D0:15:SER:OG	2.01	0.42
17:D2:91:TYR:C	17:D2:91:TYR:HD1	2.21	0.42
22:D3:25:ARG:CD	22:D3:29:GLN:NE2	2.82	0.42
12:DP:81:VAL:HG23	22:D3:7:LEU:HD21	2.01	0.42
26:D4:16:CYS:SG	26:D4:17:GLY:N	2.92	0.42
26:D4:12:ALA:HB2	26:D4:29:PRO:HA	2.00	0.42
28:D6:17:LYS:HG3	28:D6:18:ARG:H	1.83	0.42
55:DA:1312:U:H4'	55:DA:1313:U:O5'	2.18	0.42
55:DA:1786:A:N1	55:DA:2606:C:C1'	2.82	0.42
55:DA:1910:G:O2'	55:DA:1911:U:H5'	2.19	0.42
55:DA:1982:C:H2'	55:DA:1982:C:O2	2.18	0.42
55:DA:919:G:N2	55:DA:2269:A:OP2	2.48	0.42
55:DA:2735:G:H2'	55:DA:2736:G:C8	2.53	0.42
55:DA:297:C:O2'	55:DA:298:G:H5'	2.19	0.42
55:DA:319:C:H2'	55:DA:320:A:C8	2.54	0.42
55:DA:37:C:H4'	55:DA:451:C:OP1	2.18	0.42
55:DA:479:A:H4'	55:DA:480:A:H5'	2.01	0.42
55:DA:781:A:H2'	55:DA:1777:U:O2'	2.19	0.42
55:DA:91:A:H2'	55:DA:92:G:O5'	2.19	0.42
55:DA:932:G:H4'	55:DA:933:A:O5'	2.19	0.42
5:DF:117:ARG:HG3	5:DF:117:ARG:HH11	1.83	0.42
5:DF:178:PRO:HB3	5:DF:198:ALA:HB1	2.00	0.42
6:DG:47:LYS:NZ	6:DG:81:LYS:HG2	2.34	0.42
56:DJ:7:ARG:O	56:DJ:8:ILE:HB	2.19	0.42
8:DK:72:LEU:CD1	8:DK:107:VAL:HG11	2.43	0.42
8:DK:112:LYS:HG2	8:DK:112:LYS:H	1.67	0.42
8:DK:121:LYS:O	8:DK:122:GLU:HB2	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:DK:93:THR:HG22	8:DK:119:PRO:CB	2.39	0.42
10:DN:7:TYR:HE1	10:DN:20:MET:CE	2.27	0.42
10:DN:87:ILE:CD1	10:DN:91:LEU:HD12	2.48	0.42
14:DQ:18:ILE:HD12	14:DQ:88:ASP:HA	2.00	0.42
14:DQ:69:VAL:HG13	14:DQ:101:LEU:CD2	2.49	0.42
15:DR:110:ILE:CG2	15:DR:111:ARG:NH1	2.82	0.42
15:DR:19:LEU:HA	15:DR:20:PRO:HD3	1.79	0.42
15:DR:3:ARG:O	15:DR:5:ALA:N	2.51	0.42
21:DV:190:GLU:H	21:DV:193:GLU:H	1.66	0.42
57:DY:63:LEU:HD23	57:DY:63:LEU:N	2.24	0.42
57:DY:7:VAL:CG1	57:DY:8:GLU:N	2.44	0.42
57:DY:91:LYS:CA	57:DY:94:VAL:HB	2.46	0.42
16:A1:50:ARG:HG2	16:A1:53:ARG:NH2	2.33	0.42
17:A2:76:LYS:HD2	17:A2:80:GLN:O	2.19	0.42
26:A4:24:THR:O	26:A4:25:TYR:CB	2.67	0.42
26:A4:18:CYS:HB3	26:A4:36:CYS:HB2	1.82	0.42
26:A4:56:VAL:CG1	26:A4:57:GLU:N	2.81	0.42
27:A5:33:CYS:HB2	27:A5:36:CYS:HB2	2.00	0.42
28:A6:13:CYS:O	28:A6:14:THR:HB	2.19	0.42
30:A8:34:TRP:C	30:A8:36:LYS:H	2.21	0.42
1:AA:1213:A:O2'	1:AA:1214:A:H5'	2.19	0.42
1:AA:125:G:OP2	29:A7:19:ARG:HD3	2.19	0.42
1:AA:1312:U:H3'	19:AT:63:LYS:HZ2	1.84	0.42
1:AA:1419:A:O2'	1:AA:1420:U:C6	2.72	0.42
1:AA:1554:A:H5'	1:AA:1555:G:OP2	2.19	0.42
1:AA:1771:C:O5'	1:AA:1771:C:H6	2.02	0.42
1:AA:1941:C:N4	1:AA:1965:C:H5'	2.34	0.42
1:AA:2287:A:C2'	1:AA:2288:A:O5'	2.67	0.42
1:AA:26:G:C6	1:AA:27:G:N1	2.88	0.42
1:AA:460:A:H3'	1:AA:461:C:C6	2.54	0.42
1:AA:654(A):A:N1	1:AA:654(T):A:N1	2.67	0.42
1:AA:895:U:C3'	1:AA:895:U:C2	3.02	0.42
1:AA:846:C:C4	1:AA:930:U:C5	3.07	0.42
1:AA:98:G:O2'	1:AA:99:U:H5''	2.19	0.42
3:AD:137:PRO:O	3:AD:138:VAL:C	2.58	0.42
3:AD:158:ALA:N	3:AD:161:THR:OG1	2.51	0.42
3:AD:34:VAL:O	3:AD:35:LYS:HG3	2.19	0.42
4:AE:134:ILE:CG1	4:AE:134:ILE:O	2.67	0.42
8:AK:5:LEU:HD21	8:AK:12:LEU:HB3	2.00	0.42
8:AK:82:ARG:HH11	8:AK:146:ALA:HB2	1.84	0.42
9:AM:38:HIS:CG	9:AM:39:ARG:N	2.87	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:AM:45:ASN:ND2	9:AM:45:ASN:N	2.64	0.42
14:AQ:30:ARG:HG3	14:AQ:97:ARG:NH2	2.34	0.42
20:AU:97:ARG:N	20:AU:97:ARG:HD3	2.20	0.42
53:B1:43:U:C6	53:B1:43:U:H5'	2.46	0.42
53:B1:50:U:C3'	53:B1:51:U:H5'	2.50	0.42
31:BA:1148:U:O2'	31:BA:1149:C:H5'	2.19	0.42
31:BA:1327:C:OP1	51:BX:12:LYS:NZ	2.52	0.42
31:BA:1354:C:O2'	31:BA:1355:G:H5'	2.18	0.42
31:BA:563:A:H1'	31:BA:566:G:O2'	2.19	0.42
31:BA:676:A:C4	31:BA:677:U:C5	3.07	0.42
52:BD:54:U:H2'	52:BD:55:U:O4'	2.20	0.42
32:BE:102:LEU:CD1	32:BE:102:LEU:N	2.82	0.42
33:BF:135:LYS:O	33:BF:139:GLN:HB2	2.18	0.42
33:BF:35:GLU:HG2	33:BF:39:ILE:HD11	2.01	0.42
34:BG:29:PRO:HD2	34:BG:30:LYS:CE	2.49	0.42
34:BG:34:GLU:C	34:BG:35:ARG:CG	2.84	0.42
35:BH:78:HIS:ND1	38:BK:107:LEU:CD1	2.81	0.42
36:BI:54:LYS:HA	36:BI:54:LYS:HZ2	1.84	0.42
40:BM:98:ILE:O	40:BM:99:LYS:HB2	2.19	0.42
31:BA:1329:A:OP1	43:BP:26:GLY:HA3	2.19	0.42
45:BR:3:ILE:CB	45:BR:38:ARG:HH21	2.32	0.42
48:BU:37:VAL:O	48:BU:38:GLU:C	2.57	0.42
48:BU:53:ARG:NH1	48:BU:60:ALA:HB2	2.34	0.42
50:BW:43:LEU:O	50:BW:47:GLY:N	2.51	0.42
31:BA:186:C:C1'	50:BW:81:LYS:HZ1	2.32	0.42
50:BW:96:GLY:O	50:BW:97:ALA:O	2.37	0.42
31:BA:1353:G:H5''	51:BX:13:ILE:CG2	2.50	0.42
54:CA:1126:U:C5	54:CA:1127:G:C4	3.08	0.42
54:CA:1189:C:H4'	33:CF:10:PHE:CE1	2.55	0.42
54:CA:1284:C:H2'	54:CA:1285:A:N7	2.34	0.42
54:CA:1443:G:C4'	54:CA:1446:A:OP2	2.66	0.42
54:CA:186(C):G:C6	54:CA:191(E):G:N1	2.87	0.42
54:CA:241:C:O2'	54:CA:242:C:H5'	2.19	0.42
54:CA:312:C:N4	54:CA:313:A:N6	2.68	0.42
54:CA:37:U:H2'	54:CA:38:G:C8	2.50	0.42
52:CB:4:C:C2	52:CB:5:G:C8	3.07	0.42
52:CB:70:G:H2'	52:CB:71:G:C8	2.54	0.42
52:CD:46:G:O3'	52:CD:47:U:H4'	2.18	0.42
33:CF:29:TYR:C	33:CF:29:TYR:HD2	2.23	0.42
34:CG:6:GLY:O	34:CG:7:PRO:C	2.57	0.42
35:CH:80:ILE:C	35:CH:81:GLU:HG3	2.40	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:CH:89:ILE:HG12	35:CH:91:LEU:HD13	2.00	0.42
36:CI:53:ALA:O	36:CI:55:ASP:N	2.51	0.42
37:CJ:85:TYR:CD1	37:CJ:154:TYR:CE1	3.07	0.42
41:CN:78:GLN:O	41:CN:103:LEU:HD13	2.19	0.42
54:CA:537:G:C5'	42:CO:113:ARG:HH12	2.28	0.42
43:CP:62:ASN:OD1	26:D4:49:PHE:CD2	2.70	0.42
44:CQ:29:ARG:HG3	44:CQ:29:ARG:HH11	1.84	0.42
54:CA:137:C:H1'	46:CS:63:GLY:HA2	2.00	0.42
54:CA:1318:A:H1'	49:CV:37:ARG:NH2	2.34	0.42
13:D0:38:VAL:HG22	13:D0:112:ALA:HB2	2.01	0.42
13:D0:57:ARG:O	13:D0:59:ASP:N	2.52	0.42
16:D1:24:TYR:HB2	16:D1:29:SER:HB3	2.02	0.42
26:D4:24:THR:OG1	26:D4:25:TYR:N	2.49	0.42
28:D6:36:LEU:CD1	28:D6:50:ARG:NH1	2.82	0.42
55:DA:1057:A:O2'	55:DA:1058:U:OP1	2.37	0.42
55:DA:1236:G:C4'	55:DA:1237:A:OP1	2.53	0.42
55:DA:1423:G:OP1	55:DA:1492:G:O2'	2.37	0.42
55:DA:1518:C:H2'	55:DA:1519:G:H8	1.84	0.42
55:DA:1607:C:H4'	55:DA:1608:A:O5'	2.20	0.42
55:DA:1948:G:C2'	55:DA:1949:G:H5'	2.49	0.42
55:DA:189:G:H2'	55:DA:205:G:N2	2.34	0.42
55:DA:2067:G:C4'	55:DA:2068:U:OP2	2.61	0.42
55:DA:208:C:H2'	55:DA:209:C:C6	2.53	0.42
55:DA:2134:A:N6	55:DA:2157:G:O2'	2.51	0.42
55:DA:2146:C:H4'	55:DA:2147:G:C5	2.54	0.42
55:DA:2168:G:C2'	55:DA:2169:A:OP1	2.67	0.42
55:DA:2287:A:C5	55:DA:2289:G:C5	3.07	0.42
55:DA:2311:A:H3'	55:DA:2312:U:H5	1.75	0.42
55:DA:2555:U:H2'	55:DA:2556:C:O4'	2.19	0.42
55:DA:2590:A:H2'	55:DA:2591:C:H6	1.83	0.42
55:DA:279:C:H2'	55:DA:280:C:C6	2.53	0.42
55:DA:2832:U:H4'	55:DA:2833:G:C5'	2.36	0.42
55:DA:2870:C:C2'	55:DA:2871:C:H5'	2.49	0.42
55:DA:301:G:C4	55:DA:302:C:C5	3.08	0.42
55:DA:35:G:H2'	55:DA:36:G:O4'	2.18	0.42
55:DA:422:A:H2'	55:DA:423:A:C8	2.54	0.42
55:DA:443:A:H3'	5:DF:45:ARG:HH11	1.83	0.42
55:DA:553:U:H2'	55:DA:554:U:O4'	2.19	0.42
55:DA:588:U:O4	55:DA:670:A:O2'	2.29	0.42
55:DA:608:A:C8	55:DA:621:A:N6	2.87	0.42
55:DA:654(F):C:O2'	55:DA:654(G):C:OP1	2.35	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:654(S):G:H2'	55:DA:654(T):A:N9	2.33	0.42
55:DA:726:G:O2'	55:DA:727:A:P	2.77	0.42
55:DA:887:A:O2'	55:DA:888:C:P	2.78	0.42
55:DA:895:U:O2	55:DA:895:U:H2'	2.19	0.42
55:DA:940:G:H2'	55:DA:941:A:O4'	2.19	0.42
4:DE:116:VAL:O	4:DE:117:MET:CB	2.67	0.42
4:DE:59:VAL:O	4:DE:60:ASN:CG	2.58	0.42
4:DE:93:VAL:C	4:DE:95:ILE:H	2.22	0.42
5:DF:181:LEU:HD22	5:DF:181:LEU:HA	1.80	0.42
6:DG:14:GLU:O	6:DG:17:PRO:HG2	2.19	0.42
7:DH:80:SER:C	7:DH:81:GLU:OE1	2.58	0.42
9:DM:39:ARG:C	9:DM:41:ASP:H	2.22	0.42
11:DO:61:ARG:HE	30:D8:24:ALA:HB2	1.84	0.42
12:DP:3:MET:HB3	12:DP:93:TYR:CE1	2.55	0.42
14:DQ:26:LEU:HD22	14:DQ:87:PHE:CD1	2.54	0.42
14:DQ:71:ARG:HG2	14:DQ:104:GLY:CA	2.36	0.42
14:DQ:49:VAL:CG1	14:DQ:76:LYS:HB2	2.49	0.42
19:DT:10:ALA:O	19:DT:28:PHE:HB3	2.19	0.42
21:DV:116:VAL:HG13	21:DV:117:LEU:H	1.84	0.42
21:DV:119:GLU:O	21:DV:121:HIS:N	2.52	0.42
25:DX:2:PRO:O	25:DX:39:ASP:HB2	2.18	0.42
57:DY:135:ARG:CA	57:DY:138:LEU:HB3	2.45	0.42
57:DY:72:ASP:O	57:DY:112:LEU:CG	2.67	0.42
23:DZ:85:LEU:HA	23:DZ:87:PRO:HD2	2.01	0.42
1:AA:2707:G:C5'	13:A0:68:ARG:NH2	2.80	0.42
17:A2:58:VAL:HB	17:A2:98:GLU:CB	2.49	0.42
28:A6:18:ARG:HH11	28:A6:18:ARG:HG3	1.84	0.42
30:A8:8:LYS:HB3	30:A8:12:LYS:HE3	2.00	0.42
1:AA:1101:U:O2'	1:AA:1102:C:H5'	2.20	0.42
1:AA:585:G:C6	1:AA:1253:A:OP1	2.71	0.42
1:AA:1261:C:H2'	1:AA:1262:A:O5'	2.20	0.42
1:AA:1340:U:H2'	1:AA:1341:U:OP1	2.17	0.42
1:AA:1895:C:H1'	31:BA:702:A:H61	1.84	0.42
1:AA:2088:G:H2'	1:AA:2089:U:O4'	2.19	0.42
1:AA:211:A:H2'	1:AA:212:G:O4'	2.19	0.42
1:AA:215:G:H4'	1:AA:216:A:H4'	2.01	0.42
1:AA:2225:A:O2'	1:AA:2226:C:OP2	2.37	0.42
1:AA:2277:G:H5''	12:AP:85:LYS:HB3	2.02	0.42
1:AA:2335:A:O2'	1:AA:2336:A:H8	2.03	0.42
1:AA:2462:U:H2'	1:AA:2463:C:C6	2.54	0.42
1:AA:2750:A:C5'	1:AA:2751:G:OP2	2.67	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2756:U:O4'	1:AA:2757:A:C8	2.72	0.42
1:AA:339:U:H2'	1:AA:339:U:O2	2.19	0.42
1:AA:531:C:N4	1:AA:2035:G:C6	2.87	0.42
1:AA:585:G:O2'	1:AA:1254:A:N6	2.49	0.42
1:AA:654(S):G:N1	1:AA:654(T):A:C2	2.87	0.42
1:AA:753:C:H6	1:AA:753:C:O5'	2.02	0.42
1:AA:897:C:C2'	1:AA:898:C:H5'	2.45	0.42
1:AA:977:G:C6	1:AA:987:G:C6	3.08	0.42
2:AB:47:C:O2	2:AB:47:C:H2'	2.18	0.42
3:AD:186:HIS:HD2	3:AD:188:GLU:H	1.66	0.42
5:AF:118:ALA:HA	5:AF:122:LYS:O	2.19	0.42
5:AF:157:VAL:HG12	5:AF:178:PRO:HA	2.00	0.42
5:AF:40:GLN:OE1	5:AF:182:ASN:HB2	2.19	0.42
8:AK:140:LEU:HD12	8:AK:140:LEU:HA	1.84	0.42
10:AN:11:ALA:HB2	10:AN:64:ARG:NH2	2.34	0.42
12:AP:141:GLN:HB3	12:AP:141:GLN:HE21	1.60	0.42
12:AP:42:ILE:HD13	12:AP:97:VAL:CB	2.49	0.42
12:AP:77:LYS:HA	12:AP:78:PRO:HD3	1.86	0.42
15:AR:52:ILE:HG12	15:AR:61:PHE:HB2	2.00	0.42
18:AS:8:ARG:O	18:AS:9:TYR:HB2	2.18	0.42
21:AV:102:LEU:HD12	21:AV:121:HIS:O	2.19	0.42
21:AV:110:GLY:N	21:AV:143:GLY:CA	2.63	0.42
21:AV:110:GLY:CA	21:AV:143:GLY:HA2	2.48	0.42
23:AZ:73:LEU:O	23:AZ:76:ARG:HG2	2.20	0.42
53:B1:33:G:O2'	53:B1:34:G:H5'	2.20	0.42
53:B1:32:A:H2'	53:B1:33:G:O4'	2.18	0.42
31:BA:1206:G:C4'	33:BF:194:GLY:H	2.33	0.42
31:BA:1251:A:O2'	31:BA:1252:A:H5'	2.19	0.42
31:BA:1366:C:H2'	31:BA:1367:C:C6	2.54	0.42
31:BA:321:A:C2	31:BA:333:G:C2	3.07	0.42
31:BA:374:A:C6	31:BA:375:U:C4	3.07	0.42
31:BA:376:G:H5''	46:BS:5:ARG:HD3	2.01	0.42
31:BA:597:G:H2'	31:BA:598:U:H5'	2.00	0.42
31:BA:680:C:O2'	31:BA:681:C:H5'	2.19	0.42
31:BA:743:U:H2'	31:BA:744:C:C6	2.54	0.42
31:BA:662:G:O2'	31:BA:836:G:H5'	2.19	0.42
52:BD:67:C:H2'	52:BD:68:C:H6	1.84	0.42
32:BE:107:THR:C	32:BE:109:SER:H	2.23	0.42
32:BE:21:ARG:CZ	32:BE:39:ILE:HG13	2.49	0.42
33:BF:164:ARG:HG2	33:BF:165:THR:N	2.32	0.42
31:BA:490:G:P	34:BG:132:ARG:HH22	2.42	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:BG:14:ARG:CG	34:BG:14:ARG:HH11	2.32	0.42
35:BH:15:ARG:HH12	53:B1:55:U:P	2.41	0.42
36:BI:21:LEU:O	36:BI:24:GLU:HB3	2.20	0.42
37:BJ:12:LEU:HD23	37:BJ:28:ASN:ND2	2.34	0.42
37:BJ:15:ASP:OD2	37:BJ:44:TYR:OH	2.36	0.42
31:BA:1233:G:OP2	39:BL:124:GLN:HG3	2.20	0.42
39:BL:17:VAL:HG22	39:BL:63:ILE:CG1	2.49	0.42
31:BA:695:A:OP1	41:BN:52:GLY:HA3	2.19	0.42
45:BR:21:ASP:OD2	45:BR:21:ASP:C	2.57	0.42
46:BS:19:ILE:CG2	46:BS:36:ILE:HG13	2.49	0.42
48:BU:31:LEU:CD2	48:BU:31:LEU:N	2.77	0.42
49:BV:11:VAL:HA	49:BV:38:SER:HA	2.00	0.42
50:BW:100:ILE:O	50:BW:102:GLY:N	2.43	0.42
54:CA:1124:G:H21	54:CA:1280:A:N6	2.17	0.42
54:CA:130:A:H1'	54:CA:264:U:H4'	2.01	0.42
54:CA:815:A:O2'	54:CA:1527:C:C1'	2.67	0.42
54:CA:1542:U:H6	54:CA:1542:U:H2'	1.56	0.42
54:CA:267:C:P	47:CT:67:LYS:HB2	2.59	0.42
54:CA:31:G:O2'	54:CA:32:A:OP1	2.37	0.42
54:CA:437:U:O2'	54:CA:438:G:H5'	2.19	0.42
54:CA:489:C:H2'	54:CA:490:G:C8	2.54	0.42
54:CA:56:U:H2'	54:CA:57:G:H8	1.81	0.42
54:CA:64:G:H4'	54:CA:65:U:O5'	2.19	0.42
54:CA:894:G:C6	54:CA:895:G:C6	3.08	0.42
54:CA:89:U:H2'	54:CA:90:C:O4'	2.20	0.42
54:CA:980:C:H5''	54:CA:981:U:C5	2.54	0.42
52:CD:58:A:HO2'	52:CD:59:U:P	2.42	0.42
32:CE:25:ASN:O	32:CE:27:LYS:N	2.52	0.42
34:CG:119:GLN:O	34:CG:123:HIS:CD2	2.72	0.42
34:CG:14:ARG:HD3	34:CG:14:ARG:O	2.19	0.42
35:CH:102:ALA:HB2	35:CH:120:THR:OG1	2.18	0.42
38:CK:112:LEU:HD12	38:CK:114:THR:CG2	2.49	0.42
38:CK:33:GLU:HG2	38:CK:59:LEU:CD1	2.47	0.42
38:CK:48:TYR:O	38:CK:49:GLU:HB3	2.19	0.42
42:CO:127:GLU:N	42:CO:127:GLU:CD	2.72	0.42
42:CO:43:VAL:CG2	42:CO:44:THR:N	2.81	0.42
33:CF:29:TYR:OH	44:CQ:54:PRO:HD2	2.19	0.42
46:CS:4:ILE:H	46:CS:4:ILE:HD12	1.81	0.42
48:CU:22:VAL:O	48:CU:25:THR:N	2.52	0.42
48:CU:30:ASP:C	48:CU:32:ARG:N	2.72	0.42
48:CU:31:LEU:O	48:CU:65:ILE:HG22	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:CV:64:GLU:HA	49:CV:67:VAL:CG2	2.49	0.42
49:CV:67:VAL:O	49:CV:67:VAL:CG1	2.67	0.42
50:CW:57:ARG:NH1	50:CW:102:GLY:CA	2.83	0.42
13:D0:75:LEU:HD13	13:D0:75:LEU:O	2.20	0.42
27:D5:48:GLU:HG3	27:D5:59:GLU:CB	2.46	0.42
55:DA:1332:G:N2	55:DA:1610:A:C8	2.84	0.42
55:DA:1359:A:OP2	55:DA:1359:A:C8	2.72	0.42
55:DA:1682:G:H2'	55:DA:1683:C:H6	1.80	0.42
55:DA:1707:G:H2'	55:DA:1708:C:H6	1.84	0.42
55:DA:1889:A:H1'	55:DA:2087:G:O4'	2.19	0.42
55:DA:226:G:HO2'	55:DA:227:A:H8	1.61	0.42
55:DA:236:C:H2'	55:DA:237:C:C6	2.54	0.42
55:DA:2444:G:OP2	5:DF:68:LYS:NZ	2.51	0.42
55:DA:746:A:C5	55:DA:2611:U:H5''	2.54	0.42
55:DA:2649:U:H2'	55:DA:2650:U:H6	1.83	0.42
55:DA:2702:U:C2'	55:DA:2703:C:OP2	2.67	0.42
55:DA:2848:G:O2'	55:DA:2867:G:N2	2.52	0.42
55:DA:466:A:H2	55:DA:795:C:O2	2.02	0.42
55:DA:609(A):G:C4	55:DA:610:C:C5	3.07	0.42
55:DA:654(R):C:C6	55:DA:654(R):C:O5'	2.67	0.42
55:DA:730:C:H2'	55:DA:731:C:H6	1.82	0.42
2:DB:34:U:H5''	2:DB:35:U:OP1	2.19	0.42
55:DA:1788:C:OP1	3:DD:222:ARG:NH2	2.52	0.42
3:DD:32:SER:HA	3:DD:35:LYS:O	2.18	0.42
4:DE:54:GLN:CA	4:DE:54:GLN:NE2	2.81	0.42
6:DG:81:LYS:CD	6:DG:81:LYS:N	2.78	0.42
7:DH:24:VAL:CG2	7:DH:35:VAL:HB	2.49	0.42
56:DI:5:ILE:O	56:DI:6:GLU:HB2	2.18	0.42
56:DJ:4:ASP:O	56:DJ:7:ARG:O	2.36	0.42
58:DL:13:PRO:HG2	58:DL:14:ALA:N	2.34	0.42
10:DN:104:ARG:NH1	15:DR:36:GLU:CG	2.81	0.42
11:DO:47:ASP:OD2	11:DO:49:ARG:HG2	2.20	0.42
12:DP:110:THR:HB	12:DP:112:GLU:CG	2.47	0.42
12:DP:34:LEU:HB2	12:DP:118:LEU:HD22	2.00	0.42
20:DU:50:ARG:C	20:DU:52:SER:H	2.22	0.42
21:DV:112:ARG:O	21:DV:113:ALA:HB3	2.20	0.42
25:DX:36:VAL:O	25:DX:36:VAL:HG23	2.18	0.42
57:DY:74:LEU:CG	57:DY:120:LYS:HA	2.45	0.42
57:DY:142:LEU:CD1	57:DY:143:GLN:N	2.52	0.42
13:A0:101:ALA:HB2	27:A5:44:THR:OG1	2.20	0.42
28:A6:15:GLU:HG3	28:A6:47:THR:HG21	2.00	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1462:C:H2'	1:AA:1463:C:C6	2.54	0.42
1:AA:1473:G:H2'	1:AA:1474:C:H6	1.84	0.42
1:AA:1644:C:H2'	1:AA:1644:C:O2	2.19	0.42
1:AA:1685:C:H2'	1:AA:1686:C:H6	1.84	0.42
1:AA:228:A:H2'	1:AA:230:U:O4'	2.20	0.42
1:AA:2675:A:H61	1:AA:2732:G:H1	1.68	0.42
1:AA:335:C:O2'	1:AA:336:C:H5'	2.19	0.42
1:AA:28:A:C4	1:AA:513:A:C8	3.07	0.42
2:AB:9:G:H5'	14:AQ:25:ARG:NH2	2.34	0.42
3:AD:108:PRO:HA	3:AD:196:VAL:O	2.19	0.42
4:AE:137:HIS:CB	4:AE:138:PRO:CD	2.97	0.42
4:AE:4:ILE:HG13	4:AE:28:ALA:HB1	2.00	0.42
4:AE:51:PHE:O	4:AE:74:PRO:HB2	2.19	0.42
5:AF:51:THR:HG21	5:AF:92:PRO:HD2	2.01	0.42
8:AK:110:ASP:OD2	8:AK:113:ARG:CB	2.64	0.42
8:AK:5:LEU:HD23	8:AK:9:LEU:HD21	2.02	0.42
12:AP:21:THR:H	12:AP:98:LYS:HB3	1.84	0.42
19:AT:11:PRO:HD2	24:AW:40:SER:OG	2.19	0.42
20:AU:75:ILE:HG12	20:AU:76:CYS:H	1.85	0.42
21:AV:158:PRO:O	21:AV:159:PRO:C	2.57	0.42
21:AV:7:ALA:C	21:AV:8:TYR:HD2	2.23	0.42
23:AZ:78:LYS:CD	23:AZ:80:LEU:HD11	2.49	0.42
31:BA:1004:A:H2'	31:BA:1005:A:N3	2.34	0.42
31:BA:232:G:H1'	31:BA:262:A:N1	2.34	0.42
31:BA:327:A:O2'	31:BA:329:A:H5'	2.20	0.42
31:BA:577:G:O2'	31:BA:816:A:H2'	2.20	0.42
31:BA:671:G:H2'	31:BA:672:U:H6	1.84	0.42
31:BA:890:G:O2'	31:BA:891:U:C5	2.70	0.42
52:BC:1:G:H2'	52:BC:2:C:H6	1.84	0.42
52:BC:68:C:H2'	52:BC:69:G:H8	1.84	0.42
52:BC:7:A:O2'	52:BC:8:U:P	2.77	0.42
32:BE:18:GLY:O	32:BE:19:HIS:ND1	2.48	0.42
32:BE:41:ILE:CD1	32:BE:41:ILE:N	2.82	0.42
34:BG:175:SER:OG	34:BG:184:LYS:HB2	2.20	0.42
38:BK:123:GLU:O	38:BK:127:LEU:HD22	2.19	0.42
40:BM:24:VAL:HG21	40:BM:37:PRO:HG3	2.01	0.42
43:BP:99:ARG:O	43:BP:100:GLY:C	2.57	0.42
43:BP:86:CYS:O	43:BP:89:GLY:N	2.52	0.42
44:BQ:51:GLY:C	44:BQ:53:LEU:H	2.23	0.42
45:BR:72:ARG:NH1	45:BR:73:GLU:OE2	2.52	0.42
45:BR:75:PRO:O	45:BR:79:ARG:HG3	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:BT:101:ARG:HG2	47:BT:101:ARG:NH2	2.35	0.42
31:BA:130:A:C8	47:BT:63:ARG:HG3	2.54	0.42
47:BT:66:SER:OG	47:BT:69:LYS:HB3	2.20	0.42
31:BA:986:A:H4'	49:BV:55:LYS:HG3	2.01	0.42
54:CA:103:C:C4	54:CA:104:G:N7	2.88	0.42
54:CA:1129:C:N4	54:CA:1141:C:H41	2.18	0.42
54:CA:191(B):G:C2'	54:CA:191(C):G:H5'	2.48	0.42
54:CA:869:G:H4'	54:CA:872:A:N9	2.33	0.42
52:CB:19:G:N2	52:CB:56:C:N3	2.67	0.42
32:CE:111:ARG:NH2	32:CE:114:ARG:HG2	2.35	0.42
32:CE:189:ASP:OD1	32:CE:205:ASP:CG	2.57	0.42
32:CE:21:ARG:O	32:CE:22:LYS:HB2	2.19	0.42
32:CE:30:ARG:HG2	32:CE:30:ARG:H	1.66	0.42
33:CF:43:LEU:O	33:CF:47:LEU:HB3	2.19	0.42
34:CG:61:LYS:HD2	34:CG:206:PHE:CE2	2.55	0.42
36:CI:29:ALA:O	36:CI:30:LEU:C	2.58	0.42
54:CA:1343:G:H1'	39:CL:121:ARG:NH1	2.33	0.42
43:CP:87:TYR:C	43:CP:89:GLY:N	2.73	0.42
45:CR:63:ARG:O	45:CR:67:LEU:HG	2.19	0.42
45:CR:4:THR:HG1	45:CR:7:GLU:HB2	1.81	0.42
46:CS:48:TRP:HZ2	46:CS:76:GLN:OE1	2.02	0.42
46:CS:8:ARG:O	46:CS:9:PHE:CD2	2.70	0.42
16:D1:57:PHE:O	16:D1:58:ARG:C	2.57	0.42
26:D4:50:VAL:O	26:D4:51:ASP:C	2.57	0.42
28:D6:27:LYS:C	28:D6:28:ARG:HG2	2.39	0.42
55:DA:593:G:H1'	30:D8:4:MET:HE1	1.99	0.42
55:DA:594:U:P	30:D8:61:LEU:HD22	2.60	0.42
30:D8:64:TYR:HB3	30:D8:65:GLU:H	1.43	0.42
55:DA:1020:A:H4'	55:DA:1021:A:O5'	2.18	0.42
55:DA:1071:G:H8	55:DA:1071:G:O5'	2.01	0.42
55:DA:1082:U:H2'	57:DY:41:ARG:NE	2.34	0.42
55:DA:1173:G:H1'	55:DA:1175:U:C6	2.54	0.42
55:DA:1367:A:H5'	55:DA:1368:G:OP2	2.19	0.42
55:DA:1535:U:H2'	55:DA:1536:A:O5'	2.20	0.42
55:DA:1583:A:H2'	55:DA:1583:A:N3	2.34	0.42
55:DA:1810:A:H8	55:DA:1810:A:O5'	2.02	0.42
55:DA:2213:U:H6	55:DA:2213:U:O5'	2.01	0.42
55:DA:2627:G:N3	55:DA:2781:A:H2	2.17	0.42
55:DA:2848:G:HO2'	55:DA:2849:U:P	2.42	0.42
55:DA:433:C:H2'	55:DA:434:U:C6	2.54	0.42
55:DA:614:U:C2'	55:DA:615:G:OP1	2.67	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:654(I):C:O2	55:DA:654(I):C:C2'	2.66	0.42
3:DD:122:ASP:O	3:DD:123:ALA:O	2.37	0.42
3:DD:174:ILE:HD12	3:DD:174:ILE:N	2.35	0.42
3:DD:183:ARG:CG	3:DD:183:ARG:NH1	2.76	0.42
4:DE:200:GLU:O	4:DE:201:THR:C	2.57	0.42
4:DE:24:THR:OG1	4:DE:188:VAL:HG12	2.18	0.42
5:DF:133:ASN:O	5:DF:135:LYS:HB2	2.20	0.42
55:DA:1248:G:N2	5:DF:88:VAL:CG2	2.83	0.42
7:DH:143:GLN:O	7:DH:146:ALA:HB3	2.19	0.42
7:DH:45:VAL:O	7:DH:45:VAL:HG13	2.19	0.42
56:DI:28:LYS:O	56:DJ:2:ALA:HB1	2.19	0.42
8:DK:29:TYR:CD1	8:DK:33:ARG:NE	2.86	0.42
58:DL:9:LYS:N	58:DL:9:LYS:CD	2.53	0.42
9:DM:4:TYR:OH	9:DM:7:LYS:HE2	2.19	0.42
11:DO:115:LEU:HB2	11:DO:131:SER:CB	2.44	0.42
15:DR:133:GLU:O	15:DR:135:ALA:N	2.52	0.42
20:DU:94:LYS:NZ	20:DU:101:LYS:HZ1	2.17	0.42
21:DV:172:ALA:O	21:DV:173:ALA:HB2	2.18	0.42
21:DV:198:LYS:O	21:DV:199:LYS:HB3	2.19	0.42
25:DX:10:LYS:O	25:DX:53:LEU:HD22	2.19	0.42
25:DX:5:LYS:CE	25:DX:34:GLU:OE1	2.66	0.42
55:DA:1055:G:O3'	57:DY:35:LYS:CD	2.68	0.42
57:DY:36:GLU:O	57:DY:38:HIS:CE1	2.64	0.42
57:DY:88:ALA:C	56:DJ:15:ALA:CB	2.87	0.42
13:A0:59:ASP:OD1	13:A0:61:HIS:HB3	2.19	0.42
16:A1:111:GLU:O	16:A1:112:ARG:C	2.57	0.42
6:AG:6:ALA:HB3	26:A4:23:GLU:OE1	2.19	0.42
1:AA:11:G:O2'	1:AA:12:U:H5'	2.20	0.42
1:AA:1206:G:OP2	1:AA:1206:G:C8	2.73	0.42
1:AA:1210:A:N3	1:AA:1212:G:N2	2.67	0.42
1:AA:1336:A:H2'	1:AA:1337:G:H8	1.84	0.42
1:AA:1507:A:N3	1:AA:1508:A:H1'	2.34	0.42
1:AA:1813:G:H1'	3:AD:50:THR:HG1	1.83	0.42
1:AA:1961:C:H5'	31:BA:1484:C:O2'	2.20	0.42
1:AA:206:U:H6	1:AA:206:U:O5'	2.02	0.42
1:AA:2313:C:H2'	1:AA:2314:C:H6	1.84	0.42
1:AA:2313:C:O4'	6:AG:40:ASN:OD1	2.37	0.42
1:AA:2617:C:C4	1:AA:2618:G:N7	2.88	0.42
1:AA:2712:U:H1'	1:AA:2712(A):A:N7	2.35	0.42
1:AA:548:A:H2'	1:AA:549:G:C5'	2.49	0.42
1:AA:631:A:H2'	1:AA:632:A:C8	2.54	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:893:C:H4'	1:AA:894:C:OP1	2.20	0.42
4:AE:55:ASN:C	4:AE:57:LYS:N	2.73	0.42
6:AG:106:LEU:HA	6:AG:110:ALA:HB3	2.02	0.42
6:AG:47:LYS:HG3	6:AG:82:LEU:CD2	2.50	0.42
6:AG:89:GLY:C	6:AG:90:LEU:HD22	2.40	0.42
7:AH:132:ARG:O	7:AH:133:VAL:HG23	2.18	0.42
7:AH:6:ARG:C	7:AH:8:PRO:HD2	2.39	0.42
8:AK:82:ARG:HG3	8:AK:82:ARG:NH1	2.30	0.42
10:AN:119:PRO:HB2	15:AR:68:TYR:CE2	2.54	0.42
10:AN:120:GLU:CD	10:AN:122:LEU:HD21	2.40	0.42
11:AO:124:LYS:HE3	11:AO:145:PRO:HD3	2.02	0.42
15:AR:106:SER:O	15:AR:107:ASP:OD1	2.37	0.42
15:AR:132:LYS:CB	15:AR:136:GLN:HE22	2.33	0.42
18:AS:78:GLU:OE2	18:AS:99:ARG:HD2	2.19	0.42
20:AU:24:VAL:HG12	20:AU:25:GLY:N	2.35	0.42
21:AV:92:SER:C	21:AV:130:PRO:HG2	2.38	0.42
24:AW:47:ASN:O	24:AW:50:ILE:CD1	2.68	0.42
53:B1:53:U:C1'	53:B1:54:U:OP1	2.67	0.42
31:BA:1034:G:N2	31:BA:1035:A:C5	2.87	0.42
31:BA:1154:G:N3	31:BA:1155:G:C8	2.87	0.42
31:BA:1203:C:O5'	31:BA:1203:C:H6	2.03	0.42
31:BA:15:G:H1'	35:BH:19:MET:HE2	2.00	0.42
31:BA:549:C:C2	31:BA:550:G:C8	3.08	0.42
31:BA:790:A:C2	31:BA:1497:G:H5''	2.54	0.42
31:BA:899:C:H2'	31:BA:900:A:O4'	2.19	0.42
22:A3:11:ARG:NH1	52:BC:63:G:O3'	2.52	0.42
52:BD:21:A:C2	52:BD:46:G:O6	2.72	0.42
33:BF:199:LYS:HB3	33:BF:201:TYR:HE1	1.85	0.42
33:BF:91:LEU:C	33:BF:93:LYS:H	2.22	0.42
34:BG:162:LEU:HD13	34:BG:181:MET:HG2	2.01	0.42
34:BG:79:PHE:CE1	34:BG:204:ILE:HD13	2.54	0.42
34:BG:16:GLY:CA	34:BG:33:MET:HE1	2.37	0.42
35:BH:18:ARG:HG2	35:BH:19:MET:H	1.84	0.42
37:BJ:16:LEU:O	37:BJ:17:VAL:CG2	2.67	0.42
39:BL:27:THR:HG23	39:BL:31:GLN:H	1.84	0.42
41:BN:23:ALA:HB1	41:BN:88:GLY:N	2.34	0.42
42:BO:22:SER:C	42:BO:24:VAL:N	2.73	0.42
42:BO:55:VAL:HG22	42:BO:69:TYR:HA	2.02	0.42
43:BP:80:ARG:NH1	43:BP:80:ARG:HB3	2.34	0.42
31:BA:740:U:O3'	45:BR:39:LEU:HD23	2.19	0.42
46:BS:49:LEU:CD1	46:BS:73:LEU:HD22	2.48	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:BU:36:ASN:ND2	48:BU:39:VAL:CG2	2.83	0.42
53:C1:36:G:H5'	53:C1:37:G:OP2	2.19	0.42
54:CA:112:G:C2'	54:CA:113:G:H5'	2.49	0.42
54:CA:1197:G:H5'	54:CA:1197:G:C8	2.54	0.42
54:CA:119:A:O2'	54:CA:120:A:OP2	2.24	0.42
54:CA:1275:A:O2'	54:CA:1276:G:H5'	2.19	0.42
54:CA:978:A:C2	54:CA:1319:A:H1'	2.54	0.42
54:CA:1465:C:OP2	15:DR:108:ARG:NH1	2.42	0.42
54:CA:1508:G:O2'	54:CA:1509:C:H5'	2.19	0.42
54:CA:419:C:C4	54:CA:420:U:H5	2.38	0.42
54:CA:939:G:H5''	37:CJ:102:ARG:HH22	1.80	0.42
54:CA:1074:G:O4'	32:CE:104:ASN:HB2	2.20	0.42
32:CE:105:PHE:HE1	32:CE:152:PHE:CE1	2.38	0.42
32:CE:25:ASN:OD1	32:CE:27:LYS:HB2	2.19	0.42
32:CE:70:PHE:HB3	32:CE:81:VAL:CG1	2.49	0.42
34:CG:135:LEU:N	34:CG:135:LEU:HD22	2.32	0.42
34:CG:13:ARG:HB2	34:CG:33:MET:CE	2.50	0.42
34:CG:147:ALA:HB2	34:CG:182:LYS:HA	2.02	0.42
34:CG:161:ASN:O	34:CG:162:LEU:C	2.57	0.42
34:CG:170:VAL:HG22	34:CG:171:GLY:N	2.27	0.42
35:CH:48:ALA:HB1	35:CH:49:PRO:CD	2.49	0.42
37:CJ:106:GLN:O	37:CJ:110:GLN:HG3	2.19	0.42
37:CJ:57:GLU:O	37:CJ:58:PRO:C	2.58	0.42
38:CK:63:LEU:HD22	38:CK:63:LEU:H	1.85	0.42
39:CL:56:LEU:CD2	39:CL:57:GLY:N	2.83	0.42
41:CN:120:ARG:HA	41:CN:121:PRO:HD3	1.92	0.42
43:CP:87:TYR:HA	43:CP:90:LEU:CG	2.48	0.42
46:CS:55:ARG:O	46:CS:56:ALA:C	2.57	0.42
49:CV:30:LEU:HD13	49:CV:30:LEU:N	2.34	0.42
49:CV:42:PRO:O	49:CV:45:VAL:HG22	2.20	0.42
50:CW:99:LEU:O	50:CW:100:ILE:HB	2.19	0.42
13:D0:90:ARG:HH21	13:D0:118:GLU:HA	1.83	0.42
28:D6:15:GLU:OE2	28:D6:44:ARG:NH2	2.53	0.42
30:D8:8:LYS:HB3	30:D8:12:LYS:HE3	2.01	0.42
55:DA:1057:A:C6	55:DA:1086:A:H2	2.37	0.42
55:DA:1025:G:C4	55:DA:1135:C:H1'	2.55	0.42
55:DA:1343:G:N3	55:DA:1343:G:H2'	2.33	0.42
55:DA:1542:G:H5''	55:DA:1543:A:OP2	2.19	0.42
55:DA:1830:C:O2'	55:DA:1831:G:H5'	2.19	0.42
55:DA:2129:C:H2'	55:DA:2130:U:H5'	2.01	0.42
55:DA:2131:G:OP1	55:DA:2133:G:OP2	2.36	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:198:C:H5'	55:DA:2244:U:OP1	2.19	0.42
55:DA:2266:A:H4'	55:DA:2267:A:C2	2.55	0.42
55:DA:2431:U:C2	55:DA:2433:A:OP2	2.72	0.42
55:DA:2603:G:H4'	55:DA:2603:G:OP2	2.20	0.42
55:DA:2814:C:C6	55:DA:2815:C:C5	3.08	0.42
55:DA:654(S):G:H2'	55:DA:654(T):A:O4'	2.20	0.42
55:DA:740:U:H2'	55:DA:741:G:H8	1.85	0.42
55:DA:784:A:C5	3:DD:229:VAL:CG2	3.02	0.42
55:DA:880:G:C6	55:DA:881:G:N7	2.87	0.42
55:DA:989:G:N7	25:DX:13:ILE:HD12	2.34	0.42
55:DA:990:A:C6	55:DA:1186:G:H1'	2.54	0.42
2:DB:14:U:H4'	2:DB:106:G:H21	1.83	0.42
4:DE:34:VAL:O	4:DE:35:GLN:CB	2.68	0.42
4:DE:77:ILE:O	4:DE:78:LEU:C	2.55	0.42
6:DG:63:ILE:HG22	6:DG:144:ILE:HD11	2.02	0.42
6:DG:35:GLU:O	6:DG:36:LYS:HB3	2.19	0.42
6:DG:58:GLN:HE22	6:DG:148:MET:HE1	1.84	0.42
7:DH:111:HIS:CE1	7:DH:112:PRO:O	2.73	0.42
7:DH:85:LYS:HD2	7:DH:85:LYS:HA	1.71	0.42
7:DH:88:LEU:O	7:DH:163:TYR:N	2.43	0.42
7:DH:89:ILE:O	7:DH:90:LYS:C	2.58	0.42
57:DY:125:LEU:CD1	56:DJ:20:LEU:HD21	2.49	0.42
8:DK:81:VAL:HG22	8:DK:143:SER:O	2.19	0.42
58:DL:19:PRO:CD	58:DL:38:VAL:HG11	2.44	0.42
10:DN:26:LYS:HB2	10:DN:30:ALA:CB	2.48	0.42
11:DO:139:LYS:HG3	11:DO:144:GLU:OE2	2.19	0.42
12:DP:38:GLU:OE2	12:DP:128:LYS:N	2.53	0.42
12:DP:48:GLU:HA	12:DP:48:GLU:OE1	2.19	0.42
15:DR:29:ARG:NH1	15:DR:89:VAL:HG11	2.35	0.42
20:DU:86:ARG:HA	20:DU:86:ARG:HD2	1.68	0.42
21:DV:117:LEU:O	21:DV:118:GLN:O	2.38	0.42
25:DX:7:LYS:HG3	25:DX:34:GLU:CG	2.49	0.42
57:DY:13:LEU:O	57:DY:14:LYS:HE3	2.18	0.42
57:DY:25:PHE:CG	57:DY:82:PHE:CE2	2.93	0.42
16:A1:92:ARG:CG	16:A1:94:ASN:HB3	2.37	0.42
17:A2:4:ILE:HA	17:A2:12:TYR:O	2.19	0.42
17:A2:78:LYS:HB3	17:A2:78:LYS:HE2	1.82	0.42
17:A2:79:VAL:HG23	17:A2:80:GLN:N	2.33	0.42
26:A4:15:ILE:CD1	26:A4:15:ILE:H	2.32	0.42
27:A5:51:TYR:HB3	27:A5:52:TYR:H	1.45	0.42
1:AA:1085:A:O2'	1:AA:1086:A:OP1	2.36	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1246:A:H5''	11:AO:15:ARG:HH12	1.84	0.42
1:AA:13:A:O2'	1:AA:14:A:N7	2.53	0.42
1:AA:1423:G:H2'	1:AA:1424:G:H8	1.85	0.42
1:AA:1694:C:O2'	1:AA:1695:G:H5''	2.19	0.42
1:AA:1729:A:C6	1:AA:1731:G:C5	3.08	0.42
1:AA:1837:C:O2	1:AA:1927:A:H2	2.02	0.42
1:AA:2025:C:H2'	1:AA:2026:C:C6	2.55	0.42
1:AA:2258:C:C2'	1:AA:2427:C:OP2	2.68	0.42
1:AA:2657:A:H2'	1:AA:2658:C:C5'	2.46	0.42
1:AA:2657:A:C2'	1:AA:2658:C:H5'	2.48	0.42
1:AA:2711:A:C8	1:AA:2714:G:H1'	2.54	0.42
1:AA:27:G:O2'	1:AA:28:A:O5'	2.38	0.42
1:AA:2814:C:C6	1:AA:2815:C:C5	3.07	0.42
1:AA:2861:G:C4	1:AA:2862:G:C8	3.08	0.42
1:AA:2864:G:H2'	1:AA:2865:U:C6	2.55	0.42
1:AA:387:U:C4'	1:AA:388:G:O5'	2.63	0.42
1:AA:698:C:O2'	1:AA:734:A:N6	2.51	0.42
3:AD:17:THR:O	3:AD:211:ARG:NH2	2.52	0.42
5:AF:21:ALA:HB3	5:AF:23:ASP:OD2	2.19	0.42
5:AF:41:LEU:O	5:AF:44:ARG:HG2	2.19	0.42
7:AH:122:THR:O	7:AH:123:PHE:CD2	2.72	0.42
7:AH:93:GLY:C	7:AH:94:TYR:HD1	2.23	0.42
12:AP:84:GLY:HA2	22:A3:10:THR:HG21	2.01	0.42
20:AU:6:HIS:NE2	20:AU:72:VAL:CG2	2.83	0.42
21:AV:56:VAL:C	21:AV:57:ILE:HD12	2.40	0.42
21:AV:60:GLU:O	21:AV:65:GLN:C	2.58	0.42
25:AX:30:ARG:O	25:AX:33:GLN:HB3	2.20	0.42
31:BA:1000:A:O2'	31:BA:1001:G:H5''	2.19	0.42
31:BA:1032:A:N7	31:BA:1032(A):G:H1'	2.35	0.42
31:BA:1064:G:O4'	31:BA:1066:C:H1'	2.20	0.42
31:BA:1232:U:P	39:BL:126:SER:HG	2.42	0.42
31:BA:1362:C:H2'	31:BA:1362(A):C:H5''	2.00	0.42
31:BA:1408:A:C5	31:BA:1409:C:C5	3.08	0.42
31:BA:1524:C:OP1	41:BN:120:ARG:NH1	2.38	0.42
31:BA:1528:U:HO2'	31:BA:1529:G:H5''	1.77	0.42
31:BA:210:U:H1'	31:BA:216:G:C8	2.55	0.42
31:BA:872:A:C4	31:BA:874:G:N7	2.88	0.42
52:BD:21:A:H61	52:BD:46:G:H22	1.66	0.42
32:BE:153:ARG:HB2	32:BE:154:LEU:H	1.56	0.42
32:BE:33:TYR:HB2	32:BE:43:ASP:HA	2.02	0.42
31:BA:612:C:O3'	34:BG:84:LYS:NZ	2.53	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:BH:7:GLU:OE1	35:BH:37:ARG:NH2	2.52	0.42
37:BJ:69:VAL:HA	37:BJ:138:LYS:HD2	2.01	0.42
37:BJ:44:TYR:C	37:BJ:46:ALA:N	2.71	0.42
38:BK:86:ILE:CB	38:BK:133:LEU:HD22	2.50	0.42
40:BM:75:ILE:CG1	40:BM:76:ASN:H	2.31	0.42
42:BO:43:VAL:HG23	42:BO:44:THR:N	2.34	0.42
43:BP:3:ARG:HD2	43:BP:9:ILE:CD1	2.29	0.42
40:BM:52:GLY:O	44:BQ:42:ILE:HD11	2.19	0.42
46:BS:40:ASP:O	46:BS:48:TRP:HB2	2.19	0.42
31:BA:323:U:O4'	50:BW:19:SER:HB2	2.20	0.42
50:BW:72:LEU:HD23	50:BW:72:LEU:C	2.40	0.42
54:CA:109:A:C8	54:CA:326:G:H2'	2.54	0.42
54:CA:1157:A:OP1	54:CA:1157:A:O4'	2.37	0.42
54:CA:141:A:C1'	54:CA:182:U:O2	2.62	0.42
54:CA:197:A:C6	54:CA:221:C:H4'	2.54	0.42
54:CA:407:G:OP1	34:CG:115:ARG:CZ	2.67	0.42
54:CA:642:A:H2'	54:CA:643:C:C6	2.54	0.42
54:CA:671:G:H2'	54:CA:672:U:C6	2.53	0.42
54:CA:715:A:H5''	54:CA:805:C:H1'	2.02	0.42
54:CA:913:A:O2'	54:CA:914:A:P	2.77	0.42
54:CA:953:G:N7	43:CP:104:ARG:NH2	2.59	0.42
54:CA:976:G:H2'	54:CA:1362:C:N3	2.34	0.42
52:CD:27:G:H2'	52:CD:28:G:C8	2.54	0.42
32:CE:29:ALA:CA	32:CE:32:ILE:HG22	2.50	0.42
32:CE:88:ALA:HB2	32:CE:219:VAL:CG1	2.49	0.42
36:CI:15:ASP:OD1	36:CI:17:SER:HB3	2.20	0.42
37:CJ:26:PHE:HA	37:CJ:101:LEU:HD13	2.02	0.42
37:CJ:138:LYS:HE2	37:CJ:142:GLU:OE2	2.19	0.42
37:CJ:59:LEU:O	37:CJ:62:PHE:HB3	2.19	0.42
37:CJ:65:ALA:O	37:CJ:69:VAL:HG23	2.19	0.42
35:CH:152:ARG:HB3	38:CK:43:GLY:HA3	2.00	0.42
40:CM:51:ARG:HG3	40:CM:60:ARG:O	2.20	0.42
54:CA:538:G:OP1	42:CO:113:ARG:HD2	2.19	0.42
45:CR:62:GLN:O	45:CR:63:ARG:C	2.57	0.42
48:CU:58:LEU:HB3	48:CU:62:GLU:HB3	2.01	0.42
50:CW:36:LEU:HB3	50:CW:59:ALA:HB2	2.02	0.42
17:D2:44:LYS:O	17:D2:46:VAL:HG12	2.20	0.42
26:D4:49:PHE:N	26:D4:49:PHE:CD1	2.84	0.42
26:D4:61:ARG:C	26:D4:63:TYR:N	2.70	0.42
30:D8:42:ARG:HH11	30:D8:42:ARG:HG2	1.84	0.42
30:D8:6:THR:HG22	30:D8:63:PRO:CD	2.50	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1092:C:H2'	55:DA:1093:G:H5'	2.01	0.42
55:DA:1281:G:H5'	55:DA:1281:G:H8	1.84	0.42
55:DA:1384:A:H1'	55:DA:1405:U:O4'	2.20	0.42
55:DA:1548:C:H2'	55:DA:1549:C:H6	1.84	0.42
55:DA:1735:C:C6	55:DA:1735:C:C5'	2.98	0.42
55:DA:1907:G:H2'	55:DA:1908:C:C6	2.55	0.42
55:DA:1925:C:N3	55:DA:1929:G:N2	2.67	0.42
55:DA:2287:A:C2'	55:DA:2288:A:H3'	2.49	0.42
55:DA:2439:A:H1'	55:DA:2587:A:OP1	2.19	0.42
55:DA:2640:G:C6	55:DA:2775:A:C2	3.08	0.42
55:DA:590:A:OP1	5:DF:95:ARG:NH1	2.52	0.42
55:DA:897:C:O2	55:DA:898:C:H1'	2.20	0.42
55:DA:943:U:OP2	11:DO:36:LYS:NZ	2.52	0.42
55:DA:962:G:H2'	55:DA:963:U:H6	1.85	0.42
4:DE:137:HIS:CB	4:DE:138:PRO:HD2	2.47	0.42
6:DG:53:LEU:CD2	6:DG:54:GLU:N	2.77	0.42
6:DG:61:ALA:HA	6:DG:64:THR:HG23	2.02	0.42
7:DH:19:VAL:CG1	7:DH:20:ALA:H	2.32	0.42
56:DJ:10:GLU:OE2	56:DJ:19:GLU:OE2	2.37	0.42
8:DK:11:ASN:C	8:DK:12:LEU:HD13	2.40	0.42
8:DK:78:THR:HG22	8:DK:141:LYS:HD2	2.00	0.42
8:DK:96:ASP:C	8:DK:98:ALA:N	2.71	0.42
58:DL:52:ILE:HG13	58:DL:53:VAL:H	1.82	0.42
9:DM:39:ARG:HH11	9:DM:39:ARG:CB	2.32	0.42
55:DA:2565:A:H62	10:DN:28:SER:CB	2.33	0.42
11:DO:127:ALA:HB3	11:DO:130:PHE:CE2	2.54	0.42
14:DQ:52:SER:O	14:DQ:56:LEU:HD22	2.19	0.42
15:DR:105:LEU:O	15:DR:105:LEU:CG	2.63	0.42
15:DR:76:PHE:HA	15:DR:77:PRO:HD3	1.77	0.42
15:DR:33:LYS:HG3	15:DR:82:LEU:O	2.20	0.42
18:DS:95:ILE:HG13	18:DS:95:ILE:H	1.71	0.42
19:DT:21:PHE:N	19:DT:21:PHE:CD2	2.88	0.42
20:DU:50:ARG:CA	20:DU:53:PRO:HD2	2.49	0.42
20:DU:42:VAL:HG12	20:DU:65:ALA:H	1.83	0.42
21:DV:35:ARG:HH11	21:DV:35:ARG:HB2	1.84	0.42
21:DV:6:LYS:HB2	21:DV:7:ALA:H	1.66	0.42
57:DY:4:LYS:HB3	57:DY:5:ARG:CD	2.49	0.42
57:DY:73:GLY:HA3	57:DY:112:LEU:HD12	1.86	0.42
57:DY:74:LEU:HD11	57:DY:75:GLN:HG2	2.00	0.42
13:A0:18:LEU:CD1	13:A0:22:ARG:NE	2.82	0.42
13:A0:41:ALA:C	13:A0:43:GLU:N	2.73	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:A0:101:ALA:HB2	27:A5:44:THR:HB	2.02	0.42
1:AA:1195:G:C2'	1:AA:1196:C:H5'	2.50	0.42
1:AA:1210:A:H5''	1:AA:1211:U:O5'	2.20	0.42
1:AA:141:A:H5''	1:AA:141(A):C:C5	2.55	0.42
1:AA:1699:G:C6	1:AA:1763:G:N3	2.87	0.42
1:AA:1878:G:H2'	1:AA:1879:C:C6	2.55	0.42
1:AA:1912:A:C8	1:AA:1918:A:C2	3.08	0.42
1:AA:2187:G:C2'	1:AA:2188:C:H5'	2.50	0.42
1:AA:2195:C:O2'	1:AA:2196:C:H5'	2.20	0.42
1:AA:363(F):A:C5'	1:AA:364:C:OP1	2.61	0.42
1:AA:783:A:H3'	1:AA:783:A:C8	2.54	0.42
1:AA:855:G:H2'	1:AA:856:C:H6	1.85	0.42
2:AB:82:G:C2'	2:AB:83:G:H5'	2.49	0.42
3:AD:132:PRO:HD2	3:AD:135:PHE:HD1	1.84	0.42
3:AD:35:LYS:HE3	3:AD:63:ARG:C	2.40	0.42
3:AD:75:ILE:HG21	3:AD:99:ASP:HB2	2.00	0.42
4:AE:201:THR:C	4:AE:202:LYS:HE3	2.40	0.42
4:AE:35:GLN:HB2	4:AE:48:GLN:NE2	2.35	0.42
4:AE:32:PRO:HD2	4:AE:51:PHE:H	1.85	0.42
5:AF:114:VAL:O	5:AF:115:ALA:C	2.57	0.42
5:AF:178:PRO:HB3	5:AF:198:ALA:CB	2.48	0.42
6:AG:115:ARG:CZ	6:AG:115:ARG:HB3	2.49	0.42
7:AH:87:LEU:O	7:AH:131:VAL:HB	2.19	0.42
7:AH:13:LYS:HB3	7:AH:13:LYS:HZ2	1.83	0.42
7:AH:149:ARG:C	7:AH:151:ILE:N	2.73	0.42
7:AH:155:SER:O	7:AH:156:ALA:C	2.57	0.42
8:AK:40:THR:O	8:AK:44:LEU:HG	2.20	0.42
9:AM:131:GLN:NE2	9:AM:132:ALA:CB	2.83	0.42
10:AN:104:ARG:HH22	10:AN:107:ARG:HH21	1.66	0.42
10:AN:23:ARG:HH11	10:AN:23:ARG:CG	2.33	0.42
1:AA:958:U:OP2	12:AP:14:ARG:NH1	2.53	0.42
12:AP:1:MET:O	12:AP:3:MET:N	2.53	0.42
12:AP:66:ILE:C	12:AP:68:ILE:H	2.22	0.42
12:AP:66:ILE:O	12:AP:68:ILE:N	2.53	0.42
14:AQ:78:LEU:HD21	14:AQ:108:GLY:HA3	2.02	0.42
15:AR:12:SER:O	15:AR:15:VAL:HG22	2.19	0.42
15:AR:20:PRO:HD2	15:AR:86:ILE:HG23	2.02	0.42
18:AS:20:VAL:CG2	18:AS:21:VAL:N	2.82	0.42
20:AU:84:ARG:CZ	20:AU:97:ARG:HB2	2.48	0.42
21:AV:118:GLN:HE21	21:AV:118:GLN:HA	1.82	0.42
21:AV:12:GLY:C	21:AV:13:GLU:CG	2.87	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:AV:141:VAL:HG22	21:AV:144:LEU:CB	2.50	0.42
31:BA:1051:C:H2'	31:BA:1052:U:C6	2.55	0.42
31:BA:1101:A:H4'	31:BA:1102:A:H4'	2.00	0.42
31:BA:1179:A:H2'	31:BA:1180:A:O4'	2.19	0.42
31:BA:1425:U:H2'	31:BA:1426:C:H6	1.84	0.42
31:BA:1517:G:H2'	31:BA:1518:A:H8	1.85	0.42
31:BA:1524:C:H2'	31:BA:1525:G:C8	2.55	0.42
31:BA:201:C:C4	31:BA:209:U:C6	3.07	0.42
31:BA:231:G:O2'	31:BA:232:G:H5'	2.19	0.42
31:BA:31:G:C2'	31:BA:32:A:O5'	2.68	0.42
31:BA:346:G:N2	31:BA:347:G:C8	2.87	0.42
31:BA:357:G:O2'	31:BA:358:U:H5'	2.19	0.42
31:BA:397:A:H2'	31:BA:399:G:OP2	2.20	0.42
31:BA:511:C:O2'	31:BA:512:U:OP2	2.30	0.42
31:BA:673:G:H2'	31:BA:674:G:H8	1.74	0.42
31:BA:692:U:H5	41:BN:26:ASN:OD1	2.02	0.42
31:BA:581:G:N1	31:BA:759:A:OP2	2.51	0.42
52:BB:18:G:O2'	52:BB:19:G:P	2.77	0.42
52:BC:74:C:H2'	52:BC:74:C:O2	2.20	0.42
32:BE:236:TYR:HB2	32:BE:239:VAL:HB	2.02	0.42
33:BF:113:ALA:O	33:BF:115:LEU:N	2.52	0.42
33:BF:156:ARG:HD2	33:BF:193:TYR:CD1	2.55	0.42
35:BH:80:ILE:HG22	38:BK:104:ARG:NH2	2.34	0.42
37:BJ:50:ILE:C	37:BJ:52:GLU:N	2.73	0.42
38:BK:83:ILE:O	38:BK:83:ILE:HG23	2.18	0.42
39:BL:46:ALA:C	39:BL:48:GLU:H	2.21	0.42
26:A4:34:GLU:OE1	43:BP:3:ARG:CB	2.67	0.42
43:BP:23:TYR:CD1	43:BP:71:ARG:CZ	3.02	0.42
31:BA:1309:G:O2'	43:BP:77:ASN:ND2	2.52	0.42
43:BP:90:LEU:CD1	49:BV:78:ARG:HE	2.32	0.42
50:BW:13:LEU:H	50:BW:13:LEU:HD12	1.82	0.42
50:BW:64:ASP:OD1	50:BW:81:LYS:HD2	2.20	0.42
54:CA:1051:C:H2'	54:CA:1052:U:C6	2.54	0.42
54:CA:1191:A:P	33:CF:3:ASN:ND2	2.93	0.42
54:CA:119:A:H4'	54:CA:120:A:O5'	2.19	0.42
54:CA:1308:U:H5''	43:CP:98:VAL:HB	2.01	0.42
54:CA:181:G:O2'	54:CA:182:U:H6	2.03	0.42
54:CA:129(A):G:H1'	54:CA:190:G:H5'	2.02	0.42
54:CA:342:C:H2'	54:CA:343:U:O4'	2.18	0.42
54:CA:48:C:H6	54:CA:365:U:O4	2.03	0.42
54:CA:632:A:H3'	54:CA:633:G:C8	2.50	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:658:G:C4	54:CA:659:U:C5	3.08	0.42
54:CA:74:C:N4	54:CA:75:C:C4	2.87	0.42
52:CC:14:A:H2'	52:CC:15:G:H5'	2.02	0.42
32:CE:215:LEU:O	32:CE:219:VAL:HG23	2.20	0.42
33:CF:110:ASN:O	33:CF:141:VAL:HG13	2.20	0.42
33:CF:173:VAL:N	33:CF:174:PRO:HD3	2.34	0.42
33:CF:78:GLY:HA3	33:CF:83:ARG:CB	2.50	0.42
34:CG:112:VAL:HG13	34:CG:113:SER:N	2.31	0.42
34:CG:129:ASN:H	34:CG:145:GLU:HB2	1.79	0.42
34:CG:3:ARG:O	34:CG:4:TYR:C	2.57	0.42
54:CA:1081:G:P	35:CH:16:THR:OG1	2.78	0.42
38:CK:83:ILE:CG1	38:CK:137:VAL:HG22	2.45	0.42
38:CK:34:GLU:HB3	38:CK:118:VAL:CG2	2.41	0.42
41:CN:102:GLY:O	41:CN:103:LEU:C	2.58	0.42
43:CP:108:ARG:HH11	43:CP:108:ARG:HG3	1.85	0.42
50:CW:97:ALA:O	50:CW:99:LEU:N	2.45	0.42
16:D1:76:TYR:C	16:D1:76:TYR:CD2	2.92	0.42
17:D2:35:LEU:N	17:D2:35:LEU:CD2	2.81	0.42
17:D2:47:VAL:HG13	17:D2:48:GLY:H	1.85	0.42
22:D3:25:ARG:HA	22:D3:29:GLN:HE22	1.82	0.42
29:D7:8:ASN:ND2	29:D7:10:ARG:H	2.18	0.42
55:DA:1062:G:C1'	55:DA:1088:A:C5	3.03	0.42
55:DA:1079:C:C6	55:DA:1079:C:C5'	3.02	0.42
55:DA:1080:A:N3	58:DL:126:MET:HG3	2.35	0.42
55:DA:1107:G:H4'	57:DY:30:GLN:HE22	1.84	0.42
55:DA:1575:C:H2'	55:DA:1576:U:C6	2.55	0.42
55:DA:171:G:H2'	55:DA:172:C:H6	1.84	0.42
55:DA:1267:U:C5	55:DA:2012:G:C2	3.08	0.42
55:DA:2127:G:H21	55:DA:2173:A:H1'	1.84	0.42
55:DA:2308:G:H2'	55:DA:2310:A:C2	2.54	0.42
55:DA:2754:U:H5'	55:DA:2755:C:P	2.59	0.42
55:DA:2770:G:C5'	55:DA:2771:C:OP2	2.68	0.42
55:DA:2787:C:H1'	4:DE:62:PRO:CD	2.41	0.42
55:DA:324:A:N6	55:DA:338:G:O2'	2.53	0.42
55:DA:511:U:O4	55:DA:512:G:C2	2.73	0.42
55:DA:588:U:C2	5:DF:90:PHE:CD1	3.07	0.42
55:DA:729:G:C6	3:DD:208:LYS:HB2	2.55	0.42
55:DA:747:U:O2'	18:DS:88:ARG:HG3	2.19	0.42
55:DA:753:C:O5'	55:DA:753:C:H6	2.02	0.42
55:DA:886:C:O2	55:DA:887:A:C6	2.72	0.42
55:DA:931:G:H2'	55:DA:932:G:H5''	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:DD:147:LEU:HD12	3:DD:147:LEU:HA	1.72	0.42
3:DD:85:ASP:HB2	3:DD:92:ILE:CG1	2.49	0.42
55:DA:1817:G:P	3:DD:88:ARG:HH22	2.42	0.42
4:DE:107:THR:O	4:DE:107:THR:HG22	2.19	0.42
4:DE:175:VAL:O	4:DE:175:VAL:HG22	2.19	0.42
4:DE:52:LEU:HD13	4:DE:75:VAL:HG23	2.01	0.42
5:DF:59:TYR:N	5:DF:59:TYR:CD2	2.85	0.42
6:DG:126:ASP:OD2	6:DG:130:ASN:HB2	2.20	0.42
6:DG:2:PRO:C	6:DG:4:ASP:N	2.73	0.42
8:DK:81:VAL:O	8:DK:83:ALA:N	2.52	0.42
9:DM:26:LEU:O	9:DM:27:ALA:C	2.57	0.42
9:DM:95:PRO:O	9:DM:96:GLU:CG	2.68	0.42
11:DO:115:LEU:C	11:DO:115:LEU:HD12	2.39	0.42
14:DQ:111:GLU:C	14:DQ:112:PHE:CD1	2.93	0.42
18:DS:11:ARG:HH21	18:DS:99:ARG:N	2.18	0.42
20:DU:48:ALA:CB	20:DU:61:ILE:HD13	2.49	0.42
21:DV:141:VAL:C	21:DV:143:GLY:N	2.72	0.42
57:DY:69:PRO:N	57:DY:114:GLY:O	2.52	0.42
57:DY:22:GLY:O	57:DY:67:GLY:O	2.38	0.42
57:DY:29:TYR:CE2	57:DY:32:LEU:CD2	3.02	0.42
57:DY:38:HIS:CG	57:DY:40:LEU:HB3	2.55	0.42
57:DY:54:ALA:HB3	57:DY:58:LEU:HD21	2.02	0.42
57:DY:71:LEU:CD2	57:DY:72:ASP:N	2.64	0.42
23:DZ:40:ARG:NH2	23:DZ:42:GLN:HG2	2.35	0.42
23:DZ:8:SER:HB3	23:DZ:66:HIS:CG	2.54	0.42
1:AA:1278:A:H4'	13:A0:34:ILE:HD11	2.00	0.42
16:A1:97:ASP:O	16:A1:98:LEU:O	2.37	0.42
17:A2:23:GLU:O	17:A2:92:THR:HB	2.20	0.42
22:A3:43:THR:HG23	22:A3:46:LYS:HE2	2.01	0.42
1:AA:1431:U:O2'	1:AA:1432:C:H5'	2.19	0.42
1:AA:1906:G:C2	1:AA:1925:C:C2	3.08	0.42
1:AA:1952:A:N6	1:AA:1953:A:N1	2.68	0.42
1:AA:2285:C:H5''	1:AA:2286:A:OP2	2.19	0.42
1:AA:2415:G:C5	1:AA:2416:C:C5	3.08	0.42
1:AA:2443:C:H2'	1:AA:2444:G:H8	1.85	0.42
1:AA:2477:C:H3'	1:AA:2477:C:O2	2.20	0.42
1:AA:2534:A:H2'	1:AA:2535:G:O4'	2.20	0.42
1:AA:445:C:O2'	1:AA:446:G:H5'	2.19	0.42
1:AA:531:C:C5	1:AA:2035:G:C2	3.08	0.42
1:AA:675:A:N6	1:AA:676:A:N6	2.67	0.42
1:AA:685:A:N1	1:AA:787:U:H1'	2.34	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:688:U:H5'	1:AA:1780:A:N1	2.35	0.42
1:AA:750:A:C2	1:AA:753:C:C6	3.08	0.42
1:AA:755:C:H2'	1:AA:756:C:H6	1.83	0.42
1:AA:811:U:O2'	1:AA:1250:G:H2'	2.20	0.42
1:AA:954:G:H2'	1:AA:2274:A:H2	1.84	0.42
1:AA:959:A:N1	1:AA:960:A:C2	2.88	0.42
2:AB:109:G:C4	2:AB:110:G:C8	3.08	0.42
3:AD:147:LEU:HD11	3:AD:183:ARG:NH1	2.35	0.42
1:AA:1566:A:OP1	3:AD:211:ARG:NH1	2.52	0.42
3:AD:24:ILE:O	3:AD:25:THR:HG22	2.19	0.42
7:AH:4:ILE:HD11	7:AH:7:LEU:CB	2.49	0.42
8:AK:101:LEU:O	8:AK:102:SER:HB3	2.19	0.42
8:AK:7:GLU:HG3	8:AK:9:LEU:H	1.84	0.42
8:AK:81:VAL:HG12	8:AK:82:ARG:O	2.19	0.42
9:AM:43:THR:O	9:AM:46:VAL:HG12	2.19	0.42
11:AO:101:VAL:CG1	11:AO:102:ARG:N	2.83	0.42
11:AO:48:PRO:O	11:AO:49:ARG:C	2.58	0.42
12:AP:54:MET:HG3	12:AP:54:MET:H	1.65	0.42
12:AP:42:ILE:CD1	12:AP:97:VAL:HB	2.49	0.42
14:AQ:103:GLU:OE1	14:AQ:103:GLU:N	2.53	0.42
15:AR:90:GLN:CA	15:AR:90:GLN:NE2	2.74	0.42
18:AS:65:LEU:HD13	18:AS:67:ASP:HB2	2.00	0.42
19:AT:35:THR:O	19:AT:39:ILE:HD13	2.20	0.42
21:AV:120:ILE:O	21:AV:121:HIS:CB	2.68	0.42
2:AB:77:U:P	21:AV:19:ARG:HH22	2.43	0.42
21:AV:53:ILE:HA	21:AV:70:LEU:HD21	2.01	0.42
24:AW:40:SER:C	24:AW:42:GLY:N	2.71	0.42
24:AW:43:GLN:O	24:AW:44:LEU:CG	2.62	0.42
31:BA:1054:C:N4	52:BB:34:G:H1'	2.35	0.42
31:BA:1211:U:H5'	31:BA:1212:U:OP1	2.20	0.42
31:BA:1240:U:C2'	31:BA:1241:G:OP1	2.68	0.42
31:BA:233:C:O2'	31:BA:234:C:H5'	2.20	0.42
31:BA:316:G:C2	31:BA:338:A:C2	3.07	0.42
31:BA:160:A:C1'	31:BA:344:A:C5	3.03	0.42
31:BA:487:A:H2'	31:BA:488:C:O4'	2.20	0.42
31:BA:554:C:H2'	31:BA:555:C:C6	2.55	0.42
31:BA:562:C:O2'	42:BO:15:ARG:CB	2.53	0.42
31:BA:652:U:O2'	31:BA:653:A:C5'	2.67	0.42
31:BA:22:G:H4'	31:BA:885:G:C8	2.54	0.42
31:BA:887:G:C3'	31:BA:888:G:C5'	2.94	0.42
52:BB:58:A:O2'	52:BB:59:U:P	2.78	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:BE:32:ILE:HG12	32:BE:40:HIS:CD2	2.55	0.42
32:BE:70:PHE:N	32:BE:70:PHE:CD1	2.88	0.42
33:BF:115:LEU:O	33:BF:118:GLN:N	2.52	0.42
33:BF:41:GLY:O	33:BF:45:LYS:HE3	2.19	0.42
34:BG:141:ARG:HB3	34:BG:142:PRO:CD	2.50	0.42
34:BG:19:LEU:HD12	34:BG:19:LEU:H	1.85	0.42
37:BJ:79:ARG:HG3	37:BJ:84:ASN:ND2	2.35	0.42
37:BJ:85:TYR:CE1	37:BJ:154:TYR:HE1	2.38	0.42
42:BO:20:LYS:H	42:BO:20:LYS:HD3	1.85	0.42
43:BP:73:GLU:OE1	43:BP:77:ASN:ND2	2.50	0.42
44:BQ:34:TYR:CD1	44:BQ:34:TYR:N	2.87	0.42
45:BR:3:ILE:HG22	45:BR:38:ARG:CZ	2.49	0.42
47:BT:15:MET:HE3	47:BT:18:THR:HG22	2.02	0.42
48:BU:84:LYS:CE	48:BU:84:LYS:HA	2.41	0.42
50:BW:41:ILE:HG13	50:BW:42:GLN:N	2.35	0.42
53:C1:55:U:O2'	53:C1:56:U:N1	2.53	0.42
54:CA:1265:G:H2'	54:CA:1266:G:O4'	2.20	0.42
54:CA:1290:G:N3	54:CA:1290:G:H2'	2.34	0.42
54:CA:133:U:OP1	50:CW:74:LYS:NZ	2.51	0.42
54:CA:1352:C:H2'	54:CA:1353:G:H8	1.76	0.42
54:CA:1427:U:H2'	54:CA:1428:A:C8	2.55	0.42
54:CA:177:C:P	50:CW:65:LYS:HE2	2.60	0.42
54:CA:225:C:O2'	54:CA:226:G:H5'	2.19	0.42
54:CA:372:C:H2'	54:CA:372:C:O2	2.19	0.42
54:CA:390:C:O5'	54:CA:390:C:H6	2.03	0.42
54:CA:405:U:H5''	54:CA:495:A:C2	2.53	0.42
54:CA:556:C:H2'	54:CA:557:G:C5'	2.50	0.42
54:CA:568:G:O6	42:CO:5:PRO:CD	2.63	0.42
54:CA:598:U:H2'	54:CA:599:C:H6	1.84	0.42
54:CA:697:U:H2'	54:CA:698:G:H5'	2.01	0.42
54:CA:742:G:O2'	54:CA:743:U:H5'	2.20	0.42
54:CA:92:G:H5'	54:CA:92:G:H8	1.85	0.42
54:CA:92:G:H2'	54:CA:93:U:C6	2.55	0.42
32:CE:235:SER:C	32:CE:237:ALA:N	2.69	0.42
32:CE:78:GLN:O	32:CE:79:ASP:C	2.58	0.42
32:CE:86:GLU:O	32:CE:88:ALA:N	2.49	0.42
33:CF:131:ARG:CG	33:CF:131:ARG:HH11	2.30	0.42
34:CG:102:ASP:HB3	34:CG:136:PRO:CB	2.49	0.42
34:CG:157:LEU:CD1	34:CG:161:ASN:HD21	2.32	0.42
34:CG:42:GLN:HG2	34:CG:42:GLN:O	2.19	0.42
35:CH:137:GLU:N	35:CH:140:ARG:HH12	2.18	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:CH:92:LYS:O	35:CH:118:ILE:HD12	2.18	0.42
36:CI:74:ASP:O	36:CI:75:LEU:C	2.58	0.42
37:CJ:109:ASN:OD1	37:CJ:119:ARG:NH2	2.53	0.42
37:CJ:25:ALA:HA	37:CJ:28:ASN:HD22	1.84	0.42
42:CO:64:TYR:HB3	42:CO:65:GLU:H	1.55	0.42
44:CQ:36:PHE:HD1	44:CQ:37:PHE:N	2.18	0.42
44:CQ:51:GLY:C	44:CQ:53:LEU:N	2.72	0.42
46:CS:74:LEU:HB3	46:CS:80:PHE:HE1	1.84	0.42
17:D2:24:LYS:HA	17:D2:92:THR:HG23	2.01	0.42
27:D5:55:ARG:C	27:D5:57:VAL:N	2.73	0.42
30:D8:38:GLY:O	30:D8:42:ARG:HB3	2.20	0.42
55:DA:1082:U:C3'	58:DL:117:THR:CG2	2.98	0.42
55:DA:1142(A):A:C4	55:DA:1144:G:C8	3.08	0.42
55:DA:1644:C:O2	55:DA:1644:C:C2'	2.63	0.42
55:DA:165:U:H3'	55:DA:165:U:O2	2.20	0.42
55:DA:16:G:C2	55:DA:17:G:C8	3.07	0.42
55:DA:165:U:C2	55:DA:171:G:C8	3.07	0.42
55:DA:2228:G:H2'	55:DA:2229:C:C6	2.55	0.42
55:DA:2348:U:O4	55:DA:2382:G:C2	2.73	0.42
55:DA:2540:C:H2'	55:DA:2541:A:H5'	2.02	0.42
55:DA:526:A:C2	55:DA:2625:G:N3	2.88	0.42
55:DA:2814:C:C5	55:DA:2815:C:C5	3.08	0.42
54:CA:1443:G:N2	55:DA:2864:G:OP1	2.44	0.42
55:DA:695:G:H4'	55:DA:1380:G:H5'	2.02	0.42
55:DA:833:U:H2'	55:DA:834:C:H6	1.85	0.42
55:DA:846:C:O2'	55:DA:847:U:OP2	2.32	0.42
55:DA:846:C:O2'	55:DA:847:U:P	2.77	0.42
3:DD:165:ILE:HG23	3:DD:173:VAL:HG21	2.01	0.42
55:DA:1797:C:O2'	3:DD:259:THR:HB	2.19	0.42
3:DD:39:LYS:HB2	3:DD:62:TYR:HB2	2.01	0.42
4:DE:196:VAL:C	4:DE:197:ILE:CG2	2.88	0.42
4:DE:27:LEU:HD23	4:DE:27:LEU:O	2.20	0.42
5:DF:9:ILE:CD1	5:DF:125:LEU:HG	2.37	0.42
5:DF:153:SER:OG	5:DF:190:GLU:N	2.53	0.42
5:DF:117:ARG:NH2	5:DF:189:THR:O	2.53	0.42
6:DG:99:MET:HE2	6:DG:103:LEU:HD12	2.01	0.42
6:DG:7:LEU:CD2	6:DG:176:LEU:HD22	2.36	0.42
6:DG:122:PRO:HG3	6:DG:182:LYS:OXT	2.20	0.42
8:DK:128:LEU:HD13	8:DK:128:LEU:C	2.39	0.42
8:DK:56:LYS:HG3	8:DK:57:ARG:N	2.34	0.42
58:DL:112:MET:HE3	58:DL:118:THR:HA	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2562:U:C1'	10:DN:23:ARG:HH11	2.26	0.42
11:DO:1:MET:HE3	11:DO:5:ASP:HB3	2.01	0.42
15:DR:107:ASP:O	15:DR:109:GLU:N	2.53	0.42
15:DR:49:VAL:CG1	15:DR:49:VAL:O	2.68	0.42
18:DS:55:ALA:O	18:DS:56:ALA:C	2.57	0.42
19:DT:52:VAL:HG12	19:DT:52:VAL:O	2.19	0.42
21:DV:106:GLY:O	21:DV:107:THR:CG2	2.67	0.42
24:DW:71:ASN:O	24:DW:72:ALA:OXT	2.37	0.42
25:DX:16:PRO:HB2	25:DX:18:ASP:OD1	2.20	0.42
57:DY:71:LEU:CB	57:DY:113:GLN:CA	2.96	0.42
23:DZ:91:LYS:O	23:DZ:92:LYS:C	2.58	0.42
16:A1:21:ALA:O	16:A1:22:LYS:C	2.58	0.42
22:A3:65:GLY:HA3	22:A3:81:VAL:CG1	2.50	0.42
27:A5:45:VAL:CG1	27:A5:56:LYS:HG3	2.50	0.42
28:A6:36:LEU:O	28:A6:37:ARG:C	2.59	0.42
30:A8:16:ILE:HB	30:A8:65:GLU:CA	2.45	0.42
1:AA:1007:C:H5''	9:AM:35:ARG:HH11	1.85	0.42
1:AA:1309:G:C2'	1:AA:1310:G:H5'	2.49	0.42
1:AA:1848:A:C4	1:AA:1849:G:C8	3.08	0.42
1:AA:1793:C:O2	1:AA:1900:A:H2	2.03	0.42
1:AA:2287:A:C5	1:AA:2289:G:C5	3.08	0.42
1:AA:2305:A:N1	6:AG:154:GLY:N	2.61	0.42
1:AA:2759:G:N2	7:AH:139:GLN:OE1	2.52	0.42
1:AA:371:A:HO2'	1:AA:372:G:P	2.43	0.42
1:AA:404:C:C2'	1:AA:405:U:OP2	2.68	0.42
1:AA:747:U:C5	27:A5:3:LYS:HB2	2.55	0.42
1:AA:949:C:O2'	1:AA:950:G:H5'	2.20	0.42
1:AA:966:G:C6	1:AA:967:C:N4	2.88	0.42
5:AF:126:VAL:HG23	5:AF:127:GLU:N	2.34	0.42
5:AF:170:LEU:H	5:AF:170:LEU:HD12	1.85	0.42
5:AF:25:PRO:O	5:AF:26:ALA:CB	2.68	0.42
6:AG:97:ASP:O	6:AG:100:TRP:HB2	2.18	0.42
6:AG:41:GLN:HB3	6:AG:43:LEU:HD13	2.01	0.42
7:AH:53:GLU:HA	7:AH:65:HIS:CD2	2.55	0.42
8:AK:58:LEU:C	8:AK:60:GLU:N	2.73	0.42
11:AO:64:LYS:HE3	30:A8:30:ARG:NH1	2.33	0.42
12:AP:63:LYS:HB2	12:AP:63:LYS:HZ3	1.85	0.42
14:AQ:78:LEU:HD21	14:AQ:108:GLY:CA	2.50	0.42
15:AR:50:ILE:HD12	15:AR:99:LEU:CD1	2.50	0.42
15:AR:26:ASP:HB2	15:AR:90:GLN:O	2.20	0.42
19:AT:55:ASN:HB2	19:AT:80:ILE:HG12	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:AU:94:LYS:HZ2	20:AU:101:LYS:NZ	2.15	0.42
1:AA:84:A:H3'	20:AU:8:LYS:HD2	2.01	0.42
21:AV:186:GLU:O	21:AV:187:ALA:CB	2.65	0.42
21:AV:40:ASP:OD1	21:AV:42:VAL:HB	2.20	0.42
24:AW:64:LEU:O	24:AW:67:LYS:HB2	2.19	0.42
31:BA:960:U:N3	31:BA:1225:A:C8	2.78	0.42
31:BA:1297:C:O2'	31:BA:1298:C:OP2	2.38	0.42
31:BA:1426:C:H2'	31:BA:1427:U:C6	2.55	0.42
31:BA:791:G:C6	31:BA:792:A:N7	2.87	0.42
31:BA:792:A:H1'	31:BA:794:A:N7	2.35	0.42
31:BA:924:C:H2'	31:BA:925:G:H8	1.85	0.42
31:BA:934:C:O2'	31:BA:935:A:P	2.77	0.42
31:BA:1054:C:N4	52:BB:34:G:C1'	2.83	0.42
32:BE:28:PHE:CE1	32:BE:31:TYR:HB2	2.55	0.42
33:BF:34:LEU:HD12	33:BF:34:LEU:O	2.20	0.42
33:BF:63:ASN:CG	33:BF:64:VAL:H	2.23	0.42
35:BH:147:ASP:HA	35:BH:150:ARG:HB3	2.01	0.42
38:BK:102:ARG:O	38:BK:104:ARG:N	2.53	0.42
39:BL:70:LYS:O	39:BL:74:ILE:HG13	2.19	0.42
40:BM:6:ILE:HG22	40:BM:98:ILE:HG13	2.02	0.42
41:BN:92:GLU:C	41:BN:94:ALA:H	2.22	0.42
43:BP:51:ALA:O	43:BP:54:VAL:N	2.52	0.42
43:BP:81:LEU:HD13	43:BP:88:ARG:CG	2.50	0.42
31:BA:1317:C:OP1	44:BQ:17:LYS:HG2	2.20	0.42
54:CA:1028(A):C:H2'	54:CA:1028(B):C:O4'	2.19	0.42
54:CA:1054:C:O2'	54:CA:1055:A:O5'	2.37	0.42
54:CA:1176:A:N6	54:CA:1177:G:C5	2.87	0.42
54:CA:1213:A:N7	54:CA:1215:G:C6	2.87	0.42
54:CA:1363:A:H5''	54:CA:1364:U:OP1	2.20	0.42
54:CA:1469:G:H2'	54:CA:1470:G:H8	1.84	0.42
54:CA:269:C:H2'	54:CA:270:A:C8	2.55	0.42
54:CA:366:C:O2'	54:CA:367:U:O5'	2.37	0.42
54:CA:407:G:O2'	54:CA:408:A:H5'	2.20	0.42
54:CA:508:C:H4'	54:CA:509:A:O5'	2.20	0.42
54:CA:602:A:H2'	54:CA:603:U:H6	1.84	0.42
54:CA:631:G:O2'	54:CA:632:A:C4	2.72	0.42
54:CA:689:C:H2'	54:CA:690:G:C5'	2.47	0.42
54:CA:934:C:N4	54:CA:1344:C:C2	2.88	0.42
54:CA:968:A:C4'	54:CA:969:A:OP2	2.67	0.42
52:CC:44:G:H5''	52:CC:45:U:C5	2.55	0.42
52:CD:19:G:C2	55:DA:2112:G:N2	2.84	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:CF:140:ARG:CZ	33:CF:140:ARG:HB2	2.49	0.42
35:CH:14:ARG:NH1	35:CH:129:ILE:HD12	2.34	0.42
35:CH:28:PHE:O	35:CH:47:LYS:HA	2.20	0.42
38:CK:14:ARG:O	38:CK:18:ARG:HD3	2.20	0.42
39:CL:117:HIS:NE2	39:CL:123:PRO:HA	2.35	0.42
54:CA:967:C:H4'	39:CL:125:TYR:OH	2.20	0.42
41:CN:12:ARG:HG2	41:CN:13:GLN:N	2.35	0.42
54:CA:706:A:O4'	41:CN:29:ILE:HD13	2.19	0.42
43:CP:53:VAL:HG12	43:CP:57:ARG:HH12	1.84	0.42
47:CT:14:LYS:HB2	47:CT:14:LYS:NZ	2.34	0.42
50:CW:43:LEU:CA	50:CW:46:GLU:HB3	2.42	0.42
13:D0:38:VAL:CG1	13:D0:42:LYS:HE3	2.50	0.42
30:D8:29:LYS:HG3	30:D8:30:ARG:N	2.35	0.42
55:DA:1077:A:H2'	55:DA:1078:U:C5'	2.50	0.42
55:DA:1084:A:N1	55:DA:1085:A:N6	2.68	0.42
55:DA:1086:A:C5'	55:DA:1103:A:N6	2.76	0.42
55:DA:1085:A:C3'	55:DA:1086:A:N7	2.83	0.42
55:DA:1438:U:O2'	55:DA:1439:A:H5'	2.20	0.42
55:DA:1796:U:H2'	55:DA:1797:C:H6	1.78	0.42
55:DA:1830:C:H2'	55:DA:1831:G:H8	1.85	0.42
55:DA:1834:U:O2	55:DA:1970:A:C8	2.72	0.42
55:DA:271(C):U:C2'	55:DA:271:G:OP1	2.67	0.42
55:DA:2852:G:C6	55:DA:2853:C:C4	3.07	0.42
55:DA:44:A:O5'	55:DA:44:A:H8	2.03	0.42
3:DD:231:HIS:ND1	3:DD:232:PRO:HD2	2.34	0.42
4:DE:103:ASP:OD1	4:DE:201:THR:HG23	2.20	0.42
4:DE:197:ILE:HD11	4:DE:199:ARG:HH12	1.85	0.42
5:DF:59:TYR:HD1	5:DF:78:ILE:HB	1.83	0.42
6:DG:122:PRO:HG3	6:DG:182:LYS:C	2.41	0.42
7:DH:105:LEU:CD2	7:DH:113:VAL:HB	2.50	0.42
7:DH:12:PRO:CG	7:DH:13:LYS:N	2.83	0.42
7:DH:136:ILE:CG2	7:DH:136:ILE:O	2.68	0.42
7:DH:94:TYR:CD1	7:DH:94:TYR:N	2.88	0.42
56:DI:3:LEU:O	56:DI:5:ILE:N	2.53	0.42
8:DK:29:TYR:CE1	8:DK:33:ARG:NE	2.88	0.42
58:DL:78:ILE:N	58:DL:78:ILE:CD1	2.83	0.42
10:DN:101:PRO:HA	10:DN:120:GLU:O	2.20	0.42
12:DP:134:ARG:NH2	21:DV:122:ARG:NH1	2.68	0.42
12:DP:88:GLY:O	12:DP:89:ASN:C	2.56	0.42
14:DQ:19:LYS:HB3	14:DQ:20:ARG:H	1.76	0.42
18:DS:48:ALA:O	18:DS:49:LYS:C	2.58	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:DU:6:HIS:O	20:DU:7:VAL:HG13	2.19	0.42
57:DY:138:LEU:HD12	57:DY:139:VAL:C	2.37	0.42
57:DY:56:ASN:HA	57:DY:60:ARG:HG3	1.99	0.42
57:DY:71:LEU:CA	57:DY:113:GLN:CA	2.97	0.42
23:DZ:24:ALA:HA	23:DZ:32:LYS:HD3	2.01	0.42
23:DZ:82:LEU:CD2	23:DZ:82:LEU:H	2.31	0.42
13:A0:74:LYS:CE	13:A0:77:ARG:HH21	2.27	0.42
16:A1:92:ARG:NH1	17:A2:11:GLN:NE2	2.68	0.42
28:A6:10:LEU:HG	30:A8:34:TRP:CE2	2.55	0.42
28:A6:15:GLU:HG2	28:A6:16:CYS:N	2.35	0.42
1:AA:2285:C:C4	28:A6:27:LYS:HE3	2.54	0.42
1:AA:578:A:H5'	1:AA:1254:A:OP1	2.20	0.42
1:AA:137(A):G:H2'	1:AA:139:G:C8	2.55	0.42
1:AA:1396:U:O2	1:AA:1396:U:C2'	2.66	0.42
1:AA:1421:G:C2	1:AA:1422:G:C8	3.08	0.42
1:AA:1479:G:H5'	1:AA:1558:A:H2	1.85	0.42
1:AA:1642:G:O2'	1:AA:1643:G:H5'	2.20	0.42
1:AA:1698:A:O2'	1:AA:1699:G:C5'	2.54	0.42
1:AA:1726:G:C2'	1:AA:1727:U:H5'	2.50	0.42
1:AA:1863:G:H2'	1:AA:1864:U:O4'	2.20	0.42
1:AA:1956:U:C5	1:AA:1957:C:C5	3.08	0.42
1:AA:532:A:N6	1:AA:2020:A:H1'	2.35	0.42
1:AA:2087:G:C2'	1:AA:2088:G:H5'	2.50	0.42
1:AA:2247:A:H2'	1:AA:2248:C:C6	2.55	0.42
1:AA:2285:C:H5'	1:AA:2286:A:OP2	2.20	0.42
1:AA:2307:G:O2'	1:AA:2308:G:N7	2.53	0.42
1:AA:2346:A:H61	28:A6:28:ARG:HH22	1.68	0.42
1:AA:2354:G:C2	1:AA:2355:C:C6	3.07	0.42
1:AA:240:G:C2'	1:AA:257:A:H61	2.32	0.42
1:AA:792:G:H2'	1:AA:2440:C:N3	2.35	0.42
1:AA:2665:A:C2	1:AA:2666:C:C5	3.08	0.42
1:AA:2768:C:N4	1:AA:2769:C:N4	2.67	0.42
1:AA:2865:U:C4	1:AA:2866:U:N3	2.85	0.42
1:AA:2868:A:H2'	1:AA:2869:G:C8	2.55	0.42
1:AA:316:C:O2	1:AA:316:C:H2'	2.19	0.42
1:AA:528:A:H2	1:AA:2043:C:C5'	2.33	0.42
1:AA:559:G:C2'	1:AA:560:C:H5'	2.50	0.42
1:AA:708:C:H2'	1:AA:708:C:O2	2.19	0.42
1:AA:738:G:H2'	1:AA:739:G:O4'	2.20	0.42
1:AA:895:U:C2	1:AA:895:U:H3'	2.55	0.42
1:AA:964:C:O5'	1:AA:964:C:H6	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:AF:180:GLY:O	5:AF:181:LEU:C	2.57	0.42
6:AG:20:ILE:O	6:AG:24:GLY:N	2.53	0.42
6:AG:35:GLU:O	6:AG:36:LYS:HB3	2.20	0.42
6:AG:48:GLU:HA	6:AG:48:GLU:OE2	2.20	0.42
7:AH:105:LEU:HD22	7:AH:105:LEU:O	2.20	0.42
8:AK:97:ILE:N	8:AK:97:ILE:HD12	2.34	0.42
10:AN:35:VAL:HG12	10:AN:36:GLY:N	2.35	0.42
11:AO:52:GLU:HB3	11:AO:55:ARG:HD2	2.02	0.42
11:AO:59:LEU:O	11:AO:61:ARG:N	2.53	0.42
15:AR:51:ARG:NH1	15:AR:51:ARG:CG	2.82	0.42
19:AT:28:PHE:CE1	19:AT:92:LEU:HD11	2.55	0.42
19:AT:81:VAL:O	19:AT:81:VAL:HG23	2.19	0.42
20:AU:72:VAL:O	20:AU:73:ARG:CB	2.66	0.42
20:AU:84:ARG:N	20:AU:95:LYS:O	2.47	0.42
21:AV:177:PRO:O	21:AV:178:GLU:CG	2.68	0.42
21:AV:52:SER:C	21:AV:53:ILE:HG13	2.41	0.42
24:AW:49:LYS:O	24:AW:52:ASP:HB3	2.20	0.42
31:BA:1004:A:C2'	31:BA:1005:A:O4'	2.68	0.42
31:BA:1032(A):G:H2'	31:BA:1032(B):G:H8	1.85	0.42
31:BA:119:A:H4'	31:BA:120:A:O5'	2.20	0.42
31:BA:1357:A:N6	31:BA:1363:A:C2	2.88	0.42
31:BA:141:A:H1'	31:BA:182:U:O2	2.20	0.42
31:BA:1476:G:H2'	31:BA:1477:C:C6	2.55	0.42
31:BA:1405:G:H1'	31:BA:1519:A:O4'	2.19	0.42
31:BA:445:G:C6	31:BA:490:G:C6	3.08	0.42
31:BA:447:G:H3'	31:BA:485:G:N2	2.34	0.42
31:BA:500:G:H2'	31:BA:501:C:H6	1.83	0.42
31:BA:555:C:H2'	31:BA:556:C:H6	1.84	0.42
31:BA:659:U:H2'	31:BA:660:G:O4'	2.20	0.42
31:BA:7:G:H5'	31:BA:298:A:H5'	2.01	0.42
31:BA:913:A:H1'	31:BA:914:A:C1'	2.50	0.42
31:BA:924:C:H2'	31:BA:925:G:C8	2.55	0.42
31:BA:973:G:N3	40:BM:55:LYS:CE	2.83	0.42
52:BB:67:C:H2'	52:BB:68:C:C6	2.55	0.42
32:BE:16:HIS:HD2	32:BE:210:SER:HA	1.84	0.42
32:BE:236:TYR:C	32:BE:236:TYR:CD1	2.93	0.42
35:BH:24:ARG:HG3	35:BH:26:PHE:CE2	2.55	0.42
45:BR:41:GLU:O	45:BR:44:LYS:HB2	2.20	0.42
46:BS:15:PRO:HB2	46:BS:41:PRO:HG3	2.02	0.42
49:BV:20:LEU:O	49:BV:23:ASN:HB3	2.20	0.42
49:BV:63:THR:HG23	49:BV:66:MET:HG3	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
50:BW:44:ALA:HB1	50:BW:91:LEU:CB	2.50	0.42
50:BW:82:SER:O	50:BW:86:ARG:HB2	2.20	0.42
54:CA:60:A:N6	54:CA:110:C:N3	2.64	0.42
54:CA:1125:U:C2'	54:CA:1126:U:OP2	2.68	0.42
54:CA:1535:C:H2'	54:CA:1536:C:O4'	2.20	0.42
54:CA:432:A:C8	54:CA:433:C:C5	3.08	0.42
54:CA:481:G:O2'	54:CA:482:A:O5'	2.38	0.42
54:CA:533:A:C5	54:CA:536:C:C4	3.08	0.42
54:CA:537:G:H2'	54:CA:538:G:C8	2.55	0.42
54:CA:638:G:O2'	54:CA:639:G:H5'	2.20	0.42
54:CA:949:A:C4'	54:CA:1364:U:O4	2.67	0.42
52:CB:56:C:C5	52:CB:57:G:N7	2.88	0.42
52:CC:19:G:O6	6:DG:83:ARG:NH2	2.53	0.42
52:CD:20:U:C2'	52:CD:20:U:O2	2.68	0.42
52:CD:2:C:O2'	52:CD:3:C:P	2.78	0.42
33:CF:172:ARG:O	33:CF:173:VAL:CG2	2.68	0.42
34:CG:166:LYS:HE2	34:CG:178:VAL:HG11	2.01	0.42
37:CJ:20:ASP:OD1	37:CJ:23:VAL:HG23	2.20	0.42
38:CK:49:GLU:HG3	38:CK:60:ARG:HB2	2.01	0.42
39:CL:9:ARG:CB	39:CL:14:VAL:HG22	2.49	0.42
42:CO:20:LYS:HB2	42:CO:21:LYS:H	1.65	0.42
43:CP:2:ALA:O	43:CP:4:ILE:HG12	2.20	0.42
43:CP:82:MET:O	43:CP:83:ASP:HB2	2.19	0.42
54:CA:376:G:C4'	46:CS:5:ARG:HD2	2.49	0.42
47:CT:13:ASP:C	47:CT:15:MET:N	2.73	0.42
48:CU:73:ALA:HB3	48:CU:79:LEU:HD12	2.02	0.42
50:CW:38:LYS:O	50:CW:39:LYS:C	2.58	0.42
16:D1:97:ASP:OD1	16:D1:101:ARG:NH1	2.53	0.42
55:DA:1188:U:C4'	17:D2:79:VAL:CG2	2.98	0.42
26:D4:53:GLU:O	26:D4:57:GLU:CG	2.68	0.42
26:D4:9:LEU:C	26:D4:9:LEU:HD23	2.41	0.42
28:D6:41:PRO:HD2	28:D6:46:HIS:HA	2.01	0.42
55:DA:1048:A:OP2	55:DA:1048:A:H8	2.02	0.42
55:DA:1128:A:N7	55:DA:2489:G:O2'	2.53	0.42
55:DA:1177:A:H4'	55:DA:1178:C:C5'	2.37	0.42
55:DA:1485:G:C2'	55:DA:1486:A:O5'	2.67	0.42
55:DA:1558:A:O2'	55:DA:1559:G:P	2.78	0.42
55:DA:1906:G:N1	55:DA:1925:C:O2	2.53	0.42
55:DA:2347:C:P	28:D6:39:TYR:OH	2.76	0.42
55:DA:2572:A:HO2'	55:DA:2573:C:P	2.43	0.42
55:DA:2612:C:C5	55:DA:2613:U:H5	2.38	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:506:G:H5'	55:DA:509:C:H1'	2.02	0.42
55:DA:53:A:H2'	55:DA:54:G:O4'	2.20	0.42
55:DA:802:A:C2'	55:DA:803:U:C5'	2.98	0.42
55:DA:953:A:N1	55:DA:964:C:O2	2.53	0.42
4:DE:50:GLY:H	4:DE:77:ILE:HA	1.82	0.42
5:DF:134:GLY:H	5:DF:162:LEU:CD2	2.31	0.42
5:DF:57:VAL:HG12	5:DF:58:ALA:N	2.35	0.42
6:DG:115:ARG:HG2	6:DG:115:ARG:NH1	2.34	0.42
7:DH:139:GLN:HG3	7:DH:140:LYS:N	2.35	0.42
58:DL:76:TYR:CD2	58:DL:77:LEU:CD1	3.03	0.42
9:DM:75:TYR:CD2	9:DM:76:SER:O	2.72	0.42
10:DN:48:PRO:O	10:DN:49:ARG:HG2	2.19	0.42
11:DO:13:ASN:C	11:DO:15:ARG:N	2.70	0.42
11:DO:31:ALA:C	11:DO:32:THR:HG22	2.40	0.42
55:DA:2392:A:H8	11:DO:60:MET:HG3	1.77	0.42
12:DP:2:LEU:CB	12:DP:70:PRO:CG	2.95	0.42
55:DA:1266:G:C5	18:DS:15:ARG:NH1	2.87	0.42
21:DV:108:PRO:HG2	21:DV:109:ALA:N	2.35	0.42
21:DV:20:ARG:C	21:DV:22:GLY:N	2.73	0.42
57:DY:137:GLU:C	57:DY:138:LEU:O	2.55	0.42
23:DZ:20:ARG:HH11	23:DZ:20:ARG:HG2	1.84	0.42
13:A0:18:LEU:HD11	13:A0:22:ARG:CZ	2.49	0.41
1:AA:2880:C:O2'	13:A0:90:ARG:HD3	2.20	0.41
16:A1:100:VAL:C	16:A1:102:GLU:H	2.23	0.41
16:A1:92:ARG:CB	17:A2:11:GLN:NE2	2.77	0.41
17:A2:61:VAL:CG1	17:A2:62:LEU:N	2.79	0.41
30:A8:14:VAL:HG13	30:A8:23:VAL:O	2.20	0.41
30:A8:52:LYS:N	30:A8:53:PRO:CD	2.83	0.41
30:A8:4:MET:CE	30:A8:61:LEU:HD23	2.49	0.41
1:AA:1027:A:N7	1:AA:1126:A:C2	2.88	0.41
1:AA:1608:A:O2'	1:AA:1610:A:P	2.78	0.41
1:AA:1869:G:H5'	1:AA:1870:C:P	2.60	0.41
1:AA:2072:G:C5	1:AA:2073:C:C5	3.08	0.41
1:AA:2106:G:O2'	1:AA:2107:C:H5'	2.20	0.41
1:AA:2306:C:O5'	1:AA:2307:G:C5'	2.68	0.41
1:AA:2391:G:O6	1:AA:2425:A:C8	2.66	0.41
1:AA:2880:C:O2	13:A0:93:GLY:N	2.49	0.41
1:AA:434:U:H2'	1:AA:436:C:H41	1.85	0.41
1:AA:846:C:H1'	1:AA:847:U:C6	2.54	0.41
1:AA:868:U:H2'	1:AA:869:G:O4'	2.20	0.41
2:AB:41:U:C4	6:AG:70:VAL:HG23	2.55	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:AD:28:GLU:CB	3:AD:29:PRO:CD	2.98	0.41
1:AA:606:U:OP2	5:AF:104:LYS:HE3	2.20	0.41
5:AF:8:GLN:HE22	5:AF:127:GLU:HB3	1.84	0.41
5:AF:24:LEU:O	5:AF:25:PRO:C	2.58	0.41
5:AF:93:LYS:O	5:AF:94:PRO:C	2.58	0.41
6:AG:41:GLN:N	6:AG:90:LEU:O	2.49	0.41
7:AH:101:ARG:HG3	7:AH:117:PRO:CG	2.47	0.41
12:AP:134:ARG:HH11	12:AP:134:ARG:HG2	1.83	0.41
12:AP:35:VAL:CG1	12:AP:130:LYS:HB3	2.49	0.41
12:AP:54:MET:O	12:AP:57:HIS:N	2.53	0.41
15:AR:132:LYS:O	15:AR:136:GLN:HB3	2.20	0.41
19:AT:47:PHE:HD2	19:AT:89:ILE:CG2	2.33	0.41
21:AV:133:ILE:HA	21:AV:134:PRO:HD2	1.79	0.41
21:AV:120:ILE:CG2	21:AV:171:ILE:H	2.33	0.41
21:AV:58:VAL:O	21:AV:67:LEU:O	2.38	0.41
24:AW:9:GLN:CA	24:AW:12:GLU:HB3	2.48	0.41
24:AW:32:LEU:O	24:AW:33:MET:C	2.56	0.41
23:AZ:56:GLN:HB3	23:AZ:56:GLN:HE21	1.64	0.41
31:BA:1014:A:H4'	49:BV:14:HIS:CD2	2.55	0.41
31:BA:1131:G:O2'	31:BA:1132:C:H5'	2.20	0.41
31:BA:1145:C:H4'	31:BA:1146:A:H5'	2.02	0.41
31:BA:1178:G:C8	31:BA:1180:A:OP2	2.73	0.41
31:BA:1215:G:H2'	31:BA:1216:G:H5'	2.02	0.41
31:BA:1250:A:H4'	39:BL:67:GLY:HA2	2.01	0.41
31:BA:1429:C:O2'	31:BA:1430:C:H5'	2.20	0.41
31:BA:1413:A:C2	31:BA:1488:G:C2	3.08	0.41
31:BA:1535:C:H2'	31:BA:1536:C:C5'	2.50	0.41
31:BA:189:U:C4	47:BT:72:ARG:NH1	2.77	0.41
31:BA:113:G:O4'	31:BA:354:G:H4'	2.20	0.41
31:BA:468:A:H2'	31:BA:474:G:C5'	2.50	0.41
31:BA:538:G:O2'	31:BA:539:A:H5'	2.20	0.41
31:BA:542:G:H2'	31:BA:543:C:C6	2.55	0.41
31:BA:668:G:H5'	45:BR:49:ASP:HA	2.01	0.41
31:BA:728:A:C2	31:BA:729:A:C5	3.08	0.41
31:BA:991:U:OP2	31:BA:991:U:H6	2.03	0.41
32:BE:105:PHE:O	32:BE:107:THR:N	2.53	0.41
32:BE:224:GLN:OE1	32:BE:225:ALA:N	2.53	0.41
34:BG:15:GLU:O	34:BG:16:GLY:C	2.57	0.41
34:BG:178:VAL:O	34:BG:179:GLU:CB	2.68	0.41
34:BG:13:ARG:HD2	34:BG:38:TYR:O	2.20	0.41
31:BA:511:C:C1'	34:BG:43:HIS:NE2	2.81	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:BL:89:ASN:HB2	39:BL:92:TYR:HB2	2.01	0.41
41:BN:29:ILE:HG13	41:BN:29:ILE:O	2.10	0.41
43:BP:110:ARG:NH1	43:BP:110:ARG:HG2	2.35	0.41
45:BR:29:VAL:HG11	45:BR:67:LEU:HD21	2.02	0.41
46:BS:83:GLU:HB3	46:BS:84:ALA:H	1.63	0.41
48:BU:62:GLU:HA	48:BU:65:ILE:HG13	2.01	0.41
54:CA:1093:A:O5'	54:CA:1093:A:H8	2.03	0.41
54:CA:115:G:H4'	54:CA:116:A:O5'	2.20	0.41
54:CA:1194:U:H2'	54:CA:1195:C:C6	2.55	0.41
54:CA:1342:C:H2'	54:CA:1343:G:C8	2.54	0.41
54:CA:262:A:N1	54:CA:263:A:C6	2.87	0.41
54:CA:280:C:C4'	54:CA:281:G:OP2	2.63	0.41
54:CA:386:C:C2'	54:CA:387:U:H5'	2.50	0.41
54:CA:408:A:H4'	34:CG:112:VAL:HG11	2.01	0.41
54:CA:517:G:H4'	54:CA:518:C:O5'	2.20	0.41
54:CA:652:U:O2'	54:CA:653:A:O5'	2.38	0.41
54:CA:66:G:H5'	54:CA:67:C:OP2	2.19	0.41
52:CD:16:U:O2	52:CD:18:G:H5'	2.20	0.41
33:CF:7:PRO:O	33:CF:11:ARG:HG2	2.20	0.41
33:CF:124:ILE:C	33:CF:126:ARG:N	2.72	0.41
54:CA:1112:C:H1'	33:CF:179:ARG:NH1	2.35	0.41
34:CG:166:LYS:HE3	34:CG:166:LYS:HB2	1.88	0.41
34:CG:170:VAL:HG13	34:CG:171:GLY:N	2.35	0.41
54:CA:1240:U:OP2	37:CJ:116:ALA:CB	2.68	0.41
37:CJ:151:TYR:OH	41:CN:54:ARG:HD3	2.20	0.41
37:CJ:16:LEU:CD1	39:CL:45:ALA:HB2	2.49	0.41
38:CK:104:ARG:O	38:CK:106:GLY:N	2.47	0.41
38:CK:18:ARG:NH2	38:CK:81:HIS:O	2.53	0.41
39:CL:22:GLY:O	39:CL:24:GLY:N	2.52	0.41
40:CM:15:THR:O	40:CM:16:LEU:C	2.58	0.41
40:CM:31:GLY:HA3	40:CM:81:THR:HG21	2.02	0.41
46:CS:6:LEU:HG	46:CS:17:TYR:CB	2.46	0.41
47:CT:68:ARG:HG2	47:CT:68:ARG:HH11	1.84	0.41
49:CV:69:HIS:HB3	49:CV:73:GLU:CD	2.40	0.41
6:DG:145:THR:CG2	26:D4:28:LYS:NZ	2.82	0.41
26:D4:16:CYS:HA	26:D4:33:VAL:O	2.20	0.41
28:D6:42:TRP:O	28:D6:43:CYS:HB3	2.19	0.41
30:D8:28:GLY:O	30:D8:29:LYS:C	2.58	0.41
30:D8:6:THR:O	30:D8:59:LYS:O	2.38	0.41
55:DA:85:G:N3	55:DA:103:A:C2	2.88	0.41
55:DA:1316:U:H2'	55:DA:1317:A:C8	2.55	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:182:A:H1'	55:DA:433:C:O2'	2.20	0.41
55:DA:1934:C:C2'	55:DA:1935:G:O5'	2.68	0.41
55:DA:2012:G:O3'	18:DS:96:ILE:HG13	2.20	0.41
55:DA:2135:A:N3	55:DA:2135:A:H2'	2.34	0.41
55:DA:2328:A:H2'	55:DA:2329:G:O4'	2.20	0.41
55:DA:2556:C:H2'	55:DA:2557:G:O4'	2.19	0.41
55:DA:2656:U:H3'	55:DA:2656:U:C6	2.55	0.41
55:DA:2723:C:H4'	13:D0:1:MET:CB	2.50	0.41
55:DA:51:G:H1'	55:DA:118:A:N6	2.35	0.41
55:DA:800:A:C4'	55:DA:801:G:O5'	2.66	0.41
3:DD:136:ILE:HA	3:DD:137:PRO:HD3	1.82	0.41
3:DD:35:LYS:CB	3:DD:63:ARG:HA	2.43	0.41
3:DD:79:VAL:HG11	3:DD:111:LEU:HD11	2.01	0.41
4:DE:54:GLN:CD	4:DE:54:GLN:N	2.73	0.41
5:DF:129:PHE:HA	5:DF:142:TRP:CD1	2.55	0.41
5:DF:107:LYS:CE	5:DF:206:ILE:HD13	2.50	0.41
6:DG:102:PHE:HZ	6:DG:157:ILE:HD13	1.85	0.41
6:DG:143:GLU:CG	26:D4:31:ILE:HD11	2.49	0.41
7:DH:107:VAL:HG23	7:DH:109:PHE:CE1	2.55	0.41
7:DH:84:SER:O	7:DH:133:VAL:O	2.38	0.41
56:DI:23:LEU:N	56:DI:23:LEU:CD1	2.79	0.41
56:DI:3:LEU:CD2	56:DI:7:ARG:CD	2.75	0.41
8:DK:67:ARG:HD2	8:DK:68:LEU:HD13	2.01	0.41
55:DA:1082:U:O2'	58:DL:115:LEU:HD11	2.20	0.41
58:DL:21:PRO:C	58:DL:24:GLY:C	2.79	0.41
58:DL:19:PRO:CA	58:DL:25:PRO:HD3	2.41	0.41
58:DL:78:ILE:HD12	58:DL:78:ILE:N	2.34	0.41
11:DO:112:LEU:HD13	11:DO:127:ALA:HB1	2.02	0.41
11:DO:131:SER:HB3	11:DO:134:ALA:HB2	2.02	0.41
11:DO:71:VAL:HG12	11:DO:72:PRO:HD3	1.99	0.41
11:DO:95:VAL:O	11:DO:96:THR:CG2	2.68	0.41
14:DQ:88:ASP:CG	14:DQ:89:ARG:H	2.23	0.41
20:DU:33:LYS:HG3	20:DU:34:LYS:N	2.33	0.41
20:DU:65:ALA:HA	20:DU:66:PRO:HD3	1.93	0.41
20:DU:88:LYS:C	20:DU:90:LEU:N	2.73	0.41
21:DV:135:GLU:HG3	21:DV:136:PHE:CD2	2.55	0.41
21:DV:177:PRO:O	21:DV:178:GLU:CG	2.68	0.41
57:DY:1:MET:CE	57:DY:3:ASN:HD21	2.32	0.41
16:A1:76:TYR:OH	16:A1:93:LYS:NZ	2.51	0.41
16:A1:90:VAL:O	16:A1:91:ASP:C	2.58	0.41
16:A1:95:LEU:HD12	17:A2:11:GLN:CB	2.47	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:A4:61:ARG:CG	26:A4:62:ARG:HH22	2.33	0.41
30:A8:47:LYS:O	30:A8:48:PHE:CB	2.64	0.41
1:AA:1087:G:N2	1:AA:1090:U:C5	2.89	0.41
1:AA:1228:G:OP2	16:A1:16:LYS:NZ	2.52	0.41
1:AA:1261:C:C2'	1:AA:1262:A:O5'	2.68	0.41
1:AA:1341:U:H2'	1:AA:1397:U:O2	2.20	0.41
1:AA:1417:C:H2'	1:AA:1418:G:H5'	2.02	0.41
1:AA:1419:A:O2'	1:AA:1420:U:C5	2.73	0.41
1:AA:1342:A:N1	1:AA:1602:U:O2	2.53	0.41
1:AA:1791:A:H3'	1:AA:1792:G:H8	1.85	0.41
1:AA:1932:A:N6	1:AA:1968:G:H1'	2.35	0.41
1:AA:528:A:C2	1:AA:2043:C:C5'	3.02	0.41
1:AA:2320:A:O2'	1:AA:2321:G:C2	2.70	0.41
1:AA:242:G:O2'	1:AA:243:U:C5	2.73	0.41
1:AA:270:A:OP2	1:AA:270(Y):G:N2	2.46	0.41
1:AA:2756:U:C1'	1:AA:2757:A:C8	3.04	0.41
1:AA:607:U:C5'	5:AF:103:LYS:HD2	2.50	0.41
1:AA:807:U:H2'	1:AA:808:G:H8	1.84	0.41
1:AA:881:G:H1	1:AA:895:U:H2'	1.86	0.41
1:AA:887:A:C2'	1:AA:888:C:OP1	2.68	0.41
1:AA:961:C:H5	1:AA:2456:C:C4'	2.33	0.41
2:AB:96:G:N1	2:AB:97:G:C8	2.88	0.41
4:AE:31:CYS:O	4:AE:32:PRO:O	2.37	0.41
4:AE:34:VAL:HG11	4:AE:64:LYS:CD	2.41	0.41
5:AF:203:GLN:O	5:AF:206:ILE:O	2.37	0.41
7:AH:10:PRO:O	7:AH:11:VAL:HG13	2.19	0.41
7:AH:12:PRO:O	7:AH:15:VAL:HG22	2.20	0.41
8:AK:12:LEU:HD12	8:AK:12:LEU:HA	1.78	0.41
8:AK:58:LEU:O	8:AK:60:GLU:N	2.53	0.41
10:AN:63:VAL:HG22	10:AN:83:ALA:O	2.19	0.41
11:AO:114:ILE:HG23	11:AO:127:ALA:HB2	2.01	0.41
11:AO:111:ARG:HA	11:AO:128:HIS:CE1	2.56	0.41
12:AP:86:GLY:O	12:AP:87:LYS:C	2.56	0.41
14:AQ:67:ARG:CZ	14:AQ:67:ARG:HB2	2.49	0.41
24:AW:30:ARG:NH1	24:AW:30:ARG:CG	2.83	0.41
23:AZ:56:GLN:H	23:AZ:56:GLN:CD	2.23	0.41
31:BA:1081:G:N2	31:BA:1082:G:H1'	2.35	0.41
31:BA:1216:G:C4	31:BA:1217:C:C5	3.08	0.41
31:BA:1329:A:OP1	43:BP:25:ILE:O	2.38	0.41
31:BA:1437:C:H2'	31:BA:1438:G:H8	1.85	0.41
31:BA:1452:C:H2'	31:BA:1453:G:OP2	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:243:A:H4'	31:BA:244:U:C5'	2.50	0.41
31:BA:991:U:O4	31:BA:1212:U:O2'	2.35	0.41
52:BC:18:G:C4	52:BC:58:A:C2	3.08	0.41
34:BG:107:ARG:HB3	34:BG:174:LEU:CD1	2.50	0.41
34:BG:23:GLY:C	34:BG:24:GLU:CG	2.84	0.41
35:BH:78:HIS:O	35:BH:93:PRO:HD3	2.20	0.41
36:BI:75:LEU:HD21	36:BI:79:LEU:CD1	2.50	0.41
37:BJ:36:LYS:O	37:BJ:39:ALA:N	2.51	0.41
38:BK:1:MET:HE2	38:BK:1:MET:N	2.35	0.41
31:BA:878:G:H5''	38:BK:89:PRO:HG2	2.02	0.41
45:BR:64:ARG:NH1	45:BR:68:ARG:NH2	2.65	0.41
46:BS:21:VAL:O	46:BS:33:ILE:HB	2.20	0.41
54:CA:1032(B):G:C6	54:CA:1033:G:N7	2.89	0.41
54:CA:1353:G:H2'	54:CA:1354:C:H6	1.84	0.41
54:CA:1442:G:C6	54:CA:1446:A:N6	2.87	0.41
54:CA:310:G:O2'	54:CA:311:C:H5'	2.20	0.41
54:CA:377:G:OP1	46:CS:3:LYS:CD	2.68	0.41
54:CA:653:A:H1'	38:CK:56:LYS:CD	2.47	0.41
54:CA:684:A:O2'	54:CA:685:G:H5'	2.19	0.41
54:CA:900:A:H2'	54:CA:901:A:C8	2.55	0.41
52:CB:69:G:C2'	52:CB:70:G:OP1	2.67	0.41
52:CD:14:A:C6	52:CD:15:G:C6	3.07	0.41
32:CE:168:THR:HG21	32:CE:191:ASP:CG	2.40	0.41
32:CE:44:LEU:HA	32:CE:47:THR:OG1	2.20	0.41
33:CF:25:GLY:C	33:CF:27:LYS:H	2.24	0.41
34:CG:158:ILE:HG22	34:CG:159:ARG:N	2.34	0.41
34:CG:206:PHE:HD2	34:CG:207:TYR:CD1	2.38	0.41
35:CH:100:VAL:O	35:CH:107:ARG:NH2	2.53	0.41
35:CH:101:ILE:N	35:CH:101:ILE:CD1	2.81	0.41
35:CH:59:GLY:O	35:CH:60:TYR:C	2.58	0.41
37:CJ:67:GLU:HA	37:CJ:67:GLU:OE2	2.20	0.41
40:CM:12:ASP:OD2	40:CM:12:ASP:C	2.58	0.41
42:CO:60:LEU:HD21	42:CO:64:TYR:HB2	2.01	0.41
43:CP:87:TYR:C	43:CP:89:GLY:H	2.23	0.41
44:CQ:8:GLU:O	44:CQ:10:ALA:N	2.54	0.41
45:CR:37:ASN:N	45:CR:37:ASN:ND2	2.68	0.41
54:CA:624:C:H5''	46:CS:10:GLY:O	2.20	0.41
46:CS:20:VAL:HG21	46:CS:32:TYR:CB	2.49	0.41
13:D0:27:SER:O	13:D0:30:THR:HB	2.20	0.41
13:D0:3:HIS:C	13:D0:5:LYS:H	2.15	0.41
17:D2:22:VAL:HG12	17:D2:23:GLU:N	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:D2:65:GLY:HA3	17:D2:91:TYR:CZ	2.55	0.41
22:D3:36:ILE:CD1	22:D3:36:ILE:H	2.33	0.41
22:D3:37:LEU:N	22:D3:59:LEU:O	2.47	0.41
28:D6:34:LEU:HD13	28:D6:34:LEU:N	2.25	0.41
28:D6:44:ARG:O	28:D6:45:LYS:CG	2.69	0.41
30:D8:6:THR:HG22	30:D8:63:PRO:HD3	2.03	0.41
55:DA:1087:G:C2	55:DA:1089:G:O2'	2.73	0.41
55:DA:1474:C:O2	55:DA:1474:C:H2'	2.21	0.41
55:DA:1659:U:H2'	55:DA:1660:C:O4'	2.20	0.41
55:DA:192:C:C2'	55:DA:193:U:H5'	2.48	0.41
55:DA:2258:C:H4'	55:DA:2259:G:OP2	2.16	0.41
55:DA:2309:A:O5'	55:DA:2309:A:H8	2.03	0.41
55:DA:2389:G:H5''	55:DA:2390:U:O4'	2.19	0.41
55:DA:2059:A:N6	55:DA:2503:A:H2'	2.35	0.41
55:DA:2507:C:H2'	55:DA:2508:G:O5'	2.20	0.41
55:DA:2570:G:H2'	55:DA:2571:C:C6	2.56	0.41
55:DA:2785:C:C4'	4:DE:35:GLN:HE22	2.32	0.41
55:DA:2853:C:O2'	55:DA:2854:G:H5'	2.19	0.41
55:DA:332:A:O2'	55:DA:333:G:P	2.78	0.41
55:DA:502:A:C2'	55:DA:503:A:H5'	2.49	0.41
55:DA:654(B):C:N3	55:DA:654(T):A:C2	2.88	0.41
55:DA:806:C:C5	11:DO:41:ARG:NH2	2.87	0.41
55:DA:943:U:OP1	11:DO:34:GLY:O	2.38	0.41
2:DB:51:G:H2'	2:DB:52:A:C1'	2.50	0.41
3:DD:252:TRP:O	3:DD:252:TRP:CE3	2.73	0.41
55:DA:1814:G:H4'	3:DD:51:VAL:HG21	2.02	0.41
3:DD:65:ILE:HG13	3:DD:67:PHE:CE1	2.54	0.41
5:DF:129:PHE:CD2	5:DF:163:VAL:HG21	2.55	0.41
5:DF:149:ASP:OD1	5:DF:149:ASP:N	2.49	0.41
6:DG:180:PHE:O	6:DG:182:LYS:N	2.53	0.41
6:DG:63:ILE:CG2	6:DG:144:ILE:HD11	2.50	0.41
6:DG:83:ARG:HG2	6:DG:83:ARG:HH11	1.85	0.41
7:DH:167:GLU:HA	7:DH:168:PRO:HD2	1.94	0.41
56:DI:1:MET:O	56:DI:5:ILE:CD1	2.60	0.41
58:DL:128:ALA:HB1	58:DL:132:ARG:NE	2.35	0.41
58:DL:78:ILE:HG22	58:DL:79:ARG:N	2.35	0.41
9:DM:41:ASP:OD1	9:DM:41:ASP:N	2.53	0.41
10:DN:7:TYR:CE1	10:DN:20:MET:CE	3.02	0.41
11:DO:11:GLY:O	11:DO:12:ALA:HB2	2.19	0.41
11:DO:1:MET:CE	11:DO:5:ASP:HB3	2.49	0.41
11:DO:95:VAL:C	11:DO:96:THR:HG23	2.40	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:DQ:46:VAL:CG1	14:DQ:47:THR:N	2.82	0.41
14:DQ:86:ALA:O	14:DQ:87:PHE:CB	2.64	0.41
15:DR:113:LYS:HD2	15:DR:113:LYS:HA	1.83	0.41
15:DR:118:ARG:HH21	15:DR:121:ILE:HG21	1.85	0.41
21:DV:112:ARG:H	21:DV:112:ARG:HD2	1.79	0.41
21:DV:182:LYS:CB	21:DV:183:LEU:HD23	2.50	0.41
17:A2:13:ARG:HG2	17:A2:13:ARG:HH11	1.84	0.41
26:A4:16:CYS:HB3	26:A4:19:GLY:H	1.85	0.41
1:AA:2285:C:N4	28:A6:25:LYS:HE3	2.36	0.41
1:AA:1249:U:H4'	16:A1:4:ALA:HB3	2.02	0.41
1:AA:1272:A:H3'	1:AA:1273:U:H5'	2.02	0.41
1:AA:1337:G:C2	1:AA:1338:G:C4	3.08	0.41
1:AA:13:A:O2'	1:AA:15:G:C5	2.57	0.41
1:AA:1654:A:H2	4:AE:113:PHE:CE2	2.38	0.41
1:AA:1906:G:N2	1:AA:1925:C:C2	2.88	0.41
1:AA:2336:A:H4'	1:AA:2337:G:OP1	2.18	0.41
1:AA:2415:G:H2'	1:AA:2416:C:H6	1.84	0.41
1:AA:2418:A:H2'	1:AA:2419:U:C6	2.55	0.41
1:AA:2246:G:C2	1:AA:2426:A:H1'	2.55	0.41
1:AA:244:A:C2	1:AA:255:A:C4	3.09	0.41
1:AA:273(C):C:H3'	1:AA:273(D):C:C6	2.54	0.41
1:AA:2776:A:H4'	1:AA:2777:G:C5'	2.49	0.41
1:AA:2798:C:H5	1:AA:2799:A:C6	2.38	0.41
1:AA:2839:G:H2'	1:AA:2840:C:H6	1.85	0.41
1:AA:300:A:OP2	20:AU:84:ARG:NH1	2.54	0.41
1:AA:637:A:OP1	11:AO:133:SER:HB3	2.21	0.41
1:AA:638:G:C5	1:AA:639:U:C4	3.09	0.41
1:AA:706:A:C2	1:AA:707:G:H1'	2.55	0.41
1:AA:807:U:C2	1:AA:808:G:C8	3.08	0.41
2:AB:86:G:C2'	2:AB:87:G:H5'	2.50	0.41
3:AD:33:LEU:HD23	3:AD:34:VAL:N	2.36	0.41
4:AE:178:GLU:H	4:AE:178:GLU:HG3	1.50	0.41
1:AA:2788:C:OP1	4:AE:61:ARG:NH1	2.53	0.41
5:AF:116:ASP:O	5:AF:120:GLU:HG3	2.20	0.41
6:AG:122:PRO:HG2	6:AG:123:ASN:ND2	2.35	0.41
7:AH:163:TYR:CD1	7:AH:163:TYR:N	2.77	0.41
8:AK:126:TYR:N	8:AK:126:TYR:CD1	2.88	0.41
8:AK:88:ILE:HG22	8:AK:90:GLY:N	2.35	0.41
9:AM:13:TRP:N	9:AM:13:TRP:CD1	2.88	0.41
9:AM:26:LEU:HD23	9:AM:99:LEU:CD2	2.51	0.41
10:AN:59:LYS:O	10:AN:86:ILE:HA	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:AP:56:ARG:NH2	52:BB:52:G:O3'	2.54	0.41
12:AP:89:ASN:C	12:AP:91:GLU:N	2.74	0.41
14:AQ:76:LYS:O	14:AQ:79:ALA:HB3	2.21	0.41
1:AA:494:G:O5'	18:AS:8:ARG:NH1	2.53	0.41
1:AA:138:G:N2	19:AT:50:LYS:NZ	2.69	0.41
20:AU:50:ARG:CB	20:AU:53:PRO:HG3	2.42	0.41
20:AU:99:CYS:SG	20:AU:100:ALA:N	2.90	0.41
21:AV:97:GLU:CB	21:AV:125:LEU:HD11	2.28	0.41
21:AV:94:GLU:HB3	21:AV:95:PRO:HD2	2.02	0.41
23:AZ:91:LYS:O	23:AZ:92:LYS:O	2.39	0.41
31:BA:1098:C:C2	31:BA:1099:G:C8	3.08	0.41
31:BA:1118:C:H5''	39:BL:104:ARG:HG3	2.02	0.41
31:BA:1248:A:N3	39:BL:70:LYS:NZ	2.63	0.41
31:BA:1316:G:C3'	31:BA:1317:C:H5''	2.50	0.41
31:BA:1324:A:O4'	31:BA:1362:C:H4'	2.21	0.41
31:BA:15:G:C4	31:BA:16:A:C8	3.08	0.41
31:BA:255:G:O3'	47:BT:17:LYS:HD2	2.20	0.41
31:BA:543:C:OP1	34:BG:14:ARG:HD2	2.20	0.41
31:BA:5:U:H4'	31:BA:6:G:C4	2.54	0.41
31:BA:830:G:H2'	31:BA:831:U:C6	2.55	0.41
31:BA:982:U:H5''	31:BA:983:A:OP1	2.20	0.41
52:BC:42:C:C3'	52:BC:43:C:H5''	2.49	0.41
33:BF:114:PRO:O	33:BF:118:GLN:OE1	2.37	0.41
33:BF:44:GLU:HA	33:BF:52:LEU:HD11	2.01	0.41
35:BH:41:VAL:HG22	35:BH:113:ALA:HA	2.02	0.41
35:BH:80:ILE:HG22	38:BK:104:ARG:HE	1.82	0.41
36:BI:15:ASP:O	36:BI:19:LEU:HB2	2.21	0.41
36:BI:1:MET:CE	36:BI:68:PRO:HD3	2.50	0.41
38:BK:34:GLU:OE1	38:BK:34:GLU:HA	2.20	0.41
40:BM:39:PRO:HB3	40:BM:70:ARG:HH12	1.85	0.41
40:BM:3:LYS:N	40:BM:75:ILE:HA	2.36	0.41
45:BR:53:HIS:O	45:BR:56:LEU:HB3	2.20	0.41
45:BR:56:LEU:O	45:BR:60:VAL:HG23	2.20	0.41
47:BT:40:LYS:HD3	47:BT:42:TYR:CZ	2.55	0.41
53:C1:46:U:H2'	53:C1:47:U:O4'	2.20	0.41
54:CA:1003:G:C3'	54:CA:1004:A:H5'	2.47	0.41
54:CA:1297:C:C2'	37:CJ:114:ARG:NH2	2.81	0.41
54:CA:1447:G:OP2	54:CA:1447:G:H8	2.02	0.41
54:CA:1525:G:OP1	41:CN:120:ARG:NH2	2.53	0.41
54:CA:256:U:H2'	54:CA:257:G:H8	1.82	0.41
54:CA:358:U:H2'	54:CA:359:U:C6	2.55	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:437:U:H2'	54:CA:438:G:H5'	2.02	0.41
54:CA:754:C:H3'	54:CA:754:C:O2	2.19	0.41
52:CB:75:C:H6	52:CB:75:C:H3'	1.85	0.41
52:CC:72:C:H2'	52:CC:73:A:C5'	2.50	0.41
32:CE:217:ARG:HE	32:CE:217:ARG:HB2	1.53	0.41
34:CG:187:ARG:NH2	34:CG:193:ASP:OD2	2.39	0.41
34:CG:25:ARG:C	34:CG:27:TYR:N	2.68	0.41
34:CG:52:SER:O	34:CG:53:ASP:C	2.59	0.41
37:CJ:23:VAL:HG12	37:CJ:27:ILE:HD11	2.00	0.41
37:CJ:95:ARG:HA	37:CJ:98:SER:OG	2.20	0.41
41:CN:33:THR:HG22	41:CN:39:PRO:HA	2.03	0.41
41:CN:57:THR:HG22	41:CN:60:ALA:H	1.85	0.41
43:CP:100:GLY:O	43:CP:101:GLN:O	2.38	0.41
44:CQ:42:ILE:O	44:CQ:46:GLU:HG2	2.20	0.41
46:CS:58:TYR:C	46:CS:58:TYR:CD1	2.93	0.41
47:CT:62:SER:CB	47:CT:72:ARG:HE	2.33	0.41
50:CW:26:ASN:N	50:CW:26:ASN:HD22	2.19	0.41
16:D1:110:VAL:O	16:D1:114:LYS:N	2.53	0.41
16:D1:95:LEU:HD22	17:D2:4:ILE:HD12	2.02	0.41
22:D3:49:LYS:O	22:D3:50:ASN:HB2	2.20	0.41
28:D6:26:ASN:CG	28:D6:27:LYS:N	2.73	0.41
55:DA:1084:A:H1'	57:DY:53:VAL:HG21	2.02	0.41
55:DA:1157:G:O2'	25:DX:31:LEU:HD12	2.19	0.41
55:DA:1412:A:H2'	55:DA:1413:G:H8	1.85	0.41
55:DA:152:G:H2'	55:DA:153:C:H6	1.85	0.41
55:DA:2167:U:P	55:DA:2167:U:C6	3.14	0.41
55:DA:244:A:C2	55:DA:255:A:C4	3.08	0.41
55:DA:2746:U:H2'	55:DA:2747:G:H5'	2.03	0.41
55:DA:2790:A:O2'	55:DA:2893:G:O2'	1.82	0.41
55:DA:281:G:H1'	55:DA:360:G:N2	2.34	0.41
55:DA:2867:G:O2'	55:DA:2868:A:OP2	2.33	0.41
55:DA:498:G:H21	20:DU:47:LYS:HZ1	1.65	0.41
55:DA:600:G:H2'	55:DA:601:C:C6	2.56	0.41
55:DA:795:C:O2'	55:DA:796:C:H5'	2.20	0.41
2:DB:80:U:C2'	2:DB:81:G:H5'	2.50	0.41
3:DD:72:LYS:HG3	3:DD:103:ARG:NH2	2.36	0.41
3:DD:28:GLU:CB	3:DD:29:PRO:CD	2.96	0.41
4:DE:6:GLY:O	4:DE:195:LEU:HD12	2.20	0.41
56:DI:29:GLU:HG3	56:DJ:6:GLU:CD	2.31	0.41
55:DA:1059:G:P	58:DL:4:VAL:CG1	3.08	0.41
58:DL:81:ALA:O	58:DL:82:ALA:O	2.38	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:DO:112:LEU:HD22	11:DO:113:LYS:H	1.83	0.41
19:DT:49:VAL:HG13	19:DT:87:GLN:HE21	1.80	0.41
21:DV:174:VAL:HG12	21:DV:175:VAL:H	1.84	0.41
21:DV:178:GLU:O	21:DV:179:ASP:HB2	2.20	0.41
24:DW:23:LYS:HB3	24:DW:23:LYS:HE2	1.53	0.41
55:DA:1084:A:C8	57:DY:53:VAL:CG1	3.03	0.41
16:A1:52:ARG:O	16:A1:56:ASP:HB2	2.20	0.41
16:A1:95:LEU:O	16:A1:97:ASP:N	2.52	0.41
26:A4:15:ILE:O	26:A4:15:ILE:HG22	2.18	0.41
6:AG:109:VAL:HG11	26:A4:33:VAL:HG21	2.01	0.41
26:A4:38:LYS:HB3	26:A4:39:CYS:H	1.68	0.41
26:A4:6:HIS:N	26:A4:7:PRO:HD3	2.36	0.41
27:A5:45:VAL:HG12	27:A5:56:LYS:HG3	2.02	0.41
30:A8:34:TRP:C	30:A8:36:LYS:N	2.74	0.41
1:AA:1111:A:O2'	1:AA:1112:G:C4'	2.69	0.41
1:AA:1402:C:O2'	1:AA:1403:C:H5'	2.20	0.41
1:AA:1412:A:H2'	1:AA:1413:G:H8	1.85	0.41
1:AA:1474:C:H3'	1:AA:1475:G:H8	1.85	0.41
1:AA:1731:G:H3'	1:AA:1731:G:N3	2.34	0.41
1:AA:2134:A:C2	1:AA:2159:G:H1'	2.55	0.41
1:AA:222:A:HO2'	1:AA:223:A:P	2.41	0.41
1:AA:242:G:H8	30:A8:4:MET:O	2.03	0.41
1:AA:2496:C:C2'	1:AA:2497:A:O5'	2.68	0.41
1:AA:2827:C:H5'	1:AA:2828:C:OP2	2.21	0.41
1:AA:34:C:O2'	1:AA:35:G:O5'	2.39	0.41
1:AA:720:C:H2'	1:AA:721:C:H6	1.85	0.41
1:AA:808:G:H2'	1:AA:809:G:C8	2.56	0.41
1:AA:859:G:O2'	1:AA:860:U:OP2	2.37	0.41
1:AA:887:A:O2'	43:BP:93:ARG:CG	2.68	0.41
1:AA:92:G:C5	1:AA:93:C:C5	3.08	0.41
3:AD:155:LEU:N	3:AD:155:LEU:CD1	2.80	0.41
3:AD:24:ILE:HD11	3:AD:84:TYR:HB2	2.01	0.41
4:AE:134:ILE:HB	4:AE:137:HIS:HB2	2.02	0.41
5:AF:202:PHE:CE1	5:AF:206:ILE:HD11	2.56	0.41
5:AF:36:VAL:O	5:AF:40:GLN:HG2	2.20	0.41
6:AG:125:PHE:C	6:AG:127:GLY:N	2.73	0.41
6:AG:153:ARG:HH11	6:AG:153:ARG:HG2	1.86	0.41
6:AG:16:ARG:HB3	6:AG:17:PRO:HD3	2.01	0.41
8:AK:49:ALA:C	8:AK:51:ILE:H	2.23	0.41
8:AK:82:ARG:CG	8:AK:82:ARG:NH1	2.82	0.41
9:AM:43:THR:CG2	9:AM:44:PRO:HD2	2.50	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:AN:112:MET:O	10:AN:115:VAL:N	2.52	0.41
10:AN:98:VAL:HG13	10:AN:118:ALA:HA	2.02	0.41
11:AO:135:LEU:HD13	11:AO:135:LEU:O	2.20	0.41
11:AO:1:MET:HB2	11:AO:2:LYS:H	1.66	0.41
11:AO:48:PRO:CG	11:AO:49:ARG:N	2.83	0.41
19:AT:18:TYR:HD1	19:AT:21:PHE:CE2	2.38	0.41
20:AU:30:VAL:O	20:AU:31:LEU:HB2	2.21	0.41
21:AV:130:PRO:O	21:AV:133:ILE:HD13	2.20	0.41
21:AV:38:TYR:C	21:AV:38:TYR:CD1	2.93	0.41
21:AV:60:GLU:CG	21:AV:61:LEU:N	2.78	0.41
31:BA:1010:G:N2	31:BA:1020:U:H1'	2.36	0.41
31:BA:1123:A:H4'	40:BM:36:GLY:HA3	2.03	0.41
31:BA:1157:A:N3	31:BA:1157:A:H2'	2.35	0.41
31:BA:1176:A:C6	31:BA:1177:G:C6	3.08	0.41
31:BA:1405:G:O4'	31:BA:1519:A:H4'	2.20	0.41
31:BA:106:C:O2	31:BA:379:C:H4'	2.20	0.41
31:BA:440:A:H3'	31:BA:442:C:C6	2.49	0.41
31:BA:593:G:H2'	31:BA:594:G:O4'	2.19	0.41
31:BA:701:C:O2'	31:BA:702:A:P	2.78	0.41
31:BA:664:G:N2	31:BA:741:G:H1	2.10	0.41
31:BA:853:G:O2'	31:BA:854:G:H5'	2.20	0.41
31:BA:943:U:C2'	31:BA:944:G:H5'	2.50	0.41
52:BC:42:C:H2'	52:BC:42:C:O2	2.21	0.41
52:BC:62:C:H2'	52:BC:63:G:H8	1.85	0.41
52:BD:46:G:H21	52:BD:48:C:H1'	1.85	0.41
32:BE:108:ILE:O	32:BE:108:ILE:CG2	2.68	0.41
33:BF:188:LEU:CD1	33:BF:195:VAL:HG11	2.39	0.41
33:BF:45:LYS:HB2	33:BF:46:GLU:H	1.65	0.41
34:BG:139:ARG:CG	34:BG:139:ARG:NH1	2.83	0.41
34:BG:152:SER:O	34:BG:153:ARG:C	2.58	0.41
34:BG:17:VAL:HG11	34:BG:197:PRO:HB2	2.02	0.41
37:BJ:70:LYS:HA	37:BJ:71:PRO:HD2	1.90	0.41
37:BJ:80:VAL:HG13	37:BJ:80:VAL:O	2.19	0.41
41:BN:12:ARG:HG2	41:BN:13:GLN:N	2.34	0.41
43:BP:78:ILE:O	43:BP:80:ARG:N	2.53	0.41
46:BS:20:VAL:HG21	46:BS:32:TYR:HB3	2.02	0.41
47:BT:44:ALA:HA	47:BT:71:PHE:O	2.20	0.41
47:BT:68:ARG:O	47:BT:69:LYS:CB	2.67	0.41
49:BV:41:VAL:N	49:BV:44:MET:HE3	2.16	0.41
51:BX:6:ARG:O	51:BX:12:LYS:HG3	2.21	0.41
54:CA:1232:U:H2'	54:CA:1233:G:O4'	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1244:C:O2'	54:CA:1245:A:H5'	2.20	0.41
54:CA:1288:A:H2'	54:CA:1289:A:H8	1.85	0.41
54:CA:1429:C:H2'	54:CA:1430:C:C6	2.55	0.41
54:CA:333:G:H2'	54:CA:334:C:C6	2.55	0.41
54:CA:341:C:O2'	54:CA:342:C:H5'	2.20	0.41
54:CA:630:G:OP1	54:CA:630:G:O4'	2.38	0.41
52:CC:53:G:H2'	52:CC:54:U:C6	2.55	0.41
52:CD:29:G:C2'	52:CD:30:G:H5'	2.50	0.41
33:CF:69:HIS:CD2	33:CF:69:HIS:N	2.89	0.41
34:CG:154:ASN:O	34:CG:155:LEU:C	2.58	0.41
35:CH:76:ILE:HD11	35:CH:118:ILE:HD13	2.01	0.41
36:CI:101:ALA:HA	48:CU:28:GLU:HG2	2.01	0.41
38:CK:40:ALA:C	38:CK:42:GLU:H	2.24	0.41
42:CO:28:LYS:HZ2	42:CO:33:ARG:HH22	1.65	0.41
43:CP:15:VAL:CG1	43:CP:19:LEU:HD21	2.50	0.41
45:CR:70:LEU:HD11	45:CR:77:ARG:HG3	2.03	0.41
54:CA:189:U:C2	47:CT:72:ARG:NH1	2.88	0.41
50:CW:13:LEU:HD12	50:CW:14:LYS:N	2.34	0.41
16:D1:49:HIS:O	16:D1:52:ARG:N	2.54	0.41
27:D5:4:HIS:CB	27:D5:5:PRO:HD2	2.49	0.41
30:D8:22:VAL:HB	30:D8:53:PRO:CB	2.51	0.41
30:D8:26:LYS:NZ	30:D8:47:LYS:HD3	2.35	0.41
55:DA:1056:G:N2	55:DA:1087:G:C6	2.82	0.41
55:DA:1069:A:H4'	55:DA:1070:A:H5''	2.02	0.41
55:DA:1204:A:C2	55:DA:1241:A:C2	3.08	0.41
55:DA:1496:A:H5'	55:DA:1497:U:OP1	2.20	0.41
55:DA:1505:C:H2'	55:DA:1505:C:O2	2.19	0.41
55:DA:1608:A:HO2'	55:DA:1610:A:P	2.43	0.41
55:DA:793:A:N6	55:DA:2073:C:OP1	2.54	0.41
55:DA:2213:U:H4'	23:DZ:52:ARG:HH12	1.81	0.41
55:DA:828:U:N3	55:DA:2247:A:H4'	2.32	0.41
55:DA:2327:A:H2'	55:DA:2328:A:O4'	2.21	0.41
55:DA:270(Z):U:HO2'	55:DA:271(A):C:P	2.40	0.41
55:DA:278:A:H2'	55:DA:279:C:C6	2.55	0.41
55:DA:814:C:H2'	55:DA:815:C:H6	1.85	0.41
55:DA:930:U:O4'	55:DA:930:U:O2	2.37	0.41
55:DA:994:C:H3'	16:D1:54:LYS:HE3	2.01	0.41
3:DD:10:THR:O	3:DD:11:PRO:C	2.58	0.41
4:DE:176:ILE:CD1	4:DE:176:ILE:N	2.80	0.41
4:DE:31:CYS:HB3	4:DE:49:LEU:HD23	2.02	0.41
5:DF:203:GLN:HB2	5:DF:203:GLN:HE21	1.66	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:443:A:C3'	5:DF:45:ARG:NH1	2.80	0.41
7:DH:26:VAL:CG1	7:DH:27:LYS:N	2.84	0.41
56:DI:7:ARG:HE	56:DI:8:ILE:HD13	1.81	0.41
8:DK:109:ILE:HG13	8:DK:110:ASP:N	2.36	0.41
8:DK:9:LEU:C	8:DK:10:GLU:O	2.58	0.41
8:DK:94:ALA:C	8:DK:96:ASP:N	2.73	0.41
55:DA:1082:U:O2'	58:DL:117:THR:HG23	2.21	0.41
58:DL:12:LEU:CB	58:DL:13:PRO:CA	2.84	0.41
10:DN:2:ILE:N	10:DN:2:ILE:HD13	2.36	0.41
10:DN:98:VAL:HG13	10:DN:117:LEU:CB	2.49	0.41
11:DO:36:LYS:HB3	11:DO:37:GLY:H	1.64	0.41
11:DO:3:LEU:HA	11:DO:3:LEU:HD23	1.89	0.41
11:DO:42:SER:C	11:DO:44:GLY:N	2.72	0.41
11:DO:91:PHE:HZ	11:DO:100:LEU:CD1	2.33	0.41
12:DP:23:GLY:HA2	12:DP:101:ARG:NH1	2.35	0.41
12:DP:3:MET:HA	12:DP:4:PRO:HD3	1.89	0.41
12:DP:68:ILE:HD13	12:DP:103:MET:CE	2.48	0.41
12:DP:5:ARG:O	12:DP:6:ARG:O	2.37	0.41
12:DP:80:GLU:HA	22:D3:4:LYS:HZ2	1.80	0.41
14:DQ:8:GLU:H	14:DQ:8:GLU:HG3	1.67	0.41
18:DS:12:ILE:HG23	18:DS:17:VAL:HG21	2.02	0.41
20:DU:12:THR:HG23	20:DU:26:LYS:CE	2.50	0.41
21:DV:130:PRO:O	21:DV:133:ILE:HG13	2.19	0.41
57:DY:135:ARG:NH1	57:DY:138:LEU:CG	2.74	0.41
57:DY:138:LEU:O	57:DY:139:VAL:CB	2.68	0.41
57:DY:38:HIS:O	57:DY:39:ALA:HB3	2.20	0.41
57:DY:59:ILE:O	57:DY:60:ARG:C	2.57	0.41
57:DY:75:GLN:HG3	57:DY:110:GLY:N	2.34	0.41
23:DZ:51:VAL:HG11	23:DZ:74:VAL:HG21	2.01	0.41
23:DZ:83:GLU:O	23:DZ:85:LEU:N	2.53	0.41
23:DZ:91:LYS:CG	23:DZ:92:LYS:H	2.21	0.41
13:A0:34:ILE:HA	13:A0:34:ILE:HD13	1.78	0.41
22:A3:37:LEU:HG	22:A3:60:PHE:HA	2.02	0.41
22:A3:83:PRO:O	22:A3:84:LEU:C	2.59	0.41
26:A4:15:ILE:N	26:A4:15:ILE:CD1	2.83	0.41
28:A6:44:ARG:HH11	28:A6:44:ARG:HG2	1.84	0.41
1:AA:1252:G:C2	1:AA:1253:A:C2	3.08	0.41
1:AA:1397:U:O2'	1:AA:1398:C:OP1	2.39	0.41
1:AA:1493:C:C4'	1:AA:1494:A:OP2	2.68	0.41
1:AA:1924:C:C4	1:AA:1925:C:C6	2.95	0.41
1:AA:2018:G:H2'	1:AA:2019:A:O4'	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2104:G:O2'	1:AA:2105:C:H5'	2.20	0.41
1:AA:224:G:H2'	1:AA:225:A:O4'	2.21	0.41
1:AA:2296:U:O2	1:AA:2333:A:N3	2.53	0.41
1:AA:2564:A:OP1	1:AA:2648:C:H4'	2.20	0.41
1:AA:2712(A):A:H5''	1:AA:2713:A:OP2	2.20	0.41
1:AA:277:C:O2	1:AA:277:C:H2'	2.19	0.41
1:AA:2889:C:O2	1:AA:2889:C:H2'	2.21	0.41
1:AA:404:C:HO2'	1:AA:405:U:P	2.44	0.41
1:AA:554:U:O2'	1:AA:556:G:H8	2.01	0.41
1:AA:588:U:N3	1:AA:589:C:C4	2.89	0.41
1:AA:642:G:N1	1:AA:645:C:OP2	2.54	0.41
1:AA:709:U:H2'	1:AA:710:G:H8	1.84	0.41
1:AA:736:C:O2'	1:AA:737:C:H5'	2.20	0.41
1:AA:848:G:H8	1:AA:848:G:H5'	1.85	0.41
1:AA:917:A:O2'	1:AA:918:A:H5'	2.20	0.41
1:AA:993:G:H1'	17:A2:89:GLN:OE1	2.19	0.41
2:AB:0:A:N6	2:AB:119:A:N6	2.68	0.41
2:AB:95:U:C6	2:AB:95:U:C3'	3.04	0.41
3:AD:131:LEU:HB2	3:AD:136:ILE:HD11	2.02	0.41
3:AD:48:ARG:HH11	3:AD:48:ARG:CG	2.34	0.41
6:AG:112:PRO:HB2	26:A4:37:SER:CA	2.48	0.41
6:AG:131:TYR:HB3	6:AG:159:VAL:HG11	2.02	0.41
10:AN:120:GLU:OE2	10:AN:122:LEU:HD21	2.20	0.41
10:AN:97:ARG:NH1	31:BA:339:C:OP2	2.54	0.41
11:AO:131:SER:O	11:AO:132:LYS:C	2.59	0.41
1:AA:813:U:OP2	11:AO:23:PRO:O	2.38	0.41
11:AO:91:PHE:N	11:AO:91:PHE:HD1	2.18	0.41
14:AQ:108:GLY:O	14:AQ:109:GLY:C	2.59	0.41
18:AS:20:VAL:HG23	18:AS:21:VAL:N	2.35	0.41
18:AS:57:ASN:O	18:AS:58:ALA:C	2.59	0.41
21:AV:23:LYS:HB3	21:AV:38:TYR:CD1	2.56	0.41
24:AW:53:LEU:HD22	24:AW:57:ILE:CD1	2.47	0.41
24:AW:63:VAL:O	24:AW:67:LYS:HD3	2.20	0.41
24:AW:9:GLN:O	24:AW:13:ALA:N	2.50	0.41
23:AZ:80:LEU:C	23:AZ:81:LYS:NZ	2.70	0.41
31:BA:1120:G:H2'	31:BA:1121:U:C6	2.54	0.41
31:BA:1214:C:C5'	31:BA:1215:G:OP2	2.60	0.41
31:BA:1352:C:H2'	31:BA:1353:G:H8	1.78	0.41
31:BA:236:G:H2'	31:BA:237:C:H6	1.86	0.41
31:BA:31:G:HO2'	31:BA:32:A:P	2.44	0.41
31:BA:64:G:H4'	31:BA:66:G:OP1	2.19	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:BA:801:U:H2'	31:BA:802:A:O5'	2.19	0.41
31:BA:765:G:H21	31:BA:813:U:H5	1.68	0.41
31:BA:823:G:H2'	31:BA:824:C:C6	2.56	0.41
31:BA:872:A:C4	31:BA:874:G:C8	3.08	0.41
31:BA:890:G:C2'	31:BA:891:U:OP2	2.68	0.41
31:BA:958:A:C6	31:BA:959:A:C6	3.09	0.41
52:BD:34:G:H2'	52:BD:35:A:O4'	2.21	0.41
32:BE:180:LEU:O	32:BE:181:PHE:HB2	2.21	0.41
32:BE:17:PHE:CZ	32:BE:44:LEU:HB3	2.56	0.41
33:BF:116:VAL:HG11	33:BF:141:VAL:CG2	2.49	0.41
33:BF:157:ILE:C	33:BF:159:GLY:N	2.74	0.41
36:BI:9:VAL:C	36:BI:10:LEU:HD12	2.40	0.41
37:BJ:41:ARG:O	37:BJ:45:ASP:HB2	2.20	0.41
31:BA:1148:U:H4'	39:BL:14:VAL:CG1	2.51	0.41
39:BL:56:LEU:O	39:BL:56:LEU:HD23	2.21	0.41
39:BL:15:ALA:HB2	39:BL:65:VAL:HG23	2.03	0.41
40:BM:37:PRO:HA	40:BM:72:VAL:HG22	2.03	0.41
44:BQ:15:LYS:O	44:BQ:16:PHE:O	2.39	0.41
48:BU:53:ARG:C	48:BU:55:ARG:H	2.22	0.41
49:BV:24:ALA:O	49:BV:25:LYS:HB2	2.20	0.41
31:BA:958:A:C2	49:BV:55:LYS:HB2	2.55	0.41
50:BW:67:ALA:O	50:BW:69:GLY:N	2.53	0.41
54:CA:1121:U:H2'	54:CA:1122:U:C6	2.56	0.41
54:CA:1213:A:C5	54:CA:1215:G:C4	3.09	0.41
54:CA:959:A:C2	54:CA:1221:G:N3	2.82	0.41
54:CA:1262:C:C4	54:CA:1263:C:N4	2.88	0.41
54:CA:1346:A:C5	37:CJ:10:ARG:CZ	3.03	0.41
54:CA:922:G:N3	54:CA:1398:A:H2	2.18	0.41
54:CA:230:G:H2'	54:CA:231:G:O4'	2.20	0.41
54:CA:48:C:C4'	54:CA:49:U:OP2	2.66	0.41
54:CA:649:G:C4	54:CA:650:G:C8	3.09	0.41
52:CB:31:A:H2'	52:CB:32:U:H5'	2.01	0.41
32:CE:224:GLN:O	32:CE:226:ARG:N	2.54	0.41
33:CF:76:VAL:HG21	33:CF:103:VAL:HG13	2.02	0.41
34:CG:85:LYS:HG2	34:CG:86:LYS:O	2.19	0.41
36:CI:37:VAL:CG1	36:CI:38:GLU:N	2.82	0.41
47:CT:67:LYS:CA	47:CT:70:ARG:NH1	2.73	0.41
49:CV:41:VAL:CG1	49:CV:45:VAL:N	2.83	0.41
50:CW:59:ALA:O	50:CW:63:ILE:HG13	2.20	0.41
50:CW:44:ALA:HB1	50:CW:91:LEU:HB2	2.03	0.41
13:D0:37:THR:HG1	13:D0:40:LYS:HG3	1.85	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:D0:96:ARG:HD3	13:D0:98:LEU:HD21	2.02	0.41
17:D2:98:GLU:HB3	17:D2:100:ARG:HG3	2.02	0.41
17:D2:58:VAL:HG21	17:D2:100:ARG:HH12	1.84	0.41
29:D7:15:THR:HG22	29:D7:16:HIS:CE1	2.56	0.41
30:D8:14:VAL:CG1	30:D8:15:LYS:N	2.83	0.41
55:DA:1060:U:O2'	55:DA:1061:U:P	2.78	0.41
55:DA:1061:U:O4'	55:DA:1070:A:N3	2.53	0.41
55:DA:1638:C:H2'	55:DA:1639:U:O4'	2.20	0.41
55:DA:1642:G:O2'	55:DA:1643:G:H5'	2.21	0.41
55:DA:181:A:C2	55:DA:182:A:C4	3.07	0.41
55:DA:1773:A:C5	55:DA:1829:A:H1'	2.55	0.41
55:DA:2791:C:C2	55:DA:2792:G:C8	3.08	0.41
55:DA:26:G:C6	55:DA:27:G:C2	3.08	0.41
55:DA:571:A:C8	55:DA:575:A:N6	2.89	0.41
55:DA:589:C:H2'	55:DA:590:A:C8	2.55	0.41
55:DA:675:A:O2'	55:DA:676:A:H5'	2.20	0.41
3:DD:130:ALA:HB2	3:DD:192:THR:HB	2.02	0.41
5:DF:24:LEU:O	5:DF:25:PRO:C	2.58	0.41
5:DF:68:LYS:HG2	5:DF:69:HIS:CD2	2.55	0.41
6:DG:106:LEU:HD12	6:DG:110:ALA:CB	2.50	0.41
43:CP:7:VAL:HG21	6:DG:115:ARG:NH1	2.36	0.41
6:DG:115:ARG:CG	6:DG:115:ARG:NH1	2.81	0.41
6:DG:80:PHE:C	6:DG:81:LYS:HG2	2.40	0.41
7:DH:103:LEU:HD23	7:DH:103:LEU:O	2.21	0.41
56:DJ:18:LEU:HA	56:DJ:21:LYS:CG	2.51	0.41
56:DJ:28:LYS:HG2	56:DJ:29:GLU:N	2.35	0.41
58:DL:44:ALA:O	58:DL:45:THR:C	2.59	0.41
55:DA:1952:A:C5	10:DN:22:ILE:HD12	2.56	0.41
11:DO:94:GLU:O	11:DO:95:VAL:HB	2.21	0.41
21:DV:177:PRO:O	21:DV:178:GLU:HG2	2.19	0.41
52:CB:57:G:C4'	21:DV:182:LYS:NZ	2.81	0.41
21:DV:28:MET:CE	21:DV:67:LEU:HD13	2.49	0.41
24:DW:10:LEU:HD23	24:DW:10:LEU:HA	1.77	0.41
25:DX:8:LEU:HD23	25:DX:53:LEU:O	2.21	0.41
57:DY:3:ASN:O	57:DY:5:ARG:N	2.54	0.41
57:DY:7:VAL:HG13	57:DY:8:GLU:CB	2.50	0.41
57:DY:50:ARG:CA	57:DY:83:TYR:HA	2.49	0.41
23:DZ:58:ILE:CG2	23:DZ:87:PRO:HG3	2.51	0.41
22:A3:70:GLN:NE2	22:A3:80:HIS:HE2	2.19	0.41
26:A4:4:GLY:O	26:A4:5:ILE:HB	2.20	0.41
1:AA:1108:U:H2'	1:AA:1109:C:O4'	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1362:C:C2'	1:AA:1363:C:H5'	2.51	0.41
1:AA:1398:C:O2'	1:AA:1399:C:H5'	2.21	0.41
1:AA:140:A:H1'	1:AA:1408:C:O2'	2.21	0.41
1:AA:2130:U:H1'	1:AA:2134:A:O4'	2.21	0.41
1:AA:2283:C:C6	1:AA:2389:G:H2'	2.56	0.41
1:AA:2388:A:H5'	1:AA:2389:G:OP2	2.20	0.41
1:AA:2425:A:H4'	1:AA:2426:A:O5'	2.20	0.41
1:AA:2641:G:P	9:AM:74:ARG:HH21	2.44	0.41
1:AA:2742:C:C4	1:AA:2763:G:N2	2.88	0.41
1:AA:2759:G:C6	1:AA:2760:C:C4	3.09	0.41
1:AA:297:C:O2'	1:AA:298:G:H5'	2.20	0.41
1:AA:372:G:O2'	1:AA:373:U:C5	2.73	0.41
1:AA:587:C:O2'	1:AA:588:U:P	2.79	0.41
1:AA:588:U:H1'	5:AF:90:PHE:HB3	2.03	0.41
1:AA:718:A:H8	1:AA:718:A:O5'	2.03	0.41
3:AD:11:PRO:C	3:AD:13:ARG:H	2.23	0.41
3:AD:61:LEU:HA	3:AD:61:LEU:HD12	1.80	0.41
6:AG:145:THR:OG1	6:AG:148:MET:HB2	2.20	0.41
6:AG:9:ARG:C	6:AG:11:TYR:N	2.74	0.41
7:AH:11:VAL:HG23	7:AH:11:VAL:O	2.19	0.41
8:AK:102:SER:HA	8:AK:106:GLY:CA	2.51	0.41
8:AK:79:ILE:C	8:AK:142:VAL:HG21	2.40	0.41
10:AN:64:ARG:O	10:AN:82:ASN:HA	2.21	0.41
14:AQ:26:LEU:O	14:AQ:26:LEU:CD2	2.65	0.41
18:AS:29:LEU:CD1	18:AS:51:LEU:HD11	2.51	0.41
21:AV:14:LYS:HA	21:AV:15:PRO:HD3	1.89	0.41
21:AV:177:PRO:O	21:AV:178:GLU:HG3	2.21	0.41
21:AV:33:LEU:HD23	21:AV:90:VAL:HG21	2.01	0.41
24:AW:60:LEU:HD12	24:AW:60:LEU:N	2.35	0.41
1:AA:1365:A:H5'	23:AZ:12:PRO:HG2	2.03	0.41
23:AZ:58:ILE:HG22	23:AZ:60:PHE:CE1	2.55	0.41
23:AZ:97:LEU:O	23:AZ:98:LEU:OXT	2.39	0.41
31:BA:1156:G:C3'	31:BA:1157:A:H5''	2.51	0.41
31:BA:971:G:C6	31:BA:1365:G:H5'	2.55	0.41
31:BA:518:C:C5'	31:BA:519:C:O5'	2.68	0.41
31:BA:711:G:HO2'	31:BA:712:A:H5'	1.86	0.41
31:BA:795:C:C5	31:BA:796:C:C5	3.09	0.41
31:BA:84:U:H3'	31:BA:85:U:C4	2.56	0.41
31:BA:930:C:C2'	31:BA:931:C:H5'	2.51	0.41
52:BB:16:U:H2'	52:BB:17:C:H5'	2.03	0.41
32:BE:12:GLU:O	32:BE:13:ALA:C	2.58	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:BE:83:MET:O	32:BE:86:GLU:N	2.53	0.41
33:BF:127:ARG:NH1	33:BF:127:ARG:HG2	2.34	0.41
31:BA:1206:G:H4'	33:BF:192:THR:C	2.41	0.41
33:BF:32:LEU:O	33:BF:35:GLU:CB	2.68	0.41
33:BF:94:LEU:HD12	33:BF:94:LEU:C	2.40	0.41
34:BG:60:GLU:OE1	34:BG:202:LEU:HD12	2.21	0.41
34:BG:52:SER:H	34:BG:55:ALA:HB3	1.85	0.41
36:BI:53:ALA:O	36:BI:54:LYS:HB3	2.20	0.41
36:BI:69:GLU:CD	36:BI:69:GLU:H	2.23	0.41
39:BL:99:LEU:HD22	39:BL:99:LEU:N	2.36	0.41
44:BQ:47:LEU:HB2	44:BQ:53:LEU:CD1	2.49	0.41
45:BR:64:ARG:HG2	45:BR:64:ARG:HH11	1.84	0.41
45:BR:66:LEU:O	45:BR:67:LEU:C	2.58	0.41
47:BT:34:LYS:O	47:BT:35:VAL:C	2.58	0.41
49:BV:52:TYR:HA	49:BV:57:HIS:HA	2.02	0.41
49:BV:9:VAL:O	49:BV:10:PHE:CG	2.73	0.41
50:BW:48:LYS:O	50:BW:49:ALA:C	2.59	0.41
54:CA:1148:U:H2'	54:CA:1149:C:O4'	2.21	0.41
54:CA:1200:C:H4'	54:CA:1201:A:C5'	2.42	0.41
54:CA:1389:C:H2'	54:CA:1390:U:O4'	2.21	0.41
54:CA:372:C:O2'	54:CA:373:A:P	2.78	0.41
54:CA:390:C:O3'	46:CS:28:ARG:NH2	2.52	0.41
54:CA:449:C:C5	54:CA:450:G:C4	3.08	0.41
54:CA:458:C:H2'	54:CA:464:G:H8	1.85	0.41
54:CA:506:G:C6	54:CA:507:C:C4	3.09	0.41
54:CA:545:C:O2'	54:CA:549:C:OP1	2.38	0.41
54:CA:60:A:HO2'	54:CA:61:G:P	2.43	0.41
54:CA:626:U:H2'	54:CA:627:G:C8	2.56	0.41
54:CA:642:A:N3	38:CK:113:SER:OG	2.39	0.41
54:CA:673:G:O3'	36:CI:87:ARG:NH2	2.53	0.41
54:CA:706:A:O2'	41:CN:31:THR:CG2	2.69	0.41
54:CA:778:G:O5'	54:CA:778:G:H8	2.03	0.41
54:CA:894:G:C6	54:CA:895:G:C5	3.08	0.41
54:CA:73:G:N7	54:CA:99:C:N3	2.68	0.41
52:CD:45:U:C5'	52:CD:46:G:OP1	2.63	0.41
32:CE:95:GLN:HE21	32:CE:147:LYS:CE	2.33	0.41
33:CF:84:ILE:O	33:CF:84:ILE:HG12	2.21	0.41
35:CH:118:ILE:HG13	35:CH:119:LEU:N	2.35	0.41
35:CH:76:ILE:CB	35:CH:77:PRO:HD2	2.49	0.41
42:CO:120:TYR:O	42:CO:121:GLY:C	2.59	0.41
43:CP:108:ARG:CD	43:CP:108:ARG:H	2.25	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
43:CP:123:ALA:HA	43:CP:124:PRO:HD3	1.61	0.41
43:CP:27:LYS:HB3	43:CP:31:LYS:HE3	2.03	0.41
43:CP:27:LYS:O	43:CP:31:LYS:HG3	2.20	0.41
45:CR:9:GLN:HA	45:CR:12:ILE:HD12	2.03	0.41
48:CU:26:LEU:HD13	48:CU:42:ARG:NH1	2.35	0.41
49:CV:14:HIS:CD2	49:CV:35:SER:HB2	2.51	0.41
50:CW:38:LYS:O	50:CW:41:ILE:HG13	2.21	0.41
51:CX:2:GLY:O	51:CX:4:GLY:N	2.54	0.41
13:D0:31:HIS:C	13:D0:33:ARG:H	2.24	0.41
16:D1:105:VAL:HG11	17:D2:40:LEU:HD13	2.02	0.41
26:D4:8:LYS:O	26:D4:9:LEU:HB3	2.20	0.41
29:D7:8:ASN:ND2	29:D7:8:ASN:O	2.47	0.41
55:DA:1126:A:H8	55:DA:1126:A:OP1	2.04	0.41
55:DA:1374:G:C6	55:DA:1375:C:C4	3.08	0.41
55:DA:146:G:H2'	55:DA:147:U:O4'	2.21	0.41
55:DA:1586:A:O4'	55:DA:1586:A:N3	2.53	0.41
55:DA:164:U:C5	55:DA:165:U:H5	2.39	0.41
55:DA:1813:G:O2'	3:DD:44:ASN:HA	2.21	0.41
55:DA:2321:G:N3	55:DA:2321:G:C2'	2.84	0.41
55:DA:238:C:O2'	55:DA:608:A:H1'	2.21	0.41
55:DA:2817:G:C5	55:DA:2830:G:C2	3.08	0.41
55:DA:57:C:H2'	55:DA:58:G:O4'	2.21	0.41
55:DA:653:A:H5''	55:DA:654:A:OP2	2.19	0.41
55:DA:669:G:H2'	55:DA:669:G:N3	2.35	0.41
55:DA:70:G:OP2	55:DA:70:G:H8	2.03	0.41
55:DA:857:C:C4	55:DA:858:U:C4	3.09	0.41
55:DA:860:U:O4	55:DA:917:A:H2	2.04	0.41
2:DB:42:C:O2	6:DG:93:THR:N	2.40	0.41
3:DD:122:ASP:OD2	3:DD:122:ASP:N	2.53	0.41
4:DE:34:VAL:HG23	4:DE:48:GLN:HB3	2.03	0.41
6:DG:112:PRO:HB3	26:D4:37:SER:CA	2.51	0.41
6:DG:31:VAL:HG13	6:DG:31:VAL:O	2.21	0.41
55:DA:2311:A:H8	6:DG:88:ILE:HG13	1.85	0.41
56:DJ:23:LEU:HD22	56:DJ:23:LEU:HA	1.99	0.41
58:DL:104:VAL:HG12	58:DL:105:LEU:CA	2.49	0.41
58:DL:19:PRO:HG2	58:DL:38:VAL:HG21	2.03	0.41
55:DA:1138:G:O2'	9:DM:105:GLY:HA3	2.20	0.41
9:DM:39:ARG:HH11	9:DM:39:ARG:HB3	1.83	0.41
9:DM:71:ILE:N	9:DM:71:ILE:CD1	2.70	0.41
11:DO:37:GLY:O	11:DO:38:GLN:CB	2.69	0.41
11:DO:85:LEU:O	11:DO:88:LEU:HD23	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:DP:76:LYS:HG3	12:DP:77:LYS:O	2.21	0.41
15:DR:57:PHE:CG	15:DR:58:ASN:N	2.88	0.41
15:DR:85:LYS:HE2	15:DR:87:ASP:CG	2.41	0.41
20:DU:78:ALA:HB3	20:DU:81:LYS:CE	2.49	0.41
21:DV:31:ARG:HB2	21:DV:31:ARG:HE	1.54	0.41
21:DV:72:ARG:HH11	21:DV:72:ARG:CG	2.31	0.41
24:DW:17:SER:CB	24:DW:18:PRO:CA	2.99	0.41
24:DW:69:ARG:O	24:DW:70:GLN:O	2.39	0.41
24:DW:8:LYS:CB	24:DW:8:LYS:NZ	2.84	0.41
57:DY:19:ARG:NH1	57:DY:84:GLU:CD	2.61	0.41
57:DY:48:GLY:C	57:DY:84:GLU:CB	2.83	0.41
57:DY:62:ALA:C	57:DY:63:LEU:CD2	2.77	0.41
13:A0:79:LEU:CD2	13:A0:79:LEU:C	2.89	0.41
13:A0:77:ARG:O	13:A0:80:PHE:N	2.53	0.41
16:A1:100:VAL:C	16:A1:102:GLU:N	2.74	0.41
17:A2:98:GLU:O	17:A2:99:ILE:HB	2.21	0.41
22:A3:72:ARG:CZ	22:A3:75:LEU:HD12	2.51	0.41
26:A4:40:HIS:N	26:A4:41:PRO:CD	2.83	0.41
28:A6:25:LYS:HB3	30:A8:34:TRP:CZ2	2.56	0.41
1:AA:1103:A:H2'	1:AA:1104:C:C5'	2.49	0.41
1:AA:1225:C:O2'	17:A2:85:LYS:N	2.36	0.41
1:AA:1312:U:HO2'	1:AA:1314:C:H41	1.65	0.41
1:AA:1851:U:O2'	52:BD:71:G:H1'	2.20	0.41
1:AA:1965:C:H2'	1:AA:1966:A:H8	1.86	0.41
1:AA:218:A:O5'	1:AA:218:A:H8	2.04	0.41
1:AA:2729:G:O2'	4:AE:170:LEU:HD11	2.20	0.41
1:AA:2848:G:H2'	1:AA:2867:G:H22	1.85	0.41
1:AA:456:C:H6	1:AA:456:C:H3'	1.85	0.41
1:AA:536:A:H2'	1:AA:537:C:O4'	2.21	0.41
1:AA:571:A:C6	1:AA:575:A:C8	3.08	0.41
1:AA:71:A:O2'	1:AA:72:U:OP2	2.39	0.41
1:AA:734:A:H2'	1:AA:735:A:O4'	2.20	0.41
1:AA:896:A:H2	21:AV:178:GLU:CD	2.24	0.41
2:AB:7:G:C2'	2:AB:8:U:H5''	2.50	0.41
3:AD:111:LEU:HD22	3:AD:115:GLN:NE2	2.36	0.41
3:AD:156:ALA:O	3:AD:157:ARG:HB3	2.20	0.41
3:AD:43:ARG:HB3	3:AD:54:ARG:HB2	2.03	0.41
1:AA:2784:C:C4'	4:AE:41:LYS:O	2.64	0.41
4:AE:48:GLN:O	4:AE:78:LEU:HB2	2.21	0.41
4:AE:48:GLN:CD	4:AE:78:LEU:HD12	2.40	0.41
5:AF:164:ARG:HD3	5:AF:175:THR:OG1	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:AF:192:LEU:HD21	5:AF:194:MET:HG2	2.02	0.41
5:AF:20:LEU:HD13	5:AF:199:TRP:HH2	1.85	0.41
5:AF:84:VAL:O	5:AF:85:GLY:C	2.59	0.41
6:AG:7:LEU:HD22	6:AG:100:TRP:HZ3	1.85	0.41
6:AG:114:ILE:CD1	6:AG:140:ILE:HG21	2.50	0.41
7:AH:10:PRO:HB2	7:AH:50:VAL:CG1	2.51	0.41
8:AK:10:GLU:OE2	8:AK:11:ASN:N	2.54	0.41
10:AN:101:PRO:O	10:AN:102:VAL:HG13	2.21	0.41
11:AO:38:GLN:HG2	11:AO:45:LEU:CD1	2.49	0.41
14:AQ:106:ARG:HA	14:AQ:110:LEU:HD11	2.01	0.41
14:AQ:71:ARG:O	14:AQ:75:GLU:HG3	2.19	0.41
15:AR:54:ARG:HG2	15:AR:54:ARG:HH11	1.85	0.41
15:AR:64:ARG:HA	15:AR:72:VAL:O	2.20	0.41
15:AR:80:SER:HA	15:AR:81:PRO:HD3	1.84	0.41
19:AT:64:LYS:HD3	19:AT:73:ARG:CD	2.50	0.41
21:AV:116:VAL:CG1	21:AV:117:LEU:N	2.72	0.41
21:AV:99:TYR:CE1	21:AV:125:LEU:HB2	2.56	0.41
21:AV:60:GLU:HG3	21:AV:61:LEU:O	2.21	0.41
24:AW:30:ARG:O	24:AW:31:GLU:C	2.58	0.41
24:AW:68:ARG:CG	24:AW:68:ARG:NH1	2.84	0.41
53:B1:46:U:O2'	53:B1:47:U:H5'	2.20	0.41
31:BA:1256:A:H61	31:BA:1278:U:P	2.42	0.41
31:BA:1320:C:N4	31:BA:1321:C:H41	2.19	0.41
31:BA:1359:C:H6	31:BA:1359:C:O5'	2.04	0.41
31:BA:216:G:O2'	31:BA:217:C:C6	2.50	0.41
31:BA:356:A:H2'	31:BA:357:G:O4'	2.21	0.41
31:BA:422:C:HO2'	31:BA:423:G:N2	2.17	0.41
31:BA:556:C:C2'	31:BA:557:G:H5'	2.51	0.41
31:BA:575:G:O2'	31:BA:576:G:O5'	2.35	0.41
31:BA:659:U:H2'	31:BA:660:G:C8	2.56	0.41
31:BA:679:C:H2'	31:BA:680:C:C6	2.55	0.41
31:BA:687:A:H1'	31:BA:688:G:O4'	2.21	0.41
32:BE:15:VAL:O	32:BE:15:VAL:HG12	2.19	0.41
32:BE:193:ASP:OD2	32:BE:193:ASP:O	2.37	0.41
37:BJ:35:LYS:NZ	37:BJ:38:LEU:HD22	2.36	0.41
40:BM:31:GLY:O	40:BM:78:ASN:ND2	2.53	0.41
41:BN:34:ASP:OD2	41:BN:34:ASP:C	2.59	0.41
41:BN:66:LEU:O	41:BN:67:ASP:C	2.59	0.41
45:BR:2:PRO:HG2	45:BR:3:ILE:CD1	2.50	0.41
47:BT:10:VAL:HG21	47:BT:55:ASP:HB2	2.03	0.41
54:CA:1100:C:O2'	54:CA:1101:A:H5'	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:1228:C:OP2	43:CP:108:ARG:NH2	2.54	0.41
54:CA:16:A:C2'	54:CA:17:U:H5'	2.51	0.41
54:CA:439:A:C4	54:CA:496:A:C2	3.08	0.41
54:CA:533:A:O2'	54:CA:534:U:O5'	2.39	0.41
54:CA:562:C:O2'	42:CO:15:ARG:CB	2.48	0.41
54:CA:743:U:H2'	54:CA:744:C:C6	2.55	0.41
54:CA:812:C:O2'	54:CA:813:U:H6	2.03	0.41
52:CB:28:G:H2'	52:CB:29:G:H8	1.86	0.41
52:CD:49:C:O5'	52:CD:49:C:H6	2.04	0.41
32:CE:204:ASN:ND2	32:CE:204:ASN:C	2.74	0.41
33:CF:70:VAL:HG12	33:CF:72:LYS:N	2.05	0.41
33:CF:83:ARG:C	33:CF:85:ARG:H	2.24	0.41
35:CH:48:ALA:HB1	35:CH:49:PRO:HD2	2.03	0.41
35:CH:92:LYS:HA	35:CH:93:PRO:HD2	1.92	0.41
41:CN:90:GLY:O	41:CN:91:ARG:C	2.59	0.41
43:CP:108:ARG:CG	43:CP:108:ARG:HH11	2.33	0.41
43:CP:15:VAL:HG12	43:CP:19:LEU:CD2	2.50	0.41
43:CP:40:ASN:ND2	43:CP:42:ALA:HB3	2.33	0.41
47:CT:27:PHE:HB2	47:CT:28:PRO:CD	2.50	0.41
13:D0:56:LYS:C	13:D0:58:GLY:N	2.72	0.41
13:D0:79:LEU:HA	13:D0:83:ILE:CG1	2.43	0.41
16:D1:108:GLU:OE1	16:D1:112:ARG:HG2	2.21	0.41
17:D2:6:LYS:HA	17:D2:11:GLN:HA	2.02	0.41
27:D5:13:LYS:HG2	27:D5:16:ARG:NH2	2.35	0.41
27:D5:58:LEU:N	27:D5:58:LEU:HD12	2.36	0.41
28:D6:11:LEU:HG	28:D6:51:GLU:HG3	2.02	0.41
55:DA:1059:G:N2	58:DL:126:MET:C	2.72	0.41
55:DA:1099:G:H3'	55:DA:1099:G:C8	2.55	0.41
55:DA:119:A:C4'	55:DA:120:U:OP1	2.68	0.41
55:DA:1348:G:C2'	55:DA:1349:A:C5'	2.93	0.41
55:DA:1510:A:O2'	55:DA:1511:A:C8	2.74	0.41
55:DA:1893:C:H2'	55:DA:1894:C:C5'	2.50	0.41
55:DA:2152:G:H2'	55:DA:2153:G:H8	1.86	0.41
55:DA:2115:G:H1'	55:DA:2171:A:N6	2.34	0.41
55:DA:2275:C:O2	12:DP:83:MET:HG2	2.21	0.41
55:DA:2286:A:H8	55:DA:2287:A:C6	2.38	0.41
55:DA:2377:A:H2'	55:DA:2378:A:C8	2.56	0.41
55:DA:241:A:O4'	55:DA:243:U:C6	2.73	0.41
55:DA:2473:U:H2'	55:DA:2473:U:O2	2.20	0.41
55:DA:2485:G:O2'	55:DA:2486:G:H5'	2.21	0.41
55:DA:2505:G:H2'	55:DA:2576:G:O6	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2588:G:O2'	55:DA:2589:A:H5'	2.19	0.41
55:DA:319:C:H2'	55:DA:320:A:O4'	2.20	0.41
55:DA:357:A:H2'	55:DA:358:U:C6	2.55	0.41
55:DA:422:A:C6	55:DA:423:A:C6	3.08	0.41
55:DA:709:U:H2'	55:DA:710:G:C8	2.56	0.41
55:DA:774:A:H2'	55:DA:775:G:OP2	2.21	0.41
55:DA:809:G:O2'	55:DA:810:U:H5'	2.21	0.41
3:DD:176:ARG:CG	3:DD:176:ARG:NH1	2.82	0.41
4:DE:27:LEU:HD21	15:DR:1:MET:HE1	2.03	0.41
4:DE:92:THR:HG22	4:DE:93:VAL:H	1.86	0.41
6:DG:7:LEU:HD22	6:DG:100:TRP:CE3	2.56	0.41
6:DG:111:LEU:HD13	6:DG:120:LEU:HD21	2.03	0.41
6:DG:16:ARG:NH1	6:DG:16:ARG:CG	2.75	0.41
7:DH:32:GLU:O	7:DH:33:LEU:HD23	2.20	0.41
56:DJ:18:LEU:HA	56:DJ:18:LEU:HD23	1.70	0.41
58:DL:144:VAL:HG22	58:DL:145:LYS:N	2.35	0.41
55:DA:1060:U:C5	58:DL:74:ALA:HB2	2.45	0.41
10:DN:77:ILE:CG2	10:DN:77:ILE:O	2.68	0.41
12:DP:10:ARG:HG2	12:DP:10:ARG:H	1.64	0.41
18:DS:34:ASN:O	18:DS:35:ILE:C	2.58	0.41
21:DV:33:LEU:O	21:DV:34:ASN:HB2	2.20	0.41
24:DW:43:GLN:O	24:DW:44:LEU:CG	2.69	0.41
57:DY:26:LEU:CD2	57:DY:112:LEU:HB3	2.51	0.41
57:DY:16:ASN:ND2	57:DY:25:PHE:CE2	2.86	0.41
23:DZ:63:ALA:C	23:DZ:65:SER:N	2.71	0.41
23:DZ:87:PRO:O	23:DZ:89:GLU:N	2.54	0.41
26:A4:12:ALA:HB3	26:A4:29:PRO:O	2.20	0.41
28:A6:24:GLU:O	30:A8:34:TRP:CZ3	2.74	0.41
30:A8:40:GLU:O	30:A8:41:ILE:HB	2.20	0.41
30:A8:6:THR:OG1	30:A8:8:LYS:HE3	2.21	0.41
1:AA:1066:U:H6	1:AA:1066:U:O5'	2.03	0.41
1:AA:1082:U:H6	1:AA:1082:U:O5'	2.04	0.41
1:AA:1725:G:H2'	1:AA:1725:G:N3	2.35	0.41
1:AA:1819:A:O2'	1:AA:1820:U:OP2	2.26	0.41
1:AA:1857:G:N2	1:AA:1886:C:N4	2.68	0.41
1:AA:2258:C:C4'	1:AA:2259:G:OP2	2.65	0.41
1:AA:2420:C:OP1	30:A8:34:TRP:HB3	2.21	0.41
1:AA:2576:G:H3'	1:AA:2576:G:N3	2.36	0.41
1:AA:2691:C:H5'	1:AA:2691:C:H6	1.86	0.41
1:AA:2712:U:O2'	1:AA:2712(A):A:P	2.78	0.41
1:AA:2715:C:H2'	1:AA:2716:U:H6	1.85	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2762:G:H5'	1:AA:2763:G:OP2	2.20	0.41
1:AA:2798:C:H5	1:AA:2799:A:N6	2.19	0.41
1:AA:2839:G:H4'	13:A0:49:ASP:CB	2.50	0.41
1:AA:2888:C:C2	1:AA:2889:C:C6	3.08	0.41
1:AA:714:U:H2'	1:AA:716:A:OP2	2.21	0.41
1:AA:898:C:N4	1:AA:899:A:C6	2.88	0.41
1:AA:996:A:H2'	1:AA:997:G:H8	1.86	0.41
3:AD:172:TYR:HD1	3:AD:185:VAL:C	2.24	0.41
3:AD:79:VAL:O	3:AD:79:VAL:HG12	2.19	0.41
5:AF:18:ARG:O	5:AF:19:GLU:CB	2.67	0.41
5:AF:31:HIS:NE2	5:AF:35:GLU:OE1	2.51	0.41
7:AH:107:VAL:HG23	7:AH:109:PHE:CE1	2.56	0.41
8:AK:93:THR:H	8:AK:96:ASP:HB2	1.86	0.41
9:AM:19:GLU:C	9:AM:21:LYS:H	2.24	0.41
9:AM:6:PRO:C	9:AM:7:LYS:HZ3	2.23	0.41
11:AO:95:VAL:HG23	11:AO:125:VAL:HG23	2.02	0.41
15:AR:103:ARG:O	15:AR:105:LEU:N	2.54	0.41
15:AR:124:ASP:C	15:AR:126:ALA:N	2.73	0.41
15:AR:62:THR:HG22	15:AR:75:ILE:HG23	2.02	0.41
20:AU:47:LYS:HG3	20:AU:60:PHE:CB	2.45	0.41
21:AV:102:LEU:HD21	21:AV:124:ILE:HG21	2.01	0.41
21:AV:132:ASN:OD1	21:AV:132:ASN:O	2.38	0.41
21:AV:33:LEU:CG	21:AV:34:ASN:N	2.83	0.41
31:BA:1028:C:N4	31:BA:1028(A):C:H41	2.18	0.41
31:BA:1133:G:H2'	31:BA:1134:G:H8	1.85	0.41
31:BA:158:G:H2'	31:BA:159:G:H5'	2.02	0.41
31:BA:173:U:H1'	31:BA:197:A:C6	2.55	0.41
31:BA:484:G:H4'	31:BA:485:G:O5'	2.20	0.41
31:BA:706:A:O4'	41:BN:29:ILE:HD11	2.20	0.41
31:BA:792:A:N3	31:BA:794:A:N7	2.68	0.41
31:BA:797:C:O2'	31:BA:798:G:H5'	2.19	0.41
32:BE:151:GLY:C	32:BE:153:ARG:H	2.22	0.41
32:BE:56:ARG:CG	32:BE:56:ARG:HH11	2.34	0.41
32:BE:69:LEU:C	32:BE:69:LEU:HD23	2.41	0.41
33:BF:62:ASP:HA	33:BF:97:LYS:CD	2.50	0.41
33:BF:62:ASP:HA	33:BF:97:LYS:HD2	2.02	0.41
34:BG:165:MET:CE	34:BG:168:ARG:HB2	2.48	0.41
34:BG:173:TRP:HB2	34:BG:187:ARG:O	2.21	0.41
34:BG:173:TRP:CZ3	34:BG:193:ASP:HB3	2.56	0.41
35:BH:12:LEU:CD2	35:BH:13:ILE:N	2.84	0.41
39:BL:53:VAL:HG13	39:BL:95:LYS:HE3	2.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:BN:27:ASN:CG	41:BN:28:THR:N	2.73	0.41
45:BR:39:LEU:HD13	45:BR:39:LEU:O	2.20	0.41
46:BS:51:VAL:CG1	46:BS:52:ASP:N	2.82	0.41
31:BA:1313:U:H5	49:BV:4:SER:HB2	1.85	0.41
50:BW:67:ALA:CB	50:BW:73:HIS:HA	2.51	0.41
53:C1:33:G:C2'	53:C1:34:G:C8	2.96	0.41
54:CA:1297:C:C2'	54:CA:1298:C:OP2	2.68	0.41
54:CA:1347:G:OP2	39:CL:107:ARG:HG2	2.21	0.41
54:CA:134:A:H61	46:CS:25:ARG:NH1	2.18	0.41
54:CA:1422:G:H2'	54:CA:1423:G:H8	1.86	0.41
54:CA:1476:G:H2'	54:CA:1477:C:C6	2.56	0.41
54:CA:186:C:O3'	50:CW:82:SER:HA	2.21	0.41
54:CA:877:C:H5''	38:CK:88:LYS:CD	2.51	0.41
54:CA:943:U:H2'	54:CA:944:G:H5'	2.03	0.41
52:CB:5:G:O2'	52:CB:6:G:H5'	2.21	0.41
52:CC:17:C:H6	52:CC:17:C:P	2.44	0.41
32:CE:48:MET:O	32:CE:49:GLU:C	2.59	0.41
33:CF:119:ARG:NH1	33:CF:119:ARG:HG3	2.34	0.41
33:CF:83:ARG:O	33:CF:86:VAL:HG22	2.21	0.41
34:CG:67:ILE:HG22	34:CG:68:TYR:CD1	2.56	0.41
39:CL:3:GLN:HB3	39:CL:20:ARG:CD	2.44	0.41
41:CN:103:LEU:CD2	41:CN:103:LEU:H	2.33	0.41
43:CP:15:VAL:HG11	43:CP:34:LEU:HD21	2.03	0.41
45:CR:65:ARG:HH11	45:CR:65:ARG:HB2	1.84	0.41
45:CR:79:ARG:HA	45:CR:82:ILE:CG2	2.51	0.41
46:CS:9:PHE:HB3	46:CS:10:GLY:H	1.64	0.41
54:CA:392:G:C5'	46:CS:12:LYS:HG3	2.50	0.41
49:CV:3:ARG:O	49:CV:4:SER:HB3	2.21	0.41
49:CV:41:VAL:HG12	49:CV:45:VAL:H	1.86	0.41
55:DA:1248:G:C4	16:D1:3:ARG:HG3	2.55	0.41
17:D2:15:GLU:O	17:D2:18:LEU:HB2	2.21	0.41
17:D2:44:LYS:HE2	17:D2:44:LYS:HB3	1.81	0.41
49:CV:42:PRO:CD	26:D4:63:TYR:HE2	2.33	0.41
27:D5:60:VAL:CG1	27:D5:60:VAL:OXT	2.69	0.41
28:D6:42:TRP:O	28:D6:43:CYS:CB	2.68	0.41
55:DA:1510:A:OP1	55:DA:1511:A:C5'	2.60	0.41
55:DA:1590:U:H2'	55:DA:1591:G:H8	1.84	0.41
55:DA:1668:A:H61	55:DA:1676:A:N6	2.18	0.41
55:DA:1718:G:C3'	55:DA:1725:G:C5'	2.97	0.41
55:DA:2126:A:O2'	55:DA:2127:G:C5'	2.67	0.41
55:DA:2284:C:H41	28:D6:25:LYS:NZ	2.18	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2314:C:O2'	55:DA:2315:G:H5'	2.19	0.41
55:DA:2401:U:C2'	55:DA:2402:C:H5''	2.46	0.41
55:DA:2517:C:O2'	55:DA:2518:A:H3'	2.20	0.41
55:DA:387:U:C4'	55:DA:388:G:O5'	2.63	0.41
55:DA:467:G:O2'	55:DA:468:G:H5'	2.21	0.41
55:DA:50:U:C4'	55:DA:51:G:OP2	2.59	0.41
55:DA:707:G:H2'	55:DA:708:C:O4'	2.21	0.41
55:DA:863:A:O2'	55:DA:864:G:H5'	2.21	0.41
55:DA:568:U:H5'	55:DA:945:A:C2	2.56	0.41
55:DA:974:G:N2	55:DA:989:G:O2'	2.53	0.41
2:DB:21:G:H5'	2:DB:21:G:C8	2.52	0.41
3:DD:181:GLU:HA	3:DD:272:ALA:CB	2.40	0.41
5:DF:9:ILE:HA	5:DF:10:PRO:HD2	1.86	0.41
55:DA:2304:G:N2	6:DG:156:ASP:CG	2.60	0.41
6:DG:85:GLY:O	6:DG:86:MET:CB	2.68	0.41
8:DK:29:TYR:O	8:DK:33:ARG:HB2	2.20	0.41
8:DK:99:GLU:HG2	8:DK:103:ARG:HH21	1.80	0.41
58:DL:112:MET:HG2	58:DL:120:LEU:HB2	2.02	0.41
9:DM:42:TRP:HA	9:DM:48:MET:CE	2.51	0.41
11:DO:82:GLY:HA2	11:DO:113:LYS:O	2.21	0.41
2:DB:48:A:H4'	14:DQ:95:HIS:HD2	1.86	0.41
15:DR:33:LYS:HG3	15:DR:82:LEU:C	2.41	0.41
18:DS:110:LYS:HD2	18:DS:110:LYS:HA	1.78	0.41
18:DS:15:ARG:CA	18:DS:18:ARG:HD2	2.48	0.41
18:DS:11:ARG:NH2	18:DS:99:ARG:N	2.69	0.41
21:DV:154:ASP:HB2	21:DV:155:LEU:H	1.59	0.41
57:DY:21:GLN:NE2	57:DY:21:GLN:C	2.65	0.41
23:DZ:54:ALA:HB2	23:DZ:80:LEU:CD2	2.51	0.41
13:A0:51:LEU:HD22	13:A0:66:VAL:HG13	2.02	0.41
16:A1:25:TRP:C	16:A1:25:TRP:CD1	2.94	0.41
1:AA:993:G:H4'	17:A2:70:ILE:HD12	2.03	0.41
17:A2:85:LYS:CD	17:A2:86:GLY:H	2.34	0.41
1:AA:2331:G:O4'	22:A3:42:GLY:HA3	2.20	0.41
29:A7:24:THR:O	29:A7:26:GLY:N	2.53	0.41
1:AA:116:C:H2'	1:AA:117:G:H5'	2.02	0.41
1:AA:1268:A:H2'	1:AA:1269:A:O4'	2.21	0.41
1:AA:1275:A:O2'	1:AA:1276:A:H8	2.04	0.41
1:AA:1629:U:H2'	1:AA:1630:G:C8	2.56	0.41
1:AA:2080:G:H2'	1:AA:2081:C:H6	1.86	0.41
1:AA:2291:U:OP1	1:AA:2380:C:O2'	2.36	0.41
1:AA:2776:A:O2'	1:AA:2777:G:P	2.78	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:26:G:N1	1:AA:27:G:N2	2.69	0.41
1:AA:389:G:H22	11:AO:72:PRO:HG2	1.85	0.41
1:AA:608:A:N6	1:AA:609:A:C6	2.88	0.41
1:AA:71:A:C8	1:AA:71:A:H5'	2.55	0.41
1:AA:944:G:OP1	1:AA:945:A:H3'	2.21	0.41
1:AA:982:C:H6	1:AA:982:C:O5'	2.03	0.41
3:AD:134:ARG:HG3	3:AD:135:PHE:CD2	2.55	0.41
3:AD:4:LYS:HZ2	3:AD:20:ASP:HA	1.83	0.41
7:AH:15:VAL:HG23	7:AH:17:VAL:HG23	2.02	0.41
9:AM:134:ARG:O	9:AM:134:ARG:CG	2.67	0.41
11:AO:42:SER:O	11:AO:44:GLY:N	2.54	0.41
14:AQ:7:TYR:CE2	14:AQ:91:PRO:HG3	2.56	0.41
15:AR:11:GLU:O	15:AR:14:TYR:HD1	2.04	0.41
15:AR:50:ILE:HD12	15:AR:50:ILE:HA	1.85	0.41
18:AS:6:ILE:HG23	18:AS:103:ILE:O	2.20	0.41
18:AS:88:ARG:HH11	18:AS:88:ARG:HG2	1.85	0.41
24:AW:16:LEU:O	24:AW:20:GLU:HB2	2.21	0.41
25:AX:24:LYS:O	25:AX:25:ALA:C	2.59	0.41
25:AX:6:VAL:HG23	25:AX:35:ARG:O	2.20	0.41
23:AZ:90:ILE:HG23	23:AZ:91:LYS:N	2.35	0.41
31:BA:1102:A:C6	31:BA:1103:C:N4	2.89	0.41
31:BA:1135:U:HO2'	31:BA:1136:U:H5	1.65	0.41
31:BA:389:A:C2	31:BA:390:C:H1'	2.56	0.41
31:BA:432:A:H3'	31:BA:433:C:H6	1.84	0.41
31:BA:50:A:N6	31:BA:361:G:C4'	2.84	0.41
31:BA:597:G:H2'	31:BA:598:U:C5'	2.51	0.41
31:BA:865:A:H5'	31:BA:1078:U:C4	2.56	0.41
52:BC:42:C:H2'	52:BC:43:C:C5'	2.51	0.41
32:BE:167:PRO:CG	32:BE:188:ALA:HB2	2.42	0.41
32:BE:193:ASP:OD2	32:BE:196:LEU:HG	2.21	0.41
32:BE:189:ASP:HB3	32:BE:203:GLY:O	2.20	0.41
33:BF:19:GLU:O	33:BF:19:GLU:HG2	2.21	0.41
33:BF:97:LYS:O	33:BF:99:VAL:N	2.54	0.41
35:BH:91:LEU:HD12	35:BH:120:THR:HG22	2.03	0.41
36:BI:91:VAL:CG1	48:BU:72:ARG:NH2	2.84	0.41
36:BI:92:LYS:HE2	36:BI:92:LYS:HB3	1.86	0.41
37:BJ:18:TYR:O	37:BJ:19:GLY:C	2.58	0.41
37:BJ:36:LYS:NZ	37:BJ:36:LYS:HB2	2.36	0.41
40:BM:26:ALA:HB3	40:BM:85:LEU:HD21	2.02	0.41
42:BO:69:TYR:CD1	42:BO:70:ILE:N	2.89	0.41
43:BP:63:THR:HG22	43:BP:64:TRP:N	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:BQ:22:THR:HB	44:BQ:23:ARG:H	1.79	0.41
46:BS:45:THR:C	46:BS:47:ASP:H	2.24	0.41
46:BS:66:PRO:CG	46:BS:71:ARG:HG3	2.50	0.41
36:BI:89:MET:SD	48:BU:76:LEU:HD11	2.60	0.41
50:BW:10:LEU:HD22	50:BW:11:SER:N	2.36	0.41
53:C1:32:A:OP2	53:C1:33:G:OP2	2.39	0.41
53:C1:55:U:C6	53:C1:55:U:O5'	2.74	0.41
54:CA:1004:A:C6	54:CA:1025:U:C1'	3.03	0.41
54:CA:1308:U:O2'	54:CA:1309:G:H5'	2.21	0.41
54:CA:1224:G:C6	54:CA:1322:C:H1'	2.55	0.41
54:CA:1364:U:C2'	54:CA:1364:U:O2	2.66	0.41
54:CA:427:U:O5'	54:CA:427:U:H6	2.03	0.41
54:CA:630:G:C8	54:CA:630:G:H5''	2.53	0.41
54:CA:694:A:H2'	54:CA:695:A:O4'	2.21	0.41
54:CA:578:C:O2'	54:CA:728:A:H1'	2.20	0.41
54:CA:729:A:O2'	54:CA:730:G:H5'	2.21	0.41
54:CA:973:G:N3	40:CM:55:LYS:CE	2.84	0.41
54:CA:986:A:H1'	49:CV:54:GLY:O	2.21	0.41
52:CC:66:U:H3'	52:CC:67:C:H6	1.85	0.41
52:CD:18:G:C2'	52:CD:19:G:O5'	2.69	0.41
32:CE:130:ARG:HA	32:CE:131:PRO:HD3	1.92	0.41
32:CE:80:ILE:HG21	32:CE:211:ILE:HG22	2.02	0.41
36:CI:17:SER:O	36:CI:20:ALA:HB3	2.21	0.41
36:CI:5:GLU:HG3	36:CI:93:SER:OG	2.20	0.41
39:CL:118:LYS:O	39:CL:119:ALA:CB	2.62	0.41
40:CM:38:ILE:HG13	40:CM:71:LEU:HB3	2.02	0.41
40:CM:4:ILE:O	40:CM:74:ILE:HD13	2.20	0.41
41:CN:33:THR:HG21	41:CN:37:GLY:O	2.20	0.41
44:CQ:21:TYR:OH	44:CQ:23:ARG:NH2	2.54	0.41
48:CU:23:LYS:HB2	48:CU:56:THR:O	2.20	0.41
50:CW:33:ILE:HD13	50:CW:62:LEU:HB3	2.02	0.41
50:CW:84:LEU:C	50:CW:84:LEU:HD13	2.41	0.41
16:D1:74:LEU:CD1	16:D1:74:LEU:C	2.89	0.41
26:D4:55:ARG:O	26:D4:56:VAL:C	2.59	0.41
27:D5:40:LYS:CD	27:D5:46:CYS:HB3	2.50	0.41
30:D8:4:MET:O	30:D8:62:LEU:CD1	2.69	0.41
55:DA:1068:G:H4'	55:DA:1096:A:H2	1.85	0.41
55:DA:1083:U:H2'	55:DA:1084:A:H5'	2.02	0.41
55:DA:819:A:C4	55:DA:1189:A:C2	3.08	0.41
55:DA:1376:C:O2'	55:DA:1377:G:H5'	2.21	0.41
55:DA:1669:A:H2'	55:DA:1670:C:H5'	2.01	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1671:U:O2	55:DA:1673:U:H3'	2.21	0.41
55:DA:1707:G:H2'	55:DA:1708:C:C6	2.56	0.41
55:DA:1838:C:N4	55:DA:1898:U:H2'	2.36	0.41
55:DA:563:G:C4	55:DA:2018:G:C2	3.09	0.41
55:DA:2033:A:H2'	55:DA:2035:G:OP2	2.21	0.41
55:DA:2102:U:H2'	55:DA:2103:C:H6	1.81	0.41
55:DA:2110:G:O2'	55:DA:2111:C:P	2.79	0.41
55:DA:2695:C:H2'	55:DA:2696:U:C6	2.55	0.41
55:DA:270(G):C:H2'	55:DA:270(H):C:C6	2.55	0.41
55:DA:270(M):U:H1'	55:DA:270(N):G:C5	2.55	0.41
55:DA:2887:U:C2	55:DA:2888:C:C5	3.09	0.41
55:DA:358:U:H6	55:DA:358:U:O5'	2.03	0.41
55:DA:65:C:O2'	55:DA:66:C:H5'	2.21	0.41
55:DA:784:A:N7	3:DD:229:VAL:HG22	2.35	0.41
55:DA:977:G:C6	55:DA:987:G:C6	3.09	0.41
3:DD:137:PRO:HB2	3:DD:140:THR:HG23	2.03	0.41
4:DE:60:ASN:OD1	4:DE:61:ARG:N	2.54	0.41
6:DG:49:ASP:HB3	6:DG:52:ILE:CG1	2.51	0.41
56:DI:15:ALA:O	56:DI:19:GLU:CG	2.67	0.41
8:DK:144:VAL:O	8:DK:145:VAL:HG22	2.21	0.41
8:DK:21:VAL:O	8:DK:22:LYS:C	2.59	0.41
58:DL:18:THR:HG23	58:DL:42:ASN:OD1	2.20	0.41
9:DM:35:ARG:HD3	9:DM:37:LYS:HD3	2.01	0.41
10:DN:38:VAL:HA	10:DN:61:VAL:HA	2.03	0.41
11:DO:125:VAL:HG11	11:DO:138:LEU:HD22	2.03	0.41
11:DO:34:GLY:O	11:DO:35:HIS:HB2	2.21	0.41
55:DA:195:A:H5"	11:DO:46:LYS:NZ	2.36	0.41
14:DQ:52:SER:O	14:DQ:56:LEU:HD21	2.21	0.41
14:DQ:74:ALA:O	14:DQ:75:GLU:C	2.59	0.41
18:DS:88:ARG:HD2	18:DS:88:ARG:HA	1.80	0.41
19:DT:41:ASN:N	19:DT:41:ASN:ND2	2.66	0.41
21:DV:128:VAL:CG2	21:DV:129:SER:H	2.26	0.41
21:DV:61:LEU:C	21:DV:62:PRO:O	2.58	0.41
57:DY:135:ARG:O	57:DY:139:VAL:N	2.54	0.41
55:DA:1086:A:H2	57:DY:41:ARG:HH21	1.58	0.41
23:DZ:69:LYS:O	23:DZ:73:LEU:HG	2.20	0.41
16:A1:97:ASP:C	16:A1:98:LEU:O	2.59	0.41
17:A2:18:LEU:HD23	17:A2:18:LEU:C	2.41	0.41
26:A4:18:CYS:N	26:A4:36:CYS:SG	2.93	0.41
26:A4:52:THR:HG21	43:BP:65:LYS:CE	2.50	0.41
27:A5:31:VAL:HG13	27:A5:42:PRO:HG3	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:A8:61:LEU:O	30:A8:64:TYR:N	2.53	0.41
1:AA:1071:G:O4'	1:AA:1089:G:N7	2.54	0.41
1:AA:1150:C:O2'	1:AA:1151:G:H5'	2.21	0.41
1:AA:1179:C:C2'	1:AA:1180:C:H5''	2.51	0.41
1:AA:1277:G:O2'	13:A0:24:GLN:NE2	2.45	0.41
1:AA:1342:A:N7	1:AA:1345:C:C6	2.88	0.41
1:AA:1371:G:O2'	1:AA:1372:U:C5	2.60	0.41
1:AA:1410:G:H2'	1:AA:1411:C:H6	1.81	0.41
1:AA:1515:C:H2'	1:AA:1516:U:C6	2.54	0.41
1:AA:1352:U:O2	1:AA:1570:A:H2	2.04	0.41
1:AA:1606:G:H4'	1:AA:1608:A:C2	2.56	0.41
1:AA:1766:U:H2'	1:AA:1766:U:O2	2.19	0.41
1:AA:1824:G:C2'	1:AA:1825:A:H5'	2.50	0.41
1:AA:1917:U:H2'	1:AA:1918:A:O4'	2.21	0.41
1:AA:2060:A:O2'	1:AA:2061:G:OP2	2.35	0.41
1:AA:2127:G:C2'	1:AA:2128:C:H5''	2.51	0.41
1:AA:2199:A:N1	1:AA:2226:C:N4	2.69	0.41
1:AA:2681:C:C2'	1:AA:2682:U:OP2	2.68	0.41
1:AA:270(F):U:C2	1:AA:270(G):C:C5	3.09	0.41
1:AA:270(G):C:H2'	1:AA:270(H):C:H6	1.83	0.41
1:AA:270(K):C:C3'	1:AA:270(L):U:H5''	2.51	0.41
1:AA:278:A:O2'	1:AA:279:C:P	2.78	0.41
1:AA:2892:A:N7	1:AA:2893:G:C5	2.89	0.41
1:AA:29:U:H2'	1:AA:30:G:C8	2.56	0.41
1:AA:602:G:C2	1:AA:656:G:C6	3.09	0.41
1:AA:773:U:C5'	3:AD:47:GLY:HA2	2.51	0.41
1:AA:813:U:C2	1:AA:814:C:C5	3.09	0.41
1:AA:858:U:H5''	1:AA:859:G:OP2	2.20	0.41
1:AA:858:U:O2'	1:AA:2268:A:O2'	2.30	0.41
2:AB:8:U:H2'	2:AB:9:G:H8	1.85	0.41
4:AE:33:VAL:HG12	4:AE:90:THR:H	1.85	0.41
7:AH:102:ALA:HA	7:AH:117:PRO:CD	2.42	0.41
7:AH:91:GLY:O	7:AH:92:ILE:C	2.58	0.41
8:AK:145:VAL:O	8:AK:146:ALA:C	2.60	0.41
8:AK:46:ALA:O	8:AK:50:ARG:HG2	2.21	0.41
9:AM:111:PRO:HA	9:AM:114:ARG:HH12	1.79	0.41
1:AA:1666:G:H4'	10:AN:6:THR:HG23	2.02	0.41
11:AO:127:ALA:HB3	11:AO:130:PHE:CE2	2.56	0.41
12:AP:16:ARG:HH11	12:AP:16:ARG:CG	2.33	0.41
14:AQ:12:PHE:C	14:AQ:14:VAL:H	2.24	0.41
15:AR:6:LEU:CA	15:AR:9:LEU:HD12	2.50	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:AU:8:LYS:HG3	20:AU:94:LYS:HZ1	1.86	0.41
20:AU:91:GLU:CG	20:AU:92:ASN:N	2.75	0.41
21:AV:137:ILE:HG22	21:AV:138:GLU:N	2.36	0.41
21:AV:53:ILE:O	21:AV:70:LEU:HD21	2.21	0.41
25:AX:42:ALA:O	25:AX:45:GLY:N	2.53	0.41
31:BA:1026:G:C2'	31:BA:1027:C:H5'	2.50	0.41
31:BA:1154:G:C2	31:BA:1155:G:C8	3.09	0.41
31:BA:282:A:N7	31:BA:283:C:C5	2.89	0.41
31:BA:407:G:H2'	31:BA:408:A:C8	2.56	0.41
31:BA:66:G:N3	31:BA:66:G:H2'	2.35	0.41
31:BA:941:G:H2'	31:BA:942:G:O5'	2.21	0.41
52:BC:21:A:H2'	52:BC:46:G:O6	2.20	0.41
52:BD:70:G:H2'	52:BD:71:G:H8	1.86	0.41
32:BE:24:TRP:HA	32:BE:191:ASP:HA	2.03	0.41
33:BF:63:ASN:H	33:BF:97:LYS:HD2	1.85	0.41
34:BG:29:PRO:HD2	34:BG:30:LYS:CD	2.51	0.41
35:BH:7:GLU:OE1	35:BH:37:ARG:NE	2.52	0.41
37:BJ:31:MET:SD	37:BJ:34:GLY:HA2	2.61	0.41
38:BK:1:MET:CE	38:BK:1:MET:H3	2.34	0.41
40:BM:21:GLN:O	40:BM:21:GLN:HG2	2.20	0.41
40:BM:7:LYS:HG2	40:BM:71:LEU:HB2	2.03	0.41
43:BP:45:VAL:O	43:BP:45:VAL:HG12	2.21	0.41
44:BQ:22:THR:O	44:BQ:23:ARG:CB	2.69	0.41
46:BS:38:TYR:CZ	46:BS:50:LYS:CB	3.03	0.41
26:A4:63:TYR:CD2	49:BV:41:VAL:HG13	2.56	0.41
49:BV:19:VAL:HG11	49:BV:44:MET:HB3	1.97	0.41
51:BX:9:ARG:HH21	51:BX:10:ARG:CG	2.33	0.41
53:C1:53:U:O2'	53:C1:54:U:C5'	2.67	0.41
54:CA:1127:G:H2'	54:CA:1128:C:C6	2.56	0.41
54:CA:1237:C:C1'	54:CA:1334:G:H21	2.32	0.41
54:CA:1423:G:OP1	10:DN:49:ARG:NH2	2.54	0.41
54:CA:17:U:C2	54:CA:18:C:C5	3.09	0.41
54:CA:453:A:C6	54:CA:454:C:C4	3.08	0.41
54:CA:511:C:HO2'	54:CA:512:U:H6	1.68	0.41
54:CA:652:U:O2'	54:CA:653:A:C5'	2.69	0.41
54:CA:65:U:C5	54:CA:381:C:C4	3.09	0.41
54:CA:659:U:O2'	54:CA:660:G:H5'	2.21	0.41
54:CA:820:U:C4'	54:CA:821:G:OP2	2.55	0.41
52:CB:56:C:C5	55:DA:896:A:O2'	2.72	0.41
52:CD:42:C:H5'	52:CD:42:C:H6	1.85	0.41
33:CF:115:LEU:HD23	33:CF:118:GLN:OE1	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:CG:163:GLU:OE2	34:CG:163:GLU:HA	2.20	0.41
34:CG:196:LEU:CD1	34:CG:196:LEU:N	2.80	0.41
35:CH:33:VAL:HG12	35:CH:34:VAL:N	2.35	0.41
35:CH:79:GLU:OE2	38:CK:104:ARG:HA	2.20	0.41
36:CI:82:ARG:HB2	36:CI:85:VAL:CG2	2.50	0.41
38:CK:40:ALA:C	38:CK:42:GLU:N	2.74	0.41
38:CK:43:GLY:O	38:CK:64:LYS:HD2	2.21	0.41
39:CL:99:LEU:CD2	39:CL:99:LEU:N	2.84	0.41
40:CM:16:LEU:HD23	40:CM:94:VAL:HG13	2.03	0.41
43:CP:83:ASP:N	43:CP:93:ARG:NH2	2.64	0.41
46:CS:82:GLN:HB3	46:CS:82:GLN:HE21	1.53	0.41
47:CT:33:GLY:O	47:CT:34:LYS:C	2.58	0.41
47:CT:59:ILE:HG22	47:CT:71:PHE:HB3	2.01	0.41
47:CT:62:SER:HB3	47:CT:72:ARG:NH2	2.36	0.41
47:CT:92:ARG:O	47:CT:95:TYR:HB2	2.21	0.41
48:CU:47:THR:O	48:CU:83:GLU:N	2.52	0.41
48:CU:56:THR:CB	48:CU:58:LEU:HD13	2.49	0.41
55:DA:1287:A:N6	13:D0:106:GLY:O	2.50	0.41
16:D1:33:ARG:O	16:D1:37:GLU:HB2	2.20	0.41
22:D3:70:GLN:OE1	22:D3:72:ARG:HD3	2.20	0.41
26:D4:18:CYS:HB3	26:D4:19:GLY:H	1.18	0.41
55:DA:1082:U:O2'	58:DL:117:THR:CG2	2.69	0.41
55:DA:1311:G:C2	19:DT:60:ARG:NH1	2.80	0.41
55:DA:1392:A:C6	55:DA:1393:A:C6	3.09	0.41
55:DA:139:G:N2	55:DA:1596:A:H4'	2.34	0.41
55:DA:1760:A:C6	55:DA:1761:C:N4	2.89	0.41
55:DA:1862:G:O2'	55:DA:1863:G:H5'	2.21	0.41
55:DA:2250:G:H8	55:DA:2496:C:H5''	1.85	0.41
55:DA:2359:C:H2'	55:DA:2360:A:C8	2.56	0.41
55:DA:527:C:OP2	55:DA:2779:U:O4	2.39	0.41
55:DA:2789:C:HO2'	55:DA:2790:A:C4'	2.32	0.41
55:DA:531:C:H5''	55:DA:532:A:N9	2.36	0.41
55:DA:607:U:O4	55:DA:608:A:C6	2.74	0.41
55:DA:608:A:C8	55:DA:619:G:N2	2.89	0.41
55:DA:764:A:O4'	3:DD:213:ARG:HG3	2.21	0.41
55:DA:802:A:H2'	55:DA:803:U:H5''	2.03	0.41
55:DA:885:C:O4'	55:DA:885:C:OP1	2.39	0.41
2:DB:12:C:O2	22:D3:74:ARG:NH1	2.54	0.41
2:DB:66:A:O2'	2:DB:67:G:P	2.78	0.41
3:DD:137:PRO:HB2	3:DD:140:THR:CG2	2.50	0.41
55:DA:1567:A:OP2	3:DD:86:PRO:HB3	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:DE:104:VAL:HG11	4:DE:188:VAL:CG2	2.50	0.41
5:DF:150:GLY:HA2	5:DF:172:TRP:CE3	2.56	0.41
5:DF:36:VAL:HG11	5:DF:183:VAL:HG11	2.02	0.41
5:DF:63:LYS:CE	5:DF:67:GLN:HB2	2.51	0.41
6:DG:33:ARG:H	6:DG:162:THR:HG1	1.67	0.41
6:DG:39:ILE:HB	6:DG:92:VAL:HG13	2.03	0.41
7:DH:152:ARG:CG	7:DH:153:LYS:HE3	2.50	0.41
7:DH:55:PRO:HG2	7:DH:61:HIS:CE1	2.55	0.41
56:DI:27:LEU:HB2	56:DI:28:LYS:H	1.66	0.41
56:DI:29:GLU:OE2	56:DJ:6:GLU:OE2	2.38	0.41
56:DJ:8:ILE:HA	56:DJ:11:GLU:CB	2.51	0.41
8:DK:128:LEU:HD22	8:DK:128:LEU:HA	1.94	0.41
8:DK:37:VAL:CG1	8:DK:38:LEU:N	2.84	0.41
8:DK:47:LEU:O	8:DK:51:ILE:HG13	2.20	0.41
8:DK:78:THR:HB	8:DK:79:ILE:H	1.53	0.41
9:DM:15:LEU:HA	9:DM:53:VAL:HG23	2.03	0.41
14:DQ:102:ALA:O	14:DQ:105:ALA:HB3	2.20	0.41
15:DR:80:SER:HA	15:DR:81:PRO:HD3	1.86	0.41
19:DT:83:VAL:CG1	19:DT:87:GLN:HB2	2.51	0.41
20:DU:94:LYS:CE	20:DU:101:LYS:NZ	2.84	0.41
20:DU:38:ILE:HG22	20:DU:66:PRO:CA	2.50	0.41
21:DV:111:VAL:HG21	21:DV:146:ILE:HG13	1.99	0.41
57:DY:136:ALA:C	57:DY:139:VAL:HB	2.40	0.41
57:DY:40:LEU:O	57:DY:41:ARG:HB2	2.20	0.41
23:DZ:21:ARG:O	23:DZ:32:LYS:HA	2.21	0.41
13:A0:85:PRO:C	13:A0:87:TYR:H	2.25	0.41
16:A1:105:VAL:HG23	16:A1:106:PHE:H	1.86	0.41
17:A2:1:MET:CG	17:A2:42:GLY:HA3	2.51	0.41
22:A3:31:VAL:HB	22:A3:35:ASN:ND2	2.20	0.41
26:A4:24:THR:CG2	26:A4:25:TYR:H	2.29	0.41
27:A5:3:LYS:CA	27:A5:3:LYS:HE3	2.42	0.41
28:A6:18:ARG:HE	28:A6:44:ARG:HH12	1.67	0.41
28:A6:52:VAL:O	28:A6:53:LYS:C	2.60	0.41
1:AA:1042:G:H2'	1:AA:1043:C:H6	1.85	0.41
1:AA:1278:A:H2'	1:AA:1279:G:H8	1.85	0.41
1:AA:1500:G:C5	1:AA:1501:C:C5	3.09	0.41
1:AA:15:G:O2'	1:AA:16:G:H5'	2.21	0.41
1:AA:1601:G:H2'	1:AA:1602:U:O4'	2.21	0.41
1:AA:1340:U:H1'	1:AA:1603:A:H5'	2.03	0.41
1:AA:1904:G:O2'	1:AA:1905:C:H5'	2.21	0.41
1:AA:1948:G:C5'	1:AA:1948:G:C8	2.99	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:528:A:C2	1:AA:2043:C:H4'	2.56	0.41
1:AA:1493:C:C4	1:AA:2210:G:C8	3.09	0.41
1:AA:2248:C:H3'	1:AA:2249:U:H6	1.86	0.41
1:AA:2290:G:O2'	1:AA:2381:C:H1'	2.21	0.41
1:AA:2399:G:H2'	1:AA:2400:G:O4'	2.21	0.41
1:AA:2415:G:H2'	1:AA:2416:C:C6	2.56	0.41
1:AA:2586:C:O2'	1:AA:2587:A:H5'	2.20	0.41
1:AA:2672:G:C2'	1:AA:2673:G:H5''	2.50	0.41
1:AA:2688:U:O2	1:AA:2719:G:N1	2.53	0.41
1:AA:276:A:C8	1:AA:278:A:N7	2.89	0.41
1:AA:2848:G:O2'	1:AA:2867:G:N2	2.53	0.41
1:AA:324:A:N6	1:AA:338:G:O2'	2.47	0.41
3:AD:146:GLU:HA	3:AD:153:ALA:HA	2.03	0.41
1:AA:2820:A:H8	4:AE:109:LYS:HE3	0.70	0.41
4:AE:179:GLU:CB	4:AE:181:LEU:HD23	2.50	0.41
4:AE:53:PRO:HG2	4:AE:54:GLN:N	2.27	0.41
4:AE:91:VAL:HG22	4:AE:95:ILE:HD11	2.03	0.41
6:AG:178:PHE:HB3	6:AG:180:PHE:HE1	1.86	0.41
7:AH:168:PRO:HG2	7:AH:169:VAL:N	2.34	0.41
8:AK:7:GLU:O	8:AK:9:LEU:HD23	2.20	0.41
9:AM:66:LYS:O	9:AM:68:GLU:N	2.54	0.41
10:AN:68:GLU:HB3	10:AN:78:ARG:NH1	2.36	0.41
12:AP:133:ARG:CG	12:AP:133:ARG:NH1	2.83	0.41
14:AQ:24:LEU:H	14:AQ:24:LEU:HD22	1.85	0.41
19:AT:26:TYR:O	19:AT:81:VAL:HG22	2.21	0.41
19:AT:40:LYS:HA	19:AT:51:VAL:HG11	2.03	0.41
19:AT:50:LYS:O	19:AT:51:VAL:HB	2.21	0.41
23:AZ:69:LYS:HA	23:AZ:72:GLU:HB3	2.03	0.41
23:AZ:95:LEU:HD22	23:AZ:96:LYS:HG2	2.03	0.41
31:BA:1027:C:H2'	31:BA:1028:C:H6	1.80	0.41
31:BA:1124:G:HO2'	31:BA:1125:U:P	2.44	0.41
31:BA:1288:A:H2'	31:BA:1289:A:H8	1.86	0.41
31:BA:1502:A:H2	31:BA:1505:G:C2	2.35	0.41
31:BA:815:A:O2'	31:BA:1527:C:H1'	2.21	0.41
31:BA:191(C):G:C2	31:BA:191(D):U:H1'	2.56	0.41
31:BA:412:A:C6	34:BG:35:ARG:HB3	2.56	0.41
31:BA:484:G:O2'	31:BA:485:G:P	2.79	0.41
31:BA:632:A:OP2	31:BA:632:A:H3'	2.21	0.41
31:BA:803:G:H2'	31:BA:804:U:C6	2.56	0.41
52:BB:10:G:HO2'	52:BB:11:C:P	2.43	0.41
32:BE:28:PHE:CD1	32:BE:28:PHE:O	2.74	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:BF:18:TRP:NE1	44:BQ:54:PRO:CA	2.80	0.41
33:BF:19:GLU:O	33:BF:20:SER:HB2	2.21	0.41
34:BG:20:TYR:HA	34:BG:21:LEU:HD12	2.02	0.41
37:BJ:36:LYS:O	37:BJ:39:ALA:HB3	2.21	0.41
38:BK:111:ILE:O	38:BK:134:ILE:HB	2.21	0.41
39:BL:16:ARG:HB3	39:BL:18:PHE:CE1	2.56	0.41
31:BA:778:G:O2'	41:BN:120:ARG:O	2.33	0.41
44:BQ:12:ARG:CZ	44:BQ:14:PRO:CG	2.99	0.41
46:BS:20:VAL:CG2	46:BS:32:TYR:CB	2.99	0.41
50:BW:83:ARG:HH11	50:BW:83:ARG:HG2	1.86	0.41
54:CA:1021:G:H2'	54:CA:1022:G:O4'	2.21	0.41
54:CA:1073:U:H2'	54:CA:1074:G:H8	1.86	0.41
54:CA:1533:C:H2'	54:CA:1534:A:O4'	2.21	0.41
54:CA:1542:U:C3'	54:CA:1542:U:OP2	2.68	0.41
54:CA:47:C:H5"	54:CA:48:C:OP1	2.20	0.41
54:CA:503:C:C2	54:CA:504:C:C5	3.09	0.41
54:CA:598:U:H2'	54:CA:599:C:C6	2.55	0.41
54:CA:742:G:OP2	45:CR:35:ARG:NH2	2.53	0.41
52:CB:20:U:O2	52:CB:20:U:H2'	2.20	0.41
43:CP:126:LYS:CG	52:CC:26:A:OP2	2.69	0.41
52:CD:14:A:H2'	52:CD:15:G:C8	2.56	0.41
32:CE:204:ASN:ND2	32:CE:205:ASP:N	2.67	0.41
34:CG:4:TYR:CE2	34:CG:7:PRO:O	2.74	0.41
38:CK:105:ARG:O	38:CK:105:ARG:HD3	2.21	0.41
38:CK:100:ILE:CB	38:CK:125:ARG:HH12	2.34	0.41
38:CK:21:LYS:HB2	38:CK:21:LYS:HE2	1.94	0.41
54:CA:1349:A:OP2	39:CL:118:LYS:NZ	2.54	0.41
42:CO:46:LYS:HG2	42:CO:47:LYS:H	1.85	0.41
43:CP:54:VAL:HG12	43:CP:58:GLU:OE2	2.21	0.41
45:CR:2:PRO:HB2	45:CR:3:ILE:H	1.63	0.41
45:CR:58:MET:O	45:CR:59:MET:C	2.59	0.41
48:CU:19:LYS:HD2	48:CU:19:LYS:HA	1.63	0.41
48:CU:43:PHE:O	48:CU:51:LEU:HG	2.21	0.41
50:CW:104:LEU:HD12	50:CW:105:SER:N	2.35	0.41
50:CW:12:ALA:C	50:CW:14:LYS:N	2.74	0.41
13:D0:54:LEU:HD12	13:D0:54:LEU:HA	1.71	0.41
16:D1:98:LEU:HD23	16:D1:99:ALA:N	2.36	0.41
17:D2:9:GLY:O	17:D2:10:LYS:HG3	2.21	0.41
17:D2:30:GLY:O	17:D2:60:GLU:OE2	2.39	0.41
55:DA:1027:A:C6	55:DA:1126:A:C4	3.09	0.41
55:DA:1205:U:C4'	55:DA:1206:G:OP2	2.67	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1270:C:H5''	55:DA:1271:G:C5'	2.50	0.41
55:DA:1485:G:C8	55:DA:1485:G:H5'	2.56	0.41
55:DA:1508:A:O2'	55:DA:1509:C:O5'	2.39	0.41
55:DA:1512:G:C5	55:DA:1513:C:C4	3.08	0.41
55:DA:1827:C:O2'	55:DA:1828:G:H5'	2.20	0.41
55:DA:1893:C:H2'	55:DA:1894:C:H5'	2.03	0.41
55:DA:2335:A:HO2'	55:DA:2336:A:P	2.44	0.41
55:DA:252:G:OP2	11:DO:50:ARG:NH1	2.54	0.41
55:DA:2584:U:H6	55:DA:2585:U:N3	2.19	0.41
55:DA:2683:C:H2'	55:DA:2684:U:H6	1.85	0.41
55:DA:2723:C:H4'	13:DO:1:MET:CG	2.47	0.41
55:DA:2723:C:O5'	55:DA:2723:C:H6	2.02	0.41
55:DA:345:A:O2'	55:DA:347:A:N7	2.54	0.41
55:DA:483:A:H4'	20:DU:49:VAL:HG13	2.03	0.41
55:DA:605:C:H1'	55:DA:657:U:O2'	2.21	0.41
55:DA:607:U:O2	55:DA:620:G:C8	2.73	0.41
55:DA:687:C:C2'	55:DA:687:C:O2	2.68	0.41
55:DA:695:G:N2	55:DA:696:G:H1'	2.35	0.41
55:DA:782:A:N1	3:DD:226:MET:CE	2.83	0.41
55:DA:813:U:H2'	55:DA:814:C:H6	1.82	0.41
55:DA:887:A:H1'	55:DA:889:C:C4	2.56	0.41
55:DA:928:G:H3'	55:DA:929:G:C8	2.56	0.41
2:DB:77:U:P	21:DV:19:ARG:NH2	2.92	0.41
4:DE:70:ALA:O	4:DE:71:GLY:O	2.39	0.41
5:DF:89:VAL:O	5:DF:91:GLY:N	2.49	0.41
6:DG:115:ARG:HH11	6:DG:115:ARG:CB	2.33	0.41
6:DG:54:GLU:HA	6:DG:57:ALA:HB3	2.02	0.41
7:DH:26:VAL:HG11	7:DH:33:LEU:HB2	2.03	0.41
7:DH:86:GLU:HB2	7:DH:87:LEU:H	1.56	0.41
7:DH:89:ILE:N	7:DH:89:ILE:CD1	2.83	0.41
57:DY:132:ASP:OD2	56:DJ:10:GLU:OE1	2.38	0.41
58:DL:13:PRO:CD	58:DL:14:ALA:H	2.34	0.41
58:DL:52:ILE:HG12	58:DL:76:TYR:N	2.36	0.41
58:DL:64:SER:O	58:DL:65:PHE:CB	2.61	0.41
58:DL:80:LYS:CD	58:DL:80:LYS:C	2.88	0.41
9:DM:120:LEU:HD13	9:DM:122:VAL:HG23	2.01	0.41
11:DO:112:LEU:O	11:DO:128:HIS:HB2	2.21	0.41
11:DO:131:SER:N	11:DO:134:ALA:HB3	2.36	0.41
12:DP:4:PRO:HG3	12:DP:69:PHE:HE2	1.86	0.41
14:DQ:103:GLU:C	14:DQ:105:ALA:N	2.73	0.41
15:DR:41:ARG:HG2	15:DR:41:ARG:NH1	2.34	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:DV:123:ASP:O	21:DV:124:ILE:HB	2.20	0.41
24:DW:13:ALA:O	24:DW:14:ARG:C	2.59	0.41
57:DY:93:LEU:HG	57:DY:126:ALA:HB1	2.00	0.41
16:A1:83:LEU:CD1	16:A1:88:ILE:HD11	2.52	0.40
17:A2:6:LYS:H	17:A2:37:VAL:HB	1.86	0.40
22:A3:11:ARG:H	22:A3:11:ARG:HG2	1.62	0.40
22:A3:4:LYS:O	22:A3:5:LYS:C	2.60	0.40
28:A6:9:LEU:CD2	28:A6:10:LEU:N	2.84	0.40
28:A6:15:GLU:O	28:A6:16:CYS:CB	2.69	0.40
28:A6:26:ASN:OD1	28:A6:28:ARG:N	2.45	0.40
1:AA:1515:C:C2	1:AA:1516:U:C5	3.09	0.40
1:AA:1966:A:HO2'	1:AA:1967:C:P	2.44	0.40
1:AA:1966:A:O2'	1:AA:1967:C:P	2.80	0.40
1:AA:2054:A:C2	27:A5:8:LYS:HB2	2.56	0.40
1:AA:2276:G:OP2	12:AP:84:GLY:N	2.53	0.40
1:AA:2285:C:H5''	28:A6:28:ARG:NH1	2.37	0.40
1:AA:2481:G:C2'	1:AA:2482:G:OP2	2.69	0.40
1:AA:2503:A:O2'	1:AA:2505:G:P	2.79	0.40
1:AA:1669:A:H5''	1:AA:2550:G:OP1	2.21	0.40
1:AA:256:A:C2	1:AA:257:A:C4	3.10	0.40
1:AA:2674:G:H2'	1:AA:2675:A:O4'	2.21	0.40
1:AA:1754:C:O2	1:AA:2717:G:H5'	2.21	0.40
1:AA:2789:C:HO2'	1:AA:2790:A:C4'	2.31	0.40
1:AA:2840:C:H2'	1:AA:2841:C:C6	2.57	0.40
1:AA:321:G:HO2'	1:AA:340:A:HO2'	1.63	0.40
1:AA:270:A:C2	1:AA:366:C:H4'	2.56	0.40
1:AA:370:G:HO2'	1:AA:371:A:P	2.43	0.40
1:AA:388:G:N7	1:AA:390:A:H2'	2.37	0.40
1:AA:13:A:C2	1:AA:526:A:C8	3.09	0.40
1:AA:563:G:C6	1:AA:564:C:C4	3.10	0.40
1:AA:71:A:N3	1:AA:73:A:N6	2.69	0.40
1:AA:726:G:O2'	1:AA:727:A:O5'	2.39	0.40
1:AA:728:G:C4	1:AA:730:C:C5	3.09	0.40
1:AA:792:G:O2'	1:AA:2072:G:H1'	2.22	0.40
1:AA:1792:G:C5'	3:AD:205:VAL:HG13	2.52	0.40
3:AD:242:ARG:H	3:AD:242:ARG:CD	2.17	0.40
3:AD:49:ILE:HG12	3:AD:49:ILE:O	2.19	0.40
4:AE:197:ILE:O	4:AE:197:ILE:CG1	2.69	0.40
4:AE:22:PRO:HB2	4:AE:186:GLY:HA3	2.01	0.40
4:AE:44:TYR:O	4:AE:45:THR:HB	2.20	0.40
4:AE:93:VAL:O	4:AE:95:ILE:N	2.54	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:2304:G:N2	6:AG:156:ASP:OD2	2.39	0.40
8:AK:7:GLU:HG3	8:AK:9:LEU:N	2.36	0.40
8:AK:89:TYR:HB3	54:CA:368:U:N3	2.36	0.40
9:AM:51:PHE:CZ	9:AM:119:ARG:HD2	2.56	0.40
11:AO:112:LEU:C	11:AO:112:LEU:HD22	2.41	0.40
12:AP:14:ARG:O	12:AP:15:GLY:O	2.39	0.40
15:AR:1:MET:C	15:AR:3:ARG:H	2.24	0.40
15:AR:92:GLY:HA2	15:AR:117:ASP:N	2.36	0.40
20:AU:97:ARG:HG2	20:AU:97:ARG:HH11	1.86	0.40
31:BA:1025:U:O2'	31:BA:1026:G:C5'	2.68	0.40
31:BA:1072:G:H2'	31:BA:1073:U:O4'	2.21	0.40
31:BA:1288:A:H2'	31:BA:1289:A:O4'	2.22	0.40
31:BA:1326:C:H2'	31:BA:1327:C:C6	2.55	0.40
31:BA:1350:A:H2'	31:BA:1351:U:H6	1.85	0.40
31:BA:1360:A:H2'	31:BA:1361:G:C8	2.56	0.40
31:BA:1538:C:C4	31:BA:1539:C:C5	3.08	0.40
31:BA:252:U:C2	31:BA:253:U:C5	3.09	0.40
31:BA:109:A:C8	31:BA:326:G:H2'	2.56	0.40
31:BA:406:G:H2'	31:BA:407:G:C8	2.55	0.40
31:BA:542:G:H2'	31:BA:543:C:H6	1.86	0.40
31:BA:687:A:N1	31:BA:704:A:N7	2.69	0.40
52:BD:30:G:N2	52:BD:31:A:H1'	2.37	0.40
52:BD:64:A:N3	52:BD:65:G:H1'	2.36	0.40
32:BE:216:SER:C	32:BE:218:ALA:N	2.74	0.40
32:BE:236:TYR:HA	32:BE:239:VAL:CG2	2.50	0.40
32:BE:95:GLN:O	32:BE:96:ARG:O	2.39	0.40
33:BF:164:ARG:CG	33:BF:165:THR:H	2.29	0.40
35:BH:67:VAL:HG22	35:BH:68:GLU:O	2.21	0.40
38:BK:13:ILE:O	38:BK:15:ASN:N	2.55	0.40
38:BK:26:VAL:O	38:BK:27:PRO:C	2.60	0.40
38:BK:84:ARG:HH11	38:BK:84:ARG:HG2	1.87	0.40
39:BL:39:GLY:O	39:BL:41:VAL:N	2.54	0.40
41:BN:43:SER:HB3	41:BN:68:ALA:HB2	2.02	0.40
42:BO:12:ARG:HB2	42:BO:12:ARG:HE	1.64	0.40
43:BP:20:THR:C	43:BP:22:ILE:H	2.24	0.40
43:BP:23:TYR:HB3	43:BP:67:GLU:HA	2.03	0.40
43:BP:87:TYR:CE1	43:BP:91:ARG:HD3	2.56	0.40
44:BQ:35:ARG:HG3	44:BQ:36:PHE:N	2.36	0.40
54:CA:1005:A:H3'	54:CA:1006:C:C5'	2.50	0.40
54:CA:1027:C:H2'	54:CA:1028:C:H6	1.80	0.40
54:CA:124:G:H2'	54:CA:125:U:O4'	2.20	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:CA:323:U:H2'	54:CA:324:G:O4'	2.21	0.40
54:CA:12:U:H4'	54:CA:526:C:H4'	2.04	0.40
54:CA:532:A:H2'	54:CA:533:A:OP1	2.21	0.40
54:CA:784:C:H2'	54:CA:785:G:H8	1.86	0.40
54:CA:794:A:H1'	54:CA:1506:U:O4	2.21	0.40
52:CD:14:A:C5	52:CD:22:G:C2	3.09	0.40
32:CE:203:GLY:O	32:CE:204:ASN:C	2.59	0.40
33:CF:8:ILE:HG22	33:CF:12:LEU:HD23	2.03	0.40
34:CG:108:LEU:O	34:CG:176:LEU:HD22	2.21	0.40
54:CA:620:C:C2	34:CG:135:LEU:HG	2.56	0.40
34:CG:54:TYR:HE1	34:CG:209:ARG:HH22	1.69	0.40
34:CG:6:GLY:O	34:CG:8:VAL:HG23	2.20	0.40
37:CJ:78:ARG:NH1	37:CJ:79:ARG:O	2.54	0.40
38:CK:100:ILE:HA	38:CK:101:PRO:HD3	1.76	0.40
38:CK:4:ASP:C	38:CK:4:ASP:OD2	2.59	0.40
39:CL:61:ALA:HB1	39:CL:63:ILE:CD1	2.50	0.40
40:CM:62:HIS:CD2	40:CM:62:HIS:N	2.85	0.40
41:CN:99:GLN:HG2	41:CN:105:VAL:HG11	2.03	0.40
42:CO:84:LEU:HD22	42:CO:104:VAL:HG11	2.03	0.40
45:CR:12:ILE:C	45:CR:14:GLU:N	2.74	0.40
54:CA:254:G:N2	47:CT:16:GLN:HE21	2.18	0.40
47:CT:9:VAL:O	47:CT:21:VAL:HA	2.21	0.40
47:CT:77:VAL:O	47:CT:78:GLU:CB	2.63	0.40
47:CT:86:GLU:O	47:CT:87:LYS:C	2.60	0.40
50:CW:89:ARG:HG3	50:CW:89:ARG:NH2	2.34	0.40
13:D0:1:MET:SD	13:D0:1:MET:N	2.83	0.40
13:D0:81:ASP:O	13:D0:82:GLU:CG	2.68	0.40
17:D2:35:LEU:O	17:D2:36:PRO:C	2.60	0.40
22:D3:53:MET:HE3	22:D3:57:PHE:HA	2.04	0.40
27:D5:12:SER:C	27:D5:14:ALA:N	2.75	0.40
28:D6:24:GLU:HB3	28:D6:25:LYS:H	1.46	0.40
28:D6:33:LYS:C	28:D6:35:GLU:N	2.75	0.40
30:D8:2:PRO:O	30:D8:3:LYS:C	2.59	0.40
30:D8:65:GLU:N	30:D8:65:GLU:CD	2.73	0.40
55:DA:1046:A:O4'	55:DA:1046:A:N3	2.54	0.40
55:DA:1339:G:H21	55:DA:1603:A:H1'	1.86	0.40
55:DA:1388:G:C2'	55:DA:1389:G:H5'	2.51	0.40
55:DA:1535:U:H3	55:DA:1536:A:H3'	1.86	0.40
55:DA:1575:C:H2'	55:DA:1576:U:O4'	2.21	0.40
55:DA:1778:U:H3'	55:DA:1784:A:N6	2.36	0.40
55:DA:17:G:H2'	55:DA:18:C:H6	1.86	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:1899:G:O2'	55:DA:1900:A:O5'	2.40	0.40
55:DA:1952:A:C6	10:DN:22:ILE:HD12	2.56	0.40
55:DA:2245:U:H2'	55:DA:2436:G:OP2	2.21	0.40
55:DA:2443:C:H2'	55:DA:2444:G:H8	1.86	0.40
55:DA:2657:A:C2	55:DA:2665:A:C8	3.09	0.40
55:DA:727:A:C2	3:DD:9:TYR:CD2	3.10	0.40
55:DA:728:G:H4'	3:DD:13:ARG:HD2	2.03	0.40
55:DA:894:C:C2'	55:DA:895:U:H6	2.31	0.40
2:DB:12:C:H4'	2:DB:13:A:H5''	2.02	0.40
2:DB:40:U:H3'	2:DB:41:U:H5''	2.02	0.40
3:DD:35:LYS:NZ	3:DD:104:TYR:HD1	2.18	0.40
3:DD:155:LEU:O	3:DD:156:ALA:C	2.60	0.40
3:DD:133:LEU:HG	3:DD:189:CYS:O	2.20	0.40
3:DD:258:LYS:HB2	3:DD:258:LYS:HE3	1.86	0.40
3:DD:72:LYS:HB3	3:DD:72:LYS:HE3	1.90	0.40
4:DE:29:GLY:HA2	4:DE:180:ASN:CB	2.52	0.40
4:DE:4:ILE:HG22	4:DE:4:ILE:O	2.20	0.40
7:DH:83:TYR:HA	7:DH:135:GLY:H	1.86	0.40
7:DH:166:GLY:O	7:DH:167:GLU:C	2.59	0.40
56:DI:13:SER:O	56:DI:16:THR:CG2	2.66	0.40
56:DJ:8:ILE:HD12	56:DJ:8:ILE:HA	1.73	0.40
58:DL:50:ASP:H	58:DL:53:VAL:HG22	1.82	0.40
12:DP:66:ILE:N	12:DP:104:PHE:O	2.49	0.40
14:DQ:24:LEU:CD1	14:DQ:41:ASP:HB2	2.51	0.40
18:DS:14:PRO:HG2	18:DS:78:GLU:HG3	2.03	0.40
2:DB:77:U:OP1	21:DV:19:ARG:NH2	2.54	0.40
24:DW:71:ASN:O	24:DW:72:ALA:C	2.59	0.40
57:DY:74:LEU:O	57:DY:74:LEU:HD22	2.20	0.40
57:DY:88:ALA:O	57:DY:90:ALA:N	2.55	0.40
16:A1:91:ASP:OD2	16:A1:96:ALA:CB	2.70	0.40
17:A2:1:MET:H2	17:A2:16:PRO:HD3	1.86	0.40
22:A3:14:ARG:HB3	22:A3:15:ASP:H	1.75	0.40
22:A3:26:TYR:O	22:A3:29:GLN:HB2	2.21	0.40
22:A3:54:GLY:O	22:A3:56:ASP:N	2.54	0.40
27:A5:16:ARG:CG	27:A5:16:ARG:NH1	2.82	0.40
28:A6:26:ASN:C	28:A6:28:ARG:N	2.74	0.40
1:AA:1021:A:C8	1:AA:1021:A:H3'	2.55	0.40
1:AA:1095:A:C2'	1:AA:1095:A:N3	2.84	0.40
1:AA:1140:C:C5'	1:AA:1143:A:N6	2.83	0.40
1:AA:1453:A:H5''	1:AA:1454:U:OP2	2.21	0.40
1:AA:13:A:C5'	1:AA:14:A:OP1	2.66	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1822:G:O2'	1:AA:1823:G:H5'	2.22	0.40
1:AA:1921:G:O2'	1:AA:1922:G:H5'	2.21	0.40
1:AA:2228:G:H2'	1:AA:2229:C:O4'	2.21	0.40
1:AA:287:C:H2'	1:AA:288:C:C6	2.57	0.40
2:AB:40:U:C2'	2:AB:45:A:H61	2.33	0.40
3:AD:237:GLU:O	3:AD:239:ARG:N	2.54	0.40
3:AD:266:SER:O	3:AD:267:SER:O	2.39	0.40
5:AF:8:GLN:HG3	5:AF:126:VAL:CA	2.32	0.40
6:AG:170:ARG:O	6:AG:174:GLU:HB2	2.21	0.40
6:AG:64:THR:HG23	6:AG:66:GLN:N	2.21	0.40
9:AM:4:TYR:OH	9:AM:6:PRO:HA	2.22	0.40
10:AN:10:VAL:HG21	10:AN:17:ARG:HA	2.02	0.40
11:AO:62:LEU:HD21	30:A8:25:MET:CB	2.50	0.40
11:AO:88:LEU:HD11	11:AO:95:VAL:CB	2.51	0.40
12:AP:21:THR:O	12:AP:22:LYS:O	2.39	0.40
12:AP:2:LEU:H	12:AP:2:LEU:CD1	2.34	0.40
12:AP:3:MET:HA	12:AP:4:PRO:HD3	1.88	0.40
20:AU:76:CYS:O	20:AU:77:PRO:C	2.60	0.40
21:AV:109:ALA:O	21:AV:110:GLY:C	2.59	0.40
21:AV:122:ARG:H	21:AV:122:ARG:HG3	1.70	0.40
21:AV:127:LYS:HB3	21:AV:162:GLU:HB2	2.01	0.40
21:AV:177:PRO:C	21:AV:180:VAL:H	2.24	0.40
24:AW:10:LEU:HD13	24:AW:59:ARG:HD2	2.01	0.40
31:BA:1004:A:C5	31:BA:1025:U:C2	3.08	0.40
31:BA:1095:U:OP1	31:BA:1108:G:N1	2.53	0.40
31:BA:1275:A:C2'	31:BA:1276:G:H5'	2.51	0.40
31:BA:19:C:H5''	35:BH:86:ALA:HB3	2.02	0.40
31:BA:236:G:H2'	31:BA:237:C:C6	2.57	0.40
31:BA:468:A:H2'	31:BA:474:G:H5'	2.02	0.40
31:BA:476:G:O2'	31:BA:477:G:H5'	2.22	0.40
31:BA:547:A:O2'	31:BA:548:G:O4'	2.39	0.40
31:BA:554:C:H2'	31:BA:555:C:H6	1.86	0.40
52:BB:68:C:H2'	52:BB:69:G:C8	2.56	0.40
52:BC:17:C:H6	52:BC:17:C:H3'	1.86	0.40
52:BD:58:A:N6	52:BD:61:C:C2	2.90	0.40
32:BE:165:VAL:CG2	32:BE:166:ASP:N	2.83	0.40
32:BE:19:HIS:HB3	32:BE:20:GLU:H	1.58	0.40
32:BE:92:TYR:CE2	32:BE:151:GLY:CA	3.03	0.40
32:BE:97:TRP:O	32:BE:98:LEU:C	2.59	0.40
33:BF:119:ARG:HD3	33:BF:123:GLN:NE2	2.36	0.40
33:BF:130:VAL:O	33:BF:134:ILE:HG12	2.20	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:BF:134:ILE:HG23	33:BF:151:VAL:HB	2.04	0.40
33:BF:79:ARG:NE	33:BF:79:ARG:N	2.63	0.40
33:BF:76:VAL:O	33:BF:83:ARG:HG3	2.21	0.40
34:BG:108:LEU:CG	34:BG:110:PHE:HE1	2.34	0.40
34:BG:194:LEU:HD22	34:BG:194:LEU:N	2.37	0.40
35:BH:111:GLU:C	35:BH:113:ALA:N	2.75	0.40
35:BH:111:GLU:O	35:BH:113:ALA:N	2.48	0.40
35:BH:125:SER:O	35:BH:131:ILE:HD11	2.20	0.40
36:BI:12:PRO:HD3	36:BI:58:GLY:HA2	2.02	0.40
39:BL:28:VAL:HG13	39:BL:63:ILE:O	2.21	0.40
39:BL:83:ARG:HA	39:BL:86:VAL:HG12	2.03	0.40
42:BO:61:THR:OG1	42:BO:62:SER:N	2.55	0.40
45:BR:54:ARG:CG	45:BR:58:MET:HE2	2.51	0.40
45:BR:60:VAL:O	45:BR:64:ARG:HB2	2.21	0.40
48:BU:36:ASN:ND2	48:BU:39:VAL:HB	2.35	0.40
48:BU:55:ARG:HG3	48:BU:55:ARG:HH11	1.85	0.40
53:C1:44:U:HO2'	53:C1:45:U:P	2.44	0.40
54:CA:102:G:C6	54:CA:103:C:C4	3.09	0.40
54:CA:1218:C:H2'	54:CA:1219:U:H6	1.81	0.40
54:CA:1347:G:O2'	54:CA:1348:U:C6	2.74	0.40
54:CA:14:U:H2'	54:CA:16:A:OP2	2.21	0.40
54:CA:1532:U:O2	53:C1:40:U:O4	2.39	0.40
59:CA:2207:MG:MG	46:CS:24:ALA:HB1	1.47	0.40
54:CA:342:C:C2	54:CA:348:G:N2	2.89	0.40
54:CA:486:U:O2'	54:CA:487:A:H5'	2.22	0.40
54:CA:502:G:OP1	42:CO:118:SER:N	2.39	0.40
54:CA:510:A:N3	54:CA:543:C:H1'	2.36	0.40
54:CA:54:C:H2'	54:CA:54:C:O2	2.21	0.40
33:CF:172:ARG:O	33:CF:173:VAL:HG22	2.22	0.40
34:CG:133:VAL:HG11	34:CG:138:TYR:CD1	2.55	0.40
34:CG:19:LEU:N	34:CG:19:LEU:HD23	2.36	0.40
35:CH:105:VAL:N	35:CH:106:PRO:HD2	2.37	0.40
36:CI:44:GLY:HA2	36:CI:59:TYR:CZ	2.57	0.40
37:CJ:87:VAL:HG11	37:CJ:155:ARG:HA	2.03	0.40
38:CK:29:SER:CB	38:CK:32:LYS:HE2	2.52	0.40
39:CL:59:PHE:N	39:CL:59:PHE:CD1	2.88	0.40
40:CM:32:ALA:HB2	40:CM:76:ASN:HB2	2.03	0.40
46:CS:39:TYR:CD2	46:CS:73:LEU:HD11	2.56	0.40
47:CT:27:PHE:HD1	47:CT:28:PRO:O	2.04	0.40
51:CX:15:ARG:CG	51:CX:15:ARG:NH1	2.82	0.40
4:DE:118:LYS:HE3	13:D0:1:MET:HE3	2.03	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:DA:2331:G:H4'	22:D3:42:GLY:HA3	2.04	0.40
22:D3:40:GLN:NE2	22:D3:57:PHE:O	2.50	0.40
6:DG:5:VAL:HG22	26:D4:25:TYR:CD2	2.57	0.40
26:D4:2:LYS:HD2	26:D4:2:LYS:HA	1.92	0.40
55:DA:1040:C:H2'	55:DA:1041:C:C6	2.55	0.40
55:DA:1312:U:HO2'	55:DA:1313:U:P	2.43	0.40
55:DA:1482:U:C5'	55:DA:1483:G:OP2	2.58	0.40
55:DA:1695:G:H2'	55:DA:1696:G:O4'	2.21	0.40
55:DA:1982:C:H5''	55:DA:1983:C:OP2	2.21	0.40
55:DA:528:A:C2	55:DA:2043:C:O5'	2.75	0.40
55:DA:2563:U:H2'	55:DA:2565:A:OP2	2.20	0.40
55:DA:466:A:H2'	55:DA:467:G:H5'	2.03	0.40
55:DA:630:G:N2	55:DA:633:A:C8	2.88	0.40
55:DA:704:G:O2'	55:DA:726:G:N2	2.53	0.40
2:DB:15:A:C5'	2:DB:16:G:H8	2.32	0.40
3:DD:32:SER:HA	3:DD:36:PRO:HD3	2.02	0.40
4:DE:65:GLY:O	4:DE:67:PHE:N	2.55	0.40
4:DE:77:ILE:O	4:DE:78:LEU:O	2.39	0.40
5:DF:51:THR:HG21	5:DF:92:PRO:HD2	2.03	0.40
6:DG:102:PHE:CZ	6:DG:157:ILE:HD13	2.56	0.40
8:DK:144:VAL:HG22	8:DK:145:VAL:H	1.84	0.40
8:DK:52:ARG:O	8:DK:56:LYS:HB3	2.21	0.40
8:DK:77:LEU:HD13	8:DK:78:THR:O	2.21	0.40
58:DL:18:THR:HG23	58:DL:42:ASN:CG	2.41	0.40
58:DL:19:PRO:HB3	58:DL:25:PRO:HG2	2.04	0.40
58:DL:19:PRO:HB3	58:DL:34:ILE:HG12	2.04	0.40
9:DM:42:TRP:N	16:D1:64:ARG:HH22	2.19	0.40
12:DP:23:GLY:CA	12:DP:101:ARG:NH1	2.84	0.40
12:DP:10:ARG:O	12:DP:11:LYS:HB2	2.21	0.40
12:DP:54:MET:HB3	12:DP:55:VAL:H	1.76	0.40
20:DU:42:VAL:CG1	20:DU:43:ASN:N	2.83	0.40
20:DU:48:ALA:O	20:DU:49:VAL:C	2.59	0.40
21:DV:53:ILE:H	21:DV:71:VAL:HG13	1.84	0.40
21:DV:7:ALA:O	21:DV:61:LEU:HA	2.21	0.40
25:DX:6:VAL:HG11	25:DX:47:VAL:CG1	2.51	0.40
57:DY:135:ARG:HD2	57:DY:138:LEU:CG	2.51	0.40
56:DI:22:GLN:NE2	57:DY:140:GLY:O	2.54	0.40
22:A3:72:ARG:NH1	22:A3:72:ARG:CG	2.84	0.40
26:A4:11:PRO:HA	26:A4:25:TYR:H	1.84	0.40
28:A6:31:PRO:HG2	28:A6:33:LYS:HB2	2.03	0.40
30:A8:30:ARG:O	30:A8:31:HIS:ND1	2.55	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:1049:C:H42	7:AH:2:SER:CB	2.30	0.40
1:AA:1185:C:H5''	1:AA:1186:G:OP1	2.21	0.40
1:AA:1225:C:O3'	17:A2:85:LYS:CB	2.62	0.40
1:AA:1283:G:H2'	1:AA:1285:G:OP2	2.22	0.40
1:AA:1295:C:H2'	1:AA:1296:G:C8	2.56	0.40
1:AA:1360:A:C6	1:AA:1372:U:O4	2.74	0.40
1:AA:1527:G:H5''	1:AA:1528:A:OP1	2.22	0.40
1:AA:1419:A:C8	1:AA:1579:A:N6	2.89	0.40
1:AA:2305:A:C6	6:AG:154:GLY:HA3	2.56	0.40
1:AA:2591:C:H2'	1:AA:2592:G:C8	2.56	0.40
1:AA:2692:C:O2'	1:AA:2693:A:H5'	2.22	0.40
1:AA:1758:G:C2	1:AA:2696:U:C5'	3.03	0.40
1:AA:2756:U:O4	1:AA:2759:G:O6	2.39	0.40
1:AA:2778:A:C5'	1:AA:2779:U:OP2	2.61	0.40
1:AA:447:A:H5''	1:AA:448:U:OP1	2.21	0.40
1:AA:44:A:O2'	1:AA:45:G:H5'	2.21	0.40
1:AA:498:G:H21	20:AU:47:LYS:HZ1	1.69	0.40
1:AA:622:G:C2'	1:AA:623:G:H5'	2.52	0.40
1:AA:769:G:O2'	1:AA:770:G:H5'	2.21	0.40
1:AA:986:C:H2'	1:AA:987:G:H5'	2.02	0.40
2:AB:15:A:C2'	2:AB:16:G:OP1	2.70	0.40
2:AB:81:G:C8	2:AB:96:G:N2	2.90	0.40
3:AD:142:VAL:HG23	3:AD:193:VAL:CA	2.41	0.40
3:AD:35:LYS:HD3	3:AD:63:ARG:HG3	2.03	0.40
4:AE:12:THR:HB	4:AE:13:ARG:H	1.67	0.40
4:AE:111:ARG:NE	4:AE:160:TYR:CE1	2.81	0.40
4:AE:32:PRO:HB2	4:AE:33:VAL:H	1.70	0.40
4:AE:55:ASN:O	4:AE:57:LYS:N	2.44	0.40
6:AG:4:ASP:HB3	6:AG:5:VAL:H	1.75	0.40
7:AH:143:GLN:HE22	7:AH:147:ASN:CG	2.25	0.40
8:AK:19:VAL:HG22	8:AK:20:ASP:N	2.36	0.40
9:AM:95:PRO:C	9:AM:97:ARG:H	2.24	0.40
10:AN:11:ALA:O	10:AN:98:VAL:HG23	2.21	0.40
10:AN:43:VAL:HG21	10:AN:52:VAL:CG1	2.52	0.40
11:AO:84:ASN:CB	11:AO:116:GLY:HA3	2.51	0.40
11:AO:45:LEU:HD12	11:AO:45:LEU:HA	1.78	0.40
15:AR:24:PRO:HD3	15:AR:52:ILE:HD12	2.04	0.40
19:AT:12:VAL:HG13	19:AT:27:THR:O	2.20	0.40
21:AV:127:LYS:HE3	21:AV:162:GLU:HG3	2.03	0.40
21:AV:20:ARG:NH1	21:AV:20:ARG:HG2	2.33	0.40
25:AX:23:LEU:HD11	25:AX:53:LEU:HD12	2.02	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:AX:29:ARG:HG3	25:AX:29:ARG:NH1	2.36	0.40
23:AZ:67:ILE:HB	23:AZ:68:PRO:HD3	2.04	0.40
31:BA:1028(A):C:H2'	31:BA:1028(B):C:C6	2.56	0.40
31:BA:1345:U:C2	31:BA:1377:A:C6	3.09	0.40
31:BA:1346:A:C4	31:BA:1348:U:C2	3.10	0.40
31:BA:1506:U:O2'	31:BA:1507:A:OP1	2.37	0.40
31:BA:49:U:C2	31:BA:361:G:N2	2.90	0.40
31:BA:686:U:C2	31:BA:687:A:N7	2.90	0.40
31:BA:712:A:H2'	31:BA:713:G:O4'	2.21	0.40
31:BA:720:C:H6	31:BA:720:C:O5'	2.04	0.40
31:BA:778:G:C6	31:BA:779:C:C4	3.09	0.40
31:BA:849:C:H2'	31:BA:850:U:O4'	2.21	0.40
32:BE:59:GLU:CA	32:BE:221:LEU:HD13	2.50	0.40
32:BE:52:GLU:O	32:BE:53:ARG:C	2.59	0.40
33:BF:132:ARG:HE	33:BF:136:GLN:HE22	1.68	0.40
36:BI:15:ASP:OD2	34:CG:27:TYR:OH	2.39	0.40
36:BI:35:ALA:HB2	36:BI:67:MET:HB3	2.03	0.40
37:BJ:109:ASN:HA	37:BJ:119:ARG:HH21	1.86	0.40
40:BM:78:ASN:O	40:BM:80:LYS:N	2.55	0.40
43:BP:56:LEU:HD13	43:BP:60:VAL:HG21	2.04	0.40
46:BS:1:MET:O	46:BS:3:LYS:HG3	2.22	0.40
46:BS:39:TYR:CZ	46:BS:41:PRO:HA	2.56	0.40
46:BS:56:ALA:O	46:BS:60:LEU:HG	2.22	0.40
54:CA:1245:A:OP2	51:CX:9:ARG:NH2	2.49	0.40
54:CA:363:A:H62	42:CO:28:LYS:CD	2.34	0.40
54:CA:375:U:C2	54:CA:376:G:C8	3.10	0.40
54:CA:112:G:C5'	54:CA:389:A:H4'	2.46	0.40
54:CA:401:C:H2'	54:CA:402:G:H8	1.87	0.40
54:CA:865:A:H2'	54:CA:866:C:O4'	2.22	0.40
52:CD:58:A:N6	52:CD:61:C:C1'	2.84	0.40
34:CG:62:GLN:OE1	34:CG:65:ARG:HD3	2.21	0.40
35:CH:103:GLY:C	35:CH:106:PRO:HD2	2.42	0.40
35:CH:41:VAL:HG11	35:CH:113:ALA:HB2	2.01	0.40
36:CI:52:ILE:O	36:CI:53:ALA:HB3	2.22	0.40
37:CJ:99:LEU:HD23	37:CJ:102:ARG:NH1	2.36	0.40
39:CL:112:LYS:C	39:CL:113:LYS:HD2	2.42	0.40
40:CM:89:ASP:C	40:CM:91:PRO:HD3	2.42	0.40
42:CO:28:LYS:O	42:CO:30:ALA:N	2.55	0.40
45:CR:78:TYR:CZ	45:CR:82:ILE:HD12	2.55	0.40
47:CT:10:VAL:HG13	47:CT:19:VAL:HB	2.03	0.40
50:CW:89:ARG:C	50:CW:91:LEU:H	2.24	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:D2:49:THR:O	17:D2:50:PRO:C	2.60	0.40
6:DG:67:LYS:CB	26:D4:5:ILE:HG22	2.52	0.40
55:DA:1027:A:H61	55:DA:1126:A:C1'	2.34	0.40
55:DA:1060:U:C5	58:DL:74:ALA:HB1	2.55	0.40
55:DA:1057:A:C8	55:DA:1086:A:C2'	3.01	0.40
55:DA:1107:G:H2'	55:DA:1108:U:C6	2.57	0.40
55:DA:1221:C:H2'	55:DA:1222:C:H6	1.85	0.40
55:DA:1272:A:H3'	55:DA:1273:U:H5'	2.02	0.40
55:DA:1387:C:C2	55:DA:1388:G:C8	3.09	0.40
55:DA:1523:U:H2'	55:DA:1524:G:H8	1.86	0.40
55:DA:2129:C:C2'	55:DA:2130:U:H5'	2.51	0.40
55:DA:217:G:H2'	55:DA:218:A:O4'	2.21	0.40
55:DA:2314:C:C2	55:DA:2315:G:C8	3.09	0.40
55:DA:2340:G:H2'	55:DA:2341:G:H8	1.86	0.40
55:DA:2545:G:H2'	55:DA:2546:U:O4'	2.20	0.40
55:DA:2688:U:H1'	55:DA:2721:A:H62	1.85	0.40
55:DA:2787:C:H2'	55:DA:2787:C:O2	2.21	0.40
55:DA:2838:G:H1'	13:D0:45:ARG:NH2	2.31	0.40
55:DA:31:C:H4'	55:DA:1238:G:H4'	2.03	0.40
55:DA:364:C:O2	55:DA:364:C:H2'	2.21	0.40
55:DA:448:U:O4	55:DA:583:G:H1'	2.22	0.40
55:DA:637:A:HO2'	55:DA:638:G:P	2.44	0.40
2:DB:85:G:C2	2:DB:86:G:C8	3.09	0.40
3:DD:92:ILE:HG22	3:DD:105:ILE:O	2.21	0.40
3:DD:112:GLN:OE1	3:DD:115:GLN:OE1	2.40	0.40
3:DD:241:PRO:O	3:DD:242:ARG:HB2	2.22	0.40
4:DE:103:ASP:OD2	4:DE:168:MET:HE1	2.21	0.40
8:DK:144:VAL:CG2	8:DK:145:VAL:N	2.82	0.40
58:DL:107:ILE:HD13	58:DL:107:ILE:HG21	1.78	0.40
58:DL:125:ARG:O	58:DL:128:ALA:CB	2.70	0.40
58:DL:132:ARG:HD3	58:DL:137:GLU:OE2	2.21	0.40
58:DL:49:GLY:HA3	58:DL:50:ASP:CB	2.31	0.40
58:DL:9:LYS:NZ	58:DL:9:LYS:HB3	2.37	0.40
9:DM:112:LEU:O	9:DM:114:ARG:O	2.38	0.40
9:DM:57:ALA:O	9:DM:58:ASP:CB	2.69	0.40
9:DM:57:ALA:C	9:DM:58:ASP:OD1	2.59	0.40
10:DN:3:GLN:HB2	10:DN:4:PRO:HD2	2.03	0.40
11:DO:50:ARG:HB2	11:DO:50:ARG:NH2	2.36	0.40
14:DQ:103:GLU:CA	14:DQ:103:GLU:OE1	2.70	0.40
14:DQ:106:ARG:H	14:DQ:106:ARG:HG3	1.67	0.40
15:DR:88:ILE:HD12	15:DR:88:ILE:C	2.41	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:DT:31:HIS:CD2	19:DT:33:LYS:N	2.89	0.40
20:DU:101:LYS:HG2	20:DU:101:LYS:O	2.21	0.40
21:DV:102:LEU:HD12	21:DV:122:ARG:HA	2.03	0.40
21:DV:128:VAL:CG2	21:DV:129:SER:N	2.81	0.40
57:DY:18:GLU:O	57:DY:19:ARG:CB	2.69	0.40
57:DY:17:LEU:CD1	57:DY:25:PHE:HE2	2.35	0.40
57:DY:35:LYS:HA	57:DY:35:LYS:HZ2	1.83	0.40
57:DY:58:LEU:C	57:DY:62:ALA:HB2	2.40	0.40
16:A1:101:ARG:O	16:A1:102:GLU:CG	2.70	0.40
17:A2:47:VAL:O	17:A2:47:VAL:HG13	2.22	0.40
26:A4:12:ALA:N	26:A4:24:THR:CB	2.85	0.40
26:A4:53:GLU:OE2	26:A4:58:ARG:CB	2.62	0.40
26:A4:55:ARG:CG	26:A4:55:ARG:O	2.70	0.40
27:A5:40:LYS:CE	27:A5:46:CYS:H	2.33	0.40
1:AA:2017:U:H4'	27:A5:8:LYS:O	2.20	0.40
29:A7:8:ASN:ND2	29:A7:10:ARG:N	2.68	0.40
1:AA:1000:A:H8	1:AA:1000:A:H5'	1.86	0.40
1:AA:1140:C:H1'	1:AA:1143:A:C8	2.56	0.40
1:AA:117:G:C6	1:AA:119:A:C6	3.10	0.40
1:AA:1210:A:P	1:AA:1212:G:H5'	2.61	0.40
1:AA:1263:U:O3'	27:A5:11:THR:OG1	2.40	0.40
1:AA:1309:G:H2'	1:AA:1310:G:C5'	2.52	0.40
1:AA:1889:A:O2'	1:AA:2087:G:H5'	2.21	0.40
1:AA:2157:G:O2'	1:AA:2158:A:O4'	2.40	0.40
1:AA:910:A:H2'	1:AA:2264:C:O2'	2.22	0.40
1:AA:2271:G:OP1	22:A3:18:ALA:HB1	2.22	0.40
1:AA:2275:C:O2	12:AP:83:MET:CG	2.67	0.40
1:AA:2354:G:N3	1:AA:2354:G:H2'	2.36	0.40
1:AA:2556:C:H2'	1:AA:2557:G:C5'	2.52	0.40
1:AA:2718:G:C6	1:AA:2719:G:C5	3.08	0.40
1:AA:2892:A:N6	1:AA:2893:G:C2	2.90	0.40
1:AA:307:G:H21	1:AA:330:A:N6	2.14	0.40
1:AA:602:G:N2	1:AA:656:G:C5	2.89	0.40
1:AA:663:G:C5	1:AA:664:C:C5	3.09	0.40
1:AA:721:C:H2'	1:AA:721:C:O2	2.20	0.40
1:AA:858:U:HO2'	1:AA:2268:A:C1'	2.33	0.40
1:AA:898:C:C5	1:AA:899:A:N7	2.90	0.40
6:AG:153:ARG:NH1	6:AG:153:ARG:HG2	2.37	0.40
6:AG:34:LEU:CD2	6:AG:159:VAL:HG23	2.50	0.40
6:AG:169:ALA:O	6:AG:170:ARG:C	2.59	0.40
6:AG:83:ARG:HG2	6:AG:83:ARG:HH11	1.87	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:AK:109:ILE:HB	8:AK:110:ASP:H	1.67	0.40
8:AK:142:VAL:CG2	8:AK:143:SER:N	2.55	0.40
8:AK:69:LYS:O	8:AK:73:GLU:HB2	2.22	0.40
9:AM:115:ARG:HG2	9:AM:115:ARG:NH1	2.36	0.40
9:AM:46:VAL:HG13	9:AM:48:MET:HG3	2.03	0.40
10:AN:2:ILE:CD1	10:AN:6:THR:HG21	2.44	0.40
12:AP:55:VAL:CG1	12:AP:56:ARG:N	2.84	0.40
15:AR:49:VAL:O	15:AR:49:VAL:HG13	2.21	0.40
18:AS:36:LEU:HD13	18:AS:48:ALA:CA	2.52	0.40
18:AS:92:ARG:HD3	18:AS:94:ASP:OD2	2.20	0.40
21:AV:104:PHE:C	21:AV:105:VAL:CG1	2.88	0.40
23:AZ:87:PRO:HA	23:AZ:90:ILE:CG2	2.27	0.40
53:B1:43:U:H6	53:B1:43:U:C5'	2.30	0.40
31:BA:1065:U:C2'	31:BA:1066:C:OP2	2.69	0.40
31:BA:1207:G:H2'	31:BA:1208:C:H6	1.86	0.40
31:BA:1219:U:C4	31:BA:1220:G:N7	2.89	0.40
31:BA:123:C:OP1	31:BA:311:C:O2'	2.27	0.40
31:BA:1405:G:H1'	31:BA:1519:A:C4'	2.52	0.40
31:BA:1541:U:O2'	31:BA:1542:U:H5'	2.21	0.40
31:BA:338:A:N6	31:BA:351:G:H1	2.19	0.40
31:BA:342:C:C4	31:BA:343:U:C4	3.09	0.40
31:BA:392:G:H2'	31:BA:393:A:C8	2.56	0.40
31:BA:457:C:N4	31:BA:458:C:N4	2.69	0.40
31:BA:543:C:OP1	34:BG:14:ARG:CD	2.70	0.40
31:BA:69:G:C2	31:BA:73:G:N7	2.90	0.40
52:BC:25:C:O2'	52:BC:26:A:H5'	2.21	0.40
52:BC:66:U:N3	52:BC:67:C:C4	2.90	0.40
32:BE:209:ARG:HG3	32:BE:240:GLN:NE2	2.36	0.40
33:BF:116:VAL:HG21	33:BF:141:VAL:HG22	2.03	0.40
33:BF:178:LEU:C	33:BF:180:ALA:N	2.72	0.40
34:BG:209:ARG:HG3	34:BG:209:ARG:NH1	2.36	0.40
34:BG:26:CYS:HA	34:BG:31:CYS:CA	2.51	0.40
37:BJ:50:ILE:O	37:BJ:52:GLU:N	2.55	0.40
37:BJ:6:ARG:O	37:BJ:7:ALA:C	2.60	0.40
39:BL:4:TYR:CE2	39:BL:88:TYR:CB	3.04	0.40
42:BO:79:GLU:HG3	42:BO:80:HIS:CD2	2.56	0.40
43:BP:19:LEU:HD23	43:BP:19:LEU:H	1.86	0.40
43:BP:23:TYR:O	43:BP:66:LEU:HB2	2.21	0.40
43:BP:81:LEU:HB3	43:BP:89:GLY:HA2	2.02	0.40
45:BR:6:GLU:CD	45:BR:6:GLU:H	2.24	0.40
45:BR:9:GLN:O	45:BR:10:LYS:C	2.60	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:BV:20:LEU:C	49:BV:22:LEU:N	2.73	0.40
51:BX:8:THR:CG2	51:BX:9:ARG:N	2.85	0.40
54:CA:1146:A:H8	54:CA:1146:A:H5'	1.85	0.40
54:CA:1160:G:N3	54:CA:1160:G:H2'	2.36	0.40
54:CA:1339:A:H2'	54:CA:1340:A:O4'	2.21	0.40
54:CA:177:C:O2'	54:CA:178:C:H5'	2.22	0.40
54:CA:184:G:O2'	54:CA:185:A:H5'	2.22	0.40
54:CA:436:C:H2'	54:CA:437:U:O4'	2.20	0.40
54:CA:483:C:H6	54:CA:483:C:O5'	2.05	0.40
54:CA:659:U:H2'	54:CA:660:G:C8	2.57	0.40
54:CA:96:G:C6	54:CA:97:U:O2	2.74	0.40
32:CE:100:GLY:O	32:CE:102:LEU:N	2.55	0.40
32:CE:68:ILE:O	32:CE:90:MET:HB3	2.21	0.40
33:CF:46:GLU:C	33:CF:48:TYR:H	2.25	0.40
34:CG:79:PHE:CZ	34:CG:204:ILE:HA	2.57	0.40
35:CH:105:VAL:H	35:CH:106:PRO:HD2	1.85	0.40
36:CI:3:ARG:HG3	36:CI:3:ARG:HH11	1.86	0.40
36:CI:75:LEU:HD21	36:CI:79:LEU:HD11	2.03	0.40
40:CM:12:ASP:OD2	40:CM:14:LYS:HB3	2.21	0.40
43:CP:12:ASN:N	43:CP:12:ASN:OD1	2.54	0.40
45:CR:76:GLU:C	45:CR:78:TYR:N	2.74	0.40
54:CA:958:A:C8	49:CV:55:LYS:HD2	2.56	0.40
13:D0:48:VAL:O	13:D0:49:ASP:C	2.59	0.40
16:D1:95:LEU:HD13	17:D2:4:ILE:CG2	2.51	0.40
55:DA:1163:G:P	17:D2:24:LYS:HZ1	2.45	0.40
28:D6:28:ARG:CB	28:D6:28:ARG:HH11	2.28	0.40
55:DA:1179:C:H2'	55:DA:1180:C:C4'	2.50	0.40
55:DA:1372:U:C6	55:DA:1372:U:C3'	3.04	0.40
55:DA:1448:G:O2'	55:DA:1528:A:N6	2.54	0.40
55:DA:1648:C:C2	55:DA:1649:G:C8	3.09	0.40
55:DA:1794:U:C2	55:DA:1795:C:C5	3.10	0.40
55:DA:2136:C:H6	55:DA:2136:C:O5'	2.05	0.40
55:DA:2302:G:C6	55:DA:2315:G:C6	3.10	0.40
55:DA:2498:C:O2'	55:DA:2499:C:H5'	2.22	0.40
55:DA:2562:U:C2'	55:DA:2563:U:H5'	2.51	0.40
55:DA:2584:U:O2	55:DA:2584:U:O4'	2.38	0.40
55:DA:2754:U:H5''	55:DA:2755:C:OP2	2.19	0.40
55:DA:2859:G:O2'	55:DA:2860:A:H5'	2.22	0.40
55:DA:868:U:H2'	55:DA:869:G:O4'	2.20	0.40
55:DA:878:A:C2	55:DA:879:G:C8	3.10	0.40
55:DA:885:C:N4	55:DA:892:G:C6	2.86	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:DD:10:THR:C	3:DD:11:PRO:O	2.57	0.40
3:DD:120:GLY:HA3	3:DD:122:ASP:OD2	2.22	0.40
3:DD:132:PRO:O	3:DD:133:LEU:C	2.58	0.40
3:DD:9:TYR:CZ	3:DD:13:ARG:HD3	2.56	0.40
4:DE:93:VAL:C	4:DE:95:ILE:N	2.74	0.40
5:DF:33:LEU:HD12	5:DF:33:LEU:HA	1.94	0.40
6:DG:9:ARG:C	6:DG:11:TYR:N	2.75	0.40
7:DH:121:ILE:CG2	7:DH:122:THR:N	2.84	0.40
7:DH:121:ILE:HG22	7:DH:122:THR:N	2.36	0.40
7:DH:59:ARG:CG	7:DH:59:ARG:NH1	2.83	0.40
8:DK:144:VAL:O	8:DK:145:VAL:CG2	2.69	0.40
58:DL:45:THR:O	58:DL:49:GLY:HA2	2.21	0.40
58:DL:52:ILE:CD1	58:DL:76:TYR:CB	2.53	0.40
58:DL:83:GLY:N	58:DL:99:ILE:HD13	2.36	0.40
9:DM:67:LEU:O	9:DM:88:GLU:CG	2.58	0.40
11:DO:3:LEU:C	11:DO:5:ASP:H	2.25	0.40
12:DP:52:VAL:O	12:DP:53:ALA:C	2.58	0.40
14:DQ:14:VAL:HG13	14:DQ:15:ARG:N	2.35	0.40
15:DR:34:VAL:CG1	15:DR:35:LYS:N	2.83	0.40
18:DS:18:ARG:HH11	18:DS:18:ARG:HG2	1.87	0.40
18:DS:24:ILE:CD1	18:DS:24:ILE:C	2.90	0.40
18:DS:29:LEU:HD21	18:DS:33:ARG:NE	2.36	0.40
21:DV:158:PRO:HA	21:DV:159:PRO:HD3	1.93	0.40
21:DV:82:ARG:CG	21:DV:82:ARG:HH11	2.32	0.40
57:DY:139:VAL:CG1	57:DY:140:GLY:N	2.84	0.40
57:DY:75:GLN:NE2	57:DY:75:GLN:HA	2.37	0.40
1:AA:1023:U:OP2	1:AA:1024:G:N7	2.54	0.40
1:AA:1060:U:OP1	1:AA:1060:U:H6	2.05	0.40
1:AA:110:G:C2	1:AA:111:A:C8	3.10	0.40
1:AA:1379:A:HO2'	1:AA:1380:G:P	2.43	0.40
1:AA:1409:C:O2'	1:AA:1410:G:H5'	2.21	0.40
1:AA:1439:A:H2'	1:AA:1440:G:O4'	2.21	0.40
1:AA:1516:U:C2	1:AA:1517:G:C8	3.09	0.40
1:AA:1848:A:H2'	1:AA:1849:G:O4'	2.22	0.40
1:AA:1951:U:H2'	1:AA:1953:A:OP2	2.21	0.40
1:AA:2033:A:O2'	1:AA:2034:U:OP1	2.40	0.40
1:AA:2173:A:H5''	1:AA:2174:C:H5	1.83	0.40
1:AA:2243:U:H2'	1:AA:2244:U:C6	2.56	0.40
1:AA:2321:G:N3	1:AA:2321:G:H2'	2.36	0.40
1:AA:2359:C:H2'	1:AA:2360:A:O4'	2.21	0.40
1:AA:2889:C:H3'	1:AA:2891:G:C8	2.56	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:363(A):A:C3'	1:AA:363(B):G:C5'	3.00	0.40
1:AA:396:G:O5'	1:AA:396:G:H8	2.05	0.40
1:AA:439:G:H2'	1:AA:440:G:C8	2.57	0.40
1:AA:527:C:O4'	1:AA:527:C:O2	2.35	0.40
1:AA:632:A:O2'	1:AA:2404:C:H5'	2.22	0.40
1:AA:67:U:H2'	1:AA:68:G:C8	2.54	0.40
1:AA:752:A:O2'	1:AA:753:C:P	2.76	0.40
1:AA:976:C:H4'	1:AA:1156:A:N7	2.35	0.40
2:AB:40:U:C2	26:A4:1:MET:CE	3.04	0.40
2:AB:73:A:N3	2:AB:73:A:H2'	2.36	0.40
3:AD:195:ALA:O	3:AD:196:VAL:C	2.59	0.40
3:AD:33:LEU:N	3:AD:35:LYS:O	2.41	0.40
3:AD:35:LYS:CG	3:AD:64:ILE:HG23	2.51	0.40
4:AE:179:GLU:HG3	4:AE:181:LEU:HD23	2.03	0.40
4:AE:60:ASN:ND2	4:AE:61:ARG:H	2.19	0.40
5:AF:64:ILE:HG13	5:AF:65:TRP:CG	2.57	0.40
7:AH:109:PHE:CZ	7:AH:152:ARG:HD3	2.56	0.40
7:AH:33:LEU:HD21	7:AH:136:ILE:O	2.21	0.40
7:AH:32:GLU:O	7:AH:33:LEU:HD23	2.22	0.40
8:AK:82:ARG:HE	54:CA:56:U:C4'	2.23	0.40
8:AK:97:ILE:HD12	8:AK:97:ILE:H	1.86	0.40
11:AO:85:LEU:HD23	11:AO:85:LEU:N	2.32	0.40
14:AQ:108:GLY:O	14:AQ:110:LEU:HB2	2.20	0.40
14:AQ:107:GLU:H	14:AQ:110:LEU:HD12	1.87	0.40
14:AQ:38:GLN:HG3	14:AQ:47:THR:HG21	2.03	0.40
18:AS:82:LEU:HD22	18:AS:84:ARG:HH22	1.86	0.40
20:AU:20:TYR:CZ	20:AU:42:VAL:HA	2.56	0.40
21:AV:18:LEU:HD12	21:AV:18:LEU:N	2.37	0.40
23:AZ:41:ARG:NH1	23:AZ:41:ARG:HG3	2.36	0.40
31:BA:1057:G:C2'	31:BA:1058:G:H5'	2.51	0.40
31:BA:1079:G:C6	31:BA:1080:A:N6	2.89	0.40
31:BA:1174:G:C2	31:BA:1175:G:C5	3.10	0.40
31:BA:1363:A:N3	31:BA:1365:G:O6	2.55	0.40
31:BA:1512:U:C2	31:BA:1513:A:N7	2.90	0.40
31:BA:324:G:N1	31:BA:327:A:OP2	2.54	0.40
31:BA:52:G:O2'	31:BA:53:A:H5'	2.20	0.40
31:BA:554:C:C2	31:BA:555:C:C5	3.10	0.40
31:BA:4:U:O2'	31:BA:5:U:OP1	2.35	0.40
31:BA:701:C:H4'	31:BA:702:A:H5''	2.02	0.40
31:BA:754:C:O2	31:BA:754:C:H3'	2.22	0.40
31:BA:978:A:C6	31:BA:1318:A:C5	3.09	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:BD:26:A:N6	52:BD:44:G:H1	2.18	0.40
32:BE:19:HIS:O	32:BE:20:GLU:O	2.39	0.40
32:BE:82:ARG:O	32:BE:86:GLU:HG3	2.21	0.40
33:BF:141:VAL:HG12	33:BF:141:VAL:O	2.21	0.40
34:BG:107:ARG:HB3	34:BG:174:LEU:HD13	2.03	0.40
36:BI:15:ASP:OD2	36:BI:18:GLN:HG3	2.22	0.40
38:BK:82:HIS:HB3	38:BK:138:TRP:CE2	2.56	0.40
31:BA:1199:U:C4'	40:BM:54:PHE:CE1	3.04	0.40
41:BN:91:ARG:HG2	41:BN:91:ARG:NH1	2.33	0.40
42:BO:104:VAL:O	42:BO:105:TYR:HB2	2.20	0.40
31:BA:363:A:C8	42:BO:33:ARG:NH2	2.89	0.40
43:BP:80:ARG:CZ	43:BP:80:ARG:HB3	2.51	0.40
43:BP:88:ARG:NH1	43:BP:88:ARG:HG2	2.37	0.40
48:BU:81:PHE:C	48:BU:82:THR:HG1	2.22	0.40
51:BX:24:ARG:O	51:BX:25:LYS:O	2.40	0.40
54:CA:1085:U:H4'	54:CA:1086:U:OP2	2.19	0.40
54:CA:1156:G:H3'	54:CA:1157:A:C5'	2.51	0.40
54:CA:1298:C:H4'	54:CA:1299:A:N9	2.37	0.40
54:CA:949:A:H1'	54:CA:1364:U:H3	1.87	0.40
54:CA:191:G:O2'	54:CA:192:U:H5'	2.21	0.40
54:CA:329:A:H4'	54:CA:330:C:OP1	2.17	0.40
54:CA:342:C:C4	54:CA:343:U:C4	3.10	0.40
54:CA:730:G:C5	54:CA:731:G:H1'	2.56	0.40
54:CA:817:C:C4	54:CA:819:A:H1'	2.56	0.40
54:CA:831:U:H2'	54:CA:832:C:C6	2.56	0.40
54:CA:842:C:H5'	54:CA:843:U:OP1	2.22	0.40
32:CE:18:GLY:CA	32:CE:42:ILE:HG22	2.52	0.40
32:CE:52:GLU:CG	32:CE:56:ARG:HH12	2.27	0.40
32:CE:79:ASP:O	32:CE:83:MET:HG2	2.22	0.40
33:CF:122:GLU:O	33:CF:126:ARG:HG3	2.21	0.40
33:CF:153:VAL:HG13	33:CF:196:LEU:CD1	2.51	0.40
34:CG:206:PHE:C	34:CG:208:SER:H	2.25	0.40
34:CG:53:ASP:OD2	34:CG:57:ARG:NH1	2.49	0.40
34:CG:96:LEU:HD22	34:CG:96:LEU:H	1.86	0.40
36:CI:17:SER:O	36:CI:18:GLN:C	2.60	0.40
37:CJ:132:GLY:CA	37:CJ:135:VAL:HG23	2.52	0.40
37:CJ:78:ARG:HD3	37:CJ:156:TRP:HB3	2.04	0.40
39:CL:40:LEU:CD1	39:CL:70:LYS:HG2	2.51	0.40
39:CL:61:ALA:HB1	39:CL:63:ILE:HD11	2.04	0.40
40:CM:53:PRO:HA	44:CQ:42:ILE:HD11	2.03	0.40
41:CN:128:ALA:HB1	53:C1:39:U:OP1	2.22	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:CO:117:ARG:HB3	42:CO:122:THR:O	2.21	0.40
54:CA:948:C:C4	43:CP:106:ASN:ND2	2.84	0.40
43:CP:108:ARG:O	43:CP:109:THR:C	2.59	0.40
43:CP:26:GLY:O	43:CP:27:LYS:C	2.60	0.40
33:CF:29:TYR:CD1	44:CQ:36:PHE:HE1	2.40	0.40
45:CR:10:LYS:O	45:CR:14:GLU:HB2	2.22	0.40
46:CS:20:VAL:CG2	46:CS:32:TYR:HB2	2.52	0.40
46:CS:40:ASP:OD2	46:CS:42:ARG:CB	2.68	0.40
46:CS:59:TRP:HA	46:CS:62:VAL:HG22	2.04	0.40
47:CT:40:LYS:HD2	47:CT:42:TYR:CZ	2.56	0.40
26:D4:33:VAL:O	26:D4:34:GLU:C	2.60	0.40
30:D8:23:VAL:HG11	30:D8:46:ARG:HB3	2.00	0.40
30:D8:62:LEU:HA	30:D8:62:LEU:HD23	2.00	0.40
55:DA:1058:U:H5''	58:DL:4:VAL:HG12	2.04	0.40
55:DA:1240:U:O2'	55:DA:1241:A:C5'	2.67	0.40
55:DA:1472:A:H2'	55:DA:1473:G:C8	2.55	0.40
55:DA:1483:G:C2	55:DA:1484:G:C5	3.10	0.40
55:DA:1534:G:C8	55:DA:1534:G:C5'	3.05	0.40
55:DA:2097:C:H2'	55:DA:2098:U:O4'	2.22	0.40
55:DA:2104:G:O2'	55:DA:2105:C:H5'	2.22	0.40
55:DA:2373:G:H2'	55:DA:2374:C:C6	2.56	0.40
55:DA:2547:U:H2'	55:DA:2548:G:C8	2.57	0.40
55:DA:2628:C:H1'	55:DA:2781:A:C4	2.56	0.40
55:DA:2656:U:C3'	55:DA:2656:U:C6	3.04	0.40
55:DA:2770:G:H5''	55:DA:2771:C:OP2	2.22	0.40
55:DA:2870:C:H5'	13:D0:61:HIS:CE1	2.54	0.40
55:DA:2895:U:H2'	55:DA:2896:C:O4'	2.22	0.40
55:DA:547:A:H2'	55:DA:548:A:N9	2.36	0.40
55:DA:621:A:C2	55:DA:622:G:C5	3.09	0.40
55:DA:658:C:C2	55:DA:659:C:C5	3.09	0.40
55:DA:78:A:H2'	55:DA:79:G:H8	1.87	0.40
55:DA:851:U:O4'	25:DX:46:ASN:ND2	2.55	0.40
2:DB:74:U:C2'	2:DB:75:G:C5'	2.82	0.40
3:DD:10:THR:CG2	3:DD:13:ARG:HB2	2.50	0.40
3:DD:222:ARG:NH1	3:DD:224:ALA:HB3	2.37	0.40
3:DD:48:ARG:NH1	3:DD:48:ARG:HG3	2.37	0.40
4:DE:27:LEU:HD21	15:DR:1:MET:HE2	2.02	0.40
55:DA:2784:C:H4'	4:DE:41:LYS:O	2.22	0.40
6:DG:53:LEU:HD23	6:DG:54:GLU:CA	2.52	0.40
56:DI:17:VAL:CB	56:DI:21:LYS:HE3	2.52	0.40
56:DI:7:ARG:O	56:DI:7:ARG:HG2	2.20	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:DL:136:VAL:CG2	58:DL:137:GLU:N	2.83	0.40
58:DL:34:ILE:O	58:DL:34:ILE:CG2	2.65	0.40
58:DL:52:ILE:HD11	58:DL:73:PRO:HA	2.04	0.40
9:DM:39:ARG:HH12	9:DM:41:ASP:CG	2.25	0.40
9:DM:6:PRO:C	9:DM:7:LYS:HZ2	2.23	0.40
10:DN:36:GLY:HA2	10:DN:106:LEU:CD2	2.51	0.40
10:DN:10:VAL:HG23	10:DN:17:ARG:C	2.42	0.40
10:DN:77:ILE:HG12	15:DR:74:ARG:HG3	2.04	0.40
11:DO:126:VAL:HG13	11:DO:145:PRO:CB	2.52	0.40
11:DO:113:LYS:HA	11:DO:129:ALA:O	2.22	0.40
55:DA:2404:C:O3'	11:DO:77:ARG:NH2	2.54	0.40
14:DQ:20:ARG:C	14:DQ:22:GLY:N	2.74	0.40
25:DX:7:LYS:HG3	25:DX:34:GLU:HG2	2.02	0.40
57:DY:127:GLU:HA	57:DY:127:GLU:OE2	2.21	0.40
57:DY:27:VAL:O	57:DY:28:ASN:CB	2.54	0.40
57:DY:4:LYS:O	57:DY:5:ARG:HG3	2.22	0.40
57:DY:51:LEU:HD22	57:DY:82:PHE:N	2.26	0.40
57:DY:9:LEU:HD13	57:DY:10:LEU:CA	2.46	0.40
23:DZ:47:GLN:HA	23:DZ:47:GLN:OE1	2.21	0.40

All (2) symmetry-related close contacts are listed below. The label for Atom-2 includes the symmetry operator and encoded unit-cell translations to be applied.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:AA:363:G:O6	31:BA:86:U:OP1[3_555]	2.10	0.10
16:A1:84:LYS:NZ	55:DA:654(I):C:O2'[2_465]	2.12	0.08

5.3 Torsion angles ⓘ

5.3.1 Protein backbone ⓘ

In the following table, the Percentiles column shows the percent Ramachandran outliers of the chain as a percentile score with respect to all X-ray entries followed by that with respect to entries of similar resolution.

The Analysed column shows the number of residues for which the backbone conformation was analysed, and the total number of residues.

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles
3	AD	270/276 (98%)	201 (74%)	45 (17%)	24 (9%)	1 4

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
3	DD	270/276 (98%)	193 (72%)	61 (23%)	16 (6%)	1	10
4	AE	203/206 (98%)	105 (52%)	48 (24%)	50 (25%)	0	0
4	DE	203/206 (98%)	120 (59%)	36 (18%)	47 (23%)	0	0
5	AF	206/210 (98%)	136 (66%)	41 (20%)	29 (14%)	0	1
5	DF	200/210 (95%)	157 (78%)	29 (14%)	14 (7%)	1	7
6	AG	179/182 (98%)	127 (71%)	36 (20%)	16 (9%)	1	4
6	DG	179/182 (98%)	124 (69%)	30 (17%)	25 (14%)	0	1
7	AH	168/180 (93%)	80 (48%)	51 (30%)	37 (22%)	0	0
7	DH	168/180 (93%)	94 (56%)	43 (26%)	31 (18%)	0	0
8	AK	144/148 (97%)	83 (58%)	34 (24%)	27 (19%)	0	0
8	DK	144/148 (97%)	76 (53%)	41 (28%)	27 (19%)	0	0
9	AM	136/140 (97%)	95 (70%)	25 (18%)	16 (12%)	0	1
9	DM	136/140 (97%)	106 (78%)	16 (12%)	14 (10%)	0	3
10	AN	120/122 (98%)	98 (82%)	17 (14%)	5 (4%)	3	16
10	DN	120/122 (98%)	104 (87%)	12 (10%)	4 (3%)	4	21
11	AO	148/150 (99%)	73 (49%)	30 (20%)	45 (30%)	0	0
11	DO	148/150 (99%)	88 (60%)	27 (18%)	33 (22%)	0	0
12	AP	139/141 (99%)	73 (52%)	39 (28%)	27 (19%)	0	0
12	DP	139/141 (99%)	91 (66%)	28 (20%)	20 (14%)	0	1
13	A0	115/118 (98%)	84 (73%)	20 (17%)	11 (10%)	0	3
13	D0	116/118 (98%)	87 (75%)	17 (15%)	12 (10%)	0	3
14	AQ	109/112 (97%)	68 (62%)	21 (19%)	20 (18%)	0	0
14	DQ	109/112 (97%)	70 (64%)	21 (19%)	18 (16%)	0	0
15	AR	135/146 (92%)	98 (73%)	25 (18%)	12 (9%)	1	4
15	DR	135/146 (92%)	92 (68%)	27 (20%)	16 (12%)	0	1
16	A1	115/118 (98%)	79 (69%)	27 (24%)	9 (8%)	1	5
16	D1	115/118 (98%)	96 (84%)	11 (10%)	8 (7%)	1	7
17	A2	99/101 (98%)	60 (61%)	17 (17%)	22 (22%)	0	0
17	D2	99/101 (98%)	80 (81%)	9 (9%)	10 (10%)	0	3
18	AS	111/113 (98%)	82 (74%)	21 (19%)	8 (7%)	1	6
18	DS	111/113 (98%)	87 (78%)	20 (18%)	4 (4%)	3	20

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
19	AT	90/96 (94%)	66 (73%)	17 (19%)	7 (8%)	1	5
19	DT	90/96 (94%)	75 (83%)	13 (14%)	2 (2%)	6	29
20	AU	100/110 (91%)	48 (48%)	22 (22%)	30 (30%)	0	0
20	DU	100/110 (91%)	64 (64%)	11 (11%)	25 (25%)	0	0
21	AV	185/206 (90%)	90 (49%)	45 (24%)	50 (27%)	0	0
21	DV	198/206 (96%)	102 (52%)	46 (23%)	50 (25%)	0	0
22	A3	82/85 (96%)	56 (68%)	16 (20%)	10 (12%)	0	1
22	D3	82/85 (96%)	62 (76%)	14 (17%)	6 (7%)	1	6
23	AZ	95/98 (97%)	73 (77%)	10 (10%)	12 (13%)	0	1
23	DZ	95/98 (97%)	74 (78%)	13 (14%)	8 (8%)	1	5
24	AW	67/72 (93%)	42 (63%)	17 (25%)	8 (12%)	0	1
24	DW	67/72 (93%)	52 (78%)	8 (12%)	7 (10%)	0	3
25	AX	57/60 (95%)	46 (81%)	6 (10%)	5 (9%)	1	4
25	DX	57/60 (95%)	49 (86%)	5 (9%)	3 (5%)	2	12
26	A4	69/71 (97%)	26 (38%)	14 (20%)	29 (42%)	0	0
26	D4	69/71 (97%)	35 (51%)	12 (17%)	22 (32%)	0	0
27	A5	57/60 (95%)	40 (70%)	9 (16%)	8 (14%)	0	1
27	D5	57/60 (95%)	35 (61%)	9 (16%)	13 (23%)	0	0
28	A6	43/54 (80%)	15 (35%)	12 (28%)	16 (37%)	0	0
28	D6	43/54 (80%)	19 (44%)	9 (21%)	15 (35%)	0	0
29	A7	47/49 (96%)	41 (87%)	6 (13%)	0	100	100
29	D7	47/49 (96%)	40 (85%)	5 (11%)	2 (4%)	2	16
30	A8	62/65 (95%)	37 (60%)	11 (18%)	14 (23%)	0	0
30	D8	62/65 (95%)	43 (69%)	9 (14%)	10 (16%)	0	0
32	BE	235/256 (92%)	137 (58%)	62 (26%)	36 (15%)	0	0
32	CE	235/256 (92%)	140 (60%)	58 (25%)	37 (16%)	0	0
33	BF	204/239 (85%)	135 (66%)	40 (20%)	29 (14%)	0	1
33	CF	203/239 (85%)	137 (68%)	49 (24%)	17 (8%)	1	5
34	BG	206/209 (99%)	134 (65%)	43 (21%)	29 (14%)	0	1
34	CG	206/209 (99%)	124 (60%)	59 (29%)	23 (11%)	0	2
35	BH	149/162 (92%)	115 (77%)	22 (15%)	12 (8%)	1	5

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
35	CH	149/162 (92%)	115 (77%)	26 (17%)	8 (5%)	2	12
36	BI	99/101 (98%)	79 (80%)	15 (15%)	5 (5%)	2	13
36	CI	99/101 (98%)	66 (67%)	26 (26%)	7 (7%)	1	6
37	BJ	153/156 (98%)	115 (75%)	26 (17%)	12 (8%)	1	5
37	CJ	153/156 (98%)	120 (78%)	26 (17%)	7 (5%)	2	15
38	BK	136/138 (99%)	104 (76%)	21 (15%)	11 (8%)	1	5
38	CK	136/138 (99%)	101 (74%)	30 (22%)	5 (4%)	3	19
39	BL	125/128 (98%)	89 (71%)	28 (22%)	8 (6%)	1	8
39	CL	125/128 (98%)	93 (74%)	19 (15%)	13 (10%)	0	3
40	BM	97/105 (92%)	61 (63%)	24 (25%)	12 (12%)	0	1
40	CM	97/105 (92%)	71 (73%)	19 (20%)	7 (7%)	1	6
41	BN	117/129 (91%)	87 (74%)	21 (18%)	9 (8%)	1	5
41	CN	117/129 (91%)	90 (77%)	24 (20%)	3 (3%)	5	26
42	BO	123/132 (93%)	81 (66%)	19 (15%)	23 (19%)	0	0
42	CO	123/132 (93%)	85 (69%)	17 (14%)	21 (17%)	0	0
43	BP	119/126 (94%)	80 (67%)	23 (19%)	16 (13%)	0	1
43	CP	123/126 (98%)	77 (63%)	22 (18%)	24 (20%)	0	0
44	BQ	58/61 (95%)	39 (67%)	12 (21%)	7 (12%)	0	1
44	CQ	58/61 (95%)	37 (64%)	12 (21%)	9 (16%)	0	0
45	BR	86/89 (97%)	67 (78%)	16 (19%)	3 (4%)	3	20
45	CR	86/89 (97%)	66 (77%)	14 (16%)	6 (7%)	1	7
46	BS	82/88 (93%)	57 (70%)	19 (23%)	6 (7%)	1	6
46	CS	82/88 (93%)	54 (66%)	22 (27%)	6 (7%)	1	6
47	BT	98/105 (93%)	78 (80%)	11 (11%)	9 (9%)	1	4
47	CT	98/105 (93%)	75 (76%)	17 (17%)	6 (6%)	1	9
48	BU	70/88 (80%)	45 (64%)	16 (23%)	9 (13%)	0	1
48	CU	70/88 (80%)	51 (73%)	11 (16%)	8 (11%)	0	2
49	BV	81/93 (87%)	47 (58%)	23 (28%)	11 (14%)	0	1
49	CV	86/93 (92%)	56 (65%)	16 (19%)	14 (16%)	0	0
50	BW	97/106 (92%)	55 (57%)	29 (30%)	13 (13%)	0	1
50	CW	97/106 (92%)	65 (67%)	20 (21%)	12 (12%)	0	1

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
51	BX	23/27 (85%)	16 (70%)	4 (17%)	3 (13%)	0	1
51	CX	23/27 (85%)	18 (78%)	3 (13%)	2 (9%)	1	4
56	DI	28/125 (22%)	6 (21%)	6 (21%)	16 (57%)	0	0
56	DJ	28/125 (22%)	10 (36%)	8 (29%)	10 (36%)	0	0
57	DY	143/173 (83%)	30 (21%)	36 (25%)	77 (54%)	0	0
58	DL	143/147 (97%)	46 (32%)	29 (20%)	68 (48%)	0	0
All	All	11776/12624 (93%)	7791 (66%)	2307 (20%)	1678 (14%)	0	1

All (1678) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
3	AD	3	VAL
3	AD	25	THR
3	AD	26	LYS
3	AD	28	GLU
3	AD	33	LEU
3	AD	58	HIS
3	AD	157	ARG
3	AD	196	VAL
3	AD	225	ALA
3	AD	237	GLU
3	AD	267	SER
3	AD	268	ARG
4	AE	2	LYS
4	AE	4	ILE
4	AE	9	VAL
4	AE	11	MET
4	AE	23	VAL
4	AE	32	PRO
4	AE	33	VAL
4	AE	53	PRO
4	AE	59	VAL
4	AE	60	ASN
4	AE	61	ARG
4	AE	71	GLY
4	AE	72	VAL
4	AE	77	ILE
4	AE	78	LEU
4	AE	88	GLY
4	AE	92	THR

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Mol	Chain	Res	Type
4	AE	118	LYS
4	AE	131	ALA
4	AE	132	HIS
4	AE	133	LYS
5	AF	2	LYS
5	AF	21	ALA
5	AF	25	PRO
5	AF	59	TYR
5	AF	61	GLY
5	AF	62	ARG
5	AF	89	VAL
5	AF	123	LEU
5	AF	129	PHE
5	AF	176	LEU
6	AG	14	GLU
6	AG	96	ARG
6	AG	117	PHE
7	AH	23	ARG
7	AH	50	VAL
7	AH	81	GLU
7	AH	87	LEU
7	AH	92	ILE
7	AH	126	PRO
7	AH	127	GLU
7	AH	130	ARG
7	AH	152	ARG
7	AH	153	LYS
7	AH	167	GLU
8	AK	30	LEU
8	AK	77	LEU
8	AK	78	THR
8	AK	88	ILE
8	AK	100	ALA
8	AK	102	SER
8	AK	117	GLU
8	AK	131	LYS
8	AK	142	VAL
8	AK	144	VAL
9	AM	17	ASP
9	AM	18	ALA
9	AM	50	ASP
9	AM	56	ASN

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Mol	Chain	Res	Type
9	AM	63	THR
9	AM	64	GLY
9	AM	130	HIS
9	AM	133	GLN
10	AN	48	PRO
11	AO	10	PRO
11	AO	15	ARG
11	AO	17	LYS
11	AO	19	VAL
11	AO	25	SER
11	AO	34	GLY
11	AO	35	HIS
11	AO	38	GLN
11	AO	49	ARG
11	AO	56	SER
11	AO	57	THR
11	AO	58	THR
11	AO	60	MET
11	AO	62	LEU
11	AO	64	LYS
11	AO	66	GLY
11	AO	94	GLU
11	AO	108	LYS
11	AO	110	TYR
11	AO	111	ARG
11	AO	147	LEU
12	AP	2	LEU
12	AP	11	LYS
12	AP	25	ASP
12	AP	61	GLY
12	AP	79	LEU
12	AP	90	VAL
12	AP	105	GLU
12	AP	130	LYS
12	AP	134	ARG
12	AP	138	ASP
12	AP	139	GLU
13	A0	10	LEU
13	A0	82	GLU
14	AQ	12	PHE
14	AQ	14	VAL
14	AQ	55	ALA

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Mol	Chain	Res	Type
14	AQ	56	LEU
14	AQ	86	ALA
14	AQ	87	PHE
14	AQ	88	ASP
14	AQ	89	ARG
14	AQ	107	GLU
15	AR	3	ARG
15	AR	78	LEU
15	AR	84	GLN
15	AR	86	ILE
15	AR	107	ASP
16	A1	90	VAL
16	A1	91	ASP
16	A1	98	LEU
17	A2	44	LYS
17	A2	45	THR
17	A2	47	VAL
17	A2	49	THR
17	A2	71	LEU
17	A2	72	VAL
17	A2	79	VAL
17	A2	84	LYS
17	A2	85	LYS
18	AS	45	TYR
18	AS	63	ASP
18	AS	80	PRO
19	AT	68	ARG
20	AU	17	SER
20	AU	21	LYS
20	AU	29	GLU
20	AU	49	VAL
20	AU	50	ARG
20	AU	63	LYS
20	AU	72	VAL
20	AU	77	PRO
20	AU	78	ALA
20	AU	85	VAL
20	AU	89	PHE
20	AU	90	LEU
20	AU	102	CYS
21	AV	6	LYS
21	AV	31	ARG

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Mol	Chain	Res	Type
21	AV	34	ASN
21	AV	61	LEU
21	AV	62	PRO
21	AV	65	GLN
21	AV	93	ASP
21	AV	96	VAL
21	AV	107	THR
21	AV	108	PRO
21	AV	112	ARG
21	AV	115	GLY
21	AV	117	LEU
21	AV	141	VAL
21	AV	148	ASP
21	AV	149	SER
21	AV	155	LEU
21	AV	171	ILE
21	AV	177	PRO
21	AV	179	ASP
21	AV	184	ALA
21	AV	186	GLU
22	A3	5	LYS
22	A3	7	LEU
22	A3	15	ASP
22	A3	84	LEU
23	AZ	30	VAL
23	AZ	82	LEU
23	AZ	84	GLY
23	AZ	93	GLU
24	AW	16	LEU
24	AW	17	SER
24	AW	43	GLN
24	AW	44	LEU
24	AW	47	ASN
24	AW	48	HIS
24	AW	70	GLN
26	A4	8	LYS
26	A4	12	ALA
26	A4	22	ILE
26	A4	26	SER
26	A4	34	GLU
26	A4	39	CYS
26	A4	40	HIS

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Mol	Chain	Res	Type
26	A4	42	PHE
26	A4	51	ASP
26	A4	53	GLU
26	A4	56	VAL
26	A4	66	SER
26	A4	67	TYR
26	A4	68	ARG
27	A5	4	HIS
27	A5	53	ALA
27	A5	58	LEU
28	A6	16	CYS
28	A6	19	ARG
28	A6	23	THR
28	A6	44	ARG
30	A8	31	HIS
30	A8	32	LEU
30	A8	34	TRP
30	A8	41	ILE
30	A8	48	PHE
30	A8	49	VAL
30	A8	50	LEU
30	A8	51	ALA
30	A8	61	LEU
30	A8	62	LEU
30	A8	64	TYR
32	BE	7	VAL
32	BE	39	ILE
32	BE	74	LYS
32	BE	96	ARG
32	BE	191	ASP
32	BE	194	PRO
32	BE	195	ASP
32	BE	229	VAL
32	BE	232	PRO
32	BE	238	LEU
33	BF	4	LYS
33	BF	12	LEU
33	BF	18	TRP
33	BF	45	LYS
33	BF	61	ALA
33	BF	98	ASN
33	BF	101	LEU

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Mol	Chain	Res	Type
33	BF	127	ARG
33	BF	146	ALA
33	BF	155	GLY
33	BF	165	THR
34	BG	14	ARG
34	BG	17	VAL
34	BG	20	TYR
34	BG	24	GLU
34	BG	27	TYR
34	BG	40	PRO
34	BG	150	GLU
34	BG	151	LYS
34	BG	189	PRO
34	BG	190	ASP
35	BH	146	ALA
36	BI	15	ASP
38	BK	91	ARG
38	BK	101	PRO
38	BK	103	VAL
38	BK	105	ARG
39	BL	40	LEU
40	BM	57	LYS
41	BN	106	LYS
42	BO	18	VAL
42	BO	19	ARG
42	BO	47	LYS
42	BO	63	GLY
42	BO	64	TYR
42	BO	65	GLU
42	BO	79	GLU
43	BP	4	ILE
43	BP	12	ASN
43	BP	25	ILE
43	BP	116	THR
44	BQ	14	PRO
44	BQ	16	PHE
44	BQ	23	ARG
44	BQ	29	ARG
47	BT	69	LYS
48	BU	19	LYS
48	BU	20	ALA
48	BU	21	LYS

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Mol	Chain	Res	Type
49	BV	9	VAL
49	BV	24	ALA
49	BV	30	LEU
49	BV	42	PRO
49	BV	43	GLU
49	BV	48	THR
50	BW	11	SER
50	BW	49	ALA
50	BW	71	THR
51	BX	25	LYS
32	CE	6	THR
32	CE	15	VAL
32	CE	96	ARG
32	CE	122	PHE
32	CE	126	GLU
32	CE	195	ASP
32	CE	208	ILE
32	CE	230	VAL
32	CE	235	SER
32	CE	237	ALA
33	CF	12	LEU
33	CF	79	ARG
33	CF	189	ALA
34	CG	7	PRO
34	CG	28	SER
34	CG	30	LYS
34	CG	151	LYS
34	CG	178	VAL
35	CH	21	ALA
35	CH	77	PRO
35	CH	115	VAL
35	CH	146	ALA
36	CI	42	GLU
37	CJ	53	LYS
38	CK	103	VAL
39	CL	23	ASN
39	CL	41	VAL
39	CL	55	ALA
39	CL	95	LYS
40	CM	57	LYS
41	CN	82	VAL
42	CO	18	VAL

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Mol	Chain	Res	Type
42	CO	27	LEU
42	CO	46	LYS
42	CO	47	LYS
42	CO	48	PRO
42	CO	62	SER
42	CO	64	TYR
42	CO	65	GLU
42	CO	79	GLU
42	CO	91	LYS
43	CP	4	ILE
43	CP	12	ASN
43	CP	28	ALA
43	CP	60	VAL
43	CP	67	GLU
43	CP	101	GLN
43	CP	116	THR
43	CP	118	ALA
43	CP	122	LYS
44	CQ	14	PRO
44	CQ	16	PHE
44	CQ	23	ARG
44	CQ	36	PHE
46	CS	49	LEU
47	CT	78	GLU
47	CT	81	ARG
48	CU	21	LYS
49	CV	3	ARG
49	CV	5	LEU
49	CV	45	VAL
49	CV	82	GLY
49	CV	83	HIS
50	CW	49	ALA
50	CW	95	ALA
50	CW	100	ILE
3	DD	28	GLU
3	DD	33	LEU
3	DD	122	ASP
3	DD	123	ALA
4	DE	2	LYS
4	DE	15	PHE
4	DE	19	ARG
4	DE	21	VAL

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Mol	Chain	Res	Type
4	DE	22	PRO
4	DE	35	GLN
4	DE	37	ARG
4	DE	54	GLN
4	DE	61	ARG
4	DE	62	PRO
4	DE	64	LYS
4	DE	68	ALA
4	DE	72	VAL
4	DE	79	ARG
4	DE	88	GLY
4	DE	118	LYS
4	DE	132	HIS
4	DE	154	LYS
4	DE	169	ASN
4	DE	187	ALA
5	DF	24	LEU
5	DF	66	PRO
5	DF	73	ALA
5	DF	89	VAL
5	DF	128	ALA
5	DF	134	GLY
6	DG	14	GLU
6	DG	96	ARG
7	DH	10	PRO
7	DH	12	PRO
7	DH	16	SER
7	DH	17	VAL
7	DH	20	ALA
7	DH	27	LYS
7	DH	83	TYR
7	DH	86	GLU
7	DH	92	ILE
7	DH	126	PRO
7	DH	127	GLU
7	DH	138	LYS
7	DH	153	LYS
7	DH	154	PRO
7	DH	155	SER
7	DH	156	ALA
7	DH	167	GLU
7	DH	169	VAL

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Mol	Chain	Res	Type
56	DI	3	LEU
56	DI	6	GLU
56	DI	10	GLU
56	DI	13	SER
56	DI	20	LEU
56	DI	21	LYS
56	DI	24	ILE
8	DK	13	GLY
8	DK	15	VAL
8	DK	77	LEU
8	DK	78	THR
8	DK	79	ILE
8	DK	115	ALA
8	DK	134	PRO
57	DY	2	PRO
57	DY	3	ASN
57	DY	4	LYS
57	DY	6	ASN
57	DY	8	GLU
57	DY	9	LEU
57	DY	11	ALA
57	DY	23	SER
57	DY	25	PHE
57	DY	28	ASN
57	DY	32	LEU
57	DY	43	ALA
57	DY	44	LEU
57	DY	47	ASN
57	DY	50	ARG
57	DY	53	VAL
57	DY	57	THR
57	DY	58	LEU
57	DY	62	ALA
57	DY	63	LEU
57	DY	64	LYS
57	DY	71	LEU
57	DY	74	LEU
57	DY	78	SER
57	DY	83	TYR
57	DY	88	ALA
57	DY	97	ALA
57	DY	102	LYS

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Mol	Chain	Res	Type
57	DY	104	ILE
57	DY	106	GLN
57	DY	108	LYS
57	DY	113	GLN
57	DY	117	LEU
57	DY	120	LYS
57	DY	121	ASP
57	DY	122	VAL
57	DY	123	GLU
57	DY	127	GLU
57	DY	129	PRO
57	DY	130	THR
57	DY	131	MET
57	DY	132	ASP
57	DY	133	GLU
57	DY	142	LEU
57	DY	143	GLN
58	DL	4	VAL
58	DL	7	VAL
58	DL	12	LEU
58	DL	14	ALA
58	DL	18	THR
58	DL	20	ALA
58	DL	33	ASN
58	DL	39	LYS
58	DL	48	MET
58	DL	49	GLY
58	DL	50	ASP
58	DL	52	ILE
58	DL	53	VAL
58	DL	54	PRO
58	DL	57	ILE
58	DL	60	TYR
58	DL	63	ARG
58	DL	64	SER
58	DL	66	THR
58	DL	68	VAL
58	DL	70	LYS
58	DL	72	PRO
58	DL	75	SER
58	DL	81	ALA
58	DL	82	ALA

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Mol	Chain	Res	Type
58	DL	87	GLY
58	DL	90	LYS
58	DL	102	GLU
58	DL	104	VAL
58	DL	113	PRO
58	DL	114	ASP
58	DL	116	ASN
58	DL	121	GLU
58	DL	126	MET
58	DL	136	VAL
58	DL	139	VAL
58	DL	141	ALA
58	DL	143	GLU
58	DL	146	ASP
56	DJ	4	ASP
56	DJ	8	ILE
56	DJ	18	LEU
56	DJ	19	GLU
56	DJ	29	GLU
9	DM	9	VAL
9	DM	22	THR
9	DM	58	ASP
9	DM	134	ARG
10	DN	97	ARG
11	DO	6	LEU
11	DO	10	PRO
11	DO	15	ARG
11	DO	16	ARG
11	DO	19	VAL
11	DO	25	SER
11	DO	27	HIS
11	DO	31	ALA
11	DO	36	LYS
11	DO	38	GLN
11	DO	57	THR
11	DO	67	MET
11	DO	106	LEU
11	DO	149	GLU
12	DP	19	GLY
12	DP	22	LYS
12	DP	88	GLY
12	DP	135	ASP

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Mol	Chain	Res	Type
12	DP	139	GLU
13	D0	3	HIS
13	D0	4	LEU
13	D0	58	GLY
14	DQ	11	LYS
14	DQ	14	VAL
14	DQ	19	LYS
14	DQ	61	ASN
14	DQ	88	ASP
14	DQ	89	ARG
14	DQ	107	GLU
15	DR	2	ASN
15	DR	3	ARG
15	DR	42	ILE
15	DR	58	ASN
15	DR	105	LEU
15	DR	123	GLN
15	DR	124	ASP
16	D1	90	VAL
16	D1	91	ASP
16	D1	93	LYS
16	D1	117	GLN
17	D2	45	THR
17	D2	49	THR
17	D2	79	VAL
18	DS	67	ASP
19	DT	68	ARG
20	DU	5	MET
20	DU	11	ASP
20	DU	20	TYR
20	DU	45	VAL
20	DU	49	VAL
20	DU	50	ARG
20	DU	57	GLN
20	DU	63	LYS
20	DU	77	PRO
20	DU	78	ALA
20	DU	80	GLY
21	DV	7	ALA
21	DV	107	THR
21	DV	112	ARG
21	DV	118	GLN

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Mol	Chain	Res	Type
21	DV	124	ILE
21	DV	140	ASP
21	DV	143	GLY
21	DV	146	ILE
21	DV	153	SER
21	DV	154	ASP
21	DV	155	LEU
21	DV	161	VAL
21	DV	175	VAL
21	DV	180	VAL
21	DV	184	ALA
21	DV	185	GLU
21	DV	187	ALA
21	DV	191	VAL
21	DV	192	ALA
21	DV	194	PRO
21	DV	197	ILE
21	DV	198	LYS
21	DV	199	LYS
22	D3	4	LYS
22	D3	9	SER
24	DW	16	LEU
24	DW	43	GLN
24	DW	47	ASN
24	DW	48	HIS
24	DW	70	GLN
26	D4	14	ILE
26	D4	31	ILE
26	D4	38	LYS
26	D4	39	CYS
26	D4	40	HIS
26	D4	46	GLN
26	D4	50	VAL
26	D4	51	ASP
26	D4	53	GLU
26	D4	55	ARG
26	D4	56	VAL
26	D4	57	GLU
27	D5	4	HIS
27	D5	35	GLU
27	D5	43	HIS
27	D5	47	PRO

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Mol	Chain	Res	Type
27	D5	53	ALA
28	D6	15	GLU
28	D6	19	ARG
28	D6	21	TYR
28	D6	31	PRO
28	D6	32	ASN
28	D6	43	CYS
28	D6	45	LYS
28	D6	46	HIS
29	D7	48	LYS
30	D8	7	HIS
30	D8	31	HIS
30	D8	52	LYS
30	D8	62	LEU
3	AD	144	ALA
3	AD	156	ALA
3	AD	238	GLY
4	AE	25	VAL
4	AE	50	GLY
4	AE	51	PHE
4	AE	54	GLN
4	AE	58	ARG
4	AE	64	LYS
4	AE	94	GLU
4	AE	130	GLY
4	AE	152	LYS
4	AE	187	ALA
4	AE	200	GLU
4	AE	204	ALA
5	AF	3	GLU
5	AF	5	ALA
5	AF	11	VAL
5	AF	17	ARG
5	AF	19	GLU
5	AF	66	PRO
5	AF	84	VAL
5	AF	125	LEU
5	AF	127	GLU
5	AF	128	ALA
5	AF	132	VAL
6	AG	3	LEU
6	AG	5	VAL

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Mol	Chain	Res	Type
7	AH	3	ARG
7	AH	12	PRO
7	AH	27	LYS
7	AH	83	TYR
7	AH	84	SER
7	AH	141	VAL
7	AH	155	SER
7	AH	168	PRO
7	AH	169	VAL
8	AK	12	LEU
8	AK	73	GLU
8	AK	101	LEU
8	AK	119	PRO
8	AK	140	LEU
8	AK	143	SER
8	AK	145	VAL
9	AM	2	LYS
10	AN	5	GLN
11	AO	5	ASP
11	AO	6	LEU
11	AO	16	ARG
11	AO	29	LYS
11	AO	36	LYS
11	AO	47	ASP
11	AO	90	ARG
11	AO	116	GLY
11	AO	119	GLU
11	AO	141	ALA
11	AO	146	VAL
12	AP	15	GLY
12	AP	22	LYS
12	AP	23	GLY
12	AP	88	GLY
12	AP	140	ALA
13	A0	11	ASN
13	A0	14	SER
13	A0	42	LYS
13	A0	93	GLY
13	A0	102	GLU
14	AQ	4	LEU
14	AQ	11	LYS
14	AQ	57	LYS

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Mol	Chain	Res	Type
14	AQ	96	GLY
14	AQ	109	GLY
15	AR	2	ASN
15	AR	97	ALA
15	AR	104	ASN
15	AR	115	ARG
16	A1	9	VAL
17	A2	38	LEU
17	A2	67	GLY
17	A2	80	GLN
18	AS	65	LEU
18	AS	67	ASP
18	AS	110	LYS
20	AU	3	VAL
20	AU	39	VAL
20	AU	44	ILE
20	AU	57	GLN
20	AU	59	GLY
20	AU	73	ARG
20	AU	99	CYS
21	AV	12	GLY
21	AV	13	GLU
21	AV	38	TYR
21	AV	53	ILE
21	AV	63	ASP
21	AV	64	GLY
21	AV	78	LYS
21	AV	105	VAL
21	AV	140	ASP
21	AV	142	SER
21	AV	143	GLY
21	AV	145	GLU
21	AV	146	ILE
21	AV	180	VAL
21	AV	181	GLU
21	AV	185	GLU
22	A3	3	HIS
22	A3	4	LYS
22	A3	55	ARG
22	A3	73	GLY
23	AZ	28	GLY
23	AZ	31	GLY

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Mol	Chain	Res	Type
23	AZ	92	LYS
23	AZ	96	LYS
25	AX	13	ILE
25	AX	26	LEU
26	A4	36	CYS
26	A4	38	LYS
26	A4	46	GLN
26	A4	50	VAL
27	A5	48	GLU
27	A5	56	LYS
27	A5	57	VAL
28	A6	45	LYS
30	A8	30	ARG
30	A8	40	GLU
32	BE	6	THR
32	BE	13	ALA
32	BE	18	GLY
32	BE	20	GLU
32	BE	73	THR
32	BE	87	ARG
32	BE	97	TRP
32	BE	216	SER
32	BE	217	ARG
33	BF	22	TRP
33	BF	26	LYS
33	BF	34	LEU
33	BF	47	LEU
33	BF	64	VAL
33	BF	129	ALA
33	BF	164	ARG
34	BG	4	TYR
34	BG	15	GLU
34	BG	28	SER
34	BG	171	GLY
35	BH	98	THR
35	BH	128	PRO
35	BH	129	ILE
35	BH	147	ASP
36	BI	13	ASN
37	BJ	7	ALA
37	BJ	8	GLU
37	BJ	14	PRO

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Mol	Chain	Res	Type
37	BJ	17	VAL
38	BK	133	LEU
39	BL	21	PRO
39	BL	44	VAL
39	BL	100	GLY
39	BL	121	ARG
40	BM	16	LEU
40	BM	36	GLY
40	BM	51	ARG
40	BM	59	SER
40	BM	75	ILE
41	BN	54	ARG
41	BN	128	ALA
42	BO	23	LYS
42	BO	27	LEU
42	BO	43	VAL
42	BO	51	ALA
42	BO	89	ARG
42	BO	91	LYS
42	BO	92	ASP
43	BP	7	VAL
43	BP	62	ASN
43	BP	86	CYS
43	BP	100	GLY
46	BS	64	ALA
46	BS	83	GLU
47	BT	33	GLY
47	BT	66	SER
47	BT	81	ARG
48	BU	22	VAL
49	BV	70	LYS
49	BV	82	GLY
50	BW	47	GLY
50	BW	61	SER
50	BW	68	LYS
50	BW	73	HIS
50	BW	95	ALA
50	BW	103	GLY
51	BX	3	LYS
32	CE	5	ILE
32	CE	18	GLY
32	CE	83	MET

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Mol	Chain	Res	Type
32	CE	84	GLU
32	CE	165	VAL
32	CE	194	PRO
32	CE	209	ARG
32	CE	216	SER
33	CF	24	ALA
33	CF	61	ALA
33	CF	66	VAL
33	CF	145	GLY
34	CG	40	PRO
34	CG	41	GLY
34	CG	51	PRO
34	CG	150	GLU
34	CG	155	LEU
34	CG	186	LEU
34	CG	200	GLU
35	CH	12	LEU
37	CJ	7	ALA
38	CK	129	VAL
38	CK	133	LEU
39	CL	109	VAL
40	CM	30	SER
40	CM	36	GLY
41	CN	91	ARG
42	CO	43	VAL
42	CO	116	SER
42	CO	117	ARG
42	CO	121	GLY
43	CP	6	GLY
43	CP	106	ASN
44	CQ	29	ARG
44	CQ	52	GLN
45	CR	77	ARG
45	CR	80	ALA
46	CS	69	THR
46	CS	83	GLU
47	CT	14	LYS
48	CU	20	ALA
48	CU	37	VAL
49	CV	6	LYS
49	CV	9	VAL
49	CV	26	GLY

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Mol	Chain	Res	Type
49	CV	39	THR
49	CV	41	VAL
49	CV	88	LYS
50	CW	11	SER
50	CW	96	GLY
50	CW	102	GLY
50	CW	103	GLY
51	CX	3	LYS
51	CX	9	ARG
3	DD	144	ALA
3	DD	239	ARG
3	DD	257	LEU
4	DE	4	ILE
4	DE	29	GLY
4	DE	50	GLY
4	DE	71	GLY
4	DE	78	LEU
4	DE	89	ASP
4	DE	92	THR
4	DE	185	LYS
4	DE	189	PRO
5	DF	67	GLN
5	DF	132	VAL
5	DF	181	LEU
6	DG	4	ASP
6	DG	5	VAL
6	DG	24	GLY
6	DG	97	ASP
6	DG	110	ALA
6	DG	124	SER
6	DG	137	GLU
6	DG	150	ASP
7	DH	3	ARG
7	DH	21	PRO
7	DH	87	LEU
7	DH	90	LYS
7	DH	128	PRO
7	DH	151	ILE
56	DI	9	LYS
56	DI	19	GLU
56	DI	25	ASP
8	DK	9	LEU

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Mol	Chain	Res	Type
8	DK	36	ALA
8	DK	102	SER
8	DK	104	GLN
8	DK	131	LYS
57	DY	5	ARG
57	DY	7	VAL
57	DY	16	ASN
57	DY	19	ARG
57	DY	27	VAL
57	DY	31	GLY
57	DY	34	ALA
57	DY	48	GLY
57	DY	49	ALA
57	DY	69	PRO
57	DY	72	ASP
57	DY	73	GLY
57	DY	80	VAL
57	DY	87	VAL
57	DY	89	ALA
57	DY	95	GLN
57	DY	103	GLY
57	DY	107	VAL
57	DY	138	LEU
57	DY	140	GLY
57	DY	141	VAL
57	DY	144	ALA
58	DL	17	ALA
58	DL	41	PHE
58	DL	61	ALA
58	DL	65	PHE
58	DL	77	LEU
58	DL	88	ALA
58	DL	110	GLN
58	DL	133	SER
58	DL	135	GLY
58	DL	144	VAL
58	DL	145	LYS
56	DJ	12	LEU
56	DJ	13	SER
56	DJ	16	THR
9	DM	23	LEU
9	DM	36	GLY

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Mol	Chain	Res	Type
10	DN	5	GLN
11	DO	12	ALA
11	DO	29	LYS
11	DO	65	ARG
11	DO	120	ALA
11	DO	125	VAL
11	DO	141	ALA
11	DO	148	LEU
12	DP	59	ARG
12	DP	62	GLY
12	DP	89	ASN
12	DP	99	PRO
12	DP	105	GLU
13	D0	82	GLU
13	D0	86	ARG
14	DQ	4	LEU
14	DQ	12	PHE
14	DQ	57	LYS
14	DQ	96	GLY
14	DQ	109	GLY
15	DR	20	PRO
17	D2	48	GLY
17	D2	50	PRO
17	D2	53	GLU
18	DS	66	GLU
20	DU	41	GLY
20	DU	47	LYS
20	DU	58	GLY
20	DU	91	GLU
20	DU	98	VAL
20	DU	102	CYS
21	DV	31	ARG
21	DV	60	GLU
21	DV	64	GLY
21	DV	93	ASP
21	DV	110	GLY
21	DV	136	PHE
21	DV	178	GLU
21	DV	181	GLU
21	DV	182	LYS
21	DV	188	ALA
22	D3	84	LEU

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Mol	Chain	Res	Type
23	DZ	78	LYS
23	DZ	79	GLY
23	DZ	82	LEU
23	DZ	84	GLY
23	DZ	91	LYS
24	DW	15	LYS
24	DW	42	GLY
25	DX	13	ILE
25	DX	39	ASP
26	D4	8	LYS
26	D4	18	CYS
26	D4	43	TYR
26	D4	66	SER
27	D5	45	VAL
27	D5	58	LEU
28	D6	33	LYS
28	D6	34	LEU
28	D6	35	GLU
28	D6	44	ARG
30	D8	49	VAL
30	D8	64	TYR
3	AD	159	ALA
3	AD	239	ARG
3	AD	263	ARG
4	AE	8	LYS
4	AE	15	PHE
4	AE	17	ASP
4	AE	29	GLY
4	AE	45	THR
4	AE	66	HIS
5	AF	126	VAL
6	AG	81	LYS
6	AG	97	ASP
6	AG	126	ASP
6	AG	155	MET
7	AH	8	PRO
7	AH	10	PRO
7	AH	16	SER
7	AH	21	PRO
7	AH	160	LYS
7	AH	162	ILE
8	AK	59	ALA

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Mol	Chain	Res	Type
8	AK	83	ALA
8	AK	89	TYR
9	AM	67	LEU
9	AM	127	ASP
10	AN	12	ASP
10	AN	120	GLU
11	AO	42	SER
11	AO	63	PRO
12	AP	4	PRO
12	AP	21	THR
12	AP	67	ARG
12	AP	104	PHE
13	A0	3	HIS
13	A0	88	ARG
14	AQ	61	ASN
15	AR	103	ARG
15	AR	116	ALA
16	A1	32	PHE
17	A2	2	PHE
17	A2	37	VAL
17	A2	50	PRO
17	A2	98	GLU
18	AS	32	ALA
18	AS	56	ALA
19	AT	22	ALA
19	AT	60	ARG
21	AV	7	ALA
21	AV	66	SER
21	AV	124	ILE
21	AV	136	PHE
21	AV	158	PRO
22	A3	82	ARG
23	AZ	88	LYS
25	AX	27	GLY
26	A4	25	TYR
26	A4	52	THR
26	A4	57	GLU
26	A4	65	ASP
27	A5	52	TYR
27	A5	55	ARG
28	A6	14	THR
28	A6	15	GLU

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Mol	Chain	Res	Type
28	A6	49	HIS
32	BE	98	LEU
32	BE	101	MET
32	BE	153	ARG
32	BE	154	LEU
32	BE	161	ALA
33	BF	20	SER
33	BF	100	ALA
33	BF	156	ARG
33	BF	168	ALA
33	BF	179	ARG
34	BG	25	ARG
34	BG	161	ASN
34	BG	182	LYS
34	BG	200	GLU
35	BH	104	ALA
36	BI	44	GLY
37	BJ	121	ALA
38	BK	104	ARG
38	BK	108	GLY
39	BL	54	ASP
41	BN	117	ASN
42	BO	29	GLY
42	BO	42	THR
42	BO	45	PRO
42	BO	90	VAL
43	BP	27	LYS
43	BP	84	ILE
43	BP	106	ASN
45	BR	24	SER
45	BR	88	ARG
46	BS	43	LYS
46	BS	53	VAL
46	BS	81	ARG
47	BT	12	SER
47	BT	74	LEU
47	BT	94	ASN
48	BU	31	LEU
48	BU	37	VAL
48	BU	59	SER
49	BV	81	ARG
50	BW	105	SER

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Mol	Chain	Res	Type
32	CE	75	LYS
32	CE	129	GLU
32	CE	155	LEU
32	CE	207	ALA
32	CE	217	ARG
32	CE	229	VAL
33	CF	45	LYS
33	CF	107	GLN
33	CF	117	ALA
33	CF	188	LEU
34	CG	48	ALA
34	CG	73	ARG
34	CG	79	PHE
34	CG	143	GLY
34	CG	164	ALA
34	CG	181	MET
36	CI	40	VAL
36	CI	43	LEU
36	CI	54	LYS
36	CI	70	ASP
36	CI	84	ASN
37	CJ	14	PRO
37	CJ	55	GLY
37	CJ	59	LEU
40	CM	27	ALA
40	CM	59	SER
42	CO	23	LYS
42	CO	115	LYS
43	CP	14	ARG
43	CP	30	ALA
43	CP	59	TYR
43	CP	100	GLY
43	CP	107	ALA
43	CP	124	PRO
44	CQ	44	LEU
45	CR	76	GLU
46	CS	48	TRP
46	CS	76	GLN
47	CT	30	PRO
48	CU	23	LYS
48	CU	54	ARG
48	CU	87	ARG

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Mol	Chain	Res	Type
50	CW	99	LEU
3	DD	26	LYS
3	DD	30	GLU
3	DD	111	LEU
3	DD	224	ALA
4	DE	7	VAL
4	DE	8	LYS
4	DE	53	PRO
4	DE	66	HIS
4	DE	204	ALA
6	DG	48	GLU
6	DG	113	ARG
6	DG	115	ARG
6	DG	143	GLU
7	DH	168	PRO
56	DI	5	ILE
56	DI	18	LEU
56	DI	27	LEU
8	DK	39	ALA
8	DK	87	LYS
8	DK	98	ALA
8	DK	100	ALA
8	DK	109	ILE
8	DK	110	ASP
57	DY	20	ALA
57	DY	45	LYS
57	DY	70	GLU
57	DY	77	PRO
58	DL	16	LYS
58	DL	21	PRO
58	DL	28	GLY
58	DL	40	ALA
58	DL	51	ALA
58	DL	73	PRO
58	DL	101	TRP
58	DL	111	LYS
58	DL	125	ARG
58	DL	138	VAL
9	DM	76	SER
9	DM	114	ARG
9	DM	127	ASP
10	DN	112	MET

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Mol	Chain	Res	Type
11	DO	62	LEU
11	DO	66	GLY
12	DP	6	ARG
12	DP	27	VAL
12	DP	53	ALA
12	DP	91	GLU
12	DP	104	PHE
13	D0	45	ARG
13	D0	59	ASP
13	D0	88	ARG
14	DQ	20	ARG
15	DR	39	ARG
15	DR	55	ASN
15	DR	108	ARG
15	DR	116	ALA
15	DR	134	GLU
16	D1	115	ALA
17	D2	54	GLY
18	DS	80	PRO
20	DU	48	ALA
20	DU	92	ASN
21	DV	13	GLU
21	DV	34	ASN
21	DV	51	ALA
21	DV	171	ILE
22	D3	10	THR
22	D3	15	ASP
25	DX	52	HIS
26	D4	19	GLY
26	D4	23	GLU
26	D4	42	PHE
27	D5	36	CYS
27	D5	55	ARG
30	D8	6	THR
30	D8	29	LYS
30	D8	34	TRP
30	D8	63	PRO
3	AD	64	ILE
4	AE	49	LEU
4	AE	52	LEU
4	AE	68	ALA
4	AE	179	GLU

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Mol	Chain	Res	Type
5	AF	86	GLY
5	AF	133	ASN
6	AG	85	GLY
7	AH	13	LYS
7	AH	80	SER
7	AH	102	ALA
8	AK	58	LEU
8	AK	85	GLU
9	AM	23	LEU
11	AO	4	SER
11	AO	21	ARG
11	AO	48	PRO
11	AO	104	GLY
11	AO	106	LEU
12	AP	6	ARG
12	AP	29	PHE
12	AP	31	ASP
14	AQ	97	ARG
16	A1	92	ARG
16	A1	93	LYS
16	A1	102	GLU
17	A2	87	HIS
19	AT	67	GLY
20	AU	9	LYS
20	AU	43	ASN
20	AU	91	GLU
22	A3	67	VAL
23	AZ	97	LEU
26	A4	7	PRO
26	A4	37	SER
28	A6	13	CYS
28	A6	24	GLU
32	BE	5	ILE
32	BE	19	HIS
32	BE	26	PRO
32	BE	130	ARG
32	BE	150	SER
32	BE	177	ALA
32	BE	190	THR
33	BF	82	GLU
33	BF	107	GLN
34	BG	3	ARG

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Mol	Chain	Res	Type
34	BG	56	VAL
34	BG	89	THR
34	BG	110	PHE
34	BG	181	MET
35	BH	21	ALA
35	BH	112	LEU
36	BI	12	PRO
36	BI	43	LEU
37	BJ	97	GLN
37	BJ	116	ALA
37	BJ	150	ALA
38	BK	2	LEU
38	BK	76	PRO
39	BL	111	ARG
40	BM	70	ARG
40	BM	99	LYS
41	BN	62	GLN
41	BN	89	ALA
41	BN	93	GLN
42	BO	61	THR
42	BO	123	LYS
43	BP	5	ALA
43	BP	42	ALA
43	BP	79	LYS
44	BQ	15	LYS
47	BT	68	ARG
48	BU	82	THR
50	BW	10	LEU
50	BW	25	ARG
32	CE	26	PRO
32	CE	78	GLN
32	CE	139	LYS
32	CE	225	ALA
32	CE	238	LEU
33	CF	4	LYS
33	CF	47	LEU
33	CF	81	GLY
33	CF	125	GLU
34	CG	27	TYR
34	CG	189	PRO
35	CH	128	PRO
39	CL	11	LYS

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Mol	Chain	Res	Type
39	CL	44	VAL
39	CL	54	ASP
39	CL	117	HIS
41	CN	103	LEU
43	CP	21	TYR
43	CP	45	VAL
43	CP	49	THR
43	CP	68	GLY
44	CQ	22	THR
45	CR	23	GLY
47	CT	34	LYS
49	CV	4	SER
49	CV	78	ARG
3	DD	3	VAL
3	DD	46	GLN
4	DE	57	LYS
4	DE	82	ARG
4	DE	86	PRO
4	DE	90	THR
5	DF	25	PRO
5	DF	47	GLY
5	DF	130	ALA
6	DG	36	LYS
6	DG	86	MET
6	DG	116	ASP
6	DG	181	ARG
7	DH	81	GLU
7	DH	152	ARG
56	DI	16	THR
56	DI	28	LYS
8	DK	10	GLU
8	DK	12	LEU
8	DK	18	VAL
8	DK	113	ARG
8	DK	118	LYS
8	DK	133	HIS
57	DY	24	PHE
57	DY	98	LYS
57	DY	101	PRO
58	DL	59	ILE
58	DL	117	THR
58	DL	142	PRO

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Mol	Chain	Res	Type
56	DJ	14	GLN
9	DM	135	PRO
11	DO	35	HIS
11	DO	90	ARG
11	DO	93	GLY
11	DO	95	VAL
11	DO	136	GLU
11	DO	147	LEU
12	DP	18	LYS
13	D0	42	LYS
13	D0	117	VAL
14	DQ	75	GLU
14	DQ	97	ARG
14	DQ	110	LEU
15	DR	36	GLU
15	DR	57	PHE
16	D1	24	TYR
16	D1	92	ARG
17	D2	36	PRO
18	DS	65	LEU
19	DT	91	ALA
20	DU	39	VAL
21	DV	6	LYS
21	DV	53	ILE
21	DV	114	GLY
21	DV	158	PRO
21	DV	164	ALA
21	DV	195	GLU
23	DZ	64	ALA
23	DZ	97	LEU
26	D4	24	THR
27	D5	42	PRO
27	D5	59	GLU
29	D7	47	ARG
3	AD	125	ILE
3	AD	127	VAL
3	AD	240	ALA
3	AD	271	ILE
4	AE	82	ARG
5	AF	124	LEU
6	AG	47	LYS
6	AG	86	MET

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Mol	Chain	Res	Type
6	AG	124	SER
6	AG	136	ARG
6	AG	176	LEU
6	AG	181	ARG
7	AH	17	VAL
7	AH	49	VAL
7	AH	99	VAL
7	AH	137	ASP
8	AK	50	ARG
8	AK	136	VAL
9	AM	3	THR
10	AN	68	GLU
11	AO	7	ARG
11	AO	43	GLY
12	AP	7	MET
12	AP	24	GLY
13	A0	107	ASP
14	AQ	51	ALA
14	AQ	62	LYS
14	AQ	90	GLY
14	AQ	94	TYR
15	AR	57	PHE
17	A2	62	LEU
17	A2	99	ILE
19	AT	51	VAL
19	AT	61	GLY
20	AU	58	GLY
21	AV	42	VAL
21	AV	95	PRO
21	AV	121	HIS
21	AV	163	LEU
21	AV	170	THR
23	AZ	87	PRO
26	A4	15	ILE
28	A6	35	GLU
28	A6	36	LEU
28	A6	51	GLU
28	A6	52	VAL
32	BE	52	GLU
33	BF	15	THR
33	BF	66	VAL
34	BG	31	CYS

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Mol	Chain	Res	Type
34	BG	160	GLN
35	BH	105	VAL
37	BJ	51	GLN
37	BJ	111	ARG
37	BJ	117	ALA
37	BJ	154	TYR
38	BK	14	ARG
39	BL	66	ARG
40	BM	54	PHE
41	BN	65	ALA
42	BO	26	ALA
43	BP	67	GLU
44	BQ	19	ARG
44	BQ	52	GLN
46	BS	57	ARG
49	BV	10	PHE
49	BV	72	GLY
32	CE	8	LYS
32	CE	13	ALA
32	CE	67	THR
32	CE	87	ARG
32	CE	101	MET
33	CF	84	ILE
34	CG	20	TYR
35	CH	52	PRO
35	CH	129	ILE
37	CJ	17	VAL
39	CL	21	PRO
39	CL	56	LEU
42	CO	12	ARG
42	CO	28	LYS
42	CO	51	ALA
42	CO	123	LYS
43	CP	3	ARG
43	CP	70	LEU
43	CP	121	LYS
44	CQ	12	ARG
46	CS	41	PRO
3	DD	45	ASN
3	DD	159	ALA
4	DE	33	VAL
4	DE	46	ALA

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Mol	Chain	Res	Type
5	DF	8	GLN
6	DG	3	LEU
6	DG	6	ALA
6	DG	117	PHE
7	DH	110	SER
8	DK	33	ARG
8	DK	145	VAL
57	DY	68	LEU
58	DL	99	ILE
58	DL	105	LEU
9	DM	40	PRO
10	DN	117	LEU
11	DO	109	GLY
12	DP	11	LYS
12	DP	67	ARG
13	D0	14	SER
13	D0	75	LEU
14	DQ	21	THR
15	DR	4	GLY
16	D1	9	VAL
17	D2	30	GLY
20	DU	55	TYR
23	DZ	92	LYS
26	D4	9	LEU
26	D4	28	LYS
27	D5	51	TYR
28	D6	24	GLU
28	D6	27	LYS
3	AD	106	ILE
4	AE	73	GLU
4	AE	135	HIS
4	AE	169	ASN
4	AE	189	PRO
5	AF	28	ILE
7	AH	39	PRO
8	AK	135	GLU
9	AM	110	GLY
11	AO	12	ALA
11	AO	107	LYS
16	A1	73	GLY
20	AU	22	GLY
20	AU	52	SER

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Mol	Chain	Res	Type
20	AU	55	TYR
21	AV	134	PRO
24	AW	67	LYS
25	AX	59	VAL
26	A4	23	GLU
26	A4	24	THR
26	A4	48	ARG
26	A4	70	GLY
28	A6	17	LYS
30	A8	38	GLY
32	BE	80	ILE
32	BE	83	MET
32	BE	106	LYS
32	BE	122	PHE
34	BG	90	GLY
40	BM	32	ALA
42	BO	75	HIS
42	BO	117	ARG
43	BP	38	GLY
47	BT	64	PRO
48	BU	23	LYS
51	BX	17	THR
32	CE	213	LEU
32	CE	236	TYR
34	CG	32	ALA
39	CL	127	LYS
42	CO	19	ARG
45	CR	16	ALA
50	CW	84	LEU
50	CW	97	ALA
3	DD	169	GLU
4	DE	39	PRO
4	DE	47	VAL
6	DG	82	LEU
6	DG	84	LYS
6	DG	85	GLY
6	DG	112	PRO
7	DH	55	PRO
7	DH	111	HIS
56	DI	4	ASP
8	DK	82	ARG
9	DM	97	ARG

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Mol	Chain	Res	Type
11	DO	7	ARG
11	DO	71	VAL
21	DV	105	VAL
21	DV	115	GLY
21	DV	141	VAL
21	DV	173	ALA
28	D6	30	THR
5	AF	14	PRO
7	AH	142	GLY
8	AK	109	ILE
9	AM	77	GLY
11	AO	71	VAL
20	AU	61	ILE
28	A6	48	VAL
33	BF	116	VAL
34	BG	5	ILE
35	BH	51	VAL
35	BH	59	GLY
40	BM	74	ILE
41	BN	90	GLY
45	BR	86	GLY
38	CK	83	ILE
39	CL	81	ILE
40	CM	91	PRO
49	CV	31	ILE
4	DE	190	GLY
57	DY	105	PRO
57	DY	139	VAL
9	DM	6	PRO
20	DU	3	VAL
21	DV	111	VAL
21	DV	193	GLU
5	AF	30	PRO
7	AH	151	ILE
11	AO	8	PRO
12	AP	47	ILE
17	A2	30	GLY
20	AU	7	VAL
25	AX	43	ILE
34	BG	124	GLY
35	BH	39	GLY
32	CE	80	ILE

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Mol	Chain	Res	Type
38	CK	73	ASP
47	CT	33	GLY
48	CU	22	VAL
48	CU	39	VAL
4	DE	55	ASN
4	DE	130	GLY
5	DF	30	PRO
7	DH	7	LEU
20	DU	56	PRO
20	DU	96	ILE
21	DV	94	GLU
5	AF	114	VAL
17	A2	36	PRO
20	AU	53	PRO
50	BW	97	ALA
33	CF	96	GLY
34	CG	105	VAL
37	CJ	19	GLY
45	CR	18	PHE
50	CW	63	ILE
50	CW	88	VAL
4	DE	93	VAL
9	DM	111	PRO
11	DO	47	ASP
22	D3	8	GLY
4	AE	173	VAL
7	AH	79	VAL
12	AP	52	VAL
17	A2	29	PRO
19	AT	24	GLY
21	AV	165	VAL
23	AZ	36	GLY
34	BG	178	VAL
38	BK	53	VAL
40	BM	93	GLY
36	CI	37	VAL
40	CM	37	PRO
4	DE	75	VAL
58	DL	55	VAL
58	DL	78	ILE
56	DJ	24	ILE
12	DP	81	VAL

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Mol	Chain	Res	Type
12	DP	109	VAL
14	DQ	82	ILE
17	D2	28	GLU
20	DU	52	SER
27	D5	57	VAL
13	A0	106	GLY
33	BF	41	GLY
9	AM	126	PRO
58	DL	13	PRO
4	AE	147	PRO
8	AK	132	PRO

5.3.2 Protein sidechains ⓘ

In the following table, the Percentiles column shows the percent sidechain outliers of the chain as a percentile score with respect to all X-ray entries followed by that with respect to entries of similar resolution.

The Analysed column shows the number of residues for which the sidechain conformation was analysed, and the total number of residues.

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
3	AD	214/218 (98%)	182 (85%)	32 (15%)	3	12
3	DD	214/218 (98%)	181 (85%)	33 (15%)	2	11
4	AE	165/166 (99%)	146 (88%)	19 (12%)	5	22
4	DE	165/166 (99%)	128 (78%)	37 (22%)	1	3
5	AF	165/166 (99%)	151 (92%)	14 (8%)	10	37
5	DF	161/166 (97%)	136 (84%)	25 (16%)	2	11
6	AG	155/156 (99%)	138 (89%)	17 (11%)	6	25
6	DG	155/156 (99%)	134 (86%)	21 (14%)	4	16
7	AH	142/148 (96%)	130 (92%)	12 (8%)	10	37
7	DH	142/148 (96%)	117 (82%)	25 (18%)	2	8
8	AK	122/124 (98%)	105 (86%)	17 (14%)	3	15
8	DK	122/124 (98%)	92 (75%)	30 (25%)	0	2
9	AM	117/119 (98%)	100 (86%)	17 (14%)	3	13
9	DM	117/119 (98%)	102 (87%)	15 (13%)	4	18
10	AN	100/100 (100%)	83 (83%)	17 (17%)	2	9

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
10	DN	100/100 (100%)	84 (84%)	16 (16%)	2	11
11	AO	116/116 (100%)	91 (78%)	25 (22%)	1	4
11	DO	116/116 (100%)	84 (72%)	32 (28%)	0	1
12	AP	111/111 (100%)	91 (82%)	20 (18%)	1	7
12	DP	111/111 (100%)	97 (87%)	14 (13%)	4	18
13	A0	100/101 (99%)	88 (88%)	12 (12%)	5	20
13	D0	101/101 (100%)	83 (82%)	18 (18%)	2	8
14	AQ	87/88 (99%)	76 (87%)	11 (13%)	4	18
14	DQ	87/88 (99%)	74 (85%)	13 (15%)	3	13
15	AR	120/127 (94%)	98 (82%)	22 (18%)	1	7
15	DR	120/127 (94%)	96 (80%)	24 (20%)	1	5
16	A1	93/94 (99%)	82 (88%)	11 (12%)	5	21
16	D1	93/94 (99%)	77 (83%)	16 (17%)	2	9
17	A2	82/82 (100%)	72 (88%)	10 (12%)	5	19
17	D2	82/82 (100%)	64 (78%)	18 (22%)	1	4
18	AS	92/92 (100%)	80 (87%)	12 (13%)	4	18
18	DS	92/92 (100%)	79 (86%)	13 (14%)	3	15
19	AT	74/78 (95%)	67 (90%)	7 (10%)	8	31
19	DT	74/78 (95%)	66 (89%)	8 (11%)	6	25
20	AU	85/91 (93%)	61 (72%)	24 (28%)	0	1
20	DU	85/91 (93%)	66 (78%)	19 (22%)	1	3
21	AV	164/179 (92%)	130 (79%)	34 (21%)	1	5
21	DV	173/179 (97%)	131 (76%)	42 (24%)	0	2
22	A3	66/67 (98%)	59 (89%)	7 (11%)	6	26
22	D3	66/67 (98%)	59 (89%)	7 (11%)	6	26
23	AZ	82/83 (99%)	70 (85%)	12 (15%)	3	13
23	DZ	82/83 (99%)	71 (87%)	11 (13%)	4	16
24	AW	64/67 (96%)	58 (91%)	6 (9%)	8	32
24	DW	64/67 (96%)	53 (83%)	11 (17%)	2	9
25	AX	51/52 (98%)	42 (82%)	9 (18%)	2	8
25	DX	51/52 (98%)	48 (94%)	3 (6%)	19	50

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
26	A4	63/63 (100%)	45 (71%)	18 (29%)	0	1
26	D4	63/63 (100%)	49 (78%)	14 (22%)	1	4
27	A5	51/52 (98%)	45 (88%)	6 (12%)	5	21
27	D5	51/52 (98%)	36 (71%)	15 (29%)	0	1
28	A6	44/52 (85%)	31 (70%)	13 (30%)	0	1
28	D6	44/52 (85%)	28 (64%)	16 (36%)	0	0
29	A7	42/42 (100%)	34 (81%)	8 (19%)	1	6
29	D7	42/42 (100%)	38 (90%)	4 (10%)	8	31
30	A8	54/55 (98%)	43 (80%)	11 (20%)	1	5
30	D8	54/55 (98%)	39 (72%)	15 (28%)	0	1
32	BE	205/220 (93%)	187 (91%)	18 (9%)	10	36
32	CE	205/220 (93%)	183 (89%)	22 (11%)	6	26
33	BF	160/188 (85%)	140 (88%)	20 (12%)	4	18
33	CF	159/188 (85%)	146 (92%)	13 (8%)	11	38
34	BG	180/181 (99%)	155 (86%)	25 (14%)	3	15
34	CG	180/181 (99%)	160 (89%)	20 (11%)	6	24
35	BH	116/123 (94%)	103 (89%)	13 (11%)	6	24
35	CH	116/123 (94%)	101 (87%)	15 (13%)	4	18
36	BI	90/90 (100%)	81 (90%)	9 (10%)	7	28
36	CI	90/90 (100%)	85 (94%)	5 (6%)	21	52
37	BJ	126/127 (99%)	114 (90%)	12 (10%)	8	31
37	CJ	126/127 (99%)	118 (94%)	8 (6%)	18	48
38	BK	119/119 (100%)	112 (94%)	7 (6%)	19	50
38	CK	119/119 (100%)	110 (92%)	9 (8%)	13	41
39	BL	98/99 (99%)	86 (88%)	12 (12%)	5	19
39	CL	98/99 (99%)	84 (86%)	14 (14%)	3	14
40	BM	89/92 (97%)	79 (89%)	10 (11%)	6	24
40	CM	89/92 (97%)	80 (90%)	9 (10%)	7	28
41	BN	90/99 (91%)	85 (94%)	5 (6%)	21	52
41	CN	90/99 (91%)	88 (98%)	2 (2%)	52	78
42	BO	104/109 (95%)	91 (88%)	13 (12%)	4	18

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
42	CO	104/109 (95%)	93 (89%)	11 (11%)	6	26
43	BP	97/101 (96%)	86 (89%)	11 (11%)	6	23
43	CP	100/101 (99%)	82 (82%)	18 (18%)	1	7
44	BQ	49/50 (98%)	42 (86%)	7 (14%)	3	14
44	CQ	49/50 (98%)	43 (88%)	6 (12%)	5	19
45	BR	79/80 (99%)	73 (92%)	6 (8%)	13	41
45	CR	79/80 (99%)	72 (91%)	7 (9%)	9	34
46	BS	72/74 (97%)	66 (92%)	6 (8%)	11	38
46	CS	72/74 (97%)	61 (85%)	11 (15%)	2	12
47	BT	95/97 (98%)	91 (96%)	4 (4%)	30	62
47	CT	95/97 (98%)	89 (94%)	6 (6%)	18	48
48	BU	63/77 (82%)	51 (81%)	12 (19%)	1	6
48	CU	63/77 (82%)	51 (81%)	12 (19%)	1	6
49	BV	72/80 (90%)	59 (82%)	13 (18%)	1	7
49	CV	75/80 (94%)	53 (71%)	22 (29%)	0	1
50	BW	76/82 (93%)	65 (86%)	11 (14%)	3	13
50	CW	76/82 (93%)	69 (91%)	7 (9%)	9	33
51	BX	20/22 (91%)	17 (85%)	3 (15%)	3	12
51	CX	20/22 (91%)	19 (95%)	1 (5%)	24	57
56	DI	26/90 (29%)	15 (58%)	11 (42%)	0	0
56	DJ	26/90 (29%)	14 (54%)	12 (46%)	0	0
57	DY	117/135 (87%)	47 (40%)	70 (60%)	0	0
58	DL	109/111 (98%)	52 (48%)	57 (52%)	0	0
All	All	9931/10424 (95%)	8408 (85%)	1523 (15%)	2	12

All (1523) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
3	AD	10	THR
3	AD	21	PHE
3	AD	24	ILE
3	AD	48	ARG
3	AD	49	ILE
3	AD	61	LEU

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Mol	Chain	Res	Type
3	AD	64	ILE
3	AD	65	ILE
3	AD	70	TRP
3	AD	73	VAL
3	AD	94	LEU
3	AD	95	LEU
3	AD	105	ILE
3	AD	106	ILE
3	AD	131	LEU
3	AD	138	VAL
3	AD	147	LEU
3	AD	150	LYS
3	AD	157	ARG
3	AD	161	THR
3	AD	166	GLN
3	AD	168	ARG
3	AD	192	THR
3	AD	198	ASN
3	AD	211	ARG
3	AD	237	GLU
3	AD	242	ARG
3	AD	244	ARG
3	AD	255	LYS
3	AD	260	ARG
3	AD	268	ARG
3	AD	271	ILE
4	AE	4	ILE
4	AE	5	LEU
4	AE	37	ARG
4	AE	49	LEU
4	AE	59	VAL
4	AE	78	LEU
4	AE	79	ARG
4	AE	107	THR
4	AE	119	ARG
4	AE	132	HIS
4	AE	134	ILE
4	AE	144	ARG
4	AE	154	LYS
4	AE	167	VAL
4	AE	178	GLU
4	AE	197	ILE

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Mol	Chain	Res	Type
4	AE	200	GLU
4	AE	202	LYS
4	AE	203	LYS
5	AF	7	TYR
5	AF	17	ARG
5	AF	40	GLN
5	AF	48	THR
5	AF	52	LYS
5	AF	65	TRP
5	AF	66	PRO
5	AF	68	LYS
5	AF	70	THR
5	AF	74	ARG
5	AF	88	VAL
5	AF	96	ASP
5	AF	125	LEU
5	AF	165	ARG
6	AG	7	LEU
6	AG	16	ARG
6	AG	67	LYS
6	AG	80	PHE
6	AG	88	ILE
6	AG	91	ARG
6	AG	95	ARG
6	AG	101	ILE
6	AG	115	ARG
6	AG	118	ARG
6	AG	121	ASN
6	AG	123	ASN
6	AG	135	LEU
6	AG	139	LEU
6	AG	156	ASP
6	AG	159	VAL
6	AG	174	GLU
7	AH	4	ILE
7	AH	6	ARG
7	AH	13	LYS
7	AH	32	GLU
7	AH	54	ARG
7	AH	81	GLU
7	AH	85	LYS
7	AH	89	ILE

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Mol	Chain	Res	Type
7	AH	105	LEU
7	AH	116	GLU
7	AH	143	GLN
7	AH	163	TYR
8	AK	1	MET
8	AK	5	LEU
8	AK	14	ASP
8	AK	48	GLU
8	AK	52	ARG
8	AK	54	GLN
8	AK	56	LYS
8	AK	62	LYS
8	AK	82	ARG
8	AK	99	GLU
8	AK	102	SER
8	AK	109	ILE
8	AK	117	GLU
8	AK	126	TYR
8	AK	130	TYR
8	AK	133	HIS
8	AK	135	GLU
9	AM	1	MET
9	AM	7	LYS
9	AM	32	THR
9	AM	42	TRP
9	AM	45	ASN
9	AM	48	MET
9	AM	50	ASP
9	AM	71	ILE
9	AM	87	LEU
9	AM	96	GLU
9	AM	99	LEU
9	AM	111	PRO
9	AM	112	LEU
9	AM	127	ASP
9	AM	131	GLN
9	AM	137	LYS
9	AM	138	LEU
10	AN	7	TYR
10	AN	8	LEU
10	AN	9	GLU
10	AN	24	VAL

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Mol	Chain	Res	Type
10	AN	29	ASN
10	AN	32	TYR
10	AN	38	VAL
10	AN	48	PRO
10	AN	49	ARG
10	AN	66	LYS
10	AN	69	ILE
10	AN	73	ASP
10	AN	78	ARG
10	AN	80	ASP
10	AN	97	ARG
10	AN	99	PHE
10	AN	104	ARG
11	AO	10	PRO
11	AO	21	ARG
11	AO	27	HIS
11	AO	32	THR
11	AO	36	LYS
11	AO	52	GLU
11	AO	55	ARG
11	AO	59	LEU
11	AO	62	LEU
11	AO	65	ARG
11	AO	67	MET
11	AO	75	ILE
11	AO	81	GLN
11	AO	85	LEU
11	AO	91	PHE
11	AO	98	GLU
11	AO	110	TYR
11	AO	112	LEU
11	AO	115	LEU
11	AO	117	GLU
11	AO	136	GLU
11	AO	138	LEU
11	AO	144	GLU
11	AO	147	LEU
11	AO	148	LEU
12	AP	3	MET
12	AP	8	LYS
12	AP	14	ARG
12	AP	16	ARG

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Mol	Chain	Res	Type
12	AP	22	LYS
12	AP	29	PHE
12	AP	38	GLU
12	AP	45	GLN
12	AP	56	ARG
12	AP	63	LYS
12	AP	82	ARG
12	AP	83	MET
12	AP	89	ASN
12	AP	103	MET
12	AP	110	THR
12	AP	113	GLN
12	AP	115	MET
12	AP	133	ARG
12	AP	139	GLU
12	AP	141	GLN
13	A0	11	ASN
13	A0	15	SER
13	A0	18	LEU
13	A0	34	ILE
13	A0	67	LEU
13	A0	71	GLN
13	A0	74	LYS
13	A0	76	VAL
13	A0	79	LEU
13	A0	89	ASP
13	A0	104	ARG
13	A0	118	GLU
14	AQ	12	PHE
14	AQ	15	ARG
14	AQ	19	LYS
14	AQ	20	ARG
14	AQ	23	ARG
14	AQ	36	TYR
14	AQ	56	LEU
14	AQ	57	LYS
14	AQ	58	LEU
14	AQ	103	GLU
14	AQ	110	LEU
15	AR	8	LYS
15	AR	9	LEU
15	AR	11	GLU

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Mol	Chain	Res	Type
15	AR	16	ARG
15	AR	27	THR
15	AR	29	ARG
15	AR	30	VAL
15	AR	35	LYS
15	AR	41	ARG
15	AR	51	ARG
15	AR	58	ASN
15	AR	59	THR
15	AR	74	ARG
15	AR	78	LEU
15	AR	93	ARG
15	AR	95	ARG
15	AR	98	LYS
15	AR	99	LEU
15	AR	103	ARG
15	AR	115	ARG
15	AR	124	ASP
15	AR	137	LYS
16	A1	5	LYS
16	A1	31	SER
16	A1	37	GLU
16	A1	52	ARG
16	A1	56	ASP
16	A1	64	ARG
16	A1	74	LEU
16	A1	78	THR
16	A1	79	PHE
16	A1	88	ILE
16	A1	98	LEU
17	A2	15	GLU
17	A2	28	GLU
17	A2	35	LEU
17	A2	44	LYS
17	A2	64	HIS
17	A2	74	LYS
17	A2	75	PHE
17	A2	80	GLN
17	A2	81	TYR
17	A2	82	ARG
18	AS	11	ARG
18	AS	14	PRO

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Mol	Chain	Res	Type
18	AS	37	ARG
18	AS	39	THR
18	AS	51	LEU
18	AS	59	VAL
18	AS	65	LEU
18	AS	70	TYR
18	AS	88	ARG
18	AS	96	ILE
18	AS	107	LEU
18	AS	113	LYS
19	AT	12	VAL
19	AT	48	LYS
19	AT	57	LEU
19	AT	63	LYS
19	AT	69	TYR
19	AT	76	ARG
19	AT	80	ILE
20	AU	3	VAL
20	AU	7	VAL
20	AU	12	THR
20	AU	20	TYR
20	AU	26	LYS
20	AU	28	LYS
20	AU	33	LYS
20	AU	43	ASN
20	AU	50	ARG
20	AU	55	TYR
20	AU	57	GLN
20	AU	60	PHE
20	AU	62	GLU
20	AU	63	LYS
20	AU	67	LEU
20	AU	71	LYS
20	AU	75	ILE
20	AU	76	CYS
20	AU	77	PRO
20	AU	85	VAL
20	AU	86	ARG
20	AU	95	LYS
20	AU	96	ILE
20	AU	97	ARG
21	AV	2	GLU

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Mol	Chain	Res	Type
21	AV	8	TYR
21	AV	14	LYS
21	AV	29	TYR
21	AV	38	TYR
21	AV	61	LEU
21	AV	63	ASP
21	AV	67	LEU
21	AV	72	ARG
21	AV	77	ASP
21	AV	81	ARG
21	AV	96	VAL
21	AV	105	VAL
21	AV	108	PRO
21	AV	112	ARG
21	AV	117	LEU
21	AV	118	GLN
21	AV	119	GLU
21	AV	122	ARG
21	AV	131	ARG
21	AV	140	ASP
21	AV	142	SER
21	AV	145	GLU
21	AV	146	ILE
21	AV	156	LYS
21	AV	157	LEU
21	AV	159	PRO
21	AV	161	VAL
21	AV	162	GLU
21	AV	163	LEU
21	AV	178	GLU
21	AV	182	LYS
21	AV	183	LEU
21	AV	186	GLU
22	A3	3	HIS
22	A3	5	LYS
22	A3	7	LEU
22	A3	25	ARG
22	A3	36	ILE
22	A3	64	ASP
22	A3	74	ARG
23	AZ	4	VAL
23	AZ	34	THR

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Mol	Chain	Res	Type
23	AZ	38	SER
23	AZ	56	GLN
23	AZ	78	LYS
23	AZ	80	LEU
23	AZ	81	LYS
23	AZ	83	GLU
23	AZ	87	PRO
23	AZ	90	ILE
23	AZ	91	LYS
23	AZ	95	LEU
24	AW	9	GLN
24	AW	48	HIS
24	AW	50	ILE
24	AW	53	LEU
24	AW	57	ILE
24	AW	62	THR
25	AX	4	LEU
25	AX	6	VAL
25	AX	8	LEU
25	AX	24	LYS
25	AX	38	GLU
25	AX	40	THR
25	AX	46	ASN
25	AX	52	HIS
25	AX	60	GLU
26	A4	1	MET
26	A4	2	LYS
26	A4	7	PRO
26	A4	8	LYS
26	A4	9	LEU
26	A4	18	CYS
26	A4	23	GLU
26	A4	33	VAL
26	A4	36	CYS
26	A4	38	LYS
26	A4	39	CYS
26	A4	47	GLN
26	A4	55	ARG
26	A4	58	ARG
26	A4	60	GLN
26	A4	62	ARG
26	A4	67	TYR

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Mol	Chain	Res	Type
26	A4	69	LYS
27	A5	3	LYS
27	A5	4	HIS
27	A5	29	THR
27	A5	49	CYS
27	A5	51	TYR
27	A5	52	TYR
28	A6	9	LEU
28	A6	17	LYS
28	A6	21	TYR
28	A6	25	LYS
28	A6	27	LYS
28	A6	29	ASN
28	A6	30	THR
28	A6	37	ARG
28	A6	39	TYR
28	A6	45	LYS
28	A6	46	HIS
28	A6	52	VAL
28	A6	53	LYS
29	A7	4	THR
29	A7	8	ASN
29	A7	9	ARG
29	A7	32	LYS
29	A7	43	THR
29	A7	47	ARG
29	A7	48	LYS
29	A7	49	ARG
30	A8	25	MET
30	A8	32	LEU
30	A8	34	TRP
30	A8	40	GLU
30	A8	41	ILE
30	A8	46	ARG
30	A8	48	PHE
30	A8	54	GLU
30	A8	56	GLU
30	A8	61	LEU
30	A8	64	TYR
32	BE	12	GLU
32	BE	19	HIS
32	BE	22	LYS

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Mol	Chain	Res	Type
32	BE	23	ARG
32	BE	56	ARG
32	BE	80	ILE
32	BE	92	TYR
32	BE	111	ARG
32	BE	113	HIS
32	BE	121	LEU
32	BE	137	ARG
32	BE	140	HIS
32	BE	145	LEU
32	BE	163	PHE
32	BE	178	ARG
32	BE	187	LEU
32	BE	224	GLN
32	BE	232	PRO
33	BF	5	ILE
33	BF	16	ARG
33	BF	17	ASP
33	BF	18	TRP
33	BF	27	LYS
33	BF	28	GLN
33	BF	29	TYR
33	BF	43	LEU
33	BF	62	ASP
33	BF	79	ARG
33	BF	82	GLU
33	BF	85	ARG
33	BF	101	LEU
33	BF	119	ARG
33	BF	127	ARG
33	BF	131	ARG
33	BF	156	ARG
33	BF	170	GLN
33	BF	193	TYR
33	BF	196	LEU
34	BG	19	LEU
34	BG	21	LEU
34	BG	24	GLU
34	BG	25	ARG
34	BG	26	CYS
34	BG	29	PRO
34	BG	30	LYS

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Mol	Chain	Res	Type
34	BG	36	ARG
34	BG	38	TYR
34	BG	58	LEU
34	BG	61	LYS
34	BG	86	LYS
34	BG	97	LEU
34	BG	107	ARG
34	BG	108	LEU
34	BG	110	PHE
34	BG	119	GLN
34	BG	122	ARG
34	BG	135	LEU
34	BG	138	TYR
34	BG	150	GLU
34	BG	155	LEU
34	BG	187	ARG
34	BG	191	ARG
34	BG	209	ARG
35	BH	5	ASP
35	BH	6	PHE
35	BH	12	LEU
35	BH	13	ILE
35	BH	20	GLN
35	BH	24	ARG
35	BH	26	PHE
35	BH	33	VAL
35	BH	72	GLN
35	BH	78	HIS
35	BH	79	GLU
35	BH	101	ILE
35	BH	126	ARG
36	BI	1	MET
36	BI	15	ASP
36	BI	28	ARG
36	BI	36	ARG
36	BI	54	LYS
36	BI	55	ASP
36	BI	78	GLU
36	BI	95	GLU
36	BI	98	LEU
37	BJ	8	GLU
37	BJ	12	LEU

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Mol	Chain	Res	Type
37	BJ	14	PRO
37	BJ	29	LYS
37	BJ	33	ASP
37	BJ	54	THR
37	BJ	57	GLU
37	BJ	64	GLN
37	BJ	84	ASN
37	BJ	136	LYS
37	BJ	137	LYS
37	BJ	151	TYR
38	BK	1	MET
38	BK	25	ASP
38	BK	30	ARG
38	BK	52	ASP
38	BK	63	LEU
38	BK	81	HIS
38	BK	104	ARG
39	BL	10	ARG
39	BL	12	GLU
39	BL	20	ARG
39	BL	75	ASP
39	BL	95	LYS
39	BL	102	LEU
39	BL	104	ARG
39	BL	112	LYS
39	BL	113	LYS
39	BL	114	TYR
39	BL	121	ARG
39	BL	125	TYR
40	BM	16	LEU
40	BM	22	LYS
40	BM	25	GLU
40	BM	47	PHE
40	BM	57	LYS
40	BM	62	HIS
40	BM	74	ILE
40	BM	79	ARG
40	BM	80	LYS
40	BM	96	ILE
41	BN	29	ILE
41	BN	30	VAL
41	BN	54	ARG

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Mol	Chain	Res	Type
41	BN	116	HIS
41	BN	120	ARG
42	BO	17	LYS
42	BO	20	LYS
42	BO	23	LYS
42	BO	27	LEU
42	BO	39	VAL
42	BO	41	ARG
42	BO	42	THR
42	BO	46	LYS
42	BO	47	LYS
42	BO	64	TYR
42	BO	65	GLU
42	BO	66	VAL
42	BO	84	LEU
43	BP	19	LEU
43	BP	64	TRP
43	BP	70	LEU
43	BP	73	GLU
43	BP	79	LYS
43	BP	80	ARG
43	BP	82	MET
43	BP	83	ASP
43	BP	86	CYS
43	BP	88	ARG
43	BP	108	ARG
44	BQ	6	LEU
44	BQ	8	GLU
44	BQ	15	LYS
44	BQ	18	VAL
44	BQ	29	ARG
44	BQ	44	LEU
44	BQ	61	TRP
45	BR	3	ILE
45	BR	10	LYS
45	BR	26	GLU
45	BR	38	ARG
45	BR	41	GLU
45	BR	82	ILE
46	BS	20	VAL
46	BS	27	LYS
46	BS	54	GLU

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Mol	Chain	Res	Type
46	BS	55	ARG
46	BS	69	THR
46	BS	72	ARG
47	BT	49	GLU
47	BT	52	LYS
47	BT	68	ARG
47	BT	74	LEU
48	BU	21	LYS
48	BU	23	LYS
48	BU	26	LEU
48	BU	31	LEU
48	BU	37	VAL
48	BU	38	GLU
48	BU	47	THR
48	BU	54	ARG
48	BU	58	LEU
48	BU	69	THR
48	BU	81	PHE
48	BU	87	ARG
49	BV	6	LYS
49	BV	20	LEU
49	BV	22	LEU
49	BV	23	ASN
49	BV	27	GLU
49	BV	29	ARG
49	BV	37	ARG
49	BV	39	THR
49	BV	41	VAL
49	BV	44	MET
49	BV	77	THR
49	BV	78	ARG
49	BV	86	GLU
50	BW	10	LEU
50	BW	24	LEU
50	BW	26	ASN
50	BW	29	LYS
50	BW	36	LEU
50	BW	50	GLU
50	BW	64	ASP
50	BW	68	LYS
50	BW	73	HIS
50	BW	83	ARG

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Mol	Chain	Res	Type
50	BW	93	GLU
51	BX	9	ARG
51	BX	15	ARG
51	BX	26	LYS
32	CE	5	ILE
32	CE	8	LYS
32	CE	15	VAL
32	CE	16	HIS
32	CE	24	TRP
32	CE	36	ARG
32	CE	67	THR
32	CE	75	LYS
32	CE	82	ARG
32	CE	96	ARG
32	CE	146	GLN
32	CE	158	LEU
32	CE	178	ARG
32	CE	187	LEU
32	CE	192	SER
32	CE	195	ASP
32	CE	196	LEU
32	CE	200	ILE
32	CE	204	ASN
32	CE	212	GLN
32	CE	215	LEU
32	CE	236	TYR
33	CF	5	ILE
33	CF	12	LEU
33	CF	16	ARG
33	CF	17	ASP
33	CF	21	ARG
33	CF	27	LYS
33	CF	29	TYR
33	CF	34	LEU
33	CF	85	ARG
33	CF	94	LEU
33	CF	165	THR
33	CF	188	LEU
33	CF	196	LEU
34	CG	3	ARG
34	CG	7	PRO
34	CG	10	ARG

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Mol	Chain	Res	Type
34	CG	19	LEU
34	CG	33	MET
34	CG	79	PHE
34	CG	86	LYS
34	CG	110	PHE
34	CG	114	ARG
34	CG	122	ARG
34	CG	127	THR
34	CG	131	ARG
34	CG	135	LEU
34	CG	150	GLU
34	CG	154	ASN
34	CG	190	ASP
34	CG	193	ASP
34	CG	196	LEU
34	CG	200	GLU
34	CG	201	GLN
35	CH	6	PHE
35	CH	10	MET
35	CH	12	LEU
35	CH	16	THR
35	CH	20	GLN
35	CH	26	PHE
35	CH	41	VAL
35	CH	47	LYS
35	CH	60	TYR
35	CH	73	ASN
35	CH	79	GLU
35	CH	91	LEU
35	CH	101	ILE
35	CH	131	ILE
35	CH	153	LYS
36	CI	63	TYR
36	CI	69	GLU
36	CI	83	ASP
36	CI	98	LEU
36	CI	100	ASN
37	CJ	8	GLU
37	CJ	12	LEU
37	CJ	20	ASP
37	CJ	63	LYS
37	CJ	72	ARG

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Mol	Chain	Res	Type
37	CJ	104	LEU
37	CJ	137	LYS
37	CJ	155	ARG
38	CK	1	MET
38	CK	10	LEU
38	CK	25	ASP
38	CK	26	VAL
38	CK	41	ARG
38	CK	65	TYR
38	CK	70	GLN
38	CK	82	HIS
38	CK	105	ARG
39	CL	9	ARG
39	CL	47	LEU
39	CL	48	GLU
39	CL	66	ARG
39	CL	75	ASP
39	CL	83	ARG
39	CL	95	LYS
39	CL	102	LEU
39	CL	110	GLU
39	CL	112	LYS
39	CL	113	LYS
39	CL	114	TYR
39	CL	121	ARG
39	CL	128	ARG
40	CM	22	LYS
40	CM	45	ARG
40	CM	47	PHE
40	CM	57	LYS
40	CM	62	HIS
40	CM	74	ILE
40	CM	80	LYS
40	CM	95	GLU
40	CM	96	ILE
41	CN	29	ILE
41	CN	117	ASN
42	CO	20	LYS
42	CO	27	LEU
42	CO	41	ARG
42	CO	47	LYS
42	CO	53	ARG

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Mol	Chain	Res	Type
42	CO	60	LEU
42	CO	62	SER
42	CO	83	VAL
42	CO	85	ILE
42	CO	89	ARG
42	CO	127	GLU
43	CP	3	ARG
43	CP	11	ARG
43	CP	48	LEU
43	CP	56	LEU
43	CP	57	ARG
43	CP	64	TRP
43	CP	66	LEU
43	CP	67	GLU
43	CP	69	GLU
43	CP	70	LEU
43	CP	88	ARG
43	CP	101	GLN
43	CP	108	ARG
43	CP	114	ARG
43	CP	117	VAL
43	CP	120	LYS
43	CP	125	ARG
43	CP	126	LYS
44	CQ	12	ARG
44	CQ	14	PRO
44	CQ	24	CYS
44	CQ	41	ARG
44	CQ	44	LEU
44	CQ	50	LYS
45	CR	3	ILE
45	CR	8	LYS
45	CR	26	GLU
45	CR	39	LEU
45	CR	65	ARG
45	CR	74	ASP
45	CR	77	ARG
46	CS	1	MET
46	CS	4	ILE
46	CS	32	TYR
46	CS	48	TRP
46	CS	55	ARG

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Mol	Chain	Res	Type
46	CS	67	THR
46	CS	69	THR
46	CS	72	ARG
46	CS	75	ARG
46	CS	81	ARG
46	CS	82	GLN
47	CT	12	SER
47	CT	14	LYS
47	CT	35	VAL
47	CT	52	LYS
47	CT	59	ILE
47	CT	74	LEU
48	CU	18	ARG
48	CU	26	LEU
48	CU	29	PHE
48	CU	31	LEU
48	CU	32	ARG
48	CU	44	LEU
48	CU	46	GLU
48	CU	53	ARG
48	CU	54	ARG
48	CU	55	ARG
48	CU	84	LYS
48	CU	88	LYS
49	CV	3	ARG
49	CV	4	SER
49	CV	5	LEU
49	CV	6	LYS
49	CV	7	LYS
49	CV	10	PHE
49	CV	12	ASP
49	CV	15	LEU
49	CV	25	LYS
49	CV	28	LYS
49	CV	29	ARG
49	CV	30	LEU
49	CV	37	ARG
49	CV	41	VAL
49	CV	61	TYR
49	CV	63	THR
49	CV	65	ASN
49	CV	79	THR

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Mol	Chain	Res	Type
49	CV	81	ARG
49	CV	85	LYS
49	CV	86	GLU
49	CV	88	LYS
50	CW	9	ASN
50	CW	24	LEU
50	CW	26	ASN
50	CW	51	GLU
50	CW	75	ASN
50	CW	93	GLU
50	CW	99	LEU
51	CX	6	ARG
3	DD	10	THR
3	DD	17	THR
3	DD	20	ASP
3	DD	26	LYS
3	DD	28	GLU
3	DD	32	SER
3	DD	43	ARG
3	DD	44	ASN
3	DD	46	GLN
3	DD	49	ILE
3	DD	61	LEU
3	DD	64	ILE
3	DD	65	ILE
3	DD	71	ASP
3	DD	95	LEU
3	DD	99	ASP
3	DD	106	ILE
3	DD	131	LEU
3	DD	135	PHE
3	DD	138	VAL
3	DD	157	ARG
3	DD	166	GLN
3	DD	192	THR
3	DD	212	SER
3	DD	217	ARG
3	DD	221	VAL
3	DD	226	MET
3	DD	229	VAL
3	DD	230	ASP
3	DD	237	GLU

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Mol	Chain	Res	Type
3	DD	257	LEU
3	DD	259	THR
3	DD	261	LYS
4	DE	2	LYS
4	DE	4	ILE
4	DE	13	ARG
4	DE	14	ILE
4	DE	16	ARG
4	DE	26	ILE
4	DE	27	LEU
4	DE	37	ARG
4	DE	41	LYS
4	DE	47	VAL
4	DE	49	LEU
4	DE	52	LEU
4	DE	54	GLN
4	DE	62	PRO
4	DE	63	LEU
4	DE	67	PHE
4	DE	73	GLU
4	DE	75	VAL
4	DE	79	ARG
4	DE	82	ARG
4	DE	85	ASN
4	DE	89	ASP
4	DE	98	PRO
4	DE	101	ARG
4	DE	113	PHE
4	DE	117	MET
4	DE	119	ARG
4	DE	144	ARG
4	DE	163	GLU
4	DE	166	THR
4	DE	170	LEU
4	DE	175	VAL
4	DE	178	GLU
4	DE	179	GLU
4	DE	197	ILE
4	DE	202	LYS
4	DE	203	LYS
5	DF	6	VAL
5	DF	7	TYR

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Mol	Chain	Res	Type
5	DF	25	PRO
5	DF	33	LEU
5	DF	45	ARG
5	DF	48	THR
5	DF	60	SER
5	DF	65	TRP
5	DF	66	PRO
5	DF	67	GLN
5	DF	74	ARG
5	DF	78	ILE
5	DF	104	LYS
5	DF	117	ARG
5	DF	127	GLU
5	DF	140	LEU
5	DF	158	THR
5	DF	164	ARG
5	DF	165	ARG
5	DF	169	ASN
5	DF	181	LEU
5	DF	191	ARG
5	DF	192	LEU
5	DF	203	GLN
5	DF	206	ILE
6	DG	16	ARG
6	DG	34	LEU
6	DG	45	GLU
6	DG	49	ASP
6	DG	60	LEU
6	DG	67	LYS
6	DG	71	THR
6	DG	79	ASN
6	DG	88	ILE
6	DG	90	LEU
6	DG	95	ARG
6	DG	97	ASP
6	DG	108	ASN
6	DG	112	PRO
6	DG	115	ARG
6	DG	118	ARG
6	DG	121	ASN
6	DG	139	LEU
6	DG	147	ASP

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Mol	Chain	Res	Type
6	DG	159	VAL
6	DG	174	GLU
7	DH	3	ARG
7	DH	4	ILE
7	DH	7	LEU
7	DH	12	PRO
7	DH	32	GLU
7	DH	41	MET
7	DH	42	ARG
7	DH	50	VAL
7	DH	54	ARG
7	DH	83	TYR
7	DH	85	LYS
7	DH	86	GLU
7	DH	88	LEU
7	DH	89	ILE
7	DH	104	GLU
7	DH	105	LEU
7	DH	124	GLU
7	DH	132	ARG
7	DH	143	GLN
7	DH	152	ARG
7	DH	153	LYS
7	DH	154	PRO
7	DH	155	SER
7	DH	158	HIS
7	DH	160	LYS
56	DI	7	ARG
56	DI	8	ILE
56	DI	11	GLU
56	DI	13	SER
56	DI	17	VAL
56	DI	20	LEU
56	DI	22	GLN
56	DI	23	LEU
56	DI	24	ILE
56	DI	28	LYS
56	DI	29	GLU
8	DK	1	MET
8	DK	2	LYS
8	DK	6	LEU
8	DK	9	LEU

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Mol	Chain	Res	Type
8	DK	10	GLU
8	DK	12	LEU
8	DK	25	TYR
8	DK	28	ASN
8	DK	33	ARG
8	DK	38	LEU
8	DK	41	GLU
8	DK	57	ARG
8	DK	64	GLU
8	DK	70	GLU
8	DK	71	ILE
8	DK	74	ASN
8	DK	77	LEU
8	DK	78	THR
8	DK	85	GLU
8	DK	88	ILE
8	DK	110	ASP
8	DK	113	ARG
8	DK	114	LEU
8	DK	118	LYS
8	DK	120	ILE
8	DK	126	TYR
8	DK	131	LYS
8	DK	134	PRO
8	DK	135	GLU
8	DK	139	GLN
57	DY	1	MET
57	DY	2	PRO
57	DY	5	ARG
57	DY	9	LEU
57	DY	13	LEU
57	DY	14	LYS
57	DY	15	GLU
57	DY	17	LEU
57	DY	19	ARG
57	DY	21	GLN
57	DY	24	PHE
57	DY	25	PHE
57	DY	26	LEU
57	DY	27	VAL
57	DY	29	TYR
57	DY	32	LEU

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Mol	Chain	Res	Type
57	DY	33	PRO
57	DY	35	LYS
57	DY	36	GLU
57	DY	38	HIS
57	DY	41	ARG
57	DY	42	GLN
57	DY	45	LYS
57	DY	46	GLN
57	DY	47	ASN
57	DY	50	ARG
57	DY	52	PHE
57	DY	53	VAL
57	DY	56	ASN
57	DY	58	LEU
57	DY	59	ILE
57	DY	60	ARG
57	DY	61	LEU
57	DY	63	LEU
57	DY	64	LYS
57	DY	68	LEU
57	DY	70	GLU
57	DY	71	LEU
57	DY	72	ASP
57	DY	74	LEU
57	DY	75	GLN
57	DY	82	PHE
57	DY	86	PRO
57	DY	91	LYS
57	DY	92	THR
57	DY	93	LEU
57	DY	98	LYS
57	DY	99	SER
57	DY	100	ASN
57	DY	101	PRO
57	DY	102	LYS
57	DY	104	ILE
57	DY	108	LYS
57	DY	111	LEU
57	DY	112	LEU
57	DY	115	GLN
57	DY	116	ILE
57	DY	117	LEU

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Mol	Chain	Res	Type
57	DY	120	LYS
57	DY	122	VAL
57	DY	128	LEU
57	DY	130	THR
57	DY	131	MET
57	DY	132	ASP
57	DY	133	GLU
57	DY	134	LEU
57	DY	137	GLU
57	DY	138	LEU
57	DY	139	VAL
57	DY	142	LEU
58	DL	5	VAL
58	DL	8	VAL
58	DL	9	LYS
58	DL	10	LEU
58	DL	11	GLN
58	DL	12	LEU
58	DL	13	PRO
58	DL	18	THR
58	DL	27	LEU
58	DL	29	GLN
58	DL	30	HIS
58	DL	36	GLU
58	DL	37	PHE
58	DL	41	PHE
58	DL	45	THR
58	DL	48	MET
58	DL	52	ILE
58	DL	54	PRO
58	DL	55	VAL
58	DL	57	ILE
58	DL	58	THR
58	DL	59	ILE
58	DL	60	TYR
58	DL	63	ARG
58	DL	65	PHE
58	DL	66	THR
58	DL	70	LYS
58	DL	76	TYR
58	DL	80	LYS
58	DL	85	GLU

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Mol	Chain	Res	Type
58	DL	86	LYS
58	DL	89	HIS
58	DL	90	LYS
58	DL	95	LYS
58	DL	96	VAL
58	DL	98	ARG
58	DL	100	THR
58	DL	101	TRP
58	DL	102	GLU
58	DL	103	GLN
58	DL	106	GLU
58	DL	109	LYS
58	DL	111	LYS
58	DL	112	MET
58	DL	113	PRO
58	DL	116	ASN
58	DL	117	THR
58	DL	119	ASP
58	DL	120	LEU
58	DL	121	GLU
58	DL	126	MET
58	DL	127	ILE
58	DL	130	SER
58	DL	137	GLU
58	DL	139	VAL
58	DL	143	GLU
58	DL	146	ASP
56	DJ	1	MET
56	DJ	3	LEU
56	DJ	7	ARG
56	DJ	8	ILE
56	DJ	9	LYS
56	DJ	10	GLU
56	DJ	11	GLU
56	DJ	12	LEU
56	DJ	16	THR
56	DJ	19	GLU
56	DJ	23	LEU
56	DJ	24	ILE
9	DM	2	LYS
9	DM	7	LYS
9	DM	9	VAL

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Mol	Chain	Res	Type
9	DM	35	ARG
9	DM	45	ASN
9	DM	48	MET
9	DM	53	VAL
9	DM	60	ILE
9	DM	61	ARG
9	DM	71	ILE
9	DM	87	LEU
9	DM	90	MET
9	DM	96	GLU
9	DM	127	ASP
9	DM	131	GLN
10	DN	2	ILE
10	DN	8	LEU
10	DN	9	GLU
10	DN	19	ILE
10	DN	20	MET
10	DN	23	ARG
10	DN	24	VAL
10	DN	31	LYS
10	DN	32	TYR
10	DN	38	VAL
10	DN	49	ARG
10	DN	69	ILE
10	DN	77	ILE
10	DN	78	ARG
10	DN	94	ARG
10	DN	98	VAL
11	DO	5	ASP
11	DO	16	ARG
11	DO	19	VAL
11	DO	21	ARG
11	DO	27	HIS
11	DO	29	LYS
11	DO	30	THR
11	DO	42	SER
11	DO	45	LEU
11	DO	50	ARG
11	DO	58	THR
11	DO	59	LEU
11	DO	61	ARG
11	DO	62	LEU

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Mol	Chain	Res	Type
11	DO	64	LYS
11	DO	65	ARG
11	DO	67	MET
11	DO	68	GLN
11	DO	75	ILE
11	DO	76	LYS
11	DO	81	GLN
11	DO	88	LEU
11	DO	90	ARG
11	DO	101	VAL
11	DO	105	LEU
11	DO	106	LEU
11	DO	114	ILE
11	DO	119	GLU
11	DO	126	VAL
11	DO	135	LEU
11	DO	138	LEU
11	DO	144	GLU
12	DP	10	ARG
12	DP	21	THR
12	DP	26	TYR
12	DP	45	GLN
12	DP	55	VAL
12	DP	59	ARG
12	DP	79	LEU
12	DP	81	VAL
12	DP	82	ARG
12	DP	83	MET
12	DP	96	VAL
12	DP	99	PRO
12	DP	109	VAL
12	DP	132	VAL
13	D0	6	SER
13	D0	9	LYS
13	D0	10	LEU
13	D0	16	HIS
13	D0	18	LEU
13	D0	29	LEU
13	D0	44	LEU
13	D0	54	LEU
13	D0	56	LYS
13	D0	57	ARG

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Mol	Chain	Res	Type
13	D0	60	LEU
13	D0	71	GLN
13	D0	79	LEU
13	D0	91	GLN
13	D0	96	ARG
13	D0	104	ARG
13	D0	105	ARG
13	D0	118	GLU
14	DQ	12	PHE
14	DQ	17	ARG
14	DQ	19	LYS
14	DQ	20	ARG
14	DQ	44	LYS
14	DQ	52	SER
14	DQ	56	LEU
14	DQ	58	LEU
14	DQ	69	VAL
14	DQ	89	ARG
14	DQ	103	GLU
14	DQ	106	ARG
14	DQ	111	GLU
15	DR	11	GLU
15	DR	14	TYR
15	DR	23	ARG
15	DR	26	ASP
15	DR	27	THR
15	DR	30	VAL
15	DR	42	ILE
15	DR	58	ASN
15	DR	62	THR
15	DR	65	LYS
15	DR	74	ARG
15	DR	78	LEU
15	DR	84	GLN
15	DR	86	ILE
15	DR	88	ILE
15	DR	90	GLN
15	DR	95	ARG
15	DR	99	LEU
15	DR	105	LEU
15	DR	106	SER
15	DR	111	ARG

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Mol	Chain	Res	Type
15	DR	112	ARG
15	DR	125	ARG
15	DR	128	GLU
16	D1	5	LYS
16	D1	8	VAL
16	D1	19	LYS
16	D1	27	LEU
16	D1	31	SER
16	D1	44	ASN
16	D1	60	LEU
16	D1	74	LEU
16	D1	76	TYR
16	D1	78	THR
16	D1	83	LEU
16	D1	88	ILE
16	D1	92	ARG
16	D1	98	LEU
16	D1	108	GLU
16	D1	111	GLU
17	D2	14	VAL
17	D2	19	LYS
17	D2	21	ARG
17	D2	32	THR
17	D2	35	LEU
17	D2	36	PRO
17	D2	38	LEU
17	D2	39	LEU
17	D2	44	LYS
17	D2	49	THR
17	D2	57	VAL
17	D2	64	HIS
17	D2	66	ARG
17	D2	73	SER
17	D2	89	GLN
17	D2	91	TYR
17	D2	95	LEU
17	D2	99	ILE
18	DS	1	MET
18	DS	11	ARG
18	DS	16	LYS
18	DS	42	ARG
18	DS	65	LEU

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Mol	Chain	Res	Type
18	DS	67	ASP
18	DS	69	LEU
18	DS	70	TYR
18	DS	76	VAL
18	DS	88	ARG
18	DS	96	ILE
18	DS	99	ARG
18	DS	107	LEU
19	DT	27	THR
19	DT	28	PHE
19	DT	41	ASN
19	DT	57	LEU
19	DT	65	ARG
19	DT	76	ARG
19	DT	81	VAL
19	DT	88	LYS
20	DU	14	LEU
20	DU	19	LYS
20	DU	26	LYS
20	DU	27	VAL
20	DU	34	LYS
20	DU	45	VAL
20	DU	51	VAL
20	DU	57	GLN
20	DU	61	ILE
20	DU	71	LYS
20	DU	75	ILE
20	DU	77	PRO
20	DU	86	ARG
20	DU	87	LYS
20	DU	89	PHE
20	DU	90	LEU
20	DU	95	LYS
20	DU	97	ARG
20	DU	98	VAL
21	DV	5	LEU
21	DV	6	LYS
21	DV	11	GLU
21	DV	14	LYS
21	DV	19	ARG
21	DV	38	TYR
21	DV	53	ILE

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Mol	Chain	Res	Type
21	DV	61	LEU
21	DV	76	LEU
21	DV	77	ASP
21	DV	80	ARG
21	DV	81	ARG
21	DV	87	ASP
21	DV	93	ASP
21	DV	97	GLU
21	DV	105	VAL
21	DV	111	VAL
21	DV	112	ARG
21	DV	117	LEU
21	DV	119	GLU
21	DV	122	ARG
21	DV	141	VAL
21	DV	142	SER
21	DV	148	ASP
21	DV	150	LEU
21	DV	151	HIS
21	DV	153	SER
21	DV	154	ASP
21	DV	162	GLU
21	DV	169	GLU
21	DV	177	PRO
21	DV	179	ASP
21	DV	180	VAL
21	DV	182	LYS
21	DV	183	LEU
21	DV	185	GLU
21	DV	186	GLU
21	DV	193	GLU
21	DV	196	VAL
21	DV	197	ILE
21	DV	198	LYS
21	DV	199	LYS
22	D3	10	THR
22	D3	11	ARG
22	D3	14	ARG
22	D3	36	ILE
22	D3	41	ARG
22	D3	55	ARG
22	D3	64	ASP

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Mol	Chain	Res	Type
23	DZ	41	ARG
23	DZ	51	VAL
23	DZ	56	GLN
23	DZ	57	GLU
23	DZ	76	ARG
23	DZ	78	LYS
23	DZ	80	LEU
23	DZ	81	LYS
23	DZ	83	GLU
23	DZ	91	LYS
23	DZ	95	LEU
24	DW	5	GLU
24	DW	7	ARG
24	DW	15	LYS
24	DW	16	LEU
24	DW	22	GLU
24	DW	32	LEU
24	DW	35	LEU
24	DW	47	ASN
24	DW	52	ASP
24	DW	53	LEU
24	DW	64	LEU
25	DX	8	LEU
25	DX	17	LYS
25	DX	38	GLU
26	D4	15	ILE
26	D4	16	CYS
26	D4	23	GLU
26	D4	32	TYR
26	D4	33	VAL
26	D4	36	CYS
26	D4	38	LYS
26	D4	42	PHE
26	D4	49	PHE
26	D4	55	ARG
26	D4	59	PHE
26	D4	61	ARG
26	D4	68	ARG
26	D4	69	LYS
27	D5	3	LYS
27	D5	4	HIS
27	D5	6	VAL

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Mol	Chain	Res	Type
27	D5	25	LEU
27	D5	26	THR
27	D5	29	THR
27	D5	30	LEU
27	D5	36	CYS
27	D5	37	LYS
27	D5	40	LYS
27	D5	51	TYR
27	D5	52	TYR
27	D5	56	LYS
27	D5	57	VAL
27	D5	58	LEU
28	D6	9	LEU
28	D6	10	LEU
28	D6	11	LEU
28	D6	17	LYS
28	D6	19	ARG
28	D6	24	GLU
28	D6	27	LYS
28	D6	31	PRO
28	D6	34	LEU
28	D6	36	LEU
28	D6	37	ARG
28	D6	38	LYS
28	D6	39	TYR
28	D6	40	CYS
28	D6	42	TRP
28	D6	44	ARG
29	D7	4	THR
29	D7	8	ASN
29	D7	24	THR
29	D7	30	VAL
30	D8	8	LYS
30	D8	14	VAL
30	D8	15	LYS
30	D8	17	THR
30	D8	30	ARG
30	D8	35	GLN
30	D8	43	GLN
30	D8	44	LYS
30	D8	47	LYS
30	D8	48	PHE

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Mol	Chain	Res	Type
30	D8	49	VAL
30	D8	52	LYS
30	D8	56	GLU
30	D8	63	PRO
30	D8	65	GLU

Some sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (317) such sidechains are listed below:

Mol	Chain	Res	Type
3	AD	44	ASN
3	AD	58	HIS
3	AD	116	GLN
3	AD	126	GLN
3	AD	166	GLN
3	AD	186	HIS
3	AD	198	ASN
3	AD	201	HIS
4	AE	48	GLN
4	AE	55	ASN
4	AE	60	ASN
4	AE	66	HIS
4	AE	192	ASN
5	AF	8	GLN
5	AF	75	HIS
5	AF	169	ASN
6	AG	26	GLN
6	AG	40	ASN
6	AG	41	GLN
6	AG	108	ASN
6	AG	121	ASN
6	AG	123	ASN
7	AH	143	GLN
7	AH	147	ASN
8	AK	43	ASN
8	AK	54	GLN
8	AK	105	HIS
9	AM	45	ASN
9	AM	94	HIS
9	AM	131	GLN
9	AM	133	GLN
10	AN	3	GLN
10	AN	5	GLN

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Mol	Chain	Res	Type
10	AN	82	ASN
11	AO	9	ASN
11	AO	84	ASN
12	AP	12	GLN
12	AP	89	ASN
12	AP	113	GLN
12	AP	141	GLN
13	A0	11	ASN
13	A0	23	ASN
13	A0	24	GLN
13	A0	61	HIS
13	A0	71	GLN
13	A0	91	GLN
15	AR	58	ASN
15	AR	90	GLN
15	AR	136	GLN
16	A1	49	HIS
16	A1	66	ASN
16	A1	72	HIS
16	A1	104	GLN
17	A2	11	GLN
17	A2	80	GLN
18	AS	34	ASN
18	AS	40	ASN
18	AS	57	ASN
18	AS	102	HIS
19	AT	41	ASN
19	AT	55	ASN
19	AT	87	GLN
20	AU	43	ASN
20	AU	57	GLN
21	AV	32	HIS
21	AV	34	ASN
21	AV	55	HIS
21	AV	75	ASN
21	AV	132	ASN
22	A3	29	GLN
22	A3	35	ASN
22	A3	40	GLN
22	A3	70	GLN
23	AZ	56	GLN
23	AZ	66	HIS

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Mol	Chain	Res	Type
24	AW	56	GLN
24	AW	65	ASN
25	AX	19	GLN
25	AX	33	GLN
25	AX	46	ASN
25	AX	52	HIS
26	A4	40	HIS
26	A4	46	GLN
26	A4	60	GLN
27	A5	23	HIS
27	A5	43	HIS
28	A6	29	ASN
28	A6	32	ASN
29	A7	8	ASN
32	BE	37	ASN
32	BE	40	HIS
32	BE	94	ASN
32	BE	135	GLN
32	BE	146	GLN
32	BE	212	GLN
33	BF	69	HIS
33	BF	98	ASN
33	BF	110	ASN
33	BF	123	GLN
33	BF	136	GLN
33	BF	162	GLN
33	BF	170	GLN
33	BF	181	ASN
34	BG	42	GLN
34	BG	45	GLN
34	BG	62	GLN
34	BG	119	GLN
34	BG	123	HIS
34	BG	160	GLN
35	BH	20	GLN
35	BH	65	ASN
35	BH	141	GLN
36	BI	7	ASN
36	BI	27	GLN
36	BI	32	ASN
36	BI	57	GLN
36	BI	73	ASN

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Mol	Chain	Res	Type
36	BI	100	ASN
37	BJ	13	GLN
37	BJ	28	ASN
37	BJ	97	GLN
37	BJ	106	GLN
38	BK	15	ASN
38	BK	82	HIS
39	BL	3	GLN
39	BL	58	HIS
39	BL	124	GLN
40	BM	56	HIS
40	BM	68	HIS
40	BM	78	ASN
40	BM	84	GLN
41	BN	38	ASN
41	BN	93	GLN
42	BO	8	ASN
42	BO	9	GLN
42	BO	49	ASN
42	BO	75	HIS
43	BP	77	ASN
43	BP	92	HIS
43	BP	101	GLN
44	BQ	49	HIS
45	BR	13	GLN
45	BR	37	ASN
45	BR	46	HIS
46	BS	76	GLN
47	BT	16	GLN
47	BT	26	GLN
49	BV	14	HIS
49	BV	23	ASN
49	BV	47	HIS
49	BV	56	GLN
50	BW	18	GLN
50	BW	26	ASN
32	CE	37	ASN
32	CE	40	HIS
32	CE	76	GLN
32	CE	78	GLN
32	CE	94	ASN
32	CE	95	GLN

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Mol	Chain	Res	Type
32	CE	110	GLN
32	CE	146	GLN
32	CE	204	ASN
32	CE	212	GLN
33	CF	28	GLN
33	CF	31	HIS
33	CF	69	HIS
33	CF	170	GLN
33	CF	176	HIS
33	CF	181	ASN
34	CG	77	ASN
34	CG	119	GLN
34	CG	160	GLN
34	CG	161	ASN
34	CG	201	GLN
35	CH	20	GLN
35	CH	72	GLN
35	CH	73	ASN
35	CH	78	HIS
36	CI	18	GLN
36	CI	27	GLN
36	CI	32	ASN
36	CI	64	GLN
37	CJ	37	ASN
37	CJ	86	GLN
37	CJ	106	GLN
37	CJ	122	HIS
38	CK	15	ASN
39	CL	58	HIS
39	CL	124	GLN
40	CM	68	HIS
40	CM	78	ASN
41	CN	13	GLN
41	CN	117	ASN
42	CO	9	GLN
42	CO	49	ASN
43	CP	62	ASN
43	CP	77	ASN
43	CP	92	HIS
43	CP	101	GLN
44	CQ	49	HIS
45	CR	9	GLN

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Mol	Chain	Res	Type
45	CR	37	ASN
46	CS	14	ASN
46	CS	76	GLN
46	CS	82	GLN
47	CT	16	GLN
47	CT	94	ASN
49	CV	23	ASN
49	CV	47	HIS
49	CV	65	ASN
50	CW	9	ASN
50	CW	18	GLN
50	CW	26	ASN
50	CW	90	GLN
3	DD	58	HIS
3	DD	96	HIS
3	DD	116	GLN
3	DD	126	GLN
3	DD	166	GLN
3	DD	186	HIS
3	DD	198	ASN
3	DD	201	HIS
3	DD	220	HIS
3	DD	227	ASN
4	DE	35	GLN
4	DE	135	HIS
4	DE	143	ASN
4	DE	159	HIS
4	DE	192	ASN
5	DF	40	GLN
5	DF	67	GLN
5	DF	69	HIS
5	DF	133	ASN
5	DF	169	ASN
5	DF	203	GLN
5	DF	204	ASN
6	DG	26	GLN
6	DG	41	GLN
6	DG	58	GLN
6	DG	79	ASN
6	DG	121	ASN
6	DG	123	ASN
7	DH	139	GLN

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Mol	Chain	Res	Type
7	DH	147	ASN
8	DK	43	ASN
8	DK	74	ASN
8	DK	105	HIS
8	DK	139	GLN
57	DY	3	ASN
57	DY	21	GLN
57	DY	28	ASN
57	DY	46	GLN
57	DY	47	ASN
57	DY	95	GLN
57	DY	113	GLN
9	DM	45	ASN
9	DM	69	GLN
9	DM	94	HIS
9	DM	128	HIS
9	DM	130	HIS
9	DM	131	GLN
10	DN	82	ASN
11	DO	9	ASN
11	DO	68	GLN
11	DO	81	GLN
12	DP	12	GLN
12	DP	89	ASN
12	DP	141	GLN
13	D0	3	HIS
13	D0	13	HIS
13	D0	16	HIS
13	D0	23	ASN
13	D0	24	GLN
13	D0	53	HIS
13	D0	61	HIS
13	D0	91	GLN
14	DQ	34	HIS
15	DR	43	GLN
15	DR	55	ASN
15	DR	84	GLN
15	DR	90	GLN
15	DR	136	GLN
16	D1	49	HIS
16	D1	71	GLN
16	D1	72	HIS

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Mol	Chain	Res	Type
16	D1	94	ASN
17	D2	11	GLN
17	D2	89	GLN
18	DS	57	ASN
18	DS	60	ASN
18	DS	61	ASN
18	DS	102	HIS
19	DT	31	HIS
19	DT	41	ASN
19	DT	55	ASN
19	DT	87	GLN
20	DU	57	GLN
21	DV	32	HIS
21	DV	55	HIS
21	DV	75	ASN
22	D3	29	GLN
22	D3	35	ASN
23	DZ	56	GLN
23	DZ	66	HIS
24	DW	56	GLN
24	DW	65	ASN
25	DX	19	GLN
25	DX	46	ASN
25	DX	52	HIS
26	D4	6	HIS
26	D4	47	GLN
26	D4	60	GLN
27	D5	4	HIS
27	D5	43	HIS
28	D6	20	ASN
29	D7	8	ASN
29	D7	36	GLN

5.3.3 RNA [i](#)

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	AA	2908/2909 (99%)	771 (26%)	318 (10%)
2	AB	121/122 (99%)	24 (19%)	6 (4%)
2	DB	121/122 (99%)	27 (22%)	5 (4%)
31	BA	1516/1516 (100%)	331 (21%)	130 (8%)
52	BB	74/76 (97%)	28 (37%)	8 (10%)

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Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
52	BC	74/76 (97%)	19 (25%)	6 (8%)
52	BD	74/76 (97%)	15 (20%)	3 (4%)
52	CB	74/76 (97%)	25 (33%)	8 (10%)
52	CC	74/76 (97%)	18 (24%)	7 (9%)
52	CD	74/76 (97%)	23 (31%)	5 (6%)
53	B1	29/30 (96%)	12 (41%)	4 (13%)
53	C1	29/30 (96%)	10 (34%)	4 (13%)
54	CA	1514/1515 (99%)	321 (21%)	138 (9%)
55	DA	2911/2912 (99%)	806 (27%)	374 (12%)
All	All	9593/9612 (99%)	2430 (25%)	1016 (10%)

All (2430) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	AA	5	A
1	AA	13	A
1	AA	14	A
1	AA	28	A
1	AA	34	C
1	AA	35	G
1	AA	46	C
1	AA	49	A
1	AA	50	U
1	AA	51	G
1	AA	60	G
1	AA	61	G
1	AA	69	C
1	AA	70	G
1	AA	71	A
1	AA	72	U
1	AA	73	A
1	AA	74	A
1	AA	75	G
1	AA	83	G
1	AA	84	A
1	AA	85	G
1	AA	88	G
1	AA	90	U
1	AA	91	A
1	AA	93	C
1	AA	99	U
1	AA	101	G

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Mol	Chain	Res	Type
1	AA	102	G
1	AA	119	A
1	AA	120	U
1	AA	121	G
1	AA	126	A
1	AA	129	C
1	AA	137	C
1	AA	138	G
1	AA	140	A
1	AA	155	C
1	AA	161	U
1	AA	162	U
1	AA	171	G
1	AA	174	C
1	AA	175	G
1	AA	178	G
1	AA	196	A
1	AA	204	A
1	AA	205	G
1	AA	206	U
1	AA	215	G
1	AA	216	A
1	AA	221	A
1	AA	222	A
1	AA	223	A
1	AA	228	A
1	AA	229	A
1	AA	232	G
1	AA	233	A
1	AA	241	A
1	AA	242	G
1	AA	243	U
1	AA	248	G
1	AA	249	C
1	AA	250	G
1	AA	261	G
1	AA	265	A
1	AA	266	G
1	AA	267	C
1	AA	270(B)	A
1	AA	270(K)	C
1	AA	270(L)	U

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Mol	Chain	Res	Type
1	AA	270(M)	U
1	AA	270(N)	G
1	AA	270(O)	U
1	AA	270(P)	C
1	AA	270(Z)	U
1	AA	271(C)	U
1	AA	271	G
1	AA	273(D)	C
1	AA	274	G
1	AA	278	A
1	AA	279	C
1	AA	283	A
1	AA	284	U
1	AA	288	C
1	AA	289	A
1	AA	301	G
1	AA	311	A
1	AA	312	G
1	AA	316	C
1	AA	322	A
1	AA	323	G
1	AA	324	A
1	AA	329	G
1	AA	330	A
1	AA	332	A
1	AA	333	G
1	AA	346	A
1	AA	352	G
1	AA	353	G
1	AA	356	G
1	AA	363	G
1	AA	363(A)	A
1	AA	363(B)	G
1	AA	363(E)	U
1	AA	363(F)	A
1	AA	364	C
1	AA	371	A
1	AA	372	G
1	AA	385	C
1	AA	386	G
1	AA	387	U
1	AA	388	G

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Mol	Chain	Res	Type
1	AA	390	A
1	AA	391	G
1	AA	395	U
1	AA	396	G
1	AA	404	C
1	AA	405	U
1	AA	406	G
1	AA	411	G
1	AA	412	A
1	AA	422	A
1	AA	428	A
1	AA	435	C
1	AA	442	G
1	AA	443	A
1	AA	444	C
1	AA	446	G
1	AA	448	U
1	AA	449	A
1	AA	455	C
1	AA	456	C
1	AA	457	A
1	AA	458	G
1	AA	459	U
1	AA	470	A
1	AA	475	U
1	AA	480	A
1	AA	481	G
1	AA	482	A
1	AA	494	G
1	AA	504	U
1	AA	505	A
1	AA	507	A
1	AA	508	G
1	AA	509	C
1	AA	512	G
1	AA	527	C
1	AA	530	G
1	AA	531	C
1	AA	533	G
1	AA	563	G
1	AA	573	G
1	AA	574	C

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Mol	Chain	Res	Type
1	AA	575	A
1	AA	588	U
1	AA	604	G
1	AA	607	U
1	AA	612	G
1	AA	614	U
1	AA	615	G
1	AA	616	A
1	AA	617	G
1	AA	620	G
1	AA	621	A
1	AA	622	G
1	AA	627	A
1	AA	628	G
1	AA	638	G
1	AA	645	C
1	AA	646	A
1	AA	654	A
1	AA	654(E)	C
1	AA	654(F)	C
1	AA	654(G)	C
1	AA	654(H)	G
1	AA	654(I)	C
1	AA	654(J)	A
1	AA	654(K)	C
1	AA	654(L)	G
1	AA	654(N)	G
1	AA	654(S)	G
1	AA	654(T)	A
1	AA	657	U
1	AA	666	G
1	AA	668	G
1	AA	670	A
1	AA	671	C
1	AA	686	G
1	AA	687	C
1	AA	705	A
1	AA	707	G
1	AA	708	C
1	AA	722	A
1	AA	727	A
1	AA	730	C

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Mol	Chain	Res	Type
1	AA	739	G
1	AA	740	U
1	AA	746	A
1	AA	747	U
1	AA	753	C
1	AA	763	G
1	AA	775	G
1	AA	776	G
1	AA	777	A
1	AA	782	A
1	AA	784	A
1	AA	785	G
1	AA	789	A
1	AA	790	C
1	AA	791	C
1	AA	792	G
1	AA	793	A
1	AA	800	A
1	AA	801	G
1	AA	802	A
1	AA	805	G
1	AA	806	C
1	AA	811	U
1	AA	812	C
1	AA	819	A
1	AA	827	U
1	AA	828	U
1	AA	830	G
1	AA	831	G
1	AA	847	U
1	AA	848	G
1	AA	857	C
1	AA	859	G
1	AA	860	U
1	AA	866	A
1	AA	878	A
1	AA	882	G
1	AA	883	G
1	AA	884	C
1	AA	886	C
1	AA	888	C
1	AA	889	C

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Mol	Chain	Res	Type
1	AA	894	C
1	AA	895	U
1	AA	896	A
1	AA	897	C
1	AA	898	C
1	AA	899	A
1	AA	901	A
1	AA	910	A
1	AA	917	A
1	AA	918	A
1	AA	919	G
1	AA	926	A
1	AA	931	G
1	AA	932	G
1	AA	933	A
1	AA	941	A
1	AA	944	G
1	AA	945	A
1	AA	946	G
1	AA	947	G
1	AA	955	C
1	AA	956	G
1	AA	957	A
1	AA	958	U
1	AA	959	A
1	AA	961	C
1	AA	962	G
1	AA	973	A
1	AA	974	G
1	AA	983	A
1	AA	986	C
1	AA	989	G
1	AA	990	A
1	AA	991	C
1	AA	996	A
1	AA	1000	A
1	AA	1008	C
1	AA	1009	A
1	AA	1011	G
1	AA	1012	U
1	AA	1013	C
1	AA	1020	A

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Mol	Chain	Res	Type
1	AA	1021	A
1	AA	1022	G
1	AA	1023	U
1	AA	1025	G
1	AA	1026	U
1	AA	1027	A
1	AA	1044	G
1	AA	1045	A
1	AA	1046	A
1	AA	1047	G
1	AA	1048	A
1	AA	1054	A
1	AA	1056	G
1	AA	1060	U
1	AA	1061	U
1	AA	1070	A
1	AA	1071	G
1	AA	1086	A
1	AA	1087	G
1	AA	1088	A
1	AA	1090	U
1	AA	1095	A
1	AA	1096	A
1	AA	1097	U
1	AA	1099	G
1	AA	1103	A
1	AA	1112	G
1	AA	1126	A
1	AA	1127	A
1	AA	1130	U
1	AA	1131	G
1	AA	1135	C
1	AA	1136	G
1	AA	1142	U
1	AA	1142(A)	A
1	AA	1144	G
1	AA	1155	A
1	AA	1156	A
1	AA	1157	G
1	AA	1170	G
1	AA	1171	G
1	AA	1173	G

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Mol	Chain	Res	Type
1	AA	1174	A
1	AA	1175	U
1	AA	1176	G
1	AA	1177	A
1	AA	1178	C
1	AA	1180	C
1	AA	1195	G
1	AA	1204	A
1	AA	1205	U
1	AA	1206	G
1	AA	1211	U
1	AA	1212	G
1	AA	1213	A
1	AA	1220	A
1	AA	1221	C
1	AA	1225	C
1	AA	1236	G
1	AA	1237	A
1	AA	1238	G
1	AA	1244	G
1	AA	1247	A
1	AA	1248	G
1	AA	1249	U
1	AA	1251	C
1	AA	1252	G
1	AA	1253	A
1	AA	1254	A
1	AA	1255	U
1	AA	1256	G
1	AA	1265	A
1	AA	1266	G
1	AA	1271	G
1	AA	1272	A
1	AA	1273	U
1	AA	1276	A
1	AA	1281	G
1	AA	1289	C
1	AA	1300	U
1	AA	1301	A
1	AA	1302	A
1	AA	1303	G
1	AA	1313	U

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Mol	Chain	Res	Type
1	AA	1314	C
1	AA	1319	G
1	AA	1321	A
1	AA	1325	G
1	AA	1326	U
1	AA	1329	U
1	AA	1330	C
1	AA	1332	G
1	AA	1333	C
1	AA	1341	U
1	AA	1342	A
1	AA	1343	G
1	AA	1344	G
1	AA	1345	C
1	AA	1349	A
1	AA	1359	A
1	AA	1367	A
1	AA	1372	U
1	AA	1379	A
1	AA	1380	G
1	AA	1384	A
1	AA	1385	G
1	AA	1396	U
1	AA	1397	U
1	AA	1398	C
1	AA	1407	C
1	AA	1416	G
1	AA	1419	A
1	AA	1420	U
1	AA	1427	A
1	AA	1428	C
1	AA	1444(A)	A
1	AA	1449	A
1	AA	1449(A)	G
1	AA	1451	C
1	AA	1453	A
1	AA	1454	U
1	AA	1455	G
1	AA	1459	G
1	AA	1461	G
1	AA	1467	C
1	AA	1471	A

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Mol	Chain	Res	Type
1	AA	1475	G
1	AA	1478	G
1	AA	1482	U
1	AA	1483	G
1	AA	1485	G
1	AA	1488	G
1	AA	1490	A
1	AA	1491	G
1	AA	1493	C
1	AA	1494	A
1	AA	1497	U
1	AA	1498	C
1	AA	1502	C
1	AA	1505	C
1	AA	1508	A
1	AA	1509	C
1	AA	1510	A
1	AA	1522	G
1	AA	1534	G
1	AA	1535	U
1	AA	1537	C
1	AA	1538	G
1	AA	1543	A
1	AA	1544	C
1	AA	1545	A
1	AA	1554	A
1	AA	1555	G
1	AA	1558	A
1	AA	1559	G
1	AA	1560	G
1	AA	1565	C
1	AA	1566	A
1	AA	1567	A
1	AA	1569	A
1	AA	1578	U
1	AA	1579	A
1	AA	1585	C
1	AA	1586	A
1	AA	1588	C
1	AA	1602	U
1	AA	1603	A
1	AA	1607	C

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Mol	Chain	Res	Type
1	AA	1608	A
1	AA	1611	C
1	AA	1616	A
1	AA	1617	C
1	AA	1618	A
1	AA	1619	G
1	AA	1632	A
1	AA	1635	G
1	AA	1640	C
1	AA	1647	G
1	AA	1648	C
1	AA	1653	G
1	AA	1654	A
1	AA	1667	G
1	AA	1668	A
1	AA	1669	A
1	AA	1674	G
1	AA	1675	C
1	AA	1681	G
1	AA	1694	C
1	AA	1695	G
1	AA	1696	G
1	AA	1698	A
1	AA	1699	G
1	AA	1700	A
1	AA	1701	A
1	AA	1706	U
1	AA	1707	G
1	AA	1725	G
1	AA	1729	A
1	AA	1731	G
1	AA	1735	C
1	AA	1756	G
1	AA	1758	G
1	AA	1759	A
1	AA	1761	C
1	AA	1762	A
1	AA	1763	G
1	AA	1764	G
1	AA	1773	A
1	AA	1780	A
1	AA	1781	C

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Mol	Chain	Res	Type
1	AA	1782	C
1	AA	1784	A
1	AA	1785	A
1	AA	1787	A
1	AA	1791	A
1	AA	1800	C
1	AA	1801	G
1	AA	1802	A
1	AA	1815	A
1	AA	1816	G
1	AA	1820	U
1	AA	1821	A
1	AA	1822	G
1	AA	1827	C
1	AA	1828	G
1	AA	1829	A
1	AA	1835	G
1	AA	1839	G
1	AA	1847	A
1	AA	1848	A
1	AA	1858	G
1	AA	1869	G
1	AA	1870	C
1	AA	1878	G
1	AA	1880	C
1	AA	1882	C
1	AA	1888	G
1	AA	1889	A
1	AA	1900	A
1	AA	1906	G
1	AA	1914	C
1	AA	1917	U
1	AA	1919	A
1	AA	1925	C
1	AA	1929	G
1	AA	1930	G
1	AA	1931	U
1	AA	1937	A
1	AA	1938	A
1	AA	1939	U
1	AA	1940	U
1	AA	1941	C

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Mol	Chain	Res	Type
1	AA	1943	U
1	AA	1944	U
1	AA	1948	G
1	AA	1955	U
1	AA	1956	U
1	AA	1963	U
1	AA	1964	G
1	AA	1965	C
1	AA	1966	A
1	AA	1967	C
1	AA	1970	A
1	AA	1971	A
1	AA	1972	A
1	AA	1980	G
1	AA	1981	A
1	AA	1982	C
1	AA	1987	G
1	AA	1993	U
1	AA	1996	C
1	AA	1997	G
1	AA	2019	A
1	AA	2022	U
1	AA	2023	G
1	AA	2027	G
1	AA	2031	A
1	AA	2032	G
1	AA	2033	A
1	AA	2034	U
1	AA	2036	C
1	AA	2043	C
1	AA	2051	A
1	AA	2052	G
1	AA	2055	C
1	AA	2056	G
1	AA	2059	A
1	AA	2060	A
1	AA	2061	G
1	AA	2062	A
1	AA	2063	C
1	AA	2067	G
1	AA	2068	U
1	AA	2069	G

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Mol	Chain	Res	Type
1	AA	2092	U
1	AA	2093	G
1	AA	2099	U
1	AA	2108	C
1	AA	2111	C
1	AA	2112	G
1	AA	2113	U
1	AA	2114	A
1	AA	2115	G
1	AA	2118	U
1	AA	2119	A
1	AA	2120	G
1	AA	2126	A
1	AA	2127	G
1	AA	2128	C
1	AA	2130	U
1	AA	2132	U
1	AA	2133	G
1	AA	2136	C
1	AA	2146	C
1	AA	2147	G
1	AA	2159	G
1	AA	2166	G
1	AA	2168	G
1	AA	2172	U
1	AA	2173	A
1	AA	2190	G
1	AA	2192	G
1	AA	2193	G
1	AA	2198	A
1	AA	2210	G
1	AA	2211	G
1	AA	2212	A
1	AA	2215	G
1	AA	2226	C
1	AA	2239	G
1	AA	2249	U
1	AA	2250	G
1	AA	2251	G
1	AA	2259	G
1	AA	2266	A
1	AA	2267	A

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Mol	Chain	Res	Type
1	AA	2268	A
1	AA	2275	C
1	AA	2276	G
1	AA	2279	G
1	AA	2283	C
1	AA	2289	G
1	AA	2297	C
1	AA	2307	G
1	AA	2308	G
1	AA	2311	A
1	AA	2312	U
1	AA	2319	G
1	AA	2320	A
1	AA	2322	A
1	AA	2325	G
1	AA	2334	G
1	AA	2335	A
1	AA	2337	G
1	AA	2345	G
1	AA	2346	A
1	AA	2347	C
1	AA	2349	G
1	AA	2350	C
1	AA	2354	G
1	AA	2383	G
1	AA	2384	G
1	AA	2385	C
1	AA	2388	A
1	AA	2390	U
1	AA	2392	A
1	AA	2400	G
1	AA	2402	C
1	AA	2406	U
1	AA	2407	G
1	AA	2423	U
1	AA	2424	C
1	AA	2425	A
1	AA	2426	A
1	AA	2427	C
1	AA	2428	G
1	AA	2429	G
1	AA	2430	A

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Mol	Chain	Res	Type
1	AA	2434	A
1	AA	2435	A
1	AA	2439	A
1	AA	2440	C
1	AA	2441	C
1	AA	2447	G
1	AA	2448	A
1	AA	2449	U
1	AA	2450	A
1	AA	2458	G
1	AA	2459	A
1	AA	2469	A
1	AA	2470	G
1	AA	2476	A
1	AA	2478	A
1	AA	2482	G
1	AA	2484	G
1	AA	2490	G
1	AA	2491	U
1	AA	2497	A
1	AA	2498	C
1	AA	2502	G
1	AA	2503	A
1	AA	2504	U
1	AA	2505	G
1	AA	2506	U
1	AA	2507	C
1	AA	2518	A
1	AA	2519	U
1	AA	2520	C
1	AA	2523	G
1	AA	2529	G
1	AA	2531	A
1	AA	2534	A
1	AA	2543	G
1	AA	2552	U
1	AA	2554	U
1	AA	2566	A
1	AA	2567	G
1	AA	2573	C
1	AA	2574	G
1	AA	2581	G

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Mol	Chain	Res	Type
1	AA	2582	G
1	AA	2585	U
1	AA	2586	C
1	AA	2602	A
1	AA	2603	G
1	AA	2609	U
1	AA	2610	C
1	AA	2611	U
1	AA	2612	C
1	AA	2613	U
1	AA	2614	A
1	AA	2615	U
1	AA	2629	A
1	AA	2630	G
1	AA	2637	U
1	AA	2645	G
1	AA	2646	C
1	AA	2654	A
1	AA	2655	G
1	AA	2656	U
1	AA	2665	A
1	AA	2673	G
1	AA	2682	U
1	AA	2690	C
1	AA	2691	C
1	AA	2702	U
1	AA	2703	C
1	AA	2712	U
1	AA	2712(A)	A
1	AA	2713	A
1	AA	2714	G
1	AA	2725	A
1	AA	2733	A
1	AA	2748	A
1	AA	2750	A
1	AA	2751	G
1	AA	2752	C
1	AA	2754	U
1	AA	2756	U
1	AA	2757	A
1	AA	2759	G
1	AA	2762	G

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Mol	Chain	Res	Type
1	AA	2763	G
1	AA	2765	A
1	AA	2766	G
1	AA	2777	G
1	AA	2778	A
1	AA	2779	U
1	AA	2780	G
1	AA	2781	A
1	AA	2790	A
1	AA	2791	C
1	AA	2797	U
1	AA	2799	A
1	AA	2808	U
1	AA	2820	A
1	AA	2833	G
1	AA	2834	G
1	AA	2836	U
1	AA	2848	G
1	AA	2849	U
1	AA	2850	A
1	AA	2860	A
1	AA	2866	U
1	AA	2867	G
1	AA	2868	A
1	AA	2872	G
1	AA	2874	C
1	AA	2879	C
1	AA	2880	C
1	AA	2898	U
2	AB	8	U
2	AB	12	C
2	AB	13	A
2	AB	14	U
2	AB	15	A
2	AB	16	G
2	AB	25	A
2	AB	31	C
2	AB	35	U
2	AB	41	U
2	AB	42	C
2	AB	45	A
2	AB	52	A

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Mol	Chain	Res	Type
2	AB	53	A
2	AB	57	A
2	AB	67	G
2	AB	73	A
2	AB	75	G
2	AB	88	C
2	AB	89	G
2	AB	89(A)	A
2	AB	96	G
2	AB	109	G
2	AB	112	G
31	BA	5	U
31	BA	6	G
31	BA	31	G
31	BA	32	A
31	BA	39	G
31	BA	48	C
31	BA	49	U
31	BA	50	A
31	BA	51	A
31	BA	61	G
31	BA	65	U
31	BA	66	G
31	BA	76	G
31	BA	78	G
31	BA	81	G
31	BA	84	U
31	BA	85	U
31	BA	86	U
31	BA	87	A
31	BA	90	C
31	BA	91	C
31	BA	92	G
31	BA	96	G
31	BA	101	A
31	BA	109	A
31	BA	110	C
31	BA	116	A
31	BA	120	A
31	BA	121	C
31	BA	122	G
31	BA	129(A)	G

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Mol	Chain	Res	Type
31	BA	130	A
31	BA	131	C
31	BA	163	C
31	BA	169	C
31	BA	173	U
31	BA	174	C
31	BA	182	U
31	BA	189	U
31	BA	190	G
31	BA	191(D)	U
31	BA	191(E)	G
31	BA	191(F)	U
31	BA	195	A
31	BA	197	A
31	BA	198	G
31	BA	201	C
31	BA	208	U
31	BA	209	U
31	BA	210	U
31	BA	216	G
31	BA	244	U
31	BA	245	C
31	BA	247	G
31	BA	251	G
31	BA	252	U
31	BA	265	G
31	BA	266	G
31	BA	267	C
31	BA	275	G
31	BA	279	A
31	BA	280	C
31	BA	281	G
31	BA	282	A
31	BA	289	G
31	BA	306	G
31	BA	316	G
31	BA	328	C
31	BA	329	A
31	BA	330	C
31	BA	332	G
31	BA	344	A
31	BA	345	C

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Mol	Chain	Res	Type
31	BA	346	G
31	BA	350	G
31	BA	352	C
31	BA	353	A
31	BA	354	G
31	BA	367	U
31	BA	368	U
31	BA	373	A
31	BA	389	A
31	BA	397	A
31	BA	398	C
31	BA	411	A
31	BA	412	A
31	BA	413	G
31	BA	414	A
31	BA	421	U
31	BA	422	C
31	BA	423	G
31	BA	428	G
31	BA	429	U
31	BA	430	A
31	BA	438	G
31	BA	439	A
31	BA	442	C
31	BA	451	A
31	BA	465	A
31	BA	467	G
31	BA	478	A
31	BA	482	A
31	BA	484	G
31	BA	485	G
31	BA	494	U
31	BA	495	A
31	BA	496	A
31	BA	497	U
31	BA	500	G
31	BA	505	G
31	BA	508	C
31	BA	509	A
31	BA	510	A
31	BA	511	C
31	BA	517	G

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Mol	Chain	Res	Type
31	BA	518	C
31	BA	519	C
31	BA	527	G
31	BA	530	G
31	BA	531	U
31	BA	532	A
31	BA	533	A
31	BA	534	U
31	BA	536	C
31	BA	548	G
31	BA	559	A
31	BA	560	U
31	BA	561	U
31	BA	562	C
31	BA	563	A
31	BA	566	G
31	BA	567	G
31	BA	572	A
31	BA	573	A
31	BA	575	G
31	BA	576	G
31	BA	577	G
31	BA	596	C
31	BA	598	U
31	BA	630	G
31	BA	632	A
31	BA	633	G
31	BA	642	A
31	BA	653	A
31	BA	665	A
31	BA	687	A
31	BA	688	G
31	BA	697	U
31	BA	701	C
31	BA	702	A
31	BA	703	G
31	BA	704	A
31	BA	721	G
31	BA	724	G
31	BA	731	G
31	BA	749	C
31	BA	753	A

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Mol	Chain	Res	Type
31	BA	754	C
31	BA	755	G
31	BA	793	U
31	BA	801	U
31	BA	802	A
31	BA	813	U
31	BA	815	A
31	BA	816	A
31	BA	817	C
31	BA	818	G
31	BA	819	A
31	BA	820	U
31	BA	821	G
31	BA	828	A
31	BA	841	U
31	BA	842	C
31	BA	843	U
31	BA	848	C
31	BA	855	G
31	BA	859	A
31	BA	871	U
31	BA	872	A
31	BA	873	A
31	BA	874	G
31	BA	885	G
31	BA	888	G
31	BA	889	A
31	BA	890	G
31	BA	914	A
31	BA	926	G
31	BA	927	G
31	BA	934	C
31	BA	935	A
31	BA	960	U
31	BA	961	U
31	BA	966	G
31	BA	968	A
31	BA	969	A
31	BA	974	A
31	BA	976	G
31	BA	977	A
31	BA	978	A

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Mol	Chain	Res	Type
31	BA	982	U
31	BA	983	A
31	BA	991	U
31	BA	992	U
31	BA	993	G
31	BA	994	A
31	BA	1001	G
31	BA	1004	A
31	BA	1005	A
31	BA	1008	C
31	BA	1021	G
31	BA	1024	G
31	BA	1025	U
31	BA	1028	C
31	BA	1028(A)	C
31	BA	1029	G
31	BA	1032(A)	G
31	BA	1036	G
31	BA	1042	G
31	BA	1050	G
31	BA	1053	G
31	BA	1054	C
31	BA	1055	A
31	BA	1064	G
31	BA	1065	U
31	BA	1066	C
31	BA	1068	G
31	BA	1085	U
31	BA	1086	U
31	BA	1094	G
31	BA	1095	U
31	BA	1101	A
31	BA	1102	A
31	BA	1117	G
31	BA	1118	C
31	BA	1124	G
31	BA	1125	U
31	BA	1127	G
31	BA	1128	C
31	BA	1130	A
31	BA	1131	G
31	BA	1137	C

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Mol	Chain	Res	Type
31	BA	1138	G
31	BA	1139	G
31	BA	1140	C
31	BA	1146	A
31	BA	1157	A
31	BA	1158	C
31	BA	1159	U
31	BA	1160	G
31	BA	1161	C
31	BA	1178	G
31	BA	1181	G
31	BA	1183	A
31	BA	1184	G
31	BA	1187	G
31	BA	1192	C
31	BA	1196	U
31	BA	1197	G
31	BA	1200	C
31	BA	1201	A
31	BA	1202	G
31	BA	1212	U
31	BA	1215	G
31	BA	1224	G
31	BA	1225	A
31	BA	1226	C
31	BA	1227	A
31	BA	1238	A
31	BA	1240	U
31	BA	1256	A
31	BA	1257	U
31	BA	1273	G
31	BA	1278	U
31	BA	1280	A
31	BA	1282	C
31	BA	1285	A
31	BA	1286	A
31	BA	1287	A
31	BA	1297	C
31	BA	1298	C
31	BA	1299	A
31	BA	1303	C
31	BA	1305	G

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Mol	Chain	Res	Type
31	BA	1317	C
31	BA	1320	C
31	BA	1321	C
31	BA	1322	C
31	BA	1323	G
31	BA	1331	G
31	BA	1335	C
31	BA	1337	G
31	BA	1338	G
31	BA	1345	U
31	BA	1347	G
31	BA	1348	U
31	BA	1362(A)	C
31	BA	1363	A
31	BA	1364	U
31	BA	1365	G
31	BA	1381	U
31	BA	1394	A
31	BA	1395	C
31	BA	1396	A
31	BA	1397	C
31	BA	1398	A
31	BA	1400	C
31	BA	1401	G
31	BA	1419	G
31	BA	1442	G
31	BA	1443	G
31	BA	1446	A
31	BA	1450	U
31	BA	1451	A
31	BA	1452	C
31	BA	1453	G
31	BA	1454	G
31	BA	1492	A
31	BA	1499	A
31	BA	1502	A
31	BA	1503	A
31	BA	1504	G
31	BA	1506	U
31	BA	1507	A
31	BA	1517	G
31	BA	1520	G

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Mol	Chain	Res	Type
31	BA	1528	U
31	BA	1529	G
31	BA	1530	G
31	BA	1533	C
31	BA	1538	C
31	BA	1541	U
52	BD	8	U
52	BD	9	A
52	BD	10	G
52	BD	16	U
52	BD	17	C
52	BD	20	U
52	BD	22	G
52	BD	36	A
52	BD	42	C
52	BD	46	G
52	BD	49	C
52	BD	58	A
52	BD	59	U
52	BD	61	C
52	BD	73	A
52	BB	8	U
52	BB	9	A
52	BB	10	G
52	BB	11	C
52	BB	16	U
52	BB	17	C
52	BB	18	G
52	BB	19	G
52	BB	20	U
52	BB	21	A
52	BB	22	G
52	BB	24	G
52	BB	25	C
52	BB	26	A
52	BB	27	G
52	BB	41	C
52	BB	44	G
52	BB	46	G
52	BB	47	U
52	BB	49	C
52	BB	58	A

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Mol	Chain	Res	Type
52	BB	59	U
52	BB	61	C
52	BB	70	G
52	BB	73	A
52	BB	74	C
52	BB	75	C
52	BB	76	A
52	BC	8	U
52	BC	9	A
52	BC	10	G
52	BC	17	C
52	BC	18	G
52	BC	19	G
52	BC	21	A
52	BC	22	G
52	BC	29	G
52	BC	43	C
52	BC	44	G
52	BC	46	G
52	BC	47	U
52	BC	48	C
52	BC	49	C
52	BC	58	A
52	BC	59	U
52	BC	61	C
52	BC	76	A
53	B1	32	A
53	B1	37	G
53	B1	42	U
53	B1	43	U
53	B1	49	U
53	B1	51	U
53	B1	52	U
53	B1	53	U
53	B1	54	U
53	B1	55	U
53	B1	56	U
53	B1	57	U
54	CA	9	G
54	CA	13	U
54	CA	14	U
54	CA	31	G

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Mol	Chain	Res	Type
54	CA	32	A
54	CA	39	G
54	CA	48	C
54	CA	49	U
54	CA	50	A
54	CA	51	A
54	CA	61	G
54	CA	64	G
54	CA	65	U
54	CA	66	G
54	CA	76	G
54	CA	78	G
54	CA	84	U
54	CA	85	U
54	CA	86	U
54	CA	87	A
54	CA	88	C
54	CA	89	U
54	CA	90	C
54	CA	91	C
54	CA	92	G
54	CA	95	G
54	CA	96	G
54	CA	97	U
54	CA	101	A
54	CA	109	A
54	CA	110	C
54	CA	116	A
54	CA	120	A
54	CA	121	C
54	CA	122	G
54	CA	130	A
54	CA	131	C
54	CA	144	G
54	CA	147	G
54	CA	163	C
54	CA	172	A
54	CA	173	U
54	CA	174	C
54	CA	183	G
54	CA	190	G
54	CA	191(A)	G

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Mol	Chain	Res	Type
54	CA	195	A
54	CA	197	A
54	CA	198	G
54	CA	209	U
54	CA	231	G
54	CA	244	U
54	CA	245	C
54	CA	247	G
54	CA	251	G
54	CA	252	U
54	CA	266	G
54	CA	267	C
54	CA	275	G
54	CA	281	G
54	CA	289	G
54	CA	305	G
54	CA	306	G
54	CA	315	A
54	CA	316	G
54	CA	321	A
54	CA	328	C
54	CA	329	A
54	CA	330	C
54	CA	332	G
54	CA	344	A
54	CA	345	C
54	CA	346	G
54	CA	352	C
54	CA	353	A
54	CA	354	G
54	CA	367	U
54	CA	368	U
54	CA	373	A
54	CA	388	G
54	CA	389	A
54	CA	397	A
54	CA	411	A
54	CA	412	A
54	CA	413	G
54	CA	414	A
54	CA	422	C
54	CA	423	G

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Mol	Chain	Res	Type
54	CA	428	G
54	CA	429	U
54	CA	430	A
54	CA	451	A
54	CA	482	A
54	CA	484	G
54	CA	485	G
54	CA	486	U
54	CA	495	A
54	CA	496	A
54	CA	497	U
54	CA	500	G
54	CA	508	C
54	CA	509	A
54	CA	510	A
54	CA	511	C
54	CA	517	G
54	CA	518	C
54	CA	519	C
54	CA	527	G
54	CA	531	U
54	CA	532	A
54	CA	533	A
54	CA	534	U
54	CA	536	C
54	CA	548	G
54	CA	559	A
54	CA	560	U
54	CA	561	U
54	CA	563	A
54	CA	566	G
54	CA	567	G
54	CA	572	A
54	CA	573	A
54	CA	575	G
54	CA	576	G
54	CA	577	G
54	CA	596	C
54	CA	616	G
54	CA	629	G
54	CA	630	G
54	CA	631	G

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Mol	Chain	Res	Type
54	CA	632	A
54	CA	642	A
54	CA	665	A
54	CA	666	G
54	CA	688	G
54	CA	701	C
54	CA	702	A
54	CA	703	G
54	CA	704	A
54	CA	721	G
54	CA	722	A
54	CA	731	G
54	CA	749	C
54	CA	754	C
54	CA	755	G
54	CA	791	G
54	CA	793	U
54	CA	794	A
54	CA	801	U
54	CA	802	A
54	CA	813	U
54	CA	815	A
54	CA	816	A
54	CA	817	C
54	CA	818	G
54	CA	819	A
54	CA	820	U
54	CA	821	G
54	CA	828	A
54	CA	841	U
54	CA	843	U
54	CA	848	C
54	CA	859	A
54	CA	871	U
54	CA	872	A
54	CA	873	A
54	CA	874	G
54	CA	885	G
54	CA	889	A
54	CA	890	G
54	CA	891	U
54	CA	902	G

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Mol	Chain	Res	Type
54	CA	914	A
54	CA	920	U
54	CA	926	G
54	CA	927	G
54	CA	934	C
54	CA	935	A
54	CA	960	U
54	CA	961	U
54	CA	966	G
54	CA	968	A
54	CA	969	A
54	CA	971	G
54	CA	972	C
54	CA	974	A
54	CA	975	A
54	CA	976	G
54	CA	977	A
54	CA	978	A
54	CA	982	U
54	CA	983	A
54	CA	991	U
54	CA	992	U
54	CA	993	G
54	CA	994	A
54	CA	1001	G
54	CA	1002	G
54	CA	1003	G
54	CA	1004	A
54	CA	1005	A
54	CA	1008	C
54	CA	1024	G
54	CA	1025	U
54	CA	1028	C
54	CA	1028(A)	C
54	CA	1029	G
54	CA	1032(A)	G
54	CA	1036	G
54	CA	1040	U
54	CA	1050	G
54	CA	1054	C
54	CA	1055	A
54	CA	1064	G

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Mol	Chain	Res	Type
54	CA	1065	U
54	CA	1066	C
54	CA	1068	G
54	CA	1085	U
54	CA	1086	U
54	CA	1094	G
54	CA	1095	U
54	CA	1101	A
54	CA	1102	A
54	CA	1117	G
54	CA	1124	G
54	CA	1125	U
54	CA	1126	U
54	CA	1127	G
54	CA	1130	A
54	CA	1131	G
54	CA	1136	U
54	CA	1137	C
54	CA	1138	G
54	CA	1139	G
54	CA	1140	C
54	CA	1146	A
54	CA	1157	A
54	CA	1158	C
54	CA	1159	U
54	CA	1160	G
54	CA	1177	G
54	CA	1178	G
54	CA	1179	A
54	CA	1181	G
54	CA	1182	G
54	CA	1183	A
54	CA	1191	A
54	CA	1196	U
54	CA	1197	G
54	CA	1200	C
54	CA	1201	A
54	CA	1202	G
54	CA	1212	U
54	CA	1215	G
54	CA	1224	G
54	CA	1225	A

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Mol	Chain	Res	Type
54	CA	1227	A
54	CA	1238	A
54	CA	1240	U
54	CA	1241	G
54	CA	1256	A
54	CA	1257	U
54	CA	1273	G
54	CA	1280	A
54	CA	1281	U
54	CA	1282	C
54	CA	1285	A
54	CA	1286	A
54	CA	1287	A
54	CA	1298	C
54	CA	1299	A
54	CA	1301	U
54	CA	1303	C
54	CA	1305	G
54	CA	1317	C
54	CA	1320	C
54	CA	1322	C
54	CA	1331	G
54	CA	1335	C
54	CA	1336	C
54	CA	1338	G
54	CA	1346	A
54	CA	1347	G
54	CA	1348	U
54	CA	1362(A)	C
54	CA	1363	A
54	CA	1364	U
54	CA	1365	G
54	CA	1370	G
54	CA	1381	U
54	CA	1395	C
54	CA	1396	A
54	CA	1397	C
54	CA	1398	A
54	CA	1400	C
54	CA	1401	G
54	CA	1419	G
54	CA	1442	G

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Mol	Chain	Res	Type
54	CA	1443	G
54	CA	1446	A
54	CA	1447	G
54	CA	1450	U
54	CA	1452	C
54	CA	1453	G
54	CA	1454	G
54	CA	1487	G
54	CA	1492	A
54	CA	1499	A
54	CA	1502	A
54	CA	1503	A
54	CA	1504	G
54	CA	1505	G
54	CA	1506	U
54	CA	1507	A
54	CA	1517	G
54	CA	1520	G
54	CA	1529	G
54	CA	1530	G
54	CA	1531	A
54	CA	1535	C
54	CA	1542	U
52	CD	2	C
52	CD	3	C
52	CD	8	U
52	CD	9	A
52	CD	13	C
52	CD	14	A
52	CD	17	C
52	CD	19	G
52	CD	21	A
52	CD	22	G
52	CD	42	C
52	CD	44	G
52	CD	45	U
52	CD	46	G
52	CD	47	U
52	CD	48	C
52	CD	49	C
52	CD	55	U
52	CD	56	C

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Mol	Chain	Res	Type
52	CD	59	U
52	CD	61	C
52	CD	73	A
52	CD	76	A
52	CB	8	U
52	CB	9	A
52	CB	10	G
52	CB	11	C
52	CB	16	U
52	CB	17	C
52	CB	18	G
52	CB	19	G
52	CB	20	U
52	CB	21	A
52	CB	22	G
52	CB	24	G
52	CB	26	A
52	CB	27	G
52	CB	41	C
52	CB	44	G
52	CB	47	U
52	CB	49	C
52	CB	58	A
52	CB	59	U
52	CB	70	G
52	CB	71	G
52	CB	74	C
52	CB	75	C
52	CB	76	A
52	CC	7	A
52	CC	8	U
52	CC	9	A
52	CC	10	G
52	CC	17	C
52	CC	18	G
52	CC	19	G
52	CC	20	U
52	CC	21	A
52	CC	29	G
52	CC	36	A
52	CC	45	U
52	CC	47	U

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Mol	Chain	Res	Type
52	CC	48	C
52	CC	58	A
52	CC	59	U
52	CC	61	C
52	CC	76	A
53	C1	32	A
53	C1	37	G
53	C1	40	U
53	C1	43	U
53	C1	45	U
53	C1	46	U
53	C1	53	U
53	C1	54	U
53	C1	56	U
53	C1	57	U
55	DA	5	A
55	DA	13	A
55	DA	14	A
55	DA	28	A
55	DA	34	C
55	DA	35	G
55	DA	46	C
55	DA	50	U
55	DA	51	G
55	DA	52	A
55	DA	63	U
55	DA	64	A
55	DA	70	G
55	DA	71	A
55	DA	72	U
55	DA	73	A
55	DA	74	A
55	DA	75	G
55	DA	84	A
55	DA	85	G
55	DA	88	G
55	DA	90	U
55	DA	91	A
55	DA	99	U
55	DA	101	G
55	DA	102	G
55	DA	118	A

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Mol	Chain	Res	Type
55	DA	119	A
55	DA	120	U
55	DA	121	G
55	DA	126	A
55	DA	138	G
55	DA	140	A
55	DA	165	U
55	DA	196	A
55	DA	197	A
55	DA	204	A
55	DA	205	G
55	DA	206	U
55	DA	214	G
55	DA	215	G
55	DA	216	A
55	DA	221	A
55	DA	222	A
55	DA	223	A
55	DA	227	A
55	DA	229	A
55	DA	230	U
55	DA	232	G
55	DA	233	A
55	DA	241	A
55	DA	242	G
55	DA	243	U
55	DA	248	G
55	DA	249	C
55	DA	250	G
55	DA	265	A
55	DA	266	G
55	DA	269	U
55	DA	270(L)	U
55	DA	270(M)	U
55	DA	270(O)	U
55	DA	270(P)	C
55	DA	271(A)	C
55	DA	271(C)	U
55	DA	271	G
55	DA	274	G
55	DA	275	G
55	DA	277	C

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Mol	Chain	Res	Type
55	DA	279	C
55	DA	284	U
55	DA	299	A
55	DA	301	G
55	DA	305	U
55	DA	311	A
55	DA	321	G
55	DA	322	A
55	DA	323	G
55	DA	324	A
55	DA	329	G
55	DA	330	A
55	DA	332	A
55	DA	333	G
55	DA	345	A
55	DA	346	A
55	DA	352	G
55	DA	353	G
55	DA	363	G
55	DA	364	C
55	DA	371	A
55	DA	372	G
55	DA	373	U
55	DA	386	G
55	DA	387	U
55	DA	388	G
55	DA	390	A
55	DA	391	G
55	DA	396	G
55	DA	403	U
55	DA	404	C
55	DA	405	U
55	DA	406	G
55	DA	411	G
55	DA	412	A
55	DA	421	U
55	DA	428	A
55	DA	434	U
55	DA	435	C
55	DA	442	G
55	DA	443	A
55	DA	444	C

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Mol	Chain	Res	Type
55	DA	446	G
55	DA	447	A
55	DA	448	U
55	DA	449	A
55	DA	454	A
55	DA	455	C
55	DA	456	C
55	DA	457	A
55	DA	458	G
55	DA	470	A
55	DA	475	U
55	DA	479	A
55	DA	480	A
55	DA	481	G
55	DA	482	A
55	DA	504	U
55	DA	505	A
55	DA	506	G
55	DA	508	G
55	DA	509	C
55	DA	526	A
55	DA	527	C
55	DA	528	A
55	DA	529	A
55	DA	531	C
55	DA	532	A
55	DA	533	G
55	DA	537	C
55	DA	539	G
55	DA	546	C
55	DA	549	G
55	DA	563	G
55	DA	572	A
55	DA	573	G
55	DA	574	C
55	DA	575	A
55	DA	586	A
55	DA	588	U
55	DA	603	A
55	DA	604	G
55	DA	607	U
55	DA	614	U

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Mol	Chain	Res	Type
55	DA	615	G
55	DA	616	A
55	DA	617	G
55	DA	621	A
55	DA	627	A
55	DA	628	G
55	DA	638	G
55	DA	644	A
55	DA	645	C
55	DA	646	A
55	DA	650	C
55	DA	651	G
55	DA	654	A
55	DA	654(E)	C
55	DA	654(F)	C
55	DA	654(G)	C
55	DA	654(H)	G
55	DA	654(I)	C
55	DA	654(J)	A
55	DA	654(K)	C
55	DA	654(L)	G
55	DA	654(N)	G
55	DA	654(S)	G
55	DA	654(T)	A
55	DA	670	A
55	DA	671	C
55	DA	686	G
55	DA	687	C
55	DA	702	G
55	DA	705	A
55	DA	722	A
55	DA	726	G
55	DA	730	C
55	DA	739	G
55	DA	747	U
55	DA	753	C
55	DA	762	U
55	DA	763	G
55	DA	775	G
55	DA	776	G
55	DA	777	A
55	DA	782	A

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Mol	Chain	Res	Type
55	DA	784	A
55	DA	785	G
55	DA	789	A
55	DA	790	C
55	DA	791	C
55	DA	792	G
55	DA	793	A
55	DA	794	G
55	DA	800	A
55	DA	801	G
55	DA	803	U
55	DA	805	G
55	DA	806	C
55	DA	811	U
55	DA	812	C
55	DA	819	A
55	DA	827	U
55	DA	828	U
55	DA	830	G
55	DA	831	G
55	DA	845	G
55	DA	847	U
55	DA	857	C
55	DA	858	U
55	DA	859	G
55	DA	860	U
55	DA	865	C
55	DA	866	A
55	DA	871	U
55	DA	878	A
55	DA	880	G
55	DA	881	G
55	DA	882	G
55	DA	883	G
55	DA	884	C
55	DA	885	C
55	DA	886	C
55	DA	887	A
55	DA	888	C
55	DA	889	C
55	DA	892	G
55	DA	893	C

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Mol	Chain	Res	Type
55	DA	896	A
55	DA	897	C
55	DA	898	C
55	DA	899	A
55	DA	900	A
55	DA	901	A
55	DA	906	G
55	DA	910	A
55	DA	913	U
55	DA	914	C
55	DA	917	A
55	DA	919	G
55	DA	930	U
55	DA	931	G
55	DA	933	A
55	DA	941	A
55	DA	946	G
55	DA	957	A
55	DA	958	U
55	DA	959	A
55	DA	961	C
55	DA	962	G
55	DA	973	A
55	DA	974	G
55	DA	975	G
55	DA	983	A
55	DA	989	G
55	DA	990	A
55	DA	991	C
55	DA	996	A
55	DA	1005	C
55	DA	1008	C
55	DA	1009	A
55	DA	1011	G
55	DA	1012	U
55	DA	1013	C
55	DA	1020	A
55	DA	1021	A
55	DA	1022	G
55	DA	1023	U
55	DA	1025	G
55	DA	1026	U

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Mol	Chain	Res	Type
55	DA	1027	A
55	DA	1034	G
55	DA	1044	G
55	DA	1045	A
55	DA	1046	A
55	DA	1047	G
55	DA	1048	A
55	DA	1049	C
55	DA	1050	A
55	DA	1054	A
55	DA	1055	G
55	DA	1056	G
55	DA	1057	A
55	DA	1058	U
55	DA	1059	G
55	DA	1060	U
55	DA	1061	U
55	DA	1062	G
55	DA	1066	U
55	DA	1067	A
55	DA	1068	G
55	DA	1070	A
55	DA	1071	G
55	DA	1074	G
55	DA	1076	C
55	DA	1077	A
55	DA	1078	U
55	DA	1079	C
55	DA	1080	A
55	DA	1082	U
55	DA	1084	A
55	DA	1085	A
55	DA	1086	A
55	DA	1088	A
55	DA	1090	U
55	DA	1092	C
55	DA	1093	G
55	DA	1095	A
55	DA	1096	A
55	DA	1097	U
55	DA	1099	G
55	DA	1103	A

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Mol	Chain	Res	Type
55	DA	1104	C
55	DA	1110	G
55	DA	1126	A
55	DA	1127	A
55	DA	1128	A
55	DA	1131	G
55	DA	1135	C
55	DA	1136	G
55	DA	1142	U
55	DA	1142(A)	A
55	DA	1143	A
55	DA	1144	G
55	DA	1152	C
55	DA	1155	A
55	DA	1156	A
55	DA	1157	G
55	DA	1170	G
55	DA	1173	G
55	DA	1174	A
55	DA	1175	U
55	DA	1176	G
55	DA	1178	C
55	DA	1179	C
55	DA	1180	C
55	DA	1181	C
55	DA	1195	G
55	DA	1204	A
55	DA	1205	U
55	DA	1206	G
55	DA	1211	U
55	DA	1212	G
55	DA	1213	A
55	DA	1220	A
55	DA	1221	C
55	DA	1236	G
55	DA	1237	A
55	DA	1238	G
55	DA	1241	A
55	DA	1242	A
55	DA	1247	A
55	DA	1248	G
55	DA	1249	U

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Mol	Chain	Res	Type
55	DA	1251	C
55	DA	1252	G
55	DA	1253	A
55	DA	1254	A
55	DA	1256	G
55	DA	1265	A
55	DA	1266	G
55	DA	1269	A
55	DA	1271	G
55	DA	1272	A
55	DA	1273	U
55	DA	1275	A
55	DA	1276	A
55	DA	1281	G
55	DA	1286	A
55	DA	1288	U
55	DA	1289	C
55	DA	1300	U
55	DA	1301	A
55	DA	1302	A
55	DA	1303	G
55	DA	1311	G
55	DA	1313	U
55	DA	1314	C
55	DA	1319	G
55	DA	1320	C
55	DA	1321	A
55	DA	1325	G
55	DA	1326	U
55	DA	1327	C
55	DA	1329	U
55	DA	1330	C
55	DA	1333	C
55	DA	1334	G
55	DA	1340	U
55	DA	1341	U
55	DA	1342	A
55	DA	1343	G
55	DA	1344	G
55	DA	1345	C
55	DA	1349	A
55	DA	1359	A

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Mol	Chain	Res	Type
55	DA	1360	A
55	DA	1365	A
55	DA	1368	G
55	DA	1372	U
55	DA	1379	A
55	DA	1380	G
55	DA	1384	A
55	DA	1385	G
55	DA	1386	C
55	DA	1389	G
55	DA	1395	A
55	DA	1396	U
55	DA	1397	U
55	DA	1398	C
55	DA	1406	U
55	DA	1407	C
55	DA	1411	C
55	DA	1416	G
55	DA	1419	A
55	DA	1420	U
55	DA	1421	G
55	DA	1427	A
55	DA	1428	C
55	DA	1429	G
55	DA	1444(A)	A
55	DA	1449	A
55	DA	1449(A)	G
55	DA	1451	C
55	DA	1453	A
55	DA	1454	U
55	DA	1455	G
55	DA	1459	G
55	DA	1460	A
55	DA	1461	G
55	DA	1467	C
55	DA	1471	A
55	DA	1475	G
55	DA	1482	U
55	DA	1483	G
55	DA	1485	G
55	DA	1490	A
55	DA	1491	G

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Mol	Chain	Res	Type
55	DA	1493	C
55	DA	1494	A
55	DA	1497	U
55	DA	1505	C
55	DA	1506	C
55	DA	1508	A
55	DA	1510	A
55	DA	1511	A
55	DA	1514	U
55	DA	1527	G
55	DA	1533	C
55	DA	1534	G
55	DA	1535	U
55	DA	1536	A
55	DA	1537	C
55	DA	1543	A
55	DA	1544	C
55	DA	1545	A
55	DA	1555	G
55	DA	1558	A
55	DA	1559	G
55	DA	1560	G
55	DA	1565	C
55	DA	1566	A
55	DA	1567	A
55	DA	1568	G
55	DA	1569	A
55	DA	1578	U
55	DA	1579	A
55	DA	1585	C
55	DA	1586	A
55	DA	1598	C
55	DA	1603	A
55	DA	1607	C
55	DA	1608	A
55	DA	1609	A
55	DA	1611	C
55	DA	1615	C
55	DA	1616	A
55	DA	1617	C
55	DA	1618	A
55	DA	1619	G

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Mol	Chain	Res	Type
55	DA	1634	A
55	DA	1635	G
55	DA	1640	C
55	DA	1647	G
55	DA	1648	C
55	DA	1652	A
55	DA	1653	G
55	DA	1654	A
55	DA	1667	G
55	DA	1668	A
55	DA	1674	G
55	DA	1675	C
55	DA	1676	A
55	DA	1682	G
55	DA	1693	U
55	DA	1694	C
55	DA	1695	G
55	DA	1698	A
55	DA	1699	G
55	DA	1700	A
55	DA	1706	U
55	DA	1707	G
55	DA	1725	G
55	DA	1729	A
55	DA	1730	U
55	DA	1731	G
55	DA	1733	G
55	DA	1735	C
55	DA	1742	C
55	DA	1743	G
55	DA	1758	G
55	DA	1759	A
55	DA	1763	G
55	DA	1764	G
55	DA	1773	A
55	DA	1780	A
55	DA	1781	C
55	DA	1782	C
55	DA	1785	A
55	DA	1786	A
55	DA	1787	A
55	DA	1791	A

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Mol	Chain	Res	Type
55	DA	1799	G
55	DA	1800	C
55	DA	1801	G
55	DA	1802	A
55	DA	1815	A
55	DA	1816	G
55	DA	1819	A
55	DA	1820	U
55	DA	1821	A
55	DA	1828	G
55	DA	1829	A
55	DA	1833	U
55	DA	1835	G
55	DA	1839	G
55	DA	1847	A
55	DA	1848	A
55	DA	1858	G
55	DA	1869	G
55	DA	1870	C
55	DA	1878	G
55	DA	1880	C
55	DA	1882	C
55	DA	1885	A
55	DA	1888	G
55	DA	1900	A
55	DA	1906	G
55	DA	1912	A
55	DA	1913	A
55	DA	1914	C
55	DA	1918	A
55	DA	1919	A
55	DA	1930	G
55	DA	1931	U
55	DA	1934	C
55	DA	1935	G
55	DA	1937	A
55	DA	1938	A
55	DA	1939	U
55	DA	1940	U
55	DA	1941	C
55	DA	1943	U
55	DA	1944	U

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Mol	Chain	Res	Type
55	DA	1945	G
55	DA	1954	G
55	DA	1955	U
55	DA	1956	U
55	DA	1963	U
55	DA	1964	G
55	DA	1965	C
55	DA	1966	A
55	DA	1967	C
55	DA	1969	A
55	DA	1970	A
55	DA	1971	A
55	DA	1972	A
55	DA	1980	G
55	DA	1981	A
55	DA	1982	C
55	DA	1986	A
55	DA	1993	U
55	DA	1997	G
55	DA	2020	A
55	DA	2021	C
55	DA	2022	U
55	DA	2023	G
55	DA	2031	A
55	DA	2032	G
55	DA	2033	A
55	DA	2034	U
55	DA	2036	C
55	DA	2043	C
55	DA	2044	C
55	DA	2051	A
55	DA	2052	G
55	DA	2055	C
55	DA	2056	G
55	DA	2059	A
55	DA	2060	A
55	DA	2061	G
55	DA	2062	A
55	DA	2067	G
55	DA	2068	U
55	DA	2069	G
55	DA	2092	U

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Mol	Chain	Res	Type
55	DA	2093	G
55	DA	2099	U
55	DA	2111	C
55	DA	2113	U
55	DA	2114	A
55	DA	2115	G
55	DA	2116	G
55	DA	2118	U
55	DA	2120	G
55	DA	2126	A
55	DA	2127	G
55	DA	2128	C
55	DA	2132	U
55	DA	2133	G
55	DA	2135	A
55	DA	2136	C
55	DA	2146	C
55	DA	2147	G
55	DA	2159	G
55	DA	2166	G
55	DA	2167	U
55	DA	2168	G
55	DA	2171	A
55	DA	2173	A
55	DA	2176	A
55	DA	2190	G
55	DA	2198	A
55	DA	2199	A
55	DA	2210	G
55	DA	2211	G
55	DA	2212	A
55	DA	2213	U
55	DA	2215	G
55	DA	2225	A
55	DA	2226	C
55	DA	2239	G
55	DA	2245	U
55	DA	2251	G
55	DA	2259	G
55	DA	2275	C
55	DA	2283	C
55	DA	2286	A

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Mol	Chain	Res	Type
55	DA	2288	A
55	DA	2289	G
55	DA	2290	G
55	DA	2296	U
55	DA	2297	C
55	DA	2305	A
55	DA	2307	G
55	DA	2308	G
55	DA	2309	A
55	DA	2311	A
55	DA	2319	G
55	DA	2320	A
55	DA	2321	G
55	DA	2325	G
55	DA	2326	C
55	DA	2334	G
55	DA	2335	A
55	DA	2336	A
55	DA	2337	G
55	DA	2345	G
55	DA	2346	A
55	DA	2347	C
55	DA	2350	C
55	DA	2382	G
55	DA	2383	G
55	DA	2384	G
55	DA	2385	C
55	DA	2390	U
55	DA	2392	A
55	DA	2402	C
55	DA	2403	C
55	DA	2406	U
55	DA	2407	G
55	DA	2424	C
55	DA	2425	A
55	DA	2426	A
55	DA	2427	C
55	DA	2429	G
55	DA	2430	A
55	DA	2432	A
55	DA	2435	A
55	DA	2439	A

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Mol	Chain	Res	Type
55	DA	2440	C
55	DA	2441	C
55	DA	2447	G
55	DA	2448	A
55	DA	2449	U
55	DA	2450	A
55	DA	2458	G
55	DA	2459	A
55	DA	2469	A
55	DA	2470	G
55	DA	2474	C
55	DA	2475	C
55	DA	2476	A
55	DA	2478	A
55	DA	2482	G
55	DA	2483	C
55	DA	2484	G
55	DA	2491	U
55	DA	2497	A
55	DA	2502	G
55	DA	2503	A
55	DA	2504	U
55	DA	2505	G
55	DA	2506	U
55	DA	2507	C
55	DA	2508	G
55	DA	2517	C
55	DA	2518	A
55	DA	2519	U
55	DA	2520	C
55	DA	2529	G
55	DA	2531	A
55	DA	2534	A
55	DA	2542	A
55	DA	2543	G
55	DA	2554	U
55	DA	2567	G
55	DA	2573	C
55	DA	2581	G
55	DA	2582	G
55	DA	2585	U
55	DA	2586	C

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Mol	Chain	Res	Type
55	DA	2599	G
55	DA	2602	A
55	DA	2609	U
55	DA	2610	C
55	DA	2611	U
55	DA	2612	C
55	DA	2613	U
55	DA	2614	A
55	DA	2615	U
55	DA	2629	A
55	DA	2645	G
55	DA	2646	C
55	DA	2654	A
55	DA	2655	G
55	DA	2665	A
55	DA	2673	G
55	DA	2675	A
55	DA	2682	U
55	DA	2690	C
55	DA	2691	C
55	DA	2702	U
55	DA	2703	C
55	DA	2712	U
55	DA	2712(A)	A
55	DA	2713	A
55	DA	2714	G
55	DA	2726	U
55	DA	2730	C
55	DA	2733	A
55	DA	2750	A
55	DA	2751	G
55	DA	2752	C
55	DA	2756	U
55	DA	2757	A
55	DA	2758	A
55	DA	2761	G
55	DA	2765	A
55	DA	2766	G
55	DA	2776	A
55	DA	2777	G
55	DA	2778	A
55	DA	2779	U

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Mol	Chain	Res	Type
55	DA	2780	G
55	DA	2781	A
55	DA	2790	A
55	DA	2791	C
55	DA	2793	G
55	DA	2797	U
55	DA	2799	A
55	DA	2820	A
55	DA	2833	G
55	DA	2834	G
55	DA	2835	A
55	DA	2836	U
55	DA	2845	G
55	DA	2848	G
55	DA	2849	U
55	DA	2866	U
55	DA	2867	G
55	DA	2872	G
55	DA	2874	C
55	DA	2879	C
55	DA	2880	C
55	DA	2892	A
55	DA	2894	G
2	DB	10	C
2	DB	12	C
2	DB	13	A
2	DB	15	A
2	DB	21	G
2	DB	24	G
2	DB	25	A
2	DB	27	C
2	DB	34	U
2	DB	35	U
2	DB	40	U
2	DB	41	U
2	DB	42	C
2	DB	45	A
2	DB	52	A
2	DB	53	A
2	DB	56	G
2	DB	57	A
2	DB	66	A

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Mol	Chain	Res	Type
2	DB	67	G
2	DB	73	A
2	DB	75	G
2	DB	81	G
2	DB	88	C
2	DB	89	G
2	DB	109	G
2	DB	112	G

All (1016) RNA pucker outliers are listed below:

Mol	Chain	Res	Type
1	AA	13	A
1	AA	27	G
1	AA	33	U
1	AA	34	C
1	AA	49	A
1	AA	51	G
1	AA	60	G
1	AA	70	G
1	AA	71	A
1	AA	72	U
1	AA	74	A
1	AA	83	G
1	AA	84	A
1	AA	119	A
1	AA	120	U
1	AA	125	G
1	AA	128	C
1	AA	139	G
1	AA	177	G
1	AA	196	A
1	AA	199	A
1	AA	204	A
1	AA	205	G
1	AA	214	G
1	AA	215	G
1	AA	221	A
1	AA	222	A
1	AA	227	A
1	AA	241	A
1	AA	242	G

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Mol	Chain	Res	Type
1	AA	249	C
1	AA	265	A
1	AA	271(B)	G
1	AA	278	A
1	AA	283	A
1	AA	301	G
1	AA	321	G
1	AA	322	A
1	AA	323	G
1	AA	329	G
1	AA	331	A
1	AA	332	A
1	AA	345	A
1	AA	352	G
1	AA	363(F)	A
1	AA	370	G
1	AA	371	A
1	AA	387	U
1	AA	390	A
1	AA	403	U
1	AA	404	C
1	AA	411	G
1	AA	421	U
1	AA	434	U
1	AA	442	G
1	AA	446	G
1	AA	447	A
1	AA	448	U
1	AA	454	A
1	AA	455	C
1	AA	456	C
1	AA	458	G
1	AA	474	G
1	AA	479	A
1	AA	503	A
1	AA	506	G
1	AA	508	G
1	AA	527	C
1	AA	529	A
1	AA	531	C
1	AA	532	A
1	AA	571	A

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Mol	Chain	Res	Type
1	AA	573	G
1	AA	574	C
1	AA	587	C
1	AA	603	A
1	AA	616	A
1	AA	620	G
1	AA	627	A
1	AA	637	A
1	AA	654(F)	C
1	AA	654(J)	A
1	AA	654(M)	C
1	AA	654(S)	G
1	AA	669	G
1	AA	670	A
1	AA	685	A
1	AA	686	G
1	AA	704	G
1	AA	726	G
1	AA	728	G
1	AA	739	G
1	AA	746	A
1	AA	752	A
1	AA	762	U
1	AA	764	A
1	AA	775	G
1	AA	776	G
1	AA	788	A
1	AA	789	A
1	AA	790	C
1	AA	791	C
1	AA	792	G
1	AA	793	A
1	AA	800	A
1	AA	801	G
1	AA	805	G
1	AA	811	U
1	AA	829	A
1	AA	830	G
1	AA	846	C
1	AA	856	C
1	AA	858	U
1	AA	859	G

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Mol	Chain	Res	Type
1	AA	865	C
1	AA	877	U
1	AA	888	C
1	AA	893	C
1	AA	896	A
1	AA	913	U
1	AA	930	U
1	AA	932	G
1	AA	945	A
1	AA	957	A
1	AA	961	C
1	AA	973	A
1	AA	974(A)	C
1	AA	989	G
1	AA	1008	C
1	AA	1020	A
1	AA	1022	G
1	AA	1060	U
1	AA	1085	A
1	AA	1126	A
1	AA	1130	U
1	AA	1156	A
1	AA	1171	G
1	AA	1173	G
1	AA	1204	A
1	AA	1205	U
1	AA	1210	A
1	AA	1212	G
1	AA	1236	G
1	AA	1237	A
1	AA	1247	A
1	AA	1248	G
1	AA	1250	G
1	AA	1251	C
1	AA	1272	A
1	AA	1275	A
1	AA	1286	A
1	AA	1288	U
1	AA	1300	U
1	AA	1301	A
1	AA	1302	A
1	AA	1312	U

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Mol	Chain	Res	Type
1	AA	1320	C
1	AA	1325	G
1	AA	1329	U
1	AA	1332	G
1	AA	1340	U
1	AA	1344	G
1	AA	1372	U
1	AA	1378	A
1	AA	1379	A
1	AA	1385	G
1	AA	1396	U
1	AA	1397	U
1	AA	1427	A
1	AA	1428	C
1	AA	1451	C
1	AA	1453	A
1	AA	1454	U
1	AA	1490	A
1	AA	1493	C
1	AA	1497	U
1	AA	1554	A
1	AA	1558	A
1	AA	1565	C
1	AA	1566	A
1	AA	1567	A
1	AA	1602	U
1	AA	1607	C
1	AA	1608	A
1	AA	1609	A
1	AA	1610	A
1	AA	1615	C
1	AA	1617	C
1	AA	1618	A
1	AA	1634	A
1	AA	1647	G
1	AA	1652	A
1	AA	1653	G
1	AA	1668	A
1	AA	1674	G
1	AA	1681	G
1	AA	1693	U
1	AA	1694	C

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Mol	Chain	Res	Type
1	AA	1698	A
1	AA	1699	G
1	AA	1706	U
1	AA	1758	G
1	AA	1762	A
1	AA	1780	A
1	AA	1784	A
1	AA	1786	A
1	AA	1799	G
1	AA	1800	C
1	AA	1801	G
1	AA	1815	A
1	AA	1818	U
1	AA	1819	A
1	AA	1820	U
1	AA	1900	A
1	AA	1918	A
1	AA	1929	G
1	AA	1930	G
1	AA	1936	A
1	AA	1937	A
1	AA	1938	A
1	AA	1939	U
1	AA	1940	U
1	AA	1943	U
1	AA	1944	U
1	AA	1954	G
1	AA	1955	U
1	AA	1962	C
1	AA	1963	U
1	AA	1964	G
1	AA	1966	A
1	AA	1970	A
1	AA	1980	G
1	AA	1992	G
1	AA	1996	C
1	AA	2021	C
1	AA	2022	U
1	AA	2032	G
1	AA	2033	A
1	AA	2035	G
1	AA	2051	A

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Mol	Chain	Res	Type
1	AA	2060	A
1	AA	2061	G
1	AA	2062	A
1	AA	2067	G
1	AA	2092	U
1	AA	2110	G
1	AA	2126	A
1	AA	2158	A
1	AA	2191	G
1	AA	2210	G
1	AA	2211	G
1	AA	2225	A
1	AA	2238	G
1	AA	2249	U
1	AA	2250	G
1	AA	2258	C
1	AA	2266	A
1	AA	2275	C
1	AA	2282	G
1	AA	2286	A
1	AA	2288	A
1	AA	2296	U
1	AA	2311	A
1	AA	2319	G
1	AA	2334	G
1	AA	2336	A
1	AA	2344	U
1	AA	2345	G
1	AA	2346	A
1	AA	2384	G
1	AA	2391	G
1	AA	2405	G
1	AA	2406	U
1	AA	2422	A
1	AA	2423	U
1	AA	2425	A
1	AA	2426	A
1	AA	2439	A
1	AA	2447	G
1	AA	2448	A
1	AA	2449	U
1	AA	2458	G

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Mol	Chain	Res	Type
1	AA	2481	G
1	AA	2490	G
1	AA	2497	A
1	AA	2503	A
1	AA	2506	U
1	AA	2517	C
1	AA	2518	A
1	AA	2519	U
1	AA	2566	A
1	AA	2572	A
1	AA	2581	G
1	AA	2602	A
1	AA	2609	U
1	AA	2610	C
1	AA	2613	U
1	AA	2614	A
1	AA	2655	G
1	AA	2681	C
1	AA	2689	U
1	AA	2690	C
1	AA	2712	U
1	AA	2713	A
1	AA	2756	U
1	AA	2776	A
1	AA	2778	A
1	AA	2779	U
1	AA	2780	G
1	AA	2790	A
1	AA	2835	A
1	AA	2848	G
1	AA	2849	U
1	AA	2859	G
1	AA	2866	U
1	AA	2867	G
1	AA	2873	A
1	AA	2879	C
2	AB	11	C
2	AB	12	C
2	AB	34	U
2	AB	56	G
2	AB	66	A
2	AB	108	C

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Mol	Chain	Res	Type
31	BA	4	U
31	BA	5	U
31	BA	7	G
31	BA	13	U
31	BA	30	U
31	BA	31	G
31	BA	47	C
31	BA	48	C
31	BA	49	U
31	BA	50	A
31	BA	51	A
31	BA	60	A
31	BA	65	U
31	BA	86	U
31	BA	89	U
31	BA	109	A
31	BA	115	G
31	BA	119	A
31	BA	121	C
31	BA	129(A)	G
31	BA	173	U
31	BA	181	G
31	BA	197	A
31	BA	201	C
31	BA	208	U
31	BA	209	U
31	BA	210	U
31	BA	243	A
31	BA	244	U
31	BA	246	A
31	BA	250	A
31	BA	251	G
31	BA	274	A
31	BA	279	A
31	BA	280	C
31	BA	305	G
31	BA	327	A
31	BA	328	C
31	BA	329	A
31	BA	351	G
31	BA	366	C
31	BA	367	U

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Mol	Chain	Res	Type
31	BA	372	C
31	BA	388	G
31	BA	410	G
31	BA	412	A
31	BA	428	G
31	BA	429	U
31	BA	451	A
31	BA	481	G
31	BA	484	G
31	BA	495	A
31	BA	498	A
31	BA	508	C
31	BA	509	A
31	BA	518	C
31	BA	531	U
31	BA	532	A
31	BA	533	A
31	BA	535	A
31	BA	547	A
31	BA	559	A
31	BA	560	U
31	BA	562	C
31	BA	566	G
31	BA	575	G
31	BA	576	G
31	BA	631	G
31	BA	632	A
31	BA	641	U
31	BA	653	A
31	BA	687	A
31	BA	701	C
31	BA	748	C
31	BA	753	A
31	BA	792	A
31	BA	812	C
31	BA	815	A
31	BA	817	C
31	BA	818	G
31	BA	819	A
31	BA	820	U
31	BA	870	U
31	BA	871	U

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Mol	Chain	Res	Type
31	BA	872	A
31	BA	873	A
31	BA	913	A
31	BA	934	C
31	BA	965	A
31	BA	968	A
31	BA	975	A
31	BA	982	U
31	BA	992	U
31	BA	1023	G
31	BA	1027	C
31	BA	1049	U
31	BA	1065	U
31	BA	1067	A
31	BA	1085	U
31	BA	1101	A
31	BA	1124	G
31	BA	1126	U
31	BA	1139	G
31	BA	1182	G
31	BA	1201	A
31	BA	1214	C
31	BA	1224	G
31	BA	1226	C
31	BA	1239	A
31	BA	1256	A
31	BA	1281	U
31	BA	1285	A
31	BA	1297	C
31	BA	1302	U
31	BA	1346	A
31	BA	1347	G
31	BA	1363	A
31	BA	1380	U
31	BA	1394	A
31	BA	1396	A
31	BA	1399	C
31	BA	1400	C
31	BA	1452	C
31	BA	1498	U
31	BA	1502	A
31	BA	1503	A

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Mol	Chain	Res	Type
31	BA	1504	G
31	BA	1506	U
31	BA	1528	U
31	BA	1529	G
52	BD	8	U
52	BD	9	A
52	BD	58	A
52	BB	7	A
52	BB	8	U
52	BB	9	A
52	BB	10	G
52	BB	18	G
52	BB	19	G
52	BB	58	A
52	BB	74	C
52	BC	7	A
52	BC	9	A
52	BC	18	G
52	BC	43	C
52	BC	46	G
52	BC	58	A
53	B1	42	U
53	B1	51	U
53	B1	53	U
53	B1	56	U
54	CA	7	G
54	CA	30	U
54	CA	31	G
54	CA	47	C
54	CA	48	C
54	CA	49	U
54	CA	50	A
54	CA	51	A
54	CA	60	A
54	CA	64	G
54	CA	65	U
54	CA	85	U
54	CA	89	U
54	CA	109	A
54	CA	115	G
54	CA	119	A
54	CA	121	C

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Mol	Chain	Res	Type
54	CA	173	U
54	CA	197	A
54	CA	243	A
54	CA	244	U
54	CA	246	A
54	CA	250	A
54	CA	251	G
54	CA	266	G
54	CA	274	A
54	CA	279	A
54	CA	280	C
54	CA	281	G
54	CA	305	G
54	CA	315	A
54	CA	327	A
54	CA	328	C
54	CA	329	A
54	CA	351	G
54	CA	366	C
54	CA	367	U
54	CA	372	C
54	CA	388	G
54	CA	410	G
54	CA	412	A
54	CA	428	G
54	CA	429	U
54	CA	451	A
54	CA	481	G
54	CA	484	G
54	CA	485	G
54	CA	495	A
54	CA	496	A
54	CA	498	A
54	CA	508	C
54	CA	509	A
54	CA	511	C
54	CA	517	G
54	CA	530	G
54	CA	531	U
54	CA	533	A
54	CA	535	A
54	CA	547	A

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Mol	Chain	Res	Type
54	CA	559	A
54	CA	560	U
54	CA	562	C
54	CA	566	G
54	CA	575	G
54	CA	576	G
54	CA	595	G
54	CA	630	G
54	CA	631	G
54	CA	641	U
54	CA	653	A
54	CA	687	A
54	CA	701	C
54	CA	703	G
54	CA	717	C
54	CA	721	G
54	CA	748	C
54	CA	753	A
54	CA	792	A
54	CA	812	C
54	CA	815	A
54	CA	817	C
54	CA	819	A
54	CA	820	U
54	CA	871	U
54	CA	873	A
54	CA	884	U
54	CA	889	A
54	CA	890	G
54	CA	913	A
54	CA	934	C
54	CA	960	U
54	CA	965	A
54	CA	968	A
54	CA	971	G
54	CA	975	A
54	CA	976	G
54	CA	982	U
54	CA	992	U
54	CA	993	G
54	CA	1027	C
54	CA	1049	U

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Mol	Chain	Res	Type
54	CA	1054	C
54	CA	1064	G
54	CA	1065	U
54	CA	1067	A
54	CA	1085	U
54	CA	1094	G
54	CA	1101	A
54	CA	1139	G
54	CA	1159	U
54	CA	1182	G
54	CA	1200	C
54	CA	1201	A
54	CA	1214	C
54	CA	1224	G
54	CA	1226	C
54	CA	1239	A
54	CA	1280	A
54	CA	1285	A
54	CA	1297	C
54	CA	1300	G
54	CA	1345	U
54	CA	1347	G
54	CA	1363	A
54	CA	1380	U
54	CA	1394	A
54	CA	1396	A
54	CA	1399	C
54	CA	1400	C
54	CA	1443	G
54	CA	1452	C
54	CA	1498	U
54	CA	1502	A
54	CA	1503	A
54	CA	1504	G
54	CA	1506	U
54	CA	1528	U
54	CA	1529	G
52	CD	2	C
52	CD	8	U
52	CD	18	G
52	CD	45	U
52	CD	58	A

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Mol	Chain	Res	Type
52	CB	7	A
52	CB	8	U
52	CB	9	A
52	CB	10	G
52	CB	18	G
52	CB	19	G
52	CB	58	A
52	CB	74	C
52	CC	7	A
52	CC	9	A
52	CC	19	G
52	CC	20	U
52	CC	46	G
52	CC	48	C
52	CC	58	A
53	C1	31	A
53	C1	42	U
53	C1	44	U
53	C1	56	U
55	DA	13	A
55	DA	27	G
55	DA	33	U
55	DA	34	C
55	DA	49	A
55	DA	50	U
55	DA	51	G
55	DA	60	G
55	DA	63	U
55	DA	70	G
55	DA	71	A
55	DA	72	U
55	DA	74	A
55	DA	84	A
55	DA	90	U
55	DA	99	U
55	DA	101	G
55	DA	119	A
55	DA	120	U
55	DA	125	G
55	DA	139	G
55	DA	177	G
55	DA	196	A

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Mol	Chain	Res	Type
55	DA	199	A
55	DA	204	A
55	DA	205	G
55	DA	215	G
55	DA	221	A
55	DA	222	A
55	DA	226	G
55	DA	227	A
55	DA	228	A
55	DA	229	A
55	DA	232	G
55	DA	241	A
55	DA	242	G
55	DA	249	C
55	DA	265	A
55	DA	270(Z)	U
55	DA	271(B)	G
55	DA	278	A
55	DA	283	A
55	DA	301	G
55	DA	311	A
55	DA	321	G
55	DA	322	A
55	DA	323	G
55	DA	329	G
55	DA	331	A
55	DA	332	A
55	DA	345	A
55	DA	352	G
55	DA	363(F)	A
55	DA	370	G
55	DA	371	A
55	DA	372	G
55	DA	386	G
55	DA	387	U
55	DA	390	A
55	DA	403	U
55	DA	404	C
55	DA	405	U
55	DA	411	G
55	DA	421	U
55	DA	434	U

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Mol	Chain	Res	Type
55	DA	442	G
55	DA	446	G
55	DA	447	A
55	DA	448	U
55	DA	454	A
55	DA	455	C
55	DA	457	A
55	DA	458	G
55	DA	474	G
55	DA	479	A
55	DA	481	G
55	DA	503	A
55	DA	506	G
55	DA	508	G
55	DA	527	C
55	DA	528	A
55	DA	529	A
55	DA	530	G
55	DA	531	C
55	DA	532	A
55	DA	562	U
55	DA	571	A
55	DA	573	G
55	DA	574	C
55	DA	587	C
55	DA	603	A
55	DA	614	U
55	DA	616	A
55	DA	620	G
55	DA	627	A
55	DA	637	A
55	DA	654(F)	C
55	DA	654(I)	C
55	DA	654(J)	A
55	DA	654(M)	C
55	DA	654(S)	G
55	DA	669	G
55	DA	670	A
55	DA	685	A
55	DA	686	G
55	DA	704	G
55	DA	726	G

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Mol	Chain	Res	Type
55	DA	739	G
55	DA	746	A
55	DA	752	A
55	DA	762	U
55	DA	764	A
55	DA	775	G
55	DA	776	G
55	DA	788	A
55	DA	789	A
55	DA	790	C
55	DA	791	C
55	DA	792	G
55	DA	793	A
55	DA	800	A
55	DA	801	G
55	DA	805	G
55	DA	811	U
55	DA	829	A
55	DA	830	G
55	DA	846	C
55	DA	856	C
55	DA	858	U
55	DA	859	G
55	DA	865	C
55	DA	877	U
55	DA	884	C
55	DA	887	A
55	DA	888	C
55	DA	895	U
55	DA	896	A
55	DA	913	U
55	DA	930	U
55	DA	932	G
55	DA	945	A
55	DA	957	A
55	DA	961	C
55	DA	973	A
55	DA	974(A)	C
55	DA	989	G
55	DA	995	C
55	DA	1008	C
55	DA	1011	G

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Mol	Chain	Res	Type
55	DA	1012	U
55	DA	1020	A
55	DA	1022	G
55	DA	1025	G
55	DA	1026	U
55	DA	1033	U
55	DA	1056	G
55	DA	1057	A
55	DA	1060	U
55	DA	1085	A
55	DA	1126	A
55	DA	1128	A
55	DA	1130	U
55	DA	1131	G
55	DA	1142(A)	A
55	DA	1143	A
55	DA	1156	A
55	DA	1173	G
55	DA	1175	U
55	DA	1178	C
55	DA	1204	A
55	DA	1205	U
55	DA	1210	A
55	DA	1211	U
55	DA	1212	G
55	DA	1220	A
55	DA	1236	G
55	DA	1237	A
55	DA	1247	A
55	DA	1250	G
55	DA	1251	C
55	DA	1252	G
55	DA	1253	A
55	DA	1265	A
55	DA	1266	G
55	DA	1272	A
55	DA	1275	A
55	DA	1286	A
55	DA	1288	U
55	DA	1300	U
55	DA	1301	A
55	DA	1302	A

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Mol	Chain	Res	Type
55	DA	1312	U
55	DA	1320	C
55	DA	1324	G
55	DA	1325	G
55	DA	1329	U
55	DA	1332	G
55	DA	1340	U
55	DA	1341	U
55	DA	1342	A
55	DA	1344	G
55	DA	1378	A
55	DA	1379	A
55	DA	1385	G
55	DA	1396	U
55	DA	1397	U
55	DA	1427	A
55	DA	1428	C
55	DA	1451	C
55	DA	1453	A
55	DA	1454	U
55	DA	1458	C
55	DA	1490	A
55	DA	1493	C
55	DA	1497	U
55	DA	1543	A
55	DA	1544	C
55	DA	1554	A
55	DA	1558	A
55	DA	1559	G
55	DA	1565	C
55	DA	1566	A
55	DA	1567	A
55	DA	1602	U
55	DA	1607	C
55	DA	1608	A
55	DA	1609	A
55	DA	1610	A
55	DA	1615	C
55	DA	1616	A
55	DA	1617	C
55	DA	1618	A
55	DA	1634	A

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Mol	Chain	Res	Type
55	DA	1647	G
55	DA	1652	A
55	DA	1653	G
55	DA	1667	G
55	DA	1674	G
55	DA	1681	G
55	DA	1693	U
55	DA	1694	C
55	DA	1698	A
55	DA	1699	G
55	DA	1706	U
55	DA	1758	G
55	DA	1762	A
55	DA	1780	A
55	DA	1781	C
55	DA	1784	A
55	DA	1786	A
55	DA	1799	G
55	DA	1800	C
55	DA	1801	G
55	DA	1815	A
55	DA	1818	U
55	DA	1819	A
55	DA	1820	U
55	DA	1828	G
55	DA	1838	C
55	DA	1899	G
55	DA	1900	A
55	DA	1912	A
55	DA	1913	A
55	DA	1918	A
55	DA	1925	C
55	DA	1929	G
55	DA	1930	G
55	DA	1936	A
55	DA	1937	A
55	DA	1939	U
55	DA	1940	U
55	DA	1943	U
55	DA	1944	U
55	DA	1954	G
55	DA	1955	U

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Mol	Chain	Res	Type
55	DA	1962	C
55	DA	1964	G
55	DA	1966	A
55	DA	1970	A
55	DA	1980	G
55	DA	1992	G
55	DA	1996	C
55	DA	2021	C
55	DA	2022	U
55	DA	2032	G
55	DA	2033	A
55	DA	2035	G
55	DA	2051	A
55	DA	2060	A
55	DA	2061	G
55	DA	2067	G
55	DA	2092	U
55	DA	2110	G
55	DA	2126	A
55	DA	2158	A
55	DA	2197	U
55	DA	2198	A
55	DA	2210	G
55	DA	2211	G
55	DA	2225	A
55	DA	2238	G
55	DA	2249	U
55	DA	2250	G
55	DA	2258	C
55	DA	2266	A
55	DA	2275	C
55	DA	2282	G
55	DA	2286	A
55	DA	2288	A
55	DA	2296	U
55	DA	2307	G
55	DA	2308	G
55	DA	2311	A
55	DA	2318	G
55	DA	2319	G
55	DA	2320	A
55	DA	2334	G

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Mol	Chain	Res	Type
55	DA	2335	A
55	DA	2336	A
55	DA	2344	U
55	DA	2345	G
55	DA	2346	A
55	DA	2382	G
55	DA	2384	G
55	DA	2391	G
55	DA	2405	G
55	DA	2423	U
55	DA	2425	A
55	DA	2426	A
55	DA	2439	A
55	DA	2447	G
55	DA	2448	A
55	DA	2449	U
55	DA	2458	G
55	DA	2468	G
55	DA	2481	G
55	DA	2490	G
55	DA	2503	A
55	DA	2506	U
55	DA	2507	C
55	DA	2517	C
55	DA	2519	U
55	DA	2566	A
55	DA	2572	A
55	DA	2581	G
55	DA	2585	U
55	DA	2602	A
55	DA	2609	U
55	DA	2610	C
55	DA	2613	U
55	DA	2614	A
55	DA	2638	G
55	DA	2645	G
55	DA	2654	A
55	DA	2681	C
55	DA	2689	U
55	DA	2690	C
55	DA	2712	U
55	DA	2713	A

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Mol	Chain	Res	Type
55	DA	2725	A
55	DA	2726	U
55	DA	2732	G
55	DA	2750	A
55	DA	2756	U
55	DA	2776	A
55	DA	2778	A
55	DA	2790	A
55	DA	2820	A
55	DA	2832	U
55	DA	2835	A
55	DA	2848	G
55	DA	2866	U
55	DA	2873	A
55	DA	2879	C
2	DB	12	C
2	DB	24	G
2	DB	34	U
2	DB	56	G
2	DB	66	A

5.4 Non-standard residues in protein, DNA, RNA chains ⓘ

6 non-standard protein/DNA/RNA residues are modelled in this entry.

In the following table, the Counts columns list the number of bonds (or angles) for which Mogul statistics could be retrieved, the number of bonds (or angles) that are observed in the model and the number of bonds (or angles) that are defined in the Chemical Component Dictionary. The Link column lists molecule types, if any, to which the group is linked. The Z score for a bond length (or angle) is the number of standard deviations the observed value is removed from the expected value. A bond length (or angle) with $|Z| > 2$ is considered an outlier worth inspection. RMSZ is the root-mean-square of all Z scores of the bond lengths (or angles).

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
52	MIA	BB	37	52	24,31,32	1.86	4 (16%)	26,44,47	3.09	6 (23%)
52	MIA	CD	37	52	24,31,32	1.94	4 (16%)	26,44,47	2.84	6 (23%)
52	MIA	CB	37	52	24,31,32	1.78	3 (12%)	26,44,47	3.35	5 (19%)
52	MIA	CC	37	52	24,31,32	1.89	4 (16%)	26,44,47	2.81	6 (23%)
52	MIA	BC	37	52	24,31,32	1.63	2 (8%)	26,44,47	2.81	6 (23%)
52	MIA	BD	37	52	24,31,32	2.00	4 (16%)	26,44,47	3.09	6 (23%)

In the following table, the Chirals column lists the number of chiral outliers, the number of chiral centers analysed, the number of these observed in the model and the number defined in the Chemical Component Dictionary. Similar counts are reported in the Torsion and Rings columns. '-' means no outliers of that kind were identified.

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
52	MIA	BB	37	52	-	2/11/33/34	0/3/3/3
52	MIA	CD	37	52	-	4/11/33/34	0/3/3/3
52	MIA	CB	37	52	-	4/11/33/34	0/3/3/3
52	MIA	CC	37	52	-	0/11/33/34	0/3/3/3
52	MIA	BC	37	52	-	2/11/33/34	0/3/3/3
52	MIA	BD	37	52	-	2/11/33/34	0/3/3/3

All (21) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
52	CB	37	MIA	C13-C14	6.49	1.51	1.32
52	CC	37	MIA	C13-C14	6.48	1.51	1.32
52	BD	37	MIA	C13-C14	6.43	1.50	1.32
52	BB	37	MIA	C13-C14	6.41	1.50	1.32
52	CD	37	MIA	C13-C14	6.40	1.50	1.32
52	BC	37	MIA	C13-C14	6.33	1.50	1.32
52	BD	37	MIA	C2-S10	5.77	1.80	1.75
52	CD	37	MIA	C2-S10	5.34	1.80	1.75
52	CC	37	MIA	C2-S10	4.71	1.79	1.75
52	BB	37	MIA	C2-S10	4.31	1.79	1.75
52	CB	37	MIA	C2-S10	3.84	1.79	1.75
52	BC	37	MIA	C12-C13	-3.13	1.34	1.48
52	BD	37	MIA	C12-C13	-3.03	1.34	1.48
52	CC	37	MIA	C12-C13	-2.93	1.34	1.48
52	BB	37	MIA	C12-C13	-2.88	1.35	1.48
52	CD	37	MIA	C12-C13	-2.81	1.35	1.48
52	CB	37	MIA	C12-C13	-2.79	1.35	1.48
52	BD	37	MIA	C6-N1	2.43	1.36	1.32
52	CD	37	MIA	C6-N1	2.23	1.35	1.32
52	CC	37	MIA	C6-N1	2.20	1.35	1.32
52	BB	37	MIA	C8-N7	-2.04	1.31	1.34

All (35) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
52	CB	37	MIA	C12-N6-C6	-12.70	103.74	122.55
52	BB	37	MIA	C11-S10-C2	10.55	110.15	102.27

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
52	BD	37	MIA	C11-S10-C2	10.29	109.95	102.27
52	CD	37	MIA	C11-S10-C2	10.25	109.92	102.27
52	BC	37	MIA	C12-N6-C6	-9.94	107.82	122.55
52	BD	37	MIA	C12-N6-C6	-9.59	108.35	122.55
52	CC	37	MIA	C11-S10-C2	9.42	109.30	102.27
52	CB	37	MIA	C11-S10-C2	9.40	109.29	102.27
52	BB	37	MIA	C12-N6-C6	-8.78	109.55	122.55
52	CC	37	MIA	C12-N6-C6	-8.25	110.33	122.55
52	CD	37	MIA	C12-N6-C6	-7.71	111.12	122.55
52	BC	37	MIA	C11-S10-C2	7.48	107.86	102.27
52	CB	37	MIA	C5-C6-N1	-3.97	117.52	120.81
52	CD	37	MIA	C5-C6-N1	-3.94	117.54	120.81
52	BD	37	MIA	C5-C6-N1	-3.83	117.63	120.81
52	BD	37	MIA	C12-C13-C14	-3.81	119.73	127.14
52	BC	37	MIA	C5-C6-N1	-3.73	117.72	120.81
52	BB	37	MIA	C5-C6-N1	-3.70	117.74	120.81
52	CC	37	MIA	C5-C6-N1	-3.65	117.78	120.81
52	BC	37	MIA	C12-C13-C14	-3.46	120.41	127.14
52	BB	37	MIA	C2-N3-C4	-3.30	110.78	115.32
52	BC	37	MIA	C2-N3-C4	-3.30	110.78	115.32
52	CB	37	MIA	C2-N3-C4	-3.29	110.79	115.32
52	BB	37	MIA	C12-C13-C14	-3.27	120.79	127.14
52	CC	37	MIA	C2-N3-C4	-3.24	110.86	115.32
52	CD	37	MIA	C2-N3-C4	-3.18	110.94	115.32
52	BD	37	MIA	C2-N3-C4	-3.15	110.98	115.32
52	CC	37	MIA	C12-C13-C14	-3.05	121.21	127.14
52	CD	37	MIA	C12-C13-C14	-2.66	121.96	127.14
52	CC	37	MIA	C16-C14-C15	2.61	120.36	114.60
52	BB	37	MIA	C16-C14-C15	2.59	120.33	114.60
52	BC	37	MIA	C16-C14-C15	2.49	120.11	114.60
52	CD	37	MIA	C16-C14-C15	2.39	119.89	114.60
52	CB	37	MIA	C16-C14-C15	2.35	119.79	114.60
52	BD	37	MIA	C16-C14-C15	2.16	119.37	114.60

There are no chirality outliers.

All (14) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
52	BB	37	MIA	N6-C12-C13-C14
52	CD	37	MIA	N1-C2-S10-C11
52	CD	37	MIA	N3-C2-S10-C11
52	CB	37	MIA	N3-C2-S10-C11

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Mol	Chain	Res	Type	Atoms
52	CB	37	MIA	N6-C12-C13-C14
52	BC	37	MIA	N1-C2-S10-C11
52	BD	37	MIA	N1-C2-S10-C11
52	BD	37	MIA	N3-C2-S10-C11
52	BC	37	MIA	N3-C2-S10-C11
52	BB	37	MIA	C3'-C4'-C5'-O5'
52	CD	37	MIA	C3'-C4'-C5'-O5'
52	CB	37	MIA	N1-C6-N6-C12
52	CB	37	MIA	C5-C6-N6-C12
52	CD	37	MIA	N6-C12-C13-C14

There are no ring outliers.

6 monomers are involved in 25 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
52	BB	37	MIA	2	0
52	CD	37	MIA	7	0
52	CB	37	MIA	6	0
52	CC	37	MIA	1	0
52	BC	37	MIA	1	0
52	BD	37	MIA	8	0

5.5 Carbohydrates [i](#)

There are no carbohydrates in this entry.

5.6 Ligand geometry [i](#)

Of 5351 ligands modelled in this entry, 5351 are monoatomic - leaving 0 for Mogul analysis.

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

There are no torsion outliers.

There are no ring outliers.

No monomer is involved in short contacts.

5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

The following chains have linkage breaks:

Mol	Chain	Number of breaks
31	BA	1

All chain breaks are listed below:

Model	Chain	Residue-1	Atom-1	Residue-2	Atom-2	Distance (Å)
1	BA	889:A	O3'	890:G	P	1.30

6 Fit of model and data ⓘ

6.1 Protein, DNA and RNA chains ⓘ

In the following table, the column labelled ‘#RSRZ> 2’ contains the number (and percentage) of RSRZ outliers, followed by percent RSRZ outliers for the chain as percentile scores relative to all X-ray entries and entries of similar resolution. The OWAB column contains the minimum, median, 95th percentile and maximum values of the occupancy-weighted average B-factor per residue. The column labelled ‘Q< 0.9’ lists the number of (and percentage) of residues with an average occupancy less than 0.9.

Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
1	AA	2909/2909 (100%)	-0.16	74 (2%) 57 34	24, 64, 192, 200	0
2	AB	122/122 (100%)	-0.26	2 (1%) 72 51	71, 95, 122, 159	0
2	DB	122/122 (100%)	-0.36	1 (0%) 86 72	43, 62, 79, 132	0
3	AD	272/276 (98%)	0.24	4 (1%) 73 54	29, 49, 71, 94	0
3	DD	272/276 (98%)	0.00	2 (0%) 87 75	15, 35, 58, 78	0
4	AE	205/206 (99%)	1.02	40 (19%) 1 0	41, 77, 115, 121	0
4	DE	205/206 (99%)	0.37	9 (4%) 34 17	16, 49, 92, 103	0
5	AF	208/210 (99%)	0.40	16 (7%) 13 5	33, 67, 130, 143	0
5	DF	202/210 (96%)	-0.20	1 (0%) 91 81	10, 43, 77, 88	0
6	AG	181/182 (99%)	1.04	35 (19%) 1 0	86, 105, 124, 138	0
6	DG	181/182 (99%)	0.20	6 (3%) 46 24	55, 70, 98, 109	0
7	AH	170/180 (94%)	2.75	92 (54%) 0 0	124, 158, 180, 187	0
7	DH	170/180 (94%)	0.34	7 (4%) 37 18	48, 84, 96, 100	0
8	AK	146/148 (98%)	0.47	11 (7%) 14 5	58, 93, 117, 128	0
8	DK	146/148 (98%)	0.61	13 (8%) 9 3	46, 93, 108, 114	0
9	AM	138/140 (98%)	1.36	31 (22%) 0 0	59, 83, 108, 111	0
9	DM	138/140 (98%)	-0.03	2 (1%) 75 56	31, 48, 85, 96	0
10	AN	122/122 (100%)	1.07	19 (15%) 2 1	42, 68, 81, 87	0
10	DN	122/122 (100%)	0.19	1 (0%) 86 72	24, 44, 58, 66	0
11	AO	150/150 (100%)	0.32	7 (4%) 31 15	43, 83, 109, 137	0
11	DO	150/150 (100%)	0.04	4 (2%) 54 31	23, 54, 87, 117	0
12	AP	141/141 (100%)	1.40	35 (24%) 0 0	49, 84, 112, 122	0
12	DP	141/141 (100%)	-0.03	3 (2%) 63 43	28, 48, 77, 95	0
13	A0	117/118 (99%)	-0.06	0 100 100	41, 60, 80, 91	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
13	D0	118/118 (100%)	-0.41	0 100 100	21, 47, 68, 79	0
14	AQ	111/112 (99%)	0.10	4 (3%) 42 22	71, 97, 127, 137	0
14	DQ	111/112 (99%)	0.28	8 (7%) 15 6	44, 63, 84, 97	0
15	AR	137/146 (93%)	0.09	6 (4%) 34 17	57, 77, 123, 139	0
15	DR	137/146 (93%)	0.28	5 (3%) 42 22	39, 59, 103, 134	0
16	A1	117/118 (99%)	1.06	20 (17%) 1 0	42, 76, 109, 122	0
16	D1	117/118 (99%)	-0.34	2 (1%) 70 49	25, 39, 64, 101	0
17	A2	101/101 (100%)	1.46	34 (33%) 0 0	41, 94, 110, 113	0
17	D2	101/101 (100%)	-0.09	3 (2%) 50 27	18, 60, 77, 83	0
18	AS	113/113 (100%)	0.40	3 (2%) 54 31	39, 57, 84, 106	0
18	DS	113/113 (100%)	-0.35	1 (0%) 84 69	28, 40, 64, 109	0
19	AT	92/96 (95%)	0.19	3 (3%) 46 24	45, 62, 86, 90	0
19	DT	92/96 (95%)	-0.34	0 100 100	24, 41, 60, 65	0
20	AU	102/110 (92%)	1.09	17 (16%) 1 1	58, 83, 144, 149	0
20	DU	102/110 (92%)	-0.07	5 (4%) 29 14	44, 71, 114, 123	0
21	AV	187/206 (90%)	2.65	106 (56%) 0 0	96, 128, 171, 184	0
21	DV	200/206 (97%)	0.92	36 (18%) 1 0	48, 97, 155, 163	0
22	A3	84/85 (98%)	0.24	6 (7%) 16 6	52, 71, 99, 110	0
22	D3	84/85 (98%)	-0.11	3 (3%) 42 22	25, 44, 80, 96	0
23	AZ	97/98 (98%)	0.49	9 (9%) 8 3	39, 61, 102, 111	0
23	DZ	97/98 (98%)	0.33	4 (4%) 37 18	23, 50, 98, 120	0
24	AW	69/72 (95%)	0.09	3 (4%) 35 17	48, 71, 115, 141	0
24	DW	69/72 (95%)	-0.17	1 (1%) 75 56	29, 53, 84, 104	0
25	AX	59/60 (98%)	1.49	14 (23%) 0 0	60, 81, 101, 112	0
25	DX	59/60 (98%)	-0.14	1 (1%) 70 49	32, 50, 73, 89	0
26	A4	71/71 (100%)	3.13	49 (69%) 0 0	126, 151, 157, 158	0
26	D4	71/71 (100%)	1.10	13 (18%) 1 0	95, 125, 138, 143	0
27	A5	59/60 (98%)	0.78	8 (13%) 3 1	45, 65, 140, 145	0
27	D5	59/60 (98%)	0.53	10 (16%) 1 0	19, 57, 159, 165	0
28	A6	45/54 (83%)	1.86	18 (40%) 0 0	126, 140, 154, 156	0
28	D6	45/54 (83%)	3.82	33 (73%) 0 0	86, 120, 140, 143	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
29	A7	49/49 (100%)	0.11	2 (4%) 37 18	30, 40, 98, 125	0
29	D7	49/49 (100%)	-0.30	2 (4%) 37 18	16, 25, 82, 109	0
30	A8	64/65 (98%)	0.44	3 (4%) 31 15	45, 64, 95, 120	0
30	D8	64/65 (98%)	-0.11	0 100 100	25, 39, 55, 98	0
31	BA	1516/1516 (100%)	-0.51	8 (0%) 91 81	32, 82, 155, 200	0
32	BE	237/256 (92%)	0.44	21 (8%) 9 3	83, 120, 159, 171	0
32	CE	237/256 (92%)	0.31	15 (6%) 20 8	65, 104, 144, 155	0
33	BF	206/239 (86%)	0.52	18 (8%) 10 4	82, 108, 137, 147	0
33	CF	205/239 (85%)	0.85	29 (14%) 2 1	54, 85, 115, 122	0
34	BG	208/209 (99%)	0.02	4 (1%) 66 46	57, 76, 93, 103	0
34	CG	208/209 (99%)	0.20	5 (2%) 59 37	62, 80, 99, 104	0
35	BH	151/162 (93%)	-0.21	1 (0%) 87 75	68, 82, 104, 117	0
35	CH	151/162 (93%)	-0.02	1 (0%) 87 75	49, 69, 99, 114	0
36	BI	101/101 (100%)	0.68	11 (10%) 5 2	55, 73, 90, 109	0
36	CI	101/101 (100%)	0.25	1 (0%) 82 67	47, 73, 83, 109	0
37	BJ	155/156 (99%)	0.48	7 (4%) 33 16	69, 93, 115, 124	0
37	CJ	155/156 (99%)	0.19	9 (5%) 23 10	59, 85, 104, 117	0
38	BK	138/138 (100%)	-0.34	0 100 100	73, 85, 101, 109	0
38	CK	138/138 (100%)	-0.33	0 100 100	56, 74, 86, 89	0
39	BL	127/128 (99%)	-0.20	4 (3%) 49 26	81, 114, 130, 141	0
39	CL	127/128 (99%)	-0.29	1 (0%) 86 72	58, 105, 125, 134	0
40	BM	99/105 (94%)	-0.30	2 (2%) 65 44	87, 121, 137, 139	0
40	CM	99/105 (94%)	0.31	7 (7%) 16 6	57, 106, 131, 133	0
41	BN	119/129 (92%)	1.11	25 (21%) 1 0	54, 77, 104, 125	0
41	CN	119/129 (92%)	0.85	15 (12%) 3 1	50, 70, 102, 120	0
42	BO	125/132 (94%)	0.70	12 (9%) 8 2	54, 74, 91, 124	0
42	CO	125/132 (94%)	0.51	13 (10%) 6 2	38, 51, 79, 111	0
43	BP	121/126 (96%)	0.47	13 (10%) 6 2	82, 118, 131, 135	0
43	CP	125/126 (99%)	-0.02	5 (4%) 38 19	57, 92, 111, 123	0
44	BQ	60/61 (98%)	-0.21	0 100 100	78, 99, 117, 122	0
44	CQ	60/61 (98%)	-0.30	0 100 100	65, 76, 88, 99	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
45	BR	88/89 (98%)	-0.05	1 (1%) 80 64	52, 74, 102, 112	0
45	CR	88/89 (98%)	-0.14	1 (1%) 80 64	43, 65, 93, 95	0
46	BS	84/88 (95%)	-0.66	0 100 100	57, 71, 92, 113	0
46	CS	84/88 (95%)	-0.65	1 (1%) 79 61	53, 78, 103, 128	0
47	BT	100/105 (95%)	-0.06	3 (3%) 50 27	60, 80, 106, 128	0
47	CT	100/105 (95%)	-0.31	1 (1%) 82 67	56, 77, 94, 110	0
48	BU	72/88 (81%)	0.64	9 (12%) 3 1	59, 75, 115, 121	0
48	CU	72/88 (81%)	0.45	3 (4%) 36 18	56, 68, 105, 117	0
49	BV	83/93 (89%)	0.50	9 (10%) 5 2	112, 124, 144, 151	0
49	CV	88/93 (94%)	0.28	6 (6%) 17 7	80, 98, 123, 141	0
50	BW	99/106 (93%)	-0.44	1 (1%) 82 67	61, 88, 128, 132	0
50	CW	99/106 (93%)	-0.45	0 100 100	60, 85, 120, 124	0
51	BX	25/27 (92%)	-0.44	0 100 100	92, 114, 122, 125	0
51	CX	25/27 (92%)	-0.53	0 100 100	84, 93, 110, 118	0
52	BB	75/76 (98%)	1.15	19 (25%) 0 0	82, 170, 186, 188	0
52	BC	75/76 (98%)	-0.06	1 (1%) 77 59	52, 85, 121, 139	0
52	BD	75/76 (98%)	0.22	4 (5%) 26 12	60, 156, 184, 192	0
52	CB	75/76 (98%)	1.08	15 (20%) 1 0	49, 148, 176, 180	0
52	CC	75/76 (98%)	0.04	2 (2%) 54 31	33, 67, 101, 114	0
52	CD	75/76 (98%)	0.60	8 (10%) 6 2	45, 151, 177, 181	0
53	B1	30/30 (100%)	0.17	1 (3%) 46 24	67, 153, 197, 198	0
53	C1	30/30 (100%)	1.00	6 (20%) 1 0	45, 141, 192, 194	0
54	CA	1515/1515 (100%)	-0.42	7 (0%) 91 81	27, 72, 153, 200	0
55	DA	2912/2912 (100%)	-0.15	53 (1%) 68 47	11, 42, 174, 200	0
56	DI	30/125 (24%)	1.40	6 (20%) 1 0	121, 127, 132, 143	0
56	DJ	30/125 (24%)	1.41	7 (23%) 0 0	124, 129, 137, 138	0
57	DY	145/173 (83%)	0.83	27 (18%) 1 0	132, 157, 169, 177	0
58	DL	145/147 (98%)	2.11	66 (45%) 0 0	128, 199, 200, 200	0
All	All	21582/22236 (97%)	0.13	1331 (6%) 20 9	10, 73, 158, 200	0

All (1331) RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
55	DA	654(L)	G	17.7
55	DA	654(K)	C	16.8
1	AA	2901	C	15.9
21	AV	187	ALA	15.5
1	AA	654(I)	C	13.7
21	AV	186	GLU	13.7
1	AA	654(J)	A	13.1
1	AA	654(L)	G	13.0
55	DA	654(J)	A	12.5
27	D5	54	GLY	12.4
55	DA	654(I)	C	11.6
1	AA	2797	U	11.4
55	DA	163	U	11.1
7	AH	48	GLY	10.8
1	AA	2900	A	10.7
21	AV	184	ALA	10.6
7	AH	18	GLU	10.2
1	AA	2902	C	10.1
1	AA	654(M)	C	9.7
16	A1	91	ASP	9.6
1	AA	654(H)	G	9.6
7	AH	32	GLU	9.5
1	AA	654(K)	C	9.3
28	D6	22	ALA	9.3
7	AH	24	VAL	9.1
1	AA	2798	C	8.8
7	AH	31	GLY	8.8
7	AH	43	VAL	8.8
20	AU	46	LYS	8.7
1	AA	1176	G	8.7
1	AA	2899	G	8.7
9	AM	1	MET	8.6
7	AH	128	PRO	8.5
1	AA	4	C	8.5
21	AV	179	ASP	8.4
26	A4	64	GLY	8.4
55	DA	2901	C	8.4
54	CA	86	U	8.3
1	AA	3	U	8.2
55	DA	2798	C	8.2
20	AU	48	ALA	8.1
21	AV	185	GLU	8.0
1	AA	2799	A	8.0

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Mol	Chain	Res	Type	RSRZ
21	AV	121	HIS	7.9
28	D6	29	ASN	7.9
1	AA	1067	A	7.9
21	AV	180	VAL	7.8
1	AA	2	G	7.6
4	AE	69	LYS	7.6
7	AH	30	LYS	7.5
58	DL	147	ALA	7.5
12	AP	33	GLY	7.5
58	DL	96	VAL	7.4
12	AP	91	GLU	7.4
20	AU	47	LYS	7.4
28	D6	13	CYS	7.4
17	A2	45	THR	7.4
7	AH	99	VAL	7.3
28	D6	49	HIS	7.3
55	DA	654(O)	G	7.3
43	BP	7	VAL	7.3
1	AA	654(G)	C	7.2
26	A4	40	HIS	7.2
15	DR	2	ASN	7.2
4	DE	205	ALA	7.1
7	AH	29	PRO	7.1
27	A5	2	ALA	7.0
7	AH	105	LEU	7.0
8	AK	146	ALA	7.0
16	A1	118	GLY	6.9
28	D6	42	TRP	6.9
57	DY	28	ASN	6.9
58	DL	87	GLY	6.9
26	A4	45	GLY	6.9
24	AW	72	ALA	6.8
55	DA	654(H)	G	6.8
52	BB	71	G	6.8
55	DA	654(M)	C	6.7
7	AH	52	VAL	6.7
57	DY	135	ARG	6.7
58	DL	97	GLY	6.7
28	D6	23	THR	6.6
21	AV	181	GLU	6.6
7	AH	53	GLU	6.5
7	AH	35	VAL	6.5

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Mol	Chain	Res	Type	RSRZ
52	BB	76	A	6.5
52	CB	71	G	6.5
21	AV	176	PRO	6.5
55	DA	164	U	6.4
12	AP	141	GLN	6.4
16	D1	117	GLN	6.4
28	D6	20	ASN	6.4
5	AF	10	PRO	6.3
20	AU	49	VAL	6.3
55	DA	654(N)	G	6.3
20	AU	52	SER	6.2
7	AH	23	ARG	6.2
1	AA	1066	U	6.2
55	DA	654(P)	G	6.2
1	AA	654(F)	C	6.2
20	AU	102	CYS	6.1
58	DL	146	ASP	6.1
27	A5	59	GLU	6.1
7	AH	47	GLU	6.0
57	DY	12	THR	6.0
14	AQ	2	ALA	6.0
9	AM	134	ARG	5.9
28	D6	26	ASN	5.9
42	BO	129	ALA	5.9
4	AE	205	ALA	5.9
58	DL	51	ALA	5.9
55	DA	2902	C	5.9
26	A4	44	THR	5.9
54	CA	85	U	5.9
11	DO	149	GLU	5.9
58	DL	50	ASP	5.8
30	A8	65	GLU	5.8
21	AV	107	THR	5.8
52	BB	72	C	5.8
18	DS	113	LYS	5.8
58	DL	48	MET	5.8
43	BP	84	ILE	5.8
21	DV	148	ASP	5.8
42	BO	128	ALA	5.8
58	DL	88	ALA	5.8
1	AA	2795	G	5.8
28	A6	41	PRO	5.7

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Mol	Chain	Res	Type	RSRZ
7	AH	17	VAL	5.7
26	D4	30	GLU	5.7
7	AH	49	VAL	5.7
41	BN	12	ARG	5.6
7	AH	44	VAL	5.6
42	CO	129	ALA	5.6
7	AH	103	LEU	5.6
32	CE	231	GLU	5.6
5	AF	14	PRO	5.6
21	AV	155	LEU	5.6
7	AH	107	VAL	5.6
21	AV	168	GLU	5.6
21	AV	183	LEU	5.5
7	AH	39	PRO	5.5
4	AE	76	ARG	5.5
6	AG	182	LYS	5.5
25	AX	29	ARG	5.5
26	A4	13	ARG	5.5
1	AA	2801	A	5.5
17	D2	36	PRO	5.5
26	A4	29	PRO	5.5
58	DL	77	LEU	5.4
26	A4	39	CYS	5.4
7	AH	98	LEU	5.4
58	DL	94	GLU	5.4
28	A6	14	THR	5.4
48	BU	88	LYS	5.4
11	AO	150	ALA	5.4
26	A4	37	SER	5.4
7	AH	125	VAL	5.3
21	AV	79	ARG	5.3
21	DV	177	PRO	5.3
21	AV	113	ALA	5.3
7	AH	25	LYS	5.3
16	A1	89	GLU	5.3
55	DA	654(F)	C	5.3
28	D6	43	CYS	5.3
1	AA	654(B)	C	5.3
55	DA	4	C	5.3
17	A2	36	PRO	5.3
28	D6	53	LYS	5.3
21	AV	150	LEU	5.2

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Mol	Chain	Res	Type	RSRZ
32	CE	233	SER	5.2
7	AH	34	GLU	5.2
52	BB	17	C	5.2
55	DA	654(G)	C	5.2
28	D6	21	TYR	5.2
28	D6	32	ASN	5.2
6	AG	2	PRO	5.2
49	BV	86	GLU	5.2
55	DA	654(R)	C	5.2
28	D6	14	THR	5.2
28	A6	42	TRP	5.2
57	DY	83	TYR	5.1
28	D6	41	PRO	5.1
52	CD	12	U	5.1
1	AA	2790	A	5.1
21	AV	170	THR	5.1
26	A4	14	ILE	5.1
26	A4	49	PHE	5.1
20	AU	50	ARG	5.1
12	AP	37	LEU	5.1
21	AV	146	ILE	5.1
55	DA	654(E)	C	5.1
26	A4	46	GLN	5.1
21	AV	182	LYS	5.0
22	D3	85	ALA	5.0
14	DQ	109	GLY	5.0
7	AH	26	VAL	5.0
30	A8	64	TYR	5.0
26	A4	28	LYS	5.0
21	AV	149	SER	4.9
1	AA	1093	G	4.9
16	A1	117	GLN	4.9
7	DH	83	TYR	4.9
24	AW	71	ASN	4.9
21	AV	82	ARG	4.9
21	DV	195	GLU	4.9
21	AV	165	VAL	4.9
55	DA	277	C	4.9
5	AF	12	LEU	4.9
54	CA	84	U	4.9
55	DA	1	G	4.9
53	C1	56	U	4.9

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Mol	Chain	Res	Type	RSRZ
55	DA	654(Q)	C	4.9
6	AG	39	ILE	4.8
5	AF	1	MET	4.8
26	A4	30	GLU	4.8
28	D6	12	GLU	4.8
7	AH	11	VAL	4.8
3	AD	26	LYS	4.8
41	CN	11	LYS	4.8
49	CV	87	ALA	4.8
27	D5	58	LEU	4.8
1	AA	1098	A	4.8
12	AP	65	PHE	4.7
7	AH	33	LEU	4.7
16	A1	90	VAL	4.7
27	D5	59	GLU	4.7
58	DL	141	ALA	4.7
58	DL	95	LYS	4.7
21	AV	177	PRO	4.7
15	AR	2	ASN	4.7
7	AH	123	PHE	4.7
7	AH	101	ARG	4.7
32	BE	232	PRO	4.7
26	A4	27	THR	4.7
28	D6	34	LEU	4.7
55	DA	2797	U	4.7
26	A4	55	ARG	4.7
15	DR	106	SER	4.7
41	CN	12	ARG	4.7
21	AV	97	GLU	4.7
12	AP	32	TYR	4.7
27	D5	2	ALA	4.7
7	AH	126	PRO	4.6
41	BN	13	GLN	4.6
25	AX	60	GLU	4.6
21	AV	2	GLU	4.6
55	DA	2799	A	4.6
21	AV	173	ALA	4.6
7	AH	19	VAL	4.6
17	A2	101	GLY	4.6
21	AV	122	ARG	4.6
21	DV	181	GLU	4.6
21	AV	162	GLU	4.6

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Mol	Chain	Res	Type	RSRZ
55	DA	2899	G	4.6
7	AH	129	THR	4.6
7	DH	101	ARG	4.6
58	DL	64	SER	4.6
14	AQ	108	GLY	4.5
17	A2	40	LEU	4.5
12	AP	23	GLY	4.5
43	BP	85	GLY	4.5
6	AG	152	LEU	4.5
26	A4	18	CYS	4.5
4	AE	70	ALA	4.5
1	AA	1177	A	4.5
58	DL	45	THR	4.5
12	AP	36	ALA	4.5
26	A4	65	ASP	4.5
4	AE	59	VAL	4.5
7	AH	84	SER	4.5
27	D5	53	ALA	4.5
58	DL	90	LYS	4.5
1	AA	2896	C	4.4
23	DZ	96	LYS	4.4
1	AA	1095	A	4.4
55	DA	2801	A	4.4
43	CP	125	ARG	4.4
28	A6	13	CYS	4.4
16	D1	118	GLY	4.4
41	CN	129	SER	4.4
1	AA	1068	G	4.4
12	AP	1	MET	4.4
20	AU	86	ARG	4.4
12	AP	90	VAL	4.4
58	DL	85	GLU	4.4
21	AV	118	GLN	4.4
1	AA	654(D)	G	4.3
17	A2	57	VAL	4.3
24	AW	43	GLN	4.3
26	A4	63	TYR	4.3
12	AP	104	PHE	4.3
26	A4	42	PHE	4.3
8	DK	139	GLN	4.3
9	AM	133	GLN	4.3
28	D6	40	CYS	4.3

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Mol	Chain	Res	Type	RSRZ
48	CU	17	SER	4.3
7	AH	114	VAL	4.3
7	AH	96	ALA	4.3
28	D6	47	THR	4.2
7	AH	148	ILE	4.2
43	BP	8	GLU	4.2
33	CF	88	ARG	4.2
23	DZ	94	LEU	4.2
37	BJ	156	TRP	4.2
48	CU	88	LYS	4.2
58	DL	21	PRO	4.2
7	DH	34	GLU	4.2
21	DV	188	ALA	4.2
21	DV	194	PRO	4.2
33	CF	90	GLU	4.2
4	AE	54	GLN	4.2
26	A4	31	ILE	4.2
27	A5	53	ALA	4.2
52	CB	17	C	4.2
28	D6	19	ARG	4.2
7	AH	81	GLU	4.2
21	AV	172	ALA	4.2
5	AF	13	SER	4.2
32	BE	219	VAL	4.2
16	A1	85	LYS	4.2
17	A2	1	MET	4.2
52	CB	45	U	4.2
55	DA	2900	A	4.2
21	AV	175	VAL	4.2
58	DL	89	HIS	4.2
49	BV	43	GLU	4.2
3	DD	26	LYS	4.2
9	AM	9	VAL	4.2
9	AM	43	THR	4.2
4	AE	77	ILE	4.2
55	DA	2795	G	4.1
21	AV	114	GLY	4.1
26	A4	33	VAL	4.1
9	AM	8	GLN	4.1
7	AH	116	GLU	4.1
12	AP	103	MET	4.1
52	CD	6	G	4.1

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Mol	Chain	Res	Type	RSRZ
1	AA	2898	U	4.1
7	AH	46	GLU	4.1
56	DJ	4	ASP	4.1
49	BV	44	MET	4.1
4	AE	72	VAL	4.1
37	CJ	139	GLU	4.1
29	A7	49	ARG	4.1
52	CB	73	A	4.1
21	AV	161	VAL	4.1
15	DR	1	MET	4.1
21	AV	68	PRO	4.1
9	AM	10	GLU	4.1
20	AU	51	VAL	4.1
52	CB	70	G	4.1
55	DA	896	A	4.0
58	DL	138	VAL	4.0
1	AA	2793	G	4.0
58	DL	13	PRO	4.0
21	AV	178	GLU	4.0
28	A6	25	LYS	4.0
47	BT	101	ARG	4.0
2	AB	1(M)	A	4.0
11	AO	149	GLU	4.0
41	CN	128	ALA	4.0
57	DY	116	ILE	4.0
26	A4	7	PRO	4.0
52	BD	17	C	4.0
21	AV	50	GLN	3.9
21	AV	169	GLU	3.9
55	DA	888	C	3.9
21	AV	72	ARG	3.9
21	DV	147	GLY	3.9
28	D6	18	ARG	3.9
7	AH	140	LYS	3.9
7	AH	102	ALA	3.9
21	AV	156	LYS	3.9
7	AH	82	GLY	3.9
52	BB	3	C	3.9
21	DV	179	ASP	3.9
14	DQ	108	GLY	3.9
33	CF	66	VAL	3.9
8	DK	146	ALA	3.9

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Mol	Chain	Res	Type	RSRZ
17	A2	32	THR	3.9
55	DA	887	A	3.9
12	AP	105	GLU	3.9
55	DA	3	U	3.9
58	DL	44	ALA	3.9
7	AH	16	SER	3.9
17	A2	14	VAL	3.9
21	AV	142	SER	3.8
8	AK	76	THR	3.8
26	A4	12	ALA	3.8
58	DL	14	ALA	3.8
12	AP	31	ASP	3.8
1	AA	1509	C	3.8
7	AH	37	VAL	3.8
33	BF	46	GLU	3.8
41	BN	35	PRO	3.8
58	DL	27	LEU	3.8
27	D5	57	VAL	3.8
7	DH	23	ARG	3.8
21	AV	152	ALA	3.8
41	BN	129	SER	3.8
58	DL	47	ASN	3.8
26	A4	10	VAL	3.8
21	DV	184	ALA	3.8
17	A2	99	ILE	3.8
6	AG	178	PHE	3.8
7	AH	124	GLU	3.8
57	DY	129	PRO	3.8
2	AB	88	C	3.8
27	D5	60	VAL	3.7
52	CB	72	C	3.7
26	A4	9	LEU	3.7
26	A4	43	TYR	3.7
20	AU	60	PHE	3.7
28	D6	51	GLU	3.7
58	DL	49	GLY	3.7
15	DR	135	ALA	3.7
7	AH	95	ARG	3.7
36	CI	101	ALA	3.7
5	AF	11	VAL	3.7
9	AM	37	LYS	3.7
28	D6	45	LYS	3.7

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Mol	Chain	Res	Type	RSRZ
12	AP	106	VAL	3.7
9	AM	36	GLY	3.7
21	AV	164	ALA	3.7
58	DL	16	LYS	3.7
10	AN	42	SER	3.7
57	DY	108	LYS	3.7
26	A4	36	CYS	3.7
33	CF	166	GLU	3.7
52	CB	76	A	3.7
25	AX	53	LEU	3.6
4	DE	204	ALA	3.6
8	AK	145	VAL	3.6
58	DL	104	VAL	3.6
41	CN	16	SER	3.6
55	DA	165	U	3.6
1	AA	2794	C	3.6
25	AX	30	ARG	3.6
21	AV	171	ILE	3.6
52	CD	5	G	3.6
52	BD	12	U	3.6
36	BI	1	MET	3.6
26	A4	32	TYR	3.6
48	BU	17	SER	3.6
28	A6	49	HIS	3.6
27	A5	52	TYR	3.6
33	CF	101	LEU	3.6
10	AN	81	ASP	3.6
1	AA	1065	U	3.6
28	A6	50	ARG	3.6
1	AA	654(O)	G	3.6
12	AP	63	LYS	3.5
6	AG	118	ARG	3.5
28	A6	21	TYR	3.5
11	AO	92	GLU	3.5
55	DA	2794	C	3.5
58	DL	30	HIS	3.5
7	AH	28	GLY	3.5
43	BP	5	ALA	3.5
57	DY	100	ASN	3.5
28	D6	50	ARG	3.5
7	AH	40	GLU	3.5
6	AG	166	ASP	3.5

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Mol	Chain	Res	Type	RSRZ
58	DL	79	ARG	3.5
4	DE	187	ALA	3.5
7	AH	83	TYR	3.5
28	D6	39	TYR	3.5
21	AV	145	GLU	3.5
24	DW	43	GLN	3.5
21	AV	80	ARG	3.5
32	BE	4	GLU	3.5
9	AM	138	LEU	3.5
9	DM	130	HIS	3.5
8	AK	140	LEU	3.5
52	CD	13	C	3.5
32	BE	228	GLY	3.5
12	AP	99	PRO	3.4
6	DG	182	LYS	3.4
9	AM	60	ILE	3.4
16	A1	109	LEU	3.4
26	A4	52	THR	3.4
32	BE	221	LEU	3.4
57	DY	109	SER	3.4
9	AM	41	ASP	3.4
55	DA	270(L)	U	3.4
31	BA	1029	G	3.4
25	AX	28	LEU	3.4
42	CO	39	VAL	3.4
4	DE	68	ALA	3.4
16	A1	116	ALA	3.4
41	BN	109	VAL	3.4
4	AE	1	MET	3.4
1	AA	654(E)	C	3.4
21	AV	138	GLU	3.4
4	AE	85	ASN	3.4
21	AV	106	GLY	3.4
18	AS	113	LYS	3.4
5	AF	20	LEU	3.4
58	DL	43	ALA	3.4
4	AE	51	PHE	3.4
14	DQ	107	GLU	3.4
12	AP	7	MET	3.4
23	AZ	98	LEU	3.4
55	DA	276	A	3.4
9	AM	12	ARG	3.4

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Mol	Chain	Res	Type	RSRZ
21	DV	107	THR	3.4
12	AP	38	GLU	3.4
7	AH	55	PRO	3.4
56	DI	12	LEU	3.4
26	A4	16	CYS	3.4
58	DL	93	ARG	3.4
7	AH	72	ILE	3.4
7	AH	9	ILE	3.4
21	AV	115	GLY	3.4
6	AG	179	PRO	3.3
20	AU	59	GLY	3.3
53	C1	28	G	3.3
10	AN	12	ASP	3.3
32	CE	229	VAL	3.3
52	BB	75	C	3.3
14	AQ	60	GLY	3.3
55	DA	2141	G	3.3
4	AE	198	VAL	3.3
54	CA	87	A	3.3
21	DV	170	THR	3.3
29	D7	49	ARG	3.3
32	BE	36	ARG	3.3
21	AV	174	VAL	3.3
12	AP	60	ARG	3.3
6	AG	155	MET	3.3
7	AH	131	VAL	3.3
33	CF	84	ILE	3.3
21	DV	198	LYS	3.3
52	BB	73	A	3.3
4	AE	68	ALA	3.3
21	DV	197	ILE	3.3
5	AF	2	LYS	3.3
34	CG	169	LYS	3.3
33	BF	85	ARG	3.3
40	CM	91	PRO	3.3
43	BP	16	ASP	3.3
7	AH	36	PRO	3.3
17	D2	45	THR	3.3
7	AH	41	MET	3.3
26	D4	31	ILE	3.3
49	CV	86	GLU	3.3
41	CN	17	GLY	3.3

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Mol	Chain	Res	Type	RSRZ
43	BP	6	GLY	3.3
1	AA	5	A	3.3
56	DJ	22	GLN	3.3
12	AP	100	GLY	3.3
21	AV	124	ILE	3.3
33	BF	77	ILE	3.3
7	AH	45	VAL	3.3
33	CF	58	GLU	3.2
4	AE	204	ALA	3.2
26	D4	61	ARG	3.2
6	AG	82	LEU	3.2
27	A5	58	LEU	3.2
58	DL	26	ALA	3.2
1	AA	2802	G	3.2
8	DK	135	GLU	3.2
7	AH	80	SER	3.2
23	DZ	80	LEU	3.2
17	A2	46	VAL	3.2
28	D6	48	VAL	3.2
28	A6	51	GLU	3.2
53	C1	57	U	3.2
41	BN	17	GLY	3.2
58	DL	73	PRO	3.2
5	AF	133	ASN	3.2
43	CP	8	GLU	3.2
32	BE	6	THR	3.2
23	AZ	54	ALA	3.2
4	AE	29	GLY	3.2
21	DV	106	GLY	3.2
21	DV	180	VAL	3.2
42	BO	39	VAL	3.2
1	AA	1	G	3.2
1	AA	654(N)	G	3.2
7	AH	50	VAL	3.2
49	CV	89	ALA	3.2
8	DK	72	LEU	3.2
26	A4	19	GLY	3.2
55	DA	162	U	3.1
9	AM	51	PHE	3.1
7	AH	38	SER	3.1
49	BV	47	HIS	3.1
58	DL	145	LYS	3.1

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Mol	Chain	Res	Type	RSRZ
21	AV	112	ARG	3.1
21	DV	146	ILE	3.1
26	A4	71	ARG	3.1
39	BL	56	LEU	3.1
7	AH	89	ILE	3.1
9	AM	13	TRP	3.1
12	AP	20	ALA	3.1
17	A2	70	ILE	3.1
57	DY	82	PHE	3.1
21	AV	127	LYS	3.1
48	BU	23	LYS	3.1
42	CO	56	ALA	3.1
21	AV	144	LEU	3.1
23	AZ	2	SER	3.1
52	CB	2	C	3.1
4	AE	84	PHE	3.1
9	AM	40	PRO	3.1
4	AE	40	GLU	3.1
21	AV	56	VAL	3.1
28	D6	52	VAL	3.1
7	AH	170	ARG	3.1
17	A2	91	TYR	3.1
32	CE	61	LEU	3.1
58	DL	92	GLY	3.1
6	DG	2	PRO	3.1
26	A4	53	GLU	3.1
5	AF	207	GLY	3.1
10	AN	13	ASN	3.1
4	AE	74	PRO	3.1
7	AH	141	VAL	3.1
21	AV	140	ASP	3.1
58	DL	144	VAL	3.1
55	DA	5	A	3.1
23	AZ	95	LEU	3.1
58	DL	17	ALA	3.1
33	BF	88	ARG	3.1
39	CL	95	LYS	3.1
41	BN	16	SER	3.1
54	CA	345	C	3.1
17	A2	5	VAL	3.1
41	CN	36	ASP	3.1
8	AK	72	LEU	3.1

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Mol	Chain	Res	Type	RSRZ
21	AV	151	HIS	3.1
17	A2	44	LYS	3.1
49	CV	88	LYS	3.1
21	AV	71	VAL	3.0
58	DL	6	ALA	3.0
8	AK	71	ILE	3.0
28	A6	26	ASN	3.0
42	BO	65	GLU	3.0
33	BF	67	THR	3.0
41	BN	18	ARG	3.0
57	DY	90	ALA	3.0
21	AV	160	GLY	3.0
32	BE	222	ILE	3.0
26	A4	22	ILE	3.0
42	BO	64	TYR	3.0
28	A6	12	GLU	3.0
56	DI	8	ILE	3.0
6	AG	115	ARG	3.0
6	AG	116	ASP	3.0
21	AV	102	LEU	3.0
21	AV	148	ASP	3.0
11	DO	150	ALA	3.0
23	AZ	3	LYS	3.0
25	AX	8	LEU	3.0
33	CF	98	ASN	3.0
27	D5	51	TYR	3.0
41	BN	68	ALA	3.0
28	D6	24	GLU	3.0
58	DL	28	GLY	3.0
26	A4	11	PRO	3.0
12	AP	26	TYR	3.0
58	DL	20	ALA	3.0
1	AA	1064	C	3.0
52	BB	74	C	3.0
1	AA	945	A	3.0
4	AE	50	GLY	3.0
21	DV	149	SER	3.0
26	A4	70	GLY	3.0
10	AN	57	VAL	3.0
26	D4	40	HIS	3.0
1	AA	654(C)	G	3.0
55	DA	2	G	3.0

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Mol	Chain	Res	Type	RSRZ
33	CF	99	VAL	3.0
6	AG	34	LEU	3.0
8	AK	75	LEU	3.0
8	DK	140	LEU	3.0
9	AM	116	LEU	3.0
49	BV	29	ARG	3.0
34	BG	179	GLU	3.0
10	AN	51	ALA	3.0
7	AH	76	VAL	3.0
28	A6	34	LEU	3.0
10	AN	65	THR	3.0
36	BI	39	LYS	3.0
33	CF	19	GLU	3.0
33	BF	84	ILE	3.0
21	AV	86	VAL	3.0
33	CF	87	LEU	3.0
42	BO	55	VAL	3.0
55	DA	885	C	3.0
26	A4	20	ASN	3.0
33	CF	102	ASN	3.0
31	BA	1028(B)	C	2.9
6	AG	157	ILE	2.9
58	DL	25	PRO	2.9
20	AU	45	VAL	2.9
53	B1	55	U	2.9
5	AF	208	GLY	2.9
6	AG	145	THR	2.9
32	CE	133	LYS	2.9
21	AV	163	LEU	2.9
26	A4	21	VAL	2.9
3	DD	34	VAL	2.9
25	AX	55	ARG	2.9
32	BE	220	ASP	2.9
37	CJ	52	GLU	2.9
10	AN	52	VAL	2.9
21	DV	191	VAL	2.9
52	BB	70	G	2.9
28	D6	25	LYS	2.9
42	CO	28	LYS	2.9
43	BP	4	ILE	2.9
11	AO	13	ASN	2.9
43	CP	123	ALA	2.9

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Mol	Chain	Res	Type	RSRZ
21	AV	69	THR	2.9
6	AG	41	GLN	2.9
21	AV	120	ILE	2.9
21	AV	154	ASP	2.9
17	A2	56	SER	2.9
21	AV	99	TYR	2.9
58	DL	74	ALA	2.9
28	A6	20	ASN	2.9
6	AG	137	GLU	2.9
52	CD	11	C	2.9
17	A2	35	LEU	2.9
16	A1	71	GLN	2.9
26	A4	25	TYR	2.9
49	CV	61	TYR	2.9
52	BB	10	G	2.9
9	DM	134	ARG	2.9
21	AV	4	ARG	2.9
20	AU	88	LYS	2.9
4	AE	181	LEU	2.9
1	AA	654(P)	G	2.9
5	AF	7	TYR	2.9
49	BV	28	LYS	2.9
32	CE	232	PRO	2.9
7	AH	51	ARG	2.8
21	AV	95	PRO	2.8
1	AA	654(R)	C	2.8
7	AH	94	TYR	2.8
52	BD	11	C	2.8
54	CA	1030	C	2.8
55	DA	270(K)	C	2.8
25	AX	27	GLY	2.8
21	DV	105	VAL	2.8
48	BU	20	ALA	2.8
4	AE	88	GLY	2.8
4	AE	4	ILE	2.8
9	AM	14	VAL	2.8
26	A4	15	ILE	2.8
26	D4	69	LYS	2.8
32	BE	218	ALA	2.8
56	DJ	19	GLU	2.8
1	AA	1099	G	2.8
7	AH	100	GLY	2.8

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Mol	Chain	Res	Type	RSRZ
21	AV	76	LEU	2.8
21	AV	83	PRO	2.8
21	AV	123	ASP	2.8
58	DL	65	PHE	2.8
58	DL	142	PRO	2.8
42	BO	56	ALA	2.8
57	DY	99	SER	2.8
43	BP	121	LYS	2.8
4	DE	7	VAL	2.8
1	AA	654(Q)	C	2.8
7	AH	136	ILE	2.8
25	AX	47	VAL	2.8
33	BF	108	ASN	2.8
41	BN	14	VAL	2.8
4	AE	73	GLU	2.8
10	AN	56	ASP	2.8
6	AG	146	TYR	2.8
7	AH	145	ALA	2.8
27	D5	55	ARG	2.8
43	CP	122	LYS	2.8
48	BU	87	ARG	2.8
58	DL	86	LYS	2.8
4	AE	35	GLN	2.8
57	DY	115	GLN	2.8
9	AM	53	VAL	2.8
16	A1	64	ARG	2.8
21	AV	21	ALA	2.8
16	A1	88	ILE	2.8
7	AH	85	LYS	2.8
21	DV	182	LYS	2.8
52	BB	44	G	2.8
4	DE	21	VAL	2.8
15	AR	36	GLU	2.8
28	A6	40	CYS	2.8
4	AE	46	ALA	2.8
21	AV	18	LEU	2.8
48	BU	26	LEU	2.8
4	AE	34	VAL	2.8
20	AU	89	PHE	2.8
26	A4	68	ARG	2.8
42	BO	57	LYS	2.8
33	CF	104	GLN	2.7

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Mol	Chain	Res	Type	RSRZ
52	CD	17	C	2.8
28	D6	11	LEU	2.7
21	DV	172	ALA	2.7
36	BI	100	ASN	2.7
37	CJ	78	ARG	2.7
21	AV	139	VAL	2.7
4	AE	48	GLN	2.7
37	CJ	156	TRP	2.7
26	A4	56	VAL	2.7
33	CF	65	ALA	2.7
58	DL	78	ILE	2.7
12	DP	1	MET	2.7
32	BE	163	PHE	2.7
32	BE	229	VAL	2.7
52	BB	16	U	2.7
57	DY	80	VAL	2.7
21	AV	167	PRO	2.7
40	CM	33	GLN	2.7
6	AG	147	ASP	2.7
56	DI	21	LYS	2.7
27	A5	60	VAL	2.7
32	BE	70	PHE	2.7
49	BV	41	VAL	2.7
57	DY	77	PRO	2.7
4	AE	89	ASP	2.7
16	A1	111	GLU	2.7
1	AA	1092	C	2.7
8	DK	8	PRO	2.7
17	A2	26	ASP	2.7
7	AH	115	VAL	2.7
28	A6	27	LYS	2.7
32	BE	5	ILE	2.7
21	DV	196	VAL	2.7
42	CO	38	THR	2.7
4	AE	81	ILE	2.7
9	AM	6	PRO	2.7
40	CM	23	ILE	2.7
10	AN	22	ILE	2.7
4	AE	159	HIS	2.7
33	CF	103	VAL	2.7
1	AA	887	A	2.7
21	AV	24	LEU	2.7

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Mol	Chain	Res	Type	RSRZ
53	C1	55	U	2.7
58	DL	12	LEU	2.7
7	AH	104	GLU	2.7
45	BR	2	PRO	2.7
41	BN	21	ILE	2.7
6	AG	43	LEU	2.6
32	BE	215	LEU	2.6
52	BB	45	U	2.6
21	DV	145	GLU	2.6
36	BI	38	GLU	2.6
21	DV	144	LEU	2.6
4	AE	38	THR	2.6
21	DV	160	GLY	2.6
7	AH	132	ARG	2.6
1	AA	654(S)	G	2.6
21	AV	116	VAL	2.6
21	AV	119	GLU	2.6
26	A4	34	GLU	2.6
28	D6	30	THR	2.6
7	AH	71	LEU	2.6
19	AT	92	LEU	2.6
33	CF	39	ILE	2.6
56	DI	30	ALA	2.6
7	AH	13	LYS	2.6
31	BA	1030	C	2.6
32	CE	217	ARG	2.6
58	DL	15	GLY	2.6
21	AV	78	LYS	2.6
23	AZ	96	LYS	2.6
8	DK	117	GLU	2.6
12	AP	97	VAL	2.6
14	DQ	104	GLY	2.6
43	BP	88	ARG	2.6
8	AK	139	GLN	2.6
12	DP	89	ASN	2.6
17	A2	20	LEU	2.6
9	AM	136	GLU	2.6
17	A2	58	VAL	2.6
37	CJ	81	GLY	2.6
41	CN	80	VAL	2.6
33	CF	56	ASP	2.6
3	AD	236	GLY	2.6

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Mol	Chain	Res	Type	RSRZ
23	DZ	97	LEU	2.6
41	BN	128	ALA	2.6
43	BP	60	VAL	2.6
21	AV	28	MET	2.6
49	BV	25	LYS	2.6
52	BB	47	U	2.6
21	AV	20	ARG	2.6
26	D4	37	SER	2.6
34	CG	133	VAL	2.6
31	BA	1035	A	2.6
40	CM	4	ILE	2.6
7	AH	56	SER	2.6
22	A3	9	SER	2.6
7	AH	113	VAL	2.6
17	D2	37	VAL	2.6
33	CF	153	VAL	2.6
11	AO	148	LEU	2.6
14	DQ	110	LEU	2.6
32	CE	215	LEU	2.6
58	DL	99	ILE	2.6
1	AA	896	A	2.6
52	CD	14	A	2.6
7	AH	75	ALA	2.6
52	CB	20	U	2.6
56	DJ	15	ALA	2.6
21	DV	193	GLU	2.6
20	AU	44	ILE	2.6
57	DY	130	THR	2.6
26	D4	60	GLN	2.5
33	CF	60	ALA	2.5
21	AV	157	LEU	2.5
9	AM	16	ILE	2.5
26	A4	8	LYS	2.5
26	D4	28	LYS	2.5
21	AV	128	VAL	2.5
10	DN	122	LEU	2.5
25	AX	26	LEU	2.5
12	AP	68	ILE	2.5
41	BN	95	ILE	2.5
55	DA	2131	G	2.5
21	DV	121	HIS	2.5
9	AM	42	TRP	2.5

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Mol	Chain	Res	Type	RSRZ
16	A1	73	GLY	2.5
21	AV	93	ASP	2.5
21	AV	104	PHE	2.5
26	A4	47	GLN	2.5
33	BF	68	VAL	2.5
21	AV	5	LEU	2.5
16	A1	72	HIS	2.5
32	CE	222	ILE	2.5
3	AD	237	GLU	2.5
30	A8	63	PRO	2.5
56	DI	11	GLU	2.5
5	AF	22	ALA	2.5
28	D6	46	HIS	2.5
58	DL	136	VAL	2.5
40	CM	90	LEU	2.5
6	AG	75	LYS	2.5
6	DG	137	GLU	2.5
21	AV	74	VAL	2.5
21	AV	143	GLY	2.5
33	CF	76	VAL	2.5
37	BJ	65	ALA	2.5
45	CR	89	GLY	2.5
54	CA	1542	U	2.5
37	BJ	84	ASN	2.5
21	DV	192	ALA	2.5
39	BL	53	VAL	2.5
41	BN	39	PRO	2.5
57	DY	50	ARG	2.5
4	AE	177	PRO	2.5
7	AH	8	PRO	2.5
41	BN	33	THR	2.5
6	AG	62	LEU	2.5
33	CF	91	LEU	2.5
21	AV	55	HIS	2.5
41	CN	48	ILE	2.5
52	BB	11	C	2.5
23	AZ	61	ARG	2.5
26	D4	39	CYS	2.5
37	CJ	79	ARG	2.5
8	DK	75	LEU	2.5
58	DL	55	VAL	2.5
22	A3	85	ALA	2.5

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Mol	Chain	Res	Type	RSRZ
27	D5	52	TYR	2.5
42	CO	64	TYR	2.5
21	AV	110	GLY	2.4
32	CE	230	VAL	2.4
41	CN	82	VAL	2.4
12	AP	140	ALA	2.4
58	DL	128	ALA	2.4
21	AV	44	PHE	2.4
35	CH	152	ARG	2.4
17	A2	93	GLU	2.4
32	BE	231	GLU	2.4
4	AE	49	LEU	2.4
28	D6	38	LYS	2.4
43	CP	126	LYS	2.4
17	A2	27	ALA	2.4
55	DA	2132	U	2.4
26	A4	5	ILE	2.4
10	AN	11	ALA	2.4
41	CN	15	ALA	2.4
4	DE	69	LYS	2.4
8	DK	141	LYS	2.4
14	DQ	111	GLU	2.4
37	BJ	86	GLN	2.4
58	DL	36	GLU	2.4
55	DA	654(A)	A	2.4
17	A2	63	GLY	2.4
21	DV	151	HIS	2.4
7	DH	136	ILE	2.4
8	DK	74	ASN	2.4
37	BJ	88	PRO	2.4
3	AD	147	LEU	2.4
41	BN	84	VAL	2.4
52	CC	17	C	2.4
23	AZ	97	LEU	2.4
26	D4	64	GLY	2.4
58	DL	31	GLY	2.4
55	DA	886	C	2.4
15	DR	21	GLU	2.4
41	BN	99	GLN	2.4
6	AG	112	PRO	2.4
47	BT	100	LYS	2.4
1	AA	2897	U	2.4

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Mol	Chain	Res	Type	RSRZ
16	A1	113	ALA	2.4
21	AV	9	TYR	2.4
1	AA	888	C	2.4
6	AG	52	ILE	2.4
32	BE	217	ARG	2.4
52	CB	7	A	2.4
25	AX	54	VAL	2.4
11	DO	144	GLU	2.4
21	DV	63	ASP	2.4
1	AA	898	C	2.4
9	AM	52	VAL	2.4
14	DQ	83	LYS	2.4
25	DX	60	GLU	2.4
26	D4	32	TYR	2.4
36	BI	71	ARG	2.4
37	CJ	84	ASN	2.4
22	A3	2	ALA	2.4
42	BO	68	ALA	2.4
7	AH	137	ASP	2.4
22	D3	5	LYS	2.4
56	DI	4	ASP	2.4
12	AP	19	GLY	2.4
12	AP	24	GLY	2.4
17	A2	73	SER	2.4
26	D4	33	VAL	2.4
21	AV	14	LYS	2.4
33	BF	149	ALA	2.4
58	DL	135	GLY	2.4
58	DL	75	SER	2.4
28	D6	16	CYS	2.4
32	BE	46	LYS	2.4
42	BO	127	GLU	2.4
58	DL	143	GLU	2.4
16	A1	80	ILE	2.4
20	AU	65	ALA	2.4
41	BN	19	ALA	2.4
53	C1	29	G	2.3
1	AA	2138	C	2.3
4	DE	72	VAL	2.3
32	BE	7	VAL	2.3
11	AO	1	MET	2.3
1	AA	1094	U	2.3

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Mol	Chain	Res	Type	RSRZ
33	BF	144	SER	2.3
34	BG	152	SER	2.3
6	AG	142	PRO	2.3
15	AR	37	GLY	2.3
37	BJ	153	HIS	2.3
21	DV	178	GLU	2.3
33	CF	89	GLU	2.3
12	AP	66	ILE	2.3
19	AT	69	TYR	2.3
39	BL	4	TYR	2.3
41	CN	98	LEU	2.3
21	AV	147	GLY	2.3
48	BU	86	VAL	2.3
17	A2	64	HIS	2.3
1	AA	2139	C	2.3
6	DG	88	ILE	2.3
7	AH	169	VAL	2.3
34	CG	145	GLU	2.3
36	BI	31	GLU	2.3
10	AN	19	ILE	2.3
33	BF	87	LEU	2.3
48	BU	24	ALA	2.3
48	BU	34	TYR	2.3
7	DH	125	VAL	2.3
48	CU	18	ARG	2.3
6	AG	40	ASN	2.3
6	AG	108	ASN	2.3
8	DK	76	THR	2.3
12	AP	22	LYS	2.3
21	AV	117	LEU	2.3
25	AX	20	LYS	2.3
41	BN	31	THR	2.3
26	A4	35	VAL	2.3
1	AA	1100	C	2.3
4	AE	87	GLU	2.3
43	BP	11	ARG	2.3
52	CB	47	U	2.3
52	CC	44	G	2.3
16	A1	115	ALA	2.3
52	BB	9	A	2.3
20	DU	102	CYS	2.3
21	AV	91	LEU	2.3

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Mol	Chain	Res	Type	RSRZ
31	BA	1129	C	2.3
4	AE	36	ARG	2.3
5	AF	18	ARG	2.3
33	CF	169	ALA	2.3
50	BW	106	ALA	2.3
28	A6	36	LEU	2.3
7	AH	121	ILE	2.3
16	A1	104	GLN	2.3
12	DP	20	ALA	2.3
52	BB	4	C	2.3
58	DL	39	LYS	2.3
58	DL	56	GLU	2.3
21	AV	125	LEU	2.3
21	AV	159	PRO	2.3
43	BP	82	MET	2.3
34	CG	152	SER	2.3
14	DQ	76	LYS	2.3
7	AH	54	ARG	2.3
15	AR	34	VAL	2.3
55	DA	890	A	2.3
17	A2	34	GLU	2.2
7	AH	87	LEU	2.2
21	AV	59	LEU	2.2
21	AV	81	ARG	2.2
25	AX	44	ARG	2.2
37	CJ	140	ASP	2.2
57	DY	16	ASN	2.2
53	C1	53	U	2.2
8	DK	71	ILE	2.2
58	DL	22	PRO	2.2
33	CF	51	GLY	2.2
28	A6	29	ASN	2.2
55	DA	2151	G	2.2
20	DU	53	PRO	2.2
8	AK	78	THR	2.2
33	BF	102	ASN	2.2
6	AG	111	LEU	2.2
6	AG	64	THR	2.2
57	DY	89	ALA	2.2
7	DH	113	VAL	2.2
17	A2	37	VAL	2.2
17	A2	47	VAL	2.2

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Mol	Chain	Res	Type	RSRZ
20	AU	2	ARG	2.2
21	DV	171	ILE	2.2
52	BB	2	C	2.2
52	CB	13	C	2.2
36	BI	6	VAL	2.2
15	AR	40	THR	2.2
21	AV	25	PRO	2.2
6	AG	60	LEU	2.2
55	DA	654(U)	A	2.2
57	DY	26	LEU	2.2
7	AH	90	LYS	2.2
42	BO	59	ARG	2.2
12	AP	30	GLY	2.2
7	AH	117	PRO	2.2
32	CE	160	ASP	2.2
37	CJ	16	LEU	2.2
17	A2	96	ILE	2.2
40	BM	34	VAL	2.2
56	DJ	24	ILE	2.2
6	AG	69	ALA	2.2
17	A2	12	TYR	2.2
57	DY	136	ALA	2.2
6	AG	90	LEU	2.2
8	AK	77	LEU	2.2
12	AP	29	PHE	2.2
32	BE	14	GLY	2.2
1	AA	1096	A	2.2
4	DE	62	PRO	2.2
21	DV	98	MET	2.2
57	DY	118	THR	2.2
33	CF	82	GLU	2.2
42	CO	79	GLU	2.2
22	A3	7	LEU	2.2
42	BO	41	ARG	2.2
57	DY	29	TYR	2.2
58	DL	131	ALA	2.2
52	CB	23	A	2.2
56	DJ	25	ASP	2.2
41	BN	80	VAL	2.2
42	CO	40	VAL	2.2
42	CO	67	THR	2.2
1	AA	2146	C	2.2

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Mol	Chain	Res	Type	RSRZ
52	CB	69	G	2.2
32	CE	68	ILE	2.2
33	BF	56	ASP	2.2
9	AM	7	LYS	2.2
8	DK	123	LEU	2.1
17	A2	94	LEU	2.1
33	BF	101	LEU	2.1
1	AA	897	C	2.1
33	CF	100	ALA	2.1
33	BF	20	SER	2.1
1	AA	2132	U	2.1
52	CD	47	U	2.1
10	AN	58	VAL	2.1
41	BN	92	GLU	2.1
10	AN	91	LEU	2.1
17	A2	51	VAL	2.1
18	AS	109	GLU	2.1
32	CE	214	ILE	2.1
4	AE	10	GLY	2.1
8	AK	12	LEU	2.1
42	CO	61	THR	2.1
34	CG	146	ILE	2.1
41	BN	30	VAL	2.1
34	BG	37	PRO	2.1
21	AV	67	LEU	2.1
9	AM	56	ASN	2.1
47	BT	11	VAL	2.1
1	AA	654(A)	A	2.1
1	AA	2141	G	2.1
1	AA	2155	G	2.1
7	AH	22	GLY	2.1
17	A2	16	PRO	2.1
5	AF	21	ALA	2.1
6	AG	63	ILE	2.1
7	AH	97	ARG	2.1
9	AM	119	ARG	2.1
1	AA	1070	A	2.1
21	AV	136	PHE	2.1
37	BJ	62	PHE	2.1
16	A1	110	VAL	2.1
22	A3	3	HIS	2.1
35	BH	155	GLU	2.1

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Mol	Chain	Res	Type	RSRZ
19	AT	89	ILE	2.1
33	CF	67	THR	2.1
55	DA	2140	C	2.1
34	BG	169	LYS	2.1
11	DO	91	PHE	2.1
2	DB	1(M)	A	2.1
1	AA	889	C	2.1
1	AA	2792	G	2.1
22	A3	12	ASN	2.1
29	D7	48	LYS	2.1
1	AA	2803	C	2.1
52	BB	56	C	2.1
14	AQ	107	GLU	2.1
4	AE	41	LYS	2.1
15	AR	35	LYS	2.1
20	DU	49	VAL	2.1
5	DF	24	LEU	2.1
12	AP	92	GLY	2.1
22	D3	84	LEU	2.1
32	CE	228	GLY	2.1
52	BC	76	A	2.1
57	DY	92	THR	2.1
58	DL	67	PHE	2.1
9	AM	98	VAL	2.1
12	AP	102	VAL	2.1
4	AE	78	LEU	2.1
23	AZ	36	GLY	2.1
36	BI	34	GLY	2.1
6	DG	181	ARG	2.1
17	A2	3	ALA	2.1
27	A5	47	PRO	2.1
33	BF	60	ALA	2.1
58	DL	139	VAL	2.1
6	DG	52	ILE	2.1
33	BF	57	ILE	2.1
10	AN	25	LEU	2.1
55	DA	654(S)	G	2.1
21	DV	23	LYS	2.1
21	DV	108	PRO	2.1
29	A7	46	VAL	2.1
1	AA	2820	A	2.1
4	AE	3	GLY	2.1

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Mol	Chain	Res	Type	RSRZ
20	DU	52	SER	2.1
41	CN	101	SER	2.1
40	CM	89	ASP	2.1
18	AS	30	GLU	2.0
20	DU	50	ARG	2.0
28	A6	52	VAL	2.0
33	CF	201	TYR	2.0
42	CO	65	GLU	2.0
10	AN	38	VAL	2.0
39	BL	21	PRO	2.0
42	CO	128	ALA	2.0
57	DY	119	ALA	2.0
10	AN	122	LEU	2.0
26	A4	26	SER	2.0
55	DA	654(T)	A	2.0
46	CS	84	ALA	2.0
58	DL	53	VAL	2.0
10	AN	34	THR	2.0
12	AP	64	ILE	2.0
17	A2	25	LEU	2.0
1	AA	2157	G	2.0
31	BA	1032(B)	G	2.0
52	BD	5	G	2.0
52	CB	10	G	2.0
6	AG	181	ARG	2.0
42	CO	59	ARG	2.0
47	CT	101	ARG	2.0
9	AM	50	ASP	2.0
21	AV	63	ASP	2.0
21	AV	135	GLU	2.0
32	CE	95	GLN	2.0
25	AX	15	TYR	2.0
16	A1	74	LEU	2.0
26	A4	69	LYS	2.0
49	CV	15	LEU	2.0
40	BM	10	GLY	2.0
41	BN	86	GLY	2.0
41	CN	127	LYS	2.0
41	BN	91	ARG	2.0
49	BV	68	GLY	2.0
33	BF	105	GLU	2.0
6	AG	149	VAL	2.0

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Mol	Chain	Res	Type	RSRZ
21	AV	100	VAL	2.0
27	A5	51	TYR	2.0
32	BE	156	LYS	2.0
36	BI	37	VAL	2.0
41	BN	50	TYR	2.0
57	DY	76	GLY	2.0
7	AH	119	GLU	2.0
7	AH	153	LYS	2.0
9	AM	137	LYS	2.0
31	BA	841	U	2.0
21	DV	174	VAL	2.0
26	D4	63	TYR	2.0
40	CM	24	VAL	2.0
41	CN	95	ILE	2.0
11	AO	91	PHE	2.0
31	BA	1033	G	2.0
5	AF	175	THR	2.0
7	AH	159	GLU	2.0
10	AN	96	THR	2.0
56	DJ	10	GLU	2.0
55	DA	2145	C	2.0
4	AE	184	VAL	2.0
9	AM	87	LEU	2.0
33	CF	130	VAL	2.0
36	BI	36	ARG	2.0
36	BI	72	VAL	2.0
57	DY	81	VAL	2.0
58	DL	4	VAL	2.0

6.2 Non-standard residues in protein, DNA, RNA chains

In the following table, the Atoms column lists the number of modelled atoms in the group and the number defined in the chemical component dictionary. The B-factors column lists the minimum, median, 95th percentile and maximum values of B factors of atoms in the group. The column labelled 'Q< 0.9' lists the number of atoms with occupancy less than 0.9.

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(Å ²)	Q<0.9
52	MIA	BD	37	29/30	0.89	0.21	107,111,115,115	0
52	MIA	CD	37	29/30	0.92	0.27	106,108,109,110	0
52	MIA	BB	37	29/30	0.94	0.19	83,88,98,99	0
52	MIA	BC	37	29/30	0.95	0.19	70,75,80,81	0
52	MIA	CB	37	29/30	0.96	0.23	53,57,62,63	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
52	MIA	CC	37	29/30	0.97	0.21	49,53,60,61	0

6.3 Carbohydrates [i](#)

There are no carbohydrates in this entry.

6.4 Ligands [i](#)

In the following table, the Atoms column lists the number of modelled atoms in the group and the number defined in the chemical component dictionary. The B-factors column lists the minimum, median, 95th percentile and maximum values of B factors of atoms in the group. The column labelled 'Q< 0.9' lists the number of atoms with occupancy less than 0.9.

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3647	1/1	-0.04	0.80	108,108,108,108	0
59	MG	BA	1725	1/1	0.09	0.31	109,109,109,109	0
59	MG	BA	2128	1/1	0.10	0.15	108,108,108,108	0
59	MG	BA	1821	1/1	0.15	0.19	93,93,93,93	0
59	MG	BA	1857	1/1	0.18	0.27	122,122,122,122	0
59	MG	AW	101	1/1	0.18	0.64	97,97,97,97	0
59	MG	BD	116	1/1	0.23	0.37	129,129,129,129	0
59	MG	DA	4641	1/1	0.25	0.68	134,134,134,134	0
59	MG	DA	3611	1/1	0.27	0.58	115,115,115,115	0
59	MG	CA	1685	1/1	0.28	0.28	80,80,80,80	0
59	MG	AA	4130	1/1	0.29	0.25	107,107,107,107	0
59	MG	AA	4134	1/1	0.29	0.11	117,117,117,117	0
59	MG	CA	2062	1/1	0.30	0.28	85,85,85,85	0
59	MG	AA	3073	1/1	0.30	0.34	71,71,71,71	0
59	MG	AA	3402	1/1	0.30	0.17	66,66,66,66	0
59	MG	BA	2150	1/1	0.31	0.29	96,96,96,96	0
59	MG	CA	1856	1/1	0.31	0.23	81,81,81,81	0
59	MG	DA	3989	1/1	0.33	0.12	110,110,110,110	0
59	MG	CB	114	1/1	0.34	0.16	71,71,71,71	0
59	MG	BA	1724	1/1	0.34	0.34	100,100,100,100	0
59	MG	BA	1866	1/1	0.35	0.32	111,111,111,111	0
59	MG	CA	1934	1/1	0.36	0.15	110,110,110,110	0
59	MG	DA	4733	1/1	0.37	0.30	84,84,84,84	0
59	MG	CA	1852	1/1	0.37	0.29	143,143,143,143	0
59	MG	CB	121	1/1	0.37	0.15	101,101,101,101	0
59	MG	AA	4042	1/1	0.38	0.19	88,88,88,88	0
59	MG	BD	111	1/1	0.38	0.15	74,74,74,74	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4698	1/1	0.39	0.82	104,104,104,104	0
59	MG	DA	4129	1/1	0.39	0.09	109,109,109,109	0
59	MG	BA	2187	1/1	0.39	0.24	107,107,107,107	0
59	MG	CA	1887	1/1	0.39	0.32	92,92,92,92	0
59	MG	CA	1970	1/1	0.40	0.21	100,100,100,100	0
59	MG	CA	2241	1/1	0.40	0.23	98,98,98,98	0
59	MG	DA	4433	1/1	0.41	0.17	90,90,90,90	0
59	MG	BE	305	1/1	0.41	0.21	95,95,95,95	0
59	MG	AA	4133	1/1	0.41	0.50	127,127,127,127	0
59	MG	BA	1828	1/1	0.41	0.46	114,114,114,114	0
59	MG	DA	3757	1/1	0.41	0.89	128,128,128,128	0
59	MG	DV	301	1/1	0.41	0.38	108,108,108,108	0
59	MG	BA	1662	1/1	0.42	0.18	91,91,91,91	0
59	MG	DA	4627	1/1	0.42	0.23	80,80,80,80	0
59	MG	BA	1798	1/1	0.42	0.34	93,93,93,93	0
59	MG	AA	4046	1/1	0.42	0.18	111,111,111,111	0
59	MG	BA	1917	1/1	0.43	0.17	64,64,64,64	0
59	MG	BA	2275	1/1	0.44	0.17	145,145,145,145	0
59	MG	CA	2304	1/1	0.44	0.57	124,124,124,124	0
59	MG	DA	3573	1/1	0.45	0.17	84,84,84,84	0
59	MG	DA	4572	1/1	0.45	1.39	76,76,76,76	0
59	MG	AA	3423	1/1	0.45	0.48	99,99,99,99	0
59	MG	DA	3367	1/1	0.45	0.63	91,91,91,91	0
59	MG	CA	2122	1/1	0.45	0.26	123,123,123,123	0
59	MG	AA	3376	1/1	0.45	0.31	116,116,116,116	0
59	MG	CB	116	1/1	0.45	0.23	89,89,89,89	0
59	MG	AB	209	1/1	0.45	0.17	104,104,104,104	0
59	MG	AA	3848	1/1	0.46	0.23	90,90,90,90	0
59	MG	BA	2107	1/1	0.46	0.18	83,83,83,83	0
59	MG	CD	130	1/1	0.46	0.23	84,84,84,84	0
59	MG	CA	1703	1/1	0.46	0.19	49,49,49,49	0
59	MG	DA	4826	1/1	0.46	0.34	71,71,71,71	0
59	MG	DA	4995	1/1	0.46	0.07	121,121,121,121	0
59	MG	CA	1946	1/1	0.46	0.45	112,112,112,112	0
59	MG	AQ	205	1/1	0.47	0.48	94,94,94,94	0
59	MG	BD	109	1/1	0.47	0.39	139,139,139,139	0
59	MG	AA	4075	1/1	0.47	0.61	132,132,132,132	0
59	MG	CA	2314	1/1	0.47	0.30	92,92,92,92	0
59	MG	BA	1929	1/1	0.47	0.09	85,85,85,85	0
59	MG	CA	1925	1/1	0.47	0.51	115,115,115,115	0
59	MG	DA	4535	1/1	0.47	0.38	89,89,89,89	0
59	MG	DA	4951	1/1	0.48	0.34	102,102,102,102	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	2019	1/1	0.48	0.24	68,68,68,68	0
59	MG	DA	5015	1/1	0.48	0.58	102,102,102,102	0
59	MG	AA	3781	1/1	0.48	0.15	104,104,104,104	0
59	MG	DA	3628	1/1	0.48	0.35	92,92,92,92	0
59	MG	CA	2121	1/1	0.49	0.26	125,125,125,125	0
59	MG	BB	106	1/1	0.49	0.40	116,116,116,116	0
59	MG	AA	3942	1/1	0.49	0.25	85,85,85,85	0
59	MG	CA	2287	1/1	0.49	0.28	119,119,119,119	0
59	MG	CA	1955	1/1	0.49	0.18	58,58,58,58	0
59	MG	AA	3667	1/1	0.50	0.28	79,79,79,79	0
59	MG	BA	2062	1/1	0.50	0.21	75,75,75,75	0
59	MG	AA	3442	1/1	0.50	0.08	128,128,128,128	0
59	MG	AA	3903	1/1	0.50	0.19	95,95,95,95	0
59	MG	CA	2167	1/1	0.50	0.22	97,97,97,97	0
59	MG	CT	202	1/1	0.50	0.23	85,85,85,85	0
59	MG	DA	4871	1/1	0.51	0.32	86,86,86,86	0
59	MG	DB	209	1/1	0.51	0.57	77,77,77,77	0
59	MG	CS	106	1/1	0.51	0.33	77,77,77,77	0
59	MG	CD	126	1/1	0.51	0.20	88,88,88,88	0
59	MG	BA	2149	1/1	0.51	0.17	82,82,82,82	0
59	MG	DA	4783	1/1	0.51	0.38	93,93,93,93	0
59	MG	DA	3678	1/1	0.51	0.49	123,123,123,123	0
59	MG	CA	2295	1/1	0.52	0.19	110,110,110,110	0
59	MG	BC	116	1/1	0.52	0.20	107,107,107,107	0
59	MG	BA	2033	1/1	0.52	0.15	94,94,94,94	0
59	MG	AA	3904	1/1	0.52	0.18	77,77,77,77	0
59	MG	CA	2236	1/1	0.52	0.29	104,104,104,104	0
59	MG	AA	3852	1/1	0.52	0.42	67,67,67,67	0
59	MG	BA	2215	1/1	0.52	0.24	79,79,79,79	0
59	MG	DA	4115	1/1	0.52	0.35	81,81,81,81	0
59	MG	DA	4927	1/1	0.52	0.21	160,160,160,160	0
59	MG	DA	5058	1/1	0.52	0.43	103,103,103,103	0
59	MG	AA	3657	1/1	0.52	0.36	89,89,89,89	0
59	MG	BA	2043	1/1	0.53	0.17	109,109,109,109	0
59	MG	DA	3530	1/1	0.53	0.25	73,73,73,73	0
59	MG	BQ	101	1/1	0.53	0.34	66,66,66,66	0
59	MG	BA	1858	1/1	0.53	0.34	141,141,141,141	0
59	MG	AA	3361	1/1	0.53	0.19	93,93,93,93	0
59	MG	BA	2198	1/1	0.53	0.76	139,139,139,139	0
59	MG	AA	3717	1/1	0.53	0.42	106,106,106,106	0
59	MG	DA	3418	1/1	0.53	0.23	60,60,60,60	0
59	MG	DA	4593	1/1	0.53	0.28	65,65,65,65	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4007	1/1	0.53	0.14	110,110,110,110	0
59	MG	DA	5042	1/1	0.53	0.39	106,106,106,106	0
59	MG	AA	3113	1/1	0.53	0.21	85,85,85,85	0
59	MG	AA	4137	1/1	0.54	0.26	90,90,90,90	0
59	MG	BA	2213	1/1	0.54	0.47	114,114,114,114	0
59	MG	CA	2269	1/1	0.54	0.17	71,71,71,71	0
59	MG	BA	1985	1/1	0.54	0.16	105,105,105,105	0
59	MG	CA	1750	1/1	0.54	0.20	82,82,82,82	0
59	MG	DA	5067	1/1	0.54	0.32	119,119,119,119	0
59	MG	BA	2099	1/1	0.54	0.13	67,67,67,67	0
59	MG	DA	4973	1/1	0.54	0.51	115,115,115,115	0
59	MG	AA	3828	1/1	0.55	0.34	113,113,113,113	0
59	MG	AA	3303	1/1	0.55	0.24	94,94,94,94	0
59	MG	BA	1710	1/1	0.55	0.34	90,90,90,90	0
59	MG	BA	1788	1/1	0.55	0.18	53,53,53,53	0
59	MG	BA	1644	1/1	0.55	0.39	102,102,102,102	0
59	MG	AA	3819	1/1	0.55	0.26	94,94,94,94	0
59	MG	D0	203	1/1	0.55	0.57	97,97,97,97	0
59	MG	BA	2097	1/1	0.55	0.31	94,94,94,94	0
59	MG	CD	129	1/1	0.56	0.25	95,95,95,95	0
59	MG	DB	269	1/1	0.56	0.44	91,91,91,91	0
59	MG	CD	104	1/1	0.56	0.41	74,74,74,74	0
59	MG	CA	2132	1/1	0.56	0.16	80,80,80,80	0
59	MG	BA	1840	1/1	0.56	0.14	156,156,156,156	0
59	MG	AA	3567	1/1	0.56	0.18	82,82,82,82	0
59	MG	DA	4905	1/1	0.56	0.27	90,90,90,90	0
59	MG	DA	5077	1/1	0.56	0.47	102,102,102,102	0
59	MG	AA	4019	1/1	0.56	0.39	99,99,99,99	0
59	MG	AA	3868	1/1	0.56	0.19	80,80,80,80	0
59	MG	AA	3888	1/1	0.56	0.17	79,79,79,79	0
59	MG	DA	4716	1/1	0.56	0.10	77,77,77,77	0
59	MG	AA	3872	1/1	0.57	0.36	90,90,90,90	0
59	MG	AA	3869	1/1	0.57	0.27	68,68,68,68	0
59	MG	AA	3894	1/1	0.57	0.32	86,86,86,86	0
59	MG	DA	4952	1/1	0.57	0.39	86,86,86,86	0
59	MG	BA	1956	1/1	0.57	0.59	111,111,111,111	0
59	MG	AA	3408	1/1	0.57	0.29	61,61,61,61	0
59	MG	CA	2076	1/1	0.57	0.26	113,113,113,113	0
59	MG	AA	3059	1/1	0.57	0.23	68,68,68,68	0
59	MG	CA	2250	1/1	0.58	0.27	88,88,88,88	0
59	MG	CA	1694	1/1	0.58	0.17	56,56,56,56	0
59	MG	CA	2284	1/1	0.58	0.15	85,85,85,85	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	4009	1/1	0.58	0.17	99,99,99,99	0
59	MG	DA	3598	1/1	0.58	0.47	116,116,116,116	0
59	MG	BA	1899	1/1	0.58	0.14	57,57,57,57	0
59	MG	BA	2258	1/1	0.58	0.16	77,77,77,77	0
59	MG	CA	1848	1/1	0.58	0.28	100,100,100,100	0
59	MG	BA	2170	1/1	0.58	0.38	151,151,151,151	0
59	MG	BA	1804	1/1	0.58	0.09	105,105,105,105	0
59	MG	AA	3305	1/1	0.59	0.18	47,47,47,47	0
59	MG	CA	1717	1/1	0.59	0.21	67,67,67,67	0
59	MG	BA	2052	1/1	0.59	0.14	128,128,128,128	0
59	MG	CD	125	1/1	0.59	0.31	88,88,88,88	0
59	MG	DA	3955	1/1	0.59	0.37	120,120,120,120	0
59	MG	DA	4728	1/1	0.59	0.47	86,86,86,86	0
59	MG	AQ	204	1/1	0.59	0.13	80,80,80,80	0
59	MG	BA	1756	1/1	0.59	0.16	69,69,69,69	0
59	MG	BB	111	1/1	0.59	0.09	90,90,90,90	0
59	MG	DA	4842	1/1	0.59	0.29	65,65,65,65	0
59	MG	DA	4045	1/1	0.59	0.09	134,134,134,134	0
59	MG	DA	4136	1/1	0.59	0.29	47,47,47,47	0
59	MG	BA	2112	1/1	0.59	0.25	101,101,101,101	0
59	MG	AA	3807	1/1	0.59	0.17	87,87,87,87	0
59	MG	BA	2167	1/1	0.59	0.18	101,101,101,101	0
59	MG	CC	105	1/1	0.60	0.24	71,71,71,71	0
59	MG	DA	3375	1/1	0.60	0.37	64,64,64,64	0
59	MG	CA	1921	1/1	0.60	0.24	52,52,52,52	0
59	MG	DA	3984	1/1	0.60	0.40	117,117,117,117	0
59	MG	AA	4070	1/1	0.60	0.18	60,60,60,60	0
59	MG	BA	2247	1/1	0.60	0.23	82,82,82,82	0
59	MG	CA	1774	1/1	0.60	0.17	78,78,78,78	0
59	MG	BA	2230	1/1	0.60	0.28	111,111,111,111	0
59	MG	CA	1676	1/1	0.60	0.22	89,89,89,89	0
59	MG	BA	2137	1/1	0.60	0.12	119,119,119,119	0
59	MG	DF	320	1/1	0.60	0.24	58,58,58,58	0
59	MG	DA	4779	1/1	0.60	0.52	92,92,92,92	0
59	MG	DA	4672	1/1	0.60	0.36	95,95,95,95	0
59	MG	DA	4490	1/1	0.60	0.07	125,125,125,125	0
59	MG	DA	4949	1/1	0.60	0.15	71,71,71,71	0
59	MG	DA	4321	1/1	0.60	0.28	70,70,70,70	0
59	MG	DA	3535	1/1	0.60	0.24	69,69,69,69	0
59	MG	BA	1931	1/1	0.60	0.22	68,68,68,68	0
59	MG	CB	108	1/1	0.61	0.19	87,87,87,87	0
59	MG	BD	102	1/1	0.61	0.24	84,84,84,84	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3998	1/1	0.61	0.23	86,86,86,86	0
59	MG	DA	3904	1/1	0.61	0.14	105,105,105,105	0
59	MG	AA	4107	1/1	0.61	0.49	98,98,98,98	0
59	MG	AA	3570	1/1	0.61	0.22	78,78,78,78	0
59	MG	CA	1994	1/1	0.61	0.30	94,94,94,94	0
59	MG	CA	2114	1/1	0.61	0.34	111,111,111,111	0
59	MG	DA	5054	1/1	0.61	0.71	67,67,67,67	0
59	MG	AB	233	1/1	0.61	0.22	86,86,86,86	0
59	MG	BA	1663	1/1	0.61	0.16	94,94,94,94	0
59	MG	AB	226	1/1	0.61	0.14	88,88,88,88	0
59	MG	BA	1817	1/1	0.61	0.15	63,63,63,63	0
59	MG	AA	3898	1/1	0.61	0.45	112,112,112,112	0
59	MG	DA	4803	1/1	0.61	0.29	82,82,82,82	0
59	MG	AA	3591	1/1	0.61	0.22	106,106,106,106	0
59	MG	AA	3922	1/1	0.61	0.46	65,65,65,65	0
59	MG	AA	4041	1/1	0.61	0.28	101,101,101,101	0
59	MG	BA	2115	1/1	0.61	0.27	118,118,118,118	0
59	MG	CA	2290	1/1	0.61	0.26	103,103,103,103	0
59	MG	BA	1782	1/1	0.61	0.55	100,100,100,100	0
59	MG	DB	276	1/1	0.62	0.21	66,66,66,66	0
59	MG	CB	107	1/1	0.62	0.24	106,106,106,106	0
59	MG	AA	3403	1/1	0.62	0.23	60,60,60,60	0
59	MG	CA	2263	1/1	0.62	0.39	98,98,98,98	0
59	MG	BA	2138	1/1	0.62	0.18	74,74,74,74	0
59	MG	AA	3595	1/1	0.62	0.36	103,103,103,103	0
59	MG	DA	4681	1/1	0.62	0.43	118,118,118,118	0
59	MG	DA	4701	1/1	0.62	0.30	64,64,64,64	0
59	MG	BA	2036	1/1	0.62	0.27	86,86,86,86	0
59	MG	BA	1772	1/1	0.62	0.40	75,75,75,75	0
59	MG	DU	211	1/1	0.62	0.26	116,116,116,116	0
59	MG	CA	2253	1/1	0.62	0.30	85,85,85,85	0
59	MG	BA	1696	1/1	0.62	0.17	73,73,73,73	0
59	MG	AB	229	1/1	0.62	0.14	67,67,67,67	0
59	MG	BA	2037	1/1	0.62	0.12	132,132,132,132	0
59	MG	BA	2010	1/1	0.62	0.17	61,61,61,61	0
59	MG	DA	4700	1/1	0.63	0.26	75,75,75,75	0
59	MG	DA	3793	1/1	0.63	0.27	81,81,81,81	0
59	MG	BA	2243	1/1	0.63	0.55	114,114,114,114	0
59	MG	AA	4128	1/1	0.63	0.18	116,116,116,116	0
59	MG	DA	3773	1/1	0.63	0.23	70,70,70,70	0
59	MG	BF	301	1/1	0.63	0.17	104,104,104,104	0
59	MG	AA	3685	1/1	0.63	0.28	65,65,65,65	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3218	1/1	0.63	0.24	65,65,65,65	0
59	MG	CD	118	1/1	0.63	0.50	97,97,97,97	0
59	MG	AA	3565	1/1	0.63	0.36	89,89,89,89	0
59	MG	AA	3118	1/1	0.63	0.13	52,52,52,52	0
59	MG	BA	2212	1/1	0.63	0.21	91,91,91,91	0
59	MG	CK	207	1/1	0.63	0.30	80,80,80,80	0
59	MG	DA	4082	1/1	0.63	0.56	75,75,75,75	0
59	MG	CE	305	1/1	0.63	0.27	95,95,95,95	0
59	MG	AA	3824	1/1	0.64	0.21	94,94,94,94	0
59	MG	CA	2051	1/1	0.64	0.32	89,89,89,89	0
59	MG	DA	4780	1/1	0.64	0.17	62,62,62,62	0
59	MG	BA	1988	1/1	0.64	0.40	95,95,95,95	0
59	MG	CA	2057	1/1	0.64	0.15	111,111,111,111	0
59	MG	CT	201	1/1	0.64	0.23	99,99,99,99	0
59	MG	CA	1684	1/1	0.64	0.26	103,103,103,103	0
60	ZN	A4	101	1/1	0.64	0.59	200,200,200,200	0
59	MG	AA	3174	1/1	0.64	0.26	63,63,63,63	0
59	MG	DA	4809	1/1	0.64	0.47	59,59,59,59	0
59	MG	DA	4746	1/1	0.64	0.18	98,98,98,98	0
59	MG	CA	1799	1/1	0.64	0.19	85,85,85,85	0
59	MG	BA	2042	1/1	0.64	0.09	95,95,95,95	0
59	MG	DA	4657	1/1	0.64	0.24	58,58,58,58	0
59	MG	AA	3160	1/1	0.64	0.21	64,64,64,64	0
59	MG	AA	3447	1/1	0.64	0.17	71,71,71,71	0
59	MG	DA	4338	1/1	0.64	0.46	80,80,80,80	0
59	MG	AA	3340	1/1	0.64	0.15	90,90,90,90	0
59	MG	AA	4150	1/1	0.64	0.22	82,82,82,82	0
59	MG	BE	301	1/1	0.65	0.15	63,63,63,63	0
59	MG	AA	3290	1/1	0.65	0.20	78,78,78,78	0
59	MG	BA	2029	1/1	0.65	0.16	76,76,76,76	0
59	MG	AA	4044	1/1	0.65	0.42	93,93,93,93	0
59	MG	CA	1700	1/1	0.65	0.18	39,39,39,39	0
59	MG	CA	2065	1/1	0.65	0.45	87,87,87,87	0
59	MG	DA	3503	1/1	0.65	0.20	71,71,71,71	0
59	MG	BA	1632	1/1	0.65	0.33	68,68,68,68	0
59	MG	DA	4492	1/1	0.65	0.34	66,66,66,66	0
59	MG	DA	3283	1/1	0.65	0.36	87,87,87,87	0
59	MG	AA	3254	1/1	0.65	0.13	76,76,76,76	0
59	MG	CB	115	1/1	0.65	0.39	97,97,97,97	0
59	MG	AA	3878	1/1	0.65	0.21	84,84,84,84	0
59	MG	DB	261	1/1	0.65	0.35	91,91,91,91	0
59	MG	AA	3908	1/1	0.65	0.66	68,68,68,68	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4829	1/1	0.65	0.29	106,106,106,106	0
59	MG	DA	3575	1/1	0.65	0.51	112,112,112,112	0
59	MG	AA	3753	1/1	0.65	0.17	91,91,91,91	0
59	MG	AB	225	1/1	0.65	0.16	125,125,125,125	0
59	MG	DV	305	1/1	0.65	0.35	86,86,86,86	0
59	MG	AA	4099	1/1	0.65	0.30	96,96,96,96	0
59	MG	BL	201	1/1	0.66	0.20	86,86,86,86	0
59	MG	DS	207	1/1	0.66	0.35	109,109,109,109	0
59	MG	BA	1906	1/1	0.66	0.22	107,107,107,107	0
59	MG	DA	5041	1/1	0.66	0.39	57,57,57,57	0
59	MG	AA	3324	1/1	0.66	0.17	43,43,43,43	0
59	MG	CA	2002	1/1	0.66	0.09	112,112,112,112	0
59	MG	DA	3490	1/1	0.66	0.36	64,64,64,64	0
59	MG	AA	3668	1/1	0.66	0.23	71,71,71,71	0
59	MG	CA	1960	1/1	0.66	0.11	91,91,91,91	0
59	MG	AA	4004	1/1	0.66	0.28	78,78,78,78	0
59	MG	DA	3581	1/1	0.66	0.32	91,91,91,91	0
59	MG	CA	2103	1/1	0.66	0.27	111,111,111,111	0
59	MG	DA	4933	1/1	0.66	0.32	64,64,64,64	0
59	MG	CA	2318	1/1	0.66	0.18	112,112,112,112	0
59	MG	CG	312	1/1	0.66	0.29	104,104,104,104	0
59	MG	DA	4735	1/1	0.66	0.37	88,88,88,88	0
59	MG	AA	3975	1/1	0.66	0.21	76,76,76,76	0
59	MG	BA	2124	1/1	0.66	0.18	72,72,72,72	0
59	MG	AA	3508	1/1	0.66	0.21	61,61,61,61	0
59	MG	DA	3649	1/1	0.66	0.18	71,71,71,71	0
59	MG	AA	3554	1/1	0.66	0.14	86,86,86,86	0
59	MG	DA	4201	1/1	0.66	0.14	57,57,57,57	0
59	MG	CA	1843	1/1	0.67	0.13	99,99,99,99	0
59	MG	CA	2084	1/1	0.67	0.12	77,77,77,77	0
59	MG	AA	3435	1/1	0.67	0.14	69,69,69,69	0
59	MG	DA	4839	1/1	0.67	0.28	75,75,75,75	0
59	MG	CA	1980	1/1	0.67	0.38	83,83,83,83	0
59	MG	DA	4031	1/1	0.67	0.33	79,79,79,79	0
59	MG	BA	1785	1/1	0.67	0.32	67,67,67,67	0
59	MG	D0	211	1/1	0.67	0.33	89,89,89,89	0
59	MG	CA	1632	1/1	0.67	0.36	65,65,65,65	0
59	MG	AA	3434	1/1	0.67	0.12	65,65,65,65	0
59	MG	AU	204	1/1	0.67	0.29	87,87,87,87	0
59	MG	DB	252	1/1	0.67	0.29	101,101,101,101	0
59	MG	BA	2040	1/1	0.67	0.34	93,93,93,93	0
59	MG	CA	1840	1/1	0.67	0.19	84,84,84,84	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	BG	307	1/1	0.67	0.17	85,85,85,85	0
59	MG	CA	1825	1/1	0.67	0.14	90,90,90,90	0
59	MG	DA	4448	1/1	0.67	0.23	95,95,95,95	0
59	MG	DA	4668	1/1	0.67	0.56	72,72,72,72	0
59	MG	DA	4911	1/1	0.67	0.24	85,85,85,85	0
59	MG	DA	5050	1/1	0.67	0.33	83,83,83,83	0
59	MG	BA	2172	1/1	0.67	0.32	75,75,75,75	0
59	MG	AA	4073	1/1	0.67	0.11	84,84,84,84	0
59	MG	DA	3832	1/1	0.67	0.34	55,55,55,55	0
59	MG	DA	4815	1/1	0.68	0.25	53,53,53,53	0
59	MG	BA	1832	1/1	0.68	0.21	89,89,89,89	0
59	MG	AA	3719	1/1	0.68	0.49	97,97,97,97	0
59	MG	CA	2105	1/1	0.68	0.43	94,94,94,94	0
59	MG	BA	2184	1/1	0.68	0.17	101,101,101,101	0
59	MG	AA	3491	1/1	0.68	0.19	68,68,68,68	0
59	MG	BA	2153	1/1	0.68	0.11	100,100,100,100	0
59	MG	AA	3433	1/1	0.68	0.24	64,64,64,64	0
59	MG	CA	2126	1/1	0.68	0.12	59,59,59,59	0
59	MG	BA	2070	1/1	0.68	0.28	89,89,89,89	0
59	MG	DA	4810	1/1	0.68	0.26	62,62,62,62	0
59	MG	AA	4106	1/1	0.68	0.28	88,88,88,88	0
59	MG	DA	4317	1/1	0.68	0.21	61,61,61,61	0
59	MG	AA	4060	1/1	0.68	0.13	70,70,70,70	0
59	MG	AS	203	1/1	0.68	0.46	72,72,72,72	0
59	MG	DA	3741	1/1	0.68	0.42	76,76,76,76	0
59	MG	BA	2250	1/1	0.68	0.21	90,90,90,90	0
59	MG	DA	3847	1/1	0.68	0.26	81,81,81,81	0
59	MG	CA	2230	1/1	0.68	0.14	92,92,92,92	0
59	MG	BA	1995	1/1	0.68	0.17	84,84,84,84	0
59	MG	AA	3124	1/1	0.68	0.35	71,71,71,71	0
59	MG	AA	3923	1/1	0.68	0.15	107,107,107,107	0
59	MG	DA	4128	1/1	0.68	0.34	85,85,85,85	0
59	MG	DA	4795	1/1	0.68	0.25	66,66,66,66	0
59	MG	DA	4929	1/1	0.68	0.43	52,52,52,52	0
59	MG	CA	1658	1/1	0.68	0.21	48,48,48,48	0
59	MG	DA	3258	1/1	0.68	0.17	83,83,83,83	0
59	MG	D0	209	1/1	0.68	0.27	91,91,91,91	0
59	MG	AA	3552	1/1	0.68	0.25	90,90,90,90	0
59	MG	CK	208	1/1	0.68	0.21	69,69,69,69	0
59	MG	AA	3325	1/1	0.69	0.22	50,50,50,50	0
59	MG	AA	3767	1/1	0.69	0.24	78,78,78,78	0
59	MG	BA	1913	1/1	0.69	0.16	73,73,73,73	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3441	1/1	0.69	0.17	102,102,102,102	0
59	MG	BA	2161	1/1	0.69	0.19	114,114,114,114	0
59	MG	BA	2018	1/1	0.69	0.24	84,84,84,84	0
59	MG	DA	4656	1/1	0.69	0.29	71,71,71,71	0
59	MG	DA	4302	1/1	0.69	0.10	102,102,102,102	0
59	MG	BA	1639	1/1	0.69	0.27	72,72,72,72	0
59	MG	AA	3122	1/1	0.69	0.18	73,73,73,73	0
59	MG	DB	244	1/1	0.69	0.34	68,68,68,68	0
59	MG	BA	1673	1/1	0.69	0.20	73,73,73,73	0
59	MG	CE	302	1/1	0.69	0.30	98,98,98,98	0
59	MG	DB	210	1/1	0.69	0.23	89,89,89,89	0
59	MG	DA	4431	1/1	0.69	0.20	87,87,87,87	0
59	MG	AA	3821	1/1	0.69	0.17	69,69,69,69	0
59	MG	DB	263	1/1	0.69	0.24	80,80,80,80	0
59	MG	BS	101	1/1	0.69	0.13	76,76,76,76	0
59	MG	BS	106	1/1	0.69	0.33	100,100,100,100	0
59	MG	BW	207	1/1	0.69	0.16	85,85,85,85	0
59	MG	DA	4573	1/1	0.69	0.34	102,102,102,102	0
59	MG	AA	3744	1/1	0.69	0.20	64,64,64,64	0
59	MG	DA	4753	1/1	0.69	0.31	68,68,68,68	0
59	MG	DA	4461	1/1	0.69	0.37	88,88,88,88	0
59	MG	DA	4948	1/1	0.69	0.51	126,126,126,126	0
59	MG	DA	4534	1/1	0.69	0.22	118,118,118,118	0
59	MG	D6	103	1/1	0.69	0.43	63,63,63,63	0
59	MG	DY	202	1/1	0.69	0.18	91,91,91,91	0
59	MG	AA	4051	1/1	0.69	0.29	106,106,106,106	0
59	MG	BA	2122	1/1	0.69	0.19	107,107,107,107	0
59	MG	BA	1722	1/1	0.69	0.29	54,54,54,54	0
59	MG	CA	2308	1/1	0.69	0.36	107,107,107,107	0
59	MG	DA	4496	1/1	0.69	0.37	81,81,81,81	0
59	MG	AA	3438	1/1	0.69	0.08	99,99,99,99	0
59	MG	BA	2265	1/1	0.69	0.33	111,111,111,111	0
59	MG	BA	2067	1/1	0.69	0.17	95,95,95,95	0
59	MG	BA	1698	1/1	0.69	0.58	104,104,104,104	0
59	MG	AB	212	1/1	0.69	0.11	83,83,83,83	0
59	MG	CA	2303	1/1	0.69	0.26	102,102,102,102	0
59	MG	CA	1754	1/1	0.69	0.35	80,80,80,80	0
59	MG	BL	202	1/1	0.69	0.14	64,64,64,64	0
59	MG	DA	3205	1/1	0.69	0.30	65,65,65,65	0
59	MG	DA	4928	1/1	0.69	0.17	78,78,78,78	0
59	MG	DA	3582	1/1	0.69	0.25	63,63,63,63	0
59	MG	AA	3338	1/1	0.70	0.16	72,72,72,72	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3621	1/1	0.70	0.19	51,51,51,51	0
59	MG	BA	2244	1/1	0.70	0.41	113,113,113,113	0
59	MG	AA	3948	1/1	0.70	0.16	72,72,72,72	0
59	MG	CA	2099	1/1	0.70	0.11	94,94,94,94	0
59	MG	DA	3403	1/1	0.70	0.37	89,89,89,89	0
59	MG	BF	302	1/1	0.70	0.24	105,105,105,105	0
59	MG	AA	3678	1/1	0.70	0.12	49,49,49,49	0
59	MG	CA	1920	1/1	0.70	0.32	85,85,85,85	0
59	MG	DA	3370	1/1	0.70	0.28	73,73,73,73	0
59	MG	DA	4485	1/1	0.70	0.07	138,138,138,138	0
59	MG	BA	1665	1/1	0.70	0.29	61,61,61,61	0
59	MG	CW	201	1/1	0.70	0.41	89,89,89,89	0
59	MG	BA	1620	1/1	0.70	0.25	50,50,50,50	0
59	MG	BA	1769	1/1	0.70	0.29	87,87,87,87	0
59	MG	BA	1859	1/1	0.70	0.51	146,146,146,146	0
59	MG	DA	3622	1/1	0.70	0.28	69,69,69,69	0
59	MG	DA	5005	1/1	0.70	0.32	75,75,75,75	0
59	MG	DB	214	1/1	0.70	0.27	76,76,76,76	0
59	MG	BG	306	1/1	0.70	0.22	108,108,108,108	0
59	MG	DA	5065	1/1	0.70	0.39	89,89,89,89	0
59	MG	DA	5011	1/1	0.70	0.31	76,76,76,76	0
59	MG	DA	4462	1/1	0.70	0.20	74,74,74,74	0
59	MG	DA	4417	1/1	0.70	0.30	64,64,64,64	0
59	MG	BA	1915	1/1	0.70	0.33	69,69,69,69	0
59	MG	DA	4885	1/1	0.70	0.37	92,92,92,92	0
59	MG	AA	4112	1/1	0.70	0.39	89,89,89,89	0
59	MG	BA	2169	1/1	0.70	0.18	98,98,98,98	0
59	MG	BA	2267	1/1	0.70	0.26	91,91,91,91	0
59	MG	DS	201	1/1	0.70	0.32	88,88,88,88	0
59	MG	DA	4916	1/1	0.71	0.16	88,88,88,88	0
59	MG	AA	3161	1/1	0.71	0.18	63,63,63,63	0
59	MG	DA	4838	1/1	0.71	0.34	67,67,67,67	0
59	MG	AA	3333	1/1	0.71	0.52	59,59,59,59	0
59	MG	CA	2033	1/1	0.71	0.14	73,73,73,73	0
59	MG	AA	3480	1/1	0.71	0.25	70,70,70,70	0
59	MG	DA	3120	1/1	0.71	0.23	57,57,57,57	0
59	MG	DB	226	1/1	0.71	0.20	40,40,40,40	0
59	MG	CA	2077	1/1	0.71	0.14	96,96,96,96	0
59	MG	DA	5046	1/1	0.71	0.20	77,77,77,77	0
59	MG	AA	3920	1/1	0.71	0.17	85,85,85,85	0
59	MG	AA	3623	1/1	0.71	0.23	50,50,50,50	0
59	MG	AA	3564	1/1	0.71	0.24	67,67,67,67	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CG	303	1/1	0.71	0.11	73,73,73,73	0
59	MG	DA	4971	1/1	0.71	0.27	79,79,79,79	0
59	MG	DB	274	1/1	0.71	0.21	79,79,79,79	0
59	MG	DA	3795	1/1	0.71	0.34	88,88,88,88	0
59	MG	BA	2183	1/1	0.71	0.20	107,107,107,107	0
59	MG	CD	101	1/1	0.71	0.20	85,85,85,85	0
59	MG	AA	3705	1/1	0.71	0.28	87,87,87,87	0
59	MG	AB	223	1/1	0.71	0.17	112,112,112,112	0
59	MG	DA	3326	1/1	0.71	0.20	63,63,63,63	0
59	MG	AA	3995	1/1	0.71	0.33	103,103,103,103	0
59	MG	BD	125	1/1	0.71	0.30	105,105,105,105	0
59	MG	AA	3346	1/1	0.71	0.27	58,58,58,58	0
59	MG	DA	3769	1/1	0.71	0.35	72,72,72,72	0
59	MG	BA	2139	1/1	0.71	0.11	82,82,82,82	0
59	MG	DA	3658	1/1	0.71	0.26	60,60,60,60	0
59	MG	DA	3314	1/1	0.71	0.13	83,83,83,83	0
59	MG	CG	302	1/1	0.71	0.14	67,67,67,67	0
59	MG	DA	3206	1/1	0.71	0.30	76,76,76,76	0
59	MG	BQ	102	1/1	0.71	0.10	63,63,63,63	0
59	MG	DA	4723	1/1	0.71	0.17	71,71,71,71	0
59	MG	AA	3633	1/1	0.71	0.21	41,41,41,41	0
59	MG	BA	2064	1/1	0.71	0.10	90,90,90,90	0
59	MG	CA	1841	1/1	0.71	0.11	85,85,85,85	0
59	MG	AK	203	1/1	0.71	0.16	86,86,86,86	0
59	MG	BA	1968	1/1	0.71	0.50	96,96,96,96	0
59	MG	AA	3116	1/1	0.71	0.44	85,85,85,85	0
59	MG	DM	202	1/1	0.71	0.43	48,48,48,48	0
59	MG	DA	4488	1/1	0.71	0.55	89,89,89,89	0
59	MG	AA	4054	1/1	0.71	0.13	74,74,74,74	0
59	MG	DA	3113	1/1	0.71	0.28	73,73,73,73	0
59	MG	AA	3419	1/1	0.71	0.38	82,82,82,82	0
59	MG	DA	3262	1/1	0.71	0.47	72,72,72,72	0
59	MG	CA	2097	1/1	0.72	0.16	82,82,82,82	0
59	MG	DA	4769	1/1	0.72	0.24	68,68,68,68	0
59	MG	BA	2205	1/1	0.72	0.17	109,109,109,109	0
59	MG	DA	4307	1/1	0.72	0.30	57,57,57,57	0
59	MG	CA	1893	1/1	0.72	0.12	54,54,54,54	0
59	MG	AA	3692	1/1	0.72	0.26	85,85,85,85	0
59	MG	DA	4883	1/1	0.72	0.34	96,96,96,96	0
59	MG	DA	3183	1/1	0.72	0.51	97,97,97,97	0
59	MG	DA	3274	1/1	0.72	0.41	67,67,67,67	0
59	MG	AA	3542	1/1	0.72	0.19	54,54,54,54	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3307	1/1	0.72	0.20	59,59,59,59	0
59	MG	AA	4007	1/1	0.72	0.30	94,94,94,94	0
59	MG	CA	2010	1/1	0.72	0.55	104,104,104,104	0
59	MG	CA	1962	1/1	0.72	0.19	76,76,76,76	0
59	MG	AZ	101	1/1	0.72	0.23	83,83,83,83	0
59	MG	AA	3593	1/1	0.72	0.20	80,80,80,80	0
59	MG	DA	3983	1/1	0.72	0.22	57,57,57,57	0
59	MG	CB	105	1/1	0.72	0.31	76,76,76,76	0
59	MG	DA	3854	1/1	0.72	0.15	72,72,72,72	0
59	MG	BM	201	1/1	0.72	0.23	64,64,64,64	0
59	MG	CA	1828	1/1	0.72	0.35	100,100,100,100	0
59	MG	AA	3675	1/1	0.72	0.15	71,71,71,71	0
59	MG	DU	215	1/1	0.72	0.44	76,76,76,76	0
59	MG	BA	2234	1/1	0.72	0.25	108,108,108,108	0
59	MG	AA	3990	1/1	0.72	0.56	134,134,134,134	0
59	MG	BA	1635	1/1	0.72	0.35	102,102,102,102	0
59	MG	AA	4059	1/1	0.72	0.17	54,54,54,54	0
59	MG	BA	1816	1/1	0.72	0.20	71,71,71,71	0
59	MG	AA	4155	1/1	0.72	0.23	80,80,80,80	0
59	MG	AA	4153	1/1	0.72	0.29	109,109,109,109	0
59	MG	DA	4230	1/1	0.72	0.31	41,41,41,41	0
59	MG	DA	4773	1/1	0.72	0.31	81,81,81,81	0
59	MG	CA	1896	1/1	0.72	0.16	73,73,73,73	0
59	MG	BA	1757	1/1	0.72	0.27	39,39,39,39	0
59	MG	D1	206	1/1	0.72	0.28	68,68,68,68	0
59	MG	CR	101	1/1	0.72	0.19	89,89,89,89	0
59	MG	BA	1867	1/1	0.72	0.32	121,121,121,121	0
59	MG	AA	4131	1/1	0.72	0.18	99,99,99,99	0
59	MG	CA	2214	1/1	0.72	0.16	104,104,104,104	0
59	MG	DA	4784	1/1	0.72	0.16	74,74,74,74	0
59	MG	CA	2217	1/1	0.72	0.32	93,93,93,93	0
59	MG	AA	3833	1/1	0.72	0.19	80,80,80,80	0
59	MG	AA	3915	1/1	0.73	0.32	105,105,105,105	0
59	MG	DA	4872	1/1	0.73	0.58	106,106,106,106	0
59	MG	BA	1789	1/1	0.73	0.33	105,105,105,105	0
59	MG	DA	4277	1/1	0.73	0.34	66,66,66,66	0
59	MG	BA	2262	1/1	0.73	0.14	90,90,90,90	0
59	MG	DA	4904	1/1	0.73	0.31	71,71,71,71	0
59	MG	DA	4149	1/1	0.73	0.30	70,70,70,70	0
59	MG	BA	2264	1/1	0.73	0.14	79,79,79,79	0
59	MG	BA	1629	1/1	0.73	0.27	76,76,76,76	0
59	MG	DA	4053	1/1	0.73	0.23	75,75,75,75	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3265	1/1	0.73	0.40	71,71,71,71	0
59	MG	CD	123	1/1	0.73	0.25	120,120,120,120	0
59	MG	CA	2307	1/1	0.73	0.15	76,76,76,76	0
59	MG	CA	2224	1/1	0.73	0.17	83,83,83,83	0
59	MG	AA	4116	1/1	0.73	0.26	98,98,98,98	0
59	MG	BX	101	1/1	0.73	0.26	74,74,74,74	0
59	MG	BS	103	1/1	0.73	0.20	83,83,83,83	0
59	MG	DA	4366	1/1	0.73	0.22	62,62,62,62	0
59	MG	DA	4892	1/1	0.73	1.12	133,133,133,133	0
59	MG	DA	4648	1/1	0.73	0.27	61,61,61,61	0
59	MG	CA	2123	1/1	0.73	0.30	100,100,100,100	0
59	MG	AA	3883	1/1	0.73	0.32	100,100,100,100	0
59	MG	CA	1937	1/1	0.73	0.15	73,73,73,73	0
59	MG	AA	4000	1/1	0.73	0.24	91,91,91,91	0
59	MG	CA	1892	1/1	0.73	0.24	68,68,68,68	0
59	MG	CK	209	1/1	0.73	0.20	60,60,60,60	0
59	MG	BA	1612	1/1	0.73	0.49	90,90,90,90	0
59	MG	DA	5032	1/1	0.73	0.27	82,82,82,82	0
59	MG	DA	4526	1/1	0.73	0.27	53,53,53,53	0
59	MG	CA	2291	1/1	0.73	0.16	80,80,80,80	0
59	MG	CD	128	1/1	0.73	0.20	66,66,66,66	0
59	MG	BA	2003	1/1	0.73	0.17	98,98,98,98	0
59	MG	DA	4055	1/1	0.73	0.20	68,68,68,68	0
59	MG	DA	3577	1/1	0.73	0.33	101,101,101,101	0
59	MG	CA	1858	1/1	0.73	0.14	73,73,73,73	0
59	MG	AA	4151	1/1	0.73	0.13	108,108,108,108	0
59	MG	BW	205	1/1	0.73	0.18	91,91,91,91	0
59	MG	DA	4980	1/1	0.73	0.48	79,79,79,79	0
59	MG	DA	4357	1/1	0.73	0.29	53,53,53,53	0
59	MG	AA	3224	1/1	0.73	0.18	63,63,63,63	0
59	MG	CA	1890	1/1	0.73	0.09	69,69,69,69	0
59	MG	AB	219	1/1	0.73	0.15	75,75,75,75	0
59	MG	AR	201	1/1	0.73	0.11	71,71,71,71	0
59	MG	DA	4364	1/1	0.73	0.12	90,90,90,90	0
59	MG	DA	4474	1/1	0.73	0.62	54,54,54,54	0
59	MG	AA	3342	1/1	0.73	0.15	63,63,63,63	0
59	MG	AA	4036	1/1	0.73	0.10	68,68,68,68	0
59	MG	A8	102	1/1	0.74	0.20	68,68,68,68	0
59	MG	BA	2206	1/1	0.74	0.16	106,106,106,106	0
59	MG	CA	1949	1/1	0.74	0.14	60,60,60,60	0
59	MG	BA	2220	1/1	0.74	0.38	86,86,86,86	0
59	MG	DA	3377	1/1	0.74	0.33	83,83,83,83	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	4087	1/1	0.74	0.22	89,89,89,89	0
59	MG	CA	1855	1/1	0.74	0.10	79,79,79,79	0
59	MG	DA	3278	1/1	0.74	0.31	62,62,62,62	0
59	MG	CA	2117	1/1	0.74	0.34	84,84,84,84	0
59	MG	AA	3825	1/1	0.74	0.29	81,81,81,81	0
59	MG	BA	1770	1/1	0.74	0.20	67,67,67,67	0
59	MG	BD	122	1/1	0.74	0.38	100,100,100,100	0
59	MG	AA	3397	1/1	0.74	0.17	56,56,56,56	0
59	MG	BA	1936	1/1	0.74	0.25	80,80,80,80	0
59	MG	DA	4391	1/1	0.74	0.19	52,52,52,52	0
59	MG	DA	4413	1/1	0.74	0.23	65,65,65,65	0
59	MG	AA	3382	1/1	0.74	0.21	83,83,83,83	0
59	MG	BA	2276	1/1	0.74	0.41	128,128,128,128	0
59	MG	DA	4381	1/1	0.74	0.25	88,88,88,88	0
59	MG	DB	211	1/1	0.74	0.41	90,90,90,90	0
59	MG	CD	108	1/1	0.74	0.42	97,97,97,97	0
59	MG	CA	1634	1/1	0.74	0.24	91,91,91,91	0
59	MG	AO	206	1/1	0.74	0.57	53,53,53,53	0
59	MG	DA	4520	1/1	0.74	0.23	84,84,84,84	0
59	MG	DA	3859	1/1	0.74	0.31	92,92,92,92	0
59	MG	CA	1714	1/1	0.74	0.31	66,66,66,66	0
59	MG	AB	218	1/1	0.74	0.18	81,81,81,81	0
59	MG	AA	3966	1/1	0.74	0.40	75,75,75,75	0
59	MG	CA	2063	1/1	0.74	0.17	78,78,78,78	0
59	MG	CA	2312	1/1	0.74	0.25	78,78,78,78	0
59	MG	CA	2166	1/1	0.74	0.31	91,91,91,91	0
59	MG	DA	4879	1/1	0.74	0.53	62,62,62,62	0
59	MG	DB	219	1/1	0.74	0.31	70,70,70,70	0
59	MG	AA	4079	1/1	0.74	0.21	92,92,92,92	0
59	MG	BA	2182	1/1	0.74	0.17	110,110,110,110	0
59	MG	DA	4782	1/1	0.74	0.23	83,83,83,83	0
59	MG	AA	4014	1/1	0.74	0.17	84,84,84,84	0
59	MG	BA	2110	1/1	0.74	0.31	91,91,91,91	0
59	MG	BA	2022	1/1	0.74	0.10	104,104,104,104	0
59	MG	DA	4574	1/1	0.74	0.31	73,73,73,73	0
59	MG	DA	3813	1/1	0.74	0.14	96,96,96,96	0
59	MG	BA	2164	1/1	0.74	0.15	78,78,78,78	0
59	MG	BG	302	1/1	0.74	0.13	63,63,63,63	0
59	MG	AA	3946	1/1	0.74	0.07	98,98,98,98	0
59	MG	AA	3817	1/1	0.74	0.17	66,66,66,66	0
59	MG	BD	107	1/1	0.74	0.13	71,71,71,71	0
59	MG	DA	4536	1/1	0.74	0.31	64,64,64,64	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4545	1/1	0.74	0.73	90,90,90,90	0
59	MG	DA	4512	1/1	0.74	0.67	100,100,100,100	0
59	MG	AA	3563	1/1	0.74	0.23	67,67,67,67	0
59	MG	AA	3964	1/1	0.74	0.37	111,111,111,111	0
59	MG	BA	2081	1/1	0.74	0.28	69,69,69,69	0
59	MG	AA	3490	1/1	0.75	0.15	102,102,102,102	0
59	MG	DA	3487	1/1	0.75	0.21	62,62,62,62	0
59	MG	DA	3674	1/1	0.75	0.16	62,62,62,62	0
59	MG	AA	3984	1/1	0.75	0.18	98,98,98,98	0
59	MG	AA	4111	1/1	0.75	0.15	108,108,108,108	0
59	MG	DA	4154	1/1	0.75	0.36	74,74,74,74	0
59	MG	AA	3710	1/1	0.75	0.21	76,76,76,76	0
59	MG	BA	1659	1/1	0.75	0.11	71,71,71,71	0
59	MG	BC	112	1/1	0.75	0.16	57,57,57,57	0
59	MG	AA	3707	1/1	0.75	0.12	63,63,63,63	0
59	MG	CC	113	1/1	0.75	0.38	85,85,85,85	0
59	MG	CD	110	1/1	0.75	0.10	77,77,77,77	0
59	MG	BA	1926	1/1	0.75	0.11	53,53,53,53	0
59	MG	AA	3060	1/1	0.75	0.12	76,76,76,76	0
59	MG	BA	2077	1/1	0.75	0.16	74,74,74,74	0
59	MG	BA	1873	1/1	0.75	0.19	109,109,109,109	0
59	MG	CA	2137	1/1	0.75	0.19	88,88,88,88	0
59	MG	CA	1854	1/1	0.75	0.18	67,67,67,67	0
59	MG	DA	5066	1/1	0.75	0.34	61,61,61,61	0
59	MG	AA	3756	1/1	0.75	0.20	65,65,65,65	0
59	MG	BA	2014	1/1	0.75	0.14	64,64,64,64	0
59	MG	AA	3718	1/1	0.75	0.32	84,84,84,84	0
59	MG	DA	4814	1/1	0.75	0.36	114,114,114,114	0
59	MG	AA	3154	1/1	0.75	0.19	59,59,59,59	0
59	MG	DA	5052	1/1	0.75	0.23	49,49,49,49	0
59	MG	CA	2192	1/1	0.75	0.13	60,60,60,60	0
59	MG	AA	3037	1/1	0.75	0.36	107,107,107,107	0
59	MG	DA	4915	1/1	0.75	0.40	64,64,64,64	0
59	MG	DA	4297	1/1	0.75	0.38	100,100,100,100	0
59	MG	CA	2030	1/1	0.75	0.22	72,72,72,72	0
59	MG	CA	2129	1/1	0.75	0.19	75,75,75,75	0
59	MG	BA	1815	1/1	0.75	0.22	119,119,119,119	0
59	MG	DA	4190	1/1	0.75	0.08	70,70,70,70	0
59	MG	DV	302	1/1	0.75	0.15	87,87,87,87	0
59	MG	DA	3906	1/1	0.75	0.16	90,90,90,90	0
59	MG	DA	4793	1/1	0.75	0.27	86,86,86,86	0
59	MG	BA	2119	1/1	0.75	0.16	67,67,67,67	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3658	1/1	0.75	0.23	64,64,64,64	0
59	MG	AA	3698	1/1	0.75	0.18	64,64,64,64	0
59	MG	DA	4483	1/1	0.75	0.23	64,64,64,64	0
59	MG	CA	1834	1/1	0.75	0.09	67,67,67,67	0
59	MG	AA	3897	1/1	0.75	0.20	103,103,103,103	0
59	MG	CA	1894	1/1	0.75	0.23	60,60,60,60	0
59	MG	DA	3562	1/1	0.75	0.40	73,73,73,73	0
59	MG	BD	105	1/1	0.75	0.31	87,87,87,87	0
59	MG	BK	204	1/1	0.75	0.24	74,74,74,74	0
59	MG	CB	112	1/1	0.75	0.13	77,77,77,77	0
59	MG	CA	1929	1/1	0.75	0.19	40,40,40,40	0
59	MG	CA	2213	1/1	0.76	0.11	84,84,84,84	0
59	MG	AA	4015	1/1	0.76	0.39	104,104,104,104	0
59	MG	AA	4140	1/1	0.76	0.19	74,74,74,74	0
59	MG	DY	201	1/1	0.76	0.23	100,100,100,100	0
59	MG	AA	4003	1/1	0.76	0.07	109,109,109,109	0
59	MG	DA	4494	1/1	0.76	0.34	84,84,84,84	0
59	MG	BA	1774	1/1	0.76	0.19	53,53,53,53	0
59	MG	AA	3274	1/1	0.76	0.32	70,70,70,70	0
59	MG	AA	3440	1/1	0.76	0.19	75,75,75,75	0
59	MG	CA	2317	1/1	0.76	0.20	104,104,104,104	0
59	MG	DA	4454	1/1	0.76	0.54	51,51,51,51	0
59	MG	AA	3725	1/1	0.76	0.21	89,89,89,89	0
59	MG	CG	310	1/1	0.76	0.13	96,96,96,96	0
59	MG	DA	3919	1/1	0.76	0.32	76,76,76,76	0
59	MG	DA	3392	1/1	0.76	0.26	76,76,76,76	0
59	MG	CA	2184	1/1	0.76	0.38	83,83,83,83	0
59	MG	CA	2234	1/1	0.76	0.21	81,81,81,81	0
59	MG	AA	4097	1/1	0.76	0.47	109,109,109,109	0
59	MG	DA	3108	1/1	0.76	0.27	76,76,76,76	0
59	MG	DA	4742	1/1	0.76	0.17	128,128,128,128	0
59	MG	CA	1792	1/1	0.76	0.07	58,58,58,58	0
59	MG	AA	4049	1/1	0.76	0.28	115,115,115,115	0
59	MG	DA	4121	1/1	0.76	0.15	77,77,77,77	0
59	MG	DA	3162	1/1	0.76	0.24	45,45,45,45	0
59	MG	CA	1767	1/1	0.76	0.18	80,80,80,80	0
59	MG	DA	3526	1/1	0.76	0.16	46,46,46,46	0
59	MG	DA	4184	1/1	0.76	0.33	61,61,61,61	0
59	MG	AA	3245	1/1	0.76	0.28	69,69,69,69	0
59	MG	AA	3187	1/1	0.76	0.25	68,68,68,68	0
59	MG	AA	4040	1/1	0.76	0.23	67,67,67,67	0
59	MG	BA	1678	1/1	0.76	0.33	77,77,77,77	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	1709	1/1	0.76	0.17	55,55,55,55	0
59	MG	AA	3649	1/1	0.76	0.20	96,96,96,96	0
59	MG	AA	3071	1/1	0.76	0.31	16,16,16,16	0
59	MG	AA	3428	1/1	0.76	0.42	101,101,101,101	0
59	MG	AA	3728	1/1	0.76	0.19	70,70,70,70	0
59	MG	BA	1830	1/1	0.76	0.30	90,90,90,90	0
59	MG	DA	4275	1/1	0.76	0.35	78,78,78,78	0
59	MG	AA	3292	1/1	0.76	0.34	79,79,79,79	0
59	MG	AA	3726	1/1	0.76	0.16	60,60,60,60	0
59	MG	CA	2180	1/1	0.76	0.18	66,66,66,66	0
59	MG	AA	3757	1/1	0.76	0.21	56,56,56,56	0
59	MG	DA	3843	1/1	0.76	0.22	53,53,53,53	0
59	MG	BA	1851	1/1	0.76	0.29	142,142,142,142	0
59	MG	BA	2104	1/1	0.76	0.46	80,80,80,80	0
59	MG	DA	4831	1/1	0.76	0.24	82,82,82,82	0
59	MG	DA	3894	1/1	0.76	0.17	101,101,101,101	0
59	MG	BB	108	1/1	0.76	0.25	94,94,94,94	0
59	MG	DA	3219	1/1	0.76	0.22	48,48,48,48	0
59	MG	DA	4985	1/1	0.76	0.40	88,88,88,88	0
59	MG	BA	2068	1/1	0.76	0.47	90,90,90,90	0
59	MG	DA	4956	1/1	0.76	0.43	118,118,118,118	0
59	MG	DA	4739	1/1	0.76	0.37	77,77,77,77	0
59	MG	BA	2176	1/1	0.76	0.15	99,99,99,99	0
59	MG	CA	1956	1/1	0.76	0.14	74,74,74,74	0
59	MG	CC	115	1/1	0.76	0.34	69,69,69,69	0
59	MG	AA	3870	1/1	0.76	0.11	69,69,69,69	0
59	MG	DA	4620	1/1	0.76	0.24	84,84,84,84	0
59	MG	DA	3695	1/1	0.76	0.23	131,131,131,131	0
59	MG	DA	4587	1/1	0.76	0.21	71,71,71,71	0
59	MG	DA	3912	1/1	0.76	0.23	89,89,89,89	0
59	MG	BB	109	1/1	0.77	0.18	119,119,119,119	0
59	MG	DA	3654	1/1	0.77	0.17	40,40,40,40	0
59	MG	CA	1968	1/1	0.77	0.23	90,90,90,90	0
59	MG	DA	3332	1/1	0.77	0.42	82,82,82,82	0
59	MG	DB	240	1/1	0.77	0.26	84,84,84,84	0
59	MG	DB	273	1/1	0.77	0.24	100,100,100,100	0
59	MG	AA	3284	1/1	0.77	0.14	78,78,78,78	0
59	MG	DA	3130	1/1	0.77	0.26	73,73,73,73	0
59	MG	DA	4690	1/1	0.77	0.28	86,86,86,86	0
59	MG	DA	3889	1/1	0.77	0.14	80,80,80,80	0
59	MG	BA	1920	1/1	0.77	0.26	86,86,86,86	0
59	MG	DA	4650	1/1	0.77	0.30	83,83,83,83	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4757	1/1	0.77	0.17	46,46,46,46	0
59	MG	BA	1803	1/1	0.77	0.38	90,90,90,90	0
59	MG	DA	3613	1/1	0.77	0.31	68,68,68,68	0
59	MG	DA	4626	1/1	0.77	0.17	56,56,56,56	0
59	MG	AA	3743	1/1	0.77	0.43	71,71,71,71	0
59	MG	BH	205	1/1	0.77	0.10	78,78,78,78	0
59	MG	DA	4575	1/1	0.77	0.29	68,68,68,68	0
59	MG	CJ	201	1/1	0.77	0.26	89,89,89,89	0
59	MG	BD	123	1/1	0.77	0.20	82,82,82,82	0
59	MG	AB	207	1/1	0.77	0.16	74,74,74,74	0
59	MG	DA	3365	1/1	0.77	0.34	45,45,45,45	0
59	MG	DA	3887	1/1	0.77	0.36	68,68,68,68	0
59	MG	BA	2163	1/1	0.77	0.14	105,105,105,105	0
59	MG	DA	3676	1/1	0.77	0.51	70,70,70,70	0
59	MG	DA	3481	1/1	0.77	0.32	93,93,93,93	0
59	MG	DA	3626	1/1	0.77	0.19	84,84,84,84	0
59	MG	DA	3679	1/1	0.77	0.07	69,69,69,69	0
59	MG	CA	2191	1/1	0.77	0.15	99,99,99,99	0
59	MG	CA	2278	1/1	0.77	0.25	84,84,84,84	0
59	MG	CA	2109	1/1	0.77	0.11	74,74,74,74	0
59	MG	AA	3353	1/1	0.77	0.33	83,83,83,83	0
59	MG	DU	207	1/1	0.77	0.20	97,97,97,97	0
59	MG	DA	3509	1/1	0.77	0.39	47,47,47,47	0
59	MG	AB	227	1/1	0.77	0.28	82,82,82,82	0
59	MG	DS	206	1/1	0.77	0.24	60,60,60,60	0
59	MG	CA	1624	1/1	0.77	0.15	80,80,80,80	0
59	MG	DA	4015	1/1	0.77	0.40	57,57,57,57	0
59	MG	DB	231	1/1	0.77	0.49	83,83,83,83	0
59	MG	AA	4012	1/1	0.77	0.24	85,85,85,85	0
59	MG	DA	3013	1/1	0.77	0.22	50,50,50,50	0
59	MG	DA	3883	1/1	0.77	0.15	74,74,74,74	0
59	MG	CA	2282	1/1	0.77	0.20	128,128,128,128	0
59	MG	CA	2246	1/1	0.77	0.18	91,91,91,91	0
59	MG	D7	104	1/1	0.77	0.25	47,47,47,47	0
59	MG	DA	4638	1/1	0.77	0.26	56,56,56,56	0
59	MG	AA	3605	1/1	0.77	0.12	66,66,66,66	0
59	MG	DA	4442	1/1	0.77	0.19	49,49,49,49	0
59	MG	CA	2302	1/1	0.77	0.13	77,77,77,77	0
59	MG	CC	109	1/1	0.77	0.19	49,49,49,49	0
59	MG	DA	4771	1/1	0.77	0.60	92,92,92,92	0
59	MG	CA	2193	1/1	0.77	0.34	86,86,86,86	0
59	MG	DA	3344	1/1	0.77	0.62	90,90,90,90	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4084	1/1	0.77	0.27	79,79,79,79	0
59	MG	AA	3747	1/1	0.77	0.16	53,53,53,53	0
59	MG	AA	3732	1/1	0.77	0.12	76,76,76,76	0
59	MG	DA	5013	1/1	0.77	0.16	54,54,54,54	0
59	MG	AA	4023	1/1	0.77	0.12	73,73,73,73	0
59	MG	BA	1927	1/1	0.77	0.20	51,51,51,51	0
59	MG	DA	3790	1/1	0.77	0.27	66,66,66,66	0
59	MG	DA	3992	1/1	0.77	0.25	61,61,61,61	0
59	MG	DB	233	1/1	0.77	0.15	45,45,45,45	0
59	MG	DA	3633	1/1	0.77	0.29	48,48,48,48	0
59	MG	CK	205	1/1	0.77	0.16	75,75,75,75	0
59	MG	DA	3631	1/1	0.77	0.21	60,60,60,60	0
59	MG	A2	201	1/1	0.77	0.31	66,66,66,66	0
59	MG	AA	3417	1/1	0.77	0.11	58,58,58,58	0
59	MG	AA	3613	1/1	0.77	0.14	84,84,84,84	0
59	MG	AB	214	1/1	0.77	0.31	105,105,105,105	0
59	MG	BD	113	1/1	0.77	0.13	78,78,78,78	0
59	MG	AB	210	1/1	0.77	0.22	81,81,81,81	0
59	MG	DE	308	1/1	0.77	0.94	83,83,83,83	0
59	MG	CA	1699	1/1	0.77	0.12	40,40,40,40	0
59	MG	DA	3665	1/1	0.77	0.19	110,110,110,110	0
59	MG	BA	1937	1/1	0.77	0.38	71,71,71,71	0
59	MG	DA	4539	1/1	0.77	0.34	85,85,85,85	0
59	MG	AA	3287	1/1	0.77	0.14	66,66,66,66	0
59	MG	BD	126	1/1	0.77	0.41	133,133,133,133	0
59	MG	DA	4731	1/1	0.77	0.19	63,63,63,63	0
59	MG	CA	2042	1/1	0.77	0.15	57,57,57,57	0
59	MG	DA	4481	1/1	0.78	0.24	62,62,62,62	0
59	MG	DA	4279	1/1	0.78	0.12	72,72,72,72	0
59	MG	AA	3943	1/1	0.78	0.20	65,65,65,65	0
59	MG	CA	2173	1/1	0.78	0.15	71,71,71,71	0
59	MG	BA	2140	1/1	0.78	0.30	103,103,103,103	0
59	MG	CA	2313	1/1	0.78	0.34	128,128,128,128	0
59	MG	AA	3837	1/1	0.78	0.28	116,116,116,116	0
59	MG	DA	4582	1/1	0.78	0.25	69,69,69,69	0
59	MG	DA	4752	1/1	0.78	0.20	84,84,84,84	0
59	MG	AA	3422	1/1	0.78	0.14	73,73,73,73	0
59	MG	CD	109	1/1	0.78	0.12	70,70,70,70	0
59	MG	AA	3812	1/1	0.78	0.21	83,83,83,83	0
59	MG	DA	4552	1/1	0.78	0.27	83,83,83,83	0
59	MG	DA	3592	1/1	0.78	0.21	81,81,81,81	0
59	MG	AA	4083	1/1	0.78	0.16	74,74,74,74	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	1935	1/1	0.78	0.17	45,45,45,45	0
59	MG	DA	4865	1/1	0.78	0.26	67,67,67,67	0
59	MG	BA	1839	1/1	0.78	0.29	136,136,136,136	0
59	MG	DA	3408	1/1	0.78	0.14	41,41,41,41	0
59	MG	AA	4141	1/1	0.78	0.11	88,88,88,88	0
59	MG	DA	3385	1/1	0.78	0.27	85,85,85,85	0
59	MG	BA	2028	1/1	0.78	0.26	77,77,77,77	0
59	MG	AA	3396	1/1	0.78	0.17	58,58,58,58	0
59	MG	CA	1818	1/1	0.78	0.21	101,101,101,101	0
59	MG	DA	4421	1/1	0.78	0.25	64,64,64,64	0
59	MG	DA	3494	1/1	0.78	0.27	59,59,59,59	0
59	MG	CA	2208	1/1	0.78	0.15	81,81,81,81	0
59	MG	AA	4089	1/1	0.78	0.16	107,107,107,107	0
59	MG	BA	1834	1/1	0.78	0.14	70,70,70,70	0
59	MG	DA	3682	1/1	0.78	0.16	83,83,83,83	0
59	MG	DF	309	1/1	0.78	0.24	46,46,46,46	0
59	MG	DA	4834	1/1	0.78	0.26	78,78,78,78	0
59	MG	CA	1772	1/1	0.78	0.21	62,62,62,62	0
59	MG	DE	312	1/1	0.78	0.16	74,74,74,74	0
59	MG	BA	1969	1/1	0.78	0.11	71,71,71,71	0
59	MG	AA	3905	1/1	0.78	0.10	86,86,86,86	0
59	MG	DA	4717	1/1	0.78	0.33	80,80,80,80	0
59	MG	DY	204	1/1	0.78	0.14	85,85,85,85	0
59	MG	DA	4316	1/1	0.78	0.24	67,67,67,67	0
59	MG	DB	216	1/1	0.78	0.18	43,43,43,43	0
59	MG	BA	1912	1/1	0.78	0.30	82,82,82,82	0
59	MG	DA	3195	1/1	0.78	0.28	50,50,50,50	0
59	MG	CA	2047	1/1	0.78	0.12	80,80,80,80	0
59	MG	DA	3160	1/1	0.78	0.31	48,48,48,48	0
59	MG	DA	4540	1/1	0.78	0.24	88,88,88,88	0
59	MG	AA	3363	1/1	0.78	0.27	74,74,74,74	0
59	MG	D0	207	1/1	0.78	0.33	58,58,58,58	0
59	MG	AA	3775	1/1	0.78	0.48	91,91,91,91	0
59	MG	DA	4660	1/1	0.78	0.24	73,73,73,73	0
59	MG	DA	5076	1/1	0.78	0.33	85,85,85,85	0
59	MG	BA	2102	1/1	0.78	0.12	85,85,85,85	0
59	MG	DA	3766	1/1	0.78	0.24	77,77,77,77	0
59	MG	D1	201	1/1	0.78	0.24	49,49,49,49	0
59	MG	AA	3330	1/1	0.78	0.20	111,111,111,111	0
59	MG	DA	5053	1/1	0.78	0.51	101,101,101,101	0
59	MG	DA	4037	1/1	0.78	0.22	81,81,81,81	0
59	MG	DA	3624	1/1	0.78	0.27	74,74,74,74	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	4068	1/1	0.78	0.16	58,58,58,58	0
59	MG	DA	3257	1/1	0.78	0.25	66,66,66,66	0
59	MG	DA	4473	1/1	0.78	0.34	60,60,60,60	0
59	MG	BD	108	1/1	0.78	0.13	82,82,82,82	0
59	MG	DA	4744	1/1	0.78	0.30	70,70,70,70	0
59	MG	CA	2251	1/1	0.78	0.16	93,93,93,93	0
59	MG	DA	3873	1/1	0.78	0.46	119,119,119,119	0
59	MG	BM	203	1/1	0.78	0.24	91,91,91,91	0
59	MG	BA	1923	1/1	0.78	0.35	81,81,81,81	0
59	MG	BA	1814	1/1	0.78	0.12	115,115,115,115	0
59	MG	DA	3586	1/1	0.78	0.39	83,83,83,83	0
59	MG	AA	3077	1/1	0.78	0.14	63,63,63,63	0
59	MG	AU	202	1/1	0.78	0.23	76,76,76,76	0
59	MG	DB	245	1/1	0.78	0.15	59,59,59,59	0
59	MG	DA	3122	1/1	0.78	0.32	75,75,75,75	0
59	MG	AA	3321	1/1	0.78	0.17	69,69,69,69	0
59	MG	BA	1792	1/1	0.78	0.32	79,79,79,79	0
59	MG	DA	4185	1/1	0.78	0.44	88,88,88,88	0
59	MG	DW	105	1/1	0.78	0.21	81,81,81,81	0
59	MG	DA	4100	1/1	0.78	0.23	72,72,72,72	0
59	MG	BD	114	1/1	0.78	0.14	93,93,93,93	0
59	MG	CA	1997	1/1	0.78	0.46	96,96,96,96	0
59	MG	CB	101	1/1	0.78	0.33	64,64,64,64	0
59	MG	DA	4998	1/1	0.78	0.36	110,110,110,110	0
59	MG	AA	3739	1/1	0.79	0.22	86,86,86,86	0
59	MG	DO	211	1/1	0.79	0.30	53,53,53,53	0
59	MG	DA	3425	1/1	0.79	0.30	63,63,63,63	0
59	MG	CA	2023	1/1	0.79	0.36	79,79,79,79	0
59	MG	BA	2257	1/1	0.79	0.10	85,85,85,85	0
59	MG	AA	3344	1/1	0.79	0.16	76,76,76,76	0
59	MG	AG	203	1/1	0.79	0.16	97,97,97,97	0
59	MG	AA	3548	1/1	0.79	0.29	59,59,59,59	0
59	MG	BW	202	1/1	0.79	0.11	73,73,73,73	0
59	MG	BA	2188	1/1	0.79	0.37	102,102,102,102	0
59	MG	AA	4121	1/1	0.79	0.14	66,66,66,66	0
59	MG	CA	2270	1/1	0.79	0.16	68,68,68,68	0
59	MG	DA	4625	1/1	0.79	0.29	67,67,67,67	0
59	MG	BD	124	1/1	0.79	0.11	104,104,104,104	0
59	MG	BA	2191	1/1	0.79	0.13	73,73,73,73	0
59	MG	AA	3561	1/1	0.79	0.15	45,45,45,45	0
59	MG	CS	104	1/1	0.79	0.17	125,125,125,125	0
59	MG	DA	4806	1/1	0.79	0.19	59,59,59,59	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3418	1/1	0.79	0.48	91,91,91,91	0
59	MG	CA	2178	1/1	0.79	0.13	92,92,92,92	0
59	MG	BA	1994	1/1	0.79	0.15	107,107,107,107	0
59	MG	BA	1741	1/1	0.79	0.21	85,85,85,85	0
59	MG	DA	3576	1/1	0.79	0.43	102,102,102,102	0
59	MG	AA	3985	1/1	0.79	0.23	97,97,97,97	0
59	MG	DA	4993	1/1	0.79	0.21	139,139,139,139	0
59	MG	AA	4093	1/1	0.79	0.56	114,114,114,114	0
59	MG	DA	3549	1/1	0.79	0.17	40,40,40,40	0
59	MG	DA	4917	1/1	0.79	0.33	61,61,61,61	0
59	MG	AA	3927	1/1	0.79	0.17	76,76,76,76	0
59	MG	DA	3556	1/1	0.79	0.35	107,107,107,107	0
59	MG	BA	2111	1/1	0.79	0.28	79,79,79,79	0
59	MG	DA	4244	1/1	0.79	0.13	44,44,44,44	0
59	MG	DA	4874	1/1	0.79	0.45	84,84,84,84	0
59	MG	CA	1942	1/1	0.79	0.51	105,105,105,105	0
59	MG	D3	107	1/1	0.79	0.16	53,53,53,53	0
59	MG	CA	1761	1/1	0.79	0.27	65,65,65,65	0
59	MG	BA	1890	1/1	0.79	0.36	72,72,72,72	0
59	MG	BA	1928	1/1	0.79	0.14	67,67,67,67	0
59	MG	AA	3973	1/1	0.79	0.16	86,86,86,86	0
59	MG	AB	220	1/1	0.79	0.17	79,79,79,79	0
59	MG	DA	3908	1/1	0.79	0.20	124,124,124,124	0
59	MG	DA	4518	1/1	0.79	0.24	49,49,49,49	0
59	MG	BG	301	1/1	0.79	0.08	67,67,67,67	0
59	MG	DA	4265	1/1	0.79	0.31	95,95,95,95	0
59	MG	BK	201	1/1	0.79	0.34	96,96,96,96	0
59	MG	DA	4542	1/1	0.79	0.23	37,37,37,37	0
59	MG	DA	3596	1/1	0.79	0.19	76,76,76,76	0
59	MG	BA	1802	1/1	0.79	0.29	75,75,75,75	0
59	MG	DA	4823	1/1	0.79	0.24	71,71,71,71	0
59	MG	DA	4827	1/1	0.79	0.41	87,87,87,87	0
59	MG	DA	4418	1/1	0.79	0.30	96,96,96,96	0
59	MG	BA	2219	1/1	0.79	0.17	60,60,60,60	0
59	MG	DA	4924	1/1	0.79	0.18	75,75,75,75	0
59	MG	AA	3981	1/1	0.79	0.27	131,131,131,131	0
59	MG	BA	2269	1/1	0.79	0.19	80,80,80,80	0
59	MG	BK	205	1/1	0.79	0.36	86,86,86,86	0
59	MG	A7	102	1/1	0.79	0.41	59,59,59,59	0
59	MG	BA	1712	1/1	0.79	0.35	56,56,56,56	0
59	MG	DA	5033	1/1	0.79	0.11	75,75,75,75	0
59	MG	DA	5056	1/1	0.79	0.61	98,98,98,98	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	1899	1/1	0.79	0.19	49,49,49,49	0
59	MG	CA	1646	1/1	0.79	0.13	53,53,53,53	0
59	MG	DA	4940	1/1	0.79	0.32	69,69,69,69	0
59	MG	AA	3665	1/1	0.79	0.40	80,80,80,80	0
59	MG	AA	4149	1/1	0.79	0.23	125,125,125,125	0
59	MG	DA	4817	1/1	0.79	0.26	78,78,78,78	0
59	MG	AA	3416	1/1	0.79	0.35	78,78,78,78	0
59	MG	BA	1606	1/1	0.79	0.15	93,93,93,93	0
59	MG	AA	3049	1/1	0.79	0.21	48,48,48,48	0
59	MG	AZ	102	1/1	0.79	0.22	50,50,50,50	0
59	MG	DA	3872	1/1	0.79	0.15	75,75,75,75	0
59	MG	DA	3870	1/1	0.79	0.33	67,67,67,67	0
59	MG	AA	3864	1/1	0.79	0.11	48,48,48,48	0
59	MG	CB	103	1/1	0.79	0.15	53,53,53,53	0
59	MG	AG	202	1/1	0.79	0.10	83,83,83,83	0
59	MG	AA	3799	1/1	0.79	0.38	93,93,93,93	0
59	MG	DA	4886	1/1	0.79	0.41	60,60,60,60	0
59	MG	AA	3371	1/1	0.79	0.08	63,63,63,63	0
59	MG	AA	3298	1/1	0.79	0.13	38,38,38,38	0
59	MG	DA	3745	1/1	0.79	0.31	49,49,49,49	0
59	MG	BA	2130	1/1	0.79	0.13	86,86,86,86	0
59	MG	AA	4024	1/1	0.80	0.30	95,95,95,95	0
59	MG	CA	2273	1/1	0.80	0.14	65,65,65,65	0
59	MG	AA	3631	1/1	0.80	0.13	58,58,58,58	0
59	MG	A3	104	1/1	0.80	0.13	83,83,83,83	0
59	MG	CA	1835	1/1	0.80	0.27	114,114,114,114	0
59	MG	CA	2220	1/1	0.80	0.32	112,112,112,112	0
59	MG	BA	2224	1/1	0.80	0.20	83,83,83,83	0
59	MG	BA	1793	1/1	0.80	0.11	57,57,57,57	0
59	MG	DA	3856	1/1	0.80	0.26	70,70,70,70	0
59	MG	DA	3423	1/1	0.80	0.12	58,58,58,58	0
59	MG	DA	4688	1/1	0.80	0.41	83,83,83,83	0
59	MG	AA	4160	1/1	0.80	0.26	74,74,74,74	0
59	MG	BA	1605	1/1	0.80	0.10	56,56,56,56	0
59	MG	BA	1771	1/1	0.80	0.41	74,74,74,74	0
59	MG	AB	224	1/1	0.80	0.15	85,85,85,85	0
59	MG	BA	2144	1/1	0.80	0.18	75,75,75,75	0
59	MG	CA	2320	1/1	0.80	0.11	81,81,81,81	0
59	MG	DA	4504	1/1	0.80	0.36	82,82,82,82	0
59	MG	DA	4997	1/1	0.80	0.18	50,50,50,50	0
59	MG	BA	1796	1/1	0.80	0.42	77,77,77,77	0
59	MG	AA	3527	1/1	0.80	0.16	34,34,34,34	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	2035	1/1	0.80	0.10	68,68,68,68	0
59	MG	DA	4943	1/1	0.80	0.41	81,81,81,81	0
59	MG	CA	1954	1/1	0.80	0.49	90,90,90,90	0
59	MG	DA	4612	1/1	0.80	0.46	79,79,79,79	0
59	MG	BA	2209	1/1	0.80	0.11	56,56,56,56	0
59	MG	BA	1854	1/1	0.80	0.39	84,84,84,84	0
59	MG	BA	1621	1/1	0.80	0.60	120,120,120,120	0
59	MG	AA	3165	1/1	0.80	0.22	53,53,53,53	0
59	MG	DA	4549	1/1	0.80	0.13	98,98,98,98	0
59	MG	D1	202	1/1	0.80	0.24	57,57,57,57	0
59	MG	BS	105	1/1	0.80	0.11	89,89,89,89	0
59	MG	BA	1869	1/1	0.80	0.12	73,73,73,73	0
59	MG	DA	4040	1/1	0.80	0.28	59,59,59,59	0
59	MG	DA	4976	1/1	0.80	0.68	98,98,98,98	0
59	MG	CF	301	1/1	0.80	0.24	75,75,75,75	0
59	MG	DA	3473	1/1	0.80	0.19	44,44,44,44	0
59	MG	BA	1679	1/1	0.80	0.17	50,50,50,50	0
59	MG	DA	3824	1/1	0.80	0.37	73,73,73,73	0
59	MG	DA	3775	1/1	0.80	0.18	87,87,87,87	0
59	MG	AA	3526	1/1	0.80	0.14	75,75,75,75	0
59	MG	DA	3770	1/1	0.80	0.41	88,88,88,88	0
59	MG	CA	2050	1/1	0.80	0.28	80,80,80,80	0
59	MG	CB	106	1/1	0.80	0.35	97,97,97,97	0
59	MG	DA	4778	1/1	0.80	0.23	66,66,66,66	0
59	MG	BQ	103	1/1	0.80	0.17	69,69,69,69	0
59	MG	DB	228	1/1	0.80	0.65	91,91,91,91	0
59	MG	BA	1984	1/1	0.80	0.09	84,84,84,84	0
59	MG	AA	3413	1/1	0.80	0.49	95,95,95,95	0
59	MG	AA	3986	1/1	0.80	0.13	88,88,88,88	0
59	MG	DA	4404	1/1	0.80	0.18	68,68,68,68	0
59	MG	BA	1987	1/1	0.80	0.18	83,83,83,83	0
59	MG	AA	3144	1/1	0.80	0.21	69,69,69,69	0
59	MG	AA	4094	1/1	0.80	0.34	57,57,57,57	0
59	MG	DA	4863	1/1	0.80	0.22	83,83,83,83	0
59	MG	AA	3880	1/1	0.80	0.14	102,102,102,102	0
59	MG	AA	4020	1/1	0.80	0.27	85,85,85,85	0
59	MG	AA	3212	1/1	0.80	0.36	78,78,78,78	0
59	MG	BA	2157	1/1	0.80	0.13	72,72,72,72	0
59	MG	CH	202	1/1	0.80	0.12	62,62,62,62	0
59	MG	AK	202	1/1	0.80	0.11	75,75,75,75	0
59	MG	DA	3913	1/1	0.80	0.32	120,120,120,120	0
59	MG	CA	2164	1/1	0.80	0.18	103,103,103,103	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	BA	2216	1/1	0.80	0.36	91,91,91,91	0
59	MG	AA	3608	1/1	0.80	0.17	86,86,86,86	0
59	MG	CA	2210	1/1	0.80	0.11	144,144,144,144	0
59	MG	AA	3253	1/1	0.80	0.21	50,50,50,50	0
59	MG	AA	3645	1/1	0.80	0.26	46,46,46,46	0
59	MG	CA	2124	1/1	0.80	0.41	94,94,94,94	0
59	MG	DA	5012	1/1	0.80	0.25	100,100,100,100	0
59	MG	CV	103	1/1	0.80	0.14	84,84,84,84	0
59	MG	DA	4937	1/1	0.80	0.28	78,78,78,78	0
59	MG	AA	3873	1/1	0.80	0.24	87,87,87,87	0
59	MG	DG	203	1/1	0.80	0.21	64,64,64,64	0
59	MG	DA	4331	1/1	0.80	0.27	63,63,63,63	0
59	MG	A0	201	1/1	0.80	0.17	61,61,61,61	0
59	MG	DA	3525	1/1	0.80	0.18	43,43,43,43	0
59	MG	DA	4062	1/1	0.80	0.18	49,49,49,49	0
59	MG	DA	3844	1/1	0.80	0.20	109,109,109,109	0
59	MG	AA	3934	1/1	0.80	0.20	93,93,93,93	0
59	MG	AA	4126	1/1	0.80	0.31	87,87,87,87	0
59	MG	DO	218	1/1	0.80	0.19	74,74,74,74	0
59	MG	DA	4986	1/1	0.80	0.20	63,63,63,63	0
59	MG	DA	3657	1/1	0.80	0.15	74,74,74,74	0
59	MG	DA	4151	1/1	0.80	0.25	66,66,66,66	0
59	MG	CD	103	1/1	0.80	0.41	75,75,75,75	0
59	MG	DA	3511	1/1	0.80	0.30	42,42,42,42	0
59	MG	DA	4910	1/1	0.80	0.13	67,67,67,67	0
59	MG	DA	5037	1/1	0.80	0.15	84,84,84,84	0
59	MG	BA	2274	1/1	0.80	0.17	82,82,82,82	0
59	MG	AA	3497	1/1	0.80	0.26	80,80,80,80	0
59	MG	DA	4269	1/1	0.80	0.53	72,72,72,72	0
59	MG	CA	2108	1/1	0.80	0.14	78,78,78,78	0
59	MG	CA	2194	1/1	0.80	0.20	101,101,101,101	0
59	MG	BA	1930	1/1	0.80	0.24	62,62,62,62	0
59	MG	AA	3853	1/1	0.80	0.15	55,55,55,55	0
59	MG	DH	205	1/1	0.80	0.16	77,77,77,77	0
59	MG	DA	4852	1/1	0.80	0.28	142,142,142,142	0
59	MG	CA	1759	1/1	0.81	0.28	75,75,75,75	0
59	MG	BD	117	1/1	0.81	0.27	115,115,115,115	0
59	MG	DA	3968	1/1	0.81	0.30	55,55,55,55	0
59	MG	AA	3806	1/1	0.81	0.11	79,79,79,79	0
59	MG	CA	2255	1/1	0.81	0.29	95,95,95,95	0
59	MG	BA	1967	1/1	0.81	0.36	77,77,77,77	0
59	MG	CA	2186	1/1	0.81	0.18	121,121,121,121	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4263	1/1	0.81	0.18	66,66,66,66	0
59	MG	DA	4495	1/1	0.81	0.48	111,111,111,111	0
59	MG	AA	3444	1/1	0.81	0.32	81,81,81,81	0
59	MG	DA	4177	1/1	0.81	0.29	52,52,52,52	0
59	MG	BA	1755	1/1	0.81	0.15	56,56,56,56	0
59	MG	CA	1928	1/1	0.81	0.16	78,78,78,78	0
59	MG	AA	3360	1/1	0.81	0.19	95,95,95,95	0
59	MG	AA	4100	1/1	0.81	0.07	103,103,103,103	0
59	MG	DA	3371	1/1	0.81	0.25	73,73,73,73	0
59	MG	DA	3498	1/1	0.81	0.23	44,44,44,44	0
59	MG	BA	2154	1/1	0.81	0.18	73,73,73,73	0
59	MG	DA	3727	1/1	0.81	0.31	49,49,49,49	0
59	MG	DA	4658	1/1	0.81	0.26	76,76,76,76	0
59	MG	DO	216	1/1	0.81	0.35	58,58,58,58	0
59	MG	AA	3359	1/1	0.81	0.10	48,48,48,48	0
59	MG	DA	3412	1/1	0.81	0.29	68,68,68,68	0
59	MG	B1	101	1/1	0.81	0.25	60,60,60,60	0
59	MG	DA	4665	1/1	0.81	0.20	76,76,76,76	0
59	MG	DA	4781	1/1	0.81	0.46	105,105,105,105	0
59	MG	CA	2027	1/1	0.81	0.44	110,110,110,110	0
59	MG	DA	4851	1/1	0.81	0.20	139,139,139,139	0
59	MG	BA	2066	1/1	0.81	0.39	105,105,105,105	0
59	MG	DA	4934	1/1	0.81	0.30	61,61,61,61	0
59	MG	AA	3114	1/1	0.81	0.33	97,97,97,97	0
59	MG	AA	3882	1/1	0.81	0.18	77,77,77,77	0
59	MG	DA	3268	1/1	0.81	0.27	50,50,50,50	0
59	MG	DA	4802	1/1	0.81	0.33	75,75,75,75	0
59	MG	D1	208	1/1	0.81	0.19	58,58,58,58	0
59	MG	DA	4123	1/1	0.81	0.18	75,75,75,75	0
59	MG	CA	1916	1/1	0.81	0.11	61,61,61,61	0
59	MG	CA	2311	1/1	0.81	0.30	81,81,81,81	0
59	MG	AA	3805	1/1	0.81	0.33	94,94,94,94	0
59	MG	DE	311	1/1	0.81	0.24	42,42,42,42	0
59	MG	BA	1766	1/1	0.81	0.11	74,74,74,74	0
59	MG	CA	2272	1/1	0.81	0.09	147,147,147,147	0
59	MG	CA	2222	1/1	0.81	0.14	72,72,72,72	0
59	MG	BA	1762	1/1	0.81	0.29	78,78,78,78	0
59	MG	AA	3452	1/1	0.81	0.19	97,97,97,97	0
59	MG	CC	123	1/1	0.81	0.17	74,74,74,74	0
59	MG	DA	5027	1/1	0.81	0.41	85,85,85,85	0
59	MG	BI	201	1/1	0.81	0.09	47,47,47,47	0
59	MG	BA	2083	1/1	0.81	0.17	86,86,86,86	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3895	1/1	0.81	0.09	51,51,51,51	0
59	MG	CA	1966	1/1	0.81	0.19	69,69,69,69	0
59	MG	DA	4713	1/1	0.81	0.32	75,75,75,75	0
59	MG	BA	1932	1/1	0.81	0.12	107,107,107,107	0
59	MG	DA	4516	1/1	0.81	0.25	53,53,53,53	0
59	MG	CA	1963	1/1	0.81	0.11	71,71,71,71	0
59	MG	DA	4360	1/1	0.81	0.12	57,57,57,57	0
59	MG	DA	4223	1/1	0.81	0.20	60,60,60,60	0
59	MG	CA	2200	1/1	0.81	0.12	73,73,73,73	0
59	MG	DB	246	1/1	0.81	0.17	52,52,52,52	0
59	MG	AA	3928	1/1	0.81	0.14	71,71,71,71	0
59	MG	BA	1729	1/1	0.81	0.38	69,69,69,69	0
59	MG	AA	3482	1/1	0.81	0.22	58,58,58,58	0
59	MG	AA	3545	1/1	0.81	0.19	56,56,56,56	0
59	MG	DA	4215	1/1	0.81	0.23	53,53,53,53	0
59	MG	DA	4705	1/1	0.81	0.58	93,93,93,93	0
59	MG	CA	1747	1/1	0.81	0.14	51,51,51,51	0
59	MG	AA	3341	1/1	0.81	0.16	57,57,57,57	0
59	MG	AA	3997	1/1	0.81	0.16	78,78,78,78	0
59	MG	DA	3396	1/1	0.81	0.33	46,46,46,46	0
59	MG	CA	2120	1/1	0.81	0.30	84,84,84,84	0
59	MG	CA	1755	1/1	0.81	0.22	75,75,75,75	0
59	MG	BA	2238	1/1	0.81	0.14	95,95,95,95	0
59	MG	BA	1865	1/1	0.81	0.12	75,75,75,75	0
59	MG	CA	2165	1/1	0.81	0.15	101,101,101,101	0
59	MG	DA	5014	1/1	0.81	0.29	64,64,64,64	0
59	MG	DB	267	1/1	0.81	0.17	73,73,73,73	0
59	MG	CC	106	1/1	0.81	0.24	60,60,60,60	0
59	MG	DA	4006	1/1	0.81	0.42	71,71,71,71	0
59	MG	CA	2233	1/1	0.81	0.23	60,60,60,60	0
59	MG	BA	2093	1/1	0.81	0.36	106,106,106,106	0
59	MG	BA	2132	1/1	0.81	0.13	87,87,87,87	0
59	MG	CD	106	1/1	0.81	0.27	88,88,88,88	0
59	MG	DM	206	1/1	0.81	0.39	85,85,85,85	0
59	MG	AA	3841	1/1	0.81	0.25	85,85,85,85	0
59	MG	CA	2259	1/1	0.81	0.20	67,67,67,67	0
59	MG	DA	3842	1/1	0.81	0.10	54,54,54,54	0
59	MG	AU	206	1/1	0.81	0.25	75,75,75,75	0
59	MG	DA	4122	1/1	0.81	0.55	105,105,105,105	0
59	MG	BA	1949	1/1	0.81	0.32	85,85,85,85	0
59	MG	CA	2175	1/1	0.81	0.21	89,89,89,89	0
59	MG	AA	3855	1/1	0.81	0.10	72,72,72,72	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3404	1/1	0.81	0.30	65,65,65,65	0
59	MG	BA	1642	1/1	0.81	0.17	45,45,45,45	0
59	MG	DA	4605	1/1	0.81	0.38	70,70,70,70	0
59	MG	DA	4562	1/1	0.81	0.23	46,46,46,46	0
59	MG	AA	3914	1/1	0.81	0.21	78,78,78,78	0
59	MG	AA	3496	1/1	0.81	0.10	63,63,63,63	0
59	MG	DU	217	1/1	0.81	0.26	110,110,110,110	0
59	MG	AA	3474	1/1	0.81	0.16	65,65,65,65	0
59	MG	DA	4208	1/1	0.81	0.35	74,74,74,74	0
59	MG	DA	4833	1/1	0.81	0.33	76,76,76,76	0
59	MG	DA	3345	1/1	0.81	0.42	80,80,80,80	0
59	MG	CA	2139	1/1	0.81	0.09	96,96,96,96	0
59	MG	AA	3902	1/1	0.81	0.11	111,111,111,111	0
59	MG	DA	3406	1/1	0.81	0.31	59,59,59,59	0
59	MG	DA	4727	1/1	0.81	0.15	69,69,69,69	0
59	MG	DT	106	1/1	0.81	0.49	69,69,69,69	0
59	MG	DA	4890	1/1	0.81	0.39	96,96,96,96	0
59	MG	AA	4056	1/1	0.81	0.10	74,74,74,74	0
59	MG	BW	208	1/1	0.81	0.19	62,62,62,62	0
59	MG	CF	302	1/1	0.81	0.22	79,79,79,79	0
59	MG	CA	2041	1/1	0.81	0.16	64,64,64,64	0
59	MG	CA	2094	1/1	0.81	0.25	87,87,87,87	0
59	MG	DA	4134	1/1	0.81	0.28	34,34,34,34	0
59	MG	DA	3879	1/1	0.81	0.14	83,83,83,83	0
59	MG	DB	207	1/1	0.81	0.12	64,64,64,64	0
59	MG	DA	4577	1/1	0.81	0.32	57,57,57,57	0
59	MG	DA	3356	1/1	0.81	0.30	58,58,58,58	0
59	MG	CA	2149	1/1	0.81	0.39	103,103,103,103	0
59	MG	CD	111	1/1	0.81	0.40	69,69,69,69	0
59	MG	CA	1827	1/1	0.81	0.25	101,101,101,101	0
59	MG	AA	3296	1/1	0.81	0.23	55,55,55,55	0
59	MG	AA	3485	1/1	0.81	0.37	91,91,91,91	0
59	MG	CM	203	1/1	0.81	0.67	89,89,89,89	0
59	MG	DA	3452	1/1	0.82	0.29	68,68,68,68	0
59	MG	BA	1939	1/1	0.82	0.34	81,81,81,81	0
59	MG	BA	1731	1/1	0.82	0.29	46,46,46,46	0
59	MG	A5	103	1/1	0.82	0.12	54,54,54,54	0
59	MG	DA	3747	1/1	0.82	0.12	75,75,75,75	0
59	MG	CA	1864	1/1	0.82	0.28	82,82,82,82	0
59	MG	CA	2160	1/1	0.82	0.20	96,96,96,96	0
59	MG	BA	2186	1/1	0.82	0.16	79,79,79,79	0
59	MG	CA	2150	1/1	0.82	0.17	85,85,85,85	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	BA	2217	1/1	0.82	0.14	108,108,108,108	0
59	MG	DA	3180	1/1	0.82	0.12	77,77,77,77	0
59	MG	DA	3247	1/1	0.82	0.23	72,72,72,72	0
59	MG	DA	4388	1/1	0.82	0.27	80,80,80,80	0
59	MG	BA	2162	1/1	0.82	0.12	115,115,115,115	0
59	MG	CA	2049	1/1	0.82	0.39	113,113,113,113	0
59	MG	AA	3519	1/1	0.82	0.13	63,63,63,63	0
59	MG	BA	1720	1/1	0.82	0.49	95,95,95,95	0
59	MG	DA	4896	1/1	0.82	0.19	74,74,74,74	0
59	MG	BA	1647	1/1	0.82	0.21	78,78,78,78	0
59	MG	AA	3724	1/1	0.82	0.23	65,65,65,65	0
59	MG	BA	2148	1/1	0.82	0.14	110,110,110,110	0
59	MG	BB	112	1/1	0.82	0.23	107,107,107,107	0
59	MG	DA	4376	1/1	0.82	0.20	60,60,60,60	0
59	MG	DA	4570	1/1	0.82	0.23	59,59,59,59	0
59	MG	BA	1652	1/1	0.82	0.32	40,40,40,40	0
59	MG	DA	4765	1/1	0.82	0.09	120,120,120,120	0
59	MG	DS	202	1/1	0.82	0.24	71,71,71,71	0
59	MG	AF	301	1/1	0.82	0.11	73,73,73,73	0
59	MG	CA	2142	1/1	0.82	0.51	110,110,110,110	0
59	MG	DA	3857	1/1	0.82	0.48	78,78,78,78	0
59	MG	DB	271	1/1	0.82	0.68	72,72,72,72	0
59	MG	DA	4691	1/1	0.82	0.14	114,114,114,114	0
59	MG	BA	1822	1/1	0.82	0.09	97,97,97,97	0
59	MG	CA	1981	1/1	0.82	0.20	78,78,78,78	0
59	MG	DA	3190	1/1	0.82	0.27	60,60,60,60	0
59	MG	DA	3836	1/1	0.82	0.45	85,85,85,85	0
59	MG	AA	3089	1/1	0.82	0.37	70,70,70,70	0
59	MG	AA	3276	1/1	0.82	0.18	46,46,46,46	0
59	MG	BA	2021	1/1	0.82	0.12	84,84,84,84	0
59	MG	BA	1971	1/1	0.82	0.34	98,98,98,98	0
59	MG	AA	4110	1/1	0.82	0.09	102,102,102,102	0
59	MG	DA	4630	1/1	0.82	0.35	89,89,89,89	0
59	MG	DA	3309	1/1	0.82	0.28	56,56,56,56	0
59	MG	DA	3245	1/1	0.82	0.24	41,41,41,41	0
59	MG	DA	3645	1/1	0.82	0.15	61,61,61,61	0
59	MG	DA	3306	1/1	0.82	0.17	61,61,61,61	0
59	MG	BA	2136	1/1	0.82	0.11	117,117,117,117	0
59	MG	BA	1638	1/1	0.82	0.20	51,51,51,51	0
59	MG	DA	3267	1/1	0.82	0.30	72,72,72,72	0
59	MG	DA	3619	1/1	0.82	0.17	59,59,59,59	0
59	MG	AA	3822	1/1	0.82	0.34	77,77,77,77	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3804	1/1	0.82	0.15	94,94,94,94	0
59	MG	BA	2069	1/1	0.82	0.49	105,105,105,105	0
59	MG	AA	3714	1/1	0.82	0.20	70,70,70,70	0
59	MG	AA	3285	1/1	0.82	0.12	62,62,62,62	0
59	MG	CA	1912	1/1	0.82	0.41	81,81,81,81	0
59	MG	CG	307	1/1	0.82	0.14	106,106,106,106	0
59	MG	DA	4447	1/1	0.82	0.23	63,63,63,63	0
59	MG	CI	201	1/1	0.82	0.14	56,56,56,56	0
59	MG	DA	3348	1/1	0.82	0.17	54,54,54,54	0
59	MG	CA	1763	1/1	0.82	0.17	39,39,39,39	0
59	MG	DA	3191	1/1	0.82	0.20	60,60,60,60	0
59	MG	AA	3566	1/1	0.82	0.19	65,65,65,65	0
59	MG	BA	2120	1/1	0.82	0.10	67,67,67,67	0
59	MG	DA	3512	1/1	0.82	0.28	66,66,66,66	0
59	MG	DA	3121	1/1	0.82	0.30	51,51,51,51	0
59	MG	DX	102	1/1	0.82	0.14	62,62,62,62	0
59	MG	DA	4859	1/1	0.82	0.23	91,91,91,91	0
59	MG	CA	2209	1/1	0.82	0.17	68,68,68,68	0
59	MG	CA	2075	1/1	0.82	0.21	84,84,84,84	0
59	MG	AA	3820	1/1	0.82	0.37	59,59,59,59	0
59	MG	DF	318	1/1	0.82	0.25	69,69,69,69	0
59	MG	CA	2216	1/1	0.82	0.16	89,89,89,89	0
59	MG	CA	2080	1/1	0.82	0.28	83,83,83,83	0
59	MG	DL	201	1/1	0.82	0.10	94,94,94,94	0
59	MG	DA	3168	1/1	0.82	0.17	40,40,40,40	0
59	MG	DA	3660	1/1	0.82	0.19	46,46,46,46	0
59	MG	AA	3243	1/1	0.82	0.27	99,99,99,99	0
59	MG	DA	3831	1/1	0.82	0.12	61,61,61,61	0
59	MG	DA	3277	1/1	0.82	0.43	67,67,67,67	0
59	MG	DA	3891	1/1	0.82	0.13	82,82,82,82	0
59	MG	AA	3606	1/1	0.82	0.20	97,97,97,97	0
59	MG	AT	101	1/1	0.82	0.55	70,70,70,70	0
59	MG	AA	3993	1/1	0.82	0.15	91,91,91,91	0
59	MG	DA	4614	1/1	0.82	0.38	78,78,78,78	0
59	MG	BA	2118	1/1	0.82	0.26	83,83,83,83	0
59	MG	DA	3260	1/1	0.82	0.27	60,60,60,60	0
59	MG	DA	4367	1/1	0.82	0.20	65,65,65,65	0
59	MG	DA	4440	1/1	0.82	0.35	54,54,54,54	0
59	MG	BA	1656	1/1	0.82	0.18	57,57,57,57	0
59	MG	AA	3387	1/1	0.82	0.16	40,40,40,40	0
59	MG	CA	2181	1/1	0.82	0.17	73,73,73,73	0
59	MG	BA	1805	1/1	0.82	0.42	99,99,99,99	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	D2	210	1/1	0.82	0.56	98,98,98,98	0
59	MG	BA	2141	1/1	0.82	0.26	80,80,80,80	0
59	MG	DA	3748	1/1	0.82	0.26	66,66,66,66	0
59	MG	BD	103	1/1	0.82	0.17	58,58,58,58	0
59	MG	CB	113	1/1	0.82	0.18	72,72,72,72	0
59	MG	DA	4761	1/1	0.82	0.30	78,78,78,78	0
59	MG	DA	3979	1/1	0.82	0.31	82,82,82,82	0
59	MG	DA	4990	1/1	0.82	0.27	70,70,70,70	0
59	MG	DE	306	1/1	0.82	0.11	45,45,45,45	0
59	MG	AA	3701	1/1	0.82	0.13	55,55,55,55	0
59	MG	DA	4486	1/1	0.82	0.14	63,63,63,63	0
59	MG	DA	3528	1/1	0.82	0.14	45,45,45,45	0
59	MG	BW	201	1/1	0.82	0.17	88,88,88,88	0
59	MG	BA	1829	1/1	0.82	0.17	75,75,75,75	0
59	MG	CA	2297	1/1	0.82	0.17	71,71,71,71	0
59	MG	BA	2270	1/1	0.82	0.30	100,100,100,100	0
59	MG	CR	103	1/1	0.82	0.20	70,70,70,70	0
59	MG	B1	102	1/1	0.82	0.10	72,72,72,72	0
59	MG	CA	2266	1/1	0.83	0.23	73,73,73,73	0
59	MG	CA	1850	1/1	0.83	0.20	100,100,100,100	0
59	MG	AA	4016	1/1	0.83	0.37	111,111,111,111	0
59	MG	CA	1794	1/1	0.83	0.19	124,124,124,124	0
59	MG	BA	2009	1/1	0.83	0.13	63,63,63,63	0
59	MG	CA	2221	1/1	0.83	0.13	72,72,72,72	0
59	MG	AA	3309	1/1	0.83	0.14	54,54,54,54	0
59	MG	AA	3238	1/1	0.83	0.18	63,63,63,63	0
59	MG	CA	1677	1/1	0.83	0.26	89,89,89,89	0
59	MG	AA	4005	1/1	0.83	0.12	63,63,63,63	0
59	MG	DA	3739	1/1	0.83	0.17	51,51,51,51	0
59	MG	DA	4021	1/1	0.83	0.33	76,76,76,76	0
59	MG	DA	3606	1/1	0.83	0.39	88,88,88,88	0
59	MG	AA	3910	1/1	0.83	0.12	76,76,76,76	0
59	MG	DA	4978	1/1	0.83	0.28	57,57,57,57	0
59	MG	DU	218	1/1	0.83	0.49	73,73,73,73	0
59	MG	DZ	104	1/1	0.83	0.50	59,59,59,59	0
59	MG	AA	4124	1/1	0.83	0.18	107,107,107,107	0
59	MG	AA	3517	1/1	0.83	0.27	69,69,69,69	0
59	MG	DA	4060	1/1	0.83	0.09	52,52,52,52	0
59	MG	AA	4103	1/1	0.83	0.25	54,54,54,54	0
59	MG	BK	202	1/1	0.83	0.30	80,80,80,80	0
59	MG	DA	3266	1/1	0.83	0.34	57,57,57,57	0
59	MG	DA	3931	1/1	0.83	0.16	83,83,83,83	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	2169	1/1	0.83	0.36	95,95,95,95	0
59	MG	AA	3951	1/1	0.83	0.09	98,98,98,98	0
59	MG	BA	1777	1/1	0.83	0.23	57,57,57,57	0
59	MG	AA	3879	1/1	0.83	0.12	91,91,91,91	0
59	MG	CA	2258	1/1	0.83	0.18	75,75,75,75	0
59	MG	CA	1638	1/1	0.83	0.23	51,51,51,51	0
59	MG	CA	2260	1/1	0.83	0.13	67,67,67,67	0
59	MG	AA	3670	1/1	0.83	0.14	54,54,54,54	0
59	MG	BA	2001	1/1	0.83	0.34	96,96,96,96	0
59	MG	DA	4310	1/1	0.83	0.19	56,56,56,56	0
59	MG	A1	201	1/1	0.83	0.17	60,60,60,60	0
59	MG	DA	4276	1/1	0.83	1.07	75,75,75,75	0
59	MG	DA	4551	1/1	0.83	0.32	118,118,118,118	0
59	MG	DA	4180	1/1	0.83	0.23	66,66,66,66	0
59	MG	DA	4451	1/1	0.83	0.32	63,63,63,63	0
59	MG	DT	101	1/1	0.83	0.16	54,54,54,54	0
59	MG	DP	204	1/1	0.83	0.23	51,51,51,51	0
59	MG	DA	3574	1/1	0.83	0.08	71,71,71,71	0
59	MG	DA	3484	1/1	0.83	0.34	76,76,76,76	0
59	MG	DD	313	1/1	0.83	0.15	64,64,64,64	0
59	MG	AA	3796	1/1	0.83	0.19	94,94,94,94	0
59	MG	BA	1688	1/1	0.83	0.32	61,61,61,61	0
59	MG	DA	4240	1/1	0.83	0.43	70,70,70,70	0
59	MG	AB	211	1/1	0.83	0.35	77,77,77,77	0
59	MG	CA	1979	1/1	0.83	0.13	58,58,58,58	0
59	MG	BA	2151	1/1	0.83	0.20	81,81,81,81	0
59	MG	DA	4502	1/1	0.83	0.35	62,62,62,62	0
59	MG	CA	1787	1/1	0.83	0.08	59,59,59,59	0
59	MG	BA	2030	1/1	0.83	0.33	71,71,71,71	0
59	MG	DA	4284	1/1	0.83	0.22	52,52,52,52	0
59	MG	CD	112	1/1	0.83	0.13	142,142,142,142	0
59	MG	AA	4119	1/1	0.83	0.36	112,112,112,112	0
59	MG	DA	3182	1/1	0.83	0.34	65,65,65,65	0
59	MG	DA	4533	1/1	0.83	0.53	88,88,88,88	0
59	MG	DA	3641	1/1	0.83	0.23	60,60,60,60	0
59	MG	DA	3447	1/1	0.83	0.22	48,48,48,48	0
59	MG	DA	4328	1/1	0.83	0.32	59,59,59,59	0
59	MG	DA	4200	1/1	0.83	0.18	61,61,61,61	0
59	MG	DA	3407	1/1	0.83	0.33	70,70,70,70	0
59	MG	DA	4009	1/1	0.83	0.17	98,98,98,98	0
59	MG	DA	4595	1/1	0.83	0.34	84,84,84,84	0
59	MG	AA	3315	1/1	0.83	0.16	49,49,49,49	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4617	1/1	0.83	0.20	54,54,54,54	0
59	MG	AA	4082	1/1	0.83	0.17	75,75,75,75	0
59	MG	AA	3477	1/1	0.83	0.28	65,65,65,65	0
59	MG	AA	4011	1/1	0.83	0.30	122,122,122,122	0
59	MG	BA	2160	1/1	0.83	0.15	95,95,95,95	0
59	MG	DA	4186	1/1	0.83	0.24	45,45,45,45	0
59	MG	AA	3765	1/1	0.83	0.30	74,74,74,74	0
59	MG	BA	2189	1/1	0.83	0.11	140,140,140,140	0
59	MG	DA	4478	1/1	0.83	0.23	65,65,65,65	0
59	MG	AA	4064	1/1	0.83	0.11	76,76,76,76	0
59	MG	DA	3820	1/1	0.83	0.11	97,97,97,97	0
59	MG	DA	4800	1/1	0.83	0.16	49,49,49,49	0
59	MG	DA	4295	1/1	0.83	0.17	48,48,48,48	0
59	MG	CA	1951	1/1	0.83	0.08	82,82,82,82	0
59	MG	BA	2142	1/1	0.83	0.15	84,84,84,84	0
59	MG	AA	3286	1/1	0.83	0.18	67,67,67,67	0
59	MG	AA	3335	1/1	0.83	0.18	83,83,83,83	0
59	MG	BA	2208	1/1	0.83	0.35	105,105,105,105	0
59	MG	BB	110	1/1	0.83	0.19	69,69,69,69	0
59	MG	AA	3638	1/1	0.83	0.32	51,51,51,51	0
59	MG	CA	2155	1/1	0.83	0.19	65,65,65,65	0
59	MG	BW	203	1/1	0.83	0.29	125,125,125,125	0
59	MG	AA	3148	1/1	0.83	0.13	53,53,53,53	0
59	MG	DA	4629	1/1	0.83	0.26	107,107,107,107	0
59	MG	DA	4329	1/1	0.83	0.21	63,63,63,63	0
59	MG	AA	4109	1/1	0.83	0.13	81,81,81,81	0
59	MG	AA	3936	1/1	0.83	0.24	83,83,83,83	0
59	MG	DA	3605	1/1	0.83	0.14	29,29,29,29	0
59	MG	AB	221	1/1	0.83	0.17	61,61,61,61	0
59	MG	AA	4035	1/1	0.83	0.20	75,75,75,75	0
59	MG	AA	3448	1/1	0.83	0.28	93,93,93,93	0
59	MG	BA	1765	1/1	0.83	0.17	82,82,82,82	0
59	MG	DA	4398	1/1	0.83	0.24	46,46,46,46	0
59	MG	D0	208	1/1	0.83	0.24	72,72,72,72	0
59	MG	AA	3189	1/1	0.83	0.13	52,52,52,52	0
59	MG	CA	2237	1/1	0.83	0.17	62,62,62,62	0
59	MG	AA	3188	1/1	0.83	0.22	52,52,52,52	0
59	MG	BA	2087	1/1	0.83	0.20	81,81,81,81	0
59	MG	DA	4296	1/1	0.83	0.63	105,105,105,105	0
59	MG	DA	3829	1/1	0.83	0.12	66,66,66,66	0
59	MG	DA	3948	1/1	0.83	0.16	127,127,127,127	0
59	MG	CA	2045	1/1	0.83	0.18	66,66,66,66	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3514	1/1	0.83	0.32	94,94,94,94	0
59	MG	CA	1712	1/1	0.83	0.34	67,67,67,67	0
59	MG	CA	1784	1/1	0.84	0.20	54,54,54,54	0
59	MG	CA	2305	1/1	0.84	0.25	81,81,81,81	0
59	MG	BA	2113	1/1	0.84	0.30	92,92,92,92	0
59	MG	AA	3541	1/1	0.84	0.18	49,49,49,49	0
59	MG	CA	2321	1/1	0.84	0.19	90,90,90,90	0
59	MG	BA	1852	1/1	0.84	0.17	105,105,105,105	0
59	MG	DA	4693	1/1	0.84	0.30	62,62,62,62	0
59	MG	DA	3765	1/1	0.84	0.28	61,61,61,61	0
59	MG	DA	5044	1/1	0.84	0.49	93,93,93,93	0
59	MG	AA	4095	1/1	0.84	0.15	58,58,58,58	0
59	MG	CB	109	1/1	0.84	0.10	60,60,60,60	0
59	MG	AR	204	1/1	0.84	0.08	76,76,76,76	0
59	MG	AD	312	1/1	0.84	0.22	79,79,79,79	0
59	MG	DA	4004	1/1	0.84	0.62	90,90,90,90	0
59	MG	BA	1676	1/1	0.84	0.17	84,84,84,84	0
59	MG	DA	4775	1/1	0.84	0.40	86,86,86,86	0
59	MG	BA	2221	1/1	0.84	0.13	75,75,75,75	0
59	MG	AA	3407	1/1	0.84	0.43	78,78,78,78	0
59	MG	DA	3652	1/1	0.84	0.16	93,93,93,93	0
59	MG	BA	1718	1/1	0.84	0.26	43,43,43,43	0
59	MG	DA	4576	1/1	0.84	0.39	87,87,87,87	0
59	MG	AA	3065	1/1	0.84	0.17	60,60,60,60	0
59	MG	DA	4982	1/1	0.84	0.60	53,53,53,53	0
59	MG	BS	109	1/1	0.84	0.15	110,110,110,110	0
59	MG	BH	204	1/1	0.84	0.17	72,72,72,72	0
59	MG	AA	3976	1/1	0.84	0.20	86,86,86,86	0
59	MG	AA	3827	1/1	0.84	0.21	62,62,62,62	0
59	MG	BA	1827	1/1	0.84	0.17	101,101,101,101	0
59	MG	DA	3882	1/1	0.84	0.36	43,43,43,43	0
59	MG	AA	3569	1/1	0.84	0.19	75,75,75,75	0
59	MG	BA	2106	1/1	0.84	0.31	104,104,104,104	0
59	MG	CA	1940	1/1	0.84	0.26	67,67,67,67	0
59	MG	BA	2063	1/1	0.84	0.11	65,65,65,65	0
59	MG	BA	1943	1/1	0.84	0.14	72,72,72,72	0
59	MG	DA	3077	1/1	0.84	0.15	47,47,47,47	0
59	MG	AA	3766	1/1	0.84	0.20	86,86,86,86	0
59	MG	DA	3809	1/1	0.84	0.20	82,82,82,82	0
59	MG	AA	3393	1/1	0.84	0.15	63,63,63,63	0
59	MG	AA	3958	1/1	0.84	0.31	94,94,94,94	0
59	MG	AA	3421	1/1	0.84	0.12	43,43,43,43	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3697	1/1	0.84	0.28	78,78,78,78	0
59	MG	DA	4663	1/1	0.84	0.21	69,69,69,69	0
59	MG	CA	1885	1/1	0.84	0.29	49,49,49,49	0
59	MG	BA	2032	1/1	0.84	0.24	70,70,70,70	0
59	MG	DA	4409	1/1	0.84	0.25	75,75,75,75	0
59	MG	DA	4125	1/1	0.84	0.32	76,76,76,76	0
59	MG	DA	3610	1/1	0.84	0.12	45,45,45,45	0
59	MG	DA	3646	1/1	0.84	0.27	53,53,53,53	0
59	MG	CA	2219	1/1	0.84	0.22	57,57,57,57	0
59	MG	BA	2060	1/1	0.84	0.38	73,73,73,73	0
59	MG	DA	3118	1/1	0.84	0.30	69,69,69,69	0
59	MG	CA	1743	1/1	0.84	0.17	81,81,81,81	0
59	MG	BA	1905	1/1	0.84	0.11	74,74,74,74	0
59	MG	BA	2237	1/1	0.84	0.11	77,77,77,77	0
59	MG	AA	3182	1/1	0.84	0.18	60,60,60,60	0
59	MG	AA	3959	1/1	0.84	0.33	96,96,96,96	0
59	MG	CA	1851	1/1	0.84	0.10	100,100,100,100	0
59	MG	AA	3374	1/1	0.84	0.15	46,46,46,46	0
59	MG	DA	4252	1/1	0.84	0.19	60,60,60,60	0
59	MG	CA	1766	1/1	0.84	0.22	59,59,59,59	0
59	MG	DA	4544	1/1	0.84	0.30	76,76,76,76	0
59	MG	AA	3886	1/1	0.84	0.38	103,103,103,103	0
59	MG	CA	1780	1/1	0.84	0.28	56,56,56,56	0
59	MG	DA	3320	1/1	0.84	0.13	14,14,14,14	0
59	MG	AA	3575	1/1	0.84	0.31	51,51,51,51	0
59	MG	AA	4022	1/1	0.84	0.17	62,62,62,62	0
59	MG	DU	205	1/1	0.84	0.34	75,75,75,75	0
59	MG	CA	1860	1/1	0.84	0.26	107,107,107,107	0
59	MG	BA	1862	1/1	0.84	0.12	77,77,77,77	0
59	MG	DA	4080	1/1	0.84	0.34	89,89,89,89	0
59	MG	AA	3576	1/1	0.84	0.17	37,37,37,37	0
59	MG	DA	4403	1/1	0.84	0.28	83,83,83,83	0
59	MG	CA	1987	1/1	0.84	0.23	73,73,73,73	0
59	MG	DA	4734	1/1	0.84	0.40	71,71,71,71	0
59	MG	CG	309	1/1	0.84	0.36	78,78,78,78	0
59	MG	DA	3451	1/1	0.84	0.51	76,76,76,76	0
59	MG	AA	3543	1/1	0.84	0.14	77,77,77,77	0
59	MG	CA	2235	1/1	0.84	0.23	70,70,70,70	0
59	MG	DA	4379	1/1	0.84	0.15	74,74,74,74	0
59	MG	AA	3740	1/1	0.84	0.13	60,60,60,60	0
59	MG	DA	4745	1/1	0.84	0.28	53,53,53,53	0
59	MG	BK	203	1/1	0.84	0.09	69,69,69,69	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	2316	1/1	0.84	0.26	75,75,75,75	0
59	MG	CC	112	1/1	0.84	0.13	63,63,63,63	0
59	MG	AA	3831	1/1	0.84	0.13	77,77,77,77	0
59	MG	DA	4828	1/1	0.84	0.29	103,103,103,103	0
59	MG	DA	4798	1/1	0.84	0.11	72,72,72,72	0
59	MG	DA	3411	1/1	0.84	0.20	56,56,56,56	0
59	MG	DT	102	1/1	0.84	0.23	60,60,60,60	0
59	MG	AA	3320	1/1	0.84	0.20	69,69,69,69	0
59	MG	BA	2255	1/1	0.84	0.14	61,61,61,61	0
59	MG	AA	3544	1/1	0.84	0.19	62,62,62,62	0
59	MG	DA	3439	1/1	0.84	0.34	86,86,86,86	0
59	MG	CA	1680	1/1	0.84	0.23	40,40,40,40	0
59	MG	CA	1791	1/1	0.84	0.15	87,87,87,87	0
59	MG	AA	3712	1/1	0.84	0.31	57,57,57,57	0
59	MG	CD	127	1/1	0.84	0.55	101,101,101,101	0
59	MG	CA	1715	1/1	0.84	0.12	62,62,62,62	0
59	MG	DA	4597	1/1	0.84	0.15	59,59,59,59	0
59	MG	DB	212	1/1	0.84	0.27	71,71,71,71	0
59	MG	CA	1910	1/1	0.84	0.24	54,54,54,54	0
59	MG	DA	3471	1/1	0.84	0.21	27,27,27,27	0
59	MG	DA	4770	1/1	0.84	0.29	93,93,93,93	0
59	MG	AA	3102	1/1	0.84	0.15	65,65,65,65	0
59	MG	DA	3659	1/1	0.84	0.23	61,61,61,61	0
59	MG	DA	4430	1/1	0.84	0.18	83,83,83,83	0
59	MG	DA	5051	1/1	0.84	0.40	103,103,103,103	0
59	MG	BC	114	1/1	0.84	0.20	58,58,58,58	0
59	MG	DA	4776	1/1	0.84	0.07	72,72,72,72	0
59	MG	DA	3736	1/1	0.84	0.17	100,100,100,100	0
59	MG	AA	3235	1/1	0.84	0.21	48,48,48,48	0
59	MG	DA	4222	1/1	0.84	0.29	54,54,54,54	0
59	MG	CA	1730	1/1	0.84	0.10	38,38,38,38	0
59	MG	CA	2059	1/1	0.84	0.15	95,95,95,95	0
59	MG	CA	2016	1/1	0.84	0.46	80,80,80,80	0
59	MG	DA	3949	1/1	0.84	0.15	132,132,132,132	0
59	MG	BA	1780	1/1	0.84	0.48	85,85,85,85	0
59	MG	AA	3332	1/1	0.84	0.10	83,83,83,83	0
59	MG	DA	3343	1/1	0.84	0.36	58,58,58,58	0
59	MG	AA	4066	1/1	0.84	0.17	82,82,82,82	0
59	MG	DA	4498	1/1	0.84	0.08	82,82,82,82	0
59	MG	DA	4285	1/1	0.84	0.36	70,70,70,70	0
59	MG	AA	3271	1/1	0.84	0.31	83,83,83,83	0
59	MG	BA	1704	1/1	0.84	0.18	67,67,67,67	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DB	227	1/1	0.84	0.14	59,59,59,59	0
59	MG	AA	3789	1/1	0.84	0.18	55,55,55,55	0
59	MG	D2	209	1/1	0.84	0.32	74,74,74,74	0
59	MG	AA	4142	1/1	0.84	0.39	104,104,104,104	0
59	MG	DA	3208	1/1	0.84	0.26	72,72,72,72	0
59	MG	AA	3146	1/1	0.84	0.24	78,78,78,78	0
59	MG	AA	3672	1/1	0.84	0.16	55,55,55,55	0
59	MG	DA	4592	1/1	0.84	0.29	77,77,77,77	0
59	MG	DA	4066	1/1	0.84	0.11	59,59,59,59	0
59	MG	DA	3154	1/1	0.84	0.29	46,46,46,46	0
59	MG	DA	4741	1/1	0.84	0.20	62,62,62,62	0
59	MG	DA	3661	1/1	0.84	0.15	68,68,68,68	0
59	MG	AA	3350	1/1	0.84	0.12	36,36,36,36	0
59	MG	AA	3445	1/1	0.84	0.13	63,63,63,63	0
59	MG	DA	4925	1/1	0.84	0.28	73,73,73,73	0
59	MG	DA	5043	1/1	0.84	0.10	73,73,73,73	0
59	MG	DA	4724	1/1	0.84	0.24	89,89,89,89	0
59	MG	CA	2286	1/1	0.84	0.16	67,67,67,67	0
59	MG	CD	102	1/1	0.84	0.20	64,64,64,64	0
59	MG	DA	3507	1/1	0.84	0.20	66,66,66,66	0
59	MG	DA	4864	1/1	0.84	0.27	93,93,93,93	0
59	MG	DA	3761	1/1	0.84	0.15	90,90,90,90	0
59	MG	CA	2204	1/1	0.85	0.28	87,87,87,87	0
59	MG	BA	1707	1/1	0.85	0.35	59,59,59,59	0
59	MG	C1	102	1/1	0.85	0.22	68,68,68,68	0
59	MG	CA	2248	1/1	0.85	0.18	72,72,72,72	0
59	MG	AA	3885	1/1	0.85	0.19	51,51,51,51	0
59	MG	CC	111	1/1	0.85	0.31	68,68,68,68	0
59	MG	AH	201	1/1	0.85	0.09	78,78,78,78	0
59	MG	AF	306	1/1	0.85	0.17	77,77,77,77	0
59	MG	DB	268	1/1	0.85	0.25	81,81,81,81	0
59	MG	DA	4947	1/1	0.85	0.33	87,87,87,87	0
59	MG	AA	3693	1/1	0.85	0.16	52,52,52,52	0
59	MG	AA	3760	1/1	0.85	0.36	85,85,85,85	0
59	MG	AA	3752	1/1	0.85	0.21	71,71,71,71	0
59	MG	DA	3230	1/1	0.85	0.23	43,43,43,43	0
59	MG	DA	3366	1/1	0.85	0.13	35,35,35,35	0
59	MG	AA	3163	1/1	0.85	0.22	65,65,65,65	0
59	MG	AA	3219	1/1	0.85	0.30	81,81,81,81	0
59	MG	CA	1603	1/1	0.85	0.42	80,80,80,80	0
59	MG	CA	2182	1/1	0.85	0.24	78,78,78,78	0
59	MG	DA	3974	1/1	0.85	0.42	61,61,61,61	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	2092	1/1	0.85	0.43	94,94,94,94	0
59	MG	DA	3538	1/1	0.85	0.20	79,79,79,79	0
59	MG	D7	105	1/1	0.85	0.17	52,52,52,52	0
59	MG	DA	3435	1/1	0.85	0.21	49,49,49,49	0
59	MG	DA	4685	1/1	0.85	0.21	64,64,64,64	0
59	MG	DA	4477	1/1	0.85	0.19	61,61,61,61	0
59	MG	AA	4025	1/1	0.85	0.23	86,86,86,86	0
59	MG	DZ	101	1/1	0.85	0.21	87,87,87,87	0
59	MG	CA	1849	1/1	0.85	0.21	75,75,75,75	0
59	MG	CA	1846	1/1	0.85	0.13	90,90,90,90	0
59	MG	BA	1957	1/1	0.85	0.41	80,80,80,80	0
59	MG	BA	2075	1/1	0.85	0.53	101,101,101,101	0
59	MG	CA	1915	1/1	0.85	0.12	108,108,108,108	0
59	MG	AA	3432	1/1	0.85	0.30	83,83,83,83	0
59	MG	DA	3797	1/1	0.85	0.20	95,95,95,95	0
59	MG	CA	1693	1/1	0.85	0.21	60,60,60,60	0
59	MG	DA	3981	1/1	0.85	0.51	87,87,87,87	0
59	MG	AA	3935	1/1	0.85	0.14	55,55,55,55	0
59	MG	CA	1800	1/1	0.85	0.07	89,89,89,89	0
59	MG	AA	3584	1/1	0.85	0.55	85,85,85,85	0
59	MG	AA	3738	1/1	0.85	0.19	66,66,66,66	0
59	MG	BA	1894	1/1	0.85	0.47	83,83,83,83	0
59	MG	BA	1743	1/1	0.85	0.08	43,43,43,43	0
59	MG	DA	4327	1/1	0.85	0.20	51,51,51,51	0
59	MG	DU	219	1/1	0.85	0.19	72,72,72,72	0
59	MG	AA	3751	1/1	0.85	0.22	48,48,48,48	0
59	MG	CA	2162	1/1	0.85	0.09	79,79,79,79	0
59	MG	BA	1903	1/1	0.85	0.04	71,71,71,71	0
59	MG	DA	5060	1/1	0.85	0.21	64,64,64,64	0
59	MG	DA	3721	1/1	0.85	0.15	51,51,51,51	0
59	MG	DA	4042	1/1	0.85	0.34	71,71,71,71	0
59	MG	AA	3729	1/1	0.85	0.33	69,69,69,69	0
59	MG	CA	1683	1/1	0.85	0.20	47,47,47,47	0
59	MG	DA	3302	1/1	0.85	0.34	50,50,50,50	0
59	MG	AA	4017	1/1	0.85	0.14	96,96,96,96	0
59	MG	AA	3797	1/1	0.85	0.28	96,96,96,96	0
59	MG	DA	4044	1/1	0.85	0.09	107,107,107,107	0
59	MG	DA	3653	1/1	0.85	0.30	61,61,61,61	0
59	MG	AA	3367	1/1	0.85	0.28	49,49,49,49	0
59	MG	BA	1951	1/1	0.85	0.34	78,78,78,78	0
59	MG	DA	4543	1/1	0.85	0.13	60,60,60,60	0
59	MG	AA	3088	1/1	0.85	0.15	51,51,51,51	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	1813	1/1	0.85	0.33	116,116,116,116	0
59	MG	BA	1964	1/1	0.85	0.46	78,78,78,78	0
59	MG	DA	3523	1/1	0.85	0.41	82,82,82,82	0
59	MG	AA	4118	1/1	0.85	0.13	67,67,67,67	0
59	MG	DA	3722	1/1	0.85	0.20	83,83,83,83	0
59	MG	CA	1914	1/1	0.85	0.12	58,58,58,58	0
59	MG	DA	3409	1/1	0.85	0.24	44,44,44,44	0
59	MG	DA	4960	1/1	0.85	0.25	77,77,77,77	0
59	MG	CA	2007	1/1	0.85	0.31	78,78,78,78	0
59	MG	BE	304	1/1	0.85	0.38	85,85,85,85	0
59	MG	AA	3317	1/1	0.85	0.17	57,57,57,57	0
59	MG	DA	4407	1/1	0.85	0.34	85,85,85,85	0
59	MG	DA	4458	1/1	0.85	0.16	51,51,51,51	0
59	MG	AA	3354	1/1	0.85	0.07	53,53,53,53	0
59	MG	CA	1795	1/1	0.85	0.19	109,109,109,109	0
59	MG	DA	4397	1/1	0.85	0.14	56,56,56,56	0
59	MG	DA	4881	1/1	0.85	0.26	85,85,85,85	0
59	MG	AA	4050	1/1	0.85	0.09	70,70,70,70	0
59	MG	DV	303	1/1	0.85	0.18	67,67,67,67	0
59	MG	AB	206	1/1	0.85	0.16	76,76,76,76	0
59	MG	DA	5019	1/1	0.85	0.21	62,62,62,62	0
59	MG	BA	2195	1/1	0.85	0.14	67,67,67,67	0
59	MG	AA	4013	1/1	0.85	0.09	119,119,119,119	0
59	MG	DA	4837	1/1	0.85	0.18	55,55,55,55	0
59	MG	DA	4405	1/1	0.85	0.17	40,40,40,40	0
59	MG	BA	1721	1/1	0.85	0.20	71,71,71,71	0
59	MG	DA	4231	1/1	0.85	0.49	42,42,42,42	0
59	MG	DA	4386	1/1	0.85	0.18	49,49,49,49	0
59	MG	DA	4347	1/1	0.85	0.23	58,58,58,58	0
59	MG	BA	2051	1/1	0.85	0.22	108,108,108,108	0
59	MG	CA	1844	1/1	0.85	0.13	57,57,57,57	0
59	MG	AA	3198	1/1	0.85	0.20	34,34,34,34	0
59	MG	CA	1698	1/1	0.85	0.35	101,101,101,101	0
59	MG	CA	2179	1/1	0.85	0.23	80,80,80,80	0
59	MG	BA	2046	1/1	0.85	0.17	95,95,95,95	0
59	MG	AA	4001	1/1	0.85	0.25	83,83,83,83	0
59	MG	DA	3640	1/1	0.85	0.18	54,54,54,54	0
59	MG	BA	1933	1/1	0.85	0.29	63,63,63,63	0
59	MG	AA	3533	1/1	0.85	0.18	44,44,44,44	0
59	MG	CA	2187	1/1	0.85	0.42	80,80,80,80	0
59	MG	DA	4050	1/1	0.85	0.21	66,66,66,66	0
59	MG	BA	1886	1/1	0.85	0.37	64,64,64,64	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3931	1/1	0.85	0.20	86,86,86,86	0
59	MG	BA	1699	1/1	0.85	0.23	90,90,90,90	0
59	MG	DA	4173	1/1	0.85	0.18	60,60,60,60	0
59	MG	BA	1650	1/1	0.85	0.06	52,52,52,52	0
59	MG	DA	4505	1/1	0.85	0.27	97,97,97,97	0
59	MG	DB	234	1/1	0.85	0.20	64,64,64,64	0
59	MG	DH	203	1/1	0.85	0.18	78,78,78,78	0
59	MG	DA	3214	1/1	0.85	0.35	57,57,57,57	0
59	MG	DA	5074	1/1	0.85	0.09	67,67,67,67	0
59	MG	DA	3900	1/1	0.85	0.09	143,143,143,143	0
59	MG	DA	3462	1/1	0.85	0.28	56,56,56,56	0
59	MG	AA	3571	1/1	0.85	0.24	57,57,57,57	0
59	MG	AA	3912	1/1	0.85	0.37	91,91,91,91	0
59	MG	AA	3064	1/1	0.85	0.27	90,90,90,90	0
59	MG	DA	4292	1/1	0.85	0.26	62,62,62,62	0
59	MG	DA	3169	1/1	0.85	0.17	59,59,59,59	0
59	MG	AA	3792	1/1	0.85	0.53	82,82,82,82	0
59	MG	DA	3898	1/1	0.85	0.10	130,130,130,130	0
59	MG	DA	5026	1/1	0.85	1.09	42,42,42,42	0
59	MG	DA	4899	1/1	0.85	0.19	65,65,65,65	0
59	MG	AA	3773	1/1	0.85	0.21	76,76,76,76	0
59	MG	DA	3814	1/1	0.85	0.19	67,67,67,67	0
59	MG	DA	4994	1/1	0.85	0.19	98,98,98,98	0
59	MG	DA	3255	1/1	0.85	0.25	72,72,72,72	0
59	MG	CA	1752	1/1	0.85	0.16	53,53,53,53	0
59	MG	DA	4133	1/1	0.85	0.19	47,47,47,47	0
59	MG	DA	4653	1/1	0.85	0.21	69,69,69,69	0
59	MG	AA	3210	1/1	0.85	0.18	39,39,39,39	0
59	MG	CA	2154	1/1	0.85	0.16	75,75,75,75	0
59	MG	DA	5047	1/1	0.85	0.32	79,79,79,79	0
59	MG	AA	3810	1/1	0.85	0.28	85,85,85,85	0
59	MG	DA	4041	1/1	0.85	0.32	101,101,101,101	0
59	MG	DA	4611	1/1	0.85	0.20	82,82,82,82	0
59	MG	DA	4598	1/1	0.85	0.23	71,71,71,71	0
59	MG	DA	3472	1/1	0.85	0.32	55,55,55,55	0
59	MG	DA	4730	1/1	0.85	0.20	57,57,57,57	0
59	MG	BA	2084	1/1	0.85	0.29	90,90,90,90	0
59	MG	BA	2019	1/1	0.85	0.36	80,80,80,80	0
59	MG	AA	3097	1/1	0.85	0.10	73,73,73,73	0
59	MG	CA	1897	1/1	0.85	0.40	71,71,71,71	0
59	MG	CA	2152	1/1	0.85	0.39	42,42,42,42	0
59	MG	BA	1783	1/1	0.85	0.20	81,81,81,81	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3642	1/1	0.85	0.19	42,42,42,42	0
59	MG	DA	3189	1/1	0.85	0.22	26,26,26,26	0
59	MG	DA	4102	1/1	0.85	0.19	70,70,70,70	0
59	MG	DA	3998	1/1	0.86	0.38	46,46,46,46	0
59	MG	BA	2089	1/1	0.86	0.09	56,56,56,56	0
59	MG	DA	4452	1/1	0.86	0.21	50,50,50,50	0
59	MG	BA	1776	1/1	0.86	0.09	79,79,79,79	0
59	MG	CA	1990	1/1	0.86	0.14	64,64,64,64	0
59	MG	DA	4825	1/1	0.86	0.21	63,63,63,63	0
59	MG	AA	3426	1/1	0.86	0.10	64,64,64,64	0
59	MG	DA	4699	1/1	0.86	0.28	92,92,92,92	0
59	MG	AA	3194	1/1	0.86	0.14	56,56,56,56	0
59	MG	DA	4336	1/1	0.86	0.28	73,73,73,73	0
59	MG	DA	5002	1/1	0.86	0.17	89,89,89,89	0
59	MG	AA	4074	1/1	0.86	0.22	94,94,94,94	0
59	MG	CK	203	1/1	0.86	0.46	68,68,68,68	0
59	MG	BA	2146	1/1	0.86	0.27	81,81,81,81	0
59	MG	CA	2252	1/1	0.86	0.13	85,85,85,85	0
59	MG	AA	3988	1/1	0.86	0.27	64,64,64,64	0
59	MG	DA	4621	1/1	0.86	0.30	92,92,92,92	0
59	MG	DA	3643	1/1	0.86	0.24	62,62,62,62	0
59	MG	DA	4860	1/1	0.86	0.26	64,64,64,64	0
59	MG	AA	3096	1/1	0.86	0.18	67,67,67,67	0
59	MG	DA	4801	1/1	0.86	0.18	54,54,54,54	0
59	MG	DA	3518	1/1	0.86	0.21	65,65,65,65	0
59	MG	AA	3420	1/1	0.86	0.26	68,68,68,68	0
59	MG	CA	2190	1/1	0.86	0.13	62,62,62,62	0
59	MG	DA	3668	1/1	0.86	0.13	80,80,80,80	0
59	MG	DA	3567	1/1	0.86	0.17	47,47,47,47	0
59	MG	DH	202	1/1	0.86	0.16	59,59,59,59	0
59	MG	DA	3322	1/1	0.86	0.14	36,36,36,36	0
59	MG	DA	3520	1/1	0.86	0.22	95,95,95,95	0
59	MG	BA	2131	1/1	0.86	0.18	61,61,61,61	0
59	MG	AA	3733	1/1	0.86	0.23	72,72,72,72	0
59	MG	CA	1802	1/1	0.86	0.13	76,76,76,76	0
59	MG	BA	1892	1/1	0.86	0.09	75,75,75,75	0
59	MG	AA	3454	1/1	0.86	0.17	67,67,67,67	0
59	MG	AA	3484	1/1	0.86	0.10	66,66,66,66	0
59	MG	BA	2127	1/1	0.86	0.10	50,50,50,50	0
59	MG	CA	1765	1/1	0.86	0.35	56,56,56,56	0
59	MG	BA	2232	1/1	0.86	0.14	103,103,103,103	0
59	MG	CA	1859	1/1	0.86	0.28	106,106,106,106	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	2106	1/1	0.86	0.11	83,83,83,83	0
59	MG	DA	4954	1/1	0.86	0.11	73,73,73,73	0
59	MG	AA	3596	1/1	0.86	0.14	65,65,65,65	0
59	MG	BA	2166	1/1	0.86	0.12	96,96,96,96	0
59	MG	DA	4455	1/1	0.86	0.21	45,45,45,45	0
59	MG	DA	4280	1/1	0.86	0.09	91,91,91,91	0
59	MG	AA	3813	1/1	0.86	0.13	99,99,99,99	0
59	MG	DA	4513	1/1	0.86	0.16	75,75,75,75	0
59	MG	DA	4955	1/1	0.86	0.31	82,82,82,82	0
59	MG	BA	1744	1/1	0.86	0.20	54,54,54,54	0
59	MG	BA	2256	1/1	0.86	0.27	76,76,76,76	0
59	MG	DA	4559	1/1	0.86	0.26	52,52,52,52	0
59	MG	DA	4352	1/1	0.86	0.30	69,69,69,69	0
59	MG	AA	3689	1/1	0.86	0.09	61,61,61,61	0
59	MG	DA	3905	1/1	0.86	0.21	61,61,61,61	0
59	MG	DA	3311	1/1	0.86	0.19	65,65,65,65	0
59	MG	BA	1646	1/1	0.86	0.15	82,82,82,82	0
59	MG	CA	1913	1/1	0.86	0.15	52,52,52,52	0
59	MG	CA	2197	1/1	0.86	0.13	107,107,107,107	0
59	MG	CA	2185	1/1	0.86	0.11	87,87,87,87	0
59	MG	AR	205	1/1	0.86	0.25	54,54,54,54	0
59	MG	CA	2028	1/1	0.86	0.23	71,71,71,71	0
59	MG	CA	1679	1/1	0.86	0.29	60,60,60,60	0
59	MG	CA	2172	1/1	0.86	0.30	95,95,95,95	0
59	MG	CA	1633	1/1	0.86	0.52	105,105,105,105	0
59	MG	DA	3495	1/1	0.86	0.21	51,51,51,51	0
59	MG	CA	1941	1/1	0.86	0.41	91,91,91,91	0
59	MG	BA	1831	1/1	0.86	0.11	81,81,81,81	0
59	MG	DA	4524	1/1	0.86	0.41	50,50,50,50	0
59	MG	AA	4052	1/1	0.86	0.11	73,73,73,73	0
59	MG	AA	3944	1/1	0.86	0.14	71,71,71,71	0
59	MG	D3	102	1/1	0.86	0.17	47,47,47,47	0
59	MG	BA	2039	1/1	0.86	0.21	138,138,138,138	0
59	MG	CA	1701	1/1	0.86	0.20	56,56,56,56	0
59	MG	DA	3629	1/1	0.86	0.14	29,29,29,29	0
59	MG	CA	1948	1/1	0.86	0.11	97,97,97,97	0
59	MG	DA	4079	1/1	0.86	0.25	66,66,66,66	0
59	MG	CA	2017	1/1	0.86	0.22	57,57,57,57	0
59	MG	CA	1675	1/1	0.86	0.32	53,53,53,53	0
59	MG	AA	3110	1/1	0.86	0.20	59,59,59,59	0
59	MG	CA	2005	1/1	0.86	0.10	55,55,55,55	0
59	MG	DA	5038	1/1	0.86	0.21	84,84,84,84	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3989	1/1	0.86	0.16	74,74,74,74	0
59	MG	DU	208	1/1	0.86	0.21	52,52,52,52	0
59	MG	DA	4362	1/1	0.86	0.44	74,74,74,74	0
59	MG	CA	2198	1/1	0.86	0.11	81,81,81,81	0
59	MG	AA	3539	1/1	0.86	0.09	50,50,50,50	0
59	MG	AA	3460	1/1	0.86	0.13	28,28,28,28	0
59	MG	AA	3288	1/1	0.86	0.08	51,51,51,51	0
59	MG	DA	4385	1/1	0.86	0.15	55,55,55,55	0
59	MG	DO	212	1/1	0.86	0.30	44,44,44,44	0
59	MG	DA	3655	1/1	0.86	0.45	86,86,86,86	0
59	MG	CA	2170	1/1	0.86	0.11	70,70,70,70	0
59	MG	DB	254	1/1	0.86	0.28	66,66,66,66	0
59	MG	AA	3266	1/1	0.86	0.19	73,73,73,73	0
59	MG	AA	4102	1/1	0.86	0.21	90,90,90,90	0
59	MG	CA	1617	1/1	0.86	0.18	33,33,33,33	0
59	MG	DA	3572	1/1	0.86	0.22	51,51,51,51	0
59	MG	DE	309	1/1	0.86	0.45	49,49,49,49	0
59	MG	AA	3750	1/1	0.86	0.10	66,66,66,66	0
59	MG	DU	209	1/1	0.86	0.37	96,96,96,96	0
59	MG	DA	3994	1/1	0.86	0.39	48,48,48,48	0
59	MG	DA	3617	1/1	0.86	0.32	82,82,82,82	0
59	MG	AA	3938	1/1	0.86	0.24	76,76,76,76	0
59	MG	CA	1905	1/1	0.86	0.16	81,81,81,81	0
59	MG	CA	1938	1/1	0.86	0.18	87,87,87,87	0
59	MG	AA	4096	1/1	0.86	0.40	114,114,114,114	0
59	MG	D0	205	1/1	0.86	0.35	41,41,41,41	0
59	MG	CA	2118	1/1	0.86	0.18	82,82,82,82	0
59	MG	DA	3207	1/1	0.86	0.34	45,45,45,45	0
59	MG	DA	3901	1/1	0.86	0.22	30,30,30,30	0
59	MG	BA	1657	1/1	0.86	0.29	63,63,63,63	0
59	MG	AA	3749	1/1	0.86	0.36	33,33,33,33	0
59	MG	AB	216	1/1	0.86	0.23	53,53,53,53	0
59	MG	AA	4033	1/1	0.86	0.34	83,83,83,83	0
59	MG	DA	4493	1/1	0.86	0.30	107,107,107,107	0
59	MG	AA	3369	1/1	0.86	0.15	81,81,81,81	0
59	MG	AA	3524	1/1	0.86	0.14	50,50,50,50	0
59	MG	DA	4497	1/1	0.86	0.36	43,43,43,43	0
59	MG	DA	4600	1/1	0.86	0.23	57,57,57,57	0
59	MG	BA	2044	1/1	0.86	0.09	62,62,62,62	0
59	MG	CA	1770	1/1	0.86	0.16	56,56,56,56	0
59	MG	CA	1678	1/1	0.86	0.49	65,65,65,65	0
59	MG	CA	2324	1/1	0.86	0.30	89,89,89,89	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3483	1/1	0.86	0.26	61,61,61,61	0
59	MG	DA	3835	1/1	0.86	0.16	61,61,61,61	0
59	MG	AA	3736	1/1	0.86	0.18	72,72,72,72	0
59	MG	AA	3557	1/1	0.86	0.32	89,89,89,89	0
59	MG	DA	4580	1/1	0.86	0.18	69,69,69,69	0
59	MG	DA	3119	1/1	0.86	0.25	60,60,60,60	0
59	MG	BA	1981	1/1	0.86	0.11	68,68,68,68	0
59	MG	CA	2060	1/1	0.86	0.12	44,44,44,44	0
59	MG	DA	4210	1/1	0.86	0.21	53,53,53,53	0
59	MG	DA	4143	1/1	0.86	0.27	63,63,63,63	0
59	MG	BA	2225	1/1	0.86	0.13	98,98,98,98	0
59	MG	BA	1624	1/1	0.86	0.18	84,84,84,84	0
59	MG	DG	205	1/1	0.86	0.30	64,64,64,64	0
59	MG	BA	2178	1/1	0.86	0.28	141,141,141,141	0
59	MG	CA	1996	1/1	0.86	0.25	94,94,94,94	0
59	MG	BA	2242	1/1	0.86	0.18	57,57,57,57	0
59	MG	DA	3907	1/1	0.86	0.15	83,83,83,83	0
59	MG	CA	1745	1/1	0.86	0.26	56,56,56,56	0
59	MG	AA	3884	1/1	0.86	0.16	94,94,94,94	0
59	MG	AA	3180	1/1	0.86	0.14	62,62,62,62	0
59	MG	DA	4246	1/1	0.86	0.19	64,64,64,64	0
59	MG	DA	3161	1/1	0.86	0.31	55,55,55,55	0
59	MG	CC	124	1/1	0.86	0.30	77,77,77,77	0
59	MG	DA	3547	1/1	0.86	0.41	68,68,68,68	0
59	MG	DA	4983	1/1	0.86	0.34	53,53,53,53	0
59	MG	AA	3334	1/1	0.86	0.17	46,46,46,46	0
59	MG	AA	3632	1/1	0.86	0.22	43,43,43,43	0
59	MG	AB	217	1/1	0.86	0.37	87,87,87,87	0
59	MG	B1	104	1/1	0.86	0.09	116,116,116,116	0
59	MG	DA	3728	1/1	0.86	0.27	42,42,42,42	0
59	MG	DA	4159	1/1	0.86	0.18	62,62,62,62	0
59	MG	AF	305	1/1	0.86	0.27	89,89,89,89	0
59	MG	CA	2127	1/1	0.86	0.43	88,88,88,88	0
59	MG	DA	3920	1/1	0.86	0.21	50,50,50,50	0
59	MG	DB	225	1/1	0.86	0.14	76,76,76,76	0
59	MG	CA	2268	1/1	0.86	0.27	86,86,86,86	0
59	MG	DA	4961	1/1	0.86	0.21	68,68,68,68	0
59	MG	BA	2092	1/1	0.86	0.15	77,77,77,77	0
59	MG	AA	4105	1/1	0.86	0.10	106,106,106,106	0
59	MG	BA	1768	1/1	0.86	0.20	75,75,75,75	0
59	MG	CA	1919	1/1	0.86	0.13	50,50,50,50	0
59	MG	CA	1732	1/1	0.86	0.17	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3751	1/1	0.86	0.17	97,97,97,97	0
59	MG	CA	2048	1/1	0.86	0.15	77,77,77,77	0
59	MG	AA	3213	1/1	0.86	0.18	51,51,51,51	0
59	MG	BA	2152	1/1	0.86	0.20	83,83,83,83	0
59	MG	DA	3864	1/1	0.86	0.15	69,69,69,69	0
59	MG	AA	3612	1/1	0.86	0.28	106,106,106,106	0
59	MG	DA	3465	1/1	0.86	0.38	56,56,56,56	0
59	MG	BA	1784	1/1	0.86	0.09	54,54,54,54	0
59	MG	AA	3925	1/1	0.86	0.17	71,71,71,71	0
59	MG	DA	4127	1/1	0.86	0.29	70,70,70,70	0
59	MG	AA	3054	1/1	0.86	0.10	44,44,44,44	0
59	MG	AA	3628	1/1	0.86	0.35	62,62,62,62	0
59	MG	CA	1988	1/1	0.87	0.10	57,57,57,57	0
59	MG	DA	4583	1/1	0.87	0.17	51,51,51,51	0
59	MG	AA	3499	1/1	0.87	0.29	101,101,101,101	0
59	MG	BA	2207	1/1	0.87	0.42	103,103,103,103	0
59	MG	BA	2114	1/1	0.87	0.10	71,71,71,71	0
59	MG	CA	1878	1/1	0.87	0.15	26,26,26,26	0
59	MG	BA	2268	1/1	0.87	0.17	87,87,87,87	0
59	MG	AB	230	1/1	0.87	0.16	76,76,76,76	0
59	MG	AA	4090	1/1	0.87	0.11	78,78,78,78	0
59	MG	AA	4154	1/1	0.87	0.20	89,89,89,89	0
59	MG	DA	3333	1/1	0.87	0.19	78,78,78,78	0
59	MG	DA	3975	1/1	0.87	0.14	50,50,50,50	0
59	MG	DA	3424	1/1	0.87	0.12	47,47,47,47	0
59	MG	CQ	104	1/1	0.87	0.08	56,56,56,56	0
59	MG	DA	4571	1/1	0.87	0.15	61,61,61,61	0
59	MG	AA	3329	1/1	0.87	0.12	66,66,66,66	0
59	MG	AA	3830	1/1	0.87	0.08	83,83,83,83	0
59	MG	AA	3036	1/1	0.87	0.07	36,36,36,36	0
59	MG	CA	1865	1/1	0.87	0.19	67,67,67,67	0
59	MG	DA	4720	1/1	0.87	0.13	88,88,88,88	0
59	MG	BA	2096	1/1	0.87	0.27	51,51,51,51	0
59	MG	DA	4029	1/1	0.87	0.16	65,65,65,65	0
59	MG	BC	106	1/1	0.87	0.14	58,58,58,58	0
59	MG	CC	125	1/1	0.87	0.14	52,52,52,52	0
59	MG	DA	3600	1/1	0.87	0.20	59,59,59,59	0
59	MG	BA	2091	1/1	0.87	0.37	108,108,108,108	0
59	MG	AB	215	1/1	0.87	0.14	70,70,70,70	0
59	MG	DA	4422	1/1	0.87	0.31	71,71,71,71	0
59	MG	DA	4604	1/1	0.87	0.17	111,111,111,111	0
59	MG	BA	2080	1/1	0.87	0.12	61,61,61,61	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	4084	1/1	0.87	0.08	86,86,86,86	0
59	MG	AA	3529	1/1	0.87	0.15	29,29,29,29	0
59	MG	CC	118	1/1	0.87	0.24	66,66,66,66	0
59	MG	DA	3644	1/1	0.87	0.35	84,84,84,84	0
59	MG	DA	3715	1/1	0.87	0.21	75,75,75,75	0
59	MG	DA	3560	1/1	0.87	0.19	36,36,36,36	0
59	MG	BA	2272	1/1	0.87	0.23	75,75,75,75	0
59	MG	AA	3052	1/1	0.87	0.17	60,60,60,60	0
59	MG	DA	3990	1/1	0.87	0.13	31,31,31,31	0
59	MG	AA	3690	1/1	0.87	0.20	64,64,64,64	0
59	MG	BA	1959	1/1	0.87	0.21	64,64,64,64	0
59	MG	BA	1680	1/1	0.87	0.11	55,55,55,55	0
59	MG	CC	108	1/1	0.87	0.14	57,57,57,57	0
59	MG	AA	4032	1/1	0.87	0.46	102,102,102,102	0
59	MG	AA	3171	1/1	0.87	0.45	50,50,50,50	0
59	MG	C1	104	1/1	0.87	0.24	71,71,71,71	0
59	MG	DA	5063	1/1	0.87	0.39	52,52,52,52	0
59	MG	DA	3496	1/1	0.87	0.14	47,47,47,47	0
59	MG	DA	4794	1/1	0.87	0.35	86,86,86,86	0
59	MG	BA	2171	1/1	0.87	0.34	132,132,132,132	0
59	MG	DA	3522	1/1	0.87	0.35	63,63,63,63	0
59	MG	D7	101	1/1	0.87	0.33	52,52,52,52	0
59	MG	DA	4532	1/1	0.87	0.44	45,45,45,45	0
59	MG	DA	3815	1/1	0.87	0.21	83,83,83,83	0
59	MG	CA	1627	1/1	0.87	0.24	49,49,49,49	0
59	MG	AT	103	1/1	0.87	0.21	85,85,85,85	0
59	MG	A8	104	1/1	0.87	0.29	68,68,68,68	0
59	MG	DA	4345	1/1	0.87	0.12	58,58,58,58	0
59	MG	BA	2158	1/1	0.87	0.41	77,77,77,77	0
59	MG	CA	1907	1/1	0.87	0.26	66,66,66,66	0
59	MG	DD	301	1/1	0.87	0.41	45,45,45,45	0
59	MG	BA	1700	1/1	0.87	0.23	46,46,46,46	0
59	MG	AA	3540	1/1	0.87	0.27	61,61,61,61	0
59	MG	CA	1950	1/1	0.87	0.21	86,86,86,86	0
59	MG	AA	4080	1/1	0.87	0.47	71,71,71,71	0
59	MG	A7	101	1/1	0.87	0.73	77,77,77,77	0
59	MG	BA	1683	1/1	0.87	0.15	51,51,51,51	0
59	MG	A7	103	1/1	0.87	0.30	45,45,45,45	0
59	MG	DR	204	1/1	0.87	0.19	40,40,40,40	0
59	MG	DA	4218	1/1	0.87	0.10	51,51,51,51	0
59	MG	DA	4522	1/1	0.87	0.25	64,64,64,64	0
59	MG	DA	4909	1/1	0.87	0.26	70,70,70,70	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CG	306	1/1	0.87	0.11	124,124,124,124	0
59	MG	DA	4028	1/1	0.87	0.24	48,48,48,48	0
59	MG	AA	3025	1/1	0.87	0.29	38,38,38,38	0
59	MG	AA	3358	1/1	0.87	0.15	95,95,95,95	0
59	MG	BA	1934	1/1	0.87	0.54	84,84,84,84	0
59	MG	CA	1746	1/1	0.87	0.35	62,62,62,62	0
59	MG	DA	4258	1/1	0.87	0.32	64,64,64,64	0
59	MG	DA	3825	1/1	0.87	0.16	77,77,77,77	0
59	MG	BA	1748	1/1	0.87	0.30	66,66,66,66	0
59	MG	AA	3476	1/1	0.87	0.16	47,47,47,47	0
59	MG	DA	3193	1/1	0.87	0.15	71,71,71,71	0
59	MG	BA	1661	1/1	0.87	0.13	83,83,83,83	0
59	MG	CA	2043	1/1	0.87	0.51	100,100,100,100	0
59	MG	DA	3828	1/1	0.87	0.19	70,70,70,70	0
59	MG	BB	107	1/1	0.87	0.21	83,83,83,83	0
59	MG	BA	1940	1/1	0.87	0.36	79,79,79,79	0
59	MG	AG	201	1/1	0.87	0.12	86,86,86,86	0
59	MG	AA	3772	1/1	0.87	0.15	103,103,103,103	0
59	MG	DA	3690	1/1	0.87	0.27	111,111,111,111	0
59	MG	DA	4414	1/1	0.87	0.16	59,59,59,59	0
59	MG	DB	266	1/1	0.87	0.18	51,51,51,51	0
59	MG	BD	106	1/1	0.87	0.40	91,91,91,91	0
59	MG	BA	1806	1/1	0.87	0.16	78,78,78,78	0
59	MG	AA	3965	1/1	0.87	0.09	111,111,111,111	0
59	MG	CA	2138	1/1	0.87	0.42	91,91,91,91	0
59	MG	AB	234	1/1	0.87	0.25	105,105,105,105	0
59	MG	DA	4740	1/1	0.87	0.29	78,78,78,78	0
59	MG	DA	3689	1/1	0.87	0.20	67,67,67,67	0
59	MG	AA	4021	1/1	0.87	0.09	80,80,80,80	0
59	MG	DU	213	1/1	0.87	0.34	70,70,70,70	0
59	MG	CA	1817	1/1	0.87	0.14	85,85,85,85	0
59	MG	CA	1729	1/1	0.87	0.27	46,46,46,46	0
59	MG	DA	3874	1/1	0.87	0.51	69,69,69,69	0
59	MG	CA	2215	1/1	0.87	0.19	63,63,63,63	0
59	MG	AA	3439	1/1	0.87	0.15	71,71,71,71	0
59	MG	DA	4281	1/1	0.87	0.22	51,51,51,51	0
59	MG	BA	2026	1/1	0.87	0.11	128,128,128,128	0
59	MG	D0	202	1/1	0.87	0.21	62,62,62,62	0
59	MG	AA	3464	1/1	0.87	0.20	61,61,61,61	0
59	MG	CA	2156	1/1	0.87	0.32	92,92,92,92	0
59	MG	BA	2251	1/1	0.87	0.43	89,89,89,89	0
59	MG	AU	203	1/1	0.87	0.23	74,74,74,74	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	BC	115	1/1	0.87	0.13	106,106,106,106	0
59	MG	DB	213	1/1	0.87	0.27	64,64,64,64	0
59	MG	DA	4581	1/1	0.87	0.14	55,55,55,55	0
59	MG	DA	4610	1/1	0.87	0.09	114,114,114,114	0
59	MG	BA	2017	1/1	0.87	0.47	85,85,85,85	0
59	MG	DA	3957	1/1	0.87	0.14	62,62,62,62	0
59	MG	DA	4861	1/1	0.87	0.28	74,74,74,74	0
59	MG	DA	4805	1/1	0.87	0.33	74,74,74,74	0
59	MG	AA	3080	1/1	0.87	0.14	41,41,41,41	0
59	MG	DA	3142	1/1	0.87	0.28	73,73,73,73	0
59	MG	BA	2050	1/1	0.87	0.26	85,85,85,85	0
59	MG	BA	2246	1/1	0.87	0.19	125,125,125,125	0
59	MG	DA	3840	1/1	0.87	0.13	58,58,58,58	0
59	MG	DG	201	1/1	0.87	0.26	68,68,68,68	0
59	MG	DA	4234	1/1	0.87	0.28	37,37,37,37	0
59	MG	DA	4168	1/1	0.87	0.14	33,33,33,33	0
59	MG	AA	4006	1/1	0.87	0.11	65,65,65,65	0
59	MG	BA	1978	1/1	0.87	0.31	73,73,73,73	0
59	MG	AA	3779	1/1	0.87	0.13	86,86,86,86	0
59	MG	DA	4110	1/1	0.87	0.24	51,51,51,51	0
59	MG	BA	2007	1/1	0.87	0.22	106,106,106,106	0
59	MG	BA	1991	1/1	0.87	0.29	91,91,91,91	0
59	MG	AA	3583	1/1	0.87	0.43	72,72,72,72	0
59	MG	DB	215	1/1	0.87	0.32	66,66,66,66	0
59	MG	DA	4343	1/1	0.87	0.30	58,58,58,58	0
59	MG	BA	1898	1/1	0.87	0.15	49,49,49,49	0
59	MG	DA	4342	1/1	0.87	0.29	93,93,93,93	0
59	MG	AD	311	1/1	0.87	0.13	54,54,54,54	0
59	MG	AA	3960	1/1	0.87	0.17	101,101,101,101	0
59	MG	AA	3229	1/1	0.87	0.19	56,56,56,56	0
59	MG	DA	3256	1/1	0.87	0.23	50,50,50,50	0
59	MG	BA	1896	1/1	0.87	0.07	83,83,83,83	0
59	MG	CA	2201	1/1	0.87	0.09	80,80,80,80	0
59	MG	DA	4958	1/1	0.87	0.16	64,64,64,64	0
59	MG	AE	304	1/1	0.87	0.19	83,83,83,83	0
59	MG	DA	3564	1/1	0.87	0.41	83,83,83,83	0
59	MG	BA	1820	1/1	0.87	0.13	84,84,84,84	0
59	MG	AB	208	1/1	0.87	0.09	57,57,57,57	0
59	MG	DA	4349	1/1	0.87	0.31	77,77,77,77	0
59	MG	CA	1845	1/1	0.87	0.09	108,108,108,108	0
59	MG	DA	3588	1/1	0.87	0.14	54,54,54,54	0
59	MG	BA	1853	1/1	0.87	0.23	103,103,103,103	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	BA	1758	1/1	0.87	0.21	54,54,54,54	0
59	MG	DW	101	1/1	0.87	0.26	49,49,49,49	0
59	MG	CA	1961	1/1	0.87	0.30	70,70,70,70	0
59	MG	BA	1668	1/1	0.87	0.12	56,56,56,56	0
59	MG	DA	3656	1/1	0.87	0.24	81,81,81,81	0
59	MG	AA	4018	1/1	0.87	0.34	56,56,56,56	0
59	MG	AA	3863	1/1	0.87	0.18	72,72,72,72	0
59	MG	AA	3590	1/1	0.87	0.10	78,78,78,78	0
59	MG	BA	2210	1/1	0.87	0.09	78,78,78,78	0
59	MG	AA	3611	1/1	0.87	0.08	70,70,70,70	0
59	MG	DA	3623	1/1	0.87	0.33	89,89,89,89	0
59	MG	BA	2197	1/1	0.87	0.25	136,136,136,136	0
59	MG	AA	3294	1/1	0.87	0.13	64,64,64,64	0
59	MG	DA	3263	1/1	0.87	0.15	24,24,24,24	0
59	MG	CA	2309	1/1	0.87	0.31	92,92,92,92	0
59	MG	BA	2201	1/1	0.87	0.29	85,85,85,85	0
59	MG	DA	4051	1/1	0.87	0.22	94,94,94,94	0
59	MG	AA	3461	1/1	0.87	0.19	58,58,58,58	0
59	MG	BA	1888	1/1	0.87	0.28	67,67,67,67	0
59	MG	CA	2022	1/1	0.87	0.32	88,88,88,88	0
59	MG	DA	4821	1/1	0.87	0.14	54,54,54,54	0
59	MG	D1	210	1/1	0.87	0.35	66,66,66,66	0
59	MG	DA	3672	1/1	0.87	0.30	76,76,76,76	0
59	MG	AA	3795	1/1	0.87	0.13	106,106,106,106	0
59	MG	CA	2056	1/1	0.87	0.64	120,120,120,120	0
59	MG	CK	202	1/1	0.87	0.24	79,79,79,79	0
59	MG	AA	3553	1/1	0.87	0.21	73,73,73,73	0
59	MG	DA	4788	1/1	0.87	0.10	97,97,97,97	0
59	MG	AA	3835	1/1	0.87	0.15	56,56,56,56	0
59	MG	CA	1959	1/1	0.87	0.17	56,56,56,56	0
59	MG	AA	3999	1/1	0.87	0.15	57,57,57,57	0
59	MG	DA	3997	1/1	0.87	0.16	57,57,57,57	0
59	MG	AA	3366	1/1	0.87	0.09	53,53,53,53	0
59	MG	DA	4984	1/1	0.88	0.28	90,90,90,90	0
59	MG	DA	4119	1/1	0.88	0.12	47,47,47,47	0
59	MG	BA	1872	1/1	0.88	0.27	103,103,103,103	0
59	MG	DA	4054	1/1	0.88	0.62	108,108,108,108	0
59	MG	BB	105	1/1	0.88	0.10	85,85,85,85	0
59	MG	DA	3397	1/1	0.88	0.21	65,65,65,65	0
59	MG	CA	2274	1/1	0.88	0.16	54,54,54,54	0
59	MG	DA	4566	1/1	0.88	0.18	64,64,64,64	0
59	MG	CA	2293	1/1	0.88	0.13	80,80,80,80	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3384	1/1	0.88	0.32	56,56,56,56	0
59	MG	CD	107	1/1	0.88	0.16	57,57,57,57	0
59	MG	DA	4083	1/1	0.88	0.23	52,52,52,52	0
59	MG	BA	1714	1/1	0.88	0.23	69,69,69,69	0
59	MG	BA	1767	1/1	0.88	0.16	50,50,50,50	0
59	MG	DA	3634	1/1	0.88	0.24	54,54,54,54	0
59	MG	AA	3661	1/1	0.88	0.26	70,70,70,70	0
59	MG	DA	5000	1/1	0.88	0.45	71,71,71,71	0
59	MG	AA	4076	1/1	0.88	0.07	113,113,113,113	0
59	MG	AA	3713	1/1	0.88	0.20	83,83,83,83	0
59	MG	DA	3595	1/1	0.88	0.23	38,38,38,38	0
59	MG	CA	1982	1/1	0.88	0.19	59,59,59,59	0
59	MG	DA	3933	1/1	0.88	0.36	113,113,113,113	0
59	MG	AB	231	1/1	0.88	0.21	94,94,94,94	0
59	MG	DA	4649	1/1	0.88	0.20	41,41,41,41	0
59	MG	DB	253	1/1	0.88	0.15	74,74,74,74	0
59	MG	AA	3364	1/1	0.88	0.21	57,57,57,57	0
59	MG	CA	1972	1/1	0.88	0.30	68,68,68,68	0
59	MG	DA	4719	1/1	0.88	0.23	54,54,54,54	0
59	MG	DA	4935	1/1	0.88	0.70	105,105,105,105	0
59	MG	DA	3431	1/1	0.88	0.36	61,61,61,61	0
59	MG	DA	4456	1/1	0.88	0.32	59,59,59,59	0
59	MG	CA	1953	1/1	0.88	0.07	80,80,80,80	0
59	MG	AE	302	1/1	0.88	0.39	74,74,74,74	0
59	MG	DA	4155	1/1	0.88	0.34	89,89,89,89	0
59	MG	CA	1861	1/1	0.88	0.22	87,87,87,87	0
59	MG	DA	3838	1/1	0.88	0.31	80,80,80,80	0
59	MG	BE	303	1/1	0.88	0.17	94,94,94,94	0
59	MG	DA	4211	1/1	0.88	0.20	47,47,47,47	0
59	MG	DE	314	1/1	0.88	0.69	67,67,67,67	0
59	MG	AA	3022	1/1	0.88	0.24	63,63,63,63	0
59	MG	DB	238	1/1	0.88	0.44	100,100,100,100	0
59	MG	BA	1966	1/1	0.88	0.16	82,82,82,82	0
59	MG	DA	4063	1/1	0.88	0.20	52,52,52,52	0
59	MG	BA	2253	1/1	0.88	0.11	103,103,103,103	0
59	MG	DA	5031	1/1	0.88	0.34	68,68,68,68	0
59	MG	DA	3023	1/1	0.88	0.32	57,57,57,57	0
59	MG	DA	3102	1/1	0.88	0.32	61,61,61,61	0
59	MG	DA	4378	1/1	0.88	0.18	48,48,48,48	0
59	MG	DA	4820	1/1	0.88	0.22	56,56,56,56	0
59	MG	DA	3419	1/1	0.88	0.18	32,32,32,32	0
59	MG	AA	3530	1/1	0.88	0.15	50,50,50,50	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4901	1/1	0.88	0.33	84,84,84,84	0
59	MG	AA	3270	1/1	0.88	0.26	72,72,72,72	0
59	MG	DA	3485	1/1	0.88	0.33	77,77,77,77	0
59	MG	AA	3502	1/1	0.88	0.20	55,55,55,55	0
59	MG	CA	1692	1/1	0.88	0.11	51,51,51,51	0
59	MG	DS	204	1/1	0.88	0.43	75,75,75,75	0
59	MG	BA	2047	1/1	0.88	0.18	107,107,107,107	0
59	MG	CA	2004	1/1	0.88	0.13	61,61,61,61	0
59	MG	CA	2211	1/1	0.88	0.48	90,90,90,90	0
59	MG	CA	1926	1/1	0.88	0.12	88,88,88,88	0
59	MG	BA	1990	1/1	0.88	0.20	82,82,82,82	0
59	MG	DV	304	1/1	0.88	0.27	77,77,77,77	0
59	MG	AA	3362	1/1	0.88	0.11	62,62,62,62	0
59	MG	DA	4311	1/1	0.88	0.31	83,83,83,83	0
59	MG	DA	4603	1/1	0.88	0.23	55,55,55,55	0
59	MG	DA	3805	1/1	0.88	0.30	66,66,66,66	0
59	MG	AA	3917	1/1	0.88	0.30	98,98,98,98	0
59	MG	DA	3945	1/1	0.88	0.08	58,58,58,58	0
59	MG	DA	4903	1/1	0.88	0.13	69,69,69,69	0
59	MG	DA	4227	1/1	0.88	0.08	51,51,51,51	0
59	MG	DA	3863	1/1	0.88	0.23	69,69,69,69	0
59	MG	BA	2125	1/1	0.88	0.11	65,65,65,65	0
59	MG	DA	4264	1/1	0.88	0.17	58,58,58,58	0
59	MG	DA	4726	1/1	0.88	0.29	71,71,71,71	0
59	MG	DA	3570	1/1	0.88	0.38	71,71,71,71	0
59	MG	DA	3215	1/1	0.88	0.19	84,84,84,84	0
59	MG	DA	3892	1/1	0.88	0.51	46,46,46,46	0
59	MG	CA	2040	1/1	0.88	0.15	56,56,56,56	0
59	MG	AA	3770	1/1	0.88	0.11	70,70,70,70	0
59	MG	DA	4869	1/1	0.88	0.24	44,44,44,44	0
59	MG	DA	4506	1/1	0.88	0.30	65,65,65,65	0
59	MG	D5	107	1/1	0.88	0.27	59,59,59,59	0
59	MG	BA	2000	1/1	0.88	0.17	93,93,93,93	0
59	MG	AA	3414	1/1	0.88	0.29	56,56,56,56	0
59	MG	AA	3255	1/1	0.88	0.21	46,46,46,46	0
59	MG	CA	1611	1/1	0.88	0.12	39,39,39,39	0
59	MG	CD	122	1/1	0.88	0.38	84,84,84,84	0
59	MG	CA	1984	1/1	0.88	0.28	49,49,49,49	0
59	MG	DA	4760	1/1	0.88	0.23	55,55,55,55	0
59	MG	BA	2057	1/1	0.88	0.10	103,103,103,103	0
59	MG	AO	204	1/1	0.88	0.27	43,43,43,43	0
59	MG	AA	3963	1/1	0.88	0.20	62,62,62,62	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	BA	1667	1/1	0.88	0.22	62,62,62,62	0
59	MG	BA	1864	1/1	0.88	0.09	95,95,95,95	0
59	MG	DA	5028	1/1	0.88	0.18	66,66,66,66	0
59	MG	CA	1902	1/1	0.88	0.21	52,52,52,52	0
59	MG	AA	3083	1/1	0.88	0.15	33,33,33,33	0
59	MG	BA	1734	1/1	0.88	0.14	76,76,76,76	0
59	MG	CA	1702	1/1	0.88	0.14	45,45,45,45	0
59	MG	CE	306	1/1	0.88	0.22	96,96,96,96	0
59	MG	DA	3200	1/1	0.88	0.20	67,67,67,67	0
59	MG	CA	1853	1/1	0.88	0.16	61,61,61,61	0
59	MG	AA	3205	1/1	0.88	0.31	71,71,71,71	0
59	MG	CE	303	1/1	0.88	0.12	83,83,83,83	0
59	MG	DA	3493	1/1	0.88	0.13	50,50,50,50	0
59	MG	DA	4661	1/1	0.88	0.23	76,76,76,76	0
59	MG	DA	4241	1/1	0.88	0.21	55,55,55,55	0
59	MG	BA	2218	1/1	0.88	0.17	80,80,80,80	0
59	MG	CD	121	1/1	0.88	0.10	133,133,133,133	0
59	MG	DA	3474	1/1	0.88	0.26	41,41,41,41	0
59	MG	CA	2116	1/1	0.88	0.14	79,79,79,79	0
59	MG	AA	3622	1/1	0.88	0.11	43,43,43,43	0
59	MG	CB	117	1/1	0.88	0.17	68,68,68,68	0
59	MG	B1	103	1/1	0.88	0.36	67,67,67,67	0
59	MG	CA	2101	1/1	0.88	0.15	68,68,68,68	0
59	MG	AA	3437	1/1	0.88	0.13	51,51,51,51	0
59	MG	BS	104	1/1	0.88	0.16	76,76,76,76	0
59	MG	AA	3867	1/1	0.88	0.10	41,41,41,41	0
59	MG	DA	4791	1/1	0.88	0.34	69,69,69,69	0
59	MG	AA	4147	1/1	0.88	0.22	66,66,66,66	0
59	MG	DA	3184	1/1	0.88	0.27	40,40,40,40	0
59	MG	DF	311	1/1	0.88	0.28	57,57,57,57	0
59	MG	CA	1789	1/1	0.88	0.05	99,99,99,99	0
59	MG	DA	3226	1/1	0.88	0.21	44,44,44,44	0
59	MG	DA	4236	1/1	0.88	0.12	52,52,52,52	0
59	MG	DB	255	1/1	0.88	0.17	63,63,63,63	0
59	MG	CA	2277	1/1	0.88	0.13	103,103,103,103	0
59	MG	CV	102	1/1	0.88	0.15	63,63,63,63	0
59	MG	BA	1773	1/1	0.88	0.19	80,80,80,80	0
59	MG	DE	303	1/1	0.88	0.21	52,52,52,52	0
59	MG	DA	4857	1/1	0.88	0.20	56,56,56,56	0
59	MG	AA	3979	1/1	0.88	0.34	94,94,94,94	0
59	MG	DA	3136	1/1	0.88	0.27	63,63,63,63	0
59	MG	AA	3994	1/1	0.88	0.11	68,68,68,68	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	2058	1/1	0.88	0.20	73,73,73,73	0
59	MG	AA	3816	1/1	0.88	0.44	73,73,73,73	0
59	MG	DH	201	1/1	0.88	0.16	48,48,48,48	0
59	MG	DA	4313	1/1	0.88	0.22	56,56,56,56	0
59	MG	BA	2013	1/1	0.88	0.10	67,67,67,67	0
59	MG	AA	3746	1/1	0.88	0.14	66,66,66,66	0
59	MG	DA	4095	1/1	0.88	0.27	115,115,115,115	0
59	MG	AA	4088	1/1	0.88	0.13	97,97,97,97	0
59	MG	BA	1960	1/1	0.88	0.14	68,68,68,68	0
59	MG	DA	4341	1/1	0.88	0.12	59,59,59,59	0
59	MG	AA	3405	1/1	0.88	0.37	73,73,73,73	0
59	MG	BA	1841	1/1	0.88	0.06	108,108,108,108	0
59	MG	CA	1728	1/1	0.88	0.32	76,76,76,76	0
59	MG	DA	4920	1/1	0.88	0.16	61,61,61,61	0
59	MG	AA	4086	1/1	0.88	0.23	76,76,76,76	0
59	MG	DA	4996	1/1	0.88	0.18	85,85,85,85	0
59	MG	DF	303	1/1	0.88	0.17	28,28,28,28	0
59	MG	CA	1812	1/1	0.88	0.13	45,45,45,45	0
59	MG	BA	1916	1/1	0.88	0.16	56,56,56,56	0
59	MG	DA	4139	1/1	0.88	0.21	36,36,36,36	0
59	MG	DA	4387	1/1	0.88	0.33	57,57,57,57	0
59	MG	D6	102	1/1	0.88	0.15	45,45,45,45	0
59	MG	AA	3818	1/1	0.88	0.34	80,80,80,80	0
59	MG	DA	4057	1/1	0.88	0.32	87,87,87,87	0
59	MG	DA	3758	1/1	0.88	0.24	100,100,100,100	0
59	MG	DA	3513	1/1	0.88	0.32	50,50,50,50	0
59	MG	CC	103	1/1	0.88	0.12	39,39,39,39	0
59	MG	AA	3181	1/1	0.88	0.20	51,51,51,51	0
59	MG	BA	1689	1/1	0.88	0.07	64,64,64,64	0
59	MG	BA	2261	1/1	0.88	0.07	81,81,81,81	0
59	MG	DA	3492	1/1	0.88	0.25	53,53,53,53	0
59	MG	DA	3414	1/1	0.88	0.25	45,45,45,45	0
59	MG	BA	2227	1/1	0.88	0.32	72,72,72,72	0
59	MG	AA	3536	1/1	0.88	0.24	46,46,46,46	0
59	MG	AA	3637	1/1	0.88	0.12	54,54,54,54	0
59	MG	AA	3974	1/1	0.88	0.12	65,65,65,65	0
59	MG	DA	3881	1/1	0.88	0.51	86,86,86,86	0
59	MG	CA	2034	1/1	0.88	0.24	61,61,61,61	0
59	MG	D8	103	1/1	0.88	0.21	76,76,76,76	0
59	MG	DA	3347	1/1	0.88	0.27	29,29,29,29	0
59	MG	DA	3890	1/1	0.88	0.47	86,86,86,86	0
59	MG	DA	3559	1/1	0.88	0.10	23,23,23,23	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	A1	204	1/1	0.88	0.85	65,65,65,65	0
59	MG	AA	3890	1/1	0.88	0.15	103,103,103,103	0
59	MG	AA	3636	1/1	0.88	0.09	48,48,48,48	0
59	MG	D8	104	1/1	0.88	0.55	50,50,50,50	0
59	MG	DA	3713	1/1	0.88	0.36	70,70,70,70	0
59	MG	CA	1756	1/1	0.88	0.15	53,53,53,53	0
59	MG	BA	2085	1/1	0.88	0.46	102,102,102,102	0
59	MG	DW	107	1/1	0.88	0.75	70,70,70,70	0
59	MG	CA	2188	1/1	0.88	0.08	79,79,79,79	0
59	MG	C1	106	1/1	0.88	0.35	89,89,89,89	0
59	MG	BA	2173	1/1	0.88	0.13	98,98,98,98	0
59	MG	BC	101	1/1	0.88	0.13	53,53,53,53	0
59	MG	DB	229	1/1	0.88	0.17	83,83,83,83	0
59	MG	BA	1786	1/1	0.88	0.39	57,57,57,57	0
59	MG	BA	1996	1/1	0.88	0.32	86,86,86,86	0
59	MG	DA	3627	1/1	0.88	0.20	40,40,40,40	0
59	MG	CA	1648	1/1	0.88	0.17	36,36,36,36	0
59	MG	AA	3478	1/1	0.88	0.15	59,59,59,59	0
59	MG	CA	1777	1/1	0.88	0.32	49,49,49,49	0
59	MG	DA	3188	1/1	0.88	0.26	60,60,60,60	0
59	MG	BA	2179	1/1	0.88	0.14	101,101,101,101	0
59	MG	DA	3376	1/1	0.88	0.12	18,18,18,18	0
59	MG	DA	3568	1/1	0.88	0.20	54,54,54,54	0
59	MG	AA	3703	1/1	0.88	0.08	51,51,51,51	0
59	MG	AA	3356	1/1	0.88	0.12	62,62,62,62	0
59	MG	AA	3041	1/1	0.88	0.10	81,81,81,81	0
59	MG	BD	110	1/1	0.88	0.10	81,81,81,81	0
59	MG	DA	3607	1/1	0.88	0.17	79,79,79,79	0
59	MG	CA	2228	1/1	0.88	0.25	75,75,75,75	0
59	MG	DO	206	1/1	0.88	0.32	51,51,51,51	0
59	MG	AA	3839	1/1	0.88	0.20	63,63,63,63	0
59	MG	DA	4048	1/1	0.88	0.10	75,75,75,75	0
59	MG	CA	2107	1/1	0.89	0.23	73,73,73,73	0
59	MG	DD	311	1/1	0.89	0.13	65,65,65,65	0
59	MG	BA	1812	1/1	0.89	0.27	136,136,136,136	0
59	MG	DA	4077	1/1	0.89	0.20	51,51,51,51	0
59	MG	BA	1963	1/1	0.89	0.13	83,83,83,83	0
59	MG	DA	4918	1/1	0.89	0.14	48,48,48,48	0
59	MG	CA	2037	1/1	0.89	0.32	75,75,75,75	0
59	MG	DA	3827	1/1	0.89	0.14	97,97,97,97	0
59	MG	DA	4105	1/1	0.89	0.17	62,62,62,62	0
59	MG	DA	4521	1/1	0.89	0.16	47,47,47,47	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	5040	1/1	0.89	0.31	90,90,90,90	0
59	MG	BA	1715	1/1	0.89	0.28	75,75,75,75	0
59	MG	DA	3692	1/1	0.89	0.07	60,60,60,60	0
59	MG	AA	4081	1/1	0.89	0.31	49,49,49,49	0
59	MG	DA	4099	1/1	0.89	0.10	55,55,55,55	0
59	MG	CA	2011	1/1	0.89	0.16	62,62,62,62	0
59	MG	CA	2292	1/1	0.89	0.11	67,67,67,67	0
59	MG	DA	4103	1/1	0.89	0.46	85,85,85,85	0
59	MG	DA	3604	1/1	0.89	0.13	57,57,57,57	0
59	MG	DB	235	1/1	0.89	0.22	65,65,65,65	0
59	MG	DA	3714	1/1	0.89	0.25	21,21,21,21	0
59	MG	DA	5020	1/1	0.89	0.16	63,63,63,63	0
59	MG	DA	3725	1/1	0.89	0.13	22,22,22,22	0
59	MG	DA	4359	1/1	0.89	0.26	47,47,47,47	0
59	MG	CC	116	1/1	0.89	0.10	66,66,66,66	0
59	MG	BD	120	1/1	0.89	0.46	90,90,90,90	0
59	MG	BA	1919	1/1	0.89	0.30	59,59,59,59	0
59	MG	DA	4634	1/1	0.89	0.44	82,82,82,82	0
59	MG	CK	210	1/1	0.89	0.19	71,71,71,71	0
59	MG	DS	203	1/1	0.89	0.28	54,54,54,54	0
59	MG	CA	1666	1/1	0.89	0.36	60,60,60,60	0
59	MG	DA	4590	1/1	0.89	0.55	81,81,81,81	0
59	MG	CA	1650	1/1	0.89	0.23	36,36,36,36	0
59	MG	DA	4898	1/1	0.89	0.25	58,58,58,58	0
59	MG	DA	3561	1/1	0.89	0.20	51,51,51,51	0
59	MG	BA	2233	1/1	0.89	0.14	100,100,100,100	0
59	MG	DA	4989	1/1	0.89	0.25	66,66,66,66	0
59	MG	AA	3771	1/1	0.89	0.37	79,79,79,79	0
59	MG	DA	5061	1/1	0.89	0.33	71,71,71,71	0
59	MG	DA	3862	1/1	0.89	0.16	71,71,71,71	0
59	MG	CA	2086	1/1	0.89	0.22	83,83,83,83	0
59	MG	AA	3680	1/1	0.89	0.34	71,71,71,71	0
59	MG	AA	4002	1/1	0.89	0.12	93,93,93,93	0
59	MG	DA	3324	1/1	0.89	0.15	42,42,42,42	0
59	MG	AA	3406	1/1	0.89	0.24	32,32,32,32	0
59	MG	CA	2242	1/1	0.89	0.16	84,84,84,84	0
59	MG	BA	1808	1/1	0.89	0.17	70,70,70,70	0
59	MG	DA	3822	1/1	0.89	0.30	55,55,55,55	0
59	MG	DB	222	1/1	0.89	0.30	43,43,43,43	0
59	MG	DA	3693	1/1	0.89	0.13	108,108,108,108	0
59	MG	DA	3673	1/1	0.89	0.12	61,61,61,61	0
59	MG	CE	301	1/1	0.89	0.25	61,61,61,61	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3172	1/1	0.89	0.28	37,37,37,37	0
59	MG	CA	2206	1/1	0.89	0.12	52,52,52,52	0
59	MG	BW	204	1/1	0.89	0.19	90,90,90,90	0
59	MG	AA	3159	1/1	0.89	0.19	61,61,61,61	0
59	MG	DA	3275	1/1	0.89	0.42	64,64,64,64	0
59	MG	DA	4679	1/1	0.89	0.18	47,47,47,47	0
59	MG	AA	3153	1/1	0.89	0.11	40,40,40,40	0
59	MG	DA	4957	1/1	0.89	0.07	86,86,86,86	0
59	MG	AA	3117	1/1	0.89	0.27	72,72,72,72	0
59	MG	DA	4906	1/1	0.89	0.26	96,96,96,96	0
59	MG	DA	4491	1/1	0.89	0.12	66,66,66,66	0
59	MG	DA	3181	1/1	0.89	0.25	40,40,40,40	0
59	MG	DK	201	1/1	0.89	0.24	71,71,71,71	0
59	MG	CA	1911	1/1	0.89	0.12	56,56,56,56	0
59	MG	DA	4811	1/1	0.89	0.23	97,97,97,97	0
59	MG	CA	2096	1/1	0.89	0.12	60,60,60,60	0
59	MG	DA	4315	1/1	0.89	0.10	67,67,67,67	0
59	MG	CA	2071	1/1	0.89	0.28	100,100,100,100	0
59	MG	DA	4854	1/1	0.89	0.23	66,66,66,66	0
59	MG	BA	2004	1/1	0.89	0.12	69,69,69,69	0
59	MG	CA	1995	1/1	0.89	0.13	48,48,48,48	0
59	MG	DA	4237	1/1	0.89	0.27	78,78,78,78	0
59	MG	CA	1610	1/1	0.89	0.26	48,48,48,48	0
59	MG	AA	3574	1/1	0.89	0.21	57,57,57,57	0
59	MG	DA	4508	1/1	0.89	0.33	58,58,58,58	0
59	MG	AA	3347	1/1	0.89	0.21	66,66,66,66	0
59	MG	AA	3642	1/1	0.89	0.24	56,56,56,56	0
59	MG	AA	3193	1/1	0.89	0.17	99,99,99,99	0
59	MG	AA	3379	1/1	0.89	0.09	70,70,70,70	0
59	MG	DA	4644	1/1	0.89	0.12	64,64,64,64	0
59	MG	CA	2135	1/1	0.89	0.07	92,92,92,92	0
59	MG	AV	301	1/1	0.89	0.11	67,67,67,67	0
59	MG	AA	3786	1/1	0.89	0.51	107,107,107,107	0
59	MG	DA	5070	1/1	0.89	0.17	51,51,51,51	0
59	MG	CV	101	1/1	0.89	0.09	53,53,53,53	0
59	MG	DA	3517	1/1	0.89	0.10	29,29,29,29	0
59	MG	DA	4804	1/1	0.89	0.31	69,69,69,69	0
59	MG	DA	3816	1/1	0.89	0.47	89,89,89,89	0
59	MG	BA	2088	1/1	0.89	0.33	98,98,98,98	0
59	MG	AA	3836	1/1	0.89	0.29	62,62,62,62	0
59	MG	CA	2073	1/1	0.89	0.35	41,41,41,41	0
59	MG	AA	3225	1/1	0.89	0.11	18,18,18,18	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	2061	1/1	0.89	0.21	50,50,50,50	0
59	MG	DA	4229	1/1	0.89	0.22	48,48,48,48	0
59	MG	CA	2000	1/1	0.89	0.10	99,99,99,99	0
59	MG	AA	3800	1/1	0.89	0.10	55,55,55,55	0
59	MG	DF	312	1/1	0.89	0.13	75,75,75,75	0
59	MG	AR	202	1/1	0.89	0.12	64,64,64,64	0
59	MG	CA	1783	1/1	0.89	0.14	56,56,56,56	0
59	MG	BA	1753	1/1	0.89	0.14	69,69,69,69	0
59	MG	DA	3799	1/1	0.89	0.25	83,83,83,83	0
59	MG	DA	3218	1/1	0.89	0.33	47,47,47,47	0
59	MG	BA	1986	1/1	0.89	0.09	78,78,78,78	0
59	MG	CA	2046	1/1	0.89	0.14	67,67,67,67	0
59	MG	DA	4026	1/1	0.89	0.21	55,55,55,55	0
59	MG	DA	3401	1/1	0.89	0.33	45,45,45,45	0
59	MG	AA	3660	1/1	0.89	0.29	68,68,68,68	0
59	MG	AN	201	1/1	0.89	0.34	118,118,118,118	0
59	MG	D2	208	1/1	0.89	0.36	71,71,71,71	0
59	MG	D4	101	1/1	0.89	0.36	96,96,96,96	0
59	MG	DA	3025	1/1	0.89	0.18	44,44,44,44	0
59	MG	AA	4115	1/1	0.89	0.28	73,73,73,73	0
59	MG	AA	3368	1/1	0.89	0.28	87,87,87,87	0
59	MG	CA	1612	1/1	0.89	0.17	57,57,57,57	0
59	MG	CA	2147	1/1	0.89	0.16	69,69,69,69	0
59	MG	CA	1873	1/1	0.89	0.08	51,51,51,51	0
59	MG	DA	4156	1/1	0.89	0.18	47,47,47,47	0
59	MG	DA	4945	1/1	0.89	0.25	78,78,78,78	0
59	MG	AA	4159	1/1	0.89	0.14	67,67,67,67	0
59	MG	DA	4332	1/1	0.89	0.13	53,53,53,53	0
59	MG	CA	1838	1/1	0.89	0.30	122,122,122,122	0
59	MG	DA	4325	1/1	0.89	0.42	45,45,45,45	0
59	MG	CA	1637	1/1	0.89	0.26	45,45,45,45	0
59	MG	DA	4187	1/1	0.89	0.12	46,46,46,46	0
59	MG	DA	3456	1/1	0.89	0.24	60,60,60,60	0
59	MG	AA	3572	1/1	0.89	0.15	64,64,64,64	0
59	MG	DA	5004	1/1	0.89	0.25	80,80,80,80	0
59	MG	DA	4645	1/1	0.89	0.21	59,59,59,59	0
59	MG	CA	2289	1/1	0.89	0.18	75,75,75,75	0
59	MG	AA	4063	1/1	0.89	0.16	75,75,75,75	0
59	MG	AA	3892	1/1	0.89	0.29	84,84,84,84	0
59	MG	CA	2199	1/1	0.89	0.10	106,106,106,106	0
59	MG	AA	3141	1/1	0.89	0.35	66,66,66,66	0
59	MG	DA	4636	1/1	0.89	0.20	64,64,64,64	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3005	1/1	0.89	0.20	17,17,17,17	0
59	MG	DA	3615	1/1	0.89	0.45	112,112,112,112	0
59	MG	DA	4599	1/1	0.89	0.17	50,50,50,50	0
59	MG	AA	3195	1/1	0.89	0.25	82,82,82,82	0
59	MG	CA	2064	1/1	0.89	0.25	66,66,66,66	0
59	MG	DA	4282	1/1	0.89	0.14	60,60,60,60	0
59	MG	DA	4449	1/1	0.89	0.20	64,64,64,64	0
59	MG	CK	204	1/1	0.89	0.09	60,60,60,60	0
59	MG	CA	1782	1/1	0.89	0.35	50,50,50,50	0
59	MG	D3	103	1/1	0.89	0.22	23,23,23,23	0
59	MG	AA	3220	1/1	0.89	0.17	70,70,70,70	0
59	MG	BA	1603	1/1	0.89	0.39	58,58,58,58	0
59	MG	DB	260	1/1	0.89	0.12	63,63,63,63	0
59	MG	AA	3252	1/1	0.89	0.19	38,38,38,38	0
59	MG	BA	1677	1/1	0.89	0.26	52,52,52,52	0
59	MG	DA	4900	1/1	0.89	0.45	76,76,76,76	0
59	MG	DA	4529	1/1	0.89	0.29	75,75,75,75	0
59	MG	DB	237	1/1	0.89	0.10	57,57,57,57	0
59	MG	DA	4484	1/1	0.89	0.35	50,50,50,50	0
59	MG	DA	4019	1/1	0.89	0.14	91,91,91,91	0
59	MG	BA	2241	1/1	0.89	0.22	78,78,78,78	0
59	MG	DA	4248	1/1	0.89	0.50	89,89,89,89	0
59	MG	AA	3791	1/1	0.89	0.25	65,65,65,65	0
59	MG	DA	3364	1/1	0.89	0.28	54,54,54,54	0
59	MG	DA	3095	1/1	0.89	0.29	27,27,27,27	0
59	MG	BA	2027	1/1	0.89	0.24	107,107,107,107	0
59	MG	DA	4646	1/1	0.89	0.21	69,69,69,69	0
59	MG	AA	3425	1/1	0.89	0.15	44,44,44,44	0
59	MG	DA	3405	1/1	0.89	0.22	43,43,43,43	0
59	MG	AA	3933	1/1	0.89	0.43	115,115,115,115	0
59	MG	BA	1974	1/1	0.89	0.28	109,109,109,109	0
59	MG	BA	1706	1/1	0.89	0.29	52,52,52,52	0
59	MG	DA	4273	1/1	0.89	0.28	74,74,74,74	0
59	MG	AA	3215	1/1	0.89	0.16	72,72,72,72	0
59	MG	CA	1776	1/1	0.89	0.41	62,62,62,62	0
59	MG	CA	2310	1/1	0.89	0.12	127,127,127,127	0
59	MG	DA	4262	1/1	0.89	0.24	47,47,47,47	0
59	MG	DA	4480	1/1	0.89	0.28	77,77,77,77	0
59	MG	AA	3651	1/1	0.89	0.17	41,41,41,41	0
59	MG	DA	4294	1/1	0.89	0.11	83,83,83,83	0
59	MG	BA	1737	1/1	0.89	0.27	59,59,59,59	0
59	MG	DA	4813	1/1	0.89	0.38	78,78,78,78	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	BA	2020	1/1	0.89	0.13	90,90,90,90	0
59	MG	DA	4394	1/1	0.89	0.44	69,69,69,69	0
59	MG	BA	2226	1/1	0.89	0.39	101,101,101,101	0
59	MG	CA	2018	1/1	0.89	0.33	106,106,106,106	0
59	MG	DA	3971	1/1	0.89	0.21	66,66,66,66	0
59	MG	CC	127	1/1	0.89	0.49	89,89,89,89	0
59	MG	CA	1778	1/1	0.89	0.39	69,69,69,69	0
59	MG	DA	3417	1/1	0.89	0.20	62,62,62,62	0
59	MG	DA	3954	1/1	0.89	0.44	52,52,52,52	0
59	MG	CC	114	1/1	0.89	0.12	73,73,73,73	0
59	MG	CA	2298	1/1	0.89	0.09	68,68,68,68	0
59	MG	AA	3683	1/1	0.89	0.16	46,46,46,46	0
59	MG	DA	4754	1/1	0.89	0.29	71,71,71,71	0
59	MG	DA	4266	1/1	0.89	0.13	64,64,64,64	0
59	MG	DA	4250	1/1	0.89	0.30	49,49,49,49	0
59	MG	DF	305	1/1	0.89	0.10	54,54,54,54	0
59	MG	DA	3281	1/1	0.89	0.21	38,38,38,38	0
59	MG	AA	3289	1/1	0.89	0.24	90,90,90,90	0
59	MG	DA	3446	1/1	0.89	0.24	66,66,66,66	0
59	MG	DA	3152	1/1	0.89	0.24	52,52,52,52	0
59	MG	AA	3784	1/1	0.89	0.21	64,64,64,64	0
59	MG	CA	2232	1/1	0.89	0.35	86,86,86,86	0
59	MG	DA	4622	1/1	0.89	0.37	54,54,54,54	0
59	MG	DA	3947	1/1	0.89	0.07	72,72,72,72	0
59	MG	DA	3259	1/1	0.89	0.34	60,60,60,60	0
59	MG	DA	4876	1/1	0.89	0.30	61,61,61,61	0
59	MG	CG	304	1/1	0.89	0.10	95,95,95,95	0
59	MG	DA	5069	1/1	0.89	0.14	66,66,66,66	0
59	MG	D8	105	1/1	0.89	0.35	44,44,44,44	0
59	MG	CA	2087	1/1	0.89	0.23	95,95,95,95	0
59	MG	BA	2041	1/1	0.89	0.44	91,91,91,91	0
59	MG	DA	4938	1/1	0.89	0.12	55,55,55,55	0
59	MG	CA	1708	1/1	0.89	0.19	58,58,58,58	0
59	MG	CA	2014	1/1	0.89	0.10	56,56,56,56	0
59	MG	DA	4732	1/1	0.89	0.11	71,71,71,71	0
59	MG	AA	3546	1/1	0.89	0.31	52,52,52,52	0
59	MG	CH	203	1/1	0.89	0.29	70,70,70,70	0
59	MG	AA	3068	1/1	0.89	0.10	40,40,40,40	0
59	MG	DA	4637	1/1	0.89	0.24	80,80,80,80	0
59	MG	BA	1626	1/1	0.89	0.30	68,68,68,68	0
59	MG	AF	302	1/1	0.89	0.18	69,69,69,69	0
59	MG	DA	3669	1/1	0.89	0.20	68,68,68,68	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DD	307	1/1	0.89	0.34	50,50,50,50	0
59	MG	DA	3372	1/1	0.89	0.18	54,54,54,54	0
59	MG	AA	3876	1/1	0.89	0.16	67,67,67,67	0
59	MG	CG	308	1/1	0.89	0.26	100,100,100,100	0
59	MG	AA	3808	1/1	0.89	0.17	75,75,75,75	0
59	MG	AA	4055	1/1	0.89	0.35	110,110,110,110	0
59	MG	AA	3955	1/1	0.89	0.30	66,66,66,66	0
59	MG	AA	4123	1/1	0.89	0.14	55,55,55,55	0
59	MG	AA	3982	1/1	0.89	0.11	71,71,71,71	0
59	MG	AU	205	1/1	0.90	0.18	69,69,69,69	0
59	MG	DA	5057	1/1	0.90	0.57	97,97,97,97	0
59	MG	DA	4880	1/1	0.90	0.20	110,110,110,110	0
59	MG	BA	1835	1/1	0.90	0.26	72,72,72,72	0
59	MG	AA	3056	1/1	0.90	0.13	63,63,63,63	0
59	MG	DA	3732	1/1	0.90	0.43	60,60,60,60	0
59	MG	DA	4525	1/1	0.90	0.21	108,108,108,108	0
59	MG	AK	201	1/1	0.90	0.22	64,64,64,64	0
59	MG	DA	4148	1/1	0.90	0.39	63,63,63,63	0
59	MG	DA	4326	1/1	0.90	0.28	56,56,56,56	0
59	MG	AA	3794	1/1	0.90	0.17	70,70,70,70	0
59	MG	CA	2145	1/1	0.90	0.22	80,80,80,80	0
59	MG	BA	2117	1/1	0.90	0.25	109,109,109,109	0
59	MG	AA	4078	1/1	0.90	0.11	82,82,82,82	0
59	MG	DA	5006	1/1	0.90	0.18	61,61,61,61	0
59	MG	DA	4482	1/1	0.90	0.41	80,80,80,80	0
59	MG	CA	1625	1/1	0.90	0.36	68,68,68,68	0
59	MG	AA	3748	1/1	0.90	0.09	84,84,84,84	0
59	MG	AA	3135	1/1	0.90	0.20	19,19,19,19	0
59	MG	AA	4101	1/1	0.90	0.27	99,99,99,99	0
59	MG	BA	1909	1/1	0.90	0.34	66,66,66,66	0
59	MG	CA	2265	1/1	0.90	0.09	64,64,64,64	0
59	MG	CA	2238	1/1	0.90	0.26	65,65,65,65	0
59	MG	CA	2133	1/1	0.90	0.12	67,67,67,67	0
59	MG	BA	1975	1/1	0.90	0.30	109,109,109,109	0
59	MG	BA	1922	1/1	0.90	0.11	71,71,71,71	0
59	MG	DA	4975	1/1	0.90	0.19	78,78,78,78	0
59	MG	AA	3918	1/1	0.90	0.12	90,90,90,90	0
59	MG	DA	3940	1/1	0.90	0.24	64,64,64,64	0
59	MG	DA	3294	1/1	0.90	0.16	28,28,28,28	0
59	MG	AA	4143	1/1	0.90	0.49	112,112,112,112	0
59	MG	CB	118	1/1	0.90	0.09	73,73,73,73	0
59	MG	AA	3862	1/1	0.90	0.21	75,75,75,75	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4766	1/1	0.90	0.21	67,67,67,67	0
59	MG	BD	118	1/1	0.90	0.24	107,107,107,107	0
59	MG	CA	2264	1/1	0.90	0.27	77,77,77,77	0
59	MG	DP	203	1/1	0.90	0.39	92,92,92,92	0
59	MG	DA	3684	1/1	0.90	0.42	56,56,56,56	0
59	MG	CA	1992	1/1	0.90	0.11	77,77,77,77	0
59	MG	BA	1958	1/1	0.90	0.08	80,80,80,80	0
59	MG	DA	3781	1/1	0.90	0.13	73,73,73,73	0
59	MG	AA	3067	1/1	0.90	0.16	37,37,37,37	0
59	MG	AA	3514	1/1	0.90	0.34	49,49,49,49	0
59	MG	BA	1979	1/1	0.90	0.22	73,73,73,73	0
59	MG	DA	4759	1/1	0.90	0.32	91,91,91,91	0
59	MG	AA	3604	1/1	0.90	0.09	74,74,74,74	0
59	MG	DA	4396	1/1	0.90	0.47	85,85,85,85	0
59	MG	DA	3542	1/1	0.90	0.25	51,51,51,51	0
59	MG	AB	203	1/1	0.90	0.20	37,37,37,37	0
59	MG	AA	3470	1/1	0.90	0.20	76,76,76,76	0
59	MG	DA	4849	1/1	0.90	0.25	55,55,55,55	0
59	MG	CP	202	1/1	0.90	0.16	63,63,63,63	0
59	MG	BA	2076	1/1	0.90	0.13	85,85,85,85	0
59	MG	DA	3352	1/1	0.90	0.44	80,80,80,80	0
59	MG	AA	4027	1/1	0.90	0.22	72,72,72,72	0
59	MG	AA	3239	1/1	0.90	0.07	51,51,51,51	0
59	MG	DZ	103	1/1	0.90	0.14	87,87,87,87	0
59	MG	BA	1980	1/1	0.90	0.09	102,102,102,102	0
59	MG	AA	3983	1/1	0.90	0.27	126,126,126,126	0
59	MG	DA	4737	1/1	0.90	0.22	85,85,85,85	0
59	MG	DA	3252	1/1	0.90	0.24	29,29,29,29	0
59	MG	BA	2192	1/1	0.90	0.17	102,102,102,102	0
59	MG	DA	3393	1/1	0.90	0.45	88,88,88,88	0
59	MG	DA	4503	1/1	0.90	0.28	96,96,96,96	0
59	MG	BA	2086	1/1	0.90	0.21	83,83,83,83	0
59	MG	BA	2202	1/1	0.90	0.41	87,87,87,87	0
59	MG	DA	3584	1/1	0.90	0.23	63,63,63,63	0
59	MG	DA	4038	1/1	0.90	0.54	103,103,103,103	0
59	MG	DA	4547	1/1	0.90	0.23	46,46,46,46	0
59	MG	DA	3374	1/1	0.90	0.17	84,84,84,84	0
59	MG	BA	1775	1/1	0.90	0.38	93,93,93,93	0
59	MG	AA	3913	1/1	0.90	0.14	38,38,38,38	0
59	MG	BA	1675	1/1	0.90	0.10	28,28,28,28	0
59	MG	DA	4862	1/1	0.90	0.12	66,66,66,66	0
59	MG	DA	4224	1/1	0.90	0.09	90,90,90,90	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4108	1/1	0.90	0.21	74,74,74,74	0
59	MG	DA	3123	1/1	0.90	0.42	65,65,65,65	0
59	MG	DA	3767	1/1	0.90	0.45	88,88,88,88	0
59	MG	DA	4692	1/1	0.90	0.33	74,74,74,74	0
59	MG	AA	3211	1/1	0.90	0.32	56,56,56,56	0
59	MG	BA	1738	1/1	0.90	0.28	59,59,59,59	0
59	MG	DA	4369	1/1	0.90	0.38	78,78,78,78	0
59	MG	AA	3549	1/1	0.90	0.17	44,44,44,44	0
59	MG	DA	3147	1/1	0.90	0.15	45,45,45,45	0
59	MG	BA	2012	1/1	0.90	0.41	72,72,72,72	0
59	MG	DA	3620	1/1	0.90	0.19	58,58,58,58	0
59	MG	CA	1863	1/1	0.90	0.17	66,66,66,66	0
59	MG	CA	2306	1/1	0.90	0.39	90,90,90,90	0
59	MG	AA	3145	1/1	0.90	0.25	61,61,61,61	0
59	MG	DA	4217	1/1	0.90	0.24	57,57,57,57	0
59	MG	DA	4052	1/1	0.90	0.16	178,178,178,178	0
59	MG	DA	4438	1/1	0.90	0.40	89,89,89,89	0
59	MG	AA	3679	1/1	0.90	0.42	70,70,70,70	0
59	MG	DA	3221	1/1	0.90	0.32	43,43,43,43	0
59	MG	AA	3386	1/1	0.90	0.11	51,51,51,51	0
59	MG	DA	4034	1/1	0.90	0.39	71,71,71,71	0
59	MG	DA	4365	1/1	0.90	0.31	61,61,61,61	0
59	MG	DA	4683	1/1	0.90	0.32	100,100,100,100	0
59	MG	DA	3448	1/1	0.90	0.28	40,40,40,40	0
59	MG	DA	3373	1/1	0.90	0.33	40,40,40,40	0
59	MG	BA	1778	1/1	0.90	0.37	49,49,49,49	0
59	MG	DA	4655	1/1	0.90	0.13	79,79,79,79	0
59	MG	CA	2070	1/1	0.90	0.10	68,68,68,68	0
59	MG	CA	1672	1/1	0.90	0.27	57,57,57,57	0
59	MG	CA	1661	1/1	0.90	0.31	59,59,59,59	0
59	MG	DA	4081	1/1	0.90	0.15	67,67,67,67	0
59	MG	DA	4830	1/1	0.90	0.30	64,64,64,64	0
59	MG	DA	4942	1/1	0.90	0.09	82,82,82,82	0
59	MG	D8	101	1/1	0.90	0.17	42,42,42,42	0
59	MG	DA	4510	1/1	0.90	0.38	78,78,78,78	0
59	MG	AA	3027	1/1	0.90	0.17	60,60,60,60	0
59	MG	DA	3368	1/1	0.90	0.19	31,31,31,31	0
59	MG	DA	4895	1/1	0.90	0.21	77,77,77,77	0
59	MG	DG	204	1/1	0.90	0.50	104,104,104,104	0
59	MG	DN	202	1/1	0.90	0.12	65,65,65,65	0
59	MG	CA	2148	1/1	0.90	0.13	69,69,69,69	0
59	MG	CA	2081	1/1	0.90	0.18	107,107,107,107	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4652	1/1	0.90	0.17	71,71,71,71	0
59	MG	DA	3270	1/1	0.90	0.25	90,90,90,90	0
59	MG	DA	4588	1/1	0.90	0.38	41,41,41,41	0
59	MG	BA	1750	1/1	0.90	0.29	65,65,65,65	0
59	MG	BA	1754	1/1	0.90	0.21	68,68,68,68	0
59	MG	DA	3594	1/1	0.90	0.09	49,49,49,49	0
59	MG	DA	3253	1/1	0.90	0.16	15,15,15,15	0
59	MG	AA	3856	1/1	0.90	0.32	86,86,86,86	0
59	MG	DA	3342	1/1	0.90	0.30	55,55,55,55	0
59	MG	AA	3436	1/1	0.90	0.24	127,127,127,127	0
59	MG	DA	3131	1/1	0.90	0.17	32,32,32,32	0
59	MG	AA	3525	1/1	0.90	0.13	36,36,36,36	0
59	MG	BA	2082	1/1	0.90	0.16	96,96,96,96	0
59	MG	BC	111	1/1	0.90	0.08	63,63,63,63	0
59	MG	DA	3173	1/1	0.90	0.18	36,36,36,36	0
59	MG	AA	3272	1/1	0.90	0.31	54,54,54,54	0
59	MG	DA	3050	1/1	0.90	0.11	46,46,46,46	0
59	MG	BG	305	1/1	0.90	0.23	70,70,70,70	0
59	MG	BA	2105	1/1	0.90	0.17	117,117,117,117	0
59	MG	DA	3480	1/1	0.90	0.18	46,46,46,46	0
59	MG	CA	2267	1/1	0.90	0.25	80,80,80,80	0
59	MG	CA	1771	1/1	0.90	0.21	44,44,44,44	0
59	MG	AA	3537	1/1	0.90	0.65	56,56,56,56	0
59	MG	BA	2116	1/1	0.90	0.06	54,54,54,54	0
59	MG	DA	3743	1/1	0.90	0.40	65,65,65,65	0
59	MG	CA	1857	1/1	0.90	0.27	105,105,105,105	0
59	MG	CA	2069	1/1	0.90	0.09	74,74,74,74	0
59	MG	DA	4374	1/1	0.90	0.49	77,77,77,77	0
59	MG	DQ	201	1/1	0.90	0.18	51,51,51,51	0
59	MG	DA	3143	1/1	0.90	0.37	61,61,61,61	0
59	MG	CA	1660	1/1	0.90	0.27	79,79,79,79	0
59	MG	BA	1645	1/1	0.90	0.30	88,88,88,88	0
59	MG	DA	4163	1/1	0.90	0.14	55,55,55,55	0
59	MG	DA	4245	1/1	0.90	0.18	73,73,73,73	0
59	MG	AA	3560	1/1	0.90	0.20	37,37,37,37	0
59	MG	BA	2129	1/1	0.90	0.08	75,75,75,75	0
59	MG	DA	5030	1/1	0.90	0.45	93,93,93,93	0
59	MG	BA	1924	1/1	0.90	0.45	78,78,78,78	0
59	MG	CA	2256	1/1	0.90	0.24	110,110,110,110	0
59	MG	CA	2134	1/1	0.90	0.22	89,89,89,89	0
59	MG	CA	1901	1/1	0.90	0.20	65,65,65,65	0
59	MG	DB	257	1/1	0.90	0.24	57,57,57,57	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4304	1/1	0.90	0.23	94,94,94,94	0
59	MG	AA	3103	1/1	0.90	0.17	45,45,45,45	0
59	MG	DA	4715	1/1	0.90	0.31	58,58,58,58	0
59	MG	DA	5025	1/1	0.90	0.22	100,100,100,100	0
59	MG	AA	3573	1/1	0.90	0.20	44,44,44,44	0
59	MG	AA	3278	1/1	0.90	0.28	59,59,59,59	0
59	MG	BA	1893	1/1	0.90	0.07	54,54,54,54	0
59	MG	BA	1950	1/1	0.90	0.23	62,62,62,62	0
59	MG	CA	1821	1/1	0.90	0.07	51,51,51,51	0
59	MG	AA	4146	1/1	0.90	0.22	90,90,90,90	0
59	MG	DA	5062	1/1	0.90	0.30	94,94,94,94	0
59	MG	AA	3893	1/1	0.90	0.15	67,67,67,67	0
59	MG	AA	3723	1/1	0.90	0.37	79,79,79,79	0
59	MG	BA	1633	1/1	0.90	0.20	45,45,45,45	0
59	MG	AA	3337	1/1	0.90	0.23	51,51,51,51	0
59	MG	AA	3788	1/1	0.90	0.21	87,87,87,87	0
59	MG	CA	2247	1/1	0.90	0.15	103,103,103,103	0
59	MG	AA	3578	1/1	0.90	0.15	33,33,33,33	0
59	MG	DA	5017	1/1	0.90	0.18	55,55,55,55	0
59	MG	BG	304	1/1	0.90	0.40	79,79,79,79	0
59	MG	BA	1973	1/1	0.90	0.09	81,81,81,81	0
59	MG	AA	3475	1/1	0.90	0.30	85,85,85,85	0
59	MG	DA	4792	1/1	0.90	0.21	113,113,113,113	0
59	MG	DA	4568	1/1	0.90	0.29	55,55,55,55	0
59	MG	AA	3950	1/1	0.90	0.06	63,63,63,63	0
59	MG	DA	3662	1/1	0.90	0.25	73,73,73,73	0
59	MG	BA	1813	1/1	0.90	0.10	94,94,94,94	0
59	MG	AB	201	1/1	0.90	0.14	42,42,42,42	0
59	MG	DA	4354	1/1	0.90	0.31	65,65,65,65	0
59	MG	DA	3455	1/1	0.90	0.33	58,58,58,58	0
59	MG	DD	308	1/1	0.90	0.68	53,53,53,53	0
59	MG	AA	3968	1/1	0.90	0.12	77,77,77,77	0
59	MG	CA	1726	1/1	0.90	0.18	48,48,48,48	0
59	MG	AU	201	1/1	0.90	0.20	51,51,51,51	0
59	MG	CC	120	1/1	0.90	0.28	44,44,44,44	0
59	MG	DA	4870	1/1	0.90	0.16	82,82,82,82	0
59	MG	A8	103	1/1	0.90	0.13	62,62,62,62	0
59	MG	DT	104	1/1	0.90	0.33	74,74,74,74	0
59	MG	CA	2020	1/1	0.90	0.17	67,67,67,67	0
59	MG	DT	105	1/1	0.90	0.32	84,84,84,84	0
59	MG	DA	4764	1/1	0.90	0.49	85,85,85,85	0
59	MG	DA	3603	1/1	0.90	0.29	20,20,20,20	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	5071	1/1	0.90	0.30	101,101,101,101	0
59	MG	DA	4964	1/1	0.90	0.20	73,73,73,73	0
59	MG	DA	3468	1/1	0.90	0.20	49,49,49,49	0
59	MG	D1	211	1/1	0.90	0.33	49,49,49,49	0
59	MG	DA	3531	1/1	0.90	0.23	53,53,53,53	0
59	MG	DA	3284	1/1	0.90	0.20	27,27,27,27	0
59	MG	DA	5001	1/1	0.90	0.21	75,75,75,75	0
59	MG	BA	2266	1/1	0.90	0.14	91,91,91,91	0
59	MG	AA	3769	1/1	0.90	0.09	80,80,80,80	0
59	MG	AA	3829	1/1	0.90	0.08	88,88,88,88	0
59	MG	BA	2174	1/1	0.90	0.40	125,125,125,125	0
59	MG	AA	3241	1/1	0.90	0.15	59,59,59,59	0
59	MG	DA	4117	1/1	0.90	0.29	58,58,58,58	0
59	MG	AA	3538	1/1	0.90	0.15	45,45,45,45	0
59	MG	DA	3290	1/1	0.90	0.22	33,33,33,33	0
59	MG	AA	4058	1/1	0.90	0.15	63,63,63,63	0
59	MG	AA	3044	1/1	0.90	0.32	36,36,36,36	0
59	MG	AF	303	1/1	0.90	0.12	68,68,68,68	0
59	MG	DA	4509	1/1	0.90	0.29	64,64,64,64	0
59	MG	CA	1842	1/1	0.90	0.13	78,78,78,78	0
59	MG	DA	4967	1/1	0.90	0.17	74,74,74,74	0
59	MG	AZ	103	1/1	0.90	0.13	75,75,75,75	0
59	MG	D3	106	1/1	0.90	0.36	58,58,58,58	0
59	MG	DA	3453	1/1	0.90	0.23	56,56,56,56	0
59	MG	DA	3387	1/1	0.90	0.13	47,47,47,47	0
59	MG	AA	3357	1/1	0.90	1.30	91,91,91,91	0
59	MG	BA	1745	1/1	0.90	0.25	38,38,38,38	0
59	MG	BA	2002	1/1	0.90	0.23	73,73,73,73	0
59	MG	AA	3409	1/1	0.90	0.21	56,56,56,56	0
59	MG	AA	3247	1/1	0.90	0.11	47,47,47,47	0
59	MG	DA	3875	1/1	0.90	0.33	116,116,116,116	0
59	MG	AB	204	1/1	0.90	0.07	54,54,54,54	0
59	MG	DB	258	1/1	0.90	0.38	77,77,77,77	0
59	MG	DA	3651	1/1	0.90	0.18	40,40,40,40	0
59	MG	AA	3949	1/1	0.90	0.06	71,71,71,71	0
59	MG	BA	2109	1/1	0.90	0.43	93,93,93,93	0
59	MG	DE	313	1/1	0.90	0.20	73,73,73,73	0
59	MG	AA	3106	1/1	0.90	0.18	17,17,17,17	0
59	MG	CA	1704	1/1	0.90	0.23	46,46,46,46	0
59	MG	AA	4069	1/1	0.90	0.21	80,80,80,80	0
59	MG	DA	4608	1/1	0.90	0.17	47,47,47,47	0
59	MG	DA	3390	1/1	0.90	0.17	83,83,83,83	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	BA	1955	1/1	0.90	0.30	74,74,74,74	0
59	MG	DA	4591	1/1	0.90	0.30	107,107,107,107	0
59	MG	DA	3149	1/1	0.90	0.19	18,18,18,18	0
59	MG	DA	3735	1/1	0.90	0.26	76,76,76,76	0
59	MG	CA	1974	1/1	0.90	0.08	55,55,55,55	0
59	MG	CA	2239	1/1	0.90	0.13	65,65,65,65	0
59	MG	AA	3172	1/1	0.90	0.21	67,67,67,67	0
59	MG	CA	2157	1/1	0.90	0.21	79,79,79,79	0
59	MG	DA	3229	1/1	0.90	0.10	33,33,33,33	0
59	MG	DA	3851	1/1	0.90	0.23	72,72,72,72	0
59	MG	CA	2053	1/1	0.90	0.14	74,74,74,74	0
59	MG	CA	1744	1/1	0.90	0.18	55,55,55,55	0
59	MG	BA	2133	1/1	0.90	0.15	122,122,122,122	0
59	MG	DA	4408	1/1	0.90	0.23	54,54,54,54	0
59	MG	DA	4446	1/1	0.90	0.11	54,54,54,54	0
59	MG	AA	3261	1/1	0.90	0.16	40,40,40,40	0
59	MG	CA	2279	1/1	0.90	0.22	67,67,67,67	0
59	MG	AA	3008	1/1	0.90	0.24	31,31,31,31	0
59	MG	BM	202	1/1	0.90	0.32	81,81,81,81	0
59	MG	CA	1833	1/1	0.90	0.06	57,57,57,57	0
59	MG	AA	3671	1/1	0.90	0.22	102,102,102,102	0
59	MG	DA	4164	1/1	0.90	0.13	29,29,29,29	0
59	MG	AA	3446	1/1	0.90	0.36	90,90,90,90	0
59	MG	DA	4884	1/1	0.90	0.16	73,73,73,73	0
59	MG	AA	4028	1/1	0.91	0.23	81,81,81,81	0
59	MG	DA	4738	1/1	0.91	0.10	55,55,55,55	0
59	MG	AA	3647	1/1	0.91	0.12	25,25,25,25	0
59	MG	CA	1965	1/1	0.91	0.06	81,81,81,81	0
59	MG	DA	3223	1/1	0.91	0.34	45,45,45,45	0
59	MG	DU	210	1/1	0.91	0.42	107,107,107,107	0
59	MG	AQ	201	1/1	0.91	0.18	56,56,56,56	0
59	MG	DA	3299	1/1	0.91	0.14	33,33,33,33	0
59	MG	CA	1909	1/1	0.91	0.33	69,69,69,69	0
59	MG	D5	108	1/1	0.91	0.34	63,63,63,63	0
59	MG	AD	302	1/1	0.91	0.17	71,71,71,71	0
59	MG	CA	1924	1/1	0.91	0.16	88,88,88,88	0
59	MG	BA	2035	1/1	0.91	0.15	91,91,91,91	0
59	MG	DA	4254	1/1	0.91	0.46	83,83,83,83	0
59	MG	DA	4039	1/1	0.91	0.36	75,75,75,75	0
59	MG	BA	2134	1/1	0.91	0.09	62,62,62,62	0
59	MG	DA	3960	1/1	0.91	0.17	54,54,54,54	0
59	MG	DA	3434	1/1	0.91	0.23	41,41,41,41	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	BA	2228	1/1	0.91	0.22	98,98,98,98	0
59	MG	AA	3929	1/1	0.91	0.07	104,104,104,104	0
59	MG	DA	3685	1/1	0.91	0.22	59,59,59,59	0
59	MG	DA	3340	1/1	0.91	0.24	37,37,37,37	0
59	MG	CA	1764	1/1	0.91	0.08	48,48,48,48	0
59	MG	DA	3543	1/1	0.91	0.28	62,62,62,62	0
59	MG	CA	1975	1/1	0.91	0.33	71,71,71,71	0
59	MG	DA	4330	1/1	0.91	0.14	56,56,56,56	0
59	MG	DA	4290	1/1	0.91	0.45	74,74,74,74	0
59	MG	BA	2168	1/1	0.91	0.08	73,73,73,73	0
59	MG	CA	1793	1/1	0.91	0.15	74,74,74,74	0
59	MG	DA	3463	1/1	0.91	0.39	61,61,61,61	0
59	MG	CA	1616	1/1	0.91	0.23	39,39,39,39	0
59	MG	AA	4061	1/1	0.91	0.27	79,79,79,79	0
59	MG	A1	205	1/1	0.91	0.26	79,79,79,79	0
59	MG	AA	4098	1/1	0.91	0.48	83,83,83,83	0
59	MG	BA	1863	1/1	0.91	0.05	81,81,81,81	0
59	MG	BT	202	1/1	0.91	0.10	76,76,76,76	0
59	MG	AA	3503	1/1	0.91	0.17	23,23,23,23	0
59	MG	AA	3221	1/1	0.91	0.33	63,63,63,63	0
59	MG	AA	3663	1/1	0.91	0.24	61,61,61,61	0
59	MG	CA	2140	1/1	0.91	0.07	110,110,110,110	0
59	MG	BA	1999	1/1	0.91	0.19	77,77,77,77	0
59	MG	AA	3559	1/1	0.91	0.19	50,50,50,50	0
59	MG	BA	2049	1/1	0.91	0.24	55,55,55,55	0
59	MG	AA	4166	1/1	0.91	0.16	142,142,142,142	0
59	MG	AA	3074	1/1	0.91	0.23	51,51,51,51	0
59	MG	AA	3431	1/1	0.91	0.14	53,53,53,53	0
59	MG	BB	113	1/1	0.91	0.16	142,142,142,142	0
59	MG	DA	3464	1/1	0.91	0.29	53,53,53,53	0
59	MG	CM	201	1/1	0.91	0.22	74,74,74,74	0
59	MG	DA	4268	1/1	0.91	0.18	58,58,58,58	0
59	MG	AA	3471	1/1	0.91	0.17	77,77,77,77	0
59	MG	BA	1671	1/1	0.91	0.25	50,50,50,50	0
59	MG	DA	3166	1/1	0.91	0.29	56,56,56,56	0
59	MG	BA	1897	1/1	0.91	0.21	65,65,65,65	0
59	MG	DA	4319	1/1	0.91	0.23	67,67,67,67	0
59	MG	DA	5072	1/1	0.91	0.15	58,58,58,58	0
59	MG	DA	4558	1/1	0.91	0.26	86,86,86,86	0
59	MG	DA	4193	1/1	0.91	0.59	78,78,78,78	0
59	MG	D2	204	1/1	0.91	0.47	51,51,51,51	0
59	MG	DA	3187	1/1	0.91	0.23	45,45,45,45	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AB	236	1/1	0.91	0.22	64,64,64,64	0
59	MG	DA	3995	1/1	0.91	0.29	71,71,71,71	0
59	MG	DA	4355	1/1	0.91	0.50	79,79,79,79	0
59	MG	DA	3519	1/1	0.91	0.16	36,36,36,36	0
59	MG	DA	3811	1/1	0.91	0.15	101,101,101,101	0
59	MG	CA	1971	1/1	0.91	0.16	54,54,54,54	0
59	MG	DA	5008	1/1	0.91	0.25	57,57,57,57	0
59	MG	AA	3603	1/1	0.91	0.08	80,80,80,80	0
59	MG	DD	306	1/1	0.91	0.17	51,51,51,51	0
59	MG	BA	1623	1/1	0.91	0.09	53,53,53,53	0
59	MG	DA	4088	1/1	0.91	0.33	83,83,83,83	0
59	MG	DF	313	1/1	0.91	0.16	68,68,68,68	0
59	MG	BD	121	1/1	0.91	0.27	79,79,79,79	0
59	MG	AA	3248	1/1	0.91	0.33	78,78,78,78	0
59	MG	DA	4635	1/1	0.91	0.22	71,71,71,71	0
59	MG	AA	3597	1/1	0.91	0.24	63,63,63,63	0
59	MG	BA	1607	1/1	0.91	0.25	74,74,74,74	0
59	MG	DB	275	1/1	0.91	0.15	50,50,50,50	0
59	MG	BA	2073	1/1	0.91	0.08	81,81,81,81	0
59	MG	CA	2074	1/1	0.91	0.07	61,61,61,61	0
59	MG	DA	4888	1/1	0.91	0.44	76,76,76,76	0
59	MG	BA	1849	1/1	0.91	0.14	74,74,74,74	0
59	MG	DA	3246	1/1	0.91	0.16	34,34,34,34	0
59	MG	DA	3516	1/1	0.91	0.21	53,53,53,53	0
59	MG	AD	307	1/1	0.91	0.19	51,51,51,51	0
59	MG	DA	3755	1/1	0.91	0.09	70,70,70,70	0
59	MG	DA	4377	1/1	0.91	0.17	46,46,46,46	0
59	MG	CA	1718	1/1	0.91	0.30	60,60,60,60	0
59	MG	CA	1682	1/1	0.91	0.26	36,36,36,36	0
59	MG	AA	4037	1/1	0.91	0.11	43,43,43,43	0
59	MG	BA	1910	1/1	0.91	0.09	59,59,59,59	0
59	MG	AA	3427	1/1	0.91	0.07	47,47,47,47	0
59	MG	DA	3760	1/1	0.91	0.25	78,78,78,78	0
59	MG	BA	2143	1/1	0.91	0.12	138,138,138,138	0
59	MG	DF	315	1/1	0.91	0.24	76,76,76,76	0
59	MG	DA	3896	1/1	0.91	0.14	69,69,69,69	0
59	MG	DA	4300	1/1	0.91	0.15	45,45,45,45	0
59	MG	BA	1878	1/1	0.91	0.14	75,75,75,75	0
59	MG	DA	4005	1/1	0.91	0.38	84,84,84,84	0
59	MG	DA	3860	1/1	0.91	0.31	55,55,55,55	0
59	MG	AA	3644	1/1	0.91	0.10	56,56,56,56	0
59	MG	AA	3532	1/1	0.91	0.31	87,87,87,87	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3268	1/1	0.91	0.31	60,60,60,60	0
59	MG	DA	3291	1/1	0.91	0.35	59,59,59,59	0
59	MG	DA	3458	1/1	0.91	0.38	54,54,54,54	0
59	MG	CD	116	1/1	0.91	0.15	78,78,78,78	0
59	MG	BA	2006	1/1	0.91	0.23	71,71,71,71	0
59	MG	AA	3178	1/1	0.91	0.23	41,41,41,41	0
59	MG	CA	1877	1/1	0.91	0.13	100,100,100,100	0
59	MG	BA	1935	1/1	0.91	0.22	95,95,95,95	0
59	MG	DA	4616	1/1	0.91	0.38	72,72,72,72	0
59	MG	AA	3313	1/1	0.91	0.18	83,83,83,83	0
59	MG	BA	2211	1/1	0.91	0.24	78,78,78,78	0
59	MG	DD	302	1/1	0.91	0.35	58,58,58,58	0
59	MG	AA	4067	1/1	0.91	0.19	86,86,86,86	0
59	MG	DB	236	1/1	0.91	0.15	49,49,49,49	0
59	MG	AA	3777	1/1	0.91	0.11	58,58,58,58	0
59	MG	CA	2158	1/1	0.91	0.14	58,58,58,58	0
59	MG	DB	224	1/1	0.91	0.24	51,51,51,51	0
59	MG	DA	4897	1/1	0.91	0.22	76,76,76,76	0
59	MG	BA	1726	1/1	0.91	0.25	62,62,62,62	0
59	MG	DE	310	1/1	0.91	0.27	52,52,52,52	0
59	MG	DA	4059	1/1	0.91	0.16	103,103,103,103	0
59	MG	AA	3183	1/1	0.91	0.21	29,29,29,29	0
59	MG	DA	4537	1/1	0.91	0.31	58,58,58,58	0
59	MG	DA	4247	1/1	0.91	0.17	48,48,48,48	0
59	MG	DA	4893	1/1	0.91	0.21	99,99,99,99	0
59	MG	DA	4426	1/1	0.91	0.49	75,75,75,75	0
59	MG	AA	3969	1/1	0.91	0.15	60,60,60,60	0
59	MG	BA	1654	1/1	0.91	0.29	53,53,53,53	0
59	MG	AQ	203	1/1	0.91	0.28	67,67,67,67	0
59	MG	BA	1911	1/1	0.91	0.41	76,76,76,76	0
59	MG	AA	3802	1/1	0.91	0.26	78,78,78,78	0
59	MG	CA	2031	1/1	0.91	0.22	77,77,77,77	0
59	MG	DA	3466	1/1	0.91	0.20	79,79,79,79	0
59	MG	AA	4120	1/1	0.91	0.24	87,87,87,87	0
59	MG	DA	4046	1/1	0.91	0.10	52,52,52,52	0
59	MG	DA	4541	1/1	0.91	0.30	58,58,58,58	0
59	MG	DA	3111	1/1	0.91	0.38	64,64,64,64	0
59	MG	DA	3580	1/1	0.91	0.31	58,58,58,58	0
59	MG	DA	3148	1/1	0.91	0.46	67,67,67,67	0
59	MG	DA	5055	1/1	0.91	0.27	69,69,69,69	0
59	MG	DA	3254	1/1	0.91	0.26	25,25,25,25	0
59	MG	DA	4472	1/1	0.91	0.21	56,56,56,56	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	1659	1/1	0.91	0.15	41,41,41,41	0
59	MG	AQ	202	1/1	0.91	0.12	94,94,94,94	0
59	MG	AA	4026	1/1	0.91	0.20	53,53,53,53	0
59	MG	AA	3758	1/1	0.91	0.20	65,65,65,65	0
59	MG	DA	4878	1/1	0.91	0.29	50,50,50,50	0
59	MG	AA	3639	1/1	0.91	0.30	55,55,55,55	0
59	MG	CA	2171	1/1	0.91	0.14	115,115,115,115	0
59	MG	AA	3625	1/1	0.91	0.20	64,64,64,64	0
59	MG	AA	3081	1/1	0.91	0.14	61,61,61,61	0
59	MG	DA	4615	1/1	0.91	0.29	123,123,123,123	0
59	MG	DA	3986	1/1	0.91	0.19	51,51,51,51	0
59	MG	BA	1648	1/1	0.91	0.26	50,50,50,50	0
59	MG	BA	1799	1/1	0.91	0.12	55,55,55,55	0
59	MG	CA	2102	1/1	0.91	0.17	57,57,57,57	0
59	MG	CA	1798	1/1	0.91	0.19	139,139,139,139	0
59	MG	CA	1710	1/1	0.91	0.17	30,30,30,30	0
59	MG	BA	2222	1/1	0.91	0.08	102,102,102,102	0
59	MG	DA	4922	1/1	0.91	0.32	99,99,99,99	0
59	MG	BA	2249	1/1	0.91	0.18	66,66,66,66	0
59	MG	BA	1690	1/1	0.91	0.07	66,66,66,66	0
59	MG	CA	1882	1/1	0.91	0.27	55,55,55,55	0
59	MG	DA	3539	1/1	0.91	0.24	93,93,93,93	0
59	MG	DA	4424	1/1	0.91	0.13	46,46,46,46	0
59	MG	AA	3050	1/1	0.91	0.20	59,59,59,59	0
59	MG	DA	3094	1/1	0.91	0.29	23,23,23,23	0
59	MG	BA	2235	1/1	0.91	0.09	69,69,69,69	0
59	MG	CA	2111	1/1	0.91	0.06	75,75,75,75	0
59	MG	AA	3185	1/1	0.91	0.20	29,29,29,29	0
59	MG	BA	2199	1/1	0.91	0.17	107,107,107,107	0
59	MG	BA	2196	1/1	0.91	0.20	65,65,65,65	0
59	MG	DA	4962	1/1	0.91	0.25	77,77,77,77	0
59	MG	AA	3857	1/1	0.91	0.42	65,65,65,65	0
59	MG	DM	201	1/1	0.91	0.20	29,29,29,29	0
59	MG	BA	2058	1/1	0.91	0.15	104,104,104,104	0
59	MG	BB	101	1/1	0.91	0.11	76,76,76,76	0
59	MG	DA	3850	1/1	0.91	0.26	51,51,51,51	0
59	MG	AA	3262	1/1	0.91	0.20	76,76,76,76	0
59	MG	AA	4136	1/1	0.91	0.27	71,71,71,71	0
59	MG	AA	3801	1/1	0.91	0.09	90,90,90,90	0
59	MG	DA	3536	1/1	0.91	0.32	48,48,48,48	0
59	MG	DA	4427	1/1	0.91	0.15	53,53,53,53	0
59	MG	CA	1807	1/1	0.91	0.18	63,63,63,63	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	1635	1/1	0.91	0.30	21,21,21,21	0
59	MG	BA	2023	1/1	0.91	0.20	67,67,67,67	0
59	MG	BK	206	1/1	0.91	0.22	108,108,108,108	0
59	MG	DA	5039	1/1	0.91	0.28	75,75,75,75	0
59	MG	AA	3676	1/1	0.91	0.46	81,81,81,81	0
59	MG	DA	4751	1/1	0.91	0.20	121,121,121,121	0
59	MG	DA	3186	1/1	0.91	0.16	39,39,39,39	0
59	MG	DM	203	1/1	0.91	0.14	59,59,59,59	0
60	ZN	D4	103	1/1	0.91	0.36	200,200,200,200	0
59	MG	DA	5049	1/1	0.91	0.29	80,80,80,80	0
59	MG	DA	4142	1/1	0.91	0.19	29,29,29,29	0
59	MG	AA	3208	1/1	0.91	0.17	30,30,30,30	0
59	MG	DA	4812	1/1	0.91	0.18	87,87,87,87	0
59	MG	DA	4261	1/1	0.91	0.20	65,65,65,65	0
59	MG	DA	4889	1/1	0.91	0.38	75,75,75,75	0
59	MG	DF	325	1/1	0.91	0.14	47,47,47,47	0
59	MG	DS	205	1/1	0.91	0.17	58,58,58,58	0
59	MG	AA	3674	1/1	0.91	0.24	61,61,61,61	0
59	MG	BA	1877	1/1	0.91	0.15	75,75,75,75	0
59	MG	BA	2126	1/1	0.91	0.15	74,74,74,74	0
59	MG	DM	204	1/1	0.91	1.30	67,67,67,67	0
59	MG	DA	4908	1/1	0.91	0.15	64,64,64,64	0
59	MG	DA	3341	1/1	0.91	0.39	49,49,49,49	0
59	MG	CA	1716	1/1	0.91	0.31	66,66,66,66	0
59	MG	AA	4152	1/1	0.91	0.21	77,77,77,77	0
59	MG	BA	1874	1/1	0.91	0.10	93,93,93,93	0
59	MG	AA	3179	1/1	0.91	0.16	38,38,38,38	0
59	MG	AA	3084	1/1	0.91	0.11	55,55,55,55	0
59	MG	AB	202	1/1	0.91	0.25	51,51,51,51	0
59	MG	DA	4818	1/1	0.91	0.28	80,80,80,80	0
59	MG	DA	5010	1/1	0.91	0.32	68,68,68,68	0
59	MG	AA	3682	1/1	0.91	0.13	75,75,75,75	0
59	MG	AA	3394	1/1	0.91	0.16	45,45,45,45	0
59	MG	D1	205	1/1	0.91	0.32	55,55,55,55	0
59	MG	CA	2024	1/1	0.91	0.05	96,96,96,96	0
59	MG	DA	4221	1/1	0.91	0.19	49,49,49,49	0
59	MG	CA	2227	1/1	0.91	0.45	78,78,78,78	0
59	MG	DA	3135	1/1	0.91	0.14	35,35,35,35	0
59	MG	DA	3521	1/1	0.91	0.31	61,61,61,61	0
59	MG	AW	102	1/1	0.91	0.38	56,56,56,56	0
59	MG	DF	304	1/1	0.91	0.17	29,29,29,29	0
59	MG	DA	4501	1/1	0.91	0.26	54,54,54,54	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	4162	1/1	0.91	0.29	94,94,94,94	0
59	MG	DA	4098	1/1	0.91	0.13	43,43,43,43	0
59	MG	D5	102	1/1	0.91	0.15	24,24,24,24	0
59	MG	AA	3510	1/1	0.91	0.12	26,26,26,26	0
59	MG	DA	3454	1/1	0.91	0.24	51,51,51,51	0
59	MG	AA	3776	1/1	0.91	0.14	84,84,84,84	0
59	MG	DA	4894	1/1	0.91	0.21	63,63,63,63	0
59	MG	BA	1660	1/1	0.91	0.20	101,101,101,101	0
59	MG	BA	2245	1/1	0.91	0.28	71,71,71,71	0
59	MG	D5	105	1/1	0.91	0.13	63,63,63,63	0
59	MG	DA	3711	1/1	0.91	0.25	85,85,85,85	0
59	MG	AO	203	1/1	0.91	0.11	47,47,47,47	0
59	MG	DA	4680	1/1	0.91	0.08	71,71,71,71	0
59	MG	CA	1943	1/1	0.91	0.34	73,73,73,73	0
59	MG	DA	4466	1/1	0.91	0.25	73,73,73,73	0
59	MG	AA	3991	1/1	0.91	0.22	73,73,73,73	0
59	MG	DA	3552	1/1	0.91	0.19	45,45,45,45	0
59	MG	AA	3844	1/1	0.91	0.33	94,94,94,94	0
59	MG	DA	4856	1/1	0.91	0.17	49,49,49,49	0
59	MG	DA	4089	1/1	0.91	0.13	63,63,63,63	0
59	MG	BA	1947	1/1	0.91	0.15	53,53,53,53	0
59	MG	CA	2006	1/1	0.91	0.14	67,67,67,67	0
59	MG	DA	5059	1/1	0.91	0.19	66,66,66,66	0
59	MG	BA	2180	1/1	0.91	0.33	88,88,88,88	0
59	MG	DN	201	1/1	0.91	0.27	50,50,50,50	0
59	MG	AA	3722	1/1	0.91	0.12	50,50,50,50	0
59	MG	DA	4314	1/1	0.91	0.26	67,67,67,67	0
59	MG	DA	4450	1/1	0.91	0.24	68,68,68,68	0
59	MG	DA	4914	1/1	0.91	0.21	44,44,44,44	0
59	MG	AA	4048	1/1	0.91	0.14	93,93,93,93	0
59	MG	DA	3083	1/1	0.91	0.22	9,9,9,9	0
59	MG	DA	3918	1/1	0.91	0.32	50,50,50,50	0
59	MG	DA	4675	1/1	0.91	0.28	73,73,73,73	0
59	MG	AA	3249	1/1	0.91	0.22	41,41,41,41	0
59	MG	AA	3866	1/1	0.91	0.32	96,96,96,96	0
59	MG	CA	2146	1/1	0.91	0.22	85,85,85,85	0
59	MG	DA	3821	1/1	0.91	0.20	81,81,81,81	0
59	MG	AA	3345	1/1	0.91	0.23	55,55,55,55	0
59	MG	DA	4351	1/1	0.91	0.17	43,43,43,43	0
59	MG	DO	207	1/1	0.91	0.12	37,37,37,37	0
59	MG	C1	101	1/1	0.91	0.43	156,156,156,156	0
59	MG	AA	3322	1/1	0.91	0.17	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4848	1/1	0.91	0.21	67,67,67,67	0
59	MG	DA	4662	1/1	0.91	0.33	101,101,101,101	0
59	MG	CA	1964	1/1	0.91	0.14	41,41,41,41	0
59	MG	AA	3282	1/1	0.91	0.08	64,64,64,64	0
59	MG	DA	4174	1/1	0.91	0.20	49,49,49,49	0
59	MG	CA	2151	1/1	0.91	0.14	60,60,60,60	0
59	MG	BA	1749	1/1	0.91	0.08	30,30,30,30	0
59	MG	DA	4036	1/1	0.91	0.19	55,55,55,55	0
59	MG	DA	3976	1/1	0.91	0.62	37,37,37,37	0
59	MG	AA	3143	1/1	0.91	0.11	37,37,37,37	0
59	MG	CA	1722	1/1	0.91	0.18	50,50,50,50	0
59	MG	DA	4092	1/1	0.91	0.20	99,99,99,99	0
59	MG	DR	201	1/1	0.91	0.14	72,72,72,72	0
59	MG	AS	202	1/1	0.91	0.11	63,63,63,63	0
59	MG	A6	101	1/1	0.91	0.07	55,55,55,55	0
59	MG	AA	3643	1/1	0.91	0.26	61,61,61,61	0
59	MG	AS	201	1/1	0.91	0.54	67,67,67,67	0
59	MG	BA	2203	1/1	0.91	0.19	118,118,118,118	0
59	MG	DA	3819	1/1	0.91	0.16	68,68,68,68	0
59	MG	DP	202	1/1	0.92	0.23	53,53,53,53	0
59	MG	AA	3199	1/1	0.92	0.34	75,75,75,75	0
59	MG	DF	319	1/1	0.92	0.15	67,67,67,67	0
59	MG	DW	103	1/1	0.92	0.21	78,78,78,78	0
59	MG	DA	3213	1/1	0.92	0.39	53,53,53,53	0
59	MG	DB	203	1/1	0.92	0.34	46,46,46,46	0
59	MG	DA	4832	1/1	0.92	0.17	86,86,86,86	0
59	MG	CA	1753	1/1	0.92	0.40	56,56,56,56	0
59	MG	AA	3695	1/1	0.92	0.15	49,49,49,49	0
59	MG	BC	102	1/1	0.92	0.10	78,78,78,78	0
59	MG	DA	4511	1/1	0.92	0.15	90,90,90,90	0
59	MG	CA	1785	1/1	0.92	0.29	63,63,63,63	0
59	MG	AA	3962	1/1	0.92	0.31	57,57,57,57	0
59	MG	BA	1942	1/1	0.92	0.10	63,63,63,63	0
59	MG	DA	3869	1/1	0.92	0.25	83,83,83,83	0
59	MG	DA	3232	1/1	0.92	0.26	61,61,61,61	0
59	MG	AO	202	1/1	0.92	0.13	55,55,55,55	0
59	MG	BA	1856	1/1	0.92	0.11	60,60,60,60	0
59	MG	DA	4188	1/1	0.92	0.10	39,39,39,39	0
59	MG	DA	4515	1/1	0.92	0.20	60,60,60,60	0
59	MG	CA	1688	1/1	0.92	0.18	42,42,42,42	0
59	MG	AA	3742	1/1	0.92	0.08	70,70,70,70	0
59	MG	BA	1809	1/1	0.92	0.14	114,114,114,114	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3826	1/1	0.92	0.20	92,92,92,92	0
59	MG	AA	3846	1/1	0.92	0.39	88,88,88,88	0
59	MG	DA	3019	1/1	0.92	0.34	20,20,20,20	0
59	MG	DU	214	1/1	0.92	0.27	62,62,62,62	0
59	MG	CA	2128	1/1	0.92	0.39	101,101,101,101	0
59	MG	DA	3475	1/1	0.92	0.23	51,51,51,51	0
59	MG	DA	4118	1/1	0.92	0.14	96,96,96,96	0
59	MG	DA	3158	1/1	0.92	0.10	28,28,28,28	0
59	MG	DA	4682	1/1	0.92	0.31	63,63,63,63	0
59	MG	AA	3467	1/1	0.92	0.29	55,55,55,55	0
59	MG	CA	2079	1/1	0.92	0.24	83,83,83,83	0
59	MG	DF	314	1/1	0.92	0.22	58,58,58,58	0
59	MG	DA	3566	1/1	0.92	0.10	21,21,21,21	0
59	MG	DA	4819	1/1	0.92	0.26	64,64,64,64	0
59	MG	DA	4623	1/1	0.92	0.17	47,47,47,47	0
59	MG	DA	4824	1/1	0.92	0.23	71,71,71,71	0
59	MG	DA	3768	1/1	0.92	0.15	77,77,77,77	0
59	MG	DA	4569	1/1	0.92	0.17	37,37,37,37	0
59	MG	DA	4176	1/1	0.92	0.14	53,53,53,53	0
59	MG	DA	4992	1/1	0.92	0.26	88,88,88,88	0
59	MG	CA	1830	1/1	0.92	0.04	59,59,59,59	0
59	MG	DA	4554	1/1	0.92	0.34	81,81,81,81	0
59	MG	CH	206	1/1	0.92	0.18	70,70,70,70	0
59	MG	D5	104	1/1	0.92	0.12	38,38,38,38	0
59	MG	BT	201	1/1	0.92	0.45	110,110,110,110	0
59	MG	DA	3421	1/1	0.92	0.21	56,56,56,56	0
59	MG	BA	2156	1/1	0.92	0.11	90,90,90,90	0
59	MG	DA	3637	1/1	0.92	0.28	63,63,63,63	0
59	MG	CK	211	1/1	0.92	0.17	57,57,57,57	0
59	MG	CA	1651	1/1	0.92	0.18	33,33,33,33	0
59	MG	DA	3544	1/1	0.92	0.18	51,51,51,51	0
59	MG	DA	3823	1/1	0.92	0.10	81,81,81,81	0
59	MG	DA	4589	1/1	0.92	0.35	58,58,58,58	0
59	MG	CA	2257	1/1	0.92	0.26	60,60,60,60	0
59	MG	AA	3961	1/1	0.92	0.29	54,54,54,54	0
59	MG	BA	1701	1/1	0.92	0.13	59,59,59,59	0
59	MG	DA	3074	1/1	0.92	0.22	22,22,22,22	0
59	MG	AB	222	1/1	0.92	0.12	85,85,85,85	0
59	MG	DA	3450	1/1	0.92	0.45	75,75,75,75	0
59	MG	BC	107	1/1	0.92	0.31	65,65,65,65	0
59	MG	CA	2104	1/1	0.92	0.07	67,67,67,67	0
59	MG	DA	4337	1/1	0.92	0.21	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4926	1/1	0.92	0.10	46,46,46,46	0
59	MG	BA	1674	1/1	0.92	0.11	45,45,45,45	0
59	MG	AA	3283	1/1	0.92	0.08	74,74,74,74	0
59	MG	DA	4667	1/1	0.92	0.20	46,46,46,46	0
59	MG	CA	1615	1/1	0.92	0.19	60,60,60,60	0
59	MG	AA	3919	1/1	0.92	0.08	59,59,59,59	0
59	MG	DA	3958	1/1	0.92	0.11	55,55,55,55	0
59	MG	AA	3481	1/1	0.92	0.17	61,61,61,61	0
59	MG	DA	4233	1/1	0.92	0.36	69,69,69,69	0
59	MG	DA	3686	1/1	0.92	0.19	70,70,70,70	0
59	MG	DA	3303	1/1	0.92	0.41	46,46,46,46	0
59	MG	AA	3105	1/1	0.92	0.21	50,50,50,50	0
59	MG	CN	202	1/1	0.92	0.20	64,64,64,64	0
59	MG	AA	4138	1/1	0.92	0.33	57,57,57,57	0
59	MG	DA	4758	1/1	0.92	0.33	71,71,71,71	0
59	MG	DA	4777	1/1	0.92	0.24	96,96,96,96	0
59	MG	AA	3125	1/1	0.92	0.22	51,51,51,51	0
59	MG	DA	3501	1/1	0.92	0.25	51,51,51,51	0
59	MG	AA	3137	1/1	0.92	0.23	25,25,25,25	0
59	MG	CA	2280	1/1	0.92	0.07	84,84,84,84	0
59	MG	CA	1977	1/1	0.92	0.14	64,64,64,64	0
59	MG	DA	3993	1/1	0.92	0.21	51,51,51,51	0
59	MG	CM	202	1/1	0.92	0.23	86,86,86,86	0
59	MG	DA	3746	1/1	0.92	0.18	36,36,36,36	0
59	MG	DA	3298	1/1	0.92	0.17	97,97,97,97	0
59	MG	AA	3941	1/1	0.92	0.16	57,57,57,57	0
59	MG	CA	1879	1/1	0.92	0.43	58,58,58,58	0
59	MG	AA	3823	1/1	0.92	0.17	89,89,89,89	0
59	MG	CA	2012	1/1	0.92	0.25	69,69,69,69	0
59	MG	CA	1731	1/1	0.92	0.14	56,56,56,56	0
59	MG	AA	3588	1/1	0.92	0.17	82,82,82,82	0
59	MG	DF	302	1/1	0.92	0.27	46,46,46,46	0
59	MG	DA	4166	1/1	0.92	0.32	41,41,41,41	0
59	MG	DA	3630	1/1	0.92	0.27	71,71,71,71	0
59	MG	DA	3327	1/1	0.92	0.17	48,48,48,48	0
59	MG	AA	3314	1/1	0.92	0.12	74,74,74,74	0
59	MG	BA	1833	1/1	0.92	0.22	71,71,71,71	0
59	MG	DA	4008	1/1	0.92	0.17	49,49,49,49	0
59	MG	DA	3858	1/1	0.92	0.13	49,49,49,49	0
59	MG	CA	2021	1/1	0.92	0.09	63,63,63,63	0
59	MG	DA	4340	1/1	0.92	0.26	60,60,60,60	0
59	MG	CS	105	1/1	0.92	0.14	69,69,69,69	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4514	1/1	0.92	0.27	80,80,80,80	0
59	MG	DA	3228	1/1	0.92	0.16	36,36,36,36	0
59	MG	DO	214	1/1	0.92	0.78	61,61,61,61	0
59	MG	A3	103	1/1	0.92	0.12	55,55,55,55	0
59	MG	AA	3562	1/1	0.92	0.11	55,55,55,55	0
59	MG	DA	3792	1/1	0.92	0.40	92,92,92,92	0
59	MG	DA	4468	1/1	0.92	0.27	60,60,60,60	0
59	MG	CA	1808	1/1	0.92	0.19	148,148,148,148	0
59	MG	AA	3550	1/1	0.92	0.22	82,82,82,82	0
59	MG	DA	4846	1/1	0.92	0.18	73,73,73,73	0
59	MG	AA	3018	1/1	0.92	0.16	22,22,22,22	0
59	MG	BH	202	1/1	0.92	0.10	61,61,61,61	0
59	MG	DA	3833	1/1	0.92	0.14	65,65,65,65	0
59	MG	DA	3839	1/1	0.92	0.17	54,54,54,54	0
59	MG	CA	1665	1/1	0.92	0.15	31,31,31,31	0
59	MG	DB	248	1/1	0.92	0.18	76,76,76,76	0
59	MG	DA	3987	1/1	0.92	0.20	53,53,53,53	0
59	MG	AA	3115	1/1	0.92	0.36	76,76,76,76	0
59	MG	DA	3248	1/1	0.92	0.46	70,70,70,70	0
59	MG	DA	4530	1/1	0.92	0.17	72,72,72,72	0
59	MG	DA	3422	1/1	0.92	0.15	41,41,41,41	0
59	MG	DA	4853	1/1	0.92	0.21	80,80,80,80	0
59	MG	DD	310	1/1	0.92	0.45	64,64,64,64	0
59	MG	AA	3640	1/1	0.92	0.22	45,45,45,45	0
59	MG	A3	102	1/1	0.92	0.18	83,83,83,83	0
59	MG	DA	4974	1/1	0.92	0.19	57,57,57,57	0
59	MG	AA	3921	1/1	0.92	0.09	44,44,44,44	0
59	MG	AA	3399	1/1	0.92	0.19	75,75,75,75	0
59	MG	AA	3404	1/1	0.92	0.42	77,77,77,77	0
59	MG	CA	2322	1/1	0.92	0.21	88,88,88,88	0
59	MG	BA	1636	1/1	0.92	0.29	44,44,44,44	0
59	MG	CA	1741	1/1	0.92	0.13	66,66,66,66	0
59	MG	CA	1725	1/1	0.92	0.18	55,55,55,55	0
59	MG	DA	3691	1/1	0.92	0.16	69,69,69,69	0
59	MG	CA	1829	1/1	0.92	0.07	61,61,61,61	0
59	MG	DA	4527	1/1	0.92	0.11	53,53,53,53	0
59	MG	AA	3655	1/1	0.92	0.08	50,50,50,50	0
59	MG	CT	203	1/1	0.92	0.14	54,54,54,54	0
59	MG	CA	1796	1/1	0.92	0.14	67,67,67,67	0
59	MG	DA	4557	1/1	0.92	0.09	56,56,56,56	0
59	MG	CH	205	1/1	0.92	0.18	87,87,87,87	0
59	MG	DA	4065	1/1	0.92	0.11	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	1668	1/1	0.92	0.21	88,88,88,88	0
59	MG	BA	1703	1/1	0.92	0.12	40,40,40,40	0
59	MG	AA	3120	1/1	0.92	0.40	59,59,59,59	0
59	MG	CA	1822	1/1	0.92	0.09	56,56,56,56	0
59	MG	A1	203	1/1	0.92	0.21	77,77,77,77	0
59	MG	BA	1925	1/1	0.92	0.13	55,55,55,55	0
59	MG	DA	4214	1/1	0.92	0.18	50,50,50,50	0
59	MG	AA	3568	1/1	0.92	0.20	41,41,41,41	0
59	MG	AD	310	1/1	0.92	0.26	42,42,42,42	0
59	MG	AD	303	1/1	0.92	0.51	62,62,62,62	0
59	MG	DA	4363	1/1	0.92	0.13	42,42,42,42	0
59	MG	AA	3592	1/1	0.92	0.16	63,63,63,63	0
59	MG	DA	4923	1/1	0.92	0.10	74,74,74,74	0
59	MG	BA	1695	1/1	0.92	0.18	44,44,44,44	0
59	MG	BA	1631	1/1	0.92	0.25	48,48,48,48	0
59	MG	DA	4303	1/1	0.92	0.15	47,47,47,47	0
59	MG	CA	1805	1/1	0.92	0.06	52,52,52,52	0
59	MG	AA	3521	1/1	0.92	0.15	32,32,32,32	0
59	MG	DA	4750	1/1	0.92	0.15	68,68,68,68	0
59	MG	DK	202	1/1	0.92	0.13	66,66,66,66	0
59	MG	BA	2090	1/1	0.92	0.14	119,119,119,119	0
59	MG	AB	213	1/1	0.92	0.12	60,60,60,60	0
59	MG	BH	201	1/1	0.92	0.23	55,55,55,55	0
59	MG	DA	4816	1/1	0.92	0.22	66,66,66,66	0
59	MG	DA	5007	1/1	0.92	0.45	82,82,82,82	0
59	MG	DA	4694	1/1	0.92	0.15	56,56,56,56	0
59	MG	DA	3558	1/1	0.92	0.34	64,64,64,64	0
59	MG	BA	1709	1/1	0.92	0.29	53,53,53,53	0
59	MG	BA	2094	1/1	0.92	0.14	50,50,50,50	0
59	MG	BD	101	1/1	0.92	0.35	86,86,86,86	0
59	MG	BA	1876	1/1	0.92	0.07	62,62,62,62	0
59	MG	AA	3505	1/1	0.92	0.16	55,55,55,55	0
59	MG	AA	3293	1/1	0.92	0.11	45,45,45,45	0
59	MG	BA	1742	1/1	0.92	0.25	46,46,46,46	0
59	MG	AA	3971	1/1	0.92	0.18	85,85,85,85	0
59	MG	CA	2159	1/1	0.92	0.12	46,46,46,46	0
59	MG	DA	3885	1/1	0.92	0.11	57,57,57,57	0
59	MG	C1	103	1/1	0.92	0.13	64,64,64,64	0
59	MG	CA	2115	1/1	0.92	0.29	78,78,78,78	0
59	MG	D2	206	1/1	0.92	0.21	55,55,55,55	0
59	MG	AA	3458	1/1	0.92	0.10	49,49,49,49	0
59	MG	DA	4267	1/1	0.92	0.48	72,72,72,72	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3754	1/1	0.92	0.08	107,107,107,107	0
59	MG	DA	3107	1/1	0.92	0.17	43,43,43,43	0
59	MG	DA	5018	1/1	0.92	0.27	89,89,89,89	0
59	MG	AA	3119	1/1	0.92	0.12	46,46,46,46	0
59	MG	CA	1652	1/1	0.92	0.29	39,39,39,39	0
59	MG	AA	3662	1/1	0.92	0.08	51,51,51,51	0
59	MG	DA	4787	1/1	0.92	0.35	85,85,85,85	0
59	MG	DA	3064	1/1	0.92	0.15	35,35,35,35	0
59	MG	DA	4695	1/1	0.92	0.14	52,52,52,52	0
59	MG	BA	1752	1/1	0.92	0.18	48,48,48,48	0
59	MG	DB	230	1/1	0.92	0.56	87,87,87,87	0
59	MG	DA	3325	1/1	0.92	0.24	62,62,62,62	0
59	MG	BA	2193	1/1	0.92	0.41	76,76,76,76	0
59	MG	CA	2008	1/1	0.92	0.13	56,56,56,56	0
59	MG	DA	3787	1/1	0.92	0.13	114,114,114,114	0
59	MG	D2	207	1/1	0.92	0.12	68,68,68,68	0
59	MG	CA	1738	1/1	0.92	0.29	66,66,66,66	0
59	MG	AA	3957	1/1	0.92	0.15	74,74,74,74	0
59	MG	DA	3096	1/1	0.92	0.15	7,7,7,7	0
59	MG	BD	112	1/1	0.92	0.21	78,78,78,78	0
59	MG	DA	3157	1/1	0.92	0.41	58,58,58,58	0
59	MG	DA	4836	1/1	0.92	0.13	41,41,41,41	0
59	MG	BA	1658	1/1	0.92	0.14	49,49,49,49	0
59	MG	BA	1823	1/1	0.92	0.08	84,84,84,84	0
59	MG	DA	3231	1/1	0.92	0.15	9,9,9,9	0
59	MG	AA	3384	1/1	0.92	0.27	47,47,47,47	0
59	MG	AB	235	1/1	0.92	0.08	81,81,81,81	0
59	MG	DA	3042	1/1	0.92	0.20	21,21,21,21	0
59	MG	DA	3500	1/1	0.92	0.12	57,57,57,57	0
59	MG	DA	3962	1/1	0.92	0.15	81,81,81,81	0
59	MG	DA	3222	1/1	0.92	0.09	42,42,42,42	0
59	MG	DP	201	1/1	0.92	0.19	29,29,29,29	0
59	MG	CA	1836	1/1	0.92	0.52	131,131,131,131	0
59	MG	AA	3624	1/1	0.92	0.30	38,38,38,38	0
59	MG	CA	2262	1/1	0.92	0.09	61,61,61,61	0
59	MG	CA	1628	1/1	0.92	0.36	42,42,42,42	0
59	MG	DB	217	1/1	0.92	0.25	46,46,46,46	0
59	MG	DA	4271	1/1	0.92	0.74	53,53,53,53	0
59	MG	AA	3954	1/1	0.92	0.23	71,71,71,71	0
59	MG	AA	3051	1/1	0.92	0.23	62,62,62,62	0
59	MG	DA	3911	1/1	0.92	0.12	115,115,115,115	0
59	MG	DA	4709	1/1	0.92	0.25	81,81,81,81	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3411	1/1	0.92	0.18	73,73,73,73	0
59	MG	AA	3798	1/1	0.92	0.08	47,47,47,47	0
59	MG	AA	3038	1/1	0.92	0.23	68,68,68,68	0
59	MG	BA	1945	1/1	0.92	0.14	56,56,56,56	0
59	MG	BA	2095	1/1	0.92	0.27	67,67,67,67	0
59	MG	D0	206	1/1	0.92	0.14	45,45,45,45	0
59	MG	CA	2130	1/1	0.92	0.23	117,117,117,117	0
59	MG	BA	2135	1/1	0.92	0.23	77,77,77,77	0
59	MG	BA	2005	1/1	0.92	0.04	75,75,75,75	0
59	MG	BA	2038	1/1	0.92	0.25	80,80,80,80	0
59	MG	BA	2079	1/1	0.92	0.17	99,99,99,99	0
59	MG	AA	3614	1/1	0.92	0.14	62,62,62,62	0
59	MG	CC	110	1/1	0.92	0.21	71,71,71,71	0
59	MG	DB	270	1/1	0.92	0.26	61,61,61,61	0
59	MG	D8	106	1/1	0.92	0.08	41,41,41,41	0
59	MG	BB	102	1/1	0.92	0.08	55,55,55,55	0
59	MG	BA	2229	1/1	0.92	0.12	67,67,67,67	0
59	MG	BA	1998	1/1	0.92	0.13	76,76,76,76	0
59	MG	AA	4161	1/1	0.92	0.15	68,68,68,68	0
59	MG	BA	1861	1/1	0.92	0.11	78,78,78,78	0
59	MG	DA	5035	1/1	0.92	0.24	51,51,51,51	0
59	MG	AA	3149	1/1	0.92	0.26	61,61,61,61	0
59	MG	DA	3534	1/1	0.92	0.37	49,49,49,49	0
59	MG	CA	1875	1/1	0.92	0.12	71,71,71,71	0
59	MG	CA	1823	1/1	0.92	0.09	67,67,67,67	0
59	MG	DA	3488	1/1	0.92	0.25	37,37,37,37	0
59	MG	AA	3877	1/1	0.92	0.28	109,109,109,109	0
59	MG	DA	4309	1/1	0.92	0.09	36,36,36,36	0
59	MG	DA	4253	1/1	0.92	0.27	62,62,62,62	0
59	MG	CA	2136	1/1	0.92	0.23	77,77,77,77	0
59	MG	CE	304	1/1	0.92	0.12	62,62,62,62	0
59	MG	DA	3789	1/1	0.92	0.13	45,45,45,45	0
59	MG	DB	272	1/1	0.92	0.13	81,81,81,81	0
59	MG	DA	3750	1/1	0.92	0.44	101,101,101,101	0
59	MG	DA	3202	1/1	0.92	0.20	37,37,37,37	0
59	MG	DA	3165	1/1	0.92	0.31	46,46,46,46	0
59	MG	CA	2300	1/1	0.92	0.16	119,119,119,119	0
59	MG	DA	3639	1/1	0.92	0.21	66,66,66,66	0
59	MG	CA	1654	1/1	0.92	0.22	48,48,48,48	0
59	MG	BA	1976	1/1	0.92	0.15	147,147,147,147	0
59	MG	AO	205	1/1	0.92	0.18	52,52,52,52	0
59	MG	DA	3680	1/1	0.92	0.23	76,76,76,76	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	BA	1992	1/1	0.92	0.17	95,95,95,95	0
59	MG	DA	3289	1/1	0.92	0.19	27,27,27,27	0
59	MG	BA	1615	1/1	0.92	0.19	81,81,81,81	0
59	MG	CD	115	1/1	0.92	0.50	85,85,85,85	0
59	MG	CA	2245	1/1	0.92	0.41	133,133,133,133	0
59	MG	BA	1801	1/1	0.92	0.23	40,40,40,40	0
59	MG	AA	3291	1/1	0.92	0.16	34,34,34,34	0
59	MG	DA	4912	1/1	0.92	0.14	42,42,42,42	0
59	MG	AA	3233	1/1	0.92	0.20	55,55,55,55	0
59	MG	CD	117	1/1	0.92	0.19	58,58,58,58	0
59	MG	DA	4642	1/1	0.92	0.25	57,57,57,57	0
59	MG	DA	3124	1/1	0.92	0.20	36,36,36,36	0
59	MG	CA	1604	1/1	0.92	0.24	10,10,10,10	0
59	MG	BA	2055	1/1	0.92	0.08	89,89,89,89	0
59	MG	CA	2009	1/1	0.92	0.13	61,61,61,61	0
59	MG	D5	101	1/1	0.92	0.21	31,31,31,31	0
59	MG	DR	202	1/1	0.92	0.52	60,60,60,60	0
59	MG	BA	1907	1/1	0.92	0.15	56,56,56,56	0
59	MG	AA	3865	1/1	0.92	0.19	62,62,62,62	0
59	MG	BA	1983	1/1	0.92	0.05	80,80,80,80	0
59	MG	CA	1904	1/1	0.92	0.37	74,74,74,74	0
59	MG	AA	3809	1/1	0.92	0.15	84,84,84,84	0
59	MG	CA	1727	1/1	0.92	0.23	56,56,56,56	0
59	MG	DB	204	1/1	0.92	0.10	23,23,23,23	0
59	MG	AA	3280	1/1	0.92	0.26	56,56,56,56	0
59	MG	DA	3565	1/1	0.93	0.20	55,55,55,55	0
59	MG	AA	3029	1/1	0.93	0.20	46,46,46,46	0
59	MG	DA	3486	1/1	0.93	0.20	58,58,58,58	0
59	MG	BA	1825	1/1	0.93	0.10	121,121,121,121	0
59	MG	DA	4141	1/1	0.93	0.55	66,66,66,66	0
59	MG	DA	3924	1/1	0.93	0.16	48,48,48,48	0
59	MG	BA	2223	1/1	0.93	0.30	71,71,71,71	0
59	MG	CA	1742	1/1	0.93	0.13	40,40,40,40	0
59	MG	DA	3331	1/1	0.93	0.26	68,68,68,68	0
59	MG	CA	1866	1/1	0.93	0.20	109,109,109,109	0
59	MG	AA	3392	1/1	0.93	0.14	63,63,63,63	0
59	MG	DA	4001	1/1	0.93	0.22	121,121,121,121	0
59	MG	DB	265	1/1	0.93	0.21	81,81,81,81	0
59	MG	DB	208	1/1	0.93	0.10	46,46,46,46	0
59	MG	DA	4274	1/1	0.93	0.17	41,41,41,41	0
59	MG	DA	3426	1/1	0.93	0.41	39,39,39,39	0
59	MG	DA	3046	1/1	0.93	0.30	12,12,12,12	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3453	1/1	0.93	0.09	123,123,123,123	0
59	MG	CA	2218	1/1	0.93	0.20	82,82,82,82	0
59	MG	DA	4192	1/1	0.93	0.57	65,65,65,65	0
59	MG	DA	4132	1/1	0.93	0.31	40,40,40,40	0
59	MG	DA	3524	1/1	0.93	0.07	32,32,32,32	0
59	MG	DA	3867	1/1	0.93	0.25	100,100,100,100	0
59	MG	AA	3349	1/1	0.93	0.11	42,42,42,42	0
59	MG	A7	104	1/1	0.93	0.27	88,88,88,88	0
59	MG	CA	1983	1/1	0.93	0.33	43,43,43,43	0
59	MG	AA	3978	1/1	0.93	0.48	70,70,70,70	0
59	MG	CA	2112	1/1	0.93	0.14	78,78,78,78	0
59	MG	AA	3901	1/1	0.93	0.19	52,52,52,52	0
59	MG	AA	4047	1/1	0.93	0.10	59,59,59,59	0
59	MG	DA	3355	1/1	0.93	0.31	64,64,64,64	0
59	MG	BA	1719	1/1	0.93	0.18	43,43,43,43	0
59	MG	DA	4564	1/1	0.93	0.16	39,39,39,39	0
59	MG	DA	3391	1/1	0.93	0.42	84,84,84,84	0
59	MG	DD	309	1/1	0.93	0.26	57,57,57,57	0
59	MG	DT	107	1/1	0.93	0.21	73,73,73,73	0
59	MG	CA	2072	1/1	0.93	0.30	48,48,48,48	0
59	MG	DA	3192	1/1	0.93	0.27	95,95,95,95	0
59	MG	DA	4202	1/1	0.93	0.42	69,69,69,69	0
59	MG	CA	1626	1/1	0.93	0.19	38,38,38,38	0
59	MG	AA	3700	1/1	0.93	0.20	58,58,58,58	0
59	MG	DA	4070	1/1	0.93	0.22	45,45,45,45	0
59	MG	DA	3163	1/1	0.93	0.19	39,39,39,39	0
59	MG	DA	3801	1/1	0.93	0.64	96,96,96,96	0
59	MG	CA	1737	1/1	0.93	0.20	77,77,77,77	0
59	MG	DA	3196	1/1	0.93	0.29	43,43,43,43	0
59	MG	DA	3078	1/1	0.93	0.26	17,17,17,17	0
59	MG	DA	4299	1/1	0.93	0.11	72,72,72,72	0
59	MG	DA	3943	1/1	0.93	0.08	47,47,47,47	0
59	MG	AA	3716	1/1	0.93	0.12	74,74,74,74	0
59	MG	BA	1982	1/1	0.93	0.12	96,96,96,96	0
59	MG	AA	3100	1/1	0.93	0.10	34,34,34,34	0
59	MG	BA	1736	1/1	0.93	0.14	53,53,53,53	0
59	MG	AA	3061	1/1	0.93	0.21	35,35,35,35	0
59	MG	DA	4106	1/1	0.93	0.13	65,65,65,65	0
59	MG	AA	3956	1/1	0.93	0.44	93,93,93,93	0
59	MG	DA	4260	1/1	0.93	0.18	56,56,56,56	0
59	MG	BA	1902	1/1	0.93	0.33	71,71,71,71	0
59	MG	BA	1664	1/1	0.93	0.32	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3203	1/1	0.93	0.45	59,59,59,59	0
59	MG	DA	4335	1/1	0.93	0.20	50,50,50,50	0
59	MG	DA	4930	1/1	0.93	0.26	73,73,73,73	0
59	MG	AA	3849	1/1	0.93	0.39	59,59,59,59	0
59	MG	CA	1734	1/1	0.93	0.12	38,38,38,38	0
59	MG	DA	4866	1/1	0.93	0.23	53,53,53,53	0
59	MG	BB	103	1/1	0.93	0.11	77,77,77,77	0
59	MG	A8	101	1/1	0.93	0.09	45,45,45,45	0
59	MG	DA	3510	1/1	0.93	0.43	56,56,56,56	0
59	MG	DF	310	1/1	0.93	0.12	49,49,49,49	0
59	MG	DA	4762	1/1	0.93	0.23	77,77,77,77	0
59	MG	BA	2231	1/1	0.93	0.30	89,89,89,89	0
59	MG	DA	4596	1/1	0.93	0.33	63,63,63,63	0
59	MG	DA	3363	1/1	0.93	0.07	36,36,36,36	0
59	MG	BA	1681	1/1	0.93	0.17	43,43,43,43	0
59	MG	CA	2052	1/1	0.93	0.19	54,54,54,54	0
59	MG	DA	4601	1/1	0.93	0.20	88,88,88,88	0
59	MG	DA	4689	1/1	0.93	0.12	67,67,67,67	0
59	MG	CA	2275	1/1	0.93	0.23	42,42,42,42	0
59	MG	AA	3263	1/1	0.93	0.67	57,57,57,57	0
59	MG	AA	3558	1/1	0.93	0.12	50,50,50,50	0
59	MG	DA	4207	1/1	0.93	0.34	84,84,84,84	0
59	MG	CB	111	1/1	0.93	0.15	54,54,54,54	0
59	MG	DA	4499	1/1	0.93	0.21	51,51,51,51	0
59	MG	DA	3128	1/1	0.93	0.17	31,31,31,31	0
59	MG	DB	256	1/1	0.93	0.15	65,65,65,65	0
59	MG	AA	4072	1/1	0.93	0.13	56,56,56,56	0
59	MG	CA	2153	1/1	0.93	0.12	76,76,76,76	0
59	MG	DA	3117	1/1	0.93	0.17	58,58,58,58	0
59	MG	DA	3980	1/1	0.93	0.59	65,65,65,65	0
59	MG	AA	4091	1/1	0.93	0.13	107,107,107,107	0
59	MG	BA	1811	1/1	0.93	0.04	65,65,65,65	0
59	MG	BA	2145	1/1	0.93	0.20	96,96,96,96	0
59	MG	AA	3881	1/1	0.93	0.12	72,72,72,72	0
59	MG	CA	1999	1/1	0.93	0.14	63,63,63,63	0
59	MG	DA	4624	1/1	0.93	0.12	68,68,68,68	0
59	MG	DA	3098	1/1	0.93	0.12	34,34,34,34	0
59	MG	AA	3854	1/1	0.93	0.25	71,71,71,71	0
59	MG	DA	3922	1/1	0.93	0.21	69,69,69,69	0
59	MG	DA	4843	1/1	0.93	0.30	72,72,72,72	0
59	MG	CA	1888	1/1	0.93	0.41	72,72,72,72	0
59	MG	AA	3192	1/1	0.93	0.11	39,39,39,39	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3723	1/1	0.93	0.21	19,19,19,19	0
59	MG	DA	3317	1/1	0.93	0.13	86,86,86,86	0
59	MG	DA	4676	1/1	0.93	0.46	82,82,82,82	0
59	MG	BA	2056	1/1	0.93	0.27	69,69,69,69	0
59	MG	CC	101	1/1	0.93	0.22	41,41,41,41	0
59	MG	BD	119	1/1	0.93	0.40	84,84,84,84	0
59	MG	DA	3330	1/1	0.93	0.10	70,70,70,70	0
59	MG	AA	3273	1/1	0.93	0.15	30,30,30,30	0
59	MG	AA	4062	1/1	0.93	0.14	81,81,81,81	0
59	MG	DA	4606	1/1	0.93	0.69	41,41,41,41	0
59	MG	BA	1810	1/1	0.93	0.11	93,93,93,93	0
59	MG	DA	3569	1/1	0.93	0.20	21,21,21,21	0
59	MG	AA	3295	1/1	0.93	0.34	62,62,62,62	0
59	MG	AA	3626	1/1	0.93	0.22	48,48,48,48	0
59	MG	BC	110	1/1	0.93	0.14	56,56,56,56	0
59	MG	AA	3967	1/1	0.93	0.17	63,63,63,63	0
59	MG	DA	3970	1/1	0.93	0.22	50,50,50,50	0
59	MG	DA	4392	1/1	0.93	0.14	52,52,52,52	0
59	MG	AA	3947	1/1	0.93	0.24	84,84,84,84	0
59	MG	DA	4560	1/1	0.93	0.15	61,61,61,61	0
59	MG	AA	3755	1/1	0.93	0.08	63,63,63,63	0
59	MG	DA	4786	1/1	0.93	0.14	81,81,81,81	0
59	MG	BG	303	1/1	0.93	0.10	59,59,59,59	0
59	MG	DA	4382	1/1	0.93	0.13	67,67,67,67	0
59	MG	CA	1895	1/1	0.93	0.15	58,58,58,58	0
59	MG	AA	3845	1/1	0.93	0.11	54,54,54,54	0
59	MG	DA	3479	1/1	0.93	0.29	74,74,74,74	0
59	MG	AA	3201	1/1	0.93	0.10	71,71,71,71	0
59	MG	DA	4322	1/1	0.93	0.33	65,65,65,65	0
59	MG	DA	3909	1/1	0.93	0.25	69,69,69,69	0
59	MG	DA	4868	1/1	0.93	0.13	70,70,70,70	0
59	MG	CA	1814	1/1	0.93	0.07	42,42,42,42	0
59	MG	AA	3953	1/1	0.93	0.13	60,60,60,60	0
59	MG	BB	104	1/1	0.93	0.14	71,71,71,71	0
59	MG	AA	3653	1/1	0.93	0.14	58,58,58,58	0
59	MG	BC	108	1/1	0.93	0.13	81,81,81,81	0
59	MG	DA	4979	1/1	0.93	0.26	47,47,47,47	0
59	MG	BA	1846	1/1	0.93	0.13	63,63,63,63	0
59	MG	BA	2175	1/1	0.93	0.17	81,81,81,81	0
59	MG	DA	4789	1/1	0.93	0.21	46,46,46,46	0
59	MG	DA	3591	1/1	0.93	0.15	16,16,16,16	0
59	MG	BA	1843	1/1	0.93	0.07	63,63,63,63	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3625	1/1	0.93	0.34	19,19,19,19	0
59	MG	AA	3142	1/1	0.93	0.09	12,12,12,12	0
59	MG	CA	1705	1/1	0.93	0.30	54,54,54,54	0
59	MG	BA	1684	1/1	0.93	0.13	36,36,36,36	0
59	MG	AA	3162	1/1	0.93	0.10	48,48,48,48	0
59	MG	DA	3359	1/1	0.93	0.30	43,43,43,43	0
59	MG	AA	3157	1/1	0.93	0.31	73,73,73,73	0
59	MG	AA	3887	1/1	0.93	0.17	70,70,70,70	0
59	MG	BA	1728	1/1	0.93	0.14	47,47,47,47	0
59	MG	BA	2072	1/1	0.93	0.14	72,72,72,72	0
59	MG	DA	4677	1/1	0.93	0.18	61,61,61,61	0
59	MG	DA	3853	1/1	0.93	0.24	71,71,71,71	0
59	MG	DA	4255	1/1	0.93	0.30	54,54,54,54	0
59	MG	DA	4399	1/1	0.93	0.25	47,47,47,47	0
59	MG	DA	3176	1/1	0.93	0.23	39,39,39,39	0
59	MG	AA	4135	1/1	0.93	0.14	64,64,64,64	0
59	MG	DA	3264	1/1	0.93	0.26	28,28,28,28	0
59	MG	CA	2281	1/1	0.93	0.12	100,100,100,100	0
59	MG	AP	201	1/1	0.93	0.16	95,95,95,95	0
59	MG	AA	3737	1/1	0.93	0.13	55,55,55,55	0
59	MG	CG	311	1/1	0.93	0.20	150,150,150,150	0
59	MG	AA	3803	1/1	0.93	0.17	72,72,72,72	0
59	MG	DA	4921	1/1	0.93	0.27	67,67,67,67	0
59	MG	AA	3512	1/1	0.93	0.09	31,31,31,31	0
59	MG	A1	202	1/1	0.93	0.46	61,61,61,61	0
59	MG	CA	2098	1/1	0.93	0.09	64,64,64,64	0
59	MG	DA	3336	1/1	0.93	0.13	36,36,36,36	0
59	MG	DA	4476	1/1	0.93	0.11	64,64,64,64	0
59	MG	DA	4944	1/1	0.93	0.22	56,56,56,56	0
59	MG	DA	4538	1/1	0.93	0.27	92,92,92,92	0
59	MG	A5	102	1/1	0.93	0.27	28,28,28,28	0
59	MG	CA	2283	1/1	0.93	0.11	68,68,68,68	0
59	MG	AA	3987	1/1	0.93	0.16	54,54,54,54	0
59	MG	BA	2065	1/1	0.93	0.14	73,73,73,73	0
59	MG	CA	2078	1/1	0.93	0.20	75,75,75,75	0
59	MG	DA	3100	1/1	0.93	0.19	36,36,36,36	0
59	MG	CK	206	1/1	0.93	0.21	62,62,62,62	0
59	MG	DA	3930	1/1	0.93	0.22	72,72,72,72	0
59	MG	AA	3891	1/1	0.93	0.30	105,105,105,105	0
59	MG	DA	4182	1/1	0.93	0.40	82,82,82,82	0
59	MG	CA	1922	1/1	0.93	0.09	84,84,84,84	0
59	MG	CA	1986	1/1	0.93	0.13	95,95,95,95	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4432	1/1	0.93	0.16	58,58,58,58	0
59	MG	DA	3084	1/1	0.93	0.29	10,10,10,10	0
59	MG	DA	4368	1/1	0.93	0.15	51,51,51,51	0
59	MG	AA	3782	1/1	0.93	0.43	101,101,101,101	0
59	MG	AA	3762	1/1	0.93	0.18	63,63,63,63	0
59	MG	DA	4183	1/1	0.93	0.10	53,53,53,53	0
59	MG	DA	5022	1/1	0.93	0.09	82,82,82,82	0
59	MG	DA	4707	1/1	0.93	0.31	86,86,86,86	0
59	MG	DA	4436	1/1	0.93	0.06	76,76,76,76	0
59	MG	DA	5016	1/1	0.93	0.19	106,106,106,106	0
59	MG	AA	3127	1/1	0.93	0.09	46,46,46,46	0
59	MG	BW	206	1/1	0.93	0.07	56,56,56,56	0
59	MG	DA	3476	1/1	0.93	0.10	27,27,27,27	0
59	MG	DA	3978	1/1	0.93	0.56	82,82,82,82	0
59	MG	CA	1978	1/1	0.93	0.07	93,93,93,93	0
59	MG	DA	3951	1/1	0.93	0.25	136,136,136,136	0
59	MG	CA	2066	1/1	0.93	0.32	77,77,77,77	0
59	MG	DA	4320	1/1	0.93	0.17	50,50,50,50	0
59	MG	DA	4412	1/1	0.93	0.13	82,82,82,82	0
59	MG	AA	4045	1/1	0.93	0.10	57,57,57,57	0
59	MG	AA	4039	1/1	0.93	0.16	95,95,95,95	0
59	MG	CA	1602	1/1	0.93	0.16	89,89,89,89	0
59	MG	BA	1622	1/1	0.93	0.09	64,64,64,64	0
59	MG	CA	1695	1/1	0.93	0.23	50,50,50,50	0
59	MG	DA	3318	1/1	0.93	0.14	72,72,72,72	0
59	MG	CA	1663	1/1	0.93	0.10	52,52,52,52	0
59	MG	DA	4469	1/1	0.93	0.43	52,52,52,52	0
59	MG	AA	3351	1/1	0.93	0.10	62,62,62,62	0
59	MG	DA	4919	1/1	0.93	0.31	58,58,58,58	0
59	MG	DA	4706	1/1	0.93	0.35	89,89,89,89	0
59	MG	CA	2288	1/1	0.93	0.24	74,74,74,74	0
59	MG	AA	3236	1/1	0.93	0.05	20,20,20,20	0
59	MG	CA	2038	1/1	0.93	0.09	74,74,74,74	0
59	MG	AA	3258	1/1	0.93	0.44	81,81,81,81	0
59	MG	AA	3787	1/1	0.93	0.12	70,70,70,70	0
59	MG	D1	207	1/1	0.93	0.25	64,64,64,64	0
59	MG	AA	3715	1/1	0.93	0.08	47,47,47,47	0
59	MG	BA	1875	1/1	0.93	0.08	125,125,125,125	0
59	MG	DA	3914	1/1	0.93	0.09	36,36,36,36	0
59	MG	BA	1946	1/1	0.93	0.15	70,70,70,70	0
59	MG	DF	316	1/1	0.93	0.21	84,84,84,84	0
59	MG	AA	3412	1/1	0.93	0.17	30,30,30,30	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3204	1/1	0.93	0.15	50,50,50,50	0
59	MG	AA	4104	1/1	0.93	0.37	71,71,71,71	0
59	MG	AA	3123	1/1	0.93	0.20	47,47,47,47	0
59	MG	BA	2236	1/1	0.93	0.26	82,82,82,82	0
59	MG	DA	4966	1/1	0.93	0.14	64,64,64,64	0
59	MG	DA	3878	1/1	0.93	0.26	60,60,60,60	0
59	MG	AT	104	1/1	0.93	0.29	89,89,89,89	0
59	MG	DF	321	1/1	0.93	0.50	55,55,55,55	0
59	MG	DB	264	1/1	0.93	0.13	44,44,44,44	0
59	MG	DA	4383	1/1	0.93	0.45	85,85,85,85	0
59	MG	DA	5073	1/1	0.93	0.47	127,127,127,127	0
59	MG	CB	110	1/1	0.93	0.10	39,39,39,39	0
59	MG	DA	4840	1/1	0.93	0.29	111,111,111,111	0
59	MG	AA	3900	1/1	0.93	0.32	105,105,105,105	0
59	MG	AA	3415	1/1	0.93	0.06	63,63,63,63	0
59	MG	BA	1627	1/1	0.93	0.37	60,60,60,60	0
59	MG	DA	3293	1/1	0.93	0.12	35,35,35,35	0
59	MG	DA	4305	1/1	0.93	0.19	46,46,46,46	0
59	MG	BS	102	1/1	0.93	0.06	69,69,69,69	0
59	MG	AA	3246	1/1	0.93	0.20	45,45,45,45	0
59	MG	CA	1691	1/1	0.93	0.16	33,33,33,33	0
59	MG	DA	4104	1/1	0.93	0.36	76,76,76,76	0
59	MG	BA	2181	1/1	0.93	0.17	93,93,93,93	0
59	MG	BA	2121	1/1	0.93	0.32	84,84,84,84	0
59	MG	AA	3673	1/1	0.93	0.27	56,56,56,56	0
59	MG	CA	2249	1/1	0.93	0.09	68,68,68,68	0
59	MG	AA	3308	1/1	0.93	0.10	38,38,38,38	0
59	MG	DA	3272	1/1	0.93	0.25	29,29,29,29	0
59	MG	AA	4132	1/1	0.93	0.17	68,68,68,68	0
59	MG	AA	3858	1/1	0.93	0.21	76,76,76,76	0
59	MG	AT	102	1/1	0.93	0.14	57,57,57,57	0
59	MG	CA	1642	1/1	0.93	0.10	25,25,25,25	0
59	MG	AA	3721	1/1	0.93	0.69	82,82,82,82	0
59	MG	CA	1623	1/1	0.93	0.31	49,49,49,49	0
59	MG	CD	119	1/1	0.93	0.14	60,60,60,60	0
59	MG	AA	3023	1/1	0.93	0.21	59,59,59,59	0
59	MG	BA	1730	1/1	0.93	0.12	37,37,37,37	0
59	MG	AA	3197	1/1	0.93	0.11	50,50,50,50	0
59	MG	AB	232	1/1	0.93	0.27	70,70,70,70	0
59	MG	DA	3382	1/1	0.93	0.35	47,47,47,47	0
59	MG	DA	3636	1/1	0.93	0.10	46,46,46,46	0
59	MG	DA	4093	1/1	0.93	0.25	85,85,85,85	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3618	1/1	0.93	0.15	89,89,89,89	0
59	MG	CA	1976	1/1	0.93	0.21	88,88,88,88	0
59	MG	AA	3019	1/1	0.93	0.30	31,31,31,31	0
59	MG	DA	3457	1/1	0.93	0.28	76,76,76,76	0
59	MG	CA	1657	1/1	0.93	0.19	35,35,35,35	0
59	MG	C1	105	1/1	0.93	0.43	89,89,89,89	0
59	MG	BA	2108	1/1	0.93	0.22	154,154,154,154	0
59	MG	CA	1608	1/1	0.93	0.31	48,48,48,48	0
59	MG	BA	1824	1/1	0.93	0.11	107,107,107,107	0
59	MG	DA	4528	1/1	0.93	0.15	65,65,65,65	0
59	MG	DA	4124	1/1	0.93	0.08	58,58,58,58	0
59	MG	BA	2015	1/1	0.93	0.16	71,71,71,71	0
59	MG	AA	3307	1/1	0.93	0.19	50,50,50,50	0
59	MG	AA	3977	1/1	0.93	0.32	56,56,56,56	0
59	MG	CA	1788	1/1	0.93	0.07	79,79,79,79	0
59	MG	CA	2301	1/1	0.93	0.14	55,55,55,55	0
59	MG	DA	3551	1/1	0.93	0.28	29,29,29,29	0
59	MG	D0	204	1/1	0.93	0.19	27,27,27,27	0
59	MG	AA	3972	1/1	0.93	0.65	60,60,60,60	0
59	MG	BA	1848	1/1	0.93	0.10	56,56,56,56	0
59	MG	BA	1860	1/1	0.93	0.07	60,60,60,60	0
59	MG	CA	1601	1/1	0.93	0.08	41,41,41,41	0
59	MG	AF	307	1/1	0.93	0.24	63,63,63,63	0
59	MG	BA	2165	1/1	0.93	0.10	61,61,61,61	0
59	MG	CA	2091	1/1	0.93	0.25	68,68,68,68	0
59	MG	DA	3553	1/1	0.93	0.39	89,89,89,89	0
59	MG	DA	4243	1/1	0.93	0.17	59,59,59,59	0
59	MG	DA	4678	1/1	0.93	0.13	68,68,68,68	0
59	MG	DA	4774	1/1	0.93	0.16	70,70,70,70	0
59	MG	AO	207	1/1	0.93	0.33	102,102,102,102	0
59	MG	DA	3489	1/1	0.93	0.11	41,41,41,41	0
59	MG	DA	3812	1/1	0.93	0.10	73,73,73,73	0
59	MG	AA	3264	1/1	0.93	0.15	43,43,43,43	0
59	MG	CC	104	1/1	0.93	0.08	23,23,23,23	0
59	MG	AA	3164	1/1	0.93	0.27	79,79,79,79	0
59	MG	DA	5023	1/1	0.93	0.21	62,62,62,62	0
59	MG	AA	3269	1/1	0.93	0.26	61,61,61,61	0
59	MG	CK	201	1/1	0.93	0.33	102,102,102,102	0
59	MG	DA	4073	1/1	0.93	0.33	106,106,106,106	0
59	MG	CA	1958	1/1	0.93	0.18	75,75,75,75	0
59	MG	AA	3242	1/1	0.93	0.25	56,56,56,56	0
59	MG	AA	3556	1/1	0.93	0.20	74,74,74,74	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	D3	104	1/1	0.93	0.20	32,32,32,32	0
59	MG	DA	4640	1/1	0.93	0.21	66,66,66,66	0
59	MG	DA	4991	1/1	0.93	0.21	123,123,123,123	0
59	MG	DA	3315	1/1	0.93	0.26	28,28,28,28	0
59	MG	BA	2054	1/1	0.93	0.19	68,68,68,68	0
59	MG	DA	3730	1/1	0.93	0.28	47,47,47,47	0
59	MG	DA	4519	1/1	0.93	0.13	118,118,118,118	0
59	MG	BA	1692	1/1	0.93	0.23	34,34,34,34	0
59	MG	DA	3008	1/1	0.93	0.26	10,10,10,10	0
59	MG	BA	1887	1/1	0.93	0.17	56,56,56,56	0
59	MG	DA	4578	1/1	0.93	0.16	95,95,95,95	0
59	MG	BA	1948	1/1	0.93	0.34	75,75,75,75	0
59	MG	BA	1889	1/1	0.93	0.07	37,37,37,37	0
59	MG	DA	4763	1/1	0.93	0.45	51,51,51,51	0
59	MG	DA	3378	1/1	0.93	0.36	55,55,55,55	0
59	MG	DA	4373	1/1	0.93	0.20	62,62,62,62	0
59	MG	AA	4113	1/1	0.93	0.50	92,92,92,92	0
59	MG	AA	3410	1/1	0.93	0.29	84,84,84,84	0
59	MG	DA	3548	1/1	0.93	0.24	58,58,58,58	0
59	MG	DA	4785	1/1	0.94	0.13	75,75,75,75	0
59	MG	DA	4633	1/1	0.94	0.25	66,66,66,66	0
59	MG	AA	3602	1/1	0.94	0.08	102,102,102,102	0
59	MG	AA	3302	1/1	0.94	0.13	32,32,32,32	0
59	MG	DA	5034	1/1	0.94	0.34	76,76,76,76	0
59	MG	DA	4097	1/1	0.94	0.19	64,64,64,64	0
59	MG	DA	4941	1/1	0.94	0.52	88,88,88,88	0
59	MG	BA	1649	1/1	0.94	0.25	63,63,63,63	0
59	MG	AA	4139	1/1	0.94	0.10	67,67,67,67	0
59	MG	DA	4807	1/1	0.94	0.29	68,68,68,68	0
59	MG	BA	2008	1/1	0.94	0.07	50,50,50,50	0
59	MG	DU	201	1/1	0.94	0.13	36,36,36,36	0
59	MG	BA	1625	1/1	0.94	0.28	74,74,74,74	0
59	MG	CS	102	1/1	0.94	0.15	65,65,65,65	0
59	MG	DA	4153	1/1	0.94	0.23	48,48,48,48	0
59	MG	DX	101	1/1	0.94	0.15	59,59,59,59	0
59	MG	CA	2100	1/1	0.94	0.17	40,40,40,40	0
59	MG	DA	3982	1/1	0.94	0.38	106,106,106,106	0
59	MG	DA	3054	1/1	0.94	0.15	11,11,11,11	0
59	MG	DA	3590	1/1	0.94	0.19	46,46,46,46	0
59	MG	AA	3654	1/1	0.94	0.18	59,59,59,59	0
59	MG	A6	102	1/1	0.94	0.07	60,60,60,60	0
59	MG	AA	3708	1/1	0.94	0.15	49,49,49,49	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	1720	1/1	0.94	0.38	80,80,80,80	0
59	MG	AA	3318	1/1	0.94	0.28	65,65,65,65	0
59	MG	DA	3151	1/1	0.94	0.25	49,49,49,49	0
59	MG	CR	102	1/1	0.94	0.16	43,43,43,43	0
59	MG	AA	3907	1/1	0.94	1.04	57,57,57,57	0
59	MG	CB	104	1/1	0.94	0.13	51,51,51,51	0
59	MG	DA	4371	1/1	0.94	0.27	57,57,57,57	0
59	MG	AA	3861	1/1	0.94	0.30	99,99,99,99	0
59	MG	AA	3424	1/1	0.94	0.13	54,54,54,54	0
59	MG	DA	3138	1/1	0.94	0.21	44,44,44,44	0
59	MG	DA	3884	1/1	0.94	0.27	60,60,60,60	0
59	MG	DA	4671	1/1	0.94	0.22	72,72,72,72	0
59	MG	AA	4053	1/1	0.94	0.17	63,63,63,63	0
59	MG	DA	3614	1/1	0.94	0.20	53,53,53,53	0
59	MG	DA	4259	1/1	0.94	0.18	54,54,54,54	0
59	MG	AA	4038	1/1	0.94	0.14	77,77,77,77	0
59	MG	DF	301	1/1	0.94	0.17	28,28,28,28	0
59	MG	CA	2203	1/1	0.94	0.11	84,84,84,84	0
59	MG	D6	101	1/1	0.94	0.27	63,63,63,63	0
59	MG	DA	3759	1/1	0.94	0.13	84,84,84,84	0
59	MG	CV	104	1/1	0.94	0.17	86,86,86,86	0
59	MG	AA	3016	1/1	0.94	0.14	9,9,9,9	0
59	MG	D0	201	1/1	0.94	0.30	37,37,37,37	0
59	MG	DA	4197	1/1	0.94	0.19	51,51,51,51	0
59	MG	DA	4289	1/1	0.94	0.31	53,53,53,53	0
59	MG	AA	3373	1/1	0.94	0.15	48,48,48,48	0
59	MG	CA	2323	1/1	0.94	0.31	88,88,88,88	0
59	MG	BA	1952	1/1	0.94	0.32	70,70,70,70	0
59	MG	AA	3306	1/1	0.94	0.29	49,49,49,49	0
59	MG	DA	4749	1/1	0.94	0.22	78,78,78,78	0
59	MG	AA	3627	1/1	0.94	0.28	37,37,37,37	0
59	MG	BA	2103	1/1	0.94	0.14	80,80,80,80	0
59	MG	AA	3607	1/1	0.94	0.09	94,94,94,94	0
59	MG	BA	1614	1/1	0.94	0.33	73,73,73,73	0
59	MG	CA	1806	1/1	0.94	0.13	98,98,98,98	0
59	MG	DA	3178	1/1	0.94	0.27	49,49,49,49	0
59	MG	D0	210	1/1	0.94	0.15	75,75,75,75	0
59	MG	DD	303	1/1	0.94	0.13	16,16,16,16	0
59	MG	CA	1816	1/1	0.94	0.14	86,86,86,86	0
59	MG	CA	1786	1/1	0.94	0.22	93,93,93,93	0
59	MG	DA	3269	1/1	0.94	0.09	36,36,36,36	0
59	MG	CA	2143	1/1	0.94	0.20	57,57,57,57	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4639	1/1	0.94	0.12	75,75,75,75	0
59	MG	AA	3534	1/1	0.94	0.18	32,32,32,32	0
59	MG	AA	3032	1/1	0.94	0.20	28,28,28,28	0
59	MG	DA	3762	1/1	0.94	0.18	89,89,89,89	0
59	MG	CB	102	1/1	0.94	0.10	65,65,65,65	0
59	MG	DA	4999	1/1	0.94	0.24	74,74,74,74	0
59	MG	DA	3932	1/1	0.94	0.20	53,53,53,53	0
59	MG	DA	4085	1/1	0.94	0.21	60,60,60,60	0
59	MG	DB	242	1/1	0.94	0.31	61,61,61,61	0
59	MG	AA	3652	1/1	0.94	0.09	44,44,44,44	0
59	MG	CA	1871	1/1	0.94	0.14	98,98,98,98	0
59	MG	CA	2161	1/1	0.94	0.19	96,96,96,96	0
59	MG	AA	3834	1/1	0.94	0.19	79,79,79,79	0
59	MG	BA	1997	1/1	0.94	0.40	87,87,87,87	0
59	MG	DA	3777	1/1	0.94	0.29	107,107,107,107	0
59	MG	AA	3184	1/1	0.94	0.24	49,49,49,49	0
59	MG	BA	1918	1/1	0.94	0.07	74,74,74,74	0
59	MG	CA	1947	1/1	0.94	0.12	82,82,82,82	0
59	MG	DA	3328	1/1	0.94	0.12	56,56,56,56	0
59	MG	AA	3099	1/1	0.94	0.11	33,33,33,33	0
59	MG	DA	3861	1/1	0.94	0.19	77,77,77,77	0
59	MG	CA	2013	1/1	0.94	0.11	46,46,46,46	0
59	MG	BA	1850	1/1	0.94	0.13	90,90,90,90	0
59	MG	CA	1973	1/1	0.94	0.27	65,65,65,65	0
59	MG	AA	3842	1/1	0.94	0.13	67,67,67,67	0
59	MG	DA	4301	1/1	0.94	0.16	37,37,37,37	0
59	MG	BA	1630	1/1	0.94	0.21	78,78,78,78	0
59	MG	DA	4618	1/1	0.94	0.17	61,61,61,61	0
59	MG	DA	3837	1/1	0.94	0.28	61,61,61,61	0
59	MG	DA	3395	1/1	0.94	0.37	42,42,42,42	0
59	MG	DA	3939	1/1	0.94	0.12	47,47,47,47	0
59	MG	CA	1645	1/1	0.94	0.21	74,74,74,74	0
59	MG	DA	3635	1/1	0.94	0.19	33,33,33,33	0
59	MG	BA	2011	1/1	0.94	0.19	64,64,64,64	0
59	MG	DA	3849	1/1	0.94	0.12	63,63,63,63	0
59	MG	DA	3338	1/1	0.94	0.10	49,49,49,49	0
59	MG	DA	3429	1/1	0.94	0.30	30,30,30,30	0
59	MG	AA	4127	1/1	0.94	0.20	52,52,52,52	0
59	MG	BA	1685	1/1	0.94	0.13	45,45,45,45	0
59	MG	AA	3214	1/1	0.94	0.12	56,56,56,56	0
59	MG	BA	1790	1/1	0.94	0.09	56,56,56,56	0
59	MG	DA	3164	1/1	0.94	0.27	53,53,53,53	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	1903	1/1	0.94	0.14	56,56,56,56	0
59	MG	DA	4710	1/1	0.94	0.22	102,102,102,102	0
59	MG	DA	4170	1/1	0.94	0.32	46,46,46,46	0
59	MG	DA	4195	1/1	0.94	0.39	66,66,66,66	0
59	MG	AA	3619	1/1	0.94	0.08	64,64,64,64	0
59	MG	DA	3597	1/1	0.94	0.36	23,23,23,23	0
59	MG	DA	4845	1/1	0.94	0.60	43,43,43,43	0
59	MG	AA	3840	1/1	0.94	0.12	77,77,77,77	0
59	MG	D8	109	1/1	0.94	0.15	69,69,69,69	0
59	MG	DH	204	1/1	0.94	0.38	87,87,87,87	0
59	MG	AA	3339	1/1	0.94	0.13	50,50,50,50	0
59	MG	AA	3761	1/1	0.94	0.13	40,40,40,40	0
59	MG	BA	1868	1/1	0.94	0.07	65,65,65,65	0
59	MG	AD	306	1/1	0.94	0.12	44,44,44,44	0
59	MG	AA	3745	1/1	0.94	0.18	57,57,57,57	0
59	MG	CC	119	1/1	0.94	0.16	46,46,46,46	0
59	MG	DA	4067	1/1	0.94	0.09	39,39,39,39	0
59	MG	DA	4602	1/1	0.94	0.34	71,71,71,71	0
59	MG	AA	3311	1/1	0.94	0.30	49,49,49,49	0
59	MG	CA	2032	1/1	0.94	0.24	69,69,69,69	0
59	MG	AA	3727	1/1	0.94	0.22	54,54,54,54	0
59	MG	CA	2225	1/1	0.94	0.13	70,70,70,70	0
59	MG	CA	1930	1/1	0.94	0.12	42,42,42,42	0
59	MG	AA	3598	1/1	0.94	0.18	79,79,79,79	0
59	MG	DA	3926	1/1	0.94	0.35	49,49,49,49	0
59	MG	CA	2119	1/1	0.94	0.25	76,76,76,76	0
59	MG	CC	107	1/1	0.94	0.38	56,56,56,56	0
59	MG	DA	3953	1/1	0.94	0.16	84,84,84,84	0
59	MG	DA	3150	1/1	0.94	0.30	39,39,39,39	0
59	MG	BA	1845	1/1	0.94	0.11	92,92,92,92	0
59	MG	BS	107	1/1	0.94	0.11	90,90,90,90	0
59	MG	DE	301	1/1	0.94	0.18	19,19,19,19	0
59	MG	DA	4175	1/1	0.94	0.29	59,59,59,59	0
59	MG	DB	239	1/1	0.94	0.44	81,81,81,81	0
59	MG	AA	3035	1/1	0.94	0.07	17,17,17,17	0
59	MG	DA	4712	1/1	0.94	0.45	43,43,43,43	0
59	MG	CA	1932	1/1	0.94	0.23	51,51,51,51	0
59	MG	AA	3075	1/1	0.94	0.14	33,33,33,33	0
59	MG	DA	4334	1/1	0.94	0.46	70,70,70,70	0
59	MG	DA	4507	1/1	0.94	0.16	88,88,88,88	0
59	MG	DA	4489	1/1	0.94	0.20	42,42,42,42	0
59	MG	DA	3049	1/1	0.94	0.20	25,25,25,25	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3104	1/1	0.94	0.10	42,42,42,42	0
59	MG	DA	3855	1/1	0.94	0.11	85,85,85,85	0
59	MG	BA	1962	1/1	0.94	0.13	73,73,73,73	0
59	MG	AA	3457	1/1	0.94	0.11	57,57,57,57	0
59	MG	CA	2144	1/1	0.94	0.20	61,61,61,61	0
59	MG	BA	1941	1/1	0.94	0.11	58,58,58,58	0
59	MG	BA	1891	1/1	0.94	0.09	60,60,60,60	0
59	MG	DA	4116	1/1	0.94	0.36	78,78,78,78	0
59	MG	DA	3285	1/1	0.94	0.41	56,56,56,56	0
59	MG	AA	3666	1/1	0.94	0.20	52,52,52,52	0
59	MG	DA	3415	1/1	0.94	0.46	49,49,49,49	0
59	MG	DA	4586	1/1	0.94	0.24	78,78,78,78	0
59	MG	AA	3166	1/1	0.94	0.11	65,65,65,65	0
59	MG	DO	217	1/1	0.94	0.35	52,52,52,52	0
59	MG	DA	4178	1/1	0.94	0.40	61,61,61,61	0
59	MG	AA	3443	1/1	0.94	0.21	77,77,77,77	0
59	MG	DA	4755	1/1	0.94	0.25	80,80,80,80	0
59	MG	AA	3586	1/1	0.94	0.04	41,41,41,41	0
59	MG	DA	4429	1/1	0.94	0.15	55,55,55,55	0
59	MG	AA	4114	1/1	0.94	0.16	92,92,92,92	0
59	MG	AA	3687	1/1	0.94	0.17	43,43,43,43	0
59	MG	CA	1653	1/1	0.94	0.27	47,47,47,47	0
59	MG	BA	2159	1/1	0.94	0.19	89,89,89,89	0
59	MG	AA	3811	1/1	0.94	0.11	58,58,58,58	0
59	MG	CA	2026	1/1	0.94	0.07	69,69,69,69	0
59	MG	DA	4035	1/1	0.94	0.54	52,52,52,52	0
59	MG	DA	3088	1/1	0.94	0.27	12,12,12,12	0
59	MG	AA	3429	1/1	0.94	0.14	60,60,60,60	0
59	MG	DA	3199	1/1	0.94	0.31	29,29,29,29	0
59	MG	BA	1717	1/1	0.94	0.17	55,55,55,55	0
59	MG	AA	4065	1/1	0.94	0.18	68,68,68,68	0
59	MG	BA	2034	1/1	0.94	0.21	68,68,68,68	0
59	MG	AA	3155	1/1	0.94	0.16	46,46,46,46	0
59	MG	AA	3355	1/1	0.94	0.34	48,48,48,48	0
59	MG	AA	3930	1/1	0.94	0.66	46,46,46,46	0
59	MG	DA	3220	1/1	0.94	0.16	37,37,37,37	0
59	MG	CA	1952	1/1	0.94	0.14	94,94,94,94	0
59	MG	DA	3791	1/1	0.94	0.26	109,109,109,109	0
59	MG	DA	3312	1/1	0.94	0.34	49,49,49,49	0
59	MG	CA	2261	1/1	0.94	0.13	66,66,66,66	0
59	MG	DA	3250	1/1	0.94	0.19	13,13,13,13	0
59	MG	DB	241	1/1	0.94	0.17	66,66,66,66	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4120	1/1	0.94	0.11	48,48,48,48	0
59	MG	DA	4654	1/1	0.94	0.17	69,69,69,69	0
59	MG	DA	3304	1/1	0.94	0.13	43,43,43,43	0
59	MG	DA	4014	1/1	0.94	0.22	50,50,50,50	0
59	MG	CA	1618	1/1	0.94	0.12	24,24,24,24	0
59	MG	DA	3261	1/1	0.94	0.20	34,34,34,34	0
59	MG	AA	3515	1/1	0.94	0.15	20,20,20,20	0
59	MG	AA	3451	1/1	0.94	0.11	89,89,89,89	0
59	MG	DA	3194	1/1	0.94	0.21	57,57,57,57	0
59	MG	AA	3832	1/1	0.94	0.14	63,63,63,63	0
59	MG	DA	3802	1/1	0.94	0.11	88,88,88,88	0
59	MG	BA	1882	1/1	0.94	0.29	42,42,42,42	0
59	MG	DA	3055	1/1	0.94	0.41	21,21,21,21	0
59	MG	DA	4523	1/1	0.94	0.16	51,51,51,51	0
59	MG	AA	3860	1/1	0.94	0.09	83,83,83,83	0
59	MG	DA	3132	1/1	0.94	0.37	34,34,34,34	0
59	MG	DA	3717	1/1	0.94	0.10	1,1,1,1	0
59	MG	AA	3327	1/1	0.94	0.21	45,45,45,45	0
59	MG	BA	1643	1/1	0.94	0.37	53,53,53,53	0
59	MG	CA	2294	1/1	0.94	0.14	80,80,80,80	0
59	MG	DA	4437	1/1	0.94	0.25	27,27,27,27	0
59	MG	AA	3516	1/1	0.94	0.14	35,35,35,35	0
59	MG	CA	1706	1/1	0.94	0.19	41,41,41,41	0
59	MG	DA	3234	1/1	0.94	0.24	17,17,17,17	0
59	MG	DA	3848	1/1	0.94	0.45	35,35,35,35	0
59	MG	DA	3115	1/1	0.94	0.23	34,34,34,34	0
59	MG	CA	1884	1/1	0.94	0.14	64,64,64,64	0
59	MG	AA	3684	1/1	0.94	0.17	58,58,58,58	0
59	MG	AA	3256	1/1	0.94	0.21	62,62,62,62	0
59	MG	DO	208	1/1	0.94	0.23	59,59,59,59	0
59	MG	DA	3942	1/1	0.94	0.17	64,64,64,64	0
59	MG	DU	212	1/1	0.94	0.73	75,75,75,75	0
59	MG	DZ	105	1/1	0.94	0.40	54,54,54,54	0
59	MG	DA	4126	1/1	0.94	0.25	57,57,57,57	0
59	MG	DA	3571	1/1	0.94	0.29	93,93,93,93	0
59	MG	DF	317	1/1	0.94	0.09	64,64,64,64	0
59	MG	AA	3615	1/1	0.94	0.08	47,47,47,47	0
59	MG	DA	3491	1/1	0.94	0.13	78,78,78,78	0
59	MG	AA	3702	1/1	0.94	0.17	62,62,62,62	0
59	MG	AA	3400	1/1	0.94	0.21	58,58,58,58	0
59	MG	CA	1820	1/1	0.94	0.08	67,67,67,67	0
59	MG	DA	3018	1/1	0.94	0.24	20,20,20,20	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4137	1/1	0.94	0.21	39,39,39,39	0
59	MG	DA	3081	1/1	0.94	0.34	24,24,24,24	0
59	MG	DA	4318	1/1	0.94	0.14	50,50,50,50	0
59	MG	DO	205	1/1	0.94	0.30	52,52,52,52	0
59	MG	BA	1733	1/1	0.94	0.07	46,46,46,46	0
59	MG	DA	3803	1/1	0.94	0.14	54,54,54,54	0
59	MG	DA	3756	1/1	0.94	0.29	126,126,126,126	0
59	MG	D1	209	1/1	0.94	0.27	65,65,65,65	0
59	MG	DA	4206	1/1	0.94	0.16	45,45,45,45	0
59	MG	DF	322	1/1	0.94	0.81	40,40,40,40	0
59	MG	DA	3550	1/1	0.94	0.23	96,96,96,96	0
59	MG	CA	1723	1/1	0.94	0.19	61,61,61,61	0
59	MG	CA	1931	1/1	0.94	0.23	64,64,64,64	0
59	MG	CA	1641	1/1	0.94	0.17	28,28,28,28	0
59	MG	BA	1670	1/1	0.94	0.11	50,50,50,50	0
59	MG	BH	203	1/1	0.94	0.13	89,89,89,89	0
59	MG	AA	3279	1/1	0.94	0.17	45,45,45,45	0
59	MG	DA	4736	1/1	0.94	0.17	77,77,77,77	0
59	MG	AA	3847	1/1	0.94	0.17	84,84,84,84	0
59	MG	BA	1901	1/1	0.94	0.08	82,82,82,82	0
59	MG	AA	3495	1/1	0.94	0.27	56,56,56,56	0
59	MG	DA	4970	1/1	0.94	0.35	87,87,87,87	0
59	MG	DA	3772	1/1	0.94	0.16	65,65,65,65	0
59	MG	CA	1669	1/1	0.94	0.21	40,40,40,40	0
59	MG	DA	3048	1/1	0.94	0.14	15,15,15,15	0
59	MG	CA	1674	1/1	0.94	0.37	60,60,60,60	0
59	MG	DA	3337	1/1	0.94	0.27	44,44,44,44	0
59	MG	DA	4171	1/1	0.94	0.14	45,45,45,45	0
59	MG	DA	4238	1/1	0.94	0.22	52,52,52,52	0
59	MG	CA	1740	1/1	0.94	0.27	48,48,48,48	0
59	MG	CA	1991	1/1	0.94	0.12	76,76,76,76	0
59	MG	DA	4423	1/1	0.94	0.29	73,73,73,73	0
59	MG	BA	1989	1/1	0.94	0.04	75,75,75,75	0
59	MG	AA	3735	1/1	0.94	0.07	50,50,50,50	0
59	MG	CA	1649	1/1	0.94	0.08	47,47,47,47	0
59	MG	AA	3372	1/1	0.94	0.15	52,52,52,52	0
59	MG	DA	4353	1/1	0.94	0.48	45,45,45,45	0
59	MG	CA	1993	1/1	0.94	0.08	91,91,91,91	0
59	MG	DA	3608	1/1	0.94	0.27	60,60,60,60	0
59	MG	DW	106	1/1	0.94	0.19	61,61,61,61	0
59	MG	DA	4344	1/1	0.94	0.35	63,63,63,63	0
59	MG	DB	220	1/1	0.94	0.10	79,79,79,79	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	2176	1/1	0.94	0.06	62,62,62,62	0
59	MG	BP	201	1/1	0.94	0.12	61,61,61,61	0
59	MG	DA	5024	1/1	0.94	0.15	43,43,43,43	0
59	MG	D3	101	1/1	0.94	0.31	33,33,33,33	0
59	MG	AA	4148	1/1	0.94	0.28	87,87,87,87	0
59	MG	DA	4018	1/1	0.94	0.17	75,75,75,75	0
59	MG	DW	104	1/1	0.94	0.16	49,49,49,49	0
59	MG	CA	1768	1/1	0.94	0.14	48,48,48,48	0
59	MG	CA	1862	1/1	0.94	0.10	122,122,122,122	0
59	MG	DA	4235	1/1	0.94	0.13	44,44,44,44	0
59	MG	AA	3486	1/1	0.94	0.19	60,60,60,60	0
59	MG	DA	3103	1/1	0.94	0.38	49,49,49,49	0
59	MG	DA	4012	1/1	0.94	0.35	63,63,63,63	0
59	MG	DA	3681	1/1	0.94	0.08	73,73,73,73	0
59	MG	DA	4988	1/1	0.94	0.13	64,64,64,64	0
59	MG	CA	2082	1/1	0.94	0.09	63,63,63,63	0
59	MG	DA	3818	1/1	0.94	0.11	71,71,71,71	0
59	MG	AA	4122	1/1	0.94	0.08	83,83,83,83	0
59	MG	DA	4585	1/1	0.94	0.17	76,76,76,76	0
59	MG	DA	4071	1/1	0.94	0.25	55,55,55,55	0
59	MG	DA	3349	1/1	0.94	0.31	56,56,56,56	0
59	MG	AA	3754	1/1	0.94	0.13	73,73,73,73	0
59	MG	BA	2098	1/1	0.94	0.29	73,73,73,73	0
59	MG	AA	3473	1/1	0.94	0.13	63,63,63,63	0
59	MG	AA	4092	1/1	0.94	0.16	90,90,90,90	0
59	MG	AA	3785	1/1	0.94	0.27	89,89,89,89	0
59	MG	DA	3430	1/1	0.94	0.19	60,60,60,60	0
59	MG	AA	3301	1/1	0.94	0.17	50,50,50,50	0
59	MG	CA	1749	1/1	0.94	0.13	68,68,68,68	0
59	MG	BA	2185	1/1	0.94	0.41	81,81,81,81	0
59	MG	AA	3681	1/1	0.94	0.07	65,65,65,65	0
59	MG	BA	2204	1/1	0.94	0.19	70,70,70,70	0
59	MG	DO	210	1/1	0.94	0.34	53,53,53,53	0
59	MG	DA	3090	1/1	0.94	0.15	13,13,13,13	0
59	MG	DA	3459	1/1	0.94	0.08	63,63,63,63	0
59	MG	DA	4415	1/1	0.94	0.14	74,74,74,74	0
59	MG	DA	4107	1/1	0.94	0.15	49,49,49,49	0
59	MG	DA	4096	1/1	0.94	0.38	73,73,73,73	0
59	MG	CS	101	1/1	0.94	0.09	61,61,61,61	0
59	MG	DA	3198	1/1	0.94	0.17	21,21,21,21	0
59	MG	DA	4181	1/1	0.94	0.20	81,81,81,81	0
59	MG	AA	3168	1/1	0.94	0.26	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4703	1/1	0.94	0.30	49,49,49,49	0
59	MG	DA	3313	1/1	0.94	0.26	48,48,48,48	0
59	MG	CA	1621	1/1	0.94	0.16	45,45,45,45	0
59	MG	CA	2315	1/1	0.94	0.25	94,94,94,94	0
59	MG	BA	1711	1/1	0.94	0.22	68,68,68,68	0
59	MG	BA	1705	1/1	0.94	0.41	52,52,52,52	0
59	MG	BA	2263	1/1	0.94	0.23	71,71,71,71	0
59	MG	DA	4086	1/1	0.94	0.23	98,98,98,98	0
59	MG	DB	250	1/1	0.94	0.32	74,74,74,74	0
59	MG	DA	3527	1/1	0.94	0.12	46,46,46,46	0
59	MG	AA	4157	1/1	0.94	0.10	76,76,76,76	0
59	MG	DA	3648	1/1	0.94	0.17	51,51,51,51	0
59	MG	DA	4278	1/1	0.94	0.34	62,62,62,62	0
59	MG	BA	2240	1/1	0.94	0.07	86,86,86,86	0
59	MG	AA	4057	1/1	0.94	0.20	74,74,74,74	0
59	MG	CC	122	1/1	0.94	0.23	69,69,69,69	0
59	MG	DA	3301	1/1	0.94	0.11	29,29,29,29	0
59	MG	DA	3139	1/1	0.94	0.28	45,45,45,45	0
59	MG	AA	4158	1/1	0.94	0.14	93,93,93,93	0
59	MG	DA	3288	1/1	0.94	0.13	52,52,52,52	0
59	MG	CA	1989	1/1	0.94	0.22	60,60,60,60	0
59	MG	DA	4659	1/1	0.94	0.18	68,68,68,68	0
59	MG	DA	4356	1/1	0.94	0.44	76,76,76,76	0
59	MG	DA	3321	1/1	0.94	0.29	34,34,34,34	0
59	MG	CP	203	1/1	0.94	0.23	77,77,77,77	0
59	MG	DA	3515	1/1	0.94	0.11	28,28,28,28	0
59	MG	CB	120	1/1	0.94	0.38	100,100,100,100	0
59	MG	CQ	103	1/1	0.94	0.15	51,51,51,51	0
59	MG	AA	3704	1/1	0.94	0.13	59,59,59,59	0
59	MG	DA	3469	1/1	0.94	0.28	63,63,63,63	0
59	MG	DY	203	1/1	0.94	0.18	75,75,75,75	0
59	MG	DA	3227	1/1	0.94	0.44	52,52,52,52	0
59	MG	DA	3287	1/1	0.94	0.23	78,78,78,78	0
59	MG	DA	4428	1/1	0.94	0.17	64,64,64,64	0
59	MG	DA	3400	1/1	0.94	0.24	48,48,48,48	0
59	MG	DA	4619	1/1	0.94	0.17	49,49,49,49	0
59	MG	DA	3593	1/1	0.94	0.28	29,29,29,29	0
59	MG	AA	3656	1/1	0.94	0.17	37,37,37,37	0
59	MG	BU	1601	1/1	0.94	0.09	56,56,56,56	0
59	MG	CA	2196	1/1	0.94	0.12	62,62,62,62	0
59	MG	BA	2177	1/1	0.94	0.25	67,67,67,67	0
59	MG	BA	2155	1/1	0.94	0.15	71,71,71,71	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	1815	1/1	0.94	0.09	66,66,66,66	0
59	MG	BD	104	1/1	0.94	0.25	51,51,51,51	0
59	MG	DA	3209	1/1	0.95	0.25	33,33,33,33	0
59	MG	CA	2131	1/1	0.95	0.20	61,61,61,61	0
59	MG	AA	4125	1/1	0.95	0.10	69,69,69,69	0
59	MG	CB	119	1/1	0.95	0.03	69,69,69,69	0
59	MG	DA	3540	1/1	0.95	0.22	74,74,74,74	0
59	MG	AA	3104	1/1	0.95	0.29	54,54,54,54	0
59	MG	DA	3928	1/1	0.95	0.07	49,49,49,49	0
59	MG	DA	4932	1/1	0.95	0.21	69,69,69,69	0
59	MG	DA	4033	1/1	0.95	0.22	103,103,103,103	0
59	MG	BA	1602	1/1	0.95	0.20	20,20,20,20	0
59	MG	BC	109	1/1	0.95	0.09	72,72,72,72	0
59	MG	DA	4579	1/1	0.95	0.16	81,81,81,81	0
59	MG	DA	4145	1/1	0.95	0.21	44,44,44,44	0
59	MG	AA	3992	1/1	0.95	0.14	75,75,75,75	0
59	MG	DA	4556	1/1	0.95	0.16	41,41,41,41	0
59	MG	AA	3156	1/1	0.95	0.20	55,55,55,55	0
59	MG	AA	3535	1/1	0.95	0.56	58,58,58,58	0
59	MG	BA	1687	1/1	0.95	0.06	41,41,41,41	0
59	MG	DA	3175	1/1	0.95	0.20	21,21,21,21	0
59	MG	DA	4288	1/1	0.95	0.22	72,72,72,72	0
59	MG	DA	4389	1/1	0.95	0.35	60,60,60,60	0
59	MG	DA	3806	1/1	0.95	0.25	55,55,55,55	0
59	MG	CA	1945	1/1	0.95	0.14	78,78,78,78	0
59	MG	DB	232	1/1	0.95	0.38	54,54,54,54	0
59	MG	DA	4306	1/1	0.95	0.23	45,45,45,45	0
59	MG	DO	203	1/1	0.95	0.35	32,32,32,32	0
59	MG	DA	3467	1/1	0.95	0.45	75,75,75,75	0
59	MG	DA	4011	1/1	0.95	0.17	52,52,52,52	0
59	MG	DA	3197	1/1	0.95	0.10	19,19,19,19	0
59	MG	DA	3927	1/1	0.95	0.14	66,66,66,66	0
59	MG	AA	3086	1/1	0.95	0.23	35,35,35,35	0
59	MG	DA	3779	1/1	0.95	0.35	155,155,155,155	0
59	MG	AA	3216	1/1	0.95	0.21	40,40,40,40	0
59	MG	CA	2141	1/1	0.95	0.17	60,60,60,60	0
59	MG	AA	3511	1/1	0.95	0.15	26,26,26,26	0
59	MG	DA	3073	1/1	0.95	0.21	16,16,16,16	0
59	MG	DA	4087	1/1	0.95	0.18	45,45,45,45	0
59	MG	DA	4722	1/1	0.95	0.15	109,109,109,109	0
59	MG	DA	4232	1/1	0.95	0.33	68,68,68,68	0
59	MG	DA	4687	1/1	0.95	0.17	76,76,76,76	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	1824	1/1	0.95	0.15	102,102,102,102	0
59	MG	BA	2016	1/1	0.95	0.10	78,78,78,78	0
59	MG	AD	309	1/1	0.95	0.35	48,48,48,48	0
59	MG	DA	3720	1/1	0.95	0.21	46,46,46,46	0
59	MG	DA	3752	1/1	0.95	0.11	103,103,103,103	0
59	MG	CA	1837	1/1	0.95	0.15	79,79,79,79	0
59	MG	DB	249	1/1	0.95	0.24	84,84,84,84	0
59	MG	DA	4953	1/1	0.95	0.12	63,63,63,63	0
59	MG	BA	1977	1/1	0.95	0.06	85,85,85,85	0
59	MG	DA	3934	1/1	0.95	0.08	45,45,45,45	0
59	MG	AA	3939	1/1	0.95	0.18	54,54,54,54	0
59	MG	BA	1883	1/1	0.95	0.42	64,64,64,64	0
59	MG	AA	4164	1/1	0.95	0.16	80,80,80,80	0
59	MG	DA	4567	1/1	0.95	0.11	51,51,51,51	0
59	MG	DA	4361	1/1	0.95	0.37	47,47,47,47	0
59	MG	DA	4094	1/1	0.95	0.15	41,41,41,41	0
59	MG	AD	305	1/1	0.95	0.25	53,53,53,53	0
59	MG	AA	3062	1/1	0.95	0.09	29,29,29,29	0
59	MG	DA	4287	1/1	0.95	0.20	73,73,73,73	0
59	MG	BA	1666	1/1	0.95	0.12	43,43,43,43	0
59	MG	BA	2147	1/1	0.95	0.06	107,107,107,107	0
59	MG	DA	4950	1/1	0.95	0.28	125,125,125,125	0
59	MG	DA	3502	1/1	0.95	0.58	74,74,74,74	0
59	MG	AA	3551	1/1	0.95	0.35	44,44,44,44	0
59	MG	DA	3003	1/1	0.95	0.20	11,11,11,11	0
59	MG	DA	3099	1/1	0.95	0.28	48,48,48,48	0
59	MG	AA	3774	1/1	0.95	0.18	79,79,79,79	0
59	MG	DA	3708	1/1	0.95	0.25	9,9,9,9	0
59	MG	AA	3310	1/1	0.95	0.19	47,47,47,47	0
59	MG	BA	1781	1/1	0.95	0.15	69,69,69,69	0
59	MG	DA	4339	1/1	0.95	0.27	71,71,71,71	0
59	MG	DB	259	1/1	0.95	0.35	69,69,69,69	0
59	MG	AD	304	1/1	0.95	0.08	65,65,65,65	0
59	MG	BA	1613	1/1	0.95	0.45	84,84,84,84	0
59	MG	CD	113	1/1	0.95	0.09	75,75,75,75	0
59	MG	BA	1751	1/1	0.95	0.15	62,62,62,62	0
59	MG	AA	3040	1/1	0.95	0.21	37,37,37,37	0
59	MG	DF	324	1/1	0.95	0.25	76,76,76,76	0
59	MG	DA	3010	1/1	0.95	0.15	10,10,10,10	0
59	MG	DA	3771	1/1	0.95	0.09	92,92,92,92	0
59	MG	BA	1760	1/1	0.95	0.23	85,85,85,85	0
59	MG	DA	3159	1/1	0.95	0.15	47,47,47,47	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3688	1/1	0.95	0.31	59,59,59,59	0
59	MG	BA	1697	1/1	0.95	0.43	52,52,52,52	0
59	MG	DA	3449	1/1	0.95	0.22	50,50,50,50	0
59	MG	DA	3140	1/1	0.95	0.29	26,26,26,26	0
59	MG	AA	3082	1/1	0.95	0.15	48,48,48,48	0
59	MG	DA	3351	1/1	0.95	0.22	39,39,39,39	0
59	MG	AA	3150	1/1	0.95	0.13	29,29,29,29	0
59	MG	DA	4358	1/1	0.95	0.15	56,56,56,56	0
59	MG	CA	1739	1/1	0.95	0.10	43,43,43,43	0
59	MG	BA	1954	1/1	0.95	0.19	104,104,104,104	0
59	MG	DA	4747	1/1	0.95	1.00	71,71,71,71	0
59	MG	AA	3838	1/1	0.95	0.10	102,102,102,102	0
59	MG	DA	3877	1/1	0.95	0.12	37,37,37,37	0
59	MG	DF	306	1/1	0.95	0.31	46,46,46,46	0
59	MG	DA	3808	1/1	0.95	0.12	77,77,77,77	0
59	MG	CA	2202	1/1	0.95	0.28	47,47,47,47	0
59	MG	DA	3554	1/1	0.95	0.32	51,51,51,51	0
59	MG	AA	3343	1/1	0.95	0.29	48,48,48,48	0
59	MG	AA	3635	1/1	0.95	0.13	54,54,54,54	0
59	MG	DA	3433	1/1	0.95	0.23	43,43,43,43	0
59	MG	DA	4500	1/1	0.95	0.12	53,53,53,53	0
59	MG	BA	1601	1/1	0.95	0.20	27,27,27,27	0
59	MG	BA	2024	1/1	0.95	0.10	111,111,111,111	0
59	MG	AA	3875	1/1	0.95	0.06	67,67,67,67	0
59	MG	CA	1939	1/1	0.95	0.06	81,81,81,81	0
59	MG	AA	3488	1/1	0.95	0.15	68,68,68,68	0
59	MG	BA	1708	1/1	0.95	0.25	57,57,57,57	0
59	MG	DA	4453	1/1	0.95	0.24	49,49,49,49	0
59	MG	BA	1881	1/1	0.95	0.24	46,46,46,46	0
59	MG	DA	3141	1/1	0.95	0.24	42,42,42,42	0
59	MG	AA	3580	1/1	0.95	0.08	50,50,50,50	0
59	MG	AA	3072	1/1	0.95	0.24	21,21,21,21	0
59	MG	DA	4475	1/1	0.95	0.17	50,50,50,50	0
59	MG	DA	3280	1/1	0.95	0.12	32,32,32,32	0
59	MG	AA	3711	1/1	0.95	0.26	69,69,69,69	0
59	MG	DA	3171	1/1	0.95	0.28	36,36,36,36	0
59	MG	CA	2055	1/1	0.95	0.29	60,60,60,60	0
59	MG	DA	4010	1/1	0.95	0.12	91,91,91,91	0
59	MG	DA	4548	1/1	0.95	0.12	53,53,53,53	0
59	MG	CA	1673	1/1	0.95	0.20	97,97,97,97	0
59	MG	CA	1614	1/1	0.95	0.21	41,41,41,41	0
59	MG	AA	3493	1/1	0.95	0.07	76,76,76,76	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DW	102	1/1	0.95	0.19	37,37,37,37	0
59	MG	DA	3969	1/1	0.95	0.12	48,48,48,48	0
59	MG	DA	4072	1/1	0.95	0.11	48,48,48,48	0
59	MG	DA	3146	1/1	0.95	0.22	55,55,55,55	0
59	MG	AA	3843	1/1	0.95	0.18	81,81,81,81	0
59	MG	AA	3734	1/1	0.95	0.25	49,49,49,49	0
59	MG	DA	3093	1/1	0.95	0.20	13,13,13,13	0
59	MG	AA	3094	1/1	0.95	0.14	21,21,21,21	0
59	MG	DA	3334	1/1	0.95	0.30	40,40,40,40	0
59	MG	DA	3112	1/1	0.95	0.31	49,49,49,49	0
59	MG	AA	3906	1/1	0.95	0.30	67,67,67,67	0
59	MG	CA	1681	1/1	0.95	0.18	24,24,24,24	0
59	MG	DA	3210	1/1	0.95	0.21	32,32,32,32	0
59	MG	DA	4249	1/1	0.95	0.29	55,55,55,55	0
59	MG	AA	3585	1/1	0.95	0.15	45,45,45,45	0
59	MG	DA	4425	1/1	0.95	0.27	109,109,109,109	0
59	MG	DO	202	1/1	0.95	0.22	35,35,35,35	0
59	MG	AA	3468	1/1	0.95	0.09	22,22,22,22	0
59	MG	CA	1620	1/1	0.95	0.16	39,39,39,39	0
59	MG	DA	3233	1/1	0.95	0.13	26,26,26,26	0
59	MG	AA	3176	1/1	0.95	0.20	33,33,33,33	0
59	MG	CA	1696	1/1	0.95	0.18	33,33,33,33	0
59	MG	DA	3308	1/1	0.95	0.50	75,75,75,75	0
59	MG	CA	2319	1/1	0.95	0.28	60,60,60,60	0
59	MG	DA	4032	1/1	0.95	0.21	86,86,86,86	0
59	MG	BA	1693	1/1	0.95	0.31	46,46,46,46	0
59	MG	DA	4467	1/1	0.95	0.19	50,50,50,50	0
59	MG	AA	4031	1/1	0.95	0.10	84,84,84,84	0
59	MG	DA	3144	1/1	0.95	0.20	36,36,36,36	0
59	MG	DA	3952	1/1	0.95	0.12	46,46,46,46	0
59	MG	BA	1763	1/1	0.95	0.25	57,57,57,57	0
59	MG	AA	3045	1/1	0.95	0.29	51,51,51,51	0
59	MG	BA	1908	1/1	0.95	0.30	79,79,79,79	0
59	MG	DA	3134	1/1	0.95	0.43	44,44,44,44	0
59	MG	AA	3352	1/1	0.95	0.10	28,28,28,28	0
59	MG	DA	3116	1/1	0.95	0.23	63,63,63,63	0
59	MG	DU	204	1/1	0.95	0.10	62,62,62,62	0
59	MG	AA	3909	1/1	0.95	0.42	90,90,90,90	0
59	MG	DA	4380	1/1	0.95	0.20	61,61,61,61	0
59	MG	BA	1672	1/1	0.95	0.29	39,39,39,39	0
59	MG	DA	3329	1/1	0.95	0.26	57,57,57,57	0
59	MG	DA	3357	1/1	0.95	0.15	59,59,59,59	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3202	1/1	0.95	0.20	31,31,31,31	0
59	MG	DA	3217	1/1	0.95	0.20	28,28,28,28	0
59	MG	AA	3167	1/1	0.95	0.16	61,61,61,61	0
59	MG	DA	3929	1/1	0.95	0.07	73,73,73,73	0
59	MG	DA	3350	1/1	0.95	0.14	50,50,50,50	0
59	MG	BA	1844	1/1	0.95	0.17	118,118,118,118	0
59	MG	AA	3030	1/1	0.95	0.19	45,45,45,45	0
59	MG	DA	4939	1/1	0.95	0.50	57,57,57,57	0
59	MG	DA	3335	1/1	0.95	0.15	32,32,32,32	0
59	MG	AA	3043	1/1	0.95	0.04	42,42,42,42	0
59	MG	D3	105	1/1	0.95	0.32	48,48,48,48	0
59	MG	DD	304	1/1	0.95	0.88	66,66,66,66	0
59	MG	D4	102	1/1	0.95	0.19	86,86,86,86	0
59	MG	CA	1801	1/1	0.95	0.12	44,44,44,44	0
59	MG	DA	3399	1/1	0.95	0.22	52,52,52,52	0
59	MG	DA	3177	1/1	0.95	0.24	39,39,39,39	0
59	MG	CH	204	1/1	0.95	0.27	93,93,93,93	0
59	MG	AA	3395	1/1	0.95	0.29	88,88,88,88	0
59	MG	BA	2260	1/1	0.95	0.18	77,77,77,77	0
59	MG	DA	3897	1/1	0.95	0.18	27,27,27,27	0
59	MG	CA	1664	1/1	0.95	0.11	22,22,22,22	0
59	MG	DA	3460	1/1	0.95	0.25	46,46,46,46	0
59	MG	CA	2054	1/1	0.95	0.08	62,62,62,62	0
59	MG	CA	1707	1/1	0.95	0.40	54,54,54,54	0
59	MG	DA	3438	1/1	0.95	0.18	32,32,32,32	0
59	MG	BA	1691	1/1	0.95	0.04	65,65,65,65	0
59	MG	DA	4684	1/1	0.95	0.11	36,36,36,36	0
59	MG	DA	3386	1/1	0.95	0.40	69,69,69,69	0
59	MG	AA	3055	1/1	0.95	0.17	33,33,33,33	0
59	MG	CA	1908	1/1	0.95	0.26	61,61,61,61	0
59	MG	DA	3841	1/1	0.95	0.13	50,50,50,50	0
59	MG	CA	1605	1/1	0.95	0.19	36,36,36,36	0
59	MG	BA	1818	1/1	0.95	0.32	57,57,57,57	0
59	MG	DA	4891	1/1	0.95	0.32	86,86,86,86	0
59	MG	DA	3601	1/1	0.95	0.12	1,1,1,1	0
59	MG	DA	4064	1/1	0.95	0.12	56,56,56,56	0
59	MG	AA	3494	1/1	0.95	0.10	69,69,69,69	0
59	MG	BA	1879	1/1	0.95	0.43	46,46,46,46	0
59	MG	CA	2174	1/1	0.95	0.14	100,100,100,100	0
59	MG	DA	3265	1/1	0.95	0.34	47,47,47,47	0
59	MG	DA	3416	1/1	0.95	0.17	62,62,62,62	0
59	MG	D2	201	1/1	0.95	0.18	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3076	1/1	0.95	0.11	35,35,35,35	0
59	MG	DA	4704	1/1	0.95	0.26	62,62,62,62	0
59	MG	DA	3961	1/1	0.95	0.15	41,41,41,41	0
59	MG	AA	3375	1/1	0.95	0.32	61,61,61,61	0
59	MG	DR	203	1/1	0.95	0.20	84,84,84,84	0
59	MG	BA	1871	1/1	0.95	0.03	74,74,74,74	0
59	MG	CS	103	1/1	0.95	0.10	74,74,74,74	0
59	MG	BA	1884	1/1	0.95	0.46	67,67,67,67	0
59	MG	BA	1609	1/1	0.95	0.16	36,36,36,36	0
59	MG	AA	3131	1/1	0.95	0.15	34,34,34,34	0
59	MG	AA	3031	1/1	0.95	0.06	34,34,34,34	0
59	MG	AA	3945	1/1	0.95	0.12	108,108,108,108	0
59	MG	BA	2078	1/1	0.95	0.18	79,79,79,79	0
59	MG	DM	205	1/1	0.95	1.02	51,51,51,51	0
59	MG	CA	1900	1/1	0.95	0.09	74,74,74,74	0
59	MG	BC	105	1/1	0.95	0.07	75,75,75,75	0
59	MG	AA	4108	1/1	0.95	0.13	136,136,136,136	0
59	MG	CA	1969	1/1	0.95	0.14	85,85,85,85	0
59	MG	DA	3282	1/1	0.95	0.08	12,12,12,12	0
59	MG	AA	3259	1/1	0.95	0.44	57,57,57,57	0
59	MG	CA	1667	1/1	0.95	0.25	50,50,50,50	0
59	MG	AA	3899	1/1	0.95	0.20	46,46,46,46	0
59	MG	DA	3868	1/1	0.95	0.28	72,72,72,72	0
59	MG	DA	4673	1/1	0.95	0.45	91,91,91,91	0
59	MG	BV	101	1/1	0.95	0.12	73,73,73,73	0
59	MG	DA	3916	1/1	0.95	0.34	36,36,36,36	0
59	MG	AA	3646	1/1	0.95	0.11	62,62,62,62	0
59	MG	AA	3691	1/1	0.95	0.18	79,79,79,79	0
59	MG	DA	3578	1/1	0.95	0.09	62,62,62,62	0
59	MG	DA	3718	1/1	0.95	0.08	25,25,25,25	0
59	MG	CA	1831	1/1	0.95	0.13	132,132,132,132	0
59	MG	DA	4152	1/1	0.95	0.24	54,54,54,54	0
59	MG	AA	3079	1/1	0.95	0.09	40,40,40,40	0
59	MG	BC	113	1/1	0.95	0.07	54,54,54,54	0
59	MG	DA	3719	1/1	0.95	0.25	29,29,29,29	0
59	MG	DA	3764	1/1	0.95	0.19	98,98,98,98	0
59	MG	AA	3706	1/1	0.95	0.21	44,44,44,44	0
59	MG	DA	3179	1/1	0.95	0.26	52,52,52,52	0
59	MG	DA	4607	1/1	0.95	0.16	64,64,64,64	0
59	MG	DA	3216	1/1	0.95	0.37	54,54,54,54	0
59	MG	AA	3648	1/1	0.95	0.16	43,43,43,43	0
59	MG	DA	4348	1/1	0.95	0.33	62,62,62,62	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	1826	1/1	0.95	0.05	83,83,83,83	0
59	MG	AA	3617	1/1	0.95	0.06	59,59,59,59	0
59	MG	AA	3319	1/1	0.95	0.14	85,85,85,85	0
59	MG	AA	3780	1/1	0.95	0.28	69,69,69,69	0
59	MG	AA	3469	1/1	0.95	0.22	42,42,42,42	0
59	MG	BA	2100	1/1	0.95	0.21	69,69,69,69	0
59	MG	DA	4101	1/1	0.95	0.24	82,82,82,82	0
59	MG	DA	4674	1/1	0.95	0.30	93,93,93,93	0
59	MG	AA	3874	1/1	0.95	0.12	90,90,90,90	0
59	MG	DA	3921	1/1	0.95	0.21	24,24,24,24	0
59	MG	DD	312	1/1	0.95	0.16	37,37,37,37	0
59	MG	CA	2231	1/1	0.95	0.18	95,95,95,95	0
59	MG	DA	3783	1/1	0.95	0.06	63,63,63,63	0
59	MG	DA	3664	1/1	0.95	0.24	60,60,60,60	0
59	MG	CA	2044	1/1	0.95	0.11	65,65,65,65	0
59	MG	DA	4078	1/1	0.95	0.21	65,65,65,65	0
59	MG	DA	4563	1/1	0.95	0.31	77,77,77,77	0
59	MG	CA	1781	1/1	0.95	0.16	45,45,45,45	0
59	MG	CA	1769	1/1	0.95	0.16	42,42,42,42	0
59	MG	DA	3305	1/1	0.95	0.18	21,21,21,21	0
59	MG	DA	4239	1/1	0.95	0.13	62,62,62,62	0
59	MG	DA	4286	1/1	0.95	0.24	49,49,49,49	0
59	MG	DA	4257	1/1	0.95	0.20	59,59,59,59	0
59	MG	DA	3710	1/1	0.95	0.28	14,14,14,14	0
59	MG	DA	3796	1/1	0.95	0.17	26,26,26,26	0
59	MG	BA	1944	1/1	0.95	0.37	71,71,71,71	0
59	MG	DA	3712	1/1	0.95	0.20	25,25,25,25	0
59	MG	DA	4056	1/1	0.95	0.44	62,62,62,62	0
59	MG	DA	5045	1/1	0.95	0.24	72,72,72,72	0
59	MG	DA	4003	1/1	0.95	0.38	39,39,39,39	0
59	MG	AA	3121	1/1	0.95	0.30	85,85,85,85	0
59	MG	AA	3783	1/1	0.95	0.22	86,86,86,86	0
59	MG	CD	105	1/1	0.95	0.41	62,62,62,62	0
59	MG	DA	3977	1/1	0.95	0.09	71,71,71,71	0
59	MG	DA	4470	1/1	0.95	0.16	41,41,41,41	0
59	MG	AO	201	1/1	0.95	0.16	50,50,50,50	0
59	MG	CA	2195	1/1	0.95	0.13	76,76,76,76	0
59	MG	DA	4632	1/1	0.95	0.15	71,71,71,71	0
59	MG	AA	3455	1/1	0.95	0.09	22,22,22,22	0
59	MG	BA	2048	1/1	0.95	0.28	95,95,95,95	0
59	MG	CA	1721	1/1	0.95	0.14	39,39,39,39	0
59	MG	AA	3304	1/1	0.95	0.14	36,36,36,36	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DE	315	1/1	0.95	0.37	77,77,77,77	0
59	MG	CA	1883	1/1	0.95	0.07	43,43,43,43	0
59	MG	BA	1651	1/1	0.95	0.27	72,72,72,72	0
59	MG	AA	3730	1/1	0.95	0.09	58,58,58,58	0
59	MG	BA	1608	1/1	0.95	0.12	47,47,47,47	0
59	MG	BA	2061	1/1	0.95	0.32	70,70,70,70	0
59	MG	BA	1746	1/1	0.95	0.10	40,40,40,40	0
59	MG	AA	4085	1/1	0.95	0.07	60,60,60,60	0
59	MG	DA	3388	1/1	0.95	0.12	33,33,33,33	0
59	MG	D5	103	1/1	0.95	0.19	32,32,32,32	0
59	MG	BA	2252	1/1	0.95	0.08	70,70,70,70	0
59	MG	DA	4546	1/1	0.95	0.27	67,67,67,67	0
59	MG	DA	4664	1/1	0.95	0.14	66,66,66,66	0
59	MG	DE	305	1/1	0.95	0.10	44,44,44,44	0
59	MG	DA	3946	1/1	0.95	0.13	82,82,82,82	0
59	MG	DA	3532	1/1	0.95	0.33	49,49,49,49	0
59	MG	DQ	202	1/1	0.95	0.14	42,42,42,42	0
59	MG	CA	1967	1/1	0.95	0.21	56,56,56,56	0
59	MG	CA	2163	1/1	0.95	0.19	61,61,61,61	0
59	MG	DA	4628	1/1	0.95	0.17	94,94,94,94	0
59	MG	CA	2003	1/1	0.95	0.24	77,77,77,77	0
59	MG	CA	2168	1/1	0.95	0.23	108,108,108,108	0
59	MG	BA	1847	1/1	0.95	0.09	74,74,74,74	0
59	MG	DA	4226	1/1	0.95	0.20	45,45,45,45	0
59	MG	CL	201	1/1	0.95	0.09	54,54,54,54	0
59	MG	DA	4393	1/1	0.95	0.18	53,53,53,53	0
59	MG	DA	3707	1/1	0.95	0.11	28,28,28,28	0
59	MG	DA	3902	1/1	0.95	0.10	64,64,64,64	0
59	MG	CA	1957	1/1	0.95	0.31	56,56,56,56	0
59	MG	CA	1689	1/1	0.95	0.14	58,58,58,58	0
59	MG	AA	4144	1/1	0.95	0.12	100,100,100,100	0
59	MG	DA	3700	1/1	0.95	0.28	39,39,39,39	0
59	MG	DA	4323	1/1	0.95	0.21	56,56,56,56	0
59	MG	DA	3785	1/1	0.95	0.17	88,88,88,88	0
59	MG	DA	3279	1/1	0.95	0.39	45,45,45,45	0
59	MG	DA	5068	1/1	0.95	0.16	64,64,64,64	0
59	MG	DA	3988	1/1	0.95	0.37	59,59,59,59	0
59	MG	AA	4034	1/1	0.95	0.11	100,100,100,100	0
59	MG	BA	2271	1/1	0.95	0.30	106,106,106,106	0
59	MG	DA	3286	1/1	0.95	0.42	54,54,54,54	0
59	MG	BA	1921	1/1	0.95	0.09	85,85,85,85	0
59	MG	CN	201	1/1	0.95	0.10	33,33,33,33	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4147	1/1	0.96	0.33	42,42,42,42	0
59	MG	DA	3441	1/1	0.96	0.22	43,43,43,43	0
59	MG	DA	3061	1/1	0.96	0.17	23,23,23,23	0
59	MG	DA	3609	1/1	0.96	0.14	76,76,76,76	0
59	MG	AA	3621	1/1	0.96	0.04	46,46,46,46	0
59	MG	DA	4847	1/1	0.96	0.17	117,117,117,117	0
59	MG	CA	2223	1/1	0.96	0.21	67,67,67,67	0
59	MG	CA	1656	1/1	0.96	0.39	43,43,43,43	0
59	MG	BA	2031	1/1	0.96	0.11	71,71,71,71	0
59	MG	A5	101	1/1	0.96	0.14	23,23,23,23	0
59	MG	AA	3599	1/1	0.96	0.14	110,110,110,110	0
59	MG	CC	102	1/1	0.96	0.15	28,28,28,28	0
59	MG	AA	3911	1/1	0.96	0.19	69,69,69,69	0
59	MG	AA	3093	1/1	0.96	0.11	39,39,39,39	0
59	MG	CD	124	1/1	0.96	0.45	73,73,73,73	0
59	MG	BA	1972	1/1	0.96	0.10	53,53,53,53	0
59	MG	DA	3091	1/1	0.96	0.18	31,31,31,31	0
59	MG	AA	3694	1/1	0.96	0.09	56,56,56,56	0
59	MG	DA	3041	1/1	0.96	0.26	20,20,20,20	0
59	MG	CA	1751	1/1	0.96	0.22	34,34,34,34	0
59	MG	DU	203	1/1	0.96	0.17	45,45,45,45	0
59	MG	DA	3201	1/1	0.96	0.21	45,45,45,45	0
59	MG	AA	4008	1/1	0.96	0.25	78,78,78,78	0
59	MG	DB	202	1/1	0.96	0.28	48,48,48,48	0
59	MG	AA	3952	1/1	0.96	0.19	71,71,71,71	0
59	MG	DA	3461	1/1	0.96	0.11	34,34,34,34	0
59	MG	DA	3716	1/1	0.96	0.23	13,13,13,13	0
59	MG	CA	1872	1/1	0.96	0.08	74,74,74,74	0
59	MG	CA	1881	1/1	0.96	0.30	48,48,48,48	0
59	MG	DA	3413	1/1	0.96	0.32	45,45,45,45	0
59	MG	AA	3257	1/1	0.96	0.14	36,36,36,36	0
59	MG	DA	3917	1/1	0.96	0.14	77,77,77,77	0
59	MG	AA	3620	1/1	0.96	0.10	55,55,55,55	0
59	MG	BA	1885	1/1	0.96	0.25	44,44,44,44	0
59	MG	AA	3297	1/1	0.96	0.20	58,58,58,58	0
59	MG	DA	5048	1/1	0.96	0.18	64,64,64,64	0
59	MG	BA	2190	1/1	0.96	0.11	62,62,62,62	0
59	MG	AA	3170	1/1	0.96	0.20	32,32,32,32	0
59	MG	AA	3506	1/1	0.96	0.23	27,27,27,27	0
59	MG	AA	3528	1/1	0.96	0.56	46,46,46,46	0
59	MG	DA	3533	1/1	0.96	0.14	47,47,47,47	0
59	MG	DA	3137	1/1	0.96	0.36	31,31,31,31	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3582	1/1	0.96	0.28	54,54,54,54	0
59	MG	DA	3996	1/1	0.96	0.10	63,63,63,63	0
59	MG	DA	3362	1/1	0.96	0.24	35,35,35,35	0
59	MG	DA	4459	1/1	0.96	0.33	35,35,35,35	0
59	MG	AD	313	1/1	0.96	0.18	39,39,39,39	0
59	MG	DA	3794	1/1	0.96	0.30	38,38,38,38	0
59	MG	BA	2101	1/1	0.96	0.24	124,124,124,124	0
59	MG	AA	3234	1/1	0.96	0.12	54,54,54,54	0
59	MG	DU	202	1/1	0.96	0.13	17,17,17,17	0
59	MG	DA	3085	1/1	0.96	0.38	43,43,43,43	0
59	MG	CA	1918	1/1	0.96	0.44	86,86,86,86	0
59	MG	DA	3888	1/1	0.96	0.09	55,55,55,55	0
59	MG	DB	243	1/1	0.96	0.13	53,53,53,53	0
59	MG	DA	4797	1/1	0.96	0.28	88,88,88,88	0
59	MG	CA	1906	1/1	0.96	0.23	50,50,50,50	0
59	MG	DA	3470	1/1	0.96	0.18	52,52,52,52	0
59	MG	DA	4594	1/1	0.96	0.48	62,62,62,62	0
59	MG	CA	1917	1/1	0.96	0.15	64,64,64,64	0
59	MG	DA	4841	1/1	0.96	0.27	80,80,80,80	0
59	MG	BA	2059	1/1	0.96	0.19	67,67,67,67	0
59	MG	DA	3445	1/1	0.96	0.33	59,59,59,59	0
59	MG	AA	3021	1/1	0.96	0.23	26,26,26,26	0
59	MG	AA	3328	1/1	0.96	0.15	32,32,32,32	0
59	MG	AA	3581	1/1	0.96	0.23	55,55,55,55	0
59	MG	AA	3169	1/1	0.96	0.09	11,11,11,11	0
59	MG	AA	3398	1/1	0.96	0.24	30,30,30,30	0
59	MG	DA	4308	1/1	0.96	0.17	71,71,71,71	0
59	MG	CD	114	1/1	0.96	0.28	79,79,79,79	0
59	MG	BA	1764	1/1	0.96	0.08	50,50,50,50	0
59	MG	AA	3449	1/1	0.96	0.20	82,82,82,82	0
59	MG	DA	4324	1/1	0.96	0.13	49,49,49,49	0
59	MG	AA	3260	1/1	0.96	0.19	44,44,44,44	0
59	MG	BO	201	1/1	0.96	0.10	72,72,72,72	0
59	MG	DA	3807	1/1	0.96	0.12	59,59,59,59	0
59	MG	AA	3768	1/1	0.96	0.19	79,79,79,79	0
59	MG	DA	3389	1/1	0.96	0.20	48,48,48,48	0
59	MG	DA	3671	1/1	0.96	0.41	40,40,40,40	0
59	MG	CA	2226	1/1	0.96	0.12	53,53,53,53	0
59	MG	AA	3244	1/1	0.96	0.28	55,55,55,55	0
59	MG	AA	3299	1/1	0.96	0.12	82,82,82,82	0
59	MG	DA	4855	1/1	0.96	0.12	74,74,74,74	0
59	MG	AA	3204	1/1	0.96	0.09	68,68,68,68	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	2183	1/1	0.96	0.07	62,62,62,62	0
59	MG	DA	3866	1/1	0.96	0.26	59,59,59,59	0
59	MG	AA	3518	1/1	0.96	0.06	17,17,17,17	0
59	MG	BA	1914	1/1	0.96	0.10	55,55,55,55	0
59	MG	DA	4584	1/1	0.96	0.15	40,40,40,40	0
59	MG	CA	1886	1/1	0.96	0.12	43,43,43,43	0
59	MG	DA	4913	1/1	0.96	0.22	74,74,74,74	0
59	MG	DO	204	1/1	0.96	0.30	44,44,44,44	0
59	MG	DA	4209	1/1	0.96	0.18	83,83,83,83	0
59	MG	DA	4471	1/1	0.96	0.15	46,46,46,46	0
59	MG	DA	3923	1/1	0.96	0.17	70,70,70,70	0
59	MG	BA	2200	1/1	0.96	0.14	102,102,102,102	0
59	MG	DA	3506	1/1	0.96	0.15	46,46,46,46	0
59	MG	DA	4697	1/1	0.96	0.41	76,76,76,76	0
59	MG	DA	4464	1/1	0.96	0.18	51,51,51,51	0
59	MG	CA	1622	1/1	0.96	0.15	51,51,51,51	0
59	MG	DA	4189	1/1	0.96	0.08	60,60,60,60	0
59	MG	DA	3174	1/1	0.96	0.09	30,30,30,30	0
59	MG	DA	3156	1/1	0.96	0.24	27,27,27,27	0
59	MG	DA	3638	1/1	0.96	0.17	78,78,78,78	0
59	MG	BA	1900	1/1	0.96	0.10	83,83,83,83	0
59	MG	DA	4161	1/1	0.96	0.39	61,61,61,61	0
59	MG	AA	3815	1/1	0.96	0.11	61,61,61,61	0
59	MG	BA	1694	1/1	0.96	0.26	47,47,47,47	0
59	MG	DA	3733	1/1	0.96	0.10	40,40,40,40	0
59	MG	AA	3814	1/1	0.96	0.19	67,67,67,67	0
59	MG	DA	4555	1/1	0.96	0.10	43,43,43,43	0
59	MG	CA	1773	1/1	0.96	0.12	45,45,45,45	0
59	MG	BA	1795	1/1	0.96	0.32	67,67,67,67	0
59	MG	AA	3641	1/1	0.96	0.13	46,46,46,46	0
59	MG	BA	1732	1/1	0.96	0.44	62,62,62,62	0
59	MG	BA	1618	1/1	0.96	0.08	34,34,34,34	0
59	MG	DA	3339	1/1	0.96	0.12	68,68,68,68	0
59	MG	CA	2189	1/1	0.96	0.09	77,77,77,77	0
59	MG	AA	3479	1/1	0.96	0.13	59,59,59,59	0
59	MG	DA	3029	1/1	0.96	0.22	7,7,7,7	0
59	MG	DA	4686	1/1	0.96	0.38	59,59,59,59	0
59	MG	BA	1797	1/1	0.96	0.25	63,63,63,63	0
59	MG	DA	3066	1/1	0.96	0.25	17,17,17,17	0
59	MG	AA	3004	1/1	0.96	0.18	7,7,7,7	0
59	MG	BA	1819	1/1	0.96	0.17	61,61,61,61	0
59	MG	DA	3780	1/1	0.96	0.16	32,32,32,32	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	BA	1637	1/1	0.96	0.25	57,57,57,57	0
59	MG	CA	1936	1/1	0.96	0.11	55,55,55,55	0
59	MG	DA	3237	1/1	0.96	0.14	13,13,13,13	0
59	MG	DA	4411	1/1	0.96	0.16	42,42,42,42	0
59	MG	DA	3346	1/1	0.96	0.42	49,49,49,49	0
59	MG	DA	3005	1/1	0.96	0.26	17,17,17,17	0
59	MG	BA	1619	1/1	0.96	0.31	31,31,31,31	0
59	MG	AA	3594	1/1	0.96	0.11	78,78,78,78	0
59	MG	DA	3555	1/1	0.96	0.14	49,49,49,49	0
59	MG	DA	4708	1/1	0.96	0.14	46,46,46,46	0
59	MG	DA	4799	1/1	0.96	0.46	98,98,98,98	0
59	MG	DA	3361	1/1	0.96	0.13	50,50,50,50	0
59	MG	AA	3158	1/1	0.96	0.21	55,55,55,55	0
59	MG	DA	3224	1/1	0.96	0.11	31,31,31,31	0
59	MG	DA	3240	1/1	0.96	0.22	32,32,32,32	0
59	MG	DA	3826	1/1	0.96	0.23	33,33,33,33	0
59	MG	DA	5021	1/1	0.96	0.15	112,112,112,112	0
59	MG	AA	3250	1/1	0.96	0.10	24,24,24,24	0
59	MG	CA	1869	1/1	0.96	0.14	69,69,69,69	0
59	MG	DA	4228	1/1	0.96	0.15	50,50,50,50	0
59	MG	BA	1713	1/1	0.96	0.37	48,48,48,48	0
59	MG	DA	4350	1/1	0.96	0.21	54,54,54,54	0
59	MG	AA	3889	1/1	0.96	0.10	73,73,73,73	0
59	MG	DA	4016	1/1	0.96	0.07	40,40,40,40	0
59	MG	DA	4043	1/1	0.96	0.19	82,82,82,82	0
59	MG	CA	2083	1/1	0.96	0.17	87,87,87,87	0
60	ZN	CG	301	1/1	0.96	0.24	74,74,74,74	0
59	MG	CA	2177	1/1	0.96	0.19	116,116,116,116	0
59	MG	CA	1639	1/1	0.96	0.20	43,43,43,43	0
59	MG	DA	3436	1/1	0.96	0.16	48,48,48,48	0
59	MG	BA	1669	1/1	0.96	0.25	32,32,32,32	0
59	MG	DA	3076	1/1	0.96	0.24	13,13,13,13	0
59	MG	DE	307	1/1	0.96	0.18	62,62,62,62	0
59	MG	CA	1735	1/1	0.96	0.12	42,42,42,42	0
59	MG	DA	3667	1/1	0.96	0.27	74,74,74,74	0
59	MG	CF	303	1/1	0.96	0.78	79,79,79,79	0
59	MG	DA	4972	1/1	0.96	0.20	80,80,80,80	0
59	MG	BA	1838	1/1	0.96	0.16	91,91,91,91	0
59	MG	DA	3353	1/1	0.96	0.16	22,22,22,22	0
59	MG	DA	4963	1/1	0.96	0.12	63,63,63,63	0
59	MG	CA	1803	1/1	0.96	0.12	36,36,36,36	0
59	MG	AB	228	1/1	0.96	0.10	64,64,64,64	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3545	1/1	0.96	0.11	36,36,36,36	0
59	MG	DA	3002	1/1	0.96	0.28	11,11,11,11	0
59	MG	CA	1985	1/1	0.96	0.18	61,61,61,61	0
59	MG	DA	4965	1/1	0.96	0.36	61,61,61,61	0
59	MG	AA	3579	1/1	0.96	0.14	83,83,83,83	0
59	MG	CA	1870	1/1	0.96	0.10	58,58,58,58	0
60	ZN	BQ	104	1/1	0.96	0.11	97,97,97,97	0
59	MG	CA	2205	1/1	0.96	0.33	70,70,70,70	0
59	MG	DA	4457	1/1	0.96	0.13	75,75,75,75	0
59	MG	AA	4077	1/1	0.96	0.23	67,67,67,67	0
59	MG	AA	3098	1/1	0.96	0.21	45,45,45,45	0
59	MG	DA	3941	1/1	0.96	0.09	61,61,61,61	0
59	MG	BA	1610	1/1	0.96	0.11	25,25,25,25	0
59	MG	BA	1723	1/1	0.96	0.26	63,63,63,63	0
59	MG	DA	4000	1/1	0.96	0.11	50,50,50,50	0
59	MG	CA	1944	1/1	0.96	0.08	61,61,61,61	0
59	MG	AA	4163	1/1	0.96	0.38	102,102,102,102	0
59	MG	CA	2229	1/1	0.96	0.10	109,109,109,109	0
59	MG	DA	4609	1/1	0.96	0.14	48,48,48,48	0
59	MG	AA	3555	1/1	0.96	0.15	60,60,60,60	0
59	MG	AA	3002	1/1	0.96	0.11	14,14,14,14	0
59	MG	AA	3634	1/1	0.96	0.28	52,52,52,52	0
59	MG	AD	301	1/1	0.96	0.09	36,36,36,36	0
59	MG	DA	3817	1/1	0.96	0.17	46,46,46,46	0
59	MG	CA	1790	1/1	0.96	0.09	81,81,81,81	0
59	MG	BA	1993	1/1	0.96	0.13	60,60,60,60	0
59	MG	BA	1740	1/1	0.96	0.09	42,42,42,42	0
59	MG	BP	202	1/1	0.96	0.19	50,50,50,50	0
59	MG	DA	3110	1/1	0.96	0.10	35,35,35,35	0
59	MG	DA	3235	1/1	0.96	0.13	25,25,25,25	0
59	MG	CA	2271	1/1	0.96	0.12	55,55,55,55	0
59	MG	DA	3915	1/1	0.96	0.15	28,28,28,28	0
59	MG	DA	3497	1/1	0.96	0.41	38,38,38,38	0
59	MG	DA	3039	1/1	0.96	0.19	10,10,10,10	0
59	MG	AA	3085	1/1	0.96	0.07	29,29,29,29	0
59	MG	DA	3127	1/1	0.96	0.15	32,32,32,32	0
59	MG	DB	205	1/1	0.96	0.20	27,27,27,27	0
59	MG	BA	2071	1/1	0.96	0.30	93,93,93,93	0
59	MG	DA	3069	1/1	0.96	0.29	16,16,16,16	0
59	MG	DA	3296	1/1	0.96	0.19	19,19,19,19	0
59	MG	DA	3778	1/1	0.96	0.10	18,18,18,18	0
59	MG	DA	4669	1/1	0.96	0.26	71,71,71,71	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3724	1/1	0.96	0.30	45,45,45,45	0
59	MG	DA	4146	1/1	0.96	0.34	72,72,72,72	0
59	MG	DA	4212	1/1	0.96	0.21	54,54,54,54	0
59	MG	DA	4643	1/1	0.96	0.21	59,59,59,59	0
59	MG	BS	108	1/1	0.96	0.10	91,91,91,91	0
59	MG	AA	3610	1/1	0.96	0.05	69,69,69,69	0
59	MG	CA	2299	1/1	0.96	0.19	111,111,111,111	0
59	MG	AD	308	1/1	0.96	1.02	58,58,58,58	0
59	MG	AA	3391	1/1	0.96	0.16	39,39,39,39	0
59	MG	CA	2025	1/1	0.96	0.20	95,95,95,95	0
59	MG	AA	3462	1/1	0.96	0.15	32,32,32,32	0
59	MG	AA	3042	1/1	0.96	0.16	35,35,35,35	0
59	MG	DA	3015	1/1	0.96	0.25	30,30,30,30	0
59	MG	DA	3709	1/1	0.96	0.15	46,46,46,46	0
59	MG	AA	3937	1/1	0.96	0.19	67,67,67,67	0
59	MG	DA	4075	1/1	0.96	0.10	25,25,25,25	0
59	MG	AA	3388	1/1	0.96	0.20	46,46,46,46	0
59	MG	DA	3508	1/1	0.96	0.36	56,56,56,56	0
59	MG	CA	2089	1/1	0.96	0.19	63,63,63,63	0
59	MG	DF	307	1/1	0.96	0.37	27,27,27,27	0
59	MG	AA	3601	1/1	0.96	0.10	73,73,73,73	0
59	MG	AA	3152	1/1	0.96	0.08	28,28,28,28	0
59	MG	CA	1927	1/1	0.96	0.12	62,62,62,62	0
59	MG	CA	2029	1/1	0.96	0.22	115,115,115,115	0
59	MG	DA	3360	1/1	0.96	0.28	36,36,36,36	0
59	MG	DA	4721	1/1	0.96	0.11	75,75,75,75	0
59	MG	CA	1643	1/1	0.96	0.22	51,51,51,51	0
59	MG	DA	3763	1/1	0.96	0.21	8,8,8,8	0
59	MG	DA	4114	1/1	0.96	0.20	61,61,61,61	0
59	MG	AA	3996	1/1	0.96	0.13	49,49,49,49	0
59	MG	CA	1933	1/1	0.96	0.18	60,60,60,60	0
59	MG	AA	3024	1/1	0.96	0.10	61,61,61,61	0
59	MG	AA	3014	1/1	0.96	0.14	20,20,20,20	0
59	MG	AA	3970	1/1	0.96	0.10	47,47,47,47	0
59	MG	DA	3097	1/1	0.96	0.17	21,21,21,21	0
59	MG	DA	4047	1/1	0.96	0.10	44,44,44,44	0
59	MG	DO	213	1/1	0.96	0.29	96,96,96,96	0
59	MG	CA	1797	1/1	0.96	0.14	52,52,52,52	0
59	MG	CA	1690	1/1	0.96	0.10	43,43,43,43	0
59	MG	CA	2095	1/1	0.96	0.13	64,64,64,64	0
59	MG	CA	1804	1/1	0.96	0.24	67,67,67,67	0
59	MG	DA	4416	1/1	0.96	0.25	74,74,74,74	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4390	1/1	0.96	0.24	53,53,53,53	0
59	MG	BA	1970	1/1	0.96	0.15	141,141,141,141	0
59	MG	CA	2090	1/1	0.96	0.12	69,69,69,69	0
59	MG	BA	1965	1/1	0.96	0.31	110,110,110,110	0
59	MG	DA	4822	1/1	0.96	0.09	39,39,39,39	0
59	MG	DA	4420	1/1	0.96	0.12	40,40,40,40	0
59	MG	DA	4165	1/1	0.96	0.12	35,35,35,35	0
59	MG	CA	2068	1/1	0.96	0.12	89,89,89,89	0
59	MG	AA	3871	1/1	0.96	0.15	51,51,51,51	0
59	MG	CD	120	1/1	0.96	0.06	98,98,98,98	0
59	MG	DA	3383	1/1	0.96	0.11	49,49,49,49	0
59	MG	BA	1735	1/1	0.96	0.07	40,40,40,40	0
59	MG	DZ	102	1/1	0.96	0.24	49,49,49,49	0
59	MG	DA	4131	1/1	0.96	0.25	34,34,34,34	0
59	MG	DA	3211	1/1	0.96	0.31	43,43,43,43	0
59	MG	DA	4113	1/1	0.96	0.07	45,45,45,45	0
59	MG	DA	3053	1/1	0.96	0.43	24,24,24,24	0
59	MG	DA	4179	1/1	0.96	0.16	57,57,57,57	0
59	MG	DA	3241	1/1	0.96	0.22	20,20,20,20	0
59	MG	AA	3498	1/1	0.96	0.20	52,52,52,52	0
59	MG	AA	3196	1/1	0.96	0.15	29,29,29,29	0
59	MG	D8	102	1/1	0.96	0.40	37,37,37,37	0
59	MG	CA	1630	1/1	0.96	0.15	52,52,52,52	0
59	MG	AA	3630	1/1	0.96	0.37	55,55,55,55	0
59	MG	DA	3666	1/1	0.96	0.11	53,53,53,53	0
59	MG	AA	3851	1/1	0.96	0.11	60,60,60,60	0
59	MG	BA	1880	1/1	0.96	0.16	60,60,60,60	0
59	MG	CA	2088	1/1	0.96	0.20	65,65,65,65	0
59	MG	DA	4613	1/1	0.96	0.32	64,64,64,64	0
59	MG	AA	3336	1/1	0.96	0.10	122,122,122,122	0
59	MG	DA	3935	1/1	0.96	0.21	75,75,75,75	0
59	MG	DA	3740	1/1	0.96	0.12	49,49,49,49	0
59	MG	AA	3587	1/1	0.96	0.12	55,55,55,55	0
59	MG	CC	117	1/1	0.96	0.14	45,45,45,45	0
59	MG	DA	3126	1/1	0.96	0.28	63,63,63,63	0
59	MG	DA	3786	1/1	0.96	0.18	68,68,68,68	0
59	MG	CA	2110	1/1	0.96	0.10	62,62,62,62	0
59	MG	DA	4647	1/1	0.96	0.32	84,84,84,84	0
59	MG	CA	1891	1/1	0.96	0.26	56,56,56,56	0
59	MG	D8	108	1/1	0.96	0.17	52,52,52,52	0
59	MG	AA	3659	1/1	0.96	0.17	67,67,67,67	0
59	MG	AA	3200	1/1	0.96	0.34	83,83,83,83	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	BA	1895	1/1	0.96	0.26	54,54,54,54	0
59	MG	DA	3675	1/1	0.96	0.16	44,44,44,44	0
59	MG	AA	4030	1/1	0.96	0.17	66,66,66,66	0
59	MG	AA	3699	1/1	0.96	0.32	59,59,59,59	0
59	MG	AA	4156	1/1	0.96	0.17	59,59,59,59	0
59	MG	DA	3846	1/1	0.96	0.13	64,64,64,64	0
59	MG	CA	1898	1/1	0.96	0.09	59,59,59,59	0
59	MG	DA	4790	1/1	0.96	0.12	53,53,53,53	0
59	MG	BA	2074	1/1	0.96	0.45	77,77,77,77	0
59	MG	DA	3944	1/1	0.96	0.10	72,72,72,72	0
59	MG	D1	212	1/1	0.96	0.34	53,53,53,53	0
59	MG	DA	3704	1/1	0.96	0.22	33,33,33,33	0
59	MG	DA	3442	1/1	0.96	0.14	45,45,45,45	0
59	MG	D5	106	1/1	0.96	0.36	58,58,58,58	0
59	MG	DA	3950	1/1	0.96	0.26	117,117,117,117	0
59	MG	DA	4074	1/1	0.96	0.19	54,54,54,54	0
59	MG	CA	1733	1/1	0.96	0.28	55,55,55,55	0
59	MG	DA	4030	1/1	0.96	0.26	84,84,84,84	0
59	MG	DA	3402	1/1	0.96	0.15	24,24,24,24	0
59	MG	DA	4877	1/1	0.96	0.15	77,77,77,77	0
59	MG	BA	1655	1/1	0.96	0.32	58,58,58,58	0
59	MG	CA	1697	1/1	0.96	0.26	44,44,44,44	0
59	MG	DA	3880	1/1	0.96	0.38	44,44,44,44	0
59	MG	AA	3669	1/1	0.96	0.13	44,44,44,44	0
59	MG	CA	1631	1/1	0.96	0.18	42,42,42,42	0
59	MG	DA	3114	1/1	0.96	0.33	47,47,47,47	0
59	MG	DA	3738	1/1	0.96	0.14	45,45,45,45	0
59	MG	AA	3173	1/1	0.96	0.12	14,14,14,14	0
59	MG	CA	2207	1/1	0.96	0.15	70,70,70,70	0
59	MG	AA	3385	1/1	0.96	0.18	34,34,34,34	0
59	MG	DA	3021	1/1	0.96	0.31	26,26,26,26	0
59	MG	DA	4907	1/1	0.96	0.56	57,57,57,57	0
59	MG	CA	1644	1/1	0.96	0.16	24,24,24,24	0
59	MG	AA	3380	1/1	0.96	0.09	25,25,25,25	0
59	MG	CA	1686	1/1	0.96	0.28	35,35,35,35	0
59	MG	DA	3012	1/1	0.96	0.32	19,19,19,19	0
59	MG	DA	3800	1/1	0.96	0.13	33,33,33,33	0
59	MG	CA	2036	1/1	0.96	0.05	83,83,83,83	0
59	MG	DA	3705	1/1	0.96	0.07	1,1,1,1	0
59	MG	DA	4445	1/1	0.96	0.18	21,21,21,21	0
59	MG	AA	3677	1/1	0.96	0.13	60,60,60,60	0
59	MG	AA	3331	1/1	0.96	0.20	41,41,41,41	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3036	1/1	0.96	0.31	14,14,14,14	0
59	MG	DA	4293	1/1	0.96	0.38	54,54,54,54	0
59	MG	DA	4158	1/1	0.97	0.08	47,47,47,47	0
59	MG	DA	4242	1/1	0.97	0.17	49,49,49,49	0
59	MG	AA	3277	1/1	0.97	0.07	42,42,42,42	0
59	MG	AA	4043	1/1	0.97	0.71	132,132,132,132	0
59	MG	AA	3033	1/1	0.97	0.23	41,41,41,41	0
59	MG	DA	3271	1/1	0.97	0.24	33,33,33,33	0
59	MG	DA	4196	1/1	0.97	0.17	43,43,43,43	0
59	MG	CA	2067	1/1	0.97	0.16	89,89,89,89	0
59	MG	DA	3060	1/1	0.97	0.27	19,19,19,19	0
59	MG	AA	3520	1/1	0.97	0.15	75,75,75,75	0
59	MG	DA	3394	1/1	0.97	0.31	43,43,43,43	0
59	MG	DA	3541	1/1	0.97	0.11	19,19,19,19	0
59	MG	DA	4844	1/1	0.97	0.12	58,58,58,58	0
59	MG	AA	3764	1/1	0.97	0.17	53,53,53,53	0
59	MG	DA	3427	1/1	0.97	0.12	44,44,44,44	0
59	MG	AA	3207	1/1	0.97	0.20	30,30,30,30	0
59	MG	DA	4204	1/1	0.97	0.18	45,45,45,45	0
59	MG	DA	3734	1/1	0.97	0.06	55,55,55,55	0
59	MG	DA	4410	1/1	0.97	0.26	59,59,59,59	0
59	MG	DA	4346	1/1	0.97	0.21	44,44,44,44	0
59	MG	DA	4670	1/1	0.97	0.26	61,61,61,61	0
59	MG	CA	1670	1/1	0.97	0.36	46,46,46,46	0
59	MG	BA	1682	1/1	0.97	0.28	51,51,51,51	0
59	MG	AA	3778	1/1	0.97	0.10	53,53,53,53	0
59	MG	D7	103	1/1	0.97	0.28	45,45,45,45	0
59	MG	BA	1800	1/1	0.97	0.20	66,66,66,66	0
59	MG	AA	3139	1/1	0.97	0.07	45,45,45,45	0
59	MG	DA	3587	1/1	0.97	0.24	21,21,21,21	0
59	MG	AA	3070	1/1	0.97	0.14	35,35,35,35	0
59	MG	DA	4406	1/1	0.97	0.38	54,54,54,54	0
59	MG	DA	3784	1/1	0.97	0.16	52,52,52,52	0
59	MG	AA	3370	1/1	0.97	0.29	64,64,64,64	0
59	MG	DA	3871	1/1	0.97	0.45	49,49,49,49	0
59	MG	CA	2113	1/1	0.97	0.29	68,68,68,68	0
59	MG	DA	3702	1/1	0.97	0.12	36,36,36,36	0
59	MG	DA	3967	1/1	0.97	0.10	80,80,80,80	0
59	MG	CA	2254	1/1	0.97	0.18	56,56,56,56	0
59	MG	DA	3937	1/1	0.97	0.12	67,67,67,67	0
59	MG	DA	3044	1/1	0.97	0.31	18,18,18,18	0
59	MG	BA	2239	1/1	0.97	0.13	63,63,63,63	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4017	1/1	0.97	0.19	69,69,69,69	0
59	MG	DA	4743	1/1	0.97	0.29	96,96,96,96	0
59	MG	DA	3440	1/1	0.97	0.13	57,57,57,57	0
59	MG	DA	3964	1/1	0.97	0.11	68,68,68,68	0
59	MG	AA	3101	1/1	0.97	0.29	38,38,38,38	0
59	MG	BA	1616	1/1	0.97	0.27	47,47,47,47	0
59	MG	DA	4867	1/1	0.97	0.26	62,62,62,62	0
59	MG	AA	3664	1/1	0.97	0.13	68,68,68,68	0
59	MG	DA	4400	1/1	0.97	0.24	83,83,83,83	0
59	MG	DA	4902	1/1	0.97	0.19	82,82,82,82	0
59	MG	DA	3701	1/1	0.97	0.32	10,10,10,10	0
59	MG	DA	3251	1/1	0.97	0.11	12,12,12,12	0
59	MG	CA	1811	1/1	0.97	0.06	46,46,46,46	0
59	MG	DA	3834	1/1	0.97	0.20	57,57,57,57	0
59	MG	DA	4298	1/1	0.97	0.37	62,62,62,62	0
59	MG	DA	4969	1/1	0.97	0.10	75,75,75,75	0
59	MG	AR	203	1/1	0.97	0.05	55,55,55,55	0
59	MG	DA	3537	1/1	0.97	0.11	43,43,43,43	0
59	MG	DA	3737	1/1	0.97	0.24	36,36,36,36	0
59	MG	DE	302	1/1	0.97	0.22	8,8,8,8	0
59	MG	DA	4702	1/1	0.97	0.24	48,48,48,48	0
59	MG	DA	4444	1/1	0.97	0.13	51,51,51,51	0
59	MG	DA	3504	1/1	0.97	0.26	30,30,30,30	0
59	MG	AA	3365	1/1	0.97	0.25	39,39,39,39	0
59	MG	CA	1760	1/1	0.97	0.31	69,69,69,69	0
59	MG	CA	2276	1/1	0.97	0.16	69,69,69,69	0
59	MG	DA	3729	1/1	0.97	0.23	52,52,52,52	0
59	MG	BA	1779	1/1	0.97	0.27	45,45,45,45	0
59	MG	DA	3999	1/1	0.97	0.26	83,83,83,83	0
59	MG	DA	3599	1/1	0.97	0.17	6,6,6,6	0
59	MG	AA	3092	1/1	0.97	0.18	16,16,16,16	0
59	MG	DA	4435	1/1	0.97	0.14	53,53,53,53	0
59	MG	DA	3027	1/1	0.97	0.11	15,15,15,15	0
59	MG	CA	2285	1/1	0.97	0.17	135,135,135,135	0
59	MG	AA	3020	1/1	0.97	0.19	27,27,27,27	0
59	MG	DA	3910	1/1	0.97	0.10	143,143,143,143	0
59	MG	DA	4135	1/1	0.97	0.27	59,59,59,59	0
59	MG	AA	3066	1/1	0.97	0.12	17,17,17,17	0
59	MG	AA	4029	1/1	0.97	0.06	80,80,80,80	0
59	MG	AA	3087	1/1	0.97	0.17	20,20,20,20	0
59	MG	DA	4140	1/1	0.97	0.23	47,47,47,47	0
59	MG	BA	1716	1/1	0.97	0.22	56,56,56,56	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	BA	2248	1/1	0.97	0.07	61,61,61,61	0
59	MG	DA	3845	1/1	0.97	0.12	54,54,54,54	0
59	MG	DA	4020	1/1	0.97	0.21	153,153,153,153	0
59	MG	BD	115	1/1	0.97	0.09	61,61,61,61	0
59	MG	BA	1938	1/1	0.97	0.18	69,69,69,69	0
59	MG	DA	4225	1/1	0.97	0.11	83,83,83,83	0
59	MG	CA	1867	1/1	0.97	0.10	99,99,99,99	0
59	MG	DA	4550	1/1	0.97	0.20	61,61,61,61	0
59	MG	DA	4931	1/1	0.97	0.42	41,41,41,41	0
59	MG	DA	4112	1/1	0.97	0.08	45,45,45,45	0
59	MG	CA	1609	1/1	0.97	0.15	31,31,31,31	0
59	MG	DA	4748	1/1	0.97	0.09	83,83,83,83	0
59	MG	DU	216	1/1	0.97	0.17	80,80,80,80	0
59	MG	DA	3236	1/1	0.97	0.20	20,20,20,20	0
59	MG	DA	3810	1/1	0.97	0.18	39,39,39,39	0
59	MG	DA	3075	1/1	0.97	0.24	19,19,19,19	0
59	MG	DA	3663	1/1	0.97	0.09	79,79,79,79	0
59	MG	DA	4718	1/1	0.97	0.07	69,69,69,69	0
59	MG	DA	3249	1/1	0.97	0.14	13,13,13,13	0
59	MG	DA	4384	1/1	0.97	0.28	64,64,64,64	0
59	MG	AA	3231	1/1	0.97	0.09	43,43,43,43	0
59	MG	AA	3430	1/1	0.97	0.06	81,81,81,81	0
59	MG	AA	3012	1/1	0.97	0.27	21,21,21,21	0
59	MG	CQ	102	1/1	0.97	0.19	51,51,51,51	0
59	MG	CA	1655	1/1	0.97	0.12	18,18,18,18	0
59	MG	DA	4946	1/1	0.97	0.52	62,62,62,62	0
59	MG	DA	4651	1/1	0.97	0.09	66,66,66,66	0
59	MG	AA	3686	1/1	0.97	0.15	49,49,49,49	0
59	MG	DA	3616	1/1	0.97	0.21	64,64,64,64	0
59	MG	DA	3742	1/1	0.97	0.15	65,65,65,65	0
59	MG	DA	5003	1/1	0.97	0.31	76,76,76,76	0
59	MG	DA	3410	1/1	0.97	0.09	64,64,64,64	0
59	MG	DA	3300	1/1	0.97	0.17	27,27,27,27	0
59	MG	AA	3790	1/1	0.97	0.55	60,60,60,60	0
59	MG	DA	3443	1/1	0.97	0.08	54,54,54,54	0
59	MG	DA	3696	1/1	0.97	0.16	19,19,19,19	0
59	MG	AA	3456	1/1	0.97	0.13	20,20,20,20	0
59	MG	BA	1961	1/1	0.97	0.05	49,49,49,49	0
59	MG	DA	3985	1/1	0.97	0.24	81,81,81,81	0
59	MG	AA	3463	1/1	0.97	0.07	3,3,3,3	0
59	MG	DA	3006	1/1	0.97	0.13	14,14,14,14	0
59	MG	AA	4145	1/1	0.97	0.23	77,77,77,77	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3126	1/1	0.97	0.22	58,58,58,58	0
59	MG	BA	1653	1/1	0.97	0.37	47,47,47,47	0
59	MG	CA	1736	1/1	0.97	0.17	30,30,30,30	0
59	MG	AA	3300	1/1	0.97	0.28	38,38,38,38	0
59	MG	DA	3903	1/1	0.97	0.27	42,42,42,42	0
59	MG	CA	1810	1/1	0.97	0.11	61,61,61,61	0
59	MG	AA	3450	1/1	0.97	0.18	59,59,59,59	0
59	MG	DF	323	1/1	0.97	0.72	37,37,37,37	0
59	MG	DA	3170	1/1	0.97	0.28	16,16,16,16	0
59	MG	BA	1759	1/1	0.97	0.18	77,77,77,77	0
59	MG	A3	101	1/1	0.97	0.28	40,40,40,40	0
59	MG	DA	4198	1/1	0.97	0.17	45,45,45,45	0
59	MG	DA	4061	1/1	0.97	0.09	54,54,54,54	0
59	MG	DA	4160	1/1	0.97	0.33	53,53,53,53	0
59	MG	BA	1727	1/1	0.97	0.15	52,52,52,52	0
59	MG	CA	1719	1/1	0.97	0.09	62,62,62,62	0
59	MG	AA	4129	1/1	0.97	0.10	86,86,86,86	0
59	MG	AA	3487	1/1	0.97	0.14	72,72,72,72	0
59	MG	DG	202	1/1	0.97	0.07	47,47,47,47	0
59	MG	DA	3959	1/1	0.97	0.21	59,59,59,59	0
59	MG	DA	3225	1/1	0.97	0.10	39,39,39,39	0
59	MG	DA	4027	1/1	0.97	0.28	59,59,59,59	0
59	MG	AA	3026	1/1	0.97	0.38	38,38,38,38	0
59	MG	DA	3038	1/1	0.97	0.14	13,13,13,13	0
59	MG	CG	305	1/1	0.97	0.10	80,80,80,80	0
59	MG	DA	5075	1/1	0.97	0.28	50,50,50,50	0
59	MG	AA	3132	1/1	0.97	0.27	44,44,44,44	0
59	MG	DA	3557	1/1	0.97	0.13	55,55,55,55	0
59	MG	DA	3020	1/1	0.97	0.14	21,21,21,21	0
59	MG	DA	5029	1/1	0.97	0.29	39,39,39,39	0
59	MG	DA	4402	1/1	0.97	0.48	35,35,35,35	0
59	MG	D7	102	1/1	0.97	0.23	41,41,41,41	0
59	MG	BA	1686	1/1	0.97	0.27	34,34,34,34	0
59	MG	DB	251	1/1	0.97	0.14	49,49,49,49	0
59	MG	DA	3065	1/1	0.97	0.11	6,6,6,6	0
59	MG	DA	3062	1/1	0.97	0.21	19,19,19,19	0
59	MG	DU	206	1/1	0.97	0.12	47,47,47,47	0
59	MG	AA	3028	1/1	0.97	0.12	15,15,15,15	0
59	MG	DA	3612	1/1	0.97	0.15	64,64,64,64	0
59	MG	DA	3499	1/1	0.97	0.14	70,70,70,70	0
59	MG	CA	1923	1/1	0.97	0.13	48,48,48,48	0
59	MG	AA	3010	1/1	0.97	0.22	19,19,19,19	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3004	1/1	0.97	0.17	9,9,9,9	0
59	MG	AA	3013	1/1	0.97	0.14	49,49,49,49	0
59	MG	DA	3804	1/1	0.97	0.28	36,36,36,36	0
59	MG	DB	218	1/1	0.97	0.28	44,44,44,44	0
59	MG	AA	3390	1/1	0.97	0.23	46,46,46,46	0
59	MG	BA	1634	1/1	0.97	0.27	37,37,37,37	0
59	MG	AA	3107	1/1	0.97	0.27	32,32,32,32	0
59	MG	DA	4463	1/1	0.97	0.23	62,62,62,62	0
59	MG	AA	3128	1/1	0.97	0.23	32,32,32,32	0
59	MG	AA	3237	1/1	0.97	0.07	36,36,36,36	0
59	MG	D1	204	1/1	0.97	0.23	59,59,59,59	0
59	MG	CA	1998	1/1	0.97	0.11	90,90,90,90	0
59	MG	CA	1757	1/1	0.97	0.13	26,26,26,26	0
59	MG	DA	3677	1/1	0.97	0.25	78,78,78,78	0
59	MG	DA	4553	1/1	0.97	0.20	69,69,69,69	0
59	MG	CA	1868	1/1	0.97	0.07	29,29,29,29	0
59	MG	CA	1775	1/1	0.97	0.32	51,51,51,51	0
59	MG	DA	4565	1/1	0.97	0.24	119,119,119,119	0
59	MG	BA	1641	1/1	0.97	0.14	53,53,53,53	0
59	MG	AA	3312	1/1	0.97	0.13	40,40,40,40	0
59	MG	AA	3547	1/1	0.97	0.15	47,47,47,47	0
59	MG	DA	4631	1/1	0.97	0.17	67,67,67,67	0
59	MG	BA	2053	1/1	0.97	0.13	90,90,90,90	0
59	MG	AA	3980	1/1	0.97	0.38	65,65,65,65	0
59	MG	BA	2123	1/1	0.97	0.10	90,90,90,90	0
59	MG	AA	3108	1/1	0.97	0.19	41,41,41,41	0
59	MG	CC	126	1/1	0.97	0.08	55,55,55,55	0
59	MG	AA	4117	1/1	0.97	0.07	96,96,96,96	0
59	MG	BA	1702	1/1	0.97	0.22	33,33,33,33	0
59	MG	DA	4875	1/1	0.97	0.07	88,88,88,88	0
59	MG	DA	4796	1/1	0.97	0.26	45,45,45,45	0
59	MG	AA	4165	1/1	0.97	0.10	64,64,64,64	0
59	MG	DA	3579	1/1	0.97	0.24	21,21,21,21	0
59	MG	CH	201	1/1	0.97	0.30	55,55,55,55	0
59	MG	AA	3763	1/1	0.97	0.12	48,48,48,48	0
59	MG	AA	3401	1/1	0.97	0.36	46,46,46,46	0
59	MG	CA	1839	1/1	0.97	0.13	55,55,55,55	0
59	MG	AA	3136	1/1	0.97	0.24	38,38,38,38	0
59	MG	D2	205	1/1	0.97	0.31	84,84,84,84	0
59	MG	AA	3130	1/1	0.97	0.28	24,24,24,24	0
59	MG	AA	3609	1/1	0.97	0.14	72,72,72,72	0
59	MG	AA	3090	1/1	0.97	0.17	42,42,42,42	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	CA	1847	1/1	0.97	0.13	95,95,95,95	0
59	MG	DA	3688	1/1	0.97	0.23	42,42,42,42	0
59	MG	CA	1636	1/1	0.97	0.16	16,16,16,16	0
59	MG	DA	4109	1/1	0.97	0.24	61,61,61,61	0
59	MG	AA	3629	1/1	0.97	0.25	45,45,45,45	0
59	MG	DA	3319	1/1	0.97	0.15	46,46,46,46	0
59	MG	DF	308	1/1	0.97	0.51	30,30,30,30	0
59	MG	AA	3058	1/1	0.97	0.12	25,25,25,25	0
59	MG	DS	208	1/1	0.97	0.18	54,54,54,54	0
59	MG	DE	304	1/1	0.97	0.15	47,47,47,47	0
59	MG	DO	201	1/1	0.97	0.12	32,32,32,32	0
59	MG	BA	1739	1/1	0.97	0.34	67,67,67,67	0
59	MG	AA	3063	1/1	0.97	0.14	27,27,27,27	0
59	MG	DA	4977	1/1	0.97	0.34	94,94,94,94	0
59	MG	DA	4465	1/1	0.97	0.26	48,48,48,48	0
59	MG	DA	3045	1/1	0.97	0.28	23,23,23,23	0
59	MG	BC	104	1/1	0.97	0.15	45,45,45,45	0
59	MG	BA	1761	1/1	0.97	0.26	58,58,58,58	0
59	MG	DA	4216	1/1	0.97	0.24	52,52,52,52	0
59	MG	DA	4022	1/1	0.97	0.55	41,41,41,41	0
59	MG	DA	3683	1/1	0.97	0.13	56,56,56,56	0
59	MG	AA	3275	1/1	0.97	0.21	58,58,58,58	0
59	MG	DA	4531	1/1	0.97	0.23	57,57,57,57	0
59	MG	BA	1628	1/1	0.97	0.23	49,49,49,49	0
59	MG	DA	3369	1/1	0.97	0.21	16,16,16,16	0
59	MG	DA	3788	1/1	0.97	0.10	50,50,50,50	0
59	MG	DA	3381	1/1	0.97	0.24	66,66,66,66	0
59	MG	AA	3709	1/1	0.97	0.20	66,66,66,66	0
59	MG	DA	3167	1/1	0.97	0.12	48,48,48,48	0
59	MG	AA	3151	1/1	0.97	0.19	60,60,60,60	0
59	MG	AA	3932	1/1	0.97	0.09	56,56,56,56	0
59	MG	DA	3026	1/1	0.97	0.39	17,17,17,17	0
59	MG	AA	3175	1/1	0.97	0.10	29,29,29,29	0
59	MG	DA	4199	1/1	0.97	0.43	48,48,48,48	0
59	MG	BA	1640	1/1	0.97	0.28	27,27,27,27	0
59	MG	AA	3281	1/1	0.97	0.10	45,45,45,45	0
59	MG	DA	4013	1/1	0.97	0.39	42,42,42,42	0
59	MG	DA	4968	1/1	0.97	0.21	65,65,65,65	0
59	MG	AA	3650	1/1	0.97	0.10	70,70,70,70	0
59	MG	BA	2194	1/1	0.97	0.18	40,40,40,40	0
59	MG	AA	3007	1/1	0.97	0.22	24,24,24,24	0
59	MG	DA	3243	1/1	0.97	0.29	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	4858	1/1	0.97	0.27	69,69,69,69	0
59	MG	DA	3852	1/1	0.97	0.38	55,55,55,55	0
59	MG	DA	3703	1/1	0.97	0.31	12,12,12,12	0
59	MG	DA	4756	1/1	0.97	0.18	57,57,57,57	0
59	MG	AA	3389	1/1	0.97	0.15	67,67,67,67	0
59	MG	DA	4130	1/1	0.97	0.37	29,29,29,29	0
59	MG	AA	3850	1/1	0.97	0.25	41,41,41,41	0
59	MG	DA	3310	1/1	0.97	0.13	30,30,30,30	0
59	MG	CA	1832	1/1	0.97	0.15	115,115,115,115	0
59	MG	DA	4936	1/1	0.97	0.74	52,52,52,52	0
59	MG	CC	121	1/1	0.97	0.15	49,49,49,49	0
59	MG	DA	4561	1/1	0.97	0.58	43,43,43,43	0
59	MG	BA	1787	1/1	0.97	0.23	49,49,49,49	0
59	MG	DA	3035	1/1	0.97	0.20	15,15,15,15	0
59	MG	DB	223	1/1	0.97	0.07	58,58,58,58	0
59	MG	AA	4010	1/1	0.97	0.11	74,74,74,74	0
59	MG	CA	1613	1/1	0.97	0.12	41,41,41,41	0
59	MG	DA	3238	1/1	0.97	0.17	18,18,18,18	0
59	MG	AA	3326	1/1	0.97	0.10	22,22,22,22	0
59	MG	DA	3079	1/1	0.97	0.26	37,37,37,37	0
59	MG	BA	1604	1/1	0.97	0.22	47,47,47,47	0
59	MG	AA	4071	1/1	0.97	0.17	70,70,70,70	0
59	MG	DA	4272	1/1	0.97	0.28	56,56,56,56	0
59	MG	DA	4068	1/1	0.97	0.12	50,50,50,50	0
59	MG	AA	3209	1/1	0.97	0.21	77,77,77,77	0
59	MG	CA	1880	1/1	0.97	0.09	35,35,35,35	0
59	MG	CA	2296	1/1	0.97	0.06	68,68,68,68	0
59	MG	CA	2240	1/1	0.97	0.25	70,70,70,70	0
59	MG	AA	3383	1/1	0.97	0.14	24,24,24,24	0
59	MG	DA	4150	1/1	0.97	0.18	47,47,47,47	0
59	MG	AA	3759	1/1	0.97	0.09	59,59,59,59	0
59	MG	DA	3830	1/1	0.97	0.56	39,39,39,39	0
59	MG	BA	2045	1/1	0.97	0.05	66,66,66,66	0
59	MG	CA	2212	1/1	0.97	0.15	69,69,69,69	0
59	MG	D8	107	1/1	0.97	0.23	44,44,44,44	0
59	MG	AA	3091	1/1	0.97	0.38	59,59,59,59	0
59	MG	DA	3106	1/1	0.97	0.13	56,56,56,56	0
59	MG	DA	3938	1/1	0.97	0.08	96,96,96,96	0
59	MG	DA	4256	1/1	0.97	0.41	115,115,115,115	0
59	MG	AA	3323	1/1	0.97	0.11	48,48,48,48	0
59	MG	BC	103	1/1	0.97	0.09	51,51,51,51	0
59	MG	AA	3109	1/1	0.98	0.27	40,40,40,40	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	D1	203	1/1	0.98	0.10	42,42,42,42	0
59	MG	DA	3965	1/1	0.98	0.07	78,78,78,78	0
59	MG	DA	3798	1/1	0.98	0.18	30,30,30,30	0
59	MG	AA	3217	1/1	0.98	0.44	57,57,57,57	0
59	MG	AA	3251	1/1	0.98	0.18	24,24,24,24	0
59	MG	BA	2254	1/1	0.98	0.08	53,53,53,53	0
59	MG	CA	1819	1/1	0.98	0.11	72,72,72,72	0
59	MG	DA	3731	1/1	0.98	0.09	35,35,35,35	0
59	MG	DB	221	1/1	0.98	0.17	35,35,35,35	0
59	MG	DA	4443	1/1	0.98	0.56	36,36,36,36	0
59	MG	AA	3186	1/1	0.98	0.09	19,19,19,19	0
59	MG	AA	3017	1/1	0.98	0.35	29,29,29,29	0
59	MG	DA	4205	1/1	0.98	0.16	37,37,37,37	0
59	MG	DA	4479	1/1	0.98	0.07	50,50,50,50	0
59	MG	DA	3893	1/1	0.98	0.17	7,7,7,7	0
59	MG	DA	3428	1/1	0.98	0.24	38,38,38,38	0
59	MG	DA	3699	1/1	0.98	0.29	1,1,1,1	0
59	MG	AA	3940	1/1	0.98	0.31	61,61,61,61	0
59	MG	DA	3379	1/1	0.98	0.16	36,36,36,36	0
59	MG	DA	3963	1/1	0.98	0.15	59,59,59,59	0
59	MG	CA	1619	1/1	0.98	0.16	24,24,24,24	0
59	MG	DA	4333	1/1	0.98	0.12	52,52,52,52	0
59	MG	AA	3134	1/1	0.98	0.16	22,22,22,22	0
59	MG	CA	1876	1/1	0.98	0.12	79,79,79,79	0
59	MG	DA	4069	1/1	0.98	0.11	24,24,24,24	0
59	MG	CA	1748	1/1	0.98	0.27	27,27,27,27	0
59	MG	DA	4887	1/1	0.98	0.16	87,87,87,87	0
59	MG	BA	1870	1/1	0.98	0.10	119,119,119,119	0
59	MG	DA	4439	1/1	0.98	0.10	58,58,58,58	0
59	MG	AA	3069	1/1	0.98	0.12	29,29,29,29	0
59	MG	DA	4714	1/1	0.98	0.17	87,87,87,87	0
59	MG	DA	4213	1/1	0.98	0.24	49,49,49,49	0
60	ZN	CQ	101	1/1	0.98	0.09	114,114,114,114	0
59	MG	DA	4219	1/1	0.98	0.21	60,60,60,60	0
59	MG	AA	3381	1/1	0.98	0.16	32,32,32,32	0
59	MG	CA	1809	1/1	0.98	0.24	82,82,82,82	0
59	MG	DB	201	1/1	0.98	0.35	25,25,25,25	0
59	MG	DA	3477	1/1	0.98	0.24	171,171,171,171	0
59	MG	DA	4517	1/1	0.98	0.23	97,97,97,97	0
59	MG	AA	3377	1/1	0.98	0.14	19,19,19,19	0
59	MG	DA	3936	1/1	0.98	0.08	61,61,61,61	0
59	MG	D2	202	1/1	0.98	0.20	52,52,52,52	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	BA	1794	1/1	0.98	0.22	49,49,49,49	0
59	MG	DA	3070	1/1	0.98	0.25	24,24,24,24	0
59	MG	AA	3465	1/1	0.98	0.07	28,28,28,28	0
59	MG	AA	3078	1/1	0.98	0.24	44,44,44,44	0
59	MG	DA	4981	1/1	0.98	0.41	41,41,41,41	0
59	MG	AE	303	1/1	0.98	0.23	39,39,39,39	0
59	MG	DA	4270	1/1	0.98	0.18	46,46,46,46	0
59	MG	DA	3670	1/1	0.98	0.21	78,78,78,78	0
59	MG	DA	3632	1/1	0.98	0.10	52,52,52,52	0
59	MG	BA	1855	1/1	0.98	0.23	76,76,76,76	0
59	MG	CA	1713	1/1	0.98	0.42	57,57,57,57	0
59	MG	DA	4460	1/1	0.98	0.14	61,61,61,61	0
59	MG	DA	3155	1/1	0.98	0.18	32,32,32,32	0
59	MG	DA	3398	1/1	0.98	0.19	75,75,75,75	0
59	MG	DA	4395	1/1	0.98	0.19	46,46,46,46	0
59	MG	AA	3513	1/1	0.98	0.29	28,28,28,28	0
59	MG	AF	304	1/1	0.98	0.28	42,42,42,42	0
59	MG	CA	1687	1/1	0.98	0.20	46,46,46,46	0
59	MG	BA	1826	1/1	0.98	0.08	75,75,75,75	0
59	MG	DA	4987	1/1	0.98	0.32	37,37,37,37	0
59	MG	CA	1874	1/1	0.98	0.08	55,55,55,55	0
59	MG	AA	3531	1/1	0.98	0.26	26,26,26,26	0
59	MG	DA	4203	1/1	0.98	0.14	50,50,50,50	0
59	MG	DT	103	1/1	0.98	0.21	40,40,40,40	0
59	MG	DA	4370	1/1	0.98	0.21	74,74,74,74	0
59	MG	AA	3589	1/1	0.98	0.28	139,139,139,139	0
59	MG	AA	3006	1/1	0.98	0.19	14,14,14,14	0
59	MG	CA	2001	1/1	0.98	0.22	75,75,75,75	0
59	MG	AA	3492	1/1	0.98	0.06	48,48,48,48	0
59	MG	AA	3924	1/1	0.98	0.09	79,79,79,79	0
59	MG	AA	3232	1/1	0.98	0.08	25,25,25,25	0
59	MG	AA	3046	1/1	0.98	0.34	38,38,38,38	0
59	MG	DD	305	1/1	0.98	0.68	78,78,78,78	0
59	MG	DA	3276	1/1	0.98	0.29	34,34,34,34	0
59	MG	DA	4194	1/1	0.98	0.14	37,37,37,37	0
59	MG	AA	3240	1/1	0.98	0.22	34,34,34,34	0
59	MG	DA	3273	1/1	0.98	0.15	18,18,18,18	0
59	MG	DA	3068	1/1	0.98	0.18	15,15,15,15	0
59	MG	CA	1779	1/1	0.98	0.31	38,38,38,38	0
59	MG	DA	3109	1/1	0.98	0.20	18,18,18,18	0
59	MG	CA	1889	1/1	0.98	0.12	59,59,59,59	0
59	MG	AA	3616	1/1	0.98	0.08	42,42,42,42	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AE	301	1/1	0.98	0.18	35,35,35,35	0
59	MG	DA	3444	1/1	0.98	0.36	56,56,56,56	0
59	MG	DA	4157	1/1	0.98	0.23	37,37,37,37	0
59	MG	DA	4696	1/1	0.98	0.18	57,57,57,57	0
59	MG	DA	3080	1/1	0.98	0.37	31,31,31,31	0
59	MG	AA	3720	1/1	0.98	0.55	52,52,52,52	0
59	MG	DA	3478	1/1	0.98	0.10	18,18,18,18	0
59	MG	DA	4076	1/1	0.98	0.11	27,27,27,27	0
59	MG	DA	3687	1/1	0.98	0.16	41,41,41,41	0
59	MG	DA	3040	1/1	0.98	0.23	14,14,14,14	0
59	MG	DA	4767	1/1	0.98	0.28	52,52,52,52	0
59	MG	BA	1842	1/1	0.98	0.06	92,92,92,92	0
59	MG	DA	3295	1/1	0.98	0.21	14,14,14,14	0
59	MG	DA	4169	1/1	0.98	0.29	71,71,71,71	0
59	MG	DA	3032	1/1	0.98	0.25	6,6,6,6	0
59	MG	CA	1629	1/1	0.98	0.22	34,34,34,34	0
59	MG	BA	1791	1/1	0.98	0.10	55,55,55,55	0
59	MG	DA	3726	1/1	0.98	0.08	1,1,1,1	0
59	MG	AA	3203	1/1	0.98	0.12	40,40,40,40	0
59	MG	DA	3125	1/1	0.98	0.49	37,37,37,37	0
59	MG	DO	209	1/1	0.98	0.16	60,60,60,60	0
59	MG	DA	4251	1/1	0.98	0.15	51,51,51,51	0
59	MG	DA	3212	1/1	0.98	0.20	30,30,30,30	0
59	MG	DA	3602	1/1	0.98	0.36	61,61,61,61	0
59	MG	DA	3022	1/1	0.98	0.30	12,12,12,12	0
59	MG	DA	3145	1/1	0.98	0.26	39,39,39,39	0
59	MG	DA	3482	1/1	0.98	0.12	39,39,39,39	0
59	MG	CA	1606	1/1	0.98	0.13	22,22,22,22	0
59	MG	AA	3896	1/1	0.98	0.13	99,99,99,99	0
59	MG	DA	3895	1/1	0.98	0.24	14,14,14,14	0
59	MG	AA	3741	1/1	0.98	0.22	63,63,63,63	0
59	MG	AA	3001	1/1	0.98	0.18	21,21,21,21	0
59	MG	BA	1807	1/1	0.98	0.15	63,63,63,63	0
59	MG	AA	3222	1/1	0.98	0.18	28,28,28,28	0
59	MG	DA	3354	1/1	0.98	0.16	66,66,66,66	0
59	MG	DA	5064	1/1	0.98	0.12	61,61,61,61	0
59	MG	DA	3876	1/1	0.98	0.43	52,52,52,52	0
59	MG	DA	3092	1/1	0.98	0.12	3,3,3,3	0
59	MG	DA	3105	1/1	0.98	0.19	16,16,16,16	0
59	MG	AA	3466	1/1	0.98	0.03	37,37,37,37	0
59	MG	DA	4283	1/1	0.98	0.13	63,63,63,63	0
59	MG	DA	3753	1/1	0.98	0.07	62,62,62,62	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	BA	1617	1/1	0.98	0.34	44,44,44,44	0
59	MG	BA	1611	1/1	0.98	0.16	37,37,37,37	0
59	MG	BA	2259	1/1	0.98	0.18	101,101,101,101	0
59	MG	DA	3072	1/1	0.98	0.29	9,9,9,9	0
59	MG	BA	1836	1/1	0.98	0.07	57,57,57,57	0
59	MG	DA	4768	1/1	0.98	0.13	64,64,64,64	0
59	MG	DA	3059	1/1	0.98	0.13	8,8,8,8	0
59	MG	BE	302	1/1	0.98	0.09	59,59,59,59	0
59	MG	AA	3111	1/1	0.98	0.14	46,46,46,46	0
59	MG	AA	3501	1/1	0.98	0.14	24,24,24,24	0
59	MG	AB	205	1/1	0.98	0.07	54,54,54,54	0
59	MG	DA	3865	1/1	0.98	0.14	87,87,87,87	0
59	MG	DA	3037	1/1	0.98	0.16	1,1,1,1	0
59	MG	DA	3899	1/1	0.98	0.20	8,8,8,8	0
59	MG	AA	3316	1/1	0.98	0.10	47,47,47,47	0
59	MG	DA	3782	1/1	0.98	0.10	38,38,38,38	0
59	MG	DA	4666	1/1	0.98	0.36	60,60,60,60	0
59	MG	AA	3793	1/1	0.98	0.44	63,63,63,63	0
59	MG	AA	3500	1/1	0.98	0.18	22,22,22,22	0
59	MG	DA	4372	1/1	0.98	0.15	53,53,53,53	0
59	MG	DA	3694	1/1	0.98	0.17	41,41,41,41	0
59	MG	DA	5009	1/1	0.98	0.25	66,66,66,66	0
59	MG	DA	3133	1/1	0.98	0.31	45,45,45,45	0
59	MG	AA	3206	1/1	0.98	0.10	40,40,40,40	0
59	MG	AA	3472	1/1	0.98	0.17	59,59,59,59	0
59	MG	DA	3101	1/1	0.98	0.33	45,45,45,45	0
59	MG	AA	3095	1/1	0.98	0.18	18,18,18,18	0
59	MG	DA	3546	1/1	0.98	0.28	29,29,29,29	0
59	MG	AA	3523	1/1	0.98	0.23	42,42,42,42	0
59	MG	DA	4291	1/1	0.98	0.13	58,58,58,58	0
59	MG	DA	3017	1/1	0.98	0.22	7,7,7,7	0
59	MG	AA	3230	1/1	0.98	0.17	51,51,51,51	0
59	MG	DA	3052	1/1	0.98	0.14	19,19,19,19	0
59	MG	AA	3011	1/1	0.98	0.18	17,17,17,17	0
59	MG	DA	3698	1/1	0.98	0.12	35,35,35,35	0
59	MG	CA	1662	1/1	0.98	0.06	19,19,19,19	0
59	MG	DA	3972	1/1	0.98	0.16	41,41,41,41	0
59	MG	DA	4835	1/1	0.98	0.11	65,65,65,65	0
59	MG	AA	3039	1/1	0.98	0.23	30,30,30,30	0
59	MG	DB	247	1/1	0.98	0.09	61,61,61,61	0
59	MG	DA	3744	1/1	0.98	0.11	58,58,58,58	0
59	MG	DA	4375	1/1	0.98	0.25	52,52,52,52	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3505	1/1	0.98	0.35	51,51,51,51	0
59	MG	AA	3226	1/1	0.98	0.07	68,68,68,68	0
59	MG	AA	3507	1/1	0.98	0.21	30,30,30,30	0
59	MG	DA	3011	1/1	0.98	0.35	1,1,1,1	0
59	MG	DA	3956	1/1	0.98	0.27	45,45,45,45	0
59	MG	DA	4808	1/1	0.98	0.50	61,61,61,61	0
59	MG	CA	2093	1/1	0.98	0.10	64,64,64,64	0
59	MG	AA	3034	1/1	0.98	0.12	41,41,41,41	0
59	MG	CA	2085	1/1	0.98	0.16	58,58,58,58	0
59	MG	DA	4882	1/1	0.98	0.16	74,74,74,74	0
59	MG	DA	3028	1/1	0.98	0.20	18,18,18,18	0
59	MG	DA	3071	1/1	0.98	0.21	13,13,13,13	0
59	MG	DA	3063	1/1	0.98	0.38	23,23,23,23	0
59	MG	AA	3859	1/1	0.98	0.18	80,80,80,80	0
59	MG	DA	3706	1/1	0.98	0.26	20,20,20,20	0
59	MG	AA	3600	1/1	0.98	0.09	46,46,46,46	0
59	MG	CA	1647	1/1	0.98	0.20	13,13,13,13	0
59	MG	DA	3185	1/1	0.98	0.25	54,54,54,54	0
59	MG	DA	3776	1/1	0.98	0.12	46,46,46,46	0
59	MG	DA	4419	1/1	0.98	0.14	57,57,57,57	0
59	MG	DA	3358	1/1	0.98	0.27	23,23,23,23	0
59	MG	DA	4090	1/1	0.98	0.06	52,52,52,52	0
59	MG	AA	3003	1/1	0.98	0.10	12,12,12,12	0
59	MG	DA	4024	1/1	0.98	0.07	47,47,47,47	0
59	MG	BA	1904	1/1	0.98	0.21	38,38,38,38	0
59	MG	DA	3585	1/1	0.98	0.28	10,10,10,10	0
59	MG	DA	4401	1/1	0.98	0.30	87,87,87,87	0
59	MG	CA	1607	1/1	0.98	0.21	20,20,20,20	0
59	MG	AA	3731	1/1	0.98	0.08	79,79,79,79	0
59	MG	DA	3432	1/1	0.98	0.23	46,46,46,46	0
59	MG	DA	3244	1/1	0.98	0.15	97,97,97,97	0
59	MG	BA	1953	1/1	0.98	0.11	68,68,68,68	0
59	MG	AA	3140	1/1	0.98	0.21	57,57,57,57	0
59	MG	DA	3031	1/1	0.98	0.23	3,3,3,3	0
59	MG	AA	3348	1/1	0.98	0.13	26,26,26,26	0
59	MG	AA	3378	1/1	0.98	0.12	50,50,50,50	0
59	MG	BA	2273	1/1	0.98	0.16	94,94,94,94	0
59	MG	DA	4441	1/1	0.98	0.17	62,62,62,62	0
59	MG	AA	3048	1/1	0.98	0.21	41,41,41,41	0
59	MG	AA	3047	1/1	0.98	0.08	28,28,28,28	0
59	MG	DA	3774	1/1	0.98	0.13	37,37,37,37	0
59	MG	CA	1640	1/1	0.98	0.05	28,28,28,28	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DB	262	1/1	0.98	0.45	89,89,89,89	0
59	MG	DA	3153	1/1	0.98	0.26	26,26,26,26	0
59	MG	AA	3177	1/1	0.98	0.19	26,26,26,26	0
59	MG	DA	3057	1/1	0.98	0.23	6,6,6,6	0
59	MG	DA	3129	1/1	0.99	0.22	14,14,14,14	0
59	MG	DA	3925	1/1	0.99	0.06	25,25,25,25	0
59	MG	CA	1671	1/1	0.99	0.32	24,24,24,24	0
59	MG	AA	3133	1/1	0.99	0.10	25,25,25,25	0
59	MG	DA	4725	1/1	0.99	0.21	56,56,56,56	0
59	MG	DA	4023	1/1	0.99	0.09	41,41,41,41	0
59	MG	AA	3112	1/1	0.99	0.18	102,102,102,102	0
59	MG	DA	3529	1/1	0.99	0.30	84,84,84,84	0
59	MG	AA	3191	1/1	0.99	0.28	31,31,31,31	0
59	MG	DA	4162	1/1	0.99	0.28	59,59,59,59	0
59	MG	AA	3129	1/1	0.99	0.14	31,31,31,31	0
59	MG	CA	2039	1/1	0.99	0.11	66,66,66,66	0
59	MG	AA	3504	1/1	0.99	0.16	31,31,31,31	0
59	MG	DA	4312	1/1	0.99	0.20	49,49,49,49	0
59	MG	DA	3316	1/1	0.99	0.11	43,43,43,43	0
59	MG	DA	3001	1/1	0.99	0.25	8,8,8,8	0
59	MG	AA	3227	1/1	0.99	0.14	30,30,30,30	0
59	MG	AA	3015	1/1	0.99	0.15	19,19,19,19	0
59	MG	AA	3459	1/1	0.99	0.32	30,30,30,30	0
59	MG	DA	3749	1/1	0.99	0.05	92,92,92,92	0
59	MG	DA	4434	1/1	0.99	0.18	59,59,59,59	0
59	MG	DA	4729	1/1	0.99	0.36	39,39,39,39	0
59	MG	DA	4873	1/1	0.99	0.17	49,49,49,49	0
59	MG	AA	3190	1/1	0.99	0.21	26,26,26,26	0
59	MG	DA	3016	1/1	0.99	0.17	4,4,4,4	0
59	MG	DA	3966	1/1	0.99	0.09	41,41,41,41	0
59	MG	DA	3086	1/1	0.99	0.19	17,17,17,17	0
59	MG	DA	4191	1/1	0.99	0.21	48,48,48,48	0
59	MG	DA	4850	1/1	0.99	0.15	38,38,38,38	0
59	MG	DA	3437	1/1	0.99	0.15	47,47,47,47	0
59	MG	AA	3057	1/1	0.99	0.19	49,49,49,49	0
59	MG	DA	3563	1/1	0.99	0.14	44,44,44,44	0
59	MG	DA	3323	1/1	0.99	0.21	41,41,41,41	0
59	MG	DA	4959	1/1	0.99	0.15	45,45,45,45	0
59	MG	DA	3033	1/1	0.99	0.28	13,13,13,13	0
59	MG	DA	3082	1/1	0.99	0.23	27,27,27,27	0
59	MG	AA	3696	1/1	0.99	0.15	47,47,47,47	0
59	MG	BA	2025	1/1	0.99	0.17	90,90,90,90	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	AA	3509	1/1	0.99	0.16	19,19,19,19	0
59	MG	CA	1762	1/1	0.99	0.11	49,49,49,49	0
59	MG	AA	3483	1/1	0.99	0.42	65,65,65,65	0
59	MG	DA	5036	1/1	0.99	0.22	52,52,52,52	0
59	MG	DA	4711	1/1	0.99	0.15	94,94,94,94	0
59	MG	DA	3014	1/1	0.99	0.37	7,7,7,7	0
59	MG	DD	314	1/1	0.99	0.74	32,32,32,32	0
59	MG	AA	3009	1/1	0.99	0.16	18,18,18,18	0
59	MG	DA	3087	1/1	0.99	0.17	15,15,15,15	0
59	MG	DA	3292	1/1	0.99	0.11	48,48,48,48	0
59	MG	DA	3242	1/1	0.99	0.25	18,18,18,18	0
59	MG	DA	4172	1/1	0.99	0.23	41,41,41,41	0
59	MG	DA	3297	1/1	0.99	0.20	27,27,27,27	0
59	MG	AA	3228	1/1	0.99	0.06	53,53,53,53	0
59	MG	DA	4167	1/1	0.99	0.28	44,44,44,44	0
59	MG	DA	4138	1/1	0.99	0.11	54,54,54,54	0
59	MG	DA	3056	1/1	0.99	0.21	12,12,12,12	0
59	MG	CA	2015	1/1	0.99	0.10	57,57,57,57	0
59	MG	DB	206	1/1	0.99	0.04	34,34,34,34	0
59	MG	DA	3583	1/1	0.99	0.21	3,3,3,3	0
59	MG	DA	4091	1/1	0.99	0.10	42,42,42,42	0
59	MG	DO	215	1/1	0.99	0.09	41,41,41,41	0
59	MG	BA	1837	1/1	0.99	0.10	88,88,88,88	0
59	MG	DA	3034	1/1	0.99	0.24	11,11,11,11	0
59	MG	DA	4772	1/1	0.99	0.11	46,46,46,46	0
59	MG	DA	3589	1/1	0.99	0.14	7,7,7,7	0
59	MG	DA	3239	1/1	0.99	0.35	37,37,37,37	0
59	MG	CA	2125	1/1	0.99	0.28	79,79,79,79	0
59	MG	BA	2214	1/1	0.99	0.15	90,90,90,90	0
59	MG	DA	3051	1/1	0.99	0.23	17,17,17,17	0
59	MG	CP	201	1/1	0.99	0.08	54,54,54,54	0
59	MG	CA	1758	1/1	0.99	0.12	52,52,52,52	0
59	MG	DA	4144	1/1	0.99	0.31	45,45,45,45	0
59	MG	AA	3489	1/1	0.99	0.30	63,63,63,63	0
60	ZN	BG	308	1/1	0.99	0.25	71,71,71,71	0
59	MG	AA	3916	1/1	0.99	0.15	93,93,93,93	0
59	MG	DA	3007	1/1	0.99	0.33	10,10,10,10	0
59	MG	DA	3030	1/1	0.99	0.20	3,3,3,3	0
59	MG	DA	3380	1/1	0.99	0.13	55,55,55,55	0
59	MG	DA	4049	1/1	0.99	0.20	33,33,33,33	0
59	MG	AA	3926	1/1	0.99	0.11	52,52,52,52	0
59	MG	DA	4487	1/1	0.99	0.20	94,94,94,94	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
59	MG	DA	3043	1/1	0.99	0.23	23,23,23,23	0
59	MG	DA	3420	1/1	0.99	0.32	51,51,51,51	0
59	MG	CA	1724	1/1	0.99	0.20	45,45,45,45	0
59	MG	DA	3618	1/1	0.99	0.11	14,14,14,14	0
59	MG	CA	2244	1/1	0.99	0.18	126,126,126,126	0
59	MG	AA	3577	1/1	0.99	0.08	80,80,80,80	0
59	MG	D2	203	1/1	0.99	0.19	57,57,57,57	0
59	MG	CA	2243	1/1	0.99	0.12	53,53,53,53	0
59	MG	DA	4058	1/1	0.99	0.13	81,81,81,81	0
59	MG	AA	3267	1/1	0.99	0.09	49,49,49,49	0
59	MG	DA	3991	1/1	0.99	0.07	96,96,96,96	0
59	MG	DA	3089	1/1	0.99	0.36	23,23,23,23	0
59	MG	DA	3058	1/1	0.99	0.19	15,15,15,15	0
59	MG	DA	4220	1/1	0.99	0.10	40,40,40,40	0
59	MG	AA	3223	1/1	0.99	0.15	14,14,14,14	0
59	MG	DA	3886	1/1	0.99	0.04	48,48,48,48	0
59	MG	DA	3024	1/1	0.99	0.20	9,9,9,9	0
59	MG	AA	3522	1/1	0.99	0.04	14,14,14,14	0
59	MG	AA	3138	1/1	0.99	0.17	19,19,19,19	0
59	MG	DA	4111	1/1	0.99	0.09	54,54,54,54	0
59	MG	AA	3053	1/1	0.99	0.15	30,30,30,30	0
59	MG	AA	3147	1/1	0.99	0.12	60,60,60,60	0
59	MG	DA	3973	1/1	0.99	0.13	60,60,60,60	0
59	MG	DA	4025	1/1	0.99	0.07	48,48,48,48	0
59	MG	DA	3047	1/1	0.99	0.18	1,1,1,1	0
59	MG	DA	3697	1/1	0.99	0.18	6,6,6,6	0
59	MG	CA	1711	1/1	0.99	0.13	62,62,62,62	0
59	MG	BA	1747	1/1	0.99	0.18	50,50,50,50	0
59	MG	DA	3067	1/1	0.99	0.40	14,14,14,14	0
59	MG	DA	4002	1/1	0.99	0.15	42,42,42,42	0
59	MG	DA	3009	1/1	0.99	0.18	10,10,10,10	0
59	MG	DA	3650	1/1	1.00	0.17	37,37,37,37	0

6.5 Other polymers ⓘ

There are no such residues in this entry.