



# Full wwPDB X-ray Structure Validation Report ⓘ

May 16, 2020 – 12:39 am BST

PDB ID : 4WSM  
Title : Complex of 70S ribosome with tRNA-Leu and mRNA with G-U mismatch in the first position in the A- and P-sites  
Authors : Rozov, A.; Demeshkina, N.; Yusupov, M.; Yusupova, G.  
Deposited on : 2014-10-28  
Resolution : 3.30 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](mailto: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.

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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

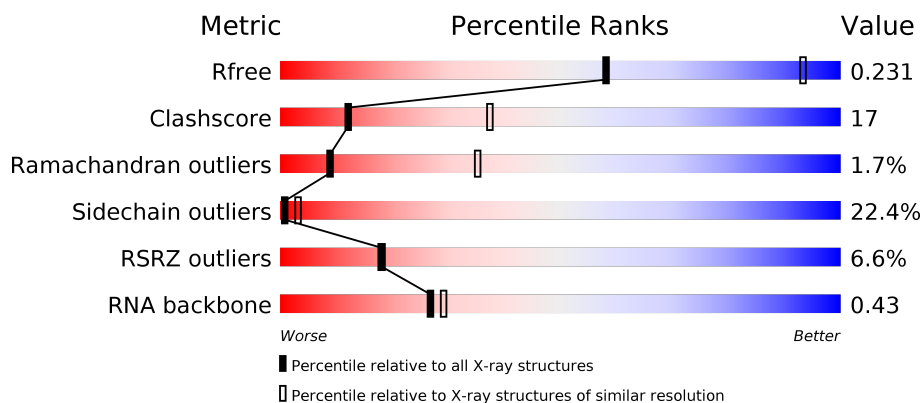
# 1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

*X-RAY DIFFRACTION*


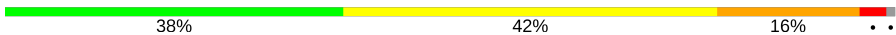


The reported resolution of this entry is 3.30 Å.

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	1149 (3.34-3.26)
Clashscore	141614	1205 (3.34-3.26)
Ramachandran outliers	138981	1183 (3.34-3.26)
Sidechain outliers	138945	1182 (3.34-3.26)
RSRZ outliers	127900	1115 (3.34-3.26)
RNA backbone	3102	1117 (3.70-2.90)

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  $\geq 3$ , 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	13	1522	
1	1G	1522	
2	12	256	
2	1E	256	

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Mol	Chain	Length	Quality of chain
3	22	239	
3	2E	239	
4	32	209	
4	3E	209	
5	42	162	
5	4E	162	
6	52	101	
6	5E	101	
7	62	156	
7	6E	156	
8	72	138	
8	7E	138	
9	82	128	
9	8E	128	
10	1A	105	
10	1I	105	
11	2A	129	
11	2I	129	
12	3A	132	
12	3I	132	
13	4A	126	
13	4I	126	
14	5A	61	
14	5I	61	
15	6A	89	

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Mol	Chain	Length	Quality of chain
15	6I	89	
16	7A	88	
16	7I	88	
17	8A	105	
17	8I	105	
18	9A	88	
18	9I	88	
19	AA	93	
19	AI	93	
20	BA	106	
20	BI	106	
21	1B	27	
21	1F	27	
22	1K	87	
22	1L	87	
23	2K	87	
23	2L	87	
24	3K	87	
24	3L	87	
25	4K	60	
25	4L	60	
26	14	2917	
26	1H	2917	
27	16	122	
27	1J	122	

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Mol	Chain	Length	Quality of chain
28	11	276	
28	19	276	
29	21	206	
29	29	206	
30	31	210	
30	39	210	
31	41	182	
31	49	182	
32	51	180	
32	59	180	
33	61	148	
33	69	148	
34	15	140	
34	58	140	
35	25	122	
35	68	122	
36	35	150	
36	78	150	
37	45	141	
37	88	141	
38	55	118	
38	98	118	
39	65	112	
39	A8	112	
40	75	146	

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Mol	Chain	Length	Quality of chain
40	B8	146	
41	85	118	
41	C8	118	
42	95	101	
42	D8	101	
43	A5	113	
43	E8	113	
44	B5	96	
44	F8	96	
45	C5	110	
45	G8	110	
46	D5	206	
46	H8	206	
47	E5	85	
47	I8	85	
48	F5	98	
48	J8	98	
49	G5	72	
49	K8	72	
50	H5	60	
50	L8	60	
51	I5	71	
51	M8	71	
52	J5	60	
52	N8	60	

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Mol	Chain	Length	Quality of chain
53	L5	49	
53	P8	49	
54	M5	65	
54	Q8	65	

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
55	MG	13	1612	-	-	-	X
55	MG	13	1627	-	-	-	X
55	MG	13	1631	-	-	-	X
55	MG	13	1672	-	-	-	X
55	MG	13	1673	-	-	-	X
55	MG	13	1677	-	-	-	X
55	MG	13	1678	-	-	-	X
55	MG	13	1683	-	-	-	X
55	MG	13	1684	-	-	-	X
55	MG	13	1686	-	-	-	X
55	MG	13	1705	-	-	-	X
55	MG	13	1713	-	-	-	X
55	MG	13	1721	-	-	-	X
55	MG	13	1727	-	-	-	X
55	MG	14	3001	-	-	-	X
55	MG	14	3035	-	-	-	X
55	MG	14	3060	-	-	-	X
55	MG	14	3096	-	-	-	X
55	MG	14	3116	-	-	-	X
55	MG	14	3119	-	-	-	X
55	MG	14	3127	-	-	-	X
55	MG	14	3130	-	-	-	X
55	MG	14	3131	-	-	-	X
55	MG	14	3152	-	-	-	X
55	MG	14	3155	-	-	-	X
55	MG	14	3159	-	-	-	X
55	MG	14	3162	-	-	-	X
55	MG	14	3171	-	-	-	X
55	MG	14	3173	-	-	-	X
55	MG	14	3175	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
55	MG	14	3200	-	-	-	X
55	MG	14	3202	-	-	-	X
55	MG	14	3204	-	-	-	X
55	MG	14	3221	-	-	-	X
55	MG	14	3229	-	-	-	X
55	MG	14	3230	-	-	-	X
55	MG	14	3233	-	-	-	X
55	MG	14	3235	-	-	-	X
55	MG	14	3247	-	-	-	X
55	MG	14	3252	-	-	-	X
55	MG	14	3265	-	-	-	X
55	MG	14	3272	-	-	-	X
55	MG	14	3275	-	-	-	X
55	MG	14	3276	-	-	-	X
55	MG	14	3292	-	-	-	X
55	MG	14	3301	-	-	-	X
55	MG	14	3306	-	-	-	X
55	MG	14	3319	-	-	-	X
55	MG	14	3323	-	-	-	X
55	MG	14	3324	-	-	-	X
55	MG	14	3352	-	-	-	X
55	MG	14	3358	-	-	-	X
55	MG	14	3367	-	-	-	X
55	MG	14	3369	-	-	-	X
55	MG	14	3372	-	-	-	X
55	MG	14	3374	-	-	-	X
55	MG	14	3381	-	-	-	X
55	MG	14	3385	-	-	-	X
55	MG	14	3386	-	-	-	X
55	MG	14	3388	-	-	-	X
55	MG	14	3393	-	-	-	X
55	MG	1G	1659	-	-	-	X
55	MG	1G	1690	-	-	-	X
55	MG	1G	1707	-	-	-	X
55	MG	1G	1708	-	-	-	X
55	MG	1G	1717	-	-	-	X
55	MG	1G	1729	-	-	-	X
55	MG	1H	3073	-	-	-	X
55	MG	1H	3086	-	-	-	X
55	MG	1H	3104	-	-	-	X
55	MG	1H	3125	-	-	-	X
55	MG	1H	3145	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
55	MG	1H	3154	-	-	-	X
55	MG	1H	3158	-	-	-	X
55	MG	1H	3164	-	-	-	X
55	MG	1H	3175	-	-	-	X
55	MG	1H	3183	-	-	-	X
55	MG	1H	3188	-	-	-	X
55	MG	1H	3195	-	-	-	X
55	MG	1H	3206	-	-	-	X
55	MG	1H	3208	-	-	-	X
55	MG	1H	3211	-	-	-	X
55	MG	1H	3229	-	-	-	X
55	MG	1H	3241	-	-	-	X
55	MG	1H	3242	-	-	-	X
55	MG	1H	3248	-	-	-	X
55	MG	1H	3250	-	-	-	X
55	MG	1H	3259	-	-	-	X
55	MG	1H	3272	-	-	-	X
55	MG	1H	3279	-	-	-	X
55	MG	1H	3283	-	-	-	X
55	MG	1H	3313	-	-	-	X
55	MG	1H	3314	-	-	-	X
55	MG	1H	3319	-	-	-	X
55	MG	1H	3324	-	-	-	X
55	MG	1H	3338	-	-	-	X
55	MG	1H	3343	-	-	-	X
55	MG	1H	3361	-	-	-	X
55	MG	1H	3368	-	-	-	X
55	MG	1H	3380	-	-	-	X
55	MG	1H	3381	-	-	-	X
55	MG	1H	3400	-	-	-	X
55	MG	1H	3402	-	-	-	X
55	MG	1H	3404	-	-	-	X
55	MG	1H	3406	-	-	-	X
55	MG	45	201	-	-	-	X
55	MG	88	304	-	-	-	X

## 2 Entry composition

There are 57 unique types of molecules in this entry. The entry contains 299429 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 16S ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
1	13	1507	Total	C	N	O	P	0	0	0
			32389	14416	5999	10467	1507			
1	1G	1513	Total	C	N	O	P	0	0	0
			32511	14472	6018	10509	1512			

- Molecule 2 is a protein called 30S ribosomal protein S2.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
2	1E	237	Total	C	N	O	S	0	0	0
			1924	1228	344	347	5			
2	12	237	Total	C	N	O	S	0	0	0
			1924	1228	344	347	5			

- Molecule 3 is a protein called 30S ribosomal protein S3.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
3	2E	205	Total	C	N	O	S	0	0	0
			1605	1011	313	280	1			
3	22	206	Total	C	N	O	S	0	0	0
			1612	1016	314	281	1			

- Molecule 4 is a protein called 30S ribosomal protein S4.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
4	3E	208	Total	C	N	O	S	0	0	0
			1702	1066	339	290	7			
4	32	208	Total	C	N	O	S	0	0	0
			1702	1066	339	290	7			

- Molecule 5 is a protein called 30S ribosomal protein S5.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
5	4E	151	Total	C	N	O	S	0	0	0
			1155	729	218	204	4			
5	42	151	Total	C	N	O	S	0	0	0
			1155	729	218	204	4			

- Molecule 6 is a protein called 30S ribosomal protein S6.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
6	5E	101	Total	C	N	O	S	0	0	0
			842	531	155	153	3			
6	52	101	Total	C	N	O	S	0	0	0
			842	531	155	153	3			

- Molecule 7 is a protein called 30S ribosomal protein S7.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
7	6E	144	Total	C	N	O	S	0	0	0
			1157	718	230	203	6			
7	62	147	Total	C	N	O	S	0	0	0
			1200	750	237	207	6			

- Molecule 8 is a protein called 30S ribosomal protein S8.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
8	7E	138	Total	C	N	O	S	0	0	0
			1115	705	215	192	3			
8	72	138	Total	C	N	O	S	0	0	0
			1115	705	215	192	3			

- Molecule 9 is a protein called 30S ribosomal protein S9.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
9	8E	127	Total	C	N	O	0	0	0
			1009	639	197	173			
9	82	124	Total	C	N	O	0	0	0
			983	624	190	169			

- Molecule 10 is a protein called 30S ribosomal protein S10.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
10	1I	99	Total	C	N	O	S	0	0	0
			801	504	157	139	1			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
10	1A	99	Total	C	N	O	S	0	0	0
			801	504	157	139	1			

- Molecule 11 is a protein called 30S ribosomal protein S11.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
11	2I	116	Total	C	N	O	S	0	0	0
			864	537	164	160	3			
11	2A	116	Total	C	N	O	S	0	0	0
			864	537	164	160	3			

- Molecule 12 is a protein called 30S ribosomal protein S12.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
12	3I	122	Total	C	N	O	S	0	0	0
			956	603	193	159	1			
12	3A	125	Total	C	N	O	S	0	0	0
			975	614	196	164	1			

- Molecule 13 is a protein called 30S ribosomal protein S13.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
13	4I	116	Total	C	N	O	S	0	0	0
			928	574	191	161	2			
13	4A	116	Total	C	N	O	S	0	0	0
			928	574	191	161	2			

- Molecule 14 is a protein called 30S ribosomal protein S14 type Z.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
14	5I	60	Total	C	N	O	S	0	0	0
			491	312	104	71	4			
14	5A	58	Total	C	N	O	S	0	0	0
			475	303	99	69	4			

- Molecule 15 is a protein called 30S ribosomal protein S15.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
15	6I	88	Total	C	N	O	S	0	0	0
			733	459	147	125	2			
15	6A	88	Total	C	N	O	S	0	0	0
			733	459	147	125	2			

- Molecule 16 is a protein called 30S ribosomal protein S16.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
16	7I	84	Total	C	N	O	S	0	0	0
			705	446	140	118	1			
16	7A	84	Total	C	N	O	S	0	0	0
			705	446	140	118	1			

- Molecule 17 is a protein called 30S ribosomal protein S17.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
17	8I	100	Total	C	N	O	S	0	0	0
			834	534	155	143	2			
17	8A	99	Total	C	N	O	S	0	0	0
			823	528	151	142	2			

- Molecule 18 is a protein called 30S ribosomal protein S18.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
18	9I	67	Total	C	N	O	0	0	0
			550	352	107	91			
18	9A	69	Total	C	N	O	0	0	0
			564	361	110	93			

- Molecule 19 is a protein called 30S ribosomal protein S19.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
19	AI	80	Total	C	N	O	S	0	0	0
			643	411	118	112	2			
19	AA	78	Total	C	N	O	S	0	0	0
			624	398	115	109	2			

- Molecule 20 is a protein called 30S ribosomal protein S20.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
20	BI	99	Total	C	N	O	S	0	0	0
			762	470	162	128	2			
20	BA	99	Total	C	N	O	S	0	0	0
			762	470	162	128	2			

- Molecule 21 is a protein called 30S ribosomal protein Thx.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
21	1F	23	Total	C	N	O	0	0	0
			199	122	48	29			
21	1B	25	Total	C	N	O	0	0	0
			217	134	52	31			

- Molecule 22 is a RNA chain called tRNA-Leu.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
22	1K	87	Total	C	N	O	P	0	0	0
			1863	831	333	612	87			
22	1L	87	Total	C	N	O	P	0	0	0
			1863	831	333	612	87			

- Molecule 23 is a RNA chain called tRNA-Leu.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
23	2K	79	Total	C	N	O	P	0	0	0
			1692	755	300	558	79			
23	2L	80	Total	C	N	O	P	0	0	0
			1712	764	303	565	80			

- Molecule 24 is a RNA chain called tRNA-Leu.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
24	3K	77	Total	C	N	O	P	0	0	0
			1646	733	292	544	77			
24	3L	75	Total	C	N	O	P	0	0	0
			1601	713	282	531	75			

- Molecule 25 is a RNA chain called mRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
25	4K	16	Total	C	N	O	P	0	0	0
			336	150	50	120	16			
25	4L	21	Total	C	N	O	P	0	0	0
			438	196	64	157	21			

- Molecule 26 is a RNA chain called 23S ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
26	1H	2912	Total	C	N	O	P	0	0	0
			62707	27911	11722	20163	2911			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
26	14	2909	Total	C	N	O	P	0	0	0
			62647	27884	11716	20139	2908			

There are 6 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
1H	161	U	-	insertion	GB 48268
1H	493	G	-	insertion	GB 48268
1H	1228	G	-	insertion	GB 48268
14	161	U	-	insertion	GB 48268
14	493	G	-	insertion	GB 48268
14	1228	G	-	insertion	GB 48268

- Molecule 27 is a RNA chain called 5S ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
27	16	122	Total	C	N	O	P	0	0	0
			2617	1166	486	844	121			
27	1J	122	Total	C	N	O	P	0	0	0
			2617	1166	486	844	121			

- Molecule 28 is a protein called 50S ribosomal protein L2.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
28	11	273	Total	C	N	O	S	0	0	0
			2126	1341	424	358	3			
28	19	273	Total	C	N	O	S	0	0	0
			2120	1338	421	358	3			

- Molecule 29 is a protein called 50S ribosomal protein L3.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
29	21	205	Total	C	N	O	S	0	0	0
			1568	991	300	271	6			
29	29	205	Total	C	N	O	S	0	0	0
			1568	991	300	271	6			

- Molecule 30 is a protein called 50S ribosomal protein L4.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
30	31	202	Total	C	N	O	S	0	0	0
			1585	1011	297	275	2			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
30	39	208	Total	C	N	O	S	0	0	0
			1627	1037	304	283	3			

- Molecule 31 is a protein called 50S ribosomal protein L5.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
31	41	181	Total	C	N	O	S	0	0	0
			1473	942	268	259	4			
31	49	181	Total	C	N	O	S	0	0	0
			1473	942	268	259	4			

- Molecule 32 is a protein called 50S ribosomal protein L6.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
32	51	174	Total	C	N	O	S	0	0	0
			1336	848	251	236	1			
32	59	170	Total	C	N	O	S	0	0	0
			1307	829	245	232	1			

- Molecule 33 is a protein called 50S ribosomal protein L9.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
33	61	146	Total	C	N	O	S	0	0	0
			1136	726	201	208	1			
33	69	146	Total	C	N	O	S	0	0	0
			1136	726	201	208	1			

- Molecule 34 is a protein called 50S ribosomal protein L13.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
34	58	138	Total	C	N	O	S	0	0	0
			1104	712	206	182	4			
34	15	138	Total	C	N	O	S	0	0	0
			1104	712	206	182	4			

- Molecule 35 is a protein called 50S ribosomal protein L14.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
35	68	122	Total	C	N	O	S	0	0	0
			932	588	171	169	4			
35	25	122	Total	C	N	O	S	0	0	0
			932	588	171	169	4			



- Molecule 36 is a protein called 50S ribosomal protein L15.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
36	78	147	Total	C	N	O	S	0	0	0
			1122	698	229	192	3			
36	35	150	Total	C	N	O	S	0	0	0
			1144	712	232	197	3			

- Molecule 37 is a protein called 50S ribosomal protein L16.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
37	88	141	Total	C	N	O	S	0	0	0
			1121	715	212	187	7			
37	45	140	Total	C	N	O	S	0	0	0
			1113	710	211	186	6			

- Molecule 38 is a protein called 50S ribosomal protein L17.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
38	98	118	Total	C	N	O	S	0	0	0
			967	604	203	159	1			
38	55	117	Total	C	N	O		0	0	0
			959	599	202	158				

- Molecule 39 is a protein called 50S ribosomal protein L18.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
39	A8	111	Total	C	N	O	0	0	0
			881	556	176	149			
39	65	111	Total	C	N	O	0	0	0
			881	556	176	149			

- Molecule 40 is a protein called 50S ribosomal protein L19.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
40	B8	129	Total	C	N	O	S	0	0	0
			1081	674	223	183	1			
40	75	137	Total	C	N	O	S	0	0	0
			1141	710	234	196	1			

- Molecule 41 is a protein called 50S ribosomal protein L20.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
41	C8	117	Total	C	N	O	S	0	0	0
			963	610	202	150	1			
41	85	117	Total	C	N	O	S	0	0	0
			963	610	202	150	1			

- Molecule 42 is a protein called 50S ribosomal protein L21.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
42	D8	101	Total	C	N	O	S	0	0	0
			778	501	142	134	1			
42	95	101	Total	C	N	O	S	0	0	0
			778	501	142	134	1			

- Molecule 43 is a protein called 50S ribosomal protein L22.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
43	E8	112	Total	C	N	O	S	0	0	0
			890	560	175	153	2			
43	A5	113	Total	C	N	O	S	0	0	0
			899	566	177	154	2			

- Molecule 44 is a protein called 50S ribosomal protein L23.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
44	F8	94	Total	C	N	O	S	0	0	0
			742	482	134	125	1			
44	B5	92	Total	C	N	O		0	0	0
			725	471	131	123				

- Molecule 45 is a protein called 50S ribosomal protein L24.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
45	G8	103	Total	C	N	O	S	0	0	0
			783	504	148	126	5			
45	C5	104	Total	C	N	O	S	0	0	0
			794	510	152	127	5			

- Molecule 46 is a protein called 50S ribosomal protein L25.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
46	H8	171	Total	C	N	O	S	0	0	0
			1373	876	247	247	3			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
46	D5	135	Total	C	N	O	S	0	0	0
			1120	720	202	195	3			

- Molecule 47 is a protein called 50S ribosomal protein L27.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
47	I8	83	Total	C	N	O	S	0	0	0
			656	407	139	109	1			
47	E5	80	Total	C	N	O	S	0	0	0
			627	388	132	106	1			

There are 4 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
I8	6	ALA	GLY	conflict	UNP P60493
I8	8	ALA	GLY	conflict	UNP P60493
E5	6	ALA	GLY	conflict	UNP P60493
E5	8	ALA	GLY	conflict	UNP P60493

- Molecule 48 is a protein called 50S ribosomal protein L28.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
48	J8	95	Total	C	N	O	S	0	0	0
			746	469	148	128	1			
48	F5	94	Total	C	N	O	S	0	0	0
			737	463	146	127	1			

- Molecule 49 is a protein called 50S ribosomal protein L29.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
49	K8	68	Total	C	N	O	S	0	0	0
			575	358	116	100	1			
49	G5	67	Total	C	N	O	S	0	0	0
			567	351	115	100	1			

- Molecule 50 is a protein called 50S ribosomal protein L30.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
50	L8	57	Total	C	N	O	0	0	0
			452	288	88	76			
50	H5	59	Total	C	N	O	0	0	0
			468	298	90	80			

- Molecule 51 is a protein called 50S ribosomal protein L31.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
51	M8	66	Total	C	N	O	S	0	0	0
			533	335	96	97	5			
51	I5	63	Total	C	N	O	S	0	0	0
			515	326	93	91	5			

- Molecule 52 is a protein called 50S ribosomal protein L32.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
52	N8	55	Total	C	N	O	S	0	0	0
			429	269	86	69	5			
52	J5	56	Total	C	N	O	S	0	0	0
			434	272	87	70	5			

- Molecule 53 is a protein called 50S ribosomal protein L34.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
53	P8	47	Total	C	N	O	S	0	0	0
			409	251	102	54	2			
53	L5	47	Total	C	N	O	S	0	0	0
			409	251	102	54	2			

- Molecule 54 is a protein called 50S ribosomal protein L35.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
54	Q8	60	Total	C	N	O	S	0	0	0
			480	306	98	74	2			
54	M5	53	Total	C	N	O	S	0	0	0
			422	270	87	63	2			

- Molecule 55 is MAGNESIUM ION (three-letter code: MG) (formula: Mg).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
55	98	1	Total	Mg	0	0
			1	1		
55	45	2	Total	Mg	0	0
			2	2		
55	P8	1	Total	Mg	0	0
			1	1		
55	85	2	Total	Mg	0	0
			2	2		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
55	32	1	Total 1	Mg 1	0	0
55	13	146	Total 146	Mg 146	0	0
55	1J	2	Total 2	Mg 2	0	0
55	16	5	Total 5	Mg 5	0	0
55	25	1	Total 1	Mg 1	0	0
55	21	2	Total 2	Mg 2	0	0
55	31	1	Total 1	Mg 1	0	0
55	3I	1	Total 1	Mg 1	0	0
55	I8	2	Total 2	Mg 2	0	0
55	5E	1	Total 1	Mg 1	0	0
55	29	3	Total 3	Mg 3	0	0
55	7A	1	Total 1	Mg 1	0	0
55	2K	3	Total 3	Mg 3	0	0
55	J8	1	Total 1	Mg 1	0	0
55	1G	143	Total 143	Mg 143	0	0
55	11	1	Total 1	Mg 1	0	0
55	1H	512	Total 512	Mg 512	0	0
55	E5	2	Total 2	Mg 2	0	0
55	88	4	Total 4	Mg 4	0	0
55	14	490	Total 490	Mg 490	0	0
55	78	1	Total 1	Mg 1	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
55	3E	1	Total 1	Mg 1	0	0
55	1K	1	Total 1	Mg 1	0	0
55	G8	1	Total 1	Mg 1	0	0
55	2L	2	Total 2	Mg 2	0	0

- Molecule 56 is ZINC ION (three-letter code: ZN) (formula: Zn).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	32	1	Total 1	Zn 1	0	0
56	3E	1	Total 1	Zn 1	0	0
56	5I	1	Total 1	Zn 1	0	0
56	5A	1	Total 1	Zn 1	0	0
56	G8	1	Total 1	Zn 1	0	0
56	C5	1	Total 1	Zn 1	0	0

- Molecule 57 is water.

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	13	74	Total 74	O 74	0	0
57	3E	1	Total 1	O 1	0	0
57	6I	1	Total 1	O 1	0	0
57	BI	1	Total 1	O 1	0	0
57	1H	552	Total 552	O 552	0	0
57	11	1	Total 1	O 1	0	0
57	21	1	Total 1	O 1	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	31	4	Total 4	O 4	0	0
57	78	2	Total 2	O 2	0	0
57	B8	1	Total 1	O 1	0	0
57	C8	2	Total 2	O 2	0	0
57	D8	1	Total 1	O 1	0	0
57	E8	1	Total 1	O 1	0	0
57	I8	3	Total 3	O 3	0	0
57	J8	1	Total 1	O 1	0	0
57	P8	1	Total 1	O 1	0	0
57	1G	73	Total 73	O 73	0	0
57	6A	2	Total 2	O 2	0	0
57	7A	1	Total 1	O 1	0	0
57	4L	2	Total 2	O 2	0	0
57	14	520	Total 520	O 520	0	0
57	19	8	Total 8	O 8	0	0
57	29	2	Total 2	O 2	0	0
57	39	4	Total 4	O 4	0	0
57	35	1	Total 1	O 1	0	0
57	55	2	Total 2	O 2	0	0
57	85	1	Total 1	O 1	0	0
57	A5	1	Total 1	O 1	0	0

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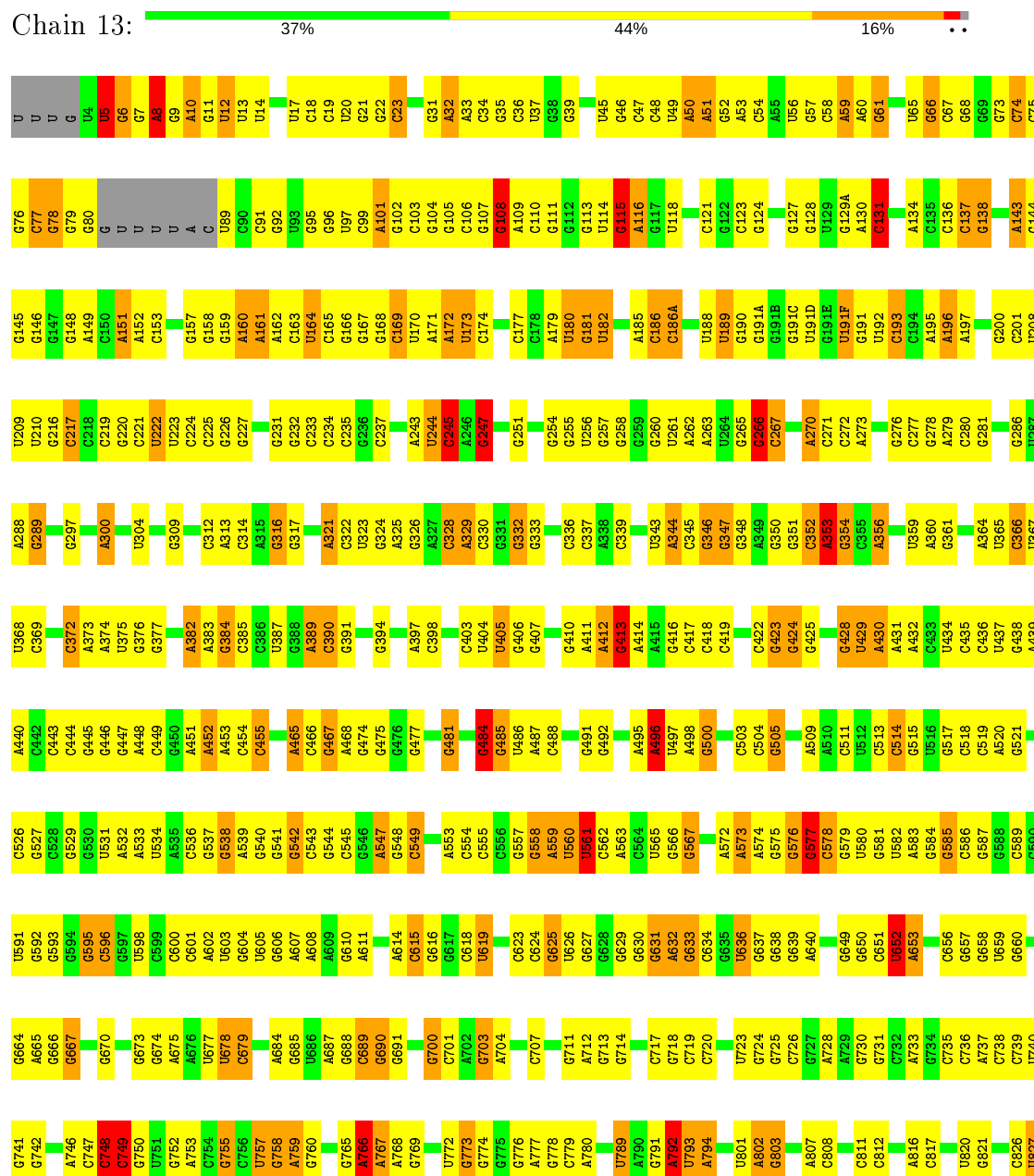
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	M5	3	Total	O	0	0
			3	3		



### 3 Residue-property plots

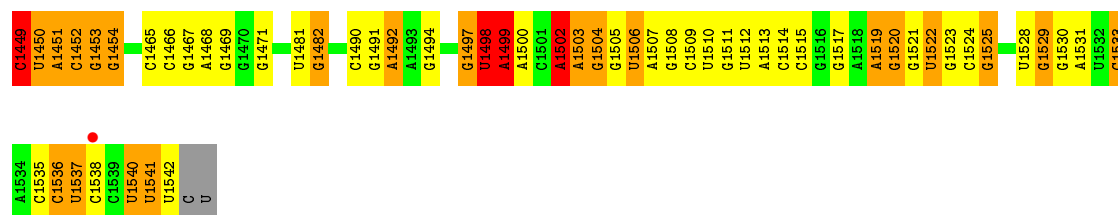
These plots are drawn for all protein, RNA and DNA chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and electron density. Residues are 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: 16S ribosomal RNA

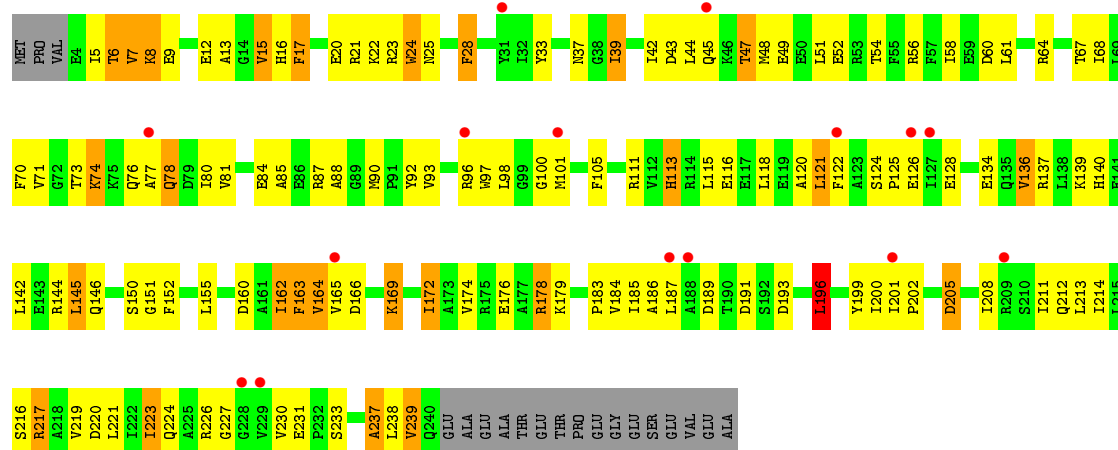




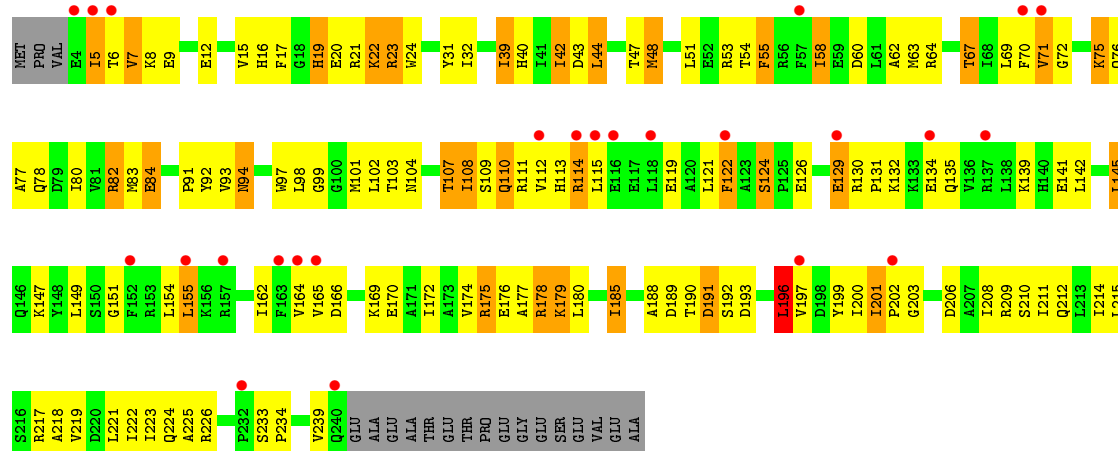
A1357	A1358	C1359	A1289	G1290	G1291	G1215	G1216	G1217	C1218	U1219	A1151	A1152	C1153	G1154	G1155	A1156	A1157	C1158	U1159	G1160	C1161	G1171	C1172	G1173	A1174	G1175	A1176	G1177	G1178	A1179	A1180	G1181	C1182	A1183	G1184	G1185	G1186	A1187	C1188	G1189	G1190	A1191	C1192	G1193	U1196	G1197	G1198	G1199	C1200	G1201	G1202	G1203	G1204	U1205	U1206	A1207	C1208	C1209	G1210	U1211	A1285	A1286	A1287	C1282	A1288																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																	
A1363	A1364	G1365	G1294	A1295	C1296	C1297	G1222	G1223	G1224	C1225	C1226	A1227	A1236	C1237	A1238	U1239	U1240	G1241	A1245	C1246	A1250	A1251	A1252	G1253	A1254	C1255	G1256	A1257	U1258	G1259	C1260	A1261	G1262	G1263	C1264	G1265	A1268	A1269	G1270	G1271	G1272	G1273	G1274	A1275	G1276	C1277	U1278	A1279	A1280	U1281	C1282	A1285	A1286	A1287	C1282	A1288																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																										
A1366	G1367	A1370	A1371	U1372	U1373	G1305	A1306	U1307	U1308	U1309	G1312	U1313	C1314	U1315	G1316	A1317	A1318	A1319	C1320	C1321	C1322	G1323	A1324	C1325	G1326	C1327	A1328	A1329	U1330	G1331	C1335	C1336	G1337	G1338	A1339	U1340	G1341	G1342	G1343	C1344	U1345	A1346	G1347	U1348	C1352	G1353	C1354	A1355	C1356																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																	
A1368	A1369	A1370	A1371	U1372	U1373	G1305	A1306	U1307	U1308	U1309	G1312	U1313	C1314	U1315	G1316	A1317	A1318	A1319	C1320	C1321	C1322	G1323	A1324	C1325	G1326	C1327	A1328	A1329	U1330	G1331	C1335	C1336	G1337	G1338	A1339	U1340	G1341	G1342	G1343	C1344	U1345	A1346	G1347	U1348	C1352	G1353	C1354	A1355	C1356																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																	
A1371	A1372	A1373	A1374	A1375	G1376	G1377	G1378	G1379	G1380	G1381	G1382	G1383	G1384	G1385	G1386	G1387	G1388	G1389	U1390	U1391	G1392	C1393	A1394	C1395	G1396	G1397	A1398	C1399	G1400	G1401	C1402	C1403	C1411	C1412	A1413	G1414	G1415	G1416	G1417	A1418	G1419	G1423	G1424	U1425	G1435	U1436	C1437	G1442	G1443	A1446	G1447	C1448																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																														
A1374	A1375	G1376	G1377	G1378	G1379	G1380	G1381	G1382	G1383	G1384	G1385	G1386	G1387	G1388	G1389	U1390	U1391	G1392	C1393	A1394	C1395	G1396	G1397	A1398	C1399	G1400	G1401	C1402	C1403	C1411	C1412	A1413	G1414	G1415	G1416	G1417	A1418	G1419	G1423	G1424	U1425	G1435	U1436	C1437	G1442	G1443	A1446	G1447	C1448																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																	
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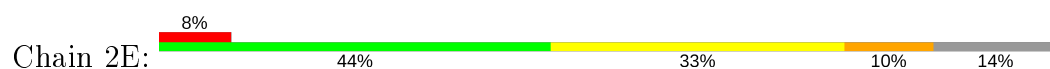
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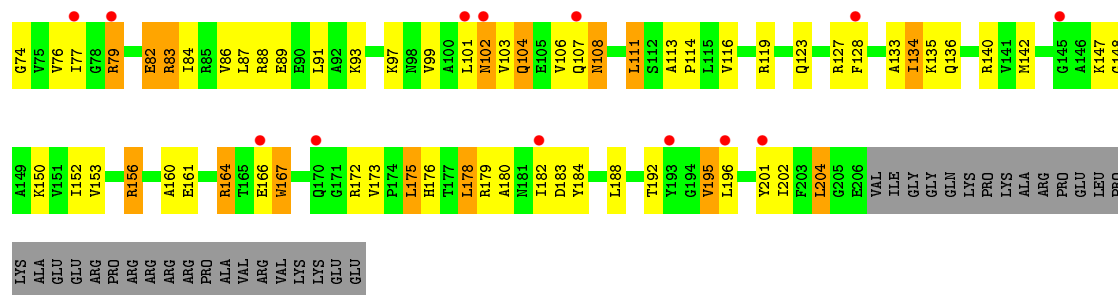


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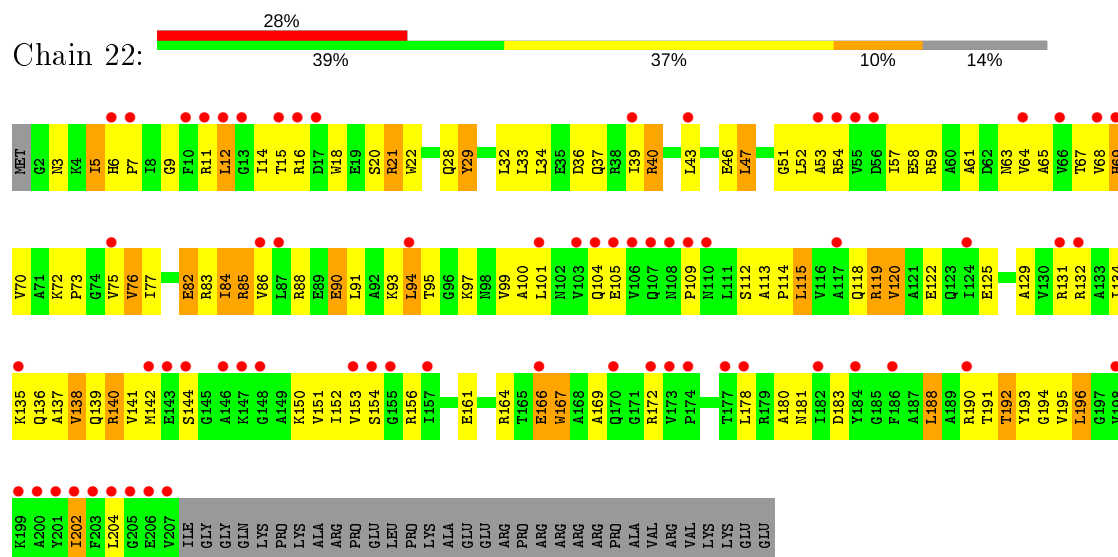


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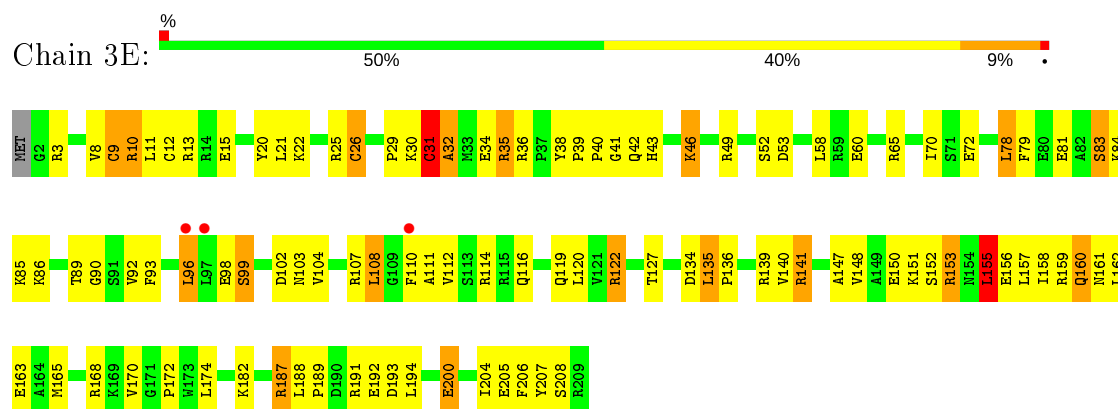




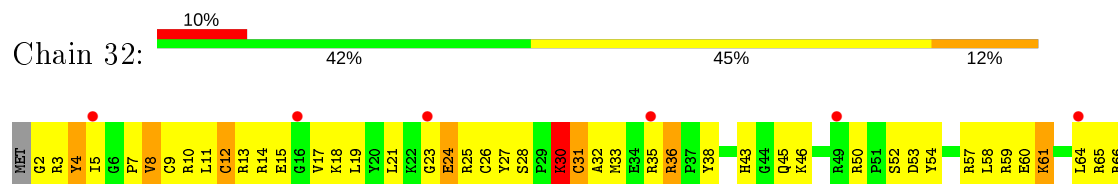
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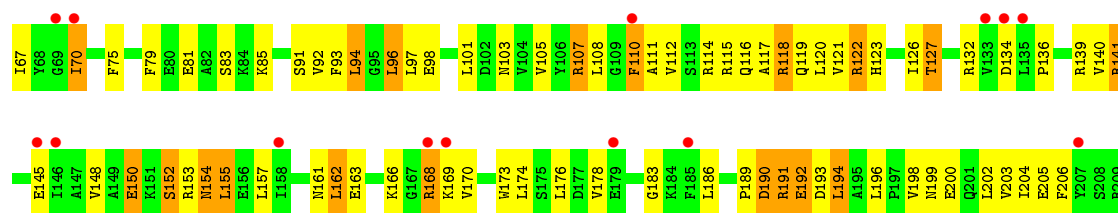


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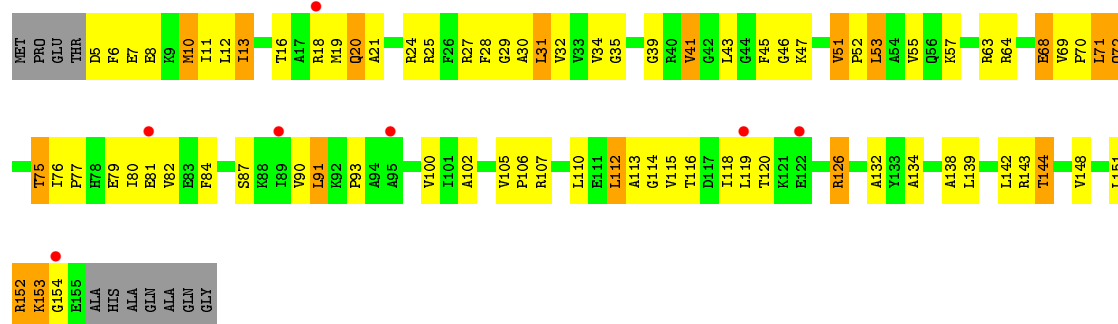
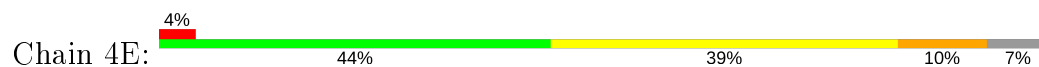


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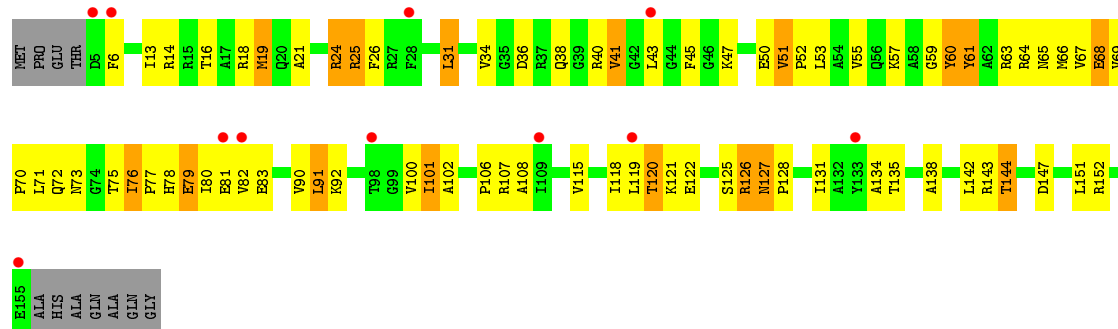




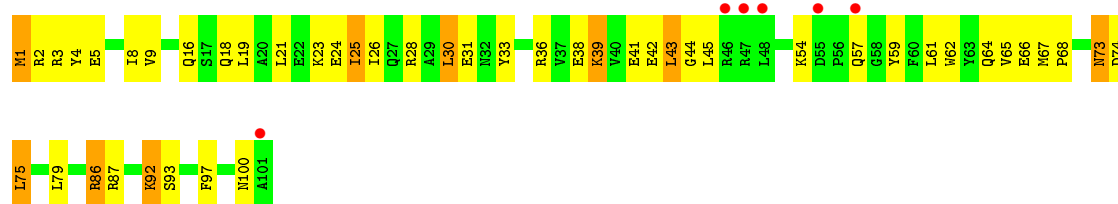
• Molecule 5: 30S ribosomal protein S5



• Molecule 5: 30S ribosomal protein S5

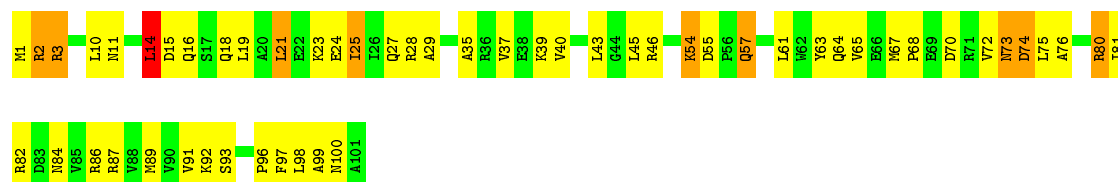


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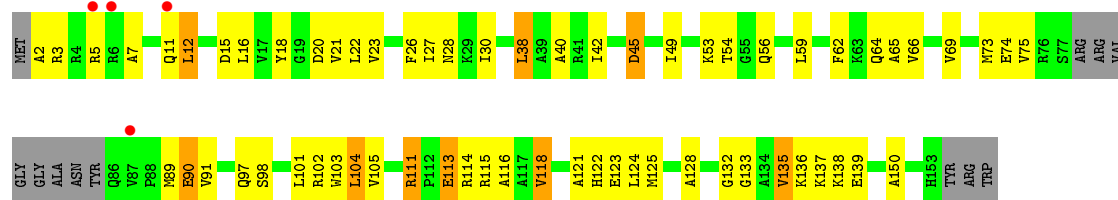


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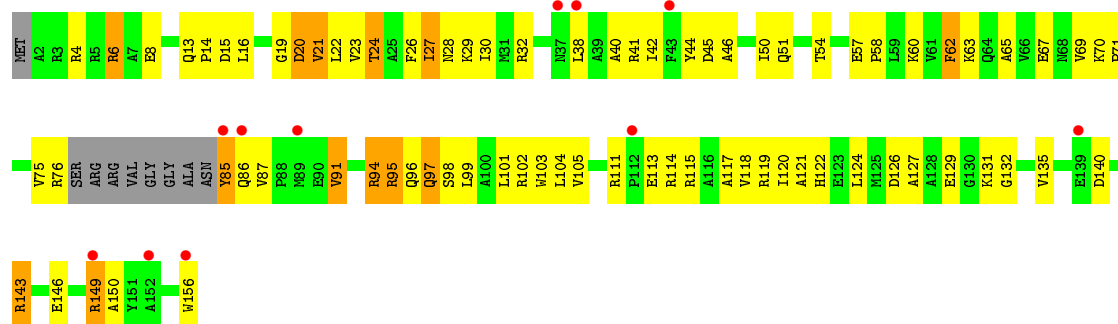
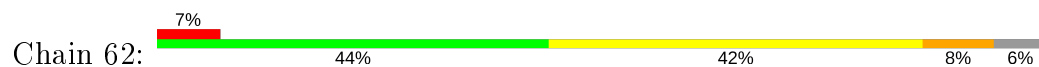




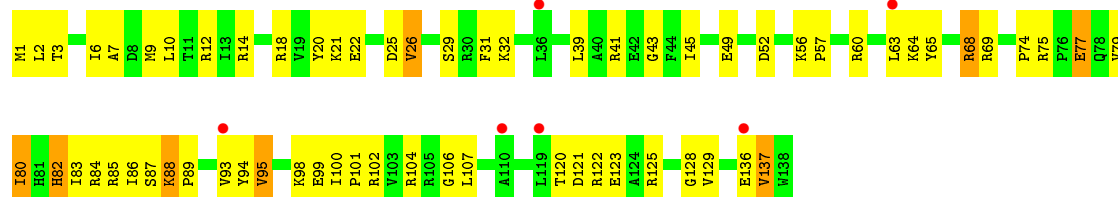
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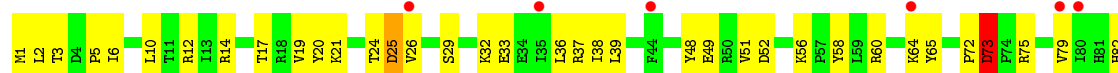
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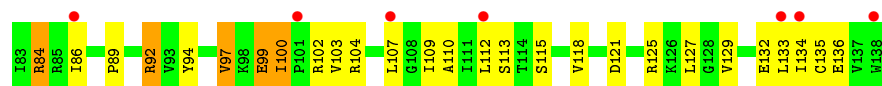


- Molecule 8: 30S ribosomal protein S8

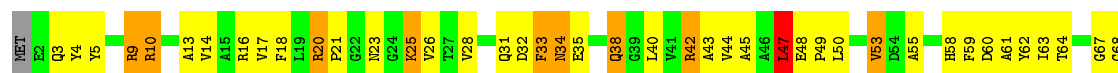


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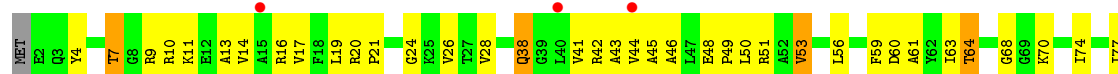




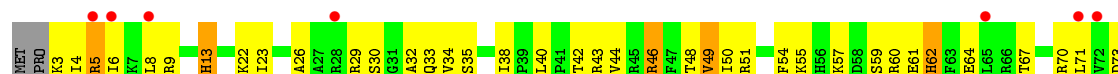
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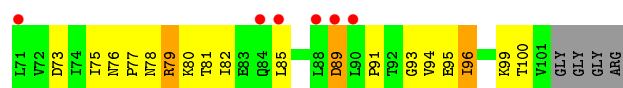
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- Molecule 10: 30S ribosomal protein S10



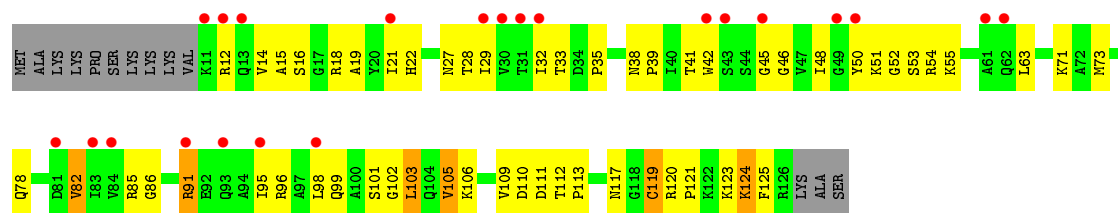
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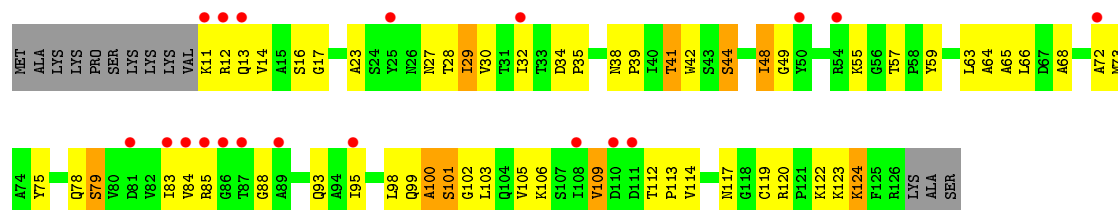
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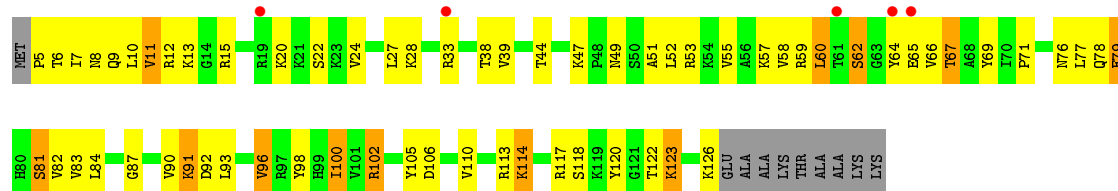
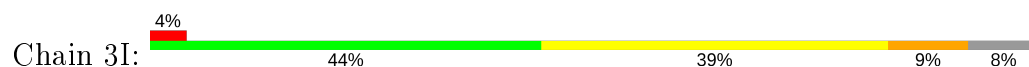




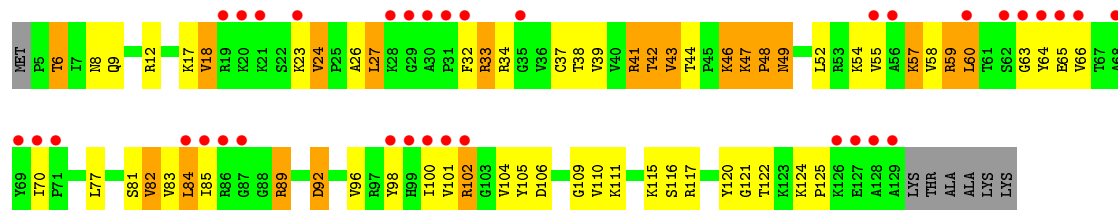
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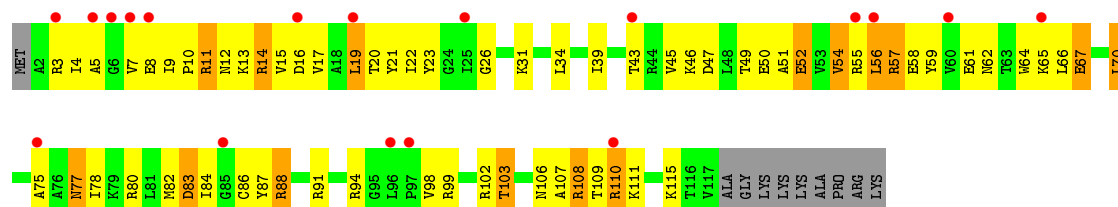
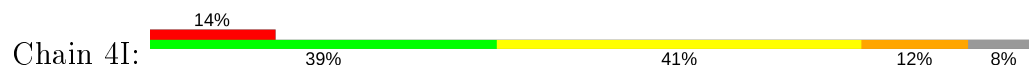
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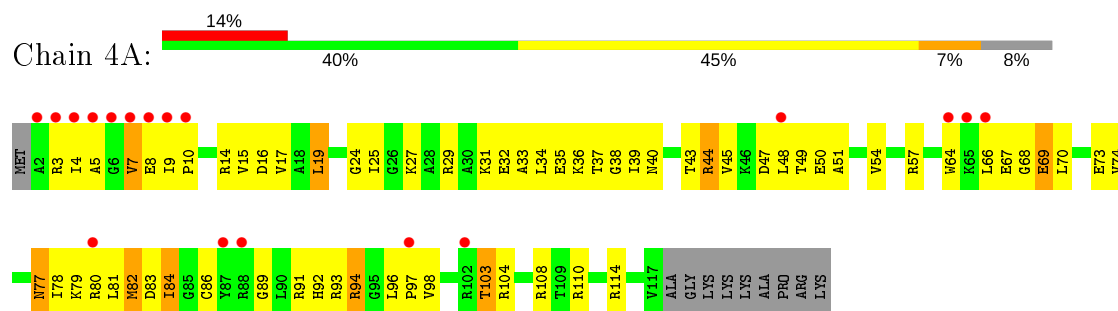
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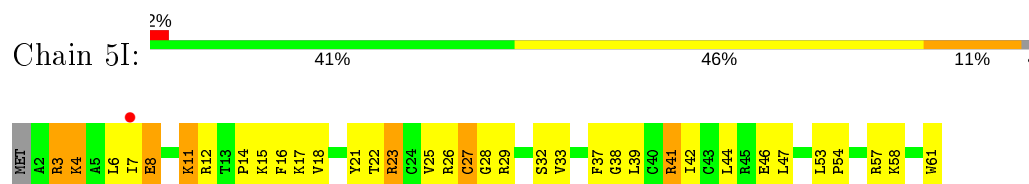
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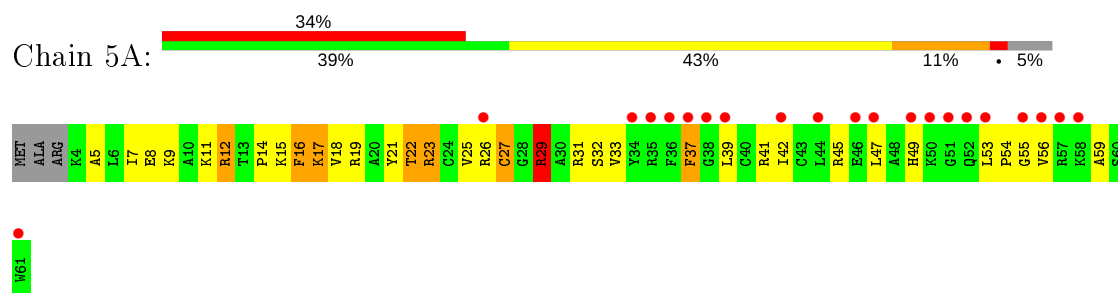
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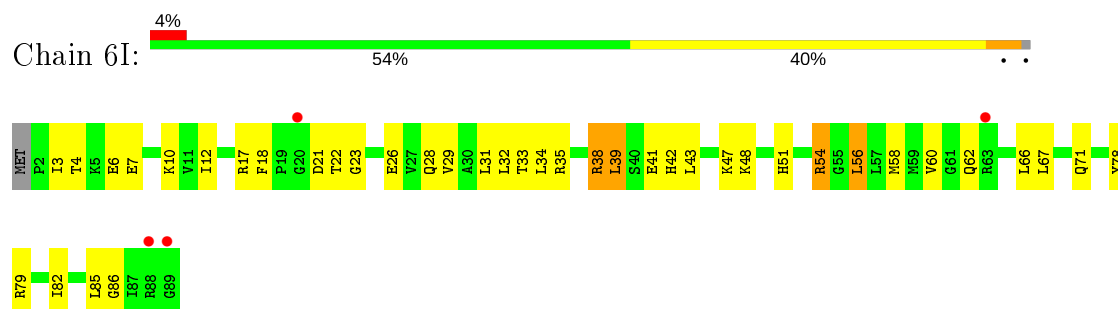
- Molecule 14: 30S ribosomal protein S14 type Z



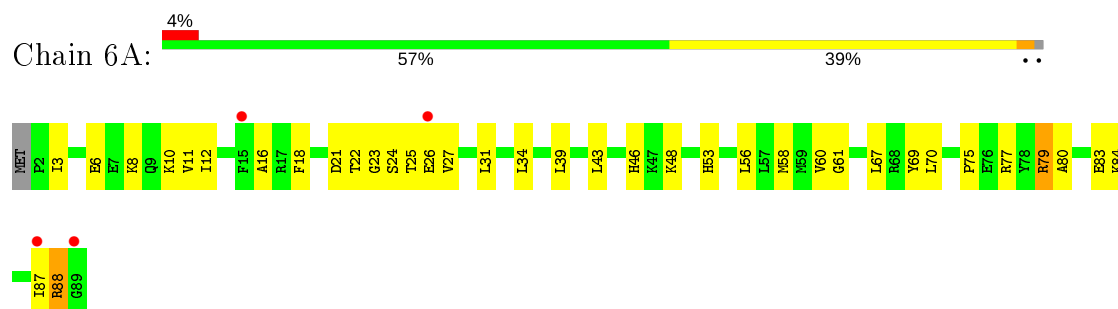
- Molecule 14: 30S ribosomal protein S14 type Z



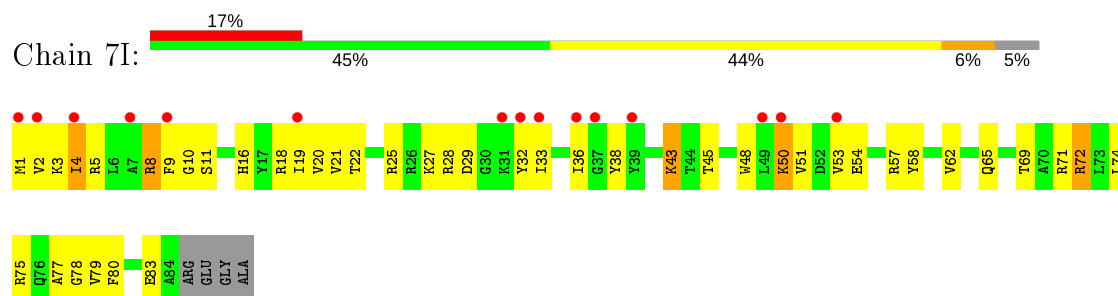
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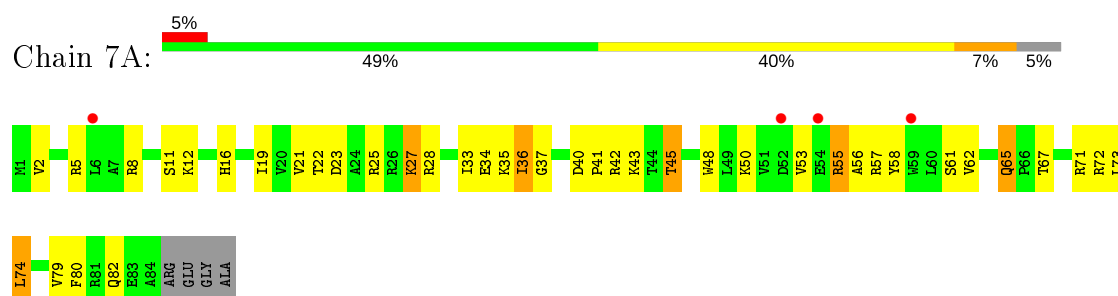
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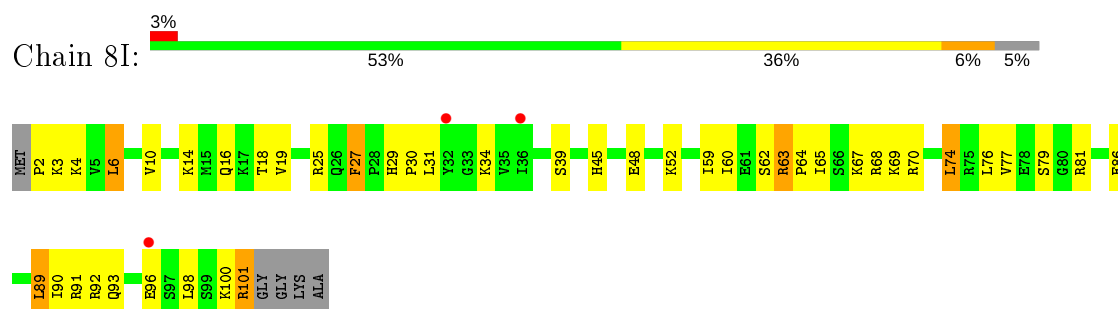
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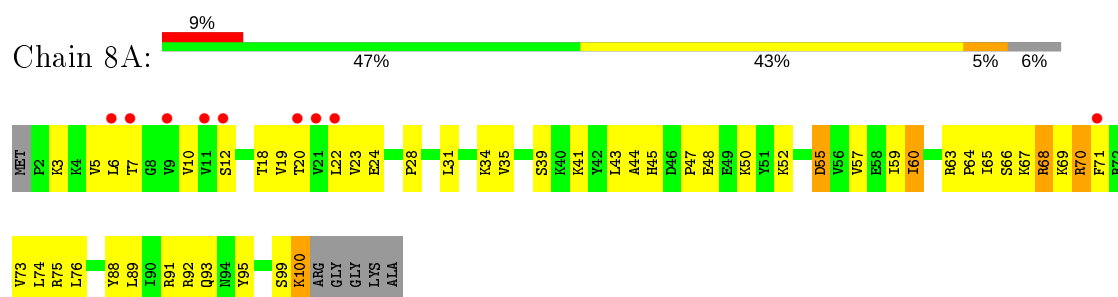
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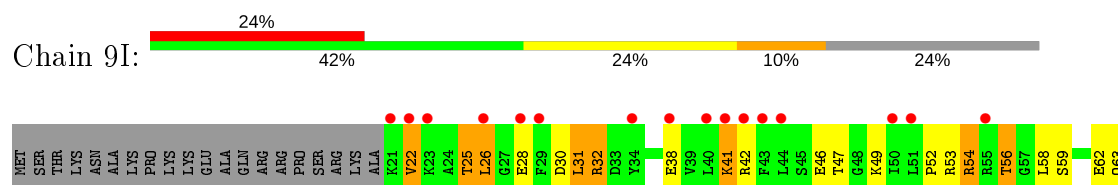
- Molecule 17: 30S ribosomal protein S17

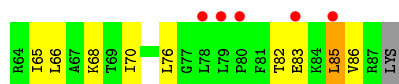


- Molecule 17: 30S ribosomal protein S17

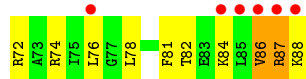
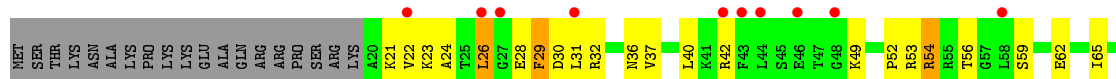
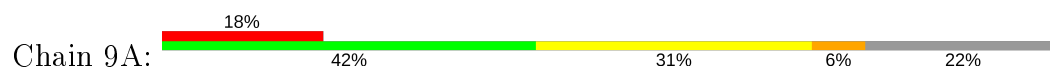


- Molecule 18: 30S ribosomal protein S18

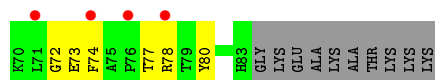
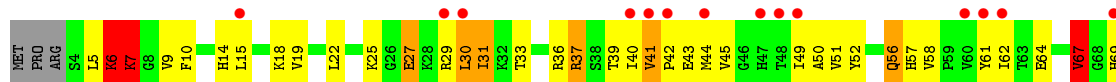




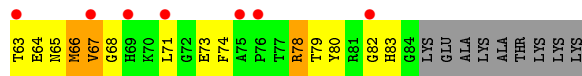
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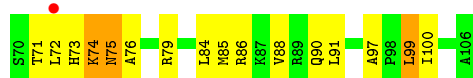
• Molecule 19: 30S ribosomal protein S19



• Molecule 19: 30S ribosomal protein S19

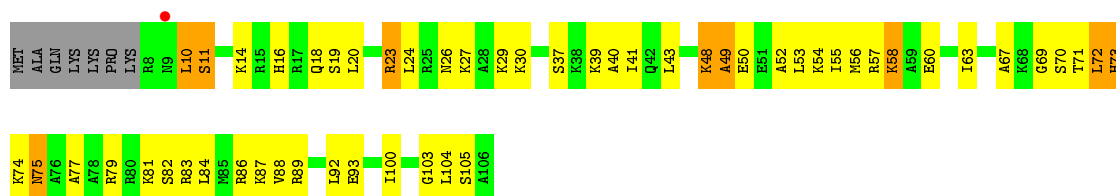


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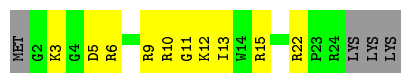


• Molecule 20: 30S ribosomal protein S20

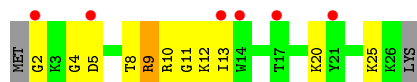




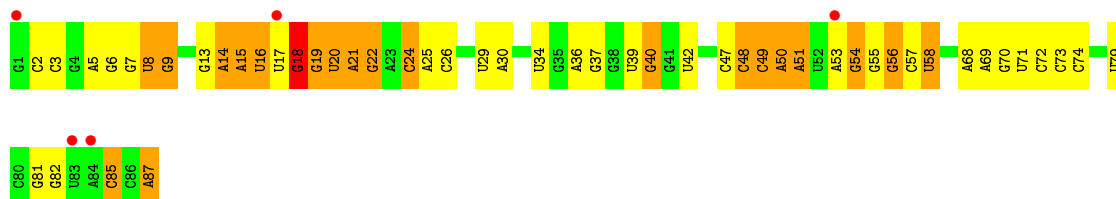
- Molecule 21: 30S ribosomal protein Thx



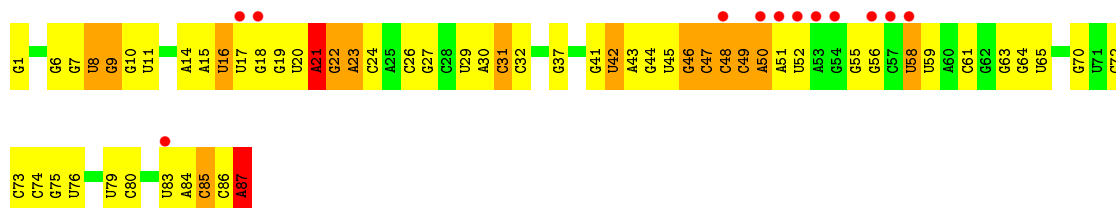
- Molecule 21: 30S ribosomal protein Thx



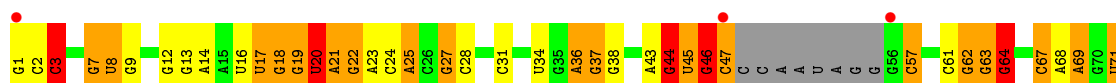
- Molecule 22: tRNA-Leu



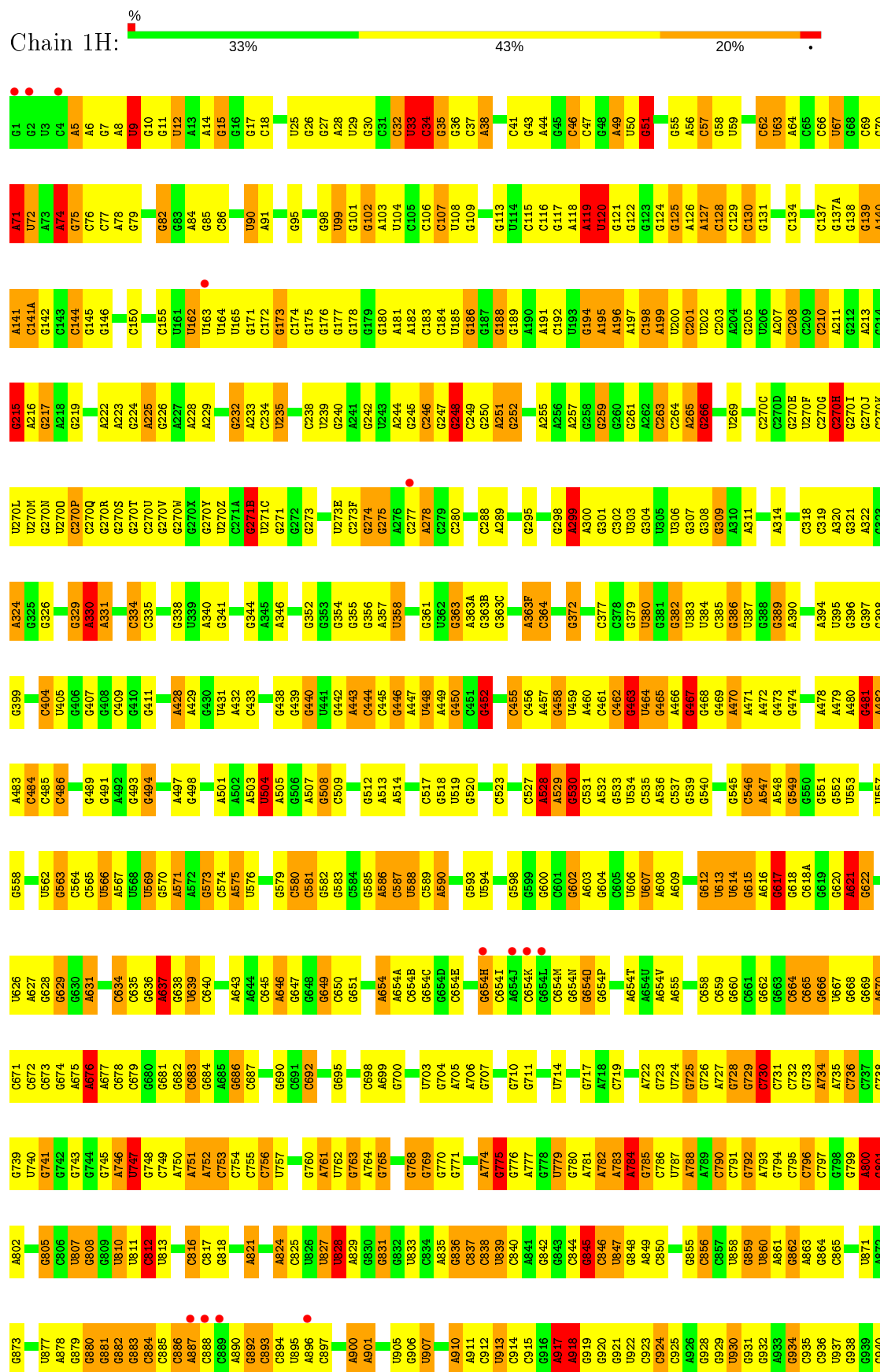
- Molecule 22: tRNA-Leu



- Molecule 23: tRNA-Leu



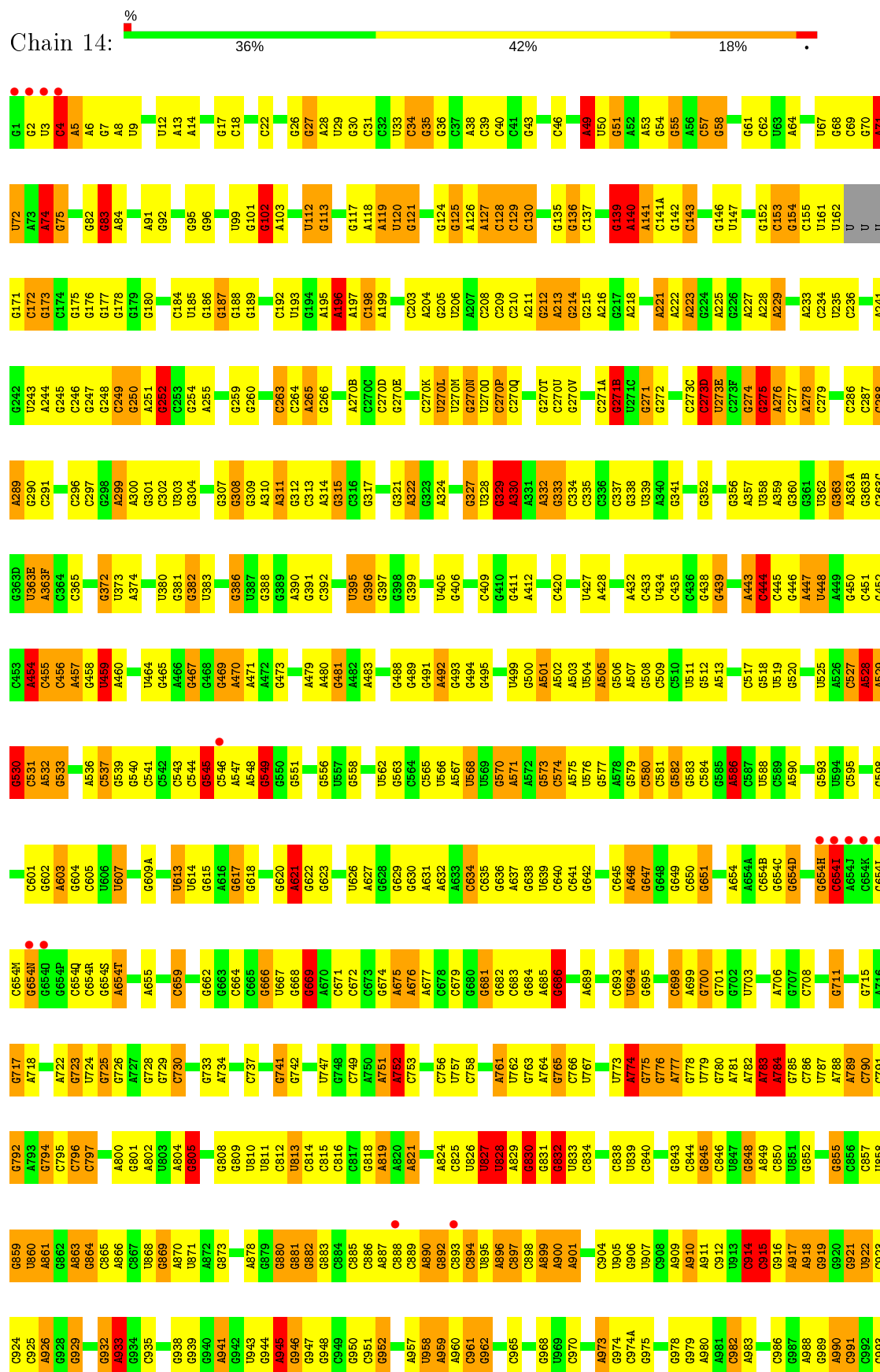




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G2011	C1930	G1842	G1681	A1609	A1610	G1543	C1476	G1343	U1273	G1190	U1113	C1049
G2012	A1931	A1843	G1682	C1611	C1611	C1544	G1479	C1345	A1274	G1193	G1114	A1050
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G2015	C1934	A1836	C1614	C1614	C1614	C1549	U1420	A1349	A1278	G1196	G1122	A1053
G2016	A1935	A1837	C1615	C1615	C1615	C1550	G1482	U1420	G1279	U1198	A1126	G1055
G2017	C1936	A1838	C1616	C1616	C1616	C1551	G1483	U1421	G1280	C1200	A1127	G1056
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G2030	C1949	C1870	A1784	C1616	C1616	C1551	G1483	U1421				
G2031	A1950	A1871	A1785	C1617	C1617	C1552	G1484	G1422				
G2032	U1951	A1872	A1786	C1618	C1618	C1553	G1485	G1423				
G2033	U1952	A1873	A1787	C1619	C1619	C1554	G1486	G1424				
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- Molecule 27: 5S ribosomal RNA

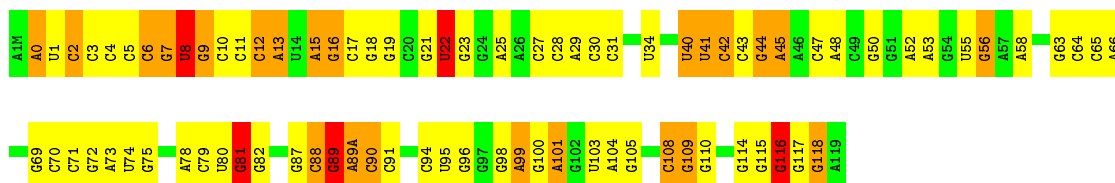
Response	Percentage
Yes	36%
No	45%
Don't know	17%





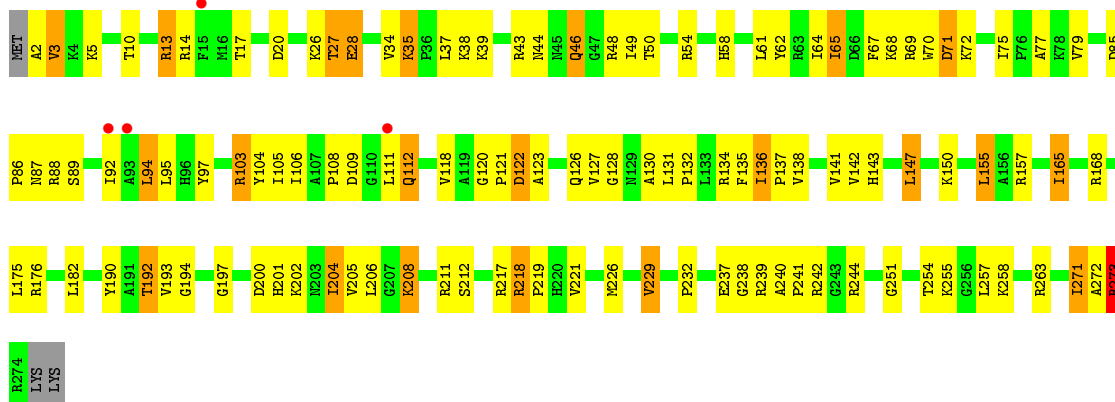
- Molecule 27: 5S ribosomal RNA

Chain 1J: 32% 45% 19%



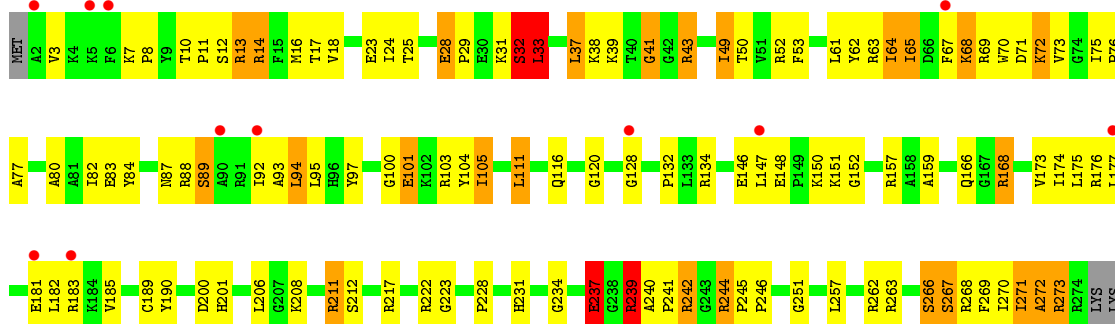
- Molecule 28: 50S ribosomal protein L2

Chain 11: 56% 35% 8%



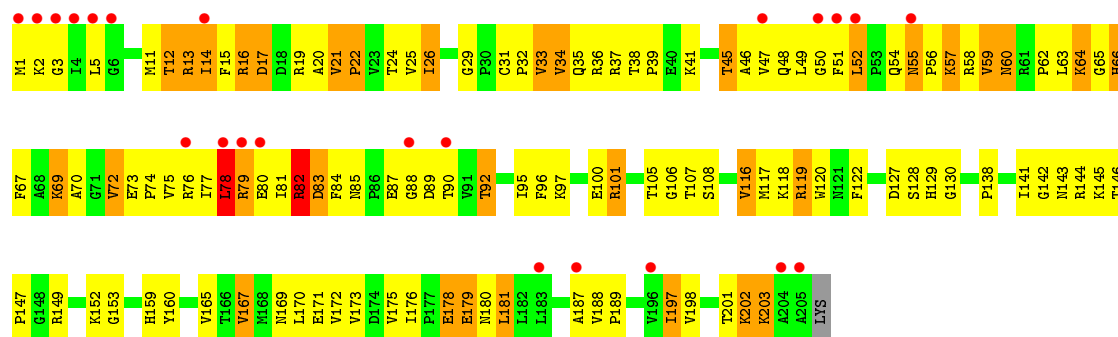
- Molecule 28: 50S ribosomal protein L2

Chain 19: 56% 33% 9%

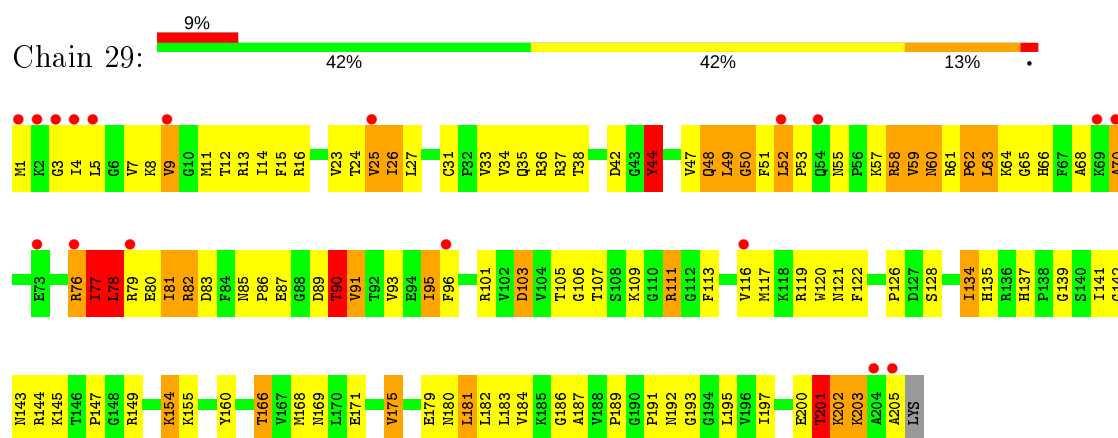


- Molecule 29: 50S ribosomal protein L3

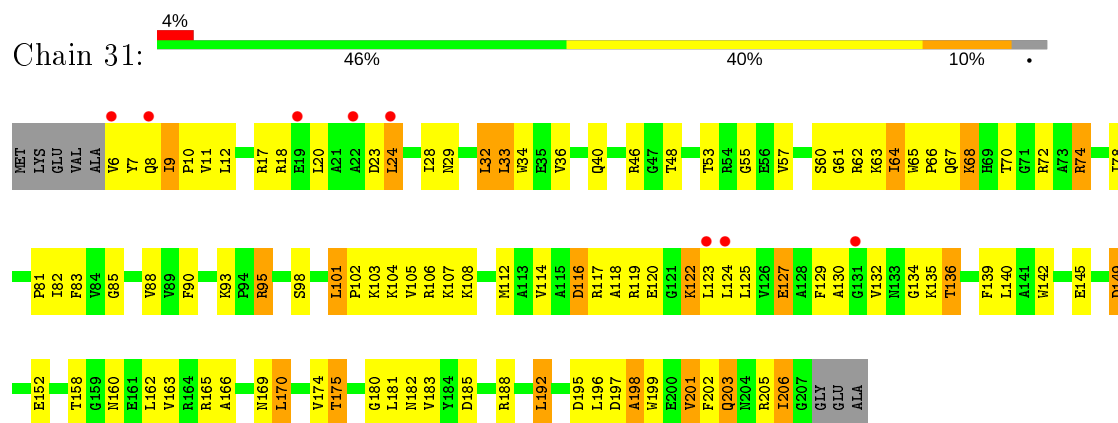
Chain 21: 39% 43% 16%



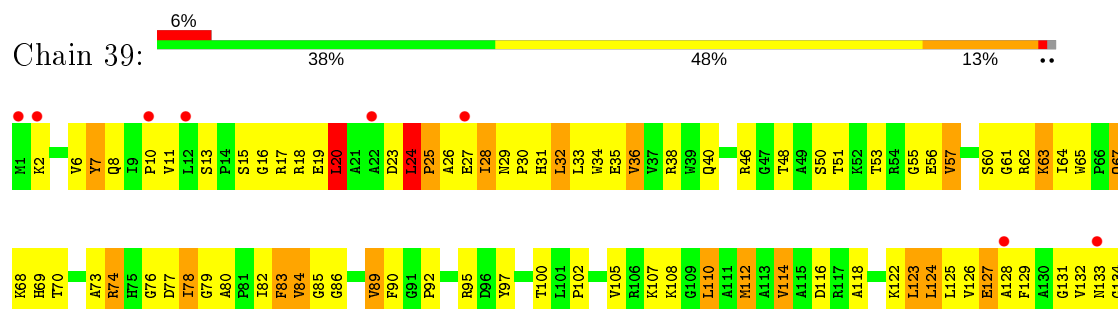
• Molecule 29: 50S ribosomal protein L3

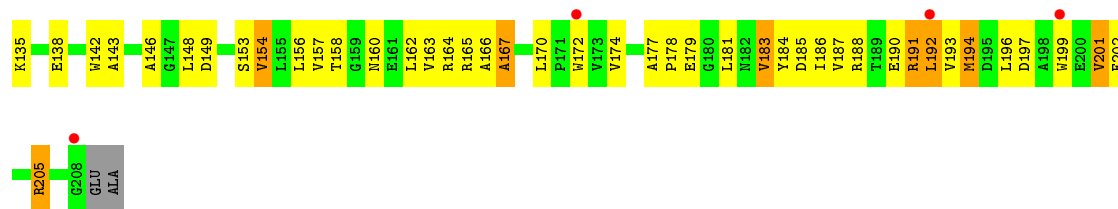


• Molecule 30: 50S ribosomal protein L4

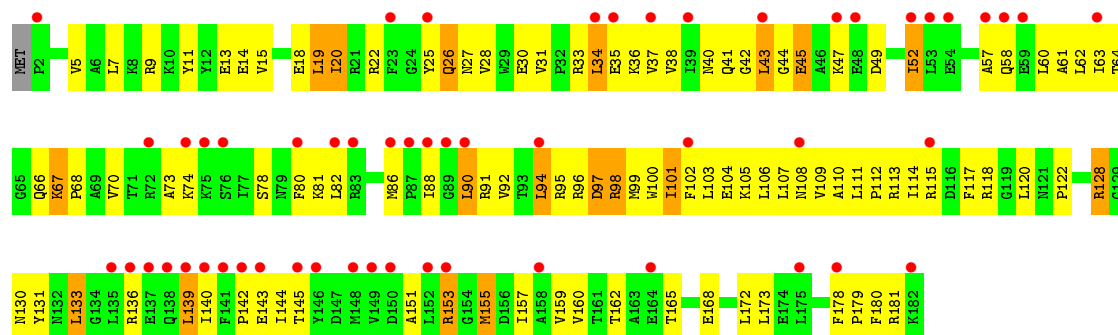


• Molecule 30: 50S ribosomal protein L4

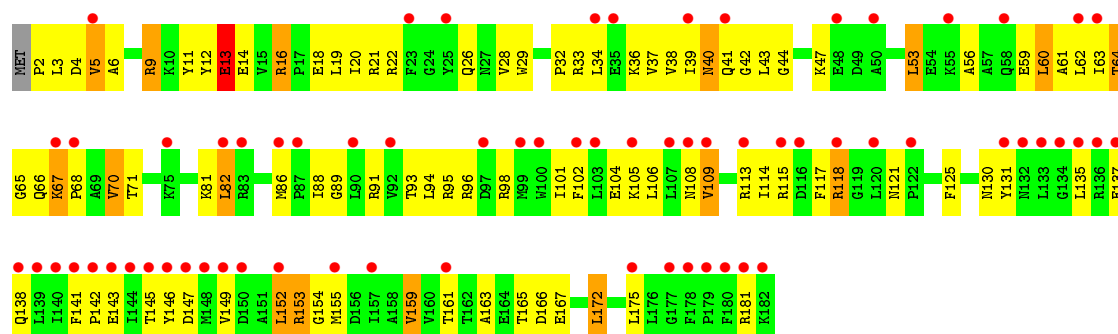




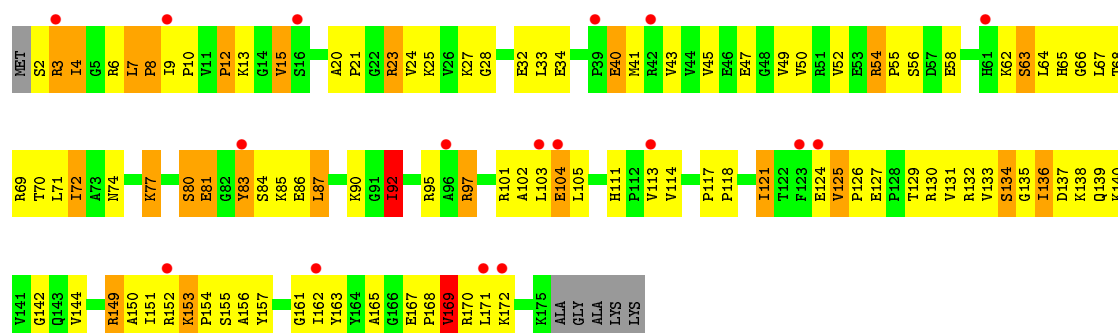
• Molecule 31: 50S ribosomal protein L5



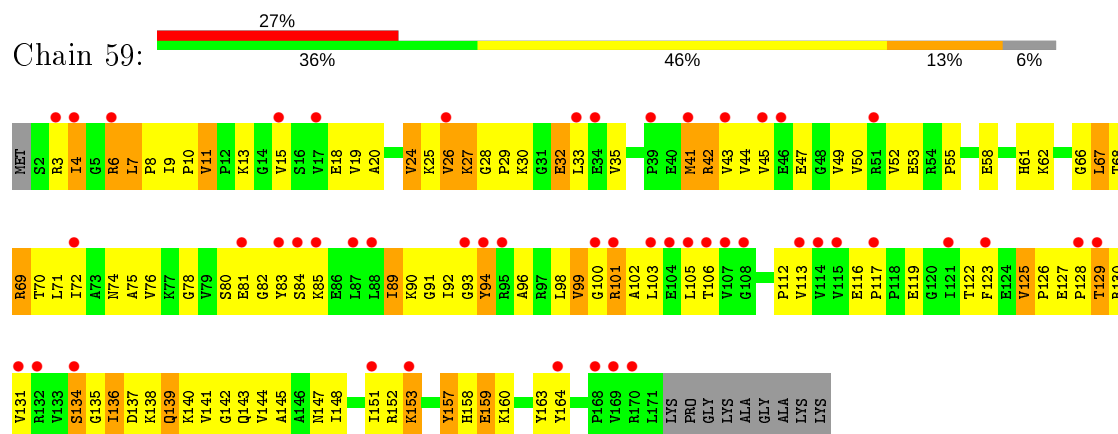
• Molecule 31: 50S ribosomal protein L5



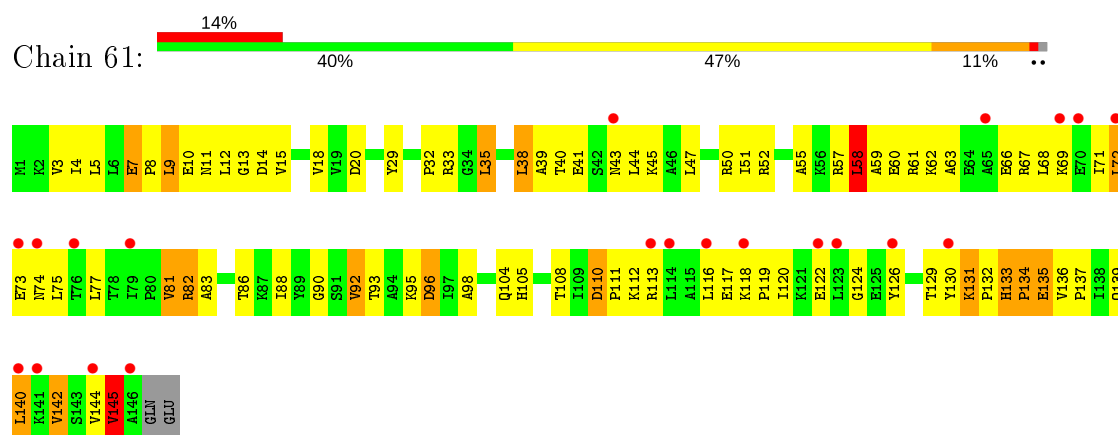
• Molecule 32: 50S ribosomal protein L6



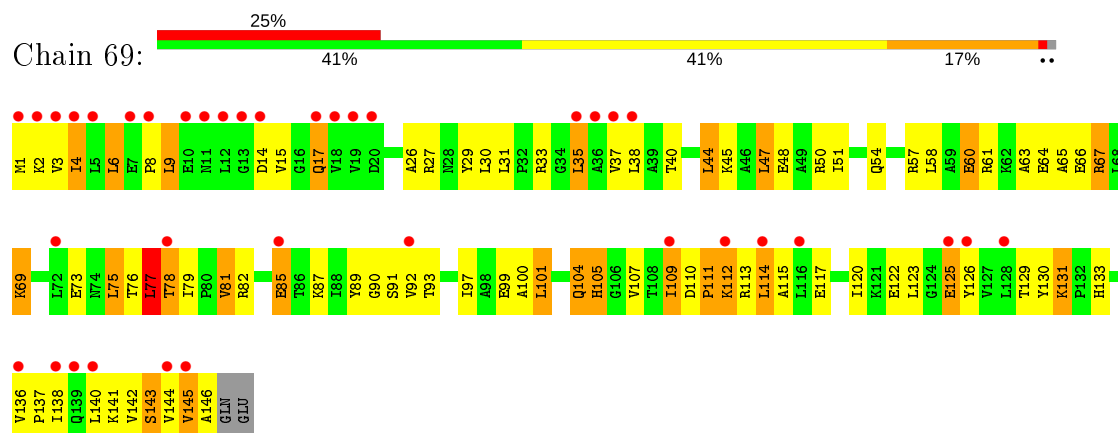
- Molecule 32: 50S ribosomal protein L6



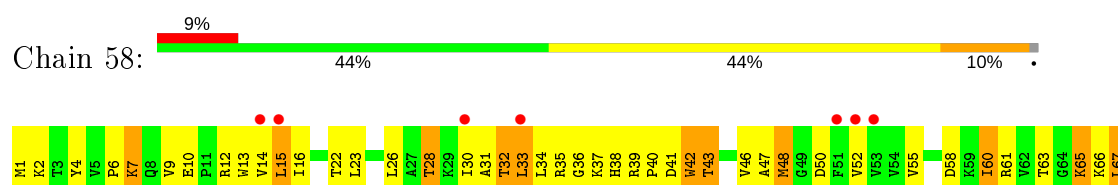
- Molecule 33: 50S ribosomal protein L9



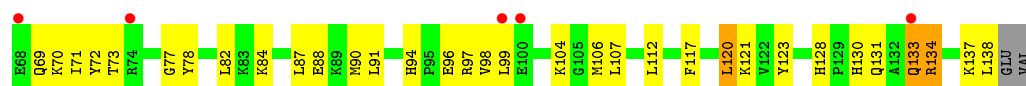
- Molecule 33: 50S ribosomal protein L9



- Molecule 34: 50S ribosomal protein L13







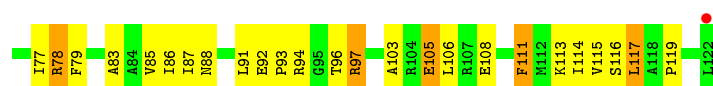
- Molecule 34: 50S ribosomal protein L13



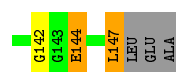
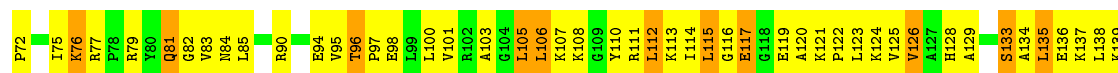
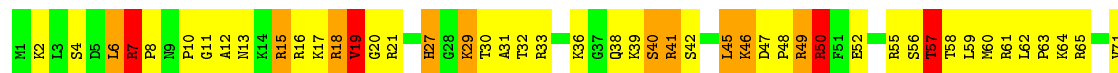
- Molecule 35: 50S ribosomal protein L14



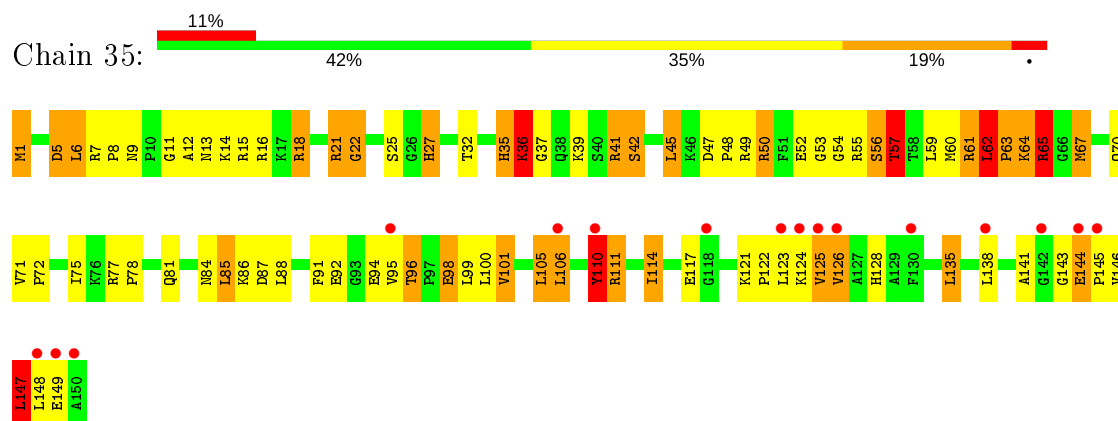
- Molecule 35: 50S ribosomal protein L14



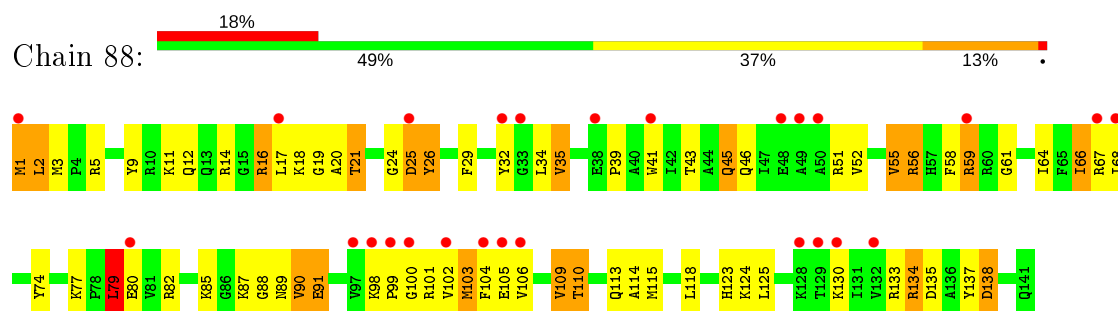
- Molecule 36: 50S ribosomal protein L15



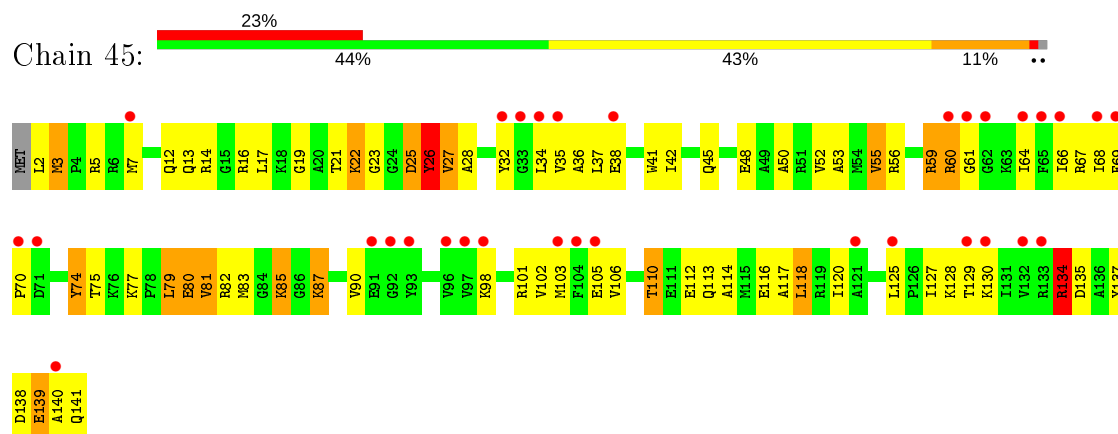
- Molecule 36: 50S ribosomal protein L15



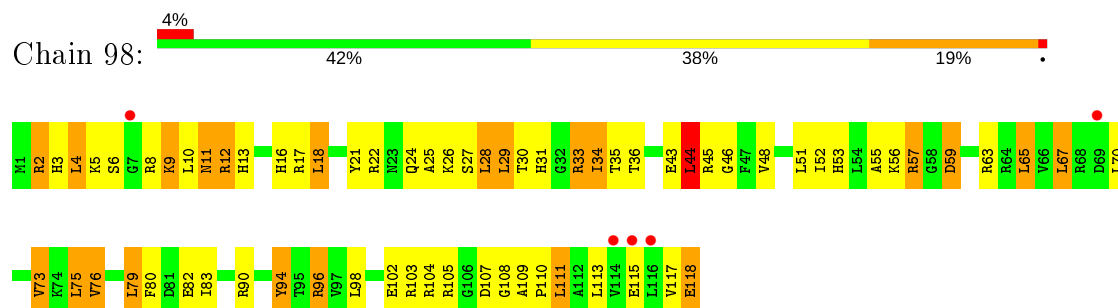
- Molecule 37: 50S ribosomal protein L16



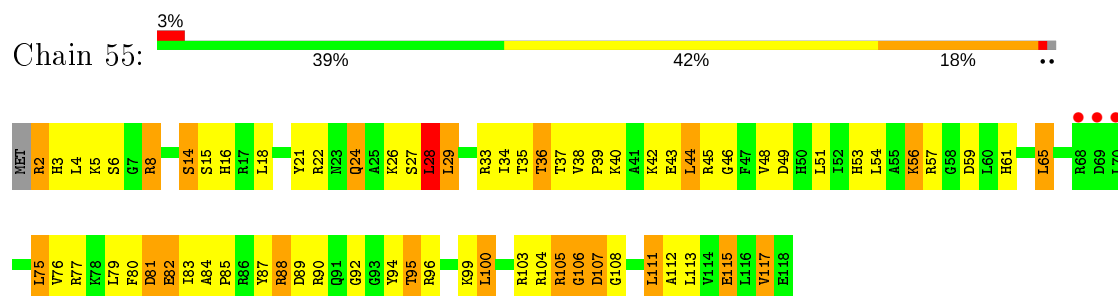
- Molecule 37: 50S ribosomal protein L16



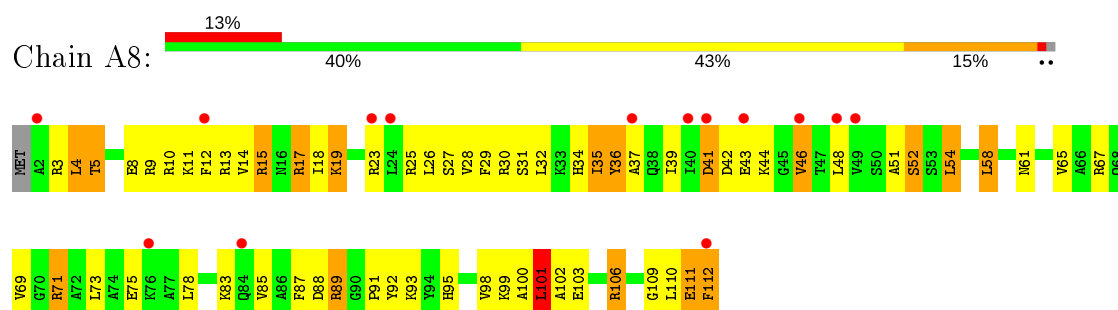
- Molecule 38: 50S ribosomal protein L17



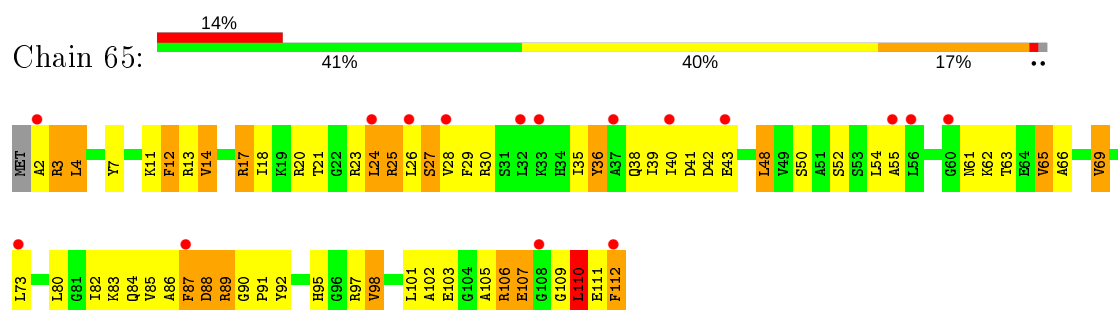
- Molecule 38: 50S ribosomal protein L17



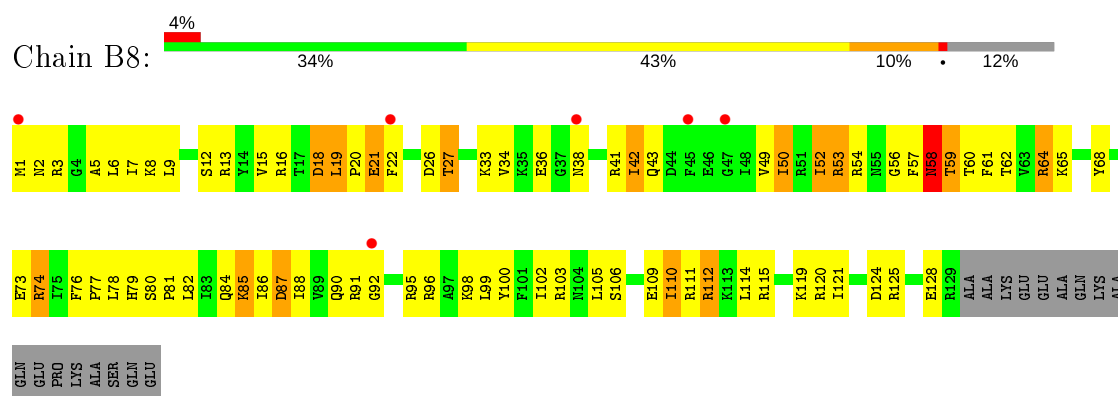
- Molecule 39: 50S ribosomal protein L18



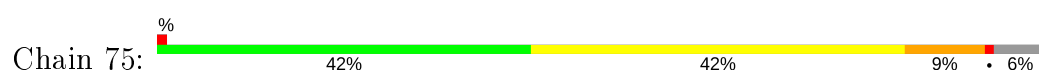
- Molecule 39: 50S ribosomal protein L18

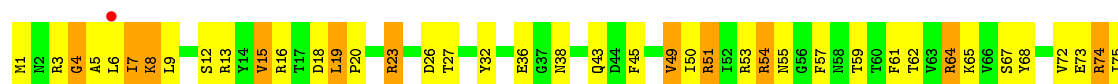


- Molecule 40: 50S ribosomal protein L19

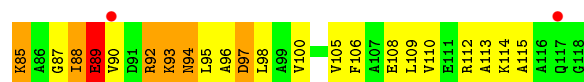


- Molecule 40: 50S ribosomal protein L19

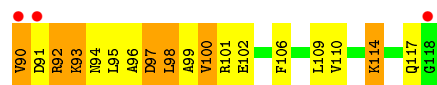
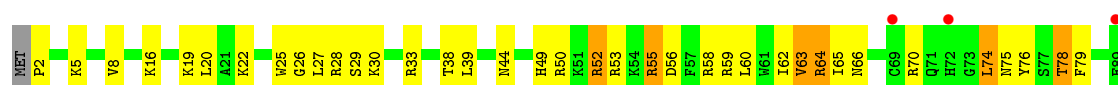




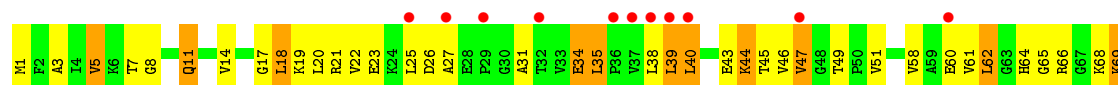
• Molecule 41: 50S ribosomal protein L20



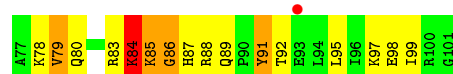
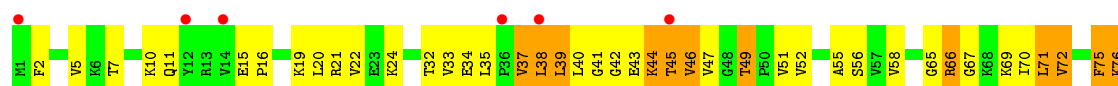
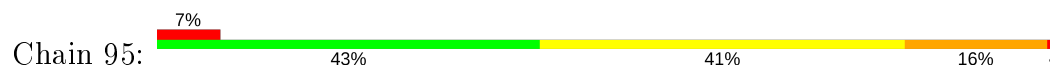
• Molecule 41: 50S ribosomal protein L20



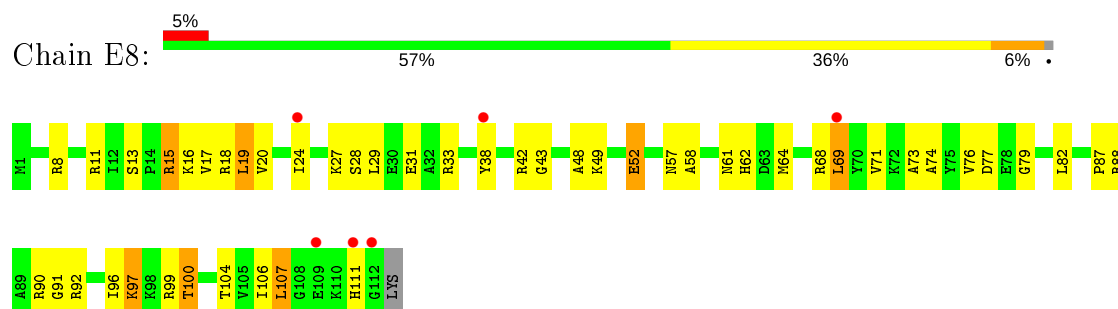
• Molecule 42: 50S ribosomal protein L21



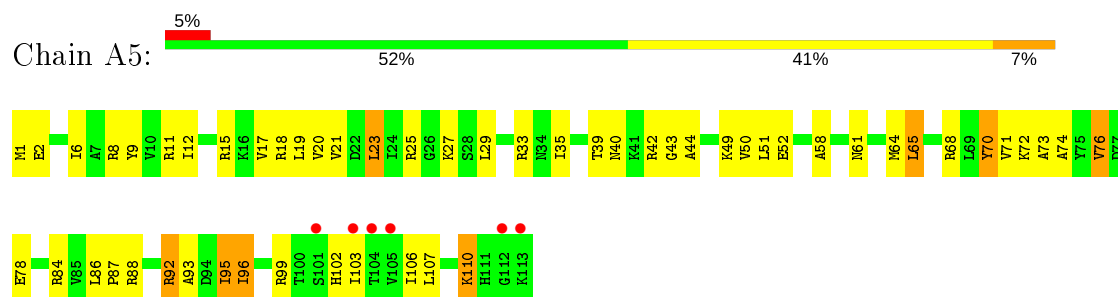
• Molecule 42: 50S ribosomal protein L21



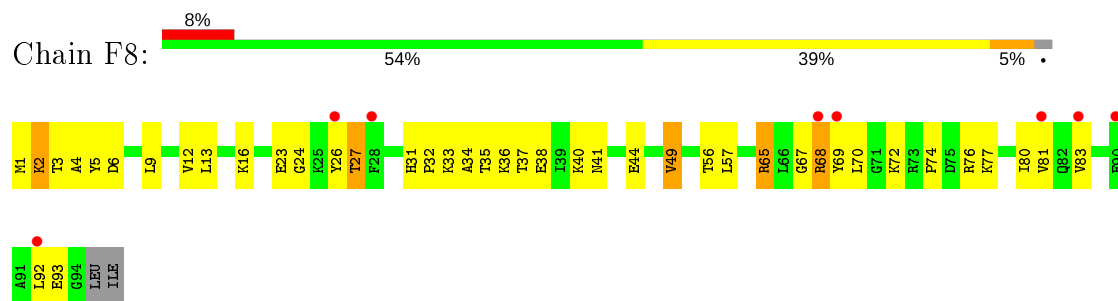
- Molecule 43: 50S ribosomal protein L22



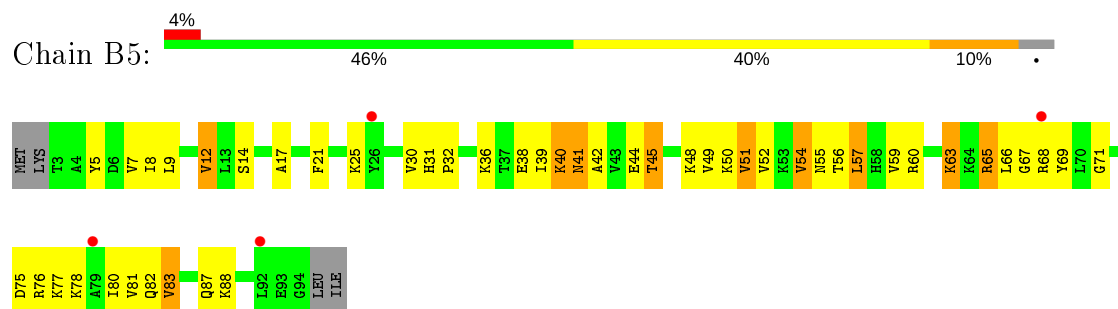
- Molecule 43: 50S ribosomal protein L22



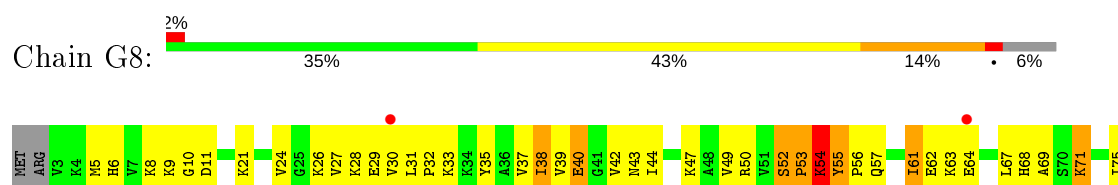
- Molecule 44: 50S ribosomal protein L23

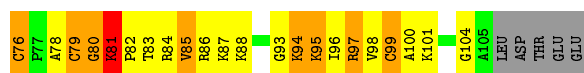


- Molecule 44: 50S ribosomal protein L23

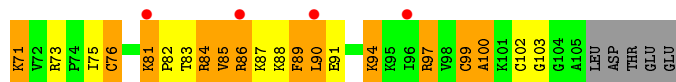
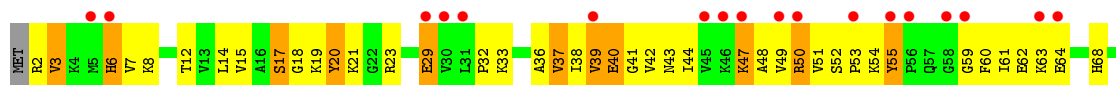


- Molecule 45: 50S ribosomal protein L24





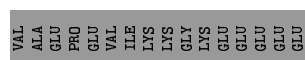
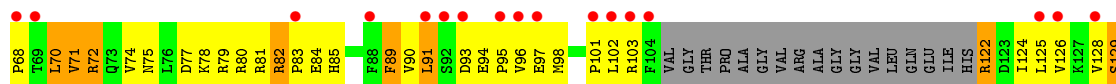
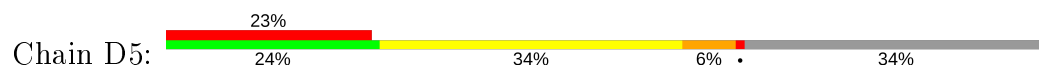
• Molecule 45: 50S ribosomal protein L24



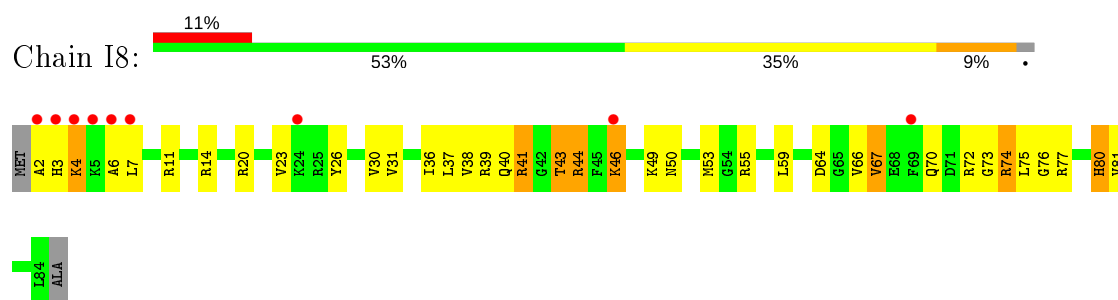
• Molecule 46: 50S ribosomal protein L25



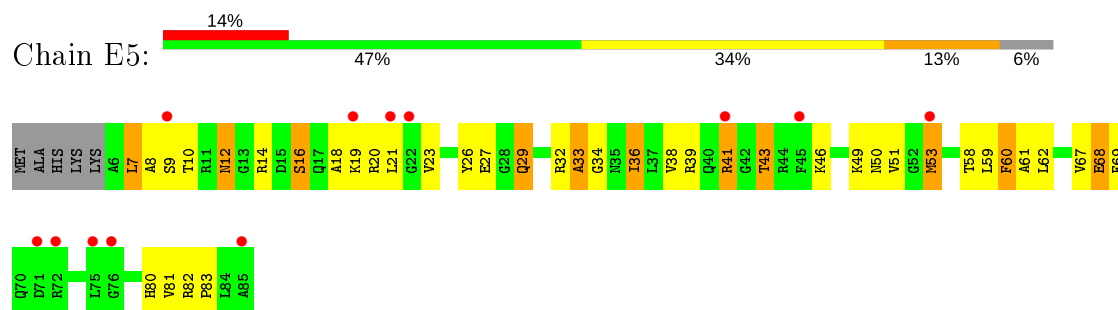
• Molecule 46: 50S ribosomal protein L25



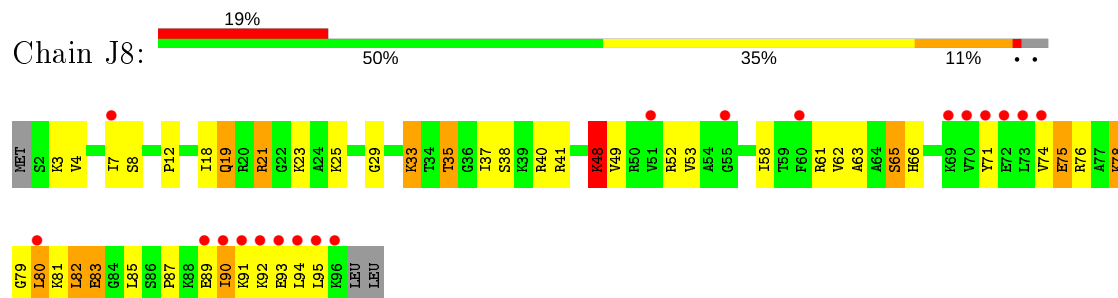
• Molecule 47: 50S ribosomal protein L27



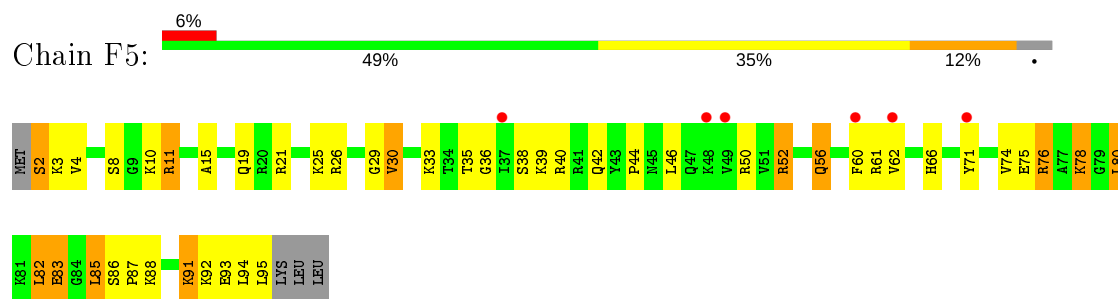
- Molecule 47: 50S ribosomal protein L27



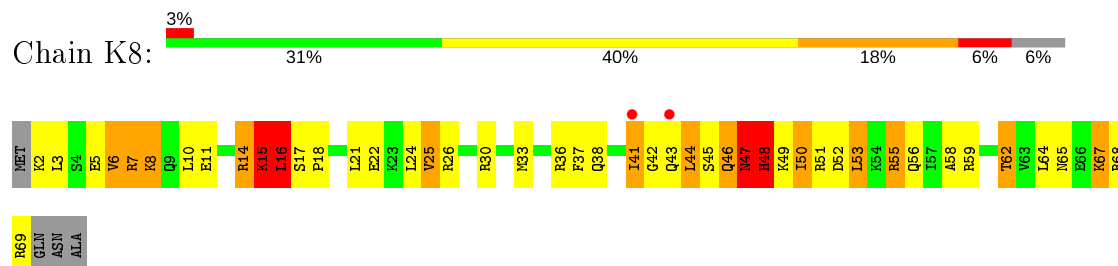
- Molecule 48: 50S ribosomal protein L28



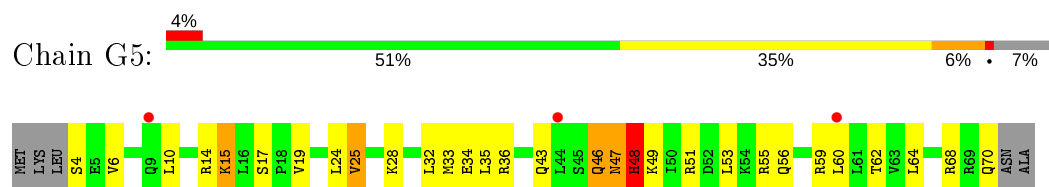
- Molecule 48: 50S ribosomal protein L28



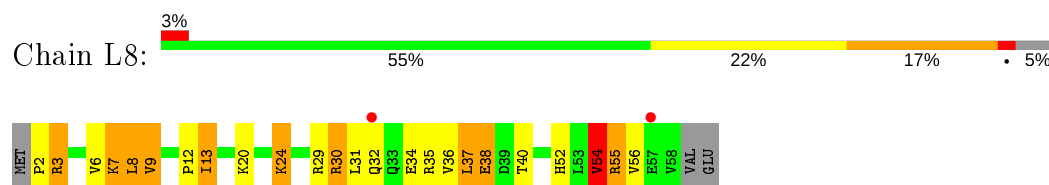
- Molecule 49: 50S ribosomal protein L29



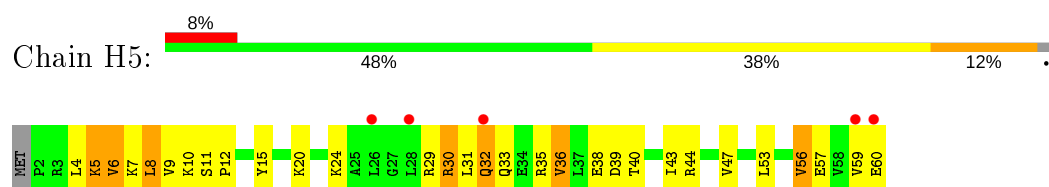
- Molecule 49: 50S ribosomal protein L29



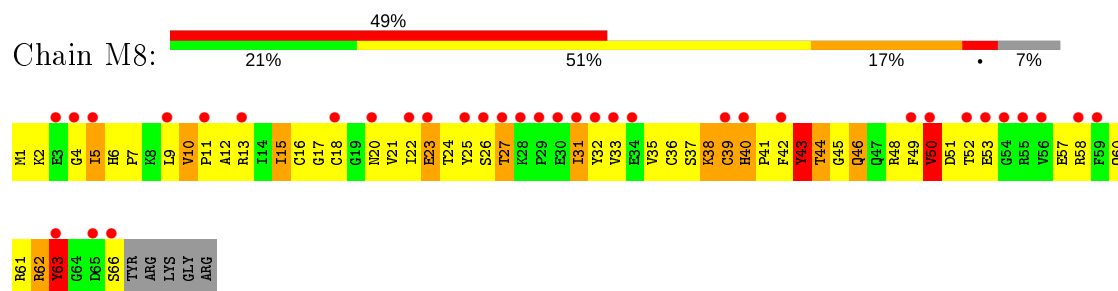
- Molecule 50: 50S ribosomal protein L30



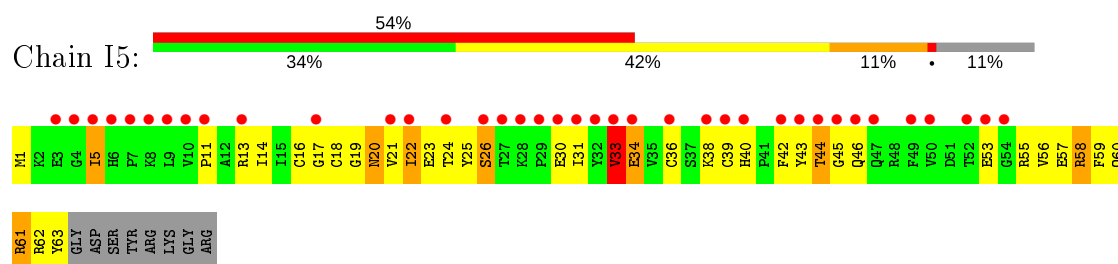
- Molecule 50: 50S ribosomal protein L30



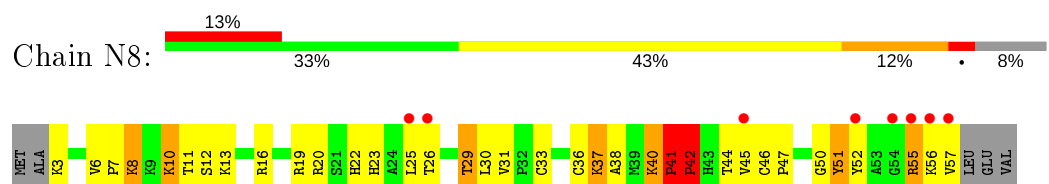
- Molecule 51: 50S ribosomal protein L31



- Molecule 51: 50S ribosomal protein L31

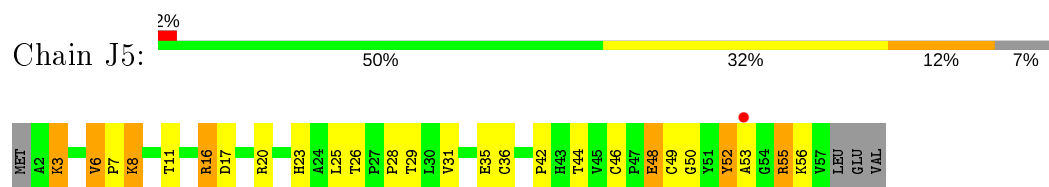


- Molecule 52: 50S ribosomal protein L32

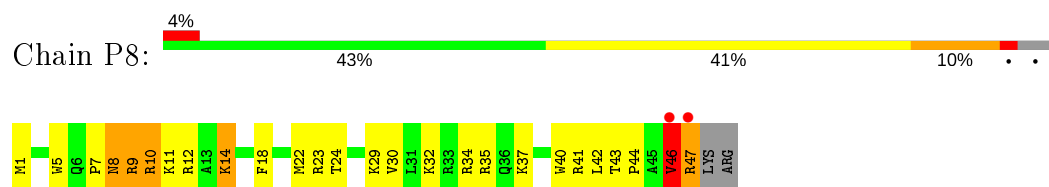




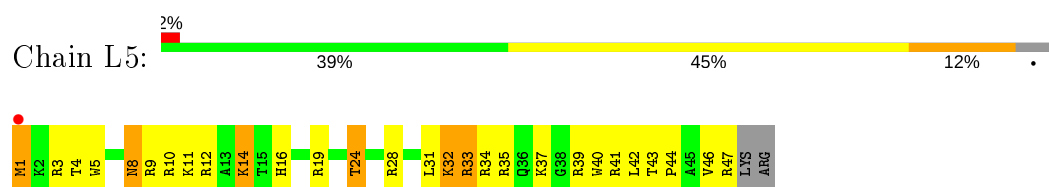
- Molecule 52: 50S ribosomal protein L32



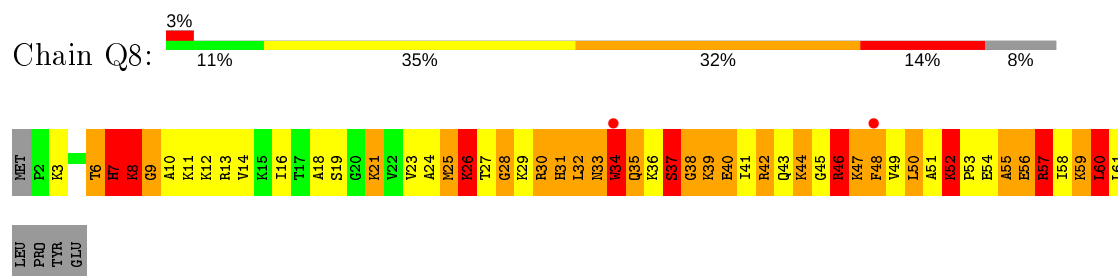
- Molecule 53: 50S ribosomal protein L34



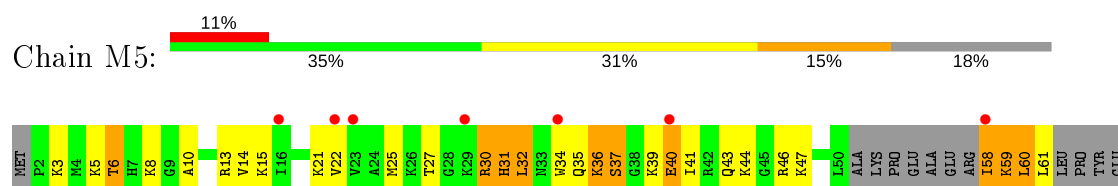
- Molecule 53: 50S ribosomal protein L34



- Molecule 54: 50S ribosomal protein L35



- Molecule 54: 50S ribosomal protein L35



## 4 Data and refinement statistics

Property	Value	Source
Space group	P 21 21 21	Depositor
Cell constants a, b, c, $\alpha$ , $\beta$ , $\gamma$	209.50Å 449.90Å 618.70Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	153.31 – 3.30 153.31 – 3.30	Depositor EDS
% Data completeness (in resolution range)	100.0 (153.31-3.30) 93.1 (153.31-3.30)	Depositor EDS
$R_{merge}$	0.54	Depositor
$R_{sym}$	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ <sup>1</sup>	1.47 (at 3.33Å)	Xtriage
Refinement program	PHENIX	Depositor
R, $R_{free}$	0.185 , (Not available) 0.185 , 0.231	Depositor DCC
$R_{free}$ test set	2000 reflections (0.23%)	wwPDB-VP
Wilson B-factor (Å <sup>2</sup> )	96.6	Xtriage
Anisotropy	0.284	Xtriage
Bulk solvent $k_{sol}$ (e/Å <sup>3</sup> ), $B_{sol}$ (Å <sup>2</sup> )	0.26 , 81.6	EDS
L-test for twinning <sup>2</sup>	$\langle  L  \rangle = 0.45$ , $\langle L^2 \rangle = 0.27$	Xtriage
Estimated twinning fraction	No twinning to report.	Xtriage
$F_o, F_c$ correlation	0.95	EDS
Total number of atoms	299429	wwPDB-VP
Average B, all atoms (Å <sup>2</sup> )	116.0	wwPDB-VP

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 1.55% of the height of the origin peak. No significant pseudotranslation is detected.*

<sup>1</sup>Intensities estimated from amplitudes.

<sup>2</sup>Theoretical 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: 5MU, MG, OMG, H2U, ZN, 1MG, PSU

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	13	0.70	4/36254 (0.0%)	1.35	317/56581 (0.6%)
1	1G	0.66	0/36390	1.30	250/56793 (0.4%)
2	12	0.39	0/1959	0.67	2/2642 (0.1%)
2	1E	0.41	0/1959	0.67	2/2642 (0.1%)
3	22	0.44	0/1636	0.65	0/2205
3	2E	0.50	0/1629	0.66	1/2195 (0.0%)
4	32	0.54	1/1732 (0.1%)	0.76	1/2318 (0.0%)
4	3E	0.63	4/1732 (0.2%)	0.75	2/2318 (0.1%)
5	42	0.50	0/1171	0.73	0/1576
5	4E	0.54	0/1171	0.70	0/1576
6	52	0.52	0/855	0.72	2/1154 (0.2%)
6	5E	0.56	0/855	0.67	0/1154
7	62	0.43	0/1218	0.60	0/1632
7	6E	0.40	0/1171	0.57	0/1567
8	72	0.47	0/1135	0.65	0/1527
8	7E	0.51	0/1135	0.72	0/1527
9	82	0.41	0/1002	0.66	0/1346
9	8E	0.43	0/1028	0.65	0/1379
10	1A	0.40	0/814	0.63	0/1095
10	1I	0.47	0/814	0.70	0/1095
11	2A	0.48	0/879	0.70	0/1187
11	2I	0.47	0/879	0.72	1/1187 (0.1%)
12	3A	0.58	0/991	0.77	0/1327
12	3I	0.62	0/972	0.83	0/1301
13	4A	0.34	0/938	0.59	0/1258
13	4I	0.45	0/938	0.65	0/1258
14	5A	0.45	0/484	0.72	0/643
14	5I	0.64	1/500 (0.2%)	0.80	0/664
15	6A	0.51	0/744	0.64	0/992
15	6I	0.51	0/744	0.71	0/992
16	7A	0.52	0/721	0.70	0/970
16	7I	0.47	0/721	0.67	0/970

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z  >5	RMSZ	# Z  >5
17	8A	0.56	0/836	0.66	0/1117
17	8I	0.52	0/847	0.73	0/1131
18	9A	0.53	0/569	0.78	0/757
18	9I	0.49	0/555	0.74	0/739
19	AA	0.39	0/638	0.66	0/860
19	AI	0.46	0/657	0.75	1/885 (0.1%)
20	BA	0.48	0/764	0.73	0/1007
20	BI	0.41	0/764	0.66	0/1007
21	1B	0.47	0/221	0.63	0/288
21	1F	0.41	0/203	0.65	0/266
22	1K	0.52	0/1988	1.13	11/3099 (0.4%)
22	1L	0.45	0/1988	1.06	9/3099 (0.3%)
23	2K	0.69	0/1746	1.39	17/2719 (0.6%)
23	2L	0.69	0/1768	1.47	21/2753 (0.8%)
24	3K	0.49	0/1838	1.12	6/2862 (0.2%)
24	3L	0.54	0/1786	1.13	4/2778 (0.1%)
25	4K	0.70	0/372	1.05	0/574
25	4L	0.60	0/485	1.03	1/749 (0.1%)
26	14	0.84	45/70167 (0.1%)	1.51	1106/109541 (1.0%)
26	1H	0.91	66/70233 (0.1%)	1.60	1430/109643 (1.3%)
27	16	0.75	0/2928	1.48	45/4568 (1.0%)
27	1J	0.71	0/2928	1.37	24/4568 (0.5%)
28	11	0.70	1/2176 (0.0%)	0.89	2/2933 (0.1%)
28	19	0.69	2/2170 (0.1%)	0.91	2/2926 (0.1%)
29	21	0.63	0/1601	0.91	2/2160 (0.1%)
29	29	0.67	0/1601	0.97	2/2160 (0.1%)
30	31	0.67	0/1620	0.82	0/2194
30	39	0.62	0/1662	0.90	0/2249
31	41	0.43	0/1498	0.66	1/2016 (0.0%)
31	49	0.37	0/1498	0.63	0/2016
32	51	0.57	0/1362	0.85	0/1841
32	59	0.38	0/1332	0.75	2/1802 (0.1%)
33	61	0.52	0/1151	0.78	2/1558 (0.1%)
33	69	0.48	0/1151	0.75	2/1558 (0.1%)
34	15	0.54	0/1131	0.71	0/1525
34	58	0.56	0/1131	0.79	1/1525 (0.1%)
35	25	0.64	0/942	0.81	2/1269 (0.2%)
35	68	0.61	0/942	0.76	0/1269
36	35	0.59	0/1161	0.99	1/1544 (0.1%)
36	78	0.64	0/1139	1.03	2/1514 (0.1%)
37	45	0.63	0/1134	0.82	0/1517
37	88	0.66	0/1142	0.92	3/1527 (0.2%)
38	55	0.62	0/973	0.87	1/1302 (0.1%)

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z  >5	RMSZ	# Z  >5
38	98	0.53	0/981	0.85	2/1312 (0.2%)
39	65	0.55	0/891	0.91	1/1187 (0.1%)
39	A8	0.55	0/891	0.81	1/1187 (0.1%)
40	75	0.61	0/1155	0.77	0/1542
40	B8	0.64	0/1095	0.79	0/1463
41	85	0.59	0/981	0.79	0/1306
41	C8	0.67	0/981	0.88	2/1306 (0.2%)
42	95	0.66	0/789	0.85	2/1057 (0.2%)
42	D8	0.59	0/789	0.83	1/1057 (0.1%)
43	A5	0.68	0/910	0.87	0/1220
43	E8	0.64	0/901	0.83	0/1209
44	B5	0.69	0/739	0.81	0/993
44	F8	0.72	0/756	0.86	0/1014
45	C5	0.68	0/807	0.90	1/1076 (0.1%)
45	G8	0.71	0/796	0.98	5/1062 (0.5%)
46	D5	0.47	0/1145	0.67	0/1547
46	H8	0.47	0/1403	0.76	2/1901 (0.1%)
47	E5	0.63	0/635	0.89	0/848
47	I8	0.69	0/665	0.82	0/885
48	F5	0.65	0/744	0.87	1/989 (0.1%)
48	J8	0.67	0/753	0.93	3/1000 (0.3%)
49	G5	0.61	0/569	0.85	0/753
49	K8	0.68	0/577	0.92	1/763 (0.1%)
50	H5	0.48	0/473	0.63	0/635
50	L8	0.66	0/457	0.79	0/613
51	I5	0.44	0/527	0.79	0/709
51	M8	0.43	0/545	0.77	0/733
52	J5	0.60	0/448	0.82	0/606
52	N8	0.64	0/443	0.84	0/599
53	L5	0.65	0/417	0.83	0/550
53	P8	0.75	0/417	0.88	0/550
54	M5	0.71	0/426	0.97	1/556 (0.2%)
54	Q8	1.05	0/486	1.47	6/638 (0.9%)
All	All	0.74	124/322561 (0.0%)	1.33	3307/483352 (0.7%)

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
2	12	0	1

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Mol	Chain	#Chirality outliers	#Planarity outliers
2	1E	0	3
4	32	0	3
4	3E	0	1
9	82	0	1
9	8E	0	1
10	1A	0	1
11	2A	0	1
12	3A	0	2
12	3I	0	1
13	4A	0	1
14	5A	0	1
19	AI	0	2
20	BA	0	2
28	11	0	1
28	19	0	3
29	21	0	4
29	29	0	8
30	31	0	1
30	39	0	6
31	49	0	1
33	61	0	3
33	69	0	3
34	15	0	1
36	35	0	7
36	78	0	2
37	45	0	5
37	88	0	3
38	55	0	2
38	98	0	1
40	75	0	2
40	B8	0	2
41	85	0	2
41	C8	0	1
42	95	0	1
43	A5	0	1
45	C5	0	4
45	G8	0	4
46	H8	0	3
47	E5	0	2
47	I8	0	2
48	F5	0	1
48	J8	0	1

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Mol	Chain	#Chirality outliers	#Planarity outliers
49	G5	0	3
49	K8	0	5
51	I5	0	1
51	M8	0	3
52	J5	0	1
52	N8	0	2
53	P8	0	1
54	M5	0	2
54	Q8	0	12
All	All	0	128

All (124) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
26	1H	774	A	N9-C4	-14.53	1.29	1.37
26	14	783	A	N9-C4	-9.44	1.32	1.37
26	1H	71	A	N9-C4	-9.18	1.32	1.37
26	14	2518	A	N9-C4	-7.93	1.33	1.37
4	3E	12	CYS	CB-SG	7.85	1.95	1.82
26	14	1950	G	C2-N3	7.73	1.39	1.32
26	1H	783	A	N7-C5	-7.61	1.34	1.39
26	1H	783	A	C5-C6	-7.49	1.34	1.41
26	1H	783	A	N9-C4	-7.41	1.33	1.37
26	14	71	A	N9-C4	-7.37	1.33	1.37
26	14	74	A	N9-C4	-7.36	1.33	1.37
26	14	2346	A	N3-C4	-7.36	1.30	1.34
26	1H	676	A	N9-C4	-7.33	1.33	1.37
26	1H	127	A	C5-C6	-7.30	1.34	1.41
26	14	774	A	N9-C4	-7.27	1.33	1.37
26	1H	784	A	C6-N1	-7.26	1.30	1.35
26	14	2506	U	C2-N3	7.24	1.42	1.37
26	1H	1950	G	C2-N3	7.13	1.38	1.32
26	1H	528	A	N9-C4	-7.04	1.33	1.37
26	1H	1786	A	N9-C4	-7.04	1.33	1.37
26	14	676	A	N9-C4	-7.00	1.33	1.37
26	1H	1899	G	N9-C4	-6.95	1.32	1.38
26	1H	821	A	N7-C5	-6.95	1.35	1.39
26	14	1332	G	N9-C4	-6.87	1.32	1.38
1	13	1523	G	C6-O6	6.82	1.30	1.24
1	13	792	A	C5-C6	-6.77	1.34	1.41
26	1H	1210	A	N9-C4	-6.65	1.33	1.37
26	14	945	A	N9-C4	-6.62	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
26	14	2287	A	N9-C4	-6.62	1.33	1.37
26	1H	2346	A	N3-C4	-6.58	1.30	1.34
14	5I	27	CYS	CB-SG	-6.58	1.71	1.82
26	1H	774	A	N7-C5	-6.47	1.35	1.39
26	1H	330	A	N9-C4	-6.46	1.33	1.37
26	14	447	A	N9-C4	-6.45	1.33	1.37
26	1H	676	A	C5-C4	6.43	1.43	1.38
4	3E	26	CYS	CB-SG	6.43	1.93	1.82
26	1H	74	A	N9-C4	-6.41	1.34	1.37
26	1H	57	C	N3-C4	-6.37	1.29	1.33
4	32	31	CYS	CB-SG	-6.36	1.71	1.82
26	1H	1378	A	N9-C4	-6.28	1.34	1.37
26	1H	945	A	N7-C5	-6.20	1.35	1.39
26	1H	774	A	N3-C4	-6.14	1.31	1.34
26	1H	2518	A	N9-C4	-6.10	1.34	1.37
4	3E	31	CYS	CB-SG	6.09	1.92	1.82
28	19	28	GLU	CG-CD	6.08	1.61	1.51
26	14	528	A	N9-C4	-6.08	1.34	1.37
26	14	945	A	C5-C6	-6.01	1.35	1.41
26	1H	676	A	N9-C8	6.00	1.42	1.37
26	14	828	U	N3-C4	-5.95	1.33	1.38
26	14	1890	A	N9-C4	-5.88	1.34	1.37
28	19	28	GLU	CB-CG	5.81	1.63	1.52
26	14	1616	A	N9-C4	-5.81	1.34	1.37
26	1H	1899	G	N9-C8	5.80	1.42	1.37
26	14	2062	A	N3-C4	5.80	1.38	1.34
26	1H	2392	A	C5-C4	5.80	1.42	1.38
26	14	774	A	C5-C6	-5.78	1.35	1.41
26	14	783	A	N7-C5	-5.72	1.35	1.39
26	1H	1977	A	C6-N1	-5.70	1.31	1.35
26	14	512	G	N9-C4	-5.69	1.33	1.38
26	1H	472	A	N3-C4	-5.69	1.31	1.34
26	1H	2287	A	N9-C4	-5.67	1.34	1.37
26	14	528	A	N3-C4	-5.64	1.31	1.34
26	14	676	A	N9-C8	5.62	1.42	1.37
26	1H	1786	A	C5-C6	-5.61	1.35	1.41
26	1H	74	A	N3-C4	-5.61	1.31	1.34
26	14	2278	A	N3-C4	-5.60	1.31	1.34
26	1H	2062	A	N3-C4	5.60	1.38	1.34
26	14	332	A	N9-C4	-5.60	1.34	1.37
26	1H	2430	A	N9-C4	-5.59	1.34	1.37
26	1H	138	G	N9-C8	5.57	1.41	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
26	1H	1786	A	N7-C5	-5.56	1.35	1.39
26	14	676	A	C5-C4	5.56	1.42	1.38
26	1H	1698	A	N3-C4	-5.55	1.31	1.34
26	14	1610	A	C5-C6	-5.53	1.36	1.41
26	1H	783	A	N3-C4	-5.51	1.31	1.34
26	1H	140	A	C5-C6	-5.51	1.36	1.41
26	1H	71	A	C5-C6	-5.50	1.36	1.41
26	14	1332	G	N3-C4	-5.50	1.31	1.35
26	14	2062	A	C6-N1	5.50	1.39	1.35
26	14	2873	A	N7-C5	-5.49	1.35	1.39
26	14	330	A	N9-C4	-5.48	1.34	1.37
1	13	974	A	N9-C4	-5.47	1.34	1.37
26	14	1678	G	N9-C4	-5.47	1.33	1.38
26	14	786	C	N3-C4	-5.47	1.30	1.33
26	1H	1021	A	N9-C4	-5.47	1.34	1.37
26	14	1617	C	N1-C6	-5.46	1.33	1.37
26	1H	667	U	N1-C2	-5.41	1.33	1.38
26	14	621	A	N9-C4	-5.41	1.34	1.37
26	1H	2602	A	N3-C4	5.37	1.38	1.34
26	1H	2713	A	C5-C4	5.37	1.42	1.38
26	1H	1829	A	C5-C4	-5.34	1.35	1.38
26	14	2490	G	N9-C8	5.34	1.41	1.37
26	14	1613	G	C6-N1	-5.33	1.35	1.39
26	1H	1363	C	N3-C4	-5.30	1.30	1.33
26	1H	449	A	N3-C4	-5.30	1.31	1.34
26	1H	1616	A	N7-C5	-5.29	1.36	1.39
28	11	28	GLU	CG-CD	5.29	1.59	1.51
26	1H	1683	C	N3-C4	-5.26	1.30	1.33
26	1H	82	G	C6-O6	5.25	1.28	1.24
4	3E	9	CYS	CB-SG	5.25	1.91	1.82
26	14	1950	G	C5-C4	5.23	1.42	1.38
26	1H	2346	A	N9-C4	-5.21	1.34	1.37
26	1H	1496	A	C5-C6	-5.20	1.36	1.41
26	1H	1558	A	N3-C4	-5.18	1.31	1.34
26	1H	768	G	C6-N1	-5.18	1.35	1.39
26	14	1142(A)	A	N9-C4	-5.18	1.34	1.37
26	14	777	A	N9-C4	-5.17	1.34	1.37
26	1H	859	G	N9-C4	-5.17	1.33	1.38
26	14	821	A	N7-C5	-5.17	1.36	1.39
1	13	792	A	N9-C4	-5.14	1.34	1.37
26	1H	1614	A	N9-C4	-5.12	1.34	1.37
26	14	1187	G	N3-C4	-5.12	1.31	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
26	1H	1332	G	N9-C4	-5.09	1.33	1.38
26	1H	1950	G	C5-C4	5.08	1.42	1.38
26	14	751	A	N9-C4	-5.08	1.34	1.37
26	1H	692	C	N3-C4	-5.08	1.30	1.33
26	1H	793	A	C5-C6	-5.06	1.36	1.41
26	1H	1300	U	N1-C2	-5.05	1.34	1.38
26	1H	1626	G	N3-C4	-5.05	1.31	1.35
26	1H	2032	G	N9-C8	-5.05	1.34	1.37
26	1H	2600	A	N3-C4	-5.05	1.31	1.34
26	14	783	A	C5-C6	-5.04	1.36	1.41
26	1H	330	A	N3-C4	-5.01	1.31	1.34
26	1H	1257	C	N3-C4	-5.01	1.30	1.33

All (3307) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	1899	G	N3-C4-N9	-18.45	114.93	126.00
26	14	1332	G	N3-C4-N9	-15.36	116.78	126.00
26	1H	676	A	C2-N3-C4	-14.95	103.12	110.60
26	1H	783	A	C5-N7-C8	-14.89	96.45	103.90
26	14	1786	A	N7-C8-N9	14.75	121.17	113.80
26	1H	1899	G	N3-C4-C5	14.68	135.94	128.60
26	1H	774	A	C2-N3-C4	-14.26	103.47	110.60
26	1H	2430	A	C2-N3-C4	-14.07	103.57	110.60
26	14	828	U	C5-C4-O4	13.91	134.25	125.90
26	1H	71	A	C2-N3-C4	-13.86	103.67	110.60
26	1H	127	A	N1-C6-N6	13.76	126.86	118.60
26	14	676	A	N1-C6-N6	13.73	126.84	118.60
26	14	676	A	C2-N3-C4	-13.71	103.74	110.60
26	1H	1332	G	C2-N3-C4	-13.70	105.05	111.90
26	1H	1786	A	C5-N7-C8	-13.70	97.05	103.90
26	1H	783	A	N1-C6-N6	13.68	126.81	118.60
26	1H	945	A	C6-C5-N7	-13.66	122.74	132.30
26	1H	1899	G	C2-N3-C4	-13.47	105.17	111.90
26	14	676	A	C5-N7-C8	-13.47	97.17	103.90
26	1H	676	A	C5-N7-C8	-13.32	97.24	103.90
26	14	741	G	O5'-P-OP1	-13.25	93.77	105.70
1	13	792	A	N1-C6-N6	13.20	126.52	118.60
26	14	774	A	C2-N3-C4	-13.11	104.05	110.60
26	1H	1786	A	N7-C8-N9	13.09	120.34	113.80
26	1H	783	A	C2-N3-C4	-12.91	104.15	110.60
26	14	74	A	C2-N3-C4	-12.90	104.15	110.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	1786	A	C8-N9-C4	-12.77	100.69	105.80
26	1H	1899	G	N3-C2-N2	-12.76	110.97	119.90
26	14	2430	A	N1-C6-N6	12.63	126.18	118.60
26	14	2518	A	C2-N3-C4	-12.46	104.37	110.60
26	1H	774	A	N3-C4-N9	-12.36	117.51	127.40
26	1H	783	A	N7-C8-N9	12.34	119.97	113.80
26	14	945	A	N1-C6-N6	12.32	125.99	118.60
26	1H	783	A	C6-C5-N7	-12.32	123.68	132.30
1	13	690	G	C6-C5-N7	-12.17	123.10	130.40
26	14	945	A	C2-N3-C4	-12.11	104.54	110.60
26	1H	2430	A	N1-C6-N6	12.09	125.85	118.60
26	1H	783	A	C4-C5-N7	12.08	116.74	110.70
26	1H	1972	A	N1-C6-N6	12.05	125.83	118.60
26	1H	774	A	N3-C4-C5	11.98	135.19	126.80
26	14	1332	G	N3-C4-C5	11.97	134.58	128.60
26	1H	945	A	N1-C6-N6	11.92	125.75	118.60
26	14	1786	A	C5-N7-C8	-11.89	97.95	103.90
26	1H	945	A	C4-C5-C6	11.80	122.90	117.00
26	1H	1210	A	C5-N7-C8	-11.80	98.00	103.90
26	14	676	A	C4-C5-N7	11.68	116.54	110.70
26	14	1678	G	C2-N3-C4	-11.68	106.06	111.90
26	1H	1786	A	C2-N3-C4	-11.68	104.76	110.60
1	13	792	A	C4-C5-N7	11.62	116.51	110.70
26	1H	2584	U	N3-C2-O2	-11.62	114.07	122.20
26	1H	74	A	C2-N3-C4	-11.49	104.86	110.60
27	16	81	G	C4-C5-N7	11.25	115.30	110.80
26	14	1332	G	C2-N3-C4	-11.17	106.31	111.90
26	1H	140	A	C5-N7-C8	-11.15	98.32	103.90
26	1H	676	A	N7-C8-N9	11.14	119.37	113.80
26	14	1786	A	C2-N3-C4	-11.13	105.03	110.60
26	14	2873	A	C5-N7-C8	-11.11	98.34	103.90
26	14	2287	A	C2-N3-C4	-11.10	105.05	110.60
26	1H	2509	G	C5-C6-O6	-11.00	122.00	128.60
26	1H	828	U	C5-C4-O4	10.89	132.43	125.90
26	14	330	A	C2-N3-C4	-10.83	105.19	110.60
26	1H	121	G	C5-C6-O6	-10.81	122.11	128.60
26	1H	1021	A	C2-N3-C4	-10.79	105.20	110.60
26	14	528	A	C2-N3-C4	-10.77	105.22	110.60
26	1H	71	A	C5-N7-C8	-10.76	98.52	103.90
26	14	2873	A	N7-C8-N9	10.75	119.17	113.80
26	1H	2430	A	C5-C6-N1	-10.75	112.33	117.70
26	14	783	A	C2-N3-C4	-10.75	105.23	110.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	1899	G	C2-N3-C4	-10.72	106.54	111.90
26	1H	1210	A	N1-C6-N6	10.70	125.02	118.60
26	1H	945	A	N7-C8-N9	10.67	119.13	113.80
26	14	1698	A	N1-C6-N6	10.65	124.99	118.60
26	1H	2609	U	C5-C6-N1	-10.63	117.39	122.70
26	14	783	A	C5-N7-C8	-10.62	98.59	103.90
26	1H	1496	A	N1-C6-N6	10.53	124.92	118.60
26	14	528	A	N1-C2-N3	10.47	134.54	129.30
26	14	2506	U	C5-C6-N1	10.46	127.93	122.70
26	14	71	A	C2-N3-C4	-10.41	105.39	110.60
26	14	71	A	N1-C6-N6	10.37	124.82	118.60
26	14	1816	G	O5'-P-OP1	-10.35	96.38	105.70
26	14	74	A	N1-C6-N6	10.34	124.80	118.60
26	14	945	A	C4-C5-N7	10.33	115.86	110.70
26	1H	2346	A	O4'-C1'-N9	10.29	116.43	108.20
26	14	945	A	C5-N7-C8	-10.29	98.76	103.90
26	14	676	A	N3-C4-C5	10.27	133.99	126.80
26	1H	1950	G	C8-N9-C4	-10.26	102.30	106.40
26	1H	1249	U	O5'-P-OP1	-10.22	96.50	105.70
26	14	2873	A	C2-N3-C4	-10.22	105.49	110.60
26	14	2490	G	C8-N9-C4	-10.20	102.32	106.40
26	1H	1332	G	C5-N7-C8	-10.16	99.22	104.30
26	1H	330	A	C2-N3-C4	-10.15	105.53	110.60
26	1H	186	G	C5-C6-O6	-10.15	122.51	128.60
26	1H	2331	G	C8-N9-C4	10.13	110.45	106.40
26	1H	684	G	C8-N9-C4	-10.13	102.35	106.40
26	1H	528	A	C2-N3-C4	-10.09	105.56	110.60
26	1H	1786	A	C8-N9-C4	-10.08	101.77	105.80
26	14	391	G	N1-C6-O6	10.08	125.95	119.90
26	1H	1204	A	O4'-C1'-N9	10.02	116.22	108.20
26	1H	71	A	N3-C4-C5	9.99	133.79	126.80
26	14	1021	A	C2-N3-C4	-9.99	105.61	110.60
26	14	774	A	N3-C4-C5	9.97	133.78	126.80
26	1H	504	U	C2-N1-C1'	9.97	129.66	117.70
26	1H	140	A	C4-C5-N7	9.96	115.68	110.70
26	1H	2503	A	N1-C6-N6	9.95	124.57	118.60
26	14	2584	U	N3-C2-O2	-9.93	115.25	122.20
26	14	767	U	C5-C4-O4	9.92	131.85	125.90
26	1H	1210	A	C4-C5-N7	9.92	115.66	110.70
26	1H	1602	U	C5-C6-N1	-9.90	117.75	122.70
26	1H	796	C	O5'-P-OP2	-9.89	96.80	105.70
26	14	2307	G	O4'-C1'-N9	9.88	116.11	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	1276	A	N1-C6-N6	9.88	124.53	118.60
26	14	2430	A	C2-N3-C4	-9.88	105.66	110.60
26	1H	783	A	C8-N9-C4	-9.86	101.86	105.80
26	1H	140	A	N1-C6-N6	9.86	124.51	118.60
26	1H	672	C	O5'-P-OP2	-9.84	96.85	105.70
26	14	74	A	C5-C6-N1	-9.80	112.80	117.70
1	13	690	G	C4-N9-C1'	9.78	139.22	126.50
26	1H	130	C	C6-N1-C2	9.78	124.21	120.30
26	14	330	A	N1-C6-N6	9.79	124.47	118.60
26	1H	2468	G	O4'-C1'-N9	9.78	116.03	108.20
26	1H	2392	A	N7-C8-N9	9.77	118.68	113.80
26	1H	2856	C	C6-N1-C2	-9.76	116.39	120.30
26	1H	2679	A	O5'-P-OP2	-9.74	96.93	105.70
26	14	2873	A	N1-C6-N6	9.73	124.44	118.60
36	78	50	ARG	NE-CZ-NH2	9.71	125.16	120.30
26	14	1950	G	N7-C8-N9	9.69	117.95	113.10
26	14	828	U	N3-C4-O4	-9.68	112.62	119.40
26	14	828	U	N3-C2-O2	-9.68	115.42	122.20
1	13	690	G	O4'-C1'-N9	9.68	115.94	108.20
26	1H	735	A	C8-N9-C4	9.67	109.67	105.80
26	1H	917	A	C2-N3-C4	-9.66	105.77	110.60
1	13	792	A	N9-C4-C5	-9.63	101.95	105.80
26	1H	121	G	C6-C5-N7	-9.61	124.64	130.40
26	1H	121	G	N3-C4-N9	9.61	131.77	126.00
26	1H	2331	G	N1-C6-O6	9.61	125.66	119.90
26	1H	2403	C	C6-N1-C2	-9.61	116.46	120.30
26	1H	2518	A	C5-N7-C8	-9.59	99.10	103.90
26	14	1496	A	N7-C8-N9	9.56	118.58	113.80
26	1H	2331	G	N9-C4-C5	-9.56	101.58	105.40
26	14	2688	U	N3-C2-O2	-9.55	115.52	122.20
39	65	110	LEU	CA-CB-CG	9.54	137.25	115.30
26	14	801	G	N1-C6-O6	-9.54	114.18	119.90
26	1H	837	C	C6-N1-C2	-9.51	116.50	120.30
26	1H	2439	A	OP1-P-O3'	9.50	126.10	105.20
26	14	574	C	C6-N1-C2	9.50	124.10	120.30
26	14	2544	G	C5-C6-O6	-9.49	122.90	128.60
26	1H	1021	A	C5-N7-C8	-9.48	99.16	103.90
26	1H	621	A	C2-N3-C4	-9.47	105.86	110.60
26	1H	676	A	N3-C4-C5	9.45	133.41	126.80
26	14	2518	A	N1-C6-N6	9.44	124.27	118.60
26	1H	2450	A	O5'-P-OP2	-9.40	97.24	105.70
26	14	933	A	N1-C6-N6	9.40	124.24	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	2392	A	C5-N7-C8	-9.38	99.21	103.90
26	1H	1698	A	C2-N3-C4	-9.37	105.92	110.60
26	1H	140	A	N7-C8-N9	9.36	118.48	113.80
26	14	1610	A	N1-C6-N6	9.36	124.22	118.60
26	1H	1899	G	N9-C4-C5	9.33	109.13	105.40
26	14	669	G	C2-N3-C4	9.33	116.56	111.90
1	13	1498	U	P-O3'-C3'	9.31	130.88	119.70
26	14	1327	C	N1-C2-O2	-9.30	113.32	118.90
26	14	71	A	C5-N7-C8	-9.29	99.26	103.90
26	14	2430	A	N1-C2-N3	9.27	133.94	129.30
26	1H	945	A	C5-N7-C8	-9.26	99.27	103.90
26	14	391	G	C6-C5-N7	-9.26	124.85	130.40
26	1H	2688	U	N3-C2-O2	-9.25	115.73	122.20
26	1H	676	A	C4-C5-N7	9.22	115.31	110.70
26	14	140	A	C5-N7-C8	-9.22	99.29	103.90
1	13	792	A	C5-C6-N6	-9.19	116.35	123.70
26	1H	1786	A	C4-C5-N7	9.17	115.29	110.70
26	1H	2440	C	O5'-P-OP1	-9.17	97.45	105.70
26	1H	127	A	C5-C6-N6	-9.16	116.37	123.70
26	1H	1931	U	N3-C2-O2	-9.15	115.79	122.20
1	13	690	G	N7-C8-N9	9.12	117.66	113.10
26	14	2012	G	C5-C6-O6	-9.12	123.13	128.60
26	1H	946	G	C8-N9-C4	9.12	110.05	106.40
26	14	2873	A	C6-C5-N7	-9.12	125.92	132.30
1	13	690	G	C4-C5-N7	9.10	114.44	110.80
26	1H	1950	G	N7-C8-N9	9.10	117.65	113.10
26	1H	1395	A	O4'-C1'-N9	9.09	115.47	108.20
1	13	623	C	C6-N1-C2	-9.06	116.67	120.30
1	1G	529	G	C5-C6-O6	-9.06	123.16	128.60
26	14	1989	G	N3-C2-N2	-9.05	113.56	119.90
1	13	792	A	O4'-C1'-N9	9.05	115.44	108.20
1	13	186	C	C6-N1-C2	-9.05	116.68	120.30
26	14	2430	A	C6-C5-N7	-9.03	125.98	132.30
26	14	2544	G	N1-C6-O6	9.03	125.31	119.90
26	1H	860	U	C4-C5-C6	9.02	125.11	119.70
26	1H	2576	G	O5'-P-OP1	-8.97	97.62	105.70
26	1H	2330	G	N1-C6-O6	8.97	125.28	119.90
26	1H	1350	C	O5'-P-OP1	-8.96	97.64	105.70
54	Q8	28	GLY	N-CA-C	-8.95	90.73	113.10
26	14	1616	A	C5-N7-C8	-8.94	99.43	103.90
26	14	2713	A	C5-N7-C8	-8.93	99.44	103.90
26	1H	461	C	N1-C2-O2	-8.91	113.55	118.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
27	16	81	G	C5-N7-C8	-8.89	99.85	104.30
22	1K	87	A	N7-C8-N9	8.86	118.23	113.80
26	1H	2567	G	O5'-P-OP1	-8.86	97.73	105.70
1	13	792	A	C5-N7-C8	-8.84	99.48	103.90
26	14	664	C	C5-C6-N1	-8.84	116.58	121.00
26	14	1496	A	C5-N7-C8	-8.84	99.48	103.90
26	1H	2710	C	C5-C6-N1	-8.82	116.59	121.00
26	1H	2346	A	N1-C2-N3	8.81	133.71	129.30
26	1H	1616	A	N7-C8-N9	8.79	118.20	113.80
26	14	1786	A	N1-C2-N3	8.79	133.69	129.30
26	14	2681	C	C5-C4-N4	8.78	126.35	120.20
26	14	686	G	O5'-P-OP2	-8.78	97.80	105.70
26	14	2506	U	C2-N1-C1'	8.78	128.24	117.70
26	1H	121	G	C4-N9-C1'	8.77	137.90	126.50
26	14	1698	A	N9-C4-C5	-8.76	102.30	105.80
26	14	2346	A	N1-C2-N3	8.76	133.68	129.30
26	14	586	A	O5'-P-OP2	-8.75	97.83	105.70
26	14	2713	A	N1-C6-N6	8.74	123.84	118.60
26	14	2477	C	N1-C2-O2	8.73	124.14	118.90
26	14	1276	A	C5-C6-N6	-8.72	116.72	123.70
26	1H	1816	G	O5'-P-OP1	-8.71	97.86	105.70
1	1G	690	G	N3-C4-C5	8.71	132.96	128.60
26	1H	784	A	N1-C6-N6	-8.71	113.38	118.60
26	1H	2258	C	O5'-P-OP1	-8.71	97.86	105.70
1	1G	690	G	N3-C4-N9	-8.70	120.78	126.00
26	14	140	A	C4-C5-N7	8.68	115.04	110.70
26	1H	71	A	C4-C5-N7	8.67	115.04	110.70
1	1G	1278	U	C2-N1-C1'	8.67	128.10	117.70
26	14	330	A	N1-C2-N3	8.66	133.63	129.30
26	14	512	G	N3-C4-N9	-8.66	120.81	126.00
26	1H	1698	A	N1-C2-N3	8.64	133.62	129.30
26	14	71	A	O4'-C1'-N9	-8.64	101.29	108.20
26	1H	859	G	N3-C4-C5	8.62	132.91	128.60
26	14	1787	A	O5'-P-OP1	-8.62	97.94	105.70
26	14	2873	A	N1-C2-N3	8.61	133.60	129.30
26	14	1396	U	N3-C2-O2	-8.61	116.18	122.20
26	1H	945	A	C4-N9-C1'	8.58	141.74	126.30
26	1H	530	G	N1-C6-O6	-8.57	114.75	119.90
26	14	1695	G	C6-C5-N7	-8.57	125.26	130.40
26	14	140	A	N1-C6-N6	8.55	123.73	118.60
26	14	1790	C	N1-C2-O2	-8.55	113.77	118.90
26	14	774	A	N1-C6-N6	8.54	123.72	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	2584	U	N1-C2-N3	8.53	120.02	114.90
26	1H	676	A	O4'-C1'-N9	8.53	115.02	108.20
26	1H	1332	G	C4-C5-N7	8.53	114.21	110.80
1	1G	1322	C	C2-N1-C1'	8.52	128.17	118.80
26	14	912	C	C6-N1-C2	-8.51	116.90	120.30
1	13	652	U	C5-C6-N1	8.51	126.95	122.70
1	1G	1281	U	C5-C6-N1	8.51	126.95	122.70
26	14	1950	G	C4-N9-C1'	8.50	137.55	126.50
26	1H	2346	A	C2-N3-C4	-8.49	106.35	110.60
26	14	1610	A	C5-C6-N6	-8.47	116.92	123.70
26	14	330	A	C6-C5-N7	-8.47	126.37	132.30
26	1H	676	A	N1-C6-N6	8.46	123.68	118.60
26	1H	679	C	C6-N1-C2	8.46	123.68	120.30
26	1H	71	A	N1-C6-N6	8.46	123.67	118.60
26	14	945	A	C6-C5-N7	-8.45	126.39	132.30
1	13	1203	C	C6-N1-C2	-8.44	116.92	120.30
26	14	566	U	C5-C6-N1	-8.44	118.48	122.70
26	14	2447	G	O4'-C1'-N9	8.44	114.95	108.20
26	1H	127	A	C4-C5-N7	8.44	114.92	110.70
26	14	2709	G	N1-C6-O6	8.44	124.96	119.90
26	1H	1496	A	N7-C8-N9	8.43	118.02	113.80
26	14	933	A	C5-N7-C8	-8.43	99.68	103.90
26	1H	2595	G	C5-C6-O6	-8.43	123.54	128.60
1	13	792	A	C6-C5-N7	-8.42	126.41	132.30
26	14	676	A	N7-C8-N9	8.41	118.00	113.80
26	1H	1616	A	C5-N7-C8	-8.40	99.70	103.90
26	1H	1496	A	C5-N7-C8	-8.39	99.71	103.90
26	1H	2287	A	C2-N3-C4	-8.38	106.41	110.60
26	1H	831	G	C8-N9-C4	8.37	109.75	106.40
26	1H	2330	G	C2-N3-C4	-8.37	107.72	111.90
26	14	783	A	C4-C5-N7	8.37	114.88	110.70
26	1H	736	C	N1-C2-O2	-8.36	113.88	118.90
26	14	1950	G	C8-N9-C4	-8.36	103.06	106.40
26	14	991	C	O5'-P-OP1	-8.36	98.18	105.70
26	1H	1616	A	C6-C5-N7	-8.34	126.46	132.30
26	1H	1614	A	N1-C6-N6	8.32	123.59	118.60
26	1H	1899	G	C8-N9-C1'	8.32	137.81	127.00
26	1H	2430	A	C6-C5-N7	-8.32	126.48	132.30
26	1H	1496	A	C6-C5-N7	-8.31	126.48	132.30
26	1H	2008	C	O5'-P-OP2	-8.30	98.23	105.70
26	1H	2595	G	C4-C5-N7	8.30	114.12	110.80
1	13	1281	U	N3-C2-O2	-8.29	116.39	122.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	2518	A	N7-C8-N9	8.29	117.95	113.80
26	14	2755	C	C2-N1-C1'	8.28	127.91	118.80
26	1H	1313	U	C2-N1-C1'	8.28	127.63	117.70
26	1H	528	A	N3-C4-N9	-8.27	120.78	127.40
26	1H	2330	G	C8-N9-C4	8.27	109.71	106.40
1	13	1523	G	C5-C6-N1	-8.26	107.37	111.50
26	1H	51	G	O5'-P-OP1	-8.26	98.26	105.70
26	14	810	U	C5-C4-O4	-8.26	120.94	125.90
26	1H	2062	A	C8-N9-C4	8.25	109.10	105.80
26	1H	1942	C	N3-C4-C5	8.25	125.20	121.90
26	1H	609	A	N1-C6-N6	8.25	123.55	118.60
26	1H	1784	A	N1-C6-N6	-8.23	113.66	118.60
1	1G	1260	C	C6-N1-C2	-8.23	117.01	120.30
26	14	2447	G	C8-N9-C1'	8.23	137.69	127.00
26	1H	121	G	N1-C6-O6	8.22	124.83	119.90
26	1H	186	G	N1-C6-O6	8.22	124.83	119.90
26	1H	512	G	O4'-C1'-N9	8.21	114.77	108.20
1	13	505	G	N1-C6-O6	8.20	124.82	119.90
26	14	1328	G	N9-C4-C5	-8.20	102.12	105.40
26	1H	2256	G	O5'-P-OP2	-8.20	98.32	105.70
1	1G	1260	C	C5-C6-N1	8.20	125.10	121.00
26	1H	2392	A	C8-N9-C4	-8.19	102.52	105.80
1	13	690	G	C5-N7-C8	-8.19	100.21	104.30
26	1H	464	U	N3-C2-O2	-8.17	116.48	122.20
26	1H	1021	A	C4-C5-N7	8.17	114.79	110.70
26	1H	140	A	C6-C5-N7	-8.16	126.58	132.30
26	1H	1602	U	C6-N1-C2	8.16	125.90	121.00
26	1H	672	C	O5'-P-OP1	8.16	120.49	110.70
26	1H	1698	A	C6-C5-N7	-8.15	126.59	132.30
26	1H	263	C	O5'-P-OP2	-8.15	98.36	105.70
26	14	1678	G	N3-C4-C5	8.15	132.68	128.60
1	13	690	G	N1-C6-O6	8.15	124.79	119.90
26	1H	1332	G	N3-C4-C5	8.15	132.67	128.60
26	1H	2259	G	N1-C6-O6	8.14	124.79	119.90
26	1H	2490	G	N3-C4-C5	8.14	132.67	128.60
26	1H	917	A	N1-C2-N3	8.14	133.37	129.30
26	14	1698	A	C4-C5-N7	8.14	114.77	110.70
1	13	247	G	N3-C4-C5	-8.13	124.53	128.60
26	1H	1496	A	C4-C5-N7	8.13	114.77	110.70
26	14	774	A	C5-N7-C8	-8.12	99.84	103.90
26	14	2518	A	N3-C4-C5	8.11	132.48	126.80
26	14	1698	A	C2-N3-C4	-8.11	106.55	110.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	2609	U	C2-N3-C4	-8.10	122.14	127.00
26	1H	138	G	C8-N9-C4	-8.09	103.16	106.40
26	14	140	A	N7-C8-N9	8.09	117.84	113.80
1	13	1305	G	C8-N9-C4	-8.08	103.17	106.40
22	1K	87	A	C5-N7-C8	-8.08	99.86	103.90
1	1G	529	G	N1-C6-O6	8.07	124.74	119.90
26	14	1332	G	N9-C4-C5	8.07	108.63	105.40
26	14	330	A	C5-N7-C8	-8.06	99.87	103.90
26	14	512	G	N3-C4-C5	8.06	132.63	128.60
26	1H	2258	C	C6-N1-C2	-8.06	117.08	120.30
26	14	2873	A	C4-C5-N7	8.05	114.73	110.70
26	1H	121	G	C8-N9-C1'	-8.04	116.54	127.00
26	1H	1611	C	C6-N1-C2	8.04	123.52	120.30
26	14	2490	G	N7-C8-N9	8.03	117.12	113.10
26	1H	247	G	C8-N9-C4	8.03	109.61	106.40
26	1H	2503	A	C5-C6-N6	-8.01	117.29	123.70
26	1H	676	A	C5-C6-N1	-8.01	113.69	117.70
26	14	2779	U	N3-C2-O2	-8.00	116.60	122.20
26	1H	1616	A	N1-C6-N6	7.99	123.40	118.60
26	1H	1528	A	C8-N9-C4	-7.99	102.61	105.80
26	1H	2565	A	C8-N9-C4	7.99	108.99	105.80
26	1H	59	U	C6-N1-C2	-7.98	116.21	121.00
26	1H	1379	A	N1-C6-N6	7.98	123.39	118.60
26	14	74	A	C5-N7-C8	-7.98	99.91	103.90
26	1H	528	A	N3-C4-C5	7.96	132.38	126.80
26	14	801	G	C5-C6-O6	7.96	133.38	128.60
26	1H	593	G	O5'-P-OP2	-7.96	98.54	105.70
26	14	241	A	O5'-P-OP2	-7.96	98.54	105.70
26	1H	774	A	C5-C6-N1	-7.95	113.72	117.70
26	1H	139	G	O5'-P-OP1	-7.95	98.54	105.70
26	1H	801	G	C5-C6-O6	7.95	133.37	128.60
26	14	2584	U	C2-N1-C1'	7.95	127.24	117.70
26	1H	74	A	N1-C2-N3	7.94	133.27	129.30
32	59	153	LYS	C-N-CD	7.94	145.07	128.40
26	1H	1829	A	O5'-P-OP1	-7.93	98.56	105.70
26	1H	828	U	N3-C2-O2	-7.93	116.65	122.20
26	14	2430	A	C4-C5-C6	7.93	120.97	117.00
26	14	2447	G	O5'-P-OP1	-7.92	98.57	105.70
26	1H	141	A	C5-N7-C8	-7.91	99.94	103.90
26	1H	729	G	C8-N9-C4	-7.91	103.24	106.40
26	1H	695	G	O5'-P-OP2	-7.90	98.59	105.70
26	1H	918	A	O5'-P-OP1	-7.89	98.60	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	208	C	C6-N1-C2	7.88	123.45	120.30
26	14	797	C	N3-C4-N4	7.88	123.51	118.00
1	13	1530	G	C5-C6-O6	-7.87	123.88	128.60
26	14	1950	G	O4'-C1'-N9	7.85	114.48	108.20
26	1H	2409	G	C4-C5-N7	7.84	113.94	110.80
26	14	71	A	C4-C5-N7	7.84	114.62	110.70
26	1H	2424	C	OP1-P-OP2	7.84	131.36	119.60
1	13	23	C	C6-N1-C2	-7.84	117.17	120.30
26	14	783	A	N1-C6-N6	7.83	123.30	118.60
26	1H	2509	G	N1-C6-O6	7.83	124.60	119.90
26	1H	140	A	C2-N3-C4	-7.83	106.68	110.60
26	14	933	A	C4-C5-N7	7.83	114.62	110.70
38	55	28	LEU	CA-CB-CG	7.83	133.31	115.30
26	1H	1363	C	C5-C6-N1	-7.83	117.08	121.00
26	1H	1026	U	N3-C2-O2	-7.83	116.72	122.20
26	14	265	A	C5-N7-C8	-7.82	99.99	103.90
1	13	802	A	N1-C6-N6	7.82	123.29	118.60
26	1H	1241	A	C5-C6-N1	-7.82	113.79	117.70
26	1H	1830	C	N1-C2-O2	-7.81	114.22	118.90
1	1G	1297	C	P-O3'-C3'	7.81	129.07	119.70
26	1H	2338	G	C5-C6-O6	-7.80	123.92	128.60
26	14	1616	A	C4-C5-N7	7.79	114.60	110.70
26	1H	824	A	O5'-P-OP1	-7.79	98.69	105.70
27	16	81	G	C6-C5-N7	-7.78	125.73	130.40
26	1H	678	C	C5-C6-N1	-7.77	117.11	121.00
26	14	2346	A	C2-N3-C4	-7.77	106.71	110.60
1	13	974	A	N1-C6-N6	7.77	123.26	118.60
26	14	801	G	C6-C5-N7	7.77	135.06	130.40
26	14	621	A	C2-N3-C4	-7.77	106.72	110.60
26	1H	127	A	C6-C5-N7	-7.75	126.87	132.30
1	13	1158	C	N1-C2-O2	7.75	123.55	118.90
26	1H	2584	U	C5-C4-O4	7.75	130.55	125.90
26	1H	127	A	C2-N3-C4	-7.74	106.73	110.60
1	1G	568	G	C8-N9-C4	-7.74	103.30	106.40
26	14	1496	A	N1-C6-N6	7.74	123.24	118.60
26	1H	2713	A	C2-N3-C4	-7.74	106.73	110.60
27	16	12	C	C5-C6-N1	-7.73	117.13	121.00
23	2K	3	C	C5-C6-N1	7.73	124.87	121.00
26	14	1260	G	N1-C6-O6	7.73	124.54	119.90
1	1G	817	C	C6-N1-C2	7.72	123.39	120.30
26	1H	684	G	N7-C8-N9	7.72	116.96	113.10
26	1H	1611	C	C5-C6-N1	-7.71	117.15	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	988	A	O5'-P-OP1	-7.70	98.77	105.70
26	14	1671	U	O5'-P-OP1	-7.70	98.77	105.70
26	1H	2447	G	C5-C6-O6	-7.69	123.99	128.60
26	1H	1544	C	N1-C2-O2	7.68	123.51	118.90
26	14	659	C	C5-C6-N1	-7.68	117.16	121.00
26	1H	1786	A	C6-C5-N7	-7.68	126.93	132.30
1	13	1053	G	P-O3'-C3'	7.67	128.91	119.70
26	14	2386	C	C6-N1-C2	7.67	123.37	120.30
26	14	621	A	C5-C6-N1	-7.67	113.86	117.70
1	1G	895	G	N1-C6-O6	7.65	124.49	119.90
54	Q8	60	LEU	CA-CB-CG	7.65	132.89	115.30
26	1H	74	A	C5-C6-N1	-7.64	113.88	117.70
26	1H	127	A	N9-C4-C5	-7.64	102.74	105.80
26	14	2447	G	C4-N9-C1'	-7.64	116.57	126.50
26	1H	856	C	O5'-P-OP1	-7.63	98.83	105.70
27	16	30	C	C6-N1-C2	-7.63	117.25	120.30
26	14	1950	G	C6-C5-N7	-7.63	125.82	130.40
1	13	690	G	C8-N9-C1'	-7.62	117.09	127.00
26	1H	2069	G	C5-C6-O6	-7.62	124.03	128.60
26	14	1275	A	N1-C6-N6	7.62	123.17	118.60
26	1H	1799	G	N3-C4-C5	-7.62	124.79	128.60
26	14	2012	G	C6-C5-N7	-7.61	125.83	130.40
26	14	2518	A	C5-N7-C8	-7.60	100.10	103.90
26	1H	1332	G	N7-C8-N9	7.60	116.90	113.10
26	1H	120	U	C4-C5-C6	7.59	124.26	119.70
26	14	2610	C	N1-C2-O2	7.59	123.45	118.90
26	1H	138	G	N7-C8-N9	7.59	116.89	113.10
26	1H	812	C	N1-C2-O2	-7.59	114.35	118.90
1	13	974	A	O4'-C1'-N9	7.59	114.27	108.20
26	14	2607	G	N1-C6-O6	7.58	124.45	119.90
26	14	2755	C	C5-C6-N1	7.58	124.79	121.00
26	1H	1989	G	N3-C2-N2	-7.58	114.59	119.90
26	14	1981	A	N1-C6-N6	7.58	123.15	118.60
27	16	60	C	C5-C6-N1	7.58	124.79	121.00
26	14	2437	U	C5-C4-O4	7.57	130.44	125.90
1	1G	527	G	C5-C6-O6	7.57	133.14	128.60
26	14	512	G	C4-N9-C1'	-7.57	116.66	126.50
26	14	1204	A	O4'-C1'-N9	7.56	114.25	108.20
26	1H	1025	G	N1-C6-O6	-7.56	115.36	119.90
26	1H	860	U	C5-C6-N1	-7.56	118.92	122.70
26	14	810	U	N3-C4-O4	7.55	124.69	119.40
26	14	74	A	C6-C5-N7	-7.55	127.01	132.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	676	A	O4'-C1'-N9	7.55	114.24	108.20
1	13	505	G	C5-C6-O6	-7.55	124.07	128.60
26	14	1328	G	C4-C5-N7	7.55	113.82	110.80
1	13	1530	G	N1-C6-O6	7.54	124.43	119.90
26	14	2275	C	P-O3'-C3'	7.54	128.75	119.70
26	14	2440	C	O5'-P-OP1	-7.54	98.91	105.70
26	14	1786	A	C6-C5-N7	-7.54	127.02	132.30
26	14	929	G	N1-C6-O6	7.53	124.42	119.90
26	1H	682	G	O5'-P-OP2	-7.53	98.93	105.70
26	1H	835	A	C6-N1-C2	-7.53	114.08	118.60
26	1H	120	U	C5-C6-N1	-7.52	118.94	122.70
26	14	801	G	O5'-P-OP2	-7.52	98.93	105.70
26	1H	2430	A	N1-C2-N3	7.51	133.05	129.30
26	1H	324	A	O5'-P-OP1	-7.50	98.95	105.70
1	1G	87	A	P-O3'-C3'	7.50	128.70	119.70
26	1H	2331	G	C5-C6-O6	-7.50	124.10	128.60
26	1H	1382	G	C5-C6-O6	-7.49	124.11	128.60
26	1H	860	U	N3-C2-O2	-7.49	116.96	122.20
26	14	2681	C	N3-C4-N4	-7.49	112.76	118.00
26	1H	494	G	C5-C6-O6	-7.48	124.11	128.60
26	14	1328	G	N3-C4-N9	7.47	130.49	126.00
26	1H	964	C	C6-N1-C2	-7.47	117.31	120.30
26	14	329	G	N1-C6-O6	-7.47	115.42	119.90
1	13	1201	A	N1-C6-N6	7.46	123.08	118.60
26	1H	831	G	N7-C8-N9	-7.46	109.37	113.10
26	1H	945	A	O4'-C1'-N9	7.46	114.17	108.20
26	1H	1972	A	C5-C6-N6	-7.46	117.73	123.70
26	1H	774	A	C5-N7-C8	-7.45	100.18	103.90
23	2L	33	U	C5-C6-N1	7.45	126.42	122.70
32	59	93	GLY	N-CA-C	7.45	131.72	113.10
26	14	1698	A	C6-C5-N7	-7.44	127.09	132.30
26	1H	1513	C	C5-C6-N1	7.44	124.72	121.00
26	14	2392	A	C5-C6-N1	-7.44	113.98	117.70
26	14	774	A	C4-C5-N7	7.44	114.42	110.70
26	14	2702	U	O4'-C1'-N1	7.44	114.15	108.20
23	2L	20	U	P-O3'-C3'	7.43	128.62	119.70
26	14	783	A	N7-C8-N9	7.43	117.52	113.80
26	14	2581	G	C8-N9-C4	-7.43	103.43	106.40
26	14	1332	G	C8-N9-C4	-7.43	103.43	106.40
1	13	1305	G	N3-C2-N2	-7.42	114.70	119.90
26	1H	1210	A	N7-C8-N9	7.42	117.51	113.80
26	14	1667	G	N1-C6-O6	7.42	124.35	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	1G	197	A	N7-C8-N9	7.41	117.51	113.80
26	14	1695	G	N3-C4-N9	7.41	130.45	126.00
26	1H	121	G	N3-C4-C5	-7.41	124.90	128.60
26	14	747	U	N3-C2-O2	7.41	127.38	122.20
23	2L	16	U	C2-N1-C1'	7.40	126.58	117.70
11	2I	102	GLY	N-CA-C	-7.40	94.60	113.10
26	1H	2503	A	N9-C4-C5	-7.40	102.84	105.80
26	1H	621	A	C5-N7-C8	-7.40	100.20	103.90
26	1H	1026	U	N1-C2-O2	7.40	127.98	122.80
26	14	1565	C	N1-C2-O2	-7.39	114.47	118.90
36	35	147	LEU	CA-CB-CG	7.39	132.30	115.30
26	1H	2392	A	C5-C6-N1	-7.39	114.00	117.70
1	1G	1322	C	C5-C6-N1	7.39	124.69	121.00
26	14	2713	A	N7-C8-N9	7.38	117.49	113.80
26	14	689	A	O5'-P-OP2	-7.38	99.06	105.70
26	14	1594	G	C8-N9-C4	-7.38	103.45	106.40
26	1H	756	C	N1-C2-O2	-7.38	114.47	118.90
26	1H	1528	A	O4'-C1'-N9	7.37	114.10	108.20
1	1G	345	C	C2-N1-C1'	7.37	126.91	118.80
26	14	1695	G	C4-N9-C1'	7.37	136.08	126.50
26	14	1260	G	C5-C6-O6	-7.37	124.18	128.60
26	1H	676	A	N3-C4-N9	-7.36	121.51	127.40
26	1H	796	C	C6-N1-C2	7.36	123.24	120.30
26	1H	1413	G	N1-C6-O6	7.36	124.31	119.90
24	3K	3	C	C6-N1-C2	-7.36	117.36	120.30
26	1H	920	G	C8-N9-C4	7.35	109.34	106.40
26	14	1332	G	C5-N7-C8	-7.35	100.62	104.30
26	1H	265	A	C5-N7-C8	-7.35	100.22	103.90
1	1G	793	U	O4'-C1'-N1	7.35	114.08	108.20
26	14	2447	G	C4-C5-N7	-7.35	107.86	110.80
26	14	2518	A	C4-C5-N7	7.35	114.37	110.70
26	1H	74	A	N7-C8-N9	7.34	117.47	113.80
1	1G	898	G	C8-N9-C4	7.34	109.34	106.40
1	13	481	G	C5-C6-N1	-7.34	107.83	111.50
1	1G	254	G	O5'-P-OP1	-7.34	99.09	105.70
26	14	2607	G	C5-C6-O6	-7.34	124.20	128.60
26	1H	783	A	C5-C6-N1	-7.34	114.03	117.70
26	1H	1606	G	C8-N9-C4	7.34	109.33	106.40
26	14	582	G	N1-C6-O6	7.34	124.30	119.90
1	1G	890	G	O4'-C1'-N9	7.33	114.06	108.20
26	14	1989	G	N9-C4-C5	7.32	108.33	105.40
1	13	1523	G	N1-C6-O6	7.32	124.29	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	1K	87	A	C8-N9-C4	-7.31	102.88	105.80
1	13	115	G	P-O3'-C3'	7.31	128.47	119.70
26	1H	945	A	C4-C5-N7	7.31	114.36	110.70
26	14	2430	A	O5'-P-OP1	-7.31	99.12	105.70
26	14	952	G	C8-N9-C4	-7.31	103.48	106.40
26	14	2518	A	O4'-C1'-N9	-7.30	102.36	108.20
26	1H	1431	U	C5-C6-N1	7.30	126.35	122.70
26	1H	1359	A	N1-C2-N3	7.30	132.95	129.30
26	1H	1899	G	C8-N9-C4	-7.30	103.48	106.40
26	1H	530	G	N3-C2-N2	7.30	125.01	119.90
29	29	78	LEU	CA-CB-CG	7.29	132.08	115.30
1	13	812	C	N3-C2-O2	-7.29	116.80	121.90
26	1H	2506	U	N1-C2-O2	7.29	127.91	122.80
26	1H	1602	U	O5'-P-OP2	7.29	119.45	110.70
26	14	613	U	N3-C2-O2	-7.29	117.10	122.20
26	1H	2591	C	N1-C2-O2	-7.29	114.53	118.90
1	13	353	A	C8-N9-C4	-7.29	102.89	105.80
26	1H	1698	A	N7-C8-N9	7.28	117.44	113.80
26	14	929	G	C6-C5-N7	-7.28	126.03	130.40
26	14	2401	U	C5-C6-N1	7.28	126.34	122.70
26	14	676	A	C5-C6-N1	-7.28	114.06	117.70
26	1H	2060	A	P-O3'-C3'	7.27	128.43	119.70
26	1H	2374	C	C5-C6-N1	-7.27	117.36	121.00
26	14	1520	U	C5-C4-O4	7.26	130.26	125.90
26	14	774	A	C5-C6-N1	-7.25	114.07	117.70
26	1H	213	A	N1-C6-N6	7.25	122.95	118.60
26	1H	1241	A	C2-N3-C4	-7.25	106.97	110.60
26	14	2477	C	C2-N1-C1'	7.25	126.77	118.80
26	1H	856	C	C6-N1-C2	-7.24	117.40	120.30
26	14	2447	G	N9-C4-C5	7.24	108.30	105.40
26	1H	71	A	N3-C4-N9	-7.24	121.61	127.40
26	14	1233	C	C6-N1-C2	-7.24	117.41	120.30
26	1H	504	U	C6-N1-C1'	-7.23	111.07	121.20
26	14	12	U	N3-C2-O2	-7.23	117.14	122.20
26	1H	2698	U	C5-C6-N1	-7.23	119.08	122.70
26	1H	1310	G	C5-C6-O6	-7.23	124.27	128.60
26	14	2880	C	C6-N1-C2	-7.22	117.41	120.30
26	1H	864	G	C5-C6-O6	-7.21	124.27	128.60
26	1H	609	A	C5-C6-N6	-7.21	117.93	123.70
27	1J	8	U	O5'-P-OP2	-7.21	99.21	105.70
26	14	774	A	N3-C4-N9	-7.21	121.63	127.40
26	14	974(A)	C	C6-N1-C2	-7.21	117.42	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	1496	A	C6-C5-N7	-7.20	127.26	132.30
26	1H	2688	U	C5-C4-O4	7.20	130.22	125.90
26	14	2430	A	C5-C6-N1	-7.20	114.10	117.70
1	13	186	C	C5-C6-N1	7.20	124.60	121.00
26	1H	2553	G	C8-N9-C1'	-7.20	117.65	127.00
26	1H	1332	G	N1-C2-N3	7.19	128.22	123.90
26	1H	38	A	C5-C6-N6	-7.19	117.95	123.70
26	14	439	G	C5-C6-O6	-7.19	124.28	128.60
26	14	567	A	C4-C5-C6	7.19	120.59	117.00
26	14	1328	G	C6-C5-N7	-7.18	126.09	130.40
27	16	81	G	C5-C6-O6	-7.18	124.29	128.60
26	14	4	C	C2-N1-C1'	7.18	126.70	118.80
26	14	2346	A	O4'-C1'-N9	7.18	113.94	108.20
26	1H	440	G	C5-C6-O6	-7.18	124.29	128.60
26	14	2622	C	C6-N1-C2	7.17	123.17	120.30
1	1G	690	G	C5-N7-C8	-7.17	100.71	104.30
26	1H	2583	G	N1-C2-N2	-7.17	109.75	116.20
1	13	860	A	N1-C6-N6	7.16	122.90	118.60
45	C5	103	GLY	N-CA-C	7.16	131.00	113.10
27	16	64	C	C6-N1-C2	7.15	123.16	120.30
26	1H	494	G	N1-C6-O6	7.15	124.19	119.90
26	1H	2490	G	C5-N7-C8	-7.15	100.72	104.30
26	1H	807	U	N3-C4-O4	7.15	124.40	119.40
26	14	945	A	N3-C4-C5	7.15	131.80	126.80
29	21	65	GLY	N-CA-C	-7.15	95.23	113.10
26	1H	2212	A	O4'-C1'-N9	7.14	113.91	108.20
28	19	272	ALA	N-CA-C	7.14	130.28	111.00
41	C8	74	LEU	CA-CB-CG	7.14	131.72	115.30
26	14	1902	C	N3-C4-N4	-7.14	113.00	118.00
1	1G	197	A	C8-N9-C4	-7.13	102.95	105.80
26	1H	1275	A	N1-C6-N6	7.13	122.88	118.60
26	14	74	A	N3-C4-C5	7.13	131.79	126.80
27	16	47	C	C6-N1-C2	7.12	123.15	120.30
27	1J	114	G	N3-C4-C5	7.12	132.16	128.60
26	1H	188	G	C8-N9-C4	7.12	109.25	106.40
27	16	70	C	C6-N1-C2	-7.12	117.45	120.30
26	1H	1559	G	N3-C4-C5	7.11	132.16	128.60
26	14	580	C	C6-N1-C2	-7.11	117.46	120.30
26	14	2755	C	C6-N1-C2	-7.11	117.46	120.30
26	1H	1332	G	C6-C5-N7	-7.11	126.13	130.40
26	1H	945	A	N1-C2-N3	7.11	132.85	129.30
1	13	1158	C	C2-N1-C1'	7.10	126.61	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	945	A	C2-N3-C4	-7.10	107.05	110.60
26	14	1695	G	C8-N9-C1'	-7.10	117.77	127.00
26	14	1634	A	N1-C6-N6	-7.10	114.34	118.60
26	1H	74	A	C5-N7-C8	-7.09	100.35	103.90
26	1H	2713	A	C5-N7-C8	-7.09	100.35	103.90
26	1H	1695	G	N3-C4-N9	7.09	130.26	126.00
26	1H	2439	A	C8-N9-C4	-7.09	102.96	105.80
26	14	2012	G	C4-C5-N7	7.08	113.63	110.80
1	13	700	G	C6-C5-N7	-7.08	126.15	130.40
26	1H	2830	G	C8-N9-C4	-7.07	103.57	106.40
26	14	752	A	C8-N9-C4	-7.07	102.97	105.80
26	14	783	A	N3-C4-C5	7.07	131.75	126.80
26	1H	736	C	O5'-P-OP1	-7.07	99.34	105.70
26	1H	1678	G	C2-N3-C4	-7.07	108.37	111.90
26	14	828	U	N1-C2-O2	7.06	127.74	122.80
26	1H	2447	G	N1-C6-O6	7.06	124.14	119.90
26	1H	2575	C	N3-C4-C5	-7.06	119.08	121.90
26	14	512	G	C8-N9-C1'	7.05	136.17	127.00
26	14	2323	G	C8-N9-C4	7.05	109.22	106.40
26	1H	698	C	C6-N1-C2	7.05	123.12	120.30
26	14	1610	A	N9-C4-C5	-7.05	102.98	105.80
26	14	2287	A	C5-C6-N1	-7.05	114.18	117.70
26	14	2688	U	C5-C4-O4	7.05	130.13	125.90
1	13	1354	C	C6-N1-C2	-7.04	117.48	120.30
26	1H	864	G	C4-C5-N7	7.04	113.62	110.80
26	14	685	A	C8-N9-C4	-7.04	102.98	105.80
26	1H	788	A	N1-C6-N6	7.03	122.82	118.60
26	1H	863	A	C5-C6-N1	7.03	121.22	117.70
26	1H	1558	A	P-O3'-C3'	7.03	128.14	119.70
26	1H	1349	A	C2-N3-C4	-7.03	107.09	110.60
26	1H	141	A	N7-C8-N9	7.03	117.31	113.80
26	14	1681	G	N3-C4-C5	7.03	132.11	128.60
26	14	2389	G	C8-N9-C4	-7.02	103.59	106.40
26	1H	2583	G	N3-C2-N2	7.02	124.81	119.90
1	1G	230	G	N1-C6-O6	7.02	124.11	119.90
26	14	801	G	N3-C4-N9	-7.01	121.79	126.00
26	14	2688	U	C4-C5-C6	7.01	123.91	119.70
26	14	621	A	C5-N7-C8	-7.00	100.40	103.90
26	1H	1968	G	C5-C6-O6	-7.00	124.40	128.60
26	1H	1204	A	C2-N3-C4	-7.00	107.10	110.60
26	1H	2287	A	N1-C2-N3	7.00	132.80	129.30
26	1H	1814	G	C4-C5-N7	7.00	113.60	110.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	13	975	A	N1-C6-N6	7.00	122.80	118.60
22	1L	85	C	C5-C4-N4	7.00	125.10	120.20
26	14	382	G	N1-C6-O6	7.00	124.10	119.90
26	14	2002	G	C5-C6-O6	-6.99	124.40	128.60
1	1G	1059	C	C5-C6-N1	6.99	124.49	121.00
26	14	621	A	N1-C6-N6	6.99	122.79	118.60
1	1G	1094	G	OP2-P-O3'	6.99	120.57	105.20
1	13	623	C	C5-C6-N1	6.98	124.49	121.00
26	1H	1899	G	N1-C2-N3	6.98	128.09	123.90
26	1H	2570	G	N1-C6-O6	6.98	124.09	119.90
26	1H	1528	A	N7-C8-N9	6.97	117.29	113.80
26	1H	871	U	N3-C4-O4	6.97	124.28	119.40
26	14	1312	U	N1-C2-O2	-6.97	117.92	122.80
26	14	1496	A	C8-N9-C4	-6.97	103.01	105.80
1	13	579	G	N1-C6-O6	6.96	124.08	119.90
26	1H	676	A	C8-N9-C4	-6.96	103.02	105.80
26	1H	1363	C	N3-C4-C5	6.96	124.69	121.90
1	1G	1278	U	N1-C2-O2	6.96	127.67	122.80
1	13	549	C	C2-N1-C1'	-6.96	111.14	118.80
26	1H	99	U	N3-C2-O2	-6.96	117.33	122.20
26	1H	1606	G	C5-C6-O6	-6.95	124.43	128.60
23	2L	27	G	C6-C5-N7	-6.95	126.23	130.40
1	13	413	G	O4'-C1'-N9	6.95	113.76	108.20
1	1G	897	C	N1-C2-O2	-6.95	114.73	118.90
26	14	1812	A	O5'-P-OP2	-6.95	99.44	105.70
26	14	1610	A	C4-C5-N7	6.94	114.17	110.70
26	1H	513	A	C6-N1-C2	-6.94	114.44	118.60
1	13	1502	A	C5-N7-C8	-6.93	100.43	103.90
26	1H	564	C	C6-N1-C2	-6.93	117.53	120.30
26	1H	945	A	C8-N9-C1'	-6.93	115.22	127.70
26	1H	1347	G	C5-C6-O6	-6.93	124.44	128.60
26	1H	1520	U	C5-C4-O4	6.93	130.06	125.90
1	13	904	C	N3-C4-C5	6.92	124.67	121.90
26	1H	2367	G	N1-C6-O6	6.92	124.05	119.90
1	1G	969	A	O5'-P-OP2	-6.92	99.48	105.70
1	1G	1346	A	P-O3'-C3'	6.92	128.00	119.70
26	14	1342	A	C2-N3-C4	-6.91	107.14	110.60
26	1H	1698	A	C5-N7-C8	-6.91	100.44	103.90
26	1H	2444	G	O5'-P-OP2	-6.91	99.48	105.70
54	Q8	52	LYS	C-N-CD	-6.91	105.39	120.60
26	1H	121	G	C4-C5-N7	6.91	113.56	110.80
26	1H	2310	A	C8-N9-C4	-6.91	103.04	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	1678	G	C5-N7-C8	-6.91	100.85	104.30
26	14	1825	A	C6-N1-C2	-6.91	114.46	118.60
26	1H	785	G	C8-N9-C4	-6.90	103.64	106.40
26	1H	2271	G	C5-C6-O6	-6.90	124.46	128.60
26	14	1602	U	O5'-P-OP2	6.90	118.98	110.70
26	1H	2295	C	C6-N1-C2	-6.90	117.54	120.30
1	13	766	A	N9-C4-C5	-6.89	103.04	105.80
26	1H	945	A	C8-N9-C4	-6.89	103.04	105.80
26	1H	1698	A	C8-N9-C4	-6.89	103.04	105.80
26	1H	1781	C	C2-N1-C1'	6.89	126.38	118.80
26	14	1899	G	N3-C4-C5	6.89	132.04	128.60
26	1H	621	A	O5'-P-OP1	-6.89	99.50	105.70
26	14	265	A	C2-N3-C4	-6.89	107.16	110.60
1	13	733	A	C8-N9-C4	6.88	108.55	105.80
26	1H	1376	C	C6-N1-C2	-6.88	117.55	120.30
26	14	1382	G	N3-C4-C5	6.88	132.04	128.60
26	14	49	A	C8-N9-C4	-6.88	103.05	105.80
26	14	2588	G	N1-C6-O6	-6.88	115.77	119.90
26	1H	1675	C	N3-C4-C5	-6.88	119.15	121.90
26	1H	2026	C	C5-C6-N1	-6.88	117.56	121.00
26	1H	1520	U	C6-N1-C2	-6.88	116.88	121.00
26	1H	2553	G	C6-C5-N7	-6.88	126.27	130.40
26	14	1661	G	O5'-P-OP2	-6.88	99.51	105.70
26	1H	2710	C	C2-N3-C4	-6.87	116.46	119.90
26	1H	2490	G	N3-C4-N9	-6.87	121.88	126.00
26	1H	2553	G	C4-N9-C1'	6.87	135.43	126.50
27	1J	114	G	C8-N9-C4	6.87	109.15	106.40
26	14	49	A	P-O3'-C3'	6.87	127.94	119.70
26	1H	197	A	OP2-P-O3'	6.86	120.30	105.20
26	14	439	G	N1-C6-O6	6.86	124.02	119.90
26	1H	1228	G	C8-N9-C4	-6.86	103.66	106.40
26	1H	821	A	C4-C5-C6	6.85	120.43	117.00
26	14	797	C	N3-C4-C5	-6.85	119.16	121.90
26	14	1616	A	O4'-C1'-N9	6.85	113.68	108.20
4	3E	12	CYS	CA-CB-SG	6.85	126.33	114.00
26	1H	1022	G	N9-C4-C5	6.85	108.14	105.40
26	14	2542	A	P-O3'-C3'	6.85	127.92	119.70
26	14	2873	A	C8-N9-C4	-6.85	103.06	105.80
26	1H	1543	A	N1-C6-N6	6.84	122.71	118.60
26	14	675	A	C4-C5-N7	6.84	114.12	110.70
26	14	2429	G	C8-N9-C4	-6.84	103.66	106.40
26	1H	189	G	C8-N9-C4	6.84	109.14	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	1G	913	A	C8-N9-C4	-6.84	103.06	105.80
26	1H	752	A	P-O3'-C3'	6.84	127.91	119.70
26	14	1914	C	N3-C2-O2	-6.84	117.11	121.90
26	14	40	C	C6-N1-C2	-6.83	117.57	120.30
1	1G	812	C	P-O3'-C3'	6.83	127.90	119.70
26	1H	1888	G	N3-C4-C5	-6.83	125.19	128.60
26	1H	2685	G	C5-C6-N1	-6.83	108.09	111.50
1	13	496	A	C8-N9-C4	-6.83	103.07	105.80
26	14	676	A	N9-C4-C5	-6.83	103.07	105.80
26	14	74	A	C4-C5-N7	6.82	114.11	110.70
26	14	1914	C	N1-C2-O2	6.82	122.99	118.90
26	1H	1298	C	C5-C6-N1	6.82	124.41	121.00
26	14	1328	G	C5-C6-O6	-6.82	124.51	128.60
26	14	1558	A	C2-N3-C4	-6.81	107.19	110.60
28	19	41	GLY	N-CA-C	-6.81	96.06	113.10
26	1H	1300	U	N1-C2-O2	-6.80	118.04	122.80
26	1H	1697	G	N1-C6-O6	6.80	123.98	119.90
45	G8	81	LYS	C-N-CD	-6.80	105.64	120.60
26	1H	850	C	C6-N1-C2	-6.80	117.58	120.30
26	1H	113	G	N1-C6-O6	6.79	123.98	119.90
26	1H	140	A	O4'-C1'-N9	6.79	113.63	108.20
1	13	766	A	C8-N9-C4	6.79	108.52	105.80
26	1H	115	C	O5'-P-OP1	-6.79	99.59	105.70
26	1H	2570	G	C5-C6-N1	-6.79	108.11	111.50
26	14	1667	G	N3-C4-C5	6.79	132.00	128.60
26	1H	1614	A	C2-N3-C4	-6.79	107.20	110.60
1	1G	529	G	C6-C5-N7	-6.79	126.33	130.40
1	1G	687	A	P-O3'-C3'	6.78	127.84	119.70
26	14	774	A	C6-N1-C2	6.78	122.67	118.60
26	14	57	C	C6-N1-C2	6.78	123.01	120.30
26	14	275	G	N3-C4-N9	6.78	130.07	126.00
22	1K	18	G	C4-N9-C1'	-6.78	117.69	126.50
26	14	914	C	N1-C2-O2	6.78	122.97	118.90
26	1H	38	A	C6-N1-C2	-6.78	114.53	118.60
26	1H	2451	A	C8-N9-C4	-6.78	103.09	105.80
26	14	2518	A	N9-C4-C5	-6.78	103.09	105.80
1	1G	449	C	C6-N1-C2	-6.77	117.59	120.30
26	1H	2444	G	C8-N9-C4	-6.77	103.69	106.40
26	14	457	A	C8-N9-C4	-6.77	103.09	105.80
1	13	690	G	C8-N9-C4	-6.77	103.69	106.40
26	14	2606	C	C6-N1-C2	6.77	123.01	120.30
23	2L	27	G	C4-C5-N7	6.77	113.51	110.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	13	449	C	C6-N1-C2	-6.76	117.59	120.30
1	13	652	U	C2-N1-C1'	6.76	125.82	117.70
26	1H	1972	A	N9-C4-C5	-6.76	103.09	105.80
1	1G	1281	U	C2-N1-C1'	6.76	125.82	117.70
26	1H	666	G	N1-C6-O6	6.76	123.95	119.90
26	1H	74	A	C6-C5-N7	-6.76	127.57	132.30
26	1H	226	G	O4'-C1'-N9	6.76	113.61	108.20
26	14	1899	G	N3-C4-N9	-6.76	121.95	126.00
26	14	2584	U	N1-C2-O2	6.76	127.53	122.80
26	1H	1379	A	C5-N7-C8	-6.75	100.52	103.90
26	1H	1950	G	C5-C6-O6	6.75	132.65	128.60
26	14	2283	C	N3-C4-N4	6.75	122.73	118.00
26	1H	2518	A	C4-C5-N7	6.75	114.08	110.70
26	1H	2062	A	N9-C4-C5	-6.75	103.10	105.80
26	1H	1950	G	N1-C6-O6	-6.75	115.85	119.90
26	1H	1210	A	C2-N3-C4	-6.74	107.23	110.60
27	1J	98	G	N3-C4-N9	6.74	130.05	126.00
26	1H	1678	G	N3-C4-C5	6.74	131.97	128.60
26	1H	1128	A	C8-N9-C4	-6.74	103.11	105.80
26	14	1011	G	O4'-C1'-N9	6.73	113.59	108.20
26	14	1603	A	C8-N9-C4	-6.73	103.11	105.80
26	14	1342	A	O4'-C1'-N9	6.73	113.58	108.20
1	1G	1112	C	C6-N1-C2	-6.73	117.61	120.30
26	14	582	G	C5-C6-O6	-6.73	124.56	128.60
1	13	1470	G	N1-C6-O6	6.73	123.94	119.90
26	14	1616	A	N1-C6-N6	6.73	122.64	118.60
26	1H	1990	C	C6-N1-C2	-6.72	117.61	120.30
26	1H	2330	G	N9-C4-C5	-6.72	102.71	105.40
26	14	1966	A	N1-C6-N6	6.72	122.63	118.60
26	1H	837	C	C5-C6-N1	6.72	124.36	121.00
26	1H	942	G	N3-C2-N2	-6.72	115.20	119.90
26	1H	263	C	N3-C2-O2	-6.72	117.20	121.90
26	1H	2377	A	C8-N9-C4	6.71	108.48	105.80
26	1H	1820	U	C5-C6-N1	-6.71	119.35	122.70
1	13	1206	G	C6-C5-N7	-6.70	126.38	130.40
26	1H	115	C	N3-C4-N4	6.70	122.69	118.00
26	1H	1972	A	C4-C5-N7	6.70	114.05	110.70
26	1H	330	A	N1-C2-N3	6.70	132.65	129.30
1	13	720	C	C6-N1-C2	-6.70	117.62	120.30
26	14	1276	A	N9-C4-C5	-6.70	103.12	105.80
1	13	1281	U	N1-C2-O2	6.70	127.49	122.80
26	14	659	C	C4-C5-C6	6.70	120.75	117.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	1G	529	G	C4-C5-N7	6.69	113.48	110.80
26	1H	2367	G	C6-C5-N7	-6.69	126.39	130.40
26	14	330	A	C4-C5-N7	6.69	114.05	110.70
26	14	574	C	N1-C2-O2	6.69	122.91	118.90
26	14	1528	A	N7-C8-N9	6.69	117.14	113.80
26	14	2433	A	N1-C2-N3	6.68	132.64	129.30
1	1G	1139	G	N3-C4-C5	6.68	131.94	128.60
1	13	758	G	N3-C4-C5	6.68	131.94	128.60
26	14	2447	G	N3-C4-N9	-6.67	122.00	126.00
26	1H	915	C	N1-C2-O2	6.67	122.90	118.90
26	1H	1647	G	O5'-P-OP1	-6.67	99.70	105.70
26	1H	1989	G	C5-C6-O6	-6.67	124.60	128.60
26	14	1373	A	C8-N9-C4	6.67	108.47	105.80
26	1H	2246	G	N3-C4-C5	-6.67	125.27	128.60
1	13	700	G	N1-C6-O6	6.66	123.90	119.90
26	14	1405	U	O5'-P-OP2	-6.66	99.70	105.70
1	13	1516	G	N3-C4-N9	-6.66	122.01	126.00
26	14	1210	A	N1-C6-N6	6.66	122.59	118.60
1	13	452	A	C8-N9-C4	6.65	108.46	105.80
1	13	186(A)	C	C6-N1-C2	-6.65	117.64	120.30
26	1H	82	G	C5-C6-N1	-6.65	108.17	111.50
26	14	574	C	N1-C2-N3	-6.65	114.54	119.20
26	1H	621	A	N1-C6-N6	6.65	122.59	118.60
26	1H	973	A	C2-N3-C4	-6.65	107.28	110.60
26	14	1774	C	C2-N1-C1'	6.65	126.11	118.80
26	14	737	C	N1-C2-O2	-6.64	114.91	118.90
26	1H	1763	G	O5'-P-OP2	-6.64	99.72	105.70
26	1H	800	A	N1-C2-N3	6.64	132.62	129.30
26	14	2012	G	N3-C4-N9	6.64	129.98	126.00
1	1G	288	A	N1-C6-N6	6.64	122.58	118.60
26	14	1471	A	C8-N9-C4	-6.63	103.15	105.80
1	13	766	A	N1-C6-N6	6.63	122.58	118.60
26	14	675	A	C5-N7-C8	-6.63	100.59	103.90
26	14	1963	U	N1-C2-O2	6.63	127.44	122.80
26	1H	2304	G	N3-C4-N9	-6.62	122.03	126.00
1	1G	518	C	N3-C2-O2	-6.62	117.26	121.90
26	1H	1612	C	N3-C4-C5	-6.62	119.25	121.90
1	1G	1128	C	C5-C6-N1	6.62	124.31	121.00
26	1H	205	G	N1-C6-O6	-6.62	115.93	119.90
1	13	1514	C	C6-N1-C2	-6.62	117.65	120.30
1	1G	527	G	N1-C6-O6	-6.62	115.93	119.90
26	14	700	G	N1-C6-O6	6.62	123.87	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	2713	A	C4-C5-N7	6.62	114.01	110.70
26	1H	2778	A	O5'-P-OP2	-6.61	99.75	105.70
1	1G	121	C	C2-N1-C1'	6.61	126.08	118.80
26	14	140	A	C6-C5-N7	-6.61	127.67	132.30
1	1G	995	C	C6-N1-C2	-6.61	117.66	120.30
26	1H	1971	A	C8-N9-C4	6.61	108.44	105.80
26	14	1613	G	N3-C4-N9	6.61	129.96	126.00
1	1G	529	G	N9-C4-C5	-6.61	102.76	105.40
26	1H	765	G	O5'-P-OP1	-6.60	99.76	105.70
26	1H	970	C	N1-C2-O2	-6.60	114.94	118.90
26	1H	1210	A	N3-C4-C5	6.60	131.42	126.80
1	1G	767	A	N1-C6-N6	6.60	122.56	118.60
26	1H	1616	A	C4-C5-N7	6.60	114.00	110.70
26	1H	49	A	C5-N7-C8	6.59	107.20	103.90
26	14	654(I)	C	C2-N1-C1'	6.59	126.05	118.80
26	1H	1784	A	C5-C6-N6	6.58	128.97	123.70
26	14	1142(A)	A	C2-N3-C4	-6.58	107.31	110.60
26	14	1989	G	N3-C4-N9	-6.58	122.05	126.00
26	1H	141	A	C4-C5-N7	6.58	113.99	110.70
1	1G	328	C	N1-C2-O2	6.58	122.85	118.90
26	1H	528	A	C5-C6-N1	-6.58	114.41	117.70
26	14	945	A	O4'-C1'-N9	6.58	113.46	108.20
26	1H	2037	G	C5-C6-N1	6.58	114.79	111.50
27	16	108	C	O4'-C1'-N1	6.58	113.46	108.20
26	14	1899	G	N1-C2-N3	6.58	127.84	123.90
26	14	1608	A	N1-C6-N6	-6.57	114.66	118.60
26	14	2443	C	N3-C4-N4	6.57	122.60	118.00
26	14	1608	A	O5'-P-OP1	-6.56	99.79	105.70
26	1H	1773	A	N1-C2-N3	6.56	132.58	129.30
1	1G	1354	C	C5-C6-N1	6.56	124.28	121.00
1	13	812	C	N1-C2-O2	6.56	122.83	118.90
1	13	1176	A	C8-N9-C4	-6.56	103.18	105.80
26	1H	2402	C	C6-N1-C2	-6.56	117.68	120.30
26	14	1564	C	C6-N1-C2	-6.56	117.68	120.30
1	13	1533	C	C6-N1-C2	-6.56	117.68	120.30
27	16	81	G	N7-C8-N9	6.55	116.38	113.10
23	2K	87	A	C8-N9-C4	6.55	108.42	105.80
26	14	312	G	O5'-P-OP1	-6.55	99.80	105.70
26	14	783	A	C6-C5-N7	-6.55	127.72	132.30
26	1H	621	A	N7-C8-N9	6.55	117.07	113.80
1	13	584	G	N1-C6-O6	-6.55	115.97	119.90
26	1H	2346	A	N7-C8-N9	6.55	117.07	113.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	13	894	G	C8-N9-C4	6.54	109.02	106.40
2	12	196	LEU	CA-CB-CG	6.54	130.34	115.30
26	14	1256	G	C5-C6-O6	-6.54	124.67	128.60
1	1G	495	A	N1-C6-N6	-6.54	114.68	118.60
1	13	1513	A	C8-N9-C4	6.54	108.41	105.80
26	1H	1378	A	C2-N3-C4	-6.54	107.33	110.60
26	1H	2518	A	N1-C6-N6	6.54	122.52	118.60
26	1H	1784	A	N1-C2-N3	6.53	132.57	129.30
26	14	1332	G	N7-C8-N9	6.53	116.37	113.10
26	14	1688	U	N1-C2-O2	-6.53	118.23	122.80
26	14	198	C	O5'-P-OP1	-6.53	99.82	105.70
26	1H	944	G	N7-C8-N9	6.53	116.36	113.10
23	2L	63	G	O4'-C1'-N9	6.53	113.42	108.20
26	1H	265	A	O4'-C1'-N9	6.53	113.42	108.20
23	2L	16	U	N1-C2-O2	6.53	127.37	122.80
26	14	933	A	C6-C5-N7	-6.53	127.73	132.30
26	1H	474	G	C8-N9-C4	-6.52	103.79	106.40
22	1L	87	A	C5-N7-C8	-6.52	100.64	103.90
27	16	74	U	C5-C4-O4	6.52	129.81	125.90
26	1H	1576	U	C6-N1-C2	-6.52	117.09	121.00
26	14	2387	U	C5-C6-N1	-6.52	119.44	122.70
26	1H	2310	A	N7-C8-N9	6.51	117.05	113.80
26	14	435	C	N1-C2-O2	6.51	122.81	118.90
26	1H	2593	U	N3-C4-C5	-6.51	110.70	114.60
26	1H	750	A	OP1-P-O3'	6.50	119.51	105.20
26	1H	1190	G	C4-C5-N7	6.50	113.40	110.80
26	1H	1376	C	N3-C4-C5	-6.50	119.30	121.90
26	14	1678	G	C5-N7-C8	-6.50	101.05	104.30
26	14	1950	G	C5-N7-C8	-6.50	101.05	104.30
26	1H	2527	C	C6-N1-C2	-6.50	117.70	120.30
1	13	874	G	N3-C4-C5	-6.50	125.35	128.60
26	14	533	G	C8-N9-C1'	-6.50	118.55	127.00
26	14	749	C	N1-C2-O2	6.50	122.80	118.90
26	14	2610	C	C6-N1-C2	6.50	122.90	120.30
26	1H	746	A	O4'-C1'-N9	6.50	113.40	108.20
26	1H	2699	C	C6-N1-C2	6.49	122.90	120.30
1	1G	46	G	N1-C6-O6	6.49	123.80	119.90
26	14	315	G	C8-N9-C4	6.49	109.00	106.40
26	1H	1781	C	N1-C2-O2	6.49	122.79	118.90
26	1H	613	U	N3-C2-O2	-6.49	117.66	122.20
26	1H	1021	A	N3-C4-C5	6.49	131.34	126.80
26	14	2072	G	N3-C2-N2	6.49	124.44	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	2526	G	N3-C4-N9	-6.49	122.11	126.00
26	14	723	G	O5'-P-OP1	6.49	118.48	110.70
1	1G	360	A	N1-C6-N6	6.48	122.49	118.60
26	1H	1559	G	C2-N3-C4	-6.48	108.66	111.90
26	14	459	U	N3-C2-O2	-6.48	117.66	122.20
1	1G	197	A	P-O3'-C3'	6.48	127.47	119.70
26	1H	330	A	O4'-C1'-N9	-6.47	103.02	108.20
26	1H	835	A	C5-C6-N6	-6.47	118.52	123.70
26	14	1204	A	C2-N3-C4	-6.47	107.36	110.60
26	1H	2468	G	C4-N9-C1'	6.47	134.91	126.50
26	1H	1698	A	O4'-C1'-N9	6.47	113.38	108.20
1	13	974	A	C4-C5-N7	6.47	113.93	110.70
26	14	1678	G	N3-C4-N9	-6.47	122.12	126.00
26	1H	265	A	N7-C8-N9	6.47	117.03	113.80
26	1H	2409	G	C6-C5-N7	-6.47	126.52	130.40
26	14	632	A	O5'-P-OP2	6.46	118.46	110.70
1	1G	121	C	N1-C2-O2	6.46	122.78	118.90
26	14	391	G	C4-N9-C1'	6.46	134.90	126.50
1	1G	353	A	N1-C6-N6	6.46	122.47	118.60
26	14	1544	C	N1-C2-O2	6.46	122.77	118.90
26	1H	186	G	N9-C4-C5	-6.45	102.82	105.40
26	1H	930	U	C5-C4-O4	6.45	129.77	125.90
26	14	1897	G	N3-C4-N9	6.45	129.87	126.00
26	1H	1021	A	N7-C8-N9	6.45	117.03	113.80
26	1H	2490	G	C4-C5-N7	6.45	113.38	110.80
26	14	533	G	C4-N9-C1'	6.45	134.88	126.50
26	14	2438	U	O5'-P-OP2	-6.45	99.90	105.70
26	1H	1520	U	N3-C2-O2	-6.45	117.69	122.20
1	1G	895	G	C6-C5-N7	-6.44	126.53	130.40
26	1H	684	G	N3-C4-C5	-6.44	125.38	128.60
26	14	1323	U	N3-C2-O2	6.44	126.71	122.20
26	1H	1320	C	C6-N1-C2	6.44	122.88	120.30
26	1H	1771	C	C5-C4-N4	-6.44	115.69	120.20
26	14	533	G	C6-C5-N7	-6.44	126.54	130.40
26	1H	2779	U	N3-C2-O2	-6.43	117.70	122.20
26	14	686	G	C4-C5-N7	6.43	113.37	110.80
26	14	945	A	N9-C4-C5	-6.43	103.23	105.80
26	14	1663	C	C6-N1-C2	6.43	122.87	120.30
26	1H	859	G	N3-C4-N9	-6.43	122.14	126.00
26	14	2287	A	N1-C2-N3	6.43	132.52	129.30
26	14	265	A	C4-C5-N7	6.43	113.92	110.70
26	1H	219	G	OP1-P-O3'	6.43	119.34	105.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	1437	C	C6-N1-C2	-6.43	117.73	120.30
26	14	2776	A	P-O3'-C3'	6.43	127.41	119.70
26	1H	1588	C	C6-N1-C2	-6.42	117.73	120.30
26	14	2581	G	N7-C8-N9	6.42	116.31	113.10
1	1G	1498	U	P-O3'-C3'	6.42	127.41	119.70
26	1H	2008	C	O5'-P-OP1	6.42	118.41	110.70
23	2L	8	U	C2-N1-C1'	6.42	125.40	117.70
1	13	739	C	N1-C2-O2	-6.42	115.05	118.90
26	1H	189	G	N1-C6-O6	6.42	123.75	119.90
26	1H	621	A	N1-C2-N3	6.42	132.51	129.30
1	1G	1322	C	C6-N1-C1'	-6.42	113.10	120.80
1	13	789	U	N3-C2-O2	-6.41	117.71	122.20
26	1H	271(B)	G	P-O3'-C3'	6.41	127.40	119.70
26	1H	915	C	N3-C2-O2	-6.41	117.41	121.90
26	1H	1391	U	N1-C2-O2	6.41	127.29	122.80
26	1H	266	G	C5-C6-O6	-6.41	124.75	128.60
26	1H	1678	G	N3-C4-N9	-6.41	122.15	126.00
26	1H	2326	C	C6-N1-C2	-6.41	117.74	120.30
26	14	676	A	C6-C5-N7	-6.41	127.81	132.30
26	14	1313	U	C2-N1-C1'	6.41	125.39	117.70
26	1H	192	C	N3-C4-C5	6.41	124.46	121.90
26	1H	2331	G	C2-N3-C4	-6.41	108.70	111.90
1	1G	180	U	C6-N1-C2	-6.41	117.16	121.00
26	1H	213	A	C5-C6-N6	-6.40	118.58	123.70
26	14	613	U	N1-C2-O2	6.40	127.28	122.80
26	1H	2655	G	O4'-C1'-N9	6.40	113.32	108.20
1	13	1305	G	N7-C8-N9	6.40	116.30	113.10
26	14	1991	U	C5-C4-O4	6.40	129.74	125.90
26	14	1547	C	C6-N1-C2	-6.39	117.74	120.30
1	1G	774	G	N1-C6-O6	6.39	123.74	119.90
26	14	71	A	N3-C4-C5	6.39	131.28	126.80
26	14	974(A)	C	N3-C2-O2	-6.39	117.43	121.90
26	14	2632	A	C8-N9-C4	6.39	108.36	105.80
23	2K	20	U	P-O3'-C3'	6.39	127.36	119.70
26	1H	330	A	C5-N7-C8	-6.38	100.71	103.90
27	16	100	G	C8-N9-C4	6.38	108.95	106.40
1	13	1530	G	C4-C5-N7	6.38	113.35	110.80
26	1H	1846	G	C5-C6-O6	-6.38	124.77	128.60
26	1H	1972	A	C6-C5-N7	-6.38	127.84	132.30
26	1H	2439	A	N7-C8-N9	6.38	116.99	113.80
26	1H	2509	G	N3-C4-N9	6.38	129.83	126.00
26	1H	1616	A	O4'-C1'-N9	6.38	113.30	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	13	578	C	C6-N1-C2	-6.38	117.75	120.30
26	1H	2346	A	C6-C5-N7	-6.37	127.84	132.30
26	14	1695	G	C4-C5-C6	6.37	122.62	118.80
26	14	2509	G	C6-C5-N7	-6.37	126.58	130.40
26	1H	2688	U	N1-C2-N3	6.37	118.72	114.90
26	1H	2824	C	C6-N1-C2	6.37	122.85	120.30
1	13	585	G	C8-N9-C4	6.37	108.95	106.40
1	1G	87	A	N1-C6-N6	6.37	122.42	118.60
26	1H	1496	A	C8-N9-C4	-6.37	103.25	105.80
26	14	1820	U	C5-C6-N1	-6.37	119.52	122.70
26	1H	1964	G	O5'-P-OP2	-6.36	99.97	105.70
26	14	1210	A	C5-N7-C8	-6.36	100.72	103.90
26	14	1970	A	N1-C6-N6	6.36	122.42	118.60
26	1H	1805	U	N3-C2-O2	-6.36	117.75	122.20
26	1H	2584	U	C4-C5-C6	6.36	123.52	119.70
27	16	79	C	C5-C6-N1	6.36	124.18	121.00
1	1G	150	C	C6-N1-C2	-6.36	117.76	120.30
26	14	2723	C	C6-N1-C2	-6.36	117.75	120.30
1	13	1362	C	N1-C2-O2	6.36	122.71	118.90
1	13	1502	A	N7-C8-N9	6.36	116.98	113.80
24	3L	59	U	P-O3'-C3'	6.36	127.33	119.70
26	14	2068	U	OP1-P-O3'	6.36	119.18	105.20
26	1H	265	A	N1-C6-N6	6.35	122.41	118.60
26	14	1917	U	C5-C6-N1	6.35	125.88	122.70
26	1H	600	G	C8-N9-C4	6.35	108.94	106.40
26	1H	1606	G	N1-C6-O6	6.35	123.71	119.90
26	1H	2593	U	N1-C2-O2	-6.35	118.36	122.80
26	1H	2346	A	C5-C6-N1	-6.34	114.53	117.70
27	16	116	G	N1-C6-O6	6.34	123.71	119.90
26	14	1298	C	O5'-P-OP2	-6.34	99.99	105.70
1	13	169	C	C6-N1-C2	-6.34	117.76	120.30
26	14	800	A	N1-C6-N6	-6.34	114.80	118.60
26	14	1629	U	N3-C4-C5	-6.34	110.80	114.60
26	1H	215	G	C5-C6-O6	-6.34	124.80	128.60
26	1H	678	C	N1-C2-O2	-6.34	115.10	118.90
1	13	1514	C	N3-C4-C5	-6.34	119.37	121.90
19	AI	30	LEU	CA-CB-CG	6.33	129.87	115.30
26	14	2688	U	C5-C6-N1	-6.33	119.53	122.70
26	1H	1798	U	C5-C4-O4	-6.33	122.10	125.90
26	1H	2771	C	C6-N1-C2	-6.33	117.77	120.30
26	1H	1363	C	C6-N1-C2	6.33	122.83	120.30
26	14	1022	G	P-O3'-C3'	6.33	127.29	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	1G	328	C	C2-N1-C1'	6.33	125.76	118.80
26	1H	801	G	O5'-P-OP2	-6.32	100.01	105.70
26	1H	1379	A	C4-C5-N7	6.32	113.86	110.70
26	1H	2504	U	N1-C2-O2	6.32	127.22	122.80
26	14	1527	G	N3-C4-C5	6.32	131.76	128.60
1	13	1125	U	P-O3'-C3'	6.32	127.28	119.70
1	13	8	A	C8-N9-C4	6.31	108.33	105.80
26	14	1681	G	C4-C5-N7	6.31	113.32	110.80
26	14	752	A	P-O3'-C3'	6.30	127.26	119.70
26	1H	1535	U	C2-N1-C1'	6.30	125.26	117.70
26	1H	667	U	N1-C2-O2	-6.30	118.39	122.80
26	1H	1675	C	OP1-P-OP2	6.30	129.04	119.60
26	1H	2674	G	N3-C4-N9	-6.29	122.22	126.00
26	1H	1950	G	O4'-C1'-N9	6.29	113.23	108.20
26	1H	2595	G	C5-N7-C8	-6.29	101.15	104.30
26	14	2027	G	C6-N1-C2	-6.29	121.33	125.10
27	1J	114	G	C4-N9-C1'	-6.29	118.32	126.50
26	1H	1626	G	C8-N9-C4	-6.29	103.89	106.40
26	14	2012	G	N9-C4-C5	-6.29	102.89	105.40
26	1H	1833	U	O5'-P-OP2	-6.29	100.04	105.70
26	1H	1021	A	N1-C6-N6	6.28	122.37	118.60
26	1H	446	G	N9-C4-C5	-6.28	102.89	105.40
26	1H	2268	A	N1-C6-N6	6.28	122.37	118.60
1	1G	898	G	N3-C4-C5	6.28	131.74	128.60
26	14	623	G	C5-C6-O6	-6.28	124.83	128.60
26	14	664	C	C4-C5-C6	6.28	120.54	117.40
26	14	1022	G	N9-C4-C5	6.28	107.91	105.40
26	14	1528	A	C8-N9-C4	-6.28	103.29	105.80
26	1H	677	A	N1-C6-N6	-6.27	114.84	118.60
26	1H	1931	U	C5-C4-O4	6.27	129.66	125.90
26	1H	2295	C	C5-C6-N1	6.27	124.14	121.00
26	1H	1396	U	N3-C2-O2	-6.27	117.81	122.20
26	1H	1817	G	C5-C6-O6	6.26	132.36	128.60
1	1G	320	C	C6-N1-C2	6.26	122.81	120.30
26	1H	1674	G	C6-C5-N7	-6.26	126.64	130.40
26	1H	2580	U	C5-C6-N1	6.26	125.83	122.70
26	1H	1940	U	N1-C2-O2	-6.26	118.42	122.80
26	14	1437	C	C5-C6-N1	6.26	124.13	121.00
26	1H	1786	A	N9-C1'-C2'	6.25	122.13	114.00
26	1H	197	A	N1-C2-N3	6.25	132.43	129.30
26	14	1999	C	C6-N1-C2	6.25	122.80	120.30
26	14	570	G	C8-N9-C4	-6.25	103.90	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	1256	G	N1-C6-O6	6.25	123.65	119.90
1	13	1413	A	N1-C6-N6	6.25	122.35	118.60
26	1H	33	U	OP1-P-O3'	6.25	118.94	105.20
26	1H	783	A	N3-C4-C5	6.25	131.17	126.80
1	1G	774	G	N9-C4-C5	-6.24	102.90	105.40
26	14	751	A	N1-C2-N3	6.24	132.42	129.30
26	1H	265	A	C6-C5-N7	-6.24	127.93	132.30
26	14	1790	C	C2-N3-C4	-6.24	116.78	119.90
26	1H	736	C	N3-C2-O2	6.24	126.27	121.90
1	13	1528	U	C5-C6-N1	-6.24	119.58	122.70
1	1G	275	G	N3-C4-N9	6.23	129.74	126.00
26	14	265	A	O4'-C1'-N9	6.23	113.19	108.20
26	14	518	G	N1-C2-N2	-6.23	110.59	116.20
26	14	1328	G	N1-C6-O6	6.23	123.64	119.90
1	13	1065	U	P-O3'-C3'	6.23	127.17	119.70
23	2K	20	U	N1-C2-O2	6.23	127.16	122.80
26	1H	2565	A	N9-C4-C5	-6.23	103.31	105.80
26	14	2726	U	C5-C4-O4	6.23	129.64	125.90
26	1H	128	C	C6-N1-C2	6.22	122.79	120.30
26	1H	621	A	O4'-C1'-N9	6.22	113.18	108.20
26	1H	1636	C	N3-C4-N4	6.22	122.36	118.00
26	1H	622	G	N3-C4-N9	6.22	129.73	126.00
26	1H	194	G	C8-N9-C4	6.22	108.89	106.40
26	14	2426	A	N7-C8-N9	6.22	116.91	113.80
1	1G	529	G	N3-C4-N9	6.22	129.73	126.00
26	1H	593	G	N1-C2-N3	6.21	127.63	123.90
26	14	2329	G	N9-C4-C5	-6.21	102.91	105.40
1	13	1158	C	N3-C2-O2	-6.21	117.55	121.90
26	1H	101	G	C8-N9-C4	6.21	108.88	106.40
26	1H	1790	C	C6-N1-C2	6.21	122.78	120.30
26	14	763	G	C8-N9-C4	-6.21	103.92	106.40
1	13	131	C	N1-C2-O2	6.20	122.62	118.90
26	1H	2424	C	N1-C2-O2	6.20	122.62	118.90
26	14	703	U	C5-C4-O4	6.20	129.62	125.90
26	1H	530	G	N1-C2-N2	-6.20	110.62	116.20
26	1H	2346	A	C4-C5-C6	6.20	120.10	117.00
26	1H	793	A	C2-N3-C4	-6.20	107.50	110.60
26	1H	829	A	C2-N3-C4	-6.20	107.50	110.60
26	1H	1426	G	C8-N9-C4	-6.20	103.92	106.40
26	14	1254	A	C6-N1-C2	-6.20	114.88	118.60
26	14	2829	C	C6-N1-C2	6.20	122.78	120.30
26	1H	115	C	C5-C4-N4	-6.19	115.86	120.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	275	G	C6-C5-N7	-6.19	126.68	130.40
26	1H	1699	G	O5'-P-OP1	-6.19	100.13	105.70
26	14	55	G	C5-C6-O6	-6.19	124.89	128.60
26	14	275	G	C5-C6-O6	-6.19	124.89	128.60
26	14	2879	C	C6-N1-C2	6.19	122.78	120.30
27	16	12	C	C4-C5-C6	6.19	120.49	117.40
26	14	1348	G	O5'-P-OP2	6.19	118.12	110.70
26	14	113	G	N1-C6-O6	6.19	123.61	119.90
34	58	120	LEU	CA-CB-CG	6.18	129.53	115.30
38	98	18	LEU	CA-CB-CG	6.18	129.52	115.30
26	14	469	G	N1-C6-O6	-6.18	116.19	119.90
26	14	1496	A	C4-C5-N7	6.18	113.79	110.70
1	13	428	G	N3-C4-N9	-6.18	122.29	126.00
26	1H	808	G	N1-C6-O6	-6.18	116.19	119.90
1	1G	1300	G	P-O3'-C3'	6.18	127.12	119.70
1	1G	1108	G	C5-C6-O6	6.18	132.31	128.60
26	14	2392	A	C6-N1-C2	6.18	122.31	118.60
26	1H	1616	A	C8-N9-C4	-6.18	103.33	105.80
26	14	265	A	N1-C6-N6	6.18	122.31	118.60
26	14	1762	A	C2-N3-C4	-6.18	107.51	110.60
1	13	880	C	C6-N1-C2	6.18	122.77	120.30
26	1H	461	C	N3-C2-O2	6.18	126.22	121.90
26	14	1807	G	N1-C6-O6	6.17	123.61	119.90
26	14	2779	U	N1-C2-O2	6.17	127.12	122.80
26	1H	1825	A	N1-C2-N3	6.17	132.39	129.30
26	1H	1535	U	N1-C2-O2	6.17	127.12	122.80
26	1H	440	G	N1-C6-O6	6.17	123.60	119.90
26	1H	945	A	C5-C6-N1	-6.17	114.62	117.70
26	1H	2007	C	N3-C2-O2	-6.17	117.58	121.90
26	14	2598	A	OP2-P-O3'	6.17	118.77	105.20
26	1H	1985	G	N1-C6-O6	-6.17	116.20	119.90
1	1G	1354	C	C6-N1-C2	-6.17	117.83	120.30
23	2K	3	C	C6-N1-C2	-6.17	117.83	120.30
1	1G	1278	U	C6-N1-C1'	-6.17	112.57	121.20
26	1H	140	A	C8-N9-C4	-6.16	103.33	105.80
26	14	974(A)	C	N3-C4-C5	-6.16	119.44	121.90
26	1H	2599	G	N3-C2-N2	-6.16	115.59	119.90
26	1H	734	A	C2-N3-C4	-6.16	107.52	110.60
26	1H	2501	C	N3-C4-C5	6.16	124.36	121.90
26	14	465	G	N9-C4-C5	6.16	107.86	105.40
26	1H	265	A	C2-N3-C4	-6.16	107.52	110.60
26	1H	1363	C	C2-N3-C4	-6.16	116.82	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	1L	87	A	N7-C8-N9	6.16	116.88	113.80
26	14	1682	G	N1-C6-O6	6.15	123.59	119.90
1	13	856	C	C6-N1-C2	-6.15	117.84	120.30
26	14	2726	U	N3-C4-O4	-6.15	115.09	119.40
27	1J	98	G	C8-N9-C1'	-6.15	119.00	127.00
26	14	2329	G	N3-C4-N9	6.15	129.69	126.00
26	1H	924	C	C6-N1-C2	6.15	122.76	120.30
26	1H	2362	G	C8-N9-C4	6.15	108.86	106.40
23	2K	72	C	C6-N1-C2	-6.15	117.84	120.30
26	1H	1604	C	N3-C4-N4	6.15	122.30	118.00
26	1H	74	A	O4'-C1'-N9	-6.14	103.28	108.20
26	1H	217	G	C4-C5-N7	-6.14	108.34	110.80
26	1H	1337	G	OP1-P-O3'	6.14	118.71	105.20
26	1H	571	A	C8-N9-C4	6.14	108.26	105.80
26	1H	1698	A	C4-C5-C6	6.14	120.07	117.00
26	14	84	A	C8-N9-C4	6.14	108.26	105.80
26	14	1395	A	C8-N9-C4	6.14	108.26	105.80
26	14	2683	C	C6-N1-C2	-6.14	117.84	120.30
26	1H	786	C	C5-C6-N1	-6.14	117.93	121.00
26	1H	1613	G	N1-C2-N2	-6.14	110.67	116.20
26	1H	2449	U	N3-C4-C5	-6.14	110.92	114.60
26	1H	659	C	C5-C6-N1	-6.14	117.93	121.00
26	1H	2562	U	C5-C6-N1	-6.14	119.63	122.70
27	16	27	C	N3-C4-C5	-6.14	119.45	121.90
1	13	811	C	C5-C6-N1	-6.13	117.93	121.00
26	1H	580	C	C6-N1-C2	-6.13	117.85	120.30
1	1G	1502	A	C5-N7-C8	-6.13	100.83	103.90
26	14	945	A	C5-C6-N1	-6.13	114.63	117.70
26	1H	2230	G	O5'-P-OP2	-6.13	100.18	105.70
26	1H	602	G	C6-C5-N7	-6.13	126.72	130.40
1	1G	1346	A	OP2-P-O3'	6.13	118.69	105.20
26	14	1276	A	C6-C5-N7	-6.13	128.01	132.30
1	13	23	C	N3-C4-C5	-6.13	119.45	121.90
1	13	974	A	C6-C5-N7	-6.13	128.01	132.30
26	1H	801	G	C4-C5-N7	-6.13	108.35	110.80
26	1H	1654	A	C8-N9-C4	-6.13	103.35	105.80
26	1H	1779	U	N1-C2-N3	-6.13	111.22	114.90
26	14	2779	U	C2-N1-C1'	6.13	125.05	117.70
1	1G	1139	G	C4-N9-C1'	-6.12	118.54	126.50
1	13	883	C	C6-N1-C2	-6.12	117.85	120.30
26	14	212	G	O5'-P-OP2	-6.12	100.19	105.70
26	14	1786	A	N9-C1'-C2'	6.12	121.96	114.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	2018	G	C8-N9-C4	-6.12	103.95	106.40
26	1H	2392	A	C4-C5-N7	6.12	113.76	110.70
26	14	2001	A	N1-C6-N6	6.12	122.27	118.60
27	1J	30	C	C6-N1-C2	-6.12	117.85	120.30
26	1H	2710	C	C4-C5-C6	6.12	120.46	117.40
26	1H	2383	G	N3-C4-N9	6.12	129.67	126.00
26	1H	2465	C	C6-N1-C2	6.12	122.75	120.30
23	2L	8	U	C5-C6-N1	6.12	125.76	122.70
26	14	1632	A	N1-C6-N6	6.12	122.27	118.60
26	1H	70	G	N3-C4-C5	-6.12	125.54	128.60
26	1H	770	G	C4-C5-N7	6.12	113.25	110.80
26	1H	2566	A	P-O3'-C3'	6.12	127.04	119.70
26	14	2404	C	N3-C4-C5	6.12	124.35	121.90
1	1G	721	G	C6-C5-N7	-6.11	126.73	130.40
1	13	576	G	N1-C6-O6	6.11	123.57	119.90
24	3K	2	C	C6-N1-C2	-6.11	117.86	120.30
26	1H	203	C	O5'-P-OP2	6.11	118.03	110.70
26	1H	816	C	N3-C2-O2	6.11	126.18	121.90
26	1H	247	G	N9-C4-C5	-6.11	102.96	105.40
26	14	1225	C	C6-N1-C2	6.11	122.74	120.30
1	13	690	G	C4-C5-C6	6.11	122.46	118.80
26	1H	1066	U	C2-N1-C1'	6.11	125.03	117.70
26	1H	1636	C	N1-C2-O2	-6.11	115.24	118.90
26	14	2477	C	N3-C2-O2	-6.11	117.63	121.90
26	1H	847	U	N1-C2-N3	6.10	118.56	114.90
26	1H	2247	A	C2-N3-C4	-6.10	107.55	110.60
26	1H	2378	A	C8-N9-C4	6.10	108.24	105.80
1	1G	536	C	N1-C2-O2	6.10	122.56	118.90
1	1G	1498	U	O4'-C1'-N1	-6.10	103.32	108.20
27	16	8	U	O5'-P-OP1	6.10	118.02	110.70
26	14	1318	C	C6-N1-C2	-6.10	117.86	120.30
26	14	2775	A	N1-C6-N6	6.10	122.26	118.60
1	13	900	A	C8-N9-C4	6.10	108.24	105.80
26	14	2702	U	C2-N1-C1'	6.10	125.02	117.70
26	1H	49	A	N7-C8-N9	-6.09	110.75	113.80
26	1H	1535	U	N3-C2-O2	-6.09	117.94	122.20
26	14	1614	A	O4'-C1'-N9	6.09	113.07	108.20
26	1H	108	U	O5'-P-OP1	-6.09	100.22	105.70
1	1G	372	C	N1-C2-O2	6.09	122.55	118.90
26	1H	186	G	C4-C5-N7	6.09	113.23	110.80
26	1H	1781	C	N3-C2-O2	-6.09	117.64	121.90
26	1H	2346	A	C4-N9-C1'	6.09	137.25	126.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	747	U	C6-N1-C2	6.09	124.65	121.00
1	13	1305	G	N9-C4-C5	6.08	107.83	105.40
1	13	1356	G	C8-N9-C4	-6.08	103.97	106.40
26	1H	2466	C	C6-N1-C2	6.08	122.73	120.30
26	14	2423	U	C6-N1-C2	6.08	124.65	121.00
26	1H	2711	A	C2-N3-C4	-6.08	107.56	110.60
26	1H	208	C	N3-C4-C5	6.08	124.33	121.90
26	1H	379	G	N3-C4-N9	6.08	129.65	126.00
26	1H	1917	U	C5-C6-N1	6.08	125.74	122.70
26	14	1422	G	N1-C6-O6	6.08	123.55	119.90
26	14	2832	U	C2-N1-C1'	-6.08	110.41	117.70
1	13	1206	G	C4-N9-C1'	6.08	134.40	126.50
26	1H	781	A	N1-C6-N6	-6.08	114.95	118.60
26	1H	1321	A	C8-N9-C4	6.08	108.23	105.80
26	14	681	G	N1-C2-N3	6.08	127.55	123.90
1	1G	652	U	O5'-P-OP1	-6.07	100.23	105.70
26	14	278	A	P-O3'-C3'	6.07	126.99	119.70
26	1H	34	C	O5'-P-OP1	-6.07	100.23	105.70
26	14	2019	A	C8-N9-C4	6.07	108.23	105.80
1	1G	576	G	C6-C5-N7	-6.07	126.76	130.40
26	14	664	C	C2-N3-C4	-6.07	116.86	119.90
26	14	675	A	N1-C6-N6	6.07	122.24	118.60
26	14	211	A	N1-C6-N6	6.07	122.24	118.60
26	14	2634	G	N1-C6-O6	6.07	123.54	119.90
26	1H	1950	G	C4-N9-C1'	6.06	134.38	126.50
26	1H	2507	C	C6-N1-C2	-6.06	117.88	120.30
1	1G	14	U	C5-C6-N1	6.06	125.73	122.70
26	14	567	A	C6-N1-C2	-6.06	114.96	118.60
1	13	792	A	C3'-C2'-C1'	-6.06	96.65	101.50
1	1G	1259	C	C6-N1-C2	-6.06	117.88	120.30
26	14	1786	A	C4-N9-C1'	6.06	137.20	126.30
26	14	567	A	N1-C2-N3	6.05	132.33	129.30
26	14	1342	A	N1-C2-N3	6.05	132.33	129.30
26	1H	194	G	C5-C6-N1	6.05	114.53	111.50
26	1H	394	A	N1-C6-N6	-6.05	114.97	118.60
26	1H	2761	G	C8-N9-C4	6.05	108.82	106.40
1	1G	1374	A	C2-N3-C4	-6.05	107.58	110.60
26	14	2387	U	C6-N1-C2	6.05	124.63	121.00
26	14	1572	A	C5-C6-N6	-6.05	118.86	123.70
26	14	715	G	C5-C6-O6	-6.05	124.97	128.60
26	14	1323	U	N1-C2-O2	-6.05	118.57	122.80
1	13	413	G	C8-N9-C4	-6.05	103.98	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	2490	G	C5-N7-C8	-6.05	101.28	104.30
26	1H	1639	U	N3-C2-O2	-6.04	117.97	122.20
26	1H	251	A	O5'-P-OP1	-6.04	100.26	105.70
26	1H	298	G	N1-C6-O6	6.04	123.53	119.90
26	1H	74	A	C4-C5-C6	6.04	120.02	117.00
26	14	505	A	C8-N9-C4	-6.04	103.38	105.80
26	14	2688	U	N1-C2-N3	6.04	118.53	114.90
26	1H	807	U	C5-C4-O4	-6.04	122.28	125.90
1	13	481	G	N1-C6-O6	6.04	123.52	119.90
1	13	898	G	C5-C6-O6	-6.04	124.98	128.60
1	13	974	A	C5-N7-C8	-6.04	100.88	103.90
26	1H	513	A	C8-N9-C4	-6.04	103.39	105.80
26	1H	1829	A	C6-N1-C2	-6.03	114.98	118.60
24	3L	57	C	N1-C2-O2	6.03	122.52	118.90
26	1H	107	C	N3-C2-O2	6.03	126.12	121.90
26	1H	1265	A	O5'-P-OP1	-6.03	100.27	105.70
1	13	5	U	P-O3'-C3'	6.03	126.93	119.70
26	1H	571	A	N9-C4-C5	-6.03	103.39	105.80
26	1H	590	A	C8-N9-C4	-6.03	103.39	105.80
26	14	1327	C	C2-N1-C1'	-6.03	112.17	118.80
26	1H	2544	G	N1-C6-O6	6.03	123.52	119.90
26	14	113	G	C6-C5-N7	-6.03	126.78	130.40
26	14	723	G	N1-C6-O6	6.03	123.52	119.90
26	14	1651	G	C6-C5-N7	-6.03	126.78	130.40
26	1H	2032	G	C4-C5-C6	6.02	122.41	118.80
26	1H	2269	A	C8-N9-C4	6.02	108.21	105.80
24	3K	87	A	O5'-P-OP1	-6.02	100.28	105.70
26	1H	1273	U	P-O3'-C3'	6.02	126.92	119.70
26	1H	1345	C	N1-C2-O2	-6.02	115.29	118.90
1	1G	1200	C	N1-C2-O2	6.02	122.51	118.90
26	14	623	G	N1-C6-O6	6.02	123.51	119.90
26	14	2092	U	C5-C4-O4	6.02	129.51	125.90
1	13	1227	A	C8-N9-C4	-6.02	103.39	105.80
26	1H	226	G	N1-C2-N2	-6.02	110.78	116.20
1	1G	898	G	N7-C8-N9	-6.02	110.09	113.10
26	14	1193	G	C4-C5-N7	6.02	113.21	110.80
26	14	830	G	C8-N9-C4	6.02	108.81	106.40
26	14	1289	C	N3-C4-C5	-6.02	119.49	121.90
1	13	1530	G	N9-C4-C5	-6.01	102.99	105.40
26	1H	2501	C	C6-N1-C2	6.01	122.71	120.30
26	1H	812	C	N3-C2-O2	6.01	126.11	121.90
1	1G	121	C	C6-N1-C1'	-6.01	113.58	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	676	A	C6-C5-N7	-6.01	128.09	132.30
26	14	2506	U	C2-N3-C4	6.01	130.61	127.00
26	1H	116	C	N3-C4-C5	-6.01	119.50	121.90
26	1H	1899	G	C4-N9-C1'	-6.01	118.69	126.50
26	1H	2499	C	C6-N1-C2	-6.01	117.90	120.30
1	13	1220	G	N1-C6-O6	6.00	123.50	119.90
26	1H	225	A	C8-N9-C4	6.00	108.20	105.80
1	13	1441	G	N3-C4-N9	-6.00	122.40	126.00
26	1H	912	C	C6-N1-C2	-6.00	117.90	120.30
26	1H	1830	C	N3-C2-O2	6.00	126.10	121.90
27	16	45	A	N7-C8-N9	6.00	116.80	113.80
1	13	656	C	C5-C6-N1	6.00	124.00	121.00
26	1H	1382	G	C6-C5-N7	-6.00	126.80	130.40
1	1G	1449	C	C2-N1-C1'	6.00	125.40	118.80
26	14	669	G	N3-C4-C5	-6.00	125.60	128.60
26	14	921	G	N3-C4-N9	-6.00	122.40	126.00
36	78	50	ARG	NE-CZ-NH1	-6.00	117.30	120.30
48	F5	36	GLY	N-CA-C	6.00	128.10	113.10
1	13	757	U	N1-C2-O2	6.00	127.00	122.80
1	13	1281	U	C2-N1-C1'	6.00	124.90	117.70
26	14	1257	C	N1-C2-O2	-6.00	115.30	118.90
26	14	1613	G	N3-C4-C5	-6.00	125.60	128.60
26	14	1889	A	N1-C6-N6	-6.00	115.00	118.60
1	13	1227	A	N7-C8-N9	6.00	116.80	113.80
26	1H	958	U	C6-N1-C2	-6.00	117.40	121.00
26	14	2060	A	C8-N9-C4	-6.00	103.40	105.80
26	1H	131	G	C8-N9-C4	5.99	108.80	106.40
26	1H	2726	U	C5-C4-O4	5.99	129.50	125.90
26	14	2066	C	C6-N1-C2	-5.99	117.90	120.30
1	1G	1059	C	C6-N1-C2	-5.99	117.90	120.30
22	1L	85	C	C2-N1-C1'	-5.99	112.21	118.80
1	13	700	G	C5-C6-O6	-5.99	125.01	128.60
24	3K	46	G	O4'-C1'-N9	5.99	112.99	108.20
37	88	2	LEU	CA-CB-CG	5.99	129.07	115.30
26	14	1950	G	N3-C2-N2	5.99	124.09	119.90
33	69	9	LEU	CA-CB-CG	5.99	129.07	115.30
26	1H	1669	A	N7-C8-N9	5.98	116.79	113.80
26	14	447	A	C2-N3-C4	-5.98	107.61	110.60
26	14	808	G	OP1-P-OP2	5.98	128.57	119.60
1	13	1206	G	N7-C8-N9	5.98	116.09	113.10
26	1H	793	A	N1-C6-N6	5.98	122.19	118.60
26	14	2575	C	C2-N3-C4	5.98	122.89	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	13	534	U	C5-C4-O4	5.98	129.49	125.90
1	13	690	G	C2-N3-C4	-5.98	108.91	111.90
26	1H	2465	C	C5-C6-N1	-5.98	118.01	121.00
26	14	1616	A	N7-C8-N9	5.98	116.79	113.80
1	13	1504	G	P-O3'-C3'	5.98	126.87	119.70
26	1H	751	A	OP1-P-OP2	-5.98	110.63	119.60
27	16	44	G	C4-N9-C1'	-5.98	118.73	126.50
1	1G	390	C	C6-N1-C2	5.98	122.69	120.30
26	14	918	A	C8-N9-C4	-5.97	103.41	105.80
26	1H	613	U	C5-C4-O4	5.97	129.48	125.90
26	1H	1606	G	N9-C4-C5	-5.97	103.01	105.40
26	1H	2324	C	C5-C6-N1	-5.97	118.02	121.00
26	14	2278	A	N1-C2-N3	5.97	132.28	129.30
26	1H	747	U	O5'-P-OP1	-5.96	100.33	105.70
26	1H	2713	A	N7-C8-N9	5.96	116.78	113.80
26	14	1612	C	C6-N1-C2	5.96	122.69	120.30
26	14	1688	U	C2-N1-C1'	-5.96	110.55	117.70
26	14	2508	G	C4-C5-N7	5.96	113.19	110.80
26	1H	2430	A	C4-C5-C6	5.96	119.98	117.00
1	13	517	G	N1-C6-O6	5.96	123.47	119.90
1	13	748	C	C5-C6-N1	5.96	123.98	121.00
26	14	1596	A	N1-C6-N6	-5.96	115.03	118.60
26	1H	2346	A	C8-N9-C4	-5.96	103.42	105.80
1	13	504	C	C6-N1-C2	-5.95	117.92	120.30
26	14	1666	G	O4'-C1'-N9	5.95	112.96	108.20
26	1H	1382	G	N1-C6-O6	5.95	123.47	119.90
26	1H	2502	G	O5'-P-OP2	5.95	117.84	110.70
1	1G	353	A	C4-C5-N7	5.95	113.67	110.70
26	14	211	A	C2-N3-C4	-5.95	107.62	110.60
26	14	2506	U	C6-N1-C2	-5.95	117.43	121.00
1	13	1497	G	N1-C6-O6	-5.95	116.33	119.90
1	1G	1519	A	C5-C6-N6	5.95	128.46	123.70
22	1L	85	C	N3-C4-N4	-5.95	113.84	118.00
26	1H	1963	U	N1-C2-O2	5.95	126.96	122.80
26	14	855	G	C8-N9-C4	-5.95	104.02	106.40
26	14	1914	C	C2-N1-C1'	5.95	125.34	118.80
26	14	2392	A	C2-N3-C4	-5.95	107.63	110.60
26	1H	2388	A	C8-N9-C4	5.95	108.18	105.80
26	14	1653	G	N3-C4-C5	-5.95	125.63	128.60
26	14	1815	A	C6-N1-C2	-5.95	115.03	118.60
26	14	1352	U	O5'-P-OP2	-5.94	100.35	105.70
26	1H	460	A	N1-C6-N6	5.94	122.17	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	1G	974	A	O4'-C1'-N9	5.94	112.95	108.20
26	1H	1308	A	N1-C6-N6	-5.94	115.03	118.60
1	1G	258	G	N1-C6-O6	5.94	123.47	119.90
26	14	577	G	C4-C5-C6	5.94	122.36	118.80
26	1H	2869	G	C8-N9-C4	-5.94	104.03	106.40
26	14	1022	G	C8-N9-C4	-5.94	104.03	106.40
26	14	2278	A	C6-N1-C2	-5.94	115.04	118.60
26	14	439	G	C6-C5-N7	-5.94	126.84	130.40
1	13	690	G	C5-C6-O6	-5.93	125.04	128.60
26	14	1820	U	C6-N1-C2	5.93	124.56	121.00
26	1H	845	G	N3-C4-C5	5.93	131.57	128.60
26	1H	2264	C	C6-N1-C2	-5.93	117.93	120.30
26	1H	2449	U	C6-N1-C2	-5.93	117.44	121.00
26	1H	1157	G	N1-C2-N3	5.93	127.46	123.90
46	H8	117	LEU	CA-CB-CG	5.93	128.94	115.30
26	1H	2856	C	C2-N1-C1'	5.93	125.32	118.80
26	14	137	C	C6-N1-C2	-5.93	117.93	120.30
26	14	929	G	N7-C8-N9	5.93	116.06	113.10
26	14	630	G	O5'-P-OP2	-5.93	100.36	105.70
1	13	789	U	C5-C4-O4	5.93	129.46	125.90
26	1H	186	G	C6-C5-N7	-5.93	126.84	130.40
26	1H	1809	A	C5-C6-N6	-5.93	118.96	123.70
37	88	79	LEU	CA-CB-CG	-5.93	101.67	115.30
26	14	2689	U	N3-C4-O4	-5.93	115.25	119.40
27	1J	103	U	C5-C6-N1	-5.93	119.74	122.70
26	1H	309	G	N3-C4-C5	-5.92	125.64	128.60
26	1H	1931	U	N1-C2-O2	5.92	126.95	122.80
45	G8	81	LYS	N-CA-C	-5.92	95.00	111.00
26	14	1432	C	N3-C2-O2	5.92	126.05	121.90
26	1H	309	G	N3-C4-N9	5.92	129.55	126.00
26	14	1572	A	N1-C6-N6	5.92	122.15	118.60
1	13	963	G	N3-C4-N9	5.92	129.55	126.00
26	1H	944	G	C8-N9-C4	-5.92	104.03	106.40
26	1H	2570	G	N3-C2-N2	-5.92	115.75	119.90
26	14	1373	A	N7-C8-N9	-5.92	110.84	113.80
26	14	2866	U	C5-C4-O4	5.92	129.45	125.90
26	1H	2610	C	C6-N1-C2	5.92	122.67	120.30
26	1H	865	C	C6-N1-C2	5.92	122.67	120.30
26	1H	2609	U	C6-N1-C2	5.92	124.55	121.00
26	14	1304	C	N3-C4-N4	-5.92	113.86	118.00
26	1H	681	G	C8-N9-C4	5.92	108.77	106.40
26	14	669	G	C4-C5-N7	-5.92	108.43	110.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	1667	G	C5-C6-O6	-5.91	125.05	128.60
26	14	2459	A	C8-N9-C4	-5.91	103.44	105.80
26	14	2023	G	O5'-P-OP2	-5.91	100.38	105.70
26	14	2818	G	C5-C6-O6	-5.91	125.05	128.60
26	1H	636	G	O5'-P-OP1	-5.91	100.38	105.70
26	1H	928	G	N1-C6-O6	5.91	123.45	119.90
26	1H	2570	G	C2-N3-C4	-5.91	108.94	111.90
26	14	1337	G	OP1-P-O3'	5.91	118.20	105.20
26	1H	517	C	C5-C4-N4	-5.91	116.06	120.20
26	1H	1000	A	C8-N9-C4	-5.91	103.44	105.80
26	1H	2275	C	OP1-P-O3'	5.91	118.20	105.20
1	13	481	G	C8-N9-C1'	-5.91	119.32	127.00
1	13	567	G	O5'-P-OP1	-5.91	100.39	105.70
1	13	975	A	C5-N7-C8	-5.91	100.95	103.90
26	1H	1384	A	N1-C6-N6	-5.91	115.06	118.60
26	1H	1915	U	N3-C2-O2	-5.91	118.07	122.20
45	G8	81	LYS	C-N-CA	5.91	146.80	122.00
26	1H	659	C	C4-C5-C6	5.90	120.35	117.40
26	14	2606	C	N3-C4-C5	5.90	124.26	121.90
26	1H	2553	G	N3-C4-N9	5.90	129.54	126.00
26	14	939	G	N1-C6-O6	5.90	123.44	119.90
26	14	1779	U	OP1-P-OP2	5.90	128.46	119.60
26	1H	2430	A	N9-C4-C5	-5.90	103.44	105.80
26	14	715	G	C4-C5-N7	5.90	113.16	110.80
27	1J	22	U	C5-C6-N1	5.90	125.65	122.70
26	1H	2402	C	N1-C2-O2	-5.90	115.36	118.90
1	1G	1301	U	C2-N1-C1'	5.90	124.78	117.70
26	14	819	A	C8-N9-C4	-5.90	103.44	105.80
26	1H	2071	A	N1-C6-N6	5.90	122.14	118.60
1	1G	1200	C	C2-N1-C1'	5.90	125.29	118.80
26	14	1323	U	N3-C4-O4	5.90	123.53	119.40
26	1H	2689	U	P-O3'-C3'	5.90	126.78	119.70
26	14	1972	A	OP2-P-O3'	5.90	118.17	105.20
27	16	54	G	N1-C6-O6	5.89	123.44	119.90
26	14	1891	G	C4-N9-C1'	5.89	134.16	126.50
26	1H	613	U	N3-C4-O4	-5.89	115.28	119.40
28	11	147	LEU	CA-CB-CG	5.89	128.85	115.30
26	14	751	A	C2-N3-C4	-5.89	107.65	110.60
1	13	1277	C	C6-N1-C2	-5.89	117.94	120.30
26	1H	2642	G	C8-N9-C4	5.89	108.76	106.40
26	14	1761	C	N1-C2-O2	-5.89	115.37	118.90
1	1G	1529	G	C8-N9-C4	-5.89	104.05	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	1276	A	C4-C5-N7	5.89	113.64	110.70
26	1H	821	A	N1-C2-N3	5.88	132.24	129.30
26	14	12	U	N1-C2-O2	5.88	126.92	122.80
26	14	211	A	C8-N9-C4	5.88	108.15	105.80
26	14	577	G	C6-C5-N7	-5.88	126.87	130.40
26	14	71	A	C6-C5-N7	-5.88	128.18	132.30
26	14	2526	G	C5-C6-N1	-5.88	108.56	111.50
1	13	366	C	N1-C2-O2	-5.88	115.37	118.90
26	1H	1135	C	N1-C2-O2	5.88	122.43	118.90
26	1H	446	G	N1-C6-O6	5.88	123.43	119.90
26	1H	1349	A	N1-C6-N6	5.88	122.12	118.60
26	14	1786	A	C4-C5-N7	5.88	113.64	110.70
26	14	1688	U	C6-N1-C1'	5.88	129.43	121.20
26	14	1966	A	C5-C6-N6	-5.87	119.00	123.70
26	14	2452	C	N3-C4-N4	5.87	122.11	118.00
26	1H	482	A	C8-N9-C4	-5.87	103.45	105.80
26	14	1193	G	N1-C6-O6	5.87	123.42	119.90
26	14	1614	A	O5'-P-OP1	-5.87	100.42	105.70
23	2L	16	U	N3-C2-O2	-5.87	118.09	122.20
26	14	2683	C	N3-C4-C5	-5.87	119.55	121.90
1	13	792	A	N9-C1'-C2'	5.87	121.63	114.00
27	16	95	U	C2-N1-C1'	-5.87	110.66	117.70
26	1H	1627	G	N1-C6-O6	-5.87	116.38	119.90
26	14	372	G	O4'-C1'-N9	5.87	112.89	108.20
26	14	1395	A	O4'-C1'-N9	5.87	112.89	108.20
1	13	748	C	P-O3'-C3'	5.86	126.74	119.70
1	1G	1126	U	P-O3'-C3'	5.86	126.73	119.70
26	14	270(B)	A	N1-C6-N6	5.86	122.12	118.60
26	14	2445	G	N3-C4-C5	-5.86	125.67	128.60
26	14	1327	C	N3-C2-O2	5.86	126.00	121.90
26	14	2425	A	O5'-P-OP2	-5.86	100.43	105.70
1	1G	963	G	N1-C2-N2	-5.85	110.93	116.20
26	14	773	U	C5-C6-N1	-5.85	119.77	122.70
26	14	1950	G	C8-N9-C1'	-5.85	119.39	127.00
26	1H	1198	U	N3-C2-O2	-5.85	118.10	122.20
26	1H	1799	G	P-O3'-C3'	5.85	126.72	119.70
26	14	1613	G	N1-C6-O6	-5.85	116.39	119.90
26	14	2403	C	N1-C2-O2	-5.85	115.39	118.90
26	14	278	A	OP1-P-O3'	5.85	118.07	105.20
26	14	2433	A	C4-C5-C6	5.85	119.93	117.00
1	13	789	U	N1-C2-N3	5.85	118.41	114.90
26	1H	486	C	N3-C2-O2	5.85	125.99	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	649	G	N1-C6-O6	5.85	123.41	119.90
26	1H	1846	G	N1-C6-O6	5.85	123.41	119.90
26	14	566	U	C6-N1-C2	5.85	124.51	121.00
26	14	1964	G	O5'-P-OP1	-5.85	100.44	105.70
26	14	2329	G	C8-N9-C4	5.85	108.74	106.40
26	14	506	G	N1-C6-O6	5.85	123.41	119.90
1	13	811	C	C2-N3-C4	-5.85	116.98	119.90
26	14	121	G	C8-N9-C1'	-5.85	119.40	127.00
27	1J	6	C	C6-N1-C2	5.85	122.64	120.30
1	1G	795	C	N1-C2-O2	-5.84	115.39	118.90
26	14	288	C	C2-N1-C1'	5.84	125.23	118.80
26	14	1528	A	C5-N7-C8	-5.84	100.98	103.90
26	14	1989	G	N1-C2-N2	5.84	121.46	116.20
1	13	1518	A	O5'-P-OP2	-5.84	100.44	105.70
1	13	185	A	C8-N9-C4	-5.84	103.46	105.80
26	1H	246	C	C6-N1-C2	5.84	122.64	120.30
26	1H	2385	C	C2-N3-C4	-5.84	116.98	119.90
26	14	1681	G	C5-N7-C8	-5.84	101.38	104.30
26	1H	1204	A	C5-C6-N1	-5.84	114.78	117.70
26	14	923	C	C6-N1-C2	-5.84	117.96	120.30
26	14	2406	U	O4'-C1'-N1	-5.84	103.53	108.20
1	13	679	C	C6-N1-C2	-5.84	117.97	120.30
26	1H	117	G	O5'-P-OP2	5.84	117.70	110.70
26	1H	829	A	C6-N1-C2	5.84	122.10	118.60
26	1H	1900	A	C5'-C4'-O4'	-5.84	102.09	109.10
26	14	2079	U	O5'-P-OP1	-5.84	100.45	105.70
26	1H	703	U	O5'-P-OP1	-5.84	100.45	105.70
26	14	1608	A	N9-C4-C5	5.84	108.14	105.80
26	14	2505	G	O5'-P-OP1	5.84	117.70	110.70
26	1H	1308	A	N9-C4-C5	5.83	108.13	105.80
26	1H	1614	A	O5'-P-OP1	-5.83	100.45	105.70
26	1H	1210	A	C5-C6-N6	-5.83	119.03	123.70
26	1H	1627	G	O5'-P-OP2	-5.83	100.45	105.70
26	1H	1273	U	O5'-P-OP1	-5.83	100.45	105.70
26	1H	1496	A	C5-C6-N6	-5.83	119.04	123.70
26	14	2413	G	C5-C6-O6	-5.83	125.10	128.60
1	13	481	G	C4-C5-C6	5.83	122.30	118.80
26	1H	774	A	C8-N9-C1'	5.83	138.19	127.70
26	1H	1006	C	N1-C2-O2	-5.83	115.40	118.90
27	1J	81	G	C4-C5-N7	5.83	113.13	110.80
26	1H	201	C	C6-N1-C2	5.83	122.63	120.30
26	1H	186	G	N3-C4-N9	5.82	129.49	126.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	2594	C	C5-C4-N4	-5.82	116.12	120.20
26	14	675	A	N9-C4-C5	-5.82	103.47	105.80
27	1J	6	C	C5-C6-N1	-5.82	118.09	121.00
26	14	796	C	C5-C6-N1	-5.82	118.09	121.00
26	1H	594	U	C5-C4-O4	5.82	129.39	125.90
26	14	439	G	C4-C5-N7	5.82	113.13	110.80
26	14	801	G	C4-C5-C6	-5.82	115.31	118.80
1	13	31	G	C8-N9-C4	5.82	108.73	106.40
1	13	505	G	C4-C5-N7	5.82	113.13	110.80
26	1H	1257	C	N1-C2-N3	5.82	123.27	119.20
26	1H	1257	C	C4-C5-C6	5.82	120.31	117.40
26	1H	1546	C	C6-N1-C2	-5.82	117.97	120.30
26	1H	2888	C	C6-N1-C2	-5.82	117.97	120.30
1	1G	345	C	C6-N1-C2	-5.82	117.97	120.30
26	14	681	G	C6-C5-N7	-5.82	126.91	130.40
26	14	866	A	O4'-C1'-N9	-5.82	103.55	108.20
33	69	77	LEU	CA-CB-CG	5.82	128.68	115.30
1	13	1502	A	N1-C2-N3	5.82	132.21	129.30
26	14	382	G	C5-C6-N1	-5.82	108.59	111.50
26	14	1673	U	OP2-P-O3'	5.82	117.99	105.20
26	14	2346	A	C6-C5-N7	-5.82	128.23	132.30
26	1H	575	A	C8-N9-C4	5.81	108.13	105.80
26	1H	138	G	C5-C6-N1	5.81	114.41	111.50
26	1H	1742	C	C6-N1-C2	-5.81	117.97	120.30
29	21	49	LEU	CA-CB-CG	-5.81	101.93	115.30
31	41	94	LEU	CA-CB-CG	5.81	128.67	115.30
26	14	275	G	C4-N9-C1'	5.81	134.06	126.50
26	14	2499	C	N3-C2-O2	-5.81	117.83	121.90
26	14	860	U	N3-C2-O2	-5.81	118.13	122.20
1	13	481	G	C6-C5-N7	-5.81	126.91	130.40
26	1H	382	G	C8-N9-C4	5.81	108.72	106.40
26	14	365	C	C6-N1-C2	-5.81	117.98	120.30
26	1H	2503	A	O5'-P-OP1	5.80	117.66	110.70
26	14	2019	A	N7-C8-N9	-5.80	110.90	113.80
26	1H	138	G	C2-N3-C4	5.80	114.80	111.90
26	1H	609	A	C4-C5-N7	5.80	113.60	110.70
26	1H	1779	U	N1-C2-O2	5.80	126.86	122.80
1	1G	980	C	C6-N1-C2	-5.80	117.98	120.30
1	1G	266	G	P-O3'-C3'	5.80	126.66	119.70
26	1H	324	A	O5'-P-OP2	5.80	117.66	110.70
26	1H	409	C	C6-N1-C2	5.80	122.62	120.30
26	14	2062	A	C4-C5-C6	-5.80	114.10	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	13	703	G	C4-C5-N7	5.80	113.12	110.80
26	1H	67	U	N3-C2-O2	-5.80	118.14	122.20
26	1H	765	G	N1-C6-O6	5.80	123.38	119.90
48	J8	79	GLY	N-CA-C	5.80	127.59	113.10
49	K8	16	LEU	N-CA-C	-5.80	95.35	111.00
1	1G	86	U	C2-N1-C1'	5.79	124.65	117.70
26	14	2067	G	N3-C4-C5	-5.79	125.70	128.60
1	13	1493	A	N1-C6-N6	5.79	122.08	118.60
22	1K	40	G	N1-C6-O6	5.79	123.38	119.90
1	1G	945	G	N1-C6-O6	5.79	123.38	119.90
26	14	391	G	C4-C5-N7	5.79	113.12	110.80
26	1H	144	C	C5-C6-N1	-5.79	118.10	121.00
26	1H	2430	A	N3-C4-C5	5.79	130.85	126.80
26	14	1696	G	O5'-P-OP2	-5.79	100.49	105.70
26	1H	195	A	N1-C6-N6	5.79	122.07	118.60
1	1G	1206	G	C6-C5-N7	-5.79	126.93	130.40
26	14	203	C	N1-C2-O2	-5.79	115.43	118.90
26	14	2447	G	C6-C5-N7	5.79	133.87	130.40
26	14	621	A	C4-C5-N7	5.79	113.59	110.70
26	14	2441	C	O5'-P-OP1	-5.79	100.49	105.70
1	13	652	U	N1-C2-O2	5.79	126.85	122.80
26	1H	2330	G	C5-C6-O6	-5.79	125.13	128.60
26	1H	2330	G	N1-C2-N3	5.79	127.37	123.90
26	1H	2330	G	C6-C5-N7	-5.79	126.93	130.40
26	1H	2503	A	N1-C2-N3	-5.79	126.41	129.30
27	16	64	C	C5-C6-N1	-5.79	118.11	121.00
27	16	102	G	N1-C6-O6	5.79	123.37	119.90
26	14	140	A	O4'-C1'-N9	5.79	112.83	108.20
26	1H	1210	A	C6-C5-N7	-5.78	128.25	132.30
1	1G	353	A	C5-N7-C8	-5.78	101.01	103.90
26	14	2390	U	C6-N1-C2	-5.78	117.53	121.00
26	14	2779	U	C6-N1-C1'	-5.78	113.10	121.20
26	1H	2392	A	C2-N3-C4	-5.78	107.71	110.60
26	14	1637	A	N1-C6-N6	5.78	122.07	118.60
1	13	956	U	C6-N1-C2	-5.78	117.53	121.00
26	1H	397	G	N3-C4-C5	5.78	131.49	128.60
26	1H	564	C	N3-C4-C5	-5.78	119.59	121.90
26	14	577	G	C8-N9-C4	-5.78	104.09	106.40
26	14	2702	U	C6-N1-C1'	-5.78	113.11	121.20
1	13	59	A	C2-N3-C4	5.78	113.49	110.60
26	1H	70	G	N1-C6-O6	-5.78	116.43	119.90
26	1H	299	A	OP2-P-O3'	5.78	117.91	105.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	1950	G	N1-C2-N2	-5.78	111.00	116.20
1	1G	1139	G	N3-C4-N9	-5.78	122.53	126.00
26	1H	743	G	N3-C2-N2	-5.78	115.86	119.90
26	1H	2527	C	C5-C6-N1	5.78	123.89	121.00
27	16	30	C	C5-C6-N1	5.77	123.89	121.00
26	14	1564	C	N3-C2-O2	-5.77	117.86	121.90
26	1H	74	A	C8-N9-C4	-5.77	103.49	105.80
1	13	131	C	N3-C2-O2	-5.77	117.86	121.90
26	14	1336	A	N1-C6-N6	-5.77	115.14	118.60
26	1H	2331	G	C6-C5-N7	-5.77	126.94	130.40
26	14	53	A	N1-C2-N3	5.77	132.18	129.30
27	1J	55	U	C5-C6-N1	5.77	125.58	122.70
23	2L	28	C	C5-C6-N1	5.77	123.88	121.00
1	13	1176	A	N7-C8-N9	5.76	116.68	113.80
26	1H	513	A	N1-C6-N6	-5.76	115.14	118.60
26	1H	856	C	C5-C6-N1	5.76	123.88	121.00
26	1H	1448	G	N1-C6-O6	5.76	123.36	119.90
26	1H	1786	A	N1-C2-N3	5.76	132.18	129.30
26	14	1828	G	O5'-P-OP2	-5.76	100.51	105.70
26	14	2012	G	N1-C6-O6	5.76	123.36	119.90
1	13	1305	G	N3-C4-N9	-5.76	122.54	126.00
1	1G	1157	A	P-O3'-C3'	5.76	126.61	119.70
26	14	776	G	O4'-C1'-N9	-5.76	103.59	108.20
54	Q8	7	HIS	N-CA-C	5.76	126.55	111.00
1	13	803	G	C5-C6-O6	5.76	132.05	128.60
26	1H	513	A	N1-C2-N3	5.76	132.18	129.30
26	1H	1035	U	C5-C4-O4	5.76	129.35	125.90
26	14	2071	A	OP1-P-OP2	-5.76	110.97	119.60
26	1H	2401	U	C6-N1-C2	-5.75	117.55	121.00
26	1H	2449	U	N3-C4-O4	5.75	123.43	119.40
26	14	676	A	C5-C6-N6	-5.75	119.10	123.70
26	14	730	C	N3-C2-O2	-5.75	117.87	121.90
26	14	2425	A	N1-C6-N6	-5.75	115.15	118.60
1	13	1197	G	C8-N9-C4	-5.75	104.10	106.40
26	1H	2468	G	N7-C8-N9	5.75	115.98	113.10
1	13	193	C	C6-N1-C2	-5.75	118.00	120.30
1	13	549	C	O4'-C1'-N1	5.75	112.80	108.20
1	13	566	G	N3-C4-N9	5.75	129.45	126.00
26	1H	2053	G	C5-C6-O6	-5.75	125.15	128.60
26	14	391	G	C8-N9-C1'	-5.75	119.52	127.00
26	14	1908	C	C6-N1-C2	-5.75	118.00	120.30
26	1H	504	U	N3-C2-O2	-5.75	118.18	122.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	1343	G	C4-N9-C1'	5.75	133.97	126.50
27	16	81	G	N9-C4-C5	-5.75	103.10	105.40
26	1H	2710	C	N1-C2-O2	-5.75	115.45	118.90
1	1G	560	U	C6-N1-C2	-5.75	117.55	121.00
26	1H	1644	C	N3-C2-O2	-5.75	117.88	121.90
26	1H	1977	A	N1-C6-N6	-5.74	115.15	118.60
26	14	1728	G	C2-N3-C4	5.74	114.77	111.90
26	1H	618(A)	C	N3-C4-C5	5.74	124.20	121.90
26	14	250	G	C8-N9-C4	-5.74	104.10	106.40
1	13	579	G	C5-C6-O6	-5.74	125.16	128.60
1	13	1515	C	N3-C4-C5	5.74	124.19	121.90
1	1G	817	C	C5-C6-N1	-5.74	118.13	121.00
26	14	545	G	N1-C6-O6	-5.74	116.46	119.90
1	13	827	U	C2-N1-C1'	5.74	124.58	117.70
26	1H	1343	G	C8-N9-C4	-5.74	104.11	106.40
26	14	1407	C	C5-C6-N1	5.74	123.87	121.00
1	13	138	G	N1-C6-O6	5.73	123.34	119.90
26	1H	2674	G	C8-N9-C1'	5.73	134.45	127.00
26	1H	513	A	N3-C4-C5	-5.73	122.79	126.80
1	1G	249	U	O5'-P-OP2	-5.73	100.54	105.70
26	1H	1261	C	N3-C2-O2	5.73	125.91	121.90
26	1H	1799	G	N3-C4-N9	5.73	129.44	126.00
26	1H	2454	G	C8-N9-C4	5.73	108.69	106.40
26	14	27	G	C5-C6-N1	-5.73	108.64	111.50
26	1H	827	U	O5'-P-OP2	-5.73	100.55	105.70
1	13	792	A	C2-N3-C4	-5.72	107.74	110.60
26	1H	576	U	OP2-P-O3'	5.72	117.79	105.20
26	14	679	C	N1-C2-O2	-5.72	115.47	118.90
26	14	2584	U	O4'-C1'-N1	5.72	112.78	108.20
1	13	758	G	C4-C5-N7	5.72	113.09	110.80
26	1H	612	G	C5-C6-O6	-5.72	125.17	128.60
39	A8	101	LEU	CA-CB-CG	5.72	128.46	115.30
26	14	362	U	N3-C2-O2	-5.72	118.19	122.20
26	14	683	C	N1-C2-O2	-5.72	115.47	118.90
23	2K	44	G	C8-N9-C4	-5.72	104.11	106.40
26	1H	2639	A	C2-N3-C4	-5.72	107.74	110.60
26	1H	1665	A	N1-C6-N6	5.72	122.03	118.60
26	14	1469	A	C8-N9-C4	-5.72	103.51	105.80
26	14	1678	G	N1-C2-N3	5.72	127.33	123.90
26	1H	309	G	C4-N9-C1'	5.72	133.93	126.50
26	1H	2509	G	C6-C5-N7	-5.72	126.97	130.40
1	1G	390	C	C5-C6-N1	-5.72	118.14	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	1G	397	A	C8-N9-C4	-5.72	103.51	105.80
26	14	1653	G	N3-C4-N9	5.72	129.43	126.00
26	14	1899	G	P-O3'-C3'	5.71	126.56	119.70
1	13	758	G	N1-C6-O6	5.71	123.33	119.90
1	13	965	A	N1-C6-N6	5.71	122.03	118.60
26	1H	917	A	C5-C6-N1	-5.71	114.84	117.70
22	1K	85	C	N1-C2-O2	5.71	122.33	118.90
26	1H	440	G	N3-C4-N9	5.71	129.43	126.00
1	13	1336	C	P-O3'-C3'	5.71	126.55	119.70
26	1H	678	C	C6-N1-C2	5.71	122.58	120.30
26	1H	1972	A	C5-N7-C8	-5.71	101.05	103.90
23	2L	27	G	N1-C6-O6	5.71	123.32	119.90
26	14	796	C	C2-N3-C4	-5.71	117.05	119.90
26	14	1028	A	OP2-P-O3'	5.71	117.75	105.20
26	1H	954	G	C4-C5-N7	-5.71	108.52	110.80
26	1H	121	G	N7-C8-N9	5.70	115.95	113.10
26	14	666	G	C8-N9-C4	5.70	108.68	106.40
26	14	1555	G	C4-N9-C1'	5.70	133.91	126.50
1	13	700	G	N3-C4-N9	5.70	129.42	126.00
26	14	1667	G	C4-C5-N7	5.70	113.08	110.80
26	1H	955	C	N3-C4-C5	-5.70	119.62	121.90
26	1H	2454	G	C4-C5-N7	-5.70	108.52	110.80
26	1H	2454	G	C5-C6-O6	5.70	132.02	128.60
26	14	1342	A	C4-C5-C6	5.70	119.85	117.00
26	1H	871	U	O5'-P-OP2	5.70	117.54	110.70
26	14	180	G	N1-C6-O6	5.70	123.32	119.90
1	13	534	U	N3-C4-O4	-5.70	115.41	119.40
26	1H	51	G	OP2-P-O3'	5.70	117.73	105.20
26	1H	831	G	C5-N7-C8	5.70	107.15	104.30
1	1G	372	C	N1-C2-N3	-5.70	115.21	119.20
1	1G	1107	C	C6-N1-C2	-5.70	118.02	120.30
1	1G	1442	G	N3-C4-C5	5.70	131.45	128.60
26	14	2413	G	N1-C6-O6	5.70	123.32	119.90
23	2K	87	A	N9-C4-C5	-5.69	103.52	105.80
26	14	808	G	C6-N1-C2	-5.69	121.68	125.10
26	1H	865	C	N3-C4-C5	5.69	124.18	121.90
26	1H	924	C	N3-C4-C5	5.69	124.18	121.90
26	1H	2025	C	C6-N1-C2	-5.69	118.02	120.30
26	14	1187	G	N1-C6-O6	5.69	123.32	119.90
26	14	1637	A	C6-C5-N7	-5.69	128.31	132.30
1	13	703	G	N1-C6-O6	5.69	123.31	119.90
24	3K	2	C	C5-C6-N1	5.69	123.84	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	138	G	C5-N7-C8	-5.69	101.45	104.30
26	1H	2346	A	C5-N7-C8	-5.69	101.06	103.90
26	1H	2350	C	C6-N1-C2	-5.69	118.02	120.30
42	D8	18	LEU	CA-CB-CG	5.69	128.39	115.30
35	25	50	GLY	N-CA-C	-5.69	98.87	113.10
26	1H	2046	G	N3-C4-C5	-5.69	125.75	128.60
26	1H	246	C	C5-C6-N1	-5.69	118.16	121.00
26	1H	679	C	N3-C2-O2	5.69	125.88	121.90
26	1H	1888	G	C4-N9-C1'	5.69	133.89	126.50
26	1H	2327	A	C2-N3-C4	5.69	113.44	110.60
26	1H	2596	U	N3-C4-C5	5.69	118.01	114.60
26	14	932	G	N3-C4-N9	-5.69	122.59	126.00
26	1H	101	G	N9-C4-C5	-5.69	103.12	105.40
54	M5	32	LEU	CA-CB-CG	5.68	128.38	115.30
26	14	545	G	O4'-C1'-N9	5.68	112.75	108.20
26	14	1474	C	C6-N1-C2	-5.68	118.03	120.30
26	14	2859	G	P-O3'-C3'	5.68	126.52	119.70
26	1H	581	C	N1-C2-O2	-5.68	115.49	118.90
26	1H	2819	G	N1-C6-O6	5.68	123.31	119.90
26	14	1613	G	N3-C2-N2	5.68	123.88	119.90
26	1H	783	A	C5-C6-N6	-5.68	119.16	123.70
26	14	1695	G	N3-C4-C5	-5.68	125.76	128.60
26	1H	2430	A	C4-C5-N7	5.67	113.54	110.70
26	14	1651	G	C4-C5-N7	5.67	113.07	110.80
26	1H	946	G	N3-C4-C5	5.67	131.44	128.60
26	1H	1940	U	N3-C2-O2	5.67	126.17	122.20
26	1H	1609	A	N1-C6-N6	-5.67	115.20	118.60
26	1H	2689	U	C2-N3-C4	-5.67	123.60	127.00
26	14	1379	A	C5-N7-C8	-5.67	101.06	103.90
26	14	2048	G	C8-N9-C4	-5.67	104.13	106.40
26	1H	134	C	N3-C2-O2	-5.67	117.93	121.90
26	1H	2619	C	C5-C4-N4	-5.67	116.23	120.20
1	13	1498	U	N3-C4-O4	5.67	123.37	119.40
26	14	2062	A	C4-N9-C1'	-5.67	116.10	126.30
26	1H	1025	G	N3-C4-C5	-5.66	125.77	128.60
26	1H	2243	U	O5'-P-OP1	5.66	117.50	110.70
26	1H	2392	A	C6-C5-N7	-5.66	128.34	132.30
26	14	2443	C	C5-C4-N4	-5.66	116.24	120.20
1	13	394	G	N3-C4-N9	-5.66	122.60	126.00
54	Q8	60	LEU	CB-CG-CD2	5.66	120.62	111.00
26	1H	729	G	C5-N7-C8	-5.66	101.47	104.30
1	1G	366	C	C6-N1-C2	5.66	122.56	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	13	1158	C	C6-N1-C2	-5.65	118.04	120.30
26	1H	127	A	C5-N7-C8	-5.65	101.07	103.90
26	1H	703	U	C5-C4-O4	5.65	129.29	125.90
1	1G	1415	G	C8-N9-C4	5.65	108.66	106.40
26	14	757	U	OP2-P-O3'	5.65	117.63	105.20
26	1H	1294	U	N3-C2-O2	5.65	126.16	122.20
26	1H	842	G	OP2-P-O3'	5.65	117.63	105.20
26	1H	859	G	N1-C6-O6	5.65	123.29	119.90
26	1H	735	A	N9-C4-C5	-5.65	103.54	105.80
26	1H	346	A	O5'-P-OP2	-5.65	100.62	105.70
22	1K	18	G	C8-N9-C1'	5.64	134.34	127.00
26	1H	191	A	N1-C2-N3	5.64	132.12	129.30
26	1H	1366	A	N1-C6-N6	5.64	121.99	118.60
26	1H	2525	G	C6-C5-N7	-5.64	127.02	130.40
26	14	2844	G	OP2-P-O3'	5.64	117.61	105.20
27	16	7	G	C8-N9-C4	5.64	108.66	106.40
26	1H	801	G	N1-C6-O6	-5.64	116.52	119.90
26	1H	1977	A	C5-C6-N6	5.64	128.21	123.70
26	1H	2409	G	C5-C6-O6	-5.64	125.22	128.60
26	1H	2593	U	N3-C4-O4	5.64	123.35	119.40
1	1G	197	A	C6-C5-N7	-5.64	128.35	132.30
26	14	465	G	C4-C5-N7	-5.64	108.55	110.80
26	1H	1368	G	C5-C6-O6	-5.64	125.22	128.60
26	1H	2419	U	OP1-P-O3'	5.64	117.60	105.20
26	1H	2525	G	N1-C6-O6	5.64	123.28	119.90
1	1G	236	G	C5-C6-N1	-5.64	108.68	111.50
26	14	728	G	C8-N9-C4	5.64	108.65	106.40
26	14	2073	C	N1-C2-O2	-5.64	115.52	118.90
1	13	1489	G	C8-N9-C4	5.63	108.65	106.40
26	1H	965	C	C5-C6-N1	5.63	123.82	121.00
26	14	33	U	C6-N1-C2	5.63	124.38	121.00
26	14	1966	A	N9-C4-C5	-5.63	103.55	105.80
1	13	1481	U	N3-C4-C5	-5.63	111.22	114.60
26	1H	1206	G	N3-C4-C5	-5.63	125.78	128.60
2	12	23	ARG	N-CA-C	-5.63	95.80	111.00
26	14	383	U	O4'-C1'-N1	5.63	112.70	108.20
26	14	726	G	O4'-C1'-N9	5.63	112.70	108.20
26	14	1312	U	N3-C2-O2	5.63	126.14	122.20
26	1H	1311	G	N3-C4-C5	5.63	131.41	128.60
26	14	4	C	C6-N1-C1'	-5.63	114.04	120.80
26	14	545	G	C4-C5-N7	-5.63	108.55	110.80
26	1H	1828	G	N3-C4-C5	5.63	131.41	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	388	G	N3-C4-N9	-5.63	122.62	126.00
26	1H	1558	A	N1-C2-N3	5.63	132.11	129.30
26	14	647	G	C8-N9-C4	-5.63	104.15	106.40
26	14	848	G	N3-C4-C5	-5.63	125.79	128.60
26	1H	1489	U	C5-C4-O4	5.62	129.28	125.90
1	13	503	C	C6-N1-C2	-5.62	118.05	120.30
26	1H	122	G	O5'-P-OP2	-5.62	100.64	105.70
26	1H	2422	A	O4'-C1'-N9	5.62	112.70	108.20
26	14	667	U	N1-C2-O2	-5.62	118.86	122.80
26	14	801	G	N3-C4-C5	5.62	131.41	128.60
1	1G	267	C	O5'-P-OP1	-5.62	100.64	105.70
26	14	757	U	N1-C2-O2	-5.62	118.86	122.80
26	14	941	A	C8-N9-C4	-5.62	103.55	105.80
26	14	1769	G	N1-C6-O6	5.62	123.27	119.90
26	14	300	A	N1-C6-N6	5.62	121.97	118.60
26	14	1977	A	C2-N3-C4	-5.62	107.79	110.60
26	1H	1022	G	C4-C5-N7	-5.62	108.55	110.80
1	13	757	U	N3-C2-O2	-5.62	118.27	122.20
26	1H	829	A	N3-C4-C5	5.62	130.73	126.80
26	1H	837	C	N3-C4-N4	5.62	121.93	118.00
26	1H	618(A)	C	C6-N1-C2	5.62	122.55	120.30
26	14	213	A	C8-N9-C4	5.62	108.05	105.80
26	1H	923	C	N3-C4-C5	-5.61	119.66	121.90
1	1G	243	A	P-O3'-C3'	5.61	126.44	119.70
26	14	675	A	C5-C6-N6	-5.61	119.21	123.70
26	1H	954	G	C6-C5-N7	5.61	133.77	130.40
4	3E	32	ALA	N-CA-C	-5.61	95.85	111.00
26	14	922	U	C5-C6-N1	5.61	125.50	122.70
26	1H	377	C	N3-C4-C5	5.61	124.14	121.90
26	1H	774	A	O5'-P-OP2	-5.61	100.65	105.70
26	1H	2475	C	C5-C6-N1	5.61	123.81	121.00
26	14	789	A	OP1-P-OP2	-5.61	111.19	119.60
26	1H	1255	U	N3-C4-O4	5.61	123.33	119.40
1	13	853	G	C8-N9-C4	5.61	108.64	106.40
26	14	1193	G	C5-C6-O6	-5.61	125.24	128.60
26	14	1313	U	C6-N1-C2	-5.61	117.64	121.00
26	14	1950	G	C4-C5-N7	5.61	113.04	110.80
26	14	1332	G	N3-C2-N2	-5.60	115.98	119.90
26	14	1787	A	C5-N7-C8	-5.60	101.10	103.90
26	1H	1638	C	O5'-P-OP2	-5.60	100.66	105.70
26	1H	2621	A	N1-C6-N6	-5.60	115.24	118.60
26	1H	2856	C	C5-C6-N1	5.60	123.80	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	1G	500	G	N3-C4-N9	5.60	129.36	126.00
1	1G	784	C	C6-N1-C2	5.60	122.54	120.30
26	14	1011	G	C4-N9-C1'	-5.60	119.22	126.50
27	16	60	C	N3-C4-N4	5.60	121.92	118.00
26	14	2429	G	OP2-P-O3'	5.60	117.52	105.20
26	14	2826	A	N9-C4-C5	5.60	108.04	105.80
1	13	1413	A	C5-N7-C8	-5.60	101.10	103.90
26	1H	1698	A	C4-N9-C1'	5.60	136.38	126.30
26	1H	1787	A	C4-C5-C6	5.60	119.80	117.00
26	14	1559	G	N1-C6-O6	5.60	123.26	119.90
26	14	1790	C	OP1-P-O3'	5.60	117.52	105.20
26	14	2335	A	O4'-C1'-N9	5.60	112.68	108.20
1	13	794	A	N1-C6-N6	5.60	121.96	118.60
26	1H	1031	G	N1-C6-O6	-5.60	116.54	119.90
1	13	266	G	P-O3'-C3'	5.59	126.41	119.70
26	14	1651	G	C5-C6-O6	-5.59	125.24	128.60
26	1H	1576	U	N3-C2-O2	-5.59	118.28	122.20
26	1H	1678	G	C4-C5-N7	5.59	113.04	110.80
26	14	391	G	C5-C6-N1	-5.59	108.70	111.50
1	13	304	U	C5-C4-O4	5.59	129.25	125.90
1	13	888	G	C8-N9-C4	5.59	108.64	106.40
26	1H	751	A	N1-C2-N3	5.59	132.10	129.30
22	1K	18	G	OP1-P-O3'	5.59	117.50	105.20
26	1H	1313	U	C6-N1-C2	-5.59	117.65	121.00
26	1H	1788	C	O5'-P-OP2	-5.59	100.67	105.70
1	1G	730	G	C4-C5-N7	-5.59	108.56	110.80
26	1H	735	A	C2-N3-C4	-5.59	107.81	110.60
26	14	141	A	O4'-C1'-N9	5.59	112.67	108.20
1	13	963	G	N1-C2-N2	-5.59	111.17	116.20
26	1H	2402	C	C6-N1-C1'	5.59	127.50	120.80
27	16	95	U	N1-C2-O2	-5.59	118.89	122.80
26	14	469	G	C5-C6-N1	5.58	114.29	111.50
26	14	1706	U	N3-C4-C5	-5.58	111.25	114.60
26	1H	807	U	C4-C5-C6	5.58	123.05	119.70
26	1H	1626	G	N9-C4-C5	5.58	107.63	105.40
26	1H	2445	G	N1-C6-O6	-5.58	116.55	119.90
45	G8	80	GLY	N-CA-C	5.58	127.06	113.10
1	1G	345	C	C5-C6-N1	5.58	123.79	121.00
26	14	698	C	C6-N1-C2	5.58	122.53	120.30
26	14	2702	U	C5-C6-N1	5.58	125.49	122.70
1	13	773	G	C4-N9-C1'	5.58	133.76	126.50
1	1G	108	G	C4-C5-N7	5.58	113.03	110.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	676	A	N3-C4-N9	-5.58	122.93	127.40
26	14	1790	C	C5-C6-N1	-5.58	118.21	121.00
26	1H	131	G	C5-C6-O6	-5.58	125.25	128.60
26	1H	781	A	P-O3'-C3'	5.58	126.39	119.70
26	1H	1446	C	C6-N1-C2	-5.58	118.07	120.30
26	1H	2606	C	OP1-P-O3'	5.58	117.47	105.20
1	1G	1519	A	C8-N9-C4	-5.58	103.57	105.80
26	14	140	A	C8-N9-C4	-5.58	103.57	105.80
26	14	582	G	C6-C5-N7	-5.58	127.05	130.40
26	14	1187	G	N3-C2-N2	-5.58	116.00	119.90
26	1H	380	U	N3-C2-O2	-5.58	118.30	122.20
26	1H	735	A	N7-C8-N9	-5.58	111.01	113.80
26	1H	953	A	C8-N9-C4	5.58	108.03	105.80
26	1H	1204	A	C5-N7-C8	-5.58	101.11	103.90
1	1G	244	U	O5'-P-OP1	-5.58	100.68	105.70
26	14	1533	C	C2-N1-C1'	5.58	124.93	118.80
27	1J	96	G	N3-C2-N2	-5.58	116.00	119.90
1	13	505	G	C6-C5-N7	-5.57	127.06	130.40
26	1H	1799	G	C2-N3-C4	5.57	114.69	111.90
26	14	1779	U	C6-N1-C1'	-5.57	113.40	121.20
26	1H	201	C	O5'-P-OP2	-5.57	100.69	105.70
26	1H	130	C	N3-C4-C5	5.57	124.13	121.90
26	1H	2071	A	OP2-P-O3'	5.57	117.45	105.20
1	13	1343	G	N1-C6-O6	5.57	123.24	119.90
1	13	1496	C	O5'-P-OP2	-5.57	100.69	105.70
26	1H	1022	G	C8-N9-C4	-5.57	104.17	106.40
26	1H	1697	G	C5-C6-O6	-5.57	125.26	128.60
26	1H	2562	U	N3-C2-O2	-5.57	118.30	122.20
26	1H	2688	U	C6-N1-C2	-5.57	117.66	121.00
1	13	748	C	C6-N1-C2	-5.57	118.07	120.30
26	14	2428	G	C5-C6-O6	5.57	131.94	128.60
42	95	86	GLY	N-CA-C	-5.57	99.19	113.10
26	1H	629	G	N1-C6-O6	-5.56	116.56	119.90
1	1G	21	G	C8-N9-C4	5.56	108.63	106.40
1	13	247	G	C8-N9-C4	-5.56	104.17	106.40
1	13	413	G	N3-C4-C5	-5.56	125.82	128.60
26	14	2433	A	C8-N9-C4	-5.56	103.58	105.80
1	13	1412	C	C6-N1-C2	5.56	122.52	120.30
26	14	2509	G	C4-N9-C1'	5.56	133.73	126.50
26	1H	845	G	C5-C6-O6	-5.56	125.26	128.60
26	1H	954	G	N1-C6-O6	-5.56	116.56	119.90
26	1H	2229	C	C6-N1-C2	5.56	122.52	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
41	C8	96	ALA	N-CA-C	-5.56	95.99	111.00
26	14	1411	C	N3-C2-O2	-5.56	118.01	121.90
26	14	1682	G	C6-C5-N7	-5.56	127.06	130.40
2	1E	196	LEU	CA-CB-CG	5.56	128.08	115.30
26	1H	1933	G	N3-C4-C5	-5.56	125.82	128.60
26	1H	2311	A	N1-C2-N3	5.56	132.08	129.30
26	14	791	C	C6-N1-C2	5.56	122.52	120.30
26	14	1681	G	N1-C6-O6	5.56	123.23	119.90
35	25	8	LEU	CA-CB-CG	5.56	128.08	115.30
26	1H	1008	C	N1-C2-O2	-5.56	115.57	118.90
26	14	2777	G	O4'-C1'-N9	-5.56	103.75	108.20
1	13	534	U	C2-N1-C1'	-5.55	111.04	117.70
26	1H	621	A	C4-C5-N7	5.55	113.48	110.70
1	1G	545	C	C6-N1-C2	-5.55	118.08	120.30
26	14	647	G	C6-C5-N7	-5.55	127.07	130.40
26	14	788	A	O5'-P-OP1	-5.55	100.70	105.70
26	14	1988	C	C6-N1-C2	5.55	122.52	120.30
26	14	2369	A	C8-N9-C4	-5.55	103.58	105.80
26	1H	38	A	N1-C6-N6	5.55	121.93	118.60
26	1H	205	G	N3-C2-N2	5.55	123.78	119.90
26	1H	622	G	C6-C5-N7	-5.55	127.07	130.40
26	1H	1989	G	O5'-P-OP1	-5.55	100.70	105.70
26	1H	2503	A	C4-C5-N7	5.55	113.48	110.70
26	1H	2724	C	N1-C2-O2	-5.55	115.57	118.90
1	1G	500	G	N3-C4-C5	-5.55	125.82	128.60
1	1G	1522	U	C6-N1-C2	-5.55	117.67	121.00
26	14	130	C	C6-N1-C2	5.55	122.52	120.30
26	14	1332	G	C8-N9-C1'	5.55	134.22	127.00
26	14	2701	C	P-O3'-C3'	5.55	126.36	119.70
26	1H	571	A	N1-C6-N6	5.55	121.93	118.60
26	14	761	A	OP1-P-O3'	5.55	117.41	105.20
26	14	470	A	C5-C6-N6	-5.55	119.26	123.70
26	14	1021	A	C5-C6-N1	-5.55	114.93	117.70
26	1H	725	G	O5'-P-OP1	-5.54	100.71	105.70
26	1H	2026	C	C6-N1-C2	5.54	122.52	120.30
26	1H	2674	G	C4-N9-C1'	-5.54	119.29	126.50
26	14	1550	C	C6-N1-C2	-5.54	118.08	120.30
26	1H	463	G	C4-C5-N7	5.54	113.02	110.80
26	1H	1598	C	OP1-P-O3'	5.54	117.40	105.20
1	1G	258	G	C6-C5-N7	-5.54	127.07	130.40
26	14	725	G	C6-C5-N7	-5.54	127.07	130.40
26	14	1825	A	C5-C6-N6	-5.54	119.27	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
27	1J	52	A	P-O3'-C3'	5.54	126.35	119.70
26	1H	462	C	N3-C4-N4	-5.54	114.12	118.00
26	14	1313	U	N3-C2-O2	-5.54	118.32	122.20
26	1H	1969	A	N1-C6-N6	-5.54	115.28	118.60
26	1H	2438	U	O5'-P-OP2	-5.54	100.72	105.70
1	1G	402	G	N1-C6-O6	5.54	123.22	119.90
26	14	196	A	O4'-C1'-N9	5.54	112.63	108.20
26	14	654(I)	C	N1-C2-O2	5.54	122.22	118.90
33	61	131	LYS	C-N-CD	-5.54	108.42	120.60
26	14	973	A	C8-N9-C4	5.54	108.01	105.80
26	14	2035	G	O4'-C1'-N9	5.54	112.63	108.20
26	1H	446	G	C6-C5-N7	-5.53	127.08	130.40
26	1H	692	C	C5-C6-N1	-5.53	118.23	121.00
26	14	970	C	O5'-P-OP1	-5.53	100.72	105.70
26	14	1858	G	C5-C6-N1	-5.53	108.73	111.50
26	14	2622	C	C5-C6-N1	-5.53	118.23	121.00
1	1G	767	A	C5-C6-N6	-5.53	119.28	123.70
26	1H	729	G	N7-C8-N9	5.53	115.87	113.10
1	1G	250	A	P-O3'-C3'	5.53	126.33	119.70
26	14	758	C	O5'-P-OP2	-5.53	100.73	105.70
26	14	1522	G	C5-C6-O6	-5.53	125.28	128.60
26	14	1989	G	C8-N9-C1'	5.53	134.19	127.00
1	13	186(A)	C	C5-C6-N1	5.53	123.76	121.00
26	1H	600	G	N7-C8-N9	-5.53	110.34	113.10
26	1H	1821	A	N1-C2-N3	5.53	132.06	129.30
26	1H	1660	C	C5-C6-N1	-5.52	118.24	121.00
27	16	38	C	N1-C2-O2	-5.52	115.59	118.90
23	2L	56	G	C4-N9-C1'	5.52	133.68	126.50
26	1H	2022	U	N3-C4-C5	5.52	117.91	114.60
26	14	968	G	N3-C4-C5	5.52	131.36	128.60
26	14	2346	A	C4-C5-C6	5.52	119.76	117.00
26	1H	1938	A	O4'-C1'-N9	5.52	112.62	108.20
26	14	1304	C	N3-C4-C5	5.52	124.11	121.90
26	1H	1327	C	O5'-P-OP1	-5.52	100.73	105.70
26	1H	2544	G	C8-N9-C4	5.52	108.61	106.40
26	1H	141	A	O4'-C1'-N9	5.52	112.61	108.20
26	1H	761	A	OP1-P-OP2	-5.52	111.32	119.60
26	1H	763	G	N3-C4-C5	-5.52	125.84	128.60
1	1G	413	G	O4'-C1'-N9	5.52	112.61	108.20
26	14	2516	G	C6-N1-C2	-5.52	121.79	125.10
26	1H	918	A	O5'-P-OP2	5.51	117.32	110.70
26	1H	1603	A	C8-N9-C4	-5.51	103.59	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	1G	1386	G	N1-C6-O6	-5.51	116.59	119.90
26	14	974(A)	C	C2-N1-C1'	5.51	124.87	118.80
26	14	1786	A	C5-C6-N1	-5.51	114.94	117.70
26	14	2001	A	C5-C6-N6	-5.51	119.29	123.70
1	13	1224	G	O4'-C1'-N9	5.51	112.61	108.20
26	1H	566	U	C5-C4-O4	-5.51	122.59	125.90
26	1H	728	G	C8-N9-C1'	-5.51	119.83	127.00
1	13	247	G	C2-N3-C4	5.51	114.66	111.90
26	1H	1797	C	C6-N1-C2	5.51	122.50	120.30
26	14	832	G	N3-C2-N2	-5.51	116.04	119.90
1	13	191(F)	U	C5-C6-N1	5.51	125.45	122.70
1	13	1504	G	O5'-P-OP2	5.51	117.31	110.70
26	1H	383	U	O4'-C1'-N1	5.51	112.61	108.20
26	1H	745	G	C6-C5-N7	-5.51	127.09	130.40
26	1H	2765	A	C2-N3-C4	-5.51	107.84	110.60
1	13	854	G	N1-C6-O6	5.51	123.20	119.90
1	13	300	A	O5'-P-OP1	-5.51	100.74	105.70
26	1H	2585	U	N1-C2-O2	5.51	126.65	122.80
26	14	330	A	N7-C8-N9	5.51	116.55	113.80
26	14	2076	U	C5-C6-N1	5.51	125.45	122.70
26	1H	622	G	C4-N9-C1'	5.50	133.66	126.50
26	14	2056	G	N3-C2-N2	-5.50	116.05	119.90
26	1H	921	G	C8-N9-C4	-5.50	104.20	106.40
26	1H	1364	G	N3-C4-N9	5.50	129.30	126.00
26	1H	1413	G	C6-C5-N7	-5.50	127.10	130.40
26	1H	1829	A	OP1-P-OP2	5.50	127.85	119.60
26	14	2426	A	C8-N9-C4	-5.50	103.60	105.80
26	1H	1604	C	C5-C4-N4	-5.50	116.35	120.20
6	52	14	LEU	CA-CB-CG	5.50	127.95	115.30
1	13	1058	G	C8-N9-C4	5.50	108.60	106.40
26	1H	2409	G	C5-N7-C8	-5.50	101.55	104.30
26	1H	2553	G	N1-C6-O6	5.50	123.20	119.90
26	14	1469	A	N7-C8-N9	5.50	116.55	113.80
1	13	266	G	O4'-C1'-N9	-5.50	103.80	108.20
1	13	578	C	C5-C6-N1	5.50	123.75	121.00
26	1H	514	A	C5-C6-N1	5.50	120.45	117.70
26	1H	2597	G	O5'-P-OP1	5.50	117.30	110.70
26	14	1963	U	N3-C2-O2	-5.50	118.35	122.20
26	14	1986	A	N1-C2-N3	5.50	132.05	129.30
26	14	2068	U	O5'-P-OP1	-5.50	100.75	105.70
26	14	2509	G	C4-C5-C6	5.50	122.10	118.80
26	14	2818	G	C8-N9-C4	5.50	108.60	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
27	1J	98	G	N9-C4-C5	-5.50	103.20	105.40
26	14	362	U	C2-N1-C1'	5.50	124.29	117.70
26	14	741	G	O5'-P-OP2	5.50	117.29	110.70
26	14	308	G	C6-C5-N7	-5.49	127.10	130.40
1	1G	46	G	C5-C6-O6	-5.49	125.31	128.60
26	14	511	U	C6-N1-C2	-5.49	117.70	121.00
26	1H	796	C	O5'-P-OP1	5.49	117.29	110.70
26	1H	840	C	C5-C6-N1	-5.49	118.25	121.00
26	1H	1675	C	O5'-P-OP1	-5.49	100.76	105.70
26	1H	1616	A	C4-N9-C1'	5.49	136.18	126.30
26	1H	2713	A	O4'-C1'-N9	-5.49	103.81	108.20
1	1G	913	A	P-O3'-C3'	5.49	126.29	119.70
26	1H	2056	G	N3-C4-N9	5.49	129.29	126.00
22	1K	85	C	N3-C4-C5	5.49	124.09	121.90
26	1H	194	G	N7-C8-N9	-5.49	110.36	113.10
26	14	2433	A	N7-C8-N9	5.49	116.54	113.80
26	14	2518	A	C5-C6-N1	-5.49	114.96	117.70
26	1H	569	U	C5-C6-N1	-5.48	119.96	122.70
37	88	24	GLY	N-CA-C	-5.48	99.39	113.10
26	14	527	C	N3-C4-N4	-5.48	114.16	118.00
26	1H	2002	G	C4-C5-N7	5.48	112.99	110.80
1	1G	1383	C	C6-N1-C2	5.48	122.49	120.30
26	14	271(B)	G	N3-C4-C5	-5.48	125.86	128.60
26	14	1637	A	C5-C6-N6	-5.48	119.31	123.70
26	1H	1537	C	C6-N1-C2	-5.48	118.11	120.30
26	1H	1858	G	P-O3'-C3'	5.48	126.28	119.70
26	1H	621	A	C6-C5-N7	-5.48	128.47	132.30
26	14	933	A	N7-C8-N9	5.48	116.54	113.80
26	14	1413	G	N1-C6-O6	5.48	123.19	119.90
26	14	2704	C	N3-C4-C5	-5.48	119.71	121.90
26	1H	1219	G	C5-C6-O6	-5.48	125.31	128.60
26	1H	2779	U	C5-C6-N1	-5.48	119.96	122.70
26	1H	2319	G	N1-C6-O6	-5.48	116.61	119.90
26	1H	1830	C	C2-N1-C1'	-5.47	112.78	118.80
1	1G	372	C	C6-N1-C1'	-5.47	114.23	120.80
26	14	2612	C	N1-C2-O2	5.47	122.18	118.90
26	1H	372	G	N1-C6-O6	-5.47	116.62	119.90
26	1H	1332	G	OP1-P-O3'	5.47	117.24	105.20
26	14	456	C	C6-N1-C2	5.47	122.49	120.30
26	14	783	A	N3-C4-N9	-5.47	123.02	127.40
26	14	2290	G	N1-C6-O6	5.47	123.18	119.90
1	13	108	G	N7-C8-N9	5.47	115.83	113.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	687	C	N3-C4-C5	5.47	124.09	121.90
26	1H	1026	U	C2-N1-C1'	5.47	124.26	117.70
26	1H	1328	G	N9-C4-C5	-5.47	103.21	105.40
26	1H	2288	A	C4-C5-N7	5.47	113.43	110.70
26	14	447	A	N1-C2-N3	5.47	132.03	129.30
26	1H	1313	U	C5-C6-N1	5.47	125.43	122.70
26	1H	828	U	N1-C2-N3	5.47	118.18	114.90
45	G8	79	CYS	N-CA-C	5.47	125.76	111.00
1	13	1178	G	C8-N9-C4	-5.46	104.21	106.40
26	1H	109	G	N3-C4-N9	5.46	129.28	126.00
26	1H	1758	G	N1-C6-O6	-5.46	116.62	119.90
26	1H	1781	C	C6-N1-C1'	-5.46	114.24	120.80
26	1H	2383	G	C8-N9-C1'	-5.46	119.90	127.00
26	1H	2719	G	OP1-P-OP2	-5.46	111.41	119.60
1	1G	20	U	N3-C4-C5	5.46	117.88	114.60
26	1H	2066	C	C6-N1-C2	-5.46	118.11	120.30
26	1H	2383	G	C4-N9-C1'	5.46	133.60	126.50
26	1H	2451	A	N9-C4-C5	5.46	107.98	105.80
26	1H	1528	A	C5-N7-C8	-5.46	101.17	103.90
26	14	275	G	C4-C5-N7	5.46	112.98	110.80
26	14	499	U	N1-C2-N3	5.46	118.18	114.90
26	14	1970	A	C4-C5-N7	5.46	113.43	110.70
26	14	2209	C	N1-C2-O2	-5.46	115.62	118.90
1	13	1203	C	C5-C6-N1	5.46	123.73	121.00
26	1H	636	G	O5'-P-OP2	5.46	117.25	110.70
26	14	1678	G	C4-C5-N7	5.46	112.98	110.80
26	1H	2078	C	O5'-P-OP2	5.46	117.25	110.70
27	16	6	C	C6-N1-C2	5.46	122.48	120.30
26	14	204	A	N1-C2-N3	5.46	132.03	129.30
26	14	1897	G	C8-N9-C1'	-5.46	119.91	127.00
1	13	286	G	C8-N9-C4	-5.45	104.22	106.40
26	1H	1626	G	N3-C2-N2	-5.45	116.08	119.90
48	J8	80	LEU	CA-CB-CG	5.45	127.84	115.30
1	1G	613	C	C6-N1-C2	-5.45	118.12	120.30
26	14	1648	C	C6-N1-C2	-5.45	118.12	120.30
1	13	278	G	N1-C6-O6	5.45	123.17	119.90
26	1H	2288	A	N1-C6-N6	5.45	121.87	118.60
38	98	107	ASP	CB-CG-OD1	-5.45	113.39	118.30
26	14	794	G	C8-N9-C4	5.45	108.58	106.40
26	1H	763	G	C4-C5-N7	-5.45	108.62	110.80
26	1H	1148	A	N7-C8-N9	-5.45	111.08	113.80
1	13	577	G	N1-C6-O6	5.45	123.17	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	1308	A	C5-C6-N6	5.45	128.06	123.70
26	1H	1382	G	C4-C5-N7	5.45	112.98	110.80
24	3L	57	C	C2-N1-C1'	5.45	124.79	118.80
26	14	1345	C	C6-N1-C2	-5.45	118.12	120.30
26	1H	211	A	N1-C6-N6	5.45	121.87	118.60
27	16	45	A	C8-N9-C4	-5.45	103.62	105.80
26	14	2624	G	N1-C6-O6	5.45	123.17	119.90
26	1H	2210	G	C8-N9-C4	-5.45	104.22	106.40
26	1H	954	G	C5-C6-O6	5.44	131.87	128.60
1	1G	1502	A	C6-C5-N7	-5.44	128.49	132.30
26	1H	301	G	OP1-P-OP2	5.44	127.77	119.60
26	1H	741	G	C8-N9-C4	-5.44	104.22	106.40
26	1H	2430	A	C5-N7-C8	-5.44	101.18	103.90
26	14	545	G	C6-C5-N7	5.44	133.67	130.40
26	14	2062	A	C8-N9-C4	5.44	107.98	105.80
26	14	2404	C	C6-N1-C2	5.44	122.48	120.30
26	1H	2304	G	C8-N9-C1'	5.44	134.07	127.00
26	14	140	A	C5-C6-N6	-5.44	119.35	123.70
1	1G	632	A	O4'-C1'-N9	5.44	112.55	108.20
26	1H	845	G	C4-N9-C1'	-5.44	119.43	126.50
1	13	975	A	O4'-C1'-N9	-5.43	103.85	108.20
26	14	1600	C	C5-C6-N1	-5.43	118.28	121.00
26	14	2620	C	C2-N1-C1'	5.43	124.78	118.80
1	13	481	G	C4-N9-C1'	5.43	133.56	126.50
1	13	1529	G	C8-N9-C4	-5.43	104.23	106.40
26	1H	634	C	O5'-P-OP2	-5.43	100.81	105.70
26	1H	1626	G	N3-C4-N9	-5.43	122.74	126.00
26	1H	2509	G	C6-N1-C2	-5.43	121.84	125.10
26	14	141	A	C5-N7-C8	-5.43	101.18	103.90
26	14	2022	U	O4'-C1'-N1	-5.43	103.86	108.20
1	13	484	G	P-O3'-C3'	5.43	126.22	119.70
1	13	974	A	N9-C4-C5	-5.43	103.63	105.80
26	1H	1554	A	O5'-P-OP2	-5.43	100.81	105.70
1	1G	1519	A	N9-C4-C5	5.43	107.97	105.80
26	14	120	U	N3-C2-O2	-5.43	118.40	122.20
26	1H	950	G	N1-C6-O6	-5.43	116.64	119.90
26	1H	1334	G	C6-C5-N7	-5.43	127.14	130.40
26	14	1266	G	C5-C6-N1	5.43	114.21	111.50
1	13	975	A	C4-C5-N7	5.43	113.41	110.70
26	1H	397	G	C2-N3-C4	-5.43	109.19	111.90
26	14	1379	A	N1-C6-N6	5.43	121.86	118.60
26	1H	856	C	C2-N1-C1'	5.42	124.77	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	1558	A	N1-C2-N3	5.42	132.01	129.30
26	14	2859	G	N3-C4-C5	-5.42	125.89	128.60
26	1H	464	U	C4-C5-C6	5.42	122.95	119.70
26	1H	2340	G	C8-N9-C4	5.42	108.57	106.40
22	1L	85	C	C6-N1-C1'	5.42	127.31	120.80
26	14	1379	A	C4-C5-N7	5.42	113.41	110.70
1	13	878	G	N3-C4-N9	5.42	129.25	126.00
26	14	155	C	N1-C2-O2	5.42	122.15	118.90
26	14	1698	A	C5-N7-C8	-5.42	101.19	103.90
26	1H	1955	U	O5'-P-OP1	-5.42	100.82	105.70
26	14	391	G	C5-C6-O6	-5.42	125.35	128.60
1	13	339	C	N1-C2-O2	-5.42	115.65	118.90
26	1H	617	G	O5'-P-OP1	5.42	117.20	110.70
26	1H	1347	G	N3-C4-N9	5.42	129.25	126.00
26	14	533	G	N3-C4-N9	5.42	129.25	126.00
26	14	1610	A	C6-C5-N7	-5.42	128.51	132.30
26	14	1774	C	C6-N1-C2	-5.42	118.13	120.30
26	1H	2590	A	O5'-P-OP1	-5.42	100.83	105.70
26	1H	162	U	C2-N1-C1'	5.41	124.20	117.70
26	14	2779	U	C2-N3-C4	-5.41	123.75	127.00
26	1H	2856	C	N3-C4-C5	-5.41	119.73	121.90
26	14	2254	C	C2-N1-C1'	-5.41	112.85	118.80
26	1H	141(A)	C	O5'-P-OP1	-5.41	100.83	105.70
26	1H	2377	A	C2-N3-C4	-5.41	107.89	110.60
26	14	689	A	O5'-P-OP1	5.41	117.19	110.70
26	14	1383	C	N3-C4-N4	5.41	121.79	118.00
26	1H	859	G	C4-N9-C1'	-5.41	119.47	126.50
26	1H	1779	U	O5'-P-OP1	-5.41	100.83	105.70
1	1G	1502	A	C4-C5-N7	5.41	113.40	110.70
26	14	2709	G	C5-C6-O6	-5.41	125.36	128.60
26	14	2826	A	N1-C6-N6	-5.41	115.36	118.60
26	14	210	C	C5-C6-N1	-5.41	118.30	121.00
26	14	915	C	N1-C2-O2	5.41	122.14	118.90
26	14	945	A	C5-C6-N6	-5.41	119.37	123.70
26	1H	464	U	N1-C2-N3	5.41	118.14	114.90
26	14	805	G	O5'-P-OP1	-5.41	100.83	105.70
26	14	2389	G	N7-C8-N9	5.41	115.80	113.10
26	1H	428	A	OP1-P-O3'	5.40	117.08	105.20
26	1H	2525	G	N9-C4-C5	-5.40	103.24	105.40
1	13	549	C	C6-N1-C1'	5.40	127.28	120.80
1	13	938	A	N1-C6-N6	5.40	121.84	118.60
1	13	1220	G	C5-C6-O6	-5.40	125.36	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
23	2K	1	G	C4-N9-C1'	5.40	133.52	126.50
26	1H	2724	C	C2-N1-C1'	-5.40	112.86	118.80
26	14	178	G	N1-C6-O6	5.40	123.14	119.90
1	13	452	A	N7-C8-N9	-5.40	111.10	113.80
26	1H	824	A	C2-N3-C4	-5.40	107.90	110.60
26	14	1999	C	C5-C4-N4	-5.40	116.42	120.20
26	14	2440	C	C2-N3-C4	5.40	122.60	119.90
26	14	2490	G	N9-C4-C5	5.40	107.56	105.40
22	1K	87	A	O4'-C1'-N9	5.40	112.52	108.20
26	1H	1391	U	N3-C2-O2	-5.40	118.42	122.20
26	1H	2296	U	N3-C4-O4	5.40	123.18	119.40
1	1G	390	C	C2-N1-C1'	-5.40	112.86	118.80
26	14	1656	C	OP2-P-O3'	5.40	117.08	105.20
26	14	83	G	N1-C6-O6	5.40	123.14	119.90
23	2K	31	C	N3-C2-O2	-5.39	118.12	121.90
26	1H	530	G	C5-C6-O6	5.39	131.84	128.60
26	1H	1301	A	C5-C6-N1	-5.39	115.00	117.70
1	1G	353	A	OP2-P-O3'	5.39	117.07	105.20
26	14	2168	G	C4-N9-C1'	5.39	133.51	126.50
26	1H	2870	C	C6-N1-C2	-5.39	118.14	120.30
1	1G	481	G	N3-C4-N9	5.39	129.24	126.00
26	14	391	G	C4-C5-C6	5.39	122.04	118.80
26	14	1790	C	C5-C4-N4	-5.39	116.43	120.20
1	13	227	G	C8-N9-C4	5.39	108.56	106.40
26	14	1698	A	O4'-C1'-N9	5.39	112.51	108.20
26	1H	1773	A	C4-C5-C6	5.39	119.69	117.00
26	14	1342	A	C4-N9-C1'	5.39	136.00	126.30
1	13	916	G	C5-C6-N1	5.39	114.19	111.50
26	1H	1695	G	N9-C4-C5	-5.39	103.25	105.40
26	14	784	A	C2-N3-C4	-5.39	107.91	110.60
26	14	1225	C	C2-N1-C1'	-5.39	112.88	118.80
26	14	2419	U	OP1-P-O3'	5.39	117.05	105.20
26	1H	117	G	C5-C6-N1	5.38	114.19	111.50
26	1H	201	C	C5-C6-N1	-5.38	118.31	121.00
1	1G	106	C	C6-N1-C2	-5.38	118.15	120.30
1	13	1512	U	C5-C6-N1	-5.38	120.01	122.70
26	1H	1178	C	P-O3'-C3'	5.38	126.16	119.70
26	1H	1771	C	N1-C2-O2	-5.38	115.67	118.90
26	1H	1952	A	OP1-P-O3'	5.38	117.04	105.20
1	13	911	U	C4-C5-C6	5.38	122.93	119.70
26	1H	847	U	N3-C2-O2	-5.38	118.43	122.20
26	1H	1379	A	C6-C5-N7	-5.38	128.53	132.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	1676	A	OP1-P-O3'	5.38	117.04	105.20
26	1H	1952	A	O5'-P-OP2	-5.38	100.86	105.70
26	1H	2449	U	C5-C6-N1	5.38	125.39	122.70
27	16	12	C	C6-N1-C2	5.38	122.45	120.30
26	14	2473	U	C2-N1-C1'	5.38	124.16	117.70
26	1H	650	C	N1-C2-O2	-5.38	115.67	118.90
26	14	2829	C	N3-C4-C5	5.38	124.05	121.90
26	1H	954	G	N9-C4-C5	5.38	107.55	105.40
26	1H	1202	C	N3-C4-C5	-5.38	119.75	121.90
26	1H	1210	A	N9-C4-C5	-5.38	103.65	105.80
1	1G	305	G	N1-C6-O6	-5.38	116.67	119.90
26	14	2253	G	C5-C6-N1	5.38	114.19	111.50
26	1H	379	G	N9-C4-C5	-5.38	103.25	105.40
26	1H	2567	G	C5-C6-O6	-5.38	125.37	128.60
26	14	1376	C	O4'-C1'-N1	5.38	112.50	108.20
26	1H	785	G	N3-C4-C5	-5.38	125.91	128.60
26	1H	1294	U	N1-C2-O2	-5.38	119.04	122.80
26	14	2526	G	N3-C4-C5	5.38	131.29	128.60
26	1H	2366	A	C8-N9-C4	-5.37	103.65	105.80
26	1H	2856	C	N3-C2-O2	-5.37	118.14	121.90
26	14	127	A	N1-C6-N6	5.37	121.83	118.60
26	1H	62	C	C6-N1-C2	5.37	122.45	120.30
1	1G	1139	G	C8-N9-C4	5.37	108.55	106.40
1	1G	1281	U	C6-N1-C2	-5.37	117.78	121.00
26	14	2511	U	C5-C4-O4	-5.37	122.68	125.90
26	14	2584	U	C6-N1-C1'	-5.37	113.68	121.20
1	13	561	U	C5-C4-O4	-5.37	122.68	125.90
1	13	700	G	N9-C4-C5	-5.37	103.25	105.40
26	1H	82	G	C4-C5-C6	5.37	122.02	118.80
26	14	794	G	N1-C2-N3	5.37	127.12	123.90
26	1H	1401	G	C8-N9-C4	-5.37	104.25	106.40
1	1G	536	C	N3-C2-O2	-5.37	118.14	121.90
26	1H	189	G	OP2-P-O3'	5.37	117.00	105.20
26	1H	1309	G	C8-N9-C4	5.37	108.55	106.40
26	1H	2468	G	C8-N9-C4	-5.37	104.25	106.40
1	1G	115	G	P-O3'-C3'	5.37	126.14	119.70
26	14	1902	C	C5-C4-N4	5.37	123.96	120.20
1	13	517	G	C5-C6-O6	-5.36	125.38	128.60
26	1H	263	C	C2-N3-C4	-5.36	117.22	119.90
26	1H	452	G	N3-C4-C5	-5.36	125.92	128.60
26	1H	1032	A	C8-N9-C4	5.36	107.94	105.80
26	14	686	G	C5-C6-O6	-5.36	125.38	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	113	G	C5-C6-O6	-5.36	125.38	128.60
1	13	529	G	N1-C6-O6	5.36	123.11	119.90
26	1H	622	G	C8-N9-C1'	-5.36	120.03	127.00
26	14	1555	G	C8-N9-C1'	-5.36	120.03	127.00
26	1H	389	G	C8-N9-C4	5.36	108.54	106.40
26	1H	671	C	C6-N1-C2	5.36	122.44	120.30
1	1G	15	G	N3-C2-N2	5.36	123.65	119.90
26	14	2276	G	O5'-P-OP1	-5.36	100.88	105.70
23	2L	27	G	C5-C6-O6	-5.35	125.39	128.60
26	1H	446	G	C4-C5-N7	5.35	112.94	110.80
26	1H	504	U	N1-C2-O2	5.35	126.55	122.80
26	1H	2240	C	N3-C4-C5	-5.35	119.76	121.90
1	13	23	C	C5-C6-N1	5.35	123.67	121.00
26	1H	486	C	N1-C2-O2	-5.35	115.69	118.90
26	1H	1610	A	C8-N9-C4	-5.35	103.66	105.80
26	14	2002	G	N1-C6-O6	5.35	123.11	119.90
26	14	2713	A	C6-C5-N7	-5.35	128.56	132.30
1	13	749	C	C2-N1-C1'	5.35	124.68	118.80
1	13	773	G	C6-C5-N7	-5.35	127.19	130.40
26	1H	684	G	C5-C6-N1	5.35	114.17	111.50
26	1H	859	G	C8-N9-C4	5.35	108.54	106.40
26	1H	1601	G	C8-N9-C4	5.35	108.54	106.40
26	1H	2504	U	C5-C4-O4	5.35	129.11	125.90
26	1H	2568	C	N3-C4-C5	5.35	124.04	121.90
26	14	465	G	C8-N9-C4	-5.35	104.26	106.40
26	14	2477	C	C6-N1-C1'	-5.35	114.38	120.80
1	13	1367	C	C6-N1-C2	-5.34	118.16	120.30
26	1H	801	G	N9-C4-C5	5.34	107.54	105.40
26	1H	810	U	C5-C4-O4	-5.34	122.69	125.90
26	1H	1190	G	C5-N7-C8	-5.34	101.63	104.30
26	1H	1364	G	N9-C4-C5	-5.34	103.26	105.40
26	1H	1998	G	C8-N9-C1'	-5.34	120.05	127.00
26	14	275	G	N1-C6-O6	5.34	123.11	119.90
26	14	1594	G	N7-C8-N9	5.34	115.77	113.10
26	14	1641	A	C6-N1-C2	-5.34	115.39	118.60
26	1H	1915	U	N1-C2-O2	5.34	126.54	122.80
26	1H	2056	G	N3-C4-C5	-5.34	125.93	128.60
1	13	893	C	N1-C2-O2	5.34	122.11	118.90
26	1H	609	A	N9-C4-C5	-5.34	103.66	105.80
26	1H	1278	A	N1-C2-N3	5.34	131.97	129.30
26	1H	1364	G	C6-C5-N7	-5.34	127.20	130.40
26	1H	1520	U	N1-C2-N3	5.34	118.10	114.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	1G	266	G	O4'-C1'-N9	-5.34	103.93	108.20
1	1G	345	C	N1-C2-O2	5.34	122.11	118.90
1	1G	827	U	N1-C2-O2	5.34	126.54	122.80
23	2L	16	U	C5-C6-N1	5.34	125.37	122.70
42	95	49	THR	C-N-CD	5.34	139.62	128.40
26	1H	961	C	O5'-P-OP1	-5.34	100.89	105.70
26	1H	2367	G	C4-C5-N7	5.34	112.94	110.80
26	14	263	C	N1-C2-O2	5.34	122.10	118.90
26	14	275	G	C8-N9-C1'	-5.34	120.06	127.00
26	1H	2830	G	N7-C8-N9	5.34	115.77	113.10
26	1H	913	U	O5'-P-OP2	-5.34	100.90	105.70
26	1H	1396	U	OP1-P-OP2	5.34	127.60	119.60
1	1G	579	G	C6-C5-N7	-5.34	127.20	130.40
26	1H	196	A	N1-C6-N6	5.33	121.80	118.60
26	1H	512	G	N3-C4-C5	5.33	131.27	128.60
26	1H	862	G	N3-C4-C5	-5.33	125.93	128.60
26	1H	1675	C	C6-N1-C2	-5.33	118.17	120.30
26	14	2490	G	N3-C4-N9	-5.33	122.80	126.00
26	1H	2595	G	C6-C5-N7	-5.33	127.20	130.40
26	1H	2827	C	C6-N1-C2	5.33	122.43	120.30
1	1G	306	G	N1-C6-O6	5.33	123.10	119.90
26	14	488	G	C5-C6-O6	-5.33	125.40	128.60
26	14	848	G	N3-C4-N9	5.33	129.20	126.00
26	14	1596	A	C5-C6-N6	5.33	127.97	123.70
26	14	1843	C	C2-N3-C4	-5.33	117.23	119.90
26	1H	1513	C	C6-N1-C2	-5.33	118.17	120.30
1	1G	733	A	O4'-C1'-N9	5.33	112.47	108.20
26	14	1695	G	N1-C2-N2	-5.33	111.40	116.20
26	14	2420	C	N1-C2-O2	-5.33	115.70	118.90
26	1H	2082	A	O5'-P-OP1	5.33	117.10	110.70
26	14	1332	G	C5-C6-O6	5.33	131.80	128.60
26	1H	840	C	N3-C4-N4	-5.33	114.27	118.00
1	1G	1390	U	C5-C4-O4	5.33	129.10	125.90
26	14	1382	G	C8-N9-C4	5.33	108.53	106.40
26	14	2607	G	O5'-P-OP2	-5.33	100.90	105.70
26	14	530	G	C5-N7-C8	-5.33	101.64	104.30
26	1H	9	U	C2-N1-C1'	5.32	124.09	117.70
26	1H	1327	C	N1-C2-O2	-5.32	115.70	118.90
26	1H	2606	C	C2-N3-C4	-5.32	117.24	119.90
26	1H	2877	G	C5-C6-N1	-5.32	108.84	111.50
1	1G	320	C	C2-N1-C1'	-5.32	112.94	118.80
1	1G	1348	U	C5-C4-O4	5.32	129.09	125.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	571	A	N1-C6-N6	5.32	121.80	118.60
26	14	1970	A	N9-C4-C5	-5.32	103.67	105.80
26	14	2827	C	C6-N1-C2	5.32	122.43	120.30
26	1H	752	A	N1-C2-N3	5.32	131.96	129.30
26	1H	1690	A	O5'-P-OP1	-5.32	100.91	105.70
26	1H	1786	A	N3-C4-C5	5.32	130.53	126.80
26	14	2463	C	C2-N1-C1'	-5.32	112.95	118.80
1	13	667	G	N3-C2-N2	-5.32	116.18	119.90
26	1H	213	A	N9-C4-C5	-5.32	103.67	105.80
26	1H	1364	G	C4-C5-N7	5.32	112.93	110.80
1	1G	1442	G	N3-C4-N9	-5.32	122.81	126.00
26	14	467	G	C8-N9-C4	5.32	108.53	106.40
26	14	488	G	N9-C4-C5	-5.32	103.27	105.40
26	14	1613	G	C5-N7-C8	5.32	106.96	104.30
26	14	188	G	OP1-P-OP2	5.32	127.58	119.60
26	14	2437	U	N1-C2-N3	5.32	118.09	114.90
1	13	405	U	C5-C6-N1	5.32	125.36	122.70
26	1H	666	G	C5-C6-O6	-5.32	125.41	128.60
26	1H	2686	G	C4-N9-C1'	5.32	133.41	126.50
26	1H	2867	G	N3-C4-N9	-5.32	122.81	126.00
26	14	2534	A	N1-C6-N6	5.32	121.79	118.60
26	14	1489	U	C2-N1-C1'	-5.31	111.32	117.70
23	2K	22	G	C4-N9-C1'	5.31	133.41	126.50
26	14	1308	A	C8-N9-C4	-5.31	103.67	105.80
26	14	1613	G	OP1-P-O3'	5.31	116.89	105.20
26	1H	946	G	N9-C4-C5	-5.31	103.28	105.40
26	1H	2228	G	C4-N9-C1'	5.31	133.40	126.50
1	1G	632	A	P-O3'-C3'	5.31	126.07	119.70
26	14	761	A	C5-N7-C8	-5.31	101.25	103.90
26	14	2596	U	OP1-P-OP2	5.31	127.56	119.60
27	1J	89	G	C4-N9-C1'	5.31	133.40	126.50
26	1H	1438	U	C2-N1-C1'	5.31	124.07	117.70
26	14	1835	G	N3-C4-N9	5.31	129.18	126.00
26	1H	770	G	N1-C6-O6	5.30	123.08	119.90
1	1G	690	G	O4'-C1'-N9	5.30	112.44	108.20
26	1H	380	U	C4-C5-C6	5.30	122.88	119.70
26	1H	1604	C	OP2-P-O3'	5.30	116.87	105.20
26	1H	2320	A	O4'-C1'-N9	5.30	112.44	108.20
1	1G	180	U	C5-C6-N1	5.30	125.35	122.70
26	1H	452	G	N1-C6-O6	-5.30	116.72	119.90
26	1H	1888	G	N3-C4-N9	5.30	129.18	126.00
26	14	2490	G	N1-C6-O6	-5.30	116.72	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	1G	332	G	C8-N9-C4	5.30	108.52	106.40
26	1H	383	U	C2-N1-C1'	-5.30	111.34	117.70
26	1H	1613	G	N3-C2-N2	5.30	123.61	119.90
26	1H	1975	G	N9-C4-C5	-5.30	103.28	105.40
26	1H	2439	A	OP2-P-O3'	-5.29	93.55	105.20
26	1H	2645	G	C5-C6-O6	5.29	131.78	128.60
26	1H	2782	G	C6-C5-N7	-5.29	127.22	130.40
1	13	186(A)	C	C2-N1-C1'	5.29	124.62	118.80
1	13	428	G	C8-N9-C1'	5.29	133.88	127.00
1	13	904	C	C2-N3-C4	-5.29	117.25	119.90
26	1H	845	G	C4-C5-N7	5.29	112.92	110.80
26	14	2622	C	N1-C2-O2	-5.29	115.72	118.90
26	14	577	G	C4-N9-C1'	5.29	133.38	126.50
26	14	784	A	N3-C4-N9	-5.29	123.17	127.40
26	1H	69	C	N3-C2-O2	-5.29	118.20	121.90
26	1H	2329	G	C4-N9-C1'	-5.29	119.63	126.50
26	14	1327	C	C6-N1-C1'	5.29	127.15	120.80
1	1G	793	U	C2-N1-C1'	-5.29	111.36	117.70
1	1G	1468	A	N1-C6-N6	5.29	121.77	118.60
26	14	273(D)	C	O5'-P-OP2	-5.29	100.94	105.70
26	1H	1695	G	C8-N9-C1'	-5.29	120.13	127.00
26	1H	2007	C	C6-N1-C2	-5.29	118.19	120.30
26	1H	2211	G	O4'-C1'-N9	-5.28	103.97	108.20
22	1L	21	A	P-O3'-C3'	5.28	126.04	119.70
23	2L	33	U	C2-N1-C1'	5.28	124.04	117.70
26	14	845	G	C2-N3-C4	-5.28	109.26	111.90
26	14	1999	C	N3-C4-C5	5.28	124.01	121.90
26	1H	1630(A)	C	C6-N1-C2	-5.28	118.19	120.30
26	1H	1786	A	N1-C6-N6	5.28	121.77	118.60
1	1G	909	A	N1-C6-N6	5.28	121.77	118.60
26	1H	271(B)	G	C4-N9-C1'	5.28	133.37	126.50
26	14	141(A)	C	C6-N1-C2	-5.28	118.19	120.30
1	13	767	A	O5'-P-OP1	-5.28	100.95	105.70
26	1H	528	A	C6-N1-C2	5.28	121.77	118.60
1	13	1513	A	N9-C4-C5	-5.28	103.69	105.80
26	1H	404	C	P-O3'-C3'	5.28	126.03	119.70
26	1H	446	G	C5-C6-O6	-5.28	125.43	128.60
26	1H	1698	A	C4-C5-N7	5.28	113.34	110.70
1	1G	963	G	N3-C2-N2	5.28	123.59	119.90
26	14	457	A	N7-C8-N9	5.28	116.44	113.80
26	14	2624	G	N7-C8-N9	-5.28	110.46	113.10
27	1J	89	G	N3-C4-N9	5.28	129.16	126.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	1676	A	C2-N3-C4	-5.27	107.96	110.60
26	1H	1956	U	N1-C2-O2	-5.27	119.11	122.80
1	13	61	G	C4-C5-N7	-5.27	108.69	110.80
26	1H	140	A	C5-C6-N6	-5.27	119.48	123.70
26	1H	678	C	C2-N3-C4	-5.27	117.27	119.90
26	1H	1299	G	C5-N7-C8	-5.27	101.66	104.30
26	14	1991	U	N3-C4-O4	-5.27	115.71	119.40
26	14	2043	C	C6-N1-C2	-5.27	118.19	120.30
1	13	538	G	C8-N9-C4	5.27	108.51	106.40
22	1K	18	G	N3-C4-C5	5.27	131.24	128.60
26	1H	2568	C	C5-C4-N4	-5.27	116.51	120.20
1	1G	890	G	C4-C5-N7	-5.27	108.69	110.80
26	14	187	G	C8-N9-C4	5.27	108.51	106.40
26	14	621	A	C6-C5-N7	-5.27	128.61	132.30
26	14	1462	C	C6-N1-C2	-5.27	118.19	120.30
26	14	2027	G	N1-C2-N3	5.27	127.06	123.90
26	1H	484	C	C6-N1-C2	-5.27	118.19	120.30
26	1H	1899	G	N1-C2-N2	5.27	120.94	116.20
1	1G	322	C	C5-C4-N4	-5.27	116.51	120.20
26	14	2313	C	C6-N1-C2	-5.27	118.19	120.30
1	13	36	C	C6-N1-C2	-5.26	118.19	120.30
1	13	542	G	N3-C2-N2	-5.26	116.21	119.90
1	1G	500	G	C6-C5-N7	-5.26	127.24	130.40
26	14	1342	A	C5-C6-N1	-5.26	115.07	117.70
26	14	2027	G	N3-C4-C5	-5.26	125.97	128.60
26	1H	141	A	C8-N9-C4	-5.26	103.69	105.80
26	1H	1325	G	N3-C4-C5	-5.26	125.97	128.60
26	1H	1790	C	C5-C6-N1	-5.26	118.37	121.00
26	1H	1799	G	OP1-P-O3'	5.26	116.78	105.20
26	14	198	C	C6-N1-C2	-5.26	118.19	120.30
26	14	850	C	C6-N1-C2	-5.26	118.19	120.30
26	14	2755	C	N1-C2-O2	5.26	122.06	118.90
26	1H	2207	C	C6-N1-C2	-5.26	118.19	120.30
26	1H	2490	G	C2-N3-C4	-5.26	109.27	111.90
26	14	1823	G	N1-C6-O6	5.26	123.06	119.90
26	14	2609	U	N3-C2-O2	-5.26	118.52	122.20
1	13	1417	G	C8-N9-C4	-5.26	104.30	106.40
26	14	647	G	C4-N9-C1'	5.26	133.34	126.50
26	14	2352	A	O5'-P-OP1	-5.26	100.97	105.70
26	14	2593	U	N3-C4-C5	-5.26	111.44	114.60
1	1G	897	C	N3-C2-O2	5.26	125.58	121.90
26	14	2442	C	N1-C2-O2	5.26	122.06	118.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	13	1052	U	N1-C2-O2	5.26	126.48	122.80
26	14	57	C	N3-C2-O2	5.26	125.58	121.90
26	14	211	A	N9-C4-C5	-5.26	103.70	105.80
26	14	1204	A	C5-N7-C8	-5.26	101.27	103.90
26	14	1602	U	N3-C4-C5	-5.26	111.45	114.60
26	14	2275	C	C6-N1-C2	-5.26	118.20	120.30
26	1H	71	A	N7-C8-N9	5.25	116.43	113.80
26	1H	481	G	O4'-C1'-N9	5.25	112.40	108.20
26	1H	664	C	C4-C5-C6	5.25	120.03	117.40
26	14	808	G	O5'-P-OP2	-5.25	100.97	105.70
1	13	561	U	N3-C2-O2	5.25	125.88	122.20
26	14	454	A	C8-N9-C4	5.25	107.90	105.80
26	14	1817	G	N1-C6-O6	5.25	123.05	119.90
26	1H	1809	A	N1-C6-N6	5.25	121.75	118.60
26	1H	1879	C	C6-N1-C2	-5.25	118.20	120.30
26	1H	2084	C	C5-C6-N1	-5.25	118.38	121.00
26	1H	2518	A	C8-N9-C4	-5.25	103.70	105.80
26	1H	2523	G	OP2-P-O3'	5.25	116.75	105.20
1	1G	414	A	O5'-P-OP2	-5.25	100.97	105.70
26	14	1275	A	C5-C6-N6	-5.25	119.50	123.70
22	1L	85	C	O4'-C1'-N1	5.25	112.40	108.20
26	14	801	G	C4-N9-C1'	-5.25	119.67	126.50
26	1H	1211	U	C6-N1-C2	5.25	124.15	121.00
26	1H	1969	A	O5'-P-OP1	-5.25	100.98	105.70
27	16	112	G	N3-C4-C5	5.25	131.22	128.60
26	14	444	C	C6-N1-C2	5.25	122.40	120.30
26	14	2769	C	O4'-C1'-N1	5.25	112.40	108.20
1	13	245	C	C6-N1-C2	-5.25	118.20	120.30
26	1H	1975	G	C8-N9-C1'	-5.25	120.18	127.00
26	1H	2015	A	OP1-P-O3'	5.25	116.74	105.20
1	1G	799	G	OP2-P-O3'	5.25	116.74	105.20
26	14	2715	C	C6-N1-C2	5.25	122.40	120.30
1	1G	380	G	N1-C6-O6	5.25	123.05	119.90
23	2L	59	U	C6-N1-C2	-5.25	117.85	121.00
26	1H	1215	G	OP1-P-O3'	5.24	116.73	105.20
26	1H	1669	A	C4-N9-C1'	5.24	135.74	126.30
26	1H	2454	G	N1-C6-O6	-5.24	116.75	119.90
1	1G	579	G	N7-C8-N9	5.24	115.72	113.10
26	14	12	U	C2-N1-C1'	5.24	123.99	117.70
26	14	391	G	N7-C8-N9	5.24	115.72	113.10
26	1H	2233	U	N1-C2-N3	5.24	118.05	114.90
1	1G	579	G	N1-C6-O6	5.24	123.05	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	2002	G	C4-C5-N7	5.24	112.90	110.80
26	1H	690	G	C8-N9-C4	5.24	108.50	106.40
26	14	143	C	C6-N1-C2	-5.24	118.20	120.30
26	14	1422	G	N3-C4-C5	5.24	131.22	128.60
26	14	1637	A	C4-C5-C6	5.24	119.62	117.00
26	14	2389	G	C5-N7-C8	-5.24	101.68	104.30
26	1H	1968	G	C4-C5-N7	5.24	112.89	110.80
26	14	2072	G	C4-C5-N7	5.24	112.89	110.80
1	13	636	U	C5-C6-N1	5.24	125.32	122.70
26	14	1604	C	C6-N1-C2	-5.24	118.20	120.30
26	1H	683	C	C5-C4-N4	-5.24	116.53	120.20
26	1H	765	G	C5-C6-O6	-5.24	125.46	128.60
1	1G	275	G	N3-C4-C5	-5.24	125.98	128.60
26	14	2377	A	C8-N9-C4	5.24	107.89	105.80
26	1H	1253	A	O5'-P-OP1	-5.23	100.99	105.70
26	1H	1897	G	C5-C6-N1	-5.23	108.88	111.50
26	1H	2719	G	C8-N9-C4	-5.23	104.31	106.40
26	14	962	G	N3-C4-N9	-5.23	122.86	126.00
26	14	2508	G	C5-N7-C8	-5.23	101.68	104.30
22	1L	31	C	C6-N1-C2	-5.23	118.21	120.30
26	14	2382	G	N3-C4-N9	5.23	129.14	126.00
1	13	247	G	N3-C4-N9	5.23	129.14	126.00
26	1H	1835	G	N3-C2-N2	5.23	123.56	119.90
1	1G	982	U	N3-C4-C5	5.23	117.74	114.60
26	14	1193	G	N9-C4-C5	-5.23	103.31	105.40
1	13	544	G	C6-C5-N7	-5.23	127.26	130.40
1	13	449	C	C2-N1-C1'	5.23	124.55	118.80
26	1H	514	A	C8-N9-C4	5.23	107.89	105.80
26	14	1800	C	N3-C4-C5	-5.23	119.81	121.90
26	14	1970	A	C5-C6-N6	-5.23	119.52	123.70
26	14	2506	U	N3-C4-O4	5.23	123.06	119.40
26	1H	676	A	N1-C2-N3	5.23	131.91	129.30
1	13	1027	C	P-O3'-C3'	5.22	125.97	119.70
26	1H	1141	U	O4'-C1'-N1	5.22	112.38	108.20
1	1G	351	G	OP2-P-O3'	5.22	116.69	105.20
26	14	1342	A	C6-C5-N7	-5.22	128.64	132.30
1	13	108	G	C5-N7-C8	-5.22	101.69	104.30
1	13	1381	U	N3-C2-O2	-5.22	118.54	122.20
26	1H	2263	C	C6-N1-C2	-5.22	118.21	120.30
1	1G	748	C	P-O3'-C3'	5.22	125.97	119.70
3	2E	204	LEU	CA-CB-CG	-5.22	103.29	115.30
23	2L	9	G	C4-N9-C1'	5.22	133.29	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	13	690	G	N1-C2-N3	5.22	127.03	123.90
26	1H	196	A	O5'-P-OP2	-5.22	101.00	105.70
26	1H	985	C	OP2-P-O3'	5.22	116.68	105.20
26	1H	2567	G	N1-C6-O6	5.22	123.03	119.90
26	14	2417	C	O5'-P-OP2	-5.22	101.00	105.70
1	13	773	G	C8-N9-C1'	-5.22	120.22	127.00
26	1H	1614	A	C4-C5-N7	5.22	113.31	110.70
26	1H	1669	A	C8-N9-C4	-5.22	103.71	105.80
26	1H	2331	G	C8-N9-C1'	-5.22	120.22	127.00
26	1H	2462	U	C6-N1-C2	-5.22	117.87	121.00
26	1H	2609	U	C4-C5-C6	5.22	122.83	119.70
26	14	641	C	C6-N1-C2	5.22	122.39	120.30
1	13	1504	G	O5'-P-OP1	-5.21	101.01	105.70
26	1H	309	G	C6-C5-N7	-5.21	127.27	130.40
26	1H	621	A	C5-C6-N1	-5.21	115.09	117.70
26	1H	838	C	C4-C5-C6	5.21	120.01	117.40
1	1G	605	U	C6-N1-C2	-5.21	117.87	121.00
26	14	329	G	O5'-P-OP2	-5.21	101.01	105.70
26	14	683	C	C5-C4-N4	-5.21	116.55	120.20
26	14	929	G	C4-C5-C6	5.21	121.93	118.80
1	1G	1128	C	C6-N1-C2	-5.21	118.22	120.30
1	1G	1498	U	C2-N1-C1'	5.21	123.96	117.70
26	14	2362	G	C8-N9-C4	5.21	108.48	106.40
26	1H	121	G	N9-C4-C5	-5.21	103.32	105.40
26	1H	741	G	C6-N1-C2	-5.21	121.97	125.10
26	14	659	C	C2-N3-C4	-5.21	117.29	119.90
26	14	1411	C	N1-C2-O2	5.21	122.03	118.90
26	14	2323	G	N1-C6-O6	5.21	123.03	119.90
26	1H	845	G	C8-N9-C1'	5.21	133.77	127.00
26	1H	871	U	C5-C6-N1	5.21	125.31	122.70
26	1H	2064	C	C4-C5-C6	5.21	120.00	117.40
26	1H	2416	C	O5'-P-OP2	-5.21	101.01	105.70
1	13	1483	A	O5'-P-OP1	-5.21	101.01	105.70
26	1H	467	G	N1-C6-O6	5.21	123.02	119.90
1	1G	275	G	C8-N9-C1'	-5.21	120.23	127.00
1	1G	324	G	C4-N9-C1'	-5.21	119.73	126.50
26	14	432	A	N1-C6-N6	5.21	121.72	118.60
26	14	2573	C	N3-C4-N4	5.21	121.64	118.00
26	1H	265	A	C4-C5-N7	5.21	113.30	110.70
26	1H	1311	G	C8-N9-C4	5.21	108.48	106.40
26	1H	2329	G	N3-C4-C5	5.21	131.20	128.60
26	1H	2383	G	C4-C5-C6	5.21	121.92	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
27	16	101	A	C8-N9-C4	5.21	107.88	105.80
26	14	74	A	N7-C8-N9	5.21	116.40	113.80
1	13	752	G	C8-N9-C1'	-5.20	120.23	127.00
26	1H	298	G	C5-C6-O6	-5.20	125.48	128.60
23	2L	35	G	OP1-P-O3'	5.20	116.65	105.20
26	14	2776	A	C8-N9-C4	-5.20	103.72	105.80
1	13	1279	A	N7-C8-N9	5.20	116.40	113.80
26	1H	953	A	N9-C4-C5	-5.20	103.72	105.80
26	14	1167	U	C2-N1-C1'	-5.20	111.46	117.70
1	13	919	A	N1-C6-N6	-5.20	115.48	118.60
26	1H	465	G	C5-C6-N1	-5.20	108.90	111.50
26	14	700	G	C6-C5-N7	-5.20	127.28	130.40
26	14	1698	A	C5-C6-N6	-5.20	119.54	123.70
26	14	1823	G	C5-C6-N1	-5.20	108.90	111.50
26	14	2518	A	C6-C5-N7	-5.20	128.66	132.30
26	14	2542	A	C8-N9-C4	5.20	107.88	105.80
26	1H	679	C	N3-C4-C5	5.20	123.98	121.90
26	1H	860	U	N1-C2-O2	5.20	126.44	122.80
26	1H	1006	C	N3-C2-O2	5.20	125.54	121.90
26	1H	2259	G	C5-C6-O6	-5.20	125.48	128.60
26	1H	2360	A	C2-N3-C4	-5.20	108.00	110.60
26	1H	2473	U	C2-N1-C1'	5.20	123.94	117.70
26	14	382	G	OP1-P-O3'	5.20	116.63	105.20
26	1H	215	G	N1-C6-O6	5.20	123.02	119.90
26	1H	829	A	C5-C6-N1	-5.20	115.10	117.70
26	14	863	A	O5'-P-OP2	-5.20	101.02	105.70
26	14	961	C	O4'-C1'-N1	5.20	112.36	108.20
26	14	2392	A	C8-N9-C4	-5.20	103.72	105.80
26	14	2619	C	N3-C4-C5	5.20	123.98	121.90
26	1H	1785	A	OP2-P-O3'	5.19	116.63	105.20
26	1H	2571	C	N3-C2-O2	-5.19	118.27	121.90
26	14	2099	U	C2-N1-C1'	5.19	123.93	117.70
1	13	776	G	N1-C6-O6	5.19	123.02	119.90
2	1E	187	LEU	CA-CB-CG	5.19	127.24	115.30
26	1H	395	U	N3-C2-O2	-5.19	118.57	122.20
46	H8	76	LEU	CA-CB-CG	5.19	127.24	115.30
26	14	1669	A	O5'-P-OP1	-5.19	101.03	105.70
26	1H	1543	A	C2-N3-C4	-5.19	108.00	110.60
27	16	29	A	C8-N9-C4	-5.19	103.72	105.80
26	14	1386	C	N3-C2-O2	5.19	125.53	121.90
26	14	2346	A	C4-N9-C1'	5.19	135.64	126.30
26	1H	1448	G	C5-C6-O6	-5.19	125.49	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	928	G	C5-C6-O6	-5.19	125.49	128.60
26	1H	1322	A	OP2-P-O3'	5.19	116.61	105.20
1	1G	218	C	C6-N1-C2	-5.19	118.22	120.30
1	1G	275	G	C4-N9-C1'	5.19	133.25	126.50
1	1G	1054	C	O5'-P-OP1	-5.19	101.03	105.70
26	14	2452	C	C5-C4-N4	-5.19	116.57	120.20
26	14	2624	G	C8-N9-C4	5.19	108.47	106.40
26	1H	107	C	N1-C2-O2	-5.19	115.79	118.90
26	1H	1400	G	C8-N9-C4	-5.19	104.33	106.40
26	1H	1636	C	C6-N1-C2	-5.19	118.23	120.30
26	14	332	A	OP2-P-O3'	5.19	116.61	105.20
26	1H	785	G	N9-C4-C5	5.18	107.47	105.40
26	1H	859	G	C2-N3-C4	-5.18	109.31	111.90
26	1H	1457	A	N1-C6-N6	5.18	121.71	118.60
26	14	784	A	P-O3'-C3'	5.18	125.92	119.70
26	1H	1022	G	P-O3'-C3'	5.18	125.92	119.70
26	1H	1963	U	N3-C2-O2	-5.18	118.57	122.20
27	16	24	G	C4-N9-C1'	5.18	133.24	126.50
1	13	50	A	P-O3'-C3'	5.18	125.92	119.70
26	1H	681	G	N9-C4-C5	-5.18	103.33	105.40
26	1H	1664	A	N1-C6-N6	-5.18	115.49	118.60
26	14	2702	U	N1-C2-N3	-5.18	111.79	114.90
1	13	625	G	C8-N9-C4	-5.18	104.33	106.40
26	1H	671	C	C5-C6-N1	-5.18	118.41	121.00
26	1H	1698	A	N9-C1'-C2'	5.18	120.73	114.00
26	1H	2234	G	C2-N3-C4	-5.18	109.31	111.90
26	1H	2339	G	C8-N9-C4	5.18	108.47	106.40
1	1G	20	U	C6-N1-C2	5.18	124.11	121.00
26	14	1342	A	N9-C1'-C2'	5.18	120.73	114.00
1	13	12	U	N3-C4-C5	-5.18	111.49	114.60
1	1G	770	C	N1-C2-O2	-5.18	115.79	118.90
26	14	1555	G	C6-C5-N7	-5.18	127.29	130.40
26	14	2169	A	N1-C6-N6	-5.18	115.49	118.60
27	1J	55	U	C6-N1-C2	-5.18	117.89	121.00
26	1H	934	G	C5-C6-O6	-5.18	125.49	128.60
26	1H	1805	U	C6-N1-C2	-5.18	117.89	121.00
1	1G	667	G	C5-C6-O6	-5.18	125.49	128.60
26	14	1996	C	O4'-C1'-N1	-5.18	104.06	108.20
26	1H	481	G	P-O3'-C3'	5.17	125.91	119.70
27	16	44	G	C8-N9-C1'	5.17	133.73	127.00
26	14	121	G	C4-N9-C1'	5.17	133.23	126.50
26	1H	1835	G	O5'-P-OP1	-5.17	101.04	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	1G	518	C	N1-C2-O2	5.17	122.00	118.90
1	1G	785	G	N1-C6-O6	5.17	123.00	119.90
1	13	1227	A	C5-N7-C8	-5.17	101.31	103.90
23	2K	46	G	O5'-P-OP2	-5.17	101.05	105.70
26	1H	1241	A	C5-N7-C8	-5.17	101.31	103.90
26	1H	2506	U	C2-N1-C1'	5.17	123.91	117.70
26	14	685	A	O4'-C1'-N9	5.17	112.34	108.20
26	14	828	U	C5-C6-N1	-5.17	120.11	122.70
26	1H	1332	G	N3-C4-N9	-5.17	122.90	126.00
1	1G	46	G	N9-C4-C5	-5.17	103.33	105.40
26	14	1799	G	N3-C4-C5	5.17	131.19	128.60
1	1G	515	G	N1-C6-O6	5.17	123.00	119.90
26	1H	210	C	C5-C6-N1	-5.17	118.42	121.00
26	1H	1135	C	C2-N1-C1'	5.17	124.48	118.80
26	1H	1603	A	OP1-P-O3'	5.17	116.57	105.20
26	1H	1846	G	C4-C5-N7	5.17	112.87	110.80
26	1H	1899	G	OP2-P-O3'	5.17	116.57	105.20
26	14	322	A	N1-C6-N6	-5.17	115.50	118.60
26	14	330	A	C4-C5-C6	5.17	119.58	117.00
1	13	1058	G	N9-C4-C5	-5.17	103.33	105.40
26	1H	358	U	C5-C6-N1	5.17	125.28	122.70
26	1H	2517	C	C4-C5-C6	5.17	119.98	117.40
26	1H	326	G	C4-C5-N7	5.16	112.87	110.80
26	1H	470	A	O5'-P-OP1	-5.16	101.05	105.70
26	1H	828	U	C6-N1-C2	-5.16	117.90	121.00
26	1H	835	A	C5-C6-N1	5.16	120.28	117.70
26	1H	839	U	C4-C5-C6	5.16	122.80	119.70
26	1H	1822	G	N1-C6-O6	5.16	123.00	119.90
26	14	252	G	N3-C2-N2	-5.16	116.29	119.90
26	14	784	A	N3-C4-C5	5.16	130.41	126.80
26	14	1903	G	OP1-P-OP2	5.16	127.34	119.60
26	1H	458	G	C8-N9-C1'	5.16	133.71	127.00
26	1H	2504	U	N3-C2-O2	-5.16	118.59	122.20
26	1H	1601	G	N9-C4-C5	-5.16	103.34	105.40
26	1H	1799	G	C8-N9-C4	-5.16	104.34	106.40
26	14	265	A	N7-C8-N9	5.16	116.38	113.80
26	14	647	G	N7-C8-N9	5.16	115.68	113.10
1	13	1503	A	OP1-P-O3'	5.16	116.55	105.20
26	1H	943	U	N1-C2-O2	-5.16	119.19	122.80
26	1H	1587	A	C8-N9-C4	-5.16	103.74	105.80
26	1H	2044	C	N3-C4-C5	-5.16	119.84	121.90
26	14	2755	C	C6-N1-C1'	-5.16	114.61	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	2840	C	C4-C5-C6	5.16	119.98	117.40
26	14	684	G	C8-N9-C4	-5.16	104.34	106.40
26	14	1712	C	C6-N1-C2	-5.16	118.24	120.30
26	14	2457	U	N3-C2-O2	-5.16	118.59	122.20
1	13	1516	G	N3-C4-C5	5.16	131.18	128.60
26	1H	165	U	C2-N1-C1'	5.16	123.89	117.70
26	1H	811	U	C2-N3-C4	-5.16	123.91	127.00
26	1H	958	U	C5-C4-O4	5.16	128.99	125.90
1	1G	324	G	N3-C4-N9	-5.16	122.91	126.00
26	14	2576	G	C2-N3-C4	5.16	114.48	111.90
26	1H	198	C	C4-C5-C6	-5.15	114.82	117.40
26	1H	828	U	C4-C5-C6	5.15	122.79	119.70
26	14	634	C	N3-C4-C5	-5.15	119.84	121.90
26	1H	178	G	C8-N9-C4	5.15	108.46	106.40
26	1H	474	G	N9-C4-C5	5.15	107.46	105.40
33	61	58	LEU	CA-CB-CG	5.15	127.15	115.30
1	1G	328	C	C6-N1-C1'	-5.15	114.62	120.80
1	1G	1498	U	N3-C2-O2	-5.15	118.59	122.20
26	14	567	A	C6-C5-N7	-5.15	128.69	132.30
1	13	219	C	C6-N1-C2	-5.15	118.24	120.30
26	1H	1695	G	N3-C2-N2	5.15	123.51	119.90
26	1H	2338	G	N1-C6-O6	5.15	122.99	119.90
6	52	21	LEU	CA-CB-CG	5.15	127.14	115.30
26	14	492	A	N1-C6-N6	5.15	121.69	118.60
26	14	695	G	C8-N9-C4	5.15	108.46	106.40
26	14	986	C	C6-N1-C2	-5.15	118.24	120.30
1	13	172	A	C8-N9-C4	-5.15	103.74	105.80
1	13	1032	A	O4'-C1'-N9	5.15	112.32	108.20
1	1G	698	G	C8-N9-C4	-5.15	104.34	106.40
1	13	865	A	N7-C8-N9	5.15	116.37	113.80
26	1H	941	A	OP2-P-O3'	5.15	116.52	105.20
26	1H	1638	C	C6-N1-C2	-5.15	118.24	120.30
26	1H	2046	G	C6-N1-C2	-5.15	122.01	125.10
1	13	428	G	N9-C4-C5	5.15	107.46	105.40
1	13	827	U	C6-N1-C1'	-5.15	114.00	121.20
1	1G	87	A	N1-C2-N3	5.15	131.87	129.30
1	1G	579	G	C4-C5-N7	5.15	112.86	110.80
26	1H	303	U	C6-N1-C2	-5.14	117.91	121.00
26	1H	783	A	C4-C5-C6	5.14	119.57	117.00
26	1H	1925	C	C6-N1-C2	-5.14	118.24	120.30
27	16	98	G	OP1-P-OP2	5.14	127.32	119.60
26	14	139	G	C8-N9-C4	-5.14	104.34	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	13	514	C	C2-N1-C1'	-5.14	113.14	118.80
1	13	578	C	N3-C4-C5	-5.14	119.84	121.90
26	14	1664	A	C8-N9-C4	-5.14	103.74	105.80
1	13	1430	C	C2-N3-C4	-5.14	117.33	119.90
26	1H	226	G	N3-C2-N2	5.14	123.50	119.90
26	1H	785	G	N1-C6-O6	-5.14	116.82	119.90
26	14	725	G	N1-C6-O6	5.14	122.98	119.90
26	1H	130	C	C5-C4-N4	-5.14	116.60	120.20
26	1H	1950	G	C5-N7-C8	-5.14	101.73	104.30
26	14	1777	U	N1-C2-N3	5.14	117.98	114.90
26	1H	1422	G	C8-N9-C4	-5.14	104.34	106.40
26	14	329	G	C5-C6-O6	5.14	131.68	128.60
26	14	694	U	N3-C2-O2	-5.14	118.60	122.20
26	14	1986	A	C2-N3-C4	-5.14	108.03	110.60
26	14	2600	A	N9-C4-C5	5.14	107.86	105.80
26	1H	2420	C	O5'-P-OP1	-5.14	101.08	105.70
27	16	81	G	N1-C6-O6	5.14	122.98	119.90
1	1G	7	G	N3-C4-N9	-5.14	122.92	126.00
26	14	329	G	N9-C4-C5	5.14	107.45	105.40
26	14	767	U	N3-C2-O2	-5.14	118.60	122.20
26	14	1648	C	N1-C2-O2	-5.14	115.82	118.90
26	1H	137	C	N1-C2-O2	5.13	121.98	118.90
26	1H	1614	A	C5-N7-C8	-5.13	101.33	103.90
26	14	62	C	N1-C2-O2	-5.13	115.82	118.90
26	14	2301	C	C6-N1-C2	-5.13	118.25	120.30
26	1H	130	C	N3-C2-O2	5.13	125.49	121.90
26	1H	839	U	C5-C4-O4	5.13	128.98	125.90
26	1H	2392	A	C6-N1-C2	5.13	121.68	118.60
26	1H	2713	A	N3-C4-C5	5.13	130.39	126.80
27	16	95	U	C6-N1-C1'	5.13	128.38	121.20
1	1G	690	G	N7-C8-N9	5.13	115.67	113.10
26	14	621	A	N3-C4-C5	5.13	130.39	126.80
26	14	1762	A	C5-N7-C8	-5.13	101.33	103.90
1	13	1374	A	O4'-C1'-N9	5.13	112.30	108.20
1	13	1430	C	N1-C2-N3	5.13	122.79	119.20
23	2L	73	C	C6-N1-C2	-5.13	118.25	120.30
26	14	1638	C	OP2-P-O3'	5.13	116.49	105.20
26	1H	144	C	C2-N3-C4	-5.13	117.33	119.90
26	1H	298	G	OP2-P-O3'	5.13	116.48	105.20
26	1H	594	U	OP2-P-O3'	5.13	116.48	105.20
26	1H	670	A	O4'-C1'-N9	-5.13	104.10	108.20
26	1H	964	C	C2-N1-C1'	5.13	124.44	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	1060	U	P-O3'-C3'	5.13	125.86	119.70
26	1H	1422	G	N3-C4-C5	-5.13	126.04	128.60
26	1H	2067	G	C6-N1-C2	-5.13	122.03	125.10
26	14	1857	G	C5-C6-N1	-5.13	108.94	111.50
26	14	2058	A	N1-C6-N6	5.13	121.67	118.60
26	1H	462	C	C6-N1-C1'	5.12	126.95	120.80
1	1G	1113	C	C5-C6-N1	5.12	123.56	121.00
26	14	1558	A	P-O3'-C3'	5.12	125.85	119.70
26	1H	2642	G	N1-C6-O6	5.12	122.97	119.90
26	1H	775	G	O4'-C1'-N9	5.12	112.30	108.20
26	1H	1043	C	C6-N1-C2	-5.12	118.25	120.30
26	1H	1949	G	C6-N1-C2	-5.12	122.03	125.10
1	1G	509	A	P-O3'-C3'	5.12	125.85	119.70
26	14	945	A	N7-C8-N9	5.12	116.36	113.80
26	1H	1206	G	C2-N3-C4	5.12	114.46	111.90
26	14	391	G	C5-N7-C8	-5.12	101.74	104.30
26	14	1609	A	O5'-P-OP2	-5.12	101.09	105.70
26	1H	779	U	OP1-P-OP2	-5.12	111.92	119.60
26	1H	1192	G	C8-N9-C4	5.12	108.45	106.40
26	1H	2029	G	C5-N7-C8	-5.12	101.74	104.30
26	14	2457	U	OP2-P-O3'	5.12	116.46	105.20
1	13	514	C	C6-N1-C2	5.12	122.35	120.30
26	1H	1622	G	C4-C5-N7	-5.12	108.75	110.80
26	1H	2642	G	N9-C4-C5	-5.12	103.35	105.40
26	14	915	C	N3-C2-O2	-5.12	118.32	121.90
26	14	1632	A	C4-C5-N7	5.12	113.26	110.70
1	13	802	A	C6-C5-N7	-5.12	128.72	132.30
28	11	273	ARG	N-CA-C	5.12	124.81	111.00
26	14	458	G	O5'-P-OP2	-5.12	101.10	105.70
26	14	1210	A	C4-C5-N7	5.12	113.26	110.70
26	1H	730	C	C6-N1-C2	-5.11	118.25	120.30
26	1H	739	G	O5'-P-OP1	5.11	116.84	110.70
26	1H	2610	C	N3-C4-C5	5.11	123.94	121.90
1	1G	230	G	C6-C5-N7	-5.11	127.33	130.40
26	14	1379	A	C2-N3-C4	-5.11	108.04	110.60
1	13	1408	A	C8-N9-C4	-5.11	103.75	105.80
26	1H	2724	C	C6-N1-C1'	5.11	126.94	120.80
26	14	1758	G	C5-C6-O6	-5.11	125.53	128.60
26	1H	811	U	N1-C2-N3	5.11	117.97	114.90
26	14	926	A	C5-C6-N6	-5.11	119.61	123.70
26	14	998	C	N1-C2-O2	5.11	121.97	118.90
26	14	2387	U	C2-N1-C1'	-5.11	111.57	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
48	J8	48	LYS	C-N-CA	5.11	134.47	121.70
26	14	912	C	C5-C6-N1	5.11	123.55	121.00
1	1G	767	A	C6-C5-N7	-5.11	128.72	132.30
1	13	1294	G	N3-C4-N9	-5.11	122.94	126.00
26	1H	47	C	N3-C4-C5	5.11	123.94	121.90
26	1H	677	A	C8-N9-C4	-5.11	103.76	105.80
26	1H	923	C	C6-N1-C2	-5.11	118.26	120.30
1	1G	1502	A	C2-N3-C4	-5.11	108.05	110.60
26	14	1601	G	OP1-P-O3'	5.11	116.43	105.20
26	14	2430	A	C5-N7-C8	-5.11	101.35	103.90
26	1H	1794	U	N1-C2-N3	5.10	117.96	114.90
26	14	558	G	C8-N9-C4	5.10	108.44	106.40
26	14	1790	C	N3-C2-O2	5.10	125.47	121.90
1	13	894	G	C4-N9-C1'	-5.10	119.87	126.50
26	1H	2287	A	O4'-C1'-N9	-5.10	104.12	108.20
26	14	236	C	N1-C2-O2	-5.10	115.84	118.90
26	14	939	G	C6-C5-N7	-5.10	127.34	130.40
26	14	2503	A	C8-N9-C4	-5.10	103.76	105.80
26	1H	2236	C	C5-C4-N4	-5.10	116.63	120.20
27	16	19	G	N3-C4-C5	5.10	131.15	128.60
1	1G	1257	U	C5-C6-N1	5.10	125.25	122.70
26	14	1167	U	N3-C2-O2	5.10	125.77	122.20
23	2K	75	G	C4-C5-N7	5.10	112.84	110.80
26	1H	1268	A	C2-N3-C4	-5.10	108.05	110.60
26	1H	1558	A	C5-C6-N6	5.10	127.78	123.70
26	1H	2491	U	N1-C2-O2	5.10	126.37	122.80
1	1G	108	G	N1-C6-O6	5.10	122.96	119.90
1	1G	413	G	C4-N9-C1'	-5.10	119.87	126.50
26	14	204	A	C6-N1-C2	-5.10	115.54	118.60
26	14	2426	A	C5-N7-C8	-5.10	101.35	103.90
27	1J	96	G	N1-C2-N2	5.10	120.79	116.20
23	2K	64	G	C8-N9-C4	-5.10	104.36	106.40
26	1H	739	G	N7-C8-N9	-5.10	110.55	113.10
26	14	621	A	C6-N1-C2	5.10	121.66	118.60
26	14	685	A	N1-C6-N6	-5.10	115.54	118.60
26	14	933	A	N9-C4-C5	-5.10	103.76	105.80
26	14	1664	A	C4-N9-C1'	5.10	135.48	126.30
26	1H	757	U	O5'-P-OP2	-5.10	101.11	105.70
26	1H	639	U	C5-C4-O4	5.09	128.96	125.90
26	1H	1751	C	N3-C2-O2	5.09	125.47	121.90
26	1H	2246	G	N3-C4-N9	5.09	129.06	126.00
26	1H	2700	C	C6-N1-C2	5.09	122.34	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	562	U	N3-C2-O2	-5.09	118.63	122.20
26	1H	467	G	N3-C4-N9	5.09	129.06	126.00
26	1H	514	A	N7-C8-N9	-5.09	111.25	113.80
26	1H	1975	G	N3-C4-N9	5.09	129.06	126.00
26	1H	2011	U	N3-C2-O2	5.09	125.76	122.20
26	1H	2565	A	N7-C8-N9	-5.09	111.25	113.80
26	1H	2388	A	N9-C4-C5	-5.09	103.76	105.80
26	1H	2779	U	N1-C2-N3	5.09	117.95	114.90
1	1G	527	G	C4-C5-N7	-5.09	108.76	110.80
26	14	229	A	O4'-C1'-N9	5.09	112.27	108.20
26	14	2374	C	C6-N1-C2	5.09	122.34	120.30
26	1H	1299	G	N7-C8-N9	5.09	115.64	113.10
26	1H	1576	U	N3-C4-C5	-5.09	111.55	114.60
26	14	783	A	C8-N9-C4	-5.09	103.77	105.80
26	1H	188	G	C5-C6-N1	5.09	114.04	111.50
26	1H	1786	A	O4'-C1'-N9	5.09	112.27	108.20
1	1G	922	G	N3-C4-C5	-5.09	126.06	128.60
1	1G	1278	U	C5-C6-N1	5.09	125.24	122.70
1	1G	1274	G	C4-N9-C1'	5.08	133.11	126.50
26	1H	945	A	C5-C6-N6	-5.08	119.63	123.70
26	1H	1616	A	C4-C5-C6	5.08	119.54	117.00
26	1H	1807	G	C5-C6-O6	-5.08	125.55	128.60
26	14	1930	G	C4-N9-C1'	-5.08	119.89	126.50
1	13	865	A	C6-C5-N7	-5.08	128.74	132.30
24	3K	18	G	OP2-P-O3'	5.08	116.38	105.20
26	1H	431	U	C5-C6-N1	5.08	125.24	122.70
26	1H	1325	G	N1-C6-O6	-5.08	116.85	119.90
26	1H	1544	C	C6-N1-C2	5.08	122.33	120.30
24	3L	83	U	N1-C1'-C2'	-5.08	106.41	112.00
26	14	1395	A	N7-C8-N9	-5.08	111.26	113.80
26	1H	213	A	C8-N9-C4	5.08	107.83	105.80
26	1H	1413	G	C5-C6-O6	-5.08	125.55	128.60
26	14	2499	C	C6-N1-C2	-5.08	118.27	120.30
1	1G	1417	G	C6-C5-N7	-5.08	127.35	130.40
26	14	1566	A	N1-C6-N6	5.08	121.65	118.60
26	14	2689	U	C5-C4-O4	5.08	128.95	125.90
26	1H	259	G	N1-C6-O6	5.08	122.95	119.90
26	14	756	C	C5-C4-N4	-5.08	116.65	120.20
1	13	874	G	N1-C6-O6	-5.08	116.85	119.90
27	16	12	C	C2-N3-C4	-5.08	117.36	119.90
26	14	22	C	N3-C4-C5	5.08	123.93	121.90
27	1J	22	U	C2-N1-C1'	5.08	123.79	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
27	1J	116	G	N3-C4-C5	5.08	131.14	128.60
1	13	1206	G	N3-C4-N9	5.07	129.04	126.00
26	1H	1265	A	C8-N9-C4	-5.07	103.77	105.80
26	1H	2636	U	O5'-P-OP1	-5.07	101.13	105.70
26	14	1730	U	C2-N1-C1'	5.07	123.79	117.70
26	14	2423	U	C5-C4-O4	-5.07	122.86	125.90
26	1H	615	G	O4'-C1'-N9	5.07	112.26	108.20
26	1H	450	G	N1-C6-O6	5.07	122.94	119.90
26	1H	678	C	C4-C5-C6	5.07	119.94	117.40
26	1H	2025	C	C2-N1-C1'	5.07	124.38	118.80
26	1H	2377	A	N9-C4-C5	-5.07	103.77	105.80
27	16	24	G	N3-C4-C5	-5.07	126.06	128.60
1	1G	383	A	N1-C6-N6	5.07	121.64	118.60
26	14	53	A	C6-N1-C2	-5.07	115.56	118.60
26	14	1306	C	O5'-P-OP1	-5.07	101.14	105.70
26	1H	828	U	N3-C4-O4	-5.07	115.85	119.40
26	1H	2082	A	C6-N1-C2	-5.07	115.56	118.60
1	1G	500	G	C4-N9-C1'	5.07	133.09	126.50
23	2L	32	C	OP1-P-O3'	5.07	116.35	105.20
26	14	2394	C	O5'-P-OP2	-5.07	101.14	105.70
26	1H	99	U	C2-N1-C1'	5.07	123.78	117.70
26	1H	395	U	N1-C2-O2	5.07	126.35	122.80
26	1H	799	G	N1-C2-N3	5.07	126.94	123.90
26	1H	1568	G	OP1-P-OP2	-5.07	112.00	119.60
1	1G	1278	U	N3-C2-O2	-5.07	118.65	122.20
26	14	2499	C	C2-N1-C1'	5.07	124.37	118.80
26	1H	670	A	C8-N9-C4	5.07	107.83	105.80
26	1H	790	C	N3-C2-O2	5.07	125.45	121.90
26	1H	1348	G	C5-C6-O6	-5.07	125.56	128.60
1	1G	596	C	C6-N1-C2	5.07	122.33	120.30
26	1H	1372	U	N3-C4-O4	5.06	122.94	119.40
26	14	2239	G	N3-C4-N9	5.06	129.04	126.00
26	1H	330	A	N3-C4-C5	5.06	130.34	126.80
26	1H	958	U	N3-C2-O2	-5.06	118.66	122.20
1	1G	232	G	C4-N9-C1'	5.06	133.08	126.50
26	14	792	G	C8-N9-C4	-5.06	104.38	106.40
26	14	1308	A	N9-C4-C5	5.06	107.82	105.80
26	1H	965	C	C6-N1-C2	-5.06	118.28	120.30
26	1H	1350	C	C6-N1-C2	5.06	122.32	120.30
26	1H	2618	G	C5-C6-O6	5.06	131.64	128.60
26	14	197	A	OP2-P-O3'	5.06	116.33	105.20
26	14	549	G	N1-C6-O6	5.06	122.94	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	933	A	C5-C6-N6	-5.06	119.65	123.70
26	14	2624	G	C5-C6-O6	-5.06	125.56	128.60
26	14	2713	A	C2-N3-C4	-5.06	108.07	110.60
1	13	898	G	N3-C4-N9	5.06	129.03	126.00
26	1H	1004	C	C6-N1-C2	-5.06	118.28	120.30
26	1H	1354	A	OP1-P-O3'	5.06	116.33	105.20
26	1H	1950	G	N3-C2-N2	5.06	123.44	119.90
26	1H	2578	G	C2-N3-C4	5.06	114.43	111.90
25	4L	51	U	C6-N1-C2	-5.06	117.97	121.00
26	14	1890	A	C8-N9-C4	5.06	107.82	105.80
26	14	2318	G	C8-N9-C4	-5.06	104.38	106.40
26	14	2604	U	C6-N1-C2	5.06	124.03	121.00
26	1H	248	G	C8-N9-C4	5.06	108.42	106.40
26	14	711	G	N1-C6-O6	5.06	122.93	119.90
26	1H	1345	C	N3-C2-O2	5.05	125.44	121.90
26	1H	2228	G	N1-C2-N2	-5.05	111.65	116.20
1	1G	944	G	N3-C4-C5	-5.05	126.07	128.60
26	14	2880	C	N3-C4-C5	-5.05	119.88	121.90
1	13	919	A	N9-C4-C5	5.05	107.82	105.80
26	14	313	C	C5-C6-N1	5.05	123.53	121.00
26	1H	774	A	C4-N9-C1'	-5.05	117.21	126.30
26	1H	2011	U	N1-C2-N3	-5.05	111.87	114.90
26	1H	2294	C	N3-C4-C5	5.05	123.92	121.90
1	1G	774	G	O4'-C1'-N9	-5.05	104.16	108.20
26	14	2447	G	P-O3'-C3'	5.05	125.76	119.70
1	13	1190	G	N1-C6-O6	5.05	122.93	119.90
26	1H	827	U	O5'-P-OP1	5.05	116.76	110.70
26	1H	1846	G	C6-C5-N7	-5.05	127.37	130.40
1	1G	701	C	N1-C2-O2	5.05	121.93	118.90
4	32	85	LYS	N-CA-C	-5.05	97.36	111.00
26	14	2168	G	N3-C4-N9	5.05	129.03	126.00
26	14	2826	A	N1-C2-N3	5.05	131.82	129.30
26	1H	2304	G	C4-N9-C1'	-5.05	119.94	126.50
26	14	27	G	N3-C4-N9	-5.05	122.97	126.00
1	13	1422	G	C8-N9-C4	5.05	108.42	106.40
26	1H	202	U	C5-C4-O4	-5.05	122.87	125.90
26	1H	1942	C	C2-N3-C4	-5.05	117.38	119.90
1	1G	1053	G	P-O3'-C3'	5.05	125.76	119.70
26	14	2634	G	C5-C6-O6	-5.05	125.57	128.60
29	29	50	GLY	N-CA-C	5.05	125.72	113.10
26	1H	32	C	C6-N1-C1'	5.04	126.85	120.80
26	1H	468	G	C4-C5-N7	5.04	112.82	110.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	1135	C	C6-N1-C1'	-5.04	114.75	120.80
26	1H	2409	G	N7-C8-N9	5.04	115.62	113.10
1	1G	689	C	O5'-P-OP1	-5.04	101.16	105.70
1	1G	1119	C	C6-N1-C2	-5.04	118.28	120.30
26	14	2573	C	C5-C4-N4	-5.04	116.67	120.20
1	13	576	G	C8-N9-C1'	-5.04	120.45	127.00
26	1H	1691	C	N1-C2-O2	-5.04	115.88	118.90
26	14	864	G	C8-N9-C4	-5.04	104.38	106.40
26	14	1629	U	N1-C2-O2	-5.04	119.27	122.80
26	14	2723	C	N3-C2-O2	-5.04	118.37	121.90
26	1H	273	G	N1-C6-O6	5.04	122.92	119.90
1	1G	360	A	C5-C6-N6	-5.04	119.67	123.70
26	14	681	G	N3-C4-N9	5.04	129.02	126.00
26	14	827	U	O5'-P-OP2	-5.04	101.16	105.70
27	1J	114	G	N7-C8-N9	-5.04	110.58	113.10
1	13	505	G	N9-C4-C5	-5.04	103.39	105.40
1	13	1498	U	OP2-P-O3'	5.04	116.28	105.20
1	13	1533	C	N3-C2-O2	-5.04	118.37	121.90
26	1H	1807	G	N1-C6-O6	5.04	122.92	119.90
26	14	1259	G	C5-C6-N1	5.04	114.02	111.50
26	14	1283	G	N1-C6-O6	5.04	122.92	119.90
26	14	2389	G	C4-C5-N7	5.04	112.81	110.80
1	13	689	C	N3-C2-O2	-5.04	118.38	121.90
26	1H	131	G	N1-C6-O6	5.04	122.92	119.90
26	1H	217	G	N9-C4-C5	5.04	107.41	105.40
26	1H	2429	G	OP2-P-O3'	5.04	116.28	105.20
26	1H	2603	G	OP1-P-O3'	5.04	116.28	105.20
26	14	1021	A	N1-C2-N3	5.04	131.82	129.30
26	14	2054	A	O5'-P-OP1	-5.04	101.17	105.70
26	14	2168	G	C8-N9-C1'	-5.04	120.45	127.00
23	2K	22	G	C6-C5-N7	-5.03	127.38	130.40
26	1H	779	U	N3-C4-C5	5.03	117.62	114.60
26	1H	1162	G	C8-N9-C4	-5.03	104.39	106.40
26	1H	1501	C	C6-N1-C2	-5.03	118.29	120.30
26	1H	2032	G	N1-C2-N3	5.03	126.92	123.90
26	14	1603	A	N7-C8-N9	5.03	116.32	113.80
1	13	575	G	C8-N9-C4	5.03	108.41	106.40
26	14	2634	G	C8-N9-C4	5.03	108.41	106.40
1	13	865	A	N1-C6-N6	5.03	121.62	118.60
26	1H	1025	G	C8-N9-C4	-5.03	104.39	106.40
26	1H	1771	C	N3-C4-N4	5.03	121.52	118.00
26	1H	2081	C	N3-C4-C5	-5.03	119.89	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	1G	686	U	O4'-C1'-N1	5.03	112.22	108.20
26	14	528	A	C5-N7-C8	-5.03	101.38	103.90
26	14	813	U	O5'-P-OP2	-5.03	101.17	105.70
1	13	1408	A	N9-C4-C5	5.03	107.81	105.80
1	13	1440	C	C5-C6-N1	-5.03	118.49	121.00
26	14	1688	U	N1-C2-N3	5.03	117.92	114.90
1	13	1052	U	N3-C2-O2	-5.03	118.68	122.20
26	1H	119	A	N1-C2-N3	5.03	131.81	129.30
26	1H	326	G	C5-C6-O6	-5.03	125.58	128.60
1	1G	197	A	C4-C5-C6	5.03	119.51	117.00
1	1G	1499	A	C8-N9-C4	5.03	107.81	105.80
26	14	1203	G	N3-C2-N2	5.03	123.42	119.90
26	14	2311	A	N7-C8-N9	5.03	116.31	113.80
26	14	2859	G	N1-C6-O6	-5.03	116.88	119.90
1	13	124	G	C5-C6-N1	-5.03	108.99	111.50
26	1H	920	G	N3-C4-C5	5.03	131.11	128.60
54	Q8	52	LYS	C-N-CA	5.03	143.11	122.00
1	1G	268	C	C6-N1-C2	-5.03	118.29	120.30
26	14	206	U	N3-C2-O2	-5.03	118.68	122.20
26	14	723	G	C8-N9-C4	5.03	108.41	106.40
26	14	1897	G	C4-N9-C1'	5.03	133.03	126.50
1	13	1413	A	C4-C5-N7	5.02	113.21	110.70
26	1H	2084	C	C4-C5-C6	5.02	119.91	117.40
1	1G	111	G	C6-C5-N7	-5.02	127.39	130.40
1	13	558	G	N1-C6-O6	5.02	122.91	119.90
23	2K	44	G	P-O3'-C3'	5.02	125.73	119.70
26	1H	270(H)	C	C5-C6-N1	5.02	123.51	121.00
26	1H	769	G	O5'-P-OP2	-5.02	101.18	105.70
26	1H	2468	G	C8-N9-C1'	-5.02	120.47	127.00
26	1H	2617	C	N3-C2-O2	5.02	125.42	121.90
26	14	1021	A	N3-C4-C5	5.02	130.32	126.80
1	13	1114	C	C5-C6-N1	5.02	123.51	121.00
26	1H	468	G	C6-C5-N7	-5.02	127.39	130.40
26	1H	1814	G	C5-C6-N1	5.02	114.01	111.50
26	1H	2337	G	C8-N9-C4	-5.02	104.39	106.40
26	14	1215	G	C8-N9-C4	-5.02	104.39	106.40
26	14	2355	C	C2-N1-C1'	5.02	124.32	118.80
26	14	2430	A	O5'-P-OP2	5.02	116.72	110.70
23	2K	81	G	C5-C6-N1	-5.02	108.99	111.50
26	1H	302	C	N1-C2-O2	5.02	121.91	118.90
26	1H	1320	C	C5-C6-N1	-5.02	118.49	121.00
26	1H	2228	G	N3-C4-C5	-5.02	126.09	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	2454	G	N7-C8-N9	-5.02	110.59	113.10
26	14	1634	A	C4-C5-N7	-5.02	108.19	110.70
26	14	1906	G	O5'-P-OP2	5.02	116.72	110.70
26	14	2789	C	C6-N1-C2	-5.02	118.29	120.30
1	13	1381	U	C2-N1-C1'	5.02	123.72	117.70
26	1H	793	A	N1-C2-N3	5.02	131.81	129.30
26	1H	2432	A	C2-N3-C4	-5.02	108.09	110.60
1	1G	99	C	N1-C2-O2	-5.02	115.89	118.90
26	1H	162	U	N1-C2-O2	5.02	126.31	122.80
1	1G	560	U	C5-C6-N1	5.02	125.21	122.70
26	1H	602	G	N1-C6-O6	5.01	122.91	119.90
26	1H	821	A	C8-N9-C4	-5.01	103.79	105.80
26	14	102	G	O5'-P-OP1	-5.01	101.19	105.70
26	1H	1263	U	N3-C2-O2	-5.01	118.69	122.20
26	14	1334	G	C6-C5-N7	-5.01	127.39	130.40
26	14	1396	U	N1-C2-O2	5.01	126.31	122.80
26	1H	70	G	P-O3'-C3'	5.01	125.71	119.70
26	1H	719	C	C6-N1-C2	-5.01	118.30	120.30
26	1H	743	G	N9-C4-C5	5.01	107.40	105.40
26	1H	1543	A	C6-C5-N7	-5.01	128.79	132.30
26	14	2226	C	C5-C6-N1	5.01	123.50	121.00
26	14	2871	C	C6-N1-C2	5.01	122.30	120.30
26	1H	71	A	O4'-C1'-N9	-5.01	104.19	108.20
26	1H	1695	G	C4-N9-C1'	5.01	133.01	126.50
26	1H	2331	G	C4-C5-N7	5.01	112.80	110.80
26	14	2137	C	C2-N1-C1'	5.01	124.31	118.80
26	14	2287	A	N3-C4-C5	5.01	130.31	126.80
26	1H	70	G	C5-C6-O6	5.01	131.60	128.60
26	1H	470	A	C6-N1-C2	-5.01	115.60	118.60
26	1H	1378	A	N3-C4-N9	-5.01	123.39	127.40
26	1H	2568	C	C2-N3-C4	-5.01	117.40	119.90
27	1J	114	G	N3-C4-N9	-5.01	123.00	126.00
26	1H	999	U	C5-C4-O4	5.00	128.90	125.90
26	1H	1025	G	C2-N3-C4	5.00	114.40	111.90
26	1H	1471	A	C8-N9-C4	-5.00	103.80	105.80
26	1H	2367	G	C2-N3-C4	-5.00	109.40	111.90
26	1H	115	C	N1-C2-O2	-5.00	115.90	118.90
26	1H	2383	G	N3-C4-C5	-5.00	126.10	128.60
1	1G	1471	G	N3-C4-N9	5.00	129.00	126.00
26	14	71	A	C5-C6-N1	-5.00	115.20	117.70
1	13	1523	G	N3-C2-N2	-5.00	116.40	119.90
26	1H	121	G	C4-C5-C6	5.00	121.80	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	1H	637	A	N1-C6-N6	5.00	121.60	118.60
26	1H	779	U	O5'-P-OP1	5.00	116.70	110.70
26	1H	1685	C	N3-C4-C5	5.00	123.90	121.90
1	1G	87	A	C5-C6-N6	-5.00	119.70	123.70
1	1G	224	C	C6-N1-C2	5.00	122.30	120.30

There are no chirality outliers.

All (128) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
28	11	273	ARG	Peptide
2	12	19	HIS	Peptide
34	15	41	ASP	Peptide
28	19	237	GLU	Peptide
28	19	32	SER	Peptide
28	19	37	LEU	Peptide
10	1A	55	LYS	Peptide
2	1E	15	VAL	Peptide
2	1E	169	LYS	Peptide
2	1E	237	ALA	Peptide
29	21	153	GLY	Peptide
29	21	57	LYS	Peptide
29	21	78	LEU	Peptide
29	21	82	ARG	Peptide
29	29	117	MET	Peptide
29	29	139	GLY	Peptide
29	29	186	GLY	Peptide
29	29	201	THR	Peptide
29	29	203	LYS	Peptide
29	29	44	TYR	Peptide
29	29	53	PRO	Peptide
29	29	70	ALA	Peptide
11	2A	49	GLY	Peptide
30	31	196	LEU	Peptide
4	32	154	ASN	Peptide
4	32	155	LEU	Peptide
4	32	30	LYS	Peptide
36	35	110	TYR	Peptide
36	35	22	GLY	Peptide
36	35	36	LYS	Peptide
36	35	5	ASP	Peptide
36	35	56	SER	Peptide

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Mol	Chain	Res	Type	Group
36	35	60	MET	Peptide
36	35	63	PRO	Peptide
30	39	127	GLU	Peptide
30	39	166	ALA	Peptide
30	39	20	LEU	Peptide
30	39	24	LEU	Peptide
30	39	26	ALA	Peptide
30	39	89	VAL	Peptide
12	3A	18	VAL	Peptide
12	3A	89	ARG	Peptide
4	3E	31	CYS	Peptide
12	3I	87	GLY	Peptide
37	45	134	ARG	Peptide
37	45	135	ASP	Peptide
37	45	137	TYR	Peptide
37	45	25	ASP	Peptide
37	45	26	TYR	Peptide
31	49	13	GLU	Peptide
13	4A	94	ARG	Peptide
38	55	105	ARG	Peptide
38	55	106	GLY	Peptide
14	5A	29	ARG	Peptide
33	61	11	ASN	Peptide
33	61	134	PRO	Peptide
33	61	82	ARG	Peptide
33	69	112	LYS	Peptide
33	69	143	SER	Peptide
33	69	85	GLU	Peptide
40	75	12	SER	Peptide
40	75	4	GLY	Peptide
36	78	11	GLY	Peptide
36	78	115	LEU	Peptide
9	82	117	HIS	Peptide
41	85	90	VAL	Peptide
41	85	98	LEU	Peptide
37	88	21	THR	Peptide
37	88	58	PHE	Peptide
37	88	79	LEU	Peptide
9	8E	47	LEU	Peptide
42	95	85	LYS	Peptide
38	98	44	LEU	Peptide
43	A5	43	GLY	Peptide

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>	<b>Group</b>
19	AI	6	LYS	Peptide
19	AI	7	LYS	Peptide
40	B8	12	SER	Peptide
40	B8	58	ASN	Peptide
20	BA	11	SER	Peptide
20	BA	72	LEU	Peptide
45	C5	100	ALA	Peptide
45	C5	39	VAL	Peptide
45	C5	81	LYS	Peptide
45	C5	90	LEU	Peptide
41	C8	115	ALA	Peptide
47	E5	7	LEU	Peptide
47	E5	83	PRO	Peptide
48	F5	78	LYS	Peptide
49	G5	15	LYS	Peptide
49	G5	17	SER	Peptide
49	G5	43	GLN	Peptide
45	G8	53	PRO	Peptide
45	G8	54	LYS	Peptide
45	G8	80	GLY	Peptide
45	G8	94	LYS	Peptide
46	H8	165	VAL	Peptide
46	H8	59	LEU	Peptide
46	H8	63	ASP	Peptide
51	I5	26	SER	Peptide
47	I8	2	ALA	Peptide
47	I8	6	ALA	Peptide
52	J5	3	LYS	Peptide
48	J8	75	GLU	Peptide
49	K8	15	LYS	Peptide
49	K8	16	LEU	Peptide
49	K8	17	SER	Peptide
49	K8	45	SER	Peptide
49	K8	46	GLN	Peptide
54	M5	35	GLN	Peptide
54	M5	40	GLU	Peptide
51	M8	4	GLY	Peptide
51	M8	40	HIS	Peptide
51	M8	43	TYR	Peptide
52	N8	41	PRO	Peptide
52	N8	42	PRO	Peptide
53	P8	46	VAL	Peptide

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Mol	Chain	Res	Type	Group
54	Q8	18	ALA	Peptide
54	Q8	26	LYS	Peptide
54	Q8	33	ASN	Peptide
54	Q8	37	SER	Peptide
54	Q8	38	GLY	Peptide
54	Q8	46	ARG	Peptide
54	Q8	48	PHE	Peptide
54	Q8	56	GLU	Peptide
54	Q8	57	ARG	Peptide
54	Q8	6	THR	Peptide
54	Q8	7	HIS	Peptide
54	Q8	9	GLY	Peptide

## 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	13	32389	0	16350	718	0
1	1G	32511	0	16414	717	0
2	12	1924	0	1975	96	0
2	1E	1924	0	1975	88	0
3	22	1612	0	1677	83	0
3	2E	1605	0	1668	69	0
4	32	1702	0	1763	104	0
4	3E	1702	0	1763	77	0
5	42	1155	0	1213	52	0
5	4E	1155	0	1213	55	0
6	52	842	0	857	32	0
6	5E	842	0	857	33	0
7	62	1200	0	1238	51	0
7	6E	1157	0	1202	45	0
8	72	1115	0	1177	46	0
8	7E	1115	0	1177	42	0
9	82	983	0	1006	76	0
9	8E	1009	0	1037	68	0
10	1A	801	0	849	56	0
10	1I	801	0	849	53	0
11	2A	864	0	881	37	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
11	2I	864	0	881	39	0
12	3A	975	0	1062	41	0
12	3I	956	0	1046	47	0
13	4A	928	0	987	52	0
13	4I	928	0	987	55	0
14	5A	475	0	511	36	0
14	5I	491	0	529	31	0
15	6A	733	0	771	31	0
15	6I	733	0	771	25	0
16	7A	705	0	725	27	0
16	7I	705	0	725	40	0
17	8A	823	0	891	32	0
17	8I	834	0	904	39	0
18	9A	564	0	631	22	0
18	9I	550	0	613	23	0
19	AA	624	0	636	42	0
19	AI	643	0	662	43	0
20	BA	762	0	861	44	0
20	BI	762	0	861	40	0
21	1B	217	0	234	11	0
21	1F	199	0	208	9	0
22	1K	1863	0	942	41	0
22	1L	1863	0	942	40	0
23	2K	1692	0	859	47	0
23	2L	1712	0	870	62	0
24	3K	1646	0	830	36	0
24	3L	1601	0	809	30	0
25	4K	336	0	168	9	0
25	4L	438	0	220	11	0
26	14	62647	0	31582	1257	0
26	1H	62707	0	31614	1349	1
27	16	2617	0	1328	60	0
27	1J	2617	0	1328	70	0
28	11	2126	0	2208	88	0
28	19	2120	0	2197	86	0
29	21	1568	0	1634	96	0
29	29	1568	0	1634	104	0
30	31	1585	0	1632	84	0
30	39	1627	0	1680	94	0
31	41	1473	0	1535	92	0
31	49	1473	0	1535	67	0
32	51	1336	0	1418	80	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
32	59	1307	0	1382	73	0
33	61	1136	0	1223	59	0
33	69	1136	0	1223	57	0
34	15	1104	0	1180	49	0
34	58	1104	0	1180	56	0
35	25	932	0	996	43	0
35	68	932	0	996	35	0
36	35	1144	0	1228	82	0
36	78	1122	0	1206	97	0
37	45	1113	0	1167	64	0
37	88	1121	0	1179	66	0
38	55	959	0	1021	51	0
38	98	967	0	1033	58	0
39	65	881	0	943	59	0
39	A8	881	0	943	57	0
40	75	1141	0	1202	54	0
40	B8	1081	0	1141	63	0
41	85	963	0	1022	61	0
41	C8	963	0	1022	60	0
42	95	778	0	852	60	0
42	D8	778	0	852	37	0
43	A5	899	0	964	34	0
43	E8	890	0	951	31	0
44	B5	725	0	778	38	0
44	F8	742	0	803	30	0
45	C5	794	0	884	52	0
45	G8	783	0	870	46	0
46	D5	1120	0	1146	54	0
46	H8	1373	0	1402	75	0
47	E5	627	0	645	42	0
47	I8	656	0	679	32	0
48	F5	737	0	813	38	0
48	J8	746	0	826	34	0
49	G5	567	0	618	20	0
49	K8	575	0	634	38	0
50	H5	468	0	518	15	1
50	L8	452	0	503	19	0
51	I5	515	0	514	35	0
51	M8	533	0	526	52	0
52	J5	434	0	454	19	0
52	N8	429	0	449	31	0
53	L5	409	0	454	21	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
53	P8	409	0	454	18	0
54	M5	422	0	484	30	0
54	Q8	480	0	549	94	0
55	11	1	0	0	0	0
55	13	146	0	0	0	0
55	14	490	0	0	0	0
55	16	5	0	0	0	0
55	1G	143	0	0	0	0
55	1H	512	0	0	0	0
55	1J	2	0	0	0	0
55	1K	1	0	0	0	0
55	21	2	0	0	0	0
55	25	1	0	0	0	0
55	29	3	0	0	0	0
55	2K	3	0	0	0	0
55	2L	2	0	0	0	0
55	31	1	0	0	0	0
55	32	1	0	0	0	0
55	3E	1	0	0	0	0
55	3I	1	0	0	0	0
55	45	2	0	0	0	0
55	5E	1	0	0	0	0
55	78	1	0	0	0	0
55	7A	1	0	0	0	0
55	85	2	0	0	0	0
55	88	4	0	0	0	0
55	98	1	0	0	0	0
55	E5	2	0	0	0	0
55	G8	1	0	0	0	0
55	I8	2	0	0	0	0
55	J8	1	0	0	0	0
55	P8	1	0	0	0	0
56	32	1	0	0	0	0
56	3E	1	0	0	0	0
56	5A	1	0	0	0	0
56	5I	1	0	0	0	0
56	C5	1	0	0	0	0
56	G8	1	0	0	0	0
57	11	1	0	0	0	0
57	13	74	0	0	11	0
57	14	520	0	0	102	0
57	19	8	0	0	2	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
57	1G	73	0	0	6	0
57	1H	552	0	0	122	0
57	21	1	0	0	1	0
57	29	2	0	0	0	0
57	31	4	0	0	1	0
57	35	1	0	0	0	0
57	39	4	0	0	0	0
57	3E	1	0	0	0	0
57	4L	2	0	0	0	0
57	55	2	0	0	0	0
57	6A	2	0	0	0	0
57	6I	1	0	0	0	0
57	78	2	0	0	0	0
57	7A	1	0	0	0	0
57	85	1	0	0	0	0
57	A5	1	0	0	0	0
57	B8	1	0	0	0	0
57	BI	1	0	0	0	0
57	C8	2	0	0	1	0
57	D8	1	0	0	0	0
57	E8	1	0	0	0	0
57	I8	3	0	0	0	0
57	J8	1	0	0	0	0
57	M5	3	0	0	0	0
57	P8	1	0	0	0	0
All	All	299429	0	199511	8202	1

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 17.

All (8202) 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)
26:1H:2032:G:N7	57:1H:4116:HOH:O	1.81	1.11
2:12:185:ILE:HG22	2:12:199:TYR:HB2	1.39	1.04
26:1H:810:U:OP1	57:1H:4123:HOH:O	1.75	1.03
26:1H:2308:G:H1	26:1H:2311:A:H2	1.07	1.02
26:1H:763:G:OP1	57:1H:3623:HOH:O	1.80	1.00
26:1H:1602:U:O4	57:1H:3791:HOH:O	1.78	1.00
28:19:37:LEU:HA	28:19:38:LYS:HB3	1.45	0.99
1:13:1305:G:H22	1:13:1331:G:H2'	1.25	0.98

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2448:A:OP1	57:14:3631:HOH:O	1.81	0.98
26:14:2701:C:H3'	26:14:2702:U:H5''	1.46	0.97
4:32:26:CYS:HA	4:32:31:CYS:HB3	1.44	0.97
1:13:1502:A:H2	1:13:1505:G:H1	0.99	0.96
26:1H:2056:G:N7	57:1H:3925:HOH:O	1.98	0.96
50:L8:8:LEU:HA	50:L8:54:VAL:HB	1.46	0.96
26:14:1647:G:OP2	57:14:3939:HOH:O	1.81	0.96
26:1H:1774:C:OP1	57:1H:3665:HOH:O	1.84	0.95
54:M5:40:GLU:H	54:M5:43:GLN:HG3	1.27	0.95
26:14:2624:G:N7	57:14:3935:HOH:O	1.98	0.94
26:1H:1388:G:N7	57:1H:3802:HOH:O	1.99	0.94
26:14:1639:U:OP1	57:14:3953:HOH:O	1.85	0.94
26:1H:330:A:HO2'	26:1H:331:A:H8	1.10	0.94
26:14:1332:G:N2	26:14:1609:A:O2'	2.01	0.94
26:1H:761:A:OP1	57:1H:3618:HOH:O	1.86	0.94
34:15:47:ALA:HB2	34:15:112:LEU:HD21	1.49	0.93
26:1H:567:A:OP1	57:1H:3603:HOH:O	1.86	0.93
32:59:6:ARG:HB2	32:59:66:GLY:HA2	1.50	0.93
26:14:1349:A:OP1	57:14:3543:HOH:O	1.84	0.93
26:1H:217:G:OP2	57:1H:3643:HOH:O	1.86	0.93
23:2L:37:G:H1	25:4L:47:U:H3	1.11	0.93
26:1H:751:A:OP1	57:1H:3745:HOH:O	1.84	0.93
30:39:53:THR:HG23	30:39:55:GLY:H	1.34	0.92
32:59:137:ASP:HB3	32:59:140:LYS:HB2	1.49	0.92
26:1H:1614:A:OP1	57:1H:3742:HOH:O	1.87	0.92
26:1H:452:G:OP2	57:1H:3740:HOH:O	1.86	0.92
26:14:1774:C:OP1	57:14:3551:HOH:O	1.88	0.92
26:14:676:A:H8	26:14:2069:G:H21	1.11	0.92
1:1G:758:G:N7	57:1G:1869:HOH:O	2.03	0.92
1:13:200:G:H1	1:13:217:C:H42	1.16	0.92
20:BA:82:SER:OG	20:BA:86:ARG:NH2	2.03	0.92
26:1H:801:G:OP2	57:1H:3907:HOH:O	1.87	0.92
26:14:741:G:OP1	57:14:3546:HOH:O	1.88	0.91
26:1H:2287:A:N6	26:1H:2344:U:H3	1.69	0.91
26:14:4:C:H42	26:14:2899:G:H22	1.10	0.91
8:7E:87:SER:HB2	8:7E:93:VAL:HB	1.52	0.91
26:14:1899:G:N2	26:14:1902:C:H41	1.68	0.91
23:2L:1:G:H1	23:2L:83:U:H3	1.18	0.91
29:21:201:THR:HG22	29:21:203:LYS:H	1.36	0.90
26:14:1613:G:N7	57:14:3608:HOH:O	2.05	0.90
3:22:32:LEU:HB3	3:22:59:ARG:HH12	1.36	0.90

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1890:A:OP2	57:1H:4002:HOH:O	1.90	0.90
1:1G:975:A:H4'	1:1G:976:G:H5''	1.52	0.90
1:13:137:C:H42	1:13:226:G:H1	0.98	0.89
26:1H:363(B):G:H2'	26:1H:363(C):G:H8	1.37	0.89
26:14:450:G:OP2	57:14:3656:HOH:O	1.87	0.89
26:14:751:A:OP1	57:14:3508:HOH:O	1.89	0.89
26:14:1496:A:H8	26:14:1577:C:HO2'	0.96	0.89
26:14:2415:G:H4'	36:35:67:MET:H	1.38	0.89
26:1H:1359:A:N1	26:1H:1372:U:N3	2.20	0.89
31:41:109:VAL:HG11	31:41:142:PRO:HD3	1.55	0.89
22:1L:46:G:H22	22:1L:58:U:H1'	1.34	0.89
26:1H:2053:G:OP1	57:1H:3670:HOH:O	1.91	0.89
26:1H:2576:G:OP1	57:1H:3671:HOH:O	1.91	0.89
26:1H:1823:G:N7	57:1H:4094:HOH:O	2.05	0.88
1:13:8:A:H62	4:3E:208:SER:HB2	1.38	0.88
26:14:1757:U:H3	26:14:1762:A:H2	1.21	0.88
1:1G:1502:A:H2	1:1G:1505:G:H1	1.20	0.88
26:14:1992:G:N7	57:14:3588:HOH:O	2.07	0.88
27:16:21:G:H1	27:16:62:C:H42	1.21	0.88
26:14:1061:U:H5'	26:14:1070:A:H1'	1.53	0.88
26:1H:450:G:OP2	57:1H:3739:HOH:O	1.91	0.88
36:78:52:GLU:HG3	36:78:57:THR:HA	1.56	0.87
26:14:2499:C:OP2	57:14:3526:HOH:O	1.89	0.87
26:14:576:U:OP1	57:14:3614:HOH:O	1.91	0.87
26:14:631:A:OP2	54:M5:47:LYS:NZ	2.07	0.87
26:14:880:G:O6	26:14:894:C:N4	2.07	0.87
26:1H:586:A:OP2	57:1H:3733:HOH:O	1.91	0.87
36:78:101:VAL:HG12	36:78:106:LEU:HD12	1.55	0.87
26:1H:731:C:OP2	57:1H:3621:HOH:O	1.93	0.87
30:31:40:GLN:HE22	30:31:182:ASN:HB2	1.38	0.87
1:1G:1178:G:H5''	9:82:93:ARG:HH22	1.37	0.87
1:1G:1348:U:H3	1:1G:1374:A:H2	1.13	0.87
26:14:1616:A:O2'	57:14:3941:HOH:O	1.93	0.87
26:1H:2096:U:H3	26:1H:2193:G:H1	1.22	0.87
54:Q8:37:SER:HA	54:Q8:40:GLU:HB3	1.57	0.87
26:14:1678:G:N2	26:14:1989:G:H22	1.73	0.87
26:1H:880:G:N2	26:1H:897:C:N3	2.22	0.87
26:1H:1062:G:N2	26:1H:1076:C:O2	2.07	0.86
1:1G:1305:G:H22	1:1G:1331:G:H2'	1.40	0.86
2:1E:17:PHE:HB3	2:1E:44:LEU:HD21	1.57	0.86
26:14:2448:A:OP2	57:14:3526:HOH:O	1.92	0.86

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:676:A:H8	26:1H:2069:G:H21	1.15	0.86
26:1H:2843:G:H1	26:1H:2874:C:H42	1.23	0.86
26:1H:571:A:OP2	57:1H:3713:HOH:O	1.93	0.86
1:1G:147:G:H1	1:1G:175:C:H42	1.24	0.86
41:85:49:HIS:HA	41:85:52:ARG:HB2	1.56	0.86
26:1H:1021:A:H62	26:1H:1141:U:H3	1.23	0.86
26:1H:607:U:H3	26:1H:621:A:H2	1.23	0.86
1:13:963:G:N3	10:1I:55:LYS:NZ	2.23	0.86
26:1H:2334:G:O6	47:I8:74:ARG:NH2	2.07	0.86
37:45:25:ASP:HB3	37:45:102:VAL:HG23	1.55	0.86
26:14:1614:A:OP1	57:14:3506:HOH:O	1.94	0.85
1:13:877:C:OP1	8:7E:88:LYS:NZ	2.09	0.85
26:1H:733:G:N7	57:1H:3838:HOH:O	2.09	0.85
26:1H:585:G:OP2	57:1H:3692:HOH:O	1.94	0.85
27:16:0:A:H62	27:16:119:A:H61	1.25	0.85
28:19:242:ARG:O	57:19:307:HOH:O	1.93	0.85
26:14:833:U:O2	36:35:55:ARG:NH1	2.10	0.85
26:1H:1899:G:H22	26:1H:1902:C:H41	1.20	0.85
39:A8:11:LYS:HD3	39:A8:91:PRO:HD3	1.58	0.85
45:G8:83:THR:HG22	45:G8:84:ARG:HG3	1.59	0.85
48:J8:83:GLU:HG2	48:J8:85:LEU:H	1.40	0.85
3:22:70:VAL:HG21	3:22:76:VAL:HG11	1.59	0.85
26:14:1614:A:OP1	57:14:3509:HOH:O	1.95	0.84
26:14:1856:G:H1	26:14:1886:C:H42	1.20	0.84
1:1G:539:A:OP2	12:3A:115:LYS:NZ	2.09	0.84
26:1H:748:G:OP2	57:1H:4036:HOH:O	1.95	0.84
26:1H:1342:A:OP2	57:1H:3789:HOH:O	1.94	0.84
48:J8:92:LYS:HA	48:J8:95:LEU:HB2	1.60	0.84
43:E8:79:GLY:HA3	43:E8:100:THR:HG22	1.60	0.84
26:14:274:G:N7	26:14:363:G:N2	2.26	0.84
1:13:1309:G:OP2	13:4I:99:ARG:NH2	2.09	0.84
1:13:177:C:OP1	20:BI:65:LYS:NZ	2.11	0.84
26:1H:1731:G:H2'	26:1H:1732:A:H8	1.42	0.84
13:4I:54:VAL:HA	13:4I:57:ARG:HB3	1.59	0.84
26:14:2598:A:OP1	57:14:3565:HOH:O	1.96	0.84
26:1H:249:C:OP1	57:1H:3617:HOH:O	1.93	0.84
22:1L:47:C:N3	22:1L:56:G:N2	2.25	0.84
26:1H:563:G:OP2	57:1H:3610:HOH:O	1.95	0.83
26:14:910:A:H62	37:45:12:GLN:HA	1.43	0.83
1:1G:1503:A:H5'	1:1G:1531:A:H1'	1.60	0.83
36:78:50:ARG:HB2	36:78:50:ARG:HH21	1.42	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1310:G:OP2	53:P8:9:ARG:NH1	2.11	0.83
1:1G:976:G:N2	1:1G:1362(A):C:OP2	2.10	0.83
26:1H:2406:U:OP1	57:1H:4139:HOH:O	1.94	0.83
26:1H:2580:U:H4'	29:21:130:GLY:HA3	1.58	0.83
1:13:1318:A:H1'	19:AI:37:ARG:HH21	1.44	0.83
26:1H:67:U:H3	26:1H:74:A:H2	1.27	0.83
26:1H:1496:A:H8	26:1H:1577:C:HO2'	1.24	0.83
26:1H:602:G:HO2'	26:1H:604:G:HO2'	1.20	0.83
26:14:857:C:H4'	47:E5:23:VAL:HG21	1.61	0.83
26:14:958:U:OP2	37:45:14:ARG:NH1	2.12	0.82
10:1A:47:PHE:HB2	10:1A:63:PHE:HB2	1.61	0.82
26:1H:49:A:N7	26:1H:120:U:H5	1.77	0.82
26:1H:2577:A:OP1	57:1H:3668:HOH:O	1.97	0.82
26:1H:780:G:H21	26:1H:783:A:H62	1.27	0.82
23:2K:2:C:H2'	23:2K:3:C:C6	2.14	0.82
26:1H:2562:U:H1'	35:68:23:ARG:HH11	1.43	0.82
30:31:9:ILE:HD11	30:31:125:LEU:H	1.45	0.82
19:AI:42:PRO:HD3	51:M8:63:TYR:HE2	1.45	0.82
26:1H:2878:U:O4	57:1H:3959:HOH:O	1.95	0.82
27:1J:48:A:H4'	39:65:95:HIS:HD2	1.45	0.82
26:14:1365:A:O2'	48:F5:11:ARG:NH2	2.12	0.82
26:14:1364:G:OP2	48:F5:2:SER:N	2.12	0.82
26:1H:1253:A:N7	57:1H:4123:HOH:O	2.13	0.82
1:13:1348:U:H3	1:13:1374:A:H2	1.27	0.82
1:13:963:G:H1	1:13:972:C:H42	1.25	0.82
22:1L:87:A:H8	26:14:2583:G:H21	1.24	0.82
2:1E:189:ASP:HB2	2:1E:205:ASP:HB3	1.62	0.82
1:1G:1340:A:O2'	23:2L:32:C:O2'	1.98	0.82
26:1H:654(B):C:H2'	26:1H:654(C):G:H8	1.44	0.82
48:J8:48:LYS:HB3	48:J8:49:VAL:HA	1.60	0.82
19:AI:40:ILE:O	51:M8:63:TYR:OH	1.98	0.82
2:1E:185:ILE:HG22	2:1E:199:TYR:HB2	1.62	0.81
26:14:1664:A:OP2	57:14:3587:HOH:O	1.96	0.81
3:22:134:ILE:HG13	3:22:153:VAL:HG21	1.63	0.81
30:39:25:PRO:HB2	30:39:27:GLU:H	1.45	0.81
26:14:993:G:OP1	41:85:50:ARG:NH2	2.12	0.81
26:1H:1187:G:O6	57:1H:3720:HOH:O	1.97	0.81
35:25:35:VAL:HG11	35:25:103:ALA:HB3	1.63	0.81
54:M5:59:LYS:HZ2	54:M5:59:LYS:HA	1.45	0.81
26:14:2597:G:O3'	57:14:3564:HOH:O	1.98	0.81
22:1K:87:A:H8	26:1H:2583:G:H21	1.24	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:399:G:OP2	57:1H:3823:HOH:O	1.96	0.81
29:21:29:GLY:H	29:21:51:PHE:HE1	1.25	0.81
8:72:73:ASP:HB2	8:72:75:ARG:HH21	1.45	0.81
46:H8:19:ARG:HH11	46:H8:84:GLU:HB2	1.45	0.81
30:31:185:ASP:OD1	30:31:188:ARG:NH1	2.13	0.81
1:13:1422:G:H5''	35:68:48:PRO:HB3	1.60	0.81
38:98:26:LYS:HG2	38:98:70:LEU:HD22	1.62	0.81
26:1H:993:G:OP1	41:C8:50:ARG:NH2	2.14	0.81
26:1H:2213:U:O2	48:J8:52:ARG:NH2	2.14	0.81
29:29:181:LEU:HD11	40:75:7:ILE:HD11	1.62	0.81
8:7E:10:LEU:HD22	8:7E:83:ILE:HD11	1.63	0.81
1:13:1435:G:H2'	1:13:1436:U:C6	2.15	0.81
26:14:1754:C:OP1	40:75:96:ARG:NH1	2.14	0.81
5:4E:142:LEU:O	5:4E:143:ARG:NH1	2.14	0.81
23:2K:2:C:H2'	23:2K:3:C:H6	1.44	0.81
26:14:2245:U:H5''	26:14:2246:G:H5'	1.63	0.80
26:14:790:C:O5'	57:14:3663:HOH:O	1.99	0.80
26:14:787:U:OP1	57:14:3664:HOH:O	1.99	0.80
27:1J:11:C:OP2	27:1J:12:C:N4	2.15	0.80
29:21:116:VAL:HG11	29:21:138:PRO:HB3	1.62	0.80
29:21:1:MET:N	29:21:83:ASP:O	2.14	0.80
3:22:138:VAL:HG23	3:22:151:VAL:HG23	1.62	0.80
1:1G:542:G:OP1	4:32:10:ARG:NH2	2.15	0.80
26:14:1434:A:H61	26:14:1558:A:H62	1.26	0.80
1:1G:928:G:O2'	1:1G:1533:C:OP1	1.99	0.80
26:14:1225:C:H4'	42:95:85:LYS:HA	1.64	0.80
26:14:780:G:H21	26:14:783:A:H62	1.29	0.80
26:1H:730:C:H3'	57:1H:3621:HOH:O	1.81	0.80
8:72:64:LYS:HG2	8:72:79:VAL:HG21	1.63	0.80
31:49:11:TYR:OH	31:49:16:ARG:NH1	2.14	0.80
26:1H:1997:G:OP2	57:1H:3786:HOH:O	2.00	0.80
29:21:55:ASN:HB3	29:21:58:ARG:HD2	1.64	0.80
26:14:1022:G:H22	26:14:1142(A):A:H2	1.30	0.80
1:1G:353:A:H8	1:1G:353:A:H5'	1.44	0.80
43:A5:92:ARG:HG2	43:A5:92:ARG:HH11	1.46	0.80
19:AA:60:VAL:HG21	19:AA:74:PHE:HB3	1.64	0.80
26:1H:1634:A:OP2	57:1H:3951:HOH:O	1.99	0.80
41:85:92:ARG:HD3	41:85:94:ASN:HB3	1.63	0.80
54:Q8:47:LYS:NZ	54:Q8:47:LYS:HA	1.97	0.80
26:1H:2837:G:N7	57:1H:3962:HOH:O	2.15	0.80
32:51:25:LYS:HG2	32:51:34:GLU:HG2	1.64	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1225:C:O2'	42:95:85:LYS:N	2.15	0.79
2:1E:21:ARG:HB2	2:1E:39:ILE:HA	1.64	0.79
26:1H:2781:A:H5''	26:1H:2782:G:H5'	1.62	0.79
26:1H:943:U:OP2	57:1H:4133:HOH:O	1.98	0.79
33:61:8:PRO:HA	33:61:14:ASP:HA	1.64	0.79
50:L8:35:ARG:HB3	50:L8:37:LEU:HD21	1.64	0.79
26:14:2498:C:OP2	57:14:3526:HOH:O	1.99	0.79
26:14:674:G:H1'	30:39:74:ARG:HD3	1.61	0.79
4:3E:9:CYS:HB3	4:3E:32:ALA:HB2	1.64	0.79
26:14:2681:C:H5	26:14:2725:A:H62	1.31	0.79
28:19:13:ARG:NH1	28:19:16:MET:SD	2.55	0.79
26:1H:382:G:O6	57:1H:4008:HOH:O	2.01	0.79
30:31:6:VAL:HG11	30:31:119:ARG:HA	1.63	0.79
12:3A:84:LEU:HD12	12:3A:104:VAL:HG11	1.63	0.79
24:3K:83:U:H2'	24:3K:84:A:H5''	1.63	0.79
26:14:2016:U:OP1	57:14:3851:HOH:O	2.00	0.79
49:K8:8:LYS:HA	49:K8:11:GLU:HB3	1.61	0.79
40:75:92:GLY:HA2	40:75:116:ALA:HA	1.64	0.79
28:11:242:ARG:H	28:11:242:ARG:HD2	1.47	0.79
40:B8:3:ARG:HB2	40:B8:6:LEU:HB2	1.63	0.79
1:1G:1255:G:O2'	1:1G:1258:G:O2'	2.00	0.79
26:1H:10:G:O2'	26:1H:2801:A:N3	2.15	0.79
30:31:66:PRO:O	30:31:67:GLN:HB3	1.82	0.79
2:12:67:THR:HG21	2:12:155:LEU:HD21	1.63	0.79
26:1H:880:G:H1	26:1H:897:C:H42	1.30	0.79
44:B5:8:ILE:O	49:G5:36:ARG:NH2	2.15	0.79
26:1H:1388:G:H2'	26:1H:1389:G:H8	1.48	0.78
26:14:2448:A:O5'	57:14:3635:HOH:O	2.01	0.78
26:14:570:G:O6	57:14:3631:HOH:O	2.01	0.78
28:11:273:ARG:O	28:11:273:ARG:NE	2.16	0.78
1:13:1452:C:O2'	1:13:1453:G:OP2	2.02	0.78
23:2L:31:C:O2	23:2L:41:G:N2	2.15	0.78
39:65:3:ARG:HH21	39:65:4:LEU:HB2	1.49	0.78
49:K8:47:ASN:O	49:K8:49:LYS:N	2.15	0.78
26:1H:1689:A:H62	26:1H:1698:A:H2	1.30	0.78
26:1H:1980:G:O2'	26:1H:1982:C:OP2	2.02	0.78
31:41:64:THR:HG22	31:41:66:GLN:H	1.47	0.78
31:49:118:ARG:HB3	31:49:181:ARG:HG3	1.66	0.78
41:85:90:VAL:HG22	42:95:39:LEU:HB2	1.63	0.78
26:14:2315:G:OP1	31:49:36:LYS:NZ	2.14	0.78
27:16:40:U:O2	27:16:45:A:N6	2.17	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:286:G:N7	57:1G:1824:HOH:O	2.17	0.78
26:1H:1643:G:N7	57:1H:3753:HOH:O	2.16	0.78
26:1H:796:C:H2'	26:1H:797:C:C6	2.18	0.78
6:5E:38:GLU:HB2	6:5E:64:GLN:HB3	1.65	0.78
26:14:1041:C:H42	26:14:1114:G:H22	1.28	0.78
1:13:1321:C:H3'	1:13:1322:C:H5''	1.66	0.78
26:1H:363(B):G:H2'	26:1H:363(C):G:C8	2.18	0.78
1:13:410:G:O6	57:13:1819:HOH:O	2.00	0.78
26:14:1030:G:OP2	37:45:128:LYS:NZ	2.16	0.78
37:45:34:LEU:HD11	37:45:129:THR:HB	1.66	0.78
1:13:975:A:H4'	1:13:976:G:H5''	1.66	0.78
26:14:1249:U:OP1	57:14:3520:HOH:O	2.02	0.78
26:14:2287:A:N6	26:14:2344:U:H3	1.81	0.78
26:1H:1534:G:H2'	26:1H:1535:U:H4'	1.65	0.78
26:1H:963:U:OP1	57:1H:3697:HOH:O	2.02	0.78
51:I5:22:ILE:HG12	51:I5:23:GLU:H	1.48	0.78
26:14:125:G:H5''	53:L5:19:ARG:HD3	1.64	0.78
24:3K:7:G:H3'	24:3K:8:U:H5''	1.65	0.77
40:75:1:MET:HB3	40:75:6:LEU:HB2	1.65	0.77
30:31:6:VAL:N	30:31:24:LEU:O	2.16	0.77
26:1H:142:G:H1'	44:F8:37:THR:HG21	1.66	0.77
45:G8:94:LYS:HA	45:G8:94:LYS:HZ3	1.50	0.77
46:D5:94:GLU:HB3	46:D5:96:VAL:HG23	1.66	0.77
26:14:1856:G:N2	26:14:1886:C:N3	2.28	0.77
1:1G:1435:G:H2'	1:1G:1436:U:C6	2.20	0.77
26:1H:2256:G:N7	57:1H:3880:HOH:O	2.16	0.77
35:25:73:ASP:OD2	40:75:32:TYR:OH	2.02	0.77
42:D8:17:GLY:N	42:D8:96:ILE:O	2.17	0.77
1:13:1003:G:N2	1:13:1037:C:N3	2.32	0.77
1:1G:987:G:N2	1:1G:1218:C:N3	2.31	0.77
29:29:33:VAL:HG12	29:29:89:ASP:HB3	1.66	0.77
3:2E:76:VAL:HG21	3:2E:103:VAL:HG21	1.65	0.77
26:1H:442:G:H1'	30:31:48:THR:HG21	1.65	0.77
5:4E:102:ALA:HB1	5:4E:106:PRO:HG2	1.67	0.77
33:61:144:VAL:HG13	33:61:145:VAL:HG23	1.67	0.77
49:G5:25:VAL:HG12	49:G5:60:LEU:HD23	1.65	0.77
2:12:54:THR:HG21	2:12:201:ILE:HD11	1.66	0.77
20:BA:82:SER:HG	20:BA:86:ARG:HH22	1.31	0.77
46:D5:19:ARG:HH11	46:D5:84:GLU:HB2	1.48	0.77
1:1G:1263:C:N4	1:1G:1272:G:O6	2.17	0.77
24:3K:13:G:H1	24:3K:23:A:HO2'	1.32	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:8E:28:VAL:HG22	9:8E:63:ILE:HB	1.65	0.77
26:14:1403:C:OP1	26:14:1522:G:N2	2.18	0.77
26:14:1828:G:OP2	57:14:3531:HOH:O	2.01	0.77
26:1H:2112:G:H22	26:1H:2169:A:H61	1.32	0.77
52:N8:42:PRO:O	52:N8:44:THR:HG22	1.85	0.77
1:13:1226:C:H2'	13:4I:103:THR:HB	1.65	0.76
26:1H:252:G:OP2	36:78:50:ARG:NH1	2.18	0.76
26:1H:259:G:O2'	26:1H:621:A:O2'	1.98	0.76
26:1H:761:A:OP2	57:1H:3836:HOH:O	2.02	0.76
1:1G:1298:C:OP2	7:62:114:ARG:NH2	2.18	0.76
45:C5:73:ARG:NH2	45:C5:81:LYS:O	2.18	0.76
54:Q8:53:PRO:HA	54:Q8:55:ALA:N	2.00	0.76
26:14:2718:G:N7	57:14:3996:HOH:O	2.18	0.76
8:7E:41:ARG:NH1	8:7E:123:GLU:OE1	2.17	0.76
28:19:181:GLU:HG3	28:19:272:ALA:HB3	1.67	0.76
1:1G:443:C:H42	1:1G:491:G:H1	1.30	0.76
10:1I:6:ILE:HG22	10:1I:98:ILE:HG12	1.68	0.76
39:65:106:ARG:O	39:65:106:ARG:NH1	2.18	0.76
39:65:84:GLN:HA	39:65:110:LEU:HD12	1.65	0.76
22:1K:49:C:N4	22:1K:55:G:O6	2.19	0.76
31:41:97:ASP:O	31:41:100:TRP:N	2.17	0.76
50:L8:12:PRO:O	50:L8:20:LYS:NZ	2.19	0.76
31:41:37:VAL:HG22	31:41:159:VAL:HG12	1.67	0.76
36:78:62:LEU:HD13	54:Q8:23:VAL:HG11	1.66	0.76
1:13:438:G:N7	57:13:1866:HOH:O	2.18	0.76
1:13:538:G:H5''	12:3I:114:LYS:HB2	1.67	0.76
2:1E:178:ARG:NH1	2:1E:196:LEU:O	2.18	0.76
29:29:50:GLY:HA2	29:29:78:LEU:HB3	1.67	0.76
4:32:157:LEU:O	4:32:161:ASN:ND2	2.16	0.76
17:8A:88:TYR:OH	17:8A:92:ARG:NH1	2.19	0.76
40:B8:50:ILE:HD11	40:B8:102:ILE:HD11	1.67	0.76
46:H8:31:ARG:NH1	46:H8:94:GLU:OE1	2.19	0.76
1:13:591:U:H2'	1:13:592:G:H8	1.51	0.76
26:1H:2038:G:O6	57:1H:3657:HOH:O	2.03	0.76
29:29:8:LYS:HB3	29:29:193:GLY:H	1.50	0.76
33:69:81:VAL:HG23	33:69:143:SER:HB2	1.68	0.76
43:E8:13:SER:HB3	43:E8:16:LYS:HD2	1.66	0.76
26:14:2837:G:N7	57:14:3966:HOH:O	2.17	0.76
26:14:517:C:OP1	52:J5:16:ARG:NH2	2.18	0.76
26:14:1627:G:OP1	57:14:3902:HOH:O	2.04	0.76
26:14:586:A:OP2	57:14:3517:HOH:O	2.03	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1454:G:OP1	20:BA:39:LYS:NZ	2.17	0.76
1:1G:152:A:N6	1:1G:170:U:O2	2.19	0.76
34:15:4:TYR:O	41:85:64:ARG:NH1	2.19	0.75
27:1J:40:U:H5	51:I5:1:MET:HB3	1.50	0.75
1:13:262:A:H2'	1:13:263:A:C8	2.21	0.75
26:14:124:G:N7	57:14:3849:HOH:O	2.20	0.75
1:1G:1047:G:H1	1:1G:1210:C:H42	1.34	0.75
26:1H:2210:G:H5'	26:1H:2211:G:N7	2.01	0.75
26:1H:2494:G:O2'	37:88:80:GLU:HB3	1.86	0.75
26:1H:141:A:H8	26:1H:1595:G:H21	1.34	0.75
34:15:14:VAL:HA	34:15:135:PRO:HD2	1.67	0.75
26:1H:2035:G:OP1	57:1H:3659:HOH:O	2.03	0.75
44:B5:36:LYS:HG2	44:B5:54:VAL:HB	1.69	0.75
28:11:71:ASP:N	28:11:71:ASP:OD1	2.17	0.75
1:13:1027:C:O2	1:13:1035:A:N6	2.19	0.75
26:1H:1525:G:H2'	26:1H:1526:G:H8	1.51	0.75
26:1H:398:G:N7	57:1H:4090:HOH:O	2.19	0.75
29:29:36:ARG:NH1	29:29:85:ASN:OD1	2.19	0.75
26:1H:730:C:OP2	57:1H:3621:HOH:O	2.05	0.75
27:1J:116:G:H5''	39:65:55:ALA:HB2	1.68	0.75
42:95:43:GLU:O	42:95:45:THR:OG1	2.04	0.75
26:14:1048:A:N6	26:14:1112:G:O2'	2.19	0.75
31:49:67:LYS:HE3	51:I5:5:ILE:HB	1.67	0.75
28:11:108:PRO:HG3	28:11:143:HIS:CE1	2.22	0.75
26:14:1771:C:HO2'	26:14:1786:A:H8	1.34	0.75
26:14:943:U:OP2	36:35:36:LYS:NZ	2.16	0.75
1:1G:1497:G:H2'	1:1G:1498:U:H5'	1.69	0.75
26:1H:860:U:H5	26:1H:917:A:C2	2.03	0.75
29:21:127:ASP:OD1	57:21:401:HOH:O	2.03	0.75
12:3I:82:VAL:HG13	12:3I:105:TYR:HB3	1.68	0.75
13:4I:58:GLU:O	13:4I:62:ASN:ND2	2.18	0.75
26:1H:910:A:H62	37:88:12:GLN:HA	1.52	0.75
30:39:25:PRO:HB3	30:39:28:ILE:HG23	1.67	0.75
47:I8:11:ARG:O	47:I8:14:ARG:NH2	2.20	0.75
1:1G:1259:C:N4	1:1G:1260:C:O2	2.19	0.74
24:3L:32:C:O2	24:3L:40:G:N1	2.19	0.74
40:75:54:ARG:HH11	40:75:54:ARG:HB3	1.49	0.74
20:BA:69:GLY:O	20:BA:73:HIS:NE2	2.20	0.74
26:14:2141:G:N1	26:14:2150:U:O2	2.19	0.74
26:1H:2035:G:OP1	57:1H:3656:HOH:O	2.04	0.74
26:1H:732:C:OP2	57:1H:3841:HOH:O	2.05	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:1L:37:G:H1	25:4L:50:U:H3	1.34	0.74
29:29:111:ARG:HH11	29:29:111:ARG:HG2	1.52	0.74
24:3K:46:G:H2'	24:3K:47:C:H5'	1.68	0.74
33:61:92:VAL:HG13	33:61:120:ILE:HG23	1.69	0.74
1:1G:278:G:N7	17:8A:92:ARG:NH2	2.35	0.74
1:1G:664:G:H22	1:1G:741:G:H1	1.34	0.74
1:1G:91:C:H2'	1:1G:92:G:H8	1.52	0.74
26:1H:1899:G:H22	26:1H:1902:C:N4	1.84	0.74
41:85:98:LEU:HB2	41:85:102:GLU:HB2	1.69	0.74
26:1H:1359:A:H2	26:1H:1372:U:O4	1.70	0.74
26:1H:2032:G:H21	29:21:146:THR:HG23	1.50	0.74
29:21:57:LYS:HG3	29:21:59:VAL:HG12	1.69	0.74
42:95:85:LYS:HB3	42:95:87:HIS:H	1.51	0.74
52:J5:49:CYS:SG	52:J5:50:GLY:N	2.61	0.74
48:F5:92:LYS:HD3	48:F5:93:GLU:H	1.51	0.74
51:M8:37:SER:OG	51:M8:42:PHE:O	2.04	0.74
19:AI:6:LYS:HE3	51:M8:62:ARG:HB2	1.69	0.74
44:B5:41:ASN:HA	44:B5:44:GLU:HB2	1.69	0.74
40:B8:52:ILE:HG23	40:B8:61:PHE:HB3	1.68	0.74
27:16:15:A:H5'	27:16:16:G:C8	2.23	0.74
26:1H:1386:C:H2'	26:1H:1387:C:H6	1.52	0.74
26:1H:958:U:H5'	37:88:14:ARG:HD3	1.68	0.74
11:2I:91:ARG:NH2	11:2I:110:ASP:OD2	2.18	0.74
33:69:77:LEU:HD13	33:69:141:LYS:HD2	1.69	0.74
1:13:137:C:N4	1:13:226:G:H1	1.82	0.74
26:14:399:G:OP2	57:14:3703:HOH:O	2.06	0.74
26:14:2255:G:H21	47:E5:9:SER:H	1.34	0.74
28:19:181:GLU:HA	28:19:272:ALA:HB1	1.69	0.74
1:1G:988:G:N2	1:1G:1217:C:O2	2.17	0.74
26:1H:1058:U:H3	26:1H:1080:A:H61	1.34	0.74
26:1H:2140:C:H42	26:1H:2151:G:H1	1.35	0.74
13:4I:39:ILE:HD12	13:4I:56:LEU:HD23	1.70	0.74
26:14:329:G:OP2	45:C5:71:LYS:NZ	2.19	0.74
47:I8:46:LYS:HE3	47:I8:76:GLY:HA3	1.70	0.74
1:1G:222:U:H2'	1:1G:223:U:C6	2.22	0.73
26:1H:399:G:OP2	57:1H:3821:HOH:O	2.06	0.73
27:1J:90:C:OP2	37:45:16:ARG:NH2	2.21	0.73
22:1L:9:G:O2'	22:1L:10:G:N7	2.19	0.73
4:32:60:GLU:HG2	4:32:202:LEU:HB2	1.69	0.73
5:42:60:TYR:HB3	5:42:64:ARG:HH21	1.51	0.73
38:55:103:ARG:NH1	38:55:108:GLY:O	2.20	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:Q8:50:LEU:C	54:Q8:52:LYS:H	1.91	0.73
26:14:54:G:O6	57:14:3945:HOH:O	2.06	0.73
26:1H:620:G:H4'	26:1H:621:A:H5''	1.71	0.73
1:13:1129:C:H4'	1:13:1130:A:H5'	1.69	0.73
1:13:1226:C:O2'	13:4I:111:LYS:NZ	2.21	0.73
26:14:249:C:OP1	57:14:3513:HOH:O	2.05	0.73
35:25:64:ARG:HB2	35:25:83:ALA:HB3	1.70	0.73
1:1G:1329:A:H5'	13:4A:29:ARG:HE	1.53	0.73
2:12:75:LYS:HA	2:12:78:GLN:HB2	1.68	0.73
38:98:55:ALA:HA	38:98:80:PHE:CE1	2.24	0.73
44:B5:55:ASN:HB2	44:B5:80:ILE:HG13	1.69	0.73
46:H8:154:ASP:OD1	46:H8:154:ASP:N	2.21	0.73
26:14:1897:G:N7	57:14:3708:HOH:O	2.21	0.73
26:14:617:G:OP1	30:39:40:GLN:NE2	2.18	0.73
26:1H:1537:C:H2'	26:1H:1538:G:O4'	1.87	0.73
36:35:85:LEU:HB3	36:35:114:ILE:HD11	1.70	0.73
42:95:21:ARG:NH2	42:95:65:GLY:O	2.22	0.73
26:14:452:G:OP2	57:14:3655:HOH:O	2.06	0.73
1:1G:987:G:H1	1:1G:1218:C:H42	1.37	0.73
1:13:411:A:C4	1:13:413:G:H1'	2.23	0.73
26:14:2327:A:H2'	26:14:2328:A:C8	2.24	0.73
26:14:2645:G:H3'	26:14:2646:C:H5'	1.71	0.73
26:14:662:G:H5''	36:35:16:ARG:HG2	1.70	0.73
26:1H:1607:C:OP1	57:1H:4098:HOH:O	2.05	0.73
13:4A:3:ARG:HG2	13:4A:9:ILE:HG12	1.69	0.73
36:78:114:ILE:HD13	36:78:125:VAL:HG11	1.70	0.73
39:A8:37:ALA:HB2	39:A8:101:LEU:HD21	1.71	0.73
45:C5:99:CYS:SG	45:C5:100:ALA:N	2.62	0.73
47:E5:32:ARG:O	47:E5:34:GLY:N	2.21	0.73
1:13:372:C:H42	1:13:389:A:H62	1.37	0.73
26:14:2520:C:H41	26:14:2542:A:H62	1.33	0.73
39:65:3:ARG:NH2	39:65:4:LEU:HB2	2.04	0.73
26:14:363:G:H2'	26:14:363(A):A:H8	1.53	0.72
26:1H:2099:U:O2	26:1H:2190:G:N1	2.18	0.72
51:I5:55:ARG:HD2	51:I5:56:VAL:HG12	1.71	0.72
1:13:1305:G:H8	1:13:1305:G:OP2	1.72	0.72
1:13:664:G:H22	1:13:741:G:H1	1.34	0.72
26:14:573:G:O2'	26:14:574:C:H3'	1.89	0.72
26:1H:945:A:OP2	57:1H:3866:HOH:O	2.07	0.72
9:82:112:LYS:HE3	9:82:118:LYS:H	1.52	0.72
26:1H:1678:G:N2	26:1H:1989:G:H22	1.87	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:59:152:ARG:HG3	32:59:153:LYS:HB2	1.71	0.72
32:59:81:GLU:O	32:59:138:LYS:NZ	2.22	0.72
37:88:89:ASN:O	37:88:91:GLU:HB2	1.89	0.72
45:C5:17:SER:HB2	45:C5:71:LYS:HD2	1.69	0.72
26:14:2355:C:H1'	47:E5:39:ARG:HH21	1.54	0.72
26:14:1525:G:H2'	26:14:1526:G:H8	1.55	0.72
26:14:2239:G:OP2	57:14:3502:HOH:O	2.08	0.72
26:14:2503:A:OP1	57:14:3614:HOH:O	2.06	0.72
26:1H:2287:A:H61	26:1H:2344:U:H3	1.37	0.72
29:29:61:ARG:O	29:29:63:LEU:N	2.20	0.72
23:2L:86:C:H3'	23:2L:87:A:H5''	1.71	0.72
40:75:65:LYS:HE3	40:75:67:SER:HB2	1.70	0.72
47:E5:21:LEU:HD11	47:E5:41:ARG:HH11	1.52	0.72
2:12:107:THR:O	2:12:110:GLN:NE2	2.22	0.72
23:2L:87:A:H2'	23:2L:87:A:N3	2.03	0.72
16:7A:53:VAL:HG13	16:7A:79:VAL:HG22	1.69	0.72
39:A8:28:VAL:HG11	39:A8:98:VAL:HG13	1.71	0.72
29:21:105:THR:HB	29:21:197:ILE:HG23	1.72	0.72
31:49:121:ASN:O	31:49:131:TYR:OH	2.07	0.72
5:4E:10:MET:HB2	5:4E:32:VAL:HG22	1.69	0.72
1:13:451:A:OP1	1:13:481:G:N2	2.21	0.72
26:14:2499:C:OP1	57:14:3631:HOH:O	2.06	0.72
26:1H:528:A:O2'	26:1H:529:A:H5''	1.90	0.72
4:32:8:VAL:O	4:32:11:LEU:N	2.20	0.72
1:13:601:C:H2'	1:13:602:A:H8	1.55	0.72
10:1I:35:SER:HB2	10:1I:73:ASP:HB2	1.72	0.72
46:H8:16:SER:O	46:H8:20:ARG:NH1	2.22	0.72
26:14:2793:G:N2	26:14:2804:C:O2	2.22	0.72
26:1H:2327:A:H2'	26:1H:2328:A:C8	2.23	0.72
34:58:30:ILE:HG23	34:58:52:VAL:HG11	1.72	0.72
19:AA:41:VAL:HG12	19:AA:42:PRO:HD2	1.70	0.72
28:11:72:LYS:HG2	28:11:103:ARG:HH21	1.53	0.72
1:13:1502:A:H2	1:13:1505:G:N1	1.82	0.72
26:14:1060:U:H4'	26:14:1061:U:H5''	1.72	0.72
26:14:1665:A:O2'	35:25:1:MET:N	2.19	0.72
26:14:491:G:H2'	26:14:492:A:C8	2.24	0.72
26:14:796:C:H2'	26:14:797:C:C6	2.25	0.72
11:2A:29:ILE:HG22	11:2A:44:SER:HB2	1.72	0.72
4:32:105:VAL:HG13	4:32:110:PHE:HB2	1.72	0.72
26:14:831:G:N7	57:14:3685:HOH:O	2.22	0.71
26:1H:1479:G:N7	26:1H:1510:A:N6	2.38	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2068:U:H3	26:1H:2430:A:H2	1.36	0.71
43:A5:72:LYS:HB3	43:A5:106:ILE:HG13	1.69	0.71
1:1G:192:U:O4'	20:BA:103:GLY:HA2	1.90	0.71
1:1G:971:G:N2	1:1G:1363:A:OP2	2.22	0.71
3:22:59:ARG:HG2	3:22:64:VAL:HG23	1.70	0.71
23:2L:71:U:H5''	23:2L:72:C:H5	1.55	0.71
32:51:86:GLU:HG3	32:51:165:ALA:HB3	1.71	0.71
17:8I:76:LEU:HD21	17:8I:79:SER:HB3	1.71	0.71
26:14:2153:G:N2	26:14:2154:G:O6	2.21	0.71
26:14:1022:G:O2'	26:14:1023:U:OP2	2.04	0.71
26:1H:2632:A:HO2'	26:1H:2811:G:HO2'	1.34	0.71
19:AA:78:ARG:HD3	19:AA:78:ARG:H	1.53	0.71
46:H8:131:ARG:NH2	46:H8:132:ASN:OD1	2.23	0.71
26:14:1250:G:OP1	57:14:3522:HOH:O	2.08	0.71
26:14:1377:G:OP2	57:14:3539:HOH:O	2.08	0.71
26:14:172:C:H2'	26:14:173:G:H8	1.54	0.71
2:1E:60:ASP:HB3	2:1E:64:ARG:HH12	1.56	0.71
26:1H:2309:A:C5	26:1H:2310:A:H8	2.08	0.71
23:2L:36:A:H2'	23:2L:37:G:H8	1.55	0.71
30:39:116:ASP:OD2	36:35:1:MET:N	2.24	0.71
34:58:96:GLU:O	34:58:98:VAL:N	2.23	0.71
26:14:270(N):G:OP1	33:69:57:ARG:NH1	2.23	0.71
26:14:1678:G:H22	26:14:1989:G:H22	1.37	0.71
26:14:2243:U:OP1	57:14:3643:HOH:O	2.07	0.71
26:1H:2884:U:O2	52:N8:52:TYR:OH	2.04	0.71
29:29:55:ASN:O	29:29:57:LYS:NZ	2.16	0.71
9:82:45:ALA:O	9:82:78:LYS:NZ	2.24	0.71
26:14:1434:A:H61	26:14:1558:A:N6	1.88	0.71
26:14:620:G:H4'	26:14:621:A:H5''	1.71	0.71
26:1H:1525:G:H2'	26:1H:1526:G:C8	2.25	0.71
30:39:123:LEU:O	30:39:125:LEU:N	2.23	0.71
40:75:51:ARG:HG2	40:75:98:LYS:HD2	1.71	0.71
26:14:1428:C:N4	26:14:1570:A:OP2	2.20	0.71
27:1J:80:U:H2'	27:1J:81:G:H21	1.56	0.71
42:95:85:LYS:HB3	42:95:87:HIS:N	2.04	0.71
20:BI:69:GLY:O	20:BI:73:HIS:NE2	2.24	0.71
44:F8:65:ARG:HG3	44:F8:67:GLY:H	1.55	0.71
1:1G:458:C:N3	1:1G:474:G:N2	2.38	0.71
1:1G:998:G:H22	1:1G:1043:C:H42	1.38	0.71
8:7E:6:ILE:HB	8:7E:85:ARG:HH12	1.56	0.71
42:95:35:LEU:O	42:95:37:VAL:HG22	1.90	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:586:A:OP2	57:14:3519:HOH:O	2.08	0.71
1:1G:827:U:H3	1:1G:872:A:H62	1.39	0.71
39:65:3:ARG:HE	39:65:4:LEU:H	1.38	0.71
26:14:1556:C:H2'	26:14:1557:C:H6	1.54	0.70
28:19:93:ALA:HB3	28:19:105:ILE:HG22	1.73	0.70
1:1G:359:U:H2'	1:1G:360:A:C8	2.26	0.70
26:1H:639:U:H3	26:1H:649:G:H1	1.36	0.70
26:1H:2636:U:OP1	29:21:79:ARG:HA	1.91	0.70
1:1G:1118:C:OP1	9:82:104:ARG:NH1	2.23	0.70
20:BI:73:HIS:HB3	20:BI:74:LYS:HD2	1.72	0.70
26:14:2469:A:H2	26:14:2481:G:H21	1.39	0.70
1:1G:45:U:H2'	1:1G:46:G:C8	2.25	0.70
11:2I:22:HIS:HB3	11:2I:29:ILE:HG23	1.73	0.70
17:8A:12:SER:HB3	17:8A:20:THR:HB	1.73	0.70
42:D8:60:GLU:HB2	42:D8:97:LYS:HE2	1.74	0.70
1:13:1454:G:OP1	20:BI:39:LYS:NZ	2.17	0.70
26:1H:307:G:N7	57:1H:3993:HOH:O	2.24	0.70
9:82:53:VAL:HG13	9:82:95:LYS:HE3	1.73	0.70
27:16:90:C:H5'	37:88:18:LYS:HA	1.73	0.70
28:11:182:LEU:H	28:11:272:ALA:HB3	1.55	0.70
1:13:166:G:H2'	1:13:167:G:H8	1.56	0.70
26:14:2111:C:N3	26:14:2118:U:O2'	2.21	0.70
1:1G:1219:U:OP1	14:5A:19:ARG:NH1	2.25	0.70
30:39:63:LYS:HE2	30:39:67:GLN:HB3	1.74	0.70
4:3E:107:ARG:HH22	4:3E:194:LEU:HD22	1.55	0.70
26:14:1864:U:OP1	26:14:2410:G:O2'	2.09	0.70
10:1A:48:THR:OG1	10:1A:62:HIS:ND1	2.21	0.70
22:1L:22:G:N2	22:1L:59:U:O4'	2.24	0.70
3:2E:91:LEU:HB2	3:2E:99:VAL:HG21	1.73	0.70
9:82:50:LEU:HB3	9:82:56:LEU:HA	1.73	0.70
28:19:68:LYS:HB3	28:19:70:TRP:CH2	2.27	0.70
2:1E:87:ARG:NH2	2:1E:220:ASP:OD1	2.24	0.70
26:1H:1164:G:H2'	26:1H:1165:U:C6	2.25	0.70
26:1H:1437:C:H2'	26:1H:1438:U:H6	1.56	0.70
39:65:27:SER:HA	39:65:88:ASP:HB2	1.73	0.70
15:6I:17:ARG:HD3	15:6I:26:GLU:HG3	1.73	0.70
40:B8:6:LEU:HA	40:B8:9:LEU:HB2	1.73	0.70
54:Q8:47:LYS:HZ2	54:Q8:47:LYS:HA	1.56	0.70
1:13:1189:C:O2	57:13:1842:HOH:O	2.05	0.70
26:14:2588:G:OP1	57:14:3567:HOH:O	2.09	0.70
36:35:94:GLU:HG3	36:35:124:LYS:HG2	1.72	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:3K:22:G:N7	24:3K:59:U:N3	2.39	0.70
9:8E:97:LYS:HD3	9:8E:102:LEU:HB2	1.73	0.70
9:8E:26:VAL:HG13	9:8E:61:ALA:HB3	1.73	0.70
26:14:1636:C:OP2	57:14:3598:HOH:O	2.08	0.70
1:1G:1240:U:H1'	7:62:38:LEU:HD21	1.73	0.70
26:1H:1839:G:OP2	57:1H:3833:HOH:O	2.09	0.70
26:1H:520:G:OP2	57:1H:3899:HOH:O	2.09	0.70
23:2K:37:G:H1	25:4K:46:U:H3	1.37	0.70
1:13:1086:U:H3	1:13:1099:G:H22	1.40	0.70
26:14:1416:G:O2'	26:14:1417:C:O5'	2.09	0.70
26:14:2557:G:H2'	26:14:2558:C:C6	2.26	0.70
26:1H:913:U:O4	57:1H:3774:HOH:O	2.07	0.70
5:4E:91:LEU:HD12	5:4E:120:THR:HG22	1.72	0.70
37:88:130:LYS:NZ	46:H8:81:ARG:HG2	2.07	0.70
33:61:110:ASP:HB2	33:61:112:LYS:H	1.57	0.70
44:B5:49:VAL:HB	44:B5:83:VAL:HG21	1.73	0.70
2:12:21:ARG:HA	2:12:39:ILE:HA	1.74	0.69
1:13:21:G:OP1	57:13:1807:HOH:O	2.10	0.69
26:1H:1253:A:N6	57:1H:4120:HOH:O	2.25	0.69
26:1H:2023:G:H5'	26:1H:2617:C:H4'	1.72	0.69
12:3I:24:VAL:HB	12:3I:27:LEU:HD12	1.73	0.69
32:51:20:ALA:HB3	32:51:23:ARG:HG3	1.74	0.69
17:8I:45:HIS:HB2	17:8I:65:ILE:HD13	1.74	0.69
28:19:76:PRO:HB2	28:19:116:GLN:OE1	1.92	0.69
26:1H:1047:G:O2'	26:1H:1111:A:N6	2.25	0.69
26:1H:429:A:OP2	57:1H:4126:HOH:O	2.09	0.69
35:25:63:VAL:HG12	35:25:106:LEU:HD11	1.73	0.69
34:58:32:THR:HG22	34:58:37:LYS:HB2	1.73	0.69
1:1G:588:G:H1	1:1G:651:C:H42	1.41	0.69
26:1H:831:G:OP1	57:1H:3705:HOH:O	2.09	0.69
32:59:26:VAL:HG12	32:59:33:LEU:H	1.54	0.69
1:1G:560:U:O2'	1:1G:561:U:OP2	2.09	0.69
26:1H:2593:U:H2'	26:1H:2594:C:C6	2.27	0.69
3:22:70:VAL:HG12	3:22:72:LYS:H	1.56	0.69
51:M8:52:THR:OG1	51:M8:53:GLU:N	2.26	0.69
1:13:624:C:H2'	1:13:625:G:H8	1.58	0.69
6:52:37:VAL:HA	6:52:65:VAL:HG12	1.75	0.69
41:C8:92:ARG:O	41:C8:94:ASN:N	2.25	0.69
26:1H:1441:G:H2'	26:1H:1442:G:H8	1.58	0.69
26:1H:674:G:H1'	30:31:74:ARG:HD3	1.73	0.69
2:1E:7:VAL:HG11	2:1E:217:ARG:HH11	1.57	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:991:U:O4	1:1G:1212:U:O2'	2.10	0.69
1:1G:992:U:H3	1:1G:1044:A:H62	1.41	0.69
26:1H:1858:G:H2'	26:1H:1883:G:H22	1.57	0.69
30:39:68:LYS:HG3	30:39:69:HIS:CE1	2.27	0.69
35:68:98:VAL:HG13	35:68:117:LEU:HB3	1.75	0.69
26:1H:102:G:OP1	49:K8:7:ARG:NH2	2.25	0.69
26:14:1382:G:N7	57:14:3696:HOH:O	2.26	0.69
26:1H:76:C:O2'	49:K8:62:THR:HG21	1.91	0.69
29:21:82:ARG:O	29:21:84:PHE:N	2.25	0.69
35:25:1:MET:HE3	35:25:67:LYS:HG2	1.74	0.69
29:29:65:GLY:HA2	29:29:66:HIS:HB3	1.75	0.69
30:31:6:VAL:HG21	30:31:119:ARG:HB2	1.74	0.69
30:39:178:PRO:HB2	30:39:201:VAL:HG11	1.75	0.69
33:61:40:THR:O	33:61:44:LEU:HB2	1.92	0.69
41:85:28:ARG:NH1	41:85:38:THR:OG1	2.26	0.69
26:14:593:G:H4'	54:M5:60:LEU:HD22	1.75	0.69
54:Q8:9:GLY:HA2	54:Q8:12:LYS:HB2	1.74	0.69
26:14:2304:G:N2	26:14:2313:C:N3	2.41	0.69
3:22:84:ILE:HD11	3:22:88:ARG:HH21	1.56	0.69
26:14:2513:G:O2'	29:29:154:LYS:NZ	2.25	0.69
31:41:41:GLN:HG2	31:41:155:MET:HB3	1.74	0.69
35:68:19:ILE:HG22	35:68:43:VAL:HA	1.73	0.69
26:1H:2485:G:H5''	37:88:46:GLN:HE21	1.58	0.69
53:L5:31:LEU:HD22	53:L5:42:LEU:HD13	1.74	0.69
1:1G:87:A:O2'	1:1G:88:C:OP2	2.10	0.69
26:1H:1434:A:H61	26:1H:1558:A:N6	1.91	0.69
10:1I:57:LYS:O	10:1I:60:ARG:NH1	2.24	0.69
22:1K:5:A:H61	22:1K:79:U:H3	1.39	0.69
3:22:94:LEU:HD12	3:22:95:THR:HG23	1.75	0.69
31:49:18:GLU:OE1	31:49:21:ARG:NH2	2.25	0.69
26:14:2104:G:O6	26:14:2185:C:N4	2.25	0.69
26:1H:1778:U:H2'	26:1H:1784:A:N6	2.08	0.69
26:1H:2127:G:N1	26:1H:2162:G:N3	2.41	0.69
26:1H:2135:A:H4'	26:1H:2160:G:H4'	1.75	0.69
7:6E:15:ASP:HB3	7:6E:20:ASP:H	1.58	0.69
20:BI:86:ARG:NH1	20:BI:86:ARG:O	2.25	0.69
26:14:1019:U:H3	26:14:1142(A):A:H62	1.40	0.68
26:14:1784:A:OP1	57:14:3549:HOH:O	2.09	0.68
27:16:3:C:H2'	27:16:4:C:C6	2.27	0.68
1:1G:448:A:P	1:1G:485:G:H22	2.15	0.68
13:4I:15:VAL:HG22	13:4I:45:VAL:HG12	1.76	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:19:69:ARG:NH2	28:19:128:GLY:O	2.26	0.68
1:1G:536:C:OP2	57:1G:1809:HOH:O	2.11	0.68
26:1H:2272:U:O4	57:1H:3954:HOH:O	2.11	0.68
34:58:13:TRP:HB2	34:58:133:GLN:HG2	1.75	0.68
34:58:33:LEU:HD12	34:58:38:HIS:CD2	2.28	0.68
26:1H:662:G:OP1	36:78:15:ARG:NE	2.24	0.68
26:14:61:G:OP1	49:G5:51:ARG:NH1	2.26	0.68
54:M5:59:LYS:C	54:M5:60:LEU:HG	2.12	0.68
1:1G:1095:U:P	1:1G:1108:G:H1	2.15	0.68
26:1H:2646:C:OP2	26:1H:2732:G:O2'	2.10	0.68
36:78:19:VAL:HB	36:78:27:HIS:HB2	1.76	0.68
52:J5:55:ARG:HH11	52:J5:55:ARG:HB2	1.58	0.68
1:13:1128:C:H5''	9:8E:16:ARG:HH22	1.58	0.68
2:1E:118:LEU:HD13	2:1E:142:LEU:HA	1.75	0.68
26:1H:918:A:N3	27:16:80:U:O2'	2.26	0.68
26:14:2472:G:H2'	26:14:2475:C:H41	1.56	0.68
26:1H:2121:G:N2	26:1H:2177:C:N3	2.41	0.68
32:51:101:ARG:NH2	32:51:121:ILE:O	2.26	0.68
26:14:252:G:OP2	36:35:50:ARG:NH2	2.24	0.68
1:1G:539:A:H2'	1:1G:540:G:C8	2.28	0.68
31:41:64:THR:HG23	31:41:94:LEU:HD13	1.76	0.68
7:62:113:GLU:HB2	7:62:119:ARG:HG2	1.75	0.68
28:11:142:VAL:HG23	28:11:193:VAL:HA	1.76	0.68
1:13:101:A:H2'	1:13:102:G:H8	1.59	0.68
26:14:1673:U:H5''	26:14:1674:G:OP2	1.93	0.68
26:14:2103:C:H2'	26:14:2104:G:C8	2.29	0.68
26:1H:1405:U:H2'	26:1H:1406:U:C6	2.29	0.68
22:1K:47:C:H42	22:1K:56:G:H1	1.41	0.68
29:29:59:VAL:HB	29:29:60:ASN:HA	1.75	0.68
34:58:65:LYS:HB3	34:58:69:GLN:HG3	1.74	0.68
54:Q8:30:ARG:NH1	54:Q8:30:ARG:HB2	2.09	0.68
1:13:618:C:H5''	1:13:619:U:H5''	1.76	0.68
1:13:735:C:H2'	1:13:736:C:H6	1.58	0.68
26:14:2588:G:OP2	57:14:3574:HOH:O	2.12	0.68
1:1G:1391:U:H2'	1:1G:1392:G:C8	2.29	0.68
26:1H:1253:A:C8	57:1H:4123:HOH:O	2.47	0.68
16:7I:22:THR:HA	16:7I:33:ILE:HG13	1.75	0.68
47:I8:38:VAL:HG12	47:I8:40:GLN:HG2	1.75	0.68
26:14:1266:G:O4'	43:A5:15:ARG:NH2	2.26	0.68
26:14:2270:G:OP2	57:14:3779:HOH:O	2.11	0.68
26:1H:1607:C:H4'	26:1H:1608:A:O5'	1.94	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:6I:39:LEU:HB3	15:6I:56:LEU:HD12	1.74	0.68
17:8A:18:THR:OG1	17:8A:69:LYS:NZ	2.20	0.68
45:G8:76:CYS:O	45:G8:78:ALA:N	2.25	0.68
26:1H:245:G:O6	54:Q8:8:LYS:NZ	2.27	0.68
1:13:328:C:H4'	1:13:329:A:H5''	1.74	0.68
26:14:1257:C:H4'	30:39:83:PHE:CD1	2.28	0.68
35:25:53:LYS:N	35:25:56:ASP:OD2	2.26	0.68
23:2L:72:C:H2'	23:2L:73:C:H6	1.59	0.68
26:1H:2292:C:OP1	39:A8:17:ARG:NH2	2.27	0.68
26:14:1174:A:H62	26:14:1177:A:H4'	1.59	0.67
26:14:2256:G:O6	57:14:3681:HOH:O	2.11	0.67
26:1H:730:C:OP2	57:1H:3618:HOH:O	2.12	0.67
27:1J:13:A:N1	27:1J:69:G:O2'	2.23	0.67
1:13:1190:G:H5''	3:2E:176:HIS:CE1	2.30	0.67
11:2I:98:LEU:O	11:2I:101:SER:OG	2.10	0.67
36:78:64:LYS:HE3	54:Q8:12:LYS:HD2	1.76	0.67
41:85:92:ARG:HH22	42:95:10:LYS:HA	1.60	0.67
51:I5:13:ARG:HG2	51:I5:22:ILE:HG13	1.76	0.67
26:14:1175:U:O2	26:14:1176:G:N2	2.27	0.67
26:14:1774:C:OP1	57:14:3555:HOH:O	2.10	0.67
26:14:528:A:C2	26:14:2042:A:H2'	2.29	0.67
23:2K:12:G:H1	23:2K:24:C:H42	1.42	0.67
23:2K:62:G:O2'	23:2K:63:G:OP2	2.12	0.67
26:14:2683:C:OP1	40:75:53:ARG:NH2	2.27	0.67
35:68:119:PRO:HB2	40:B8:68:TYR:CE2	2.29	0.67
50:H5:10:LYS:NZ	50:H5:15:TYR:OH	2.23	0.67
26:1H:1816:G:N7	28:11:35:LYS:NZ	2.42	0.67
1:13:1503:A:H5''	1:13:1531:A:H1'	1.76	0.67
26:14:1992:G:OP2	57:14:3586:HOH:O	2.12	0.67
26:14:2292:C:OP1	39:65:17:ARG:NH2	2.24	0.67
26:14:2415:G:H4'	36:35:67:MET:N	2.09	0.67
27:16:95:U:H2'	27:16:96:G:C8	2.29	0.67
37:45:25:ASP:O	46:D5:78:LYS:NZ	2.27	0.67
19:AI:5:LEU:HB3	19:AI:10:PHE:HE1	1.59	0.67
2:12:111:ARG:HH11	2:12:111:ARG:HA	1.58	0.67
26:14:1525:G:H2'	26:14:1526:G:C8	2.29	0.67
1:1G:547:A:OP2	4:32:2:GLY:N	2.26	0.67
23:2L:74:C:H2'	23:2L:75:G:C8	2.29	0.67
1:1G:376:G:H5''	16:7A:5:ARG:HD2	1.75	0.67
46:H8:117:LEU:HD22	46:H8:118:GLN:H	1.60	0.67
26:1H:2815:C:H5'	52:N8:29:THR:HG21	1.75	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:12:91:PRO:HG2	2:12:155:LEU:HG	1.75	0.67
1:13:974:A:OP2	14:5I:41:ARG:NH1	2.27	0.67
26:1H:2002:G:O6	57:1H:3940:HOH:O	2.12	0.67
40:75:108:ARG:HA	40:75:111:ARG:HG3	1.77	0.67
9:8E:13:ALA:HB2	9:8E:68:GLY:HA3	1.76	0.67
19:AA:66:MET:SD	19:AA:66:MET:N	2.67	0.67
1:1G:261:U:OP2	20:BA:79:ARG:NH2	2.28	0.67
46:D5:97:GLU:HB3	46:D5:125:LEU:HD11	1.76	0.67
46:D5:70:LEU:O	46:D5:89:PHE:N	2.23	0.67
50:H5:6:VAL:HG12	50:H5:56:VAL:HB	1.77	0.67
14:5A:45:ARG:O	14:5A:49:HIS:ND1	2.27	0.67
45:G8:97:ARG:CZ	45:G8:104:GLY:HA3	2.25	0.67
26:14:894:C:O2	26:14:896:A:N6	2.28	0.67
1:1G:929:G:H1	1:1G:1388:C:H42	1.42	0.67
26:1H:1397:U:OP2	26:1H:1398:C:N4	2.21	0.67
22:1K:30:A:N6	22:1K:42:U:O4	2.19	0.67
22:1K:49:C:N3	22:1K:55:G:N1	2.43	0.67
22:1L:15:A:H8	22:1L:16:U:C5	2.12	0.67
26:14:910:A:C5	37:45:13:GLN:HG3	2.30	0.67
26:14:2745:C:O2	32:59:139:GLN:NE2	2.28	0.67
26:14:634:C:H2'	26:14:635:C:C6	2.30	0.67
48:J8:48:LYS:HB3	48:J8:49:VAL:CA	2.24	0.67
13:4I:23:TYR:HD1	13:4I:67:GLU:HA	1.60	0.67
1:13:1063:C:H3'	1:13:1064:G:H2'	1.76	0.67
26:14:1250:G:OP2	36:35:21:ARG:NH1	2.28	0.67
26:14:2849:U:O4	40:75:23:ARG:NH2	2.28	0.67
26:14:529:A:H4'	26:14:530:G:H5'	1.77	0.67
27:16:95:U:H2'	27:16:96:G:H8	1.58	0.67
26:1H:581:C:H2'	26:1H:582:G:H8	1.58	0.67
31:41:35:GLU:HG3	31:41:36:LYS:HB2	1.76	0.67
44:B5:9:LEU:HA	49:G5:36:ARG:HH21	1.60	0.67
1:13:45:U:H2'	1:13:46:G:C8	2.30	0.66
26:14:1263:U:OP2	57:14:3820:HOH:O	2.12	0.66
26:14:2528:U:O2'	26:14:2530:A:OP1	2.12	0.66
26:1H:1496:A:H8	26:1H:1577:C:O2'	1.78	0.66
23:2L:83:U:O4	23:2L:84:A:N6	2.27	0.66
31:49:41:GLN:NE2	31:49:154:GLY:O	2.27	0.66
32:51:124:GLU:HB3	32:51:132:ARG:HB3	1.76	0.66
8:72:113:SER:HB2	8:72:134:ILE:HD11	1.75	0.66
43:A5:86:LEU:HD12	43:A5:87:PRO:HD2	1.77	0.66
1:13:1273:G:H3'	1:13:1274:G:H8	1.60	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:654(B):C:H2'	26:14:654(C):G:C8	2.29	0.66
26:1H:1357:U:OP2	57:1H:3809:HOH:O	2.12	0.66
40:75:87:ASP:N	40:75:87:ASP:OD1	2.28	0.66
51:I5:56:VAL:HG22	51:I5:57:GLU:HG3	1.76	0.66
26:14:2447:G:H3'	57:14:3633:HOH:O	1.94	0.66
1:1G:977:A:H1'	1:1G:982:U:O4	1.95	0.66
26:1H:2405:G:OP1	36:78:77:ARG:NH2	2.29	0.66
26:1H:945:A:OP1	57:1H:3872:HOH:O	2.12	0.66
10:1I:84:GLN:HE21	10:1I:88:LEU:HD22	1.61	0.66
37:45:26:TYR:O	37:45:28:ALA:N	2.28	0.66
48:J8:53:VAL:HB	48:J8:58:ILE:HD13	1.76	0.66
10:1A:3:LYS:NZ	10:1A:75:ILE:O	2.28	0.66
10:1A:91:PRO:HB2	10:1A:93:GLY:H	1.61	0.66
3:22:32:LEU:HD22	3:22:59:ARG:HH22	1.60	0.66
4:32:119:GLN:HG2	4:32:123:HIS:CD2	2.31	0.66
35:68:112:MET:HA	35:68:115:VAL:HG13	1.78	0.66
1:13:235:C:H5'	17:8I:70:ARG:HG2	1.78	0.66
45:C5:29:GLU:OE1	45:C5:29:GLU:N	2.28	0.66
46:D5:30:ASN:HD22	46:D5:90:VAL:HB	1.61	0.66
46:H8:140:ASP:OD2	46:H8:156:LYS:NZ	2.25	0.66
26:14:2497:A:O3'	57:14:3523:HOH:O	2.12	0.66
26:14:2591:C:OP1	28:19:239:ARG:HG3	1.95	0.66
26:14:71:A:H2	44:B5:31:HIS:HE2	1.41	0.66
2:1E:17:PHE:H	2:1E:17:PHE:HD1	1.44	0.66
29:21:97:LYS:N	29:21:100:GLU:OE1	2.24	0.66
29:21:24:THR:HG21	29:21:188:VAL:HG22	1.78	0.66
35:68:104:ARG:NH2	40:B8:43:GLN:OE1	2.27	0.66
28:11:38:LYS:HE2	28:11:39:LYS:O	1.96	0.66
26:14:1899:G:N2	26:14:1902:C:N4	2.44	0.66
26:14:2810:A:N6	26:14:2891:G:O2'	2.29	0.66
34:15:18:ALA:HA	34:15:21:LYS:HG3	1.78	0.66
1:1G:973:G:O3'	14:5A:41:ARG:NH2	2.27	0.66
26:1H:1503:U:H2'	26:1H:1504:C:H6	1.60	0.66
26:1H:2212:A:H1'	26:1H:2215:G:C5	2.30	0.66
26:1H:617:G:OP1	30:31:40:GLN:NE2	2.29	0.66
22:1K:47:C:N3	22:1K:56:G:N2	2.43	0.66
29:29:37:ARG:NE	29:29:42:ASP:OD2	2.26	0.66
12:3A:27:LEU:HG	12:3A:33:ARG:HG2	1.78	0.66
40:B8:50:ILE:HD13	40:B8:64:ARG:HB3	1.77	0.66
47:E5:18:ALA:HB3	47:E5:20:ARG:HH21	1.61	0.66
26:1H:489:G:N7	43:E8:49:LYS:NZ	2.42	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:259:G:H21	26:14:621:A:H8	1.40	0.66
26:1H:2134:A:OP2	26:1H:2157:G:N2	2.29	0.66
4:3E:22:LYS:HB2	4:3E:26:CYS:SG	2.36	0.66
12:3I:90:VAL:HG11	12:3I:93:LEU:HG	1.77	0.66
32:59:159:GLU:O	32:59:163:TYR:OH	2.13	0.66
1:1G:587:G:N2	1:1G:754:C:OP2	2.29	0.66
26:1H:2392:A:H2	26:1H:2424:C:H42	1.42	0.66
26:1H:529:A:H4'	26:1H:530:G:H5'	1.78	0.66
36:35:147:LEU:HG	36:35:148:LEU:H	1.60	0.66
12:3A:41:ARG:HD2	12:3A:42:THR:H	1.60	0.66
31:49:61:ALA:HB2	31:49:68:PRO:HD3	1.78	0.66
50:H5:59:VAL:HG12	50:H5:60:GLU:H	1.61	0.66
46:H8:28:MET:HG3	46:H8:37:VAL:HG11	1.78	0.66
1:13:601:C:H2'	1:13:602:A:C8	2.30	0.66
26:14:1427:A:H4'	26:14:1428:C:O4'	1.96	0.66
1:1G:359:U:H2'	1:1G:360:A:H8	1.59	0.66
43:A5:65:LEU:HD13	43:A5:68:ARG:HD3	1.78	0.66
26:14:784:A:OP1	57:14:3568:HOH:O	2.13	0.66
1:1G:1111:A:H2'	1:1G:1112:C:C6	2.31	0.66
26:1H:760:G:OP1	57:1H:3688:HOH:O	2.13	0.66
13:4I:20:THR:HG23	13:4I:26:GLY:HA3	1.78	0.66
46:H8:11:GLU:O	46:H8:36:LYS:NZ	2.28	0.66
51:M8:40:HIS:CD2	51:M8:45:GLY:HA3	2.31	0.66
54:Q8:53:PRO:HB3	54:Q8:56:GLU:H	1.61	0.66
1:13:1062:U:H2'	1:13:1063:C:C6	2.31	0.65
1:13:1305:G:H22	1:13:1331:G:C2'	2.05	0.65
26:14:1021:A:H62	26:14:1141:U:H3	1.44	0.65
1:1G:1316:G:H4'	14:5A:18:VAL:HG11	1.78	0.65
33:61:120:ILE:HD11	33:61:126:TYR:CE2	2.31	0.65
33:61:3:VAL:HG12	33:61:38:LEU:HA	1.78	0.65
33:69:75:LEU:HD22	33:69:76:THR:H	1.61	0.65
7:6E:62:PHE:HD1	7:6E:124:LEU:HD11	1.60	0.65
26:14:459:U:H5''	53:L5:40:TRP:CD2	2.32	0.65
1:13:1149:C:H2'	1:13:1150:U:C6	2.30	0.65
1:13:1182:G:H4'	1:13:1183:A:H5''	1.77	0.65
10:1A:48:THR:HG1	10:1A:62:HIS:CE1	2.13	0.65
1:1G:973:G:H1'	10:1A:55:LYS:NZ	2.12	0.65
26:1H:1341:U:O4	44:F8:16:LYS:NZ	2.30	0.65
26:1H:330:A:O2'	26:1H:331:A:H8	1.79	0.65
26:1H:33:U:H4'	26:1H:34:C:OP1	1.96	0.65
10:1I:46:ARG:HB2	10:1I:46:ARG:HH11	1.61	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2377:A:H4'	39:65:111:GLU:HG2	1.78	0.65
35:25:119:PRO:HB2	40:75:68:TYR:CE2	2.30	0.65
42:D8:34:GLU:HB2	42:D8:58:VAL:HG22	1.79	0.65
1:13:1305:G:N2	1:13:1331:G:H2'	2.05	0.65
26:14:273(C):C:H42	26:14:363(C):G:H1	1.44	0.65
1:1G:1326:C:OP1	21:1B:12:LYS:NZ	2.28	0.65
1:1G:1469:G:O6	57:1G:1831:HOH:O	2.11	0.65
26:1H:2355:C:O2	47:I8:39:ARG:NH2	2.30	0.65
3:22:7:PRO:O	3:22:11:ARG:NH1	2.28	0.65
24:3L:37:G:H22	25:4L:44:U:H3	1.43	0.65
40:B8:27:THR:HG23	40:B8:90:GLN:HB3	1.78	0.65
2:12:115:LEU:HD13	2:12:145:LEU:HB3	1.78	0.65
1:13:439:A:H2'	1:13:440:A:O4'	1.97	0.65
1:13:963:G:N2	1:13:972:C:N3	2.32	0.65
26:1H:1826:G:H4'	28:11:242:ARG:HH21	1.62	0.65
22:1L:31:C:O2	22:1L:41:G:N2	2.28	0.65
37:45:117:ALA:HA	37:45:120:ILE:HB	1.76	0.65
1:13:1149:C:H2'	1:13:1150:U:H6	1.61	0.65
26:14:1069:A:H2	26:14:1094:U:H3	1.43	0.65
26:14:1252:G:O4'	41:85:33:ARG:HD3	1.95	0.65
26:14:544:C:H2'	26:14:545:G:H5'	1.77	0.65
26:14:571:A:OP2	57:14:3987:HOH:O	2.13	0.65
21:1B:9:ARG:HG3	21:1B:10:ARG:N	2.12	0.65
26:1H:1141:U:H6	34:58:63:THR:HG1	1.42	0.65
23:2L:73:C:H2'	23:2L:74:C:C6	2.32	0.65
27:1J:42:C:O2'	31:49:67:LYS:O	2.12	0.65
32:59:26:VAL:HG21	32:59:75:ALA:HB1	1.77	0.65
36:78:138:LEU:HD12	36:78:144:GLU:HG3	1.78	0.65
8:7E:106:GLY:HA2	8:7E:122:ARG:HH12	1.60	0.65
46:D5:54:HIS:HB3	46:D5:101:PRO:HD3	1.77	0.65
46:H8:108:PRO:HB2	46:H8:112:ARG:HA	1.78	0.65
22:1L:48:C:O2'	22:1L:49:C:OP1	2.11	0.65
29:21:39:PRO:HD3	29:21:45:THR:HG23	1.78	0.65
29:29:60:ASN:HB3	29:29:61:ARG:C	2.17	0.65
11:2A:34:ASP:HB2	11:2A:35:PRO:HD2	1.77	0.65
33:69:130:TYR:HB3	33:69:136:VAL:HG13	1.78	0.65
40:75:16:ARG:HH12	40:75:83:ILE:HB	1.61	0.65
1:1G:1014:A:H4'	19:AA:14:HIS:CE1	2.31	0.65
45:G8:76:CYS:HB2	45:G8:82:PRO:HD3	1.79	0.65
46:H8:111:VAL:HG21	46:H8:146:ILE:HG13	1.78	0.65
26:1H:784:A:C5	28:11:229:VAL:HG21	2.32	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:12:178:ARG:NH1	2:12:196:LEU:O	2.27	0.65
26:14:2287:A:H62	26:14:2344:U:H3	1.43	0.65
26:1H:2577:A:H5''	26:1H:2578:G:H5'	1.78	0.65
26:1H:299:A:H5'	26:1H:300:A:OP2	1.97	0.65
26:1H:761:A:OP1	57:1H:3622:HOH:O	2.14	0.65
26:14:2444:G:OP2	30:39:68:LYS:HE2	1.97	0.65
5:4E:71:LEU:HD22	5:4E:114:GLY:HA3	1.78	0.65
22:1K:37:G:H1	25:4K:49:U:H3	1.43	0.65
18:9I:38:GLU:OE1	18:9I:41:LYS:NZ	2.29	0.65
40:B8:64:ARG:HB2	40:B8:73:GLU:HG2	1.78	0.65
46:D5:163:LEU:HD23	46:D5:163:LEU:H	1.60	0.65
1:13:21:G:OP1	57:13:1810:HOH:O	2.13	0.65
26:14:1024:G:H3'	26:14:1025:G:H5''	1.79	0.65
26:14:1939:U:OP1	26:14:2604:U:O2'	2.11	0.65
1:1G:1224:G:N1	1:1G:1322:C:H1'	2.10	0.65
1:1G:32:A:H2'	1:1G:33:A:C8	2.31	0.65
1:1G:380:G:N2	1:1G:383:A:OP2	2.30	0.65
26:1H:2168:G:OP1	26:1H:2168:G:H4'	1.97	0.65
23:2L:20:U:O2'	23:2L:21:A:OP2	2.10	0.65
26:1H:2306:C:N4	31:41:42:GLY:O	2.30	0.65
26:14:309:G:H4'	45:C5:18:GLY:HA3	1.79	0.65
26:14:2058:A:OP1	57:14:3855:HOH:O	2.15	0.65
26:14:2746:U:OP1	32:59:85:LYS:NZ	2.22	0.65
1:1G:1067:A:HO2'	1:1G:1093:A:HO2'	1.42	0.65
1:1G:1412:C:H2'	1:1G:1413:A:C8	2.32	0.65
26:1H:1388:G:H2'	26:1H:1389:G:C8	2.30	0.65
27:1J:28:C:H42	27:1J:56:G:H1	1.44	0.65
3:22:73:PRO:O	3:22:77:ILE:N	2.22	0.65
23:2L:72:C:H2'	23:2L:73:C:C6	2.32	0.65
33:61:120:ILE:HD11	33:61:126:TYR:CZ	2.31	0.65
26:14:2010:G:H5''	43:A5:42:ARG:HB2	1.78	0.65
26:14:491:G:H2'	26:14:492:A:H8	1.62	0.65
28:19:24:ILE:HA	28:19:82:ILE:HG22	1.79	0.65
1:1G:1503:A:OP1	1:1G:1531:A:O2'	2.15	0.65
4:3E:83:SER:HA	4:3E:89:THR:HG23	1.79	0.65
42:95:58:VAL:HB	42:95:98:GLU:HB2	1.79	0.65
26:1H:71:A:H2	44:F8:31:HIS:CE1	2.15	0.65
51:I5:57:GLU:O	51:I5:61:ARG:NH2	2.28	0.65
53:P8:8:ASN:OD1	53:P8:11:LYS:N	2.24	0.65
26:14:2112:G:O2'	26:14:2114:A:N6	2.27	0.64
26:14:2557:G:H2'	26:14:2558:C:H6	1.60	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:973:G:O4'	10:1A:55:LYS:HG3	1.96	0.64
26:1H:1520:U:H2'	26:1H:1521:G:O4'	1.97	0.64
26:1H:2576:G:OP1	57:1H:3667:HOH:O	2.14	0.64
37:88:104:PHE:HE2	37:88:125:LEU:HD11	1.62	0.64
46:H8:5:LEU:HD23	46:H8:47:VAL:HG21	1.79	0.64
26:14:1359:A:H62	26:14:1372:U:H3	1.45	0.64
26:1H:1249:U:OP1	57:1H:3734:HOH:O	2.14	0.64
26:1H:1997:G:OP2	57:1H:3785:HOH:O	2.15	0.64
26:1H:2255:G:OP2	57:1H:3878:HOH:O	2.14	0.64
13:4A:29:ARG:HB3	13:4A:64:TRP:CZ2	2.33	0.64
26:1H:1006:C:H1'	34:58:106:MET:HE3	1.78	0.64
1:13:468:A:H5''	16:7I:80:PHE:HB3	1.77	0.64
43:E8:57:ASN:O	43:E8:61:ASN:HB2	1.97	0.64
48:J8:87:PRO:HA	48:J8:90:ILE:HG12	1.79	0.64
2:1E:134:GLU:HA	2:1E:137:ARG:HB3	1.78	0.64
1:1G:280:C:H3'	1:1G:281:G:H5'	1.78	0.64
1:1G:636:U:H2'	1:1G:637:G:H8	1.62	0.64
26:1H:1103:A:H3'	26:1H:1104:C:H6	1.62	0.64
27:1J:15:A:H5'	27:1J:16:G:C8	2.32	0.64
40:75:125:ARG:HG3	40:75:129:ARG:HH21	1.62	0.64
9:82:28:VAL:HG22	9:82:63:ILE:HB	1.80	0.64
1:1G:1119:C:OP1	9:82:83:ARG:NH1	2.30	0.64
19:AI:40:ILE:HG23	19:AI:41:VAL:HG13	1.79	0.64
1:13:1279:A:O2'	1:13:1281:U:OP2	2.15	0.64
26:14:1485:G:H1	26:14:1504:C:H42	1.44	0.64
26:14:1828:G:OP2	57:14:3532:HOH:O	2.15	0.64
26:1H:1826:G:O2'	28:11:242:ARG:NH2	2.31	0.64
26:1H:2408:U:H2'	26:1H:2409:G:H8	1.63	0.64
48:F5:60:PHE:HE2	48:F5:91:LYS:HD3	1.63	0.64
1:13:14:U:OP2	57:13:1815:HOH:O	2.15	0.64
1:13:946:A:H2'	1:13:947:G:C8	2.33	0.64
26:14:1973:G:H2'	26:14:1974:C:C6	2.31	0.64
34:15:38:HIS:NE2	34:15:50:ASP:OD2	2.30	0.64
1:1G:1095:U:OP1	1:1G:1108:G:N2	2.30	0.64
1:1G:1356:G:H2'	1:1G:1357:A:C8	2.32	0.64
26:1H:41:C:H42	26:1H:438:G:H1	1.45	0.64
14:5A:37:PHE:HZ	14:5A:56:VAL:HG21	1.63	0.64
16:7A:22:THR:HA	16:7A:33:ILE:HG13	1.78	0.64
1:1G:1342:C:H4'	9:82:125:TYR:HB2	1.79	0.64
20:BI:71:THR:HG22	20:BI:72:LEU:H	1.61	0.64
52:N8:40:LYS:HE2	52:N8:47:PRO:HD2	1.80	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1132:C:H2'	1:13:1133:G:H8	1.60	0.64
1:13:1240:U:OP2	7:6E:116:ALA:N	2.30	0.64
1:13:148:G:H2'	1:13:149:A:H8	1.61	0.64
26:14:1794:U:H2'	26:14:1795:C:H6	1.61	0.64
26:14:2715:C:H2'	26:14:2716:U:H6	1.61	0.64
28:19:132:PRO:HG3	28:19:190:TYR:CE1	2.33	0.64
1:1G:409:G:H1	1:1G:433:C:H42	1.45	0.64
26:1H:2679:A:H4'	29:21:165:VAL:HG11	1.80	0.64
3:2E:152:ILE:HG12	3:2E:167:TRP:HB2	1.80	0.64
30:39:20:LEU:HG	30:39:199:TRP:HH2	1.63	0.64
32:51:27:LYS:HA	32:51:32:GLU:HA	1.79	0.64
6:5E:39:LYS:H	6:5E:64:GLN:HB2	1.63	0.64
36:78:36:LYS:O	36:78:40:SER:HB3	1.98	0.64
37:88:87:LYS:HG2	37:88:88:GLY:H	1.63	0.64
40:B8:5:ALA:HA	40:B8:8:LYS:HG2	1.80	0.64
47:E5:26:TYR:O	47:E5:29:GLN:HB2	1.98	0.64
43:A5:35:ILE:HG23	52:J5:28:PRO:HD2	1.80	0.64
1:13:991:U:H4'	1:13:992:U:H5''	1.78	0.64
26:14:1729:A:H2'	26:14:1731:G:H22	1.63	0.64
1:1G:1127:G:H2'	1:1G:1147:C:H42	1.62	0.64
13:4A:96:LEU:O	13:4A:110:ARG:NE	2.28	0.64
13:4A:86:CYS:HB2	19:AA:73:GLU:HB3	1.80	0.64
32:51:4:ILE:HG13	32:51:6:ARG:HB2	1.79	0.64
34:58:9:VAL:HG11	34:58:39:ARG:HH12	1.62	0.64
48:J8:18:ILE:HG12	48:J8:37:ILE:HG12	1.80	0.64
1:1G:1012:U:H3	1:1G:1017:G:H1	1.46	0.64
26:1H:29:U:H2'	26:1H:30:G:C8	2.33	0.64
54:Q8:49:VAL:HG13	54:Q8:50:LEU:O	1.96	0.64
26:14:2116:G:N1	26:14:2162:G:OP1	2.31	0.64
1:1G:413:G:O2'	1:1G:414:A:OP2	2.11	0.64
1:1G:426:G:OP1	4:32:38:TYR:OH	2.12	0.64
26:1H:2444:G:OP2	30:31:68:LYS:HE2	1.98	0.64
26:1H:547:A:H2'	26:1H:548:A:C8	2.33	0.64
26:1H:974(A):C:OP1	57:1H:3911:HOH:O	2.15	0.64
23:2L:36:A:H2'	23:2L:37:G:C8	2.33	0.64
30:39:157:VAL:HB	30:39:194:MET:HB3	1.79	0.64
44:F8:26:TYR:HD1	44:F8:92:LEU:HD12	1.63	0.64
2:12:179:LYS:HA	8:72:72:PRO:HG3	1.80	0.64
26:1H:907:U:O2'	37:88:101:ARG:NH2	2.29	0.64
29:29:103:ASP:OD1	29:29:201:THR:HG23	1.98	0.64
11:2I:85:ARG:HE	11:2I:111:ASP:HB3	1.63	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:41:64:THR:HG21	31:41:92:VAL:HG21	1.79	0.64
32:51:33:LEU:HD21	32:51:136:ILE:HB	1.80	0.64
26:14:2749:A:N1	26:14:2750:A:N6	2.46	0.63
26:14:751:A:OP1	57:14:3505:HOH:O	2.15	0.63
1:1G:413:G:HO2'	1:1G:428:G:H22	1.47	0.63
26:1H:1728:G:H8	26:1H:1732:A:H62	1.45	0.63
26:1H:1830:C:H42	26:1H:1975:G:H1	1.44	0.63
26:1H:1849:G:N2	26:1H:1893:C:O2	2.26	0.63
26:1H:2147:G:H2'	26:1H:2148:G:H4'	1.80	0.63
26:1H:635:C:O2'	26:1H:639:U:OP1	2.15	0.63
22:1L:64:G:H1	22:1L:72:C:H42	1.45	0.63
37:45:114:ALA:O	37:45:118:LEU:HB2	1.98	0.63
26:14:1111:A:H4'	32:59:3:ARG:HD3	1.80	0.63
33:61:131:LYS:HB3	33:61:132:PRO:HA	1.79	0.63
7:62:21:VAL:HG23	7:62:22:LEU:HD12	1.80	0.63
48:J8:92:LYS:O	48:J8:94:LEU:N	2.31	0.63
2:12:7:VAL:HG22	2:12:8:LYS:H	1.62	0.63
26:14:492:A:H2'	26:14:493:G:O4'	1.97	0.63
1:1G:142:G:H2'	1:1G:143:A:H8	1.61	0.63
26:1H:2419:U:O4	54:Q8:29:LYS:NZ	2.18	0.63
26:1H:2729:G:H2'	26:1H:2730:C:C6	2.34	0.63
29:29:12:THR:O	29:29:23:VAL:HG22	1.98	0.63
30:31:85:GLY:O	57:31:501:HOH:O	2.15	0.63
5:42:51:VAL:HG23	5:42:52:PRO:HD3	1.79	0.63
31:49:4:ASP:CG	31:49:5:VAL:H	2.01	0.63
27:1J:41:U:C4	31:49:70:VAL:HG23	2.32	0.63
1:13:237:C:H5''	17:8I:25:ARG:CZ	2.28	0.63
10:1A:32:ALA:HA	10:1A:76:ASN:HB2	1.80	0.63
26:1H:1187:G:OP2	57:1H:3727:HOH:O	2.15	0.63
26:1H:2502:G:H5''	26:1H:2503:A:H5''	1.81	0.63
30:31:130:ALA:H	30:31:132:VAL:HG13	1.63	0.63
12:3A:27:LEU:HB3	12:3A:33:ARG:HD3	1.79	0.63
13:4A:34:LEU:O	13:4A:38:GLY:N	2.32	0.63
33:69:3:VAL:HG12	33:69:38:LEU:HA	1.78	0.63
9:8E:112:LYS:HA	9:8E:119:ALA:HB2	1.80	0.63
48:J8:8:SER:HB3	48:J8:66:HIS:CD2	2.33	0.63
1:13:222:U:H2'	1:13:223:U:C6	2.34	0.63
26:1H:1434:A:H61	26:1H:1558:A:H62	1.46	0.63
29:21:12:THR:OG1	29:21:13:ARG:N	2.31	0.63
30:31:129:PHE:HA	30:31:142:TRP:NE1	2.13	0.63
36:35:147:LEU:CG	36:35:148:LEU:H	2.10	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:41:40:ASN:HB2	31:41:91:ARG:HG2	1.79	0.63
39:65:85:VAL:H	39:65:110:LEU:HB3	1.64	0.63
42:D8:3:ALA:HB1	42:D8:38:LEU:HD11	1.80	0.63
54:Q8:36:LYS:HG2	54:Q8:39:LYS:NZ	2.14	0.63
1:13:1028(A):C:H2'	1:13:1028(B):C:H5	1.64	0.63
1:1G:709:G:H2'	1:1G:710:G:H8	1.63	0.63
26:1H:185:U:H2'	26:1H:186:G:H8	1.62	0.63
26:1H:270(I):G:H1	26:1H:270(Q):C:H42	1.44	0.63
26:1H:2788:C:O2'	26:1H:2809:A:N3	2.30	0.63
29:29:33:VAL:HG12	29:29:89:ASP:CB	2.27	0.63
11:2A:85:ARG:HD3	11:2A:113:PRO:HD3	1.78	0.63
1:13:224:C:H2'	1:13:225:C:C6	2.33	0.63
26:14:1252:G:N3	41:85:33:ARG:HD2	2.13	0.63
28:19:182:LEU:H	28:19:272:ALA:HB2	1.63	0.63
26:1H:2392:A:OP2	54:Q8:30:ARG:NH2	2.31	0.63
33:61:69:LYS:HA	33:61:136:VAL:HB	1.80	0.63
1:1G:1291:G:OP1	7:62:41:ARG:NH2	2.32	0.63
9:82:4:TYR:HB2	9:82:19:LEU:HB2	1.78	0.63
19:AI:40:ILE:HD11	19:AI:62:ILE:HG23	1.79	0.63
46:H8:111:VAL:HG11	46:H8:146:ILE:HB	1.80	0.63
1:13:1342:C:H2'	1:13:1343:G:C8	2.34	0.63
26:14:1260:G:H2'	26:14:1261:C:C6	2.33	0.63
26:14:265:A:N6	26:14:427:U:O2'	2.31	0.63
26:1H:1165:U:H2'	26:1H:1166:C:C6	2.34	0.63
26:1H:1510:A:O2'	26:1H:1512:G:N7	2.28	0.63
26:1H:1899:G:N2	26:1H:1902:C:H5	1.96	0.63
26:1H:581:C:H2'	26:1H:582:G:C8	2.34	0.63
4:32:108:LEU:HD21	4:32:183:GLY:HA3	1.79	0.63
1:1G:1533:C:H42	25:4L:39:U:H3	1.45	0.63
18:9A:22:VAL:HG22	18:9A:23:LYS:H	1.64	0.63
46:D5:72:ARG:NH2	46:D5:97:GLU:O	2.30	0.63
36:35:64:LYS:HB3	54:M5:30:ARG:HH22	1.62	0.63
2:12:58:ILE:O	2:12:62:ALA:N	2.32	0.63
26:14:2148:G:H2'	26:14:2149:G:H8	1.64	0.63
1:1G:1142:G:H3'	1:1G:1143:G:C8	2.33	0.63
1:1G:1492:A:OP1	12:3A:46:LYS:HB2	1.99	0.63
26:1H:2176:A:H2'	26:1H:2177:C:H6	1.62	0.63
26:1H:2406:U:OP1	57:1H:4136:HOH:O	2.16	0.63
26:1H:2836:U:H2'	26:1H:2837:G:C8	2.33	0.63
30:39:24:LEU:HD12	30:39:25:PRO:HD3	1.81	0.63
32:51:86:GLU:CD	32:51:86:GLU:H	2.01	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:55:77:ARG:O	38:55:80:PHE:N	2.31	0.63
15:6A:87:ILE:HG22	15:6A:88:ARG:H	1.64	0.63
26:1H:1228:G:OP2	41:C8:16:LYS:NZ	2.30	0.63
54:Q8:35:GLN:C	54:Q8:37:SER:H	2.00	0.63
26:14:2302:G:N2	26:14:2314:C:O2	2.31	0.63
26:14:2306:C:H3'	26:14:2307:G:H5''	1.81	0.63
2:1E:8:LYS:HG2	2:1E:9:GLU:HG2	1.80	0.63
1:1G:1151:A:O2'	1:1G:1152:A:O5'	2.17	0.63
32:59:69:ARG:HA	32:59:72:ILE:HD12	1.80	0.63
1:1G:1379:G:OP1	7:62:6:ARG:NH1	2.31	0.63
9:8E:53:VAL:HG12	9:8E:55:ALA:H	1.64	0.63
38:98:104:ARG:HG3	38:98:111:LEU:HD21	1.81	0.63
19:AI:5:LEU:HD13	19:AI:10:PHE:HD1	1.64	0.63
45:C5:50:ARG:HB3	45:C5:53:PRO:HG3	1.81	0.63
53:P8:12:ARG:HH21	53:P8:44:PRO:HB3	1.64	0.63
1:13:651:C:H2'	1:13:652:U:H6	1.64	0.62
1:13:67:C:H2'	1:13:68:G:H8	1.64	0.62
26:14:443:A:H1'	26:14:1201:C:O4'	1.99	0.62
27:16:111:U:H2'	27:16:112:G:H8	1.64	0.62
21:1F:12:LYS:HB3	21:1F:22:ARG:HD2	1.81	0.62
3:2E:59:ARG:HH12	3:2E:97:LYS:HE3	1.64	0.62
30:31:40:GLN:NE2	30:31:182:ASN:HB2	2.13	0.62
13:4A:10:PRO:HG2	13:4A:45:VAL:HG11	1.81	0.62
32:51:4:ILE:HB	32:51:6:ARG:HG3	1.80	0.62
7:6E:111:ARG:NH1	7:6E:113:GLU:OE2	2.32	0.62
41:85:74:LEU:HD13	41:85:79:PHE:HB2	1.81	0.62
28:11:72:LYS:HD2	28:11:75:ILE:HD12	1.81	0.62
1:13:1106:G:H5''	3:2E:172:ARG:HG2	1.81	0.62
26:14:1204:A:O2'	26:14:1205:U:OP2	2.16	0.62
26:14:1405:U:H2'	26:14:1406:U:C6	2.34	0.62
26:14:602:G:O2'	26:14:655:A:N6	2.32	0.62
1:1G:860:A:H2'	1:1G:861:G:O4'	1.99	0.62
26:1H:1290:C:H2'	26:1H:1291:C:C6	2.34	0.62
26:1H:2079:U:O4	57:1H:3998:HOH:O	2.16	0.62
26:1H:2420:C:H41	54:Q8:29:LYS:HA	1.63	0.62
10:1I:50:ILE:HA	10:1I:60:ARG:HB3	1.81	0.62
33:69:101:LEU:HB3	33:69:105:HIS:HB2	1.80	0.62
16:7A:34:GLU:OE1	16:7A:55:ARG:NH1	2.31	0.62
1:13:413:G:N7	4:3E:35:ARG:NH1	2.47	0.62
26:14:1257:C:H4'	30:39:83:PHE:CE1	2.34	0.62
26:14:1754:C:P	40:75:96:ARG:HH12	2.21	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:287:C:H2'	26:14:288:C:H6	1.63	0.62
26:14:95:G:H4'	49:G5:46:GLN:HB2	1.81	0.62
5:4E:100:VAL:HG12	5:4E:118:ILE:HG22	1.81	0.62
44:B5:60:ARG:HA	44:B5:60:ARG:HH11	1.64	0.62
26:14:2271:G:H5''	47:E5:20:ARG:NE	2.13	0.62
48:J8:3:LYS:O	48:J8:12:PRO:HD3	1.98	0.62
1:13:1178:G:N2	1:13:1181:G:H8	1.96	0.62
26:14:2816:C:O3'	38:55:99:LYS:NZ	2.32	0.62
2:1E:88:ALA:HB2	2:1E:219:VAL:HG13	1.80	0.62
1:1G:1028:C:H42	1:1G:1033:G:H1	1.47	0.62
1:1G:673:G:H2'	1:1G:674:G:C8	2.34	0.62
26:1H:1045:A:H1'	26:1H:1047:G:N3	2.14	0.62
26:1H:77:C:H5''	49:K8:10:LEU:HD11	1.81	0.62
10:1I:49:VAL:O	10:1I:60:ARG:HB2	1.99	0.62
22:1L:41:G:H2'	22:1L:42:U:C6	2.33	0.62
26:14:2811:G:OP1	29:29:61:ARG:HB3	1.99	0.62
26:14:2690:C:OP2	38:55:14:SER:HB3	1.98	0.62
8:72:86:ILE:HG12	8:72:135:CYS:HA	1.80	0.62
16:7I:43:LYS:HA	16:7I:48:TRP:HB2	1.81	0.62
41:85:92:ARG:O	41:85:94:ASN:N	2.33	0.62
18:9I:52:PRO:HB2	18:9I:54:ARG:HD3	1.80	0.62
41:C8:8:VAL:HG23	41:C8:11:ARG:HH21	1.64	0.62
46:D5:4:ARG:HB3	46:D5:58:VAL:HB	1.81	0.62
46:D5:28:MET:HE1	46:D5:61:LEU:HD21	1.81	0.62
42:D8:19:LYS:HG3	42:D8:95:LEU:HD23	1.80	0.62
44:F8:1:MET:C	44:F8:3:THR:H	2.02	0.62
26:1H:95:G:H4'	49:K8:46:GLN:HB3	1.81	0.62
26:14:1464:C:HO2'	26:14:1528:A:H8	1.47	0.62
1:1G:978:A:O2'	1:1G:1322:C:N3	2.32	0.62
27:1J:15:A:H1'	27:1J:109:G:N9	2.14	0.62
35:25:2:ILE:HD12	35:25:6:THR:HG21	1.80	0.62
5:42:102:ALA:HB1	5:42:106:PRO:HG2	1.81	0.62
7:6E:65:ALA:HB2	7:6E:128:ALA:HB2	1.81	0.62
36:78:79:ARG:HB3	36:78:110:TYR:CD1	2.34	0.62
45:G8:39:VAL:HB	45:G8:42:VAL:HG11	1.80	0.62
48:J8:7:ILE:HD13	48:J8:62:VAL:HG11	1.81	0.62
54:Q8:32:LEU:HG	54:Q8:33:ASN:HD22	1.64	0.62
1:13:581:G:N2	1:13:760:G:N7	2.47	0.62
26:14:1106:G:H2'	26:14:1107:G:C8	2.34	0.62
29:21:77:ILE:O	29:21:79:ARG:N	2.33	0.62
3:22:11:ARG:HE	3:22:180:ALA:HB3	1.65	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:29:58:ARG:HH21	29:29:58:ARG:HG2	1.65	0.62
29:29:81:ILE:HG22	29:29:82:ARG:H	1.64	0.62
30:39:32:LEU:HD11	30:39:105:VAL:HG13	1.80	0.62
46:D5:4:ARG:NH1	46:D5:60:GLU:OE2	2.32	0.62
26:14:234:C:H2'	26:14:235:U:H6	1.63	0.62
26:14:848:G:H2'	26:14:849:A:C8	2.35	0.62
2:1E:115:LEU:HD13	2:1E:145:LEU:HB3	1.81	0.62
1:1G:407:G:O6	1:1G:435:C:N4	2.20	0.62
1:1G:892:A:O2'	1:1G:1415:G:H4'	2.00	0.62
26:1H:1430:C:H2'	26:1H:1431:U:C6	2.34	0.62
22:1L:21:A:O2'	22:1L:59:U:O4	2.16	0.62
30:39:36:VAL:HB	30:39:183:VAL:HG11	1.82	0.62
17:8A:7:THR:O	17:8A:23:VAL:HG13	2.00	0.62
38:98:27:SER:HB3	38:98:34:ILE:HD11	1.80	0.62
20:BA:16:HIS:O	20:BA:19:SER:OG	2.16	0.62
26:14:185:U:H2'	26:14:186:G:C8	2.35	0.62
26:1H:1216:G:OP2	41:C8:12:ARG:NH2	2.32	0.62
26:1H:2855:C:H2'	26:1H:2856:C:H6	1.65	0.62
3:2E:119:ARG:O	3:2E:123:GLN:HB2	2.00	0.62
6:5E:42:GLU:OE1	6:5E:59:TYR:OH	2.13	0.62
46:H8:19:ARG:NH1	46:H8:84:GLU:HB2	2.14	0.62
51:I5:58:ARG:HA	51:I5:58:ARG:NE	2.14	0.62
49:K8:47:ASN:C	49:K8:49:LYS:H	2.02	0.62
1:13:223:U:H2'	1:13:224:C:H6	1.63	0.62
26:14:1079:C:N4	26:14:1088:A:OP1	2.30	0.62
26:14:2272:U:O4	57:14:3776:HOH:O	2.11	0.62
26:14:1138:G:H21	34:15:106:MET:HE3	1.65	0.62
10:1A:3:LYS:HD2	10:1A:77:PRO:HG3	1.82	0.62
26:1H:1093:G:N2	26:1H:1097:U:OP2	2.24	0.62
26:1H:2593:U:H2'	26:1H:2594:C:H6	1.65	0.62
26:1H:607:U:OP1	30:31:102:PRO:HA	1.99	0.62
22:1L:21:A:H1'	22:1L:22:G:O5'	1.98	0.62
23:2K:61:C:N4	23:2K:62:G:O6	2.33	0.62
24:3K:7:G:H1	24:3K:77:C:H42	1.48	0.62
32:51:113:VAL:HG11	32:51:151:ILE:HD13	1.79	0.62
9:82:9:ARG:HG2	9:82:14:VAL:HG22	1.80	0.62
37:88:110:THR:HG23	37:88:113:GLN:OE1	1.99	0.62
42:95:85:LYS:HD2	42:95:87:HIS:HA	1.82	0.62
46:D5:19:ARG:NH1	46:D5:84:GLU:HB2	2.15	0.62
48:F5:85:LEU:HA	48:F5:87:PRO:HD2	1.80	0.62
26:14:1165:U:H2'	26:14:1166:C:C6	2.35	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2074:U:OP1	57:14:3503:HOH:O	2.16	0.62
1:1G:1226:C:H2'	13:4A:103:THR:HB	1.80	0.62
26:1H:1009:A:OP2	34:58:37:LYS:NZ	2.33	0.62
26:1H:2331:G:H4'	47:18:43:THR:H	1.65	0.62
37:45:139:GLU:HG2	37:45:140:ALA:H	1.65	0.62
39:65:3:ARG:HE	39:65:4:LEU:N	1.98	0.62
36:78:126:VAL:HG12	36:78:147:LEU:HD21	1.81	0.62
17:8I:76:LEU:HD11	17:8I:79:SER:HB3	1.80	0.62
38:98:12:ARG:HG2	38:98:12:ARG:HH11	1.65	0.62
40:B8:85:LYS:NZ	40:B8:87:ASP:OD1	2.19	0.62
52:N8:50:GLY:H	52:N8:56:LYS:HB2	1.64	0.62
26:1H:1309:G:H4'	53:P8:7:PRO:HB2	1.81	0.62
26:14:2295:C:H5	39:65:13:ARG:HH22	1.47	0.61
26:14:2738:A:OP2	57:14:4010:HOH:O	2.16	0.61
9:82:114:TYR:HE2	10:1A:60:ARG:H	1.46	0.61
21:1F:10:ARG:HE	21:1F:13:ILE:HD12	1.64	0.61
1:1G:413:G:O2'	1:1G:428:G:N2	2.33	0.61
27:1J:44:G:H5''	27:1J:45:A:OP1	2.00	0.61
12:3I:28:LYS:HD2	12:3I:62:SER:HB3	1.81	0.61
24:3L:30:A:N1	24:3L:43:A:N6	2.47	0.61
31:49:66:GLN:OE1	31:49:98:ARG:NH1	2.32	0.61
34:58:96:GLU:C	34:58:98:VAL:H	2.02	0.61
36:78:50:ARG:CB	36:78:50:ARG:HH21	2.12	0.61
16:7I:57:ARG:NH2	16:7I:78:GLY:O	2.33	0.61
18:9A:37:VAL:HG11	18:9A:78:LEU:HB3	1.80	0.61
40:B8:56:GLY:O	40:B8:59:THR:HG22	2.00	0.61
49:G5:56:GLN:HG2	49:G5:59:ARG:HH21	1.65	0.61
2:12:91:PRO:HG3	2:12:154:LEU:HB2	1.81	0.61
26:14:1260:G:H2'	26:14:1261:C:H6	1.65	0.61
26:14:2777:G:H5''	26:14:2778:A:H5'	1.81	0.61
26:14:581:C:H2'	26:14:582:G:C8	2.35	0.61
1:1G:688:G:H2'	1:1G:689:C:H6	1.65	0.61
26:1H:528:A:N1	26:1H:2042:A:H2'	2.15	0.61
30:31:29:ASN:H	30:31:112:MET:HE3	1.64	0.61
4:32:108:LEU:HD12	4:32:170:VAL:HG11	1.81	0.61
4:32:150:GLU:C	4:32:152:SER:H	2.03	0.61
31:41:109:VAL:HG22	51:M8:33:VAL:HG21	1.82	0.61
14:5A:21:TYR:OH	14:5A:23:ARG:NH2	2.33	0.61
26:1H:1252:G:N3	41:C8:33:ARG:HD2	2.15	0.61
54:Q8:8:LYS:HD2	54:Q8:8:LYS:H	1.65	0.61
2:12:174:VAL:HA	2:12:177:ALA:HB3	1.83	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:130:A:OP2	17:8I:63:ARG:NH2	2.33	0.61
26:14:1693:U:O2'	28:19:14:ARG:NH2	2.33	0.61
1:1G:1118:C:H1'	1:1G:1179:A:C4	2.35	0.61
1:1G:45:U:H2'	1:1G:46:G:H8	1.65	0.61
38:55:38:VAL:HG22	38:55:112:ALA:HB2	1.81	0.61
8:72:37:ARG:HH21	8:72:38:ILE:HG13	1.66	0.61
16:7A:21:VAL:HG22	16:7A:33:ILE:HB	1.82	0.61
40:B8:26:ASP:HB3	40:B8:92:GLY:H	1.64	0.61
46:H8:165:VAL:HB	46:H8:166:SER:HA	1.81	0.61
1:13:67:C:H2'	1:13:68:G:C8	2.35	0.61
26:1H:1731:G:H2'	26:1H:1732:A:C8	2.30	0.61
26:1H:2636:U:H2'	26:1H:2637:U:C6	2.35	0.61
25:4K:33:G:H2'	25:4K:34:G:C8	2.35	0.61
20:BA:49:ALA:HA	20:BA:52:ALA:HB3	1.81	0.61
44:F8:3:THR:OG1	44:F8:4:ALA:HA	2.00	0.61
54:Q8:38:GLY:H	54:Q8:41:ILE:HG13	1.66	0.61
26:14:2068:U:H3	26:14:2430:A:H2	1.47	0.61
26:14:2701:C:H3'	26:14:2702:U:C5'	2.27	0.61
26:14:581:C:H2'	26:14:582:G:H8	1.65	0.61
28:19:10:THR:OG1	28:19:13:ARG:HB2	1.99	0.61
1:1G:1449:C:H3'	1:1G:1450:U:H4'	1.82	0.61
26:1H:2784:C:H2'	26:1H:2785:C:H6	1.65	0.61
1:1G:503:C:OP2	12:3A:116:SER:HB3	2.00	0.61
37:45:22:LYS:N	37:45:23:GLY:HA3	2.13	0.61
7:62:62:PHE:HA	7:62:124:LEU:HD22	1.82	0.61
19:AA:40:ILE:O	19:AA:68:GLY:N	2.34	0.61
49:K8:42:GLY:O	49:K8:44:LEU:N	2.31	0.61
34:15:56:ASN:H	34:15:125:GLY:HA3	1.64	0.61
1:1G:91:C:H2'	1:1G:92:G:C8	2.35	0.61
1:13:1367:C:H5'	10:1I:60:ARG:HH21	1.66	0.61
31:49:135:LEU:HB2	31:49:155:MET:HG2	1.82	0.61
26:1H:1278:A:H5''	38:98:36:THR:HG22	1.82	0.61
26:14:489:G:N7	43:A5:49:LYS:NZ	2.48	0.61
46:H8:45:ASP:O	46:H8:49:ARG:HG3	2.01	0.61
1:13:560:U:H4'	1:13:561:U:H5''	1.83	0.61
2:1E:45:GLN:NE2	2:1E:49:GLU:OE1	2.33	0.61
1:1G:382:A:O5'	1:1G:382:A:H8	1.83	0.61
26:1H:1021:A:C8	26:1H:1021:A:H3'	2.35	0.61
26:1H:1891:G:O6	57:1H:4003:HOH:O	2.16	0.61
26:1H:731:C:H2'	26:1H:732:C:H6	1.64	0.61
23:2K:9:G:H21	23:2K:21:A:H8	1.49	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:3E:122:ARG:HH11	4:3E:122:ARG:HG2	1.64	0.61
31:41:133:LEU:HD11	31:41:157:ILE:HB	1.82	0.61
31:41:60:LEU:HD12	31:41:68:PRO:HG3	1.81	0.61
8:72:86:ILE:HD11	8:72:136:GLU:HG2	1.83	0.61
8:7E:95:VAL:HG12	8:7E:99:GLU:HB2	1.83	0.61
9:82:114:TYR:CD2	10:1A:60:ARG:HG3	2.35	0.61
26:1H:2379:G:O2'	39:A8:17:ARG:NH1	2.34	0.61
49:G5:47:ASN:O	49:G5:49:LYS:N	2.29	0.61
36:78:60:MET:HA	54:Q8:13:ARG:NH1	2.16	0.61
28:11:182:LEU:N	28:11:272:ALA:HB3	2.15	0.61
2:12:70:PHE:HB2	2:12:92:TYR:HB2	1.82	0.61
1:13:186:C:H2'	1:13:186(A):C:H6	1.65	0.61
1:13:973:G:H3'	1:13:974:A:H5''	1.81	0.61
26:14:639:U:H2'	26:14:640:C:C6	2.35	0.61
26:14:654(H):G:N7	26:14:654(N):G:N1	2.48	0.61
27:16:80:U:H2'	27:16:81:G:H21	1.65	0.61
26:1H:1140:C:OP1	34:58:23:LEU:HB3	2.01	0.61
26:1H:1416:G:O2'	26:1H:1417:C:O5'	2.18	0.61
12:3A:41:ARG:HH11	12:3A:41:ARG:HB3	1.66	0.61
13:4A:49:THR:HG22	13:4A:51:ALA:H	1.66	0.61
33:69:120:ILE:HG22	33:69:122:GLU:H	1.64	0.61
38:98:63:ARG:HG3	38:98:80:PHE:CE2	2.35	0.61
18:9I:47:THR:HA	18:9I:83:GLU:HB2	1.82	0.61
49:K8:15:LYS:H	49:K8:67:LYS:HE2	1.65	0.61
26:14:2074:U:H2'	26:14:2075:U:C6	2.36	0.61
26:1H:2307:G:O6	31:41:44:GLY:N	2.34	0.61
12:3I:53:ARG:HG3	12:3I:93:LEU:HD21	1.82	0.61
12:3I:71:PRO:O	12:3I:102:ARG:HD3	2.01	0.61
27:1J:90:C:P	37:45:16:ARG:HH21	2.23	0.61
13:4A:33:ALA:O	13:4A:37:THR:OG1	2.14	0.61
5:4E:51:VAL:O	5:4E:55:VAL:HG23	2.00	0.61
18:9I:85:LEU:HD12	18:9I:86:VAL:H	1.65	0.61
46:D5:60:GLU:HA	46:D5:66:SER:HA	1.83	0.61
1:13:1528:U:O2	1:13:1530:G:H5''	2.01	0.61
26:14:1358:G:N1	26:14:1372:U:OP2	2.24	0.61
26:14:1264:G:H2'	26:14:2014:A:N6	2.16	0.61
1:1G:1119:C:H42	1:1G:1154:G:H1	1.48	0.61
4:32:60:GLU:OE2	4:32:199:ASN:N	2.33	0.61
33:69:54:GLN:HA	33:69:57:ARG:HD3	1.83	0.61
9:82:63:ILE:HD11	9:82:81:ILE:HD11	1.82	0.61
49:K8:42:GLY:C	49:K8:44:LEU:H	2.04	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:352:C:O2'	1:13:354:G:OP1	2.18	0.60
26:14:1678:G:N2	26:14:1989:G:N2	2.47	0.60
26:14:2052:G:C8	29:29:141:ILE:HD11	2.35	0.60
26:14:2148:G:H2'	26:14:2149:G:C8	2.36	0.60
26:14:824:A:H1'	26:14:2358:G:N7	2.16	0.60
1:1G:981:U:O2	14:5A:31:ARG:NH2	2.33	0.60
3:2E:16:ARG:NH2	3:2E:183:ASP:OD1	2.34	0.60
36:35:52:GLU:OE2	36:35:57:THR:HA	2.01	0.60
34:58:73:THR:HG22	34:58:84:LYS:HG3	1.82	0.60
7:62:91:VAL:HG12	7:62:95:ARG:HB3	1.81	0.60
41:85:29:SER:OG	41:85:30:LYS:NZ	2.32	0.60
6:52:100:ASN:ND2	18:9A:26:LEU:O	2.33	0.60
19:AI:42:PRO:HD3	51:M8:63:TYR:CE2	2.31	0.60
40:B8:106:SER:HB2	40:B8:109:GLU:HB2	1.83	0.60
26:14:2331:G:O3'	47:E5:43:THR:HG22	2.00	0.60
43:E8:18:ARG:HD3	43:E8:76:VAL:HG22	1.81	0.60
47:I8:37:LEU:HD22	47:I8:67:VAL:HG11	1.83	0.60
1:13:833:U:H3	1:13:853:G:H1	1.48	0.60
26:14:1790:C:H2'	26:14:1791:A:C5	2.36	0.60
26:14:811:U:H2'	36:35:21:ARG:HA	1.83	0.60
1:1G:683:G:N2	1:1G:707:C:O2	2.31	0.60
10:1I:48:THR:HA	10:1I:62:HIS:CB	2.31	0.60
11:2I:15:ALA:HB1	11:2I:78:GLN:HG3	1.82	0.60
15:6A:39:LEU:HD12	15:6A:56:LEU:HB2	1.84	0.60
17:8I:18:THR:OG1	17:8I:69:LYS:NZ	2.33	0.60
43:A5:25:ARG:NH2	43:A5:74:ALA:O	2.34	0.60
52:N8:40:LYS:HZ3	52:N8:46:CYS:HB3	1.66	0.60
1:13:1004:A:O5'	1:13:1025:U:N3	2.35	0.60
1:13:1410:G:H2'	1:13:1411:C:C6	2.36	0.60
26:14:2210:G:H3'	26:14:2211:G:N2	2.15	0.60
27:16:73:A:H3'	27:16:74:U:H6	1.66	0.60
1:1G:1133:G:N2	1:1G:1141:C:N3	2.38	0.60
1:1G:985:C:H2'	1:1G:986:A:C8	2.36	0.60
26:1H:1231:G:H2'	26:1H:1232:G:C8	2.37	0.60
26:1H:1441:G:H2'	26:1H:1442:G:C8	2.36	0.60
26:1H:2341:G:H2'	26:1H:2342:C:C6	2.36	0.60
26:1H:2884:U:H2'	26:1H:2885:C:O4'	2.01	0.60
26:1H:930:U:H4'	26:1H:931:G:O5'	2.01	0.60
30:31:62:ARG:HH21	30:31:64:ILE:HD13	1.65	0.60
36:35:36:LYS:HE3	36:35:39:LYS:HB3	1.84	0.60
1:13:453:A:H4'	16:7I:72:ARG:HB2	1.83	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:82:10:ARG:HD2	9:82:105:ASP:HB3	1.83	0.60
2:12:208:ILE:HA	2:12:211:ILE:HD12	1.84	0.60
1:13:591:U:H2'	1:13:592:G:C8	2.36	0.60
1:1G:1270:C:O2'	1:1G:1313:U:O2'	2.20	0.60
4:32:118:ARG:HH21	4:32:118:ARG:HB2	1.67	0.60
49:K8:18:PRO:HA	49:K8:21:LEU:HB2	1.81	0.60
51:M8:24:THR:OG1	51:M8:25:TYR:N	2.24	0.60
54:Q8:59:LYS:HE3	54:Q8:59:LYS:H	1.67	0.60
28:11:242:ARG:N	28:11:242:ARG:HD2	2.15	0.60
1:13:864:A:H3'	1:13:865:A:C8	2.36	0.60
26:14:1963:U:H5''	26:14:1963:U:O2	2.00	0.60
26:14:274:G:H2'	26:14:275:G:H4'	1.82	0.60
10:1A:9:ARG:HE	10:1A:95:GLU:HG2	1.67	0.60
1:1G:1187:G:O5'	9:82:113:LYS:NZ	2.34	0.60
1:1G:718:G:H5'	11:2A:117:ASN:ND2	2.17	0.60
27:1J:15:A:H1'	27:1J:109:G:C4	2.37	0.60
1:1G:921:U:O2'	5:42:19:MET:O	2.14	0.60
9:82:112:LYS:HA	9:82:119:ALA:HB2	1.84	0.60
9:82:4:TYR:CZ	9:82:88:TYR:HB3	2.36	0.60
44:B5:50:LYS:HB2	44:B5:87:GLN:NE2	2.16	0.60
42:D8:1:MET:SD	42:D8:43:GLU:HB3	2.42	0.60
51:M8:23:GLU:OE1	51:M8:24:THR:N	2.34	0.60
26:14:2328:A:H2'	26:14:2329:G:C8	2.37	0.60
26:1H:2838:G:N7	57:1H:3960:HOH:O	2.30	0.60
36:35:22:GLY:HA3	36:35:25:SER:H	1.66	0.60
36:78:101:VAL:HA	36:78:105:LEU:O	2.01	0.60
8:7E:20:TYR:HE2	8:7E:75:ARG:HD2	1.66	0.60
20:BI:49:ALA:HB2	20:BI:99:LEU:HD23	1.84	0.60
45:G8:30:VAL:HG22	45:G8:37:VAL:HG12	1.84	0.60
1:13:192:U:H2'	1:13:193:C:H6	1.65	0.60
26:1H:1167:U:H2'	26:1H:1168:G:C8	2.37	0.60
29:29:3:GLY:HA3	29:29:81:ILE:HD12	1.82	0.60
36:35:39:LYS:HA	36:35:45:LEU:HD13	1.84	0.60
1:13:407:G:O4'	4:3E:119:GLN:NE2	2.34	0.60
26:14:2745:C:H4'	32:59:142:GLY:O	2.01	0.60
15:6A:11:VAL:HG21	15:6A:34:LEU:HD13	1.83	0.60
39:A8:44:LYS:HB3	39:A8:46:VAL:HB	1.84	0.60
19:AA:50:ALA:HB1	19:AA:57:HIS:HB3	1.83	0.60
40:B8:2:ASN:O	40:B8:3:ARG:HG2	2.01	0.60
1:13:1455:G:H5'	20:BI:32:ALA:HB2	1.81	0.60
26:14:1428:C:O2'	26:14:1569:A:OP2	2.15	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:172:C:H2'	26:14:173:G:C8	2.35	0.60
26:14:469:G:O6	53:L5:37:LYS:HE2	2.01	0.60
26:14:540:G:H2'	26:14:541:C:H6	1.67	0.60
26:14:1491:G:O2'	28:19:101:GLU:HB2	2.02	0.60
28:19:37:LEU:CA	28:19:38:LYS:HB3	2.27	0.60
1:1G:963:G:N2	10:1A:55:LYS:HE3	2.17	0.60
1:1G:922:G:N3	1:1G:1398:A:H2	2.00	0.60
10:1I:34:VAL:HG12	10:1I:74:ILE:HG12	1.82	0.60
29:29:25:VAL:O	29:29:26:ILE:HG12	2.02	0.60
26:14:2394:C:OP1	36:35:64:LYS:HB2	2.01	0.60
5:4E:8:GLU:OE1	5:4E:63:ARG:NH2	2.34	0.60
39:65:21:THR:HG23	39:65:23:ARG:H	1.66	0.60
16:7I:53:VAL:HG13	16:7I:79:VAL:HG22	1.82	0.60
42:95:35:LEU:HB3	42:95:37:VAL:CG1	2.32	0.60
41:C8:85:LYS:HA	41:C8:85:LYS:NZ	2.16	0.60
53:P8:46:VAL:O	53:P8:47:ARG:HB2	2.02	0.60
54:Q8:33:ASN:HB2	54:Q8:34:TRP:CE3	2.37	0.60
1:13:1053:G:H4'	1:13:1054:C:O5'	2.02	0.60
1:13:1273:G:H3'	1:13:1274:G:C8	2.36	0.60
26:14:192:C:P	57:14:3644:HOH:O	2.60	0.60
26:14:579:G:H2'	26:14:580:C:C6	2.36	0.60
10:1A:50:ILE:HA	10:1A:60:ARG:HB3	1.84	0.60
1:1G:976:G:H5'	1:1G:1358:U:O2'	2.02	0.60
26:1H:1024:G:H3'	26:1H:1025:G:H5"	1.84	0.60
26:1H:1113:U:H2'	26:1H:1114:G:C8	2.37	0.60
26:1H:232:G:H5"	26:1H:232:G:H8	1.66	0.60
27:1J:18:G:H1	27:1J:65:C:H42	1.48	0.60
35:25:47:ILE:HG13	35:25:48:PRO:HD2	1.84	0.60
3:2E:89:GLU:OE2	3:2E:93:LYS:NZ	2.35	0.60
12:3A:47:LYS:HG2	12:3A:48:PRO:HD2	1.84	0.60
12:3I:82:VAL:HG12	12:3I:106:ASP:OD2	2.02	0.60
6:52:74:ASP:N	6:52:74:ASP:OD1	2.34	0.60
34:58:26:LEU:O	34:58:30:ILE:HG13	2.01	0.60
6:5E:18:GLN:HA	6:5E:21:LEU:HB2	1.84	0.60
9:82:97:LYS:HB3	9:82:98:PRO:HD3	1.84	0.60
40:B8:26:ASP:O	40:B8:49:VAL:HG12	2.02	0.60
27:1J:40:U:C5	51:I5:1:MET:HB3	2.35	0.60
2:12:103:THR:HG23	2:12:176:GLU:HB3	1.83	0.60
1:13:1292:U:H2'	1:13:1293:G:C8	2.36	0.60
1:13:1327:C:OP2	21:1F:12:LYS:NZ	2.34	0.60
1:13:181:G:O2'	1:13:182:U:O5'	2.19	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:290:G:H2'	26:14:291:C:O4'	2.02	0.60
26:14:528:A:OP2	34:15:114:ARG:NH1	2.34	0.60
26:14:620:G:H5'	26:14:620:G:N3	2.16	0.60
34:15:42:TRP:O	41:85:64:ARG:NH2	2.34	0.60
1:1G:382:A:H2'	1:1G:383:A:H8	1.65	0.60
26:1H:2162:G:H2'	26:1H:2163:C:C6	2.36	0.60
26:1H:2693:A:H2'	26:1H:2694:G:H8	1.67	0.60
26:1H:2789:C:H2'	26:1H:2790:A:H5''	1.84	0.60
29:29:8:LYS:HG2	29:29:192:ASN:HA	1.83	0.60
32:59:7:LEU:HD12	32:59:8:PRO:HD3	1.83	0.60
14:5I:6:LEU:O	14:5I:23:ARG:NH2	2.35	0.60
41:85:92:ARG:HG2	42:95:11:GLN:OE1	2.01	0.60
53:L5:35:ARG:HG3	53:L5:42:LEU:HD11	1.84	0.60
1:13:1279:A:OP2	10:1I:9:ARG:NH1	2.35	0.59
26:1H:2291:U:O2'	26:1H:2374:C:O2	2.18	0.59
26:1H:2305:A:H1'	31:41:136:ARG:HH12	1.66	0.59
26:1H:384:U:H2'	26:1H:385:C:H6	1.67	0.59
26:1H:883:G:N2	26:1H:893:C:O2	2.33	0.59
26:1H:946:G:OP1	57:1H:4106:HOH:O	2.17	0.59
30:39:31:HIS:HB2	36:35:9:ASN:OD1	2.02	0.59
40:75:107:ASP:OD1	40:75:109:GLU:HB2	2.02	0.59
20:BA:50:GLU:HA	20:BA:100:ILE:HG21	1.84	0.59
45:C5:52:SER:HA	45:C5:55:TYR:O	2.02	0.59
1:13:1312:G:OP1	51:M8:58:ARG:NH2	2.35	0.59
54:Q8:33:ASN:HB2	54:Q8:34:TRP:CD2	2.37	0.59
1:13:37:U:O2'	1:13:500:G:H4'	2.02	0.59
1:13:619:U:H3	4:3E:134:ASP:HB2	1.65	0.59
26:14:1786:A:H2	26:14:2606:C:H1'	1.65	0.59
26:14:646:A:H2'	26:14:647:G:O4'	2.03	0.59
26:14:795:C:H2'	26:14:796:C:C6	2.37	0.59
28:19:182:LEU:H	28:19:272:ALA:CB	2.15	0.59
2:1E:74:LYS:HB2	2:1E:76:GLN:HG3	1.83	0.59
1:1G:86:U:H6	1:1G:87:A:C2	2.20	0.59
26:1H:2062:A:H62	26:1H:2503:A:H62	1.47	0.59
26:1H:270(J):G:N2	26:1H:270(Q):C:N3	2.50	0.59
26:1H:676:A:H8	26:1H:2069:G:N2	1.95	0.59
26:1H:84:A:OP2	45:G8:8:LYS:NZ	2.28	0.59
4:32:173:TRP:CD1	4:32:174:LEU:HG	2.37	0.59
30:39:27:GLU:O	30:39:28:ILE:HG12	2.01	0.59
31:49:115:ARG:NH2	31:49:137:GLU:OE1	2.35	0.59
6:52:2:ARG:HD3	6:52:92:LYS:HE3	1.82	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:59:71:LEU:O	32:59:75:ALA:N	2.35	0.59
26:1H:566:U:O4	42:D8:78:LYS:HD3	2.02	0.59
45:G8:87:LYS:H	45:G8:94:LYS:HG2	1.67	0.59
26:14:1443:G:H1	26:14:1548:C:H42	1.49	0.59
34:15:56:ASN:HA	34:15:125:GLY:H	1.66	0.59
28:19:16:MET:HE1	28:19:208:LYS:HE3	1.84	0.59
2:1E:115:LEU:HA	2:1E:118:LEU:HD12	1.84	0.59
1:1G:1536:C:H2'	1:1G:1537:U:C5	2.36	0.59
26:1H:2199:A:O3'	48:J8:48:LYS:HD3	2.02	0.59
26:1H:2400:G:H1	26:1H:2416:C:H42	1.50	0.59
26:1H:2899:G:N2	26:1H:2900:A:N1	2.50	0.59
26:1H:860:U:C5	26:1H:917:A:C2	2.90	0.59
35:25:4:PRO:O	35:25:5:GLN:HB2	2.02	0.59
3:2E:26:LYS:HG3	3:2E:27:LYS:HE2	1.83	0.59
30:31:158:THR:OG1	30:31:195:ASP:OD2	2.20	0.59
4:32:61:LYS:HZ3	4:32:206:PHE:HE2	1.48	0.59
31:41:97:ASP:O	31:41:99:MET:N	2.34	0.59
15:6I:78:TYR:CZ	15:6I:82:ILE:HD11	2.37	0.59
42:95:35:LEU:HD13	42:95:37:VAL:HG11	1.84	0.59
39:A8:93:LYS:HG2	39:A8:95:HIS:HB2	1.84	0.59
45:C5:20:TYR:CZ	45:C5:42:VAL:HA	2.37	0.59
26:1H:751:A:H5'	43:E8:90:ARG:HA	1.83	0.59
57:1H:3789:HOH:O	44:F8:56:THR:O	2.15	0.59
26:1H:480:A:OP2	45:G8:47:LYS:HD3	2.02	0.59
51:M8:17:GLY:HA3	51:M8:35:VAL:HG23	1.84	0.59
54:Q8:38:GLY:N	54:Q8:41:ILE:HG13	2.17	0.59
54:Q8:53:PRO:HB3	54:Q8:56:GLU:N	2.16	0.59
2:12:42:ILE:HD11	2:12:202:PRO:HB2	1.83	0.59
1:13:1391:U:H2'	1:13:1392:G:C8	2.38	0.59
1:13:196:A:O2'	1:13:197:A:H2'	2.03	0.59
26:14:2836:U:H2'	26:14:2837:G:C8	2.37	0.59
26:14:539:G:H2'	26:14:540:G:H8	1.67	0.59
1:1G:1305:G:H22	1:1G:1331:G:C2'	2.12	0.59
26:1H:1515:C:H2'	26:1H:1516:U:C6	2.36	0.59
26:1H:259:G:H21	26:1H:621:A:H8	1.49	0.59
27:1J:104:A:H2'	27:1J:105:G:O4'	2.03	0.59
9:8E:128:ARG:NH1	23:2K:34:U:OP2	2.27	0.59
4:3E:65:ARG:NH1	4:3E:70:ILE:O	2.34	0.59
9:82:95:LYS:HZ1	9:82:96:LEU:HD13	1.67	0.59
19:AA:23:ASN:HA	19:AA:27:GLU:HG3	1.83	0.59
1:13:789:U:H5	1:13:791:G:H3'	1.67	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1329:U:H5''	26:14:1330:C:H5	1.67	0.59
1:1G:406:G:O6	1:1G:436:C:N4	2.32	0.59
1:1G:677:U:H3	1:1G:713:G:H22	1.49	0.59
26:1H:1102:C:H2'	26:1H:1103:A:C8	2.37	0.59
26:1H:1166:C:H2'	26:1H:1167:U:C6	2.37	0.59
26:1H:1262:A:N3	52:N8:10:LYS:HE3	2.17	0.59
26:1H:1454:U:H5	38:98:73:VAL:HG13	1.67	0.59
10:1I:77:PRO:HB2	10:1I:79:ARG:HH12	1.67	0.59
13:4I:3:ARG:HE	13:4I:9:ILE:HD11	1.67	0.59
41:85:91:ASP:O	41:85:92:ARG:HG3	2.03	0.59
36:35:63:PRO:HB3	54:M5:30:ARG:NH1	2.16	0.59
26:14:1336:A:H2'	26:14:1337:G:C8	2.38	0.59
26:14:2748:A:H2'	26:14:2749:A:H8	1.66	0.59
3:2E:20:SER:OG	3:2E:40:ARG:NH2	2.33	0.59
4:32:139:ARG:HG3	4:32:139:ARG:HH11	1.67	0.59
24:3K:6:G:O2'	24:3K:7:G:O4'	2.21	0.59
38:55:38:VAL:HG12	38:55:42:LYS:HD2	1.85	0.59
14:5I:15:LYS:HG2	14:5I:16:PHE:HD2	1.67	0.59
27:1J:50:G:OP1	39:65:62:LYS:HB2	2.03	0.59
7:6E:62:PHE:CD1	7:6E:124:LEU:HD11	2.38	0.59
9:82:113:LYS:H	9:82:119:ALA:HB2	1.67	0.59
37:88:135:ASP:HB3	37:88:137:TYR:H	1.68	0.59
45:C5:15:VAL:HG12	45:C5:21:LYS:HA	1.84	0.59
2:12:5:ILE:HA	2:12:221:LEU:HD11	1.85	0.59
26:14:1534:G:H3'	26:14:1535:U:H5''	1.85	0.59
26:14:2158:A:H1'	26:14:2159:G:C8	2.37	0.59
26:14:459:U:H2'	26:14:460:A:H8	1.68	0.59
1:1G:963:G:H21	10:1A:55:LYS:HE3	1.67	0.59
2:1E:7:VAL:HG11	2:1E:217:ARG:NH1	2.17	0.59
1:1G:382:A:H2'	1:1G:383:A:C8	2.37	0.59
29:21:70:ALA:O	29:21:73:GLU:N	2.35	0.59
4:3E:150:GLU:HA	4:3E:153:ARG:HG2	1.84	0.59
31:41:113:ARG:HD2	51:M8:33:VAL:HG13	1.85	0.59
33:69:100:ALA:O	33:69:104:GLN:HB3	2.03	0.59
33:69:14:ASP:OD1	33:69:15:VAL:N	2.36	0.59
49:K8:50:ILE:HD12	49:K8:51:ARG:H	1.68	0.59
1:13:1348:U:H4'	9:8E:120:ARG:HD2	1.83	0.59
1:13:861:G:H4'	8:7E:18:ARG:NH2	2.18	0.59
26:14:1364:G:OP1	48:F5:3:LYS:HD2	2.02	0.59
26:14:1639:U:H4'	26:14:2699:C:H4'	1.85	0.59
1:1G:1134:G:C2	1:1G:1135:U:H1'	2.38	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1154:G:H2'	1:1G:1155:G:C8	2.38	0.59
26:1H:1171:G:N7	26:1H:1174:A:N6	2.51	0.59
26:1H:443:A:H1'	26:1H:1201:C:O4'	2.02	0.59
10:1I:8:LEU:HD22	10:1I:96:ILE:HG22	1.85	0.59
23:2K:19:G:O3'	23:2K:20:U:H2'	2.03	0.59
9:82:92:TYR:O	9:82:95:LYS:NZ	2.23	0.59
38:98:48:VAL:HA	38:98:51:LEU:HB2	1.83	0.59
54:Q8:53:PRO:HA	54:Q8:54:GLU:C	2.21	0.59
1:13:1034:G:N2	1:13:1035:A:N7	2.50	0.59
26:14:467:G:OP1	53:L5:33:ARG:NH1	2.36	0.59
26:14:674:G:O2'	30:39:74:ARG:HG3	2.03	0.59
27:16:30:C:H2'	27:16:31:C:H5'	1.84	0.59
10:1A:30:SER:HB2	10:1A:81:THR:HA	1.83	0.59
26:1H:1058:U:H3	26:1H:1080:A:N6	2.01	0.59
26:1H:2735:G:H2'	26:1H:2736:G:C8	2.37	0.59
29:21:2:LYS:HA	29:21:84:PHE:CD1	2.37	0.59
23:2L:44:G:H2'	23:2L:45:U:O4'	2.03	0.59
1:1G:9:G:OP1	5:42:122:GLU:HB2	2.02	0.59
5:42:24:ARG:NH1	5:42:24:ARG:HB3	2.17	0.59
32:59:81:GLU:HG2	32:59:83:TYR:H	1.68	0.59
9:82:17:VAL:HG22	9:82:63:ILE:HG12	1.85	0.59
1:1G:235:C:H5'	17:8A:70:ARG:HG2	1.85	0.59
44:B5:36:LYS:HA	44:B5:39:ILE:HD12	1.83	0.59
20:BA:54:LYS:HA	20:BA:57:ARG:HH12	1.68	0.59
49:G5:10:LEU:HD13	49:G5:59:ARG:HD2	1.85	0.59
28:11:206:LEU:O	28:11:211:ARG:HD3	2.02	0.59
1:13:1130:A:O2'	9:8E:3:GLN:NE2	2.36	0.59
1:13:359:U:H2'	1:13:360:A:C8	2.37	0.59
1:13:57:G:H2'	1:13:58:C:C6	2.38	0.59
1:13:651:C:H2'	1:13:652:U:C6	2.38	0.59
26:14:2143:C:H2'	26:14:2144:U:H4'	1.83	0.59
26:14:708:C:H42	26:14:723:G:H1	1.51	0.59
10:1A:17:ASP:OD2	10:1A:70:ARG:NH1	2.36	0.59
1:1G:1451:A:H5''	1:1G:1452:C:C5	2.38	0.59
26:1H:2883:A:H5'	26:1H:2884:U:H5'	1.84	0.59
10:1I:5:ARG:HB2	10:1I:73:ASP:OD1	2.03	0.59
3:22:150:LYS:NZ	3:22:152:ILE:HD11	2.18	0.59
30:39:7:TYR:HD1	30:39:18:ARG:H	1.51	0.59
13:4A:80:ARG:NH1	19:AA:66:MET:HG2	2.17	0.59
6:52:11:ASN:HB3	6:52:14:LEU:HD13	1.84	0.59
8:72:12:ARG:HD3	8:72:26:VAL:HG12	1.84	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1167:A:H2'	1:13:1169:A:C8	2.38	0.58
1:13:166:G:H2'	1:13:167:G:C8	2.37	0.58
26:14:1066:U:H1'	26:14:1073:A:H61	1.68	0.58
26:1H:1021:A:H8	26:1H:1021:A:H3'	1.68	0.58
26:1H:1259:G:H2'	26:1H:1260:G:H8	1.68	0.58
26:1H:1899:G:N2	26:1H:1902:C:C5	2.71	0.58
26:1H:2099:U:H2'	26:1H:2100:G:C8	2.38	0.58
3:22:112:SER:HB3	3:22:115:LEU:HB2	1.85	0.58
11:2A:41:THR:OG1	11:2A:42:TRP:N	2.36	0.58
36:35:122:PRO:HB3	36:35:141:ALA:HB1	1.85	0.58
4:3E:108:LEU:HD23	4:3E:110:PHE:HE1	1.68	0.58
13:4A:29:ARG:HD2	13:4A:64:TRP:CE2	2.38	0.58
13:4I:87:TYR:O	13:4I:91:ARG:HG2	2.02	0.58
2:12:166:ASP:OD2	2:12:169:LYS:HB2	2.02	0.58
26:14:1486:A:H2'	26:14:1487:G:C8	2.38	0.58
1:1G:1343:G:H2'	1:1G:1344:C:C6	2.38	0.58
1:1G:474:G:H2'	1:1G:475:G:C8	2.38	0.58
26:1H:2150:U:H2'	26:1H:2151:G:C8	2.38	0.58
26:1H:2309:A:C5	26:1H:2310:A:C8	2.91	0.58
26:1H:518:G:H2'	26:1H:519:U:H6	1.69	0.58
26:1H:945:A:OP2	26:1H:945:A:H4'	2.03	0.58
30:39:143:ALA:O	30:39:148:LEU:HB2	2.02	0.58
13:4I:49:THR:HB	13:4I:52:GLU:HG3	1.85	0.58
32:59:15:VAL:HG12	32:59:29:PRO:HD2	1.85	0.58
27:1J:50:G:OP1	39:65:63:THR:HG23	2.04	0.58
36:78:122:PRO:HA	36:78:142:GLY:HA3	1.84	0.58
26:14:996:A:H4'	41:85:92:ARG:CZ	2.33	0.58
26:1H:1252:G:O4'	41:C8:33:ARG:HD3	2.03	0.58
54:Q8:28:GLY:HA2	54:Q8:31:HIS:NE2	2.18	0.58
26:14:1253:A:OP1	57:14:3839:HOH:O	2.17	0.58
26:14:1342:A:H2	26:14:1602:U:H3	1.51	0.58
26:14:2371:G:O6	57:14:3861:HOH:O	2.14	0.58
1:1G:1122:U:O4	1:1G:1123:A:N6	2.37	0.58
1:1G:668:G:O2'	15:6A:46:HIS:HB3	2.04	0.58
1:1G:838:G:H1	1:1G:848:C:H42	1.51	0.58
26:1H:1512:G:H2'	26:1H:1513:C:C6	2.38	0.58
26:1H:2228:G:OP2	28:11:263:ARG:NH2	2.36	0.58
26:1H:2469:A:H61	26:1H:2481:G:H1'	1.68	0.58
23:2L:32:C:O2'	23:2L:33:U:OP1	2.15	0.58
12:3I:110:VAL:HG23	12:3I:120:TYR:HB3	1.85	0.58
13:4I:15:VAL:HG23	13:4I:43:THR:O	2.03	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:51:83:TYR:HB3	32:51:134:SER:HA	1.86	0.58
14:5I:29:ARG:HH21	14:5I:42:ILE:HD11	1.66	0.58
26:1H:2359:C:H5'	54:Q8:49:VAL:HG21	1.85	0.58
10:1A:47:PHE:O	10:1A:63:PHE:N	2.24	0.58
21:1B:2:GLY:O	21:1B:4:GLY:N	2.36	0.58
26:1H:1657:C:H2'	26:1H:1658:C:H6	1.68	0.58
26:1H:71:A:C2	44:F8:31:HIS:CE1	2.91	0.58
3:22:129:ALA:HB3	3:22:132:ARG:HB3	1.85	0.58
29:29:111:ARG:HB2	29:29:160:TYR:HB3	1.85	0.58
23:2L:29:U:H2'	23:2L:30:A:H8	1.68	0.58
26:14:943:U:OP2	36:35:36:LYS:HG3	2.03	0.58
13:4A:82:MET:SD	13:4A:83:ASP:N	2.76	0.58
7:6E:111:ARG:HD2	7:6E:123:GLU:HB2	1.85	0.58
8:72:99:GLU:OE2	8:72:100:ILE:N	2.21	0.58
36:78:19:VAL:CB	36:78:27:HIS:HB2	2.33	0.58
26:1H:2599:G:N7	28:11:237:GLU:HG2	2.19	0.58
28:11:85:ASP:OD2	28:11:88:ARG:NH1	2.27	0.58
1:13:1348:U:H2'	1:13:1349:A:H8	1.69	0.58
26:14:221:A:N6	26:14:265:A:H8	2.00	0.58
26:14:2853:C:H2'	26:14:2854:G:C8	2.38	0.58
26:14:67:U:H3	26:14:74:A:H2	1.50	0.58
34:15:67:LEU:O	34:15:88:GLU:HG3	2.03	0.58
26:1H:1386:C:H2'	26:1H:1387:C:C6	2.36	0.58
26:1H:729:G:OP2	28:11:13:ARG:NH1	2.29	0.58
26:1H:960:A:C8	26:1H:962:G:C8	2.91	0.58
4:32:24:GLU:HG2	4:32:25:ARG:H	1.67	0.58
32:59:10:PRO:HD3	32:59:50:VAL:O	2.03	0.58
33:69:77:LEU:HD12	33:69:78:THR:H	1.68	0.58
40:75:107:ASP:CG	40:75:109:GLU:HB2	2.24	0.58
16:7I:19:ILE:HB	16:7I:36:ILE:O	2.02	0.58
9:8E:21:PRO:HA	9:8E:59:PHE:HA	1.86	0.58
20:BI:86:ARG:HH12	20:BI:90:GLN:HB2	1.69	0.58
26:14:1593:G:H2'	26:14:1594:G:C8	2.39	0.58
1:1G:1043:C:H2'	1:1G:1044:A:H8	1.69	0.58
1:1G:353:A:H5'	1:1G:353:A:C8	2.34	0.58
26:1H:1858:G:O2'	26:1H:1859:A:O5'	2.22	0.58
22:1K:18:G:H4'	22:1K:19:G:OP1	2.03	0.58
36:35:47:ASP:OD2	36:35:50:ARG:NH1	2.36	0.58
30:39:18:ARG:HG2	30:39:19:GLU:H	1.67	0.58
39:A8:58:LEU:HD12	39:A8:65:VAL:HG13	1.85	0.58
19:AI:5:LEU:HD13	19:AI:10:PHE:CD1	2.38	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
46:D5:48:PHE:HA	46:D5:51:ALA:HB3	1.84	0.58
2:12:174:VAL:HG11	2:12:196:LEU:HD13	1.86	0.58
2:12:233:SER:HB2	2:12:234:PRO:HD2	1.85	0.58
1:13:758:G:O2'	1:13:759:A:H5'	2.04	0.58
1:13:975:A:H8	1:13:975:A:H5'	1.69	0.58
26:14:139:G:N2	26:14:141:A:N1	2.48	0.58
26:14:271:G:H2'	26:14:272:G:C8	2.39	0.58
26:1H:2120:G:N2	26:1H:2178:C:O2	2.32	0.58
24:3L:42:U:H2'	24:3L:43:A:H8	1.69	0.58
26:14:2873:A:C8	38:55:5:LYS:HA	2.38	0.58
29:29:9:VAL:HA	40:75:3:ARG:HG3	1.85	0.58
47:E5:21:LEU:HD11	47:E5:41:ARG:NH1	2.19	0.58
31:49:4:ASP:OD2	51:15:25:TYR:OH	2.22	0.58
1:13:1366:C:H2'	1:13:1367:C:C6	2.39	0.58
1:13:17:U:H2'	1:13:18:C:C6	2.39	0.58
21:1F:15:ARG:HH11	21:1F:15:ARG:HB2	1.68	0.58
1:1G:1142:G:H3'	1:1G:1143:G:H8	1.68	0.58
1:1G:222:U:H2'	1:1G:223:U:H6	1.67	0.58
26:1H:1264:G:OP1	52:N8:19:ARG:NH2	2.27	0.58
26:1H:1983:C:H2'	26:1H:1984:G:H5''	1.85	0.58
4:32:93:PHE:O	4:32:96:LEU:HB2	2.04	0.58
4:3E:147:ALA:HB2	4:3E:182:LYS:HB3	1.85	0.58
1:13:35:G:O2'	12:3I:118:SER:O	2.11	0.58
12:3I:60:LEU:HB2	12:3I:64:TYR:HB2	1.86	0.58
6:5E:3:ARG:NH1	6:5E:38:GLU:OE2	2.36	0.58
1:13:1239:A:O2'	7:6E:114:ARG:O	2.18	0.58
8:7E:39:LEU:HB3	8:7E:45:ILE:HG12	1.85	0.58
38:98:56:LYS:NZ	38:98:90:ARG:O	2.36	0.58
50:H5:7:LYS:HE2	50:H5:32:GLN:HG3	1.85	0.58
46:H8:3:TYR:O	46:H8:58:VAL:HG13	2.03	0.58
1:13:1349:A:H2'	1:13:1350:A:H8	1.69	0.58
1:13:864:A:H3'	1:13:865:A:H8	1.68	0.58
26:14:1534:G:H3'	26:14:1535:U:C5'	2.34	0.58
26:14:459:U:H5''	53:L5:40:TRP:CE2	2.39	0.58
1:1G:736:C:H2'	1:1G:737:A:C8	2.39	0.58
26:1H:1523:U:H2'	26:1H:1524:G:O4'	2.04	0.58
3:2E:104:GLN:OE1	3:2E:107:GLN:NE2	2.37	0.58
30:31:197:ASP:O	30:31:199:TRP:N	2.37	0.58
5:4E:79:GLU:HG3	5:4E:93:PRO:HD2	1.85	0.58
29:29:111:ARG:HA	38:55:2:ARG:HH12	1.68	0.58
7:62:67:GLU:OE2	7:62:70:LYS:NZ	2.36	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:6E:74:GLU:HG2	7:6E:91:VAL:HG22	1.85	0.58
49:K8:33:MET:HG3	49:K8:36:ARG:HH21	1.68	0.58
1:13:1128:C:O2'	1:13:1129:C:H5''	2.02	0.58
1:13:376:G:H1	1:13:387:U:H3	1.51	0.58
26:14:1388:G:H2'	26:14:1389:G:H8	1.68	0.58
26:14:1794:U:H2'	26:14:1795:C:C6	2.38	0.58
1:1G:720:C:H3'	1:1G:721:G:H2'	1.85	0.58
26:1H:1913:A:H4'	26:1H:1914:C:H5''	1.86	0.58
26:1H:2131:G:H5''	26:1H:2133:G:H4'	1.86	0.58
26:1H:2467:C:O2	37:88:124:LYS:NZ	2.37	0.58
23:2L:44:G:H8	23:2L:44:G:H5'	1.68	0.58
4:32:157:LEU:HD12	4:32:161:ASN:HD21	1.69	0.58
7:6E:118:VAL:HG23	7:6E:122:HIS:NE2	2.19	0.58
20:BA:26:ASN:O	20:BA:30:LYS:HB2	2.03	0.58
41:C8:83:LEU:HD13	41:C8:113:ALA:HB2	1.86	0.58
43:E8:29:LEU:HD21	43:E8:33:ARG:NH2	2.18	0.58
48:J8:82:LEU:HD22	48:J8:82:LEU:H	1.68	0.58
1:13:486:U:H2'	1:13:487:A:C8	2.38	0.57
1:13:861:G:H4'	8:7E:18:ARG:HH21	1.68	0.57
26:14:1418:G:OP1	26:14:1588:C:O2'	2.22	0.57
26:14:2849:U:H1'	26:14:2866:U:O2	2.04	0.57
26:14:30:G:H2'	26:14:31:C:C6	2.39	0.57
2:1E:211:ILE:HA	2:1E:214:ILE:HD12	1.86	0.57
1:1G:345:C:H1'	1:1G:346:G:C2	2.39	0.57
26:1H:1657:C:H2'	26:1H:1658:C:C6	2.38	0.57
26:1H:1814:G:O6	57:1H:3815:HOH:O	2.17	0.57
29:21:116:VAL:O	29:21:117:MET:HB3	2.04	0.57
23:2L:38:1MG:H2'	23:2L:39:PSU:C6	2.39	0.57
30:39:146:ALA:HB1	30:39:148:LEU:HG	1.86	0.57
30:39:78:ILE:HA	30:39:83:PHE:CD2	2.39	0.57
24:3K:21:A:N6	24:3K:46:G:H1'	2.19	0.57
46:D5:52:SER:O	46:D5:54:HIS:N	2.36	0.57
50:H5:39:ASP:O	50:H5:44:ARG:NH2	2.36	0.57
26:1H:2360:A:OP1	54:Q8:49:VAL:HB	2.03	0.57
1:13:1412:C:H2'	1:13:1413:A:C8	2.39	0.57
26:14:2698:U:H2'	26:14:2699:C:C6	2.38	0.57
26:14:848:G:C4	26:14:933:A:H8	2.22	0.57
26:1H:1932:A:H2'	26:1H:1933:G:O4'	2.03	0.57
26:1H:2331:G:O3'	47:I8:43:THR:HG22	2.03	0.57
11:2A:57:THR:HG22	11:2A:59:TYR:H	1.69	0.57
23:2L:38:1MG:H2'	23:2L:39:PSU:H6	1.69	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:3I:58:VAL:O	12:3I:65:GLU:HA	2.04	0.57
24:3L:31:C:O2	24:3L:41:G:N2	2.34	0.57
31:41:35:GLU:OE1	31:41:36:LYS:N	2.37	0.57
32:51:43:VAL:HB	32:51:52:VAL:HG23	1.85	0.57
32:51:83:TYR:HD1	32:51:84:SER:H	1.50	0.57
32:51:86:GLU:HG2	32:51:87:LEU:H	1.68	0.57
32:59:129:THR:HB	32:59:130:ARG:HG3	1.86	0.57
32:59:74:ASN:O	32:59:78:GLY:N	2.38	0.57
33:69:14:ASP:O	33:69:17:GLN:HB2	2.03	0.57
9:8E:97:LYS:HB2	9:8E:102:LEU:HD12	1.85	0.57
27:16:12:C:O2	47:I8:74:ARG:NH1	2.35	0.57
28:11:2:ALA:HB1	28:11:200:ASP:OD2	2.03	0.57
1:13:1006:C:O2	1:13:1023:G:N2	2.33	0.57
1:13:670:G:H21	6:5E:73:ASN:HD21	1.52	0.57
26:14:2365:G:O6	54:M5:39:LYS:NZ	2.37	0.57
26:14:2819:G:O6	57:14:4000:HOH:O	2.16	0.57
26:14:2861:G:C2	26:14:2862:G:C4	2.92	0.57
1:1G:963:G:H21	10:1A:55:LYS:CE	2.17	0.57
26:1H:2857:G:N2	26:1H:2860:A:OP2	2.31	0.57
26:1H:771:G:OP1	53:P8:10:ARG:NH1	2.38	0.57
26:1H:833:U:O2	36:78:55:ARG:NH1	2.36	0.57
16:7I:51:VAL:HG21	16:7I:74:LEU:HD21	1.85	0.57
19:AA:9:VAL:HG13	51:I5:63:TYR:CE1	2.39	0.57
41:C8:88:ILE:C	41:C8:90:VAL:H	2.08	0.57
28:11:72:LYS:HG2	28:11:103:ARG:NH2	2.19	0.57
26:14:2853:C:H2'	26:14:2854:G:H8	1.68	0.57
26:1H:1013:C:H42	26:1H:1149:G:H1	1.50	0.57
26:1H:1055:G:O2'	26:1H:1086:A:N6	2.37	0.57
26:1H:1138:G:H21	34:58:106:MET:HE3	1.70	0.57
29:21:67:PHE:O	29:21:69:LYS:HD2	2.04	0.57
31:49:161:THR:HG22	31:49:163:ALA:H	1.69	0.57
26:1H:2840:C:H5''	38:98:53:HIS:CD2	2.39	0.57
1:13:1286:A:H2'	1:13:1287:A:H4'	1.85	0.57
1:13:690:G:H2'	1:13:691:G:O4'	2.04	0.57
26:14:1188:U:H4'	42:95:79:VAL:HG12	1.87	0.57
26:14:2121:G:O6	26:14:2177:C:N4	2.37	0.57
1:1G:1106:G:H5''	3:22:172:ARG:HG2	1.86	0.57
26:1H:1538:G:H2'	26:1H:1539:G:H8	1.70	0.57
29:29:182:LEU:O	29:29:183:LEU:HD12	2.04	0.57
30:31:29:ASN:H	30:31:112:MET:CE	2.17	0.57
57:1H:3907:HOH:O	30:31:55:GLY:HA2	2.04	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:32:24:GLU:OE1	4:32:112:VAL:HG21	2.04	0.57
26:14:2414:G:H21	36:35:67:MET:CE	2.17	0.57
31:41:173:LEU:HB3	31:41:178:PHE:HD2	1.69	0.57
26:14:2873:A:H8	38:55:6:SER:H	1.49	0.57
49:K8:15:LYS:H	49:K8:15:LYS:HE3	1.70	0.57
51:M8:43:TYR:CG	51:M8:44:THR:N	2.72	0.57
1:13:108:G:H5''	1:13:109:A:H5''	1.84	0.57
1:13:1510:U:H2'	1:13:1511:G:C8	2.40	0.57
26:14:1386:C:H2'	26:14:1387:C:C6	2.39	0.57
26:14:2318:G:N2	39:65:3:ARG:HG2	2.20	0.57
26:14:244:A:C2	26:14:255:A:C4	2.92	0.57
26:14:71:A:C8	26:14:71:A:H5'	2.40	0.57
28:19:32:SER:O	28:19:33:LEU:HB2	2.04	0.57
1:1G:1126:U:C4	1:1G:1281:U:C6	2.92	0.57
1:1G:689:C:H2'	1:1G:690:G:H5'	1.87	0.57
32:51:4:ILE:HD13	32:51:4:ILE:H	1.68	0.57
32:51:3:ARG:HH21	32:51:7:LEU:HD11	1.69	0.57
6:5E:45:LEU:HD12	6:5E:59:TYR:HD2	1.69	0.57
39:65:89:ARG:O	39:65:92:TYR:N	2.38	0.57
35:68:75:SER:HB2	40:B8:74:ARG:HH12	1.69	0.57
44:F8:32:PRO:HA	44:F8:77:LYS:HD2	1.86	0.57
49:G5:25:VAL:HA	49:G5:28:LYS:HB2	1.86	0.57
51:M8:16:CYS:SG	51:M8:17:GLY:N	2.75	0.57
1:13:624:C:H2'	1:13:625:G:C8	2.39	0.57
26:14:2471:C:N4	26:14:2476:A:O2'	2.38	0.57
34:15:35:ARG:HB2	34:15:42:TRP:CZ3	2.39	0.57
28:19:17:THR:O	28:19:211:ARG:NH2	2.37	0.57
1:1G:1016:A:O2'	1:1G:1217:C:O2'	2.21	0.57
1:1G:1117:G:H4'	9:82:104:ARG:HD2	1.86	0.57
1:1G:1239:A:H4'	1:1G:1240:U:H5'	1.85	0.57
1:1G:920:U:H2'	1:1G:921:U:C6	2.39	0.57
26:1H:732:C:H3'	57:1H:3837:HOH:O	2.05	0.57
29:21:119:ARG:HD3	29:21:160:TYR:HB2	1.86	0.57
12:3I:66:VAL:HG22	12:3I:67:THR:H	1.69	0.57
24:3L:6:G:O2'	24:3L:7:G:O4'	2.23	0.57
5:4E:139:LEU:HA	5:4E:142:LEU:HD12	1.86	0.57
33:61:124:GLY:H	33:61:142:VAL:HG23	1.69	0.57
33:61:63:ALA:HA	33:61:66:GLU:HG2	1.86	0.57
9:8E:47:LEU:HD22	9:8E:47:LEU:H	1.70	0.57
38:98:12:ARG:O	38:98:17:ARG:NH2	2.37	0.57
40:B8:26:ASP:HB2	40:B8:91:ARG:HA	1.86	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
50:L8:9:VAL:HG22	50:L8:54:VAL:HA	1.86	0.57
52:N8:30:LEU:HD23	52:N8:41:PRO:HA	1.84	0.57
54:Q8:50:LEU:C	54:Q8:52:LYS:N	2.58	0.57
1:13:1227:A:OP1	19:AI:80:TYR:OH	2.16	0.57
26:14:1062:G:H21	26:14:1088:A:H2	1.52	0.57
1:1G:146:G:H2'	1:1G:147:G:H8	1.69	0.57
26:1H:1515:C:H2'	26:1H:1516:U:H6	1.69	0.57
29:21:105:THR:HG22	29:21:106:GLY:H	1.70	0.57
40:75:23:ARG:HG3	40:75:120:ARG:NH1	2.20	0.57
36:78:100:LEU:HD23	36:78:112:LEU:HD11	1.86	0.57
19:AA:63:THR:OG1	19:AA:65:ASN:O	2.15	0.57
40:B8:54:ARG:HA	40:B8:59:THR:HB	1.86	0.57
47:E5:36:ILE:HD11	47:E5:39:ARG:HG2	1.85	0.57
1:13:1440:C:H2'	1:13:1441:G:O4'	2.04	0.57
1:13:600:C:H2'	1:13:601:C:C6	2.40	0.57
1:13:614:A:H2'	1:13:615:C:C6	2.40	0.57
26:14:2138:C:O2'	26:14:2154:G:N2	2.36	0.57
26:1H:1899:G:N2	26:1H:1902:C:H41	1.97	0.57
29:29:65:GLY:C	29:29:68:ALA:HB2	2.26	0.57
34:58:96:GLU:O	34:58:98:VAL:HG12	2.04	0.57
39:65:106:ARG:NH1	39:65:107:GLU:OE1	2.37	0.57
39:65:41:ASP:HB2	39:65:48:LEU:HD21	1.87	0.57
16:7A:19:ILE:HB	16:7A:36:ILE:O	2.04	0.57
41:85:92:ARG:CD	41:85:94:ASN:HB3	2.31	0.57
38:98:33:ARG:HG3	38:98:115:GLU:HB3	1.86	0.57
47:E5:53:MET:HG3	47:E5:59:LEU:HD23	1.87	0.57
26:14:2232:U:OP2	48:F5:40:ARG:NH2	2.37	0.57
51:I5:42:PHE:O	51:I5:43:TYR:HB3	2.04	0.57
26:14:2324:C:H5''	26:14:2325:G:H5'	1.86	0.57
26:14:2319:G:N2	26:14:2334:G:OP1	2.33	0.57
26:14:2427:C:H5''	26:14:2428:G:OP1	2.05	0.57
26:14:38:A:H2'	26:14:39:C:C6	2.40	0.57
26:14:1138:G:H21	34:15:106:MET:CE	2.18	0.57
1:1G:1499:A:H1'	1:1G:1520:G:H5'	1.87	0.57
26:1H:1107:G:H2'	26:1H:1108:U:C6	2.39	0.57
26:1H:2125:G:H21	26:1H:2173:A:H62	1.52	0.57
10:1I:89:ASP:N	10:1I:89:ASP:OD1	2.37	0.57
3:22:150:LYS:HG3	3:22:169:ALA:HB2	1.87	0.57
23:2L:19:G:OP2	23:2L:19:G:H3'	2.04	0.57
12:3I:38:THR:HG22	12:3I:39:VAL:HG23	1.87	0.57
38:55:84:ALA:N	38:55:85:PRO:HD2	2.20	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:69:26:ALA:O	33:69:31:LEU:HB2	2.05	0.57
43:E8:19:LEU:HB3	52:N8:25:LEU:HD12	1.85	0.57
54:Q8:13:ARG:O	54:Q8:23:VAL:HG23	2.05	0.57
26:14:2126:A:H2	26:14:2162:G:H22	1.52	0.56
26:14:2165:G:OP2	26:14:2166:G:N2	2.38	0.56
26:14:363:G:H2'	26:14:363(A):A:C8	2.39	0.56
26:1H:2475:C:H4'	26:1H:2476:A:OP1	2.05	0.56
26:1H:665:C:H2'	26:1H:666:G:C8	2.40	0.56
23:2L:2:C:H2'	23:2L:3:C:H6	1.70	0.56
30:31:32:LEU:HD21	30:31:105:VAL:HG13	1.86	0.56
33:61:83:ALA:HB2	33:61:144:VAL:HG23	1.87	0.56
16:7A:72:ARG:HD2	16:7A:73:LEU:HD23	1.87	0.56
49:K8:15:LYS:HE3	49:K8:15:LYS:N	2.20	0.56
2:12:70:PHE:HB2	2:12:92:TYR:CB	2.35	0.56
1:13:186:C:H2'	1:13:186(A):C:C6	2.40	0.56
1:13:243:A:H4'	1:13:244:U:H5''	1.87	0.56
26:14:2314:C:H5'	31:49:38:VAL:HG11	1.85	0.56
26:14:443:A:H5''	26:14:444:C:OP1	2.05	0.56
27:16:21:G:H1	27:16:62:C:N4	1.99	0.56
1:1G:1043:C:H2'	1:1G:1044:A:C8	2.40	0.56
1:1G:1321:C:H3'	1:1G:1322:C:H5''	1.87	0.56
1:1G:345:C:O2'	1:1G:346:G:O5'	2.19	0.56
26:1H:2159:G:H2'	26:1H:2160:G:O4'	2.05	0.56
26:1H:2408:U:H2'	26:1H:2409:G:C8	2.39	0.56
26:1H:503:A:H4'	26:1H:504:U:H5'	1.87	0.56
26:14:910:A:N7	37:45:13:GLN:HG3	2.20	0.56
25:4L:35:A:H2'	25:4L:36:G:C8	2.40	0.56
18:9I:31:LEU:HD12	18:9I:65:ILE:HB	1.86	0.56
31:41:67:LYS:HE2	51:M8:6:HIS:CE1	2.40	0.56
2:12:9:GLU:HB2	2:12:217:ARG:HH12	1.69	0.56
1:13:1004:A:P	1:13:1025:U:H3	2.29	0.56
1:13:559:A:OP1	5:4E:126:ARG:NH2	2.38	0.56
26:14:1581:G:H2'	26:14:1582:C:O4'	2.05	0.56
26:14:221:A:H61	26:14:265:A:H8	1.53	0.56
28:19:228:PRO:HD3	28:19:234:GLY:O	2.05	0.56
1:1G:1028(A):C:H2'	1:1G:1028(B):C:H5''	1.87	0.56
1:1G:1086:U:H3	1:1G:1099:G:H22	1.53	0.56
26:1H:1332:G:H21	26:1H:1610:A:H8	1.54	0.56
26:1H:2306:C:H3'	26:1H:2307:G:H5'	1.87	0.56
4:3E:152:SER:HB3	4:3E:155:LEU:HD12	1.87	0.56
1:13:537:G:H5''	12:3I:113:ARG:NH1	2.21	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2873:A:H8	38:55:6:SER:N	2.03	0.56
7:62:15:ASP:O	7:62:19:GLY:HA2	2.05	0.56
9:8E:48:GLU:N	9:8E:49:PRO:HD2	2.19	0.56
38:98:10:LEU:O	38:98:12:ARG:HG2	2.03	0.56
46:D5:126:VAL:HA	46:D5:163:LEU:HA	1.86	0.56
51:I5:13:ARG:HA	51:I5:22:ILE:HB	1.87	0.56
1:13:1122:U:O4	1:13:1123:A:N6	2.38	0.56
26:14:1331:A:O2'	26:14:1332:G:H8	1.87	0.56
10:1A:48:THR:HG1	10:1A:62:HIS:HD1	1.33	0.56
1:1G:957:U:H1'	1:1G:960:U:C5	2.40	0.56
26:1H:2793:G:N2	26:1H:2804:C:O2	2.38	0.56
26:1H:330:A:H2	26:1H:1210:A:HO2'	1.53	0.56
26:1H:507:A:H5''	26:1H:508:G:H3'	1.85	0.56
12:3A:47:LYS:O	12:3A:49:ASN:N	2.37	0.56
36:78:59:LEU:HD21	54:Q8:10:ALA:HB2	1.88	0.56
17:8A:57:VAL:HG12	17:8A:76:LEU:HA	1.88	0.56
18:9I:38:GLU:HA	18:9I:41:LYS:HE3	1.85	0.56
51:M8:16:CYS:SG	51:M8:36:CYS:N	2.78	0.56
1:13:76:G:H2'	1:13:77:C:H5'	1.86	0.56
26:14:127:A:H5''	26:14:128:C:C6	2.40	0.56
26:14:1786:A:C2	26:14:2606:C:H1'	2.40	0.56
1:1G:1497:G:C2'	1:1G:1498:U:H5'	2.36	0.56
1:1G:1524:C:H2'	1:1G:1525:G:C8	2.40	0.56
1:1G:632:A:H4'	1:1G:633:G:O5'	2.05	0.56
1:1G:998:G:H22	1:1G:1043:C:N4	2.01	0.56
26:1H:1418:G:OP1	26:1H:1588:C:O2'	2.22	0.56
26:1H:780:G:H21	26:1H:783:A:N6	2.00	0.56
26:1H:782:A:H5'	26:1H:783:A:C2	2.40	0.56
3:2E:188:LEU:HD13	3:2E:195:VAL:HG11	1.87	0.56
23:2K:87:A:H2'	23:2K:87:A:N3	2.21	0.56
9:82:97:LYS:HD3	9:82:102:LEU:HB2	1.85	0.56
18:9I:53:ARG:HA	18:9I:56:THR:HG23	1.88	0.56
52:J5:46:CYS:SG	52:J5:48:GLU:HG2	2.45	0.56
26:1H:1901:A:OP2	28:11:255:LYS:HE2	2.06	0.56
1:13:173:U:H5''	1:13:197:A:O4'	2.05	0.56
26:14:1200:C:H5'	57:14:3972:HOH:O	2.04	0.56
26:14:2414:G:H21	36:35:67:MET:HE1	1.69	0.56
26:14:357:A:H8	26:14:357:A:O5'	1.89	0.56
26:14:859:G:O2'	26:14:916:G:O6	2.17	0.56
27:16:15:A:H5'	27:16:16:G:H8	1.70	0.56
1:1G:401:C:H2'	1:1G:402:G:C8	2.40	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:843:U:H3'	1:1G:848:C:O4'	2.06	0.56
26:1H:1055:G:HO2'	26:1H:1086:A:N6	2.04	0.56
26:1H:120:U:OP2	57:1H:3889:HOH:O	2.18	0.56
26:1H:862:G:OP2	57:1H:3781:HOH:O	2.18	0.56
30:39:123:LEU:O	30:39:193:VAL:HA	2.05	0.56
24:3K:17:U:O2'	24:3K:68:A:N1	2.34	0.56
7:6E:5:ARG:HG2	7:6E:7:ALA:H	1.71	0.56
39:A8:87:PHE:CE2	39:A8:102:ALA:HB2	2.40	0.56
26:1H:1266:G:O5'	43:E8:15:ARG:NH2	2.39	0.56
44:F8:12:VAL:HG13	44:F8:27:THR:O	2.06	0.56
1:13:1360:A:H2'	1:13:1361:G:O4'	2.05	0.56
26:14:1198:U:H2'	26:14:1199:U:H6	1.71	0.56
26:14:2006:C:O2'	26:14:2823:A:N3	2.37	0.56
26:14:2128:C:H42	26:14:2160:G:N2	2.04	0.56
26:14:221:A:N6	26:14:265:A:C8	2.73	0.56
26:14:698:C:O2'	26:14:734:A:N6	2.39	0.56
28:19:177:LEU:HD11	28:19:183:ARG:HD2	1.88	0.56
26:1H:1533:C:H2'	26:1H:1534:G:C2	2.41	0.56
26:1H:2091:U:O4	57:1H:4142:HOH:O	2.14	0.56
26:1H:248:G:H5'	26:1H:250:G:N7	2.19	0.56
26:1H:357:A:H2'	26:1H:358:U:H6	1.70	0.56
26:1H:818:G:H5'	26:1H:839:U:OP1	2.05	0.56
26:14:1665:A:H1'	35:25:1:MET:HG2	1.88	0.56
36:35:8:PRO:HB2	36:35:12:ALA:HB3	1.87	0.56
1:13:750:G:N3	15:6I:23:GLY:HA3	2.21	0.56
9:8E:112:LYS:HD2	9:8E:113:LYS:N	2.20	0.56
48:F5:86:SER:N	48:F5:87:PRO:HD2	2.20	0.56
37:88:85:LYS:HG3	47:I8:7:LEU:HD13	1.88	0.56
52:J5:16:ARG:HG3	52:J5:17:ASP:N	2.19	0.56
1:13:639:G:H2'	1:13:640:A:H8	1.70	0.56
26:14:1556:C:H2'	26:14:1557:C:C6	2.37	0.56
26:14:1688:U:O2	26:14:1700:A:H5'	2.05	0.56
26:14:642:G:N2	26:14:645:C:OP2	2.37	0.56
1:1G:1238:A:H62	1:1G:1301:U:H3	1.53	0.56
1:1G:1312:G:H2'	1:1G:1313:U:C6	2.40	0.56
1:1G:539:A:H2'	1:1G:540:G:H8	1.69	0.56
26:1H:1063:G:N1	26:1H:1076:C:O2'	2.38	0.56
26:1H:607:U:N3	26:1H:621:A:H2	1.97	0.56
22:1L:63:G:N2	22:1L:73:C:O2	2.29	0.56
29:21:78:LEU:O	29:21:79:ARG:HB2	2.06	0.56
1:1G:1216:G:H5''	14:5A:5:ALA:HB3	1.87	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:78:83:VAL:HG12	36:78:112:LEU:HD21	1.86	0.56
37:88:20:ALA:HB1	37:88:99:PRO:HB2	1.88	0.56
40:B8:58:ASN:C	40:B8:58:ASN:HD22	2.08	0.56
20:BI:53:LEU:O	20:BI:57:ARG:HD3	2.05	0.56
43:E8:18:ARG:HH11	43:E8:76:VAL:HG22	1.70	0.56
52:J5:6:VAL:HG22	52:J5:7:PRO:HD2	1.86	0.56
54:Q8:54:GLU:O	54:Q8:56:GLU:N	2.39	0.56
2:12:7:VAL:HG13	2:12:8:LYS:HG3	1.86	0.56
1:13:1304:G:N2	1:13:1332:A:OP2	2.30	0.56
1:13:1441:G:H21	1:13:1460:A:H62	1.52	0.56
1:13:145:G:H2'	1:13:146:G:H8	1.71	0.56
26:14:1054:A:H2'	26:14:1055:G:H8	1.70	0.56
26:14:2543:G:H2'	26:14:2544:G:C8	2.41	0.56
26:14:2761:G:H1'	32:59:143:GLN:HE22	1.70	0.56
1:1G:1297:C:OP1	13:4A:44:ARG:NH2	2.21	0.56
26:1H:2121:G:H1	26:1H:2177:C:H42	1.53	0.56
26:1H:631:A:OP1	36:78:65:ARG:NH2	2.21	0.56
22:1L:31:C:H2'	22:1L:32:C:H6	1.71	0.56
36:35:101:VAL:HA	36:35:105:LEU:O	2.06	0.56
14:5I:21:TYR:OH	14:5I:23:ARG:NH2	2.39	0.56
33:61:110:ASP:OD1	33:61:111:PRO:HA	2.06	0.56
7:62:69:VAL:HG22	7:62:135:VAL:HG23	1.87	0.56
35:68:7:TYR:HE1	35:68:20:MET:HE3	1.70	0.56
8:72:25:ASP:OD1	8:72:25:ASP:N	2.39	0.56
30:31:33:LEU:HB3	36:78:6:LEU:HD11	1.86	0.56
9:8E:26:VAL:HB	9:8E:33:PHE:HB2	1.88	0.56
48:F5:29:GLY:O	48:F5:30:VAL:HG22	2.06	0.56
26:14:752:A:H3'	53:L5:1:MET:SD	2.46	0.56
54:Q8:45:GLY:HA2	54:Q8:46:ARG:C	2.27	0.56
54:Q8:57:ARG:HD3	54:Q8:57:ARG:N	2.21	0.56
26:14:1430:C:H2'	26:14:1431:U:C6	2.41	0.56
26:14:1677:A:H2'	26:14:1678:G:C8	2.41	0.56
26:14:2156:G:N7	26:14:2157:G:N2	2.54	0.56
26:14:223:A:O2'	26:14:420:C:O2	2.19	0.56
1:1G:1143:G:H2'	1:1G:1144:G:C8	2.41	0.56
1:1G:158:G:H1	1:1G:163:C:H42	1.54	0.56
26:1H:130:C:O3'	26:1H:1349:A:H1'	2.06	0.56
26:1H:523:C:O2	26:1H:553:U:O2'	2.24	0.56
10:1I:22:LYS:NZ	10:1I:90:LEU:HB2	2.21	0.56
29:29:147:PRO:HB2	29:29:149:ARG:HG2	1.88	0.56
30:31:63:LYS:HE3	30:31:67:GLN:HB2	1.88	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2469:A:O2'	37:45:56:ARG:HG2	2.06	0.56
5:4E:72:GLN:HE22	5:4E:144:THR:HG22	1.71	0.56
33:61:117:GLU:N	33:61:117:GLU:OE2	2.38	0.56
36:78:116:GLY:H	36:78:134:ALA:HB2	1.70	0.56
26:1H:825:C:O2	36:78:55:ARG:NH2	2.39	0.56
36:78:82:GLY:HA2	36:78:113:LYS:O	2.06	0.56
17:8I:67:LYS:HA	17:8I:70:ARG:HH12	1.71	0.56
18:9A:37:VAL:HA	18:9A:40:LEU:HB2	1.87	0.56
26:14:1087:G:H2'	26:14:1089:G:H1'	1.88	0.56
26:14:2020:A:O2'	26:14:2021:C:H5'	2.05	0.56
26:14:2209:C:O2	26:14:2216:G:C2	2.59	0.56
26:14:568:U:N3	57:14:3987:HOH:O	2.32	0.56
21:1B:8:THR:HB	21:1B:11:GLY:H	1.70	0.56
1:1G:1522:U:H2'	1:1G:1523:G:H8	1.70	0.56
1:1G:464:G:C6	1:1G:466:C:H5'	2.41	0.56
26:1H:1068:G:H1'	26:1H:1096:A:H1'	1.87	0.56
26:1H:1169:G:H1	26:1H:1180:C:H42	1.53	0.56
26:1H:1503:U:H2'	26:1H:1504:C:C6	2.40	0.56
26:1H:1534:G:H22	26:1H:1538:G:H22	1.53	0.56
26:1H:660:G:H21	36:78:12:ALA:HA	1.71	0.56
1:1G:523:A:H61	12:3A:92:ASP:HB2	1.71	0.56
41:C8:88:ILE:O	41:C8:90:VAL:N	2.39	0.56
1:13:1410:G:H2'	1:13:1411:C:H6	1.72	0.55
1:13:413:G:N2	1:13:428:G:H1'	2.21	0.55
1:1G:1125:U:O4	10:1A:5:ARG:NH2	2.40	0.55
1:1G:1285:A:H4'	1:1G:1286:A:O5'	2.06	0.55
26:1H:2305:A:H1'	31:41:136:ARG:NH1	2.21	0.55
26:1H:994:C:H3'	41:C8:54:LYS:HE3	1.88	0.55
27:1J:48:A:H4'	39:65:95:HIS:CD2	2.33	0.55
22:1L:61:C:H42	22:1L:75:G:H1	1.52	0.55
30:31:67:GLN:HG3	30:31:67:GLN:O	2.05	0.55
4:32:7:PRO:HB2	4:32:10:ARG:HD2	1.88	0.55
4:32:98:GLU:OE2	4:32:103:ASN:ND2	2.34	0.55
39:65:7:TYR:HE1	39:65:11:LYS:HZ3	1.52	0.55
26:1H:2485:G:H5''	37:88:46:GLN:NE2	2.21	0.55
19:AI:40:ILE:HG12	19:AI:41:VAL:N	2.21	0.55
46:D5:52:SER:O	46:D5:52:SER:OG	2.20	0.55
42:D8:31:ALA:O	42:D8:60:GLU:HG2	2.05	0.55
47:E5:27:GLU:HG3	47:E5:68:GLU:HA	1.88	0.55
1:13:746:A:H2'	1:13:747:C:C6	2.41	0.55
1:13:748:C:H4'	1:13:749:C:O5'	2.06	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:953:G:H2'	1:13:954:G:O4'	2.06	0.55
22:1L:87:A:O3'	26:14:2506:U:O2'	2.19	0.55
26:14:333:G:H5''	26:14:334:C:OP2	2.06	0.55
28:19:75:ILE:HD12	28:19:75:ILE:H	1.70	0.55
1:1G:1347:G:N7	9:82:107:ARG:NH1	2.55	0.55
26:1H:1021:A:OP2	34:58:65:LYS:NZ	2.27	0.55
26:1H:1026:U:O2'	26:1H:1027:A:O5'	2.20	0.55
26:1H:873:G:H1'	37:88:29:PHE:HE2	1.71	0.55
27:1J:63:G:H2'	27:1J:64:C:C6	2.42	0.55
3:22:70:VAL:HG12	3:22:72:LYS:N	2.21	0.55
30:31:65:TRP:CZ3	30:31:72:ARG:HB3	2.41	0.55
30:39:57:VAL:HG11	30:39:79:GLY:HA3	1.88	0.55
5:4E:84:PHE:HB3	5:4E:134:ALA:HB2	1.88	0.55
33:69:4:ILE:HG21	33:69:47:LEU:HD13	1.88	0.55
17:8A:48:GLU:HB2	17:8A:50:LYS:HB2	1.88	0.55
53:P8:30:VAL:O	53:P8:34:ARG:HG3	2.06	0.55
1:13:673:G:H2'	1:13:674:G:C8	2.41	0.55
26:14:1060:U:H4'	26:14:1061:U:C5'	2.35	0.55
26:14:1141:U:H2'	34:15:63:THR:HG21	1.88	0.55
26:14:1174:A:N6	26:14:1177:A:H4'	2.22	0.55
27:16:73:A:H5'	27:16:74:U:OP2	2.06	0.55
1:1G:1126:U:N3	1:1G:1281:U:O4'	2.39	0.55
1:1G:158:G:N2	1:1G:163:C:N3	2.50	0.55
26:1H:1510:A:OP1	26:1H:1511:A:H5'	2.06	0.55
26:1H:1516:U:H2'	26:1H:1517:G:H8	1.71	0.55
26:1H:1533:C:H3'	26:1H:1534:G:C5'	2.36	0.55
26:1H:2611:U:H6	26:1H:2611:U:H5'	1.70	0.55
11:2A:98:LEU:HA	11:2A:101:SER:HB3	1.88	0.55
4:32:31:CYS:C	4:32:33:MET:N	2.59	0.55
12:3I:91:LYS:O	12:3I:91:LYS:HG3	2.06	0.55
37:45:35:VAL:HG12	37:45:36:ALA:H	1.70	0.55
37:45:38:GLU:HG3	37:45:127:ILE:HG22	1.89	0.55
37:45:60:ARG:HH11	37:45:61:GLY:HA2	1.71	0.55
5:4E:126:ARG:HH11	5:4E:126:ARG:HG3	1.70	0.55
6:52:86:ARG:O	6:52:87:ARG:HG2	2.07	0.55
38:55:45:ARG:HA	38:55:95:THR:HG21	1.88	0.55
7:62:20:ASP:HB3	7:62:23:VAL:HG23	1.88	0.55
1:1G:229:U:O2'	16:7A:23:ASP:OD2	2.20	0.55
16:7I:71:ARG:O	16:7I:75:ARG:N	2.39	0.55
18:9A:22:VAL:C	18:9A:24:ALA:H	2.08	0.55
45:G8:55:TYR:N	45:G8:56:PRO:HD3	2.21	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:1H:4089:HOH:O	48:J8:65:SER:HB3	2.05	0.55
54:M5:37:SER:HB2	54:M5:39:LYS:O	2.06	0.55
1:13:328:C:H4'	1:13:329:A:C5'	2.36	0.55
1:13:674:G:H2'	1:13:675:A:H8	1.71	0.55
1:13:916:G:H2'	1:13:917:G:H8	1.71	0.55
26:14:1885:A:H5'	26:14:1886:C:OP2	2.05	0.55
26:14:907:U:O2'	37:45:101:ARG:NH2	2.36	0.55
34:15:61:ARG:HA	34:15:61:ARG:CZ	2.36	0.55
26:1H:1339:G:N2	26:1H:1603:A:H1'	2.22	0.55
26:1H:1688:U:H2'	26:1H:1698:A:N6	2.22	0.55
26:1H:2103:C:H2'	26:1H:2104:G:C8	2.42	0.55
26:1H:573:G:O2'	26:1H:574:C:H3'	2.06	0.55
11:2A:48:ILE:HD11	11:2A:64:ALA:HA	1.89	0.55
23:2K:73:C:H2'	23:2K:74:C:H6	1.71	0.55
30:31:158:THR:HG23	30:31:160:ASN:H	1.70	0.55
4:32:13:ARG:C	4:32:15:GLU:H	2.09	0.55
12:3A:58:VAL:O	12:3A:65:GLU:HA	2.05	0.55
31:49:32:PRO:HB2	31:49:172:LEU:HD13	1.88	0.55
32:59:137:ASP:OD1	32:59:138:LYS:N	2.38	0.55
14:5A:29:ARG:HB3	14:5A:31:ARG:H	1.70	0.55
36:78:18:ARG:O	36:78:19:VAL:HG13	2.06	0.55
16:7I:9:PHE:HE2	16:7I:18:ARG:HH21	1.54	0.55
41:85:75:ASN:OD1	41:85:78:THR:OG1	2.18	0.55
26:1H:958:U:OP2	37:88:14:ARG:NH1	2.38	0.55
40:B8:84:GLN:HG2	40:B8:85:LYS:HG2	1.87	0.55
45:C5:68:HIS:HB2	45:C5:71:LYS:HZ3	1.71	0.55
37:45:134:ARG:HH22	46:D5:122:ARG:NH1	2.04	0.55
1:13:598:U:H4'	8:7E:94:TYR:CD2	2.42	0.55
26:14:1292:U:H2'	26:14:1293:C:C6	2.40	0.55
26:14:479:A:N3	26:14:481:G:H5''	2.21	0.55
26:14:540:G:H2'	26:14:541:C:C6	2.41	0.55
1:1G:1347:G:N2	1:1G:1373:G:H2'	2.22	0.55
26:1H:2232:U:P	48:J8:40:ARG:HH12	2.29	0.55
26:1H:760:G:C2'	26:1H:761:A:H5'	2.36	0.55
27:1J:88:C:H3'	27:1J:89:G:N7	2.21	0.55
23:2K:71:U:H5''	23:2K:72:C:C5	2.41	0.55
4:32:111:ALA:HB3	4:32:117:ALA:HB2	1.87	0.55
4:32:31:CYS:C	4:32:33:MET:H	2.09	0.55
36:35:121:LYS:HG2	36:35:122:PRO:HD2	1.88	0.55
30:39:108:LYS:O	30:39:112:MET:HB2	2.05	0.55
4:3E:102:ASP:OD1	4:3E:103:ASN:N	2.39	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:3E:107:ARG:HH12	4:3E:194:LEU:HD22	1.71	0.55
24:3L:42:U:H2'	24:3L:43:A:C8	2.42	0.55
32:59:136:ILE:HD12	32:59:137:ASP:HB2	1.87	0.55
1:1G:1375:A:H4'	7:62:29:LYS:HE3	1.89	0.55
41:C8:106:PHE:HA	41:C8:109:LEU:HD12	1.88	0.55
52:N8:40:LYS:NZ	52:N8:46:CYS:HB3	2.21	0.55
54:Q8:59:LYS:CE	54:Q8:59:LYS:H	2.19	0.55
1:13:1028:C:H42	1:13:1033:G:H22	1.53	0.55
1:13:626:U:C2	1:13:627:G:C8	2.95	0.55
26:14:1747:G:H2'	26:14:1748:G:H8	1.71	0.55
1:1G:1152:A:OP1	10:1A:68:HIS:NE2	2.39	0.55
1:1G:683:G:H2'	1:1G:684:A:C8	2.42	0.55
26:1H:1021:A:H3'	26:1H:1022:G:H5''	1.87	0.55
26:1H:2315:G:OP1	31:41:36:LYS:NZ	2.34	0.55
26:1H:71:A:H4'	26:1H:72:U:H5''	1.87	0.55
23:2L:29:U:H2'	23:2L:30:A:C8	2.42	0.55
24:3L:37:G:N2	25:4L:44:U:H3	2.05	0.55
14:5A:15:LYS:HG2	14:5A:16:PHE:CE2	2.42	0.55
39:65:28:VAL:HG11	39:65:98:VAL:HG12	1.87	0.55
39:65:7:TYR:CZ	39:65:91:PRO:HG3	2.42	0.55
38:98:9:LYS:HA	38:98:17:ARG:NE	2.22	0.55
18:9A:86:VAL:HG12	18:9A:87:ARG:NH1	2.22	0.55
36:35:63:PRO:HB3	54:M5:30:ARG:CZ	2.37	0.55
31:41:112:PRO:HB3	51:M8:37:SER:H	1.71	0.55
1:13:1023:G:H3'	1:13:1024:G:H5''	1.89	0.55
1:13:221:C:H2'	1:13:222:U:H6	1.72	0.55
26:14:113:G:H8	26:14:113:G:O5'	1.89	0.55
26:14:1771:C:H1'	26:14:1786:A:C8	2.41	0.55
26:14:2138:C:O2	26:14:2154:G:N1	2.40	0.55
26:14:830:G:H4'	26:14:831:G:OP2	2.06	0.55
1:1G:1453:G:H22	20:BA:54:LYS:HZ2	1.53	0.55
1:1G:299:G:H2'	1:1G:300:A:C8	2.42	0.55
26:1H:1494:A:H2'	26:1H:1495:A:C8	2.41	0.55
26:1H:2111:C:H5	26:1H:2147:G:C6	2.25	0.55
26:1H:2262:U:H2'	26:1H:2263:C:H6	1.72	0.55
26:1H:2882:A:OP1	38:98:96:ARG:HD3	2.06	0.55
29:21:60:ASN:OD1	29:21:62:PRO:HD2	2.06	0.55
26:14:2052:G:O4'	29:29:142:GLY:HA3	2.06	0.55
5:4E:81:GLU:HG2	5:4E:90:VAL:HG23	1.89	0.55
1:13:1316:G:H4'	14:5I:18:VAL:HG11	1.89	0.55
33:69:82:ARG:O	33:69:89:TYR:HD2	1.89	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:82:99:LEU:HB3	9:82:101:PHE:CE1	2.41	0.55
41:85:66:ASN:HB2	41:85:76:TYR:HB2	1.87	0.55
17:8A:63:ARG:HG2	17:8A:64:PRO:HD2	1.89	0.55
38:98:29:LEU:HB3	38:98:75:LEU:HD21	1.89	0.55
40:B8:50:ILE:CD1	40:B8:102:ILE:HD11	2.35	0.55
28:11:10:THR:OG1	28:11:13:ARG:HB2	2.06	0.55
1:13:498:A:H4'	1:13:500:G:OP1	2.06	0.55
26:14:1047:G:H21	26:14:1111:A:H62	1.55	0.55
26:14:1199:U:H2'	26:14:1200:C:C6	2.41	0.55
26:14:1366:A:H2'	26:14:1367:A:O4'	2.06	0.55
26:14:787:U:P	57:14:3662:HOH:O	2.63	0.55
28:19:68:LYS:HB3	28:19:70:TRP:CZ3	2.42	0.55
2:1E:76:GLN:O	2:1E:208:ILE:HG12	2.07	0.55
1:1G:448:A:OP2	1:1G:485:G:N2	2.37	0.55
26:1H:1509:C:H3'	26:1H:1510:A:H5''	1.89	0.55
26:1H:2116:G:OP2	26:1H:2166:G:O2'	2.24	0.55
26:1H:2376:A:H2'	26:1H:2377:A:O4'	2.07	0.55
26:1H:2553:G:H2'	26:1H:2554:U:H4'	1.89	0.55
11:2A:16:SER:OG	11:2A:79:SER:OG	2.23	0.55
26:14:2443:C:OP1	30:39:68:LYS:HG2	2.06	0.55
31:49:109:VAL:HG11	31:49:142:PRO:HG3	1.87	0.55
31:49:22:ARG:HH12	31:49:175:LEU:HD21	1.71	0.55
1:13:376:G:O3'	16:7I:5:ARG:NH1	2.40	0.55
41:85:92:ARG:NH2	42:95:11:GLN:H	2.05	0.55
26:14:2387:U:H1'	47:E5:41:ARG:NE	2.22	0.55
45:G8:85:VAL:HG22	45:G8:98:VAL:HB	1.88	0.55
37:88:85:LYS:CG	47:I8:7:LEU:HD13	2.37	0.55
26:14:1997:G:H5''	57:14:3984:HOH:O	2.07	0.55
26:14:2774:C:H2'	26:14:2775:A:O4'	2.06	0.55
34:15:35:ARG:HB2	34:15:42:TRP:HZ3	1.71	0.55
2:1E:174:VAL:HG11	2:1E:196:LEU:HD13	1.87	0.55
1:1G:1281:U:OP2	1:1G:1282:C:N4	2.35	0.55
1:1G:737:A:H2'	1:1G:738:C:C6	2.41	0.55
26:1H:1231:G:H2'	26:1H:1232:G:H8	1.72	0.55
26:1H:2052:G:H4'	29:21:143:ASN:O	2.06	0.55
26:1H:2481:G:O2'	26:1H:2482:G:O5'	2.25	0.55
29:21:24:THR:HG21	29:21:188:VAL:CG2	2.37	0.55
35:25:43:VAL:HG23	35:25:56:ASP:O	2.07	0.55
36:35:125:VAL:O	36:35:144:GLU:HB3	2.07	0.55
13:4I:88:ARG:HD2	13:4I:98:VAL:HG12	1.89	0.55
7:62:95:ARG:HH21	7:62:99:LEU:HD11	1.72	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1179:A:OP2	9:82:93:ARG:NH2	2.40	0.55
41:85:16:LYS:O	41:85:20:LEU:HD13	2.06	0.55
41:85:92:ARG:NH2	42:95:10:LYS:HA	2.21	0.55
17:8I:29:HIS:CD2	17:8I:30:PRO:HD2	2.42	0.55
19:AI:6:LYS:HE3	51:M8:62:ARG:HD2	1.89	0.55
54:Q8:47:LYS:HD3	54:Q8:47:LYS:C	2.28	0.55
2:12:16:HIS:HD2	2:12:210:SER:HA	1.71	0.55
1:13:1228:C:OP1	13:4I:108:ARG:NH2	2.40	0.55
26:14:2873:A:H8	38:55:5:LYS:HA	1.72	0.55
2:1E:212:GLN:O	2:1E:216:SER:OG	2.15	0.55
1:1G:411:A:H62	1:1G:413:G:H21	1.54	0.55
26:1H:2791:C:N4	26:1H:2805:G:H1	2.05	0.55
23:2L:18:OMG:P	23:2L:71:U:H3	2.30	0.55
26:14:390:A:C6	36:35:71:VAL:HG21	2.42	0.55
34:58:47:ALA:HB2	34:58:112:LEU:HD11	1.89	0.55
32:59:66:GLY:O	32:59:70:THR:OG1	2.20	0.55
39:65:42:ASP:O	39:65:43:GLU:HB2	2.06	0.55
40:75:77:PRO:HG2	40:75:80:SER:HB2	1.88	0.55
41:85:92:ARG:C	41:85:94:ASN:H	2.10	0.55
9:8E:10:ARG:HG3	9:8E:75:ASP:HB2	1.88	0.55
42:95:21:ARG:HE	42:95:91:TYR:HB3	1.71	0.55
45:C5:47:LYS:HG3	45:C5:60:PHE:CE2	2.41	0.55
45:G8:82:PRO:HG3	45:G8:97:ARG:HG3	1.89	0.55
46:H8:116:VAL:HG23	46:H8:146:ILE:HG12	1.89	0.55
28:11:17:THR:HB	28:11:205:VAL:H	1.72	0.54
2:12:109:SER:HA	2:12:112:VAL:HG23	1.89	0.54
1:13:179:A:H2'	1:13:180:U:H6	1.71	0.54
26:14:1533:C:N3	26:14:1534:G:O2'	2.33	0.54
26:14:2250:G:C6	37:45:83:MET:HB3	2.42	0.54
26:14:2294:C:P	39:65:89:ARG:HH22	2.29	0.54
26:14:2311:A:H62	31:49:44:GLY:HA3	1.70	0.54
26:14:363(F):A:OP2	26:14:363(F):A:H8	1.89	0.54
26:14:35:G:H2'	26:14:36:G:O4'	2.07	0.54
26:14:483:A:H4'	45:C5:49:VAL:HA	1.89	0.54
1:1G:186(A):C:H2'	1:1G:186(B):C:H6	1.72	0.54
1:1G:116:A:H61	1:1G:313:A:H1'	1.72	0.54
26:1H:973:A:OP2	57:1H:3717:HOH:O	2.17	0.54
29:21:70:ALA:HB1	29:21:73:GLU:HB2	1.88	0.54
29:29:25:VAL:HG12	29:29:26:ILE:H	1.72	0.54
1:13:1523:G:OP1	11:2I:123:LYS:HE3	2.06	0.54
4:32:36:ARG:HB3	4:32:38:TYR:CE1	2.41	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:39:6:VAL:HB	30:39:124:LEU:HA	1.89	0.54
31:41:131:TYR:O	31:41:159:VAL:HG22	2.07	0.54
13:4I:23:TYR:HB3	13:4I:67:GLU:HB2	1.88	0.54
13:4I:75:ALA:HA	13:4I:78:ILE:HG13	1.90	0.54
32:59:119:GLU:N	32:59:119:GLU:OE2	2.34	0.54
6:5E:9:VAL:HB	6:5E:87:ARG:HB2	1.88	0.54
7:62:97:GLN:HG3	7:62:98:SER:N	2.21	0.54
26:1H:2849:U:OP2	40:B8:95:ARG:NH1	2.40	0.54
20:BI:97:ALA:O	20:BI:99:LEU:N	2.39	0.54
47:E5:21:LEU:HD21	47:E5:41:ARG:HH12	1.71	0.54
26:1H:71:A:C2	44:F8:31:HIS:HE1	2.25	0.54
51:M8:13:ARG:HA	51:M8:24:THR:HG21	1.90	0.54
1:13:138:G:H1	1:13:225:C:H42	1.53	0.54
26:14:551:G:H1'	26:14:1220:A:C6	2.42	0.54
26:14:975:G:H1'	26:14:990:A:C2	2.42	0.54
26:14:997:G:O2'	26:14:998:C:H5'	2.07	0.54
10:1A:9:ARG:HB2	10:1A:95:GLU:HB3	1.88	0.54
2:1E:47:THR:HG23	2:1E:202:PRO:HG2	1.88	0.54
1:1G:1075:C:H4'	2:12:175:ARG:HH22	1.71	0.54
26:1H:1437:C:C2	26:1H:1438:U:C5	2.95	0.54
26:1H:2882:A:H5'	38:98:96:ARG:HG3	1.87	0.54
26:1H:654(E):C:N3	26:1H:654(P):G:N2	2.48	0.54
26:1H:924:C:H2'	26:1H:925:C:C6	2.43	0.54
27:1J:116:G:C5'	39:65:55:ALA:HB2	2.36	0.54
31:41:112:PRO:HB2	51:M8:36:CYS:HA	1.88	0.54
31:49:125:PHE:HB3	31:49:166:ASP:HB2	1.89	0.54
5:4E:20:GLN:HG2	5:4E:21:ALA:H	1.72	0.54
8:7E:104:ARG:HB3	8:7E:107:LEU:HB2	1.89	0.54
41:85:92:ARG:CZ	42:95:11:GLN:H	2.21	0.54
1:13:1368:G:H5''	9:8E:112:LYS:HB3	1.88	0.54
18:9I:26:LEU:HB3	18:9I:42:ARG:HH22	1.72	0.54
46:D5:53:ILE:HG22	46:D5:71:VAL:HG13	1.87	0.54
46:H8:134:PRO:HG3	46:H8:161:VAL:HG11	1.87	0.54
1:13:353:A:H5'	1:13:353:A:H8	1.73	0.54
1:13:992:U:H4'	1:13:993:G:O5'	2.06	0.54
26:14:2472:G:H2'	26:14:2475:C:N4	2.21	0.54
26:14:2716:U:O2'	26:14:2717:G:H5'	2.07	0.54
26:14:271:G:H2'	26:14:272:G:H8	1.72	0.54
1:1G:1142:G:H2'	1:1G:1143:G:O4'	2.07	0.54
1:1G:735:C:H2'	1:1G:736:C:H6	1.71	0.54
1:1G:974:A:P	14:5A:41:ARG:HH22	2.30	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:127:A:H5''	26:1H:128:C:C6	2.42	0.54
26:1H:935:C:H2'	26:1H:936:C:H6	1.72	0.54
22:1L:87:A:HO3'	26:14:2506:U:HO2'	1.52	0.54
29:29:89:ASP:CG	29:29:90:THR:H	2.11	0.54
23:2K:71:U:H5''	23:2K:72:C:H5	1.72	0.54
5:4E:70:PRO:O	5:4E:77:PRO:HD3	2.08	0.54
15:6I:82:ILE:O	15:6I:86:GLY:N	2.40	0.54
18:9I:66:LEU:HD11	18:9I:70:ILE:HD11	1.89	0.54
2:12:92:TYR:HE2	2:12:94:ASN:HB2	1.72	0.54
1:13:1305:G:C8	1:13:1305:G:OP2	2.57	0.54
1:13:143:A:H2	1:13:220:G:H1	1.56	0.54
1:13:434:U:H2'	1:13:435:C:C6	2.42	0.54
26:14:1048:A:H2	26:14:1112:G:H21	1.55	0.54
26:14:1149:G:H2'	26:14:1150:C:C6	2.42	0.54
26:14:1412:A:H2'	26:14:1413:G:C8	2.42	0.54
26:14:857:C:H2'	26:14:858:U:C6	2.43	0.54
1:1G:114:U:H2'	1:1G:115:G:C8	2.43	0.54
1:1G:1352:C:H2'	1:1G:1353:G:C8	2.42	0.54
1:1G:977:A:H2'	1:1G:978:A:H5'	1.90	0.54
26:1H:1060:U:C4	26:1H:1062:G:H4'	2.41	0.54
26:1H:2176:A:H2'	26:1H:2177:C:C6	2.41	0.54
26:1H:1889:A:N1	26:1H:2234:G:H1'	2.21	0.54
26:1H:722:A:H2'	26:1H:723:G:C8	2.42	0.54
26:1H:935:C:H2'	26:1H:936:C:C6	2.42	0.54
26:14:1665:A:H4'	35:25:67:LYS:HB2	1.90	0.54
32:51:169:VAL:O	32:51:170:ARG:NE	2.39	0.54
34:58:70:LYS:HE3	34:58:72:TYR:CE1	2.42	0.54
26:14:2378:A:O2'	39:65:21:THR:HG21	2.08	0.54
15:6A:87:ILE:HG22	15:6A:88:ARG:N	2.21	0.54
9:8E:125:TYR:HD1	9:8E:126:SER:N	2.05	0.54
9:8E:50:LEU:HD23	9:8E:85:LEU:HD11	1.88	0.54
43:A5:6:ILE:HG22	43:A5:8:ARG:HG3	1.89	0.54
26:14:2015:A:OP1	43:A5:92:ARG:NH2	2.40	0.54
45:C5:86:ARG:HG3	45:C5:87:LYS:N	2.21	0.54
26:1H:58:G:H5'	44:F8:74:PRO:HB3	1.88	0.54
1:13:1427:U:H2'	1:13:1428:A:C8	2.43	0.54
1:1G:171:A:H2'	1:1G:172:A:C8	2.41	0.54
1:1G:501:C:H2'	1:1G:502:G:C8	2.42	0.54
29:29:76:ARG:HG2	29:29:195:LEU:HD22	1.88	0.54
36:35:39:LYS:HG3	36:35:45:LEU:HD22	1.90	0.54
30:39:15:SER:OG	30:39:16:GLY:N	2.39	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:3E:43:HIS:O	4:3E:46:LYS:HG3	2.08	0.54
32:51:126:PRO:HB2	32:51:130:ARG:HH22	1.72	0.54
7:62:101:LEU:O	7:62:105:VAL:HG23	2.07	0.54
46:H8:105:VAL:O	46:H8:140:ASP:HA	2.08	0.54
1:13:56:U:H2'	1:13:57:G:C8	2.43	0.54
1:13:658:G:H2'	1:13:659:U:H6	1.73	0.54
1:13:859:A:H2'	1:13:860:A:O4'	2.06	0.54
26:14:1061:U:C5'	26:14:1070:A:H1'	2.32	0.54
26:14:1310:G:OP2	53:L5:9:ARG:NE	2.41	0.54
26:14:1429:G:H2'	26:14:1430:C:C6	2.42	0.54
26:14:536:A:OP1	41:85:53:ARG:NH1	2.40	0.54
28:19:208:LYS:HG3	28:19:211:ARG:H	1.73	0.54
26:1H:1259:G:H2'	26:1H:1260:G:C8	2.42	0.54
26:1H:2299:G:H1	26:1H:2318:G:H8	1.55	0.54
31:41:11:TYR:HA	31:41:15:VAL:HB	1.90	0.54
7:62:26:PHE:CD1	7:62:30:ILE:HD11	2.43	0.54
42:95:69:LYS:HG2	42:95:86:GLY:HA3	1.90	0.54
42:95:85:LYS:HD2	42:95:87:HIS:H	1.73	0.54
39:A8:42:ASP:O	39:A8:43:GLU:HB2	2.08	0.54
46:D5:79:ARG:HB3	46:D5:80:ARG:HD2	1.90	0.54
37:88:134:ARG:HE	46:H8:122:ARG:NH2	2.06	0.54
48:J8:58:ILE:HG23	48:J8:87:PRO:HG3	1.88	0.54
49:K8:52:ASP:O	49:K8:56:GLN:HB2	2.08	0.54
26:1H:1805:U:O2	28:11:50:THR:HB	2.07	0.54
1:13:1188:A:N6	57:13:1842:HOH:O	2.41	0.54
1:13:1301:U:O2'	1:13:1302:U:H5'	2.08	0.54
26:14:2138:C:H1'	26:14:2154:G:H1	1.73	0.54
26:14:607:U:H3	26:14:621:A:H2	1.54	0.54
26:14:833:U:H2'	26:14:834:C:C6	2.43	0.54
26:1H:1126:A:H4'	26:1H:1127:A:O5'	2.08	0.54
26:1H:2209:C:O2	26:1H:2216:G:C2	2.60	0.54
22:1L:21:A:H4'	22:1L:22:G:OP1	2.07	0.54
29:29:106:GLY:HA3	29:29:189:PRO:HB2	1.89	0.54
4:3E:141:ARG:HB2	4:3E:141:ARG:NH1	2.22	0.54
24:3L:22:G:H1'	24:3L:70:G:H22	1.71	0.54
13:4I:108:ARG:HH11	13:4I:108:ARG:HA	1.70	0.54
33:61:73:GLU:HG3	33:61:136:VAL:HG23	1.90	0.54
26:1H:2684:U:O2'	35:68:68:GLU:HG3	2.08	0.54
36:78:13:ASN:OD1	36:78:15:ARG:HD3	2.07	0.54
26:1H:1250:G:OP2	36:78:18:ARG:NH1	2.40	0.54
17:8A:45:HIS:HE2	17:8A:47:PRO:HB3	1.71	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:I5:18:CYS:HB2	51:I5:36:CYS:HB3	1.90	0.54
1:13:592:G:H2'	1:13:593:G:H8	1.73	0.54
1:13:664:G:N2	1:13:741:G:H1	2.04	0.54
26:14:1054:A:H2'	26:14:1055:G:C8	2.43	0.54
26:14:1327:C:O3'	38:55:105:ARG:NH2	2.41	0.54
26:14:1899:G:H21	26:14:1902:C:H41	1.53	0.54
26:14:944:G:H2'	57:14:3688:HOH:O	2.07	0.54
1:1G:960:U:H3	1:1G:1225:A:H1'	1.73	0.54
1:1G:985:C:H2'	1:1G:986:A:H8	1.72	0.54
3:22:113:ALA:HA	3:22:202:ILE:HD11	1.90	0.54
3:22:33:LEU:O	3:22:36:ASP:N	2.41	0.54
35:25:111:PHE:O	35:25:115:VAL:HG23	2.06	0.54
29:29:109:LYS:HE2	29:29:191:PRO:HA	1.90	0.54
14:5I:23:ARG:HD2	14:5I:28:GLY:O	2.08	0.54
7:62:94:ARG:H	7:62:94:ARG:HD3	1.72	0.54
33:69:101:LEU:HD12	33:69:109:ILE:HD12	1.89	0.54
42:95:76:LYS:HB2	42:95:79:VAL:HG23	1.88	0.54
39:A8:34:HIS:CE1	39:A8:54:LEU:HD23	2.43	0.54
44:B5:42:ALA:O	44:B5:45:THR:N	2.40	0.54
1:13:1469:G:H2'	1:13:1470:G:C8	2.43	0.54
26:14:1520:U:H2'	26:14:1521:G:O4'	2.07	0.54
26:14:1771:C:O2'	26:14:1786:A:H8	1.88	0.54
26:14:2462:U:H2'	26:14:2463:C:C6	2.43	0.54
26:14:2778:A:H5''	26:14:2778:A:H8	1.73	0.54
10:1A:99:LYS:HD3	10:1A:100:THR:H	1.72	0.54
1:1G:1016:A:HO2'	1:1G:1217:C:HO2'	1.53	0.54
1:1G:25:C:H2'	1:1G:26:A:H8	1.72	0.54
26:1H:1053:C:N3	26:1H:1107:G:N2	2.56	0.54
26:1H:1406:U:H2'	26:1H:1407:C:C6	2.43	0.54
26:1H:2629:A:O2'	26:1H:2630:G:H5''	2.07	0.54
24:3L:22:G:H1'	24:3L:70:G:N2	2.23	0.54
13:4I:23:TYR:CD1	13:4I:67:GLU:HA	2.41	0.54
6:52:45:LEU:HD13	6:52:57:GLN:OE1	2.08	0.54
33:61:96:ASP:OD1	33:61:96:ASP:N	2.41	0.54
7:62:115:ARG:HB3	7:62:118:VAL:HG13	1.89	0.54
9:8E:17:VAL:HA	9:8E:63:ILE:HG12	1.89	0.54
44:B5:67:GLY:C	44:B5:69:TYR:H	2.10	0.54
54:M5:36:LYS:HE2	54:M5:36:LYS:HA	1.90	0.54
26:1H:2046:G:H1'	52:N8:22:HIS:CE1	2.42	0.54
1:13:536:C:H2'	1:13:537:G:C8	2.43	0.54
28:19:75:ILE:HG22	28:19:97:TYR:HD2	1.73	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1E:52:GLU:HG2	2:1E:56:ARG:HH12	1.73	0.54
1:1G:1402:C:H2'	1:1G:1403:C:O4'	2.08	0.54
1:1G:979:C:H5''	1:1G:980:C:OP2	2.08	0.54
26:1H:1113:U:H5'	32:51:2:SER:HB2	1.90	0.54
26:1H:1316:U:H2'	26:1H:1317:A:C8	2.43	0.54
26:1H:1339:G:H21	26:1H:1603:A:H1'	1.72	0.54
26:1H:1728:G:H2'	26:1H:1731:G:O6	2.07	0.54
26:1H:2197:U:H1'	26:1H:2198:A:C8	2.43	0.54
26:1H:2580:U:H4'	29:21:130:GLY:CA	2.33	0.54
26:1H:2875:C:H2'	26:1H:2876:G:O4'	2.08	0.54
11:2I:46:GLY:HA2	11:2I:50:TYR:O	2.08	0.54
12:3A:82:VAL:HG23	12:3A:106:ASP:OD2	2.08	0.54
31:49:34:LEU:HD21	31:49:172:LEU:HD21	1.89	0.54
9:82:10:ARG:NH2	9:82:11:LYS:HE2	2.23	0.54
9:82:77:ILE:O	9:82:81:ILE:HG12	2.07	0.54
6:5E:97:PHE:N	18:9I:30:ASP:OD1	2.40	0.54
1:13:1131:G:H2'	1:13:1132:C:H6	1.72	0.53
1:13:1397:C:H4'	1:13:1398:A:OP2	2.08	0.53
1:13:891:U:H2'	1:13:892:A:H8	1.73	0.53
26:14:1579:A:H2'	26:14:1580:A:C8	2.42	0.53
26:14:1751:C:H2'	26:14:1752:C:C6	2.43	0.53
26:14:790:C:OP2	57:14:3661:HOH:O	2.19	0.53
2:1E:6:THR:OG1	2:1E:7:VAL:N	2.40	0.53
1:1G:983:A:N1	1:1G:1222:G:N2	2.54	0.53
1:1G:1481:U:H2'	1:1G:1482:G:C8	2.44	0.53
1:1G:607:A:H2'	1:1G:608:A:O4'	2.08	0.53
26:1H:1035:U:H2'	26:1H:1036:G:C8	2.43	0.53
26:1H:2481:G:HO2'	26:1H:2482:G:P	2.31	0.53
22:1K:29:U:H2'	22:1K:30:A:C8	2.43	0.53
26:14:2823:A:OP1	29:29:113:PHE:HB2	2.07	0.53
11:2I:19:ALA:HB2	11:2I:32:ILE:HG23	1.89	0.53
30:31:29:ASN:HB3	30:31:112:MET:HE1	1.91	0.53
5:4E:75:THR:OG1	5:4E:76:ILE:N	2.39	0.53
32:51:86:GLU:N	32:51:86:GLU:OE1	2.38	0.53
7:6E:62:PHE:HA	7:6E:124:LEU:HD21	1.90	0.53
37:88:14:ARG:HG2	37:88:41:TRP:HH2	1.72	0.53
39:A8:14:VAL:O	39:A8:18:ILE:HD13	2.08	0.53
41:C8:75:ASN:HB3	41:C8:77:SER:N	2.23	0.53
33:69:27:ARG:HG2	48:F5:71:TYR:CZ	2.43	0.53
26:14:2016:U:O2	52:J5:7:PRO:HG2	2.08	0.53
2:12:119:GLU:HA	2:12:122:PHE:HD2	1.73	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:631:G:H2'	1:13:632:A:N3	2.23	0.53
1:13:917:G:H2'	1:13:918:A:C8	2.44	0.53
1:13:980:C:O2	57:13:1802:HOH:O	2.18	0.53
26:14:1288:U:C2	26:14:1327:C:O2	2.61	0.53
26:14:1796:U:H2'	26:14:1797:C:C6	2.43	0.53
26:14:774:A:O2'	26:14:775:G:O5'	2.25	0.53
34:15:13:TRP:HB2	34:15:133:GLN:HB3	1.89	0.53
1:1G:1305:G:O2'	1:1G:1306:A:O5'	2.22	0.53
1:1G:652:U:H1'	1:1G:653:A:H2	1.72	0.53
26:1H:1423:G:N7	57:1H:3919:HOH:O	2.34	0.53
26:1H:1686:C:H2'	26:1H:1687:G:O4'	2.09	0.53
26:1H:2584:U:H2'	26:1H:2585:U:H2'	1.90	0.53
27:1J:44:G:H1'	27:1J:47:C:N4	2.22	0.53
22:1L:14:A:N1	22:1L:22:G:O2'	2.41	0.53
29:29:47:VAL:HG21	29:29:86:PRO:HD2	1.90	0.53
29:29:31:CYS:N	29:29:90:THR:O	2.33	0.53
1:13:718:G:H5'	11:2I:117:ASN:ND2	2.24	0.53
23:2L:62:G:O2'	23:2L:63:G:O5'	2.25	0.53
36:35:95:VAL:HA	36:35:99:LEU:HD23	1.88	0.53
24:3L:15:A:H4'	24:3L:15:A:OP1	2.07	0.53
6:52:14:LEU:HD23	6:52:19:LEU:HB2	1.89	0.53
6:52:91:VAL:HG11	18:9A:72:ARG:NH1	2.23	0.53
7:62:143:ARG:HH11	7:62:143:ARG:HB3	1.74	0.53
36:78:97:PRO:HB3	36:78:112:LEU:HD12	1.91	0.53
42:95:7:THR:HG23	42:95:22:VAL:HG21	1.89	0.53
38:98:103:ARG:HD2	38:98:108:GLY:O	2.09	0.53
26:1H:1156:A:C8	41:C8:51:LYS:HD2	2.43	0.53
41:C8:92:ARG:CZ	42:D8:11:GLN:H	2.21	0.53
45:G8:49:VAL:HG21	45:G8:55:TYR:CD2	2.43	0.53
31:41:98:ARG:NE	51:M8:1:MET:SD	2.81	0.53
54:Q8:59:LYS:CD	54:Q8:59:LYS:H	2.19	0.53
1:13:1003:G:H1	1:13:1037:C:H42	1.56	0.53
1:13:105:G:H2'	1:13:106:C:C6	2.42	0.53
1:13:1431:C:H2'	1:13:1432:G:O4'	2.09	0.53
1:13:77:C:O2'	1:13:92:G:N2	2.25	0.53
26:14:1044:G:H4'	26:14:1048:A:H1'	1.91	0.53
26:14:2111:C:H42	26:14:2147:G:H21	1.55	0.53
26:14:527:C:N4	26:14:2779:U:OP2	2.41	0.53
1:1G:1132:C:H2'	1:1G:1133:G:H8	1.74	0.53
1:1G:722:A:H2	1:1G:733:A:H61	1.56	0.53
26:1H:1244:G:OP1	36:78:7:ARG:HD3	2.08	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1467:C:H42	26:1H:1525:G:H1	1.57	0.53
26:1H:2156:G:H2'	26:1H:2157:G:C2	2.43	0.53
26:1H:586:A:OP2	57:1H:3731:HOH:O	2.19	0.53
24:3L:67:C:N3	26:14:2169:A:N6	2.56	0.53
31:41:37:VAL:O	31:41:94:LEU:HD23	2.08	0.53
1:1G:1226:C:N4	13:4A:104:ARG:HD2	2.24	0.53
13:4A:78:ILE:HD13	13:4A:92:HIS:CE1	2.44	0.53
33:61:29:TYR:C	33:61:32:PRO:HD2	2.29	0.53
15:6I:32:LEU:O	15:6I:35:ARG:N	2.42	0.53
40:75:53:ARG:O	40:75:53:ARG:HG3	2.07	0.53
40:75:96:ARG:HH11	40:75:96:ARG:HG2	1.73	0.53
36:78:52:GLU:OE2	36:78:58:THR:HG22	2.07	0.53
18:9A:31:LEU:H	18:9A:31:LEU:HD23	1.71	0.53
48:F5:91:LYS:HZ3	48:F5:91:LYS:HA	1.73	0.53
45:G8:76:CYS:SG	45:G8:97:ARG:HG2	2.49	0.53
51:I5:22:ILE:HG12	51:I5:23:GLU:N	2.22	0.53
28:11:218:ARG:HB3	28:11:219:PRO:HD2	1.91	0.53
26:14:2110:G:O2'	26:14:2120:G:OP1	2.15	0.53
26:14:314:A:H2'	26:14:315:G:C8	2.44	0.53
2:1E:136:VAL:O	2:1E:140:HIS:N	2.42	0.53
1:1G:1505:G:H4'	1:1G:1506:U:H5''	1.91	0.53
1:1G:25:C:H2'	1:1G:26:A:C8	2.43	0.53
1:1G:371:G:H1	1:1G:390:C:H42	1.55	0.53
1:1G:682:G:H1	1:1G:708:C:H42	1.55	0.53
26:1H:2324:C:H5''	26:1H:2325:G:H5'	1.90	0.53
26:1H:2479:G:O5'	26:1H:2479:G:H8	1.91	0.53
26:1H:2735:G:H2'	26:1H:2736:G:H8	1.73	0.53
10:1I:3:LYS:N	10:1I:75:ILE:O	2.41	0.53
27:1J:15:A:H1'	27:1J:109:G:C8	2.43	0.53
27:1J:94:C:H2'	27:1J:95:U:C6	2.43	0.53
29:29:35:GLN:O	29:29:48:GLN:HB2	2.09	0.53
11:2A:32:ILE:HD11	11:2A:68:ALA:HB1	1.90	0.53
36:35:59:LEU:HD21	54:M5:10:ALA:HA	1.89	0.53
30:39:102:PRO:HB2	30:39:105:VAL:HG23	1.90	0.53
31:49:71:THR:N	31:49:89:GLY:O	2.37	0.53
13:4A:79:LYS:HG3	13:4A:82:MET:SD	2.49	0.53
35:68:4:PRO:O	35:68:5:GLN:HB2	2.07	0.53
33:69:85:GLU:HA	33:69:123:LEU:HD12	1.90	0.53
9:82:63:ILE:HD13	9:82:77:ILE:HG23	1.91	0.53
17:8I:68:ARG:H	17:8I:70:ARG:NH1	2.06	0.53
43:E8:48:ALA:O	43:E8:52:GLU:HB2	2.08	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:372:C:N4	1:13:389:A:H62	2.04	0.53
26:14:829:A:N7	26:14:2247:A:O2'	2.35	0.53
34:15:42:TRP:HA	34:15:48:MET:HE1	1.91	0.53
26:1H:1316:U:H2'	26:1H:1317:A:H8	1.73	0.53
26:1H:1411:C:H2'	26:1H:1412:A:H8	1.73	0.53
26:1H:1453:A:O2'	26:1H:1454:U:H2'	2.08	0.53
26:1H:1636:C:H2'	26:1H:1637:A:C8	2.43	0.53
26:1H:906:G:OP1	37:88:26:TYR:OH	2.21	0.53
26:14:2786:U:O2	29:29:62:PRO:HB3	2.09	0.53
30:31:60:SER:OG	30:31:61:GLY:N	2.42	0.53
26:14:588:U:H1'	30:39:90:PHE:CG	2.44	0.53
4:3E:104:VAL:O	4:3E:107:ARG:N	2.41	0.53
24:3K:46:G:H2'	24:3K:47:C:C5'	2.37	0.53
34:58:6:PRO:HG3	34:58:41:ASP:HB2	1.90	0.53
7:62:143:ARG:NH1	7:62:143:ARG:HB3	2.24	0.53
1:1G:878:G:H5'	8:72:89:PRO:HG2	1.90	0.53
9:82:7:THR:H	9:82:83:ARG:HD2	1.73	0.53
26:14:64:A:O3'	44:B5:71:GLY:HA3	2.09	0.53
41:C8:97:ASP:OD1	41:C8:98:LEU:N	2.42	0.53
47:E5:29:GLN:O	47:E5:67:VAL:HG23	2.08	0.53
45:G8:93:GLY:O	45:G8:94:LYS:HB2	2.08	0.53
2:12:71:VAL:HB	2:12:164:VAL:HG13	1.90	0.53
26:14:654(R):C:N4	26:14:654(S):G:O6	2.42	0.53
1:1G:1157:A:H8	1:1G:1158:C:C5	2.27	0.53
1:1G:164:U:H2'	1:1G:165:C:C6	2.42	0.53
1:1G:377:G:H1	1:1G:386:C:H42	1.56	0.53
1:1G:410:G:N1	1:1G:431:A:OP2	2.41	0.53
1:1G:674:G:H2'	1:1G:675:A:C8	2.44	0.53
26:1H:1586:A:H3'	26:1H:1587:A:H8	1.74	0.53
26:1H:2807:G:H3'	26:1H:2808:U:H5''	1.90	0.53
1:1G:1112:C:C4	3:22:178:LEU:HD23	2.43	0.53
11:2A:13:GLN:HA	11:2A:75:TYR:O	2.08	0.53
31:41:173:LEU:HB3	31:41:178:PHE:CD2	2.42	0.53
32:59:42:ARG:NH1	32:59:53:GLU:O	2.37	0.53
40:75:4:GLY:O	40:75:7:ILE:N	2.41	0.53
9:82:51:ARG:HG2	9:82:56:LEU:HD22	1.89	0.53
19:AA:9:VAL:HG13	51:I5:63:TYR:HE1	1.73	0.53
36:78:49:ARG:HG3	54:Q8:57:ARG:HE	1.73	0.53
1:13:1139:G:H4'	1:13:1140:C:H5'	1.91	0.53
26:14:2219:G:H2'	26:14:2224:G:H5'	1.90	0.53
26:14:844:C:H3'	26:14:845:G:C8	2.44	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1521:G:H2'	1:1G:1522:U:C6	2.43	0.53
26:1H:1069:A:O2'	26:1H:1072:C:OP2	2.26	0.53
26:1H:1329:U:H5''	26:1H:1330:C:H5	1.73	0.53
26:1H:1337:G:H2'	26:1H:1338:G:H8	1.74	0.53
26:1H:1359:A:C2	26:1H:1372:U:O4	2.57	0.53
26:1H:265:A:C8	26:1H:266:G:H1'	2.44	0.53
30:31:181:LEU:O	30:31:205:ARG:NH2	2.41	0.53
26:1H:674:G:C1'	30:31:74:ARG:HD3	2.37	0.53
36:35:84:ASN:ND2	36:35:117:GLU:HB3	2.24	0.53
37:45:27:VAL:H	37:45:102:VAL:HG21	1.74	0.53
31:49:40:ASN:HB2	31:49:91:ARG:HB2	1.90	0.53
13:4A:40:ASN:HB3	13:4A:43:THR:HG23	1.90	0.53
13:4A:91:ARG:NH1	13:4A:96:LEU:HD13	2.24	0.53
6:52:25:ILE:HD12	6:52:82:ARG:HD2	1.90	0.53
39:65:39:ILE:HD12	39:65:85:VAL:HG11	1.91	0.53
9:82:43:ALA:HA	9:82:74:ILE:HD13	1.90	0.53
38:98:117:VAL:HG22	38:98:118:GLU:H	1.73	0.53
43:A5:9:TYR:H	43:A5:102:HIS:CE1	2.25	0.53
46:D5:23:LYS:HB3	46:D5:38:TYR:CD1	2.44	0.53
54:Q8:14:VAL:HB	54:Q8:21:LYS:HE2	1.90	0.53
54:Q8:7:HIS:O	54:Q8:7:HIS:ND1	2.42	0.53
2:12:84:GLU:OE2	2:12:212:GLN:NE2	2.42	0.53
2:12:44:LEU:HD23	2:12:44:LEU:H	1.74	0.53
26:14:1432:C:H2'	26:14:1433:U:O4'	2.08	0.53
26:14:1444(A):A:N3	26:14:1444(A):A:H2'	2.24	0.53
26:14:990:A:H8	26:14:990:A:H5'	1.73	0.53
1:1G:147:G:N2	1:1G:148:G:C4	2.77	0.53
1:1G:622:A:C8	1:1G:623:C:C6	2.97	0.53
26:1H:124:G:N2	26:1H:126:A:O3'	2.42	0.53
26:1H:270(P):C:H2'	26:1H:270(Q):C:C6	2.44	0.53
26:1H:50:U:H3'	26:1H:51:G:H5'	1.91	0.53
26:1H:637:A:H2'	36:78:117:GLU:OE1	2.09	0.53
10:1I:46:ARG:NH2	10:1I:64:GLU:OE1	2.42	0.53
3:2E:83:ARG:HA	3:2E:86:VAL:HG13	1.91	0.53
37:45:34:LEU:HB2	37:45:118:LEU:HD13	1.90	0.53
33:61:4:ILE:HG12	33:61:18:VAL:HG22	1.91	0.53
7:6E:73:MET:HA	7:6E:90:GLU:HA	1.91	0.53
16:7I:77:ALA:HB3	16:7I:79:VAL:HG23	1.91	0.53
9:82:85:LEU:HD12	9:82:92:TYR:CD2	2.44	0.53
37:88:20:ALA:HA	37:88:98:LYS:HB3	1.90	0.53
20:BA:26:ASN:HB3	20:BA:71:THR:OG1	2.09	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
45:C5:52:SER:HA	45:C5:55:TYR:C	2.28	0.53
2:12:104:ASN:OD1	2:12:107:THR:OG1	2.26	0.53
1:13:1107:C:C4	1:13:1108:G:C8	2.97	0.53
1:13:920:U:H2'	1:13:921:U:C6	2.44	0.53
26:14:1449(A):G:H2'	26:14:1450:C:H6	1.74	0.53
26:14:2655:G:N2	26:14:2665:A:OP2	2.42	0.53
1:1G:529:G:O6	12:3A:49:ASN:HA	2.09	0.53
1:1G:628:G:H2'	1:1G:629:G:H8	1.73	0.53
26:1H:1086:A:O2'	26:1H:1103:A:N1	2.38	0.53
26:1H:1110:G:O2'	26:1H:1111:A:O5'	2.26	0.53
26:1H:2062:A:H2'	26:1H:2062:A:N3	2.23	0.53
26:1H:692:C:O2'	28:11:38:LYS:HE3	2.09	0.53
26:1H:825:C:H5'	57:1H:3680:HOH:O	2.09	0.53
13:4I:4:ILE:HG22	13:4I:5:ALA:H	1.73	0.53
32:51:4:ILE:C	32:51:6:ARG:H	2.12	0.53
33:69:60:GLU:HG3	33:69:61:ARG:N	2.24	0.53
7:6E:38:LEU:O	7:6E:42:ILE:HG13	2.09	0.53
15:6I:29:VAL:HG11	15:6I:67:LEU:HD21	1.90	0.53
37:88:66:ILE:O	37:88:104:PHE:N	2.40	0.53
9:8E:3:GLN:OE1	9:8E:20:ARG:NH1	2.42	0.53
42:95:35:LEU:HB3	42:95:37:VAL:HG11	1.91	0.53
42:95:71:LEU:O	42:95:72:VAL:HG12	2.08	0.53
39:A8:5:THR:OG1	39:A8:8:GLU:HG2	2.08	0.53
44:B5:65:ARG:HG3	44:B5:67:GLY:H	1.72	0.53
45:C5:17:SER:OG	45:C5:18:GLY:O	2.25	0.53
48:F5:2:SER:O	48:F5:4:VAL:HG13	2.09	0.53
54:Q8:27:THR:HG22	54:Q8:42:ARG:HH21	1.74	0.53
1:13:1070:U:H2'	1:13:1071:C:H6	1.73	0.53
1:13:1131:G:H2'	1:13:1132:C:C6	2.44	0.53
1:13:1250:A:OP1	9:8E:67:GLY:N	2.30	0.53
1:13:991:U:C4	1:13:1212:U:H1'	2.44	0.53
26:14:2239:G:H5'	28:19:251:GLY:HA3	1.89	0.53
26:14:2757:A:N1	32:59:67:LEU:HD23	2.24	0.53
26:14:860:U:H1'	26:14:2268:A:H5'	1.91	0.53
27:16:7:G:H4'	39:A8:29:PHE:CD2	2.44	0.53
1:1G:1502:A:H2	1:1G:1505:G:N1	2.00	0.53
26:1H:1039:G:H1	26:1H:1116:C:H42	1.56	0.53
26:1H:1178:C:H4'	26:1H:1179:C:OP1	2.08	0.53
10:1I:48:THR:HA	10:1I:62:HIS:HB3	1.90	0.53
23:2K:63:G:C4	23:2K:64:G:C8	2.97	0.53
23:2K:7:G:OP2	23:2K:7:G:H8	1.92	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:35:27:HIS:HB3	36:35:32:THR:HG23	1.90	0.53
31:41:7:LEU:HB2	31:41:104:GLU:HB2	1.90	0.53
14:5I:15:LYS:HG2	14:5I:16:PHE:CD2	2.44	0.53
40:75:4:GLY:HA2	40:75:8:LYS:HB2	1.90	0.53
40:B8:87:ASP:OD1	40:B8:87:ASP:N	2.42	0.53
1:13:1033:G:H2'	1:13:1034:G:H8	1.74	0.52
1:13:1178:G:N2	1:13:1181:G:C8	2.76	0.52
1:13:1422:G:H1	1:13:1478:C:H42	1.55	0.52
1:13:223:U:H2'	1:13:224:C:C6	2.43	0.52
1:13:749:C:H2'	1:13:750:G:H8	1.73	0.52
26:14:1041:C:H2'	26:14:1042:G:H8	1.73	0.52
26:14:2789:C:H1'	26:14:2892:A:H2	1.74	0.52
26:14:843:G:H1	26:14:935:C:H42	1.56	0.52
27:16:15:A:H1'	27:16:109:G:N9	2.24	0.52
2:1E:70:PHE:O	2:1E:93:VAL:N	2.32	0.52
1:1G:674:G:H2'	1:1G:675:A:H8	1.72	0.52
26:1H:1437:C:H2'	26:1H:1438:U:C6	2.42	0.52
26:1H:1858:G:H1'	26:1H:1884:A:N6	2.24	0.52
26:1H:2729:G:H2'	26:1H:2730:C:H6	1.73	0.52
26:1H:277:C:H3'	26:1H:278:A:O4'	2.10	0.52
26:1H:49:A:N7	26:1H:120:U:C5	2.68	0.52
10:1I:59:SER:OG	10:1I:59:SER:O	2.21	0.52
22:1K:8:U:H1'	22:1K:22:G:H21	1.74	0.52
22:1K:87:A:O3'	26:1H:2506:U:H1'	2.08	0.52
4:32:168:ARG:HD3	4:32:169:LYS:H	1.74	0.52
31:41:151:ALA:O	31:41:153:ARG:NH1	2.42	0.52
13:4A:4:ILE:HG23	13:4A:5:ALA:H	1.74	0.52
7:6E:16:LEU:HD21	9:8E:45:ALA:HB2	1.91	0.52
19:AA:27:GLU:O	19:AA:47:HIS:NE2	2.39	0.52
48:F5:91:LYS:HA	48:F5:91:LYS:NZ	2.23	0.52
1:13:1280:A:H3'	1:13:1281:U:H5'	1.91	0.52
1:13:1513:A:H2'	1:13:1514:C:C6	2.43	0.52
1:13:554:C:H2'	1:13:555:C:C6	2.44	0.52
1:13:728:A:C5	15:6I:54:ARG:HD2	2.44	0.52
1:13:939:G:H5''	7:6E:102:ARG:NH2	2.24	0.52
26:14:1778:U:H2'	26:14:1784:A:N6	2.24	0.52
26:14:2719:G:OP2	57:14:3591:HOH:O	2.18	0.52
26:14:308:G:C8	26:14:501:A:H1'	2.44	0.52
27:16:21:G:N2	27:16:62:C:N3	2.46	0.52
1:1G:535:A:H5''	57:1G:1809:HOH:O	2.08	0.52
26:1H:176:G:O2'	26:1H:177:G:H5'	2.09	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2224:G:H4'	26:1H:2226:C:C2	2.45	0.52
26:1H:2062:A:N6	26:1H:2503:A:H62	2.07	0.52
26:1H:882:G:OP1	26:1H:882:G:H4'	2.09	0.52
26:14:1026:U:H2'	27:1J:88:C:N4	2.24	0.52
3:22:47:LEU:HD11	3:22:70:VAL:HG22	1.92	0.52
23:2L:71:U:H5''	23:2L:72:C:C5	2.40	0.52
30:31:140:LEU:HD13	30:31:170:LEU:HD21	1.91	0.52
4:32:107:ARG:HH22	4:32:196:LEU:HD21	1.73	0.52
36:35:98:GLU:HA	36:35:101:VAL:HG12	1.91	0.52
4:3E:98:GLU:HG3	4:3E:103:ASN:HD21	1.73	0.52
5:42:76:ILE:HG22	5:42:77:PRO:HD2	1.91	0.52
1:1G:994:A:C2	14:5A:5:ALA:HB2	2.44	0.52
36:78:2:LYS:HD3	36:78:4:SER:HB2	1.91	0.52
8:7E:7:ALA:HB2	8:7E:85:ARG:HD2	1.91	0.52
28:11:132:PRO:HD3	28:11:190:TYR:CZ	2.44	0.52
1:13:1243:C:H42	1:13:1294:G:H1	1.57	0.52
1:13:1469:G:H2'	1:13:1470:G:H8	1.75	0.52
1:13:221:C:H2'	1:13:222:U:C6	2.44	0.52
26:14:1278:A:OP1	38:55:36:THR:HG22	2.09	0.52
26:14:1729:A:H2'	26:14:1731:G:N2	2.24	0.52
26:14:363(E):U:H5'	26:14:363(F):A:OP2	2.10	0.52
26:14:947:G:H2'	26:14:948:G:C8	2.44	0.52
34:15:99:LEU:O	34:15:103:VAL:HG23	2.10	0.52
1:1G:1128:C:H4'	9:82:16:ARG:HH12	1.75	0.52
1:1G:841:U:O2'	1:1G:842:C:H5''	2.09	0.52
1:1G:854:G:C2	1:1G:855:G:C8	2.97	0.52
26:1H:1510:A:N3	26:1H:1510:A:H2'	2.23	0.52
26:1H:2022:U:O2'	26:1H:2617:C:H5'	2.09	0.52
26:1H:2244:U:H2'	26:1H:2245:U:O4'	2.09	0.52
26:1H:2488:A:H2'	26:1H:2489:G:O4'	2.09	0.52
26:1H:2784:C:H2'	26:1H:2785:C:C6	2.43	0.52
1:13:1152:A:O3'	10:1I:13:HIS:NE2	2.42	0.52
22:1L:75:G:H2'	22:1L:76:U:C6	2.43	0.52
29:21:5:LEU:HD12	29:21:51:PHE:HB2	1.90	0.52
29:29:68:ALA:C	29:29:70:ALA:H	2.13	0.52
23:2K:73:C:H2'	23:2K:74:C:C6	2.43	0.52
23:2L:15:A:H2'	23:2L:70:G:H1	1.75	0.52
4:3E:13:ARG:HD3	4:3E:36:ARG:O	2.09	0.52
12:3I:66:VAL:HG21	12:3I:98:TYR:HE1	1.75	0.52
32:51:20:ALA:HB1	32:51:21:PRO:HD2	1.89	0.52
40:B8:3:ARG:O	40:B8:7:ILE:N	2.36	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2262:U:P	47:E5:19:LYS:HE3	2.49	0.52
43:E8:29:LEU:O	43:E8:29:LEU:HG	2.10	0.52
46:H8:125:LEU:HG	46:H8:164:ALA:HB2	1.91	0.52
1:13:356:A:N3	1:13:368:U:O2'	2.42	0.52
1:13:792:A:H4'	1:13:793:U:O5'	2.09	0.52
26:14:1169:G:C2	26:14:1170:G:H1'	2.44	0.52
26:14:654(I):C:H5''	26:14:654(M):C:H42	1.73	0.52
27:16:29:A:H2'	27:16:30:C:O4'	2.09	0.52
28:19:242:ARG:H	28:19:242:ARG:HH11	1.58	0.52
28:19:83:GLU:HB2	28:19:92:ILE:HG13	1.92	0.52
1:1G:632:A:H1'	1:1G:633:G:OP2	2.10	0.52
35:25:71:ARG:NE	35:25:105:GLU:OE1	2.37	0.52
24:3L:9:G:H3'	24:3L:10:G:H5''	1.91	0.52
31:49:138:GLN:OE1	31:49:153:ARG:HG2	2.09	0.52
7:62:46:ALA:O	7:62:50:ILE:HG12	2.09	0.52
9:82:49:PRO:HD3	9:82:78:LYS:HZ2	1.75	0.52
41:85:66:ASN:ND2	41:85:70:ARG:HE	2.07	0.52
1:13:1117:G:H5''	9:8E:104:ARG:NH1	2.25	0.52
39:A8:109:GLY:HA3	39:A8:110:LEU:HD23	1.91	0.52
54:Q8:50:LEU:O	54:Q8:52:LYS:N	2.40	0.52
28:11:155:LEU:HD13	28:11:155:LEU:N	2.25	0.52
1:13:486:U:H2'	1:13:487:A:H8	1.75	0.52
26:14:1606:G:H5''	26:14:1607:C:OP1	2.10	0.52
26:14:185:U:H4'	26:14:218:A:H4'	1.91	0.52
26:14:1871:A:H2'	26:14:1872:A:C8	2.45	0.52
26:14:247:G:H4'	26:14:386:G:C5	2.44	0.52
26:14:2635:C:OP1	29:29:77:ILE:HB	2.08	0.52
26:14:307:G:H8	26:14:307:G:O5'	1.93	0.52
1:1G:111:G:O5'	1:1G:111:G:H8	1.92	0.52
26:1H:144:C:H2'	26:1H:145:G:H8	1.74	0.52
26:1H:1914:C:H2'	26:1H:1915:U:O4'	2.10	0.52
26:1H:2308:G:N3	26:1H:2308:G:H2'	2.25	0.52
26:1H:37:C:H2'	26:1H:38:A:C8	2.44	0.52
30:39:30:PRO:O	30:39:33:LEU:N	2.42	0.52
22:1L:64:G:O3'	37:45:56:ARG:NH2	2.43	0.52
13:4A:92:HIS:CD2	13:4A:98:VAL:HG11	2.45	0.52
32:51:104:GLU:HB2	32:51:114:VAL:HG13	1.91	0.52
33:61:62:LYS:HA	33:61:133:HIS:NE2	2.25	0.52
19:AI:6:LYS:O	19:AI:7:LYS:NZ	2.26	0.52
40:B8:121:ILE:O	40:B8:124:ASP:HB2	2.09	0.52
54:Q8:36:LYS:HG2	54:Q8:39:LYS:HZ1	1.74	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:728:G:H4'	28:11:13:ARG:HD3	1.91	0.52
1:13:270:A:H2'	1:13:271:C:O4'	2.10	0.52
1:13:736:C:H2'	1:13:737:A:H8	1.75	0.52
1:13:789:U:C5	1:13:791:G:H3'	2.45	0.52
1:13:922:G:H1'	5:4E:19:MET:HB2	1.91	0.52
26:14:193:U:H5	57:14:3644:HOH:O	1.93	0.52
26:14:662:G:OP1	36:35:14:LYS:NZ	2.20	0.52
26:1H:861:A:N3	27:16:79:C:O2'	2.43	0.52
1:1G:1148:U:H2'	1:1G:1149:C:O4'	2.10	0.52
1:1G:668:G:HO2'	15:6A:46:HIS:HB3	1.75	0.52
26:1H:1342:A:OP1	44:F8:36:LYS:NZ	2.30	0.52
26:1H:1838:C:O3'	57:1H:3834:HOH:O	2.19	0.52
26:1H:309:G:N3	26:1H:329:G:O2'	2.43	0.52
26:1H:881:G:H5'	26:1H:882:G:O5'	2.09	0.52
29:21:152:LYS:HG2	34:58:78:TYR:CE1	2.45	0.52
3:22:36:ASP:OD1	3:22:57:ILE:HG21	2.10	0.52
30:39:165:ARG:HH11	30:39:165:ARG:HB3	1.74	0.52
4:3E:29:PRO:HA	4:3E:34:GLU:HG3	1.92	0.52
37:45:26:TYR:HD1	37:45:26:TYR:O	1.93	0.52
6:5E:23:LYS:HG3	6:5E:61:LEU:HD21	1.90	0.52
38:98:2:ARG:HH11	38:98:2:ARG:HG2	1.74	0.52
26:14:483:A:H1'	45:C5:60:PHE:CE1	2.45	0.52
43:E8:17:VAL:HG13	43:E8:76:VAL:HG11	1.92	0.52
48:F5:15:ALA:O	48:F5:40:ARG:HG2	2.10	0.52
54:Q8:29:LYS:O	54:Q8:30:ARG:NH1	2.39	0.52
54:Q8:31:HIS:ND1	54:Q8:31:HIS:O	2.40	0.52
1:13:404:U:H2'	1:13:405:U:C6	2.44	0.52
1:13:973:G:O4'	10:1I:55:LYS:HB3	2.10	0.52
26:14:13:A:N1	26:14:525:U:H2'	2.25	0.52
26:14:2646:C:H2'	26:14:2647:U:O4'	2.10	0.52
26:14:57:C:H2'	26:14:58:G:O4'	2.10	0.52
28:19:166:GLN:HB3	28:19:174:ILE:HG22	1.91	0.52
1:1G:1274:G:H2'	1:1G:1275:A:C8	2.44	0.52
1:1G:173:U:O2	1:1G:197:A:N6	2.43	0.52
1:1G:313:A:H2'	1:1G:314:C:C6	2.45	0.52
1:1G:858:G:H8	1:1G:858:G:OP2	1.93	0.52
26:1H:1047:G:H2'	26:1H:1110:G:N1	2.25	0.52
26:1H:1454:U:O2'	26:1H:1455:G:N7	2.42	0.52
26:1H:1534:G:H3'	26:1H:1534:G:N3	2.24	0.52
26:1H:1332:G:N2	26:1H:1609:A:O2'	2.42	0.52
26:1H:2110:G:N2	26:1H:2180:U:O2	2.43	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2748:A:OP1	32:51:70:THR:OG1	2.18	0.52
26:1H:2843:G:H1	26:1H:2874:C:N4	2.00	0.52
26:1H:587:C:OP2	36:78:21:ARG:NH2	2.43	0.52
30:31:114:VAL:HG21	30:31:202:PHE:CZ	2.45	0.52
5:42:126:ARG:HG3	5:42:126:ARG:HH11	1.74	0.52
13:4I:39:ILE:HD13	13:4I:52:GLU:HB3	1.92	0.52
7:62:126:ASP:HB3	7:62:131:LYS:O	2.09	0.52
26:14:270(L):U:O2	33:69:50:ARG:HD2	2.10	0.52
40:75:61:PHE:CE1	40:75:76:PHE:HB2	2.45	0.52
16:7A:74:LEU:HD12	16:7A:79:VAL:HG21	1.91	0.52
26:14:1225:C:C4'	42:95:85:LYS:HA	2.36	0.52
20:BA:67:ALA:HB2	20:BA:77:ALA:HB2	1.92	0.52
2:12:82:ARG:NH1	2:12:92:TYR:OH	2.43	0.52
1:13:765:G:H5''	1:13:766:A:OP1	2.10	0.52
1:13:791:G:C6	1:13:792:A:C2	2.98	0.52
26:14:1184:G:H5'	50:H5:29:ARG:HH21	1.75	0.52
26:14:2318:G:H1	39:65:2:ALA:HA	1.73	0.52
26:14:99:U:H4'	26:14:102:G:H1'	1.91	0.52
1:1G:1028:C:N4	1:1G:1033:G:H1	2.08	0.52
26:1H:1038:C:H2'	26:1H:1039:G:O4'	2.09	0.52
26:1H:1582:C:H2'	26:1H:1583:A:O4'	2.10	0.52
26:1H:1635:G:H2'	26:1H:1636:C:C6	2.45	0.52
11:2A:100:ALA:O	11:2A:102:GLY:N	2.43	0.52
23:2K:86:C:H3'	23:2K:87:A:H5''	1.89	0.52
23:2L:86:C:H2'	23:2L:87:A:C8	2.45	0.52
36:35:111:ARG:HG3	36:35:128:HIS:CG	2.45	0.52
24:3L:5:A:H61	24:3L:79:U:H3	1.58	0.52
37:45:17:LEU:HD21	37:45:41:TRP:HE1	1.75	0.52
5:4E:28:PHE:O	5:4E:47:LYS:HA	2.10	0.52
13:4I:14:ARG:HB2	13:4I:14:ARG:HH11	1.75	0.52
38:55:37:THR:OG1	38:55:40:LYS:HE3	2.10	0.52
32:59:122:THR:OG1	32:59:134:SER:OG	2.28	0.52
6:5E:39:LYS:HD3	6:5E:64:GLN:HG3	1.91	0.52
1:1G:751:U:H4'	15:6A:24:SER:HA	1.92	0.52
7:6E:69:VAL:HG22	7:6E:135:VAL:HG13	1.90	0.52
8:72:14:ARG:O	8:72:17:THR:HG22	2.09	0.52
36:78:144:GLU:N	36:78:144:GLU:OE2	2.42	0.52
17:8A:59:ILE:HD13	17:8A:73:VAL:HA	1.92	0.52
13:4A:86:CYS:HA	19:AA:73:GLU:O	2.10	0.52
45:C5:63:LYS:HA	45:C5:63:LYS:NZ	2.25	0.52
54:Q8:35:GLN:HB3	54:Q8:37:SER:O	2.10	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:Q8:52:LYS:HA	54:Q8:54:GLU:HB2	1.91	0.52
1:13:1179:A:H2'	1:13:1180:A:O4'	2.10	0.52
1:13:312:C:H2'	1:13:313:A:H8	1.74	0.52
26:14:1609:A:OP2	57:14:3603:HOH:O	2.19	0.52
26:14:2601:C:H2'	26:14:2603:G:C8	2.45	0.52
26:14:29:U:O4	57:14:3809:HOH:O	2.19	0.52
26:14:780:G:N2	26:14:783:A:H62	2.03	0.52
10:1A:65:LEU:HD12	14:5A:55:GLY:O	2.10	0.52
2:1E:77:ALA:HB1	2:1E:165:VAL:HG11	1.90	0.52
26:1H:86:C:H4'	26:1H:104:U:H1'	1.91	0.52
26:1H:1729:A:HO2'	26:1H:1730:U:P	2.33	0.52
26:1H:2712:U:H1'	26:1H:2712(A):A:C8	2.45	0.52
26:1H:844:C:H3'	26:1H:845:G:H8	1.74	0.52
26:1H:848:G:H2'	26:1H:849:A:C8	2.45	0.52
1:13:963:G:H21	10:1I:55:LYS:CE	2.22	0.52
10:1I:92:THR:OG1	10:1I:93:GLY:N	2.43	0.52
30:39:123:LEU:HB2	30:39:192:LEU:HB3	1.90	0.52
1:13:403:C:O3'	4:3E:122:ARG:HD2	2.10	0.52
31:49:143:GLU:OE2	31:49:143:GLU:N	2.42	0.52
13:4I:3:ARG:HD3	13:4I:7:VAL:HG13	1.92	0.52
32:51:40:GLU:OE1	32:51:41:MET:N	2.43	0.52
26:14:2882:A:H5'	38:55:96:ARG:HG3	1.92	0.52
1:13:624:C:O3'	16:7I:10:GLY:HA2	2.10	0.52
9:8E:18:PHE:HD2	9:8E:62:TYR:HD2	1.58	0.52
19:AI:40:ILE:HG22	19:AI:69:HIS:O	2.10	0.52
43:E8:73:ALA:HB3	43:E8:106:ILE:HD12	1.91	0.52
53:P8:35:ARG:HG3	53:P8:42:LEU:HD11	1.92	0.52
2:12:70:PHE:O	2:12:93:VAL:N	2.39	0.52
1:13:1075:C:OP1	2:1E:179:LYS:NZ	2.27	0.52
26:14:1021:A:H2'	26:14:1023:U:H5'	1.91	0.52
26:14:1198:U:H2'	26:14:1199:U:C6	2.45	0.52
26:14:1838:C:N4	26:14:1898:U:H2'	2.25	0.52
10:1A:36:GLY:O	10:1A:38:ILE:HG13	2.10	0.52
1:1G:413:G:HO2'	1:1G:414:A:P	2.28	0.52
26:1H:1535:U:H2'	26:1H:1536:A:H5''	1.92	0.52
26:1H:1543:A:H3'	26:1H:1543:A:OP2	2.10	0.52
26:1H:2068:U:N3	26:1H:2430:A:H2	2.07	0.52
26:1H:263:C:H2'	26:1H:264:C:O4'	2.10	0.52
26:1H:389:G:H22	36:78:72:PRO:HD3	1.74	0.52
11:2A:32:ILE:HD13	11:2A:72:ALA:HB2	1.92	0.52
23:2L:74:C:H2'	23:2L:75:G:H8	1.74	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:42:6:PHE:HB2	5:42:34:VAL:HG22	1.92	0.52
15:6A:56:LEU:O	15:6A:60:VAL:HG23	2.10	0.52
9:82:118:LYS:HB3	9:82:118:LYS:HZ3	1.75	0.52
17:8I:25:ARG:CZ	17:8I:27:PHE:HE2	2.23	0.52
18:9I:66:LEU:O	18:9I:70:ILE:HG13	2.10	0.52
19:AA:29:ARG:NH1	19:AA:48:THR:H	2.08	0.52
46:D5:25:PRO:O	46:D5:85:HIS:HA	2.09	0.52
44:F8:24:GLY:O	44:F8:83:VAL:HG22	2.10	0.52
26:1H:2383:G:N7	54:Q8:34:TRP:CZ2	2.76	0.52
1:13:446:G:H1	1:13:488:C:H42	1.58	0.51
26:14:2270:G:OP2	57:14:3775:HOH:O	2.19	0.51
26:14:2666:C:C2'	26:14:2667:C:H5'	2.40	0.51
26:14:28:A:O2'	26:14:583:G:H5'	2.10	0.51
27:16:112:G:H2'	27:16:113:C:C6	2.45	0.51
2:1E:164:VAL:HG23	2:1E:186:ALA:HB2	1.91	0.51
1:1G:1179:A:H2'	1:1G:1180:A:O4'	2.10	0.51
1:1G:165:C:H2'	1:1G:166:G:C8	2.45	0.51
1:1G:551:U:H2'	1:1G:552:U:H6	1.74	0.51
26:1H:1449:A:H5'	26:1H:1449(A):G:OP2	2.10	0.51
26:1H:14:A:H3'	26:1H:15:G:H5''	1.92	0.51
26:1H:1945:G:H2'	26:1H:1946:U:C6	2.44	0.51
26:1H:445:C:H2'	26:1H:446:G:C8	2.45	0.51
26:1H:518:G:H2'	26:1H:519:U:C6	2.45	0.51
10:1I:48:THR:HA	10:1I:62:HIS:HB2	1.90	0.51
27:1J:6:C:C2	27:1J:115:G:N2	2.78	0.51
29:21:37:ARG:O	29:21:45:THR:HA	2.10	0.51
29:29:47:VAL:HG12	29:29:49:LEU:HD23	1.92	0.51
3:2E:43:LEU:HD22	3:2E:47:LEU:HD12	1.92	0.51
3:2E:83:ARG:O	3:2E:86:VAL:HG22	2.09	0.51
23:2L:32:C:H2'	23:2L:33:U:O4'	2.10	0.51
4:32:57:ARG:HE	4:32:205:GLU:HB3	1.73	0.51
24:3K:32:C:O2	24:3K:40:G:N2	2.38	0.51
31:41:34:LEU:HD23	31:41:172:LEU:HD21	1.90	0.51
1:1G:580:U:H5''	15:6A:58:MET:HG2	1.92	0.51
44:B5:55:ASN:HB2	44:B5:80:ILE:CG1	2.37	0.51
26:14:2262:U:H5	47:E5:16:SER:HG	1.56	0.51
48:F5:4:VAL:HG11	48:F5:11:ARG:NH1	2.25	0.51
28:11:136:ILE:HG22	28:11:137:PRO:HD2	1.92	0.51
2:12:175:ARG:O	2:12:179:LYS:HB2	2.09	0.51
1:13:1504:G:H3'	1:13:1504:G:P	2.51	0.51
26:14:1479:G:O2'	26:14:1558:A:H5'	2.10	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1802:A:N1	26:14:1822:G:H1'	2.25	0.51
26:14:1972:A:H2'	26:14:1973:G:H8	1.73	0.51
26:14:973:A:OP2	57:14:3989:HOH:O	2.19	0.51
1:1G:540:G:H2'	1:1G:541:G:O4'	2.09	0.51
26:1H:1203:G:H3'	26:1H:1204:A:H5''	1.91	0.51
26:1H:1709:U:H2'	26:1H:1710:C:C6	2.45	0.51
26:1H:185:U:H2'	26:1H:186:G:C8	2.44	0.51
26:1H:2656:U:H3	26:1H:2665:A:H2	1.58	0.51
26:1H:357:A:H2'	26:1H:358:U:C6	2.46	0.51
26:1H:816:C:OP2	57:1H:3724:HOH:O	2.19	0.51
29:29:105:THR:OG1	29:29:166:THR:HG23	2.10	0.51
57:14:3726:HOH:O	30:39:55:GLY:HA2	2.09	0.51
4:3E:141:ARG:HB2	4:3E:141:ARG:HH11	1.75	0.51
36:78:56:SER:O	36:78:61:ARG:HG3	2.11	0.51
16:7I:50:LYS:HD3	16:7I:51:VAL:H	1.75	0.51
2:12:19:HIS:HD2	2:12:20:GLU:HG2	1.75	0.51
1:13:1095:U:H2'	1:13:1096:C:C6	2.45	0.51
1:13:945:G:C2	1:13:946:A:C8	2.99	0.51
26:14:1041:C:N4	26:14:1114:G:H22	2.04	0.51
26:14:1505:C:H2'	26:14:1506:C:C6	2.46	0.51
26:14:2495:G:H5''	26:14:2496:C:OP2	2.09	0.51
26:14:2748:A:H2'	26:14:2749:A:C8	2.46	0.51
27:16:93:C:H2'	27:16:94:C:H6	1.74	0.51
2:1E:120:ALA:O	2:1E:124:SER:OG	2.16	0.51
1:1G:308:C:H2'	1:1G:309:G:H8	1.75	0.51
1:1G:765:G:N2	1:1G:813:U:OP2	2.43	0.51
26:1H:2392:A:H1'	36:78:61:ARG:HH11	1.75	0.51
26:1H:2693:A:H2'	26:1H:2694:G:C8	2.46	0.51
3:22:16:ARG:HH12	3:22:181:ASN:ND2	2.09	0.51
3:22:61:ALA:C	3:22:63:ASN:H	2.14	0.51
29:29:57:LYS:HA	29:29:58:ARG:O	2.11	0.51
3:2E:142:MET:SD	3:2E:148:GLY:HA2	2.50	0.51
30:39:178:PRO:HG2	30:39:179:GLU:OE1	2.10	0.51
30:39:28:ILE:HA	30:39:112:MET:HG2	1.91	0.51
14:5A:7:ILE:HA	14:5A:23:ARG:HE	1.74	0.51
7:6E:69:VAL:HG21	7:6E:104:LEU:HD21	1.92	0.51
40:75:80:SER:HB3	40:75:83:ILE:HG13	1.91	0.51
39:A8:67:ARG:HD3	39:A8:71:ARG:NH2	2.26	0.51
19:AA:29:ARG:HD3	19:AA:48:THR:OG1	2.10	0.51
53:L5:12:ARG:HH21	53:L5:44:PRO:HB3	1.75	0.51
50:L8:38:GLU:H	50:L8:38:GLU:CD	2.14	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1014:A:H4'	19:AI:14:HIS:CG	2.45	0.51
1:13:107:G:O6	20:BI:15:ARG:HG3	2.10	0.51
1:13:323:U:H2'	1:13:324:G:O4'	2.10	0.51
26:14:1014:U:H2'	26:14:1015:G:H8	1.75	0.51
26:14:128:C:H2'	26:14:129:C:H6	1.76	0.51
26:14:1810:A:H2'	26:14:1811:G:O4'	2.11	0.51
26:14:2320:A:N6	26:14:2333:A:H2'	2.26	0.51
26:14:2682:U:O2'	29:29:13:ARG:HG2	2.10	0.51
26:14:2754:U:H5'	26:14:2755:C:OP2	2.10	0.51
26:14:43:G:H1'	26:14:438:G:N2	2.25	0.51
1:1G:1095:U:H2'	1:1G:1096:C:O4'	2.11	0.51
1:1G:41:G:H2'	1:1G:42:G:C8	2.46	0.51
26:1H:1007:C:OP1	34:58:35:ARG:NH1	2.44	0.51
26:1H:1396:U:H2'	26:1H:1396:U:O2	2.08	0.51
26:1H:2815:C:O2'	52:N8:42:PRO:HG2	2.11	0.51
26:1H:2849:U:H4'	26:1H:2868:A:C2	2.46	0.51
26:1H:796:C:H2'	26:1H:797:C:H6	1.73	0.51
22:1L:10:G:H2'	22:1L:11:U:C6	2.46	0.51
29:21:119:ARG:HB3	29:21:120:TRP:CD1	2.45	0.51
30:31:103:LYS:HG2	30:31:106:ARG:HH21	1.75	0.51
26:1H:321:G:OP2	30:31:135:LYS:HG3	2.11	0.51
12:3A:100:ILE:HG22	12:3A:101:VAL:N	2.24	0.51
24:3L:2:C:H2'	24:3L:3:C:C6	2.45	0.51
32:51:85:LYS:HE3	32:51:142:GLY:HA2	1.93	0.51
7:62:15:ASP:OD1	7:62:44:TYR:OH	2.29	0.51
33:69:109:ILE:H	33:69:109:ILE:HD13	1.75	0.51
1:1G:581:G:OP1	15:6A:61:GLY:HA3	2.11	0.51
8:7E:29:SER:OG	8:7E:32:LYS:HE3	2.10	0.51
17:8A:99:SER:OG	17:8A:100:LYS:N	2.44	0.51
19:AI:41:VAL:HG11	19:AI:67:VAL:HA	1.93	0.51
45:G8:40:GLU:HA	45:G8:42:VAL:N	2.26	0.51
46:H8:48:PHE:HE1	46:H8:71:VAL:HG11	1.75	0.51
51:I5:16:CYS:HA	51:I5:33:VAL:HG22	1.91	0.51
28:11:77:ALA:HB2	28:11:97:TYR:CD1	2.45	0.51
1:13:1171:G:H2'	1:13:1172:C:C6	2.45	0.51
1:13:1226:C:H4'	19:AI:80:TYR:OH	2.09	0.51
1:13:580:U:H2'	1:13:581:G:O4'	2.11	0.51
1:13:736:C:H2'	1:13:737:A:C8	2.45	0.51
26:14:2082:A:H3'	26:14:2083:G:H8	1.74	0.51
26:14:929:G:O5'	26:14:929:G:H8	1.94	0.51
10:1A:78:ASN:OD1	10:1A:80:LYS:HB3	2.11	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1E:85:ALA:O	2:1E:90:MET:N	2.43	0.51
1:1G:1134:G:N2	1:1G:1140:C:N3	2.57	0.51
1:1G:142:G:H2'	1:1G:143:A:C8	2.44	0.51
1:1G:438:G:H4'	4:32:123:HIS:ND1	2.25	0.51
1:1G:628:G:H2'	1:1G:629:G:C8	2.46	0.51
26:1H:1184:G:H5'	50:L8:29:ARG:HH11	1.75	0.51
26:1H:1222:C:H2'	26:1H:1223:C:H6	1.75	0.51
26:1H:1312:U:H4'	26:1H:1313:U:O5'	2.10	0.51
26:1H:2210:G:H2'	26:1H:2211:G:C6	2.44	0.51
26:1H:736:C:O5'	26:1H:736:C:H6	1.93	0.51
27:1J:7:G:H3'	27:1J:8:U:H5''	1.92	0.51
29:29:11:MET:SD	29:29:24:THR:HG22	2.50	0.51
23:2K:21:A:H2	23:2K:57:C:H42	1.58	0.51
5:42:31:LEU:HD23	5:42:45:PHE:HB2	1.92	0.51
33:61:129:THR:HA	33:61:137:PRO:HA	1.92	0.51
33:61:47:LEU:O	33:61:51:ILE:HG13	2.10	0.51
18:9I:22:VAL:HA	18:9I:25:THR:OG1	2.10	0.51
46:D5:7:ALA:O	46:D5:8:TYR:CG	2.63	0.51
48:F5:87:PRO:O	48:F5:91:LYS:N	2.42	0.51
2:12:101:MET:HB2	2:12:102:LEU:HD12	1.93	0.51
1:13:1134:G:H1	1:13:1140:C:H42	1.58	0.51
1:13:1097:C:O2'	1:13:1169:A:N3	2.36	0.51
1:13:657:G:H4'	15:6I:28:GLN:HG2	1.93	0.51
26:14:1367:A:H5''	26:14:1368:G:OP2	2.10	0.51
26:14:1533:C:C4	26:14:1534:G:H1'	2.45	0.51
26:14:1819:A:H4'	26:14:1820:U:H5''	1.91	0.51
26:14:1973:G:H2'	26:14:1974:C:H6	1.74	0.51
26:14:2611:U:C4	52:J5:3:LYS:HG2	2.45	0.51
1:1G:1312:G:H2'	1:1G:1313:U:H6	1.75	0.51
26:1H:1034:G:H2'	26:1H:1035:U:O4'	2.10	0.51
26:1H:1423:G:H2'	26:1H:1424:G:H8	1.75	0.51
26:1H:1533:C:H3'	26:1H:1534:G:H5''	1.92	0.51
26:1H:274:G:H2'	26:1H:275:G:H4'	1.92	0.51
26:1H:654(B):C:H2'	26:1H:654(C):G:C8	2.34	0.51
26:1H:84:A:N1	26:1H:98:G:O2'	2.36	0.51
29:21:16:ARG:HG3	29:21:17:ASP:HB2	1.93	0.51
26:14:2575:C:H5'	29:29:143:ASN:O	2.11	0.51
37:45:19:GLY:O	37:45:98:LYS:HB3	2.10	0.51
38:55:8:ARG:HH11	38:55:39:PRO:HB3	1.74	0.51
38:98:44:LEU:HD22	38:98:48:VAL:HG13	1.93	0.51
38:98:76:VAL:HG13	38:98:80:PHE:CE2	2.45	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:AA:29:ARG:HB2	19:AA:30:LEU:HD22	1.92	0.51
40:B8:112:ARG:HA	40:B8:115:ARG:CZ	2.40	0.51
40:B8:90:GLN:HG3	40:B8:91:ARG:N	2.25	0.51
48:F5:52:ARG:HA	48:F5:56:GLN:O	2.11	0.51
48:F5:8:SER:OG	48:F5:10:LYS:HG3	2.11	0.51
45:G8:8:LYS:O	45:G8:11:ASP:HB2	2.09	0.51
46:H8:164:ALA:O	46:H8:165:VAL:HG22	2.10	0.51
1:13:411:A:N9	1:13:413:G:H1'	2.26	0.51
26:14:1159:U:O2'	26:14:1160:G:H5'	2.11	0.51
26:14:1592:C:H2'	26:14:1593:G:C8	2.45	0.51
26:14:2124:G:H1	26:14:2173:A:N6	2.09	0.51
26:14:2715:C:H2'	26:14:2716:U:C6	2.45	0.51
26:14:2776:A:OP1	26:14:2776:A:H3'	2.10	0.51
26:14:882:G:H22	26:14:894:C:N4	2.09	0.51
2:1E:93:VAL:HG21	2:1E:97:TRP:HD1	1.76	0.51
1:1G:1323:G:H2'	1:1G:1324:A:C8	2.46	0.51
26:1H:2024:G:H2'	26:1H:2025:C:H6	1.75	0.51
26:1H:2533:A:OP1	26:1H:2665:A:H1'	2.10	0.51
24:3L:44:G:N3	24:3L:44:G:H2'	2.26	0.51
31:41:112:PRO:HG2	51:M8:36:CYS:HB2	1.92	0.51
13:4A:79:LYS:O	13:4A:82:MET:N	2.41	0.51
32:59:103:LEU:HD23	32:59:103:LEU:H	1.75	0.51
6:5E:39:LYS:H	6:5E:64:GLN:CB	2.23	0.51
15:6A:16:ALA:HB1	15:6A:21:ASP:HB3	1.91	0.51
1:13:1374:A:O2'	7:6E:28:ASN:HB3	2.11	0.51
36:78:17:LYS:HE2	36:78:27:HIS:CD2	2.45	0.51
37:88:66:ILE:HG22	37:88:67:ARG:N	2.24	0.51
38:98:117:VAL:O	38:98:118:GLU:HB2	2.09	0.51
41:C8:92:ARG:NH2	42:D8:11:GLN:H	2.09	0.51
48:F5:92:LYS:O	48:F5:94:LEU:N	2.43	0.51
46:H8:81:ARG:HG3	46:H8:81:ARG:O	2.11	0.51
49:K8:48:HIS:N	49:K8:50:ILE:HD11	2.26	0.51
54:Q8:39:LYS:HA	54:Q8:42:ARG:HD3	1.92	0.51
1:13:404:U:H2'	1:13:405:U:H6	1.76	0.51
26:14:296:C:H2'	26:14:297:C:H6	1.75	0.51
26:14:399:G:OP2	57:14:3700:HOH:O	2.18	0.51
26:14:675:A:C8	26:14:804:A:C6	2.98	0.51
21:1B:9:ARG:O	21:1B:13:ILE:HG13	2.10	0.51
1:1G:1512:U:H2'	1:1G:1513:A:C8	2.46	0.51
1:1G:551:U:H2'	1:1G:552:U:C6	2.46	0.51
26:1H:2432:A:C4	48:J8:33:LYS:HG2	2.46	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:270(E):G:H1	26:1H:270(U):C:H42	1.59	0.51
26:1H:805:G:OP2	36:78:41:ARG:HG2	2.11	0.51
36:35:49:ARG:NH1	54:M5:58:ILE:HG23	2.25	0.51
31:41:20:ILE:HG23	31:41:25:TYR:HD2	1.76	0.51
32:51:90:LYS:HG3	32:51:163:TYR:CD1	2.46	0.51
34:58:78:TYR:N	34:58:78:TYR:CD1	2.79	0.51
33:61:93:THR:HA	33:61:119:PRO:HB3	1.92	0.51
33:61:39:ALA:HB1	33:61:44:LEU:HD13	1.92	0.51
33:61:68:LEU:HA	33:61:71:ILE:CG2	2.41	0.51
15:6I:12:ILE:HG12	15:6I:31:LEU:HD11	1.91	0.51
36:78:98:GLU:O	36:78:101:VAL:HG13	2.11	0.51
16:7I:8:ARG:O	16:7I:9:PHE:HD1	1.93	0.51
42:95:39:LEU:HD12	42:95:39:LEU:H	1.75	0.51
11:2A:109:VAL:HG13	18:9A:86:VAL:HG13	1.92	0.51
20:BA:37:SER:O	20:BA:41:ILE:HG23	2.11	0.51
26:1H:535:C:O3'	41:C8:53:ARG:NH1	2.43	0.51
46:D5:30:ASN:HD21	46:D5:33:LEU:HB3	1.75	0.51
45:G8:52:SER:O	45:G8:56:PRO:HA	2.11	0.51
1:13:266:G:H5''	1:13:267:C:H5	1.76	0.51
26:14:1153:C:H2'	26:14:1154:G:O4'	2.11	0.51
26:14:1485:G:H1	26:14:1504:C:N4	2.07	0.51
28:19:65:ILE:HD11	28:19:67:PHE:CE1	2.45	0.51
1:1G:1199:U:H4'	10:1A:54:PHE:CE1	2.46	0.51
2:1E:74:LYS:O	2:1E:78:GLN:HB2	2.11	0.51
1:1G:722:A:H2'	1:1G:722:A:N3	2.26	0.51
26:1H:1266:G:O4'	43:E8:15:ARG:NH2	2.44	0.51
26:1H:1523:U:O5'	26:1H:1523:U:H6	1.94	0.51
26:1H:1528:A:H2	26:1H:1542:G:C2	2.29	0.51
4:3E:99:SER:HB3	4:3E:139:ARG:HG3	1.93	0.51
12:3I:110:VAL:CG2	12:3I:120:TYR:HB3	2.40	0.51
7:62:50:ILE:HB	7:62:58:PRO:HG3	1.91	0.51
41:85:92:ARG:HD2	41:85:95:LEU:HD12	1.92	0.51
17:8A:55:ASP:OD1	17:8A:55:ASP:N	2.44	0.51
19:AI:51:VAL:O	19:AI:57:HIS:HA	2.11	0.51
19:AI:50:ALA:HB1	19:AI:57:HIS:HB3	1.93	0.51
26:14:483:A:H5'	45:C5:49:VAL:HG22	1.93	0.51
26:1H:1161:C:H4'	42:D8:8:GLY:HA2	1.92	0.51
26:14:469:G:O6	53:L5:39:ARG:NH1	2.44	0.51
54:Q8:23:VAL:HG22	54:Q8:24:ALA:N	2.26	0.51
2:12:97:TRP:HZ3	2:12:99:GLY:HA2	1.75	0.51
1:13:963:G:H5'	57:13:1840:HOH:O	2.12	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1033:U:H1'	26:14:2750:A:N1	2.25	0.51
26:14:1187:G:H8	26:14:1187:G:O5'	1.94	0.51
26:14:1784:A:H5''	57:14:3549:HOH:O	2.09	0.51
26:14:500:G:N2	26:14:502:A:H3'	2.26	0.51
26:14:1007:C:H5''	34:15:35:ARG:NH1	2.26	0.51
1:1G:323:U:H5'	20:BA:23:ARG:HB2	1.92	0.51
1:1G:980:C:H5''	1:1G:981:U:C5	2.46	0.51
26:1H:2036:C:N4	57:1H:3659:HOH:O	2.44	0.51
26:1H:589:C:H2'	26:1H:590:A:C8	2.46	0.51
1:13:1366:C:O2'	10:1I:60:ARG:NH2	2.44	0.51
29:21:35:GLN:HG3	29:21:36:ARG:N	2.26	0.51
3:22:36:ASP:HA	3:22:39:ILE:HD12	1.93	0.51
3:22:9:GLY:HA2	3:22:12:LEU:HG	1.93	0.51
4:32:148:VAL:O	4:32:152:SER:OG	2.16	0.51
6:5E:36:ARG:HH21	6:5E:38:GLU:HG2	1.75	0.51
10:1I:49:VAL:HG23	14:5I:41:ARG:HB2	1.93	0.51
7:6E:15:ASP:HB3	7:6E:20:ASP:N	2.25	0.51
40:75:113:LYS:O	40:75:114:LEU:HD23	2.11	0.51
40:75:8:LYS:HZ2	40:75:8:LYS:HA	1.76	0.51
9:82:24:GLY:HA2	9:82:59:PHE:O	2.10	0.51
38:98:10:LEU:O	38:98:12:ARG:N	2.44	0.51
42:D8:69:LYS:HB2	42:D8:88:ARG:HG2	1.91	0.51
49:K8:58:ALA:O	49:K8:62:THR:HG22	2.11	0.51
1:13:1064:G:OP1	1:13:1386:G:H4'	2.11	0.50
1:13:446:G:H2'	1:13:447:G:O4'	2.12	0.50
26:14:1159:U:H2'	26:14:1160:G:H8	1.75	0.50
26:14:1379:A:H4'	26:14:1380:G:OP2	2.11	0.50
26:14:1450:C:H2'	26:14:1451:C:C6	2.45	0.50
26:14:270(E):G:H1	26:14:270(U):C:H42	1.58	0.50
26:14:2850:A:C2	26:14:2851:A:C4	2.99	0.50
1:1G:1203:C:H2'	1:1G:1204:A:O4'	2.11	0.50
1:1G:136:C:O2'	16:7A:65:GLN:OE1	2.29	0.50
1:1G:999:U:O4	1:1G:1000:A:N6	2.42	0.50
26:1H:1265:A:H3'	52:N8:19:ARG:NH1	2.25	0.50
26:1H:1602:U:H5	57:1H:3792:HOH:O	1.93	0.50
26:1H:2146:C:H4'	26:1H:2147:G:N7	2.26	0.50
26:1H:1999:C:H5''	26:1H:2723:C:O2'	2.11	0.50
26:1H:2789:C:H1'	26:1H:2892:A:H2	1.76	0.50
26:1H:2791:C:N4	26:1H:2893:G:O6	2.44	0.50
26:1H:34:C:OP2	26:1H:34:C:H6	1.93	0.50
26:1H:844:C:H3'	26:1H:845:G:C8	2.46	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
27:1J:13:A:H2'	27:1J:70:C:O2'	2.11	0.50
29:21:45:THR:O	29:21:83:ASP:N	2.45	0.50
4:32:108:LEU:HD13	4:32:174:LEU:HB3	1.93	0.50
4:32:61:LYS:HA	4:32:203:VAL:HG22	1.93	0.50
32:51:4:ILE:HG13	32:51:6:ARG:NE	2.26	0.50
3:2E:13:GLY:HA3	14:5I:57:ARG:HD2	1.93	0.50
40:B8:100:TYR:HD1	40:B8:103:ARG:NH2	2.08	0.50
47:I8:50:ASN:HB2	47:I8:81:VAL:O	2.10	0.50
2:12:131:PRO:HG2	2:12:134:GLU:HB2	1.93	0.50
2:12:19:HIS:CD2	2:12:20:GLU:HG2	2.46	0.50
1:13:1079:G:H2'	1:13:1080:A:C8	2.46	0.50
1:13:1189:C:OP1	10:1I:51:ARG:NH2	2.44	0.50
26:14:1850:G:H2'	26:14:1851:U:O4'	2.11	0.50
26:14:2074:U:P	57:14:3503:HOH:O	2.69	0.50
26:14:2540:C:H2'	26:14:2541:A:O4'	2.11	0.50
26:14:900:A:H3'	26:14:901:A:H8	1.76	0.50
34:15:28:THR:HG22	34:15:29:LYS:HE3	1.93	0.50
27:16:3:C:H2'	27:16:4:C:H6	1.75	0.50
28:19:181:GLU:HG3	28:19:272:ALA:CB	2.40	0.50
2:1E:54:THR:HG21	2:1E:201:ILE:HD11	1.93	0.50
2:1E:226:ARG:HG3	2:1E:227:GLY:H	1.76	0.50
1:1G:197:A:H1'	1:1G:198:G:O4'	2.10	0.50
1:1G:373:A:C2	1:1G:374:A:C8	2.99	0.50
26:1H:1328:G:H2'	26:1H:1330:C:C5	2.46	0.50
26:1H:1705:G:C2'	26:1H:1706:U:H5'	2.42	0.50
26:1H:818:G:H4'	26:1H:838:C:O3'	2.10	0.50
35:25:111:PHE:HB3	35:25:114:ILE:HG13	1.92	0.50
4:32:112:VAL:HG12	4:32:116:GLN:OE1	2.11	0.50
4:32:199:ASN:HB3	4:32:202:LEU:HG	1.92	0.50
4:3E:111:ALA:HB2	4:3E:120:LEU:HD12	1.93	0.50
5:4E:148:VAL:O	5:4E:151:LEU:HB2	2.10	0.50
1:1G:1538:C:H42	25:4L:34:G:H1	1.60	0.50
32:51:54:ARG:HD3	32:51:65:HIS:ND1	2.26	0.50
32:51:4:ILE:HG21	32:51:6:ARG:NH1	2.25	0.50
34:58:137:LYS:HG3	34:58:138:LEU:O	2.11	0.50
33:69:45:LYS:O	33:69:48:GLU:HB3	2.12	0.50
33:69:76:THR:HG21	33:69:140:LEU:HD12	1.93	0.50
41:85:97:ASP:OD1	41:85:98:LEU:N	2.43	0.50
26:1H:2293:C:H5''	39:A8:89:ARG:HH22	1.75	0.50
20:BA:49:ALA:HB2	20:BA:92:LEU:HD22	1.93	0.50
45:G8:97:ARG:N	45:G8:97:ARG:HD2	2.26	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:K8:59:ARG:O	49:K8:62:THR:HG23	2.11	0.50
28:11:92:ILE:HD12	28:11:104:TYR:CE1	2.46	0.50
2:12:77:ALA:HB2	2:12:211:ILE:HD13	1.93	0.50
1:13:826:C:H2'	1:13:827:U:O2	2.12	0.50
26:14:1126:A:H4'	26:14:1127:A:O5'	2.11	0.50
26:14:1516:U:H2'	26:14:1517:G:H8	1.76	0.50
26:14:34:C:O2'	26:14:35:G:OP1	2.25	0.50
26:14:890:A:H2'	26:14:892:G:H8	1.76	0.50
27:16:11:C:O5'	27:16:12:C:H5	1.94	0.50
2:1E:163:PHE:CD1	2:1E:185:ILE:HG13	2.46	0.50
26:1H:1124:C:H2'	26:1H:1125:G:O4'	2.12	0.50
26:1H:1592:C:H2'	26:1H:1593:G:H8	1.76	0.50
26:1H:172:C:H2'	26:1H:173:G:C8	2.46	0.50
26:1H:528:A:C2	26:1H:2043:C:H4'	2.46	0.50
10:1I:57:LYS:HG3	10:1I:60:ARG:HH12	1.77	0.50
22:1L:48:C:HO2'	22:1L:49:C:P	2.32	0.50
29:21:64:LYS:O	29:21:70:ALA:HB2	2.11	0.50
3:22:134:ILE:HD12	3:22:151:VAL:HG11	1.92	0.50
4:32:126:ILE:HG22	4:32:127:THR:H	1.77	0.50
26:14:805:G:OP2	36:35:41:ARG:HG2	2.12	0.50
30:39:25:PRO:C	30:39:27:GLU:N	2.64	0.50
12:3A:82:VAL:HB	12:3A:105:TYR:HB3	1.94	0.50
12:3I:83:VAL:HG21	12:3I:100:ILE:HG12	1.93	0.50
24:3K:71:U:O2'	24:3K:72:C:O4'	2.29	0.50
31:41:106:LEU:HG	31:41:107:LEU:HD23	1.93	0.50
5:4E:100:VAL:O	5:4E:107:ARG:NH2	2.45	0.50
32:51:10:PRO:HG2	32:51:50:VAL:O	2.11	0.50
32:51:152:ARG:HG3	32:51:161:GLY:HA2	1.92	0.50
38:55:59:ASP:OD2	38:55:61:HIS:HB3	2.11	0.50
26:14:1111:A:H5'	32:59:3:ARG:HD3	1.93	0.50
15:6A:75:PRO:HB2	15:6A:79:ARG:CZ	2.41	0.50
36:78:113:LYS:HA	36:78:129:ALA:O	2.11	0.50
1:1G:1371:G:OP1	9:82:11:LYS:HB3	2.12	0.50
37:88:59:ARG:C	37:88:61:GLY:H	2.14	0.50
39:A8:35:ILE:HD11	39:A8:101:LEU:HD23	1.93	0.50
19:AA:64:GLU:CD	19:AA:64:GLU:H	2.12	0.50
47:E5:49:LYS:HG2	47:E5:80:HIS:ND1	2.26	0.50
46:H8:113:ALA:N	46:H8:114:GLY:HA2	2.26	0.50
26:1H:1354:A:H5''	28:11:38:LYS:HD3	1.94	0.50
1:13:1000:A:H2'	1:13:1001:G:C8	2.47	0.50
1:13:232:G:H1'	1:13:262:A:N1	2.26	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:674:G:H2'	1:13:675:A:C8	2.47	0.50
26:14:1021:A:H3'	26:14:1021:A:C8	2.46	0.50
26:14:1021:A:H8	26:14:1021:A:H3'	1.76	0.50
26:14:1040:C:H2'	26:14:1041:C:C6	2.47	0.50
26:14:1910:G:H1	26:14:1920:C:H42	1.58	0.50
26:14:2280:G:O2'	26:14:2388:A:N1	2.35	0.50
26:14:531:C:H4'	26:14:532:A:H5''	1.93	0.50
26:14:868:U:N3	26:14:869:G:N7	2.59	0.50
1:1G:1538:C:H42	25:4L:34:G:H22	1.59	0.50
26:1H:1041:C:H42	26:1H:1114:G:H1	1.57	0.50
26:1H:1996:C:O3'	57:1H:3785:HOH:O	2.19	0.50
26:1H:2157:G:O2'	26:1H:2158:A:O5'	2.29	0.50
26:1H:270(L):U:C2	33:61:50:ARG:HG2	2.47	0.50
26:1H:646:A:H2'	26:1H:647:G:O4'	2.11	0.50
26:1H:821:A:H62	26:1H:972:G:H21	1.59	0.50
27:1J:101:A:OP2	27:1J:101:A:H8	1.94	0.50
4:32:65:ARG:HG3	4:32:70:ILE:HG22	1.93	0.50
34:58:43:THR:HG23	41:C8:64:ARG:HH22	1.76	0.50
34:58:90:MET:O	34:58:94:HIS:N	2.43	0.50
6:5E:5:GLU:HB3	6:5E:62:TRP:HE1	1.76	0.50
7:62:26:PHE:CE1	7:62:30:ILE:HD11	2.47	0.50
33:69:75:LEU:HD13	33:69:77:LEU:N	2.26	0.50
9:82:102:LEU:O	9:82:103:THR:OG1	2.25	0.50
41:85:65:ILE:HD11	41:85:93:LYS:HA	1.94	0.50
9:8E:47:LEU:HA	9:8E:49:PRO:HD2	1.93	0.50
45:C5:86:ARG:NE	45:C5:87:LYS:O	2.40	0.50
26:1H:125:G:C6	53:P8:10:ARG:HG3	2.46	0.50
28:11:70:TRP:CH2	28:11:150:LYS:HA	2.47	0.50
1:13:1148:U:H2'	1:13:1149:C:O4'	2.11	0.50
1:13:1171:G:O2'	1:13:1172:C:H5'	2.11	0.50
1:13:1366:C:H2'	1:13:1367:C:H6	1.75	0.50
26:14:1537:C:H2'	26:14:1538:G:C8	2.46	0.50
26:14:1542:G:H3'	26:14:1543:A:H5''	1.93	0.50
26:14:234:C:H2'	26:14:235:U:C6	2.44	0.50
26:14:2693:A:H2'	26:14:2694:G:C8	2.47	0.50
26:14:752:A:OP1	53:L5:3:ARG:NH2	2.40	0.50
26:14:96:G:H4'	49:G5:48:HIS:CD2	2.47	0.50
34:15:133:GLN:HG3	34:15:135:PRO:HG3	1.93	0.50
34:15:21:LYS:O	34:15:60:ILE:HG13	2.11	0.50
34:15:23:LEU:HA	34:15:60:ILE:HD11	1.94	0.50
27:16:88:C:H2'	27:16:89:G:O4'	2.12	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1E:58:ILE:O	2:1E:61:LEU:N	2.45	0.50
1:13:1327:C:P	21:1F:12:LYS:HZ1	2.35	0.50
1:1G:1387:G:H2'	1:1G:1388:C:C6	2.47	0.50
1:1G:365:U:H5''	1:1G:366:C:OP1	2.11	0.50
1:1G:422:C:O2'	1:1G:423:G:C2	2.65	0.50
26:1H:1209:G:H21	26:1H:1210:A:H62	1.58	0.50
26:1H:1264:G:H3'	26:1H:1265:A:H5''	1.93	0.50
26:1H:1606:G:OP1	57:1H:4102:HOH:O	2.19	0.50
26:1H:2336:A:H61	47:18:43:THR:HB	1.76	0.50
26:1H:527:C:H4'	26:1H:528:A:O5'	2.12	0.50
26:1H:566:U:OP1	36:78:29:LYS:NZ	2.38	0.50
26:1H:972:G:OP2	26:1H:973:A:O2'	2.27	0.50
22:1K:14:A:H61	22:1K:21:A:H2	1.59	0.50
29:21:105:THR:HG22	29:21:106:GLY:N	2.27	0.50
24:3L:84:A:H2'	24:3L:85:C:H6	1.77	0.50
6:52:11:ASN:HB3	6:52:14:LEU:CD1	2.42	0.50
38:55:29:LEU:HB3	38:55:75:LEU:HD21	1.94	0.50
26:14:2820:A:C6	38:55:4:LEU:HD11	2.47	0.50
7:6E:12:LEU:HD21	7:6E:28:ASN:ND2	2.27	0.50
18:9A:74:ARG:HB3	18:9A:81:PHE:CE1	2.46	0.50
52:J5:20:ARG:HA	52:J5:23:HIS:ND1	2.27	0.50
2:12:48:MET:HA	2:12:51:LEU:HB2	1.94	0.50
1:13:1349:A:H2'	1:13:1350:A:C8	2.47	0.50
26:14:1640:C:H2'	26:14:1641:A:C8	2.46	0.50
26:14:1731:G:N3	26:14:1731:G:H5''	2.27	0.50
26:14:2037:G:H2'	26:14:2038:G:C8	2.46	0.50
26:14:2495:G:H5'	37:45:82:ARG:HG2	1.93	0.50
26:14:2875:C:O2'	40:75:1:MET:N	2.44	0.50
26:14:358:U:H2'	26:14:359:A:C8	2.47	0.50
21:1F:5:ASP:O	21:1F:11:GLY:HA3	2.12	0.50
1:1G:1239:A:H4'	1:1G:1240:U:C5'	2.41	0.50
1:1G:276:G:O3'	17:8A:68:ARG:NH1	2.45	0.50
1:1G:373:A:N3	1:1G:374:A:C8	2.80	0.50
1:1G:518:C:H5''	1:1G:519:C:C6	2.47	0.50
26:1H:1132:A:O2'	26:1H:1133:U:H5'	2.11	0.50
26:1H:11:G:H2'	26:1H:12:U:H5'	1.94	0.50
26:1H:1729:A:O2'	26:1H:1730:U:OP1	2.27	0.50
26:1H:2210:G:H5'	26:1H:2211:G:C5	2.47	0.50
26:1H:467:G:O5'	26:1H:467:G:H8	1.94	0.50
9:8E:128:ARG:NH2	23:2K:36:A:OP2	2.41	0.50
4:32:26:CYS:HA	4:32:31:CYS:CB	2.31	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:39:185:ASP:OD1	30:39:188:ARG:NH2	2.33	0.50
24:3L:26:C:H2'	24:3L:27:G:O4'	2.12	0.50
13:4A:3:ARG:HH21	31:49:113:ARG:HH21	1.59	0.50
5:4E:82:VAL:HG21	5:4E:138:ALA:HA	1.94	0.50
38:55:106:GLY:O	38:55:107:ASP:HB3	2.10	0.50
26:14:2880:C:O2'	38:55:90:ARG:HD3	2.12	0.50
34:58:42:TRP:HA	34:58:48:MET:HE3	1.92	0.50
26:1H:1022:G:N7	34:58:66:LYS:NZ	2.59	0.50
32:59:20:ALA:HB3	32:59:24:VAL:HA	1.93	0.50
7:62:113:GLU:O	7:62:119:ARG:HD3	2.11	0.50
33:69:112:LYS:HA	33:69:114:LEU:H	1.77	0.50
16:7A:53:VAL:O	16:7A:57:ARG:HG3	2.12	0.50
26:1H:873:G:H1'	37:88:29:PHE:CE2	2.46	0.50
37:88:82:ARG:HE	47:18:4:LYS:NZ	2.10	0.50
20:BA:40:ALA:O	20:BA:43:LEU:N	2.44	0.50
20:BI:50:GLU:HG3	20:BI:100:ILE:HG12	1.92	0.50
45:G8:100:ALA:HB1	45:G8:101:LYS:CB	2.42	0.50
46:H8:4:ARG:HA	46:H8:58:VAL:HG22	1.93	0.50
1:13:1505:G:H4'	1:13:1506:U:H5''	1.92	0.50
1:13:1504:G:OP1	1:13:1507:A:H4'	2.12	0.50
1:13:376:G:H5''	16:7I:5:ARG:HB2	1.93	0.50
1:13:73:G:H2'	1:13:74:C:O4'	2.11	0.50
26:14:142:G:H2'	26:14:143:C:H6	1.76	0.50
26:14:1462:C:H2'	26:14:1463:C:H6	1.77	0.50
26:14:2113:U:OP1	26:14:2114:A:N6	2.44	0.50
26:14:2361:A:O5'	54:M5:27:THR:OG1	2.30	0.50
26:14:2343:C:HO2'	26:14:2373:G:HO2'	1.54	0.50
26:14:2418:A:H2'	26:14:2419:U:C6	2.46	0.50
26:14:459:U:H2'	26:14:460:A:C8	2.47	0.50
28:19:12:SER:HB2	28:19:208:LYS:HB3	1.93	0.50
26:14:1800:C:OP2	28:19:183:ARG:NH2	2.44	0.50
26:1H:1100:C:H2'	26:1H:1101:U:C6	2.46	0.50
26:1H:1103:A:H3'	26:1H:1104:C:C6	2.44	0.50
26:1H:1332:G:C8	26:1H:1332:G:H5'	2.47	0.50
26:1H:2052:G:C8	29:21:141:ILE:HD11	2.47	0.50
3:22:183:ASP:HB3	3:22:202:ILE:HG13	1.93	0.50
11:2I:54:ARG:NH2	24:3K:40:G:O3'	2.45	0.50
23:2K:19:G:OP1	23:2K:19:G:H3'	2.12	0.50
12:3I:5:PRO:HG2	12:3I:10:LEU:HD21	1.94	0.50
14:5A:26:ARG:HG2	14:5A:26:ARG:O	2.12	0.50
26:1H:270(O):U:O4	33:61:52:ARG:HG2	2.12	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:78:138:LEU:CD1	36:78:144:GLU:HG3	2.42	0.50
9:8E:34:ASN:O	9:8E:38:GLN:HB2	2.11	0.50
9:8E:5:TYR:OH	9:8E:16:ARG:HG2	2.11	0.50
1:13:279:A:C5	17:8I:98:LEU:HD12	2.47	0.50
40:B8:58:ASN:ND2	40:B8:58:ASN:O	2.37	0.50
45:G8:50:ARG:HB2	45:G8:50:ARG:NH1	2.27	0.50
37:88:82:ARG:HH21	47:I8:4:LYS:NZ	2.09	0.50
28:11:70:TRP:CE2	28:11:150:LYS:HD2	2.47	0.50
1:13:95:G:H3'	1:13:96:G:C8	2.47	0.50
26:14:1019:U:HO2'	26:14:1021:A:H2	1.55	0.50
26:14:1386:C:H2'	26:14:1387:C:H6	1.77	0.50
26:14:1636:C:H2'	26:14:1637:A:C8	2.46	0.50
26:14:957:A:C6	26:14:2459:A:C8	3.00	0.50
26:14:321:G:OP1	30:39:135:LYS:NZ	2.45	0.50
26:14:539:G:H2'	26:14:540:G:C8	2.45	0.50
26:14:68:G:H2'	26:14:69:C:O4'	2.11	0.50
27:16:11:C:H3'	27:16:12:C:H6	1.76	0.50
27:16:55:U:H2'	27:16:56:G:C8	2.46	0.50
2:1E:160:ASP:O	2:1E:183:PRO:HD2	2.12	0.50
1:1G:895:G:H2'	1:1G:896:C:C6	2.46	0.50
26:1H:2630:G:H2'	26:1H:2631:G:C8	2.47	0.50
23:2K:87:A:H4'	26:1H:2602:A:C6	2.47	0.50
4:32:25:ARG:O	4:32:28:SER:N	2.45	0.50
4:3E:162:LEU:HA	4:3E:165:MET:HB2	1.93	0.50
37:45:25:ASP:HB3	37:45:102:VAL:CG2	2.34	0.50
36:78:16:ARG:HG2	36:78:17:LYS:N	2.27	0.50
26:1H:2470:G:H5'	37:88:56:ARG:NH2	2.27	0.50
17:8A:66:SER:OG	17:8A:69:LYS:HB2	2.11	0.50
39:A8:41:ASP:OD2	39:A8:44:LYS:HB2	2.12	0.50
1:1G:262:A:H5'	20:BA:74:LYS:HZ3	1.76	0.50
26:1H:1614:A:N1	43:E8:91:GLY:HA2	2.27	0.50
45:G8:85:VAL:HG23	45:G8:96:ILE:HG22	1.94	0.50
54:Q8:53:PRO:HB3	54:Q8:56:GLU:HG3	1.94	0.50
28:11:109:ASP:HB2	28:11:197:GLY:CA	2.41	0.50
1:13:563:A:N7	1:13:567:G:H1'	2.26	0.50
26:14:1439:A:H2'	26:14:1440:G:O4'	2.12	0.50
26:14:2147:G:H2'	26:14:2148:G:H4'	1.94	0.50
26:14:2720:U:N3	26:14:2873:A:C2	2.78	0.50
27:16:79:C:H6	27:16:79:C:O5'	1.95	0.50
1:1G:1037:C:H2'	1:1G:1038:C:C6	2.47	0.50
1:1G:1260:C:H6	1:1G:1260:C:H3'	1.77	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1062:G:N2	26:1H:1088:A:N1	2.60	0.50
26:1H:2514:U:H3	26:1H:2570:G:H1	1.59	0.50
26:1H:2815:C:H2'	26:1H:2816:C:C6	2.46	0.50
29:21:51:PHE:O	29:21:74:PRO:HB2	2.12	0.50
35:25:69:ILE:HD12	35:25:77:ILE:O	2.12	0.50
30:31:10:PRO:O	30:31:124:LEU:HD12	2.11	0.50
4:3E:206:PHE:HD1	4:3E:207:TYR:CD1	2.30	0.50
37:45:69:PHE:CD1	37:45:70:PRO:HD2	2.47	0.50
32:51:6:ARG:HA	32:51:66:GLY:HA2	1.94	0.50
33:61:10:GLU:O	33:61:10:GLU:HG3	2.12	0.50
35:68:25:LEU:HD12	35:68:38:VAL:O	2.12	0.50
8:72:20:TYR:HA	8:72:65:TYR:CE2	2.47	0.50
1:1G:1443:G:N2	40:75:119:LYS:HB2	2.27	0.50
19:AI:39:THR:HG22	19:AI:40:ILE:H	1.76	0.50
37:45:27:VAL:HG11	46:D5:81:ARG:NH2	2.26	0.50
42:D8:38:LEU:HD21	42:D8:40:LEU:O	2.12	0.50
49:K8:64:LEU:O	49:K8:68:ARG:HG3	2.12	0.50
49:K8:65:ASN:O	49:K8:69:ARG:HG3	2.12	0.50
2:12:53:ARG:HH12	2:12:199:TYR:HA	1.77	0.49
1:13:1227:A:OP2	13:4I:111:LYS:HE3	2.11	0.49
1:13:820:U:H4'	1:13:821:G:OP2	2.12	0.49
26:14:1582:C:HO2'	26:14:1586:A:H8	1.60	0.49
23:2L:13:G:H4'	26:14:1924:C:H4'	1.94	0.49
26:14:2638:G:OP2	29:29:82:ARG:NH2	2.45	0.49
26:14:271(A):C:O2'	26:14:271(B):G:H5'	2.12	0.49
26:14:339:U:H6	26:14:339:U:O5'	1.94	0.49
34:15:4:TYR:CD2	41:85:100:VAL:HG11	2.47	0.49
2:1E:80:ILE:HD11	2:1E:208:ILE:HG23	1.94	0.49
1:1G:1058:G:H2'	1:1G:1059:C:C6	2.47	0.49
1:1G:1135:U:O2'	1:1G:1138:G:O6	2.20	0.49
1:1G:1320:C:H2'	1:1G:1321:C:O4'	2.12	0.49
1:1G:430:A:OP2	4:32:8:VAL:HG23	2.11	0.49
1:1G:804:U:H5''	1:1G:805:C:OP2	2.12	0.49
26:1H:1652:A:OP1	38:98:8:ARG:NH1	2.43	0.49
26:1H:207:A:H2'	26:1H:208:C:O4'	2.12	0.49
26:1H:2301:C:H2'	26:1H:2302:G:H8	1.77	0.49
26:1H:2734:A:H5'	26:1H:2735:G:OP2	2.12	0.49
26:1H:356:G:H2'	26:1H:357:A:H8	1.75	0.49
26:1H:911:A:H2'	37:88:9:TYR:OH	2.12	0.49
27:1J:89(A):A:H5'	27:1J:90:C:OP2	2.12	0.49
31:41:122:PRO:HB3	31:41:180:PHE:CD2	2.47	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:41:47:LYS:HD3	31:41:81:LYS:HB2	1.94	0.49
5:42:24:ARG:HH11	5:42:24:ARG:HB3	1.77	0.49
6:5E:5:GLU:HB3	6:5E:62:TRP:NE1	2.27	0.49
33:61:75:LEU:HB3	33:61:105:HIS:CD2	2.47	0.49
7:62:111:ARG:NH2	7:62:122:HIS:HB3	2.27	0.49
39:65:106:ARG:O	39:65:106:ARG:HD2	2.12	0.49
16:7A:36:ILE:HD12	16:7A:56:ALA:HB2	1.94	0.49
40:B8:74:ARG:HD3	40:B8:76:PHE:CZ	2.47	0.49
20:BA:72:LEU:O	20:BA:73:HIS:HB2	2.11	0.49
20:BA:73:HIS:CB	20:BA:74:LYS:HZ2	2.25	0.49
46:H8:73:GLN:HB2	46:H8:87:ASP:HB2	1.94	0.49
2:12:60:ASP:O	2:12:64:ARG:N	2.43	0.49
1:13:1190:G:OP1	3:2E:5:ILE:N	2.41	0.49
1:13:1348:U:N3	1:13:1374:A:H2	2.04	0.49
1:13:45:U:H2'	1:13:46:G:H8	1.76	0.49
1:13:741:G:H2'	1:13:742:G:O4'	2.11	0.49
26:14:2030:A:H4'	26:14:2031:A:C8	2.47	0.49
26:14:2331:G:O2'	26:14:2336:A:N1	2.42	0.49
26:14:2395:C:H2'	26:14:2396:G:O4'	2.12	0.49
26:14:2636:U:O2'	29:29:44:TYR:OH	2.20	0.49
26:14:579:G:H2'	26:14:580:C:H6	1.76	0.49
2:1E:17:PHE:HB2	2:1E:42:ILE:HG22	1.93	0.49
1:1G:1127:G:N3	1:1G:1127:G:H2'	2.26	0.49
1:1G:1131:G:H2'	1:1G:1132:C:H6	1.75	0.49
1:1G:300:A:H1'	1:1G:565:U:O2	2.11	0.49
26:1H:1058:U:N3	26:1H:1059:G:N7	2.61	0.49
26:1H:1590:U:H2'	26:1H:1591:G:C8	2.47	0.49
26:1H:620:G:H4'	26:1H:621:A:C5'	2.41	0.49
26:1H:900:A:H3'	26:1H:901:A:H8	1.77	0.49
29:21:147:PRO:HB2	29:21:149:ARG:HG3	1.94	0.49
23:2L:73:C:H2'	23:2L:74:C:H6	1.75	0.49
30:31:198:ALA:O	30:31:201:VAL:N	2.45	0.49
4:32:190:ASP:OD1	4:32:190:ASP:N	2.45	0.49
4:3E:155:LEU:HB3	4:3E:158:ILE:HG13	1.93	0.49
12:3I:6:THR:OG1	12:3I:9:GLN:HG3	2.12	0.49
5:42:80:ILE:HG13	8:72:104:ARG:NH2	2.27	0.49
38:55:28:LEU:HB2	38:55:34:ILE:HB	1.94	0.49
29:21:152:LYS:HD3	34:58:77:GLY:HA3	1.94	0.49
7:62:143:ARG:O	7:62:146:GLU:HB2	2.12	0.49
57:1H:4131:HOH:O	36:78:39:LYS:HB3	2.12	0.49
41:85:106:PHE:O	41:85:109:LEU:HB2	2.13	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:255:G:H1'	17:8I:16:GLN:OE1	2.11	0.49
42:95:15:GLU:HG3	42:95:16:PRO:HD2	1.93	0.49
19:AI:51:VAL:HG12	19:AI:52:TYR:H	1.77	0.49
26:14:2056:G:O3'	52:J5:8:LYS:NZ	2.45	0.49
49:K8:37:PHE:O	49:K8:41:ILE:HG22	2.13	0.49
1:13:1001:G:C6	1:13:1002:G:N7	2.81	0.49
1:13:74:C:N3	1:13:96:G:N2	2.44	0.49
26:14:1005:C:N3	26:14:1143:A:C4	2.81	0.49
26:14:1154:G:OP2	41:85:58:ARG:NH1	2.43	0.49
26:14:1176:G:H5'	26:14:1177:A:OP1	2.12	0.49
26:14:153:C:OP1	48:F5:88:LYS:NZ	2.41	0.49
26:14:1677:A:H2'	26:14:1678:G:H8	1.77	0.49
26:14:2808:U:H5''	26:14:2891:G:O6	2.12	0.49
26:14:649:G:H2'	26:14:650:C:C6	2.48	0.49
26:14:898:C:H3'	26:14:899:A:H5''	1.94	0.49
1:1G:1172:C:H2'	1:1G:1173:G:H8	1.76	0.49
26:1H:1170:G:N2	26:1H:1180:C:O2	2.45	0.49
26:1H:140:A:H8	26:1H:1408:C:HO2'	1.56	0.49
26:1H:18:C:O3'	41:C8:23:GLY:HA2	2.13	0.49
26:1H:2880:C:O2'	38:98:90:ARG:NH1	2.39	0.49
26:1H:456:C:H3'	44:F8:68:ARG:NH2	2.27	0.49
3:2E:5:ILE:HG22	10:1I:51:ARG:HH12	1.77	0.49
22:1K:15:A:H5''	22:1K:16:U:H5''	1.93	0.49
22:1K:68:A:H2'	22:1K:69:A:H5'	1.93	0.49
22:1L:7:G:H3'	22:1L:8:U:C5'	2.41	0.49
29:21:34:VAL:HG22	29:21:48:GLN:HB3	1.92	0.49
29:21:36:ARG:NH1	29:21:85:ASN:OD1	2.45	0.49
29:29:134:ILE:O	29:29:134:ILE:HD12	2.13	0.49
23:2L:42:U:O4	23:2L:43:A:N6	2.44	0.49
24:3K:83:U:C2'	24:3K:84:A:H5''	2.36	0.49
37:45:25:ASP:HB3	37:45:102:VAL:H	1.76	0.49
13:4A:57:ARG:HH12	51:I5:17:GLY:HA3	1.78	0.49
13:4I:9:ILE:HG22	13:4I:10:PRO:O	2.11	0.49
25:4K:34:G:H2'	25:4K:35:A:C8	2.47	0.49
14:5I:4:LYS:HA	14:5I:7:ILE:HG12	1.93	0.49
7:6E:11:GLN:OE1	7:6E:12:LEU:N	2.46	0.49
7:6E:65:ALA:O	7:6E:69:VAL:HG23	2.12	0.49
36:78:17:LYS:HE2	36:78:27:HIS:HD2	1.77	0.49
37:88:34:LEU:HD23	37:88:104:PHE:HD2	1.78	0.49
19:AA:27:GLU:HG2	19:AA:47:HIS:NE2	2.27	0.49
26:1H:1188:U:H5'	42:D8:79:VAL:HG22	1.93	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
46:H8:53:ILE:HG22	46:H8:71:VAL:HG13	1.93	0.49
1:13:1218:C:H2'	1:13:1219:U:C6	2.48	0.49
1:13:113:G:O2'	1:13:354:G:H5'	2.12	0.49
26:14:1486:A:H2'	26:14:1487:G:H8	1.78	0.49
26:14:34:C:HO2'	26:14:35:G:P	2.35	0.49
27:16:0:A:H62	27:16:119:A:N6	2.02	0.49
27:16:67:G:O5'	27:16:67:G:H8	1.96	0.49
28:19:71:ASP:CG	28:19:103:ARG:HH12	2.14	0.49
1:1G:1466:C:H2'	1:1G:1467:G:O4'	2.12	0.49
1:1G:1522:U:H2'	1:1G:1523:G:C8	2.47	0.49
1:1G:491:G:H2'	1:1G:492:G:O4'	2.12	0.49
26:1H:1239:G:H2'	26:1H:1240:U:O4'	2.12	0.49
26:1H:1591:G:H2'	26:1H:1592:C:C6	2.47	0.49
26:1H:2428:G:O6	57:1H:3680:HOH:O	2.19	0.49
26:1H:270(T):G:C6	26:1H:270(U):C:C4	3.00	0.49
26:1H:654(H):G:O6	26:1H:654(N):G:N2	2.45	0.49
26:1H:963:U:H5''	57:1H:3699:HOH:O	2.11	0.49
11:2I:19:ALA:O	11:2I:82:VAL:HA	2.12	0.49
23:2K:12:G:C2	23:2K:13:G:H1'	2.47	0.49
1:1G:1400:C:C2	23:2L:35:G:C2	3.00	0.49
13:4A:80:ARG:O	13:4A:84:ILE:HB	2.11	0.49
5:4E:153:LYS:HD3	5:4E:154:GLY:O	2.13	0.49
38:55:33:ARG:HG3	38:55:115:GLU:HB3	1.94	0.49
14:5A:25:VAL:O	14:5A:26:ARG:HB3	2.12	0.49
35:68:119:PRO:HB2	40:B8:68:TYR:CD2	2.47	0.49
38:98:79:LEU:HA	38:98:83:ILE:HD12	1.94	0.49
39:A8:67:ARG:HH21	39:A8:103:GLU:HG3	1.77	0.49
46:D5:75:ASN:O	46:D5:84:GLU:HG2	2.12	0.49
46:H8:48:PHE:HA	46:H8:51:ALA:HB3	1.94	0.49
28:11:65:ILE:HD11	28:11:67:PHE:CE1	2.47	0.49
1:13:1432:G:O6	57:13:1848:HOH:O	2.15	0.49
1:13:129(A):G:C2	1:13:188:U:O2'	2.65	0.49
1:13:247:G:OP2	17:8I:100:LYS:N	2.22	0.49
1:13:735:C:H2'	1:13:736:C:C6	2.43	0.49
1:13:868:C:H2'	1:13:869:G:O4'	2.12	0.49
26:14:1268:A:C2	26:14:2013:A:C4	3.00	0.49
26:14:2134:A:C2	26:14:2159:G:H1'	2.48	0.49
26:14:858:U:O2	26:14:2268:A:H2'	2.12	0.49
26:14:598:G:H1	26:14:659:C:H42	1.59	0.49
27:16:75:G:H21	46:H8:85:HIS:CE1	2.30	0.49
10:1A:4:ILE:HA	10:1A:100:THR:HG22	1.94	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1E:121:LEU:O	2:1E:139:LYS:NZ	2.37	0.49
21:1F:9:ARG:O	21:1F:13:ILE:HG13	2.12	0.49
1:1G:973:G:H1'	10:1A:55:LYS:HZ2	1.77	0.49
26:1H:1015:G:OP2	26:1H:1015:G:H8	1.96	0.49
26:1H:82:G:N2	26:1H:103:A:OP2	2.36	0.49
26:1H:1199:U:H2'	26:1H:1200:C:O4'	2.13	0.49
26:1H:1517:G:H2'	26:1H:1518:C:C6	2.48	0.49
26:1H:2630:G:H2'	26:1H:2631:G:H8	1.78	0.49
26:1H:483:A:H2'	26:1H:484:C:O4'	2.13	0.49
26:1H:723:G:H2'	26:1H:724:U:O4'	2.13	0.49
35:25:92:GLU:HG2	35:25:113:LYS:NZ	2.27	0.49
23:2L:18:OMG:H4'	23:2L:18:OMG:OP1	2.12	0.49
23:2L:24:C:H2'	23:2L:25:A:H8	1.77	0.49
30:31:9:ILE:HD11	30:31:125:LEU:N	2.20	0.49
31:49:42:GLY:O	31:49:43:LEU:HD13	2.13	0.49
15:6A:24:SER:OG	15:6A:27:VAL:HG23	2.13	0.49
7:62:16:LEU:HD13	9:82:44:VAL:HG22	1.93	0.49
39:A8:15:ARG:O	39:A8:19:LYS:HD3	2.12	0.49
19:AI:19:VAL:HG11	19:AI:44:MET:HG2	1.93	0.49
19:AI:31:ILE:HG23	19:AI:49:ILE:HG12	1.93	0.49
45:C5:39:VAL:HG23	45:C5:41:GLY:N	2.27	0.49
46:D5:39:VAL:HG21	46:D5:44:PHE:HD2	1.77	0.49
26:1H:2615:U:C2	52:N8:7:PRO:HA	2.48	0.49
26:1H:779:U:OP1	28:11:49:ILE:HG13	2.12	0.49
1:13:368:U:P	33:69:91:SER:HG	2.35	0.49
1:13:376:G:H4'	16:7I:5:ARG:NH1	2.27	0.49
1:13:413:G:H2'	1:13:428:G:H22	1.77	0.49
26:14:2199:A:OP1	48:F5:50:ARG:NH2	2.45	0.49
26:14:2238:G:OP2	57:14:3812:HOH:O	2.20	0.49
26:14:839:U:H2'	26:14:840:C:C6	2.47	0.49
26:14:921:G:C6	26:14:922:U:C4	3.01	0.49
28:19:134:ARG:O	28:19:168:ARG:NH1	2.45	0.49
1:1G:1261:A:H61	1:1G:1274:G:H1'	1.78	0.49
1:1G:145:G:N2	1:1G:177:C:O2	2.42	0.49
1:1G:995:C:H6	1:1G:995:C:O5'	1.96	0.49
26:1H:1639:U:O2'	26:1H:1640:C:H5'	2.12	0.49
26:1H:2854:G:H2'	26:1H:2855:C:C6	2.48	0.49
10:1I:33:GLN:HG2	10:1I:75:ILE:HG13	1.94	0.49
10:1I:84:GLN:O	10:1I:88:LEU:HB2	2.12	0.49
29:21:119:ARG:HG3	29:21:119:ARG:HH11	1.76	0.49
30:39:181:LEU:HD22	30:39:186:ILE:HD11	1.95	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:8:A:N7	4:3E:208:SER:HB3	2.27	0.49
12:3I:90:VAL:O	12:3I:91:LYS:HB3	2.11	0.49
32:51:6:ARG:HB3	32:51:65:HIS:CG	2.47	0.49
32:59:35:VAL:HG13	32:59:71:LEU:HD23	1.94	0.49
33:61:110:ASP:CB	33:61:112:LYS:H	2.25	0.49
8:72:100:ILE:HG21	8:72:125:ARG:HE	1.77	0.49
42:95:5:VAL:HB	42:95:37:VAL:CG1	2.42	0.49
44:B5:59:VAL:CG2	44:B5:76:ARG:HD3	2.42	0.49
28:11:112:GLN:OE1	28:11:112:GLN:N	2.46	0.49
28:11:131:LEU:HB2	28:11:136:ILE:CD1	2.43	0.49
28:11:182:LEU:HB3	28:11:271:ILE:HG13	1.94	0.49
1:13:659:U:H2'	1:13:660:G:H8	1.77	0.49
26:14:1234:U:H2'	26:14:1235:G:O4'	2.13	0.49
27:16:71:C:H2'	27:16:72:G:H8	1.76	0.49
10:1A:76:ASN:HB3	10:1A:78:ASN:HD22	1.78	0.49
1:1G:782:A:O3'	1:1G:1515:C:H4'	2.12	0.49
26:1H:1242:A:N1	36:78:4:SER:OG	2.37	0.49
26:1H:2151:G:H2'	26:1H:2152:G:C8	2.48	0.49
26:14:2773:C:OP1	29:29:166:THR:OG1	2.31	0.49
29:29:89:ASP:C	29:29:91:VAL:H	2.14	0.49
11:2I:125:PHE:N	11:2I:125:PHE:CD1	2.79	0.49
23:2L:47:C:O2'	23:2L:48:C:O5'	2.28	0.49
30:39:18:ARG:HG2	30:39:19:GLU:N	2.27	0.49
30:39:153:SER:OG	30:39:190:GLU:HB2	2.12	0.49
5:42:41:VAL:O	5:42:67:VAL:HG12	2.13	0.49
32:51:170:ARG:HE	32:51:170:ARG:HA	1.77	0.49
38:55:49:ASP:OD1	38:55:95:THR:HB	2.13	0.49
33:69:29:TYR:HD2	33:69:30:LEU:HD23	1.78	0.49
33:69:99:GLU:H	33:69:99:GLU:CD	2.15	0.49
8:72:49:GLU:O	8:72:51:VAL:HG13	2.13	0.49
36:78:136:GLU:HA	36:78:139:LYS:HE2	1.95	0.49
36:78:57:THR:HB	36:78:59:LEU:H	1.78	0.49
9:8E:118:LYS:HE2	9:8E:118:LYS:HB3	1.65	0.49
38:98:12:ARG:HD3	38:98:16:HIS:CG	2.46	0.49
38:98:57:ARG:HB3	38:98:59:ASP:OD2	2.12	0.49
49:G5:64:LEU:HD21	49:G5:68:ARG:HH11	1.78	0.49
46:H8:105:VAL:N	46:H8:139:VAL:O	2.43	0.49
1:13:234:C:H2'	1:13:235:C:H6	1.76	0.49
1:13:368:U:C6	33:69:90:GLY:HA3	2.47	0.49
1:13:429:U:O2'	4:3E:22:LYS:NZ	2.39	0.49
1:13:778:G:H1'	11:2I:119:CYS:HB3	1.93	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1045:A:H1'	26:14:1047:G:N3	2.28	0.49
26:14:1039:G:H1	26:14:1116:C:H42	1.61	0.49
26:14:1999:C:H4'	26:14:2723:C:O2	2.12	0.49
26:14:2099:U:H3	26:14:2190:G:H1	1.60	0.49
26:14:2468:G:H3'	26:14:2476:A:C2	2.48	0.49
1:1G:1084:G:H5'	1:1G:1102:A:OP2	2.13	0.49
1:1G:1386:G:C2	1:1G:1387:G:C8	3.01	0.49
1:1G:1411:C:H2'	1:1G:1412:C:C6	2.47	0.49
26:1H:1291:C:H2'	26:1H:1292:U:H6	1.78	0.49
26:1H:1728:G:H5'	26:1H:1729:A:OP2	2.12	0.49
24:3K:87:A:O2'	26:1H:2394:C:O2	2.31	0.49
26:1H:2493:U:H2'	26:1H:2494:G:O4'	2.12	0.49
26:1H:634:C:H2'	26:1H:635:C:C6	2.47	0.49
26:1H:860:U:C5	26:1H:917:A:H2	2.30	0.49
29:29:89:ASP:O	29:29:91:VAL:N	2.35	0.49
11:2I:41:THR:HG22	11:2I:42:TRP:H	1.77	0.49
12:3I:93:LEU:HB2	12:3I:96:VAL:CG1	2.43	0.49
37:45:79:LEU:H	37:45:79:LEU:HD22	1.78	0.49
1:13:1296:C:H5'	13:4I:14:ARG:HH12	1.77	0.49
34:58:121:LYS:HB2	34:58:123:TYR:HE1	1.78	0.49
33:61:75:LEU:HB3	33:61:105:HIS:HD2	1.77	0.49
33:61:124:GLY:H	33:61:142:VAL:CG2	2.25	0.49
8:72:20:TYR:CE2	8:72:75:ARG:HG2	2.47	0.49
37:88:17:LEU:HB3	37:88:39:PRO:HB2	1.95	0.49
37:88:19:GLY:O	37:88:21:THR:OG1	2.19	0.49
9:8E:43:ALA:HA	9:8E:74:ILE:HD13	1.95	0.49
26:1H:1278:A:C5'	38:98:36:THR:HG22	2.41	0.49
13:4I:84:ILE:HG13	19:AI:74:PHE:HE1	1.78	0.49
40:B8:21:GLU:OE1	40:B8:91:ARG:NH2	2.46	0.49
42:D8:65:GLY:HA3	42:D8:91:TYR:CE1	2.47	0.49
26:14:2279:G:O6	47:E5:14:ARG:HD2	2.13	0.49
47:I8:23:VAL:HA	47:I8:38:VAL:HG22	1.95	0.49
54:Q8:8:LYS:HD2	54:Q8:8:LYS:N	2.27	0.49
1:13:272:C:H2'	1:13:273:A:C8	2.48	0.49
26:14:1430:C:H2'	26:14:1431:U:H6	1.78	0.49
26:14:2056:G:N3	26:14:2056:G:H2'	2.28	0.49
26:14:2709:G:H1'	57:14:3958:HOH:O	2.12	0.49
26:14:519:U:H2'	26:14:520:G:H8	1.77	0.49
10:1A:32:ALA:HB2	10:1A:81:THR:HG21	1.94	0.49
1:1G:1127:G:N2	1:1G:1144:G:H1	2.10	0.49
26:1H:1025:G:C4	26:1H:1135:C:H1'	2.48	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1142:U:O2	26:1H:1142:U:H2'	2.13	0.49
26:1H:1509:C:H2'	26:1H:1511:A:C8	2.48	0.49
26:1H:1729:A:H8	26:1H:1730:U:H6	1.59	0.49
26:1H:2029:G:H2'	26:1H:2031:A:OP1	2.13	0.49
26:1H:270(G):C:H2'	26:1H:270(H):C:O4'	2.13	0.49
10:1I:54:PHE:CD2	10:1I:55:LYS:HG2	2.46	0.49
22:1K:56:G:H2'	22:1K:57:C:C6	2.48	0.49
29:21:96:PHE:O	29:21:175:VAL:HG11	2.12	0.49
11:2I:16:SER:O	11:2I:35:PRO:HG3	2.13	0.49
30:39:107:LYS:HE2	30:39:205:ARG:HD2	1.95	0.49
12:3I:117:ARG:HB3	12:3I:122:THR:HB	1.94	0.49
37:45:34:LEU:HD12	37:45:130:LYS:O	2.13	0.49
13:4A:81:LEU:HD23	13:4A:86:CYS:SG	2.53	0.49
13:4I:86:CYS:HB2	19:AI:73:GLU:HB3	1.93	0.49
6:52:97:PHE:O	18:9A:31:LEU:HD23	2.13	0.49
32:59:10:PRO:HD2	32:59:50:VAL:H	1.77	0.49
6:5E:92:LYS:HB2	6:5E:92:LYS:NZ	2.27	0.49
15:6A:8:LYS:O	15:6A:12:ILE:HG13	2.12	0.49
7:6E:23:VAL:O	7:6E:27:ILE:N	2.46	0.49
16:7I:77:ALA:CB	16:7I:79:VAL:HG23	2.43	0.49
38:98:30:THR:HG22	38:98:31:HIS:ND1	2.28	0.49
39:A8:10:ARG:HG3	39:A8:13:ARG:NH1	2.27	0.49
39:A8:30:ARG:HG2	39:A8:30:ARG:HH11	1.77	0.49
45:C5:87:LYS:NZ	45:C5:88:LYS:O	2.37	0.49
50:L8:7:LYS:HE3	50:L8:32:GLN:HA	1.94	0.49
52:N8:33:CYS:SG	52:N8:40:LYS:HD3	2.52	0.49
54:Q8:46:ARG:NH2	54:Q8:48:PHE:HA	2.28	0.49
2:12:124:SER:O	2:12:126:GLU:N	2.40	0.49
1:13:266:G:H5''	1:13:267:C:C5	2.48	0.49
1:13:322:C:O3'	20:BI:23:ARG:HG3	2.13	0.49
1:13:300:A:H1'	1:13:565:U:O2	2.13	0.49
26:14:83:G:N2	26:14:102:G:H2'	2.27	0.49
26:14:1972:A:H2'	26:14:1973:G:C8	2.48	0.49
26:14:2095:C:H2'	26:14:2096:U:O4'	2.13	0.49
26:14:2872:G:C4	26:14:2873:A:C2	3.01	0.49
26:14:651:G:OP2	54:M5:21:LYS:NZ	2.32	0.49
26:14:777:A:O2'	26:14:778:G:H5'	2.13	0.49
1:1G:1294:G:H2'	1:1G:1295:G:C8	2.48	0.49
1:1G:197:A:C8	1:1G:198:G:H1'	2.48	0.49
1:1G:957:U:O2'	1:1G:959:A:N7	2.34	0.49
26:1H:1359:A:H2'	26:1H:1360:A:H5'	1.95	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1605:C:H2'	26:1H:1606:G:O4'	2.13	0.49
26:1H:1903:G:OP2	28:11:241:PRO:HB2	2.12	0.49
26:1H:2236:C:H2'	26:1H:2237:G:O4'	2.13	0.49
26:1H:836:G:H2'	26:1H:837:C:C6	2.48	0.49
10:1I:23:ILE:HA	10:1I:26:ALA:HB3	1.95	0.49
22:1L:65:5MU:H71	22:1L:65:5MU:OP2	2.12	0.49
35:25:116:SER:OG	35:25:117:LEU:N	2.46	0.49
11:2I:50:TYR:O	11:2I:55:LYS:HD3	2.12	0.49
11:2I:99:GLN:HA	11:2I:105:VAL:HG11	1.95	0.49
23:2L:43:A:O2'	23:2L:44:G:H5'	2.13	0.49
4:32:111:ALA:HB2	4:32:120:LEU:HD12	1.93	0.49
4:3E:8:VAL:HG13	4:3E:21:LEU:HD12	1.95	0.49
12:3I:28:LYS:NZ	12:3I:62:SER:HB3	2.27	0.49
31:41:68:PRO:HB3	31:41:92:VAL:HB	1.95	0.49
5:4E:41:VAL:HG13	5:4E:113:ALA:HB2	1.94	0.49
1:13:1296:C:H5'	13:4I:14:ARG:NH1	2.28	0.49
32:51:8:PRO:HG2	32:51:69:ARG:NH2	2.28	0.49
26:14:2880:C:H1'	38:55:92:GLY:HA3	1.94	0.49
7:62:65:ALA:HB1	7:62:127:ALA:HB3	1.95	0.49
8:72:29:SER:H	8:72:32:LYS:HB2	1.76	0.49
44:B5:63:LYS:H	44:B5:63:LYS:CE	2.25	0.49
45:C5:87:LYS:HB3	45:C5:94:LYS:HA	1.95	0.49
46:D5:11:GLU:HG3	46:D5:12:GLY:H	1.78	0.49
27:1J:74:U:H1'	46:D5:34:ASN:HD21	1.78	0.49
47:E5:53:MET:HG3	47:E5:59:LEU:CD2	2.42	0.49
45:G8:94:LYS:HZ2	45:G8:95:LYS:H	1.61	0.49
48:J8:71:TYR:O	48:J8:74:VAL:HG12	2.13	0.49
26:1H:75:G:H4'	49:K8:55:ARG:NH1	2.28	0.49
27:16:83:G:H4'	50:L8:52:HIS:CG	2.48	0.49
53:P8:10:ARG:HG2	53:P8:14:LYS:HD3	1.95	0.49
2:12:196:LEU:HD12	2:12:197:VAL:HG23	1.95	0.48
1:13:136:C:H2'	1:13:137:C:H5''	1.95	0.48
1:13:97:U:H2'	1:13:99:C:H6	1.77	0.48
26:14:1434:A:H2'	26:14:1435:G:C8	2.48	0.48
26:14:1689:A:H62	26:14:1698:A:H2	1.60	0.48
26:14:1754:C:N3	26:14:2716:U:O2'	2.41	0.48
26:14:2121:G:H2'	26:14:2122:U:C6	2.47	0.48
26:14:2250:G:C8	26:14:2496:C:H5''	2.46	0.48
26:14:2572:A:N7	29:29:145:LYS:HG3	2.28	0.48
26:14:49:A:H5''	26:14:51:G:O4'	2.12	0.48
1:1G:975:A:H4'	1:1G:976:G:C5'	2.33	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1564:C:O2'	26:1H:1565:C:H5'	2.12	0.48
26:1H:1885:A:H2'	26:1H:1886:C:O4'	2.13	0.48
26:1H:2286:A:H4'	26:1H:2287:A:O4'	2.13	0.48
26:1H:665:C:H2'	26:1H:666:G:H8	1.78	0.48
26:1H:971:C:H2'	26:1H:972:G:O4'	2.13	0.48
29:21:78:LEU:HD23	29:21:78:LEU:O	2.13	0.48
4:32:23:GLY:H	4:32:26:CYS:HB2	1.77	0.48
30:39:79:GLY:HA2	30:39:86:GLY:HA2	1.95	0.48
4:3E:90:GLY:O	4:3E:93:PHE:HB3	2.12	0.48
24:3K:41:G:H2'	24:3K:42:U:C6	2.48	0.48
5:42:51:VAL:O	5:42:55:VAL:HG23	2.13	0.48
5:42:59:GLY:O	5:42:63:ARG:HG2	2.13	0.48
31:49:2:PRO:HG2	31:49:4:ASP:HB3	1.95	0.48
13:4I:84:ILE:HG23	13:4I:86:CYS:H	1.78	0.48
34:58:104:LYS:HB2	34:58:117:PHE:CD1	2.47	0.48
7:6E:27:ILE:HD12	7:6E:40:ALA:HA	1.95	0.48
7:6E:45:ASP:O	7:6E:49:ILE:HG12	2.13	0.48
1:13:110:C:O2'	16:7I:25:ARG:O	2.28	0.48
9:82:26:VAL:HG22	9:82:61:ALA:N	2.28	0.48
9:82:95:LYS:HZ1	9:82:96:LEU:HB2	1.78	0.48
26:14:494:G:H5'	43:A5:8:ARG:HD3	1.95	0.48
19:AI:6:LYS:O	19:AI:7:LYS:HB3	2.11	0.48
43:E8:24:ILE:HD12	43:E8:24:ILE:O	2.12	0.48
43:E8:27:LYS:HB3	43:E8:31:GLU:HG3	1.95	0.48
28:11:108:PRO:HG3	28:11:143:HIS:HE1	1.74	0.48
2:12:92:TYR:CD1	2:12:151:GLY:HA3	2.48	0.48
1:13:1190:G:OP1	3:2E:4:LYS:HA	2.12	0.48
1:13:474:G:H2'	1:13:475:G:C8	2.48	0.48
1:13:600:C:H4'	8:7E:128:GLY:O	2.13	0.48
1:13:606:G:N2	1:13:632:A:N1	2.61	0.48
1:13:963:G:H4'	57:13:1838:HOH:O	2.13	0.48
26:14:126:A:O5'	53:L5:19:ARG:HG3	2.13	0.48
26:14:1519:G:H2'	26:14:1520:U:O4'	2.14	0.48
26:14:2503:A:OP2	26:14:2503:A:H3'	2.13	0.48
26:14:495:G:N3	43:A5:61:ASN:ND2	2.59	0.48
26:14:706:A:OP1	28:19:7:LYS:NZ	2.31	0.48
26:14:861:A:N3	27:1J:79:C:O2'	2.46	0.48
26:14:880:G:C2	26:14:881:G:C8	3.01	0.48
26:14:821:A:H2'	26:14:946:G:H5''	1.94	0.48
27:16:93:C:H2'	27:16:94:C:C6	2.47	0.48
1:1G:1157:A:H8	1:1G:1158:C:C4	2.31	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1490:C:H2'	1:1G:1491:G:O4'	2.12	0.48
1:1G:243:A:H4'	1:1G:244:U:O5'	2.12	0.48
26:1H:2110:G:C6	26:1H:2120:G:N7	2.81	0.48
26:1H:2401:U:H2'	26:1H:2402:C:C1'	2.43	0.48
26:1H:2740:A:H2'	26:1H:2741:A:C8	2.48	0.48
27:1J:2:C:O2	27:1J:118:G:N2	2.37	0.48
27:1J:94:C:H2'	27:1J:95:U:H6	1.78	0.48
11:2A:16:SER:HG	11:2A:79:SER:HG	1.54	0.48
11:2I:21:ILE:O	11:2I:85:ARG:N	2.40	0.48
4:32:64:LEU:HA	4:32:67:ILE:HD12	1.95	0.48
30:39:164:ARG:O	30:39:167:ALA:HB3	2.12	0.48
37:45:80:GLU:O	37:45:81:VAL:HG22	2.13	0.48
31:49:62:LEU:HG	31:49:143:GLU:HB3	1.94	0.48
13:4A:3:ARG:HA	13:4A:8:GLU:O	2.13	0.48
13:4I:3:ARG:HG2	13:4I:9:ILE:HG12	1.94	0.48
32:51:7:LEU:N	32:51:8:PRO:HD3	2.28	0.48
34:58:73:THR:HB	34:58:82:LEU:HD11	1.94	0.48
35:68:64:ARG:HG2	35:68:79:PHE:CG	2.48	0.48
39:A8:110:LEU:HD22	39:A8:111:GLU:HB2	1.95	0.48
45:C5:17:SER:O	45:C5:21:LYS:HB2	2.13	0.48
46:H8:77:ASP:OD2	46:H8:80:ARG:NH1	2.47	0.48
52:J5:36:CYS:SG	52:J5:49:CYS:SG	3.11	0.48
26:1H:989:G:N7	50:L8:13:ILE:HD11	2.27	0.48
51:M8:40:HIS:CG	51:M8:45:GLY:HA3	2.48	0.48
54:Q8:25:MET:HB3	54:Q8:42:ARG:HB3	1.95	0.48
1:13:659:U:H2'	1:13:660:G:C8	2.48	0.48
1:13:757:U:H2'	1:13:758:G:O4'	2.14	0.48
26:14:1536:A:C8	26:14:1537:C:H1'	2.48	0.48
26:14:1917:U:H3'	26:14:1918:A:H8	1.79	0.48
26:14:2819:G:OP1	57:14:3950:HOH:O	2.20	0.48
26:14:765:G:H2'	26:14:766:C:C6	2.48	0.48
1:1G:501:C:H2'	1:1G:502:G:H8	1.78	0.48
1:1G:653:A:C8	8:72:56:LYS:HE3	2.49	0.48
1:1G:685:G:C2	1:1G:686:U:C4	3.01	0.48
1:1G:753:A:OP1	15:6A:69:TYR:OH	2.24	0.48
26:1H:1138:G:H21	34:58:106:MET:CE	2.26	0.48
26:1H:128:C:H2'	26:1H:129:C:H6	1.79	0.48
26:1H:1318:C:H5	57:1H:3767:HOH:O	1.97	0.48
26:1H:17:G:H2'	26:1H:18:C:C6	2.48	0.48
26:1H:606:U:H4'	26:1H:658:C:H4'	1.95	0.48
30:31:9:ILE:HG23	30:31:20:LEU:O	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:39:65:TRP:HZ3	30:39:73:ALA:HB3	1.78	0.48
4:3E:92:VAL:O	4:3E:96:LEU:HD22	2.12	0.48
5:42:36:ASP:CG	5:42:38:GLN:HB2	2.34	0.48
6:52:14:LEU:HB2	6:52:18:GLN:HB2	1.95	0.48
6:52:1:MET:HB3	6:52:67:MET:O	2.13	0.48
3:22:18:TRP:NE1	14:5A:53:LEU:O	2.46	0.48
7:62:146:GLU:O	7:62:149:ARG:HB2	2.13	0.48
35:68:2:ILE:HB	35:68:33:ALA:HB3	1.95	0.48
33:69:120:ILE:HD12	33:69:126:TYR:CE2	2.48	0.48
40:75:50:ILE:HD11	40:75:102:ILE:HD11	1.93	0.48
1:13:376:G:H4'	16:7I:5:ARG:HH11	1.78	0.48
16:7I:8:ARG:C	16:7I:9:PHE:HD1	2.17	0.48
1:1G:128:G:H4'	17:8A:3:LYS:HG2	1.94	0.48
17:8I:86:GLU:O	17:8I:90:ILE:HG12	2.13	0.48
26:14:2357:U:OP1	47:E5:20:ARG:HD2	2.13	0.48
2:12:115:LEU:HB2	2:12:145:LEU:HD12	1.95	0.48
2:12:23:ARG:HH21	2:12:191:ASP:HB2	1.78	0.48
1:13:1145:C:H4'	1:13:1146:A:O5'	2.12	0.48
26:14:1001:A:H2'	26:14:1002:G:O4'	2.12	0.48
26:14:1344:G:H4'	26:14:1384:A:N7	2.28	0.48
26:14:483:A:H1'	45:C5:60:PHE:HE1	1.76	0.48
27:16:11:C:H3'	27:16:12:C:C6	2.48	0.48
1:1G:769:G:H4'	1:1G:1513:A:H4'	1.95	0.48
1:1G:542:G:N2	1:1G:543:C:C2	2.81	0.48
1:1G:580:U:H2'	1:1G:581:G:O4'	2.12	0.48
26:1H:462:C:H42	26:1H:467:G:H1	1.61	0.48
26:1H:802:A:OP1	57:1H:4017:HOH:O	2.20	0.48
29:29:15:PHE:CD2	40:75:81:PRO:HD3	2.49	0.48
26:1H:1257:C:H4'	30:31:83:PHE:CD1	2.48	0.48
30:39:118:ALA:HB2	30:39:123:LEU:HD21	1.94	0.48
12:3I:53:ARG:HB3	12:3I:69:TYR:HE1	1.78	0.48
37:45:59:ARG:O	37:45:60:ARG:HG3	2.12	0.48
31:49:108:ASN:OD1	31:49:108:ASN:N	2.46	0.48
6:52:76:ALA:O	6:52:80:ARG:HB2	2.13	0.48
33:69:77:LEU:HD22	33:69:141:LYS:HE3	1.94	0.48
33:69:8:PRO:HD3	33:69:15:VAL:HG22	1.94	0.48
15:6I:39:LEU:O	15:6I:42:HIS:HB3	2.14	0.48
15:6I:71:GLN:HG2	15:6I:71:GLN:O	2.13	0.48
36:78:121:LYS:HE2	36:78:123:LEU:HD21	1.94	0.48
2:1E:178:ARG:HH21	8:7E:74:PRO:HB3	1.78	0.48
42:95:76:LYS:HD2	42:95:80:GLN:HB3	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:98:94:TYR:CD1	38:98:94:TYR:N	2.79	0.48
41:C8:6:THR:N	57:C8:202:HOH:O	2.45	0.48
46:D5:163:LEU:HD12	46:D5:165:VAL:HG23	1.95	0.48
46:D5:24:LEU:HD12	46:D5:25:PRO:O	2.13	0.48
49:K8:3:LEU:O	49:K8:6:VAL:HG13	2.13	0.48
1:13:1002:G:C5	1:13:1003:G:C8	3.01	0.48
1:13:103:C:H2'	1:13:104:G:C8	2.49	0.48
1:13:1434:A:H2'	1:13:1435:G:O4'	2.13	0.48
26:14:1007:C:H5''	34:15:35:ARG:HH11	1.78	0.48
26:14:2512:C:H4'	29:29:122:PHE:CE2	2.48	0.48
26:14:358:U:H2'	26:14:359:A:H8	1.78	0.48
26:14:446:G:P	57:14:3971:HOH:O	2.72	0.48
26:14:681:G:H2'	26:14:682:G:O4'	2.12	0.48
28:19:95:LEU:HD11	28:19:105:ILE:HD12	1.95	0.48
1:1G:1251:A:H2'	1:1G:1252:A:C8	2.48	0.48
1:1G:308:C:H2'	1:1G:309:G:C8	2.49	0.48
1:1G:790:A:H2'	1:1G:791:G:C8	2.48	0.48
26:1H:1759:A:H4'	26:1H:2715:C:O4'	2.13	0.48
26:1H:2360:A:H2'	26:1H:2361:A:O4'	2.14	0.48
26:1H:2845:G:H2'	26:1H:2846:G:C8	2.48	0.48
26:1H:722:A:C2	26:1H:723:G:C4	3.01	0.48
26:1H:775:G:C5	26:1H:794:G:C8	3.01	0.48
27:1J:15:A:H5'	27:1J:16:G:N7	2.29	0.48
3:2E:68:VAL:HG12	3:2E:70:VAL:HG13	1.94	0.48
4:32:54:TYR:CE2	4:32:58:LEU:HD12	2.49	0.48
4:32:65:ARG:HD3	4:32:75:PHE:CG	2.48	0.48
31:41:106:LEU:HD11	31:41:111:LEU:HG	1.95	0.48
5:42:101:ILE:O	5:42:120:THR:HG23	2.13	0.48
5:4E:39:GLY:HA3	5:4E:71:LEU:HD11	1.94	0.48
15:6I:39:LEU:HD13	15:6I:56:LEU:HB2	1.94	0.48
36:78:103:ALA:HB3	36:78:105:LEU:HD12	1.95	0.48
26:14:445:C:OP1	41:85:2:PRO:HA	2.13	0.48
39:A8:83:LYS:O	39:A8:110:LEU:HB2	2.13	0.48
39:A8:51:ALA:HB3	39:A8:73:LEU:HG	1.95	0.48
42:D8:5:VAL:HB	42:D8:35:LEU:HD11	1.94	0.48
48:J8:58:ILE:CG2	48:J8:87:PRO:HG3	2.44	0.48
26:1H:210:C:OP1	53:P8:29:LYS:HD2	2.14	0.48
28:11:109:ASP:HB2	28:11:197:GLY:HA3	1.96	0.48
1:13:1031:G:H2'	1:13:1032:A:H5'	1.95	0.48
1:13:1037:C:H2'	1:13:1038:C:H6	1.77	0.48
1:13:959:A:HO2'	1:13:984:C:HO2'	1.61	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2468:G:H3'	26:14:2476:A:N1	2.29	0.48
26:14:2542:A:H2'	26:14:2542:A:N3	2.27	0.48
26:14:274:G:H3'	26:14:274:G:H8	1.79	0.48
26:14:635:C:H2'	26:14:636:G:O4'	2.13	0.48
34:15:34:LEU:O	34:15:49:GLY:HA3	2.13	0.48
10:1A:24:VAL:HG21	10:1A:37:PRO:HD3	1.95	0.48
10:1A:81:THR:OG1	10:1A:82:ILE:N	2.46	0.48
21:1B:8:THR:HG22	21:1B:10:ARG:H	1.79	0.48
1:1G:1084:G:C5	1:1G:1085:U:C4	3.00	0.48
1:1G:280:C:OP1	17:8A:91:ARG:NH2	2.44	0.48
1:1G:620:C:H2'	1:1G:621:A:O4'	2.14	0.48
26:1H:1411:C:H2'	26:1H:1412:A:C8	2.49	0.48
26:1H:1835:G:H5'	26:1H:1836:C:OP2	2.13	0.48
26:1H:2492:U:H2'	26:1H:2493:U:H6	1.79	0.48
26:1H:2756:U:H4'	26:1H:2757:A:OP1	2.12	0.48
27:1J:3:C:H42	27:1J:117:G:H1	1.61	0.48
22:1K:57:C:H2'	22:1K:58:U:H4'	1.95	0.48
26:1H:2829:C:H5'	29:21:76:ARG:HH22	1.78	0.48
3:22:51:GLY:O	3:22:70:VAL:HG13	2.13	0.48
3:2E:3:ASN:O	3:2E:4:LYS:HG2	2.14	0.48
3:2E:84:ILE:HA	3:2E:87:LEU:HD12	1.95	0.48
4:32:157:LEU:HD12	4:32:161:ASN:ND2	2.28	0.48
30:39:162:LEU:H	30:39:162:LEU:HD12	1.79	0.48
24:3L:6:G:O2'	24:3L:7:G:O5'	2.27	0.48
31:41:111:LEU:HD21	31:41:120:LEU:HD21	1.96	0.48
13:4A:97:PRO:HA	13:4A:110:ARG:HD3	1.94	0.48
34:58:33:LEU:HD12	34:58:38:HIS:HD2	1.76	0.48
10:1A:62:HIS:HB3	14:5A:59:ALA:HB3	1.95	0.48
14:5I:17:LYS:HE3	14:5I:18:VAL:HG13	1.96	0.48
15:6A:43:LEU:HD11	15:6A:53:HIS:HA	1.95	0.48
15:6I:79:ARG:HB3	15:6I:79:ARG:CZ	2.43	0.48
26:1H:1754:C:OP1	40:B8:96:ARG:NH1	2.46	0.48
20:BA:54:LYS:HA	20:BA:57:ARG:NH1	2.28	0.48
20:BI:42:GLN:O	20:BI:46:GLU:HG3	2.14	0.48
41:C8:88:ILE:HD11	41:C8:112:ARG:HB3	1.95	0.48
41:C8:88:ILE:C	41:C8:90:VAL:N	2.66	0.48
44:F8:3:THR:HA	44:F8:6:ASP:OD2	2.13	0.48
46:H8:95:PRO:HB2	46:H8:127:LYS:HD2	1.96	0.48
46:H8:45:ASP:OD1	46:H8:49:ARG:NH1	2.46	0.48
26:1H:1184:G:H5'	50:L8:29:ARG:NH1	2.29	0.48
51:M8:9:LEU:N	51:M8:27:THR:OG1	2.45	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:Q8:24:ALA:HA	54:Q8:44:LYS:HB2	1.94	0.48
54:Q8:30:ARG:HB2	54:Q8:30:ARG:HH11	1.78	0.48
1:13:1004:A:C2	1:13:1025:U:H1'	2.49	0.48
1:13:1070:U:H2'	1:13:1071:C:C6	2.49	0.48
1:13:1266:G:N2	1:13:1270:C:N3	2.61	0.48
1:13:519:C:H2'	1:13:520:A:O4'	2.14	0.48
26:14:1268:A:H2'	26:14:1269:A:O4'	2.13	0.48
26:14:1510:A:H2'	26:14:1511:A:C8	2.49	0.48
26:14:2505:G:H2'	26:14:2576:G:O6	2.14	0.48
34:15:137:LYS:HD3	34:15:137:LYS:HA	1.63	0.48
1:1G:1080:A:H5''	1:1G:1081:G:OP2	2.12	0.48
1:1G:1226:C:H4'	19:AA:80:TYR:OH	2.13	0.48
1:1G:306:G:O5'	1:1G:306:G:H8	1.96	0.48
1:1G:974:A:H5'	1:1G:975:A:OP1	2.13	0.48
26:1H:1427:A:H4'	26:1H:1428:C:O4'	2.14	0.48
26:1H:1496:A:C8	26:1H:1577:C:O2'	2.59	0.48
26:1H:1562:A:H2'	26:1H:1563:G:C8	2.48	0.48
26:1H:2454:G:H1'	57:1H:4116:HOH:O	2.12	0.48
26:1H:606:U:OP2	30:31:104:LYS:NZ	2.42	0.48
27:1J:21:G:H2'	27:1J:22:U:O4'	2.13	0.48
29:21:201:THR:HG22	29:21:203:LYS:N	2.18	0.48
26:14:2784:C:H1'	29:29:37:ARG:HH21	1.78	0.48
23:2L:27:G:H1	23:2L:45:U:H3	1.60	0.48
24:3K:67:C:O2'	24:3K:68:A:H5'	2.14	0.48
5:42:57:LYS:HE2	5:42:61:TYR:OH	2.14	0.48
13:4I:52:GLU:O	13:4I:56:LEU:HB2	2.14	0.48
25:4K:32:A:H4'	25:4K:33:G:OP1	2.13	0.48
16:7I:3:LYS:O	16:7I:21:VAL:HA	2.13	0.48
37:88:109:VAL:HG12	37:88:114:ALA:HB2	1.94	0.48
26:1H:996:A:O3'	41:C8:92:ARG:HG2	2.12	0.48
53:L5:34:ARG:HG2	53:L5:39:ARG:HG3	1.94	0.48
27:16:40:U:O4	51:M8:2:LYS:N	2.47	0.48
51:M8:57:GLU:HA	51:M8:60:GLN:OE1	2.14	0.48
26:1H:458:G:C5	53:P8:37:LYS:HE3	2.48	0.48
54:Q8:26:LYS:HB2	54:Q8:26:LYS:HE3	1.68	0.48
54:Q8:32:LEU:HG	54:Q8:33:ASN:ND2	2.28	0.48
1:13:1442:G:C6	1:13:1446:A:N6	2.82	0.48
1:13:595:G:H1'	1:13:596:C:H5	1.79	0.48
1:13:909:A:H2'	1:13:910:C:O4'	2.14	0.48
26:14:848:G:C4	26:14:933:A:C8	3.02	0.48
28:19:24:ILE:HD13	28:19:84:TYR:HB2	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1322:C:O2'	1:1G:1323:G:H5'	2.14	0.48
1:1G:134:A:H61	16:7A:25:ARG:NH1	2.11	0.48
1:1G:937:A:H1'	1:1G:1379:G:N2	2.29	0.48
26:1H:144:C:H2'	26:1H:145:G:C8	2.49	0.48
26:1H:1641:A:H2'	26:1H:1642:G:O4'	2.14	0.48
26:1H:2160:G:C2	26:1H:2161:C:H1'	2.49	0.48
26:1H:2233:U:H2'	26:1H:2234:G:C8	2.49	0.48
26:1H:448:U:O4	26:1H:583:G:H1'	2.14	0.48
27:1J:72:G:O2'	27:1J:104:A:N6	2.46	0.48
22:1K:21:A:N3	22:1K:21:A:H2'	2.29	0.48
22:1K:8:U:H1'	22:1K:22:G:N2	2.28	0.48
31:41:139:LEU:HA	31:41:144:ILE:HB	1.95	0.48
38:55:33:ARG:HA	38:55:115:GLU:HA	1.95	0.48
7:6E:133:GLY:O	7:6E:136:LYS:HB2	2.14	0.48
37:88:68:ILE:HD13	37:88:103:MET:HB3	1.96	0.48
41:C8:17:ILE:HG23	41:C8:39:LEU:HD12	1.95	0.48
41:C8:75:ASN:HB3	41:C8:77:SER:H	1.78	0.48
46:D5:29:TYR:HA	46:D5:33:LEU:O	2.14	0.48
42:D8:27:ALA:HB3	42:D8:61:VAL:HG11	1.96	0.48
51:I5:13:ARG:HB3	51:I5:22:ILE:HG21	1.96	0.48
52:N8:3:LYS:HE3	52:N8:3:LYS:HB3	1.56	0.48
1:13:1016:A:H2'	1:13:1017:G:O4'	2.13	0.48
1:13:1318:A:H1'	19:AI:37:ARG:NH2	2.23	0.48
1:13:179:A:H2'	1:13:180:U:C6	2.49	0.48
26:14:1076:C:C5	26:14:1077:A:H1'	2.49	0.48
26:14:1110:G:O2'	26:14:1111:A:O4'	2.29	0.48
26:14:1425:G:H2'	26:14:1426:G:C8	2.49	0.48
26:14:1496:A:H2'	26:14:1498:C:C5	2.49	0.48
26:14:1667:G:O2'	26:14:1991:U:O4	2.27	0.48
26:14:2086:U:H2'	26:14:2087:G:C8	2.48	0.48
26:14:2346:A:C2	26:14:2383:G:C2	3.02	0.48
26:14:2068:U:N3	26:14:2430:A:H2	2.11	0.48
26:14:794:G:H2'	26:14:795:C:C6	2.48	0.48
28:19:271:ILE:O	28:19:272:ALA:HB2	2.13	0.48
1:1G:1278:U:H5'	1:1G:1279:A:H5'	1.96	0.48
1:1G:1513:A:H2'	1:1G:1514:C:C6	2.49	0.48
1:1G:390:C:H2'	1:1G:391:G:C8	2.48	0.48
1:1G:737:A:H2'	1:1G:738:C:H6	1.78	0.48
1:1G:998(A):C:H2'	1:1G:999:U:C6	2.49	0.48
26:1H:1408:C:C2	26:1H:1595:G:N2	2.82	0.48
26:1H:150:C:H42	26:1H:176:G:H1	1.60	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1786:A:C2	26:1H:2606:C:H1'	2.49	0.48
26:1H:2317:C:H2'	26:1H:2318:G:O4'	2.14	0.48
26:1H:322:A:OP2	30:31:169:ASN:HB2	2.14	0.48
29:21:16:ARG:HH21	29:21:173:VAL:HG13	1.79	0.48
29:21:167:VAL:HG21	29:21:187:ALA:CB	2.44	0.48
29:29:58:ARG:O	29:29:59:VAL:HG22	2.14	0.48
3:2E:52:LEU:HA	3:2E:70:VAL:HG12	1.96	0.48
32:51:77:LYS:O	32:51:77:LYS:HG2	2.13	0.48
33:61:57:ARG:HG2	33:61:61:ARG:HH12	1.79	0.48
7:6E:132:GLY:O	7:6E:135:VAL:HG23	2.14	0.48
7:6E:5:ARG:NE	7:6E:7:ALA:HA	2.28	0.48
40:75:107:ASP:N	40:75:107:ASP:OD1	2.45	0.48
8:7E:41:ARG:HG2	8:7E:41:ARG:O	2.14	0.48
12:3A:8:ASN:OD1	17:8A:34:LYS:HE2	2.13	0.48
18:9I:46:GLU:HG2	18:9I:85:LEU:HD13	1.95	0.48
44:B5:40:LYS:HA	44:B5:51:VAL:HG11	1.95	0.48
48:J8:87:PRO:HA	48:J8:90:ILE:CG1	2.43	0.48
52:N8:40:LYS:HG3	52:N8:47:PRO:HD2	1.95	0.48
1:13:110:C:H2'	1:13:111:G:O4'	2.14	0.48
1:13:725:G:H2'	1:13:726:C:H6	1.79	0.48
26:14:1291:C:H2'	26:14:1292:U:C6	2.49	0.48
26:14:140:A:C8	26:14:1408:C:O2'	2.66	0.48
26:14:1488:G:C6	26:14:1489:U:C2	3.02	0.48
26:14:1537:C:H2'	26:14:1538:G:H8	1.78	0.48
26:14:184:C:H2'	26:14:185:U:H6	1.79	0.48
26:14:2064:C:H2'	26:14:2065:C:C6	2.49	0.48
26:14:2533:A:O2'	26:14:2664:G:H5'	2.13	0.48
26:14:380:U:H2'	26:14:381:G:H8	1.79	0.48
26:14:529:A:H8	26:14:530:G:C6	2.32	0.48
26:14:864:G:C6	26:14:865:C:N4	2.82	0.48
1:1G:1317:C:H5''	1:1G:1318:A:OP2	2.14	0.48
1:1G:310:G:OP2	16:7A:27:LYS:NZ	2.33	0.48
1:1G:967:C:H2'	1:1G:968:A:N7	2.29	0.48
26:1H:141(A):C:H2'	26:1H:142:G:O4'	2.13	0.48
26:1H:1929:G:H4'	26:1H:1930:G:OP1	2.14	0.48
26:1H:2183:C:H2'	26:1H:2184:G:H8	1.79	0.48
26:1H:2305:A:O2'	31:41:136:ARG:NH1	2.47	0.48
26:1H:2508:G:H2'	26:1H:2509:G:H8	1.79	0.48
26:1H:2795:G:H2'	26:1H:2798:C:H5''	1.96	0.48
26:1H:662:G:H5''	36:78:16:ARG:HB2	1.95	0.48
26:1H:710:G:H2'	26:1H:711:G:C8	2.49	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:25:63:VAL:O	35:25:64:ARG:HG3	2.14	0.48
29:29:31:CYS:HB3	29:29:49:LEU:HB2	1.96	0.48
29:29:5:LEU:HD22	29:29:49:LEU:HD12	1.95	0.48
23:2K:43:A:O2'	23:2K:44:G:P	2.72	0.48
30:31:101:LEU:HD22	30:31:102:PRO:HD2	1.95	0.48
4:32:43:HIS:HA	4:32:46:LYS:HE3	1.96	0.48
26:14:662:G:H5'	36:35:14:LYS:CB	2.43	0.48
30:39:95:ARG:HG3	30:39:97:TYR:CE1	2.49	0.48
12:3I:51:ALA:O	12:3I:52:LEU:HD23	2.14	0.48
37:45:103:MET:HE3	37:45:125:LEU:HD13	1.95	0.48
13:4I:108:ARG:N	13:4I:108:ARG:HD2	2.29	0.48
32:51:102:ALA:HA	32:51:117:PRO:HD3	1.95	0.48
8:72:51:VAL:HG11	8:72:60:ARG:HE	1.78	0.48
41:85:91:ASP:OD1	41:85:96:ALA:HB2	2.14	0.48
39:A8:99:LYS:O	39:A8:103:GLU:HG2	2.14	0.48
19:AA:51:VAL:O	19:AA:57:HIS:HA	2.14	0.48
48:J8:92:LYS:HD2	48:J8:95:LEU:HD12	1.96	0.48
28:11:130:ALA:HB2	28:11:192:THR:HB	1.96	0.47
2:12:222:ILE:O	2:12:226:ARG:HB2	2.14	0.47
1:13:560:U:H4'	1:13:561:U:C5'	2.43	0.47
26:14:1047:G:N2	26:14:1111:A:H62	2.12	0.47
26:14:1181:C:H2'	26:14:1182:A:C8	2.49	0.47
26:14:2537:U:H2'	26:14:2538:C:C6	2.49	0.47
26:14:2693:A:H2'	26:14:2694:G:H8	1.79	0.47
26:14:795:C:O2'	26:14:796:C:H5'	2.14	0.47
27:16:78:A:C2	27:16:99:A:C4	3.02	0.47
2:1E:98:LEU:HB2	2:1E:101:MET:HG3	1.96	0.47
26:1H:1045:A:C2	26:1H:1111:A:C5	3.02	0.47
26:1H:1175:U:H4'	26:1H:1176:G:H4'	1.96	0.47
26:1H:198:C:C2'	26:1H:199:A:H5''	2.44	0.47
26:1H:2035:G:H4'	26:1H:2036:C:OP2	2.13	0.47
26:1H:2243:U:H2'	26:1H:2244:U:C6	2.48	0.47
26:1H:2248:C:H2'	26:1H:2249:U:O4'	2.13	0.47
26:1H:2502:G:H5''	26:1H:2503:A:C5'	2.44	0.47
26:1H:545:G:H2'	26:1H:546:C:H5''	1.96	0.47
26:1H:639:U:H2'	26:1H:640:C:C6	2.49	0.47
3:22:94:LEU:HD12	3:22:95:THR:H	1.79	0.47
3:2E:79:ARG:NH2	11:2A:99:GLN:HG2	2.29	0.47
23:2L:47:C:N4	23:2L:57:C:H42	2.12	0.47
4:3E:155:LEU:O	4:3E:157:LEU:N	2.47	0.47
24:3K:85:C:H4'	48:J8:23:LYS:HB2	1.95	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:45:36:ALA:HB2	37:45:103:MET:HE1	1.95	0.47
13:4A:70:LEU:O	13:4A:74:VAL:HG23	2.14	0.47
32:51:137:ASP:HB3	32:51:140:LYS:HB3	1.95	0.47
1:1G:750:G:N2	15:6A:23:GLY:HA3	2.29	0.47
38:98:3:HIS:O	38:98:5:LYS:N	2.47	0.47
38:98:76:VAL:HG13	38:98:80:PHE:HE2	1.77	0.47
43:E8:58:ALA:O	43:E8:64:MET:HB2	2.14	0.47
49:G5:47:ASN:C	49:G5:49:LYS:H	2.14	0.47
45:G8:99:CYS:SG	45:G8:100:ALA:N	2.86	0.47
50:L8:31:LEU:O	50:L8:32:GLN:HB2	2.14	0.47
1:13:1167:A:C6	1:13:1169:A:C6	3.02	0.47
1:13:60:A:H8	1:13:60:A:OP1	1.97	0.47
1:13:587:G:N2	1:13:755:G:C5	2.82	0.47
26:14:1041:C:H2'	26:14:1042:G:C8	2.49	0.47
26:14:528:A:H2	26:14:2043:C:C5'	2.27	0.47
26:14:2139:C:N4	26:14:2152:G:O6	2.48	0.47
26:14:671:C:H2'	26:14:672:C:C6	2.48	0.47
2:1E:174:VAL:HG13	2:1E:184:VAL:HG11	1.95	0.47
1:1G:1054:C:H6	1:1G:1196:U:HO2'	1.61	0.47
1:1G:1399:C:C2	1:1G:1502:A:N6	2.82	0.47
1:1G:56:U:H2'	1:1G:57:G:C8	2.49	0.47
26:1H:1170:G:N2	26:1H:1180:C:C2	2.83	0.47
26:1H:1372:U:H2'	26:1H:1373:A:H8	1.79	0.47
26:1H:448:U:C4	26:1H:583:G:H1'	2.50	0.47
26:1H:753:C:O2'	26:1H:754:C:H5'	2.14	0.47
10:1I:4:ILE:HG12	10:1I:100:THR:HG23	1.96	0.47
22:1K:24:C:H2'	22:1K:25:A:O4'	2.14	0.47
29:21:21:VAL:HA	29:21:22:PRO:HD3	1.45	0.47
3:22:139:GLN:NE2	3:22:142:MET:SD	2.85	0.47
11:2I:52:GLY:H	11:2I:55:LYS:HG3	1.79	0.47
23:2L:20:U:C4	23:2L:22:G:H4'	2.48	0.47
26:1H:588:U:H1'	30:31:90:PHE:CG	2.49	0.47
4:3E:108:LEU:HB3	4:3E:110:PHE:CD1	2.48	0.47
31:41:35:GLU:OE1	31:41:95:ARG:HB3	2.14	0.47
37:45:38:GLU:HG3	37:45:127:ILE:CG2	2.44	0.47
33:69:111:PRO:O	33:69:113:ARG:HB2	2.14	0.47
7:6E:73:MET:HG2	7:6E:90:GLU:HB3	1.95	0.47
15:6I:56:LEU:O	15:6I:60:VAL:HG23	2.14	0.47
16:7I:22:THR:OG1	16:7I:32:TYR:HA	2.14	0.47
38:98:44:LEU:HA	38:98:44:LEU:HD23	1.73	0.47
27:16:7:G:O5'	39:A8:29:PHE:CE2	2.66	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
45:G8:81:LYS:HB3	45:G8:82:PRO:HA	1.96	0.47
1:13:1133:G:N2	1:13:1141:C:O2	2.47	0.47
1:13:1280:A:C3'	1:13:1281:U:H5'	2.45	0.47
26:14:184:C:H2'	26:14:185:U:C6	2.48	0.47
26:14:1966:A:H4'	26:14:1967:C:OP1	2.14	0.47
26:14:2275:C:O2	37:45:85:LYS:HG3	2.14	0.47
26:14:270(D):C:H2'	26:14:270(E):G:C8	2.50	0.47
26:14:2849:U:H5'	26:14:2867:G:N2	2.29	0.47
26:14:796:C:H2'	26:14:797:C:H6	1.73	0.47
27:16:44:G:H1'	27:16:47:C:H42	1.79	0.47
1:1G:1224:G:C2	1:1G:1322:C:H1'	2.49	0.47
1:1G:393:A:OP2	16:7A:12:LYS:NZ	2.25	0.47
1:1G:7:G:H21	5:42:121:LYS:HG2	1.79	0.47
26:1H:1915:U:H2'	26:1H:1916:A:O4'	2.13	0.47
26:1H:2254:C:H3'	57:1H:3881:HOH:O	2.13	0.47
26:1H:919:G:H4'	27:16:81:G:H4'	1.95	0.47
22:1L:44:G:H2'	22:1L:45:U:O4'	2.14	0.47
29:29:120:TRP:CD1	29:29:155:LYS:HB3	2.49	0.47
30:39:181:LEU:CD2	30:39:186:ILE:HD11	2.44	0.47
24:3K:63:G:H2'	24:3K:64:G:O4'	2.15	0.47
31:41:45:GLU:H	31:41:45:GLU:HG2	1.36	0.47
13:4A:27:LYS:HE3	13:4A:31:LYS:NZ	2.30	0.47
33:61:98:ALA:HB2	33:61:111:PRO:HB3	1.97	0.47
1:13:1179:A:H4'	9:8E:103:THR:HA	1.96	0.47
18:9I:85:LEU:HD12	18:9I:86:VAL:N	2.28	0.47
26:1H:2376:A:H2	39:A8:112:PHE:HB3	1.78	0.47
45:C5:39:VAL:HG23	45:C5:41:GLY:H	1.78	0.47
45:G8:28:LYS:HZ2	45:G8:40:GLU:HG2	1.79	0.47
47:I8:49:LYS:HB2	47:I8:80:HIS:HB3	1.96	0.47
52:N8:41:PRO:HD2	52:N8:44:THR:CG2	2.44	0.47
54:Q8:33:ASN:OD1	54:Q8:34:TRP:HB2	2.13	0.47
2:12:102:LEU:HD12	2:12:102:LEU:H	1.80	0.47
1:13:1064:G:H4'	1:13:1065:U:OP1	2.13	0.47
1:13:1390:U:H2'	1:13:1391:U:H6	1.79	0.47
1:13:1441:G:H5''	1:13:1442:G:H5'	1.96	0.47
1:13:658:G:H2'	1:13:659:U:C6	2.49	0.47
26:14:1202:C:H42	26:14:1243:G:H1	1.61	0.47
26:14:1328:G:H2'	26:14:1330:C:C5	2.50	0.47
26:14:1784:A:H4'	26:14:1785:A:O5'	2.15	0.47
26:14:2477:C:H5'	26:14:2479:G:O6	2.14	0.47
2:1E:100:GLY:N	2:1E:176:GLU:OE2	2.42	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:255:G:O6	1:1G:270:A:N6	2.48	0.47
1:1G:429:U:OP1	4:32:13:ARG:NH2	2.36	0.47
26:1H:1260:G:H2'	26:1H:1261:C:C6	2.49	0.47
26:1H:1534:G:H21	26:1H:1535:U:H5''	1.79	0.47
26:1H:794:G:H2'	26:1H:795:C:C6	2.49	0.47
22:1K:53:A:H62	22:1K:54:G:H21	1.62	0.47
11:2I:41:THR:HG21	11:2I:71:LYS:HD3	1.95	0.47
4:32:8:VAL:HG12	4:32:21:LEU:HD12	1.96	0.47
38:55:88:ARG:HD2	38:55:89:ASP:OD1	2.13	0.47
1:13:982:U:H5''	14:5I:6:LEU:HD13	1.96	0.47
33:69:101:LEU:HB2	33:69:107:VAL:O	2.13	0.47
40:75:51:ARG:HD3	40:75:100:TYR:OH	2.15	0.47
17:8A:45:HIS:ND1	17:8A:65:ILE:HG21	2.29	0.47
17:8I:76:LEU:HD12	17:8I:77:VAL:H	1.80	0.47
20:BI:12:ALA:O	20:BI:15:ARG:HB2	2.12	0.47
45:C5:20:TYR:CE2	45:C5:42:VAL:HA	2.50	0.47
48:F5:8:SER:HB3	48:F5:66:HIS:CD2	2.50	0.47
53:L5:24:THR:O	53:L5:28:ARG:HG3	2.14	0.47
43:E8:38:TYR:OH	52:N8:47:PRO:HG3	2.15	0.47
28:11:68:LYS:HB3	28:11:70:TRP:CZ3	2.49	0.47
2:12:55:PHE:HD1	2:12:58:ILE:HG13	1.78	0.47
1:13:830:G:H2'	1:13:831:U:O4'	2.15	0.47
26:14:1485:G:H2'	26:14:1486:A:H8	1.79	0.47
26:14:1967:C:H2'	26:14:1968:G:O4'	2.14	0.47
10:1A:76:ASN:HB3	10:1A:78:ASN:ND2	2.29	0.47
2:1E:239:VAL:HG12	2:1E:239:VAL:O	2.14	0.47
2:1E:85:ALA:HB1	2:1E:90:MET:O	2.14	0.47
1:1G:517:G:N2	1:1G:533:A:OP2	2.41	0.47
1:1G:890:G:O2'	1:1G:906:G:O6	2.21	0.47
26:1H:1013:C:N4	26:1H:1149:G:H1	2.12	0.47
26:1H:1786:A:H2	26:1H:2606:C:H1'	1.79	0.47
26:1H:2125:G:H21	26:1H:2173:A:N6	2.10	0.47
26:1H:2287:A:C2	26:1H:2289:G:C8	3.03	0.47
26:1H:387:U:O4	57:1H:4006:HOH:O	2.19	0.47
26:1H:844:C:H2'	26:1H:845:G:O4'	2.14	0.47
1:13:972:C:O2	10:1I:55:LYS:HD3	2.14	0.47
4:32:91:SER:HA	4:32:94:LEU:HD12	1.97	0.47
4:3E:159:ARG:O	4:3E:163:GLU:N	2.46	0.47
4:3E:8:VAL:CG1	4:3E:21:LEU:HB2	2.44	0.47
31:49:181:ARG:HB3	31:49:181:ARG:NH1	2.30	0.47
7:62:91:VAL:HB	7:62:96:GLN:HG2	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:82:16:ARG:NH1	9:82:64:THR:HG21	2.29	0.47
37:88:66:ILE:HG22	37:88:67:ARG:H	1.79	0.47
43:A5:33:ARG:NH1	43:A5:52:GLU:OE2	2.48	0.47
39:A8:5:THR:HG23	39:A8:8:GLU:OE2	2.15	0.47
45:C5:75:ILE:O	45:C5:76:CYS:HB3	2.15	0.47
47:E5:23:VAL:HA	47:E5:38:VAL:HG22	1.96	0.47
43:E8:97:LYS:HE2	43:E8:99:ARG:NH2	2.28	0.47
51:I5:21:VAL:HG22	51:I5:22:ILE:H	1.79	0.47
49:K8:15:LYS:HD3	49:K8:67:LYS:HZ3	1.80	0.47
53:L5:8:ASN:C	53:L5:8:ASN:OD1	2.52	0.47
54:M5:59:LYS:NZ	54:M5:59:LYS:HA	2.24	0.47
1:13:1187:G:O5'	9:8E:113:LYS:NZ	2.34	0.47
1:13:1346:A:H5''	9:8E:120:ARG:HH12	1.80	0.47
1:13:976:G:H5'	1:13:1358:U:O2'	2.14	0.47
1:13:769:G:H4'	1:13:1513:A:H4'	1.97	0.47
1:13:684:A:N6	1:13:685:G:C6	2.83	0.47
26:14:2196:C:O2'	26:14:2197:U:H5'	2.14	0.47
26:14:274:G:H3'	26:14:274:G:C8	2.50	0.47
26:14:2872:G:C5	26:14:2873:A:C2	3.03	0.47
26:14:957:A:N6	26:14:2459:A:C8	2.83	0.47
1:1G:188:U:O2'	1:1G:189:U:H5'	2.15	0.47
1:1G:316:G:OP2	1:1G:351:G:O2'	2.31	0.47
26:1H:1072:C:H2'	26:1H:1093:G:O6	2.15	0.47
26:1H:1756:G:H1'	26:1H:1758:G:C2	2.49	0.47
26:1H:2455:G:H2'	26:1H:2456:C:C6	2.50	0.47
26:1H:654(A):A:H2	26:1H:654(T):A:N1	2.12	0.47
22:1K:48:C:N3	22:1K:56:G:N2	2.62	0.47
30:31:108:LYS:O	30:31:112:MET:HG3	2.14	0.47
4:32:24:GLU:OE2	4:32:24:GLU:N	2.47	0.47
31:41:173:LEU:HD22	31:41:178:PHE:CE2	2.50	0.47
13:4I:16:ASP:HA	13:4I:19:LEU:HD22	1.95	0.47
34:58:7:LYS:HB3	34:58:7:LYS:HE3	1.71	0.47
14:5A:17:LYS:HD2	14:5A:18:VAL:N	2.29	0.47
6:5E:1:MET:SD	6:5E:66:GLU:HG2	2.55	0.47
33:61:40:THR:HB	33:61:43:ASN:H	1.79	0.47
20:BI:35:THR:HG22	20:BI:38:LYS:NZ	2.30	0.47
45:C5:12:THR:O	45:C5:75:ILE:HB	2.15	0.47
46:D5:24:LEU:HD23	46:D5:41:LEU:HA	1.95	0.47
43:E8:82:LEU:HA	43:E8:82:LEU:HD23	1.60	0.47
52:J5:52:TYR:HD1	52:J5:53:ALA:H	1.62	0.47
54:Q8:21:LYS:HA	54:Q8:21:LYS:HD2	1.55	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:309:G:H1'	1:13:608:A:C2	2.50	0.47
1:13:412:A:H4'	1:13:413:G:O5'	2.14	0.47
1:13:689:C:P	11:2I:46:GLY:HA3	2.55	0.47
26:14:212:G:H2'	26:14:213:A:O4'	2.14	0.47
26:14:2344:U:H4'	26:14:2345:G:OP1	2.14	0.47
26:14:2352:A:C2	47:E5:33:ALA:HB1	2.50	0.47
26:14:2479:G:C6	26:14:2480:C:C4	3.03	0.47
26:14:273(C):C:H5'	26:14:273(D):C:OP2	2.15	0.47
26:14:290:G:O5'	26:14:290:G:H8	1.97	0.47
26:14:864:G:H1'	26:14:914:C:N4	2.30	0.47
27:16:119:A:H2'	27:16:119:A:N3	2.30	0.47
21:1F:15:ARG:NH1	21:1F:15:ARG:HB2	2.28	0.47
1:1G:1277:C:O2'	1:1G:1279:A:H1'	2.15	0.47
1:1G:934:C:O2'	1:1G:1344:C:OP2	2.30	0.47
26:1H:1021:A:C8	26:1H:1022:G:H5''	2.50	0.47
26:1H:1508:A:O2'	26:1H:1509:C:O5'	2.31	0.47
26:1H:1827:C:C2'	26:1H:1828:G:H5'	2.45	0.47
26:1H:300:A:H2'	26:1H:334:C:H1'	1.97	0.47
26:1H:363(F):A:H4'	26:1H:364:C:H5'	1.96	0.47
26:1H:485:C:H2'	26:1H:486:C:C6	2.49	0.47
26:1H:760:G:H2'	26:1H:761:A:H5'	1.96	0.47
22:1K:81:G:H2'	22:1K:82:G:O4'	2.14	0.47
29:21:55:ASN:CB	29:21:58:ARG:HD2	2.42	0.47
3:22:72:LYS:NZ	3:22:75:VAL:HG23	2.29	0.47
29:29:95:ILE:HG13	29:29:95:ILE:H	1.46	0.47
3:2E:74:GLY:HA2	3:2E:77:ILE:HB	1.97	0.47
23:2L:15:A:H2'	23:2L:70:G:N1	2.30	0.47
4:32:141:ARG:HG2	4:32:141:ARG:HH11	1.80	0.47
36:35:78:PRO:HA	36:35:110:TYR:CD2	2.50	0.47
4:3E:103:ASN:O	4:3E:107:ARG:HG2	2.14	0.47
4:3E:192:GLU:OE1	4:3E:192:GLU:N	2.47	0.47
24:3L:84:A:H2'	24:3L:85:C:C6	2.49	0.47
31:41:73:ALA:HB2	31:41:88:ILE:HD11	1.97	0.47
5:42:60:TYR:HB3	5:42:64:ARG:NH2	2.25	0.47
32:51:92:ILE:HG13	32:51:92:ILE:H	1.37	0.47
32:59:11:VAL:HB	32:59:13:LYS:HE2	1.97	0.47
32:59:96:ALA:H	32:59:128:PRO:HA	1.79	0.47
15:6A:39:LEU:CD1	15:6A:56:LEU:HB2	2.44	0.47
26:1H:1250:G:OP2	36:78:21:ARG:NH1	2.47	0.47
37:88:138:ASP:OD1	37:88:138:ASP:N	2.45	0.47
42:95:66:ARG:HB2	42:95:88:ARG:HB3	1.95	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:B5:50:LYS:HB2	44:B5:87:GLN:HE22	1.79	0.47
20:BA:56:MET:HG3	20:BA:84:LEU:HD22	1.96	0.47
41:C8:8:VAL:O	41:C8:12:ARG:HG3	2.15	0.47
47:E5:50:ASN:O	47:E5:62:LEU:HB2	2.15	0.47
48:F5:44:PRO:HB2	48:F5:46:LEU:CD1	2.44	0.47
28:11:2:ALA:HA	28:11:20:ASP:HB3	1.97	0.47
28:11:238:GLY:O	28:11:239:ARG:C	2.51	0.47
1:13:1269:A:H2	1:13:1312:G:N3	2.13	0.47
1:13:191(F):U:H2'	1:13:191:G:C8	2.50	0.47
26:14:996:A:N6	26:14:1160:G:C6	2.83	0.47
26:14:1588:C:H5'	26:14:1589:C:OP2	2.13	0.47
26:14:1759:A:H4'	26:14:2715:C:O4'	2.14	0.47
26:14:2889:C:H3'	26:14:2891:G:C8	2.49	0.47
26:14:481:G:OP2	45:C5:47:LYS:HD3	2.14	0.47
34:15:96:GLU:H	34:15:96:GLU:CD	2.17	0.47
27:16:32:C:C2	27:16:51:G:N2	2.82	0.47
10:1A:30:SER:CB	10:1A:81:THR:HA	2.44	0.47
2:1E:12:GLU:C	2:1E:16:HIS:HD2	2.18	0.47
1:1G:110:C:H2'	1:1G:111:G:O4'	2.14	0.47
1:1G:1278:U:H5'	1:1G:1279:A:C5'	2.44	0.47
1:1G:713:G:H2'	1:1G:714:G:C8	2.50	0.47
1:1G:652:U:C5	1:1G:752:G:N3	2.83	0.47
26:1H:1794:U:H2'	26:1H:1795:C:H6	1.78	0.47
26:1H:2271:G:H5''	47:I8:20:ARG:HG2	1.96	0.47
26:1H:2643:G:H2'	26:1H:2644:G:O4'	2.14	0.47
26:1H:439:G:H2'	26:1H:440:G:C8	2.50	0.47
26:1H:731:C:H2'	26:1H:732:C:C6	2.47	0.47
29:21:170:LEU:HD21	29:21:187:ALA:HB3	1.97	0.47
3:22:109:PRO:HB2	3:22:115:LEU:HG	1.97	0.47
23:2L:65:5MU:C4	23:2L:66:PSU:C2	3.03	0.47
30:31:149:ASP:OD1	30:31:149:ASP:N	2.42	0.47
1:1G:407:G:O2'	4:32:116:GLN:HG3	2.15	0.47
4:3E:110:PHE:HE2	4:3E:148:VAL:HG23	1.79	0.47
12:3I:55:VAL:HG12	12:3I:69:TYR:HA	1.97	0.47
12:3I:57:LYS:HE3	12:3I:65:GLU:HG2	1.96	0.47
24:3L:27:G:H2'	24:3L:28:C:O4'	2.13	0.47
39:65:86:ALA:O	39:65:87:PHE:HB2	2.15	0.47
41:85:49:HIS:O	41:85:53:ARG:N	2.48	0.47
9:8E:25:LYS:HD3	9:8E:60:ASP:HB3	1.97	0.47
41:C8:105:VAL:HG22	42:D8:45:THR:HG21	1.97	0.47
45:G8:5:MET:HE1	45:G8:32:PRO:HA	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:737:A:H2'	1:13:738:C:C6	2.49	0.47
26:14:1259:G:H2'	26:14:1260:G:C8	2.50	0.47
26:14:1482:U:H5'	26:14:1483:G:OP2	2.14	0.47
26:14:1666:G:OP1	35:25:66:LYS:HD3	2.15	0.47
26:14:1945:G:H2'	26:14:1946:U:H6	1.79	0.47
26:14:528:A:H2	26:14:2043:C:H5'	1.80	0.47
26:14:2273:A:O2'	26:14:2274:A:H5'	2.15	0.47
26:14:873:G:C2	26:14:905:U:O2	2.67	0.47
28:19:70:TRP:C	28:19:70:TRP:CD1	2.88	0.47
2:1E:5:ILE:HG13	2:1E:6:THR:H	1.79	0.47
1:1G:1162:C:H42	1:1G:1174:G:H1	1.62	0.47
1:1G:286:G:H2'	1:1G:287:U:H6	1.80	0.47
1:1G:562:C:H4'	1:1G:563:A:O5'	2.14	0.47
26:1H:1512:G:C5	26:1H:1513:C:C4	3.03	0.47
26:1H:828:U:H4'	26:1H:831:G:N1	2.30	0.47
3:22:135:LYS:O	3:22:139:GLN:HB2	2.15	0.47
35:25:68:GLU:HB3	35:25:78:ARG:HH11	1.79	0.47
29:29:111:ARG:NH1	29:29:111:ARG:HG2	2.27	0.47
29:29:120:TRP:CG	29:29:155:LYS:HB3	2.50	0.47
11:2I:45:GLY:O	11:2I:50:TYR:HB2	2.15	0.47
4:32:163:GLU:HA	4:32:166:LYS:HG3	1.96	0.47
24:3K:7:G:H3'	24:3K:8:U:C5'	2.41	0.47
31:41:26:GLN:OE1	31:41:27:ASN:HB2	2.15	0.47
31:49:16:ARG:O	31:49:20:ILE:HG13	2.15	0.47
13:4A:86:CYS:O	13:4A:89:GLY:N	2.38	0.47
13:4I:107:ALA:O	13:4I:110:ARG:N	2.47	0.47
38:55:100:LEU:HD21	38:55:113:LEU:HD22	1.97	0.47
7:6E:121:ALA:O	7:6E:125:MET:HB2	2.15	0.47
7:6E:103:TRP:CE2	7:6E:137:LYS:HD3	2.50	0.47
40:75:3:ARG:HG2	40:75:4:GLY:N	2.30	0.47
39:A8:36:TYR:N	39:A8:36:TYR:CD1	2.82	0.47
44:B5:57:LEU:N	44:B5:57:LEU:HD23	2.29	0.47
41:C8:94:ASN:HB3	41:C8:95:LEU:HD22	1.96	0.47
45:G8:97:ARG:NH1	45:G8:104:GLY:HA3	2.30	0.47
48:J8:78:LYS:HE3	48:J8:78:LYS:H	1.80	0.47
1:13:1356:G:H2'	1:13:1357:A:C8	2.49	0.47
1:13:779:C:H2'	1:13:780:A:O4'	2.15	0.47
26:14:2147:G:C5	26:14:2148:G:H1'	2.50	0.47
26:14:2356:C:H4'	47:E5:20:ARG:HG3	1.97	0.47
26:14:395:U:H2'	26:14:396:G:N7	2.29	0.47
26:14:813:U:H2'	26:14:814:C:C6	2.50	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1104:G:O5'	2:1E:111:ARG:HD2	2.15	0.47
1:1G:595:G:H1	1:1G:641:U:HO2'	1.62	0.47
1:1G:690:G:H2'	1:1G:691:G:O4'	2.15	0.47
26:1H:1204:A:H2	26:1H:1241:A:N1	2.13	0.47
26:1H:2377:A:H2'	26:1H:2378:A:C8	2.50	0.47
26:1H:2475:C:O2'	26:1H:2476:A:H5'	2.15	0.47
26:1H:2689:U:H5''	26:1H:2713:A:C2	2.50	0.47
26:1H:997:G:O2'	26:1H:998:C:H5'	2.14	0.47
27:1J:44:G:C2	27:1J:48:A:C2	3.03	0.47
22:1K:49:C:H2'	22:1K:50:A:H4'	1.96	0.47
22:1L:15:A:H8	22:1L:16:U:H5	1.60	0.47
3:2E:150:LYS:HD2	3:2E:201:TYR:HD2	1.80	0.47
5:42:92:LYS:N	5:42:119:LEU:O	2.41	0.47
5:4E:43:LEU:HD21	5:4E:132:ALA:HB1	1.97	0.47
39:65:30:ARG:HD2	39:65:97:ARG:HD3	1.96	0.47
20:BI:50:GLU:HB2	20:BI:100:ILE:HB	1.97	0.47
42:D8:79:VAL:CG1	42:D8:81:TYR:HB3	2.45	0.47
47:E5:51:VAL:N	47:E5:62:LEU:HD12	2.30	0.47
1:13:1175:G:C6	1:13:1176:A:C2	3.03	0.47
1:13:1186:G:H21	14:5I:61:TRP:HA	1.80	0.47
1:13:313:A:H2'	1:13:314:C:C6	2.50	0.47
1:13:554:C:H2'	1:13:555:C:H6	1.79	0.47
1:13:560:U:O2'	1:13:561:U:OP2	2.30	0.47
1:13:865:A:H2	1:13:918:A:H4'	1.80	0.47
26:14:1997:G:H8	57:14:3984:HOH:O	1.97	0.47
26:14:2432:A:H2'	26:14:2433:A:C8	2.50	0.47
26:14:270(P):C:H2'	26:14:270(Q):C:C6	2.50	0.47
26:14:38:A:H2'	26:14:39:C:H6	1.78	0.47
28:19:33:LEU:HD21	28:19:103:ARG:HA	1.97	0.47
2:1E:17:PHE:CD1	2:1E:17:PHE:N	2.83	0.47
1:1G:687:A:C6	1:1G:704:A:N7	2.83	0.47
1:1G:976:G:OP1	14:5A:31:ARG:HB3	2.15	0.47
26:1H:1105:U:H2'	26:1H:1106:G:H8	1.80	0.47
26:1H:1516:U:H2'	26:1H:1517:G:C8	2.50	0.47
26:1H:2359:C:H2'	26:1H:2360:A:O4'	2.15	0.47
26:1H:2474:C:H3'	26:1H:2475:C:C6	2.50	0.47
26:1H:458:G:O2'	26:1H:469:G:O6	2.24	0.47
27:1J:11:C:H3'	27:1J:12:C:C6	2.50	0.47
23:2K:18:OMG:H1'	23:2K:19:G:OP2	2.15	0.47
23:2K:44:G:O2'	23:2K:45:U:OP1	2.25	0.47
30:39:160:ASN:HB3	30:39:163:VAL:HB	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:3E:122:ARG:NH1	4:3E:122:ARG:HG2	2.30	0.47
32:59:125:VAL:HG22	32:59:126:PRO:HA	1.97	0.47
32:59:41:MET:HB3	32:59:42:ARG:H	1.61	0.47
1:1G:979:C:O2'	14:5A:19:ARG:NH2	2.47	0.47
14:5A:39:LEU:HD13	14:5A:47:LEU:HD12	1.97	0.47
14:5I:27:CYS:HB2	14:5I:29:ARG:H	1.80	0.47
8:72:94:TYR:CE1	8:72:132:GLU:HB2	2.49	0.47
9:82:95:LYS:HE2	9:82:95:LYS:HB2	1.50	0.47
26:1H:956:G:OP2	37:88:14:ARG:NH2	2.48	0.47
41:85:92:ARG:NH1	42:95:11:GLN:O	2.48	0.47
42:95:2:PHE:H	42:95:42:GLY:HA3	1.79	0.47
39:A8:99:LYS:HE2	39:A8:103:GLU:OE1	2.15	0.47
19:AI:36:ARG:HD2	19:AI:52:TYR:O	2.15	0.47
44:B5:50:LYS:H	44:B5:83:VAL:HG23	1.78	0.47
42:D8:3:ALA:CB	42:D8:38:LEU:HD11	2.43	0.47
48:J8:23:LYS:HB3	48:J8:29:GLY:HA3	1.97	0.47
54:Q8:34:TRP:HA	54:Q8:35:GLN:HA	1.55	0.47
28:11:46:GLN:C	28:11:48:ARG:H	2.19	0.46
2:12:84:GLU:O	2:12:219:VAL:HG11	2.15	0.46
1:13:1113:C:H2'	1:13:1114:C:C6	2.50	0.46
1:13:539:A:H2'	1:13:540:G:C8	2.50	0.46
26:14:1160:G:C5	26:14:1161:C:C5	3.03	0.46
26:14:1484:G:H2'	26:14:1485:G:H8	1.80	0.46
26:14:1777:U:O2'	26:14:1778:U:H5'	2.15	0.46
26:14:2136:C:H5	26:14:2156:G:H21	1.62	0.46
26:14:2674:G:H4'	35:25:30:ALA:HB2	1.97	0.46
26:14:2695:C:H2'	26:14:2696:U:H6	1.80	0.46
26:14:586:A:N1	26:14:809:G:O2'	2.39	0.46
26:14:676:A:H8	26:14:2069:G:N2	1.94	0.46
26:14:733:G:O6	26:14:761:A:C8	2.68	0.46
26:14:881:G:N7	26:14:882:G:C2	2.83	0.46
28:19:70:TRP:CH2	28:19:150:LYS:HA	2.50	0.46
28:19:16:MET:HE1	28:19:208:LYS:HG2	1.96	0.46
2:1E:22:LYS:C	2:1E:24:TRP:H	2.17	0.46
2:1E:85:ALA:HB3	2:1E:92:TYR:CD2	2.50	0.46
1:1G:187:C:O2	1:1G:191(A):G:N1	2.48	0.46
1:1G:345:C:HO2'	1:1G:346:G:P	2.36	0.46
1:1G:616:G:C2	1:1G:617:G:C8	3.03	0.46
26:1H:1283:G:N2	26:1H:1285:G:H3'	2.29	0.46
26:1H:2843:G:H2'	26:1H:2844:G:H8	1.79	0.46
27:1J:80:U:H2'	27:1J:81:G:N2	2.26	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:22:136:GLN:O	3:22:139:GLN:N	2.48	0.46
3:2E:33:LEU:O	3:2E:36:ASP:HB2	2.15	0.46
4:32:119:GLN:HE21	4:32:123:HIS:CD2	2.33	0.46
4:32:4:TYR:CE2	4:32:11:LEU:HD11	2.50	0.46
4:3E:85:LYS:CG	4:3E:86:LYS:H	2.27	0.46
4:32:57:ARG:HH22	5:42:107:ARG:CD	2.29	0.46
31:49:118:ARG:N	31:49:118:ARG:HD2	2.30	0.46
32:51:131:VAL:HG22	32:51:132:ARG:H	1.80	0.46
32:51:157:TYR:HE1	32:51:172:LYS:HB2	1.79	0.46
32:51:8:PRO:O	32:51:10:PRO:HD3	2.15	0.46
34:58:28:THR:O	34:58:31:ALA:HB3	2.15	0.46
33:61:144:VAL:HG22	33:61:145:VAL:N	2.31	0.46
9:82:113:LYS:HD3	9:82:119:ALA:HB1	1.97	0.46
17:8I:4:LYS:HG3	17:8I:6:LEU:HD21	1.96	0.46
17:8I:93:GLN:O	17:8I:96:GLU:HG3	2.16	0.46
19:AI:6:LYS:HZ3	19:AI:6:LYS:HA	1.80	0.46
20:BI:10:LEU:HD21	20:BI:12:ALA:HB3	1.97	0.46
45:C5:59:GLY:O	45:C5:61:ILE:HG13	2.15	0.46
46:D5:94:GLU:O	46:D5:129:SER:HA	2.15	0.46
48:J8:90:ILE:HA	48:J8:94:LEU:HD12	1.96	0.46
48:J8:89:GLU:O	48:J8:93:GLU:HB2	2.15	0.46
48:J8:92:LYS:H	48:J8:95:LEU:HD12	1.78	0.46
26:14:2351:G:OP2	54:M5:46:ARG:NE	2.48	0.46
1:13:926:G:C6	1:13:1505:G:C5	3.03	0.46
1:13:582:U:H2'	1:13:583:A:C8	2.50	0.46
26:14:1485:G:H2'	26:14:1486:A:C8	2.50	0.46
26:14:1542:G:O6	26:14:1543:A:N6	2.48	0.46
26:14:2210:G:H3'	26:14:2211:G:C2	2.49	0.46
26:14:2791:C:H42	26:14:2807:G:H22	1.62	0.46
26:14:307:G:N2	26:14:310:A:OP2	2.46	0.46
26:14:959:A:C6	26:14:960:A:N1	2.84	0.46
1:1G:1275:A:H2'	1:1G:1276:G:O4'	2.15	0.46
1:1G:1541:U:H2'	1:1G:1542:U:C6	2.49	0.46
1:1G:280:C:N3	17:8A:39:SER:OG	2.40	0.46
1:1G:63:C:OP1	1:1G:384:G:N2	2.42	0.46
26:1H:1593:G:H2'	26:1H:1594:G:C8	2.49	0.46
26:1H:2306:C:H3'	26:1H:2307:G:C5'	2.45	0.46
26:1H:239:U:H2'	26:1H:240:G:C8	2.50	0.46
26:1H:340:A:H2'	26:1H:341:G:O4'	2.14	0.46
26:1H:478:A:C6	26:1H:480:A:C6	3.03	0.46
29:21:33:VAL:CG1	29:21:89:ASP:HA	2.45	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:22:21:ARG:O	3:22:58:GLU:HA	2.15	0.46
35:25:13:ASN:HD21	35:25:97:ARG:H	1.62	0.46
29:29:8:LYS:HB3	29:29:193:GLY:N	2.23	0.46
26:14:617:G:H5'	30:39:40:GLN:HE22	1.80	0.46
12:3I:8:ASN:O	12:3I:12:ARG:HG3	2.14	0.46
31:41:52:ILE:HA	31:41:52:ILE:HD13	1.79	0.46
37:45:55:VAL:HG23	37:45:64:ILE:HD12	1.98	0.46
13:4A:69:GLU:O	13:4A:73:GLU:HB2	2.15	0.46
13:4A:81:LEU:O	13:4A:89:GLY:HA3	2.15	0.46
13:4I:65:LYS:O	13:4I:66:LEU:HD23	2.15	0.46
1:1G:976:G:OP1	14:5A:32:SER:N	2.48	0.46
33:61:55:ALA:HA	33:61:58:LEU:HD22	1.97	0.46
33:69:115:ALA:HB2	33:69:131:LYS:HE3	1.98	0.46
8:72:6:ILE:O	8:72:10:LEU:HG	2.14	0.46
8:72:29:SER:HB3	8:72:32:LYS:CG	2.45	0.46
40:75:1:MET:SD	40:75:6:LEU:HD13	2.55	0.46
36:78:46:LYS:O	36:78:47:ASP:HB3	2.15	0.46
37:88:133:ARG:O	37:88:134:ARG:HB2	2.15	0.46
18:9A:22:VAL:O	18:9A:23:LYS:HB3	2.15	0.46
44:F8:57:LEU:HD23	44:F8:57:LEU:N	2.30	0.46
26:1H:2016:U:H1'	52:N8:6:VAL:HG13	1.97	0.46
28:11:26:LYS:HE2	28:11:94:LEU:HD12	1.98	0.46
2:12:42:ILE:CD1	2:12:202:PRO:HB2	2.45	0.46
1:13:134:A:H61	16:7I:25:ARG:NH1	2.13	0.46
26:14:185:U:H2'	26:14:186:G:H8	1.77	0.46
26:14:2335:A:C8	26:14:2337:G:C5	3.04	0.46
26:14:26:G:C6	26:14:27:G:N1	2.84	0.46
28:19:223:GLY:HA3	28:19:231:HIS:CE1	2.50	0.46
1:1G:1101:A:H4'	1:1G:1102:A:O5'	2.15	0.46
1:1G:19:C:OP1	5:42:125:SER:OG	2.28	0.46
1:1G:584:G:O6	57:1G:1860:HOH:O	2.20	0.46
1:1G:709:G:H2'	1:1G:710:G:C8	2.48	0.46
26:1H:1021:A:H8	26:1H:1022:G:H5''	1.80	0.46
26:1H:991:C:C5	26:1H:1185:C:N4	2.83	0.46
26:1H:1401:G:H2'	26:1H:1402:C:H6	1.79	0.46
26:1H:1971:A:C4	28:11:241:PRO:HD3	2.50	0.46
26:1H:2309:A:C6	26:1H:2310:A:H8	2.32	0.46
26:1H:2680:C:H5'	29:21:189:PRO:HA	1.97	0.46
26:1H:548:A:C6	26:1H:549:G:H1'	2.51	0.46
27:1J:0:A:H2'	27:1J:1:U:C6	2.50	0.46
29:29:179:GLU:O	29:29:180:ASN:HB2	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:2E:156:ARG:HB3	3:2E:160:ALA:O	2.15	0.46
11:2I:91:ARG:O	11:2I:95:ILE:HG13	2.16	0.46
30:31:120:GLU:HB3	30:31:122:LYS:HG2	1.97	0.46
1:13:542:G:H5'	4:3E:41:GLY:HA3	1.98	0.46
37:45:25:ASP:CB	37:45:102:VAL:H	2.28	0.46
37:45:37:LEU:HD21	37:45:130:LYS:HB3	1.95	0.46
42:95:72:VAL:HG13	42:95:72:VAL:O	2.16	0.46
38:98:52:ILE:O	38:98:55:ALA:N	2.49	0.46
43:A5:110:LYS:HA	43:A5:110:LYS:HD2	1.55	0.46
19:AA:10:PHE:O	51:I5:63:TYR:OH	2.33	0.46
19:AA:35:SER:O	19:AA:71:LEU:HD12	2.15	0.46
45:G8:10:GLY:O	45:G8:26:LYS:HE2	2.15	0.46
46:H8:151:HIS:HB3	46:H8:168:GLU:HA	1.98	0.46
28:11:165:ILE:H	28:11:165:ILE:HG12	1.57	0.46
1:13:1118:C:H1'	1:13:1179:A:C4	2.50	0.46
1:13:153:C:H42	1:13:168:G:H1	1.62	0.46
1:13:455:C:H42	1:13:477:G:H1	1.64	0.46
26:14:1312:U:H4'	26:14:1313:U:O5'	2.15	0.46
26:14:1386:C:OP2	26:14:1396:U:H5	1.98	0.46
26:14:2159:G:C2	26:14:2160:G:H1'	2.51	0.46
26:14:2169:A:N3	26:14:2169:A:H3'	2.30	0.46
26:14:2515:C:O2	26:14:2570:G:C2	2.69	0.46
26:14:2596:U:H2'	26:14:2597:G:O4'	2.16	0.46
26:14:2745:C:H1'	32:59:143:GLN:HG2	1.97	0.46
26:14:666:G:H5''	36:35:47:ASP:O	2.15	0.46
27:16:73:A:N3	27:16:73:A:H2'	2.30	0.46
28:19:148:GLU:HB2	28:19:151:LYS:HD2	1.97	0.46
10:1A:40:LEU:HG	10:1A:41:PRO:HD2	1.96	0.46
2:1E:92:TYR:CE2	2:1E:151:GLY:HA3	2.50	0.46
1:1G:1022:G:N3	1:1G:1023:G:H1'	2.30	0.46
1:1G:1412:C:H2'	1:1G:1413:A:H8	1.81	0.46
1:1G:1453:G:H1	20:BA:54:LYS:HZ2	1.63	0.46
1:1G:973:G:H3'	1:1G:974:A:H5''	1.98	0.46
26:1H:1204:A:C2	26:1H:1241:A:N1	2.84	0.46
26:1H:1337:G:H2'	26:1H:1338:G:C8	2.50	0.46
26:1H:1416:G:HO2'	26:1H:1417:C:H6	1.62	0.46
26:1H:1581:G:C6	26:1H:1582:C:C4	3.04	0.46
26:1H:1709:U:H2'	26:1H:1710:C:H6	1.80	0.46
26:1H:194:G:H2'	26:1H:195:A:O4'	2.16	0.46
26:1H:2291:U:H5''	26:1H:2380:C:O2'	2.15	0.46
26:1H:2543:G:H2'	26:1H:2544:G:O4'	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2564:A:C2	26:1H:2647:U:H4'	2.49	0.46
26:1H:443:A:O2'	26:1H:1200:C:O2'	2.16	0.46
26:1H:443:A:H5''	26:1H:444:C:OP1	2.15	0.46
22:1L:14:A:H1'	22:1L:23:A:C6	2.50	0.46
29:29:5:LEU:CD2	29:29:49:LEU:HD12	2.45	0.46
11:2I:103:LEU:O	11:2I:105:VAL:HG12	2.15	0.46
4:32:150:GLU:C	4:32:152:SER:N	2.69	0.46
30:39:177:ALA:HB1	30:39:178:PRO:HD2	1.97	0.46
30:39:36:VAL:O	30:39:40:GLN:HB2	2.15	0.46
31:49:113:ARG:NH1	31:49:142:PRO:HB3	2.31	0.46
32:51:170:ARG:O	32:51:171:LEU:HD12	2.14	0.46
14:5A:47:LEU:HA	14:5A:47:LEU:HD23	1.80	0.46
1:1G:1374:A:O2'	7:62:28:ASN:HB3	2.15	0.46
35:68:75:SER:CB	40:B8:74:ARG:HH12	2.28	0.46
8:72:20:TYR:HA	8:72:65:TYR:CZ	2.51	0.46
39:A8:11:LYS:O	39:A8:15:ARG:HB2	2.15	0.46
19:AA:17:GLU:O	19:AA:21:GLU:HG2	2.15	0.46
19:AI:18:LYS:HD3	19:AI:31:ILE:HB	1.97	0.46
44:B5:57:LEU:HD11	44:B5:78:LYS:HD2	1.97	0.46
20:BA:89:ARG:NH1	20:BA:105:SER:O	2.38	0.46
51:M8:49:PHE:HD1	51:M8:50:VAL:HB	1.80	0.46
13:4I:65:LYS:HE2	51:M8:50:VAL:HG13	1.97	0.46
28:11:27:THR:HG23	28:11:28:GLU:HG2	1.97	0.46
2:12:189:ASP:HB3	2:12:203:GLY:O	2.15	0.46
1:13:1230:C:H2'	1:13:1231:G:H8	1.79	0.46
1:13:13:U:O2	1:13:914:A:H3'	2.16	0.46
1:13:163:C:H2'	1:13:164:U:C4	2.51	0.46
1:13:22:G:C6	1:13:23:C:C4	3.04	0.46
1:13:542:G:OP1	4:3E:10:ARG:NH2	2.45	0.46
26:14:1094:U:H4'	26:14:1097:U:C5	2.51	0.46
26:14:1171:G:N2	26:14:1173:G:HO2'	2.13	0.46
26:14:2101:G:N2	26:14:2189:U:O2	2.48	0.46
26:14:2308:G:O2'	26:14:2310:A:H2	1.98	0.46
26:14:2694:G:C4	26:14:2695:C:C5	3.04	0.46
26:14:382:G:H1	26:14:392:C:H42	1.64	0.46
26:14:620:G:H4'	26:14:621:A:C5'	2.42	0.46
26:14:775:G:C5	26:14:794:G:C8	3.04	0.46
26:14:815:C:H2'	26:14:816:C:H6	1.81	0.46
26:14:863:A:H2'	26:14:864:G:C8	2.50	0.46
26:14:7:G:H2'	26:14:8:A:C8	2.50	0.46
1:1G:1011:G:N1	1:1G:1019:C:O2	2.49	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1281:U:H3'	1:1G:1282:C:C5	2.50	0.46
1:1G:410:G:N1	1:1G:429:U:O2	2.48	0.46
1:1G:487:A:H2'	1:1G:488:C:O4'	2.15	0.46
1:1G:807:A:H2'	1:1G:808:C:C6	2.51	0.46
26:1H:1044:G:O2'	26:1H:1111:A:N6	2.47	0.46
26:1H:1162:G:H1'	42:D8:23:GLU:OE2	2.16	0.46
26:1H:1729:A:H8	26:1H:1730:U:C6	2.33	0.46
26:1H:2188:C:H2'	26:1H:2189:U:O4'	2.16	0.46
26:1H:26:G:C6	26:1H:27:G:N1	2.83	0.46
26:1H:329:G:H4'	26:1H:330:A:OP2	2.15	0.46
26:1H:824:A:H1'	26:1H:2358:G:N7	2.30	0.46
26:1H:884:C:O2	26:1H:892:G:N2	2.49	0.46
29:21:101:ARG:HG2	29:21:169:ASN:OD1	2.16	0.46
29:21:167:VAL:HG21	29:21:187:ALA:HB1	1.98	0.46
29:29:116:VAL:HG23	29:29:120:TRP:HD1	1.80	0.46
11:2A:27:ASN:OD1	11:2A:28:THR:N	2.45	0.46
3:2E:3:ASN:OD1	3:2E:3:ASN:N	2.48	0.46
23:2K:20:U:H1'	23:2K:21:A:OP2	2.16	0.46
30:31:116:ASP:HA	30:31:119:ARG:NH2	2.31	0.46
1:1G:438:G:H4'	4:32:123:HIS:CE1	2.51	0.46
12:3A:70:ILE:HD13	12:3A:77:LEU:HD12	1.97	0.46
4:3E:25:ARG:NH1	4:3E:30:LYS:O	2.48	0.46
1:13:562:C:H1'	12:3I:15:ARG:HB3	1.97	0.46
37:45:68:ILE:HG12	37:45:103:MET:HB2	1.98	0.46
13:4A:32:GLU:O	13:4A:35:GLU:HG2	2.15	0.46
32:51:54:ARG:HG3	32:51:56:SER:O	2.15	0.46
33:61:33:ARG:HB3	33:61:35:LEU:HD23	1.96	0.46
35:68:101:PRO:HB3	35:68:122:LEU:CD1	2.46	0.46
33:69:38:LEU:HD12	33:69:38:LEU:H	1.81	0.46
15:6I:33:THR:HG21	15:6I:85:LEU:HD22	1.98	0.46
16:7I:38:TYR:CE1	16:7I:50:LYS:HB2	2.50	0.46
1:13:1117:G:O3'	9:8E:104:ARG:HD3	2.14	0.46
9:8E:4:TYR:CZ	9:8E:88:TYR:HB2	2.51	0.46
42:95:24:LYS:HA	42:95:92:THR:OG1	2.15	0.46
45:C5:3:VAL:HG11	45:C5:32:PRO:HB2	1.97	0.46
41:C8:34:LYS:NZ	41:C8:37:GLU:OE1	2.40	0.46
41:C8:87:GLY:C	41:C8:89:GLU:H	2.19	0.46
45:G8:21:LYS:HE2	45:G8:21:LYS:HB3	1.68	0.46
54:Q8:32:LEU:CG	54:Q8:33:ASN:HD22	2.29	0.46
54:Q8:35:GLN:C	54:Q8:37:SER:N	2.68	0.46
2:12:214:ILE:O	2:12:218:ALA:HB2	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1028(A):C:H2'	1:13:1028(B):C:C5	2.47	0.46
1:13:103:C:H2'	1:13:104:G:H8	1.80	0.46
1:13:452:A:O2'	1:13:453:A:O4'	2.30	0.46
1:13:636:U:H2'	1:13:637:G:C8	2.50	0.46
26:14:1453:A:O2'	26:14:1454:U:H2'	2.15	0.46
26:14:2320:A:C6	26:14:2333:A:C8	3.04	0.46
26:14:2494:G:C4	26:14:2495:G:C8	3.04	0.46
1:1G:1288:A:H4'	21:1B:13:ILE:HD13	1.98	0.46
2:1E:238:LEU:H	2:1E:238:LEU:HD12	1.81	0.46
2:1E:60:ASP:HB3	2:1E:64:ARG:NH1	2.28	0.46
1:1G:1004:A:H3'	1:1G:1004:A:N3	2.30	0.46
1:1G:1328:C:H2'	1:1G:1329:A:C8	2.51	0.46
26:1H:1036:G:H1	26:1H:1119:C:H42	1.62	0.46
26:1H:1141:U:H6	34:58:63:THR:OG1	1.99	0.46
26:1H:1830:C:N4	26:1H:1975:G:H1	2.12	0.46
26:1H:2562:U:H1'	35:68:23:ARG:HD3	1.97	0.46
26:1H:2579:C:H2'	26:1H:2580:U:O4'	2.15	0.46
26:1H:288:C:H2'	26:1H:289:A:C8	2.51	0.46
26:1H:497:A:C5	26:1H:498:G:C8	3.04	0.46
26:1H:795:C:H2'	26:1H:796:C:C6	2.50	0.46
27:1J:70:C:H2'	27:1J:71:C:H6	1.79	0.46
3:22:47:LEU:HD12	3:22:52:LEU:HB3	1.96	0.46
36:35:77:ARG:HB2	36:35:78:PRO:HD2	1.98	0.46
31:49:53:LEU:HA	31:49:56:ALA:HB3	1.96	0.46
31:49:98:ARG:HE	31:49:98:ARG:HB2	1.49	0.46
5:4E:30:ALA:N	5:4E:46:GLY:O	2.46	0.46
13:4I:8:GLU:OE2	13:4I:22:ILE:HA	2.16	0.46
32:51:125:VAL:HG13	32:51:131:VAL:HG23	1.97	0.46
6:52:3:ARG:HB3	6:52:3:ARG:HH11	1.81	0.46
34:58:104:LYS:HB2	34:58:117:PHE:CE1	2.51	0.46
7:62:76:ARG:C	7:62:86:GLN:HB2	2.36	0.46
1:1G:1290:G:H21	9:82:70:LYS:NZ	2.13	0.46
17:8A:67:LYS:O	17:8A:68:ARG:HB3	2.16	0.46
44:B5:25:LYS:HA	44:B5:81:VAL:O	2.15	0.46
1:13:324:G:OP1	20:BI:22:ARG:HG2	2.16	0.46
44:F8:44:GLU:HG2	44:F8:49:VAL:O	2.15	0.46
46:H8:141:VAL:O	46:H8:144:LEU:HG	2.16	0.46
51:M8:39:CYS:SG	51:M8:41:PRO:HD3	2.56	0.46
2:12:114:ARG:NH1	2:12:141:GLU:OE2	2.48	0.46
2:12:165:VAL:HG23	2:12:166:ASP:N	2.31	0.46
1:13:429:U:H1'	1:13:430:A:H5''	1.96	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:443:C:H2'	1:13:444:C:H6	1.81	0.46
1:13:491:G:H2'	1:13:492:G:H8	1.81	0.46
1:13:514:C:H2'	1:13:515:G:C8	2.51	0.46
26:14:1022:G:C6	26:14:1140:C:C4	3.03	0.46
26:14:330:A:H2	26:14:1210:A:HO2'	1.59	0.46
26:14:208:C:H2'	26:14:209:C:H6	1.81	0.46
1:13:1032:A:N1	26:14:2114:A:H5''	2.30	0.46
26:14:2252:G:H2'	26:14:2253:G:O4'	2.16	0.46
26:14:2745:C:H2'	26:14:2746:U:C6	2.51	0.46
26:14:789:A:H3'	57:14:3663:HOH:O	2.16	0.46
26:14:818:G:H4'	26:14:838:C:O3'	2.15	0.46
26:14:900:A:OP1	26:14:900:A:H4'	2.14	0.46
34:15:114:ARG:O	34:15:118:LYS:HG3	2.16	0.46
1:1G:1023:G:H3'	1:1G:1024:G:H5''	1.97	0.46
1:1G:1132:C:H2'	1:1G:1133:G:C8	2.51	0.46
1:1G:1338:G:H2'	1:1G:1339:A:O4'	2.16	0.46
26:1H:2053:G:P	57:1H:3670:HOH:O	2.66	0.46
26:1H:2065:C:H2'	26:1H:2066:C:C6	2.51	0.46
26:1H:2287:A:H2	26:1H:2346:A:C2	2.34	0.46
26:1H:2360:A:C2	26:1H:2361:A:H1'	2.50	0.46
26:1H:2683:C:H5''	40:B8:53:ARG:HH12	1.81	0.46
26:1H:2764:A:N6	26:1H:2766:G:C2	2.84	0.46
26:1H:2769:C:N4	26:1H:2770:G:C5	2.84	0.46
22:1K:18:G:N2	22:1K:20:U:OP1	2.49	0.46
29:21:50:GLY:O	29:21:74:PRO:HG3	2.16	0.46
1:1G:1523:G:OP1	11:2A:123:LYS:HD2	2.15	0.46
24:3L:34:U:H2'	24:3L:36:A:OP2	2.16	0.46
31:41:106:LEU:HD12	31:41:110:ALA:HB3	1.97	0.46
37:45:81:VAL:HG23	37:45:82:ARG:N	2.31	0.46
31:49:64:THR:HA	31:49:102:PHE:HD1	1.81	0.46
32:51:150:ALA:O	32:51:153:LYS:HE2	2.16	0.46
33:69:92:VAL:O	33:69:120:ILE:HG12	2.15	0.46
15:6I:3:ILE:HD12	15:6I:34:LEU:HD23	1.97	0.46
8:7E:12:ARG:HD3	8:7E:26:VAL:HB	1.98	0.46
8:7E:43:GLY:O	8:7E:64:LYS:HD3	2.16	0.46
41:85:66:ASN:HD21	41:85:70:ARG:HE	1.62	0.46
40:B8:57:PHE:CE1	40:B8:79:HIS:HB2	2.51	0.46
41:C8:83:LEU:HD12	41:C8:88:ILE:HB	1.97	0.46
46:D5:48:PHE:CE1	46:D5:52:SER:HA	2.51	0.46
47:E5:38:VAL:HB	47:E5:59:LEU:HD12	1.98	0.46
26:1H:142:G:C1'	44:F8:37:THR:HG21	2.43	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
45:G8:61:ILE:HD13	45:G8:63:LYS:HD3	1.97	0.46
45:G8:76:CYS:C	45:G8:78:ALA:H	2.16	0.46
49:K8:2:LYS:O	49:K8:6:VAL:HG12	2.15	0.46
53:L5:10:ARG:O	53:L5:14:LYS:HB2	2.16	0.46
28:11:121:PRO:HB3	28:11:135:PHE:CE2	2.51	0.46
26:1H:1799:G:N2	28:11:155:LEU:HD12	2.31	0.46
1:13:191:G:C6	1:13:192:U:C4	3.04	0.46
1:13:243:A:H4'	1:13:244:U:C5'	2.46	0.46
1:13:347:G:N2	1:13:348:G:N3	2.64	0.46
1:13:413:G:H2'	1:13:428:G:N2	2.31	0.46
1:13:603:U:H2'	1:13:604:G:C8	2.51	0.46
26:14:1090:U:N3	26:14:1102:C:H1'	2.30	0.46
26:14:1338:G:N3	26:14:1393:A:H2	2.14	0.46
26:14:1462:C:H2'	26:14:1463:C:C6	2.51	0.46
26:14:1717:G:N2	26:14:1742:C:O2	2.44	0.46
26:14:1945:G:H2'	26:14:1946:U:C6	2.51	0.46
26:14:2607:G:H2'	26:14:2608:G:O4'	2.15	0.46
26:14:360:G:H8	26:14:360:G:O5'	1.98	0.46
26:14:914:C:N3	26:14:915:C:H1'	2.31	0.46
26:14:779:U:OP1	28:19:49:ILE:HG22	2.15	0.46
2:1E:74:LYS:NZ	2:1E:169:LYS:HG3	2.30	0.46
2:1E:8:LYS:HG2	2:1E:9:GLU:H	1.79	0.46
1:1G:1205:U:O2'	3:22:195:VAL:HG13	2.15	0.46
1:1G:1265:G:C2	1:1G:1271:G:C2	3.04	0.46
1:1G:1342:C:H4'	9:82:125:TYR:CB	2.44	0.46
1:1G:147:G:H1	1:1G:175:C:N4	2.03	0.46
1:1G:1540:U:H3	25:4L:33:G:H22	1.64	0.46
26:1H:1046:A:O2'	26:1H:1047:G:OP1	2.28	0.46
26:1H:1491:G:O2'	26:1H:1492:G:H5'	2.16	0.46
26:1H:1906:G:OP2	26:1H:1929:G:O2'	2.31	0.46
26:1H:2046:G:H1'	52:N8:22:HIS:HE1	1.81	0.46
35:25:47:ILE:HD12	35:25:47:ILE:HA	1.75	0.46
3:2E:136:GLN:HG2	3:2E:140:ARG:NH1	2.31	0.46
36:35:144:GLU:N	36:35:144:GLU:CD	2.69	0.46
24:3K:8:U:H4'	24:3K:8:U:OP1	2.16	0.46
37:45:48:GLU:O	37:45:52:VAL:HG23	2.15	0.46
5:4E:11:ILE:HD11	5:4E:31:LEU:HD13	1.98	0.46
5:4E:68:GLU:O	5:4E:70:PRO:HD3	2.15	0.46
32:51:90:LYS:HB3	32:51:90:LYS:HE2	1.75	0.46
16:7A:23:ASP:O	16:7A:25:ARG:N	2.48	0.46
42:95:69:LYS:HE3	42:95:86:GLY:HA3	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:98:102:GLU:O	38:98:111:LEU:HD12	2.15	0.46
18:9A:52:PRO:O	18:9A:56:THR:HG23	2.15	0.46
18:9I:58:LEU:HG	18:9I:62:GLU:HB3	1.97	0.46
43:A5:65:LEU:CD1	43:A5:68:ARG:HD3	2.45	0.46
19:AI:18:LYS:HG2	19:AI:22:LEU:HD13	1.98	0.46
45:G8:9:LYS:O	45:G8:27:VAL:HG22	2.16	0.46
51:I5:20:ASN:ND2	51:I5:39:CYS:SG	2.89	0.46
51:M8:13:ARG:NH1	51:M8:22:ILE:HG23	2.31	0.46
54:Q8:59:LYS:HB2	54:Q8:60:LEU:HG	1.97	0.46
2:12:22:LYS:H	2:12:22:LYS:HG3	1.60	0.46
26:14:1171:G:H4'	26:14:1173:G:OP1	2.16	0.46
26:14:1448:G:H1'	26:14:1528:A:H62	1.81	0.46
26:14:2131:G:H5''	26:14:2158:A:N1	2.31	0.46
26:14:2093:G:N2	26:14:2196:C:O2	2.40	0.46
26:14:2197:U:H1'	26:14:2198:A:C8	2.51	0.46
26:14:2658:C:H5''	32:59:158:HIS:HE1	1.80	0.46
26:14:654(C):G:H2'	26:14:654(D):G:O4'	2.16	0.46
26:14:6:A:N3	26:14:6:A:H2'	2.31	0.46
34:15:17:ASP:O	34:15:18:ALA:HB3	2.16	0.46
1:1G:1109:C:H2'	1:1G:1110:A:O4'	2.16	0.46
1:1G:15:G:H2'	1:1G:16:A:O4'	2.16	0.46
1:1G:722:A:H4'	1:1G:723:U:OP1	2.16	0.46
1:1G:980:C:H5'	1:1G:981:U:OP2	2.16	0.46
1:1G:999:U:H2'	1:1G:1000:A:C8	2.51	0.46
26:1H:1021:A:C3'	26:1H:1021:A:C8	2.99	0.46
26:1H:1021:A:N6	26:1H:1141:U:C2	2.84	0.46
26:1H:1291:C:H2'	26:1H:1292:U:C6	2.51	0.46
26:1H:1678:G:O5'	26:1H:1678:G:H8	1.99	0.46
26:1H:2323:G:H2'	26:1H:2324:C:O4'	2.16	0.46
26:1H:2553:G:H2'	26:1H:2554:U:C4'	2.46	0.46
26:1H:2815:C:H2'	26:1H:2816:C:H6	1.81	0.46
26:1H:320:A:H2'	30:31:136:THR:HG21	1.97	0.46
26:1H:534:U:H5'	41:C8:42:ALA:HB1	1.97	0.46
26:1H:846:C:C4	26:1H:930:U:C4	3.04	0.46
1:13:1152:A:H4'	10:1I:13:HIS:CD2	2.50	0.46
26:1H:2572:A:N7	29:21:145:LYS:HB2	2.31	0.46
3:22:84:ILE:HD11	3:22:88:ARG:NH2	2.28	0.46
11:2A:122:LYS:HD3	11:2A:124:LYS:HE2	1.98	0.46
11:2A:23:ALA:HB1	11:2A:88:GLY:HA3	1.98	0.46
30:31:134:GLY:H	30:31:162:LEU:HB3	1.80	0.46
4:32:176:LEU:HG	4:32:178:VAL:HG13	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:32:9:CYS:HA	4:32:12:CYS:HB2	1.97	0.46
30:39:133:ASN:HA	30:39:162:LEU:HD22	1.98	0.46
30:39:78:ILE:C	30:39:80:ALA:H	2.19	0.46
24:3K:37:G:H2'	24:3K:38:G:O4'	2.16	0.46
34:58:60:ILE:HG13	34:58:60:ILE:H	1.45	0.46
6:5E:62:TRP:CH2	6:5E:64:GLN:HG2	2.51	0.46
35:68:43:VAL:HG12	35:68:54:GLU:HA	1.98	0.46
35:68:7:TYR:CE1	35:68:20:MET:HE3	2.50	0.46
17:8A:92:ARG:HG2	17:8A:95:TYR:CE2	2.51	0.46
42:95:20:LEU:HD12	42:95:20:LEU:HA	1.64	0.46
42:95:34:GLU:OE1	42:95:56:SER:HB2	2.16	0.46
26:1H:1277:G:O2'	38:98:24:GLN:HG2	2.15	0.46
41:C8:49:HIS:HA	41:C8:52:ARG:HG2	1.98	0.46
48:F5:85:LEU:HG	48:F5:85:LEU:H	1.25	0.46
46:H8:48:PHE:CE1	46:H8:71:VAL:HG11	2.51	0.46
46:H8:59:LEU:HD23	46:H8:59:LEU:HA	1.58	0.46
2:12:72:GLY:HA2	2:12:165:VAL:CG2	2.46	0.46
1:13:280:C:N3	17:8I:39:SER:N	2.62	0.46
1:13:587:G:O5'	1:13:587:G:H8	1.98	0.46
1:13:730:G:C5	1:13:731:G:H1'	2.51	0.46
1:13:953:G:H5'	1:13:965:A:H61	1.81	0.46
26:14:1225:C:H2'	26:14:1226:G:C8	2.51	0.46
26:14:1204:A:C2	26:14:1241:A:N1	2.84	0.46
26:14:2261:C:C6	47:E5:16:SER:HB3	2.50	0.46
26:14:1956:U:H1'	26:14:2552:U:OP1	2.16	0.46
26:14:2689:U:P	26:14:2719:G:H22	2.38	0.46
26:14:2897:U:H5'	26:14:2898:U:OP2	2.16	0.46
28:19:73:VAL:HG13	28:19:120:GLY:HA3	1.96	0.46
2:1E:74:LYS:HG2	2:1E:74:LYS:H	1.50	0.46
1:1G:1052:U:O2'	1:1G:1055:A:OP2	2.13	0.46
1:1G:1397:C:O2'	1:1G:1398:A:OP1	2.27	0.46
1:1G:623:C:C4	1:1G:624:C:C5	3.04	0.46
1:1G:636:U:H2'	1:1G:637:G:C8	2.47	0.46
26:1H:1095:A:N3	26:1H:1095:A:H2'	2.31	0.46
26:1H:1358:G:O2'	26:1H:1359:A:H5''	2.16	0.46
26:1H:1771:C:H1'	26:1H:1786:A:C8	2.50	0.46
26:1H:2251:G:H2'	26:1H:2252:G:H8	1.81	0.46
26:1H:2299:G:N1	26:1H:2318:G:H8	2.14	0.46
26:1H:2629:A:OP1	26:1H:2629:A:H4'	2.16	0.46
26:1H:2690:C:H5''	26:1H:2872:G:H21	1.81	0.46
26:1H:273(F):C:H3'	26:1H:274:G:H5''	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2769:C:C4	26:1H:2770:G:C5	3.04	0.46
26:1H:55:G:H2'	26:1H:56:A:H8	1.80	0.46
29:29:203:LYS:HG2	29:29:205:ALA:CB	2.46	0.46
3:2E:147:LYS:HA	3:2E:147:LYS:HD3	1.80	0.46
3:2E:43:LEU:O	3:2E:47:LEU:HB2	2.16	0.46
4:32:27:TYR:HA	4:32:27:TYR:HD1	1.58	0.46
31:41:111:LEU:HA	31:41:114:ILE:HD12	1.97	0.46
26:14:2275:C:O2	37:45:85:LYS:HE3	2.16	0.46
34:58:40:PRO:O	41:C8:64:ARG:HG2	2.16	0.46
32:59:99:VAL:HG13	32:59:100:GLY:H	1.79	0.46
14:5I:25:VAL:HG13	14:5I:38:GLY:O	2.15	0.46
27:1J:27:C:O3'	39:65:36:TYR:OH	2.34	0.46
39:65:14:VAL:HG21	39:65:90:GLY:O	2.16	0.46
16:7A:16:HIS:CD2	16:7A:16:HIS:N	2.84	0.46
17:8A:45:HIS:NE2	17:8A:47:PRO:HB3	2.30	0.46
17:8A:59:ILE:HG23	17:8A:71:PHE:HB3	1.98	0.46
18:9A:29:PHE:CD1	18:9A:29:PHE:N	2.84	0.46
20:BI:73:HIS:O	20:BI:76:ALA:HB3	2.16	0.46
47:E5:12:ASN:HA	47:E5:14:ARG:HH21	1.81	0.46
44:F8:1:MET:C	44:F8:3:THR:N	2.67	0.46
48:J8:62:VAL:HG23	48:J8:63:ALA:O	2.16	0.46
50:L8:6:VAL:HG12	50:L8:56:VAL:HG13	1.97	0.46
51:M8:12:ALA:HB3	51:M8:24:THR:HB	1.97	0.46
52:N8:30:LEU:HA	52:N8:41:PRO:O	2.15	0.46
54:Q8:6:THR:HG22	54:Q8:59:LYS:HE3	1.97	0.46
26:1H:1820:U:O2	28:11:202:LYS:HB3	2.16	0.45
28:11:3:VAL:HG13	28:11:17:THR:HG23	1.97	0.45
2:12:132:LYS:O	2:12:135:GLN:HB2	2.15	0.45
2:12:221:LEU:HA	2:12:224:GLN:HB3	1.98	0.45
1:13:159:G:O2'	1:13:161:A:N7	2.40	0.45
1:13:329:A:C4	1:13:332:G:C5	3.04	0.45
1:13:465:A:N7	1:13:467:G:C6	2.84	0.45
1:13:772:U:H2'	1:13:773:G:H8	1.81	0.45
1:13:993:G:H4'	1:13:994:A:OP2	2.16	0.45
26:14:1361:G:H1	26:14:1370:C:H42	1.63	0.45
26:14:218:A:H2	26:14:235:U:H4'	1.80	0.45
26:14:2469:A:H2	26:14:2481:G:N2	2.09	0.45
26:14:249:C:OP1	57:14:3515:HOH:O	2.20	0.45
26:14:2536:G:C6	26:14:2537:U:C4	3.04	0.45
26:14:568:U:O4	57:14:3989:HOH:O	2.20	0.45
27:16:111:U:H2'	27:16:112:G:C8	2.49	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:1A:45:ARG:HB3	10:1A:65:LEU:HB3	1.97	0.45
1:1G:1158:C:N3	1:1G:1160:G:C8	2.84	0.45
1:1G:1541:U:O2'	1:1G:1542:U:O4'	2.24	0.45
1:1G:695:A:H2	1:1G:787:A:HO2'	1.63	0.45
1:1G:986:A:H1'	19:AA:54:GLY:O	2.16	0.45
26:1H:1087:G:N2	26:1H:1102:C:O2	2.48	0.45
26:1H:225:A:O2'	26:1H:257:A:H4'	2.16	0.45
26:1H:2492:U:H2'	26:1H:2493:U:C6	2.51	0.45
26:1H:2695:C:H2'	26:1H:2696:U:H6	1.81	0.45
29:29:11:MET:HE3	29:29:187:ALA:H	1.80	0.45
11:2A:73:MET:SD	11:2A:103:LEU:HD13	2.56	0.45
23:2L:22:G:C2	23:2L:59:U:C2	3.04	0.45
30:31:199:TRP:NE1	30:31:203:GLN:OE1	2.49	0.45
4:32:59:ARG:NH2	4:32:66:ARG:HH12	2.14	0.45
1:13:437:U:C5'	4:3E:155:LEU:HD21	2.46	0.45
13:4I:77:ASN:HA	13:4I:80:ARG:HB3	1.98	0.45
25:4K:45:U:HO2'	25:4K:46:U:P	2.39	0.45
38:55:87:TYR:HE1	38:55:117:VAL:HG12	1.80	0.45
32:59:19:VAL:HG12	32:59:20:ALA:H	1.80	0.45
39:65:102:ALA:HA	39:65:105:ALA:HB3	1.98	0.45
15:6A:26:GLU:HG3	15:6A:77:ARG:NH1	2.31	0.45
40:75:62:THR:HG22	40:75:75:ILE:HG12	1.98	0.45
18:9A:29:PHE:HD1	18:9A:29:PHE:N	2.14	0.45
39:A8:26:LEU:HD22	39:A8:87:PHE:CD2	2.51	0.45
45:C5:48:ALA:HB3	45:C5:59:GLY:HA2	1.98	0.45
46:H8:93:ASP:HA	46:H8:130:PRO:HG2	1.98	0.45
28:11:3:VAL:HG12	28:11:3:VAL:O	2.15	0.45
1:13:1304:G:C5	1:13:1305:G:C6	3.05	0.45
1:13:1390:U:H2'	1:13:1391:U:C6	2.51	0.45
1:13:827:U:C5	1:13:872:A:N1	2.84	0.45
1:13:874:G:C6	1:13:875:C:N4	2.85	0.45
26:14:1199:U:H2'	26:14:1200:C:H6	1.80	0.45
26:14:2801:A:C5'	26:14:2895:U:H4'	2.46	0.45
26:14:392:C:H5''	26:14:409:C:H5''	1.97	0.45
26:14:676:A:H1'	26:14:2443:C:H1'	1.97	0.45
27:16:11:C:OP2	27:16:12:C:N4	2.44	0.45
28:19:175:LEU:HD12	28:19:185:VAL:HG21	1.98	0.45
2:1E:223:ILE:H	2:1E:223:ILE:HG12	1.53	0.45
1:1G:141:A:H1'	1:1G:182:U:C2	2.51	0.45
1:1G:410:G:H21	1:1G:432:A:H62	1.63	0.45
26:1H:1100:C:H2'	26:1H:1101:U:H6	1.82	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1321:A:H2'	26:1H:1322:A:O4'	2.15	0.45
26:1H:1354:A:H2'	26:1H:1355:G:O4'	2.17	0.45
26:1H:1997:G:P	57:1H:3785:HOH:O	2.74	0.45
26:1H:2275:C:H5'	26:1H:2275:C:H6	1.82	0.45
26:1H:2795:G:H3'	26:1H:2797:U:C5'	2.45	0.45
26:1H:2850:A:C2	26:1H:2851:A:C4	3.04	0.45
26:1H:479:A:C2	26:1H:481:G:H5'	2.52	0.45
26:1H:74:A:H3'	49:K8:51:ARG:HH22	1.80	0.45
26:1H:975:G:H1'	26:1H:990:A:C2	2.51	0.45
35:25:87:ILE:HD12	35:25:93:PRO:HA	1.98	0.45
11:2A:17:GLY:N	11:2A:79:SER:O	2.40	0.45
4:32:150:GLU:HA	4:32:153:ARG:HG2	1.99	0.45
32:59:27:LYS:HD2	32:59:32:GLU:HG2	1.97	0.45
7:62:27:ILE:HG21	7:62:40:ALA:HA	1.98	0.45
26:14:2864:G:OP1	40:75:119:LYS:HD3	2.16	0.45
16:7I:4:ILE:HA	16:7I:20:VAL:O	2.17	0.45
9:82:118:LYS:HB3	9:82:118:LYS:NZ	2.32	0.45
9:82:48:GLU:N	9:82:49:PRO:HD2	2.31	0.45
39:A8:14:VAL:HG21	39:A8:89:ARG:HD3	1.98	0.45
19:AA:79:THR:OG1	19:AA:79:THR:O	2.27	0.45
42:D8:25:LEU:HD12	42:D8:94:LEU:HD21	1.98	0.45
41:C8:112:ARG:HH21	42:D8:47:VAL:HG13	1.82	0.45
50:H5:9:VAL:HG22	50:H5:53:LEU:O	2.16	0.45
48:J8:78:LYS:HE3	48:J8:78:LYS:N	2.30	0.45
28:11:127:VAL:HA	28:11:193:VAL:HG22	1.98	0.45
26:1H:782:A:C2	28:11:226:MET:HG2	2.52	0.45
1:13:1413:A:H2	1:13:1487:G:H22	1.64	0.45
26:14:139:G:H1'	26:14:140:A:H2	1.82	0.45
26:14:1484:G:H2'	26:14:1485:G:C8	2.51	0.45
26:14:2392:A:OP2	54:M5:31:HIS:NE2	2.44	0.45
26:14:626:U:O2	36:35:105:LEU:HD23	2.16	0.45
26:14:724:U:H2'	26:14:725:G:O4'	2.17	0.45
26:14:774:A:H2	26:14:787:U:HO2'	1.60	0.45
34:15:34:LEU:HD21	34:15:120:LEU:HD13	1.97	0.45
26:14:1500:G:O2'	28:19:100:GLY:O	2.27	0.45
28:19:32:SER:OG	28:19:32:SER:O	2.34	0.45
28:19:41:GLY:HA3	28:19:43:ARG:HE	1.81	0.45
2:1E:125:PRO:O	2:1E:126:GLU:HB2	2.17	0.45
1:1G:1145:C:H5''	1:1G:1146:A:OP1	2.17	0.45
1:1G:1305:G:HO2'	1:1G:1306:A:P	2.38	0.45
1:1G:1327:C:OP1	21:1B:20:LYS:HB3	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:842:C:H1'	1:1G:848:C:N3	2.31	0.45
26:1H:1526:G:H2'	26:1H:1527:G:O4'	2.15	0.45
26:1H:2114:A:N3	26:1H:2114:A:H2'	2.31	0.45
26:1H:308:G:H2'	26:1H:309:G:C8	2.51	0.45
29:21:143:ASN:HB2	29:21:147:PRO:HD2	1.98	0.45
3:22:47:LEU:HG	3:22:68:VAL:HG11	1.97	0.45
11:2A:29:ILE:O	11:2A:29:ILE:HG12	2.16	0.45
11:2I:38:ASN:HA	11:2I:39:PRO:HD3	1.75	0.45
23:2K:81:G:O5'	23:2K:81:G:H8	2.00	0.45
4:32:30:LYS:HD3	4:32:35:ARG:HH11	1.81	0.45
30:39:53:THR:HG22	30:39:56:GLU:CG	2.47	0.45
4:3E:25:ARG:NH1	4:3E:30:LYS:HG3	2.31	0.45
31:49:37:VAL:HG22	31:49:159:VAL:HG12	1.97	0.45
13:4I:82:MET:O	13:4I:84:ILE:N	2.47	0.45
38:55:57:ARG:HB3	38:55:59:ASP:OD1	2.16	0.45
14:5A:53:LEU:HD23	14:5A:53:LEU:HA	1.72	0.45
16:7A:58:TYR:O	16:7A:61:SER:HB3	2.16	0.45
8:7E:121:ASP:HB2	8:7E:125:ARG:NH2	2.32	0.45
1:13:875:C:H4'	8:7E:18:ARG:NH1	2.31	0.45
1:1G:1119:C:OP2	9:82:9:ARG:NH2	2.50	0.45
44:B5:44:GLU:HG3	44:B5:51:VAL:CG2	2.47	0.45
26:14:480:A:H1'	45:C5:44:ILE:HG12	1.99	0.45
41:C8:18:LEU:HD23	41:C8:18:LEU:HA	1.51	0.45
46:H8:44:PHE:CE2	46:H8:86:VAL:HG11	2.52	0.45
28:11:79:VAL:HG21	28:11:111:LEU:HD11	1.98	0.45
2:12:208:ILE:HD12	2:12:239:VAL:HG11	1.97	0.45
2:12:9:GLU:HB2	2:12:217:ARG:NH1	2.30	0.45
2:12:70:PHE:H	2:12:92:TYR:HA	1.81	0.45
1:13:130:A:N7	17:8I:63:ARG:HB2	2.32	0.45
1:13:164:U:H2'	1:13:165:C:C6	2.50	0.45
1:13:767:A:H2'	1:13:768:A:O4'	2.16	0.45
26:14:1464:C:O2'	26:14:1528:A:H8	1.98	0.45
26:14:1619:G:N7	57:14:3602:HOH:O	2.35	0.45
26:14:1701:A:H5''	26:14:1702:G:OP2	2.16	0.45
26:14:2111:C:H42	26:14:2147:G:N2	2.15	0.45
26:14:2195:C:O2'	26:14:2196:C:H5'	2.16	0.45
26:14:31:C:N4	57:14:3807:HOH:O	2.48	0.45
26:14:686:G:H1	53:L5:16:HIS:CD2	2.35	0.45
10:1A:38:ILE:O	10:1A:70:ARG:HG3	2.17	0.45
1:1G:408:A:H2'	1:1G:409:G:O4'	2.16	0.45
1:1G:818:G:O2'	1:1G:819:A:H5'	2.15	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1359:A:C2	26:1H:1372:U:C4	3.05	0.45
26:1H:2082:A:H2'	26:1H:2083:G:O4'	2.16	0.45
26:1H:2105:C:H2'	26:1H:2106:G:H8	1.81	0.45
26:1H:2576:G:H3'	57:1H:3671:HOH:O	2.15	0.45
26:1H:493:G:H2'	26:1H:494:G:O4'	2.17	0.45
22:1K:72:C:H2'	22:1K:73:C:H6	1.81	0.45
29:21:201:THR:HG22	29:21:202:LYS:N	2.31	0.45
35:25:9:GLU:HG3	35:25:18:LYS:HG3	1.99	0.45
35:25:88:ASN:HD21	35:25:92:GLU:HB2	1.81	0.45
29:29:96:PHE:O	29:29:175:VAL:HG11	2.16	0.45
11:2I:18:ARG:HG2	11:2I:33:THR:OG1	2.16	0.45
23:2K:8:U:OP1	23:2K:8:U:H4'	2.16	0.45
23:2L:16:U:H2'	23:2L:16:U:O2	2.16	0.45
30:31:114:VAL:O	30:31:118:ALA:N	2.35	0.45
4:32:13:ARG:HD3	4:32:36:ARG:O	2.16	0.45
30:39:114:VAL:HG21	30:39:202:PHE:CZ	2.51	0.45
12:3I:76:ASN:N	12:3I:76:ASN:OD1	2.32	0.45
31:41:62:LEU:O	31:41:143:GLU:HG2	2.16	0.45
31:41:49:ASP:OD2	31:41:52:ILE:HG12	2.17	0.45
31:49:64:THR:HA	31:49:102:PHE:CD1	2.51	0.45
31:49:59:GLU:HB2	31:49:153:ARG:HH21	1.81	0.45
5:4E:68:GLU:O	5:4E:68:GLU:HG3	2.17	0.45
32:59:76:VAL:O	32:59:80:SER:OG	2.21	0.45
26:1H:587:C:N3	36:78:33:ARG:NH1	2.65	0.45
8:7E:87:SER:HB2	8:7E:93:VAL:CB	2.36	0.45
16:7I:28:ARG:HG2	16:7I:29:ASP:OD1	2.16	0.45
43:A5:20:VAL:O	43:A5:23:LEU:HB2	2.16	0.45
39:A8:15:ARG:HD3	39:A8:15:ARG:HA	1.64	0.45
19:AA:66:MET:HA	19:AA:67:VAL:C	2.36	0.45
26:1H:1754:C:P	40:B8:96:ARG:HH12	2.39	0.45
20:BI:74:LYS:HB3	20:BI:75:ASN:H	1.42	0.45
26:1H:536:A:OP1	41:C8:53:ARG:NH1	2.50	0.45
47:I8:41:ARG:NE	47:I8:41:ARG:HA	2.30	0.45
48:J8:85:LEU:HA	48:J8:85:LEU:HD13	1.82	0.45
54:Q8:30:ARG:CZ	54:Q8:30:ARG:HB2	2.46	0.45
28:11:136:ILE:O	28:11:168:ARG:NH2	2.49	0.45
1:13:160:A:OP1	1:13:160:A:H8	1.98	0.45
1:13:32:A:H2'	1:13:33:A:C8	2.51	0.45
1:13:431:A:H2'	1:13:432:A:O4'	2.17	0.45
1:13:542:G:O2'	1:13:543:C:H5'	2.17	0.45
1:13:77:C:HO2'	1:13:92:G:H22	1.57	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1412:A:H2'	26:14:1413:G:H8	1.82	0.45
26:14:154:G:O6	26:14:172:C:N4	2.49	0.45
26:14:2383:G:O2'	26:14:2384:G:H5'	2.16	0.45
26:14:246:C:N4	54:M5:8:LYS:HG3	2.31	0.45
26:14:2647:U:O2	26:14:2673:G:N2	2.39	0.45
26:14:71:A:H2	44:B5:31:HIS:NE2	2.12	0.45
27:16:15:A:H1'	27:16:109:G:C4	2.51	0.45
28:19:92:ILE:HD12	28:19:104:TYR:CE1	2.50	0.45
10:1A:47:PHE:N	10:1A:63:PHE:O	2.29	0.45
10:1A:8:LEU:HD22	10:1A:20:ALA:HB2	1.97	0.45
1:1G:1126:U:H4'	1:1G:1127:G:C5	2.51	0.45
1:1G:1140:C:H2'	1:1G:1141:C:H6	1.82	0.45
1:1G:821:G:H2'	1:1G:822:C:H6	1.82	0.45
26:1H:1433:U:O2	26:1H:1561:G:C2	2.70	0.45
26:1H:1654:A:H1'	26:1H:2823:A:H5'	1.97	0.45
26:1H:2325:G:H2'	26:1H:2326:C:H6	1.81	0.45
26:1H:250:G:H2'	26:1H:251:A:C8	2.51	0.45
26:1H:2867:G:OP2	40:B8:119:LYS:NZ	2.27	0.45
26:1H:288:C:H2'	26:1H:289:A:H8	1.82	0.45
26:1H:280:C:C2	26:1H:361:G:N2	2.84	0.45
26:1H:571:A:H5'	26:1H:2030:A:N7	2.31	0.45
1:13:973:G:OP1	10:1I:57:LYS:HD3	2.17	0.45
3:22:95:THR:HB	3:22:97:LYS:HG2	1.97	0.45
29:29:31:CYS:O	29:29:91:VAL:HG22	2.16	0.45
11:2A:85:ARG:HA	11:2A:112:THR:OG1	2.17	0.45
3:2E:133:ALA:HA	3:2E:136:GLN:HB3	1.99	0.45
1:1G:543:C:OP1	4:32:14:ARG:HD2	2.16	0.45
36:35:85:LEU:HA	36:35:88:LEU:HB2	1.98	0.45
30:39:83:PHE:O	30:39:84:VAL:HB	2.15	0.45
12:3A:49:ASN:N	12:3A:49:ASN:OD1	2.49	0.45
31:41:9:ARG:O	31:41:13:GLU:HG2	2.16	0.45
5:42:18:ARG:HH21	5:42:25:ARG:HG2	1.81	0.45
32:51:136:ILE:HD12	32:51:137:ASP:H	1.81	0.45
6:5E:97:PHE:HB3	18:9I:32:ARG:HD3	1.99	0.45
33:61:133:HIS:HB2	33:61:134:PRO:HD2	1.97	0.45
36:78:133:SER:O	36:78:137:LYS:HG3	2.16	0.45
36:78:96:THR:C	36:78:98:GLU:H	2.20	0.45
8:7E:121:ASP:OD1	8:7E:121:ASP:N	2.49	0.45
1:13:468:A:H4'	16:7I:80:PHE:O	2.17	0.45
9:82:78:LYS:HE3	9:82:101:PHE:CD2	2.52	0.45
1:1G:1187:G:P	9:82:113:LYS:HZ1	2.39	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:82:119:ALA:O	9:82:120:ARG:HB2	2.15	0.45
37:88:32:TYR:O	37:88:105:GLU:HA	2.17	0.45
17:8I:63:ARG:HG3	17:8I:64:PRO:N	2.31	0.45
19:AI:40:ILE:HD11	19:AI:62:ILE:CG2	2.44	0.45
19:AI:7:LYS:NZ	19:AI:7:LYS:HB3	2.31	0.45
52:J5:16:ARG:NH1	52:J5:17:ASP:OD2	2.50	0.45
1:13:1157:A:N6	1:13:1178:G:H21	2.13	0.45
26:14:2124:G:H2'	26:14:2124:G:N3	2.32	0.45
26:14:1903:G:OP2	28:19:241:PRO:HB2	2.16	0.45
10:1A:89:ASP:OD1	10:1A:89:ASP:N	2.48	0.45
1:1G:673:G:O3'	6:52:87:ARG:NH2	2.50	0.45
26:1H:1332:G:N2	26:1H:1610:A:C8	2.85	0.45
26:1H:200:U:O2	26:1H:386:G:N2	2.49	0.45
26:1H:2250:G:OP1	37:88:85:LYS:NZ	2.45	0.45
26:1H:2275:C:C6	26:1H:2275:C:H5'	2.52	0.45
26:1H:2741:A:H2'	26:1H:2742:C:O4'	2.17	0.45
26:1H:589:C:H2'	26:1H:590:A:H8	1.81	0.45
26:1H:880:G:HO2'	26:1H:881:G:C5'	2.27	0.45
35:25:64:ARG:HG2	35:25:79:PHE:CG	2.52	0.45
26:14:2724:C:OP1	29:29:111:ARG:NH1	2.50	0.45
3:2E:79:ARG:O	3:2E:82:GLU:HB2	2.16	0.45
23:2K:9:G:N2	23:2K:21:A:H8	2.14	0.45
30:31:107:LYS:HD2	30:31:206:ILE:HA	1.98	0.45
30:39:183:VAL:O	30:39:187:VAL:HG23	2.16	0.45
12:3I:47:LYS:HE2	12:3I:47:LYS:HB3	1.68	0.45
12:3I:9:GLN:O	12:3I:13:LYS:HG2	2.17	0.45
31:41:107:LEU:HD21	31:41:178:PHE:CE1	2.51	0.45
5:42:70:PRO:HB3	5:42:144:THR:HG22	1.99	0.45
6:52:25:ILE:O	6:52:29:ALA:N	2.35	0.45
33:61:104:GLN:O	33:61:105:HIS:HB2	2.16	0.45
39:65:110:LEU:HD23	39:65:112:PHE:CZ	2.52	0.45
33:69:93:THR:O	33:69:97:ILE:HG13	2.16	0.45
36:78:19:VAL:CB	36:78:20:GLY:HA2	2.46	0.45
1:1G:1250:A:H4'	9:82:68:GLY:N	2.30	0.45
26:14:996:A:H4'	41:85:92:ARG:NE	2.31	0.45
9:8E:121:ARG:NH1	9:8E:122:ALA:O	2.50	0.45
20:BI:56:MET:HG3	20:BI:88:VAL:HG21	1.99	0.45
41:C8:110:VAL:O	41:C8:114:LYS:N	2.45	0.45
26:1H:180:G:OP2	53:P8:32:LYS:HE2	2.17	0.45
1:13:1512:U:H3	1:13:1523:G:H1	1.64	0.45
26:14:1204:A:N1	26:14:1241:A:C2	2.85	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1225:C:O3'	42:95:85:LYS:HA	2.16	0.45
26:14:1514:U:H2'	26:14:1515:C:C6	2.52	0.45
26:14:307:G:H22	26:14:310:A:P	2.39	0.45
26:14:363(B):G:H2'	26:14:363(C):G:H8	1.81	0.45
26:14:775:G:C4	26:14:794:G:C8	3.05	0.45
28:19:69:ARG:NE	28:19:105:ILE:HD11	2.31	0.45
28:19:176:ARG:HG3	28:19:182:LEU:HD22	1.99	0.45
1:1G:1125:U:H5	10:1A:5:ARG:NH1	2.14	0.45
1:1G:1263:C:H2'	1:1G:1264:C:O4'	2.17	0.45
1:1G:1274:G:H2'	1:1G:1275:A:H8	1.79	0.45
1:1G:1364:U:O2'	1:1G:1365:G:H5'	2.17	0.45
1:1G:424:G:H2'	1:1G:425:G:H8	1.82	0.45
1:1G:522:C:H2'	1:1G:523:A:O4'	2.16	0.45
1:1G:560:U:H5'	1:1G:566:G:N2	2.32	0.45
1:1G:619:U:N3	4:32:134:ASP:OD1	2.50	0.45
1:1G:666:G:H5'	1:1G:726:C:H1'	1.99	0.45
1:1G:675:A:H2'	1:1G:676:A:O4'	2.17	0.45
1:1G:980:C:H2'	14:5A:21:TYR:HE2	1.81	0.45
26:1H:1121:C:H5''	26:1H:1122:G:OP2	2.16	0.45
26:1H:182:A:H2'	26:1H:183:C:C6	2.52	0.45
26:1H:2112:G:H22	26:1H:2169:A:N6	2.09	0.45
26:1H:2245:U:H5''	26:1H:2246:G:H5'	1.99	0.45
26:1H:2312:U:H5'	31:41:88:ILE:HG13	1.99	0.45
26:1H:354:G:H2'	26:1H:355:G:H8	1.82	0.45
26:1H:6:A:H2'	26:1H:7:G:O4'	2.16	0.45
27:1J:43:C:O2	31:49:95:ARG:NH2	2.39	0.45
29:21:52:LEU:O	29:21:75:VAL:HG23	2.17	0.45
3:2E:7:PRO:O	3:2E:11:ARG:HG3	2.17	0.45
3:2E:40:ARG:O	3:2E:44:GLU:HG2	2.17	0.45
26:1H:38:A:H1'	30:31:48:THR:O	2.17	0.45
30:31:64:ILE:HG23	30:31:65:TRP:CD1	2.52	0.45
4:32:25:ARG:NH2	4:32:30:LYS:HD2	2.32	0.45
36:35:143:GLY:HA2	36:35:144:GLU:OE2	2.16	0.45
5:42:60:TYR:O	5:42:64:ARG:HG3	2.17	0.45
26:1H:2562:U:H4'	35:68:25:LEU:HD23	1.98	0.45
35:68:64:ARG:O	35:68:82:ASN:HA	2.17	0.45
37:88:25:ASP:HA	37:88:100:GLY:O	2.16	0.45
17:8A:28:PRO:HA	17:8A:35:VAL:HA	1.98	0.45
17:8I:89:LEU:HA	17:8I:89:LEU:HD13	1.69	0.45
39:A8:10:ARG:HG3	39:A8:13:ARG:HH12	1.81	0.45
19:AA:64:GLU:N	19:AA:64:GLU:OE1	2.31	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
27:1J:104:A:OP1	46:D5:72:ARG:NH1	2.50	0.45
49:G5:47:ASN:HD22	49:G5:47:ASN:H	1.63	0.45
45:G8:100:ALA:HB1	45:G8:101:LYS:HG3	1.98	0.45
51:I5:22:ILE:HD13	51:I5:22:ILE:H	1.81	0.45
26:1H:2331:G:O2'	47:I8:43:THR:HB	2.16	0.45
31:41:66:GLN:HA	51:M8:6:HIS:CE1	2.51	0.45
54:Q8:37:SER:HB2	54:Q8:40:GLU:OE1	2.17	0.45
28:11:175:LEU:HD23	28:11:175:LEU:HA	1.67	0.45
2:12:63:MET:HG3	2:12:225:ALA:HB1	1.99	0.45
1:13:1298:C:H4'	1:13:1299:A:O4'	2.16	0.45
1:13:631:G:H2'	1:13:632:A:C2	2.52	0.45
1:13:713:G:H2'	1:13:714:G:C8	2.51	0.45
26:14:198:C:H5'	26:14:2244:U:OP1	2.16	0.45
26:14:2695:C:H2'	26:14:2696:U:C6	2.52	0.45
26:14:881:G:O6	26:14:882:G:N2	2.50	0.45
26:14:950:G:H2'	26:14:951:C:O4'	2.17	0.45
27:16:29:A:H2'	27:16:30:C:C6	2.52	0.45
27:16:61:G:C6	27:16:62:C:C4	3.05	0.45
1:1G:1023:G:H5''	1:1G:1024:G:N2	2.32	0.45
1:1G:1052:U:H5''	1:1G:1053:G:OP2	2.17	0.45
1:1G:1175:G:C2	1:1G:1176:A:C5	3.04	0.45
1:1G:35:G:H2'	1:1G:36:C:C6	2.51	0.45
1:1G:410:G:H5''	1:1G:411:A:OP1	2.17	0.45
1:1G:662:G:O2'	1:1G:836:G:OP1	2.31	0.45
26:1H:1000:A:OP2	26:1H:1154:G:N1	2.34	0.45
26:1H:1319:G:C6	26:1H:1320:C:N4	2.85	0.45
26:1H:1636:C:H2'	26:1H:1637:A:H8	1.82	0.45
26:1H:2392:A:H2	26:1H:2424:C:N4	2.13	0.45
26:1H:2515:C:O2	26:1H:2570:G:C2	2.70	0.45
26:1H:602:G:O2'	26:1H:655:A:N6	2.48	0.45
26:1H:729:G:N7	28:11:208:LYS:HG3	2.31	0.45
9:8E:114:TYR:CD2	10:1I:59:SER:HA	2.52	0.45
22:1K:39:PSU:H3'	22:1K:40:G:H8	1.81	0.45
3:22:82:GLU:O	3:22:85:ARG:N	2.45	0.45
35:25:3:GLN:HB2	35:25:4:PRO:HD2	1.99	0.45
26:14:1993:U:H4'	29:29:128:SER:OG	2.17	0.45
23:2K:79:U:H2'	23:2K:80:C:C6	2.52	0.45
23:2L:2:C:H2'	23:2L:3:C:C6	2.49	0.45
26:14:826:U:H4'	36:35:55:ARG:HA	1.99	0.45
12:3I:59:ARG:HA	12:3I:65:GLU:HA	1.98	0.45
31:49:63:ILE:HG22	31:49:143:GLU:HB2	1.97	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:4E:10:MET:HB3	5:4E:10:MET:HE2	1.92	0.45
36:78:135:LEU:HD22	36:78:139:LYS:HD3	1.99	0.45
9:82:112:LYS:HE3	9:82:118:LYS:N	2.25	0.45
38:98:25:ALA:HB3	38:98:70:LEU:HD21	1.98	0.45
26:14:2432:A:C2	48:F5:35:THR:HG22	2.52	0.45
46:H8:7:ALA:HB2	46:H8:59:LEU:HD22	1.98	0.45
48:J8:19:GLN:O	48:J8:35:THR:N	2.45	0.45
53:P8:5:TRP:HA	53:P8:5:TRP:CE3	2.51	0.45
28:11:182:LEU:O	28:11:271:ILE:HG12	2.16	0.45
2:12:75:LYS:HA	2:12:78:GLN:CB	2.43	0.45
1:13:115:G:H4'	1:13:116:A:O5'	2.17	0.45
1:13:559:A:H4'	1:13:560:U:H5''	1.99	0.45
1:13:701:C:O2	1:13:703:G:N1	2.49	0.45
1:13:911:U:H2'	1:13:912:C:C6	2.52	0.45
26:14:1225:C:H4'	42:95:85:LYS:HD3	1.98	0.45
26:14:1271:G:O3'	26:14:1272:A:H4'	2.17	0.45
26:14:1421:G:C2	26:14:1422:G:C8	3.04	0.45
26:14:17:G:H2'	26:14:18:C:C6	2.52	0.45
26:14:2116:G:H1	26:14:2162:G:P	2.40	0.45
26:14:2244:U:H2'	26:14:2245:U:O4'	2.17	0.45
26:14:2749:A:O2'	32:59:62:LYS:HD3	2.17	0.45
1:1G:1039:C:H3'	1:1G:1040:U:H5''	1.98	0.45
1:1G:149:A:H2'	1:1G:150:C:C6	2.51	0.45
1:1G:411:A:N7	1:1G:413:G:N3	2.64	0.45
1:1G:719:C:H5	1:1G:720:C:C4	2.35	0.45
26:1H:1086:A:H1'	26:1H:1103:A:H61	1.81	0.45
26:1H:1169:G:H2'	26:1H:1170:G:O4'	2.16	0.45
26:1H:1467:C:N4	26:1H:1525:G:H1	2.15	0.45
26:1H:1683:C:H2'	26:1H:1684:C:H6	1.81	0.45
26:1H:1906:G:H5''	26:1H:1929:G:O2'	2.17	0.45
26:1H:2329:G:H2'	26:1H:2330:G:C8	2.52	0.45
26:1H:2628:C:H1'	26:1H:2781:A:H2'	1.99	0.45
26:1H:306:U:C5	26:1H:307:G:C5	3.05	0.45
10:1I:42:THR:HG23	10:1I:67:THR:O	2.17	0.45
27:1J:17:C:H2'	27:1J:18:G:O4'	2.17	0.45
4:32:64:LEU:HB2	4:32:198:VAL:HG11	1.99	0.45
26:14:2406:U:C2	36:35:72:PRO:HB2	2.52	0.45
30:39:32:LEU:O	30:39:36:VAL:HG22	2.17	0.45
37:45:26:TYR:O	37:45:26:TYR:CD1	2.70	0.45
37:45:66:ILE:O	37:45:67:ARG:HB2	2.17	0.45
13:4I:91:ARG:HA	13:4I:91:ARG:HD2	1.69	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:62:85:TYR:N	7:62:85:TYR:CD1	2.85	0.45
35:68:17:ARG:HH11	35:68:17:ARG:HA	1.81	0.45
1:13:667:G:H4'	15:6I:51:HIS:CE1	2.51	0.45
40:75:99:LEU:HD23	40:75:99:LEU:HA	1.51	0.45
8:7E:83:ILE:HB	8:7E:137:VAL:HG13	1.99	0.45
41:85:99:ALA:HB2	41:85:106:PHE:CD1	2.52	0.45
41:85:55:ARG:H	41:85:55:ARG:HG2	1.51	0.45
19:AA:19:VAL:HB	19:AA:20:LEU:HD22	1.98	0.45
40:B8:34:VAL:HG22	40:B8:41:ARG:HB3	1.99	0.45
46:D5:82:ARG:HA	46:D5:83:PRO:HD3	1.69	0.45
47:E5:32:ARG:HB3	47:E5:33:ALA:H	1.55	0.45
44:F8:1:MET:HG2	44:F8:2:LYS:H	1.81	0.45
46:H8:58:VAL:O	46:H8:60:GLU:N	2.50	0.45
48:J8:21:ARG:HB3	48:J8:21:ARG:HE	1.34	0.45
54:M5:32:LEU:HD12	54:M5:32:LEU:O	2.17	0.45
1:13:1409:C:H2'	1:13:1410:G:H8	1.82	0.45
1:13:1442:G:C5	1:13:1446:A:C6	3.05	0.45
1:13:51:A:C6	1:13:353:A:C2	3.06	0.45
1:13:79:G:H2'	1:13:79:G:N3	2.32	0.45
26:14:1063:G:H1	26:14:1075:C:H1'	1.81	0.45
26:14:135:G:H2'	26:14:136:G:O4'	2.17	0.45
26:14:2286:A:H4'	26:14:2287:A:O4'	2.17	0.45
26:14:2417:C:H2'	26:14:2418:A:H8	1.81	0.45
26:14:2636:U:OP1	29:29:80:GLU:HB2	2.17	0.45
26:14:2772:C:H2'	26:14:2773:C:C6	2.52	0.45
26:14:671:C:H2'	26:14:672:C:H6	1.81	0.45
26:14:780:G:H21	26:14:783:A:N6	2.06	0.45
34:15:96:GLU:O	34:15:100:GLU:HG3	2.17	0.45
1:1G:1002:G:H2'	1:1G:1003:G:C8	2.51	0.45
1:1G:1129:C:C4	1:1G:1139:G:N1	2.85	0.45
1:1G:1306:A:N6	1:1G:1331:G:H1'	2.32	0.45
1:1G:960:U:H4'	1:1G:961:U:C5'	2.47	0.45
26:1H:1857:G:O2'	26:1H:1885:A:N6	2.45	0.45
26:1H:2118:U:H5''	26:1H:2119:A:OP1	2.16	0.45
26:1H:2210:G:H2'	26:1H:2211:G:N1	2.32	0.45
26:1H:2309:A:H2'	26:1H:2310:A:O4'	2.17	0.45
26:1H:2401:U:H2'	26:1H:2402:C:H1'	1.99	0.45
26:1H:2740:A:N6	26:1H:2764:A:C8	2.85	0.45
26:1H:699:A:H2'	26:1H:700:G:O4'	2.17	0.45
22:1K:15:A:N7	22:1K:16:U:H5	2.15	0.45
26:1H:2572:A:C8	29:21:144:ARG:HD3	2.52	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:21:178:GLU:HG3	29:21:178:GLU:H	1.52	0.45
3:2E:134:ILE:HD13	3:2E:134:ILE:HA	1.63	0.45
3:2E:135:LYS:NZ	5:4E:52:PRO:HG2	2.31	0.45
26:14:671:C:OP1	36:35:42:SER:O	2.35	0.45
36:35:65:ARG:HH22	54:M5:47:LYS:HE2	1.82	0.45
31:41:26:GLN:HG3	31:41:30:GLU:OE1	2.16	0.45
5:42:68:GLU:O	5:42:70:PRO:HD3	2.16	0.45
37:45:21:THR:HG21	37:45:101:ARG:HD2	1.99	0.45
6:52:72:VAL:HG13	6:52:73:ASN:OD1	2.16	0.45
38:55:100:LEU:HD12	38:55:111:LEU:HB2	1.98	0.45
38:55:87:TYR:CE1	38:55:117:VAL:HG12	2.52	0.45
6:5E:5:GLU:HG3	6:5E:93:SER:OG	2.17	0.45
9:8E:112:LYS:HD2	9:8E:113:LYS:H	1.81	0.45
9:8E:84:ALA:O	9:8E:87:GLN:HB2	2.17	0.45
19:AA:65:ASN:OD1	19:AA:65:ASN:N	2.50	0.45
20:BI:44:ALA:HB2	20:BI:88:VAL:HG13	1.98	0.45
45:C5:40:GLU:N	45:C5:40:GLU:OE2	2.50	0.45
45:C5:6:HIS:CE1	45:C5:7:VAL:HG13	2.52	0.45
46:D5:52:SER:C	46:D5:54:HIS:H	2.20	0.45
26:1H:456:C:H3'	44:F8:68:ARG:HH22	1.82	0.45
44:B5:5:TYR:HD1	49:G5:33:MET:SD	2.40	0.45
51:I5:24:THR:HG1	51:I5:25:TYR:H	1.65	0.45
51:M8:40:HIS:CE1	51:M8:45:GLY:HA3	2.53	0.45
52:N8:11:THR:HG22	52:N8:12:SER:O	2.17	0.45
54:Q8:45:GLY:CA	54:Q8:46:ARG:C	2.86	0.45
28:11:69:ARG:NH2	28:11:128:GLY:O	2.35	0.44
1:13:652:U:HO2'	1:13:653:A:P	2.40	0.44
26:14:1485:G:N2	26:14:1504:C:N3	2.43	0.44
26:14:2507:C:H2'	26:14:2508:G:O4'	2.16	0.44
26:14:2567:G:H2'	26:14:2568:C:C6	2.52	0.44
26:14:2698:U:H2'	26:14:2699:C:H6	1.81	0.44
26:14:2784:C:H1'	29:29:37:ARG:NH2	2.31	0.44
26:14:601:C:O2	26:14:605:C:H4'	2.16	0.44
26:14:603:A:H8	26:14:604:G:H1'	1.81	0.44
28:19:28:GLU:HG3	28:19:29:PRO:HD2	1.99	0.44
26:1H:1046:A:HO2'	26:1H:1047:G:P	2.38	0.44
26:1H:10:G:H21	26:1H:2801:A:HO2'	1.62	0.44
26:1H:1543:A:C2	26:1H:1545:A:C4	3.05	0.44
26:1H:1557:C:H5''	26:1H:1558:A:OP2	2.17	0.44
26:1H:184:C:H1'	26:1H:217:G:H1'	1.98	0.44
26:1H:1889:A:H2'	26:1H:1890:A:O4'	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2157:G:HO2'	26:1H:2158:A:P	2.40	0.44
26:1H:242:G:H5'	54:Q8:60:LEU:HD13	1.98	0.44
26:1H:244:A:C2	26:1H:255:A:C4	3.05	0.44
26:1H:2636:U:H2'	26:1H:2637:U:H6	1.78	0.44
26:1H:2698:U:H2'	26:1H:2699:C:C6	2.52	0.44
26:1H:2820:A:O2'	26:1H:2821:A:OP1	2.31	0.44
26:1H:557:U:H2'	26:1H:558:G:C8	2.52	0.44
10:1I:76:ASN:HA	10:1I:77:PRO:HD2	1.80	0.44
35:25:111:PHE:N	35:25:111:PHE:CD1	2.85	0.44
5:42:135:THR:O	5:42:138:ALA:HB3	2.17	0.44
32:51:97:ARG:HB2	32:51:97:ARG:HE	1.43	0.44
32:59:94:TYR:CD1	32:59:94:TYR:N	2.83	0.44
14:5A:12:ARG:HB2	14:5A:14:PRO:HD3	1.99	0.44
1:13:1202:G:O2'	14:5I:27:CYS:HB3	2.17	0.44
14:5I:3:ARG:O	14:5I:7:ILE:HG23	2.18	0.44
39:65:18:ILE:O	39:65:21:THR:HG22	2.16	0.44
26:1H:2393:A:H5'	36:78:63:PRO:HB3	1.98	0.44
8:7E:6:ILE:O	8:7E:9:MET:N	2.50	0.44
1:13:1147:C:O2'	9:8E:16:ARG:HD3	2.18	0.44
40:B8:60:THR:HG22	40:B8:77:PRO:HA	1.98	0.44
20:BA:55:ILE:O	20:BA:58:LYS:N	2.49	0.44
47:E5:27:GLU:HB2	47:E5:69:PHE:HD1	1.82	0.44
45:G8:83:THR:HG22	45:G8:84:ARG:HE	1.83	0.44
49:K8:64:LEU:HD21	49:K8:68:ARG:NH1	2.33	0.44
51:M8:10:VAL:HG22	51:M8:11:PRO:HD2	1.99	0.44
54:Q8:34:TRP:HE3	54:Q8:34:TRP:H	1.64	0.44
54:Q8:36:LYS:HG2	54:Q8:39:LYS:HZ2	1.82	0.44
1:13:1128:C:H42	1:13:1143:G:H1	1.63	0.44
1:13:374:A:C6	1:13:375:U:C4	3.05	0.44
1:13:51:A:N7	1:13:114:U:O2'	2.48	0.44
1:13:976:G:C8	1:13:1358:U:C2	3.05	0.44
26:14:1113:U:H2'	26:14:1114:G:O4'	2.18	0.44
26:14:1665:A:N6	57:14:3584:HOH:O	2.29	0.44
26:14:2236:C:H2'	26:14:2237:G:O4'	2.17	0.44
26:14:2287:A:H2	26:14:2346:A:C2	2.36	0.44
26:14:2410:G:H2'	26:14:2411:A:O4'	2.17	0.44
26:14:2820:A:C5	38:55:4:LEU:HD11	2.52	0.44
26:14:2822:G:H2'	26:14:2823:A:H5''	2.00	0.44
26:14:464:U:H4'	53:L5:5:TRP:CZ3	2.53	0.44
26:14:699:A:H2'	26:14:700:G:O4'	2.16	0.44
1:1G:1145:C:H4'	1:1G:1146:A:O5'	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1116:C:H42	1:1G:1184:G:H1	1.64	0.44
1:1G:1399:C:H4'	1:1G:1400:C:H5''	1.99	0.44
1:1G:600:C:H2'	1:1G:601:C:C6	2.52	0.44
26:1H:106:C:H2'	26:1H:107:C:H6	1.81	0.44
27:1J:10:C:C4	27:1J:11:C:C5	3.05	0.44
22:1L:29:U:H3	22:1L:43:A:H2	1.65	0.44
3:22:113:ALA:HB3	3:22:114:PRO:HD3	1.98	0.44
3:22:141:VAL:HA	3:22:144:SER:HB2	1.99	0.44
1:1G:1190:G:OP1	3:22:5:ILE:HG23	2.18	0.44
30:31:24:LEU:HA	30:31:24:LEU:HD12	1.76	0.44
4:32:168:ARG:HD3	4:32:169:LYS:N	2.32	0.44
36:35:105:LEU:O	36:35:106:LEU:HB3	2.17	0.44
30:39:83:PHE:C	30:39:85:GLY:H	2.20	0.44
32:51:153:LYS:HD2	32:51:153:LYS:H	1.81	0.44
6:52:54:LYS:NZ	6:52:54:LYS:HB2	2.32	0.44
32:59:90:LYS:O	32:59:160:LYS:HA	2.18	0.44
32:59:4:ILE:HB	32:59:7:LEU:HD11	2.00	0.44
8:72:5:PRO:HB2	8:72:32:LYS:HE2	2.00	0.44
17:8I:67:LYS:HA	17:8I:70:ARG:NH1	2.32	0.44
38:55:103:ARG:HD3	43:A5:40:ASN:ND2	2.31	0.44
44:B5:21:PHE:CD1	44:B5:21:PHE:N	2.84	0.44
46:D5:29:TYR:HB3	46:D5:34:ASN:OD1	2.17	0.44
46:H8:5:LEU:HD13	46:H8:6:LYS:HB2	1.99	0.44
50:L8:7:LYS:HG3	50:L8:34:GLU:HG2	1.97	0.44
1:13:1374:A:C4	1:13:1375:A:C8	3.06	0.44
1:13:1511:G:H2'	1:13:1512:U:O4'	2.18	0.44
1:13:46:G:H2'	1:13:366:C:C5	2.52	0.44
1:13:540:G:H2'	1:13:541:G:O4'	2.17	0.44
1:13:614:A:H2'	1:13:615:C:H6	1.81	0.44
26:14:565:C:H4'	26:14:1253:A:C6	2.52	0.44
26:14:2367:G:O5'	26:14:2367:G:H8	2.00	0.44
26:14:2387:U:H1'	47:E5:41:ARG:HE	1.82	0.44
26:14:839:U:H2'	26:14:840:C:H6	1.82	0.44
28:19:8:PRO:HB3	28:19:14:ARG:HB2	1.99	0.44
1:1G:458:C:H42	1:1G:474:G:H1	1.65	0.44
1:1G:838:G:O5'	1:1G:838:G:H8	2.00	0.44
1:1G:877:C:O2	8:72:3:THR:OG1	2.30	0.44
26:1H:1372:U:H2'	26:1H:1373:A:C8	2.52	0.44
26:1H:1435:G:O5'	26:1H:1435:G:H8	2.00	0.44
26:1H:354:G:H2'	26:1H:355:G:C8	2.52	0.44
26:1H:463:G:H5''	26:1H:464:U:OP2	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:768:G:O2'	26:1H:1379:A:N6	2.50	0.44
26:1H:855:G:C6	26:1H:856:C:C4	3.06	0.44
26:1H:931:G:O2'	50:L8:24:LYS:HE2	2.18	0.44
29:21:176:ILE:HB	29:21:181:LEU:HB2	1.99	0.44
3:2E:77:ILE:HA	3:2E:84:ILE:CD1	2.47	0.44
30:31:63:LYS:HG2	30:31:65:TRP:O	2.17	0.44
30:39:53:THR:HG23	30:39:55:GLY:N	2.16	0.44
4:3E:191:ARG:HD3	4:3E:200:GLU:OE1	2.17	0.44
24:3K:67:C:H2'	24:3K:68:A:H8	1.82	0.44
31:49:64:THR:HB	31:49:94:LEU:HD11	1.98	0.44
32:51:157:TYR:HD1	32:51:157:TYR:HA	1.71	0.44
38:55:81:ASP:O	38:55:82:GLU:HB3	2.17	0.44
32:59:144:VAL:O	32:59:148:ILE:HG12	2.18	0.44
33:61:81:VAL:HG11	33:61:88:ILE:HD13	1.99	0.44
42:95:85:LYS:HD2	42:95:87:HIS:CA	2.46	0.44
35:68:104:ARG:NH1	40:B8:36:GLU:OE2	2.50	0.44
20:BI:55:ILE:HA	20:BI:55:ILE:HD13	1.90	0.44
45:C5:88:LYS:O	45:C5:89:PHE:HB3	2.16	0.44
42:D8:14:VAL:HB	42:D8:96:ILE:HG13	2.00	0.44
49:K8:14:ARG:HB3	49:K8:15:LYS:NZ	2.33	0.44
50:L8:35:ARG:HB3	50:L8:37:LEU:CD2	2.40	0.44
52:N8:37:LYS:HG3	52:N8:38:ALA:N	2.32	0.44
1:13:1118:C:P	9:8E:104:ARG:HH11	2.41	0.44
1:13:1128:C:H4'	9:8E:16:ARG:HH12	1.83	0.44
1:13:1252:A:H2'	1:13:1253:G:O4'	2.17	0.44
1:13:1302:U:OP2	13:4I:21:TYR:OH	2.16	0.44
1:13:1346:A:C8	1:13:1348:U:C2	3.05	0.44
26:14:2418:A:H2'	26:14:2419:U:H6	1.82	0.44
10:1A:26:ALA:C	10:1A:85:LEU:HD21	2.38	0.44
1:1G:1003:G:H2'	1:1G:1004:A:H4'	1.98	0.44
1:1G:1307:U:H2'	1:1G:1308:U:O4'	2.17	0.44
1:1G:178:C:H2'	1:1G:179:A:O4'	2.17	0.44
1:1G:186(D):C:H2'	1:1G:186(E):C:C6	2.52	0.44
26:1H:1086:A:H5''	26:1H:1087:G:OP1	2.18	0.44
26:1H:1470:G:N2	26:1H:1522:G:OP2	2.50	0.44
26:1H:1778:U:H2'	26:1H:1784:A:C6	2.52	0.44
26:1H:2287:A:H62	26:1H:2344:U:H3	1.59	0.44
26:1H:2583:G:H2'	26:1H:2584:U:O2	2.18	0.44
26:1H:2684:U:OP2	40:B8:53:ARG:NH1	2.51	0.44
26:1H:270(C):C:H42	26:1H:270(W):G:H1	1.65	0.44
3:22:125:GLU:HG2	3:22:190:ARG:O	2.16	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:22:167:TRP:H	3:22:167:TRP:HE3	1.66	0.44
1:1G:1112:C:C2	3:22:178:LEU:HB2	2.53	0.44
11:2I:86:GLY:N	11:2I:112:THR:OG1	2.32	0.44
12:3A:102:ARG:HB3	12:3A:102:ARG:HE	1.56	0.44
24:3L:9:G:N2	24:3L:11:U:O4	2.50	0.44
31:41:26:GLN:CD	31:41:27:ASN:HB2	2.37	0.44
5:42:107:ARG:HH22	5:42:108:ALA:HB2	1.83	0.44
32:51:85:LYS:O	32:51:133:VAL:HG23	2.18	0.44
34:58:12:ARG:HB3	34:58:50:ASP:OD1	2.18	0.44
7:62:132:GLY:HA3	7:62:135:VAL:HB	1.98	0.44
26:1H:2675:A:H5''	35:68:31:LYS:HD3	2.00	0.44
15:6I:4:THR:OG1	15:6I:7:GLU:HB2	2.18	0.44
36:78:84:ASN:HA	36:78:115:LEU:O	2.18	0.44
17:8I:100:LYS:CB	17:8I:101:ARG:HE	2.31	0.44
43:A5:1:MET:CE	43:A5:2:GLU:H	2.30	0.44
19:AA:53:ASN:OD1	19:AA:55:LYS:HB3	2.17	0.44
40:B8:125:ARG:O	40:B8:128:GLU:HB3	2.17	0.44
20:BA:50:GLU:HA	20:BA:100:ILE:HG12	2.00	0.44
46:H8:104:PHE:CE2	46:H8:119:GLU:HG2	2.52	0.44
1:13:1450:U:O2'	1:13:1451:A:H3'	2.18	0.44
1:13:1489:G:H2'	1:13:1490:C:O4'	2.18	0.44
1:13:191(C):G:H2'	1:13:191(D):U:O4'	2.17	0.44
1:13:418:C:H2'	1:13:419:C:H6	1.82	0.44
1:13:573:A:N3	1:13:883:C:O2'	2.44	0.44
1:13:580:U:OP1	15:6I:54:ARG:NH2	2.48	0.44
26:14:1093:G:H1	26:14:1097:U:H5''	1.82	0.44
26:14:1257:C:H2'	26:14:1258:C:C6	2.52	0.44
26:14:1342:A:H2	26:14:1602:U:N3	2.16	0.44
26:14:1542:G:O5'	26:14:1543:A:H5''	2.18	0.44
26:14:1771:C:C1'	26:14:1786:A:C8	3.01	0.44
26:14:1962:C:O2'	26:14:1964:G:OP2	2.35	0.44
26:14:2492:U:H2'	26:14:2493:U:C6	2.53	0.44
26:14:2525:G:N2	26:14:2539:C:C2	2.85	0.44
26:14:2681:C:C4	26:14:2724:C:C5	3.06	0.44
26:14:601:C:O2'	26:14:605:C:H5''	2.16	0.44
26:14:957:A:N1	26:14:2459:A:C8	2.85	0.44
28:19:146:GLU:OE2	28:19:152:GLY:N	2.43	0.44
9:82:114:TYR:HD2	10:1A:60:ARG:HG3	1.82	0.44
1:1G:1036:G:H5'	1:1G:1037:C:OP2	2.17	0.44
1:1G:569:C:H5''	1:1G:570:G:OP1	2.17	0.44
1:1G:833:U:O2'	1:1G:834:C:H5'	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1858:G:HO2'	26:1H:1859:A:P	2.40	0.44
26:1H:1899:G:O2'	26:1H:1900:A:OP2	2.31	0.44
26:1H:2428:G:H21	36:78:61:ARG:HH21	1.66	0.44
26:1H:2896:C:H2'	26:1H:2897:U:H6	1.83	0.44
29:21:119:ARG:HG3	29:21:119:ARG:NH1	2.33	0.44
29:21:179:GLU:O	29:21:180:ASN:HB2	2.18	0.44
29:21:67:PHE:C	29:21:69:LYS:H	2.20	0.44
29:29:126:PRO:O	29:29:135:HIS:HD2	2.00	0.44
3:2E:88:ARG:HA	3:2E:91:LEU:HD12	1.98	0.44
4:32:24:GLU:HG2	4:32:25:ARG:N	2.32	0.44
36:35:52:GLU:O	36:35:54:GLY:N	2.50	0.44
4:3E:107:ARG:HH11	4:3E:114:ARG:NH2	2.14	0.44
31:41:11:TYR:O	31:41:15:VAL:HB	2.18	0.44
5:42:127:ASN:HA	5:42:128:PRO:HD3	1.86	0.44
31:49:63:ILE:HD12	31:49:141:PHE:CD2	2.52	0.44
5:4E:39:GLY:HA2	5:4E:113:ALA:HB1	1.99	0.44
13:4I:11:ARG:HG3	13:4I:46:LYS:HD2	1.99	0.44
34:58:32:THR:O	34:58:36:GLY:N	2.47	0.44
33:61:29:TYR:O	33:61:32:PRO:HD2	2.17	0.44
33:61:71:ILE:HG12	33:61:72:LEU:HD13	1.99	0.44
7:6E:115:ARG:O	7:6E:118:VAL:HG13	2.17	0.44
8:7E:64:LYS:HG2	8:7E:79:VAL:HG21	2.00	0.44
16:7I:74:LEU:HD22	16:7I:79:VAL:HG21	1.99	0.44
9:8E:77:ILE:O	9:8E:81:ILE:HG12	2.17	0.44
1:13:254:G:O2'	17:8I:16:GLN:O	2.34	0.44
38:98:109:ALA:HA	38:98:110:PRO:HD2	1.89	0.44
26:1H:2376:A:C2	39:A8:112:PHE:HB3	2.52	0.44
48:F5:19:GLN:HB3	48:F5:35:THR:O	2.16	0.44
45:G8:35:TYR:CE2	45:G8:69:ALA:HB3	2.52	0.44
45:G8:76:CYS:HA	45:G8:97:ARG:HG2	1.99	0.44
46:H8:132:ASN:HD22	46:H8:134:PRO:HD3	1.82	0.44
46:H8:155:LEU:HD23	46:H8:155:LEU:HA	1.87	0.44
46:H8:50:GLN:H	46:H8:50:GLN:HG2	1.53	0.44
51:I5:23:GLU:C	51:I5:24:THR:HG1	2.20	0.44
54:M5:40:GLU:H	54:M5:43:GLN:CG	2.14	0.44
54:M5:40:GLU:N	54:M5:43:GLN:HG3	2.11	0.44
31:41:104:GLU:OE1	51:M8:23:GLU:HG3	2.18	0.44
2:12:58:ILE:H	2:12:58:ILE:HG12	1.51	0.44
1:13:390:C:H2'	1:13:391:G:C8	2.52	0.44
1:13:633:G:H5''	1:13:634:C:OP2	2.17	0.44
1:13:74:C:H2'	1:13:75:C:O4'	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:801:U:H2'	1:13:802:A:H8	1.81	0.44
26:14:1024:G:H8	26:14:1024:G:O5'	2.00	0.44
26:14:1340:U:C2	26:14:1603:A:O4'	2.71	0.44
26:14:214:G:OP1	26:14:214:G:H4'	2.17	0.44
26:14:2313:C:H2'	26:14:2314:C:C6	2.52	0.44
26:14:825:C:H4'	26:14:2428:G:C5	2.53	0.44
26:14:1255:U:OP2	26:14:2502:G:N2	2.50	0.44
26:14:945:A:H2	57:14:3690:HOH:O	2.00	0.44
28:19:80:ALA:HB3	28:19:94:LEU:HB3	1.99	0.44
2:1E:54:THR:O	2:1E:58:ILE:HG13	2.18	0.44
1:1G:1128:C:N3	1:1G:1139:G:N1	2.66	0.44
1:1G:1176:A:C2'	1:1G:1177:G:H5'	2.47	0.44
1:1G:302:G:O2'	1:1G:556:C:H5''	2.17	0.44
26:1H:139:G:N3	26:1H:141:A:N1	2.65	0.44
26:1H:2887:U:H2'	26:1H:2888:C:C6	2.52	0.44
26:1H:34:C:C6	26:1H:34:C:OP2	2.70	0.44
26:1H:740:U:H2'	26:1H:741:G:C8	2.53	0.44
1:13:963:G:C2	10:1I:55:LYS:NZ	2.84	0.44
29:21:100:GLU:O	29:21:172:VAL:HG23	2.17	0.44
29:21:20:ALA:O	29:21:21:VAL:HB	2.18	0.44
3:2E:180:ALA:HB1	3:2E:182:ILE:HG13	1.99	0.44
11:2I:33:THR:HA	11:2I:39:PRO:HA	1.99	0.44
4:32:189:PRO:HB2	4:32:194:LEU:HD21	1.99	0.44
4:32:192:GLU:H	4:32:192:GLU:HG3	1.50	0.44
36:35:35:HIS:HB3	36:35:36:LYS:H	1.53	0.44
12:3A:41:ARG:HH12	12:3A:43:VAL:HG22	1.82	0.44
24:3K:14:A:H2'	24:3K:14:A:N3	2.33	0.44
37:45:60:ARG:NH1	37:45:61:GLY:HA2	2.33	0.44
31:49:19:LEU:HD23	31:49:19:LEU:HA	1.80	0.44
13:4A:14:ARG:HA	13:4A:43:THR:O	2.17	0.44
14:5A:41:ARG:HG3	14:5A:42:ILE:HG12	1.99	0.44
33:69:143:SER:OG	33:69:145:VAL:N	2.51	0.44
40:75:26:ASP:O	40:75:49:VAL:HG13	2.17	0.44
36:78:134:ALA:O	36:78:138:LEU:HB2	2.17	0.44
36:78:94:GLU:OE1	36:78:124:LYS:HD3	2.18	0.44
37:88:1:MET:HB3	37:88:2:LEU:H	1.58	0.44
1:13:265:G:O2'	17:8I:67:LYS:N	2.51	0.44
38:98:24:GLN:OE1	38:98:36:THR:HG21	2.17	0.44
43:A5:21:VAL:HG21	43:A5:76:VAL:HG12	2.00	0.44
43:A5:61:ASN:N	43:A5:61:ASN:OD1	2.50	0.44
39:A8:3:ARG:HG2	39:A8:4:LEU:N	2.33	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:AA:49:ILE:HD11	19:AA:62:ILE:HD11	2.00	0.44
42:D8:25:LEU:H	42:D8:92:THR:CG2	2.30	0.44
43:E8:29:LEU:HD22	43:E8:69:LEU:CD1	2.48	0.44
45:G8:68:HIS:HB3	45:G8:71:LYS:HG2	2.00	0.44
46:H8:121:HIS:HB2	46:H8:171:ILE:HD12	2.00	0.44
28:11:44:ASN:O	28:11:46:GLN:O	2.35	0.44
1:13:1171:G:H8	1:13:1171:G:O5'	2.00	0.44
1:13:435:C:H2'	1:13:436:C:H6	1.83	0.44
1:13:842:C:H1'	1:13:848:C:H42	1.83	0.44
26:14:1065:U:N3	26:14:1074:G:H1'	2.32	0.44
26:14:1131:G:OP2	26:14:2515:C:H4'	2.17	0.44
26:14:1141:U:OP2	34:15:63:THR:OG1	2.23	0.44
26:14:1190:G:OP1	36:35:32:THR:HA	2.17	0.44
26:14:1259:G:H2'	26:14:1260:G:H8	1.82	0.44
26:14:1341:U:OP2	26:14:1394:U:O2'	2.22	0.44
26:14:1388:G:H2'	26:14:1389:G:C8	2.51	0.44
26:14:1423:G:C4	26:14:1424:G:C8	3.06	0.44
26:14:1480:G:H1	26:14:1513:C:H42	1.66	0.44
26:14:186:G:H2'	26:14:187:G:H8	1.83	0.44
28:19:71:ASP:OD1	28:19:103:ARG:NH1	2.49	0.44
2:1E:172:ILE:HG13	2:1E:172:ILE:H	1.47	0.44
1:1G:1111:A:H8	1:1G:1111:A:O5'	2.00	0.44
1:1G:1125:U:H2'	1:1G:1126:U:C6	2.53	0.44
1:1G:1127:G:H1'	1:1G:1148:U:O2	2.18	0.44
1:1G:1157:A:C8	1:1G:1158:C:C5	3.06	0.44
1:1G:1171:G:H2'	1:1G:1172:C:C6	2.53	0.44
1:1G:1207:G:C6	1:1G:1208:C:C4	3.06	0.44
1:1G:1315:U:H2'	1:1G:1316:G:O4'	2.17	0.44
1:1G:1500:A:H5''	1:1G:1508:G:H5''	2.00	0.44
1:1G:730:G:C5	1:1G:731:G:H1'	2.53	0.44
26:1H:1024:G:H8	26:1H:1024:G:O5'	2.01	0.44
26:1H:1252:G:H5''	57:1H:3639:HOH:O	2.17	0.44
26:1H:1417:C:H2'	26:1H:1418:G:O4'	2.17	0.44
26:1H:1535:U:O2	26:1H:1535:U:H2'	2.16	0.44
26:1H:1859:A:C2	26:1H:1884:A:H1'	2.53	0.44
26:1H:2098:U:O2	26:1H:2191:G:N2	2.36	0.44
26:1H:2576:G:P	57:1H:3669:HOH:O	2.75	0.44
26:1H:270(Y):G:C2	26:1H:270(Z):U:O4	2.71	0.44
26:1H:608:A:H1'	26:1H:621:A:N6	2.33	0.44
3:22:53:ALA:HB2	3:22:115:LEU:HD23	1.99	0.44
4:32:162:LEU:HD21	4:32:178:VAL:HG11	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:35:96:THR:HG23	36:35:99:LEU:HB3	2.00	0.44
4:3E:84:LYS:HA	4:3E:85:LYS:C	2.38	0.44
24:3K:59:U:O2'	24:3K:60:A:OP1	2.31	0.44
31:41:142:PRO:HB2	51:M8:31:ILE:HD12	2.00	0.44
23:2K:67:C:O2'	31:41:78:SER:HB2	2.17	0.44
5:42:34:VAL:O	5:42:41:VAL:HG12	2.18	0.44
31:49:61:ALA:HA	31:49:64:THR:HG23	2.00	0.44
13:4I:50:GLU:O	13:4I:54:VAL:HG23	2.18	0.44
34:58:4:TYR:CE2	41:C8:100:VAL:HG11	2.53	0.44
34:58:71:ILE:HG21	34:58:84:LYS:HG2	1.98	0.44
6:5E:8:ILE:HD12	6:5E:26:ILE:HD13	2.00	0.44
14:5I:7:ILE:HG22	14:5I:28:GLY:HA3	2.00	0.44
33:61:144:VAL:HG22	33:61:145:VAL:H	1.82	0.44
33:69:114:LEU:HD22	33:69:114:LEU:O	2.17	0.44
8:72:19:VAL:HG23	8:72:21:LYS:HB2	1.99	0.44
37:88:135:ASP:HB2	37:88:138:ASP:OD1	2.17	0.44
9:8E:78:LYS:HE2	9:8E:101:PHE:CE1	2.52	0.44
17:8I:10:VAL:HG13	17:8I:19:VAL:HB	1.99	0.44
1:13:128:G:H4'	17:8I:3:LYS:HG2	2.00	0.44
20:BA:63:ILE:HG21	20:BA:81:LYS:HG3	2.00	0.44
48:F5:76:ARG:HB2	48:F5:94:LEU:HD13	1.99	0.44
46:H8:117:LEU:HD22	46:H8:118:GLN:N	2.30	0.44
1:13:1177:G:OP1	1:13:1177:G:H4'	2.18	0.44
1:13:1308:U:OP1	13:4I:98:VAL:HG23	2.18	0.44
1:13:260:G:H2'	1:13:261:U:C6	2.53	0.44
1:13:329:A:C5	1:13:332:G:C6	3.06	0.44
1:13:589:C:H42	1:13:650:G:H1	1.65	0.44
1:13:8:A:N6	4:3E:205:GLU:O	2.51	0.44
26:14:1680:U:H2'	26:14:1681:G:O4'	2.18	0.44
26:14:2865:U:C4	26:14:2866:U:C4	3.05	0.44
26:14:301:G:C6	26:14:317:G:C6	3.05	0.44
26:14:774:A:HO2'	26:14:775:G:P	2.40	0.44
26:14:795:C:H2'	26:14:796:C:H6	1.80	0.44
26:14:868:U:C2	26:14:869:G:C8	3.06	0.44
28:19:33:LEU:HA	28:19:64:ILE:HG23	1.99	0.44
1:1G:1139:G:H22	1:1G:1143:G:N2	2.16	0.44
1:1G:1256:A:H5''	1:1G:1258:G:C4	2.52	0.44
1:1G:555:C:H2'	1:1G:556:C:C6	2.53	0.44
26:1H:1106:G:H2'	26:1H:1107:G:O4'	2.18	0.44
26:1H:2182:G:N2	26:1H:2183:C:O2	2.51	0.44
26:1H:2197:U:O2'	26:1H:2198:A:H2'	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2804:C:H2'	26:1H:2805:G:C8	2.53	0.44
26:1H:43:G:H2'	26:1H:44:A:O4'	2.18	0.44
26:1H:795:C:H2'	26:1H:796:C:H6	1.82	0.44
26:1H:886:C:H3'	26:1H:887:A:C5'	2.48	0.44
26:1H:90:U:H1'	26:1H:91:A:C8	2.52	0.44
3:22:181:ASN:OD1	3:22:204:LEU:HB2	2.18	0.44
3:2E:173:VAL:HG12	3:2E:175:LEU:HG	2.00	0.44
3:2E:4:LYS:HE3	3:2E:4:LYS:HB3	1.66	0.44
30:31:32:LEU:HA	30:31:32:LEU:HD22	1.53	0.44
36:35:135:LEU:HD23	36:35:135:LEU:HA	1.82	0.44
30:39:154:VAL:HA	30:39:191:ARG:O	2.18	0.44
30:39:170:LEU:HD22	30:39:172:TRP:NE1	2.33	0.44
30:39:67:GLN:HG3	30:39:67:GLN:O	2.18	0.44
24:3K:42:U:H2'	24:3K:43:A:H8	1.82	0.44
31:41:111:LEU:HD13	31:41:179:PRO:HG2	2.00	0.44
1:13:7:G:H2'	5:4E:119:LEU:HD22	1.98	0.44
13:4I:66:LEU:O	13:4I:70:LEU:HB2	2.17	0.44
34:58:66:LYS:O	34:58:70:LYS:HB3	2.17	0.44
1:1G:974:A:P	14:5A:41:ARG:HH12	2.41	0.44
33:61:88:ILE:HG22	33:61:90:GLY:N	2.33	0.44
33:69:4:ILE:CG2	33:69:47:LEU:HD13	2.46	0.44
8:72:92:ARG:HA	8:72:92:ARG:HD2	1.60	0.44
9:82:10:ARG:NH1	9:82:105:ASP:OD2	2.51	0.44
7:62:16:LEU:HD11	9:82:45:ALA:HB2	1.99	0.44
9:8E:16:ARG:HB2	9:8E:64:THR:OG1	2.16	0.44
41:85:50:ARG:HH22	42:95:72:VAL:HG23	1.83	0.44
42:95:85:LYS:HD2	42:95:87:HIS:N	2.32	0.44
38:98:12:ARG:HG2	38:98:12:ARG:NH1	2.31	0.44
45:C5:82:PRO:HB2	45:C5:83:THR:O	2.17	0.44
42:D8:62:LEU:HD12	42:D8:62:LEU:HA	1.80	0.44
48:F5:85:LEU:HB2	48:F5:88:LYS:HG3	2.00	0.44
45:G8:44:ILE:HD12	45:G8:62:GLU:CD	2.38	0.44
50:H5:5:LYS:HB3	50:H5:5:LYS:HE3	1.58	0.44
46:H8:30:ASN:ND2	46:H8:33:LEU:HB3	2.32	0.44
47:I8:46:LYS:HG2	47:I8:77:ARG:O	2.17	0.44
53:P8:12:ARG:NH2	53:P8:44:PRO:HB3	2.32	0.44
1:13:1036:G:H2'	1:13:1037:C:C2	2.53	0.44
1:13:109:A:C6	1:13:326:G:C6	3.05	0.44
1:13:1365:G:H2'	1:13:1366:C:H6	1.83	0.44
1:13:1387:G:H2'	1:13:1388:C:C6	2.53	0.44
1:13:256:U:H2'	1:13:257:G:O4'	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:66:G:C2	1:13:67:C:C6	3.05	0.44
1:13:711:G:H2'	1:13:712:A:H8	1.83	0.44
26:14:1035:U:H2'	26:14:1036:G:C8	2.53	0.44
26:14:243:U:OP1	54:M5:6:THR:OG1	2.30	0.44
26:14:629:G:H5''	26:14:650:C:O2'	2.18	0.44
28:19:77:ALA:HB2	28:19:97:TYR:CD2	2.53	0.44
2:1E:124:SER:HB2	2:1E:125:PRO:HD2	1.99	0.44
1:1G:1107:C:C4	1:1G:1108:G:C8	3.06	0.44
1:1G:1157:A:C8	1:1G:1158:C:C4	3.06	0.44
1:1G:1245:A:H2'	1:1G:1246:C:O4'	2.18	0.44
1:1G:1384:C:H2'	1:1G:1385:G:C8	2.53	0.44
1:1G:1508:G:H2'	1:1G:1509:C:O4'	2.17	0.44
1:1G:197:A:OP2	1:1G:197:A:H3'	2.17	0.44
1:1G:286:G:H2'	1:1G:287:U:C6	2.53	0.44
1:1G:371:G:O2'	1:1G:373:A:N7	2.44	0.44
1:1G:613:C:H2'	1:1G:614:A:O4'	2.18	0.44
1:1G:750:G:C2	15:6A:23:GLY:HA3	2.53	0.44
1:1G:763:G:H2'	1:1G:764:C:H6	1.83	0.44
1:1G:828:A:H2'	1:1G:829:G:O4'	2.18	0.44
1:1G:921:U:H2'	1:1G:922:G:O4'	2.17	0.44
26:1H:1050:A:C8	26:1H:2751:G:C5	3.06	0.44
26:1H:1094:U:HO2'	26:1H:1096:A:P	2.41	0.44
26:1H:2070:G:C2	26:1H:2442:C:C2	3.06	0.44
26:1H:787:U:H5''	26:1H:788:A:H5'	2.00	0.44
26:1H:978:G:C2'	26:1H:979:G:H5'	2.48	0.44
22:1K:15:A:H2'	22:1K:15:A:N3	2.33	0.44
22:1L:47:C:H42	22:1L:56:G:H22	1.66	0.44
29:29:145:LYS:HB3	29:29:145:LYS:HE3	1.86	0.44
30:31:160:ASN:OD1	30:31:163:VAL:HG23	2.18	0.44
12:3I:77:LEU:HD22	12:3I:81:SER:HB2	2.00	0.44
13:4A:84:ILE:HG12	19:AA:63:THR:HG21	1.99	0.44
1:13:947:G:OP1	13:4I:108:ARG:HB3	2.18	0.44
13:4I:98:VAL:HG22	13:4I:110:ARG:HH21	1.82	0.44
34:58:16:ILE:HG21	34:58:26:LEU:HD11	1.99	0.44
39:65:7:TYR:HE1	39:65:11:LYS:NZ	2.16	0.44
26:1H:833:U:H1'	36:78:55:ARG:NH1	2.33	0.44
38:98:28:LEU:O	38:98:28:LEU:HD22	2.18	0.44
38:98:29:LEU:HA	38:98:29:LEU:HD12	1.74	0.44
18:9I:32:ARG:HD2	18:9I:65:ILE:HG21	2.00	0.44
39:A8:41:ASP:HB3	39:A8:48:LEU:HD11	2.00	0.44
19:AI:49:ILE:HG22	19:AI:51:VAL:HG22	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:B8:18:ASP:OD1	40:B8:18:ASP:N	2.50	0.44
43:E8:17:VAL:HG23	43:E8:43:GLY:HA3	1.99	0.44
37:88:134:ARG:HH21	46:H8:122:ARG:CZ	2.31	0.44
51:I5:18:CYS:HB3	51:I5:19:GLY:HA2	2.00	0.44
51:M8:18:CYS:HB3	51:M8:39:CYS:CB	2.48	0.44
52:N8:20:ARG:HA	52:N8:23:HIS:ND1	2.32	0.44
1:13:1178:G:C8	1:13:1178:G:H3'	2.52	0.43
1:13:1372:U:H5''	9:8E:71:SER:HB2	2.00	0.43
1:13:1428:A:H2'	1:13:1429:C:O4'	2.17	0.43
1:13:1452:C:HO2'	1:13:1453:G:P	2.35	0.43
1:13:1486:G:H2'	1:13:1487:G:C1'	2.48	0.43
1:13:196:A:OP1	20:BI:68:LYS:NZ	2.51	0.43
1:13:547:A:H4'	1:13:548:G:O5'	2.18	0.43
1:13:639:G:H2'	1:13:640:A:C8	2.52	0.43
1:13:79:G:N2	1:13:89:U:C2	2.86	0.43
1:13:859:A:H2'	1:13:860:A:C8	2.53	0.43
1:13:908:A:H2'	1:13:909:A:C8	2.53	0.43
1:13:958:A:C6	1:13:959:A:C6	3.05	0.43
26:14:1278:A:H2'	26:14:1279:G:C8	2.53	0.43
26:14:1440:G:C4	26:14:1441:G:C8	3.06	0.43
26:14:1729:A:OP1	26:14:1729:A:H8	2.01	0.43
26:14:654(I):C:H5''	26:14:654(M):C:N4	2.33	0.43
26:14:906:G:O3'	37:45:67:ARG:NH2	2.51	0.43
10:1A:79:ARG:HB3	10:1A:79:ARG:HH11	1.83	0.43
1:1G:1055:A:H5''	1:1G:1056:U:OP2	2.18	0.43
1:1G:1277:C:O2'	1:1G:1279:A:H8	2.01	0.43
1:1G:942:G:C2	1:1G:1342:C:C2	3.06	0.43
1:1G:1357:A:N7	1:1G:1358:U:C5	2.86	0.43
1:1G:1481:U:H2'	1:1G:1482:G:H8	1.82	0.43
1:1G:458:C:N4	1:1G:464:G:O6	2.51	0.43
1:1G:616:G:N3	1:1G:617:G:C8	2.86	0.43
1:1G:924:C:O2'	1:1G:1502:A:N6	2.50	0.43
26:1H:1105:U:H2'	26:1H:1106:G:C8	2.53	0.43
26:1H:2528:U:H2'	26:1H:2530:A:O5'	2.18	0.43
26:1H:466:A:O4'	26:1H:683:C:H4'	2.18	0.43
26:1H:563:G:OP2	57:1H:3607:HOH:O	2.20	0.43
26:1H:588:U:O4	26:1H:670:A:H1'	2.18	0.43
29:21:31:CYS:HA	29:21:32:PRO:HD3	1.68	0.43
3:22:37:GLN:O	3:22:40:ARG:N	2.51	0.43
3:2E:108:ASN:HB3	3:2E:111:LEU:HB2	2.00	0.43
3:2E:19:GLU:HA	3:2E:54:ARG:HH12	1.83	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:2I:124:LYS:HG3	11:2I:125:PHE:CE1	2.53	0.43
23:2K:72:C:H2'	23:2K:73:C:H6	1.83	0.43
23:2L:87:A:C2'	23:2L:87:A:N3	2.77	0.43
30:31:139:PHE:HB2	30:31:166:ALA:HB1	2.00	0.43
30:31:93:LYS:HA	30:31:93:LYS:HD2	1.50	0.43
4:32:101:LEU:HD12	4:32:101:LEU:HA	1.84	0.43
36:35:144:GLU:HA	36:35:145:PRO:HD3	1.62	0.43
36:35:1:MET:HB2	36:35:5:ASP:OD2	2.18	0.43
1:13:437:U:H5''	4:3E:155:LEU:HD21	2.00	0.43
4:3E:162:LEU:HA	4:3E:162:LEU:HD23	1.74	0.43
4:3E:78:LEU:HB3	4:3E:93:PHE:HE1	1.83	0.43
12:3I:90:VAL:HG11	12:3I:93:LEU:CG	2.46	0.43
24:3K:16:U:H2'	24:3K:18:G:OP2	2.17	0.43
24:3K:21:A:C5	24:3K:58:U:H1'	2.52	0.43
5:42:90:VAL:O	5:42:91:LEU:HD13	2.17	0.43
1:1G:1329:A:C5'	13:4A:29:ARG:HE	2.27	0.43
25:4K:37:G:H8	25:4K:37:G:OP2	2.01	0.43
32:59:18:GLU:HB2	32:59:25:LYS:HD2	2.00	0.43
14:5I:4:LYS:O	14:5I:7:ILE:HG12	2.18	0.43
7:62:13:GLN:HA	7:62:14:PRO:HD3	1.82	0.43
39:65:109:GLY:O	39:65:110:LEU:HD22	2.18	0.43
15:6A:10:LYS:HD2	15:6A:10:LYS:HA	1.67	0.43
36:78:116:GLY:N	36:78:134:ALA:HB2	2.33	0.43
8:7E:56:LYS:HA	8:7E:57:PRO:HD3	1.83	0.43
37:88:51:ARG:HD2	37:88:66:ILE:HD11	1.99	0.43
41:C8:17:ILE:HD12	41:C8:32:PHE:HE1	1.83	0.43
47:E5:34:GLY:HA2	47:E5:61:ALA:O	2.18	0.43
26:1H:137(A):G:H1'	44:F8:41:ASN:ND2	2.32	0.43
49:G5:47:ASN:HD22	49:G5:47:ASN:N	2.13	0.43
46:H8:30:ASN:HD21	46:H8:33:LEU:HB3	1.83	0.43
54:Q8:49:VAL:O	54:Q8:49:VAL:HG22	2.18	0.43
28:11:201:HIS:O	28:11:204:ILE:HG23	2.17	0.43
2:12:48:MET:HA	2:12:51:LEU:HD12	1.99	0.43
1:13:1057:G:H2'	1:13:1058:G:O4'	2.18	0.43
1:13:1132:C:H2'	1:13:1133:G:C8	2.46	0.43
1:13:1148:U:O3'	9:8E:14:VAL:HG11	2.18	0.43
1:13:1446:A:O2'	1:13:1447:G:H5'	2.19	0.43
1:13:277:C:H5''	17:8I:68:ARG:NH2	2.32	0.43
1:13:134:A:H1'	1:13:325:A:C4	2.53	0.43
1:13:624:C:H4'	16:7I:11:SER:N	2.33	0.43
26:14:1058:U:O5'	26:14:1058:U:H6	2.01	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2520:C:HO2'	26:14:2565:A:HO2'	1.65	0.43
26:14:2637:U:C4	26:14:2638:G:C6	3.06	0.43
26:14:2676:C:H2'	26:14:2677:G:H8	1.83	0.43
34:15:35:ARG:HG2	34:15:35:ARG:H	1.66	0.43
27:16:57:A:H5'	31:41:26:GLN:OE1	2.18	0.43
10:1A:13:HIS:O	10:1A:17:ASP:HB2	2.17	0.43
2:1E:20:GLU:HG3	2:1E:191:ASP:N	2.33	0.43
2:1E:237:ALA:O	2:1E:239:VAL:HG23	2.18	0.43
1:1G:1069:C:H42	1:1G:1106:G:H1	1.67	0.43
1:1G:129(A):G:C2	1:1G:191(A):G:C8	3.06	0.43
1:1G:1355:G:H2'	1:1G:1356:G:O4'	2.18	0.43
1:1G:374:A:C6	1:1G:375:U:C4	3.06	0.43
1:1G:78:G:O6	1:1G:91:C:N4	2.50	0.43
26:1H:1224:G:OP2	42:D8:66:ARG:NH2	2.50	0.43
26:1H:769:G:H4'	26:1H:1379:A:N1	2.33	0.43
26:1H:1534:G:N2	26:1H:1538:G:H22	2.16	0.43
26:1H:1990:C:H2'	26:1H:1991:U:C6	2.53	0.43
26:1H:2023:G:H4'	26:1H:2617:C:O3'	2.18	0.43
26:1H:2102:U:H2'	26:1H:2103:C:C6	2.53	0.43
26:1H:2822:G:H2'	26:1H:2823:A:H5''	2.00	0.43
26:1H:35:G:H2'	26:1H:36:G:O4'	2.18	0.43
26:1H:579:G:H2'	26:1H:580:C:C6	2.53	0.43
26:1H:28:A:O2'	26:1H:583:G:H5'	2.18	0.43
26:1H:607:U:N3	26:1H:621:A:C2	2.78	0.43
26:1H:654:A:H3'	26:1H:654:A:N3	2.32	0.43
27:1J:74:U:H2'	27:1J:75:G:O4'	2.18	0.43
30:31:11:VAL:HG22	30:31:125:LEU:HB2	1.99	0.43
30:39:131:GLY:HA2	30:39:138:GLU:HB3	2.01	0.43
5:42:6:PHE:HD1	5:42:36:ASP:HB3	1.83	0.43
13:4I:82:MET:C	13:4I:84:ILE:H	2.20	0.43
32:51:74:ASN:ND2	32:51:138:LYS:HD3	2.34	0.43
6:52:10:LEU:HA	6:52:84:ASN:O	2.18	0.43
6:5E:21:LEU:O	6:5E:24:GLU:HB3	2.18	0.43
14:5I:21:TYR:HE2	14:5I:23:ARG:NE	2.15	0.43
33:61:5:LEU:HD12	33:61:13:GLY:O	2.18	0.43
33:69:76:THR:HG23	33:69:77:LEU:N	2.33	0.43
15:6A:67:LEU:HD23	15:6A:67:LEU:HA	1.85	0.43
40:75:23:ARG:O	40:75:26:ASP:HB2	2.17	0.43
36:78:39:LYS:HB2	36:78:45:LEU:CD2	2.47	0.43
36:78:58:THR:HG21	54:Q8:52:LYS:NZ	2.33	0.43
16:7A:34:GLU:HG2	16:7A:35:LYS:N	2.33	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:7E:9:MET:SD	8:7E:32:LYS:HG2	2.58	0.43
9:82:99:LEU:HB3	9:82:101:PHE:CD1	2.53	0.43
37:88:14:ARG:HG2	37:88:41:TRP:CH2	2.53	0.43
38:98:96:ARG:CZ	38:98:117:VAL:HG23	2.49	0.43
43:A5:78:GLU:OE2	43:A5:99:ARG:HD3	2.18	0.43
39:A8:89:ARG:HG3	39:A8:92:TYR:O	2.17	0.43
40:B8:106:SER:O	40:B8:110:ILE:HB	2.18	0.43
29:21:13:ARG:HD2	40:B8:58:ASN:HB3	2.00	0.43
20:BA:75:ASN:O	20:BA:79:ARG:HB2	2.18	0.43
20:BI:97:ALA:O	20:BI:99:LEU:HD13	2.18	0.43
46:D5:157:LEU:HA	46:D5:158:PRO:HD2	1.85	0.43
51:I5:11:PRO:HA	51:I5:25:TYR:CG	2.53	0.43
52:J5:56:LYS:HB3	52:J5:56:LYS:HE2	1.74	0.43
48:J8:48:LYS:HG3	48:J8:48:LYS:HZ3	1.75	0.43
51:M8:13:ARG:CA	51:M8:24:THR:HG21	2.49	0.43
31:41:142:PRO:HB2	51:M8:31:ILE:HB	2.00	0.43
54:Q8:16:ILE:HD13	54:Q8:56:GLU:OE2	2.18	0.43
54:Q8:21:LYS:HZ2	54:Q8:21:LYS:HB3	1.83	0.43
1:13:1159:U:O4'	1:13:1182:G:N2	2.51	0.43
1:13:1379:G:N7	7:6E:2:ALA:HB3	2.33	0.43
1:13:625:G:H4'	16:7I:16:HIS:CG	2.53	0.43
26:14:1336:A:H2'	26:14:1337:G:H8	1.83	0.43
26:14:142:G:H2'	26:14:143:C:C6	2.53	0.43
26:14:1444:G:H2'	26:14:1445:C:C5	2.53	0.43
26:14:1849:G:H2'	26:14:1850:G:H8	1.83	0.43
26:14:531:C:C5	26:14:2035:G:C2	3.06	0.43
26:14:250:G:OP2	54:M5:13:ARG:NH2	2.51	0.43
26:14:2689:U:H5''	26:14:2713:A:C2	2.53	0.43
26:14:2887:U:H2'	26:14:2888:C:C6	2.53	0.43
27:16:30:C:OP2	39:A8:32:LEU:HD21	2.18	0.43
28:19:146:GLU:HB2	28:19:189:CYS:HB3	1.99	0.43
1:1G:1004:A:H2	1:1G:1024:G:C8	2.36	0.43
1:1G:1053:G:O2'	1:1G:1054:C:OP2	2.30	0.43
1:1G:1129:C:O5'	1:1G:1130:A:H5'	2.18	0.43
1:1G:1453:G:H22	20:BA:54:LYS:NZ	2.15	0.43
1:1G:17:U:H2'	1:1G:18:C:C6	2.53	0.43
1:1G:693:G:H2'	1:1G:694:A:C8	2.53	0.43
26:1H:1242:A:C8	26:1H:1243:G:C8	3.06	0.43
26:1H:1707:G:C5	26:1H:1756:G:C6	3.06	0.43
26:1H:1845:G:OP1	28:11:258:LYS:NZ	2.43	0.43
26:1H:1878:G:H2'	26:1H:1879:C:C6	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2251:G:H2'	26:1H:2252:G:C8	2.53	0.43
26:1H:2474:C:H3'	26:1H:2475:C:H6	1.84	0.43
22:1K:87:A:C8	26:1H:2507:C:H1'	2.53	0.43
26:1H:1710:C:H4'	26:1H:2858:C:O2	2.17	0.43
26:1H:384:U:H2'	26:1H:385:C:C6	2.50	0.43
26:1H:673:C:H5''	30:31:81:PRO:HD2	2.00	0.43
10:1I:90:LEU:N	10:1I:91:PRO:HD3	2.34	0.43
27:1J:66:A:C2	27:1J:108:C:C4	3.06	0.43
27:1J:87:G:H2'	27:1J:88:C:H5'	2.00	0.43
3:22:47:LEU:HB3	3:22:52:LEU:HD13	1.99	0.43
35:25:12:ASP:HB3	35:25:85:VAL:HG23	1.98	0.43
11:2A:38:ASN:HA	11:2A:39:PRO:HD3	1.90	0.43
11:2I:27:ASN:OD1	11:2I:28:THR:N	2.52	0.43
23:2K:12:G:C6	23:2K:13:G:C4	3.07	0.43
23:2K:19:G:O6	23:2K:69:A:H5''	2.17	0.43
23:2L:31:C:H2'	23:2L:32:C:H6	1.84	0.43
36:35:36:LYS:HB3	36:35:37:GLY:H	1.39	0.43
4:3E:155:LEU:HD13	4:3E:158:ILE:HD11	2.00	0.43
4:3E:156:GLU:O	4:3E:160:GLN:HB2	2.18	0.43
4:3E:107:ARG:NH2	4:3E:194:LEU:HD22	2.26	0.43
5:42:122:GLU:O	5:42:126:ARG:NH1	2.52	0.43
31:49:4:ASP:CG	31:49:5:VAL:N	2.70	0.43
34:58:67:LEU:HD23	34:58:88:GLU:HB3	2.00	0.43
1:1G:939:G:H5'	7:62:102:ARG:CZ	2.48	0.43
39:65:24:LEU:HD12	39:65:40:ILE:O	2.18	0.43
35:68:101:PRO:HB3	35:68:122:LEU:HD12	2.00	0.43
8:72:73:ASP:HB2	8:72:75:ARG:NH2	2.21	0.43
16:7I:74:LEU:HD23	16:7I:74:LEU:HA	1.62	0.43
43:A5:70:TYR:OH	43:A5:72:LYS:HD2	2.19	0.43
39:A8:106:ARG:HG3	39:A8:106:ARG:H	1.48	0.43
1:13:1318:A:H5''	19:AI:10:PHE:CD2	2.53	0.43
20:BI:53:LEU:HA	20:BI:56:MET:HB3	1.99	0.43
45:C5:84:ARG:HE	45:C5:85:VAL:C	2.20	0.43
46:H8:48:PHE:CE1	46:H8:71:VAL:HG21	2.53	0.43
54:Q8:46:ARG:HH21	54:Q8:48:PHE:HA	1.83	0.43
28:11:147:LEU:HD13	28:11:155:LEU:HD21	1.99	0.43
28:11:263:ARG:HB2	28:11:263:ARG:HH11	1.82	0.43
28:11:70:TRP:CD1	28:11:70:TRP:C	2.91	0.43
2:12:162:ILE:O	2:12:185:ILE:HG12	2.18	0.43
1:13:1099:G:H2'	1:13:1100:C:C6	2.52	0.43
1:13:1126:U:N3	1:13:1127:G:C2	2.85	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:131:C:O2	1:13:131:C:H2'	2.18	0.43
1:13:102:G:O2'	1:13:151:A:N3	2.40	0.43
1:13:244:U:H4'	1:13:245:C:O5'	2.18	0.43
1:13:321:A:C2	1:13:333:G:C2	3.07	0.43
1:13:875:C:O2'	8:7E:14:ARG:NH1	2.48	0.43
1:13:574:A:N3	1:13:883:C:H1'	2.34	0.43
26:14:307:G:N2	26:14:309:G:H3'	2.34	0.43
26:14:273(C):C:N4	26:14:363(C):G:H1	2.12	0.43
26:14:38:A:H1'	30:39:48:THR:HB	2.00	0.43
26:14:729:G:O5'	28:19:208:LYS:NZ	2.48	0.43
26:14:676:A:H2	26:14:802:A:H61	1.61	0.43
2:1E:5:ILE:HB	2:1E:221:LEU:HD23	2.01	0.43
1:1G:101:A:C2'	1:1G:102:G:H5'	2.48	0.43
1:1G:129(A):G:C6	1:1G:188:U:H4'	2.54	0.43
1:1G:1309:G:C6	1:1G:1329:A:C2	3.06	0.43
1:1G:1330:U:H5'	13:4A:24:GLY:H	1.84	0.43
1:1G:332:G:OP2	20:BA:10:LEU:HD12	2.18	0.43
1:1G:396:G:O2'	1:1G:398:C:OP1	2.22	0.43
1:1G:552:U:H1'	12:3A:32:PHE:CE1	2.54	0.43
1:1G:758:G:H8	1:1G:758:G:O5'	2.01	0.43
1:1G:731:G:OP1	1:1G:766:A:H1'	2.19	0.43
26:1H:1263:U:H2'	26:1H:1264:G:C8	2.53	0.43
26:1H:1400:G:H2'	26:1H:1401:G:C8	2.53	0.43
26:1H:2627:G:N2	26:1H:2777:G:OP2	2.51	0.43
26:1H:2797:U:H5''	26:1H:2798:C:OP2	2.18	0.43
26:1H:749:C:H5''	57:1H:4040:HOH:O	2.17	0.43
26:1H:934:G:H2'	26:1H:935:C:H6	1.84	0.43
22:1K:34:U:H2'	22:1K:36:A:OP2	2.18	0.43
22:1K:53:A:N7	22:1K:54:G:H1'	2.33	0.43
29:21:66:HIS:O	29:21:66:HIS:ND1	2.51	0.43
29:21:92:THR:O	29:21:95:ILE:HG22	2.18	0.43
35:25:113:LYS:HG2	35:25:113:LYS:H	1.33	0.43
23:2K:27:G:OP1	23:2K:27:G:H4'	2.19	0.43
23:2K:27:G:N3	23:2K:27:G:H5''	2.33	0.43
23:2K:62:G:HO2'	23:2K:63:G:P	2.39	0.43
36:35:47:ASP:HB3	36:35:49:ARG:N	2.32	0.43
1:1G:363:A:OP1	12:3A:33:ARG:HG3	2.18	0.43
31:49:82:LEU:HA	31:49:86:MET:SD	2.59	0.43
5:4E:8:GLU:HG2	5:4E:34:VAL:HG22	2.00	0.43
32:59:138:LYS:HD2	32:59:138:LYS:HA	1.83	0.43
26:14:2531:A:H5'	32:59:157:TYR:CZ	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:5E:16:GLN:HA	6:5E:19:LEU:HD12	2.00	0.43
39:65:26:LEU:O	39:65:88:ASP:HB2	2.18	0.43
8:72:100:ILE:HG21	8:72:125:ARG:NE	2.33	0.43
40:75:50:ILE:HA	40:75:99:LEU:HB2	1.99	0.43
36:78:27:HIS:O	36:78:31:ALA:HA	2.18	0.43
36:78:2:LYS:HD3	36:78:4:SER:CB	2.47	0.43
1:13:875:C:H4'	8:7E:18:ARG:HH12	1.83	0.43
1:1G:277:C:OP1	17:8A:41:LYS:HE3	2.18	0.43
17:8I:90:ILE:O	17:8I:93:GLN:HB3	2.18	0.43
42:95:5:VAL:HB	42:95:37:VAL:HB	1.99	0.43
41:C8:90:VAL:HG22	42:D8:39:LEU:HB3	1.99	0.43
1:13:1358:U:H2'	1:13:1359:C:O4'	2.19	0.43
1:13:1417:G:C6	1:13:1482:G:C6	3.06	0.43
1:13:34:C:H2'	1:13:35:G:C8	2.53	0.43
1:13:688:G:C5	1:13:700:G:C2	3.07	0.43
1:13:740:U:O2'	1:13:741:G:H5'	2.18	0.43
1:13:791:G:H2'	1:13:792:A:H5'	2.01	0.43
26:14:1105:U:H2'	26:14:1106:G:O4'	2.19	0.43
26:14:1041:C:N3	26:14:1115:G:N2	2.66	0.43
26:14:1287:A:C5	26:14:1288:U:C4	3.07	0.43
26:14:1591:G:H2'	26:14:1592:C:O4'	2.18	0.43
26:14:2300:G:H2'	26:14:2301:C:C6	2.54	0.43
26:14:2376:A:H2'	26:14:2377:A:O4'	2.19	0.43
26:14:2432:A:C8	48:F5:33:LYS:HD2	2.54	0.43
26:14:2480:C:H5'	26:14:2481:G:OP2	2.19	0.43
26:14:2660:A:H2'	26:14:2661:G:O4'	2.19	0.43
26:14:337:C:H2'	26:14:338:G:O4'	2.19	0.43
26:14:396:G:H8	26:14:396:G:O5'	2.01	0.43
26:14:397:G:N7	57:14:3832:HOH:O	2.36	0.43
26:14:570:G:H5''	57:14:3985:HOH:O	2.17	0.43
28:19:62:TYR:HA	28:19:87:ASN:HD21	1.84	0.43
1:1G:1129:C:H4'	1:1G:1130:A:O5'	2.18	0.43
1:1G:1306:A:C6	1:1G:1307:U:C2	3.06	0.43
1:1G:1329:A:H2'	1:1G:1330:U:O4'	2.19	0.43
1:1G:1510:U:H2'	1:1G:1511:G:C8	2.53	0.43
26:1H:1358:G:N2	26:1H:1372:U:C5	2.87	0.43
26:1H:2031:A:N3	26:1H:2455:G:O2'	2.44	0.43
26:1H:2314:C:H2'	26:1H:2315:G:H8	1.83	0.43
26:1H:2340:G:O2'	26:1H:2341:G:H5'	2.18	0.43
26:1H:2481:G:O2'	26:1H:2482:G:H8	2.02	0.43
26:1H:2660:A:H2'	26:1H:2661:G:O4'	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:628:G:H2'	26:1H:629:G:C8	2.53	0.43
26:1H:812:C:H1'	26:1H:1250:G:C2	2.54	0.43
3:2E:164:ARG:HB3	3:2E:164:ARG:HE	1.38	0.43
3:2E:184:TYR:HD2	3:2E:201:TYR:CE1	2.36	0.43
11:2I:19:ALA:HA	11:2I:32:ILE:HA	1.99	0.43
23:2K:46:G:O2'	23:2K:47:C:O5'	2.37	0.43
23:2K:46:G:HO2'	23:2K:47:C:P	2.42	0.43
30:31:136:THR:O	30:31:140:LEU:HB2	2.18	0.43
4:3E:165:MET:SD	4:3E:168:ARG:HD3	2.58	0.43
5:42:147:ASP:O	5:42:151:LEU:HG	2.18	0.43
5:4E:112:LEU:HA	5:4E:112:LEU:HD22	1.72	0.43
5:4E:53:LEU:O	5:4E:57:LYS:HG2	2.19	0.43
38:55:104:ARG:O	38:55:108:GLY:HA2	2.19	0.43
14:5I:8:GLU:HA	14:5I:11:LYS:HD2	1.99	0.43
39:65:103:GLU:O	39:65:106:ARG:NE	2.52	0.43
39:65:17:ARG:HG3	39:65:17:ARG:HH11	1.82	0.43
35:68:77:ILE:HG12	35:68:78:ARG:N	2.34	0.43
33:69:129:THR:HA	33:69:137:PRO:HA	2.01	0.43
7:6E:26:PHE:O	7:6E:30:ILE:HG13	2.18	0.43
40:75:16:ARG:NH1	40:75:80:SER:O	2.45	0.43
36:78:144:GLU:N	36:78:144:GLU:CD	2.71	0.43
1:13:1351:U:O4	9:8E:118:LYS:HE3	2.18	0.43
44:B5:59:VAL:HG21	44:B5:76:ARG:HD3	2.01	0.43
44:B5:9:LEU:HA	49:G5:36:ARG:NH2	2.31	0.43
20:BI:30:LYS:HA	20:BI:33:ILE:HG12	2.01	0.43
49:G5:47:ASN:ND2	49:G5:47:ASN:N	2.66	0.43
45:G8:94:LYS:NZ	45:G8:95:LYS:H	2.15	0.43
51:M8:48:ARG:HA	51:M8:48:ARG:HD3	1.87	0.43
52:N8:45:VAL:HG12	52:N8:51:TYR:CD2	2.54	0.43
2:12:40:HIS:HB2	2:12:190:THR:HG21	2.01	0.43
1:13:1044:A:C5	1:13:1045:C:H1'	2.54	0.43
1:13:1205:U:H2'	1:13:1206:G:C8	2.54	0.43
26:14:2173:A:C2	26:14:2174:C:H1'	2.53	0.43
26:14:2461:C:H2'	26:14:2462:U:C6	2.53	0.43
26:14:2820:A:O5'	38:55:4:LEU:HD23	2.19	0.43
26:14:311:A:C8	26:14:332:A:N7	2.87	0.43
26:14:373:U:H2'	26:14:374:A:H8	1.84	0.43
26:14:783:A:H8	26:14:784:A:H4'	1.84	0.43
26:14:1141:U:H2'	34:15:63:THR:CG2	2.47	0.43
27:16:12:C:C2	47:I8:74:ARG:NH1	2.87	0.43
1:1G:1034:G:H2'	1:1G:1035:A:C8	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1063:C:H3'	1:1G:1064:G:H2'	2.00	0.43
1:1G:1436:U:H2'	1:1G:1437:C:O4'	2.18	0.43
1:1G:476:G:H2'	1:1G:477:G:C8	2.54	0.43
1:1G:4:U:H3'	1:1G:5:U:H5'	2.00	0.43
26:1H:1184:G:C5	26:1H:1185:C:C5	3.06	0.43
26:1H:1412:A:H2'	26:1H:1413:G:C8	2.54	0.43
26:1H:1533:C:C2	26:1H:1534:G:N2	2.86	0.43
26:1H:1827:C:O2'	26:1H:1828:G:H5'	2.18	0.43
26:1H:234:C:H2'	26:1H:235:U:O4'	2.19	0.43
26:1H:234:C:H2'	26:1H:235:U:H6	1.83	0.43
26:1H:2481:G:HO2'	26:1H:2482:G:H8	1.66	0.43
26:1H:2704:C:H2'	26:1H:2705:A:O4'	2.18	0.43
26:1H:271(B):G:H2'	26:1H:271(B):G:H8	1.67	0.43
26:1H:51:G:H1'	26:1H:119:A:N1	2.33	0.43
27:1J:11:C:O5'	27:1J:12:C:H5	2.00	0.43
27:1J:18:G:H1	27:1J:65:C:N4	2.14	0.43
29:21:48:GLN:OE1	29:21:77:ILE:HG21	2.18	0.43
3:22:91:LEU:HD11	3:22:101:LEU:HD12	2.00	0.43
29:29:89:ASP:OD1	29:29:90:THR:N	2.51	0.43
3:2E:62:ASP:OD1	3:2E:97:LYS:HG2	2.17	0.43
11:2I:120:ARG:HA	11:2I:121:PRO:HD3	1.83	0.43
36:35:101:VAL:HB	36:35:106:LEU:HD23	2.01	0.43
12:3A:59:ARG:NH1	12:3A:63:GLY:HA2	2.33	0.43
4:3E:135:LEU:HA	4:3E:136:PRO:HD2	1.85	0.43
12:3I:66:VAL:HG21	12:3I:98:TYR:CE1	2.54	0.43
31:41:128:ARG:HB2	31:41:128:ARG:NH2	2.33	0.43
31:41:36:LYS:HG2	31:41:38:VAL:HG23	2.00	0.43
37:45:25:ASP:OD1	37:45:25:ASP:N	2.52	0.43
31:49:117:PHE:CG	31:49:117:PHE:O	2.72	0.43
32:51:118:PRO:HD2	32:51:121:ILE:HG21	2.00	0.43
4:3E:20:TYR:CZ	6:52:15:ASP:HB3	2.53	0.43
6:52:39:LYS:HB3	6:52:39:LYS:HE3	1.55	0.43
32:59:82:GLY:HA3	32:59:135:GLY:O	2.18	0.43
6:5E:26:ILE:O	6:5E:30:LEU:HD12	2.18	0.43
14:5I:27:CYS:HB2	14:5I:29:ARG:HB2	2.00	0.43
8:72:103:VAL:HG21	8:72:109:ILE:O	2.19	0.43
7:6E:16:LEU:HD11	9:8E:42:ARG:HA	2.01	0.43
17:8I:74:LEU:HA	17:8I:74:LEU:HD22	1.83	0.43
38:98:18:LEU:HD11	38:98:22:ARG:CZ	2.49	0.43
39:A8:32:LEU:N	39:A8:32:LEU:HD23	2.33	0.43
19:AA:17:GLU:HG2	19:AA:21:GLU:OE1	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:B8:3:ARG:O	40:B8:3:ARG:HG3	2.19	0.43
20:BI:39:LYS:O	20:BI:42:GLN:HB3	2.19	0.43
26:14:189:G:OP2	48:F5:39:LYS:HE2	2.19	0.43
45:G8:9:LYS:HA	45:G8:27:VAL:HG22	2.00	0.43
51:M8:38:LYS:N	51:M8:38:LYS:HD2	2.34	0.43
28:11:85:ASP:HB2	28:11:92:ILE:HG12	2.01	0.43
2:12:76:GLN:NE2	2:12:206:ASP:OD1	2.51	0.43
1:13:1007:C:H5'	1:13:1008:C:OP2	2.19	0.43
1:13:127:G:O2'	17:8I:2:PRO:O	2.36	0.43
1:13:1328:C:H2'	1:13:1329:A:C8	2.53	0.43
1:13:234:C:H2'	1:13:235:C:C6	2.52	0.43
1:13:424:G:N2	1:13:425:G:C4	2.86	0.43
1:13:558:G:H5''	1:13:559:A:OP2	2.19	0.43
1:13:684:A:C6	1:13:685:G:C6	3.06	0.43
1:13:77:C:H2'	1:13:78:G:H5''	2.00	0.43
1:13:865:A:C2	1:13:918:A:H4'	2.54	0.43
26:14:1024:G:C3'	26:14:1025:G:H5''	2.42	0.43
26:14:330:A:H2	26:14:1210:A:O2'	2.00	0.43
26:14:1448:G:H1'	26:14:1528:A:N6	2.34	0.43
26:14:1947:C:H2'	26:14:1948:G:O4'	2.19	0.43
26:14:2744:G:C8	26:14:2755:C:C6	3.06	0.43
26:14:71:A:H4'	26:14:72:U:H5''	1.99	0.43
26:14:815:C:H2'	26:14:816:C:C6	2.53	0.43
26:14:1568:G:P	28:19:63:ARG:HH12	2.42	0.43
2:1E:111:ARG:HH11	2:1E:111:ARG:HG2	1.82	0.43
2:1E:84:GLU:O	2:1E:88:ALA:N	2.48	0.43
1:1G:1054:C:H4'	1:1G:1055:A:C5'	2.49	0.43
1:1G:1327:C:H2'	1:1G:1328:C:H6	1.83	0.43
1:1G:1348:U:H4'	9:82:120:ARG:HD2	1.99	0.43
26:1H:1167:U:C2	26:1H:1183:G:N2	2.87	0.43
26:1H:2174:C:H2'	26:1H:2175:C:C6	2.53	0.43
26:1H:2334:G:C2	39:A8:12:PHE:CD1	3.07	0.43
26:1H:2349:G:C6	26:1H:2350:C:C5	3.07	0.43
26:1H:2756:U:H1'	26:1H:2757:A:H5''	2.00	0.43
26:1H:557:U:H2'	26:1H:558:G:H8	1.83	0.43
26:1H:638:G:C5	26:1H:651:G:C2	3.06	0.43
26:1H:784:A:C8	26:1H:792:G:C5	3.06	0.43
26:1H:944:G:H5''	26:1H:945:A:H5'	2.01	0.43
22:1K:69:A:H1'	22:1K:71:U:C5	2.54	0.43
35:25:10:VAL:HG13	35:25:17:ARG:O	2.18	0.43
30:39:129:PHE:HA	30:39:142:TRP:CD1	2.53	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:674:G:C1'	30:39:74:ARG:HD3	2.41	0.43
24:3K:39:U:H2'	24:3K:40:G:H5'	2.01	0.43
5:42:152:ARG:O	8:72:64:LYS:HD3	2.19	0.43
5:42:69:VAL:O	5:42:71:LEU:N	2.51	0.43
38:55:44:LEU:HD22	38:55:48:VAL:HG23	2.00	0.43
32:59:27:LYS:HE3	32:59:28:GLY:N	2.33	0.43
32:59:58:GLU:HB2	32:59:61:HIS:CE1	2.53	0.43
9:8E:34:ASN:OD1	9:8E:34:ASN:N	2.52	0.43
20:BI:29:LYS:HB2	20:BI:29:LYS:HE3	1.65	0.43
20:BI:91:LEU:HA	20:BI:91:LEU:HD23	1.90	0.43
46:D5:124:ILE:HD11	46:D5:165:VAL:HG21	1.99	0.43
44:F8:67:GLY:C	44:F8:69:TYR:H	2.22	0.43
46:H8:40:ASP:OD2	46:H8:43:GLU:HB2	2.19	0.43
2:12:42:ILE:CG1	2:12:202:PRO:HB2	2.48	0.43
2:12:47:THR:HG23	2:12:202:PRO:HD2	2.01	0.43
1:13:1194:U:H2'	1:13:1195:C:C6	2.54	0.43
1:13:152:A:N6	1:13:170:U:C2	2.87	0.43
1:13:288:A:H2'	1:13:289:G:H4'	1.99	0.43
1:13:498:A:C6	1:13:547:A:C8	3.06	0.43
26:14:1392:A:N6	26:14:1393:A:N6	2.67	0.43
26:14:2030:A:H4'	26:14:2031:A:H8	1.84	0.43
26:14:270(T):G:C6	26:14:270(U):C:C4	3.06	0.43
26:14:662:G:H5'	36:35:14:LYS:HB2	2.00	0.43
26:14:74:A:H4'	26:14:75:G:O5'	2.19	0.43
26:14:813:U:C2	26:14:1195:G:N2	2.87	0.43
27:16:87:G:N2	27:16:89(A):A:OP2	2.47	0.43
2:1E:212:GLN:OE1	2:1E:216:SER:OG	2.37	0.43
1:1G:1206:G:C6	1:1G:1207:G:C6	3.07	0.43
1:1G:1348:U:N3	1:1G:1374:A:H2	1.97	0.43
1:1G:56:U:H2'	1:1G:57:G:H8	1.83	0.43
1:1G:833:U:H2'	1:1G:834:C:H6	1.84	0.43
1:1G:976:G:P	14:5A:32:SER:H	2.42	0.43
26:1H:1222:C:C2	26:1H:1223:C:C5	3.07	0.43
26:1H:29:U:H2'	26:1H:30:G:H8	1.80	0.43
26:1H:465:G:H8	26:1H:465:G:O5'	2.01	0.43
26:1H:569:U:C4	26:1H:570:G:C6	3.07	0.43
26:1H:638:G:H2'	26:1H:639:U:O4'	2.19	0.43
1:13:1366:C:O3'	10:1I:60:ARG:NH2	2.52	0.43
27:1J:40:U:H1'	27:1J:45:A:H61	1.83	0.43
3:22:191:THR:OG1	3:22:194:GLY:O	2.33	0.43
29:29:200:GLU:OE1	29:29:200:GLU:N	2.52	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:29:37:ARG:HD3	29:29:44:TYR:CE2	2.53	0.43
3:2E:136:GLN:HG2	3:2E:140:ARG:HH12	1.83	0.43
23:2L:72:C:C6	23:2L:73:C:H5	2.37	0.43
30:31:130:ALA:C	30:31:132:VAL:H	2.22	0.43
4:32:154:ASN:ND2	4:32:155:LEU:HG	2.33	0.43
4:32:93:PHE:CZ	4:32:97:LEU:HD11	2.53	0.43
36:35:78:PRO:HA	36:35:110:TYR:HD2	1.83	0.43
1:13:553:A:H5''	12:3I:24:VAL:HG21	2.01	0.43
1:1G:923:A:H5'	5:42:21:ALA:HB2	1.99	0.43
13:4A:54:VAL:HA	13:4A:57:ARG:HB3	2.01	0.43
1:13:133I:G:OP2	13:4I:23:TYR:HD2	2.02	0.43
26:14:2839:G:H5'	38:55:46:GLY:HA2	2.00	0.43
32:59:84:SER:OG	32:59:85:LYS:N	2.51	0.43
33:61:110:ASP:CG	33:61:130:TYR:HE1	2.22	0.43
7:62:86:GLN:N	7:62:86:GLN:OE1	2.51	0.43
39:65:12:PHE:HA	39:65:12:PHE:HD1	1.66	0.43
39:65:66:ALA:O	39:65:69:VAL:HG13	2.19	0.43
15:6A:77:ARG:HA	15:6A:80:ALA:HB3	2.01	0.43
16:7I:53:VAL:HG21	16:7I:77:ALA:HB1	2.01	0.43
9:82:112:LYS:HD3	9:82:113:LYS:O	2.19	0.43
41:85:74:LEU:HD11	41:85:110:VAL:HG13	2.01	0.43
38:98:4:LEU:HD13	38:98:4:LEU:HA	1.48	0.43
19:AA:80:TYR:CZ	19:AA:82:GLY:HA2	2.54	0.43
40:B8:102:ILE:HA	40:B8:105:LEU:HD22	2.01	0.43
20:BA:48:LYS:O	20:BA:50:GLU:N	2.51	0.43
20:BA:79:ARG:HE	20:BA:83:ARG:NH1	2.17	0.43
43:E8:24:ILE:H	43:E8:24:ILE:HG13	1.71	0.43
44:F8:31:HIS:CD2	44:F8:33:LYS:H	2.37	0.43
50:H5:39:ASP:CG	50:H5:44:ARG:HH21	2.22	0.43
27:16:12:C:H2'	47:I8:73:GLY:HA3	2.01	0.43
44:F8:5:TYR:CZ	49:K8:30:ARG:HB2	2.53	0.43
49:K8:42:GLY:C	49:K8:44:LEU:N	2.71	0.43
51:M8:40:HIS:NE2	51:M8:45:GLY:HA3	2.34	0.43
26:1H:2348:U:OP1	54:Q8:36:LYS:HB3	2.19	0.43
28:11:43:ARG:HD2	28:11:49:ILE:HG12	2.01	0.43
1:13:159:G:N2	1:13:162:A:OP2	2.51	0.43
1:13:160:A:H2'	1:13:161:A:O4'	2.19	0.43
1:13:687:A:C2'	1:13:701:C:H41	2.31	0.43
26:14:1019:U:O2'	26:14:1021:A:H2	2.02	0.43
26:14:1164:G:H2'	26:14:1165:U:C6	2.54	0.43
26:14:1537:C:C2	26:14:1538:G:N7	2.86	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1820:U:O2	28:19:201:HIS:HB3	2.19	0.43
26:14:2191:G:C5	26:14:2192:G:C8	3.05	0.43
26:14:438:G:H2'	26:14:439:G:H8	1.83	0.43
26:14:869:G:C2	26:14:909:A:C2	3.07	0.43
34:15:30:ILE:HG22	34:15:34:LEU:HD22	2.01	0.43
27:16:37:C:H2'	27:16:38:C:O4'	2.18	0.43
10:1A:94:VAL:HG12	10:1A:96:ILE:HD13	1.99	0.43
21:1B:8:THR:HB	21:1B:11:GLY:N	2.33	0.43
1:1G:434:U:H2'	1:1G:435:C:C6	2.54	0.43
1:1G:821:G:H2'	1:1G:822:C:C6	2.53	0.43
26:1H:1265:A:C8	26:1H:1267:U:C2	3.06	0.43
26:1H:1509:C:H2'	26:1H:1511:A:H8	1.84	0.43
26:1H:1530:G:O6	26:1H:1542:G:N2	2.52	0.43
26:1H:1614:A:P	57:1H:3743:HOH:O	2.76	0.43
26:1H:2173:A:OP1	26:1H:2173:A:H8	2.02	0.43
26:1H:265:A:H1'	26:1H:266:G:O4'	2.18	0.43
26:1H:2689:U:H5''	26:1H:2713:A:H2	1.84	0.43
26:1H:2697:G:H2'	26:1H:2698:U:O4'	2.18	0.43
26:1H:2747:G:H5''	32:51:70:THR:CG2	2.49	0.43
26:1H:2774:C:H2'	26:1H:2775:A:O4'	2.18	0.43
26:1H:2838:G:C6	26:1H:2839:G:C5	3.07	0.43
26:1H:2836:U:C4	26:1H:2883:A:N6	2.86	0.43
26:1H:363:G:H2'	26:1H:363(A):A:C8	2.54	0.43
26:1H:761:A:H5''	57:1H:3620:HOH:O	2.18	0.43
27:1J:40:U:OP2	27:1J:40:U:H6	2.01	0.43
27:1J:7:G:H4'	39:65:29:PHE:CD2	2.54	0.43
3:22:14:ILE:HG12	3:22:15:THR:H	1.84	0.43
35:25:113:LYS:O	35:25:117:LEU:HD23	2.19	0.43
29:29:11:MET:HA	29:29:24:THR:HA	2.00	0.43
11:2A:65:ALA:HB1	11:2A:98:LEU:HD21	2.00	0.43
3:2E:195:VAL:O	3:2E:196:LEU:HD23	2.19	0.43
3:2E:69:HIS:HA	3:2E:104:GLN:O	2.19	0.43
23:2K:62:G:O2'	23:2K:63:G:P	2.77	0.43
23:2L:63:G:N3	23:2L:64:G:C8	2.87	0.43
30:31:127:GLU:OE2	30:31:127:GLU:HA	2.19	0.43
12:3A:124:LYS:HA	12:3A:125:PRO:HD3	1.92	0.43
12:3A:24:VAL:O	12:3A:26:ALA:N	2.51	0.43
12:3A:47:LYS:CG	12:3A:48:PRO:HD2	2.47	0.43
31:41:110:ALA:HA	31:41:140:ILE:O	2.19	0.43
37:45:3:MET:HE2	37:45:3:MET:HB3	1.80	0.43
5:4E:35:GLY:HA3	5:4E:112:LEU:O	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:4L:37:G:H2'	25:4L:38:U:C6	2.53	0.43
34:58:112:LEU:HD12	34:58:112:LEU:HA	1.73	0.43
32:59:19:VAL:O	32:59:25:LYS:HE3	2.18	0.43
6:5E:33:TYR:CD2	6:5E:75:LEU:HD23	2.54	0.43
14:5I:25:VAL:HG22	14:5I:39:LEU:HD23	2.00	0.43
39:65:110:LEU:HD23	39:65:112:PHE:CE1	2.53	0.43
8:72:20:TYR:HE2	8:72:75:ARG:HG2	1.83	0.43
36:78:19:VAL:HG21	36:78:27:HIS:CG	2.53	0.43
16:7A:45:THR:O	16:7A:48:TRP:HD1	2.01	0.43
8:7E:120:THR:H	8:7E:123:GLU:HG3	1.84	0.43
41:85:66:ASN:OD1	41:85:76:TYR:N	2.46	0.43
37:88:45:GLN:H	37:88:45:GLN:CD	2.21	0.43
17:8A:6:LEU:HA	17:8A:6:LEU:HD23	1.76	0.43
42:95:84:LYS:HA	42:95:84:LYS:HD2	1.50	0.43
18:9A:22:VAL:HG12	18:9A:56:THR:HA	2.00	0.43
26:1H:1615:C:C2	43:E8:87:PRO:HG2	2.54	0.43
46:H8:14:LYS:HA	46:H8:15:PRO:HD2	1.75	0.43
46:H8:92:SER:O	46:H8:130:PRO:HG2	2.18	0.43
54:Q8:9:GLY:H	54:Q8:12:LYS:H	1.66	0.43
1:13:1346:A:H61	1:13:1374:A:H3'	1.84	0.43
1:13:1533:C:H2'	1:13:1534:A:O4'	2.18	0.43
1:13:46:G:H2'	1:13:366:C:H5	1.83	0.43
1:13:474:G:C6	1:13:475:G:C6	3.07	0.43
1:13:828:A:H2'	1:13:829:G:O4'	2.18	0.43
1:13:909:A:H3'	1:13:910:C:H6	1.84	0.43
26:14:2492:U:H2'	26:14:2493:U:H6	1.83	0.43
26:14:251:A:C5	26:14:252:G:H1'	2.54	0.43
26:14:2694:G:C5	26:14:2695:C:C5	3.07	0.43
26:14:2786:U:H2'	26:14:2787:C:O4'	2.19	0.43
26:14:321:G:C4	26:14:341:G:H4'	2.54	0.43
26:14:328:U:H4'	45:C5:68:HIS:CD2	2.54	0.43
26:14:481:G:C4	26:14:507:A:C2	3.07	0.43
26:14:947:G:H2'	26:14:948:G:H8	1.82	0.43
26:14:999:U:H5''	26:14:1154:G:O6	2.19	0.43
28:19:266:SER:O	28:19:269:PHE:HB2	2.19	0.43
2:1E:33:TYR:HB2	2:1E:43:ASP:HB2	2.00	0.43
1:1G:409:G:C6	1:1G:410:G:C4	3.07	0.43
1:1G:66:G:C2	1:1G:67:C:C6	3.07	0.43
26:1H:1168:G:C2	26:1H:1182:A:C2	3.07	0.43
26:1H:1790:C:H2'	26:1H:1791:A:C5	2.54	0.43
26:1H:1796:U:H2'	26:1H:1797:C:C6	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2165:G:N7	26:1H:2166:G:N2	2.51	0.43
22:1K:87:A:N6	26:1H:2583:G:O2'	2.52	0.43
26:1H:432:A:C6	26:1H:433:C:C4	3.07	0.43
26:1H:529:A:H8	26:1H:530:G:C6	2.36	0.43
26:1H:539:G:H2'	26:1H:540:G:H8	1.84	0.43
26:1H:586:A:P	57:1H:3735:HOH:O	2.76	0.43
26:1H:755:C:H2'	26:1H:756:C:C6	2.53	0.43
26:1H:807:U:H2'	26:1H:808:G:O4'	2.18	0.43
29:29:58:ARG:NH2	29:29:58:ARG:HG2	2.31	0.43
4:32:114:ARG:O	4:32:118:ARG:N	2.52	0.43
12:3A:60:LEU:HA	12:3A:60:LEU:HD13	1.85	0.43
12:3I:90:VAL:HG12	12:3I:91:LYS:N	2.34	0.43
31:41:108:ASN:HD22	51:M8:21:VAL:HG11	1.83	0.43
31:41:130:ASN:HB3	31:41:160:VAL:HA	2.00	0.43
5:42:83:GLU:O	5:42:83:GLU:HG3	2.18	0.43
32:51:15:VAL:HG12	32:51:28:GLY:HA3	2.01	0.43
38:55:24:GLN:OE1	38:55:36:THR:HG21	2.19	0.43
14:5A:37:PHE:CZ	14:5A:56:VAL:HG21	2.49	0.43
6:5E:18:GLN:HG2	6:5E:21:LEU:HD22	2.01	0.43
15:6I:6:GLU:H	15:6I:6:GLU:CD	2.22	0.43
43:A5:58:ALA:HB1	43:A5:64:MET:HB2	2.00	0.43
40:B8:57:PHE:O	40:B8:58:ASN:ND2	2.52	0.43
20:BA:14:LYS:HE2	20:BA:18:GLN:OE1	2.19	0.43
42:D8:43:GLU:HA	42:D8:44:LYS:HA	1.87	0.43
42:D8:76:LYS:O	42:D8:79:VAL:HG12	2.19	0.43
45:G8:95:LYS:O	45:G8:96:ILE:HG13	2.19	0.43
46:H8:4:ARG:CZ	46:H8:58:VAL:HG21	2.49	0.43
51:M8:15:ILE:HB	51:M8:32:TYR:CE2	2.53	0.43
54:Q8:6:THR:HG22	54:Q8:59:LYS:HD2	2.01	0.43
1:13:1142:G:C5	1:13:1143:G:H1'	2.54	0.42
1:13:1277:C:O2'	1:13:1279:A:H1'	2.18	0.42
1:13:232:G:C5	1:13:233:C:C5	3.07	0.42
1:13:384:G:C6	1:13:385:C:N4	2.87	0.42
1:13:755:G:O5'	1:13:755:G:H8	2.02	0.42
26:14:1005:C:H2'	26:14:1006:C:C6	2.54	0.42
26:14:130:C:O3'	26:14:1349:A:H1'	2.19	0.42
26:14:827:U:O2'	26:14:2068:U:C2	2.69	0.42
26:14:2232:U:P	48:F5:40:ARG:HH22	2.42	0.42
26:14:2402:C:H5	26:14:2415:G:H22	1.66	0.42
26:14:2656:U:H3	26:14:2665:A:H2	1.65	0.42
26:14:2801:A:H5''	26:14:2895:U:H4'	2.00	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:598:G:H5'	36:35:11:GLY:HA3	2.01	0.42
26:14:669:G:C2'	26:14:669:G:N3	2.81	0.42
26:14:1006:C:H1'	34:15:106:MET:HB3	2.01	0.42
34:15:1:MET:HB2	34:15:2:LYS:H	1.55	0.42
2:1E:162:ILE:O	2:1E:185:ILE:HG12	2.18	0.42
2:1E:21:ARG:HG3	2:1E:22:LYS:NZ	2.34	0.42
1:1G:1465:C:H2'	1:1G:1466:C:O4'	2.18	0.42
1:1G:750:G:O3'	15:6A:18:PHE:HZ	2.02	0.42
1:1G:909:A:H2'	1:1G:910:C:O4'	2.19	0.42
26:1H:1155:A:H3'	57:1H:3677:HOH:O	2.18	0.42
26:1H:1429:G:H1'	26:1H:1568:G:H1'	2.00	0.42
26:1H:174:C:H2'	26:1H:175:G:O4'	2.19	0.42
26:1H:1766:U:H2'	26:1H:1767:C:H6	1.83	0.42
26:1H:2801:A:OP2	26:1H:2895:U:O2'	2.25	0.42
26:1H:304:G:C2	26:1H:314:A:C2	3.06	0.42
26:1H:318:C:O2'	26:1H:319:C:H5'	2.18	0.42
26:1H:62:C:H5''	26:1H:63:U:OP2	2.20	0.42
26:1H:780:G:C2	26:1H:782:A:C2	3.07	0.42
10:1I:32:ALA:CB	10:1I:76:ASN:HB2	2.49	0.42
3:22:136:GLN:HA	3:22:139:GLN:HB3	2.01	0.42
3:22:119:ARG:HH22	3:22:140:ARG:HG2	1.82	0.42
3:22:154:SER:O	3:22:196:LEU:HD13	2.19	0.42
29:29:101:ARG:CZ	29:29:171:GLU:HB2	2.49	0.42
23:2K:75:G:O2'	23:2K:76:U:O5'	2.34	0.42
23:2L:21:A:C2	23:2L:46:G:C4	3.07	0.42
30:31:123:LEU:HD13	30:31:192:LEU:HB3	2.00	0.42
30:31:170:LEU:HA	30:31:170:LEU:HD13	1.85	0.42
30:31:23:ASP:CG	30:31:24:LEU:H	2.22	0.42
4:32:120:LEU:HD23	4:32:120:LEU:HA	1.81	0.42
4:32:173:TRP:O	4:32:186:LEU:HB2	2.19	0.42
36:35:88:LEU:HA	36:35:88:LEU:HD12	1.84	0.42
4:3E:148:VAL:HG21	4:3E:158:ILE:HG21	2.01	0.42
24:3L:15:A:H8	24:3L:16:U:C4	2.36	0.42
24:3L:33:U:H2'	24:3L:34:U:C5	2.54	0.42
24:3L:83:U:C2'	24:3L:84:A:H5'	2.48	0.42
13:4A:91:ARG:HB3	13:4A:98:VAL:HG12	2.01	0.42
32:51:151:ILE:H	32:51:151:ILE:HG13	1.63	0.42
38:55:76:VAL:O	38:55:80:PHE:N	2.52	0.42
32:59:30:LYS:NZ	32:59:136:ILE:O	2.52	0.42
7:62:42:ILE:HG23	7:62:117:ALA:HB2	2.01	0.42
33:69:33:ARG:HB3	33:69:35:LEU:HD21	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2848:G:C8	40:75:97:ALA:HB2	2.54	0.42
36:78:47:ASP:OD2	36:78:50:ARG:NH2	2.52	0.42
9:82:111:ARG:HB3	9:82:111:ARG:HE	1.49	0.42
41:85:99:ALA:HB2	41:85:106:PHE:CG	2.54	0.42
9:8E:18:PHE:HD2	9:8E:62:TYR:CD2	2.37	0.42
38:98:10:LEU:O	38:98:11:ASN:C	2.56	0.42
18:9A:21:LYS:HE3	18:9A:54:ARG:O	2.18	0.42
19:AA:15:LEU:HD22	19:AA:18:LYS:HB2	2.00	0.42
44:B5:49:VAL:HA	44:B5:87:GLN:OE1	2.19	0.42
40:B8:26:ASP:OD2	40:B8:120:ARG:NH2	2.40	0.42
20:BI:75:ASN:O	20:BI:79:ARG:HB2	2.18	0.42
45:C5:76:CYS:SG	45:C5:97:ARG:HG3	2.59	0.42
47:E5:60:PHE:CD1	47:E5:60:PHE:N	2.87	0.42
46:H8:4:ARG:NH1	46:H8:60:GLU:OE2	2.51	0.42
49:K8:47:ASN:HB2	49:K8:50:ILE:HD11	2.00	0.42
1:13:1316:G:N2	1:13:1318:A:H3'	2.34	0.42
1:13:1387:G:H2'	1:13:1388:C:H6	1.84	0.42
1:13:1413:A:H2'	1:13:1414:U:O4'	2.18	0.42
1:13:1442:G:H1	1:13:1461:G:H21	1.67	0.42
26:14:1540:G:H2'	26:14:1541:U:C6	2.54	0.42
26:14:1590:U:H2'	26:14:1591:G:C8	2.53	0.42
26:14:303:U:H2'	26:14:304:G:C8	2.54	0.42
26:14:380:U:H2'	26:14:381:G:C8	2.54	0.42
26:14:28:A:C2	26:14:513:A:C8	3.06	0.42
26:14:448:U:C4	26:14:583:G:H1'	2.54	0.42
27:16:109:G:C6	27:16:110:G:C5	3.07	0.42
28:19:37:LEU:O	28:19:37:LEU:HD12	2.19	0.42
28:19:52:ARG:HB2	28:19:53:PHE:CD2	2.54	0.42
1:1G:1176:A:O2'	1:1G:1177:G:H5'	2.19	0.42
1:1G:1423:G:H2'	1:1G:1424:C:C6	2.54	0.42
1:1G:186(A):C:H2'	1:1G:186(B):C:C6	2.54	0.42
1:1G:198:G:H2'	1:1G:199:G:H8	1.84	0.42
1:1G:448:A:H2'	1:1G:449:C:O2	2.19	0.42
26:1H:2093:G:C6	26:1H:2225:A:C8	3.07	0.42
26:1H:232:G:H5''	26:1H:232:G:C8	2.50	0.42
26:1H:2512:C:H4'	29:21:122:PHE:CE2	2.54	0.42
26:1H:459:U:H5''	53:P8:40:TRP:CD2	2.54	0.42
26:1H:547:A:C6	26:1H:548:A:N6	2.87	0.42
26:1H:78:A:H2'	26:1H:79:G:H8	1.84	0.42
29:21:128:SER:OG	29:21:129:HIS:N	2.50	0.42
29:21:3:GLY:HA3	29:21:81:ILE:HG21	2.00	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:29:81:ILE:O	29:29:82:ARG:HB2	2.20	0.42
23:2L:31:C:H2'	23:2L:32:C:C6	2.54	0.42
36:35:121:LYS:O	36:35:123:LEU:N	2.47	0.42
26:14:832:G:H21	36:35:53:GLY:HA3	1.83	0.42
12:3A:109:GLY:HA3	12:3A:121:GLY:O	2.19	0.42
12:3I:47:LYS:HA	12:3I:49:ASN:H	1.85	0.42
5:42:76:ILE:HG23	5:42:142:LEU:HD13	2.01	0.42
5:4E:110:LEU:HD13	5:4E:118:ILE:HG21	2.00	0.42
32:51:83:TYR:HA	32:51:135:GLY:H	1.82	0.42
33:61:7:GLU:HG2	33:61:9:LEU:HD22	2.01	0.42
35:68:29:ASN:OD1	35:68:29:ASN:N	2.44	0.42
33:69:65:ALA:O	33:69:69:LYS:N	2.52	0.42
1:1G:750:G:H21	15:6A:23:GLY:HA3	1.84	0.42
36:78:62:LEU:HB3	54:Q8:23:VAL:HG21	2.01	0.42
36:78:84:ASN:HB3	36:78:117:GLU:O	2.18	0.42
41:85:90:VAL:HA	42:95:39:LEU:HD22	2.01	0.42
39:A8:36:TYR:N	39:A8:36:TYR:HD1	2.17	0.42
1:13:1453:G:H2'	20:BI:39:LYS:HE2	2.01	0.42
45:C5:87:LYS:NZ	45:C5:89:PHE:HB3	2.33	0.42
41:C8:92:ARG:HD3	41:C8:94:ASN:HB2	2.01	0.42
41:C8:90:VAL:HG22	42:D8:39:LEU:HG	2.01	0.42
45:G8:29:GLU:HB3	45:G8:38:ILE:HG23	2.00	0.42
51:I5:14:ILE:HG12	51:I5:33:VAL:HG21	2.01	0.42
13:4A:3:ARG:HB2	51:I5:34:GLU:CG	2.48	0.42
52:J5:52:TYR:CD1	52:J5:53:ALA:N	2.86	0.42
51:M8:46:GLN:HG3	51:M8:48:ARG:HG2	2.01	0.42
54:Q8:57:ARG:H	54:Q8:57:ARG:HH11	1.66	0.42
2:12:6:THR:OG1	2:12:7:VAL:N	2.53	0.42
1:13:11:G:C5	1:13:12:U:C5	3.06	0.42
1:13:1203:C:H6	1:13:1203:C:O5'	2.02	0.42
1:13:49:U:C4	1:13:364:A:C5	3.08	0.42
26:14:1014:U:H2'	26:14:1015:G:C8	2.54	0.42
26:14:2131:G:OP2	26:14:2132:U:H5'	2.19	0.42
26:14:218:A:C2	26:14:235:U:H4'	2.54	0.42
26:14:2370:G:H2'	26:14:2371:G:O4'	2.19	0.42
26:14:2527:C:N4	26:14:2528:U:C4	2.87	0.42
26:14:455:C:N3	26:14:473:G:H5'	2.34	0.42
34:15:34:LEU:HA	34:15:34:LEU:HD12	1.71	0.42
10:1A:27:ALA:HA	10:1A:85:LEU:HD11	2.01	0.42
1:1G:1254:C:OP1	10:1A:45:ARG:HA	2.19	0.42
1:1G:113:G:H2'	1:1G:114:U:H6	1.84	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1199:U:H4'	10:1A:54:PHE:CZ	2.55	0.42
1:1G:1346:A:OP2	1:1G:1346:A:H3'	2.19	0.42
1:1G:200:G:O2'	1:1G:201:C:H5'	2.19	0.42
1:1G:514:C:H2'	1:1G:515:G:O4'	2.20	0.42
1:1G:749:C:H2'	1:1G:750:G:H8	1.84	0.42
1:1G:92:G:H2'	1:1G:93:U:O4'	2.19	0.42
26:1H:1228:G:C2'	26:1H:1229:G:H5'	2.49	0.42
26:1H:2213:U:H6	26:1H:2213:U:H2'	1.71	0.42
26:1H:2467:C:H4'	37:88:123:HIS:ND1	2.34	0.42
26:1H:2689:U:H4'	26:1H:2690:C:H5'	2.00	0.42
26:1H:483:A:C8	26:1H:484:C:C5	3.08	0.42
26:1H:998:C:H2'	26:1H:999:U:O4'	2.19	0.42
26:14:1027:A:H5'	27:1J:88:C:H41	1.84	0.42
27:1J:9:G:OP1	39:65:25:ARG:NH2	2.51	0.42
22:1K:2:C:H2'	22:1K:3:C:C6	2.54	0.42
1:1G:707:C:OP1	11:2A:85:ARG:NH1	2.52	0.42
11:2I:112:THR:HA	11:2I:113:PRO:HD3	1.90	0.42
30:31:12:LEU:HD11	30:31:17:ARG:CZ	2.49	0.42
4:32:101:LEU:HD23	4:32:121:VAL:CG1	2.50	0.42
4:32:148:VAL:HG12	4:32:152:SER:HB2	2.00	0.42
4:32:173:TRP:NE1	4:32:174:LEU:HG	2.34	0.42
30:39:132:VAL:C	30:39:134:GLY:H	2.22	0.42
12:3A:41:ARG:NH1	12:3A:41:ARG:HB3	2.33	0.42
4:3E:188:LEU:HD23	4:3E:188:LEU:HA	1.81	0.42
24:3K:20:U:H2'	24:3K:70:G:O6	2.20	0.42
24:3K:59:U:O2'	24:3K:60:A:P	2.77	0.42
24:3L:84:A:C8	24:3L:84:A:H3'	2.54	0.42
31:41:43:LEU:HG	31:41:45:GLU:HG3	2.02	0.42
26:14:871:U:OP1	37:45:5:ARG:HG2	2.19	0.42
13:4A:67:GLU:HG3	13:4A:68:GLY:H	1.84	0.42
34:58:107:LEU:HA	34:58:107:LEU:HD23	1.80	0.42
34:58:96:GLU:C	34:58:98:VAL:N	2.70	0.42
8:72:10:LEU:O	8:72:14:ARG:HB2	2.19	0.42
40:75:4:GLY:N	40:75:7:ILE:HG22	2.34	0.42
6:5E:100:ASN:O	18:9I:28:GLU:HG2	2.19	0.42
43:A5:20:VAL:HG11	43:A5:44:ALA:H	1.84	0.42
19:AA:51:VAL:HG12	19:AA:52:TYR:H	1.84	0.42
19:AI:15:LEU:HB2	19:AI:31:ILE:HD11	2.01	0.42
41:C8:27:LEU:HD22	41:C8:27:LEU:HA	1.71	0.42
46:D5:3:TYR:HB3	46:D5:47:VAL:HG13	2.01	0.42
46:D5:67:LEU:HA	46:D5:68:PRO:HD3	1.87	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:F5:75:GLU:O	48:F5:75:GLU:HG3	2.19	0.42
46:H8:24:LEU:HD22	46:H8:41:LEU:HD23	2.00	0.42
52:J5:31:VAL:HG13	52:J5:42:PRO:HG3	2.01	0.42
2:12:189:ASP:H	2:12:192:SER:HB2	1.84	0.42
2:12:32:ILE:N	2:12:43:ASP:HB2	2.34	0.42
1:13:17:U:O4'	1:13:1080:A:H1'	2.19	0.42
1:13:1120:G:N2	1:13:1153:C:O2	2.44	0.42
1:13:1292:U:H2'	1:13:1293:G:H8	1.81	0.42
1:13:313:A:H2'	1:13:314:C:H6	1.84	0.42
1:13:611:A:H61	1:13:629:G:H1	1.68	0.42
1:13:967:C:O5'	1:13:967:C:H6	2.02	0.42
26:14:1003:G:N2	26:14:1153:C:C2	2.88	0.42
26:14:1006:C:H1'	34:15:106:MET:HE3	2.00	0.42
26:14:1465:G:C4	26:14:1466:G:C8	3.06	0.42
26:14:1473:G:H2'	26:14:1474:C:O4'	2.20	0.42
26:14:1629:U:O2	26:14:2698:U:H5''	2.19	0.42
26:14:2139:C:N3	26:14:2153:G:H1'	2.34	0.42
26:14:2336:A:H61	47:E5:43:THR:HG21	1.83	0.42
26:14:2647:U:H3	26:14:2673:G:H1	1.67	0.42
26:14:2846:G:H2'	26:14:2847:U:O4'	2.20	0.42
26:14:357:A:H2'	26:14:358:U:C6	2.55	0.42
26:14:4:C:H42	26:14:2899:G:N2	1.94	0.42
26:14:590:A:OP1	30:39:95:ARG:NH1	2.52	0.42
34:15:101:HIS:O	34:15:104:LYS:HB3	2.19	0.42
10:1A:54:PHE:CD2	10:1A:55:LYS:HD2	2.54	0.42
2:1E:105:PHE:HE1	2:1E:152:PHE:CZ	2.37	0.42
1:1G:1205:U:H2'	1:1G:1206:G:C8	2.53	0.42
1:1G:551:U:O2'	1:1G:552:U:H5'	2.18	0.42
26:1H:1111:A:N3	26:1H:1112:G:H1'	2.33	0.42
26:1H:1820:U:C2	28:11:202:LYS:HD3	2.54	0.42
26:1H:2577:A:C5'	26:1H:2578:G:H5'	2.47	0.42
26:1H:307:G:N2	26:1H:309:G:H3'	2.34	0.42
26:1H:466:A:N3	26:1H:683:C:H1'	2.35	0.42
26:1H:654(H):G:N3	26:1H:654(H):G:H2'	2.34	0.42
26:1H:722:A:H2'	26:1H:723:G:H8	1.81	0.42
26:1H:883:G:N1	26:1H:893:C:N3	2.53	0.42
29:21:11:MET:HE2	29:21:11:MET:HB3	1.79	0.42
29:21:16:ARG:NH2	29:21:173:VAL:HG13	2.34	0.42
35:25:35:VAL:HG11	35:25:103:ALA:CB	2.42	0.42
11:2A:59:TYR:CZ	11:2A:63:LEU:HD21	2.55	0.42
36:35:55:ARG:HG2	36:35:56:SER:N	2.33	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:39:123:LEU:HA	30:39:192:LEU:O	2.19	0.42
30:39:10:PRO:HG2	30:39:13:SER:HB3	2.00	0.42
4:3E:150:GLU:HA	4:3E:153:ARG:CG	2.50	0.42
4:3E:188:LEU:HA	4:3E:189:PRO:HD3	1.82	0.42
31:41:113:ARG:HD3	31:41:140:ILE:O	2.19	0.42
31:49:101:ILE:O	31:49:105:LYS:HG3	2.19	0.42
5:4E:118:ILE:HG12	5:4E:119:LEU:N	2.35	0.42
32:51:4:ILE:HG12	32:51:4:ILE:O	2.18	0.42
32:51:6:ARG:HG2	32:51:66:GLY:N	2.34	0.42
26:1H:270(L):U:O2	33:61:50:ARG:HG2	2.19	0.42
35:68:25:LEU:HD21	35:68:40:VAL:HG23	2.01	0.42
35:68:64:ARG:HG2	35:68:79:PHE:CD2	2.54	0.42
7:6E:101:LEU:O	7:6E:105:VAL:HG23	2.20	0.42
1:13:933:G:OP2	7:6E:3:ARG:HB2	2.19	0.42
8:72:121:ASP:OD2	8:72:125:ARG:NH2	2.52	0.42
36:78:76:LYS:HA	36:78:76:LYS:HD3	1.79	0.42
9:82:105:ASP:OD1	9:82:107:ARG:HD3	2.20	0.42
9:8E:102:LEU:HD23	9:8E:102:LEU:HA	1.83	0.42
9:8E:9:ARG:HD2	9:8E:14:VAL:HG13	2.02	0.42
42:95:2:PHE:O	42:95:42:GLY:N	2.52	0.42
42:95:44:LYS:HG2	42:95:45:THR:OG1	2.19	0.42
26:14:993:G:N3	42:95:89:GLN:NE2	2.67	0.42
38:98:45:ARG:HB3	38:98:46:GLY:H	1.58	0.42
19:AA:12:ASP:N	19:AA:12:ASP:OD1	2.50	0.42
20:BA:73:HIS:HB3	20:BA:74:LYS:HZ2	1.84	0.42
41:C8:58:ARG:HA	41:C8:61:TRP:CE3	2.55	0.42
51:M8:15:ILE:HG23	51:M8:20:ASN:ND2	2.35	0.42
54:Q8:57:ARG:CB	54:Q8:59:LYS:HE2	2.49	0.42
2:12:91:PRO:HG2	2:12:155:LEU:CG	2.44	0.42
1:13:1009:G:N2	1:13:1010:G:H1'	2.34	0.42
1:13:102:G:C6	1:13:103:C:C4	3.07	0.42
1:13:1108:G:H2'	1:13:1108:G:N3	2.34	0.42
1:13:1170:A:H8	1:13:1170:A:O5'	2.03	0.42
1:13:1478:C:H2'	1:13:1479:C:C6	2.54	0.42
1:13:1492:A:H1'	25:4K:50:U:O2'	2.19	0.42
1:13:350:G:O2'	1:13:351:G:H5'	2.19	0.42
1:13:353:A:C8	1:13:353:A:H5'	2.53	0.42
1:13:924:C:H2'	1:13:925:G:C8	2.54	0.42
1:13:947:G:H2'	1:13:948:C:C6	2.55	0.42
26:14:1869:G:N2	26:14:1872:A:C8	2.88	0.42
26:14:273(D):C:H5'	26:14:273(E):U:OP2	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:548:A:C6	26:14:549:G:H1'	2.54	0.42
1:1G:1256:A:H62	1:1G:1277:C:H3'	1.84	0.42
1:1G:243:A:C2	1:1G:245:C:C2	3.07	0.42
1:1G:746:A:H2'	1:1G:747:C:C6	2.53	0.42
1:1G:570:G:C6	1:1G:873:A:C2	3.08	0.42
26:1H:1178:C:HO2'	26:1H:1179:C:H5	1.67	0.42
26:1H:1488:G:C5	26:1H:1489:U:C5	3.07	0.42
26:1H:1844:C:H2'	26:1H:1845:G:H8	1.85	0.42
26:1H:2024:G:H2'	26:1H:2025:C:C6	2.55	0.42
26:1H:2108:C:H2'	26:1H:2109:U:O4'	2.20	0.42
26:1H:2325:G:C4	26:1H:2326:C:C5	3.07	0.42
26:1H:2469:A:N3	26:1H:2469:A:H5'	2.34	0.42
26:1H:455:C:N3	26:1H:473:G:H5'	2.35	0.42
26:1H:562:U:O4	26:1H:2036:C:H1'	2.20	0.42
26:1H:994:C:OP1	41:C8:53:ARG:NH2	2.52	0.42
10:1I:54:PHE:HZ	10:1I:55:LYS:HZ1	1.61	0.42
27:1J:3:C:H2'	27:1J:4:C:C6	2.54	0.42
11:2A:84:VAL:HG11	11:2A:95:ILE:HD11	2.01	0.42
3:2E:16:ARG:HB2	3:2E:16:ARG:HH11	1.84	0.42
3:2E:59:ARG:NH1	3:2E:97:LYS:HE3	2.33	0.42
30:31:120:GLU:CB	30:31:122:LYS:HG2	2.50	0.42
30:31:64:ILE:HD12	30:31:64:ILE:O	2.19	0.42
36:35:110:TYR:HB3	36:35:111:ARG:H	1.66	0.42
30:39:110:LEU:O	30:39:114:VAL:HG23	2.19	0.42
12:3A:100:ILE:HG22	12:3A:101:VAL:H	1.84	0.42
12:3A:43:VAL:HG23	12:3A:55:VAL:HG11	2.02	0.42
12:3A:38:THR:HG21	12:3A:65:GLU:OE2	2.20	0.42
1:1G:523:A:N6	12:3A:92:ASP:HB2	2.33	0.42
4:3E:112:VAL:HG12	4:3E:116:GLN:OE1	2.20	0.42
4:3E:79:PHE:HE1	4:3E:204:ILE:HD12	1.84	0.42
24:3L:11:U:H2'	24:3L:12:G:N7	2.33	0.42
26:1H:2314:C:H5''	31:41:38:VAL:HG11	2.01	0.42
31:41:98:ARG:HA	31:41:101:ILE:HG23	2.02	0.42
5:42:71:LEU:HD21	5:42:115:VAL:HG22	2.02	0.42
37:45:17:LEU:HD21	37:45:41:TRP:NE1	2.33	0.42
32:51:124:GLU:O	32:51:132:ARG:N	2.53	0.42
32:51:62:LYS:HE2	32:51:62:LYS:HB3	1.79	0.42
6:52:67:MET:HB2	6:52:68:PRO:HD2	2.02	0.42
38:55:53:HIS:HA	38:55:56:LYS:HD3	2.01	0.42
32:59:26:VAL:HG12	32:59:33:LEU:N	2.30	0.42
26:14:2748:A:O2'	32:59:66:GLY:HA3	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:5E:25:ILE:HG23	6:5E:28:ARG:HH12	1.85	0.42
39:65:91:PRO:HG2	39:65:92:TYR:CE1	2.54	0.42
33:69:145:VAL:HB	33:69:146:ALA:H	1.61	0.42
33:69:51:ILE:O	33:69:54:GLN:HG2	2.19	0.42
8:72:110:ALA:HB3	8:72:121:ASP:HB3	2.02	0.42
5:4E:148:VAL:HG21	8:7E:107:LEU:HD13	2.02	0.42
1:1G:967:C:H4'	9:82:125:TYR:OH	2.19	0.42
9:82:95:LYS:NZ	9:82:96:LEU:HB2	2.34	0.42
9:82:95:LYS:HE2	9:82:96:LEU:N	2.33	0.42
9:8E:28:VAL:N	9:8E:31:GLN:O	2.26	0.42
17:8I:90:ILE:HA	17:8I:93:GLN:HB3	2.01	0.42
18:9A:87:ARG:HB3	18:9A:88:LYS:H	1.50	0.42
39:A8:110:LEU:HA	39:A8:111:GLU:HA	1.90	0.42
46:D5:31:ARG:HB2	46:D5:31:ARG:HE	1.48	0.42
26:14:2255:G:N2	47:E5:9:SER:H	2.09	0.42
47:I8:53:MET:HG3	47:I8:59:LEU:HD23	2.01	0.42
53:P8:18:PHE:CE1	53:P8:22:MET:HG3	2.54	0.42
26:1H:2239:G:H5'	28:11:251:GLY:HA3	2.01	0.42
2:12:129:GLU:HB3	2:12:130:ARG:H	1.56	0.42
1:13:1071:C:H2'	1:13:1072:G:C8	2.54	0.42
1:13:1206:G:C6	1:13:1207:G:C5	3.08	0.42
1:13:1236:A:OP1	21:1F:3:LYS:HE3	2.20	0.42
1:13:1364:U:O2'	1:13:1365:G:H5'	2.19	0.42
1:13:1399:C:C2	1:13:1502:A:N6	2.87	0.42
1:13:347:G:OP2	1:13:347:G:H8	2.03	0.42
1:13:746:A:H2'	1:13:747:C:H6	1.85	0.42
26:14:1202:C:N4	26:14:1243:G:H1	2.17	0.42
26:14:1275:A:C5	38:55:16:HIS:CD2	3.07	0.42
26:14:1963:U:H2'	26:14:1963:U:O2	2.20	0.42
26:14:2274:A:C5	26:14:2276:G:C8	3.06	0.42
26:14:2666:C:H3'	26:14:2667:C:H6	1.84	0.42
26:14:335:C:O5'	26:14:335:C:H6	2.02	0.42
26:14:444:C:OP2	41:85:2:PRO:HD3	2.19	0.42
26:14:260:G:O4'	26:14:621:A:H1'	2.20	0.42
26:14:638:G:H2'	26:14:639:U:O4'	2.19	0.42
26:14:848:G:N9	26:14:933:A:H8	2.17	0.42
26:14:996:A:H2'	26:14:997:G:H8	1.84	0.42
1:1G:1136:U:H5''	1:1G:1137:C:C5	2.55	0.42
1:1G:790:A:C6	1:1G:791:G:C6	3.08	0.42
1:1G:894:G:C6	1:1G:895:G:C6	3.08	0.42
1:1G:962:C:H1'	1:1G:1201:A:C6	2.54	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1385:G:H1'	26:1H:1386:C:C6	2.54	0.42
26:1H:1426:G:H2'	26:1H:1427:A:C8	2.55	0.42
26:1H:1671:U:H2'	26:1H:1672:C:H5''	2.02	0.42
26:1H:551:G:OP1	42:D8:68:LYS:NZ	2.52	0.42
26:1H:614:U:O4	30:31:175:THR:HG22	2.19	0.42
26:1H:817:C:C4	26:1H:818:G:C5	3.08	0.42
26:1H:934:G:H2'	26:1H:935:C:C6	2.53	0.42
3:22:131:ARG:HG3	3:22:166:GLU:OE1	2.20	0.42
3:2E:32:LEU:HD23	3:2E:32:LEU:HA	1.89	0.42
23:2K:17:H2U:HO2'	23:2K:68:A:H2	1.66	0.42
30:31:101:LEU:HA	30:31:101:LEU:HD23	1.77	0.42
12:3I:78:GLN:HB3	12:3I:79:GLU:H	1.64	0.42
24:3K:5:A:O2'	24:3K:6:G:H5'	2.20	0.42
31:41:34:LEU:HA	31:41:34:LEU:HD22	1.86	0.42
5:42:118:ILE:HG12	5:42:119:LEU:N	2.35	0.42
31:49:12:TYR:O	31:49:16:ARG:HB3	2.19	0.42
13:4A:77:ASN:HD22	13:4A:77:ASN:C	2.23	0.42
6:52:96:PRO:HB3	18:9A:30:ASP:CG	2.40	0.42
32:59:106:THR:HG22	32:59:112:PRO:HB2	2.02	0.42
32:59:141:VAL:O	32:59:145:ALA:N	2.45	0.42
33:61:75:LEU:HD23	33:61:105:HIS:CD2	2.54	0.42
33:61:68:LEU:HA	33:61:71:ILE:HG22	2.01	0.42
33:61:77:LEU:HB2	33:61:140:LEU:HB3	2.01	0.42
8:7E:21:LYS:O	8:7E:65:TYR:OH	2.30	0.42
41:85:114:LYS:H	41:85:114:LYS:HG2	1.49	0.42
37:88:16:ARG:HE	37:88:16:ARG:HB3	1.41	0.42
37:88:34:LEU:HB2	37:88:118:LEU:HD22	2.01	0.42
17:8I:18:THR:HG23	17:8I:69:LYS:HD2	2.00	0.42
39:A8:9:ARG:O	39:A8:12:PHE:N	2.52	0.42
45:C5:19:LYS:C	45:C5:21:LYS:H	2.22	0.42
47:E5:21:LEU:HD23	47:E5:21:LEU:HA	1.85	0.42
44:F8:2:LYS:HG2	49:K8:26:ARG:HE	1.84	0.42
50:H5:30:ARG:HG3	50:H5:30:ARG:H	1.57	0.42
51:I5:38:LYS:HA	51:I5:44:THR:HG21	2.02	0.42
49:K8:21:LEU:HD23	49:K8:21:LEU:HA	1.90	0.42
28:11:62:TYR:HA	28:11:87:ASN:OD1	2.20	0.42
2:12:108:ILE:HA	2:12:108:ILE:HD13	1.75	0.42
2:12:109:SER:O	2:12:113:HIS:ND1	2.52	0.42
1:13:1092:A:C6	1:13:1093:A:C6	3.08	0.42
1:13:10:A:H2'	1:13:11:G:H8	1.85	0.42
1:13:1317:C:H2'	1:13:1318:A:O4'	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:343:U:O2'	1:13:346:G:O6	2.31	0.42
1:13:484:G:O2'	1:13:485:G:OP2	2.33	0.42
26:14:1053:C:H42	26:14:1106:G:H1	1.67	0.42
26:14:1499:C:H2'	26:14:1500:G:C8	2.55	0.42
26:14:2468:G:N2	26:14:2481:G:O2'	2.38	0.42
26:14:2702:U:HO2'	26:14:2703:C:H6	1.64	0.42
26:14:2833:G:H3'	26:14:2833:G:C8	2.54	0.42
26:14:28:A:C5	26:14:29:U:C5	3.07	0.42
26:14:4:C:O2'	26:14:5:A:OP1	2.31	0.42
26:14:669:G:N3	26:14:669:G:H2'	2.34	0.42
26:14:826:U:H2'	26:14:828:U:O4'	2.19	0.42
26:14:861:A:C2	26:14:917:A:C4	3.08	0.42
26:14:880:G:H1	26:14:897:C:H42	1.66	0.42
26:14:957:A:N1	26:14:2459:A:H8	2.18	0.42
28:19:267:SER:O	28:19:268:ARG:HG2	2.19	0.42
28:19:89:SER:HB2	28:19:159:ALA:H	1.85	0.42
2:1E:67:THR:HA	2:1E:90:MET:SD	2.60	0.42
1:1G:1424:C:H2'	1:1G:1425:U:O4'	2.20	0.42
1:1G:191:G:H1'	20:BA:104:LEU:O	2.19	0.42
1:1G:568:G:N3	1:1G:574:A:H2	2.18	0.42
1:1G:743:U:H2'	1:1G:744:C:C6	2.55	0.42
1:1G:833:U:O2	1:1G:854:G:C2	2.73	0.42
26:1H:1141:U:H4'	26:1H:1142(A):A:O4'	2.19	0.42
26:1H:2592:G:C6	26:1H:2593:U:N3	2.88	0.42
27:1J:88:C:H4'	27:1J:89:G:OP1	2.17	0.42
22:1K:9:G:N3	22:1K:9:G:H2'	2.35	0.42
22:1L:15:A:N1	22:1L:59:U:N3	2.68	0.42
29:21:46:ALA:HB2	29:21:82:ARG:HA	2.02	0.42
3:2E:70:VAL:N	3:2E:106:VAL:HG23	2.35	0.42
23:2L:86:C:H3'	23:2L:87:A:C5'	2.46	0.42
30:31:32:LEU:O	30:31:36:VAL:HG23	2.19	0.42
4:32:45:GLN:C	4:32:46:LYS:HG2	2.40	0.42
4:32:70:ILE:CG2	4:32:75:PHE:HB2	2.50	0.42
30:39:63:LYS:HA	30:39:76:GLY:O	2.19	0.42
12:3A:110:VAL:CG2	12:3A:120:TYR:HB3	2.50	0.42
6:52:61:LEU:HB3	6:52:63:TYR:CE1	2.55	0.42
34:58:91:LEU:HA	34:58:91:LEU:HD23	1.71	0.42
32:59:89:ILE:HD12	32:59:130:ARG:HA	2.02	0.42
32:59:30:LYS:HZ1	32:59:136:ILE:HA	1.84	0.42
32:59:47:GLU:O	32:59:49:VAL:HG22	2.20	0.42
32:59:55:PRO:HG2	32:59:61:HIS:CD2	2.55	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2198:A:C2	33:69:29:TYR:HB2	2.55	0.42
8:72:97:VAL:HG22	8:72:129:VAL:C	2.40	0.42
36:78:27:HIS:ND1	36:78:27:HIS:N	2.68	0.42
5:4E:143:ARG:NE	8:7E:77:GLU:OE1	2.52	0.42
41:85:19:LYS:O	41:85:22:LYS:HG3	2.20	0.42
41:85:50:ARG:HG2	41:85:53:ARG:NH2	2.35	0.42
26:1H:953:A:OP2	37:88:16:ARG:HD3	2.20	0.42
9:8E:32:ASP:HB3	9:8E:35:GLU:HB3	2.01	0.42
42:95:38:LEU:HD13	42:95:55:ALA:HB1	2.02	0.42
26:1H:2378:A:H5''	39:A8:23:ARG:NH2	2.35	0.42
39:A8:30:ARG:HG2	39:A8:30:ARG:NH1	2.34	0.42
40:B8:106:SER:HB2	40:B8:109:GLU:H	1.84	0.42
20:BA:14:LYS:O	20:BA:18:GLN:HG3	2.20	0.42
43:E8:71:VAL:HA	43:E8:107:LEU:HD12	2.00	0.42
54:Q8:53:PRO:HA	54:Q8:55:ALA:H	1.81	0.42
2:12:119:GLU:HA	2:12:122:PHE:CD2	2.54	0.42
1:13:129(A):G:HO2'	1:13:189:U:H6	1.66	0.42
1:13:1342:C:H2'	1:13:1343:G:H8	1.81	0.42
1:13:297:G:H4'	1:13:557:G:H4'	2.02	0.42
26:14:1028:A:H2'	26:14:1029:A:C8	2.54	0.42
26:14:1337:G:H2'	26:14:1338:G:H8	1.85	0.42
26:14:1475:G:H5'	26:14:1476:C:OP2	2.20	0.42
26:14:1480:G:C6	26:14:1482:U:C4	3.07	0.42
26:14:1569:A:H2'	26:14:1570:A:C8	2.55	0.42
26:14:1628:G:H2'	26:14:1629:U:C6	2.55	0.42
26:14:2239:G:OP2	28:19:244:ARG:NH2	2.46	0.42
26:14:69:C:H2'	26:14:70:G:C8	2.54	0.42
26:14:741:G:H2'	26:14:742:G:C8	2.55	0.42
34:15:134:ARG:N	34:15:135:PRO:HD3	2.33	0.42
27:16:55:U:O2'	27:16:57:A:N7	2.52	0.42
26:14:1903:G:P	28:19:241:PRO:HB2	2.60	0.42
1:1G:1157:A:N6	1:1G:1177:G:H1	2.17	0.42
1:1G:1359:C:OP1	14:5A:22:THR:OG1	2.37	0.42
1:1G:446:G:H2'	1:1G:447:G:O4'	2.20	0.42
1:1G:835:U:H3	1:1G:851:G:H1	1.66	0.42
1:1G:992:U:H3	1:1G:1044:A:N6	2.12	0.42
26:1H:1444:G:C2	26:1H:1548:C:C2	3.08	0.42
26:1H:2025:C:OP1	29:21:149:ARG:HD3	2.20	0.42
26:1H:2098:U:H2'	26:1H:2099:U:O4'	2.20	0.42
26:1H:2283:C:H2'	26:1H:2284:C:O4'	2.20	0.42
26:1H:2747:G:O6	26:1H:2755:C:H5''	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2846:G:C6	26:1H:2847:U:C4	3.08	0.42
26:1H:508:G:N3	26:1H:508:G:H5''	2.34	0.42
26:1H:734:A:O2'	26:1H:1635:G:H5'	2.19	0.42
26:1H:848:G:O6	26:1H:929:G:H2'	2.19	0.42
26:1H:950:G:H2'	26:1H:951:C:C6	2.55	0.42
26:1H:998:C:OP2	41:C8:93:LYS:NZ	2.46	0.42
10:1I:77:PRO:HB2	10:1I:79:ARG:NH1	2.34	0.42
22:1L:1:G:H1	22:1L:83:U:H3	1.66	0.42
29:21:101:ARG:NH2	29:21:171:GLU:HB2	2.35	0.42
35:25:25:LEU:HB2	35:25:38:VAL:O	2.19	0.42
23:2K:18:OMG:H4'	23:2K:19:G:O5'	2.19	0.42
23:2L:62:G:N2	23:2L:63:G:N3	2.68	0.42
4:3E:8:VAL:O	4:3E:11:LEU:N	2.41	0.42
37:45:50:ALA:O	37:45:53:ALA:HB3	2.19	0.42
32:51:155:SER:OG	32:51:156:ALA:N	2.53	0.42
38:55:27:SER:HB3	38:55:34:ILE:HD11	2.00	0.42
32:59:101:ARG:HG3	32:59:102:ALA:N	2.35	0.42
3:22:29:TYR:OH	14:5A:54:PRO:HD2	2.19	0.42
7:62:150:ALA:HB1	11:2A:57:THR:HG21	2.00	0.42
8:72:33:GLU:OE1	8:72:48:TYR:HE1	2.03	0.42
40:75:18:ASP:N	40:75:18:ASP:OD1	2.53	0.42
16:7I:33:ILE:H	16:7I:33:ILE:HG13	1.68	0.42
9:82:46:ALA:HA	9:82:78:LYS:HB2	2.00	0.42
37:88:106:VAL:HG21	37:88:114:ALA:HB1	2.02	0.42
37:88:20:ALA:HB3	46:H8:79:ARG:CZ	2.49	0.42
37:88:79:LEU:HD23	37:88:79:LEU:HA	1.56	0.42
1:1G:247:G:OP2	17:8A:100:LYS:N	2.53	0.42
26:1H:1653:G:C6	38:98:9:LYS:HG3	2.55	0.42
18:9I:53:ARG:HH21	18:9I:59:SER:HA	1.84	0.42
19:AA:32:LYS:HB3	19:AA:57:HIS:ND1	2.35	0.42
40:B8:16:ARG:HE	40:B8:19:LEU:HD21	1.85	0.42
20:BA:70:SER:HA	20:BA:73:HIS:NE2	2.34	0.42
45:C5:83:THR:HG22	45:C5:84:ARG:H	1.84	0.42
46:D5:91:LEU:H	46:D5:91:LEU:HD23	1.84	0.42
47:I8:23:VAL:HG22	47:I8:38:VAL:HG22	2.02	0.42
49:K8:22:GLU:O	49:K8:25:VAL:HG12	2.20	0.42
49:K8:50:ILE:HD12	49:K8:51:ARG:N	2.33	0.42
51:M8:16:CYS:HB3	51:M8:18:CYS:SG	2.59	0.42
28:11:136:ILE:HG12	28:11:136:ILE:H	1.59	0.42
1:13:12:U:O2'	1:13:526:C:H4'	2.19	0.42
1:13:157:G:H2'	1:13:158:G:H8	1.83	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:82:G:N1	26:14:103:A:OP2	2.48	0.42
26:14:1047:G:H2'	26:14:1110:G:H22	1.85	0.42
26:14:1784:A:H4'	26:14:1785:A:C5'	2.50	0.42
26:14:2178:C:H2'	26:14:2179:C:H6	1.83	0.42
26:14:2370:G:C6	26:14:2371:G:C6	3.07	0.42
26:14:254:G:N7	54:M5:5:LYS:HE2	2.35	0.42
26:14:2615:U:C2	52:J5:7:PRO:HA	2.55	0.42
26:14:299:A:N1	26:14:322:A:O2'	2.38	0.42
26:14:654(C):G:C2	26:14:654(S):G:N1	2.88	0.42
28:19:11:PRO:C	28:19:13:ARG:H	2.23	0.42
28:19:72:LYS:HD3	28:19:97:TYR:CE2	2.55	0.42
1:1G:1023:G:N3	1:1G:1023:G:H2'	2.35	0.42
1:1G:1245:A:H8	1:1G:1245:A:O5'	2.03	0.42
1:1G:270:A:H5''	1:1G:271:C:OP2	2.19	0.42
1:1G:350:G:H5''	1:1G:350:G:H8	1.84	0.42
26:1H:1167:U:H2'	26:1H:1168:G:H8	1.84	0.42
26:1H:1558:A:H4'	26:1H:1559:G:H5'	2.02	0.42
26:1H:2125:G:H1	26:1H:2171:A:H5''	1.85	0.42
26:1H:2287:A:H2	26:1H:2346:A:H2	1.66	0.42
26:1H:306:U:H2'	26:1H:307:G:O4'	2.20	0.42
26:1H:547:A:C6	26:1H:548:A:C6	3.08	0.42
26:1H:676:A:N1	26:1H:802:A:N1	2.67	0.42
26:1H:664:C:H4'	26:1H:941:A:OP1	2.20	0.42
3:22:114:PRO:O	3:22:118:GLN:NE2	2.36	0.42
1:1G:1190:G:H3'	3:22:3:ASN:OD1	2.20	0.42
29:29:182:LEU:HA	29:29:182:LEU:HD12	1.75	0.42
23:2L:65:5MU:H73	23:2L:66:PSU:O2	2.19	0.42
26:1H:616:A:C4	30:31:180:GLY:HA2	2.55	0.42
1:1G:490:G:OP2	4:32:132:ARG:NH2	2.53	0.42
4:32:45:GLN:HG2	4:32:45:GLN:H	1.64	0.42
4:32:4:TYR:HE2	4:32:11:LEU:HD21	1.85	0.42
12:3A:59:ARG:NH2	12:3A:65:GLU:OE1	2.53	0.42
24:3L:15:A:N6	24:3L:70:G:O2'	2.52	0.42
31:49:152:LEU:HA	31:49:152:LEU:HD22	1.85	0.42
32:51:54:ARG:HA	32:51:55:PRO:HD3	1.84	0.42
7:6E:150:ALA:HB2	11:2I:50:TYR:CE2	2.54	0.42
8:7E:68:ARG:O	8:7E:68:ARG:HG3	2.18	0.42
1:1G:1291:G:O2'	9:82:38:GLN:OE1	2.27	0.42
9:8E:83:ARG:O	9:8E:86:VAL:HG12	2.19	0.42
38:98:98:LEU:HA	38:98:98:LEU:HD23	1.77	0.42
1:1G:719:C:O2'	18:9A:49:LYS:HB3	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
43:A5:95:ILE:HG12	43:A5:95:ILE:O	2.20	0.42
20:BA:20:LEU:HD23	20:BA:20:LEU:HA	1.84	0.42
45:C5:54:LYS:C	45:C5:55:TYR:CG	2.93	0.42
41:C8:82:GLY:HA3	41:C8:113:ALA:HB1	2.01	0.42
26:1H:996:A:H4'	41:C8:92:ARG:NE	2.35	0.42
26:1H:1188:U:H4'	42:D8:79:VAL:HG22	2.02	0.42
43:E8:28:SER:OG	43:E8:31:GLU:HG2	2.20	0.42
44:B5:9:LEU:CA	49:G5:36:ARG:HH21	2.31	0.42
50:H5:4:LEU:O	50:H5:36:VAL:HA	2.20	0.42
53:P8:10:ARG:O	53:P8:14:LYS:HB2	2.19	0.42
1:13:113:G:H2'	1:13:114:U:H6	1.85	0.42
1:13:1266:G:N2	1:13:1270:C:C2	2.88	0.42
1:13:1330:U:H3'	1:13:1331:G:O4'	2.19	0.42
1:13:1336:C:H4'	1:13:1337:G:O5'	2.20	0.42
1:13:390:C:O3'	16:7I:28:ARG:NH2	2.49	0.42
1:13:577:G:C8	1:13:816:A:C6	3.08	0.42
1:13:5:U:H4'	1:13:6:G:O5'	2.19	0.42
26:14:1027:A:N6	26:14:1126:A:C4	2.88	0.42
26:14:171:G:H2'	26:14:172:C:C6	2.55	0.42
26:14:1788:C:H2'	26:14:1789:A:O4'	2.20	0.42
26:14:919:G:N2	26:14:2269:A:OP2	2.53	0.42
26:14:2564:A:OP1	26:14:2648:C:H4'	2.19	0.42
26:14:2686:G:C2	26:14:2724:C:O2	2.73	0.42
26:14:603:A:H4'	26:14:604:G:O5'	2.20	0.42
26:14:780:G:H5''	26:14:781:A:P	2.60	0.42
28:19:148:GLU:CB	28:19:151:LYS:HD2	2.50	0.42
2:1E:70:PHE:HE1	2:1E:90:MET:HB2	1.85	0.42
2:1E:74:LYS:HE3	2:1E:74:LYS:HB3	1.93	0.42
1:1G:954:G:O6	1:1G:1225:A:N6	2.53	0.42
1:1G:1258:G:H2'	1:1G:1259:C:C6	2.55	0.42
1:1G:1239:A:C4	1:1G:1298:C:N4	2.88	0.42
1:1G:1413:A:H2'	1:1G:1414:U:O4'	2.19	0.42
1:1G:20:U:H2'	1:1G:21:G:O4'	2.20	0.42
1:1G:690:G:H22	11:2A:55:LYS:NZ	2.18	0.42
26:1H:1015:G:H1	26:1H:1147:C:H42	1.68	0.42
26:1H:1094:U:H1'	26:1H:1096:A:H5'	2.00	0.42
26:1H:1044:G:HO2'	26:1H:1111:A:N6	2.18	0.42
26:1H:1178:C:H1'	26:1H:1179:C:C6	2.55	0.42
26:1H:1424:G:H2'	26:1H:1425:G:O4'	2.20	0.42
26:1H:1538:G:H2'	26:1H:1539:G:C8	2.53	0.42
26:1H:1693:U:H1'	28:11:14:ARG:NH2	2.35	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2241:A:O2'	26:1H:2242:G:H5'	2.18	0.42
26:1H:2726:U:O2'	26:1H:2727:G:H8	2.02	0.42
26:1H:2795:G:H3'	26:1H:2797:U:H5''	2.01	0.42
26:1H:41:C:N4	26:1H:438:G:H1	2.13	0.42
26:1H:442:G:H4'	30:31:46:ARG:HG3	2.02	0.42
22:1L:42:U:H2'	22:1L:43:A:O4'	2.19	0.42
22:1L:79:U:H2'	22:1L:80:C:C6	2.55	0.42
3:22:32:LEU:HD22	3:22:59:ARG:NH2	2.32	0.42
35:25:117:LEU:HA	35:25:117:LEU:HD13	1.89	0.42
35:25:7:TYR:CE1	35:25:20:MET:HB2	2.55	0.42
29:29:101:ARG:HD2	29:29:169:ASN:ND2	2.35	0.42
11:2A:48:ILE:HG13	11:2A:63:LEU:HB3	2.02	0.42
23:2K:72:C:H2'	23:2K:73:C:C6	2.55	0.42
4:32:19:LEU:HB3	4:32:21:LEU:HD21	2.02	0.42
36:35:62:LEU:HA	36:35:63:PRO:HD3	1.49	0.42
13:4A:3:ARG:NH2	31:49:113:ARG:HE	2.18	0.42
31:49:66:GLN:NE2	31:49:93:THR:O	2.53	0.42
5:4E:7:GLU:N	5:4E:35:GLY:O	2.48	0.42
32:51:68:THR:O	32:51:72:ILE:HD12	2.20	0.42
6:52:61:LEU:HB3	6:52:63:TYR:HE1	1.84	0.42
32:59:116:GLU:HA	32:59:117:PRO:HD2	1.93	0.42
7:62:71:PRO:HD3	7:62:103:TRP:CZ3	2.55	0.42
33:69:125:GLU:OE2	33:69:141:LYS:NZ	2.51	0.42
15:6A:53:HIS:O	15:6A:56:LEU:N	2.52	0.42
7:6E:49:ILE:O	7:6E:53:LYS:HG3	2.20	0.42
8:72:29:SER:HB3	8:72:32:LYS:HG3	2.02	0.42
16:7A:23:ASP:OD1	16:7A:25:ARG:HG3	2.19	0.42
1:13:136:C:O2'	16:7I:65:GLN:HG3	2.20	0.42
41:85:59:ARG:O	41:85:63:VAL:HG22	2.20	0.42
41:85:92:ARG:NH1	42:95:11:GLN:H	2.17	0.42
9:8E:93:ARG:HH11	9:8E:93:ARG:HB2	1.85	0.42
42:95:35:LEU:HB3	42:95:37:VAL:HG13	2.01	0.42
19:AI:64:GLU:O	19:AI:67:VAL:HG13	2.20	0.42
44:B5:32:PRO:HA	44:B5:77:LYS:HB2	2.02	0.42
46:D5:164:ALA:O	46:D5:165:VAL:HG13	2.20	0.42
47:E5:49:LYS:O	47:E5:50:ASN:HB2	2.19	0.42
46:H8:52:SER:O	46:H8:53:ILE:HG12	2.19	0.42
51:I5:58:ARG:HH11	51:I5:61:ARG:NE	2.17	0.42
1:13:1084:G:C5	1:13:1085:U:C4	3.08	0.41
1:13:1375:A:OP1	7:6E:28:ASN:ND2	2.45	0.41
1:13:438:G:N1	1:13:495:A:OP2	2.37	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:56:U:H2'	1:13:57:G:H8	1.84	0.41
1:13:604:G:H2'	1:13:605:U:O4'	2.20	0.41
1:13:735:C:O2'	1:13:736:C:H5'	2.20	0.41
26:14:1054:A:N6	26:14:1104:C:H42	2.18	0.41
26:14:195:A:H61	26:14:198:C:H3'	1.85	0.41
26:14:208:C:H2'	26:14:209:C:C6	2.55	0.41
26:14:2290:G:H2'	26:14:2291:U:O4'	2.19	0.41
26:14:2706:G:N2	26:14:2707:G:H1'	2.35	0.41
26:14:2688:U:C5	26:14:2720:U:OP2	2.73	0.41
26:14:2778:A:C8	26:14:2778:A:H5''	2.53	0.41
26:14:2869:G:H2'	26:14:2870:C:O4'	2.20	0.41
26:14:2901:C:H2'	26:14:2902:C:H5'	2.02	0.41
26:14:492:A:C2	26:14:493:G:H1'	2.54	0.41
26:14:925:C:H2'	26:14:926:A:H8	1.85	0.41
26:14:979:G:C4	26:14:982:C:N4	2.88	0.41
2:1E:146:GLN:O	2:1E:150:SER:HB2	2.20	0.41
2:1E:97:TRP:CZ3	2:1E:172:ILE:HB	2.54	0.41
1:1G:102:G:C6	1:1G:103:C:C4	3.08	0.41
1:1G:171:A:C2	1:1G:172:A:C4	3.08	0.41
1:1G:390:C:O2'	16:7A:28:ARG:NH1	2.51	0.41
1:1G:626:U:C2	1:1G:627:G:C8	3.08	0.41
1:1G:920:U:H2'	1:1G:921:U:H6	1.82	0.41
26:1H:1166:C:H2'	26:1H:1167:U:H6	1.83	0.41
26:1H:1593:G:C2	26:1H:1594:G:C5	3.08	0.41
26:1H:1794:U:H2'	26:1H:1795:C:C6	2.54	0.41
26:1H:2287:A:C2	26:1H:2346:A:C2	3.08	0.41
26:1H:2330:G:H2'	26:1H:2331:G:O4'	2.19	0.41
26:1H:2563:U:O2'	35:68:28:SER:HB3	2.20	0.41
26:1H:71:A:OP1	26:1H:72:U:H2'	2.19	0.41
26:1H:881:G:H2'	26:1H:881:G:N3	2.35	0.41
26:1H:8:A:H2'	26:1H:9:U:O4'	2.20	0.41
3:22:120:VAL:HG21	3:22:137:ALA:HB2	2.00	0.41
1:1G:532:A:H5'	3:22:161:GLU:OE2	2.20	0.41
23:2L:47:C:HO2'	23:2L:48:C:P	2.43	0.41
4:32:174:LEU:HD23	4:32:174:LEU:HA	1.81	0.41
12:3A:57:LYS:HB2	12:3A:57:LYS:HE3	1.97	0.41
4:3E:161:ASN:O	4:3E:165:MET:HB2	2.20	0.41
31:41:118:ARG:HB2	31:41:181:ARG:NH1	2.35	0.41
31:41:122:PRO:HB3	31:41:180:PHE:HD2	1.84	0.41
31:41:57:ALA:HB2	31:41:90:LEU:HG	2.02	0.41
31:49:114:ILE:CG2	31:49:117:PHE:HB2	2.50	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:49:60:LEU:HD22	31:49:68:PRO:HB3	2.02	0.41
5:4E:152:ARG:H	5:4E:152:ARG:HG3	1.51	0.41
32:51:154:PRO:HD3	32:51:162:ILE:O	2.20	0.41
32:51:80:SER:C	32:51:81:GLU:HG3	2.39	0.41
32:59:151:ILE:O	32:59:152:ARG:HB2	2.19	0.41
6:5E:2:ARG:HB2	6:5E:4:TYR:CE1	2.55	0.41
6:5E:44:GLY:HA2	6:5E:59:TYR:CE2	2.55	0.41
8:72:36:LEU:HD23	8:72:36:LEU:HA	1.78	0.41
40:75:64:ARG:CB	40:75:73:GLU:HG2	2.49	0.41
36:78:59:LEU:O	54:Q8:13:ARG:HD2	2.20	0.41
9:82:38:GLN:O	9:82:38:GLN:NE2	2.53	0.41
17:8A:5:VAL:HG22	17:8A:60:ILE:HG12	2.02	0.41
17:8I:93:GLN:NE2	17:8I:96:GLU:OE1	2.35	0.41
42:95:44:LYS:O	42:95:46:VAL:N	2.50	0.41
18:9I:56:THR:OG1	18:9I:58:LEU:HD13	2.19	0.41
40:B8:114:LEU:HD23	40:B8:114:LEU:HA	1.62	0.41
20:BA:23:ARG:O	20:BA:27:LYS:HB2	2.19	0.41
20:BI:56:MET:CE	20:BI:85:MET:HG2	2.50	0.41
45:C5:81:LYS:HB2	45:C5:99:CYS:SG	2.60	0.41
45:C5:87:LYS:CG	45:C5:88:LYS:H	2.33	0.41
41:C8:106:PHE:O	41:C8:109:LEU:HB2	2.20	0.41
44:F8:34:ALA:HA	44:F8:38:GLU:OE1	2.20	0.41
26:1H:922:U:H1'	47:I8:26:TYR:CE2	2.55	0.41
49:K8:49:LYS:O	49:K8:53:LEU:HB2	2.20	0.41
53:L5:12:ARG:NH2	53:L5:44:PRO:HB3	2.34	0.41
31:41:112:PRO:CB	51:M8:37:SER:H	2.32	0.41
2:12:145:LEU:O	2:12:149:LEU:HB2	2.20	0.41
1:13:1103:C:H2'	1:13:1104:G:O4'	2.21	0.41
1:13:1226:C:H4'	1:13:1227:A:OP1	2.20	0.41
1:13:32:A:C2	1:13:33:A:C4	3.08	0.41
1:13:445:G:H2'	1:13:446:G:C8	2.56	0.41
26:14:1088:A:N3	26:14:1088:A:H2'	2.35	0.41
26:14:1299:G:H5'	26:14:1301:A:O4'	2.20	0.41
26:14:1396:U:H2'	26:14:1396:U:O2	2.19	0.41
26:14:2146:C:O2'	26:14:2147:G:OP2	2.35	0.41
26:14:2271:G:H2'	26:14:2272:U:C6	2.56	0.41
26:14:2772:C:H5'	29:29:168:MET:SD	2.60	0.41
26:14:2887:U:H2'	26:14:2888:C:H6	1.84	0.41
26:14:607:U:OP1	30:39:102:PRO:HA	2.20	0.41
34:15:127:ASP:OD1	34:15:127:ASP:N	2.52	0.41
1:1G:1286:A:P	21:1B:25:LYS:HE2	2.60	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1104:G:H4'	2:1E:111:ARG:NH1	2.35	0.41
1:1G:1227:A:H8	1:1G:1227:A:H3'	1.83	0.41
1:1G:1384:C:H2'	1:1G:1385:G:H8	1.85	0.41
1:1G:449:C:O4'	1:1G:449:C:O2	2.39	0.41
26:1H:1126:A:O5'	26:1H:1126:A:H8	2.03	0.41
26:1H:1266:G:OP2	52:N8:19:ARG:NH1	2.51	0.41
26:1H:1592:C:H2'	26:1H:1593:G:C8	2.53	0.41
26:1H:1899:G:O2'	26:1H:1900:A:P	2.78	0.41
26:1H:2636:U:H1'	26:1H:2783:G:N2	2.35	0.41
26:1H:2713:A:H3'	26:1H:2714:G:H5''	2.00	0.41
26:1H:2749:A:H1'	32:51:63:SER:OG	2.20	0.41
26:1H:295:G:C6	26:1H:344:G:C2	3.08	0.41
26:1H:608:A:C2	26:1H:621:A:C5	3.08	0.41
26:1H:937:U:H2'	26:1H:938:G:O4'	2.20	0.41
3:22:73:PRO:O	3:22:76:VAL:HG22	2.20	0.41
4:32:50:ARG:HG3	4:32:50:ARG:H	1.62	0.41
30:39:34:TRP:CE3	30:39:35:GLU:HG2	2.55	0.41
30:39:63:LYS:HB2	30:39:63:LYS:HE3	1.94	0.41
12:3A:82:VAL:O	12:3A:106:ASP:HB2	2.19	0.41
5:42:107:ARG:NH2	5:42:108:ALA:HB2	2.34	0.41
5:42:126:ARG:NH1	5:42:126:ARG:HG3	2.35	0.41
32:59:33:LEU:HD11	32:59:78:GLY:HA3	2.02	0.41
33:61:58:LEU:HD23	33:61:59:ALA:HB2	2.01	0.41
33:61:67:ARG:O	33:61:71:ILE:HG22	2.20	0.41
7:62:115:ARG:O	7:62:118:VAL:HG22	2.19	0.41
7:62:118:VAL:O	7:62:121:ALA:HB3	2.19	0.41
39:65:35:ILE:HB	39:65:97:ARG:NH2	2.35	0.41
39:65:80:LEU:HD23	39:65:80:LEU:HA	1.85	0.41
15:6A:25:THR:HG21	15:6A:70:LEU:HB2	2.02	0.41
1:13:878:G:H5'	8:7E:89:PRO:HG2	2.01	0.41
9:82:13:ALA:HB2	9:82:68:GLY:HA3	2.00	0.41
41:85:30:LYS:HD3	41:85:30:LYS:HA	1.76	0.41
41:85:60:LEU:O	41:85:63:VAL:HG23	2.20	0.41
41:85:62:ILE:HD11	41:85:93:LYS:HD3	2.02	0.41
37:88:74:TYR:O	37:88:90:VAL:O	2.38	0.41
18:9I:58:LEU:HD23	18:9I:63:GLN:HA	2.01	0.41
39:A8:27:SER:HA	39:A8:88:ASP:CB	2.50	0.41
40:B8:100:TYR:HB3	40:B8:103:ARG:NH1	2.34	0.41
46:D5:8:TYR:HA	46:D5:62:PRO:HD3	2.02	0.41
48:F5:91:LYS:HZ1	48:F5:95:LEU:CD1	2.33	0.41
46:H8:129:SER:HA	46:H8:130:PRO:HD3	1.76	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:I5:40:HIS:CE1	51:I5:45:GLY:HA3	2.55	0.41
2:12:211:ILE:O	2:12:215:LEU:HD23	2.21	0.41
1:13:243:A:H4'	1:13:244:U:H3'	2.01	0.41
1:13:895:G:H2'	1:13:896:C:C6	2.56	0.41
1:13:924:C:H2'	1:13:925:G:H8	1.84	0.41
26:14:2001:A:H2'	26:14:2002:G:O4'	2.21	0.41
26:14:2057:A:H2'	26:14:2058:A:C8	2.55	0.41
26:14:2233:U:H2'	26:14:2234:G:C8	2.55	0.41
26:14:270(U):C:C2	26:14:270(V):G:C8	3.08	0.41
26:14:951:C:O2'	26:14:952:G:H5'	2.20	0.41
21:1B:2:GLY:O	21:1B:5:ASP:N	2.33	0.41
2:1E:78:GLN:O	2:1E:81:VAL:HG12	2.19	0.41
1:1G:1317:C:H3'	1:1G:1318:A:C8	2.56	0.41
1:1G:588:G:H1	1:1G:651:C:N4	2.14	0.41
1:1G:646:U:H2'	1:1G:647:C:C6	2.55	0.41
1:1G:802:A:H3'	1:1G:803:G:C8	2.55	0.41
26:1H:1144:G:H2'	26:1H:1145:C:H6	1.84	0.41
26:1H:1187:G:H5''	42:D8:81:TYR:CE2	2.54	0.41
26:1H:1359:A:C2	26:1H:1372:U:N3	2.83	0.41
26:1H:1448:G:H1'	26:1H:1528:A:H62	1.85	0.41
26:1H:2270:G:OP2	57:1H:3952:HOH:O	2.22	0.41
26:1H:2314:C:H2'	26:1H:2315:G:C8	2.55	0.41
26:1H:2506:U:H2'	26:1H:2506:U:O2	2.19	0.41
26:1H:2564:A:H5''	26:1H:2565:A:OP2	2.20	0.41
3:22:109:PRO:HB2	3:22:115:LEU:CD1	2.50	0.41
3:22:21:ARG:HA	3:22:21:ARG:HE	1.84	0.41
3:22:65:ALA:HA	3:22:100:ALA:HB3	2.02	0.41
3:22:88:ARG:HG2	3:22:88:ARG:O	2.19	0.41
29:29:111:ARG:CG	29:29:111:ARG:HH11	2.27	0.41
29:29:36:ARG:HG2	29:29:47:VAL:HG22	2.03	0.41
3:2E:113:ALA:HB3	3:2E:114:PRO:HD3	2.02	0.41
3:2E:150:LYS:HD2	3:2E:201:TYR:CD2	2.56	0.41
23:2L:12:G:H1	23:2L:24:C:H42	1.68	0.41
26:1H:1257:C:H4'	30:31:83:PHE:CE1	2.56	0.41
4:32:126:ILE:HG22	4:32:127:THR:N	2.35	0.41
4:32:79:PHE:CE2	4:32:204:ILE:HD13	2.56	0.41
36:35:126:VAL:HG13	36:35:145:PRO:HG2	2.01	0.41
36:35:86:LYS:HG3	36:35:87:ASP:N	2.35	0.41
30:39:124:LEU:HD13	30:39:191:ARG:HH22	1.85	0.41
30:39:126:VAL:HG21	30:39:142:TRP:CH2	2.56	0.41
4:3E:30:LYS:HA	4:3E:35:ARG:CG	2.51	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:41:111:LEU:HD22	31:41:117:PHE:CE1	2.55	0.41
37:45:110:THR:HG23	37:45:113:GLN:OE1	2.20	0.41
31:49:166:ASP:OD1	31:49:166:ASP:N	2.51	0.41
5:4E:12:LEU:HB3	5:4E:31:LEU:HB2	2.01	0.41
5:4E:31:LEU:HD22	5:4E:43:LEU:HD11	2.02	0.41
13:4I:107:ALA:O	13:4I:109:THR:N	2.54	0.41
39:65:3:ARG:HH21	39:65:4:LEU:CB	2.25	0.41
33:69:110:ASP:HA	33:69:111:PRO:HD3	1.69	0.41
15:6A:26:GLU:OE2	15:6A:77:ARG:HB2	2.20	0.41
9:82:20:ARG:O	9:82:60:ASP:HB2	2.19	0.41
1:13:276:G:O2'	17:8I:68:ARG:NH1	2.53	0.41
19:AI:27:GLU:HG3	19:AI:27:GLU:H	1.74	0.41
45:G8:94:LYS:CA	45:G8:94:LYS:HZ3	2.27	0.41
26:1H:1159:U:P	50:L8:30:ARG:HH12	2.43	0.41
51:M8:13:ARG:HH12	51:M8:22:ILE:HG23	1.85	0.41
1:13:1315:U:H2'	1:13:1316:G:O4'	2.20	0.41
1:13:1316:G:H5''	14:5I:17:LYS:NZ	2.35	0.41
1:13:1347:G:H22	1:13:1374:A:P	2.43	0.41
1:13:452:A:H2'	1:13:453:A:C8	2.55	0.41
26:14:1022:G:HO2'	26:14:1023:U:P	2.35	0.41
26:14:1288:U:H4'	26:14:1289:C:OP2	2.20	0.41
26:14:1494:A:C2	26:14:1495:A:C4	3.08	0.41
26:14:1717:G:C6	26:14:1743:G:C6	3.08	0.41
26:14:2544:G:O5'	26:14:2544:G:H8	2.03	0.41
26:14:2755:C:HO2'	26:14:2756:U:H6	1.67	0.41
26:14:447:A:C6	26:14:454:A:C8	3.08	0.41
34:15:69:GLN:O	34:15:71:ILE:HG23	2.21	0.41
27:16:81:G:C6	27:16:82:G:C5	3.09	0.41
28:19:116:GLN:HB2	28:19:116:GLN:HE21	1.64	0.41
1:1G:236:G:H2'	1:1G:237:C:O4'	2.20	0.41
1:1G:450:G:N7	1:1G:481:G:C6	2.89	0.41
1:1G:748:C:H4'	1:1G:749:C:O5'	2.20	0.41
1:1G:750:G:H1'	15:6A:22:THR:OG1	2.20	0.41
1:1G:894:G:C6	1:1G:895:G:C5	3.09	0.41
26:1H:2271:G:C5'	47:I8:20:ARG:HG2	2.49	0.41
26:1H:2415:G:C6	26:1H:2416:C:C4	3.09	0.41
26:1H:2462:U:H2'	26:1H:2463:C:C6	2.55	0.41
26:1H:873:G:N2	26:1H:905:U:C2	2.88	0.41
26:1H:99:U:C6	26:1H:102:G:C2	3.07	0.41
27:1J:22:U:H5'	27:1J:23:G:OP2	2.20	0.41
4:32:118:ARG:O	4:32:122:ARG:HB2	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:39:110:LEU:HD12	30:39:205:ARG:HG2	2.02	0.41
12:3A:84:LEU:C	12:3A:85:ILE:HD12	2.41	0.41
4:3E:172:PRO:HB2	4:3E:187:ARG:HH12	1.86	0.41
31:41:105:LYS:HE3	31:41:105:LYS:HB2	1.81	0.41
5:42:79:GLU:OE1	8:72:104:ARG:HA	2.20	0.41
13:4I:12:ASN:HB3	13:4I:13:LYS:H	1.58	0.41
32:51:12:PRO:HB2	32:51:13:LYS:HE2	2.01	0.41
14:5I:58:LYS:HE3	14:5I:58:LYS:HB2	1.84	0.41
39:65:109:GLY:C	39:65:110:LEU:HD13	2.41	0.41
35:68:52:VAL:C	35:68:53:LYS:HD2	2.41	0.41
36:78:111:ARG:HG2	36:78:128:HIS:CD2	2.55	0.41
16:7I:38:TYR:CZ	16:7I:50:LYS:HB2	2.56	0.41
41:85:44:ASN:HD21	42:95:75:PHE:H	1.67	0.41
17:8I:100:LYS:HB2	17:8I:101:ARG:HE	1.85	0.41
19:AI:42:PRO:O	19:AI:45:VAL:HG22	2.20	0.41
1:13:1320:C:C2	19:AI:72:GLY:HA3	2.55	0.41
42:D8:89:GLN:HA	42:D8:90:PRO:HD2	1.81	0.41
46:H8:158:PRO:O	46:H8:161:VAL:HG22	2.20	0.41
47:I8:38:VAL:CG1	47:I8:40:GLN:HG2	2.46	0.41
52:N8:40:LYS:HG2	52:N8:46:CYS:SG	2.61	0.41
54:Q8:34:TRP:HE3	54:Q8:34:TRP:N	2.19	0.41
1:13:1002:G:C2	1:13:1003:G:H1'	2.56	0.41
1:13:1304:G:C6	1:13:1305:G:C6	3.09	0.41
1:13:123:C:OP1	1:13:312:C:H5'	2.21	0.41
1:13:453:A:C6	1:13:454:C:C4	3.09	0.41
1:13:802:A:H3'	1:13:803:G:C8	2.56	0.41
26:14:1019:U:OP1	26:14:1035:U:O2'	2.25	0.41
26:14:1225:C:C3'	42:95:85:LYS:HA	2.51	0.41
26:14:176:G:O2'	26:14:177:G:H5'	2.20	0.41
26:14:1856:G:H1	26:14:1886:C:N4	2.02	0.41
26:14:1856:G:C6	26:14:1857:G:C5	3.08	0.41
26:14:2123:G:N2	26:14:2176:A:C6	2.88	0.41
26:14:198:C:H4'	26:14:2243:U:O2'	2.21	0.41
2:1E:166:ASP:HB3	2:1E:169:LYS:HB2	2.01	0.41
1:1G:1300:G:O2'	1:1G:1301:U:P	2.78	0.41
1:1G:1374:A:H2'	1:1G:1375:A:H5'	2.02	0.41
26:1H:1286:A:C5	26:1H:1289:C:C4	3.09	0.41
26:1H:2121:G:H2'	26:1H:2122:U:C6	2.56	0.41
26:1H:2338:G:H2'	26:1H:2339:G:H8	1.85	0.41
26:1H:242:G:C8	54:Q8:3:LYS:HG3	2.56	0.41
26:1H:2795:G:C4	26:1H:2802:G:O6	2.74	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:565:C:H4'	57:1H:4120:HOH:O	2.19	0.41
26:1H:686:G:H4'	26:1H:686:G:OP2	2.21	0.41
26:1H:821:A:O2'	26:1H:945:A:H3'	2.20	0.41
27:1J:99:A:C4	27:1J:100:G:C8	3.08	0.41
29:21:14:ILE:HA	29:21:14:ILE:HD13	1.80	0.41
29:29:55:ASN:C	29:29:57:LYS:H	2.23	0.41
29:29:59:VAL:CB	29:29:60:ASN:HA	2.45	0.41
29:29:81:ILE:HG22	29:29:82:ARG:N	2.35	0.41
11:2A:79:SER:OG	11:2A:106:LYS:HD2	2.20	0.41
30:39:46:ARG:O	30:39:48:THR:HG23	2.20	0.41
30:39:60:SER:OG	30:39:61:GLY:N	2.53	0.41
31:49:9:ARG:HG2	31:49:13:GLU:OE2	2.20	0.41
3:2E:29:TYR:OH	14:5I:54:PRO:O	2.27	0.41
39:65:83:LYS:HE2	39:65:83:LYS:HB3	1.77	0.41
35:68:8:LEU:HD12	35:68:82:ASN:HB2	2.01	0.41
35:68:2:ILE:HD11	35:68:82:ASN:HB3	2.02	0.41
1:1G:750:G:N3	15:6A:23:GLY:HA3	2.35	0.41
40:75:26:ASP:OD1	40:75:120:ARG:NH2	2.54	0.41
40:75:4:GLY:O	40:75:5:ALA:C	2.58	0.41
8:7E:107:LEU:HD23	8:7E:107:LEU:HA	1.83	0.41
8:7E:87:SER:HB2	8:7E:93:VAL:H	1.86	0.41
37:88:87:LYS:HG2	37:88:88:GLY:N	2.34	0.41
9:8E:118:LYS:O	9:8E:119:ALA:HB3	2.20	0.41
38:98:21:TYR:OH	38:98:43:GLU:HG2	2.20	0.41
39:A8:25:ARG:O	39:A8:39:ILE:HA	2.21	0.41
44:B5:31:HIS:HA	44:B5:32:PRO:HD3	1.78	0.41
29:21:15:PHE:CD2	40:B8:81:PRO:HD3	2.56	0.41
41:C8:72:HIS:ND1	41:C8:110:VAL:HG21	2.35	0.41
46:D5:161:VAL:HB	46:D5:162:GLU:H	1.50	0.41
46:D5:17:ALA:HA	46:D5:20:ARG:HD2	2.02	0.41
46:H8:128:VAL:CG2	46:H8:161:VAL:HG12	2.50	0.41
46:H8:60:GLU:O	46:H8:61:LEU:HD23	2.21	0.41
50:L8:2:PRO:HB2	50:L8:3:ARG:H	1.59	0.41
50:L8:55:ARG:O	50:L8:55:ARG:HG3	2.20	0.41
28:11:85:ASP:OD1	28:11:86:PRO:HD2	2.20	0.41
2:12:20:GLU:H	2:12:20:GLU:HG2	1.72	0.41
2:12:5:ILE:HG12	2:12:5:ILE:O	2.21	0.41
1:13:1014:A:C2	1:13:1219:U:H1'	2.56	0.41
1:13:1357:A:C5	1:13:1358:U:C4	3.08	0.41
1:13:148:G:N2	1:13:149:A:C5	2.89	0.41
1:13:344:A:O2'	1:13:346:G:N7	2.53	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:537:G:H2'	1:13:538:G:C8	2.55	0.41
1:13:626:U:N3	1:13:627:G:N7	2.68	0.41
1:13:659:U:C2	1:13:660:G:C8	3.08	0.41
1:13:711:G:H2'	1:13:712:A:C8	2.56	0.41
1:13:97:U:H2'	1:13:99:C:C6	2.55	0.41
26:14:1429:G:C5	26:14:1568:G:C6	3.09	0.41
26:14:1574:C:H6	26:14:1574:C:O5'	2.03	0.41
26:14:1728:G:C6	26:14:1730:U:OP2	2.74	0.41
26:14:2137:C:OP2	26:14:2137:C:H6	2.03	0.41
26:14:2433:A:H5''	26:14:2434:A:P	2.61	0.41
26:14:327:G:C6	26:14:328:U:N3	2.89	0.41
26:14:634:C:H2'	26:14:635:C:H6	1.81	0.41
1:1G:1009:G:C2	1:1G:1010:G:C8	3.09	0.41
1:1G:423:G:N2	1:1G:424:G:C4	2.88	0.41
1:1G:526:C:C5	1:1G:527:G:H1'	2.55	0.41
26:1H:1069:A:H4'	26:1H:1070:A:H5''	2.02	0.41
26:1H:1108:U:O4	26:1H:1109:C:N4	2.53	0.41
26:1H:1528:A:C2	26:1H:1542:G:C2	3.08	0.41
26:1H:2590:A:H2'	26:1H:2591:C:C6	2.55	0.41
26:1H:2655:G:N2	26:1H:2665:A:OP2	2.47	0.41
26:1H:552:G:H2'	26:1H:553:U:O4'	2.21	0.41
26:1H:775:G:C2	26:1H:777:A:N6	2.88	0.41
26:1H:800:A:H4'	26:1H:801:G:O5'	2.19	0.41
10:1I:38:ILE:HG23	10:1I:71:LEU:HB3	2.01	0.41
29:21:147:PRO:HB2	29:21:149:ARG:CG	2.51	0.41
3:22:15:THR:HG21	3:22:181:ASN:HA	2.01	0.41
3:2E:8:ILE:HG23	3:2E:16:ARG:HG2	2.03	0.41
23:2L:61:C:H6	23:2L:61:C:O5'	2.04	0.41
30:39:122:LYS:HB3	30:39:191:ARG:HB2	2.02	0.41
12:3A:12:ARG:HE	12:3A:12:ARG:HB3	1.57	0.41
12:3I:123:LYS:HG2	12:3I:123:LYS:H	1.73	0.41
12:3I:51:ALA:C	12:3I:52:LEU:HD23	2.41	0.41
31:41:99:MET:HG3	31:41:100:TRP:N	2.35	0.41
5:42:131:ILE:O	5:42:134:ALA:HB3	2.20	0.41
31:49:6:ALA:N	51:I5:23:GLU:OE2	2.53	0.41
13:4A:81:LEU:HB3	13:4A:89:GLY:CA	2.50	0.41
5:4E:106:PRO:O	5:4E:110:LEU:HG	2.21	0.41
5:4E:45:PHE:CE2	5:4E:47:LYS:HD2	2.55	0.41
6:5E:41:GLU:O	6:5E:43:LEU:HD12	2.20	0.41
14:5I:39:LEU:HD13	14:5I:47:LEU:HD12	2.02	0.41
7:62:24:THR:O	7:62:27:ILE:HG12	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:6I:35:ARG:HA	15:6I:38:ARG:HB2	2.01	0.41
40:75:19:LEU:HA	40:75:20:PRO:HD3	1.84	0.41
26:1H:598:G:C1'	36:78:12:ALA:HB2	2.50	0.41
1:13:377:G:OP1	16:7I:3:LYS:HD2	2.19	0.41
17:8A:88:TYR:CE1	17:8A:92:ARG:HD3	2.55	0.41
38:98:67:LEU:HD13	38:98:67:LEU:HA	1.75	0.41
43:A5:29:LEU:O	43:A5:29:LEU:HD12	2.20	0.41
20:BA:41:ILE:HD12	20:BA:87:LYS:HE3	2.02	0.41
20:BI:26:ASN:HB2	20:BI:71:THR:HG23	2.03	0.41
20:BI:53:LEU:HD23	20:BI:100:ILE:HG22	2.02	0.41
41:C8:11:ARG:O	41:C8:15:LYS:HG3	2.21	0.41
48:F5:80:LEU:HB3	48:F5:82:LEU:HG	2.02	0.41
38:98:33:ARG:HH12	52:N8:55:ARG:HB3	1.84	0.41
26:1H:2054:A:H2'	52:N8:8:LYS:HE2	2.03	0.41
2:12:188:ALA:O	2:12:203:GLY:N	2.54	0.41
1:13:1011:G:H2'	1:13:1012:U:O4'	2.20	0.41
1:13:1004:A:H1'	1:13:1036:G:N2	2.35	0.41
1:13:1062:U:H2'	1:13:1063:C:C5	2.56	0.41
26:14:117:G:C6	26:14:119:A:C6	3.09	0.41
26:14:125:G:H4'	26:14:126:A:OP2	2.21	0.41
26:14:1421:G:C2	26:14:1422:G:N7	2.88	0.41
26:14:1574:C:H2'	26:14:1575:C:C6	2.55	0.41
26:14:2635:C:H5'	29:29:77:ILE:O	2.20	0.41
26:14:2681:C:H2'	26:14:2681:C:O2	2.21	0.41
26:14:537:C:H5'	26:14:539:G:OP2	2.21	0.41
26:14:654(S):G:C2	26:14:654(T):A:C6	3.08	0.41
26:14:921:G:C5	26:14:922:U:C5	3.09	0.41
26:14:957:A:H4'	37:45:74:TYR:OH	2.21	0.41
34:15:87:LEU:HD23	34:15:87:LEU:HA	1.85	0.41
28:19:245:PRO:HA	28:19:246:PRO:HD3	1.83	0.41
1:1G:134:A:H1'	1:1G:325:A:C5	2.56	0.41
1:1G:38:G:C2	1:1G:397:A:C2	3.09	0.41
1:1G:407:G:OP1	4:32:115:ARG:NH2	2.48	0.41
1:1G:42:G:H2'	1:1G:43:C:O4'	2.19	0.41
1:1G:731:G:O2'	1:1G:732:C:H5'	2.21	0.41
1:1G:825:G:H1	1:1G:875:C:H42	1.68	0.41
1:1G:827:U:H3	1:1G:872:A:N6	2.11	0.41
1:1G:868:C:H2'	1:1G:869:G:O4'	2.20	0.41
1:1G:928:G:C2	1:1G:1390:U:O2	2.73	0.41
26:1H:1919:A:H5''	26:1H:1920:C:OP2	2.21	0.41
26:1H:1927:A:C6	26:1H:1928:A:C6	3.08	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2071:A:H2'	26:1H:2072:G:C8	2.56	0.41
26:1H:2287:A:N3	26:1H:2289:G:C8	2.89	0.41
26:1H:2315:G:H5''	26:1H:2316:C:OP2	2.20	0.41
26:1H:1954:G:N3	26:1H:2551:C:H5''	2.35	0.41
26:1H:1783:A:C6	26:1H:2587:A:C2	3.09	0.41
26:1H:25:U:C4	26:1H:26:G:C6	3.09	0.41
26:1H:2689:U:P	26:1H:2719:G:H22	2.43	0.41
26:1H:32:C:O2'	26:1H:33:U:H5'	2.20	0.41
26:1H:57:C:H2'	26:1H:58:G:O4'	2.20	0.41
26:1H:5:A:H2'	26:1H:6:A:C8	2.55	0.41
3:22:95:THR:C	3:22:97:LYS:H	2.24	0.41
3:2E:156:ARG:HE	3:2E:156:ARG:HB3	1.48	0.41
4:32:57:ARG:HH22	5:42:107:ARG:HD3	1.85	0.41
30:39:68:LYS:HA	30:39:68:LYS:HD2	1.97	0.41
24:3K:21:A:H62	24:3K:46:G:H1'	1.86	0.41
24:3K:21:A:N7	24:3K:58:U:H1'	2.35	0.41
24:3K:63:G:O6	24:3K:72:C:N4	2.54	0.41
31:41:26:GLN:HB2	31:41:26:GLN:HE21	1.74	0.41
31:41:44:GLY:O	31:41:47:LYS:HE3	2.20	0.41
31:41:60:LEU:HD23	31:41:60:LEU:HA	1.82	0.41
31:41:61:ALA:HB2	31:41:67:LYS:HA	2.03	0.41
26:1H:2312:U:OP2	31:41:74:LYS:HD3	2.21	0.41
13:4A:3:ARG:NH1	13:4A:7:VAL:HG12	2.35	0.41
25:4L:33:G:H2'	25:4L:34:G:C8	2.55	0.41
6:52:99:ALA:HB1	18:9A:62:GLU:OE2	2.20	0.41
14:5I:37:PHE:CE1	14:5I:53:LEU:HD13	2.55	0.41
33:61:113:ARG:H	33:61:116:LEU:HD13	1.85	0.41
7:62:120:ILE:O	7:62:124:LEU:HB2	2.21	0.41
33:69:4:ILE:HG12	33:69:4:ILE:H	1.55	0.41
40:75:15:VAL:HG23	40:75:79:HIS:CE1	2.55	0.41
41:85:114:LYS:HA	41:85:117:GLN:HG3	2.03	0.41
9:8E:20:ARG:HA	9:8E:21:PRO:HD3	1.91	0.41
38:98:65:LEU:HA	38:98:65:LEU:HD12	1.71	0.41
43:A5:73:ALA:O	43:A5:106:ILE:HG12	2.21	0.41
19:AA:80:TYR:CE2	19:AA:82:GLY:HA2	2.56	0.41
40:B8:22:PHE:N	40:B8:22:PHE:CD1	2.89	0.41
40:B8:33:LYS:HG3	40:B8:82:LEU:HA	2.01	0.41
20:BA:10:LEU:HD13	20:BA:10:LEU:O	2.21	0.41
20:BA:50:GLU:HA	20:BA:100:ILE:CG2	2.50	0.41
45:C5:36:ALA:O	45:C5:37:VAL:HG13	2.21	0.41
27:16:12:C:O2'	47:I8:74:ARG:HG2	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:M5:60:LEU:HB2	54:M5:61:LEU:H	1.34	0.41
26:1H:246:C:N4	54:Q8:8:LYS:HG2	2.35	0.41
28:11:182:LEU:HB3	28:11:271:ILE:CG1	2.51	0.41
2:12:23:ARG:NH2	2:12:191:ASP:HB2	2.35	0.41
2:12:97:TRP:CZ3	2:12:99:GLY:HA2	2.55	0.41
1:13:1121:U:C4	1:13:1122:U:C4	3.09	0.41
1:13:1235:U:H2'	1:13:1236:A:O4'	2.21	0.41
1:13:257:G:C4	1:13:258:G:C8	3.09	0.41
1:13:382:A:H2'	1:13:383:A:C8	2.55	0.41
1:13:626:U:H2'	1:13:627:G:C8	2.55	0.41
1:13:963:G:H21	10:1I:55:LYS:NZ	2.18	0.41
26:14:70:G:H5''	26:14:112:U:O2	2.20	0.41
26:14:1356:G:H2'	26:14:1357:U:O4'	2.19	0.41
26:14:152:G:H2'	26:14:153:C:O4'	2.21	0.41
26:14:1787:A:N3	26:14:1787:A:H2'	2.34	0.41
26:14:638:G:C5	26:14:651:G:C2	3.09	0.41
26:14:882:G:O2'	26:14:883:G:O4'	2.36	0.41
1:1G:1002:G:H2'	1:1G:1003:G:H8	1.85	0.41
1:1G:1124:G:OP2	1:1G:1124:G:H8	2.03	0.41
1:1G:1200:C:H5'	1:1G:1201:A:H5''	2.02	0.41
1:1G:1329:A:H5''	13:4A:25:ILE:O	2.21	0.41
1:1G:1385:G:C2	1:1G:1386:G:C8	3.09	0.41
1:1G:279:A:OP2	17:8A:95:TYR:OH	2.26	0.41
1:1G:575:G:OP1	1:1G:575:G:H4'	2.21	0.41
1:1G:625:G:C5	1:1G:626:U:C5	3.08	0.41
26:1H:1055:G:H1'	26:1H:1085:A:C2	2.55	0.41
26:1H:2000:G:C2'	26:1H:2001:A:H5'	2.51	0.41
26:1H:2342:C:O2	26:1H:2374:C:H4'	2.20	0.41
26:1H:1864:U:OP1	26:1H:2411:A:H5'	2.20	0.41
26:1H:1956:U:H1'	26:1H:2552:U:OP1	2.21	0.41
26:1H:265:A:H8	26:1H:266:G:H1'	1.86	0.41
26:1H:2712:U:OP1	26:1H:2714:G:H4'	2.21	0.41
26:1H:273(F):C:H3'	26:1H:274:G:C5'	2.50	0.41
26:1H:439:G:H2'	26:1H:440:G:H8	1.85	0.41
26:1H:447:A:C8	26:1H:473:G:C6	3.09	0.41
26:1H:608:A:C4	26:1H:621:A:C6	3.09	0.41
26:1H:706:A:H2'	26:1H:707:G:O4'	2.20	0.41
26:1H:751:A:N7	57:1H:3741:HOH:O	2.53	0.41
26:1H:956:G:N2	26:1H:959:A:H3'	2.35	0.41
10:1I:22:LYS:HZ1	10:1I:90:LEU:HB2	1.84	0.41
27:1J:56:G:H8	27:1J:56:G:O5'	2.04	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:1L:10:G:H2'	22:1L:11:U:H6	1.86	0.41
22:1L:47:C:N4	22:1L:56:G:H22	2.19	0.41
35:25:103:ALA:O	35:25:106:LEU:HB2	2.20	0.41
29:29:11:MET:HG3	29:29:24:THR:N	2.36	0.41
29:29:11:MET:CE	29:29:187:ALA:H	2.34	0.41
29:29:76:ARG:CG	29:29:195:LEU:HD22	2.50	0.41
11:2A:66:LEU:HD23	11:2A:66:LEU:HA	1.84	0.41
3:2E:204:LEU:HA	3:2E:204:LEU:HD23	1.85	0.41
11:2I:73:MET:SD	11:2I:103:LEU:HD22	2.60	0.41
23:2K:38:1MG:HN21	23:2K:38:1MG:HM11	1.75	0.41
26:1H:674:G:O2'	30:31:74:ARG:HG3	2.21	0.41
4:32:134:ASP:O	4:32:136:PRO:HD3	2.21	0.41
36:35:61:ARG:HG2	36:35:61:ARG:H	1.54	0.41
30:39:64:ILE:HG12	30:39:64:ILE:H	1.53	0.41
4:3E:108:LEU:CD1	4:3E:174:LEU:HD13	2.51	0.41
13:4A:14:ARG:HG2	13:4A:44:ARG:HE	1.85	0.41
13:4A:15:VAL:O	13:4A:19:LEU:HG	2.21	0.41
32:51:87:LEU:HA	32:51:163:TYR:O	2.20	0.41
32:59:159:GLU:HB3	32:59:160:LYS:H	1.57	0.41
33:61:104:GLN:HG2	33:61:105:HIS:ND1	2.35	0.41
39:65:26:LEU:HB3	39:65:87:PHE:HA	2.01	0.41
36:78:38:GLN:O	36:78:41:ARG:HB2	2.21	0.41
26:1H:2406:U:C2	36:78:75:ILE:HD13	2.56	0.41
37:88:59:ARG:C	37:88:61:GLY:N	2.74	0.41
42:95:41:GLY:HA3	42:95:46:VAL:HG11	2.03	0.41
42:95:67:GLY:O	42:95:88:ARG:HD2	2.21	0.41
38:98:13:HIS:CE1	38:98:16:HIS:HB2	2.55	0.41
44:B5:21:PHE:HD1	44:B5:21:PHE:N	2.18	0.41
44:B5:56:THR:HB	44:B5:77:LYS:HE2	2.03	0.41
45:C5:19:LYS:O	45:C5:21:LYS:N	2.54	0.41
46:D5:30:ASN:ND2	46:D5:90:VAL:HB	2.33	0.41
48:F5:86:SER:N	48:F5:87:PRO:CD	2.84	0.41
53:L5:32:LYS:HB3	53:L5:32:LYS:HE3	1.79	0.41
31:41:61:ALA:O	51:M8:7:PRO:HG3	2.20	0.41
1:13:1171:G:H2'	1:13:1172:C:H6	1.86	0.41
1:13:1263:C:H2'	1:13:1264:C:H6	1.85	0.41
1:13:186:C:H6	1:13:186:C:H3'	1.86	0.41
1:13:276:G:C6	1:13:277:C:C4	3.08	0.41
1:13:391:G:O3'	16:7I:8:ARG:NH2	2.54	0.41
1:13:491:G:H2'	1:13:492:G:C8	2.55	0.41
1:13:495:A:C2	1:13:496:A:C5	3.09	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:626:U:H2'	1:13:627:G:H8	1.85	0.41
1:13:807:A:H2'	1:13:808:C:C6	2.55	0.41
1:13:892:A:C6	1:13:893:C:C4	3.09	0.41
26:14:1061:U:H5''	26:14:1062:G:OP2	2.21	0.41
26:14:51:G:N3	26:14:119:A:C2	2.89	0.41
26:14:1426:G:H5''	26:14:1559:G:N1	2.36	0.41
26:14:2111:C:N4	26:14:2118:U:H1'	2.36	0.41
26:14:2273:A:H2'	26:14:2274:A:C8	2.55	0.41
26:14:2360:A:H2'	26:14:2361:A:O4'	2.21	0.41
26:14:2381:C:H6	26:14:2381:C:O5'	2.03	0.41
26:14:2499:C:OP2	57:14:3528:HOH:O	2.21	0.41
26:14:2651:C:O2'	26:14:2652:C:H5'	2.20	0.41
26:14:2784:C:H2'	26:14:2785:C:C6	2.56	0.41
1:1G:1137:C:H5''	1:1G:1138:G:OP1	2.21	0.41
1:1G:1189:C:H5''	1:1G:1190:G:OP2	2.21	0.41
1:1G:1504:G:H4'	1:1G:1505:G:O4'	2.21	0.41
1:1G:613:C:H6	1:1G:613:C:H3'	1.85	0.41
26:1H:1249:U:P	57:1H:3732:HOH:O	2.78	0.41
26:1H:1519:G:H2'	26:1H:1520:U:O4'	2.20	0.41
26:1H:1992:G:C2	26:1H:1997:G:C5	3.09	0.41
26:1H:2306:C:H2'	26:1H:2307:G:H21	1.85	0.41
26:1H:2442:C:H2'	26:1H:2443:C:C6	2.56	0.41
26:1H:2626:C:H2'	26:1H:2627:G:O4'	2.21	0.41
26:1H:2688:U:H5	26:1H:2720:U:OP2	2.04	0.41
26:1H:2721:A:H1'	26:1H:2873:A:O2'	2.20	0.41
26:1H:46:C:OP2	26:1H:215:G:H2'	2.20	0.41
26:1H:612:G:C6	26:1H:613:U:C4	3.09	0.41
26:1H:817:C:H5''	57:1H:3718:HOH:O	2.21	0.41
10:1I:6:ILE:O	10:1I:71:LEU:HD12	2.21	0.41
29:21:116:VAL:HG21	29:21:122:PHE:CE2	2.56	0.41
11:2I:78:GLN:O	11:2I:103:LEU:HA	2.21	0.41
23:2L:73:C:H2'	23:2L:74:C:C5	2.55	0.41
30:39:15:SER:OG	30:39:17:ARG:N	2.52	0.41
30:39:51:THR:HG23	30:39:92:PRO:HG2	2.03	0.41
4:3E:39:PRO:HA	4:3E:40:PRO:HD3	1.74	0.41
12:3I:84:LEU:HB2	12:3I:105:TYR:CE2	2.56	0.41
1:13:10:A:OP2	5:4E:126:ARG:HD3	2.20	0.41
13:4I:39:ILE:CD1	13:4I:56:LEU:HD23	2.46	0.41
32:51:87:LEU:HB2	32:51:131:VAL:HG12	2.01	0.41
32:59:6:ARG:HB2	32:59:66:GLY:CA	2.34	0.41
6:5E:86:ARG:O	6:5E:87:ARG:HG2	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:61:9:LEU:HD21	33:61:35:LEU:CD1	2.51	0.41
7:62:70:LYS:HD3	7:62:96:GLN:HB3	2.02	0.41
7:6E:18:TYR:HB3	7:6E:59:LEU:HD12	2.01	0.41
15:6I:18:PHE:CZ	15:6I:21:ASP:HB2	2.55	0.41
26:14:2020:A:P	41:85:27:LEU:HD23	2.61	0.41
37:88:125:LEU:HA	37:88:125:LEU:HD23	1.80	0.41
37:88:55:VAL:HG12	37:88:64:ILE:HD12	2.02	0.41
37:88:90:VAL:HG23	37:88:91:GLU:H	1.86	0.41
17:8A:19:VAL:HG23	17:8A:44:ALA:HB3	2.03	0.41
9:8E:53:VAL:HG22	9:8E:92:TYR:CE1	2.56	0.41
9:8E:18:PHE:CD2	9:8E:62:TYR:HD2	2.38	0.41
42:95:51:VAL:HG12	42:95:52:VAL:N	2.36	0.41
42:95:78:LYS:O	42:95:79:VAL:HG13	2.19	0.41
43:A5:64:MET:HE2	43:A5:64:MET:HB3	1.84	0.41
26:1H:2318:G:H22	39:A8:3:ARG:H	1.66	0.41
44:B5:60:ARG:NH1	44:B5:60:ARG:HA	2.33	0.41
1:13:322:C:O2'	20:BI:23:ARG:HD2	2.20	0.41
46:H8:133:ILE:O	46:H8:135:GLU:N	2.54	0.41
1:13:1112:C:H6	1:13:1112:C:O5'	2.04	0.41
1:13:1357:A:O5'	1:13:1359:C:N4	2.54	0.41
1:13:423:G:N2	1:13:424:G:C8	2.89	0.41
1:13:52:G:H2'	1:13:53:A:H8	1.86	0.41
26:14:1343:G:C2'	26:14:1344:G:H5'	2.51	0.41
26:14:146:G:H2'	26:14:147:U:O4'	2.21	0.41
26:14:2305:A:C2	31:49:154:GLY:HA3	2.55	0.41
26:14:2388:A:C2'	26:14:2389:G:H5'	2.51	0.41
26:14:2469:A:OP1	26:14:2469:A:H4'	2.20	0.41
26:14:2659:G:O2'	26:14:2661:G:N7	2.46	0.41
26:14:2712:U:HO2'	26:14:2712(A):A:P	2.44	0.41
26:14:2720:U:H2'	26:14:2720:U:O2	2.20	0.41
26:14:373:U:H2'	26:14:374:A:C8	2.56	0.41
26:14:890:A:O2'	26:14:892:G:H5'	2.21	0.41
28:19:222:ARG:CZ	57:19:304:HOH:O	2.68	0.41
1:1G:1140:C:H2'	1:1G:1141:C:C6	2.56	0.41
1:1G:449:C:H6	16:7A:42:ARG:HD2	1.86	0.41
1:1G:559:A:H4'	1:1G:560:U:H5''	2.03	0.41
1:1G:956:U:H2'	1:1G:957:U:O4'	2.21	0.41
26:1H:1091:G:C2	26:1H:1101:U:H1'	2.56	0.41
26:1H:1653:G:H3'	38:98:2:ARG:HG3	2.02	0.41
26:1H:2131:G:C5'	26:1H:2133:G:H4'	2.50	0.41
26:1H:2591:C:H2'	26:1H:2592:G:C8	2.56	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:270(V):G:H2'	26:1H:270(W):G:O4'	2.21	0.41
22:1K:50:A:H5'	22:1K:51:A:H5''	2.03	0.41
22:1K:5:A:H2'	22:1K:6:G:O4'	2.21	0.41
26:14:1665:A:C4'	35:25:67:LYS:HB2	2.51	0.41
11:2A:29:ILE:CG2	11:2A:44:SER:HB2	2.44	0.41
7:6E:150:ALA:HB2	11:2I:50:TYR:HE2	1.86	0.41
23:2L:76:U:H2'	23:2L:77:C:H6	1.86	0.41
30:31:129:PHE:O	30:31:130:ALA:HB3	2.21	0.41
4:32:13:ARG:C	4:32:15:GLU:N	2.74	0.41
36:35:55:ARG:HG2	36:35:56:SER:H	1.86	0.41
24:3K:43:A:H2'	24:3K:43:A:N3	2.36	0.41
31:41:64:THR:HG22	31:41:66:GLN:N	2.25	0.41
32:51:12:PRO:HB2	32:51:13:LYS:H	1.70	0.41
32:59:91:GLY:HA3	32:59:160:LYS:HB3	2.03	0.41
33:69:44:LEU:HA	33:69:44:LEU:HD22	1.83	0.41
36:78:96:THR:HG23	36:78:126:VAL:HG21	2.03	0.41
16:7I:53:VAL:CG2	16:7I:77:ALA:HB1	2.50	0.41
37:88:35:VAL:HG13	37:88:130:LYS:O	2.21	0.41
1:13:1349:A:OP1	9:8E:118:LYS:HB2	2.20	0.41
1:13:719:C:O2'	18:9I:49:LYS:HB3	2.21	0.41
43:A5:84:ARG:O	43:A5:96:ILE:HD13	2.21	0.41
39:A8:67:ARG:O	39:A8:71:ARG:HG2	2.21	0.41
26:1H:2376:A:N1	39:A8:87:PHE:HB3	2.35	0.41
40:B8:19:LEU:HA	40:B8:20:PRO:HD3	1.88	0.41
46:D5:44:PHE:CE1	46:D5:48:PHE:HB2	2.55	0.41
42:D8:21:ARG:HG2	42:D8:91:TYR:CE2	2.56	0.41
45:G8:99:CYS:N	45:G8:100:ALA:HA	2.35	0.41
50:H5:11:SER:HA	50:H5:12:PRO:HD2	1.67	0.41
50:L8:13:ILE:HA	50:L8:13:ILE:HD13	1.63	0.41
54:M5:59:LYS:HG2	54:M5:59:LYS:HZ3	1.78	0.41
28:11:254:THR:O	28:11:254:THR:OG1	2.34	0.41
1:13:52:G:H2'	1:13:53:A:O4'	2.21	0.41
1:13:585:G:C6	1:13:586:C:C4	3.09	0.41
26:14:1249:U:H2'	36:35:18:ARG:HH12	1.86	0.41
26:14:1399:C:O2'	26:14:1400:G:H5'	2.20	0.41
26:14:275:G:H5''	26:14:363:G:O6	2.21	0.41
26:14:693:C:H2'	26:14:694:U:H6	1.86	0.41
28:19:111:LEU:HA	28:19:111:LEU:HD23	1.84	0.41
28:19:244:ARG:HB2	28:19:245:PRO:HD2	2.02	0.41
2:1E:21:ARG:O	2:1E:23:ARG:N	2.54	0.41
2:1E:47:THR:O	2:1E:51:LEU:N	2.34	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1055:A:N6	1:1G:1206:G:C5	2.89	0.41
1:1G:1326:C:H2'	1:1G:1327:C:C6	2.56	0.41
1:1G:468:A:H4'	16:7A:80:PHE:O	2.21	0.41
1:1G:605:U:H3	1:1G:633:G:H1	1.68	0.41
26:1H:1028:A:N3	26:1H:2486:G:O2'	2.45	0.41
26:1H:176:G:C2'	26:1H:177:G:H5'	2.51	0.41
26:1H:249:C:O2	54:Q8:12:LYS:HE3	2.21	0.41
26:1H:2854:G:C6	26:1H:2864:G:N1	2.89	0.41
26:1H:934:G:C4	26:1H:935:C:C5	3.09	0.41
26:1H:946:G:OP1	57:1H:3866:HOH:O	2.21	0.41
22:1L:31:C:H2'	22:1L:32:C:C6	2.55	0.41
22:1L:49:C:C5	22:1L:50:A:H1'	2.56	0.41
26:1H:2053:G:H5'	29:21:144:ARG:O	2.21	0.41
11:2A:12:ARG:HG2	11:2A:13:GLN:N	2.36	0.41
4:32:91:SER:OG	4:32:191:ARG:HG3	2.20	0.41
36:35:47:ASP:HB3	36:35:48:PRO:CA	2.51	0.41
4:3E:13:ARG:HD2	4:3E:38:TYR:O	2.21	0.41
31:41:117:PHE:HE1	31:41:120:LEU:HD23	1.85	0.41
5:42:143:ARG:HA	5:42:143:ARG:HD3	1.93	0.41
13:4A:50:GLU:O	13:4A:54:VAL:HG23	2.21	0.41
6:52:72:VAL:O	6:52:75:LEU:HB3	2.21	0.41
38:55:18:LEU:HD11	38:55:22:ARG:CZ	2.51	0.41
34:58:13:TRP:O	34:58:134:ARG:HA	2.20	0.41
39:65:61:ASN:CG	39:65:62:LYS:H	2.24	0.41
8:72:100:ILE:HD11	8:72:112:LEU:HD11	2.02	0.41
36:78:85:LEU:HD13	36:78:120:ALA:HB2	2.03	0.41
16:7A:40:ASP:HA	16:7A:41:PRO:HD2	1.81	0.41
9:8E:127:LYS:O	9:8E:127:LYS:HG2	2.21	0.41
18:9A:76:LEU:HA	18:9A:76:LEU:HD23	1.68	0.41
18:9I:26:LEU:HB3	18:9I:42:ARG:NH2	2.36	0.41
43:A5:29:LEU:HD21	43:A5:33:ARG:NH2	2.36	0.41
44:B5:12:VAL:HG22	44:B5:17:ALA:HB2	2.03	0.41
40:B8:54:ARG:HA	40:B8:59:THR:CB	2.51	0.41
42:D8:46:VAL:HG12	42:D8:47:VAL:H	1.86	0.41
47:E5:21:LEU:HD21	47:E5:41:ARG:NH1	2.34	0.41
47:E5:32:ARG:C	47:E5:34:GLY:H	2.18	0.41
43:E8:74:ALA:HA	43:E8:104:THR:O	2.21	0.41
26:1H:494:G:OP1	43:E8:8:ARG:HD3	2.21	0.41
46:H8:117:LEU:CD1	46:H8:117:LEU:H	2.34	0.41
37:88:85:LYS:HG3	47:I8:7:LEU:HB3	2.02	0.41
52:J5:42:PRO:O	52:J5:44:THR:HG23	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:118:U:C5	1:13:288:A:C6	3.09	0.40
1:13:11:G:C6	1:13:12:U:C4	3.09	0.40
1:13:678:U:H2'	1:13:679:C:C6	2.57	0.40
26:14:1011:G:OP2	41:85:70:ARG:NH2	2.53	0.40
26:14:1139:G:H8	26:14:1139:G:O5'	2.03	0.40
26:14:1171:G:N3	26:14:1173:G:H1'	2.35	0.40
26:14:1418:G:O5'	26:14:1418:G:H8	2.03	0.40
26:14:1449(A):G:H2'	26:14:1450:C:C6	2.54	0.40
26:14:2124:G:H1	26:14:2173:A:H61	1.69	0.40
26:14:2184:G:H2'	26:14:2185:C:C6	2.56	0.40
26:14:2311:A:H8	31:49:88:ILE:HG12	1.85	0.40
26:14:2258:C:O2'	26:14:2427:C:OP2	2.35	0.40
26:14:2676:C:H2'	26:14:2677:G:C8	2.55	0.40
26:14:960:A:C8	26:14:962:G:C8	3.09	0.40
28:19:206:LEU:HD22	28:19:211:ARG:HG2	2.02	0.40
1:1G:109:A:C6	1:1G:326:G:C6	3.09	0.40
1:1G:1206:G:H4'	3:22:192:THR:O	2.21	0.40
1:1G:1273:G:OP2	1:1G:1273:G:H8	2.04	0.40
1:1G:1523:G:C6	1:1G:1524:C:C4	3.09	0.40
1:1G:838:G:N2	1:1G:849:C:C4	2.89	0.40
1:1G:954:G:H2'	1:1G:955:U:C6	2.56	0.40
26:1H:141:A:C8	26:1H:1408:C:H1'	2.56	0.40
26:1H:1854:A:H2'	26:1H:1855:G:O4'	2.22	0.40
26:1H:1957:C:H2'	26:1H:1958:C:C6	2.57	0.40
26:1H:2422:A:N7	54:Q8:30:ARG:NH2	2.55	0.40
26:1H:2471:C:H2'	26:1H:2472:G:O4'	2.21	0.40
26:1H:2757:A:N1	32:51:67:LEU:HD22	2.36	0.40
27:1J:78:A:H2'	27:1J:79:C:O4'	2.21	0.40
26:1H:2052:G:O4'	29:21:142:GLY:HA3	2.21	0.40
3:22:6:HIS:CD2	14:5A:49:HIS:HB3	2.56	0.40
29:29:195:LEU:HD12	29:29:195:LEU:HA	1.81	0.40
29:29:201:THR:HG22	29:29:202:LYS:N	2.37	0.40
29:29:203:LYS:HG2	29:29:205:ALA:HB3	2.02	0.40
29:29:34:VAL:HG21	29:29:78:LEU:HD22	2.03	0.40
3:2E:52:LEU:HD23	3:2E:52:LEU:O	2.21	0.40
23:2K:24:C:H2'	23:2K:25:A:O4'	2.22	0.40
23:2K:87:A:H4'	26:1H:2602:A:C5	2.56	0.40
26:1H:1247:A:OP1	30:31:95:ARG:NH2	2.54	0.40
4:32:92:VAL:O	4:32:96:LEU:HD12	2.20	0.40
37:45:79:LEU:HD13	37:45:79:LEU:H	1.86	0.40
13:4A:25:ILE:HA	13:4A:25:ILE:HD13	1.84	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:4K:34:G:H2'	25:4K:35:A:H8	1.85	0.40
26:1H:2747:G:H5''	32:51:70:THR:HG21	2.03	0.40
6:52:23:LYS:HG2	6:52:61:LEU:HD21	2.03	0.40
38:55:79:LEU:HA	38:55:83:ILE:HB	2.04	0.40
34:58:46:VAL:O	34:58:47:ALA:HB3	2.21	0.40
33:61:129:THR:HG22	33:61:137:PRO:HB3	2.02	0.40
39:65:65:VAL:O	39:65:69:VAL:HG12	2.21	0.40
7:6E:22:LEU:HG	7:6E:62:PHE:HE2	1.87	0.40
15:6I:43:LEU:HA	15:6I:43:LEU:HD23	1.87	0.40
9:82:21:PRO:HA	9:82:59:PHE:HA	2.02	0.40
41:85:25:TRP:CD1	41:85:26:GLY:N	2.89	0.40
17:8I:69:LYS:C	17:8I:70:ARG:HD2	2.42	0.40
39:A8:36:TYR:HB3	39:A8:52:SER:HB3	2.04	0.40
19:AA:71:LEU:HA	19:AA:71:LEU:HD23	1.68	0.40
19:AI:40:ILE:O	19:AI:41:VAL:HG22	2.21	0.40
40:B8:26:ASP:CB	40:B8:92:GLY:H	2.32	0.40
47:E5:7:LEU:O	47:E5:8:ALA:HB2	2.22	0.40
43:E8:79:GLY:CA	43:E8:100:THR:HG22	2.39	0.40
49:G5:4:SER:N	49:G5:6:VAL:HG22	2.36	0.40
46:H8:11:GLU:HA	46:H8:36:LYS:HE3	2.03	0.40
46:H8:152:ALA:HB3	46:H8:167:PRO:HA	2.03	0.40
46:H8:23:LYS:HD3	46:H8:40:ASP:HA	2.02	0.40
46:H8:77:ASP:OD1	46:H8:80:ARG:HD2	2.21	0.40
47:I8:44:ARG:HH11	47:I8:44:ARG:HD2	1.77	0.40
49:K8:2:LYS:O	49:K8:5:GLU:HB2	2.21	0.40
51:M8:26:SER:OG	51:M8:27:THR:N	2.54	0.40
54:Q8:27:THR:HG22	54:Q8:42:ARG:HE	1.86	0.40
2:12:180:LEU:HA	2:12:180:LEU:HD23	1.72	0.40
2:12:188:ALA:N	2:12:201:ILE:O	2.50	0.40
1:13:1010:G:N2	1:13:1019:C:N3	2.63	0.40
1:13:153:C:N4	1:13:168:G:H22	2.18	0.40
1:13:316:G:N3	1:13:316:G:H2'	2.36	0.40
1:13:448:A:P	1:13:485:G:H22	2.45	0.40
1:13:677:U:H3	1:13:713:G:H22	1.69	0.40
1:13:953:G:C6	1:13:954:G:C4	3.09	0.40
1:13:960:U:H2'	1:13:960:U:O2	2.21	0.40
1:13:984:C:H2'	1:13:985:C:C6	2.56	0.40
26:14:1049:C:N3	26:14:2751:G:N1	2.58	0.40
26:14:1057:A:H2'	26:14:1058:U:O4'	2.22	0.40
26:14:2394:C:P	54:M5:30:ARG:NH1	2.94	0.40
26:14:2720:U:N3	26:14:2873:A:H2	2.19	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:275:G:H2'	26:14:276:A:C8	2.56	0.40
26:14:288:C:H2'	26:14:289:A:C8	2.56	0.40
26:14:433:C:C4	26:14:434:U:O4	2.74	0.40
26:14:993:G:O4'	42:95:87:HIS:CD2	2.74	0.40
34:15:39:ARG:C	34:15:41:ASP:H	2.24	0.40
1:1G:1240:U:H2'	7:62:32:ARG:NH2	2.35	0.40
1:1G:1323:G:H4'	1:1G:1362(A):C:C2	2.56	0.40
1:1G:1392:G:H21	1:1G:1502:A:H8	1.67	0.40
1:1G:187:C:H2'	1:1G:188:U:O4'	2.20	0.40
1:1G:370:C:H2'	1:1G:371:G:C8	2.56	0.40
1:1G:711:G:O2'	1:1G:712:A:H5'	2.21	0.40
1:1G:973:G:C1'	10:1A:55:LYS:HG3	2.51	0.40
26:1H:1342:A:N1	26:1H:1345:C:C2	2.89	0.40
26:1H:1375:C:H2'	26:1H:1376:C:H6	1.85	0.40
26:1H:1516:U:C2	26:1H:1517:G:C8	3.10	0.40
26:1H:1413:G:N2	26:1H:1589:C:O2	2.54	0.40
26:1H:1299:G:H3'	26:1H:1639:U:O4	2.21	0.40
26:1H:1683:C:H2'	26:1H:1684:C:C6	2.56	0.40
26:1H:2123:G:H2'	26:1H:2124:G:H8	1.86	0.40
26:1H:2294:C:C4	26:1H:2295:C:C5	3.09	0.40
26:1H:234:C:H2'	26:1H:235:U:C6	2.56	0.40
26:1H:2620:C:H2'	26:1H:2621:A:O4'	2.21	0.40
26:1H:270(R):G:H2'	26:1H:270(S):G:C8	2.57	0.40
26:1H:464:U:H2'	26:1H:465:G:O4'	2.21	0.40
26:1H:643:A:N1	26:1H:2369:A:O2'	2.48	0.40
26:1H:675:A:C6	26:1H:676:A:C6	3.10	0.40
26:1H:704:G:O2'	26:1H:726:G:N2	2.42	0.40
26:1H:774:A:C2	26:1H:787:U:O2'	2.73	0.40
29:21:188:VAL:HA	29:21:189:PRO:HD3	1.90	0.40
29:21:15:PHE:HA	29:21:19:ARG:O	2.21	0.40
29:21:88:GLY:O	29:21:89:ASP:HB2	2.21	0.40
3:22:61:ALA:O	3:22:63:ASN:N	2.52	0.40
3:22:69:HIS:N	3:22:69:HIS:ND1	2.70	0.40
3:22:91:LEU:HB2	3:22:99:VAL:HG21	2.03	0.40
3:22:90:GLU:HA	3:22:93:LYS:HB3	2.03	0.40
29:29:23:VAL:HA	29:29:184:VAL:O	2.21	0.40
1:1G:777:A:H2	11:2A:119:CYS:HB3	1.86	0.40
3:2E:64:VAL:HG12	3:2E:66:VAL:HG23	2.02	0.40
36:35:14:LYS:HE2	36:35:14:LYS:HB3	1.96	0.40
4:3E:8:VAL:HG13	4:3E:21:LEU:HB2	2.03	0.40
31:41:19:LEU:HA	31:41:19:LEU:HD23	1.80	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:41:18:GLU:OE1	31:41:22:ARG:HD2	2.21	0.40
27:1J:31:C:H4'	31:49:29:TRP:HH2	1.86	0.40
5:4E:13:ILE:HA	5:4E:29:GLY:O	2.22	0.40
5:4E:69:VAL:HA	5:4E:70:PRO:HD3	1.78	0.40
13:4I:49:THR:HG22	13:4I:51:ALA:H	1.86	0.40
13:4I:88:ARG:HH11	13:4I:88:ARG:HB3	1.85	0.40
32:51:153:LYS:H	32:51:153:LYS:CD	2.33	0.40
32:51:149:ARG:NH1	32:51:154:PRO:HG3	2.36	0.40
6:52:35:ALA:HB2	6:52:67:MET:HE3	2.03	0.40
34:58:15:LEU:HD13	34:58:16:ILE:N	2.36	0.40
34:58:42:TRP:O	41:C8:64:ARG:HD2	2.21	0.40
32:59:35:VAL:HG21	32:59:75:ALA:HB2	2.03	0.40
39:65:89:ARG:O	39:65:90:GLY:C	2.60	0.40
33:69:64:GLU:HG2	33:69:67:ARG:HD2	2.02	0.40
33:69:77:LEU:CD1	33:69:78:THR:H	2.31	0.40
15:6A:84:LYS:HB2	15:6A:84:LYS:HE2	1.94	0.40
7:6E:104:LEU:HD13	7:6E:104:LEU:HA	1.71	0.40
8:72:84:ARG:O	8:72:135:CYS:HB2	2.22	0.40
40:75:45:PHE:CE2	40:75:74:ARG:HB2	2.57	0.40
36:78:81:GLN:HG2	36:78:106:LEU:HD22	2.04	0.40
36:78:47:ASP:HA	36:78:48:PRO:HD3	1.73	0.40
16:7A:37:GLY:HA3	16:7A:50:LYS:O	2.21	0.40
16:7A:74:LEU:O	16:7A:79:VAL:HG23	2.21	0.40
8:7E:100:ILE:HA	8:7E:101:PRO:HD3	1.72	0.40
8:7E:82:HIS:NE2	8:7E:136:GLU:OE2	2.54	0.40
43:A5:27:LYS:O	43:A5:71:VAL:HG23	2.21	0.40
40:B8:42:ILE:HG21	40:B8:84:GLN:NE2	2.36	0.40
41:C8:20:LEU:HD23	41:C8:20:LEU:HA	1.83	0.40
46:D5:157:LEU:HA	46:D5:157:LEU:HD12	1.76	0.40
49:G5:28:LYS:HA	49:G5:28:LYS:HD3	1.91	0.40
54:M5:41:ILE:O	54:M5:44:LYS:N	2.51	0.40
28:11:142:VAL:HA	28:11:194:GLY:H	1.87	0.40
28:11:232:PRO:HB3	28:11:244:ARG:NH1	2.36	0.40
2:12:223:ILE:H	2:12:223:ILE:HG12	1.66	0.40
1:13:160:A:H1'	1:13:344:A:C5	2.56	0.40
1:13:422:C:H4'	1:13:423:G:O5'	2.21	0.40
1:13:632:A:H8	1:13:633:G:N9	2.19	0.40
1:13:718:G:H5'	11:2I:117:ASN:CG	2.42	0.40
26:14:1160:G:C6	26:14:1161:C:C4	3.09	0.40
26:14:1324:G:C5	26:14:1328:G:O6	2.73	0.40
26:14:1416:G:C2	26:14:1417:C:C4	3.09	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1677:A:O5'	26:14:1677:A:H8	2.04	0.40
26:14:1770:G:H2'	26:14:1771:C:H6	1.87	0.40
26:14:1914:C:C5	26:14:1915:U:C2	3.10	0.40
26:14:2277:G:H5''	37:45:87:LYS:HB2	2.02	0.40
26:14:227:A:C2	26:14:2407:G:H1'	2.56	0.40
26:14:2520:C:H2'	26:14:2521:C:C6	2.56	0.40
26:14:2590:A:OP2	28:19:237:GLU:HB3	2.21	0.40
26:14:196:A:O2'	26:14:805:G:O6	2.36	0.40
26:14:881:G:N3	26:14:881:G:H2'	2.37	0.40
2:1E:25:ASN:O	2:1E:28:PHE:HB2	2.21	0.40
1:1G:1129:C:C4	1:1G:1139:G:C2	3.10	0.40
1:1G:115:G:H4'	1:1G:116:A:O5'	2.21	0.40
1:1G:1192:C:C5	1:1G:1193:G:C8	3.10	0.40
1:1G:1218:C:OP2	14:5A:9:LYS:NZ	2.41	0.40
1:1G:1347:G:C5	9:82:107:ARG:NH1	2.89	0.40
1:1G:1423:G:H2'	1:1G:1424:C:H6	1.86	0.40
1:1G:355:C:C4	1:1G:356:A:N7	2.89	0.40
1:1G:514:C:C2	1:1G:515:G:C8	3.09	0.40
1:1G:567:G:H2'	1:1G:568:G:O4'	2.22	0.40
26:1H:1022:G:O6	34:58:66:LYS:NZ	2.44	0.40
26:1H:1110:G:O2'	26:1H:1111:A:C8	2.75	0.40
26:1H:1153:C:C4	26:1H:1154:G:C5	3.08	0.40
26:1H:1432:C:H2'	26:1H:1433:U:O4'	2.22	0.40
26:1H:1649:G:C6	26:1H:2009:G:C6	3.09	0.40
26:1H:2027:G:C5	26:1H:2028:U:C5	3.09	0.40
26:1H:2261:C:O2'	26:1H:2262:U:H5'	2.20	0.40
26:1H:2364:C:H2'	26:1H:2365:G:O4'	2.21	0.40
26:1H:2850:A:H3'	26:1H:2851:A:H8	1.86	0.40
26:1H:725:G:O5'	26:1H:725:G:H8	2.04	0.40
10:1I:80:LYS:O	10:1I:80:LYS:HD3	2.21	0.40
27:1J:109:G:C6	27:1J:110:G:C5	3.10	0.40
22:1K:53:A:C8	22:1K:54:G:H1'	2.56	0.40
3:22:46:GLU:H	3:22:46:GLU:HG2	1.55	0.40
35:25:61:VAL:HG13	35:25:85:VAL:HG12	2.03	0.40
30:31:34:TRP:NE1	36:78:8:PRO:HD3	2.36	0.40
5:42:76:ILE:CG2	5:42:77:PRO:HD2	2.51	0.40
5:4E:24:ARG:HE	5:4E:24:ARG:HB3	1.58	0.40
6:52:11:ASN:O	6:52:14:LEU:HD22	2.22	0.40
32:59:11:VAL:O	32:59:13:LYS:HG3	2.21	0.40
14:5A:26:ARG:O	14:5A:27:CYS:HB3	2.21	0.40
14:5I:17:LYS:HG3	14:5I:18:VAL:N	2.36	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:72:25:ASP:HB3	8:72:58:TYR:HB3	2.03	0.40
40:75:76:PHE:HA	40:75:77:PRO:HD2	1.91	0.40
36:78:49:ARG:HG3	54:Q8:57:ARG:NE	2.35	0.40
8:7E:80:ILE:H	8:7E:80:ILE:HG12	1.58	0.40
9:8E:53:VAL:HG21	9:8E:85:LEU:HD13	2.03	0.40
43:A5:29:LEU:CG	43:A5:33:ARG:HH21	2.35	0.40
20:BI:62:LEU:HA	20:BI:65:LYS:HB3	2.03	0.40
41:C8:93:LYS:HG3	41:C8:93:LYS:H	1.60	0.40
41:C8:95:LEU:N	41:C8:95:LEU:HD22	2.37	0.40
46:D5:94:GLU:OE1	46:D5:95:PRO:HD2	2.21	0.40
50:H5:8:LEU:HD13	50:H5:31:LEU:HA	2.03	0.40
54:Q8:14:VAL:HG12	54:Q8:23:VAL:HB	2.03	0.40
2:12:221:LEU:HA	2:12:224:GLN:CB	2.52	0.40
1:13:1017:G:H2'	1:13:1018:C:O4'	2.21	0.40
1:13:336:C:O2'	1:13:337:C:H5'	2.21	0.40
1:13:416:G:C5	1:13:417:C:C4	3.10	0.40
1:13:637:G:H2'	1:13:638:G:C8	2.56	0.40
1:13:926:G:C6	1:13:1505:G:C6	3.10	0.40
26:14:141:A:H1'	26:14:1408:C:O4'	2.22	0.40
26:14:141:A:C8	26:14:1408:C:H1'	2.56	0.40
26:14:1773:A:C5	26:14:1829:A:H1'	2.56	0.40
26:14:2697:G:H2'	26:14:2698:U:O4'	2.20	0.40
26:14:446:G:OP2	57:14:3971:HOH:O	2.21	0.40
26:14:579:G:C8	26:14:2017:U:C4	3.10	0.40
26:14:603:A:C8	26:14:604:G:H1'	2.56	0.40
26:14:871:U:H4'	37:45:69:PHE:CD2	2.56	0.40
34:15:29:LYS:HG2	34:15:29:LYS:H	1.71	0.40
10:1A:51:ARG:HB2	10:1A:60:ARG:HA	2.02	0.40
2:1E:21:ARG:HG2	2:1E:22:LYS:HB2	2.03	0.40
1:1G:1036:G:N7	1:1G:1037:C:N4	2.68	0.40
1:1G:149:A:O5'	1:1G:149:A:H8	2.04	0.40
1:1G:173:U:O2	1:1G:197:A:C6	2.75	0.40
1:1G:346:G:N2	1:1G:347:G:C8	2.89	0.40
1:1G:688:G:H2'	1:1G:689:C:C6	2.51	0.40
26:1H:1078:U:O2	26:1H:1088:A:H2	2.04	0.40
26:1H:1234:U:H2'	26:1H:1235:G:H5'	2.04	0.40
26:1H:1387:C:O2	26:1H:1387:C:H2'	2.21	0.40
26:1H:2320:A:H8	26:1H:2321:G:N1	2.20	0.40
26:1H:2404:C:N4	26:1H:2405:G:C6	2.90	0.40
26:1H:270(G):C:O5'	26:1H:270(G):C:H6	2.05	0.40
26:1H:438:G:O5'	26:1H:438:G:H8	2.04	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:463:G:N1	26:1H:467:G:C6	2.90	0.40
26:1H:479:A:HO2'	26:1H:481:G:H8	1.67	0.40
26:1H:861:A:C2	26:1H:917:A:C5	3.09	0.40
22:1K:6:G:H2'	22:1K:7:G:C8	2.56	0.40
22:1L:65:5MU:O5'	22:1L:65:5MU:H6	2.04	0.40
3:22:188:LEU:HA	3:22:188:LEU:HD22	1.82	0.40
29:29:134:ILE:HG22	29:29:137:HIS:CG	2.56	0.40
3:2E:77:ILE:HA	3:2E:84:ILE:HD11	2.04	0.40
23:2L:64:G:O6	23:2L:72:C:N4	2.55	0.40
30:31:28:ILE:HD13	30:31:116:ASP:HB2	2.03	0.40
36:35:92:GLU:HA	36:35:123:LEU:HD22	2.03	0.40
30:39:40:GLN:HG2	30:39:184:TYR:HB2	2.03	0.40
5:42:60:TYR:HB3	5:42:64:ARG:HE	1.86	0.40
31:49:114:ILE:HG22	31:49:117:PHE:HB2	2.03	0.40
31:49:64:THR:OG1	31:49:65:GLY:N	2.55	0.40
32:51:149:ARG:HD2	32:51:149:ARG:HH11	1.71	0.40
38:55:56:LYS:HE3	38:55:94:TYR:CE1	2.57	0.40
6:5E:67:MET:HB2	6:5E:68:PRO:HD2	2.03	0.40
33:61:135:GLU:HB2	33:61:136:VAL:H	1.64	0.40
1:13:808:C:OP1	15:6I:48:LYS:HE2	2.22	0.40
36:78:107:LYS:HB3	36:78:108:LYS:H	1.67	0.40
41:85:39:LEU:HD23	41:85:39:LEU:HA	1.80	0.40
37:88:118:LEU:HD23	37:88:118:LEU:HA	1.87	0.40
39:A8:100:ALA:HA	39:A8:103:GLU:HG2	2.04	0.40
19:AI:52:TYR:CE1	19:AI:56:GLN:HA	2.57	0.40
44:B5:7:VAL:HG12	44:B5:8:ILE:HD12	2.03	0.40
40:B8:110:ILE:O	40:B8:110:ILE:HD13	2.20	0.40
45:C5:8:LYS:HE3	45:C5:8:LYS:HB2	1.84	0.40
48:F5:35:THR:O	48:F5:35:THR:OG1	2.30	0.40
48:F5:83:GLU:N	48:F5:83:GLU:CD	2.75	0.40
46:H8:131:ARG:CZ	46:H8:131:ARG:HB3	2.51	0.40
31:49:145:THR:HG22	51:I5:31:ILE:HG21	2.04	0.40
28:11:120:GLY:HA3	28:11:122:ASP:OD1	2.22	0.40
28:11:134:ARG:HG3	28:11:135:PHE:CE1	2.56	0.40
1:13:1063:C:OP2	1:13:1064:G:O2'	2.36	0.40
1:13:49:U:C2	1:13:361:G:N2	2.90	0.40
1:13:515:G:N2	1:13:537:G:C4	2.89	0.40
1:13:545:C:H5'	4:3E:72:GLU:HB2	2.03	0.40
1:13:577:G:O2'	1:13:578:C:H5'	2.21	0.40
1:13:958:A:C6	1:13:959:A:N1	2.89	0.40
26:14:1280:G:H1	26:14:1290:C:H42	1.68	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1405:U:H2'	26:14:1406:U:H6	1.80	0.40
26:14:1647:G:P	26:14:1647:G:H3'	2.62	0.40
26:14:1859:A:N6	26:14:1883:G:O2'	2.55	0.40
26:14:263:C:H2'	26:14:264:C:O4'	2.21	0.40
26:14:2842:G:H2'	26:14:2843:G:O4'	2.21	0.40
26:14:717:G:H2'	26:14:718:A:O4'	2.21	0.40
26:14:870:A:H2'	26:14:871:U:O4'	2.21	0.40
34:15:99:LEU:O	34:15:99:LEU:HD23	2.21	0.40
10:1A:5:ARG:HB2	10:1A:73:ASP:OD1	2.22	0.40
2:1E:113:HIS:O	2:1E:116:GLU:HB2	2.21	0.40
1:1G:999:U:H2'	1:1G:1000:A:H8	1.87	0.40
1:1G:1014:A:H2'	1:1G:1015:A:C8	2.56	0.40
1:1G:1068:G:N7	1:1G:1094:G:C8	2.90	0.40
1:1G:1071:C:H2'	1:1G:1072:G:H8	1.86	0.40
1:1G:1275:A:N6	1:1G:1276:G:C6	2.90	0.40
1:1G:383:A:H5''	1:1G:384:G:OP2	2.21	0.40
1:1G:532:A:H2	3:22:156:ARG:HH22	1.70	0.40
1:1G:57:G:C5	1:1G:58:C:C4	3.10	0.40
26:1H:1260:G:H2'	26:1H:1261:C:H6	1.86	0.40
26:1H:1590:U:H2'	26:1H:1591:G:H8	1.85	0.40
26:1H:1727:U:H3	26:1H:1733:G:H1	1.70	0.40
26:1H:2085:C:H2'	26:1H:2086:U:O4'	2.22	0.40
26:1H:2291:U:H2'	26:1H:2292:C:C6	2.57	0.40
26:1H:2309:A:N7	26:1H:2310:A:C8	2.90	0.40
26:1H:2352:A:H2'	26:1H:2353:G:O4'	2.21	0.40
26:1H:2557:G:H2'	26:1H:2558:C:C6	2.56	0.40
26:1H:2619:C:O2'	26:1H:2620:C:H5'	2.21	0.40
26:1H:2636:U:OP2	29:21:79:ARG:HD2	2.21	0.40
26:1H:2802:G:H2'	26:1H:2802:G:N3	2.37	0.40
26:1H:2846:G:C5	26:1H:2847:U:C4	3.09	0.40
26:1H:390:A:C6	36:78:71:VAL:HG21	2.56	0.40
26:1H:537:C:H2'	26:1H:539:G:O4'	2.21	0.40
26:1H:598:G:O4'	36:78:12:ALA:HB2	2.21	0.40
26:1H:705:A:C8	26:1H:727:A:C2	3.10	0.40
26:1H:747:U:O2	26:1H:2014:A:H1'	2.22	0.40
26:1H:972:G:OP1	26:1H:974:G:H5'	2.21	0.40
27:1J:1:U:C2'	27:1J:2:C:H5'	2.52	0.40
22:1K:15:A:N1	22:1K:70:G:C6	2.89	0.40
29:21:26:ILE:HD11	29:21:198:VAL:HG21	2.04	0.40
3:2E:101:LEU:HD23	3:2E:102:ASN:N	2.37	0.40
3:2E:178:LEU:HD13	3:2E:178:LEU:HA	1.87	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:2I:48:ILE:HG21	11:2I:63:LEU:HD12	2.03	0.40
12:3A:6:THR:HG23	12:3A:9:GLN:OE1	2.21	0.40
4:3E:187:ARG:HH22	4:3E:193:ASP:CG	2.25	0.40
12:3I:11:VAL:HG13	17:8I:29:HIS:CE1	2.56	0.40
12:3I:39:VAL:H	12:3I:57:LYS:HB3	1.87	0.40
24:3L:3:C:H2'	24:3L:4:G:O4'	2.21	0.40
5:42:101:ILE:HD11	5:42:119:LEU:HD23	2.02	0.40
5:42:50:GLU:HB2	5:42:53:LEU:HD13	2.04	0.40
37:45:16:ARG:HB3	37:45:16:ARG:HE	1.67	0.40
5:4E:139:LEU:HA	5:4E:142:LEU:CD1	2.49	0.40
5:4E:76:ILE:O	5:4E:93:PRO:HB3	2.22	0.40
38:55:21:TYR:OH	38:55:43:GLU:HG2	2.22	0.40
38:55:65:LEU:HA	38:55:65:LEU:HD12	1.77	0.40
3:2E:13:GLY:H	14:5I:57:ARG:HD2	1.86	0.40
35:68:6:THR:HG22	35:68:7:TYR:O	2.22	0.40
33:69:63:ALA:O	33:69:66:GLU:HB2	2.21	0.40
33:69:6:LEU:HA	33:69:6:LEU:HD12	1.70	0.40
8:72:38:ILE:HD11	8:72:118:VAL:O	2.22	0.40
8:72:20:TYR:HD1	8:72:65:TYR:CD2	2.39	0.40
1:1G:1117:G:O3'	9:82:104:ARG:HD2	2.22	0.40
37:88:34:LEU:HD23	37:88:104:PHE:CD2	2.57	0.40
9:8E:111:ARG:HG3	9:8E:112:LYS:H	1.86	0.40
20:BA:75:ASN:N	20:BA:75:ASN:OD1	2.48	0.40
20:BI:10:LEU:HG	20:BI:12:ALA:H	1.86	0.40
45:C5:75:ILE:HA	45:C5:75:ILE:HD12	1.85	0.40
45:C5:90:LEU:HA	45:C5:91:GLU:HA	1.73	0.40
41:C8:34:LYS:HE3	41:C8:34:LYS:HA	2.03	0.40
41:C8:92:ARG:HD3	41:C8:94:ASN:CB	2.52	0.40
48:F5:60:PHE:CE2	48:F5:91:LYS:HD3	2.49	0.40
44:F8:1:MET:CG	44:F8:2:LYS:H	2.33	0.40
50:H5:35:ARG:HD2	50:H5:35:ARG:HH11	1.78	0.40
50:H5:43:ILE:O	50:H5:47:VAL:HG23	2.22	0.40
46:H8:116:VAL:H	46:H8:146:ILE:HG12	1.86	0.40
51:I5:33:VAL:O	51:I5:33:VAL:HG22	2.21	0.40
47:I8:72:ARG:CB	47:I8:75:LEU:HB2	2.51	0.40

All (1) 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)
26:1H:654(O):G:N2	50:H5:57:GLU:OE1[2_464]	2.18	0.02



## 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	
2	12	235/256 (92%)	198 (84%)	35 (15%)	2 (1%)	17	48
2	1E	235/256 (92%)	200 (85%)	33 (14%)	2 (1%)	17	48
3	22	204/239 (85%)	181 (89%)	23 (11%)	0	100	100
3	2E	203/239 (85%)	185 (91%)	18 (9%)	0	100	100
4	32	206/209 (99%)	175 (85%)	30 (15%)	1 (0%)	29	61
4	3E	206/209 (99%)	194 (94%)	9 (4%)	3 (2%)	10	38
5	42	149/162 (92%)	139 (93%)	10 (7%)	0	100	100
5	4E	149/162 (92%)	142 (95%)	6 (4%)	1 (1%)	22	54
6	52	99/101 (98%)	97 (98%)	2 (2%)	0	100	100
6	5E	99/101 (98%)	94 (95%)	5 (5%)	0	100	100
7	62	143/156 (92%)	137 (96%)	6 (4%)	0	100	100
7	6E	140/156 (90%)	134 (96%)	6 (4%)	0	100	100
8	72	136/138 (99%)	125 (92%)	10 (7%)	1 (1%)	22	54
8	7E	136/138 (99%)	124 (91%)	11 (8%)	1 (1%)	22	54
9	82	122/128 (95%)	109 (89%)	12 (10%)	1 (1%)	19	51
9	8E	125/128 (98%)	106 (85%)	19 (15%)	0	100	100
10	1A	97/105 (92%)	90 (93%)	7 (7%)	0	100	100
10	1I	97/105 (92%)	90 (93%)	7 (7%)	0	100	100
11	2A	114/129 (88%)	103 (90%)	8 (7%)	3 (3%)	5	27
11	2I	114/129 (88%)	102 (90%)	10 (9%)	2 (2%)	8	35
12	3A	123/132 (93%)	99 (80%)	20 (16%)	4 (3%)	4	22
12	3I	120/132 (91%)	102 (85%)	18 (15%)	0	100	100
13	4A	114/126 (90%)	95 (83%)	18 (16%)	1 (1%)	17	48
13	4I	114/126 (90%)	95 (83%)	18 (16%)	1 (1%)	17	48
14	5A	56/61 (92%)	46 (82%)	9 (16%)	1 (2%)	8	35

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
14	5I	58/61 (95%)	49 (84%)	8 (14%)	1 (2%)	9	35
15	6A	86/89 (97%)	81 (94%)	5 (6%)	0	100	100
15	6I	86/89 (97%)	80 (93%)	6 (7%)	0	100	100
16	7A	82/88 (93%)	77 (94%)	5 (6%)	0	100	100
16	7I	82/88 (93%)	76 (93%)	6 (7%)	0	100	100
17	8A	97/105 (92%)	92 (95%)	5 (5%)	0	100	100
17	8I	98/105 (93%)	91 (93%)	7 (7%)	0	100	100
18	9A	67/88 (76%)	61 (91%)	6 (9%)	0	100	100
18	9I	65/88 (74%)	63 (97%)	1 (2%)	1 (2%)	10	38
19	AA	76/93 (82%)	62 (82%)	12 (16%)	2 (3%)	5	27
19	AI	78/93 (84%)	65 (83%)	11 (14%)	2 (3%)	5	27
20	BA	97/106 (92%)	87 (90%)	8 (8%)	2 (2%)	7	31
20	BI	97/106 (92%)	82 (84%)	15 (16%)	0	100	100
21	1B	23/27 (85%)	21 (91%)	2 (9%)	0	100	100
21	1F	21/27 (78%)	20 (95%)	1 (5%)	0	100	100
28	11	271/276 (98%)	257 (95%)	10 (4%)	4 (2%)	10	38
28	19	271/276 (98%)	247 (91%)	17 (6%)	7 (3%)	5	27
29	21	203/206 (98%)	162 (80%)	32 (16%)	9 (4%)	2	16
29	29	203/206 (98%)	154 (76%)	38 (19%)	11 (5%)	2	12
30	31	200/210 (95%)	183 (92%)	15 (8%)	2 (1%)	15	46
30	39	206/210 (98%)	165 (80%)	33 (16%)	8 (4%)	3	18
31	41	179/182 (98%)	156 (87%)	20 (11%)	3 (2%)	9	35
31	49	179/182 (98%)	153 (86%)	24 (13%)	2 (1%)	14	45
32	51	172/180 (96%)	148 (86%)	17 (10%)	7 (4%)	3	17
32	59	168/180 (93%)	131 (78%)	33 (20%)	4 (2%)	6	28
33	61	144/148 (97%)	117 (81%)	24 (17%)	3 (2%)	7	31
33	69	144/148 (97%)	116 (81%)	25 (17%)	3 (2%)	7	31
34	15	136/140 (97%)	119 (88%)	16 (12%)	1 (1%)	22	54
34	58	136/140 (97%)	118 (87%)	15 (11%)	3 (2%)	6	30
35	25	120/122 (98%)	115 (96%)	5 (4%)	0	100	100
35	68	120/122 (98%)	114 (95%)	6 (5%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
36	35	148/150 (99%)	115 (78%)	24 (16%)	9 (6%)	1	10
36	78	145/150 (97%)	117 (81%)	23 (16%)	5 (3%)	3	22
37	45	138/141 (98%)	109 (79%)	27 (20%)	2 (1%)	11	38
37	88	139/141 (99%)	115 (83%)	21 (15%)	3 (2%)	6	30
38	55	115/118 (98%)	101 (88%)	12 (10%)	2 (2%)	9	35
38	98	116/118 (98%)	101 (87%)	14 (12%)	1 (1%)	17	48
39	65	109/112 (97%)	87 (80%)	20 (18%)	2 (2%)	8	35
39	A8	109/112 (97%)	92 (84%)	16 (15%)	1 (1%)	17	48
40	75	135/146 (92%)	120 (89%)	14 (10%)	1 (1%)	22	54
40	B8	127/146 (87%)	113 (89%)	14 (11%)	0	100	100
41	85	115/118 (98%)	104 (90%)	10 (9%)	1 (1%)	17	48
41	C8	115/118 (98%)	104 (90%)	8 (7%)	3 (3%)	5	27
42	95	99/101 (98%)	77 (78%)	17 (17%)	5 (5%)	2	13
42	D8	99/101 (98%)	91 (92%)	6 (6%)	2 (2%)	7	32
43	A5	111/113 (98%)	104 (94%)	5 (4%)	2 (2%)	8	35
43	E8	110/113 (97%)	101 (92%)	9 (8%)	0	100	100
44	B5	90/96 (94%)	84 (93%)	4 (4%)	2 (2%)	6	30
44	F8	92/96 (96%)	83 (90%)	7 (8%)	2 (2%)	6	30
45	C5	102/110 (93%)	73 (72%)	24 (24%)	5 (5%)	2	14
45	G8	101/110 (92%)	78 (77%)	19 (19%)	4 (4%)	3	18
46	D5	129/206 (63%)	101 (78%)	24 (19%)	4 (3%)	4	23
46	H8	169/206 (82%)	137 (81%)	26 (15%)	6 (4%)	3	20
47	E5	78/85 (92%)	68 (87%)	9 (12%)	1 (1%)	12	40
47	I8	81/85 (95%)	74 (91%)	7 (9%)	0	100	100
48	F5	92/98 (94%)	82 (89%)	9 (10%)	1 (1%)	14	45
48	J8	93/98 (95%)	84 (90%)	7 (8%)	2 (2%)	6	30
49	G5	65/72 (90%)	61 (94%)	2 (3%)	2 (3%)	4	23
49	K8	66/72 (92%)	61 (92%)	2 (3%)	3 (4%)	2	15
50	H5	57/60 (95%)	55 (96%)	2 (4%)	0	100	100
50	L8	55/60 (92%)	53 (96%)	1 (2%)	1 (2%)	8	35
51	I5	61/71 (86%)	35 (57%)	23 (38%)	3 (5%)	2	14

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
51	M8	64/71 (90%)	41 (64%)	20 (31%)	3 (5%)	2	14
52	J5	54/60 (90%)	50 (93%)	4 (7%)	0	100	100
52	N8	53/60 (88%)	46 (87%)	5 (9%)	2 (4%)	3	19
53	L5	45/49 (92%)	44 (98%)	1 (2%)	0	100	100
53	P8	45/49 (92%)	41 (91%)	3 (7%)	1 (2%)	6	30
54	M5	49/65 (75%)	41 (84%)	7 (14%)	1 (2%)	7	32
54	Q8	58/65 (89%)	34 (59%)	18 (31%)	6 (10%)	0	3
All	All	11130/11946 (93%)	9701 (87%)	1242 (11%)	187 (2%)	9	35

All (187) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
29	21	83	ASP
32	51	169	VAL
36	78	57	THR
38	98	11	ASN
46	H8	165	VAL
49	K8	48	HIS
54	Q8	51	ALA
2	12	7	VAL
9	82	118	LYS
11	2A	48	ILE
19	AA	11	VAL
20	BA	73	HIS
28	19	237	GLU
29	29	25	VAL
29	29	59	VAL
30	39	28	ILE
37	45	27	VAL
37	45	81	VAL
38	55	107	ASP
46	D5	53	ILE
46	D5	165	VAL
47	E5	33	ALA
48	F5	30	VAL
49	G5	47	ASN
54	M5	34	TRP
8	7E	86	ILE
18	9I	22	VAL
29	21	78	LEU

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Mol	Chain	Res	Type
31	41	98	ARG
32	51	168	PRO
34	58	97	ARG
37	88	66	ILE
37	88	90	VAL
41	C8	89	GLU
41	C8	93	LYS
45	G8	81	LYS
46	H8	60	GLU
51	M8	50	VAL
52	N8	41	PRO
52	N8	42	PRO
54	Q8	8	LYS
54	Q8	55	ALA
11	2A	100	ALA
12	3A	18	VAL
14	5A	29	ARG
20	BA	49	ALA
29	29	81	ILE
30	39	84	VAL
36	35	57	THR
39	65	89	ARG
45	C5	29	GLU
49	G5	48	HIS
51	I5	5	ILE
13	4I	83	ASP
19	AI	67	VAL
29	21	118	LYS
33	61	145	VAL
34	58	22	THR
36	78	42	SER
44	F8	68	ARG
45	G8	54	LYS
46	H8	6	LYS
48	J8	75	GLU
48	J8	76	ARG
49	K8	43	GLN
54	Q8	39	LYS
11	2A	101	SER
28	19	273	ARG
29	29	51	PHE
29	29	90	THR

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Mol	Chain	Res	Type
30	39	124	LEU
30	39	167	ALA
33	69	111	PRO
33	69	145	VAL
36	35	35	HIS
36	35	64	LYS
39	65	87	PHE
44	B5	68	ARG
46	D5	161	VAL
2	1E	13	ALA
4	3E	31	CYS
11	2I	91	ARG
14	5I	14	PRO
28	11	122	ASP
29	21	21	VAL
29	21	56	PRO
29	21	60	ASN
33	61	12	LEU
46	H8	59	LEU
49	K8	47	ASN
54	Q8	34	TRP
12	3A	48	PRO
29	29	9	VAL
29	29	26	ILE
30	39	128	ALA
30	39	149	ASP
32	59	92	ILE
36	35	6	LEU
36	35	65	ARG
36	35	110	TYR
43	A5	93	ALA
45	C5	17	SER
51	I5	26	SER
29	21	55	ASN
31	41	5	VAL
31	41	97	ASP
33	61	133	HIS
34	58	128	HIS
45	G8	53	PRO
51	M8	63	TYR
54	Q8	44	LYS
8	72	73	ASP

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Mol	Chain	Res	Type
28	19	33	LEU
28	19	239	ARG
29	29	91	VAL
30	39	25	PRO
31	49	47	LYS
36	35	42	SER
38	55	3	HIS
41	85	93	LYS
42	95	71	LEU
42	95	84	LYS
45	C5	20	TYR
46	D5	167	PRO
2	1E	239	VAL
4	3E	83	SER
4	3E	155	LEU
28	11	3	VAL
28	11	240	ALA
30	31	198	ALA
32	51	12	PRO
32	51	167	GLU
36	78	7	ARG
37	88	3	MET
39	A8	61	ASN
42	D8	47	VAL
44	F8	40	LYS
4	32	32	ALA
28	19	32	SER
28	19	240	ALA
40	75	107	ASP
43	A5	12	ILE
51	I5	33	VAL
11	21	82	VAL
29	21	22	PRO
32	51	8	PRO
32	51	92	ILE
36	78	95	VAL
46	H8	53	ILE
51	M8	5	ILE
53	P8	46	VAL
28	19	3	VAL
32	59	99	VAL
32	59	131	VAL

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Mol	Chain	Res	Type
34	15	128	HIS
36	35	7	ARG
5	4E	115	VAL
2	12	39	ILE
29	29	62	PRO
29	29	77	ILE
31	49	5	VAL
32	59	4	ILE
33	69	144	VAL
36	35	62	LEU
42	95	72	VAL
30	31	24	LEU
42	D8	49	THR
45	G8	76	CYS
46	H8	61	LEU
12	3A	96	VAL
29	29	52	LEU
44	B5	51	VAL
19	AI	9	VAL
28	11	123	ALA
29	21	72	VAL
32	51	127	GLU
36	78	19	VAL
41	C8	88	ILE
50	L8	54	VAL
12	3A	47	LYS
13	4A	84	ILE
19	AA	67	VAL
30	39	89	VAL
45	C5	3	VAL
42	95	37	VAL
42	95	99	ILE
45	C5	76	CYS

### 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	
2	12	205/220 (93%)	158 (77%)	47 (23%)	1	3
2	1E	205/220 (93%)	164 (80%)	41 (20%)	1	5
3	22	160/188 (85%)	122 (76%)	38 (24%)	0	2
3	2E	159/188 (85%)	120 (76%)	39 (24%)	0	2
4	32	180/181 (99%)	144 (80%)	36 (20%)	1	5
4	3E	180/181 (99%)	151 (84%)	29 (16%)	2	10
5	42	116/123 (94%)	83 (72%)	33 (28%)	0	1
5	4E	116/123 (94%)	88 (76%)	28 (24%)	0	2
6	52	90/90 (100%)	66 (73%)	24 (27%)	0	1
6	5E	90/90 (100%)	75 (83%)	15 (17%)	2	10
7	62	121/127 (95%)	94 (78%)	27 (22%)	1	3
7	6E	118/127 (93%)	98 (83%)	20 (17%)	2	9
8	72	119/119 (100%)	101 (85%)	18 (15%)	3	13
8	7E	119/119 (100%)	96 (81%)	23 (19%)	1	6
9	82	95/99 (96%)	83 (87%)	12 (13%)	4	19
9	8E	98/99 (99%)	68 (69%)	30 (31%)	0	1
10	1A	89/92 (97%)	74 (83%)	15 (17%)	2	9
10	1I	89/92 (97%)	72 (81%)	17 (19%)	1	6
11	2A	88/99 (89%)	73 (83%)	15 (17%)	2	9
11	2I	88/99 (89%)	77 (88%)	11 (12%)	4	19
12	3A	104/109 (95%)	71 (68%)	33 (32%)	0	1
12	3I	103/109 (94%)	84 (82%)	19 (18%)	1	7
13	4A	94/101 (93%)	76 (81%)	18 (19%)	1	6
13	4I	94/101 (93%)	67 (71%)	27 (29%)	0	1
14	5A	48/50 (96%)	38 (79%)	10 (21%)	1	4
14	5I	49/50 (98%)	36 (74%)	13 (26%)	0	1
15	6A	79/80 (99%)	72 (91%)	7 (9%)	9	32
15	6I	79/80 (99%)	68 (86%)	11 (14%)	3	16
16	7A	72/74 (97%)	58 (81%)	14 (19%)	1	5
16	7I	72/74 (97%)	58 (81%)	14 (19%)	1	5
17	8A	94/97 (97%)	79 (84%)	15 (16%)	2	11
17	8I	95/97 (98%)	78 (82%)	17 (18%)	2	8

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
18	9A	60/77 (78%)	46 (77%)	14 (23%)	1	3
18	9I	59/77 (77%)	48 (81%)	11 (19%)	1	7
19	AA	67/80 (84%)	55 (82%)	12 (18%)	2	8
19	AI	70/80 (88%)	53 (76%)	17 (24%)	0	2
20	BA	76/82 (93%)	64 (84%)	12 (16%)	2	11
20	BI	76/82 (93%)	61 (80%)	15 (20%)	1	5
21	1B	20/22 (91%)	19 (95%)	1 (5%)	24	55
21	1F	18/22 (82%)	17 (94%)	1 (6%)	21	52
28	11	215/218 (99%)	175 (81%)	40 (19%)	1	7
28	19	214/218 (98%)	172 (80%)	42 (20%)	1	5
29	21	165/166 (99%)	124 (75%)	41 (25%)	0	2
29	29	165/166 (99%)	126 (76%)	39 (24%)	1	3
30	31	161/166 (97%)	126 (78%)	35 (22%)	1	4
30	39	165/166 (99%)	125 (76%)	40 (24%)	0	2
31	41	155/156 (99%)	122 (79%)	33 (21%)	1	4
31	49	155/156 (99%)	123 (79%)	32 (21%)	1	4
32	51	145/148 (98%)	106 (73%)	39 (27%)	0	1
32	59	142/148 (96%)	108 (76%)	34 (24%)	0	2
33	61	122/124 (98%)	95 (78%)	27 (22%)	1	3
33	69	122/124 (98%)	89 (73%)	33 (27%)	0	1
34	15	117/119 (98%)	86 (74%)	31 (26%)	0	1
34	58	117/119 (98%)	91 (78%)	26 (22%)	1	3
35	25	100/100 (100%)	78 (78%)	22 (22%)	1	3
35	68	100/100 (100%)	77 (77%)	23 (23%)	1	3
36	35	116/116 (100%)	78 (67%)	38 (33%)	0	1
36	78	114/116 (98%)	83 (73%)	31 (27%)	0	1
37	45	110/111 (99%)	81 (74%)	29 (26%)	0	1
37	88	111/111 (100%)	89 (80%)	22 (20%)	1	5
38	55	100/101 (99%)	76 (76%)	24 (24%)	0	2
38	98	101/101 (100%)	75 (74%)	26 (26%)	0	2
39	65	87/88 (99%)	61 (70%)	26 (30%)	0	1

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
39	A8	87/88 (99%)	64 (74%)	23 (26%)	0	1
40	75	120/127 (94%)	87 (72%)	33 (28%)	0	1
40	B8	115/127 (91%)	86 (75%)	29 (25%)	0	2
41	85	93/94 (99%)	79 (85%)	14 (15%)	3	13
41	C8	93/94 (99%)	77 (83%)	16 (17%)	2	9
42	95	82/82 (100%)	61 (74%)	21 (26%)	0	2
42	D8	82/82 (100%)	60 (73%)	22 (27%)	0	1
43	A5	92/92 (100%)	74 (80%)	18 (20%)	1	5
43	E8	91/92 (99%)	74 (81%)	17 (19%)	1	7
44	B5	74/78 (95%)	56 (76%)	18 (24%)	0	2
44	F8	76/78 (97%)	62 (82%)	14 (18%)	1	7
45	C5	85/91 (93%)	61 (72%)	24 (28%)	0	1
45	G8	84/91 (92%)	61 (73%)	23 (27%)	0	1
46	D5	125/179 (70%)	94 (75%)	31 (25%)	0	2
46	H8	152/179 (85%)	118 (78%)	34 (22%)	1	3
47	E5	62/67 (92%)	48 (77%)	14 (23%)	1	3
47	I8	66/67 (98%)	50 (76%)	16 (24%)	0	2
48	F5	79/83 (95%)	60 (76%)	19 (24%)	0	2
48	J8	80/83 (96%)	62 (78%)	18 (22%)	1	3
49	G5	63/67 (94%)	49 (78%)	14 (22%)	1	3
49	K8	64/67 (96%)	46 (72%)	18 (28%)	0	1
50	H5	51/52 (98%)	39 (76%)	12 (24%)	1	3
50	L8	49/52 (94%)	36 (74%)	13 (26%)	0	1
51	I5	57/63 (90%)	44 (77%)	13 (23%)	1	3
51	M8	59/63 (94%)	42 (71%)	17 (29%)	0	1
52	J5	48/52 (92%)	37 (77%)	11 (23%)	1	3
52	N8	48/52 (92%)	35 (73%)	13 (27%)	0	1
53	L5	40/42 (95%)	28 (70%)	12 (30%)	0	1
53	P8	40/42 (95%)	29 (72%)	11 (28%)	0	1
54	M5	44/55 (80%)	31 (70%)	13 (30%)	0	1
54	Q8	50/55 (91%)	26 (52%)	24 (48%)	0	0

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles
All	All	9404/9894 (95%)	7297 (78%)	2107 (22%)	<b>1</b> <b>3</b>

All (2107) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
2	1E	6	THR
2	1E	7	VAL
2	1E	8	LYS
2	1E	15	VAL
2	1E	17	PHE
2	1E	24	TRP
2	1E	28	PHE
2	1E	37	ASN
2	1E	39	ILE
2	1E	47	THR
2	1E	48	MET
2	1E	68	ILE
2	1E	71	VAL
2	1E	73	THR
2	1E	74	LYS
2	1E	78	GLN
2	1E	96	ARG
2	1E	113	HIS
2	1E	121	LEU
2	1E	122	PHE
2	1E	128	GLU
2	1E	136	VAL
2	1E	144	ARG
2	1E	145	LEU
2	1E	155	LEU
2	1E	162	ILE
2	1E	163	PHE
2	1E	164	VAL
2	1E	172	ILE
2	1E	178	ARG
2	1E	193	ASP
2	1E	196	LEU
2	1E	200	ILE
2	1E	205	ASP
2	1E	213	LEU
2	1E	217	ARG
2	1E	223	ILE

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Mol	Chain	Res	Type
2	1E	224	GLN
2	1E	230	VAL
2	1E	231	GLU
2	1E	233	SER
3	2E	3	ASN
3	2E	5	ILE
3	2E	11	ARG
3	2E	12	LEU
3	2E	14	ILE
3	2E	16	ARG
3	2E	17	ASP
3	2E	27	LYS
3	2E	29	TYR
3	2E	30	ARG
3	2E	31	HIS
3	2E	33	LEU
3	2E	34	LEU
3	2E	52	LEU
3	2E	58	GLU
3	2E	63	ASN
3	2E	79	ARG
3	2E	82	GLU
3	2E	83	ARG
3	2E	102	ASN
3	2E	104	GLN
3	2E	108	ASN
3	2E	111	LEU
3	2E	116	VAL
3	2E	127	ARG
3	2E	128	PHE
3	2E	134	ILE
3	2E	153	VAL
3	2E	156	ARG
3	2E	161	GLU
3	2E	164	ARG
3	2E	166	GLU
3	2E	167	TRP
3	2E	175	LEU
3	2E	178	LEU
3	2E	179	ARG
3	2E	192	THR
3	2E	195	VAL

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Mol	Chain	Res	Type
3	2E	202	ILE
4	3E	3	ARG
4	3E	10	ARG
4	3E	15	GLU
4	3E	31	CYS
4	3E	35	ARG
4	3E	42	GLN
4	3E	46	LYS
4	3E	49	ARG
4	3E	52	SER
4	3E	53	ASP
4	3E	58	LEU
4	3E	60	GLU
4	3E	78	LEU
4	3E	81	GLU
4	3E	96	LEU
4	3E	99	SER
4	3E	108	LEU
4	3E	122	ARG
4	3E	127	THR
4	3E	135	LEU
4	3E	140	VAL
4	3E	141	ARG
4	3E	151	LYS
4	3E	153	ARG
4	3E	155	LEU
4	3E	160	GLN
4	3E	170	VAL
4	3E	187	ARG
4	3E	200	GLU
5	4E	5	ASP
5	4E	6	PHE
5	4E	10	MET
5	4E	13	ILE
5	4E	16	THR
5	4E	18	ARG
5	4E	20	GLN
5	4E	25	ARG
5	4E	27	ARG
5	4E	31	LEU
5	4E	41	VAL
5	4E	51	VAL

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Mol	Chain	Res	Type
5	4E	53	LEU
5	4E	64	ARG
5	4E	68	GLU
5	4E	71	LEU
5	4E	72	GLN
5	4E	75	THR
5	4E	80	ILE
5	4E	87	SER
5	4E	91	LEU
5	4E	105	VAL
5	4E	112	LEU
5	4E	116	THR
5	4E	126	ARG
5	4E	144	THR
5	4E	152	ARG
5	4E	153	LYS
6	5E	1	MET
6	5E	25	ILE
6	5E	30	LEU
6	5E	31	GLU
6	5E	39	LYS
6	5E	43	LEU
6	5E	54	LYS
6	5E	57	GLN
6	5E	65	VAL
6	5E	73	ASN
6	5E	74	ASP
6	5E	75	LEU
6	5E	79	LEU
6	5E	86	ARG
6	5E	92	LYS
7	6E	12	LEU
7	6E	21	VAL
7	6E	38	LEU
7	6E	45	ASP
7	6E	54	THR
7	6E	56	GLN
7	6E	64	GLN
7	6E	66	VAL
7	6E	75	VAL
7	6E	89	MET
7	6E	90	GLU

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Mol	Chain	Res	Type
7	6E	97	GLN
7	6E	98	SER
7	6E	104	LEU
7	6E	111	ARG
7	6E	113	GLU
7	6E	118	VAL
7	6E	135	VAL
7	6E	138	LYS
7	6E	139	GLU
8	7E	1	MET
8	7E	2	LEU
8	7E	3	THR
8	7E	22	GLU
8	7E	25	ASP
8	7E	26	VAL
8	7E	31	PHE
8	7E	49	GLU
8	7E	52	ASP
8	7E	60	ARG
8	7E	63	LEU
8	7E	68	ARG
8	7E	69	ARG
8	7E	77	GLU
8	7E	80	ILE
8	7E	82	HIS
8	7E	84	ARG
8	7E	88	LYS
8	7E	95	VAL
8	7E	98	LYS
8	7E	102	ARG
8	7E	129	VAL
8	7E	137	VAL
9	8E	9	ARG
9	8E	10	ARG
9	8E	20	ARG
9	8E	23	ASN
9	8E	25	LYS
9	8E	33	PHE
9	8E	34	ASN
9	8E	38	GLN
9	8E	40	LEU
9	8E	42	ARG

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Mol	Chain	Res	Type
9	8E	44	VAL
9	8E	47	LEU
9	8E	53	VAL
9	8E	58	HIS
9	8E	70	LYS
9	8E	79	LEU
9	8E	91	ASP
9	8E	92	TYR
9	8E	93	ARG
9	8E	95	LYS
9	8E	99	LEU
9	8E	108	VAL
9	8E	109	VAL
9	8E	112	LYS
9	8E	118	LYS
9	8E	120	ARG
9	8E	121	ARG
9	8E	124	GLN
9	8E	125	TYR
9	8E	128	ARG
10	1I	5	ARG
10	1I	13	HIS
10	1I	29	ARG
10	1I	30	SER
10	1I	40	LEU
10	1I	43	ARG
10	1I	44	VAL
10	1I	46	ARG
10	1I	49	VAL
10	1I	61	GLU
10	1I	62	HIS
10	1I	70	ARG
10	1I	88	LEU
10	1I	89	ASP
10	1I	92	THR
10	1I	96	ILE
10	1I	101	VAL
11	2I	12	ARG
11	2I	14	VAL
11	2I	51	LYS
11	2I	53	SER
11	2I	96	ARG

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Mol	Chain	Res	Type
11	2I	103	LEU
11	2I	105	VAL
11	2I	106	LYS
11	2I	109	VAL
11	2I	119	CYS
11	2I	124	LYS
12	3I	7	ILE
12	3I	11	VAL
12	3I	20	LYS
12	3I	22	SER
12	3I	33	ARG
12	3I	44	THR
12	3I	60	LEU
12	3I	62	SER
12	3I	67	THR
12	3I	79	GLU
12	3I	81	SER
12	3I	91	LYS
12	3I	92	ASP
12	3I	96	VAL
12	3I	100	ILE
12	3I	102	ARG
12	3I	114	LYS
12	3I	123	LYS
12	3I	126	LYS
13	4I	11	ARG
13	4I	14	ARG
13	4I	17	VAL
13	4I	19	LEU
13	4I	31	LYS
13	4I	34	LEU
13	4I	47	ASP
13	4I	52	GLU
13	4I	54	VAL
13	4I	55	ARG
13	4I	56	LEU
13	4I	57	ARG
13	4I	59	TYR
13	4I	61	GLU
13	4I	64	TRP
13	4I	67	GLU
13	4I	70	LEU

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Mol	Chain	Res	Type
13	4I	77	ASN
13	4I	83	ASP
13	4I	88	ARG
13	4I	94	ARG
13	4I	102	ARG
13	4I	103	THR
13	4I	106	ASN
13	4I	108	ARG
13	4I	110	ARG
13	4I	115	LYS
14	5I	3	ARG
14	5I	4	LYS
14	5I	8	GLU
14	5I	11	LYS
14	5I	12	ARG
14	5I	22	THR
14	5I	23	ARG
14	5I	26	ARG
14	5I	32	SER
14	5I	33	VAL
14	5I	41	ARG
14	5I	44	LEU
14	5I	46	GLU
15	6I	10	LYS
15	6I	22	THR
15	6I	38	ARG
15	6I	39	LEU
15	6I	41	GLU
15	6I	47	LYS
15	6I	54	ARG
15	6I	56	LEU
15	6I	58	MET
15	6I	62	GLN
15	6I	66	LEU
16	7I	1	MET
16	7I	2	VAL
16	7I	4	ILE
16	7I	8	ARG
16	7I	27	LYS
16	7I	43	LYS
16	7I	45	THR
16	7I	50	LYS

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Mol	Chain	Res	Type
16	7I	54	GLU
16	7I	58	TYR
16	7I	62	VAL
16	7I	69	THR
16	7I	72	ARG
16	7I	83	GLU
17	8I	6	LEU
17	8I	14	LYS
17	8I	27	PHE
17	8I	31	LEU
17	8I	34	LYS
17	8I	48	GLU
17	8I	52	LYS
17	8I	59	ILE
17	8I	60	ILE
17	8I	62	SER
17	8I	63	ARG
17	8I	74	LEU
17	8I	81	ARG
17	8I	89	LEU
17	8I	91	ARG
17	8I	92	ARG
17	8I	101	ARG
18	9I	25	THR
18	9I	26	LEU
18	9I	31	LEU
18	9I	32	ARG
18	9I	41	LYS
18	9I	54	ARG
18	9I	56	THR
18	9I	68	LYS
18	9I	76	LEU
18	9I	82	THR
18	9I	85	LEU
19	AI	6	LYS
19	AI	7	LYS
19	AI	25	LYS
19	AI	27	GLU
19	AI	29	ARG
19	AI	30	LEU
19	AI	31	ILE
19	AI	33	THR

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Mol	Chain	Res	Type
19	AI	37	ARG
19	AI	41	VAL
19	AI	43	GLU
19	AI	56	GLN
19	AI	58	VAL
19	AI	61	TYR
19	AI	67	VAL
19	AI	77	THR
19	AI	78	ARG
20	BI	11	SER
20	BI	13	LEU
20	BI	20	LEU
20	BI	22	ARG
20	BI	36	LEU
20	BI	51	GLU
20	BI	53	LEU
20	BI	55	ILE
20	BI	56	MET
20	BI	57	ARG
20	BI	62	LEU
20	BI	74	LYS
20	BI	75	ASN
20	BI	84	LEU
20	BI	99	LEU
21	1F	6	ARG
28	11	5	LYS
28	11	13	ARG
28	11	27	THR
28	11	34	VAL
28	11	35	LYS
28	11	37	LEU
28	11	46	GLN
28	11	54	ARG
28	11	58	HIS
28	11	61	LEU
28	11	64	ILE
28	11	65	ILE
28	11	71	ASP
28	11	89	SER
28	11	94	LEU
28	11	95	LEU
28	11	103	ARG

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Mol	Chain	Res	Type
28	11	105	ILE
28	11	106	ILE
28	11	112	GLN
28	11	118	VAL
28	11	126	GLN
28	11	136	ILE
28	11	138	VAL
28	11	141	VAL
28	11	155	LEU
28	11	157	ARG
28	11	165	ILE
28	11	176	ARG
28	11	192	THR
28	11	204	ILE
28	11	208	LYS
28	11	212	SER
28	11	217	ARG
28	11	218	ARG
28	11	221	VAL
28	11	229	VAL
28	11	257	LEU
28	11	271	ILE
28	11	273	ARG
29	21	12	THR
29	21	13	ARG
29	21	14	ILE
29	21	16	ARG
29	21	17	ASP
29	21	25	VAL
29	21	26	ILE
29	21	33	VAL
29	21	34	VAL
29	21	38	THR
29	21	41	LYS
29	21	45	THR
29	21	47	VAL
29	21	52	LEU
29	21	54	GLN
29	21	59	VAL
29	21	63	LEU
29	21	64	LYS
29	21	66	HIS

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Mol	Chain	Res	Type
29	21	69	LYS
29	21	72	VAL
29	21	78	LEU
29	21	79	ARG
29	21	80	GLU
29	21	82	ARG
29	21	87	GLU
29	21	90	THR
29	21	92	THR
29	21	101	ARG
29	21	107	THR
29	21	108	SER
29	21	116	VAL
29	21	119	ARG
29	21	159	HIS
29	21	167	VAL
29	21	178	GLU
29	21	179	GLU
29	21	181	LEU
29	21	197	ILE
29	21	202	LYS
29	21	203	LYS
30	31	7	TYR
30	31	8	GLN
30	31	9	ILE
30	31	18	ARG
30	31	32	LEU
30	31	33	LEU
30	31	53	THR
30	31	57	VAL
30	31	64	ILE
30	31	68	LYS
30	31	70	THR
30	31	74	ARG
30	31	78	ILE
30	31	82	ILE
30	31	88	VAL
30	31	95	ARG
30	31	98	SER
30	31	101	LEU
30	31	116	ASP
30	31	117	ARG

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Mol	Chain	Res	Type
30	31	122	LYS
30	31	127	GLU
30	31	136	THR
30	31	145	GLU
30	31	149	ASP
30	31	152	GLU
30	31	165	ARG
30	31	170	LEU
30	31	174	VAL
30	31	175	THR
30	31	183	VAL
30	31	192	LEU
30	31	201	VAL
30	31	203	GLN
30	31	206	ILE
31	41	14	GLU
31	41	19	LEU
31	41	20	ILE
31	41	26	GLN
31	41	28	VAL
31	41	31	VAL
31	41	33	ARG
31	41	34	LEU
31	41	43	LEU
31	41	45	GLU
31	41	52	ILE
31	41	58	GLN
31	41	63	ILE
31	41	67	LYS
31	41	70	VAL
31	41	80	PHE
31	41	82	LEU
31	41	86	MET
31	41	90	LEU
31	41	96	ARG
31	41	101	ILE
31	41	102	PHE
31	41	103	LEU
31	41	115	ARG
31	41	128	ARG
31	41	133	LEU
31	41	139	LEU

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Mol	Chain	Res	Type
31	41	145	THR
31	41	153	ARG
31	41	155	MET
31	41	162	THR
31	41	165	THR
31	41	168	GLU
32	51	3	ARG
32	51	4	ILE
32	51	7	LEU
32	51	9	ILE
32	51	15	VAL
32	51	23	ARG
32	51	24	VAL
32	51	40	GLU
32	51	45	VAL
32	51	47	GLU
32	51	49	VAL
32	51	54	ARG
32	51	58	GLU
32	51	63	SER
32	51	64	LEU
32	51	71	LEU
32	51	72	ILE
32	51	77	LYS
32	51	80	SER
32	51	81	GLU
32	51	83	TYR
32	51	87	LEU
32	51	92	ILE
32	51	95	ARG
32	51	97	ARG
32	51	103	LEU
32	51	104	GLU
32	51	105	LEU
32	51	111	HIS
32	51	121	ILE
32	51	125	VAL
32	51	129	THR
32	51	134	SER
32	51	136	ILE
32	51	139	GLN
32	51	144	VAL

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Mol	Chain	Res	Type
32	51	149	ARG
32	51	153	LYS
32	51	169	VAL
33	61	7	GLU
33	61	9	LEU
33	61	15	VAL
33	61	20	ASP
33	61	35	LEU
33	61	38	LEU
33	61	41	GLU
33	61	45	LYS
33	61	58	LEU
33	61	60	GLU
33	61	72	LEU
33	61	74	ASN
33	61	81	VAL
33	61	82	ARG
33	61	86	THR
33	61	92	VAL
33	61	95	LYS
33	61	96	ASP
33	61	108	THR
33	61	110	ASP
33	61	118	LYS
33	61	122	GLU
33	61	135	GLU
33	61	139	GLN
33	61	140	LEU
33	61	142	VAL
33	61	145	VAL
34	58	1	MET
34	58	2	LYS
34	58	7	LYS
34	58	10	GLU
34	58	14	VAL
34	58	15	LEU
34	58	28	THR
34	58	32	THR
34	58	33	LEU
34	58	34	LEU
34	58	42	TRP
34	58	43	THR

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Mol	Chain	Res	Type
34	58	48	MET
34	58	55	VAL
34	58	58	ASP
34	58	60	ILE
34	58	61	ARG
34	58	65	LYS
34	58	67	LEU
34	58	87	LEU
34	58	99	LEU
34	58	120	LEU
34	58	130	HIS
34	58	131	GLN
34	58	133	GLN
34	58	134	ARG
35	68	1	MET
35	68	3	GLN
35	68	8	LEU
35	68	9	GLU
35	68	17	ARG
35	68	19	ILE
35	68	20	MET
35	68	23	ARG
35	68	24	VAL
35	68	25	LEU
35	68	29	ASN
35	68	32	TYR
35	68	47	ILE
35	68	52	VAL
35	68	53	LYS
35	68	66	LYS
35	68	78	ARG
35	68	82	ASN
35	68	94	ARG
35	68	96	THR
35	68	98	VAL
35	68	112	MET
35	68	115	VAL
36	78	6	LEU
36	78	7	ARG
36	78	10	PRO
36	78	15	ARG
36	78	18	ARG

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Mol	Chain	Res	Type
36	78	19	VAL
36	78	27	HIS
36	78	29	LYS
36	78	30	THR
36	78	32	THR
36	78	40	SER
36	78	41	ARG
36	78	45	LEU
36	78	46	LYS
36	78	49	ARG
36	78	50	ARG
36	78	57	THR
36	78	76	LYS
36	78	81	GLN
36	78	90	ARG
36	78	96	THR
36	78	105	LEU
36	78	106	LEU
36	78	112	LEU
36	78	117	GLU
36	78	119	GLU
36	78	126	VAL
36	78	133	SER
36	78	135	LEU
36	78	144	GLU
36	78	147	LEU
37	88	1	MET
37	88	5	ARG
37	88	11	LYS
37	88	16	ARG
37	88	25	ASP
37	88	26	TYR
37	88	35	VAL
37	88	43	THR
37	88	45	GLN
37	88	52	VAL
37	88	55	VAL
37	88	56	ARG
37	88	59	ARG
37	88	77	LYS
37	88	91	GLU
37	88	102	VAL

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Mol	Chain	Res	Type
37	88	103	MET
37	88	109	VAL
37	88	110	THR
37	88	115	MET
37	88	134	ARG
37	88	138	ASP
38	98	2	ARG
38	98	4	LEU
38	98	6	SER
38	98	9	LYS
38	98	12	ARG
38	98	28	LEU
38	98	29	LEU
38	98	33	ARG
38	98	34	ILE
38	98	35	THR
38	98	44	LEU
38	98	57	ARG
38	98	59	ASP
38	98	65	LEU
38	98	67	LEU
38	98	73	VAL
38	98	75	LEU
38	98	76	VAL
38	98	79	LEU
38	98	82	GLU
38	98	94	TYR
38	98	96	ARG
38	98	105	ARG
38	98	111	LEU
38	98	113	LEU
38	98	118	GLU
39	A8	4	LEU
39	A8	5	THR
39	A8	15	ARG
39	A8	17	ARG
39	A8	19	LYS
39	A8	31	SER
39	A8	35	ILE
39	A8	36	TYR
39	A8	41	ASP
39	A8	46	VAL

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Mol	Chain	Res	Type
39	A8	52	SER
39	A8	54	LEU
39	A8	58	LEU
39	A8	69	VAL
39	A8	71	ARG
39	A8	75	GLU
39	A8	78	LEU
39	A8	85	VAL
39	A8	89	ARG
39	A8	101	LEU
39	A8	106	ARG
39	A8	111	GLU
39	A8	112	PHE
40	B8	1	MET
40	B8	13	ARG
40	B8	15	VAL
40	B8	18	ASP
40	B8	19	LEU
40	B8	21	GLU
40	B8	27	THR
40	B8	38	ASN
40	B8	42	ILE
40	B8	50	ILE
40	B8	52	ILE
40	B8	53	ARG
40	B8	58	ASN
40	B8	59	THR
40	B8	62	THR
40	B8	64	ARG
40	B8	65	LYS
40	B8	74	ARG
40	B8	78	LEU
40	B8	80	SER
40	B8	85	LYS
40	B8	86	ILE
40	B8	87	ASP
40	B8	88	ILE
40	B8	98	LYS
40	B8	99	LEU
40	B8	110	ILE
40	B8	111	ARG
40	B8	112	ARG

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Mol	Chain	Res	Type
41	C8	5	LYS
41	C8	12	ARG
41	C8	27	LEU
41	C8	34	LYS
41	C8	52	ARG
41	C8	56	ASP
41	C8	70	ARG
41	C8	74	LEU
41	C8	81	HIS
41	C8	83	LEU
41	C8	85	LYS
41	C8	89	GLU
41	C8	92	ARG
41	C8	94	ASN
41	C8	97	ASP
41	C8	108	GLU
42	D8	5	VAL
42	D8	7	THR
42	D8	11	GLN
42	D8	18	LEU
42	D8	20	LEU
42	D8	22	VAL
42	D8	26	ASP
42	D8	34	GLU
42	D8	35	LEU
42	D8	39	LEU
42	D8	40	LEU
42	D8	44	LYS
42	D8	51	VAL
42	D8	62	LEU
42	D8	64	HIS
42	D8	69	LYS
42	D8	70	ILE
42	D8	72	VAL
42	D8	79	VAL
42	D8	88	ARG
42	D8	89	GLN
42	D8	99	ILE
43	E8	11	ARG
43	E8	15	ARG
43	E8	19	LEU
43	E8	20	VAL

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Mol	Chain	Res	Type
43	E8	42	ARG
43	E8	52	GLU
43	E8	62	HIS
43	E8	68	ARG
43	E8	69	LEU
43	E8	77	ASP
43	E8	88	ARG
43	E8	92	ARG
43	E8	96	ILE
43	E8	97	LYS
43	E8	100	THR
43	E8	107	LEU
43	E8	111	HIS
44	F8	2	LYS
44	F8	9	LEU
44	F8	13	LEU
44	F8	23	GLU
44	F8	27	THR
44	F8	35	THR
44	F8	49	VAL
44	F8	65	ARG
44	F8	70	LEU
44	F8	72	LYS
44	F8	76	ARG
44	F8	80	ILE
44	F8	81	VAL
44	F8	93	GLU
45	G8	6	HIS
45	G8	24	VAL
45	G8	31	LEU
45	G8	33	LYS
45	G8	38	ILE
45	G8	40	GLU
45	G8	43	ASN
45	G8	52	SER
45	G8	54	LYS
45	G8	55	TYR
45	G8	57	GLN
45	G8	61	ILE
45	G8	64	GLU
45	G8	67	LEU
45	G8	71	LYS

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Mol	Chain	Res	Type
45	G8	75	ILE
45	G8	79	CYS
45	G8	85	VAL
45	G8	86	ARG
45	G8	88	LYS
45	G8	95	LYS
45	G8	97	ARG
45	G8	99	CYS
46	H8	1	MET
46	H8	5	LEU
46	H8	11	GLU
46	H8	13	GLU
46	H8	16	SER
46	H8	19	ARG
46	H8	37	VAL
46	H8	40	ASP
46	H8	41	LEU
46	H8	43	GLU
46	H8	50	GLN
46	H8	53	ILE
46	H8	60	GLU
46	H8	61	LEU
46	H8	65	GLN
46	H8	71	VAL
46	H8	76	LEU
46	H8	77	ASP
46	H8	81	ARG
46	H8	82	ARG
46	H8	86	VAL
46	H8	87	ASP
46	H8	91	LEU
46	H8	92	SER
46	H8	94	GLU
46	H8	105	VAL
46	H8	112	ARG
46	H8	116	VAL
46	H8	117	LEU
46	H8	123	ASP
46	H8	145	GLU
46	H8	146	ILE
46	H8	148	ASP
46	H8	154	ASP

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Mol	Chain	Res	Type
47	I8	3	HIS
47	I8	4	LYS
47	I8	30	VAL
47	I8	31	VAL
47	I8	36	ILE
47	I8	41	ARG
47	I8	43	THR
47	I8	44	ARG
47	I8	46	LYS
47	I8	55	ARG
47	I8	64	ASP
47	I8	66	VAL
47	I8	67	VAL
47	I8	70	GLN
47	I8	74	ARG
47	I8	80	HIS
48	J8	4	VAL
48	J8	19	GLN
48	J8	21	ARG
48	J8	25	LYS
48	J8	33	LYS
48	J8	35	THR
48	J8	38	SER
48	J8	41	ARG
48	J8	48	LYS
48	J8	61	ARG
48	J8	65	SER
48	J8	78	LYS
48	J8	80	LEU
48	J8	81	LYS
48	J8	82	LEU
48	J8	83	GLU
48	J8	90	ILE
48	J8	91	LYS
49	K8	6	VAL
49	K8	7	ARG
49	K8	8	LYS
49	K8	14	ARG
49	K8	15	LYS
49	K8	16	LEU
49	K8	24	LEU
49	K8	25	VAL

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
49	K8	38	GLN
49	K8	41	ILE
49	K8	44	LEU
49	K8	47	ASN
49	K8	48	HIS
49	K8	50	ILE
49	K8	53	LEU
49	K8	55	ARG
49	K8	62	THR
49	K8	67	LYS
50	L8	3	ARG
50	L8	7	LYS
50	L8	8	LEU
50	L8	9	VAL
50	L8	13	ILE
50	L8	24	LYS
50	L8	30	ARG
50	L8	36	VAL
50	L8	37	LEU
50	L8	38	GLU
50	L8	40	THR
50	L8	54	VAL
50	L8	55	ARG
51	M8	5	ILE
51	M8	10	VAL
51	M8	15	ILE
51	M8	23	GLU
51	M8	27	THR
51	M8	31	ILE
51	M8	38	LYS
51	M8	39	CYS
51	M8	43	TYR
51	M8	44	THR
51	M8	46	GLN
51	M8	50	VAL
51	M8	51	ASP
51	M8	61	ARG
51	M8	62	ARG
51	M8	63	TYR
51	M8	66	SER
52	N8	8	LYS
52	N8	10	LYS

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Mol	Chain	Res	Type
52	N8	13	LYS
52	N8	16	ARG
52	N8	26	THR
52	N8	29	THR
52	N8	31	VAL
52	N8	36	CYS
52	N8	37	LYS
52	N8	40	LYS
52	N8	51	TYR
52	N8	55	ARG
52	N8	57	VAL
53	P8	1	MET
53	P8	8	ASN
53	P8	9	ARG
53	P8	10	ARG
53	P8	14	LYS
53	P8	23	ARG
53	P8	24	THR
53	P8	41	ARG
53	P8	43	THR
53	P8	46	VAL
53	P8	47	ARG
54	Q8	8	LYS
54	Q8	11	LYS
54	Q8	19	SER
54	Q8	21	LYS
54	Q8	25	MET
54	Q8	26	LYS
54	Q8	30	ARG
54	Q8	31	HIS
54	Q8	32	LEU
54	Q8	34	TRP
54	Q8	35	GLN
54	Q8	37	SER
54	Q8	40	GLU
54	Q8	42	ARG
54	Q8	43	GLN
54	Q8	46	ARG
54	Q8	47	LYS
54	Q8	50	LEU
54	Q8	52	LYS
54	Q8	57	ARG

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Mol	Chain	Res	Type
54	Q8	58	ILE
54	Q8	59	LYS
54	Q8	60	LEU
54	Q8	61	LEU
2	12	5	ILE
2	12	12	GLU
2	12	15	VAL
2	12	17	PHE
2	12	22	LYS
2	12	24	TRP
2	12	31	TYR
2	12	42	ILE
2	12	44	LEU
2	12	48	MET
2	12	55	PHE
2	12	58	ILE
2	12	67	THR
2	12	69	LEU
2	12	71	VAL
2	12	75	LYS
2	12	80	ILE
2	12	82	ARG
2	12	83	MET
2	12	84	GLU
2	12	94	ASN
2	12	98	LEU
2	12	107	THR
2	12	108	ILE
2	12	110	GLN
2	12	114	ARG
2	12	121	LEU
2	12	122	PHE
2	12	124	SER
2	12	129	GLU
2	12	139	LYS
2	12	142	LEU
2	12	145	LEU
2	12	147	LYS
2	12	155	LEU
2	12	170	GLU
2	12	172	ILE
2	12	175	ARG

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Mol	Chain	Res	Type
2	12	178	ARG
2	12	179	LYS
2	12	185	ILE
2	12	191	ASP
2	12	193	ASP
2	12	196	LEU
2	12	200	ILE
2	12	201	ILE
2	12	209	ARG
3	22	5	ILE
3	22	12	LEU
3	22	20	SER
3	22	21	ARG
3	22	22	TRP
3	22	28	GLN
3	22	29	TYR
3	22	34	LEU
3	22	40	ARG
3	22	43	LEU
3	22	47	LEU
3	22	54	ARG
3	22	67	THR
3	22	69	HIS
3	22	76	VAL
3	22	82	GLU
3	22	83	ARG
3	22	84	ILE
3	22	85	ARG
3	22	86	VAL
3	22	90	GLU
3	22	94	LEU
3	22	104	GLN
3	22	105	GLU
3	22	115	LEU
3	22	119	ARG
3	22	120	VAL
3	22	122	GLU
3	22	138	VAL
3	22	140	ARG
3	22	164	ARG
3	22	166	GLU
3	22	167	TRP

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Mol	Chain	Res	Type
3	22	188	LEU
3	22	192	THR
3	22	193	TYR
3	22	196	LEU
3	22	202	ILE
4	32	3	ARG
4	32	4	TYR
4	32	5	ILE
4	32	8	VAL
4	32	12	CYS
4	32	17	VAL
4	32	18	LYS
4	32	24	GLU
4	32	30	LYS
4	32	36	ARG
4	32	52	SER
4	32	53	ASP
4	32	61	LYS
4	32	70	ILE
4	32	81	GLU
4	32	83	SER
4	32	94	LEU
4	32	96	LEU
4	32	107	ARG
4	32	110	PHE
4	32	118	ARG
4	32	122	ARG
4	32	127	THR
4	32	140	VAL
4	32	141	ARG
4	32	145	GLU
4	32	150	GLU
4	32	152	SER
4	32	162	LEU
4	32	168	ARG
4	32	190	ASP
4	32	191	ARG
4	32	192	GLU
4	32	193	ASP
4	32	194	LEU
4	32	200	GLU
5	42	13	ILE

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Mol	Chain	Res	Type
5	42	14	ARG
5	42	16	THR
5	42	19	MET
5	42	24	ARG
5	42	25	ARG
5	42	26	PHE
5	42	31	LEU
5	42	40	ARG
5	42	41	VAL
5	42	43	LEU
5	42	47	LYS
5	42	51	VAL
5	42	60	TYR
5	42	61	TYR
5	42	65	ASN
5	42	66	MET
5	42	68	GLU
5	42	72	GLN
5	42	73	ASN
5	42	75	THR
5	42	76	ILE
5	42	78	HIS
5	42	79	GLU
5	42	81	GLU
5	42	82	VAL
5	42	91	LEU
5	42	100	VAL
5	42	101	ILE
5	42	120	THR
5	42	126	ARG
5	42	127	ASN
5	42	144	THR
6	52	2	ARG
6	52	3	ARG
6	52	14	LEU
6	52	16	GLN
6	52	21	LEU
6	52	24	GLU
6	52	25	ILE
6	52	27	GLN
6	52	28	ARG
6	52	40	VAL

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Mol	Chain	Res	Type
6	52	43	LEU
6	52	46	ARG
6	52	54	LYS
6	52	55	ASP
6	52	57	GLN
6	52	64	GLN
6	52	70	ASP
6	52	73	ASN
6	52	74	ASP
6	52	80	ARG
6	52	81	ILE
6	52	89	MET
6	52	93	SER
6	52	98	LEU
7	62	4	ARG
7	62	6	ARG
7	62	8	GLU
7	62	20	ASP
7	62	21	VAL
7	62	24	THR
7	62	27	ILE
7	62	45	ASP
7	62	51	GLN
7	62	54	THR
7	62	57	GLU
7	62	60	LYS
7	62	62	PHE
7	62	63	LYS
7	62	75	VAL
7	62	85	TYR
7	62	87	VAL
7	62	91	VAL
7	62	94	ARG
7	62	95	ARG
7	62	97	GLN
7	62	104	LEU
7	62	129	GLU
7	62	140	ASP
7	62	143	ARG
7	62	149	ARG
7	62	156	TRP
8	72	1	MET

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Mol	Chain	Res	Type
8	72	2	LEU
8	72	24	THR
8	72	25	ASP
8	72	39	LEU
8	72	52	ASP
8	72	73	ASP
8	72	82	HIS
8	72	84	ARG
8	72	92	ARG
8	72	97	VAL
8	72	99	GLU
8	72	100	ILE
8	72	102	ARG
8	72	107	LEU
8	72	115	SER
8	72	127	LEU
8	72	133	LEU
9	82	7	THR
9	82	38	GLN
9	82	41	VAL
9	82	42	ARG
9	82	53	VAL
9	82	64	THR
9	82	85	LEU
9	82	88	TYR
9	82	91	ASP
9	82	95	LYS
9	82	102	LEU
9	82	104	ARG
10	1A	5	ARG
10	1A	13	HIS
10	1A	17	ASP
10	1A	24	VAL
10	1A	43	ARG
10	1A	48	THR
10	1A	51	ARG
10	1A	55	LYS
10	1A	59	SER
10	1A	60	ARG
10	1A	67	THR
10	1A	70	ARG
10	1A	79	ARG

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Mol	Chain	Res	Type
10	1A	89	ASP
10	1A	96	ILE
11	2A	11	LYS
11	2A	14	VAL
11	2A	29	ILE
11	2A	30	VAL
11	2A	41	THR
11	2A	44	SER
11	2A	78	GLN
11	2A	79	SER
11	2A	83	ILE
11	2A	93	GLN
11	2A	105	VAL
11	2A	109	VAL
11	2A	114	VAL
11	2A	120	ARG
11	2A	124	LYS
12	3A	6	THR
12	3A	17	LYS
12	3A	23	LYS
12	3A	24	VAL
12	3A	27	LEU
12	3A	33	ARG
12	3A	34	ARG
12	3A	37	CYS
12	3A	39	VAL
12	3A	41	ARG
12	3A	42	THR
12	3A	43	VAL
12	3A	44	THR
12	3A	46	LYS
12	3A	49	ASN
12	3A	52	LEU
12	3A	54	LYS
12	3A	57	LYS
12	3A	59	ARG
12	3A	60	LEU
12	3A	64	TYR
12	3A	66	VAL
12	3A	81	SER
12	3A	82	VAL
12	3A	83	VAL

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Mol	Chain	Res	Type
12	3A	84	LEU
12	3A	89	ARG
12	3A	92	ASP
12	3A	98	TYR
12	3A	102	ARG
12	3A	111	LYS
12	3A	117	ARG
12	3A	122	THR
13	4A	7	VAL
13	4A	16	ASP
13	4A	17	VAL
13	4A	19	LEU
13	4A	36	LYS
13	4A	39	ILE
13	4A	44	ARG
13	4A	47	ASP
13	4A	48	LEU
13	4A	66	LEU
13	4A	69	GLU
13	4A	77	ASN
13	4A	82	MET
13	4A	93	ARG
13	4A	94	ARG
13	4A	103	THR
13	4A	108	ARG
13	4A	114	ARG
14	5A	8	GLU
14	5A	11	LYS
14	5A	12	ARG
14	5A	16	PHE
14	5A	17	LYS
14	5A	22	THR
14	5A	23	ARG
14	5A	27	CYS
14	5A	33	VAL
14	5A	37	PHE
15	6A	3	ILE
15	6A	6	GLU
15	6A	31	LEU
15	6A	48	LYS
15	6A	79	ARG
15	6A	83	GLU

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Mol	Chain	Res	Type
15	6A	88	ARG
16	7A	2	VAL
16	7A	8	ARG
16	7A	11	SER
16	7A	27	LYS
16	7A	36	ILE
16	7A	43	LYS
16	7A	45	THR
16	7A	55	ARG
16	7A	62	VAL
16	7A	65	GLN
16	7A	67	THR
16	7A	71	ARG
16	7A	74	LEU
16	7A	82	GLN
17	8A	10	VAL
17	8A	22	LEU
17	8A	24	GLU
17	8A	31	LEU
17	8A	43	LEU
17	8A	52	LYS
17	8A	55	ASP
17	8A	60	ILE
17	8A	68	ARG
17	8A	70	ARG
17	8A	74	LEU
17	8A	75	ARG
17	8A	89	LEU
17	8A	93	GLN
17	8A	100	LYS
18	9A	26	LEU
18	9A	28	GLU
18	9A	29	PHE
18	9A	32	ARG
18	9A	36	ASN
18	9A	42	ARG
18	9A	53	ARG
18	9A	54	ARG
18	9A	59	SER
18	9A	65	ILE
18	9A	82	THR
18	9A	84	LYS

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Mol	Chain	Res	Type
18	9A	86	VAL
18	9A	87	ARG
19	AA	11	VAL
19	AA	23	ASN
19	AA	25	LYS
19	AA	28	LYS
19	AA	29	ARG
19	AA	37	ARG
19	AA	39	THR
19	AA	41	VAL
19	AA	53	ASN
19	AA	66	MET
19	AA	78	ARG
19	AA	83	HIS
20	BA	10	LEU
20	BA	11	SER
20	BA	23	ARG
20	BA	24	LEU
20	BA	29	LYS
20	BA	48	LYS
20	BA	53	LEU
20	BA	58	LYS
20	BA	60	GLU
20	BA	75	ASN
20	BA	88	VAL
20	BA	93	GLU
21	1B	9	ARG
28	19	13	ARG
28	19	14	ARG
28	19	18	VAL
28	19	23	GLU
28	19	25	THR
28	19	31	LYS
28	19	32	SER
28	19	33	LEU
28	19	39	LYS
28	19	43	ARG
28	19	49	ILE
28	19	50	THR
28	19	61	LEU
28	19	64	ILE
28	19	65	ILE

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Mol	Chain	Res	Type
28	19	68	LYS
28	19	72	LYS
28	19	88	ARG
28	19	89	SER
28	19	94	LEU
28	19	101	GLU
28	19	105	ILE
28	19	111	LEU
28	19	147	LEU
28	19	157	ARG
28	19	168	ARG
28	19	173	VAL
28	19	200	ASP
28	19	211	ARG
28	19	212	SER
28	19	217	ARG
28	19	239	ARG
28	19	242	ARG
28	19	244	ARG
28	19	257	LEU
28	19	262	ARG
28	19	263	ARG
28	19	266	SER
28	19	267	SER
28	19	270	ILE
28	19	271	ILE
28	19	273	ARG
29	29	1	MET
29	29	4	ILE
29	29	7	VAL
29	29	14	ILE
29	29	16	ARG
29	29	27	LEU
29	29	38	THR
29	29	44	TYR
29	29	48	GLN
29	29	49	LEU
29	29	52	LEU
29	29	58	ARG
29	29	60	ASN
29	29	63	LEU
29	29	64	LYS

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Mol	Chain	Res	Type
29	29	76	ARG
29	29	77	ILE
29	29	78	LEU
29	29	79	ARG
29	29	82	ARG
29	29	83	ASP
29	29	87	GLU
29	29	90	THR
29	29	93	VAL
29	29	95	ILE
29	29	103	ASP
29	29	107	THR
29	29	111	ARG
29	29	119	ARG
29	29	121	ASN
29	29	134	ILE
29	29	144	ARG
29	29	154	LYS
29	29	166	THR
29	29	175	VAL
29	29	181	LEU
29	29	197	ILE
29	29	201	THR
29	29	202	LYS
30	39	2	LYS
30	39	7	TYR
30	39	8	GLN
30	39	11	VAL
30	39	20	LEU
30	39	23	ASP
30	39	24	LEU
30	39	29	ASN
30	39	32	LEU
30	39	36	VAL
30	39	38	ARG
30	39	50	SER
30	39	57	VAL
30	39	62	ARG
30	39	63	LYS
30	39	67	GLN
30	39	70	THR
30	39	74	ARG

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Mol	Chain	Res	Type
30	39	77	ASP
30	39	78	ILE
30	39	82	ILE
30	39	83	PHE
30	39	100	THR
30	39	110	LEU
30	39	112	MET
30	39	114	VAL
30	39	123	LEU
30	39	127	GLU
30	39	154	VAL
30	39	156	LEU
30	39	158	THR
30	39	174	VAL
30	39	183	VAL
30	39	191	ARG
30	39	192	LEU
30	39	194	MET
30	39	196	LEU
30	39	197	ASP
30	39	201	VAL
30	39	205	ARG
31	49	3	LEU
31	49	9	ARG
31	49	13	GLU
31	49	14	GLU
31	49	16	ARG
31	49	26	GLN
31	49	28	VAL
31	49	33	ARG
31	49	39	ILE
31	49	40	ASN
31	49	53	LEU
31	49	60	LEU
31	49	64	THR
31	49	67	LYS
31	49	70	VAL
31	49	81	LYS
31	49	82	LEU
31	49	96	ARG
31	49	104	GLU
31	49	106	LEU

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Mol	Chain	Res	Type
31	49	109	VAL
31	49	118	ARG
31	49	130	ASN
31	49	146	TYR
31	49	147	ASP
31	49	149	VAL
31	49	152	LEU
31	49	153	ARG
31	49	159	VAL
31	49	165	THR
31	49	167	GLU
31	49	172	LEU
32	59	6	ARG
32	59	7	LEU
32	59	9	ILE
32	59	11	VAL
32	59	24	VAL
32	59	26	VAL
32	59	27	LYS
32	59	32	GLU
32	59	41	MET
32	59	42	ARG
32	59	43	VAL
32	59	44	VAL
32	59	45	VAL
32	59	52	VAL
32	59	67	LEU
32	59	68	THR
32	59	69	ARG
32	59	89	ILE
32	59	94	TYR
32	59	98	LEU
32	59	101	ARG
32	59	105	LEU
32	59	113	VAL
32	59	123	PHE
32	59	125	VAL
32	59	127	GLU
32	59	129	THR
32	59	134	SER
32	59	136	ILE
32	59	139	GLN

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Mol	Chain	Res	Type
32	59	147	ASN
32	59	157	TYR
32	59	159	GLU
32	59	164	TYR
33	69	1	MET
33	69	2	LYS
33	69	4	ILE
33	69	6	LEU
33	69	9	LEU
33	69	17	GLN
33	69	35	LEU
33	69	37	VAL
33	69	40	THR
33	69	44	LEU
33	69	47	LEU
33	69	58	LEU
33	69	60	GLU
33	69	67	ARG
33	69	69	LYS
33	69	73	GLU
33	69	75	LEU
33	69	77	LEU
33	69	78	THR
33	69	79	ILE
33	69	81	VAL
33	69	87	LYS
33	69	101	LEU
33	69	104	GLN
33	69	105	HIS
33	69	109	ILE
33	69	114	LEU
33	69	117	GLU
33	69	125	GLU
33	69	131	LYS
33	69	133	HIS
33	69	138	ILE
33	69	142	VAL
34	15	1	MET
34	15	9	VAL
34	15	16	ILE
34	15	28	THR
34	15	29	LYS

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Mol	Chain	Res	Type
34	15	32	THR
34	15	33	LEU
34	15	34	LEU
34	15	35	ARG
34	15	37	LYS
34	15	42	TRP
34	15	46	VAL
34	15	48	MET
34	15	56	ASN
34	15	61	ARG
34	15	63	THR
34	15	65	LYS
34	15	76	SER
34	15	85	ILE
34	15	87	LEU
34	15	93	THR
34	15	94	HIS
34	15	98	VAL
34	15	104	LYS
34	15	106	MET
34	15	112	LEU
34	15	115	ARG
34	15	121	LYS
34	15	127	ASP
34	15	134	ARG
34	15	136	GLU
35	25	5	GLN
35	25	8	LEU
35	25	10	VAL
35	25	22	ILE
35	25	24	VAL
35	25	29	ASN
35	25	35	VAL
35	25	38	VAL
35	25	44	LYS
35	25	47	ILE
35	25	49	ARG
35	25	70	LYS
35	25	78	ARG
35	25	86	ILE
35	25	91	LEU
35	25	94	ARG

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Mol	Chain	Res	Type
35	25	96	THR
35	25	97	ARG
35	25	105	GLU
35	25	108	GLU
35	25	111	PHE
35	25	117	LEU
36	35	1	MET
36	35	6	LEU
36	35	13	ASN
36	35	15	ARG
36	35	18	ARG
36	35	21	ARG
36	35	27	HIS
36	35	36	LYS
36	35	41	ARG
36	35	45	LEU
36	35	50	ARG
36	35	57	THR
36	35	61	ARG
36	35	62	LEU
36	35	65	ARG
36	35	67	MET
36	35	70	GLN
36	35	75	ILE
36	35	81	GLN
36	35	85	LEU
36	35	91	PHE
36	35	96	THR
36	35	98	GLU
36	35	100	LEU
36	35	101	VAL
36	35	105	LEU
36	35	106	LEU
36	35	110	TYR
36	35	111	ARG
36	35	114	ILE
36	35	125	VAL
36	35	126	VAL
36	35	135	LEU
36	35	138	LEU
36	35	144	GLU
36	35	146	VAL

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Mol	Chain	Res	Type
36	35	147	LEU
36	35	149	GLU
37	45	2	LEU
37	45	3	MET
37	45	7	MET
37	45	22	LYS
37	45	26	TYR
37	45	32	TYR
37	45	42	ILE
37	45	45	GLN
37	45	55	VAL
37	45	59	ARG
37	45	60	ARG
37	45	74	TYR
37	45	75	THR
37	45	77	LYS
37	45	79	LEU
37	45	80	GLU
37	45	85	LYS
37	45	87	LYS
37	45	90	VAL
37	45	105	GLU
37	45	106	VAL
37	45	110	THR
37	45	112	GLU
37	45	116	GLU
37	45	118	LEU
37	45	134	ARG
37	45	138	ASP
37	45	139	GLU
37	45	141	GLN
38	55	2	ARG
38	55	8	ARG
38	55	14	SER
38	55	15	SER
38	55	24	GLN
38	55	26	LYS
38	55	28	LEU
38	55	29	LEU
38	55	35	THR
38	55	36	THR
38	55	44	LEU

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Mol	Chain	Res	Type
38	55	51	LEU
38	55	54	LEU
38	55	56	LYS
38	55	65	LEU
38	55	75	LEU
38	55	81	ASP
38	55	82	GLU
38	55	88	ARG
38	55	95	THR
38	55	100	LEU
38	55	111	LEU
38	55	115	GLU
38	55	117	VAL
39	65	3	ARG
39	65	4	LEU
39	65	12	PHE
39	65	14	VAL
39	65	17	ARG
39	65	20	ARG
39	65	24	LEU
39	65	25	ARG
39	65	27	SER
39	65	36	TYR
39	65	38	GLN
39	65	48	LEU
39	65	50	SER
39	65	52	SER
39	65	54	LEU
39	65	65	VAL
39	65	69	VAL
39	65	73	LEU
39	65	82	ILE
39	65	88	ASP
39	65	98	VAL
39	65	101	LEU
39	65	106	ARG
39	65	107	GLU
39	65	110	LEU
39	65	112	PHE
40	75	7	ILE
40	75	8	LYS
40	75	9	LEU

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Mol	Chain	Res	Type
40	75	13	ARG
40	75	15	VAL
40	75	19	LEU
40	75	23	ARG
40	75	27	THR
40	75	36	GLU
40	75	38	ASN
40	75	43	GLN
40	75	49	VAL
40	75	51	ARG
40	75	54	ARG
40	75	55	ASN
40	75	57	PHE
40	75	59	THR
40	75	64	ARG
40	75	72	VAL
40	75	74	ARG
40	75	86	ILE
40	75	87	ASP
40	75	88	ILE
40	75	93	ARG
40	75	105	LEU
40	75	106	SER
40	75	107	ASP
40	75	112	ARG
40	75	115	ARG
40	75	121	ILE
40	75	124	ASP
40	75	125	ARG
40	75	136	GLN
41	85	5	LYS
41	85	8	VAL
41	85	52	ARG
41	85	55	ARG
41	85	56	ASP
41	85	63	VAL
41	85	64	ARG
41	85	74	LEU
41	85	78	THR
41	85	92	ARG
41	85	97	ASP
41	85	100	VAL

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Mol	Chain	Res	Type
41	85	101	ARG
41	85	114	LYS
42	95	19	LYS
42	95	32	THR
42	95	33	VAL
42	95	38	LEU
42	95	39	LEU
42	95	40	LEU
42	95	44	LYS
42	95	45	THR
42	95	46	VAL
42	95	47	VAL
42	95	49	THR
42	95	66	ARG
42	95	70	ILE
42	95	75	PHE
42	95	76	LYS
42	95	79	VAL
42	95	83	ARG
42	95	84	LYS
42	95	91	TYR
42	95	95	LEU
42	95	97	LYS
43	A5	11	ARG
43	A5	17	VAL
43	A5	18	ARG
43	A5	19	LEU
43	A5	23	LEU
43	A5	39	THR
43	A5	50	VAL
43	A5	51	LEU
43	A5	65	LEU
43	A5	70	TYR
43	A5	76	VAL
43	A5	88	ARG
43	A5	92	ARG
43	A5	95	ILE
43	A5	96	ILE
43	A5	103	ILE
43	A5	107	LEU
43	A5	110	LYS
44	B5	12	VAL

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Mol	Chain	Res	Type
44	B5	14	SER
44	B5	30	VAL
44	B5	38	GLU
44	B5	40	LYS
44	B5	41	ASN
44	B5	45	THR
44	B5	48	LYS
44	B5	52	VAL
44	B5	54	VAL
44	B5	57	LEU
44	B5	63	LYS
44	B5	65	ARG
44	B5	66	LEU
44	B5	75	ASP
44	B5	82	GLN
44	B5	83	VAL
44	B5	88	LYS
45	C5	2	ARG
45	C5	6	HIS
45	C5	14	LEU
45	C5	23	ARG
45	C5	33	LYS
45	C5	37	VAL
45	C5	38	ILE
45	C5	40	GLU
45	C5	43	ASN
45	C5	47	LYS
45	C5	50	ARG
45	C5	51	VAL
45	C5	55	TYR
45	C5	62	GLU
45	C5	64	GLU
45	C5	71	LYS
45	C5	84	ARG
45	C5	85	VAL
45	C5	86	ARG
45	C5	89	PHE
45	C5	94	LYS
45	C5	97	ARG
45	C5	99	CYS
45	C5	102	CYS
46	D5	10	ARG

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Mol	Chain	Res	Type
46	D5	16	SER
46	D5	18	LEU
46	D5	19	ARG
46	D5	28	MET
46	D5	31	ARG
46	D5	44	PHE
46	D5	45	ASP
46	D5	56	VAL
46	D5	59	LEU
46	D5	63	ASP
46	D5	70	LEU
46	D5	71	VAL
46	D5	72	ARG
46	D5	74	VAL
46	D5	77	ASP
46	D5	82	ARG
46	D5	89	PHE
46	D5	91	LEU
46	D5	93	ASP
46	D5	98	MET
46	D5	102	LEU
46	D5	103	ARG
46	D5	122	ARG
46	D5	128	VAL
46	D5	129	SER
46	D5	132	ASN
46	D5	139	VAL
46	D5	161	VAL
46	D5	165	VAL
46	D5	168	GLU
47	E5	10	THR
47	E5	12	ASN
47	E5	16	SER
47	E5	29	GLN
47	E5	36	ILE
47	E5	41	ARG
47	E5	43	THR
47	E5	46	LYS
47	E5	53	MET
47	E5	58	THR
47	E5	60	PHE
47	E5	68	GLU

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Mol	Chain	Res	Type
47	E5	81	VAL
47	E5	82	ARG
48	F5	2	SER
48	F5	11	ARG
48	F5	21	ARG
48	F5	25	LYS
48	F5	26	ARG
48	F5	38	SER
48	F5	42	GLN
48	F5	52	ARG
48	F5	56	GLN
48	F5	61	ARG
48	F5	62	VAL
48	F5	74	VAL
48	F5	76	ARG
48	F5	78	LYS
48	F5	80	LEU
48	F5	82	LEU
48	F5	83	GLU
48	F5	85	LEU
48	F5	91	LYS
49	G5	14	ARG
49	G5	15	LYS
49	G5	19	VAL
49	G5	24	LEU
49	G5	25	VAL
49	G5	32	LEU
49	G5	34	GLU
49	G5	35	LEU
49	G5	46	GLN
49	G5	48	HIS
49	G5	53	LEU
49	G5	55	ARG
49	G5	62	THR
49	G5	70	GLN
50	H5	5	LYS
50	H5	6	VAL
50	H5	8	LEU
50	H5	20	LYS
50	H5	24	LYS
50	H5	30	ARG
50	H5	32	GLN

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Mol	Chain	Res	Type
50	H5	33	GLN
50	H5	36	VAL
50	H5	38	GLU
50	H5	40	THR
50	H5	56	VAL
51	I5	20	ASN
51	I5	22	ILE
51	I5	30	GLU
51	I5	33	VAL
51	I5	34	GLU
51	I5	44	THR
51	I5	46	GLN
51	I5	53	GLU
51	I5	58	ARG
51	I5	59	PHE
51	I5	60	GLN
51	I5	61	ARG
51	I5	62	ARG
52	J5	6	VAL
52	J5	8	LYS
52	J5	11	THR
52	J5	16	ARG
52	J5	25	LEU
52	J5	26	THR
52	J5	29	THR
52	J5	35	GLU
52	J5	48	GLU
52	J5	52	TYR
52	J5	55	ARG
53	L5	1	MET
53	L5	4	THR
53	L5	8	ASN
53	L5	11	LYS
53	L5	14	LYS
53	L5	24	THR
53	L5	32	LYS
53	L5	33	ARG
53	L5	41	ARG
53	L5	43	THR
53	L5	46	VAL
53	L5	47	ARG
54	M5	3	LYS

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Mol	Chain	Res	Type
54	M5	6	THR
54	M5	14	VAL
54	M5	15	LYS
54	M5	22	VAL
54	M5	25	MET
54	M5	30	ARG
54	M5	31	HIS
54	M5	36	LYS
54	M5	37	SER
54	M5	58	ILE
54	M5	59	LYS
54	M5	60	LEU

Some sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (17) such sidechains are listed below:

Mol	Chain	Res	Type
3	2E	104	GLN
3	2E	107	GLN
3	2E	108	ASN
10	1I	84	GLN
28	11	115	GLN
31	41	58	GLN
37	88	12	GLN
44	F8	31	HIS
46	H8	73	GLN
46	H8	151	HIS
13	4A	92	HIS
18	9A	36	ASN
28	19	87	ASN
29	29	55	ASN
32	59	61	HIS
39	65	95	HIS
45	C5	6	HIS

### 5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	13	1505/1522 (98%)	327 (21%)	31 (2%)
1	1G	1512/1522 (99%)	370 (24%)	38 (2%)
22	1K	86/87 (98%)	23 (26%)	1 (1%)

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Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
22	1L	86/87 (98%)	31 (36%)	4 (4%)
23	2K	77/87 (88%)	31 (40%)	8 (10%)
23	2L	79/87 (90%)	32 (40%)	8 (10%)
24	3K	75/87 (86%)	35 (46%)	4 (5%)
24	3L	72/87 (82%)	34 (47%)	5 (6%)
25	4K	14/60 (23%)	6 (42%)	0
25	4L	19/60 (31%)	5 (26%)	0
26	14	2908/2917 (99%)	696 (23%)	40 (1%)
26	1H	2911/2917 (99%)	672 (23%)	52 (1%)
27	16	121/122 (99%)	21 (17%)	1 (0%)
27	1J	121/122 (99%)	36 (29%)	4 (3%)
All	All	9586/9764 (98%)	2319 (24%)	196 (2%)

All (2319) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	13	5	U
1	13	6	G
1	13	8	A
1	13	9	G
1	13	10	A
1	13	19	C
1	13	20	U
1	13	32	A
1	13	39	G
1	13	47	C
1	13	48	C
1	13	50	A
1	13	51	A
1	13	54	C
1	13	59	A
1	13	61	G
1	13	65	U
1	13	66	G
1	13	74	C
1	13	77	C
1	13	78	G
1	13	80	G
1	13	91	C
1	13	101	A
1	13	108	G
1	13	116	A

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Mol	Chain	Res	Type
1	13	121	C
1	13	131	C
1	13	137	C
1	13	143	A
1	13	144	G
1	13	151	A
1	13	160	A
1	13	161	A
1	13	164	U
1	13	169	C
1	13	171	A
1	13	172	A
1	13	173	U
1	13	174	C
1	13	180	U
1	13	182	U
1	13	189	U
1	13	190	G
1	13	191(A)	G
1	13	195	A
1	13	196	A
1	13	201	C
1	13	208	U
1	13	209	U
1	13	210	U
1	13	216	G
1	13	217	C
1	13	222	U
1	13	231	G
1	13	244	U
1	13	245	C
1	13	247	G
1	13	251	G
1	13	266	G
1	13	267	C
1	13	270	A
1	13	281	G
1	13	289	G
1	13	316	G
1	13	317	G
1	13	321	A
1	13	328	C

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Mol	Chain	Res	Type
1	13	329	A
1	13	330	C
1	13	332	G
1	13	344	A
1	13	345	C
1	13	346	G
1	13	347	G
1	13	352	C
1	13	353	A
1	13	354	G
1	13	356	A
1	13	365	U
1	13	367	U
1	13	369	C
1	13	372	C
1	13	373	A
1	13	382	A
1	13	384	G
1	13	389	A
1	13	390	C
1	13	397	A
1	13	398	C
1	13	406	G
1	13	412	A
1	13	413	G
1	13	414	A
1	13	423	G
1	13	424	G
1	13	429	U
1	13	430	A
1	13	455	C
1	13	465	A
1	13	466	C
1	13	467	G
1	13	485	G
1	13	496	A
1	13	497	U
1	13	500	G
1	13	505	G
1	13	509	A
1	13	511	C
1	13	513	C

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Mol	Chain	Res	Type
1	13	518	C
1	13	521	G
1	13	527	G
1	13	531	U
1	13	532	A
1	13	533	A
1	13	547	A
1	13	549	C
1	13	559	A
1	13	561	U
1	13	572	A
1	13	573	A
1	13	576	G
1	13	577	G
1	13	595	G
1	13	596	C
1	13	607	A
1	13	610	G
1	13	615	C
1	13	616	G
1	13	619	U
1	13	630	G
1	13	631	G
1	13	632	A
1	13	633	G
1	13	649	G
1	13	652	U
1	13	653	A
1	13	665	A
1	13	666	G
1	13	678	U
1	13	704	A
1	13	707	C
1	13	717	C
1	13	723	U
1	13	724	G
1	13	748	C
1	13	749	C
1	13	753	A
1	13	755	G
1	13	759	A
1	13	766	A

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Mol	Chain	Res	Type
1	13	774	G
1	13	777	A
1	13	792	A
1	13	793	U
1	13	794	A
1	13	817	C
1	13	828	A
1	13	836	G
1	13	841	U
1	13	842	C
1	13	843	U
1	13	848	C
1	13	859	A
1	13	864	A
1	13	870	U
1	13	873	A
1	13	887	G
1	13	891	U
1	13	902	G
1	13	914	A
1	13	926	G
1	13	927	G
1	13	934	C
1	13	935	A
1	13	936	C
1	13	940	C
1	13	948	C
1	13	960	U
1	13	967	C
1	13	968	A
1	13	969	A
1	13	971	G
1	13	974	A
1	13	975	A
1	13	976	G
1	13	977	A
1	13	983	A
1	13	991	U
1	13	992	U
1	13	993	G
1	13	997	U
1	13	1004	A

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Mol	Chain	Res	Type
1	13	1006	C
1	13	1007	C
1	13	1008	C
1	13	1009	G
1	13	1020	U
1	13	1021	G
1	13	1023	G
1	13	1024	G
1	13	1025	U
1	13	1026	G
1	13	1028	C
1	13	1028(B)	C
1	13	1029	G
1	13	1030	C
1	13	1032(A)	G
1	13	1033	G
1	13	1037	C
1	13	1040	U
1	13	1042	G
1	13	1053	G
1	13	1054	C
1	13	1064	G
1	13	1065	U
1	13	1066	C
1	13	1081	G
1	13	1094	G
1	13	1095	U
1	13	1101	A
1	13	1116	C
1	13	1121	U
1	13	1124	G
1	13	1125	U
1	13	1126	U
1	13	1127	G
1	13	1129	C
1	13	1130	A
1	13	1131	G
1	13	1133	G
1	13	1136	U
1	13	1137	C
1	13	1138	G
1	13	1139	G

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Mol	Chain	Res	Type
1	13	1144	G
1	13	1146	A
1	13	1152	A
1	13	1154	G
1	13	1157	A
1	13	1158	C
1	13	1159	U
1	13	1161	C
1	13	1171	G
1	13	1177	G
1	13	1178	G
1	13	1181	G
1	13	1183	A
1	13	1187	G
1	13	1189	C
1	13	1190	G
1	13	1196	U
1	13	1197	G
1	13	1201	A
1	13	1212	U
1	13	1213	A
1	13	1215	G
1	13	1218	C
1	13	1225	A
1	13	1226	C
1	13	1227	A
1	13	1238	A
1	13	1240	U
1	13	1253	G
1	13	1256	A
1	13	1257	U
1	13	1258	G
1	13	1260	C
1	13	1270	C
1	13	1273	G
1	13	1275	A
1	13	1278	U
1	13	1279	A
1	13	1280	A
1	13	1281	U
1	13	1282	C
1	13	1286	A

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Mol	Chain	Res	Type
1	13	1287	A
1	13	1295	G
1	13	1299	A
1	13	1300	G
1	13	1305	G
1	13	1320	C
1	13	1322	C
1	13	1323	G
1	13	1331	G
1	13	1336	C
1	13	1337	G
1	13	1340	A
1	13	1346	A
1	13	1347	G
1	13	1353	G
1	13	1360	A
1	13	1362(A)	C
1	13	1370	G
1	13	1379	G
1	13	1381	U
1	13	1382	C
1	13	1394	A
1	13	1398	A
1	13	1419	G
1	13	1442	G
1	13	1443	G
1	13	1446	A
1	13	1447	G
1	13	1449	C
1	13	1451	A
1	13	1452	C
1	13	1453	G
1	13	1487	G
1	13	1492	A
1	13	1497	G
1	13	1498	U
1	13	1499	A
1	13	1502	A
1	13	1504	G
1	13	1505	G
1	13	1506	U
1	13	1517	G

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Mol	Chain	Res	Type
1	13	1519	A
1	13	1529	G
1	13	1530	G
1	13	1531	A
1	13	1533	C
1	13	1536	C
1	13	1537	U
22	1K	8	U
22	1K	9	G
22	1K	13	G
22	1K	14	A
22	1K	15	A
22	1K	16	U
22	1K	17	U
22	1K	18	G
22	1K	19	G
22	1K	20	U
22	1K	21	A
22	1K	22	G
22	1K	24	C
22	1K	26	C
22	1K	48	C
22	1K	49	C
22	1K	50	A
22	1K	51	A
22	1K	54	G
22	1K	56	G
22	1K	58	U
22	1K	74	C
22	1K	85	C
23	2K	3	C
23	2K	7	G
23	2K	8	U
23	2K	14	A
23	2K	16	U
23	2K	17	H2U
23	2K	18	OMG
23	2K	19	G
23	2K	20	U
23	2K	21	A
23	2K	22	G
23	2K	23	A

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Mol	Chain	Res	Type
23	2K	25	A
23	2K	27	G
23	2K	28	C
23	2K	37	G
23	2K	44	G
23	2K	45	U
23	2K	46	G
23	2K	47	C
23	2K	57	C
23	2K	62	G
23	2K	63	G
23	2K	64	G
23	2K	67	C
23	2K	69	A
23	2K	71	U
23	2K	74	C
23	2K	75	G
23	2K	76	U
23	2K	87	A
24	3K	2	C
24	3K	8	U
24	3K	9	G
24	3K	13	G
24	3K	14	A
24	3K	15	A
24	3K	16	U
24	3K	17	U
24	3K	18	G
24	3K	19	G
24	3K	20	U
24	3K	21	A
24	3K	23	A
24	3K	24	C
24	3K	25	A
24	3K	27	G
24	3K	35	G
24	3K	38	G
24	3K	44	G
24	3K	45	U
24	3K	46	G
24	3K	47	C
24	3K	59	U

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Mol	Chain	Res	Type
24	3K	60	A
24	3K	61	C
24	3K	66	U
24	3K	69	A
24	3K	70	G
24	3K	71	U
24	3K	78	C
24	3K	79	U
24	3K	80	C
24	3K	81	G
24	3K	84	A
24	3K	87	A
25	4K	33	G
25	4K	34	G
25	4K	35	A
25	4K	37	G
25	4K	38	U
25	4K	52	U
26	1H	5	A
26	1H	9	U
26	1H	12	U
26	1H	15	G
26	1H	34	C
26	1H	35	G
26	1H	46	C
26	1H	51	G
26	1H	63	U
26	1H	64	A
26	1H	66	C
26	1H	71	A
26	1H	72	U
26	1H	74	A
26	1H	75	G
26	1H	85	G
26	1H	90	U
26	1H	102	G
26	1H	118	A
26	1H	119	A
26	1H	120	U
26	1H	125	G
26	1H	146	G
26	1H	155	C

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Mol	Chain	Res	Type
26	1H	162	U
26	1H	163	U
26	1H	164	U
26	1H	171	G
26	1H	173	G
26	1H	181	A
26	1H	188	G
26	1H	196	A
26	1H	199	A
26	1H	201	C
26	1H	215	G
26	1H	216	A
26	1H	222	A
26	1H	223	A
26	1H	224	G
26	1H	228	A
26	1H	229	A
26	1H	232	G
26	1H	233	A
26	1H	235	U
26	1H	238	C
26	1H	248	G
26	1H	252	G
26	1H	261	G
26	1H	266	G
26	1H	269	U
26	1H	270(F)	U
26	1H	270(H)	C
26	1H	270(K)	C
26	1H	270(M)	U
26	1H	270(N)	G
26	1H	270(P)	C
26	1H	271(B)	G
26	1H	271(C)	U
26	1H	271	G
26	1H	273(E)	U
26	1H	274	G
26	1H	275	G
26	1H	278	A
26	1H	299	A
26	1H	311	A
26	1H	324	A

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Mol	Chain	Res	Type
26	1H	329	G
26	1H	330	A
26	1H	331	A
26	1H	334	C
26	1H	335	C
26	1H	338	G
26	1H	352	G
26	1H	363	G
26	1H	363(F)	A
26	1H	364	C
26	1H	372	G
26	1H	380	U
26	1H	386	G
26	1H	396	G
26	1H	405	U
26	1H	407	G
26	1H	411	G
26	1H	428	A
26	1H	443	A
26	1H	444	C
26	1H	448	U
26	1H	452	G
26	1H	455	C
26	1H	457	A
26	1H	463	G
26	1H	467	G
26	1H	470	A
26	1H	471	A
26	1H	481	G
26	1H	482	A
26	1H	491	G
26	1H	501	A
26	1H	504	U
26	1H	505	A
26	1H	508	G
26	1H	509	C
26	1H	528	A
26	1H	529	A
26	1H	530	G
26	1H	531	C
26	1H	532	A
26	1H	533	G

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Mol	Chain	Res	Type
26	1H	546	C
26	1H	547	A
26	1H	549	G
26	1H	563	G
26	1H	573	G
26	1H	575	A
26	1H	586	A
26	1H	587	C
26	1H	588	U
26	1H	603	A
26	1H	607	U
26	1H	614	U
26	1H	615	G
26	1H	617	G
26	1H	618	G
26	1H	621	A
26	1H	622	G
26	1H	626	U
26	1H	627	A
26	1H	631	A
26	1H	637	A
26	1H	645	C
26	1H	646	A
26	1H	654	A
26	1H	654(H)	G
26	1H	654(I)	C
26	1H	654(K)	C
26	1H	654(M)	C
26	1H	654(O)	G
26	1H	654(V)	A
26	1H	665	C
26	1H	668	G
26	1H	669	G
26	1H	676	A
26	1H	686	G
26	1H	714	U
26	1H	717	G
26	1H	730	C
26	1H	738	G
26	1H	747	U
26	1H	752	A
26	1H	753	C

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Mol	Chain	Res	Type
26	1H	762	U
26	1H	764	A
26	1H	765	G
26	1H	775	G
26	1H	776	G
26	1H	782	A
26	1H	784	A
26	1H	785	G
26	1H	790	C
26	1H	791	C
26	1H	792	G
26	1H	801	G
26	1H	805	G
26	1H	812	C
26	1H	813	U
26	1H	827	U
26	1H	828	U
26	1H	836	G
26	1H	845	G
26	1H	846	C
26	1H	847	U
26	1H	859	G
26	1H	877	U
26	1H	878	A
26	1H	879	G
26	1H	880	G
26	1H	881	G
26	1H	882	G
26	1H	883	G
26	1H	884	C
26	1H	885	C
26	1H	886	C
26	1H	887	A
26	1H	888	C
26	1H	890	A
26	1H	892	G
26	1H	893	C
26	1H	894	C
26	1H	895	U
26	1H	896	A
26	1H	900	A
26	1H	901	A

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Mol	Chain	Res	Type
26	1H	907	U
26	1H	910	A
26	1H	914	C
26	1H	917	A
26	1H	918	A
26	1H	932	G
26	1H	940	G
26	1H	941	A
26	1H	946	G
26	1H	953	A
26	1H	957	A
26	1H	959	A
26	1H	961	C
26	1H	974	G
26	1H	974(A)	C
26	1H	983	A
26	1H	996	A
26	1H	1003	G
26	1H	1011	G
26	1H	1012	U
26	1H	1013	C
26	1H	1015	G
26	1H	1017	G
26	1H	1020	A
26	1H	1022	G
26	1H	1023	U
26	1H	1025	G
26	1H	1027	A
26	1H	1032	A
26	1H	1033	U
26	1H	1036	G
26	1H	1037	G
26	1H	1039	G
26	1H	1041	C
26	1H	1046	A
26	1H	1047	G
26	1H	1051	G
26	1H	1055	G
26	1H	1057	A
26	1H	1059	G
26	1H	1060	U
26	1H	1061	U

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Mol	Chain	Res	Type
26	1H	1062	G
26	1H	1063	G
26	1H	1064	C
26	1H	1066	U
26	1H	1068	G
26	1H	1070	A
26	1H	1071	G
26	1H	1072	C
26	1H	1074	G
26	1H	1076	C
26	1H	1077	A
26	1H	1078	U
26	1H	1080	A
26	1H	1081	U
26	1H	1082	U
26	1H	1084	A
26	1H	1085	A
26	1H	1086	A
26	1H	1087	G
26	1H	1088	A
26	1H	1090	U
26	1H	1093	G
26	1H	1095	A
26	1H	1096	A
26	1H	1097	U
26	1H	1104	C
26	1H	1111	A
26	1H	1112	G
26	1H	1121	C
26	1H	1126	A
26	1H	1127	A
26	1H	1128	A
26	1H	1129	A
26	1H	1130	U
26	1H	1135	C
26	1H	1136	G
26	1H	1139	G
26	1H	1142	U
26	1H	1142(A)	A
26	1H	1144	G
26	1H	1151	G
26	1H	1156	A

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Mol	Chain	Res	Type
26	1H	1169	G
26	1H	1170	G
26	1H	1173	G
26	1H	1174	A
26	1H	1175	U
26	1H	1176	G
26	1H	1177	A
26	1H	1178	C
26	1H	1179	C
26	1H	1190	G
26	1H	1194	A
26	1H	1195	G
26	1H	1200	C
26	1H	1204	A
26	1H	1205	U
26	1H	1218	C
26	1H	1220	A
26	1H	1221	C
26	1H	1225	C
26	1H	1240	U
26	1H	1241	A
26	1H	1244	G
26	1H	1248	G
26	1H	1253	A
26	1H	1256	G
26	1H	1265	A
26	1H	1267	U
26	1H	1269	A
26	1H	1271	G
26	1H	1272	A
26	1H	1273	U
26	1H	1274	A
26	1H	1284	A
26	1H	1287	A
26	1H	1292	U
26	1H	1300	U
26	1H	1301	A
26	1H	1305	C
26	1H	1310	G
26	1H	1313	U
26	1H	1314	C
26	1H	1321	A

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Mol	Chain	Res	Type
26	1H	1329	U
26	1H	1341	U
26	1H	1344	G
26	1H	1347	G
26	1H	1349	A
26	1H	1359	A
26	1H	1360	A
26	1H	1365	A
26	1H	1378	A
26	1H	1380	G
26	1H	1384	A
26	1H	1385	G
26	1H	1386	C
26	1H	1395	A
26	1H	1416	G
26	1H	1417	C
26	1H	1420	U
26	1H	1421	G
26	1H	1427	A
26	1H	1428	C
26	1H	1430	C
26	1H	1437	C
26	1H	1444(A)	A
26	1H	1449	A
26	1H	1453	A
26	1H	1454	U
26	1H	1455	G
26	1H	1458	C
26	1H	1459	G
26	1H	1460	A
26	1H	1461	G
26	1H	1467	C
26	1H	1471	A
26	1H	1478	G
26	1H	1483	G
26	1H	1493	C
26	1H	1497	U
26	1H	1506	C
26	1H	1508	A
26	1H	1509	C
26	1H	1510	A
26	1H	1511	A

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
26	1H	1517	G
26	1H	1520	U
26	1H	1522	G
26	1H	1534	G
26	1H	1535	U
26	1H	1536	A
26	1H	1537	C
26	1H	1538	G
26	1H	1543	A
26	1H	1544	C
26	1H	1545	A
26	1H	1554	A
26	1H	1558	A
26	1H	1559	G
26	1H	1566	A
26	1H	1569	A
26	1H	1578	U
26	1H	1580	A
26	1H	1585	C
26	1H	1586	A
26	1H	1587	A
26	1H	1608	A
26	1H	1609	A
26	1H	1610	A
26	1H	1616	A
26	1H	1617	C
26	1H	1626	G
26	1H	1647	G
26	1H	1648	C
26	1H	1651	G
26	1H	1654	A
26	1H	1672	C
26	1H	1674	G
26	1H	1675	C
26	1H	1699	G
26	1H	1706	U
26	1H	1729	A
26	1H	1730	U
26	1H	1746	G
26	1H	1756	G
26	1H	1763	G
26	1H	1764	G

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Mol	Chain	Res	Type
26	1H	1769	G
26	1H	1773	A
26	1H	1774	C
26	1H	1791	A
26	1H	1799	G
26	1H	1800	C
26	1H	1801	G
26	1H	1802	A
26	1H	1811	G
26	1H	1816	G
26	1H	1826	G
26	1H	1829	A
26	1H	1836	C
26	1H	1847	A
26	1H	1852	C
26	1H	1853	A
26	1H	1858	G
26	1H	1859	A
26	1H	1870	C
26	1H	1878	G
26	1H	1889	A
26	1H	1896	G
26	1H	1900	A
26	1H	1901	A
26	1H	1903	G
26	1H	1906	G
26	1H	1911	U
26	1H	1914	C
26	1H	1919	A
26	1H	1929	G
26	1H	1930	G
26	1H	1936	A
26	1H	1938	A
26	1H	1952	A
26	1H	1955	U
26	1H	1963	U
26	1H	1964	G
26	1H	1967	C
26	1H	1969	A
26	1H	1970	A
26	1H	1971	A
26	1H	1972	A

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
26	1H	1982	C
26	1H	1984	G
26	1H	1987	G
26	1H	1993	U
26	1H	2019	A
26	1H	2020	A
26	1H	2023	G
26	1H	2031	A
26	1H	2032	G
26	1H	2033	A
26	1H	2043	C
26	1H	2049	G
26	1H	2051	A
26	1H	2052	G
26	1H	2055	C
26	1H	2056	G
26	1H	2060	A
26	1H	2061	G
26	1H	2062	A
26	1H	2063	C
26	1H	2069	G
26	1H	2070	G
26	1H	2093	G
26	1H	2095	C
26	1H	2096	U
26	1H	2099	U
26	1H	2108	C
26	1H	2110	G
26	1H	2111	C
26	1H	2112	G
26	1H	2113	U
26	1H	2114	A
26	1H	2115	G
26	1H	2116	G
26	1H	2117	A
26	1H	2118	U
26	1H	2119	A
26	1H	2122	U
26	1H	2125	G
26	1H	2126	A
26	1H	2127	G
26	1H	2129	C

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
26	1H	2131	G
26	1H	2132	U
26	1H	2133	G
26	1H	2134	A
26	1H	2135	A
26	1H	2136	C
26	1H	2139	C
26	1H	2144	U
26	1H	2145	C
26	1H	2147	G
26	1H	2148	G
26	1H	2151	G
26	1H	2155	G
26	1H	2157	G
26	1H	2158	A
26	1H	2162	G
26	1H	2166	G
26	1H	2167	U
26	1H	2168	G
26	1H	2170	A
26	1H	2173	A
26	1H	2174	C
26	1H	2176	A
26	1H	2190	G
26	1H	2191	G
26	1H	2193	G
26	1H	2198	A
26	1H	2210	G
26	1H	2211	G
26	1H	2212	A
26	1H	2213	U
26	1H	2215	G
26	1H	2225	A
26	1H	2238	G
26	1H	2243	U
26	1H	2249	U
26	1H	2252	G
26	1H	2268	A
26	1H	2269	A
26	1H	2270	G
26	1H	2271	G
26	1H	2273	A

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Mol	Chain	Res	Type
26	1H	2275	C
26	1H	2279	G
26	1H	2280	G
26	1H	2283	C
26	1H	2287	A
26	1H	2288	A
26	1H	2298	A
26	1H	2299	G
26	1H	2305	A
26	1H	2307	G
26	1H	2308	G
26	1H	2309	A
26	1H	2310	A
26	1H	2314	C
26	1H	2315	G
26	1H	2319	G
26	1H	2320	A
26	1H	2325	G
26	1H	2327	A
26	1H	2334	G
26	1H	2336	A
26	1H	2343	C
26	1H	2347	C
26	1H	2350	C
26	1H	2351	G
26	1H	2352	A
26	1H	2364	C
26	1H	2376	A
26	1H	2377	A
26	1H	2383	G
26	1H	2385	C
26	1H	2388	A
26	1H	2406	U
26	1H	2410	G
26	1H	2424	C
26	1H	2425	A
26	1H	2428	G
26	1H	2429	G
26	1H	2430	A
26	1H	2434	A
26	1H	2439	A
26	1H	2440	C

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
26	1H	2441	C
26	1H	2446	G
26	1H	2448	A
26	1H	2457	U
26	1H	2468	G
26	1H	2469	A
26	1H	2470	G
26	1H	2475	C
26	1H	2476	A
26	1H	2478	A
26	1H	2480	C
26	1H	2482	G
26	1H	2484	G
26	1H	2489	G
26	1H	2494	G
26	1H	2502	G
26	1H	2504	U
26	1H	2505	G
26	1H	2506	U
26	1H	2507	C
26	1H	2513	G
26	1H	2518	A
26	1H	2529	G
26	1H	2537	U
26	1H	2545	G
26	1H	2551	C
26	1H	2554	U
26	1H	2556	C
26	1H	2564	A
26	1H	2566	A
26	1H	2567	G
26	1H	2569	G
26	1H	2573	C
26	1H	2574	G
26	1H	2582	G
26	1H	2602	A
26	1H	2605	U
26	1H	2609	U
26	1H	2611	U
26	1H	2612	C
26	1H	2615	U
26	1H	2629	A

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Mol	Chain	Res	Type
26	1H	2636	U
26	1H	2646	C
26	1H	2654	A
26	1H	2661	G
26	1H	2665	A
26	1H	2673	G
26	1H	2689	U
26	1H	2690	C
26	1H	2691	C
26	1H	2702	U
26	1H	2703	C
26	1H	2707	G
26	1H	2712(A)	A
26	1H	2713	A
26	1H	2714	G
26	1H	2718	G
26	1H	2719	G
26	1H	2726	U
26	1H	2733	A
26	1H	2734	A
26	1H	2736	G
26	1H	2744	G
26	1H	2752	C
26	1H	2756	U
26	1H	2757	A
26	1H	2764	A
26	1H	2765	A
26	1H	2766	G
26	1H	2778	A
26	1H	2779	U
26	1H	2787	C
26	1H	2789	C
26	1H	2791	C
26	1H	2793	G
26	1H	2794	C
26	1H	2795	G
26	1H	2797	U
26	1H	2798	C
26	1H	2801	A
26	1H	2803	C
26	1H	2807	G
26	1H	2808	U

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
26	1H	2818	G
26	1H	2820	A
26	1H	2821	A
26	1H	2825	C
26	1H	2832	U
26	1H	2833	G
26	1H	2834	G
26	1H	2835	A
26	1H	2860	A
26	1H	2872	G
26	1H	2876	G
26	1H	2877	G
26	1H	2879	C
26	1H	2892	A
26	1H	2893	G
26	1H	2895	U
26	1H	2899	G
26	1H	2900	A
27	16	0	A
27	16	13	A
27	16	15	A
27	16	24	G
27	16	25	A
27	16	27	C
27	16	33	G
27	16	38	C
27	16	40	U
27	16	41	U
27	16	45	A
27	16	54	G
27	16	56	G
27	16	58	A
27	16	73	A
27	16	77	U
27	16	85	G
27	16	105	G
27	16	107	U
27	16	109	G
27	16	115	G
1	1G	5	U
1	1G	9	G
1	1G	22	G

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Mol	Chain	Res	Type
1	1G	26	A
1	1G	32	A
1	1G	39	G
1	1G	47	C
1	1G	48	C
1	1G	50	A
1	1G	51	A
1	1G	54	C
1	1G	68	G
1	1G	76	G
1	1G	78	G
1	1G	79	G
1	1G	80	G
1	1G	81	G
1	1G	87	A
1	1G	88	C
1	1G	92	G
1	1G	99	C
1	1G	108	G
1	1G	113	G
1	1G	116	A
1	1G	121	C
1	1G	131	C
1	1G	144	G
1	1G	163	C
1	1G	169	C
1	1G	170	U
1	1G	173	U
1	1G	174	C
1	1G	182	U
1	1G	186	C
1	1G	186(F)	C
1	1G	187	C
1	1G	188	U
1	1G	189	U
1	1G	190	G
1	1G	191(A)	G
1	1G	191(D)	U
1	1G	195	A
1	1G	197	A
1	1G	198	G
1	1G	208	U

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Mol	Chain	Res	Type
1	1G	209	U
1	1G	210	U
1	1G	216	G
1	1G	220	G
1	1G	238	G
1	1G	240	C
1	1G	243	A
1	1G	244	U
1	1G	247	G
1	1G	250	A
1	1G	251	G
1	1G	262	A
1	1G	266	G
1	1G	267	C
1	1G	270	A
1	1G	281	G
1	1G	289	G
1	1G	298	A
1	1G	321	A
1	1G	328	C
1	1G	329	A
1	1G	332	G
1	1G	345	C
1	1G	346	G
1	1G	347	G
1	1G	350	G
1	1G	351	G
1	1G	352	C
1	1G	353	A
1	1G	354	G
1	1G	363	A
1	1G	365	U
1	1G	366	C
1	1G	367	U
1	1G	372	C
1	1G	373	A
1	1G	388	G
1	1G	397	A
1	1G	398	C
1	1G	406	G
1	1G	410	G
1	1G	411	A

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Mol	Chain	Res	Type
1	1G	412	A
1	1G	413	G
1	1G	414	A
1	1G	421	U
1	1G	422	C
1	1G	423	G
1	1G	424	G
1	1G	429	U
1	1G	430	A
1	1G	431	A
1	1G	439	A
1	1G	440	A
1	1G	451	A
1	1G	452	A
1	1G	466	C
1	1G	467	G
1	1G	475	G
1	1G	476	G
1	1G	482	A
1	1G	484	G
1	1G	485	G
1	1G	486	U
1	1G	495	A
1	1G	496	A
1	1G	497	U
1	1G	498	A
1	1G	505	G
1	1G	509	A
1	1G	510	A
1	1G	511	C
1	1G	518	C
1	1G	527	G
1	1G	530	G
1	1G	531	U
1	1G	532	A
1	1G	533	A
1	1G	536	C
1	1G	547	A
1	1G	552	U
1	1G	561	U
1	1G	572	A
1	1G	573	A

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Mol	Chain	Res	Type
1	1G	576	G
1	1G	577	G
1	1G	607	A
1	1G	608	A
1	1G	615	C
1	1G	618	C
1	1G	621	A
1	1G	632	A
1	1G	633	G
1	1G	651	C
1	1G	653	A
1	1G	661	G
1	1G	665	A
1	1G	671	G
1	1G	674	G
1	1G	687	A
1	1G	688	G
1	1G	701	C
1	1G	702	A
1	1G	703	G
1	1G	704	A
1	1G	722	A
1	1G	723	U
1	1G	724	G
1	1G	731	G
1	1G	749	C
1	1G	750	G
1	1G	755	G
1	1G	760	G
1	1G	762	C
1	1G	763	G
1	1G	764	C
1	1G	773	G
1	1G	774	G
1	1G	776	G
1	1G	777	A
1	1G	793	U
1	1G	794	A
1	1G	813	U
1	1G	817	C
1	1G	821	G
1	1G	828	A

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Mol	Chain	Res	Type
1	1G	834	C
1	1G	836	G
1	1G	842	C
1	1G	843	U
1	1G	848	C
1	1G	851	G
1	1G	853	G
1	1G	859	A
1	1G	873	A
1	1G	902	G
1	1G	906	G
1	1G	914	A
1	1G	916	G
1	1G	926	G
1	1G	927	G
1	1G	934	C
1	1G	935	A
1	1G	936	C
1	1G	938	A
1	1G	958	A
1	1G	960	U
1	1G	961	U
1	1G	966	G
1	1G	968	A
1	1G	969	A
1	1G	971	G
1	1G	972	C
1	1G	974	A
1	1G	975	A
1	1G	976	G
1	1G	977	A
1	1G	978	A
1	1G	980	C
1	1G	982	U
1	1G	989	C
1	1G	991	U
1	1G	992	U
1	1G	993	G
1	1G	1004	A
1	1G	1006	C
1	1G	1009	G
1	1G	1016	A

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Mol	Chain	Res	Type
1	1G	1019	C
1	1G	1021	G
1	1G	1023	G
1	1G	1024	G
1	1G	1025	U
1	1G	1026	G
1	1G	1027	C
1	1G	1028	C
1	1G	1028(B)	C
1	1G	1029	G
1	1G	1030	C
1	1G	1031	G
1	1G	1032(A)	G
1	1G	1033	G
1	1G	1038	C
1	1G	1040	U
1	1G	1043	C
1	1G	1046	A
1	1G	1053	G
1	1G	1054	C
1	1G	1055	A
1	1G	1056	U
1	1G	1064	G
1	1G	1066	C
1	1G	1080	A
1	1G	1081	G
1	1G	1082	G
1	1G	1088	G
1	1G	1094	G
1	1G	1095	U
1	1G	1096	C
1	1G	1101	A
1	1G	1108	G
1	1G	1118	C
1	1G	1124	G
1	1G	1125	U
1	1G	1127	G
1	1G	1128	C
1	1G	1129	C
1	1G	1131	G
1	1G	1135	U
1	1G	1137	C

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Mol	Chain	Res	Type
1	1G	1139	G
1	1G	1140	C
1	1G	1146	A
1	1G	1147	C
1	1G	1154	G
1	1G	1157	A
1	1G	1158	C
1	1G	1159	U
1	1G	1160	G
1	1G	1177	G
1	1G	1178	G
1	1G	1181	G
1	1G	1183	A
1	1G	1185	G
1	1G	1186	G
1	1G	1188	A
1	1G	1193	G
1	1G	1196	U
1	1G	1197	G
1	1G	1200	C
1	1G	1201	A
1	1G	1202	G
1	1G	1211	U
1	1G	1212	U
1	1G	1213	A
1	1G	1214	C
1	1G	1225	A
1	1G	1227	A
1	1G	1236	A
1	1G	1238	A
1	1G	1240	U
1	1G	1241	G
1	1G	1246	C
1	1G	1257	U
1	1G	1258	G
1	1G	1260	C
1	1G	1263	C
1	1G	1268	A
1	1G	1269	A
1	1G	1270	C
1	1G	1275	A
1	1G	1278	U

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Mol	Chain	Res	Type
1	1G	1279	A
1	1G	1280	A
1	1G	1285	A
1	1G	1286	A
1	1G	1287	A
1	1G	1288	A
1	1G	1295	G
1	1G	1297	C
1	1G	1298	C
1	1G	1299	A
1	1G	1300	G
1	1G	1301	U
1	1G	1305	G
1	1G	1306	A
1	1G	1312	G
1	1G	1317	C
1	1G	1318	A
1	1G	1319	A
1	1G	1320	C
1	1G	1322	C
1	1G	1323	G
1	1G	1331	G
1	1G	1335	C
1	1G	1336	C
1	1G	1337	G
1	1G	1340	A
1	1G	1346	A
1	1G	1347	G
1	1G	1353	G
1	1G	1354	C
1	1G	1362(A)	C
1	1G	1364	U
1	1G	1370	G
1	1G	1379	G
1	1G	1386	G
1	1G	1397	C
1	1G	1398	A
1	1G	1402	C
1	1G	1419	G
1	1G	1442	G
1	1G	1443	G
1	1G	1446	A

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Mol	Chain	Res	Type
1	1G	1447	G
1	1G	1449	C
1	1G	1450	U
1	1G	1451	A
1	1G	1452	C
1	1G	1453	G
1	1G	1454	G
1	1G	1482	G
1	1G	1492	A
1	1G	1494	G
1	1G	1497	G
1	1G	1498	U
1	1G	1499	A
1	1G	1502	A
1	1G	1503	A
1	1G	1504	G
1	1G	1506	U
1	1G	1507	A
1	1G	1517	G
1	1G	1519	A
1	1G	1520	G
1	1G	1525	G
1	1G	1528	U
1	1G	1529	G
1	1G	1530	G
1	1G	1533	C
1	1G	1535	C
1	1G	1536	C
1	1G	1537	U
1	1G	1540	U
1	1G	1541	U
22	1L	6	G
22	1L	8	U
22	1L	9	G
22	1L	16	U
22	1L	17	U
22	1L	18	G
22	1L	19	G
22	1L	20	U
22	1L	21	A
22	1L	22	G
22	1L	23	A

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Mol	Chain	Res	Type
22	1L	24	C
22	1L	26	C
22	1L	27	G
22	1L	30	A
22	1L	42	U
22	1L	46	G
22	1L	47	C
22	1L	48	C
22	1L	49	C
22	1L	50	A
22	1L	51	A
22	1L	52	U
22	1L	55	G
22	1L	58	U
22	1L	70	G
22	1L	74	C
22	1L	84	A
22	1L	85	C
22	1L	86	C
22	1L	87	A
23	2L	4	G
23	2L	6	G
23	2L	7	G
23	2L	8	U
23	2L	9	G
23	2L	11	U
23	2L	13	G
23	2L	15	A
23	2L	16	U
23	2L	17	H2U
23	2L	18	OMG
23	2L	19	G
23	2L	20	U
23	2L	21	A
23	2L	24	C
23	2L	27	G
23	2L	33	U
23	2L	34	U
23	2L	37	G
23	2L	44	G
23	2L	45	U
23	2L	46	G

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Mol	Chain	Res	Type
23	2L	47	C
23	2L	48	C
23	2L	57	C
23	2L	63	G
23	2L	64	G
23	2L	65	5MU
23	2L	69	A
23	2L	71	U
23	2L	74	C
23	2L	87	A
24	3L	5	A
24	3L	7	G
24	3L	8	U
24	3L	10	G
24	3L	11	U
24	3L	12	G
24	3L	14	A
24	3L	15	A
24	3L	16	U
24	3L	17	U
24	3L	18	G
24	3L	19	G
24	3L	20	U
24	3L	23	A
24	3L	27	G
24	3L	30	A
24	3L	32	C
24	3L	34	U
24	3L	35	G
24	3L	36	A
24	3L	38	G
24	3L	43	A
24	3L	44	G
24	3L	45	U
24	3L	58	U
24	3L	59	U
24	3L	60	A
24	3L	64	G
24	3L	69	A
24	3L	70	G
24	3L	71	U
24	3L	83	U

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Mol	Chain	Res	Type
24	3L	84	A
24	3L	85	C
25	4L	37	G
25	4L	38	U
25	4L	39	U
25	4L	50	U
25	4L	53	U
26	14	2	G
26	14	3	U
26	14	4	C
26	14	5	A
26	14	9	U
26	14	14	A
26	14	34	C
26	14	35	G
26	14	46	C
26	14	50	U
26	14	51	G
26	14	55	G
26	14	58	G
26	14	71	A
26	14	72	U
26	14	74	A
26	14	75	G
26	14	83	G
26	14	91	A
26	14	92	G
26	14	102	G
26	14	112	U
26	14	118	A
26	14	119	A
26	14	120	U
26	14	121	G
26	14	125	G
26	14	129	C
26	14	136	G
26	14	139	G
26	14	140	A
26	14	153	C
26	14	154	G
26	14	161	U
26	14	162	U

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Mol	Chain	Res	Type
26	14	172	C
26	14	173	G
26	14	175	G
26	14	196	A
26	14	199	A
26	14	205	G
26	14	214	G
26	14	215	G
26	14	216	A
26	14	221	A
26	14	222	A
26	14	223	A
26	14	225	A
26	14	228	A
26	14	229	A
26	14	233	A
26	14	245	G
26	14	248	G
26	14	249	C
26	14	252	G
26	14	266	G
26	14	270(K)	C
26	14	270(L)	U
26	14	270(M)	U
26	14	270(N)	G
26	14	270(O)	U
26	14	270(P)	C
26	14	271(B)	G
26	14	271	G
26	14	273(D)	C
26	14	273(E)	U
26	14	274	G
26	14	275	G
26	14	276	A
26	14	277	C
26	14	278	A
26	14	279	C
26	14	286	C
26	14	289	A
26	14	299	A
26	14	302	C
26	14	311	A

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Mol	Chain	Res	Type
26	14	324	A
26	14	327	G
26	14	329	G
26	14	330	A
26	14	333	G
26	14	352	G
26	14	356	G
26	14	363	G
26	14	363(E)	U
26	14	363(F)	A
26	14	372	G
26	14	386	G
26	14	395	U
26	14	396	G
26	14	405	U
26	14	406	G
26	14	411	G
26	14	412	A
26	14	428	A
26	14	443	A
26	14	444	C
26	14	448	U
26	14	451	C
26	14	454	A
26	14	455	C
26	14	456	C
26	14	457	A
26	14	459	U
26	14	470	A
26	14	471	A
26	14	481	G
26	14	501	A
26	14	504	U
26	14	505	A
26	14	508	G
26	14	509	C
26	14	528	A
26	14	529	A
26	14	530	G
26	14	531	C
26	14	532	A
26	14	533	G

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Mol	Chain	Res	Type
26	14	537	C
26	14	543	C
26	14	546	C
26	14	547	A
26	14	549	G
26	14	556	G
26	14	563	G
26	14	568	U
26	14	573	G
26	14	575	A
26	14	584	C
26	14	586	A
26	14	595	C
26	14	603	A
26	14	607	U
26	14	609(A)	G
26	14	613	U
26	14	614	U
26	14	615	G
26	14	617	G
26	14	618	G
26	14	621	A
26	14	622	G
26	14	627	A
26	14	637	A
26	14	646	A
26	14	651	G
26	14	654	A
26	14	654(D)	G
26	14	654(H)	G
26	14	654(I)	C
26	14	654(L)	G
26	14	654(N)	G
26	14	654(Q)	C
26	14	654(T)	A
26	14	668	G
26	14	669	G
26	14	677	A
26	14	686	G
26	14	701	G
26	14	711	G
26	14	717	G

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Mol	Chain	Res	Type
26	14	722	A
26	14	730	C
26	14	752	A
26	14	753	C
26	14	762	U
26	14	764	A
26	14	765	G
26	14	775	G
26	14	776	G
26	14	782	A
26	14	783	A
26	14	784	A
26	14	785	G
26	14	790	C
26	14	792	G
26	14	805	G
26	14	812	C
26	14	819	A
26	14	827	U
26	14	828	U
26	14	830	G
26	14	832	G
26	14	846	C
26	14	852	G
26	14	855	G
26	14	859	G
26	14	861	A
26	14	869	G
26	14	878	A
26	14	880	G
26	14	881	G
26	14	882	G
26	14	885	C
26	14	886	C
26	14	887	A
26	14	888	C
26	14	889	C
26	14	890	A
26	14	892	G
26	14	893	C
26	14	894	C
26	14	895	U

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Mol	Chain	Res	Type
26	14	896	A
26	14	897	C
26	14	899	A
26	14	900	A
26	14	901	A
26	14	904	C
26	14	910	A
26	14	911	A
26	14	914	C
26	14	915	C
26	14	917	A
26	14	918	A
26	14	919	G
26	14	924	C
26	14	932	G
26	14	933	A
26	14	938	G
26	14	941	A
26	14	945	A
26	14	946	G
26	14	958	U
26	14	959	A
26	14	961	C
26	14	965	C
26	14	974	G
26	14	978	G
26	14	980	A
26	14	982	C
26	14	983	A
26	14	989	G
26	14	990	A
26	14	991	C
26	14	996	A
26	14	1009	A
26	14	1010	A
26	14	1012	U
26	14	1013	C
26	14	1020	A
26	14	1022	G
26	14	1023	U
26	14	1025	G
26	14	1026	U

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
26	14	1039	G
26	14	1044	G
26	14	1045	A
26	14	1046	A
26	14	1048	A
26	14	1049	C
26	14	1051	G
26	14	1054	A
26	14	1057	A
26	14	1060	U
26	14	1061	U
26	14	1063	G
26	14	1065	U
26	14	1066	U
26	14	1067	A
26	14	1068	G
26	14	1070	A
26	14	1073	A
26	14	1077	A
26	14	1079	C
26	14	1082	U
26	14	1083	U
26	14	1085	A
26	14	1086	A
26	14	1087	G
26	14	1088	A
26	14	1089	G
26	14	1090	U
26	14	1091	G
26	14	1093	G
26	14	1095	A
26	14	1096	A
26	14	1099	G
26	14	1105	U
26	14	1111	A
26	14	1112	G
26	14	1122	G
26	14	1126	A
26	14	1127	A
26	14	1128	A
26	14	1129	A
26	14	1130	U

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
26	14	1135	C
26	14	1136	G
26	14	1139	G
26	14	1143	A
26	14	1151	G
26	14	1170	G
26	14	1171	G
26	14	1173	G
26	14	1174	A
26	14	1175	U
26	14	1177	A
26	14	1178	C
26	14	1186	G
26	14	1204	A
26	14	1205	U
26	14	1210	A
26	14	1212	G
26	14	1220	A
26	14	1236	G
26	14	1244	G
26	14	1253	A
26	14	1256	G
26	14	1265	A
26	14	1271	G
26	14	1272	A
26	14	1273	U
26	14	1278	A
26	14	1287	A
26	14	1298	C
26	14	1300	U
26	14	1301	A
26	14	1306	C
26	14	1307	A
26	14	1319	G
26	14	1321	A
26	14	1329	U
26	14	1332	G
26	14	1345	C
26	14	1349	A
26	14	1359	A
26	14	1365	A
26	14	1368	G

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Mol	Chain	Res	Type
26	14	1370	C
26	14	1380	G
26	14	1383	C
26	14	1384	A
26	14	1385	G
26	14	1386	C
26	14	1395	A
26	14	1407	C
26	14	1408	C
26	14	1416	G
26	14	1417	C
26	14	1419	A
26	14	1420	U
26	14	1421	G
26	14	1428	C
26	14	1437	C
26	14	1441	G
26	14	1444(A)	A
26	14	1445	C
26	14	1449	A
26	14	1449(A)	G
26	14	1455	G
26	14	1458	C
26	14	1459	G
26	14	1460	A
26	14	1461	G
26	14	1467	C
26	14	1471	A
26	14	1475	G
26	14	1480	G
26	14	1482	U
26	14	1483	G
26	14	1490	A
26	14	1493	C
26	14	1494	A
26	14	1508	A
26	14	1509	C
26	14	1510	A
26	14	1522	G
26	14	1529	A
26	14	1533	C
26	14	1534	G

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Mol	Chain	Res	Type
26	14	1535	U
26	14	1537	C
26	14	1543	A
26	14	1547	C
26	14	1554	A
26	14	1558	A
26	14	1559	G
26	14	1569	A
26	14	1577	C
26	14	1578	U
26	14	1585	C
26	14	1586	A
26	14	1587	A
26	14	1588	C
26	14	1598	C
26	14	1608	A
26	14	1609	A
26	14	1616	A
26	14	1631	A
26	14	1639	U
26	14	1644	C
26	14	1647	G
26	14	1648	C
26	14	1654	A
26	14	1673	U
26	14	1674	G
26	14	1675	C
26	14	1679	U
26	14	1680	U
26	14	1695	G
26	14	1697	G
26	14	1700	A
26	14	1701	A
26	14	1725	G
26	14	1726	G
26	14	1729	A
26	14	1730	U
26	14	1731	G
26	14	1732	A
26	14	1742	C
26	14	1743	G
26	14	1756	G

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Mol	Chain	Res	Type
26	14	1762	A
26	14	1763	G
26	14	1764	G
26	14	1769	G
26	14	1773	A
26	14	1777	U
26	14	1780	A
26	14	1782	C
26	14	1787	A
26	14	1791	A
26	14	1800	C
26	14	1801	G
26	14	1812	A
26	14	1816	G
26	14	1820	U
26	14	1826	G
26	14	1829	A
26	14	1847	A
26	14	1848	A
26	14	1858	G
26	14	1878	G
26	14	1884	A
26	14	1888	G
26	14	1889	A
26	14	1894	C
26	14	1900	A
26	14	1903	G
26	14	1906	G
26	14	1909	C
26	14	1913	A
26	14	1917	U
26	14	1920	C
26	14	1929	G
26	14	1930	G
26	14	1936	A
26	14	1937	A
26	14	1938	A
26	14	1955	U
26	14	1960	A
26	14	1963	U
26	14	1967	C
26	14	1969	A

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Mol	Chain	Res	Type
26	14	1970	A
26	14	1971	A
26	14	1972	A
26	14	1991	U
26	14	1992	G
26	14	1993	U
26	14	2005	A
26	14	2023	G
26	14	2025	C
26	14	2031	A
26	14	2032	G
26	14	2033	A
26	14	2043	C
26	14	2049	G
26	14	2051	A
26	14	2053	G
26	14	2054	A
26	14	2055	C
26	14	2056	G
26	14	2057	A
26	14	2059	A
26	14	2060	A
26	14	2061	G
26	14	2062	A
26	14	2063	C
26	14	2069	G
26	14	2074	U
26	14	2076	U
26	14	2093	G
26	14	2100	G
26	14	2108	C
26	14	2111	C
26	14	2112	G
26	14	2113	U
26	14	2114	A
26	14	2115	G
26	14	2116	G
26	14	2117	A
26	14	2120	G
26	14	2122	U
26	14	2123	G
26	14	2125	G

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>
26	14	2126	A
26	14	2127	G
26	14	2129	C
26	14	2131	G
26	14	2132	U
26	14	2133	G
26	14	2136	C
26	14	2137	C
26	14	2138	C
26	14	2139	C
26	14	2140	C
26	14	2144	U
26	14	2145	C
26	14	2146	C
26	14	2147	G
26	14	2148	G
26	14	2151	G
26	14	2155	G
26	14	2158	A
26	14	2160	G
26	14	2161	C
26	14	2162	G
26	14	2164	C
26	14	2166	G
26	14	2167	U
26	14	2168	G
26	14	2169	A
26	14	2171	A
26	14	2173	A
26	14	2174	C
26	14	2182	G
26	14	2189	U
26	14	2191	G
26	14	2192	G
26	14	2198	A
26	14	2210	G
26	14	2211	G
26	14	2212	A
26	14	2213	U
26	14	2215	G
26	14	2225	A
26	14	2226	C

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Mol	Chain	Res	Type
26	14	2228	G
26	14	2239	G
26	14	2245	U
26	14	2261	C
26	14	2268	A
26	14	2269	A
26	14	2275	C
26	14	2276	G
26	14	2278	A
26	14	2280	G
26	14	2283	C
26	14	2287	A
26	14	2288	A
26	14	2305	A
26	14	2307	G
26	14	2308	G
26	14	2310	A
26	14	2311	A
26	14	2317	C
26	14	2318	G
26	14	2321	G
26	14	2325	G
26	14	2334	G
26	14	2335	A
26	14	2336	A
26	14	2345	G
26	14	2346	A
26	14	2347	C
26	14	2350	C
26	14	2354	G
26	14	2355	C
26	14	2383	G
26	14	2385	C
26	14	2389	G
26	14	2392	A
26	14	2396	G
26	14	2400	G
26	14	2401	U
26	14	2402	C
26	14	2403	C
26	14	2406	U
26	14	2410	G

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Mol	Chain	Res	Type
26	14	2414	G
26	14	2422	A
26	14	2423	U
26	14	2424	C
26	14	2425	A
26	14	2428	G
26	14	2429	G
26	14	2430	A
26	14	2431	U
26	14	2434	A
26	14	2435	A
26	14	2439	A
26	14	2441	C
26	14	2448	A
26	14	2464	C
26	14	2469	A
26	14	2470	G
26	14	2475	C
26	14	2476	A
26	14	2477	C
26	14	2478	A
26	14	2480	C
26	14	2482	G
26	14	2483	C
26	14	2484	G
26	14	2494	G
26	14	2495	G
26	14	2502	G
26	14	2504	U
26	14	2505	G
26	14	2506	U
26	14	2513	G
26	14	2518	A
26	14	2525	G
26	14	2532	G
26	14	2543	G
26	14	2554	U
26	14	2566	A
26	14	2567	G
26	14	2569	G
26	14	2573	C
26	14	2585	U

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Mol	Chain	Res	Type
26	14	2591	C
26	14	2599	G
26	14	2601	C
26	14	2602	A
26	14	2609	U
26	14	2610	C
26	14	2611	U
26	14	2612	C
26	14	2613	U
26	14	2615	U
26	14	2630	G
26	14	2636	U
26	14	2641	G
26	14	2646	C
26	14	2654	A
26	14	2660	A
26	14	2664	G
26	14	2665	A
26	14	2667	C
26	14	2673	G
26	14	2679	A
26	14	2689	U
26	14	2690	C
26	14	2698	U
26	14	2702	U
26	14	2703	C
26	14	2707	G
26	14	2712(A)	A
26	14	2713	A
26	14	2714	G
26	14	2726	U
26	14	2733	A
26	14	2739	U
26	14	2744	G
26	14	2748	A
26	14	2750	A
26	14	2751	G
26	14	2752	C
26	14	2754	U
26	14	2757	A
26	14	2758	A
26	14	2759	G

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Mol	Chain	Res	Type
26	14	2762	G
26	14	2764	A
26	14	2765	A
26	14	2766	G
26	14	2777	G
26	14	2778	A
26	14	2779	U
26	14	2780	G
26	14	2790	A
26	14	2791	C
26	14	2793	G
26	14	2797	U
26	14	2798	C
26	14	2799	A
26	14	2805	G
26	14	2810	A
26	14	2818	G
26	14	2820	A
26	14	2821	A
26	14	2833	G
26	14	2835	A
26	14	2849	U
26	14	2860	A
26	14	2861	G
26	14	2872	G
26	14	2873	A
26	14	2879	C
26	14	2894	G
26	14	2896	C
26	14	2898	U
26	14	2899	G
27	1J	0	A
27	1J	2	C
27	1J	5	C
27	1J	7	G
27	1J	8	U
27	1J	9	G
27	1J	12	C
27	1J	13	A
27	1J	15	A
27	1J	16	G
27	1J	19	G

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Mol	Chain	Res	Type
27	1J	22	U
27	1J	25	A
27	1J	29	A
27	1J	34	U
27	1J	40	U
27	1J	41	U
27	1J	42	C
27	1J	45	A
27	1J	53	A
27	1J	56	G
27	1J	58	A
27	1J	73	A
27	1J	81	G
27	1J	82	G
27	1J	88	C
27	1J	89	G
27	1J	89(A)	A
27	1J	90	C
27	1J	91	C
27	1J	99	A
27	1J	101	A
27	1J	108	C
27	1J	109	G
27	1J	116	G
27	1J	118	G

All (196) RNA pucker outliers are listed below:

Mol	Chain	Res	Type
1	13	5	U
1	13	50	A
1	13	115	G
1	13	181	G
1	13	189	U
1	13	244	U
1	13	251	G
1	13	266	G
1	13	412	A
1	13	429	U
1	13	484	G
1	13	560	U
1	13	748	C

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Mol	Chain	Res	Type
1	13	793	U
1	13	992	U
1	13	1025	U
1	13	1027	C
1	13	1053	G
1	13	1064	G
1	13	1065	U
1	13	1129	C
1	13	1145	C
1	13	1177	G
1	13	1285	A
1	13	1302	U
1	13	1336	C
1	13	1397	C
1	13	1452	C
1	13	1498	U
1	13	1503	A
1	13	1504	G
22	1K	18	G
23	2K	8	U
23	2K	18	OMG
23	2K	20	U
23	2K	36	A
23	2K	44	G
23	2K	62	G
23	2K	63	G
23	2K	73	C
24	3K	18	G
24	3K	46	G
24	3K	59	U
24	3K	69	A
26	1H	33	U
26	1H	196	A
26	1H	232	G
26	1H	271(B)	G
26	1H	334	C
26	1H	404	C
26	1H	481	G
26	1H	508	G
26	1H	528	A
26	1H	587	C
26	1H	668	G

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Mol	Chain	Res	Type
26	1H	746	A
26	1H	752	A
26	1H	764	A
26	1H	800	A
26	1H	845	G
26	1H	858	U
26	1H	859	G
26	1H	880	G
26	1H	1022	G
26	1H	1026	U
26	1H	1060	U
26	1H	1085	A
26	1H	1110	G
26	1H	1178	C
26	1H	1273	U
26	1H	1312	U
26	1H	1379	A
26	1H	1396	U
26	1H	1416	G
26	1H	1508	A
26	1H	1558	A
26	1H	1608	A
26	1H	1609	A
26	1H	1647	G
26	1H	1729	A
26	1H	1762	A
26	1H	1799	G
26	1H	1858	G
26	1H	1899	G
26	1H	1900	A
26	1H	2060	A
26	1H	2157	G
26	1H	2242	G
26	1H	2272	U
26	1H	2475	C
26	1H	2481	G
26	1H	2518	A
26	1H	2566	A
26	1H	2611	U
26	1H	2689	U
26	1H	2756	U
27	16	108	C

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Mol	Chain	Res	Type
1	1G	86	U
1	1G	115	G
1	1G	197	A
1	1G	243	A
1	1G	250	A
1	1G	266	G
1	1G	327	A
1	1G	345	C
1	1G	412	A
1	1G	413	G
1	1G	429	U
1	1G	485	G
1	1G	509	A
1	1G	528	C
1	1G	560	U
1	1G	632	A
1	1G	687	A
1	1G	722	A
1	1G	723	U
1	1G	748	C
1	1G	812	C
1	1G	913	A
1	1G	992	U
1	1G	1025	U
1	1G	1053	G
1	1G	1094	G
1	1G	1126	U
1	1G	1128	C
1	1G	1145	C
1	1G	1157	A
1	1G	1285	A
1	1G	1297	C
1	1G	1298	C
1	1G	1300	G
1	1G	1305	G
1	1G	1346	A
1	1G	1453	G
1	1G	1498	U
22	1L	20	U
22	1L	21	A
22	1L	22	G
22	1L	48	C

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Mol	Chain	Res	Type
23	2L	8	U
23	2L	20	U
23	2L	32	C
23	2L	36	A
23	2L	44	G
23	2L	56	G
23	2L	63	G
23	2L	73	C
24	3L	17	U
24	3L	18	G
24	3L	59	U
24	3L	83	U
24	3L	84	A
26	14	4	C
26	14	34	C
26	14	49	A
26	14	101	G
26	14	128	C
26	14	278	A
26	14	456	C
26	14	503	A
26	14	545	G
26	14	669	G
26	14	752	A
26	14	764	A
26	14	774	A
26	14	776	G
26	14	827	U
26	14	1022	G
26	14	1085	A
26	14	1379	A
26	14	1396	U
26	14	1420	U
26	14	1558	A
26	14	1608	A
26	14	2135	A
26	14	2157	G
26	14	2210	G
26	14	2225	A
26	14	2275	C
26	14	2402	C
26	14	2406	U

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Mol	Chain	Res	Type
26	14	2423	U
26	14	2542	A
26	14	2612	C
26	14	2629	A
26	14	2689	U
26	14	2747	G
26	14	2756	U
26	14	2776	A
26	14	2778	A
26	14	2859	G
26	14	2893	G
27	1J	15	A
27	1J	44	G
27	1J	88	C
27	1J	89	G

## 5.4 Non-standard residues in protein, DNA, RNA chains ⓘ

20 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$
22	5MU	1L	65	22	15,22,23	2.11	3 (20%)	16,32,35	1.75	3 (18%)
22	5MU	1K	65	22	15,22,23	2.19	3 (20%)	16,32,35	1.81	2 (12%)
22	PSU	1K	66	22	17,21,22	1.11	1 (5%)	20,30,33	3.28	6 (30%)
23	5MU	2L	65	23	15,22,23	2.11	3 (20%)	16,32,35	1.68	2 (12%)
23	OMG	2K	18	23	18,26,27	5.86	6 (33%)	20,38,41	5.32	7 (35%)
22	1MG	1K	38	22	18,26,27	3.62	5 (27%)	19,39,42	1.79	3 (15%)
23	PSU	2K	66	23	17,21,22	1.17	1 (5%)	20,30,33	3.50	4 (20%)
22	PSU	1K	39	22	17,21,22	1.00	1 (5%)	20,30,33	3.37	6 (30%)
22	PSU	1L	66	22	17,21,22	0.95	1 (5%)	20,30,33	3.30	6 (30%)
23	PSU	2L	39	23	17,21,22	1.03	1 (5%)	20,30,33	3.26	7 (35%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z  > 2	Counts	RMSZ	# Z  > 2
23	H2U	2K	17	23	18,21,22	2.19	4 (22%)	21,30,33	2.49	5 (23%)
23	H2U	2L	17	23	18,21,22	1.93	4 (22%)	21,30,33	2.28	5 (23%)
23	PSU	2K	39	23	17,21,22	0.97	1 (5%)	20,30,33	3.16	6 (30%)
23	OMG	2L	18	23	18,26,27	5.83	7 (38%)	20,38,41	5.56	8 (40%)
22	PSU	1L	39	22	17,21,22	1.09	2 (11%)	20,30,33	3.45	6 (30%)
23	PSU	2L	66	23	17,21,22	1.09	1 (5%)	20,30,33	3.27	5 (25%)
22	1MG	1L	38	22	18,26,27	3.65	5 (27%)	19,39,42	1.64	2 (10%)
23	1MG	2L	38	23	18,26,27	3.82	5 (27%)	19,39,42	1.92	3 (15%)
23	1MG	2K	38	23	18,26,27	3.29	5 (27%)	19,39,42	1.62	3 (15%)
23	5MU	2K	65	23	15,22,23	2.16	3 (20%)	16,32,35	1.82	2 (12%)

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
22	5MU	1L	65	22	-	0/5/25/26	0/2/2/2
22	5MU	1K	65	22	-	0/5/25/26	0/2/2/2
22	PSU	1K	66	22	-	0/7/25/26	0/2/2/2
23	5MU	2L	65	23	-	3/5/25/26	0/2/2/2
23	OMG	2K	18	23	-	0/5/27/28	0/3/3/3
22	1MG	1K	38	22	-	2/3/25/26	0/3/3/3
23	PSU	2K	66	23	-	0/7/25/26	0/2/2/2
22	PSU	1K	39	22	-	1/7/25/26	0/2/2/2
22	PSU	1L	66	22	-	0/7/25/26	0/2/2/2
23	PSU	2L	39	23	-	0/7/25/26	0/2/2/2
23	H2U	2K	17	23	-	1/7/38/39	0/2/2/2
23	H2U	2L	17	23	-	2/7/38/39	0/2/2/2
23	PSU	2K	39	23	-	0/7/25/26	0/2/2/2
23	OMG	2L	18	23	-	3/5/27/28	0/3/3/3
22	PSU	1L	39	22	-	0/7/25/26	0/2/2/2
23	PSU	2L	66	23	-	0/7/25/26	0/2/2/2
22	1MG	1L	38	22	-	0/3/25/26	0/3/3/3
23	1MG	2L	38	23	-	0/3/25/26	0/3/3/3
23	1MG	2K	38	23	-	0/3/25/26	0/3/3/3
23	5MU	2K	65	23	-	0/5/25/26	0/2/2/2

All (62) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
23	2K	18	OMG	C4-N3	17.43	1.63	1.35
23	2L	18	OMG	C4-N3	16.52	1.61	1.35
23	2L	18	OMG	C8-N7	-14.82	1.08	1.34
23	2K	18	OMG	C8-N7	-14.18	1.09	1.34
23	2L	38	1MG	C2-N3	8.76	1.47	1.34
22	1K	38	1MG	C2-N3	8.42	1.47	1.34
22	1L	38	1MG	C2-N3	8.41	1.47	1.34
23	2L	38	1MG	C4-N3	8.32	1.48	1.35
23	2K	18	OMG	C5-C4	8.23	1.62	1.40
23	2L	18	OMG	C5-C4	8.17	1.62	1.40
23	2L	38	1MG	C6-C5	8.15	1.54	1.41
22	1L	38	1MG	C4-N3	7.98	1.48	1.35
23	2K	38	1MG	C2-N3	7.78	1.46	1.34
22	1L	38	1MG	C6-C5	7.74	1.53	1.41
22	1K	38	1MG	C6-C5	7.70	1.53	1.41
22	1K	38	1MG	C4-N3	7.51	1.47	1.35
23	2K	38	1MG	C6-C5	7.21	1.52	1.41
23	2K	17	H2U	C2-N1	6.94	1.45	1.35
23	2K	38	1MG	C4-N3	6.71	1.46	1.35
22	1K	65	5MU	C4-C5	5.75	1.53	1.41
23	2K	65	5MU	C4-C5	5.54	1.53	1.41
23	2L	65	5MU	C4-C5	5.44	1.53	1.41
23	2L	17	H2U	C2-N1	5.39	1.43	1.35
23	2L	38	1MG	C2-N2	5.30	1.44	1.33
22	1K	38	1MG	C2-N2	5.17	1.44	1.33
22	1L	65	5MU	C4-C5	5.17	1.52	1.41
23	2K	38	1MG	C2-N2	5.10	1.44	1.33
22	1L	38	1MG	C2-N2	5.06	1.44	1.33
22	1K	65	5MU	C2-N3	5.05	1.48	1.38
22	1L	65	5MU	C2-N3	4.89	1.47	1.38
23	2K	65	5MU	C2-N3	4.87	1.47	1.38
23	2L	65	5MU	C2-N3	4.78	1.47	1.38
23	2L	18	OMG	C6-N1	-4.59	1.25	1.33
23	2K	18	OMG	C6-N1	-4.37	1.25	1.33
23	2L	18	OMG	C2-N2	3.99	1.41	1.33
23	2L	38	1MG	C6-N1	3.77	1.43	1.38
23	2L	17	H2U	C2-N3	3.72	1.44	1.38
23	2K	17	H2U	C2-N3	3.71	1.44	1.38
23	2K	66	PSU	C4-N3	3.57	1.39	1.33
23	2L	66	PSU	C4-N3	3.51	1.39	1.33
23	2K	18	OMG	C2-N2	3.49	1.40	1.33
22	1K	38	1MG	C6-N1	3.48	1.43	1.38
22	1L	65	5MU	C4-N3	-3.43	1.27	1.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	1L	38	1MG	C6-N1	3.42	1.43	1.38
23	2K	17	H2U	C4-N3	3.40	1.43	1.37
22	1K	66	PSU	C4-N3	3.33	1.38	1.33
23	2L	17	H2U	C4-N3	3.26	1.43	1.37
23	2L	39	PSU	C4-N3	3.23	1.38	1.33
23	2K	65	5MU	C4-N3	-3.21	1.27	1.33
22	1L	66	PSU	C4-N3	2.99	1.38	1.33
23	2L	65	5MU	C4-N3	-2.93	1.27	1.33
23	2K	39	PSU	C4-N3	2.93	1.38	1.33
22	1L	39	PSU	C4-N3	2.90	1.38	1.33
22	1K	65	5MU	C4-N3	-2.83	1.28	1.33
23	2L	17	H2U	C6-N1	-2.83	1.42	1.47
22	1K	39	PSU	C4-N3	2.60	1.37	1.33
23	2K	17	H2U	C6-N1	-2.57	1.42	1.47
23	2L	18	OMG	O6-C6	-2.52	1.18	1.24
23	2K	18	OMG	O6-C6	-2.39	1.18	1.24
23	2L	18	OMG	C6-C5	-2.19	1.37	1.41
23	2K	38	1MG	C6-N1	2.06	1.41	1.38
22	1L	39	PSU	C5-C1'	2.02	1.54	1.52

All (91) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
23	2L	18	OMG	C1'-N9-C4	14.85	152.73	126.64
23	2L	18	OMG	C6-C5-C4	-14.07	107.37	120.80
23	2K	18	OMG	C6-C5-C4	-13.58	107.84	120.80
23	2K	18	OMG	C1'-N9-C4	12.98	149.45	126.64
22	1L	39	PSU	N1-C2-N3	-12.17	118.75	128.43
22	1K	39	PSU	N1-C2-N3	-11.87	118.99	128.43
22	1L	66	PSU	N1-C2-N3	-10.98	119.70	128.43
23	2K	39	PSU	N1-C2-N3	-10.87	119.79	128.43
23	2L	39	PSU	N1-C2-N3	-10.68	119.94	128.43
23	2L	66	PSU	N1-C2-N3	-10.67	119.94	128.43
22	1K	66	PSU	N1-C2-N3	-10.59	120.01	128.43
23	2K	66	PSU	N1-C2-N3	-10.51	120.07	128.43
23	2L	18	OMG	C6-N1-C2	9.54	131.09	115.93
23	2K	18	OMG	C6-N1-C2	9.42	130.90	115.93
23	2K	17	H2U	C4-N3-C2	-8.09	119.08	125.79
23	2K	66	PSU	C4-N3-C2	7.80	121.72	115.14
23	2L	17	H2U	C4-N3-C2	-7.43	119.63	125.79
23	2K	18	OMG	C4-C5-N7	-7.00	102.10	109.40
22	1L	39	PSU	C4-N3-C2	6.92	120.99	115.14

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
23	2L	66	PSU	C4-N3-C2	6.84	120.92	115.14
23	2L	18	OMG	N3-C2-N1	-6.80	118.16	127.22
23	2K	18	OMG	N3-C2-N1	-6.51	118.54	127.22
22	1K	66	PSU	C4-N3-C2	6.31	120.47	115.14
23	2L	38	1MG	C1'-N9-C4	-6.30	115.57	126.64
22	1K	39	PSU	C4-N3-C2	6.28	120.45	115.14
22	1K	38	1MG	C1'-N9-C4	-5.97	116.15	126.64
22	1L	66	PSU	C4-N3-C2	5.89	120.12	115.14
23	2K	66	PSU	C5-C4-N3	-5.87	117.80	125.36
22	1L	38	1MG	C1'-N9-C4	-5.80	116.46	126.64
23	2K	17	H2U	C5-C4-N3	5.25	122.54	116.65
23	2L	66	PSU	C5-C4-N3	-5.19	118.67	125.36
23	2L	18	OMG	C4-C5-N7	-5.11	104.08	109.40
23	2K	39	PSU	C4-N3-C2	5.05	119.41	115.14
22	1K	65	5MU	C4-N3-C2	4.97	119.34	115.14
23	2L	39	PSU	C4-N3-C2	4.94	119.31	115.14
22	1K	66	PSU	C5-C4-N3	-4.90	119.05	125.36
23	2K	65	5MU	C4-N3-C2	4.87	119.25	115.14
23	2K	38	1MG	C1'-N9-C4	-4.84	118.13	126.64
23	2K	65	5MU	C5-C6-N1	-4.79	117.03	122.19
22	1L	65	5MU	C4-N3-C2	4.78	119.18	115.14
23	2L	65	5MU	C4-N3-C2	4.78	119.17	115.14
23	2K	66	PSU	C5-C1'-C2'	-4.76	106.82	115.32
22	1K	65	5MU	C5-C6-N1	-4.75	117.08	122.19
23	2L	18	OMG	C5-C6-N1	4.52	129.62	123.43
22	1L	66	PSU	C5-C4-N3	-4.40	119.69	125.36
22	1L	65	5MU	C5-C6-N1	-4.32	117.53	122.19
23	2L	17	H2U	C5-C4-N3	4.31	121.49	116.65
23	2K	18	OMG	C5-C6-N1	4.26	129.25	123.43
23	2L	38	1MG	C2-N3-C4	3.99	119.92	115.36
23	2L	65	5MU	C5-C6-N1	-3.98	117.90	122.19
23	2L	39	PSU	C5-C4-N3	-3.92	120.31	125.36
23	2L	39	PSU	C5-C1'-C2'	-3.81	108.52	115.32
23	2K	17	H2U	N3-C2-N1	3.80	120.67	116.65
23	2L	17	H2U	N3-C2-N1	3.76	120.63	116.65
23	2K	39	PSU	C5-C4-N3	-3.76	120.52	125.36
23	2K	38	1MG	C2-N3-C4	3.67	119.55	115.36
23	2L	39	PSU	C5-C6-N1	-3.58	120.03	124.44
22	1L	39	PSU	C5-C4-N3	-3.53	120.81	125.36
23	2L	39	PSU	C6-N1-C2	3.51	121.15	115.36
22	1K	38	1MG	C2-N3-C4	3.51	119.36	115.36
22	1K	39	PSU	C6-N1-C2	3.46	121.08	115.36

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	1K	39	PSU	C5-C4-N3	-3.44	120.92	125.36
23	2K	39	PSU	C6-N1-C2	3.43	121.02	115.36
22	1K	66	PSU	C5-C1'-C2'	-3.38	109.28	115.32
22	1L	66	PSU	C6-N1-C2	3.36	120.91	115.36
23	2K	39	PSU	C5-C6-N1	-3.34	120.33	124.44
22	1L	66	PSU	C5-C1'-C2'	-3.23	109.56	115.32
22	1L	66	PSU	C5-C6-N1	-3.20	120.50	124.44
23	2L	39	PSU	O4'-C1'-C5	3.07	114.69	109.93
22	1L	39	PSU	C6-N1-C2	3.05	120.39	115.36
22	1L	38	1MG	C2-N3-C4	3.03	118.82	115.36
23	2K	39	PSU	C5-C1'-C2'	-2.99	109.99	115.32
22	1K	39	PSU	C5-C6-N1	-2.92	120.85	124.44
23	2K	17	H2U	C5-C6-N1	2.90	121.17	111.61
22	1K	66	PSU	C5-C6-N1	-2.85	120.94	124.44
22	1K	66	PSU	C6-N1-C2	2.83	120.03	115.36
23	2L	66	PSU	C6-N1-C2	2.82	120.01	115.36
23	2K	18	OMG	N2-C2-N1	2.80	121.61	117.25
23	2K	38	1MG	N2-C2-N3	2.76	121.88	117.40
23	2L	66	PSU	C5-C6-N1	-2.72	121.10	124.44
23	2L	17	H2U	C5-C6-N1	2.71	120.56	111.61
22	1L	39	PSU	O4'-C1'-C5	2.55	113.87	109.93
23	2L	38	1MG	N2-C2-N3	2.49	121.43	117.40
23	2L	17	H2U	O2-C2-N1	-2.48	119.99	123.11
23	2L	18	OMG	N2-C2-N1	2.19	120.66	117.25
22	1K	39	PSU	C5-C1'-C2'	-2.18	111.43	115.32
22	1L	39	PSU	C5-C6-N1	-2.14	121.81	124.44
22	1K	38	1MG	N2-C2-N3	2.13	120.86	117.40
23	2K	17	H2U	O2-C2-N3	-2.11	117.57	121.50
22	1L	65	5MU	C6-N1-C1'	2.10	123.96	119.24
23	2L	18	OMG	N2-C2-N3	2.08	121.18	117.79

There are no chirality outliers.

All (12) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
23	2L	18	OMG	C4'-C5'-O5'-P
23	2L	18	OMG	O4'-C4'-C5'-O5'
23	2L	18	OMG	C3'-C4'-C5'-O5'
23	2L	65	5MU	C3'-C4'-C5'-O5'
23	2L	65	5MU	O4'-C4'-C5'-O5'
22	1K	38	1MG	O4'-C4'-C5'-O5'
22	1K	38	1MG	C3'-C4'-C5'-O5'

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Mol	Chain	Res	Type	Atoms
23	2L	17	H2U	O4'-C4'-C5'-O5'
23	2L	65	5MU	C4'-C5'-O5'-P
22	1K	39	PSU	C2'-C1'-C5-C6
23	2K	17	H2U	O4'-C4'-C5'-O5'
23	2L	17	H2U	C3'-C4'-C5'-O5'

There are no ring outliers.

10 monomers are involved in 13 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
22	1L	65	5MU	2	0
23	2L	65	5MU	2	0
23	2K	18	OMG	2	0
22	1K	39	PSU	1	0
23	2L	39	PSU	2	0
23	2K	17	H2U	1	0
23	2L	18	OMG	2	0
23	2L	66	PSU	2	0
23	2L	38	1MG	2	0
23	2K	38	1MG	1	0

## 5.5 Carbohydrates [i](#)

There are no carbohydrates in this entry.

## 5.6 Ligand geometry [i](#)

Of 1340 ligands modelled in this entry, 1340 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
25	4K	1
25	4L	1

All chain breaks are listed below:

Model	Chain	Residue-1	Atom-1	Residue-2	Atom-2	Distance (Å)
1	4K	38:U	O3'	44:U	P	28.59
1	4L	40:U	O3'	44:U	P	16.09

## 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, 95<sup>th</sup> 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 <sup>2</sup> )	Q<0.9
1	13	1507/1522 (99%)	-0.64	0 100 100	70, 113, 194, 267	0
1	1G	1513/1522 (99%)	-0.67	1 (0%) 95 97	79, 120, 189, 288	0
2	12	237/256 (92%)	0.57	25 (10%) 6 6	133, 170, 199, 207	0
2	1E	237/256 (92%)	0.36	15 (6%) 20 20	116, 156, 187, 198	0
3	22	206/239 (86%)	1.52	68 (33%) 0 0	128, 150, 185, 189	0
3	2E	205/239 (85%)	0.66	20 (9%) 7 7	94, 120, 157, 169	0
4	32	208/209 (99%)	0.80	20 (9%) 8 8	100, 121, 146, 158	0
4	3E	208/209 (99%)	0.23	3 (1%) 75 75	91, 113, 135, 145	0
5	42	151/162 (93%)	0.52	11 (7%) 15 15	105, 124, 142, 169	0
5	4E	151/162 (93%)	0.36	7 (4%) 32 30	89, 108, 128, 159	0
6	52	101/101 (100%)	0.27	0 100 100	87, 107, 126, 138	0
6	5E	101/101 (100%)	0.55	6 (5%) 22 22	87, 109, 129, 140	0
7	62	147/156 (94%)	0.51	11 (7%) 14 13	123, 138, 159, 171	0
7	6E	144/156 (92%)	0.40	4 (2%) 53 51	118, 136, 154, 175	0
8	72	138/138 (100%)	0.70	13 (9%) 8 9	100, 125, 141, 147	0
8	7E	138/138 (100%)	0.45	6 (4%) 35 34	97, 116, 130, 141	0
9	82	124/128 (96%)	0.31	4 (3%) 47 46	112, 154, 172, 181	0
9	8E	127/128 (99%)	-0.02	2 (1%) 72 70	105, 145, 167, 173	0
10	1A	99/105 (94%)	0.94	16 (16%) 1 2	126, 158, 184, 196	0
10	1I	99/105 (94%)	1.07	17 (17%) 1 1	96, 144, 175, 179	0
11	2A	116/129 (89%)	1.14	19 (16%) 1 2	92, 114, 138, 155	0
11	2I	116/129 (89%)	0.96	22 (18%) 1 1	86, 117, 142, 184	0
12	3A	125/132 (94%)	1.18	35 (28%) 0 0	91, 108, 136, 189	0
12	3I	122/132 (92%)	0.49	5 (4%) 37 35	78, 86, 112, 144	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å <sup>2</sup> )	Q<0.9
13	4A	116/126 (92%)	0.98	18 (15%) 2 2	128, 165, 190, 202	0
13	4I	116/126 (92%)	0.96	18 (15%) 2 2	106, 146, 164, 175	0
14	5A	58/61 (95%)	1.29	21 (36%) 0 0	131, 143, 165, 167	0
14	5I	60/61 (98%)	0.02	1 (1%) 70 68	98, 110, 131, 142	0
15	6A	88/89 (98%)	0.26	4 (4%) 33 32	90, 113, 134, 136	0
15	6I	88/89 (98%)	0.34	4 (4%) 33 32	87, 108, 130, 139	0
16	7A	84/88 (95%)	0.42	4 (4%) 30 28	93, 108, 135, 169	0
16	7I	84/88 (95%)	0.82	15 (17%) 1 1	108, 120, 155, 187	0
17	8A	99/105 (94%)	0.52	9 (9%) 9 9	98, 112, 126, 131	0
17	8I	100/105 (95%)	0.12	3 (3%) 50 49	95, 113, 129, 133	0
18	9A	69/88 (78%)	1.37	16 (23%) 0 1	93, 115, 139, 172	0
18	9I	67/88 (76%)	1.47	21 (31%) 0 0	93, 111, 135, 144	0
19	AA	78/93 (83%)	1.43	25 (32%) 0 0	150, 182, 201, 206	0
19	AI	80/93 (86%)	1.02	18 (22%) 0 1	116, 149, 171, 175	0
20	BA	99/106 (93%)	0.08	1 (1%) 82 82	90, 115, 143, 158	0
20	BI	99/106 (93%)	0.36	3 (3%) 50 49	113, 132, 170, 175	0
21	1B	25/27 (92%)	0.98	6 (24%) 0 0	121, 142, 155, 172	0
21	1F	23/27 (85%)	0.24	0 100 100	115, 129, 133, 136	0
22	1K	83/87 (95%)	0.02	5 (6%) 21 21	95, 175, 257, 267	0
22	1L	83/87 (95%)	0.62	12 (14%) 2 2	111, 193, 298, 318	0
23	2K	73/87 (83%)	-0.06	4 (5%) 25 23	90, 130, 190, 235	0
23	2L	74/87 (85%)	0.21	5 (6%) 17 17	89, 140, 212, 243	0
24	3K	77/87 (88%)	0.04	3 (3%) 39 37	82, 248, 274, 282	0
24	3L	75/87 (86%)	-0.21	3 (4%) 38 36	82, 207, 221, 226	0
25	4K	16/60 (26%)	0.02	0 100 100	89, 147, 205, 218	0
25	4L	21/60 (35%)	0.16	2 (9%) 8 8	105, 147, 219, 221	0
26	14	2909/2917 (99%)	-0.35	34 (1%) 79 78	61, 91, 244, 328	0
26	1H	2912/2917 (99%)	-0.40	23 (0%) 86 86	55, 86, 243, 301	0
27	16	122/122 (100%)	-0.71	0 100 100	87, 113, 138, 224	0
27	1J	122/122 (100%)	-0.68	0 100 100	92, 126, 154, 207	0
28	11	273/276 (98%)	0.30	4 (1%) 73 72	55, 76, 93, 103	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å <sup>2</sup> )	Q<0.9
28	19	273/276 (98%)	0.53	11 (4%) 38 36	60, 81, 97, 113	0
29	21	205/206 (99%)	0.72	23 (11%) 5 5	63, 98, 141, 156	0
29	29	205/206 (99%)	0.68	18 (8%) 10 10	68, 96, 137, 158	0
30	31	202/210 (96%)	0.23	8 (3%) 38 36	59, 88, 127, 140	0
30	39	208/210 (99%)	0.59	12 (5%) 23 22	67, 106, 167, 200	0
31	41	181/182 (99%)	1.48	54 (29%) 0 0	113, 144, 175, 183	0
31	49	181/182 (99%)	1.84	68 (37%) 0 0	131, 162, 190, 207	0
32	51	174/180 (96%)	0.44	17 (9%) 7 7	93, 118, 133, 143	0
32	59	170/180 (94%)	1.33	49 (28%) 0 0	154, 211, 241, 256	0
33	61	146/148 (98%)	0.73	21 (14%) 2 2	86, 134, 157, 167	0
33	69	146/148 (98%)	1.10	37 (25%) 0 0	90, 135, 158, 166	0
34	15	138/140 (98%)	0.64	9 (6%) 18 18	79, 105, 142, 157	0
34	58	138/140 (98%)	0.48	12 (8%) 10 10	78, 100, 136, 154	0
35	25	122/122 (100%)	0.63	8 (6%) 18 18	77, 93, 110, 116	0
35	68	122/122 (100%)	0.72	6 (4%) 29 27	66, 89, 105, 114	0
36	35	150/150 (100%)	0.68	16 (10%) 6 5	68, 107, 146, 170	0
36	78	147/150 (98%)	-0.08	0 100 100	64, 92, 114, 126	0
37	45	140/141 (99%)	1.24	32 (22%) 0 1	77, 106, 132, 154	0
37	88	141/141 (100%)	1.02	26 (18%) 1 1	73, 95, 118, 143	0
38	55	117/118 (99%)	0.30	3 (2%) 56 53	68, 86, 107, 119	0
38	98	118/118 (100%)	0.56	5 (4%) 36 34	76, 94, 113, 126	0
39	65	111/112 (99%)	0.92	16 (14%) 2 2	91, 117, 141, 150	0
39	A8	111/112 (99%)	0.76	14 (12%) 3 3	90, 107, 127, 141	0
40	75	137/146 (93%)	0.12	2 (1%) 73 72	83, 101, 163, 193	0
40	B8	129/146 (88%)	0.44	6 (4%) 31 29	84, 103, 135, 159	0
41	85	117/118 (99%)	0.27	6 (5%) 28 26	69, 93, 134, 152	0
41	C8	117/118 (99%)	0.34	2 (1%) 70 68	65, 89, 124, 139	0
42	95	101/101 (100%)	0.58	7 (6%) 16 16	70, 121, 143, 174	0
42	D8	101/101 (100%)	1.00	15 (14%) 2 2	69, 110, 140, 150	0
43	A5	113/113 (100%)	0.61	6 (5%) 26 24	68, 80, 112, 185	0
43	E8	112/113 (99%)	0.65	6 (5%) 25 24	72, 84, 120, 154	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å <sup>2</sup> )	Q<0.9
44	B5	92/96 (95%)	0.43	4 (4%) 35 34	76, 92, 116, 127	0
44	F8	94/96 (97%)	0.65	8 (8%) 10 10	67, 81, 107, 132	0
45	C5	104/110 (94%)	1.26	22 (21%) 0 1	96, 121, 156, 170	0
45	G8	103/110 (93%)	0.32	2 (1%) 66 65	83, 103, 144, 153	0
46	D5	135/206 (65%)	1.58	47 (34%) 0 0	110, 142, 177, 191	0
46	H8	171/206 (83%)	1.04	26 (15%) 2 2	100, 140, 212, 221	0
47	E5	80/85 (94%)	0.89	12 (15%) 2 2	76, 92, 114, 143	0
47	I8	83/85 (97%)	0.75	9 (10%) 5 5	75, 88, 104, 115	0
48	F5	94/98 (95%)	0.56	6 (6%) 19 19	70, 88, 127, 146	0
48	J8	95/98 (96%)	1.02	19 (20%) 1 1	67, 85, 135, 140	0
49	G5	67/72 (93%)	0.51	3 (4%) 33 32	90, 112, 131, 146	0
49	K8	68/72 (94%)	0.23	2 (2%) 51 50	71, 90, 105, 126	0
50	H5	59/60 (98%)	0.57	5 (8%) 10 10	85, 101, 151, 166	0
50	L8	57/60 (95%)	0.29	2 (3%) 44 42	76, 94, 117, 137	0
51	I5	63/71 (88%)	2.59	38 (60%) 0 0	157, 205, 225, 231	0
51	M8	66/71 (92%)	2.18	35 (53%) 0 0	142, 184, 199, 204	0
52	J5	56/60 (93%)	0.06	1 (1%) 68 67	63, 85, 141, 149	0
52	N8	55/60 (91%)	0.80	8 (14%) 2 2	60, 100, 170, 184	0
53	L5	47/49 (95%)	0.05	1 (2%) 63 62	64, 69, 88, 95	0
53	P8	47/49 (95%)	-0.08	2 (4%) 35 34	57, 64, 82, 108	0
54	M5	53/65 (81%)	0.87	7 (13%) 3 3	68, 83, 101, 111	0
54	Q8	60/65 (92%)	0.29	2 (3%) 46 44	69, 84, 110, 124	0
All	All	20915/21710 (96%)	0.17	1388 (6%) 18 18	55, 108, 195, 328	0

All (1388) RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
31	41	2	PRO	12.7
26	14	2902	C	11.9
26	14	2797	U	10.2
36	35	150	ALA	9.4
12	3A	128	ALA	9.4
26	14	2798	C	9.3
43	A5	113	LYS	8.9

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Mol	Chain	Res	Type	RSRZ
13	4A	2	ALA	8.7
30	39	1	MET	8.4
13	4A	7	VAL	8.0
11	2I	12	ARG	7.9
26	14	2901	C	7.6
13	4A	4	ILE	7.5
26	14	654(J)	A	7.5
26	14	2	G	7.4
18	9A	88	LYS	7.2
31	41	142	PRO	7.1
45	C5	58	GLY	7.1
26	14	1	G	7.1
46	D5	2	GLU	7.1
31	49	182	LYS	6.9
43	A5	112	GLY	6.9
31	49	137	GLU	6.7
13	4A	5	ALA	6.6
51	M8	55	ARG	6.6
31	41	137	GLU	6.5
32	59	103	LEU	6.4
12	3A	64	TYR	6.4
51	M8	52	THR	6.3
33	61	146	ALA	6.3
46	D5	1	MET	6.3
22	1L	51	A	6.3
3	22	207	VAL	6.3
51	I5	40	HIS	6.2
31	49	139	LEU	6.2
22	1L	53	A	6.1
26	14	654(K)	C	6.0
45	C5	59	GLY	6.0
51	I5	9	LEU	5.9
51	I5	42	PHE	5.9
23	2L	56	G	5.8
42	D8	36	PRO	5.8
26	14	2900	A	5.8
30	39	12	LEU	5.8
26	14	2799	A	5.8
31	49	116	ASP	5.7
3	22	143	GLU	5.7
26	14	2795	G	5.7
52	N8	57	VAL	5.7

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>	<b>RSRZ</b>
2	12	115	LEU	5.6
32	59	105	LEU	5.6
40	B8	1	MET	5.6
14	5A	39	LEU	5.6
13	4I	5	ALA	5.6
26	1H	1	G	5.5
3	22	199	LYS	5.4
51	I5	5	ILE	5.4
31	49	145	THR	5.4
51	I5	39	CYS	5.4
31	49	178	PHE	5.4
13	4I	6	GLY	5.4
46	D5	96	VAL	5.3
26	1H	654(K)	C	5.3
51	I5	46	GLN	5.3
14	5A	53	LEU	5.3
14	5A	52	GLN	5.3
32	59	115	VAL	5.3
31	49	138	GLN	5.2
31	49	90	LEU	5.2
51	M8	13	ARG	5.2
26	1H	2902	C	5.2
51	M8	30	GLU	5.1
13	4A	9	ILE	5.1
39	A8	2	ALA	5.1
22	1L	52	U	5.1
32	59	4	ILE	5.1
3	22	108	ASN	5.0
48	J8	95	LEU	5.0
46	H8	155	LEU	5.0
13	4A	6	GLY	5.0
31	49	177	GLY	5.0
51	I5	29	PRO	5.0
37	45	68	ILE	4.9
41	C8	117	GLN	4.9
29	21	205	ALA	4.9
46	H8	1	MET	4.9
26	14	654(L)	G	4.9
48	J8	91	LYS	4.9
31	41	75	LYS	4.8
6	5E	101	ALA	4.8
51	I5	10	VAL	4.8

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Mol	Chain	Res	Type	RSRZ
2	12	5	ILE	4.8
11	2A	12	ARG	4.8
31	41	138	GLN	4.8
32	59	107	VAL	4.8
12	3A	129	ALA	4.7
51	M8	42	PHE	4.7
23	2L	47	C	4.7
51	M8	32	TYR	4.7
19	AA	62	ILE	4.7
31	49	108	ASN	4.7
51	M8	40	HIS	4.7
36	35	149	GLU	4.7
37	88	25	ASP	4.6
46	D5	95	PRO	4.6
3	22	204	LEU	4.6
51	I5	45	GLY	4.6
6	5E	46	ARG	4.6
46	D5	163	LEU	4.6
18	9A	85	LEU	4.6
12	3A	127	GLU	4.6
3	22	101	LEU	4.5
46	D5	68	PRO	4.5
51	M8	31	ILE	4.5
48	J8	96	LYS	4.5
42	D8	38	LEU	4.5
32	59	93	GLY	4.5
46	D5	53	ILE	4.5
3	22	206	GLU	4.5
54	Q8	34	TRP	4.5
31	49	97	ASP	4.5
51	I5	27	THR	4.4
13	4A	3	ARG	4.4
19	AI	61	TYR	4.4
26	14	2801	A	4.4
26	14	3	U	4.4
3	22	87	LEU	4.4
3	22	155	GLY	4.4
46	H8	95	PRO	4.4
10	1A	64	GLU	4.4
47	I8	3	HIS	4.4
3	22	68	VAL	4.3
47	E5	76	GLY	4.4

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Mol	Chain	Res	Type	RSRZ
35	25	1	MET	4.3
30	31	6	VAL	4.3
36	35	110	TYR	4.3
46	D5	55	HIS	4.3
46	D5	69	THR	4.3
37	45	60	ARG	4.3
29	29	70	ALA	4.3
51	I5	31	ILE	4.3
11	2I	11	LYS	4.3
45	C5	53	PRO	4.3
51	I5	32	TYR	4.3
33	69	18	VAL	4.3
46	D5	52	SER	4.3
32	59	106	THR	4.3
45	C5	55	TYR	4.3
31	49	179	PRO	4.3
3	22	146	ALA	4.3
32	59	100	GLY	4.3
12	3A	28	LYS	4.2
33	69	72	LEU	4.2
31	49	180	PHE	4.2
31	49	146	TYR	4.2
31	41	182	LYS	4.2
19	AA	67	VAL	4.2
22	1K	17	U	4.2
49	K8	43	GLN	4.2
51	I5	44	THR	4.2
38	55	69	ASP	4.2
46	D5	91	LEU	4.2
3	22	104	GLN	4.2
51	I5	36	CYS	4.2
46	H8	113	ALA	4.2
3	22	10	PHE	4.2
31	49	41	GLN	4.2
39	65	108	GLY	4.2
19	AI	78	ARG	4.1
51	I5	28	LYS	4.1
46	D5	134	PRO	4.1
3	22	147	LYS	4.1
33	69	139	GLN	4.1
13	4I	7	VAL	4.1
33	69	11	ASN	4.1

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Mol	Chain	Res	Type	RSRZ
36	35	148	LEU	4.1
29	29	205	ALA	4.1
2	1E	31	TYR	4.1
48	J8	92	LYS	4.0
45	C5	90	LEU	4.0
29	29	69	LYS	4.0
31	49	181	ARG	4.0
10	1I	101	VAL	4.0
32	59	46	GLU	4.0
31	41	90	LEU	4.0
3	22	144	SER	4.0
31	41	34	LEU	4.0
14	5A	37	PHE	4.0
18	9A	46	GLU	4.0
31	49	48	GLU	4.0
26	14	4	C	4.0
3	22	17	ASP	4.0
18	9A	42	ARG	4.0
29	29	54	GLN	4.0
33	61	113	ARG	4.0
11	2I	13	GLN	4.0
51	M8	29	PRO	4.0
5	42	155	GLU	4.0
10	1A	88	LEU	4.0
11	2A	110	ASP	4.0
40	75	6	LEU	4.0
30	39	172	TRP	4.0
22	1L	17	U	4.0
18	9A	43	PHE	4.0
12	3A	32	PHE	4.0
19	AA	49	ILE	4.0
46	H8	96	VAL	4.0
31	49	136	ARG	4.0
23	2L	48	C	4.0
46	H8	5	LEU	4.0
16	7I	53	VAL	3.9
26	14	1509	C	3.9
10	1I	98	ILE	3.9
32	59	39	PRO	3.9
32	59	114	VAL	3.9
51	I5	54	GLY	3.9
15	6A	15	PHE	3.9

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Mol	Chain	Res	Type	RSRZ
46	D5	137	ILE	3.9
37	45	61	GLY	3.9
4	32	110	PHE	3.9
39	65	33	LYS	3.9
33	61	118	LYS	3.9
45	C5	50	ARG	3.8
19	AA	9	VAL	3.8
33	69	12	LEU	3.8
3	22	107	GLN	3.8
3	22	12	LEU	3.8
4	32	134	ASP	3.8
31	41	88	ILE	3.8
48	J8	80	LEU	3.8
39	A8	41	ASP	3.8
18	9I	42	ARG	3.8
32	59	164	TYR	3.8
42	95	36	PRO	3.8
33	69	3	VAL	3.8
48	J8	94	LEU	3.8
39	65	37	ALA	3.8
46	D5	93	ASP	3.8
5	42	6	PHE	3.8
31	49	120	LEU	3.8
3	22	203	PHE	3.8
2	12	165	VAL	3.8
3	22	198	VAL	3.8
31	41	146	TYR	3.8
19	AA	71	LEU	3.8
11	2A	50	TYR	3.7
46	D5	50	GLN	3.7
37	45	104	PHE	3.7
39	65	87	PHE	3.7
13	4A	97	PRO	3.7
32	59	117	PRO	3.7
2	1E	229	VAL	3.7
3	22	55	VAL	3.7
2	12	134	GLU	3.7
46	D5	138	GLU	3.7
26	1H	896	A	3.7
48	J8	93	GLU	3.7
51	I5	30	GLU	3.7
20	BI	18	GLN	3.7

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>	<b>RSRZ</b>
31	49	39	ILE	3.7
32	59	84	SER	3.7
4	32	133	VAL	3.7
8	72	44	PHE	3.7
31	49	87	PRO	3.7
33	69	85	GLU	3.7
45	C5	29	GLU	3.7
22	1L	57	C	3.7
26	14	888	C	3.7
31	41	94	LEU	3.7
37	88	104	PHE	3.7
39	A8	49	VAL	3.7
18	9I	26	LEU	3.7
37	45	140	ALA	3.7
46	D5	54	HIS	3.7
13	4A	87	TYR	3.7
2	12	112	VAL	3.7
48	J8	70	VAL	3.7
51	M8	28	LYS	3.7
48	J8	69	LYS	3.7
10	1I	6	ILE	3.7
26	14	654(I)	C	3.7
34	15	51	PHE	3.7
29	21	196	VAL	3.7
7	6E	5	ARG	3.7
11	2A	89	ALA	3.6
6	5E	57	GLN	3.6
19	AI	48	THR	3.6
3	22	105	GLU	3.6
19	AA	47	HIS	3.6
32	51	123	PHE	3.6
51	M8	58	ARG	3.6
36	35	95	VAL	3.6
31	49	34	LEU	3.6
31	49	135	LEU	3.6
43	E8	111	HIS	3.6
26	1H	2476	A	3.6
37	45	103	MET	3.6
30	39	2	LYS	3.6
3	22	16	ARG	3.6
7	62	86	GLN	3.6
37	45	7	MET	3.6

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Mol	Chain	Res	Type	RSRZ
19	AI	41	VAL	3.6
31	41	150	ASP	3.6
43	E8	24	ILE	3.6
18	9A	26	LEU	3.6
44	B5	92	LEU	3.6
39	65	60	GLY	3.6
14	5A	50	LYS	3.6
3	22	109	PRO	3.5
32	59	17	VAL	3.5
34	15	1	MET	3.5
31	49	62	LEU	3.5
13	4A	102	ARG	3.5
14	5A	38	GLY	3.5
16	7I	32	TYR	3.5
17	8A	7	THR	3.5
13	4I	8	GLU	3.5
37	88	105	GLU	3.5
26	1H	2900	A	3.5
10	1I	94	VAL	3.5
3	22	53	ALA	3.5
13	4I	96	LEU	3.5
47	I8	2	ALA	3.5
37	45	129	THR	3.5
4	32	69	GLY	3.5
19	AI	44	MET	3.5
31	49	175	LEU	3.5
26	1H	163	U	3.5
2	12	163	PHE	3.5
3	2E	66	VAL	3.5
31	41	139	LEU	3.5
19	AI	76	PRO	3.5
51	I5	47	GLN	3.5
42	D8	98	GLU	3.5
51	M8	25	TYR	3.5
10	1A	65	LEU	3.5
3	22	172	ARG	3.5
2	12	240	GLN	3.5
19	AA	30	LEU	3.5
14	5A	51	GLY	3.5
31	49	147	ASP	3.4
17	8A	22	LEU	3.4
33	61	70	GLU	3.4

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Mol	Chain	Res	Type	RSRZ
31	41	52	ILE	3.4
46	H8	147	GLY	3.4
18	9A	87	ARG	3.4
10	1I	8	LEU	3.4
51	M8	66	SER	3.4
3	22	142	MET	3.4
31	41	72	ARG	3.4
37	45	130	LYS	3.4
26	1H	888	C	3.4
46	D5	51	ALA	3.4
12	3A	55	VAL	3.4
3	22	154	SER	3.4
18	9A	44	LEU	3.4
33	69	116	LEU	3.4
6	5E	55	ASP	3.4
31	49	58	GLN	3.4
29	29	96	PHE	3.4
3	22	103	VAL	3.4
26	14	654(O)	G	3.4
36	35	123	LEU	3.4
26	1H	2901	C	3.4
37	88	99	PRO	3.3
39	A8	112	PHE	3.3
37	88	33	GLY	3.3
37	45	105	GLU	3.3
12	3A	87	GLY	3.3
33	61	122	GLU	3.3
36	35	144	GLU	3.3
11	2I	31	THR	3.3
39	65	112	PHE	3.3
7	62	85	TYR	3.3
12	3I	64	TYR	3.3
18	9I	78	LEU	3.3
14	5A	56	VAL	3.3
28	19	2	ALA	3.3
32	59	101	ARG	3.3
3	22	166	GLU	3.3
4	32	23	GLY	3.3
12	3A	69	TYR	3.3
42	95	12	TYR	3.3
10	1I	5	ARG	3.3
19	AA	82	GLY	3.3

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Mol	Chain	Res	Type	RSRZ
2	12	116	GLU	3.3
31	41	143	GLU	3.3
31	49	142	PRO	3.3
37	45	69	PHE	3.3
51	I5	34	GLU	3.3
11	2A	13	GLN	3.3
12	3A	68	ALA	3.3
42	D8	99	ILE	3.3
26	1H	654(L)	G	3.2
32	59	169	VAL	3.2
2	12	4	GLU	3.2
22	1K	84	A	3.2
36	35	118	GLY	3.2
4	3E	110	PHE	3.2
26	1H	4	C	3.2
18	9I	28	GLU	3.2
30	39	208	GLY	3.2
7	62	38	LEU	3.2
51	I5	24	THR	3.2
41	85	90	VAL	3.2
32	59	34	GLU	3.2
33	69	17	GLN	3.2
46	H8	128	VAL	3.2
10	1I	95	GLU	3.2
51	M8	65	ASP	3.2
45	C5	46	LYS	3.2
41	85	69	CYS	3.2
23	2K	1	G	3.2
37	88	38	GLU	3.2
13	4A	66	LEU	3.2
3	22	106	VAL	3.2
46	H8	149	SER	3.2
31	41	153	ARG	3.2
16	7I	49	LEU	3.2
17	8A	11	VAL	3.2
18	9A	84	LYS	3.2
10	1A	61	GLU	3.2
29	21	90	THR	3.2
51	I5	52	THR	3.2
18	9I	21	LYS	3.2
19	AA	41	VAL	3.2
28	19	147	LEU	3.2

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Mol	Chain	Res	Type	RSRZ
13	4I	65	LYS	3.2
31	41	23	PHE	3.2
39	65	2	ALA	3.2
47	E5	71	ASP	3.2
3	22	153	VAL	3.2
3	2E	166	GLU	3.2
31	41	48	GLU	3.2
32	59	6	ARG	3.2
38	98	115	GLU	3.2
46	D5	126	VAL	3.2
32	59	168	PRO	3.2
3	22	131	ARG	3.2
31	49	102	PHE	3.2
46	H8	104	PHE	3.2
32	59	43	VAL	3.2
37	88	129	THR	3.2
12	3A	100	ILE	3.1
35	68	63	VAL	3.1
51	M8	3	GLU	3.1
8	7E	63	LEU	3.1
2	12	114	ARG	3.1
52	N8	55	ARG	3.1
19	AI	74	PHE	3.1
32	59	104	GLU	3.1
33	69	19	VAL	3.1
46	D5	132	ASN	3.1
42	95	1	MET	3.1
33	69	145	VAL	3.1
51	I5	8	LYS	3.1
22	1L	54	G	3.1
18	9I	29	PHE	3.1
47	E5	85	ALA	3.1
10	1A	85	LEU	3.1
33	69	140	LEU	3.1
37	88	1	MET	3.1
29	21	204	ALA	3.1
3	22	178	LEU	3.1
18	9A	31	LEU	3.1
35	68	122	LEU	3.1
51	I5	17	GLY	3.1
4	32	179	GLU	3.1
31	49	150	ASP	3.1

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<b>Mol</b>	<b>Chain</b>	<b>Res</b>	<b>Type</b>	<b>RSRZ</b>
51	M8	53	GLU	3.1
31	41	82	LEU	3.1
13	4A	65	LYS	3.1
47	I8	5	LYS	3.1
16	7I	4	ILE	3.1
29	29	5	LEU	3.1
4	32	70	ILE	3.1
31	41	25	TYR	3.1
31	41	58	GLN	3.1
31	41	89	GLY	3.1
36	35	126	VAL	3.1
22	1L	18	G	3.1
33	61	140	LEU	3.1
48	J8	74	VAL	3.1
51	M8	34	GLU	3.1
52	N8	56	LYS	3.1
29	21	79	ARG	3.1
51	M8	54	GLY	3.1
19	AI	29	ARG	3.1
18	9I	23	LYS	3.0
32	59	45	VAL	3.0
31	49	83	ARG	3.0
48	J8	73	LEU	3.0
12	3A	31	PRO	3.0
51	M8	11	PRO	3.0
51	I5	4	GLY	3.0
13	4A	8	GLU	3.0
12	3A	84	LEU	3.0
13	4I	56	LEU	3.0
32	59	41	MET	3.0
3	2E	201	TYR	3.0
29	21	88	GLY	3.0
38	98	7	GLY	3.0
46	D5	133	ILE	3.0
51	I5	43	TYR	3.0
3	2E	68	VAL	3.0
3	22	66	VAL	3.0
31	41	37	VAL	3.0
39	A8	48	LEU	3.0
42	D8	40	LEU	3.0
30	39	133	ASN	3.0
10	1A	10	GLY	3.0

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Mol	Chain	Res	Type	RSRZ
12	3I	61	THR	3.0
28	19	181	GLU	3.0
13	4A	10	PRO	3.0
6	5E	47	ARG	3.0
41	85	118	GLY	3.0
26	1H	887	A	3.0
46	H8	146	ILE	3.0
14	5A	49	HIS	3.0
31	49	107	LEU	3.0
32	59	87	LEU	3.0
29	21	1	MET	3.0
11	2I	43	SER	3.0
21	1B	13	ILE	3.0
19	AA	22	LEU	3.0
28	11	111	LEU	3.0
31	41	102	PHE	3.0
54	Q8	48	PHE	3.0
11	2I	50	TYR	3.0
26	1H	2799	A	3.0
8	72	133	LEU	3.0
9	8E	126	SER	3.0
2	12	129	GLU	3.0
13	4A	64	TRP	3.0
33	69	138	ILE	3.0
45	C5	39	VAL	3.0
51	I5	11	PRO	3.0
54	M5	40	GLU	3.0
23	2K	56	G	2.9
46	D5	125	LEU	2.9
51	I5	21	VAL	2.9
33	69	14	ASP	2.9
31	49	161	THR	2.9
2	1E	127	ILE	2.9
33	69	4	ILE	2.9
10	1I	71	LEU	2.9
10	1A	90	LEU	2.9
32	51	172	LYS	2.9
31	41	149	VAL	2.9
31	49	140	ILE	2.9
19	AI	30	LEU	2.9
26	14	1095	A	2.9
31	41	80	PHE	2.9

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Mol	Chain	Res	Type	RSRZ
11	2A	11	LYS	2.9
3	22	177	THR	2.9
31	49	122	PRO	2.9
37	45	97	VAL	2.9
46	D5	128	VAL	2.9
15	6I	88	ARG	2.9
37	45	65	PHE	2.9
5	4E	89	ILE	2.9
37	88	32	TYR	2.9
31	41	35	GLU	2.9
31	41	59	GLU	2.9
10	1I	74	ILE	2.9
22	1L	50	A	2.9
33	61	144	VAL	2.9
51	M8	9	LEU	2.9
53	P8	46	VAL	2.9
3	22	190	ARG	2.9
3	22	157	ILE	2.9
8	72	86	ILE	2.9
13	4A	88	ARG	2.9
28	19	90	ALA	2.9
26	14	1065	U	2.9
51	M8	63	TYR	2.9
4	32	16	GLY	2.9
18	9I	51	LEU	2.9
44	F8	81	VAL	2.9
18	9I	41	LYS	2.9
3	22	6	HIS	2.9
22	1K	83	U	2.9
11	2A	95	ILE	2.9
31	49	155	MET	2.9
46	H8	102	LEU	2.9
50	H5	59	VAL	2.9
17	8I	32	TYR	2.9
34	15	72	TYR	2.9
26	1H	1075	C	2.9
31	41	87	PRO	2.9
46	D5	101	PRO	2.9
7	62	156	TRP	2.9
24	3L	38	G	2.9
38	98	114	VAL	2.9
29	21	3	GLY	2.9

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Mol	Chain	Res	Type	RSRZ
8	72	107	LEU	2.9
18	9I	40	LEU	2.9
37	45	66	ILE	2.9
46	D5	5	LEU	2.9
51	M8	56	VAL	2.9
18	9I	83	GLU	2.9
33	69	10	GLU	2.9
46	H8	79	ARG	2.9
33	69	2	LYS	2.9
31	41	86	MET	2.8
2	1E	122	PHE	2.8
45	C5	49	VAL	2.8
32	51	16	SER	2.8
33	69	125	GLU	2.8
10	1I	65	LEU	2.8
12	3A	99	HIS	2.8
37	45	32	TYR	2.8
46	D5	3	TYR	2.8
19	AI	71	LEU	2.8
31	49	152	LEU	2.8
44	F8	92	LEU	2.8
3	22	7	PRO	2.8
51	I5	7	PRO	2.8
14	5A	34	TYR	2.8
51	I5	26	SER	2.8
52	N8	52	TYR	2.8
8	7E	119	LEU	2.8
19	AA	48	THR	2.8
26	14	654(H)	G	2.8
37	45	34	LEU	2.8
51	I5	38	LYS	2.8
2	12	232	PRO	2.8
3	22	174	PRO	2.8
33	61	73	GLU	2.8
6	5E	48	LEU	2.8
16	7A	6	LEU	2.8
2	12	57	PHE	2.8
18	9I	43	PHE	2.8
10	1A	89	ASP	2.8
41	85	91	ASP	2.8
31	49	113	ARG	2.8
33	61	130	TYR	2.8

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Mol	Chain	Res	Type	RSRZ
3	2E	102	ASN	2.8
31	49	109	VAL	2.8
3	2E	72	LYS	2.8
32	59	95	ARG	2.8
18	9I	79	LEU	2.8
3	22	202	ILE	2.8
31	41	39	ILE	2.8
31	49	157	ILE	2.8
37	45	38	GLU	2.8
37	88	98	LYS	2.8
30	31	19	GLU	2.8
50	H5	28	LEU	2.8
32	59	94	TYR	2.8
37	45	93	TYR	2.8
32	59	123	PHE	2.8
33	69	144	VAL	2.8
35	68	52	VAL	2.8
51	I5	33	VAL	2.8
39	A8	43	GLU	2.7
50	L8	57	GLU	2.7
14	5A	57	ARG	2.7
11	2I	30	VAL	2.7
33	69	78	THR	2.7
33	69	20	ASP	2.7
46	H8	6	LYS	2.7
47	E5	21	LEU	2.7
9	82	115	GLY	2.7
8	7E	93	VAL	2.7
19	AA	75	ALA	2.7
21	1B	2	GLY	2.7
22	1L	58	U	2.7
31	49	134	GLY	2.7
40	B8	47	GLY	2.7
46	D5	136	PHE	2.7
28	19	177	LEU	2.7
45	C5	45	VAL	2.7
46	H8	88	PHE	2.7
3	2E	57	ILE	2.7
29	21	80	GLU	2.7
36	35	145	PRO	2.7
41	85	72	HIS	2.7
26	1H	1095	A	2.7

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Mol	Chain	Res	Type	RSRZ
32	59	33	LEU	2.7
33	61	116	LEU	2.7
44	B5	68	ARG	2.7
2	12	122	PHE	2.7
17	8A	9	VAL	2.7
36	35	130	PHE	2.7
11	2A	83	ILE	2.7
17	8I	36	ILE	2.7
3	22	205	GLY	2.7
10	1I	96	ILE	2.7
12	3A	60	LEU	2.7
4	32	185	PHE	2.7
3	22	184	TYR	2.7
33	61	65	ALA	2.7
48	J8	51	VAL	2.7
29	21	4	ILE	2.7
23	2L	81	G	2.7
32	59	128	PRO	2.7
50	H5	60	GLU	2.7
51	M8	22	ILE	2.7
10	1A	54	PHE	2.7
29	29	1	MET	2.7
14	5A	55	GLY	2.7
11	2A	87	THR	2.7
31	41	57	ALA	2.7
12	3A	35	GLY	2.7
13	4I	97	PRO	2.7
32	59	26	VAL	2.7
37	88	132	VAL	2.7
54	M5	34	TRP	2.7
51	M8	59	PHE	2.7
31	49	149	VAL	2.7
34	58	100	GLU	2.7
51	I5	53	GLU	2.7
11	2A	72	ALA	2.7
39	A8	40	ILE	2.7
8	72	112	LEU	2.7
20	BI	72	LEU	2.7
33	69	35	LEU	2.7
33	69	112	LYS	2.7
45	C5	81	LYS	2.7
49	G5	60	LEU	2.7

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Mol	Chain	Res	Type	RSRZ
37	45	33	GLY	2.7
10	1I	75	ILE	2.7
11	2I	83	ILE	2.7
11	2A	32	ILE	2.7
16	7I	31	LYS	2.6
31	41	145	THR	2.6
34	15	8	GLN	2.6
45	G8	64	GLU	2.6
51	M8	50	VAL	2.6
3	22	200	ALA	2.6
11	2A	108	ILE	2.6
22	1K	1	G	2.6
14	5A	36	PHE	2.6
5	42	98	THR	2.6
7	6E	11	GLN	2.6
46	D5	88	PHE	2.6
12	3A	63	GLY	2.6
33	69	1	MET	2.6
46	H8	161	VAL	2.6
13	4I	19	LEU	2.6
35	68	53	LYS	2.6
45	C5	63	LYS	2.6
31	41	115	ARG	2.6
31	41	148	MET	2.6
31	49	143	GLU	2.6
2	1E	188	ALA	2.6
4	32	146	ILE	2.6
4	32	207	TYR	2.6
18	9A	58	LEU	2.6
30	31	22	ALA	2.6
33	61	72	LEU	2.6
29	21	51	PHE	2.6
31	41	141	PHE	2.6
32	59	170	ARG	2.6
41	C8	90	VAL	2.6
46	D5	135	GLU	2.6
33	61	141	LYS	2.6
37	45	121	ALA	2.6
26	14	2898	U	2.6
3	2E	101	LEU	2.6
30	31	124	LEU	2.6
3	2E	128	PHE	2.6

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Mol	Chain	Res	Type	RSRZ
11	2A	25	TYR	2.6
16	7I	9	PHE	2.6
31	49	115	ARG	2.6
51	M8	27	THR	2.6
7	6E	87	VAL	2.6
43	E8	112	GLY	2.6
18	9I	50	ILE	2.6
33	69	128	LEU	2.6
39	65	73	LEU	2.6
52	N8	25	LEU	2.6
13	4I	110	ARG	2.6
20	BA	9	ASN	2.6
18	9I	22	VAL	2.6
48	F5	49	VAL	2.6
54	M5	58	ILE	2.6
2	12	6	THR	2.6
13	4I	85	GLY	2.6
26	14	2899	G	2.6
26	1H	654(J)	A	2.6
19	AI	49	ILE	2.6
29	21	183	LEU	2.6
39	A8	24	LEU	2.6
39	A8	84	GLN	2.6
30	39	22	ALA	2.6
12	3A	23	LYS	2.6
46	D5	12	GLY	2.6
46	D5	162	GLU	2.6
31	49	103	LEU	2.6
29	29	204	ALA	2.6
31	49	25	TYR	2.6
31	41	83	ARG	2.6
32	59	51	ARG	2.6
2	12	202	PRO	2.6
4	32	135	LEU	2.6
31	41	152	LEU	2.6
35	25	122	LEU	2.6
45	C5	5	MET	2.6
31	49	141	PHE	2.5
46	D5	164	ALA	2.5
12	3A	66	VAL	2.5
12	3A	126	LYS	2.5
19	AI	47	HIS	2.5

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Mol	Chain	Res	Type	RSRZ
44	F8	68	ARG	2.5
11	2I	29	ILE	2.5
39	65	40	ILE	2.5
29	29	76	ARG	2.5
42	D8	37	VAL	2.5
51	M8	33	VAL	2.5
4	32	5	ILE	2.5
12	3A	101	VAL	2.5
29	21	6	GLY	2.5
34	58	14	VAL	2.5
31	41	135	LEU	2.5
10	1I	28	ARG	2.5
12	3A	19	ARG	2.5
53	P8	47	ARG	2.5
31	49	105	LYS	2.5
42	D8	25	LEU	2.5
12	3A	86	ARG	2.5
32	59	121	ILE	2.5
29	21	187	ALA	2.5
54	M5	23	VAL	2.5
26	1H	2798	C	2.5
31	49	23	PHE	2.5
17	8A	21	VAL	2.5
11	2A	85	ARG	2.5
26	1H	1176	G	2.5
25	4L	40	U	2.5
47	I8	4	LYS	2.5
3	22	132	ARG	2.5
36	35	142	GLY	2.5
48	J8	60	PHE	2.5
51	M8	5	ILE	2.5
2	1E	228	GLY	2.5
3	22	13	GLY	2.5
18	9I	44	LEU	2.5
29	21	50	GLY	2.5
33	69	114	LEU	2.5
47	I8	6	ALA	2.5
35	25	42	SER	2.5
19	AI	40	ILE	2.5
17	8I	96	GLU	2.5
19	AA	44	MET	2.5
10	1I	100	THR	2.5

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Mol	Chain	Res	Type	RSRZ
26	14	2793	G	2.5
32	51	171	LEU	2.5
36	35	125	VAL	2.5
39	65	55	ALA	2.5
52	N8	54	GLY	2.5
8	72	64	LYS	2.5
31	49	75	LYS	2.5
37	88	130	LYS	2.5
22	1K	53	A	2.5
37	88	68	ILE	2.5
43	A5	103	ILE	2.5
22	1L	83	U	2.5
31	49	50	ALA	2.5
33	61	123	LEU	2.5
35	25	65	THR	2.5
38	98	69	ASP	2.5
31	41	76	SER	2.5
51	I5	3	GLU	2.5
21	1B	21	TYR	2.4
19	AI	62	ILE	2.4
32	59	3	ARG	2.4
31	49	99	MET	2.4
16	7I	37	GLY	2.4
46	H8	126	VAL	2.4
47	E5	75	LEU	2.4
2	12	152	PHE	2.4
32	59	132	ARG	2.4
30	39	10	PRO	2.4
34	58	99	LEU	2.4
34	15	87	LEU	2.4
36	35	106	LEU	2.4
7	6E	6	ARG	2.4
2	1E	45	GLN	2.4
24	3L	19	G	2.4
28	11	92	ILE	2.4
45	C5	47	LYS	2.4
29	29	52	LEU	2.4
30	39	192	LEU	2.4
31	49	86	MET	2.4
34	58	15	LEU	2.4
37	88	106	VAL	2.4
37	45	132	VAL	2.4

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Mol	Chain	Res	Type	RSRZ
46	H8	18	LEU	2.4
12	3I	19	ARG	2.4
47	E5	72	ARG	2.4
3	22	186	PHE	2.4
8	72	134	ILE	2.4
31	49	144	ILE	2.4
45	C5	64	GLU	2.4
45	C5	56	PRO	2.4
31	49	5	VAL	2.4
32	51	3	ARG	2.4
32	59	88	LEU	2.4
3	22	69	HIS	2.4
26	14	1084	A	2.4
37	45	71	ASP	2.4
12	3A	62	SER	2.4
30	39	199	TRP	2.4
33	69	36	ALA	2.4
5	4E	81	GLU	2.4
3	22	94	LEU	2.4
13	4I	55	ARG	2.4
46	D5	56	VAL	2.4
4	32	169	LYS	2.4
26	14	2792	G	2.4
2	12	164	VAL	2.4
3	22	173	VAL	2.4
31	41	53	LEU	2.4
19	AI	42	PRO	2.4
19	AA	69	HIS	2.4
11	2I	42	TRP	2.4
24	3K	21	A	2.4
11	2A	84	VAL	2.4
19	AI	60	VAL	2.4
19	AA	45	VAL	2.4
31	49	133	LEU	2.4
29	29	2	LYS	2.4
32	59	15	VAL	2.4
3	22	110	ASN	2.4
46	H8	62	PRO	2.4
11	2A	111	ASP	2.4
51	I5	13	ARG	2.4
14	5A	44	LEU	2.4
14	5A	47	LEU	2.4

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Mol	Chain	Res	Type	RSRZ
34	15	84	LYS	2.4
37	45	125	LEU	2.4
47	I8	46	LYS	2.4
2	12	157	ARG	2.4
32	59	129	THR	2.4
24	3L	35	G	2.4
29	21	78	LEU	2.4
50	H5	26	LEU	2.4
29	29	25	VAL	2.4
40	75	106	SER	2.4
1	1G	1538	C	2.4
7	62	37	ASN	2.4
18	9I	38	GLU	2.4
42	D8	29	PRO	2.4
2	1E	201	ILE	2.4
31	41	140	ILE	2.4
45	C5	6	HIS	2.4
2	12	71	VAL	2.4
22	1L	56	G	2.4
26	14	2802	G	2.4
37	45	62	GLY	2.4
32	51	39	PRO	2.4
37	88	48	GLU	2.4
3	22	15	THR	2.4
3	22	182	ILE	2.4
36	35	138	LEU	2.4
47	I8	7	LEU	2.4
18	9A	22	VAL	2.3
42	95	14	VAL	2.3
47	E5	22	GLY	2.3
35	68	1	MET	2.3
12	3I	65	GLU	2.3
18	9I	34	TYR	2.3
37	88	97	VAL	2.3
35	25	37	ASP	2.3
37	45	98	LYS	2.3
2	1E	126	GLU	2.3
19	AA	43	GLU	2.3
8	72	80	ILE	2.3
10	1A	16	LEU	2.3
12	3A	85	ILE	2.3
49	G5	44	LEU	2.3

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Mol	Chain	Res	Type	RSRZ
8	72	138	TRP	2.3
15	6A	89	GLY	2.3
26	1H	654(H)	G	2.3
42	95	45	THR	2.3
9	8E	110	GLU	2.3
16	7A	52	ASP	2.3
3	22	201	TYR	2.3
16	7I	33	ILE	2.3
23	2K	87	A	2.3
29	21	14	ILE	2.3
38	55	68	ARG	2.3
2	12	197	VAL	2.3
33	69	37	VAL	2.3
46	D5	165	VAL	2.3
48	F5	60	PHE	2.3
33	61	76	THR	2.3
46	H8	162	GLU	2.3
48	J8	89	GLU	2.3
51	M8	18	CYS	2.3
29	21	76	ARG	2.3
12	3A	70	ILE	2.3
31	41	43	LEU	2.3
42	95	38	LEU	2.3
44	F8	69	TYR	2.3
51	I5	50	VAL	2.3
10	1A	67	THR	2.3
2	1E	77	ALA	2.3
3	22	117	ALA	2.3
28	19	5	LYS	2.3
45	C5	31	LEU	2.3
54	M5	29	LYS	2.3
12	3A	29	GLY	2.3
18	9I	80	PRO	2.3
19	AA	76	PRO	2.3
22	1L	48	C	2.3
33	69	92	VAL	2.3
43	A5	101	SER	2.3
12	3A	65	GLU	2.3
34	58	74	ARG	2.3
46	D5	97	GLU	2.3
29	21	55	ASN	2.3
51	I5	22	ILE	2.3

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Mol	Chain	Res	Type	RSRZ
52	J5	53	ALA	2.3
54	M5	16	ILE	2.3
18	9A	86	VAL	2.3
11	2A	86	GLY	2.3
40	B8	45	PHE	2.3
26	1H	889	C	2.3
31	49	148	MET	2.3
19	AI	69	HIS	2.3
46	D5	102	LEU	2.3
19	AA	63	THR	2.3
7	62	43	PHE	2.3
29	29	3	GLY	2.3
8	7E	36	LEU	2.3
3	2E	182	ILE	2.3
19	AA	40	ILE	2.3
2	1E	165	VAL	2.3
8	72	26	VAL	2.3
8	72	79	VAL	2.3
12	3A	102	ARG	2.3
14	5A	35	ARG	2.3
37	45	35	VAL	2.3
47	E5	45	PHE	2.3
46	H8	97	GLU	2.3
42	D8	94	LEU	2.3
3	2E	61	ALA	2.3
5	4E	95	ALA	2.3
10	1I	72	VAL	2.3
13	4I	3	ARG	2.3
13	4I	75	ALA	2.3
14	5A	46	GLU	2.3
39	65	43	GLU	2.3
9	82	40	LEU	2.3
30	31	24	LEU	2.3
34	58	33	LEU	2.3
32	59	72	ILE	2.3
30	39	128	ALA	2.3
31	49	131	TYR	2.3
32	59	153	LYS	2.3
42	D8	97	LYS	2.3
15	6I	89	GLY	2.3
26	1H	2797	U	2.3
3	2E	196	LEU	2.3

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Mol	Chain	Res	Type	RSRZ
4	32	49	ARG	2.3
5	42	43	LEU	2.3
37	88	17	LEU	2.3
29	29	4	ILE	2.3
19	AA	50	ALA	2.3
33	69	136	VAL	2.3
5	4E	122	GLU	2.2
46	D5	59	LEU	2.2
51	M8	49	PHE	2.2
12	3A	56	ALA	2.2
32	59	134	SER	2.2
37	88	50	ALA	2.2
46	H8	9	TYR	2.2
37	88	59	ARG	2.2
19	AI	15	LEU	2.2
37	88	128	LYS	2.2
34	15	85	ILE	2.2
29	29	116	VAL	2.2
43	A5	105	VAL	2.2
48	F5	62	VAL	2.2
23	2K	47	C	2.2
26	14	2794	C	2.2
28	11	93	ALA	2.2
32	59	108	GLY	2.2
4	32	35	ARG	2.2
4	32	64	LEU	2.2
16	7I	50	LYS	2.2
16	7I	1	MET	2.2
15	6A	87	ILE	2.2
28	19	6	PHE	2.2
10	1I	77	PRO	2.2
37	45	96	VAL	2.2
46	D5	27	VAL	2.2
5	42	81	GLU	2.2
11	2I	93	GLN	2.2
32	51	104	GLU	2.2
32	51	152	ARG	2.2
34	58	133	GLN	2.2
42	D8	60	GLU	2.2
44	F8	90	GLU	2.2
50	L8	32	GLN	2.2
19	AA	20	LEU	2.2

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Mol	Chain	Res	Type	RSRZ
2	1E	96	ARG	2.2
19	AA	12	ASP	2.2
19	AA	13	ASP	2.2
3	22	135	LYS	2.2
30	31	131	GLY	2.2
31	49	55	LYS	2.2
32	51	83	TYR	2.2
34	15	83	LYS	2.2
40	B8	92	GLY	2.2
44	B5	26	TYR	2.2
7	62	89	MET	2.2
29	21	47	VAL	2.2
32	59	151	ILE	2.2
33	69	109	ILE	2.2
37	45	133	ARG	2.2
46	H8	90	VAL	2.2
13	4I	16	ASP	2.2
11	2I	61	ALA	2.2
43	A5	104	THR	2.2
44	F8	26	TYR	2.2
5	4E	119	LEU	2.2
29	21	5	LEU	2.2
11	2I	95	ILE	2.2
37	88	102	VAL	2.2
39	A8	46	VAL	2.2
3	22	148	GLY	2.2
16	7I	39	TYR	2.2
30	31	8	GLN	2.2
32	51	96	ALA	2.2
10	1I	97	GLU	2.2
33	69	7	GLU	2.2
34	58	68	GLU	2.2
45	C5	30	VAL	2.2
39	A8	37	ALA	2.2
4	32	158	ILE	2.2
13	4I	60	VAL	2.2
32	59	131	VAL	2.2
3	2E	71	ALA	2.2
31	49	82	LEU	2.2
21	1B	5	ASP	2.2
33	61	43	ASN	2.2
39	65	28	VAL	2.2

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Mol	Chain	Res	Type	RSRZ
46	D5	28	MET	2.2
2	12	118	LEU	2.2
7	62	149	ARG	2.2
31	41	136	ARG	2.2
31	49	67	LYS	2.2
37	88	67	ARG	2.2
3	22	86	VAL	2.2
42	D8	47	VAL	2.2
28	19	128	GLY	2.2
3	2E	193	TYR	2.2
3	22	54	ARG	2.2
4	3E	96	LEU	2.2
33	61	69	LYS	2.2
31	41	158	ALA	2.2
24	3K	35	G	2.2
28	19	67	PHE	2.2
14	5A	42	ILE	2.2
51	M8	39	CYS	2.2
37	88	80	GLU	2.2
37	45	91	GLU	2.2
43	E8	109	GLU	2.2
11	2I	49	GLY	2.2
31	41	47	LYS	2.2
47	E5	19	LYS	2.2
47	E5	41	ARG	2.2
2	1E	187	LEU	2.1
17	8A	6	LEU	2.1
33	69	5	LEU	2.1
38	98	116	LEU	2.1
39	65	32	LEU	2.1
9	82	15	ALA	2.1
3	22	170	GLN	2.1
34	58	51	PHE	2.1
29	29	9	VAL	2.1
32	51	42	ARG	2.1
39	A8	23	ARG	2.1
46	H8	171	ILE	2.1
19	AA	46	GLY	2.1
10	1A	62	HIS	2.1
11	2I	98	LEU	2.1
52	N8	26	THR	2.1
42	D8	27	ALA	2.1

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Mol	Chain	Res	Type	RSRZ
44	B5	79	ALA	2.1
3	22	11	ARG	2.1
4	32	145	GLU	2.1
8	72	101	PRO	2.1
10	1A	34	VAL	2.1
13	4I	25	ILE	2.1
16	7A	54	GLU	2.1
31	41	164	GLU	2.1
35	68	108	GLU	2.1
46	D5	103	ARG	2.1
48	J8	90	ILE	2.1
5	4E	154	GLY	2.1
11	2I	81	ASP	2.1
33	61	114	LEU	2.1
25	4L	34	G	2.1
8	7E	136	GLU	2.1
30	39	27	GLU	2.1
31	41	63	ILE	2.1
31	49	92	VAL	2.1
46	H8	133	ILE	2.1
3	2E	145	GLY	2.1
18	9A	48	GLY	2.1
33	69	8	PRO	2.1
37	45	70	PRO	2.1
46	D5	62	PRO	2.1
43	E8	69	LEU	2.1
8	7E	110	ALA	2.1
12	3A	21	LYS	2.1
13	4I	43	THR	2.1
21	1B	14	TRP	2.1
39	A8	76	LYS	2.1
3	2E	170	GLN	2.1
5	42	109	ILE	2.1
28	19	92	ILE	2.1
31	41	54	GLU	2.1
32	51	9	ILE	2.1
32	51	113	VAL	2.1
41	85	89	GLU	2.1
48	J8	7	ILE	2.1
5	42	5	ASP	2.1
16	7I	7	ALA	2.1
26	14	893	C	2.1

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Mol	Chain	Res	Type	RSRZ
45	C5	86	ARG	2.1
46	D5	131	ARG	2.1
33	61	74	ASN	2.1
42	95	93	GLU	2.1
30	31	123	LEU	2.1
33	69	13	GLY	2.1
31	49	68	PRO	2.1
38	55	70	LEU	2.1
53	L5	1	MET	2.1
29	21	2	LYS	2.1
11	2A	81	ASP	2.1
31	41	178	PHE	2.1
16	7I	2	VAL	2.1
32	51	61	HIS	2.1
48	J8	72	GLU	2.1
16	7I	36	ILE	2.1
21	1B	17	THR	2.1
49	K8	41	ILE	2.1
10	1A	71	LEU	2.1
15	6I	20	GLY	2.1
49	G5	9	GLN	2.1
13	4A	80	ARG	2.1
39	65	24	LEU	2.1
39	65	26	LEU	2.1
42	D8	39	LEU	2.1
2	12	70	PHE	2.1
33	61	126	TYR	2.1
39	A8	12	PHE	2.1
47	E5	9	SER	2.1
3	22	64	VAL	2.1
15	6A	26	GLU	2.1
8	72	35	ILE	2.1
14	5A	61	TRP	2.1
14	5A	26	ARG	2.1
26	1H	2	G	2.1
10	1A	47	PHE	2.1
28	11	15	PHE	2.1
46	D5	9	TYR	2.1
46	D5	92	SER	2.1
2	1E	209	ARG	2.1
3	2E	77	ILE	2.1
3	22	124	ILE	2.1

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Mol	Chain	Res	Type	RSRZ
14	5A	58	LYS	2.1
18	9I	55	ARG	2.1
28	19	183	ARG	2.1
31	49	63	ILE	2.1
31	49	118	ARG	2.1
36	35	124	LYS	2.1
37	45	64	ILE	2.1
45	C5	96	ILE	2.1
29	21	52	LEU	2.1
32	51	103	LEU	2.1
50	H5	32	GLN	2.1
31	49	132	ASN	2.1
35	25	34	THR	2.1
40	B8	22	PHE	2.1
51	I5	49	PHE	2.1
26	1H	277	C	2.1
14	5I	7	ILE	2.1
54	M5	22	VAL	2.1
5	42	119	LEU	2.1
10	1A	84	GLN	2.1
37	88	41	TRP	2.1
37	45	92	GLY	2.1
46	D5	104	PHE	2.1
7	62	112	PRO	2.1
12	3A	98	TYR	2.1
11	2I	91	ARG	2.1
31	41	74	LYS	2.1
43	E8	38	TYR	2.1
18	9I	85	LEU	2.1
31	41	175	LEU	2.1
34	15	116	LEU	2.1
46	H8	91	LEU	2.1
11	2I	62	GLN	2.1
7	62	139	GLU	2.1
31	41	108	ASN	2.1
32	59	81	GLU	2.1
40	B8	38	ASN	2.1
48	F5	48	LYS	2.1
48	F5	71	TYR	2.1
3	2E	64	VAL	2.1
9	82	44	VAL	2.1
34	58	52	VAL	2.1

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Mol	Chain	Res	Type	RSRZ
45	G8	30	VAL	2.1
26	14	1046	A	2.1
26	14	1177	A	2.1
39	65	56	LEU	2.1
2	1E	101	MET	2.1
3	2E	107	GLN	2.0
12	3A	20	LYS	2.0
15	6I	63	ARG	2.0
29	29	79	ARG	2.0
31	49	35	GLU	2.0
32	51	124	GLU	2.0
5	42	82	VAL	2.0
7	62	152	ALA	2.0
37	88	49	ALA	2.0
3	22	43	LEU	2.0
11	2I	32	ILE	2.0
46	D5	67	LEU	2.0
5	42	28	PHE	2.0
3	22	56	ASP	2.0
11	2A	54	ARG	2.0
17	8A	12	SER	2.0
31	49	100	TRP	2.0
33	69	126	TYR	2.0
34	58	53	VAL	2.0
44	F8	83	VAL	2.0
18	9A	27	GLY	2.0
18	9A	76	LEU	2.0
32	51	162	ILE	2.0
35	25	19	ILE	2.0
37	88	100	GLY	2.0
4	32	168	ARG	2.0
5	4E	18	ARG	2.0
44	F8	28	PHE	2.0
16	7A	59	TRP	2.0
19	AA	14	HIS	2.0
51	M8	23	GLU	2.0
51	I5	6	HIS	2.0
23	2L	57	C	2.0
26	14	546	C	2.0
32	59	85	LYS	2.0
48	F5	37	ILE	2.0
46	D5	83	PRO	2.0

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Mol	Chain	Res	Type	RSRZ
17	8A	71	PHE	2.0
47	I8	69	PHE	2.0
51	M8	26	SER	2.0
2	12	155	LEU	2.0
4	3E	97	LEU	2.0
11	2I	21	ILE	2.0
12	3A	30	ALA	2.0
48	J8	71	TYR	2.0
16	7I	19	ILE	2.0
33	69	38	LEU	2.0
11	2I	45	GLY	2.0
20	BI	47	GLY	2.0
48	J8	55	GLY	2.0
51	M8	4	GLY	2.0
35	25	45	GLU	2.0
47	E5	53	MET	2.0
11	2I	84	VAL	2.0
32	59	113	VAL	2.0
52	N8	45	VAL	2.0
2	12	137	ARG	2.0
3	22	39	ILE	2.0
12	3I	33	ARG	2.0
13	4A	48	LEU	2.0
33	61	79	ILE	2.0
34	58	30	ILE	2.0
24	3K	13	G	2.0
26	14	654(N)	G	2.0
12	3A	71	PRO	2.0
17	8A	20	THR	2.0
29	29	73	GLU	2.0
42	D8	32	THR	2.0
47	I8	24	LYS	2.0
51	M8	20	ASN	2.0
3	2E	79	ARG	2.0
3	22	75	VAL	2.0
5	42	133	TYR	2.0
32	59	83	TYR	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, 95<sup>th</sup> 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 <sup>2</sup> )	Q<0.9
23	H2U	2K	17	20/21	0.86	0.17	125,135,173,181	0
22	PSU	1L	66	20/21	0.89	0.14	123,141,157,157	0
23	OMG	2K	18	24/25	0.89	0.14	125,139,151,157	0
23	H2U	2L	17	20/21	0.90	0.16	129,137,180,190	0
23	PSU	2K	66	20/21	0.91	0.10	117,125,133,135	0
22	PSU	1K	66	20/21	0.92	0.14	113,134,150,151	0
22	5MU	1K	65	21/22	0.92	0.15	109,126,139,143	0
22	PSU	1L	39	20/21	0.92	0.15	112,122,132,132	0
23	OMG	2L	18	24/25	0.93	0.16	136,144,154,157	0
22	PSU	1K	39	20/21	0.93	0.15	97,105,110,111	0
23	PSU	2L	66	20/21	0.93	0.11	120,131,136,140	0
23	PSU	2K	39	20/21	0.94	0.21	83,96,104,110	0
23	5MU	2L	65	21/22	0.94	0.13	116,128,142,149	0
22	1MG	1K	38	24/25	0.95	0.15	93,100,106,107	0
23	PSU	2L	39	20/21	0.95	0.13	95,108,114,117	0
22	1MG	1L	38	24/25	0.95	0.15	106,120,124,127	0
22	5MU	1L	65	21/22	0.96	0.16	126,135,149,154	0
23	1MG	2L	38	24/25	0.96	0.13	103,110,121,121	0
23	1MG	2K	38	24/25	0.96	0.15	87,96,107,111	0
23	5MU	2K	65	21/22	0.97	0.08	119,125,131,134	0

### 6.3 Carbohydrates ⓘ

There are no carbohydrates in this entry.

### 6.4 Ligands ⓘ

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, 95<sup>th</sup> 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 <sup>2</sup> )	Q<0.9
55	MG	13	1673	1/1	0.14	0.50	145,145,145,145	0
55	MG	13	1676	1/1	0.26	0.29	103,103,103,103	0
55	MG	1G	1724	1/1	0.28	0.16	99,99,99,99	0
55	MG	1H	3335	1/1	0.32	0.31	101,101,101,101	0
55	MG	1H	3203	1/1	0.35	0.27	110,110,110,110	0
55	MG	1H	3175	1/1	0.35	0.68	112,112,112,112	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	1H	3145	1/1	0.40	0.51	83,83,83,83	0
55	MG	14	3252	1/1	0.41	0.73	97,97,97,97	0
55	MG	1H	3193	1/1	0.41	0.32	116,116,116,116	0
55	MG	1H	3048	1/1	0.41	0.33	87,87,87,87	0
55	MG	14	3229	1/1	0.41	0.57	77,77,77,77	0
55	MG	1G	1689	1/1	0.43	0.23	103,103,103,103	0
55	MG	14	3367	1/1	0.44	0.40	102,102,102,102	0
55	MG	13	1706	1/1	0.46	0.38	95,95,95,95	0
55	MG	1G	1719	1/1	0.47	0.31	118,118,118,118	0
55	MG	14	3306	1/1	0.49	0.42	95,95,95,95	0
55	MG	2K	102	1/1	0.49	0.36	118,118,118,118	0
55	MG	14	3276	1/1	0.50	0.51	86,86,86,86	0
55	MG	1G	1717	1/1	0.51	0.81	99,99,99,99	0
55	MG	14	3130	1/1	0.51	0.49	102,102,102,102	0
55	MG	1H	3400	1/1	0.51	0.45	94,94,94,94	0
55	MG	13	1620	1/1	0.52	0.22	92,92,92,92	0
55	MG	13	1713	1/1	0.54	0.42	99,99,99,99	0
55	MG	14	3129	1/1	0.54	0.33	86,86,86,86	0
55	MG	13	1702	1/1	0.54	0.22	96,96,96,96	0
55	MG	1G	1601	1/1	0.54	0.35	91,91,91,91	0
55	MG	13	1686	1/1	0.55	0.57	93,93,93,93	0
55	MG	1H	3188	1/1	0.55	0.47	92,92,92,92	0
55	MG	1H	3313	1/1	0.55	0.45	98,98,98,98	0
55	MG	1H	3259	1/1	0.56	0.99	98,98,98,98	0
55	MG	1G	1729	1/1	0.56	0.52	89,89,89,89	0
55	MG	14	3322	1/1	0.56	0.19	93,93,93,93	0
55	MG	1H	3248	1/1	0.56	0.42	106,106,106,106	0
55	MG	14	3171	1/1	0.56	0.45	103,103,103,103	0
55	MG	1H	3183	1/1	0.56	0.47	90,90,90,90	0
55	MG	14	3223	1/1	0.57	0.40	118,118,118,118	0
55	MG	1H	3494	1/1	0.58	0.15	125,125,125,125	0
55	MG	14	3369	1/1	0.58	0.68	105,105,105,105	0
55	MG	1H	3038	1/1	0.58	0.16	96,96,96,96	0
55	MG	14	3159	1/1	0.59	0.41	102,102,102,102	0
55	MG	1H	3338	1/1	0.59	0.61	82,82,82,82	0
55	MG	14	3301	1/1	0.60	0.43	90,90,90,90	0
55	MG	13	1670	1/1	0.60	0.29	99,99,99,99	0
55	MG	1G	1680	1/1	0.60	0.32	88,88,88,88	0
55	MG	1H	3324	1/1	0.60	0.44	89,89,89,89	0
55	MG	14	3364	1/1	0.61	0.27	89,89,89,89	0
55	MG	1H	3495	1/1	0.61	0.07	112,112,112,112	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	14	3247	1/1	0.61	0.50	78,78,78,78	0
55	MG	1G	1738	1/1	0.62	0.07	137,137,137,137	0
55	MG	1H	3033	1/1	0.62	0.24	92,92,92,92	0
55	MG	1H	3499	1/1	0.62	0.19	103,103,103,103	0
55	MG	29	303	1/1	0.63	0.28	79,79,79,79	0
55	MG	21	302	1/1	0.63	0.23	77,77,77,77	0
55	MG	1G	1655	1/1	0.63	0.33	108,108,108,108	0
55	MG	1G	1657	1/1	0.63	0.22	92,92,92,92	0
55	MG	14	3096	1/1	0.64	0.57	62,62,62,62	0
55	MG	88	304	1/1	0.64	0.90	77,77,77,77	0
55	MG	1H	3073	1/1	0.64	0.66	68,68,68,68	0
55	MG	14	3400	1/1	0.64	0.10	91,91,91,91	0
55	MG	14	3235	1/1	0.64	0.51	91,91,91,91	0
55	MG	14	3119	1/1	0.65	0.41	90,90,90,90	0
55	MG	14	3120	1/1	0.65	0.29	69,69,69,69	0
55	MG	1H	3242	1/1	0.65	0.47	97,97,97,97	0
55	MG	1H	3368	1/1	0.65	0.42	88,88,88,88	0
55	MG	1H	3279	1/1	0.65	0.66	86,86,86,86	0
55	MG	13	1722	1/1	0.65	0.26	106,106,106,106	0
55	MG	14	3035	1/1	0.66	0.71	91,91,91,91	0
55	MG	1H	3241	1/1	0.66	0.42	92,92,92,92	0
55	MG	1H	3266	1/1	0.66	0.23	79,79,79,79	0
55	MG	14	3374	1/1	0.66	0.43	102,102,102,102	0
55	MG	1H	3262	1/1	0.66	0.25	93,93,93,93	0
55	MG	1G	1674	1/1	0.67	0.25	113,113,113,113	0
55	MG	14	3233	1/1	0.67	0.69	91,91,91,91	0
55	MG	1G	1702	1/1	0.67	0.30	95,95,95,95	0
55	MG	14	3382	1/1	0.67	0.35	91,91,91,91	0
55	MG	1G	1625	1/1	0.68	0.38	99,99,99,99	0
55	MG	1H	3314	1/1	0.68	0.69	92,92,92,92	0
55	MG	14	3175	1/1	0.68	0.89	95,95,95,95	0
56	ZN	G8	202	1/1	0.68	0.36	205,205,205,205	0
55	MG	1H	3319	1/1	0.68	0.53	102,102,102,102	0
55	MG	1H	3198	1/1	0.68	0.34	113,113,113,113	0
55	MG	1H	3226	1/1	0.68	0.23	93,93,93,93	0
55	MG	1G	1693	1/1	0.68	0.34	109,109,109,109	0
55	MG	14	3230	1/1	0.68	0.46	85,85,85,85	0
55	MG	1H	3283	1/1	0.68	0.95	85,85,85,85	0
55	MG	1G	1742	1/1	0.69	0.09	131,131,131,131	0
55	MG	14	3381	1/1	0.69	0.41	95,95,95,95	0
55	MG	14	3264	1/1	0.69	0.37	78,78,78,78	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	1H	3354	1/1	0.69	0.21	107,107,107,107	0
55	MG	1G	1707	1/1	0.69	0.57	92,92,92,92	0
55	MG	1G	1701	1/1	0.69	0.18	119,119,119,119	0
55	MG	1H	3144	1/1	0.69	0.13	93,93,93,93	0
55	MG	1H	3169	1/1	0.69	0.34	90,90,90,90	0
55	MG	14	3388	1/1	0.70	0.72	94,94,94,94	0
55	MG	1H	3404	1/1	0.70	0.62	83,83,83,83	0
55	MG	1H	3391	1/1	0.70	0.25	78,78,78,78	0
55	MG	1H	3380	1/1	0.70	0.47	100,100,100,100	0
55	MG	14	3395	1/1	0.70	0.21	98,98,98,98	0
55	MG	1H	3206	1/1	0.70	0.59	69,69,69,69	0
55	MG	1H	3342	1/1	0.70	0.16	73,73,73,73	0
55	MG	14	3204	1/1	0.70	0.56	89,89,89,89	0
55	MG	14	3173	1/1	0.70	0.44	76,76,76,76	0
55	MG	14	3236	1/1	0.70	0.32	119,119,119,119	0
55	MG	14	3270	1/1	0.70	0.36	87,87,87,87	0
55	MG	14	3319	1/1	0.71	0.44	89,89,89,89	0
55	MG	14	3358	1/1	0.71	0.55	89,89,89,89	0
55	MG	14	3352	1/1	0.71	0.50	99,99,99,99	0
55	MG	14	3243	1/1	0.71	0.39	75,75,75,75	0
55	MG	1H	3292	1/1	0.71	0.31	90,90,90,90	0
55	MG	14	3393	1/1	0.71	0.70	79,79,79,79	0
55	MG	13	1684	1/1	0.72	0.41	98,98,98,98	0
55	MG	14	3202	1/1	0.72	0.53	85,85,85,85	0
55	MG	13	1718	1/1	0.72	0.28	78,78,78,78	0
55	MG	1H	3272	1/1	0.72	0.54	85,85,85,85	0
55	MG	13	1727	1/1	0.72	0.46	90,90,90,90	0
55	MG	25	201	1/1	0.72	0.32	107,107,107,107	0
55	MG	13	1730	1/1	0.72	0.15	95,95,95,95	0
55	MG	1G	1659	1/1	0.72	0.61	83,83,83,83	0
55	MG	1G	1613	1/1	0.72	0.21	92,92,92,92	0
55	MG	14	3280	1/1	0.72	0.17	87,87,87,87	0
55	MG	13	1721	1/1	0.72	0.40	84,84,84,84	0
55	MG	13	1657	1/1	0.73	0.23	90,90,90,90	0
55	MG	1H	3384	1/1	0.73	0.36	79,79,79,79	0
55	MG	13	1627	1/1	0.73	0.80	85,85,85,85	0
55	MG	1G	1645	1/1	0.73	0.22	80,80,80,80	0
55	MG	14	3272	1/1	0.73	0.43	90,90,90,90	0
55	MG	13	1631	1/1	0.73	0.65	80,80,80,80	0
55	MG	13	1649	1/1	0.73	0.35	91,91,91,91	0
55	MG	1H	3261	1/1	0.73	0.30	85,85,85,85	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	14	3432	1/1	0.74	0.12	93,93,93,93	0
55	MG	13	1705	1/1	0.74	0.61	102,102,102,102	0
55	MG	1G	1696	1/1	0.74	0.25	94,94,94,94	0
55	MG	1H	3337	1/1	0.74	0.31	82,82,82,82	0
55	MG	1G	1708	1/1	0.74	0.58	66,66,66,66	0
55	MG	13	1698	1/1	0.74	0.39	89,89,89,89	0
55	MG	14	3228	1/1	0.74	0.14	78,78,78,78	0
55	MG	14	3238	1/1	0.74	0.16	98,98,98,98	0
55	MG	14	3232	1/1	0.74	0.27	82,82,82,82	0
55	MG	1G	1712	1/1	0.74	0.12	87,87,87,87	0
55	MG	1H	3441	1/1	0.75	0.09	103,103,103,103	0
55	MG	1H	3479	1/1	0.75	0.07	103,103,103,103	0
55	MG	13	1658	1/1	0.75	0.39	82,82,82,82	0
55	MG	14	3463	1/1	0.75	0.09	119,119,119,119	0
55	MG	1H	3490	1/1	0.75	0.12	128,128,128,128	0
55	MG	14	3323	1/1	0.75	0.41	79,79,79,79	0
55	MG	14	3307	1/1	0.75	0.31	91,91,91,91	0
55	MG	1H	3406	1/1	0.75	0.56	87,87,87,87	0
55	MG	13	1612	1/1	0.75	0.70	74,74,74,74	0
55	MG	14	3221	1/1	0.76	0.64	92,92,92,92	0
55	MG	13	1619	1/1	0.76	0.24	111,111,111,111	0
55	MG	14	3187	1/1	0.76	0.31	81,81,81,81	0
55	MG	1H	3194	1/1	0.76	0.30	86,86,86,86	0
55	MG	14	3115	1/1	0.76	0.39	77,77,77,77	0
55	MG	14	3386	1/1	0.76	0.51	79,79,79,79	0
55	MG	1H	3208	1/1	0.76	0.44	96,96,96,96	0
55	MG	14	3127	1/1	0.76	0.89	76,76,76,76	0
55	MG	1H	3229	1/1	0.76	0.45	82,82,82,82	0
55	MG	1H	3343	1/1	0.76	0.77	90,90,90,90	0
55	MG	14	3001	1/1	0.76	0.71	81,81,81,81	0
55	MG	1H	3449	1/1	0.76	0.13	110,110,110,110	0
55	MG	1H	3158	1/1	0.76	0.47	88,88,88,88	0
55	MG	1H	3395	1/1	0.76	0.28	100,100,100,100	0
55	MG	1H	3352	1/1	0.76	0.23	87,87,87,87	0
55	MG	13	1743	1/1	0.77	0.09	142,142,142,142	0
55	MG	1H	3125	1/1	0.77	0.60	73,73,73,73	0
55	MG	13	1675	1/1	0.77	0.32	87,87,87,87	0
55	MG	1H	3361	1/1	0.77	0.73	78,78,78,78	0
55	MG	14	3216	1/1	0.77	0.18	77,77,77,77	0
55	MG	14	3375	1/1	0.77	0.40	81,81,81,81	0
55	MG	13	1674	1/1	0.77	0.20	112,112,112,112	0
55	MG	1H	3223	1/1	0.77	0.31	68,68,68,68	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	13	1695	1/1	0.77	0.29	92,92,92,92	0
55	MG	13	1677	1/1	0.77	0.49	77,77,77,77	0
55	MG	1H	3171	1/1	0.77	0.24	80,80,80,80	0
55	MG	1H	3132	1/1	0.77	0.29	63,63,63,63	0
55	MG	14	3131	1/1	0.77	0.49	93,93,93,93	0
55	MG	1H	3392	1/1	0.77	0.30	86,86,86,86	0
55	MG	1H	3347	1/1	0.77	0.23	80,80,80,80	0
55	MG	14	3275	1/1	0.77	0.47	96,96,96,96	0
55	MG	1H	3164	1/1	0.77	0.50	66,66,66,66	0
55	MG	13	1696	1/1	0.78	0.23	73,73,73,73	0
55	MG	14	3054	1/1	0.78	0.34	72,72,72,72	0
55	MG	1G	1632	1/1	0.78	0.28	81,81,81,81	0
55	MG	1H	3104	1/1	0.78	0.64	77,77,77,77	0
55	MG	14	3116	1/1	0.78	0.43	65,65,65,65	0
55	MG	14	3265	1/1	0.78	0.69	88,88,88,88	0
55	MG	1G	1705	1/1	0.78	0.38	105,105,105,105	0
55	MG	14	3292	1/1	0.78	0.59	79,79,79,79	0
55	MG	1H	3154	1/1	0.78	0.42	81,81,81,81	0
55	MG	1H	3299	1/1	0.78	0.20	79,79,79,79	0
55	MG	14	3390	1/1	0.78	0.29	70,70,70,70	0
55	MG	14	3152	1/1	0.78	0.44	99,99,99,99	0
55	MG	14	3162	1/1	0.78	0.97	94,94,94,94	0
55	MG	14	3324	1/1	0.78	0.43	80,80,80,80	0
55	MG	14	3385	1/1	0.78	0.42	97,97,97,97	0
55	MG	14	3126	1/1	0.78	0.14	83,83,83,83	0
55	MG	13	1678	1/1	0.78	0.76	85,85,85,85	0
55	MG	14	3155	1/1	0.78	0.41	72,72,72,72	0
55	MG	1H	3213	1/1	0.78	0.25	55,55,55,55	0
55	MG	14	3372	1/1	0.78	0.47	77,77,77,77	0
55	MG	1H	3381	1/1	0.78	0.89	79,79,79,79	0
55	MG	13	1683	1/1	0.79	0.74	83,83,83,83	0
55	MG	14	3486	1/1	0.79	0.07	138,138,138,138	0
55	MG	1H	3295	1/1	0.79	0.34	87,87,87,87	0
55	MG	1H	3307	1/1	0.79	0.35	83,83,83,83	0
55	MG	1H	3086	1/1	0.79	0.50	84,84,84,84	0
55	MG	1H	3250	1/1	0.79	1.87	97,97,97,97	0
55	MG	1G	1690	1/1	0.79	0.41	97,97,97,97	0
55	MG	1H	3423	1/1	0.79	0.13	110,110,110,110	0
55	MG	1H	3493	1/1	0.79	0.20	122,122,122,122	0
55	MG	13	1622	1/1	0.79	0.30	85,85,85,85	0
55	MG	13	1703	1/1	0.79	0.29	85,85,85,85	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	14	3192	1/1	0.79	0.29	102,102,102,102	0
55	MG	1H	3402	1/1	0.79	0.51	86,86,86,86	0
55	MG	13	1729	1/1	0.79	0.10	122,122,122,122	0
55	MG	45	201	1/1	0.79	0.53	76,76,76,76	0
55	MG	14	3193	1/1	0.79	0.36	83,83,83,83	0
55	MG	1H	3195	1/1	0.79	0.54	84,84,84,84	0
55	MG	13	1726	1/1	0.80	0.60	93,93,93,93	0
55	MG	1H	3166	1/1	0.80	0.33	83,83,83,83	0
55	MG	1G	1670	1/1	0.80	0.33	117,117,117,117	0
55	MG	1H	3451	1/1	0.80	0.07	105,105,105,105	0
55	MG	14	3217	1/1	0.80	0.32	98,98,98,98	0
55	MG	1H	3211	1/1	0.80	0.49	104,104,104,104	0
55	MG	14	3200	1/1	0.80	0.50	77,77,77,77	0
55	MG	13	1672	1/1	0.80	0.43	89,89,89,89	0
55	MG	14	3060	1/1	0.80	0.71	80,80,80,80	0
55	MG	14	3312	1/1	0.80	0.40	96,96,96,96	0
55	MG	1H	3225	1/1	0.80	0.39	80,80,80,80	0
55	MG	14	3308	1/1	0.81	0.39	89,89,89,89	0
55	MG	1H	3036	1/1	0.81	0.40	77,77,77,77	0
55	MG	14	3311	1/1	0.81	0.45	103,103,103,103	0
55	MG	14	3454	1/1	0.81	0.04	111,111,111,111	0
55	MG	1H	3323	1/1	0.81	0.22	83,83,83,83	0
55	MG	13	1669	1/1	0.81	0.65	97,97,97,97	0
55	MG	14	3253	1/1	0.81	0.34	98,98,98,98	0
55	MG	1G	1672	1/1	0.81	0.27	95,95,95,95	0
55	MG	1H	3186	1/1	0.81	0.15	76,76,76,76	0
55	MG	1G	1728	1/1	0.81	0.26	92,92,92,92	0
55	MG	1H	3390	1/1	0.81	0.46	86,86,86,86	0
55	MG	1H	3336	1/1	0.81	0.27	70,70,70,70	0
55	MG	1H	3065	1/1	0.81	0.35	80,80,80,80	0
55	MG	1H	3346	1/1	0.81	0.28	105,105,105,105	0
55	MG	14	3456	1/1	0.81	0.13	82,82,82,82	0
55	MG	14	3074	1/1	0.81	0.39	53,53,53,53	0
55	MG	1G	1725	1/1	0.81	0.56	104,104,104,104	0
55	MG	1H	3190	1/1	0.81	0.33	71,71,71,71	0
55	MG	14	3226	1/1	0.81	0.41	92,92,92,92	0
55	MG	14	3133	1/1	0.81	0.85	76,76,76,76	0
55	MG	14	3359	1/1	0.81	0.43	90,90,90,90	0
55	MG	13	1723	1/1	0.82	0.47	88,88,88,88	0
55	MG	1H	3228	1/1	0.82	0.17	68,68,68,68	0
55	MG	1H	3062	1/1	0.82	0.38	70,70,70,70	0
55	MG	14	3391	1/1	0.82	0.58	90,90,90,90	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	14	3285	1/1	0.82	0.33	72,72,72,72	0
55	MG	14	3378	1/1	0.82	0.32	83,83,83,83	0
55	MG	1G	1700	1/1	0.82	0.45	99,99,99,99	0
55	MG	14	3340	1/1	0.82	0.27	66,66,66,66	0
55	MG	1H	3204	1/1	0.82	0.34	83,83,83,83	0
55	MG	14	3309	1/1	0.82	0.29	95,95,95,95	0
55	MG	1G	1654	1/1	0.82	0.59	97,97,97,97	0
55	MG	1H	3388	1/1	0.82	0.45	90,90,90,90	0
55	MG	14	3026	1/1	0.82	0.10	83,83,83,83	0
55	MG	1H	3267	1/1	0.82	0.46	85,85,85,85	0
55	MG	1H	3257	1/1	0.82	0.34	82,82,82,82	0
55	MG	11	301	1/1	0.82	0.34	53,53,53,53	0
55	MG	1H	3411	1/1	0.82	0.16	87,87,87,87	0
55	MG	14	3145	1/1	0.82	0.37	79,79,79,79	0
55	MG	88	303	1/1	0.82	0.40	71,71,71,71	0
55	MG	7A	101	1/1	0.82	0.17	85,85,85,85	0
55	MG	14	3310	1/1	0.82	0.24	84,84,84,84	0
55	MG	14	3342	1/1	0.82	0.59	99,99,99,99	0
55	MG	14	3313	1/1	0.82	0.70	82,82,82,82	0
55	MG	1H	3382	1/1	0.82	0.58	73,73,73,73	0
55	MG	14	3318	1/1	0.83	0.25	77,77,77,77	0
55	MG	14	3389	1/1	0.83	0.36	77,77,77,77	0
55	MG	1H	3090	1/1	0.83	0.50	77,77,77,77	0
55	MG	14	3167	1/1	0.83	0.24	50,50,50,50	0
55	MG	1G	1631	1/1	0.83	0.54	88,88,88,88	0
55	MG	14	3110	1/1	0.83	0.36	74,74,74,74	0
55	MG	1H	3371	1/1	0.83	0.28	84,84,84,84	0
55	MG	13	1679	1/1	0.83	0.16	135,135,135,135	0
55	MG	13	1688	1/1	0.83	0.18	92,92,92,92	0
55	MG	1H	3394	1/1	0.83	0.67	82,82,82,82	0
55	MG	1H	3377	1/1	0.83	0.39	87,87,87,87	0
55	MG	14	3273	1/1	0.83	0.97	87,87,87,87	0
55	MG	14	3261	1/1	0.83	0.47	91,91,91,91	0
56	ZN	C5	201	1/1	0.83	0.41	201,201,201,201	0
55	MG	16	201	1/1	0.83	0.20	97,97,97,97	0
55	MG	1H	3293	1/1	0.83	0.44	78,78,78,78	0
55	MG	14	3246	1/1	0.83	0.34	97,97,97,97	0
55	MG	1H	3294	1/1	0.83	0.31	96,96,96,96	0
55	MG	14	3189	1/1	0.83	0.22	91,91,91,91	0
55	MG	1G	1658	1/1	0.83	0.77	78,78,78,78	0
55	MG	13	1662	1/1	0.83	0.12	98,98,98,98	0
55	MG	1H	3374	1/1	0.83	0.31	89,89,89,89	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	1H	3232	1/1	0.84	0.14	90,90,90,90	0
55	MG	14	3075	1/1	0.84	0.18	70,70,70,70	0
55	MG	1G	1727	1/1	0.84	0.30	98,98,98,98	0
55	MG	14	3282	1/1	0.84	0.47	89,89,89,89	0
55	MG	14	3485	1/1	0.84	0.18	101,101,101,101	0
55	MG	1H	3401	1/1	0.84	0.37	80,80,80,80	0
55	MG	E5	202	1/1	0.84	0.79	92,92,92,92	0
55	MG	14	3164	1/1	0.84	0.46	101,101,101,101	0
55	MG	14	3032	1/1	0.84	0.48	71,71,71,71	0
55	MG	14	3465	1/1	0.84	0.20	115,115,115,115	0
55	MG	13	1689	1/1	0.84	0.18	92,92,92,92	0
55	MG	1H	3332	1/1	0.84	0.96	78,78,78,78	0
55	MG	13	1719	1/1	0.84	0.41	89,89,89,89	0
55	MG	13	1710	1/1	0.84	0.55	80,80,80,80	0
55	MG	1G	1675	1/1	0.84	0.16	102,102,102,102	0
55	MG	1H	3405	1/1	0.84	0.95	89,89,89,89	0
55	MG	14	3098	1/1	0.84	0.43	87,87,87,87	0
55	MG	13	1740	1/1	0.84	0.15	106,106,106,106	0
55	MG	1H	3285	1/1	0.84	0.72	83,83,83,83	0
55	MG	14	3269	1/1	0.84	0.32	93,93,93,93	0
55	MG	14	3101	1/1	0.84	0.15	89,89,89,89	0
55	MG	14	3293	1/1	0.84	0.36	72,72,72,72	0
55	MG	14	3320	1/1	0.84	0.38	81,81,81,81	0
55	MG	14	3086	1/1	0.84	0.30	77,77,77,77	0
55	MG	13	1704	1/1	0.84	0.41	93,93,93,93	0
55	MG	1H	3116	1/1	0.84	0.31	56,56,56,56	0
55	MG	14	3422	1/1	0.84	0.13	68,68,68,68	0
55	MG	1G	1642	1/1	0.84	0.20	89,89,89,89	0
55	MG	85	202	1/1	0.84	0.37	79,79,79,79	0
55	MG	1H	3234	1/1	0.84	0.21	78,78,78,78	0
55	MG	14	3446	1/1	0.84	0.06	122,122,122,122	0
55	MG	14	3371	1/1	0.84	0.57	91,91,91,91	0
55	MG	14	3111	1/1	0.85	0.39	56,56,56,56	0
55	MG	14	3020	1/1	0.85	0.48	80,80,80,80	0
55	MG	29	301	1/1	0.85	0.42	88,88,88,88	0
55	MG	14	3489	1/1	0.85	0.07	99,99,99,99	0
55	MG	1H	3268	1/1	0.85	0.28	66,66,66,66	0
55	MG	1H	3070	1/1	0.85	0.70	73,73,73,73	0
55	MG	14	3208	1/1	0.85	0.39	74,74,74,74	0
55	MG	32	301	1/1	0.85	0.43	86,86,86,86	0
55	MG	1G	1637	1/1	0.85	0.46	84,84,84,84	0
55	MG	14	3433	1/1	0.85	0.10	91,91,91,91	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	16	205	1/1	0.85	0.40	87,87,87,87	0
55	MG	13	1717	1/1	0.85	0.37	100,100,100,100	0
55	MG	14	3124	1/1	0.85	0.24	77,77,77,77	0
55	MG	1H	3243	1/1	0.85	0.56	69,69,69,69	0
55	MG	1H	3040	1/1	0.85	0.20	69,69,69,69	0
55	MG	1G	1698	1/1	0.85	0.24	112,112,112,112	0
55	MG	14	3315	1/1	0.85	0.36	92,92,92,92	0
55	MG	14	3043	1/1	0.85	0.92	83,83,83,83	0
55	MG	14	3370	1/1	0.85	0.56	88,88,88,88	0
55	MG	1H	3465	1/1	0.85	0.11	108,108,108,108	0
55	MG	1H	3161	1/1	0.85	0.39	80,80,80,80	0
55	MG	14	3464	1/1	0.85	0.18	116,116,116,116	0
55	MG	14	3244	1/1	0.85	0.78	75,75,75,75	0
55	MG	1H	3333	1/1	0.85	0.64	85,85,85,85	0
55	MG	13	1655	1/1	0.85	0.30	77,77,77,77	0
55	MG	14	3149	1/1	0.85	0.37	77,77,77,77	0
55	MG	1H	3274	1/1	0.85	0.61	83,83,83,83	0
55	MG	1G	1679	1/1	0.85	0.27	88,88,88,88	0
55	MG	13	1741	1/1	0.85	0.05	128,128,128,128	0
55	MG	13	1720	1/1	0.85	0.30	94,94,94,94	0
55	MG	1H	3174	1/1	0.85	0.41	74,74,74,74	0
55	MG	14	3239	1/1	0.85	0.12	90,90,90,90	0
55	MG	14	3461	1/1	0.85	0.06	117,117,117,117	0
55	MG	14	3084	1/1	0.85	0.42	86,86,86,86	0
55	MG	1H	3276	1/1	0.85	0.22	100,100,100,100	0
55	MG	1H	3181	1/1	0.85	0.79	92,92,92,92	0
55	MG	1H	3256	1/1	0.85	0.28	82,82,82,82	0
55	MG	1H	3487	1/1	0.85	0.08	102,102,102,102	0
55	MG	1H	3238	1/1	0.85	0.38	77,77,77,77	0
55	MG	1H	3508	1/1	0.85	0.43	103,103,103,103	0
55	MG	1H	3155	1/1	0.85	0.16	67,67,67,67	0
55	MG	14	3326	1/1	0.85	0.42	88,88,88,88	0
55	MG	1H	3325	1/1	0.86	0.47	92,92,92,92	0
55	MG	1H	3235	1/1	0.86	0.61	88,88,88,88	0
55	MG	1H	3172	1/1	0.86	0.34	103,103,103,103	0
55	MG	1H	3264	1/1	0.86	0.24	86,86,86,86	0
55	MG	14	3296	1/1	0.86	0.63	94,94,94,94	0
55	MG	14	3305	1/1	0.86	0.52	89,89,89,89	0
55	MG	1H	3124	1/1	0.86	0.40	68,68,68,68	0
55	MG	1G	1610	1/1	0.86	0.29	86,86,86,86	0
55	MG	1H	3138	1/1	0.86	0.37	74,74,74,74	0
55	MG	13	1660	1/1	0.86	0.17	99,99,99,99	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	1H	3134	1/1	0.86	0.56	67,67,67,67	0
55	MG	14	3304	1/1	0.86	0.45	109,109,109,109	0
55	MG	1G	1668	1/1	0.86	0.49	96,96,96,96	0
55	MG	14	3470	1/1	0.86	0.16	119,119,119,119	0
55	MG	1H	3126	1/1	0.86	0.71	84,84,84,84	0
55	MG	13	1690	1/1	0.86	0.61	90,90,90,90	0
55	MG	14	3078	1/1	0.86	0.61	68,68,68,68	0
55	MG	1H	3446	1/1	0.86	0.11	83,83,83,83	0
55	MG	1H	3383	1/1	0.86	0.26	90,90,90,90	0
55	MG	1H	3327	1/1	0.86	0.55	83,83,83,83	0
55	MG	1G	1713	1/1	0.86	0.26	100,100,100,100	0
55	MG	14	3321	1/1	0.86	0.59	110,110,110,110	0
55	MG	14	3128	1/1	0.86	0.53	89,89,89,89	0
55	MG	1H	3497	1/1	0.86	0.09	113,113,113,113	0
55	MG	14	3082	1/1	0.86	0.45	76,76,76,76	0
55	MG	1H	3376	1/1	0.86	0.42	71,71,71,71	0
55	MG	13	1711	1/1	0.86	0.15	107,107,107,107	0
55	MG	14	3423	1/1	0.86	0.08	99,99,99,99	0
55	MG	1H	3165	1/1	0.86	0.41	76,76,76,76	0
55	MG	1H	3478	1/1	0.86	0.05	118,118,118,118	0
55	MG	13	1645	1/1	0.86	0.48	84,84,84,84	0
55	MG	1H	3317	1/1	0.86	0.19	79,79,79,79	0
55	MG	1H	3240	1/1	0.86	0.49	74,74,74,74	0
55	MG	1H	3043	1/1	0.86	0.33	83,83,83,83	0
55	MG	14	3150	1/1	0.86	0.25	92,92,92,92	0
55	MG	14	3438	1/1	0.86	0.26	117,117,117,117	0
55	MG	1H	3422	1/1	0.86	0.07	85,85,85,85	0
55	MG	14	3387	1/1	0.86	0.86	92,92,92,92	0
55	MG	1H	3491	1/1	0.86	0.04	122,122,122,122	0
55	MG	1H	3269	1/1	0.86	0.32	99,99,99,99	0
55	MG	13	1666	1/1	0.86	0.47	78,78,78,78	0
55	MG	1J	202	1/1	0.86	0.13	107,107,107,107	0
55	MG	1H	3298	1/1	0.86	0.32	68,68,68,68	0
55	MG	14	3140	1/1	0.87	0.26	108,108,108,108	0
55	MG	14	3274	1/1	0.87	0.25	83,83,83,83	0
55	MG	1H	3170	1/1	0.87	0.40	75,75,75,75	0
55	MG	14	3295	1/1	0.87	0.30	84,84,84,84	0
55	MG	1G	1697	1/1	0.87	0.24	88,88,88,88	0
55	MG	14	3459	1/1	0.87	0.10	122,122,122,122	0
55	MG	1G	1673	1/1	0.87	0.14	97,97,97,97	0
55	MG	1H	3133	1/1	0.87	0.40	79,79,79,79	0
55	MG	1H	3504	1/1	0.87	0.15	95,95,95,95	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	14	3384	1/1	0.87	0.24	92,92,92,92	0
55	MG	14	3250	1/1	0.87	0.19	87,87,87,87	0
55	MG	14	3278	1/1	0.87	0.51	69,69,69,69	0
55	MG	14	3317	1/1	0.87	0.22	88,88,88,88	0
55	MG	3E	301	1/1	0.87	0.47	109,109,109,109	0
55	MG	14	3263	1/1	0.87	0.44	92,92,92,92	0
55	MG	13	1667	1/1	0.87	0.32	88,88,88,88	0
55	MG	13	1642	1/1	0.87	0.13	85,85,85,85	0
55	MG	13	1744	1/1	0.87	0.08	117,117,117,117	0
55	MG	14	3134	1/1	0.87	0.27	76,76,76,76	0
55	MG	1H	3167	1/1	0.87	0.24	75,75,75,75	0
55	MG	1H	3233	1/1	0.87	0.48	100,100,100,100	0
55	MG	14	3357	1/1	0.87	0.38	93,93,93,93	0
55	MG	1G	1662	1/1	0.87	0.24	98,98,98,98	0
55	MG	P8	101	1/1	0.87	0.45	72,72,72,72	0
55	MG	1H	3398	1/1	0.87	0.17	109,109,109,109	0
55	MG	1G	1639	1/1	0.87	0.77	79,79,79,79	0
55	MG	1G	1609	1/1	0.87	0.27	101,101,101,101	0
55	MG	1H	3287	1/1	0.87	0.28	75,75,75,75	0
55	MG	13	1712	1/1	0.87	0.62	86,86,86,86	0
55	MG	1G	1720	1/1	0.87	0.28	102,102,102,102	0
55	MG	1H	3263	1/1	0.87	0.43	67,67,67,67	0
55	MG	1H	3153	1/1	0.87	0.23	79,79,79,79	0
55	MG	14	3259	1/1	0.87	0.78	88,88,88,88	0
55	MG	1H	3373	1/1	0.87	0.38	85,85,85,85	0
55	MG	14	3231	1/1	0.87	0.49	79,79,79,79	0
55	MG	1G	1726	1/1	0.88	0.54	90,90,90,90	0
55	MG	1H	3329	1/1	0.88	0.39	62,62,62,62	0
55	MG	1H	3399	1/1	0.88	0.48	94,94,94,94	0
55	MG	1G	1651	1/1	0.88	0.09	93,93,93,93	0
55	MG	13	1680	1/1	0.88	0.22	93,93,93,93	0
55	MG	1H	3356	1/1	0.88	0.32	100,100,100,100	0
55	MG	1H	3218	1/1	0.88	0.53	52,52,52,52	0
55	MG	14	3362	1/1	0.88	0.22	77,77,77,77	0
55	MG	14	3348	1/1	0.88	0.35	86,86,86,86	0
55	MG	16	204	1/1	0.88	0.18	103,103,103,103	0
55	MG	13	1665	1/1	0.88	0.11	103,103,103,103	0
55	MG	1H	3428	1/1	0.88	0.10	85,85,85,85	0
55	MG	1H	3389	1/1	0.88	0.54	99,99,99,99	0
55	MG	13	1647	1/1	0.88	0.31	93,93,93,93	0
55	MG	14	3363	1/1	0.88	0.32	67,67,67,67	0
55	MG	14	3103	1/1	0.88	0.18	73,73,73,73	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	13	1643	1/1	0.88	0.12	104,104,104,104	0
55	MG	14	3156	1/1	0.88	0.42	82,82,82,82	0
55	MG	1H	3397	1/1	0.88	0.40	76,76,76,76	0
55	MG	13	1635	1/1	0.88	0.28	101,101,101,101	0
55	MG	14	3188	1/1	0.88	0.27	82,82,82,82	0
55	MG	14	3198	1/1	0.88	0.14	95,95,95,95	0
55	MG	14	3368	1/1	0.88	0.44	91,91,91,91	0
55	MG	1H	3305	1/1	0.88	0.36	96,96,96,96	0
55	MG	1H	3085	1/1	0.88	0.34	75,75,75,75	0
55	MG	88	302	1/1	0.88	0.18	95,95,95,95	0
55	MG	1H	3315	1/1	0.88	0.40	92,92,92,92	0
55	MG	1G	1628	1/1	0.88	0.47	93,93,93,93	0
55	MG	13	1731	1/1	0.88	0.19	82,82,82,82	0
55	MG	1H	3099	1/1	0.88	0.46	59,59,59,59	0
55	MG	14	3290	1/1	0.88	0.46	76,76,76,76	0
55	MG	1G	1716	1/1	0.88	0.20	98,98,98,98	0
55	MG	1G	1634	1/1	0.88	0.43	77,77,77,77	0
55	MG	13	1737	1/1	0.88	0.11	114,114,114,114	0
55	MG	1G	1663	1/1	0.88	0.56	75,75,75,75	0
55	MG	1H	3210	1/1	0.88	0.46	82,82,82,82	0
55	MG	14	3478	1/1	0.88	0.04	109,109,109,109	0
55	MG	1H	3245	1/1	0.88	0.64	81,81,81,81	0
55	MG	14	3105	1/1	0.88	0.18	90,90,90,90	0
55	MG	1H	3408	1/1	0.88	0.26	82,82,82,82	0
55	MG	1G	1714	1/1	0.89	0.38	94,94,94,94	0
55	MG	1H	3414	1/1	0.89	0.58	97,97,97,97	0
55	MG	1H	3063	1/1	0.89	0.45	90,90,90,90	0
55	MG	2K	103	1/1	0.89	0.14	143,143,143,143	0
55	MG	14	3379	1/1	0.89	0.45	107,107,107,107	0
55	MG	14	3053	1/1	0.89	0.34	69,69,69,69	0
55	MG	14	3205	1/1	0.89	0.59	83,83,83,83	0
55	MG	14	3460	1/1	0.89	0.14	114,114,114,114	0
55	MG	1H	3207	1/1	0.89	0.32	86,86,86,86	0
55	MG	13	1732	1/1	0.89	0.06	113,113,113,113	0
55	MG	1G	1627	1/1	0.89	0.46	88,88,88,88	0
55	MG	1H	3179	1/1	0.89	0.28	71,71,71,71	0
55	MG	1H	3230	1/1	0.89	0.50	84,84,84,84	0
55	MG	1H	3191	1/1	0.89	0.13	75,75,75,75	0
55	MG	13	1644	1/1	0.89	0.88	91,91,91,91	0
55	MG	1H	3202	1/1	0.89	0.72	71,71,71,71	0
55	MG	1G	1614	1/1	0.89	0.30	98,98,98,98	0
55	MG	1G	1653	1/1	0.89	0.16	97,97,97,97	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	14	3112	1/1	0.89	0.20	77,77,77,77	0
55	MG	1G	1633	1/1	0.89	0.70	73,73,73,73	0
55	MG	13	1641	1/1	0.89	0.32	75,75,75,75	0
55	MG	1H	3492	1/1	0.89	0.20	101,101,101,101	0
55	MG	1H	3349	1/1	0.89	0.15	66,66,66,66	0
55	MG	1H	3289	1/1	0.89	0.18	80,80,80,80	0
55	MG	1H	3034	1/1	0.89	0.22	54,54,54,54	0
55	MG	1H	3463	1/1	0.89	0.07	91,91,91,91	0
55	MG	1H	3302	1/1	0.89	0.46	86,86,86,86	0
55	MG	13	1715	1/1	0.89	0.59	88,88,88,88	0
55	MG	1G	1669	1/1	0.89	0.43	100,100,100,100	0
55	MG	I8	102	1/1	0.89	0.12	108,108,108,108	0
55	MG	1H	3511	1/1	0.89	0.09	75,75,75,75	0
55	MG	14	3118	1/1	0.89	0.19	69,69,69,69	0
55	MG	1H	3173	1/1	0.89	0.40	70,70,70,70	0
55	MG	13	1694	1/1	0.89	0.76	88,88,88,88	0
55	MG	1H	3413	1/1	0.89	0.24	89,89,89,89	0
55	MG	1H	3162	1/1	0.89	0.47	89,89,89,89	0
55	MG	14	3099	1/1	0.89	0.29	57,57,57,57	0
55	MG	14	3121	1/1	0.89	0.42	69,69,69,69	0
55	MG	1H	3510	1/1	0.89	0.48	65,65,65,65	0
55	MG	1G	1732	1/1	0.89	0.07	123,123,123,123	0
55	MG	1H	3081	1/1	0.89	0.53	85,85,85,85	0
55	MG	14	3286	1/1	0.89	0.54	91,91,91,91	0
55	MG	14	3176	1/1	0.89	0.71	63,63,63,63	0
55	MG	1H	3341	1/1	0.89	0.41	83,83,83,83	0
55	MG	14	3453	1/1	0.90	0.14	91,91,91,91	0
55	MG	1H	3222	1/1	0.90	0.19	91,91,91,91	0
55	MG	1G	1739	1/1	0.90	0.24	107,107,107,107	0
55	MG	13	1616	1/1	0.90	0.21	102,102,102,102	0
55	MG	85	201	1/1	0.90	0.22	83,83,83,83	0
55	MG	13	1682	1/1	0.90	0.25	108,108,108,108	0
55	MG	1H	3253	1/1	0.90	0.28	90,90,90,90	0
55	MG	14	3028	1/1	0.90	0.46	46,46,46,46	0
55	MG	14	3294	1/1	0.90	0.52	69,69,69,69	0
55	MG	14	3210	1/1	0.90	0.26	86,86,86,86	0
55	MG	13	1639	1/1	0.90	0.17	85,85,85,85	0
55	MG	14	3219	1/1	0.90	0.37	86,86,86,86	0
55	MG	1H	3339	1/1	0.90	0.23	88,88,88,88	0
55	MG	14	3289	1/1	0.90	0.34	98,98,98,98	0
55	MG	14	3445	1/1	0.90	0.10	97,97,97,97	0
55	MG	14	3211	1/1	0.90	0.54	74,74,74,74	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	1H	3025	1/1	0.90	0.24	55,55,55,55	0
55	MG	1G	1649	1/1	0.90	0.20	109,109,109,109	0
55	MG	1H	3296	1/1	0.90	0.25	84,84,84,84	0
55	MG	14	3227	1/1	0.90	0.14	90,90,90,90	0
55	MG	1H	3199	1/1	0.90	0.16	74,74,74,74	0
55	MG	14	3237	1/1	0.90	0.16	65,65,65,65	0
55	MG	13	1699	1/1	0.90	0.61	72,72,72,72	0
55	MG	1H	3450	1/1	0.90	0.06	74,74,74,74	0
55	MG	1G	1623	1/1	0.90	0.47	89,89,89,89	0
55	MG	13	1659	1/1	0.90	0.68	91,91,91,91	0
55	MG	13	1735	1/1	0.90	0.07	111,111,111,111	0
55	MG	1H	3273	1/1	0.90	0.24	93,93,93,93	0
55	MG	1H	3278	1/1	0.90	0.42	84,84,84,84	0
55	MG	1H	3464	1/1	0.90	0.07	92,92,92,92	0
55	MG	1G	1715	1/1	0.90	0.28	83,83,83,83	0
55	MG	14	3143	1/1	0.90	0.72	64,64,64,64	0
55	MG	13	1650	1/1	0.90	0.70	88,88,88,88	0
55	MG	1H	3200	1/1	0.90	0.32	68,68,68,68	0
55	MG	14	3351	1/1	0.90	0.42	104,104,104,104	0
55	MG	1H	3265	1/1	0.90	0.33	65,65,65,65	0
55	MG	14	3401	1/1	0.90	0.15	78,78,78,78	0
55	MG	1H	3330	1/1	0.90	0.37	75,75,75,75	0
55	MG	1G	1722	1/1	0.90	0.32	84,84,84,84	0
55	MG	14	3434	1/1	0.90	0.18	89,89,89,89	0
55	MG	98	201	1/1	0.90	0.62	71,71,71,71	0
55	MG	1G	1704	1/1	0.90	0.18	78,78,78,78	0
55	MG	13	1736	1/1	0.90	0.07	111,111,111,111	0
55	MG	14	3380	1/1	0.90	0.78	93,93,93,93	0
55	MG	1G	1703	1/1	0.90	0.35	106,106,106,106	0
55	MG	1H	3180	1/1	0.90	0.20	79,79,79,79	0
55	MG	14	3431	1/1	0.90	0.06	86,86,86,86	0
55	MG	1H	3037	1/1	0.90	0.24	94,94,94,94	0
55	MG	1H	3270	1/1	0.90	0.25	99,99,99,99	0
55	MG	1H	3475	1/1	0.90	0.09	92,92,92,92	0
55	MG	1H	3151	1/1	0.90	0.27	81,81,81,81	0
55	MG	1K	101	1/1	0.90	0.27	90,90,90,90	0
55	MG	1H	3348	1/1	0.91	0.35	88,88,88,88	0
55	MG	1G	1711	1/1	0.91	0.42	88,88,88,88	0
55	MG	1H	3498	1/1	0.91	0.20	100,100,100,100	0
55	MG	1G	1677	1/1	0.91	0.40	105,105,105,105	0
55	MG	13	1714	1/1	0.91	0.08	103,103,103,103	0
55	MG	1G	1648	1/1	0.91	0.39	82,82,82,82	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	1H	3300	1/1	0.91	0.49	91,91,91,91	0
55	MG	1H	3140	1/1	0.91	0.32	82,82,82,82	0
55	MG	14	3242	1/1	0.91	0.24	98,98,98,98	0
55	MG	14	3123	1/1	0.91	0.41	63,63,63,63	0
55	MG	1H	3237	1/1	0.91	0.25	74,74,74,74	0
55	MG	1H	3045	1/1	0.91	0.18	64,64,64,64	0
55	MG	1H	3312	1/1	0.91	0.17	88,88,88,88	0
55	MG	1H	3318	1/1	0.91	0.18	50,50,50,50	0
55	MG	1H	3288	1/1	0.91	0.53	68,68,68,68	0
55	MG	14	3070	1/1	0.91	0.23	57,57,57,57	0
55	MG	1H	3160	1/1	0.91	0.38	88,88,88,88	0
55	MG	1H	3184	1/1	0.91	0.41	75,75,75,75	0
55	MG	1H	3092	1/1	0.91	0.43	73,73,73,73	0
55	MG	14	3249	1/1	0.91	0.12	100,100,100,100	0
55	MG	1H	3429	1/1	0.91	0.08	91,91,91,91	0
55	MG	14	3125	1/1	0.91	0.47	75,75,75,75	0
55	MG	78	201	1/1	0.91	0.21	74,74,74,74	0
55	MG	16	203	1/1	0.91	0.46	102,102,102,102	0
55	MG	1H	3367	1/1	0.91	0.14	77,77,77,77	0
55	MG	1H	3059	1/1	0.91	0.59	69,69,69,69	0
55	MG	1H	3474	1/1	0.91	0.09	102,102,102,102	0
55	MG	14	3373	1/1	0.91	1.13	74,74,74,74	0
55	MG	14	3163	1/1	0.91	0.33	81,81,81,81	0
55	MG	1G	1635	1/1	0.91	0.50	82,82,82,82	0
55	MG	1G	1647	1/1	0.91	0.20	92,92,92,92	0
55	MG	14	3467	1/1	0.91	0.09	97,97,97,97	0
55	MG	1H	3219	1/1	0.91	0.28	64,64,64,64	0
55	MG	1H	3290	1/1	0.91	0.28	71,71,71,71	0
55	MG	14	3403	1/1	0.91	0.06	96,96,96,96	0
55	MG	1H	3056	1/1	0.91	0.34	75,75,75,75	0
55	MG	13	1742	1/1	0.91	0.11	111,111,111,111	0
55	MG	1H	3214	1/1	0.91	0.15	56,56,56,56	0
55	MG	1H	3340	1/1	0.91	0.14	75,75,75,75	0
55	MG	14	3271	1/1	0.91	0.80	75,75,75,75	0
55	MG	1H	3331	1/1	0.91	0.14	68,68,68,68	0
55	MG	1H	3303	1/1	0.91	0.43	91,91,91,91	0
55	MG	14	3104	1/1	0.91	0.32	87,87,87,87	0
55	MG	1H	3071	1/1	0.91	0.37	74,74,74,74	0
55	MG	14	3196	1/1	0.91	0.47	65,65,65,65	0
55	MG	1H	3216	1/1	0.91	0.41	65,65,65,65	0
55	MG	14	3248	1/1	0.91	0.78	74,74,74,74	0
55	MG	1H	3500	1/1	0.91	0.07	105,105,105,105	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	14	3474	1/1	0.91	0.06	107,107,107,107	0
55	MG	13	1671	1/1	0.91	0.04	106,106,106,106	0
55	MG	13	1613	1/1	0.91	0.54	92,92,92,92	0
55	MG	1H	3375	1/1	0.91	0.18	70,70,70,70	0
55	MG	1G	1740	1/1	0.91	0.06	112,112,112,112	0
55	MG	1H	3480	1/1	0.91	0.10	118,118,118,118	0
55	MG	14	3141	1/1	0.91	0.45	78,78,78,78	0
55	MG	1H	3044	1/1	0.91	0.23	63,63,63,63	0
55	MG	1H	3275	1/1	0.91	0.24	89,89,89,89	0
55	MG	1G	1604	1/1	0.91	0.30	79,79,79,79	0
55	MG	1H	3350	1/1	0.91	0.26	79,79,79,79	0
55	MG	1H	3407	1/1	0.91	0.82	102,102,102,102	0
55	MG	1H	3481	1/1	0.91	0.06	116,116,116,116	0
55	MG	14	3154	1/1	0.91	0.47	81,81,81,81	0
55	MG	14	3281	1/1	0.91	0.20	79,79,79,79	0
55	MG	14	3350	1/1	0.92	0.41	84,84,84,84	0
55	MG	1H	3410	1/1	0.92	0.41	85,85,85,85	0
55	MG	1H	3369	1/1	0.92	0.20	86,86,86,86	0
55	MG	14	3343	1/1	0.92	0.32	75,75,75,75	0
55	MG	1H	3468	1/1	0.92	0.13	106,106,106,106	0
55	MG	14	3335	1/1	0.92	0.97	86,86,86,86	0
55	MG	13	1700	1/1	0.92	0.23	87,87,87,87	0
55	MG	1G	1640	1/1	0.92	0.52	90,90,90,90	0
55	MG	1G	1612	1/1	0.92	0.10	88,88,88,88	0
55	MG	14	3022	1/1	0.92	0.09	81,81,81,81	0
55	MG	1H	3344	1/1	0.92	0.34	85,85,85,85	0
55	MG	14	3097	1/1	0.92	0.41	91,91,91,91	0
55	MG	14	3339	1/1	0.92	0.42	71,71,71,71	0
55	MG	14	3354	1/1	0.92	0.77	76,76,76,76	0
55	MG	14	3027	1/1	0.92	0.25	85,85,85,85	0
55	MG	1H	3357	1/1	0.92	0.14	78,78,78,78	0
55	MG	1H	3484	1/1	0.92	0.25	104,104,104,104	0
55	MG	14	3314	1/1	0.92	0.21	90,90,90,90	0
55	MG	14	3366	1/1	0.92	0.24	89,89,89,89	0
55	MG	1H	3470	1/1	0.92	0.14	79,79,79,79	0
55	MG	1G	1665	1/1	0.92	0.16	108,108,108,108	0
55	MG	14	3303	1/1	0.92	0.41	71,71,71,71	0
55	MG	14	3288	1/1	0.92	0.74	89,89,89,89	0
55	MG	14	3297	1/1	0.92	0.34	85,85,85,85	0
55	MG	1G	1723	1/1	0.92	0.08	87,87,87,87	0
55	MG	14	3428	1/1	0.92	0.05	95,95,95,95	0
55	MG	1H	3358	1/1	0.92	0.12	81,81,81,81	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	1H	3306	1/1	0.92	0.33	84,84,84,84	0
55	MG	13	1685	1/1	0.92	0.45	103,103,103,103	0
55	MG	14	3030	1/1	0.92	0.28	70,70,70,70	0
55	MG	1H	3260	1/1	0.92	0.47	85,85,85,85	0
55	MG	14	3177	1/1	0.92	0.21	75,75,75,75	0
55	MG	1G	1629	1/1	0.92	0.30	89,89,89,89	0
55	MG	13	1611	1/1	0.92	0.31	115,115,115,115	0
55	MG	14	3144	1/1	0.92	0.43	61,61,61,61	0
55	MG	1H	3176	1/1	0.92	0.47	59,59,59,59	0
55	MG	14	3046	1/1	0.92	0.35	67,67,67,67	0
55	MG	14	3087	1/1	0.92	0.34	84,84,84,84	0
55	MG	14	3284	1/1	0.92	0.33	79,79,79,79	0
55	MG	14	3185	1/1	0.92	0.24	74,74,74,74	0
55	MG	13	1738	1/1	0.92	0.07	100,100,100,100	0
55	MG	14	3191	1/1	0.92	0.30	62,62,62,62	0
55	MG	14	3142	1/1	0.92	0.49	77,77,77,77	0
55	MG	1H	3372	1/1	0.92	0.97	75,75,75,75	0
55	MG	14	3457	1/1	0.92	0.05	103,103,103,103	0
55	MG	14	3056	1/1	0.92	0.53	71,71,71,71	0
55	MG	1H	3456	1/1	0.92	0.19	88,88,88,88	0
55	MG	1H	3316	1/1	0.92	0.44	87,87,87,87	0
55	MG	14	3241	1/1	0.92	0.23	83,83,83,83	0
55	MG	14	3360	1/1	0.92	0.58	93,93,93,93	0
55	MG	1H	3351	1/1	0.92	0.19	86,86,86,86	0
55	MG	1G	1664	1/1	0.92	0.32	95,95,95,95	0
55	MG	1G	1666	1/1	0.92	0.42	71,71,71,71	0
55	MG	14	3148	1/1	0.92	0.49	72,72,72,72	0
55	MG	14	3089	1/1	0.92	0.48	89,89,89,89	0
55	MG	13	1739	1/1	0.92	0.08	111,111,111,111	0
55	MG	1H	3334	1/1	0.92	0.23	75,75,75,75	0
55	MG	1H	3425	1/1	0.92	0.16	62,62,62,62	0
55	MG	14	3418	1/1	0.92	0.13	72,72,72,72	0
55	MG	1G	1743	1/1	0.92	0.18	98,98,98,98	0
55	MG	14	3483	1/1	0.92	0.12	104,104,104,104	0
55	MG	13	1648	1/1	0.92	0.25	96,96,96,96	0
55	MG	13	1633	1/1	0.93	0.41	59,59,59,59	0
55	MG	1H	3113	1/1	0.93	0.18	78,78,78,78	0
55	MG	1G	1615	1/1	0.93	0.20	84,84,84,84	0
55	MG	14	3209	1/1	0.93	0.23	91,91,91,91	0
55	MG	1G	1618	1/1	0.93	0.39	76,76,76,76	0
55	MG	1G	1616	1/1	0.93	0.53	62,62,62,62	0
55	MG	14	3316	1/1	0.93	0.31	78,78,78,78	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	14	3080	1/1	0.93	0.44	65,65,65,65	0
55	MG	1G	1667	1/1	0.93	0.20	84,84,84,84	0
55	MG	14	3036	1/1	0.93	0.28	71,71,71,71	0
55	MG	14	3283	1/1	0.93	0.56	101,101,101,101	0
55	MG	1H	3217	1/1	0.93	0.22	56,56,56,56	0
55	MG	1H	3353	1/1	0.93	0.45	75,75,75,75	0
55	MG	14	3442	1/1	0.93	0.07	109,109,109,109	0
55	MG	1H	3438	1/1	0.93	0.08	75,75,75,75	0
55	MG	1G	1683	1/1	0.93	0.16	132,132,132,132	0
55	MG	1H	3476	1/1	0.93	0.07	119,119,119,119	0
55	MG	1H	3107	1/1	0.93	0.85	76,76,76,76	0
55	MG	1G	1606	1/1	0.93	0.13	89,89,89,89	0
55	MG	1H	3112	1/1	0.93	0.69	69,69,69,69	0
55	MG	13	1651	1/1	0.93	0.12	91,91,91,91	0
55	MG	13	1664	1/1	0.93	0.58	76,76,76,76	0
55	MG	14	3122	1/1	0.93	0.61	84,84,84,84	0
55	MG	14	3245	1/1	0.93	0.44	81,81,81,81	0
55	MG	14	3468	1/1	0.93	0.32	107,107,107,107	0
55	MG	14	3016	1/1	0.93	0.43	74,74,74,74	0
55	MG	13	1654	1/1	0.93	0.29	78,78,78,78	0
55	MG	14	3137	1/1	0.93	0.51	77,77,77,77	0
55	MG	21	301	1/1	0.93	0.36	64,64,64,64	0
55	MG	14	3055	1/1	0.93	0.59	68,68,68,68	0
55	MG	1G	1660	1/1	0.93	0.21	82,82,82,82	0
55	MG	13	1716	1/1	0.93	0.27	103,103,103,103	0
55	MG	14	3405	1/1	0.93	0.09	70,70,70,70	0
55	MG	14	3341	1/1	0.93	0.41	77,77,77,77	0
55	MG	1G	1636	1/1	0.93	0.20	93,93,93,93	0
55	MG	1G	1706	1/1	0.93	0.38	91,91,91,91	0
55	MG	14	3291	1/1	0.93	0.52	91,91,91,91	0
55	MG	14	3012	1/1	0.93	0.41	48,48,48,48	0
55	MG	14	3332	1/1	0.93	0.35	80,80,80,80	0
55	MG	14	3416	1/1	0.93	0.11	96,96,96,96	0
55	MG	1G	1687	1/1	0.93	0.40	101,101,101,101	0
55	MG	1H	3322	1/1	0.93	0.27	109,109,109,109	0
55	MG	1H	3328	1/1	0.93	0.33	70,70,70,70	0
55	MG	14	3325	1/1	0.93	0.14	72,72,72,72	0
55	MG	1H	3308	1/1	0.93	0.55	77,77,77,77	0
55	MG	14	3448	1/1	0.93	0.11	109,109,109,109	0
55	MG	1G	1710	1/1	0.93	0.40	87,87,87,87	0
55	MG	14	3268	1/1	0.93	1.03	69,69,69,69	0
55	MG	1H	3311	1/1	0.93	0.11	65,65,65,65	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	13	1634	1/1	0.93	0.41	77,77,77,77	0
55	MG	14	3014	1/1	0.93	0.21	77,77,77,77	0
55	MG	13	1708	1/1	0.93	0.41	81,81,81,81	0
55	MG	1H	3143	1/1	0.93	0.31	96,96,96,96	0
55	MG	14	3414	1/1	0.93	0.13	85,85,85,85	0
55	MG	1H	3156	1/1	0.93	0.51	76,76,76,76	0
55	MG	14	3002	1/1	0.93	0.52	61,61,61,61	0
55	MG	14	3353	1/1	0.93	0.64	84,84,84,84	0
55	MG	14	3277	1/1	0.93	0.66	74,74,74,74	0
55	MG	1G	1695	1/1	0.93	0.22	95,95,95,95	0
55	MG	14	3287	1/1	0.93	0.08	98,98,98,98	0
55	MG	14	3444	1/1	0.93	0.06	106,106,106,106	0
55	MG	14	3023	1/1	0.93	0.26	56,56,56,56	0
55	MG	14	3427	1/1	0.93	0.09	97,97,97,97	0
55	MG	1H	3117	1/1	0.93	0.57	74,74,74,74	0
55	MG	14	3199	1/1	0.93	0.28	75,75,75,75	0
55	MG	45	202	1/1	0.93	0.15	99,99,99,99	0
55	MG	14	3225	1/1	0.93	0.18	87,87,87,87	0
55	MG	14	3455	1/1	0.93	0.13	121,121,121,121	0
55	MG	14	3337	1/1	0.93	0.36	64,64,64,64	0
55	MG	13	1607	1/1	0.93	0.18	77,77,77,77	0
55	MG	1G	1735	1/1	0.93	0.06	97,97,97,97	0
55	MG	14	3168	1/1	0.93	0.39	66,66,66,66	0
55	MG	1G	1652	1/1	0.93	0.28	99,99,99,99	0
55	MG	14	3408	1/1	0.93	0.15	63,63,63,63	0
55	MG	14	3109	1/1	0.93	0.48	66,66,66,66	0
55	MG	1H	3018	1/1	0.93	0.54	46,46,46,46	0
55	MG	1H	3122	1/1	0.93	0.14	68,68,68,68	0
55	MG	1H	3385	1/1	0.93	0.22	106,106,106,106	0
55	MG	14	3487	1/1	0.93	0.12	103,103,103,103	0
55	MG	14	3481	1/1	0.93	0.11	78,78,78,78	0
55	MG	13	1629	1/1	0.93	0.30	65,65,65,65	0
55	MG	1H	3486	1/1	0.94	0.07	129,129,129,129	0
55	MG	1H	3282	1/1	0.94	0.34	89,89,89,89	0
55	MG	14	3182	1/1	0.94	0.30	78,78,78,78	0
55	MG	1G	1692	1/1	0.94	0.10	100,100,100,100	0
55	MG	14	3038	1/1	0.94	0.58	57,57,57,57	0
55	MG	1H	3426	1/1	0.94	0.14	62,62,62,62	0
55	MG	1H	3393	1/1	0.94	0.22	58,58,58,58	0
55	MG	14	3336	1/1	0.94	0.39	70,70,70,70	0
55	MG	14	3328	1/1	0.94	0.40	38,38,38,38	0
55	MG	1H	3224	1/1	0.94	0.23	80,80,80,80	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	1H	3157	1/1	0.94	0.68	60,60,60,60	0
55	MG	13	1640	1/1	0.94	0.30	73,73,73,73	0
55	MG	14	3071	1/1	0.94	0.57	48,48,48,48	0
55	MG	1H	3227	1/1	0.94	0.27	61,61,61,61	0
55	MG	14	3392	1/1	0.94	0.16	88,88,88,88	0
55	MG	14	3334	1/1	0.94	0.35	76,76,76,76	0
55	MG	14	3452	1/1	0.94	0.17	94,94,94,94	0
55	MG	14	3068	1/1	0.94	0.52	57,57,57,57	0
55	MG	1H	3008	1/1	0.94	0.26	64,64,64,64	0
55	MG	14	3215	1/1	0.94	0.32	77,77,77,77	0
55	MG	1H	3069	1/1	0.94	0.32	69,69,69,69	0
55	MG	1H	3094	1/1	0.94	0.38	73,73,73,73	0
55	MG	14	3365	1/1	0.94	0.40	89,89,89,89	0
55	MG	1H	3074	1/1	0.94	0.31	59,59,59,59	0
55	MG	1H	3147	1/1	0.94	0.33	81,81,81,81	0
55	MG	1G	1661	1/1	0.94	0.46	65,65,65,65	0
55	MG	13	1623	1/1	0.94	0.46	94,94,94,94	0
55	MG	1H	3448	1/1	0.94	0.06	86,86,86,86	0
55	MG	14	3394	1/1	0.94	0.33	90,90,90,90	0
55	MG	14	3440	1/1	0.94	0.04	105,105,105,105	0
55	MG	14	3224	1/1	0.94	0.48	73,73,73,73	0
55	MG	1H	3178	1/1	0.94	0.50	69,69,69,69	0
55	MG	14	3106	1/1	0.94	0.22	80,80,80,80	0
55	MG	1H	3076	1/1	0.94	0.21	67,67,67,67	0
55	MG	1H	3482	1/1	0.94	0.06	83,83,83,83	0
55	MG	13	1615	1/1	0.94	0.24	88,88,88,88	0
55	MG	1H	3115	1/1	0.94	0.26	73,73,73,73	0
55	MG	1G	1671	1/1	0.94	0.51	84,84,84,84	0
55	MG	1H	3212	1/1	0.94	0.36	71,71,71,71	0
55	MG	1H	3320	1/1	0.94	0.23	82,82,82,82	0
55	MG	1H	3066	1/1	0.94	0.26	86,86,86,86	0
55	MG	14	3234	1/1	0.94	0.33	76,76,76,76	0
55	MG	1G	1721	1/1	0.94	0.17	102,102,102,102	0
55	MG	1H	3403	1/1	0.94	0.56	79,79,79,79	0
55	MG	13	1618	1/1	0.94	0.22	124,124,124,124	0
55	MG	1H	3280	1/1	0.94	0.18	92,92,92,92	0
55	MG	1H	3244	1/1	0.94	0.26	92,92,92,92	0
55	MG	1H	3462	1/1	0.94	0.09	91,91,91,91	0
55	MG	14	3033	1/1	0.94	0.32	77,77,77,77	0
55	MG	1H	3039	1/1	0.94	0.40	66,66,66,66	0
55	MG	14	3409	1/1	0.94	0.16	83,83,83,83	0
55	MG	14	3160	1/1	0.94	0.28	98,98,98,98	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	14	3413	1/1	0.94	0.07	100,100,100,100	0
55	MG	13	1637	1/1	0.94	0.41	72,72,72,72	0
55	MG	1H	3163	1/1	0.94	0.39	83,83,83,83	0
55	MG	14	3266	1/1	0.94	0.20	69,69,69,69	0
55	MG	1H	3473	1/1	0.94	0.12	100,100,100,100	0
55	MG	1G	1685	1/1	0.94	0.44	92,92,92,92	0
55	MG	18	101	1/1	0.94	0.36	55,55,55,55	0
55	MG	14	3201	1/1	0.94	0.31	68,68,68,68	0
55	MG	14	3139	1/1	0.94	0.31	98,98,98,98	0
55	MG	1H	3477	1/1	0.94	0.05	92,92,92,92	0
55	MG	1H	3412	1/1	0.94	0.14	103,103,103,103	0
55	MG	13	1617	1/1	0.94	0.32	83,83,83,83	0
55	MG	1H	3445	1/1	0.94	0.10	86,86,86,86	0
55	MG	1H	3029	1/1	0.94	0.46	56,56,56,56	0
55	MG	1H	3024	1/1	0.94	0.44	58,58,58,58	0
55	MG	14	3262	1/1	0.94	0.36	89,89,89,89	0
55	MG	1H	3447	1/1	0.94	0.20	77,77,77,77	0
55	MG	1H	3387	1/1	0.94	0.46	73,73,73,73	0
55	MG	14	3203	1/1	0.94	0.31	84,84,84,84	0
55	MG	14	3279	1/1	0.94	0.38	80,80,80,80	0
55	MG	1G	1630	1/1	0.94	0.37	91,91,91,91	0
55	MG	14	3441	1/1	0.94	0.03	102,102,102,102	0
55	MG	31	400	1/1	0.94	0.23	65,65,65,65	0
55	MG	1H	3503	1/1	0.94	0.22	76,76,76,76	0
55	MG	1H	3453	1/1	0.94	0.08	89,89,89,89	0
55	MG	14	3450	1/1	0.94	0.08	65,65,65,65	0
55	MG	1H	3152	1/1	0.94	0.41	54,54,54,54	0
55	MG	1G	1737	1/1	0.94	0.06	93,93,93,93	0
55	MG	1H	3079	1/1	0.94	0.24	79,79,79,79	0
55	MG	16	202	1/1	0.94	0.23	90,90,90,90	0
55	MG	1H	3046	1/1	0.94	0.24	56,56,56,56	0
55	MG	1H	3501	1/1	0.94	0.07	105,105,105,105	0
55	MG	14	3135	1/1	0.94	0.39	75,75,75,75	0
55	MG	14	3147	1/1	0.94	0.42	80,80,80,80	0
55	MG	14	3158	1/1	0.94	0.41	68,68,68,68	0
55	MG	1H	3051	1/1	0.94	0.22	74,74,74,74	0
55	MG	1H	3189	1/1	0.94	0.41	85,85,85,85	0
55	MG	1G	1620	1/1	0.94	0.26	88,88,88,88	0
55	MG	1H	3421	1/1	0.94	0.11	66,66,66,66	0
55	MG	1H	3041	1/1	0.94	0.21	72,72,72,72	0
55	MG	14	3251	1/1	0.94	0.54	74,74,74,74	0
55	MG	14	3349	1/1	0.94	0.32	95,95,95,95	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	13	1663	1/1	0.94	0.57	93,93,93,93	0
55	MG	1H	3485	1/1	0.94	0.07	84,84,84,84	0
55	MG	1H	3215	1/1	0.94	0.18	45,45,45,45	0
55	MG	14	3113	1/1	0.95	0.26	76,76,76,76	0
55	MG	1H	3444	1/1	0.95	0.04	94,94,94,94	0
55	MG	1G	1641	1/1	0.95	0.19	91,91,91,91	0
55	MG	1H	3196	1/1	0.95	0.41	77,77,77,77	0
55	MG	14	3267	1/1	0.95	0.25	74,74,74,74	0
55	MG	14	3355	1/1	0.95	0.55	71,71,71,71	0
55	MG	13	1687	1/1	0.95	0.74	90,90,90,90	0
55	MG	14	3437	1/1	0.95	0.12	88,88,88,88	0
55	MG	14	3333	1/1	0.95	0.55	58,58,58,58	0
55	MG	1H	3284	1/1	0.95	0.21	69,69,69,69	0
55	MG	14	3466	1/1	0.95	0.12	103,103,103,103	0
55	MG	1H	3496	1/1	0.95	0.20	99,99,99,99	0
55	MG	14	3338	1/1	0.95	0.25	66,66,66,66	0
55	MG	1H	3068	1/1	0.95	0.27	83,83,83,83	0
55	MG	13	1701	1/1	0.95	0.41	83,83,83,83	0
55	MG	14	3079	1/1	0.95	0.56	46,46,46,46	0
55	MG	1H	3042	1/1	0.95	0.32	84,84,84,84	0
55	MG	14	3429	1/1	0.95	0.12	72,72,72,72	0
55	MG	1H	3168	1/1	0.95	0.30	72,72,72,72	0
55	MG	1H	3460	1/1	0.95	0.15	72,72,72,72	0
55	MG	1G	1718	1/1	0.95	0.40	76,76,76,76	0
55	MG	13	1653	1/1	0.95	0.65	79,79,79,79	0
55	MG	14	3136	1/1	0.95	0.36	71,71,71,71	0
55	MG	14	3376	1/1	0.95	0.21	59,59,59,59	0
55	MG	1H	3271	1/1	0.95	0.36	77,77,77,77	0
55	MG	14	3174	1/1	0.95	0.52	61,61,61,61	0
55	MG	1H	3251	1/1	0.95	0.20	85,85,85,85	0
55	MG	13	1724	1/1	0.95	0.46	63,63,63,63	0
55	MG	1G	1681	1/1	0.95	0.17	86,86,86,86	0
55	MG	13	1681	1/1	0.95	0.27	91,91,91,91	0
55	MG	14	3424	1/1	0.95	0.11	72,72,72,72	0
55	MG	14	3059	1/1	0.95	0.37	54,54,54,54	0
55	MG	1H	3095	1/1	0.95	0.17	58,58,58,58	0
55	MG	13	1707	1/1	0.95	0.44	92,92,92,92	0
55	MG	13	1661	1/1	0.95	0.32	89,89,89,89	0
55	MG	1H	3142	1/1	0.95	0.21	68,68,68,68	0
55	MG	14	3471	1/1	0.95	0.29	113,113,113,113	0
55	MG	14	3073	1/1	0.95	0.34	70,70,70,70	0
55	MG	1H	3100	1/1	0.95	0.30	65,65,65,65	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	14	3003	1/1	0.95	0.27	63,63,63,63	0
55	MG	1H	3321	1/1	0.95	0.47	80,80,80,80	0
55	MG	1G	1650	1/1	0.95	0.22	92,92,92,92	0
55	MG	1H	3281	1/1	0.95	0.36	65,65,65,65	0
55	MG	14	3044	1/1	0.95	0.75	79,79,79,79	0
55	MG	14	3153	1/1	0.95	0.65	71,71,71,71	0
55	MG	1H	3109	1/1	0.95	0.69	39,39,39,39	0
55	MG	1H	3017	1/1	0.95	0.47	56,56,56,56	0
55	MG	14	3404	1/1	0.95	0.21	105,105,105,105	0
55	MG	14	3039	1/1	0.95	0.41	63,63,63,63	0
55	MG	14	3157	1/1	0.95	0.56	83,83,83,83	0
55	MG	14	3190	1/1	0.95	0.67	75,75,75,75	0
55	MG	1H	3472	1/1	0.95	0.06	86,86,86,86	0
55	MG	1H	3120	1/1	0.95	0.42	53,53,53,53	0
55	MG	1H	3286	1/1	0.95	0.29	94,94,94,94	0
55	MG	1H	3509	1/1	0.95	0.46	47,47,47,47	0
55	MG	1H	3083	1/1	0.95	0.36	55,55,55,55	0
55	MG	14	3377	1/1	0.95	0.45	82,82,82,82	0
55	MG	13	1646	1/1	0.95	0.71	76,76,76,76	0
55	MG	1H	3457	1/1	0.95	0.11	79,79,79,79	0
55	MG	2L	102	1/1	0.95	0.34	71,71,71,71	0
55	MG	1H	3118	1/1	0.95	0.39	74,74,74,74	0
55	MG	13	1624	1/1	0.95	0.37	92,92,92,92	0
55	MG	1H	3304	1/1	0.95	0.58	82,82,82,82	0
55	MG	13	1728	1/1	0.95	0.29	80,80,80,80	0
55	MG	1G	1741	1/1	0.95	0.07	105,105,105,105	0
55	MG	14	3040	1/1	0.95	0.81	59,59,59,59	0
55	MG	1H	3187	1/1	0.95	0.30	66,66,66,66	0
55	MG	13	1621	1/1	0.95	0.39	75,75,75,75	0
55	MG	1H	3231	1/1	0.95	0.16	90,90,90,90	0
55	MG	3I	201	1/1	0.95	0.18	68,68,68,68	0
55	MG	1H	3489	1/1	0.95	0.09	97,97,97,97	0
55	MG	13	1697	1/1	0.95	0.52	67,67,67,67	0
55	MG	1H	3249	1/1	0.95	0.38	80,80,80,80	0
55	MG	13	1609	1/1	0.95	0.29	80,80,80,80	0
55	MG	1H	3379	1/1	0.95	0.71	69,69,69,69	0
55	MG	1G	1676	1/1	0.95	0.30	93,93,93,93	0
55	MG	14	3412	1/1	0.95	0.11	77,77,77,77	0
55	MG	1H	3432	1/1	0.95	0.09	58,58,58,58	0
55	MG	13	1626	1/1	0.95	0.57	68,68,68,68	0
55	MG	14	3181	1/1	0.95	0.23	78,78,78,78	0
55	MG	14	3045	1/1	0.95	0.52	55,55,55,55	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	1H	3252	1/1	0.95	0.41	79,79,79,79	0
55	MG	1H	3192	1/1	0.95	0.49	88,88,88,88	0
55	MG	1G	1682	1/1	0.95	0.15	80,80,80,80	0
55	MG	1G	1617	1/1	0.95	0.58	72,72,72,72	0
55	MG	14	3019	1/1	0.95	0.26	80,80,80,80	0
55	MG	1H	3466	1/1	0.95	0.08	77,77,77,77	0
55	MG	1H	3309	1/1	0.95	0.34	78,78,78,78	0
55	MG	1H	3416	1/1	0.95	0.10	60,60,60,60	0
55	MG	1H	3386	1/1	0.95	0.58	87,87,87,87	0
55	MG	1H	3057	1/1	0.95	0.52	73,73,73,73	0
55	MG	14	3361	1/1	0.95	0.21	88,88,88,88	0
55	MG	1H	3424	1/1	0.95	0.13	72,72,72,72	0
55	MG	1H	3067	1/1	0.95	0.44	68,68,68,68	0
55	MG	1H	3050	1/1	0.95	0.53	65,65,65,65	0
55	MG	1G	1736	1/1	0.95	0.07	114,114,114,114	0
55	MG	1H	3363	1/1	0.95	0.67	54,54,54,54	0
55	MG	14	3004	1/1	0.95	0.28	59,59,59,59	0
55	MG	E5	201	1/1	0.95	0.24	62,62,62,62	0
55	MG	1H	3435	1/1	0.96	0.08	80,80,80,80	0
55	MG	1H	3047	1/1	0.96	0.10	96,96,96,96	0
55	MG	14	3473	1/1	0.96	0.17	61,61,61,61	0
55	MG	1H	3032	1/1	0.96	0.35	79,79,79,79	0
55	MG	14	3345	1/1	0.96	0.15	89,89,89,89	0
55	MG	14	3255	1/1	0.96	0.09	90,90,90,90	0
55	MG	13	1692	1/1	0.96	0.56	52,52,52,52	0
55	MG	14	3331	1/1	0.96	0.42	63,63,63,63	0
55	MG	1H	3201	1/1	0.96	0.23	62,62,62,62	0
55	MG	14	3161	1/1	0.96	0.50	74,74,74,74	0
55	MG	1G	1694	1/1	0.96	0.18	80,80,80,80	0
55	MG	14	3480	1/1	0.96	0.04	92,92,92,92	0
55	MG	14	3085	1/1	0.96	0.41	58,58,58,58	0
55	MG	14	3165	1/1	0.96	0.25	61,61,61,61	0
55	MG	14	3057	1/1	0.96	0.49	45,45,45,45	0
55	MG	1H	3430	1/1	0.96	0.10	87,87,87,87	0
55	MG	14	3062	1/1	0.96	0.35	71,71,71,71	0
55	MG	1H	3101	1/1	0.96	0.49	60,60,60,60	0
55	MG	1H	3247	1/1	0.96	0.44	77,77,77,77	0
55	MG	1H	3362	1/1	0.96	0.38	56,56,56,56	0
55	MG	14	3411	1/1	0.96	0.10	88,88,88,88	0
55	MG	13	1610	1/1	0.96	0.22	90,90,90,90	0
55	MG	13	1625	1/1	0.96	0.50	78,78,78,78	0
55	MG	1H	3258	1/1	0.96	0.11	92,92,92,92	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	14	3398	1/1	0.96	0.17	66,66,66,66	0
55	MG	1H	3409	1/1	0.96	0.20	75,75,75,75	0
55	MG	1G	1656	1/1	0.96	0.41	74,74,74,74	0
55	MG	1H	3102	1/1	0.96	0.38	64,64,64,64	0
55	MG	1H	3105	1/1	0.96	0.49	50,50,50,50	0
55	MG	1H	3096	1/1	0.96	0.28	62,62,62,62	0
55	MG	14	3031	1/1	0.96	0.45	66,66,66,66	0
55	MG	14	3439	1/1	0.96	0.06	105,105,105,105	0
55	MG	14	3212	1/1	0.96	0.37	88,88,88,88	0
55	MG	1H	3013	1/1	0.96	0.52	54,54,54,54	0
55	MG	14	3169	1/1	0.96	0.10	55,55,55,55	0
55	MG	88	301	1/1	0.96	0.29	82,82,82,82	0
55	MG	1H	3087	1/1	0.96	0.20	76,76,76,76	0
55	MG	1H	3130	1/1	0.96	0.53	70,70,70,70	0
55	MG	14	3138	1/1	0.96	0.59	83,83,83,83	0
55	MG	14	3482	1/1	0.96	0.11	81,81,81,81	0
55	MG	1G	1734	1/1	0.96	0.07	108,108,108,108	0
55	MG	14	3258	1/1	0.96	0.23	101,101,101,101	0
55	MG	1H	3366	1/1	0.96	0.16	76,76,76,76	0
55	MG	1H	3058	1/1	0.96	0.27	67,67,67,67	0
55	MG	14	3420	1/1	0.96	0.13	91,91,91,91	0
55	MG	13	1656	1/1	0.96	0.41	63,63,63,63	0
55	MG	1G	1638	1/1	0.96	0.38	95,95,95,95	0
55	MG	14	3443	1/1	0.96	0.11	74,74,74,74	0
55	MG	1H	3010	1/1	0.96	0.43	48,48,48,48	0
55	MG	1H	3082	1/1	0.96	0.17	70,70,70,70	0
55	MG	13	1709	1/1	0.96	0.30	99,99,99,99	0
55	MG	1H	3052	1/1	0.96	0.38	67,67,67,67	0
55	MG	1H	3149	1/1	0.96	0.65	74,74,74,74	0
55	MG	14	3435	1/1	0.96	0.09	117,117,117,117	0
55	MG	14	3451	1/1	0.96	0.10	91,91,91,91	0
55	MG	1H	3007	1/1	0.96	0.38	50,50,50,50	0
55	MG	14	3260	1/1	0.96	0.10	95,95,95,95	0
55	MG	14	3034	1/1	0.96	0.61	60,60,60,60	0
55	MG	1H	3089	1/1	0.96	0.39	71,71,71,71	0
55	MG	14	3029	1/1	0.96	0.49	63,63,63,63	0
55	MG	1H	3364	1/1	0.96	0.56	75,75,75,75	0
55	MG	13	1636	1/1	0.96	0.21	73,73,73,73	0
55	MG	13	1725	1/1	0.96	0.57	71,71,71,71	0
55	MG	14	3092	1/1	0.96	0.83	64,64,64,64	0
55	MG	14	3402	1/1	0.96	0.13	63,63,63,63	0
55	MG	13	1608	1/1	0.96	0.55	64,64,64,64	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	1H	3139	1/1	0.96	0.18	63,63,63,63	0
55	MG	1H	3129	1/1	0.96	0.38	61,61,61,61	0
55	MG	14	3058	1/1	0.96	0.51	53,53,53,53	0
55	MG	1H	3220	1/1	0.96	0.39	55,55,55,55	0
55	MG	1H	3455	1/1	0.96	0.12	62,62,62,62	0
55	MG	13	1604	1/1	0.96	0.30	71,71,71,71	0
55	MG	1G	1684	1/1	0.96	0.10	109,109,109,109	0
55	MG	14	3490	1/1	0.96	0.54	81,81,81,81	0
55	MG	14	3183	1/1	0.96	0.28	64,64,64,64	0
55	MG	1H	3123	1/1	0.96	0.30	64,64,64,64	0
55	MG	14	3117	1/1	0.96	0.19	73,73,73,73	0
55	MG	1H	3469	1/1	0.96	0.05	107,107,107,107	0
55	MG	1H	3028	1/1	0.96	0.32	52,52,52,52	0
55	MG	14	3166	1/1	0.96	0.28	61,61,61,61	0
55	MG	14	3132	1/1	0.96	0.16	78,78,78,78	0
56	ZN	5A	101	1/1	0.96	0.09	125,125,125,125	0
55	MG	1H	3254	1/1	0.96	0.29	56,56,56,56	0
55	MG	1H	3291	1/1	0.96	0.35	117,117,117,117	0
55	MG	1G	1644	1/1	0.96	0.27	96,96,96,96	0
55	MG	14	3298	1/1	0.96	0.45	54,54,54,54	0
55	MG	14	3049	1/1	0.96	0.25	80,80,80,80	0
55	MG	13	1746	1/1	0.96	0.20	100,100,100,100	0
55	MG	1H	3121	1/1	0.96	0.54	54,54,54,54	0
55	MG	1H	3009	1/1	0.96	0.32	56,56,56,56	0
55	MG	1H	3365	1/1	0.96	0.28	46,46,46,46	0
55	MG	14	3195	1/1	0.96	0.47	69,69,69,69	0
55	MG	1G	1686	1/1	0.96	0.26	108,108,108,108	0
55	MG	14	3042	1/1	0.97	0.38	47,47,47,47	0
55	MG	1H	3360	1/1	0.97	0.37	70,70,70,70	0
55	MG	G8	201	1/1	0.97	0.30	65,65,65,65	0
55	MG	1H	3454	1/1	0.97	0.15	94,94,94,94	0
55	MG	1G	1607	1/1	0.97	0.20	86,86,86,86	0
55	MG	1H	3185	1/1	0.97	0.42	66,66,66,66	0
55	MG	14	3179	1/1	0.97	0.38	90,90,90,90	0
55	MG	14	3088	1/1	0.97	0.32	73,73,73,73	0
55	MG	14	3356	1/1	0.97	0.27	77,77,77,77	0
55	MG	14	3406	1/1	0.97	0.13	62,62,62,62	0
55	MG	13	1652	1/1	0.97	0.34	94,94,94,94	0
55	MG	14	3050	1/1	0.97	0.33	72,72,72,72	0
55	MG	14	3184	1/1	0.97	0.50	59,59,59,59	0
55	MG	1H	3310	1/1	0.97	0.35	52,52,52,52	0
55	MG	1H	3023	1/1	0.97	0.46	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	14	3458	1/1	0.97	0.09	91,91,91,91	0
55	MG	14	3425	1/1	0.97	0.11	66,66,66,66	0
55	MG	1H	3461	1/1	0.97	0.10	67,67,67,67	0
55	MG	14	3094	1/1	0.97	0.30	60,60,60,60	0
55	MG	1H	3277	1/1	0.97	0.30	74,74,74,74	0
55	MG	1H	3031	1/1	0.97	0.31	57,57,57,57	0
55	MG	1H	3345	1/1	0.97	0.17	70,70,70,70	0
55	MG	1H	3442	1/1	0.97	0.14	77,77,77,77	0
55	MG	14	3477	1/1	0.97	0.13	63,63,63,63	0
55	MG	1H	3434	1/1	0.97	0.10	72,72,72,72	0
55	MG	1H	3431	1/1	0.97	0.06	74,74,74,74	0
55	MG	14	3479	1/1	0.97	0.09	78,78,78,78	0
55	MG	1H	3053	1/1	0.97	0.18	72,72,72,72	0
55	MG	14	3051	1/1	0.97	0.45	55,55,55,55	0
55	MG	14	3114	1/1	0.97	0.50	81,81,81,81	0
55	MG	14	3449	1/1	0.97	0.10	82,82,82,82	0
55	MG	14	3430	1/1	0.97	0.08	69,69,69,69	0
55	MG	1H	3433	1/1	0.97	0.08	67,67,67,67	0
55	MG	1H	3080	1/1	0.97	0.37	89,89,89,89	0
55	MG	1H	3035	1/1	0.97	0.32	82,82,82,82	0
55	MG	13	1691	1/1	0.97	0.37	89,89,89,89	0
55	MG	14	3186	1/1	0.97	0.18	69,69,69,69	0
55	MG	1H	3370	1/1	0.97	0.29	75,75,75,75	0
55	MG	1H	3467	1/1	0.97	0.11	67,67,67,67	0
55	MG	13	1734	1/1	0.97	0.14	93,93,93,93	0
55	MG	1G	1678	1/1	0.97	0.29	97,97,97,97	0
55	MG	14	3347	1/1	0.97	0.33	84,84,84,84	0
55	MG	1H	3148	1/1	0.97	0.31	75,75,75,75	0
55	MG	1H	3061	1/1	0.97	0.19	75,75,75,75	0
55	MG	14	3383	1/1	0.97	0.21	95,95,95,95	0
55	MG	14	3475	1/1	0.97	0.16	104,104,104,104	0
55	MG	13	1733	1/1	0.97	0.10	95,95,95,95	0
55	MG	1H	3436	1/1	0.97	0.10	77,77,77,77	0
55	MG	14	3206	1/1	0.97	0.23	92,92,92,92	0
55	MG	14	3072	1/1	0.97	0.52	52,52,52,52	0
55	MG	14	3476	1/1	0.97	0.07	65,65,65,65	0
55	MG	14	3010	1/1	0.97	0.35	54,54,54,54	0
55	MG	1H	3205	1/1	0.97	0.43	75,75,75,75	0
55	MG	14	3100	1/1	0.97	0.28	77,77,77,77	0
55	MG	1H	3197	1/1	0.97	0.48	48,48,48,48	0
55	MG	14	3488	1/1	0.97	0.57	64,64,64,64	0
55	MG	1H	3001	1/1	0.97	0.46	51,51,51,51	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	14	3329	1/1	0.97	0.58	58,58,58,58	0
55	MG	14	3194	1/1	0.97	0.15	66,66,66,66	0
55	MG	1G	1646	1/1	0.97	0.19	109,109,109,109	0
55	MG	1H	3246	1/1	0.97	0.30	90,90,90,90	0
55	MG	1H	3141	1/1	0.97	0.23	50,50,50,50	0
55	MG	1H	3159	1/1	0.97	0.26	76,76,76,76	0
55	MG	14	3011	1/1	0.97	0.39	48,48,48,48	0
55	MG	1G	1643	1/1	0.97	0.27	78,78,78,78	0
55	MG	14	3462	1/1	0.97	0.17	103,103,103,103	0
55	MG	13	1638	1/1	0.97	0.39	74,74,74,74	0
55	MG	1H	3137	1/1	0.97	0.10	72,72,72,72	0
55	MG	14	3472	1/1	0.97	0.05	103,103,103,103	0
55	MG	13	1630	1/1	0.97	0.27	65,65,65,65	0
55	MG	14	3093	1/1	0.97	0.32	60,60,60,60	0
55	MG	1G	1733	1/1	0.97	0.05	93,93,93,93	0
55	MG	1H	3127	1/1	0.97	0.40	58,58,58,58	0
55	MG	14	3207	1/1	0.97	0.48	92,92,92,92	0
55	MG	1H	3182	1/1	0.97	0.58	59,59,59,59	0
55	MG	1H	3297	1/1	0.97	0.29	87,87,87,87	0
55	MG	14	3063	1/1	0.97	0.34	69,69,69,69	0
55	MG	1H	3110	1/1	0.97	0.24	54,54,54,54	0
55	MG	14	3240	1/1	0.97	0.13	68,68,68,68	0
55	MG	14	3170	1/1	0.97	0.52	67,67,67,67	0
55	MG	14	3218	1/1	0.97	0.35	57,57,57,57	0
55	MG	1G	1626	1/1	0.97	0.34	80,80,80,80	0
55	MG	14	3214	1/1	0.97	0.38	78,78,78,78	0
55	MG	14	3180	1/1	0.97	0.28	82,82,82,82	0
55	MG	1H	3452	1/1	0.97	0.11	91,91,91,91	0
55	MG	1H	3022	1/1	0.97	0.37	57,57,57,57	0
55	MG	2L	101	1/1	0.97	0.39	82,82,82,82	0
55	MG	14	3469	1/1	0.97	0.07	96,96,96,96	0
55	MG	1H	3415	1/1	0.97	0.16	59,59,59,59	0
55	MG	1H	3136	1/1	0.97	0.47	72,72,72,72	0
55	MG	1H	3014	1/1	0.97	0.50	40,40,40,40	0
55	MG	14	3484	1/1	0.97	0.05	95,95,95,95	0
55	MG	14	3067	1/1	0.97	0.38	54,54,54,54	0
55	MG	14	3399	1/1	0.97	0.15	75,75,75,75	0
55	MG	1H	3049	1/1	0.97	0.35	89,89,89,89	0
55	MG	14	3407	1/1	0.97	0.16	66,66,66,66	0
55	MG	14	3083	1/1	0.97	0.42	68,68,68,68	0
55	MG	14	3327	1/1	0.97	0.10	98,98,98,98	0
55	MG	1H	3507	1/1	0.97	0.12	100,100,100,100	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	1H	3135	1/1	0.97	0.55	61,61,61,61	0
55	MG	13	1603	1/1	0.97	0.26	73,73,73,73	0
55	MG	1H	3420	1/1	0.97	0.06	99,99,99,99	0
55	MG	14	3077	1/1	0.97	0.46	63,63,63,63	0
55	MG	1H	3396	1/1	0.97	0.18	98,98,98,98	0
55	MG	14	3447	1/1	0.97	0.07	95,95,95,95	0
55	MG	14	3008	1/1	0.97	0.37	61,61,61,61	0
55	MG	1H	3150	1/1	0.97	0.46	84,84,84,84	0
55	MG	13	1614	1/1	0.97	0.27	92,92,92,92	0
55	MG	14	3041	1/1	0.97	0.33	53,53,53,53	0
55	MG	14	3024	1/1	0.97	0.38	79,79,79,79	0
55	MG	1G	1731	1/1	0.97	0.15	113,113,113,113	0
55	MG	1G	1603	1/1	0.98	0.49	65,65,65,65	0
55	MG	14	3069	1/1	0.98	0.36	65,65,65,65	0
55	MG	1H	3019	1/1	0.98	0.56	51,51,51,51	0
55	MG	1G	1709	1/1	0.98	0.26	81,81,81,81	0
55	MG	1H	3078	1/1	0.98	0.42	54,54,54,54	0
55	MG	1H	3427	1/1	0.98	0.08	91,91,91,91	0
55	MG	1H	3098	1/1	0.98	0.47	40,40,40,40	0
55	MG	13	1606	1/1	0.98	0.29	89,89,89,89	0
55	MG	13	1605	1/1	0.98	0.21	86,86,86,86	0
55	MG	1H	3439	1/1	0.98	0.11	59,59,59,59	0
56	ZN	32	302	1/1	0.98	0.34	100,100,100,100	0
55	MG	1H	3419	1/1	0.98	0.10	88,88,88,88	0
55	MG	1H	3458	1/1	0.98	0.06	67,67,67,67	0
55	MG	1H	3506	1/1	0.98	0.08	81,81,81,81	0
55	MG	14	3146	1/1	0.98	0.40	73,73,73,73	0
55	MG	14	3197	1/1	0.98	0.29	82,82,82,82	0
55	MG	1H	3221	1/1	0.98	0.70	57,57,57,57	0
55	MG	1H	3326	1/1	0.98	0.49	51,51,51,51	0
55	MG	1G	1622	1/1	0.98	0.37	95,95,95,95	0
55	MG	1H	3418	1/1	0.98	0.17	71,71,71,71	0
55	MG	13	1745	1/1	0.98	0.13	88,88,88,88	0
55	MG	1H	3359	1/1	0.98	0.59	45,45,45,45	0
55	MG	14	3047	1/1	0.98	0.25	67,67,67,67	0
55	MG	14	3090	1/1	0.98	0.43	58,58,58,58	0
56	ZN	3E	302	1/1	0.98	0.36	86,86,86,86	0
55	MG	14	3081	1/1	0.98	0.53	72,72,72,72	0
55	MG	1H	3097	1/1	0.98	0.47	48,48,48,48	0
55	MG	14	3300	1/1	0.98	0.28	64,64,64,64	0
55	MG	1H	3131	1/1	0.98	0.46	65,65,65,65	0
55	MG	13	1693	1/1	0.98	0.61	62,62,62,62	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	1J	201	1/1	0.98	0.18	115,115,115,115	0
55	MG	14	3091	1/1	0.98	0.58	53,53,53,53	0
55	MG	1H	3301	1/1	0.98	0.41	73,73,73,73	0
55	MG	14	3017	1/1	0.98	0.27	70,70,70,70	0
55	MG	1H	3016	1/1	0.98	0.26	70,70,70,70	0
55	MG	14	3095	1/1	0.98	0.36	54,54,54,54	0
55	MG	1H	3505	1/1	0.98	0.09	64,64,64,64	0
55	MG	1H	3072	1/1	0.98	0.31	33,33,33,33	0
55	MG	1G	1621	1/1	0.98	0.28	80,80,80,80	0
55	MG	1H	3239	1/1	0.98	0.35	93,93,93,93	0
55	MG	14	3065	1/1	0.98	0.58	47,47,47,47	0
55	MG	1G	1608	1/1	0.98	0.20	99,99,99,99	0
55	MG	13	1668	1/1	0.98	0.38	78,78,78,78	0
55	MG	14	3005	1/1	0.98	0.29	62,62,62,62	0
55	MG	29	302	1/1	0.98	0.41	65,65,65,65	0
55	MG	13	1602	1/1	0.98	0.28	85,85,85,85	0
55	MG	J8	101	1/1	0.98	0.47	65,65,65,65	0
55	MG	14	3346	1/1	0.98	0.14	62,62,62,62	0
55	MG	13	1632	1/1	0.98	0.45	74,74,74,74	0
55	MG	14	3006	1/1	0.98	0.42	54,54,54,54	0
55	MG	1G	1688	1/1	0.98	0.28	103,103,103,103	0
55	MG	1H	3417	1/1	0.98	0.15	70,70,70,70	0
55	MG	1H	3483	1/1	0.98	0.08	86,86,86,86	0
55	MG	1H	3002	1/1	0.98	0.39	53,53,53,53	0
55	MG	14	3172	1/1	0.98	0.38	69,69,69,69	0
55	MG	1G	1611	1/1	0.98	0.10	122,122,122,122	0
55	MG	14	3108	1/1	0.98	0.30	61,61,61,61	0
55	MG	14	3061	1/1	0.98	0.46	62,62,62,62	0
55	MG	14	3102	1/1	0.98	0.17	62,62,62,62	0
55	MG	1H	3119	1/1	0.98	0.71	75,75,75,75	0
55	MG	1H	3502	1/1	0.98	0.10	60,60,60,60	0
55	MG	14	3222	1/1	0.98	0.07	73,73,73,73	0
55	MG	1H	3488	1/1	0.98	0.10	66,66,66,66	0
55	MG	1H	3004	1/1	0.98	0.39	60,60,60,60	0
55	MG	14	3018	1/1	0.98	0.41	52,52,52,52	0
55	MG	1G	1699	1/1	0.98	0.39	100,100,100,100	0
55	MG	1H	3064	1/1	0.98	0.32	95,95,95,95	0
55	MG	1H	3236	1/1	0.98	0.18	72,72,72,72	0
55	MG	14	3410	1/1	0.98	0.13	68,68,68,68	0
55	MG	14	3421	1/1	0.98	0.20	55,55,55,55	0
55	MG	14	3076	1/1	0.98	0.42	45,45,45,45	0
55	MG	1H	3006	1/1	0.98	0.40	53,53,53,53	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	14	3009	1/1	0.98	0.44	62,62,62,62	0
55	MG	1G	1624	1/1	0.98	0.25	103,103,103,103	0
55	MG	14	3178	1/1	0.98	0.20	89,89,89,89	0
55	MG	1H	3111	1/1	0.98	0.56	51,51,51,51	0
55	MG	14	3344	1/1	0.98	0.15	73,73,73,73	0
55	MG	1H	3108	1/1	0.98	0.68	58,58,58,58	0
55	MG	14	3107	1/1	0.98	0.58	50,50,50,50	0
55	MG	1H	3255	1/1	0.98	0.21	72,72,72,72	0
55	MG	1H	3088	1/1	0.98	0.48	44,44,44,44	0
55	MG	5E	201	1/1	0.98	0.20	92,92,92,92	0
55	MG	1H	3011	1/1	0.98	0.33	49,49,49,49	0
55	MG	13	1601	1/1	0.98	0.36	63,63,63,63	0
55	MG	1H	3114	1/1	0.98	0.33	69,69,69,69	0
55	MG	1H	3128	1/1	0.98	0.47	66,66,66,66	0
55	MG	14	3426	1/1	0.98	0.11	84,84,84,84	0
55	MG	1H	3020	1/1	0.98	0.36	68,68,68,68	0
55	MG	1H	3103	1/1	0.98	0.43	48,48,48,48	0
55	MG	1H	3378	1/1	0.98	0.20	84,84,84,84	0
55	MG	1H	3060	1/1	0.98	0.35	59,59,59,59	0
55	MG	14	3151	1/1	0.98	0.41	68,68,68,68	0
55	MG	13	1628	1/1	0.98	0.47	64,64,64,64	0
55	MG	1G	1619	1/1	0.98	0.21	105,105,105,105	0
55	MG	14	3025	1/1	0.98	0.27	82,82,82,82	0
55	MG	14	3220	1/1	0.98	0.30	80,80,80,80	0
55	MG	1H	3054	1/1	0.98	0.52	51,51,51,51	0
55	MG	1H	3471	1/1	0.98	0.14	90,90,90,90	0
55	MG	1H	3106	1/1	0.98	0.22	47,47,47,47	0
55	MG	14	3048	1/1	0.98	0.44	66,66,66,66	0
55	MG	1H	3091	1/1	0.98	0.46	58,58,58,58	0
55	MG	14	3037	1/1	0.98	0.43	58,58,58,58	0
55	MG	14	3257	1/1	0.98	0.34	78,78,78,78	0
55	MG	14	3213	1/1	0.98	0.72	59,59,59,59	0
55	MG	1G	1730	1/1	0.98	0.13	77,77,77,77	0
55	MG	1H	3443	1/1	0.98	0.04	90,90,90,90	0
55	MG	14	3256	1/1	0.98	0.14	95,95,95,95	0
55	MG	1H	3146	1/1	0.98	0.29	64,64,64,64	0
55	MG	14	3436	1/1	0.98	0.08	69,69,69,69	0
55	MG	1H	3084	1/1	0.98	0.38	65,65,65,65	0
55	MG	14	3417	1/1	0.98	0.14	72,72,72,72	0
55	MG	1G	1602	1/1	0.98	0.38	80,80,80,80	0
55	MG	14	3021	1/1	0.99	0.41	60,60,60,60	0
55	MG	1H	3021	1/1	0.99	0.38	43,43,43,43	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors( $\text{\AA}^2$ )	Q<0.9
55	MG	14	3299	1/1	0.99	0.43	64,64,64,64	0
55	MG	1H	3440	1/1	0.99	0.10	66,66,66,66	0
55	MG	1G	1691	1/1	0.99	0.09	98,98,98,98	0
55	MG	1H	3512	1/1	0.99	0.31	72,72,72,72	0
55	MG	14	3396	1/1	0.99	0.11	79,79,79,79	0
55	MG	1H	3005	1/1	0.99	0.47	45,45,45,45	0
55	MG	14	3015	1/1	0.99	0.38	69,69,69,69	0
55	MG	14	3254	1/1	0.99	0.10	117,117,117,117	0
55	MG	1H	3027	1/1	0.99	0.43	51,51,51,51	0
55	MG	1H	3015	1/1	0.99	0.52	53,53,53,53	0
55	MG	1H	3003	1/1	0.99	0.45	58,58,58,58	0
55	MG	1H	3012	1/1	0.99	0.29	52,52,52,52	0
55	MG	14	3419	1/1	0.99	0.14	56,56,56,56	0
55	MG	14	3397	1/1	0.99	0.15	64,64,64,64	0
56	ZN	5I	101	1/1	0.99	0.10	109,109,109,109	0
55	MG	1H	3026	1/1	0.99	0.26	53,53,53,53	0
55	MG	14	3330	1/1	0.99	0.49	43,43,43,43	0
55	MG	1H	3177	1/1	0.99	0.28	93,93,93,93	0
55	MG	14	3013	1/1	0.99	0.30	66,66,66,66	0
55	MG	1H	3459	1/1	0.99	0.06	75,75,75,75	0
55	MG	1H	3055	1/1	0.99	0.25	66,66,66,66	0
55	MG	14	3302	1/1	0.99	0.16	85,85,85,85	0
55	MG	1H	3075	1/1	0.99	0.30	69,69,69,69	0
55	MG	14	3007	1/1	0.99	0.50	68,68,68,68	0
55	MG	14	3064	1/1	0.99	0.58	54,54,54,54	0
55	MG	1H	3077	1/1	0.99	0.28	65,65,65,65	0
55	MG	1H	3093	1/1	0.99	0.18	60,60,60,60	0
55	MG	1H	3437	1/1	0.99	0.13	60,60,60,60	0
55	MG	1H	3209	1/1	0.99	0.51	63,63,63,63	0
55	MG	1H	3030	1/1	0.99	0.37	63,63,63,63	0
55	MG	2K	101	1/1	0.99	0.28	85,85,85,85	0
55	MG	14	3415	1/1	0.99	0.09	71,71,71,71	0
55	MG	1H	3355	1/1	0.99	0.37	62,62,62,62	0
55	MG	14	3052	1/1	0.99	0.23	62,62,62,62	0
55	MG	14	3066	1/1	0.99	0.42	62,62,62,62	0
55	MG	1G	1605	1/1	0.99	0.23	85,85,85,85	0

## 6.5 Other polymers

There are no such residues in this entry.