



Full wwPDB X-ray Structure Validation Report ⓘ

Feb 16, 2017 – 11:02 am GMT

PDB ID : 4U56
Title : Crystal structure of Blasticidin S bound to the yeast 80S ribosome
Authors : Garreau de Loubresse, N.; Prokhorova, I.; Yusupova, G.; Yusupov, M.
Deposited on : 2014-07-24
Resolution : 3.45 Å(reported)

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

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

A user guide is available at

<http://wwpdb.org/validation/2016/XrayValidationReportHelp>

with specific help available everywhere you see the ⓘ symbol.

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

MolProbity : 4.02b-467
Mogul : 1.7.2 (RC1), CSD as538be (2017)
Xtriage (Phenix) : 1.9-1692
EDS : **FAILED**
Percentile statistics : 20161228.v01 (using entries in the PDB archive December 28th 2016)
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : recalc28986

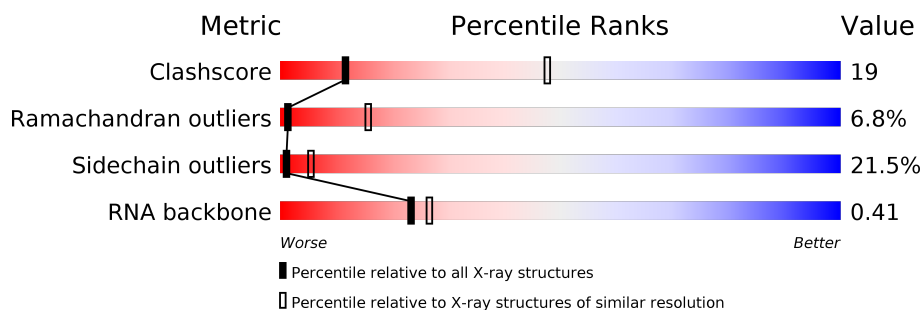
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

The reported resolution of this entry is 3.45 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))
Clashscore	112137	1040 (3.52-3.40)
Ramachandran outliers	110173	1009 (3.52-3.40)
Sidechain outliers	110143	1010 (3.52-3.40)
RNA backbone	2435	1020 (4.02-2.86)

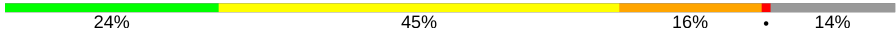



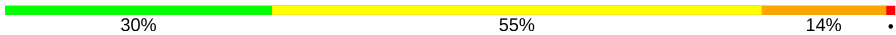



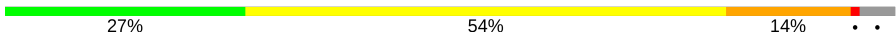

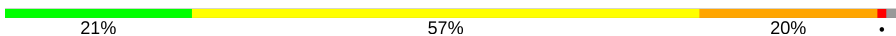

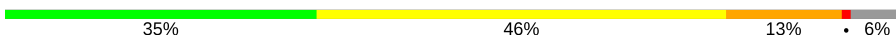

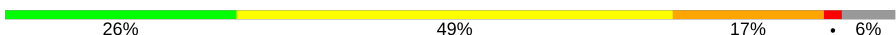

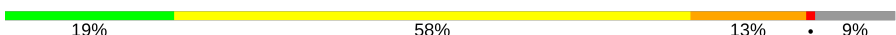



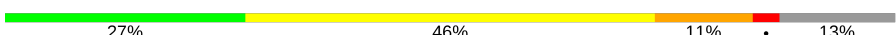




The table below summarises the geometric issues observed across the polymeric chains and their fit to the electron density. The red, orange, yellow and green segments on the lower bar indicate the fraction of residues that contain outliers for ≥ 3 , 2, 1 and 0 types of geometric quality criteria. A grey 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.

Note EDS failed to run properly.

Mol	Chain	Length	Quality of chain
1	2	1800	
1	6	1800	
2	S0	251	
2	s0	251	
3	S1	254	
3	s1	254	


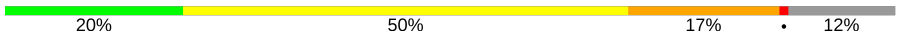

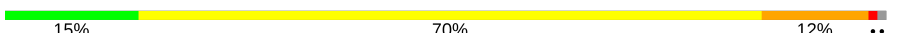

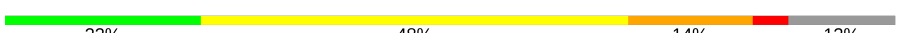





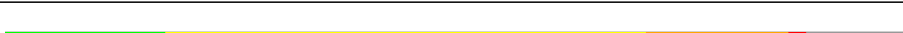


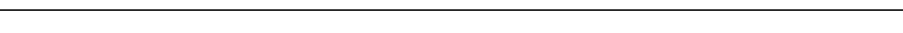



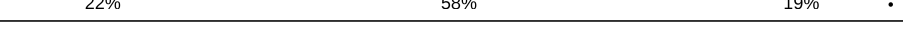
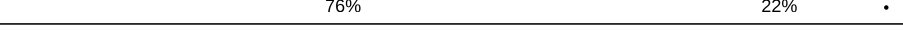




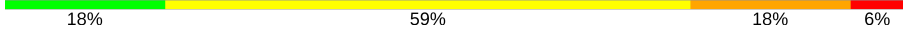
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Mol	Chain	Length	Quality of chain
4	S2	253	
4	s2	253	
5	S3	239	
5	s3	239	
6	S4	260	
6	s4	260	
7	S5	224	
7	s5	224	
8	S6	236	
8	s6	236	
9	S7	189	
9	s7	189	
10	S8	200	
10	s8	200	
11	S9	196	
11	s9	196	
12	C0	105	
12	c0	105	
13	C1	155	
13	c1	155	
14	C2	142	
14	c2	142	
15	C3	150	
15	c3	150	
16	C4	136	



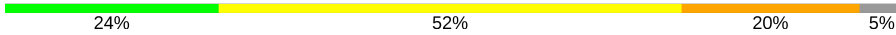

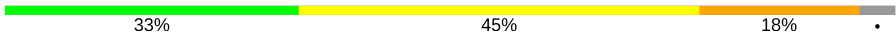

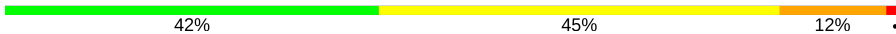
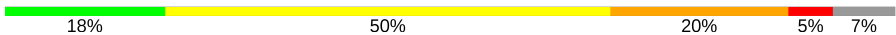

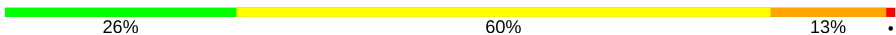





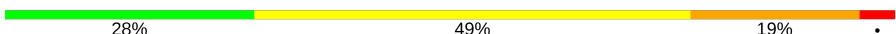

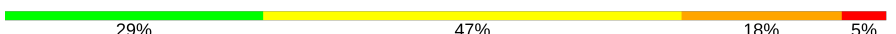
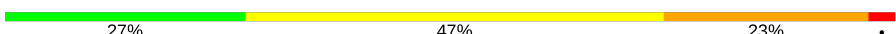
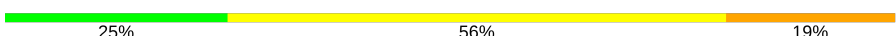





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Mol	Chain	Length	Quality of chain
16	c4	136	
17	C5	141	
17	c5	141	
18	C6	142	
18	c6	142	
19	C7	136	
19	c7	136	
20	C8	145	
20	c8	145	
21	C9	143	
21	c9	143	
22	D0	120	
22	d0	120	
23	D1	87	
23	d1	87	
24	D2	129	
24	d2	129	
25	D3	144	
25	d3	144	
26	D4	134	
26	d4	134	
27	D5	107	
27	d5	107	
28	D6	97	
28	d6	97	





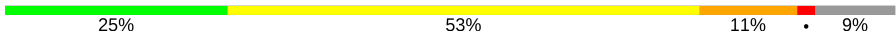

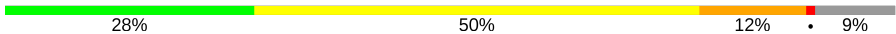

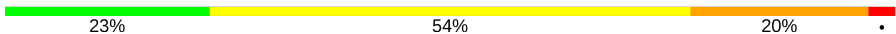

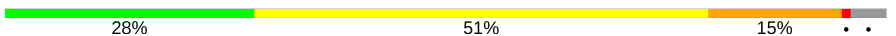

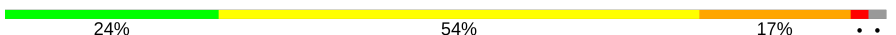





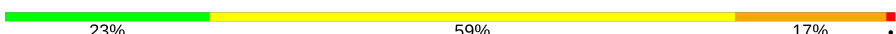

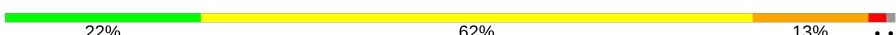




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Mol	Chain	Length	Quality of chain
29	D7	81	
29	d7	81	
30	D8	66	
30	d8	66	
31	D9	55	
31	d9	55	
32	E0	60	
33	E1	76	
33	e1	76	
34	SR	318	
34	sR	318	
35	SM	273	
35	sM	273	
36	1	3396	
36	5	3396	
37	3	121	
37	7	121	
38	4	158	
38	8	158	
39	L2	253	
39	l2	253	
40	L3	386	
40	l3	386	
41	L4	361	
41	l4	361	


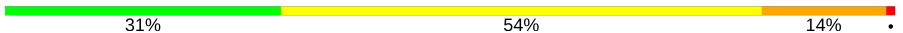



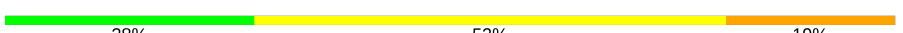





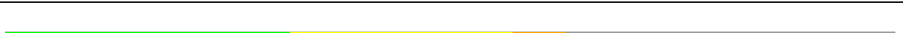


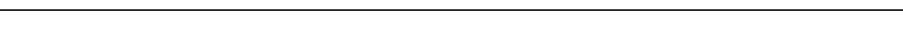

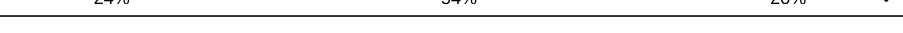


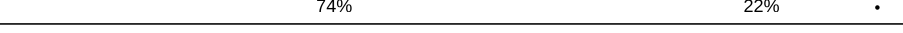


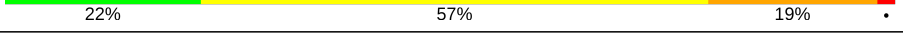

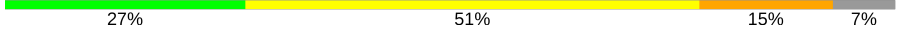
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Mol	Chain	Length	Quality of chain
42	L5	296	
42	l5	296	
43	L6	175	
43	l6	175	
44	L7	243	
44	l7	243	
45	L8	255	
45	l8	255	
46	L9	191	
46	l9	191	
47	M0	220	
47	m0	220	
48	M1	173	
48	m1	173	
49	M3	198	
49	m3	198	
50	M4	137	
50	m4	137	
51	M5	203	
51	m5	203	
52	M6	198	
52	m6	198	
53	M7	183	
53	m7	183	
54	M8	185	





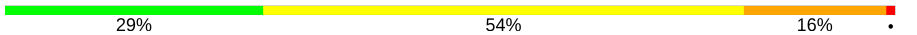

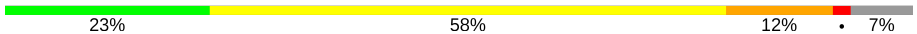

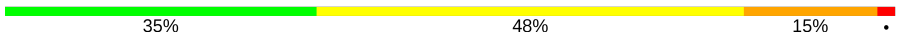

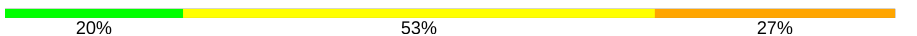

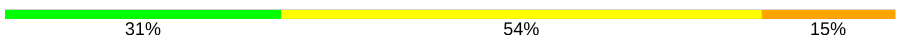







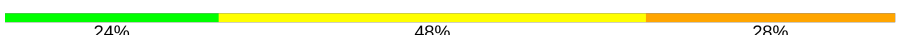



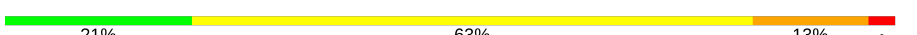
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Mol	Chain	Length	Quality of chain
54	m8	185	
55	M9	188	
55	m9	188	
56	N0	172	
56	n0	172	
57	N1	159	
57	n1	159	
58	N2	120	
58	n2	120	
59	N3	136	
59	n3	136	
60	N4	155	
60	n4	155	
61	N5	141	
61	n5	141	
62	N6	126	
62	n6	126	
63	N7	135	
63	n7	135	
64	N8	148	
64	n8	148	
65	N9	58	
65	n9	58	
66	O0	104	
66	o0	104	




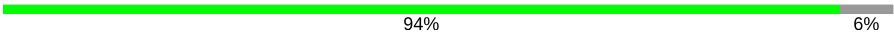


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Mol	Chain	Length	Quality of chain
67	O1	112	
67	o1	112	
68	O2	129	
68	o2	129	
69	O3	106	
69	o3	106	
70	O4	120	
70	o4	120	
71	O5	119	
71	o5	119	
72	O6	99	
72	o6	99	
73	O7	87	
73	o7	87	
74	O8	77	
74	o8	77	
75	O9	50	
75	o9	50	
76	Q0	52	
76	q0	52	
77	Q1	25	
77	q1	25	
78	Q2	105	
78	q2	105	
79	Q3	91	

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Mol	Chain	Length	Quality of chain
79	q3	91	
80	e0	62	
81	p0	311	
82	m2	160	
83	p1	47	
84	p2	46	

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
86	OHX	1	3951	-	-	X	-
86	OHX	1	3965	-	-	X	-
86	OHX	1	3998	-	-	X	-
86	OHX	1	4023	-	-	X	-
86	OHX	1	4027	-	-	X	-
86	OHX	1	4040	-	-	X	-
86	OHX	1	4050	-	-	X	-
86	OHX	1	4143	-	-	X	-
86	OHX	1	4153	-	-	X	-
86	OHX	1	4160	-	-	X	-
86	OHX	1	4169	-	-	X	-
86	OHX	1	4178	-	-	X	-
86	OHX	1	4194	-	-	X	-
86	OHX	2	2031	-	-	X	-
86	OHX	2	2043	-	-	X	-
86	OHX	2	2074	-	-	X	-
86	OHX	2	2098	-	-	X	-
86	OHX	2	2131	-	-	X	-
86	OHX	2	2146	-	-	X	-
86	OHX	2	2162	-	-	X	-
86	OHX	5	3935	-	-	X	-
86	OHX	5	3966	-	-	X	-
86	OHX	5	3971	-	-	X	-
86	OHX	5	3994	-	-	X	-
86	OHX	5	3995	-	-	X	-
86	OHX	5	4004	-	-	X	-
86	OHX	5	4014	-	-	X	-

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
86	OHX	5	4029	-	-	X	-
86	OHX	5	4060	-	-	X	-
86	OHX	5	4085	-	-	X	-
86	OHX	5	4193	-	-	X	-
86	OHX	5	4195	-	-	X	-
86	OHX	5	4212	-	-	X	-
86	OHX	5	4228	-	-	X	-
86	OHX	5	4232	-	-	X	-
86	OHX	5	4238	-	-	X	-
86	OHX	6	2059	-	-	X	-
86	OHX	6	2120	-	-	X	-
86	OHX	6	2147	-	-	X	-
86	OHX	6	2171	-	-	X	-
86	OHX	7	220	-	-	X	-
86	OHX	7	228	-	-	X	-
86	OHX	8	216	-	-	X	-
86	OHX	8	223	-	-	X	-
86	OHX	O1	202	-	-	X	-
86	OHX	O7	104	-	-	X	-
87	ZN	Q2	501	-	-	X	-

2 Entry composition

There are 88 unique types of molecules in this entry. The entry contains 411214 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 18S ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
1	2	1750	Total	C	N	O	P	0	0	0
			37283	16668	6591	12274	1750			
1	6	1795	Total	C	N	O	P	0	0	0
			38238	17095	6758	12590	1795			

- Molecule 2 is a protein called 40S ribosomal protein S0-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
2	S0	206	Total	C	N	O	S	0	0	0
			1577	1014	278	283	2			
2	s0	206	Total	C	N	O	S	0	0	0
			1583	1017	281	283	2			

- Molecule 3 is a protein called 40S ribosomal protein S1-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
3	S1	214	Total	C	N	O	S	0	0	0
			1709	1084	310	311	4			
3	s1	216	Total	C	N	O	S	0	0	0
			1722	1091	312	315	4			

- Molecule 4 is a protein called 40S ribosomal protein S2.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
4	S2	217	Total	C	N	O	S	0	0	0
			1635	1047	289	297	2			
4	s2	217	Total	C	N	O	S	0	0	0
			1635	1047	289	297	2			

- Molecule 5 is a protein called 40S ribosomal protein S3.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
5	S3	223	Total	C	N	O	S	0	0	0
			1734	1101	313	314	6			
5	s3	223	Total	C	N	O	S	0	0	0
			1734	1101	313	314	6			

- Molecule 6 is a protein called 40S ribosomal protein S4-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
6	S4	260	Total	C	N	O	S	0	0	0
			2068	1316	389	360	3			
6	s4	260	Total	C	N	O	S	0	0	0
			2068	1316	389	360	3			

- Molecule 7 is a protein called 40S ribosomal protein S5.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
7	S5	206	Total	C	N	O	S	0	0	0
			1609	1007	300	299	3			
7	s5	206	Total	C	N	O	S	0	0	0
			1609	1007	300	299	3			

- Molecule 8 is a protein called 40S ribosomal protein S6-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
8	S6	226	Total	C	N	O	S	0	0	0
			1799	1129	346	321	3			
8	s6	218	Total	C	N	O	S	0	0	0
			1755	1102	337	313	3			

- Molecule 9 is a protein called 40S ribosomal protein S7-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
9	S7	184	Total	C	N	O		0	0	0
			1481	951	265	265				
9	s7	186	Total	C	N	O		0	0	0
			1491	957	267	267				

- Molecule 10 is a protein called 40S ribosomal protein S8-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
10	S8	188	Total	C	N	O	S	0	0	0
			1489	925	298	264	2			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
10	s8	188	Total	C	N	O	S	0	0	0
			1489	925	298	264	2			

- Molecule 11 is a protein called 40S ribosomal protein S9-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
11	S9	185	Total	C	N	O	S	0	0	0
			1494	943	289	261	1			
11	s9	185	Total	C	N	O	S	0	0	0
			1494	943	289	261	1			

- Molecule 12 is a protein called 40S ribosomal protein S10-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
12	C0	96	Total	C	N	O	S	0	0	0
			773	500	126	145	2			
12	c0	96	Total	C	N	O	S	0	0	0
			762	491	125	144	2			

There are 2 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
C0	89	ALA	GLY	conflict	UNP Q08745
c0	89	ALA	GLY	conflict	UNP Q08745

- Molecule 13 is a protein called 40S ribosomal protein S11-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
13	C1	155	Total	C	N	O	S	0	0	0
			1214	775	230	206	3			
13	c1	146	Total	C	N	O	S	0	0	0
			1168	747	221	197	3			

There are 2 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
C1	147	ALA	GLY	conflict	UNP P0CX47
c1	147	ALA	GLY	conflict	UNP P0CX47

- Molecule 14 is a protein called 40S ribosomal protein S12.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
14	C2	124	Total	C	N	O	S	0	0	0
			892	562	156	172	2			
14	c2	124	Total	C	N	O	S	0	0	0
			892	562	156	172	2			

There are 4 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
C2	104	ALA	GLY	conflict	UNP P48589
C2	110	ALA	GLY	conflict	UNP P48589
c2	104	ALA	GLY	conflict	UNP P48589
c2	110	ALA	GLY	conflict	UNP P48589

- Molecule 15 is a protein called 40S ribosomal protein S13.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
15	C3	150	Total	C	N	O	S	0	0	0
			1192	759	224	207	2			
15	c3	150	Total	C	N	O	S	0	0	0
			1192	759	224	207	2			

- Molecule 16 is a protein called 40S ribosomal protein S14-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
16	C4	127	Total	C	N	O	S	0	0	0
			891	545	182	163	1			
16	c4	128	Total	C	N	O	S	0	0	0
			949	582	188	176	3			

- Molecule 17 is a protein called 40S ribosomal protein S15.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
17	C5	124	Total	C	N	O	S	0	0	0
			977	622	182	166	7			
17	c5	135	Total	C	N	O	S	0	0	0
			1039	658	196	178	7			

There are 2 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
C5	137	SER	ARG	conflict	UNP Q01855
c5	137	SER	ARG	conflict	UNP Q01855

- Molecule 18 is a protein called 40S ribosomal protein S16-A.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
18	C6	141	Total	C	N	O	0	0	0
			1105	708	203	194			
18	c6	142	Total	C	N	O	0	0	0
			1111	711	204	196			

- Molecule 19 is a protein called 40S ribosomal protein S17-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
19	C7	120	Total	C	N	O	S	0	0	0
			926	577	177	170	2			
19	c7	117	Total	C	N	O	S	0	0	0
			906	563	174	167	2			

- Molecule 20 is a protein called 40S ribosomal protein S18-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
20	C8	145	Total	C	N	O	S	0	0	0
			1192	743	237	210	2			
20	c8	145	Total	C	N	O	S	0	0	0
			1192	743	237	210	2			

- Molecule 21 is a protein called 40S ribosomal protein S19-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
21	C9	143	Total	C	N	O	S	0	0	0
			1112	694	208	208	2			
21	c9	143	Total	C	N	O	S	0	0	0
			1112	694	208	208	2			

- Molecule 22 is a protein called 40S ribosomal protein S20.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
22	D0	107	Total	C	N	O	S	0	0	0
			855	539	156	159	1			
22	d0	110	Total	C	N	O	S	0	0	0
			882	554	161	166	1			

- Molecule 23 is a protein called 40S ribosomal protein S21-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
23	D1	87	Total	C	N	O	S	0	0	0
			684	420	125	137	2			
23	d1	87	Total	C	N	O	S	0	0	0
			684	420	125	137	2			

- Molecule 24 is a protein called 40S ribosomal protein S22-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
24	D2	129	Total	C	N	O	S	0	0	0
			1021	650	188	180	3			
24	d2	129	Total	C	N	O	S	0	0	0
			1021	650	188	180	3			

- Molecule 25 is a protein called 40S ribosomal protein S23-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
25	D3	144	Total	C	N	O	S	0	0	0
			1121	708	220	191	2			
25	d3	144	Total	C	N	O	S	0	0	0
			1121	708	220	191	2			

- Molecule 26 is a protein called 40S ribosomal protein S24-A.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
26	D4	134	Total	C	N	O	0	0	0
			1073	676	208	189			
26	d4	134	Total	C	N	O	0	0	0
			1073	676	208	189			

- Molecule 27 is a protein called 40S ribosomal protein S25-A.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
27	D5	70	Total	C	N	O	0	0	0
			563	360	104	99			
27	d5	69	Total	C	N	O	0	0	0
			558	357	103	98			

- Molecule 28 is a protein called 40S ribosomal protein S26-B.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
28	D6	97	Total	C	N	O	S	0	0	0
			769	475	160	129	5			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
28	d6	97	Total	C	N	O	S	0	0	0
			769	475	160	129	5			

- Molecule 29 is a protein called 40S ribosomal protein S27-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
29	D7	81	Total	C	N	O	S	0	0	0
			610	382	110	113	5			
29	d7	81	Total	C	N	O	S	0	0	0
			610	382	110	113	5			

- Molecule 30 is a protein called 40S ribosomal protein S28-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
30	D8	63	Total	C	N	O	S	0	0	0
			497	306	99	91	1			
30	d8	63	Total	C	N	O	S	0	0	0
			497	306	99	91	1			

- Molecule 31 is a protein called 40S ribosomal protein S29-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
31	D9	53	Total	C	N	O	S	0	0	0
			442	274	92	72	4			
31	d9	53	Total	C	N	O	S	0	0	0
			442	274	92	72	4			

- Molecule 32 is a protein called 40S ribosomal protein S30-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
32	E0	60	Total	C	N	O	S	0	0	0
			475	299	98	77	1			

- Molecule 33 is a protein called Ubiquitin-40S ribosomal protein S31.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
33	E1	71	Total	C	N	O	S	0	0	0
			566	362	106	94	4			
33	e1	76	Total	C	N	O	S	0	0	0
			608	388	117	99	4			

- Molecule 34 is a protein called Guanine nucleotide-binding protein subunit beta-like protein.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
34	SR	318	Total	C	N	O	S	0	0	0
			2441	1544	419	470	8			
34	sR	318	Total	C	N	O	S	0	0	0
			2442	1544	418	472	8			

- Molecule 35 is a protein called Suppressor protein STM1.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
35	SM	159	Total	C	N	O		0	0	0
			1104	652	221	231				
35	sM	104	Total	C	N	O		0	0	0
			679	402	140	137				

- Molecule 36 is a RNA chain called 25S ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
36	1	3149	Total	C	N	O	P	0	0	0
			67355	30086	12142	21978	3149			
36	5	3150	Total	C	N	O	P	0	0	0
			67376	30095	12145	21987	3149			

- Molecule 37 is a RNA chain called 5S ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
37	3	121	Total	C	N	O	P	0	0	0
			2579	1152	461	845	121			
37	7	121	Total	C	N	O	P	0	0	0
			2579	1152	461	845	121			

- Molecule 38 is a RNA chain called 5.8S ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
38	4	158	Total	C	N	O	P	0	0	0
			3353	1500	586	1109	158			
38	8	158	Total	C	N	O	P	0	0	0
			3353	1500	586	1109	158			

- Molecule 39 is a protein called 60S ribosomal protein L2-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
39	L2	252	Total	C	N	O	S	0	0	0
			1914	1191	388	334	1			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
39	l2	252	Total	C	N	O	S	0	0	0
			1912	1190	388	333	1			

- Molecule 40 is a protein called 60S ribosomal protein L3.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
40	L3	386	Total	C	N	O	S	0	0	0
			3075	1950	584	533	8			
40	l3	386	Total	C	N	O	S	0	0	0
			3075	1950	584	533	8			

- Molecule 41 is a protein called 60S ribosomal protein L4-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
41	L4	361	Total	C	N	O	S	0	0	0
			2748	1729	522	494	3			
41	l4	361	Total	C	N	O	S	0	0	0
			2748	1729	522	494	3			

- Molecule 42 is a protein called 60S ribosomal protein L5.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
42	L5	296	Total	C	N	O	S	0	0	0
			2375	1501	414	458	2			
42	l5	294	Total	C	N	O	S	0	0	0
			2359	1489	412	456	2			

- Molecule 43 is a protein called 60S ribosomal protein L6-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
43	L6	156	Total	C	N	O	S	0	0	0
			1239	800	222	216	1			
43	l6	157	Total	C	N	O	S	0	0	0
			1248	806	224	217	1			

- Molecule 44 is a protein called 60S ribosomal protein L7-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
44	L7	222	Total	C	N	O	S	0	0	0
			1784	1151	324	308	1			
44	l7	223	Total	C	N	O	S	0	0	0
			1791	1155	325	310	1			

- Molecule 45 is a protein called 60S ribosomal protein L8-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
45	L8	233	Total	C	N	O	S	0	0	0
			1804	1151	323	327	3			
45	l8	231	Total	C	N	O	S	0	0	0
			1763	1130	316	314	3			

- Molecule 46 is a protein called 60S ribosomal protein L9-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
46	L9	191	Total	C	N	O	S	0	0	0
			1518	963	274	277	4			
46	l9	191	Total	C	N	O	S	0	0	0
			1518	963	274	277	4			

- Molecule 47 is a protein called 60S ribosomal protein L10.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
47	M0	211	Total	C	N	O	S	0	0	0
			1705	1083	322	294	6			
47	m0	213	Total	C	N	O	S	0	0	0
			1722	1094	325	297	6			

- Molecule 48 is a protein called 60S ribosomal protein L11-B.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
48	M1	169	Total	C	N	O	S	0	0	0
			1353	847	253	249	4			
48	m1	169	Total	C	N	O	S	0	0	0
			1353	847	253	249	4			

- Molecule 49 is a protein called 60S ribosomal protein L13-A.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
49	M3	193	Total	C	N	O	0	0	0
			1543	962	315	266			
49	m3	194	Total	C	N	O	0	0	0
			1548	965	316	267			

- Molecule 50 is a protein called 60S ribosomal protein L14-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
50	M4	136	Total	C	N	O	S	0	0	0
			1053	675	199	177	2			
50	m4	137	Total	C	N	O	S	0	0	0
			1059	678	200	179	2			

- Molecule 51 is a protein called 60S ribosomal protein L15-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
51	M5	203	Total	C	N	O	S	0	0	0
			1720	1077	361	281	1			
51	m5	203	Total	C	N	O	S	0	0	0
			1720	1077	361	281	1			

- Molecule 52 is a protein called 60S ribosomal protein L16-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
52	M6	197	Total	C	N	O	S	0	0	0
			1555	1003	289	262	1			
52	m6	197	Total	C	N	O	S	0	0	0
			1555	1003	289	262	1			

- Molecule 53 is a protein called 60S ribosomal protein L17-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
53	M7	183	Total	C	N	O	S	0	0	0
			1420	882	281	257				
53	m7	155	Total	C	N	O	S	0	0	0
			1227	764	238	225				

- Molecule 54 is a protein called 60S ribosomal protein L18-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
54	M8	185	Total	C	N	O	S	0	0	0
			1441	908	290	241	2			
54	m8	185	Total	C	N	O	S	0	0	0
			1441	908	290	241	2			

- Molecule 55 is a protein called 60S ribosomal protein L19-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
55	M9	188	Total	C	N	O	S	0	0	0
			1521	935	326	260				

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Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
55	m9	188	Total	C	N	O	0	0	0
			1521	935	326	260			

- Molecule 56 is a protein called 60S ribosomal protein L20-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
56	N0	172	Total	C	N	O	S	0	0	0
			1445	930	267	244	4			
56	n0	172	Total	C	N	O	S	0	0	0
			1445	930	267	244	4			

- Molecule 57 is a protein called 60S ribosomal protein L21-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
57	N1	159	Total	C	N	O	S	0	0	0
			1276	805	246	221	4			
57	n1	159	Total	C	N	O	S	0	0	0
			1276	805	246	221	4			

- Molecule 58 is a protein called 60S ribosomal protein L22-A.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
58	N2	100	Total	C	N	O	0	0	0
			796	516	131	149			
58	n2	98	Total	C	N	O	0	0	0
			778	505	127	146			

- Molecule 59 is a protein called 60S ribosomal protein L23-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
59	N3	136	Total	C	N	O	S	0	0	0
			1003	628	189	179	7			
59	n3	136	Total	C	N	O	S	0	0	0
			1003	628	189	179	7			

- Molecule 60 is a protein called 60S ribosomal protein L24-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
60	N4	98	Total	C	N	O	S	0	0	0
			699	443	137	118	1			
60	n4	135	Total	C	N	O	S	0	0	0
			1038	651	206	180	1			

- Molecule 61 is a protein called 60S ribosomal protein L25.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
61	N5	121	Total	C	N	O	S	0	0	0
			964	620	169	173	2			
61	n5	120	Total	C	N	O	S	0	0	0
			959	617	168	172	2			

- Molecule 62 is a protein called 60S ribosomal protein L26-A.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
62	N6	126	Total	C	N	O	0	0	0
			993	625	192	176			
62	n6	126	Total	C	N	O	0	0	0
			993	625	192	176			

- Molecule 63 is a protein called 60S ribosomal protein L27-A.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
63	N7	135	Total	C	N	O	0	0	0
			1092	710	202	180			
63	n7	135	Total	C	N	O	0	0	0
			1092	710	202	180			

- Molecule 64 is a protein called 60S ribosomal protein L28.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
64	N8	148	Total	C	N	O	S	0	0	0
			1173	749	231	190	3			
64	n8	148	Total	C	N	O	S	0	0	0
			1173	749	231	190	3			

- Molecule 65 is a protein called 60S ribosomal protein L29.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
65	N9	58	Total	C	N	O	0	0	0
			462	289	100	73			
65	n9	58	Total	C	N	O	0	0	0
			462	289	100	73			

- Molecule 66 is a protein called 60S ribosomal protein L30.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
66	O0	97	Total	C	N	O	S	0	0	0
			743	479	124	139	1			
66	o0	100	Total	C	N	O	S	0	0	0
			767	492	128	146	1			

- Molecule 67 is a protein called 60S ribosomal protein L31-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
67	O1	109	Total	C	N	O	S	0	0	0
			876	556	167	152	1			
67	o1	109	Total	C	N	O	S	0	0	0
			883	559	167	156	1			

- Molecule 68 is a protein called 60S ribosomal protein L32.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
68	O2	127	Total	C	N	O	S	0	0	0
			1020	647	205	167	1			
68	o2	127	Total	C	N	O	S	0	0	0
			1020	647	205	167	1			

- Molecule 69 is a protein called 60S ribosomal protein L33-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
69	O3	106	Total	C	N	O	S	0	0	0
			850	540	165	144	1			
69	o3	106	Total	C	N	O	S	0	0	0
			850	540	165	144	1			

- Molecule 70 is a protein called 60S ribosomal protein L34-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
70	O4	112	Total	C	N	O	S	0	0	0
			880	545	179	152	4			
70	o4	112	Total	C	N	O	S	0	0	0
			880	545	179	152	4			

There are 2 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
O4	121	LYS	-	expression tag	UNP P87262
o4	121	LYS	-	expression tag	UNP P87262

- Molecule 71 is a protein called 60S ribosomal protein L35-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
71	O5	119	Total	C	N	O	S	0	0	0
			969	615	186	167	1			
71	o5	119	Total	C	N	O	S	0	0	0
			965	612	185	167	1			

- Molecule 72 is a protein called 60S ribosomal protein L36-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
72	O6	99	Total	C	N	O	S	0	0	0
			771	481	156	132	2			
72	o6	99	Total	C	N	O	S	0	0	0
			770	481	156	131	2			

- Molecule 73 is a protein called 60S ribosomal protein L37-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
73	O7	87	Total	C	N	O	S	0	0	0
			681	414	148	114	5			
73	o7	87	Total	C	N	O	S	0	0	0
			681	414	148	114	5			

- Molecule 74 is a protein called 60S ribosomal protein L38.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
74	O8	77	Total	C	N	O	0	0	0
			612	391	115	106			
74	o8	77	Total	C	N	O	0	0	0
			608	388	114	106			

- Molecule 75 is a protein called 60S ribosomal protein L39.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
75	O9	50	Total	C	N	O	S	0	0	0
			436	272	97	65	2			
75	o9	50	Total	C	N	O	S	0	0	0
			436	272	97	65	2			

- Molecule 76 is a protein called Ubiquitin-60S ribosomal protein L40.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
76	Q0	52	Total	C	N	O	S	0	0	0
			417	259	86	67	5			
76	q0	52	Total	C	N	O	S	0	0	0
			417	259	86	67	5			

- Molecule 77 is a protein called 60S ribosomal protein L41-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
77	Q1	25	Total	C	N	O	S	0	0	0
			233	142	63	27	1			
77	q1	25	Total	C	N	O	S	0	0	0
			233	142	63	27	1			

- Molecule 78 is a protein called 60S ribosomal protein L42-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
78	Q2	105	Total	C	N	O	S	0	0	0
			847	534	170	138	5			
78	q2	105	Total	C	N	O	S	0	0	0
			847	534	170	138	5			

- Molecule 79 is a protein called 60S ribosomal protein L43-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
79	Q3	91	Total	C	N	O	S	0	0	0
			694	429	138	121	6			
79	q3	91	Total	C	N	O	S	0	0	0
			694	429	138	121	6			

- Molecule 80 is a protein called 40S ribosomal protein S30-A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
80	e0	62	Total	C	N	O	S	0	0	0
			491	309	101	80	1			

- Molecule 81 is a protein called 60S acidic ribosomal protein P0.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
81	p0	143	Total	C	N	O	S	0	0	0
			1076	686	192	195	3			

- Molecule 82 is a protein called unknown protein chain m2.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
82	m2	150	Total	C	N	O	0	0	0
			750	450	150	150			

- Molecule 83 is a protein called unknown protein chain p1.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
83	p1	47	Total	C	N	O	0	0	0
			235	141	47	47			

- Molecule 84 is a protein called unknown protein chain p2.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
84	p2	46	Total	C	N	O	0	0	0
			230	138	46	46			

- Molecule 85 is MAGNESIUM ION (three-letter code: MG) (formula: Mg).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
85	L7	2	Total	Mg	0	0
			2	2		
85	m6	2	Total	Mg	0	0
			2	2		
85	n8	3	Total	Mg	0	0
			3	3		
85	N5	1	Total	Mg	0	0
			1	1		
85	6	147	Total	Mg	0	0
			147	147		
85	sM	1	Total	Mg	0	0
			1	1		
85	m5	5	Total	Mg	0	0
			5	5		
85	l3	3	Total	Mg	0	0
			3	3		
85	C1	1	Total	Mg	0	0
			1	1		
85	M1	1	Total	Mg	0	0
			1	1		
85	d6	1	Total	Mg	0	0
			1	1		
85	2	122	Total	Mg	0	0
			122	122		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
85	n0	1	Total 1	Mg 1	0	0
85	L4	3	Total 3	Mg 3	0	0
85	l7	1	Total 1	Mg 1	0	0
85	M5	2	Total 2	Mg 2	0	0
85	c9	1	Total 1	Mg 1	0	0
85	L8	1	Total 1	Mg 1	0	0
85	D3	1	Total 1	Mg 1	0	0
85	M9	1	Total 1	Mg 1	0	0
85	q0	2	Total 2	Mg 2	0	0
85	SM	1	Total 1	Mg 1	0	0
85	o4	3	Total 3	Mg 3	0	0
85	M0	3	Total 3	Mg 3	0	0
85	c1	1	Total 1	Mg 1	0	0
85	n6	2	Total 2	Mg 2	0	0
85	5	495	Total 495	Mg 495	0	0
85	c8	1	Total 1	Mg 1	0	0
85	O7	2	Total 2	Mg 2	0	0
85	Q2	1	Total 1	Mg 1	0	0
85	n9	2	Total 2	Mg 2	0	0
85	1	462	Total 462	Mg 462	0	0
85	c4	1	Total 1	Mg 1	0	0

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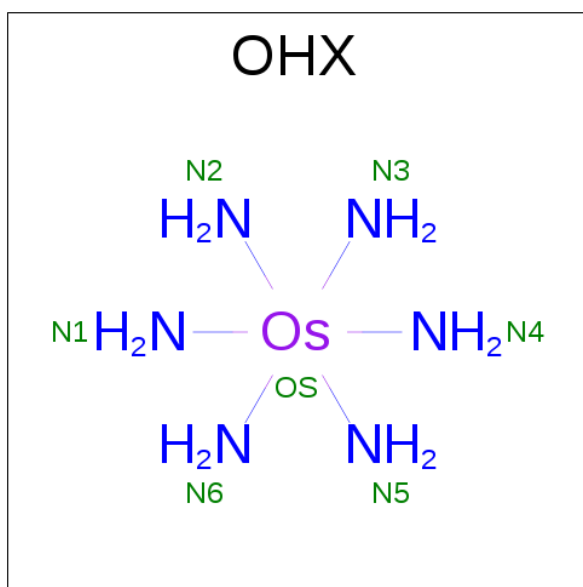
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
85	D0	1	Total 1	Mg 1	0	0
85	S8	1	Total 1	Mg 1	0	0
85	l2	3	Total 3	Mg 3	0	0
85	O2	2	Total 2	Mg 2	0	0
85	o7	1	Total 1	Mg 1	0	0
85	o3	2	Total 2	Mg 2	0	0
85	d3	3	Total 3	Mg 3	0	0
85	M3	2	Total 2	Mg 2	0	0
85	N3	3	Total 3	Mg 3	0	0
85	4	23	Total 23	Mg 23	0	0
85	D4	1	Total 1	Mg 1	0	0
85	S4	1	Total 1	Mg 1	0	0
85	L2	2	Total 2	Mg 2	0	0
85	m1	1	Total 1	Mg 1	0	0
85	l5	3	Total 3	Mg 3	0	0
85	d0	1	Total 1	Mg 1	0	0
85	M7	5	Total 5	Mg 5	0	0
85	m4	1	Total 1	Mg 1	0	0
85	N8	5	Total 5	Mg 5	0	0
85	s1	1	Total 1	Mg 1	0	0
85	l9	1	Total 1	Mg 1	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
85	O1	1	Total 1	Mg 1	0	0
85	s8	1	Total 1	Mg 1	0	0
85	c7	1	Total 1	Mg 1	0	0
85	7	17	Total 17	Mg 17	0	0
85	n3	1	Total 1	Mg 1	0	0
85	q1	1	Total 1	Mg 1	0	0
85	L3	4	Total 4	Mg 4	0	0
85	O5	1	Total 1	Mg 1	0	0
85	N6	2	Total 2	Mg 2	0	0
85	8	14	Total 14	Mg 14	0	0
85	l4	1	Total 1	Mg 1	0	0
85	M6	1	Total 1	Mg 1	0	0
85	N0	1	Total 1	Mg 1	0	0
85	m0	1	Total 1	Mg 1	0	0
85	3	15	Total 15	Mg 15	0	0
85	m7	6	Total 6	Mg 6	0	0

- Molecule 86 is osmium (III) hexammine (three-letter code: OHX) (formula: H₁₂N₆Os).



Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
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86	2	1	Total	N	Os	0	0
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86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	2	1	Total	N	Os	0	0
			7	6	1		
86	S8	1	Total	N	Os	0	0
			7	6	1		
86	C3	1	Total	N	Os	0	0
			7	6	1		
86	C5	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	C8	1	Total	N	Os	0	0
			7	6	1		
86	D3	1	Total	N	Os	0	0
			7	6	1		
86	D9	1	Total	N	Os	0	0
			7	6	1		
86	SR	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
			7	6	1		
86	1	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	1	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	1	1	Total	N	Os	0	0
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86	1	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	L3	1	Total	N	Os	0	0
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86	L4	1	Total	N	Os	0	0
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86	M0	1	Total	N	Os	0	0
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86	M5	1	Total	N	Os	0	0
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86	M7	1	Total	N	Os	0	0
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86	M9	1	Total	N	Os	0	0
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86	O3	1	Total	N	Os	0	0
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86	O7	1	Total	N	Os	0	0
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86	Q2	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	6	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	6	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	6	1	Total	N	Os	0	0
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86	6	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	6	1	Total 7	N 6	Os 1	0	0
86	6	1	Total 7	N 6	Os 1	0	0
86	6	1	Total 7	N 6	Os 1	0	0
86	6	1	Total 7	N 6	Os 1	0	0
86	6	1	Total 7	N 6	Os 1	0	0
86	6	1	Total 7	N 6	Os 1	0	0
86	6	1	Total 7	N 6	Os 1	0	0
86	6	1	Total 7	N 6	Os 1	0	0
86	6	1	Total 7	N 6	Os 1	0	0
86	6	1	Total 7	N 6	Os 1	0	0
86	s1	1	Total 7	N 6	Os 1	0	0
86	s1	1	Total 7	N 6	Os 1	0	0
86	s4	1	Total 7	N 6	Os 1	0	0
86	s8	1	Total 7	N 6	Os 1	0	0
86	s9	1	Total 7	N 6	Os 1	0	0
86	c3	1	Total 7	N 6	Os 1	0	0
86	c5	1	Total 7	N 6	Os 1	0	0
86	c8	1	Total 7	N 6	Os 1	0	0
86	d4	1	Total 7	N 6	Os 1	0	0
86	d9	1	Total 7	N 6	Os 1	0	0
86	sR	1	Total 7	N 6	Os 1	0	0

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	5	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	5	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
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86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
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			7	6	1		
86	5	1	Total	N	Os	0	0
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			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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86	5	1	Total	N	Os	0	0
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			7	6	1		
86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
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			7	6	1		
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			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
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			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
			7	6	1		
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			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
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			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
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86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	5	1	Total	N	Os	0	0
			7	6	1		
86	7	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	7	1	Total	N	Os	0	0
			7	6	1		
86	7	1	Total	N	Os	0	0
			7	6	1		
86	7	1	Total	N	Os	0	0
			7	6	1		
86	7	1	Total	N	Os	0	0
			7	6	1		
86	7	1	Total	N	Os	0	0
			7	6	1		
86	7	1	Total	N	Os	0	0
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86	7	1	Total	N	Os	0	0
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86	7	1	Total	N	Os	0	0
			7	6	1		
86	7	1	Total	N	Os	0	0
			7	6	1		
86	8	1	Total	N	Os	0	0
			7	6	1		
86	8	1	Total	N	Os	0	0
			7	6	1		
86	8	1	Total	N	Os	0	0
			7	6	1		
86	8	1	Total	N	Os	0	0
			7	6	1		
86	8	1	Total	N	Os	0	0
			7	6	1		
86	8	1	Total	N	Os	0	0
			7	6	1		
86	8	1	Total	N	Os	0	0
			7	6	1		
86	8	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	8	1	Total	N	Os	0	0
			7	6	1		
86	8	1	Total	N	Os	0	0
			7	6	1		
86	8	1	Total	N	Os	0	0
			7	6	1		
86	8	1	Total	N	Os	0	0
			7	6	1		
86	8	1	Total	N	Os	0	0
			7	6	1		
86	l3	1	Total	N	Os	0	0
			7	6	1		
86	l3	1	Total	N	Os	0	0
			7	6	1		
86	l4	1	Total	N	Os	0	0
			7	6	1		
86	l4	1	Total	N	Os	0	0
			7	6	1		
86	l5	1	Total	N	Os	0	0
			7	6	1		
86	l5	1	Total	N	Os	0	0
			7	6	1		
86	l5	1	Total	N	Os	0	0
			7	6	1		
86	l9	1	Total	N	Os	0	0
			7	6	1		
86	m0	1	Total	N	Os	0	0
			7	6	1		
86	m0	1	Total	N	Os	0	0
			7	6	1		
86	m1	1	Total	N	Os	0	0
			7	6	1		
86	m4	1	Total	N	Os	0	0
			7	6	1		
86	m5	1	Total	N	Os	0	0
			7	6	1		
86	m6	1	Total	N	Os	0	0
			7	6	1		
86	m7	1	Total	N	Os	0	0
			7	6	1		
86	n1	1	Total	N	Os	0	0
			7	6	1		

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
86	n3	1	Total	N	Os	0	0
			7	6	1		
86	n9	1	Total	N	Os	0	0
			7	6	1		
86	o3	1	Total	N	Os	0	0
			7	6	1		
86	o7	1	Total	N	Os	0	0
			7	6	1		
86	q2	1	Total	N	Os	0	0
			7	6	1		

- Molecule 87 is ZINC ION (three-letter code: ZN) (formula: Zn).

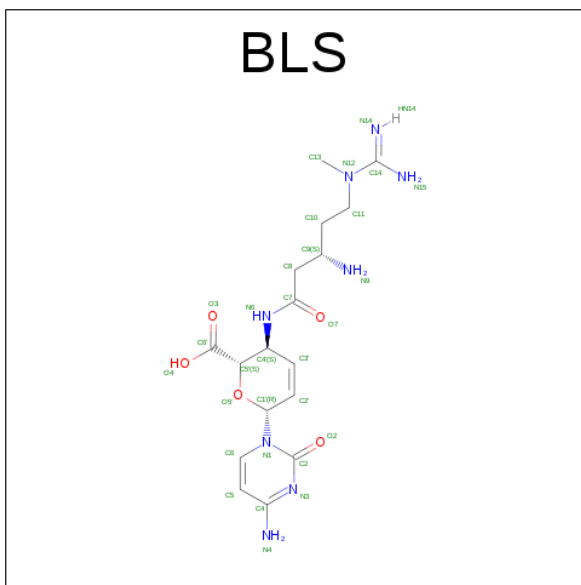
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
87	q0	1	Total	Zn	0	0
			1	1		
87	D6	1	Total	Zn	0	0
			1	1		
87	Q2	1	Total	Zn	0	0
			1	1		
87	e1	1	Total	Zn	0	0
			1	1		
87	Q3	1	Total	Zn	0	0
			1	1		
87	D9	1	Total	Zn	0	0
			1	1		
87	E1	1	Total	Zn	0	0
			1	1		
87	Q0	1	Total	Zn	0	0
			1	1		
87	d7	1	Total	Zn	0	0
			1	1		
87	q3	1	Total	Zn	0	0
			1	1		
87	d9	1	Total	Zn	0	0
			1	1		
87	D7	1	Total	Zn	0	0
			1	1		
87	d6	1	Total	Zn	0	0
			1	1		
87	o7	1	Total	Zn	0	0
			1	1		

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Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
87	O7	1	Total Zn 1 1	0	0
87	q2	1	Total Zn 1 1	0	0

- Molecule 88 is BLASTICIDIN S (three-letter code: BLS) (formula: $\text{C}_{17}\text{H}_{26}\text{N}_8\text{O}_5$).



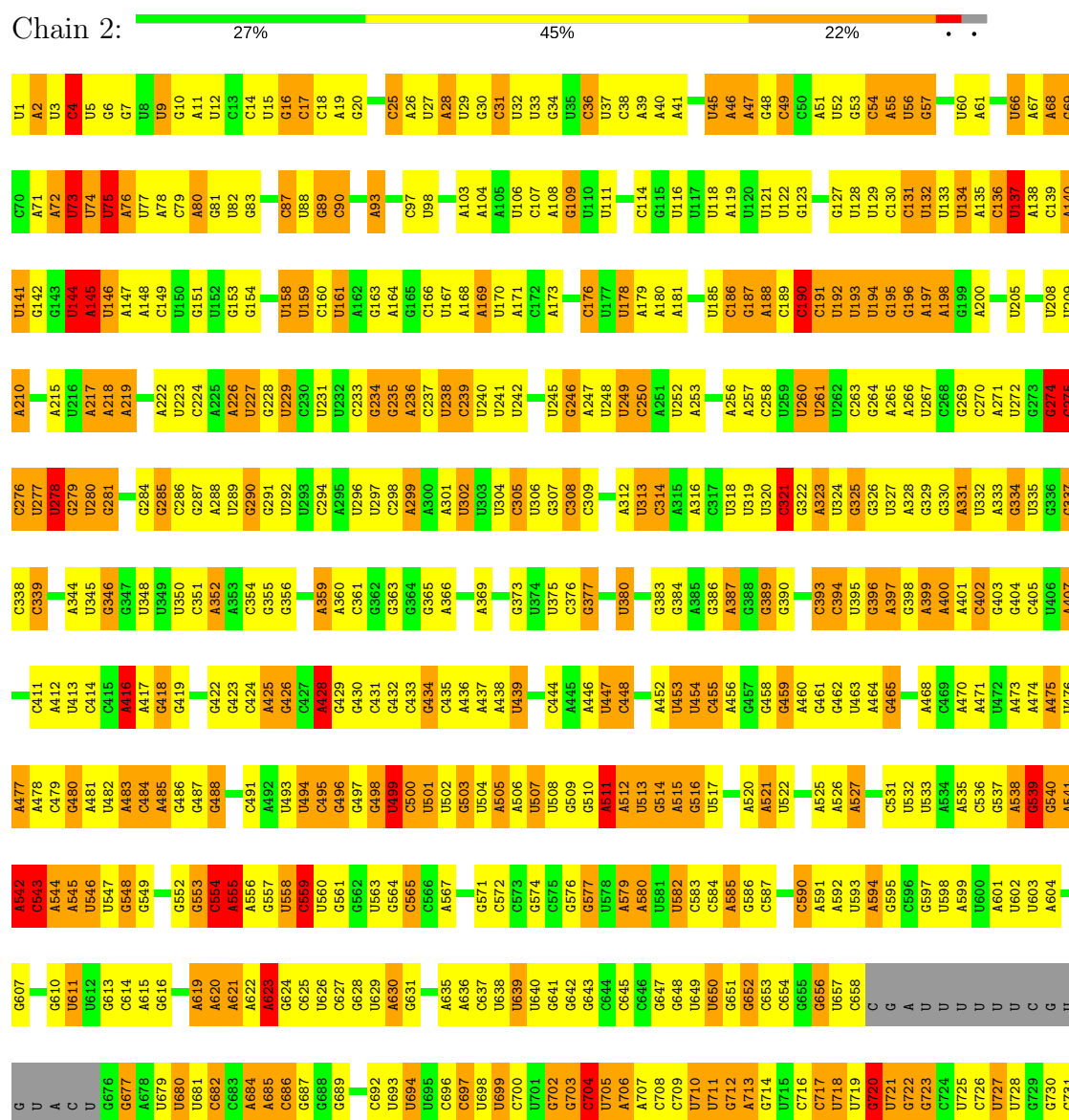
Mol	Chain	Residues	Atoms				ZeroOcc	AltConf
88	1	1	Total 30	C 17	N 8	O 5	0	0
88	5	1	Total 30	C 17	N 8	O 5	0	0

3 Residue-property plots

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

Note EDS failed to run properly.

• Molecule 1: 18S ribosomal RNA





C1796	
A1797	
U1798	
U	
A	

• Molecule 1: 18S ribosomal RNA

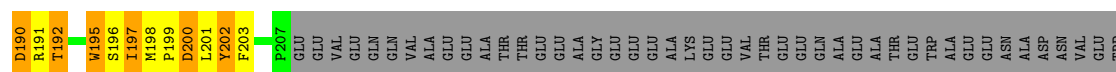
Chain 6: 

U1	G63	U132	A197	U272	C339	C409	A477	A542	G607	C674	U744	U813	U882	U945	U1
U3	U64	U	A198	G273	U340	C410	C478	C543	U608	U675	U745	A814	C883	C950	U3
U5	A65	A	G199	G274	A41	C411	C479	A544	U609	G676	U746	G815	C884	C951	U5
U6	U66		A200	C275	C342			A545	G610	G677		G816	G985	A952	U6
G6	A67	C136	U207	C276	C343	A416	U482	U546	U612	U678	A752	C817	U886	A953	U6
G7	A68	U137	U208	U277	A344	A417	A483	U547	U613	U680	A753	C818	A887	G953	G7
U8	G69	A138	U209	G278	U345	A418	C484	G548	G614	U681	A754	G819	U888	U954	U8
U9		C139	U210	G279	G346	G418	A485	G549	C614	U682	A755	U820	U889	A955	U9
A11	U72	A140	U212	U280	G347	G419	U486	A550	A615	C682	U756	U821	C990	C956	A11
G10	U73	U141	G281	G281	U348	A420	G487	G551		C683	A757	U822	A991	G957	G10
A11	U74	G142	G282	U283	U351	A421	G488	G552	G619	A684	U758	G823	A892	U958	A11
U12	U75	G143	G214	U283	C351	G422	C489	G553	A620	A685	U759	G824	U953	U959	U12
C13	A76	U144	G215	G284	A352	G423	C490	C554	A621	C686	U760	U825	U954	U960	C13
U15	U77	U145	U216	G285		C424	C491	A555	A622	G687	G761	U826	G986	U961	U15
G16	A78	U146	A217	C286	G356	A425	A492	A556	A623	G688	A762	C827	U986	C962	G16
C17	C79		A218	G287	G357	G426	U493	G557	G624	G689	G763	U828	C997	A963	C17
C18		C149	A219	U287	U358	U427	U494	U558	G625	G690	U764	U829	A898	U964	C18
A19	U82	U150	A220	G291	A359	A428	C495	C559	U626	C691	G765	U830	C999	U965	A19
G20	G83	G151	A221	U292	A360	G429	C496	C560	C627		U766	U831	A900	A966	G20
G23	A84	U152	A222	U293	C361	G430	C497	U563	U628	U694	U767	U832	A901	A967	G23
U24	A85	G153	U223	C294	G362	C431	C498	G564	U629	U695	G768	U833	G902	U968	U24
C25	A86	G154		A295	G363	G432	U499	C565	A630	C696	A769	C834	U903	C969	C25
U26	C87	U155	A226	U296	G364	C433	C500	C566	U633	U696	A770	U835	A904	U970	U26
A26	U88	A156	U227	U297		G434	U501	A567	U634	U698	A771	U836	A905	A971	A26
U27	G89	U157	G228	C298	U368	C435	U502	G568	U634	U699	G772	U837	A906	U976	U27
A28	C90	U158	U229	A299	A369	G436	G503	C569	U635	U700	C773	U838	A907	A977	A28
U29	G91	U159	C230	A300	A370	A437	U504	A570	A636	C701	A774	U839	U907	A978	U29
G30	A92	C160	U231	A301	G371	A438	A505	G571	C637	G702	G775		C910	A979	G30
U39	A93	U161	U232	U302	G372	U439	A506		U638	G703	G776	U843	U911	A979	U39
C31	U94	A152	C233		G373		U507	G574	U639	C704	G778		U912	A980	C31
	G95	C163	G234	U306	U374	C443		C575	U640	U705	U779	G846	U913	U981	
G34	G96		G235	G307	U374	C444	U508	C576	U641	U706	U780	G847	G914	U982	G34
U35	C97	C166	A236	C308	G377		G509	C577	U642	A706	U781	C848	A915	U983	U35
C36	U167	U166	C237	C309		U447	G510	U578	U643	C709	U782	C849	U916	A984	C36
U37	A168	A168		C310	U380	C448	A511	C579	C644	U710	G783		U917	U985	U37
C38	C99	A169	U240	U311	C381	C449	U512	A580		U711		G853	U918	U986	C38
A39	A100	A169	U241	A312	C382	U450	G514	U581	G647		U788	U854	A919	U987	A39
A40	U101	U170	U242	U313	G384	A451	A515	U582	U648	G714	U789	C855	U920	C990	A40
A41	U102	A171	U242	C314	C384	A452	G516	C583	U649	U715	U790	A856	U921	U991	A41
G42	A103	C172		A315	A385	U453	U517	C584	U650	C717	U791	C857	U922	A992	G42
A43	A104	A173	U245	A316	G386	U454	A518	A585	U651	U718	U792	C858	A923	A993	A43
U44	A105		G246		A387	C455	C519	C586	G652	U719	U793	A859	C924	C994	U44
U45	U108	C176	A247	U320		A456	A520	C587	C653	U720	U794	U860	G925	A995	U45
A46	G109	U178	U248	C321	G390	G457	A521	U588	U654	U721	U795	U861	A926	U996	A46
A47	U110	A179	C250	A322	A391	U522	U522	U588	G655	U722	U796	A862	C927	C997	A47
G48	U111	A180	A251	A323	G392	G523	G523	A591		G723	G797	A863	C931	A998	G48
C49	U112	A181	U252	U324	C393	U524	U524	A592	C658	U724	U798		C932	U999	C49
C50	U113			G325		A525	A525	U593	U725	U725	U799		U932	C1000	C50
A51	C114	U185	A256	G326	A397	U463	A526	U593	U726	U726	U800	G866	U933	A1001	A51
U52	G115	C186	U257	U327	A398	A464	A527	G595	A661	U727	U801	C867	U934	U1001	U52
G53	U116	C187	A328	A328	A399	U466	A527	C596	U662	U729	G801	C868	U935	A1002	G53
U54	U117	A188	U260	G329	A400	G467	C530	U597	U665	G730	A804	C870	G936	U1003	U54
A55	U118	C189	U261	G330	A401		C531	U598	U666	G731	U805	C871	G937	U1004	A55
U56	A119	C190		A331	C402	A470	C531	A599	U667	U732	U806	C872	C938	A1005	U56
G57	U120	C191	A265	U332	G403	A471	C536	U600	C668	U733	U807	U873	A939	U999	G57
U58	G123	U192	A266	A333	G404	U472	G537	U600	C669	U734	U808	C874	A940	U1009	U58
C59	U123	A180	U267	G334	G404	A473	A538	U603	U670	U735	U809	C875	C1010	C1009	C59
U60	U194	C199	C268		U406	A474	G539	U604	G	C741	G810	C876	G942	U1011	U60
A61	A126	G195		C337	A408	U475	G540	A605	U	U742	A811	C877	C943	U1012	A61
C62		U196	A271	C338	U409	U476	A606	A606	A672	U743	A812		C944	U1013	C62



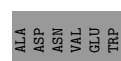
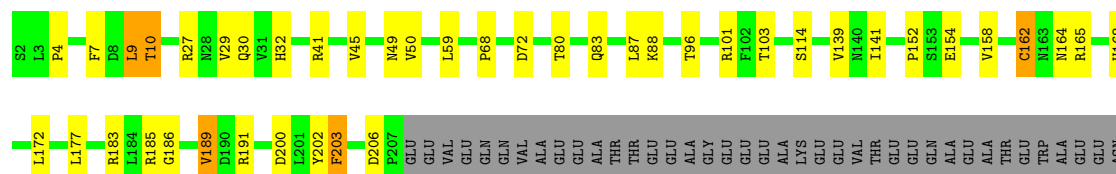
Frequency	Percentage
Daily	17%
Often	45%
Sometimes	18%
Rarely	2%
Never	18%





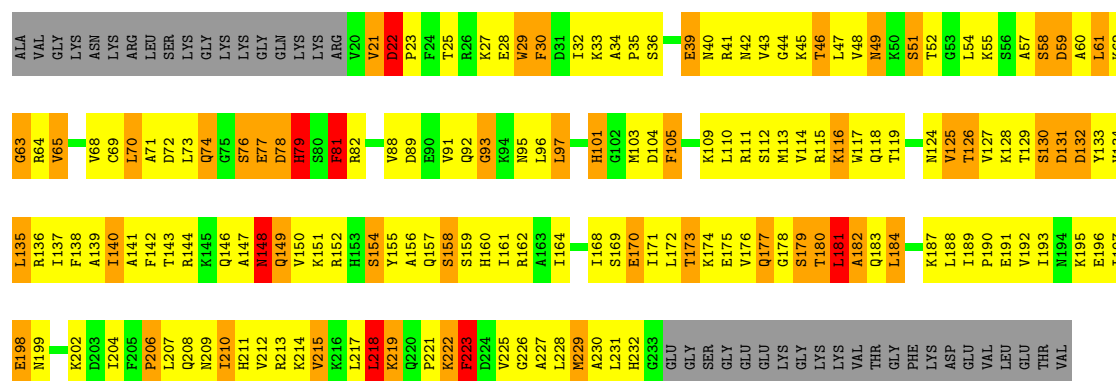
• Molecule 2: 40S ribosomal protein S0-A

Chain s0: 65% 15% 18%



• Molecule 3: 40S ribosomal protein S1-A

Chain S1: 19% 44% 18% 16%



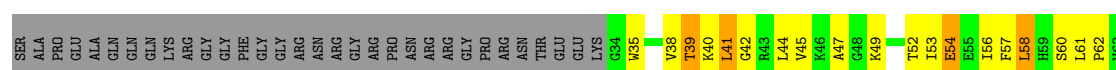
• Molecule 3: 40S ribosomal protein S1-A

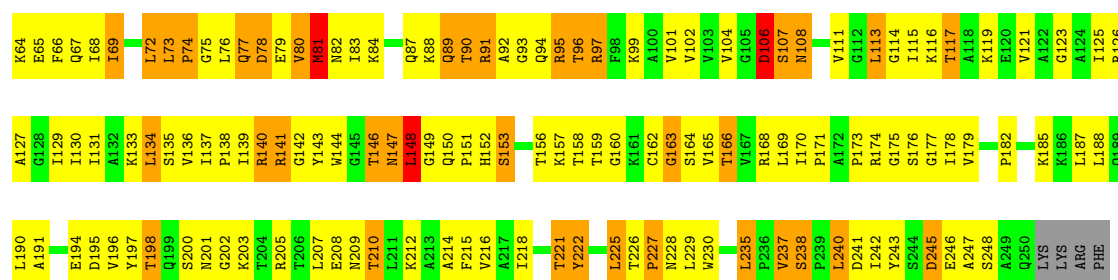
Chain s1: 64% 20% 15%



• Molecule 4: 40S ribosomal protein S2

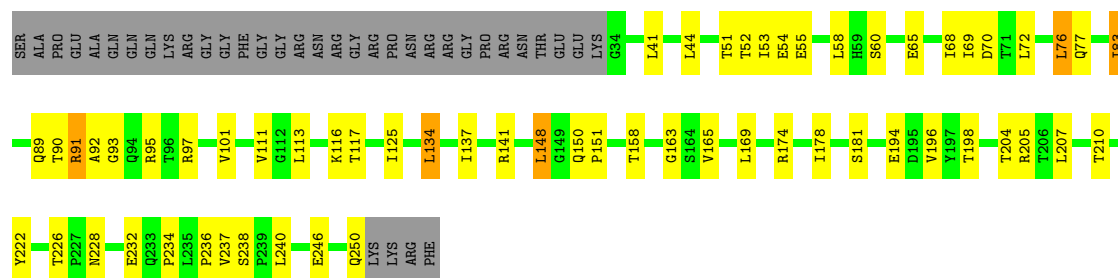
Chain S2: 24% 45% 16% 14%





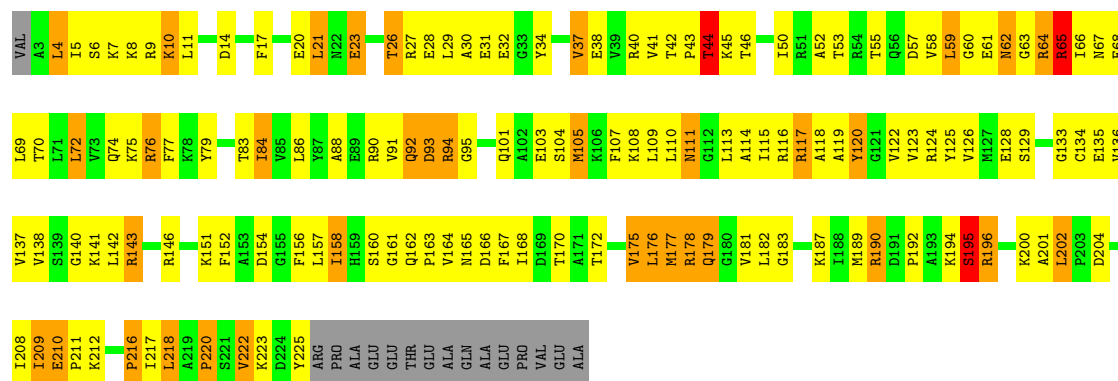
• Molecule 4: 40S ribosomal protein S2

Chain s2: 62% 22% 14%



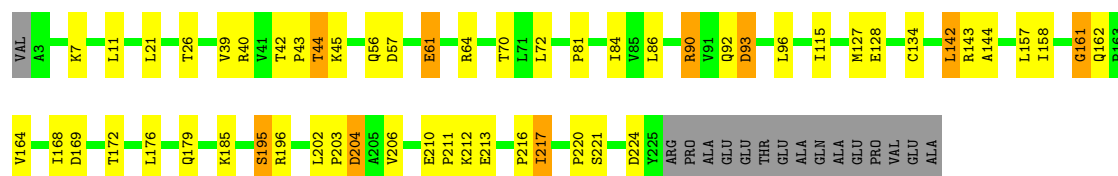
• Molecule 5: 40S ribosomal protein S3

Chain S3: 32% 46% 15% 7%



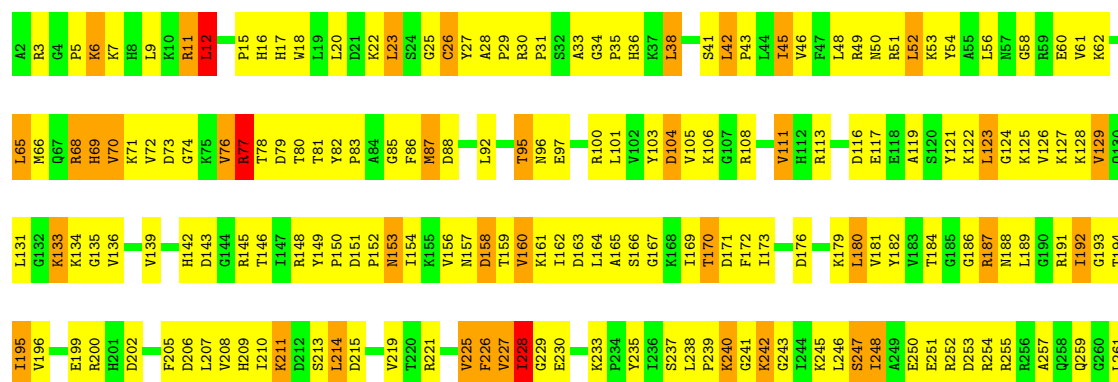
• Molecule 5: 40S ribosomal protein S3

Chain s3: 70% 20% 7%



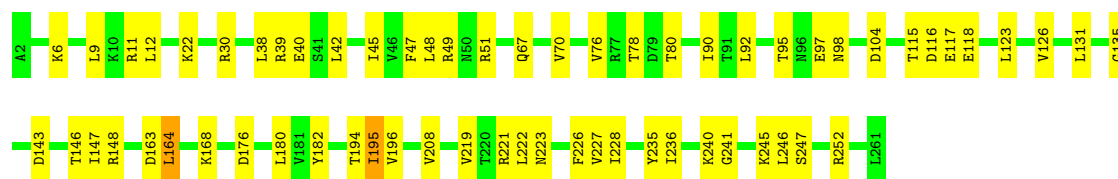
• Molecule 6: 40S ribosomal protein S4-A

Chain S4: 30% 55% 14%



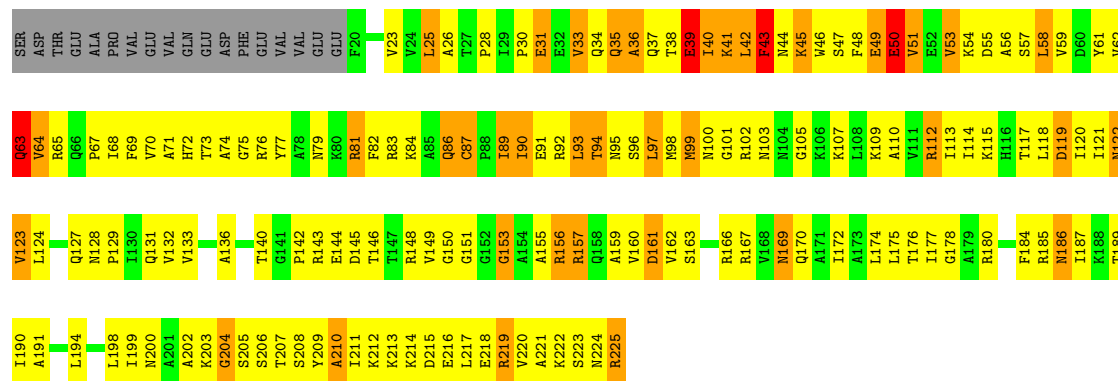
• Molecule 6: 40S ribosomal protein S4-A

Chain s4: 76% 23%



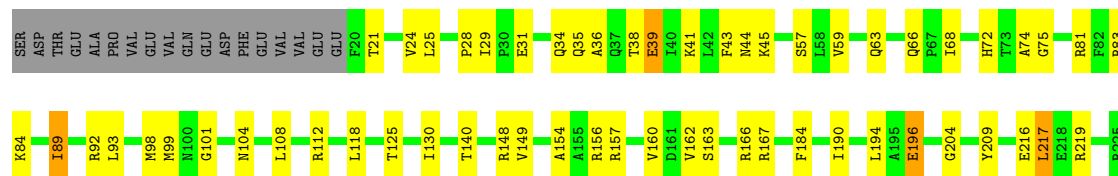
• Molecule 7: 40S ribosomal protein S5

Chain S5: 21% 52% 17% 8%



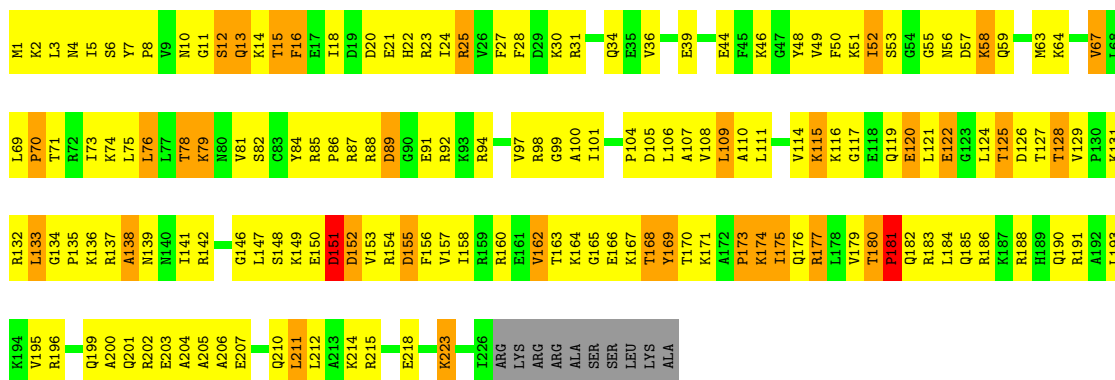
• Molecule 7: 40S ribosomal protein S5

Chain s5: 66% 24% 8%



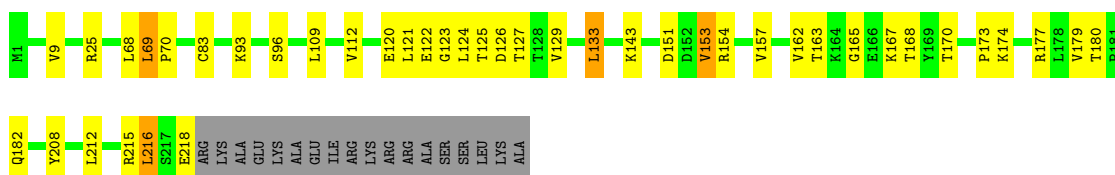
• Molecule 8: 40S ribosomal protein S6-A

Chain S6: 27% 54% 14%



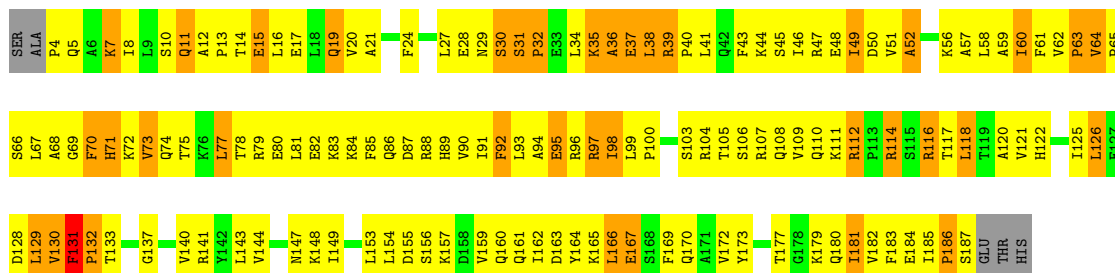
• Molecule 8: 40S ribosomal protein S6-A

Chain s6: 75% 16% 8%



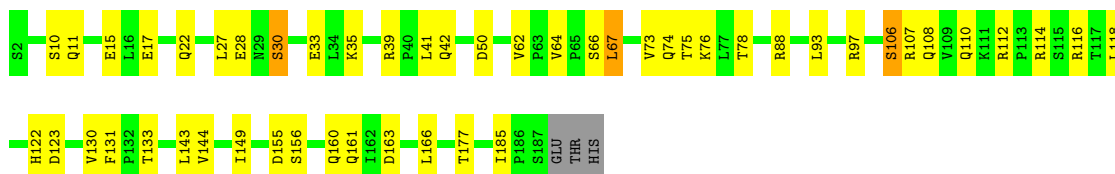
• Molecule 9: 40S ribosomal protein S7-A

Chain S7: 21% 57% 20%



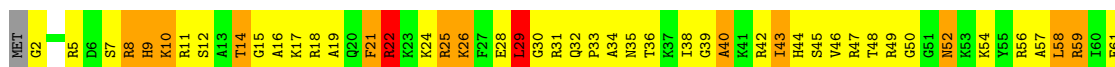
• Molecule 9: 40S ribosomal protein S7-A

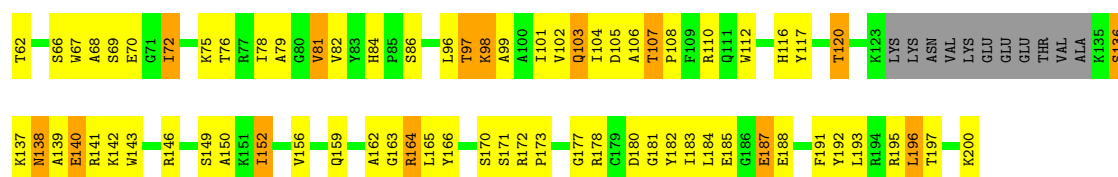
Chain s7: 72% 25%



• Molecule 10: 40S ribosomal protein S8-A

Chain S8: 35% 46% 13% 6%





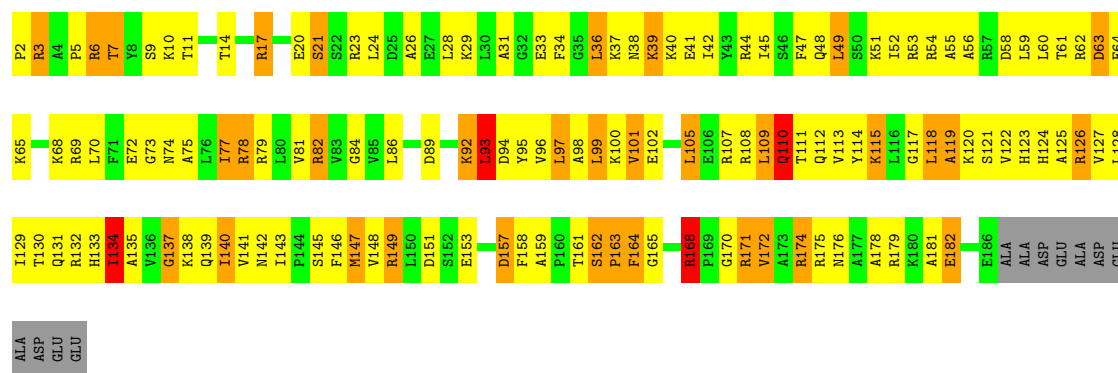
• Molecule 10: 40S ribosomal protein S8-A

Chain s8: 80% 14% 6%



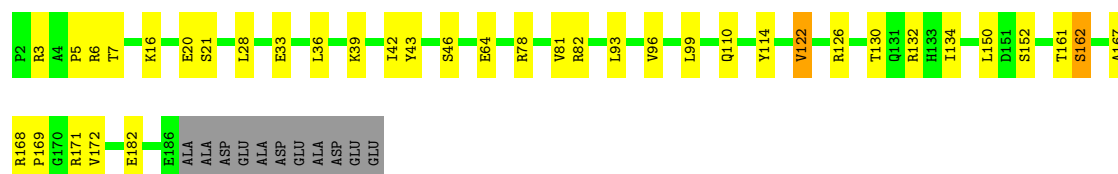
• Molecule 11: 40S ribosomal protein S9-A

Chain S9: 26% 49% 17% 6%



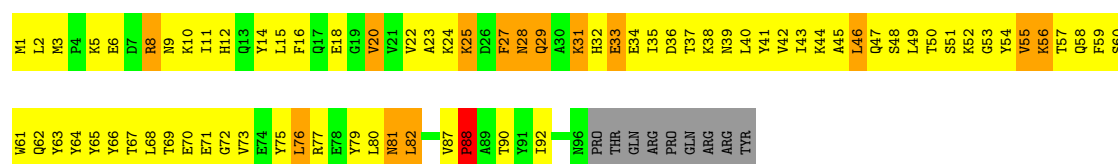
• Molecule 11: 40S ribosomal protein S9-A

Chain s9: 75% 18% 6%



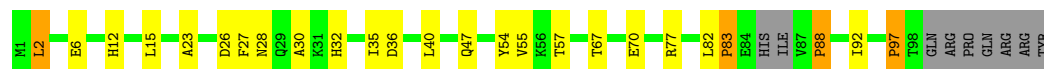
• Molecule 12: 40S ribosomal protein S10-A

Chain C0: 19% 58% 13% 9%



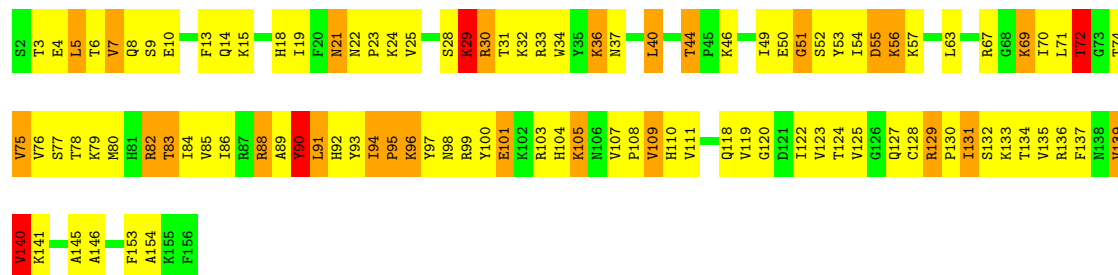
- Molecule 12: 40S ribosomal protein S10-A

Chain c0:  68% 20% 9%



- Molecule 13: 40S ribosomal protein S11-A

Chain C1: 33% 48% 16%

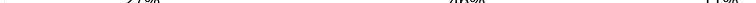


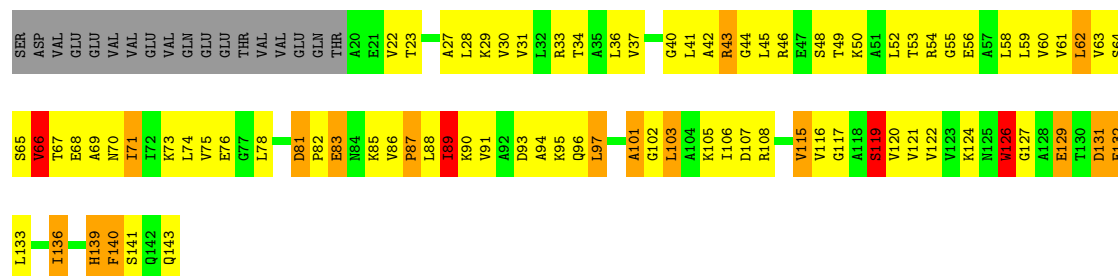
- Molecule 13: 40S ribosomal protein S11-A

Chain c1: 76% 18% 6%



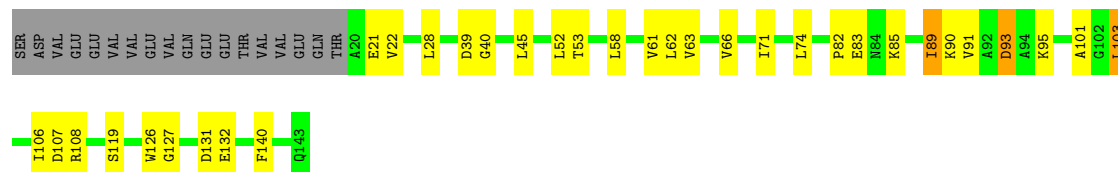
- Molecule 14: 40S ribosomal protein S12

Chain C2: 

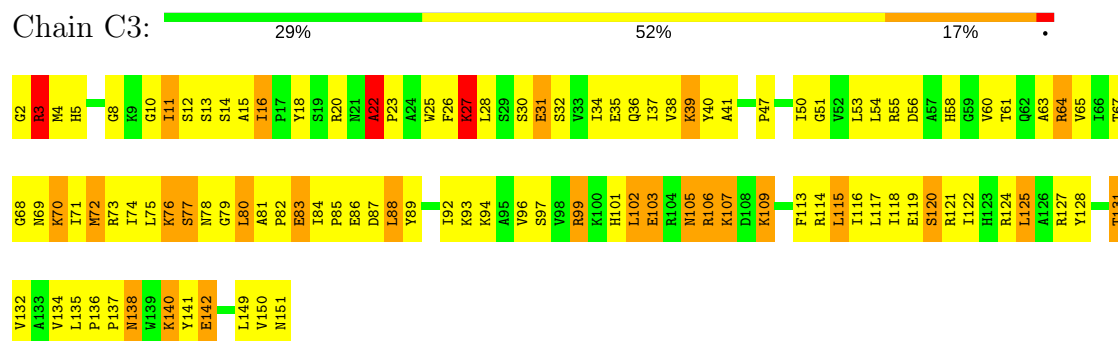


- Molecule 14: 40S ribosomal protein S12

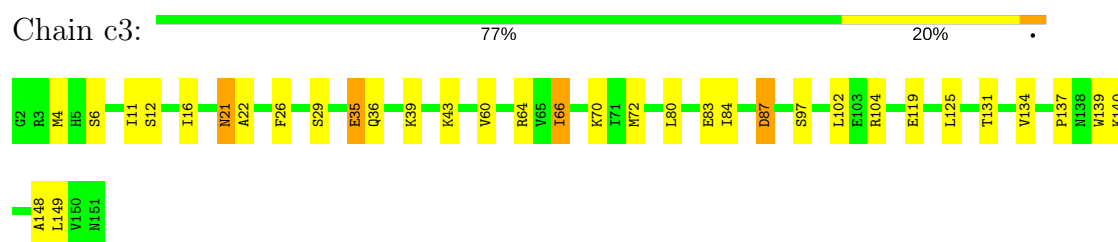
Chain c2:  63% 22% • 13%



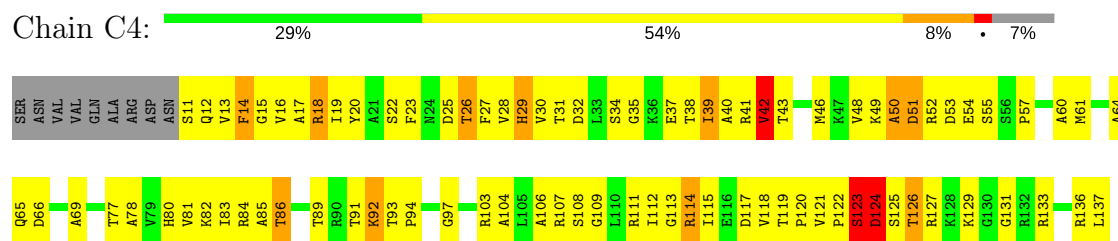
- Molecule 15: 40S ribosomal protein S13



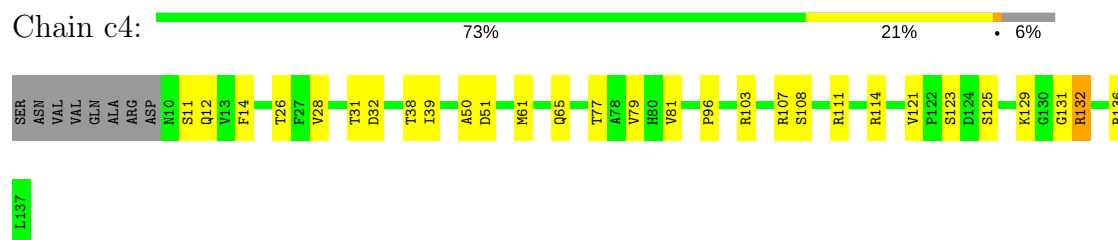
- Molecule 15: 40S ribosomal protein S13



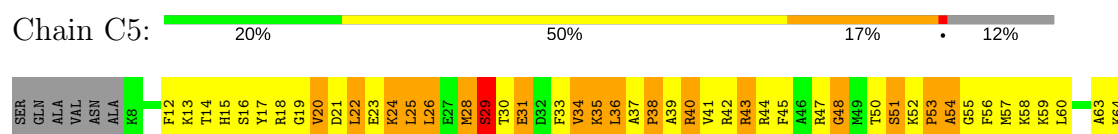
- Molecule 16: 40S ribosomal protein S14-A

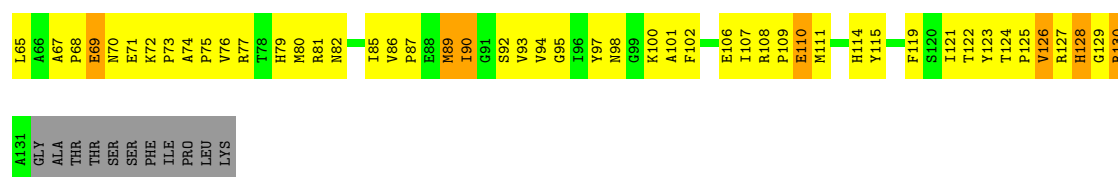


- Molecule 16: 40S ribosomal protein S14-A

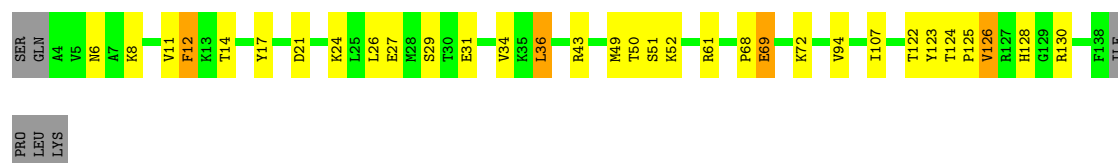


- Molecule 17: 40S ribosomal protein S15

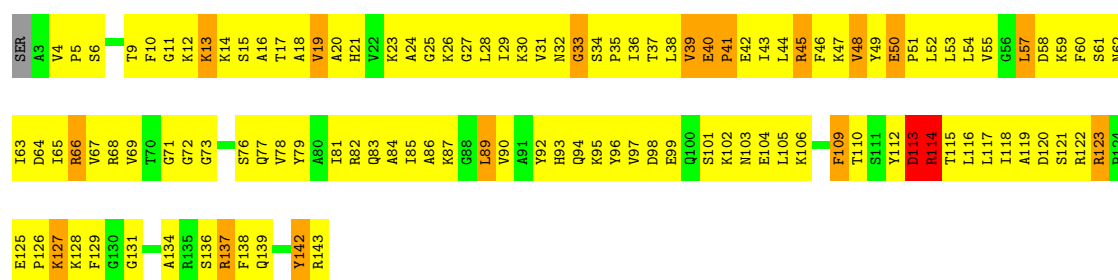




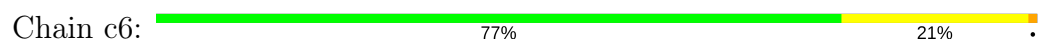
- Molecule 17: 40S ribosomal protein S15



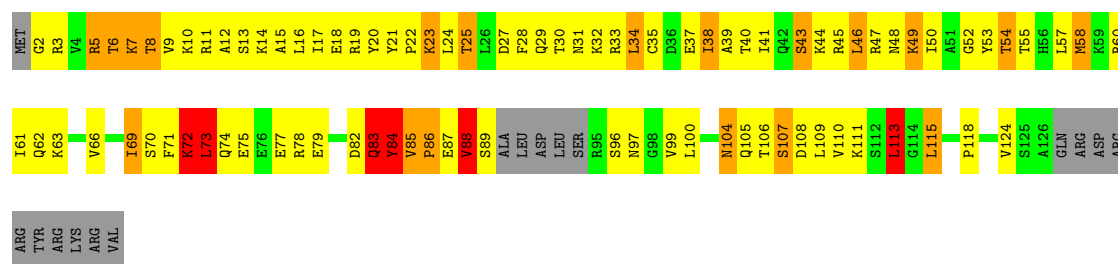
- Molecule 18: 40S ribosomal protein S16-A



- Molecule 18: 40S ribosomal protein S16-A

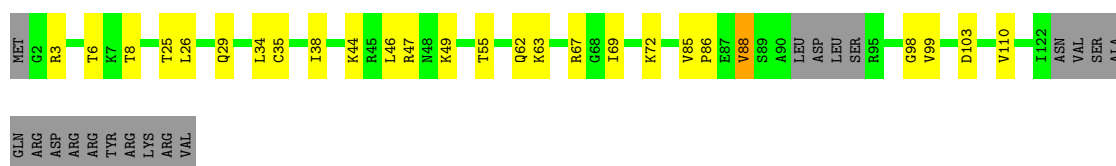


- Molecule 19: 40S ribosomal protein S17-A



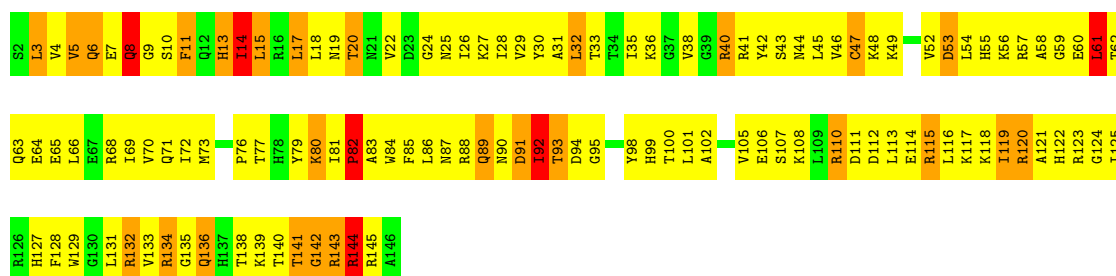
- Molecule 19: 40S ribosomal protein S17-A



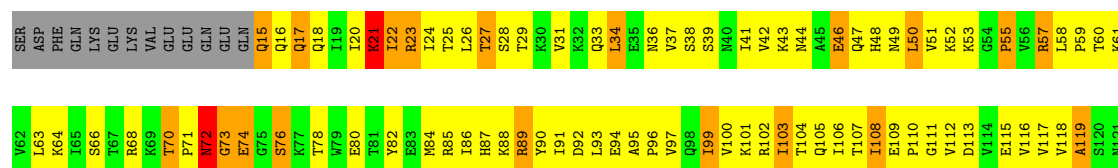


- Molecule 20: 40S ribosomal protein S18-A

Chain C8: 16% 62% 18%

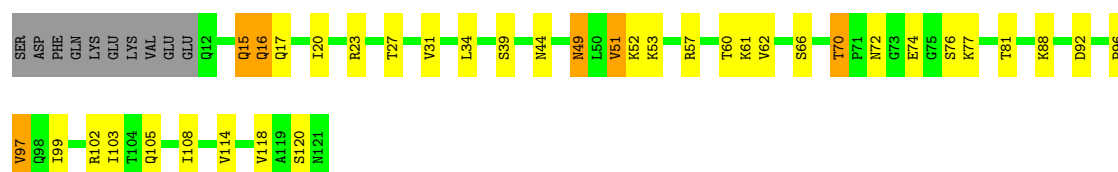


Chain D0: 

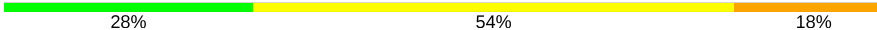


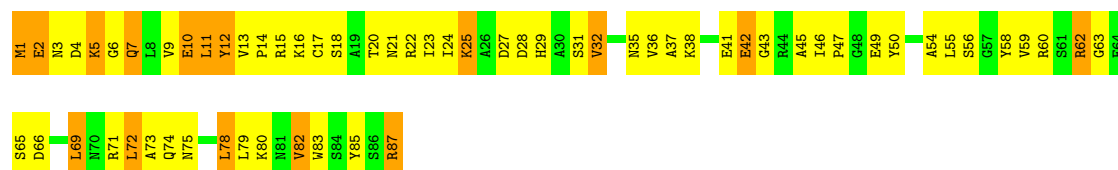
• Molecule 22: 40S ribosomal protein S20

Chain d0: 




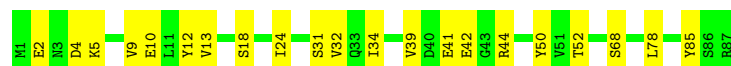
• Molecule 23: 40S ribosomal protein S21-A

Chain D1: 




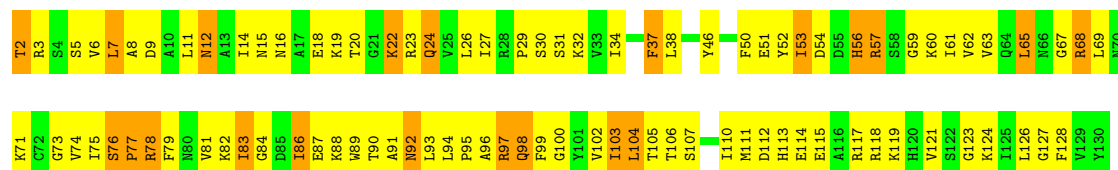
• Molecule 23: 40S ribosomal protein S21-A

Chain d1: 




• Molecule 24: 40S ribosomal protein S22-A

Chain D2: 

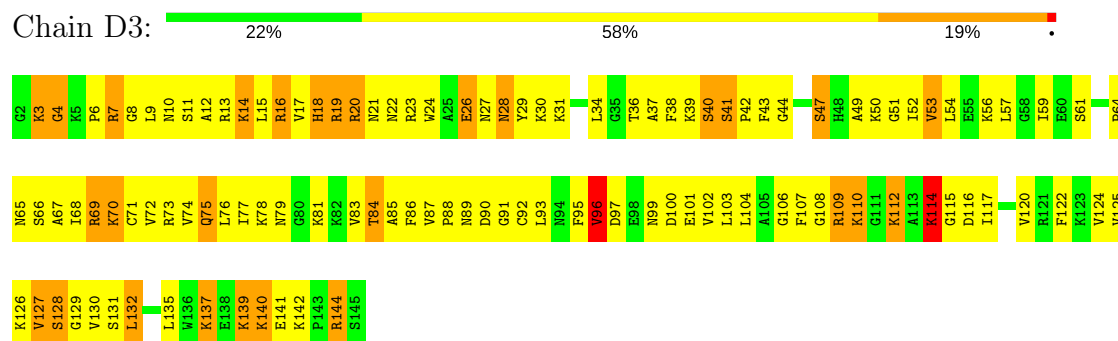


• Molecule 24: 40S ribosomal protein S22-A

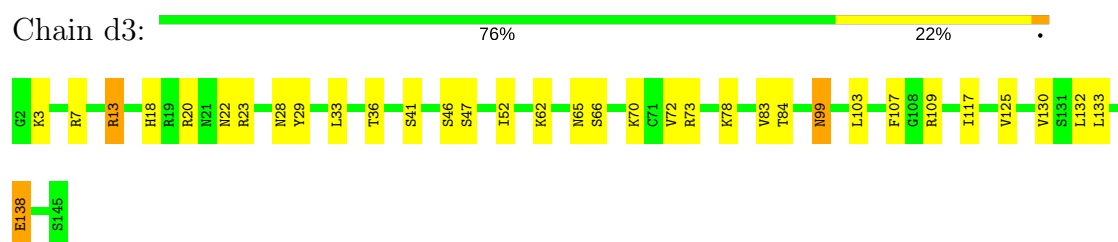
Chain d2: 



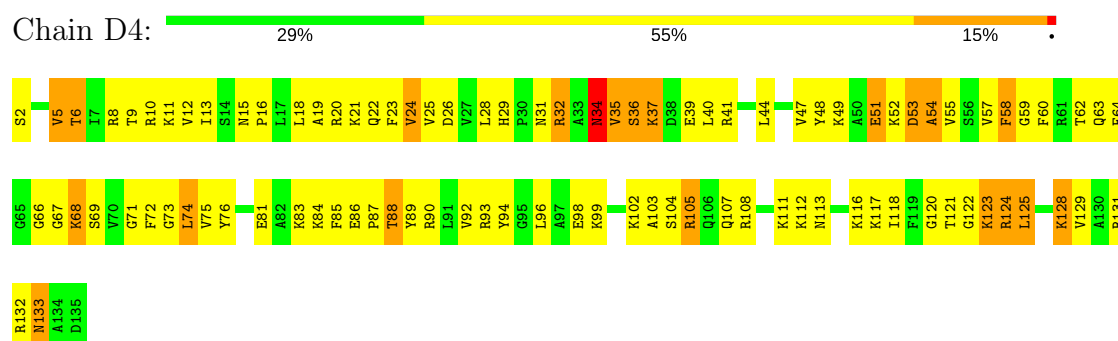
- Molecule 25: 40S ribosomal protein S23-A



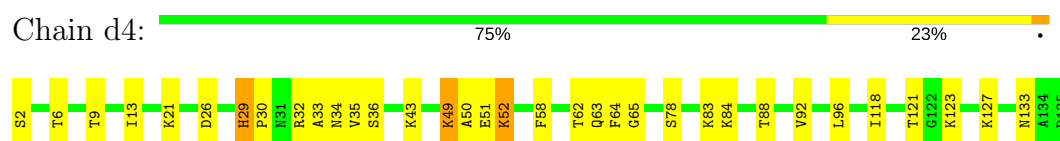
- Molecule 25: 40S ribosomal protein S23-A



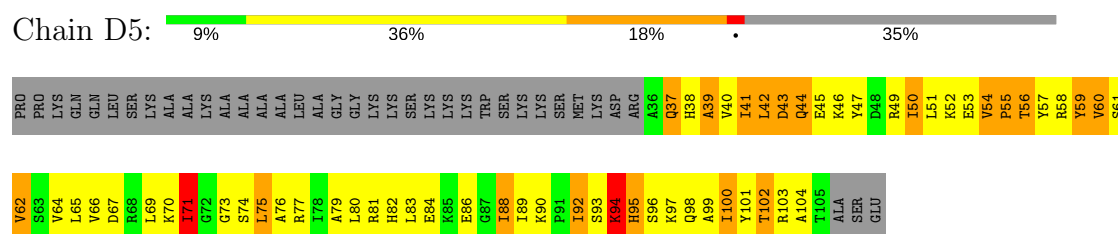
- Molecule 26: 40S ribosomal protein S24-A



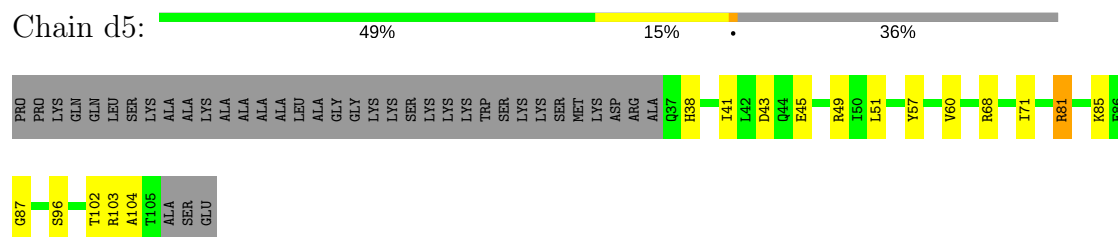
- Molecule 26: 40S ribosomal protein S24-A



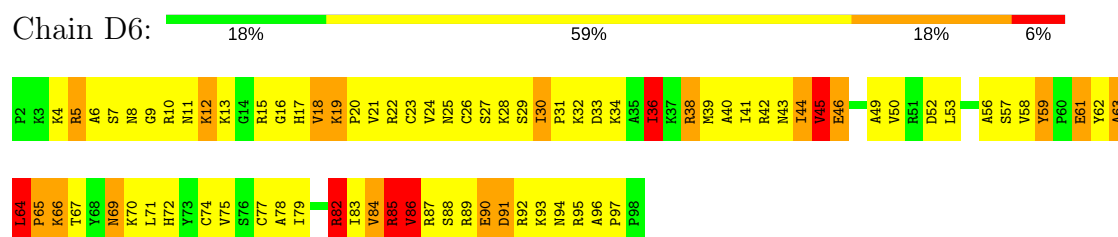
- Molecule 27: 40S ribosomal protein S25-A



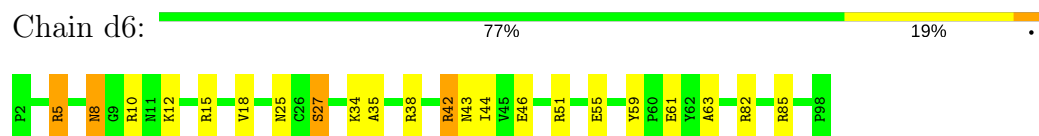
- Molecule 27: 40S ribosomal protein S25-A



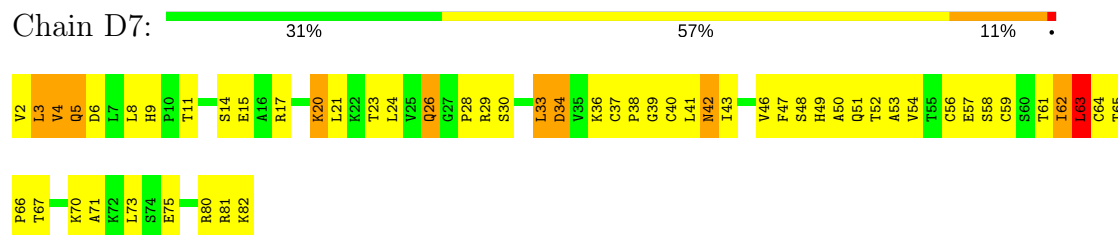
- Molecule 28: 40S ribosomal protein S26-B



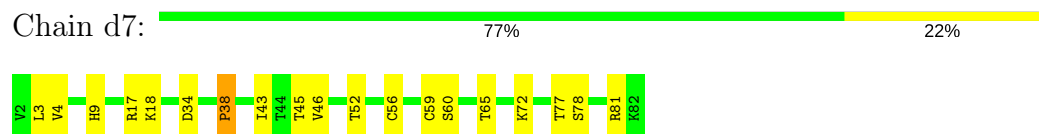
- Molecule 28: 40S ribosomal protein S26-B



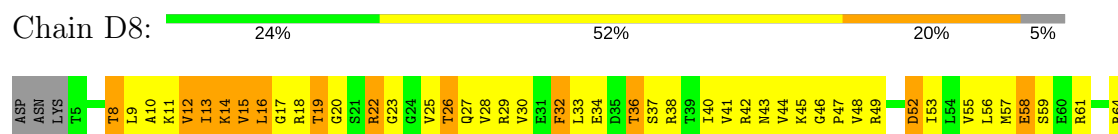
- Molecule 29: 40S ribosomal protein S27-A



- Molecule 29: 40S ribosomal protein S27-A



- Molecule 30: 40S ribosomal protein S28-A





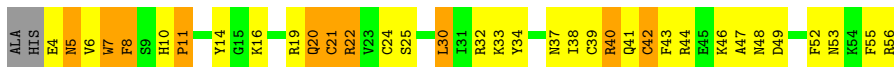
- Molecule 30: 40S ribosomal protein S28-A

Chain d8: 74% 17% 5% 5%



- Molecule 31: 40S ribosomal protein S29-A

Chain D9: 33% 45% 18% .



- Molecule 31: 40S ribosomal protein S29-A

Chain d9: 71% 24% . .



- Molecule 32: 40S ribosomal protein S30-A

Chain E0: 42% 45% 12% .



- Molecule 33: Ubiquitin-40S ribosomal protein S31

Chain E1: 18% 50% 20% 5% 7%



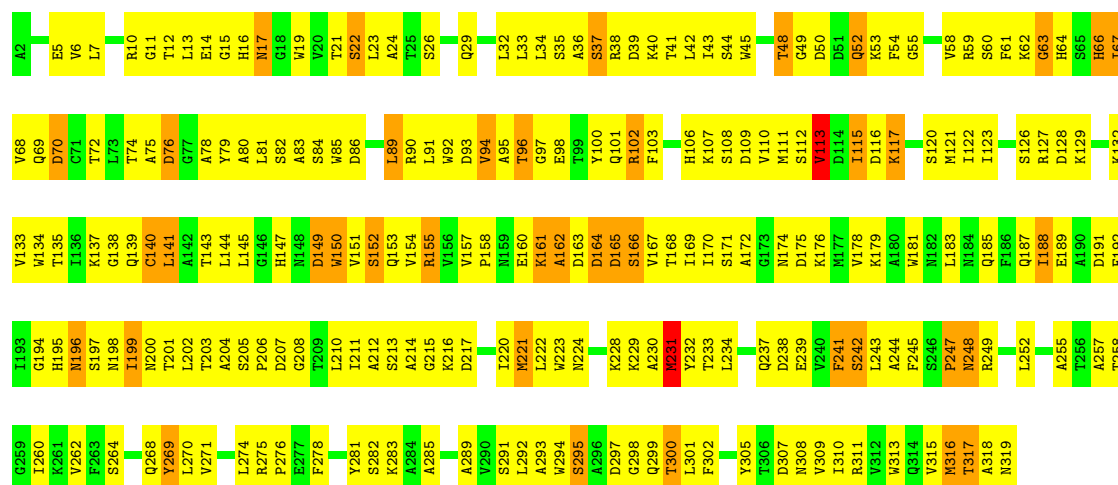
- Molecule 33: Ubiquitin-40S ribosomal protein S31

Chain e1: 57% 38% . .



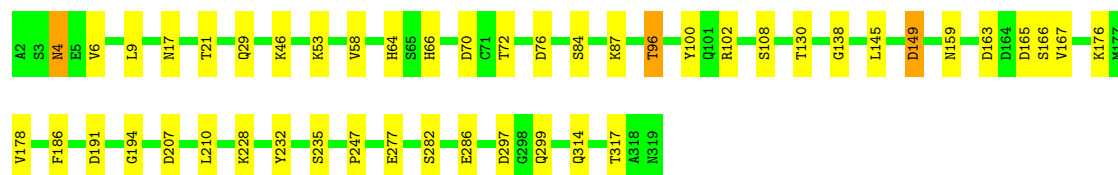
- Molecule 34: Guanine nucleotide-binding protein subunit beta-like protein

Chain SR: 26% 60% 13% .



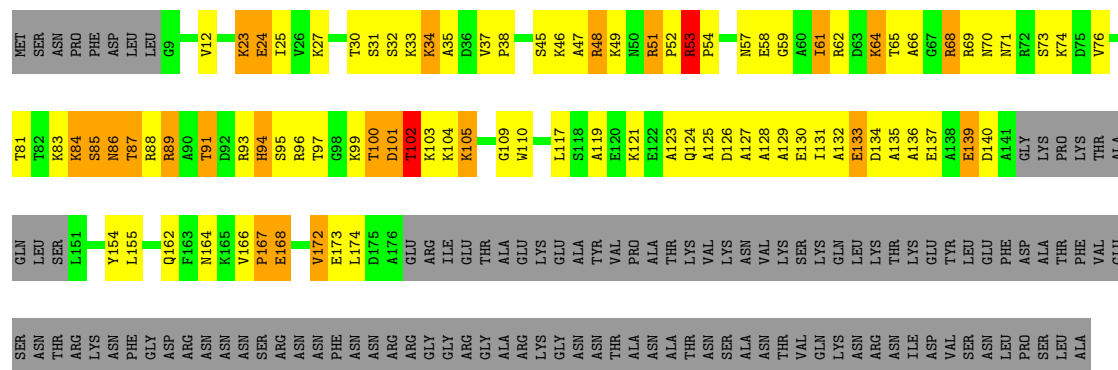
- Molecule 34: Guanine nucleotide-binding protein subunit beta-like protein

Chain sR: 85% 14% •



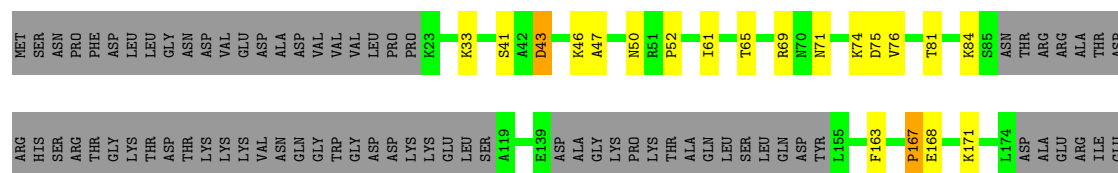
- Molecule 35: Suppressor protein STM1

Chain SM: 25% 24% 8% 42% •



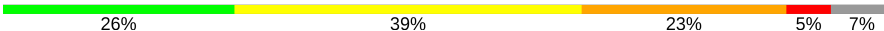
- Molecule 35: Suppressor protein STM1

Chain sM: 31% 7% 62% •



THR	GLY	ALA	GLU	LYS	GLU	ALA	TYR	VAL	PRO	THR	LYS	VAL	LYS	ASN	VAL	LYS	SER	LYS	GLN	LEU	LEU	PHE	ASP	ALA	THR	THR	PHE	VAL	GLU	SER	ASN	THR	LYS	ARG	LYS	ASN	PHE	GLY	ASP	ARG	ASN	ASN	ASN	ASN	SER	ARG	ASN	ASN	PHE	ASN	ARG	GLY	ARG	GLY	ARG
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• Molecule 36: 25S ribosomal RNA

Chain 1: 

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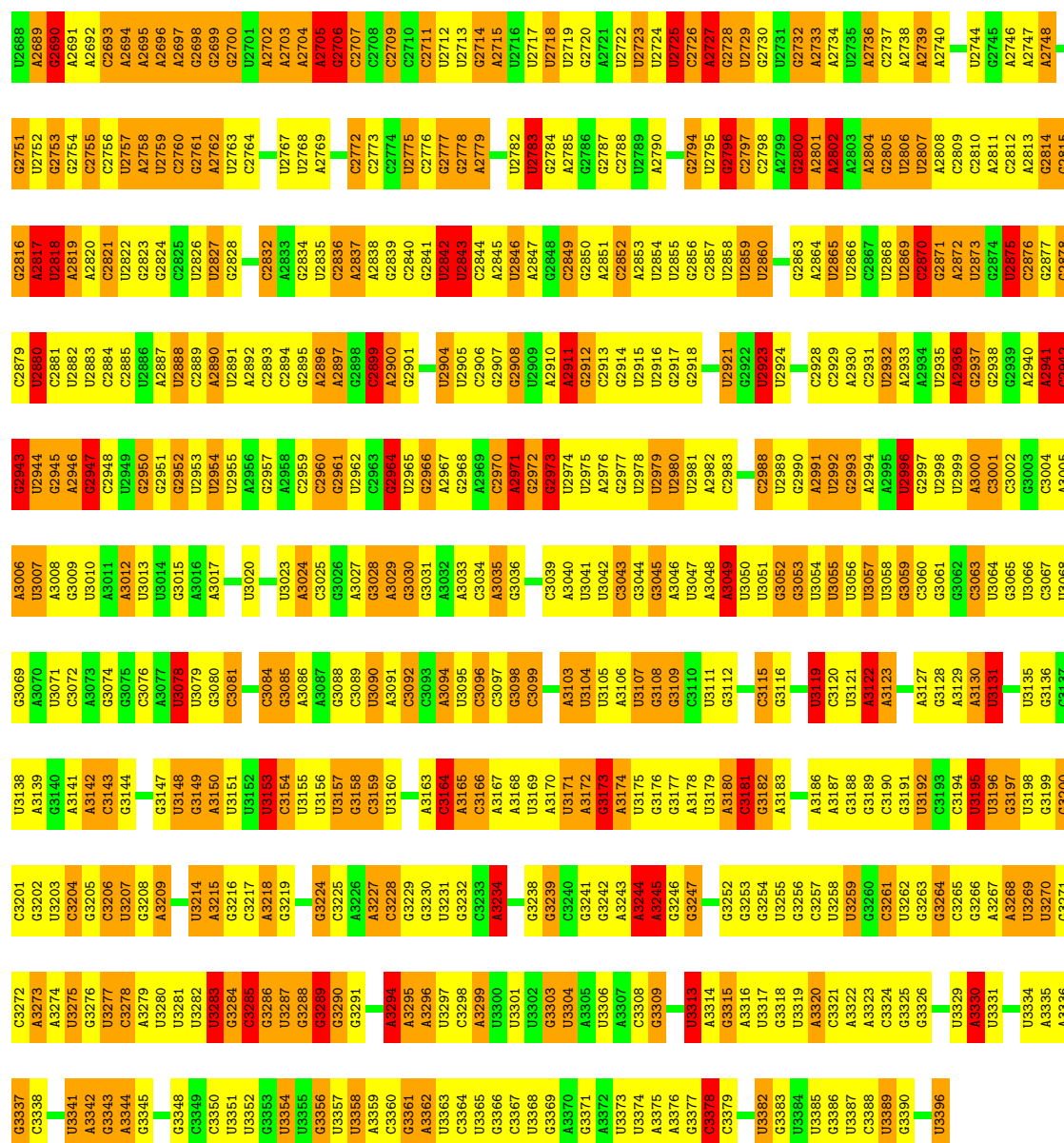
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A1945	A1809	U1601	A1601	U1534	C1469	C1404	A1343	A1273	U1208	G1140	U1071	G1005	G941
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U1955	A1814	U1606	U1606	G1538	A1474		U1348	U1278	A1212	U1145	U1077	G1011	U946
G1954	U1815	U1607	U1607	G1541	A1475	C1411	G1349	C1279	A1217	G1147	U1077	G1012	U947
U1956	A1816	A1608	A1608	G1542	U1476	C1412	U1350	C1280		U1078	U1078	U1013	C948
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C	U1620	U1620	U1620	U1558	U1488	U1425	G1362	A1303	U1235	A1165			C968
C	A1621	A1621	A1621	U1559	U1489	C1426	G1363	C1296	G1236	G1166			C969
U	G1624	G1624	G1624	G1560	A1490	G1427	A1366	A1307	G1237	U1170			G970
U	A1625	A1625	A1625	C1562	A1491	G1428	G1367	A1308	G1238	A1170			A971
C	U1626	U1626	U1626	C1563	G1492	G1429	U1368	A1309	C1239	G1171			A972
C	U1627	U1627	U1627	U1564	U1493	U1430	A1369	U1309	G1240	G1172			A973
C	G1628	G1628	G1628	G1565	U1494	A1431	A1370	U1310	U1241				G974
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C	C1701	C1701	C1701	U1567	C1496	A1433	G1372	G1313	G1243	C1176			U976
C	U1702	U1702	U1702	U1568	C1497	U1434	C1373	U1314	A1244	G1177			C977
C	C1703	C1703	C1703	U1569	U1498	A1435	A1373	U1315	A1245	A1178			G978
C	U1704	U1704	U1704	U1570	C1499	U1436	G1374	U1316	G1246	A1179			U979
C	G1705	G1705	G1705	U1571	U1500	U1437	G1375	C1317	U1247	A1180			A980
A	G1775	G1775	G1775	U1572	G1501	U1438	G1376	G1318	G1248	U1181			U981
G	U1780	U1780	U1780	U1573		U1439	G1377	U1319	G1249	U1182			C982
C	G1781	G1781	G1781	A1504	A1504	G1440	U1378	U1314	A1245	A1179			A983
C	G1782	A1714	A1714	C1505	C1505	G1441	G1379	U1315	G1246	A1179			G984
C	U1783	U1715	U1715	U1506	U1506	U1442	G1380	U1316	U1247	A1180			U985
C	G1784	U1716	U1716	C1507	C1507	G1443	A1381	A1317	U1248	U1181			U986
C	U1785	U1717	U1717	C1508	C1508	U1444	G1382	G1318	G1249	U1182			U987
C	G1786	U1718	U1718	A1509	A1509	U1445	G1383	G1319	G1250	A1183			A988
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C	C1788	U1720	U1720	C1581	C1581		U1385	G1321	A1252	U1051			G1000
C	U1789	U1721	U1721	U1511	U1511		G1386	U1322	U1253	G1186			G1001
C	G1790	U1722	U1722	U1512	U1512	A1449	A1386	U1323	U1254	C1187			A1002
C	U1791	U1723	U1723	G1513	G1513	C1450	G1387	G1324	C1255	A1053			
C	C1724	U1724	U1724	G1514	G1514	C1451	U1388	U1325	G1256	U1056			G991
C	U1725	A1655	A1655	A1515	A1515	C1452	G1389	A1326	G1257	U1123			A992
C	C1726	C1657	C1657	U1516	U1516	U1455	C1391	C1327	U1258	U1124			C993
C	G1727	U1658	U1658	G1517	G1517	A1456	G1392	C1328		G1125			G994
C	U1728	U1659	U1659	U1518	U1518		G1393	C1329	G1262	G1126			U995
C	U1729	A1589	A1589	G1519	G1519	C1459	A1394	A1330	A1263	A1197			A996
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C	A1731	C1592	C1592	G1521	G1521	A1461	G1396	C1332	U1265	A1130			A998
C	U1732	U1522	U1522	U1522	U1522	U1462	C1397	C1333	G1266	G1131			C999
C	G1733	A1524	A1524	U1523	U1523	U1463	U1398	U1334	U1267	C1201			A1065
C	U1734	U1595	U1595	A1524	A1524	U1464	U1399	C1339	U1268	A1202			G1066
C	G1735	C1596	C1596	U1596	U1596	G1465	A1401	G1340	U1269	A1203			G1001
C	A1667	U1597	U1597	C1527	C1527	G1466			U1270	A1204			A1002
C	G1668	G1598	G1598										

U2974	A2945	U2719	U2650	G2585	U2517	G2391	A2324	U2260	C2196	C2132	G
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G2977	G2848	U2722	C2653	U2588		G2394		C2263	U2199	U2135	G
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U2979	G2850	U2724	A2656	G2592	A2523	G2396	C2333	C2265	G2201	U2137	U
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A2987	G2857	U2731	G2663	G2602	G2530	G2403	C2340	G2273	A2208	A2144	U
C2988	U2858	U2732	C2664	G2603	C2531	A2404	C2341	U2274	U2209	A2145	C
U2989	U2859	A2733	U2665	G2604	U2532	C2405	U2342	A2275	G2210	A2146	U
C2990	A2860	U2734	C2666	U2605	G2533	C2406	C2343	G2276		A2147	G
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U2992	U2862	U2736	G2672	G2607	A2535	U2408	U2344	C2278	A2150		G
G2993	G2863	C2737	A2673	G2608	G2536	G2409		A2279	G2151	G2150	C
A2994	U2864	A2738	A2674	A2609	U2537	U2410	C2350	A2280	C2152		C
U2995	U2865	A2739	C2675	G2610	U2538	U2411	U2351	A2281	U2217	U2153	G
C2996	G2866	A2740	A2676	U2611	C2539	G2412	G2352	U2282	G2218	U2154	A
G2997	U2867	U2741	G2677	U2612	A2540	A2413	G2353	G2283	A2219		C
	U2868	U2742	G2678	U2613	U2541	G2414	C2354	C2284	G2220	G2155	C
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G3002	U2871	G2745	U2681	C2616	U2544	U2417	A2357	C2287	A2223	A2158	C
C3003	A2872	A2746	C2682	U2617	C2545	G2418	A2358	C2288	U2225	C2163	A
G3004	U2873	A2747	U2683	G2618	G2546	A2419	C2359	U2289	U2226	A2164	G
A3005	G2874	U2748	C2684	U2619	A2547	C2420	C2360	C2290	U2227	G2165	U
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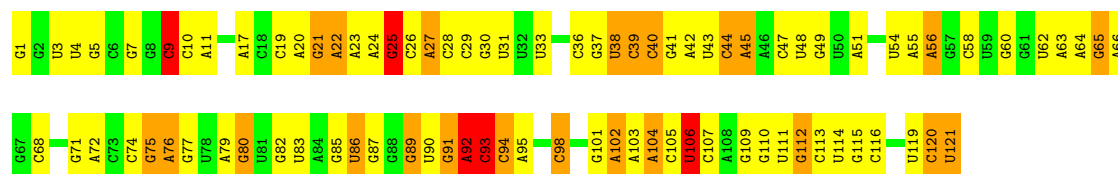
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PDB
PROTEIN DATA BANK

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		U	U	A2321		G2194	C2128	G	G	U1874	A1805	U1737	U1667
		U	A	A2322	A2259	U2195	U2129	U	U	G1875	A1806	U1738	U1668
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		U	U	G2364	G2302	G2237			C	G1915	U1848	U1780	U1712
		A	A	C2365	A2303	G2238	U2176	A2106	U	U1916	U1849	U1781	U1713
									C	C1917	U1850	U1782	U1714



• Molecule 37: 5S ribosomal RNA

Chain 3: 28% 49% 19%



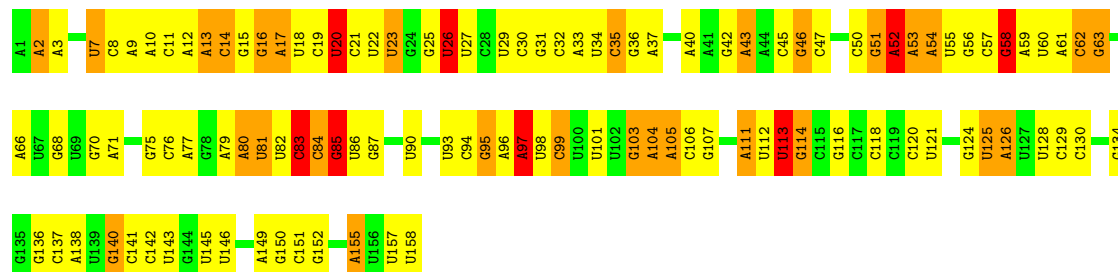
• Molecule 37: 5S ribosomal RNA

Chain 7: 27% 38% 27% 7%





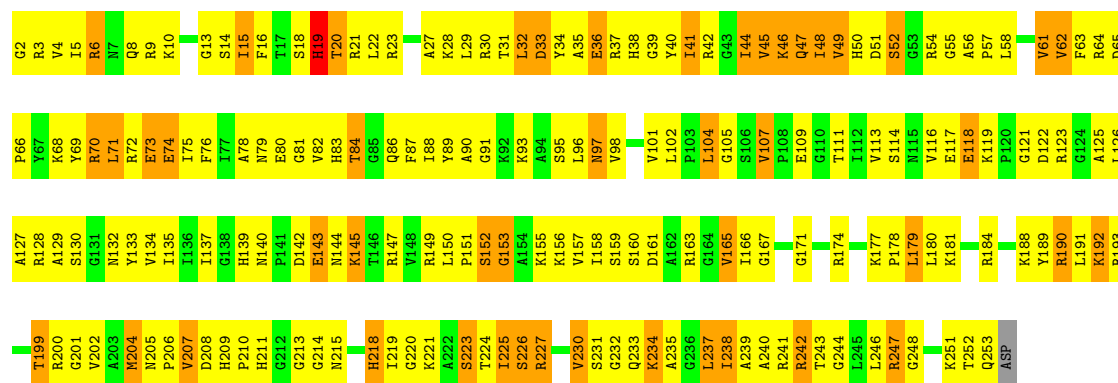
• Molecule 38: 5.8S ribosomal RNA



• Molecule 38: 5.8S ribosomal RNA

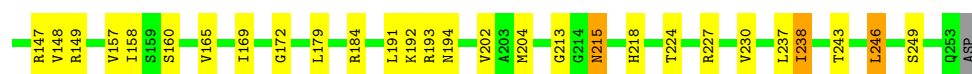


• Molecule 39: 60S ribosomal protein L2-A



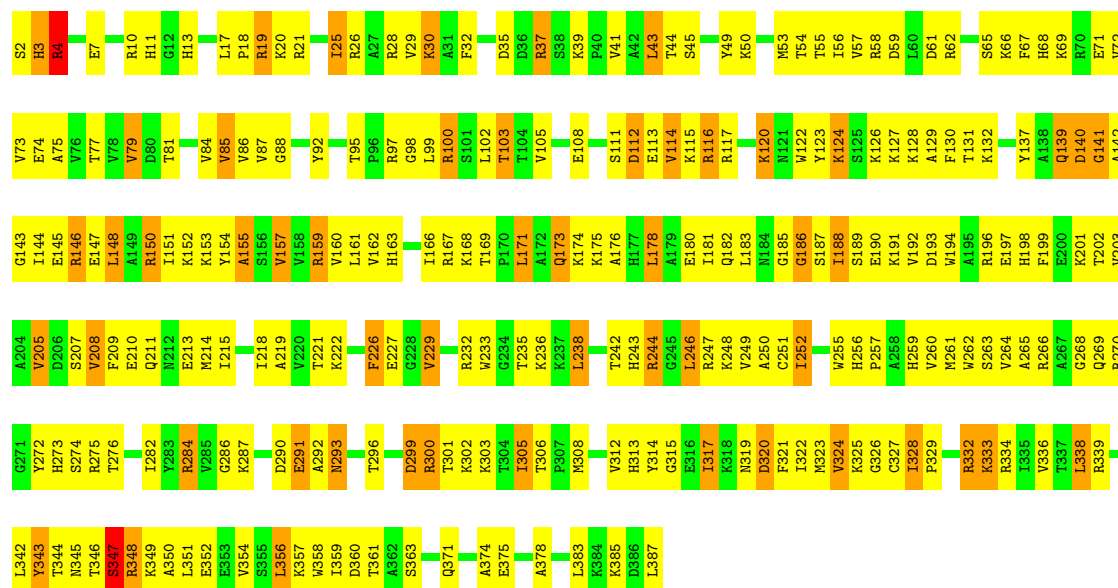
• Molecule 39: 60S ribosomal protein L2-A





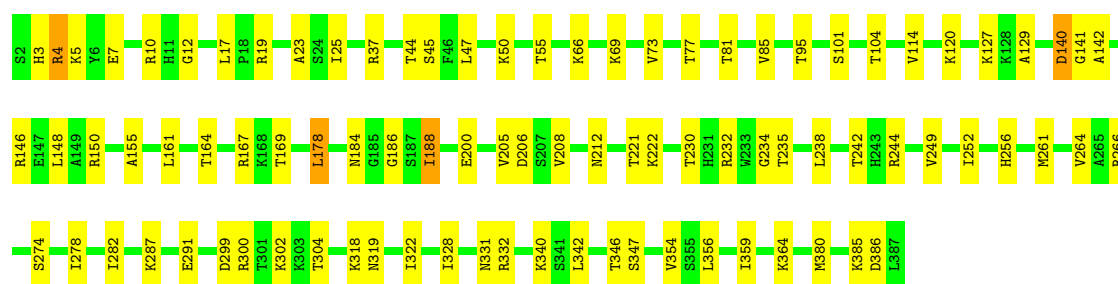
• Molecule 40: 60S ribosomal protein L3

Chain L3: 35% 51% 14%



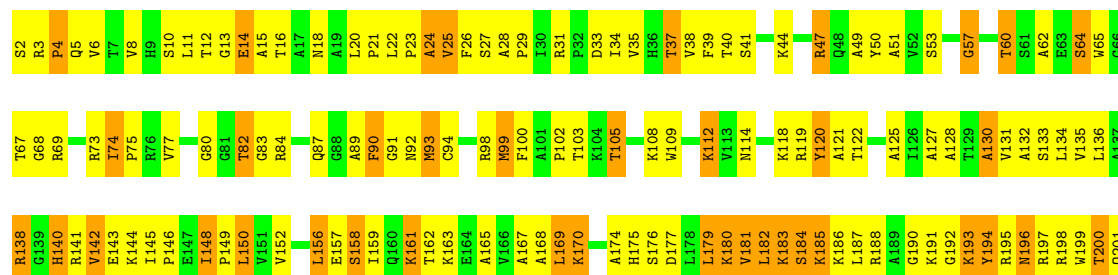
• Molecule 40: 60S ribosomal protein L3

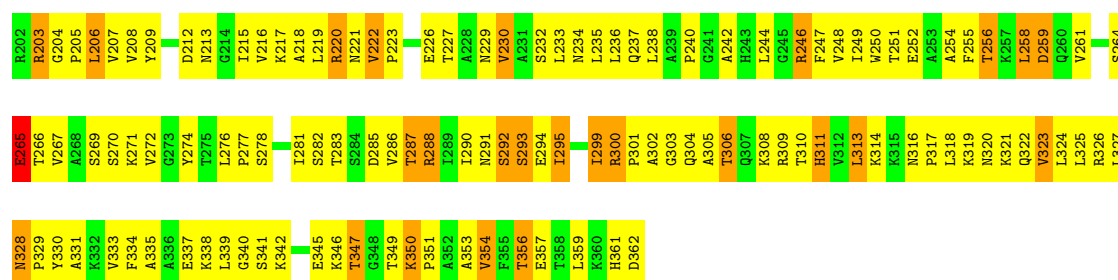
Chain l3: 77% 22%



• Molecule 41: 60S ribosomal protein L4-A

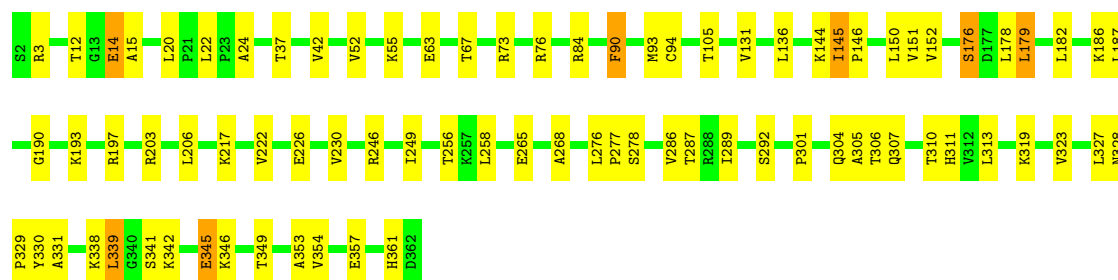
Chain L4: 28% 54% 18%





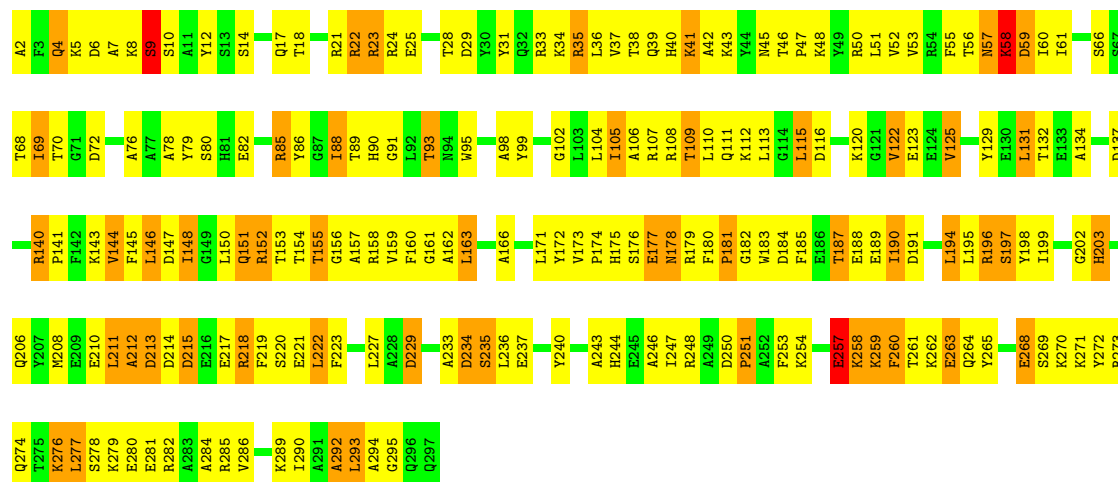
• Molecule 41: 60S ribosomal protein L4-A

Chain 14: 77% 21%



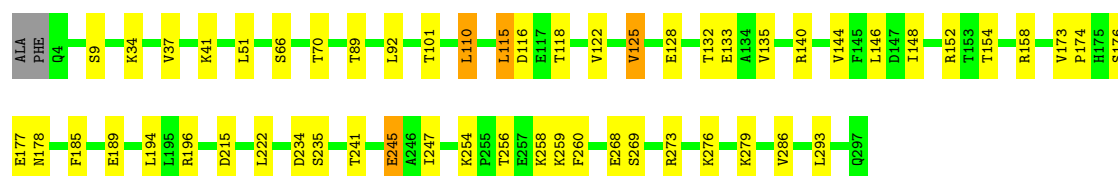
• Molecule 42: 60S ribosomal protein L5

Chain L5: 31% 50% 18%



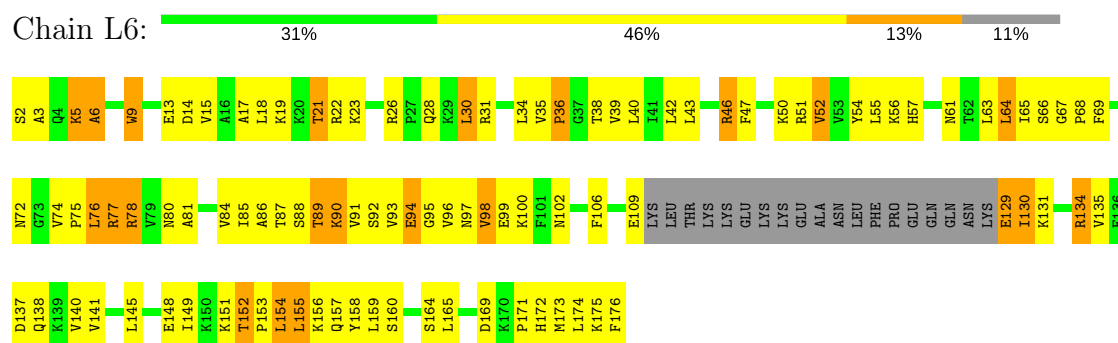
• Molecule 42: 60S ribosomal protein L5

Chain 15: 81% 17%



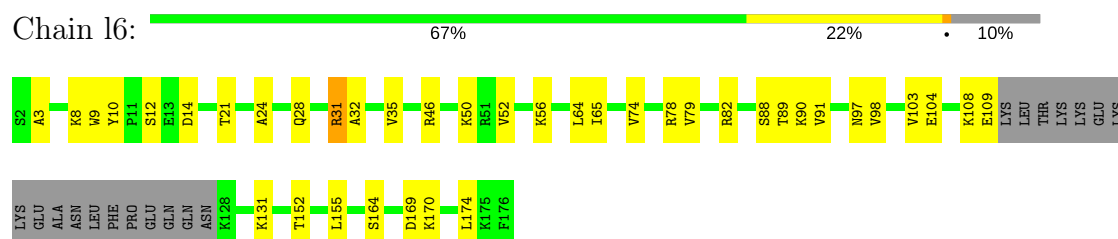
• Molecule 43: 60S ribosomal protein L6-A

Chain L6:



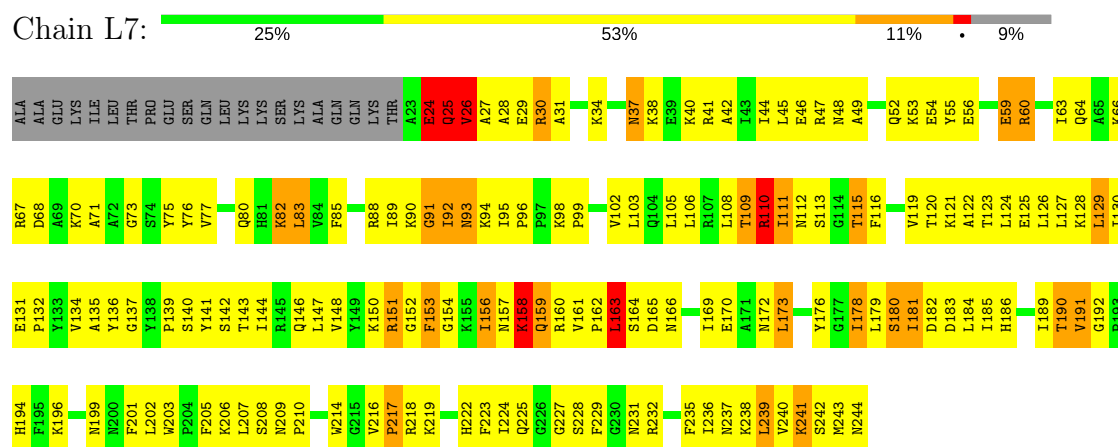
- Molecule 43: 60S ribosomal protein L6-A

Chain 16:



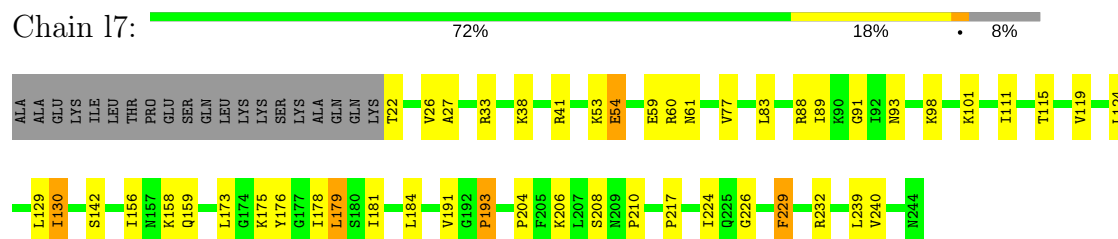
- Molecule 44: 60S ribosomal protein L7-A

Chain L7:



- Molecule 44: 60S ribosomal protein L7-A

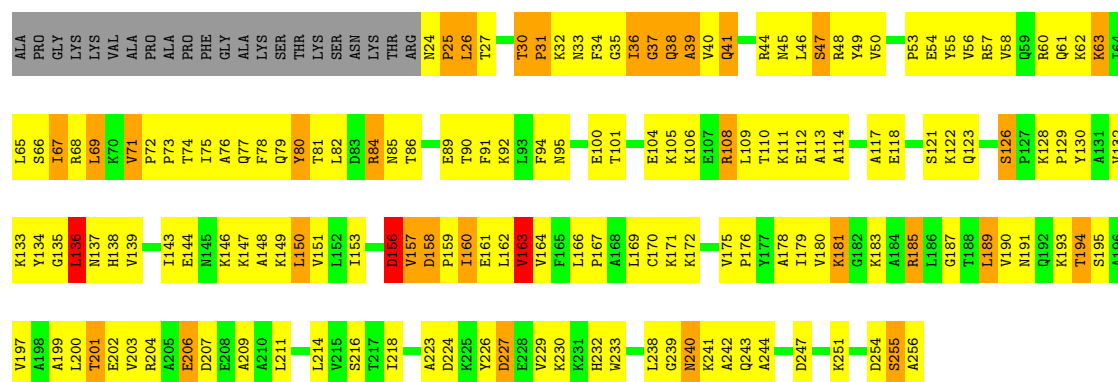
Chain 17:



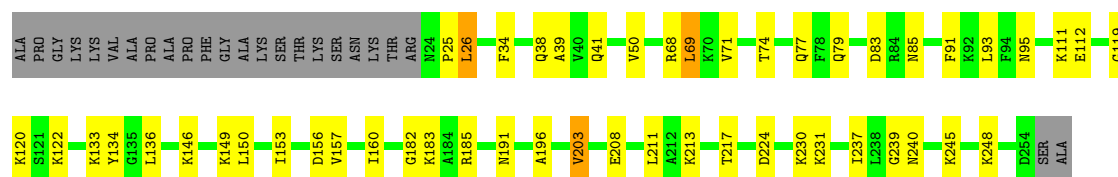
- Molecule 45: 60S ribosomal protein L8-A

Chain L8:

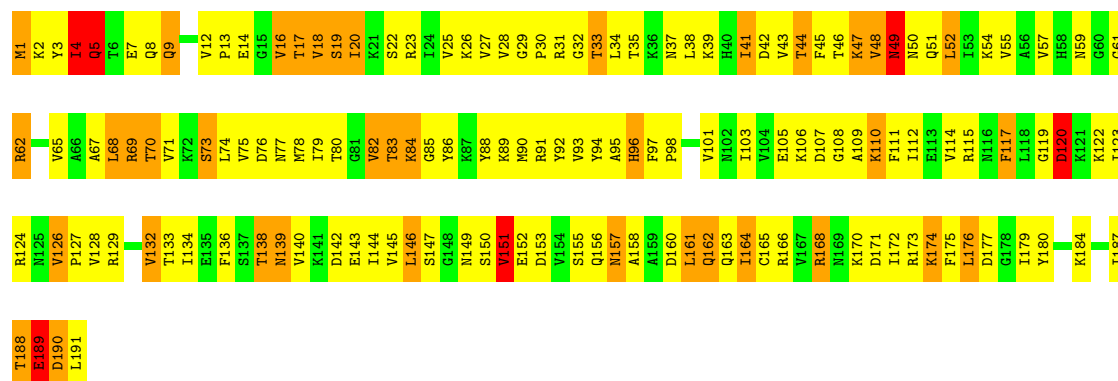




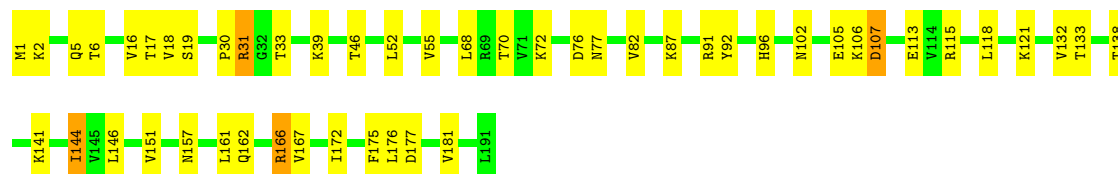
• Molecule 45: 60S ribosomal protein L8-A



• Molecule 46: 60S ribosomal protein L9-A

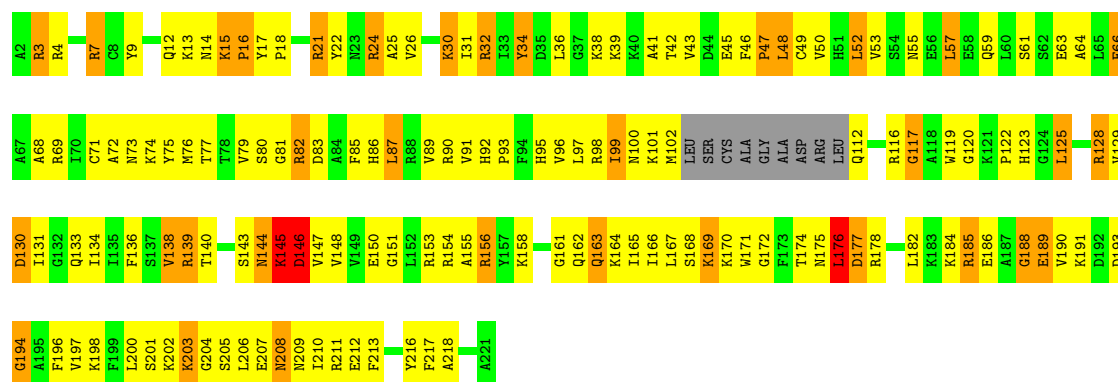


• Molecule 46: 60S ribosomal protein L9-A



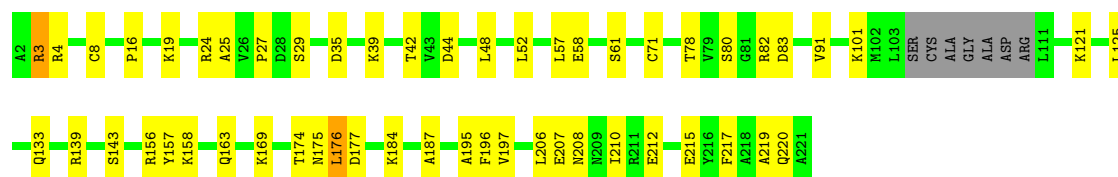
• Molecule 47: 60S ribosomal protein L10





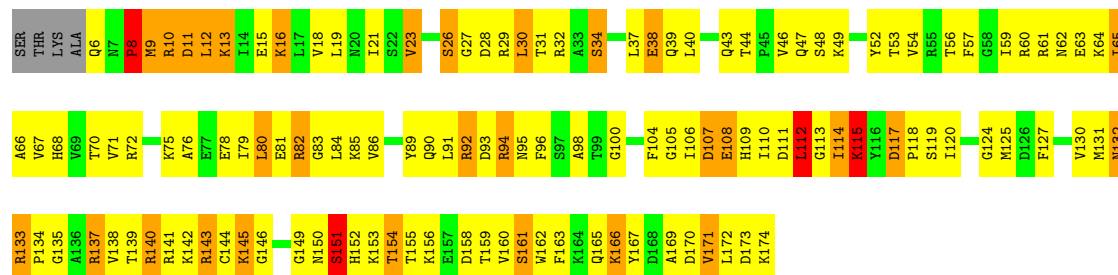
• Molecule 47: 60S ribosomal protein L10

Chain m0: 73% 23% ..



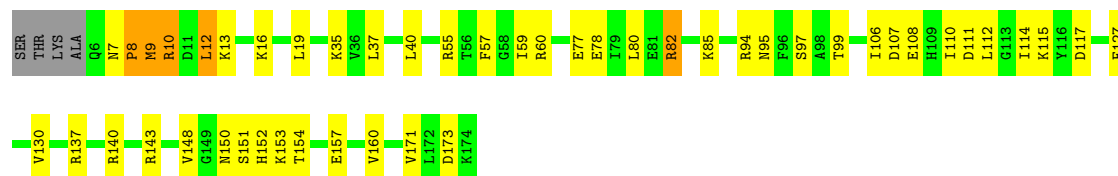
• Molecule 48: 60S ribosomal protein L11-B

Chain M1: 24% 54% 17% ..



• Molecule 48: 60S ribosomal protein L11-B

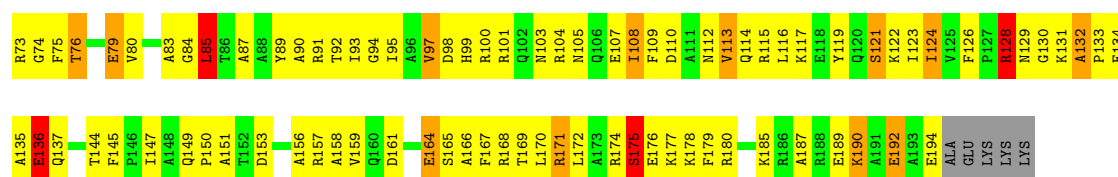
Chain m1: 70% 25% ..



• Molecule 49: 60S ribosomal protein L13-A

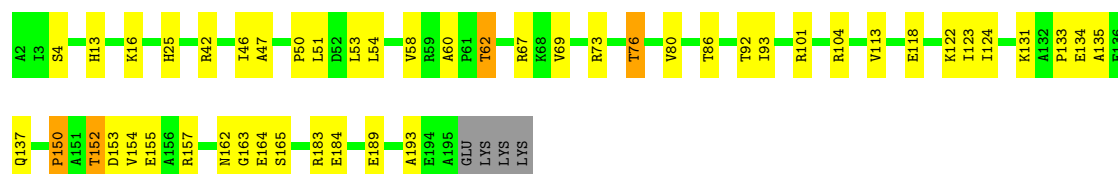
Chain M3: 30% 52% 12% ..





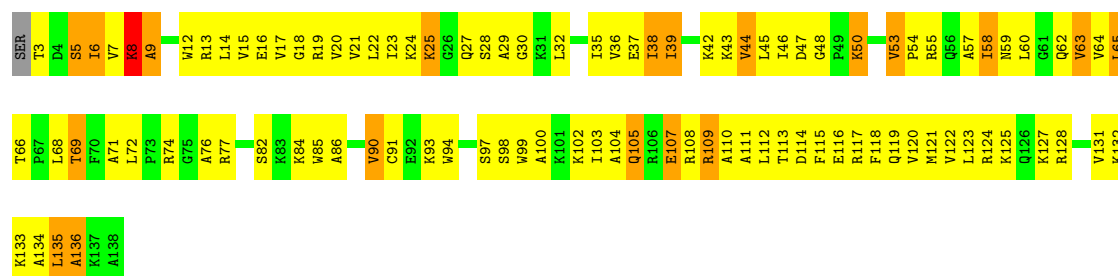
• Molecule 49: 60S ribosomal protein L13-A

Chain m3: 74% 22%



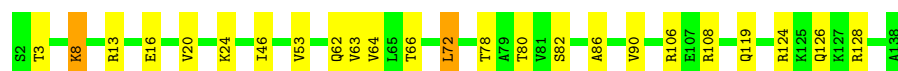
• Molecule 50: 60S ribosomal protein L14-A

Chain M4: 26% 58% 14%



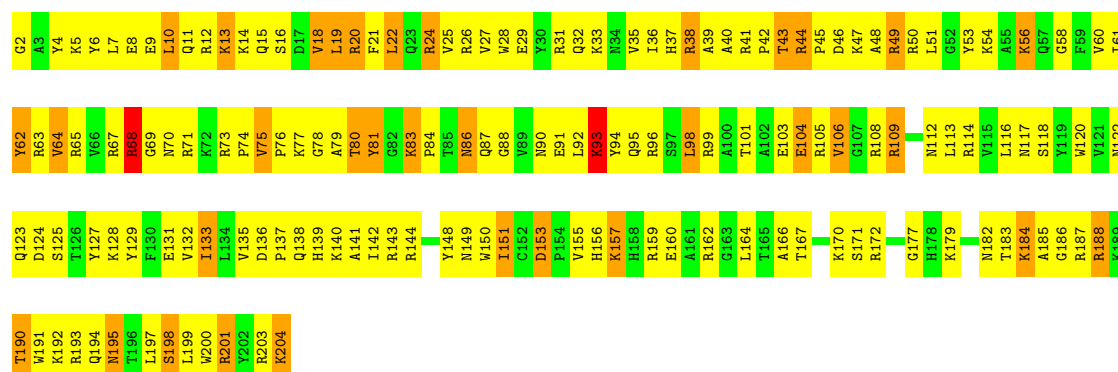
• Molecule 50: 60S ribosomal protein L14-A

Chain m4: 82% 16%

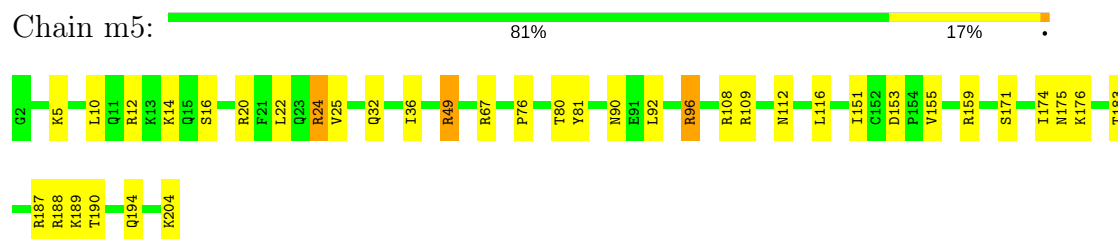


• Molecule 51: 60S ribosomal protein L15-A

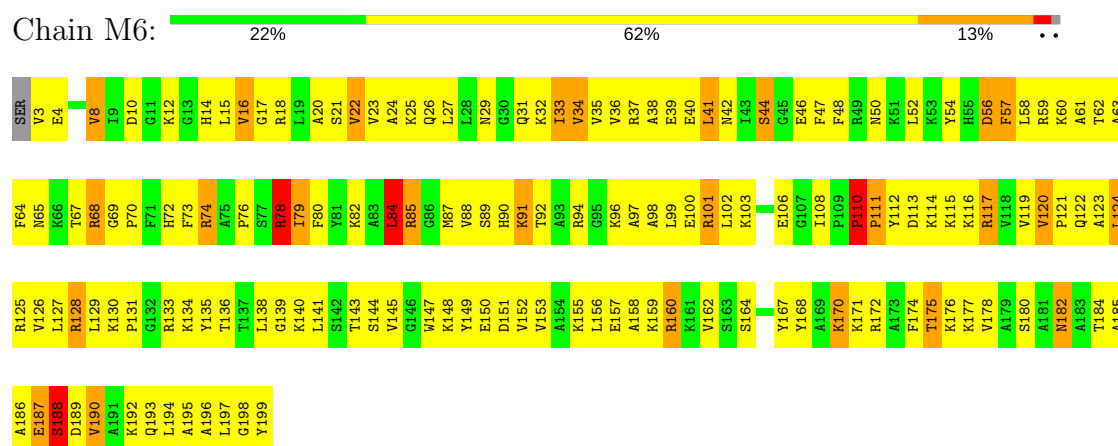
Chain M5: 23% 59% 17%



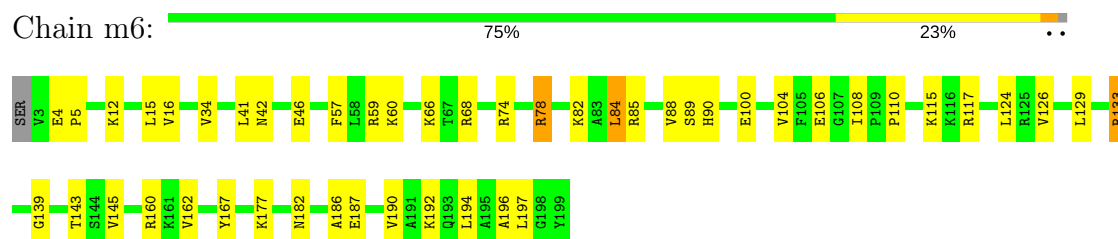
- Molecule 51: 60S ribosomal protein L15-A



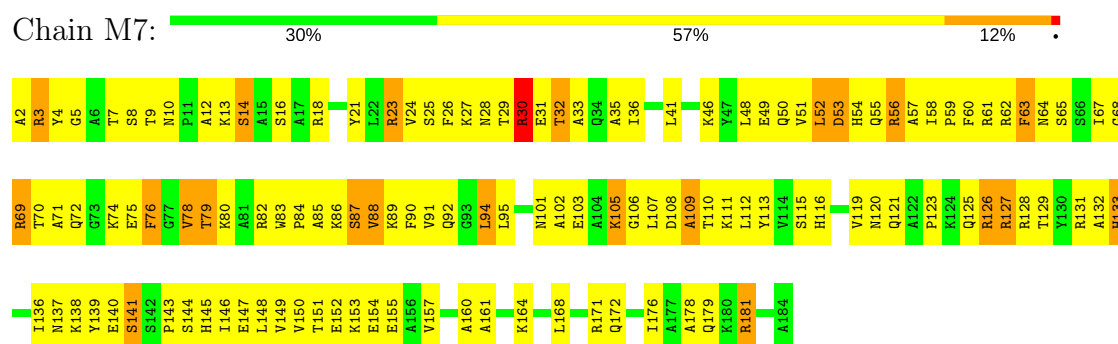
- Molecule 52: 60S ribosomal protein L16-A



- Molecule 52: 60S ribosomal protein L16-A

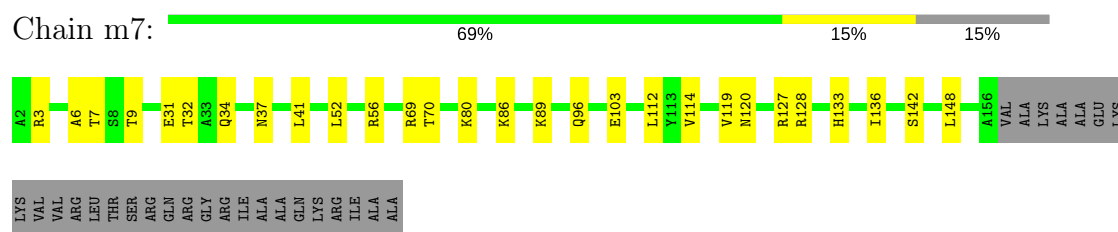


- Molecule 53: 60S ribosomal protein L17-A



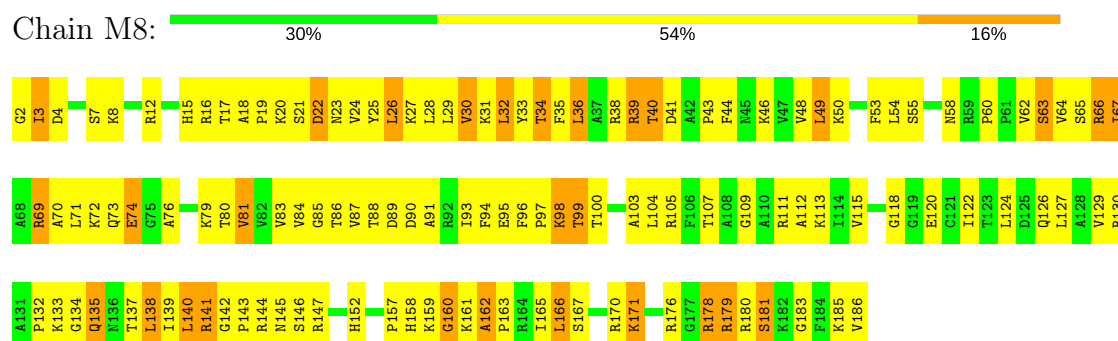
- Molecule 53: 60S ribosomal protein L17-A

Chain m7:



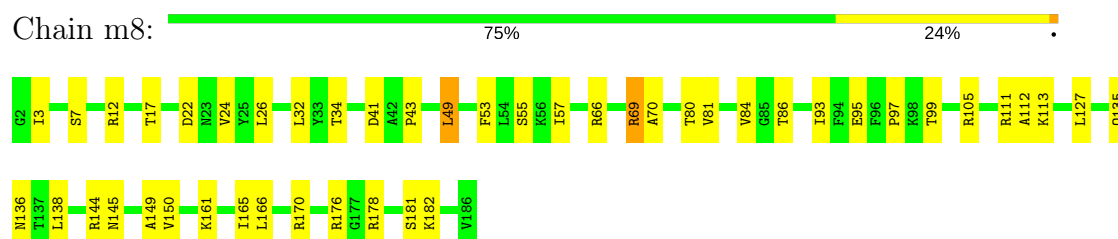
• Molecule 54: 60S ribosomal protein L18-A

Chain M8:



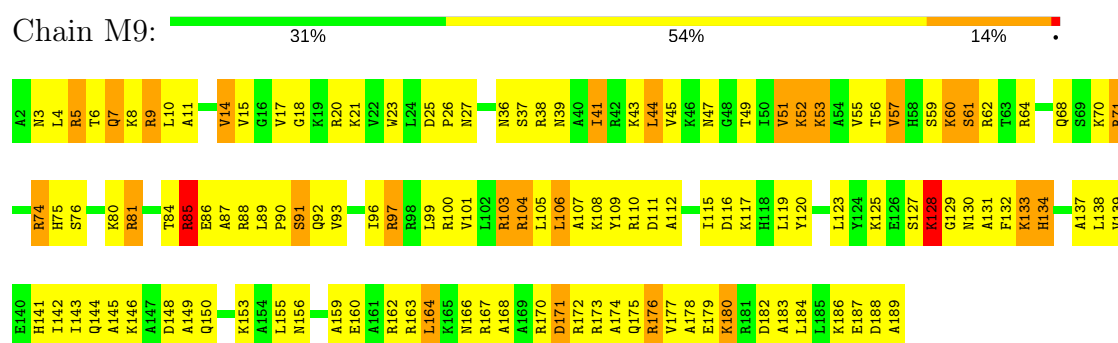
• Molecule 54: 60S ribosomal protein L18-A

Chain m8:



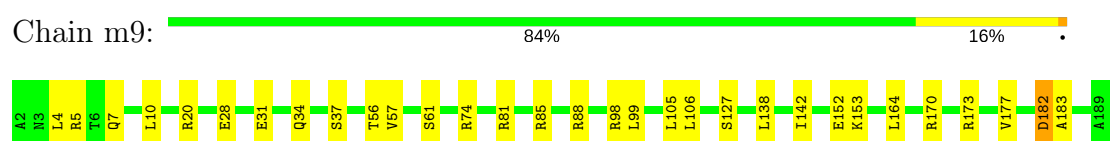
• Molecule 55: 60S ribosomal protein L19-A

Chain M9:

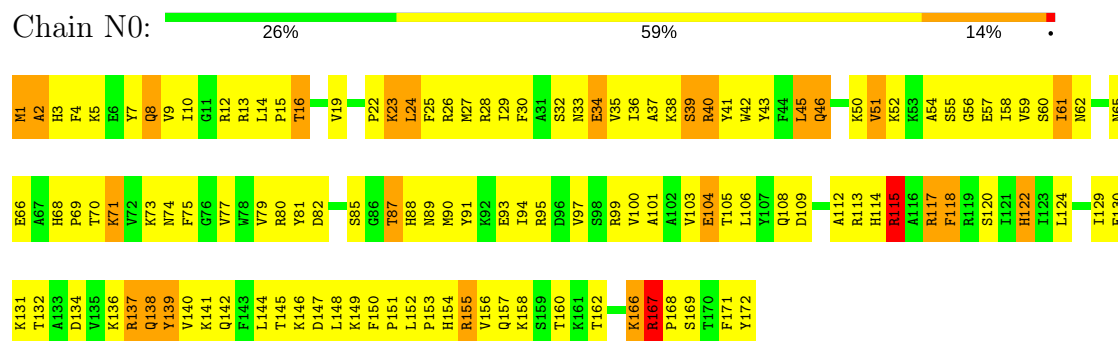


• Molecule 55: 60S ribosomal protein L19-A

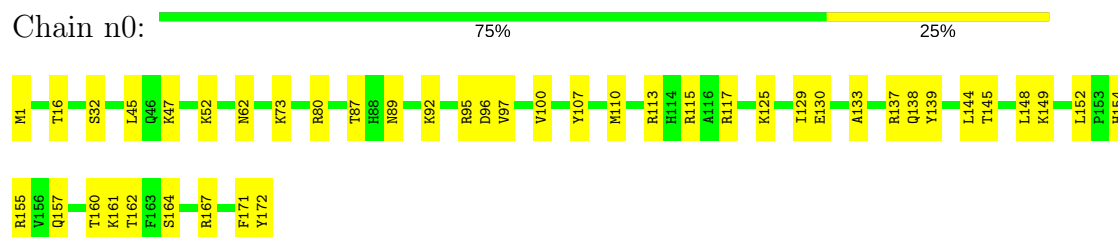
Chain m9:



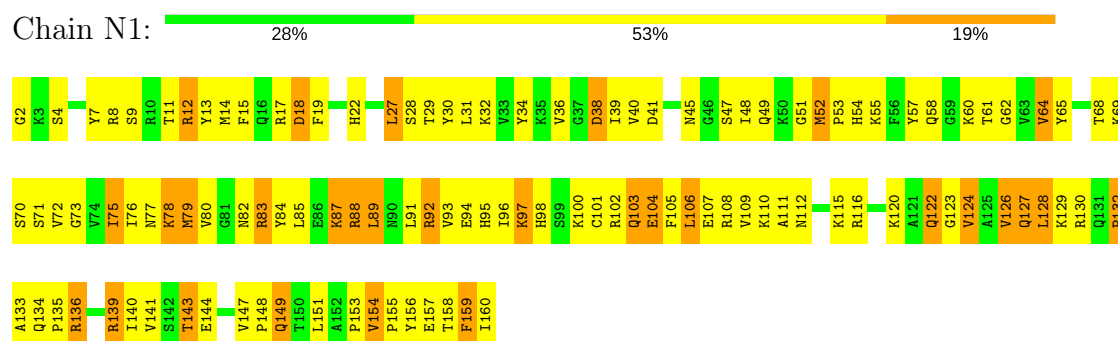
• Molecule 56: 60S ribosomal protein L20-A



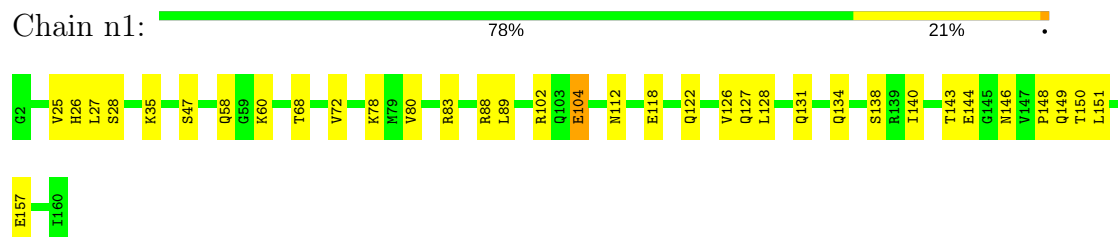
• Molecule 56: 60S ribosomal protein L20-A



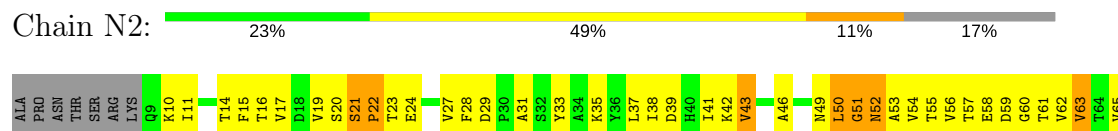
• Molecule 57: 60S ribosomal protein L21-A

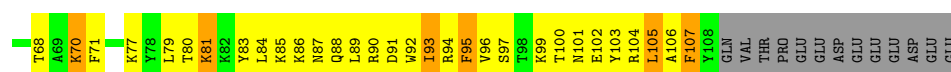


• Molecule 57: 60S ribosomal protein L21-A



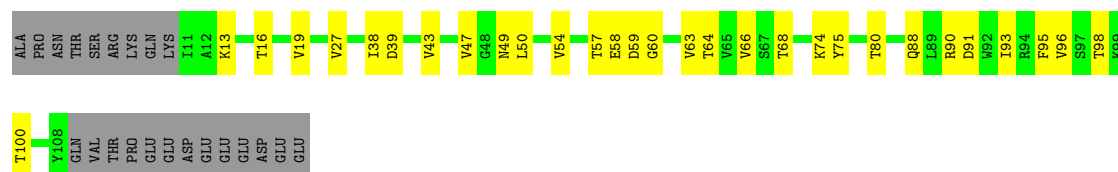
• Molecule 58: 60S ribosomal protein L22-A





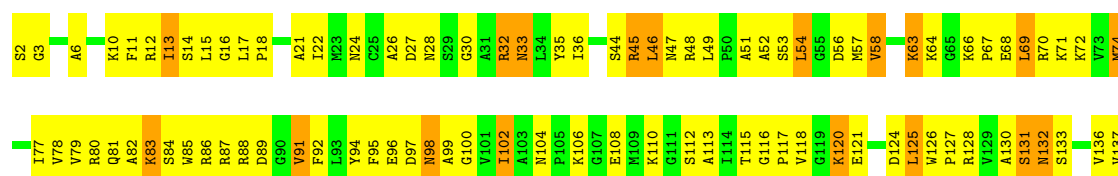
• Molecule 58: 60S ribosomal protein L22-A

Chain n2: 57% 25% 18%



• Molecule 59: 60S ribosomal protein L23-A

Chain N3: 32% 54% 13%



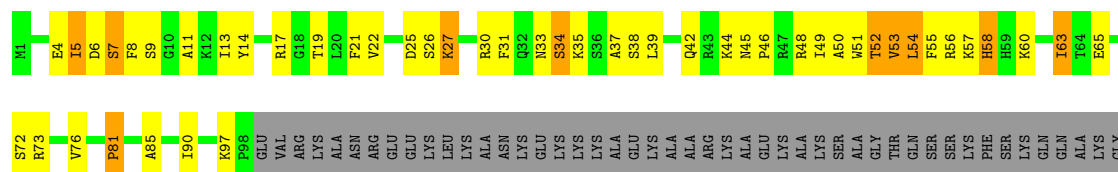
• Molecule 59: 60S ribosomal protein L23-A

Chain n3: 81% 18%



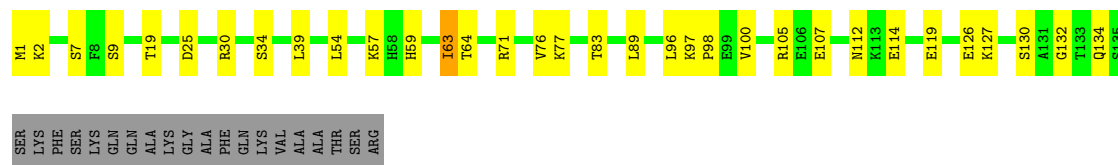
• Molecule 60: 60S ribosomal protein L24-A

Chain N4: 32% 25% 6% 37%



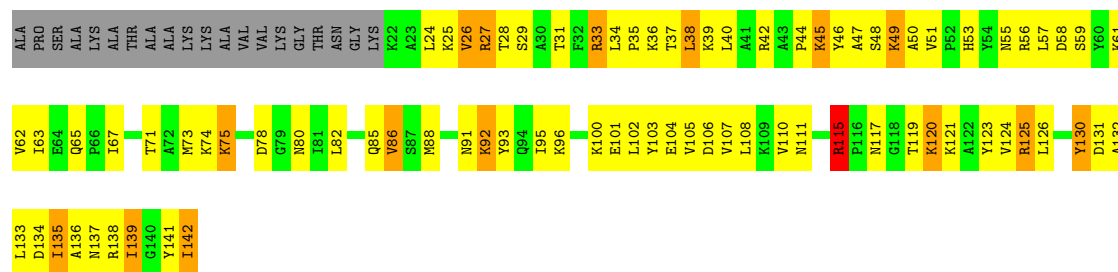
• Molecule 60: 60S ribosomal protein L24-A

Chain n4: 66% 21% 13%



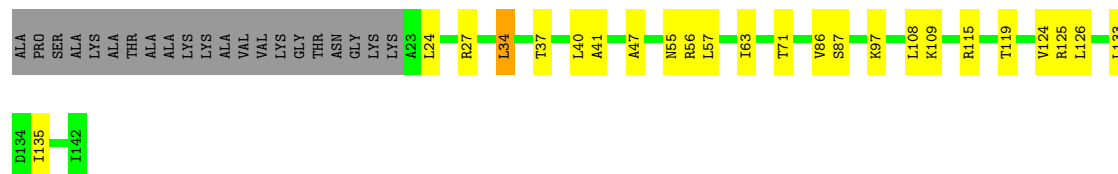
- Molecule 61: 60S ribosomal protein L25

Chain N5:  28% 47% 11% • 14%



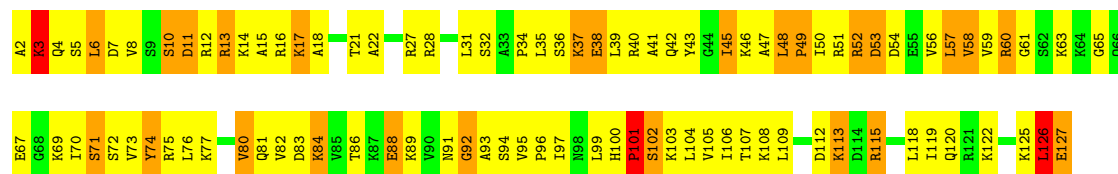
- Molecule 61: 60S ribosomal protein L25

Chain n5:  68% 16% 15%




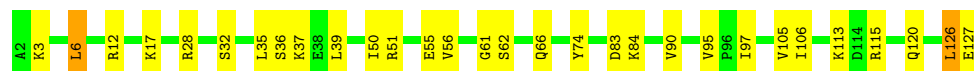
- Molecule 62: 60S ribosomal protein L26-A

Chain N6: 



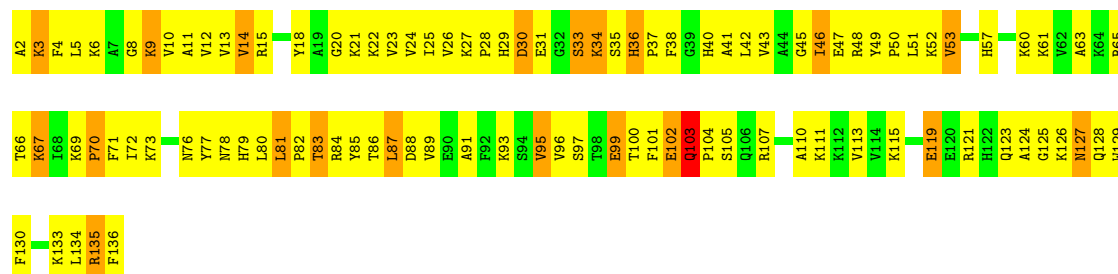
- Molecule 62: 60S ribosomal protein L26-A

Chain n6:  76% 22%



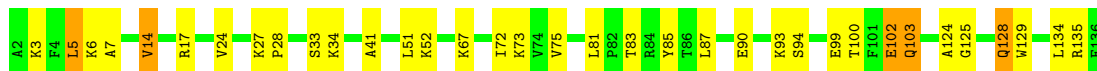
- Molecule 63: 60S ribosomal protein L27-A

Chain N7:  24% 60% 15%




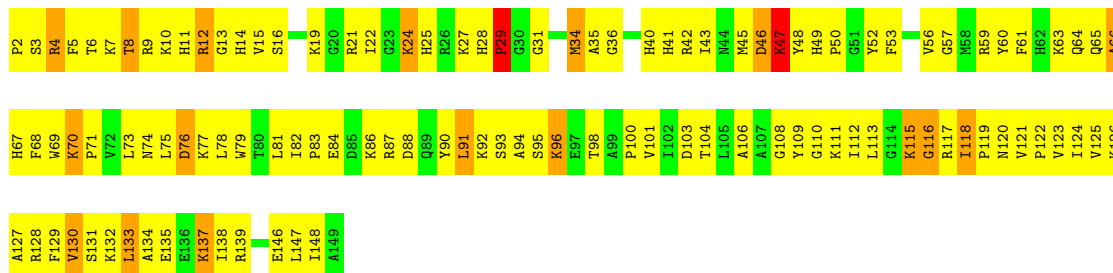
- Molecule 63: 60S ribosomal protein L27-A

Chain n7: 




- Molecule 64: 60S ribosomal protein L28

Chain N8: 



- Molecule 64: 60S ribosomal protein L28

Chain n8: 



- Molecule 65: 60S ribosomal protein L29

Chain N9: 



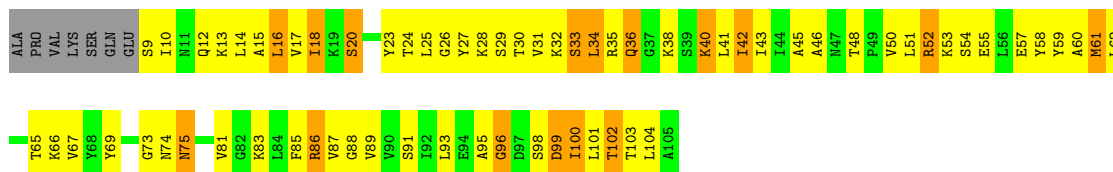
- Molecule 65: 60S ribosomal protein L29

Chain n9: 




- Molecule 66: 60S ribosomal protein L30

Chain O0: 



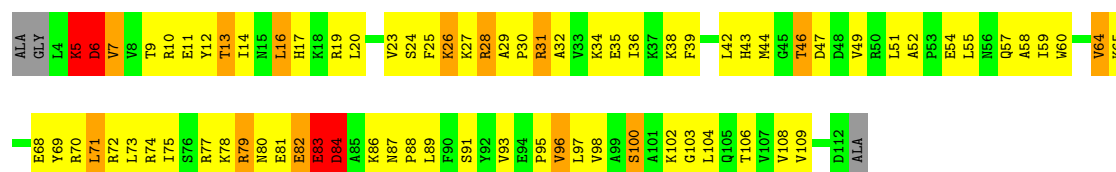
- Molecule 66: 60S ribosomal protein L30

Chain o0: 



- Molecule 67: 60S ribosomal protein L31-A

Chain O1: 



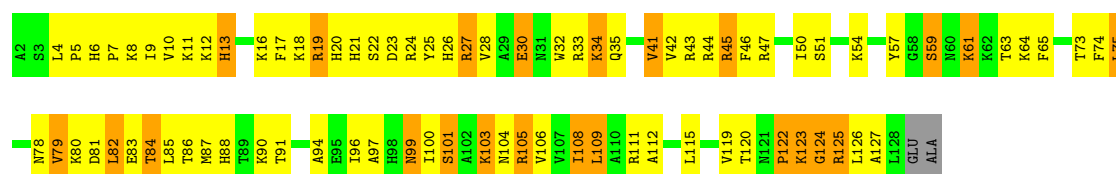
- Molecule 67: 60S ribosomal protein L31-A

Chain o1: 



- Molecule 68: 60S ribosomal protein L32

Chain O2: 



- Molecule 68: 60S ribosomal protein L32

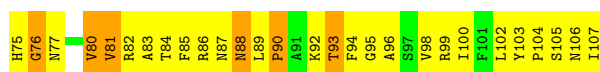
Chain o2: 



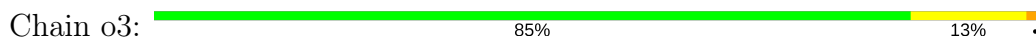
- Molecule 69: 60S ribosomal protein L33-A

Chain O3: 





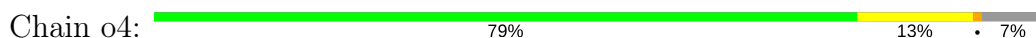
- Molecule 69: 60S ribosomal protein L33-A



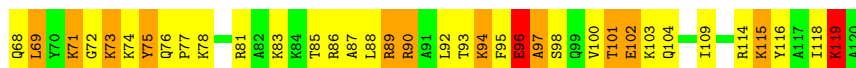
- Molecule 70: 60S ribosomal protein L34-A



- Molecule 70: 60S ribosomal protein L34-A



- Molecule 71: 60S ribosomal protein L35-A



- Molecule 71: 60S ribosomal protein L35-A



- Molecule 72: 60S ribosomal protein L36-A





- Molecule 72: 60S ribosomal protein L36-A

Chain o6: 70% 29%



- Molecule 73: 60S ribosomal protein L37-A

Chain O7: 31% 54% 15%



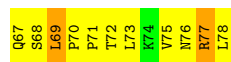
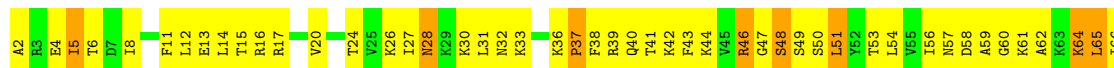
- Molecule 73: 60S ribosomal protein L37-A

Chain o7: 77% 21%



- Molecule 74: 60S ribosomal protein L38

Chain O8: 23% 64% 13%



- Molecule 74: 60S ribosomal protein L38

Chain o8: 75% 23%



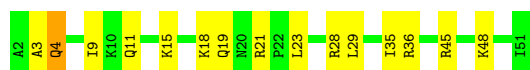
- Molecule 75: 60S ribosomal protein L39

Chain O9: 32% 56% 12%



- Molecule 75: 60S ribosomal protein L39

Chain o9:  70% 28% .



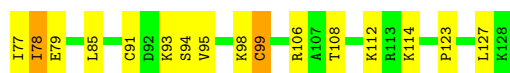
- Molecule 76: Ubiquitin-60S ribosomal protein L40

Chain Q0:  33% 52% 12% .



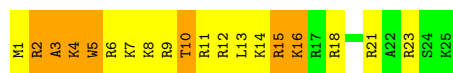
- Molecule 76: Ubiquitin-60S ribosomal protein L40

Chain q0:  69% 27% .



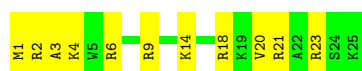
- Molecule 77: 60S ribosomal protein L41-A

Chain Q1:  24% 48% 28%

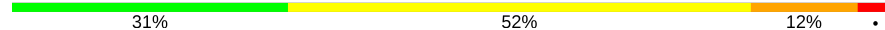


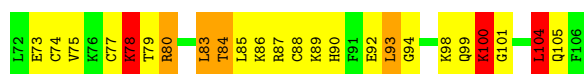
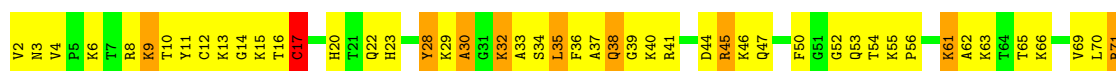
- Molecule 77: 60S ribosomal protein L41-A

Chain q1:  56% 44%




- Molecule 78: 60S ribosomal protein L42-A

Chain Q2:  31% 52% 12% .

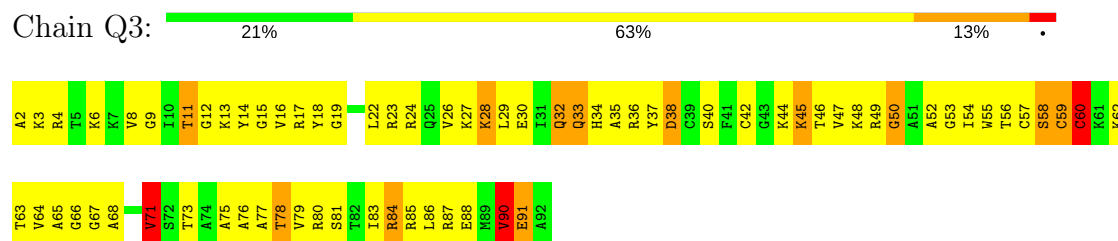


- Molecule 78: 60S ribosomal protein L42-A

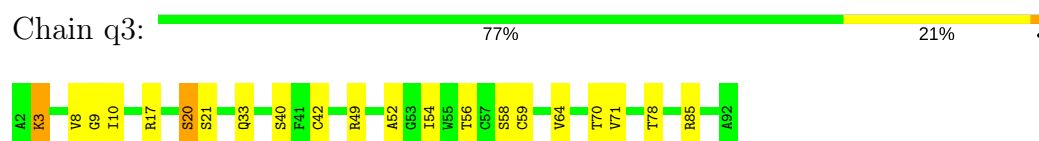
Chain q2:  76% 21% .



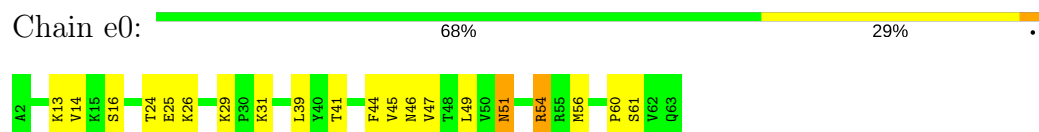
- Molecule 79: 60S ribosomal protein L43-A



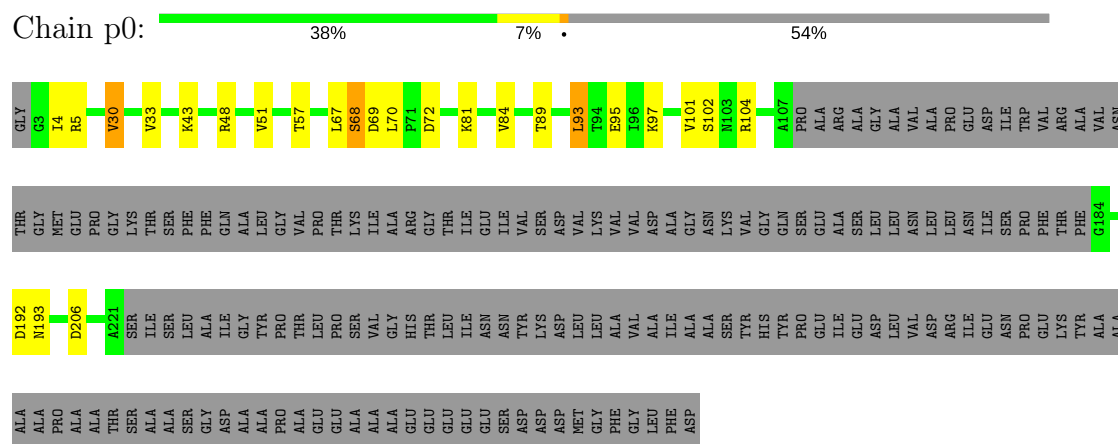
- Molecule 79: 60S ribosomal protein L43-A



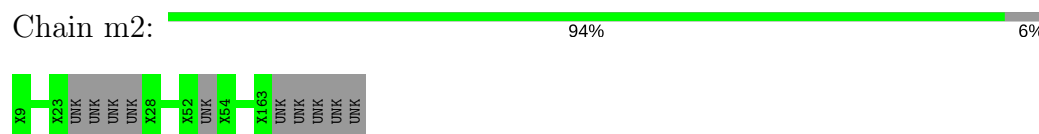
- Molecule 80: 40S ribosomal protein S30-A



- Molecule 81: 60S acidic ribosomal protein P0



- Molecule 82: unknown protein chain m2



- Molecule 83: unknown protein chain p1



There are no outlier residues recorded for this chain.

- Molecule 84: unknown protein chain p2

Chain p2:  100%

There are no outlier residues recorded for this chain.

4 Data and refinement statistics

EDS failed to run properly - this section is therefore incomplete.

Property	Value	Source
Space group	P 1 21 1	Depositor
Cell constants a, b, c, α , β , γ	435.64Å 286.76Å 303.26Å 90.00° 98.72° 90.00°	Depositor
Resolution (Å)	299.76 – 3.45	Depositor
% Data completeness (in resolution range)	100.0 (299.76-3.45)	Depositor
R_{merge}	0.45	Depositor
R_{sym}	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ ¹	1.38 (at 3.41Å)	Xtriage
Refinement program	PHENIX (phenix.refine: dev_1702)	Depositor
R, R_{free}	0.196 , 0.260	Depositor
Wilson B-factor (Å ²)	93.1	Xtriage
Anisotropy	0.148	Xtriage
L-test for twinning ²	$\langle L \rangle = 0.44$, $\langle L^2 \rangle = 0.27$	Xtriage
Estimated twinning fraction	No twinning to report.	Xtriage
Total number of atoms	411214	wwPDB-VP
Average B, all atoms (Å ²)	81.0	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.*

¹Intensities estimated from amplitudes.

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

5 Model quality ⓘ

5.1 Standard geometry ⓘ

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

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	2	0.74	4/41698 (0.0%)	1.34	362/64972 (0.6%)
1	6	0.83	14/42765 (0.0%)	1.39	467/66634 (0.7%)
2	S0	0.45	0/1617	0.68	0/2215
2	s0	0.47	0/1623	0.67	0/2222
3	S1	0.39	0/1735	0.66	1/2335 (0.0%)
3	s1	0.49	0/1748	0.68	0/2352
4	S2	0.49	0/1665	0.66	0/2263
4	s2	0.55	0/1665	0.77	0/2263
5	S3	0.50	0/1759	0.66	0/2368
5	s3	0.46	0/1759	0.64	1/2368 (0.0%)
6	S4	0.52	0/2109	0.73	0/2839
6	s4	0.53	0/2109	0.72	0/2839
7	S5	0.42	0/1629	0.64	0/2202
7	s5	0.45	0/1629	0.69	0/2202
8	S6	0.49	0/1823	0.65	0/2439
8	s6	0.59	1/1779 (0.1%)	0.72	1/2379 (0.0%)
9	S7	0.44	0/1506	0.68	0/2028
9	s7	0.45	0/1516	0.69	0/2043
10	S8	0.56	0/1514	0.74	1/2021 (0.0%)
10	s8	0.55	0/1514	0.71	0/2021
11	S9	0.48	0/1519	0.68	1/2035 (0.0%)
11	s9	0.54	0/1519	0.71	0/2035
12	C0	0.46	0/790	0.68	1/1069 (0.1%)
12	c0	0.41	0/777	0.69	3/1049 (0.3%)
13	C1	0.59	0/1240	0.76	1/1675 (0.1%)
13	c1	0.59	0/1194	0.76	0/1610
14	C2	0.41	0/900	0.65	0/1224
14	c2	0.32	0/900	0.60	0/1224
15	C3	0.52	0/1215	0.70	2/1638 (0.1%)
15	c3	0.55	0/1215	0.69	0/1638
16	C4	0.40	0/901	0.64	0/1217
16	c4	0.50	0/960	0.70	0/1290

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
17	C5	0.52	0/998	0.72	0/1341
17	c5	0.49	0/1060	0.69	1/1426 (0.1%)
18	C6	0.48	0/1125	0.68	2/1510 (0.1%)
18	c6	0.49	0/1131	0.71	0/1518
19	C7	0.48	0/935	0.68	1/1254 (0.1%)
19	c7	0.48	0/914	0.69	0/1224
20	C8	0.47	0/1211	0.70	0/1628
20	c8	0.50	0/1211	0.72	2/1628 (0.1%)
21	C9	0.44	0/1130	0.62	0/1517
21	c9	0.51	0/1130	0.67	0/1517
22	D0	0.45	0/865	0.71	1/1169 (0.1%)
22	d0	0.46	0/892	0.68	0/1205
23	D1	0.47	0/693	0.64	0/935
23	d1	0.51	0/693	0.68	0/935
24	D2	0.49	0/1038	0.71	0/1395
24	d2	0.59	0/1038	0.76	1/1395 (0.1%)
25	D3	0.61	0/1139	0.75	0/1518
25	d3	0.70	0/1139	0.83	1/1518 (0.1%)
26	D4	0.46	0/1087	0.65	0/1449
26	d4	0.55	0/1087	0.73	0/1449
27	D5	0.44	0/571	0.75	0/768
27	d5	0.45	0/566	0.68	0/761
28	D6	0.47	0/782	0.66	0/1047
28	d6	0.54	0/782	0.79	1/1047 (0.1%)
29	D7	0.42	0/620	0.65	0/838
29	d7	0.47	0/620	0.71	0/838
30	D8	0.42	0/499	0.64	0/670
30	d8	0.43	0/499	0.62	0/670
31	D9	0.53	0/452	0.67	0/600
31	d9	0.64	0/452	0.69	0/600
32	E0	0.48	0/483	0.65	0/643
33	E1	0.49	0/577	0.80	0/770
33	e1	0.42	0/619	0.75	1/822 (0.1%)
34	SR	0.40	0/2494	0.63	0/3393
34	sR	0.39	0/2495	0.58	0/3395
35	SM	0.53	0/1113	0.71	2/1502 (0.1%)
35	sM	0.49	0/682	0.69	1/921 (0.1%)
36	1	1.12	138/75394 (0.2%)	1.68	1938/117545 (1.6%)
36	5	1.11	157/75414 (0.2%)	1.68	1917/117575 (1.6%)
37	3	0.99	2/2883 (0.1%)	1.47	39/4491 (0.9%)
37	7	1.11	3/2883 (0.1%)	1.76	76/4491 (1.7%)
38	4	1.02	1/3746 (0.0%)	1.59	63/5832 (1.1%)
38	8	0.90	2/3746 (0.1%)	1.47	43/5832 (0.7%)

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
39	L2	0.68	0/1948	0.81	1/2617 (0.0%)
39	l2	0.60	0/1946	0.82	2/2614 (0.1%)
40	L3	0.74	0/3146	0.83	0/4228
40	l3	0.78	0/3146	0.86	2/4228 (0.0%)
41	L4	0.78	1/2800 (0.0%)	0.90	1/3790 (0.0%)
41	l4	0.73	1/2800 (0.0%)	0.88	3/3790 (0.1%)
42	L5	0.61	0/2425	0.73	0/3271
42	l5	0.72	0/2408	0.83	1/3248 (0.0%)
43	L6	0.78	0/1260	0.85	0/1694
43	l6	0.81	0/1269	0.87	0/1705
44	L7	0.83	0/1821	0.91	3/2451 (0.1%)
44	l7	0.87	0/1828	0.97	5/2461 (0.2%)
45	L8	0.54	0/1836	0.71	2/2481 (0.1%)
45	l8	0.48	0/1795	0.68	1/2429 (0.0%)
46	L9	0.69	0/1539	0.81	1/2073 (0.0%)
46	l9	0.76	0/1539	0.84	2/2073 (0.1%)
47	M0	0.76	0/1741	0.84	3/2335 (0.1%)
47	m0	0.76	1/1758 (0.1%)	0.82	0/2358
48	M1	0.56	0/1374	0.74	1/1842 (0.1%)
48	m1	0.70	0/1374	0.80	1/1842 (0.1%)
49	M3	0.72	0/1568	0.86	2/2106 (0.1%)
49	m3	0.64	0/1573	0.78	0/2113
50	M4	0.81	0/1068	0.88	1/1438 (0.1%)
50	m4	0.82	0/1074	0.84	1/1446 (0.1%)
51	M5	0.75	0/1757	0.86	2/2354 (0.1%)
51	m5	0.61	0/1757	0.79	2/2354 (0.1%)
52	M6	0.85	0/1585	0.92	4/2128 (0.2%)
52	m6	0.96	0/1585	0.98	4/2128 (0.2%)
53	M7	0.76	0/1443	0.83	0/1944
53	m7	0.83	0/1250	0.87	1/1683 (0.1%)
54	M8	0.75	0/1465	0.90	4/1965 (0.2%)
54	m8	0.72	0/1465	0.86	1/1965 (0.1%)
55	M9	0.60	0/1538	0.75	0/2050
55	m9	0.58	0/1538	0.69	0/2050
56	N0	0.78	0/1481	0.85	1/1990 (0.1%)
56	n0	0.88	0/1481	0.90	2/1990 (0.1%)
57	N1	0.74	0/1300	0.82	1/1743 (0.1%)
57	n1	0.81	1/1300 (0.1%)	0.88	0/1743
58	N2	0.48	0/812	0.63	0/1099
58	n2	0.49	0/794	0.68	0/1076
59	N3	0.69	0/1018	0.81	0/1369
59	n3	0.80	0/1018	0.82	0/1369
60	N4	0.62	0/712	0.74	0/958

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
60	n4	0.65	0/1052	0.75	0/1398
61	N5	0.64	0/979	0.79	1/1321 (0.1%)
61	n5	0.60	0/974	0.77	1/1314 (0.1%)
62	N6	0.72	0/1004	0.86	1/1341 (0.1%)
62	n6	0.62	0/1004	0.81	0/1341
63	N7	0.54	0/1118	0.68	0/1497
63	n7	0.48	0/1118	0.66	0/1497
64	N8	0.73	0/1204	0.85	0/1612
64	n8	0.73	0/1204	0.84	1/1612 (0.1%)
65	N9	0.69	0/473	0.83	1/629 (0.2%)
65	n9	0.71	0/473	0.88	0/629
66	O0	0.49	0/751	0.68	0/1008
66	o0	0.53	0/775	0.69	0/1040
67	O1	0.65	0/890	0.80	0/1196
67	o1	0.71	0/897	0.86	2/1205 (0.2%)
68	O2	0.79	0/1041	0.87	0/1394
68	o2	0.87	0/1041	0.89	0/1394
69	O3	0.86	0/868	0.94	0/1168
69	o3	0.87	0/868	0.90	0/1168
70	O4	0.59	0/890	0.77	1/1189 (0.1%)
70	o4	0.52	0/890	0.74	0/1189
71	O5	0.67	0/978	0.78	0/1301
71	o5	0.58	0/974	0.77	2/1297 (0.2%)
72	O6	0.61	0/778	0.77	0/1034
72	o6	0.58	0/777	0.68	0/1033
73	O7	0.74	0/696	0.87	1/923 (0.1%)
73	o7	0.64	0/696	0.79	0/923
74	O8	0.53	0/618	0.66	0/826
74	o8	0.45	0/614	0.66	0/822
75	O9	0.64	0/443	0.82	0/588
75	o9	0.62	0/443	0.78	0/588
76	Q0	0.79	0/423	0.93	0/562
76	q0	0.90	1/423 (0.2%)	0.98	0/562
77	Q1	0.67	0/234	0.87	0/300
77	q1	0.58	0/234	0.89	0/300
78	Q2	0.83	1/860 (0.1%)	0.84	0/1136
78	q2	0.73	1/860 (0.1%)	0.83	2/1136 (0.2%)
79	Q3	0.71	0/701	0.83	0/934
79	q3	0.68	0/701	0.78	0/934
80	e0	0.51	0/499	0.75	0/665
81	p0	0.48	0/1091	0.64	0/1472
All	All	0.87	329/430072 (0.1%)	1.32	4999/631360 (0.8%)

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
3	S1	0	1
7	s5	0	2
9	S7	0	1
9	s7	0	1
10	S8	0	1
11	S9	0	1
16	C4	0	1
18	C6	0	1
18	c6	0	1
19	C7	0	1
20	c8	0	1
22	d0	0	1
26	d4	0	1
27	D5	0	2
33	E1	0	1
39	L2	0	1
39	l2	0	1
40	L3	0	1
40	l3	0	1
42	L5	0	1
43	L6	0	1
44	l7	0	1
45	L8	0	1
47	M0	0	1
48	M1	0	1
48	m1	0	1
50	M4	0	1
52	M6	0	1
56	n0	0	2
64	n8	0	1
65	N9	0	1
67	O1	0	1
69	o3	0	1
80	e0	0	1
All	All	0	37

All (329) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
78	Q2	17	CYS	CB-SG	12.22	2.03	1.82
36	5	1152	G	N9-C4	-12.11	1.28	1.38
36	5	2145	A	N7-C5	-9.34	1.33	1.39
78	q2	17	CYS	CB-SG	9.04	1.97	1.82
36	5	3040	A	N9-C4	-8.70	1.32	1.37
36	5	1303	A	N9-C4	-8.37	1.32	1.37
36	5	1152	G	N3-C4	-8.29	1.29	1.35
37	3	66	A	N9-C4	-7.99	1.33	1.37
36	5	3040	A	N3-C4	-7.90	1.30	1.34
36	1	3000	A	N9-C4	-7.85	1.33	1.37
36	5	1159	A	N9-C4	-7.83	1.33	1.37
36	1	645	A	C6-N1	-7.71	1.30	1.35
36	1	2748	A	N9-C4	-7.70	1.33	1.37
36	1	1304	A	N9-C4	-7.60	1.33	1.37
36	1	3273	A	N3-C4	-7.58	1.30	1.34
36	1	2714	G	N9-C4	-7.49	1.31	1.38
36	5	607	A	N3-C4	-7.38	1.30	1.34
36	1	338	A	N7-C5	-7.35	1.34	1.39
47	m0	8	CYS	CB-SG	-7.33	1.69	1.82
36	5	706	A	N9-C4	-7.28	1.33	1.37
1	6	359	A	N9-C4	-7.17	1.33	1.37
36	5	924	G	N9-C4	-7.11	1.32	1.38
36	5	2879	C	N1-C6	-7.08	1.32	1.37
1	6	65	A	N9-C4	-7.02	1.33	1.37
36	5	1370	G	C6-N1	-7.02	1.34	1.39
36	1	1116	G	C5-C4	-7.00	1.33	1.38
36	5	3130	A	N3-C4	-6.98	1.30	1.34
36	5	397	A	N3-C4	-6.93	1.30	1.34
36	5	1135	A	N3-C4	-6.93	1.30	1.34
1	6	100	A	N7-C5	-6.91	1.35	1.39
76	q0	99	CYS	CB-SG	-6.91	1.70	1.82
36	5	522	A	N7-C5	-6.89	1.35	1.39
36	5	1205	A	N7-C5	-6.83	1.35	1.39
36	5	1432	C	N3-C4	-6.80	1.29	1.33
36	5	874	U	N1-C2	-6.75	1.32	1.38
36	5	936	A	N7-C5	-6.75	1.35	1.39
36	5	2860	U	N1-C2	6.74	1.44	1.38
36	5	924	G	N3-C4	-6.72	1.30	1.35
36	1	3091	A	N9-C4	-6.71	1.33	1.37
36	5	1192	C	N1-C2	6.68	1.46	1.40
36	5	2910	A	N3-C4	-6.68	1.30	1.34
36	1	2355	G	N7-C5	-6.66	1.35	1.39
36	5	1332	A	N7-C5	-6.63	1.35	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	1	806	A	N9-C4	-6.59	1.33	1.37
36	5	375	A	N9-C4	-6.59	1.33	1.37
36	1	1165	A	N9-C4	-6.56	1.33	1.37
36	5	2726	C	N3-C4	-6.54	1.29	1.33
36	5	2639	G	N7-C5	-6.54	1.35	1.39
36	5	1432	C	N1-C6	-6.52	1.33	1.37
36	1	1149	G	N3-C4	-6.51	1.30	1.35
36	1	342	A	N9-C4	-6.49	1.33	1.37
36	5	2368	A	N7-C5	-6.44	1.35	1.39
1	6	417	A	N9-C4	6.41	1.41	1.37
36	1	2823	G	N9-C8	-6.36	1.33	1.37
36	1	66	A	N9-C4	-6.31	1.34	1.37
36	1	972	A	C5-C4	-6.30	1.34	1.38
36	5	2358	A	N9-C4	-6.28	1.34	1.37
1	6	1746	A	N7-C5	-6.25	1.35	1.39
36	5	61	A	C6-N1	-6.25	1.31	1.35
36	1	71	A	N3-C4	-6.23	1.31	1.34
36	5	924	G	C2-N3	-6.22	1.27	1.32
36	5	2640	A	N9-C4	-6.22	1.34	1.37
36	1	1154	A	N7-C5	-6.21	1.35	1.39
36	5	647	A	N7-C5	-6.21	1.35	1.39
38	8	104	A	N9-C4	-6.20	1.34	1.37
36	1	2200	U	C4-O4	6.17	1.28	1.23
36	1	3006	A	N9-C4	-6.17	1.34	1.37
36	5	1172	G	N3-C4	-6.17	1.31	1.35
1	2	793	A	N9-C4	6.15	1.41	1.37
36	5	808	A	N3-C4	-6.15	1.31	1.34
37	7	102	A	N9-C4	-6.14	1.34	1.37
36	1	969	C	N1-C6	-6.13	1.33	1.37
36	1	2762	A	N3-C4	-6.12	1.31	1.34
36	1	936	A	N9-C4	-6.12	1.34	1.37
36	5	423	A	C5-C6	-6.12	1.35	1.41
36	1	1330	A	N9-C4	-6.11	1.34	1.37
36	5	2704	A	N9-C4	-6.11	1.34	1.37
36	5	1152	G	C5-C6	-6.10	1.36	1.42
36	1	1103	A	N7-C5	6.09	1.43	1.39
36	1	2679	A	N9-C4	-6.09	1.34	1.37
36	1	1406	A	C5-C6	-6.09	1.35	1.41
36	1	2363	A	N9-C4	-6.09	1.34	1.37
36	1	585	A	C6-N1	-6.08	1.31	1.35
36	1	1394	A	N9-C4	-6.07	1.34	1.37
36	1	1103	A	N9-C4	6.07	1.41	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	5	2971	A	N9-C4	6.06	1.41	1.37
37	7	85	G	N7-C5	-6.05	1.35	1.39
36	1	71	A	N9-C4	-6.03	1.34	1.37
36	1	355	A	N9-C4	-6.02	1.34	1.37
36	1	2409	G	N9-C8	-6.02	1.33	1.37
36	5	1348	U	N1-C2	6.00	1.44	1.38
36	1	647	A	N7-C5	-6.00	1.35	1.39
36	5	1103	A	N3-C4	5.99	1.38	1.34
36	5	519	A	C5-C6	-5.99	1.35	1.41
36	1	3006	A	N3-C4	-5.97	1.31	1.34
36	5	2932	U	C2-N3	-5.97	1.33	1.37
36	5	1149	G	N3-C4	-5.96	1.31	1.35
36	5	2138	A	N7-C5	-5.96	1.35	1.39
1	2	1124	A	N9-C4	-5.95	1.34	1.37
36	1	1103	A	N3-C4	5.94	1.38	1.34
36	5	343	U	N1-C2	-5.94	1.33	1.38
36	1	638	C	N1-C6	-5.92	1.33	1.37
36	5	1467	A	N9-C4	-5.92	1.34	1.37
41	L4	199	TRP	CB-CG	-5.91	1.39	1.50
36	1	953	G	C5-C6	-5.91	1.36	1.42
36	1	2656	A	N7-C5	-5.91	1.35	1.39
36	5	807	A	N7-C5	-5.89	1.35	1.39
36	5	2377	G	C6-N1	-5.88	1.35	1.39
36	1	1476	G	C6-N1	-5.88	1.35	1.39
36	5	522	A	C5-C6	-5.87	1.35	1.41
36	5	1370	G	N3-C4	-5.86	1.31	1.35
36	1	668	G	N3-C4	-5.86	1.31	1.35
36	5	2282	U	N1-C2	-5.86	1.33	1.38
36	5	2643	A	N9-C4	-5.85	1.34	1.37
36	1	659	G	C5-C4	-5.85	1.34	1.38
36	1	2207	A	N9-C4	5.84	1.41	1.37
36	5	1130	A	C5-C4	-5.83	1.34	1.38
36	1	1148	G	N7-C5	-5.83	1.35	1.39
36	5	2994	A	N3-C4	-5.82	1.31	1.34
36	5	917	A	N7-C5	-5.80	1.35	1.39
36	5	921	A	N3-C4	-5.79	1.31	1.34
36	5	1432	C	C2-N3	-5.79	1.31	1.35
36	1	45	A	N3-C4	-5.78	1.31	1.34
36	1	2333	C	N3-C4	-5.78	1.29	1.33
36	1	676	G	N9-C4	5.76	1.42	1.38
36	5	1306	G	C5-C6	-5.74	1.36	1.42
36	1	2986	U	N1-C2	-5.72	1.33	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	1	1047	A	N3-C4	-5.72	1.31	1.34
36	1	2147	A	N9-C4	-5.72	1.34	1.37
36	5	1159	A	N3-C4	-5.72	1.31	1.34
36	5	40	A	N7-C5	-5.72	1.35	1.39
36	5	3245	A	C5-C6	-5.72	1.35	1.41
36	1	695	C	N1-C6	-5.72	1.33	1.37
36	5	652	G	N3-C4	-5.72	1.31	1.35
36	1	2200	U	C2-N3	5.71	1.41	1.37
36	5	1849	C	N1-C2	-5.70	1.34	1.40
37	7	96	U	N1-C2	-5.70	1.33	1.38
36	5	1135	A	C6-N1	-5.70	1.31	1.35
36	1	1369	A	N7-C5	-5.70	1.35	1.39
36	5	884	A	N9-C4	-5.70	1.34	1.37
36	5	931	C	N3-C4	-5.69	1.29	1.33
36	5	652	G	N1-C2	-5.68	1.33	1.37
36	1	947	G	N7-C5	-5.67	1.35	1.39
36	1	2313	A	N9-C4	-5.67	1.34	1.37
36	1	928	C	N3-C4	-5.67	1.29	1.33
36	5	2295	A	C5-C6	-5.65	1.35	1.41
36	5	2851	A	N9-C4	-5.65	1.34	1.37
36	1	970	A	N3-C4	-5.64	1.31	1.34
36	1	653	A	C5-C6	-5.64	1.35	1.41
36	5	2970	C	N1-C6	-5.63	1.33	1.37
36	5	1152	G	N7-C5	-5.62	1.35	1.39
57	n1	104	GLU	CB-CG	5.61	1.62	1.52
36	1	2352	A	N9-C4	5.61	1.41	1.37
36	5	2248	C	N1-C6	-5.61	1.33	1.37
36	5	2113	A	N9-C4	-5.60	1.34	1.37
36	5	2377	G	N1-C2	-5.59	1.33	1.37
36	5	2966	G	N7-C5	-5.58	1.35	1.39
36	1	585	A	N3-C4	-5.58	1.31	1.34
36	1	2983	C	N3-C4	-5.58	1.30	1.33
36	1	979	U	N1-C2	5.58	1.43	1.38
36	1	908	G	N7-C5	-5.56	1.35	1.39
36	5	1462	A	N9-C4	-5.56	1.34	1.37
36	5	1902	G	N3-C4	-5.56	1.31	1.35
36	5	880	G	N7-C5	-5.55	1.35	1.39
36	5	2279	A	N9-C4	-5.55	1.34	1.37
36	1	187	A	N9-C4	5.55	1.41	1.37
36	1	1119	C	N3-C4	-5.55	1.30	1.33
36	1	2812	C	N1-C6	-5.55	1.33	1.37
36	1	2422	C	N3-C4	-5.54	1.30	1.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	5	1200	A	N7-C5	-5.54	1.35	1.39
36	1	1371	G	N3-C4	-5.54	1.31	1.35
36	1	940	G	C6-N1	-5.53	1.35	1.39
36	1	2296	A	N9-C4	-5.52	1.34	1.37
36	1	1891	A	N9-C4	-5.48	1.34	1.37
36	5	3000	A	N9-C4	-5.48	1.34	1.37
36	5	3209	A	C5-C4	5.48	1.42	1.38
36	1	1304	A	N3-C4	-5.48	1.31	1.34
36	1	218	G	N9-C8	-5.47	1.34	1.37
36	1	342	A	N3-C4	-5.47	1.31	1.34
36	1	1119	C	N1-C6	-5.47	1.33	1.37
36	5	3094	A	N9-C4	-5.47	1.34	1.37
36	1	1159	A	N9-C4	-5.46	1.34	1.37
36	5	3052	G	C6-N1	-5.46	1.35	1.39
36	1	100	A	N3-C4	-5.46	1.31	1.34
36	1	925	A	N3-C4	-5.46	1.31	1.34
36	5	672	A	N3-C4	-5.45	1.31	1.34
36	5	2840	C	N1-C6	-5.44	1.33	1.37
1	6	1137	A	N9-C4	-5.43	1.34	1.37
36	5	2626	A	N9-C4	-5.43	1.34	1.37
36	5	1429	G	N9-C4	-5.43	1.33	1.38
36	5	1143	A	N9-C4	-5.42	1.34	1.37
36	5	1370	G	N9-C8	-5.42	1.34	1.37
36	5	1081	U	N1-C2	5.42	1.43	1.38
36	5	2386	A	N7-C5	-5.41	1.36	1.39
36	5	2131	A	N7-C5	-5.40	1.36	1.39
36	5	942	U	C4-O4	5.40	1.27	1.23
36	5	3091	A	N3-C4	-5.40	1.31	1.34
36	1	2401	A	N3-C4	5.39	1.38	1.34
36	5	2872	A	N9-C4	-5.39	1.34	1.37
36	1	1153	A	C6-N1	-5.39	1.31	1.35
36	5	34	A	N9-C4	-5.37	1.34	1.37
36	5	1432	C	N1-C2	-5.37	1.34	1.40
36	5	3209	A	C5-C6	5.37	1.45	1.41
36	5	1174	G	C5-C4	-5.37	1.34	1.38
36	1	585	A	N9-C4	-5.37	1.34	1.37
1	6	337	G	C2-N3	5.36	1.37	1.32
36	5	2891	U	C2-N3	-5.36	1.34	1.37
1	6	542	A	N7-C5	-5.35	1.36	1.39
36	5	2386	A	C5-C6	-5.35	1.36	1.41
36	5	2610	G	N7-C5	-5.35	1.36	1.39
36	1	2386	A	N7-C5	-5.35	1.36	1.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	1	645	A	C6-N6	-5.34	1.29	1.33
36	1	1143	A	N7-C5	-5.34	1.36	1.39
36	5	947	G	N3-C4	-5.34	1.31	1.35
36	5	2910	A	N9-C4	-5.33	1.34	1.37
36	5	519	A	N7-C5	-5.33	1.36	1.39
36	5	655	C	N3-C4	-5.33	1.30	1.33
36	5	349	A	C5-C4	-5.33	1.35	1.38
36	1	635	G	C5-C4	-5.33	1.34	1.38
36	1	1714	A	N9-C4	-5.33	1.34	1.37
36	1	1351	U	N1-C2	5.32	1.43	1.38
36	5	2873	U	N1-C2	5.31	1.43	1.38
36	5	2918	G	N7-C5	-5.31	1.36	1.39
36	5	965	A	C6-N1	-5.30	1.31	1.35
36	1	98	G	N3-C4	-5.30	1.31	1.35
36	1	965	A	N9-C4	-5.30	1.34	1.37
36	1	1116	G	C6-N1	-5.29	1.35	1.39
36	5	43	A	N9-C4	-5.28	1.34	1.37
36	1	218	G	N7-C5	-5.28	1.36	1.39
41	14	94	CYS	CB-SG	-5.28	1.73	1.81
36	1	921	A	N3-C4	-5.28	1.31	1.34
36	1	2811	A	C6-N1	-5.27	1.31	1.35
36	1	1116	G	N3-C4	-5.26	1.31	1.35
36	1	642	U	C4-O4	5.26	1.27	1.23
36	1	3209	A	C5-C4	5.26	1.42	1.38
36	1	610	G	N9-C4	-5.25	1.33	1.38
36	1	2875	U	N1-C2	5.25	1.43	1.38
36	1	1149	G	N9-C8	-5.25	1.34	1.37
36	5	929	A	N3-C4	-5.25	1.31	1.34
36	1	1148	G	N9-C8	-5.24	1.34	1.37
36	5	3245	A	N3-C4	-5.24	1.31	1.34
36	5	3209	A	N7-C5	5.24	1.42	1.39
36	1	2244	A	N3-C4	-5.24	1.31	1.34
36	5	2976	A	N3-C4	-5.24	1.31	1.34
36	1	2386	A	C5-C6	-5.23	1.36	1.41
36	1	2627	C	N1-C6	-5.23	1.34	1.37
36	5	2131	A	C5-C6	-5.23	1.36	1.41
36	5	1061	A	N3-C4	-5.23	1.31	1.34
36	5	3081	C	N3-C4	-5.22	1.30	1.33
36	5	2302	G	C6-N1	-5.22	1.35	1.39
36	1	39	A	N9-C4	-5.21	1.34	1.37
36	5	2866	U	N1-C2	5.21	1.43	1.38
36	1	2986	U	C2-N3	-5.21	1.34	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	1	2524	A	N9-C4	-5.21	1.34	1.37
36	5	3091	A	C6-N1	-5.21	1.31	1.35
1	2	971	A	N7-C5	-5.20	1.36	1.39
36	1	1165	A	N3-C4	-5.20	1.31	1.34
36	5	887	G	C5-C6	-5.20	1.37	1.42
36	5	2910	A	C5-C4	-5.20	1.35	1.38
1	6	390	G	N7-C5	-5.20	1.36	1.39
1	6	436	A	N7-C5	-5.20	1.36	1.39
36	1	98	G	C6-N1	-5.20	1.35	1.39
36	1	2797	C	N1-C6	-5.19	1.34	1.37
36	1	670	C	N1-C6	-5.19	1.34	1.37
36	5	2813	A	N7-C5	-5.19	1.36	1.39
36	1	2232	A	N9-C4	-5.17	1.34	1.37
36	5	1107	C	N1-C6	-5.17	1.34	1.37
36	5	980	A	C5-C6	5.17	1.45	1.41
36	5	2610	G	N3-C4	-5.17	1.31	1.35
36	5	2917	G	C5-C4	-5.16	1.34	1.38
36	1	970	A	N9-C4	-5.16	1.34	1.37
36	1	1001	G	C6-N1	5.16	1.43	1.39
36	5	2403	G	C6-N1	5.15	1.43	1.39
36	1	360	G	N7-C5	-5.15	1.36	1.39
1	6	1746	A	N9-C4	-5.15	1.34	1.37
37	3	39	C	N3-C4	-5.15	1.30	1.33
1	6	384	G	N9-C4	-5.15	1.33	1.38
36	1	2682	C	N3-C4	-5.14	1.30	1.33
36	5	3012	A	N9-C4	-5.14	1.34	1.37
36	5	921	A	C6-N1	-5.13	1.31	1.35
36	1	35	A	N3-C4	-5.13	1.31	1.34
36	1	1116	G	N7-C5	-5.12	1.36	1.39
36	5	2814	G	C6-N1	-5.12	1.35	1.39
36	1	2762	A	C6-N1	-5.11	1.31	1.35
1	2	863	A	N9-C4	-5.11	1.34	1.37
36	1	2365	C	N1-C6	-5.11	1.34	1.37
36	5	424	G	N7-C5	-5.09	1.36	1.39
36	5	3294	A	N3-C4	-5.09	1.31	1.34
36	5	607	A	C6-N1	-5.09	1.31	1.35
36	5	2698	G	N9-C4	-5.09	1.33	1.38
36	1	817	A	N9-C4	5.08	1.40	1.37
36	5	2145	A	C5-C6	-5.08	1.36	1.41
36	5	1126	G	N7-C5	-5.08	1.36	1.39
36	5	3008	A	N3-C4	-5.08	1.31	1.34
36	1	1153	A	C5-C6	-5.08	1.36	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
36	1	1903	U	C4-O4	5.08	1.27	1.23
36	1	2620	G	C5-C4	-5.07	1.34	1.38
36	5	957	C	N3-C4	-5.07	1.30	1.33
1	6	1746	A	C5-C4	-5.07	1.35	1.38
36	5	1134	G	N3-C4	-5.06	1.31	1.35
36	1	2377	G	N9-C4	-5.06	1.33	1.38
36	5	1332	A	C5-C6	-5.06	1.36	1.41
38	4	37	A	N3-C4	5.06	1.37	1.34
36	5	2385	G	N9-C4	-5.06	1.33	1.38
36	1	584	G	N7-C5	-5.06	1.36	1.39
36	1	2309	A	C5-C6	-5.05	1.36	1.41
36	1	962	A	N3-C4	-5.05	1.31	1.34
36	1	2247	G	N7-C5	-5.05	1.36	1.39
36	5	1200	A	C5-C6	-5.05	1.36	1.41
36	5	1161	G	N3-C4	5.04	1.39	1.35
36	5	2826	U	C2-N3	-5.04	1.34	1.37
36	1	806	A	C5-C4	-5.04	1.35	1.38
36	5	2338	C	N1-C6	-5.04	1.34	1.37
36	5	567	G	N7-C5	-5.04	1.36	1.39
36	1	1330	A	N3-C4	-5.04	1.31	1.34
36	1	2601	A	N9-C4	-5.03	1.34	1.37
36	5	2703	A	N7-C5	-5.03	1.36	1.39
36	5	2911	A	N7-C5	-5.03	1.36	1.39
36	1	343	U	C2-N3	-5.02	1.34	1.37
36	5	2359	C	N1-C6	-5.02	1.34	1.37
38	8	20	U	N1-C2	-5.02	1.34	1.38
1	6	100	A	C5-C6	-5.02	1.36	1.41
36	1	1192	C	N1-C2	5.01	1.45	1.40
36	5	650	C	N3-C4	-5.01	1.30	1.33
36	5	1189	C	N1-C6	-5.01	1.34	1.37
36	1	2358	A	C5-C6	-5.01	1.36	1.41
8	s6	83	CYS	CB-SG	-5.01	1.73	1.81
36	1	1890	U	N1-C2	-5.01	1.34	1.38
36	1	2178	A	N3-C4	-5.01	1.31	1.34

All (4999) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	1152	G	N3-C4-N9	-19.38	114.37	126.00
36	5	1152	G	N3-C4-C5	17.87	137.53	128.60
36	5	1152	G	C2-N3-C4	-16.63	103.58	111.90
36	1	645	A	N1-C6-N6	-16.45	108.73	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	343	U	O5'-P-OP2	-14.33	92.80	105.70
36	5	1437	C	C6-N1-C2	-13.92	114.73	120.30
36	5	2818	U	O5'-P-OP1	-13.76	93.32	105.70
36	5	1192	C	N1-C2-O2	13.27	126.86	118.90
36	5	1200	A	N1-C6-N6	12.86	126.31	118.60
36	1	2714	G	N3-C4-C5	12.71	134.95	128.60
36	1	959	C	C6-N1-C2	12.63	125.35	120.30
36	5	1129	A	O5'-P-OP2	-12.62	94.34	105.70
36	5	2760	C	N3-C4-C5	12.60	126.94	121.90
36	1	2819	A	O5'-P-OP2	-12.44	94.51	105.70
36	5	1587	A	C8-N9-C4	12.21	110.68	105.80
36	1	369	A	C8-N9-C4	-12.17	100.93	105.80
36	5	1897	G	N1-C6-O6	12.09	127.15	119.90
36	5	1433	A	C8-N9-C4	-11.97	101.01	105.80
36	5	1192	C	N3-C2-O2	-11.95	113.54	121.90
36	5	344	A	O5'-P-OP1	-11.92	94.97	105.70
36	5	715	A	N1-C6-N6	-11.85	111.49	118.60
36	5	917	A	O5'-P-OP2	-11.75	95.13	105.70
36	5	940	G	O5'-P-OP1	-11.69	95.18	105.70
36	5	1152	G	C5-N7-C8	-11.67	98.47	104.30
36	1	1433	A	C8-N9-C4	-11.55	101.18	105.80
36	1	424	G	C5-C6-O6	-11.46	121.73	128.60
36	1	2374	C	C6-N1-C2	-11.45	115.72	120.30
36	5	211	A	O5'-P-OP1	-11.39	95.45	105.70
36	5	2283	G	N1-C6-O6	11.35	126.71	119.90
36	5	424	G	C5-C6-O6	-11.30	121.82	128.60
36	5	2129	U	O5'-P-OP1	-11.27	95.56	105.70
36	1	2996	U	C2-N1-C1'	11.19	131.12	117.70
36	1	3092	C	C6-N1-C2	11.11	124.74	120.30
36	1	2814	G	O5'-P-OP1	-11.10	95.71	105.70
36	1	963	G	O5'-P-OP2	-11.09	95.72	105.70
36	1	1166	G	N1-C6-O6	11.08	126.55	119.90
36	1	645	A	N9-C4-C5	11.06	110.23	105.80
36	5	1311	G	O5'-P-OP2	-10.98	95.82	105.70
36	5	923	C	C6-N1-C2	10.97	124.69	120.30
36	1	1166	G	C5-C6-O6	-10.92	122.05	128.60
36	5	2760	C	C6-N1-C2	10.90	124.66	120.30
36	5	2400	G	N1-C6-O6	10.87	126.42	119.90
1	2	553	G	N1-C6-O6	10.85	126.41	119.90
36	1	2714	G	N3-C4-N9	-10.81	119.52	126.00
36	1	1151	U	C6-N1-C2	-10.80	114.52	121.00
36	1	2374	C	N3-C2-O2	-10.80	114.34	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	2709	C	C6-N1-C2	10.75	124.60	120.30
36	1	2344	U	O5'-P-OP2	-10.72	96.05	105.70
1	2	1082	C	C2-N1-C1'	10.71	130.58	118.80
36	5	924	G	N3-C4-N9	-10.68	119.59	126.00
1	6	1537	C	C6-N1-C2	-10.67	116.03	120.30
36	1	637	C	N1-C2-O2	10.67	125.30	118.90
36	5	3245	A	C2-N3-C4	-10.65	105.27	110.60
36	1	1369	A	O5'-P-OP1	-10.64	96.12	105.70
36	1	2899	C	C6-N1-C2	-10.62	116.05	120.30
36	1	1323	G	C5-C6-O6	-10.62	122.23	128.60
37	7	93	C	O5'-P-OP2	-10.61	96.15	105.70
36	5	2607	G	N1-C6-O6	10.59	126.25	119.90
36	5	907	G	C5-C6-O6	-10.56	122.26	128.60
36	5	1306	G	C5-C6-O6	-10.53	122.28	128.60
36	1	1323	G	C4-C5-N7	10.50	115.00	110.80
36	1	1303	A	N1-C6-N6	10.49	124.90	118.60
36	1	940	G	N1-C6-O6	-10.46	113.62	119.90
36	5	1152	G	N1-C6-O6	10.45	126.17	119.90
38	4	13	A	O5'-P-OP1	-10.44	96.31	105.70
1	6	1773	C	C6-N1-C2	-10.41	116.14	120.30
36	1	2355	G	N1-C6-O6	10.40	126.14	119.90
36	5	422	A	O5'-P-OP2	-10.40	96.34	105.70
1	2	577	G	N1-C6-O6	10.38	126.13	119.90
36	5	2655	U	O5'-P-OP2	-10.37	96.36	105.70
36	1	979	U	N3-C2-O2	-10.35	114.95	122.20
36	5	567	G	C6-C5-N7	-10.31	124.21	130.40
36	5	2880	U	C6-N1-C2	-10.30	114.82	121.00
36	1	1307	G	N9-C4-C5	10.28	109.51	105.40
36	5	2707	C	C6-N1-C2	10.27	124.41	120.30
36	5	887	G	C4-C5-N7	10.22	114.89	110.80
37	7	1	G	N3-C4-C5	-10.22	123.49	128.60
36	1	2869	U	O5'-P-OP1	-10.18	96.54	105.70
36	5	1149	G	N1-C6-O6	10.14	125.98	119.90
36	1	2200	U	N3-C4-O4	10.10	126.47	119.40
36	1	676	G	N3-C4-N9	10.09	132.06	126.00
36	5	2607	G	C5-C6-O6	-10.09	122.55	128.60
36	5	1507	G	O5'-P-OP1	-10.03	96.67	105.70
36	5	924	G	N3-C4-C5	9.99	133.60	128.60
36	5	410	U	C6-N1-C2	-9.99	115.01	121.00
36	5	567	G	N1-C6-O6	9.99	125.89	119.90
36	5	715	A	N9-C4-C5	9.99	109.80	105.80
36	5	1604	G	C4-N9-C1'	9.98	139.48	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2899	C	C2-N1-C1'	9.97	129.77	118.80
36	1	3344	A	N7-C8-N9	9.97	118.78	113.80
36	1	3022	G	C5-C6-O6	-9.95	122.63	128.60
36	5	2726	C	C6-N1-C2	-9.94	116.32	120.30
36	5	1306	G	C4-C5-N7	9.92	114.77	110.80
36	1	676	G	N3-C4-C5	-9.87	123.67	128.60
36	5	693	A	O5'-P-OP1	-9.84	96.84	105.70
36	5	1200	A	C6-C5-N7	-9.84	125.41	132.30
36	1	2412	G	O5'-P-OP2	-9.84	96.85	105.70
36	1	1151	U	C5-C6-N1	9.83	127.62	122.70
36	1	2395	G	N1-C6-O6	9.83	125.80	119.90
36	1	921	A	N1-C6-N6	-9.74	112.75	118.60
36	5	1604	G	C8-N9-C1'	-9.74	114.34	127.00
36	5	874	U	O5'-P-OP1	-9.72	96.95	105.70
36	5	3092	C	C6-N1-C2	9.71	124.18	120.30
36	5	2385	G	O5'-P-OP1	-9.70	96.97	105.70
36	1	709	A	N1-C6-N6	9.68	124.41	118.60
36	5	1200	A	C4-C5-C6	9.67	121.84	117.00
36	1	2881	C	C6-N1-C2	9.64	124.16	120.30
36	1	2996	U	C6-N1-C1'	-9.63	107.71	121.20
1	2	1794	A	O5'-P-OP1	-9.62	97.04	105.70
36	1	3382	U	C2-N1-C1'	9.62	129.24	117.70
36	5	2797	C	C6-N1-C2	-9.62	116.45	120.30
36	5	812	G	O5'-P-OP2	-9.61	97.05	105.70
36	5	720	A	N1-C6-N6	9.60	124.36	118.60
36	1	1192	C	C6-N1-C2	-9.59	116.46	120.30
36	1	3217	C	C2-N1-C1'	9.58	129.34	118.80
36	5	424	G	N1-C6-O6	9.58	125.65	119.90
36	5	2272	G	O4'-C1'-N9	9.56	115.85	108.20
36	1	1166	G	C4-C5-N7	9.56	114.62	110.80
36	5	3245	A	N1-C6-N6	9.55	124.33	118.60
36	5	2385	G	N1-C6-O6	9.52	125.61	119.90
36	1	1365	G	O5'-P-OP1	-9.51	97.14	105.70
36	5	2197	C	C6-N1-C2	9.50	124.10	120.30
1	6	1773	C	N3-C4-C5	-9.47	118.11	121.90
36	1	2287	C	C6-N1-C2	-9.46	116.52	120.30
36	5	1869	C	C6-N1-C2	9.45	124.08	120.30
36	1	1307	G	P-O3'-C3'	9.44	131.03	119.70
36	1	1160	C	C6-N1-C2	9.44	124.07	120.30
36	1	580	C	N1-C2-O2	-9.43	113.24	118.90
36	1	1442	U	N3-C2-O2	9.43	128.80	122.20
36	5	3245	A	C6-C5-N7	-9.42	125.70	132.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
37	7	1	G	N3-C4-N9	9.42	131.65	126.00
36	5	1152	G	C8-N9-C1'	9.40	139.23	127.00
36	1	421	G	N1-C6-O6	-9.40	114.26	119.90
36	5	519	A	N1-C6-N6	9.40	124.24	118.60
36	5	1896	A	O5'-P-OP1	-9.40	97.24	105.70
36	1	2882	U	O5'-P-OP2	-9.40	97.24	105.70
36	1	2200	U	N3-C4-C5	-9.37	108.98	114.60
36	1	948	C	C6-N1-C2	9.36	124.04	120.30
36	1	3204	C	N3-C2-O2	-9.35	115.35	121.90
36	1	676	G	C4-N9-C1'	9.33	138.63	126.50
36	1	2247	G	N1-C6-O6	9.33	125.50	119.90
36	1	1001	G	N9-C4-C5	-9.32	101.67	105.40
36	5	1055	A	O5'-P-OP2	-9.30	97.33	105.70
36	1	1495	U	C5-C6-N1	-9.29	118.06	122.70
36	1	1367	G	C8-N9-C4	9.28	110.11	106.40
36	5	1447	G	O5'-P-OP1	-9.27	97.36	105.70
36	5	1149	G	N3-C2-N2	-9.25	113.42	119.90
36	5	1152	G	C5-C6-N1	-9.23	106.88	111.50
36	1	2406	C	C6-N1-C2	9.22	123.99	120.30
36	1	2968	G	N1-C6-O6	9.22	125.43	119.90
36	1	358	G	N1-C6-O6	9.21	125.43	119.90
36	1	1389	G	N9-C4-C5	-9.21	101.72	105.40
36	1	901	G	N1-C6-O6	9.21	125.42	119.90
1	2	1773	C	C6-N1-C2	-9.20	116.62	120.30
36	1	1314	C	C6-N1-C2	-9.20	116.62	120.30
36	5	668	G	C8-N9-C4	9.18	110.07	106.40
38	4	53	A	N1-C6-N6	-9.17	113.10	118.60
36	1	3362	A	C8-N9-C4	-9.15	102.14	105.80
1	6	337	G	N3-C4-C5	-9.13	124.04	128.60
36	5	2282	U	C5-C6-N1	-9.13	118.14	122.70
36	5	1433	A	N9-C4-C5	9.10	109.44	105.80
36	1	1904	C	C6-N1-C2	-9.10	116.66	120.30
36	1	921	A	N9-C4-C5	9.05	109.42	105.80
36	5	2392	C	C6-N1-C2	9.05	123.92	120.30
36	1	1949	G	O5'-P-OP1	-9.04	97.57	105.70
36	5	2389	C	O5'-P-OP1	-9.04	97.57	105.70
36	5	2416	U	O5'-P-OP2	-9.03	97.58	105.70
38	8	26	U	N3-C2-O2	-9.02	115.89	122.20
36	5	800	G	C6-C5-N7	-9.02	124.99	130.40
36	1	690	A	N1-C6-N6	-9.01	113.19	118.60
36	1	2355	G	C6-C5-N7	-9.01	125.00	130.40
36	1	39	A	N1-C6-N6	-9.01	113.20	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2639	G	C6-C5-N7	-9.00	125.00	130.40
36	5	2413	A	C8-N9-C4	9.00	109.40	105.80
36	1	421	G	C5-C6-N1	8.99	116.00	111.50
36	5	2971	A	C2-N3-C4	8.99	115.10	110.60
36	5	1437	C	C5-C6-N1	8.98	125.49	121.00
36	1	1001	G	C5-C6-O6	-8.97	123.22	128.60
36	1	2395	G	C5-C6-O6	-8.96	123.22	128.60
36	1	3344	A	C8-N9-C4	-8.96	102.22	105.80
36	1	3243	A	N1-C6-N6	8.95	123.97	118.60
36	1	3362	A	N7-C8-N9	8.95	118.28	113.80
36	5	1417	G	C8-N9-C4	-8.94	102.82	106.40
36	5	1124	U	N3-C2-O2	-8.93	115.95	122.20
36	1	1311	G	C5-C6-O6	-8.93	123.24	128.60
36	5	965	A	C8-N9-C4	-8.93	102.23	105.80
36	5	2283	G	C5-C6-O6	-8.93	123.24	128.60
36	5	967	A	C8-N9-C4	-8.93	102.23	105.80
36	1	1314	C	C5-C6-N1	8.92	125.46	121.00
36	5	2322	C	C6-N1-C2	-8.92	116.73	120.30
36	5	1849	C	N1-C2-O2	-8.92	113.55	118.90
36	5	2973	G	O5'-P-OP1	-8.90	97.69	105.70
36	5	907	G	N9-C4-C5	-8.89	101.84	105.40
36	1	608	A	N1-C6-N6	8.87	123.92	118.60
36	5	3154	C	N1-C2-O2	8.87	124.22	118.90
36	1	2995	A	O5'-P-OP2	-8.86	97.73	105.70
36	1	2885	C	C6-N1-C2	8.85	123.84	120.30
36	5	942	U	N3-C4-O4	8.85	125.60	119.40
36	5	1897	G	C5-C6-N1	-8.84	107.08	111.50
36	5	2327	U	C6-N1-C2	8.84	126.30	121.00
36	5	398	A	O5'-P-OP2	-8.84	97.75	105.70
36	1	1843	C	C6-N1-C2	-8.83	116.77	120.30
36	1	1001	G	N1-C6-O6	8.83	125.20	119.90
36	1	609	G	C5-C6-O6	-8.82	123.31	128.60
36	1	1323	G	N9-C4-C5	-8.82	101.87	105.40
36	5	2637	A	O5'-P-OP1	-8.82	97.76	105.70
36	5	3177	G	C2-N3-C4	-8.82	107.49	111.90
36	1	1124	U	OP1-P-O3'	8.82	124.61	105.20
38	4	94	C	C6-N1-C2	8.81	123.83	120.30
1	6	1744	A	N1-C6-N6	8.81	123.89	118.60
36	1	1323	G	N1-C6-O6	8.79	125.18	119.90
1	2	426	G	C8-N9-C1'	-8.79	115.58	127.00
36	5	2285	C	C6-N1-C2	-8.78	116.79	120.30
1	6	390	G	N1-C6-O6	8.78	125.17	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	1301	A	N1-C6-N6	8.76	123.86	118.60
36	1	933	A	N1-C6-N6	8.76	123.85	118.60
36	1	2400	G	N1-C6-O6	8.75	125.15	119.90
1	6	338	C	C6-N1-C2	-8.72	116.81	120.30
1	6	1614	A	C2-N3-C4	-8.72	106.24	110.60
36	1	2726	C	N3-C2-O2	-8.72	115.80	121.90
36	5	2936	A	C2-N3-C4	8.70	114.95	110.60
36	1	2741	C	N1-C2-O2	8.68	124.11	118.90
36	5	803	C	C6-N1-C2	-8.68	116.83	120.30
36	1	66	A	O5'-P-OP1	-8.67	97.90	105.70
36	1	1848	G	N1-C6-O6	-8.66	114.70	119.90
36	1	3154	C	C6-N1-C2	-8.66	116.83	120.30
36	1	2417	U	N1-C2-O2	-8.66	116.74	122.80
36	1	1192	C	N3-C4-C5	-8.65	118.44	121.90
36	5	414	U	O5'-P-OP2	-8.65	97.91	105.70
36	1	3217	C	N3-C2-O2	-8.65	115.85	121.90
36	5	1149	G	C5-C6-N1	-8.65	107.18	111.50
36	1	1433	A	N9-C4-C5	8.64	109.25	105.80
36	1	711	A	C8-N9-C4	8.61	109.25	105.80
1	2	971	A	C4-C5-C6	8.60	121.30	117.00
36	1	2726	C	C5-C4-N4	8.60	126.22	120.20
1	2	830	U	N3-C2-O2	-8.59	116.18	122.20
36	1	2818	U	O5'-P-OP2	-8.59	97.97	105.70
36	5	82	C	N3-C4-C5	-8.58	118.47	121.90
36	1	3022	G	N1-C6-O6	8.58	125.05	119.90
36	5	1367	G	C5-C6-N1	-8.56	107.22	111.50
1	2	453	U	C2-N1-C1'	8.55	127.96	117.70
36	5	1306	G	N1-C6-O6	8.54	125.02	119.90
36	1	809	G	C5-C6-O6	-8.53	123.48	128.60
36	5	2327	U	C5-C6-N1	-8.53	118.44	122.70
1	6	553	G	N1-C6-O6	8.53	125.02	119.90
36	5	3108	G	C5-C6-N1	-8.53	107.24	111.50
1	2	426	G	C4-N9-C1'	8.52	137.58	126.50
36	1	635	G	C5-C6-N1	8.52	115.76	111.50
37	7	5	G	C8-N9-C4	8.52	109.81	106.40
1	2	1212	G	N1-C6-O6	8.52	125.01	119.90
36	5	607	A	N1-C6-N6	-8.52	113.49	118.60
36	1	406	G	O4'-C1'-N9	8.51	115.01	108.20
36	5	2843	U	N3-C2-O2	-8.51	116.25	122.20
36	5	3002	C	C6-N1-C2	8.51	123.70	120.30
36	5	2866	U	N3-C2-O2	-8.50	116.25	122.20
36	1	860	G	N1-C6-O6	8.49	125.00	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	1179	A	O5'-P-OP1	-8.49	98.06	105.70
36	5	800	G	C5-C6-O6	-8.48	123.51	128.60
36	5	2421	U	N1-C2-N3	8.48	119.99	114.90
36	5	2726	C	N3-C2-O2	-8.47	115.97	121.90
36	5	2726	C	C5-C4-N4	8.47	126.13	120.20
36	1	645	A	C5-C6-N6	8.47	130.48	123.70
36	5	515	C	C6-N1-C2	8.47	123.69	120.30
36	1	2811	A	C6-N1-C2	-8.46	113.52	118.60
36	5	655	C	C6-N1-C2	-8.45	116.92	120.30
36	1	718	G	N7-C8-N9	8.45	117.33	113.10
36	1	1049	C	O5'-P-OP1	-8.45	98.10	105.70
36	1	1437	C	C6-N1-C2	-8.45	116.92	120.30
36	1	979	U	C6-N1-C2	-8.44	115.94	121.00
36	1	780	A	O4'-C1'-N9	-8.43	101.45	108.20
52	M6	78	ARG	NE-CZ-NH1	8.43	124.52	120.30
36	1	3344	A	C6-C5-N7	-8.43	126.40	132.30
36	5	2290	C	C6-N1-C2	8.43	123.67	120.30
1	2	1761	U	C5-C4-O4	8.43	130.96	125.90
36	5	2383	C	C6-N1-C2	-8.43	116.93	120.30
36	1	2875	U	N3-C2-O2	-8.43	116.30	122.20
36	5	226	C	C6-N1-C2	-8.43	116.93	120.30
36	1	953	G	C4-C5-N7	8.42	114.17	110.80
1	2	73	U	O4'-C1'-N1	8.41	114.93	108.20
36	1	1104	G	O5'-P-OP1	-8.40	98.14	105.70
1	6	95	G	N1-C6-O6	-8.40	114.86	119.90
36	5	1319	G	C8-N9-C4	8.40	109.76	106.40
36	1	1792	C	C6-N1-C2	-8.40	116.94	120.30
36	1	1450	G	N1-C6-O6	8.38	124.93	119.90
36	5	1200	A	C5-C6-N6	-8.38	117.00	123.70
36	5	672	A	N9-C4-C5	8.36	109.14	105.80
36	5	1437	C	C2-N1-C1'	8.37	128.00	118.80
36	5	2341	A	C8-N9-C4	8.36	109.14	105.80
36	5	2827	U	N1-C2-O2	8.36	128.65	122.80
36	5	345	G	N1-C6-O6	8.35	124.91	119.90
1	2	380	U	N3-C2-O2	-8.35	116.35	122.20
36	1	2374	C	C2-N1-C1'	8.35	127.98	118.80
36	5	817	A	O5'-P-OP1	-8.34	98.19	105.70
36	1	648	C	C2-N1-C1'	8.34	127.97	118.80
36	5	2142	A	N1-C6-N6	-8.34	113.60	118.60
38	8	17	A	N1-C6-N6	8.33	123.60	118.60
36	1	959	C	C5-C6-N1	-8.33	116.84	121.00
36	5	2295	A	C5-C6-N6	-8.33	117.04	123.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
39	12	246	LEU	CA-CB-CG	8.33	134.46	115.30
1	2	577	G	C4-C5-N7	8.32	114.13	110.80
36	1	287	G	C5-C6-N1	-8.32	107.34	111.50
36	5	2398	A	N1-C6-N6	-8.32	113.61	118.60
36	1	2881	C	N3-C4-C5	8.32	125.23	121.90
36	1	2959	C	N1-C2-O2	-8.31	113.91	118.90
36	5	3172	A	N1-C6-N6	8.30	123.58	118.60
36	1	1351	U	N3-C2-O2	-8.30	116.39	122.20
36	5	888	A	N1-C6-N6	8.29	123.58	118.60
1	6	583	C	C6-N1-C2	-8.29	116.98	120.30
36	5	48	A	C5-C6-N1	8.29	121.84	117.70
36	5	1437	C	N3-C4-C5	-8.29	118.59	121.90
36	5	2157	G	C8-N9-C4	8.28	109.71	106.40
36	1	1192	C	N1-C2-O2	8.28	123.87	118.90
36	5	2777	G	N3-C4-N9	-8.28	121.03	126.00
36	5	718	G	C4-N9-C1'	8.27	137.26	126.50
36	1	612	U	C5-C4-O4	8.27	130.86	125.90
36	5	1012	G	C4-N9-C1'	-8.27	115.75	126.50
36	1	2679	A	C2-N3-C4	-8.27	106.47	110.60
36	5	3141	A	O5'-P-OP1	-8.27	98.26	105.70
36	1	609	G	O5'-P-OP2	-8.27	98.26	105.70
36	5	1404	G	C8-N9-C4	8.27	109.71	106.40
36	1	2797	C	N3-C4-C5	-8.25	118.60	121.90
36	1	2714	G	C2-N3-C4	-8.24	107.78	111.90
1	2	1039	A	O4'-C1'-N9	8.24	114.79	108.20
36	5	2364	G	N1-C6-O6	-8.23	114.96	119.90
36	1	282	G	C8-N9-C4	-8.23	103.11	106.40
36	1	2606	G	C4-N9-C1'	8.23	137.19	126.50
44	17	229	PHE	CB-CG-CD1	8.23	126.56	120.80
36	1	690	A	N9-C4-C5	8.22	109.09	105.80
36	5	410	U	N3-C4-C5	-8.22	109.67	114.60
36	1	3022	G	C4-C5-N7	8.22	114.09	110.80
1	6	1537	C	N3-C4-C5	-8.21	118.62	121.90
1	2	1212	G	C5-C6-O6	-8.20	123.68	128.60
36	1	1323	G	C6-C5-N7	-8.20	125.48	130.40
36	1	28	C	C6-N1-C2	8.20	123.58	120.30
1	2	190	C	O4'-C1'-N1	8.19	114.75	108.20
36	1	2610	G	N1-C6-O6	8.19	124.81	119.90
36	5	3154	C	C2-N1-C1'	8.19	127.81	118.80
36	5	1607	U	P-O3'-C3'	8.19	129.53	119.70
36	1	1508	C	C6-N1-C2	-8.19	117.03	120.30
36	1	229	G	N3-C2-N2	-8.18	114.17	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2996	U	N1-C2-O2	8.18	128.53	122.80
36	1	1389	G	C4-C5-N7	8.18	114.07	110.80
36	1	2175	U	N1-C2-N3	8.18	119.81	114.90
36	5	942	U	N3-C4-C5	-8.17	109.70	114.60
36	5	2420	C	N1-C2-O2	-8.17	114.00	118.90
36	1	3143	C	C6-N1-C2	8.16	123.56	120.30
36	5	3181	C	C6-N1-C2	-8.16	117.03	120.30
36	1	3181	C	C6-N1-C2	-8.16	117.03	120.30
36	5	907	G	C4-C5-N7	8.16	114.06	110.80
36	5	2635	A	O5'-P-OP2	-8.15	98.37	105.70
36	1	2811	A	N1-C6-N6	-8.14	113.72	118.60
36	1	699	A	C2-N3-C4	-8.14	106.53	110.60
37	7	105	C	N3-C4-C5	-8.14	118.65	121.90
1	2	577	G	C5-C6-O6	-8.13	123.72	128.60
36	1	1414	G	C5-C6-O6	-8.13	123.72	128.60
36	5	1604	G	N3-C4-N9	8.12	130.88	126.00
36	5	1178	G	C8-N9-C4	-8.12	103.15	106.40
36	5	1152	G	N3-C2-N2	-8.12	114.22	119.90
36	5	1150	A	O5'-P-OP2	-8.11	98.40	105.70
36	5	1149	G	C4-C5-C6	8.11	123.67	118.80
36	1	2811	A	C5-C6-N1	8.10	121.75	117.70
36	1	1595	U	C5-C6-N1	-8.10	118.65	122.70
36	1	635	G	C5-C6-O6	-8.09	123.74	128.60
36	5	2736	A	N1-C6-N6	8.09	123.45	118.60
36	5	1452	A	N1-C6-N6	8.09	123.45	118.60
36	1	1113	G	C8-N9-C4	-8.08	103.17	106.40
36	1	3278	C	N1-C2-O2	8.08	123.75	118.90
36	5	226	C	N3-C4-C5	-8.08	118.67	121.90
36	5	1152	G	O5'-P-OP1	-8.08	98.42	105.70
36	1	2279	A	N1-C6-N6	8.08	123.45	118.60
1	2	1761	U	C6-N1-C2	-8.07	116.16	121.00
36	5	2885	C	N1-C2-O2	-8.07	114.06	118.90
36	5	1184	A	N1-C6-N6	-8.07	113.76	118.60
36	5	670	C	N3-C4-C5	-8.06	118.68	121.90
36	1	1207	G	C5-C6-O6	-8.05	123.77	128.60
36	5	2864	A	N1-C6-N6	8.05	123.43	118.60
36	1	2819	A	C8-N9-C4	-8.04	102.59	105.80
36	5	1142	G	N3-C4-C5	-8.04	124.58	128.60
36	5	694	C	C6-N1-C2	-8.03	117.09	120.30
36	5	3029	A	N1-C6-N6	-8.03	113.78	118.60
36	1	637	C	O5'-P-OP1	-8.03	98.47	105.70
1	6	337	G	C2-N3-C4	8.03	115.92	111.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	1901	A	N1-C6-N6	-8.03	113.78	118.60
36	5	2777	G	C4-C5-N7	-8.02	107.59	110.80
36	1	355	A	C2-N3-C4	-8.02	106.59	110.60
36	5	2819	A	O5'-P-OP2	-8.02	98.49	105.70
36	5	2842	U	O5'-P-OP1	-8.02	98.49	105.70
1	2	1636	C	C6-N1-C2	-8.01	117.10	120.30
37	7	80	G	C6-C5-N7	-8.01	125.59	130.40
36	1	2380	U	N1-C2-N3	8.00	119.70	114.90
36	5	2777	G	N9-C4-C5	8.00	108.60	105.40
36	5	1192	C	C2-N1-C1'	8.00	127.60	118.80
36	5	3245	A	C5-C6-N1	-8.00	113.70	117.70
36	5	1008	U	C5-C6-N1	-7.99	118.70	122.70
36	5	2880	U	C5-C6-N1	7.99	126.70	122.70
36	1	1113	G	N7-C8-N9	7.98	117.09	113.10
36	5	798	G	C8-N9-C4	-7.98	103.21	106.40
36	5	1377	G	N3-C4-C5	-7.97	124.61	128.60
36	5	2113	A	C8-N9-C4	7.97	108.99	105.80
36	1	2636	A	C8-N9-C4	-7.97	102.61	105.80
36	1	218	G	C5-C6-O6	-7.97	123.82	128.60
1	6	630	A	C8-N9-C4	7.96	108.99	105.80
36	5	826	G	N1-C6-O6	7.96	124.68	119.90
36	1	2379	U	N1-C2-O2	-7.96	117.23	122.80
36	5	567	G	C4-C5-N7	7.96	113.98	110.80
36	5	1321	G	N1-C6-O6	7.95	124.67	119.90
36	1	1166	G	C6-C5-N7	-7.95	125.63	130.40
36	5	874	U	C2-N1-C1'	-7.95	108.16	117.70
1	2	93	A	C8-N9-C4	-7.95	102.62	105.80
36	1	676	G	C8-N9-C1'	-7.94	116.68	127.00
36	5	923	C	N3-C4-C5	7.94	125.08	121.90
36	1	1124	U	N3-C2-O2	-7.93	116.64	122.20
36	5	1161	G	O5'-P-OP1	-7.93	98.56	105.70
36	5	2552	C	N1-C2-O2	7.93	123.66	118.90
36	1	914	A	N1-C6-N6	-7.93	113.84	118.60
36	5	2295	A	C4-C5-N7	7.93	114.67	110.70
36	5	2707	C	N3-C4-C5	7.93	125.07	121.90
36	1	595	G	O5'-P-OP1	-7.92	98.57	105.70
36	5	1483	G	O4'-C1'-N9	7.92	114.54	108.20
36	1	2836	C	C4-C5-C6	7.92	121.36	117.40
36	5	2811	A	O5'-P-OP1	-7.91	98.58	105.70
36	1	2795	U	O5'-P-OP1	-7.91	98.58	105.70
36	5	2199	G	C8-N9-C4	-7.91	103.24	106.40
36	5	2959	C	N3-C4-C5	-7.91	118.74	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2960	C	C6-N1-C2	-7.90	117.14	120.30
1	6	163	G	C2-N3-C4	-7.90	107.95	111.90
36	1	2916	U	N1-C2-N3	-7.89	110.16	114.90
36	5	2145	A	C8-N9-C4	-7.89	102.64	105.80
36	1	326	U	O5'-P-OP2	-7.89	98.60	105.70
36	5	800	G	C4-C5-N7	7.88	113.95	110.80
36	1	2374	C	N1-C2-O2	7.88	123.63	118.90
36	1	94	G	O5'-P-OP1	-7.88	98.61	105.70
36	1	2726	C	C6-N1-C2	-7.87	117.15	120.30
36	5	1371	G	C8-N9-C4	7.87	109.55	106.40
36	5	1900	A	N1-C6-N6	7.87	123.32	118.60
36	5	2383	C	C5-C6-N1	7.87	124.93	121.00
36	1	2409	G	N3-C4-C5	-7.86	124.67	128.60
36	1	24	G	C5-C6-N1	-7.85	107.57	111.50
36	5	776	U	N3-C2-O2	-7.85	116.70	122.20
36	1	991	G	C8-N9-C4	-7.85	103.26	106.40
36	5	1443	G	N1-C6-O6	7.85	124.61	119.90
36	5	661	G	C8-N9-C4	-7.84	103.27	106.40
36	1	2852	C	C6-N1-C2	7.83	123.43	120.30
36	5	1301	A	C6-C5-N7	-7.83	126.82	132.30
36	5	522	A	N1-C6-N6	7.82	123.29	118.60
36	5	364	G	N1-C6-O6	7.82	124.59	119.90
36	1	2899	C	N3-C2-O2	-7.82	116.43	121.90
36	5	887	G	C5-C6-O6	-7.81	123.91	128.60
1	6	782	U	N3-C2-O2	-7.81	116.73	122.20
36	5	924	G	C5-C6-N1	-7.81	107.59	111.50
36	1	2356	A	N1-C6-N6	7.81	123.28	118.60
36	1	3217	C	C6-N1-C2	-7.80	117.18	120.30
36	1	2617	U	C5-C4-O4	7.80	130.58	125.90
1	2	639	U	N1-C2-O2	7.79	128.26	122.80
36	5	1592	G	C8-N9-C4	-7.79	103.28	106.40
1	6	798	C	C6-N1-C2	7.79	123.41	120.30
36	1	718	G	C8-N9-C4	-7.78	103.29	106.40
36	5	2373	A	O5'-P-OP1	-7.78	98.70	105.70
36	1	3212	C	C6-N1-C2	7.77	123.41	120.30
36	5	2403	G	C8-N9-C4	7.76	109.50	106.40
36	5	668	G	N7-C8-N9	-7.76	109.22	113.10
36	5	2392	C	N3-C4-C5	7.76	125.00	121.90
36	1	933	A	C5-N7-C8	-7.76	100.02	103.90
36	1	2281	A	O5'-P-OP2	-7.76	98.72	105.70
36	5	2199	G	N7-C8-N9	7.76	116.98	113.10
36	1	2662	G	N1-C6-O6	7.75	124.55	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	1187	C	N3-C4-C5	7.75	125.00	121.90
1	6	1150	G	N3-C4-C5	7.75	132.48	128.60
36	5	960	U	N3-C2-O2	-7.75	116.77	122.20
36	1	689	U	N1-C2-O2	7.75	128.22	122.80
38	8	25	G	N3-C4-C5	-7.75	124.73	128.60
36	5	980	A	N1-C6-N6	-7.74	113.95	118.60
36	5	2994	A	C6-N1-C2	-7.74	113.95	118.60
36	1	590	G	C5-C6-O6	-7.74	123.95	128.60
36	1	2194	G	N1-C6-O6	7.74	124.54	119.90
36	5	2295	A	C5-N7-C8	-7.74	100.03	103.90
36	5	2797	C	N3-C4-C5	-7.73	118.81	121.90
36	1	948	C	C5-C6-N1	-7.73	117.13	121.00
36	5	388	G	C8-N9-C4	-7.72	103.31	106.40
36	5	1148	G	C5-C6-O6	-7.72	123.97	128.60
36	1	145	G	C4-C5-N7	7.72	113.89	110.80
36	1	2730	G	N3-C4-N9	-7.72	121.37	126.00
36	1	1351	U	N1-C2-O2	7.72	128.20	122.80
1	2	830	U	N1-C2-O2	7.71	128.20	122.80
36	5	1127	G	O5'-P-OP2	-7.71	98.77	105.70
36	1	689	U	N3-C2-O2	-7.70	116.81	122.20
36	1	2617	U	N3-C2-O2	-7.70	116.81	122.20
36	1	644	G	C5-C6-O6	7.70	133.22	128.60
36	1	957	C	N1-C2-O2	-7.70	114.28	118.90
36	1	637	C	N3-C2-O2	-7.70	116.51	121.90
36	5	2906	C	C6-N1-C2	7.69	123.38	120.30
36	1	398	A	C8-N9-C4	7.69	108.88	105.80
36	1	676	G	C6-C5-N7	-7.69	125.79	130.40
36	5	3149	G	C5-C6-N1	-7.69	107.66	111.50
1	2	1340	U	C5-C4-O4	7.69	130.51	125.90
1	2	1636	C	N3-C4-C5	-7.69	118.83	121.90
36	1	1001	G	N3-C4-N9	7.69	130.61	126.00
36	5	82	C	C4-C5-C6	7.68	121.24	117.40
1	6	337	G	N3-C4-N9	7.68	130.61	126.00
36	1	2968	G	C6-C5-N7	-7.68	125.79	130.40
1	2	1539	G	C4-N9-C1'	7.68	136.48	126.50
36	1	1113	G	C5-N7-C8	-7.67	100.46	104.30
37	7	1	G	C4-N9-C1'	7.67	136.47	126.50
36	5	2733	A	O5'-P-OP2	-7.67	98.80	105.70
36	1	3054	U	N3-C4-C5	-7.66	110.00	114.60
36	5	2869	U	C5-C4-O4	7.66	130.50	125.90
36	5	3078	U	N3-C2-O2	-7.66	116.84	122.20
36	1	1414	G	N1-C6-O6	7.65	124.49	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	304	G	C5-C6-O6	7.65	133.19	128.60
36	1	2697	A	N1-C6-N6	-7.65	114.01	118.60
38	4	26	U	N3-C2-O2	-7.65	116.85	122.20
36	5	2271	A	N1-C6-N6	-7.65	114.01	118.60
36	1	1904	C	N3-C4-C5	-7.65	118.84	121.90
36	1	3201	C	C6-N1-C2	-7.64	117.24	120.30
38	4	52	A	N1-C6-N6	-7.64	114.02	118.60
1	2	1200	G	N1-C6-O6	7.64	124.48	119.90
36	1	287	G	N1-C6-O6	7.64	124.48	119.90
36	5	1152	G	C4-N9-C1'	-7.64	116.57	126.50
36	1	808	A	N1-C6-N6	7.63	123.18	118.60
36	1	232	G	N3-C4-C5	-7.63	124.78	128.60
36	1	2948	C	C6-N1-C2	7.63	123.35	120.30
36	1	1607	U	P-O3'-C3'	7.63	128.85	119.70
36	1	1175	C	C6-N1-C2	7.62	123.35	120.30
36	1	3344	A	C5-N7-C8	-7.62	100.09	103.90
36	5	2865	U	C5-C6-N1	7.62	126.51	122.70
36	1	1888	U	N3-C2-O2	-7.62	116.87	122.20
44	17	232	ARG	NE-CZ-NH1	-7.62	116.49	120.30
36	1	1149	G	C4-C5-C6	7.61	123.37	118.80
36	1	1307	G	N3-C4-N9	-7.61	121.44	126.00
36	1	2912	G	O5'-P-OP1	-7.61	98.86	105.70
1	6	621	A	O5'-P-OP1	-7.61	98.85	105.70
36	5	2629	U	C5-C4-O4	-7.61	121.34	125.90
36	5	2817	A	C2-N3-C4	7.61	114.40	110.60
36	1	638	C	O5'-P-OP2	-7.60	98.86	105.70
36	5	1171	G	O5'-P-OP2	-7.60	98.86	105.70
1	6	390	G	C6-C5-N7	-7.60	125.84	130.40
36	5	2364	G	C5-C6-O6	7.60	133.16	128.60
36	1	1367	G	N9-C4-C5	-7.60	102.36	105.40
36	1	963	G	O5'-P-OP1	7.59	119.81	110.70
36	1	1495	U	N1-C2-N3	7.59	119.46	114.90
1	2	17	C	C6-N1-C2	-7.59	117.27	120.30
36	1	645	A	C4-C5-N7	-7.59	106.91	110.70
36	5	2942	C	C5-C6-N1	7.59	124.79	121.00
36	5	2353	G	N1-C6-O6	7.58	124.45	119.90
36	1	2418	G	OP1-P-O3'	7.58	121.88	105.20
36	1	397	A	N7-C8-N9	-7.58	110.01	113.80
36	1	1386	A	N1-C6-N6	7.58	123.15	118.60
36	1	1389	G	C5-C6-O6	-7.58	124.05	128.60
36	5	2413	A	O5'-P-OP1	-7.58	98.88	105.70
36	5	2761	G	C5-C6-N1	7.58	115.29	111.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2190	U	N3-C4-C5	-7.57	110.06	114.60
36	1	937	G	N1-C6-O6	7.57	124.44	119.90
36	1	2549	G	N3-C4-N9	7.57	130.54	126.00
36	1	3344	A	O4'-C1'-N9	7.57	114.26	108.20
1	6	1744	A	C4-C5-N7	7.57	114.48	110.70
36	5	519	A	C5-C6-N6	-7.57	117.64	123.70
36	5	2421	U	N1-C2-O2	-7.57	117.50	122.80
36	5	1127	G	N1-C6-O6	-7.57	115.36	119.90
36	5	672	A	C8-N9-C4	-7.57	102.77	105.80
36	5	1306	G	C5-N7-C8	-7.56	100.52	104.30
36	5	514	G	N1-C6-O6	7.56	124.44	119.90
36	5	1190	A	C8-N9-C4	-7.56	102.78	105.80
36	5	437	G	C8-N9-C4	-7.56	103.38	106.40
36	1	1004	U	C6-N1-C2	-7.56	116.47	121.00
36	1	3382	U	C6-N1-C1'	-7.56	110.62	121.20
36	5	2375	G	O4'-C1'-N9	7.56	114.25	108.20
36	1	287	G	O5'-P-OP1	-7.55	98.90	105.70
36	1	2117	A	O5'-P-OP1	-7.55	98.91	105.70
36	1	2827	U	C5-C4-O4	7.55	130.43	125.90
36	1	358	G	C4-C5-N7	7.55	113.82	110.80
36	1	2870	C	C2-N1-C1'	-7.55	110.50	118.80
36	5	2616	C	O5'-P-OP1	-7.54	98.91	105.70
36	1	880	G	O4'-C1'-N9	7.54	114.23	108.20
36	1	611	A	O5'-P-OP2	-7.54	98.92	105.70
36	1	913	A	C8-N9-C4	-7.54	102.79	105.80
1	6	779	U	N1-C2-O2	7.54	128.07	122.80
36	1	1192	C	N3-C2-O2	-7.53	116.63	121.90
36	1	690	A	C5-C6-N6	7.53	129.72	123.70
36	1	1888	U	N1-C2-N3	7.53	119.42	114.90
36	1	2358	A	N1-C6-N6	7.53	123.12	118.60
1	2	779	U	O4'-C1'-N1	7.53	114.22	108.20
36	1	715	A	O5'-P-OP2	-7.52	98.93	105.70
36	5	364	G	C4-C5-N7	7.52	113.81	110.80
36	1	1435	A	C8-N9-C4	-7.52	102.79	105.80
36	1	283	G	O4'-C1'-N9	-7.52	102.19	108.20
36	5	806	A	C8-N9-C4	7.52	108.81	105.80
36	5	1012	G	N3-C4-N9	-7.52	121.49	126.00
36	5	1198	C	C2-N1-C1'	-7.51	110.53	118.80
36	5	800	G	N1-C6-O6	7.51	124.41	119.90
36	1	1329	U	C2-N1-C1'	7.50	126.70	117.70
1	2	971	A	C8-N9-C4	-7.50	102.80	105.80
36	1	3022	G	N9-C4-C5	-7.50	102.40	105.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	1321	G	C6-C5-N7	-7.49	125.91	130.40
36	1	1115	G	C8-N9-C4	-7.49	103.40	106.40
36	1	2296	A	C8-N9-C4	7.49	108.80	105.80
36	5	1160	C	C2-N1-C1'	-7.49	110.56	118.80
36	1	3110	C	C6-N1-C2	-7.49	117.30	120.30
36	5	2866	U	N1-C2-O2	7.49	128.04	122.80
36	1	2643	A	N1-C6-N6	7.49	123.09	118.60
36	5	2186	U	O5'-P-OP2	-7.49	98.96	105.70
36	1	727	G	N3-C4-C5	-7.48	124.86	128.60
1	6	337	G	C4-N9-C1'	7.48	136.23	126.50
36	5	2777	G	C5-C6-O6	7.48	133.09	128.60
78	q2	93	LEU	CA-CB-CG	7.48	132.50	115.30
36	1	2309	A	N1-C6-N6	7.48	123.08	118.60
1	6	1600	A	O4'-C1'-N9	7.48	114.18	108.20
36	5	1917	C	C6-N1-C2	7.47	123.29	120.30
36	1	1389	G	N3-C4-N9	7.47	130.48	126.00
38	4	81	U	N3-C2-O2	-7.47	116.97	122.20
36	1	940	G	O5'-P-OP1	-7.47	98.98	105.70
1	2	1489	U	N3-C2-O2	-7.47	116.97	122.20
36	1	3078	U	N3-C2-O2	-7.47	116.97	122.20
1	6	543	C	C6-N1-C2	-7.47	117.31	120.30
36	5	877	C	N3-C4-C5	7.47	124.89	121.90
36	5	2890	A	O5'-P-OP1	-7.46	98.98	105.70
36	1	2344	U	C5-C6-N1	-7.46	118.97	122.70
36	1	709	A	C5-C6-N6	-7.46	117.74	123.70
36	1	2619	G	O5'-P-OP1	-7.46	98.99	105.70
36	1	1838	G	C5-C6-O6	-7.45	124.13	128.60
36	1	2298	U	C5-C4-O4	7.45	130.37	125.90
1	6	542	A	C8-N9-C4	-7.45	102.82	105.80
37	7	55	A	O5'-P-OP1	-7.45	99.00	105.70
36	5	1152	G	C8-N9-C4	-7.44	103.42	106.40
36	1	49	A	C8-N9-C4	7.44	108.78	105.80
36	1	14	U	O5'-P-OP2	-7.44	99.00	105.70
36	5	3245	A	C5-N7-C8	-7.44	100.18	103.90
36	1	2889	C	N1-C2-O2	7.44	123.36	118.90
10	S8	29	LEU	CA-CB-CG	7.44	132.41	115.30
36	5	670	C	C6-N1-C2	-7.44	117.33	120.30
36	5	1160	C	N1-C2-O2	-7.44	114.44	118.90
36	5	1012	G	N3-C4-C5	7.43	132.32	128.60
36	1	1450	G	C5-C6-O6	-7.43	124.14	128.60
36	1	2585	G	N3-C4-C5	-7.43	124.89	128.60
36	5	587	U	C6-N1-C2	7.43	125.46	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	1156	C	C6-N1-C2	-7.43	117.33	120.30
36	1	714	G	C6-C5-N7	-7.42	125.95	130.40
36	1	2247	G	C6-C5-N7	-7.42	125.95	130.40
36	5	2189	U	O5'-P-OP1	-7.42	99.02	105.70
36	1	2376	G	C5-N7-C8	-7.42	100.59	104.30
36	5	2400	G	C8-N9-C4	7.42	109.37	106.40
36	1	1151	U	N3-C4-O4	7.42	124.59	119.40
1	6	453	U	C2-N1-C1'	7.42	126.60	117.70
1	6	435	C	N1-C2-O2	7.41	123.35	118.90
36	5	1126	G	C8-N9-C4	-7.41	103.44	106.40
36	5	2197	C	N1-C2-N3	-7.41	114.01	119.20
36	5	1849	C	N3-C2-O2	7.41	127.09	121.90
36	1	1484	U	P-O3'-C3'	7.41	128.59	119.70
51	m5	24	ARG	NE-CZ-NH1	7.41	124.00	120.30
1	6	524	U	O5'-P-OP2	-7.40	99.04	105.70
1	2	736	C	C5-C6-N1	7.40	124.70	121.00
36	1	1595	U	C2-N1-C1'	-7.40	108.82	117.70
36	5	1317	A	N1-C2-N3	-7.40	125.60	129.30
36	1	1166	G	N9-C4-C5	-7.40	102.44	105.40
36	1	2279	A	C5-C6-N6	-7.39	117.78	123.70
36	5	2550	U	N3-C2-O2	-7.39	117.03	122.20
1	2	1280	C	C6-N1-C2	-7.39	117.34	120.30
36	1	318	A	O5'-P-OP1	-7.39	99.05	105.70
36	5	2345	A	N1-C6-N6	7.39	123.03	118.60
36	1	1522	U	C2-N3-C4	-7.38	122.57	127.00
36	5	2981	U	N3-C2-O2	-7.38	117.03	122.20
37	7	40	C	C6-N1-C2	7.38	123.25	120.30
36	5	776	U	C5-C6-N1	-7.38	119.01	122.70
37	7	93	C	C6-N1-C2	-7.38	117.35	120.30
36	1	1556	C	N1-C2-O2	7.37	123.32	118.90
36	1	2179	C	N1-C2-O2	7.37	123.32	118.90
36	1	1407	A	O5'-P-OP2	-7.37	99.06	105.70
36	5	2199	G	C5-N7-C8	-7.37	100.61	104.30
36	1	2403	G	N1-C6-O6	7.37	124.32	119.90
36	5	2281	A	N9-C4-C5	-7.37	102.85	105.80
36	1	1151	U	N3-C4-C5	-7.36	110.18	114.60
36	1	397	A	C8-N9-C4	7.36	108.74	105.80
36	5	715	A	C5-C6-N6	7.35	129.58	123.70
36	5	1212	A	C5-C6-N6	-7.35	117.82	123.70
36	1	974	G	C5-C6-N1	7.35	115.18	111.50
36	5	3197	G	N3-C2-N2	-7.35	114.75	119.90
36	5	3309	G	C4-N9-C1'	7.35	136.05	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	901	G	C5-C6-N1	-7.34	107.83	111.50
36	1	2606	G	C8-N9-C1'	-7.34	117.45	127.00
36	1	2822	U	O5'-P-OP1	-7.34	99.09	105.70
36	5	1321	G	C4-C5-N7	7.34	113.74	110.80
36	5	2699	G	N1-C6-O6	7.34	124.31	119.90
36	1	1201	C	C5-C6-N1	7.34	124.67	121.00
36	1	388	G	C8-N9-C4	-7.34	103.46	106.40
36	1	870	G	N9-C4-C5	7.34	108.34	105.40
36	5	98	G	N3-C4-N9	7.34	130.40	126.00
36	1	432	G	C5-C6-N1	-7.33	107.83	111.50
36	5	1868	G	C6-C5-N7	-7.33	126.00	130.40
38	4	99	C	N1-C2-O2	7.33	123.30	118.90
36	5	718	G	C6-C5-N7	-7.32	126.01	130.40
36	5	339	C	N3-C2-O2	7.32	127.02	121.90
1	2	1082	C	C6-N1-C2	-7.32	117.37	120.30
1	2	1082	C	C6-N1-C1'	-7.32	112.02	120.80
36	5	2899	C	C6-N1-C2	-7.32	117.37	120.30
36	5	2976	A	N9-C4-C5	7.32	108.73	105.80
36	1	3362	A	O4'-C1'-N9	7.32	114.05	108.20
36	5	197	G	C4-C5-N7	7.32	113.73	110.80
36	5	2351	U	N3-C2-O2	-7.32	117.08	122.20
37	7	51	A	C6-C5-N7	-7.32	127.18	132.30
36	5	1449	A	N9-C4-C5	-7.31	102.88	105.80
36	5	2610	G	C8-N9-C4	-7.31	103.47	106.40
1	2	1761	U	N3-C2-O2	-7.31	117.08	122.20
36	1	631	U	N3-C2-O2	7.31	127.32	122.20
38	4	83	C	N1-C2-O2	7.31	123.29	118.90
36	5	1848	G	C8-N9-C4	7.31	109.32	106.40
36	5	646	A	C2-N3-C4	-7.31	106.95	110.60
36	1	1303	A	C5-C6-N6	-7.30	117.86	123.70
36	1	1403	C	C6-N1-C2	7.30	123.22	120.30
36	1	711	A	N7-C8-N9	-7.30	110.15	113.80
1	2	1241	G	C6-C5-N7	-7.30	126.02	130.40
36	1	51	A	C5-N7-C8	-7.30	100.25	103.90
36	5	1347	U	C2-N1-C1'	-7.30	108.94	117.70
36	1	2989	U	N3-C4-O4	7.30	124.51	119.40
36	5	437	G	N9-C4-C5	7.30	108.32	105.40
36	1	2624	G	C5-C6-O6	-7.30	124.22	128.60
1	6	542	A	O5'-P-OP1	-7.29	99.14	105.70
36	5	2893	C	N3-C4-C5	-7.29	118.98	121.90
36	5	3130	A	N1-C2-N3	7.29	132.94	129.30
36	5	1316	C	N1-C2-O2	-7.28	114.53	118.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	1392	G	C8-N9-C4	7.28	109.31	106.40
36	5	1404	G	C2-N3-C4	-7.28	108.26	111.90
36	5	2723	U	C6-N1-C2	-7.28	116.63	121.00
36	5	961	C	O5'-P-OP1	-7.28	99.15	105.70
36	1	51	A	N1-C6-N6	7.27	122.97	118.60
1	2	970	A	N1-C6-N6	7.27	122.96	118.60
36	1	716	A	N9-C4-C5	-7.27	102.89	105.80
1	2	1731	A	N1-C6-N6	7.27	122.96	118.60
36	1	3078	U	N1-C2-O2	7.27	127.89	122.80
36	5	2936	A	N1-C6-N6	-7.26	114.24	118.60
36	5	2385	G	C2-N3-C4	-7.26	108.27	111.90
36	1	1493	G	N1-C6-O6	-7.26	115.54	119.90
36	5	2863	G	C5-C6-O6	7.26	132.96	128.60
36	1	2935	U	C5-C6-N1	7.25	126.33	122.70
36	5	2283	G	C4-C5-N7	7.25	113.70	110.80
36	5	304	G	N1-C6-O6	-7.25	115.55	119.90
36	5	2351	U	C6-N1-C2	-7.25	116.65	121.00
36	5	1404	G	N3-C4-C5	7.25	132.22	128.60
36	1	637	C	P-O3'-C3'	7.25	128.40	119.70
36	5	2391	G	C5-C6-N1	7.25	115.12	111.50
36	5	423	A	N9-C4-C5	-7.24	102.90	105.80
1	2	970	A	C5-C6-N6	-7.24	117.91	123.70
1	6	543	C	N1-C2-O2	7.24	123.24	118.90
36	1	558	U	C2-N1-C1'	7.24	126.38	117.70
36	1	803	C	C2-N1-C1'	7.24	126.76	118.80
36	5	1169	A	C8-N9-C4	7.24	108.69	105.80
36	1	2960	C	N3-C2-O2	-7.23	116.84	121.90
36	5	1006	A	O5'-P-OP2	-7.23	99.19	105.70
36	5	1468	A	N1-C6-N6	7.23	122.94	118.60
36	1	1437	C	C2-N1-C1'	7.22	126.75	118.80
1	2	1241	G	N7-C8-N9	7.22	116.71	113.10
36	1	1192	C	C2-N1-C1'	7.22	126.74	118.80
36	5	806	A	C4-C5-C6	-7.22	113.39	117.00
36	5	3278	C	C6-N1-C2	7.22	123.19	120.30
36	5	1452	A	N9-C4-C5	-7.22	102.91	105.80
36	1	692	A	C4-C5-C6	7.21	120.61	117.00
36	5	612	U	OP2-P-O3'	7.21	121.07	105.20
36	1	906	A	N1-C6-N6	7.21	122.92	118.60
36	5	360	G	C8-N9-C4	-7.21	103.52	106.40
1	2	1489	U	C2-N1-C1'	7.21	126.35	117.70
36	1	2369	G	C8-N9-C4	-7.20	103.52	106.40
36	5	2156	C	C6-N1-C2	7.20	123.18	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	3217	C	N1-C2-O2	7.20	123.22	118.90
36	1	947	G	C6-C5-N7	-7.20	126.08	130.40
36	1	2175	U	C5-C4-O4	7.20	130.22	125.90
38	4	93	U	C5-C4-O4	7.20	130.22	125.90
36	1	1733	G	N3-C4-C5	-7.19	125.00	128.60
38	4	66	A	C8-N9-C4	-7.19	102.92	105.80
36	1	3000	A	C2-N3-C4	-7.19	107.00	110.60
36	5	3131	U	N3-C4-C5	7.19	118.91	114.60
36	1	105	C	C6-N1-C2	7.19	123.17	120.30
36	5	1158	A	O5'-P-OP2	-7.19	99.23	105.70
38	8	20	U	N1-C2-O2	-7.19	117.77	122.80
36	1	690	A	C4-C5-N7	-7.18	107.11	110.70
36	1	912	G	N3-C4-N9	7.18	130.31	126.00
36	5	2419	A	C8-N9-C4	-7.18	102.93	105.80
37	7	48	U	N3-C4-O4	7.18	124.42	119.40
36	5	1114	U	OP1-P-O3'	7.17	120.98	105.20
1	6	1130	G	C8-N9-C4	7.17	109.27	106.40
36	5	2758	A	N9-C4-C5	7.17	108.67	105.80
36	5	720	A	C6-C5-N7	-7.17	127.28	132.30
36	1	2882	U	N3-C2-O2	-7.17	117.19	122.20
36	5	1010	G	O5'-P-OP2	-7.17	99.25	105.70
37	7	69	C	C6-N1-C2	7.17	123.17	120.30
36	1	586	C	N1-C2-O2	-7.16	114.60	118.90
36	5	1198	C	C6-N1-C1'	7.16	129.40	120.80
38	8	4	C	N3-C2-O2	-7.16	116.89	121.90
36	5	3091	A	O5'-P-OP1	-7.16	99.25	105.70
1	6	1744	A	N9-C4-C5	-7.16	102.94	105.80
36	5	948	C	C6-N1-C2	7.16	123.16	120.30
36	1	860	G	C5-C6-O6	-7.16	124.31	128.60
36	1	3054	U	C6-N1-C2	-7.16	116.70	121.00
36	5	1373	A	N1-C6-N6	7.15	122.89	118.60
36	1	833	G	C4-C5-N7	-7.15	107.94	110.80
36	1	2920	U	C5-C6-N1	-7.15	119.12	122.70
36	1	1831	U	C6-N1-C2	-7.15	116.71	121.00
36	5	48	A	C6-N1-C2	-7.15	114.31	118.60
36	1	2871	G	O5'-P-OP2	-7.14	99.27	105.70
36	1	1480	G	N3-C4-C5	7.14	132.17	128.60
1	2	639	U	N3-C2-O2	-7.14	117.20	122.20
36	1	586	C	N3-C2-O2	7.14	126.90	121.90
36	1	2870	C	C6-N1-C1'	7.14	129.37	120.80
1	6	979	A	C8-N9-C4	-7.14	102.94	105.80
36	5	2709	C	C5-C6-N1	-7.14	117.43	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	2	1541	G	N3-C4-C5	-7.14	125.03	128.60
36	5	1301	A	C5-C6-N6	-7.14	117.99	123.70
36	5	2816	G	C8-N9-C4	7.14	109.25	106.40
36	1	590	G	N1-C6-O6	7.13	124.18	119.90
36	5	205	C	O5'-P-OP1	-7.13	99.28	105.70
36	5	905	U	O5'-P-OP1	-7.13	99.28	105.70
36	5	2400	G	N3-C4-C5	7.13	132.17	128.60
1	2	346	G	C8-N9-C4	7.13	109.25	106.40
36	1	1373	A	O5'-P-OP2	-7.13	99.28	105.70
36	5	1364	C	OP2-P-O3'	7.13	120.89	105.20
36	5	2281	A	C8-N9-C4	7.13	108.65	105.80
36	5	715	A	O4'-C1'-N9	7.13	113.90	108.20
36	5	2662	G	N3-C4-N9	7.13	130.28	126.00
36	1	123	A	N1-C6-N6	7.13	122.88	118.60
36	5	921	A	OP2-P-O3'	7.13	120.88	105.20
36	5	1587	A	N7-C8-N9	-7.13	110.24	113.80
38	8	48	A	C5-N7-C8	-7.13	100.34	103.90
36	5	3309	G	N3-C4-C5	-7.12	125.04	128.60
36	5	887	G	N9-C4-C5	-7.12	102.55	105.40
1	6	136	C	C2-N1-C1'	7.12	126.63	118.80
36	1	684	G	N1-C6-O6	7.12	124.17	119.90
36	1	1885	U	C5-C6-N1	-7.12	119.14	122.70
1	6	55	A	C8-N9-C4	-7.12	102.95	105.80
36	5	364	G	C6-C5-N7	-7.12	126.13	130.40
1	6	1032	G	C8-N9-C4	7.12	109.25	106.40
36	5	2316	G	N3-C4-C5	-7.12	125.04	128.60
1	2	559	C	C6-N1-C2	-7.12	117.45	120.30
49	M3	85	LEU	CA-CB-CG	7.12	131.66	115.30
37	7	101	G	C4-C5-N7	7.12	113.65	110.80
1	2	553	G	C5-C6-O6	-7.11	124.33	128.60
36	5	1317	A	C2-N3-C4	7.11	114.16	110.60
36	5	3000	A	C8-N9-C4	7.11	108.64	105.80
36	1	3362	A	C6-C5-N7	-7.11	127.32	132.30
36	1	421	G	N3-C2-N2	7.11	124.88	119.90
1	6	1744	A	C5-C6-N6	-7.11	118.01	123.70
12	c0	97	PRO	N-CA-CB	7.11	111.83	103.30
1	6	1653	C	N1-C2-O2	-7.10	114.64	118.90
36	1	808	A	C5-C6-N6	-7.10	118.02	123.70
36	5	1163	A	O5'-P-OP2	-7.10	99.31	105.70
36	1	645	A	C8-N9-C4	-7.09	102.96	105.80
36	1	2827	U	N1-C2-N3	7.09	119.16	114.90
36	5	3315	G	C6-C5-N7	-7.09	126.14	130.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2874	G	C5-C6-N1	-7.09	107.96	111.50
36	1	612	U	N3-C4-O4	-7.09	114.44	119.40
36	1	1269	U	C2-N1-C1'	7.09	126.20	117.70
36	1	2329	C	N3-C4-C5	7.09	124.73	121.90
36	5	519	A	C4-C5-N7	7.09	114.24	110.70
36	5	2283	G	O5'-P-OP2	-7.09	99.32	105.70
36	1	1157	G	N1-C2-N3	7.08	128.15	123.90
37	7	101	G	C6-C5-N7	-7.08	126.15	130.40
36	1	1421	G	OP2-P-O3'	7.08	120.79	105.20
36	5	781	G	N1-C6-O6	-7.08	115.65	119.90
36	1	2298	U	N3-C4-O4	-7.08	114.44	119.40
36	5	610	G	C8-N9-C4	-7.08	103.57	106.40
37	3	93	C	N3-C4-C5	7.08	124.73	121.90
1	2	1182	U	N3-C2-O2	-7.08	117.25	122.20
36	5	1496	C	C2-N1-C1'	7.08	126.58	118.80
36	1	1328	C	C6-N1-C2	-7.08	117.47	120.30
1	6	1172	G	N3-C4-C5	-7.08	125.06	128.60
36	1	912	G	N3-C4-C5	-7.07	125.06	128.60
48	M1	112	LEU	CA-CB-CG	7.07	131.56	115.30
36	5	3092	C	C5-C6-N1	-7.07	117.46	121.00
36	5	2739	A	C8-N9-C4	7.07	108.63	105.80
1	2	1456	C	C6-N1-C2	-7.07	117.47	120.30
36	1	878	G	C5-C6-O6	7.07	132.84	128.60
36	1	3092	C	O5'-P-OP1	-7.07	99.34	105.70
1	6	158	U	P-O3'-C3'	7.07	128.18	119.70
1	2	1458	G	C6-C5-N7	-7.06	126.16	130.40
36	1	809	G	N1-C6-O6	7.06	124.14	119.90
1	6	901	G	C4-C5-N7	7.06	113.62	110.80
36	5	2724	U	C6-N1-C2	-7.06	116.76	121.00
36	1	2860	U	N1-C2-O2	7.06	127.74	122.80
36	1	3204	C	C6-N1-C2	-7.06	117.48	120.30
36	1	2287	C	N3-C4-C5	-7.06	119.08	121.90
38	8	26	U	N1-C2-O2	7.06	127.74	122.80
36	1	2400	G	C4-C5-N7	7.06	113.62	110.80
36	5	3107	U	N3-C4-O4	7.05	124.34	119.40
36	1	870	G	C8-N9-C4	-7.05	103.58	106.40
36	5	197	G	C6-C5-N7	-7.05	126.17	130.40
36	5	1592	G	N9-C4-C5	7.05	108.22	105.40
1	2	1128	C	O5'-P-OP1	-7.05	99.36	105.70
36	1	2393	G	C5-C6-O6	-7.05	124.37	128.60
36	1	953	G	N1-C6-O6	7.05	124.13	119.90
36	5	842	G	C4-C5-N7	7.05	113.62	110.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	910	G	C5-C6-N1	-7.04	107.98	111.50
36	1	3085	G	N3-C2-N2	-7.04	114.97	119.90
36	5	2359	C	C6-N1-C2	7.04	123.12	120.30
36	1	2983	C	O5'-P-OP1	-7.04	99.36	105.70
36	5	2157	G	N3-C4-C5	7.04	132.12	128.60
36	5	2346	C	N3-C4-N4	7.04	122.92	118.00
36	1	727	G	C8-N9-C4	-7.03	103.59	106.40
36	5	412	G	C8-N9-C4	-7.03	103.59	106.40
36	1	2689	A	N1-C6-N6	-7.03	114.38	118.60
36	5	1179	A	N1-C2-N3	7.03	132.82	129.30
36	5	1194	G	C8-N9-C4	-7.03	103.59	106.40
36	5	2169	G	N1-C6-O6	-7.03	115.68	119.90
36	1	2606	G	N3-C4-C5	-7.03	125.09	128.60
1	2	354	C	N3-C4-C5	-7.03	119.09	121.90
36	1	1834	U	N3-C4-C5	-7.03	110.38	114.60
36	1	2636	A	N7-C8-N9	7.02	117.31	113.80
1	6	1537	C	C6-N1-C1'	7.02	129.23	120.80
36	5	1147	G	C8-N9-C4	-7.02	103.59	106.40
36	1	1165	A	C8-N9-C4	7.02	108.61	105.80
36	1	1878	G	O4'-C1'-N9	-7.02	102.58	108.20
1	6	25	C	P-O3'-C3'	7.02	128.12	119.70
36	5	1370	G	N1-C2-N3	7.02	128.11	123.90
15	C3	22	ALA	C-N-CD	-7.01	105.17	120.60
36	1	1149	G	N1-C2-N3	7.01	128.11	123.90
36	1	1307	G	C8-N9-C4	-7.01	103.59	106.40
36	1	1329	U	C6-N1-C2	-7.01	116.79	121.00
1	6	53	G	N1-C2-N3	7.01	128.11	123.90
36	1	358	G	C6-C5-N7	-7.01	126.19	130.40
1	6	553	G	C5-C6-O6	-7.01	124.39	128.60
36	5	423	A	C8-N9-C4	7.01	108.60	105.80
36	5	651	G	OP2-P-O3'	7.01	120.62	105.20
36	5	874	U	C6-N1-C2	7.01	125.21	121.00
36	5	934	G	N3-C4-N9	7.01	130.21	126.00
36	5	1321	G	C5-C6-O6	-7.01	124.40	128.60
36	1	2356	A	C5-C6-N6	-7.00	118.10	123.70
36	5	1199	C	O5'-P-OP2	-7.00	99.40	105.70
36	5	2849	C	N3-C4-N4	7.00	122.90	118.00
36	1	2764	C	N3-C4-C5	-7.00	119.10	121.90
1	6	1634	C	C2-N1-C1'	7.00	126.50	118.80
36	5	519	A	C6-C5-N7	-7.00	127.40	132.30
36	5	2323	G	C8-N9-C4	-7.00	103.60	106.40
36	1	1595	U	C6-N1-C2	7.00	125.20	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2916	U	N1-C2-O2	7.00	127.70	122.80
36	1	1904	C	N3-C4-N4	6.99	122.89	118.00
36	5	1206	G	N3-C4-C5	-6.99	125.10	128.60
36	5	87	U	N3-C2-O2	-6.99	117.31	122.20
36	1	229	G	N1-C6-O6	6.99	124.09	119.90
36	1	2684	C	N3-C4-C5	-6.99	119.10	121.90
1	2	16	G	N3-C4-N9	6.99	130.19	126.00
1	6	610	G	C8-N9-C1'	-6.99	117.92	127.00
45	L8	189	LEU	CA-CB-CG	6.98	131.36	115.30
18	C6	40	GLU	C-N-CD	-6.98	105.24	120.60
36	5	661	G	N7-C8-N9	6.98	116.59	113.10
36	5	638	C	C6-N1-C2	-6.98	117.51	120.30
36	1	1365	G	N3-C4-C5	-6.98	125.11	128.60
1	6	173	A	N1-C6-N6	6.98	122.79	118.60
36	5	2695	A	C6-N1-C2	-6.98	114.42	118.60
36	5	2620	G	N3-C2-N2	6.97	124.78	119.90
36	5	2993	G	C4-C5-N7	6.97	113.59	110.80
36	1	3311	C	C6-N1-C2	6.97	123.09	120.30
1	6	426	G	N3-C4-C5	-6.97	125.11	128.60
36	5	2160	G	N1-C6-O6	6.97	124.08	119.90
36	1	3050	U	N1-C2-O2	6.97	127.68	122.80
36	1	59	G	N1-C6-O6	6.96	124.08	119.90
1	6	343	C	N1-C2-O2	-6.96	114.72	118.90
36	5	728	G	N3-C4-N9	6.96	130.18	126.00
36	1	1820	U	P-O3'-C3'	6.96	128.05	119.70
36	1	2374	C	N3-C4-C5	-6.96	119.12	121.90
36	5	2936	A	N9-C4-C5	6.96	108.58	105.80
36	1	424	G	C5-C6-N1	6.96	114.98	111.50
1	2	1568	C	P-O3'-C3'	6.96	128.05	119.70
1	2	1657	U	O4'-C1'-N1	6.96	113.77	108.20
36	5	2405	C	C6-N1-C2	-6.96	117.52	120.30
36	1	884	A	N1-C6-N6	6.96	122.77	118.60
36	1	3085	G	N1-C6-O6	6.96	124.07	119.90
1	6	47	A	O5'-P-OP1	-6.96	99.44	105.70
36	5	1556	C	N1-C2-O2	6.96	123.07	118.90
36	5	2976	A	N1-C6-N6	-6.96	114.43	118.60
1	2	1773	C	N3-C4-C5	-6.95	119.12	121.90
36	1	1522	U	C5-C4-O4	-6.95	121.73	125.90
36	1	785	G	C2-N3-C4	6.95	115.38	111.90
36	5	2783	U	N1-C2-O2	6.95	127.67	122.80
36	1	1157	G	C8-N9-C4	-6.95	103.62	106.40
37	3	80	G	C8-N9-C4	-6.95	103.62	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	4	140	G	C8-N9-C4	-6.95	103.62	106.40
36	5	36	C	C2-N3-C4	-6.95	116.43	119.90
36	5	2817	A	C5-C6-N6	-6.95	118.14	123.70
36	5	972	A	C4-C5-C6	6.95	120.47	117.00
1	2	1180	C	N1-C2-O2	6.94	123.07	118.90
1	2	553	G	C6-C5-N7	-6.94	126.24	130.40
36	1	2986	U	N1-C2-O2	-6.94	117.94	122.80
36	5	2653	C	N1-C2-O2	-6.94	114.74	118.90
36	5	567	G	C5-C6-O6	-6.94	124.44	128.60
56	n0	113	ARG	NE-CZ-NH1	-6.93	116.83	120.30
36	1	1433	A	N1-C6-N6	-6.93	114.44	118.60
36	1	1556	C	C2-N1-C1'	6.93	126.42	118.80
36	5	2858	U	N3-C4-O4	6.93	124.25	119.40
1	2	73	U	OP1-P-O3'	6.93	120.44	105.20
36	5	2950	G	C6-C5-N7	-6.93	126.24	130.40
36	5	2992	U	C5-C6-N1	6.93	126.16	122.70
36	1	2917	G	C5-C6-N1	6.92	114.96	111.50
36	5	2943	G	C6-C5-N7	-6.92	126.25	130.40
36	5	1319	G	N7-C8-N9	-6.92	109.64	113.10
1	6	1629	G	N3-C4-C5	-6.92	125.14	128.60
36	5	2281	A	N1-C6-N6	6.92	122.75	118.60
36	1	1899	G	C8-N9-C4	-6.92	103.63	106.40
52	m6	78	ARG	NE-CZ-NH2	-6.92	116.84	120.30
36	1	1450	G	C5-N7-C8	-6.91	100.84	104.30
36	5	2966	G	C8-N9-C4	-6.91	103.63	106.40
38	8	14	C	O5'-P-OP2	-6.91	99.48	105.70
36	1	2385	G	N3-C4-C5	6.91	132.06	128.60
36	5	3374	U	C6-N1-C2	6.91	125.15	121.00
36	1	2726	C	N3-C4-N4	-6.91	113.16	118.00
36	5	2723	U	C5-C6-N1	6.91	126.15	122.70
36	1	640	U	C6-N1-C2	-6.91	116.86	121.00
36	1	640	U	C5-C6-N1	6.91	126.15	122.70
36	5	3035	A	C8-N9-C4	6.91	108.56	105.80
36	5	2629	U	N3-C2-O2	6.90	127.03	122.20
37	3	82	G	N3-C4-C5	-6.90	125.15	128.60
36	5	1124	U	C5-C4-O4	6.90	130.04	125.90
36	1	713	U	C5-C6-N1	-6.90	119.25	122.70
36	1	2311	G	C8-N9-C4	-6.90	103.64	106.40
36	1	895	A	C6-C5-N7	-6.89	127.47	132.30
1	6	103	A	O4'-C1'-N9	6.89	113.71	108.20
36	5	940	G	N1-C6-O6	-6.89	115.77	119.90
38	8	100	U	C5-C6-N1	6.89	126.15	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	1077	U	C2-N1-C1'	-6.89	109.43	117.70
36	1	2938	G	N1-C2-N3	6.89	128.03	123.90
36	5	640	U	N3-C4-O4	6.89	124.22	119.40
37	3	105	C	OP2-P-O3'	6.88	120.34	105.20
36	5	3378	C	O5'-P-OP2	-6.88	99.50	105.70
36	1	1303	A	N9-C4-C5	-6.88	103.05	105.80
36	1	3210	A	O5'-P-OP2	-6.88	99.51	105.70
1	6	619	A	N1-C6-N6	-6.88	114.47	118.60
36	1	1897	G	C5-C6-O6	-6.88	124.47	128.60
36	5	1152	G	N7-C8-N9	6.88	116.54	113.10
36	5	2885	C	C5-C4-N4	-6.88	115.39	120.20
1	2	1274	C	C6-N1-C2	-6.88	117.55	120.30
38	4	29	U	N3-C4-O4	6.87	124.21	119.40
36	5	1604	G	N3-C4-C5	-6.87	125.16	128.60
36	5	2160	G	C6-C5-N7	-6.87	126.28	130.40
36	5	360	G	C5-C6-O6	6.87	132.72	128.60
36	5	2801	A	C8-N9-C4	6.87	108.55	105.80
36	5	36	C	N3-C4-C5	6.87	124.65	121.90
36	5	1513	G	N3-C4-C5	-6.87	125.17	128.60
36	5	2239	G	N1-C6-O6	6.87	124.02	119.90
36	5	2620	G	N9-C4-C5	-6.87	102.65	105.40
36	5	1115	G	N1-C6-O6	6.86	124.02	119.90
36	5	1367	G	C4-C5-C6	6.86	122.92	118.80
36	1	3054	U	N1-C2-N3	6.86	119.01	114.90
1	6	605	A	O5'-P-OP2	-6.86	99.53	105.70
52	M6	110	PRO	C-N-CD	-6.86	105.52	120.60
36	5	1192	C	C6-N1-C1'	-6.86	112.57	120.80
36	5	1604	G	C6-C5-N7	-6.86	126.29	130.40
36	5	2249	G	C8-N9-C4	-6.86	103.66	106.40
36	5	3326	G	C8-N9-C4	6.86	109.14	106.40
36	1	2688	U	C6-N1-C1'	-6.85	111.60	121.20
36	5	2842	U	O5'-P-OP2	6.85	118.92	110.70
36	5	1182	A	C5-N7-C8	-6.85	100.47	103.90
36	1	283	G	C5-N7-C8	-6.85	100.88	104.30
36	1	988	U	C5-C6-N1	-6.85	119.28	122.70
36	1	1312	C	O5'-P-OP1	-6.85	99.54	105.70
36	1	1437	C	N3-C4-C5	-6.85	119.16	121.90
36	1	3219	G	C5-C6-O6	-6.85	124.49	128.60
36	5	2531	C	N1-C2-O2	6.85	123.01	118.90
37	7	9	C	C6-N1-C2	-6.84	117.56	120.30
36	1	2606	G	C4-C5-C6	6.84	122.90	118.80
36	5	2620	G	C4-C5-N7	6.84	113.54	110.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2385	G	N3-C4-N9	-6.84	121.90	126.00
36	5	410	U	C5-C6-N1	6.84	126.12	122.70
36	5	3172	A	C5-C6-N6	-6.84	118.23	123.70
36	1	2392	C	C6-N1-C2	6.84	123.03	120.30
36	1	2283	G	N1-C6-O6	6.84	124.00	119.90
1	6	310	C	C6-N1-C2	-6.84	117.57	120.30
1	2	1560	U	N3-C2-O2	-6.83	117.42	122.20
1	6	160	C	N3-C2-O2	-6.83	117.11	121.90
36	5	666	A	N1-C6-N6	-6.83	114.50	118.60
36	5	2364	G	N9-C4-C5	6.83	108.13	105.40
36	5	104	G	C8-N9-C4	6.83	109.13	106.40
36	1	1191	U	O5'-P-OP1	-6.83	99.55	105.70
36	1	2699	G	N1-C6-O6	6.83	124.00	119.90
36	5	1370	G	C4-C5-N7	-6.82	108.07	110.80
36	5	1897	G	C6-C5-N7	-6.82	126.31	130.40
36	5	2331	C	O5'-P-OP1	-6.82	99.56	105.70
1	2	90	C	C6-N1-C2	-6.82	117.57	120.30
36	1	2853	A	C2-N3-C4	-6.82	107.19	110.60
36	5	934	G	C5-C6-N1	6.82	114.91	111.50
36	5	1434	G	O5'-P-OP2	-6.82	99.56	105.70
1	2	992	A	C5-N7-C8	-6.82	100.49	103.90
36	1	1428	A	N1-C2-N3	6.81	132.71	129.30
36	1	1175	C	C5-C6-N1	-6.81	117.59	121.00
1	6	176	C	N3-C2-O2	-6.81	117.14	121.90
36	5	2893	C	C6-N1-C2	-6.81	117.58	120.30
37	7	45	A	N1-C6-N6	-6.81	114.52	118.60
36	5	942	U	C4-C5-C6	6.80	123.78	119.70
36	5	2552	C	C2-N1-C1'	6.80	126.28	118.80
36	5	2856	G	N3-C2-N2	-6.80	115.14	119.90
36	5	3154	C	N3-C2-O2	-6.80	117.14	121.90
36	1	2846	U	N3-C2-O2	-6.80	117.44	122.20
36	1	608	A	C4-C5-C6	6.80	120.40	117.00
36	1	3278	C	N3-C4-C5	-6.80	119.18	121.90
36	5	2907	G	C5-N7-C8	-6.80	100.90	104.30
36	5	204	A	C8-N9-C4	-6.79	103.08	105.80
36	1	1115	G	N7-C8-N9	6.79	116.50	113.10
36	1	3344	A	N1-C6-N6	6.79	122.67	118.60
36	5	1449	A	N1-C6-N6	6.79	122.67	118.60
36	5	3225	C	C6-N1-C2	-6.79	117.58	120.30
36	1	332	C	C6-N1-C2	6.79	123.02	120.30
56	n0	144	LEU	CA-CB-CG	-6.79	99.69	115.30
36	5	339	C	N1-C2-O2	-6.79	114.83	118.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	2915	U	N3-C2-O2	6.79	126.95	122.20
36	1	2622	C	C6-N1-C2	-6.78	117.59	120.30
38	8	25	G	N3-C4-N9	6.78	130.07	126.00
36	1	1838	G	N1-C6-O6	6.78	123.97	119.90
1	6	250	C	C5-C6-N1	6.78	124.39	121.00
36	1	291	C	C6-N1-C2	-6.78	117.59	120.30
36	5	1550	C	C6-N1-C2	-6.78	117.59	120.30
36	5	1416	C	N3-C2-O2	-6.78	117.16	121.90
36	1	2237	C	C6-N1-C2	6.77	123.01	120.30
36	1	2688	U	C2-N1-C1'	6.77	125.83	117.70
36	1	3319	U	N3-C2-O2	-6.77	117.46	122.20
36	5	970	A	N9-C4-C5	-6.77	103.09	105.80
36	5	1710	C	C6-N1-C2	6.77	123.01	120.30
36	1	2326	A	N1-C6-N6	-6.77	114.54	118.60
36	1	2617	U	N1-C2-N3	6.77	118.96	114.90
36	1	3142	A	N1-C2-N3	6.77	132.68	129.30
36	5	2880	U	N3-C2-O2	-6.77	117.46	122.20
1	2	959	U	N3-C2-O2	-6.77	117.46	122.20
1	6	536	C	C6-N1-C2	-6.77	117.59	120.30
36	1	2662	G	N9-C4-C5	-6.76	102.69	105.40
36	5	2726	C	N1-C2-N3	6.76	123.94	119.20
36	5	715	A	C8-N9-C4	-6.76	103.09	105.80
36	5	1885	U	C6-N1-C2	6.76	125.06	121.00
1	2	1241	G	C4-N9-C1'	6.76	135.29	126.50
36	1	905	U	N1-C2-N3	6.76	118.96	114.90
1	6	1028	C	C6-N1-C2	6.76	123.00	120.30
36	5	1161	G	O5'-P-OP2	6.76	118.81	110.70
36	5	1348	U	N3-C2-O2	-6.76	117.47	122.20
38	4	53	A	C5-C6-N1	6.76	121.08	117.70
36	5	3382	U	N3-C2-O2	-6.76	117.47	122.20
36	5	907	G	N3-C4-N9	6.76	130.06	126.00
36	1	1366	A	N1-C6-N6	-6.76	114.55	118.60
38	8	99	C	C6-N1-C2	6.76	123.00	120.30
36	5	656	A	N1-C6-N6	6.75	122.65	118.60
36	5	1841	A	O5'-P-OP1	-6.75	99.62	105.70
36	5	1433	A	N7-C8-N9	6.75	117.18	113.80
38	8	17	A	C5-C6-N6	-6.75	118.30	123.70
1	6	543	C	N3-C2-O2	-6.75	117.17	121.90
1	6	1550	A	C8-N9-C4	-6.75	103.10	105.80
36	5	644	G	C8-N9-C4	-6.75	103.70	106.40
36	1	1885	U	C6-N1-C2	6.74	125.05	121.00
36	5	2865	U	C4-C5-C6	-6.74	115.65	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	970	A	C5-N7-C8	-6.74	100.53	103.90
36	5	917	A	C8-N9-C4	-6.74	103.10	105.80
36	1	519	A	C4-C5-C6	6.74	120.37	117.00
36	1	2860	U	C4-C5-C6	-6.74	115.66	119.70
36	5	718	G	O4'-C1'-N9	6.74	113.59	108.20
1	2	1200	G	C6-C5-N7	-6.74	126.36	130.40
36	5	2964	G	C5-C6-O6	6.74	132.64	128.60
12	C0	88	PRO	N-CA-CB	6.74	111.38	103.30
36	1	397	A	N1-C6-N6	-6.74	114.56	118.60
36	1	1846	C	N1-C2-O2	-6.74	114.86	118.90
1	6	1725	U	N3-C2-O2	-6.74	117.49	122.20
36	1	677	A	OP1-P-OP2	-6.73	109.50	119.60
36	5	1752	A	N1-C6-N6	6.73	122.64	118.60
36	5	3108	G	C2-N3-C4	-6.73	108.53	111.90
36	1	1306	G	N1-C6-O6	6.73	123.94	119.90
24	d2	93	LEU	CA-CB-CG	6.73	130.77	115.30
36	1	1292	C	C6-N1-C2	6.72	122.99	120.30
36	1	2836	C	C5-C4-N4	6.72	124.91	120.20
36	1	108	A	N1-C6-N6	6.72	122.63	118.60
36	1	1077	U	C5-C6-N1	-6.72	119.34	122.70
36	1	1340	G	C5-C6-O6	-6.72	124.57	128.60
36	1	2549	G	N1-C2-N2	-6.72	110.15	116.20
1	6	1782	A	C8-N9-C4	-6.72	103.11	105.80
36	5	1339	C	O5'-P-OP1	-6.72	99.65	105.70
36	1	1899	G	N9-C4-C5	6.72	108.09	105.40
36	1	864	G	O5'-P-OP1	-6.72	99.66	105.70
36	1	905	U	N1-C2-O2	-6.72	118.10	122.80
36	1	2787	G	C6-C5-N7	-6.72	126.37	130.40
36	1	921	A	C4-C5-N7	-6.71	107.34	110.70
36	1	514	G	N1-C6-O6	6.71	123.93	119.90
36	1	1165	A	C2-N3-C4	-6.71	107.24	110.60
36	1	2875	U	N1-C2-O2	6.71	127.50	122.80
36	5	46	U	N1-C2-O2	-6.71	118.10	122.80
36	5	725	G	C5-C6-N1	-6.71	108.14	111.50
36	1	709	A	N9-C4-C5	-6.71	103.12	105.80
37	7	49	G	N3-C4-N9	-6.71	121.97	126.00
36	1	212	G	N1-C6-O6	6.71	123.92	119.90
36	5	1900	A	C5-C6-N6	-6.71	118.33	123.70
36	1	421	G	N3-C4-C5	-6.70	125.25	128.60
36	5	2953	U	N3-C4-O4	6.70	124.09	119.40
36	1	1371	G	C8-N9-C4	6.70	109.08	106.40
36	1	1835	A	C8-N9-C4	6.70	108.48	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	1452	A	C4-C5-N7	6.70	114.05	110.70
37	7	1	G	N1-C2-N2	-6.70	110.17	116.20
36	5	788	C	OP2-P-O3'	6.70	119.94	105.20
1	2	453	U	N3-C2-O2	-6.70	117.51	122.20
36	1	39	A	C5-C6-N6	6.70	129.06	123.70
36	5	2966	G	C4-C5-C6	6.70	122.82	118.80
36	5	2552	C	N3-C2-O2	-6.69	117.22	121.90
1	2	1194	A	C8-N9-C4	-6.69	103.12	105.80
36	5	3326	G	N9-C4-C5	-6.69	102.72	105.40
36	1	906	A	C4-C5-C6	6.69	120.34	117.00
36	1	3278	C	N3-C2-O2	-6.69	117.22	121.90
36	1	591	G	O5'-P-OP1	6.68	118.72	110.70
36	1	3306	U	N3-C4-O4	-6.68	114.72	119.40
36	1	96	G	C2-N3-C4	-6.68	108.56	111.90
36	1	643	U	N1-C2-O2	6.68	127.48	122.80
36	1	1198	C	C6-N1-C2	-6.68	117.63	120.30
36	5	1152	G	N9-C4-C5	6.68	108.07	105.40
1	6	1285	U	C2-N1-C1'	6.68	125.71	117.70
1	2	577	G	C6-C5-N7	-6.68	126.39	130.40
36	1	1065	A	O5'-P-OP1	-6.67	99.69	105.70
36	1	864	G	N1-C6-O6	-6.67	115.90	119.90
36	1	3134	A	C8-N9-C4	6.67	108.47	105.80
36	5	3245	A	C4-C5-C6	6.67	120.33	117.00
36	1	2400	G	C5-C6-O6	-6.67	124.60	128.60
36	5	40	A	N1-C6-N6	6.67	122.60	118.60
36	5	798	G	N7-C8-N9	6.67	116.43	113.10
36	5	3024	A	C2-N3-C4	-6.66	107.27	110.60
36	1	2606	G	N3-C4-N9	6.66	130.00	126.00
1	2	16	G	C6-C5-N7	-6.66	126.41	130.40
1	6	1481	C	N3-C2-O2	-6.66	117.24	121.90
1	6	1751	C	C5-C6-N1	-6.66	117.67	121.00
36	5	3144	G	N3-C4-C5	-6.66	125.27	128.60
36	1	2647	A	C8-N9-C4	-6.66	103.14	105.80
1	2	1654	G	N3-C4-C5	-6.66	125.27	128.60
36	1	983	A	C6-N1-C2	-6.66	114.61	118.60
36	5	2645	G	C5-C6-O6	6.66	132.59	128.60
36	1	803	C	C6-N1-C1'	-6.65	112.82	120.80
36	1	969	C	N1-C2-O2	-6.65	114.91	118.90
36	1	1450	G	C4-C5-N7	6.65	113.46	110.80
36	5	371	G	N1-C6-O6	-6.65	115.91	119.90
36	1	913	A	O5'-P-OP1	-6.65	99.72	105.70
36	5	2698	G	C8-N9-C4	6.65	109.06	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	2	36	C	C6-N1-C2	6.65	122.96	120.30
1	2	75	U	N1-C2-O2	6.65	127.45	122.80
36	5	2372	A	P-O3'-C3'	6.65	127.68	119.70
36	1	638	C	N1-C2-O2	6.64	122.89	118.90
36	1	2615	G	C5-C6-O6	-6.64	124.61	128.60
36	5	523	A	N1-C6-N6	-6.64	114.61	118.60
1	2	1458	G	C4-C5-N7	6.64	113.46	110.80
1	6	473	A	N1-C6-N6	-6.64	114.62	118.60
36	1	2329	C	O5'-P-OP2	-6.64	99.73	105.70
1	6	619	A	N9-C4-C5	6.64	108.45	105.80
36	5	410	U	N3-C4-O4	6.64	124.05	119.40
36	5	803	C	C5-C6-N1	6.64	124.32	121.00
36	5	709	A	C8-N9-C4	6.63	108.45	105.80
36	1	300	G	N1-C6-O6	-6.63	115.92	119.90
36	1	2708	C	C6-N1-C2	6.63	122.95	120.30
36	5	414	U	N3-C2-O2	6.63	126.84	122.20
36	5	2758	A	C8-N9-C4	-6.63	103.15	105.80
36	1	2279	A	C4-C5-N7	6.63	114.01	110.70
37	3	91	G	C5-C6-O6	-6.63	124.62	128.60
36	5	2827	U	N3-C2-O2	-6.63	117.56	122.20
36	1	2921	U	O5'-P-OP1	6.63	118.65	110.70
37	7	82	G	C5-C6-O6	-6.63	124.62	128.60
38	4	93	U	N3-C4-O4	-6.62	114.76	119.40
36	5	822	G	O5'-P-OP1	-6.62	99.74	105.70
36	5	2620	G	C5-C6-N1	6.62	114.81	111.50
36	5	2966	G	N3-C4-C5	-6.62	125.29	128.60
36	5	3295	A	C5-C6-N1	6.62	121.01	117.70
1	6	1074	G	N1-C6-O6	6.62	123.87	119.90
36	1	2884	C	C6-N1-C2	6.61	122.94	120.30
36	1	2621	G	N3-C2-N2	-6.61	115.27	119.90
36	5	1588	A	N1-C6-N6	-6.61	114.63	118.60
36	1	3275	U	C6-N1-C2	-6.61	117.03	121.00
36	1	65	A	P-O3'-C3'	6.61	127.63	119.70
36	1	1307	G	N1-C6-O6	-6.61	115.94	119.90
36	5	2412	G	N3-C4-C5	-6.61	125.30	128.60
36	5	2783	U	N3-C2-O2	-6.60	117.58	122.20
36	1	424	G	C4-C5-N7	6.60	113.44	110.80
36	1	2434	U	N3-C2-O2	-6.60	117.58	122.20
1	6	904	G	N3-C4-C5	-6.60	125.30	128.60
36	5	1149	G	C6-C5-N7	-6.60	126.44	130.40
36	5	1416	C	N1-C2-O2	6.60	122.86	118.90
36	5	2423	U	N3-C2-O2	-6.60	117.58	122.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	1103	A	C8-N9-C4	6.60	108.44	105.80
36	5	2167	A	C8-N9-C4	-6.60	103.16	105.80
36	5	2639	G	C6-C5-N7	-6.60	126.44	130.40
36	1	2174	G	C8-N9-C4	-6.59	103.76	106.40
1	2	623	A	O5'-P-OP1	-6.59	99.77	105.70
1	6	359	A	N3-C4-C5	6.59	131.41	126.80
1	6	1773	C	N3-C4-N4	6.59	122.61	118.00
36	1	3275	U	C5-C6-N1	6.59	126.00	122.70
36	5	424	G	C4-C5-N7	6.59	113.44	110.80
36	1	2416	U	C5-C6-N1	6.59	126.00	122.70
36	1	2310	U	O5'-P-OP1	-6.59	99.77	105.70
36	5	728	G	C6-C5-N7	-6.59	126.45	130.40
36	5	931	C	N3-C4-C5	6.58	124.53	121.90
36	5	2849	C	O5'-P-OP2	-6.58	99.78	105.70
36	5	2966	G	C6-C5-N7	-6.58	126.45	130.40
38	8	46	G	O5'-P-OP1	-6.58	99.77	105.70
36	1	663	C	N3-C4-C5	-6.58	119.27	121.90
36	5	1379	G	C6-C5-N7	-6.58	126.45	130.40
36	1	1400	G	C4-N9-C1'	6.58	135.05	126.50
36	1	3092	C	C5-C6-N1	-6.58	117.71	121.00
62	N6	126	LEU	CA-CB-CG	6.58	130.43	115.30
36	5	91	G	N9-C4-C5	-6.58	102.77	105.40
1	2	1490	C	O5'-P-OP1	-6.58	99.78	105.70
36	1	685	G	C8-N9-C4	6.58	109.03	106.40
36	1	2395	G	C4-C5-N7	6.58	113.43	110.80
36	1	2525	G	C6-C5-N7	-6.58	126.45	130.40
36	5	2403	G	N1-C6-O6	6.58	123.85	119.90
36	5	2434	U	O5'-P-OP2	-6.58	99.78	105.70
36	1	1429	G	N1-C2-N3	6.58	127.85	123.90
1	2	402	C	C6-N1-C2	6.57	122.93	120.30
36	1	1139	G	O5'-P-OP1	-6.57	99.78	105.70
36	1	3278	C	C6-N1-C2	-6.57	117.67	120.30
36	5	1370	G	N3-C4-C5	-6.57	125.31	128.60
36	1	1054	A	O5'-P-OP1	6.57	118.58	110.70
36	1	1906	G	N1-C6-O6	6.57	123.84	119.90
36	1	2821	C	O5'-P-OP1	-6.57	99.79	105.70
36	1	358	G	C5-C6-O6	-6.57	124.66	128.60
36	5	2947	G	C5-C6-O6	-6.57	124.66	128.60
1	2	830	U	C2-N1-C1'	6.56	125.58	117.70
36	1	1389	G	C6-C5-N7	-6.56	126.46	130.40
36	5	2832	C	C6-N1-C2	6.56	122.92	120.30
36	5	2763	U	OP2-P-O3'	6.56	119.63	105.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	6	553	G	C6-C5-N7	-6.56	126.46	130.40
36	5	1589	A	C8-N9-C4	-6.56	103.18	105.80
36	1	685	G	N1-C6-O6	6.56	123.83	119.90
36	1	1495	U	C4-C5-C6	6.56	123.63	119.70
36	1	2556	C	N1-C2-O2	6.56	122.83	118.90
36	5	424	G	O5'-P-OP1	6.56	118.57	110.70
36	5	1440	G	C5-C6-O6	6.56	132.53	128.60
1	6	272	U	N3-C2-O2	-6.56	117.61	122.20
36	1	3278	C	C2-N3-C4	6.55	123.18	119.90
36	5	586	C	N1-C2-O2	-6.55	114.97	118.90
36	1	608	A	C6-C5-N7	-6.55	127.71	132.30
36	5	970	A	N1-C6-N6	6.55	122.53	118.60
36	1	49	A	N1-C6-N6	6.55	122.53	118.60
36	5	641	C	O5'-P-OP1	-6.55	99.81	105.70
38	4	113	U	N1-C2-N3	6.55	118.83	114.90
1	6	272	U	C2-N1-C1'	6.55	125.56	117.70
1	6	1031	U	N1-C2-O2	-6.55	118.22	122.80
36	1	784	A	N1-C6-N6	6.54	122.53	118.60
36	1	1849	C	N3-C4-C5	6.54	124.52	121.90
36	5	1379	G	N1-C6-O6	6.54	123.83	119.90
36	1	1544	G	C5-C6-O6	-6.54	124.67	128.60
36	1	221	A	O5'-P-OP2	-6.54	99.81	105.70
36	5	633	C	N3-C4-C5	-6.54	119.28	121.90
36	5	2832	C	O5'-P-OP1	-6.54	99.81	105.70
36	5	3103	A	O5'-P-OP2	-6.54	99.81	105.70
36	5	1148	G	N9-C4-C5	-6.54	102.78	105.40
36	5	3195	U	OP1-P-O3'	6.54	119.59	105.20
36	1	39	A	N9-C4-C5	6.54	108.41	105.80
36	1	29	C	N3-C4-N4	6.53	122.57	118.00
36	1	1848	G	N3-C2-N2	6.53	124.47	119.90
46	L9	176	LEU	CA-CB-CG	-6.53	100.28	115.30
36	5	1417	G	N7-C8-N9	6.53	116.37	113.10
1	2	407	A	N1-C6-N6	-6.53	114.68	118.60
36	1	2755	C	O5'-P-OP1	-6.53	99.82	105.70
37	3	104	A	N1-C6-N6	-6.53	114.68	118.60
36	5	1012	G	C8-N9-C1'	6.53	135.49	127.00
36	5	1317	A	C4-C5-N7	6.53	113.97	110.70
37	7	51	A	N1-C6-N6	6.53	122.52	118.60
36	1	612	U	N1-C2-N3	6.53	118.82	114.90
1	6	325	G	C8-N9-C4	-6.53	103.79	106.40
1	2	1467	C	C6-N1-C2	-6.53	117.69	120.30
36	1	922	U	N1-C2-O2	6.53	127.37	122.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	1857	C	C6-N1-C2	6.53	122.91	120.30
1	6	144	U	N3-C2-O2	-6.52	117.63	122.20
1	6	431	C	C2-N1-C1'	-6.52	111.63	118.80
36	5	3245	A	C4-C5-N7	6.52	113.96	110.70
36	1	993	G	N3-C4-C5	-6.52	125.34	128.60
36	1	2247	G	C4-N9-C1'	6.52	134.98	126.50
36	5	808	A	C8-N9-C4	-6.52	103.19	105.80
36	5	1133	A	N1-C6-N6	6.52	122.51	118.60
36	5	2933	A	N1-C6-N6	6.52	122.51	118.60
36	5	304	G	N9-C4-C5	6.52	108.01	105.40
36	5	800	G	N3-C4-N9	6.52	129.91	126.00
36	5	2282	U	C6-N1-C2	6.52	124.91	121.00
36	1	410	U	C6-N1-C2	-6.52	117.09	121.00
36	5	3015	G	N1-C6-O6	6.52	123.81	119.90
38	4	46	G	C5-C6-O6	6.51	132.51	128.60
36	1	3191	G	O5'-P-OP2	-6.51	99.84	105.70
36	1	197	G	C5-C6-O6	-6.51	124.69	128.60
36	5	587	U	C5-C4-O4	-6.51	121.99	125.90
36	5	942	U	N1-C2-O2	-6.51	118.24	122.80
36	5	2419	A	O5'-P-OP2	6.51	118.51	110.70
1	2	426	G	C6-C5-N7	-6.51	126.49	130.40
36	1	131	C	C6-N1-C2	-6.51	117.70	120.30
36	1	1400	G	N3-C4-C5	-6.51	125.34	128.60
36	5	3269	U	P-O3'-C3'	6.51	127.51	119.70
36	1	2136	C	N3-C4-C5	-6.51	119.30	121.90
36	5	3108	G	N1-C2-N3	6.50	127.80	123.90
36	1	339	C	N3-C4-C5	6.50	124.50	121.90
36	5	1159	A	C2-N3-C4	-6.50	107.35	110.60
36	5	1160	C	C6-N1-C1'	6.50	128.60	120.80
36	5	2892	A	N1-C6-N6	6.50	122.50	118.60
36	1	3212	C	C2-N1-C1'	-6.50	111.65	118.80
36	5	87	U	C6-N1-C2	-6.50	117.10	121.00
1	2	1082	C	N1-C2-O2	6.50	122.80	118.90
36	1	1518	U	C4-C5-C6	6.50	123.60	119.70
36	5	1130	A	C5-C6-N1	6.50	120.95	117.70
36	1	895	A	C5-N7-C8	-6.49	100.65	103.90
36	1	2422	C	N3-C2-O2	-6.49	117.36	121.90
36	5	720	A	N9-C4-C5	-6.49	103.20	105.80
36	5	2662	G	C8-N9-C1'	-6.49	118.56	127.00
36	1	2760	C	N1-C2-O2	-6.49	115.01	118.90
36	5	2875	U	N1-C1'-C2'	-6.49	104.86	112.00
36	1	1604	G	C4-N9-C1'	6.49	134.94	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2899	C	N3-C4-N4	6.49	122.54	118.00
36	5	1303	A	C8-N9-C4	6.49	108.39	105.80
1	6	965	U	N1-C2-O2	6.48	127.34	122.80
1	6	1747	G	O5'-P-OP2	-6.48	99.87	105.70
36	5	718	G	C8-N9-C1'	-6.48	118.57	127.00
36	5	2709	C	N3-C4-C5	6.48	124.49	121.90
36	1	225	C	N3-C4-C5	-6.48	119.31	121.90
36	1	1846	C	N3-C4-C5	-6.48	119.31	121.90
36	1	3214	U	N3-C2-O2	-6.48	117.66	122.20
1	6	542	A	N7-C8-N9	6.48	117.04	113.80
36	5	1135	A	N1-C6-N6	-6.48	114.71	118.60
38	4	29	U	C5-C4-O4	-6.48	122.01	125.90
36	5	1075	A	C8-N9-C4	6.48	108.39	105.80
36	5	2285	C	C6-N1-C1'	6.48	128.57	120.80
36	1	1279	C	C6-N1-C2	-6.48	117.71	120.30
36	5	2699	G	C5-C6-O6	-6.47	124.72	128.60
36	1	1332	A	O5'-P-OP1	-6.47	99.88	105.70
36	5	319	A	C8-N9-C4	6.47	108.39	105.80
1	2	334	G	N3-C4-N9	-6.47	122.12	126.00
36	1	651	G	N3-C4-N9	6.47	129.88	126.00
1	6	65	A	C2-N3-C4	-6.47	107.36	110.60
36	5	2585	G	N3-C4-N9	6.47	129.88	126.00
1	2	396	G	N3-C4-C5	6.47	131.83	128.60
36	5	793	C	C6-N1-C2	-6.47	117.71	120.30
36	1	421	G	C2-N3-C4	6.46	115.13	111.90
36	1	1367	G	N1-C6-O6	6.46	123.78	119.90
1	6	334	G	N3-C2-N2	-6.46	115.38	119.90
36	5	2307	G	N1-C6-O6	-6.46	116.02	119.90
36	1	1008	U	C5-C6-N1	-6.46	119.47	122.70
38	4	16	G	C8-N9-C4	6.46	108.98	106.40
1	6	334	G	N3-C4-N9	-6.46	122.12	126.00
36	5	2316	G	C4-C5-N7	-6.46	108.22	110.80
36	5	3214	U	N3-C2-O2	-6.46	117.68	122.20
1	2	323	A	O5'-P-OP2	-6.46	99.89	105.70
37	3	58	C	C6-N1-C2	-6.46	117.72	120.30
37	7	51	A	N1-C2-N3	6.46	132.53	129.30
36	1	369	A	N7-C8-N9	6.46	117.03	113.80
36	1	1886	A	N9-C4-C5	6.46	108.38	105.80
1	6	1564	U	C6-N1-C2	6.46	124.87	121.00
36	5	3313	U	O5'-P-OP2	-6.46	99.89	105.70
36	1	1005	G	C8-N9-C4	6.45	108.98	106.40
36	1	1404	G	C4-C5-N7	-6.45	108.22	110.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	3387	U	C5-C6-N1	-6.45	119.47	122.70
36	1	940	G	C5-C6-N1	6.45	114.73	111.50
36	5	720	A	C4-C5-N7	6.45	113.93	110.70
37	7	92	A	N1-C6-N6	6.45	122.47	118.60
36	1	145	G	C5-C6-O6	-6.45	124.73	128.60
36	1	1901	A	C2-N3-C4	6.45	113.83	110.60
36	1	2772	C	O4'-C1'-N1	6.45	113.36	108.20
36	5	1301	A	C4-C5-C6	6.45	120.22	117.00
36	1	39	A	N3-C4-N9	-6.45	122.24	127.40
36	1	645	A	C6-N1-C2	-6.45	114.73	118.60
36	5	781	G	N3-C4-C5	-6.45	125.38	128.60
36	1	218	G	N1-C6-O6	6.45	123.77	119.90
36	1	809	G	C8-N9-C4	6.45	108.98	106.40
36	5	976	U	O5'-P-OP2	-6.45	99.90	105.70
1	2	543	C	N1-C2-O2	6.44	122.77	118.90
36	1	1429	G	C4-C5-N7	-6.44	108.22	110.80
1	6	1751	C	C6-N1-C2	6.44	122.88	120.30
36	5	421	G	C4-N9-C1'	6.44	134.87	126.50
1	2	577	G	C5-N7-C8	-6.44	101.08	104.30
36	1	658	G	C8-N9-C1'	-6.44	118.63	127.00
36	5	1126	G	C4-C5-C6	6.44	122.66	118.80
36	5	2392	C	C5-C6-N1	-6.44	117.78	121.00
37	7	75	G	C6-C5-N7	-6.44	126.54	130.40
1	2	1761	U	N1-C2-N3	6.44	118.76	114.90
1	6	1629	G	C8-N9-C4	-6.44	103.83	106.40
1	2	1280	C	N3-C4-C5	-6.43	119.33	121.90
36	1	676	G	N3-C2-N2	6.43	124.40	119.90
36	1	924	G	O5'-P-OP1	-6.43	99.91	105.70
36	1	1807	G	C4-N9-C1'	6.43	134.86	126.50
36	1	2875	U	P-O3'-C3'	-6.43	111.98	119.70
1	6	1645	G	N1-C6-O6	-6.43	116.04	119.90
36	5	1851	G	C5-C6-N1	-6.43	108.28	111.50
36	5	3303	G	O5'-P-OP2	-6.43	99.91	105.70
36	5	1369	A	C5-C6-N1	-6.43	114.48	117.70
36	5	2624	G	C6-C5-N7	-6.43	126.54	130.40
36	5	2805	G	N1-C6-O6	6.43	123.76	119.90
36	5	2647	A	N1-C6-N6	-6.43	114.74	118.60
36	5	1443	G	C4-C5-N7	6.43	113.37	110.80
36	5	2278	C	N1-C2-O2	6.42	122.75	118.90
36	5	3244	A	O4'-C1'-N9	-6.42	103.06	108.20
36	5	800	G	N9-C4-C5	-6.42	102.83	105.40
36	5	220	G	O5'-P-OP2	-6.42	99.92	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	1452	A	C5-C6-N6	-6.42	118.56	123.70
36	5	2334	U	C6-N1-C2	-6.42	117.15	121.00
36	1	3379	C	O5'-P-OP2	-6.42	99.92	105.70
36	5	2420	C	N3-C2-O2	6.42	126.39	121.90
1	2	1539	G	C6-C5-N7	-6.42	126.55	130.40
36	1	2353	G	N1-C6-O6	6.42	123.75	119.90
1	6	272	U	C6-N1-C2	-6.42	117.15	121.00
36	5	936	A	O4'-C1'-N9	6.42	113.33	108.20
36	1	727	G	N1-C6-O6	-6.42	116.05	119.90
36	1	857	G	N1-C6-O6	6.41	123.75	119.90
36	1	2355	G	C4-C5-N7	6.41	113.36	110.80
36	5	2942	C	N3-C4-N4	6.41	122.49	118.00
36	5	3107	U	OP2-P-O3'	6.41	119.31	105.20
36	1	2889	C	N3-C2-O2	-6.41	117.41	121.90
36	1	2987	A	N1-C6-N6	6.41	122.45	118.60
36	1	890	C	N3-C4-C5	6.41	124.46	121.90
38	4	81	U	N1-C2-O2	6.41	127.29	122.80
36	5	1434	G	C8-N9-C4	-6.41	103.84	106.40
36	1	895	A	C4-C5-N7	6.41	113.91	110.70
1	6	1790	A	C8-N9-C4	6.41	108.36	105.80
36	5	1851	G	N1-C6-O6	6.41	123.75	119.90
36	1	2699	G	C6-C5-N7	-6.41	126.56	130.40
36	5	1481	A	C8-N9-C4	-6.41	103.24	105.80
36	5	3153	U	N1-C2-O2	6.40	127.28	122.80
36	1	1054	A	O5'-P-OP2	-6.40	99.94	105.70
36	5	916	G	O5'-P-OP1	-6.40	99.94	105.70
36	1	809	G	N9-C4-C5	-6.40	102.84	105.40
36	1	2860	U	C2-N1-C1'	6.40	125.38	117.70
36	5	2695	A	C5-C6-N1	6.40	120.90	117.70
36	5	3218	A	N1-C6-N6	6.40	122.44	118.60
1	6	298	C	C6-N1-C2	-6.40	117.74	120.30
36	5	84	U	N1-C2-O2	6.40	127.28	122.80
1	2	1116	A	N1-C6-N6	6.40	122.44	118.60
36	1	1536	G	N3-C2-N2	-6.40	115.42	119.90
36	5	345	G	C5-C6-O6	-6.39	124.76	128.60
36	5	968	G	C6-C5-N7	-6.39	126.56	130.40
36	5	3362	A	O4'-C1'-N9	6.39	113.32	108.20
36	5	874	U	C5-C6-N1	-6.39	119.50	122.70
36	5	2158	A	C5-C6-N1	6.39	120.90	117.70
36	1	2844	C	C6-N1-C2	6.39	122.86	120.30
36	5	2129	U	C5-C4-O4	6.39	129.74	125.90
36	1	39	A	C6-C5-N7	6.39	136.77	132.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	109	A	OP1-P-O3'	6.39	119.26	105.20
35	sM	167	PRO	N-CA-CB	6.39	110.97	103.30
36	5	3209	A	O4'-C1'-N9	6.39	113.31	108.20
36	1	720	A	N1-C6-N6	6.39	122.43	118.60
36	1	1001	G	C8-N9-C4	6.39	108.95	106.40
36	1	632	G	C6-C5-N7	-6.39	126.57	130.40
36	1	937	G	C8-N9-C4	6.39	108.95	106.40
36	1	3003	G	C8-N9-C4	6.39	108.95	106.40
36	5	1514	G	N3-C4-N9	6.39	129.83	126.00
36	1	2383	C	C5-C6-N1	6.38	124.19	121.00
36	5	429	U	C5-C6-N1	-6.38	119.51	122.70
36	5	867	G	N1-C6-O6	6.38	123.73	119.90
1	6	18	C	C6-N1-C2	-6.38	117.75	120.30
36	5	1161	G	N3-C4-N9	6.38	129.83	126.00
1	2	1297	G	N9-C4-C5	-6.38	102.85	105.40
36	1	933	A	C5-C6-N6	-6.38	118.60	123.70
36	1	1329	U	C5-C6-N1	6.38	125.89	122.70
36	1	2606	G	C6-C5-N7	-6.38	126.57	130.40
36	5	414	U	N1-C2-O2	-6.38	118.34	122.80
36	5	799	G	N1-C6-O6	-6.38	116.07	119.90
1	6	1169	G	C8-N9-C4	-6.38	103.85	106.40
36	5	1161	G	C5-C6-N1	6.38	114.69	111.50
37	7	51	A	C4-C5-C6	6.38	120.19	117.00
36	1	968	G	C6-C5-N7	-6.37	126.58	130.40
36	1	1792	C	N1-C2-O2	-6.37	115.08	118.90
36	5	2736	A	C6-C5-N7	-6.37	127.84	132.30
36	5	2308	C	N3-C2-O2	-6.37	117.44	121.90
36	1	111	C	N3-C2-O2	6.37	126.36	121.90
1	6	478	A	N1-C6-N6	6.37	122.42	118.60
1	2	351	C	N3-C2-O2	-6.37	117.44	121.90
36	1	2620	G	N1-C2-N3	-6.37	120.08	123.90
36	5	1599	G	C8-N9-C4	6.37	108.95	106.40
36	5	2968	G	O4'-C1'-N9	-6.37	103.11	108.20
36	5	3107	U	C6-N1-C2	-6.37	117.18	121.00
36	5	1497	C	N3-C2-O2	6.37	126.36	121.90
36	1	86	G	C4-N9-C1'	-6.36	118.23	126.50
36	5	2890	A	C8-N9-C4	-6.36	103.25	105.80
50	m4	72	LEU	CA-CB-CG	6.36	129.94	115.30
36	1	283	G	N7-C8-N9	6.36	116.28	113.10
36	1	906	A	C5-C6-N6	-6.36	118.61	123.70
36	5	3208	G	C6-C5-N7	-6.36	126.58	130.40
1	2	1170	G	C4-N9-C1'	6.36	134.77	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	3192	U	N3-C4-O4	6.36	123.85	119.40
1	2	553	G	C4-C5-N7	6.36	113.34	110.80
36	1	2937	G	C8-N9-C4	6.35	108.94	106.40
1	6	553	G	C4-C5-N7	6.35	113.34	110.80
36	5	2421	U	C6-N1-C2	-6.35	117.19	121.00
36	1	2662	G	C5-C6-O6	-6.35	124.79	128.60
36	1	1349	G	N3-C4-N9	6.35	129.81	126.00
36	5	2856	G	N1-C6-O6	6.35	123.71	119.90
1	2	16	G	N3-C4-C5	-6.35	125.43	128.60
36	1	2190	U	C4-C5-C6	6.35	123.51	119.70
36	1	66	A	O5'-P-OP2	6.34	118.31	110.70
1	2	1657	U	C4-C5-C6	6.34	123.51	119.70
36	1	2620	G	N1-C2-N2	6.34	121.91	116.20
1	6	1606	C	O5'-P-OP2	-6.34	99.99	105.70
36	5	358	G	N3-C4-C5	6.34	131.77	128.60
36	1	2549	G	N3-C2-N2	6.34	124.34	119.90
36	1	2877	G	N9-C4-C5	6.34	107.94	105.40
1	6	1560	U	N3-C2-O2	-6.34	117.76	122.20
1	2	1458	G	C4-N9-C1'	6.34	134.74	126.50
36	1	2923	U	C6-N1-C2	6.34	124.80	121.00
28	d6	42	ARG	NE-CZ-NH2	6.34	123.47	120.30
36	5	1408	G	OP2-P-O3'	6.34	119.14	105.20
45	l8	69	LEU	CA-CB-CG	6.34	129.88	115.30
36	1	3140	G	C4-C5-N7	6.34	113.33	110.80
36	1	3214	U	O4'-C1'-N1	6.34	113.27	108.20
1	6	410	A	N1-C6-N6	6.34	122.40	118.60
1	6	467	G	C8-N9-C4	6.34	108.94	106.40
36	5	170	G	C4-N9-C1'	6.34	134.74	126.50
1	6	943	C	O5'-P-OP1	-6.33	100.00	105.70
36	1	1305	U	C5-C4-O4	6.33	129.70	125.90
36	1	2167	A	O5'-P-OP1	-6.33	100.00	105.70
36	5	595	G	C8-N9-C4	-6.33	103.87	106.40
36	5	3295	A	N1-C6-N6	-6.33	114.80	118.60
36	5	1158	A	N1-C6-N6	6.33	122.40	118.60
36	5	2129	U	N3-C2-O2	-6.33	117.77	122.20
36	5	2531	C	N3-C2-O2	-6.33	117.47	121.90
35	SM	167	PRO	N-CA-CB	6.33	110.89	103.30
36	1	2355	G	C5-C6-O6	-6.33	124.80	128.60
36	5	2377	G	N1-C6-O6	-6.33	116.10	119.90
36	5	2899	C	N3-C2-O2	-6.33	117.47	121.90
36	5	3315	G	C4-C5-C6	6.33	122.60	118.80
36	1	878	G	C4-C5-N7	-6.32	108.27	110.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	2	1363	U	N1-C2-O2	6.32	127.23	122.80
36	1	961	C	C6-N1-C2	6.32	122.83	120.30
36	1	1428	A	C6-N1-C2	-6.32	114.81	118.60
36	5	2976	A	C5-C6-N6	6.32	128.76	123.70
1	2	992	A	N1-C6-N6	6.32	122.39	118.60
36	1	3269	U	C6-N1-C2	-6.32	117.21	121.00
36	1	230	U	N1-C2-N3	6.32	118.69	114.90
36	1	2373	A	O5'-P-OP1	-6.32	100.01	105.70
1	6	1428	G	C8-N9-C4	-6.32	103.87	106.40
36	5	204	A	N9-C4-C5	6.32	108.33	105.80
36	5	869	G	C5-C6-O6	-6.32	124.81	128.60
36	5	2692	A	C8-N9-C4	-6.32	103.27	105.80
36	1	1166	G	C5-N7-C8	-6.31	101.14	104.30
52	m6	84	LEU	CA-CB-CG	-6.31	100.78	115.30
37	7	35	C	N1-C2-O2	6.31	122.69	118.90
36	1	840	C	C5-C4-N4	-6.31	115.78	120.20
36	1	1851	G	C4-C5-N7	6.31	113.32	110.80
36	1	833	G	N1-C6-O6	-6.31	116.11	119.90
36	1	1891	A	C8-N9-C4	6.31	108.32	105.80
36	5	2403	G	C5-N7-C8	6.31	107.45	104.30
36	5	2945	G	N3-C4-N9	6.31	129.79	126.00
36	1	364	G	C5-C6-O6	-6.31	124.82	128.60
36	1	639	G	N1-C6-O6	6.31	123.69	119.90
36	1	3319	U	N1-C2-O2	6.31	127.22	122.80
36	5	718	G	N3-C4-C5	-6.31	125.45	128.60
36	1	2828	G	N3-C2-N2	6.31	124.31	119.90
1	6	1672	G	N3-C4-N9	6.30	129.78	126.00
36	5	514	G	C5-C6-O6	-6.30	124.82	128.60
36	5	2868	U	N1-C2-O2	6.30	127.21	122.80
36	1	232	G	N3-C4-N9	6.30	129.78	126.00
36	1	776	U	C5-C4-O4	6.30	129.68	125.90
36	5	2354	C	N1-C2-O2	-6.30	115.12	118.90
36	5	2938	G	C5-C6-O6	-6.30	124.82	128.60
36	5	1152	G	C4-C5-N7	6.29	113.32	110.80
36	1	3209	A	N1-C6-N6	6.29	122.38	118.60
39	L2	237	LEU	CA-CB-CG	-6.29	100.82	115.30
36	5	400	G	O5'-P-OP1	-6.29	100.04	105.70
36	5	1012	G	C6-C5-N7	6.29	134.18	130.40
36	5	1474	A	C8-N9-C4	6.29	108.32	105.80
36	1	73	C	N1-C2-O2	-6.29	115.12	118.90
36	5	2279	A	O4'-C1'-N9	6.29	113.23	108.20
36	5	2907	G	N3-C4-N9	-6.29	122.22	126.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	189	G	N1-C6-O6	-6.29	116.13	119.90
36	5	1126	G	C6-C5-N7	-6.29	126.63	130.40
36	5	2531	C	C2-N1-C1'	6.29	125.72	118.80
36	1	2261	G	N3-C4-N9	6.29	129.77	126.00
36	1	3242	G	C4-N9-C1'	-6.29	118.33	126.50
36	1	3306	U	N3-C2-O2	-6.29	117.80	122.20
37	7	6	C	C6-N1-C2	6.29	122.81	120.30
1	2	507	U	N1-C2-O2	6.29	127.20	122.80
1	2	863	A	C8-N9-C4	6.29	108.31	105.80
36	1	2930	A	C8-N9-C4	6.29	108.31	105.80
36	1	1349	G	N3-C4-C5	-6.28	125.46	128.60
36	1	2307	G	OP2-P-O3'	6.28	119.02	105.20
36	5	197	G	N3-C4-N9	6.28	129.77	126.00
36	1	1433	A	N7-C8-N9	6.28	116.94	113.80
36	5	718	G	C8-N9-C4	-6.28	103.89	106.40
36	1	650	C	O5'-P-OP1	-6.28	100.05	105.70
36	5	1744	G	C4-N9-C1'	6.28	134.66	126.50
36	5	2748	A	N1-C6-N6	6.28	122.37	118.60
1	2	1124	A	C2-N3-C4	-6.28	107.46	110.60
36	1	648	C	C5-C4-N4	-6.28	115.81	120.20
36	1	698	U	O4'-C1'-N1	6.28	113.22	108.20
36	1	1362	G	C8-N9-C4	6.28	108.91	106.40
1	6	360	A	N1-C6-N6	6.28	122.37	118.60
1	2	1340	U	N3-C2-O2	-6.28	117.81	122.20
1	2	1600	A	N1-C6-N6	6.28	122.36	118.60
36	1	2661	G	N1-C6-O6	6.28	123.67	119.90
36	5	1116	G	N3-C4-C5	-6.28	125.46	128.60
36	5	2160	G	C5-C6-O6	-6.28	124.83	128.60
36	1	2601	A	C8-N9-C4	6.27	108.31	105.80
36	1	1209	G	C4-C5-N7	-6.27	108.29	110.80
1	6	325	G	N9-C4-C5	6.27	107.91	105.40
36	5	2156	C	C2-N1-C1'	-6.27	111.90	118.80
36	1	1371	G	C5-C6-N1	-6.27	108.36	111.50
1	2	305	C	C6-N1-C2	-6.27	117.79	120.30
36	1	1377	G	C5-C6-O6	-6.27	124.84	128.60
36	5	648	C	N1-C2-O2	-6.27	115.14	118.90
36	5	2246	G	C8-N9-C4	-6.27	103.89	106.40
36	1	1381	A	N1-C6-N6	6.27	122.36	118.60
36	1	3362	A	C4-N9-C1'	6.27	137.58	126.30
36	5	2950	G	C4-C5-N7	6.27	113.31	110.80
36	5	1370	G	C5-C6-O6	6.27	132.36	128.60
1	6	542	A	C4-N9-C1'	6.26	137.57	126.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	6	1798	U	C2-N1-C1'	6.26	125.22	117.70
36	5	1443	G	C5-C6-O6	-6.26	124.84	128.60
36	5	2376	G	C8-N9-C4	-6.26	103.89	106.40
36	1	2812	C	C2-N3-C4	-6.26	116.77	119.90
36	5	1171	G	N3-C4-N9	-6.26	122.24	126.00
36	1	999	G	C5-C6-O6	-6.26	124.84	128.60
1	2	970	A	N9-C4-C5	-6.26	103.30	105.80
36	1	1581	C	C6-N1-C2	-6.26	117.80	120.30
36	1	2306	C	N1-C2-O2	6.26	122.65	118.90
1	6	558	U	N1-C2-O2	6.26	127.18	122.80
36	1	2376	G	C4-C5-N7	6.25	113.30	110.80
38	4	143	U	O5'-P-OP1	-6.25	100.07	105.70
36	5	1370	G	C4-C5-C6	6.25	122.55	118.80
1	6	861	U	O5'-P-OP2	-6.25	100.07	105.70
36	5	922	U	C5-C6-N1	-6.25	119.57	122.70
36	5	2385	G	N3-C4-C5	6.25	131.73	128.60
36	1	2616	C	N3-C2-O2	-6.25	117.53	121.90
36	5	562	C	C2-N1-C1'	6.25	125.67	118.80
36	1	2725	U	C6-N1-C2	6.25	124.75	121.00
36	5	1793	C	N1-C2-O2	-6.25	115.15	118.90
36	5	2145	A	O5'-P-OP1	-6.25	100.08	105.70
36	5	2991	A	C6-N1-C2	-6.25	114.85	118.60
37	3	75	G	O5'-P-OP1	-6.25	100.08	105.70
36	5	2727	A	N1-C6-N6	-6.25	114.85	118.60
1	2	416	A	C8-N9-C4	6.24	108.30	105.80
36	1	810	A	C5-C6-N1	6.24	120.82	117.70
36	1	2245	C	C6-N1-C2	-6.24	117.80	120.30
36	5	739	G	O5'-P-OP1	-6.24	100.08	105.70
36	1	62	A	O5'-P-OP2	-6.24	100.08	105.70
36	1	1886	A	N1-C6-N6	-6.24	114.86	118.60
1	6	337	G	O4'-C1'-N9	-6.24	103.21	108.20
1	6	904	G	N3-C4-N9	6.24	129.74	126.00
36	5	2626	A	C2-N3-C4	-6.24	107.48	110.60
1	2	16	G	C4-N9-C1'	6.24	134.61	126.50
36	5	2957	G	N3-C4-N9	-6.24	122.26	126.00
1	2	1170	G	C8-N9-C1'	-6.24	118.89	127.00
36	1	3099	C	O5'-P-OP1	-6.24	100.09	105.70
1	2	1739	C	C6-N1-C2	6.23	122.79	120.30
36	1	983	A	N1-C2-N3	6.23	132.42	129.30
1	2	6	G	C8-N9-C4	-6.23	103.91	106.40
36	1	3181	C	N3-C2-O2	-6.23	117.54	121.90
36	1	3362	A	C4-C5-C6	6.23	120.11	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	2370	G	C8-N9-C4	-6.23	103.91	106.40
1	6	779	U	N3-C2-O2	-6.23	117.84	122.20
36	1	1512	U	O5'-P-OP2	-6.22	100.10	105.70
36	1	2620	G	C4-N9-C1'	-6.22	118.41	126.50
1	6	1130	G	N7-C8-N9	-6.22	109.99	113.10
36	5	2662	G	C4-N9-C1'	6.22	134.59	126.50
36	1	924	G	N3-C2-N2	6.22	124.25	119.90
1	6	427	C	N1-C2-O2	-6.22	115.17	118.90
36	1	366	A	N1-C6-N6	-6.22	114.87	118.60
36	1	2385	G	C4-N9-C1'	-6.22	118.41	126.50
1	6	1634	C	N1-C2-O2	6.22	122.63	118.90
36	1	676	G	C4-C5-C6	6.22	122.53	118.80
36	1	718	G	C4-N9-C1'	6.22	134.59	126.50
36	1	933	A	C6-C5-N7	-6.22	127.95	132.30
36	1	1585	C	C5-C4-N4	-6.22	115.85	120.20
36	5	3144	G	N3-C4-N9	6.22	129.73	126.00
36	5	2947	G	C4-C5-N7	6.22	113.29	110.80
36	1	620	U	N3-C2-O2	-6.21	117.85	122.20
38	4	53	A	C6-C5-N7	6.21	136.65	132.30
36	1	1149	G	C5-C6-N1	-6.21	108.39	111.50
36	1	2283	G	C4-C5-N7	6.21	113.28	110.80
36	1	3182	G	C6-C5-N7	-6.21	126.67	130.40
36	5	1661	G	N1-C6-O6	6.21	123.63	119.90
1	2	1082	C	N3-C2-O2	-6.21	117.55	121.90
36	1	369	A	N9-C4-C5	6.21	108.28	105.80
36	1	3375	A	N9-C4-C5	6.21	108.28	105.80
36	1	24	G	C4-C5-C6	6.21	122.52	118.80
36	1	2342	U	C5-C6-N1	-6.21	119.60	122.70
36	1	714	G	N1-C6-O6	6.20	123.62	119.90
1	6	444	C	C6-N1-C2	6.20	122.78	120.30
36	5	1294	A	N1-C6-N6	-6.20	114.88	118.60
36	1	2194	G	C6-C5-N7	-6.20	126.68	130.40
36	5	1915	A	C8-N9-C4	6.20	108.28	105.80
36	1	698	U	C5-C4-O4	6.20	129.62	125.90
36	1	1118	C	C6-N1-C2	-6.20	117.82	120.30
36	1	1145	G	O5'-P-OP2	-6.20	100.12	105.70
36	1	1377	G	C4-C5-N7	6.20	113.28	110.80
36	1	3302	U	C5-C6-N1	-6.20	119.60	122.70
37	3	111	U	O5'-P-OP1	-6.20	100.12	105.70
1	6	542	A	C4-C5-C6	6.20	120.10	117.00
1	6	1137	A	C8-N9-C4	6.20	108.28	105.80
36	5	46	U	N3-C2-O2	6.20	126.54	122.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	715	A	C4-C5-N7	-6.20	107.60	110.70
36	5	1133	A	C5-C6-N6	-6.20	118.74	123.70
36	5	1376	C	C6-N1-C2	-6.20	117.82	120.30
36	1	341	G	C8-N9-C4	-6.20	103.92	106.40
36	1	1899	G	C5-C6-O6	6.20	132.32	128.60
36	1	2409	G	N3-C4-N9	6.20	129.72	126.00
1	6	619	A	C8-N9-C4	-6.20	103.32	105.80
36	5	888	A	C5-C6-N6	-6.20	118.74	123.70
36	1	2662	G	C4-C5-N7	6.20	113.28	110.80
1	6	425	A	C8-N9-C4	-6.20	103.32	105.80
36	5	2400	G	C5-C6-N1	-6.20	108.40	111.50
36	1	1125	U	O5'-P-OP1	-6.19	100.12	105.70
36	1	1178	G	N1-C2-N2	-6.19	110.62	116.20
36	5	1381	A	C2-N3-C4	-6.19	107.50	110.60
36	1	2416	U	C6-N1-C2	-6.19	117.28	121.00
36	5	904	A	C5-C6-N1	6.19	120.80	117.70
36	5	1317	A	C5-N7-C8	-6.19	100.80	103.90
36	5	1370	G	N9-C4-C5	6.19	107.88	105.40
36	5	2954	U	C2-N1-C1'	6.19	125.13	117.70
1	2	453	U	N1-C2-O2	6.19	127.13	122.80
36	1	1429	G	N3-C4-C5	-6.19	125.50	128.60
1	6	512	A	N1-C6-N6	6.19	122.31	118.60
36	5	610	G	N3-C4-C5	-6.19	125.50	128.60
36	5	706	A	C8-N9-C4	6.19	108.28	105.80
36	5	1879	A	O5'-P-OP1	6.19	118.13	110.70
36	5	2617	U	N3-C4-C5	-6.19	110.89	114.60
36	5	3028	G	N3-C2-N2	6.19	124.23	119.90
37	7	79	A	N1-C6-N6	6.19	122.31	118.60
36	5	410	U	N1-C2-O2	-6.19	118.47	122.80
36	1	645	A	C5-C6-N1	6.19	120.79	117.70
36	1	2675	C	N1-C2-O2	6.19	122.61	118.90
36	1	1901	A	N1-C6-N6	-6.18	114.89	118.60
1	6	402	C	N1-C2-O2	6.18	122.61	118.90
36	5	2727	A	O5'-P-OP2	-6.18	100.13	105.70
36	5	3294	A	N9-C4-C5	6.18	108.27	105.80
1	2	354	C	C6-N1-C2	-6.18	117.83	120.30
36	5	2283	G	N3-C4-C5	6.18	131.69	128.60
36	5	3143	C	N1-C2-O2	-6.18	115.19	118.90
1	2	55	A	C8-N9-C4	-6.18	103.33	105.80
1	2	1297	G	C8-N9-C4	6.18	108.87	106.40
36	1	1852	G	N1-C6-O6	6.18	123.61	119.90
1	6	1672	G	N1-C2-N2	-6.18	110.64	116.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	2978	U	C4-C5-C6	6.18	123.41	119.70
37	7	88	G	N3-C4-C5	-6.18	125.51	128.60
1	6	158	U	OP1-P-O3'	6.18	118.79	105.20
36	5	2891	U	N3-C4-C5	6.18	118.31	114.60
36	1	2151	C	C6-N1-C2	6.18	122.77	120.30
36	1	2662	G	C6-C5-N7	-6.18	126.69	130.40
36	5	1208	U	C5-C4-O4	6.18	129.60	125.90
37	7	5	G	N7-C8-N9	-6.18	110.01	113.10
36	1	3074	G	N3-C4-C5	6.17	131.69	128.60
1	6	467	G	N3-C4-N9	6.17	129.70	126.00
1	6	1085	G	N3-C4-N9	-6.17	122.30	126.00
36	1	590	G	C4-C5-N7	6.17	113.27	110.80
1	2	992	A	C2-N3-C4	-6.17	107.51	110.60
36	5	2702	A	N1-C2-N3	6.17	132.38	129.30
36	5	2817	A	N1-C6-N6	6.17	122.30	118.60
36	5	3154	C	C6-N1-C2	-6.17	117.83	120.30
36	1	2758	A	C8-N9-C4	6.17	108.27	105.80
36	5	3053	G	N1-C6-O6	6.17	123.60	119.90
1	6	435	C	C6-N1-C2	6.17	122.77	120.30
36	5	1377	G	N1-C6-O6	-6.17	116.20	119.90
36	5	3108	G	C4-C5-C6	6.17	122.50	118.80
1	6	337	G	C8-N9-C1'	-6.17	118.98	127.00
36	5	929	A	O5'-P-OP2	-6.17	100.15	105.70
36	5	1517	G	N3-C4-N9	-6.17	122.30	126.00
36	5	2690	G	C4-N9-C1'	6.17	134.52	126.50
1	2	765	G	O4'-C1'-N9	-6.16	103.27	108.20
36	1	2788	C	N3-C2-O2	6.16	126.21	121.90
36	1	2968	G	C5-C6-O6	-6.16	124.90	128.60
36	5	973	A	N1-C6-N6	6.16	122.30	118.60
37	7	100	C	C6-N1-C2	6.16	122.77	120.30
36	5	804	C	N3-C2-O2	6.16	126.21	121.90
36	5	1519	G	N1-C6-O6	6.16	123.60	119.90
36	5	1347	U	O5'-P-OP2	-6.16	100.16	105.70
36	5	1373	A	C5-C6-N6	-6.16	118.77	123.70
36	5	3094	A	C8-N9-C4	6.16	108.26	105.80
1	2	1600	A	C2-N3-C4	-6.16	107.52	110.60
36	1	360	G	C4-C5-N7	6.16	113.26	110.80
36	5	2693	C	C6-N1-C2	6.16	122.76	120.30
36	5	2812	C	N3-C4-C5	6.16	124.36	121.90
36	5	360	G	C4-C5-C6	6.16	122.49	118.80
39	12	237	LEU	CA-CB-CG	-6.16	101.14	115.30
1	2	1489	U	N1-C2-O2	6.16	127.11	122.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	360	G	C6-C5-N7	-6.16	126.71	130.40
36	1	1660	C	C6-N1-C2	-6.16	117.84	120.30
36	5	1001	G	O5'-P-OP1	-6.16	100.16	105.70
36	5	2754	G	N1-C2-N2	-6.16	110.66	116.20
1	2	1560	U	C5-C4-O4	6.15	129.59	125.90
36	1	2787	G	C4-C5-N7	6.15	113.26	110.80
1	2	1004	U	N3-C2-O2	-6.15	117.89	122.20
36	1	1736	G	N1-C6-O6	6.15	123.59	119.90
36	1	2341	A	O5'-P-OP2	-6.15	100.16	105.70
36	5	965	A	N9-C4-C5	6.15	108.26	105.80
36	1	1507	G	C6-C5-N7	-6.15	126.71	130.40
36	1	1904	C	C2-N1-C1'	6.15	125.57	118.80
36	1	1079	A	C8-N9-C4	6.15	108.26	105.80
36	1	56	G	C5-C6-N1	6.15	114.57	111.50
36	1	229	G	O5'-P-OP2	6.15	118.08	110.70
36	1	1365	G	C8-N9-C4	-6.15	103.94	106.40
36	1	2166	A	N1-C6-N6	6.15	122.29	118.60
36	1	2306	C	C2-N1-C1'	6.15	125.56	118.80
36	1	2608	G	N1-C6-O6	6.15	123.59	119.90
36	1	3133	C	C6-N1-C2	-6.15	117.84	120.30
1	6	1700	C	C2-N1-C1'	6.15	125.56	118.80
36	5	92	G	C4-C5-N7	-6.15	108.34	110.80
36	5	1894	U	C5-C6-N1	-6.15	119.63	122.70
36	1	1807	G	N3-C4-C5	-6.14	125.53	128.60
36	1	2276	G	C6-C5-N7	-6.14	126.71	130.40
1	6	1600	A	N9-C1'-C2'	6.14	121.99	114.00
36	1	51	A	C4-C5-N7	6.14	113.77	110.70
36	5	2849	C	N3-C2-O2	6.14	126.20	121.90
36	5	2954	U	O4'-C1'-N1	6.14	113.11	108.20
36	1	968	G	C4-N9-C1'	6.14	134.48	126.50
36	5	1321	G	C5-N7-C8	-6.14	101.23	104.30
1	2	553	G	N9-C4-C5	-6.14	102.94	105.40
1	2	554	C	N3-C4-C5	-6.14	119.44	121.90
36	1	229	G	N1-C2-N2	6.14	121.72	116.20
36	1	1224	C	C6-N1-C2	-6.14	117.84	120.30
36	5	2961	G	O5'-P-OP2	-6.14	100.17	105.70
36	5	3180	A	N1-C2-N3	6.14	132.37	129.30
36	5	2624	G	N7-C8-N9	6.14	116.17	113.10
36	5	2772	C	P-O3'-C3'	6.14	127.07	119.70
36	5	3362	A	C2-N3-C4	-6.14	107.53	110.60
36	1	2870	C	N3-C4-C5	6.14	124.35	121.90
36	5	1889	G	C4-C5-N7	6.14	113.25	110.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
33	e1	100	LEU	CA-CB-CG	6.13	129.41	115.30
36	1	1929	G	N9-C4-C5	-6.13	102.95	105.40
1	6	1031	U	N3-C2-O2	6.13	126.49	122.20
36	1	1412	G	C6-C5-N7	-6.13	126.72	130.40
1	6	453	U	N3-C2-O2	-6.13	117.91	122.20
36	1	2836	C	N3-C4-C5	-6.13	119.45	121.90
36	5	424	G	C6-C5-N7	-6.13	126.72	130.40
36	5	1016	C	O5'-P-OP1	-6.13	100.19	105.70
36	1	716	A	C4-C5-N7	6.12	113.76	110.70
36	1	1601	U	C5-C4-O4	6.12	129.57	125.90
36	1	1793	C	C6-N1-C2	6.12	122.75	120.30
36	1	3172	A	N9-C4-C5	6.12	108.25	105.80
1	6	1119	G	C8-N9-C4	-6.12	103.95	106.40
1	6	1631	A	O5'-P-OP2	-6.12	100.19	105.70
36	5	522	A	C6-C5-N7	-6.12	128.01	132.30
36	5	826	G	C5-C6-O6	-6.12	124.92	128.60
36	5	1386	A	C5-C6-N1	-6.12	114.64	117.70
36	1	642	U	N3-C4-C5	-6.12	110.93	114.60
36	1	679	U	N3-C4-O4	-6.12	115.11	119.40
1	6	1600	A	N1-C2-N3	6.12	132.36	129.30
36	5	681	U	C2-N1-C1'	6.12	125.05	117.70
36	5	2623	G	C8-N9-C4	6.12	108.85	106.40
36	5	3080	G	N1-C6-O6	6.12	123.57	119.90
36	1	633	C	C4-C5-C6	6.12	120.46	117.40
36	5	511	G	N3-C2-N2	-6.12	115.62	119.90
36	1	3057	U	N3-C2-O2	-6.12	117.92	122.20
36	5	30	G	O5'-P-OP2	6.12	118.04	110.70
36	5	1435	A	C6-N1-C2	-6.12	114.93	118.60
36	5	2142	A	C5-C6-N1	6.12	120.76	117.70
38	8	84	C	C6-N1-C2	-6.12	117.85	120.30
36	5	2661	G	N3-C4-N9	6.12	129.67	126.00
36	1	953	G	N9-C4-C5	-6.12	102.95	105.40
36	1	2374	C	C4-C5-C6	6.12	120.46	117.40
1	6	1130	G	N3-C4-C5	6.11	131.66	128.60
36	5	883	A	N9-C4-C5	6.11	108.25	105.80
1	2	1198	G	C8-N9-C4	-6.11	103.95	106.40
36	5	907	G	C5-C6-N1	6.11	114.56	111.50
36	5	3245	A	N1-C2-N3	6.11	132.36	129.30
36	1	1792	C	C6-N1-C1'	6.11	128.13	120.80
1	6	402	C	O5'-P-OP2	-6.11	100.20	105.70
36	5	2968	G	O5'-P-OP1	-6.11	100.20	105.70
1	2	1137	A	C2-N3-C4	-6.11	107.55	110.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	2	1297	G	N1-C6-O6	6.11	123.56	119.90
36	1	3089	C	C6-N1-C2	-6.11	117.86	120.30
36	5	1384	U	C6-N1-C2	-6.11	117.34	121.00
36	1	28	C	N3-C4-C5	6.11	124.34	121.90
36	1	519	A	N1-C6-N6	6.11	122.26	118.60
36	1	1171	G	N9-C4-C5	6.11	107.84	105.40
36	1	3098	G	O5'-P-OP1	-6.11	100.20	105.70
36	5	2968	G	C8-N9-C1'	-6.11	119.06	127.00
36	1	3133	C	C5-C6-N1	6.10	124.05	121.00
36	5	1790	G	N1-C6-O6	6.10	123.56	119.90
36	5	967	A	N7-C8-N9	6.10	116.85	113.80
36	5	1117	G	N1-C6-O6	6.10	123.56	119.90
36	1	1116	G	N9-C4-C5	6.10	107.84	105.40
36	1	3217	C	C6-N1-C1'	-6.10	113.48	120.80
1	6	603	U	N3-C2-O2	6.10	126.47	122.20
36	1	1141	C	C5-C6-N1	-6.10	117.95	121.00
36	1	2143	A	C5-N7-C8	-6.10	100.85	103.90
1	6	171	A	O5'-P-OP2	-6.10	100.21	105.70
1	6	553	G	C5-N7-C8	-6.10	101.25	104.30
36	5	1193	A	N1-C2-N3	6.10	132.35	129.30
36	5	2413	A	N9-C4-C5	-6.10	103.36	105.80
36	5	3298	C	O5'-P-OP1	-6.10	100.21	105.70
36	1	648	C	C6-N1-C2	-6.09	117.86	120.30
1	6	1128	C	O5'-P-OP1	-6.09	100.22	105.70
38	8	26	U	C6-N1-C2	-6.09	117.34	121.00
1	2	425	A	C4-C5-N7	6.09	113.75	110.70
36	1	1100	U	N3-C4-C5	6.09	118.26	114.60
36	5	2411	U	N3-C4-O4	-6.09	115.14	119.40
36	1	625	G	N1-C6-O6	6.09	123.56	119.90
37	7	13	A	C5-C6-N1	6.09	120.75	117.70
1	2	144	U	N3-C2-O2	-6.09	117.94	122.20
36	1	699	A	N3-C4-N9	-6.09	122.53	127.40
36	1	2625	C	N1-C2-O2	-6.09	115.25	118.90
36	5	3095	U	N1-C2-N3	6.09	118.55	114.90
36	1	2177	G	N3-C4-N9	6.09	129.65	126.00
36	5	25	U	C5-C6-N1	-6.09	119.66	122.70
36	5	2282	U	N1-C2-O2	-6.09	118.54	122.80
36	1	876	A	N1-C2-N3	-6.09	126.26	129.30
36	1	1362	G	OP2-P-O3'	6.09	118.59	105.20
36	1	1445	U	N1-C2-N3	6.09	118.55	114.90
1	2	545	A	OP1-P-O3'	6.08	118.59	105.20
36	1	3128	G	N3-C4-N9	6.08	129.65	126.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	6	1130	G	C4-N9-C1'	-6.08	118.59	126.50
36	5	3208	G	N9-C4-C5	-6.08	102.97	105.40
36	1	1371	G	N1-C6-O6	6.08	123.55	119.90
36	1	2774	C	C2-N1-C1'	6.08	125.49	118.80
1	6	117	U	C6-N1-C2	-6.08	117.35	121.00
1	2	1212	G	C4-C5-N7	6.08	113.23	110.80
36	5	1909	A	C2-N3-C4	-6.08	107.56	110.60
11	S9	93	LEU	CA-CB-CG	6.08	129.28	115.30
36	1	2273	G	C4-N9-C1'	-6.08	118.60	126.50
36	1	2860	U	C6-N1-C1'	-6.08	112.69	121.20
1	6	610	G	C4-N9-C1'	6.08	134.40	126.50
36	5	197	G	C4-N9-C1'	6.08	134.40	126.50
36	5	1379	G	N9-C4-C5	-6.08	102.97	105.40
37	7	1	G	C8-N9-C1'	-6.08	119.10	127.00
1	2	1539	G	C8-N9-C1'	-6.08	119.10	127.00
36	1	2415	C	OP1-P-O3'	6.08	118.57	105.20
36	5	650	C	OP2-P-O3'	6.08	118.57	105.20
36	5	1208	U	N1-C2-O2	6.08	127.05	122.80
36	5	2818	U	N3-C2-O2	6.08	126.45	122.20
1	6	408	C	C6-N1-C2	-6.08	117.87	120.30
36	5	2167	A	C2-N3-C4	6.08	113.64	110.60
36	5	3362	A	C5-N7-C8	-6.08	100.86	103.90
1	2	380	U	N1-C2-O2	6.07	127.05	122.80
36	1	2648	G	N9-C1'-C2'	-6.07	105.32	112.00
36	5	1551	C	C6-N1-C2	-6.07	117.87	120.30
36	5	425	G	O5'-P-OP1	6.07	117.99	110.70
36	1	335	G	O5'-P-OP2	6.07	117.99	110.70
1	6	1698	G	P-O3'-C3'	6.07	126.98	119.70
36	1	2372	A	N1-C6-N6	6.07	122.24	118.60
36	1	2978	U	C5-C6-N1	-6.07	119.67	122.70
1	6	1765	A	N1-C6-N6	-6.07	114.96	118.60
36	5	709	A	N9-C4-C5	-6.07	103.37	105.80
41	14	20	LEU	CA-CB-CG	-6.07	101.34	115.30
36	1	2852	C	C5-C4-N4	-6.07	115.95	120.20
1	2	137	U	O5'-P-OP1	-6.06	100.24	105.70
36	5	360	G	C5-C6-N1	-6.06	108.47	111.50
36	5	2295	A	C5-C6-N1	6.06	120.73	117.70
36	1	953	G	OP1-P-O3'	6.06	118.54	105.20
36	5	2964	G	N1-C6-O6	-6.06	116.26	119.90
1	2	1212	G	C6-C5-N7	-6.06	126.76	130.40
36	1	1325	U	C6-N1-C2	6.06	124.64	121.00
1	6	1620	C	C6-N1-C2	-6.06	117.88	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	725	G	C4-N9-C1'	6.06	134.38	126.50
36	5	887	G	N1-C6-O6	6.06	123.53	119.90
36	1	1074	U	O5'-P-OP1	-6.06	100.25	105.70
36	5	998	A	O5'-P-OP1	-6.06	100.25	105.70
36	5	2145	A	C4-C5-C6	6.06	120.03	117.00
36	1	965	A	C5-N7-C8	-6.06	100.87	103.90
1	6	176	C	N1-C2-O2	6.05	122.53	118.90
36	5	3084	C	C6-N1-C2	6.05	122.72	120.30
36	5	3252	G	C8-N9-C4	6.05	108.82	106.40
36	1	2610	G	C2-N3-C4	-6.05	108.87	111.90
36	5	2843	U	N1-C2-O2	6.05	127.04	122.80
36	1	947	G	C4-C5-C6	6.05	122.43	118.80
36	1	1419	A	O5'-P-OP2	-6.05	100.25	105.70
38	4	13	A	OP1-P-OP2	6.05	128.68	119.60
38	4	26	U	N1-C2-O2	6.05	127.04	122.80
36	1	283	G	C8-N9-C4	-6.05	103.98	106.40
1	6	85	A	C8-N9-C4	-6.05	103.38	105.80
36	5	2182	A	OP1-P-O3'	6.05	118.50	105.20
36	5	2794	G	C8-N9-C4	6.05	108.82	106.40
1	2	346	G	N9-C4-C5	-6.04	102.98	105.40
36	1	2609	A	O5'-P-OP1	6.04	117.95	110.70
36	5	1744	G	N3-C4-N9	6.04	129.63	126.00
36	5	2295	A	N1-C6-N6	6.04	122.23	118.60
36	1	2409	G	C6-N1-C2	-6.04	121.47	125.10
36	1	2513	U	P-O3'-C3'	6.04	126.95	119.70
36	1	3277	U	N3-C2-O2	-6.04	117.97	122.20
36	5	2391	G	C2-N3-C4	6.04	114.92	111.90
36	5	2943	G	C4-C5-C6	6.04	122.42	118.80
37	7	1	G	N3-C2-N2	6.04	124.13	119.90
36	1	3093	C	N3-C4-C5	6.04	124.32	121.90
36	1	2695	A	O4'-C1'-N9	6.04	113.03	108.20
36	5	419	G	O5'-P-OP1	6.04	117.95	110.70
36	5	637	C	O5'-P-OP1	-6.04	100.27	105.70
36	5	1443	G	N9-C4-C5	-6.04	102.98	105.40
36	5	2775	U	C2-N1-C1'	-6.04	110.45	117.70
36	1	2836	C	N3-C2-O2	-6.04	117.67	121.90
36	5	1182	A	N1-C6-N6	6.04	122.22	118.60
36	5	1450	G	N1-C6-O6	6.04	123.52	119.90
36	1	1129	A	C5-C6-N6	-6.04	118.87	123.70
36	1	2991	A	C2-N3-C4	-6.04	107.58	110.60
36	1	1406	A	N1-C6-N6	6.03	122.22	118.60
36	1	2279	A	N9-C4-C5	-6.03	103.39	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	1004	U	C6-N1-C2	-6.03	117.38	121.00
36	1	2774	C	N3-C4-N4	6.03	122.22	118.00
1	6	1672	G	N3-C4-C5	-6.03	125.58	128.60
36	5	809	G	C8-N9-C4	6.03	108.81	106.40
36	5	3309	G	C8-N9-C1'	-6.03	119.16	127.00
1	2	1541	G	C4-N9-C1'	6.03	134.34	126.50
36	1	2949	U	C6-N1-C2	6.03	124.62	121.00
36	1	1545	A	N1-C2-N3	6.03	132.31	129.30
36	1	29	C	C5-C4-N4	-6.03	115.98	120.20
36	5	43	A	O4'-C1'-N9	6.03	113.02	108.20
36	5	71	A	O5'-P-OP1	-6.03	100.28	105.70
36	1	3344	A	C4-N9-C1'	6.02	137.14	126.30
36	5	718	G	N3-C4-N9	6.02	129.61	126.00
36	5	2940	A	N1-C6-N6	-6.02	114.99	118.60
36	1	716	A	C8-N9-C4	6.02	108.21	105.80
36	1	718	G	C6-C5-N7	-6.02	126.79	130.40
36	1	3312	U	C5-C6-N1	-6.02	119.69	122.70
1	6	1619	C	C6-N1-C2	-6.02	117.89	120.30
1	6	1774	G	N1-C6-O6	-6.02	116.29	119.90
36	5	2160	G	N3-C4-N9	6.02	129.61	126.00
36	5	1857	C	C2-N1-C1'	6.02	125.42	118.80
36	5	3116	G	C4-C5-N7	6.02	113.21	110.80
37	7	26	C	C4-C5-C6	6.02	120.41	117.40
36	1	1451	C	N3-C4-C5	6.02	124.31	121.90
36	5	2403	G	N7-C8-N9	-6.02	110.09	113.10
36	5	3116	G	N9-C4-C5	-6.02	102.99	105.40
1	2	704	C	O4'-C1'-N1	6.02	113.01	108.20
36	1	921	A	C6-N1-C2	-6.02	114.99	118.60
36	5	1497	C	C5-C4-N4	-6.02	115.99	120.20
36	5	1592	G	C4-C5-N7	-6.02	108.39	110.80
36	1	25	U	N1-C2-O2	-6.02	118.59	122.80
36	1	2414	G	C5-C6-N1	-6.02	108.49	111.50
1	6	557	G	N3-C4-C5	-6.02	125.59	128.60
1	2	1539	G	N3-C4-N9	6.01	129.61	126.00
36	1	99	A	O4'-C1'-N9	6.01	113.01	108.20
36	1	658	G	C4-N9-C1'	6.01	134.32	126.50
36	1	1076	C	C6-N1-C2	6.01	122.71	120.30
36	5	1882	G	C4-C5-N7	6.01	113.20	110.80
1	2	1363	U	N3-C2-O2	-6.01	117.99	122.20
36	1	2787	G	C5-C6-O6	-6.01	124.99	128.60
36	5	578	A	O5'-P-OP2	6.01	117.92	110.70
1	2	1241	G	C4-C5-N7	6.01	113.20	110.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	2411	U	C5-C6-N1	-6.01	119.69	122.70
36	1	197	G	C4-C5-N7	6.01	113.20	110.80
36	1	651	G	N9-C4-C5	-6.01	103.00	105.40
36	5	1582	C	C6-N1-C2	-6.01	117.90	120.30
36	5	2680	A	N1-C6-N6	-6.01	114.99	118.60
36	5	1081	U	N1-C2-O2	6.01	127.01	122.80
36	5	2145	A	C6-C5-N7	-6.01	128.09	132.30
36	5	2818	U	C4-C5-C6	-6.01	116.09	119.70
36	1	1522	U	C6-N1-C2	6.01	124.60	121.00
1	2	507	U	N3-C2-O2	-6.00	118.00	122.20
1	6	901	G	N1-C6-O6	6.00	123.50	119.90
36	1	342	A	O5'-P-OP2	-6.00	100.30	105.70
36	1	816	A	C5-C6-N1	6.00	120.70	117.70
36	1	1863	G	C4-C5-N7	6.00	113.20	110.80
36	5	2351	U	N1-C2-N3	6.00	118.50	114.90
36	1	3370	A	O5'-P-OP2	-6.00	100.30	105.70
1	6	1389	C	N3-C2-O2	-6.00	117.70	121.90
1	6	163	G	N3-C4-N9	-6.00	122.40	126.00
1	2	1541	G	N3-C4-N9	6.00	129.60	126.00
36	1	2993	G	OP1-P-OP2	6.00	128.59	119.60
1	6	1389	C	N1-C2-O2	6.00	122.50	118.90
36	5	437	G	N3-C4-N9	-6.00	122.40	126.00
36	5	3080	G	C5-C6-O6	-6.00	125.00	128.60
36	5	1374	G	O5'-P-OP2	-6.00	100.30	105.70
36	1	75	G	N1-C6-O6	5.99	123.50	119.90
36	1	285	A	C8-N9-C4	5.99	108.20	105.80
36	1	2639	G	N1-C6-O6	5.99	123.50	119.90
1	6	937	C	N3-C2-O2	-5.99	117.70	121.90
1	6	1606	C	N1-C2-O2	5.99	122.50	118.90
36	5	816	A	C2-N3-C4	5.99	113.60	110.60
36	5	2400	G	C2-N3-C4	-5.99	108.90	111.90
36	5	2615	G	C2-N3-C4	-5.99	108.90	111.90
36	1	1416	C	C5-C6-N1	-5.99	118.00	121.00
36	1	2802	A	N9-C4-C5	5.99	108.20	105.80
1	2	1600	A	C5-C6-N1	-5.99	114.70	117.70
36	1	342	A	N9-C4-C5	5.99	108.20	105.80
36	1	2658	G	C4-C5-N7	-5.99	108.41	110.80
36	5	651	G	N3-C4-C5	-5.99	125.61	128.60
36	5	1897	G	C2-N3-C4	-5.99	108.91	111.90
36	5	2968	G	C4-N9-C1'	5.99	134.29	126.50
36	1	189	G	N1-C6-O6	-5.99	116.31	119.90
36	1	1323	G	C5-N7-C8	-5.99	101.31	104.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2261	G	N3-C4-C5	-5.99	125.61	128.60
36	1	2283	G	C5-N7-C8	-5.99	101.31	104.30
36	1	2643	A	C2-N3-C4	-5.99	107.61	110.60
36	1	2647	A	C6-N1-C2	-5.99	115.01	118.60
36	5	660	A	N1-C6-N6	-5.99	115.01	118.60
36	5	1182	A	C4-C5-N7	5.99	113.69	110.70
36	5	1897	G	N3-C2-N2	-5.99	115.71	119.90
36	1	1020	G	C5-C6-O6	-5.98	125.01	128.60
36	1	1429	G	N1-C2-N2	-5.98	110.81	116.20
36	1	965	A	C2-N3-C4	-5.98	107.61	110.60
36	1	1186	G	N3-C4-N9	5.98	129.59	126.00
1	6	54	C	C6-N1-C2	-5.98	117.91	120.30
36	5	330	G	N1-C6-O6	5.98	123.49	119.90
36	5	1128	U	C5-C6-N1	-5.98	119.71	122.70
36	5	2403	G	C8-N9-C1'	-5.98	119.22	127.00
36	1	1369	A	C2-N3-C4	-5.98	107.61	110.60
36	5	725	G	C8-N9-C1'	-5.98	119.23	127.00
36	5	931	C	N3-C4-N4	-5.98	113.81	118.00
38	8	32	C	C6-N1-C2	5.98	122.69	120.30
1	2	1733	C	N3-C4-N4	5.98	122.19	118.00
1	2	348	U	O5'-P-OP2	-5.98	100.32	105.70
36	1	2647	A	N3-C4-C5	-5.98	122.62	126.80
1	6	781	U	C2-N1-C1'	5.98	124.87	117.70
36	5	2368	A	C8-N9-C4	-5.98	103.41	105.80
36	1	3312	U	C6-N1-C2	5.98	124.59	121.00
36	1	1851	G	C5-N7-C8	-5.97	101.31	104.30
36	1	1906	G	N3-C2-N2	-5.97	115.72	119.90
36	1	1908	A	N1-C6-N6	5.97	122.19	118.60
36	1	2126	A	C8-N9-C4	5.97	108.19	105.80
36	5	1885	U	C2-N1-C1'	-5.97	110.53	117.70
36	5	1917	C	C5-C6-N1	-5.97	118.01	121.00
36	1	1890	U	N3-C2-O2	5.97	126.38	122.20
36	1	2688	U	N1-C2-N3	-5.97	111.32	114.90
36	5	659	G	N3-C2-N2	5.97	124.08	119.90
36	5	923	C	C5-C4-N4	-5.97	116.02	120.20
36	5	2255	A	O5'-P-OP1	-5.97	100.33	105.70
36	5	1148	G	C8-N9-C4	5.97	108.79	106.40
36	1	558	U	N1-C2-O2	5.97	126.98	122.80
36	1	3266	G	N9-C4-C5	5.97	107.79	105.40
1	6	1731	A	N1-C6-N6	-5.96	115.02	118.60
36	5	1923	C	O5'-P-OP1	-5.96	100.33	105.70
36	5	2661	G	C5-C6-O6	-5.96	125.02	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	2943	G	C5-C6-N1	-5.96	108.52	111.50
36	5	1404	G	N1-C6-O6	5.96	123.48	119.90
1	6	426	G	C4-N9-C1'	5.96	134.25	126.50
36	5	1406	A	C6-N1-C2	-5.96	115.02	118.60
1	2	1539	G	N3-C4-C5	-5.96	125.62	128.60
36	5	1212	A	N1-C6-N6	5.96	122.18	118.60
36	5	1347	U	N3-C2-O2	5.96	126.37	122.20
1	2	704	C	N1-C2-O2	5.96	122.47	118.90
36	1	2148	U	O5'-P-OP1	-5.96	100.34	105.70
36	5	672	A	N1-C6-N6	-5.96	115.02	118.60
36	1	1442	U	N3-C4-O4	5.96	123.57	119.40
36	1	1716	U	P-O3'-C3'	5.96	126.85	119.70
36	5	1592	G	C5-C6-N1	-5.96	108.52	111.50
36	5	3245	A	O4'-C1'-N9	5.96	112.97	108.20
36	1	123	A	C4-C5-N7	5.96	113.68	110.70
36	1	2836	C	N1-C2-N3	5.96	123.37	119.20
1	6	51	A	C8-N9-C4	5.95	108.18	105.80
1	6	938	G	N3-C4-C5	-5.95	125.62	128.60
1	6	1058	U	OP1-P-O3'	5.95	118.30	105.20
36	5	210	U	C5-C4-O4	5.95	129.47	125.90
36	5	2955	U	OP2-P-O3'	5.95	118.30	105.20
38	4	20	U	C5-C6-N1	-5.95	119.72	122.70
36	5	2608	G	OP2-P-O3'	5.95	118.29	105.20
36	5	2726	C	N3-C4-N4	-5.95	113.83	118.00
36	1	1393	A	N1-C6-N6	-5.95	115.03	118.60
36	1	3277	U	C5-C4-O4	5.95	129.47	125.90
36	1	936	A	C8-N9-C4	5.95	108.18	105.80
36	1	1522	U	C5-C6-N1	-5.95	119.73	122.70
1	6	1097	U	P-O3'-C3'	5.95	126.84	119.70
36	5	2914	G	N3-C4-C5	-5.95	125.63	128.60
1	2	1733	C	N3-C2-O2	5.95	126.06	121.90
1	2	87	C	C6-N1-C2	-5.95	117.92	120.30
36	1	2200	U	C4-C5-C6	5.95	123.27	119.70
36	5	119	U	N3-C4-O4	-5.95	115.24	119.40
36	5	197	G	C8-N9-C1'	-5.95	119.27	127.00
36	5	438	A	C8-N9-C4	5.95	108.18	105.80
37	3	80	G	C6-C5-N7	-5.94	126.83	130.40
36	5	2760	C	C5-C4-N4	-5.94	116.04	120.20
1	2	1761	U	P-O3'-C3'	5.94	126.83	119.70
36	1	693	A	N1-C6-N6	5.94	122.17	118.60
36	1	1851	G	C5-C6-O6	-5.94	125.03	128.60
36	5	2863	G	N1-C6-O6	-5.94	116.33	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	2888	U	C5-C4-O4	-5.94	122.33	125.90
36	1	895	A	N1-C6-N6	5.94	122.16	118.60
36	1	1306	G	C2-N3-C4	-5.94	108.93	111.90
36	1	3310	A	C8-N9-C4	5.94	108.18	105.80
37	3	94	C	N3-C2-O2	5.94	126.06	121.90
36	5	1166	G	N3-C4-C5	5.94	131.57	128.60
36	5	1179	A	C2-N3-C4	-5.94	107.63	110.60
36	5	2757	U	C6-N1-C2	-5.94	117.44	121.00
36	5	1124	U	C6-N1-C2	-5.94	117.44	121.00
36	1	2991	A	N1-C2-N3	5.94	132.27	129.30
1	6	20	G	C2-N3-C4	-5.94	108.93	111.90
1	6	1058	U	P-O3'-C3'	5.94	126.83	119.70
36	5	2416	U	N3-C2-O2	-5.94	118.04	122.20
38	8	17	A	C4-C5-N7	5.94	113.67	110.70
36	1	1323	G	N3-C4-N9	5.94	129.56	126.00
1	6	194	U	C2-N1-C1'	5.93	124.82	117.70
36	5	922	U	N3-C2-O2	-5.93	118.05	122.20
36	5	1065	A	C8-N9-C4	5.93	108.17	105.80
36	5	1611	G	N1-C6-O6	5.93	123.46	119.90
36	1	2830	G	N1-C6-O6	5.93	123.46	119.90
36	1	2942	C	C2-N1-C1'	-5.93	112.28	118.80
36	5	974	G	C8-N9-C4	-5.93	104.03	106.40
37	7	44	C	C4-C5-C6	5.93	120.37	117.40
1	2	51	A	C8-N9-C4	5.93	108.17	105.80
50	M4	30	GLY	N-CA-C	-5.93	98.27	113.10
36	5	632	G	N3-C4-N9	5.93	129.56	126.00
36	5	709	A	N1-C6-N6	5.93	122.16	118.60
36	1	1536	G	N1-C2-N2	5.93	121.54	116.20
1	2	499	U	P-O3'-C3'	5.93	126.81	119.70
36	5	1179	A	OP1-P-OP2	-5.93	110.71	119.60
36	1	691	A	N1-C6-N6	5.93	122.16	118.60
36	1	1416	C	N3-C4-C5	5.93	124.27	121.90
36	1	1437	C	C5-C6-N1	5.93	123.96	121.00
1	6	443	C	N3-C4-N4	5.93	122.15	118.00
37	7	82	G	N1-C6-O6	5.93	123.46	119.90
1	2	1052	U	C2-N1-C1'	5.92	124.81	117.70
36	1	2683	U	C6-N1-C2	5.92	124.56	121.00
1	6	542	A	P-O3'-C3'	5.92	126.81	119.70
36	1	2621	G	N1-C6-O6	5.92	123.45	119.90
36	5	2818	U	N1-C2-N3	-5.92	111.35	114.90
36	5	2944	U	N3-C4-O4	-5.92	115.25	119.40
1	2	1560	U	C6-N1-C2	-5.92	117.45	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	968	G	N3-C4-N9	5.92	129.55	126.00
36	1	2300	G	O5'-P-OP1	-5.92	100.37	105.70
1	6	1568	C	P-O3'-C3'	5.92	126.81	119.70
36	5	2705	A	C8-N9-C4	5.92	108.17	105.80
1	2	425	A	N1-C6-N6	5.92	122.15	118.60
36	5	86	G	N3-C4-C5	-5.92	125.64	128.60
36	1	833	G	C5-N7-C8	5.92	107.26	104.30
36	1	2112	U	P-O3'-C3'	5.92	126.80	119.70
36	1	2408	U	O5'-P-OP1	-5.92	100.37	105.70
36	5	814	U	C6-N1-C2	-5.92	117.45	121.00
36	5	3200	G	N1-C6-O6	5.92	123.45	119.90
36	1	2725	U	C5-C6-N1	-5.92	119.74	122.70
1	6	1614	A	C5-C6-N1	-5.92	114.74	117.70
1	2	1241	G	C8-N9-C4	-5.91	104.03	106.40
36	5	1181	U	N3-C4-C5	-5.91	111.05	114.60
36	5	1335	C	C5-C4-N4	-5.91	116.06	120.20
1	6	412	A	C8-N9-C4	-5.91	103.44	105.80
36	5	2128	C	C6-N1-C2	-5.91	117.94	120.30
36	1	2175	U	C4-C5-C6	5.91	123.25	119.70
36	1	3344	A	C4-C5-N7	5.91	113.65	110.70
36	5	1316	C	N3-C4-N4	5.91	122.14	118.00
1	2	1100	G	C4-N9-C1'	5.91	134.18	126.50
1	2	1768	G	N9-C4-C5	5.91	107.76	105.40
36	1	1179	A	C2-N3-C4	-5.91	107.65	110.60
1	6	66	U	P-O3'-C3'	5.91	126.79	119.70
36	5	2846	U	N3-C2-O2	-5.91	118.06	122.20
36	1	300	G	C5-C6-O6	5.91	132.14	128.60
36	1	1419	A	C6-N1-C2	-5.91	115.06	118.60
36	1	2333	C	N3-C4-N4	-5.91	113.87	118.00
36	1	2336	U	N3-C2-O2	-5.91	118.07	122.20
1	6	1306	C	C5-C6-N1	5.91	123.95	121.00
36	5	991	G	N1-C6-O6	-5.91	116.36	119.90
36	5	1845	G	N3-C4-N9	5.91	129.54	126.00
36	5	2361	A	C8-N9-C4	-5.91	103.44	105.80
36	5	2888	U	O5'-P-OP1	-5.91	100.39	105.70
36	1	787	G	N3-C4-C5	-5.90	125.65	128.60
36	1	789	A	N1-C6-N6	-5.90	115.06	118.60
36	1	2370	G	C5-C6-N1	-5.90	108.55	111.50
1	6	1132	A	O5'-P-OP2	-5.90	100.39	105.70
36	5	522	A	C5-C6-N6	-5.90	118.98	123.70
36	5	994	G	N3-C4-C5	-5.90	125.65	128.60
36	1	2917	G	C2-N3-C4	5.90	114.85	111.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	1306	G	C6-C5-N7	-5.90	126.86	130.40
36	5	2802	A	N1-C2-N3	-5.90	126.35	129.30
36	5	3362	A	N1-C6-N6	5.90	122.14	118.60
1	2	610	G	C8-N9-C1'	-5.90	119.33	127.00
1	6	1010	C	N3-C4-C5	-5.90	119.54	121.90
36	5	2111	G	N3-C4-N9	-5.90	122.46	126.00
36	1	1269	U	N1-C2-O2	5.90	126.93	122.80
36	1	2777	G	C8-N9-C4	-5.90	104.04	106.40
1	2	377	G	C8-N9-C4	5.89	108.76	106.40
36	5	1466	G	C5-C6-O6	-5.89	125.06	128.60
36	1	3344	A	C4-C5-C6	5.89	119.95	117.00
36	5	431	U	C6-N1-C2	5.89	124.53	121.00
36	5	3131	U	C6-N1-C2	5.89	124.54	121.00
42	15	110	LEU	CA-CB-CG	5.89	128.85	115.30
1	2	971	A	N1-C2-N3	5.89	132.25	129.30
36	1	1802	C	C5-C6-N1	5.89	123.94	121.00
36	1	2915	U	N3-C4-O4	5.89	123.52	119.40
37	3	40	C	C6-N1-C2	-5.89	117.94	120.30
1	6	1208	A	O4'-C1'-N9	5.89	112.91	108.20
36	1	335	G	N3-C4-N9	-5.89	122.47	126.00
36	1	2393	G	C4-C5-N7	5.89	113.16	110.80
1	6	1730	A	N9-C4-C5	5.89	108.16	105.80
1	6	448	C	C6-N1-C2	-5.89	117.94	120.30
1	6	455	C	C5-C6-N1	5.89	123.94	121.00
36	5	3303	G	C8-N9-C4	5.89	108.75	106.40
36	5	656	A	C5-C6-N6	-5.88	118.99	123.70
36	5	887	G	C5-N7-C8	-5.88	101.36	104.30
36	5	2715	A	OP2-P-O3'	5.88	118.15	105.20
36	5	2942	C	C6-N1-C2	-5.88	117.95	120.30
36	1	1745	C	O5'-P-OP2	-5.88	100.41	105.70
1	6	1533	C	C6-N1-C2	-5.88	117.95	120.30
36	5	968	G	N1-C6-O6	5.88	123.43	119.90
36	5	1149	G	C2-N3-C4	-5.88	108.96	111.90
36	5	2231	C	O4'-C1'-N1	5.88	112.91	108.20
36	5	3285	C	N1-C2-O2	5.88	122.43	118.90
36	1	212	G	C8-N9-C1'	-5.88	119.36	127.00
36	1	344	A	N1-C6-N6	-5.88	115.07	118.60
36	1	1694	U	N1-C2-O2	5.88	126.91	122.80
36	1	2890	A	N9-C4-C5	5.88	108.15	105.80
36	1	3214	U	C6-N1-C2	-5.88	117.47	121.00
36	5	2585	G	N3-C4-C5	-5.88	125.66	128.60
36	5	3299	A	N9-C4-C5	5.88	108.15	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	2	927	C	O5'-P-OP1	-5.88	100.41	105.70
36	1	348	A	C8-N9-C4	5.88	108.15	105.80
36	1	350	C	C2-N1-C1'	5.88	125.26	118.80
36	1	364	G	C4-C5-N7	5.88	113.15	110.80
36	1	682	U	N3-C4-O4	-5.88	115.29	119.40
36	1	857	G	C5-C6-N1	-5.88	108.56	111.50
1	6	1174	C	N1-C2-O2	5.88	122.43	118.90
36	5	2385	G	C5-C6-N1	-5.88	108.56	111.50
36	5	2992	U	C2-N1-C1'	5.88	124.75	117.70
36	1	1836	C	N1-C2-O2	5.88	122.42	118.90
36	1	188	U	N1-C2-O2	-5.87	118.69	122.80
36	1	214	G	C5-C6-O6	-5.87	125.08	128.60
36	1	648	C	N3-C2-O2	-5.87	117.79	121.90
36	1	937	G	C5-C6-O6	-5.87	125.08	128.60
36	1	2860	U	N1-C2-N3	-5.87	111.38	114.90
36	1	2876	C	N1-C2-O2	-5.87	115.38	118.90
1	6	334	G	C8-N9-C1'	5.87	134.63	127.00
36	5	356	C	C6-N1-C2	-5.87	117.95	120.30
1	6	1791	A	N1-C6-N6	5.87	122.12	118.60
36	5	36	C	C5-C4-N4	-5.87	116.09	120.20
36	5	2211	U	N3-C2-O2	-5.87	118.09	122.20
36	1	435	C	C6-N1-C2	5.87	122.65	120.30
36	1	2309	A	C5-C6-N6	-5.87	119.00	123.70
36	1	2356	A	C6-C5-N7	-5.87	128.19	132.30
1	6	1118	G	C8-N9-C4	5.87	108.75	106.40
36	1	86	G	C8-N9-C1'	5.87	134.63	127.00
36	5	397	A	N1-C6-N6	-5.87	115.08	118.60
36	5	2907	G	N3-C4-C5	5.87	131.53	128.60
36	5	3036	G	C5-C6-O6	5.87	132.12	128.60
36	1	2877	G	C8-N9-C4	-5.86	104.06	106.40
47	M0	167	LEU	CA-CB-CG	5.86	128.79	115.30
36	1	335	G	N3-C4-C5	5.86	131.53	128.60
36	5	1481	A	P-O3'-C3'	5.86	126.73	119.70
36	5	2904	U	C6-N1-C1'	-5.86	113.00	121.20
1	2	1110	G	C8-N9-C4	5.86	108.74	106.40
36	1	1366	A	N9-C4-C5	5.86	108.14	105.80
36	1	2356	A	C4-C5-N7	5.86	113.63	110.70
36	1	3303	G	C5-C6-O6	-5.86	125.09	128.60
36	5	1514	G	C6-C5-N7	-5.86	126.89	130.40
36	5	2281	A	C5-C6-N6	-5.86	119.02	123.70
1	2	793	A	C8-N9-C4	-5.85	103.46	105.80
1	2	1003	A	O4'-C1'-N9	5.85	112.88	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	817	A	N9-C1'-C2'	5.85	121.61	114.00
36	1	2860	U	C5-C6-N1	5.85	125.63	122.70
36	1	1733	G	N3-C4-N9	5.85	129.51	126.00
36	1	2688	U	C5-C4-O4	-5.85	122.39	125.90
36	1	3270	U	O5'-P-OP1	-5.85	100.43	105.70
36	5	3303	G	N3-C4-C5	5.85	131.52	128.60
36	5	658	G	C6-C5-N7	-5.85	126.89	130.40
36	5	2706	G	O5'-P-OP2	-5.85	100.44	105.70
38	8	42	G	N1-C6-O6	5.85	123.41	119.90
36	1	2828	G	N3-C4-N9	5.85	129.51	126.00
36	5	2337	C	N1-C2-O2	5.85	122.41	118.90
1	2	734	A	OP1-P-O3'	5.84	118.06	105.20
36	1	864	G	C5-C6-O6	5.84	132.11	128.60
1	6	1421	A	C8-N9-C4	5.84	108.14	105.80
36	5	328	U	C5-C4-O4	5.84	129.41	125.90
36	5	2411	U	C6-N1-C2	5.84	124.51	121.00
36	5	2817	A	N3-C4-N9	5.84	132.08	127.40
36	5	3049	A	C8-N9-C4	5.84	108.14	105.80
38	8	10	A	C2-N3-C4	-5.84	107.68	110.60
36	1	1124	U	C5-C4-O4	5.84	129.41	125.90
36	1	1126	G	N1-C6-O6	5.84	123.41	119.90
36	5	2345	A	C6-C5-N7	-5.84	128.21	132.30
36	5	2629	U	N1-C2-O2	-5.84	118.71	122.80
36	5	40	A	C6-C5-N7	-5.84	128.21	132.30
36	5	644	G	N3-C4-C5	-5.84	125.68	128.60
1	2	1745	G	N3-C4-N9	5.84	129.50	126.00
36	1	36	C	N3-C4-C5	5.84	124.23	121.90
36	1	394	G	N3-C4-N9	-5.84	122.50	126.00
36	1	1536	G	C5-C6-O6	-5.84	125.10	128.60
36	1	1855	U	C2-N1-C1'	5.84	124.70	117.70
36	5	1394	A	N1-C6-N6	-5.84	115.10	118.60
1	6	1614	A	N1-C6-N6	5.83	122.10	118.60
36	5	2572	C	C2-N1-C1'	5.83	125.22	118.80
47	M0	176	LEU	CA-CB-CG	5.83	128.71	115.30
36	5	959	C	N1-C2-O2	-5.83	115.40	118.90
36	5	1379	G	C5-C6-O6	-5.83	125.10	128.60
36	5	1855	U	N3-C2-O2	-5.83	118.12	122.20
36	5	2700	G	C8-N9-C4	5.83	108.73	106.40
36	1	1311	G	N1-C6-O6	5.83	123.40	119.90
36	5	1855	U	C6-N1-C2	-5.83	117.50	121.00
37	7	96	U	N1-C2-O2	-5.83	118.72	122.80
36	1	1313	G	C5-C6-O6	-5.83	125.10	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	6	65	A	C8-N9-C4	5.83	108.13	105.80
1	6	421	A	C8-N9-C4	5.83	108.13	105.80
1	6	886	U	N3-C2-O2	-5.83	118.12	122.20
36	5	2632	G	N1-C6-O6	5.83	123.40	119.90
54	M8	166	LEU	CA-CB-CG	-5.83	101.90	115.30
36	5	3315	G	N1-C6-O6	5.83	123.40	119.90
1	2	959	U	N1-C2-O2	5.83	126.88	122.80
36	1	974	G	N3-C4-C5	-5.83	125.69	128.60
36	1	1411	C	N3-C2-O2	-5.83	117.82	121.90
36	5	2820	A	C8-N9-C4	-5.83	103.47	105.80
36	1	690	A	N1-C2-N3	5.82	132.21	129.30
36	1	1923	C	C6-N1-C2	5.82	122.63	120.30
36	5	1301	A	N3-C4-N9	5.82	132.06	127.40
36	1	422	A	C8-N9-C4	-5.82	103.47	105.80
38	8	12	A	N1-C6-N6	5.82	122.09	118.60
1	2	447	U	N3-C4-O4	5.82	123.47	119.40
1	2	1100	G	C8-N9-C1'	-5.82	119.43	127.00
36	1	933	A	N1-C2-N3	5.82	132.21	129.30
36	1	2572	C	N1-C2-O2	5.82	122.39	118.90
36	5	519	A	C5-N7-C8	-5.82	100.99	103.90
36	1	1369	A	C4-C5-C6	5.82	119.91	117.00
1	6	1285	U	N3-C2-O2	-5.82	118.13	122.20
36	5	2190	U	N1-C2-N3	5.82	118.39	114.90
36	1	339	C	C2-N3-C4	-5.82	116.99	119.90
36	1	2975	U	N1-C2-O2	5.82	126.87	122.80
36	5	2137	U	N3-C4-C5	5.82	118.09	114.60
1	2	1590	G	N1-C6-O6	-5.82	116.41	119.90
36	1	908	G	C8-N9-C1'	-5.82	119.44	127.00
36	5	826	G	C4-C5-N7	5.82	113.13	110.80
36	1	3362	A	C5-N7-C8	-5.81	100.99	103.90
36	5	720	A	C5-C6-N6	-5.81	119.05	123.70
36	1	2935	U	C2-N3-C4	5.81	130.49	127.00
1	6	13	C	C6-N1-C2	-5.81	117.98	120.30
36	5	969	C	N1-C2-O2	-5.81	115.41	118.90
36	5	1458	U	O5'-P-OP1	-5.81	100.47	105.70
36	5	2388	U	N1-C2-N3	5.81	118.39	114.90
36	1	1179	A	OP2-P-O3'	5.81	117.98	105.20
36	1	2273	G	N7-C8-N9	-5.81	110.19	113.10
36	1	3204	C	N1-C2-O2	5.81	122.39	118.90
36	5	1878	G	P-O3'-C3'	5.81	126.67	119.70
36	5	2929	C	C5-C4-N4	-5.81	116.13	120.20
36	5	3107	U	N3-C4-C5	-5.81	111.11	114.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	2	1105	C	C6-N1-C2	-5.81	117.98	120.30
36	1	809	G	C4-C5-N7	5.81	113.12	110.80
36	5	511	G	N3-C4-N9	-5.81	122.52	126.00
36	5	531	G	O5'-P-OP1	-5.81	100.47	105.70
36	5	896	A	O5'-P-OP2	-5.81	100.47	105.70
54	m8	127	LEU	CA-CB-CG	5.81	128.66	115.30
1	2	73	U	P-O3'-C3'	5.81	126.67	119.70
36	1	213	A	N1-C6-N6	5.81	122.08	118.60
36	1	2555	G	O5'-P-OP2	-5.81	100.47	105.70
1	6	117	U	N3-C2-O2	-5.81	118.14	122.20
1	6	368	U	O5'-P-OP2	-5.81	100.47	105.70
1	2	521	A	C8-N9-C4	-5.80	103.48	105.80
36	1	2397	A	C5-N7-C8	-5.80	101.00	103.90
36	1	2954	U	OP1-P-O3'	5.80	117.97	105.20
36	1	3000	A	C8-N9-C4	5.80	108.12	105.80
36	5	676	G	N3-C4-C5	-5.80	125.70	128.60
36	5	2231	C	C2-N1-C1'	5.80	125.19	118.80
36	5	2801	A	N7-C8-N9	-5.80	110.90	113.80
36	1	1733	G	C8-N9-C4	-5.80	104.08	106.40
36	5	2911	A	O5'-P-OP2	-5.80	100.48	105.70
36	1	1492	G	N3-C4-N9	5.80	129.48	126.00
36	5	366	A	N1-C6-N6	5.80	122.08	118.60
36	5	2392	C	C2-N1-C1'	-5.80	112.42	118.80
36	5	3150	A	N1-C6-N6	5.80	122.08	118.60
1	2	331	A	N1-C6-N6	-5.80	115.12	118.60
1	2	1077	C	C5-C6-N1	5.80	123.90	121.00
36	1	609	G	N1-C6-O6	5.80	123.38	119.90
1	6	901	G	C5-C6-O6	-5.80	125.12	128.60
36	1	776	U	N1-C2-N3	5.80	118.38	114.90
36	1	2434	U	C4-C5-C6	5.80	123.18	119.70
36	5	3108	G	N1-C6-O6	5.80	123.38	119.90
37	7	115	G	C5-C6-O6	-5.80	125.12	128.60
36	5	2800	G	C8-N9-C4	-5.79	104.08	106.40
36	1	339	C	N3-C4-N4	-5.79	113.94	118.00
36	1	2554	A	C8-N9-C4	5.79	108.12	105.80
36	1	2836	C	C6-N1-C2	-5.79	117.98	120.30
1	6	782	U	N1-C2-O2	5.79	126.86	122.80
36	5	804	C	N1-C2-O2	-5.79	115.42	118.90
37	7	3	U	C5-C6-N1	-5.79	119.80	122.70
36	1	343	U	N3-C2-O2	-5.79	118.15	122.20
36	1	682	U	C5-C4-O4	5.79	129.38	125.90
1	6	402	C	O4'-C1'-N1	5.79	112.83	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	6	1115	U	C5-C6-N1	-5.79	119.80	122.70
36	5	1704	A	C8-N9-C4	5.79	108.12	105.80
36	5	2354	C	N3-C4-C5	-5.79	119.58	121.90
70	O4	51	LEU	CA-CB-CG	5.79	128.62	115.30
36	1	1365	G	N3-C4-N9	5.79	129.47	126.00
36	1	3002	C	C6-N1-C2	5.79	122.61	120.30
36	5	2412	G	N3-C4-N9	5.79	129.47	126.00
36	5	2760	C	C2-N3-C4	-5.79	117.00	119.90
36	1	1154	A	C4-C5-C6	5.79	119.89	117.00
36	1	2896	A	N1-C2-N3	5.79	132.19	129.30
36	1	906	A	O4'-C1'-N9	-5.79	103.57	108.20
36	5	567	G	C5-N7-C8	-5.79	101.41	104.30
36	5	2112	U	N1-C2-O2	-5.79	118.75	122.80
36	5	2185	G	C2-N3-C4	-5.79	109.01	111.90
36	5	2364	G	C4-C5-N7	-5.79	108.49	110.80
36	5	2627	C	C4-C5-C6	5.79	120.29	117.40
36	5	3180	A	C6-N1-C2	-5.79	115.13	118.60
36	5	2515	A	N1-C6-N6	-5.78	115.13	118.60
36	5	3010	U	C5-C4-O4	5.78	129.37	125.90
1	2	325	G	N1-C6-O6	5.78	123.37	119.90
1	2	590	C	C6-N1-C2	-5.78	117.99	120.30
36	1	1129	A	N1-C6-N6	5.78	122.07	118.60
36	5	400	G	N9-C4-C5	5.78	107.71	105.40
36	1	1131	G	C6-C5-N7	-5.78	126.93	130.40
36	1	1839	A	O5'-P-OP1	-5.78	100.50	105.70
36	1	2949	U	N3-C2-O2	5.78	126.25	122.20
36	5	647	A	C8-N9-C4	-5.78	103.49	105.80
1	2	93	A	N9-C4-C5	5.78	108.11	105.80
36	1	1303	A	C6-C5-N7	-5.78	128.25	132.30
36	1	1422	G	C6-C5-N7	-5.78	126.93	130.40
36	1	2200	U	N1-C2-O2	-5.78	118.76	122.80
1	6	976	G	C4-C5-N7	5.78	113.11	110.80
36	5	1317	A	N7-C8-N9	5.78	116.69	113.80
36	1	688	G	N3-C4-C5	-5.78	125.71	128.60
36	1	803	C	C5-C4-N4	-5.78	116.16	120.20
36	1	1190	A	C4-C5-C6	5.78	119.89	117.00
36	1	1321	G	N9-C4-C5	5.78	107.71	105.40
36	5	725	G	C4-C5-C6	5.77	122.26	118.80
36	5	2271	A	C2-N3-C4	5.77	113.49	110.60
36	5	2700	G	N9-C4-C5	-5.77	103.09	105.40
36	1	1446	A	C8-N9-C4	-5.77	103.49	105.80
36	1	86	G	O5'-P-OP2	-5.77	100.51	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	345	G	C6-C5-N7	-5.77	126.94	130.40
36	5	2724	U	N3-C4-C5	-5.77	111.14	114.60
36	5	3181	C	N3-C4-C5	-5.77	119.59	121.90
36	1	633	C	N3-C4-C5	-5.77	119.59	121.90
36	1	2688	U	N1-C2-O2	5.77	126.84	122.80
36	1	2919	A	C2-N3-C4	-5.77	107.72	110.60
36	5	1417	G	N3-C4-C5	-5.77	125.72	128.60
1	2	51	A	O5'-P-OP1	-5.77	100.51	105.70
36	1	102	C	O5'-P-OP1	5.77	117.62	110.70
36	1	1905	G	OP2-P-O3'	5.77	117.89	105.20
1	2	465	G	O5'-P-OP1	-5.76	100.51	105.70
36	1	642	U	O5'-P-OP2	-5.76	100.51	105.70
36	1	1442	U	N1-C2-O2	-5.76	118.77	122.80
36	1	2324	A	N1-C6-N6	-5.76	115.14	118.60
36	1	2868	U	N3-C2-O2	-5.76	118.16	122.20
36	5	561	C	N1-C2-O2	-5.76	115.44	118.90
36	5	1062	A	O5'-P-OP2	-5.76	100.51	105.70
36	1	999	G	C5-C6-N1	5.76	114.38	111.50
36	1	2996	U	C5-C6-N1	5.76	125.58	122.70
1	6	334	G	C4-N9-C1'	-5.76	119.01	126.50
36	1	1939	G	C4-N9-C1'	5.76	133.99	126.50
36	1	2365	C	C6-N1-C2	5.76	122.60	120.30
36	5	643	U	N3-C4-O4	5.76	123.43	119.40
36	5	1142	G	C4-N9-C1'	5.76	133.99	126.50
36	1	123	A	C5-C6-N6	-5.76	119.09	123.70
36	1	335	G	C5-N7-C8	-5.76	101.42	104.30
36	1	2664	C	C6-N1-C2	-5.76	118.00	120.30
36	5	2142	A	C2-N3-C4	5.76	113.48	110.60
1	2	425	A	C5-N7-C8	-5.76	101.02	103.90
36	1	968	G	N3-C4-C5	-5.76	125.72	128.60
36	5	1208	U	N3-C2-O2	-5.76	118.17	122.20
36	5	1193	A	C2-N3-C4	-5.75	107.72	110.60
38	8	96	A	C5-C6-N6	-5.75	119.10	123.70
37	3	91	G	N1-C6-O6	5.75	123.35	119.90
36	5	938	C	OP1-P-O3'	5.75	117.86	105.20
36	5	1886	A	C8-N9-C4	-5.75	103.50	105.80
36	5	2411	U	C2-N1-C1'	-5.75	110.80	117.70
1	2	28	A	C5-C6-N1	5.75	120.58	117.70
36	1	1296	C	C6-N1-C2	-5.75	118.00	120.30
36	1	2403	G	C4-C5-C6	5.75	122.25	118.80
36	5	1142	G	N3-C4-N9	5.75	129.45	126.00
36	5	2112	U	P-O3'-C3'	5.75	126.60	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	8	84	C	N3-C4-C5	-5.75	119.60	121.90
1	2	610	G	C4-N9-C1'	5.75	133.97	126.50
1	6	957	G	N1-C6-O6	5.75	123.35	119.90
37	7	73	C	OP1-P-O3'	5.75	117.85	105.20
36	1	1084	A	N1-C6-N6	5.75	122.05	118.60
36	1	2774	C	C5-C4-N4	-5.75	116.17	120.20
36	5	2805	G	C5-C6-O6	-5.75	125.15	128.60
36	1	2697	A	N9-C4-C5	5.75	108.10	105.80
36	1	3028	G	C8-N9-C4	-5.75	104.10	106.40
36	5	1371	G	N7-C8-N9	-5.75	110.23	113.10
36	5	2621	G	N3-C2-N2	-5.75	115.88	119.90
36	1	570	A	N1-C6-N6	-5.74	115.15	118.60
36	1	716	A	N1-C6-N6	5.74	122.05	118.60
1	6	87	C	C6-N1-C2	-5.74	118.00	120.30
36	5	651	G	C6-C5-N7	-5.74	126.95	130.40
36	5	1124	U	N1-C2-N3	5.74	118.35	114.90
36	1	1001	G	C8-N9-C1'	-5.74	119.54	127.00
36	1	3214	U	C5-C4-O4	5.74	129.34	125.90
36	1	1171	G	N3-C4-N9	-5.74	122.56	126.00
1	6	1594	G	O5'-P-OP1	-5.74	100.53	105.70
37	7	48	U	C5-C4-O4	-5.74	122.46	125.90
1	2	555	A	C8-N9-C4	-5.74	103.50	105.80
36	1	2370	G	C5-C6-O6	5.74	132.04	128.60
36	1	3266	G	C5-C6-O6	5.74	132.04	128.60
36	5	1885	U	C5-C6-N1	-5.74	119.83	122.70
1	2	736	C	C6-N1-C2	-5.74	118.01	120.30
36	1	375	A	OP1-P-O3'	5.74	117.82	105.20
36	1	1330	A	C5-C6-N1	-5.74	114.83	117.70
36	1	2406	C	C5-C4-N4	-5.74	116.19	120.20
36	1	2647	A	C5-C6-N6	-5.74	119.11	123.70
1	6	1645	G	N9-C4-C5	5.74	107.69	105.40
36	5	994	G	C2-N3-C4	5.74	114.77	111.90
36	5	2234	G	C8-N9-C4	5.74	108.69	106.40
36	5	1048	A	N7-C8-N9	-5.73	110.93	113.80
36	1	3181	C	N3-C4-C5	-5.73	119.61	121.90
36	5	1793	C	N3-C2-O2	5.73	125.91	121.90
36	5	2818	U	C5-C4-O4	-5.73	122.46	125.90
1	2	1753	A	N1-C6-N6	5.73	122.04	118.60
36	1	2938	G	C6-N1-C2	-5.73	121.66	125.10
36	1	3112	G	N3-C4-N9	5.73	129.44	126.00
37	3	66	A	C8-N9-C4	5.73	108.09	105.80
1	6	815	G	C4-C5-N7	5.73	113.09	110.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	6	1575	G	N1-C6-O6	-5.73	116.46	119.90
36	5	400	G	C5-C6-O6	5.73	132.04	128.60
36	1	1303	A	C4-C5-N7	5.73	113.56	110.70
36	1	2287	C	N1-C2-O2	-5.73	115.46	118.90
1	6	95	G	C2-N3-C4	5.73	114.76	111.90
1	6	187	G	P-O3'-C3'	5.73	126.57	119.70
36	5	508	U	C6-N1-C2	-5.73	117.56	121.00
36	5	1190	A	N1-C6-N6	-5.73	115.16	118.60
1	6	422	G	C8-N9-C4	-5.73	104.11	106.40
36	5	3270	U	O5'-P-OP1	-5.73	100.55	105.70
1	2	720	G	OP1-P-O3'	5.72	117.80	105.20
36	1	363	G	C8-N9-C4	-5.72	104.11	106.40
37	7	101	G	N1-C6-O6	5.72	123.33	119.90
36	1	1279	C	C5-C6-N1	5.72	123.86	121.00
36	1	1419	A	O5'-P-OP1	5.72	117.57	110.70
36	1	2829	U	N3-C4-O4	5.72	123.41	119.40
36	1	145	G	C6-C5-N7	-5.72	126.97	130.40
36	1	895	A	C2-N3-C4	-5.72	107.74	110.60
1	6	1340	U	C5-C4-O4	5.72	129.33	125.90
36	1	997	A	C8-N9-C4	-5.72	103.51	105.80
36	1	2180	G	C2-N3-C4	-5.72	109.04	111.90
36	1	2639	G	C8-N9-C1'	-5.72	119.56	127.00
36	1	3063	C	C2-N3-C4	-5.72	117.04	119.90
1	6	390	G	C5-C6-O6	-5.72	125.17	128.60
12	c0	83	PRO	N-CA-CB	5.72	110.16	103.30
36	1	342	A	C8-N9-C4	-5.72	103.51	105.80
36	1	2890	A	C8-N9-C4	-5.72	103.51	105.80
36	5	3129	A	O4'-C1'-N9	5.72	112.78	108.20
3	S1	181	LEU	CA-CB-CG	5.72	128.45	115.30
36	1	1445	U	C2-N3-C4	-5.72	123.57	127.00
36	1	3143	C	N3-C2-O2	5.72	125.90	121.90
37	7	37	G	C5-C6-O6	-5.72	125.17	128.60
1	2	730	G	C4-N9-C1'	5.71	133.93	126.50
36	1	2300	G	C5-C6-N1	-5.71	108.64	111.50
36	1	2903	A	N1-C6-N6	5.71	122.03	118.60
37	3	33	U	N1-C2-O2	5.71	126.80	122.80
8	s6	133	LEU	CA-CB-CG	5.71	128.44	115.30
36	5	2897	A	C6-N1-C2	-5.71	115.17	118.60
36	5	3150	A	C2-N3-C4	-5.71	107.74	110.60
36	1	974	G	N3-C4-N9	5.71	129.43	126.00
36	1	2924	U	O5'-P-OP1	-5.71	100.56	105.70
36	1	2237	C	N1-C2-O2	5.71	122.33	118.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	6	1490	C	C6-N1-C2	-5.71	118.02	120.30
36	5	282	G	N1-C2-N2	-5.71	111.06	116.20
36	5	366	A	C8-N9-C4	5.71	108.08	105.80
38	8	87	G	O4'-C1'-N9	5.71	112.77	108.20
1	2	542	A	O4'-C1'-N9	5.71	112.77	108.20
1	6	67	A	N1-C6-N6	5.71	122.03	118.60
1	6	338	C	C5-C6-N1	5.71	123.86	121.00
1	2	1622	G	N3-C4-C5	5.71	131.45	128.60
36	1	1316	C	N3-C4-N4	5.71	122.00	118.00
36	1	1445	U	N1-C2-O2	-5.71	118.80	122.80
36	1	1952	G	N3-C4-C5	-5.71	125.75	128.60
36	1	2525	G	N3-C4-N9	5.71	129.43	126.00
36	5	934	G	C5-C6-O6	-5.71	125.17	128.60
36	5	1913	A	C4-C5-C6	5.71	119.85	117.00
36	5	2400	G	C5-C6-O6	-5.71	125.17	128.60
36	5	3029	A	C5-C6-N6	5.71	128.27	123.70
36	1	760	G	O4'-C1'-N9	5.71	112.77	108.20
36	1	918	C	C2-N1-C1'	-5.71	112.52	118.80
36	1	2827	U	C6-N1-C1'	5.71	129.19	121.20
1	6	65	A	N3-C4-C5	5.71	130.79	126.80
1	6	938	G	N3-C4-N9	5.71	129.42	126.00
36	5	2729	U	O5'-P-OP2	5.71	117.55	110.70
37	7	103	A	OP2-P-O3'	5.71	117.75	105.20
36	1	1010	G	C5-C6-O6	-5.71	125.18	128.60
36	1	1308	A	O5'-P-OP1	-5.70	100.57	105.70
36	5	1604	G	C4-C5-C6	5.70	122.22	118.80
36	1	2366	C	O5'-P-OP2	-5.70	100.57	105.70
36	1	2916	U	C5-C6-N1	5.70	125.55	122.70
36	1	3057	U	N1-C2-N3	5.70	118.32	114.90
36	5	1198	C	N3-C4-N4	-5.70	114.01	118.00
1	2	1455	G	C5-C6-N1	-5.70	108.65	111.50
36	1	773	G	N9-C4-C5	5.70	107.68	105.40
36	5	2419	A	N7-C8-N9	5.70	116.65	113.80
1	2	511	A	C8-N9-C4	-5.70	103.52	105.80
1	6	1637	C	C2-N1-C1'	5.70	125.07	118.80
36	5	808	A	N9-C4-C5	5.70	108.08	105.80
36	5	1332	A	C8-N9-C1'	-5.70	117.44	127.70
36	5	1404	G	C5-C6-N1	-5.70	108.65	111.50
1	6	1117	U	O5'-P-OP2	-5.70	100.57	105.70
36	1	924	G	C4-C5-N7	5.70	113.08	110.80
36	1	2329	C	N1-C2-O2	-5.70	115.48	118.90
36	5	2130	G	N9-C1'-C2'	-5.70	105.73	112.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
38	8	80	A	C8-N9-C4	-5.70	103.52	105.80
36	1	2632	G	C2-N3-C4	-5.69	109.05	111.90
36	1	3180	A	C8-N9-C4	5.69	108.08	105.80
73	O7	65	ARG	NE-CZ-NH1	5.69	123.15	120.30
1	6	1133	A	N1-C6-N6	5.69	122.02	118.60
36	5	2849	C	N1-C2-O2	-5.69	115.48	118.90
36	5	3179	U	N3-C2-O2	-5.69	118.21	122.20
52	m6	78	ARG	NE-CZ-NH1	5.69	123.15	120.30
36	1	2869	U	O5'-P-OP2	5.69	117.53	110.70
36	5	694	C	N3-C2-O2	-5.69	117.92	121.90
36	1	2620	G	C2-N3-C4	5.69	114.75	111.90
36	1	3093	C	C6-N1-C2	5.69	122.58	120.30
1	6	1000	C	C2-N1-C1'	5.69	125.06	118.80
36	5	1417	G	OP1-P-OP2	-5.69	111.07	119.60
36	5	1744	G	C8-N9-C1'	-5.69	119.60	127.00
36	5	1868	G	N1-C6-O6	5.69	123.31	119.90
36	5	3028	G	N1-C2-N2	-5.69	111.08	116.20
38	4	9	A	O5'-P-OP2	-5.69	100.58	105.70
1	2	1749	A	N1-C2-N3	5.69	132.14	129.30
36	1	967	A	O5'-P-OP1	-5.69	100.58	105.70
36	1	1187	C	C4-C5-C6	-5.69	114.56	117.40
36	1	2554	A	P-O3'-C3'	5.69	126.52	119.70
1	6	517	U	C5-C4-O4	5.69	129.31	125.90
36	5	1319	G	C5-N7-C8	5.69	107.14	104.30
36	5	1556	C	C6-N1-C2	-5.69	118.03	120.30
36	5	2129	U	C6-N1-C2	-5.69	117.59	121.00
36	5	2687	G	N3-C4-N9	5.69	129.41	126.00
36	5	2923	U	C6-N1-C2	-5.69	117.59	121.00
36	5	3195	U	P-O3'-C3'	5.69	126.53	119.70
1	2	1241	G	O4'-C1'-N9	5.69	112.75	108.20
1	2	1145	U	N1-C2-O2	-5.68	118.82	122.80
36	1	780	A	N1-C2-N3	5.68	132.14	129.30
36	1	1294	A	O4'-C1'-N9	5.68	112.75	108.20
36	1	2403	G	N3-C4-N9	5.68	129.41	126.00
36	1	2403	G	C8-N9-C1'	-5.68	119.61	127.00
36	1	3015	G	C5-C6-O6	-5.68	125.19	128.60
36	5	883	A	N1-C6-N6	-5.68	115.19	118.60
36	5	1927	G	C8-N9-C1'	-5.68	119.61	127.00
36	5	2376	G	N7-C8-N9	5.68	115.94	113.10
36	5	3144	G	N1-C2-N2	-5.68	111.08	116.20
36	1	637	C	C6-N1-C1'	-5.68	113.98	120.80
36	5	1879	A	C6-C5-N7	-5.68	128.32	132.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	2	163	G	C4-N9-C1'	5.68	133.88	126.50
1	2	1773	C	C5-C6-N1	5.68	123.84	121.00
36	1	591	G	O5'-P-OP2	-5.68	100.59	105.70
36	1	776	U	N3-C2-O2	-5.68	118.22	122.20
36	1	953	G	C6-C5-N7	-5.68	126.99	130.40
36	1	2877	G	C4-C5-N7	-5.68	108.53	110.80
36	1	3269	U	N3-C2-O2	-5.68	118.22	122.20
1	6	1653	C	N3-C4-C5	-5.68	119.63	121.90
36	5	2875	U	C5-C6-N1	5.68	125.54	122.70
36	1	2976	A	N9-C4-C5	5.68	108.07	105.80
37	3	82	G	N3-C4-N9	5.68	129.41	126.00
36	5	1450	G	N3-C2-N2	-5.68	115.92	119.90
1	2	1747	G	C5-C6-N1	-5.68	108.66	111.50
36	1	54	C	O5'-P-OP2	5.68	117.51	110.70
36	5	888	A	C4-C5-N7	5.68	113.54	110.70
36	5	2113	A	N7-C8-N9	-5.68	110.96	113.80
36	1	1863	G	C6-C5-N7	-5.67	127.00	130.40
36	5	938	C	N3-C4-C5	5.67	124.17	121.90
1	6	997	G	C8-N9-C4	5.67	108.67	106.40
36	5	1205	A	C8-N9-C4	-5.67	103.53	105.80
36	5	2717	U	C2-N1-C1'	-5.67	110.89	117.70
36	1	948	C	C2-N1-C1'	-5.67	112.56	118.80
36	1	979	U	N1-C2-N3	5.67	118.30	114.90
36	1	3182	G	N1-C2-N2	-5.67	111.10	116.20
37	7	82	G	N9-C4-C5	-5.67	103.13	105.40
36	1	3022	G	C6-C5-N7	-5.67	127.00	130.40
36	5	116	A	O4'-C1'-N9	5.67	112.73	108.20
36	5	1184	A	N7-C8-N9	-5.67	110.97	113.80
36	5	2754	G	N3-C2-N2	5.67	123.87	119.90
36	1	1544	G	N3-C4-N9	5.67	129.40	126.00
36	1	358	G	N9-C4-C5	-5.66	103.14	105.40
36	1	2124	G	C5-C6-O6	-5.66	125.20	128.60
36	1	2905	U	N1-C2-O2	-5.66	118.84	122.80
36	1	3172	A	N1-C6-N6	-5.66	115.20	118.60
36	5	882	A	C6-N1-C2	-5.66	115.20	118.60
36	5	1927	G	C4-N9-C1'	5.66	133.86	126.50
36	5	2240	G	C8-N9-C4	-5.66	104.14	106.40
1	2	539	G	N7-C8-N9	5.66	115.93	113.10
36	1	997	A	O5'-P-OP2	-5.66	100.61	105.70
36	1	1604	G	N3-C4-C5	-5.66	125.77	128.60
36	1	2727	A	N1-C6-N6	-5.66	115.20	118.60
36	1	2728	G	N9-C4-C5	-5.66	103.14	105.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
44	L7	202	LEU	CB-CG-CD2	-5.66	101.38	111.00
1	6	1115	U	C6-N1-C2	5.66	124.40	121.00
36	5	91	G	C8-N9-C4	5.66	108.66	106.40
36	5	400	G	N1-C6-O6	-5.66	116.50	119.90
36	5	914	A	C4-C5-C6	5.66	119.83	117.00
36	1	3139	A	C8-N9-C4	-5.66	103.54	105.80
1	6	530	C	C6-N1-C2	5.66	122.56	120.30
36	5	304	G	C4-C5-N7	-5.66	108.54	110.80
36	1	362	U	C5-C6-N1	-5.66	119.87	122.70
36	1	1124	U	N1-C2-O2	5.66	126.76	122.80
36	1	1796	G	N3-C4-C5	-5.66	125.77	128.60
36	1	1415	U	C5-C6-N1	-5.65	119.87	122.70
36	5	2806	U	O5'-P-OP2	-5.65	100.61	105.70
36	1	416	A	N1-C6-N6	5.65	121.99	118.60
36	1	769	G	N9-C4-C5	-5.65	103.14	105.40
36	1	1683	A	N1-C6-N6	5.65	121.99	118.60
36	1	2279	A	C5-N7-C8	-5.65	101.07	103.90
36	1	2342	U	N1-C2-O2	-5.65	118.84	122.80
36	1	2369	G	O5'-P-OP1	-5.65	100.61	105.70
36	5	397	A	N9-C4-C5	5.65	108.06	105.80
36	5	1212	A	N3-C4-N9	5.65	131.92	127.40
1	2	1241	G	C5-N7-C8	-5.65	101.47	104.30
36	1	328	U	N3-C2-O2	-5.65	118.24	122.20
36	1	684	G	C5-C6-O6	-5.65	125.21	128.60
36	1	1412	G	N1-C6-O6	5.65	123.29	119.90
56	N0	115	ARG	NE-CZ-NH2	-5.65	117.47	120.30
36	5	1592	G	C5-C6-O6	5.65	131.99	128.60
36	5	2380	U	O5'-P-OP2	-5.65	100.61	105.70
36	1	424	G	N1-C6-O6	5.65	123.29	119.90
36	1	691	A	C6-C5-N7	-5.65	128.35	132.30
36	5	423	A	N7-C8-N9	-5.65	110.97	113.80
36	1	1041	U	O5'-P-OP2	-5.65	100.62	105.70
1	6	543	C	N3-C4-N4	-5.65	114.05	118.00
1	6	575	C	N3-C4-C5	5.65	124.16	121.90
36	5	1147	G	N3-C4-C5	-5.65	125.78	128.60
36	5	2663	G	C5-C6-O6	-5.65	125.21	128.60
36	5	2690	G	C8-N9-C1'	-5.65	119.66	127.00
1	6	42	G	O4'-C1'-N9	5.65	112.72	108.20
36	5	644	G	C4-N9-C1'	5.64	133.84	126.50
36	5	971	G	OP2-P-O3'	5.64	117.62	105.20
36	1	320	G	O5'-P-OP2	-5.64	100.62	105.70
36	1	785	G	C8-N9-C4	-5.64	104.14	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	1391	C	C5-C4-N4	-5.64	116.25	120.20
36	1	3382	U	N1-C2-O2	5.64	126.75	122.80
36	5	947	G	C5-C6-N1	-5.64	108.68	111.50
1	2	507	U	C2-N1-C1'	5.64	124.47	117.70
36	1	933	A	N7-C8-N9	5.64	116.62	113.80
36	1	953	G	C5-C6-O6	-5.64	125.22	128.60
36	5	794	U	O5'-P-OP2	-5.64	100.62	105.70
36	5	2662	G	N3-C4-C5	-5.64	125.78	128.60
36	5	3020	U	N3-C4-O4	5.64	123.35	119.40
46	19	166	ARG	NE-CZ-NH2	5.64	123.12	120.30
1	2	407	A	N9-C4-C5	5.64	108.06	105.80
36	5	1496	C	C5-C6-N1	5.64	123.82	121.00
1	2	794	U	N3-C2-O2	-5.64	118.25	122.20
36	1	2395	G	C5-N7-C8	-5.64	101.48	104.30
36	1	2639	G	C4-N9-C1'	5.64	133.83	126.50
36	1	2899	C	O4'-C1'-N1	5.64	112.71	108.20
36	1	3150	A	N1-C6-N6	5.64	121.98	118.60
1	6	261	U	N1-C2-O2	5.64	126.75	122.80
36	5	2892	A	C4-C5-C6	5.64	119.82	117.00
36	5	3090	U	N3-C2-O2	-5.64	118.25	122.20
36	5	3218	A	C4-C5-N7	5.64	113.52	110.70
1	2	145	A	C8-N9-C4	-5.63	103.55	105.80
36	1	1883	A	C6-N1-C2	-5.63	115.22	118.60
36	5	1438	U	C2-N1-C1'	5.63	124.46	117.70
36	5	2231	C	C6-N1-C2	-5.63	118.05	120.30
36	5	2664	C	OP2-P-O3'	5.63	117.59	105.20
36	5	27	C	C6-N1-C2	-5.63	118.05	120.30
36	1	1445	U	C5-C6-N1	-5.63	119.88	122.70
36	1	3243	A	C4-C5-N7	5.63	113.52	110.70
36	5	660	A	C5-C6-N6	5.63	128.21	123.70
38	8	4	C	N1-C2-O2	5.63	122.28	118.90
36	1	102	C	OP2-P-O3'	5.63	117.59	105.20
38	4	103	G	N3-C4-C5	-5.63	125.78	128.60
1	6	1614	A	C4-C5-N7	5.63	113.51	110.70
36	5	1449	A	C5-C6-N6	-5.63	119.20	123.70
36	5	2304	C	N1-C2-O2	-5.63	115.52	118.90
36	1	666	A	C8-N9-C4	5.63	108.05	105.80
36	1	1845	G	OP2-P-O3'	5.63	117.58	105.20
1	6	606	A	N1-C6-N6	5.63	121.98	118.60
36	5	1117	G	C4-C5-N7	5.63	113.05	110.80
36	5	1312	C	N3-C4-C5	-5.63	119.65	121.90
36	5	1315	U	C6-N1-C2	5.63	124.38	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	1369	A	N1-C2-N3	5.62	132.11	129.30
36	1	3243	A	C5-C6-N6	-5.62	119.20	123.70
36	5	1509	A	N1-C6-N6	5.62	121.97	118.60
36	5	2197	C	N3-C2-O2	5.62	125.84	121.90
36	1	558	U	C6-N1-C1'	-5.62	113.33	121.20
36	5	98	G	N3-C4-C5	-5.62	125.79	128.60
36	5	2385	G	N3-C2-N2	-5.62	115.96	119.90
36	5	2531	C	C6-N1-C2	-5.62	118.05	120.30
71	o5	69	LEU	CA-CB-CG	5.62	128.23	115.30
36	1	506	U	N1-C2-O2	-5.62	118.86	122.80
36	1	2658	G	O5'-P-OP2	-5.62	100.64	105.70
36	5	501	A	N1-C6-N6	-5.62	115.23	118.60
36	5	2320	A	C2-N3-C4	-5.62	107.79	110.60
36	5	2410	U	C2-N3-C4	-5.62	123.63	127.00
36	1	979	U	N1-C2-O2	5.62	126.73	122.80
36	5	2873	U	N3-C2-O2	-5.62	118.27	122.20
71	o5	89	ARG	NE-CZ-NH1	5.62	123.11	120.30
36	1	3375	A	C5-C6-N6	5.62	128.19	123.70
1	6	176	C	C2-N1-C1'	5.62	124.98	118.80
36	5	1296	C	C5-C4-N4	-5.62	116.27	120.20
36	5	1599	G	N1-C6-O6	5.62	123.27	119.90
36	1	262	U	N3-C2-O2	5.62	126.13	122.20
36	1	425	G	N1-C2-N2	-5.62	111.14	116.20
36	1	614	C	N3-C4-C5	5.62	124.15	121.90
36	5	2271	A	C6-C5-N7	5.62	136.23	132.30
36	5	2707	C	C5-C6-N1	-5.62	118.19	121.00
36	5	3088	G	N1-C6-O6	5.62	123.27	119.90
36	5	3374	U	C5-C6-N1	-5.62	119.89	122.70
36	1	212	G	N3-C4-N9	5.62	129.37	126.00
1	6	306	U	C5-C6-N1	-5.62	119.89	122.70
36	5	1475	A	O5'-P-OP2	-5.62	100.65	105.70
36	1	25	U	N3-C4-O4	5.61	123.33	119.40
36	1	285	A	N9-C4-C5	-5.61	103.56	105.80
36	1	1020	G	N9-C4-C5	-5.61	103.15	105.40
36	1	2899	C	C6-N1-C1'	-5.61	114.06	120.80
36	5	1421	G	O5'-P-OP1	-5.61	100.65	105.70
36	1	2247	G	C8-N9-C1'	-5.61	119.70	127.00
36	5	137	G	C6-C5-N7	-5.61	127.03	130.40
36	5	790	U	N1-C2-N3	5.61	118.27	114.90
1	6	273	G	O5'-P-OP1	-5.61	100.65	105.70
36	5	360	G	N3-C4-C5	-5.61	125.80	128.60
36	5	1046	A	N1-C2-N3	5.61	132.11	129.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	1379	G	C8-N9-C1'	-5.61	119.71	127.00
36	5	2585	G	C2-N3-C4	5.61	114.70	111.90
36	5	2727	A	N9-C4-C5	5.61	108.04	105.80
36	5	2813	A	C4-C5-C6	5.61	119.81	117.00
36	1	2341	A	C8-N9-C4	5.61	108.04	105.80
36	1	2899	C	C4-C5-C6	5.61	120.20	117.40
38	4	22	U	N1-C2-N3	5.61	118.26	114.90
36	1	1890	U	C6-N1-C2	5.60	124.36	121.00
1	6	1789	G	C5-C6-O6	-5.60	125.24	128.60
36	1	1373	A	C8-N9-C4	-5.60	103.56	105.80
36	1	2730	G	C8-N9-C1'	5.60	134.28	127.00
1	6	390	G	O5'-P-OP1	-5.60	100.66	105.70
12	c0	88	PRO	N-CA-CB	5.60	110.02	103.30
36	5	1513	G	C8-N9-C4	-5.60	104.16	106.40
37	7	7	G	N1-C6-O6	-5.60	116.54	119.90
64	n8	73	LEU	CA-CB-CG	5.60	128.19	115.30
1	2	1733	C	C5-C4-N4	-5.60	116.28	120.20
36	1	1045	C	OP2-P-O3'	5.60	117.52	105.20
1	2	1456	C	O4'-C1'-N1	5.60	112.68	108.20
36	1	3269	U	C5-C4-O4	5.60	129.26	125.90
36	5	1178	G	C4-C5-C6	5.60	122.16	118.80
67	o1	20	LEU	CA-CB-CG	-5.60	102.42	115.30
36	1	1100	U	C2-N3-C4	-5.60	123.64	127.00
36	1	1139	G	C5-C6-O6	5.60	131.96	128.60
36	1	2130	G	C8-N9-C4	-5.60	104.16	106.40
36	1	3204	C	C2-N1-C1'	5.60	124.96	118.80
36	5	750	G	OP2-P-O3'	5.60	117.51	105.20
1	2	80	A	O5'-P-OP1	-5.60	100.66	105.70
1	6	160	C	N1-C2-O2	5.60	122.26	118.90
36	1	304	G	C4-C5-N7	-5.59	108.56	110.80
38	4	40	A	C5-C6-N6	-5.59	119.22	123.70
1	6	30	G	N3-C4-N9	-5.59	122.64	126.00
1	6	1280	C	C6-N1-C2	-5.59	118.06	120.30
36	5	644	G	C5-C6-O6	5.59	131.96	128.60
36	5	919	U	O5'-P-OP2	-5.59	100.67	105.70
36	5	2732	G	N1-C6-O6	5.59	123.26	119.90
38	8	84	C	O4'-C1'-N1	5.59	112.67	108.20
36	1	2747	A	C8-N9-C4	-5.59	103.56	105.80
36	5	633	C	C4-C5-C6	5.59	120.20	117.40
1	2	1180	C	N3-C2-O2	-5.59	117.99	121.90
36	1	648	C	C6-N1-C1'	-5.59	114.09	120.80
36	1	1807	G	C6-C5-N7	-5.59	127.05	130.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2935	U	N3-C4-C5	-5.59	111.25	114.60
1	6	144	U	C6-N1-C2	-5.59	117.65	121.00
1	6	453	U	N1-C2-O2	5.59	126.71	122.80
36	5	406	G	N1-C6-O6	-5.59	116.55	119.90
36	5	2818	U	C5-C6-N1	5.59	125.50	122.70
40	l3	178	LEU	CA-CB-CG	5.59	128.16	115.30
18	C6	40	GLU	C-N-CA	5.59	145.48	122.00
36	1	884	A	OP1-P-O3'	5.59	117.50	105.20
36	1	1122	U	N1-C2-N3	5.59	118.25	114.90
36	1	2606	G	N1-C2-N2	-5.59	111.17	116.20
36	1	2727	A	C4-C5-N7	-5.59	107.91	110.70
36	1	2948	C	N3-C4-C5	5.59	124.14	121.90
36	5	596	C	C6-N1-C2	-5.59	118.06	120.30
36	1	51	A	N7-C8-N9	5.59	116.59	113.80
36	1	880	G	N7-C8-N9	-5.59	110.31	113.10
36	5	780	A	O5'-P-OP1	-5.59	100.67	105.70
36	5	887	G	C6-C5-N7	-5.59	127.05	130.40
36	5	2946	A	OP2-P-O3'	5.59	117.49	105.20
36	1	1838	G	N9-C4-C5	-5.58	103.17	105.40
36	1	2276	G	C5-C6-O6	-5.58	125.25	128.60
1	6	1644	C	N3-C2-O2	-5.58	117.99	121.90
36	5	1468	A	N9-C4-C5	-5.58	103.57	105.80
1	2	275	C	C6-N1-C2	-5.58	118.07	120.30
1	2	1735	U	C5-C4-O4	5.58	129.25	125.90
36	1	3112	G	N3-C4-C5	-5.58	125.81	128.60
36	1	885	U	C6-N1-C2	5.58	124.35	121.00
36	1	1307	G	C6-C5-N7	5.58	133.75	130.40
36	1	3015	G	N1-C6-O6	5.58	123.25	119.90
1	6	163	G	C5-N7-C8	-5.58	101.51	104.30
1	2	1146	G	C8-N9-C4	-5.58	104.17	106.40
36	1	1709	C	N3-C4-C5	-5.58	119.67	121.90
1	6	1537	C	O4'-C1'-N1	5.58	112.66	108.20
36	5	2820	A	N7-C8-N9	5.58	116.59	113.80
36	5	3119	U	N3-C4-O4	5.58	123.31	119.40
1	2	389	G	C8-N9-C4	-5.58	104.17	106.40
36	1	857	G	C6-C5-N7	-5.58	127.05	130.40
36	5	1628	C	C6-N1-C2	-5.58	118.07	120.30
36	5	3013	U	N1-C2-O2	5.58	126.70	122.80
36	1	632	G	C4-C5-N7	5.58	113.03	110.80
36	5	1480	G	N1-C6-O6	5.58	123.25	119.90
36	5	2362	C	O5'-P-OP2	-5.58	100.68	105.70
36	5	2718	U	N1-C2-N3	5.58	118.25	114.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	2	704	C	C2-N1-C1'	5.57	124.93	118.80
36	1	86	G	O5'-P-OP1	5.57	117.39	110.70
36	1	1517	G	O5'-P-OP2	-5.57	100.68	105.70
36	1	1831	U	C5-C6-N1	5.57	125.49	122.70
1	6	91	G	N3-C4-C5	5.57	131.39	128.60
36	5	873	C	P-O3'-C3'	5.57	126.39	119.70
36	5	2993	G	N1-C6-O6	5.57	123.25	119.90
36	5	2891	U	N3-C2-O2	-5.57	118.30	122.20
36	1	22	G	C8-N9-C4	-5.57	104.17	106.40
1	6	1614	A	C5-N7-C8	-5.57	101.11	103.90
36	5	822	G	N3-C4-C5	5.57	131.38	128.60
36	5	841	A	C8-N9-C4	5.57	108.03	105.80
36	5	2199	G	C4-C5-N7	5.57	113.03	110.80
36	1	3140	G	N1-C6-O6	5.57	123.24	119.90
1	6	1560	U	N1-C2-O2	5.57	126.70	122.80
1	2	985	G	N3-C4-C5	-5.57	125.82	128.60
36	1	884	A	C6-C5-N7	-5.57	128.40	132.30
36	1	913	A	N7-C8-N9	5.57	116.58	113.80
36	5	574	U	C6-N1-C2	5.57	124.34	121.00
36	5	970	A	C8-N9-C4	5.57	108.03	105.80
36	5	1181	U	C4-C5-C6	5.57	123.04	119.70
36	5	2940	A	N9-C4-C5	5.57	108.03	105.80
36	1	123	A	C5-N7-C8	-5.57	101.12	103.90
36	1	2643	A	N9-C4-C5	-5.57	103.57	105.80
20	c8	18	LEU	CA-CB-CG	5.57	128.10	115.30
36	5	1331	U	N3-C2-O2	-5.57	118.30	122.20
36	5	2858	U	N3-C4-C5	-5.57	111.26	114.60
36	1	942	U	OP1-P-OP2	-5.56	111.25	119.60
36	1	655	C	N3-C4-C5	-5.56	119.67	121.90
36	1	1495	U	C2-N3-C4	-5.56	123.66	127.00
36	5	567	G	C4-N9-C1'	5.56	133.73	126.50
36	5	907	G	C8-N9-C4	5.56	108.62	106.40
36	5	1205	A	C4-C5-C6	5.56	119.78	117.00
36	5	1556	C	C2-N1-C1'	5.56	124.92	118.80
37	7	30	G	C4-N9-C1'	5.56	133.73	126.50
1	2	720	G	P-O3'-C3'	5.56	126.37	119.70
1	2	1240	U	O5'-P-OP2	-5.56	100.70	105.70
1	2	1541	G	C8-N9-C4	-5.56	104.18	106.40
36	1	898	U	N1-C2-N3	-5.56	111.56	114.90
1	6	1645	G	C6-C5-N7	5.56	133.74	130.40
1	6	1744	A	C5-N7-C8	-5.56	101.12	103.90
36	5	137	G	N1-C6-O6	5.56	123.24	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	2274	U	C2-N1-C1'	5.56	124.37	117.70
36	5	2928	C	C5-C4-N4	-5.56	116.31	120.20
36	1	1851	G	C6-C5-N7	-5.56	127.07	130.40
36	5	3130	A	O5'-P-OP1	-5.56	100.70	105.70
19	C7	73	LEU	CA-CB-CG	5.55	128.07	115.30
1	6	1672	G	N3-C2-N2	5.55	123.79	119.90
1	6	1781	A	C8-N9-C4	-5.55	103.58	105.80
36	5	141	C	N1-C2-O2	-5.55	115.57	118.90
36	5	562	C	C6-N1-C2	-5.55	118.08	120.30
36	5	630	A	C8-N9-C4	5.55	108.02	105.80
36	5	1450	G	N3-C4-C5	5.55	131.38	128.60
36	1	27	C	C6-N1-C2	-5.55	118.08	120.30
36	1	612	U	C5-C6-N1	-5.55	119.92	122.70
36	1	1126	G	C6-C5-N7	-5.55	127.07	130.40
36	5	3215	A	N1-C6-N6	5.55	121.93	118.60
36	1	2192	C	O5'-P-OP2	-5.55	100.71	105.70
36	5	947	G	O5'-P-OP2	-5.55	100.70	105.70
36	5	1002	A	C2-N3-C4	-5.55	107.83	110.60
36	5	1911	A	C5-C6-N1	-5.55	114.92	117.70
36	5	2607	G	P-O3'-C3'	5.55	126.36	119.70
36	1	198	A	N1-C6-N6	5.55	121.93	118.60
36	5	1178	G	N7-C8-N9	5.55	115.87	113.10
36	1	345	G	C8-N9-C1'	-5.55	119.79	127.00
36	1	2288	G	C4-N9-C1'	5.55	133.71	126.50
37	3	91	G	N3-C2-N2	-5.55	116.02	119.90
36	5	1151	U	N3-C4-C5	-5.55	111.27	114.60
36	1	776	U	C4-C5-C6	5.54	123.03	119.70
36	1	2395	G	C6-C5-N7	-5.54	127.07	130.40
1	6	1653	C	N3-C4-N4	5.54	121.88	118.00
36	1	1366	A	OP2-P-O3'	5.54	117.39	105.20
36	1	2820	A	OP1-P-OP2	-5.54	111.29	119.60
38	4	85	G	C8-N9-C4	-5.54	104.18	106.40
36	5	2818	U	P-O3'-C3'	5.54	126.35	119.70
1	2	1611	A	C2-N3-C4	-5.54	107.83	110.60
36	1	1903	U	N3-C4-C5	-5.54	111.28	114.60
45	L8	158	ASP	C-N-CD	5.54	140.04	128.40
36	5	1310	G	N3-C4-N9	5.54	129.32	126.00
36	1	1049	C	N3-C4-C5	5.54	124.12	121.90
36	1	2134	G	C5-C6-O6	5.54	131.92	128.60
36	1	2276	G	C4-C5-N7	5.54	113.02	110.80
36	5	1485	G	C8-N9-C4	5.54	108.62	106.40
1	2	1059	U	O5'-P-OP1	-5.54	100.72	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	907	G	C5-C6-O6	-5.54	125.28	128.60
36	1	968	G	C8-N9-C1'	-5.54	119.80	127.00
36	5	1367	G	C6-C5-N7	-5.54	127.08	130.40
36	5	2826	U	N3-C2-O2	-5.54	118.32	122.20
36	1	649	A	C2-N3-C4	-5.54	107.83	110.60
36	1	1152	G	O5'-P-OP1	-5.54	100.72	105.70
36	5	1906	G	C6-C5-N7	-5.54	127.08	130.40
36	1	49	A	N9-C4-C5	-5.53	103.59	105.80
36	1	794	U	O5'-P-OP2	-5.53	100.72	105.70
36	1	1518	U	N3-C4-O4	5.53	123.27	119.40
36	1	59	G	C6-C5-N7	-5.53	127.08	130.40
36	1	1514	G	C4-N9-C1'	5.53	133.69	126.50
36	1	2699	G	C5-C6-O6	-5.53	125.28	128.60
36	5	412	G	N7-C8-N9	5.53	115.87	113.10
36	5	567	G	N9-C4-C5	-5.53	103.19	105.40
36	5	1590	G	O5'-P-OP1	-5.53	100.72	105.70
36	5	2904	U	C2-N1-C1'	5.53	124.34	117.70
36	5	3294	A	N1-C6-N6	-5.53	115.28	118.60
1	2	1156	C	C6-N1-C2	5.53	122.51	120.30
36	1	1331	U	C5-C6-N1	-5.53	119.94	122.70
36	1	2190	U	C6-N1-C2	-5.53	117.68	121.00
1	6	576	G	N1-C6-O6	5.53	123.22	119.90
36	5	924	G	C2-N3-C4	-5.53	109.14	111.90
36	5	2190	U	C6-N1-C2	-5.53	117.68	121.00
36	5	3202	G	N1-C6-O6	-5.53	116.58	119.90
36	5	3239	G	O5'-P-OP2	-5.53	100.72	105.70
36	1	716	A	C5-C6-N6	-5.53	119.28	123.70
36	1	773	G	C4-C5-N7	-5.53	108.59	110.80
36	1	1416	C	C2-N3-C4	-5.53	117.14	119.90
1	6	604	A	C8-N9-C4	-5.53	103.59	105.80
36	5	1889	G	N1-C6-O6	5.53	123.22	119.90
15	C3	22	ALA	C-N-CA	5.53	145.21	122.00
36	1	1329	U	N3-C4-O4	5.53	123.27	119.40
36	1	2174	G	N7-C8-N9	5.53	115.86	113.10
36	5	658	G	C8-N9-C4	-5.53	104.19	106.40
36	5	2816	G	O4'-C1'-N9	5.53	112.62	108.20
36	1	421	G	N3-C4-N9	5.52	129.31	126.00
36	1	816	A	C8-N9-C4	-5.52	103.59	105.80
36	1	2386	A	C4-C5-N7	5.52	113.46	110.70
1	6	1031	U	C2-N1-C1'	-5.52	111.07	117.70
36	1	22	G	N3-C4-C5	-5.52	125.84	128.60
36	1	423	A	OP2-P-O3'	5.52	117.35	105.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	682	U	C2-N1-C1'	-5.52	111.07	117.70
36	1	2277	C	C4-C5-C6	5.52	120.16	117.40
36	5	835	G	N1-C6-O6	-5.52	116.59	119.90
36	5	2912	G	OP1-P-OP2	5.52	127.89	119.60
36	1	3106	A	N1-C6-N6	5.52	121.91	118.60
36	1	229	G	C5-C6-O6	-5.52	125.29	128.60
36	1	1379	G	N1-C2-N3	5.52	127.21	123.90
36	1	1394	A	C4-C5-C6	-5.52	114.24	117.00
36	1	3046	A	C2-N3-C4	-5.52	107.84	110.60
1	6	1074	G	C5-C6-N1	-5.52	108.74	111.50
36	5	754	G	N1-C6-O6	-5.52	116.59	119.90
36	5	1405	U	C5-C4-O4	5.52	129.21	125.90
36	5	1444	G	OP2-P-O3'	5.52	117.34	105.20
36	5	2359	C	N3-C4-N4	5.52	121.86	118.00
36	5	2409	G	C5-C6-O6	-5.52	125.29	128.60
36	5	2552	C	C6-N1-C1'	-5.52	114.18	120.80
36	1	1057	A	C4-C5-C6	-5.52	114.24	117.00
36	1	1484	U	OP1-P-O3'	5.52	117.34	105.20
1	6	467	G	C8-N9-C1'	-5.52	119.83	127.00
36	5	607	A	N9-C4-C5	5.52	108.01	105.80
36	5	2123	G	N3-C4-C5	-5.52	125.84	128.60
36	5	3396	U	O4'-C1'-N1	5.52	112.61	108.20
37	7	28	C	O5'-P-OP1	-5.52	100.73	105.70
36	1	2408	U	N3-C2-O2	-5.52	118.34	122.20
36	5	1332	A	C4-N9-C1'	5.52	136.23	126.30
40	l3	4	ARG	NE-CZ-NH1	5.52	123.06	120.30
36	1	866	A	N1-C6-N6	5.51	121.91	118.60
36	1	2837	A	N1-C6-N6	5.51	121.91	118.60
1	6	168	A	C8-N9-C4	-5.51	103.59	105.80
36	1	1352	A	OP1-P-O3'	5.51	117.33	105.20
36	1	1429	G	C6-N1-C2	-5.51	121.79	125.10
36	1	2680	A	N1-C6-N6	5.51	121.91	118.60
36	5	2947	G	N1-C6-O6	5.51	123.21	119.90
36	1	187	A	C8-N9-C4	-5.51	103.59	105.80
1	6	338	C	N3-C4-C5	-5.51	119.70	121.90
36	5	668	G	N1-C6-O6	-5.51	116.59	119.90
1	2	822	U	C6-N1-C2	-5.51	117.69	121.00
38	4	94	C	N3-C4-C5	5.51	124.10	121.90
36	5	3109	G	C4-C5-N7	5.51	113.00	110.80
36	1	345	G	C4-C5-C6	5.51	122.10	118.80
36	5	964	G	C5-C6-O6	-5.51	125.30	128.60
36	5	2629	U	N3-C4-O4	5.51	123.25	119.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	2777	G	C6-C5-N7	5.51	133.70	130.40
1	2	1176	G	C6-C5-N7	-5.50	127.10	130.40
36	1	883	A	C6-N1-C2	-5.50	115.30	118.60
36	1	1441	G	N3-C4-N9	-5.50	122.70	126.00
36	1	1791	C	N3-C4-C5	-5.50	119.70	121.90
38	4	25	G	C4-C5-N7	-5.50	108.60	110.80
36	5	936	A	C4-C5-C6	5.50	119.75	117.00
36	1	53	G	N3-C4-C5	-5.50	125.85	128.60
1	6	1314	U	N1-C2-O2	5.50	126.65	122.80
36	5	1899	G	N9-C4-C5	5.50	107.60	105.40
36	5	2169	G	C2-N3-C4	5.50	114.65	111.90
36	1	1139	G	C4-C5-N7	-5.50	108.60	110.80
36	1	1148	G	C4-C5-C6	5.50	122.10	118.80
36	1	1307	G	C5-C6-O6	5.50	131.90	128.60
36	1	1417	G	N3-C4-N9	-5.50	122.70	126.00
36	1	2856	G	C5-C6-N1	-5.50	108.75	111.50
36	1	3201	C	N3-C4-C5	-5.50	119.70	121.90
36	5	658	G	C4-C5-N7	5.50	113.00	110.80
36	5	1367	G	C4-N9-C1'	5.50	133.65	126.50
36	5	1845	G	N3-C4-C5	-5.50	125.85	128.60
36	1	2412	G	C5-C6-N1	5.50	114.25	111.50
36	5	1331	U	N1-C2-N3	5.50	118.20	114.90
36	1	3390	G	N1-C6-O6	5.50	123.20	119.90
36	1	1868	G	N3-C4-N9	5.50	129.30	126.00
36	1	1940	G	N1-C2-N2	-5.50	111.25	116.20
1	6	1117	U	N3-C4-C5	-5.50	111.30	114.60
1	6	1648	A	C8-N9-C4	5.50	108.00	105.80
36	5	1409	G	N3-C4-C5	-5.50	125.85	128.60
1	6	1463	C	C6-N1-C2	5.50	122.50	120.30
36	5	2860	U	N1-C2-O2	5.50	126.65	122.80
37	7	32	U	C6-N1-C2	5.50	124.30	121.00
1	2	1645	G	N3-C4-N9	5.49	129.30	126.00
36	1	519	A	C6-C5-N7	-5.49	128.46	132.30
36	1	883	A	N1-C2-N3	5.49	132.05	129.30
38	4	52	A	N9-C4-C5	5.49	108.00	105.80
54	M8	49	LEU	CA-CB-CG	5.49	127.94	115.30
36	5	931	C	N3-C2-O2	-5.49	118.06	121.90
36	5	1384	U	C2-N1-C1'	5.49	124.29	117.70
36	5	1587	A	N9-C4-C5	-5.49	103.60	105.80
36	5	3181	C	N3-C2-O2	-5.49	118.06	121.90
36	1	818	C	N3-C4-C5	-5.49	119.70	121.90
36	1	954	U	C6-N1-C2	-5.49	117.70	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	2	210	A	N1-C6-N6	5.49	121.89	118.60
1	6	630	A	N7-C8-N9	-5.49	111.06	113.80
36	5	1354	G	C6-C5-N7	-5.49	127.11	130.40
36	5	3208	G	N1-C6-O6	5.49	123.19	119.90
36	1	1733	G	C4-N9-C1'	5.49	133.64	126.50
36	1	2376	G	C8-N9-C4	-5.49	104.20	106.40
36	5	3149	G	N3-C4-N9	-5.49	122.71	126.00
38	8	108	C	C6-N1-C2	-5.49	118.11	120.30
1	2	428	A	C8-N9-C4	-5.49	103.61	105.80
36	1	662	U	N3-C2-O2	-5.49	118.36	122.20
36	1	908	G	O4'-C1'-N9	-5.49	103.81	108.20
36	1	154	U	C2-N1-C1'	-5.49	111.12	117.70
36	1	636	C	C6-N1-C1'	-5.49	114.22	120.80
1	6	637	C	C6-N1-C2	-5.49	118.11	120.30
36	5	610	G	C2-N3-C4	5.49	114.64	111.90
38	8	42	G	C5-C6-O6	-5.49	125.31	128.60
1	2	1418	G	C4-N9-C1'	5.48	133.63	126.50
1	2	1109	G	C5-C6-N1	-5.48	108.76	111.50
36	1	3140	G	C6-C5-N7	-5.48	127.11	130.40
1	6	569	C	N3-C4-C5	-5.48	119.71	121.90
1	6	1742	U	C2-N1-C1'	5.48	124.28	117.70
36	5	942	U	N1-C2-N3	5.48	118.19	114.90
36	5	1900	A	C4-C5-N7	5.48	113.44	110.70
36	5	2624	G	N1-C6-O6	5.48	123.19	119.90
38	8	4	C	C2-N1-C1'	5.48	124.83	118.80
36	1	1375	G	N7-C8-N9	5.48	115.84	113.10
36	1	2624	G	C4-C5-N7	5.48	112.99	110.80
44	17	179	LEU	CA-CB-CG	5.48	127.91	115.30
36	1	787	G	C2-N3-C4	5.48	114.64	111.90
36	5	941	G	OP1-P-O3'	5.48	117.25	105.20
36	1	1844	C	C2-N1-C1'	-5.48	112.78	118.80
36	5	2894	C	N3-C4-C5	5.48	124.09	121.90
36	1	898	U	N1-C2-O2	5.48	126.63	122.80
36	1	1295	G	O5'-P-OP1	-5.48	100.77	105.70
36	1	85	A	C5-C6-N1	-5.47	114.96	117.70
36	1	964	G	N1-C2-N2	5.47	121.13	116.20
36	1	2134	G	N1-C6-O6	-5.47	116.61	119.90
36	1	2358	A	C5-C6-N6	-5.47	119.32	123.70
1	6	384	G	C2-N3-C4	-5.47	109.16	111.90
36	5	1332	A	C4-C5-C6	5.47	119.74	117.00
36	5	1384	U	N3-C2-O2	-5.47	118.37	122.20
36	5	2699	G	C8-N9-C4	5.47	108.59	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	345	G	C4-N9-C1'	5.47	133.61	126.50
36	1	648	C	N3-C4-N4	5.47	121.83	118.00
36	1	1201	C	N1-C2-O2	5.47	122.18	118.90
1	6	7	G	C8-N9-C4	-5.47	104.21	106.40
1	6	1764	C	N3-C4-C5	5.47	124.09	121.90
36	5	907	G	N1-C6-O6	5.47	123.18	119.90
36	5	3245	A	N7-C8-N9	5.47	116.54	113.80
36	5	2160	G	N9-C4-C5	-5.47	103.21	105.40
36	1	1855	U	C5-C6-N1	5.47	125.43	122.70
36	1	2201	G	O5'-P-OP2	-5.47	100.78	105.70
36	1	2643	A	OP2-P-O3'	5.47	117.23	105.20
1	6	272	U	N1-C2-O2	5.47	126.63	122.80
36	5	424	G	N9-C4-C5	-5.47	103.21	105.40
36	5	93	C	C6-N1-C2	5.47	122.49	120.30
36	5	1869	C	C2-N1-C1'	-5.47	112.78	118.80
36	5	2936	A	C5-C6-N1	5.47	120.43	117.70
36	1	938	C	C2-N1-C1'	5.47	124.81	118.80
36	1	3050	U	N3-C2-O2	-5.47	118.37	122.20
44	L7	163	LEU	CA-CB-CG	-5.47	102.73	115.30
36	5	917	A	N7-C8-N9	5.47	116.53	113.80
36	5	2398	A	N9-C4-C5	5.47	107.99	105.80
36	5	2711	C	C6-N1-C2	-5.47	118.11	120.30
36	5	2824	G	C2-N3-C4	-5.47	109.17	111.90
36	1	908	G	C4-N9-C1'	5.46	133.60	126.50
38	4	101	U	C2-N1-C1'	5.46	124.26	117.70
1	6	323	A	N1-C6-N6	-5.46	115.32	118.60
1	6	1340	U	N3-C2-O2	-5.46	118.38	122.20
36	5	1317	A	C5-C6-N6	-5.46	119.33	123.70
36	5	3149	G	C2-N3-C4	-5.46	109.17	111.90
44	17	83	LEU	CA-CB-CG	5.46	127.87	115.30
36	5	197	G	N9-C4-C5	-5.46	103.22	105.40
36	5	358	G	N3-C4-N9	-5.46	122.72	126.00
36	5	2884	C	N3-C4-N4	5.46	121.82	118.00
1	2	6	G	N3-C4-C5	-5.46	125.87	128.60
36	1	75	G	O5'-P-OP2	-5.46	100.78	105.70
36	1	1766	G	C4-N9-C1'	5.46	133.60	126.50
65	N9	20	GLY	N-CA-C	5.46	126.75	113.10
36	5	716	A	C8-N9-C4	5.46	107.98	105.80
1	2	1200	G	C5-C6-O6	-5.46	125.32	128.60
36	1	2380	U	C2-N3-C4	-5.46	123.72	127.00
36	5	2549	G	C6-C5-N7	-5.46	127.12	130.40
37	7	116	C	C6-N1-C2	-5.46	118.12	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	2	1458	G	C8-N9-C1'	-5.46	119.90	127.00
1	2	1778	G	C5-N7-C8	-5.46	101.57	104.30
36	1	2639	G	C4-C5-N7	5.46	112.98	110.80
38	4	83	C	C2-N1-C1'	5.46	124.81	118.80
1	6	1742	U	C5-C6-N1	5.46	125.43	122.70
1	6	1791	A	C5-C6-N6	-5.46	119.33	123.70
25	d3	33	LEU	CB-CG-CD1	-5.46	101.72	111.00
36	5	3343	G	N3-C4-C5	-5.46	125.87	128.60
37	7	104	A	N1-C6-N6	5.46	121.88	118.60
36	5	208	C	C6-N1-C2	-5.46	118.12	120.30
36	5	3278	C	N3-C4-C5	5.46	124.08	121.90
36	1	96	G	N1-C2-N3	5.45	127.17	123.90
36	1	1162	U	N3-C2-O2	-5.45	118.38	122.20
36	1	1190	A	C4-N9-C1'	5.45	136.12	126.30
36	1	1917	C	C6-N1-C2	5.45	122.48	120.30
36	1	2625	C	O5'-P-OP1	-5.45	100.79	105.70
37	3	83	U	N1-C1'-C2'	-5.45	106.00	112.00
1	6	542	A	C6-C5-N7	-5.45	128.48	132.30
1	6	570	A	O5'-P-OP2	-5.45	100.79	105.70
36	5	776	U	N1-C2-N3	5.45	118.17	114.90
36	5	2360	C	C5-C6-N1	5.45	123.73	121.00
1	2	318	U	C2-N1-C1'	-5.45	111.16	117.70
1	6	1619	C	C5-C6-N1	5.45	123.73	121.00
36	5	1377	G	N3-C4-N9	5.45	129.27	126.00
1	2	554	C	C2-N3-C4	5.45	122.62	119.90
1	2	1194	A	N7-C8-N9	5.45	116.53	113.80
36	1	404	G	O5'-P-OP2	-5.45	100.79	105.70
36	1	512	U	C5-C6-N1	-5.45	119.97	122.70
36	1	2143	A	N1-C6-N6	5.45	121.87	118.60
1	6	403	G	N3-C4-N9	-5.45	122.73	126.00
1	6	1274	C	C2-N1-C1'	5.45	124.80	118.80
36	5	920	A	O5'-P-OP1	5.45	117.24	110.70
36	5	2132	C	C6-N1-C2	-5.45	118.12	120.30
36	5	2751	G	C8-N9-C4	-5.45	104.22	106.40
36	5	3116	G	C5-C6-O6	-5.45	125.33	128.60
36	1	2662	G	C2-N3-C4	-5.45	109.18	111.90
36	1	2687	G	N1-C6-O6	-5.45	116.63	119.90
38	4	40	A	N1-C6-N6	5.45	121.87	118.60
1	6	1560	U	C2-N1-C1'	5.45	124.24	117.70
36	5	816	A	N1-C6-N6	-5.45	115.33	118.60
36	5	2123	G	C2-N3-C4	5.45	114.62	111.90
36	1	580	C	N3-C2-O2	5.45	125.71	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	1192	C	C2-N3-C4	5.45	122.62	119.90
36	5	61	A	C5-C6-N6	5.45	128.06	123.70
36	5	210	U	C5-C6-N1	-5.45	119.98	122.70
36	5	1048	A	C5-N7-C8	5.45	106.62	103.90
36	5	2992	U	N1-C2-O2	5.45	126.61	122.80
37	7	32	U	N3-C2-O2	5.45	126.01	122.20
1	6	1600	A	C5-N7-C8	-5.44	101.18	103.90
41	14	179	LEU	CA-CB-CG	5.44	127.82	115.30
1	2	992	A	C4-C5-N7	5.44	113.42	110.70
36	1	1389	G	N3-C2-N2	5.44	123.71	119.90
36	1	2342	U	C2-N1-C1'	-5.44	111.17	117.70
36	1	2976	A	N1-C6-N6	-5.44	115.33	118.60
36	5	698	U	C5-C4-O4	5.44	129.16	125.90
36	5	903	U	C5-C4-O4	5.44	129.17	125.90
36	5	1130	A	C2-N3-C4	5.44	113.32	110.60
37	7	41	G	N1-C6-O6	5.44	123.17	119.90
38	8	40	A	C8-N9-C4	-5.44	103.62	105.80
1	2	1014	G	N1-C6-O6	5.44	123.16	119.90
36	1	380	U	C6-N1-C2	-5.44	117.74	121.00
36	1	876	A	N9-C4-C5	-5.44	103.62	105.80
36	1	1473	G	C8-N9-C4	5.44	108.58	106.40
36	1	2273	G	C8-N9-C4	5.44	108.58	106.40
1	6	68	A	N1-C6-N6	5.44	121.86	118.60
1	6	1648	A	N1-C6-N6	5.44	121.86	118.60
20	c8	15	LEU	CA-CB-CG	5.44	127.81	115.30
36	5	1159	A	N3-C4-C5	5.44	130.61	126.80
36	1	22	G	C6-N1-C2	-5.44	121.84	125.10
36	1	1126	G	C5-C6-O6	-5.44	125.34	128.60
36	1	2808	A	C4-C5-N7	5.44	113.42	110.70
36	1	3364	C	O5'-P-OP1	-5.44	100.81	105.70
36	5	1373	A	OP2-P-O3'	5.44	117.17	105.20
36	1	717	C	C6-N1-C2	-5.44	118.12	120.30
36	1	816	A	C2-N3-C4	5.44	113.32	110.60
36	1	1547	G	N7-C8-N9	-5.44	110.38	113.10
36	1	2730	G	N3-C4-C5	5.44	131.32	128.60
1	6	100	A	N1-C6-N6	5.44	121.86	118.60
36	5	406	G	C5-C6-O6	5.44	131.86	128.60
36	5	3007	U	N3-C4-C5	5.44	117.86	114.60
36	5	3107	U	N1-C2-N3	5.44	118.16	114.90
36	5	3144	G	C4-N9-C1'	5.44	133.57	126.50
37	7	80	G	C4-N9-C1'	5.44	133.57	126.50
1	6	29	U	C4-C5-C6	5.44	122.96	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	2878	G	C5-C6-N1	5.44	114.22	111.50
36	5	2918	G	C8-N9-C4	-5.44	104.22	106.40
36	1	1306	G	C5-C6-N1	-5.43	108.78	111.50
36	1	2650	U	N1-C2-O2	-5.43	119.00	122.80
36	5	61	A	C2-N3-C4	-5.43	107.88	110.60
36	5	1211	U	N3-C4-C5	5.43	117.86	114.60
36	5	1476	G	N3-C2-N2	5.43	123.70	119.90
36	5	2748	A	N9-C4-C5	-5.43	103.63	105.80
1	2	577	G	N9-C4-C5	-5.43	103.23	105.40
36	1	1492	G	N3-C2-N2	5.43	123.70	119.90
36	1	2606	G	N3-C2-N2	5.43	123.70	119.90
1	6	2	A	C8-N9-C4	5.43	107.97	105.80
1	6	103	A	P-O3'-C3'	5.43	126.22	119.70
1	6	1297	G	C5-C6-O6	5.43	131.86	128.60
36	5	1305	U	N3-C4-C5	-5.43	111.34	114.60
36	5	2346	C	C5-C4-N4	-5.43	116.40	120.20
36	5	3043	C	C6-N1-C2	5.43	122.47	120.30
36	1	1495	U	N1-C2-O2	-5.43	119.00	122.80
36	5	1203	A	C5-C6-N6	-5.43	119.36	123.70
36	5	1561	G	O4'-C1'-N9	5.43	112.55	108.20
36	5	1885	U	N3-C2-O2	5.43	126.00	122.20
1	2	305	C	C2-N1-C1'	5.43	124.77	118.80
1	2	399	A	C8-N9-C4	5.43	107.97	105.80
1	2	1559	A	O4'-C1'-N9	5.43	112.54	108.20
36	1	2772	C	C2-N1-C1'	5.43	124.77	118.80
38	4	40	A	C6-C5-N7	-5.43	128.50	132.30
38	4	51	G	N1-C6-O6	5.43	123.16	119.90
1	6	29	U	N3-C2-O2	-5.43	118.40	122.20
36	5	1012	G	C8-N9-C4	5.43	108.57	106.40
1	2	334	G	N3-C4-C5	5.43	131.31	128.60
36	1	145	G	N1-C6-O6	5.43	123.16	119.90
36	5	666	A	C4-C5-N7	-5.43	107.99	110.70
38	8	17	A	N9-C4-C5	-5.43	103.63	105.80
36	1	364	G	N1-C6-O6	5.42	123.15	119.90
37	3	80	G	N7-C8-N9	5.42	115.81	113.10
36	5	3116	G	N1-C6-O6	5.42	123.16	119.90
36	1	2898	G	O4'-C1'-N9	-5.42	103.86	108.20
1	6	1760	G	N1-C6-O6	5.42	123.15	119.90
36	5	968	G	C8-N9-C1'	-5.42	119.95	127.00
36	5	1452	A	C8-N9-C4	5.42	107.97	105.80
36	5	1919	G	C8-N9-C4	-5.42	104.23	106.40
36	5	2145	A	C6-N1-C2	-5.42	115.35	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	1124	U	C6-N1-C2	-5.42	117.75	121.00
36	1	2376	G	N7-C8-N9	5.42	115.81	113.10
1	6	1614	A	O4'-C1'-N9	5.42	112.54	108.20
36	5	1056	U	OP2-P-O3'	5.42	117.12	105.20
36	5	2316	G	C5-C6-O6	5.42	131.85	128.60
1	6	163	G	N3-C4-C5	5.42	131.31	128.60
44	17	229	PHE	CB-CG-CD2	-5.42	117.01	120.80
36	1	169	U	OP1-P-O3'	5.42	117.12	105.20
36	1	558	U	N3-C2-O2	-5.42	118.41	122.20
36	1	1316	C	O5'-P-OP2	-5.42	100.82	105.70
1	6	17	C	C6-N1-C2	-5.42	118.13	120.30
1	6	778	G	C8-N9-C4	5.42	108.57	106.40
36	5	3309	G	C4-C5-C6	5.42	122.05	118.80
36	1	1205	A	C2-N3-C4	-5.42	107.89	110.60
36	1	2851	A	C5-C6-N1	-5.42	114.99	117.70
1	6	66	U	OP1-P-O3'	5.42	117.11	105.20
36	5	921	A	N9-C4-C5	5.42	107.97	105.80
36	5	934	G	C4-C5-N7	5.42	112.97	110.80
36	1	655	C	C6-N1-C2	-5.42	118.13	120.30
36	1	986	U	O5'-P-OP1	-5.42	100.83	105.70
36	1	1314	C	C4-C5-C6	-5.42	114.69	117.40
36	1	2763	U	O5'-P-OP2	-5.42	100.83	105.70
1	6	904	G	C4-N9-C1'	5.42	133.54	126.50
36	5	2804	A	N1-C2-N3	5.42	132.01	129.30
36	1	784	A	O4'-C1'-N9	5.41	112.53	108.20
36	1	988	U	C6-N1-C2	5.41	124.25	121.00
36	1	2983	C	C5-C6-N1	-5.41	118.29	121.00
36	5	388	G	N7-C8-N9	5.41	115.81	113.10
36	5	647	A	C4-C5-C6	5.41	119.71	117.00
36	5	1161	G	N3-C4-C5	-5.41	125.89	128.60
36	5	2941	A	C6-N1-C2	-5.41	115.35	118.60
36	1	2152	A	C5-C6-N1	5.41	120.41	117.70
36	5	351	A	N1-C6-N6	5.41	121.85	118.60
36	5	421	G	C6-C5-N7	-5.41	127.15	130.40
1	6	323	A	C8-N9-C4	-5.41	103.64	105.80
36	5	2801	A	C5-C6-N1	5.41	120.41	117.70
36	5	3315	G	C4-N9-C1'	5.41	133.53	126.50
1	2	973	A	N1-C2-N3	5.41	132.00	129.30
36	1	883	A	C8-N9-C4	-5.41	103.64	105.80
36	1	1414	G	C4-C5-N7	5.41	112.96	110.80
36	5	2283	G	N9-C4-C5	-5.41	103.24	105.40
36	1	928	C	C6-N1-C2	-5.41	118.14	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	2901	G	O5'-P-OP2	-5.41	100.83	105.70
1	2	351	C	N1-C2-O2	5.41	122.14	118.90
1	2	453	U	C6-N1-C1'	-5.41	113.63	121.20
1	2	1489	U	C6-N1-C2	-5.41	117.76	121.00
36	1	1939	G	C4-C5-C6	5.41	122.04	118.80
1	6	275	C	N1-C2-O2	5.41	122.14	118.90
1	6	1700	C	N1-C2-O2	5.41	122.14	118.90
36	5	1406	A	C8-N9-C4	-5.41	103.64	105.80
36	5	2649	A	C5-C6-N1	5.41	120.40	117.70
36	5	3182	G	OP1-P-OP2	-5.41	111.49	119.60
36	1	2760	C	C6-N1-C2	5.40	122.46	120.30
36	5	2893	C	N3-C4-N4	5.40	121.78	118.00
36	1	941	G	C8-N9-C4	-5.40	104.24	106.40
36	1	2372	A	C5-C6-N6	-5.40	119.38	123.70
36	1	3104	U	C6-N1-C2	5.40	124.24	121.00
36	1	3128	G	N9-C4-C5	-5.40	103.24	105.40
36	5	1338	C	C5-C4-N4	-5.40	116.42	120.20
36	5	2620	G	N3-C4-N9	5.40	129.24	126.00
61	n5	34	LEU	CA-CB-CG	5.40	127.73	115.30
1	2	1488	G	C8-N9-C4	-5.40	104.24	106.40
36	1	400	G	C8-N9-C4	-5.40	104.24	106.40
36	1	901	G	C6-C5-N7	-5.40	127.16	130.40
36	1	1341	U	C4-C5-C6	5.40	122.94	119.70
1	6	64	U	C5-C6-N1	-5.40	120.00	122.70
1	6	268	C	N1-C2-O2	5.40	122.14	118.90
1	6	558	U	O4'-C1'-N1	5.40	112.52	108.20
36	5	651	G	N3-C4-N9	5.40	129.24	126.00
36	5	1149	G	N1-C2-N3	5.40	127.14	123.90
36	1	922	U	N3-C4-C5	-5.40	111.36	114.60
36	5	2996	U	N1-C2-O2	5.40	126.58	122.80
37	7	9	C	O5'-P-OP1	-5.40	100.84	105.70
1	2	1793	G	N3-C4-C5	-5.40	125.90	128.60
36	1	2194	G	C5-C6-O6	-5.40	125.36	128.60
36	1	2404	A	C2-N3-C4	5.40	113.30	110.60
36	1	2653	C	N3-C4-C5	5.40	124.06	121.90
36	1	2817	A	C2-N3-C4	5.40	113.30	110.60
36	5	170	G	C8-N9-C1'	-5.40	119.98	127.00
36	5	647	A	C5-C6-N1	-5.40	115.00	117.70
36	5	1832	C	C6-N1-C2	5.40	122.46	120.30
36	5	3148	U	C5-C6-N1	-5.40	120.00	122.70
38	8	80	A	N7-C8-N9	5.40	116.50	113.80
1	2	638	U	O4'-C1'-N1	5.40	112.52	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	6	1012	U	N3-C4-C5	5.40	117.84	114.60
36	1	304	G	C6-C5-N7	5.39	133.64	130.40
36	1	331	G	N3-C2-N2	-5.39	116.12	119.90
36	1	1190	A	O4'-C1'-N9	-5.39	103.89	108.20
36	1	2176	U	O5'-P-OP1	-5.39	100.84	105.70
36	1	2647	A	C4-C5-C6	5.39	119.70	117.00
36	1	2719	U	C2-N1-C1'	-5.39	111.22	117.70
36	5	498	A	O5'-P-OP2	-5.39	100.84	105.70
36	5	1182	A	C5-C6-N6	-5.39	119.38	123.70
38	4	99	C	C6-N1-C2	5.39	122.46	120.30
1	6	334	G	N1-C2-N2	5.39	121.05	116.20
36	5	2704	A	N1-C6-N6	5.39	121.83	118.60
36	1	880	G	C8-N9-C4	5.39	108.56	106.40
36	1	1077	U	C6-N1-C2	5.39	124.23	121.00
36	1	1473	G	N9-C4-C5	-5.39	103.24	105.40
36	1	1836	C	C2-N1-C1'	5.39	124.73	118.80
36	5	776	U	C2-N3-C4	-5.39	123.77	127.00
36	5	2872	A	N3-C4-C5	5.39	130.57	126.80
37	7	5	G	C2-N3-C4	-5.39	109.20	111.90
1	2	1653	C	N3-C4-C5	-5.39	119.74	121.90
36	1	111	C	N1-C2-O2	-5.39	115.67	118.90
36	1	217	U	OP1-P-O3'	5.39	117.06	105.20
36	1	3266	G	N1-C6-O6	-5.39	116.67	119.90
1	6	60	U	C5-C6-N1	5.39	125.39	122.70
36	5	1611	G	C6-C5-N7	-5.39	127.17	130.40
36	5	2632	G	C2-N3-C4	-5.39	109.20	111.90
36	5	2699	G	N9-C4-C5	-5.39	103.24	105.40
36	5	3154	C	C6-N1-C1'	-5.39	114.33	120.80
48	m1	37	LEU	CA-CB-CG	-5.39	102.91	115.30
36	1	1868	G	N3-C2-N2	5.39	123.67	119.90
36	1	2970	C	C6-N1-C2	5.39	122.45	120.30
36	5	3076	C	C6-N1-C2	-5.39	118.14	120.30
1	2	1639	C	C2-N1-C1'	5.39	124.72	118.80
36	1	2321	A	N1-C6-N6	-5.39	115.37	118.60
36	5	2892	A	O5'-P-OP2	-5.39	100.85	105.70
36	1	1191	U	C5-C6-N1	-5.38	120.01	122.70
36	1	1497	C	N3-C4-C5	-5.38	119.75	121.90
36	1	2383	C	C4-C5-C6	-5.38	114.71	117.40
1	6	108	A	C4-C5-C6	5.38	119.69	117.00
36	5	2360	C	N3-C4-N4	5.38	121.77	118.00
36	5	2864	A	C6-C5-N7	-5.38	128.53	132.30
36	5	3264	G	O5'-P-OP1	-5.38	100.85	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	335	G	C4-C5-N7	5.38	112.95	110.80
36	1	1610	G	N1-C6-O6	5.38	123.13	119.90
1	6	1172	G	N3-C4-N9	5.38	129.23	126.00
36	1	1146	C	C5-C4-N4	-5.38	116.43	120.20
36	1	1929	G	C4-C5-N7	5.38	112.95	110.80
36	5	1161	G	C2-N3-C4	5.38	114.59	111.90
36	5	1519	G	C5-C6-O6	-5.38	125.37	128.60
36	5	2379	U	N3-C2-O2	-5.38	118.43	122.20
36	5	3183	A	N9-C4-C5	5.38	107.95	105.80
36	1	3002	C	C5-C6-N1	-5.38	118.31	121.00
1	6	384	G	N3-C4-N9	-5.38	122.77	126.00
38	8	64	U	C2-N1-C1'	5.38	124.16	117.70
36	1	2443	A	N1-C6-N6	5.38	121.83	118.60
36	5	924	G	O4'-C1'-N9	-5.38	103.90	108.20
36	5	1302	A	C5-C6-N1	-5.38	115.01	117.70
36	5	1830	G	C5-C6-N1	-5.38	108.81	111.50
36	5	1848	G	C5-C6-N1	5.38	114.19	111.50
36	5	2524	A	N7-C8-N9	5.38	116.49	113.80
36	5	2937	G	N3-C2-N2	-5.38	116.14	119.90
36	1	1352	A	P-O3'-C3'	5.38	126.15	119.70
36	1	1746	U	C6-N1-C2	-5.38	117.77	121.00
36	1	1863	G	C5-C6-O6	-5.38	125.38	128.60
36	5	1851	G	C6-C5-N7	-5.38	127.17	130.40
36	5	1878	G	C4-N9-C1'	5.38	133.49	126.50
36	5	1888	U	N1-C2-N3	5.38	118.13	114.90
36	5	2820	A	C5-N7-C8	-5.38	101.21	103.90
36	5	3033	A	C8-N9-C4	5.38	107.95	105.80
1	6	100	A	C4-C5-C6	5.38	119.69	117.00
36	5	1155	C	C5-C4-N4	-5.38	116.44	120.20
36	5	1407	A	O5'-P-OP2	-5.38	100.86	105.70
1	2	1284	C	N3-C2-O2	-5.37	118.14	121.90
36	1	1407	A	O5'-P-OP1	5.37	117.15	110.70
36	1	1792	C	N3-C4-C5	-5.37	119.75	121.90
36	1	2869	U	OP2-P-O3'	5.37	117.02	105.20
36	5	82	C	C6-N1-C2	-5.37	118.15	120.30
36	1	154	U	C5-C6-N1	-5.37	120.01	122.70
36	1	715	A	O4'-C1'-N9	5.37	112.50	108.20
36	1	936	A	N3-C4-C5	5.37	130.56	126.80
36	1	2616	C	N1-C2-O2	5.37	122.12	118.90
36	5	888	A	N9-C4-C5	-5.37	103.65	105.80
36	5	1556	C	N3-C2-O2	-5.37	118.14	121.90
36	5	3177	G	N1-C2-N3	5.37	127.12	123.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	80	G	N3-C4-N9	5.37	129.22	126.00
36	5	664	U	N3-C4-C5	-5.37	111.38	114.60
36	1	984	G	OP2-P-O3'	5.37	117.01	105.20
36	1	1315	U	C5-C6-N1	-5.37	120.02	122.70
1	6	1202	A	C2-N3-C4	5.37	113.28	110.60
36	5	1127	G	C5-C6-O6	5.37	131.82	128.60
36	5	1462	A	C2-N3-C4	-5.37	107.92	110.60
36	5	51	A	C4-C5-C6	5.37	119.68	117.00
36	5	646	A	N1-C2-N3	5.37	131.98	129.30
36	5	1296	C	N3-C4-N4	5.37	121.76	118.00
1	2	1568	C	C6-N1-C2	-5.37	118.15	120.30
1	6	1634	C	N3-C2-O2	-5.37	118.14	121.90
36	5	361	A	C8-N9-C4	5.37	107.95	105.80
36	5	3371	G	C5-C6-N1	-5.37	108.82	111.50
36	1	197	G	C5-N7-C8	-5.36	101.62	104.30
37	7	75	G	C4-N9-C1'	5.36	133.47	126.50
36	1	938	C	C6-N1-C2	-5.36	118.16	120.30
36	5	3218	A	C5-N7-C8	-5.36	101.22	103.90
36	1	1437	C	N3-C4-N4	5.36	121.75	118.00
47	MO	24	ARG	NE-CZ-NH1	5.36	122.98	120.30
1	6	1039	A	O4'-C1'-N9	5.36	112.49	108.20
36	5	329	U	N1-C2-O2	5.36	126.55	122.80
36	5	3108	G	C6-C5-N7	-5.36	127.18	130.40
36	1	1144	U	O5'-P-OP1	-5.36	100.88	105.70
36	1	2639	G	C4-C5-C6	5.36	122.02	118.80
1	6	1730	A	N1-C2-N3	5.36	131.98	129.30
36	5	2297	U	C2-N1-C1'	-5.36	111.27	117.70
36	1	1340	G	C5-C6-N1	5.36	114.18	111.50
36	5	1372	C	N3-C4-C5	-5.36	119.76	121.90
36	5	2624	G	C8-N9-C4	-5.36	104.26	106.40
36	5	2628	A	O5'-P-OP1	5.36	117.13	110.70
1	2	1100	G	N3-C4-N9	5.36	129.21	126.00
1	2	1274	C	N3-C2-O2	-5.36	118.15	121.90
36	1	1431	G	N1-C6-O6	-5.36	116.69	119.90
38	4	51	G	C5-C6-O6	-5.36	125.39	128.60
1	6	1285	U	C6-N1-C2	-5.36	117.79	121.00
1	6	1600	A	N7-C8-N9	5.36	116.48	113.80
36	5	1846	C	C5-C6-N1	-5.36	118.32	121.00
36	1	977	C	C6-N1-C2	5.35	122.44	120.30
36	1	1445	U	C2-N1-C1'	-5.35	111.28	117.70
36	5	360	G	C4-C5-N7	-5.35	108.66	110.80
36	5	1481	A	N7-C8-N9	5.35	116.48	113.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	326	U	N3-C4-O4	5.35	123.15	119.40
36	1	586	C	C6-N1-C2	5.35	122.44	120.30
36	5	405	U	C5-C6-N1	5.35	125.38	122.70
36	5	2976	A	OP2-P-O3'	5.35	116.98	105.20
36	5	3388	C	C5-C6-N1	-5.35	118.32	121.00
36	1	2807	U	C5-C4-O4	-5.35	122.69	125.90
36	5	656	A	N9-C4-C5	-5.35	103.66	105.80
36	1	608	A	C5-C6-N1	-5.35	115.03	117.70
36	1	1505	C	OP2-P-O3'	5.35	116.97	105.20
36	1	1834	U	C4-C5-C6	5.35	122.91	119.70
36	1	2407	C	N1-C2-O2	-5.35	115.69	118.90
1	6	391	A	N7-C8-N9	-5.35	111.13	113.80
1	6	1614	A	N3-C4-C5	5.35	130.54	126.80
1	6	1648	A	N9-C4-C5	-5.35	103.66	105.80
36	5	2301	U	N1-C2-N3	-5.35	111.69	114.90
36	5	2550	U	C5-C4-O4	5.35	129.11	125.90
37	7	80	G	N3-C4-N9	5.35	129.21	126.00
1	2	1216	C	O5'-P-OP2	-5.35	100.89	105.70
1	2	1785	U	N1-C2-N3	5.35	118.11	114.90
36	5	3000	A	N1-C6-N6	5.35	121.81	118.60
1	6	1486	G	O5'-P-OP2	-5.35	100.89	105.70
36	5	1116	G	C4-N9-C1'	5.35	133.45	126.50
36	1	324	A	N1-C2-N3	5.34	131.97	129.30
36	1	1835	A	O5'-P-OP1	-5.34	100.89	105.70
36	1	2385	G	C8-N9-C1'	5.34	133.95	127.00
37	3	25	G	N3-C4-N9	5.34	129.21	126.00
36	5	96	G	C8-N9-C4	5.34	108.54	106.40
36	5	1326	A	N7-C8-N9	-5.34	111.13	113.80
36	5	3154	C	C5-C6-N1	5.34	123.67	121.00
36	1	1434	G	C8-N9-C4	-5.34	104.26	106.40
36	5	364	G	N9-C4-C5	-5.34	103.26	105.40
1	2	321	C	N1-C2-O2	5.34	122.11	118.90
36	1	211	A	O5'-P-OP1	-5.34	100.89	105.70
36	1	922	U	C5-C4-O4	5.34	129.10	125.90
36	1	1064	A	O4'-C1'-N9	-5.34	103.93	108.20
36	1	2367	A	C5-C6-N1	5.34	120.37	117.70
1	6	1258	U	N3-C2-O2	-5.34	118.46	122.20
36	5	513	G	C5-N7-C8	5.34	106.97	104.30
36	5	1064	A	P-O3'-C3'	5.34	126.11	119.70
36	5	2375	G	N1-C6-O6	-5.34	116.69	119.90
36	1	277	G	C8-N9-C4	-5.34	104.26	106.40
36	1	1365	G	N1-C2-N2	-5.34	111.39	116.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2812	C	C4-C5-C6	5.34	120.07	117.40
36	1	2959	C	N3-C2-O2	5.34	125.64	121.90
36	5	280	U	O5'-P-OP2	-5.34	100.89	105.70
36	5	388	G	C6-C5-N7	-5.34	127.20	130.40
36	5	920	A	OP1-P-OP2	-5.34	111.59	119.60
36	5	1301	A	N3-C4-C5	-5.34	123.06	126.80
36	5	2818	U	C5'-C4'-O4'	-5.34	102.69	109.10
1	2	1057	U	C2-N1-C1'	5.34	124.11	117.70
36	1	1001	G	C6-C5-N7	-5.34	127.20	130.40
36	1	3128	G	C8-N9-C1'	-5.34	120.06	127.00
1	6	407	A	O5'-P-OP1	5.34	117.11	110.70
1	6	886	U	O4'-C1'-N1	5.34	112.47	108.20
36	5	2850	G	OP1-P-OP2	5.34	127.61	119.60
1	2	1171	A	N1-C6-N6	-5.34	115.40	118.60
1	2	1782	A	C5-C6-N6	5.34	127.97	123.70
36	1	286	U	N3-C2-O2	-5.34	118.46	122.20
36	1	2764	C	N3-C4-N4	5.34	121.73	118.00
36	1	2899	C	N3-C4-C5	-5.34	119.77	121.90
36	5	61	A	N1-C2-N3	5.34	131.97	129.30
36	5	1340	G	C8-N9-C4	5.34	108.53	106.40
36	5	1519	G	C4-C5-N7	5.34	112.94	110.80
36	1	1741	A	N1-C2-N3	5.33	131.97	129.30
36	1	2144	A	O4'-C1'-N9	5.33	112.47	108.20
38	8	15	G	N1-C6-O6	-5.33	116.70	119.90
36	1	658	G	OP2-P-O3'	5.33	116.93	105.20
36	1	1507	G	N1-C6-O6	5.33	123.10	119.90
1	6	407	A	C8-N9-C4	-5.33	103.67	105.80
36	5	2341	A	N7-C8-N9	-5.33	111.13	113.80
36	1	636	C	C5-C4-N4	-5.33	116.47	120.20
36	1	648	C	O5'-P-OP1	-5.33	100.90	105.70
36	1	804	C	C5-C4-N4	-5.33	116.47	120.20
36	1	1520	G	C5-N7-C8	5.33	106.97	104.30
36	1	2288	G	C8-N9-C1'	-5.33	120.07	127.00
36	1	2617	U	C4-C5-C6	5.33	122.90	119.70
36	5	973	A	C6-C5-N7	-5.33	128.57	132.30
36	5	1129	A	C5-N7-C8	-5.33	101.23	103.90
36	5	2406	C	N1-C2-O2	-5.33	115.70	118.90
36	5	2993	G	C5-C6-O6	-5.33	125.40	128.60
1	2	704	C	C6-N1-C2	-5.33	118.17	120.30
1	2	1031	U	OP2-P-O3'	5.33	116.93	105.20
36	5	218	G	O5'-P-OP2	-5.33	100.90	105.70
36	5	3029	A	N9-C4-C5	5.33	107.93	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	24	G	C8-N9-C1'	-5.33	120.07	127.00
36	1	2832	C	C5-C6-N1	-5.33	118.34	121.00
36	5	999	G	C8-N9-C4	5.33	108.53	106.40
36	5	1744	G	N3-C4-C5	-5.33	125.94	128.60
37	7	82	G	C4-C5-N7	5.33	112.93	110.80
1	6	542	A	O4'-C1'-N9	5.33	112.46	108.20
1	6	935	U	C6-N1-C2	-5.33	117.80	121.00
36	5	1184	A	C6-C5-N7	5.33	136.03	132.30
36	1	682	U	C6-N1-C1'	5.33	128.66	121.20
36	1	1400	G	N3-C4-N9	5.33	129.19	126.00
36	1	2646	C	N3-C4-C5	5.33	124.03	121.90
36	1	2811	A	N9-C4-C5	5.33	107.93	105.80
1	6	1199	G	N3-C4-C5	5.33	131.26	128.60
36	5	668	G	C5-N7-C8	5.33	106.96	104.30
36	5	869	G	N9-C4-C5	-5.33	103.27	105.40
36	5	2352	A	N1-C6-N6	5.33	121.80	118.60
36	5	2758	A	C4-C5-N7	-5.33	108.04	110.70
36	5	2994	A	N3-C4-C5	-5.33	123.07	126.80
1	6	29	U	N3-C4-C5	-5.32	111.41	114.60
1	6	55	A	N7-C8-N9	5.32	116.46	113.80
1	6	934	C	N3-C2-O2	-5.32	118.17	121.90
36	5	1127	G	N3-C2-N2	5.32	123.63	119.90
36	5	1405	U	O5'-P-OP2	-5.32	100.91	105.70
36	5	3109	G	N1-C6-O6	5.32	123.09	119.90
36	5	3295	A	C6-N1-C2	-5.32	115.41	118.60
1	6	1382	A	O4'-C1'-N9	5.32	112.46	108.20
1	2	305	C	C5-C6-N1	5.32	123.66	121.00
36	1	1331	U	O5'-P-OP2	-5.32	100.91	105.70
36	1	1341	U	N3-C4-C5	-5.32	111.41	114.60
36	1	1792	C	O4'-C1'-N1	5.32	112.46	108.20
36	1	2996	U	C5-C4-O4	-5.32	122.71	125.90
1	6	576	G	C6-C5-N7	-5.32	127.21	130.40
36	5	2407	C	N3-C4-C5	5.32	124.03	121.90
36	5	2813	A	N1-C6-N6	5.32	121.79	118.60
36	5	2827	U	C2-N1-C1'	5.32	124.08	117.70
1	2	1490	C	C6-N1-C2	-5.32	118.17	120.30
36	1	954	U	N1-C2-N3	5.32	118.09	114.90
36	5	2711	C	N3-C4-N4	5.32	121.72	118.00
1	2	274	G	C4-N9-C1'	5.32	133.41	126.50
1	2	394	C	C6-N1-C2	-5.32	118.17	120.30
36	1	720	A	C6-C5-N7	-5.32	128.58	132.30
36	1	2412	G	C8-N9-C4	-5.32	104.27	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2916	U	C2-N3-C4	5.32	130.19	127.00
37	3	82	G	N1-C2-N2	-5.32	111.42	116.20
1	6	536	C	C5-C6-N1	5.32	123.66	121.00
36	5	1116	G	O5'-P-OP1	-5.32	100.91	105.70
36	5	3173	G	OP1-P-OP2	5.32	127.58	119.60
1	2	75	U	C2-N1-C1'	5.32	124.08	117.70
36	1	878	G	N1-C2-N3	5.32	127.09	123.90
36	1	1074	U	N3-C4-C5	5.32	117.79	114.60
36	1	1520	G	N7-C8-N9	-5.32	110.44	113.10
1	6	1127	G	C6-C5-N7	-5.32	127.21	130.40
36	1	3224	G	C2-N3-C4	-5.31	109.24	111.90
37	7	93	C	N3-C2-O2	-5.31	118.18	121.90
36	1	509	U	O5'-P-OP1	5.31	117.08	110.70
36	1	635	G	C6-N1-C2	-5.31	121.91	125.10
36	1	2196	C	N3-C4-C5	5.31	124.03	121.90
1	6	934	C	N1-C2-O2	5.31	122.09	118.90
1	6	1119	G	N9-C4-C5	5.31	107.53	105.40
36	5	416	A	OP2-P-O3'	5.31	116.89	105.20
36	5	523	A	C5-C6-N6	5.31	127.95	123.70
36	5	834	U	N3-C4-C5	5.31	117.79	114.60
36	1	884	A	C5-C6-N6	-5.31	119.45	123.70
36	1	714	G	N3-C4-N9	5.31	129.19	126.00
36	1	1417	G	O4'-C1'-N9	-5.31	103.95	108.20
36	1	2245	C	N1-C2-O2	5.31	122.09	118.90
36	1	2924	U	C5-C6-N1	-5.31	120.05	122.70
36	1	2989	U	C6-N1-C2	-5.31	117.81	121.00
36	5	2864	A	C5-C6-N6	-5.31	119.45	123.70
36	5	3055	U	N3-C2-O2	5.31	125.92	122.20
36	1	2727	A	C2-N3-C4	5.31	113.25	110.60
54	M8	138	LEU	CA-CB-CG	5.31	127.51	115.30
1	6	65	A	N9-C4-C5	-5.31	103.68	105.80
1	6	314	C	N3-C4-C5	-5.31	119.78	121.90
1	6	1600	A	OP1-P-O3'	5.31	116.88	105.20
36	5	1354	G	N1-C6-O6	5.31	123.08	119.90
36	5	1413	G	C8-N9-C4	5.31	108.52	106.40
36	5	1770	G	C4-N9-C1'	5.31	133.40	126.50
36	5	2299	A	O5'-P-OP2	-5.31	100.92	105.70
36	5	2670	G	O5'-P-OP2	-5.31	100.92	105.70
36	5	2856	G	C5-C6-O6	-5.31	125.42	128.60
36	5	3204	C	C5-C6-N1	-5.31	118.35	121.00
36	1	2324	A	N9-C4-C5	5.31	107.92	105.80
36	1	2899	C	OP2-P-O3'	5.31	116.87	105.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	6	541	A	C8-N9-C4	-5.31	103.68	105.80
36	1	224	C	N1-C2-O2	-5.30	115.72	118.90
36	1	1581	C	N1-C2-O2	5.30	122.08	118.90
36	1	3270	U	C2-N1-C1'	-5.30	111.33	117.70
1	6	20	G	N3-C4-C5	5.30	131.25	128.60
36	5	877	C	N3-C4-N4	-5.30	114.29	118.00
36	5	953	G	C8-N9-C4	5.30	108.52	106.40
36	5	3153	U	C2-N1-C1'	5.30	124.06	117.70
1	6	1112	G	N3-C2-N2	5.30	123.61	119.90
36	5	2283	G	C5-N7-C8	-5.30	101.65	104.30
1	2	1488	G	N7-C8-N9	5.30	115.75	113.10
36	1	2953	U	N3-C4-O4	5.30	123.11	119.40
1	6	447	U	C6-N1-C2	-5.30	117.82	121.00
36	5	421	G	C8-N9-C4	-5.30	104.28	106.40
36	5	629	U	N1-C2-N3	5.30	118.08	114.90
36	5	873	C	N1-C2-O2	5.30	122.08	118.90
36	5	2607	G	C6-C5-N7	-5.30	127.22	130.40
36	1	3011	A	C8-N9-C4	5.30	107.92	105.80
36	5	1298	C	C6-N1-C2	-5.30	118.18	120.30
36	5	2123	G	C5-N7-C8	5.30	106.95	104.30
36	5	2966	G	C4-N9-C1'	5.30	133.39	126.50
36	1	679	U	N1-C2-O2	5.30	126.51	122.80
36	1	2358	A	C2-N3-C4	-5.30	107.95	110.60
36	5	2617	U	N3-C4-O4	5.30	123.11	119.40
38	8	104	A	C8-N9-C4	5.30	107.92	105.80
36	1	642	U	N1-C2-N3	5.30	118.08	114.90
36	1	2356	A	C5-N7-C8	-5.30	101.25	103.90
36	1	2968	G	C4-C5-N7	5.30	112.92	110.80
1	6	1573	A	P-O3'-C3'	5.30	126.06	119.70
36	5	27	C	N1-C2-O2	-5.30	115.72	118.90
36	5	1528	G	C4-N9-C1'	5.30	133.38	126.50
36	5	2959	C	OP2-P-O3'	5.30	116.85	105.20
38	8	96	A	N1-C6-N6	5.30	121.78	118.60
36	1	2850	G	C6-C5-N7	-5.29	127.22	130.40
38	4	16	G	N7-C8-N9	-5.29	110.45	113.10
38	4	20	U	N1-C2-N3	5.29	118.08	114.90
36	5	2817	A	N3-C4-C5	-5.29	123.09	126.80
36	1	1383	G	C8-N9-C4	-5.29	104.28	106.40
36	5	645	A	C5-C6-N1	5.29	120.35	117.70
36	5	1116	G	C8-N9-C1'	-5.29	120.12	127.00
36	5	1476	G	N1-C2-N2	-5.29	111.44	116.20
36	5	2744	U	N3-C2-O2	-5.29	118.50	122.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	2878	G	C2-N3-C4	5.29	114.55	111.90
36	5	3107	U	C2-N1-C1'	5.29	124.05	117.70
36	5	3301	U	C6-N1-C2	5.29	124.18	121.00
36	1	1005	G	N7-C8-N9	-5.29	110.45	113.10
36	1	2874	G	C4-C5-C6	5.29	121.97	118.80
37	3	9	C	N3-C4-C5	-5.29	119.78	121.90
1	6	272	U	C5-C6-N1	5.29	125.35	122.70
36	5	1369	A	C4-C5-C6	5.29	119.65	117.00
1	2	331	A	N9-C4-C5	5.29	107.92	105.80
36	1	1389	G	N1-C6-O6	5.29	123.07	119.90
36	5	2900	A	N1-C6-N6	-5.29	115.43	118.60
36	5	2930	A	C8-N9-C4	5.29	107.92	105.80
38	8	17	A	C6-C5-N7	-5.29	128.60	132.30
1	2	950	C	C6-N1-C2	-5.29	118.18	120.30
1	2	1168	U	OP1-P-O3'	5.29	116.83	105.20
36	1	59	G	C5-C6-O6	-5.29	125.43	128.60
36	1	2927	C	N1-C2-O2	-5.29	115.73	118.90
36	5	84	U	N3-C2-O2	-5.29	118.50	122.20
36	5	3261	C	C6-N1-C2	-5.29	118.19	120.30
67	o1	97	LEU	CA-CB-CG	5.29	127.47	115.30
36	1	2611	U	N3-C4-C5	-5.29	111.43	114.60
36	1	2787	G	N1-C6-O6	5.29	123.07	119.90
36	1	2875	U	C6-N1-C2	-5.29	117.83	121.00
36	5	2754	G	C8-N9-C4	5.29	108.52	106.40
1	2	1401	A	N1-C6-N6	-5.29	115.43	118.60
36	1	687	U	OP2-P-O3'	5.29	116.83	105.20
36	1	906	A	C6-N1-C2	-5.29	115.43	118.60
36	1	970	A	C4-C5-N7	5.29	113.34	110.70
36	1	1607	U	N3-C2-O2	-5.29	118.50	122.20
36	1	2247	G	C5-C6-N1	-5.29	108.86	111.50
36	1	2400	G	N3-C4-C5	5.29	131.24	128.60
36	1	2779	A	C8-N9-C4	5.29	107.91	105.80
36	5	1605	A	O4'-C1'-N9	5.29	112.43	108.20
36	1	336	A	N9-C4-C5	-5.28	103.69	105.80
36	1	410	U	N1-C2-N3	5.28	118.07	114.90
36	1	1402	C	N3-C4-C5	5.28	124.01	121.90
52	M6	133	ARG	NE-CZ-NH2	-5.28	117.66	120.30
1	6	405	C	C6-N1-C2	5.28	122.41	120.30
36	5	804	C	C6-N1-C2	5.28	122.41	120.30
36	5	1896	A	C5-N7-C8	-5.28	101.26	103.90
36	1	2122	G	O5'-P-OP2	-5.28	100.95	105.70
36	1	2169	G	N1-C6-O6	-5.28	116.73	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2403	G	C6-C5-N7	-5.28	127.23	130.40
36	1	2731	U	OP2-P-O3'	5.28	116.82	105.20
36	1	2812	C	C5-C6-N1	-5.28	118.36	121.00
36	5	437	G	N3-C2-N2	-5.28	116.20	119.90
36	5	2679	A	N7-C8-N9	-5.28	111.16	113.80
37	7	105	C	C4-C5-C6	5.28	120.04	117.40
36	1	53	G	C8-N9-C1'	-5.28	120.14	127.00
36	1	343	U	N1-C2-N3	5.28	118.07	114.90
36	1	2323	G	C5-C6-N1	5.28	114.14	111.50
36	5	1081	U	C5-C6-N1	5.28	125.34	122.70
37	7	1	G	C6-C5-N7	-5.28	127.23	130.40
37	7	73	C	C2-N1-C1'	5.28	124.61	118.80
36	1	1393	A	N1-C2-N3	5.28	131.94	129.30
37	3	38	U	C5-C6-N1	5.28	125.34	122.70
38	4	14	C	C5-C6-N1	-5.28	118.36	121.00
36	5	63	A	N1-C6-N6	5.28	121.77	118.60
37	7	49	G	N3-C4-C5	5.28	131.24	128.60
36	1	1904	C	C5-C6-N1	5.28	123.64	121.00
36	1	2132	C	O5'-P-OP2	-5.28	100.95	105.70
36	5	2398	A	C4-C5-N7	-5.28	108.06	110.70
37	7	14	U	C6-N1-C2	5.28	124.17	121.00
36	1	2586	G	N1-C6-O6	-5.27	116.74	119.90
36	5	1181	U	N1-C2-N3	5.27	118.06	114.90
36	5	1501	U	C5-C4-O4	5.27	129.06	125.90
36	5	2889	C	OP1-P-OP2	-5.27	111.69	119.60
36	5	2953	U	C5-C4-O4	-5.27	122.74	125.90
1	2	1137	A	N3-C4-C5	5.27	130.49	126.80
36	1	885	U	OP1-P-O3'	5.27	116.80	105.20
36	1	979	U	O4'-C1'-N1	5.27	112.42	108.20
36	1	1192	C	C5-C6-N1	5.27	123.64	121.00
36	1	2537	U	P-O3'-C3'	5.27	126.03	119.70
36	5	31	C	OP2-P-O3'	5.27	116.80	105.20
36	5	3164	C	O4'-C1'-N1	5.27	112.42	108.20
52	m6	133	ARG	NE-CZ-NH2	-5.27	117.66	120.30
36	1	721	G	C8-N9-C4	-5.27	104.29	106.40
36	1	1119	C	C5-C4-N4	5.27	123.89	120.20
1	6	85	A	N9-C4-C5	5.27	107.91	105.80
36	1	3000	A	N3-C4-C5	5.27	130.49	126.80
1	6	390	G	C4-C5-N7	5.27	112.91	110.80
36	5	2142	A	C6-N1-C2	-5.27	115.44	118.60
36	5	3035	A	N9-C4-C5	-5.27	103.69	105.80
36	5	3067	C	N3-C2-O2	-5.27	118.21	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	2	863	A	N3-C4-C5	5.27	130.49	126.80
36	1	1306	G	N3-C2-N2	-5.27	116.21	119.90
36	1	1604	G	C8-N9-C1'	-5.27	120.15	127.00
36	5	776	U	N3-C4-O4	-5.27	115.71	119.40
36	1	921	A	C5-C6-N6	5.27	127.91	123.70
36	1	1217	A	OP2-P-O3'	5.27	116.79	105.20
36	1	1422	G	N1-C6-O6	5.27	123.06	119.90
37	3	120	C	N3-C2-O2	-5.27	118.21	121.90
17	c5	36	LEU	CA-CB-CG	5.27	127.41	115.30
36	5	2775	U	C5-C4-O4	5.27	129.06	125.90
36	5	3092	C	C2-N3-C4	-5.27	117.27	119.90
36	1	2602	G	C4-C5-N7	-5.26	108.69	110.80
36	1	2630	C	O5'-P-OP1	-5.26	100.96	105.70
1	6	1130	G	C6-C5-N7	5.26	133.56	130.40
36	5	317	A	N1-C6-N6	-5.26	115.44	118.60
36	5	934	G	C2-N3-C4	5.26	114.53	111.90
36	5	1599	G	N9-C4-C5	-5.26	103.29	105.40
36	5	2524	A	C5-N7-C8	-5.26	101.27	103.90
36	1	594	U	C5-C4-O4	5.26	129.06	125.90
36	1	2643	A	O5'-P-OP1	-5.26	100.97	105.70
38	4	25	G	N9-C4-C5	5.26	107.50	105.40
36	5	644	G	N9-C4-C5	5.26	107.50	105.40
36	5	1332	A	C6-C5-N7	-5.26	128.62	132.30
36	5	2855	U	N1-C2-O2	-5.26	119.12	122.80
36	5	3389	U	C5-C4-O4	5.26	129.06	125.90
1	2	25	C	C2-N1-C1'	5.26	124.59	118.80
1	2	1274	C	C2-N1-C1'	5.26	124.58	118.80
22	D0	34	LEU	CA-CB-CG	5.26	127.40	115.30
36	1	2400	G	C6-C5-N7	-5.26	127.24	130.40
36	5	1839	A	O5'-P-OP1	-5.26	100.97	105.70
1	2	829	A	P-O3'-C3'	5.26	126.01	119.70
36	1	1748	G	C6-C5-N7	-5.26	127.25	130.40
36	1	2136	C	C5-C4-N4	5.26	123.88	120.20
36	1	2365	C	C5-C6-N1	-5.26	118.37	121.00
1	6	1521	G	N1-C6-O6	-5.26	116.75	119.90
36	5	508	U	C5-C6-N1	5.26	125.33	122.70
36	5	1075	A	N7-C8-N9	-5.26	111.17	113.80
36	5	1513	G	N3-C4-N9	5.26	129.15	126.00
36	5	1514	G	OP1-P-O3'	5.26	116.77	105.20
36	1	871	U	OP2-P-O3'	5.25	116.76	105.20
1	6	192	U	P-O3'-C3'	5.25	126.01	119.70
1	6	1550	A	N7-C8-N9	5.25	116.43	113.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	1169	A	O5'-P-OP2	-5.25	100.97	105.70
36	5	2826	U	C5-C6-N1	-5.25	120.07	122.70
36	5	2892	A	C6-C5-N7	-5.25	128.62	132.30
1	2	1297	G	C5-C6-O6	-5.25	125.45	128.60
36	1	1807	G	C8-N9-C4	-5.25	104.30	106.40
36	1	2633	U	O5'-P-OP2	5.25	117.00	110.70
36	1	2889	C	N3-C4-C5	5.25	124.00	121.90
1	6	767	U	N3-C2-O2	-5.25	118.52	122.20
36	5	639	G	O5'-P-OP1	-5.25	100.97	105.70
36	1	396	A	C8-N9-C4	-5.25	103.70	105.80
36	1	1429	G	C5-N7-C8	5.25	106.93	104.30
36	1	2942	C	C6-N1-C1'	5.25	127.10	120.80
1	6	455	C	C6-N1-C2	-5.25	118.20	120.30
1	6	1773	C	C5-C6-N1	5.25	123.63	121.00
36	1	2626	A	N1-C6-N6	-5.25	115.45	118.60
38	4	26	U	C2-N1-C1'	5.25	124.00	117.70
36	5	699	A	C2-N3-C4	-5.25	107.97	110.60
36	5	1054	A	OP2-P-O3'	5.25	116.75	105.20
38	8	97	A	C8-N9-C4	5.25	107.90	105.80
1	2	163	G	O4'-C1'-N9	5.25	112.40	108.20
1	2	1636	C	N3-C4-N4	5.25	121.67	118.00
36	1	1413	G	N3-C2-N2	-5.25	116.22	119.90
36	1	1495	U	C2-N1-C1'	-5.25	111.40	117.70
1	6	1600	A	C2-N3-C4	-5.25	107.98	110.60
36	5	1856	C	C6-N1-C2	-5.25	118.20	120.30
1	2	278	U	N1-C2-O2	5.25	126.47	122.80
37	3	92	A	C5-C6-N1	-5.25	115.08	117.70
37	3	93	C	N1-C2-O2	5.25	122.05	118.90
36	5	869	G	C4-C5-N7	5.25	112.90	110.80
36	5	999	G	C5-C6-O6	-5.25	125.45	128.60
36	5	1116	G	N1-C2-N2	-5.25	111.48	116.20
36	5	1190	A	N7-C8-N9	5.25	116.42	113.80
1	6	687	G	N3-C4-N9	-5.25	122.85	126.00
36	5	520	U	C2-N1-C1'	-5.25	111.41	117.70
36	5	842	G	C5-C6-O6	-5.25	125.45	128.60
1	2	734	A	P-O3'-C3'	5.24	125.99	119.70
36	1	2370	G	C2-N3-C4	-5.24	109.28	111.90
1	6	620	A	C4-C5-C6	5.24	119.62	117.00
36	5	1443	G	C6-C5-N7	-5.24	127.25	130.40
36	1	2247	G	C4-C5-C6	5.24	121.94	118.80
36	5	2950	G	N1-C6-O6	5.24	123.05	119.90
36	1	670	C	N3-C2-O2	-5.24	118.23	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	908	G	C6-C5-N7	-5.24	127.26	130.40
36	5	886	C	C6-N1-C2	5.24	122.40	120.30
38	8	34	U	C5-C6-N1	-5.24	120.08	122.70
36	1	688	G	N3-C4-N9	5.24	129.14	126.00
36	1	1444	G	N1-C6-O6	5.24	123.04	119.90
1	6	391	A	C8-N9-C4	5.24	107.89	105.80
1	6	779	U	N1-C1'-C2'	5.24	120.81	114.00
36	5	1716	U	P-O3'-C3'	5.24	125.99	119.70
37	7	49	G	N3-C2-N2	-5.24	116.23	119.90
36	5	801	A	OP1-P-O3'	5.24	116.72	105.20
36	5	1348	U	C6-N1-C2	-5.24	117.86	121.00
36	5	2329	C	C6-N1-C2	5.24	122.39	120.30
36	5	2777	G	C5-C6-N1	-5.24	108.88	111.50
36	1	557	A	C8-N9-C4	5.24	107.89	105.80
36	1	870	G	N3-C4-N9	-5.24	122.86	126.00
36	1	2746	A	C2-N3-C4	-5.24	107.98	110.60
36	1	2983	C	O4'-C1'-N1	5.24	112.39	108.20
1	6	758	U	N3-C2-O2	-5.24	118.54	122.20
1	6	1608	U	C6-N1-C2	-5.24	117.86	121.00
36	5	1481	A	N1-C2-N3	5.24	131.92	129.30
36	5	1846	C	N3-C2-O2	-5.24	118.23	121.90
38	8	15	G	C5-C6-O6	5.24	131.74	128.60
36	1	3195	U	P-O3'-C3'	5.23	125.98	119.70
36	1	3375	A	OP1-P-O3'	5.23	116.72	105.20
36	5	2736	A	C4-C5-C6	5.23	119.62	117.00
36	1	1152	G	N3-C4-C5	-5.23	125.98	128.60
36	5	1865	A	C2-N3-C4	-5.23	107.98	110.60
1	2	501	U	P-O3'-C3'	5.23	125.98	119.70
1	2	1258	U	C2-N1-C1'	5.23	123.98	117.70
1	2	1462	G	C4-C5-N7	5.23	112.89	110.80
36	1	27	C	OP1-P-OP2	5.23	127.45	119.60
36	1	692	A	C5-C6-N1	-5.23	115.08	117.70
36	1	1327	C	N1-C2-O2	-5.23	115.76	118.90
36	1	2308	C	N3-C4-C5	5.23	123.99	121.90
36	1	2726	C	N1-C2-O2	5.23	122.04	118.90
1	6	1481	C	C6-N1-C2	-5.23	118.21	120.30
1	6	1697	G	N3-C4-C5	-5.23	125.98	128.60
36	5	197	G	N3-C2-N2	5.23	123.56	119.90
36	5	295	A	O5'-P-OP1	-5.23	100.99	105.70
36	5	1213	G	C2-N3-C4	5.23	114.52	111.90
36	5	1899	G	C5-C6-O6	5.23	131.74	128.60
36	1	2624	G	N1-C6-O6	5.23	123.04	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	3182	G	N3-C4-N9	5.23	129.14	126.00
38	4	95	G	C4-N9-C1'	-5.23	119.70	126.50
1	6	543	C	C5-C4-N4	5.23	123.86	120.20
36	5	1177	G	N1-C2-N3	5.23	127.04	123.90
36	5	1186	G	C8-N9-C4	-5.23	104.31	106.40
36	5	1374	G	C8-N9-C4	5.23	108.49	106.40
36	1	2752	U	N1-C2-N3	5.23	118.04	114.90
36	1	2762	A	N1-C6-N6	-5.23	115.46	118.60
1	6	360	A	C5-C6-N6	-5.23	119.52	123.70
1	6	405	C	C5-C6-N1	-5.23	118.39	121.00
36	5	1200	A	N3-C4-N9	5.23	131.58	127.40
36	5	1440	G	N1-C2-N3	5.23	127.04	123.90
36	5	2278	C	N3-C4-C5	5.23	123.99	121.90
36	5	2945	G	N3-C2-N2	5.23	123.56	119.90
36	5	3104	U	O5'-P-OP2	-5.23	101.00	105.70
36	1	594	U	N3-C4-C5	-5.23	111.46	114.60
36	1	1504	A	N1-C6-N6	-5.23	115.46	118.60
36	5	807	A	C8-N9-C4	-5.23	103.71	105.80
36	5	1159	A	C5-N7-C8	-5.23	101.29	103.90
36	1	2946	A	C4-C5-C6	5.22	119.61	117.00
36	1	3100	U	C6-N1-C2	5.22	124.14	121.00
36	1	3302	U	C2-N3-C4	-5.22	123.86	127.00
36	1	3319	U	C2-N1-C1'	5.22	123.97	117.70
36	5	1462	A	N3-C4-C5	5.22	130.46	126.80
1	2	795	U	C4-C5-C6	5.22	122.83	119.70
36	1	1381	A	C4-C5-C6	5.22	119.61	117.00
36	1	1939	G	C8-N9-C1'	-5.22	120.21	127.00
1	6	1715	G	C4-N9-C1'	5.22	133.29	126.50
36	5	1445	U	C2-N3-C4	-5.22	123.87	127.00
1	2	426	G	N3-C4-N9	5.22	129.13	126.00
36	1	1371	G	C2-N3-C4	-5.22	109.29	111.90
36	1	2422	C	C2-N3-C4	-5.22	117.29	119.90
36	5	190	U	N1-C2-O2	5.22	126.45	122.80
36	5	1183	C	C5-C6-N1	-5.22	118.39	121.00
36	5	1891	A	N1-C6-N6	5.22	121.73	118.60
36	5	3017	A	C2-N3-C4	-5.22	107.99	110.60
37	7	101	G	C5-N7-C8	-5.22	101.69	104.30
36	1	1152	G	N3-C4-N9	5.22	129.13	126.00
36	1	1603	A	C8-N9-C4	-5.22	103.71	105.80
36	1	2143	A	C4-C5-N7	5.22	113.31	110.70
36	1	2808	A	N1-C6-N6	5.22	121.73	118.60
38	4	93	U	C2-N1-C1'	-5.22	111.44	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	6	1798	U	C5-C6-N1	5.22	125.31	122.70
36	5	34	A	OP2-P-O3'	5.22	116.69	105.20
36	5	3301	U	C5-C6-N1	-5.22	120.09	122.70
36	1	2423	U	C5-C6-N1	5.22	125.31	122.70
36	5	1477	A	C6-N1-C2	-5.22	115.47	118.60
36	5	1917	C	C2-N1-C1'	-5.22	113.06	118.80
36	5	2870	C	C2-N1-C1'	-5.22	113.06	118.80
1	2	970	A	C4-C5-N7	5.22	113.31	110.70
1	2	1614	A	N1-C6-N6	5.22	121.73	118.60
36	1	1882	G	C5-C6-O6	-5.22	125.47	128.60
36	1	2752	U	C5-C4-O4	5.22	129.03	125.90
38	4	101	U	C6-N1-C2	-5.22	117.87	121.00
1	6	438	A	O5'-P-OP1	-5.22	101.00	105.70
1	6	1458	G	C4-N9-C1'	5.22	133.28	126.50
36	5	1352	A	OP1-P-O3'	5.22	116.67	105.20
36	5	1359	C	C6-N1-C2	5.22	122.39	120.30
36	5	2852	C	C6-N1-C2	5.22	122.39	120.30
36	1	1097	G	P-O3'-C3'	5.21	125.96	119.70
36	1	2138	A	C8-N9-C4	-5.21	103.71	105.80
36	1	2386	A	C8-N9-C4	-5.21	103.72	105.80
1	6	576	G	C5-C6-O6	-5.21	125.47	128.60
36	5	1368	U	N3-C4-O4	5.21	123.05	119.40
36	5	1404	G	N9-C4-C5	-5.21	103.31	105.40
1	2	278	U	N3-C2-O2	-5.21	118.55	122.20
36	1	1498	A	C6-N1-C2	-5.21	115.47	118.60
36	5	2832	C	C5-C6-N1	-5.21	118.39	121.00
1	2	559	C	C5-C6-N1	5.21	123.61	121.00
36	1	1296	C	N1-C2-O2	-5.21	115.77	118.90
1	6	1537	C	N1-C2-O2	-5.21	115.77	118.90
36	5	400	G	C6-C5-N7	5.21	133.53	130.40
36	5	1900	A	N9-C4-C5	-5.21	103.72	105.80
36	5	2801	A	O4'-C1'-N9	5.21	112.37	108.20
1	2	553	G	C5-N7-C8	-5.21	101.70	104.30
1	2	1555	A	C8-N9-C4	-5.21	103.72	105.80
36	1	222	A	N9-C4-C5	-5.21	103.72	105.80
36	1	2913	C	N3-C2-O2	-5.21	118.25	121.90
36	5	2836	C	N3-C4-C5	-5.21	119.82	121.90
1	2	89	G	O5'-P-OP1	-5.21	101.01	105.70
1	2	1082	C	C5-C6-N1	5.21	123.60	121.00
36	1	2612	U	C2-N1-C1'	-5.21	111.45	117.70
36	1	3118	C	C6-N1-C2	-5.21	118.22	120.30
36	1	3242	G	C6-C5-N7	5.21	133.52	130.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	6	311	U	N3-C2-O2	-5.21	118.56	122.20
36	5	718	G	C4-C5-C6	5.21	121.92	118.80
36	5	1367	G	C8-N9-C1'	-5.21	120.23	127.00
36	5	2167	A	N3-C4-C5	-5.21	123.16	126.80
36	5	2649	A	C2-N3-C4	5.21	113.20	110.60
36	5	2805	G	C6-C5-N7	-5.21	127.28	130.40
36	5	3028	G	N3-C4-N9	5.21	129.12	126.00
36	1	1149	G	C4-C5-N7	-5.21	108.72	110.80
1	6	44	U	N3-C2-O2	5.21	125.84	122.20
1	6	1120	U	N3-C2-O2	-5.21	118.56	122.20
36	5	1480	G	O4'-C1'-N9	5.21	112.36	108.20
1	2	1275	A	C4-C5-C6	5.20	119.60	117.00
36	1	1394	A	N3-C4-C5	5.20	130.44	126.80
36	1	2387	A	C8-N9-C4	5.20	107.88	105.80
36	1	2808	A	C5-N7-C8	-5.20	101.30	103.90
1	6	610	G	N3-C4-N9	5.20	129.12	126.00
1	6	1437	U	C5-C6-N1	-5.20	120.10	122.70
36	5	1321	G	C2-N3-C4	-5.20	109.30	111.90
36	1	1501	U	C6-N1-C2	5.20	124.12	121.00
36	5	757	C	C5-C6-N1	-5.20	118.40	121.00
36	5	1881	A	C6-N1-C2	-5.20	115.48	118.60
36	5	2761	G	C6-N1-C2	-5.20	121.98	125.10
38	8	114	G	N1-C6-O6	5.20	123.02	119.90
1	2	822	U	C5-C6-N1	5.20	125.30	122.70
36	1	1724	U	OP1-P-O3'	5.20	116.64	105.20
36	1	3093	C	C2-N1-C1'	-5.20	113.08	118.80
1	6	913	G	C5-C6-O6	-5.20	125.48	128.60
36	5	651	G	C4-N9-C1'	5.20	133.26	126.50
36	5	1377	G	C5-N7-C8	5.20	106.90	104.30
36	5	2601	A	C8-N9-C4	5.20	107.88	105.80
36	5	2757	U	N3-C4-O4	5.20	123.04	119.40
36	5	2759	U	C6-N1-C2	-5.20	117.88	121.00
1	2	1473	U	N1-C2-O2	5.20	126.44	122.80
1	2	1600	A	P-O3'-C3'	5.20	125.94	119.70
36	1	1411	C	OP1-P-O3'	5.20	116.64	105.20
36	1	2101	C	P-O3'-C3'	5.20	125.94	119.70
36	5	1913	A	N1-C6-N6	5.20	121.72	118.60
36	5	3099	C	N1-C2-O2	-5.20	115.78	118.90
36	1	2801	A	OP1-P-O3'	5.20	116.63	105.20
1	2	1616	G	C6-C5-N7	-5.20	127.28	130.40
36	1	940	G	C6-C5-N7	5.20	133.52	130.40
36	1	1178	G	N3-C4-N9	5.20	129.12	126.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	1888	U	C2-N3-C4	-5.20	123.88	127.00
36	1	1912	U	C5-C4-O4	5.20	129.02	125.90
36	5	1185	C	N3-C4-N4	5.20	121.64	118.00
36	5	1878	G	C8-N9-C1'	-5.20	120.25	127.00
36	5	2705	A	N9-C4-C5	-5.20	103.72	105.80
36	5	2816	G	N9-C4-C5	-5.20	103.32	105.40
36	5	2924	U	C6-N1-C2	5.20	124.12	121.00
1	2	974	A	C8-N9-C4	5.19	107.88	105.80
1	2	1639	C	N3-C4-N4	5.19	121.64	118.00
36	1	993	G	N3-C4-N9	5.19	129.12	126.00
36	1	2812	C	OP2-P-O3'	5.19	116.63	105.20
37	3	92	A	C2-N3-C4	-5.19	108.00	110.60
36	5	50	U	N3-C2-O2	-5.19	118.56	122.20
36	5	2361	A	OP2-P-O3'	5.19	116.63	105.20
36	1	806	A	C8-N9-C4	5.19	107.88	105.80
36	1	1851	G	N1-C6-O6	5.19	123.02	119.90
1	6	1150	G	C4-N9-C1'	-5.19	119.75	126.50
36	5	1203	A	N1-C6-N6	5.19	121.72	118.60
36	1	70	A	O5'-P-OP2	5.19	116.93	110.70
36	1	1847	A	N1-C2-N3	5.19	131.90	129.30
36	1	1853	U	N3-C4-O4	-5.19	115.77	119.40
1	6	139	C	N3-C4-C5	5.19	123.98	121.90
1	6	1629	G	OP2-P-O3'	5.19	116.62	105.20
36	5	110	G	N7-C8-N9	-5.19	110.50	113.10
36	5	1450	G	C5-N7-C8	-5.19	101.70	104.30
36	5	2728	G	O4'-C1'-N9	5.19	112.35	108.20
36	1	2434	U	C5-C6-N1	-5.19	120.11	122.70
36	5	209	A	N1-C2-N3	5.19	131.90	129.30
36	5	523	A	C4-C5-N7	-5.19	108.11	110.70
36	5	2761	G	C8-N9-C1'	5.19	133.75	127.00
36	5	2821	C	N1-C2-O2	5.19	122.01	118.90
36	5	3192	U	C2-N1-C1'	-5.19	111.47	117.70
37	7	5	G	N3-C4-C5	5.19	131.19	128.60
1	2	56	U	N1-C2-O2	5.19	126.43	122.80
36	1	196	G	O5'-P-OP1	-5.19	101.03	105.70
36	1	645	A	N1-C2-N3	5.19	131.89	129.30
36	1	652	G	N1-C2-N2	-5.19	111.53	116.20
36	1	3005	A	OP1-P-OP2	5.19	127.38	119.60
36	5	2274	U	C5-C6-N1	5.19	125.29	122.70
36	5	2697	A	N1-C6-N6	5.19	121.71	118.60
36	5	2980	U	N1-C2-N3	5.19	118.01	114.90
36	5	3283	U	O4'-C1'-N1	5.19	112.35	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	2	1184	A	N1-C6-N6	5.19	121.71	118.60
36	1	965	A	N7-C8-N9	5.19	116.39	113.80
36	1	1449	A	N1-C6-N6	-5.19	115.49	118.60
41	14	339	LEU	CA-CB-CG	5.19	127.23	115.30
36	1	890	C	N3-C2-O2	-5.18	118.27	121.90
36	1	1001	G	C4-C5-N7	5.18	112.87	110.80
36	1	2201	G	C6-C5-N7	-5.18	127.29	130.40
36	1	2679	A	O4'-C1'-N9	5.18	112.35	108.20
38	4	40	A	N3-C4-N9	5.18	131.55	127.40
38	4	58	G	C6-C5-N7	-5.18	127.29	130.40
1	6	297	U	N1-C2-O2	5.18	126.43	122.80
1	6	1569	A	C4-N9-C1'	5.18	135.63	126.30
36	5	1011	A	C8-N9-C4	-5.18	103.73	105.80
36	5	1114	U	O5'-P-OP2	-5.18	101.03	105.70
36	5	1137	C	N1-C2-O2	5.18	122.01	118.90
36	1	3011	A	N7-C8-N9	-5.18	111.21	113.80
36	1	3375	A	N1-C6-N6	-5.18	115.49	118.60
36	5	878	G	C8-N9-C4	-5.18	104.33	106.40
36	5	2353	G	C6-C5-N7	-5.18	127.29	130.40
36	5	2639	G	C8-N9-C4	-5.18	104.33	106.40
37	7	67	G	N1-C6-O6	5.18	123.01	119.90
1	2	768	C	N3-C4-C5	-5.18	119.83	121.90
1	6	17	C	N3-C2-O2	-5.18	118.27	121.90
36	5	436	A	N7-C8-N9	5.18	116.39	113.80
36	5	1190	A	O4'-C1'-N9	-5.18	104.06	108.20
38	8	25	G	OP2-P-O3'	-5.18	93.80	105.20
1	2	1600	A	OP1-P-O3'	5.18	116.59	105.20
36	1	928	C	N3-C2-O2	-5.18	118.27	121.90
36	1	1748	G	N1-C6-O6	5.18	123.01	119.90
36	1	3178	A	C8-N9-C4	5.18	107.87	105.80
36	1	3203	U	OP2-P-O3'	5.18	116.60	105.20
36	5	1417	G	C6-C5-N7	-5.18	127.29	130.40
36	5	3063	C	N1-C2-O2	-5.18	115.79	118.90
1	2	1745	G	C5-C6-O6	-5.18	125.49	128.60
36	5	842	G	C6-C5-N7	-5.18	127.29	130.40
36	5	2344	U	C5-C6-N1	-5.18	120.11	122.70
36	5	2709	C	C2-N3-C4	-5.18	117.31	119.90
36	5	3326	G	N1-C6-O6	5.18	123.01	119.90
1	2	543	C	N3-C2-O2	-5.18	118.28	121.90
36	1	1902	G	C4-C5-N7	5.18	112.87	110.80
36	1	2896	A	C2-N3-C4	-5.18	108.01	110.60
36	1	2984	C	C5-C4-N4	5.18	123.82	120.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	6	39	A	N9-C4-C5	5.18	107.87	105.80
1	6	384	G	N3-C4-C5	5.18	131.19	128.60
36	5	413	U	C5-C4-O4	-5.18	122.80	125.90
36	5	645	A	C8-N9-C4	-5.18	103.73	105.80
36	5	2281	A	C4-C5-N7	5.18	113.29	110.70
1	2	1596	C	N3-C2-O2	-5.17	118.28	121.90
36	1	1008	U	C6-N1-C2	5.17	124.11	121.00
36	1	1367	G	C8-N9-C1'	-5.17	120.27	127.00
1	6	467	G	N9-C4-C5	-5.17	103.33	105.40
36	5	1392	G	N7-C8-N9	-5.17	110.51	113.10
36	5	1903	U	N3-C4-C5	-5.17	111.50	114.60
36	5	2334	U	N3-C2-O2	-5.17	118.58	122.20
36	5	2878	G	OP1-P-OP2	-5.17	111.84	119.60
36	5	3009	G	C8-N9-C4	-5.17	104.33	106.40
36	5	3197	G	N3-C4-N9	-5.17	122.89	126.00
36	1	426	G	C8-N9-C4	-5.17	104.33	106.40
37	3	33	U	C2-N1-C1'	5.17	123.91	117.70
1	2	1615	C	P-O3'-C3'	5.17	125.91	119.70
36	1	222	A	O5'-P-OP2	-5.17	101.05	105.70
36	1	650	C	C6-N1-C2	5.17	122.37	120.30
36	1	1908	A	C6-C5-N7	-5.17	128.68	132.30
36	1	3144	G	N1-C6-O6	5.17	123.00	119.90
36	5	661	G	C4-N9-C1'	5.17	133.22	126.50
36	5	680	G	C4-N9-C1'	-5.17	119.78	126.50
36	5	934	G	N3-C4-C5	-5.17	126.01	128.60
36	5	1049	C	N3-C4-C5	-5.17	119.83	121.90
36	5	1107	C	C5-C6-N1	-5.17	118.41	121.00
36	5	1184	A	C8-N9-C4	5.17	107.87	105.80
36	5	1783	U	C5-C4-O4	5.17	129.00	125.90
36	5	3183	A	N1-C6-N6	-5.17	115.50	118.60
36	1	285	A	C2-N3-C4	-5.17	108.02	110.60
36	1	1113	G	C4-C5-N7	5.17	112.87	110.80
36	1	1379	G	C2-N3-C4	-5.17	109.31	111.90
1	2	576	G	N3-C4-C5	5.17	131.18	128.60
36	1	233	C	C5-C6-N1	-5.17	118.42	121.00
36	1	329	U	C2-N1-C1'	5.17	123.90	117.70
36	1	917	A	N9-C4-C5	5.17	107.87	105.80
36	1	1381	A	C6-C5-N7	-5.17	128.68	132.30
36	1	1394	A	C8-N9-C4	5.17	107.87	105.80
36	5	637	C	C4-C5-C6	-5.17	114.82	117.40
36	5	1159	A	N3-C4-N9	-5.17	123.27	127.40
36	1	170	G	O5'-P-OP1	-5.17	101.05	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	639	G	N9-C1'-C2'	-5.17	106.32	112.00
36	1	897	U	N3-C4-O4	-5.17	115.78	119.40
36	1	933	A	C4-C5-N7	5.17	113.28	110.70
36	1	1929	G	N3-C2-N2	5.17	123.52	119.90
1	6	1455	G	N1-C6-O6	5.17	123.00	119.90
36	5	406	G	O4'-C1'-N9	5.17	112.33	108.20
36	5	1794	G	N3-C4-C5	5.17	131.18	128.60
36	5	2379	U	C4-C5-C6	5.17	122.80	119.70
36	5	2978	U	C5-C6-N1	-5.17	120.12	122.70
36	5	3109	G	C5-N7-C8	-5.17	101.72	104.30
1	2	1428	G	O5'-P-OP1	-5.17	101.05	105.70
36	1	906	A	C6-C5-N7	-5.17	128.68	132.30
36	1	1301	A	C8-N9-C4	-5.17	103.73	105.80
1	6	76	A	OP1-P-O3'	5.17	116.56	105.20
36	5	1607	U	OP1-P-O3'	5.17	116.56	105.20
36	5	2211	U	C5-C4-O4	5.17	129.00	125.90
36	5	2641	U	C4-C5-C6	5.17	122.80	119.70
36	5	3330	A	C8-N9-C4	-5.17	103.73	105.80
38	8	16	G	O5'-P-OP1	-5.17	101.05	105.70
1	2	4	C	C6-N1-C2	-5.16	118.23	120.30
35	SM	134	ASP	CB-CG-OD2	5.16	122.95	118.30
36	1	699	A	N1-C2-N3	5.16	131.88	129.30
36	1	1643	A	N1-C6-N6	5.16	121.70	118.60
36	1	2148	U	O5'-P-OP2	5.16	116.90	110.70
36	1	2806	U	N1-C2-O2	-5.16	119.19	122.80
1	6	901	G	N9-C4-C5	-5.16	103.33	105.40
1	6	1187	U	C5-C6-N1	5.16	125.28	122.70
36	5	513	G	C4-C5-N7	-5.16	108.73	110.80
36	5	937	G	N3-C4-N9	5.16	129.10	126.00
36	1	498	A	C4-C5-C6	5.16	119.58	117.00
36	1	2622	C	N3-C4-C5	-5.16	119.83	121.90
36	1	3242	G	C8-N9-C1'	5.16	133.71	127.00
36	5	709	A	C5-C6-N6	-5.16	119.57	123.70
36	1	2306	C	C6-N1-C1'	-5.16	114.61	120.80
36	1	2612	U	C5-C6-N1	-5.16	120.12	122.70
1	6	1697	G	C2-N3-C4	5.16	114.48	111.90
36	5	917	A	C5-N7-C8	-5.16	101.32	103.90
36	5	1661	G	C5-C6-O6	-5.16	125.50	128.60
36	5	2310	U	N3-C2-O2	-5.16	118.59	122.20
36	5	3289	G	C8-N9-C4	-5.16	104.34	106.40
36	1	2245	C	C5-C6-N1	5.16	123.58	121.00
36	1	2283	G	C5-C6-O6	-5.16	125.50	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
37	7	75	G	C8-N9-C1'	-5.16	120.29	127.00
36	5	716	A	N9-C4-C5	-5.16	103.74	105.80
36	5	1852	G	C8-N9-C4	-5.16	104.34	106.40
36	1	123	A	C6-C5-N7	-5.16	128.69	132.30
1	6	217	A	P-O3'-C3'	5.16	125.89	119.70
1	6	1793	G	N1-C6-O6	-5.16	116.81	119.90
36	5	297	G	O4'-C1'-N9	5.16	112.33	108.20
36	5	1151	U	N3-C4-O4	5.16	123.01	119.40
36	5	1839	A	N7-C8-N9	-5.16	111.22	113.80
36	1	961	C	O4'-C1'-N1	-5.15	104.08	108.20
1	6	155	U	N3-C2-O2	-5.15	118.59	122.20
36	5	359	U	C5-C6-N1	5.15	125.28	122.70
36	5	979	U	O4'-C1'-N1	5.15	112.32	108.20
46	19	176	LEU	CA-CB-CG	-5.15	103.45	115.30
1	2	937	C	N1-C2-O2	5.15	121.99	118.90
36	1	2631	U	C2-N3-C4	-5.15	123.91	127.00
36	5	1832	C	C2-N1-C1'	-5.15	113.13	118.80
36	5	2316	G	N3-C4-N9	5.15	129.09	126.00
36	5	2351	U	N3-C4-O4	-5.15	115.79	119.40
37	7	87	G	C5-C6-N1	-5.15	108.92	111.50
78	q2	104	LEU	CA-CB-CG	5.15	127.15	115.30
36	1	199	A	O4'-C1'-N9	5.15	112.32	108.20
36	1	676	G	N1-C2-N2	-5.15	111.57	116.20
36	1	1103	A	N9-C4-C5	-5.15	103.74	105.80
1	2	599	A	OP2-P-O3'	5.15	116.53	105.20
36	1	335	G	C8-N9-C1'	5.15	133.69	127.00
36	1	908	G	N9-C4-C5	-5.15	103.34	105.40
36	1	914	A	C5-N7-C8	5.15	106.47	103.90
36	1	2427	U	N3-C2-O2	-5.15	118.60	122.20
36	1	2627	C	C5-C6-N1	-5.15	118.43	121.00
36	1	2848	G	OP2-P-O3'	5.15	116.53	105.20
37	3	98	C	N1-C2-O2	-5.15	115.81	118.90
38	4	27	U	N1-C2-O2	5.15	126.40	122.80
1	6	470	A	C8-N9-C4	-5.15	103.74	105.80
36	5	1101	G	C8-N9-C4	5.15	108.46	106.40
36	5	1368	U	O5'-P-OP1	-5.15	101.07	105.70
36	5	2973	G	C8-N9-C4	-5.15	104.34	106.40
36	5	3028	G	C6-C5-N7	-5.15	127.31	130.40
36	5	2807	U	N3-C4-O4	5.15	123.00	119.40
36	1	2261	G	C4-N9-C1'	5.14	133.19	126.50
1	6	6	G	O5'-P-OP1	-5.14	101.07	105.70
1	6	209	U	N1-C2-O2	-5.14	119.20	122.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	411	U	O5'-P-OP2	-5.14	101.07	105.70
36	5	991	G	C5-C6-O6	5.14	131.69	128.60
36	5	2195	C	C5-C4-N4	-5.14	116.60	120.20
36	5	2777	G	C8-N9-C1'	5.14	133.69	127.00
36	5	2982	A	C2-N3-C4	5.14	113.17	110.60
36	1	309	U	N3-C2-O2	-5.14	118.60	122.20
36	1	722	G	C2-N3-C4	-5.14	109.33	111.90
36	1	775	A	C2-N3-C4	-5.14	108.03	110.60
36	1	1889	G	C6-C5-N7	-5.14	127.31	130.40
36	1	2747	A	N9-C4-C5	5.14	107.86	105.80
36	1	2918	G	C4-N9-C1'	5.14	133.19	126.50
1	6	337	G	N3-C2-N2	5.14	123.50	119.90
36	5	1744	G	C6-C5-N7	-5.14	127.31	130.40
36	1	2632	G	C6-C5-N7	-5.14	127.31	130.40
36	1	49	A	C2-N3-C4	-5.14	108.03	110.60
36	1	934	G	C5-C6-O6	-5.14	125.52	128.60
36	1	2531	C	C2-N1-C1'	5.14	124.45	118.80
36	1	2595	A	O4'-C1'-N9	5.14	112.31	108.20
1	6	1662	G	O5'-P-OP2	-5.14	101.07	105.70
36	5	712	G	O5'-P-OP2	-5.14	101.08	105.70
36	5	1359	C	C5-C6-N1	-5.14	118.43	121.00
36	5	1406	A	N9-C4-C5	5.14	107.86	105.80
36	5	2855	U	N1-C2-N3	5.14	117.98	114.90
37	7	8	G	N1-C2-N2	-5.14	111.57	116.20
36	5	967	A	N9-C4-C5	5.14	107.86	105.80
36	5	2938	G	C8-N9-C4	-5.14	104.34	106.40
36	1	780	A	C4-C5-C6	5.14	119.57	117.00
36	1	923	C	C6-N1-C2	5.14	122.35	120.30
36	1	1492	G	N9-C4-C5	-5.14	103.34	105.40
1	6	1297	G	N1-C6-O6	-5.14	116.82	119.90
36	5	22	G	N1-C6-O6	5.14	122.98	119.90
36	5	2322	C	N3-C4-N4	5.14	121.60	118.00
36	1	62	A	O5'-P-OP1	5.13	116.86	110.70
36	1	3173	G	C4-N9-C1'	5.13	133.18	126.50
1	6	1016	C	C6-N1-C2	-5.13	118.25	120.30
1	6	1481	C	N1-C2-O2	5.13	121.98	118.90
36	5	1049	C	C6-N1-C2	-5.13	118.25	120.30
36	5	2885	C	N3-C4-N4	5.13	121.59	118.00
36	5	3122	A	O5'-P-OP1	-5.13	101.08	105.70
1	2	1611	A	N1-C2-N3	5.13	131.87	129.30
36	1	860	G	N3-C2-N2	-5.13	116.31	119.90
36	1	1581	C	C2-N1-C1'	5.13	124.45	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
61	N5	115	ARG	NE-CZ-NH1	5.13	122.87	120.30
36	5	2282	U	N3-C2-O2	5.13	125.79	122.20
36	1	2905	U	N3-C2-O2	5.13	125.79	122.20
36	1	3178	A	N1-C6-N6	5.13	121.68	118.60
1	6	976	G	C6-C5-N7	-5.13	127.32	130.40
1	6	1081	A	O4'-C1'-N9	5.13	112.31	108.20
5	s3	161	GLY	N-CA-C	5.13	125.93	113.10
36	1	2261	G	C8-N9-C1'	-5.13	120.33	127.00
36	1	2363	A	C5-C6-N1	-5.13	115.14	117.70
36	1	2370	G	C4-C5-N7	-5.13	108.75	110.80
36	5	2183	A	C8-N9-C4	5.13	107.85	105.80
36	5	2303	A	C5-C6-N6	-5.13	119.60	123.70
1	2	1284	C	N1-C2-O2	5.13	121.98	118.90
36	1	110	G	C6-C5-N7	-5.13	127.32	130.40
36	1	340	C	N3-C2-O2	-5.13	118.31	121.90
36	1	698	U	C6-N1-C1'	5.13	128.38	121.20
1	6	477	A	N1-C6-N6	5.13	121.68	118.60
36	5	995	U	C5-C6-N1	-5.13	120.14	122.70
36	5	1500	G	C5-C6-O6	-5.13	125.52	128.60
36	1	725	G	C5-C6-O6	-5.13	125.52	128.60
36	1	757	C	N1-C2-O2	-5.13	115.82	118.90
36	1	2911	A	N1-C6-N6	-5.13	115.52	118.60
1	6	647	G	N3-C4-N9	-5.13	122.92	126.00
1	6	1036	A	O5'-P-OP2	-5.13	101.09	105.70
1	6	1146	G	N1-C6-O6	5.13	122.97	119.90
1	6	1663	G	C8-N9-C4	-5.13	104.35	106.40
36	5	2526	C	N1-C2-O2	5.13	121.98	118.90
36	5	3294	A	N1-C2-N3	5.13	131.86	129.30
36	5	3362	A	C4-C5-N7	5.13	113.26	110.70
13	C1	90	TYR	CA-CB-CG	5.12	123.14	113.40
36	1	1912	U	N3-C4-C5	-5.12	111.53	114.60
38	4	95	G	N3-C4-C5	5.12	131.16	128.60
1	6	539	G	O4'-C1'-N9	-5.12	104.10	108.20
1	6	1028	C	N3-C4-C5	5.12	123.95	121.90
36	5	1213	G	C5-C6-N1	5.12	114.06	111.50
1	2	1778	G	C4-C5-N7	5.12	112.85	110.80
36	1	780	A	C6-C5-N7	-5.12	128.71	132.30
36	1	1154	A	O5'-P-OP1	-5.12	101.09	105.70
41	L4	57	GLY	N-CA-C	-5.12	100.29	113.10
1	2	1127	G	C5-C6-N1	-5.12	108.94	111.50
36	1	947	G	C4-N9-C1'	5.12	133.16	126.50
36	1	2610	G	C6-C5-N7	-5.12	127.33	130.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	3155	U	N1-C2-O2	5.12	126.39	122.80
36	5	1115	G	C6-C5-N7	-5.12	127.33	130.40
36	5	1317	A	C8-N9-C4	-5.12	103.75	105.80
36	5	1927	G	N3-C4-N9	5.12	129.07	126.00
36	1	404	G	C8-N9-C4	-5.12	104.35	106.40
36	5	2141	U	N1-C2-N3	5.12	117.97	114.90
36	5	2413	A	N7-C8-N9	-5.12	111.24	113.80
36	1	99	A	C5'-C4'-O4'	5.12	115.24	109.10
36	1	880	G	N1-C6-O6	-5.12	116.83	119.90
36	1	1404	G	C5-C6-O6	5.12	131.67	128.60
36	1	1556	C	N3-C2-O2	-5.12	118.32	121.90
36	1	2858	U	C5-C4-O4	5.12	128.97	125.90
49	M3	67	ARG	NE-CZ-NH1	-5.12	117.74	120.30
1	6	275	C	C2-N1-C1'	5.12	124.43	118.80
36	5	728	G	C8-N9-C1'	-5.12	120.35	127.00
36	5	895	A	C2-N3-C4	-5.12	108.04	110.60
36	5	1147	G	N7-C8-N9	5.12	115.66	113.10
36	5	1178	G	C5-C6-N1	-5.12	108.94	111.50
36	5	1620	U	N3-C2-O2	-5.12	118.62	122.20
36	5	2385	G	C8-N9-C4	5.12	108.45	106.40
36	5	2675	C	C6-N1-C2	-5.12	118.25	120.30
36	1	2209	U	C5-C6-N1	5.12	125.26	122.70
36	5	661	G	O5'-P-OP1	-5.12	101.09	105.70
36	5	1853	U	N1-C2-O2	-5.12	119.22	122.80
36	5	2794	G	N9-C4-C5	-5.12	103.35	105.40
36	5	2885	C	N3-C2-O2	5.12	125.48	121.90
36	1	18	G	N3-C4-C5	5.12	131.16	128.60
36	1	517	G	N1-C6-O6	-5.12	116.83	119.90
36	1	637	C	C2-N1-C1'	5.12	124.43	118.80
36	1	1321	G	O4'-C1'-N9	5.12	112.29	108.20
36	1	1375	G	C6-C5-N7	-5.12	127.33	130.40
36	1	1694	U	N3-C2-O2	-5.12	118.62	122.20
36	1	2727	A	C5-N7-C8	5.12	106.46	103.90
36	1	2828	G	C8-N9-C1'	-5.12	120.35	127.00
38	4	7	U	N1-C2-O2	-5.12	119.22	122.80
38	4	66	A	N7-C8-N9	5.12	116.36	113.80
36	5	938	C	C5-C4-N4	-5.12	116.62	120.20
36	5	1342	C	C2-N3-C4	-5.12	117.34	119.90
36	5	2994	A	C5-C6-N1	5.12	120.26	117.70
36	1	1010	G	N1-C6-O6	5.11	122.97	119.90
36	1	2807	U	C5-C6-N1	5.11	125.26	122.70
1	6	116	U	N1-C2-N3	5.11	117.97	114.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	609	G	O5'-P-OP2	-5.11	101.10	105.70
36	5	1119	C	OP2-P-O3'	5.11	116.45	105.20
36	5	1432	C	C6-N1-C2	-5.11	118.25	120.30
36	5	2672	G	C5-C6-O6	-5.11	125.53	128.60
53	m7	41	LEU	CA-CB-CG	5.11	127.06	115.30
1	2	75	U	N3-C2-O2	-5.11	118.62	122.20
36	1	2524	A	O4'-C1'-N9	5.11	112.29	108.20
1	6	25	C	OP2-P-O3'	5.11	116.45	105.20
1	6	39	A	O4'-C1'-N9	5.11	112.29	108.20
36	5	437	G	N7-C8-N9	5.11	115.66	113.10
36	5	2237	C	C6-N1-C2	5.11	122.34	120.30
36	5	3030	G	O5'-P-OP1	-5.11	101.10	105.70
1	2	1748	G	C2-N3-C4	-5.11	109.34	111.90
36	1	1048	A	N1-C6-N6	-5.11	115.53	118.60
36	1	1417	G	N3-C4-C5	5.11	131.16	128.60
36	1	1496	C	C2-N1-C1'	5.11	124.42	118.80
36	1	2647	A	C6-C5-N7	-5.11	128.72	132.30
36	5	1214	U	C2-N1-C1'	5.11	123.83	117.70
36	5	1548	C	N3-C2-O2	5.11	125.48	121.90
38	4	60	U	N3-C2-O2	-5.11	118.62	122.20
36	5	1902	G	N1-C2-N3	5.11	126.97	123.90
36	1	282	G	C2'-C3'-O3'	5.11	121.87	113.70
36	1	394	G	C8-N9-C1'	5.11	133.64	127.00
36	1	638	C	C6-N1-C1'	-5.11	114.67	120.80
36	1	863	C	C5-C4-N4	-5.11	116.62	120.20
36	1	1661	G	N1-C2-N2	-5.11	111.60	116.20
1	6	603	U	N3-C4-O4	5.11	122.98	119.40
1	6	1517	U	C2-N1-C1'	5.11	123.83	117.70
36	5	2145	A	N7-C8-N9	5.11	116.35	113.80
51	m5	96	ARG	NE-CZ-NH1	5.11	122.85	120.30
36	1	75	G	O5'-P-OP1	5.11	116.83	110.70
36	1	1385	C	N3-C4-C5	-5.11	119.86	121.90
36	5	87	U	C2-N1-C1'	5.11	123.83	117.70
36	5	726	G	N3-C4-C5	5.11	131.15	128.60
36	5	726	G	N3-C4-N9	-5.11	122.94	126.00
36	5	1450	G	N1-C2-N2	5.11	120.80	116.20
36	5	3098	G	N9-C4-C5	5.11	107.44	105.40
36	5	3214	U	O4'-C1'-N1	5.11	112.28	108.20
1	2	1412	G	C8-N9-C4	5.10	108.44	106.40
36	1	1472	U	N1-C2-O2	-5.10	119.23	122.80
1	6	457	G	N1-C6-O6	5.10	122.96	119.90
36	5	1667	A	C8-N9-C4	5.10	107.84	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2190	U	N1-C2-N3	5.10	117.96	114.90
1	6	1196	A	N1-C6-N6	-5.10	115.54	118.60
36	5	879	U	C5-C4-O4	-5.10	122.84	125.90
36	5	1331	U	C2-N3-C4	-5.10	123.94	127.00
36	5	1335	C	N3-C4-N4	5.10	121.57	118.00
36	5	3055	U	C5-C4-O4	-5.10	122.84	125.90
36	5	3344	A	C8-N9-C4	-5.10	103.76	105.80
37	7	37	G	C4-C5-N7	5.10	112.84	110.80
36	1	2732	G	N1-C6-O6	-5.10	116.84	119.90
36	5	1142	G	C8-N9-C4	-5.10	104.36	106.40
36	5	1375	G	OP2-P-O3'	5.10	116.42	105.20
36	5	3354	U	N3-C2-O2	-5.10	118.63	122.20
1	2	1116	A	C2-N3-C4	-5.10	108.05	110.60
36	1	968	G	C4-C5-C6	5.10	121.86	118.80
51	M5	93	LYS	N-CA-C	5.10	124.77	111.00
36	5	364	G	C5-C6-O6	-5.10	125.54	128.60
36	5	567	G	C8-N9-C1'	-5.10	120.37	127.00
36	5	2827	U	C6-N1-C1'	-5.10	114.06	121.20
36	5	3361	G	N3-C4-C5	-5.10	126.05	128.60
36	1	2354	C	N3-C4-C5	-5.10	119.86	121.90
37	3	119	U	C5-C6-N1	-5.10	120.15	122.70
38	4	105	A	C8-N9-C4	-5.10	103.76	105.80
36	5	1379	G	N3-C4-N9	5.10	129.06	126.00
36	5	3040	A	N1-C2-N3	5.10	131.85	129.30
38	4	95	G	N3-C4-N9	-5.10	122.94	126.00
36	5	2859	U	C5-C6-N1	-5.10	120.15	122.70
1	2	318	U	N3-C2-O2	5.09	125.77	122.20
36	1	189	G	N9-C4-C5	5.09	107.44	105.40
36	1	282	G	N7-C8-N9	5.09	115.65	113.10
36	1	663	C	C4-C5-C6	5.09	119.95	117.40
36	1	1400	G	C8-N9-C1'	-5.09	120.38	127.00
36	1	1404	G	C5-N7-C8	5.09	106.85	104.30
36	1	2519	A	N1-C6-N6	5.09	121.66	118.60
36	5	400	G	N3-C4-N9	-5.09	122.94	126.00
36	5	1858	A	C2-N3-C4	5.09	113.15	110.60
36	5	2123	G	N3-C4-N9	5.09	129.06	126.00
36	5	2565	U	O4'-C1'-N1	5.09	112.28	108.20
36	5	2755	C	O5'-P-OP1	-5.09	101.11	105.70
1	2	1633	A	N1-C6-N6	-5.09	115.54	118.60
36	1	1106	G	N3-C2-N2	-5.09	116.33	119.90
36	1	1307	G	N3-C2-N2	-5.09	116.33	119.90
36	1	2370	G	O5'-P-OP1	-5.09	101.12	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	349	A	C5-C6-N1	5.09	120.25	117.70
36	1	895	A	N7-C8-N9	5.09	116.35	113.80
36	1	2175	U	N3-C2-O2	-5.09	118.64	122.20
36	1	3140	G	C5-C6-O6	-5.09	125.55	128.60
51	M5	68	ARG	NE-CZ-NH1	5.09	122.85	120.30
1	6	346	G	N3-C4-C5	-5.09	126.05	128.60
36	5	813	G	N9-C4-C5	-5.09	103.36	105.40
36	5	2928	C	N3-C4-N4	5.09	121.56	118.00
36	5	1473	G	C8-N9-C4	5.09	108.44	106.40
36	5	1546	A	C4-C5-C6	5.09	119.55	117.00
36	5	2624	G	C4-C5-N7	5.09	112.84	110.80
36	5	2695	A	O4'-C1'-N9	5.09	112.27	108.20
36	5	3096	C	N1-C2-O2	-5.09	115.85	118.90
1	2	1644	C	N3-C2-O2	-5.09	118.34	121.90
36	1	32	U	N3-C4-O4	5.09	122.96	119.40
36	1	320	G	OP2-P-O3'	5.09	116.39	105.20
36	1	1386	A	C6-C5-N7	-5.09	128.74	132.30
1	6	1595	U	N3-C4-C5	-5.09	111.55	114.60
36	5	220	G	C4-C5-N7	5.09	112.83	110.80
1	2	139	C	C6-N1-C2	-5.09	118.27	120.30
36	1	2572	C	N3-C2-O2	-5.09	118.34	121.90
36	5	1143	A	N1-C6-N6	5.09	121.65	118.60
36	5	2938	G	N1-C6-O6	5.09	122.95	119.90
36	1	1192	C	O5'-P-OP2	-5.08	101.12	105.70
36	1	3243	A	C6-C5-N7	-5.08	128.74	132.30
36	5	1838	G	N1-C6-O6	5.08	122.95	119.90
36	1	38	U	N1-C2-N3	-5.08	111.85	114.90
36	1	1364	C	OP2-P-O3'	5.08	116.38	105.20
36	5	757	C	C2-N3-C4	-5.08	117.36	119.90
36	1	2153	U	C6-N1-C2	-5.08	117.95	121.00
36	1	2309	A	C4-C5-N7	5.08	113.24	110.70
36	1	2601	A	N1-C6-N6	5.08	121.65	118.60
36	1	3121	U	OP1-P-O3'	5.08	116.38	105.20
36	5	1043	C	OP2-P-O3'	5.08	116.38	105.20
36	5	1169	A	N7-C8-N9	-5.08	111.26	113.80
36	5	3294	A	C8-N9-C4	-5.08	103.77	105.80
37	3	80	G	C4-C5-C6	5.08	121.85	118.80
1	6	614	C	N1-C2-O2	5.08	121.95	118.90
1	2	1539	G	C8-N9-C4	-5.08	104.37	106.40
1	2	1773	C	N1-C2-O2	-5.08	115.85	118.90
36	1	281	G	C8-N9-C4	-5.08	104.37	106.40
36	1	669	U	N3-C4-C5	5.08	117.65	114.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2549	G	N3-C4-C5	-5.08	126.06	128.60
36	5	110	G	C5-N7-C8	5.08	106.84	104.30
36	5	968	G	C4-N9-C1'	5.08	133.10	126.50
36	5	1373	A	C6-C5-N7	-5.08	128.75	132.30
36	5	2673	A	N1-C6-N6	-5.08	115.55	118.60
36	1	1012	G	C8-N9-C4	5.08	108.43	106.40
36	1	1145	G	C2-N3-C4	5.08	114.44	111.90
36	1	1844	C	N1-C2-O2	-5.08	115.85	118.90
36	1	3206	C	N1-C2-O2	5.08	121.95	118.90
36	5	574	U	C5-C6-N1	-5.08	120.16	122.70
1	2	1169	G	N1-C6-O6	-5.08	116.86	119.90
1	2	1297	G	C4-C5-N7	5.08	112.83	110.80
36	1	1190	A	C8-N9-C1'	-5.08	118.56	127.70
54	M8	140	LEU	CA-CB-CG	-5.08	103.62	115.30
36	5	999	G	N9-C4-C5	-5.08	103.37	105.40
1	2	16	G	C8-N9-C1'	-5.07	120.41	127.00
36	1	1137	C	C6-N1-C2	5.07	122.33	120.30
36	1	1307	G	C4-C5-N7	-5.07	108.77	110.80
36	1	1309	U	N3-C2-O2	-5.07	118.65	122.20
36	1	1892	G	C4-C5-N7	-5.07	108.77	110.80
36	1	3182	G	C4-C5-N7	5.07	112.83	110.80
1	6	965	U	C4-C5-C6	-5.07	116.66	119.70
36	5	692	A	O5'-P-OP1	-5.07	101.13	105.70
36	5	2198	A	C5-N7-C8	-5.07	101.36	103.90
37	7	52	G	C8-N9-C4	5.07	108.43	106.40
36	1	1581	C	N3-C2-O2	-5.07	118.35	121.90
36	5	86	G	N3-C4-N9	5.07	129.04	126.00
36	1	1515	A	N1-C6-N6	5.07	121.64	118.60
1	6	557	G	N1-C6-O6	-5.07	116.86	119.90
36	5	39	A	C5-N7-C8	-5.07	101.36	103.90
36	5	2726	C	C4-C5-C6	5.07	119.94	117.40
36	5	2748	A	C5-C6-N6	-5.07	119.64	123.70
37	3	89	G	N3-C4-N9	5.07	129.04	126.00
1	6	1130	G	N3-C4-N9	-5.07	122.96	126.00
36	5	1211	U	C4-C5-C6	-5.07	116.66	119.70
36	5	2908	G	N3-C4-N9	-5.07	122.96	126.00
1	2	913	G	P-O3'-C3'	5.07	125.78	119.70
1	2	1600	A	C5-N7-C8	-5.07	101.37	103.90
36	1	614	C	C6-N1-C2	5.07	122.33	120.30
36	1	941	G	C6-C5-N7	-5.07	127.36	130.40
36	1	1939	G	N3-C4-C5	-5.07	126.07	128.60
36	1	2257	C	O4'-C1'-N1	5.07	112.25	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2311	G	N9-C4-C5	5.07	107.43	105.40
36	1	2937	G	N7-C8-N9	-5.07	110.57	113.10
36	1	3173	G	C8-N9-C1'	-5.07	120.41	127.00
36	1	3375	A	C5'-C4'-C3'	-5.07	107.89	116.00
1	6	7	G	N3-C4-C5	-5.07	126.07	128.60
1	6	359	A	N3-C4-N9	-5.07	123.35	127.40
36	5	991	G	C4-C5-N7	-5.07	108.77	110.80
36	5	1193	A	C4-C5-C6	5.07	119.53	117.00
36	5	1306	G	N7-C8-N9	5.07	115.63	113.10
36	1	2400	G	OP2-P-O3'	5.07	116.35	105.20
36	1	2683	U	C5-C6-N1	-5.07	120.17	122.70
37	3	58	C	N3-C2-O2	-5.07	118.35	121.90
1	6	985	G	C2-N3-C4	-5.07	109.37	111.90
36	5	396	A	C5-C6-N1	-5.07	115.17	117.70
36	5	1177	G	N9-C4-C5	5.07	107.43	105.40
36	5	3315	G	N1-C2-N3	5.07	126.94	123.90
36	1	1119	C	C2-N1-C1'	-5.06	113.23	118.80
36	1	1389	G	C8-N9-C4	5.06	108.43	106.40
36	1	3277	U	N1-C2-O2	5.06	126.34	122.80
36	5	83	U	N3-C2-O2	-5.06	118.66	122.20
36	1	1207	G	N1-C6-O6	5.06	122.94	119.90
1	6	431	C	N1-C2-O2	-5.06	115.86	118.90
36	5	1101	G	N1-C2-N2	-5.06	111.64	116.20
36	5	1507	G	N1-C2-N2	-5.06	111.64	116.20
37	7	35	C	N3-C2-O2	-5.06	118.36	121.90
36	1	634	C	C5-C6-N1	-5.06	118.47	121.00
38	4	112	U	C2-N1-C1'	-5.06	111.63	117.70
1	6	1653	C	C6-N1-C2	-5.06	118.28	120.30
36	5	2932	U	O5'-P-OP2	-5.06	101.14	105.70
36	1	1587	A	N1-C6-N6	-5.06	115.56	118.60
36	1	2860	U	N3-C4-C5	5.06	117.64	114.60
36	1	3054	U	N3-C2-O2	-5.06	118.66	122.20
36	1	3243	A	N9-C4-C5	-5.06	103.78	105.80
1	6	620	A	C8-N9-C4	-5.06	103.78	105.80
1	6	1354	G	C4-N9-C1'	5.06	133.08	126.50
36	5	922	U	C2-N3-C4	-5.06	123.96	127.00
36	5	1665	C	N3-C2-O2	-5.06	118.36	121.90
36	5	3296	A	OP2-P-O3'	5.06	116.33	105.20
36	1	940	G	C5-C6-O6	5.06	131.63	128.60
36	1	2688	U	O4'-C1'-N1	-5.06	104.15	108.20
36	1	2989	U	C2-N1-C1'	5.06	123.77	117.70
38	4	52	A	C4-C5-N7	-5.06	108.17	110.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	5	96	G	OP1-P-OP2	5.06	127.19	119.60
36	5	927	C	O5'-P-OP1	-5.06	101.15	105.70
36	5	2377	G	N3-C2-N2	5.06	123.44	119.90
36	5	3247	G	C8-N9-C4	5.06	108.42	106.40
1	2	933	A	C8-N9-C4	-5.06	103.78	105.80
1	2	1778	G	N7-C8-N9	5.06	115.63	113.10
36	5	524	U	O5'-P-OP2	-5.06	101.15	105.70
36	1	1084	A	C6-C5-N7	-5.05	128.76	132.30
36	1	1401	A	C2-N3-C4	-5.05	108.07	110.60
36	1	1451	C	N1-C2-O2	5.05	121.93	118.90
36	1	1820	U	OP2-P-O3'	5.05	116.32	105.20
36	1	1851	G	N7-C8-N9	5.05	115.63	113.10
36	1	2194	G	C4-C5-N7	5.05	112.82	110.80
36	1	2415	C	C2-N1-C1'	-5.05	113.24	118.80
36	1	2897	A	C8-N9-C4	5.05	107.82	105.80
1	6	1274	C	C6-N1-C1'	-5.05	114.73	120.80
36	5	331	G	O5'-P-OP1	-5.05	101.15	105.70
36	5	666	A	C5-C6-N6	5.05	127.74	123.70
36	5	852	U	OP2-P-O3'	5.05	116.32	105.20
36	5	1370	G	N1-C6-O6	-5.05	116.87	119.90
36	5	1420	C	C2-N1-C1'	-5.05	113.24	118.80
36	5	1502	C	N1-C2-O2	5.05	121.93	118.90
1	2	9	U	O5'-P-OP1	-5.05	101.15	105.70
36	1	1157	G	C6-N1-C2	-5.05	122.07	125.10
36	1	2804	A	N1-C2-N3	5.05	131.83	129.30
36	5	1178	G	O5'-P-OP1	-5.05	101.15	105.70
36	5	1452	A	C5-N7-C8	-5.05	101.37	103.90
36	5	2195	C	N3-C4-N4	5.05	121.54	118.00
36	5	2857	C	N3-C4-C5	5.05	123.92	121.90
1	6	351	C	O4'-C1'-N1	-5.05	104.16	108.20
1	6	565	C	C2-N3-C4	-5.05	117.38	119.90
36	5	411	U	C2-N1-C1'	-5.05	111.64	117.70
36	5	691	A	OP1-P-O3'	5.05	116.31	105.20
1	2	1370	U	P-O3'-C3'	5.05	125.76	119.70
1	2	1615	C	C6-N1-C2	-5.05	118.28	120.30
36	1	2865	U	N1-C2-O2	5.05	126.33	122.80
36	1	3178	A	N9-C4-C5	-5.05	103.78	105.80
1	6	1744	A	C8-N9-C4	5.05	107.82	105.80
36	5	2374	C	O5'-P-OP2	-5.05	101.16	105.70
36	5	2888	U	C2-N1-C1'	5.05	123.76	117.70
36	1	996	A	OP2-P-O3'	5.05	116.31	105.20
36	1	1351	U	C2-N1-C1'	5.05	123.76	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2395	G	OP2-P-O3'	5.05	116.31	105.20
38	4	85	G	N7-C8-N9	5.05	115.62	113.10
1	6	1150	G	N3-C4-N9	-5.05	122.97	126.00
36	5	283	G	O4'-C1'-N9	-5.05	104.16	108.20
36	5	2796	G	N3-C4-C5	-5.05	126.08	128.60
36	5	2849	C	N3-C4-C5	-5.05	119.88	121.90
36	5	3164	C	C6-N1-C2	5.05	122.32	120.30
1	2	318	U	C6-N1-C2	5.05	124.03	121.00
36	1	349	A	OP2-P-O3'	5.05	116.30	105.20
36	1	1227	C	C5-C6-N1	5.05	123.52	121.00
36	1	1592	G	C5-C6-N1	-5.05	108.98	111.50
36	1	2714	G	C4-C5-C6	-5.05	115.77	118.80
1	6	344	A	C8-N9-C4	5.05	107.82	105.80
1	2	1535	U	O5'-P-OP1	5.04	116.75	110.70
36	1	304	G	N3-C2-N2	-5.04	116.37	119.90
36	1	1440	G	C8-N9-C4	5.04	108.42	106.40
36	5	523	A	N9-C4-C5	5.04	107.82	105.80
36	1	511	G	C6-C5-N7	-5.04	127.37	130.40
36	1	1198	C	N3-C4-C5	-5.04	119.88	121.90
36	1	1311	G	C4-C5-N7	5.04	112.82	110.80
36	1	1323	G	C8-N9-C4	5.04	108.42	106.40
36	1	1446	A	N9-C4-C5	5.04	107.82	105.80
36	1	2978	U	O4'-C1'-N1	5.04	112.23	108.20
1	6	20	G	N1-C6-O6	5.04	122.93	119.90
36	5	2952	G	C8-N9-C4	-5.04	104.38	106.40
36	5	3091	A	C4-C5-C6	5.04	119.52	117.00
1	2	1486	G	C4-N9-C1'	5.04	133.05	126.50
36	1	342	A	N3-C4-N9	-5.04	123.37	127.40
36	1	1850	A	O5'-P-OP2	-5.04	101.16	105.70
36	1	2989	U	C5-C6-N1	5.04	125.22	122.70
1	6	41	A	N1-C6-N6	5.04	121.62	118.60
1	6	1285	U	N1-C2-O2	5.04	126.33	122.80
1	6	1539	G	O4'-C1'-N9	-5.04	104.17	108.20
36	5	2897	A	O4'-C1'-N9	5.04	112.23	108.20
36	1	2525	G	C4-N9-C1'	5.04	133.05	126.50
1	6	2	A	N1-C6-N6	5.04	121.62	118.60
1	6	711	U	C2-N1-C1'	5.04	123.75	117.70
36	5	639	G	C6-C5-N7	-5.04	127.38	130.40
36	5	3149	G	O5'-P-OP1	5.04	116.75	110.70
1	6	154	G	C8-N9-C4	-5.04	104.39	106.40
36	5	970	A	C4-C5-N7	5.04	113.22	110.70
36	5	3085	G	C5-C6-N1	-5.04	108.98	111.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	2	1052	U	N1-C2-O2	5.04	126.33	122.80
36	1	808	A	C8-N9-C4	5.04	107.81	105.80
36	1	691	A	C4-C5-C6	5.04	119.52	117.00
36	1	1143	A	N9-C4-C5	5.04	107.81	105.80
36	1	2684	C	C4-C5-C6	5.04	119.92	117.40
36	1	2989	U	C5-C4-O4	-5.04	122.88	125.90
36	5	41	G	O5'-P-OP2	-5.04	101.17	105.70
36	5	2403	G	N3-C4-N9	5.04	129.02	126.00
36	1	358	G	C2-N3-C4	-5.03	109.38	111.90
36	1	695	C	O5'-P-OP1	-5.03	101.17	105.70
36	1	936	A	C2-N3-C4	-5.03	108.08	110.60
36	1	1142	G	O5'-P-OP2	-5.03	101.17	105.70
36	1	1172	G	N1-C2-N2	5.03	120.73	116.20
36	1	1874	A	N1-C2-N3	5.03	131.82	129.30
36	1	2197	C	C6-N1-C2	5.03	122.31	120.30
36	1	2371	G	O5'-P-OP2	-5.03	101.17	105.70
36	1	2661	G	C6-C5-N7	-5.03	127.38	130.40
36	1	2687	G	C5-C6-O6	5.03	131.62	128.60
36	1	2824	G	C5-C6-N1	-5.03	108.98	111.50
36	1	2936	A	C8-N9-C4	-5.03	103.79	105.80
38	4	97	A	N1-C2-N3	5.03	131.82	129.30
1	6	530	C	C2-N1-C1'	-5.03	113.26	118.80
36	5	994	G	OP1-P-O3'	5.03	116.27	105.20
36	5	1494	U	O5'-P-OP2	5.03	116.74	110.70
36	5	2725	U	C4-C5-C6	-5.03	116.68	119.70
36	5	3164	C	N3-C4-C5	5.03	123.91	121.90
1	2	875	G	C6-C5-N7	-5.03	127.38	130.40
1	2	1412	G	N7-C8-N9	-5.03	110.58	113.10
1	2	1793	G	O4'-C1'-N9	5.03	112.23	108.20
36	1	551	A	O4'-C1'-N9	5.03	112.23	108.20
36	1	1112	A	N1-C2-N3	5.03	131.82	129.30
36	1	1518	U	C5-C6-N1	-5.03	120.18	122.70
36	1	2881	C	C2-N1-C1'	-5.03	113.27	118.80
36	1	3091	A	C5-N7-C8	-5.03	101.38	103.90
36	1	3228	C	C2-N1-C1'	5.03	124.33	118.80
37	3	105	C	C6-N1-C2	5.03	122.31	120.30
38	4	10	A	N7-C8-N9	-5.03	111.28	113.80
38	4	103	G	N3-C4-N9	5.03	129.02	126.00
36	5	339	C	C2-N1-C1'	-5.03	113.26	118.80
36	5	2322	C	C5-C6-N1	5.03	123.52	121.00
1	2	334	G	C2-N3-C4	-5.03	109.38	111.90
36	1	1544	G	N1-C6-O6	5.03	122.92	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	2321	A	N9-C4-C5	5.03	107.81	105.80
36	1	2627	C	OP2-P-O3'	5.03	116.27	105.20
44	L7	110	ARG	NE-CZ-NH2	-5.03	117.78	120.30
36	5	821	U	N3-C2-O2	-5.03	118.68	122.20
36	5	1374	G	N3-C4-C5	5.03	131.12	128.60
36	5	2160	G	C8-N9-C1'	-5.03	120.46	127.00
1	6	577	G	N3-C4-C5	5.03	131.12	128.60
36	5	842	G	N9-C4-C5	-5.03	103.39	105.40
36	5	3045	G	N3-C2-N2	-5.03	116.38	119.90
36	1	2343	C	OP2-P-O3'	5.03	116.26	105.20
36	1	2857	C	N1-C2-O2	5.03	121.92	118.90
36	5	2834	G	O5'-P-OP2	-5.03	101.18	105.70
36	5	2871	G	C5-C6-N1	5.03	114.01	111.50
36	5	2891	U	N1-C2-O2	5.03	126.32	122.80
36	5	2900	A	C5-C6-N6	5.03	127.72	123.70
1	2	49	C	C6-N1-C2	-5.03	118.29	120.30
36	1	1807	G	C8-N9-C1'	-5.03	120.47	127.00
36	1	2614	G	C8-N9-C4	5.03	108.41	106.40
36	1	2918	G	C4-C5-C6	5.03	121.82	118.80
36	1	3138	U	OP2-P-O3'	5.03	116.25	105.20
52	M6	84	LEU	CB-CG-CD2	-5.03	102.46	111.00
36	5	1148	G	C5-C6-N1	5.03	114.01	111.50
36	5	1889	G	C6-C5-N7	-5.03	127.39	130.40
1	6	969	C	N1-C2-O2	-5.02	115.89	118.90
36	5	2849	C	C5-C6-N1	5.02	123.51	121.00
36	1	1896	A	N7-C8-N9	5.02	116.31	113.80
37	3	87	G	OP2-P-O3'	5.02	116.25	105.20
36	5	413	U	N3-C4-O4	5.02	122.92	119.40
36	5	607	A	C5-C6-N6	5.02	127.72	123.70
36	5	2947	G	C5-N7-C8	-5.02	101.79	104.30
36	1	647	A	C2-N3-C4	-5.02	108.09	110.60
36	5	727	G	O5'-P-OP1	-5.02	101.18	105.70
36	1	2145	A	C4-C5-C6	5.02	119.51	117.00
36	1	2585	G	C8-N9-C4	-5.02	104.39	106.40
36	5	1292	C	N3-C2-O2	5.02	125.41	121.90
36	5	2347	U	C6-N1-C2	-5.02	117.99	121.00
1	2	1554	U	N1-C2-O2	-5.02	119.29	122.80
36	1	51	A	C2-N3-C4	-5.02	108.09	110.60
36	1	86	G	C4-C5-C6	-5.02	115.79	118.80
36	1	145	G	C5-N7-C8	-5.02	101.79	104.30
36	1	709	A	C4-C5-N7	5.02	113.21	110.70
36	1	1404	G	N7-C8-N9	-5.02	110.59	113.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	1694	U	C2-N1-C1'	5.02	123.72	117.70
36	5	2415	C	OP1-P-O3'	5.02	116.24	105.20
1	2	587	C	C2-N1-C1'	5.02	124.32	118.80
1	2	736	C	C2-N1-C1'	5.02	124.32	118.80
1	2	1479	A	N1-C6-N6	5.02	121.61	118.60
36	1	1725	C	C6-N1-C2	5.02	122.31	120.30
37	3	106	U	C5-C6-N1	5.02	125.21	122.70
36	5	658	G	C5-N7-C8	-5.02	101.79	104.30
36	5	2777	G	OP1-P-O3'	5.02	116.23	105.20
36	5	2813	A	C6-C5-N7	-5.02	128.79	132.30
36	5	2843	U	C2-N1-C1'	5.02	123.72	117.70
1	2	852	C	N1-C2-O2	5.01	121.91	118.90
36	1	766	U	C6-N1-C2	-5.01	117.99	121.00
37	3	45	A	N1-C6-N6	-5.01	115.59	118.60
1	6	1317	C	C6-N1-C2	-5.01	118.29	120.30
1	6	1696	G	P-O3'-C3'	5.01	125.72	119.70
36	5	8	C	C6-N1-C2	-5.01	118.29	120.30
36	5	912	G	OP2-P-O3'	5.01	116.23	105.20
36	5	1662	G	C6-C5-N7	-5.01	127.39	130.40
36	5	2815	G	C8-N9-C4	5.01	108.41	106.40
36	5	3337	G	C8-N9-C1'	-5.01	120.48	127.00
1	2	425	A	C5-C6-N6	-5.01	119.69	123.70
36	1	1184	A	O5'-P-OP2	-5.01	101.19	105.70
36	1	1435	A	N7-C8-N9	5.01	116.31	113.80
36	1	2610	G	C5-C6-O6	-5.01	125.59	128.60
36	5	1362	G	OP2-P-O3'	5.01	116.23	105.20
36	5	1920	U	C5-C4-O4	5.01	128.91	125.90
1	2	93	A	N7-C8-N9	5.01	116.31	113.80
1	2	163	G	C8-N9-C4	-5.01	104.39	106.40
36	1	80	G	N1-C2-N2	-5.01	111.69	116.20
36	1	1940	G	N1-C2-N3	5.01	126.91	123.90
36	1	2379	U	N3-C4-O4	5.01	122.91	119.40
36	1	2920	U	C2-N3-C4	-5.01	123.99	127.00
36	1	3134	A	N9-C4-C5	-5.01	103.80	105.80
57	N1	89	LEU	CA-CB-CG	5.01	126.83	115.30
1	6	364	G	C8-N9-C4	5.01	108.41	106.40
36	5	2321	A	OP2-P-O3'	5.01	116.23	105.20
36	5	2892	A	C2-N3-C4	-5.01	108.09	110.60
36	5	2946	A	C2-N3-C4	-5.01	108.09	110.60
36	5	3214	U	N1-C2-O2	5.01	126.31	122.80
36	5	3313	U	OP1-P-OP2	5.01	127.12	119.60
1	2	109	G	N1-C6-O6	5.01	122.91	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
36	1	636	C	C5-C6-N1	-5.01	118.50	121.00
36	1	703	G	C8-N9-C4	5.01	108.40	106.40
1	6	1127	G	C2-N3-C4	-5.01	109.40	111.90
1	6	1198	G	N3-C4-N9	-5.01	122.99	126.00
36	5	98	G	N3-C2-N2	5.01	123.41	119.90
36	5	1906	G	C4-C5-N7	5.01	112.80	110.80
36	5	2340	U	O5'-P-OP1	-5.01	101.19	105.70
36	1	2549	G	N9-C4-C5	-5.01	103.40	105.40
37	3	56	A	C8-N9-C4	5.01	107.80	105.80
38	4	27	U	N3-C2-O2	-5.01	118.69	122.20
36	5	660	A	C4-C5-N7	-5.01	108.20	110.70
1	6	96	G	OP2-P-O3'	5.01	116.21	105.20
36	5	567	G	C4-C5-C6	5.01	121.80	118.80
36	5	655	C	N1-C2-N3	5.01	122.70	119.20
36	5	1190	A	N9-C4-C5	5.01	107.80	105.80
36	5	1292	C	N3-C4-C5	5.01	123.90	121.90
36	5	3001	C	C6-N1-C2	5.01	122.30	120.30
36	1	1137	C	C5-C4-N4	-5.00	116.70	120.20
37	3	86	U	C2-N3-C4	-5.00	124.00	127.00
36	5	2627	C	N3-C4-C5	-5.00	119.90	121.90
36	5	3181	C	N1-C2-N3	5.00	122.70	119.20
36	1	1848	G	N1-C2-N2	-5.00	111.70	116.20
1	6	1725	U	C2-N1-C1'	5.00	123.70	117.70
36	5	1395	G	C6-C5-N7	-5.00	127.40	130.40
36	5	1695	U	N3-C2-O2	-5.00	118.70	122.20
36	5	2152	A	O4'-C1'-N9	5.00	112.20	108.20
36	5	2572	C	N1-C2-O2	5.00	121.90	118.90
36	5	2978	U	O4'-C1'-N1	5.00	112.20	108.20
36	5	3039	C	C6-N1-C2	-5.00	118.30	120.30
1	2	1595	U	O4'-C1'-N1	5.00	112.20	108.20
36	1	546	C	C6-N1-C2	-5.00	118.30	120.30
36	1	609	G	N3-C4-N9	5.00	129.00	126.00
36	1	1126	G	N9-C4-C5	-5.00	103.40	105.40
36	1	2647	A	O5'-P-OP2	5.00	116.70	110.70
1	6	620	A	OP2-P-O3'	5.00	116.20	105.20
1	6	913	G	N1-C6-O6	5.00	122.90	119.90
36	5	1009	A	O5'-P-OP1	5.00	116.70	110.70
36	5	3234	A	O4'-C1'-N9	5.00	112.20	108.20
36	5	3309	G	N3-C4-N9	5.00	129.00	126.00

There are no chirality outliers.

All (37) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
16	C4	123	SER	Peptide
18	C6	113	ASP	Peptide
19	C7	84	TYR	Peptide
27	D5	54	VAL	Peptide
27	D5	94	LYS	Peptide
33	E1	146	SER	Peptide
39	L2	19	HIS	Peptide
40	L3	248	LYS	Peptide
42	L5	257	GLU	Peptide
43	L6	17	ALA	Peptide
45	L8	30	THR	Peptide
47	M0	188	GLY	Peptide
48	M1	8	PRO	Peptide
50	M4	112	LEU	Peptide
52	M6	110	PRO	Peptide
65	N9	19	ASN	Peptide
67	O1	5	LYS	Peptide
3	S1	177	GLN	Peptide
9	S7	131	PHE	Peptide
10	S8	8	ARG	Peptide
11	S9	137	GLY	Peptide
18	c6	40	GLU	Peptide
20	c8	144	ARG	Peptide
22	d0	70	THR	Peptide
26	d4	29	HIS	Peptide
80	e0	61	SER	Peptide
39	l2	215	ASN	Peptide
40	l3	234	GLY	Peptide
44	l7	226	GLY	Peptide
48	m1	8	PRO	Peptide
56	n0	133	ALA	Peptide
56	n0	152	LEU	Peptide
64	n8	66	ALA	Peptide
69	o3	105	SER	Peptide
7	s5	44	ASN	Peptide
7	s5	99	MET	Peptide
9	s7	30	SER	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	2	37283	0	18758	1135	1
1	6	38238	0	19241	1143	0
2	S0	1577	0	1567	185	0
2	s0	1583	0	1578	0	0
3	S1	1709	0	1784	191	0
3	s1	1722	0	1793	0	0
4	S2	1635	0	1723	161	0
4	s2	1635	0	1723	0	0
5	S3	1734	0	1817	155	0
5	s3	1734	0	1817	0	0
6	S4	2068	0	2154	199	0
6	s4	2068	0	2154	0	0
7	S5	1609	0	1675	198	0
7	s5	1609	0	1675	0	0
8	S6	1799	0	1879	192	0
8	s6	1755	0	1846	0	0
9	S7	1481	0	1572	154	0
9	s7	1491	0	1578	0	0
10	S8	1489	0	1525	137	0
10	s8	1489	0	1525	0	0
11	S9	1494	0	1573	160	0
11	s9	1494	0	1573	0	0
12	C0	773	0	729	74	0
12	c0	762	0	699	0	0
13	C1	1214	0	1259	116	0
13	c1	1168	0	1231	0	0
14	C2	892	0	891	76	0
14	c2	892	0	891	0	0
15	C3	1192	0	1255	133	0
15	c3	1192	0	1255	0	0
16	C4	891	0	883	100	0
16	c4	949	0	985	0	0
17	C5	977	0	1002	115	0
17	c5	1039	0	1050	0	0
18	C6	1105	0	1166	137	0
18	c6	1111	0	1171	0	0
19	C7	926	0	930	107	0
19	c7	906	0	909	0	0
20	C8	1192	0	1222	147	0
20	c8	1192	0	1222	0	0
21	C9	1112	0	1124	140	0
21	c9	1112	0	1124	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
22	D0	855	0	917	105	0
22	d0	882	0	939	0	0
23	D1	684	0	672	88	0
23	d1	684	0	672	0	0
24	D2	1021	0	1060	111	0
24	d2	1021	0	1060	0	0
25	D3	1121	0	1196	127	0
25	d3	1121	0	1196	0	0
26	D4	1073	0	1132	103	0
26	d4	1073	0	1132	0	0
27	D5	563	0	603	68	0
27	d5	558	0	598	0	0
28	D6	769	0	814	123	0
28	d6	769	0	814	0	0
29	D7	610	0	631	59	0
29	d7	610	0	632	0	0
30	D8	497	0	535	55	0
30	d8	497	0	535	0	0
31	D9	442	0	428	42	0
31	d9	442	0	428	0	0
32	E0	475	0	525	40	0
33	E1	566	0	602	65	0
33	e1	608	0	656	0	0
34	SR	2441	0	2397	221	0
34	sR	2442	0	2392	0	0
35	SM	1104	0	996	87	0
35	sM	679	0	603	0	0
36	1	67355	0	33845	1713	0
36	5	67376	0	33855	1695	1
37	3	2579	0	1304	81	0
37	7	2579	0	1303	73	0
38	4	3353	0	1695	85	0
38	8	3353	0	1695	98	0
39	L2	1914	0	1981	203	0
39	l2	1912	0	1976	0	0
40	L3	3075	0	3142	284	0
40	l3	3075	0	3142	0	0
41	L4	2748	0	2859	285	0
41	l4	2748	0	2859	0	0
42	L5	2375	0	2325	238	0
42	l5	2359	0	2311	0	0
43	L6	1239	0	1326	110	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
43	l6	1248	0	1339	0	0
44	L7	1784	0	1862	193	0
44	l7	1791	0	1869	0	0
45	L8	1804	0	1877	174	0
45	l8	1763	0	1819	0	0
46	L9	1518	0	1587	171	0
46	l9	1518	0	1587	0	0
47	M0	1705	0	1736	164	0
47	m0	1722	0	1755	0	0
48	M1	1353	0	1383	140	0
48	m1	1353	0	1383	0	0
49	M3	1543	0	1608	163	0
49	m3	1548	0	1613	0	0
50	M4	1053	0	1149	114	0
50	m4	1059	0	1154	0	0
51	M5	1720	0	1779	203	0
51	m5	1720	0	1779	0	0
52	M6	1555	0	1659	181	0
52	m6	1555	0	1659	0	0
53	M7	1420	0	1437	139	0
53	m7	1227	0	1236	0	0
54	M8	1441	0	1543	131	0
54	m8	1441	0	1543	0	0
55	M9	1521	0	1616	150	0
55	m9	1521	0	1617	0	0
56	N0	1445	0	1487	161	0
56	n0	1445	0	1487	0	0
57	N1	1276	0	1323	142	0
57	n1	1276	0	1323	0	0
58	N2	796	0	812	61	0
58	n2	778	0	791	0	0
59	N3	1003	0	1048	107	0
59	n3	1003	0	1048	0	0
60	N4	699	0	640	44	0
60	n4	1038	0	1071	0	0
61	N5	964	0	1025	84	0
61	n5	959	0	1023	0	0
62	N6	993	0	1081	110	0
62	n6	993	0	1081	0	0
63	N7	1092	0	1155	100	0
63	n7	1092	0	1155	0	0
64	N8	1173	0	1215	141	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
64	n8	1173	0	1215	0	0
65	N9	462	0	491	47	0
65	n9	462	0	491	0	0
66	O0	743	0	797	82	0
66	o0	767	0	816	0	0
67	O1	876	0	912	79	0
67	o1	883	0	918	0	0
68	O2	1020	0	1090	95	0
68	o2	1020	0	1090	0	0
69	O3	850	0	880	81	0
69	o3	850	0	880	0	0
70	O4	880	0	945	113	0
70	o4	880	0	945	0	0
71	O5	969	0	1078	87	0
71	o5	965	0	1067	0	0
72	O6	771	0	849	84	0
72	o6	770	0	846	0	0
73	O7	681	0	683	76	0
73	o7	681	0	683	0	0
74	O8	612	0	682	63	0
74	o8	608	0	671	0	0
75	O9	436	0	475	44	0
75	o9	436	0	475	0	0
76	Q0	417	0	457	47	0
76	q0	417	0	456	0	0
77	Q1	233	0	284	30	0
77	q1	233	0	284	0	0
78	Q2	847	0	916	83	0
78	q2	847	0	914	0	0
79	Q3	694	0	734	80	0
79	q3	694	0	734	0	0
80	e0	491	0	542	0	0
81	p0	1076	0	1040	0	0
82	m2	750	0	179	0	0
83	p1	235	0	50	0	0
84	p2	230	0	51	0	0
85	1	462	0	0	0	0
85	2	122	0	0	0	0
85	3	15	0	0	0	0
85	4	23	0	0	0	0
85	5	495	0	0	0	0
85	6	147	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
85	7	17	0	0	0	0
85	8	14	0	0	0	0
85	C1	1	0	0	0	0
85	D0	1	0	0	0	0
85	D3	1	0	0	0	0
85	D4	1	0	0	0	0
85	L2	2	0	0	0	0
85	L3	4	0	0	0	0
85	L4	3	0	0	0	0
85	L7	2	0	0	0	0
85	L8	1	0	0	0	0
85	M0	3	0	0	0	0
85	M1	1	0	0	0	0
85	M3	2	0	0	0	0
85	M5	2	0	0	0	0
85	M6	1	0	0	0	0
85	M7	5	0	0	0	0
85	M9	1	0	0	0	0
85	N0	1	0	0	0	0
85	N3	3	0	0	0	0
85	N5	1	0	0	0	0
85	N6	2	0	0	0	0
85	N8	5	0	0	0	0
85	O1	1	0	0	0	0
85	O2	2	0	0	0	0
85	O5	1	0	0	0	0
85	O7	2	0	0	0	0
85	Q2	1	0	0	0	0
85	S4	1	0	0	0	0
85	S8	1	0	0	0	0
85	SM	1	0	0	0	0
85	c1	1	0	0	0	0
85	c4	1	0	0	0	0
85	c7	1	0	0	0	0
85	c8	1	0	0	0	0
85	c9	1	0	0	0	0
85	d0	1	0	0	0	0
85	d3	3	0	0	0	0
85	d6	1	0	0	0	0
85	l2	3	0	0	0	0
85	l3	3	0	0	0	0
85	l4	1	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
85	l5	3	0	0	0	0
85	l7	1	0	0	0	0
85	l9	1	0	0	0	0
85	m0	1	0	0	0	0
85	m1	1	0	0	0	0
85	m4	1	0	0	0	0
85	m5	5	0	0	0	0
85	m6	2	0	0	0	0
85	m7	6	0	0	0	0
85	n0	1	0	0	0	0
85	n3	1	0	0	0	0
85	n6	2	0	0	0	0
85	n8	3	0	0	0	0
85	n9	2	0	0	0	0
85	o3	2	0	0	0	0
85	o4	3	0	0	0	0
85	o7	1	0	0	0	0
85	q0	2	0	0	0	0
85	q1	1	0	0	0	0
85	s1	1	0	0	0	0
85	s8	1	0	0	0	0
85	sM	1	0	0	0	0
86	1	2478	0	0	204	0
86	2	1106	0	0	106	0
86	3	70	0	0	4	0
86	4	105	0	0	6	0
86	5	2492	0	0	217	0
86	6	1113	0	0	108	0
86	7	84	0	0	12	0
86	8	105	0	0	10	0
86	C3	7	0	0	1	0
86	C5	7	0	0	3	0
86	C8	7	0	0	0	0
86	D3	7	0	0	2	0
86	D9	7	0	0	0	0
86	L3	14	0	0	2	0
86	L4	7	0	0	2	0
86	M0	7	0	0	1	0
86	M5	7	0	0	1	0
86	M7	14	0	0	2	0
86	M9	7	0	0	1	0
86	N9	7	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
86	O1	7	0	0	7	0
86	O3	7	0	0	1	0
86	O7	14	0	0	7	0
86	Q2	7	0	0	2	0
86	S8	7	0	0	1	0
86	SR	7	0	0	0	0
86	c3	7	0	0	0	0
86	c5	7	0	0	0	0
86	c8	7	0	0	0	0
86	d4	7	0	0	0	0
86	d9	7	0	0	0	0
86	l3	14	0	0	0	0
86	l4	14	0	0	0	0
86	l5	21	0	0	0	0
86	l9	7	0	0	0	0
86	m0	14	0	0	0	0
86	m1	7	0	0	0	0
86	m4	7	0	0	0	0
86	m5	7	0	0	0	0
86	m6	7	0	0	0	0
86	m7	7	0	0	0	0
86	n1	7	0	0	0	0
86	n3	7	0	0	0	0
86	n9	7	0	0	0	0
86	o3	7	0	0	0	0
86	o7	7	0	0	0	0
86	q2	7	0	0	0	0
86	s1	14	0	0	0	0
86	s4	7	0	0	0	0
86	s8	7	0	0	0	0
86	s9	7	0	0	0	0
86	sR	7	0	0	0	0
87	D6	1	0	0	0	0
87	D7	1	0	0	0	0
87	D9	1	0	0	0	0
87	E1	1	0	0	0	0
87	O7	1	0	0	0	0
87	Q0	1	0	0	1	0
87	Q2	1	0	0	2	0
87	Q3	1	0	0	0	0
87	d6	1	0	0	0	0
87	d7	1	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
87	d9	1	0	0	0	0
87	e1	1	0	0	0	0
87	o7	1	0	0	0	0
87	q0	1	0	0	0	0
87	q2	1	0	0	0	0
87	q3	1	0	0	0	0
88	1	30	0	24	2	0
88	5	30	0	25	2	0
All	All	411214	0	297334	13265	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 19.

All (13265) 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)
78:Q2:17:CYS:SG	78:Q2:17:CYS:CB	2.03	1.47
78:Q2:17:CYS:CB	87:Q2:501:ZN:ZN	0.98	1.42
78:Q2:17:CYS:SG	87:Q2:501:ZN:ZN	1.30	1.18
36:5:2273:G:O6	86:5:4193:OHX:N5	1.90	1.05
40:L3:41:VAL:HA	40:L3:185:GLY:HA3	1.53	1.04
8:S6:98:ARG:NH2	8:S6:101:ILE:O	2.22	0.99
1:6:1636:C:H4'	1:6:1637:C:H5'	1.45	0.98
2:S0:41:ARG:HE	2:S0:45:VAL:HB	1.26	0.98
50:M4:55:ARG:NH2	50:M4:76:ALA:O	2.99	0.98
70:O4:80:ARG:HH11	70:O4:88:ARG:HH22	1.04	0.97
67:O1:44:MET:HB2	67:O1:46:THR:HG22	1.46	0.97
36:1:2836:C:H5	36:1:2852:C:H42	1.13	0.97
52:M6:160:ARG:NH2	36:5:3182:G:OP1	280.03	0.97
18:C6:73:GLY:H	18:C6:76:SER:HB2	1.29	0.97
36:1:2206:G:H1	36:1:2237:C:H42	1.11	0.97
26:D4:112:LYS:HE2	26:D4:116:LYS:HD2	1.46	0.96
70:O4:74:ARG:NH2	36:5:1639:C:OP2	200.65	0.96
20:C8:145:ARG:HB3	35:SM:68:ARG:HH12	4.37	0.96
69:O3:18:ARG:HD3	36:5:1178:G:H5''	237.65	0.95
36:1:3166:C:H42	36:1:3284:G:H1	1.13	0.95
46:L9:105:GLU:HG3	46:L9:109:ALA:H	1.29	0.94
36:1:3182:G:OP1	52:M6:160:ARG:NH2	2.00	0.94
36:1:439:C:H3'	36:1:440:A:H8	1.31	0.94
61:N5:115:ARG:HD3	61:N5:121:LYS:HE3	2.10	0.94
4:S2:170:ILE:HB	4:S2:197:TYR:HB2	1.48	0.94

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:343:U:OP2	86:5:3917:OHX:N3	2.01	0.94
39:L2:193:ARG:NH2	36:5:2181:C:OP1	197.94	0.94
1:6:1370:U:H4'	1:6:1371:A:H4'	1.49	0.94
8:S6:120:GLU:HG3	8:S6:125:THR:HB	1.49	0.94
1:6:1595:U:H3	1:6:1600:A:H2	1.14	0.94
43:L6:31:ARG:HH11	69:O3:107:ILE:HG22	5.23	0.94
7:S5:94:THR:HG22	7:S5:114:ILE:HG13	1.91	0.94
40:L3:227:GLU:HG3	40:L3:270:ARG:HE	3.74	0.94
36:5:3274:A:H3'	36:5:3275:U:H5''	1.45	0.93
78:Q2:17:CYS:SG	78:Q2:77:CYS:HB3	2.08	0.93
13:C1:99:ARG:HB3	25:D3:12:ALA:HB2	3.34	0.93
36:1:2736:A:OP1	57:N1:92:ARG:NH1	2.01	0.93
1:6:1280:C:H2'	1:6:1281:G:H8	1.33	0.93
1:6:1688:U:H3	1:6:1713:G:H1	1.05	0.93
10:S8:162:ALA:HA	36:1:3353:G:H5'	1.50	0.93
36:5:2371:G:O6	86:5:3902:OHX:N6	2.02	0.93
1:2:140:A:N6	1:2:281:G:OP1	2.01	0.92
15:C3:101:HIS:O	15:C3:105:ASN:ND2	2.02	0.92
34:SR:38:ARG:HG2	34:SR:67:ILE:HG23	1.51	0.92
36:1:1240:A:H61	36:1:1244:A:H5''	1.35	0.92
1:2:991:G:OP2	86:2:2131:OHX:N1	2.02	0.92
1:2:158:U:O2'	1:2:160:C:OP2	1.89	0.91
62:N6:52:ARG:O	62:N6:54:ASP:N	2.02	0.91
39:L2:225:ILE:HG21	39:L2:234:LYS:HA	1.52	0.91
41:L4:329:PRO:O	41:L4:331:ALA:N	3.53	0.91
1:2:1595:U:H3	1:2:1600:A:H2	1.17	0.90
36:1:439:C:H3'	36:1:440:A:C8	2.06	0.90
28:D6:79:ILE:HA	28:D6:84:VAL:HG11	1.51	0.90
37:7:86:U:O2	86:7:221:OHX:N4	2.06	0.89
47:M0:170:LYS:HA	47:M0:177:ASP:HA	1.53	0.89
1:2:237:C:H5''	1:2:238:U:H5'	1.52	0.89
4:S2:164:SER:HB3	1:6:1086:A:H5'	371.74	0.89
36:1:964:G:HO2'	64:N8:41:HIS:HE2	1.15	0.89
73:O7:21:ARG:NH2	73:O7:41:ALA:O	2.06	0.89
1:6:57:G:O6	1:6:90:C:N4	2.05	0.89
51:M5:68:ARG:HH11	51:M5:68:ARG:HG2	1.38	0.89
42:L5:68:THR:HG22	42:L5:70:THR:H	1.36	0.89
78:Q2:38:GLN:HE21	78:Q2:38:GLN:HA	1.79	0.88
2:S0:39:ASN:HD22	19:C7:105:GLN:HG2	5.06	0.88
6:S4:79:ASP:HB3	6:S4:82:TYR:HB2	1.53	0.88
1:6:1011:G:OP2	86:6:2120:OHX:N3	2.06	0.88

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:73:C:N3	49:M3:59:ARG:NH1	2.21	0.88
86:1:3865:OHX:N5	38:4:2:A:OP2	2.05	0.88
1:6:230:C:N3	1:6:235:G:N2	2.21	0.88
21:C9:38:LYS:NZ	21:C9:43:ASN:O	2.07	0.88
43:L6:93:VAL:O	43:L6:95:GLY:N	2.07	0.88
36:1:1514:G:O2'	75:O9:45:ARG:NH2	2.06	0.88
49:M3:166:ALA:HB1	64:N8:147:LEU:HD21	4.03	0.88
36:1:655:C:H2'	36:1:656:A:H8	1.38	0.88
20:C8:49:LYS:NZ	20:C8:79:TYR:O	2.07	0.88
78:Q2:17:CYS:SG	78:Q2:77:CYS:CB	2.62	0.87
36:1:1639:C:OP2	70:O4:74:ARG:NH2	2.08	0.87
1:6:754:A:N6	1:6:793:A:N7	2.22	0.87
34:SR:149:ASP:HB2	34:SR:175:ASP:HB3	2.42	0.87
16:C4:107:ARG:NH1	28:D6:52:ASP:OD1	3.50	0.87
1:2:823:G:H2'	1:2:824:G:H8	1.37	0.87
71:O5:64:GLU:HA	71:O5:67:ARG:HB3	1.54	0.87
20:C8:145:ARG:HB2	35:SM:68:ARG:HH22	1.40	0.87
36:1:1268:G:N2	36:1:1269:U:O4	2.08	0.87
24:D2:2:THR:N	1:6:1034:C:HO2'	338.47	0.87
63:N7:83:THR:HG23	63:N7:85:TYR:H	1.39	0.87
1:2:1339:C:O2'	1:2:1341:A:N7	2.07	0.86
51:M5:49:ARG:NH2	36:5:115:A:OP1	101.56	0.86
43:L6:26:ARG:NH1	36:5:503:C:OP1	254.58	0.86
17:C5:130:ARG:NH2	35:SM:65:THR:O	2.71	0.86
36:1:824:C:H5''	39:L2:21:ARG:HD3	1.56	0.86
36:5:2311:G:OP2	86:5:4193:OHX:N1	2.09	0.86
6:S4:104:ASP:HB2	6:S4:108:ARG:H	1.39	0.86
6:S4:49:ARG:NH1	1:6:448:C:OP2	380.72	0.86
6:S4:199:GLU:HB2	6:S4:207:LEU:HB2	1.55	0.86
42:L5:265:TYR:OH	37:7:121:U:OP2	311.67	0.86
34:SR:153:GLN:NE2	34:SR:202:LEU:O	2.09	0.86
21:C9:89:ARG:NH2	1:6:1562:G:OP1	377.57	0.86
15:C3:103:GLU:HA	15:C3:106:ARG:HH22	1.41	0.85
29:D7:56:CYS:SG	29:D7:57:GLU:N	2.62	0.85
41:L4:150:LEU:HD13	41:L4:249:ILE:HG23	2.22	0.85
41:L4:20:LEU:HD11	41:L4:252:GLU:HG3	1.58	0.85
36:1:3376:A:OP2	86:1:3899:OHX:N5	2.08	0.85
13:C1:92:HIS:CG	13:C1:93:TYR:H	1.94	0.85
48:M1:94:ARG:O	48:M1:96:PHE:N	3.30	0.85
1:2:1228:G:H1	14:C2:67:THR:HB	1.40	0.85
16:C4:85:ALA:H	16:C4:119:THR:HG22	1.39	0.85

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:D6:42:ARG:HH21	28:D6:42:ARG:HB2	4.05	0.85
1:6:452:A:OP2	86:6:2061:OHX:N1	2.08	0.85
1:6:895:G:H1	1:6:917:U:H3	1.24	0.85
56:N0:91:TYR:O	56:N0:137:ARG:NH1	2.09	0.85
36:1:1238:C:N4	36:1:1245:A:OP2	2.10	0.85
36:5:1077:U:N3	36:5:1082:U:O4	2.10	0.85
36:5:3165:A:H61	36:5:3285:C:H42	1.19	0.85
34:SR:35:SER:HG	34:SR:45:TRP:HE1	1.25	0.85
36:5:1009:A:N6	36:5:1041:U:O4	2.09	0.85
11:S9:109:LEU:HB2	11:S9:146:PHE:HB3	1.56	0.85
1:2:1014:G:OP1	86:2:2024:OHX:N3	2.11	0.84
42:L5:276:LYS:HB3	37:7:61:G:H5''	326.99	0.84
1:2:127:G:N7	8:S6:202:ARG:NH2	2.25	0.84
5:S3:208:ILE:HD12	19:C7:16:LEU:HD21	1.59	0.84
3:S1:61:LEU:HG	3:S1:64:ARG:HH21	1.42	0.84
33:E1:134:ASN:H	1:6:1251:U:H4'	442.07	0.84
38:8:79:A:H3'	38:8:80:A:C8	2.12	0.84
47:M0:156:ARG:O	47:M0:158:LYS:N	3.11	0.84
40:L3:150:ARG:HG2	40:L3:150:ARG:HH11	2.33	0.84
36:1:1567:U:O2	36:1:1571:A:N6	2.09	0.84
1:6:991:G:OP2	86:6:2171:OHX:N2	2.11	0.84
54:M8:34:THR:HG22	54:M8:49:LEU:HD21	3.92	0.84
6:S4:117:GLU:O	6:S4:119:ALA:N	3.24	0.84
57:N1:28:SER:OG	37:7:9:C:OP1	266.80	0.83
1:2:1202:A:OP1	86:2:2110:OHX:N1	2.11	0.83
45:L8:108:ARG:HA	45:L8:111:LYS:HD2	3.40	0.83
1:2:823:G:H2'	1:2:824:G:C8	2.12	0.83
36:1:912:G:OP2	39:L2:9:ARG:NH1	2.11	0.83
74:O8:46:ARG:NH2	36:5:1613:A:OP2	131.51	0.83
58:N2:99:LYS:HB2	58:N2:102:GLU:HB2	1.59	0.83
1:2:933:A:OP1	28:D6:70:LYS:NZ	2.11	0.83
42:L5:106:ALA:HB2	42:L5:166:ALA:HA	1.58	0.83
27:D5:65:LEU:HB3	27:D5:71:ILE:HD12	1.60	0.83
36:1:1824:U:H4'	74:O8:17:ARG:HH12	1.43	0.83
34:SR:169:ILE:HG12	34:SR:183:LEU:HD21	4.02	0.83
17:C5:43:ARG:NH1	1:6:1553:G:N7	400.98	0.83
32:E0:50:VAL:HA	32:E0:54:ARG:HA	3.79	0.83
72:O6:4:LYS:O	72:O6:16:LYS:NZ	2.59	0.83
54:M8:66:ARG:NH2	36:5:744:A:OP1	166.79	0.83
14:C2:75:VAL:HG11	14:C2:120:VAL:HG11	3.17	0.83
36:1:2534:G:H1	36:1:2545:C:H42	1.27	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:7:77:G:N2	37:7:102:A:OP2	2.12	0.82
41:L4:226:GLU:OE1	41:L4:237:GLN:NE2	2.16	0.82
40:L3:169:THR:HG23	40:L3:171:LEU:H	3.00	0.82
45:L8:126:SER:O	36:5:120:G:N2	94.41	0.82
54:M8:180:ARG:HH11	54:M8:185:LYS:HB3	1.43	0.82
41:L4:299:ILE:HG23	54:M8:39:ARG:HB3	3.63	0.82
1:2:789:A:O2'	6:S4:106:LYS:NZ	2.13	0.82
6:S4:45:ILE:HB	6:S4:80:THR:HG22	2.60	0.82
36:1:2656:A:H4'	78:Q2:98:LYS:HD2	1.60	0.82
78:Q2:71:ARG:HE	78:Q2:80:ARG:HD3	4.06	0.82
34:SR:102:ARG:NH2	1:6:1341:A:O2'	459.18	0.82
36:1:1591:G:O2'	36:1:1799:A:N1	2.12	0.82
38:4:70:G:O6	86:O7:104:OHX:N4	2.13	0.82
41:L4:60:THR:HG22	41:L4:62:ALA:H	3.22	0.82
45:L8:148:ALA:O	45:L8:149:LYS:NZ	5.67	0.82
1:6:1041:G:OP1	86:6:2175:OHX:N4	2.13	0.82
44:L7:123:THR:HA	44:L7:126:LEU:HD12	1.59	0.82
1:2:79:C:H1'	8:S6:174:LYS:HD3	1.60	0.82
36:1:1095:U:H4'	36:1:1096:U:H5'	1.61	0.82
20:C8:91:ASP:HB3	20:C8:95:GLY:H	1.97	0.82
61:N5:47:ALA:HB3	71:O5:77:PRO:HG3	1.62	0.82
74:O8:58:ASP:HB3	74:O8:61:LYS:HG3	4.21	0.82
36:5:3227:A:H2'	36:5:3228:C:H5'	1.61	0.82
46:L9:28:VAL:HG13	46:L9:33:THR:HB	1.87	0.82
61:N5:115:ARG:HH11	61:N5:115:ARG:HG3	1.87	0.82
1:6:1307:U:O4	1:6:1318:G:N2	2.12	0.82
52:M6:33:ILE:HG22	52:M6:102:LEU:HD13	1.61	0.82
59:N3:30:GLY:HA3	59:N3:66:LYS:HD2	1.62	0.82
69:O3:21:ARG:HG3	69:O3:21:ARG:HH11	1.44	0.82
36:1:1024:G:N7	86:1:4162:OHX:N6	2.28	0.81
26:D4:29:HIS:HB3	26:D4:32:ARG:HB2	6.12	0.81
1:6:1010:C:OP2	86:6:2171:OHX:N3	2.13	0.81
12:C0:32:HIS:CD2	12:C0:33:GLU:H	4.09	0.81
36:5:1555:U:O4	36:5:1557:A:N6	2.13	0.81
26:D4:121:THR:OG1	1:6:149:C:OP1	336.07	0.81
31:D9:19:ARG:NH2	1:6:1597:A:OP1	408.59	0.81
59:N3:87:ARG:HH12	59:N3:137:VAL:HG11	1.95	0.81
39:L2:128:ARG:NH1	36:5:2177:G:OP2	198.65	0.81
36:1:971:G:OP1	54:M8:8:LYS:NZ	2.11	0.81
50:M4:133:LYS:NZ	36:5:3227:A:O2'	302.37	0.81
8:S6:190:GLN:NE2	1:6:265:A:N7	333.84	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:477:A:H2'	1:2:478:A:H8	1.45	0.81
1:2:657:U:O2	1:2:677:G:N2	2.13	0.81
64:N8:21:ARG:NH2	36:5:640:U:OP1	182.12	0.81
1:2:885:G:H21	16:C4:123:SER:HB2	1.43	0.81
74:O8:24:THR:HG22	74:O8:76:ASN:HB3	1.63	0.81
6:S4:52:LEU:HB3	6:S4:54:TYR:HD2	1.43	0.81
36:1:1466:G:O6	86:1:3871:OHX:N4	2.13	0.81
36:1:1560:G:N2	36:1:1579:C:O2	2.12	0.81
36:1:3243:A:H4'	40:L3:95:THR:HG22	1.62	0.81
39:L2:30:ARG:NH2	39:L2:33:ASP:OD2	2.13	0.81
36:1:3375:A:O2'	36:1:3378:C:OP2	1.99	0.81
17:C5:43:ARG:NH2	1:6:1552:U:OP2	403.47	0.81
35:SM:23:LYS:HD2	35:SM:23:LYS:H	1.44	0.81
39:L2:29:LEU:O	39:L2:123:ARG:NH2	2.90	0.81
55:M9:23:TRP:CH2	55:M9:25:ASP:HB3	2.15	0.81
1:2:9:U:O4	86:2:2155:OHX:N6	2.14	0.81
1:6:1151:A:O2'	1:6:1766:A:N7	2.12	0.81
4:S2:237:VAL:HB	4:S2:242:ILE:HD11	2.30	0.81
3:S1:34:ALA:HB3	3:S1:41:ARG:HA	1.61	0.80
45:L8:109:LEU:O	45:L8:113:ALA:N	2.13	0.80
10:S8:103:GLN:HG2	10:S8:164:ARG:HG2	1.63	0.80
36:1:329:U:OP2	86:1:4038:OHX:N4	2.14	0.80
3:S1:29:TRP:HE1	3:S1:47:LEU:HG	1.44	0.80
56:N0:89:ASN:HD21	57:N1:156:TYR:HB3	1.47	0.80
1:2:1010:C:OP2	86:2:2131:OHX:N6	2.15	0.80
21:C9:57:ARG:HH21	21:C9:101:ASN:HD21	6.44	0.80
40:L3:252:ILE:HG13	40:L3:266:ARG:HH21	1.43	0.80
37:3:49:G:N7	42:L5:58:LYS:HG3	1.97	0.80
68:O2:105:ARG:HE	68:O2:124:GLY:HA3	1.45	0.80
17:C5:37:ALA:O	17:C5:42:ARG:NH1	4.28	0.80
64:N8:88:ASP:HA	64:N8:91:LEU:HB2	2.74	0.80
2:S0:183:ARG:HG3	2:S0:188:LEU:HD12	4.39	0.80
36:5:2836:C:H5	36:5:2852:C:H42	1.27	0.80
36:5:2236:G:OP1	86:5:4244:OHX:N3	2.15	0.80
36:5:742:G:N7	86:5:3996:OHX:N4	2.30	0.80
1:6:1160:A:H2'	1:6:1161:C:C6	2.17	0.80
25:D3:112:LYS:NZ	1:6:18:C:O3'	347.19	0.80
8:S6:135:PRO:HB2	8:S6:141:ILE:HG13	1.64	0.80
74:O8:42:LYS:NZ	36:5:1750:A:OP2	141.86	0.80
22:D0:58:LEU:HD12	22:D0:88:LYS:HD2	1.64	0.80
36:5:332:C:O2	38:8:31:G:N2	2.14	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:D2:15:ASN:HD21	24:D2:71:LYS:HG3	3.77	0.80
41:L4:82:THR:O	41:L4:84:ARG:N	2.14	0.80
53:M7:50:GLN:OE1	53:M7:56:ARG:NH2	2.11	0.80
6:S4:191:ARG:HH11	6:S4:245:LYS:HD3	1.47	0.80
1:2:1435:G:N7	12:C0:25:LYS:NZ	2.30	0.79
1:6:1294:G:O6	86:6:2068:OHX:N5	2.14	0.79
66:O0:16:LEU:HB2	66:O0:98:SER:HB2	3.60	0.79
69:O3:20:LYS:HG2	69:O3:21:ARG:HG3	4.33	0.79
3:S1:92:GLN:HG2	3:S1:97:LEU:HD21	6.73	0.79
1:2:1280:C:H2'	1:2:1281:G:H8	1.47	0.79
13:C1:72:THR:HG22	13:C1:124:THR:HG23	1.65	0.79
1:2:1498:G:H5''	21:C9:72:GLY:HA3	1.64	0.79
61:N5:132:ALA:HA	61:N5:135:ILE:HG22	3.76	0.79
5:S3:70:THR:HG23	5:S3:86:LEU:HB2	2.22	0.79
36:1:3166:C:N3	36:1:3284:G:N2	2.27	0.79
36:5:272:G:OP2	86:5:4068:OHX:N6	2.15	0.79
47:M0:66:GLU:OE1	47:M0:69:ARG:NH2	2.15	0.79
59:N3:74:MET:HG3	59:N3:102:ILE:HD13	1.64	0.79
11:S9:133:HIS:NE2	1:6:513:U:OP1	447.85	0.79
1:6:826:U:O4	86:6:2065:OHX:N3	2.15	0.79
33:E1:144:CYS:HB3	33:E1:147:VAL:HG13	1.64	0.79
46:L9:96:HIS:CD2	36:5:3024:A:H5''	339.63	0.79
49:M3:47:ALA:HB1	49:M3:48:PRO:HD2	1.65	0.79
52:M6:12:LYS:O	56:N0:167:ARG:NH2	2.15	0.79
34:SR:161:LYS:O	34:SR:161:LYS:HG2	1.82	0.79
36:1:356:C:OP2	86:1:4138:OHX:N1	2.15	0.79
50:M4:113:THR:HG22	50:M4:116:GLU:H	1.46	0.79
60:N4:54:LEU:H	60:N4:54:LEU:HD12	1.46	0.79
7:S5:68:ILE:HD12	7:S5:69:PHE:H	3.74	0.79
36:1:998:A:N6	36:1:1050:U:O4	2.14	0.79
59:N3:48:ARG:NH2	36:5:3043:C:OP2	251.46	0.79
51:M5:125:SER:HB3	36:5:2433:U:H1'	160.96	0.79
36:5:3035:A:OP2	86:5:4045:OHX:N5	2.16	0.79
1:6:1783:C:H2'	1:6:1784:C:H6	1.46	0.79
1:2:1471:A:OP1	7:S5:185:ARG:NH1	2.15	0.79
36:5:501:A:H2'	36:5:502:U:H6	1.46	0.79
1:6:982:U:OP1	86:6:2075:OHX:N2	2.15	0.79
50:M4:16:GLU:HB3	56:N0:149:LYS:HB3	4.02	0.79
36:1:2355:G:H4'	53:M7:139:TYR:CE2	2.17	0.79
39:L2:149:ARG:NH2	39:L2:252:THR:O	5.88	0.79
47:M0:86:HIS:HB3	47:M0:139:ARG:HG2	1.87	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:M5:31:ARG:NH1	51:M5:124:ASP:OD1	2.58	0.79
64:N8:21:ARG:NH1	36:5:1369:A:OP1	183.41	0.79
66:O0:57:GLU:OE1	66:O0:69:TYR:OH	3.29	0.79
34:SR:197:SER:HB3	34:SR:217:ASP:HB3	2.57	0.79
36:1:1473:G:OP2	55:M9:8:LYS:NZ	2.16	0.78
1:2:1794:A:OP1	86:2:2091:OHX:N4	2.16	0.78
11:S9:126:ARG:NH1	1:6:475:A:OP2	424.88	0.78
16:C4:50:ALA:O	16:C4:52:ARG:N	2.22	0.78
20:C8:82:PRO:HB2	20:C8:85:PHE:HB2	1.65	0.78
2:S0:185:ARG:HB2	23:D1:45:ALA:HB3	1.65	0.78
47:M0:171:TRP:O	47:M0:174:THR:OG1	1.99	0.78
36:1:255:A:H2'	36:1:256:G:H8	1.49	0.78
1:2:4:C:OP1	4:S2:200:SER:OG	2.01	0.78
41:L4:203:ARG:NH1	41:L4:226:GLU:OE2	2.16	0.78
46:L9:49:ASN:O	46:L9:51:GLN:N	2.16	0.78
47:M0:112:GLN:O	86:5:4238:OHX:N3	237.23	0.78
10:S8:76:THR:HG23	10:S8:108:PRO:HG2	2.73	0.78
34:SR:70:ASP:OD2	34:SR:155:ARG:NH2	2.14	0.78
42:L5:279:LYS:HE3	42:L5:282:ARG:HH11	1.46	0.78
52:M6:68:ARG:NH1	36:5:2988:C:OP1	217.68	0.78
36:5:2211:U:O4	86:5:3955:OHX:N4	2.17	0.78
15:C3:76:LYS:NZ	1:6:813:U:OP2	317.77	0.78
18:C6:112:TYR:O	18:C6:114:ARG:NH1	7.86	0.78
36:1:2732:G:OP2	86:1:4201:OHX:N2	2.17	0.78
36:5:1216:C:O2	36:5:1289:G:N2	2.16	0.78
10:S8:18:ARG:NH1	1:6:105:A:OP1	305.55	0.78
42:L5:34:LYS:HA	57:N1:27:LEU:HD21	1.65	0.78
1:2:1588:G:OP1	86:2:2116:OHX:N3	2.17	0.78
12:C0:77:ARG:HG3	12:C0:82:LEU:HD12	1.65	0.78
28:D6:79:ILE:HG12	28:D6:84:VAL:HG21	1.63	0.78
40:L3:221:THR:HG22	40:L3:273:HIS:H	3.55	0.78
41:L4:145:ILE:HD11	41:L4:148:ILE:HD12	1.63	0.78
48:M1:62:ASN:ND2	78:Q2:101:GLY:O	2.16	0.78
10:S8:39:GLY:HA2	10:S8:61:GLU:HB3	1.66	0.78
36:1:76:G:O2'	49:M3:100:ARG:NH1	2.17	0.78
9:S7:149:ILE:HG23	9:S7:180:GLN:HB3	1.63	0.78
11:S9:65:LYS:HA	11:S9:70:LEU:HD21	1.66	0.78
1:2:697:C:O2'	1:2:699:U:OP2	2.00	0.78
36:5:980:A:H2'	36:5:981:U:C2	2.19	0.78
47:M0:36:LEU:HD11	47:M0:87:LEU:HD12	4.85	0.78
7:S5:95:ASN:OD1	7:S5:107:LYS:NZ	3.28	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:347:G:N7	73:O7:55:ARG:NH2	2.32	0.78
35:SM:48:ARG:HD3	35:SM:51:ARG:HB2	1.66	0.78
34:SR:90:ARG:HH21	34:SR:102:ARG:HH21	3.83	0.78
20:C8:11:PHE:HD2	20:C8:59:GLY:HA3	1.49	0.77
30:D8:15:VAL:HA	30:D8:28:VAL:HG23	4.13	0.77
47:M0:175:ASN:OD1	47:M0:176:LEU:N	5.03	0.77
59:N3:87:ARG:HH22	59:N3:137:VAL:HG21	1.47	0.77
62:N6:61:GLY:O	62:N6:63:LYS:N	3.73	0.77
36:1:1243:G:N2	36:1:1244:A:N7	2.33	0.77
36:1:1565:G:N2	36:1:1574:C:O2	2.17	0.77
36:1:2763:U:H5'	54:M8:176:ARG:HG3	1.67	0.77
46:L9:9:GLN:HG2	46:L9:52:LEU:HD21	1.65	0.77
34:SR:249:ARG:NH1	34:SR:298:GLY:O	3.10	0.77
1:2:1297:G:N2	1:2:1300:A:OP2	2.16	0.77
1:6:1734:U:H2'	1:6:1735:U:H6	1.49	0.77
40:L3:360:ASP:OD1	40:L3:371:GLN:NE2	4.21	0.77
54:M8:43:PRO:HA	54:M8:46:LYS:HD2	1.64	0.77
57:N1:83:ARG:HD2	57:N1:85:LEU:HD21	3.00	0.77
7:S5:35:GLN:O	7:S5:37:GLN:N	2.56	0.77
9:S7:11:GLN:HG3	9:S7:13:PRO:HD2	1.67	0.77
36:1:2662:G:H2'	36:1:2663:G:C8	2.18	0.77
36:5:2255:A:H5'	36:5:2261:G:H22	1.50	0.77
22:D0:24:ILE:HG12	22:D0:116:VAL:HG13	1.67	0.77
48:M1:59:ILE:HG21	48:M1:65:ILE:HD11	1.66	0.77
62:N6:57:LEU:HD23	62:N6:67:GLU:HG2	3.18	0.77
76:Q0:106:ARG:HB2	76:Q0:106:ARG:HH11	4.40	0.77
1:2:1203:A:OP2	86:2:2110:OHX:N5	2.17	0.77
18:C6:109:PHE:O	18:C6:113:ASP:N	3.06	0.77
1:2:1325:A:OP2	19:C7:11:ARG:NH1	2.18	0.77
50:M4:55:ARG:HD3	56:N0:70:THR:HB	1.67	0.77
79:Q3:49:ARG:HD3	79:Q3:50:GLY:H	1.49	0.77
36:1:1077:U:N3	36:1:1082:U:O4	2.15	0.77
36:1:1878:G:OP1	86:1:3920:OHX:N4	2.17	0.77
36:1:2120:A:OP2	86:1:4003:OHX:N2	2.18	0.77
36:1:2233:A:OP2	86:1:4039:OHX:N5	2.16	0.77
1:2:197:A:H61	10:S8:138:ASN:HD22	1.31	0.77
36:5:2223:A:N6	36:5:2783:U:O2'	2.16	0.77
7:S5:166:ARG:NH1	7:S5:170:GLN:OE1	2.16	0.77
36:1:1015:U:O2'	36:1:1017:C:OP2	2.02	0.77
1:6:915:A:OP1	86:6:2070:OHX:N6	2.17	0.77
15:C3:47:PRO:HG3	15:C3:72:MET:HG3	5.54	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:C3:73:ARG:O	15:C3:77:SER:OG	2.02	0.77
4:S2:140:ARG:NH1	23:D1:1:MET:SD	2.58	0.77
2:S0:157:ASP:OD2	23:D1:60:ARG:NH2	2.17	0.77
41:L4:64:SER:HA	41:L4:75:PRO:HA	1.66	0.77
44:L7:93:ASN:OD1	44:L7:93:ASN:N	2.58	0.77
36:1:269:G:OP2	51:M5:44:ARG:NH2	2.18	0.77
4:S2:243:TYR:HB3	4:S2:246:GLU:HB2	1.65	0.77
36:5:651:G:O2'	36:5:1435:A:OP1	2.03	0.77
36:5:1497:C:H2'	36:5:1498:A:H8	1.47	0.77
66:O0:29:SER:HA	66:O0:32:LYS:HD3	1.65	0.77
7:S5:84:LYS:HG2	7:S5:92:ARG:HH12	1.48	0.77
36:1:2585:G:N7	45:L8:47:SER:OG	2.18	0.77
20:C8:83:ALA:O	20:C8:89:GLN:NE2	2.18	0.77
26:D4:64:PHE:O	26:D4:66:GLY:N	2.68	0.77
28:D6:95:ARG:NH1	1:6:1796:C:O2'	343.04	0.77
76:Q0:91:CYS:O	76:Q0:126:LYS:NZ	2.15	0.77
9:S7:98:ILE:HG12	9:S7:121:VAL:HG21	1.66	0.77
36:5:1540:U:OP1	86:5:4088:OHX:N2	2.17	0.76
1:6:709:C:O2	1:6:730:G:N2	2.17	0.76
46:L9:163:GLN:O	46:L9:166:ARG:HD3	1.84	0.76
47:M0:36:LEU:HD11	47:M0:69:ARG:HD3	1.67	0.76
3:S1:71:ALA:HB3	16:C4:114:ARG:HH12	2.86	0.76
24:D2:30:SER:HA	24:D2:34:ILE:HD12	1.65	0.76
50:M4:94:TRP:O	50:M4:97:SER:OG	2.56	0.76
40:L3:56:ILE:HD11	40:L3:359:ILE:HG12	1.67	0.76
41:L4:330:TYR:CZ	44:L7:49:ALA:HA	2.79	0.76
46:L9:124:ARG:HB3	46:L9:164:ILE:HD13	2.50	0.76
49:M3:124:ILE:HD11	49:M3:126:PHE:HE1	1.50	0.76
73:O7:87:SER:O	86:O7:104:OHX:N3	2.18	0.76
6:S4:151:ASP:HB3	6:S4:154:ILE:HG13	2.20	0.76
9:S7:44:LYS:NZ	9:S7:95:GLU:OE2	2.17	0.76
36:5:2537:U:O2'	36:5:2538:U:O4'	2.03	0.76
1:6:471:A:OP2	86:6:2102:OHX:N5	2.17	0.76
27:D5:60:VAL:HG22	27:D5:101:TYR:HB2	1.67	0.76
47:M0:174:THR:OG1	47:M0:175:ASN:N	4.36	0.76
36:1:31:C:OP2	51:M5:188:ARG:NH2	2.19	0.76
36:1:953:G:OP1	65:N9:15:LYS:NZ	2.15	0.76
11:S9:168:ARG:HH12	11:S9:171:ARG:HD3	4.52	0.76
36:1:2179:C:O3'	39:L2:174:ARG:NH2	2.18	0.76
37:3:17:A:OP1	42:L5:2:ALA:N	2.19	0.76
36:5:1233:G:N2	36:5:1255:C:O2	2.17	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:566:G:N7	86:5:4126:OHX:N5	2.34	0.76
33:E1:103:LEU:HD11	1:6:1252:C:H5'	454.69	0.76
42:L5:24:ARG:NH2	37:7:13:A:N3	292.17	0.76
7:S5:57:SER:O	7:S5:59:VAL:N	2.18	0.76
36:5:847:A:H2'	36:5:848:A:C8	2.21	0.76
27:D5:58:ARG:HB3	27:D5:103:ARG:HE	6.17	0.76
1:2:1508:U:O4	86:2:2031:OHX:N5	2.19	0.76
1:2:732:G:O6	86:2:2129:OHX:N5	2.19	0.76
36:5:2996:U:OP1	36:5:2996:U:H4'	1.82	0.76
61:N5:56:ARG:NH2	38:8:135:G:OP2	81.77	0.76
1:2:1429:G:H1'	22:D0:74:GLU:HG2	1.67	0.76
8:S6:115:LYS:NZ	60:N4:73:ARG:O	2.16	0.76
8:S6:153:VAL:O	8:S6:155:ASP:N	2.78	0.76
20:C8:36:LYS:NZ	1:6:1568:C:OP1	334.90	0.76
33:E1:121:CYS:HB3	33:E1:132:LEU:HD21	4.34	0.76
1:2:814:A:H5'	55:M9:170:ARG:HH22	1.51	0.76
36:1:269:G:H5''	51:M5:14:LYS:HE2	1.68	0.76
51:M5:45:PRO:O	51:M5:49:ARG:HB2	3.55	0.76
53:M7:120:ASN:OD1	53:M7:120:ASN:N	2.81	0.76
9:S7:66:SER:O	9:S7:68:ALA:N	2.91	0.76
36:1:2916:U:H1'	59:N3:44:SER:HB3	1.67	0.76
66:O0:30:THR:HG22	66:O0:91:SER:HB3	2.51	0.76
4:S2:101:VAL:HG22	4:S2:115:ILE:HG12	1.68	0.76
8:S6:78:THR:HA	8:S6:92:ARG:HG2	2.96	0.76
9:S7:177:THR:HG23	9:S7:179:LYS:H	4.76	0.76
34:SR:160:GLU:O	34:SR:162:ALA:N	2.18	0.76
1:2:1500:C:N4	1:2:1507:G:O6	2.18	0.75
1:2:229:U:H3	1:2:236:A:H61	1.33	0.75
1:2:312:A:H4'	1:2:313:U:H5''	1.68	0.75
36:5:1696:A:OP2	86:5:4180:OHX:N6	2.19	0.75
79:Q3:17:ARG:NH1	36:5:860:G:OP1	219.30	0.75
47:M0:38:LYS:HG2	47:M0:41:ALA:HB2	1.68	0.75
6:S4:123:LEU:HD23	6:S4:228:ILE:HG22	1.99	0.75
1:2:693:U:H5'	1:2:694:U:H5'	1.67	0.75
37:3:49:G:O6	42:L5:58:LYS:NZ	2.16	0.75
1:6:1239:U:O2	1:6:1246:C:N4	2.19	0.75
51:M5:96:ARG:HH11	51:M5:96:ARG:HG2	2.08	0.75
62:N6:45:ILE:HD12	62:N6:119:ILE:HG23	1.99	0.75
7:S5:206:SER:O	7:S5:212:LYS:NZ	2.46	0.75
36:5:2697:A:H2'	36:5:2698:G:C8	2.21	0.75
23:D1:17:CYS:HB2	23:D1:56:SER:HB3	2.27	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
62:N6:83:ASP:O	62:N6:84:LYS:HB2	1.86	0.75
5:S3:94:ARG:NH2	5:S3:125:TYR:OH	4.25	0.75
1:2:134:U:OP1	1:2:136:C:N4	2.20	0.75
1:2:687:G:H5'	24:D2:119:LYS:HG2	1.68	0.75
36:5:86:G:O2'	36:5:98:G:O6	2.03	0.75
17:C5:127:ARG:O	17:C5:129:GLY:N	3.97	0.75
26:D4:52:LYS:O	26:D4:54:ALA:N	2.14	0.75
41:L4:264:SER:O	41:L4:266:THR:N	2.19	0.75
70:O4:80:ARG:NH1	70:O4:88:ARG:HH22	1.81	0.75
3:S1:149:GLN:HE21	3:S1:151:LYS:HG2	3.21	0.75
36:1:1382:G:O6	36:1:1424:C:N4	2.18	0.75
13:C1:99:ARG:NH1	25:D3:7:ARG:O	2.53	0.75
4:S2:56:ILE:HG23	4:S2:61:LEU:HB2	1.66	0.75
7:S5:92:ARG:HG2	7:S5:92:ARG:HH11	2.99	0.75
9:S7:131:PHE:O	9:S7:133:THR:N	2.19	0.75
36:5:1383:G:O6	86:5:3929:OHX:N2	2.20	0.75
13:C1:95:PRO:O	13:C1:98:ASN:N	2.16	0.75
4:S2:60:SER:OG	23:D1:15:ARG:NH2	2.33	0.75
45:L8:101:THR:H	45:L8:104:GLU:HB2	1.50	0.75
48:M1:34:SER:HB2	48:M1:67:VAL:HG11	1.67	0.75
49:M3:79:GLU:OE1	49:M3:112:ASN:ND2	2.18	0.75
72:O6:26:ILE:O	72:O6:28:TYR:N	2.17	0.75
10:S8:36:THR:HG21	10:S8:173:PRO:HB2	1.69	0.75
34:SR:89:LEU:HB2	34:SR:103:PHE:HB2	1.66	0.75
36:1:266:A:OP1	51:M5:5:LYS:NZ	2.19	0.75
17:C5:115:TYR:OH	1:6:1556:A:OP1	387.62	0.75
1:6:1765:A:OP1	86:6:2126:OHX:N2	2.18	0.75
41:L4:47:ARG:NH1	41:L4:109:TRP:O	2.20	0.75
50:M4:132:LYS:HD3	36:5:3230:G:H4'	286.31	0.75
70:O4:11:ASN:OD1	70:O4:18:ASN:ND2	2.18	0.75
34:SR:52:GLN:HG2	34:SR:53:LYS:HG3	3.57	0.75
36:1:2588:U:OP1	45:L8:48:ARG:NH2	2.19	0.75
1:2:1542:G:N2	1:2:1569:A:OP2	2.18	0.75
36:5:166:C:N4	36:5:256:G:O6	2.18	0.75
1:6:833:U:O4	86:6:2100:OHX:N2	2.20	0.75
22:D0:72:ASN:HD22	22:D0:74:GLU:H	1.34	0.75
70:O4:31:ARG:NH2	36:5:1598:G:OP2	133.34	0.75
43:L6:51:ARG:NH1	50:M4:114:ASP:OD2	2.20	0.75
61:N5:105:VAL:HA	61:N5:130:TYR:CE2	2.22	0.75
36:1:2532:U:H3	36:1:2547:A:H61	1.31	0.74
36:1:655:C:H2'	36:1:656:A:C8	2.21	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:1585:U:H3	1:2:1611:A:H2	1.35	0.74
1:2:520:A:H2'	1:2:521:A:C8	2.22	0.74
36:5:2221:G:N2	36:5:2224:A:OP2	2.19	0.74
25:D3:64:PRO:O	86:6:2159:OHX:N2	361.05	0.74
49:M3:74:GLY:O	49:M3:101:ARG:NH1	2.20	0.74
59:N3:113:ALA:HA	59:N3:132:ASN:HB3	1.68	0.74
64:N8:34:MET:HB2	36:5:95:A:H5''	162.65	0.74
2:S0:76:ILE:HG12	2:S0:98:ILE:HG13	1.69	0.74
36:5:2568:C:N4	36:5:2574:G:O6	2.19	0.74
39:L2:14:SER:OG	39:L2:15:ILE:N	2.65	0.74
49:M3:157:ARG:HG2	49:M3:158:ALA:N	2.02	0.74
60:N4:9:SER:HA	60:N4:52:THR:HG22	1.68	0.74
7:S5:216:GLU:HG3	7:S5:219:ARG:HH21	4.32	0.74
36:5:2821:C:H42	36:5:2869:U:H3	1.35	0.74
1:6:990:C:OP2	86:6:2120:OHX:N2	2.20	0.74
3:S1:157:GLN:O	3:S1:159:SER:N	2.20	0.74
4:S2:38:VAL:HG13	4:S2:39:THR:HG23	1.67	0.74
6:S4:104:ASP:HB3	6:S4:106:LYS:H	2.10	0.74
6:S4:151:ASP:OD1	8:S6:215:ARG:NH1	4.06	0.74
34:SR:292:LEU:HD12	34:SR:301:LEU:HD11	1.69	0.74
36:1:1040:A:N3	47:M0:198:LYS:NZ	2.34	0.74
36:1:2121:G:O2'	36:1:2122:G:OP1	2.05	0.74
64:N8:95:SER:HA	64:N8:122:PRO:HG2	1.69	0.74
3:S1:148:ASN:ND2	1:6:1066:C:O2'	349.44	0.74
36:1:2818:U:H6	36:1:2818:U:H5'	1.51	0.74
36:1:968:G:H2'	36:1:969:C:C6	2.22	0.74
1:2:531:C:OP2	86:2:2069:OHX:N4	2.20	0.74
64:N8:14:HIS:O	64:N8:16:SER:N	2.21	0.74
36:1:1898:G:OP2	86:1:3924:OHX:N4	2.20	0.74
36:1:269:G:P	51:M5:44:ARG:HH22	2.10	0.74
1:2:176:C:OP1	86:2:2072:OHX:N3	2.20	0.74
1:6:484:C:H42	1:6:503:G:H22	1.33	0.74
45:L8:148:ALA:HA	45:L8:201:THR:HG22	2.21	0.74
67:O1:31:ARG:HB3	67:O1:31:ARG:HH11	1.53	0.74
2:S0:124:THR:HA	2:S0:146:LEU:HB2	1.70	0.74
36:5:1564:U:H2'	36:5:1565:G:C8	2.23	0.74
40:L3:169:THR:HG22	40:L3:171:LEU:H	1.52	0.74
40:L3:308:MET:HB2	40:L3:363:SER:HB2	1.69	0.74
45:L8:65:LEU:HD12	51:M5:25:VAL:HG13	1.86	0.74
36:1:2848:G:OP1	76:Q0:100:TYR:OH	2.05	0.74
1:2:1473:U:O2'	7:S5:103:ASN:ND2	2.20	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:S6:59:GLN:NE2	1:6:418:G:O2'	296.51	0.74
42:L5:285:ARG:NH1	37:7:62:U:O3'	340.93	0.74
53:M7:25:SER:O	53:M7:29:THR:HG23	1.88	0.74
1:2:1433:G:C8	31:D9:41:GLN:HG2	2.22	0.73
36:5:1487:G:H1	36:5:1855:U:H3	1.36	0.73
73:O7:45:ARG:NH2	36:5:361:A:O3'	123.78	0.73
36:5:501:A:H2'	36:5:502:U:C6	2.23	0.73
1:6:1432:U:H4'	1:6:1433:G:H5'	1.69	0.73
62:N6:5:SER:OG	62:N6:6:LEU:N	2.18	0.73
36:1:2208:A:N1	86:1:4039:OHX:N2	2.36	0.73
36:5:1015:U:O2'	36:5:1017:C:OP1	2.06	0.73
28:D6:10:ARG:HB2	28:D6:34:LYS:HA	1.71	0.73
28:D6:40:ALA:HB3	28:D6:69:ASN:HB3	5.55	0.73
6:S4:176:ASP:N	6:S4:176:ASP:OD2	2.83	0.73
8:S6:2:LYS:HB3	8:S6:108:VAL:HG12	5.62	0.73
36:1:595:G:N1	36:1:609:G:H5''	2.02	0.73
1:2:1500:C:OP1	21:C9:122:ARG:NH2	2.18	0.73
36:5:2971:A:H3'	36:5:2971:A:N3	2.02	0.73
1:6:800:U:H2'	1:6:801:G:H8	1.53	0.73
52:M6:15:LEU:HD11	52:M6:129:LEU:HD22	3.07	0.73
58:N2:35:LYS:NZ	58:N2:39:ASP:OD1	3.14	0.73
64:N8:73:LEU:HD11	64:N8:81:LEU:HD11	4.73	0.73
69:O3:86:ARG:O	86:O3:201:OHX:N1	2.22	0.73
1:2:474:A:OP2	11:S9:44:ARG:NH1	2.20	0.73
1:2:463:U:H2'	1:2:464:A:C8	2.23	0.73
36:5:3103:A:OP2	86:5:4154:OHX:N4	2.21	0.73
1:6:69:G:N1	1:6:82:U:O2	2.16	0.73
39:L2:64:ARG:NH1	45:L8:37:GLY:O	2.21	0.73
1:6:1057:U:O2'	1:6:1059:U:OP1	2.06	0.73
1:6:1211:A:H61	1:6:1452:U:H3	1.35	0.73
40:L3:274:SER:OG	36:5:3139:A:OP1	228.49	0.73
56:N0:16:THR:OG1	56:N0:19:VAL:N	2.21	0.73
64:N8:47:LYS:HE3	64:N8:48:TYR:CZ	2.22	0.73
66:O0:36:GLN:HB3	66:O0:38:LYS:HG3	1.70	0.73
36:1:2662:G:H2'	36:1:2663:G:H8	1.52	0.73
1:2:732:G:O2'	1:2:733:A:O4'	2.06	0.73
56:N0:114:HIS:CE1	36:5:1212:A:HO2'	310.60	0.73
38:8:2:A:H3'	38:8:3:A:H8	1.53	0.73
41:L4:226:GLU:OE2	41:L4:246:ARG:NH2	2.37	0.73
43:L6:64:LEU:HD22	43:L6:65:ILE:H	2.83	0.73
46:L9:126:VAL:HG21	46:L9:161:LEU:HA	2.28	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:M5:65:ARG:HB3	51:M5:127:TYR:HD1	4.23	0.73
78:Q2:14:GLY:O	78:Q2:16:THR:N	2.20	0.73
3:S1:171:ILE:HD13	3:S1:196:GLU:HG2	1.71	0.73
5:S3:57:ASP:OD2	5:S3:57:ASP:N	3.97	0.73
8:S6:12:SER:HB2	8:S6:124:LEU:HD12	1.71	0.73
36:1:529:A:H61	36:1:563:U:H3	1.36	0.73
1:2:1147:A:H2'	1:2:1148:C:H6	1.53	0.73
53:M7:139:TYR:CE2	36:5:2355:G:H4'	147.77	0.73
17:C5:52:LYS:HB2	17:C5:53:PRO:HD3	1.70	0.73
29:D7:67:THR:O	1:6:871:G:O2'	328.40	0.73
42:L5:40:HIS:HB3	42:L5:43:LYS:HG3	1.70	0.73
62:N6:3:LYS:HG3	62:N6:8:VAL:HG13	1.70	0.73
78:Q2:12:CYS:SG	78:Q2:74:CYS:SG	3.12	0.73
36:5:1565:G:N1	36:5:1574:C:N3	2.36	0.73
36:1:1073:U:H1'	65:N9:50:THR:HG22	1.69	0.73
68:O2:123:LYS:HA	68:O2:126:LEU:HB2	3.39	0.73
6:S4:68:ARG:O	6:S4:70:VAL:N	2.20	0.73
7:S5:92:ARG:NH2	7:S5:169:ASN:OD1	3.45	0.73
9:S7:122:HIS:HA	9:S7:125:ILE:HD12	2.05	0.73
36:5:953:G:H1'	36:5:1115:G:H5''	1.70	0.73
41:L4:3:ARG:HB3	41:L4:22:LEU:H	2.95	0.73
51:M5:114:ARG:HG2	51:M5:137:PRO:HG3	1.80	0.73
57:N1:84:TYR:HB2	65:N9:24:PRO:HB3	1.71	0.73
76:Q0:99:CYS:SG	87:Q0:500:ZN:ZN	1.76	0.73
3:S1:152:ARG:NH1	1:6:1799:U:O2'	342.16	0.73
3:S1:222:LYS:HD3	3:S1:223:PHE:H	1.54	0.73
3:S1:48:VAL:HG21	3:S1:61:LEU:HD22	4.35	0.73
5:S3:92:GLN:NE2	5:S3:92:GLN:O	2.22	0.73
36:1:1454:A:OP2	86:1:4205:OHX:N6	2.22	0.73
36:5:1605:A:O2'	36:5:1607:U:OP2	2.05	0.73
18:C6:41:PRO:HG2	18:C6:78:VAL:HG21	1.71	0.73
21:C9:49:ASP:OD1	21:C9:53:TRP:N	2.22	0.73
25:D3:50:LYS:HE2	1:6:435:C:H5''	351.30	0.73
40:L3:293:ASN:N	40:L3:293:ASN:OD1	2.21	0.73
43:L6:78:ARG:NH1	36:5:3272:C:OP2	246.71	0.73
47:M0:46:PHE:CD1	47:M0:140:THR:HA	2.47	0.73
3:S1:30:PHE:HB3	3:S1:96:LEU:HD22	3.04	0.73
5:S3:175:VAL:HG13	5:S3:182:LEU:HB2	1.71	0.73
1:2:1542:G:N2	1:2:1568:C:H1'	2.04	0.72
1:6:1280:C:H2'	1:6:1281:G:C8	2.20	0.72
1:6:1542:G:N2	1:6:1568:C:H1'	2.04	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:7:91:G:H2'	37:7:92:A:H8	1.53	0.72
13:C1:139:VAL:HG12	13:C1:140:VAL:H	1.54	0.72
26:D4:55:VAL:HG12	26:D4:75:VAL:HG13	7.98	0.72
42:L5:229:ASP:N	42:L5:229:ASP:OD1	2.21	0.72
36:1:1674:G:OP2	86:1:3940:OHX:N2	2.21	0.72
1:2:1488:G:H3'	1:2:1515:A:H61	1.54	0.72
40:L3:53:MET:HG2	40:L3:77:THR:HG22	1.70	0.72
41:L4:8:VAL:HB	41:L4:16:THR:HG21	4.19	0.72
46:L9:147:SER:HB2	46:L9:187:ILE:HD11	3.23	0.72
4:S2:140:ARG:NH1	23:D1:10:GLU:OE1	6.70	0.72
36:1:314:U:O4	86:1:4147:OHX:N4	2.22	0.72
36:1:3259:U:H6	36:1:3259:U:H5'	1.52	0.72
36:1:2310:U:OP1	86:1:4135:OHX:N2	2.22	0.72
1:2:1774:G:OP1	77:Q1:7:LYS:NZ	2.23	0.72
64:N8:29:PRO:HG3	36:5:937:G:H5''	174.88	0.72
1:6:1698:G:O2'	1:6:1699:G:O5'	2.06	0.72
13:C1:77:SER:HB3	13:C1:85:VAL:HB	2.56	0.72
17:C5:55:GLY:HA2	17:C5:58:LYS:HD3	1.70	0.72
18:C6:37:THR:O	18:C6:45:ARG:NH1	3.86	0.72
40:L3:10:ARG:NH2	40:L3:263:SER:O	2.22	0.72
61:N5:74:LYS:NZ	61:N5:78:ASP:OD2	2.21	0.72
3:S1:93:GLY:O	3:S1:95:ASN:N	2.65	0.72
10:S8:38:ILE:HG12	10:S8:96:LEU:HD11	5.28	0.72
37:3:77:G:N2	37:3:102:A:OP2	2.23	0.72
26:D4:112:LYS:NZ	1:6:57:G:OP1	346.71	0.72
49:M3:128:ARG:O	49:M3:130:GLY:N	2.18	0.72
71:O5:6:ALA:HA	71:O5:9:LEU:HD12	3.57	0.72
36:1:2946:A:H5''	36:1:2947:G:H5'	1.72	0.72
36:1:1231:A:OP2	86:1:4081:OHX:N6	2.23	0.72
36:5:3107:U:H2'	36:5:3108:G:C8	2.24	0.72
65:N9:17:HIS:HA	65:N9:20:GLY:HA2	1.71	0.72
36:1:1569:U:H5''	36:1:1570:U:H6	1.55	0.72
36:1:2296:A:OP1	86:1:4144:OHX:N2	2.23	0.72
1:2:1274:C:N4	35:SM:94:HIS:O	2.23	0.72
40:L3:233:TRP:CD1	40:L3:265:ALA:HB1	2.63	0.72
41:L4:204:GLY:O	41:L4:246:ARG:NH1	2.63	0.72
62:N6:100:HIS:ND1	62:N6:102:SER:OG	3.40	0.72
70:O4:80:ARG:HH11	70:O4:88:ARG:NH2	1.85	0.72
36:1:1523:U:HO2'	36:1:1607:U:HO2'	1.32	0.72
36:1:156:G:OP2	72:O6:27:SER:OG	2.05	0.72
18:C6:82:ARG:HH22	18:C6:114:ARG:HG3	3.69	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:L7:180:SER:HB2	44:L7:183:ASP:H	1.55	0.72
51:M5:136:ASP:OD2	51:M5:138:GLN:HG2	2.09	0.72
1:2:258:C:O2	10:S8:178:ARG:NH2	2.22	0.72
58:N2:51:GLY:O	58:N2:52:ASN:ND2	2.22	0.72
79:Q3:59:CYS:O	79:Q3:60:CYS:HB3	1.89	0.72
36:1:2554:A:H62	79:Q3:62:LYS:HZ2	1.38	0.72
2:S0:185:ARG:H	23:D1:45:ALA:H	2.53	0.72
11:S9:41:GLU:OE1	11:S9:126:ARG:NH2	2.23	0.72
20:C8:120:ARG:NH2	35:SM:58:GLU:OE1	2.19	0.72
36:5:2689:A:H4'	36:5:2690:G:H5'	1.72	0.72
36:5:2620:G:O6	86:5:4238:OHX:N4	2.23	0.72
1:6:755:A:H2'	1:6:756:A:H8	1.54	0.72
20:C8:84:TRP:HA	20:C8:89:GLN:HE22	1.52	0.72
1:2:1796:C:OP2	28:D6:5:ARG:NH1	2.23	0.72
41:L4:145:ILE:HD13	41:L4:150:LEU:HG	1.71	0.72
45:L8:36:ILE:HG22	45:L8:37:GLY:H	1.53	0.72
49:M3:157:ARG:HG2	49:M3:158:ALA:H	1.54	0.72
3:S1:141:ALA:HB1	3:S1:207:LEU:HD22	3.77	0.72
11:S9:157:ASP:OD1	11:S9:158:PHE:N	3.50	0.72
20:C8:73:MET:HG2	20:C8:101:LEU:HD11	4.24	0.72
46:L9:12:VAL:HB	46:L9:51:GLN:HA	1.71	0.72
47:M0:48:LEU:HA	47:M0:178:ARG:HH12	1.54	0.72
48:M1:49:LYS:HB3	48:M1:62:ASN:HA	1.71	0.72
53:M7:105:LYS:HB3	53:M7:107:LEU:HD13	5.11	0.72
69:O3:72:THR:HG23	69:O3:83:ALA:HA	1.71	0.72
5:S3:170:THR:HG22	5:S3:187:LYS:HA	6.17	0.72
86:1:4080:OHX:N4	55:M9:14:VAL:O	2.23	0.71
1:2:452:A:OP2	86:2:2038:OHX:N5	2.22	0.71
36:5:3163:A:N6	36:5:3287:U:O4	2.20	0.71
72:O6:28:TYR:O	86:5:4184:OHX:N2	103.49	0.71
36:5:495:G:O6	36:5:618:C:N4	2.20	0.71
7:S5:37:GLN:HB3	18:C6:53:LEU:HD22	1.72	0.71
21:C9:52:GLY:O	21:C9:54:PHE:N	2.20	0.71
26:D4:20:ARG:HD2	26:D4:74:LEU:HD22	3.09	0.71
46:L9:91:ARG:NH2	46:L9:140:VAL:O	7.30	0.71
36:1:745:C:H5''	54:M8:145:ASN:ND2	2.05	0.71
59:N3:15:LEU:HD13	59:N3:51:ALA:HB3	2.47	0.71
1:2:1280:C:H2'	1:2:1281:G:C8	2.25	0.71
37:7:112:G:OP2	86:7:223:OHX:N2	2.22	0.71
40:L3:166:ILE:O	40:L3:169:THR:HG22	4.31	0.71
46:L9:49:ASN:HD21	46:L9:52:LEU:HB2	1.55	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
60:N4:49:ILE:O	60:N4:52:THR:OG1	2.08	0.71
8:S6:206:ALA:O	8:S6:210:GLN:NE2	5.00	0.71
34:SR:201:THR:HB	34:SR:242:SER:H	1.55	0.71
1:2:275:C:O2	1:2:276:C:N4	2.24	0.71
1:2:320:U:H3'	1:2:321:C:H5''	1.72	0.71
1:6:1542:G:N2	1:6:1569:A:OP2	2.23	0.71
25:D3:83:VAL:HG21	25:D3:122:PHE:HE2	4.56	0.71
71:O5:101:THR:HG22	71:O5:104:GLN:H	1.55	0.71
74:O8:13:GLU:HA	74:O8:16:ARG:HH21	1.55	0.71
7:S5:200:ASN:HB3	7:S5:208:SER:HB2	4.23	0.71
36:5:1502:C:OP1	86:5:3905:OHX:N4	2.23	0.71
1:6:1799:U:H4'	1:6:1800:A:H2'	1.71	0.71
1:6:781:U:H6	1:6:782:U:H3	1.36	0.71
55:M9:90:PRO:HB2	55:M9:93:VAL:HG23	1.72	0.71
73:O7:18:LEU:HA	73:O7:25:ARG:H	1.55	0.71
74:O8:24:THR:HG23	74:O8:44:LYS:HB2	4.39	0.71
4:S2:78:ASP:OD2	4:S2:79:GLU:N	2.18	0.71
10:S8:8:ARG:NH2	10:S8:28:GLU:OE1	9.45	0.71
36:1:1750:A:H4'	36:1:1751:G:H5'	1.71	0.71
1:2:513:U:OP1	11:S9:133:HIS:NE2	2.22	0.71
28:D6:79:ILE:HD11	1:6:1795:U:H5'	334.26	0.71
40:L3:105:VAL:HG21	40:L3:148:LEU:HD13	2.09	0.71
7:S5:62:VAL:HG13	7:S5:89:ILE:HG12	1.73	0.71
34:SR:216:LYS:HA	34:SR:239:GLU:HG3	1.73	0.71
1:6:301:A:OP2	86:6:2092:OHX:N1	2.23	0.71
25:D3:61:SER:HB2	25:D3:116:ASP:HB2	1.72	0.71
40:L3:102:LEU:O	36:5:3147:G:H4'	240.57	0.71
41:L4:157:GLU:HG3	41:L4:251:THR:HG21	1.73	0.71
41:L4:200:THR:OG1	41:L4:201:GLN:N	2.22	0.71
47:M0:208:ASN:HA	47:M0:211:ARG:HD2	2.30	0.71
65:N9:23:LYS:HD2	65:N9:24:PRO:HD3	2.94	0.71
5:S3:76:ARG:NH1	5:S3:76:ARG:O	2.24	0.71
36:1:1064:A:N6	36:1:1096:U:H3	1.88	0.71
36:1:917:A:OP2	86:1:4140:OHX:N2	2.24	0.71
1:2:499:U:O2'	1:2:500:C:O5'	2.09	0.71
1:2:895:G:H1	1:2:917:U:H3	1.37	0.71
52:M6:85:ARG:NH1	36:5:2382:G:OP1	238.37	0.71
1:6:1015:U:OP1	86:6:2055:OHX:N3	2.24	0.71
25:D3:39:LYS:NZ	1:6:1742:U:OP1	314.51	0.71
52:M6:27:LEU:HD22	52:M6:101:ARG:HG3	1.72	0.71
35:SM:34:LYS:NZ	36:1:2707:C:OP1	2.22	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:1495:U:H5	36:1:1835:A:N1	1.89	0.71
36:5:23:A:OP1	86:5:3900:OHX:N4	2.23	0.71
41:L4:44:LYS:HB3	41:L4:47:ARG:HH11	2.28	0.71
42:L5:85:ARG:HH21	42:L5:254:LYS:H	1.37	0.71
44:L7:217:PRO:O	86:5:3995:OHX:N6	259.45	0.71
45:L8:100:GLU:OE2	45:L8:108:ARG:NH1	2.49	0.71
60:N4:35:LYS:HE2	60:N4:51:TRP:CZ2	2.54	0.71
61:N5:105:VAL:HG13	61:N5:130:TYR:CD2	2.25	0.71
78:Q2:71:ARG:HH21	78:Q2:80:ARG:HH11	3.67	0.71
4:S2:137:ILE:HG12	4:S2:138:PRO:HD2	1.72	0.71
36:1:2554:A:C8	36:1:2554:A:H5'	2.26	0.71
36:5:1152:G:H22	36:5:1200:A:H61	1.38	0.71
40:L3:325:LYS:NZ	36:5:3097:C:OP1	259.15	0.71
49:M3:35:ARG:HH12	36:5:685:G:P	83.36	0.71
17:C5:130:ARG:NH1	35:SM:71:ASN:OD1	2.24	0.71
22:D0:57:ARG:HD2	22:D0:89:ARG:HD3	1.73	0.71
56:N0:137:ARG:NH1	36:5:1213:G:OP1	324.82	0.71
58:N2:59:ASP:O	58:N2:61:THR:N	2.20	0.71
61:N5:56:ARG:O	61:N5:61:LYS:HD2	1.91	0.71
72:O6:58:ILE:HA	72:O6:61:ILE:HD12	1.73	0.71
36:5:1017:C:N4	36:5:2671:A:OP2	2.23	0.71
15:C3:47:PRO:HA	15:C3:50:ILE:HD12	1.73	0.71
21:C9:68:ARG:NH1	1:6:1521:G:O6	416.09	0.71
41:L4:192:GLY:O	41:L4:195:ARG:N	3.22	0.71
5:S3:66:ILE:HD11	5:S3:88:ALA:HB2	1.71	0.71
9:S7:67:LEU:HG	9:S7:94:ALA:HB2	3.94	0.71
36:1:567:G:O6	86:1:3997:OHX:N1	2.24	0.70
12:C0:16:PHE:HD2	12:C0:76:LEU:HD23	1.55	0.70
26:D4:113:ASN:HA	26:D4:116:LYS:HD3	1.73	0.70
33:E1:82:LYS:O	33:E1:84:VAL:N	4.98	0.70
59:N3:2:SER:HA	59:N3:56:ASP:HA	3.77	0.70
69:O3:48:ARG:HH11	69:O3:48:ARG:HG2	1.54	0.70
8:S6:84:TYR:OH	8:S6:91:GLU:O	2.39	0.70
11:S9:42:ILE:O	11:S9:45:ILE:N	3.51	0.70
34:SR:258:THR:O	34:SR:275:ARG:NH1	2.23	0.70
34:SR:80:ALA:HB3	34:SR:92:TRP:HB2	1.71	0.70
36:1:3294:A:H2'	36:1:3295:A:O4'	1.90	0.70
1:2:542:A:H2'	1:2:543:C:H5'	1.72	0.70
36:5:240:U:HO2'	36:5:241:G:H8	1.39	0.70
36:5:3334:U:OP2	86:5:4228:OHX:N6	2.25	0.70
1:6:1600:A:H4'	1:6:1601:G:OP1	1.91	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:D7:67:THR:OG1	29:D7:70:LYS:O	2.84	0.70
42:L5:279:LYS:HD3	42:L5:282:ARG:HB2	1.72	0.70
49:M3:46:ILE:HG23	49:M3:49:ARG:HB2	2.52	0.70
53:M7:21:TYR:H	53:M7:145:HIS:CE1	2.73	0.70
60:N4:63:ILE:O	60:N4:65:GLU:N	3.23	0.70
67:O1:44:MET:O	67:O1:46:THR:N	3.92	0.70
76:Q0:118:THR:HG23	76:Q0:120:GLN:H	3.61	0.70
36:1:1740:U:H1'	36:1:1741:A:H2	1.56	0.70
29:D7:6:ASP:OD1	29:D7:9:HIS:ND1	2.58	0.70
44:L7:228:SER:HA	44:L7:232:ARG:HH21	2.23	0.70
49:M3:187:ALA:HA	49:M3:190:LYS:HB3	1.72	0.70
63:N7:124:ALA:O	63:N7:126:LYS:N	2.25	0.70
2:S0:13:ASP:OD1	2:S0:179:ARG:NH2	2.19	0.70
3:S1:70:LEU:HA	3:S1:73:LEU:HG	1.72	0.70
34:SR:161:LYS:HE3	34:SR:164:ASP:HB2	1.73	0.70
36:1:1064:A:H4'	36:1:1065:A:O5'	1.91	0.70
36:5:442:G:N2	36:5:491:C:O2'	2.24	0.70
1:6:815:G:H5'	1:6:815:G:H8	1.55	0.70
12:C0:16:PHE:HZ	12:C0:77:ARG:HH21	1.39	0.70
26:D4:112:LYS:HZ3	26:D4:112:LYS:HB3	1.56	0.70
31:D9:19:ARG:HD3	31:D9:32:ARG:HD2	2.35	0.70
42:L5:270:LYS:HE2	42:L5:273:ARG:HA	8.51	0.70
45:L8:36:ILE:O	45:L8:38:GLN:N	2.23	0.70
45:L8:41:GLN:HG3	45:L8:44:ARG:HH12	2.10	0.70
7:S5:41:LYS:HZ3	7:S5:67:PRO:HG2	5.58	0.70
1:2:565:C:O2	86:D3:202:OHX:N5	2.24	0.70
36:5:314:U:H2'	36:5:315:C:C6	2.27	0.70
36:5:990:U:O4	86:5:4179:OHX:N6	2.25	0.70
24:D2:30:SER:HB3	24:D2:59:GLY:HA3	2.59	0.70
25:D3:6:PRO:HG3	25:D3:14:LYS:HG2	1.74	0.70
24:D2:60:LYS:NZ	29:D7:24:LEU:O	3.39	0.70
41:L4:64:SER:OG	41:L4:65:TRP:N	2.82	0.70
8:S6:52:ILE:HA	8:S6:111:LEU:HD23	1.73	0.70
36:1:1211:U:H2'	36:1:1212:A:C8	2.26	0.70
36:1:1596:C:H2'	36:1:1597:C:C6	2.26	0.70
36:1:964:G:O2'	64:N8:41:HIS:NE2	2.11	0.70
47:M0:154:ARG:NH1	36:5:2838:A:OP1	326.87	0.70
36:5:510:G:O6	86:5:4017:OHX:N2	2.24	0.70
1:6:1182:U:H2'	1:6:1184:A:OP2	1.91	0.70
12:C0:56:LYS:HB3	12:C0:67:THR:HG23	5.12	0.70
18:C6:6:SER:HB2	18:C6:23:LYS:HB3	2.70	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:S5:81:ARG:HH21	30:D8:47:PRO:HB3	1.57	0.70
41:L4:287:THR:O	41:L4:291:ASN:ND2	4.17	0.70
36:1:1051:U:H4'	57:N1:19:PHE:CD2	2.25	0.70
1:2:741:C:O2	9:S7:107:ARG:NH2	2.21	0.70
11:S9:96:VAL:HA	11:S9:99:LEU:HD22	1.73	0.70
36:1:1772:U:H5''	36:1:1773:C:H5'	1.74	0.70
1:2:480:G:H22	1:2:509:G:H1'	1.57	0.70
40:L3:347:SER:O	40:L3:349:LYS:N	2.87	0.70
54:M8:122:ILE:HG23	54:M8:126:GLN:HB2	1.84	0.70
54:M8:65:SER:HA	54:M8:93:ILE:HD13	1.74	0.70
10:S8:21:PHE:HD1	10:S8:22:ARG:HG2	4.73	0.70
36:1:1603:A:OP1	55:M9:38:ARG:NH1	2.25	0.70
36:1:3060:C:OP1	86:1:4034:OHX:N4	2.24	0.70
36:1:831:G:O6	86:1:3881:OHX:N4	2.24	0.70
1:2:645:C:H42	1:2:689:G:H1	1.38	0.70
39:L2:9:ARG:NH1	36:5:912:G:OP2	179.80	0.70
1:6:218:A:H2'	1:6:219:A:H5''	1.72	0.70
13:C1:69:LYS:H	13:C1:127:GLN:HB3	1.55	0.70
19:C7:33:ARG:NH2	34:SR:109:ASP:OD1	3.04	0.70
39:L2:143:GLU:O	39:L2:145:LYS:N	2.92	0.70
49:M3:46:ILE:HG12	49:M3:49:ARG:NH1	3.49	0.70
36:1:1646:G:O2'	36:1:1808:G:N2	2.22	0.70
1:2:324:U:OP1	13:C1:133:LYS:NZ	2.25	0.70
36:5:2975:U:OP1	86:5:4083:OHX:N3	2.25	0.70
41:L4:207:VAL:HB	41:L4:227:THR:HG22	3.05	0.70
49:M3:165:SER:O	49:M3:167:PHE:N	2.24	0.70
49:M3:79:GLU:OE1	49:M3:101:ARG:NH2	2.25	0.70
74:O8:46:ARG:NH1	74:O8:47:GLY:O	2.83	0.70
6:S4:187:ARG:HH22	1:6:753:A:H62	376.19	0.70
34:SR:299:GLN:NE2	34:SR:315:VAL:O	2.25	0.70
36:1:1569:U:H5'	36:1:1570:U:H5''	1.72	0.70
55:M9:166:ASN:HD22	55:M9:167:ARG:HG2	6.77	0.70
35:SM:97:THR:HG22	35:SM:99:LYS:HG3	1.72	0.70
36:1:2768:U:H2'	36:1:2769:A:H8	1.57	0.69
36:1:413:U:OP1	53:M7:30:ARG:NH2	2.22	0.69
26:D4:122:GLY:O	26:D4:125:LEU:N	2.44	0.69
32:E0:48:THR:OG1	32:E0:49:LEU:N	2.65	0.69
48:M1:16:LYS:HB2	48:M1:72:ARG:HG2	1.73	0.69
66:O0:10:ILE:HD11	66:O0:104:LEU:HD11	5.12	0.69
72:O6:45:ARG:NH2	72:O6:54:GLU:OE1	2.77	0.69
73:O7:88:ALA:O	86:O7:104:OHX:N1	2.24	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:368:G:OP1	86:1:3876:OHX:N1	2.25	0.69
36:1:3200:G:O6	86:1:4124:OHX:N4	2.25	0.69
36:5:3023:U:OP2	36:5:3031:G:N1	2.23	0.69
1:6:1370:U:O4	86:6:2143:OHX:N6	2.25	0.69
13:C1:71:LEU:HB3	13:C1:88:ARG:HD3	1.74	0.69
16:C4:126:THR:HG21	1:6:888:U:H1'	275.53	0.69
17:C5:65:LEU:O	86:C5:201:OHX:N2	4.29	0.69
26:D4:12:VAL:HG13	26:D4:23:PHE:HB3	1.74	0.69
40:L3:139:GLN:O	40:L3:141:GLY:N	2.30	0.69
59:N3:10:LYS:NZ	59:N3:56:ASP:OD1	2.22	0.69
60:N4:6:ASP:OD2	60:N4:8:PHE:N	2.20	0.69
2:S0:56:LYS:NZ	2:S0:159:ALA:O	2.24	0.69
3:S1:214:LYS:NZ	1:6:886:U:OP1	284.82	0.69
9:S7:126:LEU:HB2	9:S7:173:TYR:HE2	6.28	0.69
36:1:1390:A:N6	36:1:1418:A:O2'	2.26	0.69
78:Q2:41:ARG:NH1	36:5:284:A:OP2	156.42	0.69
1:6:687:G:H2'	1:6:688:G:H8	1.57	0.69
33:E1:91:ILE:HG12	33:E1:92:LYS:HG2	6.20	0.69
55:M9:76:SER:O	55:M9:81:ARG:NH1	2.25	0.69
79:Q3:15:GLY:O	79:Q3:23:ARG:NH1	4.24	0.69
6:S4:122:LYS:HD2	6:S4:164:LEU:HD21	1.75	0.69
36:1:3035:A:OP2	86:1:4070:OHX:N4	2.25	0.69
36:1:600:G:N7	86:1:4092:OHX:N1	2.39	0.69
36:1:975:C:H2'	36:1:976:U:H6	1.57	0.69
1:2:1386:G:OP2	19:C7:44:LYS:NZ	2.25	0.69
36:5:1597:C:H2'	36:5:1598:G:H8	1.58	0.69
36:5:1877:U:H5''	36:5:1878:G:H5'	1.72	0.69
14:C2:103:LEU:HG	14:C2:116:VAL:HG22	1.75	0.69
21:C9:33:TYR:HD1	21:C9:34:VAL:H	2.27	0.69
44:L7:219:LYS:O	44:L7:228:SER:HB2	2.61	0.69
51:M5:16:SER:O	51:M5:20:ARG:HG2	1.93	0.69
70:O4:74:ARG:HD3	70:O4:85:VAL:HG21	3.59	0.69
70:O4:84:CYS:O	70:O4:88:ARG:HB2	3.40	0.69
36:1:230:U:H2'	36:1:231:G:O4'	1.92	0.69
36:1:2924:U:O4	86:1:4012:OHX:N1	2.25	0.69
36:1:612:U:H2'	36:1:613:G:H8	1.56	0.69
36:5:561:C:H2'	36:5:562:C:H6	1.57	0.69
22:D0:27:THR:HB	22:D0:88:LYS:HG3	1.74	0.69
28:D6:89:ARG:HH11	28:D6:92:ARG:HH21	1.40	0.69
33:E1:126:CYS:O	33:E1:128:ALA:N	2.25	0.69
41:L4:157:GLU:HG2	41:L4:209:TYR:HB2	1.75	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:L5:144:VAL:HG12	42:L5:173:VAL:HG22	1.74	0.69
49:M3:48:PRO:HD2	71:O5:115:LYS:HD2	2.98	0.69
72:O6:10:GLY:H	72:O6:13:LYS:HG2	2.17	0.69
20:C8:132:ARG:NH2	1:6:1173:C:OP1	344.57	0.69
8:S6:13:GLN:NE2	1:6:151:G:H21	311.37	0.69
1:6:1588:G:H1	1:6:1608:U:H3	1.40	0.69
15:C3:84:ILE:HG22	15:C3:135:LEU:HD21	1.74	0.69
22:D0:26:LEU:HB2	22:D0:89:ARG:HB2	2.79	0.69
61:N5:103:TYR:HB3	61:N5:135:ILE:HD11	3.18	0.69
68:O2:22:SER:HA	68:O2:28:VAL:HB	2.78	0.69
1:2:81:G:OP2	86:2:2140:OHX:N5	2.26	0.69
36:5:2762:A:OP2	86:5:3982:OHX:N5	2.25	0.69
18:C6:93:HIS:HA	18:C6:97:VAL:HG23	1.92	0.69
41:L4:3:ARG:HH11	41:L4:22:LEU:HD22	1.57	0.69
44:L7:154:GLY:N	44:L7:161:VAL:O	2.73	0.69
78:Q2:56:PRO:HB3	36:5:2802:A:C8	184.11	0.69
34:SR:17:ASN:O	34:SR:308:ASN:ND2	3.77	0.69
34:SR:48:THR:HG22	34:SR:55:GLY:HA2	5.59	0.69
36:1:640:U:OP1	64:N8:21:ARG:NH2	2.24	0.69
1:2:1533:C:H4'	1:2:1539:G:N1	2.06	0.69
21:C9:53:TRP:HH2	21:C9:100:ILE:HD12	2.16	0.69
26:D4:5:VAL:O	26:D4:6:THR:OG1	2.11	0.69
43:L6:31:ARG:NH2	43:L6:81:ALA:O	2.44	0.69
3:S1:40:ASN:ND2	3:S1:42:ASN:O	2.25	0.69
7:S5:123:VAL:HG12	7:S5:124:LEU:HD12	1.73	0.69
36:1:1941:C:O2'	36:1:3344:A:N6	2.26	0.69
36:1:367:A:OP1	86:1:3876:OHX:N2	2.26	0.69
1:2:1277:G:H2'	1:2:1278:G:O4'	1.92	0.69
1:2:641:G:H1	1:2:693:U:H3	1.39	0.69
37:3:19:C:H2'	37:3:20:A:H8	1.57	0.69
36:5:1170:A:OP2	86:5:3995:OHX:N4	2.26	0.69
12:C0:64:TYR:OH	1:6:1435:G:O6	425.64	0.69
40:L3:10:ARG:NH1	40:L3:11:HIS:O	3.18	0.69
42:L5:120:LYS:O	42:L5:248:ARG:NH2	2.46	0.69
52:M6:22:VAL:HG11	52:M6:120:VAL:HG11	2.29	0.69
61:N5:49:LYS:O	61:N5:51:VAL:N	2.25	0.69
62:N6:57:LEU:HD23	62:N6:67:GLU:HB3	1.75	0.69
36:1:2854:U:OP2	47:M0:3:ARG:NH2	2.26	0.69
36:1:3358:U:H2'	36:1:3359:A:O4'	1.93	0.69
36:5:2818:U:H6	36:5:2818:U:H5'	1.56	0.69
36:5:438:A:N1	36:5:621:A:N6	2.38	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:C4:111:ARG:NH1	28:D6:57:SER:O	4.40	0.69
16:C4:125:SER:OG	16:C4:126:THR:N	2.26	0.69
18:C6:34:SER:HB3	18:C6:38:LEU:HD12	1.74	0.69
21:C9:73:VAL:HG11	21:C9:102:ARG:HB2	1.73	0.69
26:D4:87:PRO:HG2	26:D4:90:ARG:HG3	3.39	0.69
71:O5:64:GLU:OE1	71:O5:68:GLN:NE2	6.01	0.69
5:S3:53:THR:HG22	5:S3:91:VAL:HG11	2.75	0.69
1:2:734:A:H5''	1:2:735:C:OP1	1.93	0.69
36:5:1615:C:H2'	36:5:1616:U:H6	1.56	0.69
1:6:1114:G:O2'	1:6:1130:G:O6	2.10	0.69
1:6:1564:U:H2'	1:6:1565:C:C6	2.28	0.69
1:6:1783:C:H2'	1:6:1784:C:C6	2.27	0.69
12:C0:50:THR:HG22	12:C0:55:VAL:HG22	1.91	0.69
40:L3:187:SER:HB2	40:L3:188:ILE:HD12	3.58	0.69
45:L8:89:GLU:HA	45:L8:92:LYS:HE2	1.74	0.69
59:N3:83:LYS:HE2	59:N3:84:SER:N	2.08	0.69
2:S0:183:ARG:NH2	2:S0:191:ARG:O	2.27	0.69
7:S5:71:ALA:HB1	7:S5:91:GLU:HG3	1.76	0.69
1:2:463:U:H2'	1:2:464:A:H8	1.56	0.68
36:5:2234:G:O6	86:5:3955:OHX:N1	2.26	0.68
12:C0:53:GLY:O	12:C0:55:VAL:N	2.26	0.68
1:2:867:G:OP2	15:C3:3:ARG:NH1	2.26	0.68
15:C3:54:LEU:HB3	15:C3:60:VAL:HG11	3.27	0.68
24:D2:7:LEU:HD22	24:D2:11:LEU:HG	1.74	0.68
52:M6:54:TYR:CD2	52:M6:145:VAL:HG11	2.66	0.68
66:O0:42:ILE:HG12	66:O0:67:VAL:HG22	4.05	0.68
6:S4:146:THR:HG21	1:6:123:G:H21	342.13	0.68
10:S8:138:ASN:N	10:S8:138:ASN:OD1	2.23	0.68
36:1:1752:A:OP2	86:1:4042:OHX:N5	2.26	0.68
1:2:705:U:H2'	1:2:706:A:C8	2.27	0.68
36:5:1103:A:H3'	36:5:1104:G:H5'	1.75	0.68
63:N7:69:LYS:NZ	36:5:1632:A:OP1	192.54	0.68
69:O3:60:ARG:HD2	36:5:3275:U:C2	214.21	0.68
1:6:976:G:O6	86:6:2079:OHX:N6	2.26	0.68
41:L4:294:GLU:N	41:L4:294:GLU:OE1	2.23	0.68
51:M5:95:GLN:OE1	36:5:288:C:O2'	132.00	0.68
53:M7:67:ILE:HG22	53:M7:80:LYS:HD2	1.74	0.68
57:N1:83:ARG:NH1	57:N1:85:LEU:HD21	2.08	0.68
62:N6:35:LEU:HD11	62:N6:45:ILE:HG22	2.49	0.68
1:2:526:A:OP2	26:D4:93:ARG:NH2	2.26	0.68
40:L3:232:ARG:NH1	40:L3:269:GLN:O	2.31	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:L3:41:VAL:HA	40:L3:185:GLY:CA	2.38	0.68
40:L3:92:TYR:HB2	40:L3:157:VAL:HG13	1.73	0.68
47:M0:146:ASP:N	47:M0:146:ASP:OD1	2.27	0.68
70:O4:82:ALA:O	70:O4:85:VAL:N	2.96	0.68
7:S5:145:ASP:OD2	7:S5:146:THR:N	2.26	0.68
34:SR:42:LEU:HB2	34:SR:61:PHE:HB2	1.90	0.68
36:1:2146:C:OP1	39:L2:200:ARG:NH1	2.27	0.68
86:2:2031:OHX:N4	86:2:2146:OHX:N2	2.41	0.68
46:L9:70:THR:HG21	36:5:3122:A:N1	323.64	0.68
36:5:3194:C:N3	36:5:3197:G:N2	2.41	0.68
53:M7:120:ASN:HD22	38:8:13:A:H1'	141.18	0.68
18:C6:97:VAL:HG12	18:C6:98:ASP:H	1.58	0.68
42:L5:146:LEU:HB3	36:5:2746:A:H2	258.92	0.68
42:L5:4:GLN:N	42:L5:4:GLN:OE1	2.26	0.68
52:M6:110:PRO:HA	52:M6:113:ASP:OD2	1.94	0.68
60:N4:39:LEU:HD12	60:N4:44:LYS:HG3	2.13	0.68
36:1:18:G:OP1	71:O5:81:ARG:NH2	2.26	0.68
78:Q2:46:LYS:O	86:Q2:503:OHX:N6	2.26	0.68
8:S6:13:GLN:HE22	1:6:151:G:H21	311.03	0.68
36:1:3074:G:OP1	86:1:4034:OHX:N1	2.26	0.68
1:2:1082:C:H6	23:D1:62:ARG:HH21	1.42	0.68
1:2:356:G:OP2	86:2:2036:OHX:N6	2.27	0.68
1:6:27:U:H2'	1:6:28:A:C8	2.29	0.68
14:C2:55:GLY:N	35:SM:172:VAL:O	2.26	0.68
19:C7:105:GLN:O	19:C7:109:LEU:N	2.75	0.68
25:D3:51:GLY:HA2	25:D3:77:ILE:HG13	2.77	0.68
40:L3:236:LYS:HD2	36:5:2340:U:OP1	233.86	0.68
45:L8:130:TYR:HD1	45:L8:202:GLU:HB3	1.56	0.68
52:M6:88:VAL:HG11	52:M6:99:LEU:HD21	1.76	0.68
59:N3:81:GLN:O	59:N3:98:ASN:ND2	2.26	0.68
9:S7:95:GLU:OE1	9:S7:97:ARG:NH1	5.91	0.68
36:1:266:A:H2'	72:O6:30:LYS:HE3	1.75	0.68
36:1:595:G:H1	36:1:609:G:H5''	1.58	0.68
36:1:612:U:H2'	36:1:613:G:C8	2.29	0.68
36:5:1662:G:H1	36:5:1787:A:H61	1.40	0.68
16:C4:52:ARG:HH21	1:6:905:A:H4'	300.66	0.68
25:D3:64:PRO:O	86:D3:202:OHX:N1	2.26	0.68
42:L5:270:LYS:HG3	42:L5:273:ARG:HB2	4.64	0.68
51:M5:157:LYS:NZ	36:5:58:G:OP1	85.54	0.68
67:O1:43:HIS:O	67:O1:44:MET:HE2	4.75	0.68
9:S7:96:ARG:NH1	9:S7:128:ASP:OD2	2.26	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:2898:G:H5''	36:1:2899:C:H5'	1.76	0.68
1:2:142:G:H22	1:2:173:A:H2	1.39	0.68
1:2:702:G:O6	1:2:737:A:N6	2.26	0.68
36:5:1541:G:OP2	86:5:4088:OHX:N4	2.26	0.68
36:5:1578:C:H3'	36:5:1579:C:H6	1.59	0.68
36:5:171:G:N2	36:5:248:U:O2	2.26	0.68
36:5:2718:U:H2'	36:5:2719:U:C6	2.29	0.68
1:6:1535:U:H1'	1:6:1536:G:C2	2.29	0.68
1:6:453:U:O4	86:6:2061:OHX:N4	2.27	0.68
1:6:8:U:O2'	86:6:2071:OHX:N2	2.27	0.68
15:C3:34:ILE:HG13	15:C3:67:THR:HG21	1.75	0.68
41:L4:264:SER:C	41:L4:266:THR:H	1.96	0.68
41:L4:89:ALA:O	41:L4:91:GLY:N	2.27	0.68
45:L8:101:THR:HG23	45:L8:104:GLU:H	1.57	0.68
54:M8:176:ARG:NH1	64:N8:46:ASP:OD2	2.26	0.68
62:N6:27:ARG:HH12	62:N6:76:LEU:HA	2.41	0.68
36:1:1642:A:N3	36:1:1822:C:O2'	2.24	0.68
1:2:1795:U:O2	28:D6:10:ARG:HD2	1.93	0.68
44:L7:158:LYS:HE2	36:5:1362:G:H21	211.89	0.68
36:5:155:G:H5''	36:5:156:G:C8	2.28	0.68
36:5:655:C:H2'	36:5:656:A:H8	1.59	0.68
1:6:1767:G:OP1	1:6:1770:U:H4'	1.93	0.68
38:8:124:G:OP2	86:8:222:OHX:N2	2.27	0.68
13:C1:33:ARG:NH2	13:C1:51:GLY:O	3.04	0.68
36:1:2392:C:O2'	40:L3:266:ARG:NH2	2.25	0.68
49:M3:175:SER:O	49:M3:178:LYS:N	2.26	0.68
8:S6:57:ASP:HA	8:S6:106:LEU:HA	1.93	0.68
36:1:1819:U:O4	86:1:4036:OHX:N4	2.26	0.68
36:5:783:A:OP2	86:5:4188:OHX:N6	2.27	0.68
1:6:1688:U:O2	1:6:1713:G:N2	2.24	0.68
12:C0:31:LYS:H	12:C0:38:LYS:HA	4.12	0.68
16:C4:19:ILE:HB	16:C4:83:ILE:HD12	1.75	0.68
16:C4:31:THR:OG1	16:C4:32:ASP:N	3.20	0.68
18:C6:31:VAL:HG13	18:C6:67:VAL:HB	2.25	0.68
52:M6:61:ALA:HA	52:M6:70:PRO:HD2	1.75	0.68
68:O2:103:LYS:O	68:O2:106:VAL:HG12	1.93	0.68
71:O5:64:GLU:HA	71:O5:67:ARG:HB2	2.51	0.68
78:Q2:12:CYS:SG	78:Q2:74:CYS:CB	3.43	0.68
34:SR:169:ILE:HG13	34:SR:181:TRP:HB2	1.76	0.68
1:2:702:G:N7	86:2:2129:OHX:N2	2.42	0.68
37:3:71:G:H2'	37:3:72:A:C8	2.29	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
64:N8:10:LYS:HE3	36:5:1375:G:O6	158.58	0.68
20:C8:41:ARG:HD3	1:6:1565:C:OP1	368.78	0.68
16:C4:131:GLY:O	16:C4:133:ARG:N	3.53	0.68
18:C6:16:ALA:HB2	18:C6:72:GLY:HA3	1.76	0.68
25:D3:91:GLY:O	25:D3:93:LEU:N	2.27	0.68
40:L3:303:LYS:HD2	40:L3:361:THR:HG21	1.92	0.68
45:L8:239:GLY:O	45:L8:241:LYS:N	2.66	0.68
73:O7:55:ARG:NH2	36:5:347:G:N7	108.95	0.68
1:2:68:A:OP1	8:S6:160:ARG:NH1	2.27	0.68
36:1:3057:U:H5'	36:1:3086:A:H61	1.57	0.67
36:1:272:G:OP2	86:1:4025:OHX:N3	2.28	0.67
1:2:973:A:H2'	1:2:974:A:H8	1.58	0.67
36:5:1661:G:H2'	36:5:1662:G:C8	2.29	0.67
24:D2:11:LEU:HD12	24:D2:74:VAL:HB	1.76	0.67
46:L9:85:GLY:HA3	46:L9:187:ILE:HD12	1.75	0.67
47:M0:189:GLU:HB3	47:M0:200:LEU:HB3	1.75	0.67
63:N7:128:GLN:O	63:N7:130:PHE:N	2.66	0.67
36:1:541:U:O4	86:1:4189:OHX:N2	2.26	0.67
36:5:3255:U:H2'	36:5:3256:G:C8	2.28	0.67
36:5:685:G:N2	36:5:695:C:O2	2.18	0.67
1:6:1017:U:H2'	1:6:1018:U:C6	2.29	0.67
31:D9:5:ASN:OD1	31:D9:7:TRP:NE1	2.28	0.67
55:M9:15:VAL:HG11	55:M9:52:LYS:HG3	1.75	0.67
68:O2:94:ALA:O	68:O2:120:THR:HG23	2.24	0.67
68:O2:21:HIS:CE1	68:O2:24:ARG:HD2	2.30	0.67
2:S0:147:THR:HG21	2:S0:159:ALA:HB1	1.75	0.67
36:1:561:C:H2'	36:1:562:C:C6	2.29	0.67
1:2:1230:A:H2'	1:2:1258:U:C5	2.29	0.67
36:5:1192:C:N4	36:5:1301:A:O3'	2.27	0.67
36:5:1464:G:O2'	86:5:3905:OHX:N5	2.27	0.67
1:6:1699:G:N1	1:6:1701:A:H5''	2.09	0.67
25:D3:30:LYS:HE3	25:D3:34:LEU:HD11	1.76	0.67
46:L9:171:ASP:OD2	46:L9:173:ARG:NH1	2.27	0.67
47:M0:87:LEU:HD23	47:M0:138:VAL:HG22	4.33	0.67
56:N0:1:MET:N	56:N0:32:SER:OG	6.85	0.67
63:N7:14:VAL:HG13	70:O4:86:LYS:HG2	1.84	0.67
74:O8:8:ILE:HD12	74:O8:8:ILE:H	1.57	0.67
5:S3:34:TYR:HE2	5:S3:37:VAL:HG13	2.91	0.67
8:S6:57:ASP:HA	8:S6:107:ALA:H	1.59	0.67
36:1:1320:C:O2	56:N0:115:ARG:NH2	2.27	0.67
36:1:544:C:H1'	36:1:548:G:H22	1.58	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:837:A:OP2	79:Q3:4:ARG:NH1	2.28	0.67
38:8:92:A:H2'	38:8:93:U:O4'	1.94	0.67
13:C1:10:GLU:HG2	1:6:327:U:H1'	271.01	0.67
20:C8:125:ILE:HG12	35:SM:61:ILE:HG23	1.76	0.67
36:1:1155:C:OP1	44:L7:94:LYS:NZ	2.27	0.67
47:M0:43:VAL:HG21	47:M0:197:VAL:HB	1.77	0.67
63:N7:95:VAL:O	63:N7:100:THR:OG1	2.08	0.67
71:O5:85:THR:HB	71:O5:88:LEU:HD12	1.77	0.67
2:S0:32:HIS:O	2:S0:34:GLU:N	2.26	0.67
36:1:1724:U:H4'	36:1:1725:C:OP1	1.93	0.67
1:2:656:G:O2'	1:2:657:U:O4'	2.12	0.67
36:5:283:G:OP2	36:5:285:A:O2'	2.12	0.67
1:6:1305:U:OP2	1:6:1306:C:N4	2.26	0.67
1:6:1579:U:OP1	86:6:2182:OHX:N4	2.27	0.67
42:L5:129:TYR:OH	42:L5:175:HIS:O	2.11	0.67
42:L5:294:ALA:HB1	47:M0:217:PHE:HB3	1.76	0.67
50:M4:13:ARG:NH1	50:M4:65:LEU:O	2.28	0.67
59:N3:120:LYS:H	59:N3:137:VAL:HG23	1.60	0.67
62:N6:112:ASP:H	62:N6:115:ARG:HB2	2.07	0.67
11:S9:149:ARG:NE	1:6:765:G:N7	430.10	0.67
36:1:1230:G:H1	36:1:1279:C:H42	1.42	0.67
1:2:560:U:H2'	1:2:561:G:H8	1.59	0.67
37:3:27:A:P	42:L5:57:ASN:HD22	2.18	0.67
44:L7:90:LYS:NZ	36:5:1158:A:OP2	240.93	0.67
1:6:1003:A:O2'	1:6:1005:A:N6	2.18	0.67
1:6:1672:G:H2'	1:6:1673:G:C8	2.29	0.67
39:L2:6:ARG:NH2	39:L2:199:THR:O	2.26	0.67
41:L4:122:THR:HG22	41:L4:235:LEU:HB2	1.75	0.67
46:L9:86:TYR:CE2	46:L9:151:VAL:HG22	2.29	0.67
46:L9:49:ASN:ND2	46:L9:52:LEU:HB2	2.08	0.67
51:M5:8:GLU:HG3	51:M5:50:ARG:HH12	3.46	0.67
68:O2:22:SER:HA	68:O2:28:VAL:HG12	1.76	0.67
70:O4:74:ARG:HG2	70:O4:75:ALA:N	3.02	0.67
4:S2:45:VAL:HG21	4:S2:68:ILE:HG23	1.95	0.67
10:S8:141:ARG:NH2	1:6:196:G:N7	281.70	0.67
36:1:2623:G:H2'	36:1:2624:G:H8	1.60	0.67
36:1:371:G:O6	86:1:4177:OHX:N4	2.28	0.67
1:6:151:G:H1	1:6:163:G:H1	1.43	0.67
18:C6:66:ARG:HH21	18:C6:68:ARG:HG2	5.64	0.67
42:L5:43:LYS:O	42:L5:46:THR:OG1	2.11	0.67
36:1:75:G:H5''	49:M3:58:VAL:HG22	1.76	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
58:N2:94:ARG:HH21	58:N2:96:VAL:HG22	1.60	0.67
4:S2:116:LYS:NZ	4:S2:117:THR:O	3.69	0.67
37:3:75:G:O2'	37:3:104:A:N6	2.26	0.67
56:N0:115:ARG:NH2	36:5:1320:C:O2	288.79	0.67
36:5:784:A:O2'	36:5:785:G:OP2	2.12	0.67
16:C4:20:TYR:HB3	16:C4:27:PHE:HB2	1.76	0.67
1:2:57:G:OP1	26:D4:112:LYS:HE3	1.94	0.67
86:1:3951:OHX:N6	44:L7:217:PRO:O	2.28	0.67
50:M4:39:ILE:HB	50:M4:43:LYS:HB2	1.76	0.67
55:M9:86:GLU:OE2	55:M9:91:SER:OG	2.11	0.67
6:S4:247:SER:N	6:S4:250:GLU:OE1	2.21	0.67
36:1:3373:U:OP2	67:O1:102:LYS:HE2	1.95	0.67
1:6:755:A:H2'	1:6:756:A:C8	2.29	0.67
1:6:830:U:H2'	1:6:831:U:H5'	1.76	0.67
1:6:833:U:OP2	86:6:2202:OHX:N5	2.28	0.67
14:C2:81:ASP:O	14:C2:83:GLU:N	3.09	0.67
19:C7:7:LYS:N	1:6:1316:G:OP1	411.13	0.67
24:D2:53:ILE:HD13	29:D7:24:LEU:HD11	2.80	0.67
39:L2:227:ARG:HG2	39:L2:239:ALA:HB2	1.76	0.67
39:L2:29:LEU:HB2	39:L2:123:ARG:HA	1.77	0.67
52:M6:8:VAL:HG12	52:M6:117:ARG:HA	1.77	0.67
53:M7:155:GLU:N	53:M7:155:GLU:OE2	4.84	0.67
59:N3:108:GLU:HA	59:N3:128:ARG:HG3	1.75	0.67
64:N8:6:THR:HB	64:N8:8:THR:HG23	4.21	0.67
66:O0:66:LYS:H	66:O0:66:LYS:HD2	3.82	0.67
78:Q2:12:CYS:SG	78:Q2:17:CYS:HB3	2.80	0.67
6:S4:95:THR:O	6:S4:97:GLU:N	2.27	0.67
10:S8:57:ALA:HB2	10:S8:177:GLY:HA2	1.76	0.67
36:1:656:A:H2'	36:1:657:A:C8	2.30	0.67
36:1:729:C:H2'	36:1:730:C:H6	1.60	0.67
37:3:43:U:H4'	48:M1:140:ARG:O	1.94	0.67
1:6:158:U:O2'	1:6:160:C:OP2	2.13	0.67
29:D7:62:ILE:HG13	29:D7:63:LEU:H	1.60	0.67
44:L7:136:TYR:CZ	44:L7:231:ASN:HB2	2.30	0.67
44:L7:30:ARG:NH1	36:5:595:G:OP2	237.26	0.67
47:M0:68:ALA:HB2	47:M0:158:LYS:HG3	2.77	0.67
3:S1:137:ILE:HD11	3:S1:172:LEU:HB3	1.75	0.67
36:1:2683:U:H2'	36:1:2684:C:C6	2.30	0.66
36:5:3194:C:H2'	36:5:3195:U:H3'	1.75	0.66
1:6:1738:U:O4	86:6:2062:OHX:N5	2.28	0.66
15:C3:27:LYS:H	15:C3:27:LYS:HE3	1.59	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:C8:136:GLN:N	20:C8:136:GLN:OE1	2.27	0.66
41:L4:181:VAL:HG12	41:L4:182:LEU:H	1.60	0.66
36:1:3310:A:OP1	53:M7:74:LYS:NZ	2.27	0.66
64:N8:131:SER:HB3	64:N8:134:ALA:HB2	1.77	0.66
67:O1:55:LEU:HB2	67:O1:95:PRO:HD3	1.76	0.66
35:SM:34:LYS:HA	48:M1:61:ARG:HH12	1.60	0.66
36:1:1382:G:OP2	41:L4:188:ARG:NH1	2.26	0.66
36:1:2768:U:H2'	36:1:2769:A:C8	2.31	0.66
36:1:561:C:H2'	36:1:562:C:H6	1.60	0.66
1:2:1160:A:H2'	1:2:1161:C:C6	2.30	0.66
37:3:26:C:H5'	42:L5:56:THR:HB	1.76	0.66
36:5:2101:C:O2'	36:5:2102:U:OP1	2.12	0.66
36:5:1365:G:OP2	86:5:4023:OHX:N3	2.28	0.66
37:7:91:G:H2'	37:7:92:A:C8	2.29	0.66
13:C1:75:VAL:HG12	13:C1:120:GLY:H	1.60	0.66
19:C7:5:ARG:HB2	19:C7:10:LYS:HE2	2.17	0.66
21:C9:52:GLY:HA2	21:C9:55:TYR:HD2	1.60	0.66
27:D5:60:VAL:HG13	27:D5:101:TYR:HB2	3.08	0.66
39:L2:28:LYS:HD3	39:L2:123:ARG:HD3	1.75	0.66
40:L3:41:VAL:CA	40:L3:185:GLY:HA3	2.38	0.66
49:M3:46:ILE:HA	49:M3:49:ARG:HH11	4.25	0.66
36:1:3214:U:OP2	50:M4:128:ARG:NH2	2.27	0.66
51:M5:70:ASN:HB3	51:M5:92:LEU:O	1.95	0.66
57:N1:78:LYS:HE3	57:N1:87:LYS:HD2	1.76	0.66
7:S5:74:ALA:O	18:C6:122:ARG:NH2	2.28	0.66
36:1:2273:G:H22	36:1:2311:G:H2'	1.59	0.66
1:2:591:A:H2'	1:2:592:A:C8	2.30	0.66
15:C3:5:HIS:CE1	15:C3:121:ARG:HG3	2.30	0.66
26:D4:2:SER:N	26:D4:32:ARG:HG3	2.10	0.66
40:L3:229:VAL:HG11	40:L3:249:VAL:HG12	5.85	0.66
46:L9:163:GLN:HB3	46:L9:166:ARG:HH11	1.60	0.66
61:N5:105:VAL:HG11	61:N5:126:LEU:HD22	2.05	0.66
78:Q2:77:CYS:SG	78:Q2:79:THR:HG22	2.36	0.66
6:S4:230:GLU:HB2	6:S4:233:LYS:HE3	5.29	0.66
36:5:1070:U:O4	86:5:4106:OHX:N6	2.29	0.66
36:5:622:A:H2'	36:5:623:U:O4'	1.93	0.66
1:6:1734:U:H2'	1:6:1735:U:C6	2.30	0.66
5:S3:72:LEU:HG	12:C0:20:VAL:HG21	2.21	0.66
15:C3:93:LYS:HG3	15:C3:150:VAL:HG11	1.78	0.66
26:D4:36:SER:HB3	26:D4:39:GLU:HB2	1.76	0.66
41:L4:269:SER:O	41:L4:271:LYS:N	2.29	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:M5:65:ARG:HD2	51:M5:129:TYR:CE1	2.31	0.66
52:M6:27:LEU:HD11	52:M6:102:LEU:HB2	1.84	0.66
58:N2:50:LEU:O	58:N2:52:ASN:N	2.29	0.66
64:N8:3:SER:O	64:N8:6:THR:OG1	5.00	0.66
66:O0:13:LYS:HB3	66:O0:100:ILE:HG22	3.68	0.66
71:O5:66:VAL:HA	71:O5:69:LEU:HD12	1.75	0.66
72:O6:58:ILE:HG22	72:O6:90:MET:HG3	1.75	0.66
4:S2:80:VAL:HG13	4:S2:81:MET:H	1.60	0.66
36:1:900:G:H1'	36:1:1589:A:N6	2.11	0.66
36:1:3085:G:OP2	86:1:3879:OHX:N2	2.28	0.66
25:D3:8:GLY:O	25:D3:11:SER:OG	2.97	0.66
36:1:13:A:H5''	36:1:13:A:H8	1.60	0.66
70:O4:37:LYS:NZ	36:5:1591:G:OP1	160.80	0.66
25:D3:30:LYS:NZ	1:6:1132:A:OP1	322.07	0.66
1:6:1133:A:H2'	1:6:1134:C:O4'	1.95	0.66
1:6:658:C:N4	1:6:673:A:N1	2.44	0.66
13:C1:90:TYR:HD1	13:C1:91:LEU:N	1.93	0.66
40:L3:346:THR:HA	40:L3:351:LEU:HD21	3.18	0.66
42:L5:111:GLN:HA	42:L5:116:ASP:HB3	3.90	0.66
79:Q3:46:THR:HB	79:Q3:58:SER:HB2	1.78	0.66
3:S1:176:VAL:HG12	3:S1:177:GLN:H	1.59	0.66
34:SR:13:LEU:HB2	34:SR:310:ILE:HB	1.77	0.66
36:1:2252:A:H61	36:1:2264:U:H3	1.42	0.66
36:1:2683:U:H2'	36:1:2684:C:H6	1.61	0.66
36:1:2960:C:OP1	86:1:3996:OHX:N4	2.28	0.66
36:1:789:A:H2'	36:1:790:U:C6	2.30	0.66
1:2:1370:U:O4	86:2:2120:OHX:N1	2.28	0.66
36:5:3259:U:H5''	36:5:3261:C:H5	1.59	0.66
36:5:658:G:OP1	86:5:4087:OHX:N5	2.29	0.66
1:6:1395:G:O6	86:6:2088:OHX:N3	2.28	0.66
40:L3:100:ARG:NH2	36:5:3244:A:OP2	249.53	0.66
40:L3:183:LEU:O	40:L3:191:LYS:NZ	2.20	0.66
47:M0:81:GLY:O	47:M0:83:ASP:N	2.65	0.66
48:M1:133:ARG:HB3	48:M1:134:PRO:HD2	2.31	0.66
63:N7:3:LYS:HE3	66:O0:36:GLN:HG3	1.78	0.66
67:O1:44:MET:HB2	67:O1:46:THR:CG2	2.25	0.66
5:S3:29:LEU:HD21	5:S3:69:LEU:HD21	3.06	0.66
36:1:651:G:O2'	36:1:1435:A:OP1	2.11	0.66
36:1:155:G:H1'	72:O6:26:ILE:HD13	1.78	0.66
36:1:679:U:O4	86:1:3966:OHX:N1	2.29	0.66
86:2:2031:OHX:N6	86:2:2146:OHX:N5	2.43	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:386:G:P	10:S8:25:ARG:HH22	2.19	0.66
1:2:794:U:O2'	1:2:795:U:O2	2.13	0.66
36:5:1155:C:H2'	36:5:1156:C:H6	1.59	0.66
36:5:410:U:O4	86:5:4097:OHX:N1	2.28	0.66
13:C1:108:PRO:HG3	13:C1:134:THR:HB	2.17	0.66
20:C8:76:PRO:HG2	20:C8:86:LEU:HD21	1.77	0.66
39:L2:20:THR:O	39:L2:23:ARG:HG2	4.59	0.66
44:L7:127:LEU:O	44:L7:129:LEU:N	2.26	0.66
46:L9:4:ILE:HG23	46:L9:5:GLN:HB3	1.78	0.66
48:M1:90:GLN:HG2	48:M1:170:ASP:HB2	1.77	0.66
54:M8:63:SER:OG	54:M8:64:VAL:N	2.25	0.66
59:N3:10:LYS:HB2	59:N3:125:LEU:HD21	1.78	0.66
70:O4:46:ASP:OD2	70:O4:88:ARG:NH2	2.29	0.66
78:Q2:50:PHE:O	86:Q2:503:OHX:N2	2.29	0.66
5:S3:41:VAL:HA	5:S3:46:THR:HG23	2.86	0.66
34:SR:38:ARG:HA	34:SR:67:ILE:HA	1.77	0.66
36:1:1281:G:H2'	36:1:1282:G:H8	1.60	0.66
36:1:1786:G:H2'	36:1:1787:A:C8	2.30	0.66
86:2:2031:OHX:N3	86:2:2146:OHX:N5	2.44	0.66
68:O2:105:ARG:NH2	36:5:1412:G:OP1	145.70	0.66
1:6:882:U:H2'	1:6:883:C:C6	2.31	0.66
1:2:1173:C:OP1	20:C8:132:ARG:NH1	2.29	0.66
20:C8:139:LYS:O	20:C8:143:ARG:NH1	2.28	0.66
2:S0:154:GLU:HA	23:D1:63:GLY:HA2	1.76	0.66
25:D3:127:VAL:O	25:D3:129:GLY:N	2.29	0.66
27:D5:71:ILE:HG23	27:D5:73:GLY:H	6.76	0.66
52:M6:65:ASN:HB3	52:M6:68:ARG:HD3	1.96	0.66
2:S0:73:VAL:O	2:S0:95:ALA:HA	1.95	0.66
8:S6:132:ARG:HG2	8:S6:133:LEU:HD13	1.76	0.66
36:1:1278:A:O2'	36:1:1279:C:O5'	2.13	0.66
36:5:2402:A:OP2	86:5:4105:OHX:N3	2.29	0.66
46:L9:62:ARG:NH2	36:5:3115:C:OP1	329.77	0.66
1:6:961:U:H2'	1:6:962:C:C6	2.30	0.66
22:D0:46:GLU:O	22:D0:49:ASN:ND2	4.52	0.66
23:D1:73:ALA:HB3	23:D1:79:LEU:HD12	4.08	0.66
27:D5:55:PRO:O	27:D5:57:TYR:N	2.24	0.66
30:D8:11:LYS:HA	30:D8:53:ILE:HG12	4.82	0.66
36:1:1347:U:H4'	41:L4:305:ALA:HB2	1.78	0.66
36:1:976:U:OP1	54:M8:144:ARG:NH2	2.29	0.66
59:N3:2:SER:N	59:N3:57:MET:H	1.93	0.66
63:N7:78:ASN:OD1	66:O0:35:ARG:NH2	2.27	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:S3:222:VAL:HG23	34:SR:192:PHE:HA	2.68	0.66
5:S3:40:ARG:HG2	22:D0:110:PRO:HB3	1.78	0.66
36:1:1758:G:H1	36:1:1767:C:H42	1.42	0.65
36:1:2320:A:H2	79:Q3:16:VAL:HG12	1.59	0.65
1:2:651:G:N7	86:2:2103:OHX:N6	2.44	0.65
1:2:866:G:OP1	15:C3:2:GLY:HA2	1.97	0.65
36:5:604:G:N7	86:5:4163:OHX:N2	2.43	0.65
39:L2:21:ARG:NH1	36:5:825:U:OP1	172.40	0.65
39:L2:3:ARG:HD3	36:5:911:C:H42	179.22	0.65
1:6:116:U:H2'	1:6:117:U:H6	1.59	0.65
19:C7:8:THR:HG21	1:6:1330:G:H21	421.49	0.65
22:D0:63:LEU:HB3	31:D9:34:TYR:CE2	2.31	0.65
71:O5:89:ARG:HG2	71:O5:89:ARG:HH11	2.00	0.65
10:S8:188:GLU:OE2	13:C1:15:LYS:NZ	2.29	0.65
11:S9:38:ASN:HB3	11:S9:40:LYS:H	1.60	0.65
1:2:1015:U:OP1	86:2:2044:OHX:N3	2.28	0.65
1:2:959:U:OP1	29:D7:30:SER:OG	2.12	0.65
1:6:1255:G:O2'	1:6:1256:A:O5'	2.13	0.65
38:8:107:G:OP2	86:8:228:OHX:N1	2.29	0.65
16:C4:35:GLY:HA3	1:6:919:A:H5'	270.04	0.65
40:L3:292:ALA:HA	40:L3:303:LYS:O	1.95	0.65
44:L7:47:ARG:NH1	44:L7:183:ASP:OD2	2.28	0.65
47:M0:130:ASP:OD1	47:M0:131:ILE:N	2.82	0.65
36:1:670:C:OP1	54:M8:147:ARG:NH2	2.28	0.65
1:2:154:G:OP1	8:S6:2:LYS:NZ	2.27	0.65
9:S7:50:ASP:N	9:S7:50:ASP:OD1	2.28	0.65
34:SR:161:LYS:HB3	34:SR:164:ASP:HB3	1.78	0.65
36:1:1554:U:HO2'	36:1:1582:C:H5	1.44	0.65
36:1:2177:G:O6	86:1:3917:OHX:N2	2.30	0.65
36:1:2356:A:H61	36:1:2983:C:H5	1.43	0.65
36:1:3224:G:O6	86:1:3886:OHX:N4	2.28	0.65
36:5:1938:U:O4	86:5:3943:OHX:N1	2.29	0.65
36:5:679:U:O4	86:5:4008:OHX:N2	2.29	0.65
1:6:828:U:H2'	1:6:829:A:H5''	1.77	0.65
42:L5:265:TYR:HE1	37:7:121:U:H5''	315.70	0.65
22:D0:24:ILE:HG12	22:D0:116:VAL:HG12	3.18	0.65
28:D6:32:LYS:NZ	1:6:932:U:O2	310.83	0.65
36:1:2393:G:H4'	40:L3:252:ILE:HG12	1.77	0.65
40:L3:332:ARG:NH1	40:L3:333:LYS:HD2	3.83	0.65
45:L8:24:ASN:HB3	45:L8:25:PRO:HD2	3.50	0.65
47:M0:89:VAL:HG22	47:M0:136:PHE:CE1	2.48	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:M5:58:GLY:HA3	51:M5:142:ILE:HD11	1.78	0.65
54:M8:83:VAL:O	54:M8:85:GLY:N	2.54	0.65
62:N6:73:VAL:HA	62:N6:80:VAL:HG23	2.10	0.65
63:N7:83:THR:HG23	63:N7:85:TYR:N	2.11	0.65
3:S1:135:LEU:HB3	3:S1:217:LEU:HD12	3.86	0.65
8:S6:28:PHE:CZ	8:S6:104:PRO:HB3	3.01	0.65
19:C7:23:LYS:H	34:SR:216:LYS:HE2	1.61	0.65
36:1:1817:G:OP1	86:1:4086:OHX:N1	2.29	0.65
1:6:1745:G:O6	86:6:2077:OHX:N4	2.28	0.65
21:C9:105:LEU:HD22	21:C9:122:ARG:HG2	1.78	0.65
42:L5:261:THR:OG1	42:L5:264:GLN:HG3	1.96	0.65
44:L7:152:GLY:O	44:L7:163:LEU:HG	1.96	0.65
49:M3:48:PRO:HA	49:M3:137:GLN:HB3	1.79	0.65
51:M5:13:LYS:O	51:M5:16:SER:OG	2.12	0.65
52:M6:27:LEU:HD13	52:M6:98:ALA:O	1.97	0.65
54:M8:30:VAL:O	54:M8:34:THR:HG23	4.75	0.65
9:S7:50:ASP:HB3	9:S7:56:LYS:HG2	1.77	0.65
1:2:1600:A:H4'	1:2:1601:G:OP1	1.96	0.65
1:2:1796:C:OP1	28:D6:87:ARG:NH1	2.28	0.65
1:2:768:C:H1'	11:S9:143:ILE:HG21	1.77	0.65
44:L7:206:LYS:HB3	36:5:1334:U:H5''	235.75	0.65
36:5:2444:C:H42	36:5:2503:G:H1	1.45	0.65
36:5:419:G:N7	86:5:3898:OHX:N3	2.45	0.65
1:6:1681:A:H2'	1:6:1682:U:H5'	1.78	0.65
13:C1:95:PRO:O	13:C1:97:TYR:N	2.29	0.65
18:C6:50:GLU:OE1	18:C6:112:TYR:OH	2.11	0.65
22:D0:36:ASN:HA	22:D0:39:SER:HB2	4.68	0.65
39:L2:221:LYS:O	36:5:2245:C:H4'	218.78	0.65
42:L5:211:LEU:HB3	42:L5:219:PHE:HB2	2.21	0.65
49:M3:87:ALA:O	49:M3:90:ALA:N	2.29	0.65
52:M6:18:ARG:O	52:M6:22:VAL:HG12	1.97	0.65
53:M7:127:ARG:NH2	36:5:1508:C:OP1	138.09	0.65
36:1:743:C:O2	54:M8:141:ARG:HD3	1.97	0.65
57:N1:130:ARG:NH1	36:5:1098:A:OP2	253.59	0.65
66:O0:20:SER:OG	66:O0:96:GLY:HA3	1.95	0.65
6:S4:180:LEU:O	6:S4:228:ILE:N	2.28	0.65
10:S8:159:GLN:HE22	10:S8:166:TYR:HB2	3.14	0.65
10:S8:76:THR:HG22	10:S8:108:PRO:HG2	1.78	0.65
36:1:118:U:O2	36:1:121:A:H5'	1.96	0.65
36:1:1493:G:O6	75:O9:2:ALA:HB2	1.96	0.65
36:1:372:A:H2'	36:1:373:A:C8	2.31	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:1780:G:OP1	86:1:4048:OHX:N1	2.29	0.65
37:3:112:G:OP2	86:3:221:OHX:N1	2.30	0.65
36:5:1387:G:OP1	86:5:4195:OHX:N3	2.29	0.65
36:5:2666:C:H2'	36:5:2667:A:H5''	1.77	0.65
36:5:3159:C:H2'	36:5:3160:U:C6	2.31	0.65
15:C3:114:ARG:HD3	15:C3:117:LEU:HD12	2.58	0.65
20:C8:28:ILE:HG12	20:C8:61:LEU:HD11	4.93	0.65
21:C9:27:LYS:HB3	21:C9:111:ILE:HD11	1.78	0.65
28:D6:84:VAL:HG13	28:D6:85:ARG:H	1.62	0.65
46:L9:111:PHE:HD1	46:L9:127:PRO:HA	2.05	0.65
56:N0:9:VAL:HG22	56:N0:61:ILE:HG12	1.78	0.65
71:O5:68:GLN:HA	71:O5:71:LYS:HB2	1.78	0.65
3:S1:61:LEU:HB2	3:S1:64:ARG:HE	1.60	0.65
6:S4:100:ARG:NH2	6:S4:121:TYR:O	2.29	0.65
1:2:187:G:OP2	10:S8:142:LYS:NZ	2.27	0.65
36:1:148:G:OP2	51:M5:4:TYR:OH	2.11	0.65
36:1:1492:G:O2'	75:O9:48:LYS:NZ	2.29	0.65
36:1:2660:G:OP1	36:1:2750:U:O2'	2.15	0.65
1:2:628:G:N2	1:2:970:A:OP2	2.26	0.65
36:5:1409:G:O6	86:5:4157:OHX:N6	2.30	0.65
26:D4:37:LYS:NZ	1:6:523:G:OP2	413.72	0.65
1:2:1796:C:H5	28:D6:6:ALA:H	1.41	0.65
53:M7:29:THR:HA	53:M7:32:THR:HG23	1.78	0.65
36:1:1295:G:O2'	56:N0:115:ARG:HD3	1.97	0.65
56:N0:137:ARG:HG2	56:N0:139:TYR:CZ	2.54	0.65
59:N3:28:ASN:HD21	59:N3:112:SER:HB2	1.62	0.65
2:S0:120:LEU:HD21	2:S0:144:ILE:HD11	1.78	0.65
9:S7:38:LEU:HD23	9:S7:41:LEU:HD12	1.79	0.65
34:SR:29:GLN:HB2	34:SR:32:LEU:HB3	3.53	0.65
36:1:1276:U:OP1	86:1:4081:OHX:N4	2.30	0.65
36:1:54:C:O2'	36:1:1547:G:H1'	1.97	0.65
36:1:2538:U:O2'	36:1:2541:U:O4	2.10	0.65
36:1:3326:G:H2'	36:1:3327:G:H8	1.62	0.65
1:2:1535:U:O2'	1:2:1536:G:N3	2.28	0.65
1:2:397:A:O3'	10:S8:50:GLY:HA2	1.96	0.65
14:C2:119:SER:OG	1:6:1228:G:OP1	465.26	0.65
1:6:1665:U:O4	86:6:2123:OHX:N6	2.29	0.65
1:6:496:G:O6	1:6:497:G:N2	2.25	0.65
15:C3:26:PHE:CE2	15:C3:28:LEU:HB2	2.68	0.65
21:C9:16:ASN:OD1	21:C9:56:LYS:NZ	2.85	0.65
24:D2:67:GLY:O	24:D2:68:ARG:HG2	4.65	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:E0:55:ARG:HB3	32:E0:58:PRO:HG3	1.77	0.65
39:L2:201:GLY:O	39:L2:204:MET:HG2	1.96	0.65
41:L4:302:ALA:HB2	54:M8:39:ARG:NH2	2.12	0.65
51:M5:96:ARG:NH2	51:M5:104:GLU:OE1	4.71	0.65
55:M9:101:VAL:HA	55:M9:104:ARG:CZ	2.27	0.65
55:M9:21:LYS:O	55:M9:53:LYS:HB2	1.97	0.65
36:1:3070:A:OP1	55:M9:62:ARG:NH2	2.29	0.65
3:S1:154:SER:O	3:S1:154:SER:OG	2.43	0.65
2:S0:119:ARG:NH1	4:S2:241:ASP:OD1	3.14	0.65
8:S6:2:LYS:HB2	8:S6:108:VAL:HG22	1.78	0.65
34:SR:200:ASN:H	34:SR:215:GLY:HA2	1.62	0.65
36:1:1381:A:OP1	41:L4:197:ARG:NH1	2.29	0.65
36:1:789:A:H2'	36:1:790:U:H6	1.62	0.65
1:2:1796:C:H42	28:D6:93:LYS:HE2	1.61	0.65
1:2:66:U:H5'	8:S6:173:PRO:HA	1.79	0.65
36:5:900:G:H1'	36:5:1589:A:N6	2.11	0.65
10:S8:178:ARG:NH1	1:6:207:U:O2	288.42	0.65
1:6:899:G:N2	1:6:910:C:O2	2.18	0.65
18:C6:55:VAL:HG11	18:C6:89:LEU:HD21	3.13	0.65
25:D3:59:ILE:HG12	32:E0:4:VAL:HG22	3.93	0.65
40:L3:187:SER:O	40:L3:190:GLU:N	2.61	0.65
86:1:3951:OHX:N3	44:L7:217:PRO:O	2.30	0.65
4:S2:106:ASP:OD1	4:S2:108:ASN:N	2.28	0.65
36:1:624:G:OP2	86:1:4128:OHX:N3	2.30	0.65
36:1:979:U:H1'	36:1:980:A:C8	2.32	0.65
1:2:649:U:O2'	1:2:650:U:O5'	2.15	0.65
36:5:1578:C:H3'	36:5:1579:C:C6	2.32	0.65
36:5:2761:G:O6	36:5:2796:G:H5''	1.97	0.65
36:5:796:U:H2'	36:5:797:U:H6	1.61	0.65
1:6:800:U:H2'	1:6:801:G:C8	2.32	0.65
22:D0:71:PRO:O	22:D0:72:ASN:ND2	5.66	0.65
27:D5:66:VAL:HG22	27:D5:71:ILE:HG22	4.84	0.65
39:L2:144:ASN:ND2	39:L2:161:ASP:OD1	3.89	0.65
42:L5:68:THR:HG22	42:L5:69:ILE:H	2.37	0.65
3:S1:126:THR:HG22	3:S1:136:ARG:HE	1.61	0.65
3:S1:61:LEU:HG	3:S1:64:ARG:NH2	2.12	0.65
11:S9:102:GLU:CD	11:S9:102:GLU:H	2.52	0.65
36:1:1597:C:H42	36:1:1610:G:H1	1.43	0.64
36:1:2278:C:OP1	86:1:3950:OHX:N3	2.30	0.64
36:1:2897:A:H2'	36:1:2899:C:H5''	1.79	0.64
36:1:3246:G:O6	86:1:4103:OHX:N4	2.30	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:L4:141:ARG:NH2	36:5:1385:C:OP1	126.46	0.64
36:5:3153:U:H4'	36:5:3154:C:H5'	1.77	0.64
15:C3:107:LYS:NZ	1:6:1019:A:OP2	267.29	0.64
3:S1:77:GLU:HG3	16:C4:114:ARG:HH22	1.62	0.64
17:C5:48:GLY:O	17:C5:50:THR:N	3.33	0.64
25:D3:57:LEU:HD22	32:E0:4:VAL:HG13	4.95	0.64
41:L4:26:PHE:HA	41:L4:127:ALA:HA	2.05	0.64
41:L4:57:GLY:HA3	41:L4:98:ARG:HB2	1.79	0.64
50:M4:47:ASP:OD2	50:M4:48:GLY:N	3.01	0.64
70:O4:105:VAL:HA	70:O4:108:GLN:HB2	3.59	0.64
34:SR:140:CYS:SG	34:SR:141:LEU:N	2.70	0.64
36:1:1688:U:H2'	36:1:1689:U:C6	2.32	0.64
36:1:2897:A:OP2	76:Q0:124:LYS:NZ	2.30	0.64
1:2:1230:A:H2'	1:2:1258:U:H5	1.62	0.64
86:2:2031:OHX:N4	86:2:2146:OHX:N1	2.45	0.64
36:5:1025:A:H3'	36:5:1026:A:H4'	1.78	0.64
36:5:655:C:H2'	36:5:656:A:C8	2.32	0.64
41:L4:292:SER:OG	41:L4:293:SER:N	2.30	0.64
49:M3:126:PHE:O	71:O5:114:ARG:NH2	2.30	0.64
2:S0:74:VAL:HA	2:S0:96:THR:O	3.79	0.64
3:S1:81:PHE:HD2	3:S1:82:ARG:H	3.30	0.64
1:2:1290:U:OP1	4:S2:95:ARG:NH1	2.30	0.64
36:1:1667:A:H2'	36:1:1668:G:C8	2.32	0.64
36:1:1821:U:C4	70:O4:67:LYS:HD2	2.33	0.64
36:1:249:U:H1'	36:1:250:U:O2	1.97	0.64
36:1:85:A:O2'	86:1:4137:OHX:N6	2.30	0.64
1:2:1290:U:H2'	1:2:1291:G:C8	2.33	0.64
1:2:329:G:H5''	10:S8:98:LYS:HB3	1.79	0.64
38:4:141:C:OP1	51:M5:109:ARG:NH1	2.28	0.64
36:5:1024:G:N7	36:5:1027:A:N6	2.46	0.64
1:6:1757:G:O6	86:6:2046:OHX:N4	2.31	0.64
28:D6:38:ARG:HH21	28:D6:83:ILE:HG13	1.63	0.64
45:L8:143:ILE:HD11	45:L8:151:VAL:HG21	1.79	0.64
48:M1:96:PHE:HB2	48:M1:156:LYS:HE3	1.78	0.64
56:N0:89:ASN:ND2	57:N1:156:TYR:HB3	2.12	0.64
59:N3:33:ASN:HD21	59:N3:63:LYS:H	1.46	0.64
61:N5:82:LEU:HD11	61:N5:135:ILE:HG21	1.79	0.64
1:2:123:G:H21	6:S4:146:THR:HG21	1.62	0.64
6:S4:163:ASP:O	6:S4:165:ALA:N	2.29	0.64
36:1:1487:G:H1	36:1:1855:U:H3	1.42	0.64
1:2:1720:G:O6	86:2:2081:OHX:N5	2.31	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:775:G:N7	26:D4:11:LYS:NZ	2.45	0.64
37:3:91:G:H2'	37:3:92:A:C8	2.33	0.64
55:M9:62:ARG:NH2	36:5:3068:U:OP2	173.30	0.64
1:6:805:U:H2'	1:6:806:A:H5'	1.80	0.64
16:C4:121:VAL:O	1:6:886:U:O2'	287.88	0.64
16:C4:81:VAL:HG22	16:C4:115:ILE:HB	1.79	0.64
1:2:1550:A:P	17:C5:42:ARG:HH22	2.21	0.64
18:C6:82:ARG:HH22	18:C6:114:ARG:HB3	1.61	0.64
22:D0:33:GLN:N	22:D0:33:GLN:OE1	2.51	0.64
29:D7:61:THR:HG23	29:D7:62:ILE:H	1.62	0.64
39:L2:117:GLU:HG2	39:L2:122:ASP:H	1.62	0.64
48:M1:28:ASP:OD2	48:M1:32:ARG:NH2	8.02	0.64
76:Q0:106:ARG:HB2	76:Q0:106:ARG:NH1	4.31	0.64
7:S5:73:THR:C	7:S5:75:GLY:H	2.58	0.64
8:S6:177:ARG:NH2	1:6:143:G:N7	312.75	0.64
11:S9:108:ARG:HB3	11:S9:110:GLN:HB3	2.34	0.64
36:1:1472:U:H2'	36:1:1473:G:C8	2.32	0.64
36:1:3151:U:OP2	40:L3:132:LYS:NZ	2.31	0.64
36:1:409:A:OP2	86:1:4051:OHX:N5	2.31	0.64
36:1:855:U:H2'	36:1:856:G:O4'	1.98	0.64
1:2:1114:G:O6	86:2:2073:OHX:N5	2.30	0.64
36:5:3065:G:O6	86:5:4100:OHX:N6	2.30	0.64
1:6:86:A:OP2	86:6:2188:OHX:N1	2.31	0.64
45:L8:81:THR:HG21	45:L8:181:LYS:HD3	1.79	0.64
46:L9:88:TYR:CZ	46:L9:184:LYS:HD3	3.83	0.64
57:N1:122:GLN:HB3	57:N1:124:VAL:HG23	7.13	0.64
8:S6:20:ASP:HB3	8:S6:23:ARG:HB2	1.79	0.64
8:S6:48:TYR:CZ	8:S6:121:LEU:HD22	4.80	0.64
36:5:3085:G:OP2	86:5:3899:OHX:N1	2.31	0.64
33:E1:97:LYS:NZ	1:6:1253:U:O4	439.97	0.64
17:C5:119:PHE:HE1	20:C8:119:ILE:HG23	1.63	0.64
22:D0:44:ASN:HA	22:D0:47:GLN:HB3	2.81	0.64
2:S0:66:ALA:HB1	23:D1:50:TYR:HD1	2.96	0.64
40:L3:171:LEU:O	86:L3:405:OHX:N6	2.31	0.64
46:L9:138:THR:O	46:L9:139:ASN:ND2	2.29	0.64
46:L9:77:ASN:HB3	46:L9:151:VAL:HG21	1.80	0.64
47:M0:99:ILE:HG13	47:M0:100:ASN:N	2.13	0.64
4:S2:69:ILE:HG12	4:S2:133:LYS:HB3	1.80	0.64
36:1:1221:A:H3'	36:1:1222:G:H5'	1.80	0.64
36:1:1783:U:H2'	36:1:1784:G:C8	2.33	0.64
1:2:7:G:O6	4:S2:205:ARG:NH2	2.31	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:L7:105:LEU:HD22	36:5:1101:G:H1'	234.51	0.64
36:5:1808:G:O6	86:5:4018:OHX:N3	2.31	0.64
36:5:2526:C:H1'	36:5:2588:U:H5''	1.79	0.64
36:5:2822:U:OP2	86:5:3946:OHX:N1	2.30	0.64
52:M6:168:TYR:OH	36:5:3190:C:OP1	301.44	0.64
12:C0:2:LEU:HD22	1:6:1258:U:H4'	433.79	0.64
17:C5:123:TYR:H	17:C5:123:TYR:HD1	2.47	0.64
19:C7:34:LEU:HD13	19:C7:38:ILE:HD12	6.91	0.64
39:L2:118:GLU:HG3	39:L2:126:LEU:HD21	2.78	0.64
54:M8:54:LEU:HD13	54:M8:58:ASN:HB3	1.80	0.64
55:M9:11:ALA:O	55:M9:15:VAL:HG23	1.97	0.64
63:N7:33:SER:OG	63:N7:34:LYS:N	2.85	0.64
78:Q2:10:THR:OG1	78:Q2:11:TYR:N	2.30	0.64
2:S0:9:LEU:HD11	2:S0:14:ALA:HB2	1.80	0.64
34:SR:220:ILE:HB	34:SR:234:LEU:HB2	2.18	0.64
36:1:269:G:H5'	51:M5:120:TRP:CE3	2.33	0.64
36:1:70:A:N1	36:1:313:A:O2'	2.24	0.64
1:2:480:G:N2	1:2:509:G:H1'	2.13	0.64
37:3:5:G:OP1	48:M1:143:ARG:NH2	2.27	0.64
44:L7:70:LYS:NZ	36:5:519:A:OP2	314.37	0.64
36:5:789:A:H2'	36:5:790:U:H6	1.63	0.64
24:D2:32:LYS:HG3	1:6:637:C:OP1	364.37	0.64
39:L2:49:VAL:HG22	39:L2:50:HIS:H	2.03	0.64
52:M6:42:ASN:OD1	52:M6:125:ARG:HD3	2.12	0.64
57:N1:45:ASN:OD1	57:N1:47:SER:OG	2.12	0.64
36:1:2630:C:C5	57:N1:4:SER:HB2	2.33	0.64
67:O1:17:HIS:CG	67:O1:69:TYR:HD1	2.16	0.64
76:Q0:99:CYS:HB3	76:Q0:114:LYS:HD3	4.03	0.64
4:S2:142:GLY:HA2	4:S2:151:PRO:HB3	1.80	0.64
5:S3:164:VAL:HG13	5:S3:168:ILE:HD11	1.80	0.64
6:S4:160:VAL:HG13	6:S4:169:ILE:HG23	2.64	0.64
10:S8:9:HIS:HD2	10:S8:10:LYS:HB2	1.62	0.64
36:1:2898:G:H5''	36:1:2899:C:C5'	2.28	0.64
1:2:1241:G:H1'	17:C5:79:HIS:CG	2.32	0.64
1:2:1274:C:H5	35:SM:96:ARG:H	1.45	0.64
1:2:1795:U:H4'	28:D6:84:VAL:HG23	1.79	0.64
65:N9:50:THR:HB	36:5:1073:U:H1'	205.42	0.64
39:L2:207:VAL:HG11	36:5:916:G:C6	185.56	0.64
1:6:218:A:H61	1:6:829:A:H2	1.45	0.64
1:2:246:G:H1'	13:C1:40:LEU:HD13	1.80	0.64
27:D5:59:TYR:HE2	27:D5:61:SER:HB3	1.62	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:L2:174:ARG:NH2	36:5:2179:C:O2'	214.72	0.64
36:1:2202:C:O2'	39:L2:240:ALA:O	2.13	0.64
39:L2:54:ARG:HG2	39:L2:56:ALA:H	1.62	0.64
52:M6:102:LEU:HD12	52:M6:103:LYS:N	2.12	0.64
67:O1:13:THR:HG22	67:O1:72:ARG:HH21	5.10	0.64
2:S0:21:ASN:O	2:S0:163:ASN:ND2	2.30	0.64
2:S0:69:ASN:HB3	2:S0:71:GLU:HG2	1.80	0.64
3:S1:109:LYS:HG3	3:S1:113:MET:HE3	1.78	0.64
36:1:12:A:OP1	86:1:4200:OHX:N6	2.31	0.64
1:6:1316:G:HO2'	1:6:1401:A:HO2'	1.46	0.64
1:6:1696:G:N2	1:6:1704:U:H3	1.96	0.64
1:6:246:G:H2'	1:6:247:A:C8	2.33	0.64
49:M3:132:ALA:O	49:M3:134:GLU:N	3.26	0.64
52:M6:76:PRO:HB3	52:M6:138:LEU:HD23	1.80	0.64
68:O2:81:ASP:O	68:O2:84:THR:HG23	1.98	0.64
9:S7:73:VAL:O	9:S7:75:THR:N	2.54	0.64
11:S9:62:ARG:O	11:S9:69:ARG:NH1	2.31	0.64
36:1:1033:U:H2'	36:1:1034:U:C6	2.34	0.63
36:1:2416:U:H2'	36:1:2417:U:C6	2.33	0.63
36:1:2734:A:OP1	86:1:4001:OHX:N3	2.30	0.63
36:1:3189:G:H2'	36:1:3190:C:O4'	1.98	0.63
1:2:1672:G:H2'	1:2:1673:G:C8	2.33	0.63
1:2:301:A:OP2	86:2:2063:OHX:N2	2.31	0.63
36:5:1501:U:H3	36:5:1515:A:H61	1.43	0.63
36:5:956:U:H2'	36:5:957:C:H6	1.62	0.63
39:L2:79:ASN:ND2	39:L2:165:VAL:HG22	2.36	0.63
40:L3:250:ALA:HB3	36:5:2880:U:H1'	224.01	0.63
41:L4:300:ARG:HH11	41:L4:300:ARG:HB2	1.63	0.63
42:L5:55:PHE:CZ	42:L5:158:ARG:HG3	2.33	0.63
47:M0:95:HIS:CD2	47:M0:128:ARG:HE	2.16	0.63
52:M6:85:ARG:HD3	52:M6:90:HIS:CG	2.32	0.63
56:N0:1:MET:HE1	56:N0:32:SER:N	2.13	0.63
43:L6:31:ARG:NH1	69:O3:107:ILE:HG22	5.79	0.63
6:S4:103:TYR:O	6:S4:182:TYR:OH	2.16	0.63
9:S7:14:THR:OG1	9:S7:15:GLU:N	2.31	0.63
11:S9:92:LYS:HB2	11:S9:95:TYR:HD2	9.55	0.63
36:1:1563:C:O2	36:1:1577:G:N2	2.20	0.63
36:1:2534:G:O6	86:1:3991:OHX:N6	2.31	0.63
36:1:1196:C:O2	86:1:3988:OHX:N2	2.31	0.63
1:2:1273:G:HO2'	1:2:1430:U:H5	1.45	0.63
36:5:507:U:H2'	36:5:508:U:C6	2.33	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:841:A:H2'	36:5:842:G:C8	2.32	0.63
36:5:975:C:H2'	36:5:976:U:H6	1.64	0.63
77:Q1:15:ARG:NH1	1:6:1126:G:OP1	279.81	0.63
1:6:180:A:H2'	1:6:181:A:O4'	1.98	0.63
71:O5:83:LYS:HA	38:8:38:U:H5	65.77	0.63
13:C1:99:ARG:HB3	25:D3:9:LEU:O	1.97	0.63
15:C3:2:GLY:N	1:6:866:G:OP1	334.75	0.63
16:C4:111:ARG:NH2	28:D6:57:SER:O	2.31	0.63
18:C6:39:VAL:HG12	18:C6:41:PRO:HD2	6.42	0.63
18:C6:50:GLU:OE1	18:C6:114:ARG:NH1	2.29	0.63
25:D3:22:ASN:OD1	1:6:1108:G:N1	334.00	0.63
26:D4:29:HIS:O	26:D4:31:ASN:N	3.63	0.63
61:N5:103:TYR:HE1	61:N5:139:ILE:HD12	1.63	0.63
63:N7:88:ASP:HB3	63:N7:121:ARG:HH22	1.63	0.63
71:O5:38:ARG:HG3	71:O5:39:PRO:HD2	2.95	0.63
74:O8:46:ARG:HH21	74:O8:51:LEU:HB2	1.62	0.63
3:S1:130:SER:OG	3:S1:131:ASP:N	2.24	0.63
4:S2:39:THR:OG1	4:S2:65:GLU:OE2	2.16	0.63
5:S3:64:ARG:NH1	5:S3:68:GLU:OE1	3.46	0.63
6:S4:200:ARG:NH2	6:S4:202:ASP:OD1	2.31	0.63
9:S7:35:LYS:O	9:S7:37:GLU:N	2.31	0.63
35:SM:64:LYS:O	35:SM:65:THR:OG1	2.11	0.63
36:1:1014:U:H2'	36:1:1015:U:H5''	1.79	0.63
1:6:463:U:H2'	1:6:464:A:H8	1.64	0.63
1:6:886:U:H2'	1:6:887:A:H8	1.63	0.63
27:D5:54:VAL:HA	27:D5:57:TYR:CE1	2.66	0.63
43:L6:153:PRO:O	43:L6:154:LEU:HB2	2.25	0.63
44:L7:142:SER:O	44:L7:146:GLN:HG3	2.24	0.63
52:M6:14:HIS:HE1	52:M6:119:VAL:HG12	1.64	0.63
56:N0:73:LYS:NZ	56:N0:97:VAL:O	3.73	0.63
11:S9:118:LEU:HD23	11:S9:158:PHE:CE1	4.84	0.63
11:S9:163:PRO:O	11:S9:165:GLY:N	2.31	0.63
11:S9:21:SER:HA	11:S9:24:LEU:HD12	2.59	0.63
36:5:246:U:H2'	36:5:247:C:H5''	1.80	0.63
36:5:2507:C:O2'	36:5:2508:U:OP1	2.13	0.63
36:5:3074:G:OP1	86:5:4114:OHX:N4	2.31	0.63
36:5:3165:A:H61	36:5:3285:C:N4	1.95	0.63
36:5:3348:G:H1	36:5:3357:U:H3	1.46	0.63
1:6:1230:A:H2'	1:6:1258:U:H5	1.63	0.63
1:6:887:A:H2'	1:6:888:U:C6	2.34	0.63
1:2:1173:C:H3'	20:C8:141:THR:HG21	1.81	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:L3:25:ILE:HD11	40:L3:334:ARG:HE	7.16	0.63
45:L8:171:LYS:NZ	45:L8:223:ALA:O	2.60	0.63
53:M7:29:THR:O	53:M7:32:THR:N	2.32	0.63
53:M7:59:PRO:HG3	53:M7:76:PHE:CD1	2.34	0.63
73:O7:81:GLY:O	38:8:95:G:H1'	40.85	0.63
74:O8:58:ASP:OD2	74:O8:61:LYS:N	2.24	0.63
3:S1:157:GLN:HB2	3:S1:160:HIS:CE1	2.48	0.63
3:S1:181:LEU:O	3:S1:184:LEU:N	2.30	0.63
36:1:2737:C:OP1	57:N1:69:LYS:HB3	1.99	0.63
36:1:2970:C:H4'	36:1:2971:A:C6	2.33	0.63
36:1:656:A:H2'	36:1:657:A:H8	1.64	0.63
1:2:577:G:OP2	35:SM:105:LYS:NZ	2.31	0.63
36:5:1627:U:O2'	36:5:1630:U:O2	2.11	0.63
36:5:979:U:H1'	36:5:980:A:C4	2.33	0.63
1:6:890:C:H2'	1:6:891:A:C8	2.33	0.63
13:C1:139:VAL:O	13:C1:140:VAL:HB	1.97	0.63
15:C3:22:ALA:HB1	15:C3:23:PRO:HA	2.17	0.63
17:C5:48:GLY:O	17:C5:52:LYS:HD3	1.98	0.63
21:C9:33:TYR:CD1	21:C9:37:VAL:HG21	4.43	0.63
40:L3:146:ARG:NE	40:L3:146:ARG:HA	2.13	0.63
40:L3:152:LYS:HE3	40:L3:192:VAL:HG22	1.80	0.63
43:L6:30:LEU:HD22	43:L6:34:LEU:HD12	5.38	0.63
43:L6:5:LYS:HG3	43:L6:6:ALA:H	4.72	0.63
52:M6:18:ARG:NH2	36:5:1318:A:OP1	276.60	0.63
52:M6:39:GLU:OE1	52:M6:39:GLU:N	2.59	0.63
74:O8:14:LEU:HA	74:O8:17:ARG:HG3	1.79	0.63
79:Q3:59:CYS:SG	79:Q3:60:CYS:N	3.45	0.63
5:S3:7:LYS:HE3	22:D0:27:THR:HG21	1.79	0.63
7:S5:51:VAL:O	7:S5:65:ARG:NH2	2.30	0.63
10:S8:98:LYS:HB2	10:S8:172:ARG:HG3	2.67	0.63
36:1:239:G:O6	86:1:4029:OHX:N3	2.32	0.63
36:5:2344:U:H2'	36:5:2345:A:C8	2.34	0.63
36:5:637:C:H1'	36:5:638:C:C6	2.32	0.63
17:C5:122:THR:HB	1:6:1558:U:H3	366.36	0.63
1:6:168:A:H2'	1:6:169:A:C8	2.34	0.63
1:6:1696:G:H2'	1:6:1698:G:O6	1.99	0.63
17:C5:20:VAL:HG12	17:C5:24:LYS:HD2	1.80	0.63
21:C9:4:VAL:HG11	21:C9:137:ALA:HB2	1.79	0.63
28:D6:38:ARG:NH2	28:D6:83:ILE:HG21	2.14	0.63
7:S5:57:SER:HA	30:D8:53:ILE:HD13	1.81	0.63
39:L2:79:ASN:HD22	39:L2:165:VAL:HG22	2.23	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:L2:70:ARG:CZ	39:L2:72:ARG:HE	5.27	0.63
42:L5:270:LYS:O	42:L5:273:ARG:HB3	2.97	0.63
52:M6:127:LEU:HD23	56:N0:156:VAL:HG12	1.79	0.63
53:M7:24:VAL:HG12	53:M7:86:LYS:HD3	2.52	0.63
36:1:970:A:OP1	65:N9:18:ARG:NH1	2.30	0.63
65:N9:51:ALA:O	65:N9:54:LEU:N	3.05	0.63
2:S0:110:TYR:HE2	4:S2:64:LYS:HB3	1.63	0.63
3:S1:137:ILE:HG13	3:S1:172:LEU:HD13	1.81	0.63
5:S3:43:PRO:O	5:S3:44:THR:HB	1.98	0.63
36:1:2155:G:O2'	39:L2:227:ARG:NH2	2.32	0.63
1:2:1591:C:H2'	1:2:1592:A:H8	1.64	0.63
1:2:205:U:O4	86:2:2066:OHX:N3	2.31	0.63
1:2:252:U:H2'	1:2:253:A:H8	1.64	0.63
36:5:1706:C:H2'	36:5:1707:A:O4'	1.99	0.63
36:5:2404:A:H2'	36:5:2405:C:H5'	1.79	0.63
1:6:1161:C:H2'	1:6:1162:C:H6	1.64	0.63
15:C3:128:TYR:OH	1:6:964:U:OP1	325.34	0.63
26:D4:120:GLY:O	26:D4:122:GLY:N	4.13	0.63
39:L2:83:HIS:CE1	39:L2:86:GLN:HB2	2.33	0.63
41:L4:341:SER:O	41:L4:342:LYS:HB3	4.64	0.63
45:L8:224:ASP:OD1	45:L8:224:ASP:N	2.68	0.63
50:M4:38:ILE:HD13	56:N0:148:LEU:HD13	5.52	0.63
52:M6:8:VAL:HG23	52:M6:34:VAL:HG13	1.79	0.63
5:S3:37:VAL:HG12	5:S3:50:ILE:HA	3.15	0.63
8:S6:201:GLN:HG2	1:6:126:A:OP1	335.18	0.63
1:2:53:G:H2'	1:2:54:C:C6	2.33	0.63
36:5:2510:U:O2'	36:5:2511:A:H5''	1.98	0.63
1:6:463:U:H2'	1:6:464:A:C8	2.33	0.63
12:C0:56:LYS:N	12:C0:67:THR:O	2.64	0.63
16:C4:81:VAL:HG13	16:C4:115:ILE:HG21	1.79	0.63
39:L2:98:VAL:HG13	39:L2:167:GLY:HA3	2.73	0.63
40:L3:123:TYR:HE2	40:L3:124:LYS:HZ2	6.45	0.63
40:L3:56:ILE:HG12	40:L3:356:LEU:HD22	1.81	0.63
40:L3:3:HIS:O	40:L3:3:HIS:ND1	3.80	0.63
42:L5:177:GLU:O	42:L5:179:ARG:N	2.44	0.63
46:L9:93:VAL:O	46:L9:177:ASP:HA	2.28	0.63
53:M7:103:GLU:O	53:M7:106:GLY:N	3.81	0.63
64:N8:71:PRO:HB2	64:N8:109:TYR:HA	1.80	0.63
69:O3:16:TYR:CD2	69:O3:25:PRO:HA	3.26	0.63
6:S4:25:GLY:HA3	1:6:447:U:O2'	375.38	0.63
8:S6:70:PRO:HD2	8:S6:71:THR:HG23	1.80	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:156:G:OP2	72:O6:25:LYS:HB3	1.99	0.63
36:1:2553:U:O2'	70:O4:91:ARG:NE	2.29	0.63
36:1:3042:U:OP2	36:1:3092:C:N4	2.21	0.63
1:2:651:G:O6	86:2:2103:OHX:N4	2.32	0.63
1:2:52:U:H2'	1:2:53:G:H8	1.64	0.63
1:2:839:U:H5''	13:C1:28:SER:HB3	1.81	0.63
31:D9:46:LYS:O	31:D9:48:ASN:N	2.31	0.63
42:L5:289:LYS:HD3	47:M0:206:LEU:HD23	1.81	0.63
36:1:49:A:OP1	49:M3:16:LYS:NZ	2.26	0.63
52:M6:25:LYS:HD2	52:M6:29:ASN:HD21	1.64	0.63
78:Q2:17:CYS:HG	78:Q2:77:CYS:CB	2.03	0.63
3:S1:48:VAL:HG13	3:S1:61:LEU:HD21	1.81	0.63
1:2:742:U:O2	9:S7:107:ARG:NH1	2.31	0.63
36:1:1220:U:OP1	36:1:1221:A:O2'	2.16	0.62
36:1:964:G:OP1	86:1:3957:OHX:N2	2.32	0.62
37:3:64:A:H3'	47:M0:204:GLY:O	1.99	0.62
1:6:1542:G:H22	1:6:1568:C:H1'	1.64	0.62
1:6:1680:G:O6	86:6:2189:OHX:N1	2.32	0.62
2:S0:62:ARG:HD3	23:D1:37:ALA:HB3	1.81	0.62
27:D5:50:ILE:HD12	27:D5:83:LEU:HD21	1.80	0.62
46:L9:18:VAL:HB	46:L9:27:VAL:HG22	1.81	0.62
46:L9:2:LYS:NZ	46:L9:59:ASN:HD21	1.96	0.62
51:M5:42:PRO:HG3	51:M5:61:ILE:HG13	1.80	0.62
52:M6:44:SER:O	52:M6:50:ASN:ND2	2.31	0.62
71:O5:78:LYS:HA	71:O5:81:ARG:HD2	1.81	0.62
3:S1:114:VAL:HG22	3:S1:142:PHE:HZ	2.85	0.62
6:S4:207:LEU:HD23	6:S4:221:ARG:HA	2.62	0.62
7:S5:97:LEU:O	7:S5:99:MET:N	3.28	0.62
36:1:299:G:N7	86:1:4076:OHX:N2	2.46	0.62
36:1:539:C:H2'	36:1:540:U:H6	1.64	0.62
1:2:1726:G:N7	86:2:2098:OHX:N4	2.47	0.62
36:5:3295:A:H2'	36:5:3296:A:C8	2.34	0.62
1:6:140:A:N6	1:6:281:G:OP1	2.32	0.62
18:C6:50:GLU:OE2	18:C6:82:ARG:NH2	4.47	0.62
19:C7:41:ILE:HD13	19:C7:47:ARG:HA	2.35	0.62
41:L4:286:VAL:O	41:L4:290:ILE:HG12	3.66	0.62
42:L5:40:HIS:CE1	57:N1:69:LYS:HA	2.34	0.62
43:L6:171:PRO:HA	43:L6:174:LEU:HD12	3.20	0.62
44:L7:121:LYS:HB2	57:N1:133:ALA:HB3	1.81	0.62
62:N6:28:ARG:HB2	62:N6:75:ARG:NH2	2.14	0.62
63:N7:29:HIS:O	63:N7:31:GLU:N	2.32	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
67:O1:57:GLN:NE2	36:5:1474:A:O2'	143.20	0.62
74:O8:40:GLN:NE2	74:O8:57:ASN:OD1	2.30	0.62
75:O9:34:THR:O	75:O9:36:ARG:N	3.01	0.62
79:Q3:13:LYS:HE3	79:Q3:14:TYR:CZ	2.34	0.62
4:S2:84:LYS:NZ	1:6:12:U:OP1	384.51	0.62
5:S3:116:ARG:O	5:S3:120:TYR:HB2	1.99	0.62
7:S5:174:LEU:O	7:S5:178:GLY:N	3.02	0.62
8:S6:67:VAL:HG21	8:S6:99:GLY:HA2	1.79	0.62
34:SR:90:ARG:HH21	34:SR:102:ARG:NH2	4.23	0.62
36:1:3152:U:O2'	36:1:3153:U:H5'	2.00	0.62
36:1:847:A:H2'	36:1:848:A:C8	2.34	0.62
1:2:482:U:H2'	1:2:483:A:H8	1.64	0.62
36:5:1659:U:H3	36:5:1790:G:H1	1.46	0.62
59:N3:49:LEU:HD12	36:5:2338:C:H1'	246.83	0.62
57:N1:92:ARG:NH1	36:5:2736:A:OP1	234.26	0.62
13:C1:90:TYR:OH	1:6:307:G:OP1	328.06	0.62
23:D1:55:LEU:HD11	23:D1:69:LEU:HG	2.92	0.62
31:D9:24:CYS:O	31:D9:25:SER:OG	2.13	0.62
33:E1:135:HIS:HB2	33:E1:138:ARG:HB3	1.80	0.62
39:L2:201:GLY:HA2	39:L2:204:MET:SD	3.40	0.62
39:L2:35:ALA:HA	45:L8:36:ILE:HD13	2.09	0.62
48:M1:52:TYR:HA	48:M1:61:ARG:HB2	1.80	0.62
52:M6:110:PRO:O	52:M6:113:ASP:N	5.00	0.62
59:N3:48:ARG:HG2	36:5:2339:C:P	247.11	0.62
72:O6:81:THR:HA	72:O6:84:LYS:HE3	4.58	0.62
79:Q3:49:ARG:HD3	79:Q3:50:GLY:N	2.14	0.62
7:S5:143:ARG:HG3	7:S5:167:ARG:NH1	4.22	0.62
11:S9:38:ASN:HB2	11:S9:41:GLU:HG3	1.81	0.62
1:2:1619:C:H2'	1:2:1620:C:H6	1.64	0.62
36:5:2233:A:OP2	86:5:3955:OHX:N5	2.31	0.62
18:C6:40:GLU:HA	18:C6:42:GLU:N	2.15	0.62
41:L4:311:HIS:NE2	41:L4:314:LYS:HA	2.14	0.62
46:L9:90:MET:HB2	46:L9:144:ILE:HG22	2.18	0.62
46:L9:163:GLN:OE1	46:L9:166:ARG:NH1	2.32	0.62
49:M3:73:ARG:NH2	36:5:77:A:N7	79.78	0.62
52:M6:22:VAL:HG21	52:M6:120:VAL:HG11	1.79	0.62
77:Q1:6:ARG:O	77:Q1:10:THR:OG1	2.16	0.62
77:Q1:2:ARG:HG2	77:Q1:5:TRP:CD1	3.80	0.62
3:S1:134:VAL:N	3:S1:219:LYS:O	2.90	0.62
3:S1:51:SER:HB3	3:S1:57:ALA:H	2.15	0.62
36:1:1599:G:H1	36:1:1608:C:H42	1.48	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:2185:G:O2'	36:1:2314:U:OP2	2.17	0.62
86:1:4194:OHX:N6	86:O1:202:OHX:N5	2.48	0.62
1:6:151:G:H2'	1:6:152:U:H6	1.64	0.62
1:6:191:C:O2'	1:6:192:U:O5'	2.14	0.62
15:C3:64:ARG:HD2	15:C3:70:LYS:HD2	5.46	0.62
17:C5:122:THR:HG21	1:6:1455:G:OP1	370.26	0.62
19:C7:88:VAL:HG22	19:C7:89:SER:H	1.64	0.62
20:C8:30:TYR:O	20:C8:33:THR:OG1	2.17	0.62
22:D0:57:ARG:HG3	22:D0:89:ARG:CZ	2.29	0.62
43:L6:91:VAL:HG23	43:L6:92:SER:O	3.17	0.62
53:M7:102:ALA:HA	53:M7:107:LEU:HD23	1.81	0.62
53:M7:178:ALA:HA	53:M7:181:ARG:HB3	1.82	0.62
69:O3:59:VAL:O	69:O3:61:GLY:N	2.42	0.62
3:S1:113:MET:HE3	3:S1:142:PHE:HE2	5.91	0.62
1:2:580:A:H5''	5:S3:143:ARG:HH12	1.64	0.62
36:1:2228:A:H2'	36:1:2229:A:C8	2.35	0.62
1:2:800:U:O4	86:2:2053:OHX:N5	2.32	0.62
1:2:471:A:OP2	86:2:2075:OHX:N4	2.32	0.62
1:2:264:G:N7	86:2:2034:OHX:N1	2.48	0.62
1:2:71:A:H1'	1:2:81:G:N2	2.14	0.62
36:5:1135:A:C2	36:5:1136:A:C8	2.87	0.62
36:5:1222:G:O6	86:5:4124:OHX:N1	2.32	0.62
36:5:2970:C:H4'	36:5:2971:A:N1	2.14	0.62
36:5:3362:A:C2	36:5:3363:U:C2	2.88	0.62
1:6:1081:A:H1'	1:6:1082:C:H5	1.63	0.62
16:C4:91:THR:O	16:C4:93:THR:N	2.30	0.62
22:D0:106:ILE:HG13	22:D0:107:THR:H	1.62	0.62
42:L5:39:GLN:HG3	42:L5:40:HIS:O	2.64	0.62
51:M5:14:LYS:HE2	36:5:269:G:H5''	132.73	0.62
53:M7:27:LYS:HD3	53:M7:63:PHE:HB3	1.98	0.62
54:M8:165:ILE:HD12	54:M8:167:SER:O	4.58	0.62
57:N1:19:PHE:CD2	36:5:1051:U:H4'	285.39	0.62
57:N1:38:ASP:N	57:N1:38:ASP:OD1	2.26	0.62
2:S0:14:ALA:HA	2:S0:17:LEU:HD12	1.80	0.62
3:S1:51:SER:HA	3:S1:57:ALA:H	1.64	0.62
5:S3:115:ILE:HD11	5:S3:138:VAL:HG21	1.80	0.62
8:S6:114:VAL:O	8:S6:115:LYS:HD3	1.99	0.62
34:SR:91:LEU:O	34:SR:100:TYR:N	2.26	0.62
36:1:2623:G:H2'	36:1:2624:G:C8	2.34	0.62
1:2:1650:U:H2'	1:2:1651:A:C8	2.34	0.62
1:2:824:G:O6	1:2:848:C:N4	2.31	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:1564:U:H2'	36:5:1565:G:H8	1.63	0.62
45:L8:240:ASN:ND2	36:5:2584:G:N3	185.09	0.62
1:6:615:A:O2'	1:6:621:A:N1	2.29	0.62
18:C6:11:GLY:HA2	18:C6:83:GLN:HE21	1.65	0.62
49:M3:39:ARG:NH2	36:5:686:G:OP2	74.95	0.62
62:N6:45:ILE:HD11	62:N6:122:LYS:HD3	2.04	0.62
70:O4:58:ARG:HG3	70:O4:58:ARG:HH11	2.63	0.62
10:S8:12:SER:O	10:S8:15:GLY:N	2.64	0.62
36:1:176:G:O6	36:1:242:C:N4	2.32	0.62
36:1:2442:G:N2	36:1:2505:U:H3	1.98	0.62
36:1:764:U:O4	86:1:3955:OHX:N5	2.31	0.62
1:2:1642:G:O6	86:2:2023:OHX:N6	2.32	0.62
1:2:57:G:O6	86:2:2045:OHX:N3	2.31	0.62
1:2:856:A:H1'	9:S7:64:VAL:HG11	1.81	0.62
36:5:1497:C:H2'	36:5:1498:A:C8	2.31	0.62
36:5:210:U:C2	36:5:230:U:H4'	2.34	0.62
1:6:1699:G:H22	1:6:1702:A:H5''	1.65	0.62
12:C0:6:GLU:O	12:C0:10:LYS:HG3	2.00	0.62
16:C4:81:VAL:H	16:C4:115:ILE:HG22	1.63	0.62
28:D6:10:ARG:HD3	28:D6:34:LYS:HA	2.33	0.62
41:L4:60:THR:HG21	41:L4:77:VAL:HG22	1.82	0.62
36:1:3308:C:N3	53:M7:69:ARG:NH1	2.47	0.62
78:Q2:12:CYS:SG	78:Q2:17:CYS:CB	3.18	0.62
78:Q2:73:GLU:OE1	78:Q2:80:ARG:NH1	2.92	0.62
8:S6:164:LYS:HB3	8:S6:167:LYS:HB3	1.82	0.62
10:S8:84:HIS:CE1	10:S8:86:SER:HB2	2.33	0.62
11:S9:129:ILE:HA	11:S9:134:ILE:HD12	1.82	0.62
36:5:1266:G:O6	36:5:1275:C:N4	2.27	0.62
36:5:1307:G:O2'	36:5:1308:A:N7	2.31	0.62
36:5:2696:A:H2'	36:5:2697:A:C8	2.35	0.62
36:5:3089:C:H2'	36:5:3090:U:O4'	2.00	0.62
36:5:299:G:N7	86:5:4184:OHX:N1	2.48	0.62
1:6:27:U:H2'	1:6:28:A:H8	1.63	0.62
1:6:580:A:O2'	1:6:582:U:OP1	2.18	0.62
42:L5:115:LEU:H	42:L5:115:LEU:HD22	1.64	0.62
42:L5:38:THR:HG22	57:N1:30:TYR:HB3	1.82	0.62
46:L9:22:SER:OG	46:L9:23:ARG:N	2.32	0.62
52:M6:121:PRO:HA	52:M6:124:LEU:HD22	1.99	0.62
52:M6:185:ALA:O	52:M6:188:SER:N	3.41	0.62
56:N0:66:GLU:OE1	56:N0:99:ARG:N	2.26	0.62
68:O2:96:ILE:HG21	68:O2:105:ARG:HG2	1.81	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:SM:47:ALA:HB2	36:1:2678:A:C8	2.35	0.62
36:1:2396:G:OP1	36:1:2397:A:H4'	2.00	0.62
1:2:635:A:H2'	1:2:636:A:C8	2.35	0.62
1:2:635:A:H2'	1:2:636:A:H8	1.65	0.62
1:6:538:A:H2	1:6:540:G:H22	1.48	0.62
37:7:114:U:H2'	37:7:115:G:H8	1.65	0.62
48:M1:143:ARG:NH2	37:7:5:G:OP1	291.54	0.62
33:E1:97:LYS:HE2	33:E1:98:VAL:HG12	1.81	0.62
42:L5:105:ILE:O	42:L5:109:THR:HG23	1.99	0.62
44:L7:88:ARG:NH1	44:L7:91:GLY:O	2.51	0.62
46:L9:136:PHE:CE2	46:L9:144:ILE:HG12	5.59	0.62
36:1:2902:A:OP1	46:L9:170:LYS:NZ	2.33	0.62
56:N0:12:ARG:HH12	56:N0:15:PRO:HG3	1.64	0.62
65:N9:12:GLN:NE2	36:5:953:G:OP1	210.83	0.62
71:O5:95:PHE:N	36:5:135:C:O2'	57.73	0.62
9:S7:74:GLN:O	9:S7:78:THR:OG1	2.16	0.62
36:1:2261:G:O2'	36:1:2263:C:N4	2.33	0.61
36:1:3155:U:H3'	36:1:3156:U:H4'	1.82	0.61
36:1:2123:G:N7	86:1:4195:OHX:N2	2.48	0.61
36:5:1062:A:H5''	36:5:1063:G:H5'	1.80	0.61
36:5:1530:U:OP1	86:5:3984:OHX:N1	2.33	0.61
36:5:22:G:H1'	38:8:104:A:N3	2.14	0.61
76:Q0:102:ARG:NE	36:5:2896:A:OP1	321.46	0.61
36:5:2993:G:H2'	36:5:3142:A:N6	2.14	0.61
36:5:994:G:O2'	36:5:1053:A:N6	2.24	0.61
1:6:1471:A:H62	1:6:1538:U:H3	1.48	0.61
12:C0:27:PHE:HB3	12:C0:40:LEU:HD23	1.81	0.61
17:C5:75:PRO:HA	17:C5:93:VAL:HG12	1.81	0.61
20:C8:33:THR:HA	20:C8:38:VAL:HG23	1.82	0.61
43:L6:175:LYS:HE3	50:M4:111:ALA:HA	5.56	0.61
53:M7:70:THR:OG1	53:M7:72:GLN:N	3.23	0.61
2:S0:4:PRO:HG3	23:D1:42:GLU:H	8.25	0.61
36:1:121:A:C6	45:L8:129:PRO:HG3	2.36	0.61
36:1:1383:G:O6	86:1:3873:OHX:N3	2.33	0.61
36:1:1115:G:O6	86:1:3957:OHX:N6	2.33	0.61
36:1:1789:G:O6	86:1:4165:OHX:N4	2.33	0.61
1:2:1773:C:OP1	77:Q1:3:ALA:HB3	2.00	0.61
1:2:1212:G:O6	86:2:2029:OHX:N3	2.33	0.61
36:5:2697:A:H2'	36:5:2698:G:H8	1.63	0.61
21:C9:60:SER:OG	1:6:1480:G:OP1	399.57	0.61
1:6:217:A:C8	1:6:218:A:C8	2.88	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:138:A:N6	1:6:266:A:H61	1.98	0.61
1:6:383:G:N7	86:6:2149:OHX:N5	2.47	0.61
1:6:486:G:O6	1:6:488:G:N2	2.31	0.61
1:6:895:G:H2'	1:6:896:U:C6	2.35	0.61
19:C7:71:PHE:HD1	19:C7:73:LEU:HB3	1.64	0.61
37:3:28:C:OP2	42:L5:57:ASN:ND2	2.33	0.61
43:L6:55:LEU:HD11	43:L6:66:SER:HB2	2.47	0.61
44:L7:186:HIS:O	44:L7:190:THR:HG23	2.13	0.61
50:M4:23:ILE:HD13	50:M4:63:VAL:HG23	1.82	0.61
52:M6:27:LEU:HD22	52:M6:101:ARG:HB2	2.96	0.61
55:M9:89:LEU:HD12	55:M9:90:PRO:HD2	2.09	0.61
56:N0:30:PHE:CE2	56:N0:103:VAL:HG21	2.35	0.61
68:O2:33:ARG:HG3	36:5:945:C:OP1	170.41	0.61
69:O3:26:ASN:HA	69:O3:88:ASN:OD1	3.46	0.61
3:S1:180:THR:HG22	3:S1:181:LEU:H	1.65	0.61
1:2:607:G:H5'	1:2:613:G:N2	2.16	0.61
36:5:209:A:H4'	36:5:211:A:N7	2.15	0.61
36:5:2310:U:OP1	86:5:4193:OHX:N2	2.34	0.61
36:5:499:G:H2'	36:5:500:C:H6	1.65	0.61
13:C1:79:LYS:HB3	1:6:346:G:H5'	282.79	0.61
7:S5:25:LEU:HB2	18:C6:27:GLY:O	2.66	0.61
39:L2:29:LEU:O	39:L2:123:ARG:NE	2.29	0.61
45:L8:108:ARG:O	45:L8:112:GLU:N	2.79	0.61
64:N8:76:ASP:HB2	64:N8:115:LYS:HB3	1.81	0.61
73:O7:18:LEU:HD21	75:O9:51:ILE:HB	1.80	0.61
5:S3:64:ARG:O	5:S3:66:ILE:N	2.34	0.61
8:S6:181:PRO:HA	8:S6:184:LEU:HD12	3.14	0.61
9:S7:141:ARG:HH21	9:S7:143:LEU:HD11	1.64	0.61
20:C8:120:ARG:HG2	35:SM:61:ILE:HG21	6.78	0.61
36:1:3165:A:H61	36:1:3285:C:H42	1.46	0.61
36:1:662:U:OP1	64:N8:8:THR:HG21	1.99	0.61
1:6:282:C:H2'	1:6:283:U:O4'	2.00	0.61
37:7:104:A:H5''	37:7:105:C:OP2	1.99	0.61
16:C4:29:HIS:O	16:C4:29:HIS:ND1	2.32	0.61
19:C7:71:PHE:CD1	19:C7:73:LEU:HB3	2.35	0.61
40:L3:81:THR:HG23	40:L3:205:VAL:HG21	4.09	0.61
41:L4:140:HIS:H	41:L4:180:LYS:HE3	1.65	0.61
41:L4:220:ARG:NH1	36:5:211:A:OP1	73.91	0.61
47:M0:72:ALA:HB2	47:M0:155:ALA:HB2	2.25	0.61
50:M4:128:ARG:HH11	50:M4:128:ARG:HB3	4.33	0.61
51:M5:116:LEU:HD23	51:M5:133:ILE:HD11	1.82	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:M8:44:PHE:CD2	54:M8:134:GLY:HA3	2.80	0.61
78:Q2:77:CYS:O	78:Q2:79:THR:N	2.49	0.61
5:S3:58:VAL:O	5:S3:65:ARG:HB3	2.00	0.61
7:S5:84:LYS:HD3	7:S5:92:ARG:HH22	1.65	0.61
36:1:1727:G:OP1	79:Q3:44:LYS:NZ	2.31	0.61
36:1:1413:G:N7	86:1:4118:OHX:N4	2.48	0.61
37:3:27:A:OP2	42:L5:57:ASN:HB2	2.00	0.61
36:5:3165:A:N6	36:5:3285:C:H42	1.95	0.61
1:2:905:A:H5''	16:C4:52:ARG:HD3	1.82	0.61
18:C6:21:HIS:HB2	18:C6:66:ARG:HB3	1.82	0.61
19:C7:108:ASP:HA	19:C7:111:LYS:HD3	1.83	0.61
36:1:119:U:C2	45:L8:138:HIS:CE1	2.89	0.61
45:L8:48:ARG:NH2	36:5:2588:U:OP1	183.65	0.61
51:M5:21:PHE:O	51:M5:25:VAL:HG23	2.21	0.61
59:N3:120:LYS:HD3	59:N3:121:GLU:HG3	1.81	0.61
67:O1:12:TYR:O	67:O1:72:ARG:HD2	2.00	0.61
6:S4:159:THR:HB	6:S4:227:VAL:HG23	1.82	0.61
6:S4:233:LYS:NZ	6:S4:233:LYS:O	6.84	0.61
7:S5:128:ASN:HB2	7:S5:131:GLN:HB3	1.81	0.61
34:SR:123:ILE:HG21	34:SR:169:ILE:HG21	2.60	0.61
34:SR:72:THR:HG23	34:SR:81:LEU:HB2	3.04	0.61
36:1:1103:A:N3	36:1:1103:A:H2'	2.16	0.61
36:1:1235:U:H4'	36:1:1236:G:H5'	1.82	0.61
36:1:1581:C:O2	36:1:1582:C:H5'	2.00	0.61
36:1:1780:G:H2'	36:1:1781:C:H6	1.65	0.61
36:1:2528:G:N7	86:1:4180:OHX:N3	2.48	0.61
36:1:7:C:H2'	36:1:8:C:C6	2.36	0.61
36:5:3204:C:H2'	36:5:3205:G:C8	2.35	0.61
86:5:3966:OHX:N3	86:5:4238:OHX:N5	2.49	0.61
36:5:223:U:O4	86:5:4239:OHX:N4	2.34	0.61
1:6:386:G:H2'	1:6:387:A:C8	2.34	0.61
1:6:587:C:H2'	1:6:588:U:O4'	2.00	0.61
15:C3:124:ARG:NH2	1:6:967:A:OP2	319.99	0.61
38:8:113:U:O2'	38:8:114:G:OP2	2.14	0.61
22:D0:103:ILE:HA	22:D0:106:ILE:HG22	3.06	0.61
2:S0:36:TYR:OH	23:D1:66:ASP:OD2	2.34	0.61
30:D8:44:VAL:HG21	30:D8:48:VAL:HG21	2.25	0.61
42:L5:22:ARG:NH1	42:L5:28:THR:OG1	4.72	0.61
45:L8:150:LEU:HD23	45:L8:176:PRO:HB2	2.41	0.61
45:L8:194:THR:OG1	45:L8:195:SER:N	2.32	0.61
46:L9:90:MET:HE3	46:L9:162:GLN:HB3	1.82	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:2899:C:C5	46:L9:171:ASP:HA	2.35	0.61
53:M7:127:ARG:HB3	53:M7:139:TYR:O	2.18	0.61
56:N0:139:TYR:HD2	56:N0:140:VAL:HG23	1.94	0.61
57:N1:14:MET:HE1	57:N1:55:LYS:HB2	1.80	0.61
78:Q2:71:ARG:HH21	78:Q2:80:ARG:NH1	3.50	0.61
36:1:1060:U:H2'	36:1:1061:A:H8	1.65	0.61
36:1:3349:C:H42	36:1:3356:G:H1	1.49	0.61
1:2:1358:G:H2'	1:2:1359:C:C6	2.36	0.61
1:2:1479:A:H2'	1:2:1480:G:H8	1.65	0.61
37:3:49:G:C5	42:L5:58:LYS:HG3	2.35	0.61
36:5:1024:G:N2	36:5:1026:A:OP2	2.34	0.61
36:5:1355:A:H4'	36:5:1356:U:O5'	2.00	0.61
36:5:1783:U:H2'	36:5:1784:G:C8	2.36	0.61
51:M5:68:ARG:HG3	36:5:291:C:OP1	145.52	0.61
36:5:2239:G:N7	86:5:4187:OHX:N5	2.49	0.61
21:C9:97:SER:OG	1:6:1504:G:OP1	394.89	0.61
1:6:782:U:H5''	1:6:782:U:O2	2.00	0.61
18:C6:40:GLU:HA	18:C6:42:GLU:H	1.66	0.61
23:D1:60:ARG:HA	23:D1:65:SER:HB2	2.04	0.61
40:L3:122:TRP:CE2	40:L3:127:LYS:HE3	2.36	0.61
48:M1:10:ARG:HA	48:M1:134:PRO:HD2	2.91	0.61
51:M5:140:LYS:NZ	71:O5:96:GLU:OE2	2.33	0.61
70:O4:6:THR:HG21	36:5:1487:G:H1'	141.60	0.61
2:S0:118:PRO:HG2	2:S0:141:ILE:HD13	1.99	0.61
2:S0:189:VAL:HG22	2:S0:190:ASP:H	1.65	0.61
6:S4:242:LYS:N	6:S4:242:LYS:HE3	2.16	0.61
6:S4:95:THR:HG23	6:S4:97:GLU:HG3	5.51	0.61
1:2:161:U:OP2	8:S6:87:ARG:NH2	2.33	0.61
36:1:2960:C:H2'	36:1:2961:G:C8	2.35	0.61
36:1:2970:C:H4'	36:1:2971:A:N6	2.15	0.61
36:1:3181:C:H2'	36:1:3182:G:C8	2.36	0.61
1:2:1482:C:OP2	1:2:1521:G:N2	2.31	0.61
1:2:1660:A:H2'	1:2:1661:U:H6	1.65	0.61
1:2:778:G:H3'	1:2:780:A:H2	1.64	0.61
36:5:136:G:H2'	36:5:137:G:H8	1.65	0.61
36:5:2217:U:H2'	36:5:2218:G:C8	2.36	0.61
36:5:2722:U:H2'	36:5:2723:U:H6	1.64	0.61
36:5:3066:U:O4	86:5:4100:OHX:N4	2.34	0.61
86:5:3935:OHX:N2	86:5:4228:OHX:N4	2.48	0.61
86:5:3935:OHX:N2	86:5:4228:OHX:N6	2.48	0.61
86:5:3935:OHX:N5	86:5:4228:OHX:N3	2.49	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:C1:90:TYR:CE1	13:C1:103:ARG:HB2	2.36	0.61
15:C3:127:ARG:O	15:C3:131:THR:OG1	2.18	0.61
24:D2:6:VAL:HG13	24:D2:29:PRO:HD2	1.82	0.61
47:M0:143:SER:O	47:M0:143:SER:OG	3.55	0.61
48:M1:52:TYR:HA	48:M1:61:ARG:HG3	3.05	0.61
51:M5:38:ARG:NH2	51:M5:60:VAL:HG22	2.15	0.61
56:N0:29:ILE:HD12	56:N0:40:ARG:HB3	1.82	0.61
59:N3:127:PRO:O	59:N3:131:SER:N	3.38	0.61
62:N6:56:VAL:HG21	62:N6:104:LEU:HD13	1.83	0.61
66:O0:99:ASP:HB2	66:O0:103:THR:HG23	1.83	0.61
63:N7:136:PHE:HB2	70:O4:88:ARG:HG3	2.28	0.61
74:O8:5:ILE:HG22	74:O8:54:LEU:HB2	1.81	0.61
2:S0:72:ASP:OD1	4:S2:40:LYS:NZ	3.71	0.61
11:S9:47:PHE:CZ	11:S9:51:LYS:HD3	2.59	0.61
36:1:3330:A:H2'	36:1:3331:U:H6	1.65	0.61
36:1:884:A:OP1	73:O7:5:THR:OG1	2.13	0.61
1:2:1323:C:H2'	1:2:1324:G:O4'	2.00	0.61
1:2:1606:C:H2'	1:2:1607:G:C8	2.36	0.61
1:2:585:A:N6	1:2:586:G:O6	2.33	0.61
1:6:1166:A:H2'	1:6:1167:G:O4'	2.01	0.61
1:6:488:G:H1'	1:6:500:C:H42	1.66	0.61
41:L4:193:LYS:NZ	38:8:21:C:OP1	109.02	0.61
14:C2:75:VAL:HG21	14:C2:120:VAL:HG21	1.86	0.61
25:D3:19:ARG:O	25:D3:22:ASN:N	2.34	0.61
25:D3:56:LYS:HD2	25:D3:97:ASP:HA	1.81	0.61
30:D8:10:ALA:HA	30:D8:32:PHE:HA	1.83	0.61
51:M5:184:LYS:H	51:M5:186:GLY:H	1.49	0.61
52:M6:138:LEU:O	52:M6:141:LEU:N	2.51	0.61
57:N1:82:ASN:HB3	65:N9:16:ALA:HB1	2.26	0.61
73:O7:58:THR:O	73:O7:61:THR:HG23	1.99	0.61
79:Q3:30:GLU:HA	79:Q3:33:GLN:OE1	2.00	0.61
2:S0:190:ASP:O	2:S0:192:THR:N	4.01	0.61
7:S5:110:ALA:HA	7:S5:113:ILE:HD12	1.81	0.61
36:1:687:U:OP2	49:M3:36:ARG:NH2	2.33	0.61
1:2:1147:A:H2'	1:2:1148:C:C6	2.34	0.61
36:5:2217:U:H2'	36:5:2218:G:H8	1.65	0.61
12:C0:23:ALA:HB3	12:C0:64:TYR:HB2	1.81	0.61
18:C6:112:TYR:OH	18:C6:114:ARG:NH1	2.34	0.61
1:2:1335:U:OP1	22:D0:85:ARG:NH1	2.34	0.61
26:D4:52:LYS:O	26:D4:55:VAL:N	2.34	0.61
33:E1:144:CYS:O	33:E1:146:SER:N	2.51	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:L3:296:THR:H	40:L3:299:ASP:HB3	1.66	0.61
45:L8:90:THR:HA	45:L8:214:LEU:HD21	1.82	0.61
64:N8:112:ILE:HB	64:N8:130:VAL:HG12	2.13	0.61
67:O1:88:PRO:HG2	67:O1:89:LEU:HD12	3.18	0.61
36:1:283:G:OP1	78:Q2:45:ARG:NH2	2.33	0.61
4:S2:116:LYS:HG2	4:S2:127:ALA:HB3	1.83	0.61
4:S2:179:VAL:O	4:S2:198:THR:OG1	2.14	0.61
11:S9:149:ARG:O	11:S9:151:ASP:N	2.31	0.61
35:SM:84:LYS:HG2	35:SM:86:ASN:H	1.66	0.61
36:1:138:U:H2'	36:1:139:G:C8	2.36	0.60
36:1:3353:G:O2'	36:1:3356:G:OP2	2.19	0.60
36:1:518:G:N7	36:1:572:A:N6	2.45	0.60
1:2:1680:G:O6	86:2:2109:OHX:N5	2.34	0.60
36:5:1096:U:H4'	36:5:1097:G:O5'	2.01	0.60
36:5:2436:U:H2'	36:5:2437:G:H5'	1.83	0.60
36:5:2908:G:N7	86:5:3895:OHX:N2	2.49	0.60
1:6:1004:U:H3'	1:6:1005:A:H5''	1.83	0.60
7:S5:185:ARG:NH1	1:6:1471:A:OP1	334.79	0.60
17:C5:87:PRO:HA	17:C5:90:ILE:HG13	2.72	0.60
26:D4:62:THR:HA	26:D4:69:SER:HA	1.94	0.60
41:L4:193:LYS:O	41:L4:198:ARG:HG2	3.77	0.60
50:M4:20:VAL:HG22	50:M4:66:THR:OG1	2.01	0.60
50:M4:65:LEU:HB2	56:N0:172:TYR:CE1	2.36	0.60
62:N6:13:ARG:HD3	38:8:24:G:OP2	86.37	0.60
63:N7:95:VAL:HG13	63:N7:110:ALA:HB1	1.83	0.60
70:O4:44:CYS:SG	70:O4:81:CYS:HB3	2.41	0.60
72:O6:95:ALA:HA	72:O6:99:ARG:HB2	1.83	0.60
79:Q3:73:THR:HG22	79:Q3:75:ALA:H	4.98	0.60
5:S3:105:MET:HG3	5:S3:122:VAL:HG21	1.82	0.60
7:S5:112:ARG:NH1	18:C6:43:ILE:HD11	2.16	0.60
34:SR:289:ALA:HA	34:SR:305:TYR:HA	1.88	0.60
37:3:101:G:OP2	56:N0:52:LYS:NZ	2.34	0.60
36:5:2722:U:H2'	36:5:2723:U:C6	2.36	0.60
36:5:1744:G:O6	86:5:4095:OHX:N1	2.34	0.60
1:6:1650:U:H2'	1:6:1651:A:C8	2.36	0.60
14:C2:76:GLU:OE2	14:C2:90:LYS:NZ	2.34	0.60
1:2:1132:A:OP1	25:D3:30:LYS:HE2	2.01	0.60
33:E1:109:ASP:O	33:E1:111:GLU:N	2.34	0.60
41:L4:237:GLN:O	41:L4:246:ARG:HG3	2.01	0.60
41:L4:346:LYS:HD2	41:L4:347:THR:H	6.09	0.60
42:L5:211:LEU:O	42:L5:215:ASP:N	3.43	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:L5:270:LYS:HG3	42:L5:273:ARG:CB	4.88	0.60
48:M1:9:MET:O	48:M1:9:MET:HG3	2.00	0.60
51:M5:143:ARG:NH2	71:O5:92:LEU:HD23	2.72	0.60
52:M6:110:PRO:O	52:M6:111:PRO:C	3.55	0.60
66:O0:25:LEU:HD22	66:O0:87:VAL:HG21	2.82	0.60
43:L6:164:SER:OG	69:O3:4:SER:HB2	2.00	0.60
7:S5:121:ILE:HD11	7:S5:198:LEU:HD13	1.83	0.60
8:S6:63:MET:HE1	8:S6:106:LEU:HD13	1.83	0.60
36:1:1743:G:H2'	36:1:1744:G:H8	1.66	0.60
36:1:2921:U:H2'	36:1:2923:U:OP2	2.01	0.60
36:1:3164:C:H1'	36:1:3165:A:H5'	1.83	0.60
36:1:77:A:H5'	49:M3:100:ARG:NH1	2.17	0.60
86:2:2031:OHX:N3	86:2:2146:OHX:N1	2.48	0.60
1:2:280:U:O2'	1:2:281:G:OP2	2.16	0.60
1:2:792:U:C2'	1:2:793:A:H5'	2.30	0.60
38:4:124:G:H1	38:4:129:C:H42	1.47	0.60
36:5:3181:C:H2'	36:5:3182:G:C8	2.35	0.60
8:S6:13:GLN:HE22	1:6:151:G:N2	311.27	0.60
1:6:1533:C:H4'	1:6:1539:G:N1	2.16	0.60
15:C3:88:LEU:O	15:C3:92:ILE:HG13	2.00	0.60
21:C9:42:GLY:HA2	21:C9:84:LYS:HG3	1.82	0.60
27:D5:47:TYR:CZ	27:D5:51:LEU:HD11	2.95	0.60
31:D9:33:LYS:HD3	31:D9:34:TYR:CZ	2.68	0.60
36:1:1388:U:H5	41:L4:186:LYS:HZ2	1.47	0.60
44:L7:165:ASP:OD2	44:L7:166:ASN:N	2.79	0.60
48:M1:100:GLY:HA3	48:M1:154:THR:HB	3.27	0.60
49:M3:92:THR:O	49:M3:92:THR:OG1	3.52	0.60
51:M5:118:SER:HB3	51:M5:132:VAL:HG13	1.83	0.60
58:N2:17:VAL:HG22	58:N2:103:TYR:HB2	1.83	0.60
8:S6:98:ARG:NE	8:S6:105:ASP:OD1	2.55	0.60
36:1:1554:U:H4'	36:1:1555:U:H5'	1.82	0.60
1:2:912:U:H5'	1:2:913:G:H2'	1.83	0.60
36:5:1778:G:O2'	36:5:1780:G:OP2	2.17	0.60
36:5:1863:G:N1	36:5:1866:C:OP2	2.34	0.60
55:M9:20:ARG:NH2	36:5:1874:A:N7	148.18	0.60
36:5:2516:U:O2'	36:5:2595:A:N1	2.31	0.60
72:O6:86:LYS:NZ	36:5:296:A:OP1	140.45	0.60
36:5:2249:G:OP1	86:5:4193:OHX:N6	2.34	0.60
36:5:796:U:H2'	36:5:797:U:C6	2.36	0.60
1:6:151:G:N2	1:6:163:G:N2	2.49	0.60
10:S8:10:LYS:HE2	1:6:339:C:OP2	286.04	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:737:A:H2'	1:6:738:G:C8	2.37	0.60
12:C0:32:HIS:CG	12:C0:33:GLU:H	4.00	0.60
13:C1:92:HIS:CD2	13:C1:93:TYR:H	2.19	0.60
19:C7:28:PHE:HA	19:C7:55:THR:HG21	2.88	0.60
1:2:574:G:O6	25:D3:65:ASN:ND2	2.33	0.60
26:D4:81:GLU:O	26:D4:85:PHE:N	3.42	0.60
39:L2:105:GLY:HA2	39:L2:139:HIS:CE1	4.28	0.60
44:L7:42:ALA:O	44:L7:45:LEU:N	2.34	0.60
49:M3:165:SER:OG	49:M3:167:PHE:HB3	2.01	0.60
53:M7:88:VAL:HG12	53:M7:92:GLN:HG3	4.81	0.60
54:M8:180:ARG:NH1	54:M8:185:LYS:HB3	2.14	0.60
59:N3:18:PRO:HA	59:N3:51:ALA:HA	1.81	0.60
61:N5:132:ALA:O	61:N5:136:ALA:N	2.72	0.60
8:S6:4:ASN:HA	8:S6:15:THR:HG22	3.35	0.60
9:S7:70:PHE:O	9:S7:74:GLN:HB2	2.46	0.60
34:SR:255:ALA:HB2	34:SR:292:LEU:HD22	1.82	0.60
36:1:595:G:C8	36:1:609:G:C6	2.89	0.60
37:3:71:G:H2'	37:3:72:A:H8	1.64	0.60
36:5:1858:A:O2'	36:5:1859:A:OP2	2.19	0.60
36:5:1934:G:O6	86:5:3908:OHX:N2	2.34	0.60
36:5:3194:C:C2	36:5:3197:G:N2	2.69	0.60
17:C5:22:LEU:HA	17:C5:25:LEU:HD12	2.48	0.60
39:L2:209:HIS:CD2	39:L2:211:HIS:H	2.19	0.60
44:L7:116:PHE:HB2	44:L7:199:ASN:OD1	2.19	0.60
48:M1:57:PHE:HD2	36:5:2680:A:C4	309.96	0.60
41:L4:362:ASP:H	56:N0:26:ARG:HH12	4.57	0.60
65:N9:20:GLY:O	65:N9:21:ILE:HB	2.37	0.60
66:O0:42:ILE:HD11	66:O0:67:VAL:HG22	1.82	0.60
68:O2:21:HIS:ND1	68:O2:24:ARG:HD2	2.15	0.60
69:O3:72:THR:HG21	69:O3:84:THR:HG23	3.33	0.60
2:S0:41:ARG:HH11	2:S0:45:VAL:HG21	3.03	0.60
3:S1:142:PHE:O	3:S1:208:GLN:N	2.33	0.60
6:S4:129:VAL:HB	6:S4:139:VAL:HG12	1.83	0.60
6:S4:191:ARG:NH1	6:S4:245:LYS:HD3	2.16	0.60
36:1:2413:A:H2'	36:1:2414:G:C8	2.36	0.60
1:2:1239:U:O2	1:2:1246:C:N4	2.35	0.60
1:2:1350:U:H2'	1:2:1351:G:C8	2.35	0.60
1:2:190:C:O2'	1:2:191:C:OP2	2.13	0.60
1:2:590:C:H5''	32:E0:43:ARG:HH12	1.66	0.60
36:5:999:G:C6	36:5:1000:C:N4	2.69	0.60
75:O9:44:TRP:HB3	36:5:1494:U:OP1	111.75	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:3337:G:H2'	36:5:3338:C:C6	2.37	0.60
36:5:581:U:O4	86:5:4017:OHX:N6	2.35	0.60
36:5:720:A:C2	36:5:784:A:H5'	2.36	0.60
10:S8:138:ASN:ND2	1:6:197:A:N1	278.53	0.60
17:C5:47:ARG:HH21	1:6:1555:A:P	404.81	0.60
18:C6:82:ARG:HH12	18:C6:114:ARG:HB2	5.09	0.60
42:L5:52:VAL:HG22	42:L5:147:ASP:HB3	1.83	0.60
45:L8:240:ASN:HA	45:L8:243:GLN:HB2	1.82	0.60
46:L9:70:THR:HB	36:5:3112:G:O2'	328.90	0.60
47:M0:4:ARG:NH2	47:M0:99:ILE:HG22	6.28	0.60
43:L6:89:THR:HG21	50:M4:115:PHE:HB2	2.04	0.60
52:M6:113:ASP:OD1	52:M6:114:LYS:HG3	2.01	0.60
58:N2:49:ASN:O	58:N2:51:GLY:N	2.36	0.60
5:S3:107:PHE:O	5:S3:111:ASN:ND2	2.35	0.60
6:S4:51:ARG:HB3	6:S4:111:VAL:HG22	1.83	0.60
9:S7:50:ASP:HA	9:S7:56:LYS:HA	1.82	0.60
36:5:2387:A:C2	36:5:2388:U:H1'	2.37	0.60
1:6:53:G:H1	1:6:427:C:H42	1.49	0.60
25:D3:68:ILE:O	25:D3:70:LYS:NZ	2.34	0.60
31:D9:6:VAL:O	31:D9:8:PHE:N	4.28	0.60
41:L4:144:LYS:HG2	41:L4:145:ILE:H	5.13	0.60
36:1:1334:U:OP1	44:L7:206:LYS:HE3	2.02	0.60
47:M0:174:THR:HG22	47:M0:176:LEU:H	1.66	0.60
51:M5:11:GLN:O	51:M5:14:LYS:NZ	4.09	0.60
51:M5:65:ARG:HB3	51:M5:127:TYR:CD1	4.32	0.60
51:M5:24:ARG:HH11	51:M5:24:ARG:HG2	4.35	0.60
51:M5:44:ARG:HB3	51:M5:47:LYS:HB3	1.83	0.60
52:M6:157:GLU:OE1	52:M6:160:ARG:NH1	2.80	0.60
53:M7:48:LEU:HD22	53:M7:88:VAL:HG13	2.49	0.60
53:M7:48:LEU:HD13	53:M7:88:VAL:HG13	1.83	0.60
68:O2:100:ILE:O	68:O2:105:ARG:NH1	2.32	0.60
75:O9:23:LEU:O	75:O9:25:GLN:NE2	2.34	0.60
9:S7:103:SER:OG	9:S7:104:ARG:N	3.32	0.60
11:S9:131:GLN:O	11:S9:132:ARG:HG2	3.16	0.60
36:1:1129:A:OP1	47:M0:13:LYS:NZ	2.25	0.60
36:1:873:C:H5''	36:1:874:U:O5'	2.02	0.60
1:2:1498:G:OP1	21:C9:75:LYS:HD3	2.02	0.60
36:5:1110:U:H2'	36:5:1111:U:C6	2.36	0.60
36:5:2807:U:O2'	36:5:2809:C:OP1	2.15	0.60
36:5:498:A:O2'	36:5:3273:A:N1	2.29	0.60
36:5:2964:G:N7	86:5:3975:OHX:N6	2.50	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:491:C:N4	1:6:496:G:O6	2.35	0.60
24:D2:96:ALA:HB3	24:D2:99:PHE:CE1	3.27	0.60
30:D8:52:ASP:N	30:D8:52:ASP:OD1	2.34	0.60
32:E0:16:SER:OG	32:E0:17:GLN:N	2.88	0.60
45:L8:86:THR:O	45:L8:90:THR:HG23	5.32	0.60
46:L9:129:ARG:HD2	46:L9:157:ASN:HB2	1.83	0.60
46:L9:176:LEU:O	46:L9:180:TYR:OH	2.58	0.60
37:3:44:C:OP2	48:M1:137:ARG:NH2	2.34	0.60
48:M1:162:TRP:CZ2	48:M1:166:LYS:HD2	2.36	0.60
51:M5:172:ARG:NH1	36:5:29:C:O3'	106.06	0.60
53:M7:116:HIS:NE2	53:M7:147:GLU:OE2	2.31	0.60
54:M8:80:THR:HG22	54:M8:100:THR:HB	1.82	0.60
36:1:1456:A:N7	67:O1:26:LYS:NZ	2.49	0.60
67:O1:44:MET:HB3	67:O1:77:ARG:HD3	1.84	0.60
70:O4:109:THR:HA	70:O4:112:ALA:HB3	2.70	0.60
11:S9:172:VAL:HG22	1:6:511:A:H5''	459.57	0.60
36:1:13:A:H5'	36:1:14:U:OP2	2.02	0.60
36:1:2225:U:H2'	36:1:2226:U:C6	2.36	0.60
1:2:73:U:H1'	1:2:74:U:H5'	1.84	0.60
36:5:656:A:H2'	36:5:657:A:C8	2.37	0.60
1:6:1658:G:H5'	1:6:1659:A:OP2	2.01	0.60
13:C1:109:VAL:HG12	13:C1:137:PHE:HB2	3.62	0.60
14:C2:61:VAL:HG22	14:C2:97:LEU:HD11	1.81	0.60
19:C7:32:LYS:HG3	19:C7:47:ARG:HD3	1.84	0.60
20:C8:14:ILE:H	20:C8:24:GLY:H	1.48	0.60
20:C8:66:LEU:O	20:C8:70:VAL:HG23	2.27	0.60
21:C9:65:ILE:HD13	21:C9:71:VAL:HG23	1.84	0.60
22:D0:15:GLN:O	22:D0:16:GLN:NE2	3.84	0.60
25:D3:16:ARG:HE	25:D3:20:ARG:HH12	4.08	0.60
28:D6:87:ARG:NH2	28:D6:94:ASN:O	2.47	0.60
42:L5:56:THR:O	42:L5:58:LYS:N	2.35	0.60
43:L6:52:VAL:HG11	43:L6:65:ILE:HD12	1.84	0.60
44:L7:27:ALA:O	44:L7:31:ALA:N	2.34	0.60
45:L8:156:ASP:OD2	45:L8:156:ASP:N	2.22	0.60
52:M6:130:LYS:HG3	52:M6:131:PRO:HD2	2.17	0.60
66:O0:99:ASP:O	66:O0:102:THR:N	2.33	0.60
86:1:4194:OHX:N2	86:O1:202:OHX:N5	2.49	0.60
67:O1:82:GLU:O	67:O1:84:ASP:N	2.34	0.60
68:O2:104:ASN:O	68:O2:108:ILE:HG13	2.37	0.60
78:Q2:104:LEU:HD22	78:Q2:104:LEU:H	1.67	0.60
4:S2:82:ASN:OD1	4:S2:83:ILE:N	3.20	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:918:C:C2'	36:1:919:U:H5'	2.32	0.60
1:2:130:C:O2'	1:2:131:C:OP1	2.17	0.60
1:2:1561:U:H2'	1:2:1562:G:H8	1.67	0.60
1:2:209:U:H2'	1:2:210:A:C8	2.37	0.60
1:2:885:G:N2	16:C4:123:SER:HB2	2.14	0.60
36:5:1232:C:C5	36:5:1261:G:H2'	2.36	0.60
36:5:1952:G:N1	36:5:1953:G:N7	2.50	0.60
86:5:3935:OHX:N5	86:5:4228:OHX:N6	2.50	0.60
13:C1:96:LYS:NZ	1:6:374:U:OP1	347.82	0.60
44:L7:180:SER:H	44:L7:183:ASP:HB2	1.66	0.60
47:M0:24:ARG:HH11	47:M0:24:ARG:HG3	1.67	0.60
48:M1:60:ARG:HH11	78:Q2:105:GLN:HA	5.63	0.60
49:M3:75:PHE:H	49:M3:97:VAL:HA	1.76	0.60
50:M4:121:MET:O	50:M4:125:LYS:HG2	2.14	0.60
68:O2:41:VAL:HG12	68:O2:46:PHE:HB2	5.87	0.60
72:O6:26:ILE:HG22	72:O6:29:LYS:HD3	1.83	0.60
4:S2:165:VAL:HG11	4:S2:210:THR:HG23	1.82	0.60
8:S6:14:LYS:HB3	8:S6:124:LEU:HD22	2.57	0.60
34:SR:121:MET:SD	34:SR:183:LEU:HD13	2.42	0.60
34:SR:37:SER:OG	34:SR:38:ARG:N	2.33	0.60
36:1:677:A:H4'	36:1:678:G:O5'	2.02	0.59
36:5:1103:A:H3'	36:5:1104:G:C5'	2.32	0.59
36:5:1284:C:O2'	36:5:1285:G:H5'	2.01	0.59
36:5:3107:U:H2'	36:5:3108:G:H8	1.67	0.59
49:M3:101:ARG:HB2	36:5:76:G:N7	84.46	0.59
1:6:196:G:N3	1:6:197:A:H1'	2.17	0.59
1:6:477:A:H2'	1:6:478:A:H8	1.66	0.59
6:S4:187:ARG:NH2	1:6:753:A:H62	375.50	0.59
1:6:894:U:H2'	1:6:895:G:C8	2.37	0.59
18:C6:42:GLU:HG3	18:C6:43:ILE:HD13	3.69	0.59
20:C8:87:ASN:OD1	20:C8:88:ARG:N	2.35	0.59
29:D7:37:CYS:O	29:D7:39:GLY:N	2.35	0.59
39:L2:152:SER:OG	39:L2:153:GLY:N	2.27	0.59
40:L3:218:ILE:HG13	40:L3:276:THR:HG23	2.75	0.59
47:M0:16:PRO:C	47:M0:18:PRO:HD3	2.22	0.59
69:O3:58:GLU:HG2	69:O3:62:SER:O	3.40	0.59
73:O7:60:GLY:O	86:O7:105:OHX:N6	2.35	0.59
78:Q2:12:CYS:SG	78:Q2:74:CYS:HB2	3.02	0.59
79:Q3:8:VAL:O	79:Q3:11:THR:HB	2.02	0.59
36:1:1352:A:H4'	36:1:1353:U:OP1	2.01	0.59
36:1:255:A:H2'	36:1:256:G:C8	2.35	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:2909:U:O2'	36:1:3105:U:O2	2.16	0.59
1:2:1535:U:H5''	7:S5:187:ILE:HD11	1.83	0.59
36:5:1033:U:H2'	36:5:1034:U:H5'	1.84	0.59
39:L2:69:TYR:CE1	36:5:2557:A:H5''	192.26	0.59
78:Q2:41:ARG:NH2	36:5:2785:A:O2'	159.98	0.59
86:5:3935:OHX:N1	86:5:4228:OHX:N4	2.50	0.59
36:5:600:G:N2	36:5:603:A:OP2	2.34	0.59
41:L4:118:LYS:NZ	36:5:681:U:O4	108.93	0.59
18:C6:125:GLU:HG2	18:C6:126:PRO:HD2	2.23	0.59
18:C6:12:LYS:HG2	18:C6:17:THR:HA	2.63	0.59
20:C8:135:GLY:N	20:C8:136:GLN:OE1	2.35	0.59
1:2:1153:G:H5'	28:D6:85:ARG:HD2	1.82	0.59
41:L4:269:SER:O	41:L4:269:SER:OG	2.32	0.59
43:L6:40:LEU:HD13	43:L6:84:VAL:HG11	1.93	0.59
47:M0:12:GLN:HE21	47:M0:128:ARG:NH1	2.00	0.59
51:M5:68:ARG:NH1	51:M5:128:LYS:HE3	2.73	0.59
55:M9:81:ARG:HD3	55:M9:88:ARG:NH1	4.06	0.59
56:N0:138:GLN:O	56:N0:140:VAL:N	2.74	0.59
61:N5:86:VAL:HG11	61:N5:95:ILE:HD11	1.91	0.59
62:N6:57:LEU:HB3	62:N6:105:VAL:HG12	1.84	0.59
67:O1:47:ASP:O	67:O1:49:VAL:HG23	4.01	0.59
75:O9:3:ALA:O	75:O9:4:GLN:HB2	2.02	0.59
5:S3:66:ILE:O	5:S3:70:THR:OG1	2.96	0.59
7:S5:36:ALA:HB1	7:S5:42:LEU:HD21	1.84	0.59
10:S8:171:SER:HB3	10:S8:180:ASP:H	3.20	0.59
1:2:169:A:H5''	8:S6:176:GLN:HG2	1.85	0.59
36:5:1944:U:H2'	36:5:1945:A:H8	1.68	0.59
36:5:3197:G:H2'	36:5:3198:U:H5''	1.83	0.59
36:5:549:U:O4	86:5:4006:OHX:N4	2.34	0.59
1:6:223:U:H3	1:6:838:G:H1	1.50	0.59
14:C2:31:VAL:HG21	14:C2:136:ILE:HD11	3.53	0.59
17:C5:47:ARG:NH2	1:6:1555:A:OP2	405.33	0.59
17:C5:81:ARG:NH1	17:C5:97:TYR:O	2.35	0.59
17:C5:19:GLY:N	20:C8:93:THR:O	2.36	0.59
44:L7:88:ARG:HD2	44:L7:90:LYS:O	2.02	0.59
46:L9:129:ARG:O	46:L9:132:VAL:HG13	2.40	0.59
57:N1:71:SER:HB2	57:N1:91:LEU:O	3.97	0.59
64:N8:66:ALA:HB1	64:N8:69:TRP:HB2	4.44	0.59
86:1:4194:OHX:N4	86:O1:202:OHX:N1	2.50	0.59
5:S3:38:GLU:OE1	5:S3:40:ARG:NE	2.35	0.59
7:S5:161:ASP:O	30:D8:44:VAL:HA	2.02	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:S6:98:ARG:HD3	8:S6:99:GLY:H	1.67	0.59
36:1:2359:C:H2'	36:1:2360:C:C6	2.37	0.59
36:1:3316:A:O2'	36:1:3317:U:OP2	2.18	0.59
36:1:900:G:H2'	36:1:901:G:C8	2.38	0.59
36:1:1831:U:O2'	38:4:114:G:OP1	2.16	0.59
36:5:1644:C:H5'	36:5:1645:U:H5''	1.83	0.59
36:5:2442:G:H22	36:5:2506:U:H3	1.48	0.59
36:5:2837:A:OP2	36:5:2837:A:H8	1.84	0.59
36:5:283:G:O6	36:5:304:G:H1'	2.03	0.59
36:5:3174:A:H2'	36:5:3175:U:H5'	1.83	0.59
36:5:3224:G:N7	86:5:3991:OHX:N6	2.49	0.59
1:6:245:U:O4	86:6:2122:OHX:N4	2.35	0.59
14:C2:97:LEU:O	14:C2:101:ALA:N	2.35	0.59
17:C5:92:SER:HB2	17:C5:107:ILE:HD11	7.64	0.59
20:C8:145:ARG:HB3	35:SM:68:ARG:NH1	4.71	0.59
2:S0:66:ALA:HB1	23:D1:50:TYR:CD1	3.51	0.59
13:C1:99:ARG:HB2	25:D3:12:ALA:HB2	1.84	0.59
40:L3:114:VAL:HG22	40:L3:163:HIS:CE1	2.90	0.59
41:L4:181:VAL:O	41:L4:183:LYS:N	2.35	0.59
47:M0:61:SER:OG	36:5:2854:U:OP1	292.29	0.59
49:M3:3:ILE:HG12	64:N8:34:MET:CE	2.77	0.59
50:M4:37:GLU:HG2	50:M4:38:ILE:H	1.67	0.59
51:M5:27:VAL:HB	51:M5:122:ASN:ND2	2.16	0.59
54:M8:165:ILE:HD13	54:M8:166:LEU:H	4.46	0.59
58:N2:16:THR:HG23	58:N2:102:GLU:HA	4.12	0.59
2:S0:41:ARG:HE	2:S0:45:VAL:HG21	2.79	0.59
7:S5:121:ILE:HA	7:S5:199:ILE:HD11	1.85	0.59
7:S5:40:ILE:HD13	7:S5:42:LEU:HB3	1.83	0.59
7:S5:41:LYS:HG2	7:S5:69:PHE:CZ	4.80	0.59
8:S6:176:GLN:HG2	1:6:169:A:H5'	328.87	0.59
9:S7:58:LEU:HB2	9:S7:90:VAL:HG22	3.26	0.59
10:S8:36:THR:HA	10:S8:58:LEU:HA	2.46	0.59
36:1:1234:G:H1	36:1:1254:C:H42	1.48	0.59
36:1:2383:C:H2'	36:1:2384:A:H5'	1.83	0.59
36:1:29:C:H4'	36:1:62:A:H4'	1.85	0.59
36:1:1464:G:OP2	86:1:4194:OHX:N5	2.35	0.59
36:1:979:U:H1'	36:1:980:A:N9	2.17	0.59
1:2:1481:C:O2'	1:2:1482:C:O5'	2.19	0.59
1:2:1765:A:H5'	1:2:1767:G:N7	2.18	0.59
1:2:350:U:O2	1:2:352:A:C6	2.56	0.59
42:L5:140:ARG:NH2	36:5:1080:A:OP2	228.77	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:1345:G:N7	86:5:4060:OHX:N5	2.51	0.59
36:5:436:A:OP2	36:5:436:A:H8	1.86	0.59
6:S4:108:ARG:NH2	1:6:789:A:OP1	391.92	0.59
6:S4:6:LYS:HD3	1:6:95:G:OP1	343.05	0.59
16:C4:19:ILE:HG23	16:C4:28:VAL:HG22	1.83	0.59
20:C8:117:LYS:HE2	20:C8:128:PHE:HB2	2.25	0.59
20:C8:4:VAL:HG21	27:D5:82:HIS:CG	3.15	0.59
21:C9:65:ILE:HD13	21:C9:114:VAL:HG11	5.89	0.59
22:D0:106:ILE:HG23	22:D0:107:THR:HG23	1.84	0.59
24:D2:86:ILE:HD12	24:D2:87:GLU:H	1.67	0.59
25:D3:57:LEU:HD11	25:D3:73:ARG:HG3	1.85	0.59
42:L5:196:ARG:NH2	42:L5:237:GLU:OE2	2.28	0.59
41:L4:316:ASN:ND2	44:L7:150:LYS:HG3	2.17	0.59
48:M1:60:ARG:NH1	78:Q2:104:LEU:O	5.00	0.59
50:M4:134:ALA:O	50:M4:136:ALA:N	2.36	0.59
53:M7:126:ARG:HA	53:M7:140:GLU:HG2	2.17	0.59
63:N7:41:ALA:HB2	63:N7:77:TYR:HE1	1.67	0.59
36:1:2643:A:H5'	65:N9:6:ASN:ND2	2.17	0.59
72:O6:9:ILE:HA	72:O6:13:LYS:HG2	1.83	0.59
72:O6:94:ILE:HG22	72:O6:99:ARG:HE	1.68	0.59
76:Q0:97:ARG:HG3	76:Q0:120:GLN:O	2.02	0.59
3:S1:168:ILE:HG12	3:S1:197:ILE:HD12	1.85	0.59
5:S3:59:LEU:HA	5:S3:66:ILE:HG13	1.83	0.59
11:S9:75:ALA:O	11:S9:78:ARG:HB3	3.55	0.59
36:1:2108:C:H1'	36:1:3344:A:C8	2.38	0.59
1:2:1031:U:H4'	1:2:1032:G:OP2	2.01	0.59
1:2:1483:A:H2'	1:2:1484:G:C8	2.38	0.59
1:2:260:U:H3'	1:2:261:U:C5'	2.33	0.59
36:5:2227:C:H2'	36:5:2228:A:C8	2.37	0.59
36:5:567:G:O6	86:5:4126:OHX:N2	2.35	0.59
1:6:1150:G:O6	86:6:2114:OHX:N5	2.35	0.59
8:S6:94:ARG:NH1	1:6:1673:G:OP1	286.37	0.59
32:E0:56:MET:HG2	1:6:556:A:H4'	415.93	0.59
1:6:729:G:O2'	1:6:730:G:O5'	2.20	0.59
17:C5:110:GLU:HG3	20:C8:119:ILE:HD11	1.84	0.59
18:C6:113:ASP:HA	18:C6:116:LEU:HD23	1.83	0.59
18:C6:18:ALA:HB2	18:C6:69:VAL:HG13	2.29	0.59
21:C9:113:ILE:HA	21:C9:128:GLY:HA3	2.70	0.59
23:D1:28:ASP:O	23:D1:31:SER:HB3	3.75	0.59
30:D8:25:VAL:HG11	30:D8:66:LEU:HD12	1.85	0.59
41:L4:229:ASN:OD1	41:L4:230:VAL:N	2.84	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:L5:233:ALA:O	42:L5:235:SER:N	2.35	0.59
61:N5:51:VAL:HG22	71:O5:66:VAL:HG21	2.59	0.59
73:O7:21:ARG:HG3	73:O7:39:TYR:CD2	3.78	0.59
3:S1:128:LYS:NZ	3:S1:132:ASP:HB3	2.17	0.59
4:S2:90:THR:OG1	4:S2:91:ARG:N	4.23	0.59
6:S4:186:GLY:HA3	1:6:753:A:OP1	369.59	0.59
8:S6:126:ASP:OD2	8:S6:127:THR:N	2.35	0.59
9:S7:14:THR:HG22	9:S7:17:GLU:OE1	2.26	0.59
9:S7:86:GLN:HG2	9:S7:87:ASP:H	1.68	0.59
36:1:1723:A:OP2	55:M9:103:ARG:NH2	2.35	0.59
36:1:1204:A:H2	36:1:2834:G:N3	2.00	0.59
36:1:2960:C:H2'	36:1:2961:G:H8	1.67	0.59
36:1:735:A:H2'	36:1:736:A:H8	1.66	0.59
1:2:28:A:H2'	1:2:29:U:C6	2.37	0.59
37:3:106:U:H2'	37:3:107:C:C6	2.38	0.59
38:4:137:C:OP2	86:4:234:OHX:N5	2.36	0.59
36:5:3242:G:H5''	36:5:3245:A:C8	2.37	0.59
1:6:1335:U:H2'	1:6:1336:A:C8	2.37	0.59
25:D3:20:ARG:HD2	1:6:311:U:OP2	326.49	0.59
22:D0:52:LYS:HB2	22:D0:92:ASP:O	2.02	0.59
26:D4:25:VAL:N	26:D4:71:GLY:O	2.30	0.59
51:M5:190:THR:O	51:M5:194:GLN:HG2	2.13	0.59
52:M6:54:TYR:HE2	52:M6:58:LEU:HD13	1.67	0.59
54:M8:141:ARG:HD3	36:5:743:C:O2	174.92	0.59
68:O2:97:ALA:HB3	68:O2:100:ILE:HG12	1.85	0.59
74:O8:26:LYS:HE3	36:5:1751:G:C8	128.34	0.59
3:S1:171:ILE:HD12	3:S1:197:ILE:HD13	1.83	0.59
4:S2:140:ARG:HB2	4:S2:222:TYR:CE1	2.37	0.59
5:S3:67:ASN:O	5:S3:70:THR:OG1	2.18	0.59
11:S9:72:GLU:OE2	1:6:761:G:O2'	398.21	0.59
36:1:3238:G:N7	86:1:3961:OHX:N4	2.51	0.59
36:1:709:A:OP1	54:M8:179:ARG:NH2	2.36	0.59
1:2:1175:U:H2'	1:2:1176:G:C8	2.38	0.59
1:2:66:U:O2	8:S6:160:ARG:NE	2.35	0.59
1:2:703:G:H2'	1:2:704:C:H5'	1.83	0.59
36:5:1785:U:H2'	36:5:1786:G:C8	2.37	0.59
36:5:2250:G:O6	86:5:3940:OHX:N6	2.36	0.59
36:5:618:C:H2'	36:5:619:A:C8	2.37	0.59
1:6:890:C:H2'	1:6:891:A:H8	1.68	0.59
37:7:78:U:OP1	86:7:220:OHX:N5	2.35	0.59
17:C5:126:VAL:HG13	17:C5:127:ARG:H	1.68	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:C9:105:LEU:HB3	21:C9:122:ARG:HH21	1.68	0.59
26:D4:8:ARG:NH1	26:D4:26:ASP:OD1	2.35	0.59
32:E0:55:ARG:NH2	1:6:558:U:OP2	416.94	0.59
33:E1:86:THR:HG23	33:E1:87:THR:H	4.38	0.59
47:M0:47:PRO:HB3	47:M0:171:TRP:CZ2	2.38	0.59
47:M0:216:TYR:CG	47:M0:217:PHE:N	2.70	0.59
49:M3:180:ARG:HD3	72:O6:11:LEU:HD21	1.85	0.59
51:M5:150:TRP:HZ3	51:M5:156:HIS:CD2	2.20	0.59
36:1:670:C:P	54:M8:147:ARG:NH2	2.75	0.59
62:N6:32:SER:HA	62:N6:49:PRO:HA	2.31	0.59
62:N6:71:SER:OG	62:N6:72:SER:N	2.33	0.59
54:M8:157:PRO:HG3	64:N8:47:LYS:HB2	1.83	0.59
67:O1:26:LYS:HE2	67:O1:64:VAL:HG11	5.15	0.59
19:C7:33:ARG:HD2	34:SR:109:ASP:OD2	2.03	0.59
34:SR:123:ILE:HG22	34:SR:133:VAL:HG22	1.92	0.59
34:SR:179:LYS:HG2	34:SR:191:ASP:OD1	2.03	0.59
36:1:1593:A:N3	36:1:1615:C:O2'	2.35	0.59
36:1:2443:A:O2'	36:1:2444:C:OP2	2.17	0.59
36:1:2679:A:O2'	48:M1:52:TYR:OH	2.19	0.59
36:1:290:G:H2'	36:1:291:C:C6	2.37	0.59
36:1:1942:U:O2'	36:1:3345:G:O2'	2.17	0.59
36:1:425:G:O6	86:1:3867:OHX:N6	2.36	0.59
1:2:1591:C:H2'	1:2:1592:A:C8	2.37	0.59
13:C1:21:ASN:HB3	13:C1:32:LYS:HD3	5.66	0.59
18:C6:46:PHE:HA	18:C6:49:TYR:HB2	2.30	0.59
20:C8:116:LEU:HD23	20:C8:116:LEU:H	3.02	0.59
20:C8:72:ILE:HA	20:C8:79:TYR:CE2	4.02	0.59
21:C9:105:LEU:HD13	21:C9:122:ARG:NE	2.18	0.59
42:L5:134:ALA:HB2	42:L5:141:PRO:HD3	2.35	0.59
63:N7:26:VAL:O	63:N7:93:LYS:NZ	2.30	0.59
49:M3:170:LEU:HD11	64:N8:147:LEU:HD21	1.85	0.59
78:Q2:28:TYR:HB3	78:Q2:69:VAL:HB	2.09	0.59
9:S7:10:SER:HB3	9:S7:43:PHE:O	2.03	0.59
36:1:3384:U:H2'	36:1:3385:U:C6	2.38	0.59
1:2:1537:C:N4	1:2:1572:G:H1	2.01	0.59
1:2:545:A:H4'	1:2:546:U:OP1	2.03	0.59
1:2:647:G:N2	1:2:687:G:H22	2.01	0.59
1:2:740:A:H2'	1:2:741:C:H5''	1.85	0.59
37:3:28:C:C4	37:3:29:C:C2	2.91	0.59
36:5:653:A:H1'	36:5:2360:C:C2	2.38	0.59
36:5:2322:C:OP1	86:5:4155:OHX:N6	2.36	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
86:5:4014:OHX:N5	86:5:4212:OHX:N1	2.50	0.59
36:5:2211:U:OP2	86:5:4218:OHX:N1	2.36	0.59
1:6:138:A:H2'	1:6:139:C:H5'	1.85	0.59
77:Q1:3:ALA:HB3	1:6:1773:C:OP1	313.15	0.59
37:7:9:C:H5''	37:7:10:C:OP2	2.03	0.59
16:C4:85:ALA:HB2	16:C4:94:PRO:HA	2.40	0.59
19:C7:7:LYS:O	19:C7:11:ARG:HB2	2.03	0.59
22:D0:80:GLU:OE1	31:D9:44:ARG:HD2	4.05	0.59
24:D2:29:PRO:O	24:D2:30:SER:HB3	2.02	0.59
30:D8:12:VAL:HA	30:D8:30:VAL:HG12	1.85	0.59
43:L6:97:ASN:OD1	43:L6:99:GLU:HG2	2.02	0.59
44:L7:222:HIS:ND1	44:L7:224:ILE:HG13	2.45	0.59
46:L9:90:MET:O	46:L9:91:ARG:HG2	3.39	0.59
48:M1:149:GLY:O	48:M1:153:LYS:HD2	4.39	0.59
50:M4:22:LEU:HD13	50:M4:32:LEU:HD23	1.95	0.59
52:M6:186:ALA:O	52:M6:187:GLU:HB2	2.01	0.59
44:L7:109:THR:HG22	54:M8:4:ASP:HB3	1.85	0.59
65:N9:16:ALA:O	65:N9:20:GLY:HA3	3.89	0.59
70:O4:43:LYS:HD3	70:O4:50:ALA:HB2	1.85	0.59
2:S0:182:LEU:HB3	2:S0:188:LEU:HD23	1.85	0.59
6:S4:252:ARG:NH2	6:S4:252:ARG:HB3	4.68	0.59
10:S8:82:VAL:HG12	10:S8:101:ILE:HG22	2.44	0.59
36:1:1783:U:H2'	36:1:1784:G:H8	1.68	0.58
36:1:2875:U:H2'	36:1:2875:U:O2	2.03	0.58
36:1:3242:G:N2	36:1:3245:A:OP2	2.35	0.58
36:1:860:G:C5	39:L2:181:LYS:HB2	2.38	0.58
36:1:975:C:H2'	36:1:976:U:C6	2.37	0.58
1:2:278:U:H4'	1:2:279:G:O5'	2.01	0.58
36:5:2177:G:O6	86:5:3968:OHX:N1	2.35	0.58
1:6:1498:G:H1	1:6:1509:C:H42	1.51	0.58
1:6:1324:G:N7	86:6:2103:OHX:N2	2.51	0.58
1:6:322:G:OP1	86:6:2106:OHX:N5	2.36	0.58
20:C8:42:TYR:HE2	20:C8:73:MET:HG3	4.15	0.58
27:D5:54:VAL:HA	27:D5:57:TYR:CZ	3.43	0.58
47:M0:194:GLY:HA3	36:5:1010:G:N3	336.24	0.58
51:M5:65:ARG:HD2	51:M5:129:TYR:HE1	1.67	0.58
37:3:98:C:OP1	56:N0:39:SER:OG	2.20	0.58
62:N6:47:ALA:O	62:N6:122:LYS:NZ	4.05	0.58
77:Q1:23:ARG:NH2	36:5:2303:A:OP1	270.86	0.58
4:S2:108:ASN:O	4:S2:108:ASN:ND2	2.35	0.58
9:S7:69:GLY:HA2	9:S7:72:LYS:HD2	1.85	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:132:C:H2'	36:1:133:U:H5''	1.86	0.58
36:1:1569:U:H5''	36:1:1570:U:C6	2.37	0.58
1:2:1537:C:H42	1:2:1572:G:H22	1.51	0.58
1:2:1169:G:O2'	1:2:1576:A:N6	2.35	0.58
1:2:187:G:H4'	1:2:188:A:OP1	2.02	0.58
1:2:623:A:OP1	86:2:2157:OHX:N2	2.36	0.58
44:L7:196:LYS:HE3	36:5:1100:U:OP2	245.36	0.58
60:N4:17:ARG:HE	36:5:3050:U:H5''	239.38	0.58
36:5:1170:A:OP2	86:5:3995:OHX:N6	2.36	0.58
1:6:1159:C:H5''	1:6:1160:A:H5'	1.83	0.58
1:6:872:G:H2'	1:6:873:U:O4'	2.04	0.58
13:C1:97:TYR:O	13:C1:99:ARG:HG2	2.03	0.58
29:D7:36:LYS:HB3	29:D7:43:ILE:HG22	1.85	0.58
41:L4:5:GLN:NE2	41:L4:21:PRO:HB3	2.18	0.58
42:L5:182:GLY:HA2	42:L5:194:LEU:HD13	2.50	0.58
61:N5:57:LEU:HA	61:N5:61:LYS:HG2	3.79	0.58
70:O4:44:CYS:HA	70:O4:51:LEU:HD21	5.03	0.58
71:O5:47:VAL:O	71:O5:51:ILE:HG13	2.31	0.58
6:S4:52:LEU:HB3	6:S4:54:TYR:CD2	2.33	0.58
6:S4:50:ASN:O	6:S4:53:LYS:NZ	2.35	0.58
10:S8:97:THR:OG1	10:S8:98:LYS:O	3.19	0.58
34:SR:22:SER:OG	34:SR:70:ASP:HA	2.91	0.58
36:1:2274:U:OP2	86:1:3958:OHX:N4	2.36	0.58
1:2:1754:A:H4'	1:2:1755:A:O4'	2.03	0.58
1:2:248:U:OP1	86:2:2092:OHX:N6	2.36	0.58
1:2:103:A:O3'	1:2:308:C:N4	2.36	0.58
1:2:375:U:OP1	25:D3:23:ARG:NH2	2.36	0.58
1:2:383:G:N7	86:2:2130:OHX:N4	2.51	0.58
1:2:56:U:H4'	1:2:57:G:H5'	1.85	0.58
1:2:759:U:OP1	86:2:2160:OHX:N1	2.36	0.58
36:5:1348:U:H5''	36:5:1355:A:N6	2.18	0.58
36:5:1500:G:H2'	36:5:1501:U:O4'	2.03	0.58
36:5:2705:A:OP2	86:5:3892:OHX:N2	2.36	0.58
36:5:108:A:O2'	36:5:323:A:N1	2.37	0.58
36:5:864:G:OP2	86:5:3909:OHX:N4	2.36	0.58
1:6:67:A:N6	1:6:83:G:O2'	2.36	0.58
1:2:1241:G:OP1	17:C5:77:ARG:NH2	2.35	0.58
19:C7:72:LYS:O	19:C7:75:GLU:N	2.35	0.58
28:D6:43:ASN:HB3	28:D6:46:GLU:H	4.15	0.58
39:L2:90:ALA:HB2	39:L2:101:VAL:HG13	3.07	0.58
43:L6:40:LEU:HB3	43:L6:84:VAL:HG11	2.35	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:M3:75:PHE:O	49:M3:79:GLU:HB2	2.17	0.58
59:N3:48:ARG:HH22	36:5:3043:C:P	250.75	0.58
64:N8:116:GLY:HA2	64:N8:137:LYS:NZ	2.18	0.58
64:N8:126:LYS:HA	64:N8:146:GLU:O	2.04	0.58
66:O0:13:LYS:O	66:O0:17:VAL:HG23	2.10	0.58
74:O8:48:SER:OG	74:O8:49:SER:N	4.56	0.58
79:Q3:36:ARG:NH2	79:Q3:46:THR:HG22	2.18	0.58
3:S1:183:GLN:O	3:S1:187:LYS:N	2.36	0.58
6:S4:240:LYS:HE2	6:S4:240:LYS:H	1.68	0.58
6:S4:9:LEU:HB3	6:S4:30:ARG:HB2	3.74	0.58
7:S5:36:ALA:HB1	7:S5:42:LEU:HD12	4.89	0.58
9:S7:46:ILE:HA	9:S7:59:ALA:O	2.79	0.58
11:S9:109:LEU:CB	11:S9:146:PHE:HB3	2.33	0.58
34:SR:22:SER:OG	34:SR:69:GLN:O	6.48	0.58
36:1:1840:U:OP2	86:1:3971:OHX:N5	2.36	0.58
36:1:1913:A:N3	36:1:2120:A:H2'	2.19	0.58
36:1:1317:A:OP1	86:1:4059:OHX:N2	2.37	0.58
1:2:1041:G:OP1	86:2:2149:OHX:N5	2.36	0.58
1:2:1469:A:H2'	1:2:1470:C:C6	2.39	0.58
1:2:66:U:H1'	8:S6:160:ARG:HH21	1.69	0.58
39:L2:156:LYS:NZ	36:5:2158:A:OP2	205.11	0.58
36:5:2661:G:H1	36:5:2709:C:H42	1.52	0.58
36:5:712:G:H2'	36:5:713:U:C6	2.38	0.58
1:6:1524:A:H2'	1:6:1525:A:C8	2.38	0.58
21:C9:57:ARG:HG2	21:C9:104:VAL:HG21	1.85	0.58
23:D1:3:ASN:OD1	23:D1:7:GLN:HB3	3.28	0.58
46:L9:109:ALA:HB1	46:L9:111:PHE:CE2	2.38	0.58
48:M1:92:ARG:HG3	48:M1:172:LEU:HB2	5.20	0.58
49:M3:52:ASP:OD1	49:M3:52:ASP:N	2.35	0.58
36:1:1874:A:H5''	55:M9:18:GLY:HA3	1.85	0.58
57:N1:69:LYS:HE2	57:N1:70:SER:HB3	4.98	0.58
63:N7:89:VAL:O	63:N7:91:ALA:N	3.04	0.58
2:S0:103:THR:O	2:S0:106:SER:OG	2.42	0.58
3:S1:128:LYS:HE3	3:S1:132:ASP:OD1	2.03	0.58
36:1:1595:U:H4'	36:1:1595:U:OP1	2.02	0.58
36:1:1942:U:HO2'	36:1:3345:G:HO2'	1.50	0.58
36:1:799:G:O6	86:1:3974:OHX:N5	2.36	0.58
1:2:778:G:H22	26:D4:10:ARG:NH1	2.02	0.58
1:2:802:G:H21	24:D2:107:SER:HB3	1.67	0.58
38:4:103:G:O6	86:4:226:OHX:N4	2.36	0.58
36:5:1241:U:O2'	36:5:1242:G:O5'	2.19	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:2439:A:N6	36:5:2508:U:H3	2.01	0.58
36:5:2698:G:H2'	36:5:2699:G:H8	1.67	0.58
36:5:2794:G:H1'	36:5:2795:U:C6	2.39	0.58
36:5:2895:G:H2'	36:5:2896:A:H5''	1.86	0.58
1:6:1017:U:H2'	1:6:1018:U:H6	1.69	0.58
21:C9:33:TYR:HD1	21:C9:37:VAL:HG21	4.62	0.58
27:D5:92:ILE:HD11	27:D5:102:THR:OG1	6.35	0.58
39:L2:32:LEU:HD13	39:L2:37:ARG:HD3	1.84	0.58
40:L3:43:LEU:HG	40:L3:181:ILE:HG21	2.49	0.58
37:3:62:U:OP1	42:L5:277:LEU:HB2	2.04	0.58
42:L5:86:TYR:HB3	42:L5:246:ALA:HB3	1.85	0.58
46:L9:174:LYS:HB2	76:Q0:127:LEU:HD11	1.85	0.58
49:M3:166:ALA:N	64:N8:135:GLU:OE1	3.86	0.58
52:M6:84:LEU:HD13	52:M6:102:LEU:HD21	1.85	0.58
62:N6:112:ASP:HB2	62:N6:115:ARG:HB2	1.85	0.58
69:O3:75:HIS:HB2	69:O3:82:ARG:HG3	4.08	0.58
74:O8:58:ASP:OD2	74:O8:60:GLY:N	2.36	0.58
78:Q2:17:CYS:HG	78:Q2:74:CYS:HG	1.44	0.58
6:S4:139:VAL:HG13	6:S4:150:PRO:HG3	1.84	0.58
11:S9:49:LEU:HA	11:S9:52:ILE:HD12	2.96	0.58
36:1:2121:G:H2'	36:1:2122:G:H4'	1.84	0.58
36:1:263:C:H2'	36:1:264:G:O4'	2.03	0.58
36:1:73:C:O2	49:M3:59:ARG:HD3	2.04	0.58
1:2:1157:A:O2'	1:2:1158:C:OP1	2.16	0.58
1:2:1231:U:H4'	1:2:1258:U:H6	1.68	0.58
1:2:190:C:N4	1:2:196:G:C6	2.72	0.58
1:2:930:A:OP1	28:D6:32:LYS:NZ	2.25	0.58
1:2:959:U:H5'	29:D7:28:PRO:HB3	1.84	0.58
36:5:1593:A:N3	36:5:1615:C:O2'	2.34	0.58
36:5:1701:C:H2'	36:5:1702:U:O4'	2.03	0.58
36:5:2136:C:O2'	36:5:2137:U:H5'	2.02	0.58
42:L5:36:LEU:HD23	36:5:2748:A:N3	254.67	0.58
1:6:116:U:H2'	1:6:117:U:C6	2.37	0.58
1:6:513:U:H2'	1:6:514:G:C8	2.37	0.58
19:C7:14:LYS:HG3	19:C7:69:ILE:HG22	4.07	0.58
20:C8:52:VAL:HG21	20:C8:69:ILE:HD11	1.85	0.58
29:D7:59:CYS:O	29:D7:61:THR:N	2.72	0.58
40:L3:171:LEU:HD21	40:L3:333:LYS:HG2	1.86	0.58
41:L4:361:HIS:CG	41:L4:362:ASP:N	2.90	0.58
41:L4:82:THR:HG23	41:L4:84:ARG:N	4.17	0.58
37:3:11:A:H8	42:L5:18:THR:HG1	1.47	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:L7:34:LYS:HA	44:L7:37:ASN:HB2	1.85	0.58
49:M3:133:PRO:O	49:M3:135:ALA:N	3.42	0.58
55:M9:70:LYS:HG3	55:M9:75:HIS:HB2	2.94	0.58
57:N1:2:GLY:N	36:5:2626:A:O5'	233.15	0.58
57:N1:88:ARG:HD3	65:N9:33:LYS:NZ	4.55	0.58
69:O3:2:ALA:HB2	36:5:3216:G:OP2	264.75	0.58
64:N8:147:LEU:HD12	72:O6:7:ILE:HD11	7.09	0.58
4:S2:187:LEU:HD23	4:S2:188:LEU:HD23	4.21	0.58
10:S8:8:ARG:NH2	10:S8:19:ALA:O	2.37	0.58
11:S9:134:ILE:HA	11:S9:158:PHE:HA	1.84	0.58
36:1:619:A:H5''	36:1:620:U:OP1	2.04	0.58
1:2:2:A:H5'	1:2:2:A:H8	1.69	0.58
36:5:1401:A:H2'	36:5:1402:C:H6	1.68	0.58
36:5:1711:C:H2'	36:5:1712:G:O4'	2.03	0.58
36:5:2271:A:H2'	36:5:2272:G:O4'	2.04	0.58
36:5:2818:U:C6	36:5:2818:U:H5'	2.38	0.58
36:5:3299:A:H61	36:5:3315:G:H1	1.50	0.58
1:6:1535:U:O2'	1:6:1536:G:O5'	2.20	0.58
1:6:333:A:C6	1:6:334:G:C6	2.91	0.58
1:6:67:A:O2'	1:6:69:G:OP1	2.10	0.58
15:C3:63:ALA:HB3	15:C3:71:ILE:HG12	3.35	0.58
16:C4:29:HIS:HB3	16:C4:41:ARG:HG3	1.86	0.58
20:C8:112:ASP:O	20:C8:115:ARG:HB3	2.38	0.58
22:D0:95:ALA:HB1	22:D0:99:ILE:HG21	1.84	0.58
51:M5:35:VAL:HG13	51:M5:65:ARG:HB3	1.86	0.58
52:M6:89:SER:O	52:M6:91:LYS:N	2.57	0.58
54:M8:64:VAL:HG22	54:M8:96:PHE:CE2	2.38	0.58
58:N2:56:VAL:HG22	58:N2:65:VAL:HG22	1.85	0.58
60:N4:46:PRO:HB2	60:N4:54:LEU:HD22	1.84	0.58
60:N4:9:SER:O	60:N4:9:SER:OG	3.82	0.58
61:N5:75:LYS:HD3	61:N5:123:TYR:HE1	1.69	0.58
39:L2:48:ILE:HD11	79:Q3:54:ILE:HG12	1.86	0.58
4:S2:140:ARG:HH12	23:D1:1:MET:HB3	1.67	0.58
4:S2:156:THR:HA	24:D2:96:ALA:HB2	2.66	0.58
5:S3:44:THR:HG22	5:S3:45:LYS:HG3	1.85	0.58
6:S4:195:ILE:HA	6:S4:210:ILE:HD13	1.85	0.58
36:1:3243:A:C8	52:M6:156:LEU:HD22	2.38	0.58
36:1:300:G:O6	86:1:4147:OHX:N1	2.36	0.58
1:2:1504:G:H2'	1:2:1505:A:C8	2.38	0.58
1:2:1657:U:H5	36:1:2125:A:O3'	1.87	0.58
86:2:2031:OHX:N6	86:2:2146:OHX:N2	2.51	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:306:U:H2'	1:2:307:G:H8	1.69	0.58
1:2:990:C:O2'	16:C4:127:ARG:HG2	2.03	0.58
56:N0:90:MET:CG	36:5:1213:G:H4'	317.20	0.58
36:5:1222:G:H8	36:5:1222:G:OP2	1.87	0.58
36:5:2991:A:O2'	36:5:3309:G:N7	2.36	0.58
59:N3:14:SER:OG	36:5:3094:A:OP1	251.62	0.58
50:M4:128:ARG:NH2	36:5:3214:U:OP2	280.37	0.58
36:5:499:G:H2'	36:5:500:C:C6	2.39	0.58
1:6:1273:G:O5'	1:6:1274:C:H3'	2.04	0.58
18:C6:82:ARG:NH2	18:C6:114:ARG:HB3	2.18	0.58
1:2:1367:G:O5'	21:C9:7:ARG:NH1	2.36	0.58
24:D2:5:SER:O	24:D2:7:LEU:N	3.88	0.58
39:L2:3:ARG:HG2	39:L2:4:VAL:H	1.69	0.58
47:M0:86:HIS:HB3	47:M0:139:ARG:CG	2.67	0.58
49:M3:83:ALA:HB2	49:M3:113:VAL:HG12	1.85	0.58
36:1:1544:G:H5'	51:M5:67:ARG:HE	1.69	0.58
53:M7:127:ARG:O	53:M7:139:TYR:N	2.36	0.58
36:1:1722:U:OP1	55:M9:100:ARG:HD3	2.03	0.58
67:O1:16:LEU:O	67:O1:20:LEU:N	2.59	0.58
67:O1:26:LYS:HA	67:O1:64:VAL:HG21	3.36	0.58
39:L2:80:GLU:OE1	79:Q3:73:THR:HG22	2.03	0.58
36:1:1108:U:H2'	36:1:1109:U:C6	2.38	0.58
36:1:2216:G:OP1	72:O6:75:LYS:NZ	2.30	0.58
36:1:2366:C:H42	36:1:2381:G:H1	1.51	0.58
36:1:3181:C:H2'	36:1:3182:G:H8	1.69	0.58
36:1:709:A:H8	36:1:709:A:O5'	1.87	0.58
1:2:1002:G:N2	1:2:1760:G:O3'	2.35	0.58
37:3:112:G:H2'	37:3:113:C:C6	2.38	0.58
36:5:2207:A:H2'	36:5:2208:A:O4'	2.04	0.58
36:5:268:A:O2'	36:5:269:G:OP2	2.20	0.58
36:5:2921:U:H2'	36:5:2923:U:OP2	2.04	0.58
38:8:126:A:O2'	38:8:128:U:OP2	2.16	0.58
13:C1:6:THR:O	13:C1:8:GLN:N	2.37	0.58
14:C2:131:ASP:HB2	14:C2:132:GLU:CD	2.25	0.58
14:C2:33:ARG:HA	14:C2:36:LEU:HD12	1.84	0.58
1:2:1795:U:O4	28:D6:9:GLY:HA2	2.04	0.58
42:L5:166:ALA:HB1	42:L5:171:LEU:HD12	1.85	0.58
51:M5:9:GLU:HG3	72:O6:44:VAL:HG11	1.85	0.58
36:1:3268:A:OP2	53:M7:181:ARG:NH1	2.37	0.58
41:L4:302:ALA:HB2	54:M8:39:ARG:NH1	2.54	0.58
70:O4:99:LYS:O	70:O4:103:LYS:HG2	2.11	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
71:O5:21:LEU:HD11	71:O5:25:LYS:HE3	2.80	0.58
72:O6:25:LYS:HB2	72:O6:28:TYR:HD2	1.69	0.58
5:S3:75:LYS:HB3	12:C0:22:VAL:HG22	2.76	0.58
9:S7:98:ILE:HD13	9:S7:118:LEU:HD22	1.85	0.58
11:S9:159:ALA:HB3	11:S9:162:SER:HB3	3.90	0.58
1:2:577:G:C6	35:SM:99:LYS:HD3	2.39	0.58
34:SR:214:ALA:HB2	34:SR:220:ILE:HA	1.85	0.58
36:1:1144:U:OP1	36:1:1367:G:O2'	2.18	0.58
36:1:1901:A:H5''	36:1:1902:G:OP2	2.04	0.58
1:2:1022:C:O2'	1:2:1125:A:N1	2.34	0.58
1:2:1146:G:C6	1:2:1147:A:C6	2.92	0.58
1:2:1427:A:OP2	35:SM:93:ARG:NH1	2.36	0.58
1:2:66:U:H5	8:S6:173:PRO:HG3	1.69	0.58
37:3:80:G:OP2	86:3:224:OHX:N6	2.37	0.58
38:4:62:C:O2	86:4:229:OHX:N5	2.37	0.58
58:N2:104:ARG:NH2	36:5:1758:G:H5'	119.93	0.58
1:6:1238:A:H2'	1:6:1239:U:H5'	1.84	0.58
1:6:500:C:O2'	1:6:501:U:O4'	2.22	0.58
14:C2:50:LYS:O	14:C2:54:ARG:HG2	2.04	0.58
14:C2:54:ARG:O	14:C2:85:LYS:NZ	2.35	0.58
15:C3:116:ILE:O	15:C3:120:SER:OG	2.22	0.58
19:C7:52:GLY:O	19:C7:55:THR:OG1	4.08	0.58
21:C9:86:ARG:NH1	21:C9:90:PRO:O	2.52	0.58
47:M0:3:ARG:NH2	36:5:2854:U:OP2	290.96	0.58
47:M0:42:THR:HG23	47:M0:45:GLU:HB2	2.31	0.58
49:M3:56:PRO:HG3	49:M3:74:GLY:O	2.05	0.58
51:M5:150:TRP:CH2	51:M5:151:ILE:HD12	4.56	0.58
54:M8:33:TYR:HA	54:M8:36:LEU:HB2	1.86	0.58
54:M8:62:VAL:HB	54:M8:83:VAL:HG11	3.08	0.58
67:O1:78:LYS:HG2	67:O1:79:ARG:HH21	1.69	0.58
75:O9:3:ALA:O	75:O9:5:LYS:N	5.07	0.58
4:S2:101:VAL:HG23	4:S2:115:ILE:HG23	2.67	0.58
7:S5:64:VAL:HG23	7:S5:89:ILE:HD11	1.86	0.58
36:1:120:G:N2	45:L8:126:SER:HB2	2.19	0.57
36:1:1321:G:O3'	56:N0:117:ARG:NH2	2.37	0.57
36:1:1355:A:H4'	36:1:1356:U:O5'	2.03	0.57
36:1:1420:C:OP2	41:L4:193:LYS:NZ	2.23	0.57
36:1:53:G:OP1	73:O7:48:ASN:HB2	2.04	0.57
1:2:992:A:H2	1:2:1012:U:H3	1.48	0.57
1:2:1370:U:H4'	1:2:1371:A:O5'	2.03	0.57
1:2:190:C:N4	1:2:196:G:O6	2.37	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:350:C:N3	36:5:368:G:H5'	2.18	0.57
1:6:250:C:H2'	1:6:251:A:C8	2.39	0.57
1:6:607:G:H5'	1:6:613:G:N2	2.17	0.57
15:C3:16:ILE:HD12	1:6:959:U:H4'	347.47	0.57
39:L2:137:ILE:HG12	39:L2:147:ARG:HG3	4.19	0.57
41:L4:35:VAL:HG21	41:L4:244:LEU:HD21	1.84	0.57
42:L5:279:LYS:HD3	42:L5:282:ARG:HH22	5.62	0.57
44:L7:214:TRP:CE2	44:L7:219:LYS:HD2	2.39	0.57
44:L7:132:PRO:HA	44:L7:229:PHE:CD2	2.68	0.57
66:O0:30:THR:HG21	66:O0:89:VAL:HG22	2.97	0.57
43:L6:13:GLU:OE2	68:O2:91:THR:HB	4.71	0.57
69:O3:71:VAL:HA	69:O3:83:ALA:HB2	1.85	0.57
73:O7:72:ARG:NH1	38:8:95:G:OP2	52.41	0.57
4:S2:106:ASP:OD1	4:S2:107:SER:N	2.52	0.57
4:S2:225:LEU:HD13	24:D2:68:ARG:HA	1.84	0.57
11:S9:113:VAL:HG21	11:S9:134:ILE:HG21	3.82	0.57
11:S9:58:ASP:O	11:S9:61:THR:OG1	4.01	0.57
36:1:3269:U:H5'	36:1:3269:U:O2	2.05	0.57
86:1:4194:OHX:N4	86:O1:202:OHX:N3	2.52	0.57
1:2:1208:A:H5'	1:2:1209:C:OP2	2.04	0.57
1:2:1489:U:OP2	5:S3:9:ARG:NH2	2.38	0.57
1:2:1761:U:O2'	1:2:1762:A:OP2	2.20	0.57
36:5:1228:C:H2'	36:5:1229:G:C8	2.38	0.57
55:M9:143:ILE:HG12	36:5:2093:A:P	251.99	0.57
36:5:2115:G:H22	36:5:2120:A:H1'	1.69	0.57
54:M8:93:ILE:HG23	36:5:784:A:C6	149.55	0.57
32:E0:37:ARG:NH1	1:6:478:A:OP1	441.12	0.57
1:6:489:C:O2'	1:6:490:C:O5'	2.22	0.57
15:C3:15:ALA:O	1:6:959:U:H5''	352.06	0.57
20:C8:99:HIS:HD2	20:C8:101:LEU:HD21	1.69	0.57
5:S3:8:LYS:HG2	22:D0:63:LEU:HD21	2.55	0.57
40:L3:88:GLY:O	40:L3:161:LEU:N	2.52	0.57
41:L4:271:LYS:HB2	41:L4:274:TYR:HB2	2.31	0.57
46:L9:105:GLU:OE2	46:L9:108:GLY:HA2	2.04	0.57
49:M3:18:TRP:C	49:M3:20:GLU:H	2.08	0.57
54:M8:96:PHE:CG	54:M8:97:PRO:HD2	2.48	0.57
58:N2:50:LEU:HB3	58:N2:54:VAL:HG23	1.87	0.57
5:S3:166:ASP:O	5:S3:190:ARG:NH2	2.82	0.57
8:S6:164:LYS:N	8:S6:167:LYS:O	2.31	0.57
8:S6:185:GLN:HA	8:S6:188:ARG:NH1	2.19	0.57
9:S7:132:PRO:O	9:S7:133:THR:OG1	4.86	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:S9:114:TYR:HA	11:S9:119:ALA:HB3	1.86	0.57
11:S9:123:HIS:CE1	32:E0:37:ARG:HD2	3.65	0.57
36:1:1844:C:O2	73:O7:9:GLY:HA2	2.03	0.57
36:1:582:G:O6	86:1:4169:OHX:N2	2.37	0.57
36:1:646:A:H2'	36:1:647:A:O4'	2.04	0.57
36:1:824:C:H2'	36:1:825:U:C6	2.39	0.57
1:2:647:G:H2'	1:2:648:G:H8	1.69	0.57
36:5:3191:G:O6	86:5:4139:OHX:N6	2.37	0.57
36:5:438:A:H2'	36:5:494:G:N2	2.19	0.57
1:6:484:C:N4	1:6:503:G:H22	2.02	0.57
28:D6:46:GLU:HG2	28:D6:49:ALA:HB2	1.85	0.57
41:L4:5:GLN:HE22	41:L4:21:PRO:HB3	1.69	0.57
42:L5:9:SER:OG	42:L5:10:SER:N	2.36	0.57
46:L9:20:ILE:HD12	46:L9:45:PHE:CD1	2.39	0.57
66:O0:31:VAL:HA	66:O0:34:LEU:HB2	1.86	0.57
67:O1:44:MET:O	67:O1:46:THR:HG22	4.14	0.57
72:O6:54:GLU:O	72:O6:58:ILE:HG23	2.05	0.57
78:Q2:99:GLN:HG2	78:Q2:100:LYS:H	1.70	0.57
10:S8:140:GLU:HA	10:S8:143:TRP:HB2	3.12	0.57
11:S9:163:PRO:HG2	11:S9:164:PHE:HD2	1.70	0.57
11:S9:17:ARG:O	11:S9:23:ARG:NH2	2.37	0.57
36:1:1362:G:H4'	44:L7:159:GLN:O	2.05	0.57
36:1:1874:A:OP2	55:M9:20:ARG:NH1	2.35	0.57
1:2:321:C:N4	1:2:1667:A:OP1	2.37	0.57
1:2:186:C:H3'	1:2:187:G:H8	1.70	0.57
57:N1:129:LYS:HB2	36:5:1098:A:O5'	252.25	0.57
51:M5:4:TYR:OH	36:5:148:G:OP2	110.18	0.57
36:5:2362:C:H42	36:5:2376:G:H1	1.50	0.57
39:L2:230:VAL:HG21	36:5:2424:A:N1	183.70	0.57
36:5:731:U:H2'	36:5:732:C:C6	2.40	0.57
18:C6:14:LYS:HE2	1:6:1584:G:N7	396.48	0.57
1:6:922:G:H2'	1:6:923:A:H8	1.70	0.57
12:C0:41:TYR:O	12:C0:45:ALA:N	2.91	0.57
15:C3:88:LEU:HG	15:C3:125:LEU:HD13	2.50	0.57
18:C6:143:ARG:NH1	1:6:1191:U:H5'	350.58	0.57
1:2:1383:G:H1'	22:D0:57:ARG:NH1	2.20	0.57
14:C2:50:LYS:NZ	33:E1:129:GLY:O	2.25	0.57
47:M0:117:GLY:O	86:M0:304:OHX:N3	2.37	0.57
49:M3:135:ALA:O	49:M3:136:GLU:HB3	2.04	0.57
51:M5:65:ARG:HG2	51:M5:127:TYR:CD1	2.39	0.57
51:M5:21:PHE:HD2	51:M5:22:LEU:HD13	2.26	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:S6:158:ILE:HG23	60:N4:85:ALA:HB2	2.83	0.57
70:O4:56:THR:O	70:O4:56:THR:OG1	2.17	0.57
38:4:58:G:N7	73:O7:63:ARG:NH1	2.50	0.57
75:O9:6:SER:OG	75:O9:9:ILE:HG12	2.05	0.57
76:Q0:110:CYS:SG	76:Q0:111:ARG:N	3.16	0.57
78:Q2:47:GLN:NE2	78:Q2:53:GLN:OE1	2.78	0.57
5:S3:179:GLN:O	5:S3:179:GLN:NE2	2.36	0.57
8:S6:122:GLU:O	8:S6:124:LEU:N	2.63	0.57
8:S6:137:ARG:O	8:S6:141:ILE:HD12	2.52	0.57
8:S6:171:LYS:HZ3	1:6:68:A:P	349.40	0.57
36:1:2725:U:H5''	36:1:2726:C:OP2	2.04	0.57
36:1:3316:A:OP1	36:1:3318:G:N2	2.38	0.57
1:2:1788:G:P	16:C4:127:ARG:HH12	2.27	0.57
36:5:1152:G:N2	36:5:1200:A:H61	2.02	0.57
36:5:869:G:N2	36:5:890:C:O2	2.37	0.57
1:6:1175:U:H2'	1:6:1176:G:C8	2.40	0.57
1:6:45:U:H5''	1:6:46:A:OP2	2.05	0.57
1:6:846:G:H2'	1:6:847:A:O4'	2.05	0.57
18:C6:24:ALA:HB2	18:C6:92:TYR:OH	2.05	0.57
22:D0:96:PRO:O	22:D0:100:VAL:HG23	2.26	0.57
25:D3:14:LYS:HZ1	1:6:1105:C:P	326.03	0.57
25:D3:14:LYS:HE3	25:D3:18:HIS:CE1	4.21	0.57
39:L2:104:LEU:O	39:L2:107:VAL:HG22	3.44	0.57
36:1:911:C:H42	39:L2:3:ARG:HD3	1.70	0.57
41:L4:10:SER:OG	41:L4:13:GLY:O	2.16	0.57
42:L5:88:ILE:HD12	42:L5:240:TYR:CE1	4.39	0.57
44:L7:24:GLU:O	44:L7:26:VAL:N	2.33	0.57
47:M0:98:ARG:HB3	47:M0:120:GLY:HA3	2.72	0.57
48:M1:93:ASP:O	48:M1:156:LYS:NZ	2.26	0.57
55:M9:125:LYS:NZ	36:5:1720:U:O4	241.50	0.57
59:N3:33:ASN:HD21	59:N3:63:LYS:N	2.02	0.57
66:O0:53:LYS:HZ2	66:O0:69:TYR:HE2	4.34	0.57
5:S3:103:GLU:HG3	5:S3:107:PHE:CE2	3.55	0.57
9:S7:114:ARG:O	9:S7:117:THR:HB	3.43	0.57
11:S9:34:PHE:HD1	11:S9:111:THR:HG21	1.69	0.57
11:S9:126:ARG:HA	11:S9:129:ILE:HD12	3.32	0.57
34:SR:13:LEU:HD12	34:SR:310:ILE:HG21	2.21	0.57
36:1:1716:U:O2'	36:1:1717:U:H4'	2.04	0.57
1:2:1165:G:C6	1:2:1166:A:C6	2.93	0.57
1:2:1424:A:H1'	4:S2:92:ALA:HB1	1.85	0.57
1:2:560:U:H2'	1:2:561:G:C8	2.39	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:761:G:OP1	11:S9:54:ARG:NH1	2.38	0.57
41:L4:221:ASN:ND2	36:5:211:A:H5''	80.04	0.57
36:5:2254:U:H2'	36:5:2261:G:N2	2.18	0.57
36:5:422:A:C2	36:5:2363:A:H4'	2.40	0.57
36:5:2897:A:H2'	36:5:2899:C:H5''	1.87	0.57
7:S5:103:ASN:OD1	1:6:1473:U:O2'	357.94	0.57
1:6:484:C:H42	1:6:503:G:N2	2.01	0.57
1:6:831:U:O2'	1:6:832:U:H5'	2.05	0.57
17:C5:64:LYS:HA	17:C5:73:PRO:HB3	1.85	0.57
18:C6:115:THR:O	18:C6:117:LEU:N	3.41	0.57
19:C7:13:SER:HA	19:C7:54:THR:HG22	2.36	0.57
21:C9:86:ARG:HB2	21:C9:89:ARG:HG3	3.83	0.57
24:D2:82:LYS:O	24:D2:84:GLY:N	2.30	0.57
39:L2:159:SER:O	39:L2:161:ASP:N	2.79	0.57
45:L8:241:LYS:HD3	36:5:2586:G:C8	184.17	0.57
63:N7:42:LEU:HD23	63:N7:96:VAL:HG12	3.54	0.57
70:O4:91:ARG:HG3	70:O4:95:ILE:HD13	1.87	0.57
36:1:2303:A:P	77:Q1:23:ARG:HH22	2.27	0.57
78:Q2:38:GLN:NE2	78:Q2:38:GLN:HA	2.54	0.57
4:S2:127:ALA:O	4:S2:131:ILE:HG13	2.04	0.57
5:S3:104:SER:OG	5:S3:105:MET:N	2.37	0.57
7:S5:98:MET:HE1	7:S5:105:GLY:HA2	1.86	0.57
9:S7:153:LEU:HD22	9:S7:184:GLU:HB3	1.86	0.57
1:2:333:A:H5'	10:S8:48:THR:HB	1.87	0.57
36:1:3049:A:C2	40:L3:75:ALA:HB2	2.40	0.57
1:2:1677:C:H2'	1:2:1678:A:O4'	2.05	0.57
1:2:52:U:H2'	1:2:53:G:C8	2.40	0.57
1:2:782:U:H5''	1:2:783:G:H5''	1.86	0.57
1:2:806:A:N6	9:S7:104:ARG:HH22	2.03	0.57
1:6:492:A:H2'	1:6:493:U:H5''	1.85	0.57
56:N0:52:LYS:NZ	37:7:101:G:OP2	281.44	0.57
15:C3:67:THR:O	15:C3:69:ASN:N	2.32	0.57
19:C7:35:CYS:O	19:C7:39:ALA:N	2.34	0.57
22:D0:25:THR:HB	22:D0:115:GLU:HG2	5.14	0.57
30:D8:36:THR:OG1	30:D8:37:SER:N	2.36	0.57
33:E1:109:ASP:HB2	33:E1:113:LYS:HG2	1.87	0.57
39:L2:65:ASP:OD2	39:L2:68:LYS:N	3.08	0.57
41:L4:16:THR:HG22	41:L4:18:ASN:H	1.69	0.57
45:L8:226:TYR:O	45:L8:229:VAL:N	2.37	0.57
36:1:2384:A:N1	52:M6:96:LYS:HE2	2.19	0.57
54:M8:26:LEU:HD23	54:M8:29:LEU:HD12	3.29	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:213:A:H2'	62:N6:10:SER:HB3	1.86	0.57
71:O5:49:LYS:HE3	71:O5:49:LYS:HA	1.87	0.57
2:S0:41:ARG:NE	2:S0:45:VAL:HB	2.09	0.57
6:S4:252:ARG:HA	6:S4:255:ARG:HG3	5.29	0.57
35:SM:73:SER:OG	35:SM:74:LYS:N	2.37	0.57
34:SR:5:GLU:HA	34:SR:317:THR:HA	2.73	0.57
36:1:1103:A:OP2	36:1:1103:A:H4'	2.03	0.57
36:1:2532:U:H3	36:1:2547:A:N6	2.02	0.57
36:1:2824:G:O6	86:1:3897:OHX:N4	2.37	0.57
36:1:610:G:O6	41:L4:309:ARG:NH2	2.38	0.57
1:2:1773:C:H2'	1:2:1774:G:C8	2.39	0.57
44:L7:94:LYS:NZ	36:5:1155:C:OP1	233.73	0.57
36:5:1200:A:H5'	36:5:1201:C:O5'	2.05	0.57
1:6:1213:G:O2'	1:6:1244:A:N6	2.23	0.57
1:6:1282:U:OP1	86:6:2137:OHX:N4	2.38	0.57
13:C1:37:ASN:HA	13:C1:44:THR:HG21	1.85	0.57
20:C8:30:TYR:CE2	20:C8:40:ARG:HD2	2.58	0.57
20:C8:53:ASP:OD2	20:C8:55:HIS:HB2	3.46	0.57
22:D0:44:ASN:OD1	22:D0:102:ARG:NH2	6.81	0.57
22:D0:82:TYR:HB3	31:D9:52:PHE:HB3	2.42	0.57
28:D6:87:ARG:CZ	28:D6:92:ARG:HA	2.78	0.57
29:D7:49:HIS:CE1	29:D7:70:LYS:HG2	2.86	0.57
40:L3:81:THR:HG22	40:L3:321:PHE:HA	4.41	0.57
45:L8:121:SER:O	45:L8:123:GLN:N	2.46	0.57
45:L8:130:TYR:CD1	45:L8:202:GLU:HB3	2.39	0.57
51:M5:98:LEU:HD22	51:M5:128:LYS:NZ	5.31	0.57
51:M5:45:PRO:O	51:M5:49:ARG:HB3	2.05	0.57
52:M6:15:LEU:HD12	52:M6:125:ARG:HA	2.67	0.57
55:M9:186:LYS:HG2	55:M9:187:GLU:HG3	1.87	0.57
58:N2:94:ARG:HG2	58:N2:96:VAL:HG22	5.50	0.57
60:N4:6:ASP:OD2	60:N4:7:SER:N	2.38	0.57
62:N6:115:ARG:O	62:N6:119:ILE:HG13	2.25	0.57
2:S0:26:ALA:HB3	2:S0:149:LEU:HB2	2.29	0.57
8:S6:180:THR:HG22	8:S6:181:PRO:HD2	1.87	0.57
36:1:1016:C:H1'	36:1:1028:U:C2	2.40	0.57
36:1:2157:G:O6	39:L2:152:SER:N	2.38	0.57
36:1:2656:A:C8	36:1:2658:G:C8	2.93	0.57
36:1:3028:G:H2'	36:1:3029:A:O4'	2.04	0.57
36:1:3200:G:C6	36:1:3201:C:C4	2.92	0.57
36:1:1363:A:OP2	86:1:4040:OHX:N6	2.38	0.57
36:1:56:G:H2'	36:1:57:A:H5''	1.87	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:1366:U:OP1	18:C6:30:LYS:HD2	2.05	0.57
1:2:55:A:H4'	1:2:459:G:OP1	2.05	0.57
1:2:826:U:H2'	1:2:827:C:C6	2.40	0.57
38:4:57:C:N4	73:O7:63:ARG:HH11	2.03	0.57
36:5:1581:C:OP2	36:5:1581:C:H4'	2.04	0.57
36:5:1952:G:H1	36:5:2094:C:H42	1.51	0.57
36:5:252:U:H4'	36:5:253:A:H5''	1.87	0.57
36:5:3279:A:H2'	36:5:3280:U:H5'	1.86	0.57
1:6:926:A:H2'	1:6:927:C:C6	2.40	0.57
24:D2:3:ARG:NH1	24:D2:9:ASP:OD2	3.83	0.57
39:L2:105:GLY:HA3	39:L2:160:SER:HB3	2.49	0.57
63:N7:73:LYS:HE3	36:5:1637:A:OP1	211.02	0.57
65:N9:3:LYS:HE3	36:5:2618:G:O4'	228.52	0.57
67:O1:10:ARG:HG2	67:O1:108:VAL:HA	1.86	0.57
67:O1:83:GLU:OE2	86:O1:202:OHX:N4	2.37	0.57
2:S0:164:ASN:HA	2:S0:170:ILE:HD11	1.87	0.57
3:S1:128:LYS:HG3	3:S1:134:VAL:HG22	1.86	0.57
8:S6:154:ARG:HD3	1:6:78:A:C8	341.42	0.57
34:SR:115:ILE:HD12	34:SR:122:ILE:HG12	1.85	0.57
36:1:1240:A:H3'	36:1:1241:U:C5'	2.35	0.57
36:1:1506:A:H1'	36:1:1848:G:O6	2.05	0.57
1:2:1475:A:H2'	1:2:1476:C:O4'	2.04	0.57
1:2:491:C:N3	1:2:496:G:N2	2.50	0.57
1:2:717:C:H2'	1:2:718:U:H5''	1.85	0.57
36:5:2689:A:C8	36:5:2702:A:C6	2.92	0.57
36:5:789:A:H2'	36:5:790:U:C6	2.40	0.57
16:C4:48:VAL:HG22	16:C4:49:LYS:H	1.99	0.57
21:C9:61:VAL:HG21	21:C9:104:VAL:HG11	1.86	0.57
22:D0:70:THR:HG23	1:6:1280:C:O2'	389.35	0.57
25:D3:6:PRO:HD2	25:D3:15:LEU:HD21	1.87	0.57
26:D4:88:THR:O	26:D4:92:VAL:HG22	3.85	0.57
29:D7:47:PHE:HE1	29:D7:49:HIS:HB2	1.70	0.57
33:E1:108:VAL:HB	33:E1:114:VAL:HG22	1.86	0.57
40:L3:159:ARG:HD3	40:L3:180:GLU:HB3	1.84	0.57
40:L3:79:VAL:HG22	40:L3:81:THR:OG1	2.05	0.57
41:L4:142:VAL:O	41:L4:144:LYS:N	2.37	0.57
42:L5:244:HIS:O	42:L5:248:ARG:HG3	2.05	0.57
43:L6:31:ARG:NH1	69:O3:107:ILE:O	2.38	0.57
45:L8:195:SER:O	45:L8:197:VAL:N	2.70	0.57
47:M0:12:GLN:NE2	47:M0:128:ARG:HG2	2.20	0.57
49:M3:159:VAL:HB	64:N8:96:LYS:HG2	1.86	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:M3:76:THR:HG22	49:M3:101:ARG:HB3	1.87	0.57
58:N2:23:THR:HA	58:N2:28:PHE:HB3	1.86	0.57
59:N3:13:ILE:HG12	59:N3:53:SER:HB2	1.86	0.57
63:N7:12:VAL:HB	63:N7:81:LEU:HB3	3.79	0.57
70:O4:105:VAL:HG12	70:O4:106:LYS:HG2	1.86	0.57
76:Q0:93:LYS:HD2	76:Q0:102:ARG:HG2	1.87	0.57
2:S0:163:ASN:HB3	2:S0:169:SER:OG	3.34	0.57
6:S4:68:ARG:HD3	6:S4:76:VAL:HG11	2.50	0.57
11:S9:125:ALA:HA	11:S9:128:LEU:HD12	3.75	0.57
36:1:2107:A:C2	36:1:3344:A:H8	2.23	0.56
36:1:2800:G:O6	64:N8:42:ARG:NH2	2.37	0.56
36:1:3085:G:H5''	36:1:3086:A:OP1	2.05	0.56
1:2:1002:G:H22	1:2:1761:U:P	2.27	0.56
1:2:416:A:H4'	1:2:417:A:OP2	2.04	0.56
36:5:1060:U:H2'	36:5:1061:A:C8	2.40	0.56
36:5:2775:U:H2'	36:5:2776:C:H6	1.69	0.56
1:6:328:A:H2'	1:6:329:G:C8	2.40	0.56
1:6:43:A:O2'	1:6:99:C:OP1	2.20	0.56
15:C3:114:ARG:HG3	1:6:952:A:O2'	299.75	0.56
18:C6:113:ASP:O	18:C6:114:ARG:HB2	2.03	0.56
21:C9:31:PRO:HG3	21:C9:103:LYS:HD3	1.87	0.56
39:L2:204:MET:HB2	39:L2:208:ASP:HB2	1.86	0.56
42:L5:106:ALA:O	42:L5:110:LEU:HD22	3.60	0.56
43:L6:154:LEU:HD23	43:L6:157:GLN:HB2	1.85	0.56
43:L6:46:ARG:HH11	43:L6:46:ARG:CG	2.91	0.56
43:L6:46:ARG:HH11	43:L6:46:ARG:HG3	2.95	0.56
44:L7:127:LEU:HD13	44:L7:136:TYR:CE2	4.14	0.56
45:L8:108:ARG:O	45:L8:112:GLU:HG2	2.05	0.56
36:1:716:A:N6	64:N8:117:ARG:HG3	2.19	0.56
49:M3:167:PHE:CE1	64:N8:132:LYS:HB2	2.40	0.56
38:4:52:A:H4'	75:O9:19:GLN:HA	1.87	0.56
5:S3:91:VAL:O	5:S3:93:ASP:N	3.27	0.56
7:S5:84:LYS:NZ	1:6:1614:A:OP2	366.89	0.56
7:S5:90:ILE:O	7:S5:94:THR:HG23	2.51	0.56
11:S9:60:LEU:HD21	11:S9:93:LEU:HG	1.85	0.56
17:C5:129:GLY:HA3	35:SM:74:LYS:HG2	4.85	0.56
34:SR:81:LEU:HD23	34:SR:91:LEU:HA	2.43	0.56
1:2:306:U:H2'	1:2:307:G:C8	2.40	0.56
1:2:485:A:H2'	1:2:486:G:O4'	2.05	0.56
38:4:52:A:H61	75:O9:35:ILE:HD12	1.70	0.56
36:5:1668:G:C6	36:5:1669:C:C4	2.93	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:252:U:H4'	36:5:253:A:C5'	2.35	0.56
36:5:2882:U:H2'	36:5:2883:U:C6	2.40	0.56
36:5:3264:G:N2	36:5:3265:C:H1'	2.20	0.56
36:5:228:U:OP2	86:5:4128:OHX:N2	2.38	0.56
36:5:917:A:OP2	86:5:4209:OHX:N1	2.38	0.56
1:6:1042:G:N2	1:6:1077:C:O2	2.38	0.56
7:S5:109:LYS:HE2	1:6:1473:U:H4'	362.45	0.56
1:6:1724:U:O4	86:6:2091:OHX:N5	2.38	0.56
1:6:696:C:H4'	1:6:697:C:C6	2.40	0.56
42:L5:265:TYR:CE1	37:7:121:U:H5''	315.16	0.56
20:C8:26:ILE:HG12	20:C8:31:ALA:HB2	3.65	0.56
25:D3:83:VAL:HG21	25:D3:122:PHE:CE2	4.52	0.56
28:D6:44:ILE:HD13	28:D6:65:PRO:HG2	4.25	0.56
39:L2:211:HIS:CD2	39:L2:219:ILE:HG23	3.46	0.56
40:L3:123:TYR:CZ	40:L3:124:LYS:HG3	2.40	0.56
41:L4:206:LEU:HD12	41:L4:248:VAL:HG22	2.02	0.56
41:L4:234:ASN:OD1	41:L4:236:LEU:N	3.45	0.56
43:L6:169:ASP:HB3	43:L6:174:LEU:HD11	2.69	0.56
44:L7:229:PHE:HD1	44:L7:229:PHE:C	2.52	0.56
48:M1:11:ASP:O	48:M1:12:LEU:HB3	4.58	0.56
50:M4:77:ARG:NH1	36:5:562:C:OP2	345.98	0.56
53:M7:32:THR:HG21	53:M7:87:SER:HB3	1.86	0.56
59:N3:44:SER:HB3	36:5:2916:U:H1'	262.49	0.56
64:N8:94:ALA:CB	64:N8:121:VAL:HG13	2.35	0.56
79:Q3:49:ARG:CD	79:Q3:50:GLY:H	2.18	0.56
2:S0:60:ALA:O	2:S0:64:ILE:HG13	2.41	0.56
8:S6:200:ALA:O	8:S6:203:GLU:N	2.81	0.56
10:S8:171:SER:OG	10:S8:180:ASP:N	2.28	0.56
36:1:2433:U:H1'	51:M5:125:SER:HB3	1.86	0.56
36:1:3309:G:O6	40:L3:21:ARG:NH2	2.38	0.56
36:1:924:G:OP1	86:1:4140:OHX:N5	2.38	0.56
36:1:3050:U:OP2	86:1:4178:OHX:N2	2.39	0.56
1:2:1:U:O2	1:2:369:A:H2'	2.04	0.56
73:O7:3:LYS:HG2	36:5:2138:A:O2'	173.00	0.56
40:L3:21:ARG:NH2	36:5:3309:G:O6	198.91	0.56
86:5:4014:OHX:N5	86:5:4212:OHX:N2	2.53	0.56
36:5:956:U:H2'	36:5:957:C:C6	2.40	0.56
1:6:151:G:H22	1:6:163:G:N2	2.03	0.56
47:M0:206:LEU:HD22	37:7:64:A:O5'	343.49	0.56
12:C0:32:HIS:HA	12:C0:39:ASN:HD22	4.83	0.56
16:C4:15:GLY:N	16:C4:78:ALA:O	2.39	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:L5:270:LYS:HB3	37:7:1:G:O2'	321.95	0.56
45:L8:117:ALA:HB1	36:5:252:U:H5'	66.76	0.56
46:L9:92:TYR:CG	46:L9:142:ASP:HB3	2.97	0.56
53:M7:110:THR:OG1	53:M7:111:LYS:N	2.37	0.56
53:M7:129:THR:HG23	53:M7:139:TYR:HB2	1.86	0.56
4:S2:157:LYS:HG2	4:S2:170:ILE:HG23	3.19	0.56
7:S5:208:SER:O	7:S5:210:ALA:N	3.22	0.56
8:S6:78:THR:HG22	8:S6:79:LYS:H	1.69	0.56
36:1:2163:C:O2	36:1:2171:G:N2	2.30	0.56
36:1:2244:A:H5''	39:L2:243:THR:OG1	2.05	0.56
36:1:3327:G:C2	36:1:3328:G:C8	2.94	0.56
1:2:1327:C:C2	1:2:1328:G:C8	2.93	0.56
1:2:218:A:O2'	1:2:219:A:OP1	2.20	0.56
1:2:260:U:H3'	1:2:261:U:H5''	1.87	0.56
1:2:514:G:O2'	1:2:515:A:H5'	2.06	0.56
37:3:62:U:H5''	42:L5:277:LEU:HD22	1.87	0.56
36:5:434:U:O5'	36:5:434:U:H6	1.88	0.56
36:5:760:G:H1'	36:5:770:G:N2	2.21	0.56
8:S6:13:GLN:CD	1:6:151:G:H21	311.76	0.56
1:6:699:U:O4	86:6:2073:OHX:N1	2.37	0.56
13:C1:29:LYS:O	13:C1:31:THR:N	2.35	0.56
14:C2:48:SER:OG	14:C2:120:VAL:O	3.31	0.56
19:C7:104:ASN:O	19:C7:106:THR:N	3.12	0.56
20:C8:88:ARG:NH1	20:C8:112:ASP:OD2	3.57	0.56
1:2:572:C:H5''	25:D3:109:ARG:NH2	2.20	0.56
33:E1:127:GLY:O	33:E1:129:GLY:N	2.37	0.56
39:L2:242:ARG:NH1	39:L2:244:GLY:O	2.33	0.56
40:L3:215:ILE:HD13	40:L3:282:ILE:HD11	2.18	0.56
36:1:578:A:H2'	41:L4:334:PHE:HD2	1.70	0.56
44:L7:166:ASN:OD1	44:L7:181:ILE:N	2.39	0.56
48:M1:26:SER:HB3	48:M1:64:LYS:O	2.05	0.56
51:M5:96:ARG:NH1	51:M5:96:ARG:HG2	2.57	0.56
52:M6:12:LYS:HG2	52:M6:40:GLU:HB3	4.26	0.56
60:N4:4:GLU:HG2	60:N4:30:ARG:HD2	1.87	0.56
63:N7:88:ASP:OD1	63:N7:89:VAL:N	2.39	0.56
73:O7:76:ASN:ND2	38:8:95:G:OP1	46.62	0.56
73:O7:76:ASN:O	73:O7:79:GLN:HG3	2.19	0.56
79:Q3:56:THR:HA	79:Q3:63:THR:HA	1.87	0.56
2:S0:124:THR:HG22	2:S0:174:TRP:HE1	2.23	0.56
6:S4:18:TRP:HE3	6:S4:20:LEU:HD11	1.70	0.56
7:S5:33:VAL:HG13	7:S5:37:GLN:OE1	3.08	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:S5:55:ASP:HB3	7:S5:58:LEU:HD12	1.88	0.56
36:1:1306:G:C6	52:M6:62:THR:HA	2.40	0.56
36:1:2107:A:H2	36:1:3344:A:H8	1.54	0.56
36:1:2167:A:H8	36:1:2167:A:O5'	1.89	0.56
1:2:1337:A:H5'	1:2:1338:C:OP2	2.06	0.56
1:2:17:C:H2'	1:2:18:C:C6	2.39	0.56
36:5:1506:A:H1'	36:5:1848:G:O6	2.05	0.56
1:6:1657:U:H4'	1:6:1658:G:OP2	2.06	0.56
1:6:1725:U:H2'	1:6:1726:G:O4'	2.06	0.56
1:6:484:C:H42	1:6:503:G:H1	1.54	0.56
26:D4:5:VAL:HG12	26:D4:6:THR:H	1.70	0.56
50:M4:119:GLN:O	50:M4:123:LEU:HG	3.17	0.56
52:M6:56:ASP:HA	52:M6:59:ARG:HD2	3.21	0.56
53:M7:67:ILE:CG2	53:M7:80:LYS:HB3	2.35	0.56
58:N2:21:SER:HB3	58:N2:107:PHE:HB2	3.58	0.56
64:N8:6:THR:HG23	64:N8:8:THR:HG23	1.87	0.56
36:1:1729:A:OP1	66:O0:88:GLY:N	2.38	0.56
68:O2:79:VAL:HG13	68:O2:111:ARG:HG2	2.73	0.56
69:O3:49:ILE:O	69:O3:69:GLY:N	2.38	0.56
70:O4:42:PRO:HD3	70:O4:56:THR:HG22	3.29	0.56
78:Q2:55:LYS:HD2	78:Q2:56:PRO:HD3	1.87	0.56
3:S1:169:SER:O	3:S1:173:THR:OG1	2.24	0.56
8:S6:139:ASN:HA	8:S6:142:ARG:HB2	2.35	0.56
9:S7:137:GLY:HA2	15:C3:18:TYR:CZ	2.40	0.56
11:S9:109:LEU:HD13	11:S9:129:ILE:HD13	1.88	0.56
36:1:321:C:H2'	36:1:322:U:H6	1.70	0.56
1:2:1228:G:P	14:C2:119:SER:HB3	2.45	0.56
1:2:1660:A:H2'	1:2:1661:U:C6	2.41	0.56
36:5:1021:G:O6	36:5:1030:A:N6	2.39	0.56
36:5:1249:G:H2'	36:5:1250:G:H8	1.70	0.56
36:5:563:U:H2'	36:5:564:G:H8	1.70	0.56
36:5:852:U:O2'	36:5:853:G:H5'	2.05	0.56
1:6:621:A:HO2'	1:6:1106:U:HO2'	1.53	0.56
1:6:158:U:O4	1:6:420:A:H4'	2.06	0.56
18:C6:113:ASP:OD2	18:C6:116:LEU:N	2.34	0.56
19:C7:31:ASN:HD21	19:C7:55:THR:HG22	3.78	0.56
19:C7:41:ILE:HG12	19:C7:50:ILE:HD12	1.87	0.56
24:D2:103:ILE:HA	24:D2:112:ASP:HA	2.11	0.56
24:D2:22:LYS:HA	29:D7:3:LEU:HD22	1.87	0.56
26:D4:29:HIS:CE1	26:D4:34:ASN:H	2.23	0.56
28:D6:82:ARG:HB2	28:D6:85:ARG:NE	9.34	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:C3:55:ARG:HD3	29:D7:47:PHE:CG	2.41	0.56
40:L3:65:SER:O	40:L3:67:PHE:N	2.72	0.56
42:L5:21:ARG:O	42:L5:25:GLU:HG3	2.06	0.56
46:L9:43:VAL:HG23	46:L9:57:VAL:HG23	5.91	0.56
46:L9:48:VAL:HG11	46:L9:52:LEU:HD13	1.87	0.56
48:M1:92:ARG:HB2	48:M1:94:ARG:HG2	1.88	0.56
52:M6:115:LYS:O	52:M6:117:ARG:NH1	2.39	0.56
54:M8:67:ILE:HG12	54:M8:81:VAL:HG21	1.86	0.56
36:1:1095:U:C2	57:N1:127:GLN:HG2	2.41	0.56
36:1:2339:C:P	59:N3:48:ARG:HG3	2.46	0.56
63:N7:11:ALA:HB1	63:N7:80:LEU:HD22	2.88	0.56
64:N8:74:ASN:CG	64:N8:115:LYS:HB2	2.26	0.56
64:N8:125:VAL:HG21	64:N8:138:ILE:HD13	2.47	0.56
69:O3:38:PRO:HG2	69:O3:39:GLN:OE1	4.27	0.56
69:O3:88:ASN:OD1	69:O3:88:ASN:N	2.37	0.56
69:O3:8:TYR:HB2	69:O3:100:ILE:O	2.05	0.56
70:O4:84:CYS:O	70:O4:88:ARG:HB3	2.06	0.56
2:S0:139:VAL:HG23	4:S2:62:PRO:HG3	2.01	0.56
6:S4:143:ASP:OD1	6:S4:145:ARG:NE	2.33	0.56
9:S7:114:ARG:NH2	1:6:637:C:O2	352.96	0.56
11:S9:107:ARG:HH22	11:S9:153:GLU:HG3	4.00	0.56
35:SM:37:VAL:HG12	35:SM:38:PRO:O	2.06	0.56
34:SR:70:ASP:HB3	34:SR:113:VAL:HG12	3.06	0.56
36:1:2107:A:H2	36:1:3344:A:C8	2.23	0.56
36:1:2320:A:C2	79:Q3:16:VAL:HG12	2.40	0.56
1:2:1341:A:O2'	34:SR:102:ARG:NH2	2.39	0.56
36:5:434:U:H2'	36:5:435:C:C6	2.39	0.56
36:5:883:A:H2'	36:5:921:A:C2	2.41	0.56
71:O5:49:LYS:NZ	38:8:63:G:O2'	52.27	0.56
17:C5:127:ARG:H	35:SM:71:ASN:HD21	2.89	0.56
20:C8:35:ILE:HB	20:C8:38:VAL:HG22	1.86	0.56
20:C8:94:ASP:OD1	20:C8:98:TYR:OH	2.12	0.56
21:C9:86:ARG:HG3	21:C9:86:ARG:HH11	1.71	0.56
23:D1:71:ARG:HG2	23:D1:75:ASN:ND2	2.21	0.56
23:D1:5:LYS:O	23:D1:7:GLN:N	2.39	0.56
25:D3:28:ASN:N	25:D3:28:ASN:OD1	2.33	0.56
28:D6:38:ARG:HH21	28:D6:83:ILE:HG21	1.71	0.56
46:L9:94:TYR:CE2	46:L9:98:PRO:HA	2.71	0.56
47:M0:185:ARG:HH21	47:M0:186:GLU:HG3	1.69	0.56
47:M0:210:ILE:HA	47:M0:217:PHE:CE2	2.64	0.56
52:M6:195:ALA:O	52:M6:198:GLY:N	2.47	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
62:N6:18:ALA:O	62:N6:22:ALA:HB2	2.06	0.56
70:O4:8:ARG:HB2	70:O4:34:HIS:NE2	2.21	0.56
3:S1:157:GLN:HB2	3:S1:160:HIS:ND1	2.20	0.56
6:S4:127:LYS:HA	6:S4:127:LYS:HE2	5.12	0.56
6:S4:18:TRP:HH2	6:S4:31:PRO:HD3	2.04	0.56
34:SR:214:ALA:HB2	34:SR:220:ILE:HG23	1.88	0.56
36:1:3278:C:H2'	36:1:3278:C:O2	2.04	0.56
36:1:3254:G:O6	86:1:4050:OHX:N5	2.39	0.56
1:2:1067:C:H2'	1:2:1068:C:C6	2.41	0.56
1:2:1240:U:H2'	1:2:1241:G:H5''	1.86	0.56
1:2:189:C:H2'	1:2:190:C:H5'	1.87	0.56
36:5:1063:G:OP2	36:5:1097:G:H5''	2.06	0.56
36:5:1919:G:N7	86:5:4066:OHX:N4	2.54	0.56
36:5:894:G:N2	36:5:1660:C:OP1	2.38	0.56
19:C7:5:ARG:NH1	1:6:1402:G:OP2	409.60	0.56
22:D0:22:ILE:HG22	22:D0:93:LEU:HB2	2.07	0.56
27:D5:93:SER:OG	27:D5:94:LYS:N	2.38	0.56
28:D6:42:ARG:HH21	28:D6:42:ARG:CB	4.80	0.56
28:D6:84:VAL:HG13	28:D6:85:ARG:N	2.20	0.56
30:D8:12:VAL:HG22	30:D8:28:VAL:HG11	1.87	0.56
40:L3:387:LEU:O	86:L3:406:OHX:N4	2.39	0.56
41:L4:261:VAL:O	41:L4:271:LYS:HE2	3.13	0.56
42:L5:61:ILE:HG12	42:L5:79:TYR:CD1	2.41	0.56
46:L9:188:THR:HG22	46:L9:189:GLU:H	4.79	0.56
56:N0:141:LYS:HA	56:N0:144:LEU:HD12	1.87	0.56
50:M4:38:ILE:HD11	56:N0:150:PHE:CE2	2.40	0.56
36:1:1612:A:H5''	74:O8:51:LEU:HD23	1.87	0.56
36:1:999:G:O2'	36:1:1000:C:H5'	2.06	0.56
36:1:1599:G:OP1	86:1:4080:OHX:N5	2.39	0.56
36:1:1721:U:O4	55:M9:128:LYS:NZ	2.38	0.56
36:1:2192:C:O2'	36:1:2312:A:N1	2.29	0.56
36:1:2503:G:H1'	36:1:2504:U:H5	1.70	0.56
36:1:3192:U:O4	86:1:4124:OHX:N1	2.39	0.56
36:1:3276:G:N7	53:M7:171:ARG:NH1	2.54	0.56
1:2:1490:C:H4'	1:2:1491:U:OP1	2.06	0.56
1:2:1649:G:N7	86:2:2050:OHX:N1	2.54	0.56
1:2:365:G:N7	86:2:2105:OHX:N5	2.53	0.56
1:2:396:G:H22	1:2:399:A:H5'	1.71	0.56
36:5:2147:A:H2'	36:5:2148:U:O4'	2.06	0.56
36:5:345:G:OP1	36:5:1429:G:N1	2.35	0.56
86:5:3935:OHX:N1	86:5:4228:OHX:N3	2.54	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:1232:U:H2'	1:6:1233:G:O4'	2.06	0.56
1:6:1316:G:O2'	1:6:1401:A:O2'	2.19	0.56
1:6:345:U:H1'	1:6:346:G:C8	2.41	0.56
37:7:28:C:H1'	37:7:55:A:H61	1.71	0.56
41:L4:51:ALA:HB3	38:8:27:U:H4'	109.49	0.56
18:C6:83:GLN:HE22	18:C6:119:ALA:HB2	1.70	0.56
20:C8:63:GLN:HA	20:C8:66:LEU:HD12	1.87	0.56
25:D3:23:ARG:O	25:D3:26:GLU:HB2	2.06	0.56
36:1:2548:C:OP2	39:L2:93:LYS:NZ	2.38	0.56
42:L5:279:LYS:HG2	42:L5:282:ARG:NH1	2.21	0.56
44:L7:89:ILE:HG22	44:L7:219:LYS:HE3	1.87	0.56
44:L7:229:PHE:CD1	44:L7:229:PHE:C	3.01	0.56
52:M6:171:LYS:O	52:M6:174:PHE:HB3	2.73	0.56
53:M7:25:SER:HB3	53:M7:28:ASN:HB2	1.87	0.56
56:N0:155:ARG:NH1	36:5:3206:C:O2	310.62	0.56
56:N0:77:VAL:HG11	56:N0:106:LEU:HG	4.47	0.56
65:N9:49:GLY:HA2	65:N9:52:LYS:HB3	2.77	0.56
74:O8:46:ARG:HG3	74:O8:47:GLY:O	2.59	0.56
79:Q3:44:LYS:HE2	36:5:1727:G:P	233.55	0.56
79:Q3:86:LEU:O	79:Q3:90:VAL:HG13	5.74	0.56
4:S2:175:GLY:O	11:S9:53:ARG:NH2	3.12	0.56
4:S2:95:ARG:HB3	4:S2:97:ARG:HD3	1.88	0.56
5:S3:101:GLN:HA	5:S3:104:SER:HB3	2.55	0.56
7:S5:25:LEU:HD22	7:S5:25:LEU:H	1.71	0.56
7:S5:40:ILE:HG12	7:S5:41:LYS:N	2.19	0.56
8:S6:7:TYR:CE1	8:S6:125:THR:HA	3.54	0.56
9:S7:13:PRO:HB2	9:S7:14:THR:HB	1.87	0.56
10:S8:84:HIS:HE1	10:S8:86:SER:HB2	1.71	0.56
36:1:3215:A:O5'	50:M4:121:MET:HE1	2.06	0.56
36:1:2841:G:OP2	86:1:4139:OHX:N2	2.39	0.56
1:2:1059:U:O2'	1:2:1060:U:N3	2.39	0.56
1:2:141:U:H5	8:S6:179:VAL:HG23	1.71	0.56
1:2:1599:C:O2	86:2:2110:OHX:N1	2.39	0.56
36:5:1066:G:OP1	86:5:4222:OHX:N2	2.37	0.56
1:6:1225:U:O2	1:6:1230:A:O2'	2.24	0.56
1:6:578:U:O2	86:6:2154:OHX:N5	2.39	0.56
1:6:755:A:O2'	1:6:756:A:OP1	2.17	0.56
12:C0:23:ALA:O	12:C0:24:LYS:HB3	4.62	0.56
22:D0:118:VAL:HG13	22:D0:119:ALA:H	3.16	0.56
43:L6:130:ILE:HG21	43:L6:135:VAL:HG23	1.87	0.56
48:M1:143:ARG:O	48:M1:144:CYS:HB2	2.59	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:M1:38:GLU:C	48:M1:40:LEU:H	2.09	0.56
48:M1:47:GLN:OE1	48:M1:64:LYS:HD2	3.86	0.56
50:M4:59:ASN:OD1	50:M4:60:LEU:N	2.39	0.56
53:M7:116:HIS:O	53:M7:149:VAL:N	2.39	0.56
55:M9:84:THR:O	55:M9:87:ALA:N	2.33	0.56
36:1:534:U:O2	56:N0:146:LYS:HA	2.06	0.56
59:N3:17:LEU:HD21	59:N3:98:ASN:ND2	2.21	0.56
62:N6:28:ARG:HB2	62:N6:75:ARG:HH21	1.71	0.56
66:O0:101:LEU:H	66:O0:101:LEU:HD22	3.22	0.56
69:O3:21:ARG:HG3	69:O3:21:ARG:NH1	2.15	0.56
71:O5:21:LEU:HD11	71:O5:55:LEU:HD21	1.88	0.56
7:S5:222:LYS:HG2	7:S5:225:ARG:NH2	2.21	0.56
36:1:2533:G:H2'	36:1:2534:G:O4'	2.06	0.56
36:1:1171:G:N7	86:1:3951:OHX:N5	2.53	0.56
36:1:796:U:H2'	36:1:797:U:C6	2.41	0.56
1:2:1476:C:H2'	1:2:1477:G:C8	2.39	0.56
1:2:453:U:O4	86:2:2038:OHX:N5	2.39	0.56
1:2:992:A:H2'	1:2:993:A:H5'	1.87	0.56
36:5:394:G:O2'	36:5:396:A:N7	2.26	0.56
1:6:1380:U:H2'	1:6:1381:U:C6	2.41	0.56
1:6:1579:U:H2'	1:6:1580:C:C6	2.40	0.56
86:6:2120:OHX:N2	86:6:2171:OHX:N1	2.53	0.56
1:6:250:C:H5'	1:6:250:C:H6	1.71	0.56
1:6:2:A:C8	1:6:370:A:H1'	2.40	0.56
1:6:809:A:C6	1:6:810:G:C6	2.94	0.56
13:C1:109:VAL:HG21	13:C1:139:VAL:H	1.70	0.56
20:C8:27:LYS:H	20:C8:57:ARG:NH2	2.04	0.56
1:2:611:U:OP1	25:D3:19:ARG:NH2	2.38	0.56
32:E0:20:LYS:HD2	32:E0:21:VAL:H	4.19	0.56
40:L3:113:GLU:HB3	40:L3:176:ALA:HB2	1.88	0.56
40:L3:44:THR:OG1	40:L3:182:GLN:O	2.37	0.56
46:L9:106:LYS:O	46:L9:109:ALA:HB2	2.07	0.56
48:M1:114:ILE:HG22	48:M1:115:LYS:O	3.07	0.56
51:M5:116:LEU:HD12	51:M5:151:ILE:HD13	4.99	0.56
51:M5:188:ARG:NH2	36:5:31:C:OP2	121.72	0.56
53:M7:24:VAL:HG12	53:M7:86:LYS:HG2	1.88	0.56
36:1:944:C:H4'	68:O2:33:ARG:CZ	2.35	0.56
73:O7:72:ARG:O	73:O7:75:LYS:N	2.38	0.56
3:S1:149:GLN:NE2	3:S1:151:LYS:HG2	3.96	0.56
3:S1:62:LYS:O	3:S1:64:ARG:N	2.39	0.56
5:S3:167:PHE:O	5:S3:190:ARG:HG2	4.89	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:S4:171:ASP:OD1	6:S4:172:PHE:N	2.39	0.56
1:2:512:A:OP2	11:S9:172:VAL:HG13	2.06	0.56
36:1:197:G:H2'	36:1:198:A:C8	2.41	0.55
1:2:1745:G:O6	86:2:2085:OHX:N6	2.39	0.55
1:2:770:A:OP2	86:2:2138:OHX:N6	2.40	0.55
1:2:604:A:OP2	86:2:2168:OHX:N5	2.39	0.55
1:2:808:U:H2'	1:2:809:A:C8	2.40	0.55
37:3:47:C:H2'	37:3:48:U:H6	1.72	0.55
36:5:1307:G:C2	36:5:1308:A:C2	2.94	0.55
36:5:2192:C:O2'	36:5:2312:A:N1	2.24	0.55
36:5:2726:C:O2'	36:5:2727:A:H2'	2.05	0.55
37:7:107:C:H2'	37:7:108:A:C8	2.41	0.55
18:C6:66:ARG:HD2	18:C6:68:ARG:HG3	1.87	0.55
20:C8:81:ILE:HG23	20:C8:82:PRO:HD2	1.87	0.55
23:D1:27:ASP:HA	23:D1:29:HIS:NE2	2.21	0.55
1:2:567:A:O2'	25:D3:90:ASP:OD2	2.19	0.55
39:L2:117:GLU:HB3	39:L2:119:LYS:O	2.05	0.55
41:L4:327:LEU:HD11	44:L7:165:ASP:HA	2.73	0.55
42:L5:278:SER:O	42:L5:280:GLU:N	2.95	0.55
44:L7:40:LYS:HD3	44:L7:170:GLU:OE1	2.06	0.55
47:M0:38:LYS:HG3	47:M0:41:ALA:HB2	3.23	0.55
53:M7:116:HIS:HB3	53:M7:149:VAL:HB	2.06	0.55
57:N1:82:ASN:HA	65:N9:21:ILE:HD12	2.64	0.55
36:1:1609:C:H5''	61:N5:125:ARG:NH1	2.21	0.55
63:N7:25:ILE:HG13	63:N7:43:VAL:HG12	3.32	0.55
67:O1:74:ARG:HH12	67:O1:109:VAL:HG11	2.06	0.55
67:O1:84:ASP:N	67:O1:84:ASP:OD1	2.77	0.55
71:O5:86:ARG:HG3	71:O5:90:ARG:CZ	3.38	0.55
51:M5:88:GLY:HA2	78:Q2:50:PHE:CE1	2.41	0.55
5:S3:11:LEU:HD12	22:D0:86:ILE:HG12	1.94	0.55
5:S3:162:GLN:N	5:S3:163:PRO:HD2	2.77	0.55
7:S5:56:ALA:O	7:S5:57:SER:OG	2.23	0.55
35:SM:102:THR:HG23	35:SM:105:LYS:HB2	1.88	0.55
36:1:2503:G:HO2'	36:1:2504:U:H5	1.53	0.55
36:1:2986:U:H2'	36:1:2987:A:H8	1.71	0.55
36:1:407:A:C2	38:4:17:A:H1'	2.40	0.55
1:2:1338:C:OP1	18:C6:12:LYS:NZ	2.37	0.55
1:2:1756:A:O5'	1:2:1756:A:H8	1.89	0.55
1:2:1795:U:H5'	28:D6:79:ILE:HD11	1.88	0.55
1:2:40:A:H2'	1:2:41:A:O4'	2.07	0.55
1:2:615:A:H2'	1:2:616:G:H8	1.71	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:3274:A:H3'	36:5:3275:U:C5'	2.27	0.55
36:5:1662:G:O6	86:5:3912:OHX:N1	2.38	0.55
36:5:528:U:H2'	36:5:529:A:C8	2.41	0.55
1:6:1309:C:H2'	1:6:1310:U:O4'	2.05	0.55
1:6:1489:U:H5'	1:6:1494:C:H1'	1.87	0.55
1:6:1:U:C4	1:6:369:A:C6	2.94	0.55
1:6:921:U:O4	86:6:2179:OHX:N3	2.39	0.55
13:C1:94:ILE:HD11	25:D3:16:ARG:HH21	3.57	0.55
1:2:951:A:H1'	15:C3:101:HIS:CD2	2.41	0.55
17:C5:127:ARG:HE	17:C5:130:ARG:HE	7.94	0.55
20:C8:13:HIS:O	20:C8:14:ILE:HG22	4.08	0.55
21:C9:14:PHE:HD2	21:C9:15:ILE:HD12	1.71	0.55
25:D3:84:THR:O	25:D3:120:VAL:HG13	2.07	0.55
26:D4:84:LYS:HD2	26:D4:85:PHE:CE2	2.41	0.55
30:D8:32:PHE:O	30:D8:34:GLU:N	3.11	0.55
36:1:1362:G:O2'	44:L7:159:GLN:O	2.16	0.55
44:L7:217:PRO:O	86:5:3995:OHX:N3	259.78	0.55
46:L9:174:LYS:HG3	46:L9:175:PHE:CD2	2.41	0.55
47:M0:169:LYS:HD3	57:N1:159:PHE:HA	1.88	0.55
47:M0:191:LYS:O	47:M0:197:VAL:HG22	2.52	0.55
62:N6:34:PRO:HA	62:N6:47:ALA:HA	1.89	0.55
64:N8:104:THR:HG21	64:N8:112:ILE:HD11	2.77	0.55
66:O0:27:TYR:OH	66:O0:55:GLU:OE1	2.19	0.55
73:O7:18:LEU:HA	73:O7:25:ARG:N	2.22	0.55
74:O8:65:LEU:HD23	74:O8:68:SER:HB2	1.88	0.55
36:1:1927:G:P	79:Q3:6:LYS:H	2.29	0.55
4:S2:174:ARG:HH12	11:S9:94:ASP:HB3	1.71	0.55
36:1:1227:C:H42	36:1:1282:G:H1	1.53	0.55
1:2:107:C:H42	1:2:307:G:H1	1.54	0.55
1:2:1239:U:O4	86:2:2046:OHX:N2	2.39	0.55
1:2:1619:C:H2'	1:2:1620:C:C6	2.41	0.55
36:5:1901:A:H5''	36:5:1902:G:OP2	2.07	0.55
36:5:2768:U:H2'	36:5:2769:A:C8	2.42	0.55
36:5:3044:G:H2'	36:5:3045:G:C8	2.41	0.55
36:5:325:A:H5''	36:5:326:U:OP2	2.06	0.55
1:6:235:G:H2'	1:6:236:A:H8	1.70	0.55
1:6:419:G:N7	86:6:2117:OHX:N1	2.54	0.55
1:6:542:A:C8	1:6:543:C:H2'	2.42	0.55
16:C4:61:MET:O	16:C4:65:GLN:N	2.39	0.55
18:C6:110:THR:HA	18:C6:113:ASP:HB2	2.28	0.55
22:D0:53:LYS:HB2	22:D0:92:ASP:HB2	2.73	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:D1:31:SER:HB2	23:D1:55:LEU:O	2.06	0.55
11:S9:123:HIS:ND1	32:E0:37:ARG:HD2	4.05	0.55
40:L3:161:LEU:HD22	40:L3:178:LEU:HD11	1.88	0.55
40:L3:97:ARG:NH1	36:5:3244:A:N1	245.28	0.55
41:L4:82:THR:HG23	41:L4:84:ARG:H	3.95	0.55
42:L5:144:VAL:O	42:L5:173:VAL:HG13	2.06	0.55
42:L5:156:GLY:HA2	42:L5:181:PRO:HD3	1.92	0.55
61:N5:80:ASN:HD21	61:N5:126:LEU:HB2	1.71	0.55
62:N6:58:VAL:HB	62:N6:63:LYS:O	2.56	0.55
66:O0:99:ASP:N	66:O0:99:ASP:OD2	2.39	0.55
2:S0:126:PRO:HB2	2:S0:152:PRO:HG2	2.65	0.55
6:S4:131:LEU:HD11	6:S4:135:GLY:HA2	1.87	0.55
8:S6:76:LEU:HD22	8:S6:92:ARG:HB3	1.89	0.55
36:1:1276:U:OP1	86:1:4081:OHX:N1	2.40	0.55
36:1:3254:G:H2'	36:1:3255:U:O4'	2.07	0.55
36:1:579:G:H4'	36:1:579:G:OP2	2.06	0.55
37:3:24:A:C2'	37:3:25:G:H5'	2.37	0.55
45:L8:105:LYS:NZ	36:5:123:A:OP1	92.21	0.55
36:5:1765:U:OP1	36:5:1765:U:H4'	2.06	0.55
49:M3:14:PHE:CE1	36:5:665:A:H1'	133.34	0.55
1:6:1273:G:H4'	1:6:1274:C:H5''	1.89	0.55
1:6:301:A:H2'	1:6:302:U:O4'	2.05	0.55
1:6:403:G:H8	1:6:403:G:P	2.29	0.55
1:6:700:C:H2'	1:6:701:U:C6	2.41	0.55
19:C7:41:ILE:HG22	19:C7:43:SER:H	1.72	0.55
24:D2:27:ILE:HB	24:D2:61:ILE:HB	4.46	0.55
26:D4:49:LYS:N	26:D4:49:LYS:HD3	2.80	0.55
47:M0:200:LEU:HD23	47:M0:202:LYS:HE3	1.87	0.55
47:M0:93:PRO:HB2	47:M0:125:LEU:HB2	4.48	0.55
48:M1:47:GLN:HG2	48:M1:67:VAL:HG12	1.88	0.55
51:M5:106:VAL:HG11	51:M5:132:VAL:HG21	1.89	0.55
54:M8:157:PRO:HA	54:M8:186:VAL:HG12	1.88	0.55
56:N0:131:LYS:HG3	56:N0:134:ASP:OD2	2.05	0.55
56:N0:82:ASP:OD1	56:N0:87:THR:HB	2.06	0.55
57:N1:7:TYR:OH	57:N1:54:HIS:HB2	2.07	0.55
58:N2:27:VAL:HG21	58:N2:107:PHE:HE1	1.72	0.55
59:N3:33:ASN:HD21	59:N3:64:LYS:H	1.54	0.55
64:N8:60:TYR:CG	64:N8:63:LYS:HE3	2.40	0.55
72:O6:89:GLU:O	72:O6:93:ILE:HG12	2.06	0.55
77:Q1:23:ARG:HH22	36:5:2303:A:P	269.98	0.55
79:Q3:56:THR:HG22	79:Q3:63:THR:HG23	1.88	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:S0:125:ASP:O	2:S0:128:SER:N	2.39	0.55
3:S1:181:LEU:O	3:S1:183:GLN:N	2.39	0.55
4:S2:177:GLY:HA2	4:S2:194:GLU:O	2.06	0.55
35:SM:85:SER:O	35:SM:87:THR:N	2.39	0.55
36:1:3078:U:O2	36:1:3078:U:H2'	2.06	0.55
36:1:317:A:C2	36:1:318:A:C4	2.94	0.55
86:1:3998:OHX:N3	86:1:4169:OHX:N5	2.55	0.55
1:2:505:A:N3	1:2:505:A:H2'	2.21	0.55
1:2:625:C:H2'	1:2:626:U:C6	2.41	0.55
1:2:929:A:H1'	16:C4:124:ASP:H	1.71	0.55
38:4:104:A:C8	38:4:105:A:C8	2.95	0.55
36:1:2585:G:N3	38:4:151:C:H5	2.05	0.55
39:L2:204:MET:HG2	36:5:914:A:C2	195.59	0.55
13:C1:33:ARG:HH21	13:C1:33:ARG:HG3	2.22	0.55
13:C1:92:HIS:CG	13:C1:93:TYR:N	2.66	0.55
1:2:1460:A:C8	17:C5:128:HIS:HB3	2.42	0.55
7:S5:72:HIS:O	18:C6:79:TYR:OH	2.24	0.55
48:M1:107:ASP:HA	48:M1:124:GLY:HA2	2.21	0.55
36:1:1915:A:H5''	55:M9:84:THR:HG22	1.88	0.55
58:N2:87:ASN:HB2	58:N2:89:LEU:HD11	4.05	0.55
62:N6:109:LEU:HD22	62:N6:115:ARG:NH1	2.20	0.55
71:O5:57:VAL:HA	71:O5:60:GLU:HG3	5.26	0.55
76:Q0:98:LYS:HD2	76:Q0:118:THR:HG21	1.88	0.55
2:S0:79:ARG:O	2:S0:83:GLN:NE2	4.21	0.55
6:S4:31:PRO:HB2	6:S4:38:LEU:HD22	1.88	0.55
6:S4:49:ARG:HH11	6:S4:50:ASN:HD21	1.53	0.55
36:1:2376:G:H2'	36:1:2377:G:C8	2.42	0.55
36:1:2514:U:OP1	36:1:2514:U:H6	1.90	0.55
36:1:3165:A:H2'	36:1:3166:C:C6	2.42	0.55
36:1:402:A:OP1	75:O9:36:ARG:NH2	2.40	0.55
1:2:1041:G:H2'	1:2:1042:G:C8	2.41	0.55
1:2:82:U:H2'	1:2:83:G:O4'	2.05	0.55
37:3:3:U:H2'	37:3:4:U:C6	2.42	0.55
36:5:1192:C:H41	36:5:1302:A:P	2.30	0.55
36:5:1560:G:H2'	36:5:1561:G:C8	2.42	0.55
36:5:1700:G:H1	36:5:1745:C:H42	1.55	0.55
36:5:1819:U:H2'	36:5:1820:U:H5'	1.88	0.55
36:5:304:G:H5'	36:5:304:G:N3	2.20	0.55
36:5:59:G:H4'	36:5:60:A:H4'	1.89	0.55
1:6:1339:C:O2'	1:6:1341:A:N7	2.40	0.55
35:SM:68:ARG:HD3	1:6:1460:A:P	335.35	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:1511:U:H2'	1:6:1512:G:C8	2.42	0.55
1:6:463:U:OP1	86:6:2204:OHX:N1	2.40	0.55
1:6:675:U:H2'	1:6:676:G:C8	2.41	0.55
38:8:122:U:H2'	38:8:123:G:H8	1.71	0.55
16:C4:84:ARG:HB2	16:C4:118:VAL:HG23	1.88	0.55
19:C7:47:ARG:NH1	19:C7:48:ASN:OD1	3.27	0.55
25:D3:12:ALA:O	25:D3:16:ARG:HG3	2.05	0.55
39:L2:233:GLN:O	39:L2:235:ALA:N	2.39	0.55
36:1:2415:C:OP1	39:L2:2:GLY:HA2	2.06	0.55
42:L5:69:ILE:HD12	42:L5:69:ILE:H	5.07	0.55
36:1:3267:A:H2'	43:L6:69:PHE:CZ	2.41	0.55
46:L9:101:VAL:HG22	46:L9:114:VAL:HG22	2.98	0.55
46:L9:86:TYR:CZ	46:L9:151:VAL:HG22	3.08	0.55
47:M0:4:ARG:NH1	36:5:2828:G:O2'	264.47	0.55
61:N5:51:VAL:HG11	71:O5:62:GLN:HB3	2.84	0.55
36:1:1729:A:N1	79:Q3:42:CYS:HB3	2.22	0.55
4:S2:38:VAL:HG22	4:S2:39:THR:H	1.71	0.55
7:S5:156:ARG:HH11	7:S5:156:ARG:HB2	1.72	0.55
9:S7:141:ARG:NH2	9:S7:143:LEU:HD11	2.22	0.55
1:2:40:A:OP1	11:S9:3:ARG:NH1	2.39	0.55
35:SM:102:THR:OG1	35:SM:103:LYS:N	2.40	0.55
36:1:1434:G:O2'	36:1:1435:A:H5'	2.07	0.55
36:1:1657:C:C5	36:1:1797:A:H5''	2.42	0.55
36:1:2224:A:N1	36:1:2783:U:O2'	2.35	0.55
1:2:1588:G:H1	1:2:1608:U:H3	1.54	0.55
1:2:763:G:C6	1:2:764:U:C4	2.95	0.55
37:3:89:G:N2	37:3:92:A:OP2	2.40	0.55
64:N8:22:ILE:HG13	36:5:1114:U:H5''	191.44	0.55
36:5:1549:U:H2'	36:5:1550:C:C6	2.42	0.55
36:5:1597:C:H2'	36:5:1598:G:C8	2.38	0.55
36:5:247:C:C4	36:5:248:U:H1'	2.42	0.55
45:L8:241:LYS:HB2	36:5:2586:G:N7	184.19	0.55
1:6:1335:U:H2'	1:6:1336:A:H8	1.72	0.55
17:C5:122:THR:OG1	1:6:1454:G:O3'	369.03	0.55
1:6:1595:U:N3	1:6:1600:A:H2	1.93	0.55
9:S7:104:ARG:NH1	1:6:745:U:O4	353.82	0.55
14:C2:36:LEU:HD11	14:C2:101:ALA:O	2.07	0.55
16:C4:29:HIS:CG	16:C4:41:ARG:HG3	3.48	0.55
19:C7:6:THR:OG1	19:C7:7:LYS:N	2.39	0.55
22:D0:27:THR:HG23	22:D0:113:ASP:HB3	1.97	0.55
29:D7:47:PHE:CE1	29:D7:49:HIS:HB2	2.42	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:D7:54:VAL:O	29:D7:63:LEU:HB2	2.07	0.55
1:2:1234:A:O2'	33:E1:146:SER:HB3	2.06	0.55
43:L6:131:LYS:HE2	43:L6:131:LYS:HA	5.38	0.55
43:L6:51:ARG:HD3	43:L6:158:TYR:CZ	2.42	0.55
49:M3:64:LYS:HG3	64:N8:69:TRP:CG	2.50	0.55
49:M3:68:LYS:HE3	49:M3:149:GLN:NE2	5.60	0.55
52:M6:115:LYS:HG2	36:5:3178:A:N3	260.07	0.55
56:N0:74:ASN:HD21	56:N0:144:LEU:HD21	1.72	0.55
61:N5:86:VAL:HG21	61:N5:95:ILE:HG12	2.49	0.55
65:N9:46:ALA:O	65:N9:50:THR:HG23	2.06	0.55
73:O7:14:LYS:HD2	75:O9:51:ILE:HD11	1.89	0.55
78:Q2:28:TYR:CE1	78:Q2:30:ALA:HA	5.49	0.55
2:S0:75:ALA:HB1	2:S0:86:VAL:HG12	1.88	0.55
1:2:1145:U:O2'	4:S2:89:GLN:O	2.17	0.55
6:S4:105:VAL:HG22	6:S4:243:GLY:HA2	1.87	0.55
6:S4:163:ASP:OD1	6:S4:164:LEU:N	4.75	0.55
6:S4:194:THR:O	6:S4:195:ILE:HB	2.07	0.55
35:SM:83:LYS:NZ	1:6:1178:G:H4'	339.08	0.55
36:1:1780:G:H2'	36:1:1781:C:C6	2.40	0.55
36:1:2723:U:H2'	36:1:2724:U:C6	2.41	0.55
36:1:291:C:OP1	51:M5:68:ARG:NH1	2.40	0.55
36:1:2942:C:O2	86:1:4130:OHX:N3	2.40	0.55
36:1:361:A:H5'	73:O7:35:SER:OG	2.06	0.55
36:1:603:A:H2'	36:1:604:G:O4'	2.06	0.55
36:1:735:A:H2'	36:1:736:A:C8	2.40	0.55
1:2:234:G:C6	1:2:235:G:H1'	2.42	0.55
53:M7:67:ILE:HD11	36:5:1447:G:H3'	164.82	0.55
36:5:2796:G:H4'	36:5:2798:C:C6	2.42	0.55
31:D9:14:TYR:HE1	1:6:1553:G:H4'	406.97	0.55
12:C0:34:GLU:O	12:C0:35:ILE:HB	4.30	0.55
15:C3:128:TYR:O	15:C3:132:VAL:HG22	2.52	0.55
15:C3:13:SER:O	29:D7:20:LYS:NZ	3.72	0.55
19:C7:32:LYS:HD2	19:C7:47:ARG:HH11	1.71	0.55
19:C7:82:ASP:O	19:C7:83:GLN:HB2	2.07	0.55
4:S2:144:TRP:CZ2	24:D2:97:ARG:HD2	3.37	0.55
25:D3:13:ARG:HA	25:D3:16:ARG:HD3	1.88	0.55
40:L3:287:LYS:HA	40:L3:320:ASP:HB3	1.89	0.55
41:L4:302:ALA:HB2	54:M8:39:ARG:CZ	2.36	0.55
44:L7:83:LEU:HD21	44:L7:116:PHE:HD1	1.71	0.55
44:L7:182:ASP:HA	44:L7:185:ILE:HD12	1.88	0.55
45:L8:147:LYS:O	45:L8:201:THR:HB	2.06	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:M1:75:LYS:O	48:M1:79:ILE:HG13	2.34	0.55
51:M5:124:ASP:OD2	51:M5:125:SER:N	2.39	0.55
51:M5:46:ASP:OD1	51:M5:50:ARG:NH2	2.25	0.55
53:M7:31:GLU:CG	53:M7:60:PHE:HA	3.46	0.55
64:N8:64:GLN:HG3	64:N8:67:HIS:CD2	2.42	0.55
69:O3:37:THR:HG23	69:O3:40:ASP:HB2	1.88	0.55
2:S0:66:ALA:HB2	23:D1:37:ALA:HB2	2.39	0.55
3:S1:147:ALA:O	3:S1:148:ASN:HB3	2.06	0.55
4:S2:170:ILE:O	4:S2:196:VAL:HG23	2.47	0.55
6:S4:161:LYS:HB3	6:S4:170:THR:O	4.25	0.55
7:S5:163:SER:O	7:S5:167:ARG:HB2	2.07	0.55
7:S5:53:VAL:HB	7:S5:59:VAL:HG22	1.88	0.55
8:S6:132:ARG:HB3	8:S6:133:LEU:HD12	4.74	0.55
8:S6:148:SER:O	8:S6:151:ASP:HB2	4.27	0.55
5:S3:115:ILE:HG21	35:SM:110:TRP:HA	1.87	0.55
35:SM:54:PRO:HB2	35:SM:59:GLY:HA2	2.48	0.55
36:1:1039:U:H2'	36:1:1040:A:C8	2.42	0.55
36:1:1660:C:H2'	36:1:1661:G:H8	1.71	0.55
36:1:3115:C:OP1	36:1:3115:C:H6	1.90	0.55
36:1:3318:G:OP2	36:1:3318:G:H2'	2.06	0.55
36:1:437:G:H2'	36:1:438:A:O4'	2.07	0.55
36:1:507:U:H2'	36:1:508:U:C6	2.42	0.55
36:1:769:G:N1	36:1:770:G:C6	2.75	0.55
1:2:1222:C:H2'	1:2:1223:A:H8	1.71	0.55
1:2:138:A:N6	1:2:266:A:H61	2.05	0.55
1:2:979:A:N3	1:2:1775:U:O2'	2.39	0.55
1:2:996:U:H3	1:2:1008:G:H1	1.54	0.55
56:N0:137:ARG:HH12	36:5:1213:G:P	324.09	0.55
36:5:1262:G:H5''	36:5:1263:A:OP2	2.07	0.55
36:5:1555:U:H5'	36:5:1556:C:OP2	2.07	0.55
70:O4:60:ARG:HG3	36:5:1802:C:H4'	153.22	0.55
36:5:2263:C:OP1	86:5:3949:OHX:N2	2.40	0.55
1:6:1018:U:H2'	1:6:1019:A:C8	2.41	0.55
1:6:1321:A:H4'	1:6:1322:A:O5'	2.06	0.55
1:6:431:C:H2'	1:6:432:G:H8	1.71	0.55
1:6:815:G:C8	1:6:815:G:H5'	2.40	0.55
1:2:306:U:H5''	13:C1:90:TYR:CE2	2.42	0.55
25:D3:70:LYS:HB3	25:D3:93:LEU:HD22	2.06	0.55
26:D4:129:VAL:O	26:D4:132:ARG:HB3	2.06	0.55
40:L3:262:TRP:HB3	52:M6:65:ASN:HA	2.20	0.55
40:L3:313:HIS:O	40:L3:333:LYS:HE3	3.13	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:L4:156:LEU:HD13	41:L4:215:ILE:HG12	2.46	0.55
43:L6:54:TYR:CE2	43:L6:63:LEU:HD22	2.42	0.55
45:L8:94:PHE:CZ	45:L8:200:LEU:HG	2.42	0.55
45:L8:243:GLN:O	45:L8:247:ASP:N	2.33	0.55
45:L8:62:LYS:NZ	51:M5:29:GLU:OE1	2.34	0.55
55:M9:128:LYS:HE2	36:5:1723:A:OP1	233.17	0.55
78:Q2:28:TYR:HE1	78:Q2:30:ALA:HA	6.10	0.55
5:S3:34:TYR:CE2	5:S3:37:VAL:HG13	3.71	0.55
36:1:1696:A:OP2	86:1:4155:OHX:N3	2.40	0.55
36:1:2404:A:H2'	36:1:2404:A:N3	2.22	0.55
36:1:3344:A:H2	36:1:3361:G:H21	1.55	0.55
36:1:619:A:H4'	36:1:620:U:O4'	2.07	0.55
1:2:918:U:H2'	1:2:919:A:H8	1.71	0.55
38:4:53:A:H2'	38:4:54:A:H8	1.72	0.55
38:4:79:A:C3'	38:4:80:A:H4'	2.37	0.55
45:L8:193:LYS:NZ	36:5:145:G:OP2	112.87	0.55
36:5:1556:C:H5''	36:5:2169:G:N2	2.21	0.55
36:5:1616:U:H2'	36:5:1617:G:C8	2.41	0.55
36:5:1715:A:H4'	36:5:1716:U:OP1	2.07	0.55
36:5:2623:G:C4	36:5:2624:G:C8	2.95	0.55
36:5:2651:G:H4'	36:5:2652:U:OP2	2.07	0.55
40:L3:250:ALA:HB1	36:5:2947:G:C2	219.68	0.55
36:5:407:A:C2	38:8:17:A:H1'	2.42	0.55
36:5:508:U:H2'	36:5:509:U:H6	1.72	0.55
18:C6:73:GLY:HA3	1:6:1608:U:O3'	397.81	0.55
1:6:820:U:O2'	1:6:821:U:H5''	2.07	0.55
37:7:106:U:H2'	37:7:107:C:O4'	2.07	0.55
13:C1:83:THR:HA	13:C1:111:VAL:HG12	2.36	0.55
18:C6:49:TYR:O	18:C6:53:LEU:HG	2.06	0.55
1:2:1369:U:OP2	21:C9:69:LYS:NZ	2.39	0.55
23:D1:24:ILE:HD11	23:D1:56:SER:HA	4.48	0.55
24:D2:86:ILE:HD12	24:D2:87:GLU:N	2.22	0.55
43:L6:72:ASN:ND2	43:L6:159:LEU:O	3.04	0.55
44:L7:216:VAL:HG11	44:L7:227:GLY:HA3	3.80	0.55
46:L9:161:LEU:HD22	46:L9:179:ILE:HD12	1.88	0.55
49:M3:63:VAL:HG22	36:5:72:C:H5'	113.20	0.55
50:M4:50:LYS:HD3	50:M4:85:TRP:CD1	2.42	0.55
51:M5:116:LEU:HB3	51:M5:133:ILE:HG12	1.89	0.55
51:M5:183:THR:OG1	51:M5:184:LYS:N	3.51	0.55
54:M8:71:LEU:HD13	54:M8:99:THR:HG21	1.88	0.55
57:N1:12:ARG:HD3	57:N1:13:TYR:CZ	3.05	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
61:N5:108:LEU:HD12	61:N5:125:ARG:HD2	1.89	0.55
62:N6:27:ARG:NH1	62:N6:76:LEU:HA	2.76	0.55
78:Q2:83:LEU:HD22	78:Q2:84:THR:H	1.72	0.55
3:S1:126:THR:HA	3:S1:136:ARG:HA	2.59	0.55
4:S2:125:ILE:O	4:S2:129:ILE:HG13	2.37	0.55
5:S3:136:VAL:HB	5:S3:152:PHE:HB2	1.88	0.55
6:S4:45:ILE:HA	6:S4:61:VAL:HG11	2.02	0.55
36:1:2916:U:C2'	36:1:2917:G:H5'	2.37	0.54
86:1:4194:OHX:N2	86:O1:202:OHX:N1	2.55	0.54
1:2:1370:U:H4'	1:2:1371:A:C5'	2.37	0.54
1:2:1648:A:H2'	1:2:1649:G:C8	2.41	0.54
36:5:1031:C:H2'	36:5:1032:C:H5''	1.89	0.54
36:5:1613:A:H2'	36:5:1614:C:C6	2.42	0.54
36:5:2440:G:H2'	36:5:2441:A:C8	2.41	0.54
36:5:2795:U:O2	36:5:2800:G:O2'	2.21	0.54
36:5:2881:C:H2'	36:5:2882:U:H6	1.71	0.54
1:6:1174:C:C4	1:6:1175:U:C4	2.95	0.54
1:6:1251:U:O2'	1:6:1252:C:H5''	2.07	0.54
1:6:1337:A:OP1	86:6:2178:OHX:N5	2.38	0.54
1:6:1591:C:H2'	1:6:1592:A:C8	2.42	0.54
86:6:2120:OHX:N6	86:6:2171:OHX:N3	2.55	0.54
7:S5:28:PRO:HB2	18:C6:37:THR:HG21	4.98	0.54
20:C8:4:VAL:HG21	27:D5:82:HIS:CD2	4.16	0.54
23:D1:87:ARG:O	29:D7:11:THR:HG23	3.06	0.54
25:D3:125:VAL:HG12	25:D3:126:LYS:HG3	2.25	0.54
26:D4:123:LYS:H	26:D4:123:LYS:HZ2	1.54	0.54
27:D5:83:LEU:HB2	27:D5:89:ILE:HD13	1.90	0.54
28:D6:26:CYS:HB3	28:D6:77:CYS:SG	2.46	0.54
28:D6:75:VAL:HA	28:D6:78:ALA:HB3	1.89	0.54
29:D7:46:VAL:HG13	29:D7:54:VAL:HG21	2.13	0.54
39:L2:79:ASN:O	39:L2:82:VAL:HG13	2.07	0.54
41:L4:37:THR:OG1	41:L4:38:VAL:N	2.40	0.54
42:L5:33:ARG:NH1	37:7:7:G:OP1	270.26	0.54
43:L6:96:VAL:HG12	43:L6:98:VAL:HG23	1.89	0.54
44:L7:232:ARG:O	44:L7:235:PHE:HB2	2.07	0.54
45:L8:163:VAL:O	45:L8:166:LEU:HB2	2.46	0.54
45:L8:161:GLU:HA	45:L8:164:VAL:HG22	1.92	0.54
47:M0:46:PHE:HD1	47:M0:140:THR:HA	2.39	0.54
50:M4:19:ARG:HA	50:M4:69:THR:HG22	2.54	0.54
51:M5:112:ASN:OD1	38:8:140:G:N2	102.67	0.54
51:M5:187:ARG:O	51:M5:190:THR:HG23	2.20	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:N0:33:ASN:O	56:N0:35:VAL:N	2.40	0.54
56:N0:124:LEU:HD23	57:N1:153:PRO:HG2	1.87	0.54
61:N5:121:LYS:HD3	61:N5:123:TYR:CZ	2.96	0.54
61:N5:82:LEU:O	61:N5:124:VAL:HG23	2.84	0.54
69:O3:13:HIS:O	69:O3:95:GLY:N	2.41	0.54
36:1:1489:A:OP1	70:O4:10:ARG:NH1	2.41	0.54
75:O9:23:LEU:HD22	75:O9:24:PRO:HD2	1.89	0.54
76:Q0:81:SER:O	76:Q0:84:ALA:HB3	3.54	0.54
77:Q1:9:ARG:HG3	77:Q1:9:ARG:HH11	3.37	0.54
3:S1:178:GLY:HA3	3:S1:187:LYS:HZ1	1.71	0.54
4:S2:80:VAL:O	4:S2:102:VAL:HA	2.07	0.54
10:S8:76:THR:HG21	10:S8:105:ASP:O	5.14	0.54
11:S9:11:THR:HG23	1:6:472:U:H5''	398.70	0.54
11:S9:38:ASN:HB3	11:S9:40:LYS:N	2.21	0.54
11:S9:64:GLU:HG3	11:S9:69:ARG:CZ	3.13	0.54
36:1:1766:G:OP2	36:1:1766:G:H8	1.91	0.54
36:1:209:A:H4'	36:1:211:A:C8	2.42	0.54
36:1:247:C:H2'	36:1:248:U:C6	2.42	0.54
36:1:665:A:H1'	49:M3:14:PHE:CE1	2.43	0.54
1:2:1595:U:N3	1:2:1600:A:H2	1.98	0.54
1:2:413:U:H2'	1:2:414:C:C6	2.43	0.54
1:2:936:G:O6	28:D6:15:ARG:NH1	2.41	0.54
1:2:97:C:H2'	1:2:98:U:C6	2.42	0.54
36:5:3279:A:C2'	36:5:3280:U:H5'	2.38	0.54
40:L3:334:ARG:NH2	36:5:3304:U:O2'	212.99	0.54
36:5:1878:G:OP1	86:5:3950:OHX:N5	2.40	0.54
36:5:1860:G:O6	86:5:4047:OHX:N2	2.41	0.54
1:6:978:A:H2'	1:6:979:A:O4'	2.07	0.54
19:C7:105:GLN:CD	19:C7:105:GLN:H	2.11	0.54
28:D6:87:ARG:NH2	28:D6:91:ASP:O	2.85	0.54
33:E1:135:HIS:HB2	33:E1:138:ARG:CB	2.37	0.54
43:L6:130:ILE:CG2	43:L6:135:VAL:HG23	2.37	0.54
46:L9:45:PHE:CE1	46:L9:55:VAL:HG13	3.51	0.54
49:M3:64:LYS:HG3	64:N8:69:TRP:CD1	2.42	0.54
52:M6:14:HIS:CE1	52:M6:119:VAL:HG12	2.40	0.54
56:N0:167:ARG:HG3	56:N0:168:PRO:HD2	1.89	0.54
56:N0:52:LYS:HE3	56:N0:54:ALA:HB3	1.89	0.54
64:N8:74:ASN:HB2	64:N8:76:ASP:HB3	2.52	0.54
64:N8:75:LEU:HD12	64:N8:137:LYS:HD3	1.88	0.54
70:O4:38:LEU:HD12	70:O4:38:LEU:H	4.05	0.54
72:O6:10:GLY:N	72:O6:13:LYS:HG2	2.64	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
73:O7:52:LYS:HG2	73:O7:56:ARG:NH2	2.23	0.54
3:S1:129:THR:HG22	3:S1:176:VAL:HG13	1.89	0.54
4:S2:90:THR:O	4:S2:93:GLY:N	2.24	0.54
6:S4:152:PRO:O	6:S4:154:ILE:N	2.40	0.54
10:S8:120:THR:O	86:S8:302:OHX:N4	5.99	0.54
10:S8:187:GLU:HG3	13:C1:30:ARG:HH12	1.72	0.54
36:1:3022:G:O2'	36:1:3031:G:O6	2.21	0.54
36:1:3248:C:O5'	36:1:3248:C:H6	1.91	0.54
36:1:23:A:OP1	86:1:3862:OHX:N5	2.40	0.54
86:1:3998:OHX:N6	86:1:4169:OHX:N1	2.56	0.54
1:2:1281:G:O3'	22:D0:76:SER:OG	2.24	0.54
1:2:1486:G:C8	1:2:1487:A:C8	2.96	0.54
1:2:1535:U:OP1	1:2:1535:U:H4'	2.07	0.54
1:2:891:A:H2'	1:2:892:A:C8	2.42	0.54
36:5:1049:C:H2'	36:5:1050:U:H6	1.71	0.54
36:5:1627:U:H2'	36:5:1814:A:N6	2.23	0.54
36:5:1840:U:OP2	86:5:4033:OHX:N4	2.40	0.54
36:5:2112:U:O2	86:5:3969:OHX:N1	2.40	0.54
36:5:568:G:N7	86:5:3932:OHX:N6	2.55	0.54
36:5:400:G:H4'	36:5:401:U:O5'	2.07	0.54
36:5:801:A:O2'	86:5:4022:OHX:N1	2.40	0.54
36:5:867:G:C6	36:5:868:C:C4	2.95	0.54
1:6:25:C:H4'	1:6:25:C:OP2	2.07	0.54
26:D4:10:ARG:HD3	1:6:780:A:C2	432.55	0.54
13:C1:46:LYS:HE2	1:6:846:G:N2	311.60	0.54
1:6:992:A:OP1	86:6:2053:OHX:N1	2.40	0.54
19:C7:71:PHE:O	19:C7:73:LEU:N	2.36	0.54
22:D0:57:ARG:HG3	22:D0:89:ARG:NE	2.50	0.54
1:2:1034:C:HO2'	24:D2:2:THR:N	2.04	0.54
26:D4:104:SER:O	26:D4:107:GLN:N	2.41	0.54
28:D6:26:CYS:O	28:D6:27:SER:HB2	2.35	0.54
28:D6:12:LYS:HB2	28:D6:33:ASP:OD2	2.07	0.54
28:D6:23:CYS:HB2	28:D6:74:CYS:HB3	1.89	0.54
39:L2:6:ARG:HH12	39:L2:199:THR:H	1.54	0.54
41:L4:148:ILE:HA	41:L4:149:PRO:C	2.60	0.54
44:L7:147:LEU:HD22	44:L7:205:PHE:CD1	3.67	0.54
45:L8:134:TYR:CD2	45:L8:190:VAL:HG21	2.42	0.54
47:M0:86:HIS:O	47:M0:138:VAL:HA	2.24	0.54
48:M1:117:ASP:O	48:M1:120:ILE:HG22	2.07	0.54
48:M1:54:VAL:HG12	48:M1:57:PHE:H	1.72	0.54
50:M4:24:LYS:HG2	50:M4:62:GLN:O	2.07	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:L4:362:ASP:H	56:N0:26:ARG:NH1	4.97	0.54
57:N1:8:ARG:HG3	36:5:2757:U:H4'	239.14	0.54
79:Q3:37:TYR:HB2	79:Q3:47:VAL:HB	1.89	0.54
4:S2:53:ILE:HD12	4:S2:53:ILE:H	4.06	0.54
7:S5:153:GLY:O	7:S5:155:ALA:N	2.89	0.54
7:S5:57:SER:OG	7:S5:58:LEU:N	2.81	0.54
8:S6:134:GLY:HA3	8:S6:158:ILE:HG13	6.48	0.54
1:2:332:U:OP2	10:S8:56:ARG:NH2	2.39	0.54
36:1:2674:A:C2	48:M1:124:GLY:HA3	2.42	0.54
36:1:2836:C:H5	36:1:2852:C:N4	1.94	0.54
36:1:2881:C:H2'	36:1:2882:U:H6	1.72	0.54
36:1:578:A:H5''	36:1:579:G:O5'	2.06	0.54
36:5:1768:U:H2'	36:5:1769:G:O4'	2.07	0.54
36:5:2425:G:H2'	36:5:2426:U:O4'	2.07	0.54
36:5:1686:U:H1'	36:5:3069:G:N2	2.22	0.54
36:5:3241:G:H2'	36:5:3245:A:C8	2.42	0.54
1:6:1594:G:OP2	1:6:1596:C:N4	2.40	0.54
1:6:417:A:H5'	1:6:418:G:C5	2.42	0.54
1:6:609:U:H4'	1:6:610:G:O5'	2.07	0.54
15:C3:56:ASP:N	15:C3:56:ASP:OD1	3.70	0.54
2:S0:63:ILE:HG12	23:D1:36:VAL:HG22	2.67	0.54
31:D9:20:GLN:HB2	31:D9:25:SER:HA	2.20	0.54
42:L5:22:ARG:HG2	42:L5:28:THR:OG1	2.08	0.54
42:L5:233:ALA:HA	42:L5:236:LEU:HD13	1.88	0.54
44:L7:26:VAL:C	44:L7:28:ALA:H	3.22	0.54
46:L9:61:GLY:O	46:L9:65:VAL:HG23	2.08	0.54
48:M1:54:VAL:HG11	48:M1:57:PHE:CG	2.43	0.54
50:M4:120:VAL:HG22	52:M6:197:LEU:HD13	2.46	0.54
50:M4:24:LYS:N	50:M4:62:GLN:O	2.40	0.54
51:M5:53:TYR:O	51:M5:54:LYS:HD2	2.07	0.54
52:M6:94:ARG:N	36:5:632:G:OP1	222.50	0.54
54:M8:62:VAL:O	54:M8:87:VAL:HA	2.46	0.54
55:M9:20:ARG:NH1	36:5:1873:U:OP2	147.48	0.54
76:Q0:110:CYS:SG	76:Q0:112:LYS:HB2	2.48	0.54
2:S0:169:SER:O	2:S0:173:ILE:HG12	2.06	0.54
9:S7:4:PRO:HD2	9:S7:5:GLN:HG2	1.89	0.54
34:SR:79:TYR:HE1	34:SR:100:TYR:CE1	3.81	0.54
34:SR:223:TRP:HE3	34:SR:230:ALA:HA	1.73	0.54
34:SR:282:SER:N	1:6:1394:G:OP1	417.06	0.54
36:1:1060:U:H2'	36:1:1061:A:C8	2.43	0.54
36:1:3242:G:N2	36:1:3245:A:H5''	2.22	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:39:A:H5''	64:N8:35:ALA:HB2	1.88	0.54
1:2:1675:C:H1'	10:S8:32:GLN:OE1	2.07	0.54
1:2:195:G:H2'	1:2:196:G:H5'	1.90	0.54
36:5:1791:C:H2'	36:5:1792:C:C6	2.42	0.54
36:5:653:A:H1'	36:5:2360:C:O2	2.07	0.54
16:C4:111:ARG:HH11	16:C4:111:ARG:HB3	5.07	0.54
17:C5:51:SER:C	17:C5:53:PRO:HD2	4.38	0.54
20:C8:26:ILE:HG23	20:C8:31:ALA:HB2	1.89	0.54
24:D2:118:ARG:HH11	24:D2:118:ARG:HB2	2.10	0.54
25:D3:24:TRP:HE3	25:D3:30:LYS:HD2	1.73	0.54
1:2:1102:G:OP2	25:D3:7:ARG:NH1	2.40	0.54
28:D6:23:CYS:O	28:D6:25:ASN:N	3.37	0.54
41:L4:22:LEU:HD21	41:L4:26:PHE:HB2	1.89	0.54
41:L4:311:HIS:CE1	41:L4:314:LYS:HA	2.86	0.54
41:L4:350:LYS:HG3	41:L4:351:PRO:HD2	1.89	0.54
41:L4:44:LYS:HB3	41:L4:47:ARG:NH1	2.51	0.54
36:1:2663:G:H5'	42:L5:152:ARG:HD3	1.89	0.54
43:L6:152:THR:O	43:L6:152:THR:OG1	2.40	0.54
38:4:155:A:H5'	45:L8:185:ARG:CZ	2.37	0.54
52:M6:36:VAL:HB	52:M6:108:ILE:HG12	1.89	0.54
53:M7:59:PRO:HB3	53:M7:78:VAL:HG21	1.88	0.54
56:N0:12:ARG:NH1	56:N0:15:PRO:HG3	2.23	0.54
63:N7:38:PHE:O	63:N7:40:HIS:ND1	2.40	0.54
68:O2:88:HIS:HB3	68:O2:91:THR:HG22	4.66	0.54
70:O4:52:GLN:HB3	36:5:1639:C:OP1	197.26	0.54
73:O7:14:LYS:HE2	75:O9:51:ILE:HG12	2.13	0.54
5:S3:119:ALA:O	5:S3:123:VAL:HG23	2.07	0.54
36:1:1478:C:H2'	36:1:1479:U:C6	2.42	0.54
36:1:3375:A:O2'	36:1:3378:C:H5'	2.08	0.54
1:2:1068:C:H2'	1:2:1069:A:C8	2.43	0.54
1:2:1358:G:H2'	1:2:1359:C:H6	1.71	0.54
1:2:1165:G:N2	1:2:1581:C:O2	2.41	0.54
1:2:1648:A:H2'	1:2:1649:G:H8	1.73	0.54
1:2:75:U:N3	1:2:76:A:N3	2.56	0.54
52:M6:18:ARG:NH1	36:5:1315:U:OP1	277.52	0.54
36:5:2249:G:C8	36:5:2249:G:H3'	2.43	0.54
36:5:2256:A:OP2	36:5:2256:A:H8	1.91	0.54
36:5:1930:A:O2'	86:5:3920:OHX:N3	2.39	0.54
39:L2:181:LYS:NZ	36:5:860:G:O5'	212.89	0.54
36:5:985:U:H2'	36:5:986:U:H6	1.72	0.54
1:6:694:U:H3'	1:6:695:U:O2	2.07	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:73:U:H2'	1:6:74:U:C6	2.43	0.54
42:L5:203:HIS:NE2	37:7:47:C:OP1	281.51	0.54
7:S5:189:THR:OG1	27:D5:98:GLN:OE1	2.24	0.54
39:L2:41:ILE:HG22	39:L2:90:ALA:HB3	1.90	0.54
42:L5:39:GLN:OE1	42:L5:40:HIS:N	2.34	0.54
48:M1:161:SER:OG	48:M1:162:TRP:N	2.39	0.54
51:M5:193:ARG:C	51:M5:195:ASN:H	2.10	0.54
51:M5:38:ARG:HD3	38:8:142:C:OP1	115.46	0.54
52:M6:102:LEU:HD12	52:M6:103:LYS:H	1.71	0.54
59:N3:96:GLU:HG2	59:N3:96:GLU:O	3.41	0.54
62:N6:17:LYS:HD3	62:N6:21:THR:HG21	1.89	0.54
43:L6:86:ALA:H	69:O3:107:ILE:HG21	4.69	0.54
71:O5:101:THR:HG22	71:O5:104:GLN:N	2.23	0.54
45:L8:144:GLU:OE1	72:O6:36:ARG:NH2	3.05	0.54
78:Q2:35:LEU:HD13	78:Q2:36:PHE:CE1	2.43	0.54
3:S1:175:GLU:HG3	3:S1:193:ILE:HD12	1.89	0.54
6:S4:16:HIS:O	6:S4:18:TRP:N	2.40	0.54
9:S7:12:ALA:HB3	9:S7:13:PRO:HD3	1.88	0.54
35:SM:131:ILE:O	35:SM:135:ALA:N	3.00	0.54
36:1:138:U:H2'	36:1:139:G:H8	1.71	0.54
36:1:1687:U:H2'	58:N2:70:LYS:NZ	2.23	0.54
36:1:2554:A:H62	79:Q3:62:LYS:NZ	2.04	0.54
36:1:3094:A:H2'	36:1:3095:U:C6	2.43	0.54
36:1:41:G:N2	36:1:2803:A:N7	2.56	0.54
1:2:542:A:C8	1:2:543:C:H3'	2.43	0.54
36:5:1072:G:H2'	36:5:1073:U:C6	2.42	0.54
45:L8:147:LYS:HD3	36:5:117:U:O4	106.00	0.54
36:5:119:U:H4'	36:5:120:G:H3'	1.89	0.54
36:5:739:G:O6	86:5:3959:OHX:N6	2.41	0.54
1:6:1690:G:H1	1:6:1711:C:H42	1.55	0.54
1:6:58:U:O2'	1:6:451:A:N3	2.33	0.54
71:O5:86:ARG:NE	38:8:37:A:OP2	83.26	0.54
18:C6:17:THR:O	18:C6:68:ARG:NH2	7.04	0.54
2:S0:200:ASP:HB2	19:C7:85:VAL:HG13	1.89	0.54
27:D5:61:SER:H	27:D5:64:VAL:HB	1.72	0.54
40:L3:123:TYR:CZ	40:L3:124:LYS:HD3	3.11	0.54
40:L3:339:ARG:CZ	40:L3:342:LEU:HD21	3.33	0.54
36:1:1382:G:P	41:L4:188:ARG:HH12	2.31	0.54
42:L5:257:GLU:O	42:L5:258:LYS:HB2	2.08	0.54
48:M1:8:PRO:HG2	48:M1:9:MET:HB3	1.89	0.54
51:M5:183:THR:O	51:M5:184:LYS:HB3	3.16	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:N0:93:GLU:OE1	56:N0:137:ARG:HB2	2.86	0.54
60:N4:4:GLU:O	60:N4:13:ILE:N	2.35	0.54
64:N8:66:ALA:HA	64:N8:69:TRP:N	4.29	0.54
67:O1:7:VAL:HG13	67:O1:77:ARG:O	3.04	0.54
70:O4:24:LYS:HB3	36:5:1695:U:H5'	152.03	0.54
2:S0:70:PRO:HB2	2:S0:94:GLY:HA3	1.90	0.54
4:S2:227:PRO:HA	4:S2:230:TRP:CE2	2.43	0.54
7:S5:133:VAL:HA	7:S5:198:LEU:HD22	2.26	0.54
36:1:345:G:OP1	36:1:1429:G:N1	2.39	0.54
36:1:498:A:P	69:O3:86:ARG:HH21	2.30	0.54
1:2:1498:G:C2'	1:2:1499:G:H5'	2.38	0.54
1:2:432:G:C6	1:2:433:C:C4	2.96	0.54
1:2:434:G:N2	1:2:436:A:H3'	2.22	0.54
1:2:814:A:O2'	1:2:816:G:OP2	2.26	0.54
1:2:959:U:O2	1:2:959:U:H2'	2.08	0.54
37:3:106:U:H2'	37:3:107:C:H6	1.73	0.54
36:5:2523:A:O2'	36:5:2587:U:H1'	2.08	0.54
1:6:151:G:H2'	1:6:152:U:C6	2.42	0.54
1:6:1203:A:C4	1:6:1556:A:C2	2.96	0.54
26:D4:105:ARG:HB2	1:6:443:C:OP2	372.13	0.54
3:S1:136:ARG:NH1	1:6:885:G:OP1	274.96	0.54
38:8:122:U:H2'	38:8:123:G:C8	2.43	0.54
16:C4:26:THR:HG21	16:C4:97:GLY:HA3	1.88	0.54
21:C9:108:LEU:HB3	21:C9:114:VAL:HG22	5.98	0.54
21:C9:23:GLN:HG2	21:C9:55:TYR:CG	2.43	0.54
23:D1:35:ASN:HB3	23:D1:50:TYR:CD1	2.42	0.54
24:D2:83:ILE:O	24:D2:86:ILE:HD11	2.08	0.54
25:D3:100:ASP:O	25:D3:101:GLU:HB3	4.75	0.54
40:L3:84:VAL:HG22	40:L3:162:VAL:HB	1.88	0.54
43:L6:54:TYR:HA	43:L6:65:ILE:CD1	5.83	0.54
44:L7:169:ILE:HD12	44:L7:181:ILE:HA	1.89	0.54
56:N0:1:MET:SD	56:N0:36:ILE:HG21	2.48	0.54
61:N5:141:TYR:O	61:N5:142:ILE:HG13	3.95	0.54
73:O7:67:LEU:O	73:O7:69:HIS:N	3.01	0.54
3:S1:131:ASP:O	3:S1:133:TYR:N	2.32	0.54
8:S6:153:VAL:HG11	8:S6:175:ILE:HG21	1.88	0.54
11:S9:81:VAL:HG22	11:S9:86:LEU:HB3	1.90	0.54
34:SR:198:ASN:O	34:SR:215:GLY:HA3	2.30	0.54
36:1:1047:A:C6	36:1:1048:A:C6	2.96	0.54
36:1:1667:A:H2'	36:1:1668:G:H8	1.72	0.54
36:1:1919:G:N7	86:1:4008:OHX:N5	2.56	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:207:U:H2'	36:1:208:C:H6	1.72	0.54
36:1:2712:U:H2'	36:1:2713:U:C6	2.43	0.54
36:1:563:U:H2'	36:1:564:G:H8	1.72	0.54
1:2:1350:U:H2'	1:2:1351:G:H8	1.73	0.54
1:2:118:U:OP1	86:2:2134:OHX:N3	2.41	0.54
1:2:843:U:H2'	1:2:844:A:C8	2.43	0.54
38:4:19:C:H2'	38:4:20:U:O4'	2.07	0.54
38:4:83:C:H1'	38:4:85:G:N2	2.23	0.54
36:5:1236:G:N2	36:5:1244:A:OP1	2.39	0.54
36:5:1659:U:H2'	36:5:1660:C:C6	2.42	0.54
36:5:2561:A:O2'	36:5:2562:A:H5''	2.08	0.54
52:M6:156:LEU:HD13	36:5:3243:A:C8	263.77	0.54
86:5:4014:OHX:N6	86:5:4212:OHX:N2	2.55	0.54
36:5:2732:G:OP2	86:5:4214:OHX:N1	2.41	0.54
36:5:644:G:H2'	36:5:2372:A:N7	2.23	0.54
36:5:842:G:H1	36:5:851:C:H42	1.56	0.54
1:6:1645:G:OP2	86:6:2183:OHX:N3	2.40	0.54
15:C3:138:ASN:O	15:C3:140:LYS:N	3.40	0.54
15:C3:65:VAL:C	15:C3:67:THR:H	3.03	0.54
17:C5:15:HIS:H	17:C5:22:LEU:HD22	4.84	0.54
19:C7:12:ALA:O	19:C7:15:ALA:HB3	2.50	0.54
23:D1:71:ARG:HG3	23:D1:83:TRP:CZ2	2.89	0.54
1:2:601:A:OP1	25:D3:110:LYS:HD3	2.08	0.54
24:D2:57:ARG:NH2	29:D7:26:GLN:OE1	3.31	0.54
40:L3:58:ARG:HA	40:L3:357:LYS:HG3	1.90	0.54
43:L6:52:VAL:HG22	43:L6:67:GLY:HA2	2.33	0.54
47:M0:156:ARG:NH1	47:M0:163:GLN:O	3.03	0.54
65:N9:14:ARG:HH21	65:N9:18:ARG:HD3	4.43	0.54
67:O1:75:ILE:HG12	67:O1:93:VAL:HG22	1.90	0.54
70:O4:47:CYS:SG	70:O4:48:GLY:N	2.80	0.54
6:S4:182:TYR:HA	6:S4:192:ILE:HG23	1.90	0.54
8:S6:64:LYS:HB2	8:S6:97:VAL:HG21	1.90	0.54
11:S9:26:ALA:HA	11:S9:29:LYS:HD3	4.40	0.54
36:1:1076:C:O3'	65:N9:38:LYS:NZ	2.40	0.54
36:1:1190:A:H4'	36:1:1191:U:OP1	2.06	0.54
36:1:2718:U:H2'	36:1:2719:U:C6	2.43	0.54
1:2:1068:C:H2'	1:2:1069:A:H8	1.72	0.54
1:2:11:A:C2'	1:2:12:U:H5'	2.38	0.54
1:2:67:A:O2'	1:2:69:G:OP1	2.19	0.54
37:3:22:A:H2'	37:3:23:A:C8	2.43	0.54
36:5:736:A:C5	36:5:737:G:H1'	2.42	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:1037:C:H2'	1:6:1038:U:C6	2.43	0.54
1:6:1263:G:C2	1:6:1264:G:H1'	2.43	0.54
1:6:1579:U:H2'	1:6:1580:C:H6	1.73	0.54
10:S8:142:LYS:NZ	1:6:186:C:OP2	277.15	0.54
1:6:482:U:H3	1:6:505:A:H61	1.55	0.54
1:6:569:C:H2'	1:6:570:A:O4'	2.07	0.54
14:C2:103:LEU:HD23	14:C2:115:VAL:HA	2.34	0.54
15:C3:12:SER:O	15:C3:13:SER:HB3	2.08	0.54
16:C4:80:HIS:ND1	16:C4:113:GLY:O	2.40	0.54
18:C6:68:ARG:NE	18:C6:68:ARG:O	4.65	0.54
22:D0:16:GLN:HB2	22:D0:17:GLN:HE21	1.73	0.54
24:D2:53:ILE:HB	24:D2:60:LYS:HB2	4.78	0.54
39:L2:102:LEU:HD22	39:L2:166:ILE:HD11	1.90	0.54
40:L3:53:MET:HE1	40:L3:327:CYS:CB	2.66	0.54
43:L6:40:LEU:HB3	43:L6:84:VAL:HG13	1.90	0.54
36:1:1101:G:OP2	44:L7:196:LYS:HE3	2.08	0.54
48:M1:12:LEU:HD12	48:M1:131:MET:HE3	1.90	0.54
51:M5:139:HIS:HB3	51:M5:142:ILE:HD13	1.90	0.54
55:M9:84:THR:O	55:M9:86:GLU:N	2.41	0.54
57:N1:104:GLU:HG3	57:N1:105:PHE:N	2.22	0.54
57:N1:8:ARG:HD2	36:5:2756:C:O2'	243.11	0.54
58:N2:41:ILE:HG12	58:N2:79:LEU:HD13	1.90	0.54
59:N3:80:ARG:NH1	59:N3:117:PRO:O	2.36	0.54
63:N7:53:VAL:HA	63:N7:57:HIS:CD2	2.43	0.54
64:N8:59:ARG:NH1	36:5:90:C:OP1	152.03	0.54
67:O1:25:PHE:O	67:O1:27:LYS:N	2.60	0.54
74:O8:62:ALA:O	74:O8:66:ILE:HG13	2.07	0.54
2:S0:49:ASN:HD22	2:S0:52:LYS:HE3	4.12	0.54
3:S1:104:ASP:HA	3:S1:214:LYS:HG3	1.89	0.54
3:S1:41:ARG:HH22	3:S1:97:LEU:HD21	1.73	0.54
6:S4:108:ARG:HH22	1:6:788:A:H3'	394.46	0.54
6:S4:159:THR:HG23	6:S4:173:ILE:HD13	1.88	0.54
11:S9:109:LEU:HD22	11:S9:113:VAL:HG23	1.90	0.54
36:1:1609:C:H5''	61:N5:125:ARG:HH11	1.73	0.53
36:1:2273:G:N2	36:1:2311:G:H2'	2.23	0.53
36:1:2586:G:N7	45:L8:241:LYS:HB2	2.23	0.53
36:1:2687:G:OP1	42:L5:8:LYS:NZ	2.25	0.53
36:1:3393:U:H2'	36:1:3394:U:C6	2.43	0.53
36:1:1170:A:OP2	86:1:3951:OHX:N3	2.40	0.53
36:1:608:A:C6	43:L6:22:ARG:HD3	2.43	0.53
36:1:945:C:H2'	36:1:946:U:H6	1.73	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:985:U:H2'	36:1:986:U:H6	1.73	0.53
1:2:615:A:H2'	1:2:616:G:C8	2.43	0.53
1:2:947:U:H2'	1:2:948:G:C8	2.43	0.53
1:2:978:A:H2'	1:2:979:A:O4'	2.07	0.53
36:5:290:G:H2'	36:5:291:C:C6	2.43	0.53
1:6:138:A:H62	1:6:266:A:H61	1.55	0.53
31:D9:14:TYR:CE1	1:6:1553:G:H4'	406.60	0.53
1:6:1638:G:C2	1:6:1639:C:H1'	2.43	0.53
1:6:769:A:OP2	86:6:2138:OHX:N4	2.41	0.53
1:6:489:C:O2'	1:6:490:C:O4'	2.25	0.53
1:6:75:U:O2'	1:6:76:A:O4'	2.26	0.53
38:8:81:U:H1'	38:8:82:U:H5'	1.90	0.53
22:D0:48:HIS:HE1	22:D0:102:ARG:NH2	10.29	0.53
2:S0:66:ALA:O	23:D1:50:TYR:HE1	1.90	0.53
27:D5:41:ILE:O	27:D5:75:LEU:HD13	2.07	0.53
27:D5:82:HIS:O	27:D5:82:HIS:ND1	2.41	0.53
28:D6:45:VAL:HB	28:D6:49:ALA:HB3	1.90	0.53
40:L3:282:ILE:HG23	40:L3:322:ILE:HG23	1.90	0.53
40:L3:219:ALA:HB2	40:L3:336:VAL:HG13	2.70	0.53
41:L4:50:TYR:CD2	41:L4:109:TRP:HH2	2.62	0.53
42:L5:279:LYS:HA	42:L5:282:ARG:HB2	1.90	0.53
46:L9:75:VAL:HA	46:L9:78:MET:HG3	1.88	0.53
52:M6:32:LYS:HA	52:M6:101:ARG:HB2	1.88	0.53
60:N4:25:ASP:OD2	60:N4:25:ASP:N	4.14	0.53
60:N4:39:LEU:O	60:N4:42:GLN:N	3.22	0.53
64:N8:88:ASP:N	64:N8:88:ASP:OD2	4.21	0.53
63:N7:136:PHE:CE1	70:O4:89:ILE:HG12	3.32	0.53
72:O6:35:ASN:OD1	72:O6:35:ASN:N	2.85	0.53
79:Q3:38:ASP:HA	79:Q3:45:LYS:HA	1.90	0.53
4:S2:212:LYS:O	4:S2:216:VAL:HG23	2.07	0.53
8:S6:139:ASN:OD1	8:S6:142:ARG:NH1	2.38	0.53
36:1:3049:A:OP2	86:1:4178:OHX:N1	2.42	0.53
1:2:1157:A:HO2'	1:2:1158:C:P	2.31	0.53
1:2:1261:G:H2'	1:2:1262:U:C6	2.42	0.53
1:2:1308:G:C6	1:2:1309:C:C4	2.96	0.53
1:2:511:A:P	11:S9:176:ASN:HD22	2.32	0.53
1:2:539:G:OP2	1:2:539:G:H8	1.91	0.53
1:2:826:U:H2'	1:2:827:C:H6	1.74	0.53
36:5:1093:A:OP1	36:5:1093:A:H4'	2.07	0.53
36:5:2144:A:C4	36:5:2281:A:C6	2.96	0.53
36:5:2396:G:OP1	36:5:2397:A:H4'	2.07	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:L2:213:GLY:HA2	36:5:2967:A:H5''	205.73	0.53
36:5:863:C:OP1	86:5:3909:OHX:N3	2.42	0.53
36:5:392:G:O6	86:5:4061:OHX:N3	2.41	0.53
68:O2:33:ARG:HH11	36:5:944:C:H4'	161.69	0.53
1:6:1000:C:H5	1:6:1002:G:H3'	1.73	0.53
1:6:163:G:O5'	1:6:163:G:H8	1.91	0.53
1:6:274:G:N2	1:6:282:C:O2	2.29	0.53
25:D3:69:ARG:NH2	1:6:568:G:N7	366.09	0.53
1:6:891:A:H2'	1:6:892:A:H8	1.72	0.53
48:M1:137:ARG:HG2	37:7:28:C:H5''	308.75	0.53
17:C5:28:MET:O	17:C5:29:SER:HB3	2.08	0.53
21:C9:28:LEU:HD22	21:C9:30:VAL:HG22	1.90	0.53
24:D2:96:ALA:HB3	24:D2:99:PHE:HE1	2.68	0.53
26:D4:29:HIS:CD2	26:D4:29:HIS:N	3.58	0.53
39:L2:96:LEU:HD23	79:Q3:83:ILE:HG23	1.90	0.53
36:1:3138:U:OP2	40:L3:30:LYS:HD3	2.08	0.53
43:L6:176:PHE:HA	50:M4:114:ASP:HB2	2.81	0.53
36:1:148:G:N7	45:L8:137:ASN:ND2	2.56	0.53
54:M8:69:ARG:O	54:M8:72:LYS:HG3	4.06	0.53
54:M8:96:PHE:CD2	54:M8:97:PRO:HD2	2.51	0.53
55:M9:81:ARG:HD2	55:M9:88:ARG:CZ	2.38	0.53
56:N0:33:ASN:C	56:N0:35:VAL:H	2.10	0.53
62:N6:58:VAL:HA	62:N6:104:LEU:HD23	1.91	0.53
62:N6:32:SER:HB2	62:N6:49:PRO:HA	4.17	0.53
62:N6:56:VAL:CG2	62:N6:104:LEU:HB3	2.37	0.53
63:N7:34:LYS:O	63:N7:37:PRO:HG3	5.07	0.53
3:S1:105:PHE:HB3	3:S1:110:LEU:HD11	1.89	0.53
6:S4:239:PRO:O	6:S4:241:GLY:N	3.58	0.53
10:S8:192:TYR:O	10:S8:196:LEU:HB2	2.07	0.53
34:SR:19:TRP:HB2	34:SR:38:ARG:HG3	2.03	0.53
36:1:1916:U:O3'	55:M9:85:ARG:NH2	2.41	0.53
36:1:1134:G:O2'	36:1:2642:A:N3	2.32	0.53
36:1:2770:G:O2'	36:1:2771:U:H5'	2.09	0.53
36:1:2762:A:OP2	86:1:3927:OHX:N4	2.41	0.53
36:1:829:U:H3	36:1:895:A:N6	2.06	0.53
1:2:1067:C:OP1	3:S1:150:VAL:HB	2.08	0.53
1:2:1578:U:O2'	1:2:1579:U:H5'	2.09	0.53
1:2:1586:A:OP1	18:C6:136:SER:N	2.30	0.53
1:2:846:G:H2'	1:2:847:A:O4'	2.09	0.53
36:5:1049:C:H2'	36:5:1050:U:C6	2.43	0.53
36:5:1070:U:C4	36:5:1071:U:C4	2.96	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:1595:U:C2	36:5:1596:C:C5	2.96	0.53
36:5:2518:C:H2'	36:5:2519:A:C8	2.42	0.53
39:L2:69:TYR:HE1	36:5:2557:A:H5''	193.14	0.53
36:5:2875:U:H3	36:5:2952:G:H22	1.56	0.53
36:5:27:C:O2'	36:5:327:A:N3	2.38	0.53
20:C8:140:THR:OG1	1:6:1175:U:O4	352.38	0.53
86:6:2120:OHX:N2	86:6:2171:OHX:N5	2.56	0.53
1:6:922:G:H2'	1:6:923:A:C8	2.43	0.53
15:C3:30:SER:O	15:C3:34:ILE:HG13	2.93	0.53
19:C7:25:THR:O	19:C7:31:ASN:ND2	2.88	0.53
19:C7:24:LEU:HD23	19:C7:34:LEU:HD13	1.89	0.53
24:D2:103:ILE:HG12	24:D2:126:LEU:HB2	1.90	0.53
1:2:54:C:O5'	26:D4:113:ASN:ND2	2.42	0.53
27:D5:61:SER:HB2	27:D5:99:ALA:HB3	1.90	0.53
28:D6:53:LEU:O	28:D6:57:SER:OG	2.21	0.53
40:L3:205:VAL:HG11	40:L3:322:ILE:HD13	3.65	0.53
41:L4:188:ARG:O	41:L4:193:LYS:HE3	2.09	0.53
41:L4:345:GLU:O	41:L4:346:LYS:HB2	4.74	0.53
45:L8:84:ARG:HH12	45:L8:181:LYS:HZ1	1.55	0.53
49:M3:73:ARG:NH1	36:5:110:G:OP2	75.38	0.53
51:M5:99:ARG:HD3	51:M5:167:THR:HB	2.20	0.53
54:M8:89:ASP:OD1	54:M8:90:ASP:N	3.15	0.53
55:M9:17:VAL:HG21	55:M9:52:LYS:HD3	1.89	0.53
56:N0:80:ARG:HB3	56:N0:122:HIS:HB2	1.90	0.53
65:N9:47:LEU:HA	65:N9:50:THR:HG22	3.76	0.53
66:O0:32:LYS:HG3	66:O0:35:ARG:HH21	3.19	0.53
69:O3:6:ARG:HG3	69:O3:8:TYR:CD1	2.64	0.53
5:S3:64:ARG:HA	5:S3:67:ASN:HB2	1.90	0.53
5:S3:64:ARG:HG2	5:S3:65:ARG:H	2.38	0.53
5:S3:143:ARG:HB3	35:SM:109:GLY:HA3	1.90	0.53
35:SM:130:GLU:HA	35:SM:133:GLU:OE1	2.07	0.53
36:1:1063:G:N7	36:1:1097:G:H2'	2.24	0.53
36:1:1780:G:C4	36:1:1781:C:C5	2.97	0.53
36:1:2317:A:OP2	86:1:4065:OHX:N6	2.41	0.53
36:1:2822:U:OP2	86:1:3897:OHX:N5	2.42	0.53
36:1:3118:C:H4'	76:Q0:106:ARG:HH22	1.73	0.53
36:1:915:A:H2'	36:1:915:A:N3	2.23	0.53
1:2:17:C:H4'	1:2:1109:G:C8	2.44	0.53
1:2:577:G:H1	35:SM:99:LYS:HA	1.73	0.53
1:2:844:A:H2'	1:2:845:G:C8	2.44	0.53
36:5:1107:C:H2'	36:5:1108:U:H6	1.73	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:114:A:H2'	36:5:115:A:O4'	2.08	0.53
36:5:209:A:H4'	36:5:211:A:C8	2.43	0.53
56:N0:65:ASN:N	36:5:519:A:N1	318.49	0.53
36:5:595:G:C8	36:5:609:G:C6	2.96	0.53
64:N8:22:ILE:HG22	36:5:642:U:OP1	191.72	0.53
1:6:1013:A:H2'	1:6:1014:G:O4'	2.09	0.53
1:6:1449:U:H2'	1:6:1450:U:C6	2.43	0.53
1:6:1557:U:OP2	1:6:1559:A:O2'	2.21	0.53
1:6:1518:C:OP2	86:6:2143:OHX:N1	2.41	0.53
1:6:52:U:H2'	1:6:53:G:C8	2.43	0.53
17:C5:21:ASP:O	17:C5:25:LEU:N	3.77	0.53
19:C7:22:PRO:HA	34:SR:216:LYS:NZ	2.24	0.53
20:C8:17:LEU:O	20:C8:19:ASN:N	3.09	0.53
22:D0:28:SER:OG	22:D0:111:GLY:O	2.17	0.53
22:D0:28:SER:HB2	22:D0:112:VAL:HA	1.91	0.53
1:2:1100:G:O2'	24:D2:76:SER:N	2.41	0.53
25:D3:130:VAL:HG21	25:D3:135:LEU:HD21	1.89	0.53
26:D4:83:LYS:HE3	26:D4:96:LEU:HB3	2.63	0.53
40:L3:19:ARG:HB3	40:L3:232:ARG:NH1	2.82	0.53
43:L6:14:ASP:N	43:L6:14:ASP:OD2	4.04	0.53
44:L7:136:TYR:O	44:L7:231:ASN:HA	2.08	0.53
44:L7:240:VAL:O	44:L7:242:SER:N	2.42	0.53
44:L7:25:GLN:H	44:L7:28:ALA:HB3	1.72	0.53
46:L9:162:GLN:HB2	46:L9:179:ILE:O	2.09	0.53
48:M1:117:ASP:OD2	48:M1:119:SER:HB3	2.15	0.53
48:M1:43:GLN:NE2	48:M1:70:THR:O	2.33	0.53
52:M6:159:LYS:O	52:M6:162:VAL:HB	2.42	0.53
53:M7:112:LEU:HA	53:M7:151:THR:O	2.53	0.53
53:M7:136:ILE:O	53:M7:137:ASN:ND2	2.43	0.53
53:M7:29:THR:HG22	53:M7:87:SER:OG	2.08	0.53
54:M8:44:PHE:CD1	54:M8:139:ILE:HD11	2.44	0.53
55:M9:134:HIS:CE1	55:M9:137:ALA:HB2	2.44	0.53
59:N3:12:ARG:HG2	59:N3:13:ILE:H	2.90	0.53
60:N4:33:ASN:OD1	60:N4:35:LYS:HB3	2.08	0.53
61:N5:141:TYR:O	61:N5:142:ILE:HB	2.09	0.53
65:N9:39:PHE:O	65:N9:43:HIS:N	2.74	0.53
66:O0:53:LYS:HE2	36:5:2552:C:H5	241.09	0.53
74:O8:56:ILE:HG22	74:O8:58:ASP:HB3	1.91	0.53
79:Q3:84:ARG:O	79:Q3:88:GLU:HG2	2.09	0.53
35:SM:27:LYS:HD2	48:M1:68:HIS:CE1	4.43	0.53
36:1:2724:U:OP1	57:N1:57:TYR:OH	2.24	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:3088:G:H2'	36:1:3089:C:C6	2.43	0.53
36:1:511:G:H2'	36:1:512:U:O4'	2.09	0.53
36:1:5:G:H2'	36:1:6:A:O4'	2.08	0.53
1:2:1265:G:C2	1:2:1266:U:H1'	2.44	0.53
37:3:94:C:H2'	37:3:95:A:H8	1.72	0.53
36:5:1317:A:O2'	36:5:1318:A:H3'	2.07	0.53
36:5:1348:U:H5	36:5:1355:A:C8	2.26	0.53
53:M7:23:ARG:NH2	36:5:1505:C:OP1	128.50	0.53
49:M3:128:ARG:NH2	36:5:168:U:O2'	39.30	0.53
36:5:3306:U:O2'	36:5:3308:C:OP2	2.23	0.53
53:M7:3:ARG:HD2	36:5:398:A:H5''	123.06	0.53
86:5:4014:OHX:N3	86:5:4212:OHX:N1	2.55	0.53
41:L4:73:ARG:HH11	36:5:805:G:H1'	163.98	0.53
1:6:1414:U:O2'	86:6:2047:OHX:N5	2.41	0.53
1:6:241:U:H2'	1:6:242:U:C6	2.43	0.53
1:6:271:A:H1'	1:6:285:G:N2	2.24	0.53
11:S9:149:ARG:HD2	1:6:765:G:O6	431.73	0.53
38:8:142:C:H2'	38:8:143:U:C6	2.43	0.53
13:C1:33:ARG:NH1	13:C1:53:TYR:O	2.95	0.53
16:C4:51:ASP:OD1	1:6:902:G:N1	283.42	0.53
20:C8:25:ASN:HB2	27:D5:40:VAL:HG11	1.90	0.53
7:S5:225:ARG:HH22	30:D8:57:MET:HB2	4.65	0.53
33:E1:98:VAL:HG22	33:E1:99:LYS:H	1.74	0.53
41:L4:320:ASN:OD1	41:L4:323:VAL:HG12	3.16	0.53
46:L9:92:TYR:HB2	46:L9:142:ASP:HB3	1.88	0.53
47:M0:85:PHE:CB	47:M0:140:THR:HG22	2.78	0.53
50:M4:20:VAL:HG13	50:M4:68:LEU:O	2.07	0.53
54:M8:25:TYR:HA	54:M8:28:LEU:HD12	2.61	0.53
58:N2:90:ARG:NH1	58:N2:90:ARG:HB3	4.64	0.53
64:N8:90:TYR:O	64:N8:94:ALA:HB2	2.09	0.53
69:O3:8:TYR:CZ	69:O3:99:ARG:HD3	2.63	0.53
72:O6:58:ILE:HG12	72:O6:59:ASP:N	2.23	0.53
36:1:852:U:C5	79:Q3:2:ALA:HA	2.43	0.53
2:S0:30:GLN:NE2	2:S0:149:LEU:HD13	2.24	0.53
2:S0:175:TYR:HE1	2:S0:197:ILE:HG22	3.90	0.53
4:S2:95:ARG:HD2	4:S2:95:ARG:N	2.24	0.53
7:S5:59:VAL:C	7:S5:61:TYR:H	2.11	0.53
9:S7:155:ASP:OD2	9:S7:156:SER:N	2.35	0.53
35:SM:124:GLN:O	35:SM:127:ALA:N	2.41	0.53
36:1:2359:C:H2'	36:1:2360:C:H6	1.73	0.53
36:1:1055:A:N6	36:1:2637:A:OP2	2.41	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:1933:A:OP2	86:1:3877:OHX:N6	2.41	0.53
36:5:119:U:H4'	36:5:120:G:H5''	1.89	0.53
36:5:1782:U:H2'	36:5:1783:U:C6	2.43	0.53
36:5:1785:U:H2'	36:5:1786:G:H8	1.72	0.53
36:5:2623:G:H2'	36:5:2624:G:H8	1.73	0.53
36:5:3199:G:C2	36:5:3200:G:C8	2.97	0.53
36:5:332:C:N3	38:8:31:G:N1	2.42	0.53
36:5:553:U:H2'	36:5:554:A:O4'	2.09	0.53
1:6:1417:A:OP1	86:6:2086:OHX:N4	2.42	0.53
1:6:407:A:H2'	1:6:408:C:C6	2.44	0.53
1:6:417:A:O5'	1:6:417:A:H8	1.92	0.53
8:S6:173:PRO:HA	1:6:66:U:H5'	340.48	0.53
15:C3:5:HIS:HE1	15:C3:121:ARG:HG3	1.74	0.53
19:C7:5:ARG:O	19:C7:10:LYS:HE3	2.08	0.53
22:D0:46:GLU:HA	22:D0:49:ASN:HA	3.38	0.53
22:D0:58:LEU:HD13	22:D0:88:LYS:HB3	3.45	0.53
1:2:1793:G:O2'	28:D6:5:ARG:NH2	2.42	0.53
39:L2:5:ILE:HG12	39:L2:8:GLN:HG3	2.29	0.53
40:L3:227:GLU:HG2	40:L3:270:ARG:HD3	1.89	0.53
43:L6:39:VAL:O	43:L6:40:LEU:HD23	3.52	0.53
45:L8:254:ASP:O	45:L8:256:ALA:N	2.42	0.53
45:L8:63:LYS:O	45:L8:66:SER:N	3.17	0.53
52:M6:54:TYR:CE2	52:M6:58:LEU:HD13	2.44	0.53
57:N1:102:ARG:O	57:N1:106:LEU:HG	4.69	0.53
60:N4:25:ASP:OD2	60:N4:27:LYS:HB2	2.09	0.53
61:N5:130:TYR:CD1	61:N5:130:TYR:N	2.77	0.53
62:N6:100:HIS:CD2	62:N6:101:PRO:HD2	2.67	0.53
71:O5:95:PHE:O	71:O5:97:ALA:N	2.39	0.53
2:S0:32:HIS:C	2:S0:34:GLU:H	2.10	0.53
3:S1:206:PRO:O	3:S1:207:LEU:HB2	2.09	0.53
3:S1:218:LEU:HD23	3:S1:219:LYS:N	2.24	0.53
6:S4:252:ARG:HH21	6:S4:252:ARG:HB3	4.53	0.53
1:2:164:A:H1'	8:S6:13:GLN:HE22	1.74	0.53
10:S8:138:ASN:OD1	1:6:189:C:N4	273.16	0.53
11:S9:171:ARG:HH11	11:S9:174:ARG:HB3	4.30	0.53
34:SR:176:LYS:HG2	34:SR:197:SER:O	2.08	0.53
36:1:675:C:O2'	36:1:679:U:OP1	2.25	0.53
36:1:915:A:O2'	36:1:917:A:OP1	2.15	0.53
1:2:138:A:O2'	8:S6:149:LYS:NZ	2.42	0.53
1:2:1600:A:O2'	1:2:1602:C:N4	2.42	0.53
1:2:191:C:O2'	1:2:192:U:O5'	2.27	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:458:G:OP2	26:D4:105:ARG:NH2	2.41	0.53
1:2:61:A:C8	1:2:269:G:O2'	2.61	0.53
60:N4:48:ARG:NH1	36:5:2112:U:OP2	232.90	0.53
36:5:2319:U:O4	86:5:3989:OHX:N2	2.42	0.53
36:5:561:C:H2'	36:5:562:C:C6	2.41	0.53
49:M3:73:ARG:HD2	36:5:76:G:H3'	81.41	0.53
10:S8:33:PRO:HB3	1:6:330:G:O2'	274.94	0.53
1:6:363:G:OP1	86:6:2111:OHX:N1	2.41	0.53
1:6:570:A:H5''	1:6:571:G:OP2	2.09	0.53
1:2:1228:G:N1	14:C2:67:THR:HB	2.18	0.53
18:C6:71:GLY:HA2	1:6:1483:A:H4'	410.68	0.53
20:C8:113:LEU:HD21	20:C8:127:HIS:CE1	2.43	0.53
20:C8:91:ASP:O	20:C8:92:ILE:HB	2.07	0.53
23:D1:56:SER:OG	23:D1:59:VAL:HG23	2.24	0.53
39:L2:133:TYR:HE1	39:L2:135:ILE:HD11	1.73	0.53
47:M0:38:LYS:HB2	47:M0:83:ASP:HA	2.26	0.53
50:M4:115:PHE:O	50:M4:119:GLN:HG3	2.08	0.53
50:M4:125:LYS:HA	50:M4:128:ARG:NH1	4.47	0.53
50:M4:24:LYS:HE2	50:M4:25:LYS:NZ	2.24	0.53
51:M5:68:ARG:NH2	36:5:292:U:OP2	150.33	0.53
55:M9:108:LYS:O	55:M9:112:ALA:N	2.37	0.53
56:N0:139:TYR:CD2	56:N0:140:VAL:HG23	2.47	0.53
60:N4:34:SER:HA	60:N4:37:ALA:HB3	1.89	0.53
61:N5:105:VAL:HG21	61:N5:135:ILE:HG13	2.78	0.53
63:N7:10:VAL:HB	63:N7:83:THR:CG2	2.39	0.53
68:O2:4:LEU:HD12	68:O2:5:PRO:HD3	2.69	0.53
53:M7:172:GLN:NE2	69:O3:61:GLY:HA3	2.23	0.53
69:O3:8:TYR:CE1	69:O3:99:ARG:HD3	2.87	0.53
70:O4:58:ARG:O	70:O4:61:GLN:HB2	2.43	0.53
2:S0:117:GLU:O	4:S2:40:LYS:NZ	2.42	0.53
3:S1:81:PHE:CE1	3:S1:109:LYS:HE2	2.44	0.53
5:S3:91:VAL:HG23	5:S3:92:GLN:OE1	2.08	0.53
6:S4:208:VAL:HG21	6:S4:225:VAL:HG21	2.62	0.53
6:S4:23:LEU:H	6:S4:23:LEU:HD22	1.71	0.53
9:S7:111:LYS:O	9:S7:112:ARG:HB2	2.08	0.53
11:S9:146:PHE:O	11:S9:147:MET:HB2	2.36	0.53
34:SR:22:SER:HB2	34:SR:70:ASP:HA	1.90	0.53
36:1:1724:U:H1'	36:1:1725:C:C6	2.44	0.53
36:1:1807:G:C6	36:1:1808:G:N1	2.77	0.53
36:1:2355:G:H4'	53:M7:139:TYR:CZ	2.43	0.53
36:1:3039:C:H2'	36:1:3040:A:H8	1.74	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:3039:C:H2'	36:1:3040:A:C8	2.44	0.53
36:1:3317:U:H4'	36:1:3318:G:O5'	2.09	0.53
36:1:830:A:OP1	86:1:4005:OHX:N4	2.42	0.53
36:1:708:G:H5''	36:1:708:G:H8	1.74	0.53
36:1:86:G:O2'	49:M3:11:LYS:HD3	2.09	0.53
36:1:829:U:H3	36:1:895:A:H62	1.57	0.53
36:5:1113:G:O2'	36:5:1369:A:N3	2.34	0.53
36:5:1685:C:H2'	36:5:1686:U:H6	1.74	0.53
36:5:1451:C:H42	36:5:2353:G:H1	1.57	0.53
46:L9:44:THR:HG22	36:5:3186:A:C2	327.77	0.53
36:5:3294:A:H2'	36:5:3295:A:O4'	2.09	0.53
36:5:34:A:H2'	36:5:35:A:C8	2.44	0.53
36:5:2846:U:O2	86:5:4046:OHX:N5	2.42	0.53
36:5:438:A:H2'	36:5:494:G:H21	1.74	0.53
36:5:563:U:H2'	36:5:564:G:C8	2.44	0.53
1:6:333:A:N6	1:6:334:G:O6	2.42	0.53
37:7:23:A:H2'	37:7:24:A:C8	2.44	0.53
18:C6:71:GLY:O	18:C6:77:GLN:NE2	2.42	0.53
24:D2:11:LEU:O	24:D2:15:ASN:HB2	3.32	0.53
25:D3:87:VAL:HG12	25:D3:92:CYS:HB3	1.89	0.53
26:D4:20:ARG:HB3	26:D4:76:TYR:CD2	2.90	0.53
40:L3:108:GLU:HB2	40:L3:137:TYR:CD1	2.44	0.53
41:L4:11:LEU:HD13	41:L4:159:ILE:HD11	1.90	0.53
36:1:2703:A:N6	42:L5:23:ARG:HG2	2.24	0.53
42:L5:281:GLU:O	42:L5:284:ALA:HB3	3.67	0.53
45:L8:100:GLU:OE2	45:L8:105:LYS:HD2	2.08	0.53
45:L8:56:VAL:O	45:L8:60:ARG:HG3	2.08	0.53
46:L9:29:GLY:HA3	46:L9:82:VAL:HG13	1.91	0.53
50:M4:22:LEU:HD22	50:M4:94:TRP:CH2	2.67	0.53
36:1:1327:C:O3'	69:O3:76:GLY:HA2	2.09	0.53
70:O4:3:GLN:HB3	70:O4:30:LEU:HD12	1.90	0.53
6:S4:159:THR:HG22	6:S4:173:ILE:HB	2.26	0.53
6:S4:163:ASP:HB3	6:S4:166:SER:O	2.09	0.53
7:S5:44:ASN:OD1	7:S5:70:VAL:HG12	2.09	0.53
8:S6:53:SER:OG	8:S6:110:ALA:O	2.22	0.53
8:S6:3:LEU:HD22	8:S6:111:LEU:HD11	3.20	0.53
9:S7:117:THR:HG23	1:6:639:U:P	364.80	0.53
34:SR:64:HIS:ND1	34:SR:86:ASP:OD2	2.34	0.53
36:1:1004:U:C2	36:1:1005:G:C8	2.97	0.53
36:1:1506:A:C4	36:1:1513:G:N2	2.77	0.53
36:1:1547:G:H2'	36:1:1548:C:C6	2.44	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:2225:U:H2'	36:1:2226:U:H6	1.74	0.53
36:1:2641:U:H5''	36:1:2642:A:OP1	2.09	0.53
36:1:2766:U:O4	86:1:4033:OHX:N2	2.41	0.53
36:1:718:G:C2	36:1:721:G:H1'	2.44	0.53
36:1:92:G:H5'	36:1:93:C:H5''	1.89	0.53
36:5:1806:A:H2'	36:5:1807:G:O4'	2.08	0.53
53:M7:80:LYS:NZ	36:5:2389:C:OP1	179.03	0.53
36:5:3084:C:H2'	36:5:3085:G:O4'	2.08	0.53
36:5:679:U:O2'	36:5:788:C:O2	2.27	0.53
18:C6:66:ARG:NH1	1:6:1351:G:OP1	436.32	0.53
1:6:196:G:C2	1:6:197:A:H1'	2.42	0.53
1:6:719:U:N3	1:6:721:U:H5	2.06	0.53
15:C3:73:ARG:HD3	1:6:859:A:C6	330.44	0.53
15:C3:36:GLN:HB3	15:C3:54:LEU:HD21	1.91	0.53
20:C8:123:ARG:NH1	1:6:1546:G:OP1	358.47	0.53
20:C8:46:VAL:HG22	20:C8:72:ILE:HG22	1.90	0.53
27:D5:49:ARG:NH2	27:D5:53:GLU:OE2	2.41	0.53
41:L4:285:ASP:OD1	41:L4:288:ARG:HB2	2.08	0.53
41:L4:93:MET:HB2	36:5:658:G:N2	145.58	0.53
42:L5:158:ARG:HB2	42:L5:158:ARG:HH21	6.16	0.53
48:M1:53:THR:OG1	48:M1:60:ARG:HA	2.09	0.53
54:M8:94:PHE:O	54:M8:96:PHE:N	3.44	0.53
55:M9:127:SER:HB3	55:M9:132:PHE:HD2	1.74	0.53
55:M9:68:GLN:HA	55:M9:71:ARG:HG2	4.29	0.53
56:N0:155:ARG:HD3	56:N0:172:TYR:CD2	4.09	0.53
59:N3:108:GLU:HG2	59:N3:128:ARG:NH1	2.24	0.53
66:O0:66:LYS:N	66:O0:66:LYS:HD2	4.20	0.53
74:O8:61:LYS:O	74:O8:65:LEU:HB2	2.09	0.53
79:Q3:28:LYS:HG2	79:Q3:29:LEU:HG	1.91	0.53
3:S1:27:LYS:HD3	3:S1:47:LEU:HD22	4.26	0.53
3:S1:52:THR:OG1	3:S1:55:LYS:HB3	4.63	0.53
10:S8:116:HIS:CD2	10:S8:146:ARG:HD3	3.91	0.53
11:S9:117:GLY:O	11:S9:119:ALA:N	2.71	0.53
4:S2:175:GLY:HA3	11:S9:53:ARG:HH22	1.74	0.53
36:1:1109:U:H2'	36:1:1110:U:C6	2.44	0.53
36:1:1686:U:O2	36:1:1688:U:H1'	2.08	0.53
36:1:2258:U:H2'	36:1:2259:A:O4'	2.09	0.53
36:1:2381:G:C2'	36:1:2382:G:H5'	2.39	0.53
36:1:2997:G:H1'	36:1:3396:U:H5'	1.91	0.53
36:1:3066:U:O4	86:1:4131:OHX:N5	2.42	0.53
36:1:3203:U:H2'	36:1:3204:C:C6	2.44	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:973:A:P	54:M8:12:ARG:HH12	2.32	0.53
1:2:992:A:O2'	1:2:1785:U:O2	2.26	0.53
1:2:384:G:OP1	86:2:2063:OHX:N5	2.42	0.53
1:2:73:U:H4'	1:2:74:U:OP1	2.09	0.53
38:4:3:A:OP1	86:4:236:OHX:N2	2.42	0.53
36:5:1806:A:OP2	86:5:4018:OHX:N5	2.42	0.53
43:L6:23:LYS:HE3	36:5:503:C:O2	238.39	0.53
1:6:1080:U:O2'	1:6:1081:A:H5'	2.09	0.53
1:6:1148:C:O2'	1:6:1149:G:H5'	2.09	0.53
1:6:1697:G:H8	1:6:1705:C:N3	2.07	0.53
11:S9:133:HIS:CE1	1:6:513:U:H5'	446.17	0.53
42:L5:269:SER:OG	37:7:1:G:N3	316.12	0.53
12:C0:43:ILE:O	12:C0:47:GLN:HB3	3.38	0.53
13:C1:57:LYS:HD3	13:C1:131:ILE:HG23	2.06	0.53
16:C4:13:VAL:N	16:C4:77:THR:OG1	2.42	0.53
24:D2:88:LYS:NZ	1:6:371:G:O3'	372.12	0.53
13:C1:101:GLU:OE1	25:D3:16:ARG:NH2	2.41	0.53
27:D5:93:SER:HB3	27:D5:100:ILE:HB	1.91	0.53
40:L3:117:ARG:HA	40:L3:175:LYS:HD2	4.28	0.53
40:L3:221:THR:HG22	40:L3:272:TYR:N	3.09	0.53
40:L3:4:ARG:HB3	36:5:2882:U:O4	242.51	0.53
40:L3:68:HIS:CD2	40:L3:69:LYS:HG3	2.44	0.53
44:L7:102:VAL:HG13	44:L7:126:LEU:HD22	2.27	0.53
44:L7:110:ARG:CZ	54:M8:3:ILE:HD12	4.34	0.53
44:L7:108:LEU:HD21	44:L7:115:THR:HG23	1.91	0.53
44:L7:52:GLN:O	44:L7:56:GLU:HG2	2.09	0.53
47:M0:3:ARG:CZ	47:M0:63:GLU:HG3	2.38	0.53
48:M1:139:THR:HG22	48:M1:146:GLY:O	2.87	0.53
51:M5:91:GLU:O	51:M5:93:LYS:NZ	3.93	0.53
52:M6:110:PRO:O	52:M6:112:TYR:N	3.14	0.53
52:M6:3:VAL:HG13	52:M6:4:GLU:N	2.24	0.53
53:M7:108:ASP:O	53:M7:111:LYS:N	2.40	0.53
53:M7:50:GLN:O	53:M7:53:ASP:N	2.31	0.53
59:N3:13:ILE:HD12	59:N3:85:TRP:CG	2.44	0.53
49:M3:3:ILE:HG12	64:N8:34:MET:HE1	2.73	0.53
67:O1:103:GLY:HA2	36:5:3325:G:H5''	179.60	0.53
36:1:3275:U:H5'	69:O3:68:TRP:HE1	1.73	0.53
71:O5:24:LEU:HB3	71:O5:51:ILE:HG12	2.19	0.53
5:S3:50:ILE:HB	5:S3:88:ALA:HA	1.91	0.53
6:S4:97:GLU:HG2	6:S4:113:ARG:HH22	6.97	0.53
6:S4:92:LEU:O	6:S4:95:THR:HG22	6.14	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:S5:44:ASN:OD1	7:S5:115:LYS:HB2	3.77	0.53
8:S6:25:ARG:HA	8:S6:28:PHE:CD1	2.44	0.53
9:S7:63:PRO:O	9:S7:64:VAL:HG23	2.09	0.53
36:1:2121:G:HO2'	36:1:2122:G:P	2.31	0.52
36:1:2379:U:H2'	36:1:2380:U:H6	1.72	0.52
36:1:1941:C:H1'	36:1:3362:A:H8	1.73	0.52
36:1:849:C:O2'	36:1:850:U:H5'	2.09	0.52
1:2:1166:A:H2'	1:2:1167:G:O4'	2.10	0.52
1:2:744:U:H5'	1:2:745:U:OP2	2.09	0.52
37:3:3:U:H2'	37:3:4:U:H6	1.74	0.52
36:1:1618:G:H4'	38:4:129:C:H1'	1.91	0.52
36:5:1580:A:OP1	36:5:2522:G:N2	2.40	0.52
70:O4:102:LYS:NZ	36:5:2552:C:OP1	231.12	0.52
36:5:507:U:H2'	36:5:508:U:H6	1.72	0.52
36:5:656:A:H2'	36:5:657:A:H8	1.73	0.52
36:5:731:U:H2'	36:5:732:C:H6	1.72	0.52
4:S2:168:ARG:NE	1:6:1098:U:OP2	385.20	0.52
11:S9:7:THR:HG21	1:6:758:U:OP1	384.41	0.52
1:6:871:G:H2'	1:6:872:G:C8	2.44	0.52
26:D4:120:GLY:HA2	1:6:85:A:O3'	336.02	0.52
40:L3:346:THR:O	40:L3:348:ARG:N	2.63	0.52
42:L5:261:THR:HG23	42:L5:264:GLN:OE1	2.09	0.52
42:L5:286:VAL:HG13	47:M0:206:LEU:HD22	1.91	0.52
46:L9:20:ILE:HG12	46:L9:25:VAL:HG22	5.22	0.52
47:M0:31:ILE:HA	47:M0:66:GLU:HB2	1.91	0.52
52:M6:171:LYS:O	52:M6:175:THR:HG23	3.69	0.52
72:O6:26:ILE:O	72:O6:29:LYS:N	2.42	0.52
2:S0:90:ALA:HB1	2:S0:95:ALA:O	3.34	0.52
3:S1:217:LEU:O	3:S1:218:LEU:HB3	2.09	0.52
6:S4:36:HIS:CG	6:S4:85:GLY:HA3	2.44	0.52
9:S7:77:LEU:HD22	9:S7:81:LEU:HD11	1.92	0.52
34:SR:16:HIS:CE1	34:SR:37:SER:HB3	2.44	0.52
34:SR:11:GLY:HA3	34:SR:54:PHE:HB2	1.89	0.52
36:1:109:A:H8	36:1:109:A:O5'	1.92	0.52
36:1:1564:U:H2'	36:1:1565:G:C8	2.43	0.52
36:1:1661:G:H2'	36:1:1662:G:C8	2.44	0.52
36:1:1809:A:H2'	36:1:1810:A:O4'	2.08	0.52
36:1:1818:U:H2'	36:1:1819:U:O4'	2.09	0.52
36:1:971:G:H2'	36:1:972:A:O4'	2.09	0.52
1:2:1061:A:H2'	1:2:1062:A:H5'	1.91	0.52
1:2:1586:A:H1'	1:2:1611:A:N6	2.24	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:737:A:HO2'	1:2:738:G:H8	1.55	0.52
1:2:932:U:OP2	3:S1:155:TYR:OH	2.27	0.52
1:2:97:C:H2'	1:2:98:U:H6	1.74	0.52
36:5:1944:U:H2'	36:5:1945:A:C8	2.44	0.52
36:5:3269:U:H5'	36:5:3271:G:O4'	2.09	0.52
36:5:339:C:OP1	36:5:1380:G:O2'	2.27	0.52
1:6:276:C:O2'	1:6:277:U:H5''	2.08	0.52
25:D3:19:ARG:NH1	1:6:610:G:H21	343.05	0.52
1:6:876:G:H2'	1:6:936:G:N2	2.24	0.52
14:C2:56:GLU:OE1	14:C2:124:LYS:NZ	2.97	0.52
1:2:901:G:N2	16:C4:54:GLU:OE1	2.43	0.52
22:D0:72:ASN:HD22	22:D0:73:GLY:H	1.56	0.52
25:D3:69:ARG:NH1	25:D3:116:ASP:OD1	2.48	0.52
27:D5:41:ILE:HG13	27:D5:42:LEU:HG	1.91	0.52
27:D5:38:HIS:HE1	27:D5:70:LYS:HA	1.72	0.52
23:D1:71:ARG:HH11	29:D7:4:VAL:HG11	2.62	0.52
40:L3:290:ASP:C	40:L3:292:ALA:H	4.05	0.52
40:L3:286:GLY:HA3	40:L3:321:PHE:CZ	2.72	0.52
41:L4:316:ASN:HD22	41:L4:319:LYS:HE2	1.75	0.52
43:L6:175:LYS:O	50:M4:117:ARG:NH2	2.43	0.52
46:L9:41:ILE:HD12	46:L9:71:VAL:HG21	1.91	0.52
48:M1:150:ASN:C	48:M1:151:SER:O	4.34	0.52
50:M4:17:VAL:HG11	50:M4:74:ARG:HA	2.08	0.52
51:M5:172:ARG:HH11	36:5:30:G:P	107.89	0.52
51:M5:4:TYR:O	51:M5:7:LEU:N	2.42	0.52
55:M9:25:ASP:OD1	55:M9:25:ASP:N	2.40	0.52
61:N5:28:THR:OG1	61:N5:29:SER:N	2.69	0.52
64:N8:127:ALA:O	64:N8:148:ILE:HG12	2.35	0.52
68:O2:94:ALA:HB3	68:O2:119:VAL:HG22	1.90	0.52
70:O4:41:ARG:HB2	70:O4:43:LYS:HE3	4.91	0.52
73:O7:10:LYS:HB2	36:5:818:C:H5''	155.59	0.52
73:O7:52:LYS:HA	73:O7:55:ARG:HD2	1.91	0.52
79:Q3:4:ARG:NH2	36:5:838:G:O6	237.51	0.52
2:S0:110:TYR:CE2	4:S2:64:LYS:HB3	2.42	0.52
2:S0:195:TRP:CE2	2:S0:197:ILE:HG13	2.44	0.52
3:S1:195:LYS:HA	3:S1:198:GLU:HB3	1.92	0.52
5:S3:27:ARG:O	12:C0:58:GLN:NE2	3.19	0.52
7:S5:99:MET:HG3	7:S5:180:ARG:HH21	1.73	0.52
7:S5:186:ASN:OD1	7:S5:187:ILE:N	2.91	0.52
8:S6:87:ARG:NH2	1:6:161:U:OP2	315.81	0.52
36:1:1064:A:H5''	36:1:1066:G:O4'	2.09	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:1708:C:H2'	36:1:1709:C:C6	2.44	0.52
36:1:2240:G:H2'	36:1:2241:U:O4'	2.10	0.52
36:1:2413:A:H2'	36:1:2414:G:H8	1.74	0.52
36:1:2516:U:O2'	36:1:2595:A:N6	2.41	0.52
1:2:1104:U:OP1	25:D3:14:LYS:NZ	2.35	0.52
1:2:1275:A:N3	5:S3:141:LYS:NZ	2.50	0.52
36:5:1049:C:C2	36:5:1050:U:C5	2.98	0.52
36:5:1112:A:H5''	36:5:1113:G:OP2	2.09	0.52
36:5:1132:C:HO2'	36:5:2865:U:HO2'	1.57	0.52
36:5:1155:C:H2'	36:5:1156:C:C6	2.42	0.52
36:5:1565:G:N2	36:5:1566:A:H1'	2.24	0.52
36:5:1584:U:H2'	36:5:1585:C:C6	2.44	0.52
36:5:2594:C:H2'	36:5:2595:A:O4'	2.09	0.52
36:5:286:U:H2'	36:5:287:G:C8	2.44	0.52
36:5:3287:U:H2'	36:5:3288:G:H5'	1.90	0.52
1:6:1489:U:C4	1:6:1513:G:C6	2.97	0.52
1:6:1496:U:HO2'	1:6:1519:U:HO2'	1.52	0.52
13:C1:23:PRO:O	13:C1:25:VAL:N	3.35	0.52
24:D2:105:THR:OG1	24:D2:126:LEU:HG	2.08	0.52
24:D2:86:ILE:HD12	24:D2:87:GLU:HG3	1.91	0.52
26:D4:35:VAL:HG13	26:D4:36:SER:H	1.87	0.52
30:D8:11:LYS:HE2	30:D8:33:LEU:HD21	1.91	0.52
46:L9:48:VAL:HG21	46:L9:52:LEU:HD13	2.36	0.52
47:M0:210:ILE:HG12	47:M0:217:PHE:CZ	2.44	0.52
48:M1:48:SER:HB2	48:M1:66:ALA:O	2.24	0.52
49:M3:18:TRP:O	49:M3:20:GLU:N	2.42	0.52
55:M9:172:ARG:O	55:M9:176:ARG:HG2	2.29	0.52
55:M9:4:LEU:HB3	55:M9:7:GLN:HG2	4.45	0.52
55:M9:88:ARG:O	36:5:1779:C:N4	207.17	0.52
57:N1:96:ILE:HG22	57:N1:97:LYS:O	4.17	0.52
64:N8:111:LYS:HA	64:N8:129:PHE:O	2.51	0.52
66:O0:26:GLY:O	66:O0:30:THR:HG23	2.10	0.52
68:O2:123:LYS:HA	68:O2:126:LEU:HD12	2.67	0.52
2:S0:179:ARG:HH11	2:S0:183:ARG:HH11	1.57	0.52
5:S3:26:THR:HG23	5:S3:34:TYR:HD1	2.38	0.52
6:S4:95:THR:HG23	6:S4:97:GLU:H	5.11	0.52
8:S6:157:VAL:HG11	8:S6:175:ILE:HD11	2.90	0.52
10:S8:197:THR:HA	10:S8:200:LYS:HB2	1.91	0.52
10:S8:36:THR:HB	10:S8:57:ALA:O	2.08	0.52
34:SR:166:SER:O	34:SR:166:SER:OG	2.24	0.52
36:1:1019:G:H2'	36:1:1020:G:O4'	2.09	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:2700:G:O2'	36:1:2705:A:N1	2.32	0.52
1:2:1650:U:H2'	1:2:1651:A:H8	1.72	0.52
1:2:386:G:C6	1:2:387:A:N6	2.77	0.52
1:2:952:A:O2'	15:C3:114:ARG:HG3	2.09	0.52
36:5:136:G:H2'	36:5:137:G:C8	2.45	0.52
36:5:662:U:H2'	36:5:663:C:C6	2.45	0.52
1:6:1257:U:O2'	1:6:1258:U:O5'	2.27	0.52
1:6:1372:U:H2'	1:6:1373:C:C6	2.45	0.52
8:S6:133:LEU:HD13	1:6:166:C:O2	327.16	0.52
11:S9:54:ARG:HG3	1:6:1:U:C4	393.02	0.52
1:6:1726:G:N7	86:6:2147:OHX:N5	2.58	0.52
11:S9:133:HIS:HE1	1:6:512:A:O2'	447.80	0.52
1:6:82:U:H2'	1:6:83:G:O4'	2.09	0.52
1:6:903:U:H2'	1:6:905:A:OP2	2.10	0.52
13:C1:128:CYS:O	13:C1:129:ARG:HB3	4.33	0.52
14:C2:88:LEU:O	14:C2:89:ILE:HB	2.47	0.52
17:C5:85:ILE:HG13	17:C5:114:HIS:O	2.64	0.52
21:C9:53:TRP:CH2	21:C9:100:ILE:HD12	2.89	0.52
24:D2:18:GLU:HG2	24:D2:65:LEU:HG	1.91	0.52
29:D7:50:ALA:HB1	29:D7:52:THR:O	2.10	0.52
29:D7:61:THR:HG23	29:D7:62:ILE:N	2.25	0.52
33:E1:108:VAL:HG12	33:E1:114:VAL:HG13	2.18	0.52
39:L2:5:ILE:HD11	39:L2:232:GLY:HA2	3.11	0.52
36:1:3316:A:N6	40:L3:124:LYS:HG2	2.24	0.52
40:L3:139:GLN:HE22	40:L3:143:GLY:H	2.99	0.52
40:L3:187:SER:O	40:L3:189:SER:N	2.96	0.52
41:L4:25:VAL:O	41:L4:28:ALA:N	2.38	0.52
41:L4:269:SER:C	41:L4:271:LYS:H	2.83	0.52
42:L5:274:GLN:OE1	37:7:60:G:N2	332.79	0.52
42:L5:56:THR:OG1	42:L5:59:ASP:HB3	2.09	0.52
42:L5:85:ARG:HH21	42:L5:254:LYS:N	2.04	0.52
45:L8:134:TYR:CD2	45:L8:190:VAL:HG11	4.08	0.52
45:L8:160:ILE:HG23	45:L8:164:VAL:CG1	5.69	0.52
45:L8:85:ASN:N	45:L8:85:ASN:OD1	4.16	0.52
51:M5:73:ARG:HG2	51:M5:75:VAL:HG22	2.28	0.52
55:M9:168:ALA:HB1	55:M9:172:ARG:CZ	2.40	0.52
62:N6:71:SER:OG	62:N6:83:ASP:N	4.28	0.52
67:O1:46:THR:HG21	67:O1:91:SER:OG	2.09	0.52
70:O4:59:PRO:HB3	36:5:1654:A:N3	165.82	0.52
72:O6:83:ALA:O	72:O6:87:VAL:HG23	2.10	0.52
75:O9:21:ARG:HD3	75:O9:22:PRO:O	2.09	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:S4:126:VAL:HG12	6:S4:158:ASP:O	2.09	0.52
7:S5:109:LYS:O	7:S5:113:ILE:HG13	2.09	0.52
8:S6:5:ILE:HA	8:S6:111:LEU:O	2.09	0.52
8:S6:164:LYS:O	8:S6:166:GLU:N	2.43	0.52
11:S9:107:ARG:O	11:S9:147:MET:HA	2.09	0.52
17:C5:119:PHE:HA	35:SM:57:ASN:ND2	2.86	0.52
36:1:1014:U:C2'	36:1:1015:U:H5''	2.39	0.52
36:1:1144:U:H1'	36:1:1145:G:C8	2.45	0.52
36:1:839:C:H4'	36:1:1724:U:H3'	1.91	0.52
36:1:2273:G:O6	86:1:4135:OHX:N5	2.43	0.52
36:1:244:G:C2	36:1:245:U:H1'	2.44	0.52
36:1:2510:U:O2'	36:1:2511:A:H8	1.93	0.52
36:1:2714:G:H4'	36:1:2715:A:O5'	2.09	0.52
36:1:386:A:H2'	36:1:387:A:O4'	2.09	0.52
36:1:918:C:H2'	36:1:919:U:H5'	1.92	0.52
1:2:1222:C:H2'	1:2:1223:A:C8	2.44	0.52
1:2:1449:U:H2'	1:2:1450:U:C6	2.45	0.52
1:2:274:G:C2	1:2:275:C:H1'	2.44	0.52
1:2:331:A:H5'	10:S8:33:PRO:HA	1.91	0.52
1:2:218:A:N6	1:2:844:A:H1'	2.24	0.52
36:5:2603:G:O6	86:5:3897:OHX:N1	2.43	0.52
36:5:3122:A:H2'	36:5:3123:A:H5'	1.92	0.52
36:5:2827:U:O4	86:5:3894:OHX:N6	2.42	0.52
36:5:2767:U:O4	86:5:4113:OHX:N3	2.42	0.52
37:7:4:U:H2'	37:7:5:G:C8	2.44	0.52
12:C0:81:ASN:HB2	14:C2:37:VAL:HG13	1.92	0.52
16:C4:23:PHE:HE2	16:C4:91:THR:HG21	1.74	0.52
21:C9:117:SER:OG	21:C9:118:PRO:O	2.26	0.52
21:C9:141:GLU:C	21:C9:143:ASP:H	3.41	0.52
29:D7:28:PRO:HB3	1:6:959:U:H5'	352.08	0.52
40:L3:152:LYS:HD3	40:L3:189:SER:HA	1.92	0.52
40:L3:92:TYR:CE1	40:L3:159:ARG:HD2	2.62	0.52
41:L4:340:GLY:HA3	36:5:577:C:O2'	283.43	0.52
42:L5:140:ARG:HD3	36:5:1080:A:OP1	226.21	0.52
47:M0:76:MET:HE3	47:M0:148:VAL:HA	2.34	0.52
47:M0:17:TYR:O	47:M0:96:VAL:HB	2.09	0.52
57:N1:68:THR:HG21	36:5:2736:A:O2'	223.00	0.52
59:N3:66:LYS:O	59:N3:68:GLU:N	2.42	0.52
60:N4:4:GLU:HG3	60:N4:5:ILE:H	1.75	0.52
63:N7:103:GLN:HB3	63:N7:105:SER:OG	3.71	0.52
64:N8:128:ARG:HB2	72:O6:8:ALA:HB2	4.55	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
69:O3:70:LYS:O	69:O3:70:LYS:HG2	2.09	0.52
70:O4:44:CYS:HB3	70:O4:49:SER:H	3.43	0.52
74:O8:14:LEU:HD23	74:O8:17:ARG:HD3	1.90	0.52
66:O0:86:ARG:NH1	79:Q3:44:LYS:HE3	5.10	0.52
4:S2:175:GLY:HA3	11:S9:97:LEU:O	3.43	0.52
9:S7:77:LEU:O	9:S7:81:LEU:HB2	2.72	0.52
34:SR:165:ASP:O	34:SR:167:VAL:N	2.43	0.52
36:1:1004:U:N3	36:1:1005:G:N7	2.58	0.52
36:1:1481:A:H2'	36:1:1481:A:N3	2.25	0.52
36:1:2932:U:O2	36:1:2934:A:H8	1.93	0.52
36:1:549:U:H2'	36:1:550:A:C8	2.44	0.52
1:2:1188:G:O2'	1:2:1430:U:OP1	2.23	0.52
1:2:641:G:H2'	1:2:642:G:C8	2.44	0.52
1:2:793:A:OP2	1:2:793:A:H8	1.91	0.52
1:2:993:A:OP1	1:2:1777:G:N2	2.40	0.52
36:5:1414:G:O6	86:5:4142:OHX:N1	2.42	0.52
36:5:1843:C:H2'	36:5:1844:C:H6	1.74	0.52
36:5:2912:G:H1'	36:5:3131:U:OP1	2.09	0.52
36:5:2960:C:OP1	86:5:3965:OHX:N5	2.42	0.52
86:5:3971:OHX:N6	86:5:4193:OHX:N5	2.58	0.52
1:6:235:G:H2'	1:6:236:A:C8	2.44	0.52
1:6:240:U:H4'	1:6:241:U:OP2	2.09	0.52
1:6:36:C:C2	1:6:473:A:C2	2.97	0.52
86:7:220:OHX:N4	86:7:228:OHX:N6	2.57	0.52
17:C5:55:GLY:O	17:C5:58:LYS:HB2	2.10	0.52
2:S0:88:LYS:NZ	19:C7:82:ASP:OD2	2.40	0.52
27:D5:42:LEU:HD12	27:D5:43:ASP:H	1.74	0.52
28:D6:58:VAL:HG22	28:D6:59:TYR:H	4.03	0.52
33:E1:144:CYS:HB3	33:E1:147:VAL:HG12	3.10	0.52
40:L3:150:ARG:HH11	40:L3:150:ARG:CG	2.77	0.52
40:L3:213:GLU:H	40:L3:282:ILE:HB	3.28	0.52
41:L4:209:TYR:O	41:L4:230:VAL:HG22	2.41	0.52
42:L5:104:LEU:HD11	42:L5:108:ARG:NH2	2.81	0.52
48:M1:40:LEU:HD11	48:M1:79:ILE:HG23	2.67	0.52
54:M8:24:VAL:HA	54:M8:27:LYS:HE3	1.91	0.52
54:M8:44:PHE:O	54:M8:48:VAL:HG23	2.08	0.52
56:N0:40:ARG:O	56:N0:43:TYR:HB3	2.09	0.52
59:N3:127:PRO:O	59:N3:130:ALA:HB3	2.09	0.52
63:N7:51:LEU:HD23	63:N7:52:LYS:HG2	8.26	0.52
79:Q3:52:ALA:HB1	79:Q3:68:ALA:HA	1.91	0.52
2:S0:125:ASP:OD1	2:S0:127:ARG:HB3	2.10	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:S2:44:LEU:HD22	4:S2:49:LYS:HB2	3.23	0.52
4:S2:88:LYS:HB3	4:S2:95:ARG:HD2	4.92	0.52
6:S4:86:PHE:O	6:S4:87:MET:HB2	2.09	0.52
36:1:1899:G:N7	86:1:3924:OHX:N3	2.58	0.52
36:1:2339:C:OP2	59:N3:48:ARG:HG3	2.10	0.52
36:1:2445:A:H61	36:1:2502:A:H2	1.57	0.52
36:1:2534:G:H2'	36:1:2535:A:H8	1.74	0.52
36:1:3280:U:O2'	36:1:3281:U:OP2	2.25	0.52
36:1:2169:G:O6	86:1:3905:OHX:N4	2.43	0.52
1:2:1414:U:O2'	86:2:2025:OHX:N4	2.43	0.52
36:5:736:A:C6	36:5:737:G:H1'	2.44	0.52
1:6:515:A:H2'	1:6:516:G:O4'	2.10	0.52
1:6:696:C:H4'	1:6:697:C:H6	1.75	0.52
1:6:811:A:C2	1:6:858:G:H1'	2.44	0.52
19:C7:47:ARG:HD2	19:C7:48:ASN:OD1	2.10	0.52
19:C7:99:VAL:CB	19:C7:118:PRO:HB2	2.40	0.52
5:S3:7:LYS:HD3	22:D0:88:LYS:HE2	1.91	0.52
26:D4:57:VAL:HA	26:D4:73:GLY:HA2	1.92	0.52
28:D6:24:VAL:HG11	28:D6:71:LEU:HD12	1.91	0.52
40:L3:361:THR:HG22	40:L3:371:GLN:OE1	2.38	0.52
41:L4:120:TYR:CD2	41:L4:277:PRO:HG3	3.07	0.52
42:L5:292:ALA:O	42:L5:295:GLY:N	2.43	0.52
42:L5:85:ARG:HH12	42:L5:254:LYS:H	4.58	0.52
47:M0:99:ILE:CD1	47:M0:101:LYS:HB2	5.50	0.52
49:M3:90:ALA:HB1	49:M3:95:ILE:HD12	2.52	0.52
36:1:1544:G:H5'	51:M5:67:ARG:NE	2.23	0.52
52:M6:78:ARG:CG	52:M6:78:ARG:HH11	2.22	0.52
56:N0:75:PHE:HB2	56:N0:94:ILE:O	2.35	0.52
63:N7:46:ILE:HD11	63:N7:49:TYR:CG	2.45	0.52
86:1:4194:OHX:N6	86:O1:202:OHX:N3	2.57	0.52
5:S3:74:GLN:HG3	5:S3:79:TYR:HB2	1.92	0.52
6:S4:43:PRO:HA	6:S4:82:TYR:O	2.10	0.52
7:S5:144:GLU:HG3	7:S5:221:ALA:HB1	1.91	0.52
8:S6:49:VAL:HB	8:S6:115:LYS:HG3	4.83	0.52
9:S7:11:GLN:OE1	9:S7:12:ALA:N	2.34	0.52
36:1:109:A:H4'	36:1:110:G:OP1	2.08	0.52
36:1:1660:C:H2'	36:1:1661:G:C8	2.45	0.52
36:1:2190:U:C4	36:1:2191:U:C4	2.98	0.52
36:1:2352:A:OP1	53:M7:82:ARG:HD3	2.10	0.52
36:1:550:A:N6	36:1:551:A:H62	2.07	0.52
1:2:1313:A:H2'	1:2:1315:U:H5'	1.92	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:1527:C:H2'	1:2:1528:U:H6	1.75	0.52
1:2:1395:G:O6	86:2:2097:OHX:N6	2.43	0.52
65:N9:10:HIS:NE2	36:5:1139:G:O6	225.63	0.52
36:5:142:C:H2'	36:5:143:G:O4'	2.10	0.52
36:5:1440:G:H2'	36:5:1441:G:O4'	2.10	0.52
45:L8:136:LEU:HB2	36:5:147:U:H5'	116.54	0.52
36:5:2308:C:O2	86:5:4234:OHX:N1	2.43	0.52
36:5:257:U:H2'	36:5:258:G:C8	2.45	0.52
36:5:2881:C:H2'	36:5:2882:U:C6	2.44	0.52
36:5:2896:A:H5'	36:5:2896:A:H8	1.74	0.52
1:6:1489:U:O2'	1:6:1490:C:OP2	2.28	0.52
30:D8:18:ARG:NH1	1:6:1616:G:H4'	364.89	0.52
8:S6:136:LYS:NZ	1:6:66:U:OP1	336.64	0.52
21:C9:16:ASN:HA	21:C9:56:LYS:NZ	2.29	0.52
23:D1:25:LYS:HB2	23:D1:28:ASP:HB2	5.28	0.52
23:D1:36:VAL:HG11	23:D1:78:LEU:HD13	1.92	0.52
24:D2:107:SER:HA	1:6:804:A:C8	367.59	0.52
25:D3:61:SER:OG	25:D3:67:ALA:N	2.75	0.52
26:D4:51:GLU:O	26:D4:53:ASP:N	3.68	0.52
28:D6:44:ILE:HD12	28:D6:45:VAL:N	2.25	0.52
40:L3:187:SER:OG	40:L3:188:ILE:HD12	2.10	0.52
40:L3:18:PRO:HG2	40:L3:20:LYS:HD2	2.24	0.52
42:L5:40:HIS:HB3	42:L5:43:LYS:HD2	3.81	0.52
45:L8:156:ASP:O	45:L8:183:LYS:HE3	7.81	0.52
45:L8:200:LEU:HD13	45:L8:203:VAL:HG23	3.30	0.52
45:L8:73:PRO:HD3	45:L8:233:TRP:CG	3.57	0.52
48:M1:160:VAL:O	48:M1:163:PHE:N	3.13	0.52
51:M5:199:LEU:HD13	51:M5:203:ARG:CZ	3.08	0.52
54:M8:24:VAL:O	54:M8:28:LEU:HG	2.10	0.52
57:N1:101:CYS:HB3	36:5:990:U:H1'	251.41	0.52
58:N2:43:VAL:HG11	58:N2:50:LEU:HA	1.92	0.52
63:N7:13:VAL:HG12	63:N7:14:VAL:H	3.00	0.52
63:N7:76:ASN:HB3	63:N7:79:HIS:ND1	2.25	0.52
5:S3:101:GLN:O	5:S3:104:SER:OG	2.15	0.52
5:S3:111:ASN:HD22	5:S3:111:ASN:N	2.07	0.52
34:SR:307:ASP:O	34:SR:309:VAL:HG23	2.32	0.52
34:SR:5:GLU:HG2	34:SR:317:THR:HG23	3.83	0.52
36:1:1245:A:H3'	36:1:1246:G:H5''	1.92	0.52
36:1:1461:A:H2'	36:1:1462:A:C8	2.45	0.52
36:1:1699:A:H2'	36:1:1700:G:C8	2.45	0.52
36:1:265:A:H5''	36:1:266:A:OP2	2.09	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:2778:G:H2'	36:1:2779:A:H5'	1.91	0.52
36:1:824:C:H2'	36:1:825:U:H6	1.74	0.52
36:1:94:G:H2'	36:1:95:A:C8	2.45	0.52
1:2:1107:G:H3'	1:2:1108:G:H21	1.73	0.52
1:2:1158:C:OP2	86:2:2173:OHX:N5	2.43	0.52
1:2:795:U:C5	1:2:796:A:C8	2.98	0.52
1:2:973:A:H2'	1:2:974:A:C8	2.43	0.52
37:3:79:A:C2	37:3:102:A:C4	2.98	0.52
36:5:1401:A:C2	36:5:1411:C:C2	2.97	0.52
36:5:3060:C:H2'	36:5:3061:G:O4'	2.10	0.52
36:5:300:G:O6	86:5:4186:OHX:N2	2.43	0.52
36:5:359:U:H4'	36:5:817:A:N6	2.25	0.52
1:6:882:U:H2'	1:6:883:C:H6	1.72	0.52
14:C2:43:ARG:HA	14:C2:121:VAL:HG12	3.21	0.52
21:C9:10:ALA:HB3	21:C9:13:ASP:HB2	1.92	0.52
5:S3:8:LYS:HE2	22:D0:61:LYS:HD3	1.91	0.52
1:2:637:C:OP1	24:D2:32:LYS:HG3	2.08	0.52
25:D3:96:VAL:HG12	25:D3:127:VAL:HG21	4.55	0.52
26:D4:76:TYR:OH	26:D4:86:GLU:OE2	2.20	0.52
1:2:871:G:O2'	29:D7:66:PRO:HB2	2.10	0.52
30:D8:26:THR:O	30:D8:44:VAL:HG22	2.40	0.52
39:L2:209:HIS:HD2	39:L2:210:PRO:N	2.07	0.52
39:L2:3:ARG:HD3	36:5:911:C:N4	178.86	0.52
40:L3:103:THR:HG21	40:L3:147:GLU:OE2	2.50	0.52
41:L4:185:LYS:HZ1	41:L4:201:GLN:HG2	1.74	0.52
42:L5:60:ILE:HB	42:L5:80:SER:HB2	1.91	0.52
43:L6:76:LEU:HD12	43:L6:138:GLN:HA	1.91	0.52
43:L6:165:LEU:HD11	69:O3:102:LEU:HD11	1.91	0.52
49:M3:123:ILE:HG22	71:O5:118:ILE:HG12	2.51	0.52
52:M6:190:VAL:HG23	52:M6:194:LEU:HD12	1.91	0.52
54:M8:63:SER:OG	54:M8:65:SER:N	2.42	0.52
54:M8:83:VAL:HB	54:M8:103:ALA:HB2	1.92	0.52
50:M4:60:LEU:HD13	56:N0:152:LEU:HD11	1.92	0.52
44:L7:73:GLY:O	57:N1:143:THR:HB	2.10	0.52
59:N3:80:ARG:HE	59:N3:97:ASP:CG	2.13	0.52
62:N6:59:VAL:HG22	62:N6:103:LYS:O	6.29	0.52
71:O5:7:TYR:HA	71:O5:10:ARG:HE	2.87	0.52
36:1:155:G:H1'	72:O6:26:ILE:CD1	2.39	0.52
36:1:3120:C:H3'	76:Q0:111:ARG:HH21	1.75	0.52
2:S0:147:THR:O	2:S0:161:PRO:HA	4.75	0.52
6:S4:158:ASP:OD1	6:S4:158:ASP:N	2.42	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:S9:93:LEU:HA	11:S9:96:VAL:HG13	1.91	0.52
35:SM:128:ALA:O	35:SM:131:ILE:HG22	2.10	0.52
34:SR:175:ASP:N	34:SR:175:ASP:OD1	2.42	0.52
34:SR:203:THR:OG1	34:SR:204:ALA:N	2.40	0.52
36:1:1708:C:H2'	36:1:1709:C:H6	1.75	0.52
36:1:1821:U:C2	70:O4:67:LYS:HB2	2.45	0.52
36:1:3139:A:H8	36:1:3139:A:H5''	1.75	0.52
36:1:542:G:H1	36:1:549:U:H3	1.56	0.52
1:2:1173:C:H2'	1:2:1174:C:H6	1.75	0.52
1:2:1183:A:C4	17:C5:100:LYS:HD3	2.44	0.52
1:2:1214:U:OP1	1:2:1246:C:H1'	2.10	0.52
1:2:1433:G:H2'	1:2:1434:U:C6	2.45	0.52
1:2:487:G:H3'	1:2:488:G:H5''	1.91	0.52
1:2:702:G:N2	1:2:703:G:H1'	2.25	0.52
1:2:778:G:H3'	1:2:780:A:C2	2.44	0.52
38:4:47:C:H1'	38:4:61:A:H2'	1.93	0.52
36:5:1596:C:H2'	36:5:1597:C:C6	2.45	0.52
36:5:1615:C:H2'	36:5:1616:U:C6	2.43	0.52
36:5:1514:G:C6	36:5:1841:A:C5	2.97	0.52
36:5:2106:A:H2'	36:5:2107:A:H8	1.75	0.52
36:5:2694:A:N6	36:5:2695:A:N6	2.57	0.52
51:M5:44:ARG:NH1	36:5:269:G:OP1	124.20	0.52
86:5:4014:OHX:N6	86:5:4212:OHX:N4	2.58	0.52
1:6:1001:A:C6	1:6:1002:G:C6	2.97	0.52
1:6:1488:G:O2'	1:6:1494:C:O2	2.20	0.52
1:6:1482:C:OP2	1:6:1521:G:N1	2.42	0.52
1:6:1714:A:H2'	1:6:1715:G:O4'	2.10	0.52
10:S8:52:ASN:OD1	86:6:2136:OHX:N3	310.81	0.52
1:6:25:C:H1'	1:6:26:A:OP2	2.10	0.52
1:6:431:C:H2'	1:6:432:G:C8	2.45	0.52
8:S6:171:LYS:NZ	1:6:68:A:OP2	350.33	0.52
1:6:76:A:H2'	1:6:76:A:N3	2.23	0.52
13:C1:90:TYR:HD1	13:C1:90:TYR:C	2.14	0.52
40:L3:347:SER:HB2	40:L3:350:ALA:HB2	4.15	0.52
41:L4:158:SER:HA	41:L4:213:ASN:O	2.10	0.52
41:L4:89:ALA:C	41:L4:91:GLY:H	2.13	0.52
36:1:1385:C:HO2'	43:L6:2:SER:N	2.07	0.52
46:L9:20:ILE:HG23	46:L9:25:VAL:HA	1.92	0.52
49:M3:121:SER:OG	49:M3:121:SER:O	3.49	0.52
54:M8:135:GLN:OE1	54:M8:135:GLN:N	2.42	0.52
55:M9:146:LYS:O	55:M9:149:ALA:N	2.61	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:N0:1:MET:HE2	56:N0:4:PHE:CE1	2.45	0.52
57:N1:100:LYS:HB3	36:5:990:U:H4'	258.59	0.52
63:N7:23:VAL:HG12	63:N7:45:GLY:HA3	2.18	0.52
63:N7:29:HIS:CE1	63:N7:42:LEU:HD11	2.45	0.52
79:Q3:9:GLY:O	36:5:836:A:O2'	235.60	0.52
2:S0:112:THR:HG22	2:S0:115:PHE:HB2	2.05	0.52
3:S1:29:TRP:CZ3	3:S1:45:LYS:HD2	2.45	0.52
5:S3:162:GLN:O	5:S3:165:ASN:N	2.42	0.52
7:S5:43:PHE:CZ	7:S5:90:ILE:HG21	2.45	0.52
10:S8:103:GLN:HB3	10:S8:164:ARG:HB3	4.84	0.52
10:S8:76:THR:HG22	10:S8:105:ASP:HB3	2.17	0.52
11:S9:68:LYS:O	11:S9:72:GLU:HB2	2.74	0.52
36:1:1069:C:H2'	36:1:1070:U:C6	2.45	0.51
36:1:1098:A:O2'	57:N1:132:PRO:HD3	2.10	0.51
36:1:1908:A:H8	36:1:1908:A:O5'	1.92	0.51
1:2:263:C:N4	1:2:264:G:O6	2.43	0.51
36:5:1688:U:H2'	36:5:1689:U:C6	2.46	0.51
36:5:2404:A:H2'	36:5:2405:C:C5'	2.40	0.51
4:S2:159:THR:HG21	1:6:1097:U:O3'	384.89	0.51
1:6:1317:C:H2'	1:6:1318:G:O4'	2.10	0.51
1:6:447:U:C4	1:6:448:C:C4	2.98	0.51
1:6:639:U:H1'	1:6:640:U:C5	2.46	0.51
1:6:804:A:H2'	1:6:805:U:C6	2.45	0.51
1:6:920:U:H2'	1:6:921:U:O4'	2.09	0.51
1:6:995:A:H2'	1:6:996:U:O4'	2.10	0.51
86:7:220:OHX:N1	86:7:228:OHX:N5	2.58	0.51
37:7:37:G:H2'	37:7:38:U:H5''	1.92	0.51
14:C2:139:HIS:ND1	14:C2:139:HIS:O	2.43	0.51
15:C3:64:ARG:HG2	15:C3:64:ARG:HH11	3.95	0.51
19:C7:53:TYR:CZ	19:C7:57:LEU:HD21	2.45	0.51
21:C9:72:GLY:HA3	1:6:1498:G:H5''	421.56	0.51
36:1:2748:A:N3	42:L5:36:LEU:HD23	2.25	0.51
46:L9:156:GLN:HG3	46:L9:160:ASP:OD2	3.02	0.51
47:M0:48:LEU:HD11	47:M0:145:LYS:HG3	1.90	0.51
45:L8:137:ASN:HB3	51:M5:2:GLY:HA2	1.92	0.51
52:M6:158:ALA:O	52:M6:162:VAL:HG23	2.38	0.51
53:M7:87:SER:O	53:M7:90:PHE:N	2.40	0.51
54:M8:60:PRO:HG2	54:M8:142:GLY:HA3	3.20	0.51
57:N1:65:TYR:CE2	57:N1:88:ARG:HB2	2.46	0.51
67:O1:19:ARG:HD3	67:O1:35:GLU:HG2	1.92	0.51
78:Q2:12:CYS:HB3	78:Q2:17:CYS:HB3	2.21	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:S0:74:VAL:CG2	2:S0:118:PRO:HB3	2.41	0.51
7:S5:30:PRO:O	7:S5:34:GLN:N	2.36	0.51
9:S7:56:LYS:HB2	9:S7:88:ARG:HD3	1.91	0.51
9:S7:57:ALA:HA	9:S7:89:HIS:O	2.10	0.51
10:S8:62:THR:HB	10:S8:75:LYS:HE2	3.30	0.51
11:S9:47:PHE:CE1	11:S9:51:LYS:HD3	3.30	0.51
36:1:1941:C:OP2	55:M9:74:ARG:HG2	2.10	0.51
36:1:3113:A:OP1	46:L9:73:SER:OG	2.27	0.51
1:2:1125:A:C5	1:2:1126:G:H1'	2.44	0.51
1:2:1165:G:O6	1:2:1166:A:N6	2.43	0.51
1:2:1381:U:H1'	1:2:1516:A:N6	2.25	0.51
1:2:950:C:H2'	1:2:951:A:C8	2.45	0.51
36:5:1317:A:OP1	86:5:4092:OHX:N1	2.44	0.51
36:5:247:C:N3	36:5:248:U:H1'	2.25	0.51
36:5:3375:A:OP2	86:5:3951:OHX:N3	2.43	0.51
86:5:4014:OHX:N3	86:5:4212:OHX:N4	2.58	0.51
36:5:437:G:H22	36:5:622:A:H61	1.58	0.51
25:D3:14:LYS:NZ	1:6:1105:C:OP2	327.36	0.51
1:6:1244:A:O2'	1:6:1245:G:O5'	2.18	0.51
1:6:1381:U:OP1	86:6:2178:OHX:N6	2.44	0.51
13:C1:33:ARG:HH22	13:C1:52:SER:HA	1.97	0.51
18:C6:47:LYS:NZ	18:C6:114:ARG:HD3	4.34	0.51
1:2:1235:C:H2'	33:E1:138:ARG:HH21	1.75	0.51
33:E1:91:ILE:HG12	33:E1:92:LYS:H	1.74	0.51
39:L2:224:THR:HA	39:L2:237:LEU:O	2.78	0.51
40:L3:59:ASP:OD1	40:L3:71:GLU:HG2	2.60	0.51
41:L4:353:ALA:O	41:L4:357:GLU:HG3	2.10	0.51
42:L5:98:ALA:O	42:L5:162:ALA:HA	2.77	0.51
44:L7:127:LEU:C	44:L7:129:LEU:H	2.13	0.51
44:L7:237:ASN:O	44:L7:241:LYS:HB2	2.10	0.51
45:L8:53:PRO:HD2	45:L8:56:VAL:HG21	1.90	0.51
46:L9:91:ARG:HH21	46:L9:140:VAL:HG12	4.97	0.51
48:M1:16:LYS:HE3	48:M1:130:VAL:HG11	1.91	0.51
48:M1:34:SER:HA	48:M1:67:VAL:HG21	1.91	0.51
50:M4:99:TRP:CD1	50:M4:103:ILE:HD13	4.21	0.51
52:M6:182:ASN:O	52:M6:185:ALA:N	2.54	0.51
53:M7:111:LYS:HB2	53:M7:152:GLU:HG2	2.82	0.51
56:N0:155:ARG:HD3	56:N0:172:TYR:CG	4.33	0.51
56:N0:77:VAL:HG11	56:N0:106:LEU:HD22	1.92	0.51
61:N5:58:ASP:OD1	71:O5:25:LYS:NZ	3.20	0.51
63:N7:135:ARG:NH2	63:N7:135:ARG:HB3	5.01	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
63:N7:61:LYS:O	63:N7:65:ARG:N	2.60	0.51
66:O0:13:LYS:NZ	66:O0:99:ASP:OD2	2.87	0.51
68:O2:123:LYS:O	68:O2:125:ARG:N	2.43	0.51
69:O3:14:LEU:HD21	69:O3:31:LYS:HB3	2.63	0.51
72:O6:26:ILE:CD1	36:5:155:G:H1'	88.23	0.51
2:S0:157:ASP:OD2	23:D1:65:SER:OG	2.27	0.51
2:S0:55:GLU:HG2	23:D1:79:LEU:HD23	4.61	0.51
5:S3:63:GLY:O	5:S3:67:ASN:HB2	2.09	0.51
6:S4:11:ARG:O	6:S4:12:LEU:HB2	2.09	0.51
36:1:1233:G:H22	36:1:1255:C:N4	2.07	0.51
36:1:129:U:O4	86:1:3883:OHX:N5	2.44	0.51
36:1:1545:A:H2	36:1:1548:C:OP2	1.93	0.51
36:1:162:G:H1	36:1:259:C:H42	1.59	0.51
36:1:1760:A:N7	36:1:1761:C:N4	2.58	0.51
36:1:2316:G:C6	36:1:2317:A:C5	2.98	0.51
1:2:1291:G:O5'	1:2:1291:G:H8	1.93	0.51
1:2:1601:G:C5	21:C9:89:ARG:HD2	2.45	0.51
1:2:1670:G:N7	86:2:2122:OHX:N5	2.59	0.51
1:2:352:A:OP2	1:2:352:A:H8	1.92	0.51
36:5:1208:U:O2	36:5:1208:U:H2'	2.09	0.51
36:5:1366:A:C2	36:5:1367:G:C4	2.98	0.51
36:5:2286:U:C4	36:5:2288:G:H1'	2.45	0.51
36:5:2444:C:N4	36:5:2503:G:H1	2.09	0.51
36:5:2545:C:H2'	36:5:2546:C:H5'	1.93	0.51
36:5:2712:U:H2'	36:5:2713:U:C6	2.45	0.51
62:N6:10:SER:N	36:5:336:A:OP1	78.78	0.51
36:5:629:U:H2'	36:5:630:A:C8	2.45	0.51
42:L5:158:ARG:HB2	37:7:46:A:OP1	279.06	0.51
1:2:246:G:C2	13:C1:40:LEU:HD22	2.46	0.51
13:C1:93:TYR:CG	13:C1:94:ILE:N	2.78	0.51
14:C2:71:ILE:O	14:C2:75:VAL:HG23	2.09	0.51
21:C9:28:LEU:CD1	21:C9:29:GLU:H	2.24	0.51
25:D3:7:ARG:HD2	1:6:1102:G:OP2	352.23	0.51
28:D6:87:ARG:NH1	1:6:1796:C:OP1	345.18	0.51
28:D6:90:GLU:H	28:D6:90:GLU:CD	4.56	0.51
39:L2:62:VAL:HA	39:L2:73:GLU:HA	2.31	0.51
40:L3:375:GLU:O	40:L3:378:ALA:HB3	2.09	0.51
42:L5:109:THR:OG1	42:L5:110:LEU:N	2.92	0.51
46:L9:1:MET:SD	56:N0:138:GLN:HG2	2.50	0.51
46:L9:8:GLN:HG2	46:L9:68:LEU:HD13	1.92	0.51
50:M4:13:ARG:HD2	50:M4:65:LEU:O	2.86	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:M5:36:ILE:HG21	51:M5:109:ARG:HG2	1.91	0.51
53:M7:67:ILE:HG23	53:M7:68:GLY:N	2.25	0.51
61:N5:105:VAL:HG11	61:N5:126:LEU:HD13	1.92	0.51
61:N5:56:ARG:HG2	38:8:134:G:OP1	78.40	0.51
64:N8:116:GLY:HA2	64:N8:137:LYS:HZ1	1.73	0.51
67:O1:78:LYS:HB3	67:O1:79:ARG:HE	1.75	0.51
70:O4:22:VAL:HG11	36:5:1668:G:O2'	158.48	0.51
74:O8:17:ARG:HB2	74:O8:20:VAL:HG23	1.91	0.51
2:S0:172:LEU:O	2:S0:176:LEU:HG	2.11	0.51
4:S2:228:ASN:N	4:S2:228:ASN:OD1	2.78	0.51
7:S5:149:VAL:HG12	7:S5:156:ARG:HG3	5.01	0.51
7:S5:40:ILE:HG23	7:S5:42:LEU:HG	4.84	0.51
8:S6:190:GLN:OE1	8:S6:190:GLN:HA	2.97	0.51
1:2:323:A:OP2	10:S8:10:LYS:HA	2.11	0.51
10:S8:10:LYS:HG2	13:C1:133:LYS:HD2	3.31	0.51
36:1:135:C:C2	71:O5:94:LYS:HG3	2.46	0.51
36:1:1895:A:O2'	36:1:3053:G:H4'	2.10	0.51
36:1:2206:G:H8	36:1:2206:G:OP2	1.93	0.51
36:1:2221:G:N2	36:1:2223:A:H3'	2.25	0.51
36:1:3295:A:OP2	40:L3:126:LYS:N	2.42	0.51
36:1:3332:U:H2'	36:1:3333:G:O4'	2.10	0.51
1:2:1211:A:H61	1:2:1452:U:H3	1.57	0.51
1:2:1483:A:H61	1:2:1591:C:H1'	1.74	0.51
86:2:2043:OHX:N1	86:2:2098:OHX:N3	2.59	0.51
1:2:642:G:H2'	1:2:643:G:H8	1.76	0.51
1:2:912:U:H5'	1:2:913:G:C8	2.45	0.51
36:5:1168:U:O2'	36:5:1169:A:H5'	2.10	0.51
36:5:1908:A:H2'	36:5:1909:A:O4'	2.09	0.51
45:L8:241:LYS:HB2	36:5:2586:G:C5	183.90	0.51
36:5:2248:C:OP2	86:5:3971:OHX:N6	2.44	0.51
36:5:437:G:H22	36:5:622:A:N6	2.08	0.51
36:5:937:G:N3	36:5:963:G:H1'	2.25	0.51
35:SM:68:ARG:HG2	1:6:1460:A:OP1	336.01	0.51
1:6:195:G:H2'	1:6:196:G:H5'	1.93	0.51
1:6:274:G:H2'	1:6:275:C:H6	1.75	0.51
56:N0:43:TYR:OH	37:7:96:U:OP1	294.26	0.51
38:8:157:U:H2'	38:8:158:U:C6	2.46	0.51
15:C3:115:LEU:HD23	15:C3:115:LEU:O	4.37	0.51
5:S3:210:GLU:OE1	19:C7:19:ARG:HD3	5.04	0.51
28:D6:75:VAL:O	28:D6:79:ILE:N	2.40	0.51
30:D8:49:ARG:HG2	30:D8:52:ASP:OD1	2.56	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:E1:141:CYS:SG	33:E1:143:LYS:HB3	3.02	0.51
42:L5:146:LEU:HD13	42:L5:148:ILE:HD13	5.22	0.51
42:L5:155:THR:HG23	37:7:36:C:H5'	269.33	0.51
44:L7:132:PRO:HA	44:L7:229:PHE:CE2	3.00	0.51
48:M1:133:ARG:HD2	48:M1:152:HIS:O	2.11	0.51
48:M1:10:ARG:HH21	48:M1:152:HIS:H	4.58	0.51
48:M1:60:ARG:O	48:M1:63:GLU:HB2	2.11	0.51
49:M3:58:VAL:HG12	49:M3:70:ARG:O	2.10	0.51
50:M4:24:LYS:HG3	50:M4:25:LYS:HD3	1.91	0.51
52:M6:46:GLU:HG2	52:M6:48:PHE:H	1.75	0.51
55:M9:20:ARG:HD2	36:5:1874:A:OP2	141.88	0.51
60:N4:50:ALA:HA	60:N4:55:PHE:CG	2.75	0.51
62:N6:120:GLN:NE2	62:N6:126:LEU:HA	7.15	0.51
67:O1:13:THR:HB	67:O1:72:ARG:HD3	2.23	0.51
67:O1:52:ALA:HB3	67:O1:55:LEU:HB2	2.21	0.51
68:O2:82:LEU:HD11	68:O2:112:ALA:HB2	1.93	0.51
68:O2:47:ARG:NH1	36:5:634:C:H4'	218.47	0.51
70:O4:41:ARG:NH1	70:O4:50:ALA:HB1	4.77	0.51
78:Q2:99:GLN:HG2	78:Q2:100:LYS:N	2.25	0.51
6:S4:104:ASP:HB2	6:S4:108:ARG:O	2.52	0.51
1:2:178:U:OP1	8:S6:191:ARG:NH2	2.44	0.51
9:S7:44:LYS:HD2	9:S7:63:PRO:HA	3.46	0.51
34:SR:126:SER:OG	34:SR:127:ARG:N	2.42	0.51
34:SR:12:THR:HG22	34:SR:311:ARG:HG2	1.91	0.51
36:1:1114:U:OP2	86:1:3957:OHX:N4	2.44	0.51
36:1:1155:C:H2'	36:1:1156:C:C6	2.45	0.51
36:1:1162:U:H4'	68:O2:57:TYR:CD1	2.45	0.51
36:1:1233:G:H22	36:1:1255:C:H42	1.58	0.51
36:1:1317:A:O2'	36:1:1318:A:H3'	2.10	0.51
36:1:1524:A:OP1	61:N5:92:LYS:NZ	2.43	0.51
36:1:1740:U:H1'	36:1:1741:A:C2	2.42	0.51
36:1:2529:A:H2'	36:1:2530:G:O4'	2.10	0.51
36:1:497:C:O3'	69:O3:86:ARG:NH2	2.44	0.51
36:1:748:U:H2'	36:1:749:C:H6	1.76	0.51
36:1:841:A:OP2	86:1:4172:OHX:N2	2.43	0.51
36:1:898:U:H2'	36:1:899:U:O4'	2.11	0.51
1:2:1157:A:H61	1:2:1621:U:H3	1.59	0.51
1:2:67:A:N6	1:2:83:G:O2'	2.43	0.51
37:3:68:C:OP1	42:L5:14:SER:OG	2.28	0.51
36:5:1014:U:H2'	36:5:1015:U:H5'	1.93	0.51
36:5:1152:G:H22	36:5:1200:A:N6	2.07	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:1238:C:H2'	36:5:1239:C:C6	2.45	0.51
41:L4:197:ARG:NH1	36:5:1381:A:OP1	108.95	0.51
36:5:1579:C:H2'	36:5:1580:A:H8	1.75	0.51
36:5:128:G:O6	86:5:3925:OHX:N4	2.43	0.51
1:6:717:C:O2'	1:6:718:U:OP1	2.22	0.51
37:7:3:U:H2'	37:7:4:U:C6	2.46	0.51
12:C0:72:GLY:O	12:C0:75:TYR:N	2.51	0.51
14:C2:28:LEU:HD11	14:C2:89:ILE:HG21	2.15	0.51
20:C8:17:LEU:HD12	20:C8:18:LEU:HD23	1.93	0.51
23:D1:32:VAL:HB	23:D1:60:ARG:HD2	3.19	0.51
24:D2:2:THR:OG1	24:D2:3:ARG:N	2.42	0.51
33:E1:102:VAL:O	33:E1:104:SER:N	2.73	0.51
39:L2:171:GLY:O	79:Q3:68:ALA:HB2	2.61	0.51
40:L3:193:ASP:O	40:L3:197:GLU:HG3	2.10	0.51
42:L5:148:ILE:HG23	42:L5:151:GLN:HB3	4.88	0.51
44:L7:90:LYS:HG2	44:L7:95:ILE:HD11	2.76	0.51
45:L8:63:LYS:O	45:L8:67:ILE:HG13	2.11	0.51
48:M1:57:PHE:N	48:M1:57:PHE:CD1	2.90	0.51
49:M3:54:LEU:HD12	49:M3:75:PHE:CE2	2.45	0.51
52:M6:20:ALA:O	52:M6:24:ALA:N	2.37	0.51
54:M8:158:HIS:H	54:M8:186:VAL:HG11	1.91	0.51
56:N0:30:PHE:CD2	56:N0:103:VAL:HG21	2.45	0.51
44:L7:80:GLN:HG3	57:N1:136:ARG:HB2	4.23	0.51
59:N3:17:LEU:O	59:N3:52:ALA:N	2.32	0.51
59:N3:24:ASN:O	59:N3:99:ALA:HA	2.30	0.51
36:1:426:G:H5'	68:O2:50:ILE:HG22	1.92	0.51
70:O4:63:ALA:HB2	36:5:1803:C:H5'	157.88	0.51
73:O7:37:CYS:O	73:O7:45:ARG:N	2.35	0.51
4:S2:44:LEU:HD11	4:S2:243:TYR:HB2	3.24	0.51
36:1:109:A:O2'	36:1:323:A:N6	2.44	0.51
36:1:139:G:H2'	36:1:140:C:O4'	2.11	0.51
36:1:1445:U:H5''	36:1:1446:A:OP2	2.11	0.51
36:1:2558:U:O2'	36:1:2559:U:H5'	2.10	0.51
36:1:3027:A:H2'	36:1:3028:G:O4'	2.10	0.51
36:1:955:U:H2'	36:1:956:U:C6	2.45	0.51
1:2:1065:A:N3	3:S1:146:GLN:NE2	2.56	0.51
52:M6:130:LYS:HA	36:5:1316:C:C5	296.36	0.51
36:5:1471:U:H2'	36:5:1472:U:C6	2.44	0.51
36:5:2359:C:H2'	36:5:2360:C:H6	1.76	0.51
36:5:345:G:H1'	38:8:24:G:H22	1.75	0.51
36:5:754:G:H2'	36:5:755:A:H8	1.76	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:1603:U:H2'	1:6:1604:U:H6	1.74	0.51
1:6:938:G:N7	86:6:2105:OHX:N3	2.59	0.51
86:6:2120:OHX:N4	86:6:2171:OHX:N3	2.59	0.51
1:6:509:G:H2'	1:6:510:G:O4'	2.10	0.51
79:Q3:24:ARG:NH2	1:6:982:U:O3'	249.32	0.51
12:C0:6:GLU:OE1	12:C0:10:LYS:HD2	2.11	0.51
16:C4:25:ASP:OD1	16:C4:26:THR:N	2.84	0.51
23:D1:85:TYR:CD1	29:D7:6:ASP:HB2	2.86	0.51
39:L2:117:GLU:OE1	39:L2:163:ARG:NH2	2.40	0.51
40:L3:130:PHE:CE1	36:5:3149:G:H4'	221.44	0.51
41:L4:93:MET:HB2	36:5:658:G:H21	145.14	0.51
42:L5:211:LEU:O	42:L5:214:ASP:N	3.49	0.51
45:L8:45:ASN:ND2	45:L8:47:SER:H	2.07	0.51
47:M0:177:ASP:OD2	47:M0:177:ASP:N	3.51	0.51
48:M1:57:PHE:N	48:M1:57:PHE:HD1	2.40	0.51
50:M4:32:LEU:HD11	50:M4:94:TRP:CD1	2.46	0.51
51:M5:106:VAL:O	51:M5:109:ARG:HB3	2.09	0.51
52:M6:10:ASP:HA	52:M6:36:VAL:HG23	1.92	0.51
52:M6:141:LEU:O	52:M6:144:SER:HB3	3.79	0.51
52:M6:62:THR:HB	52:M6:65:ASN:O	2.10	0.51
52:M6:73:PHE:HB3	52:M6:78:ARG:HG2	1.92	0.51
54:M8:16:ARG:HH12	54:M8:55:SER:HB3	1.73	0.51
62:N6:37:LYS:HG2	62:N6:38:GLU:H	1.74	0.51
62:N6:52:ARG:HG2	62:N6:53:ASP:N	3.14	0.51
68:O2:123:LYS:HA	68:O2:126:LEU:HG	1.91	0.51
68:O2:50:ILE:HG13	68:O2:50:ILE:O	2.36	0.51
69:O3:26:ASN:O	69:O3:84:THR:HG22	2.11	0.51
78:Q2:47:GLN:NE2	78:Q2:53:GLN:HA	2.37	0.51
36:1:860:G:H5''	79:Q3:17:ARG:HH12	1.76	0.51
79:Q3:73:THR:HB	79:Q3:76:ALA:H	4.96	0.51
3:S1:140:ILE:O	3:S1:210:ILE:HA	2.11	0.51
4:S2:89:GLN:OE1	4:S2:94:GLN:HG2	2.10	0.51
1:2:753:A:H5'	6:S4:221:ARG:HG3	1.93	0.51
10:S8:12:SER:HB3	10:S8:16:ALA:HB3	2.72	0.51
10:S8:31:ARG:HH21	10:S8:48:THR:HG22	2.52	0.51
34:SR:135:THR:HG23	34:SR:139:GLN:O	5.27	0.51
34:SR:185:GLN:HB3	34:SR:187:GLN:HE22	1.75	0.51
36:1:1949:G:OP1	55:M9:104:ARG:NH1	2.43	0.51
36:1:2555:G:N3	70:O4:92:ALA:HA	2.26	0.51
86:1:3965:OHX:N3	86:1:4153:OHX:N4	2.59	0.51
36:1:3087:A:OP1	86:1:4178:OHX:N5	2.44	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:607:A:H4'	36:1:608:A:OP2	2.09	0.51
1:2:1076:A:H2'	1:2:1077:C:C6	2.46	0.51
1:2:1575:G:H2'	1:2:1576:A:C8	2.46	0.51
1:2:1643:U:H2'	1:2:1644:C:O4'	2.11	0.51
1:2:285:G:H2'	1:2:286:C:C6	2.46	0.51
36:5:982:C:H42	36:5:1101:G:H1	1.58	0.51
36:5:172:G:H2'	36:5:172:G:N3	2.26	0.51
36:5:2407:C:H1'	36:5:2818:U:O2	2.11	0.51
36:5:2407:C:H1'	36:5:2818:U:C2	2.45	0.51
36:5:2992:U:H2'	36:5:2993:G:O4'	2.11	0.51
36:5:549:U:H2'	36:5:550:A:C8	2.46	0.51
64:N8:16:SER:HA	36:5:942:U:N3	169.54	0.51
1:6:1180:C:H2'	1:6:1181:U:O4'	2.10	0.51
1:6:1274:C:H4'	1:6:1275:A:O5'	2.10	0.51
1:6:1491:U:H5'	1:6:1492:A:OP1	2.11	0.51
1:6:1647:U:H2'	1:6:1648:A:C8	2.44	0.51
1:6:886:U:H2'	1:6:887:A:C8	2.43	0.51
16:C4:111:ARG:HA	28:D6:56:ALA:O	2.22	0.51
17:C5:110:GLU:HG2	20:C8:119:ILE:HD13	4.65	0.51
1:2:1565:C:OP1	20:C8:41:ARG:HG3	2.10	0.51
25:D3:52:ILE:O	25:D3:74:VAL:HA	2.10	0.51
26:D4:20:ARG:HD3	26:D4:76:TYR:CE2	2.96	0.51
40:L3:152:LYS:HG3	40:L3:192:VAL:HG11	3.36	0.51
41:L4:264:SER:C	41:L4:266:THR:N	2.62	0.51
41:L4:300:ARG:O	41:L4:302:ALA:N	3.76	0.51
42:L5:131:LEU:HD22	42:L5:131:LEU:H	1.76	0.51
45:L8:204:ARG:HB3	45:L8:206:GLU:HG3	1.92	0.51
46:L9:20:ILE:HG23	46:L9:25:VAL:HG13	1.93	0.51
48:M1:60:ARG:HB2	48:M1:60:ARG:HH21	5.13	0.51
53:M7:41:LEU:O	53:M7:41:LEU:HD22	2.11	0.51
55:M9:105:LEU:HD11	55:M9:139:VAL:HG23	1.93	0.51
55:M9:109:TYR:C	55:M9:115:ILE:HG22	2.31	0.51
55:M9:39:ASN:O	55:M9:43:LYS:HG2	3.83	0.51
55:M9:9:ARG:HH11	55:M9:9:ARG:HG3	1.76	0.51
59:N3:120:LYS:H	59:N3:137:VAL:HG22	3.08	0.51
62:N6:36:SER:OG	62:N6:38:GLU:HB3	2.11	0.51
62:N6:37:LYS:CD	62:N6:37:LYS:H	2.93	0.51
64:N8:47:LYS:O	64:N8:49:HIS:N	3.39	0.51
65:N9:18:ARG:CZ	65:N9:18:ARG:HB3	2.41	0.51
68:O2:109:LEU:HD22	68:O2:119:VAL:HG21	2.45	0.51
70:O4:46:ASP:HB3	70:O4:84:CYS:SG	2.50	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
72:O6:52:PRO:O	72:O6:55:ARG:HB2	2.77	0.51
73:O7:63:ARG:NH2	38:8:58:G:O6	78.94	0.51
73:O7:64:MET:O	73:O7:68:LYS:HG3	2.11	0.51
2:S0:140:ASN:ND2	23:D1:29:HIS:HA	2.26	0.51
3:S1:170:GLU:O	3:S1:174:LYS:HG3	2.11	0.51
5:S3:194:LYS:O	5:S3:196:ARG:N	3.03	0.51
6:S4:131:LEU:HD13	6:S4:135:GLY:HA2	2.43	0.51
1:2:1611:A:O2'	7:S5:95:ASN:O	2.26	0.51
11:S9:77:ILE:HG23	11:S9:86:LEU:HD23	3.88	0.51
36:1:2539:C:H5'	36:1:2541:U:O4	2.10	0.51
36:1:2534:G:N2	36:1:2545:C:N3	2.45	0.51
36:1:3259:U:C6	36:1:3259:U:H5'	2.41	0.51
36:1:535:G:C6	36:1:555:U:N3	2.79	0.51
1:2:45:U:O2'	1:2:46:A:H2'	2.11	0.51
1:2:623:A:OP2	86:2:2157:OHX:N4	2.44	0.51
36:5:1716:U:O2'	36:5:1717:U:H4'	2.11	0.51
36:5:1952:G:O6	36:5:2094:C:N4	2.44	0.51
36:5:2694:A:C6	36:5:2695:A:C6	2.99	0.51
36:5:2947:G:OP2	36:5:2947:G:H4'	2.09	0.51
36:5:421:G:O6	36:5:2383:C:O2'	2.15	0.51
36:5:637:C:C2	36:5:638:C:C5	2.98	0.51
36:5:945:C:H2'	36:5:946:U:C6	2.46	0.51
1:6:1124:A:H2'	1:6:1125:A:C8	2.46	0.51
1:6:1333:C:O2	1:6:1419:G:C2	2.64	0.51
1:6:643:G:H1	1:6:691:C:H42	1.59	0.51
1:6:737:A:H2'	1:6:738:G:H8	1.75	0.51
13:C1:14:GLN:HB3	13:C1:54:ILE:HG13	3.63	0.51
15:C3:105:ASN:ND2	15:C3:105:ASN:H	2.08	0.51
18:C6:120:ASP:OD1	18:C6:122:ARG:HG3	2.90	0.51
18:C6:47:LYS:HE2	18:C6:114:ARG:CZ	2.41	0.51
22:D0:24:ILE:O	22:D0:90:TYR:HA	2.11	0.51
29:D7:64:CYS:HB3	29:D7:73:LEU:HD12	4.78	0.51
39:L2:227:ARG:HG2	39:L2:239:ALA:CB	2.38	0.51
42:L5:190:ILE:HG13	42:L5:191:ASP:N	2.24	0.51
42:L5:198:TYR:CE1	42:L5:203:HIS:CD2	2.99	0.51
42:L5:23:ARG:O	42:L5:23:ARG:HD2	4.03	0.51
45:L8:101:THR:N	45:L8:104:GLU:OE2	2.43	0.51
50:M4:99:TRP:HE1	50:M4:103:ILE:HD13	3.42	0.51
50:M4:25:LYS:HD3	50:M4:62:GLN:HB3	1.91	0.51
57:N1:109:VAL:O	57:N1:112:ASN:N	2.44	0.51
58:N2:59:ASP:N	58:N2:62:VAL:O	2.40	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
63:N7:9:LYS:HD2	63:N7:83:THR:O	2.10	0.51
64:N8:25:HIS:C	64:N8:25:HIS:CD2	3.81	0.51
63:N7:3:LYS:HE3	66:O0:36:GLN:HA	1.92	0.51
67:O1:54:GLU:HA	67:O1:57:GLN:OE1	4.61	0.51
68:O2:99:ASN:N	68:O2:99:ASN:OD1	2.43	0.51
3:S1:144:ARG:HB3	3:S1:208:GLN:HB3	1.93	0.51
3:S1:175:GLU:O	3:S1:187:LYS:NZ	2.43	0.51
3:S1:32:ILE:HB	3:S1:44:GLY:H	1.74	0.51
3:S1:62:LYS:HD2	3:S1:91:VAL:HG11	1.92	0.51
8:S6:25:ARG:HG3	8:S6:28:PHE:HD1	1.76	0.51
36:1:1240:A:N6	36:1:1244:A:H5''	2.16	0.51
36:1:1573:G:N3	36:1:1573:G:H2'	2.26	0.51
36:1:2526:C:O2'	45:L8:241:LYS:NZ	2.41	0.51
36:1:944:C:H4'	68:O2:33:ARG:NH1	2.26	0.51
1:2:1234:A:O2'	1:2:1235:C:O5'	2.28	0.51
1:2:792:U:H2'	1:2:793:A:H5'	1.91	0.51
36:5:1521:G:C2	36:5:1522:U:H5	2.29	0.51
36:5:2317:A:OP2	86:5:4182:OHX:N4	2.43	0.51
36:5:3296:A:H2'	36:5:3297:U:H6	1.76	0.51
36:5:2400:G:OP1	86:5:4105:OHX:N1	2.44	0.51
38:8:113:U:HO2'	38:8:114:G:P	2.31	0.51
15:C3:12:SER:HB3	1:6:956:C:O5'	337.13	0.51
18:C6:127:LYS:HA	18:C6:134:ALA:HA	1.92	0.51
20:C8:42:TYR:HA	20:C8:85:PHE:HE1	1.75	0.51
22:D0:25:THR:HG23	22:D0:88:LYS:HD3	1.93	0.51
7:S5:61:TYR:OH	30:D8:52:ASP:OD1	3.95	0.51
40:L3:81:THR:HG23	40:L3:81:THR:O	3.98	0.51
41:L4:182:LEU:O	41:L4:184:SER:N	2.42	0.51
41:L4:99:MET:SD	41:L4:102:PRO:HA	2.51	0.51
44:L7:209:ASN:HD22	36:5:1168:U:H1'	243.44	0.51
44:L7:214:TRP:CD2	44:L7:219:LYS:HD2	2.46	0.51
44:L7:92:ILE:HD12	44:L7:92:ILE:O	2.11	0.51
45:L8:91:PHE:HE1	45:L8:185:ARG:HD2	1.75	0.51
47:M0:77:THR:O	47:M0:81:GLY:N	2.41	0.51
47:M0:97:LEU:O	47:M0:123:HIS:N	2.42	0.51
52:M6:189:ASP:O	52:M6:193:GLN:HG3	2.16	0.51
54:M8:53:PHE:N	54:M8:53:PHE:CD1	2.85	0.51
59:N3:33:ASN:ND2	59:N3:64:LYS:H	2.09	0.51
59:N3:80:ARG:HB2	59:N3:99:ALA:HB3	1.93	0.51
70:O4:25:THR:OG1	70:O4:27:GLY:N	3.34	0.51
71:O5:94:LYS:O	71:O5:98:SER:OG	3.36	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:S8:69:SER:OG	10:S8:185:GLU:OE2	2.77	0.51
34:SR:79:TYR:HE1	34:SR:100:TYR:HE1	3.48	0.51
34:SR:112:SER:OG	34:SR:153:GLN:HA	2.11	0.51
36:1:1075:A:C5	65:N9:45:HIS:CD2	2.99	0.51
36:1:1712:G:N2	36:1:1731:A:OP2	2.35	0.51
36:1:2294:U:OP1	59:N3:70:ARG:NH2	2.42	0.51
36:1:2307:G:O2'	36:1:2310:U:OP2	2.28	0.51
36:1:3199:G:C2	36:1:3200:G:C8	2.99	0.51
36:1:1752:A:OP2	86:1:4042:OHX:N3	2.43	0.51
1:2:1428:G:H8	1:2:1428:G:H5'	1.76	0.51
1:2:396:G:H22	1:2:399:A:C5'	2.24	0.51
38:4:53:A:H2'	38:4:54:A:C8	2.46	0.51
36:5:1077:U:H2'	36:5:1078:U:H6	1.76	0.51
36:5:1176:C:H2'	36:5:1177:G:N2	2.26	0.51
36:5:1597:C:H42	36:5:1610:G:H1	1.57	0.51
36:5:2768:U:H2'	36:5:2769:A:H8	1.76	0.51
52:M6:115:LYS:HG2	36:5:3178:A:C2	259.42	0.51
36:5:2942:C:O2	86:5:4104:OHX:N2	2.44	0.51
1:6:1058:U:H4'	1:6:1059:U:OP1	2.09	0.51
1:6:1140:G:OP2	86:6:2071:OHX:N3	2.44	0.51
17:C5:122:THR:HB	1:6:1558:U:N3	366.49	0.51
38:8:79:A:H2'	38:8:80:A:O4'	2.11	0.51
1:2:861:U:O3'	15:C3:20:ARG:NH2	2.42	0.51
19:C7:109:LEU:O	19:C7:113:LEU:HB2	5.16	0.51
22:D0:104:THR:HG22	22:D0:116:VAL:HG21	1.93	0.51
24:D2:22:LYS:HG3	29:D7:3:LEU:HA	1.93	0.51
40:L3:153:LYS:HD3	40:L3:154:TYR:CZ	2.45	0.51
41:L4:131:VAL:O	41:L4:135:VAL:HG23	3.24	0.51
42:L5:269:SER:OG	42:L5:270:LYS:N	4.43	0.51
42:L5:279:LYS:HE3	42:L5:282:ARG:HD2	1.93	0.51
44:L7:157:ASN:O	44:L7:159:GLN:HG2	2.11	0.51
48:M1:16:LYS:HG3	48:M1:130:VAL:HG13	4.25	0.51
50:M4:17:VAL:HG22	50:M4:36:VAL:O	2.11	0.51
51:M5:11:GLN:O	51:M5:11:GLN:HG3	2.11	0.51
56:N0:14:LEU:HG	56:N0:55:SER:O	2.10	0.51
41:L4:354:VAL:HG11	57:N1:143:THR:HG21	1.93	0.51
61:N5:100:LYS:HE3	61:N5:106:ASP:OD2	2.18	0.51
61:N5:31:THR:HB	61:N5:33:ARG:NH2	2.26	0.51
62:N6:63:LYS:HE3	62:N6:97:ILE:HD13	1.93	0.51
72:O6:90:MET:O	72:O6:93:ILE:N	2.44	0.51
2:S0:7:PHE:HZ	23:D1:43:GLY:HA2	1.97	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:S5:208:SER:O	7:S5:211:ILE:N	3.80	0.51
7:S5:43:PHE:CG	7:S5:44:ASN:N	2.90	0.51
8:S6:10:ASN:HB3	8:S6:128:THR:HA	2.05	0.51
9:S7:35:LYS:HG2	9:S7:36:ALA:H	1.75	0.51
34:SR:264:SER:HB3	34:SR:269:TYR:CE1	2.46	0.51
34:SR:49:GLY:N	34:SR:54:PHE:O	2.43	0.51
36:1:1236:G:N2	36:1:1272:C:OP1	2.40	0.50
36:1:1412:G:OP1	68:O2:105:ARG:NH2	2.44	0.50
36:1:1586:G:OP1	86:1:3935:OHX:N5	2.44	0.50
36:1:181:U:O3'	73:O7:75:LYS:HD3	2.10	0.50
36:1:224:C:H2'	36:1:225:C:H6	1.76	0.50
36:1:2503:G:H1'	36:1:2504:U:C5	2.46	0.50
36:1:3174:A:C6	36:1:3175:U:N3	2.79	0.50
36:1:3206:C:H5''	36:1:3207:U:O5'	2.11	0.50
36:1:3348:G:H1	36:1:3357:U:H3	1.58	0.50
36:1:345:G:OP1	36:1:1429:G:N2	2.42	0.50
1:2:74:U:H1'	1:2:75:U:H5''	1.92	0.50
36:5:2936:A:H2'	36:5:2937:G:C8	2.47	0.50
1:6:425:A:H8	1:6:425:A:H5'	1.76	0.50
1:6:763:G:C6	1:6:764:U:C4	2.99	0.50
15:C3:94:LYS:HG2	15:C3:118:ILE:HD13	2.17	0.50
19:C7:6:THR:HG23	19:C7:9:VAL:HG23	1.93	0.50
21:C9:135:ILE:HA	21:C9:138:GLN:HG3	2.30	0.50
22:D0:37:VAL:O	22:D0:41:ILE:HD13	2.11	0.50
40:L3:247:ARG:NH2	36:5:2341:A:OP1	219.03	0.50
41:L4:138:ARG:NH1	41:L4:138:ARG:O	2.44	0.50
41:L4:361:HIS:CD2	41:L4:362:ASP:HB2	4.04	0.50
43:L6:40:LEU:HB3	43:L6:84:VAL:CG1	2.52	0.50
41:L4:327:LEU:HD21	44:L7:164:SER:HA	1.92	0.50
46:L9:91:ARG:NH2	46:L9:140:VAL:HG12	4.64	0.50
46:L9:150:SER:HB3	46:L9:153:ASP:HB2	1.93	0.50
47:M0:42:THR:HG23	47:M0:45:GLU:H	1.76	0.50
50:M4:14:LEU:H	50:M4:19:ARG:NH1	2.22	0.50
50:M4:22:LEU:HB3	50:M4:64:VAL:HG13	3.25	0.50
50:M4:99:TRP:NE1	50:M4:103:ILE:HD13	3.92	0.50
51:M5:191:TRP:O	51:M5:194:GLN:N	2.44	0.50
56:N0:68:HIS:O	56:N0:73:LYS:NZ	2.41	0.50
57:N1:14:MET:SD	57:N1:58:GLN:HG2	2.81	0.50
59:N3:36:ILE:HG23	59:N3:58:VAL:HG21	1.92	0.50
79:Q3:33:GLN:HG2	79:Q3:34:HIS:N	2.26	0.50
6:S4:16:HIS:C	6:S4:18:TRP:H	2.14	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:S5:51:VAL:O	7:S5:65:ARG:NH1	3.84	0.50
10:S8:10:LYS:HG2	13:C1:133:LYS:HE3	1.93	0.50
34:SR:143:THR:HG22	34:SR:145:LEU:HD21	1.93	0.50
36:1:593:C:C4	36:1:594:U:C5	2.99	0.50
36:1:703:G:C6	36:1:704:U:C4	3.00	0.50
1:2:1335:U:P	22:D0:85:ARG:HH12	2.34	0.50
1:2:1748:G:O6	86:2:2104:OHX:N4	2.44	0.50
37:3:76:A:OP2	86:3:216:OHX:N5	2.44	0.50
36:5:65:A:C4	36:5:110:G:N7	2.79	0.50
36:5:1602:A:H2'	36:5:1603:A:C8	2.47	0.50
36:5:1700:G:H1	36:5:1745:C:N4	2.09	0.50
36:5:1792:C:H5''	36:5:1793:C:P	2.51	0.50
39:L2:244:GLY:HA2	36:5:2243:A:H3'	233.67	0.50
35:SM:31:SER:OG	36:5:2667:A:OP1	288.59	0.50
1:6:12:U:H1'	1:6:1300:A:N3	2.27	0.50
1:6:1333:C:O2	1:6:1419:G:N2	2.45	0.50
1:6:1427:A:O2'	1:6:1428:G:OP1	2.24	0.50
1:6:1733:C:H2'	1:6:1734:U:C6	2.47	0.50
16:C4:127:ARG:HG3	28:D6:22:ARG:HH22	1.77	0.50
31:D9:22:ARG:HH21	31:D9:37:ASN:HD22	3.82	0.50
39:L2:20:THR:HA	39:L2:23:ARG:HD2	1.93	0.50
40:L3:43:LEU:HB2	40:L3:208:VAL:HG11	1.93	0.50
40:L3:75:ALA:HB2	36:5:3049:A:C2	246.26	0.50
41:L4:328:ASN:OD1	41:L4:330:TYR:HB3	2.42	0.50
36:1:2687:G:P	42:L5:8:LYS:HZ1	2.32	0.50
43:L6:172:HIS:CD2	43:L6:173:MET:HG2	2.46	0.50
43:L6:64:LEU:O	43:L6:65:ILE:HD13	5.49	0.50
44:L7:186:HIS:CE1	44:L7:190:THR:HG21	3.31	0.50
44:L7:25:GLN:HA	44:L7:29:GLU:H	1.76	0.50
44:L7:80:GLN:HB2	57:N1:135:PRO:HB2	1.92	0.50
49:M3:110:ASP:HA	49:M3:113:VAL:HG23	1.92	0.50
50:M4:48:GLY:HA3	50:M4:53:VAL:HG13	2.28	0.50
51:M5:140:LYS:O	51:M5:142:ILE:N	2.45	0.50
51:M5:28:TRP:O	51:M5:32:GLN:HG2	2.10	0.50
52:M6:72:HIS:O	52:M6:74:ARG:HD3	2.51	0.50
36:1:1313:G:OP1	52:M6:82:LYS:HD3	2.11	0.50
53:M7:48:LEU:O	53:M7:52:LEU:HD22	2.87	0.50
54:M8:115:VAL:O	54:M8:118:GLY:N	2.39	0.50
60:N4:52:THR:O	60:N4:56:ARG:HG3	2.11	0.50
62:N6:35:LEU:HA	62:N6:106:ILE:HB	1.93	0.50
66:O0:95:ALA:HB2	66:O0:101:LEU:HD21	4.34	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
68:O2:105:ARG:O	68:O2:109:LEU:HB2	2.41	0.50
36:1:135:C:N3	71:O5:94:LYS:HG3	2.26	0.50
72:O6:26:ILE:C	72:O6:28:TYR:N	2.63	0.50
73:O7:17:THR:HG22	73:O7:18:LEU:H	1.76	0.50
79:Q3:17:ARG:HB3	79:Q3:18:TYR:CD1	3.19	0.50
3:S1:23:PRO:HG2	1:6:896:U:H5'	254.72	0.50
5:S3:202:LEU:O	5:S3:204:ASP:N	2.99	0.50
6:S4:22:LYS:O	6:S4:23:LEU:HD13	3.61	0.50
34:SR:157:VAL:HB	34:SR:168:THR:HG22	4.28	0.50
34:SR:5:GLU:OE1	34:SR:249:ARG:NE	4.96	0.50
36:1:2146:C:O2'	36:1:2147:A:H5'	2.11	0.50
36:1:2278:C:C2	36:1:2307:G:N2	2.79	0.50
36:1:2422:C:O5'	78:Q2:52:GLY:HA2	2.10	0.50
36:1:2592:G:H4'	36:1:2594:C:C2	2.47	0.50
36:1:3134:A:OP1	86:1:3894:OHX:N4	2.44	0.50
36:1:529:A:N6	36:1:563:U:H3	2.06	0.50
36:1:945:C:H2'	36:1:946:U:C6	2.46	0.50
1:2:1474:G:H2'	1:2:1475:A:C8	2.46	0.50
1:2:1537:C:O2'	1:2:1540:G:O6	2.25	0.50
36:5:1195:A:H1'	36:5:1319:G:H4'	1.92	0.50
36:5:2427:U:H2'	36:5:2428:U:C6	2.47	0.50
36:5:3055:U:H1'	36:5:3057:U:OP2	2.12	0.50
36:5:787:G:H2'	36:5:788:C:C6	2.46	0.50
1:6:1188:G:O2'	1:6:1430:U:OP1	2.28	0.50
27:D5:74:SER:OG	1:6:1534:G:OP2	344.99	0.50
1:6:1699:G:H22	1:6:1701:A:H3'	1.76	0.50
1:6:328:A:H2'	1:6:329:G:H8	1.76	0.50
1:6:990:C:H2'	1:6:991:G:O4'	2.12	0.50
13:C1:75:VAL:CG1	13:C1:119:VAL:HA	2.41	0.50
15:C3:54:LEU:HB3	15:C3:60:VAL:CG1	3.68	0.50
18:C6:38:LEU:O	18:C6:40:GLU:N	2.45	0.50
24:D2:102:VAL:O	24:D2:113:HIS:HB3	2.90	0.50
32:E0:28:LYS:HZ1	1:6:542:A:H61	428.90	0.50
40:L3:86:VAL:HB	40:L3:198:HIS:O	2.11	0.50
8:S6:22:HIS:CD2	40:L3:300:ARG:HH22	2.30	0.50
40:L3:77:THR:OG1	40:L3:324:VAL:HG12	2.11	0.50
42:L5:61:ILE:HG12	42:L5:79:TYR:HD1	1.75	0.50
44:L7:191:VAL:HG12	44:L7:192:GLY:H	3.64	0.50
44:L7:224:ILE:HG23	56:N0:36:ILE:HG12	2.31	0.50
45:L8:101:THR:CG2	45:L8:104:GLU:H	2.22	0.50
49:M3:46:ILE:HG12	49:M3:49:ARG:CZ	3.42	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:M5:79:ALA:HB1	51:M5:81:TYR:CZ	2.46	0.50
52:M6:24:ALA:HB1	52:M6:88:VAL:HG23	2.88	0.50
54:M8:170:ARG:O	54:M8:171:LYS:HB2	3.83	0.50
59:N3:133:SER:O	86:6:2117:OHX:N3	295.78	0.50
70:O4:104:VAL:HA	70:O4:107:GLU:HB2	1.93	0.50
70:O4:66:SER:O	70:O4:70:LYS:HE3	3.92	0.50
51:M5:6:TYR:CD2	72:O6:40:VAL:HG13	2.78	0.50
74:O8:30:LYS:HB2	74:O8:38:PHE:CE2	2.46	0.50
6:S4:181:VAL:HG11	6:S4:225:VAL:HG13	2.93	0.50
6:S4:246:LEU:HD21	6:S4:254:ARG:CZ	2.41	0.50
6:S4:9:LEU:CB	6:S4:30:ARG:HB2	3.51	0.50
1:2:395:U:O2'	8:S6:89:ASP:HB3	2.11	0.50
9:S7:126:LEU:HB2	9:S7:173:TYR:CE2	6.37	0.50
9:S7:63:PRO:O	9:S7:64:VAL:HB	3.36	0.50
9:S7:61:PHE:HE1	9:S7:93:LEU:HD12	2.62	0.50
10:S8:44:HIS:O	10:S8:56:ARG:N	2.90	0.50
34:SR:289:ALA:HB2	34:SR:305:TYR:CE2	2.65	0.50
36:1:123:A:C6	36:1:150:A:C5	3.00	0.50
36:1:1904:C:N4	36:1:1905:G:O6	2.44	0.50
86:1:3998:OHX:N3	86:1:4169:OHX:N1	2.60	0.50
1:2:936:G:N7	28:D6:15:ARG:NH1	2.49	0.50
38:4:124:G:H3'	38:4:125:U:C5'	2.37	0.50
36:5:2385:G:O6	86:5:3926:OHX:N4	2.45	0.50
36:5:240:U:O2'	36:5:241:G:H8	1.94	0.50
36:5:3257:C:H2'	36:5:3258:U:O4'	2.12	0.50
36:5:980:A:H2'	36:5:981:U:N1	2.27	0.50
1:6:1649:G:N7	86:6:2109:OHX:N2	2.58	0.50
1:6:1239:U:O4	86:6:2096:OHX:N1	2.44	0.50
1:6:139:C:C4	1:6:266:A:C2	3.00	0.50
86:7:220:OHX:N1	86:7:228:OHX:N2	2.59	0.50
13:C1:111:VAL:HB	13:C1:139:VAL:HG21	1.92	0.50
2:S0:185:ARG:HH21	23:D1:47:PRO:HD3	4.05	0.50
41:L4:5:GLN:HA	41:L4:21:PRO:HA	1.92	0.50
43:L6:149:ILE:HG23	43:L6:155:LEU:HD13	2.38	0.50
43:L6:64:LEU:HD22	43:L6:65:ILE:N	2.85	0.50
43:L6:65:ILE:O	43:L6:76:LEU:HA	2.41	0.50
45:L8:134:TYR:CG	45:L8:190:VAL:HG11	3.30	0.50
47:M0:188:GLY:HA3	47:M0:216:TYR:HD1	1.76	0.50
48:M1:11:ASP:N	48:M1:11:ASP:OD1	2.41	0.50
51:M5:177:GLY:HA2	36:5:68:C:O3'	110.73	0.50
55:M9:41:ILE:O	55:M9:45:VAL:HG23	2.11	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:N0:12:ARG:HG3	56:N0:13:ARG:O	2.42	0.50
56:N0:146:LYS:HG3	56:N0:147:ASP:N	2.25	0.50
66:O0:23:TYR:OH	66:O0:83:LYS:HE2	4.21	0.50
68:O2:26:HIS:O	68:O2:28:VAL:N	2.44	0.50
72:O6:53:TYR:O	72:O6:57:LEU:N	2.61	0.50
2:S0:56:LYS:HZ2	2:S0:158:VAL:HA	3.49	0.50
2:S0:197:ILE:HG21	2:S0:201:LEU:HD22	1.93	0.50
36:1:1878:G:C2'	36:1:1879:A:H5'	2.42	0.50
36:1:2971:A:H3'	36:1:2971:A:N3	2.26	0.50
36:1:363:G:H2'	36:1:364:G:O4'	2.12	0.50
1:2:130:C:HO2'	1:2:131:C:P	2.33	0.50
1:2:138:A:H61	1:2:266:A:H61	1.59	0.50
1:2:872:G:O6	86:2:2126:OHX:N3	2.45	0.50
36:5:1192:C:N4	36:5:1302:A:P	2.85	0.50
36:5:2111:G:H4'	36:5:2112:U:OP2	2.12	0.50
53:M7:69:ARG:NH2	36:5:2991:A:N3	194.78	0.50
36:5:873:C:H5''	36:5:874:U:O5'	2.12	0.50
1:6:1092:A:O2'	1:6:1093:A:H3'	2.10	0.50
7:S5:166:ARG:NH2	1:6:1163:A:O3'	348.16	0.50
1:6:1698:G:N2	1:6:1699:G:N7	2.58	0.50
25:D3:65:ASN:ND2	1:6:574:G:O6	365.36	0.50
1:6:647:G:H1	1:6:687:G:H22	1.57	0.50
86:8:216:OHX:N2	86:8:223:OHX:N1	2.59	0.50
13:C1:90:TYR:CD1	13:C1:90:TYR:C	2.84	0.50
17:C5:69:GLU:OE1	86:C5:201:OHX:N4	2.44	0.50
22:D0:72:ASN:OD1	22:D0:73:GLY:N	4.23	0.50
25:D3:47:SER:HB3	1:6:600:U:H1'	354.57	0.50
26:D4:108:ARG:O	26:D4:111:LYS:HB3	2.12	0.50
26:D4:63:GLN:OE1	26:D4:64:PHE:N	3.48	0.50
27:D5:49:ARG:O	27:D5:53:GLU:HB2	2.28	0.50
27:D5:54:VAL:HG13	27:D5:57:TYR:HD1	1.76	0.50
40:L3:57:VAL:HG23	40:L3:358:TRP:HE3	1.76	0.50
49:M3:54:LEU:HD12	49:M3:75:PHE:CZ	2.47	0.50
50:M4:19:ARG:NH2	50:M4:69:THR:HG23	3.22	0.50
55:M9:141:HIS:O	55:M9:145:ALA:N	2.42	0.50
57:N1:53:PRO:HB3	57:N1:91:LEU:HD22	2.37	0.50
59:N3:33:ASN:ND2	59:N3:63:LYS:H	2.10	0.50
62:N6:39:LEU:HD12	62:N6:106:ILE:HG22	1.93	0.50
63:N7:15:ARG:HH11	63:N7:15:ARG:HG3	2.60	0.50
64:N8:82:ILE:HB	64:N8:87:ARG:HG3	1.94	0.50
76:Q0:96:CYS:C	76:Q0:98:LYS:H	2.15	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:S7:52:ALA:HB3	9:S7:167:GLU:OE1	4.12	0.50
10:S8:170:SER:OG	10:S8:181:GLY:HA2	2.12	0.50
34:SR:123:ILE:HG21	34:SR:169:ILE:HD13	2.03	0.50
34:SR:249:ARG:HH12	34:SR:315:VAL:HG21	3.93	0.50
36:1:1162:U:H4'	68:O2:57:TYR:CE1	2.47	0.50
36:1:1511:U:H5''	36:1:1512:U:H5	1.77	0.50
36:1:1919:G:H1'	36:1:1934:G:N2	2.27	0.50
36:1:2689:A:H2'	36:1:2689:A:N3	2.26	0.50
1:2:1316:G:HO2'	1:2:1401:A:HO2'	1.59	0.50
1:2:1460:A:OP2	35:SM:68:ARG:HD3	2.11	0.50
1:2:252:U:H2'	1:2:253:A:C8	2.45	0.50
1:2:462:G:N7	86:2:2143:OHX:N1	2.60	0.50
1:2:784:C:H2'	1:2:785:U:O4'	2.12	0.50
38:4:46:G:OP2	75:O9:15:LYS:NZ	2.40	0.50
36:5:1323:G:C2'	36:5:1324:U:H5'	2.42	0.50
36:5:2344:U:H2'	36:5:2345:A:H8	1.76	0.50
36:5:2524:A:O2'	36:5:2525:G:OP2	2.26	0.50
36:5:2641:U:H5''	36:5:2642:A:OP1	2.12	0.50
86:5:3966:OHX:N1	86:5:4238:OHX:N5	2.60	0.50
1:6:1079:U:H2'	1:6:1080:U:O4'	2.12	0.50
5:S3:162:GLN:HG3	1:6:1333:C:O4'	426.96	0.50
1:6:1626:U:O4	86:6:2201:OHX:N5	2.45	0.50
1:6:641:G:H2'	1:6:642:G:H8	1.76	0.50
1:6:754:A:N6	1:6:793:A:H62	2.09	0.50
14:C2:62:LEU:HB3	14:C2:75:VAL:HG11	1.93	0.50
17:C5:110:GLU:H	17:C5:110:GLU:CD	2.40	0.50
23:D1:71:ARG:HG2	23:D1:75:ASN:HD21	1.76	0.50
27:D5:84:GLU:HB2	27:D5:89:ILE:HD11	1.94	0.50
16:C4:112:ILE:H	28:D6:57:SER:HA	1.76	0.50
39:L2:246:LEU:HD23	39:L2:248:GLY:N	7.27	0.50
39:L2:51:ASP:HB3	39:L2:54:ARG:HD2	3.28	0.50
39:L2:96:LEU:O	79:Q3:87:ARG:NH1	2.45	0.50
41:L4:271:LYS:HB2	41:L4:274:TYR:CB	2.94	0.50
42:L5:107:ARG:HH12	42:L5:120:LYS:HA	2.20	0.50
36:1:2748:A:O2'	42:L5:48:LYS:HE2	2.11	0.50
47:M0:171:TRP:CE3	47:M0:178:ARG:HD2	3.62	0.50
47:M0:96:VAL:HG11	47:M0:122:PRO:HB3	2.23	0.50
48:M1:89:TYR:HB3	48:M1:169:ALA:CB	2.41	0.50
52:M6:108:ILE:HG12	52:M6:108:ILE:O	4.72	0.50
52:M6:33:ILE:HG22	52:M6:102:LEU:CD1	2.37	0.50
57:N1:110:LYS:NZ	36:5:1068:C:OP1	245.78	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
67:O1:19:ARG:HD3	67:O1:35:GLU:CG	2.41	0.50
69:O3:48:ARG:CG	69:O3:48:ARG:HH11	2.24	0.50
71:O5:71:LYS:HE3	71:O5:72:GLY:N	2.27	0.50
71:O5:93:THR:OG1	71:O5:96:GLU:HG3	2.28	0.50
73:O7:39:TYR:CD2	73:O7:40:PRO:HA	2.58	0.50
2:S0:202:TYR:H	2:S0:202:TYR:HD2	2.03	0.50
3:S1:48:VAL:HG11	3:S1:61:LEU:HB2	6.07	0.50
7:S5:225:ARG:NH2	30:D8:58:GLU:H	5.06	0.50
9:S7:62:VAL:O	9:S7:95:GLU:HB3	2.11	0.50
10:S8:136:SER:HB3	10:S8:139:ALA:HB3	1.94	0.50
35:SM:88:ARG:HG2	35:SM:91:THR:OG1	2.12	0.50
34:SR:161:LYS:O	34:SR:161:LYS:CG	2.55	0.50
34:SR:171:SER:N	34:SR:179:LYS:O	2.44	0.50
36:1:2400:G:H5''	36:1:2401:A:OP2	2.12	0.50
36:1:247:C:H2'	36:1:248:U:H6	1.76	0.50
36:1:3302:U:H3	36:1:3312:U:H3	1.59	0.50
86:1:3951:OHX:N4	44:L7:217:PRO:HA	2.27	0.50
36:1:534:U:O4	56:N0:144:LEU:HD23	2.12	0.50
1:2:918:U:H2'	1:2:919:A:C8	2.47	0.50
47:M0:119:TRP:HZ3	36:5:1126:G:H5''	257.51	0.50
36:5:1291:A:H2'	36:5:1292:C:O4'	2.12	0.50
36:5:132:C:N4	36:5:134:U:C4	2.79	0.50
36:5:2444:C:H42	36:5:2503:G:H22	1.59	0.50
1:6:1395:G:H1	1:6:1403:C:H42	1.60	0.50
1:6:487:G:H3'	1:6:488:G:C5'	2.41	0.50
15:C3:20:ARG:NH2	24:D2:56:HIS:HB3	2.26	0.50
16:C4:117:ASP:OD2	16:C4:119:THR:OG1	2.19	0.50
17:C5:22:LEU:HD21	17:C5:109:PRO:HB3	1.94	0.50
17:C5:53:PRO:O	17:C5:56:PHE:HB3	2.12	0.50
17:C5:89:MET:O	17:C5:107:ILE:HG13	2.76	0.50
28:D6:36:ILE:HD12	28:D6:78:ALA:HB1	1.92	0.50
42:L5:60:ILE:H	42:L5:80:SER:HB3	1.77	0.50
44:L7:158:LYS:HD2	44:L7:159:GLN:H	2.01	0.50
46:L9:106:LYS:HG3	46:L9:107:ASP:OD1	3.94	0.50
46:L9:136:PHE:HE2	46:L9:144:ILE:HG12	5.71	0.50
46:L9:31:ARG:HD3	46:L9:149:ASN:OD1	2.77	0.50
46:L9:49:ASN:ND2	46:L9:49:ASN:O	2.45	0.50
46:L9:91:ARG:HH21	46:L9:91:ARG:HG3	1.77	0.50
49:M3:115:ARG:NH1	49:M3:147:ILE:HG12	2.27	0.50
53:M7:95:LEU:HD23	53:M7:148:LEU:HD11	2.88	0.50
36:1:1608:C:H5''	61:N5:111:ASN:ND2	2.27	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
61:N5:96:LYS:HE3	61:N5:107:VAL:HB	1.93	0.50
66:O0:30:THR:HB	66:O0:91:SER:HB2	1.93	0.50
67:O1:23:VAL:O	67:O1:28:ARG:NH1	2.93	0.50
69:O3:90:PRO:O	69:O3:92:LYS:N	2.42	0.50
74:O8:17:ARG:NH2	36:5:1824:U:O3'	139.37	0.50
2:S0:125:ASP:O	2:S0:127:ARG:N	2.45	0.50
2:S0:42:PRO:O	2:S0:43:ASP:HB2	2.12	0.50
2:S0:72:ASP:OD2	2:S0:72:ASP:N	3.10	0.50
4:S2:140:ARG:HB2	4:S2:222:TYR:CD1	2.60	0.50
5:S3:192:PRO:HB2	5:S3:201:ALA:HA	2.89	0.50
5:S3:6:SER:HB2	1:6:1514:U:H1'	439.08	0.50
5:S3:20:GLU:OE2	5:S3:76:ARG:NH2	3.38	0.50
6:S4:38:LEU:O	6:S4:41:SER:OG	4.57	0.50
36:1:1281:G:H2'	36:1:1282:G:C8	2.44	0.50
36:1:1439:U:H2'	36:1:1440:G:H8	1.77	0.50
36:1:2371:G:O6	86:1:3864:OHX:N3	2.45	0.50
36:1:2782:U:OP1	49:M3:185:LYS:NZ	2.45	0.50
36:1:3166:C:H2'	36:1:3167:A:O4'	2.12	0.50
36:1:3218:A:HO2'	36:1:3278:C:H5	1.60	0.50
36:1:780:A:H8	36:1:780:A:O5'	1.95	0.50
36:1:7:C:H2'	36:1:8:C:H6	1.76	0.50
1:2:1335:U:H5'	22:D0:85:ARG:HH22	1.77	0.50
1:2:1789:G:H8	1:2:1789:G:H5''	1.77	0.50
1:2:192:U:HO2'	1:2:193:U:P	2.35	0.50
1:2:484:C:N4	1:2:503:G:H22	2.10	0.50
1:2:558:U:O2'	1:2:559:C:O5'	2.26	0.50
1:2:863:A:OP1	24:D2:57:ARG:NE	2.41	0.50
36:5:1000:C:C2	36:5:1045:C:N4	2.80	0.50
36:5:150:A:H2'	36:5:151:A:H5'	1.94	0.50
36:5:2546:C:H2'	36:5:2547:A:H8	1.76	0.50
36:5:414:U:H2'	36:5:415:G:C8	2.46	0.50
36:5:975:C:H2'	36:5:976:U:C6	2.46	0.50
1:6:1358:G:H2'	1:6:1359:C:H6	1.76	0.50
1:6:15:U:H2'	1:6:16:G:O4'	2.12	0.50
26:D4:8:ARG:HD2	1:6:780:A:C2	439.82	0.50
37:7:25:G:H2'	37:7:26:C:O4'	2.12	0.50
12:C0:11:ILE:HD11	12:C0:42:VAL:HG22	1.94	0.50
15:C3:31:GLU:OE1	15:C3:31:GLU:N	5.68	0.50
15:C3:47:PRO:HG3	15:C3:75:LEU:HD22	1.94	0.50
17:C5:18:ARG:HD3	20:C8:90:ASN:OD1	2.12	0.50
23:D1:9:VAL:HG22	23:D1:10:GLU:H	2.20	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:525:A:H5''	26:D4:89:TYR:CE1	2.47	0.50
7:S5:143:ARG:HG2	30:D8:55:VAL:HB	2.07	0.50
31:D9:38:ILE:HG22	31:D9:42:CYS:HB3	2.17	0.50
32:E0:13:LYS:O	32:E0:16:SER:N	2.44	0.50
40:L3:159:ARG:HG2	40:L3:182:GLN:HA	3.67	0.50
46:L9:105:GLU:HB2	46:L9:110:LYS:H	1.75	0.50
47:M0:63:GLU:H	47:M0:63:GLU:CD	2.15	0.50
48:M1:91:LEU:O	48:M1:172:LEU:N	4.42	0.50
53:M7:131:ARG:HH11	53:M7:131:ARG:HG3	1.77	0.50
53:M7:53:ASP:O	86:M7:207:OHX:N3	2.44	0.50
57:N1:38:ASP:O	57:N1:64:VAL:HG23	2.12	0.50
57:N1:57:TYR:CD1	57:N1:89:LEU:HD21	2.46	0.50
61:N5:38:LEU:HD22	61:N5:40:LEU:HD13	3.30	0.50
64:N8:70:LYS:HE2	64:N8:129:PHE:CD2	2.47	0.50
64:N8:8:THR:HG21	36:5:662:U:OP1	149.33	0.50
36:1:1485:G:N2	70:O4:4:ARG:HD2	2.27	0.50
78:Q2:47:GLN:HE22	78:Q2:53:GLN:HA	1.76	0.50
36:1:860:G:H5''	79:Q3:17:ARG:NH1	2.27	0.50
3:S1:23:PRO:O	3:S1:27:LYS:HG3	2.47	0.50
6:S4:123:LEU:HD21	6:S4:235:TYR:HB3	1.93	0.50
10:S8:163:GLY:HA3	36:1:3354:U:H1'	1.94	0.50
10:S8:36:THR:HG23	10:S8:96:LEU:O	2.39	0.50
34:SR:101:GLN:HG2	34:SR:137:LYS:O	2.12	0.50
34:SR:13:LEU:O	34:SR:309:VAL:HG13	2.11	0.50
36:1:122:A:OP1	36:1:122:A:H3'	2.12	0.50
36:1:1334:U:O2'	44:L7:151:ARG:NH2	2.45	0.50
36:1:1472:U:H2'	36:1:1473:G:H8	1.77	0.50
36:1:177:U:C4	36:1:178:U:C4	3.00	0.50
36:1:1952:G:H5'	36:1:1953:G:OP2	2.12	0.50
36:1:2842:U:C5	36:1:2843:U:C5	3.00	0.50
86:1:4050:OHX:N2	86:1:4160:OHX:N1	2.60	0.50
36:1:901:G:H2'	36:1:902:G:H8	1.77	0.50
36:1:900:G:H2'	36:1:901:G:H8	1.77	0.50
1:2:1274:C:H41	35:SM:95:SER:HA	1.77	0.50
1:2:1429:G:H2'	1:2:1430:U:C6	2.46	0.50
1:2:1684:U:O2	1:2:1718:G:N2	2.45	0.50
1:2:38:C:C2'	1:2:39:A:H5'	2.42	0.50
1:2:436:A:O5'	1:2:436:A:H8	1.94	0.50
37:3:23:A:HO2'	37:3:121:U:HO3'	1.55	0.50
37:3:37:G:N2	37:3:41:G:O4'	2.45	0.50
36:5:1340:G:C6	36:5:1341:U:C4	3.00	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:123:A:C6	36:5:150:A:C5	3.00	0.50
70:O4:7:PHE:HE2	36:5:1856:C:H1'	152.95	0.50
39:L2:152:SER:N	36:5:2157:G:O6	218.99	0.50
36:5:2661:G:H1	36:5:2709:C:N4	2.09	0.50
36:5:2758:A:C8	36:5:2759:U:C5	3.00	0.50
40:L3:275:ARG:NH1	36:5:3045:G:O3'	234.46	0.50
36:5:3266:G:C6	36:5:3267:A:C6	3.00	0.50
36:5:537:A:C2	36:5:557:A:C4	2.99	0.50
1:6:138:A:H62	1:6:266:A:N6	2.10	0.50
1:6:192:U:H1'	1:6:193:U:C4	2.46	0.50
1:6:30:G:H2'	1:6:31:C:C6	2.47	0.50
12:C0:23:ALA:CB	12:C0:64:TYR:HB2	2.74	0.50
14:C2:59:LEU:HA	14:C2:87:PRO:HB2	1.94	0.50
17:C5:34:VAL:HG11	17:C5:45:PHE:CZ	4.39	0.50
17:C5:18:ARG:HD2	17:C5:36:LEU:O	2.71	0.50
18:C6:136:SER:O	18:C6:137:ARG:NH2	2.45	0.50
19:C7:21:TYR:HE2	19:C7:61:ILE:HG21	2.34	0.50
22:D0:72:ASN:ND2	22:D0:73:GLY:H	2.09	0.50
23:D1:74:GLN:HG2	23:D1:79:LEU:CB	4.80	0.50
25:D3:14:LYS:HA	25:D3:17:VAL:HG12	4.39	0.50
31:D9:39:CYS:O	31:D9:43:PHE:N	2.55	0.50
31:D9:49:ASP:OD1	31:D9:49:ASP:N	3.91	0.50
39:L2:10:LYS:HA	39:L2:16:PHE:CD2	2.46	0.50
40:L3:108:GLU:HB2	40:L3:137:TYR:CE1	2.47	0.50
40:L3:159:ARG:HB3	40:L3:182:GLN:HA	1.93	0.50
46:L9:161:LEU:HD22	46:L9:161:LEU:O	2.72	0.50
47:M0:68:ALA:HB2	47:M0:158:LYS:HB2	1.94	0.50
49:M3:119:TYR:HD1	49:M3:145:PHE:CZ	3.38	0.50
49:M3:171:ARG:O	49:M3:174:ARG:HB3	2.12	0.50
49:M3:79:GLU:OE2	49:M3:103:ASN:ND2	2.30	0.50
51:M5:27:VAL:HB	51:M5:122:ASN:HD21	1.76	0.50
52:M6:3:VAL:HG22	52:M6:4:GLU:HG3	1.93	0.50
54:M8:133:LYS:HB2	54:M8:135:GLN:NE2	3.19	0.50
55:M9:106:LEU:HD22	55:M9:138:LEU:HD11	2.52	0.50
57:N1:108:ARG:O	57:N1:112:ASN:N	3.22	0.50
58:N2:20:SER:OG	58:N2:21:SER:N	2.44	0.50
64:N8:7:LYS:O	64:N8:10:LYS:N	2.37	0.50
65:N9:14:ARG:NH2	65:N9:18:ARG:HD3	4.05	0.50
68:O2:20:HIS:HB3	68:O2:35:GLN:OE1	2.12	0.50
2:S0:122:ILE:HA	2:S0:144:ILE:O	2.12	0.50
2:S0:153:SER:O	2:S0:156:VAL:HG22	3.25	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:S2:163:GLY:HA3	4:S2:209:ASN:ND2	2.27	0.50
8:S6:148:SER:O	8:S6:150:GLU:N	2.42	0.50
1:2:142:G:N7	8:S6:177:ARG:NH1	2.58	0.50
9:S7:184:GLU:OE1	9:S7:187:SER:OG	9.24	0.50
34:SR:21:THR:HG23	34:SR:37:SER:HA	3.90	0.50
36:1:1110:U:H2'	36:1:1111:U:C6	2.47	0.49
36:1:1176:C:OP1	52:M6:25:LYS:HE3	2.12	0.49
36:1:1343:A:H2'	36:1:1344:G:C8	2.47	0.49
36:1:1478:C:H2'	36:1:1479:U:H6	1.76	0.49
36:1:1521:G:C2	36:1:1522:U:H5	2.30	0.49
36:1:439:C:H5'	36:1:440:A:OP2	2.12	0.49
1:2:123:G:N2	6:S4:146:THR:HG21	2.27	0.49
1:2:477:A:OP1	32:E0:30:PRO:HA	2.12	0.49
36:5:1060:U:H2'	36:5:1061:A:H8	1.75	0.49
36:5:1068:C:H2'	36:5:1069:C:C6	2.47	0.49
36:5:1239:C:H3'	36:5:1240:A:H8	1.76	0.49
69:O3:19:SER:HB3	36:5:1330:A:OP1	232.79	0.49
36:5:1743:G:H2'	36:5:1744:G:C8	2.47	0.49
36:5:873:C:H4'	36:5:1908:A:H5'	1.94	0.49
36:5:2283:G:N3	36:5:2285:C:N4	2.60	0.49
53:M7:139:TYR:CE1	36:5:2355:G:H5'	143.71	0.49
45:L8:38:GLN:HB2	36:5:2557:A:H2	207.85	0.49
36:5:3234:A:H8	36:5:3234:A:OP2	1.94	0.49
1:6:1467:C:H2'	1:6:1468:U:C6	2.46	0.49
1:6:1670:G:O6	86:6:2190:OHX:N4	2.44	0.49
1:6:837:G:O6	86:6:2100:OHX:N1	2.45	0.49
12:C0:46:LEU:O	12:C0:50:THR:HG23	2.19	0.49
13:C1:78:THR:HG21	13:C1:118:GLN:HA	2.92	0.49
14:C2:129:GLU:HA	14:C2:133:LEU:HD22	1.94	0.49
15:C3:150:VAL:HG12	15:C3:151:ASN:OD1	2.11	0.49
17:C5:22:LEU:HD23	17:C5:23:GLU:H	5.17	0.49
20:C8:14:ILE:HA	20:C8:22:VAL:O	2.12	0.49
21:C9:33:TYR:OH	21:C9:99:SER:OG	2.29	0.49
21:C9:49:ASP:HB3	21:C9:53:TRP:HB3	1.94	0.49
1:2:1798:U:C5	28:D6:38:ARG:NH2	2.80	0.49
39:L2:140:ASN:OD1	39:L2:142:ASP:HB3	5.43	0.49
40:L3:21:ARG:HG2	40:L3:269:GLN:HG2	1.94	0.49
41:L4:216:VAL:HG23	41:L4:217:LYS:HG2	1.94	0.49
44:L7:134:VAL:O	44:L7:229:PHE:HA	2.58	0.49
45:L8:161:GLU:CD	51:M5:26:ARG:HH22	2.88	0.49
45:L8:163:VAL:HG22	45:L8:166:LEU:HD12	1.93	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:M1:160:VAL:HG12	48:M1:161:SER:N	2.90	0.49
49:M3:121:SER:OG	49:M3:122:LYS:N	2.44	0.49
51:M5:39:ALA:HB3	51:M5:61:ILE:HG22	2.63	0.49
36:1:784:A:C6	54:M8:93:ILE:HG22	2.47	0.49
62:N6:58:VAL:O	62:N6:65:GLY:N	3.12	0.49
66:O0:15:ALA:O	66:O0:18:ILE:HG22	2.12	0.49
70:O4:25:THR:HG23	70:O4:29:ILE:O	3.17	0.49
71:O5:86:ARG:O	71:O5:90:ARG:NE	2.68	0.49
73:O7:37:CYS:SG	73:O7:38:GLY:N	2.85	0.49
77:Q1:4:LYS:HG2	77:Q1:5:TRP:CE3	2.47	0.49
79:Q3:45:LYS:O	79:Q3:45:LYS:HG3	2.12	0.49
2:S0:136:ALA:HA	2:S0:141:ILE:HD11	3.92	0.49
3:S1:226:GLY:HA2	36:5:2536:A:H4'	257.64	0.49
4:S2:90:THR:C	4:S2:92:ALA:H	2.55	0.49
6:S4:124:GLY:HA2	6:S4:142:HIS:CE1	3.53	0.49
6:S4:180:LEU:N	6:S4:229:GLY:O	3.00	0.49
7:S5:117:THR:O	7:S5:120:ILE:N	3.01	0.49
34:SR:232:TYR:HD2	34:SR:232:TYR:H	2.46	0.49
34:SR:248:ASN:HD21	34:SR:298:GLY:HA3	2.43	0.49
36:1:1381:A:C2	36:1:1426:C:C2	3.00	0.49
36:1:1626:U:H2'	36:1:1627:U:C6	2.47	0.49
36:1:1686:U:OP1	58:N2:42:LYS:NZ	2.32	0.49
36:1:2767:U:OP2	86:1:4129:OHX:N2	2.45	0.49
36:1:2795:U:OP1	78:Q2:62:ALA:N	2.42	0.49
36:1:3012:A:C2	36:1:3043:C:H1'	2.47	0.49
1:2:1492:A:C4	1:2:1493:A:C8	3.00	0.49
1:2:1500:C:H5'	21:C9:106:GLN:NE2	2.26	0.49
1:2:968:U:H5''	1:2:1033:C:O2'	2.11	0.49
36:5:1228:C:H2'	36:5:1229:G:H8	1.76	0.49
36:5:513:G:H2'	36:5:514:G:O4'	2.12	0.49
1:6:1003:A:HO2'	1:6:1005:A:H62	1.54	0.49
1:6:1334:U:C4	1:6:1335:U:C4	3.00	0.49
34:SR:63:GLY:HA2	1:6:1341:A:OP1	451.28	0.49
1:6:1392:U:H2'	1:6:1393:C:C6	2.45	0.49
86:6:2104:OHX:N5	86:6:2190:OHX:N6	2.60	0.49
14:C2:45:LEU:O	14:C2:49:THR:HG23	2.12	0.49
15:C3:132:VAL:HG23	15:C3:134:VAL:HG13	1.92	0.49
16:C4:106:ALA:HA	16:C4:112:ILE:HD11	1.94	0.49
16:C4:17:ALA:HA	16:C4:30:VAL:HG22	4.88	0.49
16:C4:57:PRO:HA	16:C4:60:ALA:HB3	1.92	0.49
17:C5:12:PHE:HD1	17:C5:13:LYS:H	1.60	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:C7:37:GLU:HG3	34:SR:150:TRP:HE1	1.77	0.49
20:C8:44:ASN:HD21	20:C8:48:LYS:HE3	1.77	0.49
21:C9:16:ASN:HA	21:C9:56:LYS:HZ3	1.77	0.49
21:C9:73:VAL:HG21	21:C9:102:ARG:HG3	3.35	0.49
24:D2:50:PHE:HB3	24:D2:63:VAL:HG22	1.93	0.49
25:D3:49:ALA:O	25:D3:104:LEU:HB2	2.11	0.49
26:D4:23:PHE:HE2	26:D4:75:VAL:HG12	1.77	0.49
26:D4:83:LYS:HE2	26:D4:96:LEU:HB3	1.93	0.49
27:D5:44:GLN:O	27:D5:47:TYR:HB3	2.48	0.49
24:D2:22:LYS:HZ3	29:D7:3:LEU:H	1.59	0.49
29:D7:54:VAL:HG12	29:D7:63:LEU:HB2	3.15	0.49
40:L3:229:VAL:HG13	40:L3:235:THR:HG21	3.08	0.49
41:L4:209:TYR:CE2	41:L4:212:ASP:HB2	2.48	0.49
41:L4:3:ARG:NE	41:L4:22:LEU:O	2.44	0.49
42:L5:110:LEU:HA	42:L5:113:LEU:HB2	3.50	0.49
42:L5:85:ARG:HD2	42:L5:86:TYR:CZ	2.46	0.49
44:L7:140:SER:O	44:L7:143:THR:N	2.43	0.49
44:L7:141:TYR:HA	44:L7:144:ILE:HD12	3.02	0.49
45:L8:78:PHE:C	45:L8:80:TYR:H	2.32	0.49
46:L9:37:ASN:OD1	46:L9:39:LYS:HB2	2.12	0.49
48:M1:60:ARG:HB2	48:M1:60:ARG:NH2	4.69	0.49
49:M3:17:HIS:HB3	49:M3:20:GLU:HG3	3.37	0.49
49:M3:46:ILE:HA	49:M3:49:ARG:NH1	4.30	0.49
52:M6:185:ALA:O	52:M6:187:GLU:N	3.62	0.49
53:M7:26:PHE:CE2	53:M7:121:GLN:HG2	2.47	0.49
36:1:1348:U:OP2	54:M8:38:ARG:NH2	2.45	0.49
55:M9:15:VAL:CG1	55:M9:52:LYS:HG3	2.42	0.49
55:M9:8:LYS:O	55:M9:11:ALA:HB3	2.12	0.49
56:N0:12:ARG:HH11	56:N0:22:PRO:HD2	1.76	0.49
36:1:1682:U:O4	58:N2:90:ARG:NH1	2.45	0.49
61:N5:34:LEU:HB2	36:5:1558:A:H1'	141.72	0.49
63:N7:53:VAL:HA	63:N7:57:HIS:HD2	1.77	0.49
63:N7:8:GLY:HA2	63:N7:25:ILE:O	3.88	0.49
64:N8:90:TYR:HA	64:N8:93:SER:HB3	5.58	0.49
66:O0:41:LEU:HD22	66:O0:66:LYS:O	2.12	0.49
67:O1:11:GLU:HG2	67:O1:74:ARG:HB2	1.94	0.49
2:S0:74:VAL:HG23	2:S0:118:PRO:HB3	1.92	0.49
3:S1:137:ILE:HG22	3:S1:215:VAL:CG2	2.42	0.49
3:S1:70:LEU:O	3:S1:74:GLN:HB2	2.11	0.49
4:S2:102:VAL:O	4:S2:113:LEU:HD23	2.37	0.49
35:SM:162:GLN:O	35:SM:164:ASN:N	2.77	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:C2:127:GLY:HA2	35:SM:168:GLU:HA	1.93	0.49
35:SM:64:LYS:O	35:SM:66:ALA:N	3.76	0.49
34:SR:134:TRP:HA	34:SR:140:CYS:HA	1.94	0.49
36:1:1916:U:H2'	36:1:1917:C:O4'	2.12	0.49
36:1:2137:U:C6	36:1:2141:U:C4	3.00	0.49
36:1:313:A:C6	36:1:314:U:C4	3.00	0.49
36:1:796:U:H2'	36:1:797:U:H6	1.77	0.49
1:2:1527:C:H2'	1:2:1528:U:C6	2.46	0.49
1:2:1764:C:H3'	1:2:1767:G:N7	2.27	0.49
1:2:1485:C:OP1	86:2:2099:OHX:N6	2.45	0.49
1:2:818:C:N4	1:2:819:G:O6	2.42	0.49
36:1:1419:A:H5'	38:4:20:U:O3'	2.12	0.49
56:N0:139:TYR:OH	36:5:1213:G:OP1	323.39	0.49
36:5:2171:G:H2'	36:5:2172:A:H8	1.77	0.49
36:5:2542:U:H1'	36:5:2543:U:C5	2.47	0.49
36:5:550:A:H2'	36:5:551:A:C8	2.48	0.49
36:5:795:G:O6	86:5:3928:OHX:N6	2.45	0.49
1:6:1189:A:N6	1:6:1190:C:H41	2.11	0.49
16:C4:136:ARG:HD2	1:6:1769:U:O2	302.49	0.49
1:6:9:U:O4	86:6:2146:OHX:N3	2.45	0.49
10:S8:5:ARG:NH1	1:6:332:U:O2'	300.17	0.49
1:6:938:G:O6	86:6:2105:OHX:N3	2.45	0.49
12:C0:16:PHE:O	12:C0:88:PRO:HA	2.12	0.49
15:C3:65:VAL:O	15:C3:67:THR:N	3.57	0.49
1:2:1601:G:N2	21:C9:88:VAL:HG22	2.27	0.49
23:D1:16:LYS:HD2	23:D1:21:ASN:O	2.61	0.49
23:D1:18:SER:HG	23:D1:54:ALA:H	2.32	0.49
39:L2:86:GLN:HG2	39:L2:88:ILE:HD11	2.71	0.49
41:L4:125:ALA:HB1	41:L4:238:LEU:HB3	1.94	0.49
41:L4:145:ILE:O	86:L4:404:OHX:N5	2.45	0.49
42:L5:140:ARG:HB2	36:5:1080:A:OP1	228.28	0.49
42:L5:148:ILE:HG13	42:L5:159:VAL:HG21	1.93	0.49
47:M0:16:PRO:O	47:M0:18:PRO:HD3	2.11	0.49
36:1:1048:A:H2'	47:M0:22:TYR:CE1	2.48	0.49
50:M4:125:LYS:HA	50:M4:128:ARG:HH12	4.81	0.49
50:M4:19:ARG:HB3	50:M4:35:ILE:HD12	2.20	0.49
50:M4:36:VAL:HB	50:M4:45:LEU:HD23	2.22	0.49
59:N3:87:ARG:HG3	59:N3:89:ASP:OD1	2.13	0.49
63:N7:46:ILE:HD11	63:N7:49:TYR:CD2	2.52	0.49
70:O4:86:LYS:O	70:O4:90:ILE:HG12	2.12	0.49
2:S0:175:TYR:CD1	2:S0:199:PRO:HA	2.47	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:S1:127:VAL:HG21	3:S1:173:THR:HG22	1.94	0.49
3:S1:70:LEU:HD13	3:S1:71:ALA:N	2.27	0.49
4:S2:116:LYS:HG3	4:S2:117:THR:N	2.80	0.49
6:S4:134:LYS:O	6:S4:136:VAL:HG23	3.33	0.49
6:S4:154:ILE:HG12	6:S4:172:PHE:CD2	3.33	0.49
6:S4:26:CYS:HB3	11:S9:2:PRO:O	2.12	0.49
11:S9:6:ARG:HB2	11:S9:6:ARG:HH11	2.05	0.49
1:2:1402:G:H2'	1:2:1403:C:O4'	2.13	0.49
1:2:1536:G:N1	1:2:1538:U:O2	2.45	0.49
1:2:285:G:H2'	1:2:286:C:H6	1.77	0.49
1:2:66:U:C5	8:S6:173:PRO:HG3	2.47	0.49
36:5:1276:U:H2'	36:5:1277:C:C6	2.47	0.49
36:5:1317:A:C4	36:5:1319:G:N7	2.80	0.49
68:O2:64:LYS:NZ	36:5:1405:U:OP1	180.97	0.49
36:5:2661:G:H2'	36:5:2662:G:H8	1.77	0.49
36:5:3329:U:H2'	36:5:3330:A:H5''	1.93	0.49
1:6:1553:G:O2'	1:6:1555:A:N7	2.45	0.49
13:C1:133:LYS:NZ	1:6:324:U:OP1	292.76	0.49
15:C3:87:ASP:HB3	15:C3:125:LEU:HD11	4.51	0.49
15:C3:47:PRO:HG2	15:C3:86:GLU:OE2	4.01	0.49
16:C4:50:ALA:HB3	16:C4:53:ASP:HB2	1.94	0.49
19:C7:86:PRO:HG2	19:C7:88:VAL:HA	9.51	0.49
40:L3:53:MET:HE1	40:L3:327:CYS:HB3	2.61	0.49
41:L4:25:VAL:O	41:L4:27:SER:N	2.45	0.49
42:L5:41:LYS:HA	42:L5:41:LYS:HE3	3.70	0.49
45:L8:91:PHE:O	45:L8:95:ASN:HB2	2.53	0.49
46:L9:49:ASN:C	46:L9:51:GLN:H	2.13	0.49
47:M0:80:SER:HB3	47:M0:147:VAL:HG11	1.95	0.49
49:M3:105:ASN:ND2	49:M3:108:ILE:HG12	4.04	0.49
53:M7:67:ILE:HG22	53:M7:80:LYS:HB3	1.93	0.49
55:M9:123:LEU:O	55:M9:127:SER:OG	2.82	0.49
56:N0:134:ASP:O	56:N0:136:LYS:HG2	2.47	0.49
41:L4:362:ASP:C	56:N0:26:ARG:HH12	2.99	0.49
56:N0:89:ASN:OD1	57:N1:156:TYR:N	2.58	0.49
8:S6:131:LYS:HB2	60:N4:81:PRO:O	2.11	0.49
66:O0:100:ILE:HG13	66:O0:101:LEU:HD13	7.46	0.49
66:O0:16:LEU:HD12	66:O0:98:SER:N	2.27	0.49
67:O1:36:ILE:O	67:O1:39:PHE:HB3	2.13	0.49
68:O2:105:ARG:HE	68:O2:124:GLY:CA	2.23	0.49
71:O5:41:LEU:O	71:O5:44:ILE:HG22	4.08	0.49
61:N5:45:LYS:HG2	71:O5:75:TYR:CD2	2.47	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:S0:75:ALA:HB1	2:S0:174:TRP:CH2	3.46	0.49
2:S0:41:ARG:HB2	2:S0:45:VAL:HB	4.52	0.49
3:S1:139:ALA:HB2	3:S1:172:LEU:HD11	2.15	0.49
4:S2:81:MET:HB2	4:S2:101:VAL:HG12	2.32	0.49
4:S2:162:CYS:O	4:S2:165:VAL:N	2.45	0.49
4:S2:176:SER:HB2	4:S2:195:ASP:HB3	2.67	0.49
5:S3:30:ALA:C	5:S3:32:GLU:H	2.14	0.49
8:S6:105:ASP:OD2	8:S6:105:ASP:N	2.74	0.49
11:S9:111:THR:O	11:S9:114:TYR:N	2.82	0.49
36:1:1018:G:N7	36:1:1035:G:N2	2.61	0.49
36:1:1245:A:N6	36:1:1272:C:O2'	2.45	0.49
36:1:1580:A:H5'	36:1:2522:G:N7	2.27	0.49
36:1:290:G:H2'	36:1:291:C:H6	1.77	0.49
36:1:3386:G:H2'	36:1:3387:U:C6	2.48	0.49
1:2:1738:U:H2'	1:2:1739:C:C6	2.48	0.49
1:2:69:G:H1	1:2:82:U:H3	1.59	0.49
36:5:1470:U:OP1	86:5:3950:OHX:N6	2.46	0.49
36:5:2335:G:N2	36:5:2339:C:O2	2.38	0.49
36:5:70:A:N1	36:5:313:A:O2'	2.39	0.49
1:6:1031:U:H4'	1:6:1032:G:OP2	2.12	0.49
38:8:83:C:H4'	38:8:85:G:N3	2.28	0.49
12:C0:24:LYS:HB2	12:C0:63:TYR:CE1	2.62	0.49
16:C4:31:THR:OG1	16:C4:34:SER:O	5.95	0.49
17:C5:15:HIS:CG	17:C5:16:SER:H	3.15	0.49
19:C7:106:THR:O	19:C7:110:VAL:HG22	3.40	0.49
19:C7:6:THR:OG1	19:C7:8:THR:HG23	5.74	0.49
21:C9:65:ILE:HD11	21:C9:76:LEU:HD11	1.95	0.49
28:D6:66:LYS:HD3	28:D6:66:LYS:H	1.78	0.49
36:1:2177:G:OP2	39:L2:128:ARG:NH1	2.43	0.49
39:L2:46:LYS:HD2	39:L2:62:VAL:HG11	2.89	0.49
40:L3:213:GLU:HB2	40:L3:282:ILE:HG13	5.07	0.49
40:L3:227:GLU:HG3	40:L3:270:ARG:NE	4.24	0.49
44:L7:180:SER:HB2	44:L7:183:ASP:N	2.26	0.49
44:L7:210:PRO:HD3	44:L7:243:MET:HG2	1.95	0.49
44:L7:85:PHE:H	44:L7:139:PRO:HD3	1.77	0.49
45:L8:34:PHE:H	45:L8:39:ALA:HB3	4.83	0.49
47:M0:73:ASN:O	47:M0:77:THR:HG23	2.12	0.49
48:M1:82:ARG:CG	48:M1:112:LEU:HB2	2.43	0.49
52:M6:92:THR:O	52:M6:96:LYS:HG3	2.78	0.49
36:1:976:U:P	54:M8:144:ARG:HH22	2.36	0.49
54:M8:69:ARG:O	54:M8:71:LEU:N	3.46	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
63:N7:88:ASP:HB3	63:N7:121:ARG:NH2	2.27	0.49
71:O5:85:THR:HG22	71:O5:87:ALA:H	1.77	0.49
76:Q0:122:ARG:HH11	76:Q0:122:ARG:HG3	1.77	0.49
78:Q2:9:LYS:HD2	36:5:2713:U:OP1	227.68	0.49
3:S1:23:PRO:HB2	3:S1:27:LYS:HE2	5.39	0.49
4:S2:78:ASP:HA	4:S2:104:VAL:HG12	1.93	0.49
7:S5:225:ARG:CZ	30:D8:58:GLU:HB2	4.99	0.49
7:S5:94:THR:OG1	7:S5:95:ASN:N	2.46	0.49
1:2:803:A:N3	9:S7:104:ARG:NE	2.60	0.49
9:S7:182:VAL:HG12	9:S7:183:PHE:H	1.77	0.49
9:S7:71:HIS:CG	9:S7:131:PHE:HZ	2.30	0.49
11:S9:121:SER:HB3	11:S9:124:HIS:HB2	1.94	0.49
11:S9:171:ARG:NH1	11:S9:174:ARG:HG3	2.27	0.49
35:SM:123:ALA:O	35:SM:126:ASP:HB2	2.11	0.49
35:SM:64:LYS:HD3	35:SM:64:LYS:H	1.77	0.49
36:1:1650:G:O6	86:1:4134:OHX:N2	2.46	0.49
36:1:2252:A:N6	36:1:2264:U:H3	2.09	0.49
36:1:3295:A:H2'	36:1:3296:A:C8	2.48	0.49
36:1:3325:G:H5''	67:O1:103:GLY:HA2	1.95	0.49
86:1:3998:OHX:N6	86:1:4169:OHX:N5	2.60	0.49
1:2:159:U:O2'	8:S6:87:ARG:NH1	2.46	0.49
1:2:87:C:H1'	1:2:168:A:N1	2.27	0.49
1:2:1153:G:N7	86:2:2167:OHX:N1	2.60	0.49
1:2:751:G:H2'	1:2:752:A:C8	2.48	0.49
38:4:103:G:C6	38:4:105:A:C6	3.01	0.49
36:5:1119:C:H2'	36:5:1120:A:C8	2.47	0.49
1:6:1629:G:H2'	1:6:1630:U:O4'	2.13	0.49
1:6:228:G:H22	1:6:237:C:N4	2.10	0.49
1:6:323:A:H61	1:6:345:U:H3	1.59	0.49
1:6:754:A:OP1	1:6:754:A:H4'	2.11	0.49
38:8:2:A:H3'	38:8:3:A:C8	2.41	0.49
20:C8:108:LYS:HA	20:C8:111:ASP:HB2	2.53	0.49
21:C9:27:LYS:HD2	21:C9:27:LYS:O	6.07	0.49
25:D3:87:VAL:HG22	25:D3:124:VAL:HG21	1.93	0.49
36:1:2180:G:P	39:L2:174:ARG:HH22	2.35	0.49
45:L8:161:GLU:OE1	51:M5:26:ARG:NH1	2.89	0.49
46:L9:171:ASP:OD1	46:L9:173:ARG:HD3	2.12	0.49
46:L9:86:TYR:CE1	46:L9:151:VAL:HG13	2.48	0.49
51:M5:24:ARG:HH11	51:M5:24:ARG:CG	3.96	0.49
59:N3:13:ILE:HG13	59:N3:85:TRP:CG	3.89	0.49
62:N6:4:GLN:HB2	36:5:229:G:H5''	69.23	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
67:O1:31:ARG:HB3	67:O1:31:ARG:NH1	2.26	0.49
67:O1:80:ASN:OD1	67:O1:81:GLU:N	2.46	0.49
67:O1:9:THR:HG21	67:O1:74:ARG:HH11	1.78	0.49
70:O4:9:ARG:HG2	70:O4:34:HIS:NE2	4.86	0.49
71:O5:62:GLN:O	71:O5:65:ALA:HB3	2.12	0.49
73:O7:87:SER:O	86:O7:104:OHX:N4	2.46	0.49
36:1:1845:G:O2'	73:O7:5:THR:HG22	2.11	0.49
78:Q2:3:ASN:O	36:5:2655:U:H5'	238.70	0.49
2:S0:26:ALA:HB1	2:S0:29:VAL:HG13	1.94	0.49
4:S2:130:ILE:O	4:S2:134:LEU:HD22	2.12	0.49
5:S3:195:SER:HB2	5:S3:200:LYS:HG3	1.94	0.49
6:S4:123:LEU:HA	6:S4:160:VAL:O	2.12	0.49
6:S4:182:TYR:CE1	6:S4:192:ILE:HD11	3.28	0.49
7:S5:56:ALA:O	7:S5:58:LEU:N	3.53	0.49
7:S5:89:ILE:HD12	7:S5:90:ILE:H	2.10	0.49
9:S7:131:PHE:HB3	9:S7:132:PRO:CD	2.42	0.49
1:2:209:U:H5'	10:S8:171:SER:HB3	1.94	0.49
11:S9:163:PRO:HG2	11:S9:164:PHE:CD2	2.48	0.49
36:1:1045:C:O2'	36:1:1046:A:H5'	2.12	0.49
36:1:1062:A:H4'	57:N1:105:PHE:CD2	2.48	0.49
36:1:1278:A:O2'	36:1:1279:C:H6	1.95	0.49
36:1:3273:A:OP2	43:L6:77:ARG:NH1	2.42	0.49
36:1:599:C:H5''	36:1:600:G:OP2	2.12	0.49
1:2:1244:A:O2'	1:2:1245:G:OP1	2.29	0.49
1:2:1396:U:H2'	1:2:1397:U:C6	2.48	0.49
1:2:1636:C:C2	1:2:1765:A:N6	2.81	0.49
86:2:2043:OHX:N1	86:2:2098:OHX:N5	2.60	0.49
1:2:77:U:H4'	1:2:78:A:O5'	2.13	0.49
1:2:795:U:H5	1:2:796:A:C4	2.30	0.49
36:5:1741:A:C6	36:5:1742:U:C2	3.00	0.49
36:5:2211:U:H5	36:5:2234:G:O6	1.96	0.49
36:5:237:G:C2	36:5:238:A:C8	3.01	0.49
36:5:3296:A:H2'	36:5:3297:U:C6	2.47	0.49
36:5:3357:U:O2'	36:5:3358:U:OP1	2.31	0.49
86:5:3966:OHX:N3	86:5:4238:OHX:N2	2.60	0.49
1:6:1529:C:H2'	1:6:1530:C:C6	2.48	0.49
1:6:1153:G:H1	1:6:1625:C:H42	1.61	0.49
1:6:357:G:OP2	86:6:2074:OHX:N6	2.46	0.49
1:6:488:G:N2	1:6:499:U:H3	2.11	0.49
1:6:689:G:H2'	1:6:690:G:O4'	2.13	0.49
1:6:74:U:C4	1:6:76:A:H5'	2.48	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:8:15:G:C6	38:8:16:G:N1	2.81	0.49
36:5:59:G:H2'	38:8:33:A:O2'	2.12	0.49
28:D6:44:ILE:HD12	28:D6:45:VAL:H	1.78	0.49
33:E1:106:TYR:CZ	33:E1:131:PHE:HZ	2.86	0.49
41:L4:128:ALA:HB1	41:L4:134:LEU:HD12	1.95	0.49
41:L4:232:SER:O	41:L4:233:LEU:HB2	2.21	0.49
41:L4:299:ILE:HG22	41:L4:300:ARG:O	2.11	0.49
42:L5:76:ALA:HB3	42:L5:109:THR:HG22	2.34	0.49
42:L5:265:TYR:HE1	37:7:121:U:C5'	316.56	0.49
42:L5:268:GLU:HA	42:L5:271:LYS:HE2	1.94	0.49
44:L7:158:LYS:O	44:L7:160:ARG:N	2.42	0.49
46:L9:90:MET:HB2	46:L9:144:ILE:CG2	2.43	0.49
46:L9:89:LYS:HG2	46:L9:145:VAL:HG22	1.97	0.49
52:M6:177:LYS:O	52:M6:180:SER:N	2.46	0.49
53:M7:70:THR:OG1	53:M7:71:ALA:N	2.63	0.49
54:M8:60:PRO:HG3	54:M8:144:ARG:HA	2.10	0.49
41:L4:300:ARG:NE	54:M8:39:ARG:HA	2.27	0.49
55:M9:6:THR:O	55:M9:10:LEU:HB2	2.13	0.49
60:N4:14:TYR:O	60:N4:17:ARG:HB3	2.13	0.49
60:N4:4:GLU:HG3	60:N4:30:ARG:CZ	4.43	0.49
3:S1:27:LYS:HA	3:S1:49:ASN:HA	1.94	0.49
5:S3:10:LYS:HG3	5:S3:11:LEU:HD23	2.52	0.49
6:S4:49:ARG:NH1	6:S4:50:ASN:HD21	2.10	0.49
1:2:803:A:H1'	9:S7:104:ARG:HE	1.77	0.49
10:S8:171:SER:HG	10:S8:180:ASP:H	1.56	0.49
11:S9:146:PHE:HZ	1:6:765:G:N1	432.09	0.49
36:1:1017:C:O2'	36:1:1018:G:OP2	2.31	0.49
36:1:2352:A:H5''	53:M7:83:TRP:O	2.11	0.49
36:1:2666:C:OP2	36:1:2687:G:N1	2.39	0.49
36:1:358:G:N2	36:1:361:A:OP2	2.35	0.49
36:1:65:A:H4'	36:1:66:A:O5'	2.13	0.49
36:1:888:A:H2'	36:1:889:U:O4'	2.11	0.49
1:2:194:U:O2'	1:2:195:G:O2'	2.24	0.49
1:2:641:G:H2'	1:2:642:G:H8	1.77	0.49
1:2:711:U:H1'	1:2:712:G:H5'	1.94	0.49
1:2:844:A:H2'	1:2:845:G:H8	1.77	0.49
37:3:25:G:H2'	37:3:26:C:O4'	2.11	0.49
65:N9:7:HIS:O	36:5:1135:A:H5'	226.51	0.49
36:5:1439:U:C2	36:5:1440:G:C8	3.01	0.49
70:O4:72:VAL:HG11	36:5:1639:C:H5''	192.91	0.49
36:5:2134:G:C2	36:5:2135:U:C6	3.01	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:235:A:H2'	36:5:236:G:O4'	2.11	0.49
36:5:856:G:C6	36:5:857:G:N1	2.80	0.49
68:O2:33:ARG:HD3	36:5:944:C:H4'	163.58	0.49
1:6:1228:G:OP2	1:6:1228:G:H4'	2.13	0.49
1:6:198:A:C2'	1:6:199:G:H5'	2.41	0.49
1:6:434:G:O6	86:6:2081:OHX:N2	2.45	0.49
1:6:230:C:H42	1:6:235:G:H1	1.61	0.49
1:6:585:A:H2'	1:6:586:G:C8	2.48	0.49
6:S4:6:LYS:HA	1:6:94:U:H4'	341.73	0.49
86:7:220:OHX:N4	86:7:228:OHX:N2	2.61	0.49
13:C1:91:LEU:HD22	13:C1:101:GLU:O	2.13	0.49
13:C1:83:THR:HB	13:C1:110:HIS:HA	1.95	0.49
14:C2:42:ALA:N	14:C2:122:VAL:O	2.87	0.49
16:C4:42:VAL:HA	16:C4:46:MET:SD	2.53	0.49
17:C5:68:PRO:HG2	17:C5:71:GLU:HB3	1.95	0.49
18:C6:139:GLN:HG3	1:6:1579:U:H1'	356.29	0.49
19:C7:71:PHE:CE2	19:C7:74:GLN:HB2	5.81	0.49
21:C9:76:LEU:O	21:C9:80:TYR:HD2	2.89	0.49
33:E1:146:SER:HB3	1:6:1234:A:H4'	434.90	0.49
36:1:2157:G:C6	39:L2:151:PRO:HD2	2.48	0.49
40:L3:112:ASP:O	40:L3:116:ARG:HB2	2.54	0.49
36:1:2392:C:H1'	40:L3:266:ARG:NH1	2.27	0.49
41:L4:180:LYS:NZ	41:L4:203:ARG:O	2.45	0.49
41:L4:251:THR:O	41:L4:254:ALA:HB3	2.13	0.49
43:L6:148:GLU:OE2	43:L6:151:LYS:HE3	2.12	0.49
45:L8:84:ARG:HH22	45:L8:181:LYS:NZ	2.10	0.49
46:L9:19:SER:HB3	50:M4:6:ILE:HB	2.66	0.49
47:M0:9:TYR:CG	47:M0:97:LEU:HD13	2.47	0.49
49:M3:168:ARG:HD3	49:M3:172:LEU:HD11	3.08	0.49
52:M6:41:LEU:HB2	52:M6:138:LEU:HD22	1.94	0.49
53:M7:54:HIS:HA	53:M7:83:TRP:CD1	2.48	0.49
55:M9:105:LEU:HD22	55:M9:138:LEU:HD22	1.95	0.49
56:N0:1:MET:HB2	56:N0:118:PHE:CD1	2.48	0.49
62:N6:11:ASP:HB3	62:N6:14:LYS:HG3	4.20	0.49
68:O2:74:PHE:CD2	68:O2:85:LEU:HD21	2.47	0.49
74:O8:12:LEU:HD11	74:O8:68:SER:O	4.49	0.49
76:Q0:99:CYS:O	76:Q0:100:TYR:HB2	2.13	0.49
79:Q3:18:TYR:O	79:Q3:22:LEU:HD12	2.35	0.49
3:S1:81:PHE:CD2	3:S1:82:ARG:HG3	2.48	0.49
7:S5:120:ILE:O	7:S5:124:LEU:HD13	3.15	0.49
9:S7:103:SER:O	9:S7:106:SER:HB3	2.13	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:S9:78:ARG:HH22	11:S9:82:ARG:HH21	1.61	0.49
36:1:1015:U:O4	36:1:1035:G:N2	2.46	0.49
36:1:2380:U:H2'	36:1:2381:G:H8	1.77	0.49
36:1:2407:C:H6	36:1:2407:C:O5'	1.96	0.49
36:1:3230:G:H2'	36:1:3231:U:O4'	2.12	0.49
1:2:1628:U:H2'	1:2:1629:G:C8	2.48	0.49
1:2:1793:G:H4'	1:2:1794:A:OP1	2.12	0.49
86:2:2043:OHX:N2	86:2:2098:OHX:N5	2.61	0.49
1:2:494:U:O2'	1:2:495:C:O5'	2.26	0.49
1:2:681:U:C4	1:2:682:C:H5	2.30	0.49
1:2:702:G:H21	1:2:703:G:H1'	1.77	0.49
1:2:712:G:H2'	1:2:713:A:O4'	2.12	0.49
36:5:1149:G:N1	36:5:1155:C:N4	2.60	0.49
36:5:257:U:H2'	36:5:258:G:H8	1.78	0.49
1:6:1392:U:H2'	1:6:1393:C:H6	1.78	0.49
1:6:1595:U:H3'	1:6:1596:C:O2	2.13	0.49
1:6:1685:G:N2	1:6:1717:G:H1'	2.28	0.49
4:S2:203:LYS:NZ	1:6:16:G:O6	367.36	0.49
1:6:848:C:H2'	1:6:849:C:C6	2.47	0.49
1:2:1401:A:O3'	19:C7:10:LYS:NZ	2.45	0.49
19:C7:27:ASP:OD2	19:C7:30:THR:N	2.42	0.49
20:C8:91:ASP:OD1	20:C8:93:THR:OG1	2.29	0.49
21:C9:38:LYS:HE3	1:6:1564:U:OP1	379.18	0.49
22:D0:27:THR:HB	22:D0:88:LYS:HG2	2.97	0.49
26:D4:122:GLY:O	26:D4:124:ARG:N	2.98	0.49
27:D5:38:HIS:CE1	27:D5:70:LYS:HA	2.48	0.49
28:D6:22:ARG:NH2	28:D6:27:SER:O	4.78	0.49
40:L3:211:GLN:NE2	40:L3:284:ARG:HA	2.27	0.49
40:L3:56:ILE:HG22	40:L3:74:GLU:HB2	2.18	0.49
42:L5:176:SER:OG	36:5:2747:A:OP1	244.30	0.49
47:M0:85:PHE:HA	47:M0:140:THR:HG22	2.03	0.49
48:M1:29:ARG:HA	48:M1:32:ARG:NH2	2.73	0.49
51:M5:44:ARG:HH11	51:M5:47:LYS:HG2	1.78	0.49
54:M8:64:VAL:HG13	54:M8:93:ILE:HD11	1.95	0.49
36:1:2756:C:O4'	57:N1:49:GLN:HG2	2.13	0.49
58:N2:95:PHE:HE1	58:N2:103:TYR:CD1	5.93	0.49
45:L8:45:ASN:OD1	61:N5:26:VAL:HA	2.13	0.49
64:N8:116:GLY:O	64:N8:137:LYS:NZ	5.30	0.49
73:O7:87:SER:O	86:O7:104:OHX:N1	2.46	0.49
5:S3:202:LEU:C	5:S3:204:ASP:H	2.69	0.49
6:S4:163:ASP:HB3	6:S4:167:GLY:O	4.69	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:S7:31:SER:O	9:S7:35:LYS:HE3	2.12	0.49
10:S8:22:ARG:CZ	10:S8:25:ARG:HD2	3.46	0.49
36:1:1655:G:O2'	36:1:1800:A:N6	2.44	0.49
36:1:213:A:N6	36:1:227:G:O2'	2.46	0.49
36:1:2303:A:OP1	77:Q1:23:ARG:NH2	2.46	0.49
36:1:2370:G:C6	36:1:2371:G:C6	3.00	0.49
36:1:847:A:H2'	36:1:848:A:H8	1.76	0.49
1:2:1151:A:H61	1:2:1627:U:H3	1.61	0.49
1:2:1480:G:H3'	1:2:1481:C:C6	2.47	0.49
1:2:1497:U:OP2	86:2:2031:OHX:N1	2.46	0.49
1:2:1765:A:C8	1:2:1768:G:N2	2.81	0.49
1:2:908:U:H5''	1:2:909:U:OP2	2.13	0.49
37:3:36:C:O2'	37:3:37:G:H5'	2.13	0.49
36:5:1299:U:H2'	36:5:1300:G:O4'	2.13	0.49
36:5:1618:G:H2'	36:5:1619:A:O4'	2.13	0.49
36:5:181:U:H1'	36:5:236:G:N2	2.27	0.49
39:L2:130:SER:OG	36:5:2179:C:O2'	215.84	0.49
36:5:2201:G:H2'	36:5:2202:C:H6	1.78	0.49
36:5:2647:A:N6	36:5:2648:G:C5	2.81	0.49
36:5:293:C:H2'	36:5:294:U:O4'	2.13	0.49
54:M8:107:THR:HG21	36:5:676:G:H3'	136.19	0.49
1:6:1194:A:H2'	1:6:1195:C:H5'	1.93	0.49
1:6:1200:G:H4'	1:6:1201:G:C5'	2.43	0.49
1:6:1588:G:OP1	86:6:2124:OHX:N2	2.46	0.49
86:6:2120:OHX:N6	86:6:2171:OHX:N5	2.59	0.49
1:6:482:U:O4	1:6:483:A:N6	2.46	0.49
1:6:581:U:H6	1:6:581:U:H3'	1.78	0.49
1:6:723:G:H5'	1:6:724:C:OP2	2.13	0.49
12:C0:38:LYS:HB2	12:C0:41:TYR:CD2	3.40	0.49
21:C9:54:PHE:CE2	21:C9:104:VAL:HG22	2.48	0.49
26:D4:113:ASN:O	26:D4:116:LYS:HB2	2.54	0.49
26:D4:19:ALA:HB1	26:D4:81:GLU:HG2	1.95	0.49
27:D5:62:VAL:O	27:D5:66:VAL:HG23	2.13	0.49
45:L8:133:LYS:HD2	45:L8:138:HIS:HE1	1.82	0.49
45:L8:139:VAL:O	45:L8:143:ILE:HG13	2.64	0.49
41:L4:105:THR:O	49:M3:26:PHE:HZ	1.96	0.49
1:2:814:A:C5'	55:M9:170:ARG:HH22	2.22	0.49
55:M9:6:THR:HG23	55:M9:9:ARG:HH12	5.51	0.49
56:N0:151:PRO:C	56:N0:153:PRO:HD3	2.75	0.49
61:N5:115:ARG:HH11	61:N5:115:ARG:CG	2.43	0.49
61:N5:67:ILE:CD1	61:N5:121:LYS:HG3	2.43	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
64:N8:36:GLY:HA3	64:N8:40:HIS:CE1	2.48	0.49
64:N8:73:LEU:HD22	64:N8:109:TYR:CE1	2.47	0.49
66:O0:74:ASN:OD1	66:O0:74:ASN:N	2.68	0.49
67:O1:32:ALA:O	67:O1:36:ILE:HD12	2.12	0.49
68:O2:43:ARG:HH11	68:O2:43:ARG:HG2	2.00	0.49
69:O3:96:ALA:HB2	36:5:3173:G:C2	230.44	0.49
36:1:816:A:OP1	73:O7:15:SER:OG	2.31	0.49
77:Q1:3:ALA:O	77:Q1:6:ARG:N	2.71	0.49
3:S1:164:ILE:O	3:S1:168:ILE:HG13	2.64	0.49
4:S2:144:TRP:CE2	4:S2:173:PRO:HG3	2.48	0.49
4:S2:179:VAL:HG11	1:6:2:A:H3'	392.20	0.49
6:S4:131:LEU:O	1:6:252:U:H5'	327.67	0.49
7:S5:94:THR:O	7:S5:97:LEU:N	2.39	0.49
9:S7:173:TYR:O	9:S7:177:THR:HG22	4.17	0.49
10:S8:117:TYR:CD1	10:S8:150:ALA:HB2	2.48	0.49
10:S8:8:ARG:HD3	10:S8:21:PHE:HD1	1.78	0.49
11:S9:37:LYS:HE2	1:6:594:A:OP2	413.45	0.49
34:SR:165:ASP:O	34:SR:166:SER:HB2	4.78	0.49
34:SR:239:GLU:O	34:SR:257:ALA:N	3.58	0.49
36:1:1722:U:H5''	55:M9:99:LEU:HD12	1.95	0.48
36:1:2434:U:H5	36:1:2594:C:OP2	1.96	0.48
36:1:2732:G:C6	36:1:2733:A:C5	3.01	0.48
36:1:295:A:OP1	72:O6:53:TYR:HE2	1.96	0.48
36:1:56:G:H1'	51:M5:162:ARG:HG3	1.94	0.48
36:1:650:C:O5'	36:1:650:C:H6	1.96	0.48
36:1:817:A:H2'	36:1:920:A:C2	2.48	0.48
36:1:926:A:H2'	36:1:927:C:C6	2.48	0.48
1:2:1370:U:H4'	1:2:1371:A:H5'	1.95	0.48
1:2:291:G:H2'	1:2:292:U:C6	2.47	0.48
1:2:307:G:OP1	13:C1:90:TYR:OH	2.31	0.48
1:2:927:C:H1'	16:C4:125:SER:HB2	1.93	0.48
56:N0:90:MET:HG2	36:5:1213:G:H4'	317.56	0.48
36:5:1340:G:H2'	36:5:1341:U:H6	1.77	0.48
36:5:2203:U:H1'	36:5:2240:G:N2	2.28	0.48
36:5:3121:U:H4'	36:5:3122:A:OP1	2.12	0.48
86:5:3994:OHX:N4	86:5:4085:OHX:N2	2.60	0.48
36:5:495:G:H2'	36:5:496:C:O4'	2.12	0.48
36:5:978:G:N2	36:5:1104:G:C5	2.81	0.48
1:6:1186:U:H2'	1:6:1187:U:O4'	2.13	0.48
1:6:1661:U:H2'	1:6:1662:G:C8	2.48	0.48
1:6:591:A:H2'	1:6:592:A:C8	2.48	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:7:23:A:C6	37:7:24:A:C6	3.01	0.48
37:7:57:G:H3'	37:7:58:C:C6	2.48	0.48
13:C1:86:ILE:HD11	13:C1:109:VAL:HG11	4.58	0.48
14:C2:119:SER:OG	14:C2:120:VAL:N	2.46	0.48
20:C8:105:VAL:HG13	20:C8:106:GLU:H	2.45	0.48
17:C5:108:ARG:HH21	20:C8:119:ILE:HD12	3.86	0.48
21:C9:5:SER:OG	21:C9:6:VAL:N	2.46	0.48
22:D0:20:ILE:HG22	22:D0:21:LYS:H	5.27	0.48
1:2:1095:U:O3'	24:D2:19:LYS:NZ	2.45	0.48
31:D9:10:HIS:CG	31:D9:11:PRO:HD2	2.48	0.48
33:E1:103:LEU:HA	33:E1:105:TYR:HD2	3.68	0.48
41:L4:140:HIS:NE2	41:L4:246:ARG:HG2	3.77	0.48
42:L5:181:PRO:HG2	42:L5:195:LEU:HD13	1.93	0.48
43:L6:80:ASN:OD1	43:L6:81:ALA:N	2.72	0.48
43:L6:97:ASN:O	43:L6:98:VAL:HB	2.16	0.48
51:M5:36:ILE:HG13	51:M5:64:VAL:HG23	2.96	0.48
52:M6:98:ALA:HA	52:M6:101:ARG:NH1	2.30	0.48
54:M8:16:ARG:NH1	54:M8:55:SER:HB3	2.28	0.48
56:N0:104:GLU:O	56:N0:108:GLN:HG2	2.13	0.48
57:N1:77:ASN:HB3	57:N1:84:TYR:HD2	1.77	0.48
58:N2:29:ASP:OD1	58:N2:31:ALA:HB3	2.13	0.48
61:N5:115:ARG:HG3	61:N5:115:ARG:NH1	2.45	0.48
61:N5:121:LYS:HD3	61:N5:123:TYR:CE1	3.79	0.48
63:N7:37:PRO:HD2	63:N7:38:PHE:CD1	2.48	0.48
63:N7:97:SER:H	63:N7:100:THR:HG1	2.75	0.48
67:O1:72:ARG:O	67:O1:96:VAL:HG13	2.13	0.48
2:S0:175:TYR:HD2	2:S0:176:LEU:HD23	1.78	0.48
4:S2:162:CYS:SG	4:S2:212:LYS:HE2	2.52	0.48
7:S5:63:GLN:H	7:S5:89:ILE:HG13	1.78	0.48
8:S6:186:ARG:O	8:S6:190:GLN:HG2	2.12	0.48
9:S7:91:ILE:HD12	9:S7:92:PHE:H	2.83	0.48
11:S9:171:ARG:HE	11:S9:174:ARG:HB2	4.82	0.48
35:SM:58:GLU:OE2	35:SM:62:ARG:HD2	5.34	0.48
34:SR:135:THR:O	34:SR:138:GLY:N	2.41	0.48
36:1:1673:G:C5	36:1:1775:G:C2	3.01	0.48
36:1:2733:A:H2'	36:1:2734:A:O4'	2.14	0.48
36:1:3138:U:H2'	36:1:3139:A:H5''	1.95	0.48
1:2:1242:A:OP1	17:C5:59:LYS:NZ	2.45	0.48
1:2:1590:G:H2'	1:2:1591:C:C6	2.49	0.48
1:2:249:U:H3'	1:2:250:C:H5'	1.93	0.48
36:5:1246:G:C4	36:5:1264:G:C2	3.02	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:54:C:O2'	36:5:1547:G:H1'	2.13	0.48
36:5:2298:U:O4	36:5:2923:U:H5	1.96	0.48
36:5:2567:C:N4	36:5:2568:C:H41	2.11	0.48
36:5:2676:A:H4'	36:5:2677:G:O5'	2.12	0.48
1:6:1097:U:H4'	1:6:1098:U:O5'	2.12	0.48
37:7:23:A:O2'	37:7:121:U:O3'	2.16	0.48
38:8:10:A:H2'	38:8:11:C:C6	2.47	0.48
17:C5:111:MET:HG2	20:C8:119:ILE:HD11	4.70	0.48
19:C7:19:ARG:HG3	19:C7:20:TYR:CE1	2.48	0.48
20:C8:88:ARG:CZ	20:C8:108:LYS:HE2	2.42	0.48
20:C8:32:LEU:HD11	20:C8:47:CYS:SG	2.53	0.48
20:C8:3:LEU:HD23	20:C8:5:VAL:HG13	1.95	0.48
39:L2:244:GLY:N	36:5:2244:A:OP1	230.73	0.48
40:L3:209:PHE:HB3	40:L3:282:ILE:CD1	2.43	0.48
41:L4:6:VAL:N	41:L4:20:LEU:O	2.32	0.48
43:L6:80:ASN:HB2	36:5:3272:C:O2	247.23	0.48
46:L9:33:THR:O	46:L9:34:LEU:HD23	2.12	0.48
46:L9:92:TYR:N	46:L9:92:TYR:CD1	2.79	0.48
46:L9:9:GLN:HG2	46:L9:54:LYS:HD3	4.34	0.48
47:M0:150:GLU:O	47:M0:150:GLU:HG3	2.14	0.48
48:M1:81:GLU:O	48:M1:84:LEU:N	3.32	0.48
51:M5:140:LYS:HA	51:M5:143:ARG:HB2	3.14	0.48
51:M5:48:ALA:C	51:M5:53:TYR:HB3	2.70	0.48
61:N5:96:LYS:O	61:N5:100:LYS:HB2	2.67	0.48
62:N6:42:GLN:HE21	62:N6:127:GLU:HB3	1.79	0.48
63:N7:51:LEU:HB2	63:N7:65:ARG:HH11	1.78	0.48
79:Q3:75:ALA:O	79:Q3:79:VAL:HG23	2.47	0.48
2:S0:155:PHE:O	23:D1:60:ARG:NH2	3.45	0.48
4:S2:114:GLY:C	4:S2:115:ILE:HG13	2.32	0.48
5:S3:133:GLY:HA3	5:S3:156:PHE:O	2.13	0.48
7:S5:140:THR:HA	7:S5:214:LYS:HD2	1.94	0.48
8:S6:3:LEU:HD23	8:S6:109:LEU:HB3	3.23	0.48
34:SR:112:SER:HB2	34:SR:153:GLN:HA	2.91	0.48
36:1:1450:G:H1	36:1:2354:C:H42	1.60	0.48
36:1:1495:U:C5	36:1:1835:A:N1	2.77	0.48
36:1:2116:G:N3	36:1:2116:G:H5'	2.28	0.48
36:1:3039:C:OP1	40:L3:62:ARG:NH1	2.47	0.48
36:1:3150:A:OP1	40:L3:132:LYS:HB2	2.14	0.48
36:1:3336:A:O5'	36:1:3336:A:H8	1.96	0.48
36:1:677:A:OP1	54:M8:89:ASP:HB3	2.13	0.48
1:2:1402:G:H2'	1:2:1403:C:H6	1.78	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:1478:G:OP1	21:C9:39:THR:OG1	2.31	0.48
36:5:1404:G:N2	36:5:1407:A:OP2	2.39	0.48
36:5:2214:A:H8	36:5:2214:A:O5'	1.95	0.48
36:5:2363:A:O2'	36:5:2364:G:H5'	2.13	0.48
36:5:2805:G:N3	36:5:2967:A:H2	2.11	0.48
36:5:437:G:N2	36:5:622:A:H61	2.11	0.48
36:5:819:U:H6	36:5:819:U:O5'	1.96	0.48
1:6:1117:U:H2'	1:6:1118:G:C8	2.48	0.48
1:6:1230:A:H2'	1:6:1258:U:C5	2.46	0.48
1:6:194:U:H2'	1:6:195:G:H4'	1.95	0.48
11:S9:44:ARG:NH1	1:6:473:A:OP1	410.05	0.48
1:6:722:G:O2'	1:6:723:G:H5''	2.13	0.48
9:S7:107:ARG:NH2	1:6:741:C:O2	349.10	0.48
1:6:870:C:H42	1:6:957:G:H1	1.62	0.48
15:C3:71:ILE:HA	15:C3:74:ILE:HG13	1.95	0.48
17:C5:38:PRO:HA	1:6:1549:C:OP1	384.33	0.48
17:C5:60:LEU:HD23	17:C5:76:VAL:HG21	2.24	0.48
17:C5:75:PRO:HA	17:C5:93:VAL:HB	3.10	0.48
20:C8:11:PHE:CE1	27:D5:41:ILE:HG12	2.86	0.48
20:C8:69:ILE:O	20:C8:73:MET:HG3	2.13	0.48
9:S7:140:VAL:HB	24:D2:52:TYR:HB3	2.17	0.48
25:D3:109:ARG:HB3	25:D3:109:ARG:NH1	4.82	0.48
25:D3:26:GLU:HB3	25:D3:29:TYR:HB3	1.95	0.48
27:D5:59:TYR:HD2	27:D5:60:VAL:N	2.11	0.48
27:D5:71:ILE:HG21	27:D5:76:ALA:HB2	3.05	0.48
28:D6:36:ILE:HD12	28:D6:36:ILE:H	5.53	0.48
31:D9:41:GLN:HB3	1:6:1433:G:C4	403.69	0.48
31:D9:4:GLU:N	31:D9:4:GLU:OE1	4.97	0.48
41:L4:330:TYR:O	41:L4:333:VAL:HG13	2.13	0.48
36:1:2402:A:H2'	41:L4:67:THR:OG1	2.13	0.48
42:L5:50:ARG:NH1	42:L5:72:ASP:OD2	2.46	0.48
44:L7:88:ARG:CZ	44:L7:103:LEU:HD13	2.43	0.48
45:L8:91:PHE:CZ	45:L8:185:ARG:HD3	4.85	0.48
46:L9:101:VAL:HG12	46:L9:136:PHE:CZ	2.48	0.48
49:M3:46:ILE:HD13	49:M3:49:ARG:HB2	5.75	0.48
51:M5:150:TRP:CZ3	36:5:321:C:H5''	88.97	0.48
52:M6:21:SER:OG	36:5:1175:C:O4'	257.68	0.48
54:M8:120:GLU:OE2	54:M8:130:ARG:NH2	2.77	0.48
56:N0:12:ARG:HB3	56:N0:24:LEU:HD23	2.11	0.48
56:N0:151:PRO:O	56:N0:152:LEU:HD23	2.71	0.48
57:N1:54:HIS:CE1	57:N1:55:LYS:HG2	2.48	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
61:N5:131:ASP:OD2	61:N5:133:LEU:N	3.16	0.48
64:N8:63:LYS:HE2	64:N8:68:PHE:CE2	2.82	0.48
67:O1:72:ARG:NE	67:O1:104:LEU:HD12	2.28	0.48
68:O2:44:ARG:NH1	36:5:1145:G:OP1	206.40	0.48
72:O6:91:ASN:HA	72:O6:94:ILE:HD12	1.95	0.48
76:Q0:77:ILE:HG23	76:Q0:78:ILE:HG22	5.81	0.48
2:S0:175:TYR:OH	2:S0:195:TRP:HB3	2.82	0.48
3:S1:131:ASP:HB3	3:S1:180:THR:CG2	2.44	0.48
7:S5:184:PHE:CZ	7:S5:185:ARG:HG3	2.49	0.48
10:S8:159:GLN:HB2	10:S8:165:LEU:HD23	1.94	0.48
36:1:1791:C:H2'	36:1:1792:C:C6	2.49	0.48
36:1:2269:U:C2	36:1:2272:G:C2	3.00	0.48
36:1:2392:C:H5''	36:1:2393:G:OP2	2.13	0.48
36:1:2877:G:H2'	36:1:2878:G:O4'	2.14	0.48
36:1:3151:U:H4'	36:1:3294:A:H1'	1.95	0.48
36:1:374:A:HO2'	36:1:376:G:H8	1.59	0.48
1:2:1082:C:N4	1:2:1091:A:H62	2.10	0.48
1:2:1388:A:H4'	1:2:1389:C:O5'	2.14	0.48
1:2:142:G:N2	1:2:173:A:H2	2.09	0.48
1:2:71:A:H2'	1:2:72:A:O4'	2.13	0.48
36:5:2252:A:H3'	36:5:2253:G:H5''	1.95	0.48
36:5:2659:G:H4'	36:5:2751:G:O2'	2.13	0.48
36:5:3096:C:H2'	36:5:3097:C:C6	2.47	0.48
86:5:3966:OHX:N1	86:5:4238:OHX:N2	2.61	0.48
36:5:441:U:H2'	36:5:442:G:C8	2.48	0.48
36:5:644:G:OP1	36:5:1142:G:O2'	2.29	0.48
36:5:697:A:H2'	36:5:698:U:O4'	2.12	0.48
36:5:767:U:H1'	36:5:768:C:H6	1.78	0.48
1:6:484:C:N4	1:6:503:G:H1	2.11	0.48
1:6:538:A:H2	1:6:540:G:N2	2.11	0.48
1:6:619:A:H5'	1:6:620:A:OP2	2.13	0.48
1:6:621:A:N3	1:6:1107:G:H1'	2.27	0.48
1:6:830:U:C2'	1:6:831:U:H5'	2.42	0.48
1:6:938:G:N2	1:6:941:A:OP2	2.39	0.48
13:C1:6:THR:HB	13:C1:9:SER:HB3	1.95	0.48
15:C3:92:ILE:O	15:C3:96:VAL:HG23	2.13	0.48
20:C8:18:LEU:HD23	20:C8:35:ILE:HD13	3.63	0.48
24:D2:27:ILE:CG1	24:D2:61:ILE:HB	2.44	0.48
28:D6:6:ALA:HB3	1:6:1796:C:H5	346.55	0.48
39:L2:130:SER:OG	39:L2:174:ARG:NH2	3.68	0.48
42:L5:143:LYS:HA	42:L5:172:TYR:HB3	2.76	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:L5:180:PHE:HB3	42:L5:195:LEU:HD22	1.95	0.48
42:L5:86:TYR:CE1	42:L5:247:ILE:HA	2.74	0.48
42:L5:61:ILE:HG23	42:L5:79:TYR:CE1	2.66	0.48
43:L6:93:VAL:HG12	43:L6:93:VAL:O	3.01	0.48
44:L7:178:ILE:HA	44:L7:183:ASP:HB3	1.95	0.48
41:L4:342:LYS:HD3	44:L7:56:GLU:OE2	2.89	0.48
47:M0:210:ILE:HD13	47:M0:217:PHE:CE1	3.23	0.48
48:M1:40:LEU:HD12	48:M1:40:LEU:O	3.87	0.48
52:M6:56:ASP:O	52:M6:59:ARG:HG2	2.33	0.48
54:M8:104:LEU:O	54:M8:105:ARG:HG3	2.55	0.48
54:M8:185:LYS:HD3	54:M8:186:VAL:HG23	3.92	0.48
36:1:1321:G:N2	56:N0:112:ALA:HB2	2.28	0.48
58:N2:58:GLU:HG2	58:N2:59:ASP:N	4.64	0.48
61:N5:55:ASN:OD1	61:N5:56:ARG:N	2.45	0.48
61:N5:57:LEU:HD23	61:N5:61:LYS:HG2	5.79	0.48
63:N7:2:ALA:O	63:N7:4:PHE:N	2.46	0.48
66:O0:54:SER:HA	66:O0:57:GLU:OE2	3.31	0.48
69:O3:51:TYR:CE2	69:O3:53:TYR:HB3	3.35	0.48
70:O4:54:ILE:HD11	70:O4:78:GLY:HA2	2.58	0.48
71:O5:42:PRO:O	71:O5:44:ILE:N	2.98	0.48
3:S1:39:GLU:HB3	3:S1:73:LEU:O	2.12	0.48
3:S1:91:VAL:HG23	3:S1:96:LEU:HB3	1.94	0.48
5:S3:17:PHE:HE1	5:S3:79:TYR:HE2	4.51	0.48
10:S8:62:THR:HA	10:S8:76:THR:O	2.50	0.48
35:SM:84:LYS:H	35:SM:84:LYS:HD2	1.77	0.48
34:SR:274:LEU:HD13	34:SR:313:TRP:CD2	2.48	0.48
36:1:1245:A:C3'	36:1:1246:G:H5''	2.44	0.48
36:1:1638:A:N3	36:1:1709:C:H1'	2.28	0.48
36:1:1815:U:O2'	36:1:1816:A:OP2	2.27	0.48
36:1:2298:U:O4	36:1:2923:U:H5	1.96	0.48
36:1:2390:A:C5	36:1:2391:G:C8	3.01	0.48
36:1:3009:G:C5	36:1:3010:U:C5	3.01	0.48
36:1:1941:C:H1'	36:1:3362:A:C8	2.48	0.48
36:1:1486:G:N7	86:1:4152:OHX:N2	2.61	0.48
86:1:3965:OHX:N5	86:1:4153:OHX:N2	2.61	0.48
36:1:47:C:N4	36:1:48:A:C6	2.82	0.48
36:1:864:G:O6	36:1:893:C:H3'	2.13	0.48
36:1:887:G:H2'	36:1:888:A:C8	2.48	0.48
1:2:1039:A:H5'	1:2:1082:C:H5	1.77	0.48
1:2:772:G:H21	1:2:774:A:H1'	1.78	0.48
1:2:794:U:O2'	1:2:795:U:C2	2.66	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:992:A:C2	1:2:1012:U:N3	2.75	0.48
38:4:107:G:C2	38:4:116:G:C5	3.01	0.48
38:4:85:G:H3'	38:4:85:G:C8	2.48	0.48
43:L6:2:SER:N	36:5:1385:C:O2	134.53	0.48
36:5:1554:U:H4'	36:5:1555:U:OP1	2.13	0.48
36:5:2157:G:N2	36:5:2177:G:O2'	2.47	0.48
36:5:3174:A:C2'	36:5:3175:U:H5'	2.42	0.48
36:5:732:C:H2'	36:5:733:G:O4'	2.13	0.48
1:6:1050:G:N2	1:6:1068:C:O2	2.46	0.48
1:6:119:A:H1'	1:6:397:A:C5	2.48	0.48
14:C2:67:THR:HB	1:6:1228:G:N7	459.75	0.48
1:6:139:C:H4'	1:6:140:A:O5'	2.12	0.48
1:6:1666:U:C4	1:6:1736:G:C2	3.02	0.48
12:C0:8:ARG:HG3	12:C0:12:HIS:HD1	1.77	0.48
14:C2:66:VAL:HG11	14:C2:71:ILE:HD12	1.96	0.48
21:C9:79:LEU:O	21:C9:79:LEU:HG	3.14	0.48
25:D3:86:PHE:O	25:D3:88:PRO:HD3	2.13	0.48
33:E1:83:LYS:O	33:E1:84:VAL:HG12	2.13	0.48
39:L2:238:ILE:C	39:L2:240:ALA:H	2.81	0.48
40:L3:226:PHE:CE1	40:L3:268:GLY:HA2	4.26	0.48
42:L5:102:GLY:O	42:L5:106:ALA:N	2.37	0.48
42:L5:8:LYS:HG2	42:L5:12:TYR:CE2	4.21	0.48
42:L5:212:ALA:HB2	42:L5:219:PHE:CG	4.15	0.48
37:3:121:U:C2	42:L5:268:GLU:HB3	2.48	0.48
47:M0:76:MET:O	47:M0:80:SER:N	2.42	0.48
49:M3:177:LYS:HA	72:O6:11:LEU:HD13	1.96	0.48
49:M3:18:TRP:C	49:M3:20:GLU:N	2.66	0.48
52:M6:3:VAL:HG13	52:M6:4:GLU:H	1.79	0.48
52:M6:47:PHE:HA	52:M6:136:THR:OG1	2.12	0.48
52:M6:48:PHE:O	52:M6:52:LEU:HG	2.89	0.48
36:1:1307:G:O4'	52:M6:60:LYS:HE3	2.12	0.48
53:M7:95:LEU:HD23	53:M7:148:LEU:CD1	3.02	0.48
37:3:76:A:O2'	56:N0:50:LYS:NZ	2.47	0.48
59:N3:79:VAL:HB	59:N3:118:VAL:HG13	1.95	0.48
74:O8:59:ALA:O	74:O8:62:ALA:HB3	2.12	0.48
1:2:1643:U:H5'	77:Q1:9:ARG:NH2	2.28	0.48
2:S0:178:ALA:HA	2:S0:181:VAL:HG22	2.40	0.48
3:S1:137:ILE:HD13	3:S1:172:LEU:HD22	3.52	0.48
1:2:1290:U:P	4:S2:95:ARG:HH12	2.37	0.48
5:S3:138:VAL:HA	5:S3:183:GLY:O	2.62	0.48
6:S4:45:ILE:HG12	6:S4:45:ILE:O	2.14	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:S5:48:PHE:HD1	7:S5:64:VAL:HG13	1.78	0.48
7:S5:73:THR:HG23	18:C6:114:ARG:CD	2.43	0.48
7:S5:77:TYR:HA	7:S5:83:ARG:HG2	4.63	0.48
8:S6:182:GLN:NE2	8:S6:185:GLN:OE1	2.47	0.48
11:S9:112:GLN:HG3	11:S9:148:VAL:HG21	1.94	0.48
36:1:1443:G:O6	86:1:3970:OHX:N3	2.46	0.48
36:1:1633:C:H2'	36:1:1634:G:C8	2.48	0.48
36:1:1721:U:O2'	36:1:1723:A:N7	2.42	0.48
36:1:3227:A:C6	36:1:3228:C:N3	2.81	0.48
36:1:3384:U:H2'	36:1:3385:U:H6	1.79	0.48
36:1:2405:C:O3'	88:1:4211:BLS:H82	2.13	0.48
36:1:856:G:C6	36:1:857:G:N1	2.81	0.48
1:2:1183:A:C6	1:2:1184:A:N1	2.82	0.48
1:2:32:U:OP1	25:D3:139:LYS:NZ	2.42	0.48
1:2:544:A:H5''	1:2:545:A:OP2	2.12	0.48
36:5:229:G:C2	36:5:230:U:C2	3.02	0.48
36:5:2533:G:H2'	36:5:2534:G:C8	2.48	0.48
36:5:2698:G:H2'	36:5:2699:G:C8	2.47	0.48
36:5:3356:G:C6	36:5:3357:U:C4	3.02	0.48
1:6:1136:U:O2'	1:6:1137:A:H5'	2.14	0.48
1:6:1211:A:N6	1:6:1452:U:H3	2.08	0.48
8:S6:53:SER:OG	1:6:163:G:H4'	294.84	0.48
1:6:348:U:O4	86:6:2163:OHX:N4	2.46	0.48
1:6:53:G:H2'	1:6:54:C:C6	2.49	0.48
1:6:950:C:H2'	1:6:951:A:C8	2.49	0.48
37:7:3:U:H2'	37:7:4:U:H6	1.78	0.48
36:5:419:G:N2	38:8:5:U:C2	2.82	0.48
12:C0:14:TYR:CE1	12:C0:18:GLU:HG3	2.68	0.48
13:C1:130:PRO:HG3	13:C1:136:ARG:HD2	5.45	0.48
17:C5:22:LEU:HD13	17:C5:26:LEU:HD11	1.96	0.48
17:C5:34:VAL:HG11	17:C5:45:PHE:CE1	4.26	0.48
18:C6:47:LYS:HZ3	18:C6:114:ARG:NE	2.12	0.48
19:C7:20:TYR:CZ	19:C7:38:ILE:HG13	2.48	0.48
20:C8:65:GLU:HG2	20:C8:68:ARG:NH2	3.71	0.48
21:C9:35:ASP:OD2	21:C9:36:ILE:HG23	3.65	0.48
27:D5:54:VAL:HG13	27:D5:57:TYR:CD1	2.48	0.48
28:D6:87:ARG:NH2	28:D6:92:ARG:O	2.47	0.48
29:D7:14:SER:HA	29:D7:17:ARG:HE	1.79	0.48
31:D9:48:ASN:HD22	31:D9:53:ASN:HD21	1.62	0.48
36:1:2948:C:O2'	40:L3:242:THR:HA	2.13	0.48
45:L8:68:ARG:O	45:L8:69:LEU:HB2	4.48	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
46:L9:134:ILE:CD1	46:L9:146:LEU:HD23	2.44	0.48
47:M0:55:ASN:ND2	47:M0:162:GLN:OE1	3.14	0.48
47:M0:76:MET:CE	47:M0:148:VAL:HA	2.63	0.48
49:M3:16:LYS:O	49:M3:17:HIS:HB2	4.54	0.48
50:M4:32:LEU:HD11	50:M4:94:TRP:CG	2.82	0.48
50:M4:71:ALA:O	50:M4:84:LYS:HD3	2.14	0.48
53:M7:2:ALA:O	53:M7:3:ARG:HB2	2.30	0.48
36:1:2763:U:C5'	54:M8:176:ARG:HG3	2.42	0.48
57:N1:31:LEU:HD23	57:N1:31:LEU:HA	1.61	0.48
59:N3:17:LEU:HD11	59:N3:98:ASN:HB3	1.95	0.48
60:N4:6:ASP:HA	60:N4:30:ARG:O	2.14	0.48
62:N6:74:TYR:CE2	62:N6:77:LYS:HD2	4.08	0.48
62:N6:71:SER:N	62:N6:81:GLN:O	2.80	0.48
68:O2:80:LYS:HD3	36:5:1386:A:OP1	135.67	0.48
79:Q3:13:LYS:NZ	79:Q3:30:GLU:OE1	3.01	0.48
2:S0:52:LYS:HA	23:D1:82:VAL:HG22	1.93	0.48
3:S1:117:TRP:HA	3:S1:155:TYR:HE1	4.17	0.48
3:S1:88:VAL:HG11	3:S1:96:LEU:HD12	1.95	0.48
4:S2:141:ARG:HG2	23:D1:10:GLU:OE1	2.13	0.48
6:S4:179:LYS:N	6:S4:194:THR:O	2.46	0.48
7:S5:177:ILE:HA	7:S5:180:ARG:NH1	2.28	0.48
34:SR:199:ILE:HA	34:SR:215:GLY:HA3	1.95	0.48
34:SR:228:LYS:O	34:SR:229:LYS:HG3	2.12	0.48
34:SR:274:LEU:O	34:SR:276:PRO:HD3	3.22	0.48
36:1:1237:G:H2'	36:1:1237:G:N3	2.27	0.48
36:1:1687:U:H2'	58:N2:70:LYS:HZ3	1.79	0.48
36:1:1770:G:H5'	36:1:1771:C:OP2	2.13	0.48
36:1:1449:A:C2	36:1:2356:A:C4	3.01	0.48
36:1:2902:A:H2'	36:1:2903:A:O4'	2.13	0.48
36:1:3006:A:H2'	36:1:3007:U:O4'	2.14	0.48
36:1:657:A:OP1	86:1:4167:OHX:N5	2.46	0.48
36:1:594:U:H2'	36:1:609:G:O6	2.13	0.48
36:1:708:G:H5'	36:1:709:A:OP2	2.14	0.48
36:1:75:G:H5''	49:M3:58:VAL:CG2	2.42	0.48
1:2:1559:A:C6	20:C8:134:ARG:HD2	2.48	0.48
1:2:400:A:H8	10:S8:24:LYS:O	1.97	0.48
36:5:1418:A:H4'	36:5:1419:A:OP1	2.14	0.48
36:5:1631:C:H5''	36:5:1632:A:H5''	1.95	0.48
36:5:198:A:N3	36:5:218:G:O2'	2.44	0.48
86:5:3966:OHX:N4	86:5:4238:OHX:N2	2.61	0.48
49:M3:59:ARG:HD3	36:5:73:C:C2	92.82	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:1368:G:C5	1:6:1369:U:C5	3.02	0.48
1:6:1699:G:C2	1:6:1701:A:H5''	2.49	0.48
1:6:192:U:O2'	1:6:193:U:O5'	2.28	0.48
1:6:28:A:H2'	1:6:29:U:O4'	2.13	0.48
1:6:386:G:C6	1:6:387:A:N6	2.82	0.48
1:6:43:A:H5''	1:6:437:A:N1	2.28	0.48
25:D3:22:ASN:O	1:6:609:U:H5	336.96	0.48
1:6:717:C:H42	1:6:720:G:H1	1.62	0.48
51:M5:112:ASN:OD1	38:8:141:C:H1'	104.37	0.48
13:C1:14:GLN:OE1	1:6:327:U:H4'	277.38	0.48
14:C2:93:ASP:HB3	14:C2:96:GLN:HB2	4.17	0.48
17:C5:108:ARG:NH2	20:C8:119:ILE:HD12	4.65	0.48
22:D0:44:ASN:CG	22:D0:102:ARG:HH21	7.26	0.48
24:D2:14:ILE:HD11	24:D2:27:ILE:HD11	4.68	0.48
25:D3:53:VAL:HG23	25:D3:100:ASP:O	2.13	0.48
26:D4:9:THR:HG21	26:D4:48:TYR:OH	2.13	0.48
24:D2:24:GLN:OE1	29:D7:4:VAL:HA	2.12	0.48
39:L2:113:VAL:HG12	39:L2:166:ILE:HA	1.95	0.48
41:L4:269:SER:OG	41:L4:271:LYS:HG3	3.13	0.48
41:L4:330:TYR:O	41:L4:333:VAL:N	2.58	0.48
38:4:26:U:H5'	41:L4:53:SER:HB2	1.96	0.48
44:L7:137:GLY:HA3	44:L7:236:ILE:HB	1.95	0.48
45:L8:251:LYS:O	45:L8:255:SER:HB2	2.14	0.48
46:L9:120:ASP:OD1	46:L9:124:ARG:NH2	2.46	0.48
47:M0:75:TYR:HD1	47:M0:151:GLY:HA2	2.85	0.48
52:M6:125:ARG:HD2	52:M6:135:TYR:CD2	3.42	0.48
53:M7:109:ALA:HA	53:M7:112:LEU:HD22	3.78	0.48
55:M9:23:TRP:HB3	55:M9:51:VAL:HG22	1.96	0.48
57:N1:75:ILE:O	57:N1:75:ILE:HD13	2.14	0.48
58:N2:90:ARG:O	58:N2:91:ASP:HB2	2.36	0.48
61:N5:137:ASN:OD1	61:N5:137:ASN:N	2.41	0.48
49:M3:67:ARG:NH2	64:N8:108:GLY:HA2	3.13	0.48
64:N8:91:LEU:HD12	64:N8:121:VAL:HG21	1.96	0.48
66:O0:98:SER:OG	66:O0:100:ILE:HG13	2.13	0.48
68:O2:12:LYS:HD3	68:O2:57:TYR:O	2.17	0.48
72:O6:74:LYS:HG3	72:O6:80:PHE:HA	1.95	0.48
72:O6:95:ALA:O	72:O6:99:ARG:HB2	2.13	0.48
78:Q2:8:ARG:O	78:Q2:23:HIS:N	2.57	0.48
2:S0:88:LYS:O	2:S0:202:TYR:OH	2.28	0.48
3:S1:227:ALA:HA	3:S1:230:ALA:HB3	2.20	0.48
4:S2:169:LEU:HD11	4:S2:188:LEU:HD11	4.46	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:S4:173:ILE:HD11	6:S4:235:TYR:HD1	2.96	0.48
11:S9:31:ALA:HA	11:S9:36:LEU:HD12	2.56	0.48
11:S9:44:ARG:O	11:S9:48:GLN:HG3	2.14	0.48
34:SR:248:ASN:ND2	34:SR:297:ASP:O	2.45	0.48
34:SR:64:HIS:HD1	34:SR:86:ASP:CG	2.15	0.48
36:1:1781:C:H2'	36:1:1782:U:C6	2.48	0.48
36:1:1852:G:N7	86:1:3971:OHX:N3	2.62	0.48
36:1:2765:C:O3'	78:Q2:39:GLY:HA3	2.14	0.48
36:1:620:U:C4	36:1:622:A:N1	2.81	0.48
36:1:968:G:H2'	36:1:969:C:H6	1.76	0.48
1:2:1153:G:H2'	1:2:1154:G:O4'	2.14	0.48
1:2:1238:A:H2'	1:2:1239:U:O4'	2.13	0.48
1:2:1287:A:H4'	1:2:1288:G:OP1	2.13	0.48
1:2:1335:U:H3	1:2:1416:G:H1	1.62	0.48
1:2:136:C:H4'	1:2:137:U:OP1	2.12	0.48
1:2:1794:A:H1'	28:D6:79:ILE:HD12	1.94	0.48
35:SM:25:ILE:HG13	37:3:39:C:H5'	1.95	0.48
36:5:2621:G:H2'	36:5:2622:C:H6	1.79	0.48
78:Q2:98:LYS:HD2	36:5:2656:A:O5'	251.31	0.48
65:N9:12:GLN:NE2	36:5:954:U:H1'	212.32	0.48
1:6:1597:A:H2'	1:6:1598:U:O4'	2.13	0.48
1:6:256:A:H2'	1:6:257:A:O4'	2.13	0.48
1:6:312:A:C5	1:6:314:C:C4	3.02	0.48
1:6:450:U:O2'	1:6:451:A:H5'	2.14	0.48
6:S4:66:MET:HG3	1:6:454:U:N1	375.21	0.48
1:6:219:A:N6	1:6:843:U:C2	2.82	0.48
1:6:862:A:C2	1:6:963:A:C4	3.02	0.48
14:C2:41:LEU:O	14:C2:43:ARG:HG2	4.61	0.48
18:C6:131:GLY:HA3	18:C6:136:SER:O	2.21	0.48
20:C8:107:SER:O	20:C8:110:ARG:HB2	2.13	0.48
22:D0:68:ARG:NH2	22:D0:70:THR:OG1	4.88	0.48
7:S5:112:ARG:HD3	27:D5:95:HIS:NE2	3.54	0.48
32:E0:40:TYR:HD1	32:E0:41:THR:HG23	4.01	0.48
33:E1:117:LEU:HD22	33:E1:118:ARG:NH1	4.97	0.48
36:1:860:G:O5'	39:L2:181:LYS:NZ	2.44	0.48
40:L3:255:TRP:CD1	36:5:2395:G:H5''	216.54	0.48
41:L4:169:LEU:HD22	41:L4:249:ILE:HD12	3.50	0.48
42:L5:153:THR:HG23	42:L5:160:PHE:CZ	2.49	0.48
44:L7:217:PRO:HA	86:5:3995:OHX:N5	262.86	0.48
45:L8:160:ILE:O	45:L8:164:VAL:HG13	2.22	0.48
45:L8:81:THR:HG21	45:L8:181:LYS:NZ	3.11	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
46:L9:134:ILE:HD12	46:L9:146:LEU:HD23	1.95	0.48
46:L9:86:TYR:CE1	46:L9:151:VAL:HG22	4.05	0.48
46:L9:13:PRO:HG2	46:L9:16:VAL:HG13	1.96	0.48
50:M4:113:THR:HB	50:M4:116:GLU:HG3	1.95	0.48
51:M5:120:TRP:CE3	36:5:269:G:H5'	132.93	0.48
52:M6:116:LYS:HG3	52:M6:117:ARG:N	2.29	0.48
52:M6:8:VAL:HG22	52:M6:34:VAL:HG13	2.90	0.48
55:M9:90:PRO:HG2	55:M9:93:VAL:HG23	2.94	0.48
56:N0:80:ARG:HB2	56:N0:124:LEU:HD11	2.46	0.48
57:N1:8:ARG:HH11	57:N1:15:PHE:HD1	2.54	0.48
59:N3:86:ARG:HD2	59:N3:92:PHE:CZ	2.49	0.48
59:N3:94:TYR:CZ	60:N4:21:PHE:HB2	2.49	0.48
67:O1:16:LEU:HD12	67:O1:16:LEU:HA	1.39	0.48
69:O3:45:LEU:HA	69:O3:71:VAL:HG12	2.21	0.48
3:S1:158:SER:HA	3:S1:161:ILE:HD12	2.43	0.48
4:S2:245:ASP:OD1	4:S2:245:ASP:N	2.45	0.48
5:S3:116:ARG:CZ	5:S3:116:ARG:HB2	5.17	0.48
5:S3:65:ARG:O	5:S3:69:LEU:HB2	2.79	0.48
6:S4:29:PRO:O	6:S4:31:PRO:HD3	2.13	0.48
7:S5:205:SER:C	7:S5:207:THR:H	2.17	0.48
7:S5:82:PHE:CZ	30:D8:49:ARG:HB3	2.48	0.48
10:S8:156:VAL:O	10:S8:159:GLN:HB2	3.55	0.48
34:SR:115:ILE:HG13	34:SR:116:ASP:N	2.29	0.48
34:SR:68:VAL:HA	34:SR:84:SER:HB2	2.65	0.48
36:1:1094:U:H1'	36:1:1096:U:H2'	1.95	0.48
36:1:2613:U:O2	36:1:2804:A:C8	2.67	0.48
36:1:3019:U:C4	36:1:3020:U:C4	3.02	0.48
36:1:915:A:C5	36:1:917:A:H1'	2.48	0.48
36:1:962:A:N1	36:1:2814:G:O2'	2.40	0.48
1:2:1107:G:O2'	1:2:1108:G:H5'	2.14	0.48
1:2:387:A:H8	1:2:387:A:OP2	1.96	0.48
37:3:20:A:C2	37:3:21:G:C4	3.02	0.48
36:5:1072:G:H2'	36:5:1073:U:H6	1.78	0.48
36:5:1764:U:H2'	36:5:1765:U:O4'	2.13	0.48
36:5:1841:A:H5'	36:5:1849:C:OP1	2.14	0.48
36:5:287:G:H2'	36:5:288:C:H6	1.79	0.48
36:5:760:G:H1'	36:5:770:G:H22	1.78	0.48
1:6:1202:A:H2'	1:6:1203:A:H5''	1.94	0.48
1:6:550:A:OP2	86:6:2049:OHX:N2	2.47	0.48
38:8:74:U:O2	86:8:219:OHX:N5	2.47	0.48
38:8:78:G:H2'	38:8:79:A:O4'	2.14	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:C0:56:LYS:CG	12:C0:67:THR:HB	2.43	0.48
19:C7:96:SER:HA	19:C7:97:ASN:HA	1.57	0.48
23:D1:55:LEU:HD13	23:D1:65:SER:OG	3.19	0.48
25:D3:24:TRP:CE3	25:D3:30:LYS:HG3	4.09	0.48
25:D3:89:ASN:HB2	25:D3:92:CYS:SG	2.59	0.48
26:D4:121:THR:HG22	26:D4:123:LYS:HB2	8.39	0.48
27:D5:38:HIS:HA	27:D5:70:LYS:HG2	9.47	0.48
29:D7:29:ARG:HG3	29:D7:29:ARG:HH11	2.27	0.48
30:D8:9:LEU:HD12	30:D8:34:GLU:OE2	2.13	0.48
39:L2:30:ARG:HA	39:L2:74:GLU:OE2	2.13	0.48
41:L4:334:PHE:CE1	41:L4:339:LEU:HB3	2.49	0.48
42:L5:122:VAL:HG23	42:L5:125:VAL:H	1.79	0.48
42:L5:41:LYS:HG2	57:N1:93:VAL:HG11	1.96	0.48
43:L6:18:LEU:N	43:L6:18:LEU:HD22	2.28	0.48
43:L6:42:LEU:HD11	43:L6:52:VAL:HG21	2.93	0.48
45:L8:41:GLN:CG	45:L8:44:ARG:HH12	3.00	0.48
48:M1:85:LYS:NZ	48:M1:85:LYS:HB2	2.29	0.48
49:M3:3:ILE:HG21	64:N8:45:MET:HE2	4.89	0.48
51:M5:150:TRP:CZ3	51:M5:156:HIS:CD2	3.01	0.48
52:M6:57:PHE:CE2	52:M6:72:HIS:HD2	2.43	0.48
55:M9:80:LYS:HE2	36:5:1940:G:OP1	206.80	0.48
56:N0:152:LEU:HA	56:N0:152:LEU:HD23	1.90	0.48
56:N0:23:LYS:HB3	56:N0:23:LYS:HE3	1.72	0.48
56:N0:41:TYR:O	56:N0:45:LEU:HB2	2.55	0.48
56:N0:24:LEU:HD22	56:N0:59:VAL:HG21	2.90	0.48
57:N1:75:ILE:HD13	57:N1:88:ARG:HD2	8.08	0.48
59:N3:81:GLN:HE21	59:N3:85:TRP:HE3	2.69	0.48
63:N7:84:ARG:N	66:O0:58:TYR:OH	3.01	0.48
74:O8:4:GLU:HG2	74:O8:5:ILE:H	2.46	0.48
75:O9:10:LYS:HA	75:O9:13:MET:HE3	2.38	0.48
2:S0:168:HIS:O	2:S0:172:LEU:HB2	2.41	0.48
2:S0:57:LEU:HG	2:S0:177:LEU:HD23	1.94	0.48
3:S1:127:VAL:HG22	3:S1:128:LYS:H	1.79	0.48
3:S1:160:HIS:O	3:S1:164:ILE:HG13	2.32	0.48
6:S4:49:ARG:HG3	6:S4:50:ASN:N	4.81	0.48
7:S5:156:ARG:NH1	7:S5:156:ARG:HB2	2.28	0.48
7:S5:42:LEU:HB2	7:S5:46:TRP:O	2.14	0.48
7:S5:77:TYR:HD2	7:S5:83:ARG:HG3	6.17	0.48
10:S8:171:SER:HB3	10:S8:180:ASP:HB2	2.39	0.48
35:SM:48:ARG:HE	36:1:1017:C:H5'	1.78	0.48
34:SR:172:ALA:HB2	34:SR:202:LEU:HD22	1.96	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:SR:294:TRP:CZ3	34:SR:301:LEU:HB2	2.49	0.48
36:1:1307:G:H1'	36:1:1308:A:C8	2.49	0.48
36:1:1764:U:H5''	55:M9:43:LYS:NZ	2.29	0.48
36:1:2442:G:H22	36:1:2505:U:H3	1.62	0.48
36:1:2986:U:H2'	36:1:2987:A:C8	2.49	0.48
36:1:3355:U:H3'	36:1:3356:G:H5''	1.96	0.48
36:1:348:A:H1'	36:1:352:A:N3	2.28	0.48
36:1:872:U:H2'	36:1:873:C:C6	2.48	0.48
1:2:1535:U:O4	7:S5:186:ASN:N	2.47	0.48
86:2:2043:OHX:N2	86:2:2098:OHX:N6	2.62	0.48
1:2:28:A:H2'	1:2:29:U:H6	1.76	0.48
1:2:447:U:O3'	6:S4:11:ARG:NH2	2.37	0.48
1:2:704:C:H3'	1:2:704:C:OP2	2.14	0.48
38:4:21:C:OP1	41:L4:193:LYS:HD2	2.14	0.48
38:4:36:G:C8	71:O5:86:ARG:HD2	2.49	0.48
36:5:1179:A:H5''	36:5:1180:A:OP2	2.13	0.48
70:O4:43:LYS:O	36:5:1653:G:H4'	186.49	0.48
36:5:2278:C:OP1	86:5:4085:OHX:N6	2.47	0.48
42:L5:23:ARG:NH2	36:5:2703:A:OP2	283.85	0.48
36:5:1116:G:N2	36:5:2817:A:O4'	2.47	0.48
36:5:2965:U:C5	36:5:2966:G:C5	3.02	0.48
36:5:3200:G:C6	36:5:3201:C:C4	3.02	0.48
86:5:3971:OHX:N2	86:5:4193:OHX:N5	2.62	0.48
1:6:1091:A:H4'	1:6:1092:A:O5'	2.14	0.48
1:6:1398:U:H4'	1:6:1399:C:OP2	2.14	0.48
1:6:1640:C:H6	1:6:1640:C:O5'	1.97	0.48
1:6:221:A:C2'	1:6:222:A:H5'	2.44	0.48
1:6:373:G:H2'	1:6:374:U:H6	1.78	0.48
1:6:541:A:OP1	1:6:541:A:H8	1.96	0.48
1:6:855:A:C2	1:6:857:U:H1'	2.49	0.48
13:C1:94:ILE:HD13	25:D3:16:ARG:HD3	4.28	0.48
19:C7:16:LEU:HD12	19:C7:54:THR:HG21	2.64	0.48
19:C7:17:ILE:HG23	19:C7:58:MET:HE2	3.66	0.48
26:D4:118:ILE:HD13	26:D4:125:LEU:HD22	4.62	0.48
27:D5:59:TYR:HE1	27:D5:100:ILE:HG23	6.36	0.48
29:D7:50:ALA:HB2	29:D7:71:ALA:HB2	1.96	0.48
39:L2:155:LYS:HE2	39:L2:253:GLN:HA	8.05	0.48
39:L2:204:MET:HB3	39:L2:208:ASP:HB2	2.80	0.48
40:L3:169:THR:HG23	40:L3:314:TYR:OH	2.13	0.48
40:L3:361:THR:HG23	40:L3:371:GLN:O	2.78	0.48
42:L5:140:ARG:HH21	36:5:1080:A:P	228.37	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:L5:148:ILE:HA	42:L5:148:ILE:HD12	4.50	0.48
43:L6:88:SER:N	43:L6:176:PHE:O	3.73	0.48
44:L7:38:LYS:HA	44:L7:38:LYS:HD2	4.47	0.48
45:L8:128:LYS:HG3	36:5:120:G:C5	98.24	0.48
47:M0:191:LYS:HB3	47:M0:213:PHE:CE2	2.49	0.48
48:M1:21:ILE:HG12	48:M1:125:MET:HB3	4.69	0.48
54:M8:80:THR:O	54:M8:137:THR:HA	2.80	0.48
55:M9:96:ILE:HG22	55:M9:100:ARG:HD2	3.97	0.48
63:N7:46:ILE:HG13	63:N7:46:ILE:O	2.13	0.48
67:O1:70:ARG:HE	67:O1:102:LYS:HZ3	4.29	0.48
69:O3:24:ASN:HD21	69:O3:27:VAL:HG23	2.19	0.48
70:O4:74:ARG:HG2	70:O4:75:ALA:H	3.40	0.48
74:O8:28:ASN:ND2	74:O8:40:GLN:OE1	4.37	0.48
75:O9:7:PHE:CE2	38:8:113:U:C4	98.47	0.48
2:S0:110:TYR:O	2:S0:112:THR:N	2.47	0.48
2:S0:147:THR:HB	2:S0:151:SER:HB2	2.89	0.48
2:S0:167:LYS:HG2	2:S0:168:HIS:CE1	2.48	0.48
3:S1:195:LYS:O	3:S1:199:ASN:N	2.39	0.48
5:S3:142:LEU:HD13	5:S3:182:LEU:HD11	1.96	0.48
6:S4:36:HIS:HD2	6:S4:83:PRO:O	2.97	0.48
7:S5:120:ILE:HG23	27:D5:59:TYR:CE1	2.49	0.48
7:S5:222:LYS:HA	7:S5:225:ARG:HD2	3.22	0.48
7:S5:41:LYS:HE2	7:S5:41:LYS:HB3	2.57	0.48
7:S5:84:LYS:HG2	7:S5:92:ARG:CZ	3.09	0.48
11:S9:127:VAL:O	11:S9:131:GLN:HB2	2.61	0.48
34:SR:128:ASP:N	34:SR:128:ASP:OD1	2.42	0.48
36:1:1103:A:N6	36:1:1363:A:O2'	2.47	0.47
36:1:1166:G:O6	86:1:3857:OHX:N4	2.47	0.47
36:1:1847:A:O2'	36:1:1848:G:H5''	2.14	0.47
36:1:2115:G:H22	36:1:2120:A:H1'	1.79	0.47
86:1:4027:OHX:N4	86:1:4040:OHX:N3	2.62	0.47
86:1:4050:OHX:N6	86:1:4160:OHX:N3	2.62	0.47
36:1:642:U:H2'	36:1:644:G:OP2	2.14	0.47
1:2:1752:U:OP2	86:2:2057:OHX:N2	2.46	0.47
1:2:1795:U:H3'	28:D6:5:ARG:NH1	2.29	0.47
1:2:1662:G:O6	86:2:2040:OHX:N3	2.47	0.47
1:2:111:U:C2	1:2:304:U:C4	3.01	0.47
37:3:115:G:H2'	37:3:116:C:H6	1.78	0.47
37:3:92:A:C4	37:3:93:C:H1'	2.49	0.47
36:5:2546:C:H2'	36:5:2547:A:C8	2.49	0.47
36:5:2951:G:H21	36:5:2952:G:H1'	1.79	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:2841:G:OP2	86:5:4133:OHX:N1	2.46	0.47
36:5:1790:G:O6	86:5:4192:OHX:N4	2.47	0.47
86:5:3971:OHX:N2	86:5:4193:OHX:N1	2.61	0.47
1:6:1679:G:C6	1:6:1680:G:N1	2.82	0.47
1:6:1708:U:H2'	1:6:1709:C:C6	2.48	0.47
1:6:296:U:H2'	1:6:297:U:C6	2.48	0.47
1:6:524:U:C2	1:6:526:A:OP2	2.67	0.47
1:6:53:G:H1	1:6:427:C:N4	2.11	0.47
37:7:64:A:H5'	37:7:65:G:H5''	1.96	0.47
71:O5:83:LYS:HA	38:8:38:U:C5	66.54	0.47
38:8:71:A:H4'	38:8:72:A:O5'	2.13	0.47
18:C6:69:VAL:HG11	18:C6:77:GLN:HB3	1.96	0.47
22:D0:95:ALA:HB1	22:D0:99:ILE:HG13	3.57	0.47
28:D6:44:ILE:HG12	28:D6:65:PRO:HD2	1.95	0.47
7:S5:57:SER:HB3	30:D8:53:ILE:HB	1.95	0.47
39:L2:130:SER:HG	39:L2:174:ARG:HH21	3.19	0.47
40:L3:221:THR:HG22	40:L3:222:LYS:O	2.14	0.47
40:L3:255:TRP:O	40:L3:255:TRP:HD1	1.95	0.47
40:L3:214:MET:HG3	40:L3:350:ALA:HB1	1.96	0.47
41:L4:10:SER:N	41:L4:13:GLY:O	3.12	0.47
44:L7:25:GLN:HG2	44:L7:29:GLU:HB2	1.94	0.47
46:L9:133:THR:HG23	46:L9:147:SER:O	3.08	0.47
36:1:2854:U:P	47:M0:3:ARG:HH22	2.37	0.47
48:M1:109:HIS:HD2	48:M1:114:ILE:HG21	1.79	0.47
48:M1:8:PRO:HD2	48:M1:10:ARG:HG3	1.96	0.47
48:M1:30:LEU:HD21	48:M1:66:ALA:HA	1.96	0.47
49:M3:104:ARG:C	72:O6:20:MET:HB2	2.35	0.47
49:M3:161:ASP:OD1	64:N8:139:ARG:NH1	3.70	0.47
51:M5:105:ARG:NH1	36:5:1545:A:N7	134.47	0.47
51:M5:49:ARG:HH11	51:M5:49:ARG:HB2	1.78	0.47
52:M6:147:TRP:NE1	52:M6:149:TYR:HB2	2.29	0.47
53:M7:151:THR:HG22	53:M7:152:GLU:O	2.49	0.47
55:M9:81:ARG:HD3	55:M9:88:ARG:HH12	4.11	0.47
58:N2:20:SER:O	58:N2:24:GLU:HG2	2.32	0.47
36:1:1901:A:O2'	59:N3:49:LEU:HD21	2.14	0.47
40:L3:67:PHE:CE1	59:N3:88:ARG:HB2	2.76	0.47
36:1:1608:C:H5''	61:N5:111:ASN:HD22	1.79	0.47
61:N5:58:ASP:O	61:N5:62:VAL:HG23	2.97	0.47
62:N6:5:SER:C	62:N6:7:ASP:H	3.31	0.47
63:N7:135:ARG:CZ	63:N7:135:ARG:HB3	4.79	0.47
67:O1:36:ILE:HD13	67:O1:59:ILE:HD11	1.94	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
68:O2:64:LYS:O	68:O2:65:PHE:HB2	2.13	0.47
69:O3:85:PHE:CD1	69:O3:89:LEU:HD21	3.21	0.47
70:O4:106:LYS:HD3	70:O4:109:THR:OG1	2.14	0.47
73:O7:52:LYS:HG2	73:O7:56:ARG:CZ	2.43	0.47
36:1:3118:C:H4'	76:Q0:106:ARG:HH12	1.79	0.47
76:Q0:96:CYS:SG	76:Q0:98:LYS:HB2	2.66	0.47
78:Q2:28:TYR:C	78:Q2:28:TYR:CD2	2.87	0.47
3:S1:184:LEU:HA	3:S1:187:LYS:HB2	1.96	0.47
4:S2:178:ILE:HG21	4:S2:185:LYS:HA	1.96	0.47
4:S2:42:GLY:HA2	4:S2:68:ILE:HD11	1.96	0.47
4:S2:65:GLU:HG3	4:S2:68:ILE:HD11	4.67	0.47
4:S2:75:GLY:O	4:S2:77:GLN:N	3.71	0.47
6:S4:35:PRO:HD2	6:S4:83:PRO:HG2	1.95	0.47
8:S6:7:TYR:CD1	8:S6:125:THR:HA	3.23	0.47
8:S6:64:LYS:O	8:S6:67:VAL:HG22	2.14	0.47
34:SR:89:LEU:O	34:SR:103:PHE:HD2	1.97	0.47
36:1:1307:G:H5'	52:M6:60:LYS:NZ	2.29	0.47
36:1:1633:C:H2'	36:1:1634:G:H8	1.79	0.47
36:1:1950:U:H3	36:1:2096:A:H2	1.62	0.47
36:1:2157:G:N2	36:1:2178:A:OP2	2.42	0.47
36:1:227:G:H2'	36:1:228:U:H6	1.79	0.47
36:1:2984:C:H2'	36:1:2985:C:H6	1.78	0.47
36:1:3026:G:O6	86:1:3932:OHX:N4	2.46	0.47
36:1:3356:G:H2'	36:1:3357:U:O4'	2.14	0.47
86:1:4076:OHX:N1	72:O6:28:TYR:O	2.47	0.47
36:1:612:U:O5'	43:L6:21:THR:HG21	2.14	0.47
1:2:226:A:H2'	1:2:227:U:H5'	1.97	0.47
1:2:820:U:H2'	1:2:821:U:H4'	1.96	0.47
38:4:124:G:OP2	86:4:232:OHX:N4	2.47	0.47
36:5:1074:U:O2'	36:5:1075:A:H2'	2.14	0.47
36:5:2107:A:C2	36:5:2108:C:C2	3.02	0.47
36:5:2215:A:H2'	36:5:2216:G:O4'	2.14	0.47
36:5:3170:A:H2'	36:5:3171:U:H5'	1.97	0.47
36:5:3323:A:H8	36:5:3323:A:O5'	1.97	0.47
1:6:1311:U:O4	86:6:2184:OHX:N4	2.47	0.47
1:6:1388:A:C6	1:6:1412:G:C6	3.03	0.47
31:D9:24:CYS:HB2	1:6:1434:U:H4'	410.59	0.47
1:6:1742:U:C2	1:6:1743:U:C6	3.03	0.47
1:6:992:A:O2'	1:6:1785:U:O2	2.32	0.47
11:S9:172:VAL:HG13	1:6:512:A:OP2	455.87	0.47
12:C0:28:ASN:N	12:C0:28:ASN:OD1	2.46	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:C2:48:SER:O	14:C2:52:LEU:HD23	2.14	0.47
16:C4:126:THR:O	16:C4:126:THR:OG1	2.29	0.47
25:D3:37:ALA:HB3	25:D3:38:PHE:CD2	4.01	0.47
16:C4:114:ARG:HA	28:D6:62:TYR:CZ	2.50	0.47
28:D6:82:ARG:HB2	28:D6:85:ARG:HE	9.08	0.47
22:D0:82:TYR:OH	31:D9:44:ARG:HG2	2.56	0.47
33:E1:103:LEU:HA	33:E1:103:LEU:HD23	1.67	0.47
41:L4:167:ALA:HA	41:L4:170:LYS:HB2	1.97	0.47
41:L4:179:LEU:CD2	41:L4:183:LYS:HG3	4.01	0.47
41:L4:5:GLN:HA	41:L4:20:LEU:O	2.13	0.47
43:L6:6:ALA:HA	68:O2:74:PHE:HE1	1.79	0.47
44:L7:112:ASN:O	44:L7:207:LEU:HB2	2.44	0.47
45:L8:238:LEU:HD23	45:L8:242:ALA:O	2.14	0.47
45:L8:45:ASN:HD21	45:L8:47:SER:HB3	1.79	0.47
46:L9:122:LYS:HG3	46:L9:123:ILE:N	2.84	0.47
51:M5:120:TRP:CZ2	51:M5:122:ASN:HA	2.49	0.47
51:M5:68:ARG:HH11	51:M5:128:LYS:HE3	2.87	0.47
53:M7:5:GLY:HA3	53:M7:116:HIS:CE1	2.50	0.47
53:M7:31:GLU:HG2	53:M7:60:PHE:HA	4.43	0.47
57:N1:34:TYR:CD1	57:N1:98:HIS:CE1	3.27	0.47
61:N5:132:ALA:CA	61:N5:135:ILE:HG22	4.31	0.47
63:N7:80:LEU:HD23	63:N7:80:LEU:HA	2.83	0.47
36:1:283:G:O2'	64:N8:59:ARG:NH1	2.46	0.47
70:O4:37:LYS:HG3	70:O4:58:ARG:NH2	5.53	0.47
71:O5:18:ALA:O	71:O5:22:VAL:HG23	2.14	0.47
77:Q1:9:ARG:HH11	77:Q1:9:ARG:CG	3.44	0.47
2:S0:179:ARG:CD	2:S0:183:ARG:HD2	2.44	0.47
5:S3:34:TYR:OH	5:S3:37:VAL:HG22	2.13	0.47
5:S3:68:GLU:OE2	12:C0:67:THR:HG23	2.14	0.47
7:S5:136:ALA:O	7:S5:140:THR:OG1	3.33	0.47
10:S8:11:ARG:HD3	10:S8:15:GLY:O	2.13	0.47
35:SM:35:ALA:HB1	35:SM:37:VAL:HG23	2.35	0.47
36:1:1949:G:OP1	55:M9:104:ARG:NH2	2.47	0.47
36:1:2661:G:H1	36:1:2709:C:H42	1.61	0.47
36:1:308:A:H5'	36:1:2223:A:O2'	2.14	0.47
36:1:3310:A:H2'	36:1:3311:C:H5'	1.96	0.47
1:2:1255:G:O6	14:C2:46:ARG:HD3	2.13	0.47
1:2:304:U:H2'	1:2:305:C:H6	1.78	0.47
1:2:477:A:H2'	1:2:478:A:C8	2.37	0.47
1:2:517:U:H3	1:2:535:A:H61	1.60	0.47
36:5:1170:A:O5'	36:5:1170:A:H8	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:1363:A:H2'	36:5:1364:C:C6	2.50	0.47
36:5:144:A:N6	36:5:145:G:C2	2.82	0.47
75:O9:2:ALA:HA	36:5:1491:A:H62	125.79	0.47
36:5:268:A:N1	36:5:295:A:H5'	2.29	0.47
36:5:2693:C:H1'	36:5:2706:G:H5''	1.95	0.47
36:5:3194:C:H42	36:5:3197:G:H1	1.62	0.47
36:5:597:G:H2'	36:5:598:A:H8	1.78	0.47
1:6:1312:A:O5'	1:6:1312:A:H8	1.97	0.47
1:6:1381:U:H1'	1:6:1516:A:N6	2.29	0.47
6:S4:248:ILE:HD12	1:6:789:A:H2	398.62	0.47
9:S7:111:LYS:HA	1:6:810:G:C2	341.69	0.47
12:C0:10:LYS:NZ	12:C0:36:ASP:HB3	2.63	0.47
17:C5:34:VAL:HG23	17:C5:41:VAL:HG12	1.95	0.47
18:C6:101:SER:O	18:C6:104:GLU:N	3.10	0.47
19:C7:27:ASP:OD1	34:SR:38:ARG:NH1	2.41	0.47
21:C9:63:ARG:O	21:C9:67:MET:HG3	5.16	0.47
2:S0:32:HIS:HE1	23:D1:87:ARG:HH12	1.62	0.47
25:D3:51:GLY:HA3	25:D3:76:LEU:HA	2.91	0.47
1:2:864:U:O2	29:D7:21:LEU:HB3	2.15	0.47
1:2:1597:A:C8	31:D9:14:TYR:HD2	2.32	0.47
39:L2:209:HIS:CD2	39:L2:210:PRO:N	2.83	0.47
39:L2:242:ARG:HD3	39:L2:246:LEU:HD21	1.96	0.47
39:L2:45:VAL:CG1	39:L2:86:GLN:HB3	2.44	0.47
40:L3:4:ARG:HB2	40:L3:4:ARG:HH11	5.05	0.47
41:L4:207:VAL:HG22	41:L4:249:ILE:HB	1.95	0.47
41:L4:40:THR:HG21	36:5:1426:C:H5'	130.21	0.47
42:L5:187:THR:OG1	42:L5:189:GLU:HB2	2.13	0.47
42:L5:289:LYS:O	42:L5:292:ALA:HB3	2.87	0.47
42:L5:45:ASN:O	42:L5:47:PRO:HD3	2.71	0.47
44:L7:127:LEU:HD22	44:L7:136:TYR:CE2	2.48	0.47
44:L7:224:ILE:HD13	56:N0:39:SER:HB2	1.95	0.47
46:L9:162:GLN:HG3	46:L9:163:GLN:N	2.45	0.47
47:M0:30:LYS:HA	47:M0:30:LYS:HE3	1.96	0.47
48:M1:110:ILE:O	48:M1:112:LEU:N	3.18	0.47
48:M1:38:GLU:O	48:M1:40:LEU:N	2.41	0.47
49:M3:178:LYS:HD3	49:M3:179:PHE:CZ	3.49	0.47
50:M4:99:TRP:O	50:M4:103:ILE:HG13	2.13	0.47
51:M5:184:LYS:HG2	51:M5:185:ALA:N	3.25	0.47
51:M5:80:THR:HG21	51:M5:86:ASN:O	2.14	0.47
51:M5:98:LEU:HD22	51:M5:128:LYS:HZ3	5.98	0.47
53:M7:60:PHE:CE2	53:M7:82:ARG:HB2	2.49	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:N0:115:ARG:N	56:N0:115:ARG:HD2	2.28	0.47
47:M0:162:GLN:HE22	56:N0:85:SER:HB2	1.78	0.47
57:N1:32:LYS:HE3	57:N1:98:HIS:HD2	7.12	0.47
60:N4:6:ASP:HB2	60:N4:11:ALA:HB3	3.53	0.47
61:N5:115:ARG:NH1	61:N5:119:THR:OG1	2.47	0.47
63:N7:51:LEU:HB3	63:N7:65:ARG:HD3	4.02	0.47
63:N7:76:ASN:OD1	63:N7:77:TYR:N	2.47	0.47
64:N8:28:HIS:HD2	36:5:936:A:OP1	161.32	0.47
65:N9:25:LYS:N	65:N9:25:LYS:HE3	2.29	0.47
66:O0:74:ASN:HB2	66:O0:86:ARG:HB2	3.29	0.47
36:1:1456:A:C6	67:O1:64:VAL:HG22	2.49	0.47
36:1:430:U:H4'	69:O3:67:MET:HE1	1.96	0.47
70:O4:102:LYS:HB3	70:O4:103:LYS:HE3	1.96	0.47
36:1:1590:G:OP1	70:O4:17:SER:OG	2.31	0.47
71:O5:4:VAL:CG2	71:O5:9:LEU:HD11	2.96	0.47
71:O5:53:CYS:O	71:O5:57:VAL:HG23	2.14	0.47
36:1:1925:U:O2	79:Q3:19:GLY:HA2	2.14	0.47
5:S3:134:CYS:N	5:S3:157:LEU:HD11	2.29	0.47
6:S4:176:ASP:HB2	6:S4:179:LYS:HE2	1.96	0.47
8:S6:174:LYS:HG3	1:6:79:C:H1'	342.98	0.47
9:S7:130:VAL:HG11	9:S7:154:LEU:HD21	3.24	0.47
9:S7:31:SER:HB3	9:S7:32:PRO:HD3	1.96	0.47
11:S9:178:ALA:O	11:S9:181:ALA:HB3	4.87	0.47
35:SM:30:THR:O	36:1:2666:C:H5''	2.14	0.47
34:SR:40:LYS:HA	34:SR:68:VAL:HG23	2.45	0.47
36:1:1595:U:O2'	36:1:1596:C:H5''	2.14	0.47
36:1:2232:A:H2'	36:1:2233:A:C8	2.49	0.47
36:1:2255:A:OP1	86:1:3926:OHX:N3	2.48	0.47
1:2:1015:U:H5''	1:2:1016:C:OP2	2.15	0.47
1:2:1393:C:H2'	1:2:1394:G:O4'	2.14	0.47
1:2:1407:U:H2'	1:2:1408:G:O4'	2.15	0.47
1:2:1440:C:H2'	1:2:1441:C:C6	2.50	0.47
1:2:89:G:C6	1:2:90:C:C4	3.03	0.47
1:2:93:A:C6	1:2:398:G:C6	3.02	0.47
42:L5:174:PRO:O	36:5:2747:A:H4'	247.01	0.47
36:5:94:G:H2'	36:5:95:A:C8	2.49	0.47
1:6:1325:A:H2'	1:6:1326:A:H8	1.79	0.47
10:S8:25:ARG:HB3	1:6:400:A:O5'	312.10	0.47
1:6:521:A:H2'	1:6:522:U:O4'	2.14	0.47
37:7:32:U:H4'	37:7:33:U:OP1	2.13	0.47
18:C6:78:VAL:O	18:C6:81:ILE:HG12	2.53	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:C8:29:VAL:O	20:C8:43:SER:OG	2.14	0.47
21:C9:14:PHE:CD2	21:C9:15:ILE:HD12	2.49	0.47
21:C9:40:SER:HB2	21:C9:96:ALA:HA	2.02	0.47
22:D0:59:PRO:HG3	1:6:1381:U:H4'	436.48	0.47
23:D1:71:ARG:HE	29:D7:4:VAL:HG11	2.41	0.47
23:D1:71:ARG:HG3	23:D1:83:TRP:CH2	2.50	0.47
24:D2:12:ASN:O	24:D2:16:ASN:N	2.86	0.47
15:C3:18:TYR:O	24:D2:56:HIS:HD2	1.97	0.47
26:D4:123:LYS:HZ3	26:D4:124:ARG:H	1.62	0.47
28:D6:91:ASP:OD1	28:D6:91:ASP:N	2.48	0.47
39:L2:15:ILE:HG22	39:L2:16:PHE:CD2	6.77	0.47
40:L3:140:ASP:OD2	40:L3:141:GLY:N	3.89	0.47
41:L4:208:VAL:HG11	41:L4:230:VAL:HG13	2.12	0.47
41:L4:23:PRO:HB3	41:L4:259:ASP:OD1	2.14	0.47
42:L5:109:THR:O	42:L5:113:LEU:HB2	2.14	0.47
45:L8:143:ILE:HD11	45:L8:151:VAL:HG11	2.62	0.47
45:L8:71:VAL:HG22	45:L8:76:ALA:HB2	2.94	0.47
46:L9:105:GLU:HA	46:L9:109:ALA:HB3	1.97	0.47
47:M0:76:MET:HE3	47:M0:148:VAL:HG13	1.96	0.47
48:M1:90:GLN:OE1	48:M1:172:LEU:HD21	2.91	0.47
50:M4:17:VAL:HG21	50:M4:74:ARG:HB2	2.03	0.47
50:M4:50:LYS:HE3	50:M4:86:ALA:HB2	1.95	0.47
36:1:1545:A:N7	51:M5:105:ARG:NH1	2.63	0.47
51:M5:83:LYS:O	51:M5:87:GLN:HG3	2.72	0.47
54:M8:66:ARG:HE	54:M8:143:PRO:HD3	2.61	0.47
55:M9:130:ASN:C	55:M9:132:PHE:H	2.18	0.47
55:M9:182:ASP:O	55:M9:184:LEU:N	4.10	0.47
59:N3:26:ALA:O	59:N3:115:THR:N	2.28	0.47
60:N4:4:GLU:HG3	60:N4:30:ARG:NH1	3.96	0.47
67:O1:29:ALA:HB3	67:O1:30:PRO:HD3	2.10	0.47
68:O2:20:HIS:O	68:O2:21:HIS:HB2	2.14	0.47
68:O2:24:ARG:HD3	68:O2:25:TYR:CZ	2.50	0.47
71:O5:45:LYS:O	71:O5:49:LYS:HG2	2.53	0.47
76:Q0:95:VAL:HG21	76:Q0:122:ARG:NH1	3.40	0.47
2:S0:50:VAL:HA	2:S0:53:THR:OG1	2.14	0.47
5:S3:4:LEU:O	5:S3:5:ILE:HD13	2.39	0.47
5:S3:76:ARG:HG3	5:S3:77:PHE:CD1	6.59	0.47
6:S4:66:MET:HB3	1:6:454:U:C4	377.07	0.47
7:S5:99:MET:HG3	7:S5:180:ARG:NH2	2.29	0.47
8:S6:31:ARG:HD2	8:S6:34:GLN:NE2	2.28	0.47
10:S8:184:LEU:HD21	10:S8:192:TYR:CD2	2.50	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:S8:61:GLU:HG2	10:S8:62:THR:HG23	3.48	0.47
36:1:1456:A:N1	36:1:1476:G:O2'	2.33	0.47
36:1:1496:C:C2	36:1:1521:G:N2	2.83	0.47
36:1:199:A:C4	36:1:201:A:C8	3.02	0.47
36:1:2854:U:OP1	47:M0:64:ALA:HB2	2.15	0.47
36:1:370:U:C4	36:1:371:G:C6	3.03	0.47
36:1:812:G:N7	86:1:3977:OHX:N1	2.62	0.47
36:1:92:G:H5'	36:1:93:C:C5'	2.45	0.47
1:2:335:U:O2'	13:C1:129:ARG:HD2	2.14	0.47
36:5:1284:C:O2'	36:5:1285:G:OP1	2.30	0.47
51:M5:56:LYS:NZ	36:5:150:A:OP1	87.20	0.47
36:5:176:G:H2'	36:5:177:U:C6	2.49	0.47
36:5:195:U:H2'	36:5:196:G:O4'	2.13	0.47
36:5:2725:U:H5''	36:5:2726:C:OP2	2.15	0.47
36:5:2806:U:H2'	36:5:2807:U:H6	1.80	0.47
36:5:3181:C:H2'	36:5:3182:G:H8	1.78	0.47
36:5:3358:U:H2'	36:5:3359:A:C8	2.49	0.47
17:C5:97:TYR:OH	1:6:1211:A:N3	380.22	0.47
1:6:1698:G:H1'	1:6:1699:G:OP1	2.14	0.47
25:D3:108:GLY:HA2	1:6:600:U:OP2	358.12	0.47
1:6:755:A:HO2'	1:6:756:A:P	2.34	0.47
37:7:85:G:O6	86:7:222:OHX:N5	2.47	0.47
86:8:216:OHX:N6	86:8:223:OHX:N4	2.62	0.47
15:C3:89:TYR:CE2	15:C3:150:VAL:HG13	2.49	0.47
16:C4:43:THR:OG1	16:C4:46:MET:HG3	2.15	0.47
17:C5:28:MET:HE2	17:C5:29:SER:N	2.30	0.47
18:C6:79:TYR:HA	18:C6:82:ARG:HD3	1.96	0.47
1:2:1389:C:H4'	19:C7:49:LYS:HA	1.97	0.47
22:D0:50:LEU:HD13	22:D0:95:ALA:HB2	4.09	0.47
24:D2:77:PRO:O	24:D2:79:PHE:N	2.47	0.47
39:L2:144:ASN:O	39:L2:160:SER:N	3.76	0.47
39:L2:70:ARG:CZ	39:L2:72:ARG:NE	5.53	0.47
40:L3:174:LYS:N	36:5:3314:A:OP1	203.57	0.47
41:L4:317:PRO:O	41:L4:324:LEU:HB2	2.40	0.47
42:L5:108:ARG:NH2	42:L5:251:PRO:O	3.15	0.47
43:L6:56:LYS:HB2	43:L6:98:VAL:CG1	2.44	0.47
43:L6:72:ASN:HB3	43:L6:160:SER:HA	2.34	0.47
43:L6:78:ARG:HD3	43:L6:106:PHE:CD2	2.49	0.47
44:L7:131:GLU:O	44:L7:229:PHE:HB2	2.14	0.47
45:L8:25:PRO:HB2	45:L8:26:LEU:HD12	1.97	0.47
50:M4:38:ILE:O	56:N0:95:ARG:NH2	3.21	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:M5:31:ARG:HG3	51:M5:129:TYR:OH	3.14	0.47
53:M7:141:SER:O	53:M7:143:PRO:HD3	2.28	0.47
53:M7:18:ARG:O	53:M7:94:LEU:HD11	4.15	0.47
55:M9:90:PRO:HG2	55:M9:93:VAL:CG2	3.26	0.47
57:N1:62:GLY:HA3	57:N1:76:ILE:HD13	2.25	0.47
58:N2:43:VAL:HB	58:N2:49:ASN:HB3	1.97	0.47
62:N6:36:SER:HB3	62:N6:106:ILE:O	2.13	0.47
62:N6:5:SER:O	62:N6:7:ASP:N	4.40	0.47
63:N7:115:LYS:O	63:N7:119:GLU:HB2	3.08	0.47
66:O0:13:LYS:HE3	66:O0:103:THR:HG21	2.69	0.47
67:O1:25:PHE:O	67:O1:28:ARG:N	2.74	0.47
68:O2:79:VAL:O	68:O2:83:GLU:HG3	3.47	0.47
69:O3:103:TYR:HA	69:O3:105:SER:N	2.30	0.47
76:Q0:114:LYS:HG2	76:Q0:115:CYS:N	3.25	0.47
2:S0:90:ALA:HB3	2:S0:97:PRO:HG3	1.96	0.47
9:S7:173:TYR:CE1	9:S7:181:ILE:HD11	3.86	0.47
11:S9:127:VAL:HG12	11:S9:131:GLN:NE2	2.29	0.47
36:1:142:C:H2'	36:1:143:G:O4'	2.15	0.47
36:1:1769:G:N7	86:1:4166:OHX:N2	2.62	0.47
36:1:198:A:C6	36:1:219:A:C6	3.02	0.47
36:1:421:G:C8	36:1:2365:C:C6	3.02	0.47
36:1:2884:C:H2'	36:1:2885:C:H6	1.80	0.47
36:1:2932:U:H2'	36:1:2933:A:H5''	1.97	0.47
36:1:3279:A:N6	36:1:3280:U:O4	2.48	0.47
36:1:3326:G:H2'	36:1:3327:G:C8	2.48	0.47
36:1:3335:A:C2	36:1:3336:A:C4	3.03	0.47
36:1:692:A:C4	36:1:693:A:C8	3.02	0.47
1:2:106:U:H2'	1:2:107:C:O4'	2.15	0.47
1:2:108:A:H2'	1:2:109:G:C8	2.49	0.47
1:2:1410:A:H2'	1:2:1411:A:O4'	2.14	0.47
1:2:247:A:O2'	1:2:248:U:H5'	2.14	0.47
1:2:47:A:N7	1:2:98:U:O2'	2.44	0.47
36:5:1020:G:H2'	36:5:1021:G:O4'	2.15	0.47
71:O5:95:PHE:CG	36:5:136:G:H5'	61.86	0.47
36:5:1434:G:OP1	36:5:1437:C:N4	2.48	0.47
62:N6:2:ALA:N	36:5:213:A:H5''	79.86	0.47
36:5:2882:U:H2'	36:5:2883:U:O4'	2.15	0.47
36:5:767:U:H1'	36:5:768:C:C6	2.48	0.47
36:5:40:A:C6	36:5:937:G:C2	3.02	0.47
36:5:982:C:N4	36:5:1101:G:H1	2.13	0.47
1:6:1263:G:H2'	1:6:1264:G:O4'	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:1614:A:C6	1:6:1615:C:C4	3.02	0.47
1:6:1314:U:OP1	86:6:2184:OHX:N1	2.48	0.47
1:6:246:G:C2	1:6:247:A:C4	3.03	0.47
16:C4:122:PRO:HB3	1:6:887:A:H1'	283.67	0.47
1:6:899:G:H2'	1:6:900:A:C8	2.49	0.47
13:C1:86:ILE:HD11	13:C1:125:VAL:HG11	4.29	0.47
13:C1:69:LYS:HB3	13:C1:71:LEU:HD21	3.27	0.47
15:C3:84:ILE:H	15:C3:84:ILE:HD13	4.56	0.47
17:C5:100:LYS:HB3	1:6:1183:A:C2	370.51	0.47
18:C6:58:ASP:OD2	18:C6:59:LYS:HD2	6.08	0.47
1:2:1389:C:O2'	19:C7:52:GLY:HA3	2.15	0.47
20:C8:45:LEU:HD11	21:C9:36:ILE:HG22	1.95	0.47
21:C9:125:SER:OG	21:C9:128:GLY:N	2.48	0.47
24:D2:94:LEU:HD11	24:D2:102:VAL:HG23	1.96	0.47
1:2:778:G:H22	26:D4:10:ARG:CZ	2.26	0.47
27:D5:43:ASP:O	27:D5:45:GLU:N	2.75	0.47
28:D6:30:ILE:HG13	28:D6:31:PRO:HD2	2.16	0.47
29:D7:63:LEU:HD23	29:D7:63:LEU:HA	1.80	0.47
40:L3:111:SER:O	40:L3:114:VAL:HG23	2.42	0.47
40:L3:92:TYR:HE1	40:L3:159:ARG:HD2	2.01	0.47
40:L3:199:PHE:C	40:L3:201:LYS:H	2.42	0.47
40:L3:284:ARG:HB3	40:L3:323:MET:HB3	2.48	0.47
40:L3:79:VAL:HG11	40:L3:338:LEU:HD21	1.96	0.47
43:L6:75:PRO:O	43:L6:77:ARG:HB2	3.55	0.47
44:L7:222:HIS:O	44:L7:227:GLY:N	2.36	0.47
45:L8:49:TYR:O	36:5:2523:A:H2'	170.67	0.47
46:L9:2:LYS:HZ1	46:L9:59:ASN:HD21	1.62	0.47
37:3:64:A:H5''	47:M0:206:LEU:H	1.79	0.47
36:1:2680:A:C2	48:M1:57:PHE:HB3	2.49	0.47
48:M1:57:PHE:HB3	36:5:2680:A:C2	309.31	0.47
49:M3:27:ASP:O	49:M3:31:LYS:HB2	3.02	0.47
51:M5:13:LYS:O	51:M5:19:LEU:HD22	2.80	0.47
52:M6:12:LYS:HD3	52:M6:37:ARG:NH2	2.30	0.47
53:M7:24:VAL:HG12	53:M7:86:LYS:CD	2.85	0.47
56:N0:87:THR:H	56:N0:88:HIS:CE1	3.82	0.47
67:O1:87:ASN:OD1	67:O1:88:PRO:HD2	2.14	0.47
69:O3:49:ILE:HD13	69:O3:100:ILE:HG13	2.85	0.47
77:Q1:14:LYS:O	77:Q1:18:ARG:HB2	2.14	0.47
2:S0:195:TRP:NE1	2:S0:197:ILE:HD13	4.96	0.47
3:S1:109:LYS:O	3:S1:112:SER:OG	2.22	0.47
3:S1:128:LYS:HZ2	3:S1:132:ASP:HB3	1.77	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:S1:129:THR:OG1	3:S1:131:ASP:OD1	6.78	0.47
5:S3:61:GLU:O	5:S3:63:GLY:N	2.47	0.47
6:S4:22:LYS:HD3	1:6:757:A:H4'	379.10	0.47
7:S5:114:ILE:O	7:S5:118:LEU:HD12	2.14	0.47
7:S5:203:LYS:HD2	7:S5:203:LYS:HA	1.83	0.47
8:S6:25:ARG:HG3	8:S6:28:PHE:CD1	2.50	0.47
8:S6:88:ARG:NH2	1:6:401:A:N1	322.98	0.47
8:S6:98:ARG:HD3	8:S6:99:GLY:N	2.32	0.47
11:S9:51:LYS:O	11:S9:54:ARG:HB3	2.15	0.47
36:1:1593:A:C6	36:1:1594:A:C6	3.02	0.47
36:1:1615:C:H2'	36:1:1616:U:C6	2.50	0.47
36:1:1826:C:H2'	36:1:1827:C:C6	2.49	0.47
36:1:2444:C:OP2	36:1:2445:A:H2'	2.14	0.47
36:1:2512:C:N4	36:1:2593:A:OP2	2.46	0.47
36:1:2943:G:C8	40:L3:2:SER:HB3	2.50	0.47
36:1:3346:U:H3	36:1:3359:A:N6	2.12	0.47
86:1:3965:OHX:N6	86:1:4153:OHX:N2	2.62	0.47
36:1:2754:G:OP2	86:1:4001:OHX:N6	2.47	0.47
1:2:411:C:H2'	1:2:412:A:O4'	2.14	0.47
1:2:452:A:H3'	1:2:453:U:C5	2.50	0.47
1:2:4:C:H2'	1:2:5:U:C6	2.49	0.47
1:2:626:U:H2'	1:2:627:C:H6	1.79	0.47
1:2:793:A:H5''	1:2:794:U:C5	2.50	0.47
37:3:91:G:H2'	37:3:92:A:H8	1.78	0.47
71:O5:95:PHE:CD2	36:5:136:G:H5'	63.13	0.47
36:5:1696:A:H2'	36:5:1697:A:C8	2.49	0.47
57:N1:49:GLN:HG2	36:5:2756:C:O4'	247.05	0.47
36:5:2787:G:OP2	86:5:4028:OHX:N6	2.48	0.47
40:L3:120:LYS:NZ	36:5:3001:C:OP1	203.58	0.47
36:5:3163:A:O2'	36:5:3164:C:H5'	2.15	0.47
36:5:702:C:O2'	36:5:788:C:H5''	2.14	0.47
1:6:1394:G:C4	1:6:1405:G:N2	2.83	0.47
1:6:1564:U:H2'	1:6:1565:C:H6	1.76	0.47
1:6:1651:A:H8	1:6:1651:A:O5'	1.97	0.47
1:6:221:A:O2'	1:6:222:A:H5'	2.15	0.47
15:C3:76:LYS:NZ	1:6:813:U:H5'	319.40	0.47
1:6:918:U:H2'	1:6:919:A:H8	1.80	0.47
37:7:110:G:C6	37:7:111:U:C4	3.03	0.47
13:C1:76:VAL:HG12	13:C1:85:VAL:O	2.15	0.47
15:C3:47:PRO:HB3	15:C3:71:ILE:HG22	1.97	0.47
19:C7:107:SER:O	19:C7:110:VAL:HG23	2.59	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:C9:79:LEU:HD23	21:C9:80:TYR:CE2	3.65	0.47
23:D1:79:LEU:HD13	23:D1:82:VAL:HG11	2.83	0.47
26:D4:54:ALA:O	26:D4:76:TYR:N	2.35	0.47
14:C2:46:ARG:HB2	33:E1:103:LEU:HD12	1.97	0.47
33:E1:127:GLY:C	33:E1:129:GLY:H	2.18	0.47
39:L2:34:TYR:CD1	39:L2:38:HIS:HD2	2.75	0.47
40:L3:114:VAL:HG13	40:L3:163:HIS:CG	2.74	0.47
41:L4:244:LEU:HD23	41:L4:244:LEU:HA	1.92	0.47
41:L4:6:VAL:HG21	41:L4:255:PHE:CZ	2.50	0.47
42:L5:51:LEU:N	42:L5:145:PHE:O	2.57	0.47
42:L5:261:THR:C	42:L5:263:GLU:H	3.19	0.47
45:L8:180:VAL:HG22	45:L8:181:LYS:H	2.68	0.47
50:M4:14:LEU:HA	50:M4:14:LEU:HD23	1.71	0.47
51:M5:148:TYR:O	51:M5:150:TRP:N	2.46	0.47
59:N3:28:ASN:ND2	59:N3:112:SER:HB2	2.27	0.47
61:N5:135:ILE:O	61:N5:135:ILE:HD13	3.40	0.47
63:N7:48:ARG:HB3	63:N7:69:LYS:HB3	2.07	0.47
64:N8:103:ASP:HB3	64:N8:106:ALA:HB3	1.94	0.47
65:N9:6:ASN:O	65:N9:7:HIS:HB2	2.14	0.47
65:N9:8:THR:HG23	65:N9:9:ALA:N	3.66	0.47
69:O3:60:ARG:CZ	69:O3:60:ARG:HB3	2.45	0.47
76:Q0:95:VAL:HG21	76:Q0:122:ARG:CZ	3.59	0.47
2:S0:10:THR:OG1	2:S0:12:GLU:OE2	2.29	0.47
2:S0:142:PRO:HG3	23:D1:32:VAL:HG22	2.32	0.47
2:S0:175:TYR:OH	2:S0:197:ILE:O	2.86	0.47
2:S0:27:ARG:HG2	2:S0:44:GLY:O	2.14	0.47
4:S2:54:GLU:OE1	23:D1:11:LEU:HD13	4.18	0.47
6:S4:104:ASP:HB3	6:S4:105:VAL:H	1.43	0.47
7:S5:225:ARG:NH2	30:D8:58:GLU:HB2	5.19	0.47
7:S5:73:THR:HG23	18:C6:114:ARG:HE	4.42	0.47
9:S7:117:THR:O	9:S7:120:ALA:N	2.47	0.47
9:S7:131:PHE:C	9:S7:133:THR:H	2.16	0.47
10:S8:171:SER:CB	10:S8:180:ASP:H	2.86	0.47
11:S9:168:ARG:NH1	11:S9:171:ARG:HH21	4.73	0.47
11:S9:171:ARG:NE	11:S9:174:ARG:HB2	4.47	0.47
34:SR:132:LYS:HB3	34:SR:140:CYS:SG	3.29	0.47
34:SR:23:LEU:HD22	34:SR:33:LEU:HD21	2.76	0.47
34:SR:67:ILE:O	34:SR:84:SER:OG	2.18	0.47
34:SR:69:GLN:HG2	34:SR:111:MET:SD	4.11	0.47
36:1:1094:U:H6	36:1:1094:U:H3'	1.80	0.47
36:1:1677:G:OP2	58:N2:103:TYR:OH	2.25	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:2904:U:H2'	36:1:2905:U:C6	2.50	0.47
36:1:3000:A:H2'	36:1:3001:C:C6	2.49	0.47
36:1:3045:G:O3'	40:L3:275:ARG:NH1	2.48	0.47
36:1:371:G:H4'	36:1:396:A:N1	2.30	0.47
86:1:3882:OHX:N5	57:N1:13:TYR:O	2.47	0.47
36:1:706:A:H4'	36:1:781:G:O2'	2.15	0.47
1:2:1202:A:H2'	1:2:1203:A:H5''	1.97	0.47
1:2:144:U:O2'	1:2:145:A:H5'	2.15	0.47
1:2:1541:G:C5	1:2:1542:G:C6	3.02	0.47
1:2:332:U:P	10:S8:56:ARG:HH22	2.38	0.47
1:2:48:G:O2'	1:2:49:C:H5'	2.15	0.47
1:2:755:A:H2'	1:2:756:A:O4'	2.14	0.47
1:2:763:G:C6	1:2:764:U:N3	2.83	0.47
1:2:868:G:H1	1:2:960:U:H3	1.61	0.47
48:M1:174:LYS:HE3	36:5:1016:C:H42	358.06	0.47
70:O4:7:PHE:CE2	36:5:1856:C:H1'	152.27	0.47
36:5:2288:G:H5''	36:5:2289:U:OP2	2.14	0.47
36:5:2621:G:H2'	36:5:2622:C:C6	2.50	0.47
36:5:3045:G:H1	36:5:3096:C:H42	1.63	0.47
36:5:668:G:OP1	86:5:4136:OHX:N1	2.47	0.47
36:5:646:A:H2'	36:5:647:A:O4'	2.14	0.47
36:5:818:C:N3	36:5:920:A:H5'	2.28	0.47
1:6:190:C:O2'	1:6:191:C:O5'	2.33	0.47
1:6:197:A:H2'	1:6:198:A:C8	2.50	0.47
1:6:515:A:C8	1:6:516:G:C8	3.03	0.47
13:C1:8:GLN:OE1	13:C1:14:GLN:N	2.46	0.47
17:C5:30:THR:O	17:C5:33:PHE:N	2.52	0.47
18:C6:14:LYS:HB3	18:C6:15:SER:H	1.46	0.47
26:D4:94:TYR:HB2	26:D4:96:LEU:HG	4.16	0.47
31:D9:21:CYS:HA	31:D9:30:LEU:HD21	2.60	0.47
39:L2:101:VAL:HB	39:L2:165:VAL:HB	2.51	0.47
39:L2:188:LYS:NZ	39:L2:189:TYR:OH	4.79	0.47
41:L4:181:VAL:HG12	41:L4:182:LEU:N	2.26	0.47
41:L4:326:ARG:O	44:L7:41:ARG:NH2	3.73	0.47
45:L8:238:LEU:HB3	45:L8:243:GLN:HG2	1.97	0.47
46:L9:146:LEU:HD12	46:L9:146:LEU:N	2.90	0.47
46:L9:96:HIS:O	46:L9:98:PRO:HD3	2.15	0.47
49:M3:59:ARG:NH1	49:M3:66:ASN:O	2.77	0.47
46:L9:20:ILE:HG13	50:M4:7:VAL:HG13	1.96	0.47
51:M5:135:VAL:O	51:M5:137:PRO:HD3	2.15	0.47
51:M5:35:VAL:O	51:M5:64:VAL:HA	2.21	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:M6:73:PHE:CD1	52:M6:78:ARG:HG2	2.55	0.47
56:N0:16:THR:HG23	56:N0:19:VAL:HB	1.96	0.47
41:L4:359:LEU:HA	56:N0:8:GLN:OE1	2.53	0.47
60:N4:54:LEU:H	60:N4:54:LEU:CD1	2.23	0.47
63:N7:66:THR:HG21	63:N7:123:GLN:HE21	5.12	0.47
65:N9:56:ALA:O	65:N9:59:LYS:HB3	2.15	0.47
66:O0:95:ALA:HB1	66:O0:98:SER:HB3	1.97	0.47
69:O3:39:GLN:O	69:O3:41:ALA:N	2.48	0.47
71:O5:73:LYS:HE2	71:O5:73:LYS:HB3	1.78	0.47
78:Q2:70:LEU:N	78:Q2:83:LEU:O	2.72	0.47
2:S0:144:ILE:HG23	2:S0:158:VAL:HG13	1.97	0.47
3:S1:27:LYS:HG2	3:S1:49:ASN:OD1	2.14	0.47
4:S2:158:THR:HG21	4:S2:221:THR:HG23	1.96	0.47
5:S3:61:GLU:C	5:S3:63:GLY:H	2.18	0.47
8:S6:158:ILE:HA	8:S6:158:ILE:HD12	1.81	0.47
8:S6:7:TYR:CE1	8:S6:125:THR:HG23	2.50	0.47
5:S3:222:VAL:HB	34:SR:192:PHE:HA	1.97	0.47
5:S3:216:PRO:HB2	34:SR:196:ASN:OD1	3.39	0.47
34:SR:269:TYR:CG	34:SR:270:LEU:N	2.81	0.47
34:SR:90:ARG:NH2	1:6:1341:A:H4'	457.37	0.47
36:1:201:A:H2'	36:1:202:G:H8	1.79	0.47
36:1:2287:C:C2	36:1:2298:U:O4'	2.68	0.47
36:1:2842:U:C4	36:1:2843:U:C4	3.02	0.47
36:1:3353:G:O2'	36:1:3354:U:OP1	2.33	0.47
36:1:929:A:H2'	36:1:930:U:C6	2.50	0.47
1:2:1000:C:N4	1:2:1003:A:OP2	2.47	0.47
1:2:1046:G:OP1	3:S1:157:GLN:NE2	2.48	0.47
1:2:1039:A:N7	1:2:1091:A:C5	2.83	0.47
1:2:1345:A:H2'	1:2:1348:A:H62	1.79	0.47
1:2:1776:A:H2'	1:2:1777:G:C8	2.50	0.47
1:2:386:G:N1	1:2:387:A:C6	2.83	0.47
62:N6:16:ARG:NH1	36:5:216:G:OP1	83.51	0.47
36:5:3286:G:H2'	36:5:3287:U:O4'	2.14	0.47
36:5:1078:U:O4	86:5:3992:OHX:N5	2.48	0.47
36:5:678:G:O6	86:5:4008:OHX:N5	2.48	0.47
1:6:1077:C:H2'	1:6:1078:C:C6	2.50	0.47
10:S8:2:GLY:HA2	1:6:1729:C:O2'	288.33	0.47
1:6:16:G:H2'	1:6:17:C:C6	2.50	0.47
1:6:485:A:C2	1:6:486:G:H1'	2.49	0.47
1:6:836:U:H2'	1:6:837:G:C8	2.49	0.47
37:7:114:U:H2'	37:7:115:G:C8	2.49	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:C1:124:THR:O	13:C1:140:VAL:HG12	2.15	0.47
13:C1:63:LEU:HD23	13:C1:63:LEU:HA	2.49	0.47
14:C2:94:ALA:HB1	14:C2:119:SER:H	1.79	0.47
18:C6:131:GLY:HA2	18:C6:138:PHE:CD1	3.30	0.47
18:C6:99:GLU:OE2	34:SR:60:SER:HB2	2.40	0.47
19:C7:44:LYS:O	19:C7:47:ARG:HB3	2.72	0.47
20:C8:133:VAL:O	20:C8:135:GLY:N	2.48	0.47
24:D2:24:GLN:NE2	29:D7:5:GLN:H	2.13	0.47
1:2:863:A:OP1	24:D2:57:ARG:NH2	2.46	0.47
25:D3:99:ASN:N	25:D3:99:ASN:OD1	3.85	0.47
29:D7:46:VAL:HG12	29:D7:47:PHE:H	2.12	0.47
39:L2:32:LEU:HD23	39:L2:163:ARG:CZ	2.43	0.47
40:L3:54:THR:OG1	40:L3:360:ASP:HB3	2.15	0.47
43:L6:76:LEU:HD11	43:L6:141:VAL:HG21	2.95	0.47
45:L8:239:GLY:O	45:L8:242:ALA:N	2.43	0.47
45:L8:75:ILE:HD11	51:M5:18:VAL:HG23	1.95	0.47
46:L9:32:GLY:O	46:L9:82:VAL:HG22	2.70	0.47
48:M1:142:LYS:HA	48:M1:142:LYS:HD3	2.75	0.47
49:M3:71:ALA:HA	49:M3:147:ILE:HD12	1.97	0.47
52:M6:68:ARG:HG2	52:M6:68:ARG:H	1.41	0.47
52:M6:85:ARG:HD3	52:M6:90:HIS:CD2	2.49	0.47
53:M7:26:PHE:CD1	53:M7:121:GLN:HG3	5.10	0.47
55:M9:81:ARG:HD2	55:M9:88:ARG:NH2	2.29	0.47
56:N0:2:ALA:HB3	56:N0:32:SER:HB3	1.97	0.47
57:N1:68:THR:HG23	57:N1:69:LYS:N	4.69	0.47
61:N5:101:GLU:HG2	61:N5:102:LEU:HD23	4.23	0.47
62:N6:125:LYS:O	62:N6:126:LEU:HG	2.15	0.47
63:N7:25:ILE:HA	63:N7:43:VAL:HG12	1.97	0.47
64:N8:12:ARG:NH1	36:5:1431:G:OP2	148.02	0.47
64:N8:70:LYS:N	64:N8:71:PRO:HD3	2.44	0.47
68:O2:100:ILE:HB	68:O2:105:ARG:NH1	2.30	0.47
73:O7:21:ARG:HD2	73:O7:37:CYS:SG	2.54	0.47
78:Q2:2:VAL:N	78:Q2:90:HIS:O	2.47	0.47
2:S0:49:ASN:ND2	2:S0:52:LYS:HE3	3.75	0.47
4:S2:121:VAL:O	4:S2:125:ILE:HG13	2.15	0.47
4:S2:81:MET:HB2	4:S2:101:VAL:O	2.13	0.47
5:S3:179:GLN:HE21	5:S3:179:GLN:C	2.18	0.47
5:S3:74:GLN:HA	5:S3:79:TYR:HB2	2.94	0.47
6:S4:105:VAL:HG11	6:S4:245:LYS:H	3.13	0.47
6:S4:208:VAL:HG11	6:S4:225:VAL:HG21	1.95	0.47
6:S4:85:GLY:O	6:S4:88:ASP:HB2	3.04	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:S7:162:ILE:O	9:S7:166:LEU:HD13	2.14	0.47
9:S7:40:PRO:HG2	9:S7:41:LEU:HD23	3.91	0.47
10:S8:39:GLY:CA	10:S8:61:GLU:HB3	2.41	0.47
34:SR:281:TYR:HB3	34:SR:285:ALA:HB3	2.31	0.47
34:SR:260:ILE:HD13	34:SR:292:LEU:HD13	1.97	0.47
34:SR:93:ASP:HB2	34:SR:100:TYR:CE1	2.49	0.47
36:1:1577:G:H2'	36:1:1578:C:O4'	2.15	0.47
36:1:2341:A:OP2	40:L3:247:ARG:NH2	2.47	0.47
36:1:2421:U:H2'	36:1:2422:C:O4'	2.15	0.47
36:1:2944:U:H1'	40:L3:251:CYS:SG	2.54	0.47
36:1:3216:G:C4	36:1:3259:U:C4	3.03	0.47
36:1:517:G:P	44:L7:60:ARG:HH22	2.38	0.47
36:1:563:U:H2'	36:1:564:G:C8	2.49	0.47
36:1:697:A:H2'	36:1:698:U:O4'	2.15	0.47
1:2:1130:G:OP2	86:2:2073:OHX:N2	2.48	0.47
1:2:121:U:H1'	6:S4:33:ALA:HB3	1.96	0.47
1:2:1402:G:H2'	1:2:1403:C:C6	2.49	0.47
1:2:623:A:OP1	86:2:2157:OHX:N1	2.48	0.47
1:2:481:A:H61	1:2:505:A:H62	1.62	0.47
1:2:68:A:OP1	8:S6:160:ARG:NH2	2.46	0.47
36:5:1237:G:H2'	36:5:1237:G:N3	2.29	0.47
36:5:1560:G:N2	36:5:1580:A:N7	2.62	0.47
36:5:1843:C:H2'	36:5:1844:C:C6	2.49	0.47
55:M9:56:THR:HG22	36:5:1872:C:O3'	154.06	0.47
36:5:3343:G:N2	36:5:3361:G:H2'	2.29	0.47
1:6:1182:U:O2	1:6:1184:A:H8	1.97	0.47
1:6:433:C:H5''	1:6:434:G:OP2	2.15	0.47
11:S9:2:PRO:HD2	1:6:461:G:OP1	360.74	0.47
14:C2:81:ASP:HA	14:C2:82:PRO:HD3	1.65	0.47
18:C6:103:ASN:HA	18:C6:106:LYS:HB2	3.43	0.47
18:C6:31:VAL:O	18:C6:33:GLY:N	2.59	0.47
18:C6:49:TYR:HB3	18:C6:53:LEU:HD11	1.97	0.47
19:C7:86:PRO:HB2	19:C7:88:VAL:H	6.21	0.47
21:C9:30:VAL:HG12	21:C9:54:PHE:CG	2.50	0.47
24:D2:6:VAL:HG22	24:D2:29:PRO:HD2	3.19	0.47
25:D3:131:SER:O	25:D3:135:LEU:HG	2.28	0.47
31:D9:41:GLN:N	31:D9:41:GLN:OE1	2.48	0.47
40:L3:262:TRP:N	52:M6:64:PHE:O	2.48	0.47
40:L3:53:MET:HE1	40:L3:327:CYS:HB2	1.96	0.47
41:L4:234:ASN:OD1	41:L4:236:LEU:HD12	2.14	0.47
47:M0:77:THR:C	47:M0:79:VAL:H	2.17	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:M0:99:ILE:HG13	47:M0:100:ASN:H	1.78	0.47
48:M1:23:VAL:HG12	48:M1:29:ARG:HD3	1.97	0.47
49:M3:80:VAL:HG11	49:M3:87:ALA:HB2	2.83	0.47
51:M5:31:ARG:HD3	51:M5:129:TYR:OH	2.15	0.47
51:M5:137:PRO:HG2	51:M5:138:GLN:NE2	3.22	0.47
51:M5:197:LEU:HG	51:M5:199:LEU:HD21	1.97	0.47
51:M5:68:ARG:NH1	51:M5:68:ARG:HG2	2.14	0.47
52:M6:58:LEU:HD12	52:M6:58:LEU:HA	2.27	0.47
36:1:1507:G:N7	53:M7:129:THR:HB	2.30	0.47
54:M8:133:LYS:HB2	54:M8:135:GLN:HE22	2.66	0.47
55:M9:145:ALA:O	55:M9:148:ASP:HB2	2.55	0.47
56:N0:45:LEU:HA	56:N0:45:LEU:HD22	1.56	0.47
59:N3:12:ARG:HG2	59:N3:13:ILE:N	2.84	0.47
59:N3:53:SER:N	59:N3:56:ASP:OD2	2.79	0.47
64:N8:3:SER:O	64:N8:6:THR:HG22	2.15	0.47
64:N8:47:LYS:HG3	64:N8:48:TYR:N	3.90	0.47
57:N1:88:ARG:HD3	65:N9:33:LYS:HZ1	5.39	0.47
66:O0:18:ILE:HD13	66:O0:81:VAL:HB	1.95	0.47
72:O6:43:LEU:HA	72:O6:43:LEU:HD22	2.16	0.47
75:O9:25:GLN:O	75:O9:28:ARG:HG3	3.06	0.47
77:Q1:8:LYS:HD3	77:Q1:12:ARG:HH21	1.80	0.47
8:S6:200:ALA:HA	8:S6:203:GLU:HG3	4.04	0.47
34:SR:117:LYS:H	34:SR:117:LYS:HD2	1.79	0.47
36:1:1069:C:H2'	36:1:1070:U:H6	1.80	0.47
36:1:1095:U:N3	57:N1:127:GLN:HG2	2.30	0.47
36:1:1301:A:H4'	36:1:1302:A:O5'	2.15	0.47
36:1:1794:G:H4'	39:L2:191:LEU:HD13	1.96	0.47
36:1:2380:U:C2	36:1:2381:G:C8	3.02	0.47
36:1:2380:U:H2'	36:1:2381:G:C8	2.50	0.47
36:1:2438:A:H2'	36:1:2439:A:C8	2.50	0.47
36:1:2776:C:H5'	36:1:2777:G:C2	2.50	0.47
36:1:2947:G:H4'	36:1:2947:G:OP2	2.14	0.47
36:1:325:A:H5''	36:1:326:U:OP2	2.15	0.47
36:1:2165:G:OP1	86:1:3994:OHX:N6	2.48	0.47
36:1:992:A:N1	36:1:993:G:C6	2.83	0.47
1:2:363:G:OP1	86:2:2077:OHX:N2	2.48	0.47
1:2:2:A:H5'	1:2:2:A:C8	2.49	0.47
1:2:344:A:H2'	1:2:345:U:C6	2.50	0.47
1:2:398:G:P	10:S8:47:ARG:HH12	2.36	0.47
1:2:482:U:H2'	1:2:483:A:C8	2.48	0.47
1:2:620:A:O2'	1:2:621:A:H5'	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:679:U:H2'	1:2:680:U:C6	2.49	0.47
36:5:112:U:O2'	36:5:113:C:OP2	2.30	0.47
36:5:1877:U:C5'	36:5:1878:G:H5'	2.45	0.47
36:5:2278:C:O2'	36:5:2279:A:H5''	2.15	0.47
36:5:2441:A:H61	36:5:2507:C:N4	2.12	0.47
79:Q3:62:LYS:NZ	36:5:2554:A:N7	218.78	0.47
36:5:313:A:C6	36:5:314:U:C4	3.02	0.47
88:5:4248:BLS:H111	88:5:4248:BLS:H81	1.73	0.47
16:C4:129:LYS:NZ	1:6:1009:U:OP2	282.17	0.47
1:6:370:A:H5''	1:6:371:G:OP2	2.15	0.47
1:6:492:A:C2'	1:6:493:U:H5''	2.44	0.47
1:6:74:U:O2	86:6:2192:OHX:N2	2.48	0.47
1:6:805:U:C2'	1:6:806:A:H5'	2.44	0.47
37:7:57:G:H3'	37:7:58:C:H6	1.80	0.47
14:C2:124:LYS:O	14:C2:126:TRP:N	2.44	0.47
15:C3:128:TYR:HA	15:C3:131:THR:HB	3.60	0.47
16:C4:49:LYS:HD3	16:C4:49:LYS:HA	1.70	0.47
20:C8:117:LYS:O	20:C8:119:ILE:N	2.48	0.47
22:D0:16:GLN:HB2	22:D0:17:GLN:NE2	2.30	0.47
23:D1:71:ARG:O	23:D1:75:ASN:ND2	2.47	0.47
27:D5:55:PRO:HG2	27:D5:56:THR:HG23	1.97	0.47
27:D5:66:VAL:HA	27:D5:70:LYS:O	2.15	0.47
1:2:1153:G:OP1	28:D6:85:ARG:NH1	2.47	0.47
30:D8:27:GLN:HE22	30:D8:64:ARG:NH1	4.90	0.47
32:E0:37:ARG:O	32:E0:41:THR:HG23	2.15	0.47
40:L3:218:ILE:CG1	40:L3:276:THR:HG23	3.00	0.47
40:L3:84:VAL:CG2	40:L3:162:VAL:HB	2.45	0.47
42:L5:269:SER:O	42:L5:272:TYR:HD2	3.33	0.47
42:L5:58:LYS:HD2	42:L5:93:THR:HG21	1.97	0.47
44:L7:106:LEU:HA	44:L7:106:LEU:HD23	1.83	0.47
36:1:1334:U:H5''	44:L7:206:LYS:HB3	1.97	0.47
46:L9:17:THR:OG1	46:L9:17:THR:O	2.87	0.47
46:L9:41:ILE:HD13	46:L9:41:ILE:O	2.15	0.47
47:M0:129:VAL:HG13	47:M0:133:GLN:HG2	2.48	0.47
49:M3:3:ILE:HG12	64:N8:34:MET:HE2	2.05	0.47
49:M3:89:TYR:CE1	49:M3:93:ILE:HG13	2.50	0.47
52:M6:195:ALA:O	52:M6:197:LEU:N	2.85	0.47
55:M9:166:ASN:ND2	55:M9:167:ARG:HG2	6.21	0.47
41:L4:362:ASP:N	56:N0:26:ARG:HH12	3.94	0.47
56:N0:46:GLN:HG3	56:N0:51:VAL:O	2.14	0.47
57:N1:17:ARG:HG2	36:5:2700:G:H5''	266.54	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:N1:34:TYR:HE1	57:N1:98:HIS:CD2	3.36	0.47
59:N3:80:ARG:HH12	59:N3:116:GLY:HA3	2.61	0.47
59:N3:74:MET:HG3	59:N3:102:ILE:CD1	2.41	0.47
59:N3:89:ASP:OD1	59:N3:91:VAL:HG13	2.15	0.47
68:O2:86:THR:HG23	68:O2:115:LEU:HD22	2.93	0.47
70:O4:6:THR:HG22	36:5:1486:G:N2	146.69	0.47
71:O5:85:THR:O	71:O5:89:ARG:HB2	2.14	0.47
72:O6:58:ILE:HA	72:O6:61:ILE:HB	2.87	0.47
74:O8:41:THR:HG21	74:O8:62:ALA:HB2	1.97	0.47
75:O9:2:ALA:HA	36:5:1491:A:N6	125.57	0.47
79:Q3:2:ALA:HB2	36:5:853:G:N7	252.30	0.47
2:S0:162:CYS:HB3	2:S0:173:ILE:HG13	1.97	0.47
3:S1:157:GLN:H	3:S1:160:HIS:HB2	1.79	0.47
3:S1:189:ILE:HG13	3:S1:189:ILE:H	2.15	0.47
6:S4:181:VAL:O	6:S4:192:ILE:HA	2.45	0.47
6:S4:71:LYS:HD2	6:S4:74:GLY:O	4.44	0.47
6:S4:72:VAL:HB	6:S4:77:ARG:HG3	3.91	0.47
8:S6:28:PHE:CZ	8:S6:104:PRO:HG3	2.49	0.47
8:S6:76:LEU:HD23	8:S6:76:LEU:HA	1.71	0.47
9:S7:35:LYS:HZ2	9:S7:36:ALA:H	1.63	0.47
11:S9:175:ARG:HD3	11:S9:179:ARG:CZ	2.45	0.47
20:C8:145:ARG:HD3	35:SM:68:ARG:NH2	3.30	0.47
36:1:1605:A:O2'	36:1:1607:U:OP2	2.25	0.46
36:1:1743:G:C4	36:1:1744:G:C8	3.03	0.46
36:1:2108:C:O2'	36:1:3362:A:N6	2.48	0.46
36:1:2218:G:H2'	36:1:2219:A:H8	1.79	0.46
36:1:2747:A:OP1	42:L5:176:SER:OG	2.30	0.46
36:1:3006:A:C2	36:1:3141:A:C4	3.03	0.46
36:1:3191:G:C2	36:1:3202:G:C2	3.03	0.46
36:1:86:G:O2'	36:1:98:G:O6	2.27	0.46
1:2:1536:G:C5	1:2:1538:U:H1'	2.50	0.46
1:2:1558:U:H3'	1:2:1559:A:H4'	1.97	0.46
1:2:1682:U:O2'	1:2:1683:C:H5'	2.15	0.46
1:2:1773:C:H2'	1:2:1774:G:H8	1.77	0.46
1:2:542:A:H8	1:2:543:C:H3'	1.78	0.46
1:2:720:G:H1'	1:2:721:U:H5''	1.98	0.46
38:4:62:C:H4'	38:4:63:G:O5'	2.16	0.46
38:4:99:C:OP1	61:N5:53:HIS:NE2	2.48	0.46
57:N1:130:ARG:HD3	36:5:1098:A:OP2	254.57	0.46
36:5:2124:G:C2	36:5:2330:C:C2	3.03	0.46
36:5:2128:C:OP1	86:5:4085:OHX:N3	2.48	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:2236:G:O5'	36:5:2236:G:H8	1.97	0.46
36:5:287:G:H2'	36:5:288:C:C6	2.49	0.46
36:5:3151:U:H4'	36:5:3294:A:H1'	1.96	0.46
36:5:3170:A:C2'	36:5:3171:U:H5'	2.44	0.46
36:5:549:U:H2'	36:5:550:A:H8	1.80	0.46
36:5:987:U:H2'	36:5:988:U:C6	2.50	0.46
1:6:219:A:C6	1:6:843:U:H1'	2.51	0.46
1:6:479:C:O2	1:6:510:G:N2	2.48	0.46
16:C4:43:THR:OG1	1:6:900:A:OP1	278.62	0.46
86:5:4087:OHX:N1	38:8:18:U:OP1	2.48	0.46
12:C0:15:LEU:HG	12:C0:68:LEU:HD22	1.97	0.46
15:C3:99:ARG:O	15:C3:103:GLU:HG2	2.15	0.46
1:2:887:A:H1'	16:C4:122:PRO:HB3	1.97	0.46
18:C6:113:ASP:CG	18:C6:115:THR:H	2.18	0.46
20:C8:117:LYS:C	20:C8:119:ILE:H	2.18	0.46
24:D2:71:LYS:NZ	1:6:1099:U:H5''	375.22	0.46
25:D3:137:LYS:O	25:D3:139:LYS:N	4.66	0.46
39:L2:90:ALA:CB	39:L2:101:VAL:HG13	3.64	0.46
40:L3:137:TYR:CE2	40:L3:144:ILE:HG13	2.49	0.46
8:S6:22:HIS:CG	40:L3:300:ARG:HH12	2.33	0.46
40:L3:43:LEU:HD22	40:L3:203:VAL:HG11	1.96	0.46
42:L5:222:LEU:HG	42:L5:222:LEU:H	1.46	0.46
42:L5:86:TYR:CD1	42:L5:247:ILE:HA	2.72	0.46
42:L5:51:LEU:HB2	42:L5:144:VAL:HG13	2.49	0.46
43:L6:46:ARG:NH2	36:5:3268:A:OP1	244.40	0.46
45:L8:36:ILE:HG22	45:L8:37:GLY:N	2.25	0.46
47:M0:200:LEU:HB2	47:M0:213:PHE:CD2	3.34	0.46
47:M0:216:TYR:CD2	47:M0:217:PHE:N	2.83	0.46
49:M3:156:ALA:HA	64:N8:101:VAL:HG23	2.99	0.46
50:M4:94:TRP:CE2	50:M4:100:ALA:HB2	2.68	0.46
52:M6:124:LEU:O	52:M6:128:ARG:HB2	2.15	0.46
52:M6:65:ASN:OD1	52:M6:67:THR:HB	2.66	0.46
52:M6:58:LEU:HA	52:M6:72:HIS:CD2	2.50	0.46
53:M7:125:GLN:HB2	53:M7:141:SER:HB2	2.16	0.46
1:2:850:A:OP1	55:M9:162:ARG:HG2	2.16	0.46
56:N0:114:HIS:CE1	36:5:1212:A:H1'	310.57	0.46
57:N1:34:TYR:HD1	57:N1:98:HIS:CE1	3.34	0.46
58:N2:33:TYR:CE2	58:N2:63:VAL:HG21	2.54	0.46
58:N2:90:ARG:HH11	58:N2:90:ARG:HB3	4.53	0.46
64:N8:104:THR:O	64:N8:109:TYR:HB2	2.75	0.46
66:O0:61:MET:HG3	66:O0:62:LEU:N	2.29	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
72:O6:70:ARG:HD3	72:O6:84:LYS:HG3	2.24	0.46
2:S0:71:GLU:O	2:S0:96:THR:HG22	2.82	0.46
3:S1:219:LYS:HE2	3:S1:219:LYS:HA	1.96	0.46
4:S2:107:SER:HA	4:S2:190:LEU:O	2.92	0.46
5:S3:134:CYS:SG	5:S3:135:GLU:N	2.88	0.46
9:S7:131:PHE:HB3	9:S7:132:PRO:HD3	1.97	0.46
9:S7:156:SER:OG	9:S7:186:PRO:HG2	2.14	0.46
34:SR:23:LEU:HD12	34:SR:291:SER:O	4.14	0.46
36:1:1635:G:N2	36:1:1638:A:OP2	2.43	0.46
36:1:1701:C:H2'	36:1:1702:U:O4'	2.14	0.46
1:2:1187:U:O2'	1:2:1188:G:H5'	2.15	0.46
1:2:1573:A:H5'	1:2:1574:G:N2	2.31	0.46
1:2:237:C:H4'	1:2:238:U:C6	2.50	0.46
1:2:333:A:C6	1:2:334:G:C6	3.04	0.46
1:2:755:A:HO2'	1:2:756:A:P	2.38	0.46
1:2:986:G:H2'	1:2:987:G:O4'	2.15	0.46
37:3:4:U:H2'	37:3:5:G:C8	2.50	0.46
36:5:1074:U:O3'	36:5:1075:A:H8	1.98	0.46
36:5:1773:C:H2'	36:5:1774:C:H6	1.79	0.46
36:5:290:G:H2'	36:5:291:C:H6	1.80	0.46
36:5:3337:G:H2'	36:5:3338:C:H6	1.80	0.46
36:5:2187:G:OP2	86:5:3965:OHX:N4	2.48	0.46
1:6:1220:C:H6	1:6:1220:C:OP2	1.98	0.46
1:6:1541:G:C6	1:6:1542:G:C6	3.04	0.46
1:6:212:U:OP2	86:6:2125:OHX:N1	2.49	0.46
1:6:217:A:O2'	1:6:218:A:H8	1.98	0.46
1:6:993:A:H5''	1:6:994:G:OP2	2.14	0.46
73:O7:70:VAL:HG11	38:8:35:C:H5'	71.37	0.46
15:C3:101:HIS:CE1	15:C3:105:ASN:HB2	4.75	0.46
15:C3:23:PRO:HD2	15:C3:26:PHE:HB3	1.98	0.46
17:C5:77:ARG:HB3	17:C5:102:PHE:CE1	3.50	0.46
20:C8:15:LEU:HD22	20:C8:22:VAL:O	5.12	0.46
23:D1:62:ARG:HB3	23:D1:62:ARG:HH11	1.80	0.46
24:D2:5:SER:HB3	24:D2:8:ALA:HB2	2.22	0.46
28:D6:30:ILE:HD11	28:D6:34:LYS:O	2.15	0.46
28:D6:8:ASN:OD1	28:D6:8:ASN:N	2.84	0.46
29:D7:61:THR:O	29:D7:62:ILE:HB	2.15	0.46
25:D3:70:LYS:HE3	32:E0:8:LEU:HA	2.81	0.46
33:E1:91:ILE:HG12	33:E1:92:LYS:N	2.31	0.46
39:L2:190:ARG:HG2	39:L2:191:LEU:HD12	1.96	0.46
36:1:3304:U:O3'	40:L3:334:ARG:NH2	2.48	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
46:L9:26:LYS:HA	46:L9:35:THR:HG22	1.97	0.46
48:M1:18:VAL:HG22	48:M1:70:THR:HG23	3.70	0.46
49:M3:3:ILE:HD13	64:N8:45:MET:HE3	4.22	0.46
50:M4:45:LEU:HA	50:M4:57:ALA:HA	2.44	0.46
51:M5:150:TRP:CH2	51:M5:151:ILE:HG12	2.50	0.46
51:M5:54:LYS:O	51:M5:56:LYS:N	3.22	0.46
36:1:291:C:OP1	51:M5:68:ARG:HB3	2.15	0.46
36:1:289:A:O2'	51:M5:93:LYS:O	2.32	0.46
52:M6:23:VAL:HG13	52:M6:33:ILE:HG21	1.96	0.46
53:M7:155:GLU:CD	53:M7:155:GLU:H	4.12	0.46
55:M9:127:SER:C	55:M9:129:GLY:H	2.17	0.46
58:N2:51:GLY:C	58:N2:52:ASN:HD22	2.18	0.46
61:N5:75:LYS:HD3	61:N5:123:TYR:CE1	2.48	0.46
63:N7:101:PHE:HB3	63:N7:107:ARG:HH21	1.81	0.46
63:N7:5:LEU:HD21	63:N7:82:PRO:HB3	1.97	0.46
64:N8:133:LEU:HD22	64:N8:133:LEU:HA	2.09	0.46
66:O0:102:THR:O	66:O0:102:THR:OG1	2.34	0.46
66:O0:17:VAL:HG13	66:O0:95:ALA:HA	1.97	0.46
71:O5:85:THR:HG22	71:O5:87:ALA:HB3	2.69	0.46
72:O6:37:THR:O	72:O6:41:ARG:HB2	2.15	0.46
2:S0:6:THR:C	2:S0:8:ASP:H	2.18	0.46
5:S3:177:MET:HG3	5:S3:182:LEU:HD12	1.96	0.46
7:S5:209:TYR:O	7:S5:213:LYS:HG2	2.15	0.46
8:S6:73:ILE:HD12	8:S6:75:LEU:HD21	1.98	0.46
9:S7:143:LEU:HA	9:S7:143:LEU:HD22	2.86	0.46
10:S8:82:VAL:CG1	10:S8:101:ILE:HG22	3.06	0.46
34:SR:203:THR:HG21	34:SR:244:ALA:H	1.79	0.46
34:SR:216:LYS:HA	34:SR:239:GLU:CG	2.48	0.46
36:1:1062:A:H4'	57:N1:105:PHE:CE2	2.51	0.46
36:1:1444:G:C6	36:1:1445:U:C2	3.03	0.46
36:1:1820:U:H1'	36:1:1821:U:OP2	2.14	0.46
36:1:2307:G:H4'	36:1:2308:C:OP2	2.15	0.46
36:1:2378:C:H2'	36:1:2379:U:C6	2.51	0.46
36:1:3043:C:H2'	36:1:3044:G:O4'	2.15	0.46
36:1:3118:C:H4'	76:Q0:106:ARG:NH2	2.30	0.46
86:1:4027:OHX:N2	86:1:4040:OHX:N1	2.64	0.46
36:1:629:U:H2'	36:1:630:A:C8	2.51	0.46
36:1:660:A:H5''	41:L4:100:PHE:CD1	2.50	0.46
1:2:1368:G:C5	1:2:1369:U:C4	3.03	0.46
1:2:1518:C:OP1	86:2:2120:OHX:N5	2.48	0.46
1:2:538:A:H8	1:2:543:C:N4	2.12	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:753:A:H4'	6:S4:221:ARG:HE	1.79	0.46
1:2:79:C:H4'	8:S6:173:PRO:O	2.15	0.46
1:2:820:U:H3'	1:2:821:U:H5''	1.96	0.46
1:2:967:A:H2'	1:2:968:U:O4'	2.15	0.46
36:5:1025:A:H2'	36:5:1025:A:N3	2.31	0.46
36:5:1448:U:O2'	36:5:1449:A:H5'	2.15	0.46
36:5:1631:C:H5''	36:5:1632:A:C5'	2.45	0.46
36:5:1877:U:OP2	86:5:3950:OHX:N1	2.48	0.46
36:5:198:A:C6	36:5:219:A:C5	3.04	0.46
36:5:173:G:N1	36:5:246:U:C2	2.83	0.46
54:M8:181:SER:HB3	36:5:2790:A:OP2	183.57	0.46
36:5:415:G:H2'	36:5:416:A:H8	1.79	0.46
36:5:1814:A:OP1	86:5:4175:OHX:N3	2.48	0.46
1:6:1185:U:C2	1:6:1458:G:N7	2.83	0.46
21:C9:79:LEU:HD13	1:6:1523:G:H8	406.25	0.46
1:6:52:U:H2'	1:6:53:G:H8	1.80	0.46
1:6:109:G:O2'	1:6:796:A:N1	2.42	0.46
37:7:4:U:H2'	37:7:5:G:H8	1.80	0.46
38:8:16:G:O6	86:8:215:OHX:N6	2.48	0.46
13:C1:56:LYS:HG3	13:C1:56:LYS:H	1.59	0.46
18:C6:79:TYR:O	18:C6:82:ARG:HG2	2.38	0.46
20:C8:64:GLU:C	20:C8:66:LEU:H	3.12	0.46
21:C9:74:GLY:O	21:C9:77:ASN:N	2.73	0.46
22:D0:101:LYS:HA	22:D0:101:LYS:HD3	4.55	0.46
25:D3:54:LEU:HD21	25:D3:75:GLN:HG3	1.97	0.46
28:D6:82:ARG:HH21	1:6:1152:A:H5''	330.29	0.46
31:D9:5:ASN:CG	31:D9:7:TRP:HE1	2.17	0.46
32:E0:43:ARG:HB3	32:E0:44:PHE:CD2	2.50	0.46
39:L2:52:SER:HB3	39:L2:191:LEU:HD12	5.81	0.46
40:L3:356:LEU:HG	40:L3:356:LEU:H	1.82	0.46
36:1:1361:U:OP1	41:L4:309:ARG:HG2	2.15	0.46
46:L9:48:VAL:HG13	46:L9:52:LEU:HB3	1.96	0.46
50:M4:20:VAL:HG22	50:M4:68:LEU:HB2	2.50	0.46
50:M4:93:LYS:HE3	50:M4:93:LYS:HB2	1.71	0.46
51:M5:190:THR:O	51:M5:193:ARG:HB3	2.38	0.46
52:M6:54:TYR:CE2	52:M6:145:VAL:HG11	3.37	0.46
58:N2:83:TYR:O	58:N2:86:LYS:N	2.38	0.46
59:N3:22:ILE:HD13	59:N3:35:TYR:HA	1.97	0.46
63:N7:81:LEU:HD22	63:N7:81:LEU:HA	1.83	0.46
72:O6:51:SER:HB2	72:O6:52:PRO:HD2	1.98	0.46
73:O7:67:LEU:C	73:O7:69:HIS:H	2.90	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
74:O8:15:THR:HB	74:O8:70:PRO:HG2	1.97	0.46
36:1:1834:U:OP1	75:O9:5:LYS:HE3	2.15	0.46
3:S1:70:LEU:HD11	3:S1:79:HIS:ND1	2.30	0.46
4:S2:147:ASN:O	4:S2:149:GLY:N	3.86	0.46
4:S2:66:PHE:CE2	4:S2:67:GLN:HG3	2.51	0.46
5:S3:195:SER:O	5:S3:196:ARG:HG2	2.15	0.46
7:S5:43:PHE:H	7:S5:46:TRP:H	2.48	0.46
8:S6:1:MET:N	8:S6:18:ILE:O	2.86	0.46
9:S7:160:GLN:HA	9:S7:163:ASP:OD2	2.32	0.46
9:S7:48:GLU:HG2	9:S7:56:LYS:HD3	2.86	0.46
34:SR:83:ALA:HB1	34:SR:110:VAL:HG12	1.98	0.46
36:1:155:G:H5'	36:1:156:G:C8	2.51	0.46
36:1:217:U:O2	62:N6:103:LYS:NZ	2.45	0.46
36:1:2565:U:H2'	36:1:2566:C:C6	2.49	0.46
36:1:3087:A:H2'	36:1:3088:G:C8	2.51	0.46
36:1:3153:U:H1'	36:1:3158:G:C5	2.50	0.46
86:1:3988:OHX:N5	37:3:86:U:O2	2.49	0.46
36:1:992:A:N6	36:1:993:G:O6	2.48	0.46
1:2:1009:U:OP1	16:C4:129:LYS:NZ	2.48	0.46
1:2:1252:C:N4	33:E1:97:LYS:HE3	2.30	0.46
1:2:1207:C:H42	1:2:1456:C:H41	1.64	0.46
86:2:2043:OHX:N4	86:2:2098:OHX:N6	2.63	0.46
1:2:583:C:OP1	86:2:2026:OHX:N3	2.49	0.46
1:2:838:G:C6	1:2:839:U:C4	3.04	0.46
36:5:2186:U:H2'	36:5:2187:G:O4'	2.16	0.46
36:5:3288:G:O2'	36:5:3289:G:OP2	2.27	0.46
36:5:2264:U:OP1	86:5:3949:OHX:N4	2.49	0.46
36:5:3192:U:O4	86:5:4139:OHX:N6	2.48	0.46
1:6:1322:A:H2'	1:6:1323:C:C6	2.50	0.46
21:C9:69:LYS:NZ	1:6:1368:G:OP1	436.98	0.46
1:6:542:A:H1'	1:6:543:C:H5'	1.97	0.46
1:6:763:G:H2'	1:6:764:U:C6	2.50	0.46
1:6:886:U:C2	1:6:887:A:C8	3.04	0.46
13:C1:75:VAL:HG12	13:C1:120:GLY:N	2.29	0.46
15:C3:34:ILE:HA	15:C3:37:ILE:HD12	3.58	0.46
15:C3:88:LEU:HD22	15:C3:92:ILE:HD11	1.97	0.46
19:C7:84:TYR:O	19:C7:86:PRO:HD3	2.15	0.46
20:C8:80:LYS:NZ	20:C8:80:LYS:HA	2.30	0.46
24:D2:15:ASN:O	24:D2:19:LYS:HG3	3.26	0.46
39:L2:86:GLN:HG2	39:L2:88:ILE:CD1	3.09	0.46
40:L3:205:VAL:C	40:L3:207:SER:H	2.28	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:L3:244:ARG:NH1	40:L3:244:ARG:HB3	2.61	0.46
40:L3:320:ASP:N	40:L3:320:ASP:OD2	2.31	0.46
41:L4:150:LEU:HB3	41:L4:249:ILE:HG23	1.98	0.46
36:1:210:U:OP1	41:L4:161:LYS:HD2	2.16	0.46
43:L6:19:LYS:HB3	43:L6:19:LYS:HE3	1.70	0.46
43:L6:52:VAL:HG13	43:L6:65:ILE:HG23	4.55	0.46
44:L7:77:VAL:HG22	57:N1:139:ARG:HG2	1.97	0.46
45:L8:110:THR:O	45:L8:114:ALA:HB3	2.85	0.46
45:L8:33:ASN:O	45:L8:39:ALA:HB3	2.16	0.46
47:M0:48:LEU:HD22	47:M0:49:CYS:N	2.30	0.46
52:M6:124:LEU:HD12	52:M6:124:LEU:HA	1.61	0.46
53:M7:4:TYR:CE1	53:M7:16:SER:HB2	3.05	0.46
55:M9:108:LYS:HA	55:M9:111:ASP:HB2	1.97	0.46
58:N2:59:ASP:OD2	58:N2:59:ASP:N	4.37	0.46
59:N3:102:ILE:HD11	59:N3:110:LYS:HD3	1.97	0.46
59:N3:54:LEU:HA	59:N3:54:LEU:HD12	1.58	0.46
59:N3:87:ARG:HH12	59:N3:137:VAL:CG1	2.81	0.46
59:N3:87:ARG:HB2	59:N3:89:ASP:OD1	4.32	0.46
63:N7:95:VAL:HG11	63:N7:113:VAL:HG21	4.59	0.46
64:N8:74:ASN:ND2	64:N8:113:LEU:HB2	2.30	0.46
70:O4:41:ARG:CZ	36:5:1739:U:H1'	188.37	0.46
36:1:2802:A:C8	78:Q2:56:PRO:HA	2.51	0.46
79:Q3:49:ARG:CG	79:Q3:50:GLY:H	2.29	0.46
3:S1:113:MET:HE3	3:S1:142:PHE:CE2	6.44	0.46
3:S1:118:GLN:OE1	3:S1:208:GLN:NE2	3.04	0.46
4:S2:164:SER:O	4:S2:202:GLY:HA3	2.94	0.46
5:S3:124:ARG:NH2	35:SM:128:ALA:HB2	9.54	0.46
6:S4:15:PRO:HD2	6:S4:18:TRP:CZ3	2.50	0.46
7:S5:73:THR:C	7:S5:75:GLY:N	3.14	0.46
9:S7:91:ILE:HG12	9:S7:129:LEU:HD23	1.98	0.46
1:2:197:A:N6	10:S8:138:ASN:HD22	2.05	0.46
36:1:1246:G:H2'	36:1:1247:U:O4'	2.15	0.46
36:1:1312:C:O2	52:M6:87:MET:HE2	2.15	0.46
36:1:1949:G:H2'	36:1:1950:U:C6	2.51	0.46
86:1:4023:OHX:N2	86:1:4143:OHX:N5	2.63	0.46
36:1:664:U:H2'	36:1:665:A:C8	2.50	0.46
36:1:970:A:H1'	36:1:1112:A:N1	2.30	0.46
1:2:1039:A:N6	1:2:1091:A:C2	2.84	0.46
1:2:1291:G:H5'	4:S2:119:LYS:HE2	1.95	0.46
1:2:1477:G:N2	1:2:1530:C:O2	2.48	0.46
1:2:1553:G:N2	1:2:1555:A:H3'	2.31	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:1615:C:H4'	1:2:1616:G:O5'	2.15	0.46
41:L4:180:LYS:HA	36:5:1386:A:N3	118.24	0.46
74:O8:2:ALA:N	36:5:1613:A:OP1	139.05	0.46
55:M9:39:ASN:ND2	36:5:1765:U:OP2	93.85	0.46
36:5:180:C:C2'	36:5:181:U:H5'	2.46	0.46
78:Q2:33:ALA:HA	36:5:2767:U:OP1	184.77	0.46
36:5:2806:U:C2	36:5:2807:U:C5	3.03	0.46
86:5:4029:OHX:N6	86:5:4232:OHX:N2	2.64	0.46
1:6:1182:U:N3	1:6:1185:U:OP2	2.42	0.46
1:6:492:A:H1'	1:6:496:G:H1	1.81	0.46
1:6:542:A:H1'	1:6:543:C:P	2.55	0.46
1:6:720:G:N3	1:6:720:G:H5''	2.31	0.46
1:6:93:A:C6	1:6:398:G:C6	3.03	0.46
16:C4:31:THR:HA	16:C4:38:THR:HA	2.34	0.46
17:C5:130:ARG:CD	35:SM:74:LYS:HG2	2.45	0.46
17:C5:25:LEU:HA	17:C5:28:MET:SD	2.56	0.46
19:C7:71:PHE:HE1	19:C7:73:LEU:HD22	1.80	0.46
1:2:1525:A:OP1	21:C9:93:HIS:ND1	2.48	0.46
24:D2:111:MET:HG3	24:D2:112:ASP:O	4.28	0.46
25:D3:79:ASN:OD1	25:D3:81:LYS:HG3	2.23	0.46
25:D3:52:ILE:HG22	25:D3:99:ASN:HA	3.37	0.46
26:D4:104:SER:O	26:D4:108:ARG:N	2.75	0.46
27:D5:42:LEU:O	27:D5:46:LYS:HB2	2.15	0.46
24:D2:53:ILE:HD11	29:D7:24:LEU:O	2.16	0.46
39:L2:104:LEU:HD22	39:L2:158:ILE:HD11	2.75	0.46
39:L2:219:ILE:HD13	39:L2:223:SER:HB2	1.97	0.46
41:L4:192:GLY:HA2	41:L4:195:ARG:HB2	1.97	0.46
43:L6:47:PHE:HE2	43:L6:77:ARG:NE	2.50	0.46
44:L7:217:PRO:HG2	44:L7:218:ARG:H	1.81	0.46
48:M1:155:THR:OG1	48:M1:158:ASP:HB2	2.15	0.46
50:M4:39:ILE:HG13	50:M4:44:VAL:HA	1.97	0.46
52:M6:174:PHE:O	52:M6:177:LYS:N	2.87	0.46
53:M7:13:LYS:O	53:M7:151:THR:HA	2.65	0.46
53:M7:13:LYS:HD2	53:M7:152:GLU:OE1	2.15	0.46
54:M8:166:LEU:HA	54:M8:166:LEU:HD23	1.44	0.46
54:M8:16:ARG:HG3	36:5:974:G:H5'	173.33	0.46
54:M8:35:PHE:CZ	54:M8:39:ARG:HG3	3.32	0.46
41:L4:300:ARG:HE	54:M8:39:ARG:HA	1.80	0.46
55:M9:149:ALA:O	55:M9:150:GLN:NE2	7.00	0.46
55:M9:4:LEU:HA	55:M9:7:GLN:NE2	4.28	0.46
59:N3:33:ASN:HD21	59:N3:64:LYS:N	2.12	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
65:N9:7:HIS:CG	65:N9:8:THR:N	2.84	0.46
67:O1:51:LEU:HB3	67:O1:55:LEU:HD12	1.98	0.46
69:O3:47:LYS:HA	69:O3:104:PRO:HD2	2.04	0.46
74:O8:8:ILE:HG12	74:O8:61:LYS:NZ	2.31	0.46
79:Q3:55:TRP:CD2	79:Q3:71:VAL:HG13	2.51	0.46
6:S4:158:ASP:HB3	6:S4:173:ILE:O	2.16	0.46
6:S4:184:THR:C	6:S4:189:LEU:HD13	3.08	0.46
6:S4:72:VAL:HG12	6:S4:73:ASP:HB2	2.75	0.46
7:S5:215:ASP:O	7:S5:219:ARG:HB2	3.04	0.46
10:S8:50:GLY:O	10:S8:52:ASN:ND2	2.31	0.46
34:SR:260:ILE:HB	34:SR:274:LEU:HB2	1.98	0.46
34:SR:34:LEU:HD13	34:SR:94:VAL:HG22	1.98	0.46
36:1:1278:A:HO2'	36:1:1279:C:C5'	2.28	0.46
36:1:1323:G:O3'	56:N0:2:ALA:HA	2.16	0.46
36:1:1669:C:OP1	70:O4:24:LYS:HE2	2.16	0.46
36:1:2278:C:C2'	36:1:2279:A:H5''	2.45	0.46
36:1:2386:A:H2'	36:1:2387:A:O4'	2.16	0.46
36:1:2778:G:C2'	36:1:2779:A:H5'	2.45	0.46
36:1:2942:C:H5''	36:1:2943:G:H5''	1.96	0.46
36:1:2987:A:H2'	36:1:2988:C:C6	2.51	0.46
36:1:3358:U:H2'	36:1:3359:A:C1'	2.45	0.46
36:1:352:A:H61	36:1:365:A:H5''	1.81	0.46
1:2:1383:G:H1'	22:D0:57:ARG:HH12	1.79	0.46
1:2:1440:C:H2'	1:2:1441:C:H6	1.80	0.46
1:2:329:G:O6	86:2:2106:OHX:N6	2.49	0.46
1:2:500:C:C5	1:2:501:U:C4	3.04	0.46
1:2:833:U:H5'	1:2:834:G:H5''	1.96	0.46
38:4:57:C:O2'	38:4:58:G:H5'	2.16	0.46
36:5:1025:A:H5'	36:5:1026:A:OP2	2.15	0.46
36:5:2304:C:C5	36:5:2305:G:C6	3.04	0.46
36:5:2437:G:C6	36:5:2511:A:C6	3.03	0.46
36:5:3165:A:N1	36:5:3285:C:N3	2.64	0.46
36:5:1898:G:OP2	86:5:3938:OHX:N5	2.48	0.46
22:D0:23:ARG:NH2	1:6:1347:U:OP2	458.51	0.46
19:C7:44:LYS:NZ	1:6:1386:G:OP2	442.53	0.46
1:6:1406:A:H2'	1:6:1407:U:H6	1.81	0.46
27:D5:77:ARG:NH1	1:6:1533:C:OP2	352.72	0.46
1:6:1534:G:H4'	1:6:1536:G:O6	2.15	0.46
1:6:97:C:O2	1:6:425:A:O2'	2.25	0.46
13:C1:104:HIS:O	13:C1:105:LYS:HG2	4.39	0.46
13:C1:7:VAL:HB	13:C1:8:GLN:H	3.82	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:C1:91:LEU:HD13	13:C1:92:HIS:N	2.30	0.46
20:C8:73:MET:HB3	20:C8:101:LEU:HD11	1.97	0.46
21:C9:121:GLY:O	21:C9:122:ARG:HD3	2.16	0.46
21:C9:21:PHE:HD2	21:C9:22:LEU:HD13	1.80	0.46
21:C9:57:ARG:HH22	21:C9:80:TYR:HB3	3.37	0.46
25:D3:91:GLY:N	32:E0:12:GLY:HA2	2.31	0.46
30:D8:13:ILE:HG13	30:D8:29:ARG:O	2.16	0.46
30:D8:44:VAL:HG21	30:D8:48:VAL:CG2	2.77	0.46
39:L2:44:ILE:HD12	39:L2:62:VAL:O	2.16	0.46
40:L3:20:LYS:HG3	40:L3:21:ARG:O	2.16	0.46
40:L3:293:ASN:HD22	40:L3:305:ILE:HG22	5.90	0.46
40:L3:62:ARG:CZ	40:L3:349:LYS:HD2	2.45	0.46
41:L4:80:GLY:O	41:L4:82:THR:HG22	2.15	0.46
42:L5:50:ARG:HB3	42:L5:147:ASP:HB2	1.98	0.46
44:L7:60:ARG:HH22	36:5:517:G:P	306.03	0.46
45:L8:178:ALA:HB2	45:L8:218:ILE:HG23	1.96	0.46
47:M0:76:MET:HE1	47:M0:138:VAL:HG11	1.98	0.46
47:M0:34:TYR:N	47:M0:34:TYR:CD1	2.84	0.46
50:M4:6:ILE:HA	50:M4:6:ILE:HD13	1.74	0.46
51:M5:43:THR:OG1	51:M5:131:GLU:OE2	2.30	0.46
55:M9:138:LEU:O	55:M9:142:ILE:HG13	2.15	0.46
55:M9:175:GLN:HA	55:M9:178:ALA:HB3	1.97	0.46
56:N0:138:GLN:C	56:N0:140:VAL:H	2.66	0.46
56:N0:46:GLN:HG2	56:N0:51:VAL:O	3.22	0.46
56:N0:42:TRP:O	56:N0:46:GLN:HG3	4.32	0.46
58:N2:17:VAL:HB	58:N2:63:VAL:HG23	2.52	0.46
59:N3:120:LYS:O	59:N3:124:ASP:HB2	2.45	0.46
70:O4:81:CYS:HG	70:O4:84:CYS:HG	2.08	0.46
51:M5:6:TYR:CE2	72:O6:40:VAL:HG22	4.13	0.46
74:O8:70:PRO:HA	74:O8:71:PRO:HD2	1.67	0.46
75:O9:8:ARG:O	75:O9:11:GLN:HB3	2.88	0.46
78:Q2:63:LYS:HE3	36:5:2760:C:N3	216.92	0.46
3:S1:135:LEU:HD13	3:S1:137:ILE:HG23	1.98	0.46
3:S1:218:LEU:HD23	3:S1:219:LYS:H	1.80	0.46
6:S4:18:TRP:CE3	6:S4:20:LEU:HD11	2.49	0.46
8:S6:163:THR:HG22	8:S6:168:THR:HG22	1.97	0.46
1:2:127:G:C6	8:S6:195:VAL:HG13	2.51	0.46
11:S9:108:ARG:HB2	11:S9:111:THR:HG23	2.58	0.46
36:1:1139:G:C5	36:1:1140:G:C8	3.04	0.46
36:1:1547:G:OP1	51:M5:108:ARG:NH2	2.46	0.46
36:1:2206:G:N2	36:1:2237:C:N3	2.47	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:2397:A:C2	36:1:2873:U:H5'	2.50	0.46
36:1:2417:U:H2'	36:1:2606:G:N2	2.31	0.46
36:1:2988:C:P	52:M6:68:ARG:NH1	2.88	0.46
36:1:3152:U:O2	86:1:4141:OHX:N4	2.49	0.46
36:1:3386:G:H2'	36:1:3387:U:H6	1.80	0.46
1:2:1388:A:HO2'	1:2:1411:A:H2	1.63	0.46
1:2:276:C:O2'	1:2:277:U:H5''	2.15	0.46
1:2:431:C:H3'	1:2:432:G:H8	1.80	0.46
1:2:516:G:N2	1:2:537:G:H1'	2.31	0.46
1:2:681:U:C4	1:2:682:C:C5	3.04	0.46
37:3:9:C:H5''	37:3:10:C:OP2	2.16	0.46
36:5:118:U:O2	36:5:121:A:H5'	2.16	0.46
36:5:1263:A:H2'	36:5:1263:A:N3	2.30	0.46
36:5:201:A:H2'	36:5:202:G:H8	1.80	0.46
36:5:2509:U:H2'	36:5:2510:U:C5'	2.45	0.46
76:Q0:112:LYS:NZ	36:5:3107:U:P	304.52	0.46
86:5:3994:OHX:N3	86:5:4085:OHX:N5	2.63	0.46
36:5:409:A:OP2	86:5:4097:OHX:N3	2.48	0.46
36:5:373:A:OP1	86:5:4145:OHX:N6	2.49	0.46
86:5:3971:OHX:N4	86:5:4193:OHX:N3	2.64	0.46
36:5:799:G:H2'	36:5:801:A:N7	2.31	0.46
53:M7:133:HIS:HA	36:5:883:A:H5'	164.00	0.46
1:6:1354:G:C6	1:6:1355:C:C4	3.04	0.46
1:6:1358:G:H2'	1:6:1359:C:C6	2.51	0.46
1:6:1268:G:H1'	1:6:1448:G:H5''	1.97	0.46
1:6:1614:A:C6	1:6:1615:C:N4	2.84	0.46
1:6:37:U:O2'	1:6:770:A:N1	2.40	0.46
1:6:790:U:H2'	1:6:791:A:H8	1.81	0.46
37:7:92:A:H5''	37:7:93:C:OP2	2.15	0.46
13:C1:109:VAL:HA	13:C1:135:VAL:HG13	1.98	0.46
1:2:1788:G:P	16:C4:127:ARG:HH22	2.36	0.46
19:C7:100:LEU:H	19:C7:118:PRO:HG2	1.80	0.46
20:C8:27:LYS:O	20:C8:31:ALA:N	2.87	0.46
23:D1:79:LEU:HD23	23:D1:79:LEU:HA	1.70	0.46
24:D2:20:THR:OG1	24:D2:22:LYS:HD3	2.27	0.46
25:D3:79:ASN:CG	25:D3:81:LYS:HG3	2.62	0.46
29:D7:80:ARG:HG2	29:D7:81:ARG:N	2.39	0.46
1:2:1235:C:C2	33:E1:138:ARG:NH2	2.84	0.46
41:L4:290:ILE:HG23	54:M8:35:PHE:CE2	2.50	0.46
49:M3:64:LYS:HD3	49:M3:65:TYR:CE1	3.13	0.46
52:M6:15:LEU:O	52:M6:18:ARG:N	2.47	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:M8:21:SER:OG	54:M8:22:ASP:N	2.46	0.46
63:N7:5:LEU:HD22	63:N7:25:ILE:HD12	1.98	0.46
64:N8:74:ASN:ND2	64:N8:115:LYS:HB2	2.31	0.46
68:O2:97:ALA:O	68:O2:100:ILE:HG12	2.62	0.46
71:O5:9:LEU:O	71:O5:12:LYS:N	2.47	0.46
73:O7:27:PHE:HA	73:O7:34:CYS:HA	1.97	0.46
2:S0:9:LEU:HD21	2:S0:14:ALA:HB2	1.96	0.46
3:S1:190:PRO:HG2	3:S1:192:VAL:HG23	2.11	0.46
3:S1:33:LYS:HB3	3:S1:232:HIS:HE1	7.58	0.46
2:S0:140:ASN:OD1	4:S2:62:PRO:HD3	2.21	0.46
7:S5:166:ARG:HD2	30:D8:46:GLY:CA	2.46	0.46
7:S5:216:GLU:OE2	7:S5:219:ARG:NH2	2.49	0.46
7:S5:91:GLU:HA	7:S5:94:THR:HG23	1.97	0.46
8:S6:175:ILE:H	8:S6:175:ILE:HG12	1.55	0.46
35:SM:86:ASN:HD21	35:SM:89:ARG:HB2	1.80	0.46
34:SR:91:LEU:HG	34:SR:100:TYR:HB2	2.27	0.46
34:SR:6:VAL:HG22	34:SR:7:LEU:H	1.81	0.46
36:1:1020:G:O6	36:1:1032:C:N4	2.39	0.46
36:1:1340:G:H2'	36:1:1341:U:H6	1.81	0.46
36:1:1439:U:H2'	36:1:1440:G:C8	2.50	0.46
36:1:1801:U:O5'	36:1:1801:U:H6	1.99	0.46
36:1:2424:A:H2'	36:1:2425:G:O4'	2.16	0.46
36:1:2991:A:P	40:L3:20:LYS:HB2	2.56	0.46
36:1:3024:A:H4'	46:L9:97:PHE:CE2	2.51	0.46
36:1:3087:A:P	86:1:4178:OHX:N5	2.89	0.46
36:1:386:A:C5	36:1:387:A:H1'	2.51	0.46
36:1:1734:G:N7	86:1:3908:OHX:N5	2.63	0.46
36:1:3155:U:O2	86:1:4141:OHX:N3	2.49	0.46
36:1:568:G:N7	86:1:3936:OHX:N4	2.64	0.46
36:1:729:C:H2'	36:1:730:C:C6	2.45	0.46
1:2:1550:A:H2'	1:2:1551:U:C6	2.51	0.46
1:2:1203:A:C4	1:2:1556:A:C2	3.04	0.46
1:2:540:G:H4'	1:2:541:A:H3'	1.97	0.46
38:4:55:U:O2	86:4:229:OHX:N6	2.49	0.46
38:4:56:G:H2'	38:4:57:C:C6	2.51	0.46
56:N0:137:ARG:NH1	36:5:1213:G:P	324.51	0.46
56:N0:115:ARG:NH1	36:5:1296:C:H5'	291.37	0.46
36:5:1439:U:H2'	36:5:1440:G:C8	2.51	0.46
36:5:2232:A:O2'	36:5:2429:G:H5'	2.15	0.46
36:5:2534:G:H1	36:5:2545:C:H42	1.63	0.46
36:5:3006:A:H2'	36:5:3007:U:O4'	2.15	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:M5:96:ARG:HD2	36:5:31:C:H4'	123.69	0.46
36:5:415:G:H2'	36:5:416:A:C8	2.51	0.46
36:5:550:A:C6	36:5:551:A:C6	3.04	0.46
1:6:1446:A:O2'	1:6:1447:C:H5''	2.15	0.46
1:6:1488:G:H3'	1:6:1515:A:H61	1.79	0.46
10:S8:137:LYS:NZ	1:6:192:U:O4	264.55	0.46
1:6:439:U:O4'	1:6:465:G:N2	2.48	0.46
42:L5:155:THR:HG23	37:7:36:C:C5'	270.14	0.46
38:8:145:U:H2'	38:8:146:U:O4'	2.16	0.46
38:8:88:A:H5''	38:8:89:A:OP2	2.16	0.46
14:C2:136:ILE:HA	14:C2:139:HIS:HB3	1.98	0.46
14:C2:45:LEU:HB2	1:6:1228:G:OP1	462.87	0.46
15:C3:39:LYS:HD2	15:C3:40:TYR:CD1	5.55	0.46
15:C3:71:ILE:HD12	1:6:961:U:H5''	328.97	0.46
18:C6:35:PRO:HG2	18:C6:38:LEU:HG	1.98	0.46
21:C9:10:ALA:O	21:C9:13:ASP:N	2.49	0.46
23:D1:32:VAL:HB	23:D1:60:ARG:HD3	1.96	0.46
24:D2:37:PHE:CE2	24:D2:103:ILE:HD11	3.34	0.46
25:D3:104:LEU:HD23	25:D3:104:LEU:HA	1.77	0.46
13:C1:100:TYR:HB2	25:D3:10:ASN:OD1	2.16	0.46
28:D6:9:GLY:O	28:D6:10:ARG:HG3	2.16	0.46
28:D6:26:CYS:HB2	28:D6:28:LYS:H	3.85	0.46
28:D6:4:LYS:HE2	28:D6:5:ARG:NH2	2.31	0.46
30:D8:25:VAL:HG11	30:D8:43:ASN:HB3	2.64	0.46
32:E0:20:LYS:HD2	32:E0:21:VAL:N	4.62	0.46
39:L2:96:LEU:HD11	39:L2:107:VAL:HG12	1.98	0.46
40:L3:332:ARG:O	40:L3:333:LYS:HB2	2.20	0.46
43:L6:102:ASN:N	43:L6:102:ASN:OD1	3.81	0.46
44:L7:25:GLN:N	44:L7:28:ALA:HB3	2.31	0.46
45:L8:63:LYS:HG3	45:L8:67:ILE:HD11	1.98	0.46
46:L9:101:VAL:HG12	46:L9:136:PHE:HZ	1.81	0.46
47:M0:134:ILE:HD11	57:N1:160:ILE:HD11	1.96	0.46
49:M3:168:ARG:HG3	49:M3:172:LEU:HG	2.43	0.46
51:M5:10:LEU:HD22	51:M5:19:LEU:HD13	1.97	0.46
53:M7:171:ARG:HB2	53:M7:171:ARG:HH11	1.80	0.46
57:N1:57:TYR:CE1	57:N1:89:LEU:HD11	2.51	0.46
61:N5:24:LEU:HB3	61:N5:25:LYS:H	1.33	0.46
61:N5:40:LEU:HA	61:N5:40:LEU:HD13	2.42	0.46
63:N7:11:ALA:HB1	63:N7:80:LEU:HB3	1.97	0.46
66:O0:46:ALA:O	66:O0:48:THR:N	3.25	0.46
66:O0:65:THR:O	66:O0:67:VAL:HG23	2.99	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
69:O3:107:ILE:HD12	69:O3:107:ILE:HA	3.42	0.46
69:O3:85:PHE:CZ	69:O3:89:LEU:HD11	2.60	0.46
70:O4:94:LEU:HA	70:O4:94:LEU:HD23	2.03	0.46
72:O6:70:ARG:HD3	72:O6:84:LYS:CG	3.00	0.46
3:S1:196:GLU:O	3:S1:199:ASN:HB2	2.16	0.46
5:S3:62:ASN:OD1	5:S3:62:ASN:N	2.47	0.46
7:S5:220:VAL:HA	7:S5:223:SER:HB3	1.97	0.46
7:S5:84:LYS:O	7:S5:92:ARG:HD2	4.10	0.46
8:S6:73:ILE:HD11	8:S6:75:LEU:HD21	2.86	0.46
9:S7:105:THR:O	9:S7:107:ARG:N	3.81	0.46
9:S7:15:GLU:O	9:S7:19:GLN:HG3	3.48	0.46
9:S7:170:GLN:HG2	9:S7:181:ILE:HG22	1.97	0.46
9:S7:4:PRO:HA	9:S7:7:LYS:HD3	1.98	0.46
10:S8:184:LEU:HD12	10:S8:184:LEU:HA	1.65	0.46
11:S9:124:HIS:CE1	11:S9:128:LEU:HD11	2.77	0.46
34:SR:187:GLN:HG2	34:SR:187:GLN:O	2.16	0.46
34:SR:205:SER:O	34:SR:208:GLY:N	3.08	0.46
34:SR:262:VAL:O	34:SR:271:VAL:N	2.79	0.46
36:1:1614:C:H2'	36:1:1615:C:H6	1.81	0.46
36:1:1658:G:O4'	36:1:1796:G:H2'	2.15	0.46
86:1:4027:OHX:N6	86:1:4040:OHX:N3	2.64	0.46
36:1:2767:U:O4	86:1:4033:OHX:N6	2.49	0.46
36:1:966:U:N3	36:1:967:A:N7	2.63	0.46
1:2:131:C:O2'	1:2:132:U:OP1	2.28	0.46
1:2:1530:C:H2'	1:2:1531:G:O4'	2.16	0.46
1:2:197:A:H61	10:S8:138:ASN:ND2	2.07	0.46
1:2:761:G:O2'	1:2:789:A:N6	2.49	0.46
36:5:1220:U:O2	36:5:1222:G:N1	2.49	0.46
39:L2:40:TYR:O	36:5:2550:U:H5	212.31	0.46
36:5:2738:A:H2'	36:5:2739:A:C8	2.51	0.46
36:5:2960:C:H2'	36:5:2961:G:C8	2.51	0.46
36:5:355:A:H2'	36:5:356:C:O4'	2.16	0.46
36:5:1151:U:OP1	86:5:4206:OHX:N1	2.48	0.46
1:6:1315:U:C4	1:6:1316:G:N7	2.84	0.46
1:6:1742:U:C4	1:6:1743:U:C5	3.03	0.46
1:6:76:A:H3'	86:6:2192:OHX:N1	2.31	0.46
36:5:1055:A:H4'	37:7:100:C:O2	2.15	0.46
14:C2:67:THR:C	14:C2:69:ALA:H	2.23	0.46
16:C4:16:VAL:HG22	16:C4:32:ASP:O	4.42	0.46
17:C5:17:TYR:HB2	17:C5:25:LEU:HD11	2.61	0.46
17:C5:86:VAL:O	17:C5:89:MET:HG3	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:D6:66:LYS:HE2	28:D6:66:LYS:HB2	1.78	0.46
14:C2:46:ARG:HE	33:E1:102:VAL:HG21	5.43	0.46
33:E1:136:LYS:O	33:E1:138:ARG:HB2	2.16	0.46
39:L2:179:LEU:O	39:L2:184:ARG:HG3	2.16	0.46
40:L3:128:LYS:O	40:L3:131:THR:HG23	2.34	0.46
40:L3:148:LEU:HD12	40:L3:148:LEU:HA	2.22	0.46
40:L3:243:HIS:ND1	40:L3:244:ARG:N	2.63	0.46
41:L4:31:ARG:HB3	41:L4:34:ILE:HG13	1.98	0.46
42:L5:88:ILE:HD11	42:L5:240:TYR:CD1	2.51	0.46
44:L7:82:LYS:O	44:L7:119:VAL:HG23	2.53	0.46
44:L7:160:ARG:HG3	44:L7:203:TRP:CD2	3.16	0.46
44:L7:49:ALA:O	44:L7:52:GLN:HB3	2.87	0.46
45:L8:136:LEU:HD11	45:L8:162:LEU:O	2.16	0.46
46:L9:7:GLU:HG2	46:L9:8:GLN:N	2.29	0.46
48:M1:15:GLU:HB3	48:M1:130:VAL:HG22	2.66	0.46
51:M5:69:GLY:O	36:5:290:G:H4'	145.71	0.46
53:M7:10:ASN:OD1	53:M7:12:ALA:HB3	2.15	0.46
57:N1:147:VAL:HA	57:N1:148:PRO:HD3	2.01	0.46
57:N1:97:LYS:HG2	57:N1:98:HIS:H	3.69	0.46
62:N6:91:ASN:O	62:N6:93:ALA:N	2.49	0.46
64:N8:94:ALA:HB1	64:N8:121:VAL:HG13	1.98	0.46
36:1:1431:G:N7	64:N8:9:ARG:NH2	2.63	0.46
65:N9:40:ARG:O	65:N9:41:ARG:C	2.72	0.46
67:O1:9:THR:HG22	67:O1:109:VAL:HB	2.88	0.46
72:O6:11:LEU:HA	72:O6:11:LEU:HD12	1.80	0.46
73:O7:11:ARG:HB3	73:O7:11:ARG:HH11	1.80	0.46
2:S0:167:LYS:HG2	2:S0:168:HIS:CD2	2.51	0.46
4:S2:177:GLY:C	4:S2:195:ASP:HA	3.02	0.46
6:S4:159:THR:CG2	6:S4:173:ILE:HB	2.76	0.46
8:S6:121:LEU:H	8:S6:125:THR:HG1	2.50	0.46
8:S6:39:GLU:OE2	8:S6:46:LYS:HG3	3.41	0.46
8:S6:58:LYS:O	8:S6:59:GLN:HB2	2.16	0.46
11:S9:64:GLU:HG3	11:S9:69:ARG:NH2	3.04	0.46
36:1:1230:G:H1	36:1:1279:C:N4	2.09	0.46
36:1:1620:U:H2'	36:1:1621:A:C8	2.51	0.46
36:1:1758:G:N2	36:1:1767:C:N3	2.48	0.46
36:1:2621:G:C6	36:1:2622:C:C4	3.03	0.46
36:1:2812:C:H2'	36:1:2813:A:H8	1.81	0.46
36:1:290:G:H4'	51:M5:69:GLY:O	2.15	0.46
86:1:3965:OHX:N3	86:1:4153:OHX:N1	2.63	0.46
36:1:703:G:C5	36:1:704:U:C5	3.04	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:906:A:OP1	86:1:3994:OHX:N1	2.49	0.46
1:2:1231:U:C4	1:2:1255:G:N2	2.84	0.46
1:2:154:G:H5'	8:S6:108:VAL:HG21	1.97	0.46
1:2:1783:C:C5	77:Q1:5:TRP:CD1	3.04	0.46
36:5:1239:C:H3'	36:5:1240:A:C8	2.51	0.46
36:5:1614:C:H2'	36:5:1615:C:C6	2.50	0.46
36:5:188:U:H1'	36:5:208:C:H1'	1.98	0.46
36:5:2378:C:H2'	36:5:2378:C:O2	2.16	0.46
76:Q0:102:ARG:HH21	36:5:2896:A:P	318.78	0.46
36:5:2947:G:N2	36:5:2948:C:C2	2.84	0.46
36:5:2948:C:H6	36:5:2948:C:O5'	1.98	0.46
36:5:3194:C:O2'	36:5:3195:U:H5'	2.16	0.46
36:5:3348:G:C2	36:5:3358:U:C2	3.04	0.46
36:5:1940:G:N2	36:5:3362:A:H8	2.14	0.46
36:5:614:C:H2'	36:5:615:U:O4'	2.15	0.46
37:7:3:U:O2'	37:7:4:U:H5'	2.16	0.46
14:C2:64:SER:OG	14:C2:65:SER:N	2.49	0.46
15:C3:119:GLU:O	15:C3:122:ILE:HB	2.16	0.46
15:C3:94:LYS:HE2	1:6:952:A:OP1	298.36	0.46
16:C4:108:SER:OG	16:C4:109:GLY:N	3.28	0.46
16:C4:16:VAL:O	16:C4:30:VAL:HA	2.18	0.46
17:C5:74:ALA:HA	17:C5:75:PRO:HD3	2.22	0.46
19:C7:104:ASN:OD1	19:C7:104:ASN:N	3.52	0.46
19:C7:53:TYR:CE1	19:C7:57:LEU:HD21	2.50	0.46
19:C7:57:LEU:O	19:C7:61:ILE:HG13	2.16	0.46
24:D2:81:VAL:N	24:D2:123:GLY:O	2.97	0.46
27:D5:61:SER:H	27:D5:64:VAL:CG2	2.81	0.46
7:S5:166:ARG:HD2	30:D8:46:GLY:HA2	1.98	0.46
30:D8:48:VAL:HG13	30:D8:52:ASP:OD2	2.15	0.46
33:E1:139:LEU:HD13	33:E1:152:ALA:H	1.80	0.46
39:L2:61:VAL:HG21	39:L2:76:PHE:CD2	2.50	0.46
36:1:3011:A:C5	40:L3:13:HIS:CD2	3.04	0.46
40:L3:221:THR:CG2	40:L3:273:HIS:H	3.53	0.46
40:L3:65:SER:C	40:L3:67:PHE:H	2.55	0.46
41:L4:23:PRO:HG2	41:L4:26:PHE:HE2	1.81	0.46
43:L6:18:LEU:HA	43:L6:18:LEU:HD23	4.31	0.46
43:L6:50:LYS:HE2	43:L6:72:ASN:HB2	4.56	0.46
43:L6:94:GLU:HA	43:L6:94:GLU:OE1	3.39	0.46
44:L7:151:ARG:HD2	44:L7:244:ASN:OD1	2.56	0.46
45:L8:159:PRO:HB2	45:L8:161:GLU:OE2	3.48	0.46
46:L9:30:PRO:HG2	46:L9:83:THR:HA	3.64	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:M1:105:GLY:HA3	36:5:2674:A:H5''	333.58	0.46
50:M4:5:SER:O	50:M4:6:ILE:HB	2.15	0.46
51:M5:98:LEU:O	51:M5:101:THR:N	2.78	0.46
36:1:3191:G:H5''	52:M6:176:LYS:HE2	1.97	0.46
52:M6:79:ILE:HG21	52:M6:138:LEU:HD11	1.98	0.46
56:N0:100:VAL:HG13	56:N0:101:ALA:N	2.31	0.46
57:N1:82:ASN:OD1	57:N1:82:ASN:N	2.66	0.46
58:N2:33:TYR:OH	58:N2:80:THR:HG21	2.15	0.46
59:N3:80:ARG:NH1	59:N3:116:GLY:HA3	2.92	0.46
62:N6:39:LEU:HD13	62:N6:43:TYR:HE2	1.81	0.46
62:N6:88:GLU:OE1	62:N6:88:GLU:N	4.20	0.46
66:O0:52:ARG:HB2	66:O0:52:ARG:HE	1.51	0.46
69:O3:73:ARG:HE	69:O3:82:ARG:CZ	2.71	0.46
36:1:316:U:O2'	72:O6:30:LYS:HG3	2.16	0.46
73:O7:28:HIS:CG	73:O7:31:LYS:HB2	3.24	0.46
2:S0:12:GLU:HA	2:S0:15:GLN:HG3	2.81	0.46
2:S0:58:VAL:O	2:S0:61:ALA:HB3	2.48	0.46
6:S4:46:VAL:O	6:S4:50:ASN:HB2	2.30	0.46
7:S5:129:PRO:O	7:S5:133:VAL:HG23	2.16	0.46
34:SR:84:SER:OG	34:SR:85:TRP:N	2.49	0.46
36:1:1583:A:N7	36:1:1584:U:C2	2.84	0.45
36:1:2134:G:C2	36:1:2135:U:C6	3.03	0.45
36:1:2714:G:O6	36:1:2741:C:N3	2.49	0.45
36:1:2771:U:O2'	36:1:2772:C:O4'	2.27	0.45
36:1:2916:U:H2'	36:1:2917:G:H5'	1.97	0.45
36:1:3160:U:H2'	36:1:3161:C:C6	2.51	0.45
36:1:3166:C:N4	36:1:3284:G:H1	1.95	0.45
36:1:537:A:H2'	36:1:538:G:O4'	2.16	0.45
36:1:650:C:O2'	36:1:651:G:H5'	2.16	0.45
36:1:709:A:C1'	64:N8:57:GLY:HA2	2.46	0.45
1:2:119:A:H1'	1:2:397:A:C4	2.50	0.45
1:2:1565:C:H2'	1:2:1566:U:O4'	2.16	0.45
1:2:1789:G:C8	1:2:1789:G:H5''	2.51	0.45
86:2:2043:OHX:N4	86:2:2098:OHX:N3	2.65	0.45
1:2:460:A:H5'	1:2:461:G:OP2	2.17	0.45
1:2:439:U:O5'	1:2:465:G:N2	2.49	0.45
1:2:20:G:H5'	1:2:571:G:C8	2.51	0.45
1:2:739:G:O6	86:2:2096:OHX:N4	2.48	0.45
1:2:862:A:OP1	15:C3:20:ARG:NE	2.24	0.45
1:2:941:A:C5	1:2:942:G:H1'	2.51	0.45
38:4:17:A:C2	38:4:18:U:H1'	2.51	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:4:35:C:H5''	73:O7:70:VAL:HG11	1.98	0.45
36:5:1104:G:H2'	36:5:1105:A:C8	2.52	0.45
36:5:1439:U:H2'	36:5:1440:G:O4'	2.16	0.45
36:5:2180:G:H2'	36:5:2181:C:C6	2.51	0.45
53:M7:139:TYR:CZ	36:5:2355:G:H4'	146.99	0.45
36:5:1447:G:O2'	36:5:2355:G:O6	2.32	0.45
36:5:300:G:H2'	36:5:301:G:H8	1.81	0.45
15:C3:11:ILE:HD11	1:6:1072:C:H4'	349.73	0.45
1:6:1347:U:O2	1:6:1516:A:H2'	2.16	0.45
28:D6:87:ARG:HD2	1:6:1797:A:C6	345.05	0.45
1:6:565:C:N3	86:6:2159:OHX:N4	2.63	0.45
37:7:57:G:C8	37:7:58:C:C5	3.04	0.45
12:C0:49:LEU:HD13	12:C0:52:LYS:HE3	4.93	0.45
13:C1:54:ILE:HG22	13:C1:55:ASP:N	2.31	0.45
14:C2:30:VAL:O	14:C2:34:THR:HG23	2.40	0.45
15:C3:121:ARG:NH1	1:6:868:G:OP1	312.21	0.45
1:2:928:U:H4'	16:C4:124:ASP:OD1	2.16	0.45
17:C5:22:LEU:O	17:C5:26:LEU:HD13	2.15	0.45
20:C8:52:VAL:HG13	20:C8:61:LEU:HD21	3.12	0.45
21:C9:18:TYR:O	21:C9:22:LEU:HD22	2.16	0.45
26:D4:40:LEU:O	26:D4:44:LEU:HD12	2.48	0.45
27:D5:76:ALA:O	27:D5:79:ALA:HB3	2.93	0.45
29:D7:34:ASP:OD1	29:D7:34:ASP:N	2.49	0.45
44:L7:113:SER:HA	44:L7:205:PHE:O	2.47	0.45
45:L8:178:ALA:HB2	45:L8:218:ILE:HD13	3.06	0.45
46:L9:76:ASP:O	46:L9:80:THR:HG23	2.16	0.45
47:M0:144:ASN:O	47:M0:145:LYS:C	2.60	0.45
47:M0:201:SER:OG	47:M0:203:LYS:O	2.27	0.45
47:M0:30:LYS:HG3	47:M0:63:GLU:HB3	3.92	0.45
48:M1:173:ASP:HB3	48:M1:174:LYS:H	1.81	0.45
49:M3:124:ILE:HD11	49:M3:126:PHE:CE1	2.41	0.45
49:M3:149:GLN:HA	49:M3:150:PRO:HD3	1.50	0.45
50:M4:68:LEU:N	50:M4:68:LEU:HD23	2.78	0.45
53:M7:30:ARG:NH1	53:M7:31:GLU:OE2	2.97	0.45
57:N1:39:ILE:HD12	57:N1:102:ARG:HB2	2.29	0.45
57:N1:12:ARG:HD3	57:N1:13:TYR:CE1	2.86	0.45
57:N1:7:TYR:CE2	57:N1:54:HIS:HD2	2.53	0.45
61:N5:100:LYS:HE3	61:N5:106:ASP:HA	1.97	0.45
68:O2:96:ILE:HD11	68:O2:109:LEU:HD23	1.98	0.45
68:O2:115:LEU:HA	68:O2:115:LEU:HD23	1.73	0.45
68:O2:6:HIS:HA	68:O2:7:PRO:HD2	3.01	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
69:O3:29:LEU:HD22	69:O3:75:HIS:CD2	2.78	0.45
71:O5:4:VAL:HG21	71:O5:9:LEU:HD11	2.25	0.45
36:1:1493:G:C6	75:O9:2:ALA:HB2	2.50	0.45
2:S0:170:ILE:H	2:S0:170:ILE:HD12	1.81	0.45
2:S0:179:ARG:O	2:S0:183:ARG:HG3	2.16	0.45
3:S1:127:VAL:HG13	3:S1:176:VAL:HG11	1.99	0.45
3:S1:222:LYS:HD3	3:S1:223:PHE:N	2.28	0.45
4:S2:49:LYS:HB3	4:S2:243:TYR:CE2	2.51	0.45
8:S6:204:ALA:HA	8:S6:207:GLU:HB3	2.22	0.45
10:S8:50:GLY:HA2	1:6:397:A:O3'	315.24	0.45
34:SR:147:HIS:CD2	34:SR:151:VAL:HG22	2.51	0.45
36:1:1063:G:O2'	36:1:1097:G:N2	2.48	0.45
36:1:1202:A:C2	36:1:2857:C:H5'	2.52	0.45
36:1:1625:A:H4'	36:1:1643:A:C6	2.51	0.45
36:1:1836:C:N4	75:O9:3:ALA:HB2	2.31	0.45
36:1:2209:U:H6	36:1:2209:U:OP2	1.98	0.45
36:1:2381:G:C2	36:1:2382:G:C8	3.04	0.45
36:1:2769:A:C2'	36:1:2770:G:H5'	2.46	0.45
36:1:921:A:N6	36:1:1846:C:OP2	2.48	0.45
1:2:1252:C:H41	33:E1:97:LYS:HE3	1.80	0.45
1:2:1062:A:OP2	86:2:2165:OHX:N4	2.49	0.45
1:2:327:U:C2	1:2:328:A:C8	3.04	0.45
1:2:416:A:H5'	1:2:417:A:N7	2.31	0.45
1:2:423:G:N7	86:2:2107:OHX:N3	2.65	0.45
1:2:601:A:H2'	1:2:602:U:O4'	2.16	0.45
1:2:906:A:H2'	1:2:907:A:O4'	2.17	0.45
38:4:120:C:C4	38:4:121:U:C4	3.04	0.45
38:4:97:A:C2	38:4:98:U:C2	3.05	0.45
36:5:100:A:O2'	36:5:101:G:H5'	2.16	0.45
36:5:1077:U:H2'	36:5:1078:U:C6	2.52	0.45
36:5:1192:C:N4	36:5:1302:A:OP2	2.45	0.45
67:O1:57:GLN:HG2	36:5:1475:A:H4'	146.92	0.45
36:5:1919:G:C6	36:5:1920:U:C4	3.04	0.45
36:5:2222:A:N6	36:5:2783:U:H1'	2.32	0.45
36:5:265:A:H5''	36:5:266:A:OP2	2.16	0.45
36:5:2720:G:O6	36:5:2737:C:N4	2.49	0.45
36:5:2945:G:O2'	36:5:2948:C:OP2	2.22	0.45
67:O1:17:HIS:NE2	36:5:3059:G:OP1	189.16	0.45
36:5:3216:G:O6	36:5:3259:U:H2'	2.16	0.45
36:5:343:U:H4'	36:5:344:A:OP2	2.16	0.45
36:5:600:G:H5'	36:5:601:U:OP2	2.17	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:916:G:H4'	36:5:917:A:O5'	2.15	0.45
1:6:1238:A:H2'	1:6:1239:U:C5'	2.46	0.45
8:S6:179:VAL:HG21	1:6:140:A:H1'	328.34	0.45
1:6:1491:U:O2'	1:6:1492:A:H5''	2.15	0.45
1:6:1554:U:H3'	1:6:1555:A:H8	1.81	0.45
1:6:1731:A:H5''	1:6:1732:A:OP2	2.16	0.45
1:6:1:U:O2	1:6:369:A:H2'	2.16	0.45
38:8:121:U:O2'	38:8:122:U:H5'	2.15	0.45
15:C3:54:LEU:HA	15:C3:58:HIS:HB2	1.97	0.45
18:C6:47:LYS:HA	18:C6:47:LYS:HD2	1.88	0.45
27:D5:39:ALA:HB1	27:D5:71:ILE:C	2.37	0.45
39:L2:143:GLU:O	39:L2:145:LYS:HG2	2.16	0.45
39:L2:218:HIS:O	39:L2:218:HIS:ND1	2.45	0.45
40:L3:345:ASN:ND2	40:L3:347:SER:HB2	2.32	0.45
40:L3:65:SER:C	40:L3:67:PHE:N	2.90	0.45
46:L9:92:TYR:CD1	46:L9:142:ASP:HB3	3.93	0.45
47:M0:34:TYR:N	47:M0:34:TYR:HD1	2.13	0.45
49:M3:24:VAL:HG22	51:M5:199:LEU:HB2	4.97	0.45
58:N2:42:LYS:HA	58:N2:46:ALA:O	2.97	0.45
61:N5:135:ILE:O	61:N5:139:ILE:HG22	2.15	0.45
38:4:71:A:O2'	62:N6:52:ARG:NH2	2.48	0.45
67:O1:20:LEU:HD23	67:O1:20:LEU:HA	1.75	0.45
72:O6:26:ILE:HD12	72:O6:27:SER:H	1.82	0.45
72:O6:98:ARG:HB3	72:O6:99:ARG:HG3	4.94	0.45
73:O7:63:ARG:O	73:O7:64:MET:HB2	2.16	0.45
38:4:75:G:C8	75:O9:30:ARG:HG2	2.51	0.45
78:Q2:22:GLN:O	78:Q2:75:VAL:HG22	2.23	0.45
3:S1:59:ASP:C	3:S1:61:LEU:H	3.63	0.45
4:S2:162:CYS:O	4:S2:164:SER:N	2.50	0.45
6:S4:62:LYS:O	6:S4:66:MET:HG2	2.16	0.45
7:S5:119:ASP:O	7:S5:123:VAL:HG23	4.64	0.45
7:S5:216:GLU:CD	7:S5:219:ARG:HD2	2.37	0.45
8:S6:214:LYS:O	8:S6:218:GLU:HG3	2.49	0.45
11:S9:101:VAL:HG23	11:S9:102:GLU:OE2	2.17	0.45
11:S9:78:ARG:HH22	11:S9:82:ARG:NH2	2.13	0.45
34:SR:6:VAL:HG12	34:SR:316:MET:O	3.97	0.45
34:SR:42:LEU:HD21	34:SR:82:SER:HB3	1.99	0.45
36:1:1326:A:H2'	36:1:1327:C:O4'	2.17	0.45
36:1:1557:A:H5''	45:L8:54:GLU:OE1	2.16	0.45
36:1:3282:U:H2'	36:1:3283:U:O4'	2.16	0.45
36:1:3163:A:N6	36:1:3288:G:H1	2.14	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
86:1:3965:OHX:N6	86:1:4153:OHX:N4	2.64	0.45
36:1:564:G:H2'	36:1:565:U:C6	2.51	0.45
36:1:573:C:H2'	36:1:574:U:C6	2.51	0.45
1:2:141:U:C5	8:S6:179:VAL:HG23	2.50	0.45
1:2:1764:C:C5	1:2:1767:G:C4	3.04	0.45
1:2:422:G:OP1	86:2:2041:OHX:N6	2.50	0.45
1:2:344:A:H2'	1:2:345:U:H6	1.81	0.45
1:2:488:G:OP1	1:2:488:G:H4'	2.16	0.45
36:5:1047:A:C6	36:5:1048:A:C6	3.05	0.45
36:5:1131:G:H4'	36:5:1132:C:OP2	2.17	0.45
36:5:1632:A:C6	36:5:1644:C:C4	3.05	0.45
36:5:1648:A:H2'	36:5:1649:U:O4'	2.16	0.45
36:5:1936:A:H2'	36:5:1937:U:O4'	2.17	0.45
36:5:2152:A:H2'	36:5:2153:U:C6	2.51	0.45
36:5:230:U:H2'	36:5:231:G:O4'	2.17	0.45
36:5:2361:A:C6	36:5:2362:C:C4	3.03	0.45
36:5:310:U:H2'	36:5:311:C:O4'	2.17	0.45
36:5:3199:G:O6	86:5:4139:OHX:N5	2.50	0.45
36:5:953:G:C8	36:5:1117:G:C8	3.05	0.45
1:6:138:A:C2'	1:6:139:C:H5'	2.45	0.45
1:6:1756:A:H8	1:6:1756:A:OP2	2.00	0.45
1:6:291:G:H2'	1:6:292:U:C6	2.52	0.45
1:6:294:C:H2'	1:6:295:A:H8	1.81	0.45
1:6:639:U:H5	1:6:695:U:C5	2.35	0.45
14:C2:44:GLY:O	14:C2:46:ARG:N	3.68	0.45
14:C2:52:LEU:HD12	14:C2:78:LEU:O	2.16	0.45
15:C3:107:LYS:HD2	15:C3:107:LYS:HA	1.83	0.45
17:C5:82:ASN:H	17:C5:82:ASN:HD22	1.64	0.45
22:D0:93:LEU:HA	22:D0:93:LEU:HD23	1.70	0.45
23:D1:41:GLU:OE2	23:D1:41:GLU:N	3.22	0.45
23:D1:60:ARG:O	23:D1:63:GLY:N	3.32	0.45
39:L2:117:GLU:CD	39:L2:121:GLY:H	2.19	0.45
39:L2:15:ILE:HD12	39:L2:15:ILE:HA	4.61	0.45
39:L2:192:LYS:NZ	36:5:2181:C:OP1	201.04	0.45
39:L2:48:ILE:HD13	39:L2:57:PRO:HB2	1.98	0.45
40:L3:112:ASP:HA	40:L3:115:LYS:HB2	2.29	0.45
36:1:3003:G:H4'	40:L3:180:GLU:OE1	2.15	0.45
40:L3:77:THR:HG23	40:L3:326:GLY:O	2.17	0.45
42:L5:178:ASN:HA	42:L5:183:TRP:CG	2.80	0.45
43:L6:69:PHE:HB2	43:L6:138:GLN:NE2	2.58	0.45
44:L7:66:LYS:HG3	44:L7:76:TYR:CD2	2.50	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
45:L8:41:GLN:HG3	45:L8:44:ARG:NH1	2.94	0.45
52:M6:58:LEU:HD11	52:M6:74:ARG:HH21	1.82	0.45
64:N8:75:LEU:O	64:N8:77:LYS:N	2.89	0.45
66:O0:45:ALA:O	66:O0:48:THR:HG22	2.17	0.45
70:O4:42:PRO:HG3	70:O4:62:TYR:OH	3.14	0.45
70:O4:47:CYS:HB3	70:O4:84:CYS:SG	2.56	0.45
71:O5:21:LEU:O	71:O5:24:LEU:N	3.18	0.45
74:O8:58:ASP:HB3	74:O8:61:LYS:CG	4.29	0.45
75:O9:35:ILE:H	75:O9:35:ILE:HD12	3.96	0.45
78:Q2:40:LYS:HE3	78:Q2:44:ASP:OD2	2.17	0.45
79:Q3:85:ARG:NH2	1:6:923:A:OP1	249.62	0.45
3:S1:125:VAL:HG11	3:S1:173:THR:HG23	3.93	0.45
4:S2:139:ILE:HD12	4:S2:191:ALA:HB1	1.97	0.45
4:S2:78:ASP:O	4:S2:79:GLU:HB3	2.15	0.45
5:S3:177:MET:HG3	5:S3:178:ARG:H	4.71	0.45
5:S3:43:PRO:O	5:S3:44:THR:HG22	3.82	0.45
6:S4:108:ARG:H	6:S4:108:ARG:HG2	3.79	0.45
7:S5:102:ARG:HG3	7:S5:103:ASN:H	1.81	0.45
8:S6:10:ASN:ND2	8:S6:128:THR:HG22	4.76	0.45
9:S7:41:LEU:HB3	9:S7:70:PHE:CE2	4.10	0.45
11:S9:6:ARG:HA	11:S9:6:ARG:HD3	1.56	0.45
34:SR:103:PHE:CZ	34:SR:122:ILE:HD12	2.51	0.45
34:SR:115:ILE:HG13	34:SR:121:MET:O	2.45	0.45
36:1:1118:C:O2	36:1:1154:A:H2	1.99	0.45
36:1:1313:G:N3	36:1:1318:A:H2	2.14	0.45
36:1:1341:U:H2'	36:1:1342:C:H6	1.81	0.45
36:1:1547:G:OP2	51:M5:105:ARG:NH1	2.48	0.45
36:1:1573:G:N2	36:1:1574:C:O2'	2.49	0.45
36:1:1722:U:H2'	36:1:1723:A:O4'	2.17	0.45
36:1:2205:U:H5'	36:1:2206:G:OP2	2.17	0.45
36:1:259:C:H2'	36:1:260:C:H6	1.82	0.45
36:1:3204:C:OP1	50:M4:98:SER:OG	2.31	0.45
86:1:4023:OHX:N2	86:1:4143:OHX:N1	2.65	0.45
1:2:1017:U:H2'	1:2:1018:U:C6	2.52	0.45
1:2:1163:A:N6	1:2:1164:G:C6	2.84	0.45
1:2:1317:C:O2	1:2:1400:A:H2	1.99	0.45
1:2:1383:G:OP1	22:D0:87:HIS:ND1	2.47	0.45
1:2:1608:U:H2'	1:2:1609:U:H6	1.82	0.45
1:2:1793:G:H1'	1:2:1794:A:H2'	1.98	0.45
1:2:274:G:H3'	1:2:275:C:C6	2.51	0.45
1:2:795:U:C5	1:2:796:A:C5	3.04	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:4:84:C:H5'	38:4:85:G:C5	2.52	0.45
36:5:1794:G:O2'	36:5:1795:U:H5'	2.15	0.45
36:5:2658:G:C6	36:5:2659:G:N7	2.85	0.45
78:Q2:45:ARG:NH2	36:5:283:G:OP2	146.70	0.45
36:5:3027:A:OP1	36:5:3027:A:H8	1.99	0.45
36:5:3096:C:H2'	36:5:3097:C:H6	1.82	0.45
36:5:3112:G:N7	86:5:3910:OHX:N6	2.63	0.45
36:5:3165:A:H2'	36:5:3166:C:C6	2.51	0.45
1:6:1388:A:C5	1:6:1411:A:C6	3.05	0.45
21:C9:68:ARG:NH1	1:6:1523:G:N7	414.67	0.45
11:S9:124:HIS:HD2	1:6:479:C:H5'	453.50	0.45
38:8:141:C:OP1	86:8:229:OHX:N6	2.50	0.45
12:C0:56:LYS:HG2	12:C0:67:THR:HB	1.98	0.45
13:C1:13:PHE:CE2	13:C1:15:LYS:HB3	2.51	0.45
14:C2:55:GLY:HA2	14:C2:85:LYS:HD3	1.97	0.45
18:C6:23:LYS:O	18:C6:64:ASP:N	2.42	0.45
21:C9:37:VAL:HG11	21:C9:100:ILE:HD11	2.33	0.45
23:D1:72:LEU:HD23	23:D1:72:LEU:HA	2.93	0.45
24:D2:77:PRO:HD3	25:D3:7:ARG:HB2	5.61	0.45
40:L3:315:GLY:HA2	36:5:3379:C:H4'	214.82	0.45
41:L4:118:LYS:O	41:L4:121:ALA:HB3	2.27	0.45
41:L4:84:ARG:O	41:L4:87:GLN:HB2	2.93	0.45
43:L6:175:LYS:O	43:L6:176:PHE:HB2	4.14	0.45
44:L7:173:LEU:HD11	44:L7:201:PHE:HB2	2.95	0.45
45:L8:134:TYR:CG	45:L8:190:VAL:HG21	2.51	0.45
46:L9:110:LYS:HB3	46:L9:128:VAL:HB	2.11	0.45
46:L9:38:LEU:HA	46:L9:38:LEU:HD23	2.01	0.45
47:M0:196:PHE:CG	47:M0:197:VAL:N	2.84	0.45
48:M1:115:LYS:HB2	48:M1:115:LYS:NZ	2.31	0.45
48:M1:89:TYR:O	48:M1:169:ALA:HB1	2.27	0.45
35:SM:24:GLU:HG2	48:M1:64:LYS:HZ3	1.82	0.45
48:M1:78:GLU:O	48:M1:82:ARG:HB3	2.16	0.45
50:M4:120:VAL:O	50:M4:124:ARG:HG3	2.17	0.45
53:M7:3:ARG:CG	53:M7:3:ARG:HH21	3.86	0.45
36:1:973:A:OP2	54:M8:12:ARG:NH1	2.49	0.45
58:N2:19:VAL:O	58:N2:22:PRO:HD2	2.33	0.45
59:N3:24:ASN:HB2	59:N3:98:ASN:O	2.97	0.45
62:N6:2:ALA:N	36:5:213:A:OP1	80.74	0.45
63:N7:27:LYS:HD2	63:N7:28:PRO:HD2	1.98	0.45
64:N8:110:GLY:O	64:N8:129:PHE:HB2	2.63	0.45
64:N8:2:PRO:N	64:N8:5:PHE:HD2	2.15	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
66:O0:34:LEU:HD13	66:O0:34:LEU:HA	2.96	0.45
67:O1:31:ARG:O	67:O1:35:GLU:HB2	2.19	0.45
69:O3:102:LEU:HA	69:O3:102:LEU:HD23	1.65	0.45
71:O5:76:GLN:O	71:O5:81:ARG:NH1	2.49	0.45
72:O6:56:ARG:HH22	72:O6:76:ARG:NH1	2.21	0.45
74:O8:68:SER:O	74:O8:69:LEU:HD23	5.92	0.45
77:Q1:4:LYS:O	77:Q1:7:LYS:HB3	2.15	0.45
2:S0:154:GLU:OE1	2:S0:154:GLU:N	2.34	0.45
3:S1:69:CYS:SG	16:C4:114:ARG:NH1	3.70	0.45
4:S2:123:GLY:HA2	4:S2:126:ARG:HH12	3.02	0.45
5:S3:42:THR:HG23	5:S3:45:LYS:O	4.78	0.45
5:S3:57:ASP:O	5:S3:65:ARG:HG2	4.58	0.45
11:S9:137:GLY:C	11:S9:139:GLN:H	2.19	0.45
36:1:1170:A:H2'	36:1:1171:G:O4'	2.16	0.45
36:1:12:A:H1'	61:N5:37:THR:HG21	1.99	0.45
36:1:1498:A:H2'	36:1:1499:C:C6	2.51	0.45
36:1:2261:G:H21	36:1:2262:A:N6	2.15	0.45
36:1:184:U:H3	36:1:232:G:H1	1.63	0.45
36:1:2571:U:OP1	36:1:2571:U:H2'	2.16	0.45
36:1:288:C:H2'	36:1:289:A:C8	2.52	0.45
36:1:637:C:H2'	36:1:637:C:H6	1.40	0.45
1:2:1098:U:H4'	1:2:1099:U:OP2	2.16	0.45
1:2:1211:A:N6	1:2:1452:U:H3	2.14	0.45
1:2:1746:A:H2'	1:2:1747:G:O4'	2.16	0.45
1:2:1796:C:H4'	1:2:1797:A:OP2	2.16	0.45
1:2:580:A:O2'	1:2:582:U:OP1	2.35	0.45
1:2:751:G:H2'	1:2:752:A:H8	1.81	0.45
39:L2:70:ARG:HB3	36:5:1650:G:H5''	179.40	0.45
36:5:1783:U:H2'	36:5:1784:G:H8	1.79	0.45
36:5:2359:C:O2'	36:5:2360:C:H5'	2.17	0.45
36:5:2400:G:H5''	36:5:2401:A:OP2	2.17	0.45
36:5:277:G:H2'	36:5:278:U:C6	2.52	0.45
51:M5:98:LEU:HD13	36:5:290:G:OP1	136.00	0.45
36:5:2374:C:N4	36:5:2941:A:N3	2.64	0.45
36:5:3167:A:H2'	36:5:3168:A:O4'	2.17	0.45
36:5:3287:U:H2'	36:5:3288:G:C5'	2.46	0.45
36:5:508:U:H2'	36:5:509:U:C6	2.50	0.45
36:5:63:A:H8	36:5:63:A:O5'	1.98	0.45
36:5:726:G:H5'	36:5:727:G:P	2.56	0.45
36:5:912:G:C2	36:5:914:A:C2	3.05	0.45
1:6:1039:A:O2'	1:6:1040:G:OP2	2.31	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:1143:A:H2'	1:6:1144:U:H6	1.81	0.45
1:6:1258:U:C4	1:6:1259:U:C4	3.04	0.45
1:6:1207:C:N3	1:6:1456:C:H5	2.15	0.45
1:6:1531:G:C6	1:6:1532:U:C4	3.05	0.45
1:6:1774:G:H2'	1:6:1775:U:O4'	2.16	0.45
1:6:538:A:C8	1:6:543:C:N4	2.73	0.45
32:E0:13:LYS:HE3	1:6:566:C:O2	374.61	0.45
1:6:686:C:H2'	1:6:687:G:C8	2.51	0.45
37:7:8:G:C6	37:7:9:C:C4	3.04	0.45
86:8:216:OHX:N2	86:8:223:OHX:N4	2.64	0.45
38:8:82:U:H2'	38:8:83:C:H5'	1.97	0.45
12:C0:49:LEU:HB3	12:C0:55:VAL:CG1	2.46	0.45
12:C0:70:GLU:O	12:C0:73:VAL:HG22	5.01	0.45
16:C4:117:ASP:OD1	16:C4:118:VAL:N	2.50	0.45
17:C5:87:PRO:O	17:C5:90:ILE:HB	2.16	0.45
1:2:1340:U:O4	18:C6:9:THR:HA	2.16	0.45
18:C6:129:PHE:CE1	22:D0:78:THR:HA	2.52	0.45
23:D1:13:VAL:HA	23:D1:14:PRO:HD3	1.83	0.45
28:D6:10:ARG:NH1	28:D6:36:ILE:HA	2.34	0.45
28:D6:84:VAL:O	28:D6:86:VAL:N	2.38	0.45
39:L2:64:ARG:HH22	45:L8:38:GLN:HA	3.06	0.45
41:L4:341:SER:O	41:L4:342:LYS:CB	4.25	0.45
42:L5:111:GLN:HG3	42:L5:116:ASP:OD2	2.17	0.45
42:L5:22:ARG:NH2	42:L5:28:THR:OG1	2.49	0.45
44:L7:176:TYR:HB3	44:L7:194:HIS:CD2	3.50	0.45
47:M0:14:ASN:O	47:M0:128:ARG:NH2	2.49	0.45
50:M4:108:ARG:NH2	52:M6:196:ALA:O	2.50	0.45
50:M4:113:THR:HG22	50:M4:116:GLU:HB2	3.20	0.45
49:M3:28:GLN:HB3	51:M5:201:ARG:HD3	2.30	0.45
55:M9:106:LEU:HB3	55:M9:120:TYR:HE1	2.01	0.45
55:M9:109:TYR:OH	55:M9:139:VAL:HG22	2.15	0.45
36:1:2631:U:OP2	57:N1:4:SER:OG	2.35	0.45
59:N3:127:PRO:HA	59:N3:130:ALA:HB3	2.65	0.45
63:N7:97:SER:HB2	63:N7:99:GLU:HG3	1.98	0.45
65:N9:43:HIS:CE1	65:N9:47:LEU:HD11	3.06	0.45
75:O9:45:ARG:NH2	36:5:1841:A:N3	126.61	0.45
79:Q3:26:VAL:HG13	79:Q3:30:GLU:HG3	2.09	0.45
79:Q3:77:ALA:C	79:Q3:79:VAL:H	2.69	0.45
2:S0:112:THR:CG2	2:S0:115:PHE:HB2	2.85	0.45
2:S0:88:LYS:HA	2:S0:88:LYS:HE2	1.98	0.45
3:S1:82:ARG:HH11	3:S1:82:ARG:HB2	4.81	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:S2:84:LYS:HE3	4:S2:84:LYS:HB2	4.22	0.45
5:S3:46:THR:O	5:S3:84:ILE:HG12	2.38	0.45
8:S6:76:LEU:HD13	8:S6:92:ARG:HD2	1.98	0.45
1:2:197:A:N1	10:S8:138:ASN:ND2	2.64	0.45
10:S8:12:SER:O	10:S8:14:THR:N	3.21	0.45
34:SR:201:THR:HB	34:SR:241:PHE:O	4.30	0.45
34:SR:38:ARG:HB3	34:SR:67:ILE:HG12	1.97	0.45
36:1:114:A:O2'	51:M5:50:ARG:HB3	2.17	0.45
36:1:1461:A:O2'	36:1:1462:A:H5'	2.16	0.45
36:1:1530:U:OP1	86:1:3935:OHX:N2	2.49	0.45
36:1:1602:A:H5''	55:M9:38:ARG:HG3	1.98	0.45
36:1:2633:U:H2'	36:1:2634:U:O4'	2.17	0.45
36:1:2775:U:H2'	36:1:2776:C:C6	2.51	0.45
36:1:283:G:O6	36:1:304:G:H1'	2.17	0.45
36:1:2961:G:C6	36:1:2962:U:C4	3.05	0.45
36:1:3178:A:H5''	36:1:3179:U:OP1	2.17	0.45
36:1:63:A:H2'	36:1:64:G:O4'	2.16	0.45
36:1:718:G:N2	36:1:721:G:H1'	2.31	0.45
1:2:1178:G:H2'	1:2:1179:G:O4'	2.16	0.45
86:2:2036:OHX:N2	10:S8:17:LYS:O	2.50	0.45
1:2:411:C:O2	1:2:423:G:N2	2.49	0.45
1:2:546:U:H2'	1:2:547:U:H6	1.81	0.45
36:5:1336:U:O2'	36:5:1337:A:H5'	2.17	0.45
36:5:1348:U:H5'	36:5:1349:G:OP1	2.17	0.45
36:5:1549:U:H2'	36:5:1550:C:H6	1.82	0.45
36:5:1672:U:O2	36:5:1776:G:C2	2.69	0.45
36:5:1838:G:O5'	36:5:1838:G:H8	1.99	0.45
53:M7:68:GLY:HA3	36:5:2350:C:H5''	175.50	0.45
36:5:253:A:HO2'	36:5:254:A:H8	1.62	0.45
36:5:114:A:C2	36:5:267:G:C4	3.04	0.45
57:N1:8:ARG:NH2	36:5:2756:C:O2	248.65	0.45
43:L6:69:PHE:CZ	36:5:3267:A:H2'	258.40	0.45
36:5:3119:U:OP2	86:5:3910:OHX:N3	2.50	0.45
86:5:3994:OHX:N6	86:5:4085:OHX:N5	2.65	0.45
49:M3:171:ARG:HD3	36:5:770:G:OP1	143.42	0.45
64:N8:12:ARG:O	36:5:944:C:H5'	162.60	0.45
1:6:100:A:C2	1:6:101:U:C2	3.04	0.45
1:6:1769:U:OP2	86:6:2144:OHX:N2	2.50	0.45
1:6:701:U:H2'	1:6:702:G:H8	1.81	0.45
1:6:74:U:H5''	1:6:75:U:OP2	2.16	0.45
1:6:915:A:H5''	1:6:916:U:OP2	2.17	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:C2:68:GLU:C	14:C2:70:ASN:H	2.19	0.45
15:C3:94:LYS:O	15:C3:97:SER:N	2.94	0.45
16:C4:114:ARG:HB2	28:D6:59:TYR:CE2	2.51	0.45
16:C4:20:TYR:OH	16:C4:86:THR:HA	2.16	0.45
19:C7:63:LYS:HD3	19:C7:63:LYS:HA	1.63	0.45
21:C9:45:MET:HE3	21:C9:46:PRO:HD2	2.01	0.45
23:D1:38:LYS:HB2	23:D1:38:LYS:HE3	4.73	0.45
24:D2:52:TYR:HE2	24:D2:54:ASP:HA	3.51	0.45
24:D2:6:VAL:HG12	24:D2:7:LEU:N	3.08	0.45
25:D3:41:SER:HA	25:D3:42:PRO:HD3	1.75	0.45
12:C0:38:LYS:NZ	31:D9:4:GLU:HG3	2.32	0.45
32:E0:30:PRO:O	32:E0:35:TYR:HB2	2.17	0.45
33:E1:103:LEU:HD13	33:E1:106:TYR:HE1	1.81	0.45
33:E1:111:GLU:HA	33:E1:112:GLY:HA2	1.78	0.45
40:L3:345:ASN:HD21	40:L3:347:SER:HB2	1.82	0.45
41:L4:33:ASP:O	41:L4:37:THR:HG23	2.17	0.45
43:L6:131:LYS:O	43:L6:134:ARG:N	2.45	0.45
44:L7:158:LYS:CG	44:L7:159:GLN:H	2.88	0.45
44:L7:24:GLU:C	44:L7:26:VAL:H	2.19	0.45
45:L8:26:LEU:HD11	63:N7:123:GLN:NE2	5.07	0.45
46:L9:175:PHE:N	46:L9:175:PHE:CD2	3.29	0.45
47:M0:41:ALA:O	47:M0:139:ARG:NH2	2.46	0.45
49:M3:128:ARG:HD3	71:O5:114:ARG:CZ	2.47	0.45
50:M4:133:LYS:NZ	36:5:3227:A:HO2'	302.50	0.45
50:M4:18:GLY:O	50:M4:69:THR:HA	2.50	0.45
50:M4:24:LYS:HE2	50:M4:25:LYS:HZ2	1.81	0.45
53:M7:78:VAL:HG12	53:M7:80:LYS:H	1.82	0.45
56:N0:100:VAL:HG13	56:N0:101:ALA:H	1.80	0.45
57:N1:96:ILE:HA	57:N1:96:ILE:HD13	1.59	0.45
58:N2:89:LEU:HD23	58:N2:92:TRP:CE3	2.52	0.45
36:1:2916:U:C1'	59:N3:44:SER:HB3	2.42	0.45
62:N6:100:HIS:CE1	62:N6:102:SER:HG	4.62	0.45
70:O4:51:LEU:HA	70:O4:51:LEU:HD23	3.90	0.45
71:O5:21:LEU:HD22	71:O5:25:LYS:HG3	1.99	0.45
73:O7:50:GLY:O	73:O7:53:ALA:HB3	2.17	0.45
2:S0:165:ARG:HD3	2:S0:165:ARG:HA	1.80	0.45
3:S1:209:ASN:O	3:S1:210:ILE:HB	2.17	0.45
3:S1:76:SER:OG	3:S1:78:ASP:OD1	2.29	0.45
4:S2:137:ILE:HD12	4:S2:215:PHE:CZ	5.45	0.45
4:S2:153:SER:OG	4:S2:171:PRO:HA	2.58	0.45
6:S4:213:SER:OG	6:S4:214:LEU:HD12	4.74	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:S4:92:LEU:HB2	6:S4:95:THR:CG2	4.20	0.45
7:S5:184:PHE:CD1	7:S5:185:ARG:HG3	3.06	0.45
1:2:339:C:OP2	10:S8:10:LYS:NZ	2.49	0.45
10:S8:165:LEU:HB3	10:S8:183:ILE:HD13	3.74	0.45
11:S9:115:LYS:HD2	11:S9:115:LYS:HA	1.66	0.45
34:SR:171:SER:OG	34:SR:179:LYS:HB2	2.15	0.45
36:1:1341:U:H2'	36:1:1342:C:C6	2.52	0.45
36:1:1366:A:N3	36:1:1366:A:H2'	2.31	0.45
36:1:1427:U:OP2	64:N8:4:ARG:NH2	2.47	0.45
36:1:1838:G:H4'	36:1:1839:A:N3	2.32	0.45
36:1:2249:G:C8	36:1:2249:G:H3'	2.51	0.45
36:1:2544:U:H2'	36:1:2545:C:C6	2.52	0.45
36:1:891:G:H2'	36:1:892:U:O4'	2.17	0.45
1:2:1201:G:H22	1:2:1600:A:H5''	1.81	0.45
1:2:393:C:H4'	1:2:1673:G:O2'	2.16	0.45
1:2:169:A:OP1	8:S6:137:ARG:HG3	2.16	0.45
1:2:651:G:C2	1:2:684:A:C6	3.05	0.45
57:N1:127:GLN:HG3	36:5:1095:U:H3	260.46	0.45
36:5:1134:G:C2	36:5:1135:A:N7	2.84	0.45
36:5:1235:U:H4'	36:5:1236:G:H5'	1.98	0.45
36:5:2429:G:C2	36:5:2601:A:C2	3.05	0.45
78:Q2:56:PRO:HB2	36:5:2802:A:H5'	187.48	0.45
36:5:2881:C:C2	36:5:2882:U:C5	3.05	0.45
36:5:3364:C:OP1	86:5:3935:OHX:N1	2.49	0.45
36:5:1656:A:O2'	86:5:4172:OHX:N2	2.49	0.45
36:5:624:G:C2	36:5:625:G:C5	3.04	0.45
36:5:814:U:C2	36:5:815:G:C8	3.04	0.45
1:6:1124:A:O2'	1:6:1125:A:H5'	2.17	0.45
1:6:1450:U:H2'	1:6:1451:C:C6	2.52	0.45
1:6:309:C:H2'	1:6:310:C:H6	1.82	0.45
1:6:417:A:H5'	1:6:418:G:C4	2.52	0.45
1:6:95:G:H2'	1:6:95:G:N3	2.31	0.45
1:6:868:G:H1	1:6:960:U:H3	1.65	0.45
38:8:116:G:H1	38:8:137:C:H42	1.65	0.45
13:C1:72:THR:HA	13:C1:124:THR:HA	1.98	0.45
13:C1:55:ASP:HB2	13:C1:82:ARG:CZ	2.47	0.45
15:C3:54:LEU:HB3	15:C3:60:VAL:HG21	1.98	0.45
24:D2:103:ILE:O	24:D2:103:ILE:HG12	2.16	0.45
24:D2:46:TYR:CD1	24:D2:69:LEU:HD13	2.52	0.45
26:D4:123:LYS:HZ2	26:D4:123:LYS:HG3	1.31	0.45
7:S5:57:SER:HB2	30:D8:53:ILE:O	3.17	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:1235:C:H2'	33:E1:138:ARG:NH2	2.32	0.45
39:L2:125:ALA:C	39:L2:126:LEU:HD23	2.84	0.45
40:L3:185:GLY:O	40:L3:191:LYS:NZ	2.97	0.45
41:L4:182:LEU:C	41:L4:184:SER:H	2.20	0.45
41:L4:193:LYS:HB2	41:L4:193:LYS:HE3	1.50	0.45
41:L4:169:LEU:HD12	41:L4:219:LEU:HD21	1.99	0.45
41:L4:313:LEU:HA	41:L4:313:LEU:HD22	1.71	0.45
42:L5:211:LEU:HD13	42:L5:219:PHE:HA	3.38	0.45
45:L8:146:LYS:O	45:L8:146:LYS:HE2	6.74	0.45
46:L9:103:ILE:HG13	46:L9:136:PHE:HE2	1.82	0.45
47:M0:189:GLU:HB3	47:M0:200:LEU:CB	2.46	0.45
48:M1:81:GLU:O	48:M1:83:GLY:N	3.17	0.45
49:M3:2:ALA:N	64:N8:31:GLY:O	4.19	0.45
51:M5:137:PRO:O	51:M5:143:ARG:NH1	2.50	0.45
51:M5:183:THR:O	51:M5:183:THR:HG23	2.17	0.45
54:M8:16:ARG:NH2	54:M8:20:LYS:HB2	2.85	0.45
66:O0:99:ASP:O	66:O0:101:LEU:N	2.74	0.45
67:O1:75:ILE:HG23	67:O1:93:VAL:HG13	4.75	0.45
38:4:113:U:N3	75:O9:7:PHE:CE2	2.85	0.45
79:Q3:53:GLY:HA2	79:Q3:67:GLY:O	3.13	0.45
3:S1:130:SER:OG	3:S1:180:THR:N	2.50	0.45
4:S2:53:ILE:O	4:S2:56:ILE:N	2.50	0.45
10:S8:11:ARG:NH1	10:S8:15:GLY:O	3.07	0.45
11:S9:110:GLN:HE22	11:S9:126:ARG:HG2	1.81	0.45
34:SR:164:ASP:O	34:SR:166:SER:N	3.26	0.45
34:SR:95:ALA:O	34:SR:96:THR:HG22	5.35	0.45
36:1:1340:G:H2'	36:1:1341:U:C6	2.52	0.45
36:1:1517:G:OP1	75:O9:41:ARG:NH2	2.29	0.45
36:1:1578:C:H3'	36:1:1579:C:C6	2.51	0.45
36:1:1554:U:O2'	36:1:1582:C:H5	1.98	0.45
36:1:246:U:H2'	36:1:247:C:H6	1.82	0.45
36:1:3068:U:OP1	55:M9:59:SER:OG	2.32	0.45
36:1:1184:A:OP2	86:1:4059:OHX:N3	2.49	0.45
36:1:595:G:H2'	36:1:596:C:C6	2.51	0.45
1:2:1081:A:O2'	1:2:1082:C:O2	2.35	0.45
1:2:1291:G:N2	1:2:1324:G:H22	2.14	0.45
1:2:1595:U:H5	1:2:1596:C:C5	2.34	0.45
1:2:48:G:C6	1:2:432:G:C2	3.05	0.45
1:2:55:A:C2	1:2:403:G:C5	3.05	0.45
1:2:800:U:H2'	1:2:801:G:H8	1.82	0.45
1:2:892:A:C6	1:2:893:U:C4	3.04	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:915:A:H2'	1:2:915:A:N3	2.31	0.45
36:5:1238:C:H2'	36:5:1239:C:H6	1.82	0.45
36:5:138:U:H2'	36:5:139:G:H8	1.82	0.45
36:5:2213:A:N1	36:5:2429:G:H1'	2.32	0.45
36:5:2518:C:H2'	36:5:2519:A:H8	1.81	0.45
36:5:2877:G:H2'	36:5:2878:G:O4'	2.16	0.45
36:5:3254:G:C6	36:5:3255:U:C4	3.05	0.45
36:5:3330:A:H8	36:5:3330:A:H5''	1.81	0.45
36:5:607:A:N3	36:5:607:A:H2'	2.32	0.45
36:5:900:G:H2'	36:5:901:G:C8	2.51	0.45
1:6:1050:G:O6	86:6:2195:OHX:N4	2.49	0.45
1:6:1172:G:H2'	1:6:1173:C:O4'	2.17	0.45
1:6:119:A:OP2	1:6:120:U:H5	1.99	0.45
19:C7:49:LYS:HA	1:6:1389:C:H4'	422.94	0.45
31:D9:33:LYS:HE3	1:6:1594:G:H5'	410.12	0.45
1:6:1030:A:C5	1:6:1792:G:C6	3.05	0.45
1:6:1314:U:OP2	86:6:2184:OHX:N4	2.50	0.45
1:6:83:G:H8	1:6:83:G:O5'	2.00	0.45
15:C3:20:ARG:NE	1:6:862:A:OP1	356.87	0.45
37:7:112:G:H2'	37:7:113:C:C6	2.51	0.45
13:C1:83:THR:HG21	1:6:325:G:H4'	290.27	0.45
15:C3:70:LYS:HE2	15:C3:70:LYS:HB3	4.37	0.45
16:C4:41:ARG:O	16:C4:42:VAL:HG22	2.16	0.45
16:C4:84:ARG:HB3	16:C4:118:VAL:HG23	3.25	0.45
20:C8:11:PHE:CD2	20:C8:59:GLY:HA3	2.39	0.45
26:D4:63:GLN:HG3	26:D4:64:PHE:O	2.55	0.45
16:C4:112:ILE:O	28:D6:58:VAL:HG22	2.17	0.45
33:E1:88:PRO:HA	33:E1:89:LYS:HA	4.66	0.45
39:L2:95:SER:O	39:L2:97:ASN:N	3.49	0.45
41:L4:301:PRO:O	41:L4:302:ALA:HB3	4.61	0.45
41:L4:316:ASN:OD1	41:L4:318:LEU:HB2	2.17	0.45
41:L4:3:ARG:HA	41:L4:4:PRO:HD2	1.54	0.45
42:L5:95:TRP:CH2	42:L5:161:GLY:HA2	2.52	0.45
44:L7:38:LYS:HE3	44:L7:38:LYS:HB2	1.69	0.45
36:1:2561:A:N1	45:L8:32:LYS:HB2	2.32	0.45
45:L8:73:PRO:O	45:L8:77:GLN:N	4.88	0.45
49:M3:192:GLU:O	49:M3:194:GLU:N	2.60	0.45
52:M6:94:ARG:O	52:M6:97:ALA:HB3	2.50	0.45
53:M7:101:ASN:O	53:M7:105:LYS:HG3	2.17	0.45
61:N5:82:LEU:HD11	61:N5:135:ILE:HD12	3.52	0.45
62:N6:39:LEU:O	62:N6:43:TYR:N	3.04	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:M3:64:LYS:HA	64:N8:69:TRP:CE3	3.17	0.45
66:O0:16:LEU:HA	66:O0:16:LEU:HD22	1.73	0.45
70:O4:24:LYS:HB2	70:O4:24:LYS:HE3	1.78	0.45
73:O7:72:ARG:HH11	38:8:94:C:H3'	49.90	0.45
74:O8:16:ARG:HB3	74:O8:17:ARG:H	4.44	0.45
75:O9:5:LYS:HD3	75:O9:13:MET:HE1	1.98	0.45
76:Q0:79:GLU:HA	76:Q0:80:PRO:HD3	1.75	0.45
77:Q1:21:ARG:HH11	1:6:1654:G:P	283.67	0.45
3:S1:116:LYS:HG3	1:6:931:C:H5''	321.44	0.45
3:S1:28:GLU:HG3	3:S1:29:TRP:H	1.81	0.45
3:S1:40:ASN:OD1	3:S1:73:LEU:HA	2.16	0.45
1:2:3:U:H5''	4:S2:198:THR:O	2.15	0.45
7:S5:142:PRO:HG2	7:S5:170:GLN:NE2	2.74	0.45
7:S5:143:ARG:HD3	30:D8:55:VAL:HG11	2.13	0.45
35:SM:66:ALA:O	35:SM:70:ASN:HB2	2.17	0.45
34:SR:152:SER:OG	34:SR:153:GLN:N	2.48	0.45
34:SR:66:HIS:HB3	34:SR:85:TRP:HB2	1.97	0.45
36:1:1002:A:H2'	36:1:1003:A:H8	1.81	0.45
36:1:1048:A:H2'	47:M0:22:TYR:CZ	2.52	0.45
36:1:1077:U:O2'	36:1:1078:U:H5'	2.16	0.45
36:1:1354:G:O5'	36:1:1354:G:H8	2.00	0.45
36:1:1565:G:H2'	36:1:1566:A:H8	1.82	0.45
36:1:1607:U:O2'	36:1:1608:C:H5'	2.17	0.45
36:1:1877:U:OP2	86:1:3920:OHX:N2	2.50	0.45
36:1:1902:G:C6	36:1:1903:U:C2	3.05	0.45
36:1:1907:C:C5	36:1:1908:A:C5	3.05	0.45
36:1:2112:U:O2'	86:1:3952:OHX:N1	2.50	0.45
36:1:2399:A:N6	36:1:2400:G:C6	2.85	0.45
36:1:2799:A:H1'	64:N8:42:ARG:NH2	2.31	0.45
36:1:3107:U:OP1	76:Q0:114:LYS:NZ	2.36	0.45
86:1:3862:OHX:N2	73:O7:46:SER:HB3	2.32	0.45
36:1:835:G:O2'	36:1:857:G:N2	2.37	0.45
1:2:1062:A:H2'	1:2:1063:U:O4'	2.17	0.45
1:2:1241:G:H1'	17:C5:79:HIS:CD2	2.51	0.45
1:2:1542:G:H5''	21:C9:88:VAL:N	2.32	0.45
86:2:2089:OHX:N3	86:2:2131:OHX:N6	2.65	0.45
38:4:11:C:H2'	38:4:12:A:O4'	2.17	0.45
36:5:1807:G:C6	36:5:1808:G:N1	2.85	0.45
36:5:2209:U:H4'	36:5:2210:G:OP1	2.15	0.45
36:5:2516:U:O2	36:5:2594:C:N4	2.49	0.45
36:5:2822:U:H2'	36:5:2823:G:O4'	2.17	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:3051:U:C2	36:5:3052:G:C8	3.04	0.45
36:5:3189:G:C2	36:5:3190:C:C2	3.05	0.45
36:5:3375:A:C8	36:5:3378:C:H5	2.35	0.45
36:5:784:A:HO2'	36:5:785:G:P	2.34	0.45
36:5:900:G:C6	36:5:901:G:C6	3.05	0.45
36:5:916:G:N7	36:5:924:G:C6	2.85	0.45
36:5:979:U:H4'	36:5:980:A:H5'	1.98	0.45
1:6:1360:A:C3'	1:6:1361:U:H4'	2.47	0.45
1:6:1514:U:H4'	1:6:1515:A:C4	2.52	0.45
1:6:1716:C:O2'	1:6:1717:G:O5'	2.29	0.45
38:8:41:A:N7	38:8:42:G:C8	2.85	0.45
24:D2:7:LEU:HD11	24:D2:37:PHE:HD2	1.82	0.45
20:C8:25:ASN:O	27:D5:40:VAL:HG21	3.90	0.45
27:D5:80:LEU:HD23	27:D5:80:LEU:HA	1.73	0.45
28:D6:10:ARG:HH11	28:D6:36:ILE:HG13	5.26	0.45
39:L2:237:LEU:HG	39:L2:237:LEU:H	2.19	0.45
39:L2:65:ASP:HB3	39:L2:68:LYS:O	2.20	0.45
40:L3:371:GLN:HG3	60:N4:14:TYR:CE1	3.92	0.45
42:L5:184:ASP:OD2	42:L5:187:THR:OG1	3.60	0.45
42:L5:40:HIS:ND1	57:N1:69:LYS:HA	2.72	0.45
43:L6:172:HIS:HB3	69:O3:44:TYR:CE2	2.52	0.45
44:L7:80:GLN:OE1	57:N1:136:ARG:HB2	5.19	0.45
46:L9:92:TYR:N	46:L9:92:TYR:CD2	4.08	0.45
51:M5:49:ARG:HA	51:M5:53:TYR:HB3	1.98	0.45
52:M6:125:ARG:HG3	52:M6:129:LEU:HD23	3.50	0.45
53:M7:41:LEU:N	53:M7:113:TYR:HA	2.32	0.45
55:M9:99:LEU:HD11	55:M9:103:ARG:CZ	2.47	0.45
55:M9:36:ASN:OD1	55:M9:37:SER:N	2.49	0.45
55:M9:47:ASN:OD1	55:M9:49:THR:HG23	6.31	0.45
55:M9:60:LYS:O	55:M9:64:ARG:HG3	2.17	0.45
44:L7:75:TYR:CD1	56:N0:60:SER:HB2	2.60	0.45
57:N1:108:ARG:O	57:N1:112:ASN:HB2	3.02	0.45
59:N3:66:LYS:HB2	59:N3:69:LEU:HD22	1.99	0.45
62:N6:56:VAL:HG23	62:N6:106:ILE:HA	2.79	0.45
64:N8:135:GLU:O	64:N8:135:GLU:HG2	2.51	0.45
64:N8:74:ASN:CB	64:N8:76:ASP:HB3	2.94	0.45
71:O5:54:VAL:O	71:O5:58:ILE:HG13	2.63	0.45
71:O5:57:VAL:O	71:O5:61:GLN:HG3	2.42	0.45
72:O6:29:LYS:O	36:5:266:A:N6	100.21	0.45
2:S0:151:SER:HA	2:S0:152:PRO:HD2	2.09	0.45
3:S1:97:LEU:HA	3:S1:97:LEU:HD13	1.78	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:S2:54:GLU:H	4:S2:54:GLU:HG2	1.52	0.45
5:S3:168:ILE:HG22	5:S3:189:MET:HB2	1.98	0.45
6:S4:136:VAL:HG11	6:S4:148:ARG:CZ	2.47	0.45
11:S9:65:LYS:HA	11:S9:70:LEU:HD11	1.99	0.45
35:SM:102:THR:CG2	35:SM:105:LYS:HB2	2.47	0.45
35:SM:61:ILE:H	35:SM:61:ILE:HG13	1.47	0.45
34:SR:210:LEU:HA	34:SR:210:LEU:HD23	1.79	0.45
34:SR:95:ALA:O	34:SR:97:GLY:N	4.87	0.45
36:1:160:G:O6	86:1:4191:OHX:N6	2.50	0.45
36:1:2657:A:C2	36:1:2694:A:C8	3.05	0.45
36:1:2948:C:H2'	36:1:2949:U:H6	1.81	0.45
86:1:4027:OHX:N4	86:1:4040:OHX:N1	2.65	0.45
36:1:600:G:H5'	36:1:601:U:OP2	2.17	0.45
36:1:846:A:H3'	36:1:847:A:H8	1.81	0.45
36:1:936:A:OP1	64:N8:28:HIS:ND1	2.39	0.45
1:2:1014:G:H2'	1:2:1015:U:O4'	2.16	0.45
1:2:1145:U:C4	1:2:1146:G:N7	2.86	0.45
1:2:1530:C:OP2	27:D5:95:HIS:ND1	2.27	0.45
1:2:1718:G:H2'	1:2:1719:A:O4'	2.17	0.45
1:2:180:A:H2'	1:2:181:A:O4'	2.17	0.45
1:2:289:U:H2'	1:2:290:G:O4'	2.17	0.45
1:2:435:C:H2'	1:2:436:A:C8	2.52	0.45
1:2:902:G:H8	1:2:902:G:O5'	2.00	0.45
36:5:1018:G:H2'	36:5:1019:G:O4'	2.17	0.45
36:5:1171:G:O6	86:5:3995:OHX:N1	2.50	0.45
36:5:1340:G:C5	36:5:1341:U:C5	3.05	0.45
36:5:2971:A:H5''	36:5:2972:G:C5'	2.47	0.45
36:5:374:A:N3	36:5:376:G:H5''	2.32	0.45
36:5:847:A:H2'	36:5:848:A:H8	1.75	0.45
36:5:949:C:O2'	36:5:971:G:OP1	2.26	0.45
1:6:1192:C:H3'	1:6:1193:A:H2'	1.99	0.45
1:6:1512:G:C5	1:6:1513:G:N7	2.85	0.45
27:D5:96:SER:HB3	1:6:1530:C:OP2	371.43	0.45
1:6:907:A:H1'	1:6:997:G:O2'	2.17	0.45
86:7:220:OHX:N3	86:7:228:OHX:N6	2.64	0.45
36:5:22:G:N2	38:8:35:C:O2	2.46	0.45
13:C1:86:ILE:HG13	13:C1:107:VAL:O	4.11	0.45
14:C2:30:VAL:HG11	14:C2:132:GLU:OE1	2.70	0.45
16:C4:117:ASP:OD1	16:C4:119:THR:HG23	2.17	0.45
1:2:927:C:H1'	16:C4:125:SER:CB	2.47	0.45
17:C5:53:PRO:HG2	17:C5:57:MET:HG3	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:C8:134:ARG:HG2	1:6:1559:A:C4	362.76	0.45
22:D0:23:ARG:HD2	22:D0:90:TYR:CD1	2.52	0.45
29:D7:50:ALA:O	29:D7:51:GLN:HB2	2.17	0.45
30:D8:36:THR:O	30:D8:37:SER:OG	2.27	0.45
33:E1:91:ILE:HD11	33:E1:92:LYS:HZ3	11.10	0.45
41:L4:215:ILE:HA	41:L4:218:ALA:HB3	3.02	0.45
41:L4:286:VAL:HG11	54:M8:31:LYS:HE3	4.78	0.45
41:L4:325:LEU:HD23	41:L4:325:LEU:HA	2.13	0.45
41:L4:49:ALA:HA	41:L4:109:TRP:CZ2	2.81	0.45
42:L5:268:GLU:HG3	42:L5:269:SER:N	3.59	0.45
44:L7:127:LEU:HA	44:L7:127:LEU:HD23	3.30	0.45
46:L9:88:TYR:CE2	46:L9:184:LYS:HG2	2.52	0.45
49:M3:168:ARG:HA	49:M3:171:ARG:HB2	1.99	0.45
50:M4:39:ILE:O	56:N0:95:ARG:HD3	2.17	0.45
51:M5:37:HIS:NE2	51:M5:63:ARG:NH1	2.65	0.45
52:M6:34:VAL:HB	52:M6:103:LYS:HB2	2.24	0.45
52:M6:117:ARG:HG2	52:M6:117:ARG:H	1.55	0.45
52:M6:138:LEU:O	52:M6:140:LYS:N	2.50	0.45
52:M6:172:ARG:HD2	36:5:3191:G:P	307.23	0.45
53:M7:69:ARG:HG2	53:M7:79:THR:OG1	4.32	0.45
54:M8:62:VAL:HG13	54:M8:66:ARG:HD3	1.98	0.45
54:M8:73:GLN:HB3	54:M8:76:ALA:HB3	1.98	0.45
55:M9:171:ASP:O	55:M9:175:GLN:NE2	2.49	0.45
55:M9:97:ARG:HH21	36:5:1779:C:H5'	205.68	0.45
56:N0:109:ASP:O	56:N0:112:ALA:N	3.03	0.45
56:N0:124:LEU:HA	56:N0:124:LEU:HD23	1.69	0.45
56:N0:148:LEU:HD22	56:N0:149:LYS:N	5.34	0.45
56:N0:166:LYS:HB3	56:N0:167:ARG:H	1.54	0.45
57:N1:40:VAL:HB	57:N1:96:ILE:HG23	2.15	0.45
61:N5:55:ASN:ND2	38:8:133:G:H5''	80.31	0.45
62:N6:27:ARG:NH1	62:N6:75:ARG:O	2.49	0.45
63:N7:42:LEU:HD23	63:N7:101:PHE:HE1	1.82	0.45
64:N8:118:ILE:H	64:N8:118:ILE:HD13	1.82	0.45
64:N8:34:MET:HA	64:N8:34:MET:HE2	2.97	0.45
68:O2:22:SER:HA	68:O2:28:VAL:CG1	2.46	0.45
70:O4:74:ARG:NH2	70:O4:82:ALA:HB2	3.12	0.45
63:N7:136:PHE:CD1	70:O4:89:ILE:HG12	3.39	0.45
3:S1:81:PHE:HE1	3:S1:109:LYS:HE2	1.81	0.45
5:S3:167:PHE:HA	5:S3:190:ARG:NE	2.32	0.45
5:S3:209:ILE:O	19:C7:20:TYR:OH	2.90	0.45
5:S3:20:GLU:HG3	12:C0:61:TRP:CG	2.51	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:S5:205:SER:O	7:S5:207:THR:N	2.50	0.45
6:S4:148:ARG:NH1	8:S6:201:GLN:OE1	2.46	0.45
9:S7:91:ILE:HA	9:S7:91:ILE:HD12	1.74	0.45
10:S8:34:ALA:HB2	10:S8:56:ARG:HG3	1.99	0.45
11:S9:54:ARG:HH11	11:S9:55:ALA:HB2	1.81	0.45
34:SR:127:ARG:C	34:SR:129:LYS:H	2.74	0.45
36:1:1699:A:H2'	36:1:1700:G:H8	1.81	0.44
36:1:2305:G:N2	36:1:2305:G:OP2	2.43	0.44
36:1:2337:C:H2'	36:1:2338:C:H6	1.81	0.44
36:1:238:A:C2'	36:1:239:G:H5'	2.47	0.44
36:1:409:A:H61	38:4:15:G:H1'	1.82	0.44
36:1:558:U:O5'	36:1:558:U:H6	2.00	0.44
36:1:613:G:C6	36:1:614:C:C4	3.05	0.44
36:1:744:A:H2'	36:1:745:C:O4'	2.17	0.44
1:2:1378:U:H2'	1:2:1379:C:O4'	2.18	0.44
1:2:1459:C:H42	20:C8:139:LYS:HE2	1.82	0.44
1:2:301:A:C6	1:2:302:U:C4	3.05	0.44
1:2:627:C:H2'	1:2:628:G:O4'	2.18	0.44
1:2:863:A:H2'	1:2:865:A:H8	1.81	0.44
1:2:869:A:H2'	1:2:870:C:O4'	2.16	0.44
37:3:109:G:C6	37:3:110:G:C5	3.05	0.44
36:5:1114:U:O2'	36:5:1115:G:H5'	2.17	0.44
36:5:1210:U:H2'	36:5:1211:U:C6	2.51	0.44
36:5:1675:G:O6	86:5:3963:OHX:N4	2.51	0.44
36:5:2279:A:C2	36:5:2288:G:C6	3.06	0.44
36:5:786:A:H4'	36:5:787:G:OP1	2.16	0.44
1:6:1239:U:O4	86:6:2096:OHX:N2	2.50	0.44
21:C9:91:TYR:OH	1:6:1469:A:OP1	365.11	0.44
1:6:489:C:C4	1:6:498:G:N2	2.85	0.44
37:7:47:C:O2'	37:7:48:U:H5'	2.17	0.44
13:C1:33:ARG:HD2	13:C1:49:ILE:O	3.47	0.44
1:2:867:G:H5'	15:C3:4:MET:HE3	1.98	0.44
16:C4:120:PRO:HB2	1:6:887:A:H5''	284.03	0.44
16:C4:123:SER:O	16:C4:124:ASP:HB2	4.89	0.44
16:C4:64:ALA:HB3	16:C4:104:ALA:HB3	1.99	0.44
18:C6:47:LYS:HZ3	18:C6:114:ARG:HE	1.65	0.44
19:C7:27:ASP:HB3	19:C7:30:THR:HG22	1.98	0.44
20:C8:49:LYS:NZ	20:C8:80:LYS:HB2	3.19	0.44
21:C9:114:VAL:HG23	21:C9:123:ARG:O	2.17	0.44
21:C9:40:SER:CB	21:C9:96:ALA:HA	2.47	0.44
24:D2:76:SER:HA	24:D2:77:PRO:C	5.15	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:D3:61:SER:HB3	25:D3:69:ARG:HD2	1.99	0.44
26:D4:123:LYS:HE2	26:D4:123:LYS:HB3	5.06	0.44
28:D6:5:ARG:HG2	1:6:1796:C:C6	343.19	0.44
31:D9:22:ARG:HD2	31:D9:38:ILE:HG12	1.99	0.44
33:E1:144:CYS:SG	33:E1:147:VAL:HG12	5.21	0.44
40:L3:205:VAL:O	40:L3:207:SER:N	2.67	0.44
40:L3:256:HIS:HA	40:L3:257:PRO:C	2.45	0.44
40:L3:221:THR:O	40:L3:272:TYR:HA	2.47	0.44
40:L3:305:ILE:HG12	40:L3:321:PHE:CZ	2.52	0.44
41:L4:128:ALA:O	41:L4:131:VAL:HG23	3.19	0.44
36:1:933:A:C2	41:L4:98:ARG:NH1	2.85	0.44
42:L5:234:ASP:OD2	42:L5:234:ASP:N	2.50	0.44
42:L5:29:ASP:O	42:L5:31:TYR:N	3.08	0.44
42:L5:34:LYS:HD2	57:N1:30:TYR:CZ	2.59	0.44
43:L6:169:ASP:OD1	43:L6:169:ASP:N	3.14	0.44
43:L6:57:HIS:NE2	43:L6:61:ASN:HA	2.31	0.44
45:L8:143:ILE:HG23	45:L8:175:VAL:HG21	1.99	0.44
45:L8:49:TYR:HD2	36:5:2587:U:H4'	178.11	0.44
45:L8:61:GLN:HB3	51:M5:28:TRP:CH2	2.51	0.44
47:M0:150:GLU:OE2	47:M0:153:ARG:NH2	3.88	0.44
49:M3:116:LEU:HA	49:M3:116:LEU:HD23	2.14	0.44
49:M3:165:SER:HB3	49:M3:168:ARG:HB3	1.99	0.44
49:M3:178:LYS:HD3	49:M3:179:PHE:CE2	3.02	0.44
36:1:328:U:P	49:M3:23:LYS:HE3	2.58	0.44
51:M5:197:LEU:HD12	51:M5:197:LEU:HA	2.03	0.44
51:M5:49:ARG:HB2	51:M5:49:ARG:NH1	2.32	0.44
53:M7:176:ILE:O	53:M7:179:GLN:HB2	2.17	0.44
59:N3:53:SER:O	59:N3:54:LEU:C	2.66	0.44
62:N6:112:ASP:HB2	62:N6:115:ARG:H	1.81	0.44
63:N7:3:LYS:HD3	66:O0:35:ARG:O	2.91	0.44
64:N8:83:PRO:HG2	64:N8:86:LYS:HB2	4.72	0.44
68:O2:20:HIS:ND1	68:O2:42:VAL:HG21	2.33	0.44
69:O3:87:ASN:OD1	69:O3:87:ASN:N	2.50	0.44
70:O4:59:PRO:HA	70:O4:62:TYR:HD2	1.82	0.44
49:M3:128:ARG:NH1	71:O5:109:ILE:O	5.08	0.44
74:O8:14:LEU:O	74:O8:17:ARG:HB2	2.89	0.44
2:S0:179:ARG:HH11	2:S0:183:ARG:NH1	2.14	0.44
3:S1:62:LYS:HB2	3:S1:62:LYS:HE2	1.87	0.44
4:S2:57:PHE:CZ	4:S2:138:PRO:HD3	2.59	0.44
5:S3:45:LYS:HB2	5:S3:45:LYS:HE2	1.83	0.44
7:S5:119:ASP:O	7:S5:122:ASN:ND2	2.40	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:S5:62:VAL:HG13	7:S5:89:ILE:HG21	2.57	0.44
8:S6:20:ASP:OD2	8:S6:22:HIS:N	5.06	0.44
10:S8:40:ALA:O	10:S8:59:ARG:HB3	2.17	0.44
11:S9:69:ARG:O	11:S9:73:GLY:HA3	2.17	0.44
34:SR:170:ILE:HG21	34:SR:211:ILE:HD13	2.87	0.44
34:SR:176:LYS:HD3	34:SR:196:ASN:C	2.37	0.44
34:SR:201:THR:HB	34:SR:242:SER:N	2.26	0.44
36:1:1362:G:O2'	36:1:1363:A:H5'	2.17	0.44
36:1:1840:U:O2	36:1:1850:A:H5'	2.17	0.44
36:1:1845:G:N2	36:1:1851:G:C4	2.85	0.44
36:1:2366:C:N4	36:1:2381:G:H1	2.14	0.44
1:2:1053:G:C6	1:2:1054:U:C4	3.05	0.44
1:2:1086:A:H5''	1:2:1087:A:OP2	2.18	0.44
1:2:121:U:C4	1:2:122:U:C4	3.05	0.44
1:2:145:A:O2'	1:2:146:U:O5'	2.23	0.44
1:2:1546:G:OP1	20:C8:123:ARG:HD2	2.17	0.44
1:2:1657:U:H4'	1:2:1658:G:O5'	2.17	0.44
1:2:33:U:O4	86:2:2055:OHX:N3	2.50	0.44
1:2:129:U:O4	1:2:264:G:H2'	2.17	0.44
1:2:732:G:N3	1:2:732:G:H2'	2.31	0.44
1:2:706:A:C6	1:2:734:A:N6	2.85	0.44
1:2:88:U:H4'	1:2:171:A:O4'	2.17	0.44
1:2:932:U:H4'	1:2:933:A:O4'	2.18	0.44
38:4:111:A:OP2	73:O7:32:LYS:HE3	2.18	0.44
36:1:2585:G:C2	38:4:151:C:H5	2.35	0.44
38:4:42:G:C4	38:4:43:A:C8	3.05	0.44
38:4:68:G:OP1	73:O7:86:ALA:N	2.49	0.44
57:N1:105:PHE:CE2	36:5:1062:A:H4'	243.83	0.44
36:5:1522:U:H4'	36:5:1523:U:OP2	2.17	0.44
36:5:1763:U:H3'	36:5:1764:U:C6	2.53	0.44
36:5:1784:G:H2'	36:5:1785:U:O4'	2.17	0.44
36:5:2137:U:C6	36:5:2141:U:C5	3.06	0.44
36:5:198:A:C6	36:5:219:A:C6	3.05	0.44
36:5:2407:C:O2	36:5:2818:U:N3	2.41	0.44
36:5:3078:U:O2'	86:5:4190:OHX:N1	2.50	0.44
40:L3:129:ALA:O	36:5:3150:A:H5'	211.47	0.44
36:5:3160:U:C2	36:5:3291:G:C2	3.05	0.44
36:5:3255:U:H2'	36:5:3256:G:H8	1.79	0.44
40:L3:173:GLN:HB2	36:5:3313:U:O3'	203.16	0.44
36:5:1752:A:OP2	86:5:4074:OHX:N3	2.51	0.44
36:5:1670:C:OP1	86:5:4229:OHX:N3	2.51	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:720:A:O2'	36:5:784:A:OP2	2.24	0.44
1:6:1192:C:OP2	1:6:1193:A:O2'	2.30	0.44
21:C9:75:LYS:HE2	1:6:1520:U:OP2	418.89	0.44
1:6:1570:A:C6	1:6:1571:C:C2	3.05	0.44
32:E0:2:ALA:HA	1:6:1647:U:O2	329.41	0.44
1:6:356:G:OP2	86:6:2074:OHX:N5	2.51	0.44
26:D4:62:THR:HG23	1:6:531:C:O2	421.87	0.44
1:6:640:U:H2'	1:6:641:G:C8	2.52	0.44
8:S6:171:LYS:NZ	1:6:67:A:OP1	349.04	0.44
1:6:861:U:H5'	1:6:862:A:OP2	2.17	0.44
15:C3:14:SER:OG	1:6:958:U:H2'	340.58	0.44
1:6:624:G:N2	1:6:976:G:H1'	2.32	0.44
12:C0:14:TYR:CZ	12:C0:18:GLU:HG3	2.52	0.44
14:C2:89:ILE:O	14:C2:89:ILE:HD13	4.51	0.44
18:C6:116:LEU:HA	18:C6:116:LEU:HD23	3.90	0.44
18:C6:15:SER:OG	18:C6:72:GLY:N	2.94	0.44
18:C6:82:ARG:NH1	18:C6:114:ARG:HB2	5.38	0.44
21:C9:28:LEU:HB3	21:C9:29:GLU:H	3.72	0.44
23:D1:14:PRO:HB2	23:D1:23:ILE:HG23	2.00	0.44
25:D3:17:VAL:HG23	25:D3:20:ARG:NH2	4.34	0.44
26:D4:103:ALA:O	26:D4:108:ARG:HG3	4.78	0.44
28:D6:17:HIS:ND1	28:D6:18:VAL:O	2.50	0.44
28:D6:10:ARG:HD3	28:D6:34:LYS:CA	2.89	0.44
40:L3:85:VAL:O	40:L3:162:VAL:HA	2.36	0.44
40:L3:238:LEU:HB2	40:L3:246:LEU:O	2.17	0.44
40:L3:286:GLY:HA3	40:L3:321:PHE:CE2	2.70	0.44
40:L3:81:THR:CG2	40:L3:81:THR:O	3.33	0.44
41:L4:349:THR:HG21	44:L7:64:GLN:NE2	2.33	0.44
42:L5:233:ALA:O	42:L5:236:LEU:N	2.39	0.44
42:L5:37:VAL:HG12	57:N1:31:LEU:HD21	1.98	0.44
46:L9:47:LYS:HZ3	50:M4:6:ILE:H	1.63	0.44
51:M5:104:GLU:O	51:M5:108:ARG:HG3	2.16	0.44
51:M5:160:GLU:OE1	51:M5:160:GLU:N	2.82	0.44
52:M6:85:ARG:HD3	52:M6:90:HIS:ND1	2.84	0.44
54:M8:64:VAL:HG12	54:M8:90:ASP:H	1.83	0.44
57:N1:17:ARG:O	57:N1:18:ASP:HB2	2.18	0.44
58:N2:22:PRO:HG3	58:N2:105:LEU:HB3	1.99	0.44
61:N5:105:VAL:HG13	61:N5:130:TYR:CD1	3.75	0.44
62:N6:36:SER:OG	62:N6:38:GLU:HB2	3.59	0.44
69:O3:6:ARG:HD2	69:O3:8:TYR:O	2.33	0.44
70:O4:20:ILE:HD11	70:O4:34:HIS:CE1	2.52	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
72:O6:68:ARG:O	72:O6:72:VAL:HG23	2.16	0.44
74:O8:65:LEU:O	74:O8:69:LEU:HD22	2.16	0.44
5:S3:158:ILE:HD13	5:S3:158:ILE:H	1.87	0.44
5:S3:17:PHE:CE1	5:S3:79:TYR:HE2	5.05	0.44
7:S5:30:PRO:HD2	7:S5:33:VAL:HG11	1.98	0.44
7:S5:73:THR:HG23	18:C6:114:ARG:HD2	1.99	0.44
9:S7:49:ILE:O	9:S7:57:ALA:N	2.50	0.44
11:S9:171:ARG:NH1	11:S9:174:ARG:HB3	3.75	0.44
11:S9:17:ARG:HD3	11:S9:20:GLU:OE2	2.17	0.44
34:SR:43:ILE:HD13	34:SR:60:SER:HA	1.98	0.44
34:SR:40:LYS:HB3	34:SR:64:HIS:O	3.03	0.44
36:1:1302:A:N1	36:1:2832:C:O2'	2.44	0.44
36:1:1307:G:H4'	36:1:1308:A:O5'	2.18	0.44
36:1:1488:G:O2'	70:O4:10:ARG:O	2.34	0.44
36:1:1619:A:C2	36:1:1826:C:C2	3.06	0.44
36:1:1715:A:H4'	36:1:1716:U:OP1	2.17	0.44
36:1:2107:A:C2	36:1:2108:C:C2	3.06	0.44
36:1:250:U:H5''	36:1:251:G:H5''	1.99	0.44
36:1:2603:G:H2'	36:1:2604:U:O4'	2.18	0.44
36:1:268:A:C4	51:M5:12:ARG:HG2	2.52	0.44
36:1:3046:A:C5	36:1:3047:U:C5	3.05	0.44
36:1:3082:C:H2'	36:1:3083:G:C8	2.52	0.44
36:1:3084:C:OP2	86:1:3879:OHX:N5	2.50	0.44
36:1:3131:U:H2'	36:1:3132:C:C6	2.53	0.44
36:1:3310:A:C2'	36:1:3311:C:H5'	2.48	0.44
86:1:4023:OHX:N4	86:1:4143:OHX:N3	2.65	0.44
36:1:748:U:H2'	36:1:749:C:C6	2.51	0.44
1:2:1734:U:H2'	1:2:1735:U:H6	1.82	0.44
1:2:198:A:H2'	1:2:198:A:N3	2.31	0.44
1:2:314:C:C2	1:2:355:G:C2	3.05	0.44
37:3:90:U:C4	37:3:91:G:C5	3.05	0.44
38:4:98:U:H2'	38:4:99:C:H5'	1.99	0.44
36:5:1341:U:H2'	36:5:1342:C:H6	1.82	0.44
36:5:951:A:P	36:5:1367:G:H22	2.40	0.44
36:5:1804:A:H2'	36:5:1805:C:C6	2.53	0.44
36:5:2106:A:H2'	36:5:2107:A:C8	2.53	0.44
86:5:3966:OHX:N1	86:5:4238:OHX:N1	2.65	0.44
69:O3:86:ARG:HH12	36:5:498:A:H5'	215.23	0.44
36:5:692:A:C4	36:5:693:A:C8	3.06	0.44
36:5:920:A:OP1	36:5:922:U:C5	2.71	0.44
1:6:1187:U:O5'	1:6:1187:U:H6	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:1255:G:H4'	1:6:1256:A:OP1	2.16	0.44
1:6:157:A:O5'	1:6:157:A:H8	2.00	0.44
1:6:293:U:OP2	86:6:2136:OHX:N2	2.49	0.44
1:6:451:A:OP2	86:6:2061:OHX:N3	2.50	0.44
1:6:526:A:N6	1:6:527:A:C6	2.85	0.44
1:6:809:A:N6	1:6:810:G:O6	2.50	0.44
86:8:216:OHX:N6	86:8:223:OHX:N3	2.65	0.44
16:C4:85:ALA:N	16:C4:119:THR:HG22	2.21	0.44
18:C6:114:ARG:O	18:C6:115:THR:OG1	2.30	0.44
18:C6:32:ASN:N	18:C6:67:VAL:O	2.67	0.44
26:D4:15:ASN:ND2	26:D4:18:LEU:HB2	2.32	0.44
28:D6:36:ILE:HD12	28:D6:78:ALA:CB	2.47	0.44
29:D7:23:THR:HG21	29:D7:29:ARG:HH21	1.83	0.44
30:D8:12:VAL:HG11	30:D8:49:ARG:O	2.17	0.44
30:D8:19:THR:HG22	30:D8:20:GLY:H	1.83	0.44
40:L3:339:ARG:NH2	40:L3:342:LEU:HD11	3.13	0.44
41:L4:13:GLY:O	41:L4:14:GLU:HG2	4.90	0.44
42:L5:131:LEU:HD11	42:L5:175:HIS:H	3.53	0.44
45:L8:244:ALA:HA	45:L8:247:ASP:HB2	2.36	0.44
46:L9:68:LEU:O	46:L9:71:VAL:N	2.99	0.44
48:M1:15:GLU:HG2	48:M1:16:LYS:HE2	1.99	0.44
36:1:2356:A:OP1	53:M7:138:LYS:NZ	2.49	0.44
53:M7:94:LEU:CD2	53:M7:146:ILE:HB	3.52	0.44
54:M8:23:ASN:OD1	54:M8:25:TYR:N	2.50	0.44
55:M9:43:LYS:O	55:M9:47:ASN:HB2	5.51	0.44
67:O1:44:MET:HB3	67:O1:77:ARG:NH1	3.57	0.44
68:O2:75:LEU:HD23	68:O2:75:LEU:HA	1.84	0.44
70:O4:44:CYS:HB3	70:O4:47:CYS:HG	1.83	0.44
2:S0:102:PHE:HZ	2:S0:107:PHE:CE1	2.35	0.44
1:2:1064:G:O2'	3:S1:204:ILE:O	2.34	0.44
5:S3:28:GLU:HG3	5:S3:29:LEU:HD23	3.44	0.44
6:S4:181:VAL:CG1	6:S4:225:VAL:HG13	3.39	0.44
6:S4:31:PRO:HG2	6:S4:38:LEU:HD13	2.00	0.44
6:S4:3:ARG:O	1:6:93:A:H1'	326.94	0.44
7:S5:96:SER:HB3	7:S5:176:THR:HG21	3.08	0.44
9:S7:133:THR:HG21	9:S7:159:VAL:HA	2.92	0.44
36:1:1149:G:H21	36:1:1199:C:N4	2.15	0.44
36:1:1831:U:P	61:N5:92:LYS:HD3	2.57	0.44
36:1:2112:U:HO2'	36:1:2113:A:P	2.40	0.44
36:1:2265:C:H2'	36:1:2266:U:O4'	2.17	0.44
36:1:3026:G:O2'	36:1:3028:G:N7	2.33	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:3131:U:O2'	36:1:3132:C:H5'	2.18	0.44
36:1:3344:A:H2	36:1:3361:G:N2	2.16	0.44
36:1:350:C:N3	36:1:367:A:H2'	2.33	0.44
36:1:372:A:H2'	36:1:373:A:H8	1.82	0.44
1:2:1295:G:C6	1:2:1303:U:C2	3.06	0.44
1:2:1354:G:H5'	1:2:1355:C:OP2	2.17	0.44
1:2:1460:A:C5	35:SM:76:VAL:HG22	2.52	0.44
1:2:1766:A:H5''	86:2:2091:OHX:N6	2.32	0.44
1:2:245:U:O4	86:2:2092:OHX:N5	2.51	0.44
1:2:330:G:H2'	1:2:331:A:O4'	2.17	0.44
1:2:705:U:H2'	1:2:706:A:N7	2.33	0.44
36:1:59:G:H2'	38:4:33:A:O2'	2.17	0.44
36:5:1249:G:H2'	36:5:1250:G:C8	2.52	0.44
36:5:1650:G:H1	36:5:1805:C:H42	1.65	0.44
36:5:201:A:H2'	36:5:202:G:C8	2.52	0.44
36:5:24:G:H2'	36:5:25:U:O4'	2.17	0.44
36:5:2609:A:H2'	36:5:2610:G:H8	1.82	0.44
36:5:2973:G:N7	86:5:4112:OHX:N1	2.66	0.44
40:L3:98:GLY:HA3	36:5:3005:A:H5'	248.39	0.44
36:5:2993:G:H2'	36:5:3142:A:H61	1.81	0.44
36:5:314:U:O4	86:5:4186:OHX:N5	2.49	0.44
36:5:607:A:H4'	36:5:608:A:OP2	2.17	0.44
36:5:855:U:C4	36:5:856:G:C6	3.06	0.44
1:6:1117:U:H2'	1:6:1118:G:H8	1.82	0.44
20:C8:142:GLY:HA2	1:6:1173:C:OP1	342.70	0.44
1:6:390:G:H2'	1:6:391:A:O4'	2.18	0.44
1:6:867:G:O6	86:6:2057:OHX:N1	2.50	0.44
51:M5:38:ARG:NH2	38:8:143:U:OP1	108.62	0.44
38:8:65:A:H2'	38:8:66:A:O4'	2.17	0.44
12:C0:12:HIS:NE2	12:C0:49:LEU:HD21	2.32	0.44
12:C0:27:PHE:HD2	1:6:1217:A:C2	420.59	0.44
12:C0:49:LEU:HB3	12:C0:55:VAL:HG13	2.68	0.44
14:C2:140:PHE:HD2	14:C2:140:PHE:HA	1.88	0.44
15:C3:3:ARG:HB2	15:C3:8:GLY:O	2.18	0.44
7:S5:34:GLN:HA	18:C6:57:LEU:HD21	2.84	0.44
20:C8:134:ARG:HG2	1:6:1559:A:C5	362.81	0.44
21:C9:20:SER:O	21:C9:24:ARG:HG3	3.95	0.44
22:D0:50:LEU:HG	22:D0:94:GLU:O	2.17	0.44
23:D1:4:ASP:OD1	23:D1:5:LYS:HD2	3.34	0.44
25:D3:69:ARG:HD3	25:D3:117:ILE:HG12	1.98	0.44
28:D6:12:LYS:HB3	28:D6:13:LYS:H	4.48	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:D6:82:ARG:NH2	1:6:1153:G:OP2	332.01	0.44
29:D7:15:GLU:OE2	29:D7:24:LEU:N	2.44	0.44
30:D8:57:MET:O	30:D8:59:SER:N	4.90	0.44
39:L2:36:GLU:OE2	39:L2:163:ARG:NH1	2.97	0.44
36:1:2185:G:P	39:L2:201:GLY:H	2.40	0.44
39:L2:57:PRO:HG2	39:L2:78:ALA:HB3	2.09	0.44
40:L3:142:ALA:O	40:L3:145:GLU:N	2.61	0.44
41:L4:192:GLY:O	41:L4:194:TYR:N	3.22	0.44
41:L4:236:LEU:HA	41:L4:236:LEU:HD23	2.57	0.44
41:L4:35:VAL:HG11	41:L4:244:LEU:HD21	2.00	0.44
41:L4:24:ALA:HB2	41:L4:264:SER:HB3	1.99	0.44
41:L4:119:ARG:HG2	41:L4:274:TYR:CE2	2.52	0.44
41:L4:295:ILE:HD12	54:M8:132:PRO:HG3	2.00	0.44
41:L4:67:THR:HG23	41:L4:68:GLY:N	3.23	0.44
42:L5:99:TYR:HA	42:L5:161:GLY:O	2.68	0.44
42:L5:184:ASP:OD2	42:L5:187:THR:HG23	2.17	0.44
42:L5:261:THR:O	42:L5:263:GLU:N	3.58	0.44
44:L7:222:HIS:CE1	44:L7:224:ILE:HG13	2.56	0.44
44:L7:142:SER:HG	44:L7:241:LYS:HZ1	2.71	0.44
45:L8:111:LYS:HG2	45:L8:112:GLU:N	4.72	0.44
45:L8:156:ASP:HB2	45:L8:157:VAL:H	1.65	0.44
45:L8:84:ARG:HH22	45:L8:181:LYS:HZ3	1.65	0.44
49:M3:151:ALA:O	49:M3:153:ASP:N	4.18	0.44
51:M5:183:THR:HA	51:M5:187:ARG:H	3.26	0.44
53:M7:26:PHE:CD2	53:M7:121:GLN:HG2	2.53	0.44
57:N1:92:ARG:O	57:N1:94:GLU:N	2.51	0.44
58:N2:53:ALA:HB1	58:N2:68:THR:HG22	1.99	0.44
59:N3:27:ASP:CG	59:N3:28:ASN:H	3.24	0.44
59:N3:81:GLN:NE2	59:N3:85:TRP:CE3	3.55	0.44
62:N6:104:LEU:HA	62:N6:104:LEU:HD23	1.75	0.44
62:N6:88:GLU:HG3	62:N6:94:SER:OG	3.38	0.44
64:N8:11:HIS:O	64:N8:14:HIS:HB2	2.17	0.44
66:O0:57:GLU:O	66:O0:60:ALA:HB3	2.33	0.44
66:O0:42:ILE:CG1	66:O0:67:VAL:HG22	3.77	0.44
66:O0:73:GLY:O	66:O0:75:ASN:N	2.51	0.44
67:O1:24:SER:HB3	36:5:2345:A:H5''	178.70	0.44
68:O2:120:THR:O	68:O2:122:PRO:HD3	2.17	0.44
78:Q2:47:GLN:OE1	78:Q2:54:THR:OG1	2.39	0.44
2:S0:73:VAL:O	2:S0:95:ALA:HB1	3.57	0.44
3:S1:143:THR:HA	3:S1:207:LEU:HA	1.99	0.44
3:S1:231:LEU:HA	3:S1:231:LEU:HD23	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:S1:58:SER:O	3:S1:60:ALA:N	2.51	0.44
3:S1:97:LEU:HG	3:S1:232:HIS:CE1	2.52	0.44
4:S2:57:PHE:CE1	4:S2:138:PRO:HD3	2.52	0.44
5:S3:110:LEU:HD23	5:S3:110:LEU:HA	1.83	0.44
5:S3:167:PHE:O	5:S3:190:ARG:HD2	2.17	0.44
6:S4:156:VAL:O	6:S4:157:ASN:HB2	2.17	0.44
6:S4:246:LEU:HD23	6:S4:250:GLU:HB3	2.00	0.44
6:S4:34:GLY:HA3	6:S4:35:PRO:HD3	1.59	0.44
7:S5:177:ILE:HA	7:S5:180:ARG:HH11	1.82	0.44
7:S5:217:LEU:HA	7:S5:217:LEU:HD23	2.31	0.44
7:S5:35:GLN:HB3	7:S5:36:ALA:H	1.88	0.44
1:2:164:A:H1'	8:S6:13:GLN:NE2	2.32	0.44
1:2:66:U:C5'	8:S6:173:PRO:HA	2.46	0.44
10:S8:104:ILE:O	10:S8:164:ARG:HA	2.59	0.44
10:S8:106:ALA:HB2	10:S8:165:LEU:HG	1.98	0.44
11:S9:129:ILE:O	11:S9:142:ASN:HA	2.50	0.44
11:S9:31:ALA:HA	11:S9:36:LEU:HG	2.00	0.44
36:1:1148:G:C6	36:1:1149:G:N7	2.86	0.44
36:1:1340:G:OP2	68:O2:61:LYS:HE3	2.17	0.44
36:1:1554:U:H4'	36:1:1555:U:C5'	2.48	0.44
36:1:1613:A:P	74:O8:46:ARG:HH22	2.41	0.44
36:1:2117:A:C8	36:1:3064:U:O2	2.71	0.44
36:1:2383:C:C2'	36:1:2384:A:H5'	2.48	0.44
36:1:36:C:OP2	51:M5:83:LYS:HE2	2.17	0.44
36:1:58:G:OP1	51:M5:157:LYS:NZ	2.39	0.44
36:1:75:G:O3'	49:M3:70:ARG:NH1	2.46	0.44
1:2:1160:A:OP2	18:C6:142:TYR:OH	2.28	0.44
1:2:1433:G:H2'	1:2:1434:U:H6	1.83	0.44
1:2:1434:U:H5''	1:2:1435:G:OP1	2.18	0.44
1:2:1636:C:C2	1:2:1638:G:C5	3.05	0.44
1:2:563:U:O2'	1:2:564:G:H5'	2.17	0.44
1:2:722:G:H3'	1:2:723:G:H5'	1.99	0.44
1:2:887:A:H61	1:2:925:G:H1	1.65	0.44
36:5:1106:G:C5	36:5:1107:C:C5	3.05	0.44
36:5:150:A:C2'	36:5:151:A:H5'	2.47	0.44
36:5:1692:U:C4	36:5:1693:C:N4	2.86	0.44
36:5:2256:A:OP2	36:5:2256:A:H2'	2.18	0.44
36:5:2369:G:H2'	36:5:2370:G:C8	2.53	0.44
36:5:2378:C:C2	36:5:2379:U:C5	3.05	0.44
76:Q0:112:LYS:HZ3	36:5:3107:U:P	304.05	0.44
36:5:2730:G:OP2	86:5:3952:OHX:N4	2.51	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:330:G:OP2	86:5:4043:OHX:N1	2.50	0.44
36:5:601:U:H2'	36:5:602:A:O4'	2.16	0.44
1:6:1053:G:C2	1:6:1067:C:C2	3.05	0.44
1:6:1161:C:H2'	1:6:1162:C:C6	2.49	0.44
1:6:1350:U:H2'	1:6:1351:G:H8	1.83	0.44
1:6:1743:U:H2'	1:6:1744:A:H8	1.82	0.44
86:6:2104:OHX:N1	86:6:2190:OHX:N4	2.66	0.44
1:6:544:A:H8	1:6:544:A:O5'	2.01	0.44
1:6:643:G:H1	1:6:691:C:N4	2.15	0.44
1:6:961:U:H2'	1:6:962:C:H6	1.80	0.44
36:5:1618:G:H4'	38:8:129:C:H1'	1.98	0.44
38:8:132:G:C6	38:8:133:G:N7	2.86	0.44
12:C0:44:LYS:HA	12:C0:44:LYS:HD2	2.95	0.44
14:C2:95:LYS:HB2	14:C2:95:LYS:HE3	4.31	0.44
19:C7:58:MET:HA	19:C7:61:ILE:HB	3.32	0.44
21:C9:33:TYR:HE1	21:C9:100:ILE:HD13	2.27	0.44
22:D0:58:LEU:HD23	1:6:1516:A:C8	444.34	0.44
25:D3:27:ASN:O	25:D3:31:LYS:HG2	2.17	0.44
27:D5:41:ILE:HG23	27:D5:42:LEU:H	1.82	0.44
33:E1:98:VAL:HG13	33:E1:99:LYS:N	2.32	0.44
40:L3:255:TRP:CD1	40:L3:255:TRP:C	2.90	0.44
40:L3:59:ASP:OD1	40:L3:357:LYS:NZ	3.35	0.44
41:L4:144:LYS:H	41:L4:144:LYS:CE	4.99	0.44
42:L5:113:LEU:HA	42:L5:113:LEU:HD12	1.77	0.44
42:L5:211:LEU:HD22	42:L5:211:LEU:HA	1.75	0.44
43:L6:30:LEU:HA	43:L6:30:LEU:HD22	1.74	0.44
44:L7:102:VAL:HG12	44:L7:130:ILE:HD12	3.76	0.44
41:L4:327:LEU:CD1	44:L7:165:ASP:HA	3.14	0.44
44:L7:173:LEU:HB3	44:L7:178:ILE:HB	2.15	0.44
45:L8:33:ASN:O	36:5:2549:G:C6	208.93	0.44
46:L9:13:PRO:HG2	46:L9:16:VAL:HG11	2.91	0.44
46:L9:43:VAL:CG2	46:L9:55:VAL:HG12	5.10	0.44
47:M0:55:ASN:HA	47:M0:131:ILE:HG23	2.00	0.44
47:M0:138:VAL:HG11	47:M0:148:VAL:HG13	3.23	0.44
47:M0:21:ARG:NH2	47:M0:22:TYR:OH	2.50	0.44
49:M3:84:GLY:O	49:M3:85:LEU:HB3	2.66	0.44
49:M3:97:VAL:HG12	49:M3:98:ASP:H	1.82	0.44
50:M4:21:VAL:HB	50:M4:63:VAL:HG13	2.00	0.44
51:M5:50:ARG:HH11	36:5:267:G:H4'	111.33	0.44
53:M7:25:SER:CB	53:M7:28:ASN:HB2	2.47	0.44
54:M8:18:ALA:HB1	54:M8:19:PRO:HD2	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:M9:175:GLN:O	55:M9:179:GLU:N	2.50	0.44
55:M9:43:LYS:HG3	36:5:1764:U:H5''	94.19	0.44
56:N0:42:TRP:HA	56:N0:42:TRP:CE3	2.53	0.44
57:N1:128:LEU:H	57:N1:128:LEU:HD12	1.93	0.44
63:N7:47:GLU:OE2	63:N7:69:LYS:HE2	3.48	0.44
63:N7:15:ARG:HB2	63:N7:79:HIS:CD2	2.53	0.44
63:N7:87:LEU:HB2	63:N7:127:ASN:ND2	2.32	0.44
68:O2:11:LYS:O	68:O2:13:HIS:N	3.01	0.44
70:O4:3:GLN:HG2	70:O4:4:ARG:N	2.35	0.44
77:Q1:2:ARG:HD2	77:Q1:4:LYS:HB3	3.63	0.44
3:S1:140:ILE:HG22	3:S1:211:HIS:HB2	1.99	0.44
6:S4:193:GLY:O	6:S4:210:ILE:HG23	2.17	0.44
6:S4:173:ILE:HD11	6:S4:235:TYR:CE1	2.52	0.44
6:S4:26:CYS:HB2	6:S4:27:TYR:CE2	2.52	0.44
8:S6:136:LYS:HG3	8:S6:173:PRO:HB3	2.00	0.44
8:S6:13:GLN:OE1	1:6:151:G:N2	311.88	0.44
8:S6:87:ARG:NH1	1:6:159:U:O2'	321.92	0.44
9:S7:39:ARG:HG3	9:S7:40:PRO:HD3	1.97	0.44
10:S8:70:GLU:OE2	13:C1:24:LYS:NZ	2.35	0.44
11:S9:105:LEU:HA	11:S9:105:LEU:HD12	1.88	0.44
11:S9:10:LYS:HE3	11:S9:10:LYS:HB2	1.60	0.44
35:SM:64:LYS:C	35:SM:66:ALA:H	2.47	0.44
34:SR:122:ILE:HB	34:SR:134:TRP:HB2	1.99	0.44
36:1:1826:C:H2'	36:1:1827:C:H6	1.82	0.44
36:1:1842:A:H4'	36:1:1843:C:OP2	2.18	0.44
36:1:1864:A:OP1	55:M9:88:ARG:NH1	2.51	0.44
36:1:282:G:C8	36:1:282:G:H3'	2.53	0.44
36:1:3049:A:C6	36:1:3050:U:C2	3.05	0.44
36:1:597:G:C2	36:1:598:A:C8	3.06	0.44
1:2:1064:G:H2'	1:2:1065:A:C8	2.52	0.44
1:2:1156:C:C2'	1:2:1157:A:H5'	2.47	0.44
1:2:1395:G:N2	1:2:1404:C:O2	2.51	0.44
1:2:552:G:C6	1:2:553:G:C6	3.05	0.44
1:2:804:A:N3	24:D2:105:THR:HG22	2.33	0.44
1:2:970:A:H5''	1:2:970:A:H8	1.83	0.44
38:4:118:C:C2	38:4:136:G:C2	3.05	0.44
38:4:124:G:N2	38:4:130:C:C2	2.86	0.44
36:5:1277:C:H2'	36:5:1278:A:C8	2.53	0.44
36:5:1389:G:N2	36:5:1390:A:N1	2.65	0.44
68:O2:59:SER:OG	36:5:1405:U:OP2	184.38	0.44
36:5:1731:A:C5	36:5:1732:U:C5	3.06	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:2299:A:OP2	86:5:3953:OHX:N1	2.49	0.44
36:5:2440:G:HO2'	36:5:2441:A:P	2.40	0.44
36:5:2689:A:C4'	36:5:2690:G:H5'	2.44	0.44
36:5:2697:A:C2	36:5:2698:G:C5	3.05	0.44
36:5:2779:A:H8	36:5:2779:A:H5'	1.82	0.44
36:5:3195:U:H1'	36:5:3196:U:OP1	2.18	0.44
36:5:3385:U:H2'	36:5:3386:G:H8	1.83	0.44
51:M5:204:LYS:HE3	36:5:683:U:OP2	105.52	0.44
36:5:701:G:H2'	36:5:702:C:C6	2.53	0.44
36:5:817:A:H2'	36:5:920:A:C2	2.53	0.44
36:5:82:C:H2'	36:5:83:U:O4'	2.17	0.44
1:6:1138:A:H2'	1:6:1139:A:H8	1.83	0.44
1:6:1163:A:N6	1:6:1164:G:C6	2.85	0.44
1:6:1545:A:O2'	1:6:1546:G:H5'	2.18	0.44
1:6:25:C:O2	86:6:2107:OHX:N5	2.50	0.44
1:6:398:G:O5'	1:6:398:G:H8	2.01	0.44
1:6:565:C:C2	86:6:2159:OHX:N4	2.86	0.44
1:6:586:G:C6	1:6:587:C:C4	3.06	0.44
62:N6:113:LYS:HE2	38:8:85:G:N7	15.16	0.44
10:S8:70:GLU:CD	13:C1:24:LYS:HZ1	2.17	0.44
14:C2:43:ARG:HH12	14:C2:102:GLY:HA3	1.82	0.44
15:C3:78:ASN:C	15:C3:80:LEU:H	2.21	0.44
16:C4:133:ARG:HB2	16:C4:136:ARG:HH21	1.82	0.44
18:C6:102:LYS:O	18:C6:106:LYS:N	3.30	0.44
19:C7:69:ILE:HG12	19:C7:69:ILE:H	1.52	0.44
19:C7:74:GLN:HA	19:C7:77:GLU:HB2	1.98	0.44
22:D0:102:ARG:HG3	22:D0:103:ILE:N	4.51	0.44
22:D0:28:SER:HB3	22:D0:34:LEU:HD23	2.00	0.44
22:D0:38:SER:O	22:D0:42:VAL:HG23	2.18	0.44
25:D3:74:VAL:N	25:D3:83:VAL:O	2.49	0.44
27:D5:46:LYS:O	27:D5:50:ILE:HG13	4.71	0.44
33:E1:135:HIS:ND1	33:E1:138:ARG:HD2	2.33	0.44
39:L2:220:GLY:O	39:L2:221:LYS:HG3	2.58	0.44
39:L2:39:GLY:O	39:L2:91:GLY:HA3	2.18	0.44
43:L6:6:ALA:HA	68:O2:74:PHE:CE1	2.52	0.44
44:L7:105:LEU:HA	44:L7:105:LEU:HD23	1.66	0.44
44:L7:127:LEU:C	44:L7:129:LEU:N	2.69	0.44
44:L7:77:VAL:HG13	57:N1:139:ARG:O	3.01	0.44
47:M0:185:ARG:HA	47:M0:190:VAL:HB	1.99	0.44
47:M0:57:LEU:HA	47:M0:57:LEU:HD22	1.74	0.44
37:3:40:C:O2'	48:M1:72:ARG:HD2	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:M3:109:PHE:O	49:M3:112:ASN:HB2	2.18	0.44
49:M3:124:ILE:HD11	49:M3:126:PHE:CZ	3.13	0.44
51:M5:122:ASN:OD1	51:M5:123:GLN:N	2.50	0.44
51:M5:68:ARG:HA	51:M5:98:LEU:HD21	2.91	0.44
52:M6:25:LYS:HG3	36:5:1175:C:H5'	254.31	0.44
54:M8:161:LYS:O	54:M8:162:ALA:HB3	2.18	0.44
54:M8:36:LEU:O	54:M8:40:THR:OG1	2.39	0.44
56:N0:8:GLN:OE1	56:N0:26:ARG:NE	2.41	0.44
57:N1:157:GLU:HB3	57:N1:159:PHE:CE1	2.75	0.44
63:N7:18:TYR:CE1	63:N7:71:PHE:CG	3.05	0.44
66:O0:29:SER:O	66:O0:33:SER:HB3	2.49	0.44
71:O5:102:GLU:HA	71:O5:102:GLU:OE2	2.16	0.44
71:O5:119:LYS:HA	71:O5:119:LYS:HD2	1.83	0.44
75:O9:41:ARG:HH22	36:5:1517:G:P	97.78	0.44
76:Q0:108:THR:HB	76:Q0:109:ASN:OD1	3.23	0.44
78:Q2:36:PHE:N	78:Q2:36:PHE:CD1	2.85	0.44
2:S0:167:LYS:HG2	2:S0:168:HIS:NE2	2.32	0.44
2:S0:179:ARG:HG3	2:S0:195:TRP:CE3	2.53	0.44
2:S0:70:PRO:O	2:S0:94:GLY:HA3	2.17	0.44
3:S1:119:THR:HB	3:S1:143:THR:CG2	2.66	0.44
3:S1:61:LEU:HD22	3:S1:61:LEU:H	1.82	0.44
4:S2:133:LYS:O	4:S2:136:VAL:HG23	2.20	0.44
10:S8:29:LEU:HD23	10:S8:30:GLY:CA	2.48	0.44
11:S9:64:GLU:HA	11:S9:69:ARG:HD3	2.31	0.44
34:SR:206:PRO:HG2	34:SR:247:PRO:HA	2.56	0.44
34:SR:269:TYR:H	34:SR:269:TYR:HD1	1.65	0.44
36:1:1048:A:N1	36:1:2646:C:O2'	2.40	0.44
36:1:1348:U:O2	36:1:1349:G:N2	2.51	0.44
36:1:1544:G:O6	86:1:4052:OHX:N4	2.50	0.44
36:1:1661:G:N2	36:1:1789:G:H1'	2.32	0.44
36:1:1668:G:H2'	36:1:1669:C:O4'	2.16	0.44
36:1:2289:U:H2'	36:1:2290:C:H6	1.83	0.44
1:2:1268:G:C2	1:2:1270:G:N7	2.86	0.44
1:2:1335:U:H1'	31:D9:56:ARG:HH11	1.82	0.44
1:2:1643:U:H5'	77:Q1:9:ARG:HH22	1.83	0.44
1:2:73:U:O2'	1:2:74:U:C2	2.69	0.44
37:3:24:A:H2'	37:3:25:G:H5'	2.00	0.44
36:5:3108:G:H2'	36:5:3109:G:O5'	2.17	0.44
36:5:3283:U:H2'	36:5:3284:G:C8	2.53	0.44
36:5:3320:A:C2	36:5:3321:C:C2	3.06	0.44
36:5:2734:A:OP1	86:5:4040:OHX:N6	2.50	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:1314:U:O2'	1:6:1315:U:OP2	2.35	0.44
1:6:1614:A:O2'	1:6:1615:C:H5'	2.18	0.44
1:6:1747:G:H2'	1:6:1748:G:O4'	2.17	0.44
1:6:187:G:N2	1:6:198:A:H62	2.16	0.44
1:6:47:A:H4'	1:6:48:G:C5'	2.48	0.44
1:6:891:A:H2'	1:6:892:A:C8	2.51	0.44
12:C0:2:LEU:HB3	12:C0:3:MET:H	4.27	0.44
15:C3:114:ARG:HA	15:C3:114:ARG:HD3	1.65	0.44
15:C3:12:SER:HB3	15:C3:13:SER:H	1.62	0.44
1:2:966:A:H5''	15:C3:4:MET:CE	2.48	0.44
17:C5:72:LYS:HG2	17:C5:72:LYS:H	3.25	0.44
17:C5:77:ARG:HA	17:C5:95:GLY:HA3	2.37	0.44
20:C8:134:ARG:HD2	1:6:1545:A:OP2	358.72	0.44
21:C9:105:LEU:HD13	21:C9:122:ARG:HD3	2.85	0.44
21:C9:136:ALA:O	21:C9:140:LEU:HD12	2.17	0.44
26:D4:57:VAL:HG22	26:D4:60:PHE:CE2	2.52	0.44
28:D6:4:LYS:HE2	28:D6:4:LYS:HB3	4.23	0.44
33:E1:137:ASP:HB2	33:E1:138:ARG:H	1.58	0.44
39:L2:29:LEU:H	39:L2:123:ARG:HB3	2.06	0.44
41:L4:219:LEU:O	41:L4:222:VAL:HG13	2.65	0.44
36:1:1080:A:OP2	42:L5:140:ARG:NH2	2.51	0.44
43:L6:165:LEU:HA	43:L6:165:LEU:HD23	1.75	0.44
44:L7:59:GLU:O	44:L7:63:ILE:HG13	2.18	0.44
45:L8:207:ASP:O	45:L8:209:ALA:N	3.69	0.44
46:L9:51:GLN:OE1	46:L9:51:GLN:N	3.21	0.44
46:L9:86:TYR:CZ	46:L9:151:VAL:HG13	2.52	0.44
48:M1:141:ARG:O	48:M1:145:LYS:HE2	2.18	0.44
50:M4:99:TRP:O	50:M4:99:TRP:HD1	2.13	0.44
51:M5:150:TRP:O	51:M5:153:ASP:HB2	2.42	0.44
55:M9:92:GLN:O	55:M9:96:ILE:HG13	2.25	0.44
55:M9:99:LEU:HD11	55:M9:103:ARG:NH1	2.33	0.44
57:N1:102:ARG:O	57:N1:106:LEU:HD22	2.18	0.44
57:N1:17:ARG:HG2	57:N1:22:HIS:HA	1.99	0.44
64:N8:2:PRO:O	64:N8:4:ARG:N	2.51	0.44
70:O4:46:ASP:CG	70:O4:80:ARG:HD2	2.38	0.44
73:O7:52:LYS:O	73:O7:56:ARG:HG3	2.17	0.44
73:O7:69:HIS:O	73:O7:73:ARG:HG3	2.18	0.44
2:S0:200:ASP:HA	2:S0:203:PHE:CE2	2.53	0.44
3:S1:130:SER:HB3	3:S1:179:SER:HA	2.00	0.44
4:S2:152:HIS:HA	4:S2:194:GLU:HG3	1.99	0.44
4:S2:57:PHE:CD2	4:S2:138:PRO:HB3	3.41	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:S2:72:LEU:HD13	4:S2:72:LEU:HA	1.72	0.44
5:S3:212:LYS:HB2	5:S3:212:LYS:NZ	3.89	0.44
6:S4:242:LYS:H	6:S4:242:LYS:HE3	1.80	0.44
8:S6:63:MET:HA	8:S6:98:ARG:O	2.28	0.44
8:S6:6:SER:O	8:S6:8:PRO:HD3	2.31	0.44
10:S8:26:LYS:O	10:S8:29:LEU:HB3	2.18	0.44
11:S9:45:ILE:HG22	11:S9:101:VAL:HG12	1.99	0.44
11:S9:113:VAL:O	11:S9:118:LEU:HB2	4.26	0.44
34:SR:147:HIS:HD2	34:SR:151:VAL:HG22	1.83	0.44
34:SR:178:VAL:O	34:SR:192:PHE:HB2	2.53	0.44
34:SR:220:ILE:O	34:SR:234:LEU:N	2.81	0.44
34:SR:222:LEU:HA	34:SR:222:LEU:HD13	2.13	0.44
36:1:1108:U:H2'	36:1:1109:U:H6	1.79	0.44
36:1:1322:U:P	56:N0:117:ARG:HH21	2.41	0.44
36:1:1480:G:H4'	36:1:1481:A:OP1	2.18	0.44
36:1:165:A:H2'	36:1:166:C:C6	2.53	0.44
36:1:2215:A:H2'	36:1:2216:G:O4'	2.17	0.44
36:1:968:G:C6	36:1:969:C:N4	2.86	0.44
1:2:1039:A:O2'	1:2:1040:G:OP2	2.35	0.44
1:2:1172:G:H21	21:C9:88:VAL:HG23	1.83	0.44
1:2:1311:U:C2	1:2:1315:U:C4	3.06	0.44
1:2:149:C:OP1	26:D4:121:THR:OG1	2.33	0.44
1:2:1621:U:H2'	1:2:1622:G:C8	2.53	0.44
1:2:986:G:OP2	39:L2:251:LYS:NZ	2.35	0.44
37:3:55:A:H2'	37:3:56:A:O4'	2.18	0.44
36:5:1027:A:N7	36:5:1029:G:C2	2.86	0.44
36:5:1256:G:H2'	36:5:1257:C:O4'	2.17	0.44
70:O4:59:PRO:HD3	36:5:1654:A:O2'	168.29	0.44
36:5:1926:C:H5'	36:5:1927:G:C5	2.53	0.44
36:5:2197:C:H4'	36:5:2198:A:H8	1.82	0.44
36:5:2359:C:H6	36:5:2359:C:O5'	2.00	0.44
36:5:2413:A:H2'	36:5:2414:G:C8	2.53	0.44
36:5:2530:G:H2'	36:5:2531:C:H5''	2.00	0.44
36:5:3047:U:O2'	36:5:3048:A:H5'	2.18	0.44
36:5:2942:C:O2	86:5:4048:OHX:N6	2.51	0.44
36:5:3377:G:O6	86:5:4082:OHX:N1	2.50	0.44
36:5:414:U:H2'	36:5:415:G:H8	1.82	0.44
36:5:747:A:H2'	36:5:748:U:O4'	2.18	0.44
36:5:951:A:OP2	36:5:1367:G:N2	2.42	0.44
36:5:999:G:H2'	36:5:1000:C:C6	2.53	0.44
1:6:1349:G:O2'	1:6:1379:C:N3	2.34	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:C6:122:ARG:HB3	1:6:1584:G:H5''	397.13	0.44
1:6:215:A:C6	1:6:216:U:C4	3.05	0.44
1:6:1:U:N3	1:6:369:A:C5	2.86	0.44
1:6:66:U:O2'	1:6:67:A:H5''	2.16	0.44
1:6:813:U:H3'	1:6:814:A:H4'	2.00	0.44
1:6:955:A:H2'	1:6:956:C:O4'	2.18	0.44
13:C1:23:PRO:C	13:C1:25:VAL:H	2.65	0.44
16:C4:16:VAL:HG23	16:C4:31:THR:HG23	5.71	0.44
17:C5:15:HIS:CG	17:C5:16:SER:N	3.12	0.44
17:C5:97:TYR:HD2	17:C5:102:PHE:CE1	2.36	0.44
18:C6:11:GLY:N	18:C6:18:ALA:O	2.87	0.44
1:2:1459:C:N4	20:C8:139:LYS:HE2	2.32	0.44
23:D1:71:ARG:HG3	23:D1:83:TRP:CE2	3.48	0.44
23:D1:74:GLN:HG2	23:D1:79:LEU:HB3	4.40	0.44
25:D3:19:ARG:NH2	1:6:611:U:OP1	345.78	0.44
32:E0:46:ASN:HD21	32:E0:48:THR:HG22	4.62	0.44
39:L2:36:GLU:CD	39:L2:163:ARG:HH11	2.38	0.44
39:L2:68:LYS:HD3	39:L2:70:ARG:HG2	2.00	0.44
41:L4:205:PRO:HB3	41:L4:247:PHE:CD2	2.59	0.44
41:L4:265:GLU:OE2	41:L4:266:THR:HG23	2.17	0.44
41:L4:361:HIS:CE1	41:L4:362:ASP:HB2	3.95	0.44
42:L5:220:SER:OG	42:L5:220:SER:O	3.63	0.44
45:L8:143:ILE:HD13	45:L8:170:CYS:SG	2.63	0.44
45:L8:34:PHE:H	45:L8:39:ALA:CB	5.76	0.44
45:L8:33:ASN:O	45:L8:35:GLY:N	3.59	0.44
45:L8:94:PHE:CE1	45:L8:200:LEU:HG	2.53	0.44
46:L9:117:PHE:CE1	46:L9:165:CYS:HB3	3.29	0.44
46:L9:189:GLU:HB3	46:L9:190:ASP:H	1.58	0.44
46:L9:89:LYS:NZ	46:L9:191:LEU:HD12	15.49	0.44
47:M0:36:LEU:HD12	47:M0:87:LEU:HB3	2.85	0.44
48:M1:110:ILE:HG22	48:M1:115:LYS:O	2.18	0.44
49:M3:67:ARG:HH22	64:N8:108:GLY:HA2	3.32	0.44
49:M3:25:HIS:HD2	51:M5:199:LEU:O	2.07	0.44
51:M5:44:ARG:NH1	51:M5:47:LYS:HG2	2.33	0.44
52:M6:27:LEU:HD11	52:M6:102:LEU:CB	2.73	0.44
54:M8:109:GLY:O	54:M8:112:ALA:HB3	2.77	0.44
59:N3:13:ILE:HD11	59:N3:81:GLN:NE2	2.32	0.44
60:N4:57:LYS:HB2	60:N4:57:LYS:HE3	1.87	0.44
62:N6:95:VAL:HA	62:N6:96:PRO:HD3	1.67	0.44
64:N8:90:TYR:CG	64:N8:100:PRO:HG3	2.53	0.44
64:N8:64:GLN:O	64:N8:67:HIS:HB2	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
66:O0:50:VAL:HG11	36:5:2552:C:H2'	233.93	0.44
73:O7:3:LYS:HB3	36:5:2138:A:C5	170.60	0.44
76:Q0:92:ASP:C	76:Q0:93:LYS:HG2	2.81	0.44
79:Q3:78:THR:HA	79:Q3:81:SER:OG	2.18	0.44
2:S0:102:PHE:O	2:S0:103:THR:HB	2.18	0.44
2:S0:70:PRO:O	2:S0:95:ALA:N	2.36	0.44
2:S0:8:ASP:HB3	2:S0:9:LEU:H	2.20	0.44
4:S2:126:ARG:NH1	4:S2:126:ARG:HB3	5.02	0.44
5:S3:103:GLU:HG3	5:S3:107:PHE:CZ	4.03	0.44
5:S3:21:LEU:HD23	5:S3:21:LEU:HA	1.73	0.44
5:S3:52:ALA:O	5:S3:90:ARG:HA	2.17	0.44
6:S4:196:VAL:HG21	6:S4:211:LYS:HB3	1.98	0.44
6:S4:46:VAL:HG13	6:S4:50:ASN:OD1	2.17	0.44
1:2:1167:G:OP1	7:S5:101:GLY:HA3	2.18	0.44
7:S5:77:TYR:CE1	7:S5:87:CYS:HB2	2.53	0.44
34:SR:16:HIS:ND1	34:SR:37:SER:HB3	2.33	0.44
36:1:1534:A:H2'	36:1:1535:A:C8	2.52	0.44
36:1:1664:G:C4	36:1:1786:G:N2	2.86	0.44
36:1:3047:U:O2'	36:1:3048:A:H5'	2.17	0.44
36:1:3182:G:N2	36:1:3189:G:OP1	2.51	0.44
86:1:3965:OHX:N5	86:1:4153:OHX:N1	2.66	0.44
86:1:4128:OHX:N5	86:1:4161:OHX:N6	2.66	0.44
36:1:651:G:C6	36:1:652:G:C6	3.06	0.44
36:1:863:C:H2'	36:1:864:G:O4'	2.17	0.44
1:2:1199:G:P	31:D9:40:ARG:HH21	2.41	0.44
1:2:1215:C:H2'	1:2:1216:C:C6	2.53	0.44
1:2:1387:G:O2'	1:2:1410:A:N6	2.50	0.44
1:2:407:A:O2'	1:2:1671:A:N3	2.44	0.44
1:2:1002:G:N1	1:2:1761:U:OP1	2.39	0.44
1:2:176:C:N4	1:2:266:A:OP2	2.51	0.44
1:2:320:U:H3'	1:2:321:C:C5'	2.45	0.44
1:2:548:G:H2'	1:2:549:G:O4'	2.18	0.44
1:2:565:C:OP2	1:2:577:G:O2'	2.26	0.44
1:2:577:G:H3'	1:2:577:G:C8	2.53	0.44
1:2:704:C:H4'	1:2:705:U:OP1	2.16	0.44
38:4:140:G:H2'	38:4:141:C:O4'	2.18	0.44
38:4:31:G:O2'	38:4:32:C:H5'	2.18	0.44
38:4:52:A:C5	38:4:53:A:C8	3.05	0.44
57:N1:127:GLN:HA	36:5:1095:U:O2	256.73	0.44
36:5:137:G:C2	36:5:138:U:N3	2.86	0.44
36:5:2314:U:O4	86:5:3971:OHX:N5	2.51	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:2623:G:H2'	36:5:2624:G:C8	2.53	0.44
36:5:2623:G:H2'	36:5:2624:G:O4'	2.18	0.44
36:5:776:U:C5	36:5:2719:U:O2	2.71	0.44
36:5:3053:G:C5	36:5:3054:U:C4	3.05	0.44
36:5:314:U:H2'	36:5:315:C:H6	1.76	0.44
36:5:186:U:OP2	86:5:3903:OHX:N4	2.51	0.44
36:5:659:G:H2'	36:5:660:A:N7	2.32	0.44
49:M3:100:ARG:NH1	36:5:76:G:O2'	84.13	0.44
36:5:807:A:H61	36:5:934:G:H22	1.66	0.44
1:6:760:A:OP2	86:6:2083:OHX:N5	2.50	0.44
1:6:294:C:C2	1:6:295:A:C8	3.06	0.44
1:6:63:G:C5	1:6:64:U:C5	3.06	0.44
38:8:83:C:C4'	38:8:85:G:H21	2.30	0.44
38:8:88:A:H3'	38:8:89:A:C8	2.53	0.44
13:C1:29:LYS:NZ	13:C1:32:LYS:HA	2.33	0.44
18:C6:19:VAL:O	18:C6:67:VAL:HA	2.18	0.44
19:C7:24:LEU:HD21	19:C7:34:LEU:HD12	4.24	0.44
19:C7:18:GLU:HG2	19:C7:70:SER:O	4.02	0.44
17:C5:111:MET:HG2	20:C8:119:ILE:CG1	3.58	0.44
24:D2:30:SER:HB2	24:D2:61:ILE:HG13	1.99	0.44
25:D3:92:CYS:HA	25:D3:95:PHE:CD2	2.52	0.44
26:D4:123:LYS:N	26:D4:123:LYS:HZ2	2.16	0.44
28:D6:28:LYS:HG2	28:D6:29:SER:O	2.18	0.44
44:L7:121:LYS:HE2	44:L7:125:GLU:CD	3.30	0.44
44:L7:66:LYS:HG3	44:L7:76:TYR:HD2	1.82	0.44
46:L9:119:GLY:O	46:L9:120:ASP:C	2.55	0.44
47:M0:50:VAL:HG21	47:M0:148:VAL:HG11	3.79	0.44
48:M1:96:PHE:HB2	48:M1:156:LYS:HE2	3.32	0.44
48:M1:92:ARG:NH2	48:M1:94:ARG:HH11	4.84	0.44
49:M3:9:ILE:HD12	64:N8:52:TYR:CE1	3.13	0.44
50:M4:8:LYS:HB3	50:M4:9:ALA:H	1.32	0.44
51:M5:67:ARG:O	51:M5:98:LEU:HD11	2.18	0.44
52:M6:119:VAL:O	52:M6:121:PRO:HD3	2.18	0.44
52:M6:124:LEU:HD23	56:N0:168:PRO:HG3	2.00	0.44
52:M6:38:ALA:O	52:M6:41:LEU:HB2	2.42	0.44
53:M7:25:SER:OG	53:M7:28:ASN:HB2	2.32	0.44
53:M7:31:GLU:HG3	53:M7:60:PHE:HA	3.29	0.44
55:M9:180:LYS:HA	55:M9:183:ALA:HB3	2.00	0.44
55:M9:7:GLN:N	55:M9:7:GLN:OE1	4.16	0.44
56:N0:34:GLU:O	56:N0:38:LYS:HG3	2.18	0.44
56:N0:5:LYS:HB2	56:N0:7:TYR:CE2	2.58	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:N0:91:TYR:OH	56:N0:93:GLU:OE2	2.23	0.44
60:N4:58:HIS:O	60:N4:60:LYS:N	3.44	0.44
63:N7:70:PRO:HG3	63:N7:115:LYS:HB2	1.99	0.44
66:O0:40:LYS:HB3	66:O0:101:LEU:HD11	2.00	0.44
66:O0:30:THR:HG22	66:O0:91:SER:CB	2.99	0.44
66:O0:98:SER:OG	66:O0:99:ASP:N	2.50	0.44
67:O1:55:LEU:O	67:O1:58:ALA:HB3	2.41	0.44
68:O2:9:ILE:HG12	68:O2:63:THR:HB	1.99	0.44
72:O6:94:ILE:CG2	72:O6:99:ARG:HE	2.31	0.44
75:O9:10:LYS:HD3	36:5:1833:G:H5''	108.75	0.44
3:S1:36:SER:HB2	3:S1:231:LEU:O	2.18	0.44
4:S2:65:GLU:HB2	4:S2:68:ILE:HG13	2.79	0.44
6:S4:133:LYS:O	6:S4:134:LYS:HB2	2.18	0.44
6:S4:182:TYR:HB2	6:S4:228:ILE:HD11	1.99	0.44
6:S4:6:LYS:HE2	6:S4:6:LYS:HB2	4.23	0.44
8:S6:14:LYS:HD3	8:S6:16:PHE:CZ	3.46	0.44
8:S6:162:VAL:HG21	8:S6:171:LYS:HG3	5.60	0.44
9:S7:148:LYS:O	9:S7:149:ILE:HG13	3.59	0.44
9:S7:15:GLU:HB2	9:S7:16:LEU:H	1.63	0.44
9:S7:17:GLU:HG3	9:S7:46:ILE:HG13	3.26	0.44
10:S8:193:LEU:HD23	10:S8:193:LEU:HA	1.84	0.44
17:C5:127:ARG:N	35:SM:71:ASN:HD21	3.24	0.44
34:SR:158:PRO:O	34:SR:208:GLY:HA3	2.18	0.44
34:SR:22:SER:CB	34:SR:70:ASP:HA	2.47	0.44
36:1:1456:A:N6	67:O1:64:VAL:HG22	2.33	0.43
36:1:1511:U:H3'	36:1:1512:U:H6	1.83	0.43
36:1:2191:U:H2'	36:1:2192:C:O4'	2.17	0.43
36:1:2213:A:H2'	36:1:2214:A:C8	2.52	0.43
36:1:284:A:OP2	78:Q2:41:ARG:HD2	2.18	0.43
36:1:2883:U:H2'	36:1:2884:C:C6	2.53	0.43
36:1:3060:C:H1'	36:1:3332:U:H1'	1.99	0.43
36:1:1685:C:C2	36:1:3070:A:C2	3.06	0.43
36:1:670:C:OP2	54:M8:147:ARG:NH2	2.51	0.43
1:2:1163:A:C6	1:2:1164:G:C5	3.06	0.43
1:2:1172:G:C6	1:2:1173:C:C4	3.06	0.43
1:2:1776:A:C2	1:2:1786:G:C6	3.06	0.43
1:2:448:C:OP1	6:S4:29:PRO:HD3	2.18	0.43
1:2:476:U:C5	32:E0:31:LYS:HB2	2.52	0.43
1:2:766:U:H3'	1:2:768:C:OP2	2.18	0.43
1:2:888:U:H1'	16:C4:126:THR:HG21	2.00	0.43
36:5:1138:U:H2'	36:5:1139:G:O4'	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:M6:82:LYS:HD3	36:5:1313:G:OP1	254.59	0.43
36:5:1440:G:C4	36:5:1441:G:C8	3.06	0.43
36:5:199:A:C4	36:5:201:A:C8	3.06	0.43
36:5:201:A:OP2	86:5:3980:OHX:N1	2.50	0.43
36:5:2171:G:C2	36:5:2172:A:N7	2.86	0.43
36:5:2248:C:H2'	36:5:2273:G:C8	2.53	0.43
36:5:2764:C:O5'	36:5:2764:C:H6	2.01	0.43
36:5:305:U:C5	36:5:2776:C:H1'	2.52	0.43
86:5:3971:OHX:N4	86:5:4193:OHX:N1	2.66	0.43
36:5:423:A:H2'	36:5:424:G:O4'	2.18	0.43
51:M5:84:PRO:HB2	36:5:44:U:OP1	167.13	0.43
36:5:572:A:C5	36:5:573:C:C5	3.05	0.43
36:5:675:C:O2'	36:5:679:U:OP1	2.29	0.43
36:5:687:U:H2'	36:5:688:G:C8	2.52	0.43
36:5:912:G:O5'	36:5:912:G:H8	2.01	0.43
36:5:993:G:C5	36:5:2637:A:C2	3.06	0.43
1:6:1001:A:H2'	1:6:1002:G:C8	2.53	0.43
1:6:1110:G:N2	1:6:1136:U:H1'	2.33	0.43
1:6:1727:G:H2'	1:6:1728:A:C8	2.53	0.43
11:S9:2:PRO:HB3	1:6:381:C:OP1	361.00	0.43
45:L8:181:LYS:HG2	38:8:154:C:H5''	148.96	0.43
12:C0:32:HIS:CG	12:C0:33:GLU:N	3.67	0.43
17:C5:69:GLU:OE1	17:C5:70:ASN:N	4.94	0.43
18:C6:60:PHE:C	18:C6:62:ASN:H	2.88	0.43
18:C6:20:ALA:HB2	18:C6:84:ALA:HB1	2.30	0.43
18:C6:94:GLN:OE1	34:SR:62:LYS:NZ	2.51	0.43
20:C8:113:LEU:HA	20:C8:113:LEU:HD23	1.41	0.43
21:C9:112:GLY:O	21:C9:125:SER:OG	3.59	0.43
21:C9:93:HIS:O	21:C9:94:ILE:HD12	2.17	0.43
25:D3:51:GLY:HA3	25:D3:74:VAL:CG1	2.48	0.43
7:S5:123:VAL:HG13	27:D5:102:THR:HG23	4.70	0.43
28:D6:62:TYR:O	28:D6:63:ALA:HB2	2.50	0.43
28:D6:87:ARG:HB3	28:D6:91:ASP:HB2	1.99	0.43
39:L2:70:ARG:NE	39:L2:72:ARG:HE	5.16	0.43
40:L3:291:GLU:OE1	40:L3:292:ALA:N	2.51	0.43
40:L3:29:VAL:HG11	40:L3:32:PHE:CE1	2.90	0.43
40:L3:361:THR:CG2	40:L3:371:GLN:HB3	2.47	0.43
40:L3:383:LEU:HD23	40:L3:383:LEU:HA	2.53	0.43
40:L3:53:MET:CG	40:L3:77:THR:HG22	2.45	0.43
41:L4:51:ALA:CB	41:L4:105:THR:HG22	4.58	0.43
41:L4:165:ALA:O	41:L4:168:ALA:HB3	2.37	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:L4:276:LEU:HD12	41:L4:276:LEU:H	3.44	0.43
36:1:364:G:OP1	41:L4:60:THR:HG22	2.17	0.43
42:L5:51:LEU:O	42:L5:147:ASP:N	2.42	0.43
44:L7:173:LEU:HD12	44:L7:173:LEU:HA	1.47	0.43
45:L8:166:LEU:HA	45:L8:166:LEU:HD23	1.83	0.43
48:M1:104:PHE:O	48:M1:127:PHE:HB2	3.07	0.43
49:M3:89:TYR:O	49:M3:92:THR:HG23	3.55	0.43
51:M5:49:ARG:NH1	36:5:149:U:OP2	100.97	0.43
56:N0:87:THR:O	56:N0:88:HIS:ND1	2.66	0.43
57:N1:51:GLY:HA3	57:N1:92:ARG:HG3	4.30	0.43
57:N1:77:ASN:HB3	57:N1:84:TYR:CD2	2.52	0.43
59:N3:69:LEU:HA	59:N3:69:LEU:HD12	1.80	0.43
62:N6:125:LYS:HG3	62:N6:126:LEU:H	1.83	0.43
62:N6:60:ARG:HD3	62:N6:60:ARG:HA	1.71	0.43
63:N7:27:LYS:HD2	63:N7:27:LYS:HA	2.13	0.43
36:1:936:A:OP2	64:N8:27:LYS:HE2	2.18	0.43
64:N8:56:VAL:HG23	64:N8:57:GLY:N	2.32	0.43
67:O1:71:LEU:HD22	67:O1:71:LEU:HA	2.29	0.43
68:O2:16:LYS:O	68:O2:17:PHE:HB2	4.67	0.43
69:O3:89:LEU:HA	69:O3:90:PRO:HD3	2.05	0.43
70:O4:82:ALA:O	70:O4:85:VAL:HG22	4.80	0.43
71:O5:31:LEU:HD21	71:O5:43:LYS:HG2	3.00	0.43
61:N5:46:TYR:CD2	71:O5:75:TYR:HB3	2.53	0.43
74:O8:77:ARG:O	74:O8:78:LEU:HB2	2.17	0.43
79:Q3:24:ARG:O	79:Q3:27:LYS:HB3	2.18	0.43
5:S3:109:LEU:HD23	5:S3:109:LEU:HA	1.56	0.43
5:S3:176:LEU:HD12	5:S3:176:LEU:H	1.85	0.43
6:S4:88:ASP:HB2	6:S4:101:LEU:HD12	3.18	0.43
6:S4:246:LEU:HB2	6:S4:251:GLU:HG3	1.99	0.43
8:S6:85:ARG:HA	8:S6:86:PRO:HD3	1.72	0.43
10:S8:86:SER:O	1:6:341:A:O2'	259.51	0.43
11:S9:161:THR:O	11:S9:162:SER:OG	2.29	0.43
35:SM:32:SER:OG	35:SM:33:LYS:N	4.22	0.43
1:2:577:G:C5	35:SM:99:LYS:HD3	2.52	0.43
34:SR:26:SER:OG	34:SR:75:ALA:O	2.34	0.43
36:1:1047:A:N3	36:1:2633:U:O2'	2.49	0.43
36:1:1668:G:C2	36:1:1669:C:C2	3.06	0.43
36:1:2173:U:H2'	36:1:2174:G:C8	2.53	0.43
36:1:2278:C:H2'	36:1:2279:A:H5''	1.99	0.43
36:1:2144:A:H1'	36:1:2281:A:N6	2.33	0.43
36:1:2353:G:C6	36:1:2354:C:C4	3.06	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:2743:A:C6	36:1:2744:U:C4	3.06	0.43
36:1:2911:A:H4'	36:1:2912:G:C8	2.52	0.43
36:1:3045:G:H2'	36:1:3046:A:O4'	2.19	0.43
36:1:3121:U:C4	36:1:3124:G:O6	2.71	0.43
36:1:608:A:N6	43:L6:22:ARG:HD3	2.33	0.43
36:1:945:C:OP1	68:O2:33:ARG:HG3	2.18	0.43
1:2:1106:U:H2'	1:2:1107:G:H8	1.82	0.43
1:2:1367:G:N7	86:2:2108:OHX:N6	2.66	0.43
1:2:1489:U:C5	1:2:1513:G:C6	3.06	0.43
1:2:187:G:H3'	10:S8:138:ASN:ND2	2.33	0.43
1:2:239:C:H6	1:2:239:C:H2'	1.69	0.43
1:2:30:G:H2'	1:2:31:C:C6	2.52	0.43
1:2:393:C:H2'	1:2:394:C:H6	1.83	0.43
38:4:17:A:H5''	38:4:18:U:OP2	2.18	0.43
36:5:138:U:H2'	36:5:139:G:C8	2.52	0.43
36:5:1552:G:O2'	36:5:1553:U:H5'	2.18	0.43
36:5:1809:A:H2'	36:5:1810:A:O4'	2.18	0.43
36:5:2321:A:C6	36:5:2322:C:N3	2.85	0.43
36:5:277:G:H2'	36:5:278:U:H6	1.82	0.43
36:5:701:G:C5	36:5:702:C:C4	3.06	0.43
1:6:1244:A:N3	1:6:1244:A:H3'	2.33	0.43
1:6:1298:U:H2'	1:6:1299:G:O4'	2.17	0.43
21:C9:102:ARG:NH2	1:6:1502:G:N7	405.80	0.43
18:C6:13:LYS:NZ	1:6:1584:G:O6	398.94	0.43
1:6:40:A:H2'	1:6:41:A:O4'	2.17	0.43
1:6:430:G:N2	1:6:431:C:C2	2.86	0.43
32:E0:31:LYS:NZ	1:6:544:A:O3'	419.04	0.43
1:6:63:G:C6	1:6:64:U:C5	3.05	0.43
1:6:724:C:H2'	1:6:725:U:C6	2.54	0.43
38:8:39:G:H1'	38:8:105:A:N1	2.33	0.43
13:C1:37:ASN:HA	13:C1:44:THR:CG2	2.48	0.43
13:C1:91:LEU:HD22	13:C1:92:HIS:H	1.82	0.43
15:C3:55:ARG:HD2	15:C3:56:ASP:OD1	4.16	0.43
24:D2:50:PHE:HB3	24:D2:63:VAL:HA	2.25	0.43
25:D3:114:LYS:HB3	25:D3:115:GLY:H	1.57	0.43
26:D4:67:GLY:O	26:D4:68:LYS:HB2	2.19	0.43
26:D4:84:LYS:HG3	26:D4:85:PHE:N	2.33	0.43
28:D6:12:LYS:HE2	28:D6:16:GLY:O	2.18	0.43
33:E1:90:LYS:HG3	33:E1:90:LYS:H	4.45	0.43
39:L2:19:HIS:CE1	39:L2:193:ARG:HA	2.53	0.43
39:L2:238:ILE:HB	39:L2:239:ALA:H	3.99	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:L2:96:LEU:HG	39:L2:166:ILE:HD13	2.00	0.43
40:L3:199:PHE:O	40:L3:201:LYS:N	3.02	0.43
40:L3:266:ARG:HB3	40:L3:266:ARG:HE	3.95	0.43
42:L5:202:GLY:O	42:L5:206:GLN:HG3	5.28	0.43
45:L8:238:LEU:HB3	45:L8:242:ALA:HB3	3.68	0.43
45:L8:55:TYR:CE2	45:L8:56:VAL:HG23	3.11	0.43
47:M0:182:LEU:O	47:M0:186:GLU:N	2.40	0.43
47:M0:193:ASP:CG	47:M0:198:LYS:HE3	2.38	0.43
47:M0:205:SER:O	47:M0:209:ASN:HB2	2.17	0.43
50:M4:127:LYS:O	50:M4:131:VAL:HG23	2.63	0.43
52:M6:62:THR:H	52:M6:69:GLY:HA3	1.82	0.43
52:M6:84:LEU:O	52:M6:87:MET:N	2.34	0.43
54:M8:62:VAL:HG11	54:M8:83:VAL:HG21	3.22	0.43
58:N2:93:ILE:HA	58:N2:106:ALA:O	2.17	0.43
62:N6:36:SER:HB2	62:N6:37:LYS:HE2	3.67	0.43
72:O6:9:ILE:HD13	72:O6:10:GLY:N	4.54	0.43
72:O6:34:SER:OG	72:O6:37:THR:OG1	2.21	0.43
75:O9:41:ARG:HG3	75:O9:42:ARG:N	2.33	0.43
2:S0:179:ARG:HD3	2:S0:183:ARG:HD2	2.00	0.43
2:S0:13:ASP:CG	2:S0:179:ARG:HH22	2.25	0.43
4:S2:152:HIS:HB2	4:S2:194:GLU:HB2	2.00	0.43
5:S3:141:LYS:HG2	5:S3:141:LYS:H	2.56	0.43
5:S3:79:TYR:HB3	5:S3:83:THR:OG1	2.18	0.43
9:S7:82:GLU:OE1	9:S7:164:TYR:OH	3.37	0.43
10:S8:43:ILE:O	10:S8:44:HIS:CD2	2.71	0.43
10:S8:98:LYS:HG2	10:S8:99:ALA:N	3.70	0.43
11:S9:182:GLU:HG3	11:S9:182:GLU:H	2.87	0.43
11:S9:49:LEU:O	11:S9:52:ILE:N	3.63	0.43
34:SR:174:ASN:OD1	34:SR:198:ASN:HB3	2.47	0.43
34:SR:217:ASP:OD1	34:SR:217:ASP:N	2.51	0.43
34:SR:67:ILE:HB	34:SR:85:TRP:CG	2.53	0.43
36:1:1578:C:H3'	36:1:1579:C:H6	1.82	0.43
36:1:1944:U:H2'	36:1:1945:A:H8	1.84	0.43
36:1:913:A:H2	36:1:2134:G:N3	2.16	0.43
36:1:2352:A:N6	36:1:2353:G:C6	2.87	0.43
36:1:3297:U:C4	36:1:3298:C:C5	3.06	0.43
36:1:428:A:H2'	36:1:429:U:C6	2.52	0.43
36:1:603:A:C5	36:1:604:G:H1'	2.53	0.43
36:1:993:G:N3	36:1:2637:A:H2'	2.34	0.43
1:2:1128:C:H2'	1:2:1129:U:O4'	2.19	0.43
1:2:1253:U:O2'	33:E1:143:LYS:HA	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:1417:A:H2'	1:2:1418:G:O4'	2.18	0.43
1:2:1450:U:OP1	86:2:2061:OHX:N5	2.50	0.43
1:2:1498:G:C2	1:2:1510:U:O2	2.72	0.43
1:2:344:A:C6	1:2:345:U:C4	3.06	0.43
1:2:811:A:H1'	1:2:858:G:H21	1.84	0.43
1:2:912:U:H5'	1:2:913:G:H8	1.82	0.43
38:4:85:G:H3'	38:4:85:G:H8	1.83	0.43
36:5:1046:A:H2'	36:5:1049:C:C5	2.53	0.43
36:5:1193:A:H2'	36:5:1194:G:O4'	2.19	0.43
36:5:1234:G:OP2	36:5:1235:U:H3'	2.17	0.43
36:5:1556:C:H2'	36:5:2169:G:N1	2.32	0.43
36:5:2376:G:C6	36:5:2377:G:O6	2.71	0.43
36:5:2657:A:C2	36:5:2694:A:C8	3.06	0.43
36:5:3028:G:H2'	36:5:3029:A:O4'	2.18	0.43
36:5:3288:G:O2'	36:5:3289:G:P	2.76	0.43
36:5:902:G:C5	36:5:903:U:C5	3.07	0.43
1:6:1009:U:H2'	1:6:1010:C:C6	2.54	0.43
1:6:1688:U:H2'	1:6:1689:A:C8	2.53	0.43
1:6:198:A:H2'	1:6:199:G:H5'	2.00	0.43
13:C1:19:ILE:HD13	86:6:2125:OHX:N3	295.57	0.43
1:6:1202:A:OP1	86:6:2130:OHX:N2	2.50	0.43
12:C0:29:GLN:HE21	12:C0:29:GLN:HB3	1.60	0.43
13:C1:57:LYS:HB2	13:C1:110:HIS:NE2	2.32	0.43
14:C2:42:ALA:HB3	14:C2:122:VAL:HB	2.61	0.43
15:C3:81:ALA:HA	15:C3:82:PRO:HD2	2.10	0.43
18:C6:25:GLY:N	18:C6:63:ILE:HA	2.86	0.43
18:C6:95:LYS:O	34:SR:59:ARG:NH2	2.51	0.43
19:C7:32:LYS:O	19:C7:35:CYS:HB2	2.51	0.43
20:C8:125:ILE:O	20:C8:129:TRP:N	3.26	0.43
20:C8:14:ILE:HD12	20:C8:14:ILE:HA	4.55	0.43
20:C8:11:PHE:HA	20:C8:59:GLY:O	2.48	0.43
22:D0:21:LYS:HE2	22:D0:21:LYS:HB2	1.62	0.43
24:D2:82:LYS:C	24:D2:84:GLY:H	2.20	0.43
25:D3:139:LYS:O	25:D3:140:LYS:HG3	2.18	0.43
27:D5:90:LYS:HD2	27:D5:104:ALA:HA	2.00	0.43
27:D5:52:LYS:HD3	27:D5:52:LYS:HA	1.84	0.43
28:D6:26:CYS:CB	28:D6:77:CYS:SG	3.06	0.43
30:D8:8:THR:HB	30:D8:56:LEU:HB2	2.87	0.43
31:D9:19:ARG:HD3	31:D9:32:ARG:CD	3.22	0.43
33:E1:126:CYS:HB3	33:E1:130:VAL:HG11	1.99	0.43
36:1:2203:U:H4'	39:L2:241:ARG:HB3	1.99	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:L3:140:ASP:CG	40:L3:141:GLY:H	2.22	0.43
40:L3:227:GLU:CG	40:L3:270:ARG:HE	3.04	0.43
40:L3:299:ASP:O	40:L3:300:ARG:HB2	2.19	0.43
40:L3:58:ARG:O	40:L3:71:GLU:HA	2.43	0.43
41:L4:300:ARG:NH1	41:L4:300:ARG:HB2	2.31	0.43
41:L4:328:ASN:O	41:L4:329:PRO:O	4.15	0.43
42:L5:40:HIS:CE1	42:L5:42:ALA:HB3	2.77	0.43
43:L6:72:ASN:OD1	43:L6:74:VAL:HG23	2.87	0.43
45:L8:73:PRO:HB2	45:L8:230:LYS:O	2.18	0.43
45:L8:91:PHE:CZ	45:L8:189:LEU:HD22	4.71	0.43
46:L9:146:LEU:HD12	46:L9:146:LEU:H	2.36	0.43
46:L9:3:TYR:HA	56:N0:142:GLN:OE1	2.18	0.43
46:L9:74:LEU:HD23	46:L9:74:LEU:HA	1.63	0.43
47:M0:22:TYR:CZ	36:5:1048:A:H2'	267.46	0.43
48:M1:23:VAL:CG1	48:M1:29:ARG:HD3	2.48	0.43
54:M8:159:LYS:HB3	54:M8:159:LYS:HE2	1.82	0.43
54:M8:178:ARG:HA	54:M8:178:ARG:HD3	1.81	0.43
55:M9:115:ILE:HG13	55:M9:119:LEU:HD23	1.99	0.43
55:M9:133:LYS:HD3	55:M9:134:HIS:CD2	2.53	0.43
55:M9:27:ASN:O	86:M9:202:OHX:N6	2.51	0.43
56:N0:141:LYS:HE3	56:N0:141:LYS:HB3	4.63	0.43
50:M4:65:LEU:HB2	56:N0:172:TYR:CZ	2.53	0.43
56:N0:71:LYS:HD2	56:N0:73:LYS:HG3	2.01	0.43
47:M0:168:SER:HA	57:N1:160:ILE:HG23	2.00	0.43
57:N1:52:MET:HG3	57:N1:95:HIS:CE1	2.74	0.43
58:N2:85:LYS:HA	58:N2:85:LYS:HD2	1.70	0.43
63:N7:36:HIS:N	63:N7:37:PRO:HD3	3.08	0.43
36:1:1389:G:OP1	68:O2:101:SER:HB3	2.17	0.43
68:O2:4:LEU:H	68:O2:90:LYS:HB3	1.83	0.43
70:O4:42:PRO:HG2	70:O4:54:ILE:HG22	3.39	0.43
72:O6:62:ARG:HH12	72:O6:98:ARG:HD2	1.83	0.43
73:O7:65:ARG:HH11	73:O7:65:ARG:HG3	1.84	0.43
78:Q2:32:LYS:O	78:Q2:33:ALA:HB3	4.54	0.43
78:Q2:4:VAL:HG22	36:5:2655:U:C5	230.20	0.43
78:Q2:71:ARG:HE	78:Q2:80:ARG:NH2	2.15	0.43
2:S0:31:VAL:HG23	2:S0:150:ASP:HA	2.00	0.43
2:S0:82:GLY:O	2:S0:86:VAL:HG22	2.19	0.43
4:S2:40:LYS:HG3	4:S2:247:ALA:O	7.14	0.43
6:S4:209:HIS:HA	6:S4:219:VAL:HG22	2.00	0.43
9:S7:126:LEU:HD13	9:S7:173:TYR:CE2	3.31	0.43
9:S7:47:ARG:H	9:S7:59:ALA:HB3	1.82	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:S7:71:HIS:CG	9:S7:131:PHE:CZ	3.07	0.43
9:S7:48:GLU:OE2	9:S7:88:ARG:NH2	2.52	0.43
10:S8:178:ARG:HA	10:S8:178:ARG:HD3	1.77	0.43
36:1:1412:G:C6	36:1:1413:G:C5	3.06	0.43
36:1:1506:A:C6	36:1:1510:G:C6	3.06	0.43
36:1:1668:G:C6	36:1:1669:C:C4	3.07	0.43
36:1:2208:A:O3'	36:1:2209:U:H6	2.01	0.43
36:1:2217:U:H2'	36:1:2218:G:C8	2.53	0.43
36:1:2732:G:C6	36:1:2733:A:C4	3.06	0.43
36:1:2948:C:O2'	40:L3:242:THR:HG22	2.18	0.43
36:1:2965:U:H2'	36:1:2966:G:O4'	2.18	0.43
36:1:3051:U:C2	36:1:3052:G:C8	3.06	0.43
36:1:3275:U:O4'	69:O3:66:VAL:HG21	2.19	0.43
36:1:2790:A:O2'	86:1:3976:OHX:N1	2.52	0.43
36:1:698:U:H2'	36:1:699:A:O4'	2.19	0.43
36:1:964:G:HO2'	64:N8:41:HIS:CD2	2.32	0.43
1:2:1285:U:H6	1:2:1285:U:O5'	2.01	0.43
1:2:1499:G:C2	1:2:1500:C:C2	3.06	0.43
1:2:298:C:H5''	1:2:299:A:OP2	2.18	0.43
1:2:36:C:H2'	1:2:37:U:O4'	2.17	0.43
1:2:825:U:H2'	1:2:826:U:C6	2.53	0.43
1:2:912:U:C5'	1:2:913:G:H2'	2.47	0.43
36:5:1302:A:N1	36:5:2832:C:O2'	2.42	0.43
36:5:2406:C:H2'	36:5:2407:C:C6	2.53	0.43
36:5:2943:G:H2'	36:5:2944:U:O4'	2.17	0.43
36:5:3159:C:H2'	36:5:3160:U:H6	1.81	0.43
36:5:547:G:C5	36:5:548:G:H1'	2.53	0.43
36:5:811:U:H2'	36:5:812:G:C8	2.53	0.43
36:5:985:U:H2'	36:5:986:U:C6	2.52	0.43
1:6:1030:A:N7	1:6:1792:G:C2	2.86	0.43
1:6:1271:G:H2'	1:6:1272:U:O4'	2.18	0.43
1:6:141:U:H6	1:6:141:U:H2'	1.69	0.43
1:6:1347:U:C4	1:6:1517:U:O4	2.71	0.43
1:6:1638:G:C6	1:6:1639:C:C2	3.06	0.43
37:7:110:G:C5	37:7:111:U:C4	3.07	0.43
37:7:55:A:H2'	37:7:56:A:O4'	2.18	0.43
1:2:1232:U:H4'	12:C0:2:LEU:HD21	2.00	0.43
13:C1:124:THR:HB	13:C1:141:LYS:HB3	2.01	0.43
15:C3:53:LEU:HA	15:C3:53:LEU:HD12	2.96	0.43
16:C4:121:VAL:HA	16:C4:122:PRO:HD3	2.39	0.43
17:C5:127:ARG:HA	17:C5:130:ARG:CZ	3.30	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:C8:143:ARG:HA	20:C8:145:ARG:HG2	5.58	0.43
21:C9:76:LEU:HB3	21:C9:101:ASN:OD1	4.15	0.43
21:C9:28:LEU:HD12	21:C9:29:GLU:H	1.84	0.43
21:C9:6:VAL:HG22	21:C9:66:TYR:HE1	1.83	0.43
22:D0:99:ILE:H	22:D0:99:ILE:HG12	4.36	0.43
24:D2:11:LEU:HD11	24:D2:37:PHE:CE2	2.70	0.43
26:D4:36:SER:O	26:D4:40:LEU:HB2	2.44	0.43
27:D5:42:LEU:HD22	27:D5:47:TYR:HB2	6.81	0.43
33:E1:91:ILE:HB	1:6:1445:G:C6	387.41	0.43
41:L4:185:LYS:NZ	41:L4:201:GLN:HG2	2.32	0.43
41:L4:31:ARG:O	41:L4:34:ILE:N	2.52	0.43
41:L4:328:ASN:OD1	41:L4:330:TYR:N	2.43	0.43
42:L5:211:LEU:HA	42:L5:211:LEU:HD23	2.07	0.43
44:L7:163:LEU:HA	44:L7:163:LEU:HD23	1.75	0.43
45:L8:133:LYS:HD2	45:L8:138:HIS:CE1	2.53	0.43
46:L9:48:VAL:CG2	46:L9:52:LEU:HB3	3.24	0.43
46:L9:48:VAL:HG23	46:L9:49:ASN:H	4.07	0.43
47:M0:52:LEU:HA	47:M0:52:LEU:HD23	2.04	0.43
48:M1:107:ASP:O	48:M1:108:GLU:HG2	4.37	0.43
48:M1:9:MET:O	48:M1:11:ASP:N	3.73	0.43
50:M4:108:ARG:O	50:M4:110:ALA:N	2.51	0.43
50:M4:12:TRP:CZ2	56:N0:153:PRO:HB3	2.53	0.43
51:M5:106:VAL:O	51:M5:109:ARG:N	2.51	0.43
51:M5:172:ARG:NH1	36:5:30:G:P	107.55	0.43
54:M8:70:ALA:O	54:M8:73:GLN:HB2	2.18	0.43
55:M9:119:LEU:HA	55:M9:119:LEU:HD12	2.30	0.43
1:2:813:U:C2	55:M9:163:ARG:HD2	2.53	0.43
55:M9:174:ALA:HA	55:M9:177:VAL:HG23	4.33	0.43
56:N0:10:ILE:HG12	56:N0:26:ARG:HB2	2.88	0.43
60:N4:38:SER:O	60:N4:42:GLN:N	2.47	0.43
62:N6:102:SER:O	62:N6:103:LYS:HD3	2.18	0.43
63:N7:13:VAL:HG12	63:N7:14:VAL:N	2.92	0.43
66:O0:43:ILE:O	66:O0:89:VAL:HG23	2.18	0.43
72:O6:62:ARG:NH1	72:O6:98:ARG:NH2	6.28	0.43
2:S0:20:ALA:HB1	2:S0:168:HIS:HB2	2.99	0.43
2:S0:64:ILE:HD12	2:S0:181:VAL:HG11	2.68	0.43
6:S4:126:VAL:HG13	6:S4:158:ASP:O	3.29	0.43
1:2:1613:U:OP1	7:S5:169:ASN:HB3	2.18	0.43
7:S5:31:GLU:HA	7:S5:34:GLN:HB2	2.82	0.43
7:S5:73:THR:HG22	7:S5:74:ALA:N	2.42	0.43
8:S6:48:TYR:CD2	8:S6:117:GLY:HA3	2.88	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:S7:186:PRO:HB2	9:S7:187:SER:H	1.58	0.43
9:S7:80:GLU:HG3	9:S7:83:LYS:NZ	4.39	0.43
11:S9:123:HIS:CG	32:E0:37:ARG:HD2	3.97	0.43
35:SM:25:ILE:HA	35:SM:25:ILE:HD13	1.67	0.43
36:1:1109:U:H2'	36:1:1110:U:O4'	2.18	0.43
36:1:1155:C:H2'	36:1:1156:C:H6	1.82	0.43
36:1:1352:A:N3	36:1:1352:A:H5''	2.33	0.43
36:1:1459:C:H2'	36:1:1460:A:O4'	2.19	0.43
36:1:1541:G:H1'	36:1:1557:A:C4	2.53	0.43
36:1:1662:G:O6	86:1:3878:OHX:N2	2.51	0.43
36:1:1717:U:H2'	36:1:1718:G:C8	2.53	0.43
36:1:2254:U:H2'	36:1:2261:G:H22	1.83	0.43
36:1:2275:A:H2'	36:1:2276:G:O4'	2.18	0.43
36:1:2403:G:C2	36:1:2405:C:C4	3.06	0.43
36:1:2993:G:H2'	36:1:3142:A:N6	2.33	0.43
36:1:333:G:H1	38:4:30:C:H42	1.66	0.43
36:1:539:C:H2'	36:1:540:U:C6	2.48	0.43
1:2:1087:A:H2'	1:2:1088:A:C8	2.52	0.43
1:2:1091:A:C8	1:2:1092:A:C6	3.06	0.43
1:2:1338:C:H1'	1:2:1410:A:C4	2.53	0.43
1:2:1364:G:H4'	18:C6:26:LYS:HE2	2.00	0.43
1:2:1725:U:H2'	1:2:1726:G:O4'	2.18	0.43
1:2:17:C:HO2'	1:2:1137:A:N6	2.17	0.43
1:2:48:G:C2	1:2:49:C:C6	3.07	0.43
1:2:546:U:H2'	1:2:547:U:C6	2.54	0.43
1:2:93:A:H2'	1:2:398:G:N2	2.33	0.43
36:5:1017:C:H2'	36:5:1017:C:OP1	2.18	0.43
36:5:1030:A:H2'	36:5:1031:C:C6	2.53	0.43
36:5:132:C:N4	36:5:134:U:O4	2.51	0.43
36:5:1393:A:C8	36:5:1418:A:C6	3.07	0.43
36:5:1614:C:H2'	36:5:1615:C:H6	1.84	0.43
36:5:236:G:H2'	36:5:236:G:N3	2.32	0.43
40:L3:266:ARG:NH2	36:5:2392:C:O2'	209.34	0.43
36:5:2436:U:C2'	36:5:2437:G:H5'	2.47	0.43
36:5:2588:U:C4	36:5:2589:G:N7	2.86	0.43
36:5:320:G:O2'	36:5:321:C:H5'	2.19	0.43
36:5:415:G:H1	38:8:8:C:H42	1.66	0.43
86:5:3971:OHX:N6	86:5:4193:OHX:N3	2.65	0.43
36:5:1196:C:OP1	86:5:4231:OHX:N6	2.51	0.43
1:6:1171:A:H2'	1:6:1172:G:C8	2.52	0.43
1:6:1391:A:H2'	1:6:1392:U:C6	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:1540:G:C5	1:6:1541:G:C8	3.06	0.43
1:6:1684:U:H1'	1:6:1718:G:N2	2.34	0.43
1:6:454:U:H2'	1:6:455:C:C5	2.53	0.43
1:6:462:G:O2'	1:6:463:U:H5'	2.18	0.43
86:8:216:OHX:N5	86:8:223:OHX:N3	2.66	0.43
12:C0:1:MET:HG2	12:C0:2:LEU:N	2.34	0.43
12:C0:10:LYS:HZ2	12:C0:36:ASP:HB3	3.41	0.43
13:C1:129:ARG:HB2	1:6:115:G:C8	310.93	0.43
14:C2:88:LEU:HG	14:C2:89:ILE:H	2.91	0.43
1:2:958:U:O4	15:C3:12:SER:HB3	2.18	0.43
17:C5:122:THR:CB	1:6:1558:U:H3	366.01	0.43
19:C7:27:ASP:OD2	19:C7:30:THR:HG22	2.19	0.43
23:D1:3:ASN:CG	23:D1:7:GLN:HB3	3.51	0.43
27:D5:38:HIS:CG	27:D5:39:ALA:N	2.86	0.43
28:D6:24:VAL:HG12	28:D6:72:HIS:O	2.18	0.43
1:2:542:A:N1	32:E0:28:LYS:HD2	2.33	0.43
39:L2:3:ARG:HG2	39:L2:4:VAL:N	2.33	0.43
41:L4:74:ILE:HG22	41:L4:75:PRO:HD2	2.00	0.43
44:L7:208:SER:O	44:L7:243:MET:HB3	2.19	0.43
45:L8:65:LEU:O	45:L8:69:LEU:HD13	5.37	0.43
46:L9:2:LYS:HZ2	46:L9:59:ASN:HD21	1.65	0.43
49:M3:185:LYS:NZ	49:M3:189:GLU:OE1	5.30	0.43
36:1:3180:A:H5''	52:M6:116:LYS:HB2	2.00	0.43
53:M7:14:SER:HB3	53:M7:150:VAL:O	2.17	0.43
53:M7:57:ALA:O	53:M7:59:PRO:HD3	2.19	0.43
53:M7:61:ARG:HA	53:M7:64:ASN:ND2	3.05	0.43
55:M9:170:ARG:HA	55:M9:173:ARG:HB3	4.24	0.43
46:L9:4:ILE:HD11	56:N0:150:PHE:CD2	3.47	0.43
66:O0:40:LYS:HD3	66:O0:93:LEU:O	2.18	0.43
71:O5:86:ARG:O	71:O5:90:ARG:HG2	2.17	0.43
73:O7:84:SER:O	73:O7:85:LYS:HB3	3.76	0.43
79:Q3:56:THR:HG22	79:Q3:63:THR:CG2	2.48	0.43
3:S1:136:ARG:HB2	3:S1:218:LEU:HD11	4.81	0.43
4:S2:90:THR:N	4:S2:93:GLY:O	2.44	0.43
5:S3:115:ILE:HD12	5:S3:116:ARG:H	4.73	0.43
5:S3:60:GLY:HA3	5:S3:65:ARG:HB2	2.01	0.43
7:S5:190:ILE:O	7:S5:194:LEU:HB2	2.82	0.43
11:S9:59:LEU:HD22	11:S9:69:ARG:HA	2.00	0.43
34:SR:192:PHE:HD1	34:SR:223:TRP:CE3	2.37	0.43
36:1:1470:U:H2'	36:1:1471:U:H6	1.82	0.43
36:1:1743:G:H2'	36:1:1744:G:C8	2.51	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:2288:G:C2	36:1:2289:U:C2	3.06	0.43
36:1:2363:A:C6	36:1:2364:G:C6	3.06	0.43
36:1:2631:U:O2'	36:1:2632:G:H5'	2.19	0.43
36:1:2881:C:H42	36:1:2943:G:H1	1.65	0.43
36:1:385:A:C6	36:1:386:A:C6	3.06	0.43
86:1:4027:OHX:N2	86:1:4040:OHX:N5	2.66	0.43
36:1:586:C:H2'	36:1:587:U:O4'	2.17	0.43
1:2:1036:A:H2'	1:2:1037:C:C6	2.52	0.43
1:2:1278:G:H2'	1:2:1279:C:O4'	2.17	0.43
1:2:1442:U:H2'	1:2:1443:U:C6	2.54	0.43
1:2:1498:G:O2'	1:2:1499:G:H5'	2.18	0.43
1:2:1560:U:O4'	1:2:1560:U:O2	2.35	0.43
1:2:1793:G:O3'	1:2:1794:A:H3'	2.19	0.43
1:2:359:A:C2	25:D3:38:PHE:HB3	2.53	0.43
36:5:1002:A:N1	36:5:1051:U:C6	2.87	0.43
36:5:1130:A:C8	36:5:1132:C:C6	3.07	0.43
36:5:1567:U:HO2'	36:5:1570:U:H5	1.66	0.43
36:5:2554:A:H4'	36:5:2555:G:OP1	2.18	0.43
36:5:2711:C:H4'	86:5:4230:OHX:N1	2.33	0.43
51:M5:179:LYS:HB3	36:5:287:G:H5'	124.95	0.43
36:5:3163:A:C6	36:5:3164:C:N4	2.87	0.43
36:5:3295:A:C2	36:5:3296:A:C4	3.06	0.43
86:5:3994:OHX:N4	86:5:4085:OHX:N1	2.67	0.43
36:5:744:A:H5''	36:5:745:C:OP2	2.19	0.43
1:6:1177:C:H4'	1:6:1189:A:N1	2.33	0.43
1:6:1350:U:H2'	1:6:1351:G:C8	2.53	0.43
20:C8:135:GLY:HA2	1:6:1559:A:H5''	365.17	0.43
1:6:278:U:H2'	1:6:278:U:OP2	2.18	0.43
1:6:294:C:H2'	1:6:295:A:C8	2.54	0.43
1:6:104:A:OP2	1:6:308:C:N4	2.52	0.43
86:5:3984:OHX:N6	38:8:111:A:O2'	2.52	0.43
38:8:46:G:N2	38:8:58:G:C4	2.87	0.43
15:C3:27:LYS:HB2	15:C3:28:LEU:H	1.41	0.43
17:C5:31:GLU:O	17:C5:35:LYS:HD3	2.19	0.43
18:C6:95:LYS:HE3	18:C6:96:TYR:CZ	2.54	0.43
19:C7:24:LEU:CD2	19:C7:34:LEU:HD12	4.06	0.43
21:C9:30:VAL:HA	21:C9:31:PRO:HD2	1.87	0.43
22:D0:106:ILE:O	22:D0:107:THR:OG1	2.34	0.43
22:D0:63:LEU:HB2	22:D0:84:MET:HB3	2.39	0.43
23:D1:20:THR:HB	23:D1:22:ARG:HD3	2.00	0.43
23:D1:58:TYR:OH	24:D2:20:THR:HA	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:D2:113:HIS:O	24:D2:117:ARG:HB2	2.45	0.43
24:D2:87:GLU:HB2	24:D2:117:ARG:HH22	6.58	0.43
25:D3:132:LEU:HA	25:D3:132:LEU:HD22	1.81	0.43
26:D4:128:LYS:HA	26:D4:131:ARG:HG2	1.99	0.43
26:D4:20:ARG:CZ	26:D4:74:LEU:HD22	2.48	0.43
28:D6:23:CYS:C	28:D6:25:ASN:H	2.93	0.43
29:D7:57:GLU:HG3	29:D7:58:SER:N	2.84	0.43
32:E0:51:ASN:HB3	32:E0:53:LYS:HG2	6.03	0.43
39:L2:48:ILE:CD1	39:L2:57:PRO:HB2	2.49	0.43
39:L2:65:ASP:HA	39:L2:66:PRO:HD3	1.84	0.43
39:L2:71:LEU:HA	39:L2:71:LEU:HD12	1.72	0.43
40:L3:128:LYS:HE2	36:5:3151:U:OP1	203.22	0.43
40:L3:252:ILE:HD12	40:L3:252:ILE:HA	1.75	0.43
40:L3:67:PHE:HD1	40:L3:72:VAL:HG12	1.87	0.43
41:L4:144:LYS:H	41:L4:144:LYS:CD	4.40	0.43
41:L4:300:ARG:O	54:M8:39:ARG:NH1	2.51	0.43
41:L4:91:GLY:HA3	41:L4:94:CYS:SG	2.59	0.43
42:L5:211:LEU:HD22	42:L5:215:ASP:HB3	2.71	0.43
42:L5:41:LYS:HD2	42:L5:41:LYS:HA	1.58	0.43
44:L7:156:ILE:HD13	44:L7:172:ASN:OD1	2.19	0.43
44:L7:223:PHE:C	44:L7:225:GLN:H	3.22	0.43
46:L9:90:MET:HB2	46:L9:144:ILE:HG23	2.01	0.43
46:L9:49:ASN:C	46:L9:49:ASN:HD22	2.21	0.43
47:M0:210:ILE:HD13	47:M0:217:PHE:CZ	3.60	0.43
48:M1:82:ARG:HG3	48:M1:112:LEU:HB2	2.00	0.43
48:M1:27:GLY:O	48:M1:31:THR:HG23	3.83	0.43
50:M4:128:ARG:HG2	50:M4:128:ARG:O	2.16	0.43
45:L8:159:PRO:HG3	51:M5:43:THR:O	3.88	0.43
52:M6:147:TRP:CD1	52:M6:148:LYS:N	2.86	0.43
56:N0:30:PHE:CE1	56:N0:103:VAL:HG21	3.15	0.43
56:N0:77:VAL:HG12	56:N0:79:VAL:HG23	2.36	0.43
56:N0:93:GLU:HG2	56:N0:129:ILE:HD11	3.57	0.43
57:N1:72:VAL:CG2	57:N1:96:ILE:HG13	3.07	0.43
59:N3:27:ASP:OD2	59:N3:28:ASN:N	3.07	0.43
62:N6:109:LEU:HD22	62:N6:115:ARG:CZ	2.49	0.43
66:O0:9:SER:OG	66:O0:12:GLN:HB3	5.51	0.43
68:O2:57:TYR:CD1	36:5:1162:U:H4'	197.17	0.43
70:O4:20:ILE:HD12	70:O4:32:ALA:HB1	2.00	0.43
70:O4:57:LEU:HB3	70:O4:61:GLN:OE1	5.25	0.43
36:1:1191:U:OP1	76:Q0:113:ARG:NH2	2.52	0.43
2:S0:110:TYR:C	2:S0:112:THR:H	2.23	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:S1:32:ILE:HB	3:S1:43:VAL:HB	2.76	0.43
4:S2:68:ILE:O	4:S2:72:LEU:HB2	2.26	0.43
9:S7:35:LYS:HB3	9:S7:35:LYS:HE3	1.84	0.43
11:S9:132:ARG:HB2	11:S9:140:ILE:HD13	5.17	0.43
1:2:380:U:H5	11:S9:5:PRO:CB	2.32	0.43
34:SR:221:MET:HG2	34:SR:233:THR:HG23	4.07	0.43
36:1:1579:C:N3	36:1:1580:A:N6	2.66	0.43
36:1:3195:U:O2'	36:1:3197:G:N2	2.51	0.43
86:1:4023:OHX:N6	86:1:4143:OHX:N3	2.67	0.43
36:1:88:A:O5'	36:1:88:A:H8	2.02	0.43
1:2:1107:G:C6	1:2:1108:G:C6	3.07	0.43
1:2:1274:C:C4	35:SM:96:ARG:HG3	2.53	0.43
86:2:2074:OHX:N6	86:2:2162:OHX:N2	2.67	0.43
1:2:387:A:H5''	1:2:389:G:OP2	2.19	0.43
56:N0:90:MET:HG3	36:5:1213:G:H4'	316.45	0.43
36:5:1317:A:C2	36:5:1319:G:C6	3.06	0.43
36:5:2232:A:H2'	36:5:2233:A:O4'	2.19	0.43
36:5:2278:C:C2'	36:5:2279:A:H5''	2.49	0.43
36:5:1131:G:C4	36:5:2373:A:C2	3.07	0.43
36:5:2444:C:N4	36:5:2504:U:O4	2.52	0.43
86:5:3994:OHX:N6	86:5:4085:OHX:N2	2.66	0.43
36:5:612:U:H2'	36:5:613:G:H8	1.84	0.43
1:6:1066:C:H2'	1:6:1067:C:C6	2.54	0.43
24:D2:71:LYS:NZ	1:6:1099:U:OP1	376.38	0.43
19:C7:10:LYS:NZ	1:6:1401:A:O3'	407.46	0.43
1:6:1431:C:H1'	1:6:1437:U:O4	2.19	0.43
1:6:1491:U:H4'	1:6:1492:A:O5'	2.19	0.43
86:6:2059:OHX:N2	86:6:2147:OHX:N6	2.67	0.43
86:6:2059:OHX:N1	86:6:2147:OHX:N4	2.67	0.43
1:6:26:A:HO2'	1:6:27:U:H6	1.64	0.43
1:6:538:A:H8	1:6:543:C:H41	1.58	0.43
1:6:633:U:H2'	1:6:634:G:O4'	2.18	0.43
5:S3:23:GLU:CD	12:C0:61:TRP:HE1	2.22	0.43
15:C3:136:PRO:HA	15:C3:137:PRO:HD2	2.35	0.43
20:C8:116:LEU:HD12	20:C8:121:ALA:HB3	2.00	0.43
21:C9:135:ILE:O	21:C9:138:GLN:HB2	2.19	0.43
5:S3:11:LEU:HD12	22:D0:29:THR:HG23	3.39	0.43
28:D6:92:ARG:C	28:D6:94:ASN:H	2.22	0.43
40:L3:124:LYS:HB2	40:L3:124:LYS:HE3	1.80	0.43
40:L3:312:VAL:O	40:L3:313:HIS:HB2	2.18	0.43
40:L3:35:ASP:OD2	40:L3:37:ARG:NH1	2.48	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:L4:233:LEU:HA	41:L4:233:LEU:HD23	2.23	0.43
42:L5:152:ARG:HG3	42:L5:152:ARG:NH1	2.61	0.43
42:L5:153:THR:HG23	42:L5:160:PHE:CE2	2.54	0.43
43:L6:134:ARG:O	43:L6:137:ASP:N	2.51	0.43
44:L7:103:LEU:HA	44:L7:130:ILE:HD11	4.60	0.43
44:L7:148:VAL:HG12	44:L7:181:ILE:HD11	1.99	0.43
45:L8:109:LEU:O	45:L8:112:GLU:N	2.51	0.43
47:M0:145:LYS:HB3	47:M0:146:ASP:H	1.68	0.43
49:M3:101:ARG:HA	36:5:76:G:O6	87.34	0.43
49:M3:119:TYR:HD1	49:M3:145:PHE:CE2	2.41	0.43
49:M3:47:ALA:HB1	49:M3:48:PRO:CD	2.45	0.43
51:M5:103:GLU:OE1	51:M5:118:SER:OG	2.41	0.43
52:M6:26:GLN:OE1	52:M6:31:GLN:HG2	2.18	0.43
53:M7:65:SER:HB2	36:5:1446:A:H5"	174.30	0.43
53:M7:85:ALA:O	53:M7:89:LYS:HB2	2.85	0.43
54:M8:160:GLY:O	54:M8:161:LYS:HG2	2.19	0.43
54:M8:91:ALA:HB3	64:N8:77:LYS:HE3	4.25	0.43
56:N0:14:LEU:HD23	56:N0:14:LEU:HA	2.28	0.43
56:N0:151:PRO:O	56:N0:153:PRO:HD3	2.62	0.43
57:N1:79:MET:HA	57:N1:84:TYR:HA	2.00	0.43
57:N1:7:TYR:CZ	57:N1:54:HIS:HB2	2.54	0.43
58:N2:89:LEU:O	58:N2:93:ILE:HG13	2.19	0.43
61:N5:67:ILE:HD12	61:N5:121:LYS:HG3	2.06	0.43
61:N5:53:HIS:HB3	61:N5:56:ARG:HH21	1.83	0.43
62:N6:74:TYR:CE1	62:N6:77:LYS:HG3	2.53	0.43
63:N7:107:ARG:O	63:N7:111:LYS:HB2	3.13	0.43
63:N7:89:VAL:HG22	63:N7:93:LYS:HA	3.52	0.43
64:N8:22:ILE:H	64:N8:22:ILE:HG12	2.19	0.43
64:N8:49:HIS:N	64:N8:50:PRO:HD3	2.92	0.43
65:N9:14:ARG:NH2	65:N9:18:ARG:HH11	4.96	0.43
67:O1:13:THR:HG22	67:O1:72:ARG:NH1	2.33	0.43
71:O5:85:THR:HG22	71:O5:87:ALA:N	2.51	0.43
72:O6:53:TYR:O	72:O6:57:LEU:HB2	2.53	0.43
77:Q1:12:ARG:O	77:Q1:15:ARG:N	2.52	0.43
2:S0:34:GLU:OE2	23:D1:87:ARG:NH1	11.20	0.43
3:S1:131:ASP:HB3	3:S1:180:THR:OG1	2.19	0.43
3:S1:119:THR:HB	3:S1:143:THR:HG23	2.31	0.43
4:S2:143:TYR:OH	4:S2:151:PRO:HD3	2.20	0.43
5:S3:156:PHE:C	5:S3:157:LEU:HD12	2.39	0.43
6:S4:113:ARG:H	6:S4:113:ARG:HG3	2.60	0.43
7:S5:49:GLU:O	7:S5:51:VAL:N	2.52	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:S5:41:LYS:HE2	7:S5:67:PRO:HB2	3.57	0.43
8:S6:223:LYS:HA	8:S6:223:LYS:HD3	1.88	0.43
35:SM:135:ALA:C	35:SM:137:GLU:H	2.21	0.43
34:SR:309:VAL:HB	34:SR:311:ARG:NH1	3.03	0.43
34:SR:16:HIS:HE1	34:SR:41:THR:O	2.66	0.43
34:SR:95:ALA:C	34:SR:97:GLY:H	3.89	0.43
36:1:1230:G:N2	36:1:1279:C:N3	2.64	0.43
36:1:1802:C:H2'	36:1:1803:C:O4'	2.18	0.43
36:1:1939:G:C6	36:1:1940:G:C5	3.07	0.43
36:1:2173:U:H2'	36:1:2174:G:N7	2.34	0.43
36:1:2265:C:C2'	36:1:2266:U:H5'	2.49	0.43
36:1:2295:A:OP1	59:N3:63:LYS:NZ	2.52	0.43
36:1:277:G:OP1	86:1:3869:OHX:N5	2.51	0.43
36:1:2988:C:OP1	52:M6:68:ARG:NH1	2.51	0.43
36:1:3033:A:H2'	36:1:3034:C:H6	1.84	0.43
36:1:304:G:H2'	36:1:304:G:N3	2.34	0.43
36:1:3084:C:H2'	36:1:3085:G:O4'	2.19	0.43
36:1:381:U:O4	86:1:4057:OHX:N4	2.51	0.43
36:1:867:G:C6	36:1:868:C:C4	3.07	0.43
1:2:1483:A:H61	1:2:1591:C:C1'	2.31	0.43
86:2:2089:OHX:N5	86:2:2131:OHX:N6	2.66	0.43
1:2:376:C:H2'	1:2:377:G:H8	1.83	0.43
1:2:439:U:O4'	1:2:465:G:N2	2.52	0.43
1:2:521:A:H2'	1:2:522:U:O4'	2.18	0.43
1:2:539:G:OP2	1:2:539:G:C8	2.71	0.43
1:2:538:A:C8	1:2:543:C:N4	2.87	0.43
1:2:819:G:C6	1:2:853:G:C6	3.07	0.43
1:2:872:G:H2'	1:2:873:U:O4'	2.18	0.43
1:2:938:G:N2	1:2:941:A:OP2	2.38	0.43
37:3:79:A:OP2	86:3:218:OHX:N2	2.52	0.43
55:M9:74:ARG:NH1	36:5:1942:U:OP2	210.15	0.43
36:5:2442:G:N1	36:5:2443:A:N7	2.67	0.43
36:5:2520:A:H2'	36:5:2521:U:C6	2.53	0.43
45:L8:47:SER:HB2	36:5:2585:G:O6	168.17	0.43
36:5:2897:A:H2'	36:5:2899:C:C5'	2.48	0.43
36:5:3366:G:H2'	36:5:3367:C:C6	2.54	0.43
86:5:4004:OHX:N4	86:5:4195:OHX:N1	2.67	0.43
36:5:541:U:H2'	36:5:542:G:C8	2.54	0.43
36:5:671:U:H2'	36:5:672:A:C8	2.54	0.43
1:6:1590:G:OP2	86:6:2157:OHX:N6	2.51	0.43
1:6:1765:A:OP2	86:6:2126:OHX:N4	2.52	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:246:G:C6	1:6:247:A:C6	3.06	0.43
1:6:629:U:O2	1:6:971:A:C2	2.72	0.43
1:6:964:U:H4'	1:6:965:U:O4'	2.19	0.43
13:C1:44:THR:O	13:C1:44:THR:OG1	2.32	0.43
15:C3:23:PRO:C	15:C3:25:TRP:H	2.22	0.43
25:D3:103:LEU:HD13	25:D3:104:LEU:N	2.33	0.43
26:D4:18:LEU:HA	26:D4:18:LEU:HD23	1.82	0.43
27:D5:52:LYS:O	27:D5:55:PRO:HD3	2.18	0.43
28:D6:96:ALA:HA	28:D6:97:PRO:HD3	1.73	0.43
30:D8:16:LEU:HB2	30:D8:27:GLN:O	2.19	0.43
33:E1:113:LYS:HB3	33:E1:113:LYS:HE3	1.85	0.43
40:L3:238:LEU:HD12	40:L3:238:LEU:HA	1.72	0.43
40:L3:293:ASN:HB2	40:L3:305:ILE:H	3.43	0.43
40:L3:4:ARG:CG	40:L3:4:ARG:HH11	3.32	0.43
41:L4:23:PRO:HD3	41:L4:255:PHE:CZ	2.53	0.43
43:L6:159:LEU:HD23	43:L6:159:LEU:HA	1.59	0.43
44:L7:110:ARG:HB2	54:M8:2:GLY:O	2.18	0.43
44:L7:144:ILE:H	44:L7:144:ILE:HG13	2.19	0.43
44:L7:147:LEU:HA	44:L7:147:LEU:HD23	1.77	0.43
45:L8:46:LEU:O	45:L8:49:TYR:N	2.62	0.43
47:M0:144:ASN:O	47:M0:148:VAL:HG23	4.13	0.43
47:M0:182:LEU:HD22	47:M0:186:GLU:OE2	2.18	0.43
48:M1:60:ARG:NH1	48:M1:63:GLU:HG3	2.33	0.43
48:M1:81:GLU:HB2	48:M1:167:TYR:HE2	1.84	0.43
51:M5:60:VAL:O	51:M5:61:ILE:HD13	2.18	0.43
52:M6:16:VAL:HG12	52:M6:17:GLY:N	2.59	0.43
36:1:1175:C:O2	52:M6:87:MET:HG2	2.19	0.43
53:M7:64:ASN:O	53:M7:67:ILE:HB	2.18	0.43
53:M7:84:PRO:O	53:M7:87:SER:HB2	2.58	0.43
55:M9:41:ILE:HA	55:M9:41:ILE:HD13	3.83	0.43
36:1:3068:U:P	55:M9:59:SER:HG	2.41	0.43
56:N0:82:ASP:HB2	56:N0:120:SER:HB3	2.13	0.43
56:N0:68:HIS:HA	56:N0:69:PRO:HD3	1.80	0.43
57:N1:103:GLN:O	57:N1:107:GLU:HB2	2.58	0.43
58:N2:15:PHE:CE2	58:N2:71:PHE:HD1	2.71	0.43
59:N3:46:LEU:O	59:N3:47:ASN:HB2	2.18	0.43
60:N4:46:PRO:HB3	60:N4:54:LEU:HD13	2.00	0.43
62:N6:103:LYS:HA	62:N6:103:LYS:HD3	2.01	0.43
62:N6:108:LYS:HA	62:N6:108:LYS:HD3	2.42	0.43
62:N6:38:GLU:O	62:N6:41:ALA:HB3	3.44	0.43
62:N6:69:LYS:HB2	62:N6:69:LYS:HE3	4.26	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
63:N7:46:ILE:HD11	63:N7:49:TYR:CE2	2.84	0.43
64:N8:84:GLU:O	64:N8:87:ARG:HB2	2.25	0.43
64:N8:6:THR:HG22	64:N8:9:ARG:HG2	2.00	0.43
65:N9:43:HIS:NE2	65:N9:47:LEU:HD21	3.25	0.43
66:O0:12:GLN:O	66:O0:16:LEU:N	2.52	0.43
67:O1:23:VAL:HG12	67:O1:24:SER:O	2.19	0.43
68:O2:82:LEU:HD12	68:O2:108:ILE:CG2	4.02	0.43
36:1:1407:A:O3'	68:O2:33:ARG:NH2	2.51	0.43
36:1:1591:G:OP2	70:O4:17:SER:HB3	2.19	0.43
70:O4:44:CYS:SG	70:O4:46:ASP:HB2	2.59	0.43
70:O4:66:SER:HB2	70:O4:69:HIS:ND1	4.58	0.43
74:O8:27:ILE:H	74:O8:78:LEU:HD11	1.83	0.43
2:S0:25:GLY:N	2:S0:46:HIS:O	2.65	0.43
2:S0:86:VAL:HG12	2:S0:174:TRP:CZ2	3.27	0.43
3:S1:171:ILE:O	3:S1:175:GLU:HB2	3.57	0.43
3:S1:179:SER:O	3:S1:182:ALA:HB3	2.19	0.43
4:S2:218:ILE:HG13	4:S2:218:ILE:H	1.56	0.43
4:S2:90:THR:O	4:S2:92:ALA:N	2.52	0.43
4:S2:94:GLN:HG2	4:S2:95:ARG:H	4.42	0.43
5:S3:161:GLY:O	5:S3:164:VAL:HB	2.18	0.43
6:S4:180:LEU:HD23	6:S4:194:THR:H	1.84	0.43
6:S4:26:CYS:HB2	6:S4:27:TYR:CD2	2.70	0.43
7:S5:50:GLU:HB2	7:S5:51:VAL:H	1.70	0.43
8:S6:22:HIS:HA	8:S6:25:ARG:HB2	2.01	0.43
1:2:405:C:O2'	8:S6:92:ARG:O	2.27	0.43
9:S7:182:VAL:HG12	9:S7:183:PHE:N	2.33	0.43
9:S7:154:LEU:HD21	9:S7:183:PHE:CD1	2.53	0.43
9:S7:74:GLN:NE2	9:S7:92:PHE:HB2	3.12	0.43
1:2:1274:C:N4	35:SM:95:SER:HA	2.33	0.43
34:SR:228:LYS:HE3	34:SR:228:LYS:HB2	2.40	0.43
34:SR:276:PRO:O	34:SR:278:PHE:N	3.33	0.43
34:SR:281:TYR:HB3	34:SR:282:SER:H	1.47	0.43
36:1:1213:G:C2	36:1:1293:U:C2	3.07	0.43
36:1:1428:A:OP2	64:N8:2:PRO:HB2	2.18	0.43
36:1:1488:G:H5''	36:1:1838:G:O6	2.19	0.43
36:1:1764:U:OP1	55:M9:43:LYS:HD3	2.19	0.43
36:1:201:A:H2'	36:1:202:G:C8	2.54	0.43
36:1:2653:C:OP1	78:Q2:89:LYS:HB2	2.18	0.43
36:1:2763:U:O5'	36:1:2763:U:H6	2.01	0.43
36:1:2945:G:O2'	36:1:2948:C:OP2	2.32	0.43
36:1:3007:U:OP1	52:M6:73:PHE:HA	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:3018:C:C5	36:1:3019:U:C5	3.07	0.43
36:1:3100:U:O2	36:1:3101:G:C8	2.72	0.43
36:1:2810:C:OP1	86:1:4078:OHX:N6	2.52	0.43
36:1:520:U:C4	44:L7:67:ARG:HD3	2.54	0.43
36:1:532:A:H2	36:1:560:G:H22	1.67	0.43
36:1:634:C:H5'	69:O3:21:ARG:O	2.19	0.43
36:1:839:C:H1'	36:1:1724:U:OP1	2.18	0.43
36:1:915:A:C2'	36:1:915:A:N3	2.82	0.43
1:2:1193:A:OP1	1:2:1193:A:H8	2.01	0.43
1:2:1484:G:O4'	1:2:1607:G:H4'	2.19	0.43
1:2:1520:U:H5''	21:C9:75:LYS:HZ3	1.84	0.43
1:2:1521:G:O2'	1:2:1523:G:OP2	2.15	0.43
1:2:373:G:N7	86:2:2159:OHX:N6	2.67	0.43
1:2:398:G:OP2	10:S8:47:ARG:NH1	2.48	0.43
1:2:702:G:O2'	1:2:703:G:H8	2.00	0.43
1:2:710:U:H2'	1:2:711:U:H5'	2.01	0.43
1:2:854:U:O2'	1:2:855:A:H5'	2.19	0.43
1:2:926:A:H2'	1:2:927:C:O4'	2.19	0.43
37:3:28:C:H5''	48:M1:137:ARG:HG2	2.01	0.43
36:5:998:A:H2	36:5:1051:U:O4	2.02	0.43
36:5:1573:G:C6	36:5:1574:C:H1'	2.54	0.43
36:5:1700:G:C6	36:5:1701:C:C4	3.06	0.43
36:5:1733:G:N7	86:5:3962:OHX:N6	2.67	0.43
36:5:1734:G:O6	86:5:3962:OHX:N5	2.52	0.43
36:5:2652:U:C4	36:5:2759:U:O2	2.71	0.43
48:M1:57:PHE:CD2	36:5:2680:A:C4	310.54	0.43
36:5:3280:U:O2'	36:5:3281:U:H5''	2.18	0.43
36:5:3308:C:N4	36:5:3309:G:C6	2.87	0.43
86:5:4029:OHX:N6	86:5:4232:OHX:N5	2.67	0.43
36:5:3128:G:OP2	86:5:4154:OHX:N5	2.52	0.43
36:5:624:G:H2'	36:5:625:G:C8	2.53	0.43
1:6:1147:A:H2'	1:6:1148:C:C6	2.54	0.43
1:6:1153:G:H1	1:6:1625:C:N4	2.16	0.43
1:6:1172:G:H2'	1:6:1173:C:C6	2.54	0.43
1:6:1391:A:C8	1:6:1412:G:C6	3.06	0.43
7:S5:184:PHE:CE2	1:6:1471:A:H5'	341.11	0.43
1:6:1203:A:C6	1:6:1555:A:C6	3.07	0.43
1:6:1697:G:H2'	1:6:1697:G:N3	2.34	0.43
1:6:770:A:OP2	86:6:2138:OHX:N3	2.52	0.43
1:6:234:G:H2'	1:6:235:G:O4'	2.18	0.43
1:6:358:U:O2'	1:6:360:A:H5''	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:877:G:H5'	1:6:937:C:H1'	2.00	0.43
42:L5:33:ARG:NH2	37:7:7:G:O3'	270.05	0.43
38:8:37:A:C6	38:8:104:A:C5	3.06	0.43
12:C0:76:LEU:HD22	12:C0:76:LEU:H	1.84	0.43
13:C1:46:LYS:O	13:C1:50:GLU:HG2	4.75	0.43
15:C3:92:ILE:HA	15:C3:122:ILE:HD11	2.58	0.43
16:C4:14:PHE:HA	16:C4:78:ALA:O	2.53	0.43
18:C6:45:ARG:O	18:C6:48:VAL:HG12	2.57	0.43
18:C6:60:PHE:CZ	18:C6:89:LEU:HD13	2.54	0.43
19:C7:12:ALA:O	19:C7:15:ALA:N	2.51	0.43
20:C8:100:THR:O	20:C8:101:LEU:HD23	2.61	0.43
20:C8:117:LYS:O	20:C8:120:ARG:HG3	5.30	0.43
20:C8:131:LEU:HA	20:C8:131:LEU:HD23	2.32	0.43
21:C9:38:LYS:HD2	21:C9:40:SER:O	3.87	0.43
1:2:1601:G:H22	21:C9:88:VAL:HG22	1.82	0.43
2:S0:184:LEU:HD13	23:D1:45:ALA:HB2	2.01	0.43
24:D2:114:GLU:O	24:D2:118:ARG:HG3	2.25	0.43
25:D3:109:ARG:HB3	25:D3:109:ARG:CZ	4.75	0.43
39:L2:54:ARG:HG2	39:L2:55:GLY:O	4.64	0.43
41:L4:120:TYR:HD1	41:L4:120:TYR:O	2.02	0.43
41:L4:191:LYS:HD3	41:L4:194:TYR:OH	2.32	0.43
44:L7:140:SER:O	44:L7:144:ILE:HG13	2.69	0.43
39:L2:64:ARG:HH12	45:L8:38:GLN:HA	1.84	0.43
47:M0:170:LYS:HG3	47:M0:175:ASN:C	4.46	0.43
47:M0:92:HIS:HA	47:M0:93:PRO:HD3	1.89	0.43
50:M4:118:PHE:O	50:M4:122:VAL:HG23	2.18	0.43
50:M4:24:LYS:HG2	50:M4:62:GLN:C	2.39	0.43
51:M5:177:GLY:H	51:M5:184:LYS:NZ	2.17	0.43
53:M7:62:ARG:O	86:M7:206:OHX:N1	2.52	0.43
53:M7:94:LEU:HD13	53:M7:94:LEU:N	2.34	0.43
56:N0:14:LEU:HA	56:N0:15:PRO:HD3	2.02	0.43
56:N0:33:ASN:C	56:N0:35:VAL:N	2.72	0.43
63:N7:49:TYR:CE2	63:N7:133:LYS:HA	2.87	0.43
67:O1:17:HIS:HB2	67:O1:69:TYR:HB3	2.01	0.43
71:O5:114:ARG:O	71:O5:116:TYR:HD2	2.75	0.43
72:O6:42:SER:OG	72:O6:43:LEU:N	3.67	0.43
2:S0:109:ASN:OD1	2:S0:111:ILE:HG22	2.19	0.43
3:S1:178:GLY:HA3	3:S1:187:LYS:NZ	2.34	0.43
4:S2:160:GLY:O	4:S2:166:THR:HA	2.32	0.43
6:S4:194:THR:HG23	6:S4:211:LYS:O	3.42	0.43
7:S5:216:GLU:HG3	7:S5:219:ARG:NH2	4.99	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:S6:21:GLU:O	8:S6:25:ARG:HG3	4.13	0.43
9:S7:49:ILE:HD11	9:S7:172:VAL:HG22	2.01	0.43
11:S9:127:VAL:O	11:S9:130:THR:HG22	2.19	0.43
34:SR:161:LYS:HE3	34:SR:164:ASP:CB	2.45	0.43
36:1:1177:G:H1'	36:1:1178:G:N7	2.33	0.43
36:1:824:C:O2'	36:1:1534:A:N3	2.48	0.43
36:1:1562:C:H2'	36:1:1563:C:C6	2.54	0.43
36:1:1670:C:H1'	36:1:1780:G:N2	2.34	0.43
36:1:1927:G:OP1	79:Q3:8:VAL:HG13	2.19	0.43
36:1:2723:U:H2'	36:1:2724:U:H6	1.82	0.43
36:1:2419:A:H1'	36:1:2804:A:O4'	2.19	0.43
36:1:3113:A:H4'	46:L9:69:ARG:HB3	2.00	0.43
36:1:1119:C:OP2	86:1:3947:OHX:N1	2.52	0.43
86:1:3998:OHX:N3	86:1:4169:OHX:N3	2.67	0.43
36:1:522:A:C2	36:1:523:A:H1'	2.54	0.43
36:1:657:A:H2'	36:1:658:G:H8	1.83	0.43
1:2:1017:U:H2'	1:2:1018:U:H6	1.84	0.43
1:2:1036:A:H2'	1:2:1037:C:H6	1.82	0.43
1:2:154:G:C5'	8:S6:108:VAL:HG21	2.49	0.43
1:2:14:C:H2'	1:2:15:U:C6	2.54	0.43
1:2:170:U:H6	1:2:267:U:HO2'	1.67	0.43
1:2:651:G:H2'	1:2:652:G:O4'	2.19	0.43
37:3:30:G:C6	37:3:31:U:C4	3.07	0.43
52:M6:131:PRO:HD3	36:5:1316:C:N4	299.37	0.43
36:5:1716:U:H5'	36:5:1716:U:H6	1.83	0.43
36:5:1700:G:N2	36:5:1745:C:N3	2.63	0.43
36:5:1764:U:H3'	36:5:1765:U:H5''	2.01	0.43
36:5:2101:C:O2'	36:5:2102:U:P	2.76	0.43
36:5:2568:C:N4	36:5:2574:G:C6	2.86	0.43
36:5:2842:U:C4	36:5:2843:U:C5	3.06	0.43
36:5:2876:C:H42	36:5:2951:G:H1	1.66	0.43
36:5:3111:U:O4	36:5:3121:U:H5	2.01	0.43
36:5:3121:U:H1'	36:5:3122:A:H5''	2.01	0.43
86:5:4004:OHX:N3	86:5:4195:OHX:N5	2.67	0.43
56:N0:146:LYS:HA	36:5:534:U:O2	350.07	0.43
1:6:1248:C:H2'	1:6:1249:U:C6	2.53	0.43
1:6:1562:G:C6	1:6:1563:C:C4	3.07	0.43
1:6:1699:G:N2	1:6:1702:A:H5''	2.33	0.43
1:6:400:A:H4'	1:6:401:A:H5'	2.01	0.43
1:6:405:C:H6	1:6:405:C:O5'	2.02	0.43
38:8:83:C:H4'	38:8:85:G:H21	1.83	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:C1:90:TYR:HE1	13:C1:92:HIS:HB2	1.84	0.43
17:C5:130:ARG:HD3	35:SM:74:LYS:HG2	2.01	0.43
19:C7:46:LEU:O	19:C7:50:ILE:HG13	2.19	0.43
20:C8:72:ILE:HA	20:C8:79:TYR:HE2	4.30	0.43
21:C9:16:ASN:HA	21:C9:56:LYS:HZ1	2.48	0.43
18:C6:33:GLY:O	21:C9:7:ARG:HB3	2.83	0.43
4:S2:229:LEU:HD23	23:D1:23:ILE:HD11	2.77	0.43
2:S0:157:ASP:OD2	23:D1:60:ARG:HD2	2.19	0.43
27:D5:49:ARG:HD2	27:D5:53:GLU:OE1	4.25	0.43
30:D8:22:ARG:HA	30:D8:22:ARG:HD3	1.58	0.43
7:S5:163:SER:HB2	30:D8:48:VAL:HG22	2.01	0.43
33:E1:144:CYS:C	33:E1:146:SER:N	2.72	0.43
39:L2:129:ALA:HB3	39:L2:132:ASN:OD1	2.19	0.43
40:L3:81:THR:HG23	40:L3:205:VAL:CG2	4.48	0.43
36:1:208:C:OP2	41:L4:163:LYS:NZ	2.50	0.43
41:L4:22:LEU:CD2	41:L4:26:PHE:HB2	2.48	0.43
42:L5:250:ASP:HA	42:L5:251:PRO:HD3	1.74	0.43
42:L5:37:VAL:CG1	57:N1:31:LEU:HD21	2.49	0.43
43:L6:9:TRP:CZ2	36:5:1354:G:C6	179.26	0.43
44:L7:125:GLU:O	44:L7:129:LEU:HB2	2.19	0.43
44:L7:176:TYR:CD2	44:L7:194:HIS:CD2	3.07	0.43
44:L7:68:ASP:O	44:L7:71:ALA:HB3	2.19	0.43
45:L8:226:TYR:O	45:L8:227:ASP:C	2.56	0.43
36:1:3186:A:O2'	46:L9:42:ASP:HA	2.19	0.43
49:M3:33:VAL:HG12	49:M3:34:SER:N	2.33	0.43
50:M4:108:ARG:HH21	52:M6:197:LEU:HA	3.81	0.43
50:M4:25:LYS:CD	50:M4:62:GLN:HB3	2.49	0.43
51:M5:91:GLU:O	51:M5:93:LYS:HE2	2.19	0.43
52:M6:41:LEU:HD12	52:M6:41:LEU:HA	1.80	0.43
53:M7:27:LYS:HA	53:M7:63:PHE:CD2	2.79	0.43
53:M7:72:GLN:OE1	53:M7:83:TRP:NE1	3.08	0.43
54:M8:88:THR:HG22	54:M8:107:THR:HG21	2.00	0.43
54:M8:147:ARG:NH2	36:5:670:C:OP1	162.70	0.43
54:M8:76:ALA:HA	54:M8:79:LYS:HD3	4.43	0.43
57:N1:134:GLN:HB3	57:N1:134:GLN:HE21	4.14	0.43
57:N1:41:ASP:OD1	57:N1:61:THR:OG1	2.90	0.43
58:N2:101:ASN:O	58:N2:102:GLU:HG3	3.84	0.43
61:N5:133:LEU:HD22	61:N5:133:LEU:HA	2.24	0.43
62:N6:86:THR:HG22	62:N6:96:PRO:HA	2.01	0.43
66:O0:27:TYR:O	66:O0:31:VAL:HG23	2.19	0.43
67:O1:100:SER:OG	67:O1:102:LYS:HB3	3.11	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
68:O2:101:SER:OG	68:O2:103:LYS:HG2	2.19	0.43
69:O3:75:HIS:HB3	69:O3:80:VAL:HG12	2.00	0.43
77:Q1:4:LYS:HG2	77:Q1:5:TRP:CZ3	2.53	0.43
2:S0:146:LEU:HB3	2:S0:162:CYS:SG	3.19	0.43
2:S0:67:ILE:HA	2:S0:67:ILE:HD13	2.01	0.43
3:S1:197:ILE:HB	3:S1:210:ILE:HG21	2.55	0.43
4:S2:139:ILE:CD1	4:S2:191:ALA:HB1	2.55	0.43
4:S2:99:LYS:HB2	4:S2:117:THR:HB	2.06	0.43
1:2:1332:C:O2'	5:S3:162:GLN:HB3	2.19	0.43
6:S4:42:LEU:HA	6:S4:42:LEU:HD23	1.67	0.43
6:S4:51:ARG:HB3	6:S4:111:VAL:CG2	2.48	0.43
7:S5:93:LEU:HD22	7:S5:93:LEU:HA	1.90	0.43
8:S6:28:PHE:O	8:S6:30:LYS:HG3	3.67	0.43
9:S7:114:ARG:O	9:S7:116:ARG:N	2.47	0.43
9:S7:137:GLY:HA2	15:C3:18:TYR:CE2	2.54	0.43
10:S8:182:TYR:H	10:S8:182:TYR:HD2	3.95	0.43
36:1:1101:G:H1'	44:L7:105:LEU:CD2	2.49	0.42
36:1:111:C:O2'	36:1:112:U:H5'	2.18	0.42
36:1:1246:G:H22	36:1:1265:U:H5	1.67	0.42
36:1:1524:A:C2	36:1:1527:C:C6	3.07	0.42
36:1:2288:G:N2	36:1:2289:U:C2	2.87	0.42
36:1:2512:C:C4	36:1:2513:U:O4	2.72	0.42
36:1:2579:G:O6	86:1:3919:OHX:N2	2.52	0.42
36:1:2726:C:O2'	36:1:2727:A:H2'	2.19	0.42
36:1:3210:A:H2'	36:1:3211:C:O4'	2.19	0.42
36:1:550:A:N6	36:1:551:A:N6	2.66	0.42
36:1:679:U:H2'	36:1:680:G:C8	2.54	0.42
1:2:1080:U:H2'	1:2:1081:A:C8	2.54	0.42
1:2:1081:A:H5''	1:2:1082:C:OP1	2.18	0.42
1:2:393:C:H2'	1:2:394:C:C6	2.54	0.42
1:2:41:A:H2'	1:2:438:A:N7	2.34	0.42
1:2:462:G:C6	1:2:463:U:C4	3.07	0.42
1:2:498:G:C4	1:2:499:U:N3	2.87	0.42
1:2:4:C:H2'	1:2:5:U:H6	1.84	0.42
1:2:852:C:N4	1:2:853:G:C6	2.87	0.42
1:2:855:A:H3'	1:2:856:A:H5''	2.00	0.42
1:2:924:A:H2'	1:2:925:G:C8	2.53	0.42
37:3:103:A:N1	37:3:104:A:C4	2.87	0.42
38:4:98:U:C2'	38:4:99:C:H5'	2.49	0.42
36:5:1019:G:N1	36:5:1020:G:O6	2.51	0.42
36:5:1101:G:H2'	36:5:1102:A:O4'	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:1235:U:C4'	36:5:1236:G:H5'	2.49	0.42
36:5:1611:G:H2'	36:5:1612:A:H8	1.84	0.42
36:5:1790:G:H2'	36:5:1791:C:O4'	2.19	0.42
36:5:2567:C:H42	36:5:2568:C:H41	1.67	0.42
36:5:1941:C:O2'	36:5:3344:A:N1	2.36	0.42
1:6:1157:A:C2	1:6:1160:A:C8	3.07	0.42
19:C7:52:GLY:HA3	1:6:1389:C:O2'	423.89	0.42
1:6:1584:G:H22	1:6:1611:A:P	2.40	0.42
1:6:1733:C:O2'	1:6:1734:U:H5'	2.19	0.42
1:6:603:U:H2'	1:6:604:A:H8	1.84	0.42
12:C0:68:LEU:HD12	12:C0:69:THR:N	2.34	0.42
12:C0:8:ARG:HG3	12:C0:12:HIS:ND1	2.34	0.42
13:C1:90:TYR:CD1	13:C1:91:LEU:N	2.82	0.42
14:C2:58:LEU:HG	14:C2:126:TRP:CZ3	6.15	0.42
15:C3:18:TYR:HE1	24:D2:54:ASP:OD2	3.92	0.42
15:C3:151:ASN:O	86:C3:201:OHX:N3	4.46	0.42
21:C9:6:VAL:C	21:C9:8:ASP:H	2.45	0.42
25:D3:36:THR:HG23	25:D3:40:SER:OG	2.18	0.42
26:D4:41:ARG:HG2	26:D4:55:VAL:HG12	2.01	0.42
26:D4:34:ASN:OD1	26:D4:62:THR:HG21	4.02	0.42
27:D5:81:ARG:HH11	27:D5:81:ARG:HB2	3.19	0.42
14:C2:46:ARG:HH21	33:E1:102:VAL:HG22	7.90	0.42
39:L2:159:SER:C	39:L2:161:ASP:H	2.53	0.42
36:1:790:U:H4'	41:L4:112:LYS:O	2.19	0.42
41:L4:3:ARG:CB	41:L4:22:LEU:H	3.89	0.42
41:L4:330:TYR:O	41:L4:331:ALA:C	2.61	0.42
42:L5:218:ARG:HA	42:L5:221:GLU:OE2	2.19	0.42
44:L7:159:GLN:O	44:L7:160:ARG:HB3	2.19	0.42
44:L7:40:LYS:O	44:L7:44:ILE:HG13	2.19	0.42
46:L9:161:LEU:O	46:L9:164:ILE:HG22	2.18	0.42
47:M0:38:LYS:CG	47:M0:41:ALA:HB2	2.59	0.42
47:M0:91:VAL:HG12	47:M0:91:VAL:O	2.19	0.42
48:M1:91:LEU:HA	48:M1:91:LEU:HD23	1.87	0.42
51:M5:118:SER:HB3	51:M5:132:VAL:HA	2.61	0.42
51:M5:149:ASN:OD1	86:M5:303:OHX:N2	2.52	0.42
51:M5:155:VAL:HG22	36:5:58:G:H4'	81.92	0.42
51:M5:38:ARG:HB2	51:M5:62:TYR:CZ	2.54	0.42
53:M7:70:THR:HG23	53:M7:72:GLN:H	1.84	0.42
53:M7:88:VAL:O	53:M7:92:GLN:HG3	3.88	0.42
54:M8:178:ARG:HA	54:M8:178:ARG:HD2	2.62	0.42
41:L4:29:PRO:HB3	54:M8:25:TYR:HE2	1.84	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:854:U:O4	55:M9:173:ARG:NH2	2.52	0.42
56:N0:148:LEU:HD12	56:N0:149:LYS:N	2.34	0.42
56:N0:37:ALA:O	56:N0:40:ARG:HB2	2.19	0.42
36:1:1068:C:H5'	57:N1:110:LYS:HZ1	1.83	0.42
59:N3:77:ILE:O	59:N3:100:GLY:HA2	2.19	0.42
61:N5:91:ASN:OD1	61:N5:93:TYR:HB2	2.18	0.42
62:N6:56:VAL:HG21	62:N6:104:LEU:HB3	2.01	0.42
63:N7:50:PRO:HB3	63:N7:66:THR:O	3.33	0.42
67:O1:14:ILE:HG13	67:O1:39:PHE:CD1	3.16	0.42
69:O3:81:VAL:HG22	69:O3:82:ARG:N	3.85	0.42
72:O6:26:ILE:HD12	72:O6:27:SER:N	2.34	0.42
51:M5:15:GLN:HB3	72:O6:52:PRO:HD3	3.68	0.42
72:O6:99:ARG:H	72:O6:99:ARG:HG3	4.14	0.42
74:O8:12:LEU:C	74:O8:14:LEU:N	3.23	0.42
78:Q2:93:LEU:HD13	78:Q2:93:LEU:O	5.19	0.42
79:Q3:17:ARG:O	79:Q3:19:GLY:N	2.82	0.42
79:Q3:77:ALA:O	79:Q3:80:ARG:HB3	2.19	0.42
79:Q3:87:ARG:O	79:Q3:90:VAL:HG22	4.01	0.42
2:S0:103:THR:HA	2:S0:104:PRO:HD3	1.60	0.42
1:2:1277:G:H5'	5:S3:140:GLY:HA2	2.01	0.42
6:S4:148:ARG:H	6:S4:148:ARG:HG2	2.63	0.42
8:S6:150:GLU:O	8:S6:152:ASP:N	2.51	0.42
8:S6:5:ILE:HD13	8:S6:50:PHE:CE1	2.54	0.42
9:S7:118:LEU:N	1:6:639:U:OP1	366.91	0.42
9:S7:71:HIS:CD2	9:S7:131:PHE:HZ	2.36	0.42
9:S7:71:HIS:CE1	9:S7:131:PHE:CE1	3.18	0.42
9:S7:89:HIS:HD2	9:S7:165:LYS:HG2	1.84	0.42
11:S9:120:LYS:O	11:S9:120:LYS:HD3	5.20	0.42
11:S9:108:ARG:HH21	11:S9:145:SER:HB2	1.84	0.42
11:S9:161:THR:HB	11:S9:162:SER:H	2.09	0.42
34:SR:44:SER:O	34:SR:58:VAL:HG13	4.57	0.42
36:1:1645:U:H2'	36:1:1646:G:H5'	2.00	0.42
36:1:2130:G:C6	36:1:2323:G:O6	2.71	0.42
36:1:2771:U:H2'	36:1:2772:C:O2	2.19	0.42
36:1:2801:A:O2'	36:1:2802:A:H2'	2.19	0.42
36:1:2842:U:C5	36:1:2843:U:C4	3.07	0.42
36:1:2883:U:H2'	36:1:2884:C:H6	1.83	0.42
36:1:2932:U:O2	36:1:2934:A:C8	2.72	0.42
36:1:299:G:H2'	36:1:300:G:O4'	2.19	0.42
36:1:3092:C:H2'	36:1:3092:C:H6	1.68	0.42
36:1:956:U:OP1	86:1:4121:OHX:N1	2.52	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:627:U:H4'	36:1:1399:A:H1'	2.00	0.42
36:1:641:C:OP1	64:N8:21:ARG:HB3	2.19	0.42
36:1:693:A:H2'	36:1:694:C:C6	2.54	0.42
36:1:693:A:H2'	36:1:694:C:H6	1.84	0.42
36:1:732:C:H2'	36:1:733:G:O4'	2.19	0.42
1:2:619:A:HO2'	1:2:1140:G:HO2'	1.63	0.42
1:2:1388:A:C5	1:2:1411:A:C6	3.06	0.42
1:2:1479:A:H8	1:2:1479:A:O5'	2.01	0.42
1:2:1486:G:N7	1:2:1487:A:C8	2.88	0.42
1:2:1512:G:C6	1:2:1513:G:C6	3.07	0.42
1:2:1559:A:H5''	20:C8:135:GLY:HA3	2.01	0.42
1:2:1370:U:O4	86:2:2120:OHX:N5	2.53	0.42
1:2:217:A:H2'	1:2:217:A:OP1	2.19	0.42
1:2:67:A:O3'	1:2:68:A:H3'	2.19	0.42
1:2:726:C:H2'	1:2:727:U:C5	2.54	0.42
36:5:1349:G:H2'	36:5:1350:A:C8	2.54	0.42
36:5:1573:G:C5	36:5:1574:C:H1'	2.54	0.42
36:5:1604:G:H2'	36:5:1605:A:O4'	2.19	0.42
36:5:2413:A:O2'	36:5:2414:G:H5'	2.18	0.42
36:5:2950:G:C5	36:5:2979:U:C4	3.07	0.42
36:5:3261:C:O2'	36:5:3262:U:H5'	2.19	0.42
1:6:1342:C:C2'	1:6:1343:U:H5'	2.48	0.42
1:6:1541:G:C6	1:6:1542:G:N1	2.86	0.42
1:6:162:A:H2'	1:6:163:G:C8	2.55	0.42
1:6:1696:G:H2'	1:6:1698:G:C6	2.54	0.42
1:6:493:U:HO2'	1:6:494:U:H6	1.66	0.42
11:S9:170:GLY:HA3	1:6:512:A:OP2	455.26	0.42
38:8:106:C:H5''	38:8:108:C:OP2	2.18	0.42
13:C1:139:VAL:HG12	13:C1:140:VAL:N	2.27	0.42
14:C2:60:VAL:HG22	14:C2:122:VAL:HG22	2.00	0.42
18:C6:30:LYS:HD3	1:6:1366:U:OP1	425.96	0.42
22:D0:63:LEU:HB3	31:D9:34:TYR:HE2	1.78	0.42
1:2:636:A:H5''	24:D2:31:SER:HB3	2.01	0.42
24:D2:94:LEU:HA	24:D2:95:PRO:HD3	1.86	0.42
25:D3:76:LEU:HD23	25:D3:76:LEU:HA	2.31	0.42
26:D4:19:ALA:CB	26:D4:81:GLU:HG2	2.49	0.42
28:D6:50:VAL:HA	28:D6:53:LEU:HB2	3.16	0.42
28:D6:62:TYR:CG	28:D6:63:ALA:N	2.89	0.42
28:D6:64:LEU:HA	28:D6:65:PRO:HD3	1.76	0.42
15:C3:55:ARG:HD3	29:D7:47:PHE:CD2	2.54	0.42
39:L2:209:HIS:ND1	39:L2:210:PRO:HD2	4.22	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:L3:151:ILE:O	40:L3:155:ALA:HB3	2.27	0.42
40:L3:25:ILE:H	40:L3:25:ILE:HD13	1.85	0.42
40:L3:287:LYS:O	40:L3:290:ASP:HB3	2.19	0.42
40:L3:49:TYR:H	40:L3:79:VAL:HG23	1.84	0.42
41:L4:119:ARG:NH1	41:L4:271:LYS:HB3	3.87	0.42
36:1:338:A:OP1	41:L4:47:ARG:HA	2.18	0.42
42:L5:160:PHE:O	42:L5:180:PHE:HE1	2.01	0.42
42:L5:163:LEU:HD21	42:L5:175:HIS:CD2	4.45	0.42
42:L5:261:THR:C	42:L5:263:GLU:N	3.26	0.42
44:L7:99:PRO:HB3	44:L7:129:LEU:O	2.67	0.42
44:L7:55:TYR:CE2	44:L7:141:TYR:CE2	3.14	0.42
46:L9:77:ASN:N	46:L9:77:ASN:OD1	2.68	0.42
48:M1:132:ASN:HA	48:M1:154:THR:HG21	2.01	0.42
48:M1:6:GLN:HA	48:M1:6:GLN:NE2	2.34	0.42
51:M5:36:ILE:HG13	51:M5:64:VAL:CG2	3.41	0.42
53:M7:32:THR:HG21	53:M7:87:SER:CB	2.50	0.42
54:M8:165:ILE:HD13	54:M8:166:LEU:N	4.64	0.42
55:M9:4:LEU:O	55:M9:7:GLN:N	2.52	0.42
56:N0:154:HIS:CG	56:N0:154:HIS:O	2.71	0.42
56:N0:7:TYR:O	56:N0:28:ARG:HA	2.46	0.42
57:N1:11:THR:HG22	57:N1:55:LYS:HD2	2.01	0.42
57:N1:27:LEU:HA	57:N1:27:LEU:HD22	1.96	0.42
62:N6:100:HIS:CE1	62:N6:102:SER:HB3	2.54	0.42
63:N7:133:LYS:CE	63:N7:135:ARG:HD2	3.62	0.42
36:1:1375:G:O6	64:N8:10:LYS:HE2	2.18	0.42
64:N8:40:HIS:CD2	64:N8:41:HIS:CE1	3.07	0.42
64:N8:88:ASP:O	64:N8:92:LYS:HG2	2.19	0.42
65:N9:51:ALA:O	65:N9:53:ALA:N	2.95	0.42
67:O1:102:LYS:HA	67:O1:102:LYS:HE3	5.25	0.42
67:O1:5:LYS:O	67:O1:6:ASP:HB2	2.19	0.42
74:O8:46:ARG:NH2	74:O8:51:LEU:HB2	2.30	0.42
76:Q0:94:SER:OG	76:Q0:105:PRO:HA	2.19	0.42
39:L2:81:GLY:N	79:Q3:65:ALA:O	2.42	0.42
79:Q3:84:ARG:NH2	79:Q3:88:GLU:OE2	2.51	0.42
2:S0:157:ASP:O	2:S0:158:VAL:C	2.58	0.42
2:S0:88:LYS:O	2:S0:91:ALA:HB3	3.60	0.42
6:S4:188:ASN:HB3	6:S4:191:ARG:HG3	2.59	0.42
8:S6:179:VAL:HA	8:S6:183:ARG:HH11	2.68	0.42
9:S7:21:ALA:O	9:S7:24:PHE:HB2	2.52	0.42
34:SR:126:SER:HG	34:SR:127:ARG:H	1.67	0.42
36:1:1461:A:H2'	36:1:1462:A:H8	1.82	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:1566:A:N6	36:1:1567:U:O2	2.53	0.42
36:1:2202:C:H5''	39:L2:226:SER:HB3	2.00	0.42
36:1:2217:U:H2'	36:1:2218:G:H8	1.84	0.42
36:1:2381:G:O2'	36:1:2382:G:H5'	2.19	0.42
36:1:246:U:H2'	36:1:247:C:C6	2.54	0.42
36:1:2564:G:H2'	36:1:2565:U:O4'	2.19	0.42
36:1:2592:G:O2'	36:1:2593:A:O4'	2.37	0.42
36:1:2861:U:H2'	36:1:2862:U:O4'	2.19	0.42
36:1:3121:U:C1'	36:1:3122:A:H5''	2.49	0.42
36:1:3317:U:H5'	36:1:3318:G:C2	2.54	0.42
36:1:781:G:O6	86:1:3933:OHX:N5	2.52	0.42
86:1:4023:OHX:N6	86:1:4143:OHX:N5	2.67	0.42
36:1:1343:A:N7	86:1:4136:OHX:N2	2.67	0.42
36:1:571:U:H2'	36:1:572:A:O4'	2.20	0.42
36:1:595:G:H2'	36:1:596:C:H6	1.83	0.42
36:1:998:A:O2'	36:1:999:G:H5'	2.19	0.42
1:2:1241:G:C6	1:2:1242:A:C6	3.07	0.42
1:2:1277:G:O3'	5:S3:183:GLY:HA3	2.20	0.42
1:2:1597:A:H2'	1:2:1598:U:O4'	2.19	0.42
1:2:1585:U:N3	1:2:1611:A:H2	2.08	0.42
1:2:1653:C:C2	1:2:1748:G:N2	2.87	0.42
1:2:1795:U:H5'	28:D6:79:ILE:CD1	2.49	0.42
1:2:296:U:H2'	1:2:297:U:C6	2.54	0.42
1:2:525:A:H2'	1:2:526:A:C8	2.55	0.42
36:5:1742:U:H2'	36:5:1743:G:C8	2.54	0.42
40:L3:247:ARG:NH2	36:5:2341:A:P	219.70	0.42
36:5:2931:C:H2'	36:5:2932:U:O4'	2.20	0.42
86:5:4004:OHX:N6	86:5:4195:OHX:N5	2.68	0.42
36:5:584:G:O6	86:5:4014:OHX:N1	2.53	0.42
1:6:1799:U:O3'	1:6:1800:A:H2'	2.19	0.42
1:6:271:A:H5'	1:6:272:U:OP2	2.20	0.42
1:6:509:G:H2'	1:6:510:G:C1'	2.48	0.42
1:6:874:C:O2'	1:6:875:G:H5'	2.19	0.42
38:8:48:A:C2	38:8:51:G:N1	2.87	0.42
13:C1:5:LEU:HD22	13:C1:5:LEU:H	4.76	0.42
15:C3:3:ARG:HD3	15:C3:3:ARG:HA	1.73	0.42
3:S1:65:VAL:O	16:C4:34:SER:HA	2.19	0.42
17:C5:68:PRO:HG2	17:C5:71:GLU:CD	2.76	0.42
18:C6:10:PHE:O	18:C6:87:LYS:NZ	2.43	0.42
19:C7:23:LYS:HG2	34:SR:198:ASN:HD21	2.42	0.42
20:C8:114:GLU:HA	20:C8:117:LYS:HB2	3.14	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:C9:136:ALA:O	21:C9:139:THR:OG1	2.28	0.42
21:C9:136:ALA:O	21:C9:140:LEU:HB2	3.91	0.42
22:D0:58:LEU:HD23	1:6:1516:A:H8	444.85	0.42
1:2:1198:G:H4'	22:D0:72:ASN:O	2.19	0.42
22:D0:80:GLU:OE1	31:D9:44:ARG:NH1	2.52	0.42
4:S2:58:LEU:HA	23:D1:12:TYR:HE1	1.84	0.42
26:D4:60:PHE:O	1:6:523:G:H5'	413.92	0.42
39:L2:126:LEU:HD13	39:L2:150:LEU:HD21	2.02	0.42
39:L2:200:ARG:NH1	36:5:2147:A:OP2	212.35	0.42
40:L3:37:ARG:HG2	40:L3:186:GLY:O	2.18	0.42
40:L3:287:LYS:HB2	40:L3:290:ASP:HB2	2.01	0.42
42:L5:95:TRP:CZ2	42:L5:161:GLY:HA2	2.55	0.42
42:L5:208:MET:HG2	42:L5:223:PHE:CE2	3.15	0.42
42:L5:260:PHE:HD1	42:L5:264:GLN:NE2	2.16	0.42
47:M0:31:ILE:O	47:M0:32:ARG:HD3	2.20	0.42
47:M0:12:GLN:HA	47:M0:59:GLN:HE21	2.23	0.42
48:M1:174:LYS:HE3	36:5:1016:C:N4	357.93	0.42
50:M4:85:TRP:O	50:M4:90:VAL:HG12	2.19	0.42
52:M6:97:ALA:O	52:M6:100:GLU:HB2	2.18	0.42
53:M7:67:ILE:HD12	53:M7:67:ILE:HA	1.57	0.42
55:M9:164:LEU:O	55:M9:168:ALA:N	3.82	0.42
55:M9:186:LYS:C	55:M9:188:ASP:H	2.71	0.42
36:1:1690:C:P	55:M9:64:ARG:HH12	2.43	0.42
56:N0:155:ARG:HH21	56:N0:155:ARG:CG	2.84	0.42
57:N1:91:LEU:HD12	57:N1:96:ILE:HD11	2.00	0.42
59:N3:17:LEU:HD23	59:N3:17:LEU:HA	2.10	0.42
61:N5:34:LEU:HD22	61:N5:35:PRO:HD2	2.01	0.42
62:N6:39:LEU:HD12	62:N6:43:TYR:CE2	4.19	0.42
54:M8:94:PHE:CE2	64:N8:119:PRO:HD3	2.95	0.42
66:O0:20:SER:O	66:O0:20:SER:OG	2.34	0.42
69:O3:48:ARG:HG2	69:O3:103:TYR:O	3.91	0.42
69:O3:89:LEU:HA	69:O3:89:LEU:HD23	1.69	0.42
70:O4:11:ASN:HA	70:O4:12:PRO:HD3	1.78	0.42
70:O4:77:GLY:C	70:O4:79:SER:H	2.22	0.42
72:O6:66:GLU:HA	72:O6:69:ALA:HB3	2.42	0.42
73:O7:24:ARG:NH1	36:5:361:A:OP1	120.37	0.42
74:O8:56:ILE:HG21	74:O8:61:LYS:HB2	2.67	0.42
74:O8:61:LYS:HE3	74:O8:61:LYS:HB3	4.41	0.42
2:S0:103:THR:HA	2:S0:104:PRO:HD2	2.63	0.42
2:S0:26:ALA:HB1	2:S0:29:VAL:CG1	2.48	0.42
4:S2:175:GLY:HA3	11:S9:53:ARG:NH2	2.34	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:S2:178:ILE:HD12	4:S2:178:ILE:H	4.50	0.42
4:S2:58:LEU:HA	4:S2:58:LEU:HD23	1.66	0.42
4:S2:96:THR:OG1	4:S2:97:ARG:N	3.62	0.42
6:S4:11:ARG:HB2	6:S4:28:ALA:N	2.34	0.42
6:S4:123:LEU:HD12	6:S4:161:LYS:HA	2.01	0.42
6:S4:66:MET:HG3	1:6:454:U:C2	376.57	0.42
7:S5:210:ALA:O	7:S5:213:LYS:N	2.52	0.42
7:S5:92:ARG:HG2	7:S5:92:ARG:NH1	3.29	0.42
8:S6:148:SER:OG	8:S6:150:GLU:HB2	2.19	0.42
9:S7:129:LEU:O	9:S7:131:PHE:N	4.24	0.42
9:S7:71:HIS:CE1	9:S7:131:PHE:HE1	2.38	0.42
10:S8:67:TRP:CD1	10:S8:70:GLU:HB2	2.53	0.42
34:SR:224:ASN:HB2	34:SR:231:MET:SD	2.59	0.42
36:1:13:A:H5''	36:1:13:A:C8	2.47	0.42
36:1:1668:G:N1	36:1:1669:C:C2	2.88	0.42
36:1:2664:C:H2'	36:1:2665:U:C6	2.55	0.42
36:1:26:A:C4	36:1:330:G:C8	3.07	0.42
36:1:3077:A:N6	36:1:3080:G:C5	2.87	0.42
36:1:197:G:N2	36:1:372:A:C8	2.87	0.42
36:1:960:U:O2'	36:1:961:C:H5'	2.19	0.42
1:2:1073:G:H2'	1:2:1074:G:H5''	2.02	0.42
1:2:1642:G:O3'	77:Q1:9:ARG:NH2	2.52	0.42
1:2:455:C:H3'	1:2:456:A:H8	1.84	0.42
1:2:629:U:C4	1:2:630:A:N7	2.88	0.42
38:4:11:C:C4	38:4:12:A:N7	2.87	0.42
36:5:1039:U:H2'	36:5:1040:A:C8	2.54	0.42
36:5:1258:U:O2	36:5:1260:A:H8	2.02	0.42
64:N8:9:ARG:NH2	36:5:1431:G:N7	148.01	0.42
36:5:1549:U:H6	36:5:1549:U:OP2	2.02	0.42
36:5:1769:G:C2	36:5:1770:G:C8	3.07	0.42
36:5:1792:C:H5''	36:5:1793:C:OP2	2.20	0.42
36:5:179:C:H2'	36:5:180:C:C6	2.54	0.42
36:5:2271:A:N7	36:5:2272:G:C6	2.87	0.42
36:5:2353:G:C6	36:5:2354:C:C4	3.08	0.42
36:5:1449:A:C2	36:5:2356:A:C4	3.08	0.42
36:5:41:G:H4'	36:5:2410:U:H2'	2.02	0.42
36:5:2946:A:H5''	36:5:2947:G:H5'	2.01	0.42
36:5:2961:G:C6	36:5:2962:U:C4	3.07	0.42
36:5:3341:U:H5''	36:5:3342:A:OP2	2.19	0.42
36:5:3365:U:O2'	36:5:3366:G:H5'	2.19	0.42
36:5:372:A:C6	36:5:373:A:C6	3.08	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:M3:36:ARG:CZ	36:5:687:U:H5	74.95	0.42
36:5:897:U:H2'	36:5:898:U:C6	2.55	0.42
64:N8:28:HIS:CD2	36:5:936:A:OP1	161.14	0.42
1:6:112:A:C6	1:6:113:U:O4	2.72	0.42
1:6:1138:A:H2'	1:6:1139:A:C8	2.54	0.42
1:6:1142:A:N6	1:6:1143:A:C6	2.87	0.42
21:C9:88:VAL:CG2	1:6:1172:G:H21	356.50	0.42
1:6:1499:G:H2'	1:6:1500:C:C6	2.54	0.42
1:6:1796:C:H5'	1:6:1797:A:N7	2.34	0.42
1:6:192:U:H1'	1:6:193:U:C5	2.55	0.42
1:6:475:A:H2'	1:6:476:U:O4'	2.19	0.42
1:6:542:A:H1'	1:6:543:C:OP1	2.19	0.42
1:6:543:C:O2	1:6:543:C:O4'	2.38	0.42
1:6:794:U:H4'	1:6:795:U:OP2	2.20	0.42
1:6:825:U:O2'	1:6:826:U:H6	2.02	0.42
1:6:889:U:H4'	1:6:989:U:OP1	2.18	0.42
12:C0:6:GLU:HA	12:C0:9:ASN:HB2	3.67	0.42
17:C5:60:LEU:HA	17:C5:60:LEU:HD23	2.08	0.42
17:C5:44:ARG:NH2	17:C5:82:ASN:O	3.51	0.42
18:C6:44:LEU:O	18:C6:47:LYS:HB2	2.20	0.42
19:C7:79:GLU:HA	19:C7:82:ASP:OD2	2.20	0.42
20:C8:108:LYS:HD3	20:C8:108:LYS:HA	3.45	0.42
20:C8:17:LEU:O	20:C8:20:THR:N	2.78	0.42
21:C9:100:ILE:HD13	21:C9:100:ILE:HA	1.85	0.42
21:C9:105:LEU:HD23	21:C9:105:LEU:HA	1.70	0.42
1:2:1520:U:H5''	21:C9:75:LYS:NZ	2.34	0.42
25:D3:23:ARG:HG3	25:D3:23:ARG:HH11	2.15	0.42
26:D4:44:LEU:HA	26:D4:47:VAL:HG13	4.91	0.42
28:D6:12:LYS:HB3	28:D6:12:LYS:NZ	4.10	0.42
30:D8:32:PHE:CZ	30:D8:38:ARG:HD2	2.54	0.42
39:L2:132:ASN:OD1	36:5:2178:A:H5''	215.81	0.42
40:L3:41:VAL:HG23	40:L3:186:GLY:H	1.85	0.42
40:L3:232:ARG:HG2	40:L3:233:TRP:CD1	2.91	0.42
40:L3:25:ILE:HD13	40:L3:25:ILE:N	2.34	0.42
40:L3:322:ILE:HA	40:L3:322:ILE:HD12	2.97	0.42
40:L3:50:LYS:HE2	40:L3:328:ILE:HG22	4.15	0.42
37:3:120:C:H42	42:L5:262:LYS:HD3	1.83	0.42
44:L7:123:THR:O	44:L7:126:LEU:HB2	2.19	0.42
44:L7:40:LYS:HE2	44:L7:170:GLU:OE1	4.07	0.42
44:L7:132:PRO:HA	44:L7:229:PHE:CG	2.55	0.42
44:L7:240:VAL:C	44:L7:242:SER:H	2.22	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:M1:91:LEU:HD12	48:M1:163:PHE:CE2	2.54	0.42
51:M5:21:PHE:CD2	51:M5:22:LEU:HD13	3.18	0.42
52:M6:130:LYS:HG3	52:M6:131:PRO:CD	2.79	0.42
53:M7:119:VAL:HA	53:M7:145:HIS:O	2.60	0.42
53:M7:33:ALA:O	53:M7:35:ALA:N	3.37	0.42
53:M7:3:ARG:HH21	53:M7:3:ARG:HG2	4.37	0.42
56:N0:139:TYR:CE2	56:N0:140:VAL:HG23	2.53	0.42
57:N1:111:ALA:O	57:N1:115:LYS:HG3	2.19	0.42
57:N1:92:ARG:C	57:N1:94:GLU:H	2.23	0.42
59:N3:6:ALA:HB2	59:N3:126:TRP:CH2	2.55	0.42
38:4:23:U:H1'	62:N6:17:LYS:HG2	2.02	0.42
62:N6:42:GLN:O	62:N6:125:LYS:HG3	2.53	0.42
62:N6:82:VAL:O	62:N6:84:LYS:N	2.98	0.42
63:N7:10:VAL:HB	63:N7:83:THR:HG21	2.01	0.42
66:O0:28:LYS:HB2	36:5:1730:G:C5	240.36	0.42
70:O4:78:GLY:O	70:O4:80:ARG:N	4.75	0.42
74:O8:17:ARG:HB2	74:O8:20:VAL:CG2	2.48	0.42
74:O8:41:THR:HG1	74:O8:43:PHE:HE2	3.18	0.42
46:L9:93:VAL:HG22	76:Q0:82:LEU:HD13	2.02	0.42
78:Q2:78:LYS:HG2	78:Q2:78:LYS:O	3.36	0.42
79:Q3:29:LEU:HA	79:Q3:32:GLN:HB2	2.00	0.42
3:S1:134:VAL:HG12	3:S1:218:LEU:HB2	6.77	0.42
7:S5:42:LEU:HB2	7:S5:47:SER:HA	2.00	0.42
9:S7:108:GLN:HB2	9:S7:108:GLN:HE21	3.39	0.42
9:S7:162:ILE:HB	9:S7:169:PHE:HE2	1.85	0.42
9:S7:19:GLN:H	9:S7:19:GLN:HG2	1.61	0.42
10:S8:38:ILE:HD11	10:S8:78:ILE:HB	2.01	0.42
11:S9:110:GLN:HA	11:S9:129:ILE:HD13	2.80	0.42
34:SR:115:ILE:HB	34:SR:122:ILE:HG12	4.18	0.42
34:SR:37:SER:HB3	34:SR:39:ASP:OD1	2.90	0.42
34:SR:69:GLN:O	34:SR:83:ALA:HB3	2.19	0.42
36:1:1054:A:H5''	36:1:2637:A:N6	2.34	0.42
36:1:1204:A:H2'	36:1:1205:A:H5'	2.02	0.42
36:1:1784:G:H2'	36:1:1785:U:H6	1.84	0.42
36:1:2143:A:H3'	36:1:2143:A:C8	2.54	0.42
36:1:2267:C:H2'	36:1:2268:U:O4'	2.20	0.42
36:1:2656:A:C4	36:1:2658:G:N7	2.88	0.42
36:1:2854:U:H2'	36:1:2855:U:H6	1.84	0.42
36:1:396:A:O2'	36:1:399:A:OP1	2.33	0.42
1:2:144:U:C2	1:2:145:A:C8	3.08	0.42
1:2:1476:C:H2'	1:2:1477:G:H8	1.83	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:19:A:C2	1:2:20:G:C4	3.08	0.42
1:2:275:C:H2'	1:2:276:C:C5	2.54	0.42
1:2:641:G:N2	1:2:693:U:O2	2.49	0.42
1:2:740:A:O5'	1:2:740:A:H8	2.03	0.42
36:5:1119:C:H2'	36:5:1120:A:H8	1.85	0.42
36:5:1424:C:H2'	36:5:1425:U:O4'	2.20	0.42
55:M9:5:ARG:NH2	36:5:1471:U:OP1	123.03	0.42
36:5:1743:G:H2'	36:5:1744:G:H8	1.83	0.42
36:5:1817:G:HO2'	36:5:1818:U:H6	1.60	0.42
36:5:1481:A:O2'	36:5:1858:A:N3	2.48	0.42
36:5:2320:A:OP2	86:5:4069:OHX:N5	2.53	0.42
36:5:48:A:C8	36:5:50:U:C2	3.07	0.42
1:6:1066:C:H2'	1:6:1067:C:H6	1.84	0.42
1:6:1079:U:C4	1:6:1080:U:C4	3.07	0.42
1:6:119:A:H2'	1:6:120:U:O4'	2.20	0.42
1:6:1283:U:H2'	1:6:1284:C:C5	2.54	0.42
1:6:1512:G:C4	1:6:1513:G:C8	3.08	0.42
1:6:1660:A:H2'	1:6:1661:U:C6	2.54	0.42
1:6:23:G:C6	1:6:24:U:N3	2.88	0.42
1:6:542:A:OP2	1:6:542:A:H2'	2.20	0.42
1:6:563:U:C4	1:6:564:G:C6	3.08	0.42
1:6:607:G:OP2	1:6:613:G:N1	2.51	0.42
1:6:809:A:C6	1:6:810:G:O6	2.73	0.42
1:6:875:G:H4'	1:6:936:G:O2'	2.19	0.42
38:8:56:G:H2'	38:8:57:C:O4'	2.20	0.42
15:C3:99:ARG:O	15:C3:102:LEU:N	2.47	0.42
16:C4:83:ILE:O	16:C4:118:VAL:HG22	2.42	0.42
16:C4:66:ASP:O	16:C4:69:ALA:HB3	2.34	0.42
20:C8:116:LEU:HD13	20:C8:116:LEU:HA	1.80	0.42
20:C8:27:LYS:HB3	20:C8:27:LYS:HE2	4.51	0.42
22:D0:100:VAL:O	22:D0:104:THR:HG23	2.20	0.42
22:D0:15:GLN:O	22:D0:16:GLN:HB2	4.18	0.42
25:D3:141:GLU:OE1	25:D3:144:ARG:NH1	13.09	0.42
26:D4:36:SER:HB3	26:D4:39:GLU:HG3	3.97	0.42
30:D8:18:ARG:HA	30:D8:26:THR:HA	2.02	0.42
36:1:2242:A:H5'	39:L2:243:THR:HG23	2.01	0.42
36:1:2988:C:O4'	40:L3:260:VAL:HB	2.19	0.42
36:1:3139:A:OP1	40:L3:274:SER:HB2	2.19	0.42
40:L3:219:ALA:HB3	40:L3:329:PRO:HB2	2.01	0.42
41:L4:346:LYS:HD2	41:L4:347:THR:N	5.90	0.42
41:L4:62:ALA:HB3	41:L4:90:PHE:CE2	2.96	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:L5:179:ARG:HA	42:L5:179:ARG:HD3	2.16	0.42
42:L5:210:GLU:O	42:L5:214:ASP:HB2	2.52	0.42
43:L6:38:THR:HA	43:L6:90:LYS:HG3	2.18	0.42
47:M0:169:LYS:HD2	47:M0:169:LYS:H	2.28	0.42
36:1:2856:G:N7	47:M0:7:ARG:NH2	2.68	0.42
48:M1:37:LEU:HD23	48:M1:37:LEU:HA	1.83	0.42
49:M3:187:ALA:CA	49:M3:190:LYS:HB3	2.45	0.42
50:M4:99:TRP:C	50:M4:99:TRP:CD1	3.04	0.42
53:M7:132:ALA:O	53:M7:133:HIS:HB2	2.20	0.42
55:M9:59:SER:HB2	55:M9:61:SER:HB3	2.02	0.42
55:M9:5:ARG:HG3	55:M9:6:THR:N	3.00	0.42
58:N2:55:THR:HG22	58:N2:57:THR:HG22	7.03	0.42
58:N2:27:VAL:HG11	58:N2:92:TRP:CZ3	5.33	0.42
60:N4:58:HIS:C	60:N4:60:LYS:H	2.70	0.42
61:N5:103:TYR:O	61:N5:105:VAL:HG23	2.33	0.42
66:O0:101:LEU:N	66:O0:101:LEU:HD22	2.94	0.42
70:O4:71:THR:HG23	70:O4:78:GLY:H	1.84	0.42
71:O5:31:LEU:HD12	71:O5:31:LEU:HA	1.76	0.42
71:O5:32:LYS:O	71:O5:36:LEU:HD23	3.19	0.42
38:4:58:G:O6	73:O7:63:ARG:NH1	2.52	0.42
73:O7:67:LEU:HD23	73:O7:70:VAL:HG21	2.01	0.42
73:O7:5:THR:HA	73:O7:8:PHE:HD2	1.85	0.42
74:O8:54:LEU:HG	74:O8:56:ILE:HD11	2.69	0.42
74:O8:43:PHE:CZ	74:O8:66:ILE:HG12	3.29	0.42
78:Q2:12:CYS:CB	78:Q2:17:CYS:HB3	2.87	0.42
2:S0:57:LEU:HD21	2:S0:177:LEU:HA	2.02	0.42
3:S1:229:MET:HA	3:S1:232:HIS:ND1	2.35	0.42
3:S1:61:LEU:O	3:S1:62:LYS:NZ	2.27	0.42
2:S0:119:ARG:HH21	4:S2:240:LEU:HD23	1.85	0.42
8:S6:48:TYR:OH	8:S6:119:GLN:O	3.13	0.42
8:S6:39:GLU:CD	8:S6:46:LYS:HG3	2.84	0.42
9:S7:20:VAL:HG13	9:S7:81:LEU:HD21	4.59	0.42
10:S8:42:ARG:HB3	10:S8:59:ARG:HB2	2.04	0.42
11:S9:141:VAL:HG12	11:S9:143:ILE:H	1.84	0.42
35:SM:61:ILE:HD12	35:SM:62:ARG:N	2.34	0.42
34:SR:202:LEU:HA	34:SR:212:ALA:O	2.20	0.42
34:SR:295:SER:HB2	34:SR:300:THR:O	2.19	0.42
36:1:1064:A:N6	36:1:1096:U:N3	2.63	0.42
36:1:1719:G:H2'	36:1:1720:U:O4'	2.19	0.42
36:1:1844:C:H42	36:1:1851:G:H1	1.66	0.42
36:1:3119:U:OP2	86:1:3884:OHX:N4	2.53	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:3193:C:H2'	36:1:3194:C:H6	1.85	0.42
36:1:3284:G:H2'	36:1:3285:C:C6	2.54	0.42
36:1:3326:G:C4	36:1:3327:G:C8	3.08	0.42
36:1:3387:U:H2'	36:1:3388:C:C6	2.54	0.42
36:1:547:G:H4'	36:1:548:G:OP2	2.14	0.42
36:1:849:C:H2'	36:1:850:U:H6	1.84	0.42
1:2:1339:C:O2'	1:2:1340:U:OP1	2.37	0.42
1:2:1368:G:C4	1:2:1369:U:C5	3.07	0.42
1:2:1431:C:H3'	1:2:1432:U:H5'	2.02	0.42
1:2:1575:G:H2'	1:2:1576:A:H8	1.84	0.42
1:2:226:A:C2'	1:2:227:U:H5'	2.50	0.42
1:2:475:A:OP2	11:S9:126:ARG:NH1	2.53	0.42
1:2:850:A:C2	1:2:851:U:C2	3.07	0.42
1:2:852:C:OP1	55:M9:172:ARG:HD3	2.19	0.42
37:3:106:U:H6	37:3:106:U:OP2	2.03	0.42
36:5:1123:U:H2'	36:5:1124:U:H5'	2.01	0.42
36:5:1313:G:H2'	36:5:1314:C:H6	1.84	0.42
36:5:1613:A:H2'	36:5:1614:C:H6	1.85	0.42
36:5:1895:A:O2'	36:5:3053:G:H4'	2.20	0.42
36:5:2514:U:H6	36:5:2514:U:OP1	2.02	0.42
36:5:3188:G:C2	36:5:3205:G:N1	2.87	0.42
36:5:3234:A:OP2	36:5:3234:A:C8	2.71	0.42
36:5:3289:G:H4'	36:5:3290:G:OP1	2.19	0.42
88:5:4248:BLS:H102	88:5:4248:BLS:HN6	1.85	0.42
36:5:574:U:H2'	36:5:575:G:O4'	2.19	0.42
51:M5:198:SER:OG	36:5:82:C:OP1	107.97	0.42
1:6:1071:U:H2'	1:6:1072:C:C6	2.55	0.42
1:6:1215:C:H2'	1:6:1216:C:C6	2.54	0.42
1:6:1325:A:H2'	1:6:1326:A:C8	2.55	0.42
1:6:1759:C:H2'	1:6:1760:G:O4'	2.20	0.42
1:6:228:G:H22	1:6:237:C:H42	1.67	0.42
1:6:329:G:H2'	1:6:330:G:H8	1.84	0.42
11:S9:5:PRO:HG3	1:6:380:U:C2	368.41	0.42
73:O7:72:ARG:NH1	38:8:94:C:H3'	50.55	0.42
12:C0:59:PHE:CZ	12:C0:62:GLN:HA	2.55	0.42
13:C1:22:ASN:HA	13:C1:23:PRO:HD3	1.72	0.42
18:C6:5:PRO:HB2	18:C6:96:TYR:CE2	3.22	0.42
20:C8:101:LEU:HD23	20:C8:101:LEU:HA	2.29	0.42
24:D2:78:ARG:HG2	24:D2:78:ARG:H	1.65	0.42
25:D3:127:VAL:HG23	25:D3:128:SER:N	2.35	0.42
25:D3:24:TRP:CH2	25:D3:34:LEU:HD21	3.32	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:1105:C:H41	25:D3:4:GLY:HA3	1.84	0.42
1:2:542:A:C2	32:E0:28:LYS:HD2	2.55	0.42
1:2:545:A:OP1	32:E0:31:LYS:HE3	2.18	0.42
41:L4:144:LYS:HA	86:L4:404:OHX:N2	2.34	0.42
41:L4:161:LYS:O	41:L4:162:THR:C	2.58	0.42
41:L4:22:LEU:HA	41:L4:23:PRO:HD3	1.71	0.42
41:L4:258:LEU:HD12	41:L4:258:LEU:HA	1.87	0.42
36:1:2663:G:H4'	42:L5:152:ARG:NH1	2.35	0.42
42:L5:268:GLU:O	42:L5:268:GLU:HG3	2.19	0.42
45:L8:156:ASP:O	45:L8:157:VAL:HB	2.20	0.42
45:L8:158:ASP:HB3	45:L8:159:PRO:HD3	2.01	0.42
45:L8:91:PHE:CE1	45:L8:185:ARG:HD2	2.53	0.42
46:L9:13:PRO:HG2	46:L9:16:VAL:CG1	2.73	0.42
47:M0:22:TYR:HB3	36:5:2647:A:H4'	265.64	0.42
50:M4:50:LYS:HD3	50:M4:85:TRP:HD1	1.83	0.42
51:M5:15:GLN:HA	51:M5:20:ARG:HD2	2.20	0.42
52:M6:147:TRP:HE1	52:M6:149:TYR:HB2	1.84	0.42
53:M7:95:LEU:HD23	53:M7:95:LEU:HA	1.65	0.42
57:N1:154:VAL:HA	57:N1:155:PRO:HD3	1.90	0.42
57:N1:34:TYR:CE1	57:N1:98:HIS:NE2	3.46	0.42
62:N6:32:SER:HA	62:N6:50:ILE:H	2.25	0.42
63:N7:93:LYS:HD3	63:N7:93:LYS:HA	1.83	0.42
64:N8:13:GLY:O	64:N8:14:HIS:ND1	2.45	0.42
71:O5:78:LYS:HA	71:O5:81:ARG:HE	4.44	0.42
72:O6:93:ILE:O	72:O6:96:ALA:HB3	2.90	0.42
73:O7:52:LYS:HE2	36:5:353:G:O6	114.52	0.42
74:O8:26:LYS:HB3	74:O8:42:LYS:HB2	2.02	0.42
38:4:45:C:OP1	75:O9:12:LYS:NZ	2.52	0.42
77:Q1:16:LYS:HE2	77:Q1:16:LYS:HB3	4.70	0.42
78:Q2:3:ASN:HA	78:Q2:92:GLU:O	2.20	0.42
79:Q3:8:VAL:HG22	36:5:1927:G:OP1	245.24	0.42
2:S0:146:LEU:HA	2:S0:160:ILE:O	3.31	0.42
2:S0:92:HIS:CD2	2:S0:195:TRP:HH2	3.41	0.42
2:S0:21:ASN:HB3	2:S0:24:LEU:HD13	2.01	0.42
2:S0:42:PRO:CD	19:C7:104:ASN:HD21	7.21	0.42
3:S1:101:HIS:HA	3:S1:217:LEU:HD22	2.02	0.42
4:S2:147:ASN:HB3	23:D1:4:ASP:HA	2.01	0.42
4:S2:166:THR:O	4:S2:166:THR:OG1	2.78	0.42
4:S2:44:LEU:HA	4:S2:44:LEU:HD23	2.24	0.42
5:S3:113:LEU:HD13	5:S3:113:LEU:HA	4.07	0.42
1:2:1514:U:O2'	5:S3:5:ILE:O	2.38	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:S5:131:GLN:O	7:S5:131:GLN:HG3	2.18	0.42
7:S5:162:VAL:HG23	7:S5:166:ARG:HB3	3.07	0.42
8:S6:139:ASN:O	8:S6:142:ARG:HB2	2.20	0.42
9:S7:89:HIS:CD2	9:S7:165:LYS:HG2	2.54	0.42
10:S8:79:ALA:N	10:S8:103:GLN:O	2.50	0.42
10:S8:43:ILE:HA	10:S8:57:ALA:HA	2.01	0.42
11:S9:109:LEU:HD13	11:S9:129:ILE:CD1	2.50	0.42
11:S9:174:ARG:HA	11:S9:174:ARG:HE	1.85	0.42
34:SR:107:LYS:HB2	34:SR:128:ASP:CB	3.61	0.42
34:SR:145:LEU:HG	34:SR:145:LEU:H	2.38	0.42
34:SR:178:VAL:HB	34:SR:192:PHE:HB2	2.19	0.42
36:1:1054:A:H5''	36:1:2637:A:H61	1.85	0.42
36:1:1233:G:N1	36:1:1234:G:O6	2.52	0.42
36:1:1388:U:O2'	68:O2:99:ASN:HB2	2.19	0.42
36:1:2218:G:H2'	36:1:2219:A:C8	2.55	0.42
36:1:2257:C:H2'	36:1:2258:U:C6	2.55	0.42
36:1:2794:G:N7	86:1:3927:OHX:N2	2.67	0.42
36:1:2794:G:C5	36:1:2795:U:C4	3.06	0.42
36:1:2894:C:OP1	46:L9:168:ARG:NH2	2.52	0.42
36:1:3039:C:OP1	40:L3:65:SER:OG	2.29	0.42
36:1:3321:C:H2'	36:1:3322:A:O4'	2.19	0.42
36:1:973:A:H2'	36:1:974:G:O4'	2.19	0.42
1:2:1182:U:O2	1:2:1182:U:H2'	2.18	0.42
1:2:1297:G:O2'	1:2:1299:G:N7	2.48	0.42
1:2:147:A:C6	1:2:148:A:C2	3.08	0.42
1:2:1659:A:C2	1:2:1660:A:C5	3.07	0.42
1:2:208:U:H2'	1:2:209:U:C6	2.54	0.42
1:2:750:U:H2'	1:2:751:G:O4'	2.20	0.42
1:2:7:G:N7	4:S2:205:ARG:NH1	2.68	0.42
1:2:817:A:C6	1:2:818:C:N4	2.88	0.42
36:5:1202:A:N6	36:5:1301:A:N9	2.67	0.42
43:L6:5:LYS:NZ	36:5:1423:C:H1'	140.79	0.42
36:5:1673:G:C5	36:5:1775:G:C6	3.07	0.42
36:5:1816:A:C2'	36:5:1817:G:H5''	2.49	0.42
36:5:2808:A:O2'	86:5:3904:OHX:N6	2.53	0.42
40:L3:30:LYS:HD2	36:5:3138:U:OP2	240.70	0.42
36:5:3203:U:O2'	36:5:3204:C:H5'	2.19	0.42
36:5:528:U:H2'	36:5:529:A:H8	1.82	0.42
36:5:651:G:C6	36:5:652:G:C6	3.08	0.42
49:M3:35:ARG:NH1	36:5:685:G:OP1	81.52	0.42
36:5:846:A:OP1	36:5:846:A:H8	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:914:A:O2'	36:5:2146:C:H4'	2.19	0.42
1:6:1139:A:C5	1:6:1140:G:C8	3.07	0.42
1:6:1360:A:O3'	1:6:1361:U:H4'	2.19	0.42
1:6:140:A:OP2	1:6:140:A:H4'	2.20	0.42
1:6:694:U:H6	1:6:694:U:OP2	2.02	0.42
86:7:220:OHX:N3	86:7:228:OHX:N5	2.68	0.42
56:N0:39:SER:OG	37:7:98:C:OP1	284.24	0.42
13:C1:71:LEU:HD22	13:C1:88:ARG:NH1	2.35	0.42
14:C2:27:ALA:O	14:C2:31:VAL:HG23	2.20	0.42
17:C5:86:VAL:HB	17:C5:87:PRO:HD2	2.36	0.42
18:C6:29:ILE:HG23	18:C6:65:ILE:HG21	2.02	0.42
18:C6:40:GLU:OE2	18:C6:45:ARG:NH2	4.78	0.42
19:C7:29:GLN:HB2	19:C7:29:GLN:HE21	2.17	0.42
24:D2:73:GLY:HA3	24:D2:128:PHE:CE1	3.49	0.42
4:S2:230:TRP:CD2	24:D2:68:ARG:HD3	2.55	0.42
28:D6:36:ILE:HD12	28:D6:36:ILE:N	4.98	0.42
28:D6:44:ILE:HD12	28:D6:45:VAL:HG13	2.00	0.42
30:D8:14:LYS:O	30:D8:28:VAL:HG13	2.19	0.42
39:L2:42:ARG:HA	39:L2:88:ILE:O	2.43	0.42
40:L3:252:ILE:HG23	40:L3:260:VAL:HG22	3.17	0.42
41:L4:259:ASP:OD1	41:L4:259:ASP:N	2.53	0.42
41:L4:281:ILE:HG13	54:M8:29:LEU:HD21	2.02	0.42
42:L5:150:LEU:HD23	42:L5:150:LEU:HA	1.85	0.42
42:L5:219:PHE:HE1	42:L5:227:LEU:HD11	2.25	0.42
36:1:2703:A:H62	42:L5:23:ARG:HG2	1.85	0.42
45:L8:57:ARG:O	45:L8:61:GLN:HG3	3.88	0.42
48:M1:86:VAL:HG22	48:M1:111:ASP:O	2.24	0.42
49:M3:172:LEU:HA	49:M3:172:LEU:HD23	1.81	0.42
52:M6:121:PRO:C	52:M6:123:ALA:H	2.44	0.42
52:M6:36:VAL:HB	52:M6:108:ILE:HB	4.66	0.42
53:M7:30:ARG:HD2	53:M7:63:PHE:CE2	2.72	0.42
55:M9:164:LEU:HD22	55:M9:164:LEU:HA	2.15	0.42
55:M9:176:ARG:HA	55:M9:176:ARG:HD3	1.68	0.42
9:S7:39:ARG:CZ	55:M9:189:ALA:HB3	5.43	0.42
57:N1:126:VAL:HB	57:N1:128:LEU:HG	2.89	0.42
56:N0:25:PHE:HA	57:N1:149:GLN:O	2.62	0.42
61:N5:24:LEU:HD22	61:N5:25:LYS:H	3.34	0.42
64:N8:74:ASN:HA	64:N8:113:LEU:O	2.37	0.42
36:1:1456:A:H5'	67:O1:26:LYS:HG2	2.01	0.42
68:O2:24:ARG:HG2	68:O2:25:TYR:CE1	2.55	0.42
68:O2:8:LYS:HB2	68:O2:8:LYS:HE3	1.82	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
70:O4:6:THR:HG21	36:5:1487:G:C1'	140.84	0.42
49:M3:94:GLY:HA3	71:O5:116:TYR:CE1	3.61	0.42
74:O8:12:LEU:C	74:O8:14:LEU:H	3.12	0.42
39:L2:80:GLU:HG3	79:Q3:66:GLY:HA2	2.00	0.42
2:S0:11:PRO:O	2:S0:15:GLN:HG3	2.20	0.42
3:S1:117:TRP:HD1	3:S1:152:ARG:O	2.02	0.42
3:S1:135:LEU:HD11	3:S1:176:VAL:HG11	2.04	0.42
6:S4:73:ASP:OD2	6:S4:122:LYS:NZ	2.37	0.42
6:S4:92:LEU:HB2	6:S4:95:THR:HG21	4.16	0.42
7:S5:185:ARG:NH1	1:6:1572:G:H1'	331.27	0.42
7:S5:39:GLU:HB3	7:S5:40:ILE:H	1.65	0.42
8:S6:196:ARG:O	8:S6:199:GLN:HB2	3.10	0.42
8:S6:20:ASP:OD2	8:S6:23:ARG:N	4.28	0.42
9:S7:60:ILE:HD12	9:S7:92:PHE:CZ	2.55	0.42
11:S9:65:LYS:HA	11:S9:65:LYS:HD3	4.06	0.42
34:SR:133:VAL:O	34:SR:141:LEU:HB2	2.20	0.42
34:SR:134:TRP:CZ3	34:SR:140:CYS:HB2	2.54	0.42
34:SR:283:LYS:HB2	34:SR:283:LYS:HE3	1.67	0.42
34:SR:23:LEU:HD22	34:SR:33:LEU:HD11	4.68	0.42
36:1:1151:U:O4	36:1:1200:A:N6	2.50	0.42
36:1:1414:G:N7	86:1:4118:OHX:N2	2.67	0.42
36:1:1504:A:C5	36:1:1505:C:C5	3.08	0.42
36:1:1803:C:H2'	36:1:1804:A:C8	2.54	0.42
36:1:185:C:H2'	36:1:186:U:H6	1.84	0.42
36:1:193:C:H2'	36:1:194:U:C6	2.54	0.42
36:1:207:U:H2'	36:1:208:C:C6	2.55	0.42
36:1:2389:C:O2'	36:1:2390:A:H5'	2.19	0.42
36:1:2691:A:H2'	36:1:2692:A:O4'	2.19	0.42
36:1:2738:A:C6	36:1:2739:A:C5	3.08	0.42
36:1:2943:G:O5'	36:1:2943:G:H8	2.03	0.42
36:1:2993:G:H2'	36:1:3142:A:H61	1.85	0.42
36:1:3350:C:HO2'	36:1:3351:U:P	2.41	0.42
36:1:3020:U:O4	86:1:3982:OHX:N4	2.53	0.42
36:1:968:G:C5	36:1:969:C:C4	3.07	0.42
1:2:160:C:H2'	1:2:161:U:O4'	2.20	0.42
1:2:1791:A:H5''	28:D6:8:ASN:HD22	1.83	0.42
1:2:196:G:O2'	1:2:197:A:P	2.77	0.42
86:2:2074:OHX:N3	86:2:2162:OHX:N1	2.68	0.42
86:2:2074:OHX:N3	86:2:2162:OHX:N5	2.67	0.42
1:2:256:A:H2'	1:2:257:A:O4'	2.20	0.42
1:2:428:A:H2'	1:2:429:G:O4'	2.19	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:475:A:C6	1:2:476:U:C2	3.08	0.42
1:2:827:C:H2'	1:2:828:U:O4'	2.19	0.42
36:1:18:G:N2	38:4:142:C:C2	2.88	0.42
36:5:127:G:H2'	36:5:128:G:C8	2.55	0.42
36:5:1615:C:C2	36:5:1616:U:C5	3.08	0.42
36:5:2342:U:OP1	36:5:3089:C:O2'	2.35	0.42
39:L2:89:TYR:CZ	36:5:2551:U:C2	224.88	0.42
36:5:3227:A:C2'	36:5:3228:C:H5'	2.42	0.42
41:L4:196:ASN:ND2	36:5:337:G:OP2	91.93	0.42
36:5:2658:G:OP2	86:5:3901:OHX:N1	2.53	0.42
1:6:1041:G:N2	1:6:1042:G:C2	2.88	0.42
1:6:1164:G:H2'	1:6:1165:G:C8	2.55	0.42
1:6:1255:G:O2'	1:6:1256:A:H8	2.02	0.42
1:6:1397:U:C4	1:6:1399:C:H1'	2.54	0.42
1:6:1458:G:C2	1:6:1459:C:C4	3.08	0.42
21:C9:79:LEU:HD13	1:6:1523:G:C8	407.02	0.42
1:6:1592:A:C2	1:6:1605:G:C2	3.07	0.42
86:6:2059:OHX:N5	86:6:2147:OHX:N6	2.68	0.42
1:6:705:U:HO2'	1:6:706:A:H8	1.66	0.42
1:6:774:A:C2	1:6:775:G:C8	3.07	0.42
16:C4:52:ARG:N	1:6:906:A:OP2	292.69	0.42
38:8:68:G:C6	38:8:69:U:C4	3.08	0.42
38:8:76:C:H2'	38:8:77:A:O4'	2.20	0.42
5:S3:72:LEU:HD22	12:C0:65:TYR:HB3	2.57	0.42
12:C0:80:LEU:O	12:C0:82:LEU:N	2.53	0.42
14:C2:40:GLY:O	14:C2:124:LYS:N	3.35	0.42
15:C3:51:GLY:O	15:C3:55:ARG:N	2.53	0.42
16:C4:25:ASP:HA	16:C4:54:GLU:O	2.19	0.42
20:C8:144:ARG:HB2	20:C8:145:ARG:H	1.70	0.42
20:C8:6:GLN:O	27:D5:42:LEU:HD13	2.20	0.42
21:C9:64:HIS:O	21:C9:68:ARG:HG2	2.20	0.42
22:D0:50:LEU:HD21	22:D0:95:ALA:HB2	2.01	0.42
23:D1:74:GLN:HG2	23:D1:79:LEU:HB2	4.35	0.42
24:D2:32:LYS:HB2	24:D2:32:LYS:HE3	1.82	0.42
24:D2:90:THR:O	24:D2:94:LEU:HB2	2.20	0.42
26:D4:118:ILE:HG23	26:D4:125:LEU:HD23	2.02	0.42
27:D5:47:TYR:O	27:D5:50:ILE:HG22	2.20	0.42
30:D8:29:ARG:HA	30:D8:41:VAL:HA	2.01	0.42
39:L2:158:ILE:HG21	39:L2:158:ILE:HD13	1.97	0.42
41:L4:142:VAL:C	41:L4:144:LYS:N	2.73	0.42
43:L6:155:LEU:HA	43:L6:155:LEU:HD23	1.76	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
46:L9:157:ASN:C	46:L9:157:ASN:HD22	2.18	0.42
46:L9:13:PRO:O	46:L9:16:VAL:HG13	4.24	0.42
36:1:3122:A:N1	46:L9:70:THR:HG21	2.35	0.42
47:M0:24:ARG:O	47:M0:25:ALA:HB3	4.59	0.42
49:M3:50:PRO:O	49:M3:52:ASP:N	4.19	0.42
49:M3:57:VAL:HG13	49:M3:147:ILE:HG23	2.01	0.42
49:M3:64:LYS:HD3	49:M3:65:TYR:CZ	2.79	0.42
50:M4:24:LYS:HG3	50:M4:25:LYS:CD	2.50	0.42
50:M4:42:LYS:HE2	50:M4:42:LYS:HB3	4.01	0.42
52:M6:87:MET:HG2	36:5:1175:C:O2	252.06	0.42
55:M9:17:VAL:HG13	55:M9:21:LYS:HB2	2.02	0.42
55:M9:25:ASP:HA	55:M9:26:PRO:HD2	2.01	0.42
57:N1:160:ILE:HA	57:N1:160:ILE:HD12	2.09	0.42
57:N1:85:LEU:HD23	57:N1:85:LEU:HA	2.19	0.42
57:N1:8:ARG:O	57:N1:11:THR:HG23	2.19	0.42
62:N6:118:LEU:O	62:N6:122:LYS:HG3	2.20	0.42
63:N7:60:LYS:O	63:N7:63:ALA:HB3	3.35	0.42
64:N8:2:PRO:HG2	64:N8:5:PHE:CD2	4.72	0.42
66:O0:38:LYS:C	66:O0:93:LEU:HD23	3.48	0.42
68:O2:19:ARG:HD3	68:O2:19:ARG:HH11	1.69	0.42
69:O3:6:ARG:HG3	69:O3:8:TYR:CE1	2.55	0.42
36:1:1589:A:H4'	70:O4:11:ASN:HD22	1.85	0.42
70:O4:43:LYS:HA	70:O4:50:ALA:HA	2.01	0.42
76:Q0:82:LEU:HD23	76:Q0:82:LEU:HA	2.03	0.42
78:Q2:35:LEU:C	78:Q2:37:ALA:H	2.22	0.42
78:Q2:63:LYS:HA	78:Q2:63:LYS:HD3	2.55	0.42
79:Q3:8:VAL:H	79:Q3:8:VAL:HG13	1.62	0.42
2:S0:125:ASP:HA	2:S0:126:PRO:HD2	1.75	0.42
2:S0:195:TRP:CD2	2:S0:197:ILE:HB	4.06	0.42
6:S4:153:ASN:OD1	8:S6:215:ARG:NH1	2.53	0.42
7:S5:151:GLY:HA3	7:S5:155:ALA:HA	5.53	0.42
7:S5:57:SER:OG	7:S5:58:LEU:HG	2.20	0.42
8:S6:137:ARG:HG3	1:6:169:A:OP1	322.62	0.42
8:S6:154:ARG:C	8:S6:156:PHE:H	3.22	0.42
9:S7:61:PHE:CE1	9:S7:93:LEU:HD12	3.40	0.42
11:S9:129:ILE:HA	11:S9:134:ILE:CD1	2.49	0.42
11:S9:171:ARG:O	11:S9:175:ARG:HB2	3.26	0.42
34:SR:245:PHE:CD1	34:SR:252:LEU:HD13	2.73	0.42
36:1:1764:U:H3'	36:1:1765:U:C4'	2.50	0.42
36:1:2376:G:C6	36:1:2377:G:C6	3.08	0.42
36:1:2529:A:C2	36:1:2530:G:H1'	2.55	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:2571:U:H4'	36:1:2572:C:OP1	2.20	0.42
36:1:2659:G:C2	36:1:2712:U:O2	2.72	0.42
36:1:2721:A:O3'	65:N9:33:LYS:HE2	2.20	0.42
36:1:3034:C:O2	46:L9:122:LYS:HB3	2.20	0.42
36:1:541:U:O4	86:1:4189:OHX:N4	2.52	0.42
36:1:802:C:C2	36:1:803:C:C5	3.07	0.42
36:1:815:G:C2	36:1:926:A:C2	3.07	0.42
1:2:1043:A:C2	1:2:1076:A:C2	3.08	0.42
1:2:1308:G:C2	1:2:1309:C:C2	3.07	0.42
1:2:1542:G:H22	1:2:1568:C:H1'	1.78	0.42
1:2:192:U:O2'	1:2:193:U:O5'	2.33	0.42
86:2:2089:OHX:N1	86:2:2131:OHX:N2	2.68	0.42
1:2:325:G:H2'	1:2:326:G:H8	1.85	0.42
37:3:113:C:H2'	37:3:114:U:O4'	2.20	0.42
38:4:7:U:H2'	38:4:8:C:C6	2.54	0.42
36:5:1014:U:H4'	36:5:1014:U:OP1	2.20	0.42
35:SM:48:ARG:NH1	36:5:1017:C:H5''	337.73	0.42
36:5:1609:C:H2'	36:5:1610:G:C8	2.55	0.42
36:5:1925:U:OP2	36:5:1926:C:N4	2.52	0.42
36:5:2115:G:N3	36:5:2119:A:C2	2.87	0.42
36:5:2186:U:H5'	36:5:2314:U:OP2	2.20	0.42
36:5:1205:A:H4'	36:5:2835:U:O2'	2.19	0.42
36:5:3044:G:H2'	36:5:3045:G:H8	1.85	0.42
36:5:3105:U:H2'	36:5:3106:A:C8	2.55	0.42
36:5:3335:A:C2	36:5:3336:A:C4	3.08	0.42
36:5:501:A:C4	36:5:502:U:C5	3.08	0.42
36:5:848:A:C5	36:5:849:C:H1'	2.54	0.42
1:6:1030:A:C8	1:6:1792:G:C2	3.08	0.42
1:6:1165:G:H2'	1:6:1166:A:O4'	2.19	0.42
1:6:1230:A:C8	1:6:1258:U:C4	3.07	0.42
1:6:1327:C:H6	1:6:1327:C:O5'	2.03	0.42
1:6:1512:G:H2'	1:6:1513:G:H8	1.84	0.42
1:6:329:G:H2'	1:6:330:G:C8	2.55	0.42
1:6:680:U:C2	1:6:682:C:N4	2.88	0.42
37:7:73:C:H2'	37:7:73:C:H6	1.62	0.42
10:S8:187:GLU:HG3	13:C1:30:ARG:NH1	2.34	0.42
15:C3:86:GLU:O	15:C3:89:TYR:HB3	2.19	0.42
16:C4:18:ARG:HA	16:C4:82:LYS:O	2.72	0.42
17:C5:111:MET:HG2	20:C8:119:ILE:HG13	2.71	0.42
18:C6:14:LYS:O	18:C6:123:ARG:NH1	2.52	0.42
19:C7:15:ALA:O	19:C7:19:ARG:HG2	2.51	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:C9:33:TYR:CD1	21:C9:33:TYR:C	2.94	0.42
21:C9:23:GLN:HG3	21:C9:55:TYR:CE2	4.35	0.42
24:D2:104:LEU:O	24:D2:110:ILE:HG23	3.45	0.42
25:D3:106:GLY:O	1:6:599:A:H5''	359.15	0.42
25:D3:54:LEU:HD21	25:D3:75:GLN:HB2	2.30	0.42
26:D4:10:ARG:O	26:D4:24:VAL:HG23	2.20	0.42
26:D4:57:VAL:HG22	26:D4:60:PHE:HE2	1.84	0.42
22:D0:70:THR:O	31:D9:40:ARG:NH1	2.53	0.42
39:L2:84:THR:OG1	79:Q3:63:THR:N	2.51	0.42
36:1:2987:A:O2'	40:L3:259:HIS:HB3	2.19	0.42
44:L7:89:ILE:HD11	44:L7:135:ALA:HB3	2.01	0.42
44:L7:26:VAL:HG23	44:L7:27:ALA:H	1.85	0.42
44:L7:88:ARG:NH1	44:L7:92:ILE:HD13	2.35	0.42
45:L8:153:ILE:O	45:L8:179:ILE:HA	2.49	0.42
47:M0:200:LEU:HD12	47:M0:213:PHE:HB2	2.92	0.42
48:M1:92:ARG:HA	48:M1:171:VAL:O	2.19	0.42
48:M1:98:ALA:HA	48:M1:156:LYS:HB2	2.55	0.42
49:M3:62:THR:OG1	49:M3:62:THR:O	2.90	0.42
52:M6:76:PRO:HD3	52:M6:147:TRP:CD2	2.54	0.42
56:N0:155:ARG:HG3	56:N0:155:ARG:O	2.20	0.42
57:N1:87:LYS:HE3	57:N1:87:LYS:HB3	3.90	0.42
63:N7:103:GLN:HA	63:N7:104:PRO:HD3	1.91	0.42
63:N7:5:LEU:HD11	63:N7:30:ASP:OD2	9.07	0.42
67:O1:51:LEU:HD22	67:O1:55:LEU:HD13	4.79	0.42
68:O2:24:ARG:HG2	68:O2:25:TYR:CZ	2.55	0.42
43:L6:3:ALA:HB1	68:O2:75:LEU:HD13	2.02	0.42
75:O9:49:MET:O	75:O9:50:ASN:HB2	2.26	0.42
36:1:3120:C:H3'	76:Q0:111:ARG:NH2	2.34	0.42
77:Q1:9:ARG:HG3	77:Q1:9:ARG:NH1	3.66	0.42
36:1:2226:U:OP1	78:Q2:35:LEU:HD11	2.20	0.42
79:Q3:22:LEU:HD23	79:Q3:22:LEU:HA	2.02	0.42
79:Q3:35:ALA:HB3	79:Q3:37:TYR:HE2	3.11	0.42
2:S0:87:LEU:HA	2:S0:87:LEU:HD12	2.72	0.42
3:S1:43:VAL:HG13	3:S1:68:VAL:HG21	3.64	0.42
3:S1:29:TRP:NE1	3:S1:47:LEU:HG	2.25	0.42
3:S1:69:CYS:SG	16:C4:114:ARG:HD3	2.60	0.42
6:S4:95:THR:HG22	26:D4:16:PRO:HG2	2.02	0.42
7:S5:30:PRO:O	7:S5:33:VAL:HB	2.20	0.42
7:S5:35:GLN:C	7:S5:37:GLN:H	2.22	0.42
8:S6:30:LYS:HB3	8:S6:30:LYS:HE3	1.88	0.42
8:S6:48:TYR:CE1	8:S6:116:LYS:HA	2.55	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:S7:75:THR:HG23	9:S7:161:GLN:OE1	3.77	0.42
1:2:330:G:O2'	10:S8:33:PRO:HB3	2.20	0.42
1:2:329:G:H5'	10:S8:99:ALA:HB3	2.00	0.42
34:SR:106:HIS:CD2	34:SR:110:VAL:HG22	3.18	0.42
34:SR:84:SER:OG	34:SR:86:ASP:OD1	3.50	0.42
36:1:1351:U:H2'	36:1:1351:U:O2	2.19	0.42
36:1:1639:C:N4	70:O4:73:SER:HB2	2.35	0.42
36:1:1939:G:O5'	36:1:1939:G:H8	2.02	0.42
36:1:242:C:HO2'	36:1:243:G:P	2.43	0.42
36:1:2997:G:C6	36:1:3396:U:C4	3.08	0.42
36:1:607:A:H2'	36:1:607:A:N3	2.34	0.42
36:1:763:G:HO2'	36:1:764:U:P	2.42	0.42
36:1:812:G:H2'	36:1:813:G:O4'	2.20	0.42
36:1:852:U:H2'	36:1:853:G:H8	1.84	0.42
1:2:1046:G:C4	1:2:1073:G:N2	2.88	0.42
1:2:1390:U:P	19:C7:49:LYS:HG3	2.59	0.42
1:2:187:G:H3'	10:S8:138:ASN:HD21	1.85	0.42
1:2:276:C:H42	1:2:281:G:H1	1.68	0.42
37:3:71:G:O2'	37:3:72:A:H5'	2.20	0.42
36:5:129:U:H2'	36:5:130:A:C8	2.55	0.42
36:5:1444:G:H2'	36:5:1445:U:O4'	2.20	0.42
36:5:1805:C:H2'	36:5:1806:A:C8	2.55	0.42
36:5:1822:C:H2'	36:5:1823:A:C8	2.55	0.42
36:5:1844:C:C2	36:5:1845:G:C8	3.08	0.42
36:5:2140:U:H2'	36:5:2977:G:O2'	2.19	0.42
36:5:2185:G:O2'	36:5:2314:U:OP2	2.38	0.42
36:5:2653:C:O2	36:5:2694:A:C6	2.73	0.42
36:5:2715:A:C2	36:5:2753:G:C6	3.08	0.42
36:5:3277:U:H2'	36:5:3278:C:O4'	2.19	0.42
36:5:26:A:C2	36:5:330:G:C5	3.07	0.42
36:5:916:G:N7	36:5:924:G:C5	2.88	0.42
1:6:1073:G:H2'	1:6:1074:G:H5''	2.00	0.42
1:6:1347:U:O2	1:6:1516:A:H5'	2.20	0.42
1:6:1561:U:H4'	1:6:1599:C:H4'	2.01	0.42
1:6:1697:G:H8	1:6:1705:C:C4	2.38	0.42
1:6:1796:C:H5'	1:6:1797:A:C8	2.55	0.42
1:6:722:G:O2'	1:6:723:G:H8	2.03	0.42
15:C3:140:LYS:HD3	15:C3:142:GLU:HG3	8.05	0.42
15:C3:27:LYS:CE	15:C3:27:LYS:H	2.28	0.42
18:C6:40:GLU:HG3	18:C6:41:PRO:C	2.40	0.42
18:C6:43:ILE:HG12	18:C6:43:ILE:H	1.99	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:C8:121:ALA:O	20:C8:124:GLY:N	2.53	0.42
20:C8:42:TYR:CE2	20:C8:73:MET:HG3	5.01	0.42
20:C8:8:GLN:HB3	20:C8:9:GLY:H	3.21	0.42
21:C9:9:VAL:HG22	21:C9:140:LEU:HD11	2.02	0.42
27:D5:86:GLU:O	27:D5:88:ILE:N	3.89	0.42
32:E0:46:ASN:O	32:E0:47:VAL:HG12	2.20	0.42
32:E0:49:LEU:HD11	32:E0:55:ARG:HB2	2.01	0.42
39:L2:41:ILE:HG23	39:L2:63:PHE:CD2	2.55	0.42
40:L3:233:TRP:CG	40:L3:265:ALA:HB1	3.04	0.42
41:L4:142:VAL:C	41:L4:144:LYS:H	2.22	0.42
41:L4:321:LYS:O	41:L4:324:LEU:HB3	2.19	0.42
42:L5:131:LEU:HD13	42:L5:131:LEU:N	2.35	0.42
42:L5:158:ARG:CB	42:L5:158:ARG:HH21	6.09	0.42
43:L6:43:LEU:HD12	69:O3:105:SER:HB2	2.02	0.42
44:L7:40:LYS:HE2	44:L7:170:GLU:CD	4.88	0.42
44:L7:160:ARG:HG3	44:L7:203:TRP:CG	3.21	0.42
49:M3:95:ILE:HG12	49:M3:119:TYR:CE2	2.71	0.42
52:M6:10:ASP:HB2	52:M6:117:ARG:HB2	2.02	0.42
52:M6:48:PHE:CE1	52:M6:52:LEU:HD21	3.11	0.42
55:M9:101:VAL:HG13	55:M9:104:ARG:NH2	2.35	0.42
55:M9:105:LEU:HD23	55:M9:105:LEU:C	2.39	0.42
55:M9:14:VAL:HG21	55:M9:41:ILE:HG22	2.02	0.42
55:M9:9:ARG:NH2	36:5:1602:A:O3'	108.58	0.42
58:N2:39:ASP:OD2	58:N2:39:ASP:N	2.87	0.42
59:N3:87:ARG:HH22	59:N3:137:VAL:CG2	2.24	0.42
63:N7:22:LYS:HE3	63:N7:134:LEU:HB2	2.02	0.42
70:O4:66:SER:HB2	70:O4:69:HIS:CE1	4.66	0.42
75:O9:17:LYS:O	75:O9:19:GLN:N	3.32	0.42
2:S0:32:HIS:HD2	2:S0:153:SER:HB2	9.65	0.42
2:S0:8:ASP:C	2:S0:54:TRP:HE1	2.37	0.42
3:S1:190:PRO:HG2	3:S1:192:VAL:CG2	3.06	0.42
3:S1:225:VAL:O	3:S1:228:LEU:HB3	2.20	0.42
4:S2:41:LEU:HD11	4:S2:56:ILE:HG12	2.02	0.42
4:S2:74:PRO:O	4:S2:76:LEU:N	2.49	0.42
5:S3:124:ARG:O	5:S3:128:GLU:HB2	2.20	0.42
5:S3:160:SER:O	5:S3:164:VAL:HB	4.44	0.42
5:S3:168:ILE:H	5:S3:168:ILE:HG13	1.59	0.42
5:S3:29:LEU:HA	5:S3:32:GLU:OE1	2.20	0.42
6:S4:207:LEU:HD23	6:S4:207:LEU:HA	2.01	0.42
6:S4:221:ARG:HG2	6:S4:221:ARG:H	2.41	0.42
8:S6:147:LEU:HB3	8:S6:151:ASP:OD1	3.98	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:S7:38:LEU:HA	9:S7:38:LEU:HD23	2.40	0.42
10:S8:21:PHE:CD1	10:S8:22:ARG:HG2	4.56	0.42
1:2:471:A:O3'	11:S9:10:LYS:HA	2.20	0.42
11:S9:33:GLU:HB2	11:S9:34:PHE:CD2	4.60	0.42
11:S9:5:PRO:HG3	1:6:380:U:C4	369.10	0.42
35:SM:125:ALA:C	35:SM:127:ALA:H	3.40	0.42
34:SR:23:LEU:HB2	34:SR:293:ALA:HB3	2.02	0.42
36:1:1103:A:H1'	36:1:1104:G:OP1	2.20	0.41
36:1:2124:G:C2	36:1:2125:A:C5	3.08	0.41
36:1:2148:U:H2'	36:1:2149:A:C4	2.55	0.41
36:1:2519:A:C4	36:1:2589:G:N2	2.87	0.41
36:1:2746:A:OP2	42:L5:178:ASN:ND2	2.49	0.41
36:1:3191:G:H2'	36:1:3192:U:O4'	2.20	0.41
86:1:4023:OHX:N4	86:1:4143:OHX:N1	2.68	0.41
86:1:4200:OHX:N4	38:4:140:G:OP1	2.53	0.41
36:1:602:A:N6	36:1:603:A:N1	2.68	0.41
36:1:627:U:H2'	36:1:628:A:C8	2.55	0.41
1:2:1240:U:C2'	1:2:1241:G:H5''	2.50	0.41
1:2:1583:A:N1	1:2:1611:A:H5''	2.34	0.41
1:2:553:G:C5	1:2:554:C:C2	3.08	0.41
65:N9:38:LYS:HD3	36:5:1076:C:H4'	211.97	0.41
36:5:1151:U:C5	36:5:1152:G:C6	3.08	0.41
36:5:1241:U:HO2'	36:5:1242:G:P	2.42	0.41
36:5:171:G:H8	36:5:171:G:OP2	2.02	0.41
36:5:2562:A:C5	36:5:2563:G:C8	3.07	0.41
86:5:4060:OHX:N3	86:5:4138:OHX:N4	2.68	0.41
36:5:612:U:H2'	36:5:613:G:C8	2.55	0.41
36:5:706:A:O2'	36:5:707:U:H5'	2.19	0.41
36:5:711:A:N7	36:5:712:G:H1'	2.34	0.41
36:5:78:U:H2'	36:5:79:U:H6	1.84	0.41
21:C9:130:ARG:HB3	1:6:1358:G:H4'	430.25	0.41
1:6:265:A:C2	1:6:267:U:C4	3.08	0.41
1:6:53:G:N2	1:6:427:C:N3	2.56	0.41
1:6:626:U:H2'	1:6:627:C:H6	1.85	0.41
1:6:72:A:C6	1:6:73:U:N3	2.88	0.41
1:6:755:A:O2'	1:6:756:A:O4'	2.38	0.41
1:6:219:A:H2'	1:6:831:U:O2	2.20	0.41
1:6:999:U:O2	1:6:1006:C:N4	2.53	0.41
38:8:24:G:N2	38:8:25:G:H1'	2.35	0.41
15:C3:40:TYR:CE2	15:C3:53:LEU:HD23	3.51	0.41
20:C8:18:LEU:HD21	20:C8:70:VAL:HG13	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:C9:133:ASP:OD1	1:6:1358:G:O2'	427.31	0.41
21:C9:6:VAL:HG13	21:C9:66:TYR:CZ	3.32	0.41
22:D0:107:THR:O	22:D0:108:ILE:HD12	4.55	0.41
23:D1:73:ALA:HB1	23:D1:78:LEU:HD11	2.02	0.41
23:D1:85:TYR:HA	23:D1:85:TYR:HD2	2.15	0.41
25:D3:127:VAL:CG2	25:D3:142:LYS:HE3	2.50	0.41
29:D7:40:CYS:C	29:D7:42:ASN:H	2.45	0.41
33:E1:104:SER:O	33:E1:106:TYR:N	2.52	0.41
39:L2:227:ARG:HH11	39:L2:227:ARG:HG3	1.85	0.41
39:L2:71:LEU:HD13	36:5:1651:U:H5'	188.80	0.41
40:L3:261:MET:HE3	40:L3:261:MET:HB3	4.27	0.41
40:L3:291:GLU:OE1	40:L3:302:LYS:NZ	3.33	0.41
41:L4:6:VAL:HG21	41:L4:255:PHE:HZ	2.05	0.41
36:1:805:G:H1'	41:L4:73:ARG:HH11	1.85	0.41
42:L5:104:LEU:HB2	42:L5:247:ILE:HD13	3.73	0.41
37:3:1:G:N2	42:L5:265:TYR:HB3	2.35	0.41
42:L5:278:SER:O	42:L5:281:GLU:HG2	4.78	0.41
42:L5:31:TYR:O	42:L5:35:ARG:HD2	2.92	0.41
86:1:4190:OHX:N4	43:L6:129:GLU:HB3	2.35	0.41
44:L7:120:THR:C	44:L7:122:ALA:N	2.99	0.41
44:L7:218:ARG:NH1	37:7:86:U:O2'	257.99	0.41
46:L9:126:VAL:HA	46:L9:127:PRO:HD2	2.89	0.41
46:L9:84:LYS:HA	46:L9:188:THR:HG23	2.02	0.41
46:L9:48:VAL:O	46:L9:49:ASN:HB3	2.20	0.41
47:M0:71:CYS:SG	47:M0:155:ALA:HA	3.30	0.41
47:M0:9:TYR:CD2	47:M0:97:LEU:HD13	2.88	0.41
48:M1:8:PRO:HD2	48:M1:10:ARG:CG	2.50	0.41
49:M3:167:PHE:O	49:M3:170:LEU:HB2	2.19	0.41
51:M5:14:LYS:HA	51:M5:19:LEU:CD2	2.50	0.41
36:1:2790:A:OP2	54:M8:181:SER:HB3	2.20	0.41
54:M8:32:LEU:O	54:M8:35:PHE:HB3	2.28	0.41
56:N0:27:MET:HG2	57:N1:151:LEU:O	2.20	0.41
59:N3:74:MET:HE3	59:N3:102:ILE:HD13	2.01	0.41
59:N3:126:TRP:HA	59:N3:127:PRO:HD3	1.73	0.41
62:N6:37:LYS:HD3	62:N6:37:LYS:H	3.36	0.41
62:N6:56:VAL:HG22	62:N6:105:VAL:O	2.71	0.41
63:N7:10:VAL:HB	63:N7:83:THR:HG22	2.01	0.41
65:N9:32:LEU:HB2	65:N9:40:ARG:NH1	2.53	0.41
66:O0:83:LYS:HG2	66:O0:85:PHE:CZ	2.74	0.41
67:O1:74:ARG:NH1	67:O1:109:VAL:HG11	2.89	0.41
67:O1:35:GLU:O	67:O1:38:LYS:HB3	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
43:L6:13:GLU:OE1	68:O2:90:LYS:HB2	2.20	0.41
74:O8:12:LEU:HD13	74:O8:12:LEU:HA	4.06	0.41
75:O9:48:LYS:HA	75:O9:48:LYS:HD2	2.29	0.41
2:S0:124:THR:CG2	2:S0:174:TRP:HE1	2.48	0.41
4:S2:49:LYS:HD3	4:S2:49:LYS:HA	2.18	0.41
5:S3:163:PRO:O	5:S3:167:PHE:HD2	2.42	0.41
1:2:1437:U:H5'	5:S3:176:LEU:HD23	2.02	0.41
5:S3:210:GLU:HA	5:S3:211:PRO:HD3	2.03	0.41
5:S3:32:GLU:O	5:S3:52:ALA:HB1	2.19	0.41
1:2:454:U:H4'	6:S4:62:LYS:HE3	2.02	0.41
7:S5:157:ARG:O	7:S5:224:ASN:HB3	2.20	0.41
7:S5:160:VAL:HG12	30:D8:43:ASN:HB2	2.21	0.41
7:S5:30:PRO:HB2	7:S5:33:VAL:HB	2.02	0.41
7:S5:89:ILE:O	7:S5:93:LEU:HB2	3.99	0.41
8:S6:137:ARG:HH21	8:S6:177:ARG:HE	1.68	0.41
8:S6:137:ARG:O	8:S6:138:ALA:C	2.59	0.41
8:S6:67:VAL:HG23	8:S6:100:ALA:H	1.85	0.41
9:S7:29:ASN:O	9:S7:30:SER:OG	2.32	0.41
11:S9:92:LYS:HB2	11:S9:95:TYR:CD2	9.58	0.41
35:SM:49:LYS:HZ3	35:SM:53:ARG:HH12	13.36	0.41
36:1:1235:U:C4'	36:1:1236:G:H5'	2.48	0.41
36:1:1522:U:H4'	36:1:1523:U:OP2	2.20	0.41
36:1:2264:U:OP2	86:1:3979:OHX:N5	2.54	0.41
36:1:2144:A:C4	36:1:2281:A:C6	3.08	0.41
36:1:22:G:O4'	38:4:104:A:H1'	2.20	0.41
36:1:2886:U:C5	36:1:2911:A:C5	3.08	0.41
36:1:2943:G:H2'	36:1:2944:U:O4'	2.20	0.41
36:1:2955:U:H6	36:1:2955:U:O5'	2.03	0.41
36:1:28:C:O2'	36:1:29:C:H5'	2.21	0.41
36:1:3189:G:C2	36:1:3190:C:C2	3.09	0.41
36:1:336:A:O2'	36:1:337:G:H5'	2.20	0.41
36:1:3384:U:C2	36:1:3385:U:C5	3.09	0.41
36:1:498:A:OP1	69:O3:86:ARG:NE	2.50	0.41
36:1:985:U:H2'	36:1:986:U:C6	2.52	0.41
1:2:1219:A:O2'	12:C0:48:SER:HA	2.21	0.41
1:2:1288:G:C6	1:2:1328:G:C2	3.09	0.41
1:2:1552:U:C5	1:2:1553:G:C5	3.08	0.41
1:2:1564:U:H2'	1:2:1565:C:C6	2.56	0.41
1:2:1669:U:H2'	1:2:1670:G:O4'	2.19	0.41
1:2:980:G:O6	86:2:2044:OHX:N2	2.53	0.41
1:2:495:C:H3'	1:2:496:G:C4'	2.49	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:813:U:H5'	15:C3:76:LYS:HD3	2.02	0.41
1:2:892:A:C5	1:2:893:U:C4	3.08	0.41
1:2:895:G:H2'	1:2:896:U:C6	2.56	0.41
1:2:952:A:OP1	15:C3:94:LYS:HE2	2.20	0.41
37:3:113:C:C4	37:3:114:U:C4	3.08	0.41
37:3:92:A:C5	37:3:93:C:H1'	2.55	0.41
36:5:1432:C:O2'	36:5:1433:A:H3'	2.21	0.41
36:5:1547:G:H2'	36:5:1548:C:C6	2.55	0.41
36:5:1666:G:C6	36:5:1667:A:C6	3.08	0.41
39:L2:8:GLN:NE2	36:5:2164:A:OP1	179.84	0.41
36:5:2566:C:O2	36:5:2576:G:C2	2.74	0.41
36:5:2815:G:H5''	36:5:2816:G:OP2	2.20	0.41
36:5:3163:A:C6	36:5:3288:G:C6	3.08	0.41
86:5:3930:OHX:N5	38:8:7:U:O4	2.53	0.41
36:5:2169:G:O6	86:5:3947:OHX:N5	2.53	0.41
36:5:370:U:OP1	86:5:4161:OHX:N1	2.53	0.41
36:5:686:G:C6	36:5:687:U:C2	3.08	0.41
36:5:701:G:C6	36:5:702:C:C4	3.08	0.41
36:5:938:C:OP1	36:5:963:G:H5'	2.20	0.41
33:E1:87:THR:HB	1:6:1445:G:O6	380.59	0.41
77:Q1:21:ARG:NH1	1:6:1654:G:OP1	282.55	0.41
1:6:1779:U:H2'	1:6:1781:A:OP2	2.20	0.41
1:6:373:G:H2'	1:6:374:U:C6	2.54	0.41
1:6:473:A:N6	1:6:474:A:C2	2.87	0.41
1:6:545:A:C2	1:6:593:U:O2	2.73	0.41
1:6:30:G:C6	1:6:597:G:C6	3.08	0.41
1:6:643:G:C2	1:6:644:C:C2	3.08	0.41
1:6:771:A:C2	1:6:772:G:H1'	2.55	0.41
38:8:129:C:H2'	38:8:130:C:H6	1.85	0.41
12:C0:32:HIS:HB3	12:C0:34:GLU:O	5.27	0.41
17:C5:63:ALA:HB1	17:C5:74:ALA:HB3	2.97	0.41
17:C5:67:ALA:O	86:C5:201:OHX:N2	2.52	0.41
22:D0:105:GLN:HG3	22:D0:106:ILE:H	1.84	0.41
23:D1:24:ILE:HD12	23:D1:31:SER:HB2	5.54	0.41
24:D2:106:THR:HG21	24:D2:111:MET:HE3	2.02	0.41
24:D2:115:GLU:HA	24:D2:118:ARG:CZ	2.50	0.41
25:D3:38:PHE:N	25:D3:38:PHE:CD2	3.66	0.41
26:D4:117:LYS:O	26:D4:118:ILE:HD13	2.20	0.41
26:D4:57:VAL:HG13	26:D4:58:PHE:N	2.35	0.41
7:S5:162:VAL:HB	30:D8:45:LYS:HB3	2.07	0.41
32:E0:4:VAL:HG12	32:E0:4:VAL:O	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:L2:27:ALA:HA	39:L2:75:ILE:HG22	2.01	0.41
40:L3:85:VAL:HG22	40:L3:163:HIS:CD2	2.55	0.41
36:1:2392:C:H1'	40:L3:266:ARG:HH12	1.84	0.41
40:L3:35:ASP:OD2	40:L3:37:ARG:HD2	2.24	0.41
41:L4:303:GLY:H	36:5:1347:U:H5''	197.96	0.41
41:L4:356:THR:O	41:L4:359:LEU:N	3.18	0.41
41:L4:361:HIS:NE2	41:L4:362:ASP:HB2	3.87	0.41
42:L5:277:LEU:HA	42:L5:277:LEU:HD12	1.88	0.41
42:L5:36:LEU:HA	42:L5:36:LEU:HD23	1.95	0.41
42:L5:78:ALA:HA	42:L5:82:GLU:OE2	2.21	0.41
43:L6:31:ARG:HE	43:L6:31:ARG:HB3	1.59	0.41
44:L7:159:GLN:HA	36:5:1362:G:O2'	217.02	0.41
44:L7:239:LEU:HA	44:L7:239:LEU:HD23	2.14	0.41
46:L9:91:ARG:HD3	46:L9:143:GLU:OE2	2.21	0.41
46:L9:94:TYR:CD1	46:L9:94:TYR:N	3.00	0.41
47:M0:175:ASN:C	47:M0:176:LEU:HG	4.42	0.41
47:M0:169:LYS:O	47:M0:178:ARG:HG2	2.20	0.41
48:M1:100:GLY:O	48:M1:159:THR:HG21	2.20	0.41
48:M1:154:THR:O	48:M1:154:THR:OG1	2.37	0.41
50:M4:14:LEU:H	50:M4:19:ARG:HH11	2.32	0.41
50:M4:22:LEU:HD23	50:M4:99:TRP:CZ2	3.16	0.41
51:M5:101:THR:O	51:M5:105:ARG:HG3	2.20	0.41
52:M6:127:LEU:HD11	56:N0:168:PRO:HG3	2.02	0.41
52:M6:84:LEU:HD22	52:M6:102:LEU:HD22	3.14	0.41
54:M8:162:ALA:HA	54:M8:163:PRO:HD2	1.85	0.41
55:M9:15:VAL:H	55:M9:15:VAL:HG23	1.58	0.41
56:N0:99:ARG:O	56:N0:103:VAL:HG23	2.42	0.41
56:N0:91:TYR:CZ	56:N0:93:GLU:HG2	2.56	0.41
57:N1:116:ARG:O	57:N1:120:LYS:N	2.53	0.41
59:N3:120:LYS:H	59:N3:137:VAL:CG2	2.38	0.41
61:N5:88:MET:HG2	61:N5:120:LYS:H	2.09	0.41
61:N5:27:ARG:H	61:N5:27:ARG:HG2	2.29	0.41
62:N6:15:ALA:O	62:N6:18:ALA:HB3	3.15	0.41
62:N6:48:LEU:HD23	62:N6:49:PRO:HD2	2.01	0.41
66:O0:28:LYS:HD2	36:5:1730:G:C8	235.11	0.41
66:O0:54:SER:HA	66:O0:57:GLU:HB2	2.42	0.41
76:Q0:77:ILE:O	76:Q0:78:ILE:HG23	4.86	0.41
79:Q3:37:TYR:CD2	79:Q3:37:TYR:N	3.11	0.41
2:S0:114:SER:O	2:S0:114:SER:OG	2.81	0.41
2:S0:131:GLN:NE2	2:S0:134:LYS:HE2	4.35	0.41
2:S0:187:ALA:O	2:S0:188:LEU:HD22	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:S1:48:VAL:CG1	3:S1:61:LEU:HD21	2.48	0.41
4:S2:115:ILE:HD13	4:S2:208:GLU:OE1	3.16	0.41
4:S2:214:ALA:O	4:S2:218:ILE:HG13	2.19	0.41
4:S2:35:TRP:CD1	4:S2:35:TRP:C	2.92	0.41
4:S2:77:GLN:HG3	4:S2:77:GLN:H	2.72	0.41
6:S4:60:GLU:OE1	26:D4:20:ARG:NH1	3.35	0.41
8:S6:97:VAL:HG22	8:S6:98:ARG:O	2.20	0.41
9:S7:27:LEU:C	9:S7:29:ASN:H	2.86	0.41
9:S7:77:LEU:HD23	9:S7:77:LEU:HA	2.04	0.41
10:S8:72:ILE:HG21	10:S8:112:TRP:CZ2	2.55	0.41
10:S8:196:LEU:HA	10:S8:196:LEU:HD23	4.22	0.41
10:S8:31:ARG:HH22	10:S8:48:THR:HA	1.85	0.41
10:S8:52:ASN:OD1	10:S8:52:ASN:N	3.96	0.41
11:S9:135:ALA:HA	11:S9:139:GLN:O	3.19	0.41
11:S9:134:ILE:HG12	11:S9:135:ALA:N	2.34	0.41
35:SM:27:LYS:HD2	48:M1:68:HIS:NE2	4.54	0.41
34:SR:168:THR:HG23	34:SR:181:TRP:O	2.20	0.41
36:1:1074:U:O2'	36:1:1075:A:H2'	2.20	0.41
36:1:1230:G:O6	36:1:1231:A:N6	2.53	0.41
36:1:1325:U:H5''	36:1:1325:U:H6	1.85	0.41
36:1:1343:A:H2'	36:1:1344:G:H8	1.85	0.41
36:1:422:A:C2	36:1:2363:A:H4'	2.55	0.41
36:1:2369:G:H2'	36:1:2370:G:O4'	2.20	0.41
36:1:2374:C:C4	36:1:2941:A:C4	3.08	0.41
36:1:2504:U:O5'	36:1:2504:U:H6	2.03	0.41
36:1:2553:U:H4'	36:1:2554:A:OP2	2.21	0.41
36:1:3133:C:H2'	36:1:3134:A:O4'	2.21	0.41
36:1:3375:A:H5''	36:1:3378:C:H5	1.85	0.41
36:1:346:C:C4	36:1:348:A:C8	3.08	0.41
36:1:674:G:C6	36:1:789:A:C6	3.07	0.41
36:1:815:G:C6	36:1:906:A:C4	3.07	0.41
1:2:1666:U:C4	1:2:1736:G:C2	3.08	0.41
1:2:1798:U:C4	28:D6:38:ARG:NH2	2.88	0.41
1:2:478:A:C2	1:2:479:C:C2	3.08	0.41
1:2:593:U:OP2	11:S9:39:LYS:HB2	2.20	0.41
1:2:68:A:OP2	8:S6:171:LYS:NZ	2.53	0.41
38:4:68:G:OP2	86:O7:104:OHX:N6	2.52	0.41
38:4:79:A:H2'	38:4:80:A:O2'	2.20	0.41
36:5:1569:U:H5'	36:5:1570:U:H6	1.85	0.41
36:5:1622:U:H2'	36:5:1623:G:O4'	2.21	0.41
36:5:2259:A:C5	36:5:2260:U:C5	3.08	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:2419:A:H1'	36:5:2804:A:O4'	2.20	0.41
36:5:2438:A:C2	36:5:2439:A:C8	3.08	0.41
36:5:271:C:H2'	36:5:272:G:O4'	2.20	0.41
40:L3:343:TYR:CE1	36:5:3099:C:H4'	266.86	0.41
36:5:1940:G:N2	36:5:3362:A:C8	2.89	0.41
36:5:961:C:N3	86:5:4173:OHX:N4	2.68	0.41
1:6:1046:G:C2	1:6:1073:G:C2	3.08	0.41
1:6:1039:A:N6	1:6:1091:A:C2	2.88	0.41
1:6:1103:U:O2'	1:6:1104:U:H5'	2.20	0.41
1:6:1185:U:C2	1:6:1458:G:C8	3.08	0.41
28:D6:5:ARG:HD3	1:6:1796:C:O4'	340.71	0.41
86:6:2059:OHX:N2	86:6:2147:OHX:N4	2.68	0.41
1:6:273:G:O2'	1:6:274:G:H5'	2.20	0.41
6:S4:3:ARG:HG2	1:6:399:A:H4'	320.93	0.41
1:6:730:G:C5	1:6:731:C:C4	3.08	0.41
37:7:49:G:N3	37:7:50:U:H5	2.18	0.41
73:O7:21:ARG:HG2	38:8:103:G:H4'	104.30	0.41
38:8:15:G:O2'	38:8:16:G:H5'	2.19	0.41
73:O7:60:GLY:N	38:8:42:G:OP1	87.98	0.41
38:8:80:A:H2	38:8:83:C:H41	1.68	0.41
13:C1:131:ILE:HA	13:C1:131:ILE:HD13	1.83	0.41
15:C3:119:GLU:HA	15:C3:122:ILE:HD12	2.02	0.41
17:C5:123:TYR:HD2	1:6:1458:G:O2'	354.07	0.41
17:C5:15:HIS:NE2	17:C5:16:SER:O	2.54	0.41
18:C6:38:LEU:HA	18:C6:38:LEU:HD23	2.25	0.41
18:C6:50:GLU:N	18:C6:51:PRO:HD2	2.44	0.41
18:C6:52:LEU:HA	18:C6:60:PHE:CE1	3.24	0.41
7:S5:25:LEU:HD12	18:C6:61:SER:OG	3.15	0.41
18:C6:47:LYS:HE3	18:C6:79:TYR:OH	2.20	0.41
24:D2:79:PHE:O	24:D2:124:LYS:HA	2.42	0.41
24:D2:38:LEU:HA	24:D2:38:LEU:HD23	2.00	0.41
24:D2:51:GLU:O	24:D2:61:ILE:HA	2.20	0.41
24:D2:75:ILE:HA	24:D2:75:ILE:HD13	2.09	0.41
25:D3:71:CYS:HB3	25:D3:85:ALA:O	2.30	0.41
28:D6:23:CYS:C	28:D6:25:ASN:N	3.15	0.41
24:D2:62:VAL:HB	29:D7:8:LEU:HD21	2.03	0.41
30:D8:52:ASP:N	30:D8:52:ASP:OD2	4.05	0.41
39:L2:102:LEU:HD22	39:L2:107:VAL:HG12	4.00	0.41
39:L2:27:ALA:HB3	39:L2:128:ARG:HH22	2.19	0.41
36:1:2943:G:N7	40:L3:2:SER:HB3	2.36	0.41
40:L3:345:ASN:OD1	40:L3:347:SER:HB2	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:L4:138:ARG:HH21	41:L4:240:PRO:HB2	2.09	0.41
41:L4:329:PRO:HB2	41:L4:330:TYR:H	4.16	0.41
41:L4:337:GLU:O	41:L4:339:LEU:N	2.64	0.41
42:L5:110:LEU:O	42:L5:116:ASP:HB3	4.78	0.41
42:L5:108:ARG:O	42:L5:111:GLN:HB3	2.20	0.41
42:L5:244:HIS:HA	42:L5:247:ILE:HD12	2.02	0.41
44:L7:120:THR:C	44:L7:122:ALA:H	2.72	0.41
46:L9:155:SER:O	46:L9:158:ALA:HB3	2.20	0.41
47:M0:17:TYR:CE1	47:M0:98:ARG:HD3	2.83	0.41
49:M3:119:TYR:HA	49:M3:145:PHE:CE1	4.00	0.41
49:M3:42:ARG:O	49:M3:46:ILE:HB	2.58	0.41
45:L8:71:VAL:HB	51:M5:21:PHE:CE1	2.55	0.41
51:M5:35:VAL:HA	51:M5:65:ARG:HD3	3.05	0.41
51:M5:76:PRO:O	51:M5:78:GLY:N	2.53	0.41
52:M6:192:LYS:O	52:M6:195:ALA:HB3	2.20	0.41
52:M6:33:ILE:HD12	52:M6:33:ILE:HG23	4.43	0.41
55:M9:84:THR:C	55:M9:86:GLU:N	2.74	0.41
56:N0:30:PHE:CZ	56:N0:103:VAL:HG21	2.73	0.41
58:N2:22:PRO:HB2	58:N2:28:PHE:HB2	2.65	0.41
59:N3:13:ILE:HD12	59:N3:85:TRP:CD2	2.54	0.41
63:N7:10:VAL:O	63:N7:83:THR:HB	2.96	0.41
63:N7:63:ALA:O	63:N7:67:LYS:HD3	2.21	0.41
68:O2:23:ASP:OD1	68:O2:23:ASP:N	2.53	0.41
68:O2:78:ASN:HA	68:O2:108:ILE:HD11	2.02	0.41
69:O3:52:VAL:HG22	69:O3:66:VAL:HG22	2.78	0.41
70:O4:106:LYS:O	70:O4:110:GLU:HG3	2.20	0.41
71:O5:89:ARG:HD2	38:8:38:U:C4	68.22	0.41
73:O7:19:CYS:HB2	73:O7:34:CYS:HB2	2.02	0.41
75:O9:35:ILE:N	75:O9:35:ILE:HD12	4.47	0.41
79:Q3:3:LYS:HB3	79:Q3:3:LYS:HE2	2.42	0.41
2:S0:18:LEU:HD22	2:S0:18:LEU:HA	1.91	0.41
2:S0:28:ASN:OD1	2:S0:28:ASN:N	4.31	0.41
3:S1:78:ASP:N	3:S1:78:ASP:OD1	3.13	0.41
4:S2:61:LEU:HA	4:S2:62:PRO:HD2	1.98	0.41
6:S4:187:ARG:NH2	1:6:754:A:N7	375.29	0.41
7:S5:159:ALA:HB3	7:S5:225:ARG:HB3	3.99	0.41
8:S6:147:LEU:O	8:S6:148:SER:OG	2.26	0.41
10:S8:138:ASN:HB3	10:S8:141:ARG:HH12	1.86	0.41
11:S9:146:PHE:CZ	11:S9:149:ARG:CZ	3.03	0.41
11:S9:33:GLU:HG2	11:S9:33:GLU:H	4.22	0.41
34:SR:108:SER:H	34:SR:128:ASP:HB3	3.57	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:1052:U:H5''	36:1:1053:A:OP2	2.19	0.41
36:1:1112:A:H2'	36:1:1113:G:O4'	2.20	0.41
36:1:1517:G:H2'	36:1:1518:U:O4'	2.20	0.41
36:1:1813:A:N3	36:1:1813:A:H3'	2.35	0.41
36:1:1939:G:C5	36:1:1940:G:C8	3.08	0.41
36:1:2414:G:C6	36:1:2415:C:C4	3.09	0.41
36:1:3008:A:H2'	36:1:3009:G:C8	2.55	0.41
36:1:3237:U:H2'	36:1:3238:G:O4'	2.21	0.41
36:1:3305:A:C6	36:1:3306:U:C4	3.08	0.41
36:1:682:U:H5	41:L4:112:LYS:HD3	1.85	0.41
36:1:771:A:H2'	36:1:772:U:O4'	2.20	0.41
36:1:677:A:C8	36:1:786:A:C6	3.09	0.41
36:1:953:G:C8	36:1:1117:G:C8	3.08	0.41
1:2:1081:A:H2'	1:2:1083:G:N7	2.34	0.41
1:2:1450:U:H2'	1:2:1451:C:C6	2.55	0.41
1:2:1485:C:N3	1:2:1592:A:H1'	2.36	0.41
1:2:417:A:H4'	1:2:418:G:O5'	2.20	0.41
1:2:432:G:C5	1:2:433:C:C4	3.09	0.41
1:2:892:A:H2'	1:2:893:U:C6	2.55	0.41
36:5:1093:A:C2	36:5:1096:U:O2	2.74	0.41
68:O2:45:ARG:NH2	36:5:1366:A:O3'	200.20	0.41
36:5:1861:G:C6	36:5:1862:U:C4	3.08	0.41
36:5:1928:G:C6	36:5:1929:G:C4	3.08	0.41
36:5:2297:U:C2	36:5:2299:A:C6	3.09	0.41
36:5:2440:G:O2'	36:5:2441:A:OP1	2.32	0.41
36:5:2704:A:OP2	86:5:3892:OHX:N2	2.53	0.41
50:M4:121:MET:HE1	36:5:3215:A:H5'	274.80	0.41
73:O7:35:SER:OG	36:5:361:A:H5'	126.48	0.41
86:5:4029:OHX:N3	86:5:4232:OHX:N1	2.68	0.41
36:5:1839:A:OP2	86:5:4075:OHX:N6	2.52	0.41
1:6:1150:G:H5''	1:6:1151:A:O5'	2.20	0.41
1:6:1318:G:N7	86:6:2165:OHX:N5	2.68	0.41
1:6:1737:G:H2'	1:6:1738:U:O4'	2.19	0.41
1:6:763:G:H2'	1:6:764:U:H6	1.84	0.41
1:6:76:A:H2'	86:6:2192:OHX:N2	2.34	0.41
36:5:415:G:H1	38:8:8:C:N4	2.18	0.41
14:C2:73:LYS:NZ	33:E1:108:VAL:H	2.18	0.41
15:C3:105:ASN:ND2	15:C3:105:ASN:N	2.68	0.41
1:2:955:A:H5''	15:C3:10:GLY:HA3	2.03	0.41
16:C4:42:VAL:HG21	16:C4:66:ASP:CB	4.01	0.41
16:C4:48:VAL:HG22	16:C4:49:LYS:N	2.48	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:C8:36:LYS:HB2	20:C8:102:ALA:HA	2.01	0.41
20:C8:82:PRO:HB3	20:C8:84:TRP:CE2	2.55	0.41
15:C3:56:ASP:HA	29:D7:47:PHE:HB3	2.12	0.41
29:D7:53:ALA:HB1	29:D7:62:ILE:HD11	3.13	0.41
31:D9:19:ARG:HD3	31:D9:32:ARG:NH1	3.56	0.41
39:L2:214:GLY:O	39:L2:215:ASN:HB2	4.38	0.41
40:L3:194:TRP:CE2	40:L3:198:HIS:CE1	3.51	0.41
36:1:2989:U:O3'	40:L3:232:ARG:NH2	2.52	0.41
41:L4:191:LYS:HD3	41:L4:194:TYR:CZ	2.97	0.41
41:L4:208:VAL:HG23	41:L4:248:VAL:HG13	2.43	0.41
42:L5:55:PHE:CE1	42:L5:158:ARG:HG2	4.39	0.41
46:L9:109:ALA:O	46:L9:110:LYS:HB2	2.32	0.41
46:L9:90:MET:HA	46:L9:180:TYR:O	2.20	0.41
36:1:2646:C:H5''	47:M0:119:TRP:CD1	2.56	0.41
48:M1:80:LEU:HD23	48:M1:80:LEU:HA	1.79	0.41
50:M4:53:VAL:HA	50:M4:54:PRO:HD3	1.74	0.41
51:M5:51:LEU:HA	51:M5:51:LEU:HD23	1.80	0.41
52:M6:112:TYR:O	52:M6:114:LYS:N	3.18	0.41
36:1:3180:A:H2'	52:M6:167:TYR:HE1	1.86	0.41
44:L7:92:ILE:HD11	54:M8:4:ASP:HB2	2.01	0.41
54:M8:60:PRO:HG3	54:M8:144:ARG:CA	2.99	0.41
55:M9:156:ASN:HA	55:M9:159:ALA:HB3	2.78	0.41
55:M9:81:ARG:HG3	55:M9:88:ARG:NH1	2.35	0.41
62:N6:57:LEU:HD12	62:N6:59:VAL:HG12	5.04	0.41
62:N6:71:SER:HB3	62:N6:83:ASP:N	2.35	0.41
63:N7:87:LEU:HD12	63:N7:127:ASN:ND2	4.13	0.41
64:N8:47:LYS:HE3	64:N8:48:TYR:CE2	2.53	0.41
64:N8:96:LYS:O	64:N8:98:THR:N	2.52	0.41
36:1:1430:U:H2'	64:N8:9:ARG:NH2	2.35	0.41
65:N9:36:ASP:HA	65:N9:37:PRO:HD3	1.90	0.41
69:O3:13:HIS:ND1	69:O3:93:THR:HB	2.35	0.41
69:O3:72:THR:OG1	69:O3:73:ARG:HG2	2.60	0.41
72:O6:57:LEU:HD21	72:O6:73:ALA:HB2	2.02	0.41
73:O7:5:THR:HG23	36:5:1845:G:O2'	154.57	0.41
74:O8:69:LEU:HA	74:O8:70:PRO:HD3	1.94	0.41
76:Q0:96:CYS:O	76:Q0:98:LYS:N	2.53	0.41
78:Q2:35:LEU:HB3	78:Q2:40:LYS:HG2	2.02	0.41
3:S1:113:MET:HB2	3:S1:113:MET:HE2	4.72	0.41
6:S4:36:HIS:NE2	6:S4:88:ASP:OD2	2.53	0.41
7:S5:42:LEU:HD21	7:S5:45:LYS:HD2	2.02	0.41
7:S5:45:LYS:HD3	7:S5:45:LYS:HA	1.29	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:S6:11:GLY:O	8:S6:129:VAL:HG13	2.72	0.41
9:S7:169:PHE:O	9:S7:172:VAL:HB	3.36	0.41
9:S7:45:SER:OG	9:S7:46:ILE:N	2.54	0.41
11:S9:97:LEU:HA	11:S9:97:LEU:HD23	2.30	0.41
36:1:1488:G:C2	36:1:1489:A:C8	3.08	0.41
36:1:1538:G:N7	86:1:4132:OHX:N1	2.68	0.41
36:1:1675:G:N2	36:1:1773:C:C5	2.89	0.41
36:1:223:U:HO2'	36:1:224:C:P	2.44	0.41
36:1:2294:U:OP2	59:N3:71:LYS:HE2	2.21	0.41
36:1:3024:A:H5''	46:L9:96:HIS:CD2	2.55	0.41
36:1:3078:U:H4'	36:1:3079:U:O5'	2.21	0.41
36:1:3372:A:H2'	36:1:3373:U:O4'	2.19	0.41
86:1:4014:OHX:N6	86:1:4052:OHX:N2	2.68	0.41
36:1:590:G:C2	36:1:610:G:H2'	2.55	0.41
1:2:11:A:O2'	1:2:12:U:H5'	2.21	0.41
1:2:1362:U:H2'	1:2:1362:U:H6	1.68	0.41
1:2:1479:A:H2'	1:2:1480:G:C8	2.52	0.41
1:2:1492:A:N3	1:2:1493:A:H8	2.18	0.41
1:2:1603:U:H2'	1:2:1604:U:C6	2.55	0.41
1:2:1663:G:C2	1:2:1664:C:C2	3.08	0.41
1:2:56:U:H4'	1:2:57:G:C5'	2.50	0.41
1:2:834:G:C6	1:2:835:U:C4	3.08	0.41
37:3:42:A:H2'	37:3:43:U:H6	1.86	0.41
37:3:90:U:C5	37:3:91:G:N7	2.89	0.41
38:4:76:C:H2'	38:4:77:A:C8	2.55	0.41
36:5:1204:A:H2'	36:5:1205:A:H5'	2.03	0.41
36:5:1179:A:N3	36:5:1328:C:H1'	2.35	0.41
36:5:1338:C:H2'	36:5:1339:C:H6	1.85	0.41
36:5:2256:A:C8	36:5:2256:A:OP2	2.73	0.41
36:5:2403:G:N7	36:5:2870:C:H4'	2.35	0.41
36:5:3164:C:H2'	36:5:3164:C:OP2	2.19	0.41
36:5:2211:U:OP1	86:5:4218:OHX:N5	2.54	0.41
36:5:422:A:N3	36:5:2363:A:H4'	2.36	0.41
36:5:715:A:H4'	36:5:716:A:OP1	2.20	0.41
36:5:817:A:H4'	36:5:818:C:OP2	2.20	0.41
1:6:1160:A:H2'	1:6:1161:C:C5	2.55	0.41
1:6:1228:G:N3	1:6:1228:G:H2'	2.35	0.41
1:6:1406:A:H2'	1:6:1407:U:C6	2.55	0.41
1:6:1608:U:H2'	1:6:1609:U:H6	1.85	0.41
1:6:1147:A:N6	1:6:1629:G:O6	2.53	0.41
28:D6:5:ARG:HH12	1:6:1795:U:H3'	338.52	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:87:C:C4	1:6:88:U:C5	3.09	0.41
42:L5:260:PHE:CG	37:7:121:U:C5	320.25	0.41
71:O5:10:ARG:NH2	38:8:66:A:OP1	31.10	0.41
17:C5:18:ARG:NH1	20:C8:90:ASN:HD21	5.03	0.41
1:2:1549:C:OP1	17:C5:38:PRO:HA	2.21	0.41
18:C6:57:LEU:H	18:C6:57:LEU:HD12	1.85	0.41
18:C6:86:ALA:O	18:C6:90:VAL:HG13	2.20	0.41
21:C9:40:SER:OG	21:C9:96:ALA:HA	2.20	0.41
21:C9:34:VAL:HG23	21:C9:53:TRP:NE1	2.36	0.41
22:D0:41:ILE:HG13	22:D0:103:ILE:HD12	2.02	0.41
22:D0:29:THR:HG22	22:D0:86:ILE:HA	2.02	0.41
1:2:775:G:C5	26:D4:11:LYS:NZ	2.89	0.41
29:D7:81:ARG:HG2	29:D7:82:LYS:H	1.86	0.41
31:D9:53:ASN:HB2	31:D9:55:PHE:CE2	3.80	0.41
39:L2:177:LYS:HA	39:L2:178:PRO:HD3	2.24	0.41
39:L2:247:ARG:HG2	39:L2:247:ARG:H	1.54	0.41
40:L3:28:ARG:HH21	40:L3:30:LYS:NZ	3.32	0.41
43:L6:36:PRO:HA	43:L6:54:TYR:CD1	3.25	0.41
44:L7:98:LYS:O	44:L7:102:VAL:HG23	2.71	0.41
45:L8:109:LEU:HD13	45:L8:109:LEU:HA	4.13	0.41
45:L8:158:ASP:HA	45:L8:159:PRO:C	4.40	0.41
45:L8:214:LEU:HD12	45:L8:214:LEU:HA	1.85	0.41
47:M0:53:VAL:O	47:M0:164:LYS:N	2.73	0.41
47:M0:99:ILE:O	47:M0:120:GLY:HA2	2.86	0.41
48:M1:131:MET:HB3	48:M1:131:MET:HE3	2.54	0.41
51:M5:60:VAL:HG21	51:M5:113:LEU:HD21	2.03	0.41
51:M5:73:ARG:HA	51:M5:74:PRO:HD3	1.57	0.41
53:M7:87:SER:O	53:M7:88:VAL:C	2.59	0.41
55:M9:138:LEU:O	55:M9:138:LEU:HD23	2.20	0.41
58:N2:97:SER:HA	58:N2:103:TYR:HD1	1.85	0.41
59:N3:95:PHE:HE1	60:N4:22:VAL:HG11	1.85	0.41
62:N6:27:ARG:O	62:N6:31:LEU:HD12	2.44	0.41
62:N6:52:ARG:HA	62:N6:70:ILE:CG2	3.11	0.41
62:N6:56:VAL:HA	62:N6:107:THR:HG23	2.02	0.41
64:N8:79:TRP:CE3	64:N8:82:ILE:HD12	2.59	0.41
66:O0:100:ILE:HG13	66:O0:100:ILE:H	1.61	0.41
66:O0:34:LEU:HD23	66:O0:59:TYR:HB3	2.03	0.41
67:O1:71:LEU:HA	67:O1:71:LEU:HD23	1.55	0.41
68:O2:34:LYS:HG3	68:O2:35:GLN:N	2.52	0.41
71:O5:100:VAL:HG22	71:O5:104:GLN:HB3	2.03	0.41
71:O5:31:LEU:HD23	71:O5:41:LEU:HD21	4.46	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
71:O5:74:LYS:HD3	71:O5:75:TYR:CZ	2.56	0.41
73:O7:24:ARG:HB2	73:O7:26:SER:OG	4.04	0.41
75:O9:17:LYS:O	75:O9:20:ASN:N	3.12	0.41
79:Q3:16:VAL:HG21	36:5:1928:G:O2'	236.29	0.41
2:S0:154:GLU:O	2:S0:156:VAL:HG13	2.58	0.41
3:S1:180:THR:H	3:S1:183:GLN:HB2	5.42	0.41
3:S1:184:LEU:O	3:S1:188:LEU:HG	2.19	0.41
3:S1:206:PRO:HB2	3:S1:207:LEU:H	1.67	0.41
3:S1:36:SER:HB2	3:S1:231:LEU:HB3	2.01	0.41
4:S2:73:LEU:HA	4:S2:74:PRO:HD3	1.81	0.41
5:S3:108:LYS:O	5:S3:113:LEU:HB2	3.63	0.41
5:S3:202:LEU:HA	5:S3:202:LEU:HD13	2.28	0.41
1:2:294:C:H5'	6:S4:133:LYS:HE3	2.03	0.41
7:S5:113:ILE:HG23	7:S5:191:ALA:HB2	2.02	0.41
7:S5:129:PRO:O	7:S5:132:VAL:HB	2.99	0.41
8:S6:106:LEU:HG	8:S6:106:LEU:H	2.47	0.41
8:S6:56:ASN:HB2	8:S6:108:VAL:HG22	6.50	0.41
10:S8:34:ALA:O	10:S8:35:ASN:C	2.58	0.41
34:SR:78:ALA:O	34:SR:94:VAL:N	2.37	0.41
36:1:1209:G:H2'	36:1:1210:U:O4'	2.21	0.41
36:1:1473:G:H2'	36:1:1474:A:O4'	2.21	0.41
36:1:1670:C:H4'	36:1:1859:A:O3'	2.21	0.41
36:1:1697:A:C8	36:1:1698:C:C5	3.09	0.41
36:1:1815:U:O2'	36:1:1816:A:P	2.79	0.41
36:1:2359:C:O5'	36:1:2359:C:H6	2.02	0.41
36:1:2437:G:C5	36:1:2511:A:C2	3.09	0.41
36:1:287:G:O5'	36:1:287:G:H8	2.04	0.41
36:1:3275:U:H3'	36:1:3276:G:H5''	2.02	0.41
36:1:2248:C:OP2	86:1:3874:OHX:N3	2.54	0.41
36:1:776:U:O2	36:1:2720:G:N2	2.54	0.41
36:1:813:G:C2	36:1:814:U:C6	3.08	0.41
1:2:1154:G:N2	1:2:1155:G:H1'	2.36	0.41
1:2:1653:C:C2	1:2:1748:G:C2	3.08	0.41
1:2:685:A:O2'	1:2:686:C:OP1	2.34	0.41
1:2:779:U:O2'	1:2:780:A:H5'	2.20	0.41
37:3:115:G:H2'	37:3:116:C:C6	2.55	0.41
37:3:23:A:C6	37:3:24:A:C6	3.08	0.41
38:4:145:U:H2'	38:4:146:U:C6	2.56	0.41
38:4:79:A:H3'	38:4:80:A:H4'	2.02	0.41
36:5:1037:C:H2'	36:5:1038:C:C6	2.56	0.41
36:5:1114:U:H2'	36:5:1115:G:O4'	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:1312:C:H2'	36:5:1313:G:O4'	2.20	0.41
36:5:1460:A:H2'	36:5:1461:A:C8	2.55	0.41
36:5:1543:G:C6	36:5:1544:G:N7	2.89	0.41
36:5:1610:G:C6	36:5:1611:G:C6	3.08	0.41
36:5:189:G:C2	36:5:191:U:C4	3.09	0.41
36:5:2403:G:C2	36:5:2405:C:C4	3.09	0.41
36:5:2608:G:O2'	36:5:2609:A:H5'	2.20	0.41
36:5:3168:A:H2'	36:5:3169:U:O4'	2.20	0.41
36:5:3288:G:OP2	36:5:3288:G:H2'	2.20	0.41
36:5:1276:U:OP2	86:5:4000:OHX:N1	2.52	0.41
41:L4:335:ALA:HB1	36:5:579:G:OP2	279.40	0.41
36:5:680:G:N1	36:5:701:G:C6	2.89	0.41
36:5:891:G:OP1	86:5:3909:OHX:N6	2.54	0.41
1:6:1274:C:C4	1:6:1427:A:N7	2.88	0.41
8:S6:137:ARG:NH2	1:6:169:A:OP1	320.63	0.41
1:6:189:C:C2'	1:6:190:C:H5'	2.50	0.41
1:6:60:U:H5'	1:6:61:A:OP2	2.20	0.41
1:6:625:C:H2'	1:6:626:U:C6	2.55	0.41
1:6:808:U:H2'	1:6:809:A:C8	2.55	0.41
37:7:75:G:OP1	86:7:218:OHX:N5	2.54	0.41
37:7:76:A:OP2	86:7:218:OHX:N3	2.53	0.41
38:8:83:C:H4'	38:8:85:G:N2	2.35	0.41
13:C1:83:THR:HA	13:C1:111:VAL:H	2.22	0.41
1:2:975:C:H5'	15:C3:109:LYS:HE2	2.02	0.41
15:C3:84:ILE:HA	15:C3:85:PRO:HD3	2.29	0.41
18:C6:83:GLN:OE1	18:C6:119:ALA:HA	2.21	0.41
18:C6:93:HIS:HA	18:C6:97:VAL:CG2	2.50	0.41
19:C7:24:LEU:O	19:C7:25:THR:HG23	2.21	0.41
20:C8:136:GLN:H	20:C8:136:GLN:CD	2.24	0.41
20:C8:45:LEU:HA	20:C8:48:LYS:HG3	3.81	0.41
22:D0:105:GLN:HG3	22:D0:106:ILE:N	2.35	0.41
22:D0:118:VAL:HG22	22:D0:119:ALA:N	2.40	0.41
5:S3:11:LEU:CD1	22:D0:29:THR:HG23	3.24	0.41
22:D0:43:LYS:O	22:D0:47:GLN:N	3.38	0.41
24:D2:98:GLN:O	24:D2:99:PHE:HB3	3.71	0.41
25:D3:19:ARG:HG3	25:D3:23:ARG:HG2	2.03	0.41
25:D3:24:TRP:HE3	25:D3:30:LYS:HG3	3.40	0.41
28:D6:79:ILE:HD12	1:6:1794:A:H1'	331.14	0.41
32:E0:28:LYS:HZ1	1:6:542:A:N6	428.78	0.41
33:E1:105:TYR:CG	33:E1:118:ARG:HD3	2.56	0.41
33:E1:149:LYS:HD3	1:6:1235:C:O2'	435.16	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:822:G:H1'	39:L2:15:ILE:HG22	2.03	0.41
40:L3:139:GLN:OE1	40:L3:142:ALA:HB3	2.21	0.41
41:L4:136:LEU:HD23	41:L4:136:LEU:HA	1.72	0.41
42:L5:152:ARG:CG	42:L5:152:ARG:HH11	2.68	0.41
42:L5:88:ILE:HD11	42:L5:243:ALA:CB	3.20	0.41
44:L7:151:ARG:NH1	44:L7:244:ASN:O	3.17	0.41
44:L7:96:PRO:O	44:L7:99:PRO:HD2	2.42	0.41
46:L9:176:LEU:HD23	46:L9:176:LEU:HA	1.84	0.41
47:M0:138:VAL:HG21	47:M0:148:VAL:HG13	2.02	0.41
49:M3:101:ARG:HH21	49:M3:112:ASN:ND2	2.19	0.41
36:1:1307:G:H5'	52:M6:60:LYS:HE3	2.02	0.41
36:1:1306:G:C5	52:M6:62:THR:HA	2.55	0.41
53:M7:112:LEU:HA	53:M7:112:LEU:HD12	2.92	0.41
54:M8:139:ILE:C	54:M8:140:LEU:HD23	3.19	0.41
56:N0:1:MET:HE1	56:N0:32:SER:H	1.85	0.41
56:N0:42:TRP:HZ2	56:N0:58:ILE:HG12	3.25	0.41
62:N6:126:LEU:HB3	62:N6:127:GLU:H	4.07	0.41
62:N6:89:LYS:O	62:N6:92:GLY:N	2.86	0.41
72:O6:14:GLY:HA2	36:5:73:C:OP1	107.02	0.41
72:O6:26:ILE:H	72:O6:26:ILE:HG13	1.81	0.41
49:M3:174:ARG:HD3	72:O6:9:ILE:HD12	4.87	0.41
73:O7:18:LEU:HA	73:O7:25:ARG:HA	2.03	0.41
74:O8:5:ILE:HG23	74:O8:54:LEU:HD13	2.99	0.41
75:O9:9:ILE:HD12	75:O9:51:ILE:HG22	2.02	0.41
76:Q0:113:ARG:NH1	36:5:1298:C:O3'	291.41	0.41
77:Q1:2:ARG:HG2	77:Q1:4:LYS:HB3	2.03	0.41
78:Q2:65:THR:OG1	78:Q2:87:ARG:HD2	3.98	0.41
2:S0:180:GLU:HA	2:S0:183:ARG:HB2	2.43	0.41
3:S1:131:ASP:HB3	3:S1:180:THR:HG23	2.02	0.41
3:S1:124:ASN:HB3	3:S1:138:PHE:CD1	2.69	0.41
3:S1:29:TRP:HZ3	3:S1:45:LYS:HD2	1.84	0.41
3:S1:46:THR:OG1	3:S1:47:LEU:N	4.49	0.41
4:S2:49:LYS:HB3	4:S2:243:TYR:CE1	3.44	0.41
6:S4:125:LYS:HB2	6:S4:226:PHE:CE1	2.88	0.41
9:S7:104:ARG:HB3	1:6:742:U:O4'	353.65	0.41
11:S9:84:GLY:O	11:S9:107:ARG:HD3	2.69	0.41
35:SM:84:LYS:HG2	35:SM:86:ASN:N	2.32	0.41
36:1:1003:A:C2'	36:1:1004:U:H5'	2.50	0.41
36:1:115:A:OP2	36:1:115:A:H8	2.04	0.41
36:1:1278:A:HO2'	36:1:1279:C:P	2.41	0.41
36:1:1286:A:N3	36:1:1287:A:H1'	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:1383:G:H4'	41:L4:240:PRO:O	2.19	0.41
36:1:1394:A:H2'	36:1:1395:G:O4'	2.20	0.41
36:1:1491:A:H8	36:1:1491:A:O5'	2.04	0.41
36:1:1673:G:H2'	36:1:1674:G:H8	1.86	0.41
36:1:1681:U:O4	58:N2:81:LYS:HE3	2.21	0.41
36:1:1712:G:C6	36:1:1713:G:C6	3.09	0.41
36:1:175:C:H2'	36:1:176:G:O4'	2.21	0.41
36:1:1856:C:H2'	36:1:1857:C:C6	2.56	0.41
36:1:2517:U:H2'	36:1:2518:C:H6	1.85	0.41
36:1:279:U:H2'	36:1:280:U:C6	2.56	0.41
36:1:394:G:N1	36:1:397:A:OP2	2.54	0.41
36:1:527:A:H2'	36:1:528:U:O4'	2.21	0.41
36:1:559:A:N7	36:1:560:G:C8	2.89	0.41
36:1:597:G:N3	36:1:598:A:C8	2.89	0.41
36:1:966:U:C2	36:1:967:A:C8	3.08	0.41
36:1:8:C:H2'	36:1:9:U:O4'	2.20	0.41
1:2:1158:C:H6	1:2:1158:C:H2'	1.68	0.41
1:2:1248:C:H2'	1:2:1249:U:C6	2.56	0.41
1:2:1294:G:C2	1:2:1295:G:C8	3.09	0.41
1:2:1617:U:O2'	1:2:1618:C:H5'	2.21	0.41
1:2:1746:A:H3'	1:2:1747:G:H8	1.86	0.41
1:2:1756:A:H2'	1:2:1757:G:H8	1.85	0.41
1:2:462:G:C5	1:2:463:U:C5	3.09	0.41
37:3:38:U:H5''	37:3:38:U:H6	1.86	0.41
36:5:1363:A:OP2	86:5:4194:OHX:N3	2.54	0.41
36:5:1611:G:H2'	36:5:1612:A:C8	2.56	0.41
36:5:1641:U:O2'	36:5:1642:A:H3'	2.20	0.41
36:5:197:G:H2'	36:5:198:A:O4'	2.20	0.41
36:5:2291:A:C6	36:5:2292:U:C4	3.08	0.41
36:5:2428:U:O4	86:5:4211:OHX:N5	2.53	0.41
36:5:2507:C:HO2'	36:5:2508:U:P	2.38	0.41
36:5:2509:U:H2'	36:5:2510:U:H5'	2.01	0.41
36:5:2778:G:H2'	36:5:2779:A:H5'	2.02	0.41
36:5:2783:U:H2'	36:5:2783:U:O2	2.21	0.41
36:5:3231:U:H2'	36:5:3232:G:H8	1.85	0.41
36:5:3242:G:C6	36:5:3245:A:C2	3.08	0.41
36:5:764:U:H6	36:5:764:U:O5'	2.03	0.41
36:5:815:G:C6	36:5:906:A:C4	3.08	0.41
1:6:1231:U:H2'	1:6:1232:U:C6	2.56	0.41
30:D8:18:ARG:HH11	1:6:1616:G:H4'	364.09	0.41
1:6:1698:G:HO2'	1:6:1699:G:P	2.39	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:249:U:H3'	1:6:250:C:C5'	2.50	0.41
1:6:493:U:H2'	1:6:494:U:H5''	2.03	0.41
1:6:628:G:H8	1:6:628:G:O5'	2.04	0.41
9:S7:100:PRO:HA	1:6:639:U:H3	367.85	0.41
1:6:704:C:H2'	1:6:705:U:O4'	2.20	0.41
1:6:722:G:HO2'	1:6:723:G:H8	1.68	0.41
8:S6:169:TYR:CD1	1:6:72:A:N6	364.64	0.41
1:6:745:U:C2	1:6:807:A:C2	3.08	0.41
13:C1:34:TRP:CH2	13:C1:36:LYS:HD3	3.24	0.41
16:C4:37:GLU:HA	1:6:895:G:O2'	259.39	0.41
17:C5:54:ALA:O	17:C5:58:LYS:HG3	2.21	0.41
18:C6:82:ARG:HG2	18:C6:82:ARG:H	1.63	0.41
20:C8:6:GLN:NE2	20:C8:6:GLN:HA	2.34	0.41
21:C9:68:ARG:HB3	21:C9:69:LYS:H	3.52	0.41
22:D0:46:GLU:OE1	22:D0:46:GLU:HA	2.20	0.41
26:D4:89:TYR:O	26:D4:92:VAL:HB	2.21	0.41
7:S5:161:ASP:OD2	30:D8:42:ARG:HD2	4.44	0.41
39:L2:202:VAL:HB	36:5:2185:G:OP1	211.32	0.41
40:L3:221:THR:HB	40:L3:273:HIS:O	2.77	0.41
40:L3:84:VAL:HG22	40:L3:85:VAL:N	2.67	0.41
41:L4:10:SER:C	41:L4:12:THR:H	2.24	0.41
41:L4:209:TYR:CZ	41:L4:229:ASN:HB2	2.55	0.41
41:L4:304:GLN:O	41:L4:306:THR:N	2.45	0.41
36:1:598:A:H1'	41:L4:322:GLN:HE22	1.84	0.41
42:L5:196:ARG:O	42:L5:199:ILE:HB	2.71	0.41
42:L5:211:LEU:C	42:L5:213:ASP:H	2.24	0.41
42:L5:90:HIS:NE2	42:L5:229:ASP:OD2	3.67	0.41
44:L7:161:VAL:HG12	44:L7:162:PRO:O	2.20	0.41
36:1:2586:G:C5	45:L8:241:LYS:HB2	2.55	0.41
46:L9:106:LYS:HE3	46:L9:107:ASP:H	4.90	0.41
47:M0:98:ARG:CZ	47:M0:119:TRP:CZ3	3.03	0.41
49:M3:79:GLU:OE2	49:M3:101:ARG:NH2	3.39	0.41
49:M3:91:ARG:CZ	49:M3:97:VAL:HB	2.51	0.41
43:L6:158:TYR:CE1	50:M4:115:PHE:HA	2.56	0.41
53:M7:26:PHE:C	53:M7:26:PHE:CD1	2.94	0.41
53:M7:55:GLN:OE1	36:5:3299:A:H4'	161.52	0.41
53:M7:84:PRO:O	53:M7:88:VAL:HG23	2.21	0.41
55:M9:10:LEU:HD12	55:M9:10:LEU:HA	1.86	0.41
55:M9:96:ILE:HG22	55:M9:100:ARG:HG3	2.01	0.41
56:N0:144:LEU:HA	56:N0:144:LEU:HD23	2.36	0.41
58:N2:95:PHE:CD2	58:N2:95:PHE:C	2.93	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
60:N4:9:SER:O	60:N4:53:VAL:HG23	3.67	0.41
64:N8:19:LYS:HD2	64:N8:25:HIS:ND1	3.38	0.41
65:N9:18:ARG:H	65:N9:18:ARG:HG2	1.49	0.41
65:N9:23:LYS:HB3	65:N9:24:PRO:CD	3.02	0.41
69:O3:57:LYS:HB2	36:5:432:G:OP1	200.86	0.41
72:O6:4:LYS:HD2	72:O6:12:ASN:O	3.37	0.41
3:S1:22:ASP:HA	3:S1:23:PRO:HD3	1.93	0.41
3:S1:32:ILE:HB	3:S1:44:GLY:N	2.36	0.41
4:S2:148:LEU:HA	23:D1:4:ASP:OD2	2.61	0.41
4:S2:52:THR:HA	4:S2:72:LEU:HD12	2.03	0.41
7:S5:169:ASN:OD1	7:S5:169:ASN:N	2.48	0.41
7:S5:205:SER:O	7:S5:206:SER:HB2	3.57	0.41
7:S5:208:SER:HB3	7:S5:211:ILE:HG13	4.27	0.41
7:S5:225:ARG:NH1	30:D8:58:GLU:HG2	2.35	0.41
1:2:78:A:C8	8:S6:154:ARG:HG2	2.56	0.41
9:S7:64:VAL:HG12	9:S7:65:PRO:N	2.42	0.41
11:S9:65:LYS:HD3	11:S9:70:LEU:HD11	4.51	0.41
36:1:1246:G:N2	36:1:1265:U:H5	2.19	0.41
36:1:1269:U:H6	36:1:1272:C:H5	1.69	0.41
36:1:1577:G:H2'	36:1:1578:C:C1'	2.51	0.41
36:1:1599:G:H1	36:1:1608:C:N4	2.17	0.41
36:1:172:G:C6	36:1:173:G:C5	3.09	0.41
36:1:2247:G:H5''	36:1:2248:C:OP2	2.20	0.41
36:1:3204:C:H2'	36:1:3205:G:C8	2.56	0.41
36:1:383:G:N7	86:1:4089:OHX:N6	2.68	0.41
36:1:397:A:C2	36:1:399:A:C4	3.09	0.41
86:1:4050:OHX:N4	86:1:4160:OHX:N1	2.67	0.41
1:2:1225:U:O2	1:2:1230:A:O2'	2.38	0.41
1:2:1243:G:OP1	86:2:2046:OHX:N4	2.53	0.41
1:2:1380:U:H2'	1:2:1381:U:O4'	2.21	0.41
1:2:1542:G:H5''	21:C9:87:GLY:C	2.41	0.41
1:2:1739:C:H2'	1:2:1740:A:H8	1.85	0.41
1:2:337:G:H3'	13:C1:133:LYS:HB2	2.01	0.41
1:2:376:C:H2'	1:2:377:G:C8	2.55	0.41
1:2:377:G:O6	86:2:2077:OHX:N5	2.54	0.41
1:2:795:U:H5	1:2:796:A:C5	2.39	0.41
1:2:813:U:H4'	1:2:814:A:OP2	2.21	0.41
36:5:1239:C:H42	36:5:1249:G:H1	1.69	0.41
36:5:1348:U:C5	36:5:1355:A:C8	3.07	0.41
36:5:1521:G:C2	36:5:1522:U:C5	3.09	0.41
36:5:1692:U:O4	36:5:1693:C:N4	2.54	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:1741:A:H2'	36:5:1742:U:O4'	2.21	0.41
36:5:180:C:O2	36:5:237:G:C2	2.74	0.41
36:5:2117:A:H3'	36:5:2118:C:H6	1.86	0.41
39:L2:240:ALA:HA	36:5:2154:U:O3'	218.45	0.41
36:5:2325:G:H2'	36:5:2325:G:N3	2.36	0.41
36:5:2549:G:H2'	36:5:2549:G:H8	1.67	0.41
36:5:2888:U:C6	36:5:2911:A:N6	2.88	0.41
36:5:2943:G:O5'	36:5:2943:G:H8	2.02	0.41
36:5:3063:C:H2'	36:5:3064:U:C6	2.55	0.41
36:5:3000:A:C2	36:5:3149:G:C5	3.08	0.41
36:5:380:U:C2	36:5:390:G:N2	2.88	0.41
86:5:4049:OHX:N5	86:5:4194:OHX:N6	2.68	0.41
36:5:651:G:H2'	36:5:652:G:O4'	2.21	0.41
36:5:703:G:O2'	36:5:787:G:H4'	2.20	0.41
36:5:891:G:C6	36:5:892:U:C4	3.09	0.41
68:O2:33:ARG:NH1	36:5:944:C:H4'	161.58	0.41
36:5:971:G:O2'	36:5:972:A:H5'	2.21	0.41
1:6:1146:G:H2'	1:6:1147:A:O4'	2.20	0.41
1:6:1244:A:HO2'	1:6:1245:G:P	2.40	0.41
1:6:702:G:N7	86:6:2098:OHX:N4	2.68	0.41
1:6:452:A:H3'	1:6:453:U:C5	2.56	0.41
1:6:542:A:H8	1:6:543:C:H5'	1.85	0.41
1:6:555:A:O2'	1:6:556:A:H5'	2.21	0.41
1:6:86:A:H2'	1:6:87:C:H6	1.86	0.41
1:6:940:A:C4	1:6:941:A:C8	3.08	0.41
38:8:115:C:H2'	38:8:116:G:H5''	2.02	0.41
15:C3:26:PHE:HA	15:C3:27:LYS:HE3	2.02	0.41
17:C5:123:TYR:CE1	20:C8:122:HIS:HE1	4.55	0.41
17:C5:22:LEU:HA	17:C5:25:LEU:HB2	3.00	0.41
21:C9:131:ASP:O	21:C9:134:ARG:HB3	2.21	0.41
21:C9:88:VAL:O	1:6:1467:C:O2'	364.03	0.41
22:D0:37:VAL:HG13	22:D0:107:THR:HG22	5.02	0.41
22:D0:68:ARG:NH2	22:D0:70:THR:HG21	2.35	0.41
23:D1:5:LYS:NZ	23:D1:5:LYS:HB3	4.83	0.41
23:D1:69:LEU:HD23	23:D1:69:LEU:HA	2.31	0.41
28:D6:19:LYS:HG3	28:D6:20:PRO:HD2	2.01	0.41
31:D9:16:LYS:HG2	31:D9:16:LYS:H	2.32	0.41
39:L2:107:VAL:HG11	39:L2:111:THR:HG21	2.01	0.41
39:L2:207:VAL:HG12	36:5:2415:C:H5''	185.77	0.41
40:L3:99:LEU:O	36:5:3004:C:H4'	245.55	0.41
41:L4:174:ALA:O	41:L4:177:ASP:N	2.49	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:L4:93:MET:HE2	41:L4:94:CYS:H	4.10	0.41
41:L4:98:ARG:CZ	36:5:933:A:C2	138.98	0.41
42:L5:86:TYR:CG	42:L5:247:ILE:HG12	3.27	0.41
43:L6:34:LEU:O	43:L6:54:TYR:HE1	2.88	0.41
44:L7:210:PRO:CD	44:L7:243:MET:HG2	2.49	0.41
44:L7:41:ARG:O	44:L7:44:ILE:HB	2.67	0.41
45:L8:160:ILE:H	45:L8:160:ILE:HG12	1.53	0.41
46:L9:94:TYR:HB2	46:L9:95:ALA:H	1.62	0.41
47:M0:15:LYS:H	47:M0:15:LYS:HG2	1.67	0.41
49:M3:115:ARG:CZ	49:M3:147:ILE:HG12	2.51	0.41
49:M3:60:ALA:HA	49:M3:61:PRO:HD3	1.85	0.41
52:M6:110:PRO:HA	52:M6:113:ASP:CG	3.13	0.41
52:M6:167:TYR:O	52:M6:170:LYS:N	3.04	0.41
54:M8:8:LYS:HE2	36:5:950:G:OP1	200.19	0.41
55:M9:23:TRP:CH2	55:M9:25:ASP:HA	3.79	0.41
57:N1:132:PRO:O	57:N1:134:GLN:HG2	4.18	0.41
59:N3:13:ILE:HG12	59:N3:53:SER:CB	2.50	0.41
59:N3:82:ALA:HA	59:N3:95:PHE:H	1.86	0.41
61:N5:34:LEU:O	61:N5:34:LEU:HD13	2.38	0.41
64:N8:128:ARG:O	64:N8:129:PHE:CG	2.85	0.41
65:N9:31:SER:HA	36:5:748:U:H5''	200.32	0.41
36:1:1486:G:H21	70:O4:6:THR:HG22	1.86	0.41
63:N7:20:GLY:HA2	70:O4:89:ILE:HD13	2.42	0.41
63:N7:14:VAL:HG21	70:O4:90:ILE:HD11	2.02	0.41
72:O6:43:LEU:O	72:O6:47:ILE:HG13	2.28	0.41
73:O7:39:TYR:CG	73:O7:40:PRO:HA	2.68	0.41
74:O8:61:LYS:O	74:O8:64:LYS:HG3	5.74	0.41
2:S0:170:ILE:O	2:S0:174:TRP:HD1	2.49	0.41
2:S0:200:ASP:N	2:S0:200:ASP:OD1	2.61	0.41
2:S0:65:ALA:C	2:S0:67:ILE:H	3.16	0.41
3:S1:70:LEU:HD21	3:S1:79:HIS:NE2	2.36	0.41
3:S1:81:PHE:HB2	3:S1:82:ARG:H	1.70	0.41
4:S2:133:LYS:C	4:S2:135:SER:H	2.41	0.41
8:S6:27:PHE:CE1	8:S6:36:VAL:HG11	2.56	0.41
8:S6:48:TYR:CE2	8:S6:121:LEU:HD22	4.45	0.41
9:S7:141:ARG:HH21	9:S7:143:LEU:CD1	2.31	0.41
9:S7:84:LYS:HA	9:S7:84:LYS:HD3	1.92	0.41
10:S8:183:ILE:HG23	10:S8:185:GLU:OE1	5.86	0.41
36:1:1138:U:H2'	36:1:1139:G:O4'	2.20	0.41
36:1:1190:A:H3'	36:1:1190:A:N3	2.36	0.41
36:1:1211:U:H2'	36:1:1212:A:H8	1.78	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:1322:U:H2'	36:1:1323:G:C8	2.56	0.41
36:1:13:A:OP2	86:1:4200:OHX:N5	2.53	0.41
36:1:1389:G:N1	36:1:1419:A:N6	2.68	0.41
36:1:1482:A:H4'	36:1:1483:G:OP2	2.21	0.41
36:1:1506:A:C2	36:1:1510:G:N1	2.89	0.41
36:1:1792:C:H2'	36:1:1795:U:C5	2.56	0.41
36:1:1670:C:O2'	36:1:1860:G:OP1	2.27	0.41
36:1:2209:U:H2'	36:1:2209:U:OP2	2.21	0.41
36:1:2209:U:OP2	36:1:2209:U:C6	2.74	0.41
36:1:2278:C:C2	36:1:2307:G:C2	3.09	0.41
36:1:283:G:H2'	64:N8:61:PHE:CZ	2.55	0.41
36:1:346:C:C4	36:1:348:A:N7	2.89	0.41
36:1:812:G:C5	36:1:813:G:C8	3.09	0.41
36:1:996:A:C2	36:1:1054:A:C4	3.09	0.41
1:2:1140:G:H2'	1:2:1141:G:H8	1.85	0.41
1:2:1313:A:C2	1:2:1315:U:H4'	2.55	0.41
1:2:1418:G:N2	1:2:1419:G:C4	2.89	0.41
1:2:1641:C:H2'	1:2:1642:G:C8	2.56	0.41
1:2:269:G:C6	1:2:287:G:C6	3.09	0.41
1:2:516:G:O5'	1:2:516:G:H8	2.04	0.41
1:2:580:A:C6	1:2:583:C:C2	3.09	0.41
1:2:881:A:H2'	1:2:882:U:O4'	2.21	0.41
37:3:19:C:H2'	37:3:20:A:C8	2.47	0.41
37:3:85:G:O2'	44:L7:218:ARG:NH2	2.51	0.41
38:4:125:U:HO2'	38:4:126:A:P	2.44	0.41
38:4:149:A:H2'	38:4:150:G:C8	2.56	0.41
36:5:1208:U:H6	36:5:3115:C:H42	1.67	0.41
51:M5:105:ARG:NH1	36:5:1545:A:C5	133.45	0.41
36:5:1786:G:H2'	36:5:1787:A:C8	2.56	0.41
36:5:1802:C:H2'	36:5:1803:C:C6	2.55	0.41
36:5:2194:G:C6	36:5:2195:C:C4	3.08	0.41
36:5:2369:G:OP2	86:5:3902:OHX:N1	2.53	0.41
48:M1:68:HIS:CE1	36:5:2682:C:H5''	303.83	0.41
36:5:2904:U:H2'	36:5:2905:U:C6	2.55	0.41
36:5:2890:A:N1	36:5:2913:C:N3	2.68	0.41
69:O3:60:ARG:NH1	36:5:3275:U:O2'	219.82	0.41
36:5:865:U:C5	36:5:866:A:N7	2.89	0.41
36:5:96:G:H2'	36:5:97:U:O4'	2.21	0.41
1:6:1180:C:C4	1:6:1181:U:C4	3.09	0.41
1:6:1382:A:C4	1:6:1383:G:N7	2.88	0.41
1:6:1699:G:N2	1:6:1702:A:O4'	2.54	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:6:1720:G:O6	86:6:2093:OHX:N4	2.53	0.41
32:E0:17:GLN:OE1	1:6:563:U:H4'	384.77	0.41
38:8:83:C:H4'	38:8:85:G:C2	2.56	0.41
13:C1:105:LYS:HD2	1:6:306:U:OP2	321.69	0.41
13:C1:132:SER:OG	13:C1:135:VAL:HB	2.45	0.41
13:C1:23:PRO:C	13:C1:25:VAL:N	3.06	0.41
13:C1:74:THR:HG22	13:C1:122:ILE:HG13	2.03	0.41
14:C2:52:LEU:HD11	14:C2:60:VAL:HG21	2.03	0.41
14:C2:88:LEU:HD12	14:C2:88:LEU:HA	2.20	0.41
15:C3:55:ARG:HD2	15:C3:56:ASP:OD2	2.20	0.41
2:S0:50:VAL:HG22	19:C7:109:LEU:HD21	2.03	0.41
19:C7:20:TYR:CE1	19:C7:38:ILE:HD11	2.55	0.41
20:C8:26:ILE:O	20:C8:31:ALA:HB2	2.31	0.41
15:C3:20:ARG:CZ	24:D2:56:HIS:HB3	2.51	0.41
25:D3:41:SER:OG	25:D3:43:PHE:O	3.19	0.41
26:D4:22:GLN:HB2	26:D4:72:PHE:CZ	2.56	0.41
1:2:1597:A:C8	31:D9:14:TYR:CD2	3.08	0.41
39:L2:181:LYS:NZ	36:5:860:G:P	214.40	0.41
39:L2:21:ARG:CZ	39:L2:22:LEU:HD21	2.51	0.41
41:L4:20:LEU:HD23	41:L4:20:LEU:HA	3.04	0.41
41:L4:20:LEU:HD22	41:L4:256:THR:CG2	3.66	0.41
41:L4:26:PHE:CD1	41:L4:130:ALA:HB2	3.49	0.41
41:L4:283:THR:HB	41:L4:285:ASP:H	4.03	0.41
42:L5:122:VAL:HG23	42:L5:123:GLU:H	2.69	0.41
43:L6:78:ARG:NH1	43:L6:106:PHE:HB2	2.35	0.41
44:L7:116:PHE:CZ	44:L7:144:ILE:HG23	2.60	0.41
45:L8:187:GLY:O	45:L8:191:ASN:N	2.53	0.41
45:L8:78:PHE:CD1	45:L8:78:PHE:N	2.88	0.41
46:L9:110:LYS:HE3	46:L9:110:LYS:HB2	4.70	0.41
46:L9:124:ARG:HD2	46:L9:124:ARG:H	4.38	0.41
47:M0:169:LYS:O	47:M0:177:ASP:HA	2.63	0.41
49:M3:124:ILE:HG12	49:M3:124:ILE:O	2.17	0.41
50:M4:21:VAL:HG12	50:M4:65:LEU:HD23	2.30	0.41
50:M4:39:ILE:HG13	50:M4:44:VAL:CA	2.51	0.41
54:M8:23:ASN:C	54:M8:23:ASN:OD1	2.58	0.41
55:M9:105:LEU:HA	55:M9:105:LEU:HD22	4.32	0.41
57:N1:156:TYR:CG	57:N1:157:GLU:N	3.23	0.41
58:N2:77:LYS:HG2	58:N2:81:LYS:HE3	4.80	0.41
60:N4:55:PHE:CD1	60:N4:55:PHE:C	2.94	0.41
61:N5:65:GLN:O	61:N5:85:GLN:N	2.72	0.41
62:N6:40:ARG:HD2	62:N6:46:LYS:HA	2.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
62:N6:99:LEU:HD23	62:N6:99:LEU:HA	1.71	0.41
63:N7:36:HIS:HA	63:N7:38:PHE:CE1	3.16	0.41
66:O0:46:ALA:HB3	66:O0:73:GLY:HA2	2.02	0.41
68:O2:100:ILE:O	68:O2:105:ARG:HD2	2.21	0.41
70:O4:32:ALA:O	70:O4:33:GLN:HG3	2.20	0.41
66:O0:51:LEU:HD22	70:O4:91:ARG:CZ	2.51	0.41
72:O6:17:VAL:HG12	72:O6:18:THR:O	2.21	0.41
74:O8:78:LEU:HD13	74:O8:78:LEU:HA	1.88	0.41
2:S0:102:PHE:HZ	2:S0:107:PHE:HE1	1.68	0.41
2:S0:108:THR:HA	4:S2:64:LYS:HE3	2.02	0.41
3:S1:158:SER:O	3:S1:162:ARG:HG3	2.21	0.41
5:S3:101:GLN:O	5:S3:105:MET:N	3.22	0.41
6:S4:136:VAL:HG13	6:S4:149:TYR:CE1	2.56	0.41
6:S4:16:HIS:ND1	6:S4:16:HIS:O	3.80	0.41
6:S4:192:ILE:HD12	6:S4:238:LEU:HD13	2.13	0.41
1:2:448:C:H5'	6:S4:29:PRO:HG3	2.02	0.41
7:S5:34:GLN:O	7:S5:37:GLN:HB2	2.21	0.41
7:S5:63:GLN:HB2	7:S5:89:ILE:HG13	2.01	0.41
8:S6:202:ARG:O	8:S6:205:ALA:HB3	2.21	0.41
9:S7:15:GLU:O	9:S7:19:GLN:HG2	2.21	0.41
10:S8:105:ASP:O	10:S8:107:THR:N	2.50	0.41
11:S9:110:GLN:OE1	11:S9:126:ARG:HG2	2.21	0.41
35:SM:84:LYS:C	35:SM:86:ASN:H	2.23	0.41
34:SR:203:THR:HG23	34:SR:245:PHE:CE2	2.56	0.41
36:1:1149:G:H21	36:1:1199:C:H41	1.69	0.41
36:1:1345:G:C2	36:1:1360:C:C2	3.09	0.41
36:1:1407:A:H5'	68:O2:32:TRP:HB3	2.03	0.41
36:1:1618:G:H4'	38:4:129:C:C1'	2.51	0.41
36:1:1514:G:C2	36:1:1841:A:C6	3.09	0.41
36:1:233:C:H2'	36:1:234:G:O4'	2.21	0.41
36:1:2403:G:N3	36:1:2405:C:C4	2.89	0.41
36:1:2417:U:O2'	36:1:2418:G:H5'	2.20	0.41
36:1:25:U:O4	86:1:3862:OHX:N3	2.53	0.41
36:1:274:G:H2'	36:1:275:U:O4'	2.21	0.41
36:1:2814:G:C2	36:1:2815:G:C4	3.08	0.41
36:1:2948:C:O2'	36:1:2949:U:H5'	2.21	0.41
36:1:3090:U:O2'	36:1:3091:A:H5'	2.21	0.41
36:1:2560:C:O2	86:1:3919:OHX:N1	2.53	0.41
36:1:566:G:N7	86:1:3997:OHX:N4	2.69	0.41
36:1:595:G:OP2	44:L7:30:ARG:NH2	2.54	0.41
1:2:1661:U:C2	1:2:1662:G:C8	3.09	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:15:U:H2'	1:2:16:G:O4'	2.21	0.41
1:2:481:A:H2'	1:2:482:U:O4'	2.21	0.41
1:2:602:U:H2'	1:2:603:U:C6	2.55	0.41
1:2:855:A:C2	1:2:857:U:H1'	2.56	0.41
36:5:1347:U:H2'	36:5:1355:A:H61	1.86	0.41
36:5:1556:C:H5''	36:5:2169:G:H22	1.85	0.41
36:5:176:G:H2'	36:5:177:U:H6	1.86	0.41
73:O7:4:GLY:HA3	36:5:2139:A:H62	164.56	0.41
36:5:2255:A:H5'	36:5:2261:G:N2	2.27	0.41
36:5:2515:A:C6	36:5:2516:U:C2	3.08	0.41
36:5:2623:G:C5	36:5:2624:G:N7	2.89	0.41
36:5:157:A:N6	36:5:264:G:O2'	2.48	0.41
36:5:3135:U:C5	36:5:3136:G:C5	3.08	0.41
36:5:3191:G:C6	36:5:3192:U:N3	2.89	0.41
50:M4:121:MET:HE1	36:5:3215:A:C5'	274.24	0.41
86:5:4004:OHX:N3	86:5:4195:OHX:N1	2.69	0.41
86:5:4029:OHX:N4	86:5:4232:OHX:N1	2.69	0.41
86:5:4060:OHX:N5	86:5:4138:OHX:N2	2.69	0.41
36:5:422:A:N6	36:5:423:A:C6	2.89	0.41
44:L7:60:ARG:NH2	36:5:516:A:O3'	303.84	0.41
56:N0:132:THR:OG1	36:5:534:U:OP1	353.41	0.41
36:5:567:G:H2'	36:5:568:G:C8	2.56	0.41
36:5:765:C:H4'	36:5:766:U:OP1	2.19	0.41
1:6:956:C:H5''	1:6:1072:C:O2'	2.21	0.41
1:6:1081:A:H1'	1:6:1082:C:C5	2.51	0.41
1:6:11:A:C2'	1:6:12:U:H5'	2.50	0.41
1:6:1391:A:H2'	1:6:1392:U:H6	1.84	0.41
1:6:1450:U:O5'	1:6:1450:U:H6	2.04	0.41
17:C5:44:ARG:CD	1:6:1555:A:H5''	397.84	0.41
1:6:1576:A:H2'	1:6:1577:A:O4'	2.20	0.41
1:6:1466:G:O2'	1:6:1602:C:OP1	2.36	0.41
1:6:145:A:H61	1:6:169:A:H62	1.69	0.41
1:6:624:G:H2'	1:6:625:C:O4'	2.21	0.41
1:6:641:G:H2'	1:6:642:G:C8	2.55	0.41
48:M1:152:HIS:HE1	37:7:55:A:N3	326.16	0.41
36:5:18:G:N2	38:8:142:C:C2	2.88	0.41
15:C3:34:ILE:O	15:C3:38:VAL:HG23	2.21	0.41
15:C3:83:GLU:HG2	15:C3:83:GLU:H	1.55	0.41
17:C5:128:HIS:HB2	35:SM:71:ASN:CG	2.40	0.41
20:C8:35:ILE:HB	20:C8:38:VAL:CG2	2.50	0.41
21:C9:108:LEU:HD23	21:C9:108:LEU:HA	2.18	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:C9:141:GLU:O	21:C9:143:ASP:N	4.28	0.41
21:C9:20:SER:OG	21:C9:21:PHE:N	3.91	0.41
21:C9:64:HIS:CE1	21:C9:68:ARG:HH21	2.39	0.41
21:C9:86:ARG:HG3	21:C9:90:PRO:O	3.47	0.41
22:D0:17:GLN:HG3	22:D0:18:GLN:HG3	9.04	0.41
22:D0:47:GLN:HG2	22:D0:47:GLN:O	2.20	0.41
23:D1:20:THR:O	23:D1:21:ASN:HB2	2.64	0.41
24:D2:126:LEU:HD23	24:D2:126:LEU:HA	1.92	0.41
24:D2:53:ILE:HG12	24:D2:60:LYS:HB2	2.03	0.41
24:D2:99:PHE:N	24:D2:99:PHE:CD1	2.89	0.41
26:D4:37:LYS:O	26:D4:41:ARG:N	2.96	0.41
39:L2:204:MET:H	39:L2:204:MET:HG2	1.70	0.41
39:L2:18:SER:O	39:L2:20:THR:HG23	5.72	0.41
40:L3:123:TYR:CE2	40:L3:124:LYS:HG3	2.56	0.41
40:L3:299:ASP:OD1	40:L3:301:THR:OG1	2.31	0.41
41:L4:222:VAL:HA	41:L4:223:PRO:HD3	1.63	0.41
41:L4:209:TYR:CE2	41:L4:229:ASN:HB2	2.55	0.41
41:L4:292:SER:OG	41:L4:295:ILE:N	2.53	0.41
42:L5:102:GLY:HA2	42:L5:105:ILE:HG22	3.24	0.41
42:L5:88:ILE:HD12	42:L5:240:TYR:CD1	4.05	0.41
46:L9:91:ARG:HD3	76:Q0:82:LEU:HD21	5.12	0.41
53:M7:29:THR:HA	53:M7:32:THR:CG2	2.49	0.41
54:M8:122:ILE:CG2	54:M8:126:GLN:HB2	2.50	0.41
36:1:1347:U:H3'	54:M8:38:ARG:HH21	1.86	0.41
55:M9:117:LYS:O	55:M9:120:TYR:HB3	2.62	0.41
55:M9:168:ALA:HB1	55:M9:172:ARG:NH2	2.36	0.41
56:N0:26:ARG:HH22	56:N0:28:ARG:HD2	2.33	0.41
57:N1:73:GLY:HA2	57:N1:89:LEU:O	2.20	0.41
58:N2:28:PHE:HZ	58:N2:33:TYR:CD2	2.39	0.41
59:N3:32:ARG:HB2	59:N3:32:ARG:CZ	2.51	0.41
65:N9:22:LYS:HE2	65:N9:22:LYS:HB2	4.56	0.41
68:O2:19:ARG:CD	68:O2:28:VAL:HG12	3.78	0.41
68:O2:18:LYS:HD3	68:O2:30:GLU:OE1	2.93	0.41
69:O3:24:ASN:ND2	69:O3:27:VAL:HG23	2.81	0.41
69:O3:41:ALA:HB3	69:O3:74:THR:HG22	2.27	0.41
74:O8:11:PHE:HD1	74:O8:12:LEU:HD22	4.00	0.41
78:Q2:61:LYS:HG2	78:Q2:61:LYS:O	2.21	0.41
2:S0:38:PHE:HB2	2:S0:49:ASN:OD1	2.20	0.41
3:S1:110:LEU:HD12	3:S1:110:LEU:H	1.86	0.41
3:S1:58:SER:O	3:S1:62:LYS:NZ	2.42	0.41
4:S2:238:SER:HG	4:S2:240:LEU:HD22	1.86	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2:579:A:C8	5:S3:178:ARG:HD3	2.56	0.41
9:S7:35:LYS:CG	9:S7:36:ALA:H	2.30	0.41
10:S8:2:GLY:N	1:6:393:C:OP2	292.69	0.41
10:S8:81:VAL:HA	10:S8:102:VAL:HG12	2.61	0.41
1:2:592:A:OP2	11:S9:39:LYS:HE3	2.21	0.41
11:S9:63:ASP:C	11:S9:65:LYS:H	4.21	0.41
34:SR:33:LEU:HD22	34:SR:302:PHE:CD1	3.82	0.41
36:1:1278:A:O2'	36:1:1279:C:C6	2.74	0.41
36:1:2177:G:N2	39:L2:118:GLU:OE2	2.54	0.41
36:1:2203:U:H1'	36:1:2240:G:N2	2.36	0.41
36:1:2617:U:H4'	36:1:2644:C:C5	2.56	0.41
36:1:2688:U:H4'	36:1:2689:A:O4'	2.20	0.41
36:1:2774:C:H6	36:1:2774:C:O5'	2.04	0.41
36:1:305:U:C5	36:1:2776:C:H1'	2.56	0.41
36:1:3071:U:H2'	36:1:3072:C:O4'	2.21	0.41
86:1:4027:OHX:N6	86:1:4040:OHX:N5	2.69	0.41
36:1:507:U:H2'	36:1:508:U:H6	1.86	0.41
36:1:685:G:OP1	49:M3:35:ARG:NH1	2.51	0.41
36:1:787:G:H2'	36:1:788:C:C6	2.56	0.41
36:1:901:G:H2'	36:1:902:G:C8	2.54	0.41
1:2:934:C:N3	1:2:1077:C:H4'	2.35	0.41
1:2:1291:G:H22	1:2:1324:G:H1	1.69	0.41
1:2:1545:A:C2	1:2:1546:G:C5	3.08	0.41
1:2:1656:U:C2	1:2:1658:G:H1'	2.55	0.41
1:2:546:U:C2	1:2:547:U:C6	3.09	0.41
1:2:748:U:O2	1:2:802:G:C2	2.74	0.41
1:2:894:U:H3	1:2:918:U:H3	1.69	0.41
1:2:970:A:H5''	1:2:970:A:C8	2.56	0.41
36:5:1412:G:H8	36:5:1412:G:O5'	2.03	0.41
36:5:1679:A:C2	36:5:1680:G:C8	3.09	0.41
36:5:287:G:N2	36:5:288:C:H1'	2.36	0.41
36:5:945:C:H2'	36:5:946:U:O4'	2.21	0.41
1:6:1450:U:OP2	86:6:2128:OHX:N4	2.54	0.41
1:6:1532:U:C4	1:6:1533:C:C5	3.09	0.41
17:C5:40:ARG:NH2	1:6:1553:G:O6	394.26	0.41
1:6:1582:U:C4	1:6:1614:A:C8	3.08	0.41
1:6:1791:A:C4	1:6:1793:G:C8	3.08	0.41
1:6:377:G:O6	86:6:2111:OHX:N4	2.53	0.41
86:6:2059:OHX:N5	86:6:2147:OHX:N3	2.69	0.41
1:6:390:G:N7	1:6:407:A:N1	2.69	0.41
1:6:542:A:C8	1:6:543:C:H5'	2.56	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:D3:19:ARG:HH12	1:6:610:G:H21	343.22	0.41
26:D4:10:ARG:HD2	1:6:778:G:O6	429.74	0.41
1:6:940:A:C6	1:6:941:A:C5	3.08	0.41
37:7:92:A:C5	37:7:93:C:H1'	2.56	0.41
37:7:8:G:H2'	37:7:9:C:O4'	2.21	0.41
12:C0:32:HIS:CD2	12:C0:33:GLU:HG2	6.17	0.41
12:C0:50:THR:HG21	12:C0:57:THR:OG1	2.21	0.41
15:C3:20:ARG:HG3	15:C3:20:ARG:HH11	4.18	0.41
15:C3:88:LEU:HD23	15:C3:88:LEU:HA	2.29	0.41
18:C6:13:LYS:NZ	18:C6:14:LYS:H	2.19	0.41
19:C7:45:ARG:NH2	19:C7:49:LYS:HE2	2.36	0.41
20:C8:53:ASP:O	20:C8:56:LYS:HB2	2.21	0.41
20:C8:62:THR:O	20:C8:66:LEU:HD23	5.10	0.41
21:C9:102:ARG:O	21:C9:105:LEU:N	3.39	0.41
21:C9:15:ILE:HD13	21:C9:60:SER:HB2	5.44	0.41
28:D6:49:ALA:HB1	28:D6:53:LEU:HD23	5.16	0.41
30:D8:15:VAL:O	30:D8:17:GLY:N	2.54	0.41
31:D9:14:TYR:CD2	31:D9:14:TYR:C	3.40	0.41
1:2:1233:G:O2'	33:E1:145:HIS:HB2	2.21	0.41
40:L3:53:MET:HE2	40:L3:77:THR:HG22	2.03	0.41
41:L4:39:PHE:CG	41:L4:242:ALA:HB2	2.95	0.41
41:L4:345:GLU:HB3	41:L4:346:LYS:H	3.99	0.41
36:1:933:A:C4	41:L4:98:ARG:NH2	2.89	0.41
42:L5:111:GLN:HA	42:L5:116:ASP:CG	2.42	0.41
42:L5:197:SER:O	42:L5:198:TYR:C	2.83	0.41
42:L5:258:LYS:HG3	42:L5:258:LYS:O	3.14	0.41
44:L7:207:LEU:HD23	44:L7:207:LEU:HA	2.07	0.41
44:L7:55:TYR:HE2	44:L7:141:TYR:CE2	2.38	0.41
44:L7:82:LYS:HE2	44:L7:82:LYS:HB3	1.80	0.41
45:L8:157:VAL:HG21	45:L8:163:VAL:HG21	2.86	0.41
46:L9:112:ILE:HG21	46:L9:161:LEU:HD11	2.03	0.41
46:L9:67:ALA:HA	46:L9:70:THR:HG22	2.03	0.41
48:M1:12:LEU:HD22	48:M1:13:LYS:N	4.93	0.41
50:M4:46:ILE:HD12	50:M4:58:ILE:HG21	3.65	0.41
51:M5:99:ARG:HH22	51:M5:166:ALA:HB3	2.37	0.41
52:M6:153:VAL:O	52:M6:156:LEU:HB2	3.00	0.41
52:M6:80:PHE:CE1	52:M6:84:LEU:HD12	3.56	0.41
53:M7:123:PRO:O	53:M7:143:PRO:HG2	2.34	0.41
53:M7:3:ARG:HA	53:M7:3:ARG:HH21	5.17	0.41
53:M7:94:LEU:HD12	53:M7:94:LEU:HA	2.09	0.41
54:M8:124:LEU:O	54:M8:127:LEU:HB3	2.60	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:M9:107:ALA:O	55:M9:111:ASP:N	2.50	0.41
57:N1:126:VAL:O	57:N1:127:GLN:HB3	2.29	0.41
59:N3:21:ALA:O	59:N3:36:ILE:HG13	2.20	0.41
63:N7:5:LEU:HB3	63:N7:6:LYS:H	4.02	0.41
70:O4:99:LYS:HA	70:O4:102:LYS:HB2	2.03	0.41
78:Q2:41:ARG:HH21	36:5:2785:A:H4'	162.81	0.41
2:S0:96:THR:HA	2:S0:97:PRO:HD3	1.85	0.41
3:S1:117:TRP:HA	3:S1:155:TYR:CE1	4.44	0.41
5:S3:55:THR:C	5:S3:57:ASP:H	2.24	0.41
5:S3:95:GLY:O	5:S3:126:VAL:HG13	2.21	0.41
6:S4:122:LYS:O	6:S4:162:ILE:N	2.50	0.41
7:S5:200:ASN:O	7:S5:204:GLY:HA2	2.21	0.41
8:S6:200:ALA:C	8:S6:202:ARG:H	2.71	0.41
10:S8:70:GLU:HG3	10:S8:112:TRP:CZ3	2.56	0.41
11:S9:133:HIS:O	11:S9:134:ILE:HD13	2.21	0.41
34:SR:165:ASP:O	34:SR:167:VAL:HG23	2.21	0.41
36:1:120:G:H4'	36:1:121:A:C8	2.56	0.40
36:1:1314:C:H5'	52:M6:17:GLY:HA3	2.02	0.40
36:1:1743:G:N3	36:1:1744:G:C8	2.89	0.40
36:1:1878:G:N3	36:1:1878:G:H3'	2.36	0.40
36:1:2430:A:H2'	36:1:2431:C:O4'	2.22	0.40
36:1:246:U:O2'	36:1:247:C:H5'	2.21	0.40
36:1:1580:A:H5'	36:1:2522:G:C5	2.56	0.40
36:1:3232:G:H1	36:1:3255:U:H3	1.69	0.40
36:1:3215:A:C8	36:1:3259:U:O2	2.74	0.40
36:1:326:U:O2'	36:1:327:A:H5'	2.21	0.40
36:1:3338:C:C2	36:1:3339:A:C8	3.09	0.40
36:1:395:A:H2'	36:1:396:A:C8	2.56	0.40
36:1:639:G:C2	36:1:651:G:C2	3.09	0.40
36:1:867:G:C5	36:1:868:C:C4	3.09	0.40
1:2:1019:A:OP2	15:C3:107:LYS:HE3	2.21	0.40
1:2:1230:A:C8	1:2:1256:A:C6	3.10	0.40
1:2:1242:A:O5'	1:2:1242:A:H8	2.03	0.40
1:2:223:U:H2'	1:2:224:C:C6	2.57	0.40
1:2:614:C:H42	1:2:1107:G:H1	1.69	0.40
1:2:741:C:O2'	1:2:742:U:O4'	2.39	0.40
1:2:743:U:O2	1:2:809:A:H1'	2.21	0.40
1:2:912:U:H4'	1:2:913:G:H2'	2.03	0.40
38:4:13:A:C6	38:4:14:C:C4	3.09	0.40
38:4:76:C:H2'	38:4:77:A:H8	1.85	0.40
36:5:1104:G:H2'	36:5:1105:A:H8	1.86	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:1242:G:H2'	36:5:1243:G:O4'	2.21	0.40
36:5:1816:A:H2'	36:5:1817:G:H5''	2.03	0.40
36:5:2288:G:OP1	86:5:3953:OHX:N3	2.54	0.40
36:5:2714:G:H8	36:5:2751:G:H2'	1.86	0.40
36:5:2998:U:C4	36:5:2999:U:C4	3.09	0.40
40:L3:18:PRO:HD2	36:5:3139:A:O4'	228.00	0.40
36:5:3157:U:H3'	36:5:3158:G:C5'	2.51	0.40
36:5:3207:U:H5'	36:5:3209:A:H2	1.86	0.40
36:5:3059:G:H4'	36:5:3373:U:O2'	2.20	0.40
36:5:572:A:C4	36:5:573:C:C6	3.09	0.40
36:5:767:U:C2	36:5:768:C:C6	3.09	0.40
36:5:830:A:H2'	36:5:831:G:O4'	2.21	0.40
1:6:1106:U:H2'	1:6:1107:G:H8	1.85	0.40
1:6:1684:U:H2'	1:6:1685:G:C8	2.56	0.40
1:6:1799:U:H4'	1:6:1800:A:C2'	2.47	0.40
1:6:545:A:H4'	1:6:546:U:H5'	2.03	0.40
1:6:853:G:H2'	1:6:854:U:C6	2.56	0.40
1:6:938:G:N2	1:6:942:G:C5	2.89	0.40
37:7:28:C:C4	37:7:29:C:C2	3.08	0.40
62:N6:51:ARG:HB2	38:8:71:A:H5''	39.14	0.40
14:C2:29:LYS:O	14:C2:33:ARG:HG2	2.21	0.40
1:2:951:A:H1'	15:C3:101:HIS:CG	2.56	0.40
16:C4:30:VAL:O	16:C4:39:ILE:HG12	3.36	0.40
17:C5:82:ASN:H	17:C5:82:ASN:ND2	2.19	0.40
18:C6:41:PRO:O	18:C6:42:GLU:HB3	2.21	0.40
18:C6:94:GLN:HG3	18:C6:95:LYS:N	2.54	0.40
19:C7:23:LYS:O	19:C7:24:LEU:HB2	2.22	0.40
19:C7:31:ASN:ND2	19:C7:55:THR:HG23	2.35	0.40
19:C7:60:ARG:NH2	19:C7:66:VAL:HG13	4.22	0.40
20:C8:62:THR:O	20:C8:65:GLU:N	2.54	0.40
20:C8:82:PRO:O	20:C8:83:ALA:HB3	2.21	0.40
21:C9:20:SER:O	21:C9:24:ARG:HB2	2.21	0.40
21:C9:45:MET:O	1:6:1477:G:H5''	373.16	0.40
22:D0:52:LYS:HE3	22:D0:52:LYS:HB2	4.35	0.40
26:D4:37:LYS:HD2	26:D4:57:VAL:HG13	2.03	0.40
28:D6:25:ASN:HB3	28:D6:77:CYS:SG	2.62	0.40
28:D6:85:ARG:HA	28:D6:85:ARG:HD3	1.68	0.40
36:L1:2183:A:OP1	39:L2:10:LYS:HD2	2.20	0.40
39:L2:42:ARG:HD2	39:L2:87:PHE:CD1	2.55	0.40
39:L2:58:LEU:HD23	39:L2:58:LEU:HA	2.20	0.40
39:L2:83:HIS:O	39:L2:84:THR:C	2.59	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:L3:113:GLU:CD	40:L3:167:ARG:HD3	2.46	0.40
36:1:3003:G:P	40:L3:26:ARG:HH22	2.44	0.40
40:L3:56:ILE:HD12	40:L3:359:ILE:HA	2.03	0.40
41:L4:316:ASN:O	41:L4:319:LYS:O	4.39	0.40
44:L7:111:ILE:HG22	44:L7:111:ILE:H	1.58	0.40
44:L7:231:ASN:OD1	44:L7:231:ASN:C	2.59	0.40
41:L4:330:TYR:OH	44:L7:49:ALA:HA	3.29	0.40
45:L8:240:ASN:OD1	45:L8:240:ASN:N	2.52	0.40
45:L8:72:PRO:HA	45:L8:73:PRO:HD3	1.80	0.40
46:L9:115:ARG:HG2	46:L9:123:ILE:HG13	2.02	0.40
47:M0:24:ARG:H	47:M0:24:ARG:HG3	3.22	0.40
47:M0:47:PRO:O	47:M0:172:GLY:N	2.50	0.40
48:M1:61:ARG:O	48:M1:62:ASN:HB2	2.21	0.40
36:1:96:G:H5'	49:M3:15:ARG:CZ	2.51	0.40
49:M3:190:LYS:HE2	49:M3:190:LYS:HB2	1.89	0.40
36:1:699:A:OP1	49:M3:68:LYS:HE3	2.21	0.40
51:M5:98:LEU:HD13	51:M5:98:LEU:HA	1.83	0.40
52:M6:78:ARG:HH11	52:M6:78:ARG:HG3	1.85	0.40
53:M7:46:LYS:HE3	53:M7:46:LYS:HB2	4.35	0.40
56:N0:1:MET:HB3	56:N0:1:MET:HE2	1.83	0.40
59:N3:45:ARG:O	59:N3:46:LEU:C	2.59	0.40
61:N5:80:ASN:O	61:N5:125:ARG:HG2	2.84	0.40
62:N6:102:SER:HG	62:N6:103:LYS:HZ2	1.57	0.40
62:N6:40:ARG:CD	62:N6:46:LYS:HA	2.51	0.40
72:O6:60:LEU:HD11	72:O6:68:ARG:NE	2.36	0.40
72:O6:79:SER:OG	72:O6:82:ARG:HG3	6.32	0.40
76:Q0:78:ILE:HG22	76:Q0:79:GLU:N	2.35	0.40
78:Q2:52:GLY:HA3	36:5:2421:U:O2'	175.55	0.40
79:Q3:36:ARG:HB2	79:Q3:48:LYS:HE2	2.02	0.40
5:S3:20:GLU:HA	12:C0:61:TRP:CE2	2.56	0.40
6:S4:233:LYS:HB3	6:S4:233:LYS:HE3	4.44	0.40
6:S4:23:LEU:HD13	6:S4:23:LEU:N	2.37	0.40
9:S7:114:ARG:C	9:S7:116:ARG:H	2.22	0.40
9:S7:157:LYS:O	9:S7:159:VAL:HG13	2.22	0.40
9:S7:166:LEU:HD12	9:S7:169:PHE:CD2	2.56	0.40
10:S8:48:THR:CG2	10:S8:54:LYS:HB2	2.51	0.40
34:SR:100:TYR:HA	34:SR:100:TYR:HD2	2.09	0.40
34:SR:36:ALA:HB2	34:SR:42:LEU:HG	2.04	0.40
36:1:1039:U:H2'	36:1:1040:A:H8	1.86	0.40
36:1:1720:U:P	55:M9:110:ARG:HH12	2.44	0.40
36:1:1449:A:C2	36:1:2356:A:C5	3.09	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:2617:U:H4'	36:1:2644:C:H5	1.87	0.40
36:1:2828:G:C4	36:1:2829:U:C6	3.09	0.40
36:1:2947:G:H2'	36:1:2948:C:C6	2.56	0.40
36:1:3009:G:O6	86:1:3894:OHX:N5	2.54	0.40
36:1:3195:U:O2'	36:1:3196:U:H5'	2.20	0.40
36:1:3362:A:H2'	36:1:3363:U:O4'	2.21	0.40
36:1:361:A:O4'	36:1:814:U:H4'	2.21	0.40
36:1:583:G:OP1	86:1:3860:OHX:N2	2.55	0.40
36:1:39:A:N6	36:1:43:A:OP2	2.54	0.40
36:1:556:U:H5'	36:1:557:A:C2	2.55	0.40
1:2:1133:A:H2'	1:2:1134:C:O4'	2.22	0.40
1:2:1383:G:C2'	1:2:1384:A:H5'	2.51	0.40
1:2:1489:U:H3'	1:2:1489:U:O2	2.21	0.40
1:2:1537:C:N4	86:2:2154:OHX:N4	2.69	0.40
1:2:1586:A:H2'	1:2:1587:A:O4'	2.21	0.40
1:2:1636:C:H4'	1:2:1637:C:H5''	2.03	0.40
1:2:1759:C:H2'	1:2:1760:G:O4'	2.21	0.40
86:2:2095:OHX:N1	86:2:2115:OHX:N2	2.69	0.40
1:2:366:A:O5'	1:2:366:A:H8	2.05	0.40
1:2:545:A:H61	1:2:594:A:C5'	2.33	0.40
1:2:555:A:C8	1:2:555:A:H3'	2.56	0.40
1:2:738:G:H2'	1:2:739:G:H8	1.86	0.40
36:5:1114:U:C4	36:5:1115:G:N7	2.88	0.40
36:5:1230:G:OP2	86:5:4000:OHX:N6	2.54	0.40
36:5:1345:G:C2	36:5:1360:C:C2	3.10	0.40
36:5:1734:G:H2'	36:5:1735:G:O4'	2.21	0.40
36:5:1932:A:H5'	36:5:1933:A:OP2	2.21	0.40
36:5:2442:G:C2	36:5:2443:A:C8	3.09	0.40
36:5:171:G:C2	36:5:248:U:O2	2.74	0.40
36:5:2608:G:C2	36:5:2609:A:C8	3.09	0.40
36:5:2732:G:H2'	36:5:2733:A:O4'	2.20	0.40
36:5:2756:C:H2'	36:5:2757:U:H6	1.85	0.40
36:5:299:G:H2'	36:5:300:G:O4'	2.21	0.40
36:5:3041:U:H2'	36:5:3042:U:C6	2.55	0.40
59:N3:92:PHE:CE1	36:5:3051:U:H1'	246.02	0.40
36:5:3071:U:H2'	36:5:3072:C:O4'	2.21	0.40
36:5:3330:A:C8	36:5:3330:A:H5''	2.56	0.40
36:5:345:G:H2'	38:8:25:G:O2'	2.21	0.40
36:5:562:C:C2	36:5:563:U:C5	3.09	0.40
41:L4:310:THR:HG23	36:5:609:G:H8	224.33	0.40
49:M3:14:PHE:CZ	36:5:665:A:H1'	132.02	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:677:A:H4'	36:5:678:G:O5'	2.21	0.40
1:6:1039:A:N3	1:6:1040:G:C8	2.89	0.40
1:6:1078:C:H2'	1:6:1079:U:C6	2.56	0.40
19:C7:2:GLY:N	1:6:1312:A:N7	395.13	0.40
1:6:1334:U:H2'	1:6:1335:U:O4'	2.21	0.40
1:6:1342:C:O2'	1:6:1343:U:H5'	2.21	0.40
1:6:272:U:O2'	1:6:273:G:OP2	2.34	0.40
1:6:300:A:H2'	1:6:301:A:C8	2.56	0.40
11:S9:133:HIS:CE1	1:6:512:A:O2'	447.44	0.40
1:6:540:G:O2'	1:6:542:A:H5'	2.21	0.40
1:6:981:U:H2'	1:6:982:U:H6	1.87	0.40
37:7:43:U:C4	37:7:44:C:C4	3.09	0.40
38:8:48:A:C6	38:8:62:C:C5	3.09	0.40
14:C2:136:ILE:O	14:C2:140:PHE:HB2	2.21	0.40
16:C4:11:SER:OG	16:C4:12:GLN:N	4.66	0.40
18:C6:10:PHE:HE1	18:C6:12:LYS:HE2	1.85	0.40
18:C6:38:LEU:O	18:C6:45:ARG:NE	2.54	0.40
20:C8:119:ILE:HA	20:C8:119:ILE:HD12	2.00	0.40
23:D1:87:ARG:O	29:D7:11:THR:OG1	2.33	0.40
24:D2:27:ILE:HG12	24:D2:61:ILE:HB	2.04	0.40
24:D2:7:LEU:HD11	24:D2:37:PHE:CD2	2.55	0.40
24:D2:89:TRP:O	24:D2:92:ASN:N	2.80	0.40
32:E0:35:TYR:CE1	32:E0:39:LEU:HD23	4.12	0.40
39:L2:46:LYS:O	39:L2:47:GLN:HB2	2.22	0.40
40:L3:328:ILE:HG23	40:L3:329:PRO:O	2.20	0.40
40:L3:95:THR:O	40:L3:98:GLY:N	2.51	0.40
36:1:933:A:N1	41:L4:98:ARG:NH1	2.68	0.40
44:L7:152:GLY:C	44:L7:153:PHE:HD2	2.61	0.40
45:L8:204:ARG:HG2	45:L8:206:GLU:HG3	2.02	0.40
45:L8:71:VAL:HA	45:L8:72:PRO:HD2	1.62	0.40
37:3:39:C:O2'	48:M1:43:GLN:HB3	2.22	0.40
49:M3:114:GLN:O	49:M3:117:LYS:HB2	2.21	0.40
50:M4:135:LEU:O	50:M4:136:ALA:HB2	2.21	0.40
51:M5:86:ASN:OD1	51:M5:86:ASN:N	2.53	0.40
54:M8:152:HIS:ND1	54:M8:162:ALA:O	3.07	0.40
54:M8:170:ARG:HH21	54:M8:171:LYS:HD2	4.46	0.40
55:M9:144:GLN:HG2	55:M9:144:GLN:O	2.21	0.40
55:M9:21:LYS:HA	55:M9:53:LYS:HD2	2.03	0.40
58:N2:22:PRO:HG3	58:N2:93:ILE:HG21	2.28	0.40
64:N8:42:ARG:HG2	64:N8:46:ASP:OD1	5.12	0.40
67:O1:10:ARG:NH1	67:O1:12:TYR:OH	3.75	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:1433:A:C2	68:O2:25:TYR:CD2	3.10	0.40
69:O3:50:ALA:HB1	69:O3:66:VAL:HG11	2.04	0.40
70:O4:105:VAL:HG12	70:O4:106:LYS:N	2.35	0.40
70:O4:30:LEU:O	70:O4:31:ARG:HB2	3.27	0.40
71:O5:95:PHE:H	36:5:135:C:HO2'	58.34	0.40
72:O6:51:SER:O	72:O6:55:ARG:HG3	2.83	0.40
72:O6:91:ASN:HA	72:O6:94:ILE:HG22	4.62	0.40
74:O8:36:LYS:HA	74:O8:37:PRO:HD3	2.41	0.40
74:O8:66:ILE:HG21	74:O8:77:ARG:NH2	2.36	0.40
75:O9:9:ILE:HA	75:O9:9:ILE:HD13	2.05	0.40
2:S0:178:ALA:O	2:S0:182:LEU:HG	2.37	0.40
3:S1:36:SER:HB3	3:S1:231:LEU:O	3.68	0.40
3:S1:70:LEU:HA	3:S1:73:LEU:HB2	4.47	0.40
5:S3:60:GLY:HA3	5:S3:65:ARG:HB3	3.17	0.40
7:S5:143:ARG:O	7:S5:162:VAL:HG13	2.21	0.40
7:S5:53:VAL:O	7:S5:55:ASP:N	2.55	0.40
7:S5:99:MET:O	7:S5:100:ASN:HB2	2.85	0.40
8:S6:3:LEU:HA	8:S6:109:LEU:O	2.21	0.40
8:S6:20:ASP:O	8:S6:24:ILE:HG13	2.22	0.40
8:S6:84:TYR:CZ	8:S6:86:PRO:HA	2.56	0.40
9:S7:114:ARG:C	9:S7:116:ARG:N	2.75	0.40
9:S7:62:VAL:HG13	9:S7:70:PHE:HD2	1.86	0.40
34:SR:231:MET:HB3	34:SR:232:TYR:H	1.59	0.40
34:SR:319:ASN:N	34:SR:319:ASN:OD1	2.98	0.40
36:1:1004:U:C4	36:1:1005:G:N7	2.90	0.40
36:1:1113:G:O2'	36:1:1369:A:N3	2.37	0.40
36:1:1465:A:H2'	36:1:1466:G:O4'	2.20	0.40
36:1:2104:A:H2'	36:1:2105:G:H8	1.87	0.40
36:1:2284:C:H5''	36:1:2285:C:OP2	2.21	0.40
36:1:420:G:N2	36:1:2385:G:OP2	2.49	0.40
36:1:2421:U:C4	36:1:2422:C:C5	3.09	0.40
36:1:2616:C:O5'	36:1:2616:C:H6	2.04	0.40
36:1:2619:G:H2'	36:1:2620:G:O4'	2.22	0.40
36:1:2925:C:H2'	36:1:2926:A:O4'	2.21	0.40
36:1:3046:A:C6	36:1:3047:U:C4	3.08	0.40
36:1:3174:A:C5	36:1:3175:U:C2	3.09	0.40
88:1:4211:BLS:O7	88:1:4211:BLS:H133	2.21	0.40
36:1:437:G:O2'	36:1:438:A:H5'	2.22	0.40
36:1:591:G:N2	43:L6:18:LEU:HB3	2.36	0.40
36:1:620:U:N3	36:1:622:A:C2	2.90	0.40
36:1:62:A:N6	36:1:63:A:N1	2.69	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:1:679:U:H2'	36:1:680:G:H8	1.87	0.40
1:2:1179:G:C6	1:2:1180:C:C4	3.09	0.40
1:2:1349:G:N3	1:2:1379:C:N4	2.67	0.40
1:2:1397:U:H2'	1:2:1398:U:H5''	2.03	0.40
1:2:1460:A:C4	35:SM:76:VAL:HG22	2.56	0.40
1:2:1524:A:C6	1:2:1525:A:C6	3.10	0.40
1:2:1483:A:C2	1:2:1607:G:H1'	2.57	0.40
1:2:1649:G:H2'	1:2:1650:U:C6	2.57	0.40
1:2:584:C:OP2	86:2:2026:OHX:N6	2.55	0.40
1:2:801:G:O6	86:2:2053:OHX:N3	2.54	0.40
1:2:422:G:N7	86:2:2107:OHX:N5	2.69	0.40
1:2:681:U:O5'	1:2:681:U:H6	2.04	0.40
1:2:772:G:N2	1:2:774:A:H1'	2.36	0.40
37:3:28:C:O3'	48:M1:135:GLY:HA2	2.22	0.40
36:5:1024:G:H3'	36:5:1024:G:N3	2.36	0.40
36:5:1223:A:H8	36:5:1223:A:P	2.44	0.40
36:5:1161:G:H1'	36:5:1365:G:N2	2.36	0.40
41:L4:183:LYS:HE3	36:5:1386:A:C5	119.60	0.40
36:5:1555:U:H5	36:5:1559:A:H61	1.69	0.40
36:5:1791:C:H2'	36:5:1792:C:C5	2.56	0.40
36:5:2662:G:H2'	36:5:2663:G:O4'	2.21	0.40
36:5:2784:G:H2'	36:5:2785:A:O4'	2.22	0.40
36:5:2785:A:OP1	86:5:4164:OHX:N4	2.55	0.40
36:5:3045:G:H2'	36:5:3046:A:O4'	2.22	0.40
36:5:3170:A:C6	36:5:3171:U:C4	3.09	0.40
1:6:1130:G:OP2	86:6:2112:OHX:N1	2.55	0.40
1:6:1211:A:N6	1:6:1453:G:C6	2.89	0.40
1:6:1244:A:H4'	1:6:1245:G:OP1	2.21	0.40
1:6:1270:G:C2	1:6:1271:G:C8	3.09	0.40
1:6:1269:U:H5'	1:6:1432:U:OP2	2.21	0.40
1:6:1620:C:C2	1:6:1621:U:C5	3.10	0.40
9:S7:96:ARG:HB3	1:6:856:A:N6	366.45	0.40
15:C3:20:ARG:NH2	1:6:862:A:OP1	357.02	0.40
28:D6:11:ASN:HB3	1:6:934:C:H6	333.14	0.40
1:6:957:G:C6	1:6:958:U:C4	3.08	0.40
37:7:5:G:C6	37:7:6:C:C4	3.10	0.40
12:C0:24:LYS:HG3	12:C0:25:LYS:N	3.19	0.40
14:C2:42:ALA:HB2	14:C2:124:LYS:HD2	2.67	0.40
14:C2:41:LEU:HD13	14:C2:43:ARG:HB3	2.03	0.40
1:2:1228:G:H5'	14:C2:45:LEU:HB2	2.03	0.40
15:C3:113:PHE:O	15:C3:117:LEU:HG	2.39	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:C6:37:THR:O	18:C6:38:LEU:HD23	2.21	0.40
20:C8:133:VAL:HG13	1:6:1545:A:H5''	353.29	0.40
20:C8:86:LEU:HD12	20:C8:99:HIS:CG	4.25	0.40
24:D2:75:ILE:H	24:D2:127:GLY:HA2	2.71	0.40
25:D3:11:SER:O	25:D3:15:LEU:HG	2.20	0.40
25:D3:44:GLY:N	25:D3:78:LYS:HE3	4.34	0.40
25:D3:51:GLY:O	25:D3:101:GLU:HA	2.74	0.40
26:D4:132:ARG:HG2	26:D4:133:ASN:OD1	2.94	0.40
29:D7:33:LEU:HD13	29:D7:46:VAL:HB	2.04	0.40
29:D7:6:ASP:CG	29:D7:9:HIS:HD1	2.43	0.40
30:D8:9:LEU:HB2	30:D8:34:GLU:OE1	2.21	0.40
36:1:2940:A:N7	40:L3:2:SER:O	2.54	0.40
40:L3:65:SER:O	40:L3:68:HIS:N	2.54	0.40
41:L4:191:LYS:C	41:L4:193:LYS:N	2.75	0.40
41:L4:23:PRO:HD2	41:L4:26:PHE:CD2	2.56	0.40
41:L4:320:ASN:HB3	41:L4:323:VAL:CG1	2.52	0.40
42:L5:53:VAL:HA	42:L5:61:ILE:O	2.21	0.40
44:L7:239:LEU:HB3	44:L7:240:VAL:H	1.80	0.40
45:L8:224:ASP:C	45:L8:226:TYR:H	2.25	0.40
46:L9:67:ALA:HA	46:L9:70:THR:CG2	2.51	0.40
47:M0:150:GLU:OE1	47:M0:154:ARG:NE	2.51	0.40
48:M1:117:ASP:HA	48:M1:118:PRO:HD2	1.97	0.40
48:M1:76:ALA:O	48:M1:79:ILE:N	3.00	0.40
36:1:77:A:H5'	49:M3:100:ARG:CZ	2.51	0.40
51:M5:204:LYS:O	86:5:3972:OHX:N4	123.82	0.40
52:M6:96:LYS:HE2	36:5:2384:A:C2	220.20	0.40
53:M7:32:THR:HA	53:M7:58:ILE:HG21	2.03	0.40
53:M7:67:ILE:HD11	53:M7:82:ARG:NH1	2.36	0.40
36:1:841:A:H5'	55:M9:125:LYS:O	2.21	0.40
56:N0:13:ARG:HD3	56:N0:51:VAL:HG13	4.67	0.40
56:N0:42:TRP:CH2	56:N0:56:GLY:HA3	2.56	0.40
59:N3:10:LYS:HE2	59:N3:13:ILE:HG21	5.46	0.40
59:N3:48:ARG:HG3	59:N3:48:ARG:HH11	1.86	0.40
60:N4:39:LEU:HD12	60:N4:44:LYS:CG	3.07	0.40
60:N4:42:GLN:HB3	60:N4:44:LYS:HG2	2.03	0.40
63:N7:15:ARG:NH1	63:N7:15:ARG:HG3	3.06	0.40
66:O0:24:THR:CG2	66:O0:91:SER:HB3	2.51	0.40
68:O2:123:LYS:HA	68:O2:126:LEU:CB	3.64	0.40
69:O3:51:TYR:O	69:O3:51:TYR:CG	2.74	0.40
70:O4:31:ARG:HG3	70:O4:32:ALA:N	2.94	0.40
70:O4:41:ARG:HB2	70:O4:50:ALA:HB1	2.02	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
71:O5:64:GLU:O	71:O5:68:GLN:HG3	2.22	0.40
72:O6:34:SER:O	72:O6:38:LYS:HD3	5.10	0.40
45:L8:230:LYS:NZ	72:O6:47:ILE:O	2.50	0.40
76:Q0:96:CYS:C	76:Q0:98:LYS:N	2.75	0.40
77:Q1:1:MET:O	77:Q1:1:MET:HG3	2.20	0.40
79:Q3:35:ALA:HB3	79:Q3:37:TYR:CE2	3.46	0.40
3:S1:63:GLY:HA2	3:S1:88:VAL:O	2.20	0.40
5:S3:134:CYS:HA	5:S3:187:LYS:O	2.27	0.40
5:S3:222:VAL:HG23	34:SR:191:ASP:O	2.21	0.40
5:S3:26:THR:HA	5:S3:34:TYR:CD1	2.56	0.40
6:S4:23:LEU:HD22	6:S4:23:LEU:N	2.50	0.40
6:S4:253:ASP:O	6:S4:257:ALA:N	2.54	0.40
6:S4:58:GLY:HA2	6:S4:61:VAL:HG23	2.75	0.40
7:S5:172:ILE:HA	7:S5:175:LEU:HD12	2.02	0.40
8:S6:141:ILE:HD12	8:S6:141:ILE:H	1.87	0.40
8:S6:211:LEU:HD13	8:S6:214:LYS:NZ	9.86	0.40
9:S7:30:SER:O	9:S7:34:LEU:HD12	2.21	0.40
9:S7:41:LEU:HD13	9:S7:70:PHE:CE1	2.56	0.40
11:S9:108:ARG:HE	11:S9:145:SER:HA	1.86	0.40
34:SR:249:ARG:NH1	34:SR:315:VAL:HG21	3.69	0.40
36:1:1139:G:C6	36:1:1140:G:N7	2.89	0.40
36:1:150:A:C5	36:1:151:A:C8	3.10	0.40
36:1:2957:G:C5	36:1:2976:A:C2	3.09	0.40
36:1:3163:A:C2'	36:1:3164:C:H5'	2.51	0.40
86:1:4050:OHX:N6	86:1:4160:OHX:N5	2.68	0.40
36:1:496:C:C2'	36:1:497:C:H5'	2.52	0.40
36:1:531:G:C2	36:1:532:A:C4	3.10	0.40
36:1:544:C:O2'	36:1:548:G:N2	2.55	0.40
1:2:1274:C:C5	35:SM:95:SER:HA	2.56	0.40
1:2:526:A:H2'	1:2:527:A:O4'	2.21	0.40
1:2:648:G:N1	1:2:649:U:C4	2.89	0.40
1:2:755:A:H2'	1:2:756:A:C8	2.55	0.40
1:2:856:A:N7	9:S7:97:ARG:HB2	2.37	0.40
37:3:62:U:O3'	42:L5:285:ARG:NH1	2.54	0.40
36:1:406:G:H1'	38:4:16:G:N2	2.36	0.40
38:4:79:A:O3'	38:4:80:A:H4'	2.21	0.40
36:5:1014:U:C2'	36:5:1015:U:H5'	2.51	0.40
36:5:1093:A:C2	36:5:1096:U:C2	3.09	0.40
36:5:1340:G:C4	36:5:1341:U:C5	3.09	0.40
36:5:1440:G:C6	36:5:1441:G:C5	3.09	0.40
36:5:1624:G:H2'	36:5:1625:A:C8	2.57	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:1714:A:H2	36:5:1727:G:N3	2.20	0.40
36:5:2268:U:O2	36:5:2269:U:N3	2.55	0.40
36:5:2275:A:H2'	36:5:2276:G:O4'	2.22	0.40
36:5:238:A:H2'	36:5:239:G:C8	2.56	0.40
36:5:3046:A:C5	36:5:3047:U:C5	3.09	0.40
36:5:3119:U:H5''	36:5:3120:C:OP2	2.21	0.40
36:5:3321:C:H2'	36:5:3322:A:C8	2.57	0.40
36:5:383:G:N2	36:5:385:A:H3'	2.35	0.40
86:5:4060:OHX:N5	86:5:4138:OHX:N6	2.69	0.40
36:5:597:G:H2'	36:5:598:A:C8	2.55	0.40
49:M3:35:ARG:NH1	36:5:685:G:P	82.77	0.40
1:6:1388:A:C6	1:6:1411:A:C6	3.09	0.40
1:6:152:U:C2	1:6:163:G:N2	2.89	0.40
1:6:48:G:C6	1:6:432:G:C2	3.09	0.40
1:6:551:G:C8	1:6:582:U:C5	3.09	0.40
1:6:799:A:H2'	1:6:800:U:O4'	2.20	0.40
24:D2:105:THR:HG22	1:6:804:A:N3	366.41	0.40
1:6:817:A:H2'	1:6:818:C:O4'	2.22	0.40
1:6:838:G:C6	1:6:839:U:C4	3.10	0.40
37:7:52:G:C2	37:7:53:U:C5	3.10	0.40
38:8:73:U:C5	38:8:74:U:C5	3.09	0.40
12:C0:31:LYS:HA	12:C0:37:THR:O	2.32	0.40
12:C0:57:THR:HG23	12:C0:66:TYR:CE1	2.56	0.40
15:C3:128:TYR:CE1	1:6:964:U:H5''	323.62	0.40
17:C5:130:ARG:NH2	35:SM:70:ASN:HB3	2.36	0.40
17:C5:35:LYS:HA	17:C5:35:LYS:HD2	1.89	0.40
17:C5:34:VAL:HA	17:C5:37:ALA:HB2	2.03	0.40
20:C8:26:ILE:O	20:C8:58:ALA:HB2	2.22	0.40
21:C9:123:ARG:HG2	21:C9:124:ILE:O	2.98	0.40
21:C9:135:ILE:H	21:C9:135:ILE:HG13	1.66	0.40
21:C9:86:ARG:CG	21:C9:86:ARG:HH11	2.34	0.40
22:D0:55:PRO:HA	22:D0:91:ILE:HG12	2.03	0.40
25:D3:102:VAL:HG23	25:D3:104:LEU:HD21	3.83	0.40
26:D4:96:LEU:H	26:D4:96:LEU:HG	2.71	0.40
28:D6:7:SER:OG	28:D6:10:ARG:HA	2.21	0.40
33:E1:130:VAL:HG11	33:E1:143:LYS:HG2	2.04	0.40
39:L2:134:VAL:HG12	39:L2:150:LEU:HA	2.02	0.40
40:L3:120:LYS:HA	40:L3:120:LYS:HD2	1.94	0.40
40:L3:59:ASP:CG	40:L3:357:LYS:HZ2	2.82	0.40
41:L4:31:ARG:O	41:L4:33:ASP:N	2.55	0.40
41:L4:99:MET:HG3	41:L4:102:PRO:HB3	2.03	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:L7:144:ILE:HD12	44:L7:189:ILE:HD12	2.03	0.40
44:L7:77:VAL:HG22	57:N1:139:ARG:HB2	4.31	0.40
46:L9:1:MET:HG3	46:L9:3:TYR:CZ	2.85	0.40
47:M0:98:ARG:CZ	47:M0:119:TRP:CH2	3.19	0.40
47:M0:81:GLY:C	47:M0:83:ASP:N	2.97	0.40
48:M1:113:GLY:O	48:M1:114:ILE:HB	2.36	0.40
48:M1:144:CYS:O	48:M1:146:GLY:N	2.53	0.40
48:M1:53:THR:HA	48:M1:59:ILE:O	2.22	0.40
36:1:2680:A:N3	48:M1:59:ILE:HD13	2.37	0.40
50:M4:104:ALA:HA	50:M4:107:GLU:HB2	2.90	0.40
51:M5:164:LEU:HD23	51:M5:164:LEU:HA	2.10	0.40
51:M5:171:SER:HB3	36:5:289:A:OP1	125.13	0.40
52:M6:138:LEU:HA	52:M6:138:LEU:HD12	2.15	0.40
52:M6:38:ALA:N	52:M6:39:GLU:OE1	3.17	0.40
40:L3:261:MET:SD	52:M6:64:PHE:HA	2.62	0.40
53:M7:111:LYS:O	53:M7:153:LYS:N	2.42	0.40
54:M8:170:ARG:HD2	64:N8:56:VAL:O	2.49	0.40
54:M8:87:VAL:O	54:M8:107:THR:HG23	2.21	0.40
55:M9:44:LEU:HA	55:M9:44:LEU:HD12	1.76	0.40
57:N1:80:VAL:HG13	57:N1:85:LEU:HG	2.37	0.40
59:N3:80:ARG:NE	59:N3:97:ASP:OD2	2.40	0.40
63:N7:87:LEU:HD12	63:N7:121:ARG:NH1	2.37	0.40
64:N8:19:LYS:HB3	64:N8:25:HIS:HB2	2.04	0.40
64:N8:52:TYR:HD2	64:N8:53:PHE:CD1	2.39	0.40
67:O1:70:ARG:HE	67:O1:102:LYS:NZ	4.53	0.40
71:O5:13:SER:O	71:O5:14:LYS:C	2.79	0.40
71:O5:85:THR:CG2	71:O5:87:ALA:HB3	3.35	0.40
73:O7:28:HIS:O	73:O7:32:LYS:N	2.54	0.40
79:Q3:83:ILE:HD13	79:Q3:83:ILE:HA	2.03	0.40
3:S1:181:LEU:HD13	3:S1:181:LEU:H	1.87	0.40
6:S4:131:LEU:CD1	6:S4:135:GLY:HA2	2.52	0.40
6:S4:139:VAL:HG13	6:S4:150:PRO:HB3	2.88	0.40
6:S4:65:LEU:HG	6:S4:70:VAL:HG11	2.03	0.40
6:S4:65:LEU:HD23	6:S4:70:VAL:HG13	2.49	0.40
6:S4:92:LEU:HD12	6:S4:95:THR:HG21	5.77	0.40
7:S5:143:ARG:HB2	7:S5:218:GLU:OE2	3.27	0.40
7:S5:194:LEU:HD23	7:S5:194:LEU:HA	4.52	0.40
7:S5:59:VAL:C	7:S5:61:TYR:N	2.74	0.40
7:S5:59:VAL:O	7:S5:61:TYR:N	2.51	0.40
7:S5:63:GLN:HE21	7:S5:86:GLN:HG3	1.87	0.40
8:S6:28:PHE:HZ	8:S6:104:PRO:HB3	2.30	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:S6:55:GLY:CA	8:S6:63:MET:HG3	3.58	0.40
9:S7:114:ARG:HH11	9:S7:114:ARG:HB2	2.99	0.40
9:S7:165:LYS:H	9:S7:165:LYS:HG2	2.99	0.40
10:S8:191:PHE:O	10:S8:195:ARG:HG2	2.89	0.40
11:S9:60:LEU:HA	11:S9:60:LEU:HD23	3.98	0.40
35:SM:129:ALA:HA	35:SM:132:ALA:HB3	2.56	0.40
34:SR:112:SER:CB	34:SR:153:GLN:HA	2.51	0.40
34:SR:178:VAL:HG23	34:SR:192:PHE:O	4.35	0.40
34:SR:195:HIS:NE2	34:SR:213:SER:O	2.38	0.40
34:SR:243:LEU:O	34:SR:244:ALA:HB3	2.22	0.40
36:1:1184:A:C2	36:1:1323:G:C4	3.09	0.40
36:1:1248:C:OP1	36:1:1249:G:H8	2.05	0.40
36:1:156:G:C5	49:M3:99:HIS:CD2	3.09	0.40
36:1:2182:A:H2'	36:1:2183:A:H8	1.86	0.40
36:1:2202:C:H2'	36:1:2203:U:O4'	2.21	0.40
36:1:2208:A:H4'	36:1:2209:U:OP1	2.22	0.40
36:1:2657:A:OP1	36:1:2657:A:H8	2.05	0.40
36:1:3118:C:H4'	76:Q0:106:ARG:NH1	2.35	0.40
36:1:2320:A:OP2	86:1:4207:OHX:N5	2.54	0.40
36:1:61:A:H2'	36:1:62:A:O4'	2.21	0.40
36:1:902:G:O6	86:1:3931:OHX:N5	2.55	0.40
1:2:1268:G:C2	1:2:1270:G:C8	3.09	0.40
1:2:1770:U:H5	28:D6:28:LYS:NZ	2.19	0.40
1:2:1754:A:O2'	86:2:2057:OHX:N5	2.55	0.40
86:2:2074:OHX:N4	86:2:2162:OHX:N2	2.69	0.40
1:2:222:A:H2'	1:2:223:U:C6	2.56	0.40
1:2:476:U:H5''	1:2:477:A:O4'	2.22	0.40
1:2:982:U:H2'	1:2:983:A:C8	2.57	0.40
37:3:63:A:C2	37:3:65:G:C5	3.10	0.40
86:1:3962:OHX:N1	38:4:31:G:OP2	2.54	0.40
36:5:1459:C:H2'	36:5:1460:A:O4'	2.22	0.40
36:5:1560:G:O2'	36:5:1561:G:OP1	2.33	0.40
36:5:1757:A:H2'	36:5:1758:G:C8	2.57	0.40
36:5:1765:U:H2'	36:5:1766:G:O4'	2.21	0.40
36:5:1830:G:C5	36:5:1831:U:C5	3.10	0.40
36:5:199:A:N3	36:5:201:A:C8	2.90	0.40
36:5:2300:G:C5	36:5:2301:U:C5	3.10	0.40
36:5:2651:G:C2	36:5:2796:G:C4	3.10	0.40
36:5:2859:U:H4'	36:5:2860:U:O5'	2.21	0.40
36:5:3122:A:C2'	36:5:3123:A:H5'	2.50	0.40
36:5:3348:G:N2	36:5:3358:U:C2	2.90	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:5:358:G:N2	36:5:361:A:OP2	2.48	0.40
36:5:568:G:H2'	36:5:569:A:O4'	2.21	0.40
36:5:572:A:H2'	36:5:573:C:H6	1.86	0.40
1:6:1198:G:OP1	1:6:1199:G:H1'	2.22	0.40
1:6:1208:A:H4'	1:6:1270:G:P	2.61	0.40
1:6:1525:A:N1	1:6:1526:A:C2	2.90	0.40
1:6:1698:G:N2	1:6:1699:G:C5	2.90	0.40
1:6:275:C:N4	1:6:281:G:H1	2.20	0.40
1:6:648:G:C4	1:6:687:G:N2	2.90	0.40
38:8:113:U:H3'	38:8:113:U:O2	2.21	0.40
12:C0:48:SER:O	12:C0:51:SER:N	2.52	0.40
13:C1:70:ILE:O	13:C1:71:LEU:HD23	2.22	0.40
13:C1:78:THR:HA	13:C1:84:ILE:HG22	2.03	0.40
14:C2:87:PRO:O	14:C2:88:LEU:HB2	2.21	0.40
15:C3:38:VAL:HA	15:C3:41:ALA:HB3	2.80	0.40
18:C6:115:THR:HG23	18:C6:118:ILE:O	4.89	0.40
18:C6:29:ILE:HG22	18:C6:36:ILE:HB	3.06	0.40
19:C7:19:ARG:HG3	19:C7:20:TYR:CD1	2.57	0.40
19:C7:46:LEU:HD23	19:C7:46:LEU:HA	2.14	0.40
20:C8:123:ARG:CD	20:C8:133:VAL:HG21	2.52	0.40
20:C8:44:ASN:OD1	20:C8:48:LYS:HD2	2.22	0.40
21:C9:33:TYR:O	21:C9:35:ASP:N	3.66	0.40
25:D3:59:ILE:HG12	32:E0:4:VAL:HG13	2.04	0.40
25:D3:96:VAL:HG23	25:D3:97:ASP:H	1.86	0.40
28:D6:36:ILE:HD13	28:D6:36:ILE:O	2.21	0.40
33:E1:108:VAL:HA	33:E1:113:LYS:O	2.21	0.40
40:L3:150:ARG:NH1	40:L3:150:ARG:HG2	2.95	0.40
36:1:3378:C:O2'	40:L3:312:VAL:HA	2.21	0.40
40:L3:328:ILE:HD13	40:L3:328:ILE:HG21	1.84	0.40
40:L3:374:ALA:O	40:L3:378:ALA:N	2.53	0.40
41:L4:156:LEU:C	41:L4:158:SER:H	2.23	0.40
42:L5:105:ILE:HA	42:L5:105:ILE:HD13	1.59	0.40
42:L5:95:TRP:CH2	42:L5:181:PRO:HD3	5.20	0.40
43:L6:67:GLY:HA3	43:L6:68:PRO:C	2.42	0.40
43:L6:39:VAL:O	43:L6:87:THR:HG23	2.72	0.40
45:L8:151:VAL:HG13	45:L8:199:ALA:HB2	2.87	0.40
45:L8:90:THR:OG1	45:L8:91:PHE:N	4.56	0.40
46:L9:90:MET:SD	46:L9:146:LEU:HD11	2.61	0.40
47:M0:191:LYS:NZ	47:M0:212:GLU:OE2	4.07	0.40
48:M1:94:ARG:HB3	48:M1:95:ASN:H	4.38	0.40
49:M3:54:LEU:HD22	49:M3:54:LEU:HA	1.88	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:M3:76:THR:HG21	49:M3:103:ASN:OD1	4.51	0.40
49:M3:25:HIS:CG	51:M5:200:TRP:CZ3	3.10	0.40
52:M6:108:ILE:HG21	52:M6:160:ARG:NH1	4.75	0.40
50:M4:109:ARG:HG3	52:M6:199:TYR:CE2	2.99	0.40
52:M6:78:ARG:NH1	52:M6:78:ARG:HG3	2.37	0.40
53:M7:87:SER:O	53:M7:89:LYS:N	2.54	0.40
54:M8:36:LEU:HA	54:M8:36:LEU:HD23	1.79	0.40
55:M9:107:ALA:O	55:M9:110:ARG:N	2.71	0.40
57:N1:126:VAL:HG23	57:N1:127:GLN:H	1.86	0.40
42:L5:17:GLN:HE22	57:N1:22:HIS:HB2	3.75	0.40
58:N2:99:LYS:HD2	58:N2:102:GLU:CD	2.42	0.40
59:N3:10:LYS:HG2	59:N3:11:PHE:O	2.21	0.40
59:N3:2:SER:N	59:N3:56:ASP:OD1	4.96	0.40
59:N3:97:ASP:OD2	59:N3:99:ALA:HB2	2.21	0.40
36:1:2111:G:N3	60:N4:44:LYS:HE3	2.37	0.40
61:N5:110:VAL:HG22	61:N5:124:VAL:HG13	3.12	0.40
62:N6:12:ARG:O	62:N6:12:ARG:HD2	5.01	0.40
36:1:1430:U:H2'	64:N8:9:ARG:HH22	1.87	0.40
36:1:1433:A:H2	68:O2:25:TYR:CD2	2.40	0.40
68:O2:44:ARG:NH1	68:O2:46:PHE:CE2	4.14	0.40
70:O4:57:LEU:HD13	70:O4:61:GLN:HB3	2.40	0.40
71:O5:38:ARG:HG2	71:O5:39:PRO:HD2	2.02	0.40
74:O8:4:GLU:HG3	36:5:1747:G:H4'	149.70	0.40
78:Q2:14:GLY:C	78:Q2:16:THR:H	2.29	0.40
2:S0:111:ILE:HG21	1:6:1293:U:H1'	420.20	0.40
2:S0:69:ASN:HB3	2:S0:71:GLU:CD	2.76	0.40
3:S1:116:LYS:HB3	3:S1:117:TRP:CD1	5.26	0.40
3:S1:141:ALA:HB1	3:S1:207:LEU:CD2	3.02	0.40
3:S1:207:LEU:HD13	3:S1:210:ILE:HD11	4.97	0.40
3:S1:103:MET:HB3	3:S1:215:VAL:HG12	2.86	0.40
4:S2:115:ILE:HG21	4:S2:208:GLU:OE1	3.95	0.40
5:S3:114:ALA:HB3	5:S3:117:ARG:HB2	2.04	0.40
5:S3:44:THR:O	5:S3:45:LYS:HB2	4.23	0.40
1:2:1514:U:C4	5:S3:4:LEU:HD12	2.57	0.40
7:S5:25:LEU:N	7:S5:25:LEU:HD22	2.40	0.40
8:S6:138:ALA:O	8:S6:142:ARG:HG3	3.34	0.40
8:S6:69:LEU:HD13	8:S6:69:LEU:HA	1.87	0.40
9:S7:99:LEU:HA	9:S7:99:LEU:HD23	1.92	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)
1:2:1353:U:O2'	36:5:3165:A:OP1[2_546]	2.15	0.05

5.3 Torsion angles [i](#)

5.3.1 Protein backbone [i](#)

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	S0	204/251 (81%)	142 (70%)	37 (18%)	25 (12%)	0	5
2	s0	204/251 (81%)	150 (74%)	36 (18%)	18 (9%)	1	10
3	S1	212/254 (84%)	142 (67%)	42 (20%)	28 (13%)	0	4
3	s1	214/254 (84%)	165 (77%)	37 (17%)	12 (6%)	2	20
4	S2	215/253 (85%)	175 (81%)	23 (11%)	17 (8%)	1	12
4	s2	215/253 (85%)	169 (79%)	28 (13%)	18 (8%)	1	11
5	S3	221/239 (92%)	170 (77%)	37 (17%)	14 (6%)	1	17
5	s3	221/239 (92%)	174 (79%)	25 (11%)	22 (10%)	1	8
6	S4	258/260 (99%)	201 (78%)	43 (17%)	14 (5%)	2	21
6	s4	258/260 (99%)	206 (80%)	35 (14%)	17 (7%)	1	16
7	S5	204/224 (91%)	150 (74%)	33 (16%)	21 (10%)	0	8
7	s5	204/224 (91%)	140 (69%)	41 (20%)	23 (11%)	0	6
8	S6	224/236 (95%)	181 (81%)	33 (15%)	10 (4%)	3	26
8	s6	216/236 (92%)	172 (80%)	34 (16%)	10 (5%)	3	25
9	S7	182/189 (96%)	137 (75%)	27 (15%)	18 (10%)	1	8
9	s7	184/189 (97%)	133 (72%)	34 (18%)	17 (9%)	1	9
10	S8	184/200 (92%)	145 (79%)	29 (16%)	10 (5%)	2	21
10	s8	184/200 (92%)	149 (81%)	27 (15%)	8 (4%)	3	27
11	S9	183/196 (93%)	144 (79%)	27 (15%)	12 (7%)	1	16
11	s9	183/196 (93%)	133 (73%)	40 (22%)	10 (6%)	2	20
12	C0	94/105 (90%)	68 (72%)	19 (20%)	7 (7%)	1	13

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
12	c0	92/105 (88%)	68 (74%)	11 (12%)	13 (14%)	0	3
13	C1	153/155 (99%)	107 (70%)	27 (18%)	19 (12%)	0	5
13	c1	144/155 (93%)	114 (79%)	23 (16%)	7 (5%)	2	23
14	C2	122/142 (86%)	66 (54%)	39 (32%)	17 (14%)	0	4
14	c2	122/142 (86%)	72 (59%)	30 (25%)	20 (16%)	0	2
15	C3	148/150 (99%)	117 (79%)	24 (16%)	7 (5%)	3	25
15	c3	148/150 (99%)	107 (72%)	28 (19%)	13 (9%)	1	10
16	C4	125/136 (92%)	93 (74%)	23 (18%)	9 (7%)	1	14
16	c4	126/136 (93%)	99 (79%)	18 (14%)	9 (7%)	1	14
17	C5	122/141 (86%)	84 (69%)	24 (20%)	14 (12%)	0	6
17	c5	133/141 (94%)	92 (69%)	24 (18%)	17 (13%)	0	5
18	C6	139/142 (98%)	117 (84%)	16 (12%)	6 (4%)	3	27
18	c6	140/142 (99%)	106 (76%)	23 (16%)	11 (8%)	1	12
19	C7	116/136 (85%)	85 (73%)	18 (16%)	13 (11%)	0	6
19	c7	113/136 (83%)	84 (74%)	22 (20%)	7 (6%)	2	18
20	C8	143/145 (99%)	107 (75%)	23 (16%)	13 (9%)	1	9
20	c8	143/145 (99%)	105 (73%)	25 (18%)	13 (9%)	1	9
21	C9	141/143 (99%)	108 (77%)	27 (19%)	6 (4%)	3	27
21	c9	141/143 (99%)	114 (81%)	21 (15%)	6 (4%)	3	27
22	D0	105/120 (88%)	79 (75%)	20 (19%)	6 (6%)	2	20
22	d0	108/120 (90%)	83 (77%)	14 (13%)	11 (10%)	1	8
23	D1	85/87 (98%)	58 (68%)	19 (22%)	8 (9%)	1	9
23	d1	85/87 (98%)	68 (80%)	12 (14%)	5 (6%)	2	19
24	D2	127/129 (98%)	103 (81%)	17 (13%)	7 (6%)	2	20
24	d2	127/129 (98%)	101 (80%)	23 (18%)	3 (2%)	7	42
25	D3	142/144 (99%)	107 (75%)	22 (16%)	13 (9%)	1	9
25	d3	142/144 (99%)	115 (81%)	19 (13%)	8 (6%)	2	20
26	D4	132/134 (98%)	105 (80%)	17 (13%)	10 (8%)	1	12
26	d4	132/134 (98%)	100 (76%)	15 (11%)	17 (13%)	0	5
27	D5	68/107 (64%)	44 (65%)	14 (21%)	10 (15%)	0	3
27	d5	67/107 (63%)	47 (70%)	15 (22%)	5 (8%)	1	13

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
28	D6	95/97 (98%)	59 (62%)	23 (24%)	13 (14%)	0	4
28	d6	95/97 (98%)	70 (74%)	18 (19%)	7 (7%)	1	13
29	D7	79/81 (98%)	55 (70%)	18 (23%)	6 (8%)	1	12
29	d7	79/81 (98%)	53 (67%)	21 (27%)	5 (6%)	1	17
30	D8	61/66 (92%)	42 (69%)	14 (23%)	5 (8%)	1	11
30	d8	61/66 (92%)	43 (70%)	14 (23%)	4 (7%)	1	16
31	D9	51/55 (93%)	37 (72%)	10 (20%)	4 (8%)	1	12
31	d9	51/55 (93%)	39 (76%)	8 (16%)	4 (8%)	1	12
32	E0	58/60 (97%)	43 (74%)	10 (17%)	5 (9%)	1	11
33	E1	69/76 (91%)	32 (46%)	20 (29%)	17 (25%)	0	1
33	e1	74/76 (97%)	33 (45%)	21 (28%)	20 (27%)	0	0
34	SR	316/318 (99%)	231 (73%)	58 (18%)	27 (8%)	1	11
34	sR	316/318 (99%)	255 (81%)	49 (16%)	12 (4%)	4	30
35	SM	155/273 (57%)	99 (64%)	34 (22%)	22 (14%)	0	3
35	sM	98/273 (36%)	63 (64%)	23 (24%)	12 (12%)	0	5
39	L2	250/253 (99%)	214 (86%)	28 (11%)	8 (3%)	5	35
39	l2	250/253 (99%)	200 (80%)	32 (13%)	18 (7%)	1	14
40	L3	384/386 (100%)	324 (84%)	45 (12%)	15 (4%)	3	30
40	l3	384/386 (100%)	323 (84%)	42 (11%)	19 (5%)	2	23
41	L4	359/361 (99%)	279 (78%)	54 (15%)	26 (7%)	1	14
41	l4	359/361 (99%)	277 (77%)	56 (16%)	26 (7%)	1	14
42	L5	294/296 (99%)	237 (81%)	34 (12%)	23 (8%)	1	12
42	l5	292/296 (99%)	240 (82%)	39 (13%)	13 (4%)	3	26
43	L6	152/175 (87%)	121 (80%)	26 (17%)	5 (3%)	4	34
43	l6	153/175 (87%)	121 (79%)	25 (16%)	7 (5%)	3	25
44	L7	220/243 (90%)	183 (83%)	24 (11%)	13 (6%)	2	19
44	l7	221/243 (91%)	177 (80%)	30 (14%)	14 (6%)	1	17
45	L8	231/255 (91%)	179 (78%)	35 (15%)	17 (7%)	1	13
45	l8	229/255 (90%)	176 (77%)	36 (16%)	17 (7%)	1	13
46	L9	189/191 (99%)	150 (79%)	28 (15%)	11 (6%)	2	19
46	l9	189/191 (99%)	153 (81%)	27 (14%)	9 (5%)	2	24

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
47	M0	207/220 (94%)	167 (81%)	30 (14%)	10 (5%)	2	24
47	m0	209/220 (95%)	163 (78%)	27 (13%)	19 (9%)	1	9
48	M1	167/173 (96%)	126 (75%)	31 (19%)	10 (6%)	2	18
48	m1	167/173 (96%)	127 (76%)	25 (15%)	15 (9%)	1	9
49	M3	191/198 (96%)	140 (73%)	37 (19%)	14 (7%)	1	13
49	m3	192/198 (97%)	150 (78%)	24 (12%)	18 (9%)	1	9
50	M4	134/137 (98%)	106 (79%)	19 (14%)	9 (7%)	1	16
50	m4	135/137 (98%)	116 (86%)	16 (12%)	3 (2%)	8	43
51	M5	201/203 (99%)	170 (85%)	24 (12%)	7 (4%)	4	33
51	m5	201/203 (99%)	171 (85%)	25 (12%)	5 (2%)	6	40
52	M6	195/198 (98%)	167 (86%)	22 (11%)	6 (3%)	5	36
52	m6	195/198 (98%)	164 (84%)	23 (12%)	8 (4%)	3	29
53	M7	181/183 (99%)	142 (78%)	27 (15%)	12 (7%)	1	16
53	m7	153/183 (84%)	130 (85%)	20 (13%)	3 (2%)	9	45
54	M8	183/185 (99%)	148 (81%)	28 (15%)	7 (4%)	4	30
54	m8	183/185 (99%)	140 (76%)	30 (16%)	13 (7%)	1	14
55	M9	186/188 (99%)	160 (86%)	17 (9%)	9 (5%)	2	24
55	m9	186/188 (99%)	156 (84%)	26 (14%)	4 (2%)	8	43
56	N0	170/172 (99%)	149 (88%)	14 (8%)	7 (4%)	3	29
56	n0	170/172 (99%)	146 (86%)	19 (11%)	5 (3%)	5	37
57	N1	157/159 (99%)	136 (87%)	16 (10%)	5 (3%)	5	35
57	n1	157/159 (99%)	125 (80%)	26 (17%)	6 (4%)	4	30
58	N2	98/120 (82%)	74 (76%)	16 (16%)	8 (8%)	1	11
58	n2	96/120 (80%)	76 (79%)	17 (18%)	3 (3%)	5	36
59	N3	134/136 (98%)	114 (85%)	14 (10%)	6 (4%)	3	26
59	n3	134/136 (98%)	116 (87%)	9 (7%)	9 (7%)	1	16
60	N4	96/155 (62%)	68 (71%)	20 (21%)	8 (8%)	1	11
60	n4	133/155 (86%)	99 (74%)	21 (16%)	13 (10%)	1	8
61	N5	119/141 (84%)	98 (82%)	17 (14%)	4 (3%)	4	34
61	n5	118/141 (84%)	97 (82%)	16 (14%)	5 (4%)	3	28
62	N6	124/126 (98%)	99 (80%)	16 (13%)	9 (7%)	1	13

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
62	n6	124/126 (98%)	100 (81%)	16 (13%)	8 (6%)	1	16
63	N7	133/135 (98%)	104 (78%)	20 (15%)	9 (7%)	1	16
63	n7	133/135 (98%)	91 (68%)	30 (23%)	12 (9%)	1	9
64	N8	146/148 (99%)	114 (78%)	23 (16%)	9 (6%)	2	18
64	n8	146/148 (99%)	119 (82%)	19 (13%)	8 (6%)	2	20
65	N9	56/58 (97%)	39 (70%)	11 (20%)	6 (11%)	0	7
65	n9	56/58 (97%)	36 (64%)	11 (20%)	9 (16%)	0	2
66	O0	95/104 (91%)	76 (80%)	17 (18%)	2 (2%)	8	44
66	o0	98/104 (94%)	83 (85%)	11 (11%)	4 (4%)	3	29
67	O1	107/112 (96%)	87 (81%)	13 (12%)	7 (6%)	1	16
67	o1	107/112 (96%)	87 (81%)	12 (11%)	8 (8%)	1	13
68	O2	125/129 (97%)	99 (79%)	18 (14%)	8 (6%)	1	17
68	o2	125/129 (97%)	102 (82%)	14 (11%)	9 (7%)	1	14
69	O3	104/106 (98%)	86 (83%)	13 (12%)	5 (5%)	2	24
69	o3	104/106 (98%)	95 (91%)	7 (7%)	2 (2%)	9	46
70	O4	110/120 (92%)	92 (84%)	16 (14%)	2 (2%)	10	47
70	o4	110/120 (92%)	90 (82%)	15 (14%)	5 (4%)	3	26
71	O5	117/119 (98%)	99 (85%)	13 (11%)	5 (4%)	3	27
71	o5	117/119 (98%)	89 (76%)	22 (19%)	6 (5%)	2	22
72	O6	97/99 (98%)	70 (72%)	15 (16%)	12 (12%)	0	5
72	o6	97/99 (98%)	83 (86%)	10 (10%)	4 (4%)	3	29
73	O7	85/87 (98%)	66 (78%)	18 (21%)	1 (1%)	15	56
73	o7	85/87 (98%)	69 (81%)	12 (14%)	4 (5%)	3	25
74	O8	75/77 (97%)	61 (81%)	11 (15%)	3 (4%)	3	29
74	o8	75/77 (97%)	63 (84%)	10 (13%)	2 (3%)	6	39
75	O9	48/50 (96%)	41 (85%)	6 (12%)	1 (2%)	8	44
75	o9	48/50 (96%)	38 (79%)	6 (12%)	4 (8%)	1	11
76	Q0	50/52 (96%)	42 (84%)	3 (6%)	5 (10%)	1	8
76	q0	50/52 (96%)	40 (80%)	8 (16%)	2 (4%)	3	29
77	Q1	23/25 (92%)	16 (70%)	5 (22%)	2 (9%)	1	10
77	q1	23/25 (92%)	19 (83%)	2 (9%)	2 (9%)	1	10

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
78	Q2	103/105 (98%)	77 (75%)	18 (18%)	8 (8%)	1	12
78	q2	103/105 (98%)	87 (84%)	11 (11%)	5 (5%)	2	23
79	Q3	89/91 (98%)	74 (83%)	8 (9%)	7 (8%)	1	12
79	q3	89/91 (98%)	70 (79%)	13 (15%)	6 (7%)	1	16
80	e0	60/62 (97%)	39 (65%)	15 (25%)	6 (10%)	1	8
81	p0	139/311 (45%)	108 (78%)	24 (17%)	7 (5%)	2	23
All	All	22333/24143 (92%)	17400 (78%)	3410 (15%)	1523 (7%)	1	16

All (1523) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
2	S0	4	PRO
2	S0	39	ASN
2	S0	68	PRO
2	S0	139	VAL
2	S0	158	VAL
3	S1	49	ASN
3	S1	58	SER
3	S1	63	GLY
3	S1	79	HIS
3	S1	132	ASP
3	S1	148	ASN
3	S1	158	SER
3	S1	179	SER
3	S1	182	ALA
3	S1	206	PRO
3	S1	223	PHE
4	S2	80	VAL
4	S2	81	MET
4	S2	163	GLY
5	S3	65	ARG
5	S3	93	ASP
5	S3	220	PRO
6	S4	12	LEU
6	S4	69	HIS
6	S4	96	ASN
7	S5	35	GLN
7	S5	39	GLU
7	S5	43	PHE
7	S5	51	VAL

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Mol	Chain	Res	Type
7	S5	58	LEU
8	S6	122	GLU
8	S6	173	PRO
8	S6	174	LYS
9	S7	31	SER
9	S7	32	PRO
9	S7	63	PRO
9	S7	112	ARG
9	S7	116	ARG
9	S7	131	PHE
9	S7	186	PRO
10	S8	81	VAL
10	S8	152	ILE
11	S9	134	ILE
12	C0	87	VAL
12	C0	88	PRO
13	C1	4	GLU
13	C1	7	VAL
13	C1	55	ASP
13	C1	95	PRO
13	C1	96	LYS
13	C1	140	VAL
13	C1	146	ALA
13	C1	154	ALA
14	C2	91	VAL
15	C3	138	ASN
16	C4	42	VAL
16	C4	124	ASP
16	C4	126	THR
17	C5	125	PRO
17	C5	126	VAL
18	C6	41	PRO
18	C6	114	ARG
19	C7	83	GLN
19	C7	84	TYR
19	C7	85	VAL
19	C7	86	PRO
19	C7	88	VAL
19	C7	104	ASN
19	C7	124	VAL
20	C8	91	ASP
20	C8	92	ILE

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Mol	Chain	Res	Type
21	C9	53	TRP
21	C9	69	LYS
23	D1	6	GLY
24	D2	57	ARG
24	D2	78	ARG
24	D2	83	ILE
25	D3	70	LYS
25	D3	112	LYS
25	D3	114	LYS
25	D3	128	SER
25	D3	137	LYS
25	D3	144	ARG
26	D4	5	VAL
26	D4	53	ASP
26	D4	58	PHE
27	D5	37	GLN
27	D5	43	ASP
27	D5	56	THR
27	D5	71	ILE
28	D6	36	ILE
28	D6	45	VAL
28	D6	84	VAL
28	D6	86	VAL
29	D7	38	PRO
29	D7	62	ILE
29	D7	75	GLU
31	D9	8	PHE
31	D9	47	ALA
32	E0	13	LYS
32	E0	47	VAL
33	E1	98	VAL
33	E1	102	VAL
33	E1	103	LEU
33	E1	105	TYR
33	E1	106	TYR
33	E1	110	ALA
33	E1	111	GLU
33	E1	138	ARG
34	SR	155	ARG
34	SR	161	LYS
34	SR	188	ILE
34	SR	231	MET

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Mol	Chain	Res	Type
34	SR	318	ALA
35	SM	52	PRO
35	SM	86	ASN
35	SM	87	THR
35	SM	140	ASP
35	SM	166	VAL
35	SM	167	PRO
40	L3	140	ASP
40	L3	348	ARG
40	L3	385	LYS
41	L4	24	ALA
41	L4	83	GLY
41	L4	132	ALA
41	L4	146	PRO
41	L4	183	LYS
41	L4	265	GLU
41	L4	270	SER
41	L4	292	SER
41	L4	311	HIS
42	L5	6	ASP
42	L5	59	ASP
42	L5	178	ASN
42	L5	215	ASP
42	L5	234	ASP
42	L5	253	PHE
42	L5	258	LYS
42	L5	292	ALA
42	L5	293	LEU
43	L6	94	GLU
43	L6	98	VAL
44	L7	24	GLU
44	L7	191	VAL
45	L8	25	PRO
45	L8	31	PRO
45	L8	36	ILE
45	L8	37	GLY
45	L8	40	VAL
45	L8	156	ASP
45	L8	255	SER
46	L9	50	ASN
47	M0	145	LYS
47	M0	207	GLU

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Mol	Chain	Res	Type
48	M1	8	PRO
48	M1	145	LYS
48	M1	165	GLN
49	M3	129	ASN
50	M4	6	ILE
50	M4	8	LYS
50	M4	9	ALA
50	M4	135	LEU
50	M4	136	ALA
51	M5	40	ALA
52	M6	111	PRO
53	M7	88	VAL
53	M7	109	ALA
53	M7	157	VAL
55	M9	61	SER
55	M9	85	ARG
56	N0	2	ALA
57	N1	159	PHE
58	N2	51	GLY
59	N3	3	GLY
60	N4	26	SER
60	N4	81	PRO
61	N5	26	VAL
61	N5	50	ALA
62	N6	53	ASP
63	N7	30	ASP
63	N7	125	GLY
63	N7	129	TRP
64	N8	24	LYS
67	O1	5	LYS
67	O1	6	ASP
67	O1	83	GLU
68	O2	30	GLU
69	O3	94	PHE
71	O5	96	GLU
72	O6	27	SER
73	O7	12	HIS
75	O9	4	GLN
76	Q0	78	ILE
76	Q0	120	GLN
77	Q1	4	LYS
78	Q2	30	ALA

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Mol	Chain	Res	Type
78	Q2	100	LYS
79	Q3	58	SER
2	s0	4	PRO
2	s0	68	PRO
2	s0	158	VAL
2	s0	191	ARG
2	s0	203	PHE
2	s0	206	ASP
3	s1	132	ASP
3	s1	147	ALA
3	s1	206	PRO
3	s1	232	HIS
4	s2	76	LEU
4	s2	92	ALA
4	s2	134	LEU
4	s2	148	LEU
4	s2	150	GLN
4	s2	207	LEU
5	s3	92	GLN
5	s3	144	ALA
5	s3	161	GLY
5	s3	211	PRO
5	s3	216	PRO
5	s3	217	ILE
5	s3	220	PRO
5	s3	221	SER
6	s4	95	THR
6	s4	104	ASP
6	s4	118	GLU
6	s4	163	ASP
6	s4	195	ILE
6	s4	196	VAL
6	s4	240	LYS
7	s5	28	PRO
7	s5	35	GLN
7	s5	36	ALA
7	s5	39	GLU
7	s5	81	ARG
7	s5	154	ALA
7	s5	184	PHE
7	s5	204	GLY
7	s5	209	TYR

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Mol	Chain	Res	Type
8	s6	70	PRO
8	s6	153	VAL
8	s6	173	PRO
9	s7	64	VAL
9	s7	66	SER
9	s7	67	LEU
9	s7	106	SER
9	s7	112	ARG
9	s7	131	PHE
9	s7	163	ASP
10	s8	13	ALA
10	s8	36	THR
11	s9	5	PRO
12	c0	2	LEU
12	c0	28	ASN
12	c0	32	HIS
12	c0	83	PRO
12	c0	88	PRO
12	c0	92	ILE
12	c0	97	PRO
13	c1	114	ALA
13	c1	129	ARG
14	c2	89	ILE
14	c2	93	ASP
15	c3	60	VAL
15	c3	66	ILE
15	c3	139	TRP
16	c4	51	ASP
16	c4	132	ARG
17	c5	11	VAL
17	c5	49	MET
17	c5	51	SER
17	c5	52	LYS
17	c5	125	PRO
17	c5	128	HIS
18	c6	39	VAL
18	c6	116	LEU
19	c7	67	ARG
19	c7	88	VAL
19	c7	99	VAL
20	c8	18	LEU
20	c8	61	LEU

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Mol	Chain	Res	Type
20	c8	91	ASP
20	c8	92	ILE
22	d0	17	GLN
22	d0	51	VAL
22	d0	52	LYS
22	d0	118	VAL
23	d1	4	ASP
25	d3	70	LYS
26	d4	30	PRO
26	d4	35	VAL
26	d4	52	LYS
26	d4	63	GLN
27	d5	85	LYS
27	d5	87	GLY
27	d5	104	ALA
28	d6	63	ALA
29	d7	18	LYS
29	d7	38	PRO
29	d7	60	SER
31	d9	6	VAL
31	d9	7	TRP
31	d9	19	ARG
33	e1	83	LYS
33	e1	84	VAL
33	e1	87	THR
33	e1	92	LYS
33	e1	98	VAL
33	e1	100	LEU
33	e1	102	VAL
33	e1	103	LEU
33	e1	106	TYR
34	sR	4	ASN
34	sR	165	ASP
34	sR	282	SER
35	sM	47	ALA
35	sM	50	ASN
35	sM	163	PHE
39	l2	24	GLN
39	l2	96	LEU
39	l2	144	ASN
39	l2	160	SER
39	l2	215	ASN

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Mol	Chain	Res	Type
39	l2	238	ILE
39	l2	249	SER
40	l3	129	ALA
40	l3	140	ASP
40	l3	142	ALA
40	l3	188	ILE
40	l3	200	GLU
40	l3	235	THR
40	l3	347	SER
41	l4	14	GLU
41	l4	24	ALA
41	l4	90	PHE
41	l4	193	LYS
41	l4	268	ALA
41	l4	277	PRO
41	l4	311	HIS
41	l4	329	PRO
41	l4	338	LYS
41	l4	339	LEU
41	l4	345	GLU
42	l5	178	ASN
42	l5	279	LYS
43	l6	3	ALA
43	l6	98	VAL
44	l7	178	ILE
44	l7	193	PRO
45	l8	25	PRO
45	l8	122	LYS
45	l8	208	GLU
45	l8	240	ASN
46	l9	141	LYS
46	l9	177	ASP
47	m0	82	ARG
47	m0	156	ARG
47	m0	157	TYR
48	m1	8	PRO
48	m1	9	MET
48	m1	10	ARG
48	m1	94	ARG
48	m1	173	ASP
49	m3	25	HIS
49	m3	47	ALA

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Mol	Chain	Res	Type
49	m3	50	PRO
49	m3	133	PRO
49	m3	134	GLU
49	m3	150	PRO
49	m3	152	THR
49	m3	193	ALA
50	m4	78	THR
51	m5	76	PRO
51	m5	81	TYR
51	m5	187	ARG
52	m6	16	VAL
52	m6	90	HIS
52	m6	186	ALA
54	m8	84	VAL
54	m8	95	GLU
54	m8	112	ALA
55	m9	182	ASP
56	n0	87	THR
56	n0	139	TYR
57	n1	122	GLN
57	n1	144	GLU
57	n1	146	ASN
58	n2	50	LEU
60	n4	25	ASP
60	n4	63	ILE
60	n4	64	THR
60	n4	76	VAL
61	n5	40	LEU
61	n5	55	ASN
62	n6	6	LEU
62	n6	62	SER
62	n6	83	ASP
62	n6	84	LYS
62	n6	90	VAL
62	n6	97	ILE
63	n7	5	LEU
63	n7	7	ALA
63	n7	90	GLU
63	n7	125	GLY
63	n7	129	TRP
64	n8	76	ASP
65	n9	21	ILE

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Mol	Chain	Res	Type
65	n9	23	LYS
65	n9	24	PRO
65	n9	39	PHE
68	o2	4	LEU
68	o2	5	PRO
70	o4	33	GLN
71	o5	43	LYS
71	o5	119	LYS
72	o6	12	ASN
72	o6	98	ARG
73	o7	68	LYS
73	o7	84	SER
73	o7	85	LYS
74	o8	16	ARG
75	o9	35	ILE
79	q3	52	ALA
81	p0	93	LEU
2	S0	5	ALA
2	S0	33	GLN
2	S0	43	ASP
2	S0	64	ILE
2	S0	111	ILE
2	S0	185	ARG
2	S0	190	ASP
2	S0	202	TYR
3	S1	54	LEU
3	S1	181	LEU
3	S1	218	LEU
3	S1	221	PRO
4	S2	78	ASP
4	S2	106	ASP
4	S2	107	SER
4	S2	148	LEU
5	S3	62	ASN
5	S3	137	VAL
5	S3	218	LEU
6	S4	17	HIS
6	S4	77	ARG
6	S4	153	ASN
6	S4	205	PHE
6	S4	228	ILE
7	S5	33	VAL

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Mol	Chain	Res	Type
7	S5	50	GLU
7	S5	54	LYS
7	S5	63	GLN
7	S5	127	GLN
7	S5	153	GLY
7	S5	204	GLY
8	S6	70	PRO
8	S6	138	ALA
8	S6	152	ASP
8	S6	165	GLY
9	S7	8	ILE
9	S7	30	SER
9	S7	35	LYS
9	S7	52	ALA
9	S7	64	VAL
9	S7	73	VAL
9	S7	110	GLN
9	S7	132	PRO
10	S8	22	ARG
10	S8	40	ALA
10	S8	149	SER
11	S9	98	ALA
11	S9	164	PHE
12	C0	60	SER
13	C1	51	GLY
13	C1	72	THR
13	C1	89	ALA
13	C1	139	VAL
13	C1	145	ALA
14	C2	87	PRO
14	C2	89	ILE
14	C2	101	ALA
14	C2	106	ILE
14	C2	119	SER
14	C2	126	TRP
14	C2	131	ASP
15	C3	22	ALA
15	C3	27	LYS
16	C4	50	ALA
16	C4	92	LYS
17	C5	48	GLY
17	C5	53	PRO

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Mol	Chain	Res	Type
17	C5	80	MET
18	C6	39	VAL
19	C7	23	LYS
20	C8	7	GLU
20	C8	14	ILE
20	C8	60	GLU
20	C8	82	PRO
20	C8	142	GLY
20	C8	144	ARG
22	D0	21	LYS
22	D0	72	ASN
23	D1	12	TYR
26	D4	36	SER
26	D4	59	GLY
27	D5	44	GLN
27	D5	55	PRO
27	D5	88	ILE
28	D6	39	MET
28	D6	82	ARG
28	D6	85	ARG
29	D7	4	VAL
29	D7	63	LEU
30	D8	22	ARG
30	D8	36	THR
32	E0	3	LYS
32	E0	9	ALA
33	E1	84	VAL
33	E1	85	TYR
33	E1	87	THR
33	E1	127	GLY
33	E1	128	ALA
34	SR	24	ALA
34	SR	141	LEU
34	SR	166	SER
35	SM	101	ASP
35	SM	139	GLU
35	SM	154	TYR
39	L2	13	GLY
39	L2	234	LYS
40	L3	4	ARG
40	L3	155	ALA
40	L3	186	GLY

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Mol	Chain	Res	Type
40	L3	333	LYS
41	L4	82	THR
41	L4	90	PHE
41	L4	130	ALA
41	L4	143	GLU
41	L4	175	HIS
41	L4	293	SER
42	L5	7	ALA
42	L5	9	SER
42	L5	57	ASN
42	L5	91	GLY
42	L5	93	THR
43	L6	130	ILE
44	L7	26	VAL
44	L7	159	GLN
44	L7	241	LYS
45	L8	39	ALA
45	L8	157	VAL
46	L9	49	ASN
46	L9	151	VAL
46	L9	189	GLU
47	M0	146	ASP
47	M0	194	GLY
48	M1	39	GLN
48	M1	115	LYS
49	M3	19	GLN
49	M3	47	ALA
50	M4	109	ARG
51	M5	141	ALA
51	M5	184	LYS
52	M6	178	VAL
53	M7	160	ALA
53	M7	164	LYS
54	M8	98	LYS
55	M9	128	LYS
56	N0	118	PHE
56	N0	130	GLU
56	N0	166	LYS
57	N1	124	VAL
58	N2	11	ILE
58	N2	50	LEU
58	N2	60	GLY

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Mol	Chain	Res	Type
58	N2	70	LYS
58	N2	107	PHE
59	N3	131	SER
60	N4	72	SER
61	N5	117	ASN
62	N6	84	LYS
63	N7	102	GLU
64	N8	47	LYS
64	N8	66	ALA
65	N9	5	LYS
65	N9	24	PRO
65	N9	25	LYS
66	O0	20	SER
67	O1	84	ASP
68	O2	13	HIS
68	O2	27	ARG
69	O3	40	ASP
70	O4	77	GLY
71	O5	94	LYS
71	O5	97	ALA
71	O5	119	LYS
72	O6	3	VAL
72	O6	21	THR
72	O6	33	ALA
72	O6	64	SER
74	O8	75	VAL
78	Q2	15	LYS
78	Q2	17	CYS
78	Q2	94	GLY
79	Q3	60	CYS
79	Q3	71	VAL
79	Q3	90	VAL
2	s0	29	VAL
2	s0	189	VAL
3	s1	63	GLY
3	s1	81	PHE
3	s1	93	GLY
3	s1	223	PHE
4	s2	163	GLY
5	s3	44	THR
5	s3	142	LEU
5	s3	195	SER

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Mol	Chain	Res	Type
5	s3	203	PRO
6	s4	164	LEU
6	s4	245	LYS
7	s5	43	PHE
7	s5	75	GLY
7	s5	101	GLY
7	s5	196	GLU
7	s5	217	LEU
8	s6	9	VAL
8	s6	123	GLY
8	s6	154	ARG
8	s6	174	LYS
9	s7	15	GLU
9	s7	74	GLN
9	s7	155	ASP
9	s7	156	SER
10	s8	94	ASN
10	s8	162	ALA
10	s8	199	LYS
11	s9	99	LEU
11	s9	122	VAL
11	s9	167	ALA
13	c1	7	VAL
13	c1	24	LYS
13	c1	55	ASP
13	c1	144	ALA
14	c2	101	ALA
15	c3	21	ASN
15	c3	35	GLU
15	c3	87	ASP
15	c3	140	LYS
16	c4	32	ASP
17	c5	17	TYR
17	c5	50	THR
17	c5	69	GLU
17	c5	126	VAL
18	c6	42	GLU
18	c6	113	ASP
18	c6	115	THR
18	c6	120	ASP
19	c7	63	LYS
20	c8	4	VAL

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Mol	Chain	Res	Type
20	c8	29	VAL
20	c8	55	HIS
20	c8	90	ASN
20	c8	139	LYS
21	c9	29	GLU
21	c9	142	GLU
22	d0	15	GLN
22	d0	16	GLN
23	d1	42	GLU
25	d3	3	LYS
25	d3	47	SER
25	d3	65	ASN
25	d3	138	GLU
26	d4	32	ARG
26	d4	33	ALA
26	d4	36	SER
26	d4	51	GLU
26	d4	58	PHE
26	d4	64	PHE
26	d4	65	GLY
26	d4	84	LYS
26	d4	121	THR
26	d4	123	LYS
27	d5	38	HIS
80	e0	44	PHE
80	e0	51	ASN
33	e1	128	ALA
34	sR	149	ASP
34	sR	194	GLY
35	sM	33	LYS
35	sM	65	THR
35	sM	168	GLU
39	l2	54	ARG
39	l2	172	GLY
39	l2	194	ASN
40	l3	23	ALA
40	l3	66	LYS
40	l3	230	THR
40	l3	385	LYS
40	l3	386	ASP
41	l4	15	ALA
41	l4	145	ILE

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Mol	Chain	Res	Type
41	l4	341	SER
41	l4	342	LYS
42	l5	9	SER
42	l5	115	LEU
42	l5	256	THR
42	l5	260	PHE
43	l6	24	ALA
43	l6	32	ALA
44	l7	91	GLY
44	l7	130	ILE
45	l8	26	LEU
45	l8	112	GLU
45	l8	120	LYS
45	l8	133	LYS
45	l8	182	GLY
45	l8	203	VAL
46	l9	144	ILE
47	m0	3	ARG
47	m0	25	ALA
47	m0	83	ASP
47	m0	101	LYS
47	m0	174	THR
47	m0	187	ALA
47	m0	196	PHE
48	m1	95	ASN
48	m1	108	GLU
48	m1	114	ILE
49	m3	76	THR
49	m3	101	ARG
49	m3	122	LYS
51	m5	183	THR
52	m6	139	GLY
53	m7	34	GLN
54	m8	70	ALA
54	m8	99	THR
54	m8	181	SER
56	n0	129	ILE
58	n2	91	ASP
59	n3	54	LEU
59	n3	134	GLY
60	n4	34	SER
60	n4	71	ARG

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Mol	Chain	Res	Type
60	n4	77	LYS
60	n4	98	PRO
60	n4	126	GLU
61	n5	41	ALA
62	n6	61	GLY
63	n7	6	LYS
63	n7	102	GLU
63	n7	128	GLN
64	n8	15	VAL
64	n8	24	LYS
65	n9	6	ASN
65	n9	52	LYS
66	o0	49	PRO
67	o1	26	LYS
67	o1	45	GLY
67	o1	84	ASP
68	o2	124	GLY
70	o4	79	SER
74	o8	18	ALA
75	o9	4	GLN
78	q2	15	LYS
78	q2	17	CYS
79	q3	9	GLY
81	p0	102	SER
2	S0	30	GLN
2	S0	49	ASN
2	S0	66	ALA
2	S0	103	THR
2	S0	163	ASN
2	S0	192	THR
2	S0	195	TRP
3	S1	35	PRO
3	S1	81	PHE
3	S1	130	SER
3	S1	154	SER
3	S1	213	ARG
4	S2	91	ARG
4	S2	146	THR
4	S2	150	GLN
5	S3	31	GLU
5	S3	44	THR
5	S3	195	SER

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Mol	Chain	Res	Type
5	S3	217	ILE
6	S4	104	ASP
6	S4	195	ILE
7	S5	49	GLU
7	S5	64	VAL
7	S5	81	ARG
8	S6	151	ASP
9	S7	36	ALA
9	S7	118	LEU
10	S8	52	ASN
10	S8	68	ALA
10	S8	120	THR
11	S9	9	SER
11	S9	163	PRO
12	C0	54	TYR
12	C0	81	ASN
12	C0	90	THR
13	C1	29	LYS
13	C1	30	ARG
14	C2	22	VAL
14	C2	105	LYS
14	C2	107	ASP
15	C3	68	GLY
15	C3	149	LEU
16	C4	18	ARG
16	C4	40	ALA
17	C5	25	LEU
17	C5	29	SER
17	C5	39	ALA
17	C5	51	SER
17	C5	54	ALA
17	C5	69	GLU
17	C5	101	ALA
18	C6	113	ASP
18	C6	142	TYR
19	C7	87	GLU
19	C7	113	LEU
19	C7	115	LEU
20	C8	8	GLN
20	C8	61	LEU
20	C8	110	ARG
21	C9	31	PRO

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Mol	Chain	Res	Type
21	C9	39	THR
23	D1	2	GLU
24	D2	77	PRO
24	D2	91	ALA
24	D2	92	ASN
26	D4	6	THR
26	D4	68	LYS
27	D5	39	ALA
27	D5	94	LYS
28	D6	63	ALA
29	D7	48	SER
31	D9	20	GLN
33	E1	86	THR
33	E1	90	LYS
34	SR	22	SER
34	SR	70	ASP
34	SR	76	ASP
34	SR	140	CYS
34	SR	150	TRP
34	SR	162	ALA
34	SR	163	ASP
34	SR	194	GLY
34	SR	242	SER
35	SM	81	THR
35	SM	102	THR
39	L2	47	GLN
39	L2	152	SER
40	L3	3	HIS
40	L3	171	LEU
40	L3	299	ASP
40	L3	300	ARG
40	L3	347	SER
41	L4	15	ALA
41	L4	64	SER
41	L4	103	THR
41	L4	181	VAL
41	L4	190	GLY
42	L5	58	LYS
42	L5	157	ALA
42	L5	213	ASP
42	L5	259	LYS
42	L5	260	PHE

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Mol	Chain	Res	Type
42	L5	276	LYS
44	L7	158	LYS
45	L8	135	GLY
45	L8	227	ASP
46	L9	117	PHE
46	L9	120	ASP
47	M0	16	PRO
47	M0	82	ARG
47	M0	161	GLY
47	M0	218	ALA
48	M1	38	GLU
48	M1	151	SER
49	M3	58	VAL
49	M3	62	THR
49	M3	176	GLU
50	M4	29	ALA
50	M4	105	GLN
51	M5	94	TYR
51	M5	192	LYS
52	M6	63	ALA
52	M6	188	SER
53	M7	3	ARG
53	M7	30	ARG
53	M7	51	VAL
53	M7	161	ALA
54	M8	74	GLU
55	M9	97	ARG
57	N1	132	PRO
58	N2	84	LEU
59	N3	16	GLY
59	N3	67	PRO
60	N4	34	SER
60	N4	97	LYS
62	N6	3	LYS
62	N6	52	ARG
62	N6	92	GLY
63	N7	3	LYS
64	N8	96	LYS
67	O1	7	VAL
72	O6	34	SER
72	O6	96	ALA
72	O6	97	SER

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Mol	Chain	Res	Type
74	O8	33	LYS
76	Q0	79	GLU
76	Q0	114	LYS
78	Q2	34	SER
2	s0	7	PHE
2	s0	10	THR
2	s0	152	PRO
2	s0	186	GLY
3	s1	94	LYS
3	s1	154	SER
4	s2	83	ILE
4	s2	91	ARG
4	s2	151	PRO
4	s2	196	VAL
4	s2	204	THR
4	s2	205	ARG
4	s2	238	SER
5	s3	61	GLU
5	s3	90	ARG
5	s3	93	ASP
5	s3	179	GLN
5	s3	204	ASP
6	s4	12	LEU
7	s5	34	GLN
7	s5	84	LYS
7	s5	98	MET
11	s9	43	TYR
11	s9	64	GLU
11	s9	78	ARG
11	s9	169	PRO
12	c0	30	ALA
13	c1	28	SER
14	c2	39	ASP
14	c2	45	LEU
14	c2	106	ILE
14	c2	108	ARG
14	c2	119	SER
14	c2	131	ASP
15	c3	29	SER
15	c3	137	PRO
16	c4	12	GLN
16	c4	96	PRO

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Mol	Chain	Res	Type
17	c5	8	LYS
17	c5	130	ARG
20	c8	65	GLU
21	c9	66	TYR
22	d0	49	ASN
25	d3	66	SER
25	d3	99	ASN
26	d4	49	LYS
28	d6	5	ARG
28	d6	8	ASN
28	d6	27	SER
29	d7	59	CYS
30	d8	33	LEU
30	d8	57	MET
80	e0	56	MET
33	e1	107	LYS
33	e1	111	GLU
33	e1	112	GLY
33	e1	125	THR
33	e1	131	PHE
33	e1	136	LYS
33	e1	145	HIS
34	sR	96	THR
34	sR	163	ASP
34	sR	186	PHE
34	sR	277	GLU
35	sM	171	LYS
39	l2	17	THR
39	l2	56	ALA
39	l2	70	ARG
39	l2	127	ALA
40	l3	12	GLY
40	l3	291	GLU
40	l3	302	LYS
41	l4	330	TYR
41	l4	361	HIS
42	l5	269	SER
43	l6	10	TYR
44	l7	53	LYS
44	l7	191	VAL
45	l8	69	LEU
45	l8	237	ILE

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Mol	Chain	Res	Type
45	l8	239	GLY
46	l9	2	LYS
47	m0	78	THR
47	m0	176	LEU
47	m0	207	GLU
47	m0	219	ALA
48	m1	82	ARG
48	m1	111	ASP
48	m1	151	SER
48	m1	153	LYS
49	m3	93	ILE
49	m3	163	GLY
50	m4	8	LYS
51	m5	49	ARG
52	m6	110	PRO
52	m6	162	VAL
54	m8	41	ASP
54	m8	69	ARG
55	m9	142	ILE
59	n3	4	ASN
59	n3	27	ASP
59	n3	28	ASN
59	n3	68	GLU
59	n3	69	LEU
60	n4	130	SER
61	n5	47	ALA
62	n6	126	LEU
63	n7	14	VAL
63	n7	41	ALA
63	n7	103	GLN
64	n8	110	GLY
65	n9	25	LYS
65	n9	33	LYS
67	o1	47	ASP
67	o1	82	GLU
67	o1	83	GLU
69	o3	59	VAL
70	o4	12	PRO
71	o5	16	GLN
71	o5	81	ARG
71	o5	105	ARG
76	q0	78	ILE

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Mol	Chain	Res	Type
77	q1	3	ALA
77	q1	4	LYS
78	q2	74	CYS
78	q2	78	LYS
79	q3	20	SER
81	p0	206	ASP
2	S0	7	PHE
2	S0	78	SER
2	S0	126	PRO
3	S1	156	ALA
4	S2	39	THR
4	S2	235	LEU
5	S3	118	ALA
5	S3	129	SER
6	S4	237	SER
7	S5	31	GLU
7	S5	36	ALA
7	S5	202	ALA
7	S5	210	ALA
9	S7	98	ILE
10	S8	136	SER
11	S9	56	ALA
11	S9	100	LYS
11	S9	119	ALA
11	S9	162	SER
13	C1	3	THR
13	C1	82	ARG
13	C1	153	PHE
15	C3	3	ARG
16	C4	51	ASP
16	C4	114	ARG
17	C5	24	LYS
19	C7	73	LEU
20	C8	118	LYS
20	C8	134	ARG
21	C9	25	GLN
22	D0	17	GLN
22	D0	55	PRO
22	D0	73	GLY
23	D1	7	GLN
23	D1	82	VAL
25	D3	4	GLY

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Mol	Chain	Res	Type
25	D3	41	SER
25	D3	139	LYS
27	D5	41	ILE
28	D6	18	VAL
28	D6	61	GLU
28	D6	64	LEU
28	D6	65	PRO
30	D8	16	LEU
30	D8	61	ARG
32	E0	23	LYS
33	E1	100	LEU
33	E1	137	ASP
34	SR	15	GLY
34	SR	189	GLU
35	SM	53	ARG
35	SM	100	THR
35	SM	119	ALA
35	SM	136	ALA
35	SM	168	GLU
35	SM	172	VAL
35	SM	173	GLU
35	SM	174	LEU
39	L2	153	GLY
39	L2	231	SER
40	L3	244	ARG
41	L4	140	HIS
41	L4	182	LEU
41	L4	184	SER
42	L5	137	ASP
42	L5	212	ALA
43	L6	6	ALA
44	L7	128	LYS
45	L8	136	LEU
45	L8	163	VAL
46	L9	5	GLN
46	L9	96	HIS
46	L9	110	LYS
46	L9	190	ASP
48	M1	114	ILE
48	M1	117	ASP
49	M3	76	THR
49	M3	128	ARG

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Mol	Chain	Res	Type
49	M3	164	GLU
49	M3	175	SER
51	M5	77	LYS
51	M5	81	TYR
53	M7	63	PHE
53	M7	133	HIS
54	M8	162	ALA
55	M9	3	ASN
55	M9	53	LYS
56	N0	24	LEU
56	N0	34	GLU
59	N3	46	LEU
59	N3	54	LEU
62	N6	38	GLU
63	N7	35	SER
63	N7	103	GLN
65	N9	33	LYS
67	O1	97	LEU
68	O2	122	PRO
68	O2	127	ALA
69	O3	59	VAL
70	O4	81	CYS
71	O5	75	TYR
72	O6	32	ALA
74	O8	37	PRO
77	Q1	3	ALA
78	Q2	78	LYS
2	s0	9	LEU
2	s0	49	ASN
2	s0	103	THR
3	s1	22	ASP
3	s1	60	ALA
4	s2	234	PRO
5	s3	43	PRO
5	s3	64	ARG
6	s4	30	ARG
6	s4	90	ILE
6	s4	117	GLU
6	s4	168	LYS
6	s4	223	ASN
7	s5	29	ILE
7	s5	118	LEU

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Mol	Chain	Res	Type
8	s6	216	LEU
9	s7	11	GLN
9	s7	41	LEU
9	s7	130	VAL
9	s7	133	THR
12	c0	23	ALA
12	c0	35	ILE
12	c0	54	TYR
14	c2	21	GLU
14	c2	40	GLY
14	c2	107	ASP
14	c2	127	GLY
15	c3	26	PHE
17	c5	12	PHE
17	c5	14	THR
17	c5	68	PRO
19	c7	103	ASP
20	c8	14	ILE
22	d0	120	SER
23	d1	9	VAL
23	d1	10	GLU
24	d2	68	ARG
24	d2	99	PHE
25	d3	13	ARG
26	d4	50	ALA
27	d5	81	ARG
28	d6	34	LYS
28	d6	59	TYR
30	d8	62	GLU
31	d9	11	PRO
33	e1	81	LYS
33	e1	85	TYR
33	e1	148	TYR
34	sR	166	SER
35	sM	43	ASP
35	sM	84	LYS
39	l2	80	GLU
39	l2	213	GLY
40	l3	155	ALA
40	l3	206	ASP
41	l4	197	ARG
41	l4	301	PRO

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Mol	Chain	Res	Type
41	l4	305	ALA
42	l5	189	GLU
42	l5	215	ASP
42	l5	245	GLU
43	l6	97	ASN
44	l7	159	GLN
45	l8	39	ALA
45	l8	196	ALA
46	l9	107	ASP
47	m0	27	PRO
47	m0	175	ASN
47	m0	195	ALA
47	m0	220	GLN
48	m1	115	LYS
49	m3	60	ALA
49	m3	62	THR
49	m3	135	ALA
50	m4	86	ALA
52	m6	196	ALA
54	m8	49	LEU
54	m8	97	PRO
54	m8	105	ARG
54	m8	149	ALA
55	m9	28	GLU
55	m9	183	ALA
56	n0	110	MET
56	n0	154	HIS
57	n1	58	GLN
58	n2	60	GLY
60	n4	83	THR
64	n8	48	TYR
66	o0	46	ALA
66	o0	47	ASN
68	o2	6	HIS
68	o2	27	ARG
69	o3	60	ARG
70	o4	32	ALA
71	o5	82	ALA
73	o7	65	ARG
75	o9	3	ALA
75	o9	18	LYS
78	q2	77	CYS

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Mol	Chain	Res	Type
81	p0	68	SER
3	S1	22	ASP
3	S1	59	ASP
3	S1	101	HIS
3	S1	131	ASP
3	S1	210	ILE
4	S2	47	ALA
5	S3	72	LEU
5	S3	216	PRO
6	S4	5	PRO
6	S4	11	ARG
11	S9	110	GLN
11	S9	147	MET
14	C2	23	THR
14	C2	83	GLU
17	C5	38	PRO
18	C6	33	GLY
19	C7	72	LYS
21	C9	122	ARG
22	D0	119	ALA
23	D1	42	GLU
24	D2	100	GLY
25	D3	3	LYS
25	D3	20	ARG
25	D3	21	ASN
26	D4	54	ALA
31	D9	11	PRO
34	SR	98	GLU
34	SR	120	SER
34	SR	152	SER
34	SR	237	GLN
34	SR	247	PRO
35	SM	12	VAL
35	SM	85	SER
35	SM	155	LEU
39	L2	127	ALA
40	L3	141	GLY
40	L3	317	ILE
41	L4	14	GLU
42	L5	251	PRO
43	L6	36	PRO
44	L7	25	GLN

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Mol	Chain	Res	Type
44	L7	54	GLU
44	L7	163	LEU
44	L7	217	PRO
45	L8	80	TYR
45	L8	122	LYS
48	M1	108	GLU
49	M3	46	ILE
50	M4	28	SER
52	M6	16	VAL
53	M7	75	GLU
55	M9	131	ALA
57	N1	127	GLN
62	N6	126	LEU
63	N7	36	HIS
64	N8	76	ASP
67	O1	60	TRP
69	O3	90	PRO
72	O6	12	ASN
72	O6	36	ARG
72	O6	78	GLY
78	Q2	104	LEU
79	Q3	12	GLY
79	Q3	91	GLU
2	s0	162	CYS
5	s3	45	LYS
5	s3	56	GLN
7	s5	21	THR
7	s5	57	SER
7	s5	74	ALA
8	s6	165	GLY
10	s8	20	GLN
10	s8	52	ASN
10	s8	78	ILE
11	s9	162	SER
12	c0	26	ASP
12	c0	82	LEU
14	c2	66	VAL
14	c2	90	LYS
14	c2	91	VAL
15	c3	12	SER
15	c3	22	ALA
15	c3	148	ALA

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Mol	Chain	Res	Type
16	c4	131	GLY
17	c5	31	GLU
18	c6	61	SER
18	c6	97	VAL
19	c7	86	PRO
20	c8	60	GLU
21	c9	49	ASP
22	d0	53	LYS
22	d0	96	PRO
35	sM	46	LYS
39	l2	98	VAL
41	l4	146	PRO
41	l4	176	SER
41	l4	328	ASN
41	l4	353	ALA
42	l5	174	PRO
42	l5	258	LYS
43	l6	31	ARG
44	l7	27	ALA
44	l7	217	PRO
46	l9	30	PRO
46	l9	31	ARG
46	l9	167	VAL
48	m1	12	LEU
48	m1	117	ASP
49	m3	51	LEU
53	m7	6	ALA
57	n1	83	ARG
57	n1	148	PRO
60	n4	59	HIS
60	n4	132	GLY
61	n5	97	LYS
63	n7	124	ALA
64	n8	70	LYS
65	n9	37	PRO
67	o1	44	MET
68	o2	69	SER
68	o2	70	GLY
72	o6	4	LYS
72	o6	33	ALA
79	q3	3	LYS
79	q3	17	ARG

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Mol	Chain	Res	Type
2	S0	189	VAL
3	S1	21	VAL
4	S2	153	SER
4	S2	248	SER
7	S5	26	ALA
8	S6	146	GLY
10	S8	10	LYS
12	C0	92	ILE
13	C1	5	LEU
14	C2	108	ARG
15	C3	79	GLY
23	D1	10	GLU
23	D1	46	ILE
26	D4	34	ASN
26	D4	133	ASN
34	SR	113	VAL
41	L4	328	ASN
44	L7	91	GLY
44	L7	178	ILE
46	L9	4	ILE
47	M0	47	PRO
49	M3	136	GLU
52	M6	139	GLY
54	M8	84	VAL
54	M8	99	THR
56	N0	167	ARG
61	N5	44	PRO
64	N8	15	VAL
64	N8	70	LYS
64	N8	116	GLY
66	O0	96	GLY
68	O2	45	ARG
76	Q0	88	LYS
2	s0	139	VAL
2	s0	185	ARG
4	s2	93	GLY
4	s2	236	PRO
6	s4	241	GLY
9	s7	177	THR
11	s9	42	ILE
14	c2	103	LEU
16	c4	11	SER

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Mol	Chain	Res	Type
16	c4	50	ALA
17	c5	6	ASN
18	c6	40	GLU
19	c7	98	GLY
21	c9	118	PRO
23	d1	44	ARG
24	d2	8	ALA
26	d4	78	SER
28	d6	35	ALA
30	d8	58	GLU
80	e0	47	VAL
80	e0	54	ARG
41	l4	331	ALA
44	l7	54	GLU
45	l8	34	PHE
46	l9	172	ILE
49	m3	13	HIS
52	m6	5	PRO
53	m7	37	ASN
59	n3	3	GLY
64	n8	47	LYS
64	n8	120	ASN
67	o1	86	LYS
68	o2	17	PHE
70	o4	59	PRO
81	p0	33	VAL
2	S0	50	VAL
4	S2	74	PRO
7	S5	150	GLY
30	D8	23	GLY
34	SR	67	ILE
60	N4	63	ILE
62	N6	101	PRO
65	N9	21	ILE
68	O2	108	ILE
6	s4	135	GLY
9	s7	73	VAL
14	c2	63	VAL
14	c2	82	PRO
22	d0	97	VAL
80	e0	60	PRO
35	sM	167	PRO

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Mol	Chain	Res	Type
42	l5	125	VAL
54	m8	43	PRO
59	n3	13	ILE
68	o2	122	PRO
4	S2	182	PRO
6	S4	111	VAL
14	C2	115	VAL
34	SR	63	GLY
39	L2	15	ILE
57	N1	123	GLY
62	N6	49	PRO
68	O2	124	GLY
14	c2	22	VAL
16	c4	39	ILE
29	d7	4	VAL
66	o0	96	GLY
81	p0	101	VAL
3	S1	93	GLY
8	S6	181	PRO
14	C2	66	VAL
45	L8	30	THR
47	M0	117	GLY
49	M3	132	ALA
54	M8	160	GLY
54	M8	183	GLY
55	M9	57	VAL
58	N2	22	PRO
60	N4	90	ILE
63	N7	70	PRO
69	O3	76	GLY
79	Q3	50	GLY
4	s2	68	ILE
8	s6	69	LEU
18	c6	4	VAL
21	c9	100	ILE
34	sR	138	GLY
35	sM	52	PRO
40	l3	141	GLY
40	l3	186	GLY
28	D6	59	TYR
41	L4	4	PRO
49	M3	33	VAL

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Mol	Chain	Res	Type
64	N8	29	PRO
65	N9	29	TYR
5	s3	81	PRO
7	s5	89	ILE
18	c6	33	GLY
20	c8	69	ILE
34	sR	247	PRO
39	l2	49	VAL
44	l7	89	ILE
44	l7	204	PRO
44	l7	210	PRO
44	l7	240	VAL
45	l8	119	GLY
47	m0	16	PRO
79	q3	71	VAL
81	p0	30	VAL
11	S9	168	ARG
14	C2	117	GLY
25	D3	96	VAL
45	L8	167	PRO
55	M9	14	VAL
60	N4	76	VAL
41	l4	190	GLY
76	q0	123	PRO

5.3.2 Protein sidechains ⓘ

In the following table, the Percentiles column shows the percent sidechain outliers of the chain as a percentile score with respect to all X-ray entries followed by that with respect to entries of similar resolution.

The Analysed column shows the number of residues for which the sidechain conformation was analysed, and the total number of residues.

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
2	S0	164/209 (78%)	125 (76%)	39 (24%)	1	4
2	s0	165/209 (79%)	135 (82%)	30 (18%)	2	10
3	S1	191/223 (86%)	146 (76%)	45 (24%)	1	4
3	s1	192/223 (86%)	147 (77%)	45 (23%)	1	4
4	S2	176/204 (86%)	136 (77%)	40 (23%)	1	4

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
4	s2	176/204 (86%)	128 (73%)	48 (27%)	0	3
5	S3	182/194 (94%)	140 (77%)	42 (23%)	1	4
5	s3	182/194 (94%)	140 (77%)	42 (23%)	1	4
6	S4	221/221 (100%)	175 (79%)	46 (21%)	1	6
6	s4	221/221 (100%)	173 (78%)	48 (22%)	1	5
7	S5	173/190 (91%)	139 (80%)	34 (20%)	1	7
7	s5	173/190 (91%)	136 (79%)	37 (21%)	1	5
8	S6	188/201 (94%)	151 (80%)	37 (20%)	1	7
8	s6	187/201 (93%)	153 (82%)	34 (18%)	2	10
9	S7	165/169 (98%)	135 (82%)	30 (18%)	2	10
9	s7	165/169 (98%)	130 (79%)	35 (21%)	1	6
10	S8	150/161 (93%)	123 (82%)	27 (18%)	2	11
10	s8	150/161 (93%)	127 (85%)	23 (15%)	3	18
11	S9	158/165 (96%)	119 (75%)	39 (25%)	1	4
11	s9	158/165 (96%)	128 (81%)	30 (19%)	2	9
12	C0	77/98 (79%)	61 (79%)	16 (21%)	1	6
12	c0	73/98 (74%)	60 (82%)	13 (18%)	2	11
13	C1	129/136 (95%)	105 (81%)	24 (19%)	2	9
13	c1	129/136 (95%)	108 (84%)	21 (16%)	3	15
14	C2	88/118 (75%)	67 (76%)	21 (24%)	1	4
14	c2	88/118 (75%)	71 (81%)	17 (19%)	1	8
15	C3	127/127 (100%)	96 (76%)	31 (24%)	1	4
15	c3	127/127 (100%)	102 (80%)	25 (20%)	1	7
16	C4	81/104 (78%)	67 (83%)	14 (17%)	2	12
16	c4	97/104 (93%)	76 (78%)	21 (22%)	1	5
17	C5	101/117 (86%)	79 (78%)	22 (22%)	1	5
17	c5	103/117 (88%)	85 (82%)	18 (18%)	2	12
18	C6	117/118 (99%)	97 (83%)	20 (17%)	2	13
18	c6	118/118 (100%)	96 (81%)	22 (19%)	2	9
19	C7	94/124 (76%)	70 (74%)	24 (26%)	0	3
19	c7	92/124 (74%)	72 (78%)	20 (22%)	1	5

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
20	C8	128/128 (100%)	95 (74%)	33 (26%)	0	3
20	c8	128/128 (100%)	94 (73%)	34 (27%)	0	3
21	C9	115/115 (100%)	86 (75%)	29 (25%)	0	3
21	c9	115/115 (100%)	96 (84%)	19 (16%)	2	14
22	D0	100/113 (88%)	76 (76%)	24 (24%)	1	4
22	d0	103/113 (91%)	72 (70%)	31 (30%)	0	2
23	D1	74/74 (100%)	61 (82%)	13 (18%)	2	11
23	d1	74/74 (100%)	58 (78%)	16 (22%)	1	5
24	D2	110/110 (100%)	90 (82%)	20 (18%)	2	10
24	d2	110/110 (100%)	89 (81%)	21 (19%)	2	8
25	D3	119/119 (100%)	95 (80%)	24 (20%)	1	7
25	d3	119/119 (100%)	91 (76%)	28 (24%)	1	4
26	D4	112/112 (100%)	93 (83%)	19 (17%)	2	13
26	d4	112/112 (100%)	93 (83%)	19 (17%)	2	13
27	D5	61/88 (69%)	46 (75%)	15 (25%)	1	4
27	d5	61/88 (69%)	48 (79%)	13 (21%)	1	6
28	D6	83/83 (100%)	61 (74%)	22 (26%)	0	3
28	d6	83/83 (100%)	65 (78%)	18 (22%)	1	5
29	D7	70/70 (100%)	59 (84%)	11 (16%)	3	17
29	d7	70/70 (100%)	55 (79%)	15 (21%)	1	5
30	D8	56/59 (95%)	45 (80%)	11 (20%)	1	8
30	d8	56/59 (95%)	43 (77%)	13 (23%)	1	4
31	D9	47/48 (98%)	40 (85%)	7 (15%)	3	19
31	d9	47/48 (98%)	36 (77%)	11 (23%)	1	4
32	E0	51/51 (100%)	42 (82%)	9 (18%)	2	11
33	E1	62/66 (94%)	48 (77%)	14 (23%)	1	5
33	e1	66/66 (100%)	49 (74%)	17 (26%)	0	3
34	SR	260/261 (100%)	223 (86%)	37 (14%)	4	21
34	sR	260/261 (100%)	222 (85%)	38 (15%)	3	20
35	SM	97/228 (42%)	72 (74%)	25 (26%)	0	3
35	sM	54/228 (24%)	45 (83%)	9 (17%)	2	14

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
39	L2	193/195 (99%)	143 (74%)	50 (26%)	0	3
39	l2	192/195 (98%)	144 (75%)	48 (25%)	1	4
40	L3	320/322 (99%)	256 (80%)	64 (20%)	1	7
40	l3	320/322 (99%)	248 (78%)	72 (22%)	1	5
41	L4	288/288 (100%)	223 (77%)	65 (23%)	1	5
41	l4	288/288 (100%)	229 (80%)	59 (20%)	1	7
42	L5	244/244 (100%)	191 (78%)	53 (22%)	1	5
42	l5	243/244 (100%)	198 (82%)	45 (18%)	2	9
43	L6	134/152 (88%)	108 (81%)	26 (19%)	1	8
43	l6	135/152 (89%)	102 (76%)	33 (24%)	1	4
44	L7	186/204 (91%)	154 (83%)	32 (17%)	2	13
44	l7	187/204 (92%)	152 (81%)	35 (19%)	2	9
45	L8	187/207 (90%)	150 (80%)	37 (20%)	1	7
45	l8	177/207 (86%)	141 (80%)	36 (20%)	1	7
46	L9	171/171 (100%)	127 (74%)	44 (26%)	0	3
46	l9	171/171 (100%)	128 (75%)	43 (25%)	0	3
47	M0	177/186 (95%)	137 (77%)	40 (23%)	1	5
47	m0	179/186 (96%)	144 (80%)	35 (20%)	1	8
48	M1	147/150 (98%)	112 (76%)	35 (24%)	1	4
48	m1	147/150 (98%)	111 (76%)	36 (24%)	1	4
49	M3	154/158 (98%)	122 (79%)	32 (21%)	1	6
49	m3	154/158 (98%)	120 (78%)	34 (22%)	1	5
50	M4	107/108 (99%)	85 (79%)	22 (21%)	1	6
50	m4	108/108 (100%)	86 (80%)	22 (20%)	1	7
51	M5	175/175 (100%)	132 (75%)	43 (25%)	1	4
51	m5	175/175 (100%)	141 (81%)	34 (19%)	1	8
52	M6	160/161 (99%)	121 (76%)	39 (24%)	1	4
52	m6	160/161 (99%)	120 (75%)	40 (25%)	1	4
53	M7	140/145 (97%)	111 (79%)	29 (21%)	1	6
53	m7	125/145 (86%)	101 (81%)	24 (19%)	1	8
54	M8	150/150 (100%)	116 (77%)	34 (23%)	1	4

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
54	m8	150/150 (100%)	116 (77%)	34 (23%)	1	4
55	M9	153/153 (100%)	124 (81%)	29 (19%)	2	9
55	m9	153/153 (100%)	125 (82%)	28 (18%)	2	10
56	N0	156/156 (100%)	122 (78%)	34 (22%)	1	5
56	n0	156/156 (100%)	122 (78%)	34 (22%)	1	5
57	N1	136/136 (100%)	102 (75%)	34 (25%)	1	4
57	n1	136/136 (100%)	107 (79%)	29 (21%)	1	6
58	N2	87/106 (82%)	73 (84%)	14 (16%)	3	16
58	n2	85/106 (80%)	58 (68%)	27 (32%)	0	2
59	N3	104/104 (100%)	84 (81%)	20 (19%)	1	8
59	n3	104/104 (100%)	85 (82%)	19 (18%)	2	10
60	N4	57/129 (44%)	47 (82%)	10 (18%)	2	12
60	n4	100/129 (78%)	79 (79%)	21 (21%)	1	6
61	N5	104/117 (89%)	78 (75%)	26 (25%)	1	4
61	n5	104/117 (89%)	85 (82%)	19 (18%)	2	10
62	N6	109/109 (100%)	88 (81%)	21 (19%)	1	8
62	n6	109/109 (100%)	85 (78%)	24 (22%)	1	5
63	N7	115/115 (100%)	94 (82%)	21 (18%)	2	10
63	n7	115/115 (100%)	87 (76%)	28 (24%)	1	4
64	N8	118/118 (100%)	98 (83%)	20 (17%)	2	13
64	n8	118/118 (100%)	91 (77%)	27 (23%)	1	4
65	N9	46/46 (100%)	35 (76%)	11 (24%)	1	4
65	n9	46/46 (100%)	30 (65%)	16 (35%)	0	2
66	O0	81/87 (93%)	66 (82%)	15 (18%)	2	9
66	o0	84/87 (97%)	69 (82%)	15 (18%)	2	11
67	O1	92/96 (96%)	69 (75%)	23 (25%)	1	4
67	o1	94/96 (98%)	63 (67%)	31 (33%)	0	2
68	O2	109/110 (99%)	87 (80%)	22 (20%)	1	7
68	o2	109/110 (99%)	81 (74%)	28 (26%)	0	3
69	O3	90/90 (100%)	71 (79%)	19 (21%)	1	6
69	o3	90/90 (100%)	75 (83%)	15 (17%)	2	14

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
70	O4	95/102 (93%)	75 (79%)	20 (21%)	1	6
70	o4	95/102 (93%)	82 (86%)	13 (14%)	4	22
71	O5	104/104 (100%)	83 (80%)	21 (20%)	1	7
71	o5	103/104 (99%)	76 (74%)	27 (26%)	0	3
72	O6	81/81 (100%)	56 (69%)	25 (31%)	0	2
72	o6	80/81 (99%)	53 (66%)	27 (34%)	0	2
73	O7	70/70 (100%)	55 (79%)	15 (21%)	1	5
73	o7	70/70 (100%)	52 (74%)	18 (26%)	0	3
74	O8	68/68 (100%)	50 (74%)	18 (26%)	0	3
74	o8	67/68 (98%)	49 (73%)	18 (27%)	0	3
75	O9	45/45 (100%)	40 (89%)	5 (11%)	7	31
75	o9	45/45 (100%)	33 (73%)	12 (27%)	0	3
76	Q0	47/47 (100%)	41 (87%)	6 (13%)	5	25
76	q0	47/47 (100%)	32 (68%)	15 (32%)	0	2
77	Q1	23/23 (100%)	16 (70%)	7 (30%)	0	2
77	q1	23/23 (100%)	14 (61%)	9 (39%)	0	1
78	Q2	90/90 (100%)	67 (74%)	23 (26%)	0	3
78	q2	90/90 (100%)	70 (78%)	20 (22%)	1	5
79	Q3	71/71 (100%)	55 (78%)	16 (22%)	1	5
79	q3	71/71 (100%)	54 (76%)	17 (24%)	1	4
80	e0	53/53 (100%)	38 (72%)	15 (28%)	0	3
81	p0	105/253 (42%)	84 (80%)	21 (20%)	1	7
All	All	18728/20241 (92%)	14710 (78%)	4018 (22%)	1	5

All (4018) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
2	S0	6	THR
2	S0	10	THR
2	S0	12	GLU
2	S0	18	LEU
2	S0	32	HIS
2	S0	37	VAL
2	S0	49	ASN

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Mol	Chain	Res	Type
2	S0	50	VAL
2	S0	59	LEU
2	S0	62	ARG
2	S0	71	GLU
2	S0	78	SER
2	S0	84	ARG
2	S0	86	VAL
2	S0	88	LYS
2	S0	96	THR
2	S0	98	ILE
2	S0	103	THR
2	S0	110	TYR
2	S0	112	THR
2	S0	123	VAL
2	S0	124	THR
2	S0	140	ASN
2	S0	144	ILE
2	S0	146	LEU
2	S0	153	SER
2	S0	156	VAL
2	S0	157	ASP
2	S0	162	CYS
2	S0	168	HIS
2	S0	172	LEU
2	S0	177	LEU
2	S0	184	LEU
2	S0	185	ARG
2	S0	188	LEU
2	S0	196	SER
2	S0	197	ILE
2	S0	198	MET
2	S0	200	ASP
3	S1	21	VAL
3	S1	22	ASP
3	S1	25	THR
3	S1	29	TRP
3	S1	30	PHE
3	S1	39	GLU
3	S1	46	THR
3	S1	51	SER
3	S1	61	LEU
3	S1	65	VAL

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Mol	Chain	Res	Type
3	S1	70	LEU
3	S1	72	ASP
3	S1	74	GLN
3	S1	76	SER
3	S1	77	GLU
3	S1	78	ASP
3	S1	79	HIS
3	S1	81	PHE
3	S1	89	ASP
3	S1	97	LEU
3	S1	105	PHE
3	S1	111	ARG
3	S1	115	ARG
3	S1	116	LYS
3	S1	125	VAL
3	S1	126	THR
3	S1	135	LEU
3	S1	140	ILE
3	S1	148	ASN
3	S1	149	GLN
3	S1	170	GLU
3	S1	173	THR
3	S1	180	THR
3	S1	181	LEU
3	S1	184	LEU
3	S1	191	GLU
3	S1	198	GLU
3	S1	202	LYS
3	S1	212	VAL
3	S1	215	VAL
3	S1	218	LEU
3	S1	219	LYS
3	S1	222	LYS
3	S1	223	PHE
3	S1	229	MET
4	S2	41	LEU
4	S2	54	GLU
4	S2	58	LEU
4	S2	69	ILE
4	S2	72	LEU
4	S2	73	LEU
4	S2	77	GLN

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Mol	Chain	Res	Type
4	S2	81	MET
4	S2	87	GLN
4	S2	89	GLN
4	S2	90	THR
4	S2	95	ARG
4	S2	96	THR
4	S2	97	ARG
4	S2	106	ASP
4	S2	108	ASN
4	S2	111	VAL
4	S2	113	LEU
4	S2	117	THR
4	S2	134	LEU
4	S2	140	ARG
4	S2	141	ARG
4	S2	146	THR
4	S2	147	ASN
4	S2	148	LEU
4	S2	166	THR
4	S2	198	THR
4	S2	201	ASN
4	S2	207	LEU
4	S2	210	THR
4	S2	221	THR
4	S2	222	TYR
4	S2	225	LEU
4	S2	226	THR
4	S2	227	PRO
4	S2	235	LEU
4	S2	237	VAL
4	S2	238	SER
4	S2	240	LEU
4	S2	245	ASP
5	S3	4	LEU
5	S3	10	LYS
5	S3	14	ASP
5	S3	21	LEU
5	S3	23	GLU
5	S3	26	THR
5	S3	37	VAL
5	S3	44	THR
5	S3	59	LEU

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Mol	Chain	Res	Type
5	S3	64	ARG
5	S3	65	ARG
5	S3	76	ARG
5	S3	84	ILE
5	S3	92	GLN
5	S3	94	ARG
5	S3	105	MET
5	S3	111	ASN
5	S3	117	ARG
5	S3	120	TYR
5	S3	143	ARG
5	S3	146	ARG
5	S3	151	LYS
5	S3	154	ASP
5	S3	158	ILE
5	S3	172	THR
5	S3	175	VAL
5	S3	176	LEU
5	S3	177	MET
5	S3	178	ARG
5	S3	179	GLN
5	S3	181	VAL
5	S3	190	ARG
5	S3	195	SER
5	S3	196	ARG
5	S3	202	LEU
5	S3	209	ILE
5	S3	210	GLU
5	S3	218	LEU
5	S3	220	PRO
5	S3	222	VAL
5	S3	223	LYS
5	S3	225	TYR
6	S4	6	LYS
6	S4	7	LYS
6	S4	12	LEU
6	S4	23	LEU
6	S4	26	CYS
6	S4	38	LEU
6	S4	42	LEU
6	S4	45	ILE
6	S4	48	LEU

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Mol	Chain	Res	Type
6	S4	52	LEU
6	S4	56	LEU
6	S4	65	LEU
6	S4	68	ARG
6	S4	69	HIS
6	S4	70	VAL
6	S4	76	VAL
6	S4	77	ARG
6	S4	78	THR
6	S4	81	THR
6	S4	87	MET
6	S4	95	THR
6	S4	116	ASP
6	S4	123	LEU
6	S4	128	LYS
6	S4	129	VAL
6	S4	133	LYS
6	S4	158	ASP
6	S4	160	VAL
6	S4	170	THR
6	S4	180	LEU
6	S4	187	ARG
6	S4	192	ILE
6	S4	206	ASP
6	S4	211	LYS
6	S4	214	LEU
6	S4	215	ASP
6	S4	225	VAL
6	S4	226	PHE
6	S4	227	VAL
6	S4	228	ILE
6	S4	240	LYS
6	S4	242	LYS
6	S4	247	SER
6	S4	248	ILE
6	S4	259	GLN
6	S4	261	LEU
7	S5	23	VAL
7	S5	25	LEU
7	S5	38	THR
7	S5	39	GLU
7	S5	40	ILE

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Mol	Chain	Res	Type
7	S5	41	LYS
7	S5	42	LEU
7	S5	43	PHE
7	S5	45	LYS
7	S5	50	GLU
7	S5	53	VAL
7	S5	63	GLN
7	S5	76	ARG
7	S5	79	ASN
7	S5	86	GLN
7	S5	87	CYS
7	S5	89	ILE
7	S5	90	ILE
7	S5	93	LEU
7	S5	94	THR
7	S5	97	LEU
7	S5	99	MET
7	S5	112	ARG
7	S5	119	ASP
7	S5	122	ASN
7	S5	123	VAL
7	S5	148	ARG
7	S5	156	ARG
7	S5	157	ARG
7	S5	161	ASP
7	S5	169	ASN
7	S5	186	ASN
7	S5	219	ARG
7	S5	225	ARG
8	S6	12	SER
8	S6	13	GLN
8	S6	15	THR
8	S6	16	PHE
8	S6	25	ARG
8	S6	44	GLU
8	S6	51	LYS
8	S6	52	ILE
8	S6	58	LYS
8	S6	67	VAL
8	S6	74	LYS
8	S6	76	LEU
8	S6	78	THR

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Mol	Chain	Res	Type
8	S6	79	LYS
8	S6	81	VAL
8	S6	82	SER
8	S6	89	ASP
8	S6	109	LEU
8	S6	115	LYS
8	S6	120	GLU
8	S6	125	THR
8	S6	128	THR
8	S6	133	LEU
8	S6	151	ASP
8	S6	155	ASP
8	S6	162	VAL
8	S6	168	THR
8	S6	169	TYR
8	S6	170	THR
8	S6	175	ILE
8	S6	177	ARG
8	S6	180	THR
8	S6	181	PRO
8	S6	193	LEU
8	S6	211	LEU
8	S6	212	LEU
8	S6	223	LYS
9	S7	7	LYS
9	S7	11	GLN
9	S7	15	GLU
9	S7	19	GLN
9	S7	28	GLU
9	S7	37	GLU
9	S7	38	LEU
9	S7	39	ARG
9	S7	49	ILE
9	S7	51	VAL
9	S7	60	ILE
9	S7	70	PHE
9	S7	71	HIS
9	S7	77	LEU
9	S7	79	ARG
9	S7	85	PHE
9	S7	92	PHE
9	S7	95	GLU

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Mol	Chain	Res	Type
9	S7	97	ARG
9	S7	109	VAL
9	S7	114	ARG
9	S7	126	LEU
9	S7	129	LEU
9	S7	130	VAL
9	S7	144	VAL
9	S7	147	ASN
9	S7	166	LEU
9	S7	167	GLU
9	S7	181	ILE
9	S7	185	ILE
10	S8	7	SER
10	S8	9	HIS
10	S8	14	THR
10	S8	21	PHE
10	S8	22	ARG
10	S8	25	ARG
10	S8	26	LYS
10	S8	29	LEU
10	S8	43	ILE
10	S8	45	SER
10	S8	46	VAL
10	S8	49	ARG
10	S8	58	LEU
10	S8	59	ARG
10	S8	66	SER
10	S8	72	ILE
10	S8	97	THR
10	S8	98	LYS
10	S8	103	GLN
10	S8	107	THR
10	S8	110	ARG
10	S8	138	ASN
10	S8	140	GLU
10	S8	152	ILE
10	S8	164	ARG
10	S8	187	GLU
10	S8	196	LEU
11	S9	3	ARG
11	S9	6	ARG
11	S9	7	THR

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Mol	Chain	Res	Type
11	S9	14	THR
11	S9	17	ARG
11	S9	21	SER
11	S9	28	LEU
11	S9	36	LEU
11	S9	39	LYS
11	S9	49	LEU
11	S9	63	ASP
11	S9	74	ASN
11	S9	77	ILE
11	S9	78	ARG
11	S9	79	ARG
11	S9	82	ARG
11	S9	89	ASP
11	S9	92	LYS
11	S9	93	LEU
11	S9	97	LEU
11	S9	99	LEU
11	S9	101	VAL
11	S9	105	LEU
11	S9	109	LEU
11	S9	110	GLN
11	S9	115	LYS
11	S9	118	LEU
11	S9	122	VAL
11	S9	126	ARG
11	S9	134	ILE
11	S9	138	LYS
11	S9	140	ILE
11	S9	149	ARG
11	S9	157	ASP
11	S9	168	ARG
11	S9	171	ARG
11	S9	172	VAL
11	S9	174	ARG
11	S9	182	GLU
12	C0	5	LYS
12	C0	8	ARG
12	C0	20	VAL
12	C0	25	LYS
12	C0	27	PHE
12	C0	28	ASN

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Mol	Chain	Res	Type
12	C0	29	GLN
12	C0	31	LYS
12	C0	33	GLU
12	C0	46	LEU
12	C0	55	VAL
12	C0	56	LYS
12	C0	71	GLU
12	C0	76	LEU
12	C0	79	TYR
12	C0	82	LEU
13	C1	18	HIS
13	C1	21	ASN
13	C1	29	LYS
13	C1	36	LYS
13	C1	40	LEU
13	C1	44	THR
13	C1	56	LYS
13	C1	67	ARG
13	C1	69	LYS
13	C1	72	THR
13	C1	75	VAL
13	C1	80	MET
13	C1	83	THR
13	C1	88	ARG
13	C1	90	TYR
13	C1	91	LEU
13	C1	94	ILE
13	C1	101	GLU
13	C1	105	LYS
13	C1	109	VAL
13	C1	123	VAL
13	C1	129	ARG
13	C1	131	ILE
13	C1	140	VAL
14	C2	43	ARG
14	C2	53	THR
14	C2	62	LEU
14	C2	63	VAL
14	C2	66	VAL
14	C2	71	ILE
14	C2	74	LEU
14	C2	81	ASP

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Mol	Chain	Res	Type
14	C2	86	VAL
14	C2	89	ILE
14	C2	97	LEU
14	C2	103	LEU
14	C2	119	SER
14	C2	126	TRP
14	C2	129	GLU
14	C2	132	GLU
14	C2	136	ILE
14	C2	139	HIS
14	C2	140	PHE
14	C2	141	SER
14	C2	143	GLN
15	C3	3	ARG
15	C3	11	ILE
15	C3	16	ILE
15	C3	27	LYS
15	C3	31	GLU
15	C3	32	SER
15	C3	35	GLU
15	C3	39	LYS
15	C3	61	THR
15	C3	64	ARG
15	C3	70	LYS
15	C3	72	MET
15	C3	76	LYS
15	C3	77	SER
15	C3	80	LEU
15	C3	83	GLU
15	C3	88	LEU
15	C3	99	ARG
15	C3	102	LEU
15	C3	103	GLU
15	C3	105	ASN
15	C3	106	ARG
15	C3	107	LYS
15	C3	109	LYS
15	C3	115	LEU
15	C3	120	SER
15	C3	125	LEU
15	C3	131	THR
15	C3	140	LYS

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Mol	Chain	Res	Type
15	C3	141	TYR
15	C3	142	GLU
16	C4	14	PHE
16	C4	22	SER
16	C4	26	THR
16	C4	29	HIS
16	C4	39	ILE
16	C4	42	VAL
16	C4	55	SER
16	C4	86	THR
16	C4	89	THR
16	C4	92	LYS
16	C4	103	ARG
16	C4	123	SER
16	C4	124	ASP
16	C4	137	LEU
17	C5	14	THR
17	C5	20	VAL
17	C5	22	LEU
17	C5	26	LEU
17	C5	28	MET
17	C5	29	SER
17	C5	31	GLU
17	C5	34	VAL
17	C5	35	LYS
17	C5	36	LEU
17	C5	40	ARG
17	C5	43	ARG
17	C5	89	MET
17	C5	90	ILE
17	C5	94	VAL
17	C5	98	ASN
17	C5	106	GLU
17	C5	110	GLU
17	C5	121	ILE
17	C5	124	THR
17	C5	128	HIS
17	C5	130	ARG
18	C6	4	VAL
18	C6	13	LYS
18	C6	19	VAL
18	C6	28	LEU

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Mol	Chain	Res	Type
18	C6	45	ARG
18	C6	48	VAL
18	C6	50	GLU
18	C6	54	LEU
18	C6	57	LEU
18	C6	66	ARG
18	C6	85	ILE
18	C6	89	LEU
18	C6	105	LEU
18	C6	109	PHE
18	C6	114	ARG
18	C6	121	SER
18	C6	123	ARG
18	C6	127	LYS
18	C6	128	LYS
18	C6	137	ARG
19	C7	3	ARG
19	C7	5	ARG
19	C7	6	THR
19	C7	7	LYS
19	C7	8	THR
19	C7	25	THR
19	C7	34	LEU
19	C7	38	ILE
19	C7	40	THR
19	C7	43	SER
19	C7	46	LEU
19	C7	49	LYS
19	C7	54	THR
19	C7	58	MET
19	C7	62	GLN
19	C7	69	ILE
19	C7	72	LYS
19	C7	78	ARG
19	C7	83	GLN
19	C7	84	TYR
19	C7	88	VAL
19	C7	107	SER
19	C7	113	LEU
19	C7	115	LEU
20	C8	3	LEU
20	C8	5	VAL

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Mol	Chain	Res	Type
20	C8	6	GLN
20	C8	8	GLN
20	C8	10	SER
20	C8	11	PHE
20	C8	13	HIS
20	C8	14	ILE
20	C8	15	LEU
20	C8	17	LEU
20	C8	20	THR
20	C8	32	LEU
20	C8	40	ARG
20	C8	47	CYS
20	C8	53	ASP
20	C8	54	LEU
20	C8	61	LEU
20	C8	71	GLN
20	C8	77	THR
20	C8	80	LYS
20	C8	82	PRO
20	C8	89	GLN
20	C8	92	ILE
20	C8	93	THR
20	C8	115	ARG
20	C8	119	ILE
20	C8	120	ARG
20	C8	132	ARG
20	C8	136	GLN
20	C8	138	THR
20	C8	141	THR
20	C8	143	ARG
20	C8	144	ARG
21	C9	6	VAL
21	C9	15	ILE
21	C9	18	TYR
21	C9	22	LEU
21	C9	25	GLN
21	C9	28	LEU
21	C9	30	VAL
21	C9	33	TYR
21	C9	34	VAL
21	C9	35	ASP
21	C9	36	ILE

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Mol	Chain	Res	Type
21	C9	39	THR
21	C9	43	ASN
21	C9	57	ARG
21	C9	60	SER
21	C9	63	ARG
21	C9	67	MET
21	C9	70	GLN
21	C9	71	VAL
21	C9	73	VAL
21	C9	76	LEU
21	C9	89	ARG
21	C9	115	GLU
21	C9	122	ARG
21	C9	124	ILE
21	C9	125	SER
21	C9	130	ARG
21	C9	139	THR
21	C9	143	ASP
22	D0	15	GLN
22	D0	21	LYS
22	D0	22	ILE
22	D0	23	ARG
22	D0	27	THR
22	D0	31	VAL
22	D0	46	GLU
22	D0	50	LEU
22	D0	51	VAL
22	D0	57	ARG
22	D0	60	THR
22	D0	64	LYS
22	D0	66	SER
22	D0	70	THR
22	D0	72	ASN
22	D0	74	GLU
22	D0	76	SER
22	D0	89	ARG
22	D0	97	VAL
22	D0	99	ILE
22	D0	103	ILE
22	D0	108	ILE
22	D0	109	GLU
22	D0	117	VAL

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Mol	Chain	Res	Type
23	D1	1	MET
23	D1	2	GLU
23	D1	5	LYS
23	D1	11	LEU
23	D1	25	LYS
23	D1	32	VAL
23	D1	49	GLU
23	D1	62	ARG
23	D1	69	LEU
23	D1	72	LEU
23	D1	78	LEU
23	D1	80	LYS
23	D1	87	ARG
24	D2	2	THR
24	D2	7	LEU
24	D2	12	ASN
24	D2	22	LYS
24	D2	23	ARG
24	D2	24	GLN
24	D2	26	LEU
24	D2	37	PHE
24	D2	53	ILE
24	D2	56	HIS
24	D2	65	LEU
24	D2	68	ARG
24	D2	76	SER
24	D2	86	ILE
24	D2	93	LEU
24	D2	97	ARG
24	D2	98	GLN
24	D2	103	ILE
24	D2	104	LEU
24	D2	121	VAL
25	D3	3	LYS
25	D3	7	ARG
25	D3	14	LYS
25	D3	16	ARG
25	D3	18	HIS
25	D3	19	ARG
25	D3	26	GLU
25	D3	28	ASN
25	D3	40	SER

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Mol	Chain	Res	Type
25	D3	47	SER
25	D3	53	VAL
25	D3	66	SER
25	D3	69	ARG
25	D3	72	VAL
25	D3	75	GLN
25	D3	84	THR
25	D3	96	VAL
25	D3	107	PHE
25	D3	109	ARG
25	D3	110	LYS
25	D3	114	LYS
25	D3	127	VAL
25	D3	132	LEU
25	D3	140	LYS
26	D4	13	ILE
26	D4	21	LYS
26	D4	24	VAL
26	D4	28	LEU
26	D4	32	ARG
26	D4	34	ASN
26	D4	35	VAL
26	D4	37	LYS
26	D4	51	GLU
26	D4	74	LEU
26	D4	88	THR
26	D4	98	GLU
26	D4	99	LYS
26	D4	102	LYS
26	D4	105	ARG
26	D4	123	LYS
26	D4	124	ARG
26	D4	125	LEU
26	D4	128	LYS
27	D5	37	GLN
27	D5	42	LEU
27	D5	50	ILE
27	D5	59	TYR
27	D5	60	VAL
27	D5	62	VAL
27	D5	67	ASP
27	D5	69	LEU

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Mol	Chain	Res	Type
27	D5	71	ILE
27	D5	75	LEU
27	D5	92	ILE
27	D5	95	HIS
27	D5	97	LYS
27	D5	100	ILE
27	D5	102	THR
28	D6	5	ARG
28	D6	12	LYS
28	D6	19	LYS
28	D6	21	VAL
28	D6	30	ILE
28	D6	36	ILE
28	D6	38	ARG
28	D6	41	ILE
28	D6	44	ILE
28	D6	45	VAL
28	D6	46	GLU
28	D6	61	GLU
28	D6	64	LEU
28	D6	66	LYS
28	D6	67	THR
28	D6	69	ASN
28	D6	82	ARG
28	D6	85	ARG
28	D6	86	VAL
28	D6	88	SER
28	D6	90	GLU
28	D6	91	ASP
29	D7	2	VAL
29	D7	3	LEU
29	D7	5	GLN
29	D7	20	LYS
29	D7	26	GLN
29	D7	33	LEU
29	D7	34	ASP
29	D7	41	LEU
29	D7	42	ASN
29	D7	63	LEU
29	D7	65	THR
30	D8	8	THR
30	D8	12	VAL

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Mol	Chain	Res	Type
30	D8	13	ILE
30	D8	14	LYS
30	D8	15	VAL
30	D8	19	THR
30	D8	26	THR
30	D8	32	PHE
30	D8	40	ILE
30	D8	52	ASP
30	D8	58	GLU
31	D9	5	ASN
31	D9	7	TRP
31	D9	21	CYS
31	D9	22	ARG
31	D9	30	LEU
31	D9	40	ARG
31	D9	42	CYS
32	E0	20	LYS
32	E0	22	GLU
32	E0	28	LYS
32	E0	41	THR
32	E0	42	ARG
32	E0	47	VAL
32	E0	49	LEU
32	E0	54	ARG
32	E0	56	MET
33	E1	84	VAL
33	E1	86	THR
33	E1	89	LYS
33	E1	93	HIS
33	E1	94	LYS
33	E1	97	LYS
33	E1	102	VAL
33	E1	107	LYS
33	E1	113	LYS
33	E1	115	THR
33	E1	116	LYS
33	E1	120	GLU
33	E1	146	SER
33	E1	147	VAL
34	SR	10	ARG
34	SR	14	GLU
34	SR	17	ASN

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Mol	Chain	Res	Type
34	SR	37	SER
34	SR	48	THR
34	SR	50	ASP
34	SR	52	GLN
34	SR	66	HIS
34	SR	74	THR
34	SR	76	ASP
34	SR	89	LEU
34	SR	94	VAL
34	SR	96	THR
34	SR	102	ARG
34	SR	113	VAL
34	SR	115	ILE
34	SR	117	LYS
34	SR	144	LEU
34	SR	149	ASP
34	SR	154	VAL
34	SR	164	ASP
34	SR	165	ASP
34	SR	188	ILE
34	SR	196	ASN
34	SR	199	ILE
34	SR	207	ASP
34	SR	221	MET
34	SR	231	MET
34	SR	238	ASP
34	SR	241	PHE
34	SR	248	ASN
34	SR	268	GLN
34	SR	269	TYR
34	SR	295	SER
34	SR	300	THR
34	SR	316	MET
34	SR	317	THR
35	SM	23	LYS
35	SM	24	GLU
35	SM	34	LYS
35	SM	45	SER
35	SM	46	LYS
35	SM	48	ARG
35	SM	51	ARG
35	SM	53	ARG

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Mol	Chain	Res	Type
35	SM	61	ILE
35	SM	64	LYS
35	SM	68	ARG
35	SM	69	ARG
35	SM	84	LYS
35	SM	89	ARG
35	SM	91	THR
35	SM	94	HIS
35	SM	100	THR
35	SM	101	ASP
35	SM	102	THR
35	SM	104	LYS
35	SM	105	LYS
35	SM	117	LEU
35	SM	121	LYS
35	SM	133	GLU
35	SM	139	GLU
39	L2	6	ARG
39	L2	19	HIS
39	L2	20	THR
39	L2	31	THR
39	L2	32	LEU
39	L2	33	ASP
39	L2	36	GLU
39	L2	41	ILE
39	L2	44	ILE
39	L2	45	VAL
39	L2	46	LYS
39	L2	48	ILE
39	L2	49	VAL
39	L2	52	SER
39	L2	61	VAL
39	L2	62	VAL
39	L2	70	ARG
39	L2	71	LEU
39	L2	73	GLU
39	L2	74	GLU
39	L2	84	THR
39	L2	97	ASN
39	L2	104	LEU
39	L2	107	VAL
39	L2	109	GLU

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Mol	Chain	Res	Type
39	L2	114	SER
39	L2	116	VAL
39	L2	118	GLU
39	L2	143	GLU
39	L2	145	LYS
39	L2	157	VAL
39	L2	165	VAL
39	L2	179	LEU
39	L2	180	LEU
39	L2	190	ARG
39	L2	192	LYS
39	L2	199	THR
39	L2	204	MET
39	L2	205	ASN
39	L2	206	PRO
39	L2	207	VAL
39	L2	218	HIS
39	L2	223	SER
39	L2	225	ILE
39	L2	226	SER
39	L2	227	ARG
39	L2	230	VAL
39	L2	238	ILE
39	L2	242	ARG
39	L2	247	ARG
40	L3	4	ARG
40	L3	7	GLU
40	L3	17	LEU
40	L3	19	ARG
40	L3	25	ILE
40	L3	30	LYS
40	L3	37	ARG
40	L3	39	LYS
40	L3	43	LEU
40	L3	45	SER
40	L3	55	THR
40	L3	61	ASP
40	L3	66	LYS
40	L3	73	VAL
40	L3	79	VAL
40	L3	85	VAL
40	L3	87	VAL

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Mol	Chain	Res	Type
40	L3	100	ARG
40	L3	103	THR
40	L3	112	ASP
40	L3	114	VAL
40	L3	116	ARG
40	L3	120	LYS
40	L3	124	LYS
40	L3	139	GLN
40	L3	146	ARG
40	L3	148	LEU
40	L3	150	ARG
40	L3	157	VAL
40	L3	159	ARG
40	L3	160	VAL
40	L3	168	LYS
40	L3	173	GLN
40	L3	178	LEU
40	L3	188	ILE
40	L3	196	ARG
40	L3	202	THR
40	L3	205	VAL
40	L3	208	VAL
40	L3	210	GLU
40	L3	226	PHE
40	L3	229	VAL
40	L3	238	LEU
40	L3	246	LEU
40	L3	252	ILE
40	L3	264	VAL
40	L3	284	ARG
40	L3	291	GLU
40	L3	293	ASN
40	L3	305	ILE
40	L3	306	THR
40	L3	317	ILE
40	L3	319	ASN
40	L3	320	ASP
40	L3	324	VAL
40	L3	328	ILE
40	L3	332	ARG
40	L3	338	LEU
40	L3	343	TYR

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Mol	Chain	Res	Type
40	L3	344	THR
40	L3	347	SER
40	L3	352	GLU
40	L3	354	VAL
40	L3	356	LEU
41	L4	2	SER
41	L4	25	VAL
41	L4	37	THR
41	L4	41	SER
41	L4	47	ARG
41	L4	60	THR
41	L4	69	ARG
41	L4	74	ILE
41	L4	92	ASN
41	L4	93	MET
41	L4	99	MET
41	L4	105	THR
41	L4	108	LYS
41	L4	112	LYS
41	L4	114	ASN
41	L4	120	TYR
41	L4	133	SER
41	L4	138	ARG
41	L4	142	VAL
41	L4	148	ILE
41	L4	150	LEU
41	L4	152	VAL
41	L4	156	LEU
41	L4	158	SER
41	L4	161	LYS
41	L4	169	LEU
41	L4	170	LYS
41	L4	176	SER
41	L4	179	LEU
41	L4	180	LYS
41	L4	185	LYS
41	L4	187	LEU
41	L4	193	LYS
41	L4	194	TYR
41	L4	196	ASN
41	L4	200	THR
41	L4	203	ARG

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Mol	Chain	Res	Type
41	L4	206	LEU
41	L4	220	ARG
41	L4	222	VAL
41	L4	230	VAL
41	L4	246	ARG
41	L4	250	TRP
41	L4	256	THR
41	L4	258	LEU
41	L4	259	ASP
41	L4	265	GLU
41	L4	267	VAL
41	L4	272	VAL
41	L4	278	SER
41	L4	282	SER
41	L4	287	THR
41	L4	288	ARG
41	L4	295	ILE
41	L4	299	ILE
41	L4	300	ARG
41	L4	306	THR
41	L4	308	LYS
41	L4	313	LEU
41	L4	323	VAL
41	L4	338	LYS
41	L4	347	THR
41	L4	350	LYS
41	L4	354	VAL
41	L4	356	THR
42	L5	4	GLN
42	L5	5	LYS
42	L5	9	SER
42	L5	22	ARG
42	L5	23	ARG
42	L5	35	ARG
42	L5	41	LYS
42	L5	58	LYS
42	L5	66	SER
42	L5	69	ILE
42	L5	85	ARG
42	L5	88	ILE
42	L5	89	THR
42	L5	105	ILE

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Mol	Chain	Res	Type
42	L5	109	THR
42	L5	112	LYS
42	L5	115	LEU
42	L5	122	VAL
42	L5	125	VAL
42	L5	131	LEU
42	L5	132	THR
42	L5	140	ARG
42	L5	144	VAL
42	L5	146	LEU
42	L5	148	ILE
42	L5	151	GLN
42	L5	152	ARG
42	L5	154	THR
42	L5	155	THR
42	L5	163	LEU
42	L5	177	GLU
42	L5	181	PRO
42	L5	185	PHE
42	L5	187	THR
42	L5	188	GLU
42	L5	190	ILE
42	L5	194	LEU
42	L5	196	ARG
42	L5	197	SER
42	L5	203	HIS
42	L5	211	LEU
42	L5	217	GLU
42	L5	218	ARG
42	L5	222	LEU
42	L5	229	ASP
42	L5	235	SER
42	L5	257	GLU
42	L5	259	LYS
42	L5	263	GLU
42	L5	268	GLU
42	L5	277	LEU
42	L5	290	ILE
42	L5	293	LEU
43	L6	5	LYS
43	L6	9	TRP
43	L6	15	VAL

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Mol	Chain	Res	Type
43	L6	21	THR
43	L6	28	GLN
43	L6	30	LEU
43	L6	35	VAL
43	L6	46	ARG
43	L6	52	VAL
43	L6	64	LEU
43	L6	76	LEU
43	L6	77	ARG
43	L6	78	ARG
43	L6	85	ILE
43	L6	89	THR
43	L6	90	LYS
43	L6	100	LYS
43	L6	109	GLU
43	L6	129	GLU
43	L6	134	ARG
43	L6	140	VAL
43	L6	145	LEU
43	L6	152	THR
43	L6	154	LEU
43	L6	155	LEU
43	L6	156	LYS
44	L7	24	GLU
44	L7	25	GLN
44	L7	26	VAL
44	L7	30	ARG
44	L7	37	ASN
44	L7	46	GLU
44	L7	48	ASN
44	L7	53	LYS
44	L7	59	GLU
44	L7	60	ARG
44	L7	82	LYS
44	L7	83	LEU
44	L7	92	ILE
44	L7	93	ASN
44	L7	109	THR
44	L7	110	ARG
44	L7	111	ILE
44	L7	115	THR
44	L7	124	LEU

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Mol	Chain	Res	Type
44	L7	129	LEU
44	L7	151	ARG
44	L7	153	PHE
44	L7	156	ILE
44	L7	158	LYS
44	L7	173	LEU
44	L7	179	LEU
44	L7	180	SER
44	L7	181	ILE
44	L7	184	LEU
44	L7	190	THR
44	L7	238	LYS
44	L7	239	LEU
45	L8	26	LEU
45	L8	27	THR
45	L8	31	PRO
45	L8	38	GLN
45	L8	41	GLN
45	L8	47	SER
45	L8	50	VAL
45	L8	58	VAL
45	L8	63	LYS
45	L8	67	ILE
45	L8	69	LEU
45	L8	71	VAL
45	L8	74	THR
45	L8	79	GLN
45	L8	82	LEU
45	L8	84	ARG
45	L8	106	LYS
45	L8	108	ARG
45	L8	118	GLU
45	L8	126	SER
45	L8	132	VAL
45	L8	136	LEU
45	L8	150	LEU
45	L8	156	ASP
45	L8	160	ILE
45	L8	163	VAL
45	L8	169	LEU
45	L8	172	LYS
45	L8	181	LYS

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Mol	Chain	Res	Type
45	L8	185	ARG
45	L8	194	THR
45	L8	201	THR
45	L8	206	GLU
45	L8	211	LEU
45	L8	216	SER
45	L8	232	HIS
45	L8	240	ASN
46	L9	1	MET
46	L9	4	ILE
46	L9	5	GLN
46	L9	9	GLN
46	L9	14	GLU
46	L9	16	VAL
46	L9	17	THR
46	L9	18	VAL
46	L9	19	SER
46	L9	20	ILE
46	L9	33	THR
46	L9	41	ILE
46	L9	44	THR
46	L9	46	THR
46	L9	47	LYS
46	L9	48	VAL
46	L9	49	ASN
46	L9	52	LEU
46	L9	62	ARG
46	L9	68	LEU
46	L9	69	ARG
46	L9	70	THR
46	L9	73	SER
46	L9	79	ILE
46	L9	82	VAL
46	L9	83	THR
46	L9	84	LYS
46	L9	120	ASP
46	L9	126	VAL
46	L9	132	VAL
46	L9	138	THR
46	L9	139	ASN
46	L9	146	LEU
46	L9	151	VAL

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Mol	Chain	Res	Type
46	L9	152	GLU
46	L9	157	ASN
46	L9	161	LEU
46	L9	162	GLN
46	L9	164	ILE
46	L9	168	ARG
46	L9	172	ILE
46	L9	174	LYS
46	L9	188	THR
46	L9	189	GLU
47	M0	3	ARG
47	M0	7	ARG
47	M0	15	LYS
47	M0	21	ARG
47	M0	26	VAL
47	M0	30	LYS
47	M0	32	ARG
47	M0	34	TYR
47	M0	39	LYS
47	M0	48	LEU
47	M0	52	LEU
47	M0	57	LEU
47	M0	66	GLU
47	M0	74	LYS
47	M0	82	ARG
47	M0	87	LEU
47	M0	90	ARG
47	M0	99	ILE
47	M0	102	MET
47	M0	116	ARG
47	M0	125	LEU
47	M0	128	ARG
47	M0	130	ASP
47	M0	138	VAL
47	M0	139	ARG
47	M0	144	ASN
47	M0	145	LYS
47	M0	146	ASP
47	M0	156	ARG
47	M0	163	GLN
47	M0	165	ILE
47	M0	166	ILE

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Mol	Chain	Res	Type
47	M0	169	LYS
47	M0	176	LEU
47	M0	177	ASP
47	M0	184	LYS
47	M0	185	ARG
47	M0	189	GLU
47	M0	203	LYS
47	M0	208	ASN
48	M1	9	MET
48	M1	10	ARG
48	M1	11	ASP
48	M1	12	LEU
48	M1	13	LYS
48	M1	16	LYS
48	M1	19	LEU
48	M1	23	VAL
48	M1	26	SER
48	M1	30	LEU
48	M1	34	SER
48	M1	44	THR
48	M1	46	VAL
48	M1	56	THR
48	M1	65	ILE
48	M1	71	VAL
48	M1	80	LEU
48	M1	82	ARG
48	M1	92	ARG
48	M1	94	ARG
48	M1	106	ILE
48	M1	107	ASP
48	M1	112	LEU
48	M1	115	LYS
48	M1	132	ASN
48	M1	133	ARG
48	M1	137	ARG
48	M1	138	VAL
48	M1	140	ARG
48	M1	143	ARG
48	M1	151	SER
48	M1	154	THR
48	M1	161	SER
48	M1	166	LYS

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Mol	Chain	Res	Type
48	M1	171	VAL
49	M3	10	LEU
49	M3	11	LYS
49	M3	17	HIS
49	M3	23	LYS
49	M3	24	VAL
49	M3	34	SER
49	M3	53	LEU
49	M3	54	LEU
49	M3	57	VAL
49	M3	58	VAL
49	M3	59	ARG
49	M3	62	THR
49	M3	67	ARG
49	M3	70	ARG
49	M3	79	GLU
49	M3	85	LEU
49	M3	97	VAL
49	M3	107	GLU
49	M3	108	ILE
49	M3	113	VAL
49	M3	121	SER
49	M3	124	ILE
49	M3	128	ARG
49	M3	131	LYS
49	M3	136	GLU
49	M3	144	THR
49	M3	164	GLU
49	M3	169	THR
49	M3	171	ARG
49	M3	175	SER
49	M3	190	LYS
49	M3	192	GLU
50	M4	3	THR
50	M4	5	SER
50	M4	8	LYS
50	M4	15	VAL
50	M4	25	LYS
50	M4	27	GLN
50	M4	38	ILE
50	M4	39	ILE
50	M4	44	VAL

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Mol	Chain	Res	Type
50	M4	50	LYS
50	M4	53	VAL
50	M4	58	ILE
50	M4	63	VAL
50	M4	65	LEU
50	M4	69	THR
50	M4	72	LEU
50	M4	82	SER
50	M4	90	VAL
50	M4	91	CYS
50	M4	102	LYS
50	M4	105	GLN
50	M4	107	GLU
51	M5	10	LEU
51	M5	13	LYS
51	M5	18	VAL
51	M5	19	LEU
51	M5	20	ARG
51	M5	22	LEU
51	M5	24	ARG
51	M5	33	LYS
51	M5	38	ARG
51	M5	41	ARG
51	M5	43	THR
51	M5	44	ARG
51	M5	49	ARG
51	M5	56	LYS
51	M5	62	TYR
51	M5	64	VAL
51	M5	68	ARG
51	M5	71	ARG
51	M5	75	VAL
51	M5	80	THR
51	M5	83	LYS
51	M5	86	ASN
51	M5	90	ASN
51	M5	93	LYS
51	M5	98	LEU
51	M5	104	GLU
51	M5	106	VAL
51	M5	109	ARG
51	M5	117	ASN

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Mol	Chain	Res	Type
51	M5	133	ILE
51	M5	144	ARG
51	M5	151	ILE
51	M5	153	ASP
51	M5	157	LYS
51	M5	159	ARG
51	M5	170	LYS
51	M5	182	ASN
51	M5	188	ARG
51	M5	190	THR
51	M5	195	ASN
51	M5	198	SER
51	M5	201	ARG
51	M5	204	LYS
52	M6	8	VAL
52	M6	22	VAL
52	M6	33	ILE
52	M6	34	VAL
52	M6	35	VAL
52	M6	41	LEU
52	M6	44	SER
52	M6	56	ASP
52	M6	57	PHE
52	M6	68	ARG
52	M6	74	ARG
52	M6	78	ARG
52	M6	79	ILE
52	M6	84	LEU
52	M6	85	ARG
52	M6	91	LYS
52	M6	101	ARG
52	M6	106	GLU
52	M6	117	ARG
52	M6	120	VAL
52	M6	122	GLN
52	M6	124	LEU
52	M6	126	VAL
52	M6	128	ARG
52	M6	134	LYS
52	M6	143	THR
52	M6	150	GLU
52	M6	151	ASP

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Mol	Chain	Res	Type
52	M6	152	VAL
52	M6	155	LYS
52	M6	160	ARG
52	M6	164	SER
52	M6	170	LYS
52	M6	175	THR
52	M6	182	ASN
52	M6	184	THR
52	M6	187	GLU
52	M6	188	SER
52	M6	190	VAL
53	M7	7	THR
53	M7	8	SER
53	M7	9	THR
53	M7	14	SER
53	M7	23	ARG
53	M7	30	ARG
53	M7	32	THR
53	M7	36	ILE
53	M7	49	GLU
53	M7	52	LEU
53	M7	53	ASP
53	M7	56	ARG
53	M7	69	ARG
53	M7	76	PHE
53	M7	78	VAL
53	M7	79	THR
53	M7	87	SER
53	M7	91	VAL
53	M7	94	LEU
53	M7	105	LYS
53	M7	115	SER
53	M7	126	ARG
53	M7	127	ARG
53	M7	128	ARG
53	M7	141	SER
53	M7	144	SER
53	M7	154	GLU
53	M7	168	LEU
53	M7	181	ARG
54	M8	3	ILE
54	M8	7	SER

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Mol	Chain	Res	Type
54	M8	15	HIS
54	M8	17	THR
54	M8	22	ASP
54	M8	26	LEU
54	M8	30	VAL
54	M8	32	LEU
54	M8	34	THR
54	M8	36	LEU
54	M8	39	ARG
54	M8	40	THR
54	M8	41	ASP
54	M8	50	LYS
54	M8	63	SER
54	M8	66	ARG
54	M8	67	ILE
54	M8	69	ARG
54	M8	74	GLU
54	M8	81	VAL
54	M8	86	THR
54	M8	95	GLU
54	M8	98	LYS
54	M8	111	ARG
54	M8	113	LYS
54	M8	129	VAL
54	M8	135	GLN
54	M8	138	LEU
54	M8	141	ARG
54	M8	146	SER
54	M8	171	LYS
54	M8	178	ARG
54	M8	179	ARG
54	M8	181	SER
55	M9	5	ARG
55	M9	7	GLN
55	M9	9	ARG
55	M9	41	ILE
55	M9	44	LEU
55	M9	51	VAL
55	M9	52	LYS
55	M9	55	VAL
55	M9	57	VAL
55	M9	60	LYS

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Mol	Chain	Res	Type
55	M9	71	ARG
55	M9	74	ARG
55	M9	81	ARG
55	M9	85	ARG
55	M9	91	SER
55	M9	103	ARG
55	M9	104	ARG
55	M9	106	LEU
55	M9	116	ASP
55	M9	128	LYS
55	M9	133	LYS
55	M9	134	HIS
55	M9	153	LYS
55	M9	155	LEU
55	M9	160	GLU
55	M9	164	LEU
55	M9	171	ASP
55	M9	176	ARG
55	M9	180	LYS
56	N0	1	MET
56	N0	3	HIS
56	N0	8	GLN
56	N0	16	THR
56	N0	23	LYS
56	N0	39	SER
56	N0	40	ARG
56	N0	45	LEU
56	N0	46	GLN
56	N0	51	VAL
56	N0	57	GLU
56	N0	61	ILE
56	N0	62	ASN
56	N0	71	LYS
56	N0	81	TYR
56	N0	87	THR
56	N0	104	GLU
56	N0	105	THR
56	N0	113	ARG
56	N0	115	ARG
56	N0	117	ARG
56	N0	122	HIS
56	N0	137	ARG

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Mol	Chain	Res	Type
56	N0	138	GLN
56	N0	139	TYR
56	N0	145	THR
56	N0	155	ARG
56	N0	157	GLN
56	N0	158	LYS
56	N0	160	THR
56	N0	162	THR
56	N0	167	ARG
56	N0	169	SER
56	N0	171	PHE
57	N1	9	SER
57	N1	12	ARG
57	N1	18	ASP
57	N1	27	LEU
57	N1	29	THR
57	N1	36	VAL
57	N1	38	ASP
57	N1	48	ILE
57	N1	52	MET
57	N1	60	LYS
57	N1	64	VAL
57	N1	75	ILE
57	N1	78	LYS
57	N1	79	MET
57	N1	83	ARG
57	N1	87	LYS
57	N1	88	ARG
57	N1	92	ARG
57	N1	97	LYS
57	N1	103	GLN
57	N1	104	GLU
57	N1	106	LEU
57	N1	122	GLN
57	N1	126	VAL
57	N1	128	LEU
57	N1	136	ARG
57	N1	139	ARG
57	N1	140	ILE
57	N1	141	VAL
57	N1	143	THR
57	N1	144	GLU

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Mol	Chain	Res	Type
57	N1	149	GLN
57	N1	154	VAL
57	N1	158	THR
58	N2	10	LYS
58	N2	14	THR
58	N2	21	SER
58	N2	37	LEU
58	N2	38	ILE
58	N2	43	VAL
58	N2	52	ASN
58	N2	63	VAL
58	N2	81	LYS
58	N2	88	GLN
58	N2	93	ILE
58	N2	95	PHE
58	N2	100	THR
58	N2	105	LEU
59	N3	13	ILE
59	N3	32	ARG
59	N3	33	ASN
59	N3	45	ARG
59	N3	58	VAL
59	N3	63	LYS
59	N3	69	LEU
59	N3	72	LYS
59	N3	74	MET
59	N3	78	VAL
59	N3	83	LYS
59	N3	91	VAL
59	N3	98	ASN
59	N3	102	ILE
59	N3	104	ASN
59	N3	106	LYS
59	N3	120	LYS
59	N3	125	LEU
59	N3	132	ASN
59	N3	136	VAL
60	N4	5	ILE
60	N4	7	SER
60	N4	19	THR
60	N4	27	LYS
60	N4	31	PHE

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Mol	Chain	Res	Type
60	N4	45	ASN
60	N4	52	THR
60	N4	53	VAL
60	N4	54	LEU
60	N4	58	HIS
61	N5	27	ARG
61	N5	33	ARG
61	N5	36	LYS
61	N5	38	LEU
61	N5	39	LYS
61	N5	42	ARG
61	N5	45	LYS
61	N5	48	SER
61	N5	49	LYS
61	N5	59	SER
61	N5	63	ILE
61	N5	71	THR
61	N5	73	MET
61	N5	75	LYS
61	N5	86	VAL
61	N5	92	LYS
61	N5	104	GLU
61	N5	115	ARG
61	N5	120	LYS
61	N5	125	ARG
61	N5	130	TYR
61	N5	134	ASP
61	N5	135	ILE
61	N5	138	ARG
61	N5	139	ILE
61	N5	142	ILE
62	N6	3	LYS
62	N6	6	LEU
62	N6	10	SER
62	N6	11	ASP
62	N6	13	ARG
62	N6	17	LYS
62	N6	37	LYS
62	N6	45	ILE
62	N6	48	LEU
62	N6	57	LEU
62	N6	58	VAL

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Mol	Chain	Res	Type
62	N6	60	ARG
62	N6	71	SER
62	N6	74	TYR
62	N6	80	VAL
62	N6	88	GLU
62	N6	101	PRO
62	N6	102	SER
62	N6	113	LYS
62	N6	115	ARG
62	N6	127	GLU
63	N7	9	LYS
63	N7	14	VAL
63	N7	21	LYS
63	N7	24	VAL
63	N7	33	SER
63	N7	34	LYS
63	N7	46	ILE
63	N7	53	VAL
63	N7	67	LYS
63	N7	72	ILE
63	N7	81	LEU
63	N7	83	THR
63	N7	86	THR
63	N7	87	LEU
63	N7	95	VAL
63	N7	99	GLU
63	N7	102	GLU
63	N7	103	GLN
63	N7	119	GLU
63	N7	127	ASN
63	N7	135	ARG
64	N8	4	ARG
64	N8	8	THR
64	N8	12	ARG
64	N8	24	LYS
64	N8	29	PRO
64	N8	34	MET
64	N8	43	ILE
64	N8	46	ASP
64	N8	47	LYS
64	N8	65	GLN
64	N8	78	LEU

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Mol	Chain	Res	Type
64	N8	91	LEU
64	N8	115	LYS
64	N8	118	ILE
64	N8	120	ASN
64	N8	123	VAL
64	N8	124	ILE
64	N8	130	VAL
64	N8	133	LEU
64	N8	137	LYS
65	N9	5	LYS
65	N9	8	THR
65	N9	13	THR
65	N9	14	ARG
65	N9	18	ARG
65	N9	25	LYS
65	N9	28	LYS
65	N9	36	ASP
65	N9	38	LYS
65	N9	42	ASN
65	N9	59	LYS
66	O0	14	LEU
66	O0	16	LEU
66	O0	18	ILE
66	O0	33	SER
66	O0	34	LEU
66	O0	36	GLN
66	O0	40	LYS
66	O0	42	ILE
66	O0	52	ARG
66	O0	61	MET
66	O0	75	ASN
66	O0	86	ARG
66	O0	99	ASP
66	O0	100	ILE
66	O0	102	THR
67	O1	6	ASP
67	O1	13	THR
67	O1	16	LEU
67	O1	26	LYS
67	O1	28	ARG
67	O1	31	ARG
67	O1	34	LYS

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Mol	Chain	Res	Type
67	O1	42	LEU
67	O1	46	THR
67	O1	64	VAL
67	O1	65	LYS
67	O1	68	GLU
67	O1	71	LEU
67	O1	73	LEU
67	O1	79	ARG
67	O1	82	GLU
67	O1	83	GLU
67	O1	84	ASP
67	O1	86	LYS
67	O1	96	VAL
67	O1	98	VAL
67	O1	100	SER
67	O1	106	THR
68	O2	10	VAL
68	O2	19	ARG
68	O2	27	ARG
68	O2	34	LYS
68	O2	41	VAL
68	O2	51	SER
68	O2	54	LYS
68	O2	59	SER
68	O2	61	LYS
68	O2	73	THR
68	O2	75	LEU
68	O2	79	VAL
68	O2	82	LEU
68	O2	84	THR
68	O2	87	MET
68	O2	99	ASN
68	O2	101	SER
68	O2	103	LYS
68	O2	105	ARG
68	O2	109	LEU
68	O2	123	LYS
68	O2	125	ARG
69	O3	4	SER
69	O3	15	SER
69	O3	21	ARG
69	O3	29	LEU

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Mol	Chain	Res	Type
69	O3	31	LYS
69	O3	37	THR
69	O3	45	LEU
69	O3	47	LYS
69	O3	48	ARG
69	O3	49	ILE
69	O3	59	VAL
69	O3	70	LYS
69	O3	77	ASN
69	O3	80	VAL
69	O3	81	VAL
69	O3	88	ASN
69	O3	93	THR
69	O3	98	VAL
69	O3	106	ASN
70	O4	5	VAL
70	O4	7	PHE
70	O4	8	ARG
70	O4	16	ARG
70	O4	20	ILE
70	O4	21	LYS
70	O4	29	ILE
70	O4	36	LYS
70	O4	38	LEU
70	O4	51	LEU
70	O4	52	GLN
70	O4	58	ARG
70	O4	65	VAL
70	O4	71	THR
70	O4	74	ARG
70	O4	81	CYS
70	O4	86	LYS
70	O4	88	ARG
70	O4	103	LYS
70	O4	104	VAL
71	O5	13	SER
71	O5	21	LEU
71	O5	27	GLU
71	O5	28	LEU
71	O5	31	LEU
71	O5	44	ILE
71	O5	45	LYS

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Mol	Chain	Res	Type
71	O5	46	THR
71	O5	49	LYS
71	O5	62	GLN
71	O5	69	LEU
71	O5	71	LYS
71	O5	73	LYS
71	O5	89	ARG
71	O5	90	ARG
71	O5	96	GLU
71	O5	101	THR
71	O5	102	GLU
71	O5	103	LYS
71	O5	115	LYS
71	O5	119	LYS
72	O6	9	ILE
72	O6	20	MET
72	O6	21	THR
72	O6	25	LYS
72	O6	26	ILE
72	O6	28	TYR
72	O6	30	LYS
72	O6	44	VAL
72	O6	45	ARG
72	O6	46	GLU
72	O6	57	LEU
72	O6	58	ILE
72	O6	60	LEU
72	O6	64	SER
72	O6	67	LYS
72	O6	68	ARG
72	O6	70	ARG
72	O6	74	LYS
72	O6	76	ARG
72	O6	81	THR
72	O6	82	ARG
72	O6	84	LYS
72	O6	88	GLU
72	O6	90	MET
72	O6	98	ARG
73	O7	15	SER
73	O7	18	LEU
73	O7	19	CYS

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Mol	Chain	Res	Type
73	O7	21	ARG
73	O7	24	ARG
73	O7	25	ARG
73	O7	34	CYS
73	O7	36	SER
73	O7	44	THR
73	O7	55	ARG
73	O7	58	THR
73	O7	59	THR
73	O7	67	LEU
73	O7	76	ASN
73	O7	87	SER
74	O8	5	ILE
74	O8	6	THR
74	O8	28	ASN
74	O8	31	LEU
74	O8	32	ASN
74	O8	39	ARG
74	O8	46	ARG
74	O8	48	SER
74	O8	50	SER
74	O8	51	LEU
74	O8	53	THR
74	O8	64	LYS
74	O8	65	LEU
74	O8	67	GLN
74	O8	69	LEU
74	O8	72	THR
74	O8	73	LEU
74	O8	77	ARG
75	O9	5	LYS
75	O9	21	ARG
75	O9	23	LEU
75	O9	25	GLN
75	O9	51	ILE
76	Q0	78	ILE
76	Q0	79	GLU
76	Q0	92	ASP
76	Q0	112	LYS
76	Q0	113	ARG
76	Q0	127	LEU
77	Q1	2	ARG

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Mol	Chain	Res	Type
77	Q1	5	TRP
77	Q1	10	THR
77	Q1	11	ARG
77	Q1	13	LEU
77	Q1	15	ARG
77	Q1	16	LYS
78	Q2	6	LYS
78	Q2	9	LYS
78	Q2	13	LYS
78	Q2	20	HIS
78	Q2	28	TYR
78	Q2	29	LYS
78	Q2	32	LYS
78	Q2	35	LEU
78	Q2	38	GLN
78	Q2	45	ARG
78	Q2	61	LYS
78	Q2	66	LYS
78	Q2	71	ARG
78	Q2	78	LYS
78	Q2	80	ARG
78	Q2	83	LEU
78	Q2	84	THR
78	Q2	85	LEU
78	Q2	86	LYS
78	Q2	88	CYS
78	Q2	93	LEU
78	Q2	100	LYS
78	Q2	104	LEU
79	Q3	11	THR
79	Q3	28	LYS
79	Q3	32	GLN
79	Q3	33	GLN
79	Q3	38	ASP
79	Q3	40	SER
79	Q3	45	LYS
79	Q3	57	CYS
79	Q3	59	CYS
79	Q3	60	CYS
79	Q3	64	VAL
79	Q3	71	VAL
79	Q3	78	THR

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Mol	Chain	Res	Type
79	Q3	84	ARG
79	Q3	90	VAL
79	Q3	91	GLU
2	s0	9	LEU
2	s0	10	THR
2	s0	27	ARG
2	s0	30	GLN
2	s0	32	HIS
2	s0	41	ARG
2	s0	45	VAL
2	s0	50	VAL
2	s0	59	LEU
2	s0	72	ASP
2	s0	80	THR
2	s0	83	GLN
2	s0	87	LEU
2	s0	88	LYS
2	s0	96	THR
2	s0	101	ARG
2	s0	114	SER
2	s0	141	ILE
2	s0	154	GLU
2	s0	162	CYS
2	s0	164	ASN
2	s0	165	ARG
2	s0	168	HIS
2	s0	172	LEU
2	s0	177	LEU
2	s0	183	ARG
2	s0	189	VAL
2	s0	200	ASP
2	s0	202	TYR
2	s0	203	PHE
3	s1	21	VAL
3	s1	25	THR
3	s1	47	LEU
3	s1	48	VAL
3	s1	51	SER
3	s1	54	LEU
3	s1	58	SER
3	s1	62	LYS
3	s1	64	ARG

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Mol	Chain	Res	Type
3	s1	68	VAL
3	s1	70	LEU
3	s1	72	ASP
3	s1	73	LEU
3	s1	78	ASP
3	s1	81	PHE
3	s1	82	ARG
3	s1	90	GLU
3	s1	97	LEU
3	s1	105	PHE
3	s1	111	ARG
3	s1	115	ARG
3	s1	125	VAL
3	s1	126	THR
3	s1	129	THR
3	s1	135	LEU
3	s1	151	LYS
3	s1	154	SER
3	s1	159	SER
3	s1	169	SER
3	s1	173	THR
3	s1	175	GLU
3	s1	177	GLN
3	s1	179	SER
3	s1	180	THR
3	s1	181	LEU
3	s1	183	GLN
3	s1	186	SER
3	s1	189	ILE
3	s1	203	ASP
3	s1	204	ILE
3	s1	205	PHE
3	s1	212	VAL
3	s1	217	LEU
3	s1	223	PHE
3	s1	225	VAL
4	s2	41	LEU
4	s2	44	LEU
4	s2	51	THR
4	s2	52	THR
4	s2	53	ILE
4	s2	54	GLU

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Mol	Chain	Res	Type
4	s2	55	GLU
4	s2	58	LEU
4	s2	60	SER
4	s2	65	GLU
4	s2	69	ILE
4	s2	70	ASP
4	s2	72	LEU
4	s2	76	LEU
4	s2	77	GLN
4	s2	83	ILE
4	s2	89	GLN
4	s2	90	THR
4	s2	91	ARG
4	s2	95	ARG
4	s2	97	ARG
4	s2	101	VAL
4	s2	111	VAL
4	s2	113	LEU
4	s2	116	LYS
4	s2	117	THR
4	s2	125	ILE
4	s2	134	LEU
4	s2	137	ILE
4	s2	141	ARG
4	s2	148	LEU
4	s2	158	THR
4	s2	165	VAL
4	s2	169	LEU
4	s2	174	ARG
4	s2	178	ILE
4	s2	181	SER
4	s2	194	GLU
4	s2	198	THR
4	s2	210	THR
4	s2	222	TYR
4	s2	226	THR
4	s2	228	ASN
4	s2	232	GLU
4	s2	237	VAL
4	s2	240	LEU
4	s2	246	GLU
4	s2	250	GLN

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Mol	Chain	Res	Type
5	s3	7	LYS
5	s3	11	LEU
5	s3	21	LEU
5	s3	26	THR
5	s3	39	VAL
5	s3	40	ARG
5	s3	42	THR
5	s3	44	THR
5	s3	57	ASP
5	s3	61	GLU
5	s3	70	THR
5	s3	72	LEU
5	s3	84	ILE
5	s3	86	LEU
5	s3	90	ARG
5	s3	93	ASP
5	s3	96	LEU
5	s3	115	ILE
5	s3	127	MET
5	s3	128	GLU
5	s3	134	CYS
5	s3	142	LEU
5	s3	143	ARG
5	s3	157	LEU
5	s3	158	ILE
5	s3	162	GLN
5	s3	164	VAL
5	s3	168	ILE
5	s3	169	ASP
5	s3	172	THR
5	s3	176	LEU
5	s3	185	LYS
5	s3	195	SER
5	s3	196	ARG
5	s3	202	LEU
5	s3	204	ASP
5	s3	206	VAL
5	s3	210	GLU
5	s3	212	LYS
5	s3	213	GLU
5	s3	217	ILE
5	s3	224	ASP

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Mol	Chain	Res	Type
6	s4	6	LYS
6	s4	9	LEU
6	s4	11	ARG
6	s4	22	LYS
6	s4	38	LEU
6	s4	39	ARG
6	s4	40	GLU
6	s4	42	LEU
6	s4	45	ILE
6	s4	47	PHE
6	s4	48	LEU
6	s4	49	ARG
6	s4	51	ARG
6	s4	67	GLN
6	s4	70	VAL
6	s4	76	VAL
6	s4	78	THR
6	s4	80	THR
6	s4	92	LEU
6	s4	97	GLU
6	s4	98	ASN
6	s4	115	THR
6	s4	116	ASP
6	s4	123	LEU
6	s4	126	VAL
6	s4	131	LEU
6	s4	143	ASP
6	s4	146	THR
6	s4	147	ILE
6	s4	148	ARG
6	s4	164	LEU
6	s4	176	ASP
6	s4	180	LEU
6	s4	182	TYR
6	s4	194	THR
6	s4	195	ILE
6	s4	208	VAL
6	s4	219	VAL
6	s4	221	ARG
6	s4	222	LEU
6	s4	226	PHE
6	s4	227	VAL

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Mol	Chain	Res	Type
6	s4	228	ILE
6	s4	235	TYR
6	s4	236	ILE
6	s4	246	LEU
6	s4	247	SER
6	s4	252	ARG
7	s5	24	VAL
7	s5	25	LEU
7	s5	31	GLU
7	s5	38	THR
7	s5	39	GLU
7	s5	41	LYS
7	s5	45	LYS
7	s5	59	VAL
7	s5	63	GLN
7	s5	66	GLN
7	s5	68	ILE
7	s5	72	HIS
7	s5	83	ARG
7	s5	89	ILE
7	s5	92	ARG
7	s5	93	LEU
7	s5	104	ASN
7	s5	108	LEU
7	s5	112	ARG
7	s5	125	THR
7	s5	130	ILE
7	s5	140	THR
7	s5	148	ARG
7	s5	149	VAL
7	s5	156	ARG
7	s5	157	ARG
7	s5	160	VAL
7	s5	162	VAL
7	s5	163	SER
7	s5	166	ARG
7	s5	167	ARG
7	s5	190	ILE
7	s5	194	LEU
7	s5	196	GLU
7	s5	216	GLU
7	s5	217	LEU

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Mol	Chain	Res	Type
7	s5	219	ARG
8	s6	25	ARG
8	s6	68	LEU
8	s6	69	LEU
8	s6	93	LYS
8	s6	96	SER
8	s6	109	LEU
8	s6	112	VAL
8	s6	120	GLU
8	s6	121	LEU
8	s6	122	GLU
8	s6	124	LEU
8	s6	125	THR
8	s6	126	ASP
8	s6	127	THR
8	s6	129	VAL
8	s6	133	LEU
8	s6	143	LYS
8	s6	151	ASP
8	s6	153	VAL
8	s6	157	VAL
8	s6	162	VAL
8	s6	163	THR
8	s6	167	LYS
8	s6	168	THR
8	s6	170	THR
8	s6	177	ARG
8	s6	179	VAL
8	s6	180	THR
8	s6	182	GLN
8	s6	208	TYR
8	s6	212	LEU
8	s6	215	ARG
8	s6	216	LEU
8	s6	218	GLU
9	s7	10	SER
9	s7	17	GLU
9	s7	22	GLN
9	s7	27	LEU
9	s7	28	GLU
9	s7	30	SER
9	s7	33	GLU

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Mol	Chain	Res	Type
9	s7	35	LYS
9	s7	39	ARG
9	s7	42	GLN
9	s7	50	ASP
9	s7	62	VAL
9	s7	67	LEU
9	s7	75	THR
9	s7	76	LYS
9	s7	78	THR
9	s7	88	ARG
9	s7	93	LEU
9	s7	97	ARG
9	s7	106	SER
9	s7	107	ARG
9	s7	108	GLN
9	s7	110	GLN
9	s7	114	ARG
9	s7	116	ARG
9	s7	118	LEU
9	s7	122	HIS
9	s7	123	ASP
9	s7	143	LEU
9	s7	144	VAL
9	s7	149	ILE
9	s7	160	GLN
9	s7	161	GLN
9	s7	166	LEU
9	s7	185	ILE
10	s8	6	ASP
10	s8	20	GLN
10	s8	22	ARG
10	s8	24	LYS
10	s8	25	ARG
10	s8	29	LEU
10	s8	32	GLN
10	s8	48	THR
10	s8	58	LEU
10	s8	59	ARG
10	s8	60	ILE
10	s8	76	THR
10	s8	95	THR
10	s8	97	THR

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Mol	Chain	Res	Type
10	s8	120	THR
10	s8	152	ILE
10	s8	155	SER
10	s8	172	ARG
10	s8	175	GLN
10	s8	182	TYR
10	s8	184	LEU
10	s8	185	GLU
10	s8	199	LYS
11	s9	3	ARG
11	s9	6	ARG
11	s9	7	THR
11	s9	16	LYS
11	s9	20	GLU
11	s9	21	SER
11	s9	28	LEU
11	s9	33	GLU
11	s9	36	LEU
11	s9	39	LYS
11	s9	46	SER
11	s9	81	VAL
11	s9	82	ARG
11	s9	93	LEU
11	s9	96	VAL
11	s9	110	GLN
11	s9	114	TYR
11	s9	122	VAL
11	s9	126	ARG
11	s9	130	THR
11	s9	132	ARG
11	s9	134	ILE
11	s9	150	LEU
11	s9	152	SER
11	s9	161	THR
11	s9	162	SER
11	s9	168	ARG
11	s9	171	ARG
11	s9	172	VAL
11	s9	182	GLU
12	c0	2	LEU
12	c0	6	GLU
12	c0	12	HIS

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Mol	Chain	Res	Type
12	c0	15	LEU
12	c0	27	PHE
12	c0	36	ASP
12	c0	40	LEU
12	c0	47	GLN
12	c0	55	VAL
12	c0	57	THR
12	c0	67	THR
12	c0	70	GLU
12	c0	77	ARG
13	c1	5	LEU
13	c1	9	SER
13	c1	10	GLU
13	c1	14	GLN
13	c1	26	LYS
13	c1	27	THR
13	c1	31	THR
13	c1	33	ARG
13	c1	44	THR
13	c1	49	ILE
13	c1	60	PHE
13	c1	67	ARG
13	c1	72	THR
13	c1	74	THR
13	c1	80	MET
13	c1	82	ARG
13	c1	83	THR
13	c1	99	ARG
13	c1	109	VAL
13	c1	123	VAL
13	c1	141	LYS
14	c2	28	LEU
14	c2	52	LEU
14	c2	53	THR
14	c2	58	LEU
14	c2	61	VAL
14	c2	62	LEU
14	c2	71	ILE
14	c2	74	LEU
14	c2	83	GLU
14	c2	85	LYS
14	c2	89	ILE

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Mol	Chain	Res	Type
14	c2	93	ASP
14	c2	95	LYS
14	c2	103	LEU
14	c2	126	TRP
14	c2	132	GLU
14	c2	140	PHE
15	c3	4	MET
15	c3	6	SER
15	c3	11	ILE
15	c3	16	ILE
15	c3	21	ASN
15	c3	35	GLU
15	c3	36	GLN
15	c3	39	LYS
15	c3	43	LYS
15	c3	64	ARG
15	c3	66	ILE
15	c3	70	LYS
15	c3	72	MET
15	c3	80	LEU
15	c3	83	GLU
15	c3	84	ILE
15	c3	87	ASP
15	c3	97	SER
15	c3	102	LEU
15	c3	104	ARG
15	c3	119	GLU
15	c3	125	LEU
15	c3	131	THR
15	c3	134	VAL
15	c3	149	LEU
16	c4	14	PHE
16	c4	26	THR
16	c4	28	VAL
16	c4	31	THR
16	c4	38	THR
16	c4	61	MET
16	c4	65	GLN
16	c4	77	THR
16	c4	79	VAL
16	c4	81	VAL
16	c4	103	ARG

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Mol	Chain	Res	Type
16	c4	107	ARG
16	c4	108	SER
16	c4	111	ARG
16	c4	114	ARG
16	c4	121	VAL
16	c4	123	SER
16	c4	125	SER
16	c4	129	LYS
16	c4	132	ARG
16	c4	136	ARG
17	c5	12	PHE
17	c5	21	ASP
17	c5	24	LYS
17	c5	26	LEU
17	c5	27	GLU
17	c5	29	SER
17	c5	34	VAL
17	c5	36	LEU
17	c5	43	ARG
17	c5	61	ARG
17	c5	69	GLU
17	c5	72	LYS
17	c5	94	VAL
17	c5	107	ILE
17	c5	122	THR
17	c5	123	TYR
17	c5	124	THR
17	c5	126	VAL
18	c6	15	SER
18	c6	17	THR
18	c6	23	LYS
18	c6	28	LEU
18	c6	31	VAL
18	c6	37	THR
18	c6	42	GLU
18	c6	43	ILE
18	c6	53	LEU
18	c6	54	LEU
18	c6	55	VAL
18	c6	57	LEU
18	c6	68	ARG
18	c6	69	VAL

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Mol	Chain	Res	Type
18	c6	83	GLN
18	c6	94	GLN
18	c6	98	ASP
18	c6	105	LEU
18	c6	110	THR
18	c6	114	ARG
18	c6	117	LEU
18	c6	137	ARG
19	c7	3	ARG
19	c7	6	THR
19	c7	8	THR
19	c7	25	THR
19	c7	26	LEU
19	c7	29	GLN
19	c7	34	LEU
19	c7	35	CYS
19	c7	38	ILE
19	c7	44	LYS
19	c7	46	LEU
19	c7	47	ARG
19	c7	49	LYS
19	c7	55	THR
19	c7	62	GLN
19	c7	69	ILE
19	c7	72	LYS
19	c7	85	VAL
19	c7	88	VAL
19	c7	110	VAL
20	c8	3	LEU
20	c8	4	VAL
20	c8	5	VAL
20	c8	6	GLN
20	c8	13	HIS
20	c8	15	LEU
20	c8	20	THR
20	c8	25	ASN
20	c8	26	ILE
20	c8	27	LYS
20	c8	28	ILE
20	c8	29	VAL
20	c8	33	THR
20	c8	34	THR

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Mol	Chain	Res	Type
20	c8	36	LYS
20	c8	40	ARG
20	c8	41	ARG
20	c8	54	LEU
20	c8	55	HIS
20	c8	63	GLN
20	c8	66	LEU
20	c8	85	PHE
20	c8	86	LEU
20	c8	94	ASP
20	c8	103	ASN
20	c8	110	ARG
20	c8	112	ASP
20	c8	113	LEU
20	c8	119	ILE
20	c8	138	THR
20	c8	140	THR
20	c8	141	THR
20	c8	144	ARG
20	c8	145	ARG
21	c9	6	VAL
21	c9	9	VAL
21	c9	20	SER
21	c9	27	LYS
21	c9	28	LEU
21	c9	38	LYS
21	c9	68	ARG
21	c9	70	GLN
21	c9	86	ARG
21	c9	89	ARG
21	c9	111	ILE
21	c9	116	ILE
21	c9	117	SER
21	c9	123	ARG
21	c9	139	THR
21	c9	140	LEU
21	c9	141	GLU
21	c9	142	GLU
21	c9	144	GLU
22	d0	15	GLN
22	d0	16	GLN
22	d0	20	ILE

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Mol	Chain	Res	Type
22	d0	23	ARG
22	d0	27	THR
22	d0	31	VAL
22	d0	34	LEU
22	d0	39	SER
22	d0	44	ASN
22	d0	49	ASN
22	d0	51	VAL
22	d0	57	ARG
22	d0	60	THR
22	d0	61	LYS
22	d0	62	VAL
22	d0	66	SER
22	d0	70	THR
22	d0	72	ASN
22	d0	74	GLU
22	d0	76	SER
22	d0	77	LYS
22	d0	81	THR
22	d0	88	LYS
22	d0	92	ASP
22	d0	97	VAL
22	d0	99	ILE
22	d0	102	ARG
22	d0	103	ILE
22	d0	105	GLN
22	d0	108	ILE
22	d0	114	VAL
23	d1	2	GLU
23	d1	5	LYS
23	d1	12	TYR
23	d1	13	VAL
23	d1	18	SER
23	d1	24	ILE
23	d1	31	SER
23	d1	32	VAL
23	d1	34	ILE
23	d1	39	VAL
23	d1	41	GLU
23	d1	50	TYR
23	d1	52	THR
23	d1	68	SER

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Mol	Chain	Res	Type
23	d1	78	LEU
23	d1	85	TYR
24	d2	2	THR
24	d2	6	VAL
24	d2	7	LEU
24	d2	16	ASN
24	d2	23	ARG
24	d2	24	GLN
24	d2	25	VAL
24	d2	26	LEU
24	d2	37	PHE
24	d2	49	GLU
24	d2	56	HIS
24	d2	65	LEU
24	d2	74	VAL
24	d2	93	LEU
24	d2	97	ARG
24	d2	98	GLN
24	d2	103	ILE
24	d2	117	ARG
24	d2	118	ARG
24	d2	126	LEU
24	d2	129	VAL
25	d3	7	ARG
25	d3	13	ARG
25	d3	18	HIS
25	d3	20	ARG
25	d3	22	ASN
25	d3	23	ARG
25	d3	28	ASN
25	d3	29	TYR
25	d3	36	THR
25	d3	41	SER
25	d3	46	SER
25	d3	52	ILE
25	d3	62	LYS
25	d3	72	VAL
25	d3	73	ARG
25	d3	78	LYS
25	d3	83	VAL
25	d3	84	THR
25	d3	99	ASN

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Mol	Chain	Res	Type
25	d3	103	LEU
25	d3	107	PHE
25	d3	109	ARG
25	d3	117	ILE
25	d3	125	VAL
25	d3	130	VAL
25	d3	132	LEU
25	d3	133	LEU
25	d3	138	GLU
26	d4	2	SER
26	d4	6	THR
26	d4	9	THR
26	d4	13	ILE
26	d4	21	LYS
26	d4	26	ASP
26	d4	29	HIS
26	d4	34	ASN
26	d4	43	LYS
26	d4	49	LYS
26	d4	52	LYS
26	d4	62	THR
26	d4	83	LYS
26	d4	88	THR
26	d4	92	VAL
26	d4	96	LEU
26	d4	118	ILE
26	d4	127	LYS
26	d4	133	ASN
27	d5	41	ILE
27	d5	43	ASP
27	d5	45	GLU
27	d5	49	ARG
27	d5	51	LEU
27	d5	57	TYR
27	d5	60	VAL
27	d5	68	ARG
27	d5	71	ILE
27	d5	81	ARG
27	d5	96	SER
27	d5	102	THR
27	d5	103	ARG
28	d6	5	ARG

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Mol	Chain	Res	Type
28	d6	8	ASN
28	d6	10	ARG
28	d6	12	LYS
28	d6	15	ARG
28	d6	18	VAL
28	d6	25	ASN
28	d6	27	SER
28	d6	38	ARG
28	d6	42	ARG
28	d6	43	ASN
28	d6	44	ILE
28	d6	46	GLU
28	d6	51	ARG
28	d6	55	GLU
28	d6	61	GLU
28	d6	82	ARG
28	d6	85	ARG
29	d7	3	LEU
29	d7	9	HIS
29	d7	17	ARG
29	d7	34	ASP
29	d7	38	PRO
29	d7	43	ILE
29	d7	45	THR
29	d7	46	VAL
29	d7	52	THR
29	d7	56	CYS
29	d7	65	THR
29	d7	72	LYS
29	d7	77	THR
29	d7	78	SER
29	d7	81	ARG
30	d8	13	ILE
30	d8	16	LEU
30	d8	19	THR
30	d8	22	ARG
30	d8	32	PHE
30	d8	33	LEU
30	d8	38	ARG
30	d8	48	VAL
30	d8	58	GLU
30	d8	62	GLU

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Mol	Chain	Res	Type
30	d8	64	ARG
30	d8	65	ARG
30	d8	66	LEU
31	d9	6	VAL
31	d9	10	HIS
31	d9	14	TYR
31	d9	32	ARG
31	d9	36	LEU
31	d9	38	ILE
31	d9	42	CYS
31	d9	44	ARG
31	d9	45	GLU
31	d9	49	ASP
31	d9	54	LYS
80	e0	13	LYS
80	e0	14	VAL
80	e0	16	SER
80	e0	24	THR
80	e0	25	GLU
80	e0	26	LYS
80	e0	29	LYS
80	e0	31	LYS
80	e0	39	LEU
80	e0	41	THR
80	e0	45	VAL
80	e0	46	ASN
80	e0	49	LEU
80	e0	51	ASN
80	e0	54	ARG
33	e1	80	ARG
33	e1	82	LYS
33	e1	86	THR
33	e1	90	LYS
33	e1	91	ILE
33	e1	96	LYS
33	e1	100	LEU
33	e1	102	VAL
33	e1	106	TYR
33	e1	109	ASP
33	e1	113	LYS
33	e1	118	ARG
33	e1	135	HIS

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Mol	Chain	Res	Type
33	e1	137	ASP
33	e1	147	VAL
33	e1	148	TYR
33	e1	150	VAL
34	sR	4	ASN
34	sR	6	VAL
34	sR	9	LEU
34	sR	17	ASN
34	sR	21	THR
34	sR	29	GLN
34	sR	46	LYS
34	sR	53	LYS
34	sR	58	VAL
34	sR	64	HIS
34	sR	66	HIS
34	sR	70	ASP
34	sR	72	THR
34	sR	76	ASP
34	sR	84	SER
34	sR	87	LYS
34	sR	96	THR
34	sR	100	TYR
34	sR	102	ARG
34	sR	108	SER
34	sR	130	THR
34	sR	145	LEU
34	sR	149	ASP
34	sR	159	ASN
34	sR	167	VAL
34	sR	176	LYS
34	sR	178	VAL
34	sR	191	ASP
34	sR	207	ASP
34	sR	210	LEU
34	sR	228	LYS
34	sR	232	TYR
34	sR	235	SER
34	sR	286	GLU
34	sR	297	ASP
34	sR	299	GLN
34	sR	314	GLN
34	sR	317	THR

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Mol	Chain	Res	Type
35	sM	41	SER
35	sM	43	ASP
35	sM	61	ILE
35	sM	69	ARG
35	sM	71	ASN
35	sM	74	LYS
35	sM	75	ASP
35	sM	76	VAL
35	sM	81	THR
39	l2	30	ARG
39	l2	31	THR
39	l2	32	LEU
39	l2	36	GLU
39	l2	42	ARG
39	l2	44	ILE
39	l2	45	VAL
39	l2	46	LYS
39	l2	48	ILE
39	l2	49	VAL
39	l2	61	VAL
39	l2	62	VAL
39	l2	70	ARG
39	l2	71	LEU
39	l2	75	ILE
39	l2	77	ILE
39	l2	79	ASN
39	l2	80	GLU
39	l2	82	VAL
39	l2	101	VAL
39	l2	119	LYS
39	l2	122	ASP
39	l2	128	ARG
39	l2	130	SER
39	l2	132	ASN
39	l2	137	ILE
39	l2	144	ASN
39	l2	147	ARG
39	l2	148	VAL
39	l2	149	ARG
39	l2	157	VAL
39	l2	158	ILE
39	l2	165	VAL

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Mol	Chain	Res	Type
39	l2	169	ILE
39	l2	179	LEU
39	l2	184	ARG
39	l2	191	LEU
39	l2	192	LYS
39	l2	193	ARG
39	l2	202	VAL
39	l2	204	MET
39	l2	218	HIS
39	l2	224	THR
39	l2	227	ARG
39	l2	230	VAL
39	l2	238	ILE
39	l2	243	THR
39	l2	246	LEU
40	l3	3	HIS
40	l3	4	ARG
40	l3	5	LYS
40	l3	7	GLU
40	l3	10	ARG
40	l3	17	LEU
40	l3	19	ARG
40	l3	25	ILE
40	l3	37	ARG
40	l3	44	THR
40	l3	45	SER
40	l3	47	LEU
40	l3	50	LYS
40	l3	55	THR
40	l3	69	LYS
40	l3	73	VAL
40	l3	77	THR
40	l3	81	THR
40	l3	85	VAL
40	l3	95	THR
40	l3	101	SER
40	l3	104	THR
40	l3	114	VAL
40	l3	120	LYS
40	l3	127	LYS
40	l3	140	ASP
40	l3	146	ARG

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Mol	Chain	Res	Type
40	l3	148	LEU
40	l3	150	ARG
40	l3	161	LEU
40	l3	164	THR
40	l3	167	ARG
40	l3	169	THR
40	l3	178	LEU
40	l3	184	ASN
40	l3	188	ILE
40	l3	205	VAL
40	l3	208	VAL
40	l3	212	ASN
40	l3	221	THR
40	l3	222	LYS
40	l3	232	ARG
40	l3	238	LEU
40	l3	242	THR
40	l3	244	ARG
40	l3	249	VAL
40	l3	252	ILE
40	l3	256	HIS
40	l3	261	MET
40	l3	264	VAL
40	l3	266	ARG
40	l3	274	SER
40	l3	278	ILE
40	l3	282	ILE
40	l3	287	LYS
40	l3	299	ASP
40	l3	300	ARG
40	l3	304	THR
40	l3	318	LYS
40	l3	319	ASN
40	l3	322	ILE
40	l3	328	ILE
40	l3	331	ASN
40	l3	332	ARG
40	l3	340	LYS
40	l3	342	LEU
40	l3	346	THR
40	l3	354	VAL
40	l3	356	LEU

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Mol	Chain	Res	Type
40	l3	359	ILE
40	l3	364	LYS
40	l3	380	MET
41	l4	3	ARG
41	l4	12	THR
41	l4	14	GLU
41	l4	22	LEU
41	l4	37	THR
41	l4	42	VAL
41	l4	52	VAL
41	l4	55	LYS
41	l4	63	GLU
41	l4	67	THR
41	l4	73	ARG
41	l4	76	ARG
41	l4	84	ARG
41	l4	90	PHE
41	l4	93	MET
41	l4	105	THR
41	l4	131	VAL
41	l4	136	LEU
41	l4	144	LYS
41	l4	145	ILE
41	l4	150	LEU
41	l4	151	VAL
41	l4	152	VAL
41	l4	176	SER
41	l4	178	LEU
41	l4	179	LEU
41	l4	182	LEU
41	l4	186	LYS
41	l4	187	LEU
41	l4	203	ARG
41	l4	206	LEU
41	l4	217	LYS
41	l4	222	VAL
41	l4	226	GLU
41	l4	230	VAL
41	l4	246	ARG
41	l4	249	ILE
41	l4	256	THR
41	l4	258	LEU

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Mol	Chain	Res	Type
41	14	265	GLU
41	14	276	LEU
41	14	278	SER
41	14	286	VAL
41	14	287	THR
41	14	289	ILE
41	14	292	SER
41	14	304	GLN
41	14	306	THR
41	14	307	GLN
41	14	310	THR
41	14	313	LEU
41	14	319	LYS
41	14	323	VAL
41	14	327	LEU
41	14	345	GLU
41	14	346	LYS
41	14	349	THR
41	14	354	VAL
41	14	357	GLU
42	15	34	LYS
42	15	37	VAL
42	15	41	LYS
42	15	51	LEU
42	15	66	SER
42	15	70	THR
42	15	89	THR
42	15	92	LEU
42	15	101	THR
42	15	110	LEU
42	15	115	LEU
42	15	116	ASP
42	15	118	THR
42	15	122	VAL
42	15	125	VAL
42	15	128	GLU
42	15	132	THR
42	15	133	GLU
42	15	135	VAL
42	15	140	ARG
42	15	144	VAL
42	15	146	LEU

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Mol	Chain	Res	Type
42	15	148	ILE
42	15	152	ARG
42	15	154	THR
42	15	158	ARG
42	15	173	VAL
42	15	176	SER
42	15	177	GLU
42	15	185	PHE
42	15	194	LEU
42	15	196	ARG
42	15	222	LEU
42	15	234	ASP
42	15	235	SER
42	15	241	THR
42	15	245	GLU
42	15	247	ILE
42	15	254	LYS
42	15	259	LYS
42	15	268	GLU
42	15	273	ARG
42	15	276	LYS
42	15	286	VAL
42	15	293	LEU
43	16	8	LYS
43	16	9	TRP
43	16	12	SER
43	16	14	ASP
43	16	21	THR
43	16	28	GLN
43	16	31	ARG
43	16	35	VAL
43	16	46	ARG
43	16	50	LYS
43	16	52	VAL
43	16	56	LYS
43	16	64	LEU
43	16	65	ILE
43	16	74	VAL
43	16	78	ARG
43	16	79	VAL
43	16	82	ARG
43	16	88	SER

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Mol	Chain	Res	Type
43	16	89	THR
43	16	90	LYS
43	16	91	VAL
43	16	103	VAL
43	16	104	GLU
43	16	108	LYS
43	16	109	GLU
43	16	131	LYS
43	16	152	THR
43	16	155	LEU
43	16	164	SER
43	16	169	ASP
43	16	170	LYS
43	16	174	LEU
44	17	22	THR
44	17	26	VAL
44	17	33	ARG
44	17	38	LYS
44	17	41	ARG
44	17	54	GLU
44	17	59	GLU
44	17	60	ARG
44	17	61	ASN
44	17	77	VAL
44	17	88	ARG
44	17	93	ASN
44	17	98	LYS
44	17	101	LYS
44	17	111	ILE
44	17	115	THR
44	17	119	VAL
44	17	124	LEU
44	17	129	LEU
44	17	130	ILE
44	17	142	SER
44	17	156	ILE
44	17	158	LYS
44	17	173	LEU
44	17	175	LYS
44	17	176	TYR
44	17	179	LEU
44	17	181	ILE

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Mol	Chain	Res	Type
44	17	184	LEU
44	17	193	PRO
44	17	206	LYS
44	17	208	SER
44	17	224	ILE
44	17	229	PHE
44	17	239	LEU
45	18	26	LEU
45	18	38	GLN
45	18	41	GLN
45	18	50	VAL
45	18	68	ARG
45	18	71	VAL
45	18	74	THR
45	18	77	GLN
45	18	79	GLN
45	18	83	ASP
45	18	85	ASN
45	18	91	PHE
45	18	93	LEU
45	18	95	ASN
45	18	111	LYS
45	18	134	TYR
45	18	136	LEU
45	18	146	LYS
45	18	149	LYS
45	18	150	LEU
45	18	153	ILE
45	18	156	ASP
45	18	157	VAL
45	18	160	ILE
45	18	183	LYS
45	18	185	ARG
45	18	191	ASN
45	18	203	VAL
45	18	211	LEU
45	18	213	LYS
45	18	217	THR
45	18	224	ASP
45	18	230	LYS
45	18	231	LYS
45	18	245	LYS

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Mol	Chain	Res	Type
45	18	248	LYS
46	19	1	MET
46	19	5	GLN
46	19	6	THR
46	19	16	VAL
46	19	17	THR
46	19	18	VAL
46	19	19	SER
46	19	31	ARG
46	19	33	THR
46	19	39	LYS
46	19	46	THR
46	19	52	LEU
46	19	55	VAL
46	19	68	LEU
46	19	70	THR
46	19	72	LYS
46	19	76	ASP
46	19	77	ASN
46	19	82	VAL
46	19	87	LYS
46	19	91	ARG
46	19	92	TYR
46	19	96	HIS
46	19	102	ASN
46	19	105	GLU
46	19	106	LYS
46	19	107	ASP
46	19	113	GLU
46	19	115	ARG
46	19	118	LEU
46	19	121	LYS
46	19	132	VAL
46	19	133	THR
46	19	138	THR
46	19	144	ILE
46	19	146	LEU
46	19	151	VAL
46	19	157	ASN
46	19	161	LEU
46	19	162	GLN
46	19	166	ARG

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Mol	Chain	Res	Type
46	l9	175	PHE
46	l9	181	VAL
47	m0	3	ARG
47	m0	4	ARG
47	m0	19	LYS
47	m0	24	ARG
47	m0	29	SER
47	m0	35	ASP
47	m0	39	LYS
47	m0	42	THR
47	m0	44	ASP
47	m0	48	LEU
47	m0	52	LEU
47	m0	57	LEU
47	m0	58	GLU
47	m0	61	SER
47	m0	71	CYS
47	m0	80	SER
47	m0	91	VAL
47	m0	121	LYS
47	m0	125	LEU
47	m0	133	GLN
47	m0	139	ARG
47	m0	143	SER
47	m0	158	LYS
47	m0	163	GLN
47	m0	169	LYS
47	m0	176	LEU
47	m0	177	ASP
47	m0	184	LYS
47	m0	197	VAL
47	m0	206	LEU
47	m0	208	ASN
47	m0	210	ILE
47	m0	212	GLU
47	m0	215	GLU
47	m0	217	PHE
48	m1	7	ASN
48	m1	9	MET
48	m1	10	ARG
48	m1	12	LEU
48	m1	13	LYS

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Mol	Chain	Res	Type
48	m1	16	LYS
48	m1	19	LEU
48	m1	35	LYS
48	m1	40	LEU
48	m1	55	ARG
48	m1	57	PHE
48	m1	59	ILE
48	m1	60	ARG
48	m1	77	GLU
48	m1	78	GLU
48	m1	80	LEU
48	m1	82	ARG
48	m1	85	LYS
48	m1	97	SER
48	m1	99	THR
48	m1	106	ILE
48	m1	107	ASP
48	m1	110	ILE
48	m1	112	LEU
48	m1	127	PHE
48	m1	130	VAL
48	m1	137	ARG
48	m1	140	ARG
48	m1	143	ARG
48	m1	148	VAL
48	m1	150	ASN
48	m1	152	HIS
48	m1	154	THR
48	m1	157	GLU
48	m1	160	VAL
48	m1	171	VAL
49	m3	4	SER
49	m3	16	LYS
49	m3	42	ARG
49	m3	46	ILE
49	m3	53	LEU
49	m3	54	LEU
49	m3	58	VAL
49	m3	62	THR
49	m3	67	ARG
49	m3	69	VAL
49	m3	73	ARG

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Mol	Chain	Res	Type
49	m3	76	THR
49	m3	80	VAL
49	m3	86	THR
49	m3	92	THR
49	m3	104	ARG
49	m3	113	VAL
49	m3	118	GLU
49	m3	123	ILE
49	m3	124	ILE
49	m3	131	LYS
49	m3	137	GLN
49	m3	150	PRO
49	m3	152	THR
49	m3	153	ASP
49	m3	154	VAL
49	m3	155	GLU
49	m3	157	ARG
49	m3	162	ASN
49	m3	164	GLU
49	m3	165	SER
49	m3	183	ARG
49	m3	184	GLU
49	m3	189	GLU
50	m4	3	THR
50	m4	8	LYS
50	m4	13	ARG
50	m4	16	GLU
50	m4	20	VAL
50	m4	24	LYS
50	m4	46	ILE
50	m4	53	VAL
50	m4	62	GLN
50	m4	63	VAL
50	m4	64	VAL
50	m4	66	THR
50	m4	72	LEU
50	m4	80	THR
50	m4	82	SER
50	m4	90	VAL
50	m4	106	ARG
50	m4	108	ARG
50	m4	119	GLN

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Mol	Chain	Res	Type
50	m4	124	ARG
50	m4	126	GLN
50	m4	128	ARG
51	m5	5	LYS
51	m5	10	LEU
51	m5	12	ARG
51	m5	14	LYS
51	m5	16	SER
51	m5	20	ARG
51	m5	22	LEU
51	m5	24	ARG
51	m5	25	VAL
51	m5	32	GLN
51	m5	36	ILE
51	m5	49	ARG
51	m5	67	ARG
51	m5	80	THR
51	m5	90	ASN
51	m5	92	LEU
51	m5	96	ARG
51	m5	108	ARG
51	m5	109	ARG
51	m5	112	ASN
51	m5	116	LEU
51	m5	151	ILE
51	m5	153	ASP
51	m5	155	VAL
51	m5	159	ARG
51	m5	171	SER
51	m5	174	ILE
51	m5	175	ASN
51	m5	176	LYS
51	m5	188	ARG
51	m5	189	LYS
51	m5	190	THR
51	m5	194	GLN
51	m5	204	LYS
52	m6	4	GLU
52	m6	12	LYS
52	m6	15	LEU
52	m6	34	VAL
52	m6	41	LEU

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Mol	Chain	Res	Type
52	m6	42	ASN
52	m6	46	GLU
52	m6	57	PHE
52	m6	59	ARG
52	m6	60	LYS
52	m6	66	LYS
52	m6	68	ARG
52	m6	74	ARG
52	m6	78	ARG
52	m6	82	LYS
52	m6	84	LEU
52	m6	85	ARG
52	m6	88	VAL
52	m6	89	SER
52	m6	100	GLU
52	m6	104	VAL
52	m6	106	GLU
52	m6	108	ILE
52	m6	115	LYS
52	m6	117	ARG
52	m6	124	LEU
52	m6	126	VAL
52	m6	129	LEU
52	m6	133	ARG
52	m6	143	THR
52	m6	145	VAL
52	m6	160	ARG
52	m6	167	TYR
52	m6	177	LYS
52	m6	182	ASN
52	m6	187	GLU
52	m6	190	VAL
52	m6	192	LYS
52	m6	194	LEU
52	m6	197	LEU
53	m7	3	ARG
53	m7	7	THR
53	m7	9	THR
53	m7	31	GLU
53	m7	32	THR
53	m7	52	LEU
53	m7	56	ARG

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Mol	Chain	Res	Type
53	m7	69	ARG
53	m7	70	THR
53	m7	80	LYS
53	m7	86	LYS
53	m7	89	LYS
53	m7	96	GLN
53	m7	103	GLU
53	m7	112	LEU
53	m7	114	VAL
53	m7	119	VAL
53	m7	120	ASN
53	m7	127	ARG
53	m7	128	ARG
53	m7	133	HIS
53	m7	136	ILE
53	m7	142	SER
53	m7	148	LEU
54	m8	3	ILE
54	m8	7	SER
54	m8	12	ARG
54	m8	17	THR
54	m8	22	ASP
54	m8	24	VAL
54	m8	26	LEU
54	m8	32	LEU
54	m8	34	THR
54	m8	49	LEU
54	m8	53	PHE
54	m8	55	SER
54	m8	57	ILE
54	m8	66	ARG
54	m8	69	ARG
54	m8	80	THR
54	m8	81	VAL
54	m8	86	THR
54	m8	93	ILE
54	m8	111	ARG
54	m8	113	LYS
54	m8	135	GLN
54	m8	136	ASN
54	m8	138	LEU
54	m8	144	ARG

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Mol	Chain	Res	Type
54	m8	145	ASN
54	m8	150	VAL
54	m8	161	LYS
54	m8	165	ILE
54	m8	166	LEU
54	m8	170	ARG
54	m8	176	ARG
54	m8	178	ARG
54	m8	182	LYS
55	m9	4	LEU
55	m9	5	ARG
55	m9	7	GLN
55	m9	10	LEU
55	m9	20	ARG
55	m9	31	GLU
55	m9	34	GLN
55	m9	37	SER
55	m9	56	THR
55	m9	57	VAL
55	m9	61	SER
55	m9	74	ARG
55	m9	81	ARG
55	m9	85	ARG
55	m9	88	ARG
55	m9	98	ARG
55	m9	99	LEU
55	m9	105	LEU
55	m9	106	LEU
55	m9	127	SER
55	m9	138	LEU
55	m9	152	GLU
55	m9	153	LYS
55	m9	164	LEU
55	m9	170	ARG
55	m9	173	ARG
55	m9	177	VAL
55	m9	182	ASP
56	n0	1	MET
56	n0	16	THR
56	n0	32	SER
56	n0	45	LEU
56	n0	47	LYS

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Mol	Chain	Res	Type
56	n0	52	LYS
56	n0	62	ASN
56	n0	73	LYS
56	n0	80	ARG
56	n0	89	ASN
56	n0	92	LYS
56	n0	95	ARG
56	n0	96	ASP
56	n0	97	VAL
56	n0	100	VAL
56	n0	107	TYR
56	n0	115	ARG
56	n0	117	ARG
56	n0	125	LYS
56	n0	130	GLU
56	n0	137	ARG
56	n0	138	GLN
56	n0	145	THR
56	n0	148	LEU
56	n0	149	LYS
56	n0	155	ARG
56	n0	157	GLN
56	n0	160	THR
56	n0	161	LYS
56	n0	162	THR
56	n0	164	SER
56	n0	167	ARG
56	n0	171	PHE
56	n0	172	TYR
57	n1	25	VAL
57	n1	26	HIS
57	n1	27	LEU
57	n1	28	SER
57	n1	35	LYS
57	n1	47	SER
57	n1	60	LYS
57	n1	68	THR
57	n1	72	VAL
57	n1	78	LYS
57	n1	80	VAL
57	n1	88	ARG
57	n1	89	LEU

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Mol	Chain	Res	Type
57	n1	102	ARG
57	n1	104	GLU
57	n1	112	ASN
57	n1	118	GLU
57	n1	126	VAL
57	n1	127	GLN
57	n1	128	LEU
57	n1	131	GLN
57	n1	134	GLN
57	n1	138	SER
57	n1	140	ILE
57	n1	143	THR
57	n1	149	GLN
57	n1	150	THR
57	n1	151	LEU
57	n1	157	GLU
58	n2	13	LYS
58	n2	16	THR
58	n2	19	VAL
58	n2	27	VAL
58	n2	38	ILE
58	n2	39	ASP
58	n2	43	VAL
58	n2	47	VAL
58	n2	49	ASN
58	n2	54	VAL
58	n2	57	THR
58	n2	58	GLU
58	n2	59	ASP
58	n2	63	VAL
58	n2	64	THR
58	n2	66	VAL
58	n2	68	THR
58	n2	74	LYS
58	n2	75	TYR
58	n2	80	THR
58	n2	88	GLN
58	n2	90	ARG
58	n2	93	ILE
58	n2	95	PHE
58	n2	96	VAL
58	n2	98	THR

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Mol	Chain	Res	Type
58	n2	100	THR
59	n3	4	ASN
59	n3	7	GLN
59	n3	9	THR
59	n3	10	LYS
59	n3	14	SER
59	n3	22	ILE
59	n3	23	MET
59	n3	61	THR
59	n3	69	LEU
59	n3	70	ARG
59	n3	73	VAL
59	n3	74	MET
59	n3	83	LYS
59	n3	88	ARG
59	n3	91	VAL
59	n3	93	LEU
59	n3	104	ASN
59	n3	112	SER
59	n3	120	LYS
60	n4	1	MET
60	n4	2	LYS
60	n4	7	SER
60	n4	9	SER
60	n4	19	THR
60	n4	30	ARG
60	n4	39	LEU
60	n4	54	LEU
60	n4	57	LYS
60	n4	63	ILE
60	n4	89	LEU
60	n4	96	LEU
60	n4	97	LYS
60	n4	100	VAL
60	n4	105	ARG
60	n4	107	GLU
60	n4	112	ASN
60	n4	114	GLU
60	n4	119	GLU
60	n4	127	LYS
60	n4	134	GLN
61	n5	24	LEU

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Mol	Chain	Res	Type
61	n5	27	ARG
61	n5	34	LEU
61	n5	37	THR
61	n5	56	ARG
61	n5	57	LEU
61	n5	63	ILE
61	n5	71	THR
61	n5	86	VAL
61	n5	87	SER
61	n5	108	LEU
61	n5	109	LYS
61	n5	115	ARG
61	n5	119	THR
61	n5	124	VAL
61	n5	125	ARG
61	n5	126	LEU
61	n5	133	LEU
61	n5	135	ILE
62	n6	3	LYS
62	n6	6	LEU
62	n6	12	ARG
62	n6	17	LYS
62	n6	28	ARG
62	n6	32	SER
62	n6	35	LEU
62	n6	36	SER
62	n6	37	LYS
62	n6	39	LEU
62	n6	50	ILE
62	n6	51	ARG
62	n6	55	GLU
62	n6	56	VAL
62	n6	66	GLN
62	n6	74	TYR
62	n6	95	VAL
62	n6	105	VAL
62	n6	106	ILE
62	n6	113	LYS
62	n6	115	ARG
62	n6	120	GLN
62	n6	126	LEU
62	n6	127	GLU

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Mol	Chain	Res	Type
63	n7	3	LYS
63	n7	5	LEU
63	n7	14	VAL
63	n7	17	ARG
63	n7	24	VAL
63	n7	27	LYS
63	n7	28	PRO
63	n7	33	SER
63	n7	34	LYS
63	n7	51	LEU
63	n7	52	LYS
63	n7	67	LYS
63	n7	72	ILE
63	n7	73	LYS
63	n7	75	VAL
63	n7	81	LEU
63	n7	83	THR
63	n7	85	TYR
63	n7	87	LEU
63	n7	93	LYS
63	n7	94	SER
63	n7	99	GLU
63	n7	100	THR
63	n7	102	GLU
63	n7	103	GLN
63	n7	128	GLN
63	n7	134	LEU
63	n7	135	ARG
64	n8	3	SER
64	n8	6	THR
64	n8	8	THR
64	n8	9	ARG
64	n8	10	LYS
64	n8	12	ARG
64	n8	15	VAL
64	n8	22	ILE
64	n8	24	LYS
64	n8	25	HIS
64	n8	26	ARG
64	n8	27	LYS
64	n8	43	ILE
64	n8	47	LYS

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Mol	Chain	Res	Type
64	n8	56	VAL
64	n8	60	TYR
64	n8	65	GLN
64	n8	70	LYS
64	n8	73	LEU
64	n8	88	ASP
64	n8	91	LEU
64	n8	97	GLU
64	n8	117	ARG
64	n8	128	ARG
64	n8	132	LYS
64	n8	133	LEU
64	n8	139	ARG
65	n9	3	LYS
65	n9	4	SER
65	n9	7	HIS
65	n9	8	THR
65	n9	13	THR
65	n9	14	ARG
65	n9	15	LYS
65	n9	17	HIS
65	n9	19	ASN
65	n9	21	ILE
65	n9	22	LYS
65	n9	26	THR
65	n9	31	SER
65	n9	38	LYS
65	n9	58	LYS
65	n9	59	LYS
66	o0	8	GLU
66	o0	13	LYS
66	o0	19	LYS
66	o0	32	LYS
66	o0	33	SER
66	o0	34	LEU
66	o0	51	LEU
66	o0	61	MET
66	o0	71	GLN
66	o0	74	ASN
66	o0	84	LEU
66	o0	86	ARG
66	o0	87	VAL

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Mol	Chain	Res	Type
66	o0	101	LEU
66	o0	103	THR
67	o1	6	ASP
67	o1	13	THR
67	o1	16	LEU
67	o1	24	SER
67	o1	26	LYS
67	o1	28	ARG
67	o1	31	ARG
67	o1	42	LEU
67	o1	44	MET
67	o1	46	THR
67	o1	55	LEU
67	o1	64	VAL
67	o1	67	VAL
67	o1	68	GLU
67	o1	71	LEU
67	o1	73	LEU
67	o1	75	ILE
67	o1	76	SER
67	o1	83	GLU
67	o1	87	ASN
67	o1	90	PHE
67	o1	91	SER
67	o1	93	VAL
67	o1	96	VAL
67	o1	97	LEU
67	o1	100	SER
67	o1	102	LYS
67	o1	104	LEU
67	o1	106	THR
67	o1	107	VAL
67	o1	110	GLU
68	o2	4	LEU
68	o2	6	HIS
68	o2	8	LYS
68	o2	10	VAL
68	o2	14	THR
68	o2	19	ARG
68	o2	21	HIS
68	o2	24	ARG
68	o2	27	ARG

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Mol	Chain	Res	Type
68	o2	33	ARG
68	o2	35	GLN
68	o2	40	SER
68	o2	41	VAL
68	o2	44	ARG
68	o2	45	ARG
68	o2	51	SER
68	o2	59	SER
68	o2	61	LYS
68	o2	67	SER
68	o2	73	THR
68	o2	75	LEU
68	o2	91	THR
68	o2	105	ARG
68	o2	109	LEU
68	o2	119	VAL
68	o2	120	THR
68	o2	125	ARG
68	o2	126	LEU
69	o3	9	VAL
69	o3	22	VAL
69	o3	31	LYS
69	o3	37	THR
69	o3	42	GLN
69	o3	57	LYS
69	o3	58	GLU
69	o3	59	VAL
69	o3	60	ARG
69	o3	70	LYS
69	o3	73	ARG
69	o3	74	THR
69	o3	93	THR
69	o3	97	SER
69	o3	98	VAL
70	o4	4	ARG
70	o4	10	ARG
70	o4	17	SER
70	o4	22	VAL
70	o4	25	THR
70	o4	30	LEU
70	o4	47	CYS
70	o4	58	ARG

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Mol	Chain	Res	Type
70	o4	66	SER
70	o4	79	SER
70	o4	86	LYS
70	o4	98	GLN
70	o4	104	VAL
71	o5	4	VAL
71	o5	5	LYS
71	o5	20	GLN
71	o5	21	LEU
71	o5	26	LYS
71	o5	27	GLU
71	o5	28	LEU
71	o5	30	GLU
71	o5	31	LEU
71	o5	37	SER
71	o5	38	ARG
71	o5	46	THR
71	o5	47	VAL
71	o5	48	ARG
71	o5	57	VAL
71	o5	62	GLN
71	o5	68	GLN
71	o5	69	LEU
71	o5	76	GLN
71	o5	81	ARG
71	o5	84	LYS
71	o5	89	ARG
71	o5	90	ARG
71	o5	93	THR
71	o5	107	LYS
71	o5	108	GLN
71	o5	113	GLN
72	o6	7	ILE
72	o6	9	ILE
72	o6	13	LYS
72	o6	17	VAL
72	o6	20	MET
72	o6	21	THR
72	o6	27	SER
72	o6	29	LYS
72	o6	35	ASN
72	o6	36	ARG

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Mol	Chain	Res	Type
72	o6	37	THR
72	o6	42	SER
72	o6	43	LEU
72	o6	45	ARG
72	o6	57	LEU
72	o6	58	ILE
72	o6	60	LEU
72	o6	68	ARG
72	o6	70	ARG
72	o6	74	LYS
72	o6	75	LYS
72	o6	76	ARG
72	o6	79	SER
72	o6	81	THR
72	o6	94	ILE
72	o6	98	ARG
72	o6	99	ARG
73	o7	15	SER
73	o7	17	THR
73	o7	21	ARG
73	o7	24	ARG
73	o7	25	ARG
73	o7	33	THR
73	o7	36	SER
73	o7	46	SER
73	o7	54	LYS
73	o7	55	ARG
73	o7	59	THR
73	o7	64	MET
73	o7	65	ARG
73	o7	67	LEU
73	o7	73	ARG
73	o7	75	LYS
73	o7	76	ASN
73	o7	85	LYS
74	o8	8	ILE
74	o8	12	LEU
74	o8	15	THR
74	o8	16	ARG
74	o8	17	ARG
74	o8	19	ASP
74	o8	24	THR

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Mol	Chain	Res	Type
74	o8	27	ILE
74	o8	41	THR
74	o8	46	ARG
74	o8	48	SER
74	o8	53	THR
74	o8	55	VAL
74	o8	61	LYS
74	o8	63	LYS
74	o8	64	LYS
74	o8	65	LEU
74	o8	72	THR
75	o9	4	GLN
75	o9	9	ILE
75	o9	11	GLN
75	o9	15	LYS
75	o9	19	GLN
75	o9	21	ARG
75	o9	23	LEU
75	o9	28	ARG
75	o9	29	LEU
75	o9	36	ARG
75	o9	45	ARG
75	o9	48	LYS
76	q0	77	ILE
76	q0	78	ILE
76	q0	79	GLU
76	q0	85	LEU
76	q0	91	CYS
76	q0	93	LYS
76	q0	94	SER
76	q0	95	VAL
76	q0	98	LYS
76	q0	99	CYS
76	q0	106	ARG
76	q0	108	THR
76	q0	112	LYS
76	q0	114	LYS
76	q0	127	LEU
77	q1	1	MET
77	q1	2	ARG
77	q1	6	ARG
77	q1	9	ARG

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Mol	Chain	Res	Type
77	q1	14	LYS
77	q1	18	ARG
77	q1	20	VAL
77	q1	21	ARG
77	q1	23	ARG
78	q2	7	THR
78	q2	8	ARG
78	q2	28	TYR
78	q2	38	GLN
78	q2	45	ARG
78	q2	61	LYS
78	q2	71	ARG
78	q2	72	LEU
78	q2	79	THR
78	q2	83	LEU
78	q2	84	THR
78	q2	85	LEU
78	q2	89	LYS
78	q2	92	GLU
78	q2	93	LEU
78	q2	98	LYS
78	q2	100	LYS
78	q2	104	LEU
78	q2	105	GLN
78	q2	106	PHE
79	q3	3	LYS
79	q3	8	VAL
79	q3	10	ILE
79	q3	20	SER
79	q3	21	SER
79	q3	33	GLN
79	q3	40	SER
79	q3	42	CYS
79	q3	49	ARG
79	q3	54	ILE
79	q3	56	THR
79	q3	58	SER
79	q3	59	CYS
79	q3	64	VAL
79	q3	70	THR
79	q3	78	THR
79	q3	85	ARG

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Mol	Chain	Res	Type
81	p0	4	ILE
81	p0	5	ARG
81	p0	30	VAL
81	p0	43	LYS
81	p0	48	ARG
81	p0	51	VAL
81	p0	57	THR
81	p0	67	LEU
81	p0	68	SER
81	p0	69	ASP
81	p0	70	LEU
81	p0	72	ASP
81	p0	81	LYS
81	p0	84	VAL
81	p0	89	THR
81	p0	93	LEU
81	p0	95	GLU
81	p0	97	LYS
81	p0	104	ARG
81	p0	192	ASP
81	p0	193	ASN

Some sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (73) such sidechains are listed below:

Mol	Chain	Res	Type
2	S0	32	HIS
5	S3	111	ASN
5	S3	179	GLN
6	S4	17	HIS
6	S4	50	ASN
6	S4	98	ASN
7	S5	103	ASN
9	S7	5	GLN
9	S7	71	HIS
10	S8	9	HIS
13	C1	104	HIS
15	C3	5	HIS
15	C3	105	ASN
17	C5	82	ASN
18	C6	83	GLN
20	C8	89	GLN
24	D2	24	GLN

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Mol	Chain	Res	Type
24	D2	56	HIS
29	D7	5	GLN
31	D9	48	ASN
34	SR	153	GLN
39	L2	209	HIS
40	L3	256	HIS
41	L4	5	GLN
44	L7	64	GLN
45	L8	45	ASN
46	L9	49	ASN
46	L9	59	ASN
46	L9	116	ASN
47	M0	12	GLN
49	M3	25	HIS
52	M6	29	ASN
54	M8	145	ASN
58	N2	52	ASN
59	N3	33	ASN
59	N3	98	ASN
63	N7	57	HIS
64	N8	74	ASN
65	N9	45	HIS
70	O4	18	ASN
72	O6	91	ASN
2	s0	39	ASN
2	s0	46	HIS
3	s1	148	ASN
3	s1	149	GLN
8	s6	210	GLN
11	s9	124	HIS
12	c0	32	HIS
12	c0	39	ASN
19	c7	29	GLN
20	c8	90	ASN
21	c9	101	ASN
22	d0	48	HIS
25	d3	10	ASN
25	d3	18	HIS
26	d4	77	ASN
28	d6	43	ASN
33	e1	151	ASN
34	sR	198	ASN

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Mol	Chain	Res	Type
35	sM	71	ASN
41	l4	221	ASN
45	l8	41	GLN
47	m0	12	GLN
51	m5	178	HIS
53	m7	137	ASN
57	n1	98	HIS
61	n5	55	ASN
62	n6	66	GLN
63	n7	123	GLN
64	n8	28	HIS
69	o3	88	ASN
70	o4	18	ASN
78	q2	38	GLN

5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	2	1747/1800 (97%)	541 (30%)	0
1	6	1792/1800 (99%)	494 (27%)	0
36	1	3145/3396 (92%)	774 (24%)	0
36	5	3145/3396 (92%)	801 (25%)	0
37	3	120/121 (99%)	20 (16%)	0
37	7	120/121 (99%)	28 (23%)	0
38	4	157/158 (99%)	42 (26%)	0
38	8	157/158 (99%)	44 (28%)	0
All	All	10383/10950 (94%)	2744 (26%)	0

All (2744) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	2	2	A
1	2	4	C
1	2	10	G
1	2	25	C
1	2	26	A
1	2	27	U
1	2	31	C
1	2	34	G
1	2	45	U
1	2	46	A

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Mol	Chain	Res	Type
1	2	47	A
1	2	54	C
1	2	57	G
1	2	60	U
1	2	66	U
1	2	68	A
1	2	69	G
1	2	72	A
1	2	73	U
1	2	74	U
1	2	75	U
1	2	76	A
1	2	80	A
1	2	104	A
1	2	114	C
1	2	116	U
1	2	128	U
1	2	131	C
1	2	132	U
1	2	133	U
1	2	134	U
1	2	135	A
1	2	136	C
1	2	137	U
1	2	140	A
1	2	141	U
1	2	144	U
1	2	145	A
1	2	146	U
1	2	151	G
1	2	153	G
1	2	158	U
1	2	159	U
1	2	161	U
1	2	166	C
1	2	167	U
1	2	169	A
1	2	176	C
1	2	178	U
1	2	179	A
1	2	185	U
1	2	186	C

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Mol	Chain	Res	Type
1	2	187	G
1	2	188	A
1	2	190	C
1	2	191	C
1	2	192	U
1	2	193	U
1	2	194	U
1	2	195	G
1	2	196	G
1	2	197	A
1	2	198	A
1	2	200	A
1	2	215	A
1	2	217	A
1	2	218	A
1	2	219	A
1	2	226	A
1	2	227	U
1	2	228	G
1	2	229	U
1	2	231	U
1	2	233	C
1	2	234	G
1	2	235	G
1	2	236	A
1	2	238	U
1	2	239	C
1	2	240	U
1	2	241	U
1	2	242	U
1	2	246	G
1	2	249	U
1	2	250	C
1	2	260	U
1	2	261	U
1	2	265	A
1	2	270	C
1	2	271	A
1	2	272	U
1	2	274	G
1	2	275	C
1	2	276	C

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Mol	Chain	Res	Type
1	2	277	U
1	2	278	U
1	2	279	G
1	2	280	U
1	2	281	G
1	2	284	G
1	2	285	G
1	2	288	A
1	2	290	G
1	2	299	A
1	2	302	U
1	2	308	C
1	2	309	C
1	2	313	U
1	2	314	C
1	2	316	A
1	2	319	U
1	2	321	C
1	2	322	G
1	2	337	G
1	2	338	C
1	2	339	C
1	2	346	G
1	2	352	A
1	2	359	A
1	2	360	A
1	2	361	C
1	2	387	A
1	2	390	G
1	2	393	C
1	2	397	A
1	2	400	A
1	2	401	A
1	2	402	C
1	2	404	G
1	2	416	A
1	2	418	G
1	2	419	G
1	2	424	C
1	2	425	A
1	2	426	G
1	2	428	A

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Mol	Chain	Res	Type
1	2	430	G
1	2	434	G
1	2	437	A
1	2	439	U
1	2	444	C
1	2	446	A
1	2	448	C
1	2	454	U
1	2	455	C
1	2	459	G
1	2	468	A
1	2	470	A
1	2	473	A
1	2	475	A
1	2	477	A
1	2	480	G
1	2	483	A
1	2	484	C
1	2	485	A
1	2	488	G
1	2	493	U
1	2	494	U
1	2	495	C
1	2	496	G
1	2	497	G
1	2	498	G
1	2	499	U
1	2	500	C
1	2	502	U
1	2	503	G
1	2	504	U
1	2	505	A
1	2	506	A
1	2	507	U
1	2	508	U
1	2	510	G
1	2	511	A
1	2	512	A
1	2	513	U
1	2	514	G
1	2	515	A
1	2	516	G

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Mol	Chain	Res	Type
1	2	527	A
1	2	532	U
1	2	533	U
1	2	536	C
1	2	538	A
1	2	539	G
1	2	540	G
1	2	541	A
1	2	542	A
1	2	543	C
1	2	544	A
1	2	546	U
1	2	548	G
1	2	554	C
1	2	555	A
1	2	556	A
1	2	557	G
1	2	558	U
1	2	559	C
1	2	565	C
1	2	579	A
1	2	580	A
1	2	582	U
1	2	585	A
1	2	594	A
1	2	595	G
1	2	597	G
1	2	598	U
1	2	611	U
1	2	619	A
1	2	620	A
1	2	621	A
1	2	622	A
1	2	623	A
1	2	624	G
1	2	630	A
1	2	631	G
1	2	639	U
1	2	640	U
1	2	650	U
1	2	652	G
1	2	653	C

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Mol	Chain	Res	Type
1	2	654	C
1	2	656	G
1	2	658	C
1	2	677	G
1	2	680	U
1	2	682	C
1	2	684	A
1	2	685	A
1	2	686	C
1	2	692	C
1	2	694	U
1	2	696	C
1	2	697	C
1	2	698	U
1	2	699	U
1	2	700	C
1	2	702	G
1	2	703	G
1	2	704	C
1	2	705	U
1	2	706	A
1	2	707	A
1	2	708	C
1	2	709	C
1	2	710	U
1	2	711	U
1	2	712	G
1	2	713	A
1	2	714	G
1	2	716	C
1	2	717	C
1	2	718	U
1	2	719	U
1	2	720	G
1	2	721	U
1	2	722	G
1	2	723	G
1	2	725	U
1	2	727	U
1	2	728	U
1	2	731	C
1	2	732	G

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Mol	Chain	Res	Type
1	2	733	A
1	2	734	A
1	2	735	C
1	2	736	C
1	2	737	A
1	2	738	G
1	2	742	U
1	2	745	U
1	2	754	A
1	2	755	A
1	2	756	A
1	2	765	G
1	2	766	U
1	2	771	A
1	2	774	A
1	2	775	G
1	2	778	G
1	2	779	U
1	2	781	U
1	2	782	U
1	2	783	G
1	2	784	C
1	2	787	G
1	2	789	A
1	2	793	A
1	2	794	U
1	2	795	U
1	2	796	A
1	2	806	A
1	2	807	A
1	2	811	A
1	2	812	A
1	2	814	A
1	2	815	G
1	2	816	G
1	2	818	C
1	2	819	G
1	2	820	U
1	2	821	U
1	2	823	G
1	2	824	G
1	2	829	A

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Mol	Chain	Res	Type
1	2	830	U
1	2	831	U
1	2	833	U
1	2	846	G
1	2	848	C
1	2	852	C
1	2	855	A
1	2	856	A
1	2	862	A
1	2	863	A
1	2	876	G
1	2	886	U
1	2	892	A
1	2	897	C
1	2	898	A
1	2	908	U
1	2	912	U
1	2	913	G
1	2	914	G
1	2	915	A
1	2	916	U
1	2	921	U
1	2	933	A
1	2	935	U
1	2	942	G
1	2	952	A
1	2	960	U
1	2	964	U
1	2	966	A
1	2	975	C
1	2	977	A
1	2	988	A
1	2	991	G
1	2	992	A
1	2	993	A
1	2	995	A
1	2	997	G
1	2	1001	A
1	2	1002	G
1	2	1003	A
1	2	1004	U
1	2	1005	A

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Mol	Chain	Res	Type
1	2	1011	G
1	2	1014	G
1	2	1026	A
1	2	1028	C
1	2	1032	G
1	2	1039	A
1	2	1040	G
1	2	1043	A
1	2	1052	U
1	2	1053	G
1	2	1058	U
1	2	1059	U
1	2	1060	U
1	2	1061	A
1	2	1064	G
1	2	1074	G
1	2	1076	A
1	2	1079	U
1	2	1080	U
1	2	1082	C
1	2	1086	A
1	2	1091	A
1	2	1092	A
1	2	1093	A
1	2	1096	C
1	2	1097	U
1	2	1100	G
1	2	1109	G
1	2	1138	A
1	2	1139	A
1	2	1141	G
1	2	1143	A
1	2	1146	G
1	2	1150	G
1	2	1151	A
1	2	1155	G
1	2	1157	A
1	2	1158	C
1	2	1160	A
1	2	1164	G
1	2	1167	G
1	2	1168	U

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Mol	Chain	Res	Type
1	2	1185	U
1	2	1187	U
1	2	1191	U
1	2	1194	A
1	2	1196	A
1	2	1199	G
1	2	1200	G
1	2	1202	A
1	2	1207	C
1	2	1217	A
1	2	1218	G
1	2	1226	A
1	2	1227	A
1	2	1228	G
1	2	1229	G
1	2	1235	C
1	2	1238	A
1	2	1241	G
1	2	1243	G
1	2	1244	A
1	2	1245	G
1	2	1250	U
1	2	1251	U
1	2	1258	U
1	2	1273	G
1	2	1276	U
1	2	1284	C
1	2	1285	U
1	2	1286	U
1	2	1288	G
1	2	1307	U
1	2	1314	U
1	2	1315	U
1	2	1320	U
1	2	1321	A
1	2	1332	C
1	2	1337	A
1	2	1339	C
1	2	1340	U
1	2	1341	A
1	2	1344	A
1	2	1345	A

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Mol	Chain	Res	Type
1	2	1346	A
1	2	1354	G
1	2	1355	C
1	2	1363	U
1	2	1364	G
1	2	1370	U
1	2	1371	A
1	2	1372	U
1	2	1379	C
1	2	1384	A
1	2	1388	A
1	2	1390	U
1	2	1391	A
1	2	1395	G
1	2	1398	U
1	2	1399	C
1	2	1400	A
1	2	1412	G
1	2	1413	U
1	2	1414	U
1	2	1415	U
1	2	1418	G
1	2	1427	A
1	2	1428	G
1	2	1432	U
1	2	1433	G
1	2	1445	G
1	2	1446	A
1	2	1447	C
1	2	1448	G
1	2	1457	C
1	2	1458	G
1	2	1459	C
1	2	1460	A
1	2	1461	C
1	2	1471	A
1	2	1473	U
1	2	1474	G
1	2	1478	G
1	2	1482	C
1	2	1483	A
1	2	1487	A

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Mol	Chain	Res	Type
1	2	1488	G
1	2	1489	U
1	2	1490	C
1	2	1491	U
1	2	1492	A
1	2	1493	A
1	2	1499	G
1	2	1501	C
1	2	1506	G
1	2	1514	U
1	2	1515	A
1	2	1516	A
1	2	1517	U
1	2	1521	G
1	2	1523	G
1	2	1524	A
1	2	1526	A
1	2	1533	C
1	2	1535	U
1	2	1536	G
1	2	1537	C
1	2	1538	U
1	2	1539	G
1	2	1542	G
1	2	1555	A
1	2	1557	U
1	2	1559	A
1	2	1569	A
1	2	1573	A
1	2	1574	G
1	2	1576	A
1	2	1584	G
1	2	1590	G
1	2	1596	C
1	2	1601	G
1	2	1614	A
1	2	1616	G
1	2	1619	C
1	2	1624	C
1	2	1626	U
1	2	1631	A
1	2	1634	C

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Mol	Chain	Res	Type
1	2	1637	C
1	2	1650	U
1	2	1657	U
1	2	1658	G
1	2	1666	U
1	2	1682	U
1	2	1683	C
1	2	1684	U
1	2	1731	A
1	2	1750	A
1	2	1754	A
1	2	1755	A
1	2	1756	A
1	2	1760	G
1	2	1762	A
1	2	1766	A
1	2	1769	U
1	2	1770	U
1	2	1780	G
1	2	1782	A
1	2	1783	C
1	2	1789	G
1	2	1790	A
1	2	1792	G
1	2	1793	G
1	2	1794	A
1	2	1796	C
36	1	6	A
36	1	14	U
36	1	26	A
36	1	39	A
36	1	40	A
36	1	43	A
36	1	49	A
36	1	55	G
36	1	57	A
36	1	59	G
36	1	60	A
36	1	65	A
36	1	66	A
36	1	76	G
36	1	77	A

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Mol	Chain	Res	Type
36	1	83	U
36	1	92	G
36	1	93	C
36	1	97	U
36	1	99	A
36	1	102	C
36	1	109	A
36	1	110	G
36	1	111	C
36	1	114	A
36	1	115	A
36	1	116	A
36	1	121	A
36	1	122	A
36	1	133	U
36	1	135	C
36	1	136	G
36	1	139	G
36	1	156	G
36	1	157	A
36	1	160	G
36	1	165	A
36	1	166	C
36	1	169	U
36	1	170	G
36	1	177	U
36	1	187	A
36	1	188	U
36	1	190	U
36	1	191	U
36	1	201	A
36	1	202	G
36	1	210	U
36	1	213	A
36	1	218	G
36	1	219	A
36	1	220	G
36	1	224	C
36	1	227	G
36	1	239	G
36	1	240	U
36	1	241	G

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Mol	Chain	Res	Type
36	1	243	G
36	1	245	U
36	1	246	U
36	1	247	C
36	1	249	U
36	1	250	U
36	1	251	G
36	1	252	U
36	1	269	G
36	1	283	G
36	1	286	U
36	1	295	A
36	1	298	U
36	1	315	C
36	1	316	U
36	1	323	A
36	1	326	U
36	1	329	U
36	1	339	C
36	1	350	C
36	1	352	A
36	1	374	A
36	1	375	A
36	1	376	G
36	1	398	A
36	1	399	A
36	1	401	U
36	1	402	A
36	1	403	C
36	1	404	G
36	1	421	G
36	1	422	A
36	1	424	G
36	1	438	A
36	1	439	C
36	1	440	A
36	1	495	G
36	1	497	C
36	1	503	C
36	1	510	G
36	1	518	G
36	1	520	U

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Mol	Chain	Res	Type
36	1	521	A
36	1	529	A
36	1	531	G
36	1	535	G
36	1	544	C
36	1	546	C
36	1	547	G
36	1	548	G
36	1	550	A
36	1	551	A
36	1	552	G
36	1	553	U
36	1	555	U
36	1	557	A
36	1	558	U
36	1	559	A
36	1	572	A
36	1	578	A
36	1	579	G
36	1	581	U
36	1	592	A
36	1	594	U
36	1	596	C
36	1	599	C
36	1	600	G
36	1	601	U
36	1	604	G
36	1	609	G
36	1	611	A
36	1	618	C
36	1	619	A
36	1	620	U
36	1	621	A
36	1	624	G
36	1	636	C
36	1	637	C
36	1	638	C
36	1	642	U
36	1	644	G
36	1	648	C
36	1	649	A
36	1	651	G

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Mol	Chain	Res	Type
36	1	657	A
36	1	660	A
36	1	661	G
36	1	662	U
36	1	675	C
36	1	677	A
36	1	681	U
36	1	683	U
36	1	685	G
36	1	691	A
36	1	705	A
36	1	708	G
36	1	709	A
36	1	711	A
36	1	712	G
36	1	715	A
36	1	716	A
36	1	717	C
36	1	719	U
36	1	762	U
36	1	764	U
36	1	766	U
36	1	767	U
36	1	776	U
36	1	777	U
36	1	780	A
36	1	781	G
36	1	785	G
36	1	786	A
36	1	790	U
36	1	792	G
36	1	798	G
36	1	803	C
36	1	806	A
36	1	817	A
36	1	825	U
36	1	830	A
36	1	837	A
36	1	845	G
36	1	861	C
36	1	867	G
36	1	869	G

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Mol	Chain	Res	Type
36	1	871	U
36	1	872	U
36	1	874	U
36	1	875	G
36	1	879	U
36	1	880	G
36	1	890	C
36	1	896	A
36	1	907	G
36	1	908	G
36	1	910	G
36	1	914	A
36	1	916	G
36	1	917	A
36	1	919	U
36	1	923	C
36	1	924	G
36	1	930	U
36	1	937	G
36	1	944	C
36	1	953	G
36	1	959	C
36	1	960	U
36	1	963	G
36	1	969	C
36	1	979	U
36	1	981	U
36	1	982	C
36	1	984	G
36	1	991	G
36	1	993	G
36	1	994	G
36	1	995	U
36	1	1000	C
36	1	1001	G
36	1	1002	A
36	1	1003	A
36	1	1004	U
36	1	1010	G
36	1	1017	C
36	1	1018	G
36	1	1019	G

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Mol	Chain	Res	Type
36	1	1020	G
36	1	1021	G
36	1	1024	G
36	1	1025	A
36	1	1029	G
36	1	1034	U
36	1	1036	A
36	1	1037	C
36	1	1041	U
36	1	1042	U
36	1	1047	A
36	1	1049	C
36	1	1052	U
36	1	1053	A
36	1	1063	G
36	1	1064	A
36	1	1065	A
36	1	1068	C
36	1	1072	G
36	1	1081	U
36	1	1082	U
36	1	1087	G
36	1	1093	A
36	1	1094	U
36	1	1095	U
36	1	1096	U
36	1	1097	G
36	1	1098	A
36	1	1103	A
36	1	1104	G
36	1	1117	G
36	1	1129	A
36	1	1131	G
36	1	1153	A
36	1	1159	A
36	1	1160	C
36	1	1178	G
36	1	1180	A
36	1	1181	U
36	1	1182	A
36	1	1191	U
36	1	1192	C

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Mol	Chain	Res	Type
36	1	1200	A
36	1	1201	C
36	1	1202	A
36	1	1209	G
36	1	1212	A
36	1	1213	G
36	1	1217	A
36	1	1222	G
36	1	1225	A
36	1	1226	G
36	1	1227	C
36	1	1232	C
36	1	1233	G
36	1	1236	G
36	1	1237	G
36	1	1241	U
36	1	1245	A
36	1	1246	G
36	1	1248	C
36	1	1249	G
36	1	1251	A
36	1	1253	U
36	1	1254	C
36	1	1257	C
36	1	1258	U
36	1	1262	G
36	1	1263	A
36	1	1264	G
36	1	1265	U
36	1	1267	U
36	1	1269	U
36	1	1270	A
36	1	1271	A
36	1	1272	C
36	1	1274	A
36	1	1276	U
36	1	1278	A
36	1	1279	C
36	1	1285	G
36	1	1287	A
36	1	1292	C
36	1	1305	U

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Mol	Chain	Res	Type
36	1	1307	G
36	1	1308	A
36	1	1309	U
36	1	1312	C
36	1	1313	G
36	1	1318	A
36	1	1325	U
36	1	1330	A
36	1	1331	U
36	1	1339	C
36	1	1348	U
36	1	1349	G
36	1	1350	A
36	1	1351	U
36	1	1352	A
36	1	1353	U
36	1	1355	A
36	1	1356	U
36	1	1357	G
36	1	1373	A
36	1	1386	A
36	1	1392	G
36	1	1397	C
36	1	1399	A
36	1	1400	G
36	1	1405	U
36	1	1406	A
36	1	1417	G
36	1	1418	A
36	1	1419	A
36	1	1429	G
36	1	1431	G
36	1	1433	A
36	1	1434	G
36	1	1435	A
36	1	1437	C
36	1	1446	A
36	1	1450	G
36	1	1468	A
36	1	1471	U
36	1	1472	U
36	1	1481	A

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Mol	Chain	Res	Type
36	1	1482	A
36	1	1485	G
36	1	1508	C
36	1	1511	U
36	1	1527	C
36	1	1533	U
36	1	1535	A
36	1	1548	C
36	1	1555	U
36	1	1556	C
36	1	1557	A
36	1	1560	G
36	1	1562	C
36	1	1563	C
36	1	1564	U
36	1	1566	A
36	1	1567	U
36	1	1568	U
36	1	1569	U
36	1	1570	U
36	1	1572	U
36	1	1578	C
36	1	1579	C
36	1	1580	A
36	1	1581	C
36	1	1582	C
36	1	1583	A
36	1	1587	A
36	1	1589	A
36	1	1593	A
36	1	1594	A
36	1	1605	A
36	1	1607	U
36	1	1608	C
36	1	1620	U
36	1	1621	A
36	1	1624	G
36	1	1629	U
36	1	1633	C
36	1	1639	C
36	1	1641	U
36	1	1643	A

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Mol	Chain	Res	Type
36	1	1645	U
36	1	1656	A
36	1	1657	C
36	1	1668	G
36	1	1679	A
36	1	1683	A
36	1	1687	U
36	1	1716	U
36	1	1717	U
36	1	1724	U
36	1	1725	C
36	1	1731	A
36	1	1736	G
36	1	1741	A
36	1	1742	U
36	1	1750	A
36	1	1751	G
36	1	1756	C
36	1	1761	C
36	1	1764	U
36	1	1765	U
36	1	1766	G
36	1	1767	C
36	1	1770	G
36	1	1773	C
36	1	1775	G
36	1	1780	G
36	1	1782	U
36	1	1793	C
36	1	1797	A
36	1	1810	A
36	1	1814	A
36	1	1815	U
36	1	1816	A
36	1	1817	G
36	1	1819	U
36	1	1820	U
36	1	1821	U
36	1	1835	A
36	1	1839	A
36	1	1842	A
36	1	1845	G

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Mol	Chain	Res	Type
36	1	1846	C
36	1	1849	C
36	1	1858	A
36	1	1866	C
36	1	1871	U
36	1	1879	A
36	1	1880	U
36	1	1886	A
36	1	1901	A
36	1	1906	G
36	1	1908	A
36	1	1920	U
36	1	1951	C
36	1	1952	G
36	1	1953	G
36	1	1954	G
36	1	2101	C
36	1	2102	U
36	1	2106	A
36	1	2111	G
36	1	2112	U
36	1	2113	A
36	1	2114	C
36	1	2116	G
36	1	2121	G
36	1	2122	G
36	1	2127	U
36	1	2131	A
36	1	2134	G
36	1	2139	A
36	1	2140	U
36	1	2149	A
36	1	2156	C
36	1	2158	A
36	1	2169	G
36	1	2187	G
36	1	2198	A
36	1	2205	U
36	1	2208	A
36	1	2209	U
36	1	2210	G
36	1	2213	A

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Mol	Chain	Res	Type
36	1	2223	A
36	1	2225	U
36	1	2228	A
36	1	2229	A
36	1	2239	G
36	1	2244	A
36	1	2245	C
36	1	2246	G
36	1	2247	G
36	1	2249	G
36	1	2250	G
36	1	2252	A
36	1	2255	A
36	1	2256	A
36	1	2262	A
36	1	2266	U
36	1	2272	G
36	1	2273	G
36	1	2276	G
36	1	2281	A
36	1	2282	U
36	1	2283	G
36	1	2284	C
36	1	2288	G
36	1	2295	A
36	1	2303	A
36	1	2306	C
36	1	2307	G
36	1	2308	C
36	1	2309	A
36	1	2310	U
36	1	2313	A
36	1	2314	U
36	1	2315	G
36	1	2318	U
36	1	2319	U
36	1	2324	A
36	1	2334	U
36	1	2336	U
36	1	2350	C
36	1	2352	A
36	1	2361	A

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Mol	Chain	Res	Type
36	1	2372	A
36	1	2373	A
36	1	2374	C
36	1	2375	G
36	1	2382	G
36	1	2385	G
36	1	2388	U
36	1	2393	G
36	1	2394	G
36	1	2397	A
36	1	2402	A
36	1	2403	G
36	1	2411	U
36	1	2418	G
36	1	2419	A
36	1	2427	U
36	1	2444	C
36	1	2445	A
36	1	2502	A
36	1	2503	G
36	1	2504	U
36	1	2514	U
36	1	2515	A
36	1	2522	G
36	1	2523	A
36	1	2530	G
36	1	2531	C
36	1	2532	U
36	1	2533	G
36	1	2537	U
36	1	2538	U
36	1	2539	C
36	1	2540	A
36	1	2541	U
36	1	2542	U
36	1	2543	U
36	1	2544	U
36	1	2545	C
36	1	2548	C
36	1	2549	G
36	1	2552	C
36	1	2554	A

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Mol	Chain	Res	Type
36	1	2555	G
36	1	2561	A
36	1	2568	C
36	1	2569	A
36	1	2570	U
36	1	2571	U
36	1	2572	C
36	1	2573	G
36	1	2581	U
36	1	2582	C
36	1	2585	G
36	1	2593	A
36	1	2606	G
36	1	2607	G
36	1	2614	G
36	1	2626	A
36	1	2629	U
36	1	2637	A
36	1	2638	C
36	1	2652	U
36	1	2656	A
36	1	2657	A
36	1	2660	G
36	1	2672	G
36	1	2674	A
36	1	2675	C
36	1	2677	G
36	1	2681	U
36	1	2689	A
36	1	2690	G
36	1	2691	A
36	1	2694	A
36	1	2696	A
36	1	2705	A
36	1	2714	G
36	1	2725	U
36	1	2727	A
36	1	2728	G
36	1	2729	U
36	1	2739	A
36	1	2740	A
36	1	2752	U

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Mol	Chain	Res	Type
36	1	2753	G
36	1	2762	A
36	1	2772	C
36	1	2773	C
36	1	2777	G
36	1	2778	G
36	1	2779	A
36	1	2796	G
36	1	2797	C
36	1	2800	G
36	1	2801	A
36	1	2802	A
36	1	2810	C
36	1	2814	G
36	1	2816	G
36	1	2817	A
36	1	2818	U
36	1	2821	C
36	1	2822	U
36	1	2827	U
36	1	2829	U
36	1	2839	G
36	1	2842	U
36	1	2844	C
36	1	2845	A
36	1	2847	A
36	1	2849	C
36	1	2855	U
36	1	2860	U
36	1	2862	U
36	1	2863	G
36	1	2865	U
36	1	2867	C
36	1	2871	G
36	1	2872	A
36	1	2884	C
36	1	2887	A
36	1	2889	C
36	1	2893	C
36	1	2897	A
36	1	2898	G
36	1	2899	C

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Mol	Chain	Res	Type
36	1	2911	A
36	1	2917	G
36	1	2923	U
36	1	2933	A
36	1	2935	U
36	1	2936	A
36	1	2942	C
36	1	2945	G
36	1	2947	G
36	1	2954	U
36	1	2971	A
36	1	2973	G
36	1	2974	U
36	1	2977	G
36	1	2979	U
36	1	2982	A
36	1	2983	C
36	1	2990	G
36	1	2996	U
36	1	2997	G
36	1	3012	A
36	1	3021	A
36	1	3030	G
36	1	3041	U
36	1	3049	A
36	1	3056	U
36	1	3057	U
36	1	3058	U
36	1	3059	G
36	1	3078	U
36	1	3079	U
36	1	3080	G
36	1	3085	G
36	1	3086	A
36	1	3087	A
36	1	3092	C
36	1	3093	C
36	1	3113	A
36	1	3120	C
36	1	3122	A
36	1	3130	A
36	1	3131	U

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Mol	Chain	Res	Type
36	1	3139	A
36	1	3142	A
36	1	3143	C
36	1	3151	U
36	1	3154	C
36	1	3155	U
36	1	3156	U
36	1	3157	U
36	1	3158	G
36	1	3164	C
36	1	3165	A
36	1	3167	A
36	1	3168	A
36	1	3170	A
36	1	3171	U
36	1	3173	G
36	1	3174	A
36	1	3176	G
36	1	3179	U
36	1	3181	C
36	1	3187	A
36	1	3195	U
36	1	3196	U
36	1	3198	U
36	1	3199	G
36	1	3204	C
36	1	3206	C
36	1	3207	U
36	1	3217	C
36	1	3218	A
36	1	3219	G
36	1	3228	C
36	1	3229	G
36	1	3243	A
36	1	3244	A
36	1	3245	A
36	1	3246	G
36	1	3247	G
36	1	3249	C
36	1	3253	G
36	1	3259	U
36	1	3262	U

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Mol	Chain	Res	Type
36	1	3263	G
36	1	3269	U
36	1	3270	U
36	1	3273	A
36	1	3276	G
36	1	3279	A
36	1	3281	U
36	1	3286	G
36	1	3287	U
36	1	3288	G
36	1	3289	G
36	1	3290	G
36	1	3293	U
36	1	3294	A
36	1	3295	A
36	1	3304	U
36	1	3309	G
36	1	3313	U
36	1	3316	A
36	1	3317	U
36	1	3318	G
36	1	3319	U
36	1	3320	A
36	1	3335	A
36	1	3341	U
36	1	3342	A
36	1	3345	G
36	1	3347	A
36	1	3350	C
36	1	3351	U
36	1	3352	U
36	1	3353	G
36	1	3354	U
36	1	3355	U
36	1	3356	G
36	1	3363	U
36	1	3369	G
36	1	3372	A
36	1	3375	A
36	1	3376	A
36	1	3378	C
36	1	3382	U

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Mol	Chain	Res	Type
36	1	3389	U
36	1	3390	G
36	1	3396	U
37	3	7	G
37	3	9	C
37	3	21	G
37	3	22	A
37	3	25	G
37	3	27	A
37	3	44	C
37	3	45	A
37	3	51	A
37	3	54	U
37	3	60	G
37	3	65	G
37	3	74	C
37	3	76	A
37	3	92	A
37	3	93	C
37	3	102	A
37	3	106	U
37	3	112	G
37	3	121	U
38	4	2	A
38	4	17	A
38	4	20	U
38	4	23	U
38	4	26	U
38	4	34	U
38	4	35	C
38	4	43	A
38	4	50	C
38	4	51	G
38	4	52	A
38	4	54	A
38	4	58	G
38	4	59	A
38	4	62	C
38	4	63	G
38	4	80	A
38	4	81	U
38	4	82	U

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Mol	Chain	Res	Type
38	4	83	C
38	4	84	C
38	4	85	G
38	4	86	U
38	4	87	G
38	4	90	U
38	4	95	G
38	4	96	A
38	4	97	A
38	4	104	A
38	4	106	C
38	4	111	A
38	4	113	U
38	4	114	G
38	4	125	U
38	4	126	A
38	4	128	U
38	4	134	G
38	4	138	A
38	4	152	G
38	4	155	A
38	4	157	U
38	4	158	U
1	6	2	A
1	6	4	C
1	6	6	G
1	6	11	A
1	6	17	C
1	6	25	C
1	6	26	A
1	6	27	U
1	6	34	G
1	6	38	C
1	6	45	U
1	6	46	A
1	6	47	A
1	6	50	C
1	6	57	G
1	6	61	A
1	6	66	U
1	6	67	A
1	6	68	A

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Mol	Chain	Res	Type
1	6	69	G
1	6	73	U
1	6	75	U
1	6	76	A
1	6	77	U
1	6	78	A
1	6	101	U
1	6	104	A
1	6	105	A
1	6	111	U
1	6	114	C
1	6	132	U
1	6	137	U
1	6	138	A
1	6	140	A
1	6	141	U
1	6	142	G
1	6	143	G
1	6	144	U
1	6	145	A
1	6	146	U
1	6	153	G
1	6	158	U
1	6	159	U
1	6	171	A
1	6	176	C
1	6	178	U
1	6	185	U
1	6	188	A
1	6	190	C
1	6	191	C
1	6	192	U
1	6	193	U
1	6	194	U
1	6	195	G
1	6	196	G
1	6	199	G
1	6	200	A
1	6	214	G
1	6	215	A
1	6	216	U
1	6	217	A

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Mol	Chain	Res	Type
1	6	218	A
1	6	219	A
1	6	220	A
1	6	222	A
1	6	226	A
1	6	227	U
1	6	228	G
1	6	230	C
1	6	232	U
1	6	233	C
1	6	235	G
1	6	240	U
1	6	241	U
1	6	249	U
1	6	250	C
1	6	260	U
1	6	261	U
1	6	265	A
1	6	271	A
1	6	272	U
1	6	273	G
1	6	275	C
1	6	277	U
1	6	278	U
1	6	280	U
1	6	281	G
1	6	287	G
1	6	292	U
1	6	299	A
1	6	302	U
1	6	308	C
1	6	314	C
1	6	316	A
1	6	320	U
1	6	321	C
1	6	322	G
1	6	337	G
1	6	338	C
1	6	341	A
1	6	352	A
1	6	359	A
1	6	360	A

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Mol	Chain	Res	Type
1	6	361	C
1	6	370	A
1	6	381	C
1	6	386	G
1	6	390	G
1	6	400	A
1	6	401	A
1	6	402	C
1	6	404	G
1	6	416	A
1	6	418	G
1	6	419	G
1	6	424	C
1	6	425	A
1	6	426	G
1	6	429	G
1	6	434	G
1	6	437	A
1	6	439	U
1	6	444	C
1	6	448	C
1	6	470	A
1	6	475	A
1	6	477	A
1	6	484	C
1	6	485	A
1	6	486	G
1	6	487	G
1	6	488	G
1	6	489	C
1	6	490	C
1	6	492	A
1	6	493	U
1	6	494	U
1	6	496	G
1	6	499	U
1	6	500	C
1	6	501	U
1	6	502	U
1	6	504	U
1	6	505	A
1	6	506	A

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Mol	Chain	Res	Type
1	6	507	U
1	6	508	U
1	6	510	G
1	6	511	A
1	6	512	A
1	6	513	U
1	6	515	A
1	6	518	A
1	6	519	C
1	6	527	A
1	6	536	C
1	6	539	G
1	6	540	G
1	6	541	A
1	6	542	A
1	6	543	C
1	6	544	A
1	6	548	G
1	6	554	C
1	6	557	G
1	6	558	U
1	6	559	C
1	6	565	C
1	6	570	A
1	6	574	G
1	6	575	C
1	6	578	U
1	6	579	A
1	6	580	A
1	6	582	U
1	6	584	C
1	6	588	U
1	6	594	A
1	6	595	G
1	6	611	U
1	6	619	A
1	6	620	A
1	6	622	A
1	6	623	A
1	6	634	G
1	6	635	A
1	6	637	C

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Mol	Chain	Res	Type
1	6	639	U
1	6	650	U
1	6	651	G
1	6	652	G
1	6	653	C
1	6	655	G
1	6	661	A
1	6	662	U
1	6	665	U
1	6	667	U
1	6	668	C
1	6	669	G
1	6	670	U
1	6	676	G
1	6	678	A
1	6	679	U
1	6	680	U
1	6	681	U
1	6	682	C
1	6	683	C
1	6	684	A
1	6	691	C
1	6	696	C
1	6	697	C
1	6	698	U
1	6	709	C
1	6	710	U
1	6	711	U
1	6	714	G
1	6	718	U
1	6	719	U
1	6	720	G
1	6	721	U
1	6	722	G
1	6	723	G
1	6	726	C
1	6	730	G
1	6	741	C
1	6	742	U
1	6	744	U
1	6	751	G
1	6	752	A

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Mol	Chain	Res	Type
1	6	754	A
1	6	755	A
1	6	756	A
1	6	765	G
1	6	774	A
1	6	775	G
1	6	780	A
1	6	781	U
1	6	782	U
1	6	783	G
1	6	789	A
1	6	792	U
1	6	793	A
1	6	794	U
1	6	811	A
1	6	812	A
1	6	814	A
1	6	815	G
1	6	816	G
1	6	820	U
1	6	821	U
1	6	823	G
1	6	824	G
1	6	825	U
1	6	826	U
1	6	829	A
1	6	830	U
1	6	831	U
1	6	832	U
1	6	834	G
1	6	835	U
1	6	856	A
1	6	860	U
1	6	862	A
1	6	863	A
1	6	868	G
1	6	873	U
1	6	875	G
1	6	898	A
1	6	906	A
1	6	912	U
1	6	913	G

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Mol	Chain	Res	Type
1	6	914	G
1	6	915	A
1	6	916	U
1	6	925	G
1	6	931	C
1	6	933	A
1	6	935	U
1	6	940	A
1	6	942	G
1	6	944	A
1	6	945	U
1	6	954	G
1	6	959	U
1	6	960	U
1	6	966	A
1	6	969	C
1	6	971	A
1	6	983	A
1	6	992	A
1	6	993	A
1	6	996	U
1	6	1003	A
1	6	1004	U
1	6	1005	A
1	6	1021	C
1	6	1026	A
1	6	1028	C
1	6	1029	U
1	6	1039	A
1	6	1040	G
1	6	1041	G
1	6	1042	G
1	6	1043	A
1	6	1052	U
1	6	1053	G
1	6	1057	U
1	6	1058	U
1	6	1059	U
1	6	1060	U
1	6	1071	U
1	6	1074	G
1	6	1080	U

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Mol	Chain	Res	Type
1	6	1081	A
1	6	1082	C
1	6	1092	A
1	6	1096	C
1	6	1097	U
1	6	1098	U
1	6	1100	G
1	6	1109	G
1	6	1121	C
1	6	1138	A
1	6	1151	A
1	6	1155	G
1	6	1158	C
1	6	1159	C
1	6	1160	A
1	6	1161	C
1	6	1167	G
1	6	1185	U
1	6	1194	A
1	6	1196	A
1	6	1199	G
1	6	1200	G
1	6	1202	A
1	6	1217	A
1	6	1218	G
1	6	1219	A
1	6	1220	C
1	6	1221	A
1	6	1226	A
1	6	1228	G
1	6	1229	G
1	6	1230	A
1	6	1236	A
1	6	1239	U
1	6	1241	G
1	6	1242	A
1	6	1243	G
1	6	1244	A
1	6	1245	G
1	6	1248	C
1	6	1252	C
1	6	1255	G

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Mol	Chain	Res	Type
1	6	1256	A
1	6	1257	U
1	6	1258	U
1	6	1259	U
1	6	1275	A
1	6	1285	U
1	6	1286	U
1	6	1288	G
1	6	1293	U
1	6	1307	U
1	6	1312	A
1	6	1314	U
1	6	1315	U
1	6	1316	G
1	6	1319	A
1	6	1320	U
1	6	1321	A
1	6	1343	U
1	6	1344	A
1	6	1345	A
1	6	1346	A
1	6	1354	G
1	6	1355	C
1	6	1360	A
1	6	1361	U
1	6	1362	U
1	6	1363	U
1	6	1364	G
1	6	1370	U
1	6	1371	A
1	6	1373	C
1	6	1382	A
1	6	1388	A
1	6	1390	U
1	6	1398	U
1	6	1399	C
1	6	1400	A
1	6	1402	G
1	6	1412	G
1	6	1413	U
1	6	1415	U
1	6	1422	A

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Mol	Chain	Res	Type
1	6	1427	A
1	6	1428	G
1	6	1431	C
1	6	1433	G
1	6	1445	G
1	6	1446	A
1	6	1447	C
1	6	1448	G
1	6	1459	C
1	6	1461	C
1	6	1471	A
1	6	1481	C
1	6	1482	C
1	6	1483	A
1	6	1486	G
1	6	1490	C
1	6	1491	U
1	6	1492	A
1	6	1493	A
1	6	1494	C
1	6	1496	U
1	6	1506	G
1	6	1514	U
1	6	1515	A
1	6	1516	A
1	6	1517	U
1	6	1519	U
1	6	1520	U
1	6	1521	G
1	6	1523	G
1	6	1524	A
1	6	1526	A
1	6	1531	G
1	6	1535	U
1	6	1536	G
1	6	1537	C
1	6	1538	U
1	6	1539	G
1	6	1540	G
1	6	1550	A
1	6	1554	U
1	6	1557	U

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Mol	Chain	Res	Type
1	6	1559	A
1	6	1560	U
1	6	1569	A
1	6	1573	A
1	6	1574	G
1	6	1575	G
1	6	1584	G
1	6	1590	G
1	6	1596	C
1	6	1600	A
1	6	1601	G
1	6	1619	C
1	6	1620	C
1	6	1621	U
1	6	1634	C
1	6	1637	C
1	6	1638	G
1	6	1641	C
1	6	1657	U
1	6	1658	G
1	6	1682	U
1	6	1696	G
1	6	1697	G
1	6	1698	G
1	6	1699	G
1	6	1700	C
1	6	1701	A
1	6	1702	A
1	6	1703	C
1	6	1712	A
1	6	1715	G
1	6	1716	C
1	6	1717	G
1	6	1727	G
1	6	1731	A
1	6	1735	U
1	6	1736	G
1	6	1742	U
1	6	1747	G
1	6	1760	G
1	6	1766	A
1	6	1767	G

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Mol	Chain	Res	Type
1	6	1769	U
1	6	1770	U
1	6	1780	G
1	6	1782	A
1	6	1783	C
1	6	1788	G
1	6	1792	G
1	6	1793	G
1	6	1794	A
1	6	1796	C
1	6	1798	U
1	6	1799	U
1	6	1800	A
36	5	15	C
36	5	16	A
36	5	26	A
36	5	40	A
36	5	43	A
36	5	44	U
36	5	49	A
36	5	58	G
36	5	59	G
36	5	60	A
36	5	65	A
36	5	66	A
36	5	73	C
36	5	76	G
36	5	85	A
36	5	92	G
36	5	94	G
36	5	96	G
36	5	102	C
36	5	108	A
36	5	109	A
36	5	110	G
36	5	111	C
36	5	116	A
36	5	120	G
36	5	121	A
36	5	122	A
36	5	130	A
36	5	133	U

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Mol	Chain	Res	Type
36	5	134	U
36	5	135	C
36	5	136	G
36	5	152	U
36	5	156	G
36	5	157	A
36	5	165	A
36	5	169	U
36	5	170	G
36	5	171	G
36	5	173	G
36	5	174	C
36	5	178	U
36	5	180	C
36	5	181	U
36	5	187	A
36	5	190	U
36	5	191	U
36	5	200	C
36	5	203	G
36	5	204	A
36	5	206	G
36	5	210	U
36	5	211	A
36	5	213	A
36	5	218	G
36	5	219	A
36	5	220	G
36	5	221	A
36	5	222	A
36	5	227	G
36	5	231	G
36	5	234	G
36	5	236	G
36	5	239	G
36	5	240	U
36	5	244	G
36	5	247	C
36	5	248	U
36	5	249	U
36	5	250	U
36	5	251	G

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Mol	Chain	Res	Type
36	5	252	U
36	5	253	A
36	5	254	A
36	5	259	C
36	5	269	G
36	5	284	A
36	5	286	U
36	5	294	U
36	5	295	A
36	5	311	C
36	5	321	C
36	5	322	U
36	5	323	A
36	5	328	U
36	5	329	U
36	5	333	G
36	5	334	A
36	5	338	A
36	5	339	C
36	5	349	A
36	5	350	C
36	5	351	A
36	5	361	A
36	5	366	A
36	5	376	G
36	5	377	A
36	5	384	A
36	5	395	A
36	5	398	A
36	5	399	A
36	5	401	U
36	5	402	A
36	5	403	C
36	5	404	G
36	5	417	A
36	5	419	G
36	5	421	G
36	5	422	A
36	5	426	G
36	5	436	A
36	5	437	G
36	5	439	C

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Mol	Chain	Res	Type
36	5	441	U
36	5	442	G
36	5	492	U
36	5	507	U
36	5	520	U
36	5	521	A
36	5	525	C
36	5	531	G
36	5	535	G
36	5	538	G
36	5	539	C
36	5	542	G
36	5	546	C
36	5	547	G
36	5	548	G
36	5	553	U
36	5	555	U
36	5	557	A
36	5	558	U
36	5	559	A
36	5	569	A
36	5	578	A
36	5	579	G
36	5	582	G
36	5	592	A
36	5	604	G
36	5	609	G
36	5	611	A
36	5	612	U
36	5	619	A
36	5	621	A
36	5	636	C
36	5	648	C
36	5	649	A
36	5	660	A
36	5	662	U
36	5	677	A
36	5	681	U
36	5	692	A
36	5	699	A
36	5	705	A
36	5	712	G

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Mol	Chain	Res	Type
36	5	715	A
36	5	716	A
36	5	719	U
36	5	724	U
36	5	725	G
36	5	726	G
36	5	727	G
36	5	744	A
36	5	762	U
36	5	765	C
36	5	766	U
36	5	767	U
36	5	768	C
36	5	776	U
36	5	777	U
36	5	780	A
36	5	781	G
36	5	784	A
36	5	785	G
36	5	786	A
36	5	805	G
36	5	806	A
36	5	817	A
36	5	821	U
36	5	829	U
36	5	830	A
36	5	831	G
36	5	837	A
36	5	842	G
36	5	857	G
36	5	861	C
36	5	869	G
36	5	874	U
36	5	876	A
36	5	879	U
36	5	882	A
36	5	890	C
36	5	891	G
36	5	896	A
36	5	897	U
36	5	898	U
36	5	907	G

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Mol	Chain	Res	Type
36	5	908	G
36	5	909	G
36	5	910	G
36	5	913	A
36	5	914	A
36	5	916	G
36	5	917	A
36	5	921	A
36	5	923	C
36	5	924	G
36	5	929	A
36	5	931	C
36	5	932	U
36	5	937	G
36	5	944	C
36	5	948	C
36	5	959	C
36	5	960	U
36	5	964	G
36	5	979	U
36	5	980	A
36	5	981	U
36	5	994	G
36	5	1001	G
36	5	1002	A
36	5	1006	A
36	5	1010	G
36	5	1014	U
36	5	1015	U
36	5	1017	C
36	5	1019	G
36	5	1021	G
36	5	1024	G
36	5	1025	A
36	5	1026	A
36	5	1028	U
36	5	1029	G
36	5	1032	C
36	5	1035	G
36	5	1047	A
36	5	1049	C
36	5	1057	A

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Mol	Chain	Res	Type
36	5	1059	G
36	5	1063	G
36	5	1064	A
36	5	1065	A
36	5	1072	G
36	5	1081	U
36	5	1082	U
36	5	1083	G
36	5	1093	A
36	5	1094	U
36	5	1095	U
36	5	1097	G
36	5	1098	A
36	5	1103	A
36	5	1104	G
36	5	1106	G
36	5	1112	A
36	5	1115	G
36	5	1117	G
36	5	1129	A
36	5	1131	G
36	5	1143	A
36	5	1152	G
36	5	1153	A
36	5	1155	C
36	5	1159	A
36	5	1161	G
36	5	1162	U
36	5	1169	A
36	5	1178	G
36	5	1179	A
36	5	1180	A
36	5	1181	U
36	5	1190	A
36	5	1191	U
36	5	1192	C
36	5	1193	A
36	5	1201	C
36	5	1206	G
36	5	1209	G
36	5	1220	U
36	5	1222	G

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Mol	Chain	Res	Type
36	5	1232	C
36	5	1235	U
36	5	1236	G
36	5	1237	G
36	5	1239	C
36	5	1241	U
36	5	1242	G
36	5	1244	A
36	5	1245	A
36	5	1246	G
36	5	1252	A
36	5	1258	U
36	5	1262	G
36	5	1263	A
36	5	1264	G
36	5	1266	G
36	5	1268	G
36	5	1269	U
36	5	1275	C
36	5	1281	G
36	5	1284	C
36	5	1285	G
36	5	1292	C
36	5	1305	U
36	5	1307	G
36	5	1308	A
36	5	1309	U
36	5	1314	C
36	5	1324	U
36	5	1330	A
36	5	1331	U
36	5	1332	A
36	5	1347	U
36	5	1348	U
36	5	1349	G
36	5	1351	U
36	5	1352	A
36	5	1353	U
36	5	1354	G
36	5	1355	A
36	5	1356	U
36	5	1357	G

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Mol	Chain	Res	Type
36	5	1365	G
36	5	1368	U
36	5	1370	G
36	5	1372	C
36	5	1377	G
36	5	1379	G
36	5	1380	G
36	5	1385	C
36	5	1386	A
36	5	1391	C
36	5	1392	G
36	5	1398	U
36	5	1399	A
36	5	1400	G
36	5	1418	A
36	5	1419	A
36	5	1421	G
36	5	1428	A
36	5	1431	G
36	5	1433	A
36	5	1434	G
36	5	1437	C
36	5	1440	G
36	5	1441	G
36	5	1450	G
36	5	1456	A
36	5	1465	A
36	5	1466	G
36	5	1481	A
36	5	1482	A
36	5	1483	G
36	5	1495	U
36	5	1502	C
36	5	1508	C
36	5	1511	U
36	5	1519	G
36	5	1533	U
36	5	1536	G
36	5	1549	U
36	5	1552	G
36	5	1554	U
36	5	1555	U

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Mol	Chain	Res	Type
36	5	1556	C
36	5	1557	A
36	5	1560	G
36	5	1561	G
36	5	1562	C
36	5	1564	U
36	5	1566	A
36	5	1567	U
36	5	1569	U
36	5	1570	U
36	5	1571	A
36	5	1572	U
36	5	1574	C
36	5	1575	A
36	5	1577	G
36	5	1578	C
36	5	1579	C
36	5	1581	C
36	5	1583	A
36	5	1587	A
36	5	1589	A
36	5	1598	G
36	5	1600	U
36	5	1605	A
36	5	1607	U
36	5	1608	C
36	5	1619	A
36	5	1620	U
36	5	1629	U
36	5	1639	C
36	5	1641	U
36	5	1642	A
36	5	1643	A
36	5	1644	C
36	5	1645	U
36	5	1666	G
36	5	1683	A
36	5	1684	U
36	5	1686	U
36	5	1687	U
36	5	1688	U
36	5	1713	G

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Mol	Chain	Res	Type
36	5	1714	A
36	5	1716	U
36	5	1717	U
36	5	1724	U
36	5	1729	A
36	5	1733	G
36	5	1736	G
36	5	1750	A
36	5	1751	G
36	5	1760	A
36	5	1761	C
36	5	1762	C
36	5	1764	U
36	5	1765	U
36	5	1766	G
36	5	1767	C
36	5	1769	G
36	5	1770	G
36	5	1775	G
36	5	1780	G
36	5	1793	C
36	5	1795	U
36	5	1797	A
36	5	1810	A
36	5	1814	A
36	5	1815	U
36	5	1816	A
36	5	1817	G
36	5	1818	U
36	5	1821	U
36	5	1839	A
36	5	1841	A
36	5	1842	A
36	5	1846	C
36	5	1849	C
36	5	1866	C
36	5	1871	U
36	5	1876	U
36	5	1878	G
36	5	1879	A
36	5	1880	U
36	5	1881	A

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Mol	Chain	Res	Type
36	5	1885	U
36	5	1886	A
36	5	1891	A
36	5	1893	A
36	5	1901	A
36	5	1903	U
36	5	1904	C
36	5	1906	G
36	5	1922	A
36	5	1944	U
36	5	1948	G
36	5	2100	A
36	5	2101	C
36	5	2102	U
36	5	2111	G
36	5	2112	U
36	5	2113	A
36	5	2114	C
36	5	2121	G
36	5	2122	G
36	5	2131	A
36	5	2154	U
36	5	2158	A
36	5	2169	G
36	5	2176	U
36	5	2187	G
36	5	2192	C
36	5	2198	A
36	5	2205	U
36	5	2209	U
36	5	2210	G
36	5	2223	A
36	5	2225	U
36	5	2228	A
36	5	2231	C
36	5	2234	G
36	5	2244	A
36	5	2250	G
36	5	2252	A
36	5	2253	G
36	5	2255	A
36	5	2256	A

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Mol	Chain	Res	Type
36	5	2264	U
36	5	2270	A
36	5	2272	G
36	5	2273	G
36	5	2276	G
36	5	2281	A
36	5	2282	U
36	5	2283	G
36	5	2287	C
36	5	2288	G
36	5	2291	A
36	5	2298	U
36	5	2307	G
36	5	2309	A
36	5	2310	U
36	5	2313	A
36	5	2315	G
36	5	2334	U
36	5	2335	G
36	5	2336	U
36	5	2362	C
36	5	2365	C
36	5	2372	A
36	5	2373	A
36	5	2374	C
36	5	2375	G
36	5	2379	U
36	5	2385	G
36	5	2388	U
36	5	2392	C
36	5	2393	G
36	5	2394	G
36	5	2397	A
36	5	2398	A
36	5	2401	A
36	5	2403	G
36	5	2404	A
36	5	2405	C
36	5	2410	U
36	5	2411	U
36	5	2412	G
36	5	2413	A

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Mol	Chain	Res	Type
36	5	2414	G
36	5	2418	G
36	5	2420	C
36	5	2435	G
36	5	2437	G
36	5	2438	A
36	5	2439	A
36	5	2441	A
36	5	2443	A
36	5	2444	C
36	5	2504	U
36	5	2505	U
36	5	2506	U
36	5	2507	C
36	5	2508	U
36	5	2509	U
36	5	2510	U
36	5	2511	A
36	5	2513	U
36	5	2514	U
36	5	2515	A
36	5	2523	A
36	5	2524	A
36	5	2525	G
36	5	2526	C
36	5	2530	G
36	5	2531	C
36	5	2535	A
36	5	2537	U
36	5	2538	U
36	5	2539	C
36	5	2540	A
36	5	2543	U
36	5	2545	C
36	5	2547	A
36	5	2549	G
36	5	2552	C
36	5	2554	A
36	5	2555	G
36	5	2557	A
36	5	2560	C
36	5	2562	A

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Mol	Chain	Res	Type
36	5	2566	C
36	5	2567	C
36	5	2568	C
36	5	2569	A
36	5	2570	U
36	5	2571	U
36	5	2572	C
36	5	2573	G
36	5	2574	G
36	5	2578	U
36	5	2584	G
36	5	2585	G
36	5	2586	G
36	5	2589	G
36	5	2593	A
36	5	2598	G
36	5	2600	C
36	5	2606	G
36	5	2607	G
36	5	2609	A
36	5	2614	G
36	5	2618	G
36	5	2629	U
36	5	2637	A
36	5	2638	C
36	5	2642	A
36	5	2652	U
36	5	2653	C
36	5	2656	A
36	5	2667	A
36	5	2670	G
36	5	2674	A
36	5	2675	C
36	5	2677	G
36	5	2681	U
36	5	2689	A
36	5	2690	G
36	5	2691	A
36	5	2694	A
36	5	2696	A
36	5	2705	A
36	5	2706	G

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Mol	Chain	Res	Type
36	5	2707	C
36	5	2714	G
36	5	2725	U
36	5	2727	A
36	5	2728	G
36	5	2729	U
36	5	2740	A
36	5	2752	U
36	5	2753	G
36	5	2755	C
36	5	2762	A
36	5	2772	C
36	5	2773	C
36	5	2777	G
36	5	2778	G
36	5	2779	A
36	5	2782	U
36	5	2783	U
36	5	2788	C
36	5	2796	G
36	5	2797	C
36	5	2800	G
36	5	2801	A
36	5	2802	A
36	5	2810	C
36	5	2814	G
36	5	2817	A
36	5	2818	U
36	5	2819	A
36	5	2837	A
36	5	2839	G
36	5	2842	U
36	5	2843	U
36	5	2844	C
36	5	2845	A
36	5	2847	A
36	5	2849	C
36	5	2853	A
36	5	2870	C
36	5	2871	G
36	5	2872	A
36	5	2873	U

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Mol	Chain	Res	Type
36	5	2875	U
36	5	2876	C
36	5	2880	U
36	5	2887	A
36	5	2896	A
36	5	2899	C
36	5	2900	A
36	5	2911	A
36	5	2921	U
36	5	2923	U
36	5	2935	U
36	5	2936	A
36	5	2941	A
36	5	2942	C
36	5	2943	G
36	5	2947	G
36	5	2954	U
36	5	2960	C
36	5	2964	G
36	5	2971	A
36	5	2972	G
36	5	2973	G
36	5	2974	U
36	5	2979	U
36	5	2980	U
36	5	2983	C
36	5	2988	C
36	5	2989	U
36	5	2990	G
36	5	2996	U
36	5	2997	G
36	5	3006	A
36	5	3012	A
36	5	3025	C
36	5	3030	G
36	5	3034	C
36	5	3049	A
36	5	3056	U
36	5	3057	U
36	5	3058	U
36	5	3059	G
36	5	3078	U

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Mol	Chain	Res	Type
36	5	3079	U
36	5	3081	C
36	5	3086	A
36	5	3092	C
36	5	3098	G
36	5	3104	U
36	5	3115	C
36	5	3119	U
36	5	3122	A
36	5	3123	A
36	5	3127	A
36	5	3130	A
36	5	3131	U
36	5	3142	A
36	5	3143	C
36	5	3148	U
36	5	3153	U
36	5	3155	U
36	5	3156	U
36	5	3157	U
36	5	3158	G
36	5	3159	C
36	5	3164	C
36	5	3165	A
36	5	3166	C
36	5	3171	U
36	5	3172	A
36	5	3173	G
36	5	3174	A
36	5	3176	G
36	5	3180	A
36	5	3181	C
36	5	3187	A
36	5	3195	U
36	5	3196	U
36	5	3206	C
36	5	3207	U
36	5	3217	C
36	5	3218	A
36	5	3219	G
36	5	3224	G
36	5	3227	A

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Mol	Chain	Res	Type
36	5	3228	C
36	5	3229	G
36	5	3234	A
36	5	3238	G
36	5	3239	G
36	5	3244	A
36	5	3245	A
36	5	3246	G
36	5	3247	G
36	5	3253	G
36	5	3259	U
36	5	3263	G
36	5	3268	A
36	5	3270	U
36	5	3273	A
36	5	3275	U
36	5	3276	G
36	5	3277	U
36	5	3282	U
36	5	3283	U
36	5	3284	G
36	5	3285	C
36	5	3286	G
36	5	3287	U
36	5	3288	G
36	5	3289	G
36	5	3290	G
36	5	3294	A
36	5	3303	G
36	5	3304	U
36	5	3313	U
36	5	3316	A
36	5	3317	U
36	5	3318	G
36	5	3319	U
36	5	3320	A
36	5	3324	C
36	5	3330	A
36	5	3331	U
36	5	3341	U
36	5	3342	A
36	5	3345	G

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Mol	Chain	Res	Type
36	5	3350	C
36	5	3351	U
36	5	3352	U
36	5	3354	U
36	5	3356	G
36	5	3358	U
36	5	3360	C
36	5	3368	U
36	5	3369	G
36	5	3376	A
36	5	3378	C
36	5	3382	U
36	5	3383	G
36	5	3389	U
36	5	3390	G
36	5	3396	U
37	7	2	G
37	7	7	G
37	7	9	C
37	7	22	A
37	7	29	C
37	7	38	U
37	7	39	C
37	7	41	G
37	7	49	G
37	7	51	A
37	7	53	U
37	7	54	U
37	7	55	A
37	7	57	G
37	7	60	G
37	7	65	G
37	7	73	C
37	7	74	C
37	7	76	A
37	7	84	A
37	7	92	A
37	7	93	C
37	7	102	A
37	7	104	A
37	7	110	G
37	7	112	G

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Mol	Chain	Res	Type
37	7	114	U
37	7	121	U
38	8	13	A
38	8	21	C
38	8	23	U
38	8	34	U
38	8	35	C
38	8	48	A
38	8	50	C
38	8	51	G
38	8	53	A
38	8	59	A
38	8	62	C
38	8	63	G
38	8	64	U
38	8	71	A
38	8	80	A
38	8	81	U
38	8	82	U
38	8	83	C
38	8	84	C
38	8	85	G
38	8	86	U
38	8	87	G
38	8	88	A
38	8	90	U
38	8	95	G
38	8	96	A
38	8	97	A
38	8	104	A
38	8	105	A
38	8	106	C
38	8	110	C
38	8	111	A
38	8	113	U
38	8	114	G
38	8	116	G
38	8	122	U
38	8	125	U
38	8	126	A
38	8	127	U
38	8	138	A

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Mol	Chain	Res	Type
38	8	148	G
38	8	152	G
38	8	157	U
38	8	158	U

There are no RNA pucker outliers to report.

5.4 Non-standard residues in protein, DNA, RNA chains [i](#)

There are no non-standard protein/DNA/RNA residues in this entry.

5.5 Carbohydrates [i](#)

There are no carbohydrates in this entry.

5.6 Ligand geometry [i](#)

Of 2557 ligands modelled in this entry, 1423 are monoatomic - leaving 1134 for Mogul analysis.

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	$\# Z > 2$
86	OHX	1	3857	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3858	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3859	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3860	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3861	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3862	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3863	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3864	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3865	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3866	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3867	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3868	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	1	3869	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3870	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3871	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3872	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3873	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3874	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3875	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3876	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3877	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3878	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3879	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3880	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3881	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3882	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3883	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3884	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3885	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3886	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3887	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3888	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3889	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3890	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3891	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3892	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3893	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3894	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3895	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3896	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3897	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3898	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3899	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3900	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3901	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3902	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3903	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3904	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3905	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3906	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3907	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3908	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3909	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3910	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3911	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	1	3912	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3913	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3914	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3915	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3916	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3917	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3918	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3919	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3920	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3921	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3922	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3923	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3924	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3925	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3926	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3927	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3928	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3929	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3930	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3931	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3932	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3933	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3934	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3935	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3936	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3937	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3938	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3939	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3940	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3941	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3942	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3943	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3944	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3945	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3946	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3947	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3948	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3949	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3950	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3951	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3952	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3953	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3954	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	1	3955	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3956	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3957	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3958	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3959	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3960	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3961	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3962	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3963	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3964	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3965	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3966	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3967	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3968	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3969	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3970	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3971	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3972	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3973	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3974	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3975	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3976	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3977	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3978	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3979	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3980	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3981	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3982	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3983	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3984	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3985	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3986	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3987	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3988	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3989	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3990	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3991	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3992	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3993	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3994	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3995	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3996	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3997	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	1	3998	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	3999	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4000	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4001	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4002	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4003	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4004	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4005	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4006	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4007	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4008	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4009	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4010	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4011	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4012	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4013	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4014	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4015	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4016	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4017	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4018	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4019	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4020	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4021	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4022	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4023	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4024	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4025	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4026	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4027	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4028	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4029	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4030	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4031	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4032	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4033	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4034	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4035	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4036	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4037	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4038	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4039	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4040	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	1	4041	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4042	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4043	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4044	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4045	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4046	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4047	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4048	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4049	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4050	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4051	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4052	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4053	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4054	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4055	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4056	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4057	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4058	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4059	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4060	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4061	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4062	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4063	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4064	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4065	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4066	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4067	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4068	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4069	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4070	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4071	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4072	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4073	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4074	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4075	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4076	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4077	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4078	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4079	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4080	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4081	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4082	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4083	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	1	4084	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4085	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4086	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4087	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4088	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4089	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4090	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4091	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4092	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4093	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4094	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4095	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4096	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4097	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4098	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4099	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4100	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4101	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4102	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4103	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4104	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4105	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4106	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4107	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4108	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4109	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4110	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4111	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4112	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4113	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4114	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4115	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4116	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4117	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4118	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4119	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4120	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4121	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4122	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4123	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4124	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4125	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4126	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	1	4127	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4128	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4129	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4130	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4131	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4132	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4133	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4134	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4135	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4136	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4137	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4138	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4139	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4140	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4141	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4142	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4143	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4144	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4145	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4146	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4147	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4148	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4149	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4150	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4151	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4152	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4153	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4154	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4155	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4156	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4157	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4158	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4159	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4160	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4161	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4162	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4163	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4164	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4165	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4166	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4167	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4168	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4169	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	1	4170	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4171	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4172	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4173	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4174	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4175	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4176	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4177	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4178	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4179	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4180	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4181	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4182	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4183	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4184	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4185	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4186	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4187	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4188	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4189	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4190	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4191	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4192	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4193	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4194	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4195	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4196	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4197	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4198	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4199	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4200	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4201	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4202	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4203	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4204	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4205	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4206	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4207	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4208	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4209	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	1	4210	-	0,6,6	0.00	-	0,15,15	0.00	-
88	BLS	1	4211	-	23,31,31	1.88	5 (21%)	19,43,43	1.98	6 (31%)
86	OHX	2	2023	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	2	2024	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2025	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2026	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2027	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2028	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2029	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2030	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2031	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2032	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2033	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2034	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2035	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2036	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2037	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2038	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2039	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2040	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2041	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2042	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2043	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2044	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2045	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2046	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2047	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2048	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2049	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2050	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2051	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2052	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2053	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2054	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2055	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2056	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2057	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2058	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2059	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2060	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2061	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2062	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2063	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2064	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2065	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2066	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	2	2067	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2068	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2069	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2070	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2071	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2072	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2073	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2074	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2075	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2076	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2077	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2078	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2079	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2080	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2081	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2082	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2083	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2084	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2085	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2086	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2087	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2088	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2089	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2090	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2091	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2092	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2093	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2094	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2095	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2096	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2097	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2098	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2099	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2100	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2101	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2102	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2103	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2104	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2105	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2106	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2107	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2108	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2109	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	2	2110	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2111	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2112	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2113	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2114	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2115	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2116	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2117	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2118	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2119	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2120	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2121	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2122	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2123	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2124	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2125	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2126	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2127	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2128	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2129	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2130	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2131	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2132	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2133	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2134	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2135	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2136	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2137	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2138	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2139	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2140	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2141	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2142	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2143	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2144	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2145	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2146	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2147	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2148	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2149	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2150	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2151	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2152	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	2	2153	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2154	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2155	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2156	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2157	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2158	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2159	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2160	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2161	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2162	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2163	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2164	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2165	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2166	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2167	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2168	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2169	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2170	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2171	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2172	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2173	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2174	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2175	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2176	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2177	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2178	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2179	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	2	2180	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	3	216	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	3	217	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	3	218	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	3	219	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	3	220	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	3	221	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	3	222	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	3	223	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	3	224	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	3	225	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	4	224	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	4	225	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	4	226	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	4	227	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	4	228	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	4	229	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	4	230	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	4	231	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	4	232	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	4	233	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	4	234	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	4	235	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	4	236	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	4	237	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	4	238	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3892	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3893	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3894	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3895	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3896	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3897	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3898	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3899	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3900	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3901	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3902	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3903	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3904	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3905	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3906	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3907	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3908	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3909	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3910	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3911	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3912	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3913	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3914	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3915	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3916	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3917	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3918	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3919	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3920	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3921	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3922	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3923	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3924	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	5	3925	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3926	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3927	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3928	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3929	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3930	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3931	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3932	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3933	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3934	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3935	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3936	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3937	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3938	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3939	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3940	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3941	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3942	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3943	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3944	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3945	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3946	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3947	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3948	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3949	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3950	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3951	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3952	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3953	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3954	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3955	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3956	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3957	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3958	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3959	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3960	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3961	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3962	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3963	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3964	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3965	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3966	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3967	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	5	3968	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3969	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3970	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3971	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3972	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3973	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3974	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3975	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3976	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3977	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3978	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3979	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3980	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3981	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3982	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3983	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3984	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3985	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3986	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3987	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3988	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3989	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3990	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3991	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3992	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3993	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3994	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3995	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3996	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3997	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3998	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	3999	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4000	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4001	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4002	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4003	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4004	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4005	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4006	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4007	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4008	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4009	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4010	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	5	4011	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4012	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4013	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4014	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4015	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4016	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4017	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4018	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4019	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4020	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4021	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4022	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4023	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4024	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4025	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4026	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4027	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4028	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4029	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4030	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4031	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4032	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4033	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4034	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4035	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4036	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4037	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4038	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4039	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4040	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4041	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4042	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4043	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4044	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4045	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4046	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4047	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4048	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4049	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4050	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4051	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4052	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4053	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	5	4054	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4055	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4056	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4057	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4058	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4059	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4060	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4061	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4062	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4063	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4064	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4065	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4066	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4067	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4068	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4069	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4070	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4071	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4072	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4073	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4074	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4075	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4076	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4077	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4078	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4079	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4080	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4081	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4082	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4083	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4084	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4085	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4086	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4087	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4088	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4089	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4090	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4091	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4092	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4093	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4094	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4095	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4096	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	5	4097	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4098	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4099	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4100	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4101	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4102	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4103	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4104	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4105	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4106	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4107	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4108	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4109	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4110	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4111	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4112	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4113	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4114	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4115	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4116	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4117	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4118	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4119	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4120	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4121	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4122	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4123	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4124	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4125	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4126	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4127	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4128	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4129	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4130	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4131	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4132	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4133	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4134	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4135	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4136	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4137	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4138	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4139	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	5	4140	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4141	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4142	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4143	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4144	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4145	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4146	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4147	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4148	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4149	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4150	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4151	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4152	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4153	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4154	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4155	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4156	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4157	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4158	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4159	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4160	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4161	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4162	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4163	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4164	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4165	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4166	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4167	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4168	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4169	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4170	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4171	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4172	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4173	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4174	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4175	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4176	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4177	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4178	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4179	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4180	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4181	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4182	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
86	OHX	5	4183	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4184	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4185	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4186	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4187	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4188	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4189	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4190	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4191	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4192	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4193	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4194	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4195	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4196	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4197	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4198	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4199	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4200	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4201	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4202	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4203	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4204	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4205	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4206	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4207	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4208	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4209	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4210	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4211	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4212	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4213	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4214	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4215	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4216	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4217	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4218	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4219	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4220	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4221	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4222	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4223	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4224	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4225	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	5	4226	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4227	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4228	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4229	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4230	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4231	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4232	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4233	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4234	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4235	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4236	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4237	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4238	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4239	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4240	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4241	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4242	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4243	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4244	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4245	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4246	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	5	4247	-	0,6,6	0.00	-	0,15,15	0.00	-
88	BLS	5	4248	-	23,31,31	1.44	2 (8%)	19,43,43	0.73	0
86	OHX	6	2046	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2047	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2048	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2049	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2050	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2051	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2052	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2053	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2054	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2055	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2056	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2057	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2058	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2059	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2060	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2061	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2062	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2063	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2064	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2065	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	6	2066	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2067	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2068	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2069	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2070	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2071	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2072	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2073	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2074	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2075	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2076	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2077	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2078	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2079	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2080	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2081	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2082	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2083	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2084	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2085	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2086	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2087	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2088	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2089	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2090	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2091	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2092	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2093	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2094	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2095	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2096	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2097	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2098	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2099	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2100	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2101	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2102	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2103	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2104	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2105	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2106	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2107	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2108	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	6	2109	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2110	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2111	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2112	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2113	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2114	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2115	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2116	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2117	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2118	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2119	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2120	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2121	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2122	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2123	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2124	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2125	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2126	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2127	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2128	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2129	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2130	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2131	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2132	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2133	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2134	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2135	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2136	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2137	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2138	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2139	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2140	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2141	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2142	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2143	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2144	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2145	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2146	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2147	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2148	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2149	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2150	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2151	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	6	2152	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2153	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2154	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2155	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2156	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2157	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2158	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2159	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2160	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2161	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2162	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2163	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2164	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2165	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2166	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2167	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2168	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2169	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2170	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2171	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2172	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2173	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2174	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2175	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2176	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2177	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2178	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2179	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2180	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2181	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2182	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2183	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2184	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2185	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2186	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2187	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2188	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2189	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2190	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2191	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2192	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2193	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2194	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	6	2195	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2196	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2197	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2198	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2199	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2200	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2201	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2202	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2203	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	6	2204	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	7	218	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	7	219	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	7	220	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	7	221	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	7	222	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	7	223	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	7	224	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	7	225	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	7	226	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	7	227	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	7	228	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	7	229	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	8	215	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	8	216	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	8	217	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	8	218	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	8	219	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	8	220	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	8	221	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	8	222	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	8	223	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	8	224	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	8	225	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	8	226	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	8	227	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	8	228	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	8	229	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	C3	201	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	C5	201	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	C8	201	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	D3	202	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	D9	102	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	L3	405	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	L3	406	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	L4	404	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	M0	304	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	M5	303	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	M7	206	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	M7	207	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	M9	202	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	N9	101	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	O1	202	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	O3	201	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	O7	104	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	O7	105	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	Q2	503	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	S8	302	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	SR	401	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	c3	201	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	c5	201	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	c8	202	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	d4	201	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	d9	102	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	l3	404	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	l3	405	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	l4	402	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	l4	403	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	l5	304	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	l5	305	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	l5	306	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	l9	202	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	m0	302	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	m0	303	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	m1	202	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	m4	202	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	m5	306	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	m6	203	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	m7	207	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	n1	201	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	n3	202	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	n9	103	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	o3	203	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	o7	103	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	q2	502	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	s1	302	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	s1	303	-	0,6,6	0.00	-	0,15,15	0.00	-

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
86	OHX	s4	301	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	s8	302	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	s9	201	-	0,6,6	0.00	-	0,15,15	0.00	-
86	OHX	sR	401	-	0,6,6	0.00	-	0,15,15	0.00	-

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
86	OHX	1	3857	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3858	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3859	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3860	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3861	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3862	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3863	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3864	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3865	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3866	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3867	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3868	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3869	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3870	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3871	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3872	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3873	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3874	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3875	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3876	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3877	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3878	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3879	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3880	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3881	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3882	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3883	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3884	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3885	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3886	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3887	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3888	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	1	3889	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3890	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3891	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3892	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3893	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3894	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3895	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3896	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3897	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3898	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3899	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3900	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3901	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3902	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3903	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3904	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3905	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3906	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3907	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3908	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3909	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3910	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3911	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3912	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3913	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3914	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3915	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3916	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3917	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3918	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3919	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3920	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3921	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3922	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3923	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3924	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3925	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3926	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3927	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3928	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3929	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3930	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	1	3931	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3932	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3933	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3934	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3935	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3936	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3937	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3938	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3939	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3940	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3941	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3942	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3943	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3944	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3945	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3946	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3947	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3948	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3949	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3950	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3951	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3952	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3953	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3954	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3955	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3956	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3957	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3958	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3959	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3960	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3961	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3962	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3963	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3964	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3965	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3966	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3967	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3968	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3969	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3970	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3971	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3972	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	1	3973	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3974	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3975	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3976	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3977	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3978	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3979	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3980	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3981	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3982	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3983	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3984	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3985	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3986	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3987	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3988	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3989	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3990	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3991	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3992	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3993	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3994	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3995	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3996	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3997	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3998	-	-	0/0/0/0	0/0/0/0
86	OHX	1	3999	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4000	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4001	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4002	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4003	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4004	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4005	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4006	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4007	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4008	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4009	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4010	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4011	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4012	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4013	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4014	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	1	4015	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4016	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4017	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4018	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4019	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4020	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4021	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4022	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4023	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4024	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4025	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4026	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4027	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4028	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4029	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4030	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4031	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4032	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4033	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4034	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4035	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4036	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4037	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4038	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4039	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4040	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4041	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4042	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4043	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4044	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4045	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4046	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4047	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4048	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4049	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4050	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4051	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4052	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4053	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4054	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4055	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4056	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	1	4057	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4058	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4059	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4060	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4061	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4062	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4063	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4064	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4065	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4066	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4067	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4068	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4069	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4070	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4071	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4072	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4073	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4074	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4075	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4076	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4077	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4078	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4079	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4080	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4081	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4082	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4083	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4084	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4085	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4086	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4087	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4088	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4089	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4090	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4091	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4092	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4093	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4094	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4095	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4096	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4097	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4098	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	1	4099	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4100	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4101	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4102	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4103	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4104	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4105	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4106	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4107	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4108	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4109	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4110	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4111	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4112	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4113	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4114	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4115	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4116	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4117	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4118	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4119	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4120	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4121	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4122	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4123	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4124	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4125	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4126	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4127	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4128	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4129	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4130	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4131	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4132	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4133	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4134	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4135	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4136	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4137	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4138	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4139	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4140	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	1	4141	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4142	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4143	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4144	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4145	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4146	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4147	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4148	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4149	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4150	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4151	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4152	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4153	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4154	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4155	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4156	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4157	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4158	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4159	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4160	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4161	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4162	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4163	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4164	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4165	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4166	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4167	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4168	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4169	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4170	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4171	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4172	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4173	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4174	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4175	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4176	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4177	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4178	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4179	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4180	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4181	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4182	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	1	4183	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4184	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4185	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4186	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4187	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4188	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4189	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4190	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4191	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4192	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4193	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4194	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4195	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4196	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4197	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4198	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4199	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4200	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4201	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4202	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4203	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4204	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4205	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4206	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4207	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4208	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4209	-	-	0/0/0/0	0/0/0/0
86	OHX	1	4210	-	-	0/0/0/0	0/0/0/0
88	BLS	1	4211	-	-	0/17/38/38	0/2/2/2
86	OHX	2	2023	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2024	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2025	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2026	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2027	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2028	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2029	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2030	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2031	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2032	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2033	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2034	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2035	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	2	2036	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2037	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2038	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2039	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2040	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2041	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2042	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2043	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2044	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2045	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2046	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2047	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2048	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2049	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2050	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2051	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2052	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2053	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2054	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2055	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2056	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2057	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2058	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2059	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2060	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2061	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2062	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2063	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2064	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2065	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2066	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2067	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2068	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2069	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2070	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2071	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2072	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2073	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2074	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2075	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2076	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2077	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	2	2078	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2079	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2080	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2081	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2082	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2083	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2084	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2085	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2086	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2087	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2088	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2089	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2090	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2091	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2092	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2093	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2094	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2095	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2096	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2097	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2098	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2099	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2100	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2101	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2102	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2103	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2104	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2105	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2106	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2107	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2108	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2109	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2110	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2111	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2112	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2113	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2114	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2115	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2116	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2117	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2118	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2119	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	2	2120	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2121	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2122	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2123	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2124	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2125	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2126	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2127	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2128	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2129	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2130	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2131	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2132	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2133	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2134	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2135	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2136	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2137	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2138	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2139	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2140	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2141	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2142	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2143	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2144	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2145	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2146	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2147	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2148	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2149	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2150	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2151	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2152	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2153	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2154	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2155	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2156	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2157	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2158	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2159	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2160	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2161	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	2	2162	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2163	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2164	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2165	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2166	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2167	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2168	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2169	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2170	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2171	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2172	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2173	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2174	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2175	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2176	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2177	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2178	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2179	-	-	0/0/0/0	0/0/0/0
86	OHX	2	2180	-	-	0/0/0/0	0/0/0/0
86	OHX	3	216	-	-	0/0/0/0	0/0/0/0
86	OHX	3	217	-	-	0/0/0/0	0/0/0/0
86	OHX	3	218	-	-	0/0/0/0	0/0/0/0
86	OHX	3	219	-	-	0/0/0/0	0/0/0/0
86	OHX	3	220	-	-	0/0/0/0	0/0/0/0
86	OHX	3	221	-	-	0/0/0/0	0/0/0/0
86	OHX	3	222	-	-	0/0/0/0	0/0/0/0
86	OHX	3	223	-	-	0/0/0/0	0/0/0/0
86	OHX	3	224	-	-	0/0/0/0	0/0/0/0
86	OHX	3	225	-	-	0/0/0/0	0/0/0/0
86	OHX	4	224	-	-	0/0/0/0	0/0/0/0
86	OHX	4	225	-	-	0/0/0/0	0/0/0/0
86	OHX	4	226	-	-	0/0/0/0	0/0/0/0
86	OHX	4	227	-	-	0/0/0/0	0/0/0/0
86	OHX	4	228	-	-	0/0/0/0	0/0/0/0
86	OHX	4	229	-	-	0/0/0/0	0/0/0/0
86	OHX	4	230	-	-	0/0/0/0	0/0/0/0
86	OHX	4	231	-	-	0/0/0/0	0/0/0/0
86	OHX	4	232	-	-	0/0/0/0	0/0/0/0
86	OHX	4	233	-	-	0/0/0/0	0/0/0/0
86	OHX	4	234	-	-	0/0/0/0	0/0/0/0
86	OHX	4	235	-	-	0/0/0/0	0/0/0/0
86	OHX	4	236	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	4	237	-	-	0/0/0/0	0/0/0/0
86	OHX	4	238	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3892	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3893	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3894	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3895	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3896	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3897	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3898	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3899	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3900	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3901	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3902	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3903	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3904	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3905	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3906	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3907	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3908	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3909	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3910	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3911	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3912	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3913	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3914	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3915	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3916	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3917	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3918	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3919	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3920	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3921	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3922	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3923	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3924	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3925	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3926	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3927	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3928	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3929	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3930	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3931	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	5	3932	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3933	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3934	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3935	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3936	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3937	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3938	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3939	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3940	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3941	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3942	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3943	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3944	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3945	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3946	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3947	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3948	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3949	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3950	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3951	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3952	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3953	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3954	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3955	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3956	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3957	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3958	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3959	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3960	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3961	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3962	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3963	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3964	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3965	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3966	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3967	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3968	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3969	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3970	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3971	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3972	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3973	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	5	3974	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3975	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3976	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3977	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3978	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3979	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3980	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3981	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3982	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3983	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3984	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3985	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3986	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3987	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3988	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3989	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3990	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3991	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3992	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3993	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3994	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3995	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3996	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3997	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3998	-	-	0/0/0/0	0/0/0/0
86	OHX	5	3999	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4000	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4001	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4002	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4003	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4004	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4005	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4006	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4007	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4008	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4009	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4010	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4011	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4012	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4013	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4014	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4015	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	5	4016	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4017	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4018	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4019	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4020	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4021	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4022	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4023	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4024	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4025	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4026	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4027	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4028	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4029	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4030	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4031	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4032	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4033	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4034	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4035	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4036	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4037	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4038	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4039	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4040	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4041	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4042	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4043	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4044	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4045	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4046	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4047	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4048	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4049	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4050	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4051	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4052	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4053	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4054	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4055	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4056	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4057	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	5	4058	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4059	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4060	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4061	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4062	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4063	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4064	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4065	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4066	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4067	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4068	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4069	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4070	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4071	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4072	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4073	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4074	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4075	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4076	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4077	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4078	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4079	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4080	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4081	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4082	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4083	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4084	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4085	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4086	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4087	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4088	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4089	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4090	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4091	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4092	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4093	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4094	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4095	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4096	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4097	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4098	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4099	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	5	4100	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4101	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4102	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4103	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4104	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4105	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4106	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4107	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4108	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4109	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4110	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4111	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4112	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4113	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4114	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4115	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4116	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4117	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4118	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4119	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4120	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4121	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4122	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4123	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4124	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4125	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4126	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4127	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4128	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4129	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4130	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4131	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4132	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4133	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4134	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4135	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4136	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4137	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4138	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4139	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4140	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4141	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	5	4142	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4143	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4144	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4145	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4146	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4147	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4148	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4149	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4150	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4151	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4152	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4153	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4154	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4155	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4156	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4157	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4158	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4159	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4160	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4161	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4162	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4163	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4164	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4165	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4166	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4167	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4168	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4169	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4170	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4171	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4172	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4173	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4174	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4175	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4176	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4177	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4178	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4179	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4180	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4181	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4182	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4183	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	5	4184	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4185	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4186	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4187	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4188	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4189	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4190	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4191	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4192	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4193	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4194	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4195	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4196	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4197	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4198	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4199	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4200	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4201	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4202	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4203	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4204	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4205	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4206	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4207	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4208	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4209	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4210	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4211	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4212	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4213	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4214	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4215	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4216	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4217	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4218	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4219	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4220	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4221	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4222	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4223	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4224	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4225	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	5	4226	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4227	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4228	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4229	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4230	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4231	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4232	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4233	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4234	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4235	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4236	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4237	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4238	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4239	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4240	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4241	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4242	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4243	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4244	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4245	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4246	-	-	0/0/0/0	0/0/0/0
86	OHX	5	4247	-	-	0/0/0/0	0/0/0/0
88	BLS	5	4248	-	-	0/17/38/38	0/2/2/2
86	OHX	6	2046	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2047	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2048	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2049	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2050	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2051	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2052	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2053	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2054	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2055	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2056	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2057	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2058	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2059	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2060	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2061	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2062	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2063	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2064	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	6	2065	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2066	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2067	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2068	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2069	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2070	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2071	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2072	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2073	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2074	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2075	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2076	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2077	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2078	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2079	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2080	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2081	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2082	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2083	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2084	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2085	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2086	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2087	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2088	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2089	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2090	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2091	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2092	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2093	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2094	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2095	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2096	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2097	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2098	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2099	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2100	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2101	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2102	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2103	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2104	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2105	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2106	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	6	2107	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2108	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2109	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2110	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2111	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2112	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2113	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2114	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2115	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2116	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2117	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2118	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2119	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2120	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2121	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2122	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2123	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2124	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2125	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2126	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2127	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2128	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2129	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2130	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2131	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2132	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2133	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2134	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2135	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2136	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2137	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2138	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2139	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2140	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2141	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2142	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2143	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2144	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2145	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2146	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2147	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2148	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	6	2149	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2150	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2151	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2152	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2153	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2154	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2155	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2156	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2157	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2158	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2159	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2160	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2161	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2162	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2163	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2164	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2165	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2166	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2167	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2168	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2169	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2170	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2171	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2172	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2173	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2174	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2175	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2176	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2177	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2178	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2179	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2180	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2181	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2182	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2183	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2184	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2185	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2186	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2187	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2188	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2189	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2190	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	6	2191	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2192	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2193	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2194	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2195	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2196	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2197	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2198	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2199	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2200	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2201	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2202	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2203	-	-	0/0/0/0	0/0/0/0
86	OHX	6	2204	-	-	0/0/0/0	0/0/0/0
86	OHX	7	218	-	-	0/0/0/0	0/0/0/0
86	OHX	7	219	-	-	0/0/0/0	0/0/0/0
86	OHX	7	220	-	-	0/0/0/0	0/0/0/0
86	OHX	7	221	-	-	0/0/0/0	0/0/0/0
86	OHX	7	222	-	-	0/0/0/0	0/0/0/0
86	OHX	7	223	-	-	0/0/0/0	0/0/0/0
86	OHX	7	224	-	-	0/0/0/0	0/0/0/0
86	OHX	7	225	-	-	0/0/0/0	0/0/0/0
86	OHX	7	226	-	-	0/0/0/0	0/0/0/0
86	OHX	7	227	-	-	0/0/0/0	0/0/0/0
86	OHX	7	228	-	-	0/0/0/0	0/0/0/0
86	OHX	7	229	-	-	0/0/0/0	0/0/0/0
86	OHX	8	215	-	-	0/0/0/0	0/0/0/0
86	OHX	8	216	-	-	0/0/0/0	0/0/0/0
86	OHX	8	217	-	-	0/0/0/0	0/0/0/0
86	OHX	8	218	-	-	0/0/0/0	0/0/0/0
86	OHX	8	219	-	-	0/0/0/0	0/0/0/0
86	OHX	8	220	-	-	0/0/0/0	0/0/0/0
86	OHX	8	221	-	-	0/0/0/0	0/0/0/0
86	OHX	8	222	-	-	0/0/0/0	0/0/0/0
86	OHX	8	223	-	-	0/0/0/0	0/0/0/0
86	OHX	8	224	-	-	0/0/0/0	0/0/0/0
86	OHX	8	225	-	-	0/0/0/0	0/0/0/0
86	OHX	8	226	-	-	0/0/0/0	0/0/0/0
86	OHX	8	227	-	-	0/0/0/0	0/0/0/0
86	OHX	8	228	-	-	0/0/0/0	0/0/0/0
86	OHX	8	229	-	-	0/0/0/0	0/0/0/0
86	OHX	C3	201	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	C5	201	-	-	0/0/0/0	0/0/0/0
86	OHX	C8	201	-	-	0/0/0/0	0/0/0/0
86	OHX	D3	202	-	-	0/0/0/0	0/0/0/0
86	OHX	D9	102	-	-	0/0/0/0	0/0/0/0
86	OHX	L3	405	-	-	0/0/0/0	0/0/0/0
86	OHX	L3	406	-	-	0/0/0/0	0/0/0/0
86	OHX	L4	404	-	-	0/0/0/0	0/0/0/0
86	OHX	M0	304	-	-	0/0/0/0	0/0/0/0
86	OHX	M5	303	-	-	0/0/0/0	0/0/0/0
86	OHX	M7	206	-	-	0/0/0/0	0/0/0/0
86	OHX	M7	207	-	-	0/0/0/0	0/0/0/0
86	OHX	M9	202	-	-	0/0/0/0	0/0/0/0
86	OHX	N9	101	-	-	0/0/0/0	0/0/0/0
86	OHX	O1	202	-	-	0/0/0/0	0/0/0/0
86	OHX	O3	201	-	-	0/0/0/0	0/0/0/0
86	OHX	O7	104	-	-	0/0/0/0	0/0/0/0
86	OHX	O7	105	-	-	0/0/0/0	0/0/0/0
86	OHX	Q2	503	-	-	0/0/0/0	0/0/0/0
86	OHX	S8	302	-	-	0/0/0/0	0/0/0/0
86	OHX	SR	401	-	-	0/0/0/0	0/0/0/0
86	OHX	c3	201	-	-	0/0/0/0	0/0/0/0
86	OHX	c5	201	-	-	0/0/0/0	0/0/0/0
86	OHX	c8	202	-	-	0/0/0/0	0/0/0/0
86	OHX	d4	201	-	-	0/0/0/0	0/0/0/0
86	OHX	d9	102	-	-	0/0/0/0	0/0/0/0
86	OHX	l3	404	-	-	0/0/0/0	0/0/0/0
86	OHX	l3	405	-	-	0/0/0/0	0/0/0/0
86	OHX	l4	402	-	-	0/0/0/0	0/0/0/0
86	OHX	l4	403	-	-	0/0/0/0	0/0/0/0
86	OHX	l5	304	-	-	0/0/0/0	0/0/0/0
86	OHX	l5	305	-	-	0/0/0/0	0/0/0/0
86	OHX	l5	306	-	-	0/0/0/0	0/0/0/0
86	OHX	l9	202	-	-	0/0/0/0	0/0/0/0
86	OHX	m0	302	-	-	0/0/0/0	0/0/0/0
86	OHX	m0	303	-	-	0/0/0/0	0/0/0/0
86	OHX	m1	202	-	-	0/0/0/0	0/0/0/0
86	OHX	m4	202	-	-	0/0/0/0	0/0/0/0
86	OHX	m5	306	-	-	0/0/0/0	0/0/0/0
86	OHX	m6	203	-	-	0/0/0/0	0/0/0/0
86	OHX	m7	207	-	-	0/0/0/0	0/0/0/0
86	OHX	n1	201	-	-	0/0/0/0	0/0/0/0
86	OHX	n3	202	-	-	0/0/0/0	0/0/0/0

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
86	OHX	n9	103	-	-	0/0/0/0	0/0/0/0
86	OHX	o3	203	-	-	0/0/0/0	0/0/0/0
86	OHX	o7	103	-	-	0/0/0/0	0/0/0/0
86	OHX	q2	502	-	-	0/0/0/0	0/0/0/0
86	OHX	s1	302	-	-	0/0/0/0	0/0/0/0
86	OHX	s1	303	-	-	0/0/0/0	0/0/0/0
86	OHX	s4	301	-	-	0/0/0/0	0/0/0/0
86	OHX	s8	302	-	-	0/0/0/0	0/0/0/0
86	OHX	s9	201	-	-	0/0/0/0	0/0/0/0
86	OHX	sR	401	-	-	0/0/0/0	0/0/0/0

All (7) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
88	5	4248	BLS	C11-N12	-5.16	1.36	1.47
88	1	4211	BLS	C2-N3	-4.20	1.29	1.38
88	1	4211	BLS	C6-C5	-2.57	1.32	1.38
88	5	4248	BLS	C14-N12	2.10	1.40	1.35
88	1	4211	BLS	C11-N12	3.05	1.53	1.47
88	1	4211	BLS	C5-C4	3.49	1.49	1.41
88	1	4211	BLS	C6-N1	4.07	1.41	1.35

All (6) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
88	1	4211	BLS	C11-N12-C14	-4.36	116.17	121.53
88	1	4211	BLS	C6-N1-C2	-2.78	116.78	121.28
88	1	4211	BLS	C5-C4-N3	-2.57	118.62	121.68
88	1	4211	BLS	C5'-O5'-C1'	2.30	117.17	112.26
88	1	4211	BLS	C4'-N6-C7	3.27	126.81	123.28
88	1	4211	BLS	N4-C4-N3	3.33	122.25	116.64

There are no chirality outliers.

There are no torsion outliers.

There are no ring outliers.

463 monomers are involved in 698 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
86	1	3857	OHX	1	0
86	1	3860	OHX	1	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
86	1	3862	OHX	3	0
86	1	3864	OHX	1	0
86	1	3865	OHX	1	0
86	1	3867	OHX	1	0
86	1	3869	OHX	1	0
86	1	3871	OHX	1	0
86	1	3873	OHX	1	0
86	1	3874	OHX	1	0
86	1	3876	OHX	2	0
86	1	3877	OHX	1	0
86	1	3878	OHX	1	0
86	1	3879	OHX	2	0
86	1	3881	OHX	1	0
86	1	3882	OHX	1	0
86	1	3883	OHX	1	0
86	1	3884	OHX	1	0
86	1	3886	OHX	1	0
86	1	3894	OHX	2	0
86	1	3897	OHX	2	0
86	1	3899	OHX	1	0
86	1	3905	OHX	1	0
86	1	3908	OHX	1	0
86	1	3917	OHX	1	0
86	1	3919	OHX	2	0
86	1	3920	OHX	2	0
86	1	3924	OHX	2	0
86	1	3926	OHX	1	0
86	1	3927	OHX	2	0
86	1	3931	OHX	1	0
86	1	3932	OHX	1	0
86	1	3933	OHX	1	0
86	1	3935	OHX	2	0
86	1	3936	OHX	1	0
86	1	3940	OHX	1	0
86	1	3947	OHX	1	0
86	1	3950	OHX	1	0
86	1	3951	OHX	5	0
86	1	3952	OHX	1	0
86	1	3955	OHX	1	0
86	1	3957	OHX	3	0
86	1	3958	OHX	1	0
86	1	3961	OHX	1	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
86	1	3962	OHX	1	0
86	1	3965	OHX	6	0
86	1	3966	OHX	1	0
86	1	3970	OHX	1	0
86	1	3971	OHX	2	0
86	1	3974	OHX	1	0
86	1	3976	OHX	1	0
86	1	3977	OHX	1	0
86	1	3979	OHX	1	0
86	1	3982	OHX	1	0
86	1	3988	OHX	2	0
86	1	3991	OHX	1	0
86	1	3994	OHX	2	0
86	1	3996	OHX	1	0
86	1	3997	OHX	2	0
86	1	3998	OHX	5	0
86	1	4001	OHX	2	0
86	1	4003	OHX	1	0
86	1	4005	OHX	1	0
86	1	4008	OHX	1	0
86	1	4012	OHX	1	0
86	1	4014	OHX	1	0
86	1	4023	OHX	6	0
86	1	4025	OHX	1	0
86	1	4027	OHX	6	0
86	1	4029	OHX	1	0
86	1	4033	OHX	2	0
86	1	4034	OHX	2	0
86	1	4036	OHX	1	0
86	1	4038	OHX	1	0
86	1	4039	OHX	2	0
86	1	4040	OHX	7	0
86	1	4042	OHX	2	0
86	1	4048	OHX	1	0
86	1	4050	OHX	5	0
86	1	4051	OHX	1	0
86	1	4052	OHX	2	0
86	1	4057	OHX	1	0
86	1	4059	OHX	2	0
86	1	4065	OHX	1	0
86	1	4070	OHX	1	0
86	1	4076	OHX	2	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
86	1	4078	OHX	1	0
86	1	4080	OHX	2	0
86	1	4081	OHX	3	0
86	1	4086	OHX	1	0
86	1	4089	OHX	1	0
86	1	4092	OHX	1	0
86	1	4103	OHX	1	0
86	1	4118	OHX	2	0
86	1	4121	OHX	1	0
86	1	4124	OHX	2	0
86	1	4128	OHX	2	0
86	1	4129	OHX	1	0
86	1	4130	OHX	1	0
86	1	4131	OHX	1	0
86	1	4132	OHX	1	0
86	1	4134	OHX	1	0
86	1	4135	OHX	2	0
86	1	4136	OHX	1	0
86	1	4137	OHX	1	0
86	1	4138	OHX	1	0
86	1	4139	OHX	1	0
86	1	4140	OHX	2	0
86	1	4141	OHX	2	0
86	1	4143	OHX	6	0
86	1	4144	OHX	1	0
86	1	4147	OHX	2	0
86	1	4152	OHX	1	0
86	1	4153	OHX	6	0
86	1	4155	OHX	1	0
86	1	4160	OHX	4	0
86	1	4161	OHX	1	0
86	1	4162	OHX	1	0
86	1	4165	OHX	1	0
86	1	4166	OHX	1	0
86	1	4167	OHX	1	0
86	1	4169	OHX	6	0
86	1	4172	OHX	1	0
86	1	4177	OHX	1	0
86	1	4178	OHX	4	0
86	1	4180	OHX	1	0
86	1	4189	OHX	2	0
86	1	4190	OHX	1	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
86	1	4191	OHX	1	0
86	1	4194	OHX	7	0
86	1	4195	OHX	1	0
86	1	4200	OHX	3	0
86	1	4201	OHX	1	0
86	1	4205	OHX	1	0
86	1	4207	OHX	1	0
88	1	4211	BLS	2	0
86	2	2023	OHX	1	0
86	2	2024	OHX	1	0
86	2	2025	OHX	1	0
86	2	2026	OHX	2	0
86	2	2029	OHX	1	0
86	2	2031	OHX	8	0
86	2	2034	OHX	1	0
86	2	2036	OHX	2	0
86	2	2038	OHX	2	0
86	2	2040	OHX	1	0
86	2	2041	OHX	1	0
86	2	2043	OHX	6	0
86	2	2044	OHX	2	0
86	2	2045	OHX	1	0
86	2	2046	OHX	2	0
86	2	2050	OHX	1	0
86	2	2053	OHX	2	0
86	2	2055	OHX	1	0
86	2	2057	OHX	2	0
86	2	2061	OHX	1	0
86	2	2063	OHX	2	0
86	2	2066	OHX	1	0
86	2	2069	OHX	1	0
86	2	2072	OHX	1	0
86	2	2073	OHX	2	0
86	2	2074	OHX	4	0
86	2	2075	OHX	1	0
86	2	2077	OHX	2	0
86	2	2081	OHX	1	0
86	2	2085	OHX	1	0
86	2	2089	OHX	3	0
86	2	2091	OHX	2	0
86	2	2092	OHX	2	0
86	2	2095	OHX	1	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
86	2	2096	OHX	1	0
86	2	2097	OHX	1	0
86	2	2098	OHX	7	0
86	2	2099	OHX	1	0
86	2	2103	OHX	2	0
86	2	2104	OHX	1	0
86	2	2105	OHX	1	0
86	2	2106	OHX	1	0
86	2	2107	OHX	2	0
86	2	2108	OHX	1	0
86	2	2109	OHX	1	0
86	2	2110	OHX	3	0
86	2	2115	OHX	1	0
86	2	2116	OHX	1	0
86	2	2120	OHX	3	0
86	2	2122	OHX	1	0
86	2	2126	OHX	1	0
86	2	2129	OHX	2	0
86	2	2130	OHX	1	0
86	2	2131	OHX	5	0
86	2	2134	OHX	1	0
86	2	2138	OHX	1	0
86	2	2140	OHX	1	0
86	2	2143	OHX	1	0
86	2	2146	OHX	6	0
86	2	2149	OHX	1	0
86	2	2154	OHX	1	0
86	2	2155	OHX	1	0
86	2	2157	OHX	3	0
86	2	2159	OHX	1	0
86	2	2160	OHX	1	0
86	2	2162	OHX	4	0
86	2	2165	OHX	1	0
86	2	2167	OHX	1	0
86	2	2168	OHX	1	0
86	2	2173	OHX	1	0
86	3	216	OHX	1	0
86	3	218	OHX	1	0
86	3	221	OHX	1	0
86	3	224	OHX	1	0
86	4	226	OHX	1	0
86	4	229	OHX	2	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
86	4	232	OHX	1	0
86	4	234	OHX	1	0
86	4	236	OHX	1	0
86	5	3892	OHX	2	0
86	5	3894	OHX	1	0
86	5	3895	OHX	1	0
86	5	3897	OHX	1	0
86	5	3898	OHX	1	0
86	5	3899	OHX	1	0
86	5	3900	OHX	1	0
86	5	3901	OHX	1	0
86	5	3902	OHX	2	0
86	5	3903	OHX	1	0
86	5	3904	OHX	1	0
86	5	3905	OHX	2	0
86	5	3908	OHX	1	0
86	5	3909	OHX	3	0
86	5	3910	OHX	2	0
86	5	3912	OHX	1	0
86	5	3917	OHX	1	0
86	5	3920	OHX	1	0
86	5	3925	OHX	1	0
86	5	3926	OHX	1	0
86	5	3928	OHX	1	0
86	5	3929	OHX	1	0
86	5	3930	OHX	1	0
86	5	3932	OHX	1	0
86	5	3935	OHX	7	0
86	5	3938	OHX	1	0
86	5	3940	OHX	1	0
86	5	3943	OHX	1	0
86	5	3946	OHX	1	0
86	5	3947	OHX	1	0
86	5	3949	OHX	2	0
86	5	3950	OHX	3	0
86	5	3951	OHX	1	0
86	5	3952	OHX	1	0
86	5	3953	OHX	2	0
86	5	3955	OHX	3	0
86	5	3959	OHX	1	0
86	5	3962	OHX	2	0
86	5	3963	OHX	1	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
86	5	3965	OHX	2	0
86	5	3966	OHX	6	0
86	5	3968	OHX	1	0
86	5	3969	OHX	1	0
86	5	3971	OHX	8	0
86	5	3972	OHX	1	0
86	5	3975	OHX	1	0
86	5	3980	OHX	1	0
86	5	3982	OHX	1	0
86	5	3984	OHX	2	0
86	5	3989	OHX	1	0
86	5	3991	OHX	1	0
86	5	3992	OHX	1	0
86	5	3994	OHX	5	0
86	5	3995	OHX	6	0
86	5	3996	OHX	1	0
86	5	4000	OHX	2	0
86	5	4004	OHX	4	0
86	5	4006	OHX	1	0
86	5	4008	OHX	2	0
86	5	4014	OHX	7	0
86	5	4017	OHX	2	0
86	5	4018	OHX	2	0
86	5	4022	OHX	1	0
86	5	4023	OHX	1	0
86	5	4028	OHX	1	0
86	5	4029	OHX	4	0
86	5	4033	OHX	1	0
86	5	4040	OHX	1	0
86	5	4043	OHX	1	0
86	5	4045	OHX	1	0
86	5	4046	OHX	1	0
86	5	4047	OHX	1	0
86	5	4048	OHX	1	0
86	5	4049	OHX	1	0
86	5	4060	OHX	4	0
86	5	4061	OHX	1	0
86	5	4066	OHX	1	0
86	5	4068	OHX	1	0
86	5	4069	OHX	1	0
86	5	4074	OHX	1	0
86	5	4075	OHX	1	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
86	5	4082	OHX	1	0
86	5	4083	OHX	1	0
86	5	4085	OHX	7	0
86	5	4087	OHX	2	0
86	5	4088	OHX	2	0
86	5	4092	OHX	1	0
86	5	4095	OHX	1	0
86	5	4097	OHX	2	0
86	5	4100	OHX	2	0
86	5	4104	OHX	1	0
86	5	4105	OHX	2	0
86	5	4106	OHX	1	0
86	5	4112	OHX	1	0
86	5	4113	OHX	1	0
86	5	4114	OHX	1	0
86	5	4124	OHX	1	0
86	5	4126	OHX	2	0
86	5	4128	OHX	1	0
86	5	4133	OHX	1	0
86	5	4136	OHX	1	0
86	5	4138	OHX	3	0
86	5	4139	OHX	3	0
86	5	4142	OHX	1	0
86	5	4145	OHX	1	0
86	5	4154	OHX	2	0
86	5	4155	OHX	1	0
86	5	4157	OHX	1	0
86	5	4161	OHX	1	0
86	5	4163	OHX	1	0
86	5	4164	OHX	1	0
86	5	4172	OHX	1	0
86	5	4173	OHX	1	0
86	5	4175	OHX	1	0
86	5	4179	OHX	1	0
86	5	4180	OHX	1	0
86	5	4182	OHX	1	0
86	5	4184	OHX	2	0
86	5	4186	OHX	2	0
86	5	4187	OHX	1	0
86	5	4188	OHX	1	0
86	5	4190	OHX	1	0
86	5	4192	OHX	1	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
86	5	4193	OHX	10	0
86	5	4194	OHX	2	0
86	5	4195	OHX	5	0
86	5	4206	OHX	1	0
86	5	4209	OHX	1	0
86	5	4211	OHX	1	0
86	5	4212	OHX	6	0
86	5	4214	OHX	1	0
86	5	4218	OHX	2	0
86	5	4222	OHX	1	0
86	5	4228	OHX	7	0
86	5	4229	OHX	1	0
86	5	4230	OHX	1	0
86	5	4231	OHX	1	0
86	5	4232	OHX	4	0
86	5	4234	OHX	1	0
86	5	4238	OHX	8	0
86	5	4239	OHX	1	0
86	5	4244	OHX	1	0
88	5	4248	BLS	2	0
86	6	2046	OHX	1	0
86	6	2047	OHX	1	0
86	6	2049	OHX	1	0
86	6	2053	OHX	1	0
86	6	2055	OHX	1	0
86	6	2057	OHX	1	0
86	6	2059	OHX	5	0
86	6	2061	OHX	3	0
86	6	2062	OHX	1	0
86	6	2065	OHX	1	0
86	6	2068	OHX	1	0
86	6	2070	OHX	1	0
86	6	2071	OHX	2	0
86	6	2073	OHX	1	0
86	6	2074	OHX	2	0
86	6	2075	OHX	1	0
86	6	2077	OHX	1	0
86	6	2079	OHX	1	0
86	6	2081	OHX	1	0
86	6	2083	OHX	1	0
86	6	2086	OHX	1	0
86	6	2088	OHX	1	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
86	6	2091	OHX	1	0
86	6	2092	OHX	1	0
86	6	2093	OHX	1	0
86	6	2096	OHX	2	0
86	6	2098	OHX	1	0
86	6	2100	OHX	2	0
86	6	2102	OHX	1	0
86	6	2103	OHX	1	0
86	6	2104	OHX	2	0
86	6	2105	OHX	2	0
86	6	2106	OHX	1	0
86	6	2107	OHX	1	0
86	6	2109	OHX	1	0
86	6	2111	OHX	2	0
86	6	2112	OHX	1	0
86	6	2114	OHX	1	0
86	6	2117	OHX	2	0
86	6	2120	OHX	7	0
86	6	2122	OHX	1	0
86	6	2123	OHX	1	0
86	6	2124	OHX	1	0
86	6	2125	OHX	2	0
86	6	2126	OHX	2	0
86	6	2128	OHX	1	0
86	6	2130	OHX	1	0
86	6	2136	OHX	2	0
86	6	2137	OHX	1	0
86	6	2138	OHX	2	0
86	6	2143	OHX	2	0
86	6	2144	OHX	1	0
86	6	2146	OHX	1	0
86	6	2147	OHX	6	0
86	6	2149	OHX	1	0
86	6	2154	OHX	1	0
86	6	2157	OHX	1	0
86	6	2159	OHX	3	0
86	6	2163	OHX	1	0
86	6	2165	OHX	1	0
86	6	2171	OHX	7	0
86	6	2175	OHX	1	0
86	6	2178	OHX	2	0
86	6	2179	OHX	1	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
86	6	2182	OHX	1	0
86	6	2183	OHX	1	0
86	6	2184	OHX	3	0
86	6	2188	OHX	1	0
86	6	2189	OHX	1	0
86	6	2190	OHX	3	0
86	6	2192	OHX	3	0
86	6	2195	OHX	1	0
86	6	2201	OHX	1	0
86	6	2202	OHX	1	0
86	6	2204	OHX	1	0
86	7	218	OHX	2	0
86	7	220	OHX	7	0
86	7	221	OHX	1	0
86	7	222	OHX	1	0
86	7	223	OHX	1	0
86	7	228	OHX	6	0
86	8	215	OHX	1	0
86	8	216	OHX	5	0
86	8	219	OHX	1	0
86	8	222	OHX	1	0
86	8	223	OHX	5	0
86	8	228	OHX	1	0
86	8	229	OHX	1	0
86	C3	201	OHX	1	0
86	C5	201	OHX	3	0
86	D3	202	OHX	2	0
86	L3	405	OHX	1	0
86	L3	406	OHX	1	0
86	L4	404	OHX	2	0
86	M0	304	OHX	1	0
86	M5	303	OHX	1	0
86	M7	206	OHX	1	0
86	M7	207	OHX	1	0
86	M9	202	OHX	1	0
86	O1	202	OHX	7	0
86	O3	201	OHX	1	0
86	O7	104	OHX	6	0
86	O7	105	OHX	1	0
86	Q2	503	OHX	2	0
86	S8	302	OHX	1	0

5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

6 Fit of model and data

6.1 Protein, DNA and RNA chains

EDS failed to run properly - this section is therefore empty.

6.2 Non-standard residues in protein, DNA, RNA chains

EDS failed to run properly - this section is therefore empty.

6.3 Carbohydrates

EDS failed to run properly - this section is therefore empty.

6.4 Ligands

EDS failed to run properly - this section is therefore empty.

6.5 Other polymers

EDS failed to run properly - this section is therefore empty.