



Full wwPDB X-ray Structure Validation Report ⓘ

Mar 20, 2026 – 04:04 AM UTC

PDB ID : 6QNNQ / pdb_00006qnnq
Title : 70S ribosome initiation complex (IC) with experimentally assigned potassium ions
Authors : Rozov, A.; Khusainov, I.; Yusupov, M.; Yusupova, G.
Deposited on : 2019-02-11
Resolution : 3.50 Å(reported)

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

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

A user guide is available at

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

with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

<http://www.wwpdb.org/validation/2017/FAQs#types>.

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

MolProbity	:	4-5-2 with Phenix2.0
Mogul	:	2022.3.0, CSD as543be (2022)
Xtriage (Phenix)	:	2.0
EDS	:	3.0
Percentile statistics	:	20250101.v01 (using entries in the PDB archive January 1st 2025)
CCP4	:	9.0.010 (Gargrove)
Density-Fitness	:	1.0.12
Ideal geometry (proteins)	:	Engh & Huber (2001)
Ideal geometry (DNA, RNA)	:	Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP)	:	2.49

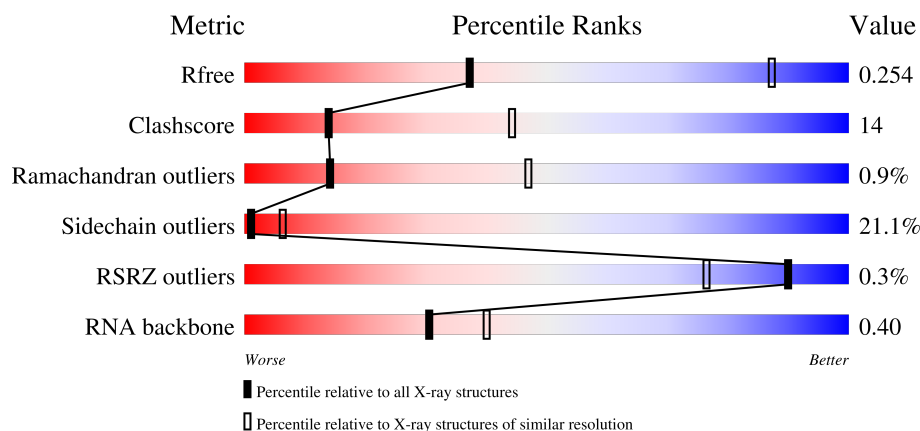
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

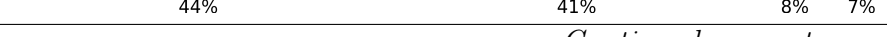
The reported resolution of this entry is 3.50 Å.

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



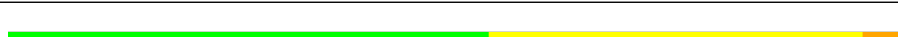
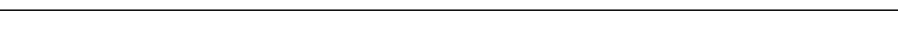
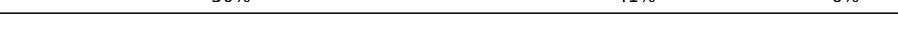
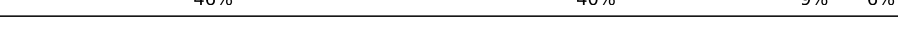
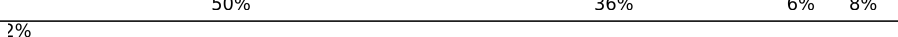
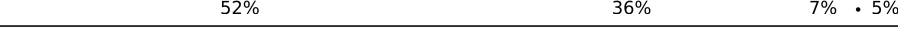
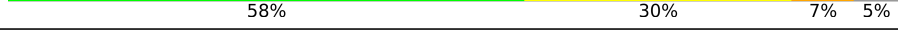
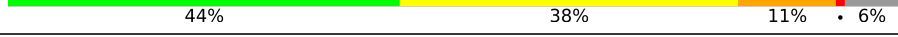
Metric	Whole archive (#Entries)	Similar resolution (#Entries, resolution range(Å))
R_{free}	180053	1085 (3.54-3.46)
Clashscore	190562	1140 (3.54-3.46)
Ramachandran outliers	187476	1113 (3.54-3.46)
Sidechain outliers	187428	1114 (3.54-3.46)
RSRZ outliers	180081	1084 (3.54-3.46)
RNA backbone	3983	1010 (3.98-3.02)

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

Mol	Chain	Length	Quality of chain
1	13	1522	
1	1G	1522	
2	12	256	
2	1E	256	

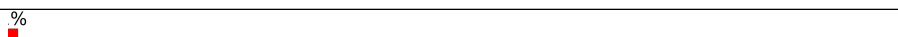
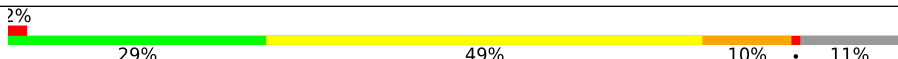
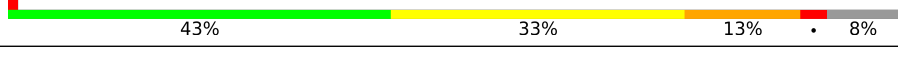
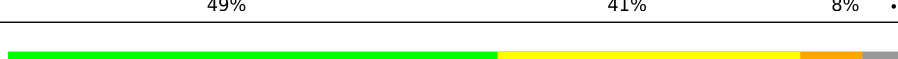
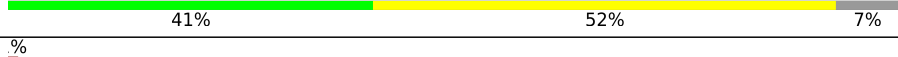
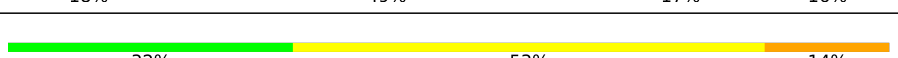
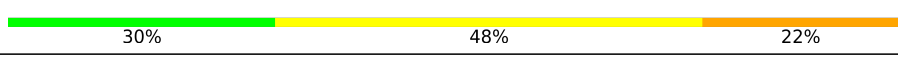
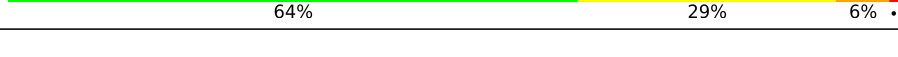
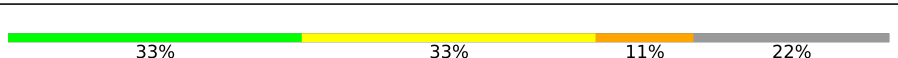
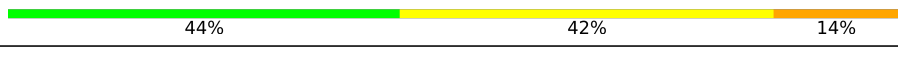
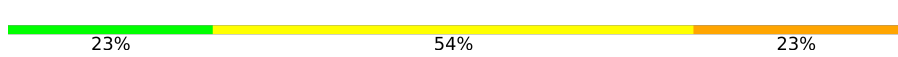
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Mol	Chain	Length	Quality of chain
3	22	239	
3	2E	239	
4	32	209	
4	3E	209	
5	42	162	
5	4E	162	
6	52	101	
6	5E	101	
7	62	156	
7	6E	156	
8	72	138	
8	7E	138	
9	82	128	
9	8E	128	
10	1A	105	
10	1I	105	
11	2A	129	
11	2I	129	
12	3A	132	
12	3I	132	
13	4A	126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

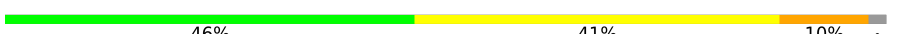
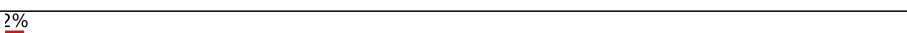
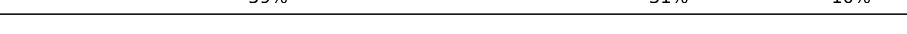
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Mol	Chain	Length	Quality of chain
15	6I	89	
16	7A	88	
16	7I	88	
17	8A	105	
17	8I	105	
18	9A	88	
18	9I	88	
19	AA	93	
19	AI	93	
20	BA	106	
20	BI	106	
21	1B	27	
21	1F	27	
22	1K	77	
22	3K	77	
22	3L	77	
23	2K	77	
24	4K	27	
24	4L	27	
25	14	2917	
25	1H	2917	
26	16	122	
26	1J	122	
27	7I	229	
28	11	276	

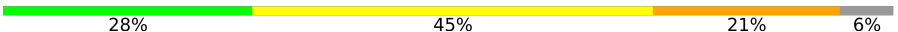
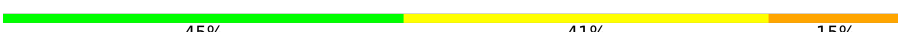
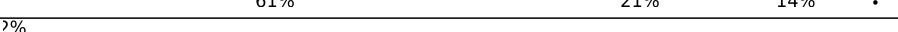
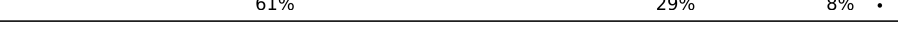
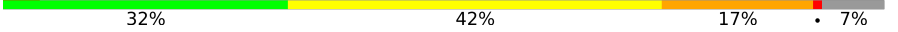
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Mol	Chain	Length	Quality of chain
28	19	276	 55% 38% 7% .
29	21	206	 61% 29% 9% .
29	29	206	 55% 34% 9% .
30	31	210	%  55% 33% 8% .
30	39	210	 46% 41% 10% .
31	41	182	 52% 39% 9% .
31	49	182	%  52% 37% 10% ..
32	51	180	 48% 38% 11% .
32	59	180	 49% 38% 10% .
33	61	148	%  42% 39% 16% ..
33	69	148	 50% 38% 11% .
34	38	173	%  35% 32% 12% . 20%
35	15	140	 59% 34% 5% .
35	58	140	 53% 35% 10% ..
36	25	122	2%  59% 31% 10%
36	68	122	 60% 37% .
37	35	150	%  42% 41% 15% ..
37	78	150	 50% 32% 16% ..
38	45	141	 50% 38% 11% ..
38	88	141	 58% 32% 9% .
39	55	118	 58% 35% 7% .
39	98	118	%  55% 33% 12%
40	65	112	 45% 48% 6% .
40	A8	112	 50% 38% 12% .
41	75	146	%  36% 43% 14% 6%

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

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Mol	Chain	Length	Quality of chain
54	K5	54	
54	O8	54	
55	L5	49	
55	P8	49	
56	M5	65	
56	Q8	65	
57	2L	77	

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
60	SF4	32	303	-	-	X	-

2 Entry composition

There are 62 unique types of molecules in this entry. The entry contains 300507 atoms, of which 0 are hydrogens and 0 are deuteriums.

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

- Molecule 1 is a RNA chain called 16S ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
1	13	1506	Total	C	N	O	P	0	0	0
			32387	14423	5999	10459	1506			
1	1G	1510	Total	C	N	O	P	0	0	0
			32470	14460	6012	10488	1510			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
2	1E	237	Total	C	N	O	S	0	0	0
			1924	1228	344	347	5			
2	12	236	Total	C	N	O	S	0	0	0
			1915	1223	343	344	5			

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

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

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

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

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
5	4E	151	Total	C	N	O	S	0	0	0
			1155	729	218	204	4			
5	42	154	Total	C	N	O	S	0	0	0
			1178	743	221	210	4			

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

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

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
7	6E	155	Total	C	N	O	S	0	0	0
			1256	781	252	217	6			
7	62	149	Total	C	N	O	S	0	0	0
			1217	759	242	210	6			

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

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

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
9	8E	124	Total	C	N	O		0	0	0
			983	624	190	169				
9	82	127	Total	C	N	O	S	0	0	0
			1002	635	193	173	1			

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

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

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

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
11	2I	117	Total	C	N	O	S	0	0	0
			864	537	162	162	3			
11	2A	119	Total	C	N	O	S	0	0	0
			889	554	169	163	3			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
12	3I	125	Total	C	N	O	S	0	0	0
			977	615	196	164	2			
12	3A	125	Total	C	N	O	S	0	0	0
			977	615	196	164	2			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
13	4I	119	Total	C	N	O	S	0	0	0
			946	585	195	164	2			
13	4A	118	Total	C	N	O	S	0	0	0
			937	579	193	163	2			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
14	5I	60	Total	C	N	O	S	0	0	0
			491	312	104	71	4			
14	5A	60	Total	C	N	O	S	0	0	0
			485	309	101	71	4			

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

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

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
16	7I	83	Total	C	N	O	S	0	0	0
			700	443	139	117	1			
16	7A	84	Total	C	N	O	S	0	0	0
			705	446	140	118	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
17	8I	100	Total	C	N	O	S	0	0	0
			834	534	155	143	2			
17	8A	99	Total	C	N	O	S	0	0	0
			819	525	150	142	2			

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

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
18	9I	70	Total	C	N	O	0	0	0
			573	367	112	94			
18	9A	71	Total	C	N	O	0	0	0
			584	373	116	95			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
19	AI	86	Total	C	N	O	S	0	0	0
			688	438	128	120	2			
19	AA	83	Total	C	N	O	S	0	0	0
			665	424	122	117	2			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
20	BI	101	Total	C	N	O	S	0	0	0
			766	473	161	130	2			
20	BA	103	Total	C	N	O	S	0	0	0
			776	479	163	132	2			

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

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

- Molecule 22 is a RNA chain called E. coli tRNA^{fMet}.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
22	1K	65	Total	C	N	O	P	0	0	0
			1385	618	249	454	64			
22	3K	77	Total	C	N	O	P	0	0	0
			1640	732	297	535	76			
22	3L	77	Total	C	N	O	P	0	0	0
			1640	732	297	535	76			

- Molecule 23 is a RNA chain called E. coli tRNA^{fMet}.

Mol	Chain	Residues	Atoms						ZeroOcc	AltConf	Trace
23	2K	77	Total	C	N	O	P	S	0	0	0
			1643	735	297	534	76	1			

- Molecule 24 is a RNA chain called mRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
24	4K	17	Total	C	N	O	P	0	0	0
			373	168	79	109	17			
24	4L	21	Total	C	N	O	P	0	0	0
			463	208	99	135	21			

- Molecule 25 is a RNA chain called 23S ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
25	1H	2879	Total	C	N	O	P	0	0	0
			62010	27605	11592	19935	2878			
25	14	2879	Total	C	N	O	P	0	0	0
			62013	27604	11596	19934	2879			

- Molecule 26 is a RNA chain called 5S ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
26	16	120	Total	C	N	O	P	0	0	0
			2573	1146	476	832	119			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
26	1J	122	Total	C	N	O	P	0	0	0
			2617	1166	486	844	121			

- Molecule 27 is a protein called 50S ribosomal protein L1.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
27	71	135	Total	C	N	O	S	0	0	0
			1049	662	197	189	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
28	11	274	Total	C	N	O	S	0	0	0
			2125	1341	422	359	3			
28	19	274	Total	C	N	O	S	0	0	0
			2124	1341	421	359	3			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
29	21	204	Total	C	N	O	S	0	0	0
			1559	985	298	270	6			
29	29	204	Total	C	N	O	S	0	0	0
			1559	985	298	270	6			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
30	31	202	Total	C	N	O	S	0	0	0
			1585	1011	297	275	2			
30	39	206	Total	C	N	O	S	0	0	0
			1619	1033	302	281	3			

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

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

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

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

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

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

- Molecule 34 is a protein called 50S ribosomal protein L10.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
34	38	139	Total	C	N	O	S	0	0	0
			1056	672	187	195	2			

- Molecule 35 is a protein called 50S ribosomal protein L13.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
35	58	138	Total	C	N	O	S	0	0	0
			1104	712	206	182	4			
35	15	138	Total	C	N	O	S	0	0	0
			1104	712	206	182	4			

- Molecule 36 is a protein called 50S ribosomal protein L14.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
36	68	122	Total	C	N	O	S	0	0	0
			932	588	171	169	4			
36	25	122	Total	C	N	O	S	0	0	0
			932	588	171	169	4			

- Molecule 37 is a protein called 50S ribosomal protein L15.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
37	78	148	Total	C	N	O	S	0	0	0
			1130	704	230	193	3			
37	35	147	Total	C	N	O	S	0	0	0
			1122	698	229	192	3			

- Molecule 38 is a protein called 50S ribosomal protein L16.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
38	88	141	Total	C	N	O	S	0	0	0
			1121	715	212	187	7			
38	45	140	Total	C	N	O	S	0	0	0
			1112	710	210	185	7			

- Molecule 39 is a protein called 50S ribosomal protein L17.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
39	98	118	Total	C	N	O	S	0	0	0
			967	604	203	159	1			
39	55	118	Total	C	N	O	S	0	0	0
			967	604	203	159	1			

- Molecule 40 is a protein called 50S ribosomal protein L18.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
40	A8	111	Total	C	N	O	0	0	0
			881	556	176	149			
40	65	111	Total	C	N	O	0	0	0
			881	556	176	149			

- Molecule 41 is a protein called 50S ribosomal protein L19.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
41	B8	137	Total	C	N	O	S	0	0	0
			1141	710	234	196	1			
41	75	137	Total	C	N	O	S	0	0	0
			1141	710	234	196	1			

- Molecule 42 is a protein called 50S ribosomal protein L20.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
42	C8	117	Total	C	N	O	S	0	0	0
			963	610	202	150	1			
42	85	117	Total	C	N	O	S	0	0	0
			963	610	202	150	1			

- Molecule 43 is a protein called 50S ribosomal protein L21.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
43	D8	100	Total	C	N	O	S	0	0	0
			774	499	141	133	1			
43	95	101	Total	C	N	O	S	0	0	0
			778	501	142	134	1			

- Molecule 44 is a protein called 50S ribosomal protein L22.

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

- Molecule 45 is a protein called 50S ribosomal protein L23.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
45	F8	95	Total	C	N	O	S	0	0	0
			747	485	135	126	1			
45	B5	93	Total	C	N	O		0	0	0
			730	474	132	124				

- Molecule 46 is a protein called 50S ribosomal protein L24.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
46	G8	107	Total	C	N	O	S	0	0	0
			805	517	151	132	5			
46	C5	109	Total	C	N	O	S	0	0	0
			825	528	153	139	5			

- Molecule 47 is a protein called 50S ribosomal protein L25.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
47	H8	174	Total	C	N	O	S	0	0	0
			1390	887	250	250	3			
47	D5	176	Total	C	N	O	S	0	0	0
			1404	897	252	252	3			

- Molecule 48 is a protein called 50S ribosomal protein L27.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
48	I8	77	Total	C	N	O	S	0	0	0
			611	378	129	103	1			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
48	E5	79	Total	C	N	O	S	0	0	0
			623	386	131	105	1			

- Molecule 49 is a protein called 50S ribosomal protein L28.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
49	J8	96	Total	C	N	O	S	0	0	0
			754	474	149	129	2			
49	F5	92	Total	C	N	O	S	0	0	0
			721	451	144	125	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
50	K8	71	Total	C	N	O	S	0	0	0
			590	367	119	103	1			
50	G5	69	Total	C	N	O	S	0	0	0
			580	358	118	103	1			

- Molecule 51 is a protein called 50S ribosomal protein L30.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
51	L8	59	Total	C	N	O	0	0	0
			468	298	90	80			
51	H5	59	Total	C	N	O	0	0	0
			468	298	90	80			

- Molecule 52 is a protein called 50S ribosomal protein L31.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
52	M8	66	Total	C	N	O	S	0	0	0
			533	335	96	97	5			
52	I5	71	Total	C	N	O	S	0	0	0
			580	364	108	103	5			

- Molecule 53 is a protein called 50S ribosomal protein L32.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
53	N8	56	Total	C	N	O	S	0	0	0
			434	272	87	70	5			
53	J5	56	Total	C	N	O	S	0	0	0
			434	272	87	70	5			

- Molecule 54 is a protein called 50S ribosomal protein L33.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
54	O8	49	Total	C	N	O	S	0	0	0
			424	264	87	69	4			
54	K5	48	Total	C	N	O	S	0	0	0
			417	259	86	68	4			

- Molecule 55 is a protein called 50S ribosomal protein L34.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
55	P8	48	Total	C	N	O	S	0	0	0
			418	257	104	55	2			
55	L5	48	Total	C	N	O	S	0	0	0
			418	257	104	55	2			

- Molecule 56 is a protein called 50S ribosomal protein L35.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
56	Q8	64	Total	C	N	O	S	0	0	0
			506	326	99	79	2			
56	M5	64	Total	C	N	O	S	0	0	0
			506	326	99	79	2			

- Molecule 57 is a RNA chain called E. coli tRNA^fMet.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace	
57	2L	77	Total	C	N	O	P	S	0	0	0
			1643	735	297	534	76	1			

- Molecule 58 is POTASSIUM ION (CCD ID: K) (formula: K) (labeled as "Ligand of Interest" by depositor).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
58	13	37	Total	K	0	1
			38	38		
58	5E	1	Total	K	0	0
			1	1		
58	5I	1	Total	K	0	0
			1	1		
58	8I	1	Total	K	0	0
			1	1		
58	1H	121	Total	K	0	1
			122	122		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
58	16	2	Total 2	K 2	0	0
58	11	1	Total 1	K 1	0	0
58	21	1	Total 1	K 1	0	0
58	31	2	Total 2	K 2	0	0
58	41	2	Total 2	K 2	0	0
58	1G	25	Total 25	K 25	0	0
58	32	1	Total 1	K 1	0	0
58	52	1	Total 1	K 1	0	0
58	3A	1	Total 1	K 1	0	0
58	BA	1	Total 1	K 1	0	0
58	14	94	Total 94	K 94	0	0
58	19	1	Total 1	K 1	0	0
58	29	2	Total 2	K 2	0	0
58	39	1	Total 1	K 1	0	0

- Molecule 59 is MAGNESIUM ION (CCD ID: MG) (formula: Mg).

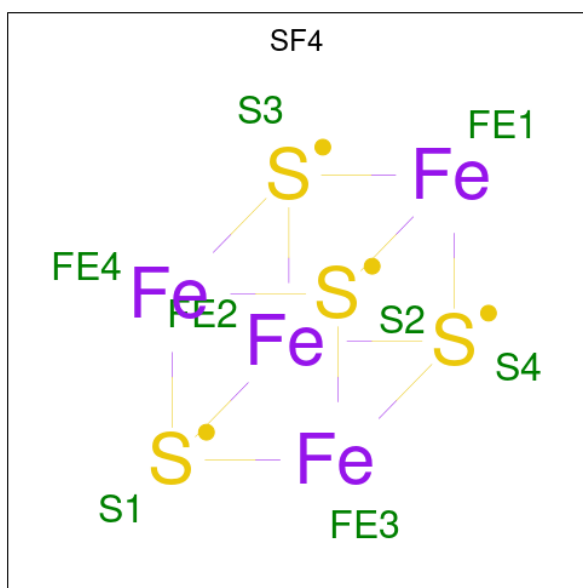
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	13	111	Total 111	Mg 111	0	0
59	3E	1	Total 1	Mg 1	0	0
59	8E	1	Total 1	Mg 1	0	0
59	4I	1	Total 1	Mg 1	0	0
59	7I	1	Total 1	Mg 1	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	BI	2	Total 2	Mg 2	0	0
59	2K	1	Total 1	Mg 1	0	0
59	1H	404	Total 405	Mg 405	0	1
59	16	8	Total 8	Mg 8	0	0
59	21	2	Total 2	Mg 2	0	0
59	41	1	Total 1	Mg 1	0	0
59	78	1	Total 1	Mg 1	0	0
59	D8	3	Total 3	Mg 3	0	0
59	G8	1	Total 1	Mg 1	0	0
59	I8	2	Total 2	Mg 2	0	0
59	J8	1	Total 1	Mg 1	0	0
59	N8	1	Total 1	Mg 1	0	0
59	Q8	1	Total 1	Mg 1	0	0
59	1G	102	Total 102	Mg 102	0	0
59	32	1	Total 1	Mg 1	0	0
59	2L	2	Total 2	Mg 2	0	0
59	14	312	Total 312	Mg 312	0	0
59	1J	5	Total 5	Mg 5	0	0
59	29	2	Total 2	Mg 2	0	0
59	55	1	Total 1	Mg 1	0	0
59	E5	1	Total 1	Mg 1	0	0

- Molecule 60 is IRON/SULFUR CLUSTER (CCD ID: SF4) (formula: Fe_4S_4).



Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
60	3E	1	Total	Fe	S	0	0
			8	4	4		
60	32	1	Total	Fe	S	0	0
			8	4	4		

- Molecule 61 is ZINC ION (CCD ID: ZN) (formula: Zn).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
61	5I	1	Total	Zn	0	0
			1	1		
61	5A	1	Total	Zn	0	0
			1	1		

- Molecule 62 is water.

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
62	13	190	Total	O	0	0
			190	190		
62	3E	2	Total	O	0	0
			2	2		
62	3I	1	Total	O	0	0
			1	1		
62	4I	2	Total	O	0	0
			2	2		
62	5I	1	Total	O	0	0
			1	1		

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Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
62	6I	1	Total O 1 1	0	0
62	7I	3	Total O 3 3	0	0
62	BI	4	Total O 4 4	0	0
62	3K	1	Total O 1 1	0	0
62	4K	2	Total O 2 2	0	0
62	1H	983	Total O 983 983	0	0
62	16	12	Total O 12 12	0	0
62	11	10	Total O 10 10	0	0
62	21	5	Total O 5 5	0	0
62	31	6	Total O 6 6	0	0
62	58	1	Total O 1 1	0	0
62	78	6	Total O 6 6	0	0
62	98	1	Total O 1 1	0	0
62	B8	2	Total O 2 2	0	0
62	C8	2	Total O 2 2	0	0
62	F8	2	Total O 2 2	0	0
62	G8	1	Total O 1 1	0	0
62	I8	5	Total O 5 5	0	0
62	J8	3	Total O 3 3	0	0
62	Q8	1	Total O 1 1	0	0
62	1G	268	Total O 268 268	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
62	32	4	Total 4	O 4	0	0
62	42	1	Total 1	O 1	0	0
62	3A	1	Total 1	O 1	0	0
62	7A	5	Total 5	O 5	0	0
62	BA	1	Total 1	O 1	0	0
62	2L	6	Total 6	O 6	0	0
62	4L	4	Total 4	O 4	0	0
62	14	681	Total 681	O 681	0	0
62	1J	16	Total 16	O 16	0	0
62	19	12	Total 12	O 12	0	0
62	29	3	Total 3	O 3	0	0
62	39	3	Total 3	O 3	0	0
62	35	3	Total 3	O 3	0	0
62	55	2	Total 2	O 2	0	0
62	75	1	Total 1	O 1	0	0
62	85	1	Total 1	O 1	0	0
62	B5	2	Total 2	O 2	0	0
62	C5	2	Total 2	O 2	0	0
62	F5	1	Total 1	O 1	0	0
62	G5	1	Total 1	O 1	0	0
62	L5	1	Total 1	O 1	0	0

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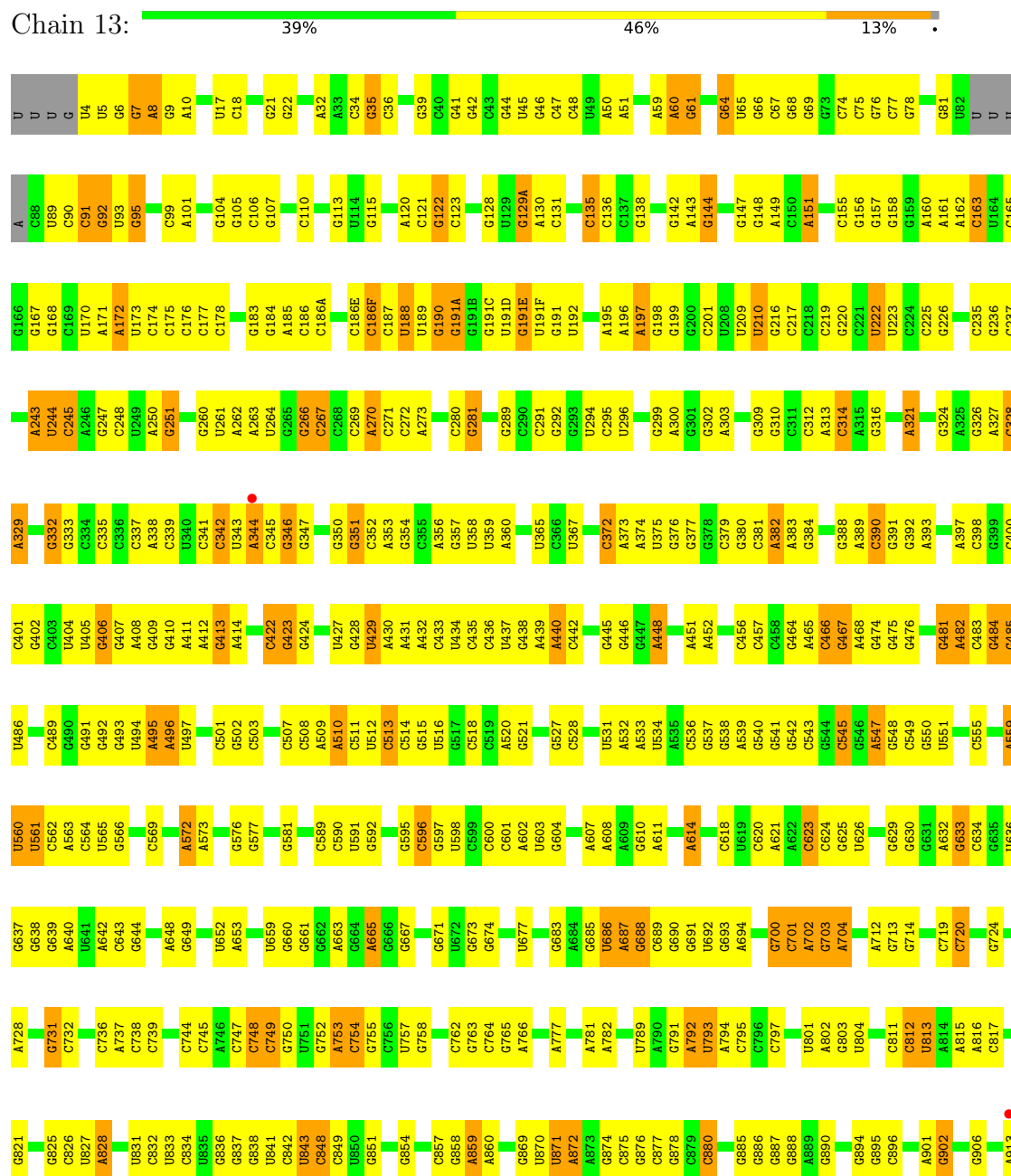
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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
62	M5	1	Total	O	0	0
			1	1		

3 Residue-property plots [i](#)

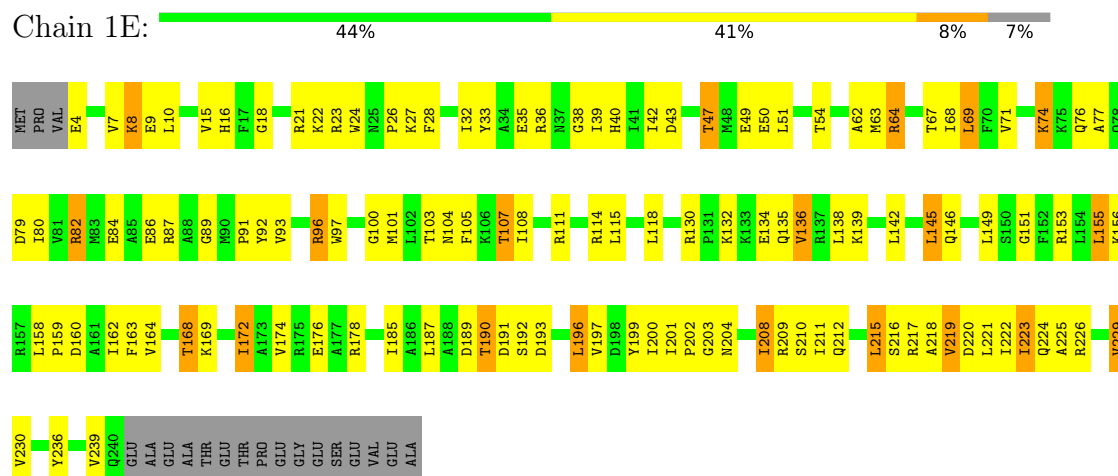
These plots are drawn for all protein, RNA, DNA and oligosaccharide chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and electron density.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

• Molecule 1: 16S ribosomal RNA

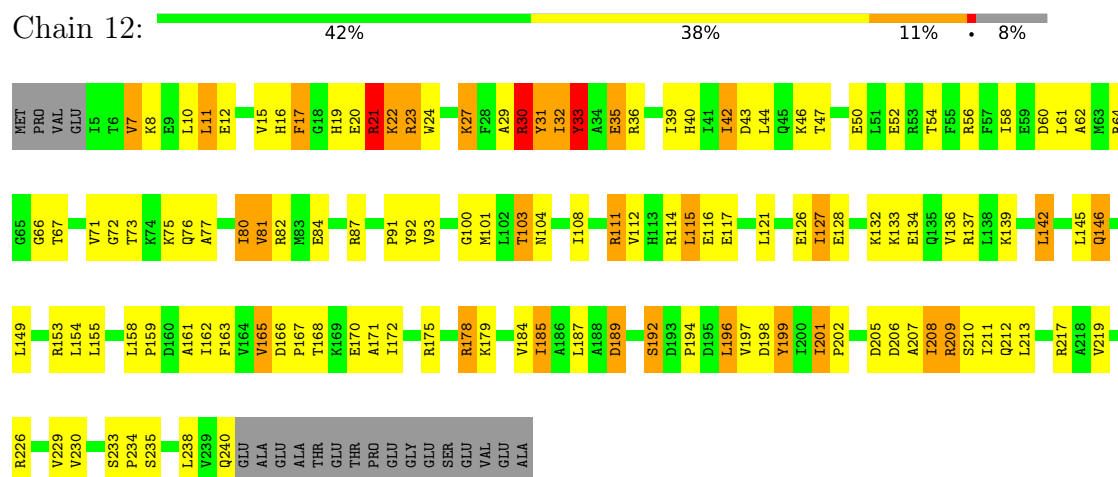


U1481	U1380	A1318	A1252	A1188	U1126	U1062	G1001	C932	G758	A694	G587	A509	C422
G1482	U1381	A1319	G1253	G1189	U1127	G1063	G1002	G933	C762	G695	G595	A510	G423
A1483	G1387	C1320	G1254	G1190	C1128	G1064	G1003	C934	G763	U686	G596	C511	G424
C1490	C1388	C1321	G1255	G1191	C1129	A1005	A1004	A935	G764	G688	G597	U516	G425
G1491	C1389	G1323	U1257	C1195	A1130	C1066	C1006	A937	G765	C689	U598	G517	G426
A1492	G1392	A1324	G1258	U1196	G1131	A1067	C1007	G942	A766	G690	C599	C518	U429
U1493	U1393	C1325	C1259	G1197	C1132	G1068	C1008	U943	A767	G691	C601	C519	A430
A1494	C1394	C1327	G1198	G1199	G1134	G1069	G1009	U944	A768	G692	G602	A520	C435
U1495	C1395	C1328	C1200	C1071	U1135	C1070	G1010	A946	G769	G693	G603	G521	C436
A1500	G1263	C1263	G1263	A1201	U1136	G1072	G1011	G947	G770	C701	A607	C522	A439
A1502	G1266	G1266	G1266	G1202	C1137	U1073	A1014	C948	G773	A702	A608	G527	A440
A1503	C1267	C1267	C1267	C1203	G1138	G1074	A1015	C949	G774	G703	G610	C442	C442
G1504	A1268	A1268	A1268	A1204	G1139	C1075	G1016	G951	A777	A532	C613	A532	G445
G1505	A1269	A1269	A1269	U1205	C1140	C1076	G1017	U952	G778	A533	C614	A533	G446
U1506	C1270	G1270	G1270	G1206	C1141	G1077	C1018	G953	C779	U534	C615	U534	G447
A1507	G1271	G1271	G1271	G1207	G1142	G1077	C1019	G954	A780	A535	C616	C536	G448
G1508	G1272	C1208	G1272	C1208	G1143	A1080	U1020	U955	C784	U536	C617	G537	A448
C1509	G1273	G1273	G1273	G1208	G1144	G1081	G1023	U956	C785	A452	C618	G538	A452
U1510	G1274	G1274	G1274	U1211	C1145	G1082	G1024	U957	G786	A453	C619	G539	A453
G1511	A1275	A1275	A1275	A1212	A1146	U1083	U1025	U960	C787	G714	U619	A539	G457
U1512	G1276	G1276	G1276	A1213	C1147	G1084	G1026	U961	C788	A715	C620	G540	C458
A1513	C1277	C1277	C1277	G1214	U1148	U1085	G1027	C962	A792	A716	A621	G541	C459
C1514	U1278	U1278	U1278	G1215	C1149	U1086	C1028	G963	C791	C717	C624	C543	G464
C1515	A1279	A1279	A1279	G1216	U1150	G1087	C1029	G964	A793	G718	C625	C544	A465
G1516	U1280	U1280	U1280	C1217	A1151	G1088	C1028B	A965	A794	C719	C626	C545	A466
G1517	A1281	A1281	A1281	C1218	G1154	U1089	G1030	G966	C797	G720	C627	C546	C466
A1518	C1282	C1282	C1282	U1219	C1155	U1090	G1031	C967	C806	A728	C628	A547	C467
A1519	G1283	G1283	G1283	G1220	G1156	U1091	G1032	A968	A807	A729	G629	G548	A468
G1520	C1284	C1284	C1284	G1221	C1157	A1092	G1033	C969	C810	G730	C630	C555	C474
G1521	U1285	U1285	U1285	G1222	A1158	U1093	A1034	A974	C811	C731	C631	C556	G475
G1526	A1286	A1286	A1286	C1223	C1159	G1094	G1035	A975	C812	C735	C632	U552	G476
U1529	U1287	U1287	U1287	U1224	U1159	U1095	G1036	A976	U813	C736	C633	A553	G477
A1531	A1288	A1288	A1288	A1225	C1160	C1096	C1037	A977	A814	A737	C634	C554	A478
U1532	C1355	C1355	C1355	G1226	G1161	C1097	G1038	A978	A815	C738	C635	C555	C479
C1533	G1362	G1362	G1362	U1232	A1167	G1104	A1042	C980	A816	C739	A653	A563	G489
C1535	U1296	U1296	U1296	G1105	C1169	A1105	G1043	U981	C817	U740	C636	C567	G490
C1536	C1297	C1297	C1297	G1106	A1170	G1106	A1044	U982	C818	G741	C637	G568	G491
U1537	C1298	C1298	C1298	C1109	G1174	C1109	C1045	A983	A819	G742	C638	C569	U494
U1540	A1299	A1299	A1299	A1110	C1175	A1110	A1046	A986	U820	U743	G661	A572	A495
U	G1300	G1300	G1300	A1238	A1176	C1111	G1047	A987	C821	C744	A663	A573	A496
U	U1301	U1301	U1301	A1239	G1177	C1112	G1048	G988	C822	C745	G664	A574	U497
C	C1302	C1302	C1302	U1240	C1178	C1113	U1049	C989	C823	A665	C665	G575	A498
U	G1303	G1303	G1303	G1241	G1179	C1114	G1050	C990	C824	C746	G500	C501	C502
U	C1304	C1304	C1304	G1242	A1179	C1115	C1051	U920	C825	C747	G501	C503	C504
	G1305	G1305	G1305	C1243	A1180	C1116	U1052	U921	A828	G750	G668	G576	G505
	A1306	A1306	A1306	A1244	G1181	G1117	G1053	U991	C829	U751	G673	G577	G506
	G1309	G1309	G1309	C1118	G1182	C1118	C1054	G993	C832	G752	G674	U582	G507
	G1314	G1314	G1314	C1119	A1183	C1119	U1056	A994	C833	A753	A675	A583	C508
	C1314	C1314	C1314	U1122	G1184	U1057	G1057	C998A	C834	C754	C675	G584	G509
	U1315	U1315	U1315	G1123	G1185	G1058	G1058	G927	U835	C755	G682	G585	C510
	G1317	G1317	G1317	G1124	G1186	G1059	C1059	G929	U836	U757	G683	C586	C511

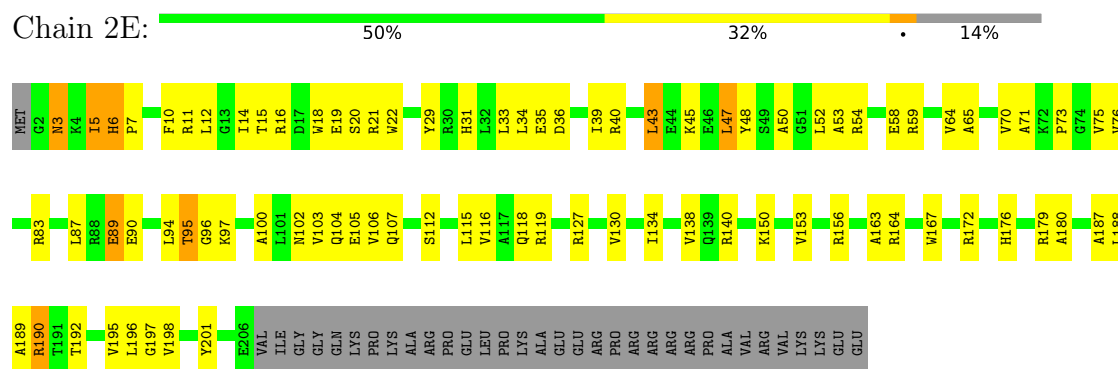
- Molecule 2: 30S ribosomal protein S2



- Molecule 2: 30S ribosomal protein S2

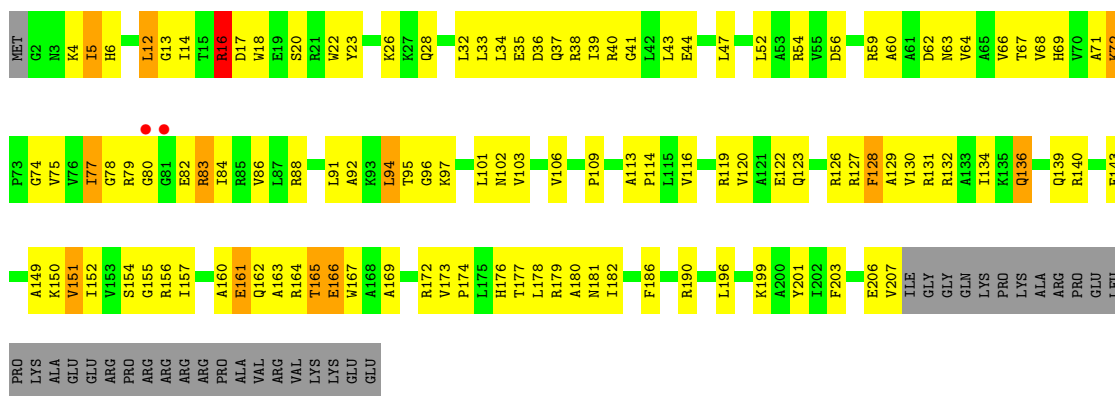


- Molecule 3: 30S ribosomal protein S3

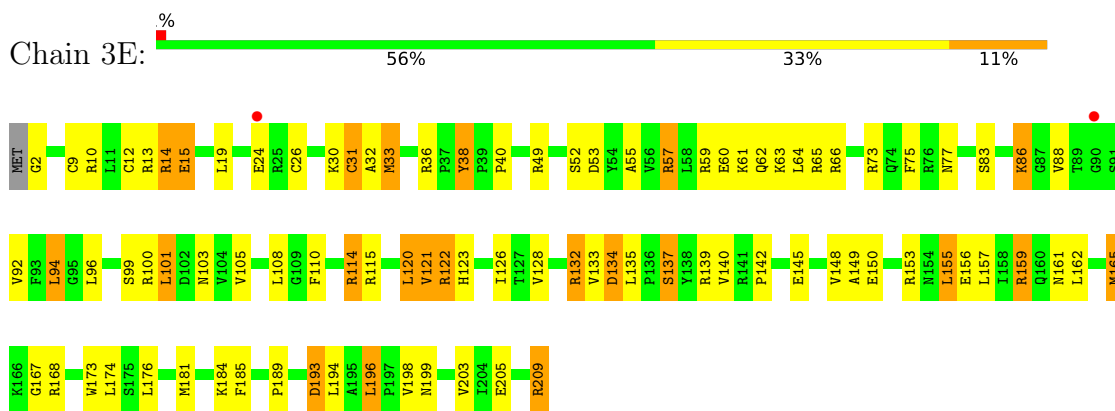


- Molecule 3: 30S ribosomal protein S3

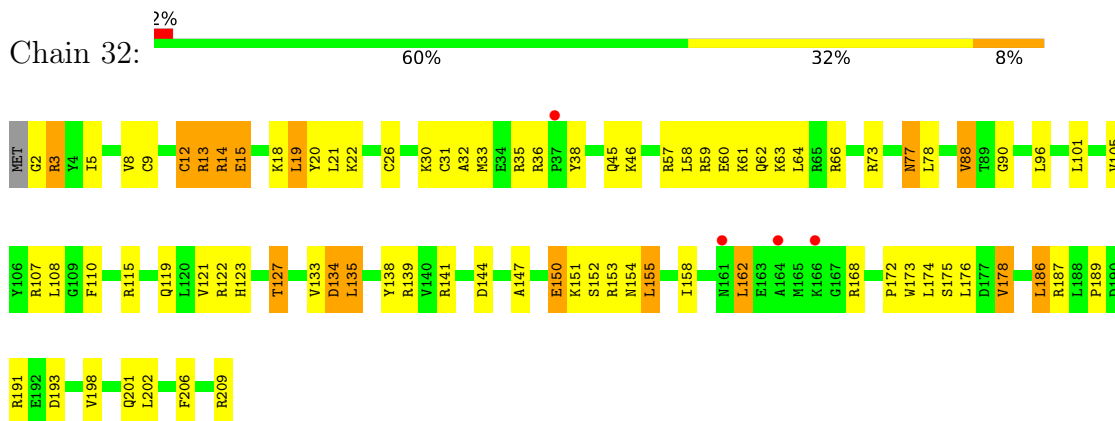




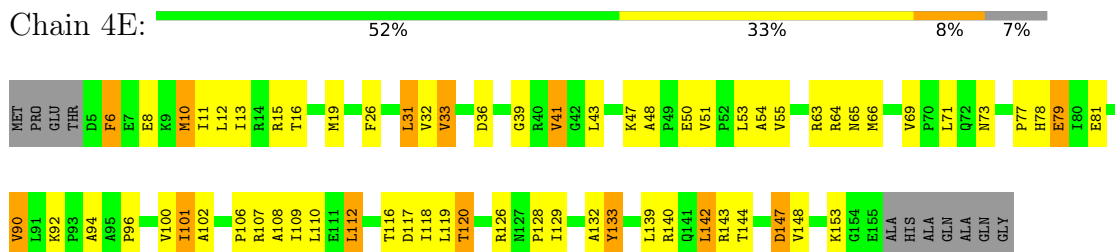
• Molecule 4: 30S ribosomal protein S4



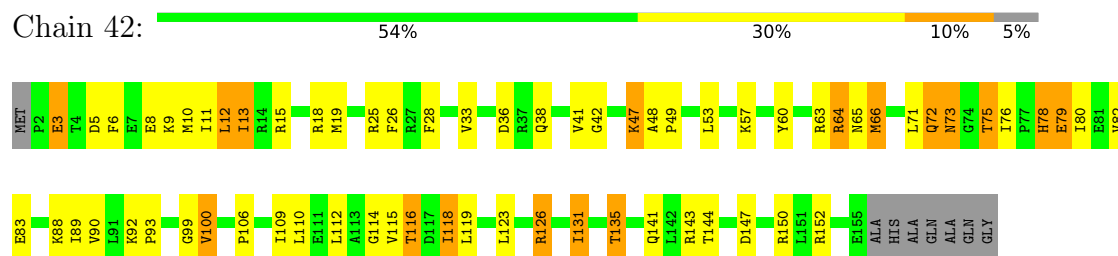
• Molecule 4: 30S ribosomal protein S4



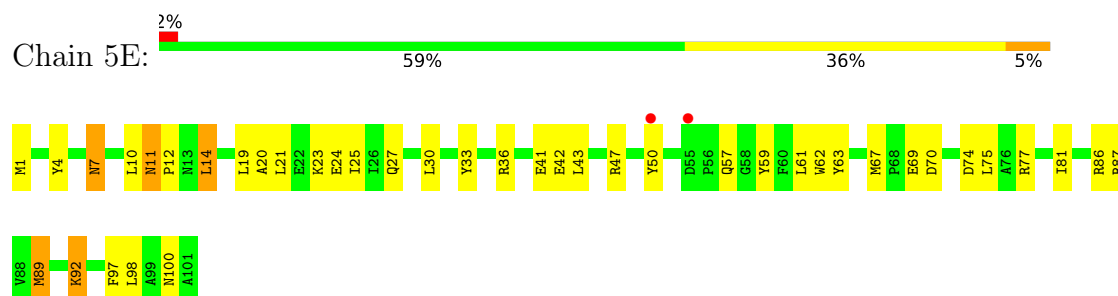
• Molecule 5: 30S ribosomal protein S5



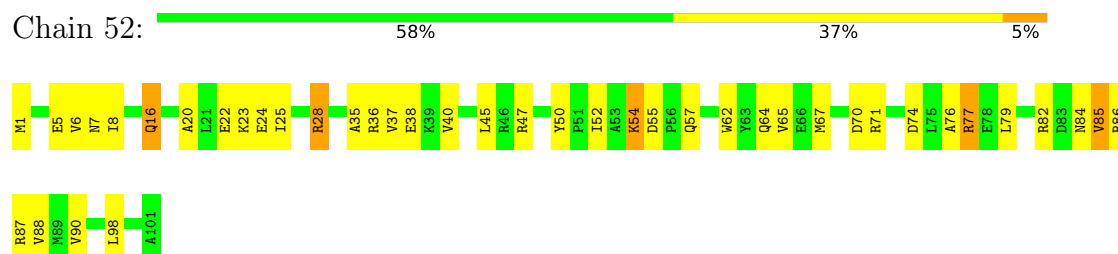
- Molecule 5: 30S ribosomal protein S5



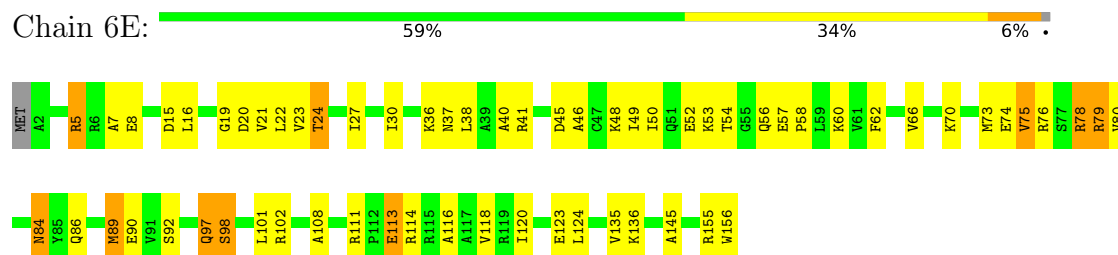
- Molecule 6: 30S ribosomal protein S6



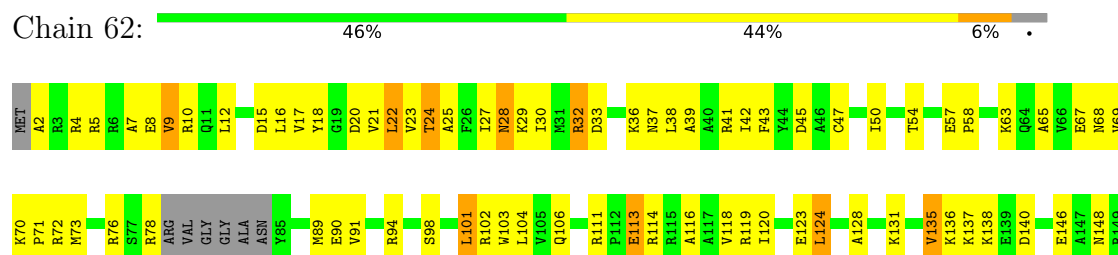
- Molecule 6: 30S ribosomal protein S6



- Molecule 7: 30S ribosomal protein S7



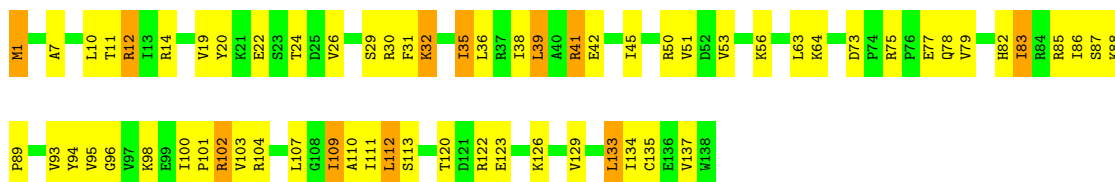
- Molecule 7: 30S ribosomal protein S7





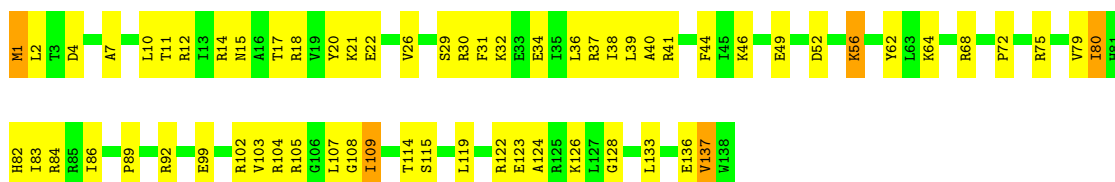
- Molecule 8: 30S ribosomal protein S8

Chain 7E: 53% 39% 8%



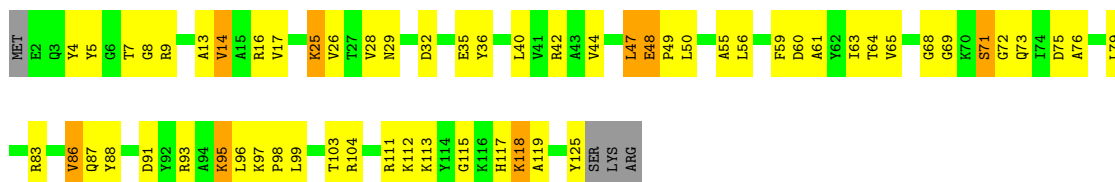
- Molecule 8: 30S ribosomal protein S8

Chain 72: 54% 42% .



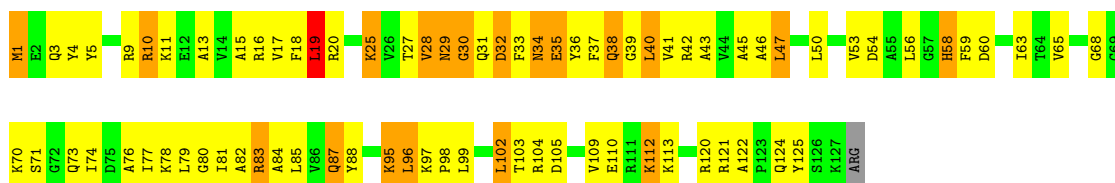
- Molecule 9: 30S ribosomal protein S9

Chain 8E: 50% 41% 6% .



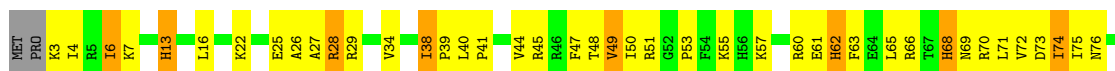
- Molecule 9: 30S ribosomal protein S9

Chain 82: 38% 46% 15% ..



- Molecule 10: 30S ribosomal protein S10

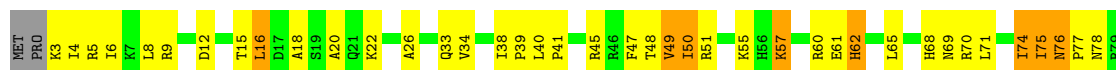
Chain 1I: 48% 38% 9% 6%





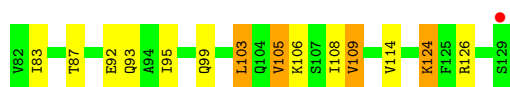
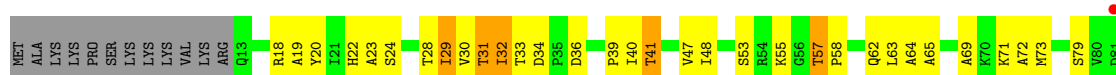
- Molecule 10: 30S ribosomal protein S10

Chain 1A: 46% 40% 9% 6%



- Molecule 11: 30S ribosomal protein S11

Chain 2I: 2% 55% 29% 7% 9%



- Molecule 11: 30S ribosomal protein S11

Chain 2A: 50% 36% 6% 8%



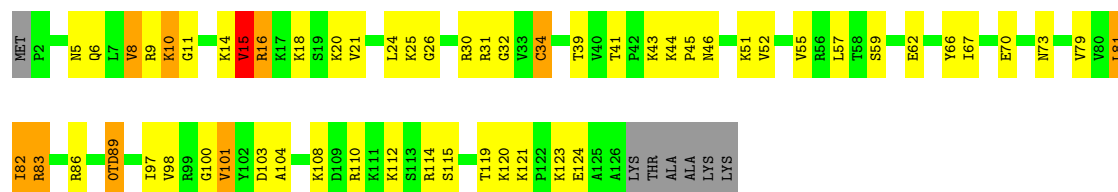
- Molecule 12: 30S ribosomal protein S12

Chain 3I: 58% 30% 7% 5%

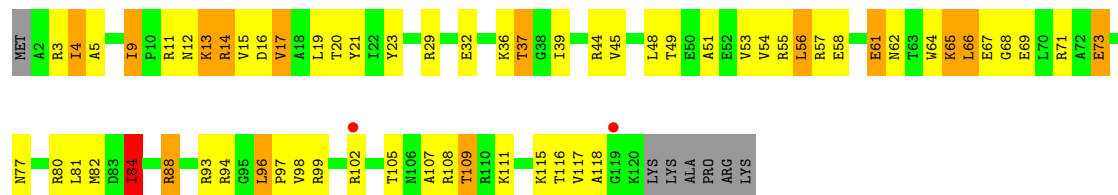
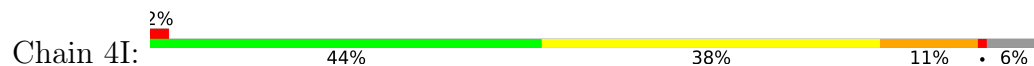


- Molecule 12: 30S ribosomal protein S12

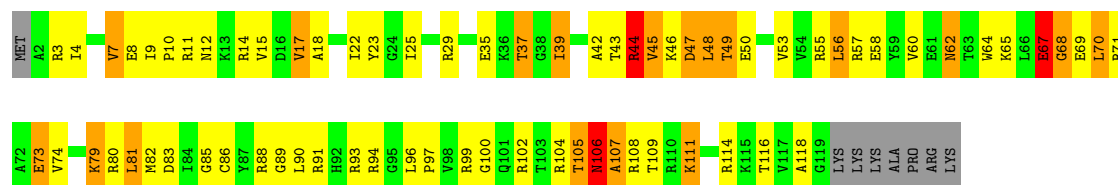
Chain 3A: 52% 36% 7% 5%



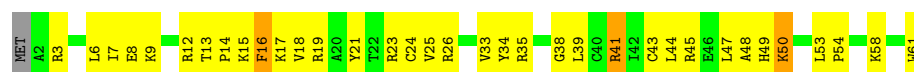
• Molecule 13: 30S ribosomal protein S13



• Molecule 13: 30S ribosomal protein S13



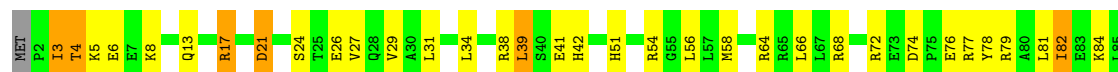
• Molecule 14: 30S ribosomal protein S14 type Z



• Molecule 14: 30S ribosomal protein S14 type Z



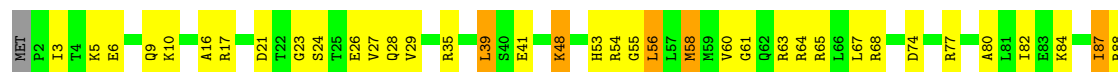
• Molecule 15: 30S ribosomal protein S15





- Molecule 15: 30S ribosomal protein S15

Chain 6A: 57% 36% 6%



- Molecule 16: 30S ribosomal protein S16

Chain 7I: 51% 36% 7% 6%



- Molecule 16: 30S ribosomal protein S16

Chain 7A: 61% 24% 10% 5%



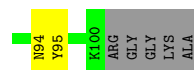
- Molecule 17: 30S ribosomal protein S17

Chain 8I: 2% 57% 32% 6% 5%

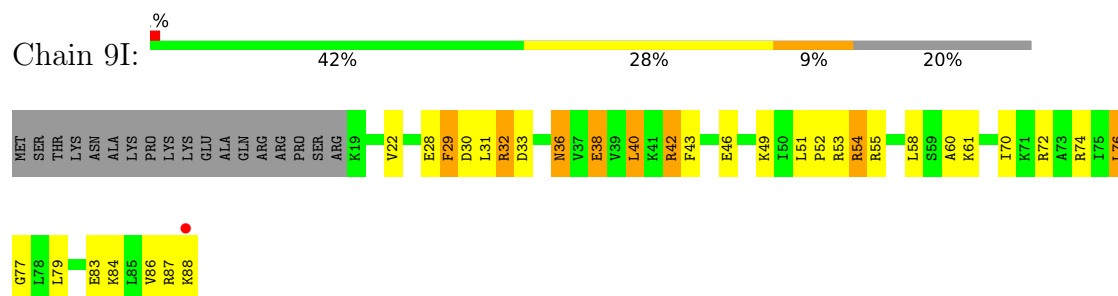


- Molecule 17: 30S ribosomal protein S17

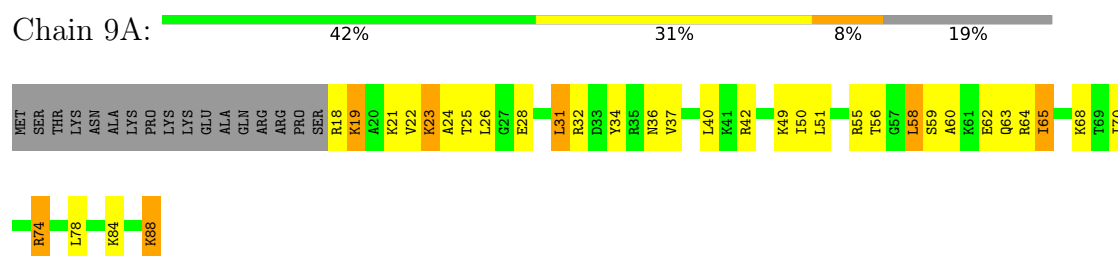
Chain 8A: 54% 36% 6%



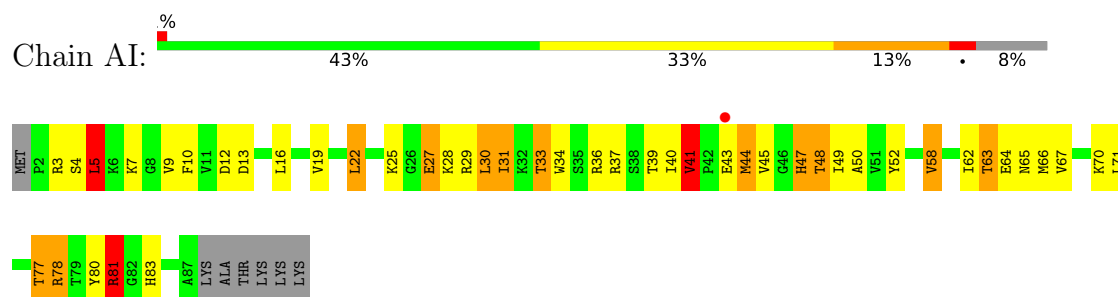
• Molecule 18: 30S ribosomal protein S18



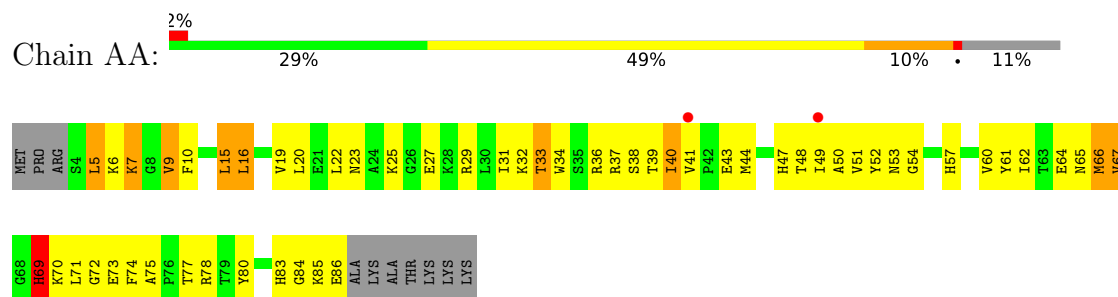
• Molecule 18: 30S ribosomal protein S18



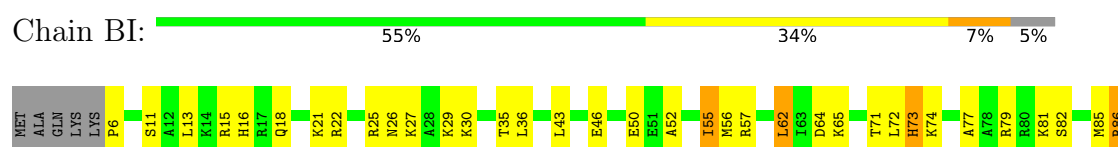
• Molecule 19: 30S ribosomal protein S19



• Molecule 19: 30S ribosomal protein S19



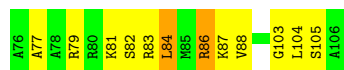
• Molecule 20: 30S ribosomal protein S20





- Molecule 20: 30S ribosomal protein S20

Chain BA: 49% 41% 8%



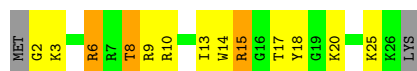
- Molecule 21: 30S ribosomal protein Thx

Chain 1F: 41% 52% 7%



- Molecule 21: 30S ribosomal protein Thx

Chain 1B: 44% 37% 11% 7%



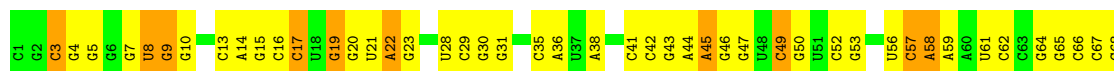
- Molecule 22: E. coli tRNA^{fMet}

Chain 1K: 18% 49% 17% 16%



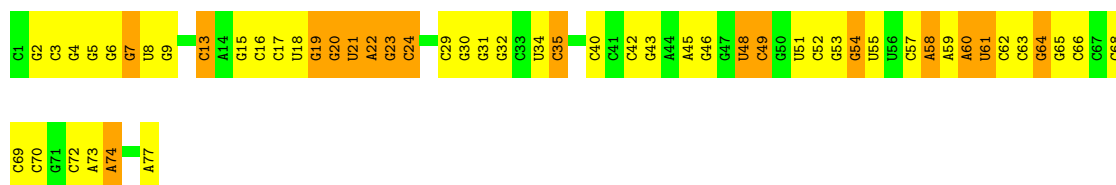
- Molecule 22: E. coli tRNA^{fMet}

Chain 3K: 32% 53% 14%



- Molecule 22: E. coli tRNA^{fMet}

Chain 3L: 30% 48% 22%



• Molecule 23: E. coli tRNA^{fMet}

Chain 2K: 64% 29% 6%



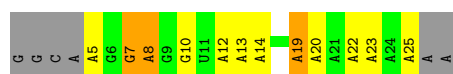
• Molecule 24: mRNA

Chain 4K: 7% 33% 22% 37%



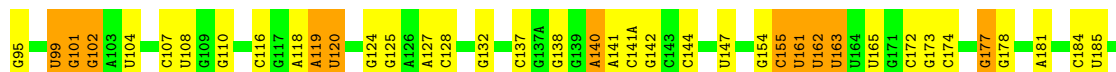
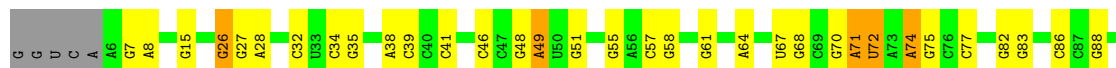
• Molecule 24: mRNA

Chain 4L: 33% 33% 11% 22%

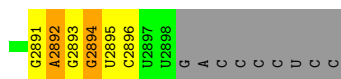


• Molecule 25: 23S ribosomal RNA

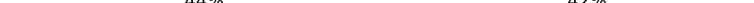
Chain 1H: 47% 39% 12%

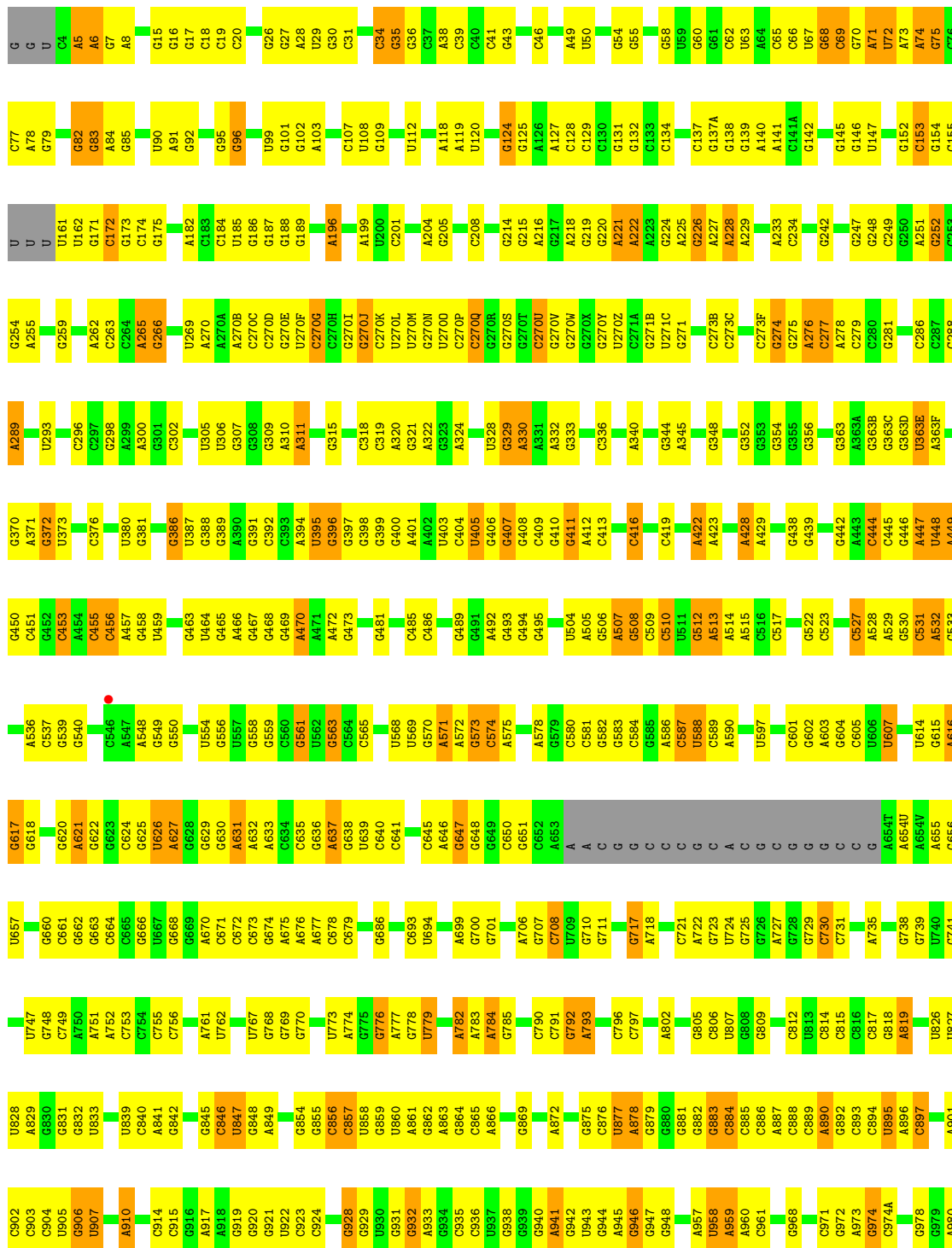




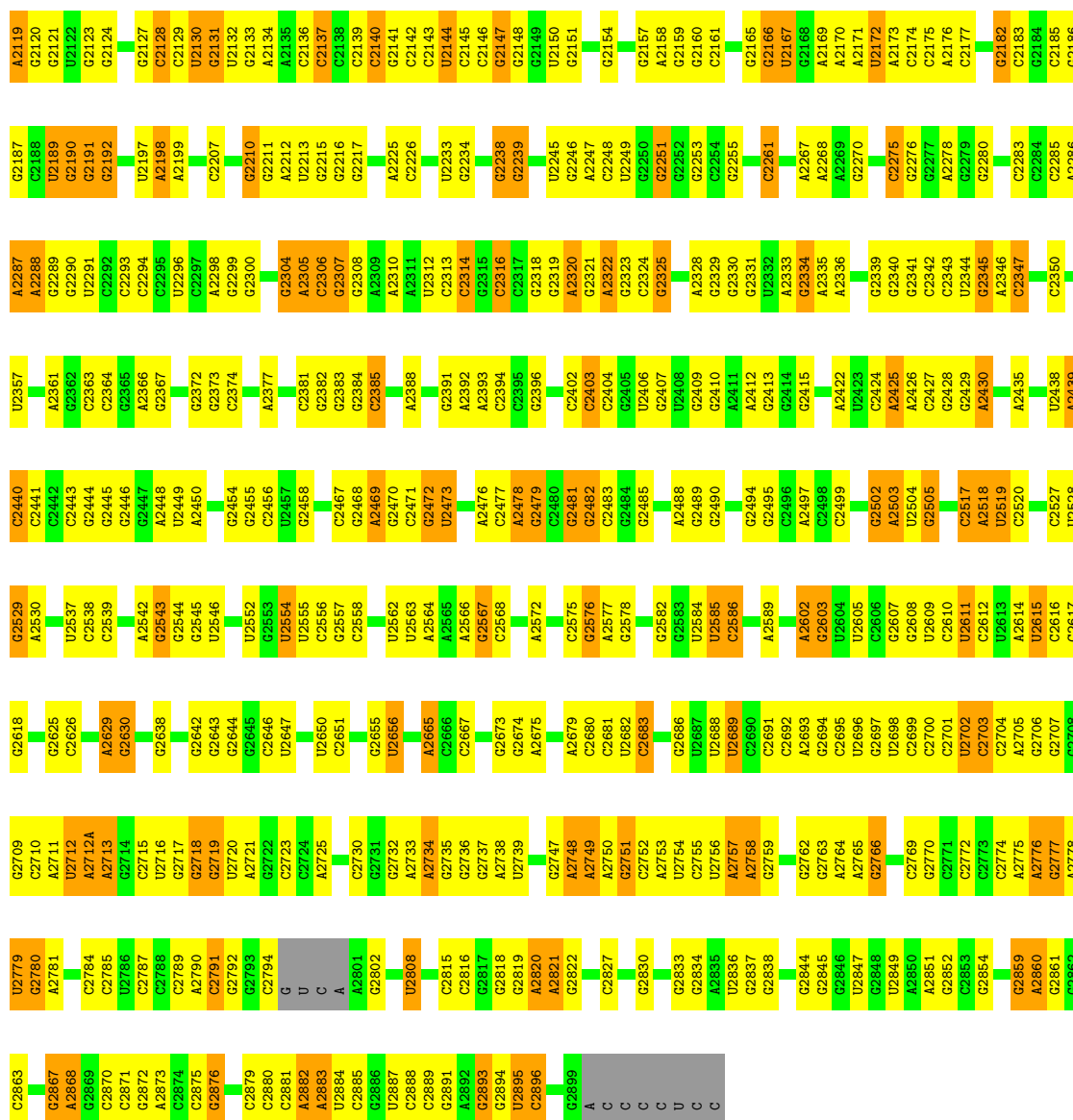


- Molecule 25: 23S ribosomal RNA

Chain 14:  44% 42% 14%

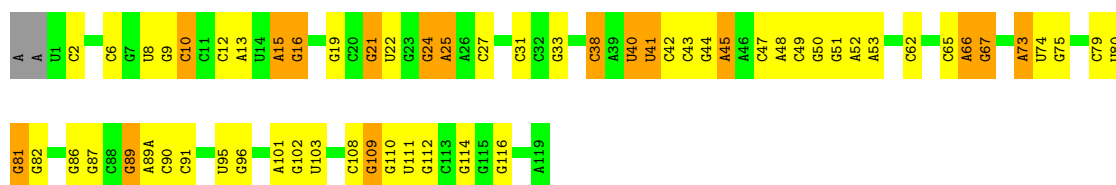


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A2054	G1964	A1791	G1695	G1595	A1427	A1349	G1266	G1187	G1120	C1052	A983
C2055	C1965	G1792	G1695	A1510	C1428	U1352	G1271	U1188	G1120	C1053	A984
G2056	A1966	C1793	U1697	C1598	C1429	U1353	A1272	G1189	G1122	A1054	C985
A2057	C1967	U1794	U1514	C1599	C1430	A1354	A1278	G1190	G1122	G1055	C986
C2058	G1982	C1795	C1515	G1603	U1431	A1354	A1278	G1191	G1122	G1056	C987
A2059	G1883	C1796	G1519	C1607	U1431	G1358	A1278	G1192	G1125	A1057	A990
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										G1114	A1048
										G1115	C1049



• Molecule 26: 5S ribosomal RNA

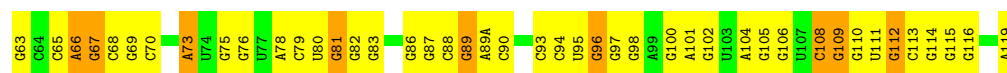
Chain 16: 49% 36% 13%



• Molecule 26: 5S ribosomal RNA

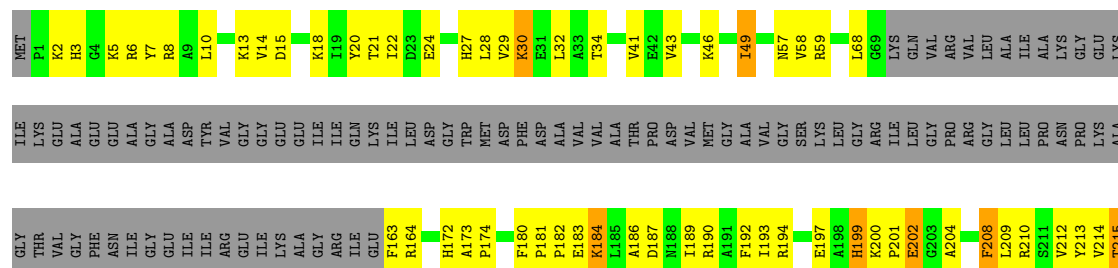
Chain 1J: 23% 54% 23%





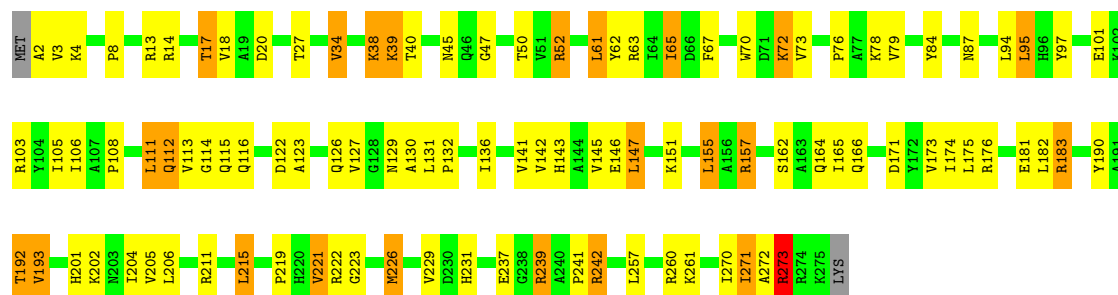
• Molecule 27: 50S ribosomal protein L1

Chain 71: 29% 26% 41%



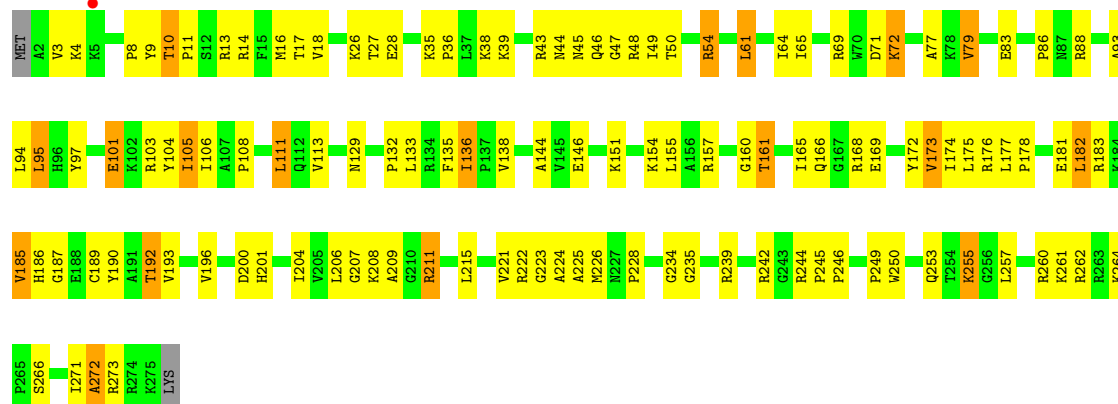
• Molecule 28: 50S ribosomal protein L2

Chain 11: 62% 29% 8%



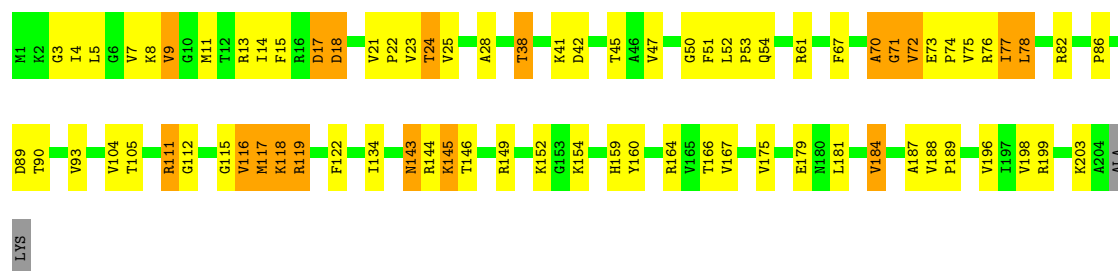
• Molecule 28: 50S ribosomal protein L2

Chain 19: 55% 38% 7%



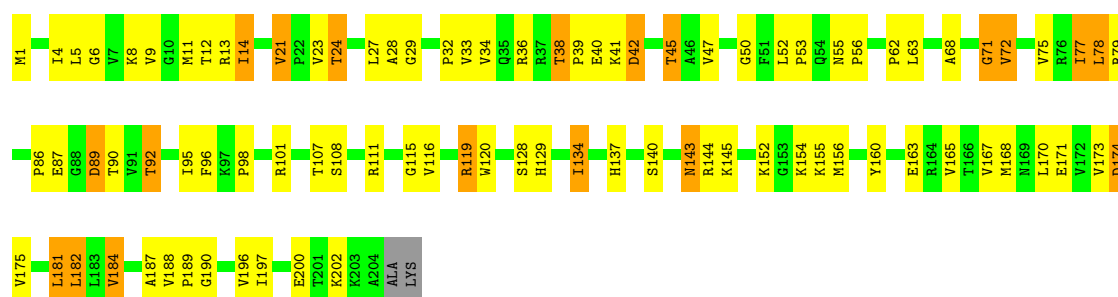
• Molecule 29: 50S ribosomal protein L3

Chain 21: 



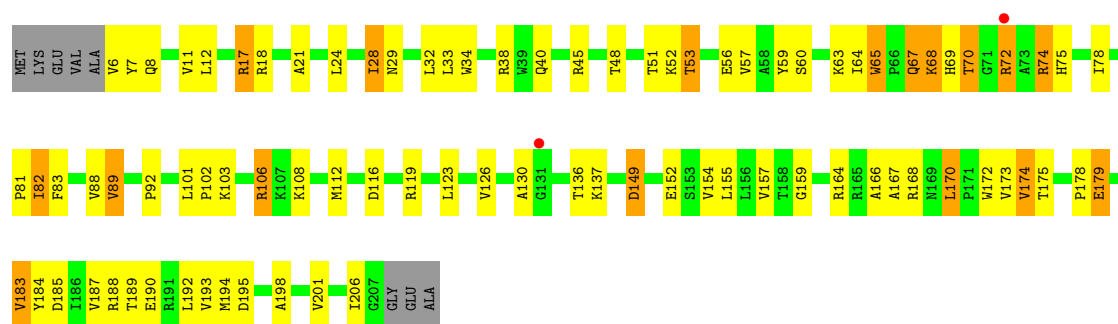
- Molecule 29: 50S ribosomal protein L3

Chain 29: 

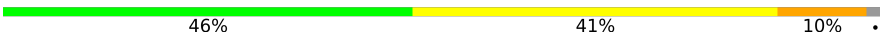


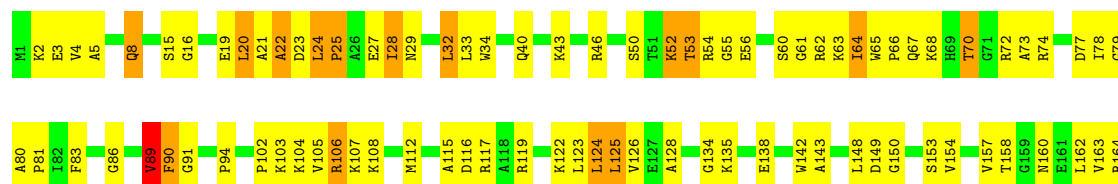
- Molecule 30: 50S ribosomal protein L4

Chain 31: 



- Molecule 30: 50S ribosomal protein L4

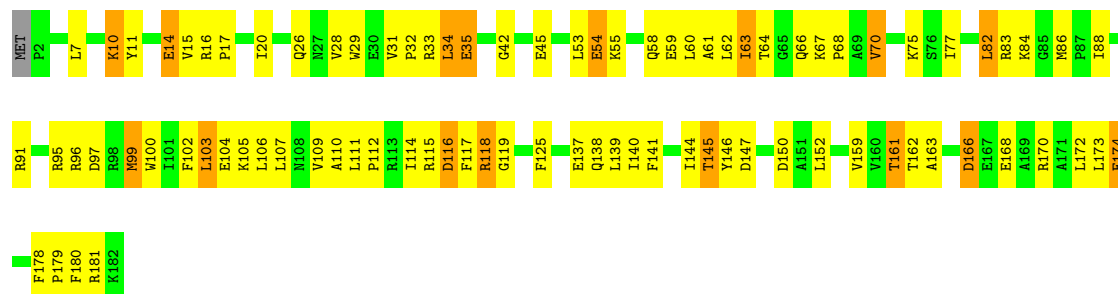
Chain 39: 





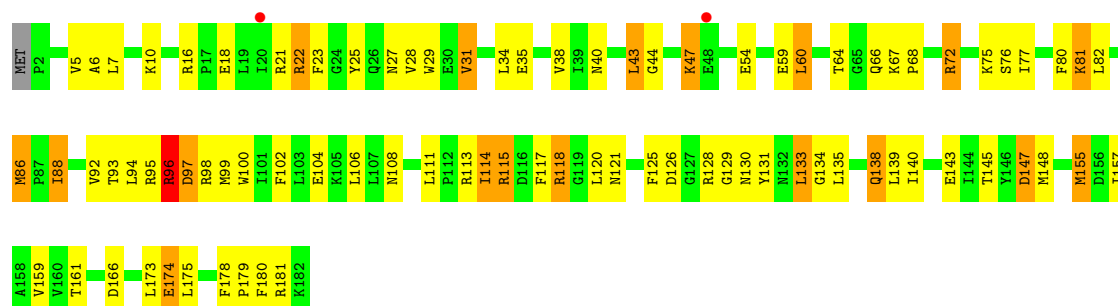
• Molecule 31: 50S ribosomal protein L5

Chain 41: 52% 39% 9%



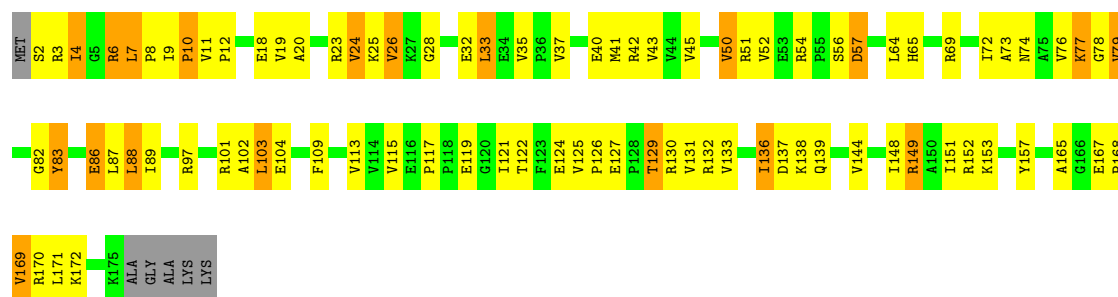
• Molecule 31: 50S ribosomal protein L5

Chain 49: 52% 37% 10%



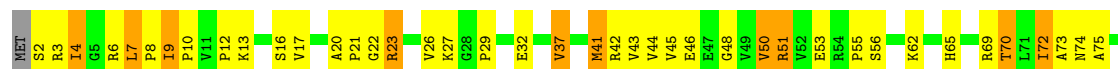
• Molecule 32: 50S ribosomal protein L6

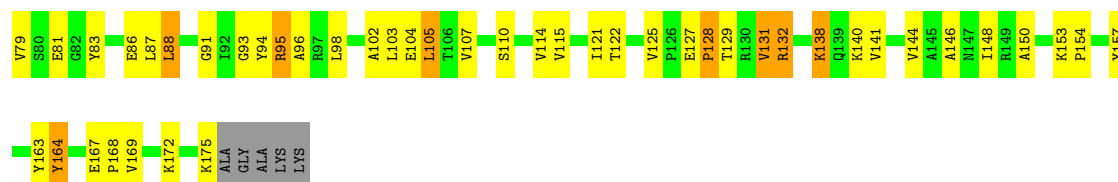
Chain 51: 48% 38% 11%



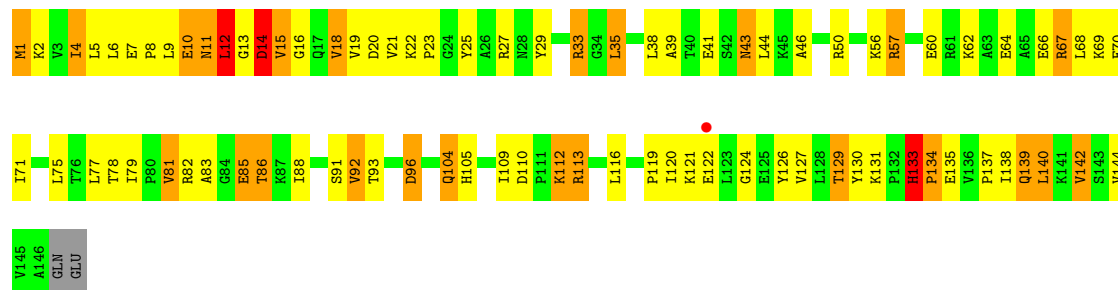
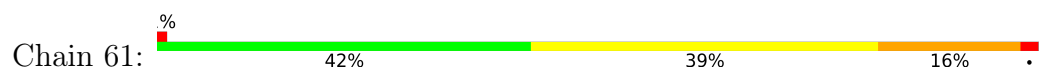
• Molecule 32: 50S ribosomal protein L6

Chain 59: 49% 38% 10%

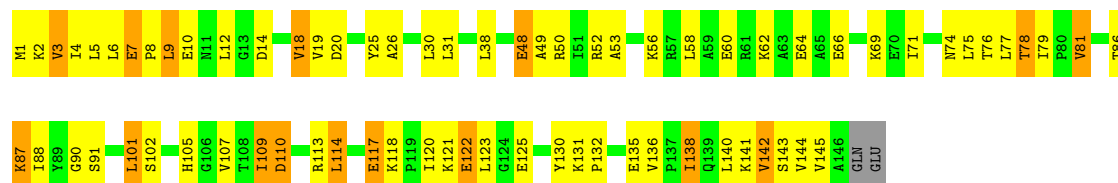




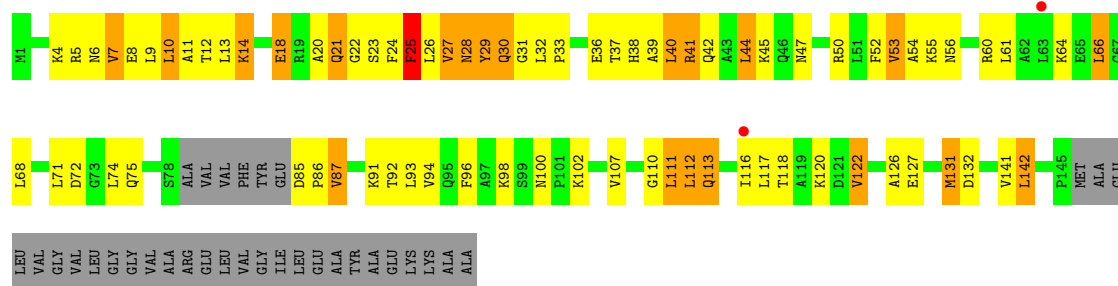
- Molecule 33: 50S ribosomal protein L9



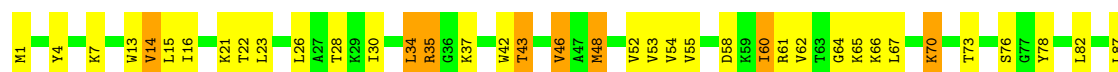
- Molecule 33: 50S ribosomal protein L9

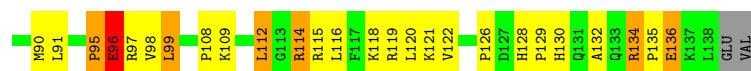


- Molecule 34: 50S ribosomal protein L10



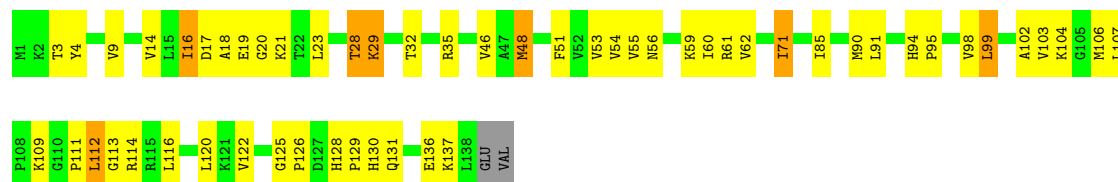
- Molecule 35: 50S ribosomal protein L13





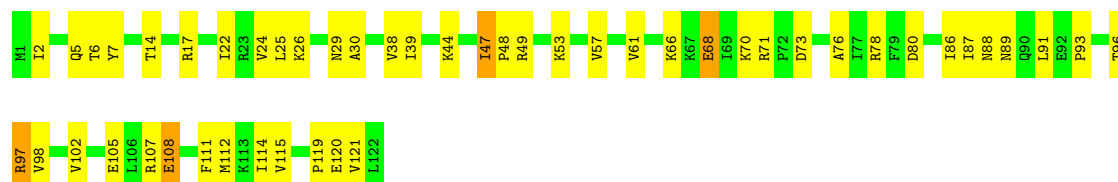
• Molecule 35: 50S ribosomal protein L13

Chain 15: 59% 34% 5% •



• Molecule 36: 50S ribosomal protein L14

Chain 68: 60% 37% •



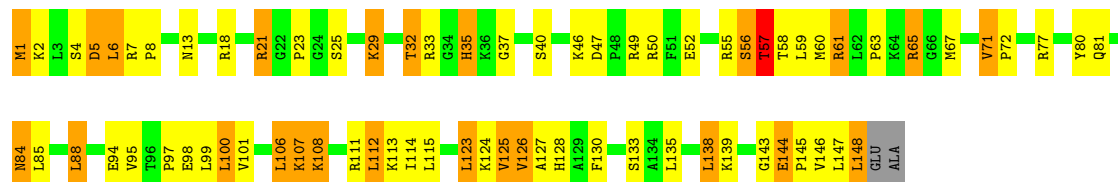
• Molecule 36: 50S ribosomal protein L14

Chain 25: 2% 59% 31% 10%



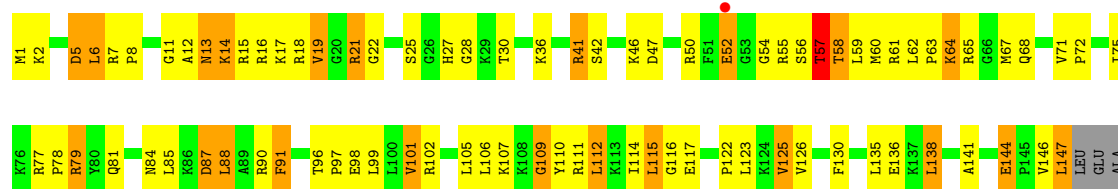
• Molecule 37: 50S ribosomal protein L15

Chain 78: 50% 32% 16% ••



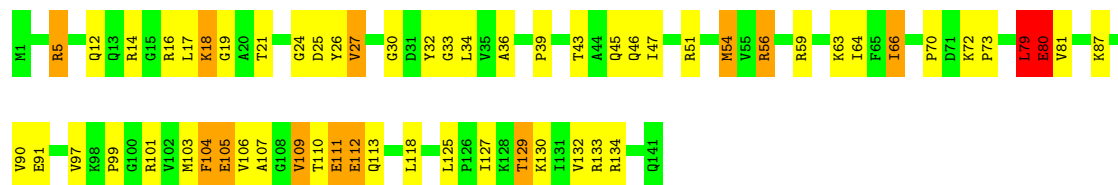
• Molecule 37: 50S ribosomal protein L15

Chain 35: 42% 41% 15% ••



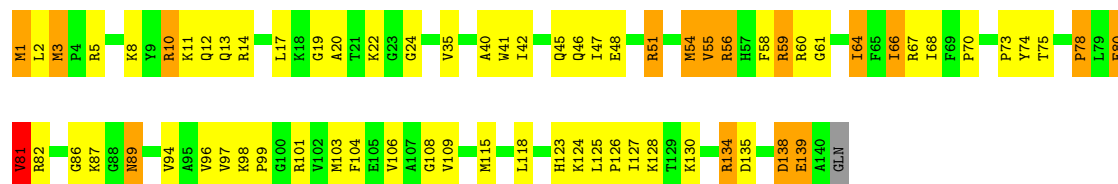
• Molecule 38: 50S ribosomal protein L16

Chain 88: 58% 32% 9%



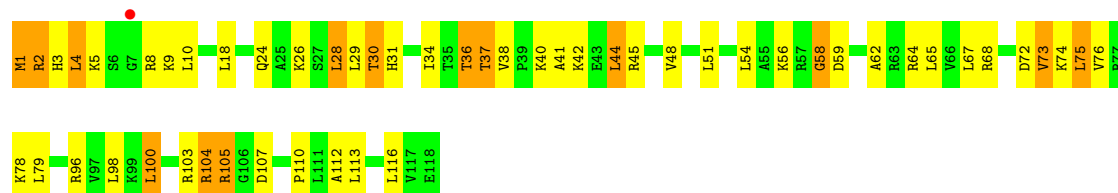
• Molecule 38: 50S ribosomal protein L16

Chain 45: 50% 38% 11%



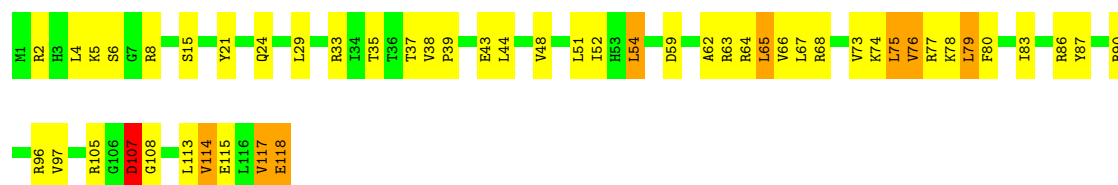
• Molecule 39: 50S ribosomal protein L17

Chain 98: 55% 33% 12%



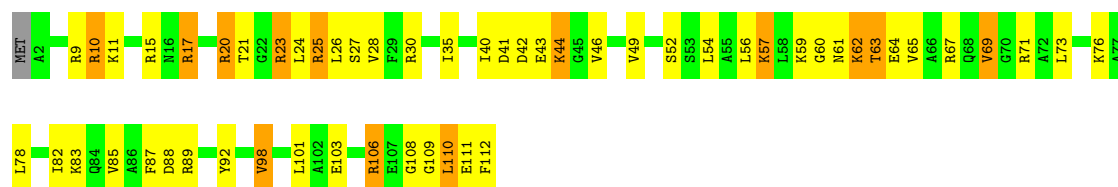
• Molecule 39: 50S ribosomal protein L17

Chain 55: 58% 35% 7%

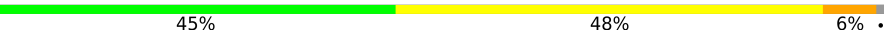


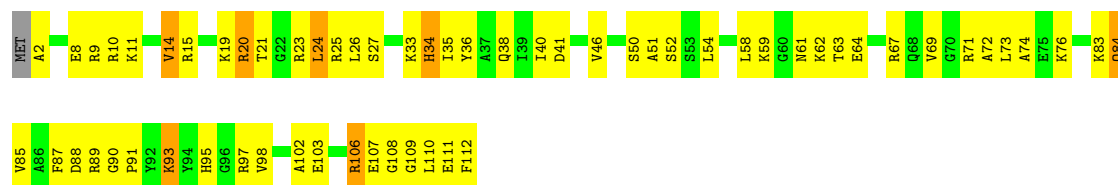
• Molecule 40: 50S ribosomal protein L18

Chain A8: 

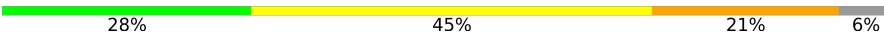


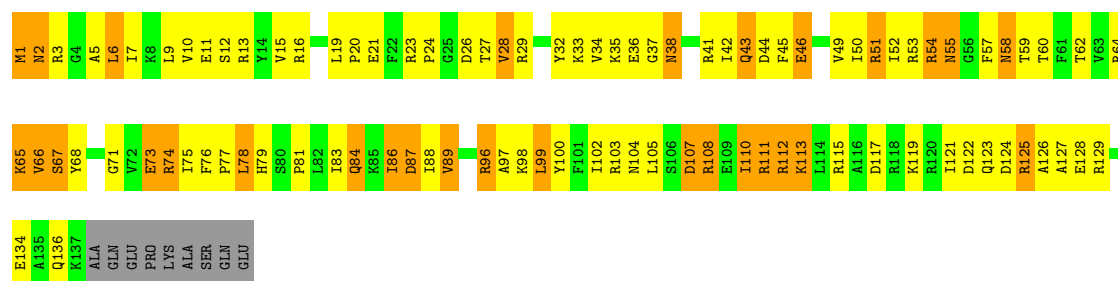
- Molecule 40: 50S ribosomal protein L18

Chain 65: 



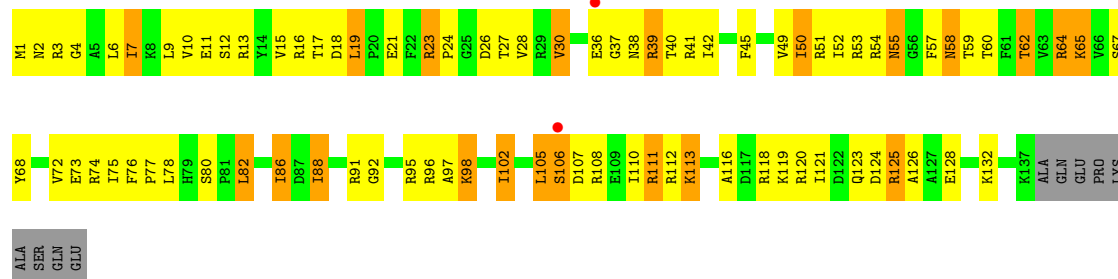
- Molecule 41: 50S ribosomal protein L19

Chain B8: 



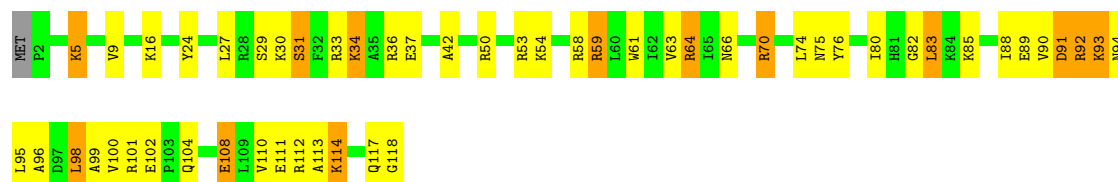
- Molecule 41: 50S ribosomal protein L19

Chain 75: 



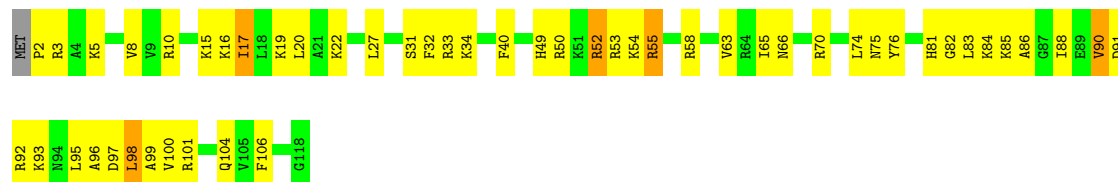
- Molecule 42: 50S ribosomal protein L20

Chain C8: 



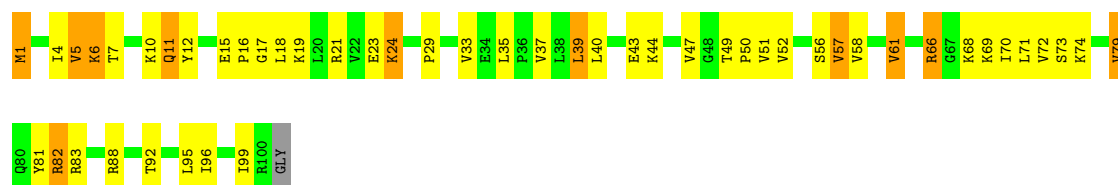
- Molecule 42: 50S ribosomal protein L20

Chain 85: 56% 39% . .



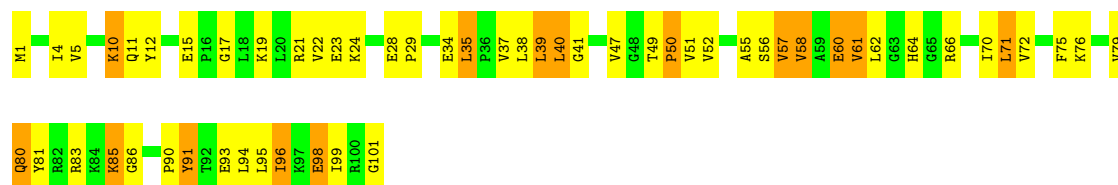
- Molecule 43: 50S ribosomal protein L21

Chain D8: 50% 39% 11% .



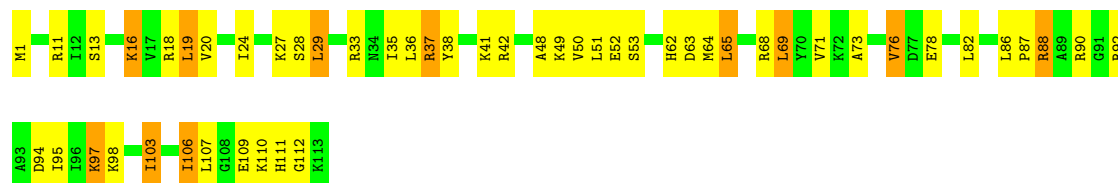
- Molecule 43: 50S ribosomal protein L21

Chain 95: 45% 41% 15%



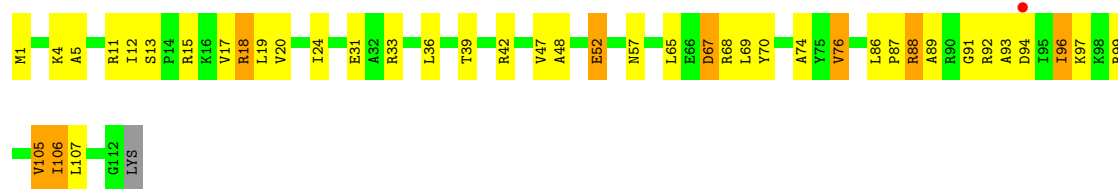
- Molecule 44: 50S ribosomal protein L22

Chain E8: 55% 35% 10%



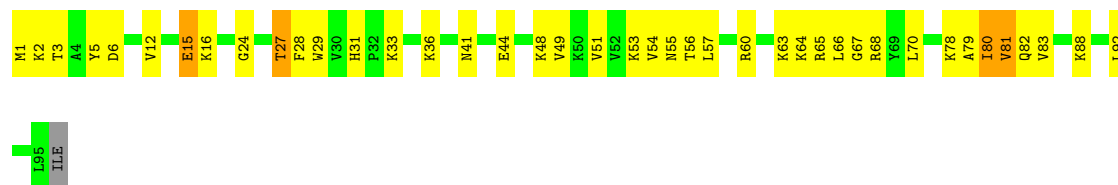
- Molecule 44: 50S ribosomal protein L22

Chain A5: 62% 30% 7% .



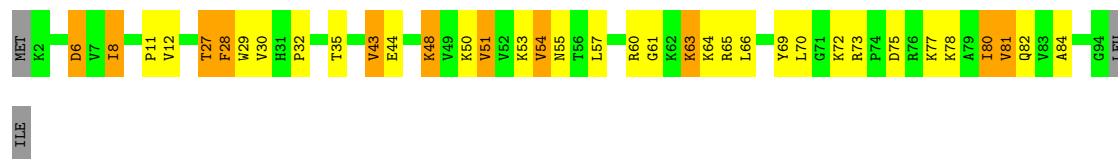
- Molecule 45: 50S ribosomal protein L23

Chain F8: ..



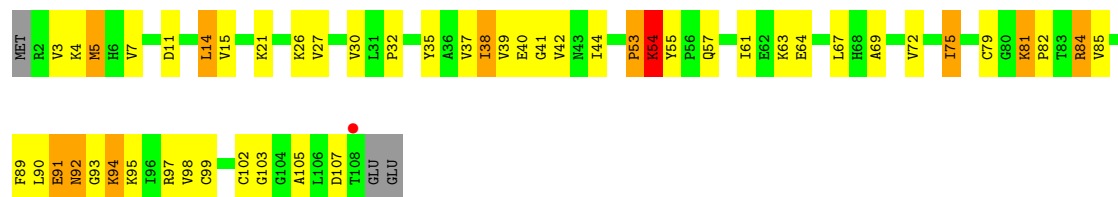
- Molecule 45: 50S ribosomal protein L23

Chain B5: ..



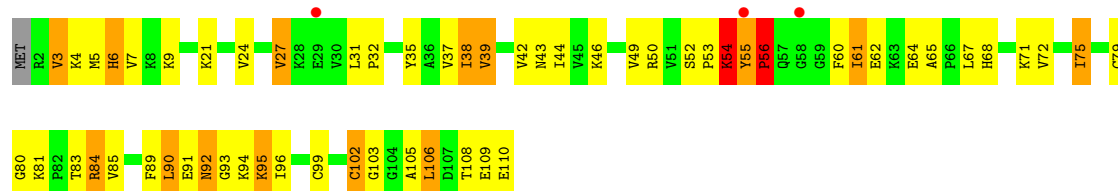
- Molecule 46: 50S ribosomal protein L24

Chain G8: ..

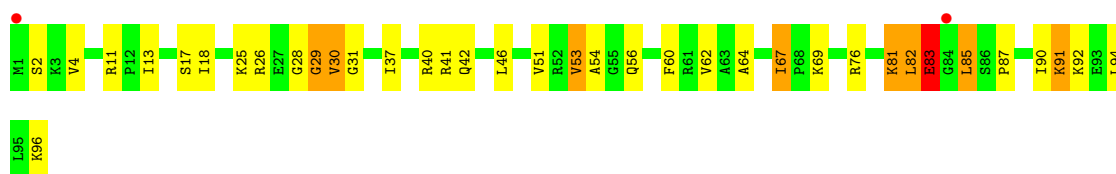


- Molecule 46: 50S ribosomal protein L24

Chain C5: ..



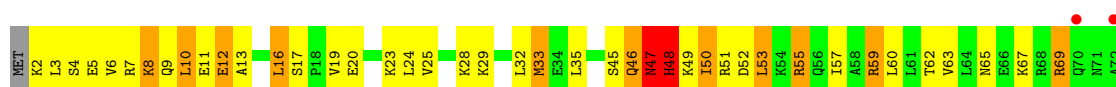
- Molecule 47: 50S ribosomal protein L25



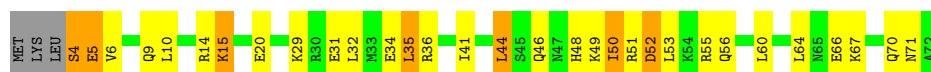
- Molecule 49: 50S ribosomal protein L28



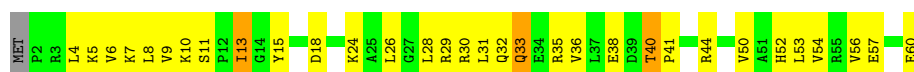
- Molecule 50: 50S ribosomal protein L29



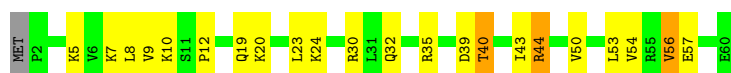
- Molecule 50: 50S ribosomal protein L29



- Molecule 51: 50S ribosomal protein L30



- Molecule 51: 50S ribosomal protein L30

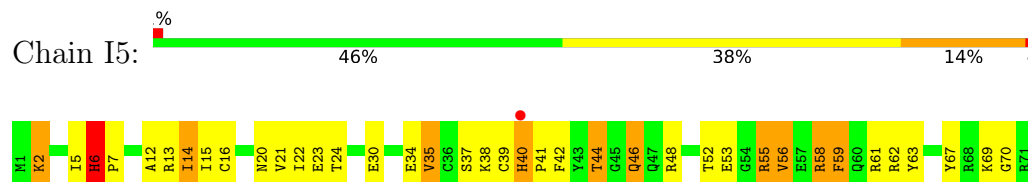


- Molecule 52: 50S ribosomal protein L31

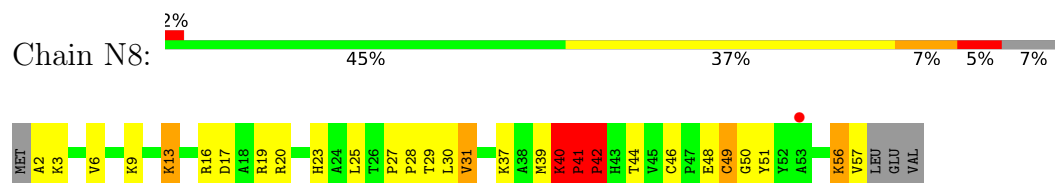


GLY
ARG

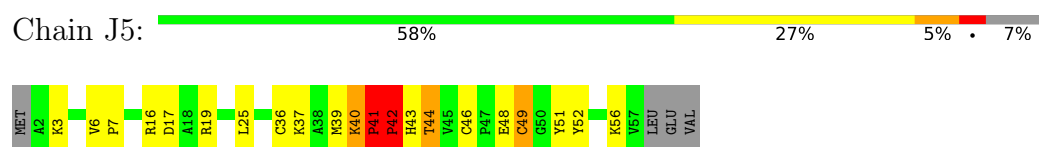
- Molecule 52: 50S ribosomal protein L31



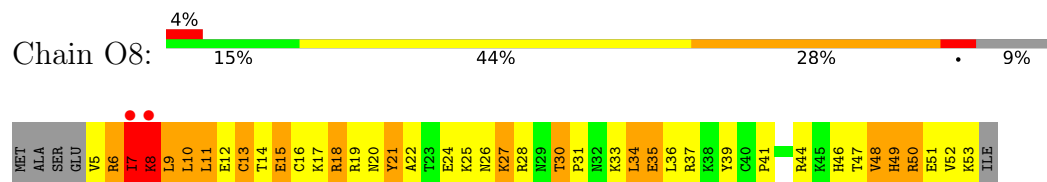
- Molecule 53: 50S ribosomal protein L32



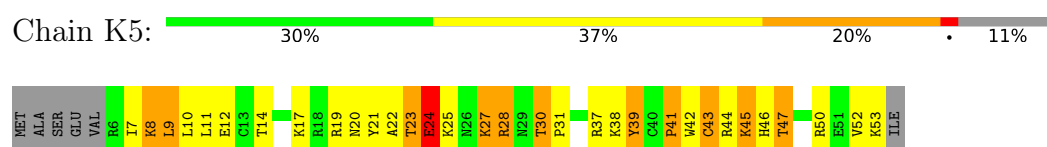
- Molecule 53: 50S ribosomal protein L32



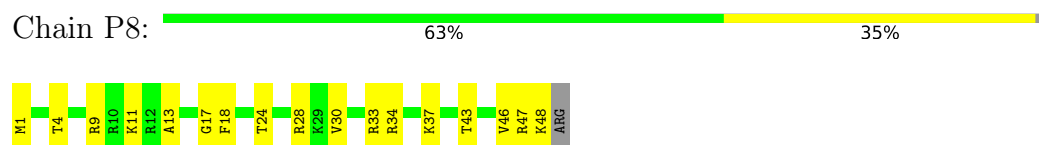
- Molecule 54: 50S ribosomal protein L33



- Molecule 54: 50S ribosomal protein L33



- Molecule 55: 50S ribosomal protein L34



- Molecule 55: 50S ribosomal protein L34

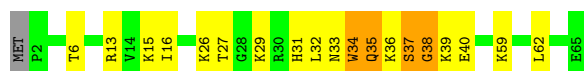




- Molecule 56: 50S ribosomal protein L35



- Molecule 56: 50S ribosomal protein L35



- Molecule 57: E. coli tRNA^{fMet}



4 Data and refinement statistics

Property	Value	Source
Space group	P 21 21 21	Depositor
Cell constants a, b, c, α , β , γ	210.46Å 452.18Å 626.12Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	226.09 – 3.50 226.09 – 3.50	Depositor EDS
% Data completeness (in resolution range)	100.0 (226.09-3.50) 100.0 (226.09-3.50)	Depositor EDS
R_{merge}	0.28	Depositor
R_{sym}	0.18	Depositor
$\langle I/\sigma(I) \rangle$ ¹	1.34 (at 3.01Å)	Xtriage
Refinement program	PHENIX 1.13_2998	Depositor
R, R_{free}	0.187 , 0.251 0.198 , 0.254	Depositor DCC
R_{free} test set	22133 reflections (1.88%)	wwPDB-VP
Wilson B-factor (Å ²)	71.7	Xtriage
Anisotropy	0.113	Xtriage
Bulk solvent k_{sol} (e/Å ³), B_{sol} (Å ²)	0.26 , 60.6	EDS
L-test for twinning ²	$\langle L \rangle = 0.40$, $\langle L^2 \rangle = 0.22$	Xtriage
Estimated twinning fraction	No twinning to report.	Xtriage
F_o, F_c correlation	0.95	EDS
Total number of atoms	300507	wwPDB-VP
Average B, all atoms (Å ²)	96.0	wwPDB-VP

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 1.33% 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: OMC, 2MG, OMG, PSU, 4OC, UR3, ZN, K, 5MC, MG, 4SU, 5MU, 2MA, OMU, M2G, 0TD, H2U, SF4, MA6, 7MG

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

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
1	13	0.17	1/35952 (0.0%)	0.39	0/56107
1	1G	0.16	1/36044 (0.0%)	0.40	3/56250 (0.0%)
2	12	0.19	0/1950	0.47	0/2630
2	1E	0.18	0/1959	0.44	0/2642
3	22	0.19	0/1636	0.50	0/2205
3	2E	0.18	0/1629	0.45	0/2195
4	32	0.17	0/1732	0.45	0/2318
4	3E	0.21	0/1732	0.45	0/2318
5	42	0.21	0/1195	0.45	0/1609
5	4E	0.16	0/1171	0.41	0/1576
6	52	0.15	0/855	0.40	0/1154
6	5E	0.15	0/855	0.39	0/1154
7	62	0.16	0/1235	0.44	0/1654
7	6E	0.14	0/1275	0.38	0/1709
8	72	0.16	0/1135	0.39	0/1527
8	7E	0.18	0/1135	0.46	0/1527
9	82	0.22	0/1021	0.64	0/1371
9	8E	0.17	0/1002	0.47	0/1346
10	1A	0.18	0/814	0.50	0/1095
10	1I	0.16	0/814	0.43	0/1095
11	2A	0.18	0/904	0.46	0/1219
11	2I	0.16	0/879	0.39	0/1188
12	3A	0.17	0/982	0.53	0/1313
12	3I	0.21	0/982	0.52	0/1313
13	4A	0.20	0/947	0.58	0/1270
13	4I	0.21	0/956	0.54	1/1281 (0.1%)
14	5A	0.16	0/494	0.49	0/657
14	5I	0.20	0/500	0.63	2/664 (0.3%)
15	6A	0.15	0/744	0.38	0/992
15	6I	0.16	0/744	0.35	0/992
16	7A	0.18	0/721	0.45	0/970

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
16	7I	0.17	0/716	0.44	0/963
17	8A	0.18	0/832	0.39	0/1113
17	8I	0.15	0/847	0.40	0/1131
18	9A	0.17	0/589	0.43	0/782
18	9I	0.17	0/578	0.42	0/768
19	AA	0.19	0/679	0.51	0/913
19	AI	0.20	0/703	0.59	0/945
20	BA	0.17	0/778	0.50	1/1028 (0.1%)
20	BI	0.18	0/768	0.60	3/1014 (0.3%)
21	1B	0.20	0/221	0.47	0/288
21	1F	0.17	0/221	0.43	0/288
22	1K	0.14	0/1547	0.38	0/2411
22	3K	0.13	0/1832	0.32	0/2855
22	3L	0.14	0/1832	0.33	0/2855
23	2K	0.19	0/1696	0.33	0/2644
24	4K	0.16	0/421	0.37	0/655
24	4L	0.14	0/523	0.38	0/815
25	14	0.18	0/69194	0.40	3/108015 (0.0%)
25	1H	0.20	0/69191	0.42	4/108008 (0.0%)
26	16	0.17	0/2878	0.38	0/4490
26	1J	0.15	0/2928	0.37	0/4568
27	7I	0.19	0/1072	0.44	0/1447
28	11	0.23	0/2175	0.55	0/2933
28	19	0.20	0/2173	0.49	2/2928 (0.1%)
29	21	0.20	0/1592	0.46	0/2149
29	29	0.19	0/1592	0.47	0/2149
30	31	0.22	0/1620	0.48	0/2194
30	39	0.21	0/1654	0.55	0/2239
31	41	0.18	0/1498	0.50	0/2016
31	49	0.18	0/1498	0.50	0/2016
32	51	0.20	0/1362	0.54	0/1841
32	59	0.27	1/1362 (0.1%)	0.47	0/1841
33	61	0.21	0/1151	0.64	0/1558
33	69	0.20	0/1151	0.56	0/1558
34	38	0.23	0/1069	0.66	0/1444
35	15	0.17	0/1131	0.42	0/1525
35	58	0.21	0/1131	0.51	0/1525
36	25	0.19	0/942	0.48	0/1269
36	68	0.19	0/942	0.47	0/1269
37	35	0.25	0/1139	0.70	0/1514
37	78	0.25	0/1147	0.73	1/1525 (0.1%)
38	45	0.22	0/1133	0.53	0/1515
38	88	0.23	0/1142	0.60	1/1527 (0.1%)

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
39	55	0.26	1/981 (0.1%)	0.50	0/1312
39	98	0.20	0/981	0.55	0/1312
40	65	0.21	0/891	0.53	0/1187
40	A8	0.22	0/891	0.63	0/1187
41	75	0.50	1/1155 (0.1%)	0.72	0/1542
41	B8	0.46	0/1155	0.68	0/1542
42	85	0.22	0/981	0.54	0/1306
42	C8	0.20	0/981	0.45	0/1306
43	95	0.24	0/789	0.61	0/1057
43	D8	0.22	0/785	0.52	0/1052
44	A5	0.19	0/901	0.47	0/1209
44	E8	0.22	0/910	0.51	0/1220
45	B5	0.20	0/744	0.45	0/1000
45	F8	0.20	0/761	0.44	0/1021
46	C5	0.25	0/838	0.66	0/1121
46	G8	0.21	0/818	0.60	0/1094
47	D5	0.26	0/1435	0.69	2/1947 (0.1%)
47	H8	0.23	0/1420	0.62	0/1925
48	E5	0.19	0/631	0.51	0/841
48	I8	0.19	0/619	0.49	0/825
49	F5	0.21	0/728	0.49	0/967
49	J8	0.37	1/761 (0.1%)	0.60	0/1010
50	G5	0.18	0/582	0.45	0/771
50	K8	0.19	0/592	0.55	0/784
51	H5	0.17	0/473	0.44	0/635
51	L8	0.19	0/473	0.40	0/635
52	I5	0.26	0/593	0.71	0/795
52	M8	0.24	0/545	0.75	2/733 (0.3%)
53	J5	0.22	0/448	0.49	0/606
53	N8	0.33	1/448 (0.2%)	0.57	0/606
54	K5	0.29	0/424	1.08	6/565 (1.1%)
54	O8	0.30	0/431	0.83	2/575 (0.3%)
55	L5	0.20	0/426	0.45	0/561
55	P8	0.24	0/426	0.48	0/561
56	M5	0.18	0/514	0.45	0/679
56	Q8	0.20	0/514	0.41	0/679
57	2L	0.18	0/1718	0.35	0/2678
All	All	0.19	7/321407 (0.0%)	0.44	33/480718 (0.0%)

Chiral center outliers are detected by calculating the chiral volume of a chiral center and verifying if the center is modelled as a planar moiety or with the opposite hand. A planarity outlier is detected by checking planarity of atoms in a peptide group, atoms in a mainchain group or atoms of a sidechain that are expected to be planar.

Mol	Chain	#Chirality outliers	#Planarity outliers
2	12	0	4
4	32	0	1
7	6E	0	1
12	3A	0	1
12	3I	0	2
13	4A	0	4
19	AA	0	1
19	AI	0	3
20	BA	0	1
20	BI	0	1
28	11	0	1
29	21	0	2
29	29	0	1
31	41	0	1
31	49	0	1
32	51	0	2
33	61	0	4
33	69	0	1
34	38	0	4
37	35	0	3
37	78	0	3
38	45	0	3
38	88	0	1
41	B8	0	1
42	85	0	1
43	95	0	1
45	B5	0	1
46	G8	0	2
47	D5	0	6
47	H8	0	4
48	E5	0	1
49	J8	0	2
50	K8	0	1
52	I5	0	1
52	M8	0	4
53	J5	0	2
53	N8	0	3
54	K5	0	4
54	O8	0	1
All	All	0	81

All (7) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
41	75	106	SER	CA-C	8.11	1.63	1.52
32	59	167	GLU	C-O	-6.74	1.20	1.23
49	J8	67	ILE	CA-CB	6.60	1.57	1.54
1	1G	1498	UR3	O3'-P	5.83	1.62	1.56
1	13	1498	UR3	O3'-P	5.79	1.62	1.56
53	N8	27	PRO	CA-C	-5.08	1.48	1.52
39	55	38	VAL	CA-CB	5.02	1.56	1.54

All (33) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
54	K5	30	THR	CA-C-N	8.01	146.22	127.00
54	K5	30	THR	C-N-CA	8.01	146.22	127.00
54	K5	30	THR	C-N-CD	-7.70	103.66	120.60
37	78	35	HIS	N-CA-C	7.30	121.14	111.28
20	BA	6	PRO	N-CA-CB	6.75	110.46	103.38
20	BI	6	PRO	N-CA-CB	6.63	110.29	103.00
54	K5	20	ASN	CA-C-N	6.59	131.13	121.31
54	K5	20	ASN	C-N-CA	6.59	131.13	121.31
13	4I	84	ILE	N-CA-C	-6.29	106.67	112.96
38	88	105	GLU	N-CA-C	5.72	117.61	110.91
54	O8	7	ILE	CA-C-N	5.70	132.42	121.54
54	O8	7	ILE	C-N-CA	5.70	132.42	121.54
20	BI	72	LEU	CA-C-N	5.61	132.25	121.54
20	BI	72	LEU	C-N-CA	5.61	132.25	121.54
28	19	272	ALA	CA-C-N	5.60	132.24	121.54
28	19	272	ALA	C-N-CA	5.60	132.24	121.54
14	5I	15	LYS	CA-C-N	5.40	131.86	121.54
14	5I	15	LYS	C-N-CA	5.40	131.86	121.54
1	1G	1498	UR3	P-O3'-C3'	5.39	126.16	119.70
47	D5	156	LYS	CA-C-N	5.37	134.89	121.80
47	D5	156	LYS	C-N-CA	5.37	134.89	121.80
25	1H	1266	G	C2'-C3'-O3'	5.26	117.39	109.50
25	14	1694	C	C4'-C3'-O3'	5.25	117.27	109.40
25	1H	271(B)	G	P-O3'-C3'	5.21	125.95	119.70
1	1G	1004	A	P-O3'-C3'	5.19	127.99	120.20
54	K5	21	TYR	N-CA-C	5.19	117.62	110.35
25	1H	1266	G	P-O3'-C3'	5.19	127.98	120.20
1	1G	1004	A	C2'-C3'-O3'	5.14	117.22	109.50
25	1H	1992	G	C2'-C3'-O3'	5.09	117.14	109.50
25	14	1992	G	C2'-C3'-O3'	5.09	117.14	109.50
52	M8	39	CYS	CA-C-N	5.05	134.13	121.80
52	M8	39	CYS	C-N-CA	5.05	134.13	121.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
25	14	1694	C	P-O3'-C3'	5.03	127.74	120.20

There are no chirality outliers.

All (81) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
28	11	239	ARG	Peptide
2	12	16	HIS	Peptide
2	12	199	TYR	Peptide
2	12	21	ARG	Peptide
2	12	230	VAL	Peptide
29	21	70	ALA	Peptide
29	21	72	VAL	Peptide
29	29	72	VAL	Peptide
4	32	88	VAL	Peptide
37	35	109	GLY	Peptide
37	35	115	LEU	Peptide
37	35	57	THR	Peptide
34	38	107	VAL	Peptide
34	38	25	PHE	Peptide
34	38	28	ASN	Peptide
34	38	53	VAL	Peptide
12	3A	101	VAL	Peptide
12	3I	23	ALA	Peptide
12	3I	44	LYS	Peptide
31	41	116	ASP	Peptide
38	45	134	ARG	Peptide
38	45	80	GLU	Peptide
38	45	81	VAL	Peptide
31	49	96	ARG	Peptide
13	4A	106	ASN	Peptide
13	4A	43	THR	Peptide
13	4A	62	ASN	Peptide
13	4A	67	GLU	Peptide
32	51	10	PRO	Peptide
32	51	82	GLY	Peptide
33	61	11	ASN	Peptide
33	61	13	GLY	Peptide
33	61	133	HIS	Peptide
33	61	14	ASP	Peptide
33	69	142	VAL	Peptide
7	6E	5	ARG	Peptide

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Mol	Chain	Res	Type	Group
37	78	35	HIS	Peptide
37	78	56	SER	Peptide
37	78	57	THR	Peptide
42	85	90	VAL	Peptide
38	88	79	LEU	Peptide
43	95	98	GLU	Peptide
19	AA	69	HIS	Peptide
19	AI	41	VAL	Peptide
19	AI	5	LEU	Peptide
19	AI	81	ARG	Peptide
45	B5	60	ARG	Peptide
41	B8	113	LYS	Peptide
20	BA	69	GLY	Peptide
20	BI	73	HIS	Peptide
47	D5	140	ASP	Peptide
47	D5	145	GLU	Peptide
47	D5	152	ALA	Peptide
47	D5	155	LEU	Peptide
47	D5	170	THR	Peptide
47	D5	52	SER	Peptide
48	E5	49	LYS	Peptide
46	G8	53	PRO	Peptide
46	G8	54	LYS	Peptide
47	H8	109	ALA	Peptide
47	H8	140	ASP	Peptide
47	H8	155	LEU	Peptide
47	H8	52	SER	Peptide
52	I5	39	CYS	Peptide
53	J5	41	PRO	Peptide
53	J5	42	PRO	Peptide
49	J8	29	GLY	Peptide
49	J8	53	VAL	Peptide
54	K5	24	GLU	Peptide
54	K5	27	LYS	Peptide
54	K5	30	THR	Peptide
54	K5	46	HIS	Peptide
50	K8	46	GLN	Peptide
52	M8	22	ILE	Peptide
52	M8	38	LYS	Peptide
52	M8	4	GLY	Peptide
52	M8	40	HIS	Peptide
53	N8	40	LYS	Peptide

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Mol	Chain	Res	Type	Group
53	N8	41	PRO	Peptide
53	N8	42	PRO	Peptide
54	O8	8	LYS	Peptide

5.2 Too-close contacts

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	13	32387	0	16370	693	0
1	1G	32470	0	16411	755	0
2	12	1915	0	1969	85	0
2	1E	1924	0	1975	71	0
3	22	1612	0	1677	63	0
3	2E	1605	0	1668	52	0
4	32	1702	0	1766	64	0
4	3E	1702	0	1766	68	0
5	42	1178	0	1234	41	0
5	4E	1155	0	1213	39	0
6	52	842	0	857	28	0
6	5E	842	0	857	21	0
7	62	1217	0	1256	55	0
7	6E	1256	0	1296	36	0
8	72	1115	0	1177	42	0
8	7E	1115	0	1177	41	0
9	82	1002	0	1025	79	0
9	8E	983	0	1006	49	0
10	1A	801	0	849	35	0
10	1I	801	0	849	43	0
11	2A	889	0	916	34	0
11	2I	864	0	878	24	0
12	3A	977	0	1061	36	0
12	3I	977	0	1061	30	0
13	4A	937	0	995	60	0
13	4I	946	0	1008	43	0
14	5A	485	0	519	25	0
14	5I	491	0	530	32	0
15	6A	733	0	771	23	0
15	6I	733	0	771	19	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
16	7A	705	0	725	20	0
16	7I	700	0	720	27	0
17	8A	819	0	880	26	0
17	8I	834	0	904	27	0
18	9A	584	0	657	27	0
18	9I	573	0	644	29	0
19	AA	665	0	684	61	0
19	AI	688	0	710	43	0
20	BA	776	0	857	32	0
20	BI	766	0	854	30	0
21	1B	217	0	234	14	0
21	1F	217	0	234	13	0
22	1K	1385	0	705	38	0
22	3K	1640	0	837	30	0
22	3L	1640	0	837	36	0
23	2K	1643	0	847	17	0
24	4K	373	0	186	8	0
24	4L	463	0	230	7	0
25	14	62013	0	31255	1097	0
25	1H	62010	0	31252	1057	0
26	16	2573	0	1305	35	0
26	1J	2617	0	1328	78	0
27	7I	1049	0	1071	52	0
28	11	2125	0	2199	64	0
28	19	2124	0	2197	91	0
29	21	1559	0	1618	49	0
29	29	1559	0	1617	58	0
30	31	1585	0	1632	73	0
30	39	1619	0	1674	82	0
31	41	1473	0	1535	64	0
31	49	1473	0	1535	65	0
32	51	1336	0	1418	58	0
32	59	1336	0	1418	48	0
33	61	1136	0	1223	43	0
33	69	1136	0	1223	40	0
34	38	1056	0	1119	50	0
35	15	1104	0	1180	33	0
35	58	1104	0	1180	42	0
36	25	932	0	996	33	0
36	68	932	0	996	33	0
37	35	1122	0	1206	78	0
37	78	1130	0	1217	65	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
38	45	1112	0	1171	57	0
38	88	1121	0	1179	35	0
39	55	967	0	1033	43	0
39	98	967	0	1033	32	0
40	65	881	0	943	47	0
40	A8	881	0	943	37	0
41	75	1141	0	1202	76	0
41	B8	1141	0	1202	89	0
42	85	963	0	1022	52	0
42	C8	963	0	1021	44	0
43	95	778	0	852	46	0
43	D8	774	0	849	32	0
44	A5	890	0	951	22	0
44	E8	899	0	964	30	0
45	B5	730	0	780	27	0
45	F8	747	0	805	27	0
46	C5	825	0	898	47	0
46	G8	805	0	881	27	0
47	D5	1404	0	1437	81	0
47	H8	1390	0	1421	60	0
48	E5	623	0	645	30	0
48	I8	611	0	631	23	0
49	F5	721	0	791	24	0
49	J8	754	0	838	25	0
50	G5	580	0	629	16	0
50	K8	590	0	643	25	0
51	H5	468	0	518	12	0
51	L8	468	0	518	15	0
52	I5	580	0	577	30	0
52	M8	533	0	526	33	0
53	J5	434	0	454	15	0
53	N8	434	0	454	29	0
54	K5	417	0	441	20	0
54	O8	424	0	450	36	0
55	L5	418	0	467	17	0
55	P8	418	0	467	10	0
56	M5	506	0	567	15	0
56	Q8	506	0	567	18	0
57	2L	1643	0	845	25	0
58	11	1	0	0	0	0
58	13	38	0	0	0	0
58	14	94	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
58	16	2	0	0	0	0
58	19	1	0	0	0	0
58	1G	25	0	0	0	0
58	1H	122	0	0	0	0
58	21	1	0	0	0	0
58	29	2	0	0	0	0
58	31	2	0	0	0	0
58	32	1	0	0	0	0
58	39	1	0	0	0	0
58	3A	1	0	0	0	0
58	41	2	0	0	0	0
58	52	1	0	0	0	0
58	5E	1	0	0	0	0
58	5I	1	0	0	0	0
58	8I	1	0	0	0	0
58	BA	1	0	0	0	0
59	13	111	0	0	0	0
59	14	312	0	0	0	0
59	16	8	0	0	0	0
59	1G	102	0	0	0	0
59	1H	405	0	0	0	0
59	1J	5	0	0	0	0
59	21	2	0	0	0	0
59	29	2	0	0	0	0
59	2K	1	0	0	0	0
59	2L	2	0	0	0	0
59	32	1	0	0	0	0
59	3E	1	0	0	0	0
59	41	1	0	0	0	0
59	4I	1	0	0	0	0
59	55	1	0	0	0	0
59	78	1	0	0	0	0
59	7I	1	0	0	0	0
59	8E	1	0	0	0	0
59	BI	2	0	0	0	0
59	D8	3	0	0	0	0
59	E5	1	0	0	0	0
59	G8	1	0	0	0	0
59	I8	2	0	0	0	0
59	J8	1	0	0	0	0
59	N8	1	0	0	0	0
59	Q8	1	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
60	32	8	0	0	2	0
60	3E	8	0	0	1	0
61	5A	1	0	0	0	0
61	5I	1	0	0	0	0
62	11	10	0	0	1	0
62	13	190	0	0	11	0
62	14	681	0	0	72	0
62	16	12	0	0	2	0
62	19	12	0	0	4	0
62	1G	268	0	0	18	0
62	1H	983	0	0	113	0
62	1J	16	0	0	4	0
62	21	5	0	0	0	0
62	29	3	0	0	0	0
62	2L	6	0	0	0	0
62	31	6	0	0	0	0
62	32	4	0	0	0	0
62	35	3	0	0	1	0
62	39	3	0	0	0	0
62	3A	1	0	0	0	0
62	3E	2	0	0	0	0
62	3I	1	0	0	0	0
62	3K	1	0	0	0	0
62	42	1	0	0	0	0
62	4I	2	0	0	0	0
62	4K	2	0	0	0	0
62	4L	4	0	0	0	0
62	55	2	0	0	0	0
62	58	1	0	0	0	0
62	5I	1	0	0	0	0
62	6I	1	0	0	0	0
62	75	1	0	0	0	0
62	78	6	0	0	1	0
62	7A	5	0	0	0	0
62	7I	3	0	0	1	0
62	85	1	0	0	0	0
62	98	1	0	0	0	0
62	B5	2	0	0	0	0
62	B8	2	0	0	0	0
62	BA	1	0	0	0	0
62	BI	4	0	0	1	0
62	C5	2	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
62	C8	2	0	0	0	0
62	F5	1	0	0	0	0
62	F8	2	0	0	0	0
62	G5	1	0	0	0	0
62	G8	1	0	0	0	0
62	I8	5	0	0	0	0
62	J8	3	0	0	0	0
62	L5	1	0	0	0	0
62	M5	1	0	0	0	0
62	Q8	1	0	0	0	0
All	All	300507	0	201662	6882	0

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 14.

All (6882) 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)
28:19:182:LEU:H	28:19:272:ALA:HB3	1.30	0.97
1:13:1129:C:H5'	1:13:1130:A:H5'	1.47	0.96
25:1H:1053:C:H42	25:1H:1106:G:H1	1.12	0.95
25:14:2245:U:H5'	25:14:2246:G:H5'	1.47	0.95
1:1G:73:G:H1	1:1G:97:U:H3	1.10	0.94
25:14:1899:G:H22	25:14:1902:C:H41	1.12	0.93
25:1H:307:G:N2	25:1H:310:A:OP2	2.01	0.93
25:1H:2415:G:H4'	37:78:67:MET:H	1.34	0.93
25:1H:2701:C:H3'	25:1H:2702:U:H5''	1.52	0.92
57:2L:8:4SU:HN3	57:2L:14:A:H62	1.18	0.91
25:14:2720:U:H3	25:14:2873:A:H2	1.12	0.90
25:14:945:A:OP1	62:14:3510:HOH:O	1.91	0.89
1:1G:664:G:H22	1:1G:741:G:H1	1.20	0.89
25:1H:2068:U:H3	25:1H:2430:A:H2	1.22	0.88
25:1H:875:G:H4'	47:H8:170:THR:HG21	1.55	0.88
28:11:182:LEU:H	28:11:272:ALA:HB3	1.40	0.87
34:38:10:LEU:HB3	34:38:53:VAL:HG11	1.56	0.87
1:13:1139:G:N1	1:13:1143:G:O6	2.07	0.87
25:14:2701:C:H3'	25:14:2702:U:H5''	1.55	0.86
25:1H:993:G:OP1	42:C8:50:ARG:NH2	2.09	0.86
34:38:98:LYS:H	34:38:102:LYS:HD3	1.40	0.86
25:1H:219:G:OP2	62:1H:3614:HOH:O	1.92	0.86
57:2L:10:G:H22	57:2L:27:G:H1'	1.40	0.86

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:676:A:H8	25:14:2069:G:H21	1.23	0.86
25:14:2068:U:H3	25:14:2430:A:H2	1.21	0.86
1:1G:1109:C:OP1	3:22:176:HIS:ND1	2.09	0.85
28:19:8:PRO:HB3	28:19:14:ARG:HB3	1.57	0.85
47:H8:140:ASP:H	47:H8:155:LEU:HG	1.40	0.85
42:85:50:ARG:HH12	43:95:72:VAL:HG11	1.41	0.85
25:1H:2723:C:H5''	39:98:1:MET:HE2	1.59	0.85
25:14:397:G:N7	62:14:3523:HOH:O	2.09	0.85
25:1H:761:A:N7	62:1H:3638:HOH:O	2.09	0.85
53:N8:40:LYS:HG2	53:N8:46:CYS:HA	1.57	0.84
25:1H:1103:A:H5'	25:1H:1104:C:H5	1.43	0.84
4:3E:156:GLU:HB2	4:3E:159:ARG:HH11	1.43	0.84
26:16:73:A:OP2	62:16:301:HOH:O	1.96	0.84
33:69:79:ILE:H	33:69:142:VAL:HG21	1.42	0.84
1:1G:928:G:H1	1:1G:1389:C:H42	1.25	0.84
25:1H:987:G:O2'	25:1H:1000:A:N3	2.11	0.83
1:13:1502:A:H2	1:13:1505:G:H1	1.26	0.83
25:1H:2112:G:H21	25:1H:2113:U:H3	1.25	0.83
37:35:19:VAL:HG13	37:35:21:ARG:H	1.43	0.83
25:14:815:C:H42	25:14:1192:G:H1	1.23	0.83
25:1H:563:G:OP2	62:1H:3616:HOH:O	1.97	0.83
25:1H:1170:G:H1	25:1H:1179:C:H42	1.26	0.82
1:1G:448:A:H62	1:1G:486:U:H3	1.23	0.82
25:1H:2615:U:OP1	62:1H:3615:HOH:O	1.96	0.82
22:3L:13:C:O2	22:3L:23:G:N2	2.12	0.82
25:14:784:A:OP2	62:14:3511:HOH:O	1.96	0.82
25:14:819:A:OP2	25:14:1187:G:N2	2.12	0.82
25:14:660:G:H21	37:35:12:ALA:HB2	1.44	0.82
25:14:1783:A:OP1	62:14:3512:HOH:O	1.97	0.82
26:16:6:C:H42	26:16:114:G:H1	1.27	0.82
22:3L:3:C:H2'	22:3L:4:G:H8	1.45	0.82
1:13:1299:A:H2'	1:13:1301:U:H1'	1.61	0.81
31:41:97:ASP:H	31:41:100:TRP:HD1	1.27	0.81
25:1H:265:A:N6	25:1H:427:U:O2'	2.13	0.81
1:1G:40:C:H42	1:1G:402:G:H1	1.27	0.81
29:29:47:VAL:HG11	29:29:86:PRO:HD2	1.61	0.81
49:F5:75:GLU:HB3	49:F5:76:ARG:HH21	1.45	0.81
1:13:559:A:H4'	1:13:560:U:H3'	1.63	0.81
38:88:12:GLN:HB2	38:88:73:PRO:HD2	1.63	0.81
25:1H:1332:G:OP1	62:1H:3617:HOH:O	1.98	0.81
41:75:118:ARG:HA	41:75:121:ILE:HB	1.61	0.81

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:C8:95:LEU:HD13	43:D8:4:ILE:HG23	1.63	0.81
25:14:273(B):C:H42	25:14:363(D):G:H1	1.29	0.80
26:16:31:C:O2	26:16:53:A:N6	2.13	0.80
1:1G:1352:C:H42	1:1G:1370:G:H1	1.28	0.80
42:C8:92:ARG:HG3	42:C8:94:ASN:HB3	1.64	0.80
16:7I:72:ARG:HA	16:7I:75:ARG:HB3	1.62	0.80
1:1G:1149:C:OP2	9:82:9:ARG:NH1	2.14	0.80
1:13:1133:G:H1	1:13:1141:C:H42	1.28	0.80
2:1E:168:THR:HG22	2:1E:192:SER:HB2	1.62	0.80
25:1H:1359:A:H62	25:1H:1372:U:H3	1.27	0.80
1:1G:1359:C:O2'	1:1G:1361:G:N7	2.13	0.80
30:39:157:VAL:HB	30:39:194:MET:HB3	1.62	0.80
30:31:101:LEU:O	30:31:106:ARG:NH1	2.14	0.80
1:13:75:C:N4	1:13:95:G:O6	2.15	0.80
25:1H:676:A:H8	25:1H:2069:G:H21	1.24	0.79
42:85:92:ARG:HH22	43:95:10:LYS:HA	1.46	0.79
47:H8:48:PHE:HA	47:H8:51:ALA:HB3	1.65	0.79
4:3E:101:LEU:HD21	4:3E:121:VAL:HG11	1.63	0.79
1:1G:1252:A:H61	1:1G:1285:A:H61	1.29	0.79
25:14:768:G:O2'	25:14:1379:A:N6	2.15	0.79
25:1H:1439:A:OP1	62:1H:3618:HOH:O	1.98	0.79
33:61:62:LYS:HE3	33:61:134:PRO:HG2	1.65	0.79
50:G5:52:ASP:N	50:G5:52:ASP:OD1	2.14	0.79
52:M8:18:CYS:SG	52:M8:19:GLY:N	2.56	0.79
25:14:1062:G:H22	25:14:1076:C:H42	1.30	0.79
40:65:62:LYS:HB3	40:65:97:ARG:HD3	1.62	0.79
26:1J:36:C:N4	26:1J:49:C:O2	2.15	0.79
25:1H:1009:A:OP2	35:58:37:LYS:NZ	2.16	0.79
26:16:101:A:OP2	62:16:302:HOH:O	1.99	0.79
25:1H:769:G:N7	62:1H:3661:HOH:O	2.15	0.79
45:B5:63:LYS:H	45:B5:63:LYS:HE3	1.48	0.78
42:85:92:ARG:HG2	43:95:11:GLN:HB2	1.64	0.78
25:1H:2712(A):A:OP1	62:1H:3621:HOH:O	2.02	0.78
49:J8:29:GLY:O	49:J8:31:GLY:N	2.16	0.78
1:1G:582:U:OP1	15:6A:68:ARG:NH2	2.15	0.78
1:1G:1119:C:OP2	9:82:9:ARG:NH2	2.17	0.78
26:1J:76:G:N7	62:1J:304:HOH:O	2.16	0.78
33:69:79:ILE:HB	33:69:142:VAL:HG11	1.65	0.78
4:3E:15:GLU:HG2	4:3E:63:LYS:HB3	1.65	0.78
25:14:2121:G:H22	25:14:2177:C:H42	1.31	0.78
39:98:2:ARG:O	39:98:4:LEU:N	2.15	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:595:G:O2'	1:1G:596:C:OP2	2.00	0.78
1:1G:1324:A:OP2	13:4A:99:ARG:NH2	2.17	0.78
25:14:1310:G:OP2	55:L5:9:ARG:NH1	2.17	0.78
1:13:1023:G:H3'	1:13:1024:G:H5''	1.64	0.78
2:1E:101:MET:HA	2:1E:108:ILE:HG13	1.63	0.78
31:41:102:PHE:HA	31:41:105:LYS:HE2	1.65	0.78
1:1G:1200:C:O2'	1:1G:1201:A:OP2	2.00	0.78
25:14:7:G:H1	25:14:2896:C:H42	1.32	0.78
37:35:55:ARG:HH22	37:35:58:THR:HG22	1.48	0.78
25:1H:1252:G:N2	42:C8:37:GLU:OE2	2.12	0.77
25:1H:746:A:O2'	25:1H:747:U:OP2	2.01	0.77
25:1H:973:A:OP2	62:1H:3620:HOH:O	2.02	0.77
41:B8:60:THR:HG22	41:B8:77:PRO:HA	1.66	0.77
7:62:113:GLU:HB2	7:62:119:ARG:HG2	1.66	0.77
46:C5:55:TYR:H	46:C5:55:TYR:HD1	1.31	0.77
25:1H:427:U:OP2	62:1H:3619:HOH:O	2.01	0.77
25:1H:819:A:OP2	25:1H:1187:G:N2	2.18	0.77
13:4A:3:ARG:HH12	31:49:113:ARG:HB3	1.49	0.77
25:14:259:G:H21	25:14:621:A:H8	1.28	0.77
47:D5:152:ALA:O	47:D5:154:ASP:N	2.16	0.77
25:1H:731:C:OP2	62:1H:3622:HOH:O	2.02	0.77
25:1H:2131:G:H4'	25:1H:2132:U:H4'	1.64	0.77
25:1H:2168:G:H21	25:1H:2169:A:H3'	1.48	0.77
25:14:2364:C:OP1	48:E5:55:ARG:NH1	2.18	0.77
1:1G:971:G:N2	1:1G:1363:A:OP2	2.17	0.77
25:14:517:C:OP1	53:J5:16:ARG:NH2	2.17	0.77
25:14:2377:A:H4'	40:65:111:GLU:HB3	1.66	0.77
11:2I:41:THR:HG21	11:2I:71:LYS:HD3	1.65	0.77
14:5A:9:LYS:HA	14:5A:12:ARG:HB3	1.67	0.77
4:3E:105:VAL:HG21	4:3E:126:ILE:HD13	1.65	0.77
22:1K:49:C:O2'	22:1K:50:G:OP2	2.01	0.77
25:1H:607:U:H3	25:1H:621:A:H2	1.32	0.77
25:1H:774:A:H2	25:1H:787:U:HO2'	1.33	0.77
1:1G:1118:C:OP1	9:82:104:ARG:NH1	2.17	0.77
47:D5:52:SER:O	47:D5:54:HIS:N	2.18	0.77
5:42:88:LYS:HB3	5:42:123:LEU:HB2	1.65	0.77
25:14:641:C:H42	25:14:647:G:H1	1.31	0.77
46:C5:91:GLU:O	46:C5:92:ASN:ND2	2.18	0.77
13:4I:39:ILE:HD12	13:4I:56:LEU:HD23	1.65	0.77
22:3L:52:C:H2'	22:3L:53:G:H8	1.50	0.77
30:39:5:ALA:HB1	30:39:125:LEU:HD21	1.67	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
46:C5:60:PHE:HD2	46:C5:62:GLU:H	1.30	0.76
25:14:154:G:O6	25:14:172:C:N4	2.18	0.76
28:19:255:LYS:HE3	28:19:255:LYS:H	1.50	0.76
35:58:95:PRO:O	35:58:97:ARG:N	2.18	0.76
25:1H:943:U:OP2	62:1H:3623:HOH:O	2.02	0.76
25:1H:1186:G:OP1	62:1H:3624:HOH:O	2.04	0.76
25:14:958:U:OP2	38:45:14:ARG:NH1	2.18	0.76
31:49:47:LYS:HA	31:49:82:LEU:HD21	1.68	0.76
1:13:380:G:N2	1:13:383:A:OP2	2.19	0.76
1:13:501:C:OP1	12:3I:114:ARG:NH2	2.19	0.76
31:41:114:ILE:HB	31:41:117:PHE:HB2	1.67	0.76
25:14:2106:G:H1	25:14:2183:C:H42	1.30	0.76
25:1H:2405:G:O2'	25:1H:2411:A:N6	2.19	0.76
1:1G:1221:G:OP1	1:1G:1320:C:N4	2.15	0.76
25:1H:2584:U:H2'	25:1H:2585:U:H2'	1.67	0.76
37:35:57:THR:HG21	37:35:60:MET:HB2	1.68	0.76
1:13:914:A:H2'	1:13:915:A:H8	1.51	0.76
1:1G:1321:C:H5''	1:1G:1322:C:H5''	1.68	0.75
10:1A:61:GLU:HB2	14:5A:58:LYS:HE2	1.66	0.75
31:49:16:ARG:HE	31:49:31:VAL:HG11	1.50	0.75
36:25:104:ARG:NH1	41:75:36:GLU:OE2	2.20	0.75
1:13:1026:G:O6	1:13:1036:G:N2	2.16	0.75
25:1H:881:G:H22	25:1H:895:U:H3	1.33	0.75
25:1H:1601:G:N7	62:1H:3686:HOH:O	2.19	0.75
25:14:741:G:OP1	62:14:3515:HOH:O	2.04	0.75
54:K5:11:LEU:HG	54:K5:53:LYS:HG3	1.69	0.75
1:13:1189:C:OP1	10:1I:51:ARG:NH2	2.20	0.75
1:13:1309:G:O6	1:13:1328:C:N4	2.18	0.75
25:1H:2624:G:N7	62:1H:3682:HOH:O	2.18	0.75
48:I8:27:GLU:HG3	48:I8:68:GLU:HA	1.68	0.75
1:13:448:A:OP2	1:13:485:G:N2	2.18	0.75
25:1H:1333:C:OP2	62:1H:3627:HOH:O	2.05	0.75
25:1H:1388:G:N7	62:1H:3685:HOH:O	2.19	0.75
1:1G:1196:U:O4	24:4L:22:A:N6	2.20	0.75
1:13:251:G:O6	1:13:271:C:N4	2.17	0.75
12:3I:34:CYS:HA	12:3I:55:VAL:HA	1.69	0.75
25:14:783:A:OP2	62:14:3511:HOH:O	2.03	0.75
25:14:2102:U:H3	25:14:2187:G:H1	1.32	0.75
25:14:2819:G:OP1	62:14:3513:HOH:O	2.04	0.75
39:55:29:LEU:HB3	39:55:75:LEU:HD11	1.67	0.75
25:1H:1496:A:H8	25:1H:1577:C:HO2'	1.35	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:16:21:G:H1	26:16:62:C:H42	1.35	0.75
46:G8:91:GLU:O	46:G8:92:ASN:ND2	2.19	0.75
35:15:23:LEU:HB3	35:15:60:ILE:HG21	1.67	0.75
48:E5:48:GLY:H	48:E5:51:VAL:HB	1.51	0.75
25:1H:489:G:N7	44:E8:49:LYS:NZ	2.33	0.75
25:14:2867:G:OP2	41:75:119:LYS:NZ	2.18	0.75
1:13:1255:G:OP2	10:1I:45:ARG:NH2	2.20	0.74
47:H8:52:SER:O	47:H8:54:HIS:N	2.20	0.74
1:1G:1286:A:H8	1:1G:1287:A:H4'	1.51	0.74
1:13:64:G:O6	1:13:99:C:N4	2.20	0.74
25:1H:805:G:OP1	62:1H:3626:HOH:O	2.04	0.74
1:1G:77:C:H42	1:1G:92:G:H1	1.35	0.74
38:45:54:MET:HE1	38:45:118:LEU:HD23	1.69	0.74
40:65:25:ARG:HG3	40:65:88:ASP:HB2	1.68	0.74
25:1H:963:U:OP1	62:1H:3625:HOH:O	2.04	0.74
1:1G:1114:C:H1'	14:5A:60:SER:HB2	1.70	0.74
25:14:770:G:OP2	62:14:3514:HOH:O	2.04	0.74
25:14:1619:G:N7	62:14:3550:HOH:O	2.19	0.74
32:51:77:LYS:HE2	32:51:138:LYS:HB2	1.70	0.74
57:2L:76:C:H3'	57:2L:77:A:H5''	1.69	0.74
19:AA:80:TYR:HE1	19:AA:83:HIS:HB2	1.50	0.74
1:13:358:U:OP1	33:69:87:LYS:NZ	2.18	0.74
2:1E:40:HIS:HB2	2:1E:190:THR:HG21	1.68	0.74
9:8E:112:LYS:HA	9:8E:119:ALA:HB2	1.67	0.74
28:11:142:VAL:HG23	28:11:193:VAL:HA	1.70	0.74
25:14:994:C:OP1	42:85:53:ARG:NH2	2.20	0.74
33:69:71:ILE:O	33:69:74:ASN:ND2	2.20	0.74
25:1H:2062:A:OP1	62:1H:3629:HOH:O	2.06	0.74
18:9A:51:LEU:HB2	18:9A:56:THR:HG23	1.70	0.74
1:13:963:G:N3	10:1I:55:LYS:NZ	2.34	0.74
25:1H:1299:G:N7	62:1H:3692:HOH:O	2.20	0.74
25:1H:1980:G:O2'	25:1H:1982:C:OP2	2.05	0.74
1:1G:895:G:H1	1:1G:904:C:H42	1.36	0.74
25:14:1728:G:N1	25:14:1730:U:OP2	2.21	0.74
25:14:2469:A:H8	38:45:56:ARG:HH21	1.35	0.74
38:45:3:MET:HE2	38:45:3:MET:H	1.53	0.74
1:13:951:G:OP2	13:4I:102:ARG:NH2	2.21	0.74
25:1H:2287:A:H62	25:1H:2344:U:H3	1.36	0.74
1:1G:1115:C:H1'	14:5A:61:TRP:HB2	1.69	0.73
15:6A:16:ALA:HB1	15:6A:21:ASP:HB3	1.67	0.73
1:13:1321:C:H5''	1:13:1322:C:H5''	1.70	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:270:A:OP1	62:1H:3628:HOH:O	2.05	0.73
1:1G:1076:C:N4	1:1G:1081:G:O6	2.17	0.73
9:82:20:ARG:HB3	9:82:60:ASP:HB2	1.68	0.73
9:82:50:LEU:HA	9:82:53:VAL:HG22	1.70	0.73
25:14:1918:A:O2'	25:14:1920:OMC:N4	2.20	0.73
36:25:75:SER:OG	41:75:74:ARG:NH1	2.21	0.73
40:65:74:ALA:HB1	40:65:107:GLU:HB2	1.70	0.73
25:1H:2343:C:O2'	25:1H:2373:G:O2'	2.06	0.73
1:13:639:G:H2'	1:13:640:A:H8	1.54	0.73
25:1H:878:A:N6	25:1H:899:A:O2'	2.21	0.73
30:39:61:GLY:HA2	30:39:77:ASP:HB3	1.70	0.73
13:4A:44:ARG:HG2	13:4A:45:VAL:H	1.52	0.73
25:14:558:G:H5''	35:15:112:LEU:HD22	1.69	0.73
25:14:1997:G:OP2	62:14:3518:HOH:O	2.06	0.73
41:B8:55:ASN:H	41:B8:59:THR:HB	1.53	0.73
51:L8:10:LYS:NZ	51:L8:15:TYR:OH	2.22	0.73
1:1G:147:G:H2'	1:1G:148:G:H8	1.51	0.73
5:42:9:LYS:HB2	5:42:112:LEU:HD11	1.70	0.73
25:14:2777:G:OP2	25:14:2781:A:O2'	2.04	0.73
1:13:745:C:OP1	1:13:851:G:O2'	2.07	0.73
25:1H:563:G:OP2	62:1H:3630:HOH:O	2.06	0.73
25:1H:2306:C:N4	31:41:42:GLY:O	2.22	0.73
41:B8:24:PRO:HB2	41:B8:99:LEU:HD21	1.70	0.73
1:1G:1324:A:H4'	1:1G:1362:C:H4'	1.70	0.73
1:1G:1343:G:H4'	9:82:122:ALA:HB3	1.71	0.73
1:1G:1422:G:H5''	36:25:48:PRO:HB3	1.69	0.73
25:14:708:C:H42	25:14:723:G:H1	1.33	0.73
54:K5:8:LYS:HB2	54:K5:27:LYS:HA	1.69	0.73
25:14:583:G:N7	62:14:3557:HOH:O	2.20	0.73
1:13:1452:C:H4'	1:13:1453:G:H5'	1.71	0.73
25:1H:2159:G:H2'	25:1H:2160:G:H8	1.54	0.73
38:88:30:GLY:HA2	38:88:107:ALA:HB2	1.70	0.73
9:82:46:ALA:HB2	9:82:74:ILE:HG23	1.69	0.73
25:14:2719:G:OP2	62:14:3517:HOH:O	2.06	0.73
25:1H:2832:U:H4'	25:1H:2833:G:H5''	1.70	0.73
51:L8:5:LYS:HB3	51:L8:57:GLU:HB3	1.69	0.73
1:1G:1162:C:H42	1:1G:1174:G:H1	1.35	0.73
25:14:640:C:H42	25:14:648:G:H1	1.36	0.73
25:14:1283:G:N2	25:14:1286:A:OP2	2.21	0.73
25:14:1800:C:OP2	28:19:183:ARG:NH2	2.21	0.73
25:1H:511:U:OP2	62:1H:3631:HOH:O	2.07	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:2346:A:O2'	54:O8:24:GLU:OE2	2.07	0.72
25:14:450:G:O6	62:14:3516:HOH:O	2.05	0.72
41:75:16:ARG:NH1	41:75:80:SER:O	2.21	0.72
1:13:1002:G:H2'	1:13:1003:G:H8	1.54	0.72
3:2E:20:SER:OG	3:2E:40:ARG:NH2	2.22	0.72
25:14:2589:A:OP1	62:14:3511:HOH:O	2.07	0.72
35:58:4:TYR:O	42:C8:64:ARG:NH1	2.21	0.72
41:B8:57:PHE:O	41:B8:58:ASN:ND2	2.22	0.72
10:1A:3:LYS:N	10:1A:74:ILE:O	2.22	0.72
10:1A:48:THR:HA	10:1A:62:HIS:HB3	1.71	0.72
25:14:273(C):C:H42	25:14:363(C):G:H1	1.36	0.72
26:1J:4:C:H42	26:1J:116:G:H1	1.35	0.72
25:1H:1728:G:H8	25:1H:1732:A:H62	1.38	0.72
25:14:2285:C:OP2	54:K5:27:LYS:NZ	2.23	0.72
47:D5:80:ARG:HB2	47:D5:82:ARG:HG2	1.72	0.72
25:1H:565:C:OP1	43:D8:82:ARG:NH2	2.22	0.72
25:1H:1016:G:N7	62:1H:3695:HOH:O	2.20	0.72
40:A8:26:LEU:HB3	40:A8:87:PHE:HA	1.72	0.72
25:14:2119:A:N6	25:14:2170:A:N7	2.37	0.72
25:1H:2524:G:N7	62:1H:3705:HOH:O	2.23	0.72
1:1G:536:C:OP1	62:1G:1804:HOH:O	2.08	0.72
1:1G:1437:C:H42	1:1G:1464:G:H1	1.34	0.72
4:32:191:ARG:NH1	4:32:191:ARG:O	2.22	0.72
25:14:729:G:OP2	28:19:13:ARG:NH1	2.22	0.72
41:B8:20:PRO:HD2	41:B8:86:ILE:HG23	1.71	0.72
1:1G:1256:A:N6	1:1G:1278:U:OP2	2.22	0.72
25:14:2472:G:N2	25:14:2529:G:O6	2.23	0.72
1:13:198:G:H1	1:13:219:C:H42	1.36	0.72
25:14:2239:G:OP2	62:14:3519:HOH:O	2.07	0.72
4:3E:120:LEU:HB3	4:3E:126:ILE:HD11	1.71	0.72
30:31:101:LEU:HD12	30:31:102:PRO:HD2	1.72	0.72
45:B5:43:VAL:HG23	45:B5:51:VAL:HG21	1.70	0.72
25:14:67:U:H3	25:14:74:A:H2	1.38	0.71
25:14:530:G:O2'	25:14:532:A:N7	2.23	0.71
25:14:2357:U:OP1	48:E5:20:ARG:NH1	2.23	0.71
25:1H:1352:U:O5'	62:1H:3635:HOH:O	2.09	0.71
1:1G:833:U:H3	1:1G:853:G:H1	1.38	0.71
47:D5:48:PHE:HA	47:D5:51:ALA:HB3	1.71	0.71
14:5I:21:TYR:HE2	14:5I:23:ARG:HE	1.36	0.71
25:1H:1828:G:OP2	62:1H:3633:HOH:O	2.07	0.71
11:2I:33:THR:HA	11:2I:39:PRO:HA	1.72	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1498:UR3:O2'	1:1G:1499:A:OP2	2.08	0.71
42:85:90:VAL:HG13	43:95:39:LEU:HB2	1.73	0.71
25:1H:958:U:OP2	38:88:14:ARG:NH1	2.22	0.71
25:1H:1042:G:N2	25:1H:1113:U:O2	2.24	0.71
38:88:79:LEU:O	38:88:81:VAL:N	2.22	0.71
25:14:456:C:H2'	45:B5:69:TYR:HE2	1.55	0.71
25:14:2706:G:N7	62:14:3567:HOH:O	2.23	0.71
46:C5:102:CYS:SG	46:C5:103:GLY:N	2.62	0.71
1:13:1320:C:H5'	19:AI:70:LYS:HG3	1.73	0.71
25:1H:467:G:OP1	55:P8:33:ARG:NH1	2.24	0.71
29:21:54:GLN:HB2	29:21:76:ARG:HB3	1.70	0.71
1:1G:1125:U:OP2	1:1G:1145:C:N4	2.23	0.71
1:1G:1268:A:N3	1:1G:1326:C:O2'	2.24	0.71
25:14:2343:C:HO2'	25:14:2373:G:HO2'	1.30	0.71
28:19:242:ARG:HG3	28:19:246:PRO:HG3	1.72	0.71
1:13:1347:G:N2	1:13:1374:A:OP2	2.19	0.71
2:1E:77:ALA:HB2	2:1E:211:ILE:HD13	1.72	0.71
25:1H:372:G:N2	25:1H:401:A:OP2	2.22	0.71
25:1H:2126:A:H2	25:1H:2162:G:H21	1.39	0.71
32:51:149:ARG:NH2	32:51:167:GLU:OE2	2.22	0.71
25:14:536:A:OP1	42:85:53:ARG:NH1	2.23	0.71
56:M5:37:SER:O	56:M5:40:GLU:N	2.20	0.71
1:13:184:G:H2'	1:13:185:A:H8	1.55	0.71
36:68:107:ARG:HA	36:68:112:MET:HE1	1.73	0.71
1:1G:736:C:H2'	1:1G:737:A:C8	2.26	0.71
25:14:1332:G:OP1	62:14:3522:HOH:O	2.09	0.71
38:45:59:ARG:HG2	38:45:61:GLY:H	1.54	0.71
25:1H:2249:U:O4	62:1H:3632:HOH:O	2.07	0.71
26:16:44:G:H1'	26:16:47:C:H42	1.55	0.71
1:13:766:A:OP2	62:13:1804:HOH:O	2.08	0.70
2:12:167:PRO:O	2:12:171:ALA:N	2.24	0.70
22:3K:22:A:H61	22:3K:47:G:H2'	1.56	0.70
25:1H:1171:G:N2	25:1H:1178:C:O2	2.24	0.70
25:1H:1899:G:H21	25:1H:1902:C:N4	1.89	0.70
31:41:141:PHE:HB2	31:41:144:ILE:HG12	1.73	0.70
1:1G:673:G:H2'	1:1G:674:G:C8	2.26	0.70
25:14:2499:C:OP2	62:14:3521:HOH:O	2.08	0.70
41:75:58:ASN:C	41:75:58:ASN:HD22	1.99	0.70
46:C5:55:TYR:CG	46:C5:56:PRO:HD3	2.26	0.70
40:A8:87:PHE:HB2	40:A8:112:PHE:CZ	2.26	0.70
25:14:584:C:OP2	42:85:10:ARG:NH2	2.23	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:1643:G:O6	62:14:3520:HOH:O	2.08	0.70
25:14:1900:A:H1'	25:14:1970:A:H2'	1.71	0.70
30:39:79:GLY:HA2	30:39:86:GLY:HA2	1.71	0.70
25:14:1487:G:H1	25:14:1502:C:H42	1.38	0.70
26:1J:15:A:H5'	26:1J:16:G:C8	2.26	0.70
1:13:210:U:O2'	1:13:216:G:N7	2.25	0.70
25:1H:1997:G:OP2	62:1H:3634:HOH:O	2.08	0.70
25:1H:2033:A:OP1	62:1H:3636:HOH:O	2.09	0.70
1:1G:376:G:H5''	16:7A:5:ARG:HB2	1.73	0.70
1:1G:728:A:H2'	1:1G:729:A:C8	2.26	0.70
2:12:54:THR:HG21	2:12:201:ILE:HD11	1.73	0.70
2:12:142:LEU:O	2:12:146:GLN:NE2	2.24	0.70
40:65:26:LEU:HB3	40:65:87:PHE:HA	1.73	0.70
15:6I:3:ILE:HG23	15:6I:38:ARG:HG3	1.72	0.70
20:BI:35:THR:HG22	62:BI:301:HOH:O	1.90	0.70
25:1H:1604:C:OP2	62:1H:3641:HOH:O	2.10	0.70
25:1H:2334:G:O6	48:I8:74:ARG:NH1	2.24	0.70
41:B8:124:ASP:H	41:B8:127:ALA:HB3	1.56	0.70
2:12:115:LEU:HG	2:12:153:ARG:HH21	1.57	0.70
25:14:2207:C:O2	28:19:151:LYS:NZ	2.25	0.70
39:55:78:LYS:HE2	39:55:83:ILE:HD11	1.74	0.70
28:11:17:THR:HB	28:11:205:VAL:H	1.57	0.70
30:31:185:ASP:OD1	30:31:188:ARG:NH1	2.23	0.70
3:22:60:ALA:O	3:22:63:ASN:ND2	2.24	0.70
25:14:761:A:N7	62:14:3571:HOH:O	2.23	0.70
25:14:2680:C:OP2	29:29:111:ARG:NH2	2.25	0.70
10:1I:48:THR:HA	10:1I:62:HIS:HB3	1.74	0.70
25:1H:1043:C:H42	25:1H:1112:G:H1	1.40	0.70
31:49:82:LEU:HD23	31:49:86:MET:HB3	1.74	0.70
1:13:1200:C:O2'	1:13:1201:A:OP2	2.07	0.70
24:4L:7:G:H2'	24:4L:8:A:H8	1.56	0.70
25:14:1057:A:N6	25:14:1088:A:OP2	2.25	0.70
25:14:1542:G:O6	25:14:1543:A:N6	2.24	0.70
25:1H:138:G:N2	45:F8:44:GLU:OE2	2.24	0.70
25:1H:587:C:OP2	37:78:21:ARG:NH2	2.24	0.70
36:68:14:THR:HG21	36:68:86:ILE:HB	1.73	0.70
11:2A:22:HIS:HB3	11:2A:29:ILE:HG23	1.72	0.70
13:4A:67:GLU:O	13:4A:69:GLU:N	2.25	0.70
25:14:1817:G:OP1	28:19:88:ARG:NH2	2.25	0.70
25:14:2107:C:H42	25:14:2182:G:H1	1.38	0.70
1:13:158:G:N2	1:13:163:C:O2	2.22	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:346:G:OP2	41:B8:41:ARG:NH1	2.25	0.69
1:13:1054:C:N4	22:1K:35:C:O2	2.25	0.69
52:M8:1:MET:SD	52:M8:6:HIS:NE2	2.63	0.69
25:14:1771:C:OP1	62:14:3525:HOH:O	2.10	0.69
1:13:837:G:H1	1:13:849:C:H42	1.40	0.69
1:13:1329:A:N7	21:1F:7:ARG:NH2	2.39	0.69
2:1E:69:LEU:HD23	2:1E:159:PRO:HG3	1.74	0.69
25:1H:265:A:O2'	25:1H:266:G:H4'	1.92	0.69
25:1H:2104:G:O3'	27:71:223:ARG:NH2	2.25	0.69
29:21:24:THR:HG23	29:21:184:VAL:HG23	1.72	0.69
11:2I:34:ASP:HB3	11:2I:40:ILE:HD11	1.73	0.69
25:1H:1479:G:H1	25:1H:1514:U:H3	1.41	0.69
1:1G:971:G:OP2	1:1G:1231:G:N2	2.23	0.69
25:14:1314:C:OP1	62:14:3522:HOH:O	2.11	0.69
1:13:1402:4OC:HM22	1:13:1403:C:H5'	1.74	0.69
25:1H:1530:G:O6	25:1H:1542:G:N2	2.25	0.69
25:1H:2705:A:OP2	62:1H:3644:HOH:O	2.11	0.69
25:1H:2788:C:O2'	25:1H:2809:A:N3	2.24	0.69
27:71:32:LEU:HD13	27:71:220:PRO:HD2	1.75	0.69
52:I5:2:LYS:HD3	52:I5:5:ILE:HG23	1.75	0.69
1:13:501:C:H2'	1:13:502:G:H8	1.56	0.69
32:51:45:VAL:HA	32:51:50:VAL:HG23	1.75	0.69
44:E8:18:ARG:HG2	44:E8:76:VAL:HG13	1.73	0.69
15:6A:80:ALA:O	15:6A:84:LYS:NZ	2.24	0.69
16:7A:40:ASP:HB3	16:7A:48:TRP:HB2	1.75	0.69
10:1I:38:ILE:HG23	10:1I:71:LEU:HB3	1.75	0.69
25:1H:77:C:H5''	50:K8:10:LEU:HD11	1.74	0.69
13:4I:107:ALA:HB3	13:4I:111:LYS:HD2	1.75	0.69
25:1H:450:G:OP2	62:1H:3637:HOH:O	2.09	0.69
25:1H:1310:G:OP2	55:P8:9:ARG:NH1	2.26	0.69
34:38:40:LEU:HD13	34:38:41:ARG:HG2	1.73	0.69
1:1G:448:A:OP2	1:1G:485:G:N2	2.26	0.69
25:14:1012:U:H3	25:14:1143:A:H2	1.39	0.69
25:14:2485:G:H5''	38:45:46:GLN:HE21	1.57	0.69
51:H5:10:LYS:HB3	51:H5:53:LEU:HD23	1.73	0.69
1:13:1216:G:OP2	14:5I:3:ARG:NH2	2.26	0.69
25:1H:1603:A:OP1	62:1H:3641:HOH:O	2.10	0.69
25:1H:2037:G:N7	62:1H:3718:HOH:O	2.26	0.69
1:1G:1104:G:O2'	2:12:111:ARG:NH2	2.26	0.69
25:14:1300:U:O2'	25:14:1635:G:OP1	2.10	0.69
26:1J:70:C:H42	26:1J:106:G:H1	1.41	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:19:242:ARG:O	62:19:402:HOH:O	2.11	0.69
31:49:77:ILE:HG22	31:49:80:PHE:H	1.58	0.69
31:49:97:ASP:HA	31:49:100:TRP:HD1	1.57	0.69
1:13:1422:G:H5''	36:68:48:PRO:HB3	1.74	0.69
11:2I:32:ILE:HD12	11:2I:72:ALA:HB2	1.75	0.69
25:1H:1370:C:OP2	62:1H:3643:HOH:O	2.10	0.69
29:21:47:VAL:HG11	29:21:86:PRO:HD2	1.74	0.69
6:5E:12:PRO:HB2	6:5E:57:GLN:HE21	1.57	0.69
28:11:181:GLU:HB2	28:11:273:ARG:HG2	1.75	0.69
1:13:107:G:N7	20:BI:15:ARG:NH1	2.41	0.68
1:13:1247:U:H3	1:13:1290:G:H1	1.41	0.68
13:4I:80:ARG:NH1	19:AI:65:ASN:O	2.25	0.68
25:1H:27:G:H22	25:1H:512:G:H2'	1.57	0.68
25:1H:2290:G:H1	25:1H:2342:C:H42	1.40	0.68
1:1G:276:G:O2'	17:8A:68:ARG:NH1	2.26	0.68
6:52:5:GLU:HB3	6:52:62:TRP:HE1	1.57	0.68
25:14:842:G:N7	62:14:3583:HOH:O	2.26	0.68
26:1J:73:A:OP2	62:1J:301:HOH:O	2.11	0.68
25:1H:509:C:OP1	62:1H:3646:HOH:O	2.11	0.68
25:1H:1654:A:OP2	39:98:1:MET:N	2.23	0.68
25:1H:1754:C:OP1	41:B8:96:ARG:NH1	2.26	0.68
32:51:101:ARG:HH12	32:51:122:THR:HG23	1.57	0.68
47:H8:108:PRO:HG2	47:H8:112:ARG:HB2	1.74	0.68
4:32:101:LEU:HB2	4:32:138:TYR:HB3	1.75	0.68
19:AA:49:ILE:HG21	19:AA:71:LEU:HD22	1.75	0.68
1:13:648:A:H2'	1:13:649:G:H8	1.57	0.68
1:13:1121:U:H3	1:13:1151:A:H61	1.40	0.68
25:1H:2002:G:OP2	62:1H:3639:HOH:O	2.10	0.68
33:61:127:VAL:HG22	33:61:139:GLN:HB3	1.74	0.68
1:1G:947:G:HO2'	1:1G:1306:A:HO2'	1.40	0.68
1:1G:1057:G:H1	1:1G:1203:C:H42	1.40	0.68
25:14:1784:A:OP2	62:14:3528:HOH:O	2.11	0.68
25:1H:392:C:OP1	62:1H:3642:HOH:O	2.10	0.68
19:AA:39:THR:HG22	19:AA:40:ILE:H	1.58	0.68
25:14:2037:G:N7	62:14:3596:HOH:O	2.27	0.68
13:4I:13:LYS:HA	13:4I:44:ARG:HD2	1.75	0.68
19:AI:19:VAL:HG11	19:AI:44:MET:HB3	1.75	0.68
25:1H:487:C:O2	44:E8:53:SER:OG	2.11	0.68
25:1H:674:G:H1'	30:31:74:ARG:HD3	1.76	0.68
1:1G:186(B):C:H1'	20:BA:105:SER:HB2	1.74	0.68
1:1G:474:G:H2'	1:1G:475:G:H8	1.59	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1321:C:H41	1:1G:1322:C:H41	1.38	0.68
4:32:175:SER:HB3	4:32:186:LEU:HD21	1.74	0.68
22:3L:9:G:N2	22:3L:46:G:N7	2.42	0.68
26:1J:116:G:H4'	40:65:54:LEU:HB3	1.75	0.68
32:59:125:VAL:HG12	32:59:128:PRO:HA	1.75	0.68
25:1H:1359:A:N6	25:1H:1372:U:H3	1.92	0.68
25:14:389:G:H22	37:35:72:PRO:HD3	1.57	0.68
25:14:1997:G:OP2	62:14:3524:HOH:O	2.10	0.68
26:1J:102:G:N3	47:D5:73:GLN:NE2	2.40	0.68
1:1G:147:G:H1	1:1G:175:C:H42	1.42	0.68
25:14:731:C:OP2	62:14:3527:HOH:O	2.11	0.68
45:B5:6:ASP:OD2	50:G5:29:LYS:NZ	2.24	0.68
25:1H:243:U:OP1	56:Q8:6:THR:OG1	2.12	0.68
25:1H:2270:G:OP2	62:1H:3645:HOH:O	2.11	0.68
31:41:109:VAL:HG21	52:M8:14:ILE:HD13	1.76	0.68
33:61:79:ILE:HG22	33:61:81:VAL:HG22	1.76	0.68
38:88:33:GLY:HA2	38:88:105:GLU:HA	1.74	0.68
40:A8:59:LYS:HD3	40:A8:60:GLY:H	1.59	0.68
25:14:370:G:OP2	62:14:3526:HOH:O	2.11	0.68
25:14:831:G:N7	62:14:3582:HOH:O	2.25	0.68
2:1E:132:LYS:HA	2:1E:135:GLN:HB2	1.76	0.68
25:1H:1221:C:OP1	43:D8:68:LYS:NZ	2.27	0.68
47:H8:4:ARG:NH1	47:H8:60:GLU:OE2	2.27	0.68
3:22:32:LEU:O	3:22:59:ARG:NH1	2.27	0.68
38:45:55:VAL:HG13	38:45:60:ARG:HA	1.75	0.68
1:13:1210:C:H2'	1:13:1211:U:H5''	1.74	0.68
1:13:1286:A:H5''	21:1F:26:LYS:HD2	1.75	0.68
30:31:198:ALA:HA	30:31:201:VAL:HG12	1.76	0.68
41:B8:28:VAL:HB	41:B8:86:ILE:HD11	1.76	0.68
1:1G:186(A):C:OP1	20:BA:86:ARG:NH1	2.27	0.68
5:42:88:LYS:HD3	5:42:123:LEU:HD12	1.76	0.68
1:13:113:G:H1	1:13:314:C:H42	1.42	0.67
1:13:997:U:H2'	1:13:998:G:C8	2.29	0.67
25:1H:2751:G:O2'	25:1H:2752:C:O5'	2.12	0.67
25:1H:2882:A:OP1	39:98:96:ARG:NH1	2.27	0.67
1:1G:1227:A:OP1	19:AA:80:TYR:OH	2.07	0.67
25:14:1828:G:OP2	62:14:3532:HOH:O	2.12	0.67
38:45:64:ILE:HG22	38:45:106:VAL:HG12	1.75	0.67
40:65:106:ARG:NH2	40:65:107:GLU:OE1	2.27	0.67
46:C5:55:TYR:HE2	46:C5:60:PHE:HE1	1.40	0.67
47:D5:107:THR:HB	47:D5:108:PRO:HD3	1.75	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:1606:G:OP1	62:1H:3653:HOH:O	2.13	0.67
25:1H:2154:G:H2'	25:1H:2155:G:H8	1.58	0.67
37:78:125:VAL:HG13	37:78:144:GLU:HB3	1.75	0.67
38:45:11:LYS:NZ	38:45:86:GLY:O	2.27	0.67
20:BI:62:LEU:HA	20:BI:65:LYS:HB2	1.76	0.67
25:1H:2683:C:OP1	41:B8:53:ARG:NH2	2.27	0.67
25:1H:2844:G:O6	62:1H:3640:HOH:O	2.10	0.67
34:38:50:ARG:HB2	34:38:50:ARG:HH11	1.58	0.67
17:8A:62:SER:HB3	17:8A:72:ARG:HH11	1.59	0.67
25:14:2820:A:OP1	39:55:2:ARG:NH2	2.19	0.67
1:13:183:G:N2	1:13:223:U:O2'	2.27	0.67
19:AI:28:LYS:HA	19:AI:47:HIS:HE1	1.60	0.67
25:1H:2499:C:N3	62:1H:3731:HOH:O	2.27	0.67
30:31:28:ILE:HG22	30:31:112:MET:HB3	1.77	0.67
4:32:15:GLU:OE1	4:32:66:ARG:NH1	2.27	0.67
22:3L:19:G:O2'	22:3L:61:U:N3	2.28	0.67
22:3L:29:C:H2'	22:3L:30:G:H8	1.59	0.67
25:14:411:G:OP1	62:14:3533:HOH:O	2.12	0.67
25:1H:1021:A:H8	25:1H:1022:G:H5''	1.60	0.67
25:1H:2680:C:H5'	29:21:189:PRO:HA	1.74	0.67
25:1H:2701:C:H3'	25:1H:2702:U:C5'	2.24	0.67
49:J8:83:GLU:HG2	49:J8:85:LEU:H	1.58	0.67
54:O8:47:THR:HG22	54:O8:48:VAL:HG12	1.76	0.67
1:1G:616:G:H1	1:1G:624:C:H42	1.40	0.67
1:1G:976:G:N2	1:1G:1362(A):C:OP2	2.19	0.67
1:1G:1127:G:H2'	1:1G:1128:C:H6	1.59	0.67
1:1G:1240:U:H2'	7:62:32:ARG:HD2	1.77	0.67
12:3A:34:CYS:HA	12:3A:55:VAL:HA	1.75	0.67
25:14:1024:G:H3'	25:14:1025:G:H5''	1.76	0.67
30:39:116:ASP:OD1	30:39:119:ARG:NH2	2.28	0.67
31:49:64:THR:HG23	31:49:66:GLN:H	1.59	0.67
4:3E:157:LEU:O	4:3E:161:ASN:ND2	2.25	0.67
25:1H:617:G:OP1	30:31:40:GLN:NE2	2.27	0.67
4:3E:53:ASP:OD1	4:3E:57:ARG:NH2	2.28	0.67
25:1H:946:G:OP1	62:1H:3651:HOH:O	2.12	0.67
39:98:72:ASP:HB3	39:98:75:LEU:HD12	1.75	0.67
47:H8:15:PRO:HA	47:H8:18:LEU:HD13	1.76	0.67
1:1G:189:U:O2	17:8A:63:ARG:NH1	2.27	0.67
25:14:1315:C:OP2	62:14:3522:HOH:O	2.11	0.67
22:1K:12:G:H1	22:1K:24:C:H42	1.43	0.67
25:1H:49:A:N7	25:1H:120:U:H5	1.93	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:93:U:H2'	1:1G:95:G:C8	2.30	0.67
1:1G:1281:U:O4	10:1A:5:ARG:NH2	2.28	0.67
25:14:2415:G:H4'	37:35:67:MET:H	1.59	0.67
25:14:2881:C:H2'	39:55:96:ARG:CZ	2.24	0.67
31:49:68:PRO:HA	31:49:92:VAL:HB	1.77	0.67
33:69:76:THR:HG21	33:69:138:ILE:HD11	1.77	0.67
1:13:1124:G:H3'	1:13:1145:C:H41	1.59	0.67
25:1H:1083:U:H2'	25:1H:1085:A:H5''	1.75	0.67
25:1H:2175:C:O2'	27:71:219:GLY:O	2.12	0.67
1:1G:1038:C:O2'	1:1G:1039:C:O5'	2.13	0.67
2:12:11:LEU:HD13	2:12:11:LEU:H	1.59	0.67
11:2A:13:GLN:HE21	11:2A:15:ALA:HB2	1.60	0.67
25:14:95:G:HO2'	50:G5:48:HIS:HD1	1.43	0.67
25:14:997:G:OP2	42:85:58:ARG:NH1	2.28	0.67
25:14:2060:A:O2'	25:14:2502:G:O4'	2.11	0.67
30:39:28:ILE:HG22	30:39:112:MET:HE3	1.77	0.67
32:59:26:VAL:HG21	32:59:75:ALA:HB1	1.76	0.67
36:25:80:ASP:OD1	41:75:64:ARG:NH2	2.27	0.67
53:J5:16:ARG:NH1	53:J5:17:ASP:OD1	2.28	0.67
5:4E:11:ILE:HD11	5:4E:31:LEU:HD12	1.76	0.67
25:1H:730:C:OP2	62:1H:3622:HOH:O	2.13	0.67
47:H8:7:ALA:HB2	47:H8:59:LEU:HD22	1.76	0.67
22:3L:63:C:N4	22:3L:64:G:O6	2.28	0.67
25:14:883:G:H22	25:14:893:C:H42	1.43	0.67
1:13:1386:G:N7	62:13:1816:HOH:O	2.28	0.66
7:6E:20:ASP:HB3	7:6E:23:VAL:HG22	1.76	0.66
25:1H:833:U:O2	37:78:55:ARG:NH1	2.28	0.66
25:1H:996:A:OP2	42:C8:92:ARG:NH2	2.28	0.66
25:1H:1906:G:H1	25:1H:1924:C:H42	1.42	0.66
34:38:93:LEU:HD22	34:38:127:GLU:HA	1.75	0.66
52:M8:9:LEU:H	52:M8:27:THR:HG23	1.59	0.66
9:82:103:THR:HG22	9:82:105:ASP:H	1.59	0.66
25:14:1110:G:O2'	25:14:1111:A:O4'	2.13	0.66
25:14:2575:C:O3'	62:14:3530:HOH:O	2.11	0.66
30:39:63:LYS:HE3	30:39:67:GLN:HB3	1.75	0.66
1:13:105:G:OP2	20:BI:18:GLN:NE2	2.27	0.66
1:1G:476:G:H2'	1:1G:477:G:H8	1.60	0.66
13:4A:73:GLU:HG3	52:I5:52:THR:HG21	1.76	0.66
25:14:1297:C:O2'	25:14:1302:A:N1	2.26	0.66
25:14:2873:A:H8	39:55:6:SER:H	1.43	0.66
30:39:125:LEU:HD23	30:39:125:LEU:H	1.59	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:85:88:ILE:HG22	43:95:49:THR:HA	1.76	0.66
25:1H:602:G:O2'	25:1H:604:G:O2'	2.14	0.66
25:1H:778:G:OP2	62:1H:3652:HOH:O	2.12	0.66
28:19:206:LEU:HD22	28:19:211:ARG:HG2	1.76	0.66
29:21:9:VAL:HG23	41:B8:3:ARG:HG2	1.77	0.66
1:1G:610:G:OP1	62:1G:1805:HOH:O	2.12	0.66
2:12:100:GLY:O	2:12:104:ASN:N	2.28	0.66
25:14:1226:G:H5'	43:95:85:LYS:HB3	1.77	0.66
11:2I:79:SER:HB2	11:2I:106:LYS:HD2	1.77	0.66
25:1H:945:A:C8	62:1H:3826:HOH:O	2.48	0.66
25:1H:1228:G:OP2	42:C8:16:LYS:NZ	2.29	0.66
25:1H:1485:G:H1	25:1H:1504:C:H42	1.42	0.66
25:1H:2421:G:OP1	54:O8:6:ARG:NH1	2.29	0.66
46:G8:14:LEU:HD23	46:G8:82:PRO:HG3	1.78	0.66
2:12:161:ALA:HB1	2:12:185:ILE:HD11	1.77	0.66
57:2L:8:4SU:HN3	57:2L:14:A:N6	1.92	0.66
25:14:2016:U:OP1	62:14:3536:HOH:O	2.14	0.66
28:19:16:MET:HG2	28:19:211:ARG:HH21	1.61	0.66
36:25:24:VAL:HA	36:25:39:ILE:HG22	1.75	0.66
42:85:92:ARG:O	42:85:95:LEU:N	2.29	0.66
1:13:1319:A:OP1	19:AI:70:LYS:NZ	2.29	0.66
1:13:1391:U:H2'	1:13:1392:G:C8	2.31	0.66
25:1H:1113:U:H5'	32:51:2:SER:HB2	1.78	0.66
25:14:1899:G:H22	25:14:1902:C:N4	1.90	0.66
30:39:40:GLN:HE22	30:39:182:ASN:HB2	1.60	0.66
12:3I:50:ARG:HB3	12:3I:66:TYR:HE1	1.61	0.66
25:1H:860:U:OP1	62:1H:3650:HOH:O	2.12	0.66
25:1H:1434:A:H61	25:1H:1558:A:N6	1.94	0.66
1:1G:1016:A:HO2'	1:1G:1217:C:HO2'	1.43	0.66
57:2L:21:U:H3'	57:2L:22:A:H5'	1.77	0.66
26:1J:18:G:H1	26:1J:65:C:H42	1.42	0.66
41:75:30:VAL:HG12	41:75:86:ILE:HD13	1.78	0.66
1:13:186(A):C:H5''	20:BI:86:ARG:HD2	1.78	0.66
1:13:1077:G:N2	1:13:1080:A:OP2	2.23	0.66
25:1H:1469:A:OP2	62:1H:3649:HOH:O	2.11	0.66
25:1H:2156:G:H2'	25:1H:2157:G:C2	2.31	0.66
30:31:155:LEU:HB2	30:31:189:THR:HG21	1.77	0.66
25:14:2392:A:H2	25:14:2424:C:H42	1.44	0.66
47:D5:146:ILE:HG23	47:D5:175:VAL:HG12	1.77	0.66
25:1H:2159:G:H2'	25:1H:2160:G:C8	2.31	0.66
16:7A:37:GLY:HA2	16:7A:50:LYS:HD3	1.77	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:138:G:H1	1:13:225:C:H42	1.42	0.66
1:13:1147:C:O2	9:8E:16:ARG:NH1	2.29	0.66
1:13:1206:G:O2'	3:2E:192:THR:O	2.12	0.66
2:1E:33:TYR:HB2	2:1E:43:ASP:HB2	1.78	0.66
5:4E:148:VAL:HG21	8:7E:107:LEU:HD22	1.78	0.66
25:1H:1899:G:H21	25:1H:1902:C:H41	1.42	0.66
27:71:194:ARG:HA	27:71:197:GLU:HB2	1.77	0.66
42:C8:92:ARG:HD2	43:D8:11:GLN:HB2	1.78	0.66
1:1G:881:G:P	12:3A:9:ARG:HH22	2.19	0.66
25:14:1156:A:OP2	62:14:3531:HOH:O	2.12	0.66
1:13:894:G:N7	62:13:1817:HOH:O	2.29	0.65
26:16:51:G:OP2	40:A8:59:LYS:NZ	2.29	0.65
43:D8:5:VAL:HG11	43:D8:57:VAL:HG11	1.77	0.65
1:1G:1071:C:H2'	1:1G:1072:G:H8	1.61	0.65
1:1G:1540:U:H3	24:4L:5:A:H61	1.44	0.65
25:14:512:G:N7	62:14:3541:HOH:O	2.28	0.65
25:14:586:A:H5'	30:39:89:VAL:HG21	1.77	0.65
25:14:2392:A:OP2	56:M5:31:HIS:NE2	2.29	0.65
30:39:143:ALA:HB1	30:39:148:LEU:HB2	1.78	0.65
1:13:1246:C:O2	1:13:1291:G:N2	2.25	0.65
3:2E:35:GLU:OE2	3:2E:97:LYS:NZ	2.30	0.65
12:3I:83:ARG:HB2	12:3I:98:VAL:HG23	1.79	0.65
25:1H:1019:U:H3	25:1H:1142(A):A:H62	1.44	0.65
25:1H:2287:A:N6	25:1H:2344:U:H3	1.94	0.65
10:1A:99:LYS:HE2	10:1A:101:VAL:HB	1.78	0.65
25:14:2873:A:H8	39:55:6:SER:N	1.93	0.65
37:35:87:ASP:OD1	37:35:87:ASP:N	2.18	0.65
7:6E:27:ILE:HD11	7:6E:40:ALA:HA	1.79	0.65
25:1H:739:G:OP1	62:1H:3654:HOH:O	2.13	0.65
25:1H:1265:A:OP2	62:1H:3657:HOH:O	2.14	0.65
25:14:311:A:OP2	62:14:3535:HOH:O	2.14	0.65
25:14:2712:U:H1'	25:14:2712(A):A:C8	2.32	0.65
33:69:7:GLU:HG3	33:69:8:PRO:HD2	1.77	0.65
4:3E:96:LEU:HD21	4:3E:139:ARG:HH12	1.61	0.65
25:1H:1972:A:H2'	25:1H:1973:G:H8	1.60	0.65
10:1A:40:LEU:HG	10:1A:41:PRO:HD2	1.76	0.65
25:14:392:C:H5''	25:14:409:C:H5''	1.78	0.65
25:14:528:A:C2	25:14:2042:A:H2'	2.32	0.65
25:14:1754:C:OP1	41:75:96:ARG:NH1	2.29	0.65
25:14:2821:A:OP2	62:14:3534:HOH:O	2.14	0.65
1:13:871:U:OP1	62:13:1805:HOH:O	2.15	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:2I:48:ILE:HD11	11:2I:64:ALA:HA	1.79	0.65
25:1H:2472:G:H22	25:1H:2477:C:H5'	1.62	0.65
1:1G:74:C:H42	1:1G:96:G:H1	1.45	0.65
1:1G:491:G:N7	62:1G:1826:HOH:O	2.28	0.65
25:14:252:G:OP2	37:35:50:ARG:NH2	2.26	0.65
48:E5:65:GLY:HA3	48:E5:83:PRO:HA	1.77	0.65
1:13:1312:G:H1	1:13:1325:C:H42	1.43	0.65
2:1E:84:GLU:HB3	2:1E:219:VAL:HG21	1.78	0.65
25:1H:1728:G:N1	25:1H:1730:U:OP2	2.29	0.65
25:1H:2334:G:H5'	40:A8:9:ARG:HG3	1.77	0.65
26:16:15:A:H1'	26:16:109:G:C4	2.30	0.65
26:16:40:U:O2'	26:16:45:A:N6	2.28	0.65
34:38:27:VAL:HG12	34:38:28:ASN:HB2	1.79	0.65
1:1G:860:A:OP2	62:1G:1807:HOH:O	2.15	0.65
25:14:1071:G:H1'	25:14:1089:G:H2'	1.77	0.65
1:13:1227:A:OP1	19:AI:80:TYR:OH	2.12	0.65
26:16:50:G:OP1	40:A8:63:THR:OG1	2.13	0.65
50:K8:47:ASN:O	50:K8:49:LYS:N	2.29	0.65
53:N8:49:CYS:HA	53:N8:56:LYS:HB2	1.79	0.65
11:2I:30:VAL:HG21	11:2I:65:ALA:HA	1.79	0.65
22:1K:31:G:H1	22:1K:41:C:H42	1.45	0.65
3:22:33:LEU:HD12	3:22:36:ASP:HB3	1.77	0.65
25:14:587:C:OP2	37:35:21:ARG:NH2	2.29	0.65
25:14:1962:5MC:O2'	25:14:1964:G:OP2	2.15	0.65
25:14:2895:U:H2'	25:14:2896:C:O4'	1.97	0.65
10:1I:7:LYS:HB3	10:1I:97:GLU:HB2	1.78	0.65
25:1H:2582:G:OP1	62:1H:3656:HOH:O	2.14	0.65
25:1H:2659:G:O2'	25:1H:2661:G:N7	2.27	0.65
27:71:181:PRO:HB2	27:71:184:LYS:HB2	1.79	0.65
1:1G:971:G:C5	1:1G:1365:G:H5'	2.32	0.65
13:4A:37:THR:O	13:4A:55:ARG:NH1	2.27	0.65
25:14:1069:A:H5''	25:14:1070:A:H5''	1.78	0.65
29:29:24:THR:HG23	29:29:184:VAL:HG23	1.79	0.65
25:1H:310:A:O2'	25:1H:311:A:H2'	1.97	0.65
25:1H:1103:A:H5'	25:1H:1104:C:C5	2.29	0.65
25:1H:1622:G:OP2	62:1H:3655:HOH:O	2.14	0.65
25:1H:2314:C:H2'	25:1H:2315:G:H8	1.62	0.65
31:41:97:ASP:H	31:41:100:TRP:CD1	2.11	0.65
1:1G:973:G:O6	1:1G:974:A:N6	2.29	0.65
1:1G:1059:C:OP1	3:22:199:LYS:NZ	2.30	0.65
3:22:92:ALA:HA	3:22:96:GLY:H	1.62	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:2270:G:OP2	62:14:3539:HOH:O	2.15	0.65
25:14:2702:U:H1'	25:14:2703:C:H5	1.62	0.65
41:75:92:GLY:HA2	41:75:116:ALA:HA	1.79	0.65
43:95:29:PRO:HA	43:95:61:VAL:HG22	1.78	0.65
19:AI:40:ILE:HG12	19:AI:41:VAL:H	1.62	0.64
25:1H:863:A:N7	62:1H:3757:HOH:O	2.30	0.64
41:B8:16:ARG:HE	41:B8:19:LEU:HD21	1.61	0.64
1:1G:346:G:OP1	41:75:41:ARG:NH2	2.30	0.64
1:1G:1026:G:N1	1:1G:1036:G:N3	2.45	0.64
2:12:8:LYS:HG2	2:12:11:LEU:HD11	1.79	0.64
19:AA:50:ALA:HB1	19:AA:57:HIS:HB3	1.78	0.64
25:14:1365:A:O2'	49:F5:11:ARG:NH2	2.24	0.64
33:69:8:PRO:HA	33:69:14:ASP:HA	1.79	0.64
12:3A:34:CYS:HB3	12:3A:55:VAL:HG22	1.80	0.64
33:69:88:ILE:HG22	33:69:90:GLY:H	1.62	0.64
25:1H:964:C:OP1	62:1H:3658:HOH:O	2.15	0.64
40:A8:24:LEU:HB2	40:A8:85:VAL:HG12	1.79	0.64
1:1G:1127:G:N2	1:1G:1145:C:O2	2.26	0.64
2:12:32:ILE:O	2:12:43:ASP:N	2.30	0.64
11:2A:62:GLN:HB2	11:2A:93:GLN:HG3	1.79	0.64
25:14:221:A:H4'	25:14:222:A:O5'	1.98	0.64
25:14:467:G:OP1	55:L5:33:ARG:NH1	2.29	0.64
25:14:1980:G:O2'	25:14:1982:C:OP2	2.15	0.64
25:14:2875:C:OP1	41:75:3:ARG:NH1	2.29	0.64
1:13:405:U:O4	4:3E:2:GLY:N	2.30	0.64
5:4E:10:MET:HB3	5:4E:32:VAL:HG22	1.79	0.64
22:3K:57:C:H2'	22:3K:58:A:H8	1.60	0.64
25:14:2495:G:O3'	38:45:81:VAL:HG11	1.97	0.64
25:14:2702:U:H1'	25:14:2703:C:C5	2.33	0.64
1:13:513:C:H42	1:13:538:G:H1	1.45	0.64
37:78:147:LEU:O	37:78:148:LEU:HB2	1.96	0.64
1:1G:260:G:OP2	20:BA:83:ARG:NH1	2.30	0.64
1:1G:321:A:H62	1:1G:328:C:H1'	1.62	0.64
1:1G:1321:C:H41	1:1G:1322:C:N4	1.94	0.64
1:1G:1325:C:H4'	21:1B:17:THR:HG21	1.80	0.64
25:14:2330:G:H1	25:14:2385:C:H42	1.45	0.64
25:14:2335:A:O2'	25:14:2336:A:O5'	2.13	0.64
47:H8:151:HIS:HD2	47:H8:170:THR:HA	1.63	0.64
1:1G:817:C:OP2	62:1G:1806:HOH:O	2.14	0.64
1:1G:986:A:N3	19:AA:52:TYR:OH	2.30	0.64
25:14:2314:C:H5'	31:49:38:VAL:HG11	1.79	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:A5:13:SER:HA	44:A5:99:ARG:HB2	1.79	0.64
1:13:160:A:H1'	1:13:344:A:N7	2.13	0.64
1:13:451:A:OP1	1:13:481:G:N2	2.31	0.64
3:2E:47:LEU:HD11	3:2E:52:LEU:HB3	1.78	0.64
19:AI:40:ILE:HG12	19:AI:41:VAL:HG13	1.80	0.64
25:1H:760:G:OP1	62:1H:3662:HOH:O	2.15	0.64
25:1H:2125:G:N2	25:1H:2172:U:OP1	2.29	0.64
25:1H:2401:U:H2'	25:1H:2402:C:H5''	1.80	0.64
39:98:1:MET:O	39:98:2:ARG:HD3	1.97	0.64
1:1G:585:G:OP1	17:8A:37:LYS:NZ	2.30	0.64
1:1G:1244:C:H2'	1:1G:1245:A:H8	1.61	0.64
2:12:32:ILE:HD11	2:12:40:HIS:HB3	1.78	0.64
25:14:132:G:H1	25:14:147:U:H3	1.46	0.64
25:14:2881:C:H2'	39:55:96:ARG:NE	2.13	0.64
49:F5:40:ARG:HH21	49:F5:42:GLN:HG2	1.62	0.64
10:1I:50:ILE:HB	14:5I:41:ARG:HE	1.62	0.64
25:1H:2196:C:OP2	62:1H:3660:HOH:O	2.15	0.64
25:1H:2552:OMU:H2'	25:1H:2554:U:OP2	1.98	0.64
31:41:29:TRP:O	31:41:33:ARG:NH1	2.30	0.64
25:14:730:C:OP2	62:14:3527:HOH:O	2.14	0.64
25:14:987:G:O2'	25:14:1000:A:N3	2.29	0.64
25:14:1346:G:OP2	62:14:3540:HOH:O	2.15	0.64
36:25:89:ASN:OD1	36:25:89:ASN:N	2.31	0.64
12:3I:67:ILE:HG23	12:3I:97:ILE:HD12	1.80	0.64
25:1H:2438:U:O3'	25:1H:2439:A:H3'	1.98	0.64
1:1G:286:G:N7	62:1G:1830:HOH:O	2.30	0.64
1:1G:947:G:O3'	13:4A:109:THR:OG1	2.16	0.64
25:14:674:G:H1'	30:39:74:ARG:HD3	1.78	0.64
25:1H:1364:G:N7	49:J8:2:SER:HB3	2.13	0.64
1:1G:452:A:OP1	16:7A:43:LYS:NZ	2.30	0.64
25:14:162:U:H4'	25:14:171:G:C4	2.33	0.64
1:13:838:G:H1	1:13:848:C:N4	1.95	0.63
17:8I:51:TYR:HE1	17:8I:76:LEU:HB2	1.63	0.63
30:31:185:ASP:HA	30:31:188:ARG:HD3	1.80	0.63
41:B8:58:ASN:C	41:B8:58:ASN:HD22	2.07	0.63
47:H8:76:LEU:HA	47:H8:83:PRO:HA	1.78	0.63
13:4A:44:ARG:HH21	13:4A:48:LEU:HD21	1.63	0.63
15:6A:39:LEU:HD12	15:6A:56:LEU:HB2	1.80	0.63
25:14:20:C:OP1	42:85:22:LYS:NZ	2.31	0.63
25:14:569:U:OP2	62:14:3537:HOH:O	2.14	0.63
33:69:48:GLU:OE1	33:69:49:ALA:N	2.31	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:A5:65:LEU:HD13	44:A5:68:ARG:HD2	1.80	0.63
1:13:372:C:H42	1:13:389:A:H62	1.46	0.63
1:13:376:G:H2'	1:13:377:G:C8	2.33	0.63
1:13:978:A:O2'	1:13:1322:C:N3	2.29	0.63
25:1H:1187:G:O6	62:1H:3648:HOH:O	2.11	0.63
25:1H:2574:G:OP1	62:1H:3663:HOH:O	2.16	0.63
1:1G:54:C:N4	1:1G:353:A:OP2	2.31	0.63
3:22:134:ILE:HD12	3:22:151:VAL:HG11	1.79	0.63
25:14:2103:C:H42	25:14:2186:G:H1	1.46	0.63
1:13:1002:G:H2'	1:13:1003:G:C8	2.32	0.63
16:7I:17:TYR:HB2	16:7I:39:TYR:HB3	1.80	0.63
25:1H:191:A:N1	62:1H:3759:HOH:O	2.30	0.63
1:1G:501:C:H2'	1:1G:502:G:C8	2.32	0.63
1:1G:686:U:O4	1:1G:703:G:O2'	2.10	0.63
37:35:85:LEU:HA	37:35:88:LEU:HD22	1.80	0.63
25:1H:484:C:H2'	25:1H:485:C:C6	2.34	0.63
25:1H:1588:C:H2'	25:1H:1589:C:H6	1.63	0.63
25:1H:1920:OMC:HM22	25:1H:1921:G:H5'	1.80	0.63
52:M8:48:ARG:HD3	52:M8:51:ASP:H	1.62	0.63
1:1G:501:C:OP2	12:3A:121:LYS:NZ	2.31	0.63
19:AA:48:THR:HG22	19:AA:61:TYR:HA	1.79	0.63
57:2L:32:G:H1	57:2L:40:C:H42	1.46	0.63
45:B5:65:ARG:HB3	45:B5:70:LEU:HA	1.79	0.63
50:G5:50:ILE:HD12	50:G5:51:ARG:H	1.63	0.63
1:13:422:C:O2'	1:13:423:G:N2	2.31	0.63
22:1K:31:G:H2'	22:1K:32:G:H8	1.63	0.63
25:1H:2471:C:N4	25:1H:2476:A:O2'	2.31	0.63
32:51:41:MET:HG3	32:51:54:ARG:HA	1.80	0.63
36:68:26:LYS:HB2	36:68:30:ALA:HB2	1.81	0.63
36:68:111:PHE:HB3	36:68:114:ILE:HG13	1.80	0.63
1:1G:946:A:O2'	1:1G:1333:A:N3	2.24	0.63
25:14:2059:A:H5'	25:14:2060:A:OP2	1.99	0.63
25:14:2116:G:OP2	25:14:2166:G:N2	2.31	0.63
32:59:146:ALA:O	32:59:150:ALA:N	2.30	0.63
47:D5:87:ASP:N	47:D5:87:ASP:OD1	2.29	0.63
25:1H:247:G:O6	56:Q8:12:LYS:NZ	2.30	0.63
25:1H:1972:A:H2'	25:1H:1973:G:C8	2.33	0.63
40:A8:20:ARG:HE	40:A8:21:THR:HG22	1.63	0.63
41:B8:111:ARG:HD3	41:B8:111:ARG:H	1.62	0.63
7:62:89:MET:HE3	7:62:90:GLU:H	1.64	0.63
16:7A:53:VAL:HG12	16:7A:79:VAL:HG22	1.79	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:55:97:VAL:HA	39:55:113:LEU:O	1.98	0.63
43:95:58:VAL:HG12	43:95:98:GLU:HB2	1.80	0.63
47:D5:138:GLU:HG3	47:D5:156:LYS:HG3	1.79	0.63
1:13:560:U:H4'	1:13:561:U:H5''	1.81	0.63
1:13:1366:C:O3'	10:11:60:ARG:NH2	2.32	0.63
4:3E:167:GLY:HA2	28:19:135:PHE:HE2	1.62	0.63
18:9I:74:ARG:HG2	18:9I:79:LEU:HB2	1.81	0.63
33:61:124:GLY:H	33:61:142:VAL:HG23	1.64	0.63
36:68:25:LEU:HB2	36:68:38:VAL:HG13	1.80	0.63
53:N8:41:PRO:HB2	53:N8:42:PRO:HD3	1.80	0.63
1:1G:1512:U:H2'	1:1G:1513:A:C8	2.34	0.63
4:3E:73:ARG:O	4:3E:77:ASN:ND2	2.26	0.63
6:5E:97:PHE:HB3	18:9I:32:ARG:HD3	1.80	0.63
18:9I:86:VAL:HG12	18:9I:87:ARG:HG2	1.80	0.63
1:1G:735:C:H2'	1:1G:736:C:H6	1.63	0.63
1:1G:779:C:H5''	11:2A:122:LYS:HG2	1.80	0.63
1:1G:1157:A:O2'	1:1G:1158:C:O5'	2.16	0.63
13:4A:79:LYS:NZ	13:4A:82:MET:SD	2.67	0.63
25:14:2585:U:O2'	25:14:2586:C:O5'	2.16	0.63
1:13:377:G:OP1	16:7I:3:LYS:NZ	2.30	0.63
1:13:936:C:H42	1:13:1379:G:H1	1.47	0.63
1:13:1152:A:H5''	10:1I:13:HIS:CD2	2.33	0.63
25:1H:99:U:H4'	25:1H:101:G:H5'	1.81	0.63
25:1H:2692:C:O2	25:1H:2847:U:O2'	2.17	0.63
39:98:26:LYS:O	39:98:30:THR:OG1	2.17	0.63
44:E8:68:ARG:HE	44:E8:112:GLY:HA2	1.64	0.63
1:1G:762:C:H2'	1:1G:763:G:H8	1.64	0.63
1:1G:963:G:N3	10:1A:55:LYS:NZ	2.46	0.63
7:62:16:LEU:O	7:62:18:TYR:N	2.32	0.63
9:82:29:ASN:O	9:82:31:GLN:N	2.32	0.63
9:82:121:ARG:NH1	9:82:122:ALA:O	2.32	0.63
25:14:783:A:H8	25:14:784:A:H4'	1.63	0.63
25:14:2394:C:H5''	37:35:64:LYS:HD2	1.81	0.63
47:D5:97:GLU:HB3	47:D5:125:LEU:HD11	1.80	0.63
25:1H:273(F):C:H3'	25:1H:274:G:H5''	1.79	0.62
44:E8:88:ARG:HB3	44:E8:92:ARG:HB2	1.80	0.62
49:J8:4:VAL:HG23	49:J8:11:ARG:HB3	1.81	0.62
1:1G:445:G:H2'	1:1G:446:G:H8	1.63	0.62
1:1G:1175:G:H2'	1:1G:1176:A:H8	1.64	0.62
19:AA:36:ARG:HG3	19:AA:72:GLY:H	1.62	0.62
25:14:62:C:H42	25:14:92:G:H1	1.47	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:15:56:ASN:H	35:15:125:GLY:H	1.47	0.62
39:55:44:LEU:HD12	39:55:114:VAL:HG11	1.78	0.62
1:13:243:A:H4'	1:13:244:U:O5'	1.99	0.62
1:13:688:G:H2'	1:13:689:C:H6	1.64	0.62
25:1H:2103:C:H2'	25:1H:2104:G:C8	2.33	0.62
33:61:5:LEU:HD11	33:61:19:VAL:HG12	1.80	0.62
33:61:96:ASP:OD1	33:61:96:ASP:N	2.32	0.62
1:1G:5:U:H4'	1:1G:6:G:O5'	1.99	0.62
1:1G:370:C:H42	1:1G:391:G:H1	1.46	0.62
1:1G:595:G:H4'	1:1G:596:C:H5'	1.81	0.62
1:1G:1298:C:H41	7:62:114:ARG:HA	1.63	0.62
3:22:77:ILE:HG23	3:22:83:ARG:HB2	1.81	0.62
19:AA:52:TYR:HB2	19:AA:57:HIS:CE1	2.34	0.62
25:14:1430:C:H2'	25:14:1431:U:C6	2.34	0.62
25:14:2844:G:H3'	25:14:2845:G:H8	1.64	0.62
36:25:71:ARG:HH21	36:25:77:ILE:HG21	1.63	0.62
1:13:1004:A:H2	1:13:1024:G:C8	2.18	0.62
2:1E:8:LYS:HD3	2:1E:8:LYS:H	1.65	0.62
2:1E:185:ILE:HG22	2:1E:199:TYR:HB2	1.81	0.62
6:5E:7:ASN:N	6:5E:7:ASN:OD1	2.31	0.62
25:1H:732:C:H3'	62:1H:3714:HOH:O	1.99	0.62
25:1H:857:C:H4'	48:I8:23:VAL:HG21	1.80	0.62
52:M8:37:SER:HA	52:M8:41:PRO:HD2	1.80	0.62
1:1G:1127:G:H2'	1:1G:1128:C:C6	2.34	0.62
25:14:451:C:OP2	62:14:3538:HOH:O	2.15	0.62
25:14:510:C:OP1	62:14:3541:HOH:O	2.15	0.62
25:14:662:G:H5''	37:35:17:LYS:HG2	1.81	0.62
28:19:101:GLU:OE2	28:19:103:ARG:NH1	2.31	0.62
32:59:41:MET:HG3	32:59:55:PRO:HD3	1.81	0.62
25:1H:1301:A:HO2'	25:1H:1302:A:H3'	1.63	0.62
37:78:52:GLU:HB3	37:78:55:ARG:HE	1.63	0.62
42:C8:92:ARG:NH1	43:D8:11:GLN:O	2.32	0.62
1:1G:359:U:H2'	1:1G:360:A:C8	2.34	0.62
13:4A:68:GLY:HA2	13:4A:71:ARG:HB2	1.81	0.62
25:14:631:A:OP1	37:35:65:ARG:NH2	2.31	0.62
25:14:994:C:O2'	25:14:996:A:OP1	2.17	0.62
25:14:1091:G:N2	25:14:1101:U:O2	2.28	0.62
25:14:2454:G:H1'	62:14:3757:HOH:O	1.99	0.62
30:39:160:ASN:OD1	30:39:163:VAL:N	2.25	0.62
41:75:26:ASP:O	41:75:49:VAL:HG12	2.00	0.62
48:E5:38:VAL:HB	48:E5:59:LEU:HB2	1.81	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1001:G:H2'	1:13:1002:G:O4'	2.00	0.62
17:8I:24:GLU:OE2	17:8I:37:LYS:NZ	2.32	0.62
25:1H:309:G:N3	25:1H:329:G:O2'	2.32	0.62
25:1H:860:U:H5	25:1H:917:A:C2	2.16	0.62
43:D8:6:LYS:HA	43:D8:11:GLN:HA	1.81	0.62
1:1G:1309:G:OP1	13:4A:88:ARG:NH1	2.29	0.62
1:1G:1318:A:H4'	19:AA:10:PHE:CD2	2.34	0.62
13:4A:81:LEU:HD13	13:4A:89:GLY:HA2	1.80	0.62
25:14:773:U:H4'	28:19:47:GLY:HA3	1.82	0.62
25:14:2111:C:H42	25:14:2147:G:H21	1.46	0.62
37:35:36:LYS:O	62:35:201:HOH:O	2.15	0.62
41:75:45:PHE:CD2	41:75:74:ARG:HD2	2.35	0.62
46:C5:55:TYR:CD2	46:C5:56:PRO:HD3	2.34	0.62
4:3E:122:ARG:HD3	4:3E:134:ASP:HB2	1.80	0.62
20:BI:85:MET:HA	20:BI:88:VAL:HG12	1.81	0.62
23:2K:32:G:H2'	23:2K:33:OMC:H6	1.64	0.62
22:3K:77:A:O2'	25:1H:2394:C:N3	2.26	0.62
40:A8:28:VAL:HG11	40:A8:98:VAL:HG13	1.82	0.62
1:1G:972:C:O3'	10:1A:57:LYS:NZ	2.32	0.62
7:62:111:ARG:NH1	7:62:123:GLU:OE1	2.32	0.62
28:19:10:THR:HG23	28:19:13:ARG:HB2	1.82	0.62
28:19:108:PRO:HG2	28:19:111:LEU:HD23	1.81	0.62
32:59:103:LEU:HG	32:59:105:LEU:HB3	1.79	0.62
1:13:501:C:H2'	1:13:502:G:C8	2.34	0.62
1:13:545:C:OP2	4:3E:62:GLN:NE2	2.33	0.62
1:13:1369:C:H2'	1:13:1370:G:C8	2.34	0.62
25:1H:1178:C:O2'	25:1H:1179:C:O5'	2.15	0.62
25:1H:2712:U:HO2'	25:1H:2712(A):A:H8	1.46	0.62
33:61:110:ASP:HB3	33:61:112:LYS:HG2	1.82	0.62
1:1G:646:U:H2'	1:1G:647:C:C6	2.35	0.62
5:42:76:ILE:HG12	5:42:118:ILE:HD11	1.81	0.62
7:62:41:ARG:NH2	9:82:39:GLY:O	2.32	0.62
25:14:998:C:OP2	42:85:58:ARG:NH2	2.33	0.62
25:14:2336:A:H61	48:E5:43:THR:HG21	1.63	0.62
25:14:2468:G:O2'	25:14:2469:A:O4'	2.16	0.62
47:D5:137:ILE:HG23	47:D5:157:LEU:HD11	1.80	0.62
1:13:184:G:H2'	1:13:185:A:C8	2.35	0.62
1:13:243:A:N6	1:13:281:G:O2'	2.33	0.62
1:13:642:A:N3	8:7E:113:SER:OG	2.28	0.62
10:1I:40:LEU:HB2	10:1I:69:ASN:HB3	1.82	0.62
25:1H:443:A:H3'	30:31:45:ARG:HH12	1.64	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:A8:83:LYS:HG2	40:A8:109:GLY:HA2	1.81	0.62
22:3L:3:C:H2'	22:3L:4:G:C8	2.33	0.62
25:14:635:C:O2'	25:14:639:U:OP1	2.17	0.62
25:14:1202:C:H42	25:14:1243:G:H1	1.45	0.62
26:1J:102:G:O6	62:1J:302:HOH:O	2.14	0.62
47:D5:61:LEU:HD11	47:D5:67:LEU:HD12	1.81	0.62
1:13:376:G:H2'	1:13:377:G:H8	1.64	0.62
1:13:457:C:O2	1:13:476:G:N2	2.33	0.62
21:1F:10:ARG:HG2	21:1F:13:ILE:HD12	1.82	0.62
1:1G:401:C:H2'	1:1G:402:G:H8	1.64	0.62
1:1G:929:G:H1	1:1G:1388:C:H42	1.48	0.62
6:52:22:GLU:OE1	6:52:84:ASN:ND2	2.32	0.62
38:45:108:GLY:HA3	47:D5:116:VAL:HG13	1.81	0.62
1:13:1328:C:OP1	21:1F:21:TYR:OH	2.14	0.62
5:4E:100:VAL:O	5:4E:107:ARG:NH2	2.33	0.62
9:8E:29:ASN:N	9:8E:63:ILE:O	2.30	0.62
25:1H:1171:G:N7	25:1H:1174:A:N6	2.48	0.62
25:1H:2731:G:O3'	29:21:203:LYS:NZ	2.33	0.62
27:71:58:VAL:HG22	27:71:199:HIS:HB3	1.80	0.62
45:F8:6:ASP:OD2	50:K8:29:LYS:NZ	2.23	0.62
1:1G:28:G:O2'	1:1G:296:U:OP1	2.15	0.62
1:1G:1001:G:O6	1:1G:1038:C:N4	2.33	0.62
2:12:121:LEU:HB3	2:12:127:ILE:HG12	1.82	0.62
13:4A:37:THR:HG22	13:4A:55:ARG:HD2	1.82	0.62
25:14:1620:G:O4'	55:L5:1:MET:N	2.32	0.62
25:14:2296:U:OP2	40:65:9:ARG:NH1	2.25	0.62
30:39:53:THR:HG23	30:39:55:GLY:H	1.63	0.62
54:K5:23:THR:OG1	54:K5:24:GLU:N	2.29	0.62
22:1K:10:G:H2'	22:1K:11:A:H8	1.65	0.61
25:1H:535:C:H42	25:1H:558:G:H1	1.46	0.61
25:1H:2347:C:OP1	54:O8:39:TYR:OH	2.06	0.61
25:1H:2389:G:H5''	25:1H:2390:U:H5'	1.82	0.61
30:31:155:LEU:HD13	30:31:174:VAL:HG12	1.81	0.61
33:61:6:LEU:HD22	33:61:35:LEU:HA	1.80	0.61
34:38:18:GLU:HB3	34:38:66:LEU:HD22	1.81	0.61
2:12:185:ILE:HG22	2:12:199:TYR:HB2	1.81	0.61
9:82:29:ASN:OD1	9:82:30:GLY:N	2.29	0.61
25:14:2816:C:O2	25:14:2883:A:O2'	2.17	0.61
25:14:2830:G:O6	62:14:3529:HOH:O	2.11	0.61
1:13:67:C:H2'	1:13:68:G:C8	2.34	0.61
1:13:1448:C:H42	1:13:1455:G:H1	1.47	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:547:A:H2'	25:1H:548:A:C8	2.35	0.61
28:11:166:GLN:HB2	28:11:174:ILE:HG22	1.81	0.61
25:14:2467:C:H4'	38:45:123:HIS:ND1	2.15	0.61
45:B5:27:THR:HB	45:B5:80:ILE:HG22	1.81	0.61
1:13:877:C:H5''	8:7E:88:LYS:HD3	1.81	0.61
19:AI:3:ARG:HE	19:AI:7:LYS:HD2	1.63	0.61
22:1K:20:G:N7	25:1H:881:G:O2'	2.29	0.61
25:1H:784:A:OP2	62:1H:3665:HOH:O	2.16	0.61
25:1H:1634:A:OP2	62:1H:3664:HOH:O	2.16	0.61
31:41:166:ASP:N	31:41:166:ASP:OD1	2.32	0.61
33:61:2:LYS:NZ	33:61:20:ASP:OD2	2.30	0.61
7:62:16:LEU:HD11	9:82:42:ARG:HA	1.81	0.61
8:72:123:GLU:HA	8:72:126:LYS:HD3	1.81	0.61
25:14:571:A:H5'	25:14:2030:A:H62	1.65	0.61
25:14:1973:G:OP2	62:14:3543:HOH:O	2.16	0.61
25:14:2129:C:O2	25:14:2159:G:N1	2.33	0.61
25:14:2425:A:H5'	25:14:2427:C:O4'	2.00	0.61
25:14:2471:C:H3'	25:14:2472:G:H8	1.65	0.61
38:45:138:ASP:N	38:45:138:ASP:OD1	2.33	0.61
42:85:98:LEU:C	42:85:100:VAL:H	2.06	0.61
54:K5:42:TRP:CG	54:K5:43:CYS:H	2.18	0.61
1:13:263:A:OP2	20:BI:79:ARG:NH1	2.33	0.61
1:13:427:U:OP2	4:3E:36:ARG:NH1	2.34	0.61
5:4E:102:ALA:HB1	5:4E:106:PRO:HG2	1.82	0.61
12:3I:112:LYS:O	12:3I:114:ARG:HG2	2.01	0.61
25:1H:995:C:OP2	42:C8:54:LYS:NZ	2.34	0.61
39:98:104:ARG:HB3	39:98:107:ASP:HB3	1.81	0.61
25:14:1246:A:P	37:35:15:ARG:HH12	2.24	0.61
25:14:2111:C:C2	25:14:2118:U:H1'	2.36	0.61
26:1J:28:C:H2'	26:1J:29:A:H8	1.66	0.61
37:35:138:LEU:HD21	37:35:144:GLU:HG3	1.82	0.61
38:45:12:GLN:HG2	38:45:73:PRO:HD2	1.82	0.61
43:95:22:VAL:HG22	43:95:23:GLU:H	1.65	0.61
1:13:339:C:OP2	36:68:97:ARG:NH1	2.33	0.61
1:13:1179:A:H4'	9:8E:103:THR:HA	1.81	0.61
1:13:1317:C:N3	19:AI:37:ARG:NH2	2.47	0.61
52:M8:37:SER:HB3	52:M8:42:PHE:CD1	2.36	0.61
1:1G:489:C:H2'	1:1G:490:G:H8	1.65	0.61
1:1G:643:C:H2'	1:1G:644:G:C8	2.36	0.61
2:12:47:THR:HG23	2:12:202:PRO:HD2	1.83	0.61
4:32:168:ARG:HB3	4:32:168:ARG:HH11	1.65	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:391:G:O2'	25:14:410:G:OP1	2.16	0.61
46:C5:38:ILE:HD11	46:C5:64:GLU:HB2	1.83	0.61
49:F5:72:GLU:O	49:F5:76:ARG:NH2	2.33	0.61
25:1H:1742:C:H5'	25:1H:1743:G:OP2	2.01	0.61
25:1H:2777:G:OP2	25:1H:2781:A:O2'	2.19	0.61
52:M8:56:VAL:HA	52:M8:60:GLN:HB3	1.82	0.61
1:1G:841:U:H5'	1:1G:842:C:H5	1.65	0.61
1:1G:1286:A:C8	1:1G:1287:A:H4'	2.35	0.61
2:12:17:PHE:CG	2:12:42:ILE:HG12	2.36	0.61
18:9A:31:LEU:HD13	18:9A:65:ILE:HD13	1.83	0.61
39:55:51:LEU:HD23	39:55:66:VAL:HG22	1.83	0.61
50:G5:32:LEU:HB2	50:G5:53:LEU:HD22	1.81	0.61
1:13:677:U:H3	1:13:713:G:H22	1.47	0.61
1:13:811:C:N3	62:13:1820:HOH:O	2.31	0.61
5:4E:50:GLU:HB3	5:4E:53:LEU:HD13	1.82	0.61
9:8E:5:TYR:HE1	9:8E:16:ARG:HG2	1.65	0.61
11:2I:62:GLN:NE2	11:2I:93:GLN:OE1	2.32	0.61
25:1H:1101:U:H2'	25:1H:1102:C:C6	2.35	0.61
25:1H:1382:G:O6	62:1H:3647:HOH:O	2.11	0.61
37:78:100:LEU:HB2	37:78:106:LEU:HB2	1.81	0.61
25:14:1645:G:H5''	25:14:1646:C:H5'	1.83	0.61
25:14:2103:C:H2'	25:14:2104:G:C8	2.35	0.61
30:39:89:VAL:O	30:39:91:GLY:N	2.33	0.61
54:K5:39:TYR:HD1	54:K5:41:PRO:HD3	1.66	0.61
3:2E:65:ALA:HA	3:2E:100:ALA:HB3	1.81	0.61
10:1I:28:ARG:NH1	10:1I:34:VAL:O	2.34	0.61
14:5I:9:LYS:HG2	14:5I:12:ARG:HH11	1.66	0.61
29:21:115:GLY:O	29:21:119:ARG:HB2	2.00	0.61
1:1G:401:C:OP1	4:32:77:ASN:ND2	2.33	0.61
2:12:189:ASP:OD1	2:12:189:ASP:N	2.27	0.61
2:12:212:GLN:NE2	2:12:234:PRO:O	2.34	0.61
15:6A:29:VAL:HG13	15:6A:63:ARG:HG3	1.83	0.61
25:14:127:A:H5''	25:14:128:C:O4'	2.00	0.61
25:14:797:C:OP2	30:39:62:ARG:HG3	2.01	0.61
25:14:1300:U:H4'	25:14:1301:A:C5'	2.31	0.61
25:14:1614:A:N6	44:A5:91:GLY:HA2	2.16	0.61
25:14:2749:A:H5'	32:59:6:ARG:HD3	1.82	0.61
30:39:4:VAL:HA	30:39:19:GLU:HB3	1.83	0.61
32:59:127:GLU:O	32:59:129:THR:N	2.34	0.61
1:13:589:C:H5''	8:7E:29:SER:HB2	1.81	0.61
1:13:1002:G:H1	1:13:1038:C:H42	1.49	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1024:G:H4'	1:13:1024:G:OP1	1.99	0.61
1:13:1296:C:OP1	13:4I:44:ARG:NH2	2.34	0.61
5:4E:139:LEU:HA	5:4E:142:LEU:HD12	1.83	0.61
7:6E:89:MET:HE1	7:6E:156:TRP:H	1.66	0.61
9:8E:50:LEU:HD22	9:8E:55:ALA:HB3	1.83	0.61
16:7I:45:THR:HG22	16:7I:46:PRO:HD2	1.83	0.61
25:1H:259:G:O2'	25:1H:621:A:O2'	2.05	0.61
41:B8:50:ILE:HD11	41:B8:102:ILE:HD11	1.81	0.61
50:K8:16:LEU:H	50:K8:67:LYS:HZ1	1.48	0.61
1:1G:310:G:H4'	16:7A:31:LYS:HE3	1.81	0.61
1:1G:1279:A:O2'	1:1G:1281:U:OP2	2.16	0.61
1:1G:1315:U:O2'	1:1G:1360:A:O2'	2.17	0.61
5:42:57:LYS:HA	5:42:60:TYR:HB2	1.83	0.61
25:14:309:G:N3	25:14:329:G:O2'	2.34	0.61
28:19:168:ARG:HA	28:19:173:VAL:HA	1.83	0.61
19:AI:40:ILE:HD11	19:AI:62:ILE:HD13	1.83	0.61
25:1H:64:A:C4	45:F8:66:LEU:HD22	2.36	0.61
33:61:29:TYR:O	33:61:33:ARG:HB2	2.00	0.61
37:78:57:THR:HG22	37:78:59:LEU:H	1.65	0.61
1:1G:60:A:N6	1:1G:110:C:N3	2.49	0.61
11:2A:11:LYS:HG3	11:2A:12:ARG:HG2	1.83	0.61
22:3L:51:U:O2	22:3L:65:G:N2	2.34	0.61
25:14:270(N):G:H5''	33:69:53:ALA:HB1	1.81	0.61
25:14:1871:A:H2'	25:14:1872:A:C8	2.36	0.61
25:14:2086:U:H2'	25:14:2087:G:C8	2.36	0.61
25:14:2748:A:N1	25:14:2749:A:N6	2.49	0.61
26:1J:101:A:OP2	62:1J:303:HOH:O	2.16	0.61
42:85:50:ARG:NH1	43:95:72:VAL:HG11	2.12	0.61
1:13:1070:U:H2'	1:13:1071:C:H6	1.65	0.60
25:1H:2002:G:OP2	39:98:9:LYS:NZ	2.28	0.60
28:11:221:VAL:HG22	28:11:226:MET:HE2	1.83	0.60
52:M8:37:SER:HB3	52:M8:42:PHE:CG	2.36	0.60
1:1G:458:C:H42	1:1G:474:G:H1	1.47	0.60
1:1G:975:A:H5'	1:1G:1363:A:N6	2.16	0.60
2:12:8:LYS:HG2	2:12:11:LEU:HD21	1.82	0.60
14:5A:21:TYR:OH	14:5A:23:ARG:NH2	2.33	0.60
25:14:751:A:OP1	62:14:3544:HOH:O	2.16	0.60
25:14:2293:C:H42	25:14:2339:G:H1	1.49	0.60
43:95:34:GLU:OE2	43:95:56:SER:OG	2.17	0.60
7:6E:5:ARG:HB3	7:6E:7:ALA:H	1.64	0.60
25:1H:811:U:H2'	37:78:21:ARG:HA	1.83	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
27:71:10:LEU:HA	27:71:13:LYS:HD2	1.84	0.60
29:21:28:ALA:HB3	29:21:93:VAL:HG12	1.82	0.60
1:1G:1077:G:N2	1:1G:1080:A:OP2	2.31	0.60
2:12:21:ARG:HA	2:12:39:ILE:HA	1.83	0.60
20:BA:73:HIS:ND1	20:BA:75:ASN:HB2	2.15	0.60
22:3L:19:G:HO2'	22:3L:61:U:H3	1.47	0.60
25:14:70:G:H21	25:14:71:A:H62	1.49	0.60
25:14:1647:G:H3'	25:14:1647:G:OP2	2.01	0.60
46:C5:84:ARG:HB2	46:C5:84:ARG:HH11	1.65	0.60
1:13:148:G:H1	1:13:174:C:H42	1.49	0.60
9:8E:42:ARG:NH1	9:8E:75:ASP:OD1	2.32	0.60
1:1G:728:A:H2'	1:1G:729:A:H8	1.66	0.60
25:14:2068:U:N3	25:14:2430:A:H2	1.96	0.60
3:2E:11:ARG:HH21	3:2E:180:ALA:HB3	1.65	0.60
25:1H:1605:C:O3'	62:1H:3668:HOH:O	2.16	0.60
1:1G:1157:A:H61	1:1G:1178:G:H21	1.48	0.60
3:22:71:ALA:HA	3:22:106:VAL:HB	1.84	0.60
57:2L:44:A:H2'	57:2L:45:A:H8	1.65	0.60
25:14:1634:A:OP1	62:14:3545:HOH:O	2.17	0.60
41:75:36:GLU:HB3	41:75:39:ARG:HH22	1.66	0.60
1:13:327:A:O2'	1:13:328:C:O4'	2.18	0.60
25:1H:2178:C:H5''	27:71:46:LYS:HD3	1.83	0.60
1:1G:67:C:H2'	1:1G:68:G:C8	2.37	0.60
1:1G:323:U:H5'	20:BA:23:ARG:HB2	1.83	0.60
8:72:82:HIS:NE2	8:72:136:GLU:OE2	2.34	0.60
25:14:2287:A:H62	25:14:2344:U:H3	1.47	0.60
25:14:2680:C:H1'	29:29:187:ALA:HB1	1.84	0.60
26:1J:95:U:H2'	26:1J:96:G:C8	2.36	0.60
28:19:175:LEU:HD12	28:19:185:VAL:HG21	1.82	0.60
29:29:12:THR:HG22	29:29:13:ARG:H	1.66	0.60
30:39:20:LEU:HD22	30:39:199:TRP:HZ3	1.66	0.60
43:95:71:LEU:HA	43:95:86:GLY:H	1.66	0.60
1:13:110:C:O2'	16:7I:25:ARG:O	2.20	0.60
1:13:191:G:O2'	20:BI:101:GLY:O	2.15	0.60
1:13:1259:C:N4	1:13:1260:C:O2	2.34	0.60
5:4E:77:PRO:HG3	5:4E:144:THR:HG22	1.84	0.60
15:6I:76:GLU:OE1	15:6I:79:ARG:NH1	2.35	0.60
25:1H:1169:G:H1	25:1H:1180:C:H42	1.50	0.60
25:1H:2847:U:OP1	41:B8:98:LYS:NZ	2.33	0.60
27:71:5:LYS:HZ1	27:71:8:ARG:HH22	1.50	0.60
33:61:2:LYS:HG3	33:61:20:ASP:HB3	1.82	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:61:57:ARG:HA	33:61:60:GLU:HB3	1.84	0.60
1:1G:976:G:O5'	1:1G:1358:U:O2'	2.19	0.60
36:25:105:GLU:OE1	36:25:105:GLU:N	2.32	0.60
1:13:1004:A:H8	1:13:1026:G:C5	2.19	0.60
14:5I:24:CYS:SG	14:5I:25:VAL:N	2.75	0.60
22:3K:28:U:H3	22:3K:45:A:H61	1.47	0.60
41:B8:29:ARG:HB2	41:B8:46:GLU:HB2	1.82	0.60
2:12:142:LEU:HD23	2:12:145:LEU:HB2	1.83	0.60
10:1A:38:ILE:HB	10:1A:71:LEU:HB3	1.83	0.60
25:14:271(B):G:H1	25:14:404:C:H42	1.48	0.60
28:19:16:MET:HG3	28:19:207:GLY:HA3	1.83	0.60
31:49:113:ARG:HG2	52:I5:35:VAL:HG23	1.84	0.60
1:13:1124:G:H3'	1:13:1145:C:N4	2.17	0.60
8:7E:85:ARG:NE	8:7E:87:SER:O	2.35	0.60
29:21:38:THR:N	29:21:42:ASP:OD2	2.32	0.60
16:7A:4:ILE:HG12	16:7A:21:VAL:HG12	1.84	0.60
25:14:2324:C:H5''	25:14:2325:G:H5'	1.83	0.60
44:A5:18:ARG:NH1	44:A5:76:VAL:O	2.35	0.60
1:13:346:G:H5''	41:B8:41:ARG:HH11	1.66	0.60
1:13:673:G:H2'	1:13:674:G:C8	2.36	0.60
25:1H:1448:G:O2'	25:1H:1529:A:N1	2.31	0.60
25:1H:2494:G:O2'	38:88:80:GLU:HB3	2.02	0.60
1:1G:1066:C:H3'	1:1G:1067:A:C8	2.37	0.60
1:1G:1290:G:H5'	7:62:38:LEU:HD11	1.83	0.60
17:8A:19:VAL:HG22	17:8A:44:ALA:HB3	1.84	0.60
25:14:1011:G:OP2	42:85:66:ASN:ND2	2.35	0.60
35:15:113:GLY:HA2	35:15:116:LEU:HD12	1.84	0.60
38:45:75:THR:HA	38:45:89:ASN:H	1.67	0.60
1:13:1028(A):C:H2'	1:13:1028(B):C:C5	2.37	0.60
1:13:1392:G:H21	1:13:1502:A:H8	1.49	0.60
22:3K:72:C:H2'	22:3K:73:A:C8	2.37	0.60
25:1H:1191:G:OP1	37:78:32:THR:OG1	2.20	0.60
25:1H:2216:G:H2'	25:1H:2217:G:H8	1.67	0.60
25:1H:2637:U:H5''	29:21:82:ARG:HH21	1.67	0.60
29:21:181:LEU:HD11	41:B8:6:LEU:HD21	1.83	0.60
1:1G:457:C:H2'	1:1G:458:C:C6	2.37	0.60
25:14:815:C:N4	25:14:1192:G:H1	1.99	0.60
25:14:833:U:H1'	37:35:55:ARG:HH11	1.65	0.60
25:14:1603:A:OP1	62:14:3542:HOH:O	2.15	0.60
1:13:1298:C:H4'	1:13:1299:A:C4	2.37	0.59
5:4E:110:LEU:HD13	5:4E:118:ILE:HG21	1.84	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:5I:9:LYS:HA	14:5I:12:ARG:HD3	1.83	0.59
17:8I:76:LEU:HD11	17:8I:79:SER:HB2	1.84	0.59
25:1H:2168:G:N2	25:1H:2169:A:H3'	2.16	0.59
44:E8:86:LEU:HD12	44:E8:87:PRO:HD2	1.83	0.59
47:H8:127:LYS:HB3	47:H8:162:GLU:HB3	1.83	0.59
54:O8:9:LEU:HD12	54:O8:26:ASN:HB3	1.83	0.59
4:32:187:ARG:NH2	4:32:193:ASP:OD2	2.36	0.59
9:82:43:ALA:HA	9:82:74:ILE:HD13	1.83	0.59
14:5A:12:ARG:HG2	14:5A:14:PRO:HD3	1.84	0.59
25:14:69:C:O2	25:14:73:A:O2'	2.17	0.59
25:14:247:G:N1	25:14:247:G:O6	2.35	0.59
31:49:60:LEU:HD13	31:49:68:PRO:HB3	1.82	0.59
1:13:390:C:H2'	1:13:391:G:C8	2.37	0.59
25:1H:110:G:O6	62:1H:3666:HOH:O	2.16	0.59
25:1H:271(B):G:H4'	25:1H:271(C):U:OP1	2.02	0.59
52:M8:38:LYS:HA	52:M8:44:THR:HB	1.84	0.59
1:1G:78:G:O6	1:1G:91:C:N4	2.36	0.59
1:1G:1001:G:H4'	1:1G:1001:G:OP1	2.02	0.59
1:1G:1289:A:OP1	21:1B:9:ARG:NH2	2.35	0.59
25:14:1316:U:H2'	25:14:1317:A:H8	1.65	0.59
25:14:1443:G:H1	25:14:1548:C:H42	1.49	0.59
25:14:2129:C:H3'	25:14:2130:U:H5''	1.83	0.59
46:C5:83:THR:OG1	46:C5:84:ARG:N	2.35	0.59
1:13:1023:G:H3'	1:13:1024:G:C5'	2.31	0.59
10:1I:27:ALA:HB2	10:1I:85:LEU:HD11	1.84	0.59
25:1H:529:A:H4'	25:1H:530:G:H5'	1.84	0.59
25:1H:581:C:H2'	25:1H:582:G:C8	2.37	0.59
25:1H:848:G:H2'	25:1H:849:A:C8	2.37	0.59
25:1H:1532:C:N3	25:1H:1539:G:N2	2.49	0.59
25:1H:2030:A:H4'	25:1H:2031:A:H8	1.66	0.59
46:G8:84:ARG:HH11	46:G8:84:ARG:HB2	1.67	0.59
2:12:72:GLY:HA3	2:12:81:VAL:HG11	1.84	0.59
25:14:2403:C:C4	25:14:2403:C:C6	2.90	0.59
41:75:28:VAL:HB	41:75:86:ILE:HD11	1.84	0.59
42:85:98:LEU:HB2	42:85:106:PHE:HD2	1.67	0.59
1:13:312:C:H2'	1:13:313:A:H8	1.67	0.59
1:13:1113:C:H2'	1:13:1114:C:H6	1.67	0.59
9:8E:13:ALA:HB2	9:8E:68:GLY:HA3	1.83	0.59
25:1H:330:A:H2	25:1H:1210:A:H2'	1.68	0.59
25:1H:821:A:H5'	25:1H:822:U:H6	1.66	0.59
25:1H:1301:A:H2'	25:1H:1302:A:H3'	1.84	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:E8:76:VAL:HB	44:E8:103:ILE:HG23	1.84	0.59
1:1G:1051:C:N3	1:1G:1207:2MG:N1	2.41	0.59
1:1G:1114:C:O2'	14:5A:60:SER:O	2.10	0.59
7:62:15:ASP:HB3	7:62:24:THR:HB	1.84	0.59
13:4A:45:VAL:O	13:4A:47:ASP:N	2.35	0.59
17:8A:90:ILE:O	17:8A:94:ASN:ND2	2.36	0.59
25:14:300:A:H1'	25:14:319:C:H1'	1.85	0.59
25:14:1636:C:H2'	25:14:1637:A:C8	2.38	0.59
1:13:831:U:H2'	1:13:832:C:H6	1.66	0.59
1:13:1008:C:N4	1:13:1021:G:O6	2.33	0.59
25:1H:1057:A:H62	25:1H:1086:A:H2'	1.66	0.59
27:71:20:TYR:HB2	27:71:224:ILE:HA	1.82	0.59
9:82:13:ALA:HB2	9:82:68:GLY:HA3	1.85	0.59
41:75:120:ARG:HA	41:75:123:GLN:HG2	1.84	0.59
46:C5:44:ILE:HA	46:C5:64:GLU:HA	1.85	0.59
51:H5:5:LYS:HE3	51:H5:7:LYS:HE3	1.85	0.59
1:13:1133:G:N2	1:13:1141:C:N3	2.43	0.59
2:1E:21:ARG:HB2	2:1E:39:ILE:HG13	1.85	0.59
3:2E:53:ALA:HB2	3:2E:115:LEU:HD21	1.85	0.59
4:3E:173:TRP:CD2	4:3E:189:PRO:HB3	2.37	0.59
17:8I:86:GLU:O	17:8I:90:ILE:HG12	2.02	0.59
32:51:54:ARG:NH2	32:51:57:ASP:OD2	2.36	0.59
1:1G:186(A):C:H5''	20:BA:86:ARG:HG3	1.85	0.59
1:1G:662:G:H1	1:1G:743:U:H3	1.49	0.59
3:22:161:GLU:CD	3:22:162:GLN:H	2.11	0.59
10:1A:8:LEU:HB2	10:1A:16:LEU:HD11	1.85	0.59
25:14:458:G:N2	25:14:470:A:OP2	2.36	0.59
25:14:670:A:H5''	37:35:42:SER:OG	2.02	0.59
25:14:2787:C:H1'	29:29:62:PRO:HG3	1.84	0.59
51:H5:12:PRO:O	51:H5:20:LYS:NZ	2.35	0.59
1:13:191:G:H1'	20:BI:105:SER:HB3	1.85	0.59
1:13:661:G:H1	1:13:744:C:H42	1.50	0.59
2:1E:47:THR:HG23	2:1E:202:PRO:HD2	1.84	0.59
25:1H:573:G:N1	25:1H:2031:A:OP2	2.29	0.59
25:1H:1981:A:OP2	62:1H:3669:HOH:O	2.16	0.59
34:38:25:PHE:HD1	34:38:26:LEU:HG	1.68	0.59
1:1G:553:A:O2'	12:3A:26:GLY:O	2.20	0.59
1:1G:971:G:H4'	1:1G:972:C:H5''	1.83	0.59
8:72:10:LEU:HD22	8:72:83:ILE:HD11	1.84	0.59
28:19:264:LYS:HG2	28:19:266:SER:H	1.68	0.59
40:65:34:HIS:O	40:65:97:ARG:NH2	2.36	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:281:G:H8	1:13:281:G:OP2	1.84	0.59
1:13:321:A:N6	1:13:329:A:OP2	2.35	0.59
1:13:1255:G:H21	1:13:1259:C:H1'	1.66	0.59
1:13:1427:U:H2'	1:13:1428:A:H8	1.68	0.59
5:4E:12:LEU:HD13	5:4E:128:PRO:HG2	1.85	0.59
13:4I:96:LEU:HD13	13:4I:97:PRO:HD2	1.84	0.59
25:1H:2698:U:H2'	25:1H:2699:C:C6	2.37	0.59
30:31:178:PRO:HB2	30:31:201:VAL:HG11	1.85	0.59
50:K8:50:ILE:HD12	50:K8:51:ARG:H	1.68	0.59
1:1G:500:G:H1	1:1G:545:C:H42	1.50	0.59
1:1G:974:A:OP2	14:5A:29:ARG:NH2	2.36	0.59
25:14:662:G:H5'	37:35:15:ARG:HB2	1.84	0.59
38:45:1:MET:SD	38:45:1:MET:N	2.75	0.59
8:7E:83:ILE:HB	8:7E:137:VAL:HG13	1.85	0.59
22:1K:38:A:O2'	25:1H:1913:A:N1	2.35	0.59
25:1H:1257:C:H4'	30:31:83:PHE:CD1	2.38	0.59
25:1H:1900:A:OP2	62:1H:3671:HOH:O	2.17	0.59
26:16:10:C:H42	26:16:110:G:H1	1.51	0.59
33:61:129:THR:HA	33:61:137:PRO:HA	1.85	0.59
47:H8:155:LEU:H	47:H8:155:LEU:HD22	1.67	0.59
1:1G:250:A:H4'	1:1G:251:G:O5'	2.02	0.59
1:1G:1070:U:OP1	5:42:18:ARG:NH2	2.35	0.59
13:4A:23:TYR:HB3	13:4A:71:ARG:HD2	1.84	0.59
25:14:273(C):C:N4	25:14:363(C):G:H1	2.00	0.59
25:14:708:C:N4	25:14:723:G:H1	2.00	0.59
25:14:848:G:H2'	25:14:849:A:C8	2.38	0.59
25:14:1689:A:H62	25:14:1698:A:H2	1.51	0.59
1:13:328:C:H4'	1:13:329:A:H5'	1.85	0.59
1:13:406:G:N2	1:13:437:U:O2	2.36	0.59
1:13:1459:C:OP1	20:BI:27:LYS:NZ	2.34	0.59
1:13:1504:G:H4'	1:13:1505:G:O5'	2.02	0.59
25:1H:860:U:H5	25:1H:917:A:H2	1.51	0.59
27:71:21:THR:H	27:71:24:GLU:HB2	1.68	0.59
1:1G:261:U:OP2	20:BA:79:ARG:NH2	2.35	0.59
1:1G:328:C:H4'	1:1G:329:A:H5'	1.84	0.59
1:1G:1049:U:H4'	1:1G:1050:G:C5'	2.33	0.59
2:12:80:ILE:HD13	2:12:212:GLN:HG2	1.85	0.59
2:12:82:ARG:NH2	2:12:92:TYR:OH	2.35	0.59
9:82:28:VAL:HG22	9:82:29:ASN:H	1.68	0.59
26:1J:45:A:O5'	31:49:95:ARG:NH1	2.36	0.59
25:1H:1407:C:H42	25:1H:1595:G:H1	1.49	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:2848:G:C8	41:B8:97:ALA:HB2	2.37	0.58
27:71:186:ALA:HA	27:71:189:ILE:HD12	1.85	0.58
33:61:91:SER:HB3	33:61:121:LYS:HE2	1.84	0.58
54:O8:7:ILE:O	54:O8:8:LYS:HB3	2.02	0.58
9:82:82:ALA:HA	9:82:85:LEU:HG	1.85	0.58
1:13:596:C:H2'	1:13:597:G:H8	1.68	0.58
25:1H:2114:A:H2'	25:1H:2168:G:H8	1.68	0.58
39:98:103:ARG:HH21	39:98:110:PRO:HB3	1.68	0.58
56:Q8:11:LYS:HB2	56:Q8:62:LEU:HD21	1.84	0.58
1:1G:145:G:H2'	1:1G:146:G:C8	2.38	0.58
1:1G:1239:A:H4'	1:1G:1240:U:H5''	1.85	0.58
1:1G:1244:C:H2'	1:1G:1245:A:C8	2.38	0.58
4:32:105:VAL:HG13	4:32:110:PHE:HB2	1.84	0.58
25:14:1550:C:OP1	25:14:1727:U:O2'	2.13	0.58
25:14:2186:G:H2'	25:14:2187:G:H8	1.68	0.58
1:13:973:G:H3'	1:13:974:A:H5''	1.85	0.58
1:13:1435:G:H2'	1:13:1436:U:C6	2.39	0.58
10:1I:26:ALA:HA	10:1I:29:ARG:HH21	1.67	0.58
25:1H:270(H):C:H2'	25:1H:270(I):G:O4'	2.02	0.58
35:58:96:GLU:O	35:58:99:LEU:N	2.34	0.58
1:1G:401:C:H2'	1:1G:402:G:C8	2.39	0.58
1:1G:662:G:H2'	1:1G:663:A:H8	1.68	0.58
1:1G:737:A:H2'	1:1G:738:C:C6	2.38	0.58
25:14:1849:G:H2'	25:14:1850:G:H8	1.68	0.58
43:95:70:ILE:HB	43:95:86:GLY:HA3	1.84	0.58
47:D5:24:LEU:HB3	47:D5:41:LEU:HD12	1.84	0.58
1:13:537:G:OP2	62:13:1807:HOH:O	2.17	0.58
1:13:1292:U:OP1	7:6E:41:ARG:NH2	2.36	0.58
1:13:1314:C:OP2	19:AI:4:SER:OG	2.21	0.58
13:4I:81:LEU:HD22	13:4I:88:ARG:HB3	1.85	0.58
15:6I:74:ASP:OD1	15:6I:77:ARG:N	2.30	0.58
16:7I:39:TYR:HD1	16:7I:49:LEU:HD13	1.66	0.58
22:3K:67:C:H2'	22:3K:68:C:C6	2.39	0.58
25:1H:620:G:H4'	25:1H:621:A:H5''	1.85	0.58
25:1H:1021:A:OP2	35:58:65:LYS:NZ	2.28	0.58
25:1H:1550:C:H2'	25:1H:1551:C:H6	1.67	0.58
37:78:71:VAL:HG13	37:78:72:PRO:HD3	1.86	0.58
38:88:34:LEU:HB2	38:88:118:LEU:HD22	1.85	0.58
7:62:69:VAL:HG22	7:62:135:VAL:HG22	1.86	0.58
28:19:44:ASN:ND2	28:19:46:GLN:OE1	2.37	0.58
43:95:21:ARG:HA	43:95:93:GLU:HA	1.85	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1065:U:C5	1:13:1190:G:H1'	2.39	0.58
2:1E:168:THR:HG21	2:1E:191:ASP:HB3	1.84	0.58
8:7E:42:GLU:HG3	8:7E:109:ILE:HD12	1.84	0.58
25:1H:1697:G:OP2	25:1H:1698:A:O2'	2.17	0.58
25:1H:2439:A:H5'	25:1H:2439:A:C8	2.39	0.58
31:41:62:LEU:HD21	52:M8:28:LYS:HD2	1.86	0.58
33:61:66:GLU:HA	33:61:69:LYS:HB3	1.86	0.58
4:32:62:GLN:HB3	4:32:66:ARG:HD2	1.85	0.58
16:7A:22:THR:HA	16:7A:33:ILE:HG12	1.85	0.58
18:9A:62:GLU:HA	18:9A:65:ILE:HD11	1.85	0.58
25:14:1171:G:O2'	25:14:1173:G:O5'	2.21	0.58
25:14:2882:A:O5'	39:55:96:ARG:NE	2.34	0.58
29:29:14:ILE:HG13	29:29:21:VAL:HG13	1.84	0.58
29:29:115:GLY:O	29:29:119:ARG:HB2	2.03	0.58
2:1E:87:ARG:NH1	2:1E:220:ASP:OD1	2.25	0.58
25:1H:2232:U:P	49:J8:40:ARG:HH12	2.25	0.58
26:16:38:C:H42	26:16:44:G:H1	1.50	0.58
30:31:152:GLU:HB3	30:31:190:GLU:HB2	1.85	0.58
32:51:7:LEU:N	32:51:8:PRO:HD3	2.19	0.58
35:58:60:ILE:H	35:58:60:ILE:HD13	1.68	0.58
42:C8:83:LEU:HD12	42:C8:88:ILE:HD12	1.85	0.58
1:1G:1166:G:N2	1:1G:1170:A:OP2	2.34	0.58
3:22:38:ARG:HD2	3:22:94:LEU:HD22	1.86	0.58
4:32:201:GLN:NE2	5:42:116:THR:O	2.37	0.58
12:3A:124:GLU:OE1	12:3A:124:GLU:N	2.27	0.58
20:BA:18:GLN:HA	20:BA:21:LYS:HE3	1.86	0.58
25:14:1332:G:H21	25:14:1610:A:H8	1.49	0.58
25:14:1997:G:H5''	62:14:3518:HOH:O	2.04	0.58
25:14:2443:C:H2'	25:14:2444:G:H8	1.68	0.58
1:13:1139:G:H4'	1:13:1140:C:H5'	1.86	0.58
1:13:1446:A:O2'	1:13:1447:G:O5'	2.20	0.58
4:3E:142:PRO:HA	4:3E:185:PHE:HD2	1.69	0.58
18:9I:22:VAL:HG13	18:9I:42:ARG:HH11	1.68	0.58
25:1H:1187:G:H5''	43:D8:81:TYR:CE1	2.39	0.58
25:1H:2298:A:H2'	25:1H:2299:G:O4'	2.03	0.58
1:1G:147:G:H2'	1:1G:148:G:C8	2.37	0.58
19:AA:40:ILE:HG13	19:AA:66:MET:HB3	1.85	0.58
21:1B:8:THR:HG22	21:1B:10:ARG:H	1.69	0.58
29:29:68:ALA:HA	29:29:71:GLY:HA2	1.86	0.58
29:29:128:SER:OG	29:29:129:HIS:N	2.37	0.58
5:4E:142:LEU:O	5:4E:143:ARG:NH1	2.35	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:68:17:ARG:HA	36:68:17:ARG:HH11	1.69	0.58
1:1G:1049:U:H4'	1:1G:1050:G:H5''	1.86	0.58
5:42:71:LEU:HD22	5:42:115:VAL:HG13	1.85	0.58
7:62:20:ASP:O	7:62:22:LEU:N	2.36	0.58
14:5A:45:ARG:O	14:5A:49:HIS:ND1	2.32	0.58
25:14:730:C:H3'	62:14:3527:HOH:O	2.03	0.58
25:14:2129:C:O2'	25:14:2160:G:N2	2.37	0.58
25:14:2287:A:N6	25:14:2344:U:H3	2.02	0.58
29:29:120:TRP:CG	29:29:155:LYS:HB3	2.39	0.58
32:59:12:PRO:HB3	32:59:48:GLY:HA3	1.86	0.58
38:45:74:TYR:O	38:45:89:ASN:HA	2.04	0.58
1:13:1372:U:H5''	9:8E:71:SER:HB3	1.84	0.58
22:3K:29:C:H2'	22:3K:30:G:H8	1.68	0.58
22:3K:35:C:H41	24:4K:14:A:H61	1.52	0.58
25:1H:67:U:H3	25:1H:74:A:H2	1.52	0.58
25:1H:1268:A:OP1	62:1H:3672:HOH:O	2.17	0.58
47:H8:108:PRO:HD2	47:H8:111:VAL:HA	1.86	0.58
1:1G:1157:A:N6	1:1G:1178:G:H21	2.01	0.58
25:14:856:C:H1'	48:E5:27:GLU:HB3	1.85	0.58
25:14:883:G:H22	25:14:893:C:N4	2.02	0.58
25:14:2119:A:N7	25:14:2170:A:N6	2.52	0.58
30:39:128:ALA:O	30:39:142:TRP:NE1	2.36	0.58
31:49:97:ASP:HA	31:49:100:TRP:CD1	2.37	0.58
1:13:753:A:H4'	1:13:754:C:O5'	2.03	0.58
1:13:826:C:H4'	8:7E:12:ARG:HG3	1.84	0.58
1:13:1081:G:H2'	1:13:1082:G:C8	2.38	0.58
5:4E:79:GLU:HB3	5:4E:92:LYS:HA	1.86	0.58
25:1H:2860:A:C8	25:1H:2861:G:H1'	2.39	0.58
30:31:103:LYS:HA	30:31:106:ARG:HG3	1.86	0.58
35:58:35:ARG:HD3	35:58:37:LYS:HD2	1.86	0.58
4:32:57:ARG:HB3	4:32:206:PHE:HB2	1.85	0.58
6:52:82:ARG:HB2	6:52:85:VAL:HG23	1.86	0.58
25:14:448:U:O4	25:14:583:G:H1'	2.04	0.58
25:14:2300:G:O6	25:14:2316:C:N4	2.36	0.58
25:14:2650:U:H2'	25:14:2651:C:H6	1.69	0.58
26:1J:80:U:H2'	26:1J:81:G:H21	1.67	0.58
26:1J:83:G:H1	26:1J:93:C:H42	1.51	0.58
37:35:79:ARG:HE	37:35:109:GLY:HA3	1.68	0.58
55:L5:9:ARG:HG3	55:L5:46:VAL:HG22	1.85	0.58
1:13:1292:U:H2'	1:13:1293:G:C8	2.39	0.57
25:1H:534:U:H5'	42:C8:42:ALA:HB1	1.85	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:636:G:N7	37:78:113:LYS:HE2	2.19	0.57
25:1H:810:U:O4	62:1H:3667:HOH:O	2.16	0.57
30:31:29:ASN:HB3	30:31:112:MET:HE1	1.85	0.57
31:41:112:PRO:HG3	52:M8:37:SER:O	2.04	0.57
43:D8:15:GLU:HG3	43:D8:16:PRO:HD2	1.85	0.57
1:1G:503:C:O2'	1:1G:510:A:N1	2.35	0.57
38:45:17:LEU:HD21	38:45:41:TRP:HE1	1.69	0.57
1:13:324:G:O2'	1:13:326:G:N7	2.37	0.57
1:13:750:G:O2'	15:6I:21:ASP:OD1	2.22	0.57
22:3K:9:G:O2'	22:3K:10:G:N7	2.37	0.57
25:1H:70:G:H21	25:1H:71:A:H62	1.50	0.57
25:1H:1423:G:H2'	25:1H:1424:G:H8	1.69	0.57
40:A8:67:ARG:NH2	40:A8:103:GLU:OE2	2.37	0.57
1:1G:567:G:O2'	62:1G:1808:HOH:O	2.17	0.57
1:1G:729:A:H2'	1:1G:730:G:H8	1.68	0.57
1:1G:1106:G:H5''	3:22:172:ARG:HG2	1.85	0.57
1:1G:1259:C:N4	1:1G:1260:C:O2	2.37	0.57
1:1G:1327:C:H2'	1:1G:1328:C:C6	2.39	0.57
7:62:36:LYS:HA	7:62:39:ALA:HB3	1.85	0.57
19:AA:39:THR:HA	19:AA:70:LYS:HD2	1.86	0.57
25:14:792:G:O2'	25:14:2440:C:N3	2.31	0.57
25:14:2023:G:OP2	25:14:2617:C:H4'	2.04	0.57
29:29:1:MET:HA	29:29:200:GLU:HB3	1.85	0.57
22:3K:57:C:H2'	22:3K:58:A:C8	2.37	0.57
27:71:27:HIS:HE1	27:71:183:GLU:HB3	1.68	0.57
33:61:113:ARG:HB3	33:61:131:LYS:HD3	1.86	0.57
41:B8:105:LEU:C	41:B8:107:ASP:H	2.12	0.57
47:H8:137:ILE:HB	47:H8:156:LYS:HE3	1.85	0.57
1:1G:814:A:O2'	1:1G:815:A:H3'	2.03	0.57
1:1G:932:C:H5'	7:62:4:ARG:HG2	1.86	0.57
25:14:451:C:C5	25:14:453:C:H5''	2.39	0.57
25:14:2121:G:H22	25:14:2177:C:N4	2.00	0.57
25:14:2577:A:H4'	53:J5:3:LYS:H	1.69	0.57
32:59:7:LEU:N	32:59:8:PRO:HD3	2.19	0.57
32:59:27:LYS:H	32:59:32:GLU:HB2	1.69	0.57
41:75:123:GLN:HA	41:75:126:ALA:HB3	1.87	0.57
42:85:90:VAL:HA	43:95:39:LEU:HD22	1.86	0.57
52:I5:15:ILE:HG13	52:I5:30:GLU:HB2	1.86	0.57
1:13:7:G:OP1	4:3E:209:ARG:NH2	2.37	0.57
1:13:342:C:H2'	1:13:343:U:O4'	2.04	0.57
25:1H:107:C:H2'	25:1H:108:U:H6	1.69	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:2210:G:H3'	25:1H:2211:G:C8	2.40	0.57
25:1H:2415:G:H1'	37:78:67:MET:HE3	1.86	0.57
33:61:116:LEU:HD11	33:61:119:PRO:HA	1.85	0.57
1:1G:390:C:H2'	1:1G:391:G:C8	2.39	0.57
1:1G:643:C:H2'	1:1G:644:G:H8	1.68	0.57
1:1G:979:C:OP1	1:1G:1223:C:N4	2.37	0.57
5:42:131:ILE:O	5:42:135:THR:OG1	2.22	0.57
7:62:9:VAL:HG22	7:62:94:ARG:HH11	1.69	0.57
22:3L:21:U:H5''	22:3L:22:A:H5''	1.86	0.57
26:1J:32:C:N3	26:1J:51:G:N2	2.52	0.57
38:45:64:ILE:H	38:45:64:ILE:HD13	1.70	0.57
1:13:280:C:N3	17:8I:39:SER:N	2.53	0.57
1:13:720:C:H5''	18:9I:52:PRO:HA	1.85	0.57
1:13:1081:G:H2'	1:13:1082:G:H8	1.70	0.57
8:7E:64:LYS:HG2	8:7E:79:VAL:HG21	1.86	0.57
19:AI:30:LEU:H	19:AI:30:LEU:HD13	1.69	0.57
22:1K:23:G:N7	22:1K:47:G:N2	2.42	0.57
25:1H:878:A:H3'	25:1H:879:G:H8	1.70	0.57
25:1H:2233:U:H2'	25:1H:2234:G:C8	2.40	0.57
31:41:32:PRO:HB2	31:41:172:LEU:HD13	1.86	0.57
34:38:53:VAL:HG13	34:38:54:ALA:HB3	1.86	0.57
1:1G:145:G:H2'	1:1G:146:G:H8	1.70	0.57
1:1G:765:G:N2	1:1G:813:U:OP2	2.37	0.57
1:1G:868:C:H2'	1:1G:869:G:O4'	2.04	0.57
1:1G:1128:C:N3	1:1G:1139:G:N2	2.53	0.57
1:1G:1305:G:OP1	21:1B:2:GLY:N	2.38	0.57
25:14:2656:U:C4	25:14:2665:A:H2	2.22	0.57
26:1J:15:A:H1'	26:1J:109:G:H1'	1.87	0.57
5:4E:43:LEU:H	5:4E:65:ASN:HB2	1.69	0.57
25:1H:1991:U:H2'	25:1H:1992:G:H5''	1.86	0.57
25:1H:2666:C:O2	32:51:152:ARG:NH2	2.37	0.57
36:68:2:ILE:HG13	36:68:6:THR:HG21	1.86	0.57
37:78:81:GLN:NE2	37:78:106:LEU:O	2.37	0.57
1:1G:1241:G:H1	1:1G:1296:C:H42	1.53	0.57
1:1G:1259:C:O2	1:1G:1283:G:O2'	2.20	0.57
13:4A:60:VAL:HG13	13:4A:64:TRP:HE1	1.70	0.57
21:1B:3:LYS:HD2	21:1B:14:TRP:HD1	1.70	0.57
29:29:8:LYS:NZ	29:29:188:VAL:O	2.34	0.57
10:1I:50:ILE:HD12	10:1I:57:LYS:HA	1.85	0.57
25:1H:177:G:H3'	25:1H:178:G:C8	2.40	0.57
37:78:59:LEU:HD11	56:Q8:10:ALA:HA	1.87	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:A8:61:ASN:O	40:A8:64:GLU:N	2.34	0.57
54:O8:41:PRO:HD2	54:O8:46:HIS:H	1.70	0.57
1:1G:1071:C:H2'	1:1G:1072:G:C8	2.40	0.57
1:1G:1086:U:H3'	1:1G:1087:G:C8	2.40	0.57
2:12:184:VAL:HG12	2:12:197:VAL:HG13	1.86	0.57
12:3A:73:ASN:ND2	12:3A:103:ASP:O	2.36	0.57
13:4A:3:ARG:NH2	31:49:113:ARG:O	2.30	0.57
25:14:666:G:H5''	37:35:47:ASP:O	2.04	0.57
25:14:1204:A:O2'	25:14:1205:U:OP2	2.23	0.57
26:1J:13:A:N1	26:1J:69:G:O2'	2.38	0.57
37:35:99:LEU:HD12	37:35:102:ARG:HE	1.68	0.57
38:45:103:MET:HB2	38:45:104:PHE:HD1	1.69	0.57
1:13:243:A:H4'	1:13:244:U:H3'	1.87	0.57
1:13:1373:G:O3'	7:6E:36:LYS:NZ	2.37	0.57
25:1H:270:A:OP2	25:1H:270(Y):G:N1	2.33	0.57
25:1H:1013:C:H42	25:1H:1149:G:H1	1.51	0.57
25:1H:1348:G:H2'	25:1H:1349:A:H5''	1.86	0.57
25:1H:2059:A:H5'	25:1H:2060:A:OP2	2.05	0.57
25:1H:2062:A:OP2	62:1H:3673:HOH:O	2.17	0.57
25:1H:2397:G:H5''	49:J8:28:GLY:HA2	1.85	0.57
1:1G:920:U:H2'	1:1G:921:U:C6	2.40	0.57
2:12:30:ARG:O	2:12:46:LYS:NZ	2.36	0.57
15:6A:67:LEU:HD11	15:6A:87:ILE:HD12	1.85	0.57
25:14:380:U:H5'	49:F5:18:ILE:HD12	1.87	0.57
25:14:2667:C:N3	32:59:110:SER:OG	2.33	0.57
30:39:104:LYS:O	30:39:108:LYS:HG2	2.04	0.57
36:25:120:GLU:OE1	41:75:67:SER:OG	2.22	0.57
43:95:49:THR:HB	43:95:50:PRO:HD3	1.85	0.57
1:13:812:C:H4'	1:13:813:U:H5'	1.86	0.57
1:13:1011:G:H2'	1:13:1012:U:O4'	2.05	0.57
25:1H:163:U:C4	49:J8:81:LYS:HE3	2.38	0.57
25:1H:196:A:OP2	37:78:46:LYS:NZ	2.38	0.57
25:1H:370:G:OP2	62:1H:3676:HOH:O	2.18	0.57
30:31:63:LYS:NZ	30:31:75:HIS:O	2.38	0.57
1:1G:474:G:H2'	1:1G:475:G:C8	2.40	0.57
1:1G:476:G:H2'	1:1G:477:G:C8	2.40	0.57
1:1G:971:G:C6	1:1G:1365:G:H5'	2.40	0.57
5:42:12:LEU:HD13	5:42:13:ILE:H	1.70	0.57
9:82:20:ARG:O	9:82:60:ASP:N	2.37	0.57
10:1A:40:LEU:HB3	10:1A:69:ASN:HB2	1.85	0.57
22:3L:42:C:H2'	22:3L:43:G:H8	1.69	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:527:C:N4	25:14:2779:U:OP2	2.37	0.57
25:14:1300:U:H4'	25:14:1301:A:H5''	1.87	0.57
25:14:1762:A:H4'	25:14:1763:G:OP2	2.05	0.57
25:14:2011:U:OP1	44:A5:42:ARG:NH1	2.38	0.57
1:13:1073:U:H2'	1:13:1074:G:H8	1.69	0.57
25:1H:833:U:H1'	37:78:55:ARG:HH11	1.70	0.57
31:41:60:LEU:O	31:41:64:THR:HG22	2.04	0.57
33:61:88:ILE:HB	33:61:121:LYS:HG3	1.86	0.57
40:A8:106:ARG:O	40:A8:106:ARG:NH1	2.33	0.57
47:H8:156:LYS:HG2	47:H8:158:PRO:HD3	1.87	0.57
1:1G:1317:C:O2	19:AA:37:ARG:NH2	2.37	0.57
4:32:141:ARG:N	4:32:144:ASP:OD2	2.37	0.57
21:1B:3:LYS:HD2	21:1B:14:TRP:CD1	2.40	0.57
25:14:946:G:H2'	25:14:947:G:H8	1.69	0.57
1:13:272:C:H2'	1:13:273:A:C8	2.40	0.56
1:13:392:G:H2'	1:13:393:A:H8	1.69	0.56
1:13:538:G:H2'	1:13:539:A:H8	1.70	0.56
25:1H:270(M):U:H1'	25:1H:270(N):G:C5	2.39	0.56
25:1H:517:C:OP1	53:N8:16:ARG:NH2	2.38	0.56
25:1H:1800:C:OP2	28:11:183:ARG:NH1	2.38	0.56
34:38:4:LYS:O	34:38:7:VAL:N	2.38	0.56
41:B8:65:LYS:NZ	41:B8:66:VAL:O	2.38	0.56
47:H8:77:ASP:OD1	47:H8:80:ARG:N	2.34	0.56
1:1G:1218:C:P	14:5A:9:LYS:HZ3	2.28	0.56
1:1G:1268:A:H2'	1:1G:1269:A:C8	2.38	0.56
11:2A:92:GLU:OE2	11:2A:96:ARG:NH1	2.38	0.56
25:14:2393:A:H5'	37:35:63:PRO:HB3	1.85	0.56
25:14:2882:A:O4'	39:55:96:ARG:NE	2.38	0.56
38:45:20:ALA:HA	38:45:99:PRO:HG2	1.86	0.56
41:75:36:GLU:O	41:75:39:ARG:NH2	2.38	0.56
52:I5:22:ILE:HG12	52:I5:23:GLU:H	1.70	0.56
25:1H:587:C:N3	37:78:33:ARG:NH1	2.53	0.56
39:98:38:VAL:HG22	39:98:112:ALA:HB2	1.87	0.56
39:98:42:LYS:HA	39:98:45:ARG:HD2	1.85	0.56
45:F8:24:GLY:O	45:F8:82:GLN:HA	2.05	0.56
51:L8:26:LEU:O	51:L8:35:ARG:NE	2.37	0.56
1:1G:750:G:N3	15:6A:23:GLY:HA3	2.20	0.56
13:4A:10:PRO:HA	13:4A:18:ALA:HB1	1.86	0.56
17:8A:55:ASP:OD1	17:8A:55:ASP:N	2.37	0.56
25:14:83:G:N2	25:14:102:G:O2'	2.37	0.56
25:14:1796:U:H2'	25:14:1797:C:C6	2.40	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:2347:C:OP1	54:K5:39:TYR:OH	2.21	0.56
25:14:2716:U:H2'	25:14:2717:G:H8	1.71	0.56
28:19:181:GLU:HB2	28:19:273:ARG:HB2	1.87	0.56
41:75:54:ARG:HA	41:75:59:THR:HG23	1.87	0.56
1:13:151:A:H62	1:13:170:U:H3	1.51	0.56
1:13:543:C:OP1	4:3E:14:ARG:HG3	2.05	0.56
1:13:1305:G:H21	1:13:1331:G:H2'	1.69	0.56
1:13:1367:C:OP1	9:8E:115:GLY:N	2.34	0.56
4:3E:49:ARG:H	4:3E:49:ARG:HD3	1.69	0.56
9:8E:93:ARG:HE	9:8E:97:LYS:HB2	1.70	0.56
25:1H:224:G:O6	25:1H:419:C:O2'	2.24	0.56
25:1H:535:C:O3'	42:C8:53:ARG:NH1	2.38	0.56
25:1H:1485:G:H1	25:1H:1504:C:N4	2.04	0.56
25:1H:1937:A:H1'	25:1H:1939:5MU:H72	1.88	0.56
25:1H:2155:G:H2'	25:1H:2156:G:H5'	1.88	0.56
25:1H:2708:G:H5'	39:98:68:ARG:HG3	1.88	0.56
28:11:126:GLN:OE1	28:11:129:ASN:ND2	2.35	0.56
37:78:126:VAL:HG13	37:78:145:PRO:HG2	1.86	0.56
47:H8:102:LEU:HD21	47:H8:124:ILE:HG22	1.86	0.56
1:1G:256:U:H3	1:1G:270:A:H61	1.53	0.56
1:1G:876:G:H4'	8:72:14:ARG:HH12	1.69	0.56
1:1G:1235:U:O2'	1:1G:1305:G:O5'	2.23	0.56
3:22:16:ARG:HH22	3:22:181:ASN:HA	1.68	0.56
9:82:34:ASN:O	9:82:36:TYR:N	2.38	0.56
25:14:82:G:H5'	25:14:296:C:H5''	1.87	0.56
25:14:907:U:OP1	38:45:24:GLY:N	2.38	0.56
26:1J:40:U:N3	26:1J:43:C:H5''	2.20	0.56
26:1J:109:G:H2'	26:1J:110:G:H8	1.71	0.56
28:19:160:GLY:H	28:19:196:VAL:HB	1.69	0.56
32:59:9:ILE:HG23	32:59:51:ARG:HG3	1.88	0.56
46:C5:5:MET:SD	46:C5:32:PRO:HA	2.45	0.56
1:13:35:G:O2'	12:3I:115:SER:O	2.19	0.56
1:13:188:U:H2'	1:13:189:U:H5''	1.86	0.56
10:1I:3:LYS:N	10:1I:74:ILE:O	2.39	0.56
25:1H:598:G:H1	25:1H:659:C:H42	1.53	0.56
25:1H:2119:A:H2	25:1H:2171:A:H1'	1.71	0.56
26:16:40:U:H3	26:16:43:C:H5''	1.71	0.56
29:21:53:PRO:HA	29:21:75:VAL:HG12	1.88	0.56
34:38:122:VAL:HA	34:38:126:ALA:HB3	1.86	0.56
35:58:54:VAL:HB	35:58:122:VAL:HG22	1.87	0.56
47:H8:69:THR:HG22	47:H8:90:VAL:HA	1.86	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:H8:79:ARG:HB3	47:H8:80:ARG:HD3	1.88	0.56
1:1G:59:A:H5''	1:1G:387:U:H5''	1.87	0.56
1:1G:518:C:H4'	1:1G:519:C:O5'	2.05	0.56
1:1G:616:G:H2'	1:1G:617:G:H8	1.69	0.56
1:1G:1055:A:N7	1:1G:1200:C:N4	2.53	0.56
1:1G:1317:C:N3	19:AA:37:ARG:NH1	2.53	0.56
25:14:74:A:H5''	25:14:75:G:O4'	2.04	0.56
25:14:2503:2MA:O2'	25:14:2505:G:OP2	2.19	0.56
32:59:88:LEU:H	32:59:164:TYR:HA	1.69	0.56
41:75:107:ASP:C	41:75:111:ARG:HH21	2.14	0.56
1:13:1286:A:H2'	1:13:1287:A:H4'	1.88	0.56
8:7E:122:ARG:O	8:7E:126:LYS:N	2.35	0.56
25:1H:270(F):U:H2'	25:1H:270(G):C:C6	2.40	0.56
25:1H:1329:U:H5''	25:1H:1330:C:H5	1.70	0.56
25:1H:2372:G:H4'	54:O8:46:HIS:CD2	2.40	0.56
33:61:9:LEU:HB3	33:61:12:LEU:O	2.06	0.56
1:1G:595:G:H4'	1:1G:596:C:C5'	2.36	0.56
2:12:71:VAL:HA	2:12:93:VAL:HB	1.88	0.56
9:82:58:HIS:HB3	9:82:59:PHE:CD1	2.40	0.56
25:14:895:U:H3'	25:14:896:A:H5'	1.85	0.56
25:14:1281:G:H5'	25:14:1282:U:OP2	2.05	0.56
25:14:1957:C:OP1	62:14:3546:HOH:O	2.18	0.56
25:14:2123:G:H2'	25:14:2124:G:H8	1.71	0.56
26:1J:66:A:N6	26:1J:108:C:OP2	2.38	0.56
41:75:64:ARG:HB2	41:75:73:GLU:HG2	1.87	0.56
1:13:59:A:N1	62:13:1825:HOH:O	2.32	0.56
1:13:1324:A:H4'	1:13:1362:C:H4'	1.87	0.56
2:1E:93:VAL:HG21	2:1E:101:MET:HE1	1.87	0.56
3:2E:83:ARG:HD2	3:2E:87:LEU:HD11	1.85	0.56
4:3E:196:LEU:HD13	4:3E:198:VAL:HB	1.88	0.56
18:9I:53:ARG:HH21	18:9I:60:ALA:N	2.04	0.56
25:1H:124:G:O6	62:1H:3679:HOH:O	2.18	0.56
25:1H:630:G:N2	25:1H:633:A:OP2	2.26	0.56
25:1H:2638:G:HO2'	25:1H:2639:A:H8	1.52	0.56
25:1H:2788:C:OP1	29:21:61:ARG:NH2	2.38	0.56
28:11:34:VAL:HG21	28:11:61:LEU:HB3	1.87	0.56
1:1G:392:G:H2'	1:1G:393:A:C8	2.41	0.56
5:42:12:LEU:HD13	5:42:13:ILE:HD13	1.88	0.56
5:42:92:LYS:N	5:42:119:LEU:O	2.34	0.56
10:1A:78:ASN:HD22	10:1A:80:LYS:HB3	1.69	0.56
25:14:451:C:H5	25:14:453:C:H5''	1.70	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:2032:G:N2	25:14:2572:A:OP2	2.38	0.56
28:19:69:ARG:HD3	28:19:105:ILE:HD11	1.86	0.56
44:A5:12:ILE:HG13	44:A5:42:ARG:HH12	1.71	0.56
46:C5:46:LYS:HA	46:C5:61:ILE:O	2.05	0.56
1:13:45:U:H2'	1:13:46:G:C8	2.40	0.56
5:4E:78:HIS:CE1	5:4E:142:LEU:HD23	2.41	0.56
17:8I:58:GLU:OE1	17:8I:75:ARG:NH1	2.39	0.56
25:1H:653:A:O2'	25:1H:654:A:OP1	2.24	0.56
25:1H:1021:A:C8	25:1H:1021:A:H3'	2.41	0.56
25:1H:1301:A:C2'	25:1H:1302:A:H3'	2.36	0.56
25:1H:1301:A:O2'	25:1H:1302:A:O5'	2.19	0.56
37:78:85:LEU:HA	37:78:88:LEU:HD22	1.88	0.56
39:98:28:LEU:HD23	39:98:34:ILE:HG12	1.88	0.56
44:E8:64:MET:HE3	44:E8:109:GLU:HG3	1.88	0.56
1:1G:404:U:H5'	4:32:122:ARG:HE	1.71	0.56
10:1A:22:LYS:O	10:1A:26:ALA:N	2.38	0.56
10:1A:34:VAL:HG22	10:1A:74:ILE:HA	1.87	0.56
19:AA:67:VAL:HG21	52:I5:59:PHE:CG	2.40	0.56
25:14:372:G:N2	25:14:401:A:OP2	2.34	0.56
25:14:607:U:OP1	30:39:103:LYS:N	2.35	0.56
25:14:2469:A:H2'	38:45:56:ARG:HE	1.71	0.56
33:69:38:LEU:HD12	33:69:38:LEU:H	1.70	0.56
41:75:125:ARG:HA	41:75:128:GLU:HB3	1.87	0.56
47:D5:40:ASP:OD1	47:D5:43:GLU:N	2.33	0.56
1:13:1414:U:H3	1:13:1486:G:H1	1.52	0.56
5:4E:43:LEU:HD21	5:4E:132:ALA:HB1	1.88	0.56
5:4E:147:ASP:N	5:4E:147:ASP:OD1	2.39	0.56
15:6I:4:THR:OG1	15:6I:6:GLU:OE2	2.23	0.56
25:1H:28:A:N6	25:1H:512:G:H1'	2.21	0.56
32:51:11:VAL:HG22	32:51:76:VAL:HG11	1.88	0.56
1:1G:539:A:OP2	12:3A:112:LYS:NZ	2.37	0.56
1:1G:636:U:H2'	1:1G:637:G:C8	2.41	0.56
1:1G:762:C:H2'	1:1G:763:G:C8	2.40	0.56
1:1G:860:A:H4'	8:72:75:ARG:HH12	1.69	0.56
1:1G:953:G:H5'	1:1G:965:A:H61	1.69	0.56
2:12:58:ILE:O	2:12:62:ALA:N	2.33	0.56
25:14:1033:U:H1'	25:14:1034:G:H5''	1.88	0.56
25:14:1864:U:OP1	25:14:2410:G:O2'	2.23	0.56
41:75:39:ARG:HG2	41:75:40:THR:H	1.70	0.56
45:B5:44:GLU:HG3	45:B5:51:VAL:HG23	1.87	0.56
46:C5:91:GLU:HG3	46:C5:92:ASN:HB3	1.88	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1301:U:O3'	13:4I:21:TYR:OH	2.21	0.56
6:5E:23:LYS:NZ	6:5E:42:GLU:OE2	2.39	0.56
13:4I:3:ARG:HA	13:4I:9:ILE:HG22	1.86	0.56
25:1H:221:A:O2'	25:1H:222:A:OP2	2.23	0.56
25:1H:484:C:H2'	25:1H:485:C:H6	1.70	0.56
25:1H:1204:A:O2'	25:1H:1205:U:O5'	2.23	0.56
25:1H:2189:U:H2'	25:1H:2190:G:C8	2.41	0.56
27:71:14:VAL:HG22	27:71:28:LEU:HD21	1.87	0.56
34:38:21:GLN:O	34:38:23:SER:N	2.39	0.56
1:1G:745:C:OP1	1:1G:851:G:O2'	2.17	0.56
1:1G:880:C:OP1	12:3A:5:ASN:ND2	2.38	0.56
1:1G:910:C:OP2	12:3A:18:LYS:NZ	2.35	0.56
1:1G:1346:A:H4'	1:1G:1347:G:O5'	2.06	0.56
2:12:32:ILE:HG23	2:12:33:TYR:H	1.71	0.56
2:12:166:ASP:HB2	2:12:205:ASP:HB2	1.88	0.56
13:4A:81:LEU:HA	13:4A:85:GLY:HA3	1.88	0.56
57:2L:4:G:H1	57:2L:70:C:H42	1.53	0.56
25:14:586:A:N1	25:14:809:G:O2'	2.31	0.56
25:14:773:U:C4'	28:19:47:GLY:HA3	2.35	0.56
25:14:910:A:H62	38:45:12:GLN:HA	1.70	0.56
25:14:1849:G:H2'	25:14:1850:G:C8	2.41	0.56
25:14:2646:C:OP2	25:14:2732:G:O2'	2.17	0.56
45:B5:53:LYS:HB3	45:B5:82:GLN:HB3	1.88	0.56
23:2K:52:C:H42	23:2K:64:G:H1	1.53	0.56
25:1H:1666:G:O6	62:1H:3670:HOH:O	2.16	0.56
25:1H:2494:G:H2'	25:1H:2495:G:H8	1.71	0.56
25:1H:2646:C:OP2	25:1H:2732:G:O2'	2.19	0.56
25:1H:2689:U:P	25:1H:2719:G:H22	2.29	0.56
34:38:20:ALA:HB1	34:38:86:PRO:HA	1.87	0.56
40:A8:109:GLY:O	40:A8:110:LEU:HD22	2.05	0.56
49:J8:53:VAL:HG12	49:J8:54:ALA:H	1.70	0.56
1:1G:1129:C:O2	1:1G:1143:G:N2	2.38	0.56
1:1G:1148:U:H1'	9:82:16:ARG:HH21	1.71	0.56
2:12:80:ILE:HD11	2:12:211:ILE:HG22	1.87	0.56
11:2A:21:ILE:HG12	11:2A:30:VAL:HG12	1.87	0.56
13:4A:65:LYS:HB3	13:4A:70:LEU:HB2	1.86	0.56
25:14:1218:C:H42	25:14:1231:G:H1	1.52	0.56
32:59:17:VAL:HG11	32:59:50:VAL:HG11	1.88	0.56
33:69:12:LEU:HG	33:69:19:VAL:HG11	1.86	0.56
37:35:101:VAL:HA	37:35:105:LEU:O	2.06	0.56
42:85:96:ALA:HA	42:85:98:LEU:HD12	1.87	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
52:I5:58:ARG:O	52:I5:62:ARG:N	2.39	0.56
1:13:643:C:H2'	1:13:644:G:C8	2.41	0.55
25:1H:363:G:H2'	25:1H:363(A):A:H8	1.71	0.55
25:1H:536:A:H61	25:1H:557:U:H3	1.54	0.55
25:1H:1292:U:H2'	25:1H:1293:C:C6	2.41	0.55
33:61:46:ALA:HB1	33:61:50:ARG:HH21	1.71	0.55
40:A8:62:LYS:HA	40:A8:65:VAL:HB	1.88	0.55
47:H8:24:LEU:HB2	47:H8:41:LEU:HG	1.88	0.55
1:1G:1008:C:O2'	1:1G:1009:G:OP1	2.23	0.55
5:42:71:LEU:HD11	5:42:114:GLY:HA3	1.88	0.55
25:14:270(V):G:H2'	25:14:270(W):G:C8	2.41	0.55
25:14:444:C:OP2	42:85:2:PRO:HD3	2.06	0.55
25:14:582:G:H2'	25:14:583:G:C8	2.42	0.55
25:14:840:C:H2'	25:14:841:A:C8	2.41	0.55
25:14:906:G:HO2'	38:45:67:ARG:HE	1.49	0.55
25:14:1011:G:H1	25:14:1150:C:H42	1.52	0.55
30:39:194:MET:HE3	30:39:199:TRP:CD1	2.40	0.55
47:D5:150:LEU:HG	47:D5:171:ILE:H	1.70	0.55
1:13:992:U:H3	1:13:1044:A:H62	1.53	0.55
1:13:1516:G:N2	1:13:1519:MA6:OP2	2.38	0.55
25:1H:1188:U:H4'	43:D8:79:VAL:HG22	1.87	0.55
27:71:59:ARG:HD2	27:71:164:ARG:HD2	1.89	0.55
41:B8:16:ARG:HD3	41:B8:19:LEU:HD11	1.86	0.55
51:L8:40:THR:HG22	51:L8:41:PRO:HD2	1.87	0.55
52:M8:16:CYS:SG	52:M8:33:VAL:HB	2.46	0.55
1:1G:812:C:N3	62:1G:1833:HOH:O	2.33	0.55
8:72:4:ASP:HB2	8:72:89:PRO:HG3	1.89	0.55
25:14:5:A:H2'	25:14:6:A:C8	2.42	0.55
25:14:242:G:N2	25:14:255:A:OP2	2.25	0.55
25:14:270(S):G:H1'	49:F5:78:LYS:HE3	1.88	0.55
25:14:946:G:H2'	25:14:947:G:C8	2.40	0.55
30:39:34:TRP:CE3	37:35:8:PRO:HB3	2.42	0.55
41:75:57:PHE:O	41:75:58:ASN:ND2	2.40	0.55
17:8I:45:HIS:HB2	17:8I:65:ILE:HD13	1.87	0.55
18:9I:52:PRO:HG2	18:9I:55:ARG:HG2	1.87	0.55
25:1H:1102:C:H2'	25:1H:1103:A:H5''	1.89	0.55
25:1H:2216:G:H2'	25:1H:2217:G:C8	2.42	0.55
30:31:167:ALA:HB1	30:31:173:VAL:HG11	1.88	0.55
55:P8:18:PHE:HA	55:P8:43:THR:HG21	1.88	0.55
1:1G:1288:A:H2'	1:1G:1289:A:H8	1.71	0.55
9:82:47:LEU:HB3	9:82:50:LEU:HB2	1.87	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:639:U:H2'	25:14:640:C:C6	2.42	0.55
25:14:2140:C:H2'	25:14:2141:G:C8	2.40	0.55
25:14:2537:U:H2'	25:14:2538:C:C6	2.41	0.55
48:E5:19:LYS:C	48:E5:20:ARG:HG2	2.31	0.55
1:13:332:G:H2'	1:13:333:G:H8	1.70	0.55
1:13:565:U:H5''	1:13:566:G:H2'	1.88	0.55
1:13:701:C:H1'	1:13:703:G:C4	2.42	0.55
25:1H:1303:G:OP1	62:1H:3674:HOH:O	2.18	0.55
25:1H:1315:C:OP2	62:1H:3617:HOH:O	2.18	0.55
26:16:15:A:H3'	26:16:16:G:H5'	1.88	0.55
26:16:24:G:O2'	26:16:25:A:O5'	2.25	0.55
31:41:64:THR:HG23	31:41:66:GLN:H	1.72	0.55
32:51:10:PRO:HB2	32:51:50:VAL:HG12	1.89	0.55
37:78:4:SER:OG	37:78:5:ASP:OD1	2.24	0.55
54:O8:8:LYS:O	54:O8:9:LEU:HB2	2.07	0.55
1:1G:794:A:H4'	1:1G:1521:G:O2'	2.05	0.55
1:1G:1129:C:H5'	1:1G:1130:A:H5'	1.88	0.55
1:1G:1150:U:O2'	10:1A:39:PRO:O	2.23	0.55
1:1G:1315:U:HO2'	1:1G:1360:A:HO2'	1.55	0.55
8:72:20:TYR:HE2	8:72:75:ARG:HD2	1.71	0.55
25:14:220:G:O2'	25:14:233:A:N3	2.38	0.55
25:14:1937:A:H1'	25:14:1939:5MU:H72	1.88	0.55
25:14:2015:A:OP1	44:A5:92:ARG:NH2	2.39	0.55
40:65:20:ARG:HG3	48:E5:49:LYS:HE2	1.88	0.55
46:C5:83:THR:HG21	46:C5:99:CYS:SG	2.47	0.55
1:13:34:C:H2'	1:13:35:G:C8	2.41	0.55
1:13:1054:C:C4	22:1K:35:C:H1'	2.42	0.55
10:1I:13:HIS:HB2	10:1I:68:HIS:CD2	2.41	0.55
18:9I:52:PRO:HB2	18:9I:54:ARG:HG2	1.88	0.55
25:1H:1053:C:N3	25:1H:1106:G:N2	2.47	0.55
25:1H:2169:A:N6	25:1H:2170:A:N1	2.55	0.55
25:1H:2415:G:H4'	37:78:67:MET:N	2.14	0.55
32:51:8:PRO:HB2	32:51:69:ARG:HH21	1.71	0.55
44:E8:13:SER:HB3	44:E8:16:LYS:HE3	1.89	0.55
1:1G:1001:G:H22	1:1G:1039:C:H42	1.54	0.55
1:1G:1145:C:H4'	1:1G:1146:A:H8	1.71	0.55
25:14:1384:A:O2'	25:14:1404:C:O2	2.22	0.55
25:14:1853:A:H2'	25:14:1854:A:C8	2.41	0.55
25:14:2456:C:H42	25:14:2495:G:H1	1.54	0.55
25:14:2629:A:H62	25:14:2895:U:H3	1.55	0.55
33:69:77:LEU:HD22	33:69:141:LYS:H	1.71	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:75:24:PRO:HA	41:75:49:VAL:HG13	1.88	0.55
41:75:108:ARG:HA	41:75:111:ARG:HE	1.71	0.55
47:D5:127:LYS:HB3	47:D5:162:GLU:HB2	1.87	0.55
52:I5:55:ARG:HH21	52:I5:56:VAL:HG12	1.71	0.55
1:13:302:G:H4'	12:3I:14:LYS:NZ	2.22	0.55
1:13:1118:C:H1'	1:13:1179:A:C5	2.42	0.55
4:3E:128:VAL:HA	4:3E:145:GLU:O	2.06	0.55
11:2I:18:ARG:O	11:2I:33:THR:N	2.40	0.55
25:1H:1824:G:OP1	28:11:52:ARG:NH1	2.40	0.55
25:1H:2751:G:H5'	32:51:4:ILE:HA	1.87	0.55
30:31:11:VAL:HB	30:31:18:ARG:HG3	1.89	0.55
33:61:39:ALA:HB1	33:61:44:LEU:HD13	1.88	0.55
34:38:87:VAL:HG11	34:38:91:LYS:HD3	1.88	0.55
1:1G:538:G:H2'	1:1G:539:A:C8	2.40	0.55
1:1G:1232:U:H5''	9:82:124:GLN:O	2.06	0.55
7:62:50:ILE:HG13	7:62:58:PRO:HB3	1.89	0.55
30:39:117:ARG:HG3	30:39:122:LYS:HB2	1.87	0.55
56:M5:35:GLN:NE2	56:M5:40:GLU:OE1	2.39	0.55
1:13:1113:C:H2'	1:13:1114:C:C6	2.41	0.55
7:6E:46:ALA:O	7:6E:50:ILE:HG12	2.07	0.55
10:1I:47:PHE:CZ	10:1I:63:PHE:HB2	2.42	0.55
25:1H:2129:C:OP1	27:71:6:ARG:NE	2.39	0.55
32:51:74:ASN:HA	32:51:77:LYS:HD3	1.88	0.55
4:32:19:LEU:HG	4:32:21:LEU:HG	1.89	0.55
4:32:150:GLU:HA	4:32:153:ARG:HG2	1.89	0.55
13:4A:39:ILE:HG13	13:4A:56:LEU:HD23	1.89	0.55
13:4A:94:ARG:HB3	13:4A:96:LEU:HG	1.87	0.55
15:6A:54:ARG:NH1	15:6A:58:MET:SD	2.79	0.55
25:14:7:G:H1	25:14:2896:C:N4	2.00	0.55
25:14:2705:A:O2'	25:14:2852:G:OP1	2.20	0.55
40:65:10:ARG:NH2	40:65:91:PRO:O	2.34	0.55
1:13:648:A:H2'	1:13:649:G:C8	2.40	0.55
8:7E:29:SER:HB3	8:7E:32:LYS:HG3	1.87	0.55
46:G8:97:ARG:O	46:G8:107:ASP:HB3	2.07	0.55
1:1G:1129:C:H4'	1:1G:1130:A:C8	2.42	0.55
1:1G:1397:C:H42	24:4L:22:A:H8	1.54	0.55
2:12:162:ILE:HD11	2:12:184:VAL:HG13	1.89	0.55
3:22:161:GLU:O	3:22:162:GLN:HB3	2.07	0.55
5:42:42:GLY:HA2	5:42:65:ASN:O	2.07	0.55
9:82:77:ILE:O	9:82:81:ILE:N	2.35	0.55
10:1A:33:GLN:H	10:1A:75:ILE:HG12	1.71	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:3A:108:LYS:H	12:3A:108:LYS:HD2	1.70	0.55
17:8A:12:SER:HB3	17:8A:20:THR:HB	1.88	0.55
25:14:458:G:O2'	25:14:469:G:O6	2.17	0.55
1:13:93:U:H2'	1:13:95:G:O4'	2.07	0.55
1:13:939:G:H1	1:13:1344:C:H42	1.54	0.55
1:13:1304:G:N1	1:13:1332:A:OP2	2.33	0.55
3:2E:71:ALA:HB2	3:2E:115:LEU:HD13	1.87	0.55
25:1H:177:G:H3'	25:1H:178:G:H8	1.72	0.55
25:1H:605:C:O2	25:1H:657:U:O2'	2.24	0.55
25:1H:888:C:O2'	25:1H:889:C:H4'	2.07	0.55
32:51:121:ILE:HG23	32:51:133:VAL:HG13	1.88	0.55
38:88:112:GLU:H	38:88:112:GLU:CD	2.15	0.55
44:E8:65:LEU:HD13	44:E8:68:ARG:HD3	1.88	0.55
1:1G:1274:G:H21	1:1G:1275:A:H62	1.53	0.55
19:AA:72:GLY:HA2	19:AA:75:ALA:HB3	1.89	0.55
25:14:404:C:O2'	25:14:405:U:OP2	2.23	0.55
25:14:531:C:OP1	25:14:561:G:N1	2.40	0.55
25:14:1035:U:H3	25:14:1120:G:H1	1.54	0.55
25:14:1430:C:H2'	25:14:1431:U:H6	1.72	0.55
30:39:102:PRO:HB2	30:39:105:VAL:HG23	1.88	0.55
1:13:413:G:H2'	1:13:428:G:N2	2.21	0.55
1:13:433:C:H2'	1:13:434:U:H6	1.71	0.55
1:13:1340:A:HO2'	23:2K:32:G:HO2'	1.44	0.55
3:2E:118:GLN:HG2	3:2E:187:ALA:HB2	1.89	0.55
25:1H:845:G:H4'	25:1H:846:C:OP1	2.06	0.55
28:11:108:PRO:HD2	28:11:111:LEU:HG	1.87	0.55
31:41:66:GLN:HA	52:M8:6:HIS:CE1	2.42	0.55
41:B8:105:LEU:O	41:B8:107:ASP:N	2.40	0.55
49:J8:87:PRO:HA	49:J8:90:ILE:HB	1.89	0.55
1:1G:426:G:OP1	4:32:36:ARG:NH1	2.40	0.55
1:1G:690:G:OP2	11:2A:27:ASN:ND2	2.34	0.55
1:1G:1274:G:N2	1:1G:1275:A:H62	2.05	0.55
6:52:36:ARG:NH2	6:52:38:GLU:OE2	2.37	0.55
19:AA:20:LEU:HA	19:AA:44:MET:HE1	1.88	0.55
25:14:2191:G:O2'	25:14:2192:G:OP1	2.25	0.55
25:14:2494:G:H2'	25:14:2495:G:H8	1.72	0.55
26:1J:15:A:H5'	26:1J:16:G:H8	1.71	0.55
29:29:89:ASP:OD1	29:29:89:ASP:N	2.36	0.55
32:59:121:ILE:HD11	32:59:140:LYS:HD2	1.89	0.55
35:15:28:THR:HG22	35:15:29:LYS:HG2	1.88	0.55
37:35:79:ARG:NE	37:35:109:GLY:HA3	2.22	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:65:102:ALA:HB1	40:65:112:PHE:HB3	1.89	0.55
1:13:639:G:H2'	1:13:640:A:C8	2.38	0.54
1:13:1313:U:OP1	19:AI:5:LEU:HB2	2.07	0.54
9:8E:5:TYR:HA	9:8E:17:VAL:O	2.07	0.54
25:1H:1423:G:H2'	25:1H:1424:G:C8	2.42	0.54
25:1H:2781:A:H5''	25:1H:2782:G:H5'	1.87	0.54
36:68:102:VAL:HG23	36:68:121:VAL:HG13	1.89	0.54
44:E8:38:TYR:CE2	53:N8:41:PRO:HD3	2.42	0.54
49:J8:81:LYS:HA	49:J8:81:LYS:NZ	2.22	0.54
1:1G:1175:G:H2'	1:1G:1176:A:C8	2.42	0.54
25:14:883:G:H1	25:14:893:C:H42	1.54	0.54
26:1J:14:U:OP2	26:1J:70:C:O2'	2.25	0.54
31:49:47:LYS:HB2	31:49:82:LEU:HD11	1.89	0.54
33:69:20:ASP:OD1	33:69:20:ASP:N	2.40	0.54
37:35:97:PRO:O	37:35:98:GLU:HG3	2.07	0.54
38:45:81:VAL:HG12	38:45:82:ARG:CZ	2.36	0.54
1:13:1273:G:H3'	1:13:1274:G:H8	1.70	0.54
1:13:1305:G:O2'	1:13:1306:A:N7	2.37	0.54
25:1H:363(F):A:N1	62:1H:3779:HOH:O	2.33	0.54
25:1H:1108:U:OP1	34:38:55:LYS:NZ	2.40	0.54
25:1H:1212:G:O2'	25:1H:1236:G:N2	2.39	0.54
27:71:43:VAL:HG22	27:71:214:VAL:HG22	1.89	0.54
32:51:88:LEU:HA	32:51:130:ARG:HA	1.87	0.54
1:1G:933:G:OP2	7:62:2:ALA:N	2.40	0.54
2:12:15:VAL:HG22	2:12:210:SER:HA	1.88	0.54
7:62:116:ALA:HA	7:62:119:ARG:HG3	1.88	0.54
8:72:44:PHE:HD1	8:72:80:ILE:HG12	1.72	0.54
8:72:64:LYS:HG2	8:72:79:VAL:HG21	1.90	0.54
9:82:65:VAL:HG21	9:82:73:GLN:HB3	1.89	0.54
12:3A:46:ASN:ND2	12:3A:89:OTD:SB	2.79	0.54
25:14:1421:G:H1	25:14:1577:C:H42	1.54	0.54
25:14:2116:G:O6	25:14:2172:U:N3	2.40	0.54
25:14:2567:G:H2'	25:14:2568:C:C6	2.42	0.54
28:19:71:ASP:OD1	28:19:103:ARG:NH2	2.39	0.54
49:F5:40:ARG:NH2	49:F5:42:GLN:HE21	2.04	0.54
1:13:235:C:H5'	17:8I:70:ARG:HG2	1.88	0.54
1:13:380:G:O2'	1:13:382:A:N7	2.36	0.54
1:13:437:U:H2'	1:13:438:G:O4'	2.07	0.54
2:1E:21:ARG:HD2	2:1E:38:GLY:HA3	1.89	0.54
25:1H:1999:C:OP1	25:1H:2723:C:O2'	2.25	0.54
25:1H:2140:C:H1'	25:1H:2152:G:N2	2.22	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:11:130:ALA:HA	28:11:192:THR:HA	1.88	0.54
32:51:103:LEU:HD13	32:51:131:VAL:HG21	1.88	0.54
41:B8:66:VAL:HA	41:B8:71:GLY:HA2	1.88	0.54
1:1G:812:C:H1'	1:1G:813:U:OP2	2.08	0.54
20:BA:54:LYS:HA	20:BA:57:ARG:HH21	1.72	0.54
25:14:34:C:HO2'	25:14:35:G:P	2.30	0.54
25:14:265:A:O2'	25:14:266:G:H4'	2.07	0.54
25:14:675:A:N3	25:14:2443:C:O2'	2.40	0.54
25:14:1074:G:N2	25:14:1075:C:O2	2.40	0.54
25:14:1757:U:H3	25:14:1762:A:H2	1.54	0.54
31:49:96:ARG:O	31:49:98:ARG:N	2.40	0.54
37:35:62:LEU:O	56:M5:13:ARG:NH1	2.40	0.54
45:B5:8:ILE:O	50:G5:36:ARG:NH2	2.41	0.54
55:L5:12:ARG:HB3	55:L5:46:VAL:HG21	1.87	0.54
1:13:739:C:O2'	15:6I:42:HIS:ND1	2.38	0.54
1:13:1274:G:N2	1:13:1275:A:H62	2.06	0.54
2:1E:80:ILE:HD11	2:1E:208:ILE:HG23	1.88	0.54
4:3E:60:GLU:OE2	4:3E:199:ASN:N	2.39	0.54
25:1H:1177:A:H4'	25:1H:1178:C:O5'	2.07	0.54
31:41:77:ILE:H	31:41:82:LEU:HB3	1.73	0.54
41:B8:57:PHE:HA	41:B8:79:HIS:CE1	2.43	0.54
50:K8:65:ASN:CG	50:K8:69:ARG:HH22	2.15	0.54
53:N8:39:MET:C	53:N8:40:LYS:HD2	2.33	0.54
1:1G:181:G:O2'	1:1G:182:U:O5'	2.26	0.54
1:1G:355:C:H1'	1:1G:388:G:H1'	1.88	0.54
1:1G:636:U:H2'	1:1G:637:G:H8	1.72	0.54
1:1G:662:G:H2'	1:1G:663:A:C8	2.42	0.54
1:1G:838:G:N2	1:1G:849:C:O2	2.40	0.54
1:1G:1086:U:H3'	1:1G:1087:G:H8	1.72	0.54
2:12:21:ARG:N	2:12:39:ILE:HG12	2.23	0.54
17:8A:74:LEU:HD12	17:8A:75:ARG:HG3	1.89	0.54
57:2L:76:C:N3	25:14:2251:OMG:N2	2.50	0.54
25:14:993:G:OP1	42:85:50:ARG:NH2	2.41	0.54
25:14:1454:U:OP1	39:55:77:ARG:NH1	2.40	0.54
25:14:1791:A:N6	25:14:1828:G:O2'	2.40	0.54
31:49:47:LYS:HE2	31:49:81:LYS:HG2	1.90	0.54
47:D5:19:ARG:NH1	47:D5:84:GLU:O	2.40	0.54
1:13:1062:U:H2'	1:13:1063:C:C6	2.43	0.54
1:13:1137:C:O2'	1:13:1138:G:N3	2.40	0.54
16:7I:21:VAL:O	16:7I:33:ILE:N	2.41	0.54
25:1H:603:A:N1	25:1H:625:G:O2'	2.41	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:16:95:U:H2'	26:16:96:G:C8	2.43	0.54
31:41:112:PRO:HB3	52:M8:37:SER:N	2.22	0.54
36:68:78:ARG:HG2	41:B8:73:GLU:HG3	1.89	0.54
41:B8:50:ILE:HG13	41:B8:64:ARG:HB2	1.88	0.54
1:1G:601:C:H42	1:1G:637:G:H1	1.56	0.54
3:22:47:LEU:HD21	3:22:68:VAL:HG11	1.89	0.54
18:9A:50:ILE:HD11	18:9A:70:ILE:HG21	1.90	0.54
57:2L:33:OMC:HM22	57:2L:34:U:H5'	1.90	0.54
22:3L:42:C:H2'	22:3L:43:G:C8	2.43	0.54
22:3L:60:A:H2'	22:3L:61:U:H5'	1.90	0.54
25:14:270(U):C:H2'	25:14:270(V):G:C8	2.42	0.54
25:14:1313:U:H4'	25:14:1332:G:H4'	1.88	0.54
25:14:2439:A:H5'	25:14:2439:A:C8	2.41	0.54
31:49:114:ILE:HG22	31:49:117:PHE:HB2	1.90	0.54
1:13:346:G:H5''	41:B8:41:ARG:NH1	2.23	0.54
1:13:1206:G:OP1	3:2E:190:ARG:NH1	2.39	0.54
12:3I:72:HIS:HA	12:3I:99:ARG:HH22	1.73	0.54
25:1H:953:A:OP2	38:88:16:ARG:HD3	2.08	0.54
25:1H:1637:A:OP2	62:1H:3683:HOH:O	2.19	0.54
25:1H:1818:U:H2'	28:11:157:ARG:HG3	1.88	0.54
37:78:37:GLY:N	62:78:301:HOH:O	2.40	0.54
37:78:56:SER:O	37:78:57:THR:OG1	2.18	0.54
51:L8:30:ARG:HE	51:L8:33:GLN:HG3	1.72	0.54
1:1G:99:C:H2'	1:1G:101:A:C8	2.42	0.54
1:1G:303:A:H2'	1:1G:304:U:O4'	2.08	0.54
1:1G:1256:A:H5'	1:1G:1257:U:OP1	2.07	0.54
2:12:15:VAL:HG23	2:12:209:ARG:HD2	1.89	0.54
11:2A:32:ILE:HD13	11:2A:72:ALA:HB2	1.90	0.54
25:14:34:C:O2'	25:14:35:G:O5'	2.24	0.54
25:14:296:C:OP1	46:C5:4:LYS:NZ	2.37	0.54
25:14:1860:G:H1	25:14:1882:C:H42	1.54	0.54
25:14:2299:G:C2	25:14:2318:G:H8	2.26	0.54
62:14:3519:HOH:O	28:19:244:ARG:NH2	2.41	0.54
29:29:120:TRP:CD1	29:29:155:LYS:HB3	2.43	0.54
32:59:93:GLY:O	32:59:95:ARG:NH2	2.41	0.54
1:13:812:C:O2'	1:13:813:U:O5'	2.24	0.54
1:13:1060:C:OP1	14:5I:45:ARG:NH2	2.38	0.54
1:13:1157:A:H1'	1:13:1158:C:N3	2.23	0.54
1:13:1233:G:O2'	1:13:1365:G:OP1	2.26	0.54
1:13:1451:A:H5''	1:13:1452:C:O5'	2.07	0.54
18:9I:51:LEU:HD22	18:9I:55:ARG:HD2	1.90	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:AI:22:LEU:HA	19:AI:25:LYS:O	2.07	0.54
32:51:86:GLU:OE2	32:51:165:ALA:N	2.39	0.54
4:32:22:LYS:HB2	60:32:303:SF4:S4	2.47	0.54
22:3L:64:G:H2'	22:3L:65:G:C8	2.42	0.54
25:14:16:G:H2'	25:14:17:G:H8	1.72	0.54
25:14:814:C:OP1	43:95:83:ARG:N	2.38	0.54
25:14:1191:G:OP1	37:35:18:ARG:NH1	2.40	0.54
26:1J:5:C:O2'	26:1J:27:C:O2	2.26	0.54
26:1J:28:C:H2'	26:1J:29:A:C8	2.42	0.54
30:39:20:LEU:HD22	30:39:199:TRP:CZ3	2.43	0.54
32:59:16:SER:OG	32:59:26:VAL:O	2.24	0.54
1:13:801:U:H2'	1:13:802:A:C8	2.43	0.54
10:1I:50:ILE:HB	14:5I:41:ARG:NE	2.22	0.54
25:1H:213:A:OP2	62:1H:3677:HOH:O	2.18	0.54
25:1H:1053:C:N4	25:1H:1106:G:H1	1.93	0.54
41:B8:107:ASP:N	41:B8:107:ASP:OD1	2.38	0.54
1:1G:304:U:H2'	1:1G:305:G:C8	2.43	0.54
1:1G:445:G:H2'	1:1G:446:G:C8	2.43	0.54
1:1G:1306:A:N6	1:1G:1331:G:O2'	2.40	0.54
25:14:630:G:N2	25:14:633:A:OP2	2.34	0.54
25:14:2305:A:H5''	31:49:134:GLY:HA3	1.89	0.54
25:14:2334:G:O6	48:E5:74:ARG:NH2	2.40	0.54
25:14:2564:A:C2	25:14:2647:U:H4'	2.42	0.54
32:59:2:SER:OG	32:59:3:ARG:N	2.40	0.54
37:35:5:ASP:HA	37:35:7:ARG:NH2	2.23	0.54
40:65:88:ASP:OD1	40:65:90:GLY:N	2.41	0.54
1:13:375:U:H5''	16:7I:69:THR:HG21	1.90	0.54
6:5E:41:GLU:HB3	6:5E:62:TRP:HB3	1.90	0.54
10:1I:51:ARG:NH2	10:1I:61:GLU:OE1	2.41	0.54
25:1H:1858:G:H2'	25:1H:1883:G:H22	1.72	0.54
25:1H:2154:G:H2'	25:1H:2155:G:C8	2.40	0.54
46:G8:38:ILE:HD11	46:G8:64:GLU:HG3	1.89	0.54
1:1G:1104:G:H4'	2:12:111:ARG:NE	2.23	0.54
2:12:29:ALA:C	2:12:31:TYR:N	2.63	0.54
4:32:152:SER:HA	4:32:155:LEU:HD23	1.90	0.54
8:72:89:PRO:HA	8:72:92:ARG:HH11	1.73	0.54
25:14:1007:C:OP1	35:15:35:ARG:NH1	2.41	0.54
25:14:2502:G:H5''	25:14:2503:2MA:H5''	1.89	0.54
25:14:2784:C:H2'	25:14:2785:C:C6	2.43	0.54
32:59:154:PRO:HB3	32:59:163:TYR:CE2	2.43	0.54
48:E5:25:ARG:HH21	48:E5:35:ASN:HB3	1.73	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
55:L5:8:ASN:HB3	55:L5:11:LYS:HB3	1.88	0.54
1:13:401:C:H2'	1:13:402:G:C8	2.43	0.54
1:13:1289:A:H3'	1:13:1290:G:H8	1.73	0.54
13:4I:17:VAL:O	13:4I:20:THR:OG1	2.25	0.54
25:1H:557:U:H2'	25:1H:558:G:C8	2.43	0.54
27:71:46:LYS:HE3	27:71:210:ARG:HB3	1.90	0.54
27:71:201:PRO:HD2	27:71:208:PHE:CE2	2.43	0.54
32:51:41:MET:HE1	32:51:64:LEU:HB2	1.90	0.54
1:1G:36:C:N4	62:1G:1848:HOH:O	2.41	0.54
1:1G:951:G:OP2	13:4A:102:ARG:NH2	2.41	0.54
1:1G:1137:C:H4'	1:1G:1138:G:O5'	2.07	0.54
6:52:25:ILE:HG21	6:52:82:ARG:HD2	1.90	0.54
19:AA:38:SER:O	19:AA:70:LYS:HB3	2.08	0.54
25:14:19:C:H2'	25:14:20:C:H6	1.73	0.54
25:14:1043:C:H42	25:14:1112:G:H22	1.56	0.54
25:14:1802:A:H2'	25:14:1803:A:C8	2.42	0.54
25:14:1991:U:H2'	25:14:1992:G:H5''	1.89	0.54
25:14:2304:G:N2	25:14:2312:U:O4	2.41	0.54
25:14:2710:C:OP1	62:14:3549:HOH:O	2.19	0.54
30:39:78:ILE:HA	30:39:83:PHE:CD2	2.43	0.54
35:15:19:GLU:HG3	35:15:59:LYS:HB3	1.91	0.54
41:75:80:SER:HB2	41:75:82:LEU:HD12	1.89	0.54
47:D5:70:LEU:O	47:D5:89:PHE:N	2.26	0.54
53:J5:41:PRO:HD2	53:J5:44:THR:HG22	1.90	0.54
1:13:572:A:OP1	62:13:1808:HOH:O	2.18	0.53
1:13:1291:G:OP1	7:6E:37:ASN:ND2	2.41	0.53
7:6E:15:ASP:HB3	7:6E:19:GLY:H	1.74	0.53
9:8E:4:TYR:CZ	9:8E:88:TYR:HB2	2.43	0.53
9:8E:61:ALA:HB1	9:8E:63:ILE:HD11	1.90	0.53
10:1I:51:ARG:HB2	10:1I:60:ARG:HA	1.90	0.53
25:1H:612:G:O2'	25:1H:616:A:N1	2.35	0.53
25:1H:2306:C:H3'	25:1H:2307:G:H5''	1.90	0.53
25:1H:2607:G:O3'	62:1H:3680:HOH:O	2.18	0.53
28:11:182:LEU:HB2	28:11:271:ILE:HG13	1.91	0.53
38:88:36:ALA:HA	38:88:129:THR:HG22	1.89	0.53
1:1G:547:A:OP1	4:32:73:ARG:NH2	2.40	0.53
1:1G:1122:U:O4	1:1G:1123:A:N6	2.42	0.53
11:2A:20:TYR:HB2	11:2A:31:THR:HG23	1.89	0.53
25:14:403:U:H4'	25:14:404:C:H5'	1.90	0.53
25:14:559:G:O2'	42:85:52:ARG:NH1	2.37	0.53
31:49:60:LEU:O	31:49:64:THR:HG22	2.08	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:35:106:LEU:HD13	37:35:112:LEU:HG	1.90	0.53
38:45:125:LEU:HD13	38:45:126:PRO:HD2	1.91	0.53
1:13:4:U:C5	8:7E:102:ARG:HG3	2.43	0.53
1:13:949:A:H61	1:13:1232:U:H3	1.56	0.53
3:2E:71:ALA:HA	3:2E:106:VAL:HB	1.90	0.53
25:1H:222:A:H3'	25:1H:421:U:H5'	1.90	0.53
25:1H:1109:C:O2'	25:1H:1110:G:OP1	2.27	0.53
25:1H:2099:U:H2'	25:1H:2100:G:H8	1.73	0.53
25:1H:2701:C:C3'	25:1H:2702:U:H5''	2.33	0.53
52:M8:48:ARG:CZ	52:M8:51:ASP:HA	2.38	0.53
1:1G:413:G:O2'	1:1G:414:A:OP2	2.26	0.53
1:1G:843:U:H3'	1:1G:848:C:O4'	2.08	0.53
19:AA:36:ARG:HD2	19:AA:72:GLY:H	1.73	0.53
25:14:782:A:N7	28:19:221:VAL:HG21	2.23	0.53
25:14:992:C:H2'	25:14:993:G:H8	1.72	0.53
25:14:1076:C:H2'	25:14:1077:A:H5'	1.89	0.53
40:65:67:ARG:O	40:65:71:ARG:HG3	2.08	0.53
50:G5:52:ASP:O	50:G5:56:GLN:N	2.36	0.53
1:13:422:C:HO2'	1:13:423:G:N2	2.06	0.53
1:13:1151:A:OP1	10:1I:41:PRO:HA	2.09	0.53
5:4E:41:VAL:HG22	5:4E:69:VAL:HG21	1.89	0.53
25:1H:184:C:H2'	25:1H:185:U:C6	2.43	0.53
25:1H:451:C:H4'	30:31:52:LYS:NZ	2.24	0.53
25:1H:2168:G:N2	25:1H:2170:A:OP2	2.41	0.53
25:1H:2310:A:H8	25:1H:2311:A:H5''	1.73	0.53
34:38:72:ASP:HB3	34:38:118:THR:O	2.08	0.53
47:H8:18:LEU:HD22	47:H8:25:PRO:HG3	1.90	0.53
3:22:165:THR:O	3:22:165:THR:OG1	2.26	0.53
6:52:35:ALA:HA	6:52:67:MET:HB3	1.89	0.53
20:BA:63:ILE:HG21	20:BA:81:LYS:HB2	1.89	0.53
25:14:5:A:H2'	25:14:6:A:H8	1.72	0.53
25:14:270(E):G:H2'	25:14:270(F):U:O4'	2.09	0.53
25:14:1343:G:H1	25:14:1404:C:H42	1.56	0.53
42:85:88:ILE:HD12	42:85:90:VAL:HB	1.90	0.53
1:13:1410:G:H2'	1:13:1411:C:C6	2.44	0.53
11:2I:22:HIS:HB3	11:2I:29:ILE:HG23	1.91	0.53
22:3K:7:G:H5''	22:3K:8:U:OP2	2.08	0.53
25:1H:1156:A:OP2	62:1H:3678:HOH:O	2.18	0.53
25:1H:1470:G:O2'	25:1H:1522:G:O6	2.19	0.53
25:1H:2172:U:H5'	25:1H:2173:A:OP2	2.09	0.53
31:41:125:PHE:HB3	31:41:166:ASP:HB3	1.90	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:68:80:ASP:OD2	41:B8:64:ARG:NH2	2.42	0.53
41:B8:108:ARG:HA	41:B8:111:ARG:NE	2.23	0.53
47:H8:23:LYS:HD3	47:H8:40:ASP:HA	1.90	0.53
47:H8:154:ASP:HB3	47:H8:155:LEU:HD22	1.89	0.53
52:M8:11:PRO:HA	52:M8:25:TYR:HA	1.90	0.53
54:O8:25:LYS:HE3	56:Q8:34:TRP:HE1	1.74	0.53
1:1G:1074:G:H4'	2:12:104:ASN:HB2	1.91	0.53
16:7A:21:VAL:HG23	16:7A:33:ILE:HB	1.89	0.53
26:1J:94:C:H2'	26:1J:95:U:O4'	2.08	0.53
40:65:63:THR:OG1	40:65:64:GLU:OE1	2.27	0.53
45:B5:8:ILE:H	45:B5:8:ILE:HD12	1.72	0.53
46:C5:106:LEU:C	46:C5:108:THR:H	2.16	0.53
47:D5:4:ARG:HD3	47:D5:4:ARG:H	1.73	0.53
51:H5:39:ASP:O	51:H5:44:ARG:NH2	2.41	0.53
54:K5:14:THR:HG22	54:K5:50:ARG:O	2.08	0.53
1:13:222:U:H2'	1:13:223:U:C6	2.44	0.53
1:13:762:C:H2'	1:13:763:G:C8	2.43	0.53
1:13:1108:G:H5'	3:2E:176:HIS:ND1	2.22	0.53
1:13:1219:U:OP1	14:5I:19:ARG:NH2	2.39	0.53
9:8E:9:ARG:HG3	9:8E:14:VAL:HG22	1.91	0.53
25:1H:127:A:H5''	25:1H:128:C:C6	2.43	0.53
25:1H:661:C:O2'	37:78:13:ASN:HA	2.09	0.53
25:1H:784:A:C5	28:11:229:VAL:HG21	2.44	0.53
25:1H:873:G:O3'	38:88:63:LYS:NZ	2.37	0.53
25:1H:945:A:N7	62:1H:3826:HOH:O	2.41	0.53
25:1H:1657:C:H2'	25:1H:1658:C:C6	2.44	0.53
30:31:183:VAL:O	30:31:187:VAL:HG23	2.08	0.53
41:B8:37:GLY:O	41:B8:38:ASN:ND2	2.26	0.53
41:B8:55:ASN:H	41:B8:59:THR:CB	2.19	0.53
1:1G:543:C:OP1	4:32:14:ARG:HG3	2.08	0.53
1:1G:947:G:O2'	1:1G:1306:A:O2'	2.15	0.53
2:12:77:ALA:HB1	2:12:165:VAL:HG11	1.91	0.53
3:22:54:ARG:HB2	3:22:69:HIS:HB2	1.90	0.53
6:52:76:ALA:HA	6:52:79:LEU:HB2	1.91	0.53
25:14:270(U):C:H2'	25:14:270(V):G:H8	1.73	0.53
25:14:363(E):U:H5'	25:14:363(F):A:OP2	2.09	0.53
25:14:1383:C:O2	62:14:3547:HOH:O	2.18	0.53
28:19:239:ARG:NE	62:19:404:HOH:O	2.31	0.53
47:D5:145:GLU:HA	47:D5:174:VAL:HG21	1.90	0.53
1:13:483:C:OP2	1:13:484:G:O2'	2.26	0.53
1:13:1302:U:C6	13:4I:17:VAL:HG21	2.43	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1314:C:H2'	1:13:1315:U:C6	2.44	0.53
1:13:1346:A:N1	1:13:1374:A:H5''	2.23	0.53
7:6E:15:ASP:HB3	7:6E:19:GLY:N	2.23	0.53
9:8E:48:GLU:OE1	9:8E:48:GLU:N	2.42	0.53
22:1K:31:G:H1	22:1K:41:C:N4	2.06	0.53
25:1H:1252:G:H1'	42:C8:33:ARG:HD3	1.91	0.53
25:1H:2068:U:N3	25:1H:2430:A:H2	2.00	0.53
49:J8:76:ARG:H	49:J8:76:ARG:HD2	1.73	0.53
1:1G:123:C:O2'	1:1G:290:C:O2	2.27	0.53
1:1G:137:C:H1'	16:7A:63:GLY:HA3	1.90	0.53
3:22:5:ILE:HG12	3:22:6:HIS:N	2.24	0.53
6:52:37:VAL:HA	6:52:65:VAL:HG12	1.90	0.53
9:82:4:TYR:HB3	9:82:87:GLN:HE22	1.73	0.53
19:AA:5:LEU:HD13	19:AA:9:VAL:HA	1.91	0.53
19:AA:44:MET:HB3	19:AA:47:HIS:CD2	2.44	0.53
25:14:920:G:H2'	25:14:921:G:C8	2.44	0.53
25:14:2019:A:H4'	42:85:34:LYS:HD2	1.90	0.53
26:1J:55:U:H4'	31:49:29:TRP:HE1	1.73	0.53
40:65:20:ARG:HE	40:65:21:THR:HG22	1.73	0.53
42:85:88:ILE:HA	43:95:49:THR:O	2.08	0.53
43:95:62:LEU:HD22	43:95:95:LEU:HB2	1.89	0.53
46:C5:91:GLU:C	46:C5:92:ASN:HD22	2.15	0.53
48:E5:18:ALA:HB3	48:E5:20:ARG:HH21	1.73	0.53
1:13:983:A:H5'	14:5I:3:ARG:NH1	2.23	0.53
1:13:1336:C:H1'	1:13:1337:G:C2	2.44	0.53
18:9I:70:ILE:O	18:9I:74:ARG:HG3	2.08	0.53
25:1H:640:C:H2'	25:1H:641:C:C6	2.44	0.53
25:1H:666:G:H4'	37:78:49:ARG:HH12	1.73	0.53
25:1H:997:G:OP1	42:C8:93:LYS:HB2	2.08	0.53
25:1H:1639:U:H2'	25:1H:1640:C:H5''	1.90	0.53
28:11:182:LEU:N	28:11:272:ALA:HB3	2.18	0.53
1:1G:928:G:O2'	1:1G:1533:C:O2'	2.23	0.53
1:1G:973:G:H3'	1:1G:974:A:H5''	1.91	0.53
2:12:67:THR:HG23	2:12:159:PRO:HA	1.91	0.53
4:32:45:GLN:HB2	4:32:46:LYS:HE2	1.90	0.53
7:62:68:ASN:ND2	7:62:128:ALA:O	2.42	0.53
25:14:1366:A:OP1	49:F5:2:SER:OG	2.25	0.53
25:14:1525:G:H2'	25:14:1526:G:H8	1.74	0.53
25:14:2837:G:H1	25:14:2881:C:H42	1.55	0.53
26:1J:113:C:H4'	40:65:46:VAL:HG22	1.90	0.53
31:49:18:GLU:HB3	31:49:175:LEU:HD13	1.91	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1252:A:H2'	1:13:1253:G:O4'	2.08	0.53
25:1H:64:A:H1'	45:F8:66:LEU:HB3	1.89	0.53
25:1H:363:G:H2'	25:1H:363(A):A:C8	2.44	0.53
25:1H:780:G:H21	25:1H:783:A:H62	1.55	0.53
25:1H:2638:G:O2'	25:1H:2639:A:H8	1.90	0.53
25:1H:2652:C:H42	25:1H:2668:G:H1	1.57	0.53
25:1H:2807:G:H3'	25:1H:2808:U:H5''	1.91	0.53
32:51:20:ALA:HB3	32:51:23:ARG:HG3	1.91	0.53
33:61:4:ILE:HG12	33:61:18:VAL:HG13	1.91	0.53
41:B8:2:ASN:HB3	41:B8:5:ALA:HB3	1.90	0.53
1:1G:824:C:H2'	1:1G:825:G:H8	1.73	0.53
1:1G:1001:G:H2'	1:1G:1002:G:C8	2.44	0.53
1:1G:1499:A:H1'	1:1G:1520:G:H5'	1.90	0.53
25:14:1058:U:H2'	25:14:1059:G:C8	2.43	0.53
25:14:1324:G:O2'	25:14:1326:U:OP2	2.23	0.53
25:14:1567:A:H5'	28:19:86:PRO:HB3	1.89	0.53
25:14:2749:A:H4'	32:59:62:LYS:HB3	1.91	0.53
32:59:86:GLU:H	32:59:86:GLU:CD	2.17	0.53
47:D5:124:ILE:HD11	47:D5:165:VAL:HG11	1.91	0.53
52:I5:40:HIS:H	52:I5:41:PRO:HD3	1.73	0.53
55:L5:13:ALA:HB2	55:L5:46:VAL:HG11	1.90	0.53
1:13:1414:U:H2'	1:13:1415:G:H8	1.74	0.53
1:13:1427:U:H2'	1:13:1428:A:C8	2.43	0.53
2:1E:115:LEU:HA	2:1E:118:LEU:HD12	1.90	0.53
3:2E:95:THR:HG22	3:2E:96:GLY:H	1.74	0.53
25:1H:287:C:H42	25:1H:354:G:H1	1.55	0.53
25:1H:818:G:OP2	62:1H:3648:HOH:O	2.17	0.53
25:1H:1204:A:H1'	25:1H:1206:G:C8	2.44	0.53
25:1H:1971:A:C4	28:11:241:PRO:HD3	2.43	0.53
25:1H:2086:U:H2'	25:1H:2087:G:C8	2.43	0.53
25:1H:2308:G:H1	25:1H:2311:A:H2	1.52	0.53
28:11:70:TRP:O	28:11:73:VAL:HG23	2.09	0.53
47:H8:140:ASP:N	47:H8:155:LEU:HG	2.18	0.53
51:L8:6:VAL:HG12	51:L8:56:VAL:HG22	1.89	0.53
1:1G:1154:G:H2'	1:1G:1155:G:H8	1.72	0.53
6:52:38:GLU:OE1	6:52:64:GLN:NE2	2.41	0.53
25:14:1434:A:H61	25:14:1558:A:H62	1.57	0.53
25:14:1783:A:OP2	62:14:3528:HOH:O	2.18	0.53
25:14:2053:G:H2'	25:14:2054:A:O4'	2.09	0.53
25:14:2730:C:O2'	29:29:168:MET:O	2.27	0.53
52:I5:46:GLN:HG2	52:I5:48:ARG:H	1.74	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:K5:42:TRP:CG	54:K5:43:CYS:N	2.77	0.53
1:13:439:A:OP2	1:13:493:G:N1	2.41	0.53
1:13:540:G:H2'	1:13:541:G:O4'	2.09	0.53
1:13:659:U:H2'	1:13:660:G:C8	2.44	0.53
1:13:757:U:H2'	1:13:758:G:O4'	2.08	0.53
1:13:811:C:O2'	1:13:901:A:N1	2.41	0.53
1:13:831:U:H2'	1:13:832:C:C6	2.42	0.53
19:AI:27:GLU:OE2	19:AI:29:ARG:NH2	2.42	0.53
20:BI:90:GLN:HA	20:BI:93:GLU:HB2	1.91	0.53
25:1H:1203:G:O2'	25:1H:1242:A:N6	2.40	0.53
25:1H:1557:C:OP2	25:1H:1558:A:O2'	2.22	0.53
28:11:108:PRO:HG3	28:11:143:HIS:CE1	2.44	0.53
31:41:83:ARG:HB2	31:41:86:MET:HG3	1.90	0.53
35:58:14:VAL:HG12	35:58:15:LEU:H	1.74	0.53
1:1G:559:A:H4'	1:1G:560:U:H5''	1.91	0.53
1:1G:954:G:H21	1:1G:1227:A:N6	2.07	0.53
12:3A:67:ILE:HG12	12:3A:97:ILE:HD12	1.91	0.53
20:BA:34:LYS:O	20:BA:38:LYS:N	2.40	0.53
25:14:38:A:H2'	25:14:39:C:C6	2.44	0.53
25:14:1055:G:N2	25:14:1085:A:N3	2.52	0.53
25:14:2197:U:H1'	25:14:2198:A:C8	2.44	0.53
25:14:2294:C:OP1	40:65:89:ARG:NH2	2.37	0.53
28:19:146:GLU:HB2	28:19:189:CYS:HB3	1.90	0.53
38:45:126:PRO:HG2	38:45:127:ILE:HD12	1.91	0.53
55:L5:5:TRP:NE1	55:L5:7:PRO:HG3	2.24	0.53
1:13:316:G:OP2	1:13:351:G:O2'	2.26	0.52
1:13:1321:C:H3'	1:13:1322:C:H5''	1.91	0.52
3:2E:95:THR:HG21	3:2E:97:LYS:HZ3	1.74	0.52
34:38:141:VAL:HG12	34:38:142:LEU:HD23	1.90	0.52
1:1G:1502:A:H4'	1:1G:1503:A:OP2	2.09	0.52
6:52:6:VAL:HG13	6:52:90:VAL:HG22	1.90	0.52
8:72:109:ILE:HG23	8:72:137:VAL:HG23	1.90	0.52
22:3L:32:G:H1	22:3L:40:C:H42	1.57	0.52
25:14:413:C:H42	25:14:2410:G:H1	1.57	0.52
25:14:602:G:O2'	25:14:655:A:N6	2.43	0.52
25:14:928:G:H5'	25:14:929:G:OP2	2.09	0.52
25:14:1062:G:N2	25:14:1076:C:H42	2.04	0.52
25:14:2046:G:H5'	53:J5:19:ARG:HG3	1.90	0.52
25:14:2391:G:H1'	25:14:2424:C:H41	1.74	0.52
41:75:108:ARG:O	41:75:111:ARG:HG2	2.09	0.52
1:13:294:U:OP1	1:13:610:G:O2'	2.22	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:503:C:OP2	12:3I:113:SER:HB3	2.08	0.52
1:13:1106:G:H5'	3:2E:172:ARG:HG2	1.90	0.52
1:13:1241:G:H2'	1:13:1242:C:C6	2.44	0.52
1:13:1302:U:H6	13:4I:17:VAL:HG21	1.74	0.52
7:6E:75:VAL:HG13	7:6E:145:ALA:HA	1.90	0.52
9:8E:28:VAL:HG22	9:8E:63:ILE:HD13	1.91	0.52
11:2I:23:ALA:HA	11:2I:28:THR:HG23	1.91	0.52
18:9I:33:ASP:OD2	18:9I:36:ASN:ND2	2.42	0.52
19:AI:48:THR:O	19:AI:48:THR:OG1	2.26	0.52
25:1H:1175:U:H4'	25:1H:1176:G:OP1	2.08	0.52
25:1H:1314:C:OP1	62:1H:3617:HOH:O	2.19	0.52
25:1H:2630:G:H1	25:1H:2788:C:H42	1.55	0.52
27:7I:215:THR:OG1	27:7I:220:PRO:O	2.14	0.52
32:5I:18:GLU:HG3	32:5I:25:LYS:HB2	1.91	0.52
41:B8:76:PHE:HB3	41:B8:83:ILE:HD11	1.90	0.52
10:1A:51:ARG:HB2	10:1A:60:ARG:HA	1.92	0.52
12:3A:44:LYS:HB2	12:3A:45:PRO:HD3	1.90	0.52
25:14:34:C:O2'	25:14:35:G:H8	1.92	0.52
25:14:328:U:H4'	46:C5:68:HIS:CE1	2.44	0.52
25:14:416:C:H42	25:14:2407:G:H1	1.56	0.52
25:14:940:G:H2'	25:14:941:A:O4'	2.09	0.52
25:14:1092:C:H42	25:14:1100:C:N4	2.07	0.52
25:14:2518:A:H4'	25:14:2519:U:OP1	2.09	0.52
25:14:2851:A:O2'	39:55:64:ARG:NH2	2.43	0.52
26:1J:42:C:H1'	31:49:92:VAL:HG23	1.91	0.52
31:49:72:ARG:HB2	31:49:72:ARG:HH11	1.75	0.52
38:45:19:GLY:H	38:45:98:LYS:HZ3	1.58	0.52
41:75:51:ARG:HG2	41:75:98:LYS:HE3	1.92	0.52
1:13:42:G:H1	1:13:400:C:H42	1.57	0.52
1:13:737:A:H2'	1:13:738:C:C6	2.44	0.52
2:1E:153:ARG:O	2:1E:156:LYS:NZ	2.40	0.52
4:3E:162:LEU:HA	4:3E:165:MET:HB3	1.91	0.52
5:4E:63:ARG:HA	5:4E:66:MET:HE2	1.90	0.52
8:7E:20:TYR:HE2	8:7E:75:ARG:HB3	1.75	0.52
8:7E:109:ILE:HG22	8:7E:137:VAL:HB	1.92	0.52
12:3I:8:VAL:HG21	17:8I:34:LYS:HD3	1.91	0.52
25:1H:1026:U:H1'	25:1H:1027:A:O5'	2.09	0.52
27:7I:216:THR:OG1	27:7I:217:THR:N	2.42	0.52
30:31:149:ASP:OD1	30:31:149:ASP:N	2.40	0.52
45:F8:5:TYR:HB3	50:K8:33:MET:HB2	1.89	0.52
1:1G:1028:C:H1'	1:1G:1034:G:H1	1.74	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:2L:44:A:H2'	57:2L:45:A:C8	2.43	0.52
25:14:330:A:H2	25:14:1210:A:HO2'	1.55	0.52
25:14:1300:U:H4'	25:14:1301:A:O5'	2.10	0.52
25:14:2557:G:H2'	25:14:2558:C:C6	2.44	0.52
29:29:53:PRO:HA	29:29:75:VAL:HA	1.90	0.52
40:65:15:ARG:HH21	40:65:90:GLY:HA2	1.75	0.52
1:13:514:C:H2'	1:13:515:G:C8	2.44	0.52
1:13:1367:C:H5'	10:1I:60:ARG:CZ	2.39	0.52
3:2E:112:SER:HB3	3:2E:115:LEU:HG	1.91	0.52
7:6E:155:ARG:O	7:6E:155:ARG:NH2	2.42	0.52
9:8E:26:VAL:HG13	9:8E:61:ALA:HB3	1.91	0.52
11:2I:19:ALA:HA	11:2I:32:ILE:HA	1.92	0.52
25:1H:660:G:O3'	30:31:38:ARG:NH2	2.43	0.52
25:1H:2480:C:N4	25:1H:2481:G:O6	2.43	0.52
25:1H:2820:A:O2'	25:1H:2821:A:OP1	2.26	0.52
27:71:10:LEU:HG	27:71:32:LEU:HA	1.92	0.52
33:61:92:VAL:HG13	33:61:120:ILE:HG23	1.90	0.52
37:78:126:VAL:HG12	37:78:147:LEU:HG	1.90	0.52
40:A8:67:ARG:O	40:A8:71:ARG:HG3	2.08	0.52
48:I8:10:THR:OG1	48:I8:12:ASN:OD1	2.27	0.52
50:K8:17:SER:O	50:K8:20:GLU:HG2	2.09	0.52
53:N8:41:PRO:HB2	53:N8:42:PRO:CD	2.39	0.52
1:1G:1043:C:H2'	1:1G:1044:A:O4'	2.10	0.52
1:1G:1305:G:H22	1:1G:1331:G:H2'	1.73	0.52
1:1G:1316:G:H4'	14:5A:18:VAL:HG11	1.91	0.52
1:1G:1342:C:H1'	9:82:124:GLN:NE2	2.24	0.52
1:1G:1350:A:O2'	7:62:33:ASP:OD1	2.23	0.52
3:22:88:ARG:HA	3:22:91:LEU:HD13	1.90	0.52
9:82:27:THR:HG21	9:82:33:PHE:H	1.74	0.52
13:4A:90:LEU:HA	13:4A:93:ARG:HG2	1.91	0.52
25:14:142:G:H4'	45:B5:35:THR:HG21	1.92	0.52
25:14:262:A:H2'	25:14:263:C:O4'	2.10	0.52
25:14:729:G:H2'	25:14:1775:U:H1'	1.90	0.52
25:14:1255:U:H5''	25:14:1256:G:H5''	1.91	0.52
25:14:1368:G:H2'	25:14:1369:G:H8	1.74	0.52
25:14:1419:A:H61	25:14:1494:A:H61	1.57	0.52
25:14:1442:G:H1	25:14:1549:C:H42	1.56	0.52
25:14:2602:A:H4'	25:14:2603:G:O5'	2.09	0.52
31:49:67:LYS:HE2	52:I5:2:LYS:HG2	1.89	0.52
1:13:1306:A:H61	1:13:1331:G:H1'	1.75	0.52
6:5E:50:TYR:OH	18:9I:74:ARG:O	2.22	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:1K:60:A:H2'	22:1K:61:U:H5'	1.91	0.52
22:3K:69:C:H2'	22:3K:70:C:C6	2.44	0.52
25:1H:140:A:H8	25:1H:1408:C:HO2'	1.58	0.52
25:1H:2636:U:H3	25:1H:2782:G:H1	1.57	0.52
28:11:8:PRO:HB3	28:11:14:ARG:HG2	1.92	0.52
33:61:43:ASN:OD1	33:61:43:ASN:N	2.42	0.52
1:1G:616:G:H2'	1:1G:617:G:C8	2.44	0.52
1:1G:684:A:N6	62:1G:1820:HOH:O	2.42	0.52
1:1G:1158:C:H2'	1:1G:1158:C:O2	2.08	0.52
1:1G:1316:G:N2	1:1G:1318:A:H3'	2.25	0.52
3:22:35:GLU:HA	3:22:38:ARG:HE	1.75	0.52
16:7A:18:ARG:HD3	16:7A:35:LYS:HD2	1.91	0.52
39:55:96:ARG:HG3	39:55:97:VAL:N	2.24	0.52
1:13:1157:A:H62	1:13:1178:G:N2	2.07	0.52
7:6E:50:ILE:HB	7:6E:58:PRO:HB3	1.92	0.52
25:1H:856:C:H42	25:1H:921:G:H1	1.57	0.52
25:1H:1434:A:H61	25:1H:1558:A:H62	1.56	0.52
27:71:213:TYR:CZ	27:71:223:ARG:HG3	2.45	0.52
29:21:70:ALA:O	29:21:72:VAL:N	2.42	0.52
36:68:29:ASN:OD1	36:68:29:ASN:N	2.43	0.52
38:88:30:GLY:CA	38:88:107:ALA:HB2	2.39	0.52
38:88:111:GLU:HG3	38:88:112:GLU:N	2.24	0.52
1:1G:279:A:OP2	17:8A:95:TYR:OH	2.17	0.52
1:1G:872:A:H4'	1:1G:873:A:OP1	2.08	0.52
11:2A:34:ASP:HB3	11:2A:40:ILE:HD11	1.92	0.52
25:14:468:G:N7	55:L5:39:ARG:NH2	2.58	0.52
25:14:589:C:H2'	25:14:590:A:C8	2.44	0.52
25:14:846:C:H4'	25:14:847:U:OP1	2.08	0.52
25:14:2123:G:H22	25:14:2176:A:H2	1.57	0.52
26:1J:75:G:H5''	47:D5:36:LYS:HE2	1.92	0.52
37:35:125:VAL:HG13	37:35:144:GLU:HB3	1.91	0.52
43:95:35:LEU:H	43:95:35:LEU:HD22	1.73	0.52
43:95:40:LEU:HD13	43:95:41:GLY:H	1.74	0.52
2:1E:8:LYS:HG2	2:1E:10:LEU:H	1.74	0.52
20:BI:50:GLU:HG3	20:BI:100:ILE:HB	1.92	0.52
25:1H:628:G:H2'	25:1H:629:G:H8	1.75	0.52
25:1H:1454:U:O2'	25:1H:1455:G:N7	2.41	0.52
43:D8:74:LYS:HB2	43:D8:83:ARG:HB2	1.90	0.52
52:M8:23:GLU:C	52:M8:25:TYR:H	2.18	0.52
1:1G:392:G:H2'	1:1G:393:A:H8	1.75	0.52
1:1G:411:A:C5	1:1G:413:G:H1'	2.44	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1226:C:OP1	13:4A:91:ARG:NH1	2.43	0.52
1:1G:1255:G:OP1	10:1A:45:ARG:NH2	2.42	0.52
9:82:10:ARG:HH21	9:82:11:LYS:HB2	1.73	0.52
25:14:95:G:O2'	50:G5:48:HIS:ND1	2.36	0.52
25:14:492:A:H2'	25:14:493:G:O4'	2.10	0.52
1:13:235:C:H2'	1:13:236:G:H8	1.73	0.52
1:13:598:U:H4'	8:7E:94:TYR:CG	2.45	0.52
1:13:1175:G:H2'	1:13:1176:A:C8	2.45	0.52
1:13:1224:G:C6	1:13:1322:C:H1'	2.45	0.52
1:13:1261:A:H1'	1:13:1283:G:H5''	1.92	0.52
1:13:1280:A:O2'	1:13:1281:U:OP1	2.22	0.52
2:1E:212:GLN:NE2	2:1E:216:SER:OG	2.38	0.52
25:1H:412:A:N7	25:1H:2411:A:H2	2.07	0.52
25:1H:774:A:H2	25:1H:787:U:O2'	1.92	0.52
25:1H:1635:G:OP1	62:1H:3681:HOH:O	2.18	0.52
29:21:4:ILE:HD13	29:21:28:ALA:HB1	1.92	0.52
37:78:52:GLU:OE2	37:78:58:THR:N	2.43	0.52
1:1G:328:C:O2	1:1G:328:C:H2'	2.07	0.52
1:1G:587:G:N2	1:1G:754:C:OP2	2.43	0.52
2:12:7:VAL:HG13	2:12:8:LYS:H	1.75	0.52
13:4A:86:CYS:HB2	19:AA:73:GLU:HB3	1.91	0.52
20:BA:25:ARG:HG2	20:BA:29:LYS:HE2	1.90	0.52
25:14:2646:C:H2'	25:14:2647:U:O4'	2.09	0.52
25:14:2747:G:H21	25:14:2757:A:H62	1.55	0.52
30:39:195:ASP:HB3	30:39:198:ALA:H	1.75	0.52
36:25:97:ARG:HH21	36:25:99:PHE:HE1	1.58	0.52
43:95:57:VAL:HG12	43:95:58:VAL:H	1.74	0.52
52:I5:40:HIS:N	52:I5:41:PRO:HD3	2.25	0.52
1:13:144:G:H1	1:13:178:C:H42	1.57	0.52
1:13:812:C:O2'	1:13:813:U:P	2.68	0.52
1:13:983:A:H1'	1:13:1049:U:O2	2.10	0.52
22:3K:43:G:H2'	22:3K:44:A:C8	2.45	0.52
25:1H:1078:U:O2'	25:1H:1079:C:OP2	2.24	0.52
25:1H:2294:C:H42	25:1H:2338:G:H1	1.58	0.52
25:1H:2439:A:H5'	25:1H:2439:A:H8	1.74	0.52
25:1H:2867:G:OP2	41:B8:119:LYS:NZ	2.43	0.52
31:41:35:GLU:HA	31:41:99:MET:SD	2.49	0.52
32:51:83:TYR:HD2	32:51:132:ARG:HH22	1.58	0.52
37:78:47:ASP:OD1	37:78:49:ARG:HG2	2.09	0.52
39:98:59:ASP:H	39:98:62:ALA:HB3	1.75	0.52
54:O8:26:ASN:OD1	54:O8:27:LYS:N	2.41	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:780:A:N7	62:1G:1822:HOH:O	2.34	0.52
1:1G:790:A:OP1	57:2L:39:A:O2'	2.26	0.52
1:1G:1139:G:H4'	1:1G:1140:C:O5'	2.09	0.52
1:1G:1271:G:H2'	1:1G:1272:G:H8	1.73	0.52
1:1G:1322:C:O2	1:1G:1322:C:H2'	2.09	0.52
1:1G:1423:G:OP1	36:25:49:ARG:NH1	2.42	0.52
13:4A:23:TYR:CD1	13:4A:70:LEU:HD11	2.45	0.52
25:14:841:A:H2'	25:14:842:G:C8	2.45	0.52
25:14:1090:U:O4	25:14:1102:C:H1'	2.10	0.52
25:14:2443:C:H2'	25:14:2444:G:C8	2.45	0.52
41:75:2:ASN:O	41:75:6:LEU:HB2	2.09	0.52
41:75:123:GLN:C	41:75:125:ARG:H	2.17	0.52
45:B5:63:LYS:HA	45:B5:72:LYS:HA	1.92	0.52
50:G5:4:SER:OG	50:G5:5:GLU:OE2	2.22	0.52
1:13:244:U:H4'	1:13:245:C:O5'	2.09	0.52
25:1H:628:G:H2'	25:1H:629:G:C8	2.45	0.52
25:1H:729:G:OP2	28:11:13:ARG:NH1	2.41	0.52
25:1H:1256:G:O2'	30:31:82:ILE:HD11	2.10	0.52
31:41:170:ARG:O	31:41:174:GLU:HB2	2.10	0.52
34:38:116:ILE:HG13	34:38:117:LEU:HD13	1.92	0.52
36:68:76:ALA:HB3	41:B8:75:ILE:HB	1.92	0.52
37:78:114:ILE:HD11	37:78:130:PHE:HD2	1.75	0.52
51:L8:50:VAL:HB	51:L8:53:LEU:HD12	1.90	0.52
2:12:112:VAL:HG13	2:12:115:LEU:HD23	1.92	0.52
9:82:4:TYR:HD2	9:82:19:LEU:HB3	1.74	0.52
20:BA:51:GLU:HA	20:BA:54:LYS:HE3	1.91	0.52
25:14:451:C:H4'	30:39:52:LYS:HE2	1.91	0.52
25:14:1488:G:H5'	25:14:1489:U:OP2	2.11	0.52
25:14:1543:A:H1'	25:14:1545:A:O4'	2.10	0.52
25:14:1674:G:N1	25:14:1989:G:O6	2.43	0.52
25:14:2133:G:H2'	25:14:2157:G:H22	1.75	0.52
36:25:35:VAL:HG21	36:25:103:ALA:HB3	1.91	0.52
42:85:49:HIS:HA	42:85:52:ARG:HB2	1.92	0.52
45:B5:32:PRO:HA	45:B5:77:LYS:HD2	1.92	0.52
49:F5:23:LYS:HD3	49:F5:29:GLY:H	1.75	0.52
56:M5:35:GLN:OE1	56:M5:36:LYS:N	2.43	0.52
1:13:156:G:H1	1:13:165:C:H42	1.58	0.51
1:13:1355:G:H2'	1:13:1356:G:C8	2.44	0.51
2:1E:96:ARG:H	2:1E:96:ARG:HD2	1.75	0.51
2:1E:223:ILE:HA	2:1E:226:ARG:HB3	1.92	0.51
13:4I:58:GLU:O	13:4I:62:ASN:ND2	2.41	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:27:G:HO2'	25:1H:28:A:H8	1.57	0.51
25:1H:2848:G:O2'	25:1H:2867:G:N2	2.40	0.51
41:B8:29:ARG:HA	41:B8:45:PHE:O	2.11	0.51
47:H8:24:LEU:N	47:H8:39:VAL:O	2.43	0.51
53:N8:31:VAL:HB	53:N8:41:PRO:O	2.11	0.51
1:1G:403:C:O2'	4:32:122:ARG:NH2	2.43	0.51
1:1G:824:C:H1'	8:72:1:MET:HE1	1.91	0.51
1:1G:983:A:N1	1:1G:1222:G:N2	2.58	0.51
1:1G:1355:G:H2'	1:1G:1356:G:C8	2.45	0.51
16:7A:29:ASP:OD1	16:7A:29:ASP:N	2.42	0.51
25:14:1033:U:H4'	25:14:1034:G:OP1	2.10	0.51
25:14:1058:U:H2'	25:14:1059:G:H8	1.73	0.51
25:14:1973:G:H2'	25:14:1974:C:C6	2.45	0.51
25:14:2716:U:H2'	25:14:2717:G:C8	2.45	0.51
29:29:27:LEU:HD12	41:75:6:LEU:HD11	1.92	0.51
37:35:146:VAL:HG22	37:35:147:LEU:H	1.74	0.51
40:65:14:VAL:HG11	40:65:89:ARG:HH12	1.75	0.51
41:75:9:LEU:O	41:75:12:SER:OG	2.18	0.51
1:13:510:A:OP2	4:3E:49:ARG:NH2	2.43	0.51
2:1E:178:ARG:NH1	2:1E:196:LEU:O	2.39	0.51
2:1E:215:LEU:HD13	2:1E:218:ALA:HB3	1.91	0.51
13:4I:65:LYS:NZ	13:4I:73:GLU:OE2	2.43	0.51
25:1H:195:A:N7	62:1H:3730:HOH:O	2.34	0.51
25:1H:380:U:H2'	25:1H:381:G:H8	1.75	0.51
25:1H:1496:A:H8	25:1H:1577:C:O2'	1.92	0.51
25:1H:2428:G:N7	62:1H:3786:HOH:O	2.34	0.51
28:11:112:GLN:H	28:11:115:GLN:NE2	2.08	0.51
30:31:157:VAL:HB	30:31:194:MET:HE3	1.91	0.51
32:51:86:GLU:CD	32:51:87:LEU:H	2.17	0.51
34:38:6:ASN:O	34:38:10:LEU:N	2.35	0.51
35:58:1:MET:HE1	42:C8:95:LEU:HD21	1.93	0.51
39:98:10:LEU:HD13	39:98:40:LYS:HG2	1.92	0.51
45:F8:67:GLY:O	45:F8:68:ARG:HG3	2.10	0.51
53:N8:40:LYS:HZ3	53:N8:46:CYS:HB3	1.74	0.51
1:1G:429:U:H3'	4:32:9:CYS:SG	2.50	0.51
8:72:41:ARG:NH2	8:72:123:GLU:OE2	2.43	0.51
9:82:4:TYR:HB2	9:82:19:LEU:HB2	1.92	0.51
17:8A:8:GLY:HA3	17:8A:21:VAL:HG12	1.92	0.51
25:14:376:C:H42	25:14:398:G:H1	1.58	0.51
25:14:797:C:OP1	30:39:60:SER:OG	2.28	0.51
25:14:1460:A:H4'	25:14:1461:G:OP2	2.11	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:15:130:HIS:ND1	35:15:130:HIS:O	2.42	0.51
36:25:63:VAL:HG12	36:25:106:LEU:HD11	1.92	0.51
47:D5:5:LEU:O	47:D5:59:LEU:HA	2.09	0.51
1:13:381:C:H2'	1:13:382:A:O4'	2.10	0.51
4:3E:64:LEU:HD13	4:3E:198:VAL:HG21	1.92	0.51
5:4E:101:ILE:O	5:4E:120:THR:OG1	2.28	0.51
10:1I:4:ILE:HB	10:1I:74:ILE:HD11	1.91	0.51
25:1H:242:G:H5'	56:Q8:64:TYR:CE2	2.46	0.51
25:1H:330:A:HO2'	25:1H:331:A:H8	1.53	0.51
25:1H:2466:C:H42	25:1H:2484:G:H1	1.58	0.51
29:21:15:PHE:CD2	41:B8:81:PRO:HD2	2.45	0.51
30:31:64:ILE:HG13	30:31:65:TRP:CD1	2.45	0.51
34:38:92:THR:HG21	34:38:126:ALA:HA	1.92	0.51
46:G8:89:PHE:CD1	46:G8:94:LYS:HB3	2.45	0.51
50:K8:47:ASN:C	50:K8:49:LYS:H	2.19	0.51
1:1G:539:A:H2'	1:1G:540:G:C8	2.45	0.51
1:1G:838:G:H1	1:1G:848:C:H42	1.57	0.51
1:1G:1435:G:H2'	1:1G:1436:U:C6	2.46	0.51
3:22:130:VAL:HG21	3:22:157:ILE:HG23	1.91	0.51
5:42:106:PRO:HA	5:42:109:ILE:HD12	1.93	0.51
25:14:270(P):C:H2'	25:14:270(Q):C:C6	2.46	0.51
25:14:626:U:H5''	25:14:627:A:H5'	1.90	0.51
25:14:1417:C:H42	25:14:1581:G:H1	1.57	0.51
25:14:2472:G:N1	25:14:2477:C:OP1	2.42	0.51
25:14:2830:G:O2'	25:14:2883:A:N1	2.42	0.51
32:59:29:PRO:HD2	32:59:79:VAL:HB	1.93	0.51
39:55:37:THR:HG22	39:55:39:PRO:HD2	1.91	0.51
1:13:659:U:H2'	1:13:660:G:H8	1.75	0.51
1:13:1396:A:H4'	1:13:1397:C:H5''	1.92	0.51
20:BI:26:ASN:O	20:BI:30:LYS:HB2	2.10	0.51
25:1H:259:G:H21	25:1H:621:A:H8	1.58	0.51
25:1H:289:A:H2'	25:1H:290:G:O4'	2.10	0.51
25:1H:526:A:N3	25:1H:2044:C:H1'	2.26	0.51
25:1H:1497:U:H5''	25:1H:1498:C:H5	1.74	0.51
25:1H:2567:G:H2'	25:1H:2568:C:C6	2.46	0.51
47:H8:10:ARG:O	47:H8:36:LYS:HB2	2.10	0.51
47:H8:163:LEU:HD22	47:H8:167:PRO:HA	1.91	0.51
50:K8:24:LEU:HD22	50:K8:60:LEU:HD11	1.92	0.51
1:1G:952:U:H2'	1:1G:953:G:C8	2.45	0.51
1:1G:1062:U:H2'	1:1G:1063:C:C6	2.45	0.51
3:22:182:ILE:HG12	3:22:203:PHE:HB2	1.91	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:3A:8:VAL:HG11	17:8A:36:ILE:HG21	1.93	0.51
25:14:883:G:N2	25:14:893:C:H42	2.06	0.51
25:14:1427:A:H4'	25:14:1428:C:O5'	2.10	0.51
29:29:38:THR:H	29:29:42:ASP:HB2	1.74	0.51
31:49:6:ALA:N	52:I5:23:GLU:OE2	2.34	0.51
35:15:128:HIS:HB2	35:15:129:PRO:HD2	1.92	0.51
37:35:84:ASN:HB3	37:35:116:GLY:HA3	1.91	0.51
39:55:118:GLU:OE2	39:55:118:GLU:N	2.41	0.51
44:A5:12:ILE:HD13	44:A5:17:VAL:HG22	1.91	0.51
47:D5:150:LEU:HD21	47:D5:171:ILE:HG23	1.91	0.51
1:13:1318:A:O2'	19:AI:37:ARG:HB2	2.10	0.51
3:2E:5:ILE:HG21	10:1I:51:ARG:HH22	1.75	0.51
25:1H:1586:A:H3'	25:1H:1587:A:C8	2.45	0.51
25:1H:2356:C:O3'	48:I8:20:ARG:HD3	2.10	0.51
54:O8:13:CYS:SG	54:O8:22:ALA:HB3	2.50	0.51
56:Q8:50:LEU:HB3	56:Q8:55:ALA:HB2	1.91	0.51
1:1G:398:C:OP2	62:1G:1809:HOH:O	2.18	0.51
1:1G:969:A:H2'	1:1G:970:C:O4'	2.11	0.51
1:1G:1306:A:H1'	1:1G:1332:A:N1	2.25	0.51
1:1G:1387:G:H2'	1:1G:1388:C:C6	2.46	0.51
1:1G:1517:G:H3'	1:1G:1518:MA6:H8	1.93	0.51
5:42:3:GLU:O	5:42:5:ASP:N	2.39	0.51
12:3A:114:ARG:NE	12:3A:120:LYS:O	2.44	0.51
22:3L:64:G:H2'	22:3L:65:G:H8	1.76	0.51
25:14:536:A:H2'	25:14:537:C:O4'	2.11	0.51
25:14:1020:A:OP1	25:14:1034:G:N2	2.38	0.51
25:14:2776:A:OP1	25:14:2776:A:H3'	2.10	0.51
25:14:2882:A:C8	39:55:96:ARG:NH2	2.79	0.51
35:15:104:LYS:HA	35:15:107:LEU:HD12	1.92	0.51
40:65:23:ARG:NH1	40:65:84:GLN:HG2	2.26	0.51
49:F5:91:LYS:HA	49:F5:91:LYS:NZ	2.25	0.51
1:13:303:A:HO2'	1:13:555:C:HO2'	1.57	0.51
1:13:947:G:O3'	13:4I:109:THR:OG1	2.29	0.51
1:13:1423:G:OP1	36:68:49:ARG:NH2	2.44	0.51
4:3E:193:ASP:OD1	4:3E:193:ASP:N	2.44	0.51
25:1H:273(C):C:H42	25:1H:363(C):G:H1	1.58	0.51
25:1H:2308:G:H22	25:1H:2311:A:H2	1.59	0.51
25:1H:2313:C:H5''	31:41:91:ARG:HD2	1.93	0.51
25:1H:2336:A:H61	48:I8:43:THR:HG21	1.75	0.51
25:1H:2577:A:OP2	53:N8:3:LYS:NZ	2.43	0.51
25:1H:2737:G:H2'	25:1H:2738:A:H8	1.75	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:41:34:LEU:HB2	31:41:172:LEU:HD21	1.92	0.51
39:98:105:ARG:C	39:98:107:ASP:H	2.19	0.51
42:C8:92:ARG:C	42:C8:94:ASN:H	2.18	0.51
54:O8:9:LEU:HD13	54:O8:10:LEU:H	1.75	0.51
1:1G:502:G:H2'	1:1G:503:C:O4'	2.11	0.51
1:1G:718:G:H5'	11:2A:117:ASN:CG	2.35	0.51
1:1G:1007:C:H3'	1:1G:1008:C:H5''	1.93	0.51
1:1G:1053:G:O6	1:1G:1199:U:H2'	2.11	0.51
7:62:120:ILE:O	7:62:124:LEU:HB2	2.10	0.51
25:14:372:G:O2'	25:14:400:G:N1	2.42	0.51
25:14:872:A:OP1	38:45:5:ARG:NH2	2.43	0.51
25:14:1114:G:H2'	25:14:1115:G:C8	2.46	0.51
25:14:2784:C:H2'	25:14:2785:C:H6	1.76	0.51
26:1J:60:C:H2'	26:1J:61:G:H8	1.75	0.51
32:59:9:ILE:HD13	32:59:51:ARG:HE	1.75	0.51
40:65:84:GLN:HA	40:65:109:GLY:CA	2.41	0.51
42:85:81:HIS:CE1	42:85:84:LYS:HD3	2.46	0.51
46:C5:54:LYS:HD2	46:C5:55:TYR:HD1	1.76	0.51
1:13:392:G:H2'	1:13:393:A:C8	2.46	0.51
1:13:992:U:H1'	1:13:993:G:OP2	2.11	0.51
22:1K:49:C:HO2'	22:1K:50:G:P	2.33	0.51
25:1H:676:A:H2	25:1H:802:A:H61	1.59	0.51
25:1H:1301:A:O2'	25:1H:1302:A:H3'	2.10	0.51
25:1H:1484:G:O6	62:1H:3684:HOH:O	2.19	0.51
25:1H:1994:C:OP1	62:1H:3687:HOH:O	2.19	0.51
25:1H:2330:G:H2'	25:1H:2331:G:O4'	2.11	0.51
25:1H:2712:U:O2'	25:1H:2712(A):A:H8	1.93	0.51
25:1H:2816:C:O2	25:1H:2883:A:O2'	2.29	0.51
27:71:49:ILE:HG12	27:71:208:PHE:HE1	1.75	0.51
31:41:7:LEU:N	31:41:104:GLU:OE2	2.35	0.51
33:61:104:GLN:O	33:61:105:HIS:ND1	2.43	0.51
49:J8:60:PHE:HB3	49:J8:62:VAL:HG13	1.92	0.51
1:1G:1203:C:H2'	1:1G:1204:A:O4'	2.09	0.51
1:1G:1376:U:H2'	1:1G:1377:A:C8	2.45	0.51
5:42:73:ASN:OD1	5:42:73:ASN:N	2.44	0.51
13:4A:12:ASN:O	13:4A:12:ASN:ND2	2.44	0.51
16:7A:15:PRO:O	16:7A:16:HIS:ND1	2.43	0.51
25:14:1002:G:H2'	25:14:1003:G:O4'	2.11	0.51
25:14:2133:G:H2'	25:14:2157:G:N2	2.25	0.51
25:14:2328:A:H2'	25:14:2329:G:C8	2.46	0.51
31:49:128:ARG:HB2	31:49:130:ASN:HD21	1.76	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:69:26:ALA:HA	33:69:30:LEU:HB2	1.92	0.51
1:13:236:G:H5''	17:8I:42:TYR:OH	2.11	0.51
1:13:1372:U:OP1	9:8E:72:GLY:N	2.42	0.51
12:3I:34:CYS:HB3	12:3I:55:VAL:HG22	1.93	0.51
25:1H:576:U:OP1	62:1H:3675:HOH:O	2.18	0.51
25:1H:2331:G:O2'	48:I8:43:THR:HG22	2.10	0.51
25:1H:2630:G:O4'	25:1H:2894:G:H1'	2.11	0.51
27:71:7:TYR:HA	27:71:10:LEU:HD13	1.93	0.51
27:71:173:ALA:HB3	27:71:192:PHE:CE1	2.46	0.51
33:61:81:VAL:HB	33:61:83:ALA:HB2	1.91	0.51
37:78:138:LEU:HD11	37:78:144:GLU:HG3	1.93	0.51
46:G8:15:VAL:HG21	46:G8:42:VAL:HG21	1.93	0.51
46:G8:35:TYR:CE2	46:G8:69:ALA:HB3	2.46	0.51
48:I8:40:GLN:HE21	48:I8:43:THR:HA	1.74	0.51
1:1G:715:A:H2'	1:1G:716:A:C8	2.46	0.51
1:1G:766:A:H2'	1:1G:767:A:O4'	2.10	0.51
1:1G:1082:G:H8	1:1G:1082:G:OP2	1.92	0.51
2:12:21:ARG:O	2:12:23:ARG:HB2	2.11	0.51
4:32:173:TRP:CZ3	4:32:193:ASP:HB3	2.46	0.51
57:2L:24:C:H2'	57:2L:25:U:C6	2.45	0.51
25:14:108:U:H2'	25:14:109:G:C8	2.45	0.51
25:14:817:C:H2'	25:14:818:G:O4'	2.11	0.51
25:14:1592:C:H2'	25:14:1593:G:C8	2.46	0.51
25:14:2052:G:H4'	29:29:143:ASN:O	2.11	0.51
25:14:2150:U:H2'	25:14:2151:G:C8	2.46	0.51
25:14:2412:A:H2'	25:14:2413:G:O4'	2.11	0.51
31:49:6:ALA:H	52:I5:23:GLU:CD	2.19	0.51
45:B5:65:ARG:HD3	45:B5:70:LEU:HG	1.93	0.51
47:D5:165:VAL:HG13	47:D5:166:SER:H	1.76	0.51
1:13:973:G:OP1	10:1I:57:LYS:HE3	2.10	0.51
1:13:1390:U:H2'	1:13:1391:U:C6	2.46	0.51
8:7E:112:LEU:HB3	8:7E:133:LEU:HA	1.91	0.51
13:4I:3:ARG:HD2	13:4I:9:ILE:HG21	1.92	0.51
16:7I:74:LEU:HA	16:7I:77:ALA:HB3	1.93	0.51
18:9I:31:LEU:H	18:9I:31:LEU:HD23	1.76	0.51
19:AI:9:VAL:HG23	19:AI:10:PHE:H	1.75	0.51
23:2K:54:G:H2'	23:2K:55:5MU:H6	1.75	0.51
25:1H:1224:G:OP2	43:D8:66:ARG:NH2	2.44	0.51
25:1H:1364:G:C5	49:J8:2:SER:HB3	2.46	0.51
25:1H:1586:A:H3'	25:1H:1587:A:H8	1.75	0.51
25:1H:1614:A:P	25:1H:1614:A:H8	2.33	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:1683:C:H2'	25:1H:1684:C:C6	2.46	0.51
25:1H:1711:C:H42	25:1H:1747:G:H1	1.57	0.51
31:41:11:TYR:O	31:41:16:ARG:HB2	2.11	0.51
1:1G:752:G:H1'	1:1G:754:C:H41	1.75	0.51
1:1G:1288:A:H2'	1:1G:1289:A:C8	2.46	0.51
2:12:61:LEU:HD11	2:12:66:GLY:HA3	1.92	0.51
25:14:1022:G:H22	25:14:1142(A):A:H2	1.58	0.51
25:14:1608:A:H1'	25:14:1610:A:OP2	2.11	0.51
25:14:1678:G:H22	25:14:1989:G:N2	2.08	0.51
25:14:2870:C:H2'	25:14:2871:C:O4'	2.10	0.51
37:35:15:ARG:HG3	37:35:16:ARG:H	1.76	0.51
37:35:97:PRO:C	37:35:99:LEU:H	2.18	0.51
47:D5:16:SER:O	47:D5:20:ARG:HG3	2.11	0.51
4:3E:103:ASN:OD1	4:3E:114:ARG:NH2	2.41	0.51
13:4I:54:VAL:HG12	13:4I:58:GLU:OE2	2.10	0.51
19:AI:31:ILE:HG23	19:AI:49:ILE:HA	1.93	0.51
25:1H:363(B):G:H2'	25:1H:363(C):G:H8	1.75	0.51
25:1H:1149:G:H2'	25:1H:1150:C:C6	2.45	0.51
25:1H:1412:A:H2'	25:1H:1413:G:C8	2.47	0.51
25:1H:2732:G:H3'	25:1H:2733:A:O4'	2.11	0.51
41:B8:107:ASP:O	41:B8:111:ARG:NH1	2.44	0.51
44:E8:24:ILE:HG21	44:E8:36:LEU:HD21	1.93	0.51
1:1G:1399:C:N3	1:1G:1502:A:N6	2.58	0.51
13:4A:91:ARG:NH2	13:4A:97:PRO:O	2.44	0.51
20:BA:26:ASN:HA	20:BA:29:LYS:HG2	1.93	0.51
25:14:676:A:H2	25:14:802:A:H61	1.56	0.51
25:14:1048:A:OP2	25:14:1109:C:N4	2.43	0.51
25:14:1085:A:O2'	25:14:1086:A:OP1	2.25	0.51
25:14:1245:G:OP1	37:35:13:ASN:ND2	2.43	0.51
25:14:2331:G:O2'	48:E5:43:THR:HG22	2.11	0.51
25:14:2471:C:H3'	25:14:2472:G:C8	2.45	0.51
30:39:25:PRO:HB3	30:39:28:ILE:HG23	1.93	0.51
35:15:20:GLY:HA2	35:15:61:ARG:HG2	1.92	0.51
54:K5:19:ARG:NH2	54:K5:52:VAL:HG21	2.26	0.51
1:13:156:G:H2'	1:13:157:G:H8	1.75	0.50
1:13:404:U:H2'	1:13:405:U:C6	2.46	0.50
1:13:474:G:H2'	1:13:475:G:C8	2.46	0.50
3:2E:11:ARG:HB3	3:2E:15:THR:HB	1.93	0.50
4:3E:108:LEU:HD11	4:3E:174:LEU:HD22	1.93	0.50
7:6E:97:GLN:O	7:6E:101:LEU:HG	2.11	0.50
10:1I:45:ARG:HB2	10:1I:65:LEU:HB3	1.92	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:1055:G:H22	25:1H:1104:C:N4	2.09	0.50
25:1H:1588:C:H2'	25:1H:1589:C:C6	2.45	0.50
25:1H:1652:A:OP1	39:98:8:ARG:NH1	2.40	0.50
47:H8:19:ARG:HD3	47:H8:19:ARG:H	1.76	0.50
48:I8:7:LEU:HB2	48:I8:11:ARG:HG2	1.93	0.50
2:12:142:LEU:HA	2:12:145:LEU:HD23	1.93	0.50
25:14:855:G:H1	25:14:922:U:H3	1.58	0.50
39:55:48:VAL:HA	39:55:51:LEU:HD12	1.92	0.50
41:75:74:ARG:HD3	41:75:76:PHE:CZ	2.45	0.50
43:95:35:LEU:HD23	43:95:37:VAL:CG2	2.41	0.50
54:K5:23:THR:HB	56:M5:36:LYS:HD2	1.91	0.50
54:K5:39:TYR:CD1	54:K5:41:PRO:HD3	2.47	0.50
1:13:431:A:H2'	1:13:432:A:O4'	2.12	0.50
1:13:1349:A:OP2	9:8E:118:LYS:NZ	2.38	0.50
4:3E:149:ALA:O	4:3E:153:ARG:N	2.44	0.50
8:7E:83:ILE:HG13	8:7E:137:VAL:HG22	1.92	0.50
22:1K:16:C:O2'	22:1K:61:U:O3'	2.24	0.50
22:3K:29:C:H2'	22:3K:30:G:C8	2.46	0.50
25:1H:520:G:H2'	25:1H:521:G:C8	2.46	0.50
25:1H:972:G:H3'	25:1H:973:A:H2'	1.94	0.50
25:1H:1071:G:H1'	25:1H:1089:G:H2'	1.92	0.50
25:1H:1754:C:OP2	41:B8:113:LYS:HE3	2.12	0.50
25:1H:2614:A:H3'	62:1H:3838:HOH:O	2.10	0.50
27:71:212:VAL:HG21	27:71:226:PRO:HB3	1.93	0.50
46:G8:99:CYS:SG	46:G8:105:ALA:HB3	2.51	0.50
1:1G:6:G:H4'	1:1G:298:A:H4'	1.92	0.50
1:1G:620:C:H2'	1:1G:621:A:O4'	2.10	0.50
1:1G:1090:U:H2'	1:1G:1091:U:C6	2.45	0.50
1:1G:1290:G:O3'	7:62:37:ASN:ND2	2.44	0.50
1:1G:1347:G:O2'	1:1G:1348:U:O5'	2.29	0.50
18:9A:58:LEU:HD12	18:9A:63:GLN:HA	1.94	0.50
20:BA:71:THR:HG22	20:BA:72:LEU:HD23	1.92	0.50
25:14:125:G:C6	55:L5:10:ARG:HG3	2.46	0.50
25:14:1693:U:O2'	28:19:14:ARG:NH2	2.44	0.50
25:14:2074:U:OP1	62:14:3551:HOH:O	2.19	0.50
25:14:2111:C:N4	25:14:2147:G:H21	2.08	0.50
25:14:2758:A:C2	25:14:2759:G:H1'	2.46	0.50
25:14:2822:G:OP2	62:14:3552:HOH:O	2.20	0.50
39:55:59:ASP:OD1	39:55:59:ASP:N	2.40	0.50
40:65:24:LEU:HD12	40:65:41:ASP:HB2	1.93	0.50
41:75:55:ASN:O	41:75:55:ASN:ND2	2.43	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1109:C:H2'	1:13:1110:A:O4'	2.11	0.50
1:13:1154:G:H2'	1:13:1155:G:O4'	2.12	0.50
1:13:1330:U:H4'	13:4I:23:TYR:HE1	1.76	0.50
8:7E:35:ILE:HG22	8:7E:111:ILE:HG21	1.93	0.50
8:7E:120:THR:OG1	8:7E:123:GLU:OE1	2.22	0.50
22:3K:3:C:H2'	22:3K:4:G:H8	1.76	0.50
25:1H:263:C:H2'	25:1H:264:C:O4'	2.12	0.50
25:1H:635:C:O2'	25:1H:639:U:OP1	2.24	0.50
25:1H:792:G:N3	25:1H:2072:G:O2'	2.41	0.50
25:1H:1107:G:OP1	34:38:56:ASN:HB2	2.11	0.50
27:71:209:LEU:HD13	27:71:226:PRO:HB2	1.92	0.50
32:51:10:PRO:HD2	32:51:50:VAL:O	2.11	0.50
35:58:53:VAL:HG22	35:58:121:LYS:HB2	1.93	0.50
36:68:25:LEU:HD12	36:68:38:VAL:HG22	1.93	0.50
36:68:68:GLU:CD	36:68:68:GLU:H	2.19	0.50
1:1G:45:U:H2'	1:1G:46:G:C8	2.46	0.50
1:1G:828:A:H2'	1:1G:829:G:O4'	2.10	0.50
1:1G:1277:C:HO2'	1:1G:1279:A:H8	1.58	0.50
1:1G:1314:C:H2'	1:1G:1315:U:C6	2.46	0.50
2:12:8:LYS:HB3	2:12:217:ARG:CZ	2.41	0.50
2:12:162:ILE:HD11	2:12:184:VAL:HA	1.92	0.50
2:12:178:ARG:HH12	8:72:68:ARG:HH22	1.59	0.50
7:62:76:ARG:HD2	7:62:89:MET:HG3	1.92	0.50
8:72:37:ARG:O	8:72:41:ARG:N	2.32	0.50
11:2A:99:GLN:HG2	11:2A:105:VAL:HG21	1.94	0.50
11:2A:123:LYS:NZ	11:2A:123:LYS:HB3	2.25	0.50
17:8A:67:LYS:O	17:8A:69:LYS:N	2.44	0.50
25:14:26:G:C6	25:14:27:G:N1	2.79	0.50
25:14:727:A:OP1	25:14:1431:U:O2'	2.28	0.50
25:14:2698:U:H2'	25:14:2699:C:C6	2.46	0.50
37:35:122:PRO:HB3	37:35:141:ALA:HB1	1.93	0.50
42:85:17:ILE:HD12	42:85:32:PHE:HE1	1.76	0.50
49:F5:75:GLU:HB3	49:F5:76:ARG:NH2	2.21	0.50
1:13:167:G:H2'	1:13:168:G:C8	2.46	0.50
1:13:792:A:H4'	1:13:793:U:O5'	2.12	0.50
1:13:1104:G:H4'	2:1E:111:ARG:NH1	2.27	0.50
1:13:1148:U:H2'	1:13:1149:C:O4'	2.12	0.50
25:1H:363(A):A:H2'	25:1H:363(B):G:C8	2.46	0.50
25:1H:1266:G:O2'	25:1H:1267:U:OP2	2.27	0.50
32:51:101:ARG:HH22	32:51:122:THR:HA	1.77	0.50
35:58:13:TRP:O	35:58:135:PRO:HD2	2.10	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:68:26:LYS:HB2	36:68:30:ALA:CB	2.42	0.50
54:08:11:LEU:HD11	54:08:51:GLU:HG3	1.93	0.50
1:1G:299:G:H2'	1:1G:300:A:C8	2.47	0.50
1:1G:882:C:OP2	12:3A:10:LYS:NZ	2.45	0.50
1:1G:1053:G:H4'	1:1G:1054:C:H5'	1.94	0.50
1:1G:1137:C:H5'	1:1G:1138:G:C2	2.47	0.50
9:82:104:ARG:HG3	9:82:104:ARG:O	2.12	0.50
11:2A:11:LYS:NZ	25:14:2144:U:OP2	2.39	0.50
12:3A:100:GLY:N	12:3A:104:ALA:O	2.35	0.50
18:9A:37:VAL:HG12	18:9A:40:LEU:HD12	1.93	0.50
19:AA:36:ARG:HG3	19:AA:71:LEU:HB2	1.93	0.50
22:3L:57:C:H42	25:14:2112:G:N2	2.10	0.50
25:14:141:A:C8	25:14:1408:C:H1'	2.47	0.50
25:14:186:G:H2'	25:14:187:G:C8	2.46	0.50
25:14:265:A:H2'	25:14:266:G:H4'	1.93	0.50
25:14:861:A:N3	26:1J:79:C:O2'	2.42	0.50
25:14:1161:C:H2'	25:14:1162:G:H8	1.76	0.50
25:14:1812:A:O2'	28:19:45:ASN:N	2.43	0.50
25:14:2186:G:H2'	25:14:2187:G:C8	2.46	0.50
25:14:2779:U:H1'	25:14:2781:A:C6	2.46	0.50
30:39:150:GLY:HA2	30:39:172:TRP:CD2	2.47	0.50
35:15:46:VAL:HG13	35:15:48:MET:HG3	1.93	0.50
50:G5:10:LEU:HD21	50:G5:14:ARG:HH21	1.77	0.50
51:H5:50:VAL:HB	51:H5:53:LEU:HD12	1.93	0.50
1:13:1186:G:H21	14:5I:61:TRP:HA	1.76	0.50
1:13:1320:C:N4	19:AI:36:ARG:HG3	2.27	0.50
3:2E:156:ARG:HG2	3:2E:163:ALA:HB2	1.94	0.50
23:2K:32:G:H2'	23:2K:33:OMC:C6	2.45	0.50
25:1H:2122:U:H2'	25:1H:2123:G:H8	1.76	0.50
1:1G:963:G:H21	10:1A:55:LYS:HE2	1.76	0.50
1:1G:1126:U:H5''	1:1G:1280:A:N7	2.26	0.50
1:1G:1517:G:H1'	25:14:1919:A:O3'	2.11	0.50
9:82:34:ASN:O	9:82:37:PHE:N	2.32	0.50
17:8A:64:PRO:HA	17:8A:70:ARG:HG3	1.94	0.50
25:14:636:G:C6	37:35:115:LEU:HD11	2.46	0.50
25:14:1139:G:O2'	25:14:1143:A:N6	2.36	0.50
25:14:2306:C:N4	25:14:2307:G:O6	2.45	0.50
31:49:114:ILE:HG12	31:49:140:ILE:HG21	1.94	0.50
33:69:58:LEU:O	33:69:62:LYS:HG2	2.12	0.50
36:25:23:ARG:HG3	36:25:24:VAL:N	2.26	0.50
54:K5:45:LYS:HG3	54:K5:47:THR:HG23	1.93	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1492:A:H4'	12:3I:44:LYS:HD3	1.93	0.50
2:1E:50:GLU:O	2:1E:54:THR:OG1	2.28	0.50
3:2E:58:GLU:O	3:2E:64:VAL:HA	2.12	0.50
13:4I:5:ALA:HB2	13:4I:61:GLU:HG2	1.93	0.50
22:3K:4:G:N2	22:3K:71:G:H1'	2.26	0.50
22:3K:19:G:O3'	22:3K:61:U:N3	2.45	0.50
25:1H:1277:G:O2'	39:98:24:GLN:OE1	2.28	0.50
25:1H:1376:C:OP2	62:1H:3691:HOH:O	2.20	0.50
25:1H:2522:U:O2'	25:1H:2647:U:OP1	2.23	0.50
31:4I:114:ILE:HG22	31:4I:115:ARG:H	1.77	0.50
34:38:4:LYS:HG3	34:38:6:ASN:HB2	1.94	0.50
35:58:30:ILE:HG23	35:58:52:VAL:HG11	1.94	0.50
35:58:114:ARG:O	35:58:115:ARG:HB3	2.10	0.50
1:1G:1187:G:OP1	9:82:113:LYS:NZ	2.40	0.50
1:1G:1347:G:N2	1:1G:1374:A:OP2	2.42	0.50
2:12:60:ASP:OD1	2:12:64:ARG:NH1	2.45	0.50
25:14:310:A:O2'	62:14:3535:HOH:O	2.20	0.50
25:14:507:A:H5'	25:14:508:G:H2'	1.92	0.50
25:14:1012:U:N3	25:14:1143:A:H2	2.09	0.50
28:19:161:THR:HG22	28:19:178:PRO:HG2	1.94	0.50
29:29:98:PRO:HG3	29:29:174:ASP:HA	1.93	0.50
30:39:15:SER:OG	30:39:16:GLY:N	2.45	0.50
33:69:2:LYS:HD3	33:69:20:ASP:HB3	1.93	0.50
1:13:688:G:H2'	1:13:689:C:C6	2.45	0.50
1:13:737:A:H2'	1:13:738:C:H6	1.76	0.50
1:13:944:G:N2	1:13:1338:G:N7	2.60	0.50
22:1K:31:G:H2'	22:1K:32:G:C8	2.44	0.50
25:1H:221:A:H4'	25:1H:222:A:O5'	2.11	0.50
25:1H:489:G:N2	25:1H:1321:A:OP1	2.42	0.50
25:1H:528:A:H2	25:1H:2043:C:H5'	1.76	0.50
25:1H:1894:C:H2'	25:1H:1895:C:H6	1.76	0.50
25:1H:2114:A:H2'	25:1H:2168:G:C8	2.46	0.50
27:71:173:ALA:HB1	27:71:174:PRO:HD2	1.92	0.50
32:51:121:ILE:HG13	32:51:144:VAL:HG21	1.94	0.50
44:E8:110:LYS:HG3	44:E8:111:HIS:H	1.77	0.50
54:O8:15:GLU:HA	54:O8:49:HIS:HA	1.93	0.50
1:1G:142:G:H2'	1:1G:143:A:C8	2.47	0.50
2:12:165:VAL:HG23	2:12:166:ASP:H	1.77	0.50
3:22:152:ILE:HB	3:22:199:LYS:HB2	1.94	0.50
9:82:19:LEU:HD12	9:82:84:ALA:HB1	1.93	0.50
10:1A:4:ILE:HG22	10:1A:6:ILE:HG23	1.94	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:3A:15:VAL:O	12:3A:16:ARG:HB2	2.11	0.50
18:9A:21:LYS:HE2	18:9A:24:ALA:HB2	1.94	0.50
22:3L:51:U:H3	22:3L:65:G:H1	1.60	0.50
25:14:2425:A:H4'	25:14:2426:A:H5''	1.94	0.50
25:14:2643:G:H2'	25:14:2644:G:O4'	2.11	0.50
29:29:4:ILE:HD11	29:29:29:GLY:HA2	1.93	0.50
29:29:23:VAL:HG12	29:29:173:VAL:HG21	1.93	0.50
33:69:114:LEU:HA	33:69:130:TYR:HB2	1.94	0.50
35:15:3:THR:HG22	35:15:4:TYR:H	1.77	0.50
46:C5:6:HIS:ND1	46:C5:7:VAL:HG13	2.27	0.50
47:D5:45:ASP:O	47:D5:49:ARG:HG2	2.12	0.50
1:13:636:U:H2'	1:13:637:G:H8	1.75	0.50
1:13:719:C:H1'	18:9I:49:LYS:HB3	1.93	0.50
2:1E:18:GLY:H	2:1E:42:ILE:HD12	1.76	0.50
3:2E:116:VAL:HA	3:2E:119:ARG:HB3	1.93	0.50
22:1K:29:C:H2'	22:1K:30:G:C8	2.46	0.50
22:1K:49:C:C2	22:1K:60:A:H1'	2.47	0.50
23:2K:76:C:C2	25:1H:2251:OMG:N2	2.77	0.50
25:1H:2702:U:H6	25:1H:2702:U:OP1	1.94	0.50
47:H8:45:ASP:O	47:H8:49:ARG:HG2	2.11	0.50
47:H8:54:HIS:HB3	47:H8:101:PRO:HD3	1.94	0.50
1:1G:44:G:O6	62:1G:1810:HOH:O	2.19	0.50
1:1G:298:A:OP1	4:32:209:ARG:NH2	2.45	0.50
1:1G:719:C:O2'	18:9A:49:LYS:HB3	2.12	0.50
1:1G:1218:C:OP2	14:5A:9:LYS:NZ	2.38	0.50
5:42:60:TYR:O	5:42:64:ARG:NE	2.40	0.50
6:52:24:GLU:OE2	6:52:28:ARG:NH1	2.32	0.50
7:62:20:ASP:O	7:62:23:VAL:N	2.44	0.50
7:62:73:MET:HE3	7:62:90:GLU:HG2	1.94	0.50
20:BA:53:LEU:HA	20:BA:56:MET:HB2	1.93	0.50
25:14:1159:U:OP1	51:H5:30:ARG:NH1	2.45	0.50
25:14:2012:G:O3'	44:A5:96:ILE:HG13	2.11	0.50
25:14:2318:G:H1	40:65:2:ALA:N	2.10	0.50
26:1J:16:G:H1	26:1J:68:C:H42	1.59	0.50
32:59:70:THR:HA	32:59:73:ALA:HB3	1.93	0.50
40:65:67:ARG:HH21	40:65:103:GLU:HG3	1.77	0.50
42:85:31:SER:HB3	42:85:34:LYS:HB2	1.93	0.50
51:H5:54:VAL:HG12	51:H5:56:VAL:HG22	1.93	0.50
1:13:235:C:H2'	1:13:236:G:C8	2.47	0.50
1:13:481:G:O2'	1:13:482:A:O5'	2.29	0.50
1:13:636:U:H2'	1:13:637:G:C8	2.46	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1358:U:H5''	14:5I:33:VAL:O	2.11	0.50
2:1E:168:THR:OG1	2:1E:169:LYS:N	2.45	0.50
10:1I:49:VAL:HG22	14:5I:41:ARG:HG3	1.94	0.50
14:5I:13:THR:N	14:5I:14:PRO:HD2	2.27	0.50
15:6I:24:SER:HB2	15:6I:27:VAL:HG23	1.94	0.50
19:AI:77:THR:HG22	19:AI:78:ARG:HG2	1.92	0.50
25:1H:270(M):U:H1'	25:1H:270(N):G:C6	2.47	0.50
25:1H:528:A:C2	25:1H:2043:C:H5'	2.47	0.50
25:1H:1287:A:N7	39:98:107:ASP:HB2	2.26	0.50
25:1H:2144:U:O2'	25:1H:2147:G:O6	2.15	0.50
25:1H:2469:A:O2'	38:88:56:ARG:HG2	2.12	0.50
30:31:34:TRP:CE3	37:78:8:PRO:HB3	2.46	0.50
30:31:68:LYS:C	30:31:70:THR:H	2.20	0.50
34:38:29:TYR:O	34:38:31:GLY:N	2.44	0.50
37:78:80:TYR:OH	37:78:111:ARG:NH1	2.45	0.50
41:B8:78:LEU:O	41:B8:78:LEU:HD13	2.12	0.50
1:1G:935:A:H2'	1:1G:936:C:C6	2.46	0.50
1:1G:1213:A:N6	1:1G:1215:G:N3	2.59	0.50
9:82:76:ALA:O	9:82:80:GLY:N	2.38	0.50
25:14:134:C:H42	25:14:145:G:H1	1.60	0.50
25:14:570:G:H2'	25:14:2030:A:C5	2.47	0.50
25:14:1252:G:N3	42:85:33:ARG:HD2	2.27	0.50
25:14:1418:G:OP1	25:14:1588:C:O2'	2.30	0.50
25:14:2815:C:H2'	25:14:2816:C:C6	2.46	0.50
29:29:1:MET:HE2	29:29:202:LYS:HE2	1.94	0.50
35:15:112:LEU:HD23	35:15:113:GLY:N	2.26	0.50
36:25:68:GLU:H	36:25:68:GLU:CD	2.19	0.50
55:L5:12:ARG:NH2	55:L5:44:PRO:HB3	2.27	0.50
1:13:1322:C:O2'	1:13:1323:G:H5'	2.12	0.49
1:13:1348:U:N3	1:13:1374:A:H2	2.09	0.49
25:1H:856:C:N4	25:1H:921:G:H1	2.10	0.49
25:1H:1429:G:H1'	25:1H:1568:G:H1'	1.94	0.49
29:21:4:ILE:HG12	29:21:5:LEU:H	1.76	0.49
32:51:4:ILE:HB	32:51:6:ARG:HD3	1.93	0.49
41:B8:33:LYS:HE3	41:B8:84:GLN:HB2	1.93	0.49
56:Q8:16:ILE:HD13	56:Q8:59:LYS:HG2	1.94	0.49
4:32:133:VAL:HG13	4:32:135:LEU:HD22	1.93	0.49
13:4A:14:ARG:HA	13:4A:44:ARG:HA	1.93	0.49
19:AA:40:ILE:HG12	19:AA:69:HIS:O	2.11	0.49
25:14:637:A:H4'	25:14:638:G:O5'	2.12	0.49
25:14:1106:G:H3'	25:14:1107:G:H8	1.77	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:2165:G:H5'	25:14:2166:G:OP2	2.12	0.49
25:14:2483:C:N3	38:45:124:LYS:NZ	2.58	0.49
25:14:2697:G:H2'	25:14:2698:U:O4'	2.12	0.49
29:29:101:ARG:CZ	29:29:171:GLU:HB2	2.41	0.49
30:39:124:LEU:O	30:39:126:VAL:HG13	2.11	0.49
41:75:60:THR:HG22	41:75:77:PRO:HA	1.93	0.49
46:C5:55:TYR:CD1	46:C5:55:TYR:N	2.79	0.49
49:F5:41:ARG:HB2	49:F5:43:TYR:CE1	2.47	0.49
1:13:192:U:H5'	20:BI:102:GLY:HA2	1.93	0.49
1:13:564:C:H5'	17:8I:32:TYR:CE1	2.47	0.49
1:13:859:A:H2'	1:13:860:A:O4'	2.12	0.49
2:1E:92:TYR:CE1	2:1E:151:GLY:HA3	2.47	0.49
25:1H:250:G:P	37:78:60:MET:HE1	2.52	0.49
25:1H:1301:A:HO2'	25:1H:1302:A:C3'	2.24	0.49
25:1H:1859:A:N6	25:1H:1883:G:O2'	2.46	0.49
25:1H:1925:C:H2'	25:1H:1926:U:O4'	2.12	0.49
25:1H:2737:G:H2'	25:1H:2738:A:C8	2.47	0.49
30:31:78:ILE:HA	30:31:83:PHE:CD2	2.46	0.49
36:68:71:ARG:NH1	41:B8:74:ARG:HH21	2.09	0.49
42:C8:27:LEU:HD12	42:C8:31:SER:HB3	1.93	0.49
1:1G:824:C:H2'	1:1G:825:G:C8	2.47	0.49
1:1G:993:G:H4'	1:1G:994:A:OP2	2.12	0.49
1:1G:1145:C:H4'	1:1G:1146:A:H5'	1.93	0.49
1:1G:1227:A:H4'	19:AA:85:LYS:HD2	1.93	0.49
2:12:42:ILE:HG13	2:12:43:ASP:N	2.26	0.49
6:52:55:ASP:HB2	6:52:86:ARG:HH12	1.77	0.49
11:2A:48:ILE:HG13	11:2A:64:ALA:HB2	1.93	0.49
25:14:137(A):G:H2'	25:14:139:G:N7	2.27	0.49
25:14:305:U:H2'	25:14:306:U:C6	2.46	0.49
25:14:588:U:H1'	30:39:90:PHE:HB3	1.94	0.49
25:14:1063:G:H2'	25:14:1063:G:N3	2.28	0.49
25:14:2681:C:H4'	25:14:2682:U:H5'	1.94	0.49
25:14:2707:G:H5'	39:55:68:ARG:HH21	1.77	0.49
26:1J:13:A:O2'	26:1J:15:A:O5'	2.31	0.49
41:75:55:ASN:HD22	41:75:58:ASN:HD21	1.59	0.49
43:95:35:LEU:O	43:95:37:VAL:HG22	2.12	0.49
51:H5:40:THR:HG23	51:H5:43:ILE:H	1.77	0.49
3:2E:89:GLU:HG3	3:2E:90:GLU:N	2.25	0.49
10:1I:34:VAL:HG22	10:1I:74:ILE:HG22	1.95	0.49
25:1H:86:C:H4'	25:1H:104:U:H1'	1.93	0.49
25:1H:270(G):C:H2'	25:1H:270(H):C:C6	2.47	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:796:C:H2'	25:1H:797:C:C6	2.47	0.49
25:1H:958:U:O2	26:16:89(A):A:H4'	2.13	0.49
25:1H:1799:G:H4'	25:1H:1800:C:O5'	2.12	0.49
25:1H:2712:U:O2'	25:1H:2712(A):A:C8	2.62	0.49
31:41:99:MET:HG3	31:41:100:TRP:N	2.26	0.49
32:51:124:GLU:HG2	32:51:126:PRO:HD3	1.94	0.49
34:38:50:ARG:HB2	34:38:50:ARG:NH1	2.26	0.49
41:B8:74:ARG:HD3	41:B8:76:PHE:CE1	2.48	0.49
43:D8:5:VAL:O	43:D8:12:TYR:N	2.44	0.49
49:J8:18:ILE:HG12	49:J8:37:ILE:HG12	1.94	0.49
52:M8:23:GLU:O	52:M8:25:TYR:N	2.43	0.49
1:1G:668:G:H4'	15:6A:48:LYS:HB2	1.92	0.49
1:1G:975:A:H4'	1:1G:976:G:O5'	2.13	0.49
1:1G:1006:C:N3	1:1G:1023:G:N2	2.59	0.49
1:1G:1109:C:H2'	1:1G:1110:A:O4'	2.12	0.49
1:1G:1223:C:OP2	19:AA:78:ARG:NH2	2.44	0.49
1:1G:1227:A:N3	19:AA:84:GLY:HA3	2.28	0.49
1:1G:1443:G:C5	41:75:118:ARG:HB2	2.47	0.49
4:32:158:ILE:HG12	4:32:162:LEU:HD12	1.93	0.49
25:14:322:A:H5'	25:14:340:A:H1'	1.94	0.49
25:14:1105:U:H2'	25:14:1106:G:H8	1.77	0.49
25:14:1191:G:H2'	25:14:1192:G:O4'	2.12	0.49
25:14:1467:C:C5	25:14:1546:C:H2'	2.47	0.49
29:29:5:LEU:HD21	29:29:79:ARG:HB2	1.94	0.49
37:35:96:THR:HG22	37:35:126:VAL:HB	1.93	0.49
46:C5:43:ASN:OD1	46:C5:65:ALA:HB3	2.12	0.49
1:13:489:C:OP1	4:3E:132:ARG:NH2	2.44	0.49
10:1I:61:GLU:HG3	14:5I:58:LYS:HE2	1.92	0.49
25:1H:177:G:N3	25:1H:177:G:H5''	2.27	0.49
25:1H:666:G:H4'	37:78:49:ARG:NH1	2.27	0.49
25:1H:1952:A:OP1	36:68:44:LYS:NZ	2.27	0.49
25:1H:2611:U:H2'	53:N8:2:ALA:O	2.11	0.49
25:1H:2776:A:H4'	25:1H:2777:G:O5'	2.12	0.49
27:71:193:ILE:HG22	27:71:194:ARG:HG3	1.94	0.49
28:11:145:VAL:HG12	28:11:146:GLU:O	2.12	0.49
31:41:54:GLU:O	31:41:58:GLN:N	2.42	0.49
33:61:77:LEU:HD11	33:61:140:LEU:HD12	1.94	0.49
36:68:107:ARG:NH1	41:B8:36:GLU:HG2	2.28	0.49
45:F8:31:HIS:CE1	45:F8:33:LYS:HB2	2.47	0.49
1:1G:266:G:H2'	1:1G:266:G:N3	2.27	0.49
1:1G:390:C:H2'	1:1G:391:G:H8	1.78	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:552:U:H4'	12:3A:83:ARG:HG2	1.95	0.49
1:1G:613:C:H2'	1:1G:614:A:O4'	2.12	0.49
1:1G:929:G:H1	1:1G:1388:C:N4	2.10	0.49
1:1G:1300:G:O2'	1:1G:1301:U:O5'	2.28	0.49
1:1G:1378:C:H5'	7:62:94:ARG:NH2	2.27	0.49
2:12:8:LYS:HD3	2:12:217:ARG:NH1	2.28	0.49
2:12:23:ARG:NH1	2:12:23:ARG:O	2.45	0.49
3:22:22:TRP:HZ2	3:22:33:LEU:HD13	1.76	0.49
8:72:104:ARG:HB3	8:72:108:GLY:N	2.27	0.49
25:14:251:A:C5	25:14:252:G:H1'	2.47	0.49
25:14:1428:C:N4	25:14:1570:A:OP2	2.43	0.49
25:14:1678:G:H22	25:14:1989:G:H22	1.61	0.49
25:14:1784:A:OP1	62:14:3555:HOH:O	2.20	0.49
25:14:2680:C:H5'	29:29:189:PRO:HA	1.94	0.49
25:14:2849:U:H4'	25:14:2868:A:C2	2.47	0.49
26:1J:78:A:H2'	26:1J:79:C:O4'	2.12	0.49
44:A5:74:ALA:HB2	44:A5:105:VAL:HG13	1.93	0.49
1:13:965:A:O2'	1:13:966:M2G:OP2	2.21	0.49
1:13:1128:C:H4'	9:8E:16:ARG:HH12	1.78	0.49
1:13:1342:C:H4'	9:8E:125:TYR:HB3	1.95	0.49
3:2E:14:ILE:HG12	3:2E:15:THR:N	2.28	0.49
8:7E:32:LYS:O	8:7E:36:LEU:HG	2.12	0.49
14:5I:48:ALA:HB2	14:5I:53:LEU:HD12	1.94	0.49
23:2K:13:C:O2'	25:1H:1924:C:H4'	2.12	0.49
25:1H:658:C:H2'	25:1H:659:C:C6	2.47	0.49
25:1H:1368:G:OP1	55:P8:28:ARG:NH2	2.44	0.49
28:11:95:LEU:HD13	28:11:97:TYR:HE1	1.77	0.49
30:31:116:ASP:OD1	30:31:119:ARG:NH2	2.45	0.49
40:A8:43:GLU:HG3	48:I8:49:LYS:HE2	1.93	0.49
54:O8:12:GLU:OE1	54:O8:12:GLU:N	2.46	0.49
1:1G:1155:G:H2'	1:1G:1156:G:H5'	1.94	0.49
1:1G:1194:U:H2'	1:1G:1195:C:C6	2.47	0.49
1:1G:1323:G:O5'	13:4A:99:ARG:NH2	2.46	0.49
8:72:30:ARG:O	8:72:34:GLU:HG2	2.11	0.49
9:82:40:LEU:HD11	9:82:70:LYS:HB3	1.93	0.49
15:6A:55:GLY:HA2	15:6A:58:MET:HG3	1.94	0.49
19:AA:16:LEU:HA	19:AA:19:VAL:HB	1.94	0.49
22:3L:31:G:H2'	22:3L:32:G:C8	2.48	0.49
25:14:65:C:H2'	25:14:66:C:C6	2.47	0.49
25:14:265:A:C2'	25:14:266:G:H4'	2.42	0.49
25:14:935:C:H2'	25:14:936:C:C6	2.47	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:1492:G:H3'	25:14:1493:C:H5'	1.95	0.49
25:14:2031:A:N3	25:14:2455:G:O2'	2.43	0.49
25:14:2836:U:H2'	25:14:2837:G:C8	2.46	0.49
26:1J:7:G:H3'	26:1J:8:U:H5''	1.95	0.49
26:1J:18:G:H2'	26:1J:19:G:C8	2.47	0.49
48:E5:36:ILE:HG13	48:E5:58:THR:HG23	1.93	0.49
52:I5:6:HIS:HB2	52:I5:7:PRO:HA	1.94	0.49
52:I5:67:TYR:CD2	52:I5:70:GLY:HA3	2.47	0.49
1:13:128:G:O2'	17:8I:3:LYS:NZ	2.30	0.49
1:13:559:A:OP1	5:4E:126:ARG:NH1	2.37	0.49
1:13:749:C:H2'	1:13:750:G:H8	1.76	0.49
1:13:833:U:H2'	1:13:834:C:C6	2.48	0.49
2:1E:209:ARG:HH11	2:1E:236:TYR:HE1	1.59	0.49
3:2E:10:PHE:HD2	3:2E:11:ARG:HD3	1.77	0.49
19:AI:36:ARG:NH1	19:AI:52:TYR:O	2.44	0.49
22:3K:8:U:O4'	22:3K:49:C:O2'	2.31	0.49
25:1H:1052:C:H42	25:1H:1107:G:H1	1.61	0.49
25:1H:1210:A:H4'	25:1H:1211:U:O5'	2.12	0.49
25:1H:1788:C:OP1	28:11:222:ARG:NH2	2.43	0.49
25:1H:2747:G:O6	25:1H:2755:C:H5''	2.12	0.49
32:51:115:VAL:HG11	32:51:148:ILE:HD11	1.95	0.49
38:88:32:TYR:CD1	38:88:133:ARG:HA	2.47	0.49
41:B8:41:ARG:HE	41:B8:43:GLN:HB2	1.77	0.49
42:C8:114:LYS:HA	42:C8:118:GLY:HA3	1.94	0.49
47:H8:5:LEU:HD23	47:H8:47:VAL:HG21	1.94	0.49
53:N8:30:LEU:HA	53:N8:41:PRO:HA	1.94	0.49
55:P8:24:THR:O	55:P8:28:ARG:HG3	2.12	0.49
1:1G:79:G:H2'	1:1G:79:G:N3	2.28	0.49
1:1G:539:A:H2'	1:1G:540:G:H8	1.78	0.49
1:1G:860:A:H2'	1:1G:861:G:O4'	2.13	0.49
1:1G:1227:A:C2	19:AA:84:GLY:HA3	2.46	0.49
11:2A:33:THR:HG22	11:2A:39:PRO:HA	1.92	0.49
19:AA:10:PHE:O	19:AA:39:THR:OG1	2.31	0.49
19:AA:33:THR:HG23	19:AA:51:VAL:HA	1.95	0.49
57:2L:26:C:H2'	57:2L:27:G:O4'	2.12	0.49
25:14:587:C:H4'	25:14:588:U:O5'	2.12	0.49
25:14:903:C:H2'	25:14:904:C:C6	2.47	0.49
25:14:1790:C:H5''	25:14:1791:A:OP1	2.12	0.49
25:14:1821:A:H2'	25:14:1822:G:H8	1.78	0.49
25:14:2050:C:H1'	29:29:156:MET:HE2	1.95	0.49
25:14:2867:G:O2'	25:14:2868:A:P	2.70	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:39:8:GLN:HG3	30:39:126:VAL:HA	1.95	0.49
30:39:61:GLY:CA	30:39:77:ASP:HB3	2.39	0.49
41:75:37:GLY:C	41:75:39:ARG:H	2.21	0.49
45:B5:6:ASP:OD1	45:B5:6:ASP:N	2.45	0.49
48:E5:32:ARG:N	48:E5:35:ASN:OD1	2.31	0.49
1:13:309:G:H2'	1:13:310:G:H8	1.77	0.49
1:13:445:G:H1	1:13:489:C:H42	1.59	0.49
1:13:1152:A:H5''	10:1I:13:HIS:NE2	2.28	0.49
1:13:1188:A:H5''	14:5I:58:LYS:NZ	2.27	0.49
3:2E:189:ALA:HB3	3:2E:196:LEU:HB2	1.95	0.49
5:4E:11:ILE:HG13	5:4E:31:LEU:HB3	1.95	0.49
16:7I:4:ILE:O	16:7I:66:PRO:HA	2.13	0.49
20:BI:64:ASP:OD2	20:BI:81:LYS:NZ	2.42	0.49
25:1H:95:G:H4'	50:K8:46:GLN:HB3	1.95	0.49
25:1H:248:G:H5'	25:1H:250:G:N7	2.28	0.49
25:1H:719:C:H2'	25:1H:720:C:H6	1.77	0.49
30:31:63:LYS:NZ	30:31:67:GLN:HB3	2.27	0.49
37:78:50:ARG:HD3	56:Q8:7:HIS:CD2	2.47	0.49
43:D8:10:LYS:NZ	43:D8:23:GLU:OE1	2.44	0.49
1:1G:278:G:N2	17:8A:95:TYR:HB3	2.27	0.49
1:1G:500:G:H2'	1:1G:501:C:C6	2.47	0.49
1:1G:1117:G:O3'	9:82:104:ARG:HD3	2.13	0.49
1:1G:1320:C:C5	19:AA:36:ARG:HD3	2.48	0.49
1:1G:1418:A:N6	1:1G:1482:G:O2'	2.45	0.49
6:52:20:ALA:HA	6:52:23:LYS:HB3	1.94	0.49
7:62:57:GLU:OE1	7:62:57:GLU:N	2.42	0.49
25:14:678:C:H2'	25:14:679:C:C6	2.48	0.49
25:14:996:A:O2'	42:85:92:ARG:NH2	2.46	0.49
25:14:1027:A:C2	25:14:2488:A:H5'	2.47	0.49
25:14:1314:C:OP1	25:14:1332:G:H5''	2.13	0.49
25:14:1358:G:N1	25:14:1372:U:OP2	2.25	0.49
25:14:1827:C:OP2	28:19:222:ARG:NH1	2.46	0.49
25:14:2718:G:H5'	62:14:3517:HOH:O	2.11	0.49
25:14:2720:U:N3	25:14:2873:A:H2	1.94	0.49
28:19:79:VAL:HG12	28:19:113:VAL:HA	1.95	0.49
1:13:495:A:H4'	1:13:496:A:OP1	2.11	0.49
1:13:939:G:H2'	1:13:940:C:C6	2.48	0.49
7:6E:22:LEU:HG	7:6E:62:PHE:HE1	1.78	0.49
15:6I:78:TYR:OH	15:6I:88:ARG:NE	2.46	0.49
19:AI:80:TYR:CG	19:AI:80:TYR:O	2.66	0.49
25:1H:458:G:O2'	25:1H:469:G:O6	2.22	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:519:U:H2'	25:1H:520:G:C8	2.47	0.49
25:1H:1417:C:H2'	25:1H:1418:G:O4'	2.12	0.49
27:71:10:LEU:HB3	27:71:220:PRO:HD3	1.95	0.49
28:11:226:MET:O	62:11:401:HOH:O	2.20	0.49
31:41:63:ILE:HG13	31:41:141:PHE:CG	2.47	0.49
45:F8:55:ASN:HB2	45:F8:80:ILE:HG12	1.94	0.49
1:1G:552:U:O2'	12:3A:83:ARG:O	2.26	0.49
1:1G:1067:A:O2'	1:1G:1068:G:OP2	2.30	0.49
2:12:179:LYS:HA	8:72:72:PRO:HG3	1.93	0.49
3:22:18:TRP:HH2	14:5A:57:ARG:HD2	1.78	0.49
13:4A:25:ILE:O	13:4A:29:ARG:HB2	2.12	0.49
18:9A:22:VAL:HG22	18:9A:23:LYS:H	1.78	0.49
22:3L:6:G:H1	22:3L:69:C:H42	1.61	0.49
25:14:71:A:H4'	25:14:72:U:O5'	2.13	0.49
25:14:189:G:O6	25:14:205:G:O2'	2.26	0.49
25:14:1222:C:H42	25:14:1229:G:H1	1.60	0.49
25:14:1278:A:H2'	25:14:1279:G:C8	2.48	0.49
25:14:1306:C:H2'	25:14:1307:A:C8	2.48	0.49
25:14:1364:G:OP2	49:F5:2:SER:N	2.46	0.49
25:14:2099:U:H3	25:14:2190:G:H1	1.60	0.49
25:14:2287:A:O2'	25:14:2288:A:H5''	2.13	0.49
35:15:112:LEU:HD23	35:15:113:GLY:H	1.78	0.49
37:35:54:GLY:C	37:35:56:SER:H	2.21	0.49
37:35:144:GLU:OE1	37:35:144:GLU:N	2.46	0.49
45:B5:64:LYS:HE2	45:B5:73:ARG:CZ	2.43	0.49
49:F5:90:ILE:HG23	49:F5:90:ILE:O	2.12	0.49
1:13:148:G:H2'	1:13:149:A:C8	2.47	0.49
7:6E:49:ILE:O	7:6E:53:LYS:HB3	2.12	0.49
25:1H:1677:A:N7	62:1H:3789:HOH:O	2.35	0.49
25:1H:2612:C:OP2	53:N8:2:ALA:HA	2.12	0.49
31:41:106:LEU:HD12	31:41:110:ALA:HB3	1.93	0.49
35:58:96:GLU:HB2	35:58:122:VAL:HG12	1.95	0.49
40:A8:42:ASP:C	40:A8:44:LYS:H	2.21	0.49
42:C8:91:ASP:O	42:C8:93:LYS:N	2.46	0.49
1:1G:545:C:OP2	4:32:62:GLN:NE2	2.46	0.49
25:14:307:G:N1	25:14:310:A:OP2	2.46	0.49
25:14:455:C:N3	25:14:472:A:H2'	2.28	0.49
30:39:60:SER:OG	30:39:60:SER:O	2.31	0.49
31:49:59:GLU:OE1	31:49:138:GLN:NE2	2.45	0.49
38:45:135:ASP:OD2	47:D5:81:ARG:NH1	2.45	0.49
47:D5:111:VAL:O	47:D5:113:ALA:N	2.46	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:921:U:O2'	5:4E:19:MET:O	2.29	0.49
2:1E:67:THR:HG21	2:1E:155:LEU:HG	1.95	0.49
11:2I:83:ILE:HG12	11:2I:109:VAL:HG23	1.94	0.49
25:1H:221:A:H5''	25:1H:221:A:H8	1.78	0.49
25:1H:1257:C:H4'	30:31:83:PHE:CE1	2.48	0.49
27:71:201:PRO:HD2	27:71:208:PHE:HE2	1.76	0.49
28:11:123:ALA:CB	28:11:131:LEU:HG	2.43	0.49
33:61:67:ARG:HE	33:61:67:ARG:HB3	1.41	0.49
38:88:64:ILE:HA	38:88:106:VAL:HG12	1.95	0.49
41:B8:46:GLU:O	41:B8:46:GLU:HG2	2.12	0.49
48:I8:36:ILE:HA	48:I8:60:PHE:HA	1.94	0.49
1:1G:543:C:H5''	4:32:14:ARG:HD2	1.95	0.49
1:1G:575:G:O4'	1:1G:881:G:N2	2.46	0.49
1:1G:1491:G:H5''	1:1G:1492:A:OP2	2.12	0.49
3:22:123:GLN:HA	3:22:126:ARG:HD2	1.95	0.49
19:AA:36:ARG:HG2	19:AA:36:ARG:O	2.12	0.49
25:14:224:G:O6	25:14:419:C:O2'	2.31	0.49
25:14:1359:A:N6	25:14:1372:U:H3	2.11	0.49
25:14:1374:G:H2'	25:14:1375:C:C6	2.48	0.49
28:19:10:THR:OG1	28:19:11:PRO:O	2.30	0.49
29:29:36:ARG:HG2	29:29:47:VAL:HG12	1.95	0.49
29:29:77:ILE:HA	29:29:77:ILE:HD13	1.70	0.49
36:25:119:PRO:HB2	41:75:68:TYR:CZ	2.48	0.49
1:13:842:C:H5'	1:13:843:U:OP1	2.13	0.48
1:13:1119:C:H2'	1:13:1120:G:C8	2.47	0.48
1:13:1431:C:H2'	1:13:1432:G:O4'	2.13	0.48
2:1E:105:PHE:HZ	2:1E:156:LYS:HA	1.77	0.48
25:1H:57:C:H2'	25:1H:58:G:O4'	2.13	0.48
25:1H:64:A:C8	45:F8:66:LEU:HD13	2.48	0.48
25:1H:247:G:H4'	25:1H:386:G:C5	2.48	0.48
25:1H:2182:G:H2'	25:1H:2183:C:C6	2.48	0.48
25:1H:2712:U:O2'	25:1H:2712(A):A:OP1	2.31	0.48
25:1H:2734:A:H3'	25:1H:2735:G:H8	1.77	0.48
31:41:119:GLY:HA3	31:41:181:ARG:HG3	1.94	0.48
32:51:127:GLU:O	32:51:129:THR:N	2.46	0.48
40:A8:61:ASN:OD1	40:A8:63:THR:OG1	2.23	0.48
43:D8:66:ARG:HD2	43:D8:88:ARG:HB2	1.94	0.48
49:J8:87:PRO:O	49:J8:91:LYS:HB2	2.13	0.48
1:1G:15:G:H1'	5:42:19:MET:HE2	1.94	0.48
1:1G:1319:A:H2'	1:1G:1323:G:C8	2.48	0.48
3:22:150:LYS:HG3	3:22:169:ALA:HB2	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:247:G:H4'	25:14:386:G:C5	2.48	0.48
25:14:247:G:H4'	25:14:386:G:C6	2.48	0.48
25:14:663:G:H2'	25:14:664:C:C6	2.48	0.48
25:14:1322:A:N1	25:14:1333:C:O2'	2.38	0.48
25:14:2291:U:O2'	25:14:2374:C:O2	2.28	0.48
25:14:2747:G:O6	25:14:2755:C:H5''	2.13	0.48
31:49:38:VAL:HG23	31:49:93:THR:HA	1.95	0.48
35:15:55:VAL:HB	35:15:126:PRO:HA	1.95	0.48
36:25:98:VAL:HG21	36:25:114:ILE:HG22	1.95	0.48
40:65:84:GLN:HA	40:65:109:GLY:HA3	1.95	0.48
44:A5:36:LEU:HD11	44:A5:47:VAL:HG12	1.95	0.48
46:C5:55:TYR:HE2	46:C5:60:PHE:CE1	2.27	0.48
1:13:600:C:H2'	1:13:601:C:H6	1.77	0.48
1:13:1037:C:H2'	1:13:1038:C:H6	1.78	0.48
1:13:1132:C:H2'	1:13:1133:G:C8	2.47	0.48
5:4E:140:ARG:O	5:4E:143:ARG:NH1	2.43	0.48
8:7E:120:THR:H	8:7E:123:GLU:HB2	1.77	0.48
11:2I:95:ILE:O	11:2I:99:GLN:HG3	2.12	0.48
25:1H:1291:C:H2'	25:1H:1292:U:C6	2.49	0.48
26:16:15:A:H1'	26:16:109:G:C5	2.47	0.48
35:58:43:THR:HB	35:58:46:VAL:HG12	1.95	0.48
41:B8:24:PRO:HA	41:B8:49:VAL:HG13	1.94	0.48
41:B8:124:ASP:N	41:B8:127:ALA:HB3	2.26	0.48
50:K8:29:LYS:HG2	50:K8:57:ILE:HD13	1.95	0.48
1:1G:8:A:H4'	1:1G:9:G:OP1	2.13	0.48
1:1G:1266:G:N2	1:1G:1269:A:OP2	2.45	0.48
2:12:103:THR:OG1	2:12:104:ASN:N	2.45	0.48
25:14:1259:G:H2'	25:14:1260:G:C8	2.47	0.48
26:1J:7:G:N2	40:65:38:GLN:OE1	2.31	0.48
47:D5:103:ARG:O	47:D5:139:VAL:HG11	2.12	0.48
1:13:643:C:H2'	1:13:644:G:H8	1.77	0.48
17:8I:90:ILE:O	17:8I:94:ASN:ND2	2.45	0.48
22:1K:10:G:H2'	22:1K:11:A:C8	2.47	0.48
25:1H:7:G:H1	25:1H:2896:C:H42	1.61	0.48
25:1H:195:A:C8	25:1H:197:A:OP1	2.67	0.48
25:1H:319:C:OP1	30:31:137:LYS:NZ	2.43	0.48
25:1H:583:G:O6	62:1H:3688:HOH:O	2.20	0.48
25:1H:862:G:H2'	25:1H:863:A:O4'	2.14	0.48
25:1H:1678:G:N2	25:1H:1989:G:H22	2.11	0.48
25:1H:2348:U:O4	25:1H:2382:G:N2	2.46	0.48
31:41:10:LYS:HG2	31:41:14:GLU:HG2	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:51:19:VAL:HA	32:51:24:VAL:HG12	1.95	0.48
33:61:110:ASP:N	33:61:130:TYR:OH	2.35	0.48
34:38:25:PHE:CD1	34:38:26:LEU:HG	2.47	0.48
35:58:132:ALA:O	35:58:134:ARG:NH1	2.46	0.48
37:78:18:ARG:NH2	37:78:21:ARG:HD2	2.29	0.48
43:D8:29:PRO:HA	43:D8:61:VAL:HG22	1.95	0.48
44:E8:19:LEU:HB3	53:N8:25:LEU:HD12	1.95	0.48
47:H8:151:HIS:CD2	47:H8:170:THR:HA	2.46	0.48
1:1G:232:G:H2'	1:1G:233:C:O4'	2.13	0.48
1:1G:302:G:H5''	12:3A:14:LYS:HE2	1.95	0.48
1:1G:353:A:H5'	1:1G:353:A:H8	1.77	0.48
1:1G:501:C:H2'	1:1G:502:G:H8	1.76	0.48
1:1G:554:C:H2'	1:1G:555:C:C6	2.48	0.48
1:1G:784:C:H4'	25:14:1837:C:OP1	2.12	0.48
1:1G:1226:C:H42	13:4A:104:ARG:HD2	1.78	0.48
1:1G:1352:C:N3	1:1G:1370:G:N2	2.53	0.48
25:14:2061:G:H5''	25:14:2503:2MA:N7	2.28	0.48
25:14:2392:A:H8	37:35:61:ARG:HB3	1.77	0.48
25:14:2503:2MA:H5'	25:14:2503:2MA:H8	1.94	0.48
28:19:77:ALA:HB2	28:19:97:TYR:CD1	2.48	0.48
46:C5:90:LEU:HD12	46:C5:91:GLU:H	1.79	0.48
1:13:148:G:H2'	1:13:149:A:H8	1.78	0.48
1:13:156:G:H2'	1:13:157:G:C8	2.49	0.48
1:13:591:U:H2'	1:13:592:G:C8	2.48	0.48
1:13:701:C:O2	1:13:703:G:N1	2.46	0.48
1:13:1498:UR3:H6	1:13:1498:UR3:O5'	2.12	0.48
23:2K:76:C:N3	25:1H:2251:OMG:N2	2.45	0.48
25:1H:2057:A:H2'	25:1H:2058:A:O4'	2.12	0.48
28:11:65:ILE:HD11	28:11:67:PHE:CE1	2.48	0.48
29:21:111:ARG:HG3	29:21:160:TYR:HD2	1.78	0.48
31:41:10:LYS:O	31:41:14:GLU:HB3	2.14	0.48
41:B8:11:GLU:OE1	41:B8:57:PHE:HB3	2.13	0.48
50:K8:25:VAL:HG23	50:K8:57:ILE:HG23	1.95	0.48
53:N8:46:CYS:H	53:N8:50:GLY:HA3	1.78	0.48
1:1G:145:G:H1	1:1G:177:C:N4	2.11	0.48
1:1G:189:U:O2'	17:8A:63:ARG:NH2	2.46	0.48
1:1G:408:A:H2'	1:1G:409:G:O4'	2.13	0.48
8:72:7:ALA:O	8:72:11:THR:OG1	2.22	0.48
25:14:17:G:H2'	25:14:18:C:C6	2.48	0.48
25:14:184:C:H2'	25:14:185:U:C6	2.49	0.48
25:14:2505:G:O6	25:14:2576:G:H2'	2.14	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:59:8:PRO:HD2	32:59:69:ARG:HE	1.78	0.48
33:69:88:ILE:HG22	33:69:90:GLY:N	2.26	0.48
33:69:109:ILE:HG22	33:69:110:ASP:H	1.78	0.48
41:75:18:ASP:OD1	41:75:18:ASP:N	2.47	0.48
48:E5:27:GLU:HB2	48:E5:69:PHE:HD2	1.78	0.48
54:K5:12:GLU:OE1	54:K5:12:GLU:N	2.45	0.48
1:13:581:G:O3'	15:6I:64:ARG:NH2	2.47	0.48
1:13:789:U:H1'	1:13:792:A:H2	1.79	0.48
14:5I:39:LEU:HD11	14:5I:47:LEU:HD12	1.95	0.48
18:9I:43:PHE:CE1	18:9I:58:LEU:HD11	2.49	0.48
25:1H:102:G:OP1	50:K8:7:ARG:NH2	2.42	0.48
25:1H:708:C:H42	25:1H:723:G:H1	1.60	0.48
31:41:16:ARG:NH2	31:41:31:VAL:HG11	2.28	0.48
31:41:152:LEU:H	31:41:152:LEU:HD12	1.79	0.48
54:O8:34:LEU:H	54:O8:34:LEU:HD22	1.78	0.48
1:1G:405:U:O4	4:32:2:GLY:N	2.46	0.48
1:1G:1255:G:O2'	1:1G:1258:G:O2'	2.14	0.48
2:12:19:HIS:NE2	2:12:20:GLU:HG2	2.28	0.48
5:42:28:PHE:O	5:42:47:LYS:HA	2.14	0.48
20:BA:74:LYS:HA	20:BA:77:ALA:HB3	1.93	0.48
25:14:1075:C:H2'	25:14:1076:C:C6	2.48	0.48
25:14:1306:C:H2'	25:14:1307:A:H8	1.79	0.48
25:14:1846:G:H5'	25:14:1847:A:OP2	2.14	0.48
25:14:2059:A:OP2	62:14:3553:HOH:O	2.20	0.48
25:14:2335:A:HO2'	25:14:2336:A:P	2.36	0.48
26:1J:16:G:H1	26:1J:68:C:N4	2.11	0.48
37:35:130:PHE:CZ	37:35:144:GLU:HB2	2.48	0.48
45:B5:11:PRO:HA	45:B5:28:PHE:HB3	1.96	0.48
47:D5:74:VAL:HG13	47:D5:86:VAL:HG22	1.96	0.48
1:13:346:G:H5''	41:B8:41:ARG:HD2	1.96	0.48
1:13:596:C:H2'	1:13:597:G:C8	2.47	0.48
4:3E:86:LYS:H	4:3E:86:LYS:HD2	1.77	0.48
12:3I:51:LYS:HG2	12:3I:72:HIS:CD2	2.49	0.48
13:4I:5:ALA:HB1	13:4I:66:LEU:HD13	1.94	0.48
22:1K:40:C:H2'	22:1K:41:C:C6	2.48	0.48
23:2K:21:H2U:H2'	23:2K:22:A:C5'	2.43	0.48
25:1H:49:A:N7	25:1H:120:U:C5	2.80	0.48
25:1H:1174:A:H2'	25:1H:1174:A:N3	2.29	0.48
25:1H:2125:G:H21	25:1H:2173:A:H62	1.61	0.48
25:1H:2347:C:C5	25:1H:2382:G:H1'	2.49	0.48
25:1H:2528:U:O3'	25:1H:2529:G:H8	1.97	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:2680:C:H1'	29:21:187:ALA:HB1	1.96	0.48
25:1H:2699:C:H2'	25:1H:2700:C:O4'	2.14	0.48
47:H8:162:GLU:HG3	47:H8:163:LEU:N	2.27	0.48
1:1G:452:A:O2'	1:1G:453:A:O5'	2.27	0.48
1:1G:988:G:H2'	1:1G:989:C:O4'	2.14	0.48
1:1G:1179:A:H2'	1:1G:1180:A:O4'	2.14	0.48
4:32:12:CYS:SG	4:32:18:LYS:HA	2.52	0.48
4:32:18:LYS:HD2	4:32:31:CYS:SG	2.53	0.48
9:82:28:VAL:HG23	9:82:63:ILE:HD13	1.96	0.48
11:2A:13:GLN:NE2	11:2A:15:ALA:HB2	2.27	0.48
25:14:1045:A:OP1	25:14:1046:A:H5'	2.13	0.48
25:14:2630:G:H1'	25:14:2894:G:N9	2.29	0.48
30:39:53:THR:HG22	30:39:56:GLU:HG3	1.94	0.48
32:59:10:PRO:HD2	32:59:50:VAL:H	1.78	0.48
51:H5:5:LYS:HD3	51:H5:57:GLU:HB3	1.95	0.48
56:M5:29:LYS:HB3	56:M5:33:ASN:OD1	2.13	0.48
1:13:456:C:H2'	1:13:457:C:O4'	2.14	0.48
1:13:713:G:H2'	1:13:714:G:C8	2.49	0.48
25:1H:270(I):G:N2	25:1H:270(R):G:N3	2.62	0.48
25:1H:764:A:H2	28:11:219:PRO:HG3	1.78	0.48
25:1H:1021:A:H8	25:1H:1021:A:H3'	1.77	0.48
25:1H:2693:A:H2'	25:1H:2694:G:H8	1.78	0.48
44:E8:82:LEU:HB2	44:E8:98:LYS:O	2.14	0.48
1:1G:664:G:P	18:9A:64:ARG:HH21	2.36	0.48
1:1G:707:C:H2'	1:1G:708:C:C6	2.49	0.48
1:1G:1350:A:N6	1:1G:1373:G:H21	2.11	0.48
7:62:5:ARG:HH21	7:62:7:ALA:HA	1.77	0.48
8:72:29:SER:HB3	8:72:32:LYS:HG3	1.95	0.48
20:BA:36:LEU:HD13	20:BA:55:ILE:HG23	1.95	0.48
22:3L:29:C:H2'	22:3L:30:G:C8	2.43	0.48
25:14:29:U:H2'	25:14:30:G:C8	2.49	0.48
25:14:41:C:H2'	25:14:43:G:H8	1.78	0.48
25:14:906:G:O2'	38:45:67:ARG:NE	2.31	0.48
25:14:1049:C:N4	25:14:2751:G:H1	2.10	0.48
25:14:1344:G:H4'	25:14:1384:A:C5	2.48	0.48
25:14:2779:U:H5'	25:14:2780:G:OP1	2.14	0.48
25:14:2837:G:H22	39:55:96:ARG:HH12	1.61	0.48
36:25:114:ILE:O	36:25:117:LEU:N	2.47	0.48
39:55:2:ARG:HB3	39:55:2:ARG:NH1	2.29	0.48
1:13:74:C:H2'	1:13:75:C:C6	2.49	0.48
1:13:337:C:H2'	1:13:338:A:C8	2.48	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1E:97:TRP:CZ3	2:1E:172:ILE:HB	2.49	0.48
11:2I:57:THR:HG22	11:2I:58:PRO:HD2	1.95	0.48
25:1H:1113:U:H2'	25:1H:1114:G:C8	2.49	0.48
25:1H:2105:C:H42	25:1H:2184:G:H1	1.62	0.48
26:16:90:C:H5'	38:88:18:LYS:HA	1.95	0.48
30:31:123:LEU:HD13	30:31:192:LEU:HB3	1.96	0.48
52:M8:3:GLU:HG3	52:M8:4:GLY:H	1.79	0.48
1:1G:740:U:H2'	1:1G:741:G:H8	1.78	0.48
1:1G:932:C:H4'	7:62:4:ARG:CZ	2.44	0.48
1:1G:1537:U:O4	24:4L:8:A:N6	2.47	0.48
10:1A:80:LYS:O	10:1A:83:GLU:HB3	2.13	0.48
17:8A:29:HIS:CG	17:8A:30:PRO:HD2	2.49	0.48
25:14:1203:G:O6	25:14:1204:A:N6	2.47	0.48
25:14:1946:U:H2'	25:14:1947:C:C6	2.48	0.48
25:14:2468:G:N2	25:14:2481:G:H2'	2.29	0.48
26:1J:113:C:H2'	26:1J:114:G:C8	2.49	0.48
28:19:172:TYR:HD1	28:19:186:HIS:HA	1.79	0.48
30:39:5:ALA:H	30:39:19:GLU:HB3	1.78	0.48
30:39:174:VAL:HG11	30:39:188:ARG:NH2	2.29	0.48
40:65:61:ASN:OD1	40:65:62:LYS:N	2.46	0.48
42:85:91:ASP:O	42:85:92:ARG:HB3	2.13	0.48
1:13:105:G:H2'	1:13:106:C:C6	2.49	0.48
1:13:895:G:H2'	1:13:896:C:C6	2.49	0.48
3:2E:50:ALA:HB1	3:2E:70:VAL:HG11	1.96	0.48
13:4I:14:ARG:HG3	13:4I:16:ASP:OD1	2.14	0.48
15:6I:82:ILE:O	15:6I:86:GLY:N	2.43	0.48
25:1H:466:A:N3	25:1H:683:C:H1'	2.29	0.48
25:1H:1332:G:H21	25:1H:1610:A:H8	1.58	0.48
30:31:67:GLN:HG3	30:31:67:GLN:O	2.13	0.48
47:H8:116:VAL:HB	47:H8:174:VAL:HG12	1.96	0.48
1:1G:142:G:H2'	1:1G:143:A:H8	1.79	0.48
1:1G:629:G:H2'	1:1G:630:G:O4'	2.13	0.48
1:1G:709:G:H2'	1:1G:710:G:H8	1.79	0.48
1:1G:719:C:O2	18:9A:50:ILE:HG12	2.14	0.48
1:1G:1074:G:O2'	1:1G:1101:A:N1	2.40	0.48
1:1G:1532:U:H4'	1:1G:1533:C:H5'	1.94	0.48
19:AA:66:MET:HE3	19:AA:66:MET:HB2	1.62	0.48
22:3L:53:G:N2	22:3L:54:G:N3	2.62	0.48
25:14:1166:C:H2'	25:14:1167:U:C6	2.49	0.48
25:14:2030:A:H4'	25:14:2031:A:C8	2.49	0.48
25:14:2478:A:H3'	25:14:2479:G:H8	1.78	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:2563:U:O2'	36:25:28:SER:HB3	2.14	0.48
25:14:2748:A:H2	25:14:2753:A:N6	2.11	0.48
47:D5:69:THR:HG22	47:D5:90:VAL:HG22	1.95	0.48
1:13:1330:U:H3'	1:13:1331:G:O4'	2.14	0.48
1:13:1350:A:H8	1:13:1350:A:O5'	1.97	0.48
3:2E:6:HIS:HB2	14:5I:49:HIS:ND1	2.28	0.48
5:4E:108:ALA:O	5:4E:112:LEU:HB2	2.14	0.48
24:4K:13:A:O2'	24:4K:14:A:H3'	2.14	0.48
25:1H:191:A:H2'	25:1H:192:C:C6	2.49	0.48
25:1H:860:U:C5	25:1H:917:A:H2	2.32	0.48
25:1H:2112:G:N2	25:1H:2113:U:H3	2.03	0.48
25:1H:2131:G:N3	25:1H:2158:A:N6	2.61	0.48
26:16:91:C:OP1	38:88:19:GLY:HA2	2.13	0.48
42:C8:110:VAL:O	42:C8:114:LYS:HG2	2.14	0.48
49:J8:92:LYS:O	49:J8:96:LYS:N	2.47	0.48
1:1G:559:A:OP1	5:42:126:ARG:NH2	2.46	0.48
1:1G:690:G:H22	11:2A:55:LYS:HE3	1.78	0.48
1:1G:817:C:O3'	1:1G:818:G:H3'	2.13	0.48
1:1G:861:G:O2'	1:1G:874:G:O2'	2.30	0.48
1:1G:1018:C:H2'	1:1G:1019:C:O4'	2.14	0.48
7:62:16:LEU:HD13	9:82:45:ALA:HB2	1.96	0.48
9:82:97:LYS:HB3	9:82:98:PRO:HD3	1.95	0.48
25:14:107:C:H2'	25:14:108:U:C6	2.48	0.48
25:14:288:C:H2'	25:14:289:A:C8	2.48	0.48
25:14:2735:G:H2'	25:14:2736:G:H8	1.79	0.48
28:19:155:LEU:HD23	28:19:177:LEU:HD22	1.96	0.48
39:55:33:ARG:HA	39:55:114:VAL:O	2.14	0.48
46:C5:55:TYR:CG	46:C5:56:PRO:CD	2.95	0.48
49:F5:4:VAL:HG11	49:F5:11:ARG:HH11	1.78	0.48
1:13:1513:A:H2'	1:13:1514:C:C6	2.49	0.47
3:2E:6:HIS:CD2	3:2E:7:PRO:HD2	2.49	0.47
4:3E:52:SER:H	4:3E:55:ALA:HB3	1.79	0.47
9:8E:5:TYR:CE1	9:8E:16:ARG:HG2	2.46	0.47
14:5I:34:TYR:O	14:5I:38:GLY:N	2.33	0.47
22:1K:58:A:C6	22:1K:59:A:H2	2.32	0.47
25:1H:715:G:H2'	25:1H:716:A:C8	2.49	0.47
28:11:123:ALA:HB3	28:11:131:LEU:HG	1.95	0.47
30:31:172:TRP:CE3	30:31:173:VAL:HG23	2.48	0.47
33:61:14:ASP:O	33:61:16:GLY:N	2.46	0.47
37:78:23:PRO:O	37:78:25:SER:N	2.46	0.47
45:F8:67:GLY:C	45:F8:68:ARG:HG3	2.39	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:99:C:H2'	1:1G:101:A:H8	1.79	0.47
1:1G:792:A:H1'	1:1G:794:A:N7	2.29	0.47
1:1G:1228:C:H2'	1:1G:1229:A:H8	1.78	0.47
1:1G:1399:C:H4'	1:1G:1400:5MC:H5''	1.96	0.47
3:22:166:GLU:HG3	3:22:167:TRP:H	1.78	0.47
9:82:4:TYR:HD2	9:82:19:LEU:CB	2.27	0.47
9:82:37:PHE:HB3	9:82:43:ALA:HB1	1.95	0.47
13:4A:58:GLU:O	13:4A:62:ASN:HB2	2.13	0.47
14:5A:27:CYS:SG	14:5A:29:ARG:HB2	2.54	0.47
22:3L:22:A:OP1	22:3L:22:A:H4'	2.13	0.47
25:14:806:C:OP2	37:35:41:ARG:HD2	2.14	0.47
25:14:1209:G:H21	25:14:1210:A:H62	1.62	0.47
25:14:2335:A:O2'	25:14:2336:A:H2'	2.14	0.47
38:45:54:MET:HE2	38:45:54:MET:HB3	1.67	0.47
43:95:60:GLU:H	43:95:60:GLU:HG2	1.41	0.47
47:D5:70:LEU:HD23	47:D5:70:LEU:HA	1.80	0.47
47:D5:145:GLU:HG2	47:D5:146:ILE:H	1.79	0.47
1:13:507:C:OP2	1:13:508:C:O2'	2.23	0.47
1:13:791:G:H2'	1:13:792:A:H5'	1.96	0.47
1:13:1367:C:OP2	9:8E:112:LYS:NZ	2.47	0.47
19:AI:13:ASP:HA	19:AI:16:LEU:HB3	1.95	0.47
25:1H:107:C:H2'	25:1H:108:U:C6	2.48	0.47
25:1H:132:G:H1	25:1H:147:U:H3	1.62	0.47
25:1H:528:A:C2	25:1H:2043:C:H4'	2.49	0.47
26:16:66:A:O2'	26:16:67:G:O5'	2.32	0.47
29:21:13:ARG:HA	29:21:22:PRO:HA	1.96	0.47
30:31:32:LEU:HD22	30:31:112:MET:HE2	1.96	0.47
32:51:6:ARG:HH21	32:51:7:LEU:HD11	1.80	0.47
34:38:14:LYS:HE2	34:38:14:LYS:HA	1.96	0.47
37:78:84:ASN:HA	37:78:115:LEU:O	2.14	0.47
41:B8:111:ARG:C	41:B8:113:LYS:H	2.22	0.47
1:1G:108:G:H5''	1:1G:109:A:C2	2.49	0.47
1:1G:541:G:H2'	1:1G:542:G:H8	1.78	0.47
1:1G:1137:C:H5'	1:1G:1138:G:C4	2.49	0.47
1:1G:1392:G:H21	1:1G:1502:A:H8	1.61	0.47
2:12:20:GLU:HG3	2:12:23:ARG:NH2	2.29	0.47
2:12:91:PRO:HG3	2:12:155:LEU:HD21	1.96	0.47
25:14:363(C):G:H2'	25:14:363(D):G:H8	1.78	0.47
25:14:563:G:OP2	62:14:3556:HOH:O	2.20	0.47
25:14:1181:C:H2'	25:14:1182:A:O4'	2.14	0.47
25:14:1429:G:H2'	25:14:1430:C:C6	2.49	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:19:133:LEU:HB2	28:19:187:GLY:HA2	1.96	0.47
36:25:13:ASN:HD21	36:25:96:THR:HG23	1.80	0.47
44:A5:4:LYS:HG3	44:A5:106:ILE:HG22	1.96	0.47
1:13:186:C:O2'	20:BI:85:MET:SD	2.68	0.47
1:13:266:G:O2'	1:13:267:C:OP2	2.33	0.47
1:13:335:C:O2'	1:13:1433:A:N3	2.45	0.47
3:2E:83:ARG:O	3:2E:87:LEU:HG	2.14	0.47
7:6E:113:GLU:HB2	7:6E:118:VAL:HG23	1.96	0.47
22:3K:16:C:H5''	22:3K:17:C:OP2	2.14	0.47
25:1H:207:A:H2'	25:1H:208:C:O4'	2.14	0.47
25:1H:974(A):C:OP1	62:1H:3690:HOH:O	2.20	0.47
25:1H:1061:U:H3'	25:1H:1062:G:H5''	1.96	0.47
25:1H:1239:G:H2'	25:1H:1240:U:O4'	2.15	0.47
25:1H:1424:G:H2'	25:1H:1425:G:O4'	2.15	0.47
25:1H:2095:C:H2'	25:1H:2096:U:O4'	2.15	0.47
25:1H:2593:U:H2'	25:1H:2594:C:C6	2.49	0.47
27:71:15:ASP:N	27:71:20:TYR:OH	2.40	0.47
44:E8:1:MET:HE3	44:E8:62:HIS:ND1	2.29	0.47
47:H8:5:LEU:HB3	47:H8:59:LEU:HD23	1.96	0.47
52:M8:42:PHE:CG	52:M8:43:TYR:N	2.82	0.47
1:1G:803:G:OP1	62:1G:1812:HOH:O	2.20	0.47
1:1G:980:C:H3'	1:1G:981:U:C6	2.48	0.47
2:12:29:ALA:C	2:12:31:TYR:H	2.22	0.47
3:22:60:ALA:HB3	3:22:63:ASN:HB2	1.95	0.47
57:2L:65:G:P	48:E5:11:ARG:HH12	2.38	0.47
25:14:1141:U:H1'	25:14:1142(A):A:C2	2.49	0.47
25:14:1819:A:OP1	28:19:161:THR:HG21	2.14	0.47
25:14:2854:G:H1	25:14:2863:C:H42	1.63	0.47
31:49:125:PHE:HA	31:49:131:TYR:HD1	1.79	0.47
32:59:138:LYS:HA	32:59:141:VAL:HG12	1.96	0.47
41:75:65:LYS:HD3	41:75:67:SER:HB2	1.96	0.47
46:C5:3:VAL:HG22	46:C5:5:MET:HE2	1.96	0.47
1:13:186(A):C:OP1	20:BI:86:ARG:NH1	2.48	0.47
1:13:356:A:H2'	1:13:357:G:H8	1.79	0.47
1:13:1036:G:H5'	1:13:1037:C:OP2	2.14	0.47
1:13:1406:U:H3'	1:13:1407:5MC:HM53	1.96	0.47
5:4E:96:PRO:HA	5:4E:117:ASP:OD2	2.14	0.47
9:8E:4:TYR:HE1	9:8E:87:GLN:HB2	1.78	0.47
25:1H:38:A:H2'	25:1H:39:C:C6	2.49	0.47
25:1H:2137:C:H42	25:1H:2154:G:H1	1.62	0.47
25:1H:2790:A:H2'	25:1H:2791:C:C5'	2.44	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
27:71:21:THR:HA	27:71:225:ASN:HB3	1.95	0.47
35:58:116:LEU:HD23	35:58:119:ARG:HD2	1.94	0.47
41:B8:108:ARG:O	41:B8:111:ARG:HG2	2.14	0.47
1:1G:56:U:H2'	1:1G:57:G:H8	1.79	0.47
1:1G:191(E):G:H2'	1:1G:191(F):U:C6	2.50	0.47
1:1G:963:G:H21	10:1A:55:LYS:CE	2.27	0.47
1:1G:1317:C:N4	14:5A:19:ARG:HH21	2.12	0.47
1:1G:1452:C:O2'	1:1G:1453:G:O5'	2.33	0.47
57:2L:1:C:N4	38:45:87:LYS:HE2	2.28	0.47
25:14:226:G:O2'	25:14:228:A:N1	2.46	0.47
25:14:307:G:H21	25:14:330:A:H62	1.62	0.47
25:14:605:C:H1'	25:14:657:U:O2'	2.14	0.47
25:14:1790:C:O2'	28:19:209:ALA:HB2	2.14	0.47
30:39:50:SER:HB2	30:39:94:PRO:HD3	1.95	0.47
31:49:43:LEU:HD12	31:49:43:LEU:HA	1.75	0.47
38:45:41:TRP:HB3	38:45:94:VAL:HB	1.96	0.47
39:55:21:TYR:CZ	39:55:43:GLU:HG2	2.50	0.47
47:D5:153:SER:C	47:D5:155:LEU:N	2.72	0.47
1:13:130:A:OP2	17:8I:63:ARG:NH2	2.43	0.47
1:13:312:C:H2'	1:13:313:A:C8	2.49	0.47
1:13:445:G:H2'	1:13:446:G:C8	2.50	0.47
1:13:1070:U:H2'	1:13:1071:C:C6	2.47	0.47
1:13:1298:C:H4'	1:13:1299:A:N9	2.29	0.47
4:3E:148:VAL:HG12	4:3E:149:ALA:H	1.79	0.47
6:5E:11:ASN:HD22	6:5E:14:LEU:HG	1.79	0.47
11:2I:95:ILE:HD12	11:2I:108:ILE:HD13	1.97	0.47
22:1K:9:G:H5'	22:1K:10:G:OP2	2.15	0.47
22:1K:14:A:H2'	22:1K:15:G:O4'	2.14	0.47
23:2K:44:A:H2'	23:2K:45:A:C8	2.49	0.47
25:1H:270(M):U:O2'	25:1H:270(N):G:O5'	2.26	0.47
25:1H:642:G:H21	25:1H:646:A:H2	1.63	0.47
25:1H:751:A:H5'	44:E8:90:ARG:HG2	1.95	0.47
25:1H:1483:G:N2	25:1H:1507:A:H1'	2.30	0.47
25:1H:2131:G:N2	25:1H:2158:A:H62	2.12	0.47
25:1H:2418:A:OP2	56:Q8:29:LYS:NZ	2.39	0.47
25:1H:2474:C:H5''	25:1H:2475:C:C5	2.49	0.47
25:1H:2851:A:O3'	39:98:64:ARG:NH2	2.48	0.47
32:51:101:ARG:NH2	32:51:122:THR:HA	2.29	0.47
34:38:74:LEU:HB2	34:38:110:GLY:O	2.13	0.47
41:B8:86:ILE:HD12	41:B8:87:ASP:N	2.29	0.47
42:C8:98:LEU:O	42:C8:102:GLU:N	2.33	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:403:C:H2'	1:1G:404:U:C6	2.49	0.47
1:1G:664:G:N2	1:1G:741:G:H1	2.01	0.47
1:1G:728:A:C5	15:6A:54:ARG:HD2	2.49	0.47
5:42:41:VAL:O	5:42:66:MET:HA	2.13	0.47
17:8A:86:GLU:O	17:8A:90:ILE:HG12	2.15	0.47
19:AA:41:VAL:C	19:AA:43:GLU:H	2.22	0.47
25:14:270(G):C:H42	25:14:270(S):G:H1	1.62	0.47
25:14:1035:U:H2'	25:14:1036:G:C8	2.49	0.47
25:14:1667:G:H8	25:14:1667:G:OP2	1.97	0.47
25:14:1717:G:H1	25:14:1742:C:H42	1.61	0.47
25:14:1935:G:H1'	25:14:1964:G:N2	2.29	0.47
26:1J:97:G:H2'	26:1J:98:G:O4'	2.15	0.47
28:19:9:TYR:CD1	28:19:10:THR:HG22	2.50	0.47
28:19:201:HIS:O	28:19:204:ILE:HG12	2.15	0.47
30:39:65:TRP:HB3	30:39:70:THR:HG21	1.97	0.47
33:69:79:ILE:HG22	33:69:81:VAL:HG22	1.95	0.47
1:13:520:A:N1	1:13:536:C:H1'	2.29	0.47
2:1E:130:ARG:HD3	2:1E:138:LEU:HD11	1.97	0.47
25:1H:500:G:N2	25:1H:503:A:O5'	2.39	0.47
25:1H:704:G:H2'	25:1H:726:G:H22	1.79	0.47
25:1H:744:G:OP1	62:1H:3689:HOH:O	2.20	0.47
25:1H:1550:C:H2'	25:1H:1551:C:C6	2.50	0.47
25:1H:2428:G:N2	37:78:61:ARG:HH21	2.12	0.47
31:41:14:GLU:O	31:41:17:PRO:HD2	2.15	0.47
32:51:9:ILE:HG13	32:51:9:ILE:O	2.15	0.47
41:B8:98:LYS:HB3	41:B8:100:TYR:CE2	2.49	0.47
1:1G:957:U:N3	1:1G:960:U:OP2	2.34	0.47
1:1G:1003:G:N2	1:1G:1038:C:N3	2.63	0.47
1:1G:1097:C:O2'	1:1G:1169:A:N3	2.45	0.47
13:4A:97:PRO:HG3	13:4A:107:ALA:HB1	1.97	0.47
19:AA:62:ILE:HG23	19:AA:66:MET:SD	2.54	0.47
25:14:849:A:N6	25:14:929:G:O2'	2.40	0.47
25:14:1138:G:H21	35:15:106:MET:HE3	1.79	0.47
25:14:1161:C:H2'	25:14:1162:G:C8	2.50	0.47
25:14:1385:G:H4'	25:14:1386:C:OP1	2.15	0.47
25:14:1543:A:C2'	25:14:1544:C:H5''	2.45	0.47
25:14:2134:A:OP2	25:14:2157:G:N2	2.48	0.47
32:59:91:GLY:HA3	32:59:94:TYR:CD2	2.49	0.47
33:69:3:VAL:HG12	33:69:38:LEU:HA	1.97	0.47
37:35:5:ASP:HA	37:35:7:ARG:HH22	1.78	0.47
40:65:87:PHE:CZ	40:65:102:ALA:HB2	2.50	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:D5:99:TYR:HB3	47:D5:123:ASP:HB3	1.97	0.47
1:13:142:G:H2'	1:13:143:A:H8	1.80	0.47
1:13:359:U:H2'	1:13:360:A:C8	2.50	0.47
1:13:405:U:H3'	1:13:406:G:H5'	1.96	0.47
1:13:991:U:O2'	1:13:992:U:O5'	2.31	0.47
1:13:1188:A:H5''	14:5I:58:LYS:HZ1	1.80	0.47
1:13:1234:C:H2'	1:13:1235:U:O4'	2.15	0.47
2:1E:22:LYS:HA	2:1E:40:HIS:NE2	2.30	0.47
3:2E:153:VAL:HA	3:2E:197:GLY:O	2.14	0.47
4:3E:26:CYS:HA	60:3E:302:SF4:S1	2.55	0.47
4:3E:101:LEU:HD23	4:3E:135:LEU:O	2.14	0.47
8:7E:77:GLU:HG2	8:7E:78:GLN:H	1.80	0.47
13:4I:93:ARG:HB3	13:4I:94:ARG:HH11	1.80	0.47
22:3K:57:C:H5''	25:1H:2169:A:H1'	1.97	0.47
25:1H:574:C:N3	29:21:145:LYS:NZ	2.47	0.47
25:1H:581:C:H2'	25:1H:582:G:H8	1.78	0.47
25:1H:1302:A:O5'	25:1H:1302:A:H8	1.98	0.47
25:1H:2285:C:C5	54:O8:27:LYS:HD2	2.50	0.47
25:1H:2372:G:H2'	25:1H:2373:G:C8	2.49	0.47
25:1H:2537:U:H2'	25:1H:2538:C:C6	2.50	0.47
25:1H:2576:G:O2'	25:1H:2579:C:OP2	2.32	0.47
27:71:200:LYS:HA	27:71:208:PHE:HE2	1.79	0.47
29:21:118:LYS:HG2	29:21:118:LYS:O	2.13	0.47
30:31:81:PRO:HB2	30:31:89:VAL:HG22	1.97	0.47
30:31:167:ALA:HA	30:31:170:LEU:HD12	1.97	0.47
32:51:101:ARG:NH2	32:51:121:ILE:O	2.47	0.47
35:58:42:TRP:HA	35:58:48:MET:HE1	1.97	0.47
37:78:46:LYS:HB3	37:78:46:LYS:HE2	1.81	0.47
42:C8:108:GLU:OE1	42:C8:112:ARG:NH1	2.36	0.47
1:1G:191:G:O2'	20:BA:103:GLY:HA2	2.15	0.47
1:1G:413:G:HO2'	1:1G:414:A:P	2.38	0.47
1:1G:537:G:H5''	12:3A:110:ARG:NH1	2.29	0.47
1:1G:1251:A:H2'	1:1G:1252:A:C8	2.50	0.47
1:1G:1317:C:H5'	1:1G:1318:A:OP2	2.15	0.47
1:1G:1416:G:OP2	62:1G:1811:HOH:O	2.20	0.47
2:12:27:LYS:HB3	2:12:194:PRO:HD2	1.95	0.47
3:22:72:LYS:O	3:22:74:GLY:N	2.47	0.47
3:22:123:GLN:O	3:22:128:PHE:HB2	2.15	0.47
5:42:147:ASP:OD1	5:42:147:ASP:N	2.47	0.47
6:52:8:ILE:HG13	6:52:88:VAL:HG22	1.97	0.47
6:52:52:ILE:HD13	6:52:87:ARG:HG3	1.95	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:62:25:ALA:O	7:62:28:ASN:ND2	2.48	0.47
8:72:12:ARG:HH11	8:72:26:VAL:HA	1.78	0.47
8:72:39:LEU:HG	8:72:44:PHE:HB2	1.97	0.47
9:82:38:GLN:HG2	9:82:39:GLY:N	2.29	0.47
25:14:67:U:H2'	25:14:68:G:H8	1.80	0.47
25:14:71:A:OP2	25:14:71:A:H3'	2.14	0.47
25:14:445:C:OP1	42:85:2:PRO:HA	2.15	0.47
25:14:466:A:H5'	55:L5:21:ARG:NH2	2.29	0.47
25:14:971:C:H2'	25:14:972:G:O4'	2.15	0.47
25:14:1257:C:H4'	30:39:83:PHE:CD1	2.49	0.47
25:14:1364:G:OP1	49:F5:3:LYS:HD2	2.14	0.47
25:14:1616:A:H8	62:14:4024:HOH:O	1.98	0.47
25:14:1882:C:H5'	25:14:1883:G:OP2	2.14	0.47
25:14:2815:C:H2'	25:14:2816:C:H6	1.79	0.47
29:29:50:GLY:HA2	29:29:78:LEU:HD12	1.96	0.47
35:15:53:VAL:HG11	35:15:128:HIS:CD2	2.50	0.47
42:85:52:ARG:HA	42:85:55:ARG:HG3	1.97	0.47
43:95:35:LEU:HD21	43:95:57:VAL:HB	1.95	0.47
43:95:76:LYS:HD2	43:95:80:GLN:O	2.14	0.47
46:C5:99:CYS:SG	46:C5:105:ALA:HB3	2.54	0.47
52:I5:12:ALA:HB1	52:I5:24:THR:N	2.29	0.47
1:13:60:A:H4'	1:13:61:G:O5'	2.15	0.47
1:13:186(A):C:H5''	20:BI:86:ARG:HH11	1.79	0.47
1:13:1397:C:O2'	1:13:1398:A:OP2	2.30	0.47
5:4E:112:LEU:HD22	5:4E:112:LEU:HA	1.77	0.47
23:2K:54:G:H2'	23:2K:55:5MU:C6	2.50	0.47
25:1H:200:U:O2	25:1H:386:G:N2	2.48	0.47
25:1H:442:G:N2	30:31:48:THR:OG1	2.47	0.47
32:51:131:VAL:HG22	32:51:132:ARG:H	1.80	0.47
36:68:87:ILE:HG22	36:68:93:PRO:HA	1.97	0.47
41:B8:1:MET:O	41:B8:1:MET:HG2	2.15	0.47
41:B8:105:LEU:HD12	41:B8:110:ILE:HG12	1.96	0.47
43:D8:68:LYS:HD3	43:D8:68:LYS:HA	1.57	0.47
45:F8:60:ARG:HH12	55:P8:47:ARG:HH22	1.63	0.47
50:K8:28:LYS:HD2	50:K8:53:LEU:HD21	1.97	0.47
1:1G:188:U:O2'	1:1G:189:U:H5'	2.15	0.47
1:1G:1150:U:O4	1:1G:1151:A:N6	2.48	0.47
2:12:207:ALA:O	2:12:211:ILE:HG13	2.14	0.47
8:72:12:ARG:HD2	8:72:26:VAL:HG23	1.97	0.47
13:4A:3:ARG:HH11	31:49:113:ARG:HE	1.62	0.47
13:4A:44:ARG:H	13:4A:44:ARG:NE	2.13	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:AA:19:VAL:O	19:AA:23:ASN:HB2	2.15	0.47
25:14:226:G:H2'	25:14:227:A:C8	2.50	0.47
25:14:270(I):G:H2'	25:14:270(J):G:C8	2.49	0.47
25:14:1025:G:O2'	25:14:1026:U:OP1	2.33	0.47
25:14:1105:U:H2'	25:14:1106:G:C8	2.49	0.47
25:14:1316:U:H2'	25:14:1317:A:C8	2.48	0.47
25:14:2849:U:OP2	41:75:95:ARG:NH1	2.48	0.47
25:14:2884:U:H2'	25:14:2885:C:O4'	2.15	0.47
28:19:255:LYS:HE3	28:19:255:LYS:N	2.25	0.47
30:39:54:ARG:HG3	30:39:80:ALA:HA	1.96	0.47
31:49:114:ILE:HD13	31:49:115:ARG:H	1.79	0.47
33:69:91:SER:HB2	33:69:121:LYS:HD2	1.95	0.47
50:G5:15:LYS:H	50:G5:67:LYS:NZ	2.12	0.47
1:13:130:A:H5'	17:8I:63:ARG:CZ	2.44	0.47
1:13:1343:G:H2'	1:13:1344:C:C6	2.49	0.47
5:4E:33:VAL:HG11	5:4E:109:ILE:HG12	1.97	0.47
19:AI:50:ALA:HA	19:AI:58:VAL:O	2.15	0.47
22:1K:27:G:H2'	22:1K:28:U:O4'	2.15	0.47
22:1K:43:G:OP2	22:1K:43:G:H8	1.98	0.47
25:1H:296:C:H2'	25:1H:297:C:H6	1.80	0.47
25:1H:517:C:OP2	53:N8:13:LYS:NZ	2.47	0.47
25:1H:558:G:H5''	35:58:112:LEU:HD22	1.97	0.47
25:1H:1083:U:OP2	34:38:47:ASN:HB2	2.14	0.47
25:1H:1092:C:H2'	25:1H:1093:G:H5'	1.97	0.47
25:1H:1265:A:H3'	53:N8:19:ARG:NH1	2.30	0.47
25:1H:1562:A:H2'	25:1H:1563:G:O4'	2.15	0.47
28:11:201:HIS:O	28:11:204:ILE:HG12	2.15	0.47
32:51:54:ARG:HD3	32:51:65:HIS:ND1	2.30	0.47
32:51:78:GLY:O	32:51:136:ILE:HG22	2.15	0.47
32:51:157:TYR:CZ	32:51:172:LYS:HG3	2.50	0.47
33:61:109:ILE:HB	33:61:130:TYR:OH	2.15	0.47
36:68:78:ARG:HH21	41:B8:103:ARG:NH2	2.13	0.47
37:78:1:MET:SD	37:78:2:LYS:N	2.84	0.47
37:78:63:PRO:HG2	56:Q8:25:MET:HB2	1.97	0.47
38:88:109:VAL:HG13	38:88:113:GLN:HB3	1.95	0.47
47:H8:35:ARG:HB3	47:H8:35:ARG:HH11	1.80	0.47
51:L8:13:ILE:HD13	51:L8:13:ILE:HA	1.70	0.47
1:1G:324:G:OP1	20:BA:22:ARG:HD2	2.14	0.47
1:1G:544:G:P	4:32:62:GLN:HE21	2.38	0.47
1:1G:885:G:H2'	1:1G:886:G:H8	1.80	0.47
1:1G:1322:C:O2'	1:1G:1323:G:H5'	2.15	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:12:50:GLU:O	2:12:54:THR:OG1	2.20	0.47
2:12:172:ILE:H	2:12:172:ILE:HD12	1.79	0.47
3:22:13:GLY:HA3	14:5A:57:ARG:NE	2.29	0.47
4:32:155:LEU:HA	4:32:155:LEU:HD13	1.67	0.47
5:42:110:LEU:HD12	5:42:118:ILE:HG21	1.97	0.47
25:14:601:C:O2	25:14:605:C:H4'	2.15	0.47
25:14:706:A:C2	25:14:707:G:H1'	2.50	0.47
25:14:1022:G:O2'	25:14:1024:G:O6	2.32	0.47
25:14:1860:G:H1	25:14:1882:C:N4	2.12	0.47
25:14:2210:G:H2'	25:14:2210:G:N3	2.28	0.47
25:14:2251:OMG:HM21	25:14:2449:U:C2	2.50	0.47
25:14:2329:G:H2'	25:14:2330:G:C8	2.50	0.47
25:14:2692:C:H2'	25:14:2693:A:H8	1.79	0.47
29:29:111:ARG:HB2	29:29:160:TYR:HB3	1.96	0.47
30:39:194:MET:HB2	30:39:198:ALA:HB3	1.97	0.47
42:85:66:ASN:O	42:85:70:ARG:HG2	2.14	0.47
47:D5:5:LEU:HD23	47:D5:47:VAL:HG21	1.97	0.47
47:D5:104:PHE:O	47:D5:107:THR:OG1	2.30	0.47
1:13:129(A):G:O6	1:13:187:C:O2'	2.20	0.47
1:13:614:A:H61	1:13:626:U:H3	1.63	0.47
1:13:878:G:H5'	8:7E:89:PRO:HG2	1.97	0.47
1:13:1288:A:N1	1:13:1371:G:H1'	2.30	0.47
2:1E:76:GLN:O	2:1E:208:ILE:HG13	2.15	0.47
7:6E:20:ASP:OD1	7:6E:21:VAL:N	2.48	0.47
20:BI:77:ALA:O	20:BI:81:LYS:HB2	2.15	0.47
25:1H:38:A:N6	25:1H:440:G:O6	2.48	0.47
25:1H:1084:A:H5''	25:1H:1085:A:H8	1.79	0.47
25:1H:1085:A:HO2'	25:1H:1086:A:P	2.35	0.47
25:1H:1257:C:H2'	25:1H:1258:C:C6	2.50	0.47
25:1H:1790:C:H5''	25:1H:1791:A:OP1	2.14	0.47
25:1H:2293:C:H42	25:1H:2339:G:H1	1.62	0.47
25:1H:2396:G:H1'	49:J8:30:VAL:HG12	1.97	0.47
28:11:2:ALA:O	28:11:20:ASP:HB2	2.15	0.47
28:11:108:PRO:HG3	28:11:143:HIS:HE1	1.79	0.47
31:41:179:PRO:HB3	52:M8:38:LYS:CD	2.44	0.47
36:68:120:GLU:OE1	41:B8:67:SER:OG	2.31	0.47
42:C8:76:TYR:CZ	42:C8:80:ILE:HG13	2.50	0.47
45:F8:60:ARG:NH2	55:P8:47:ARG:HH12	2.13	0.47
46:G8:5:MET:HB2	46:G8:5:MET:HE3	1.42	0.47
51:L8:4:LEU:O	51:L8:36:VAL:HA	2.15	0.47
1:1G:119:A:H4'	1:1G:120:A:C8	2.50	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:452:A:O2'	1:1G:453:A:O4'	2.33	0.47
1:1G:1149:C:P	9:82:9:ARG:HH11	2.38	0.47
1:1G:1177:G:H2'	1:1G:1178:G:C4	2.50	0.47
2:12:208:ILE:HD13	2:12:211:ILE:HD12	1.96	0.47
3:22:113:ALA:HB3	3:22:114:PRO:HD3	1.97	0.47
8:72:12:ARG:NH1	8:72:26:VAL:HA	2.30	0.47
25:14:1354:A:O3'	28:19:38:LYS:NZ	2.47	0.47
26:1J:15:A:OP2	26:1J:69:G:N2	2.45	0.47
48:E5:56:ASP:OD2	48:E5:58:THR:OG1	2.30	0.47
49:F5:4:VAL:HG11	49:F5:11:ARG:HD2	1.96	0.47
51:H5:35:ARG:HD2	51:H5:35:ARG:HA	1.55	0.47
52:I5:37:SER:OG	52:I5:38:LYS:N	2.46	0.47
1:13:160:A:H2'	1:13:161:A:O4'	2.15	0.46
1:13:299:G:H2'	1:13:300:A:C8	2.50	0.46
1:13:801:U:H2'	1:13:802:A:H8	1.78	0.46
1:13:1438:G:H1	1:13:1463:C:H42	1.63	0.46
9:8E:25:LYS:N	9:8E:60:ASP:OD1	2.47	0.46
9:8E:26:VAL:HG22	9:8E:61:ALA:HB3	1.95	0.46
18:9I:29:PHE:CD1	18:9I:29:PHE:N	2.83	0.46
22:3K:42:C:H2'	22:3K:43:G:C8	2.50	0.46
25:1H:500:G:N1	25:1H:503:A:OP2	2.48	0.46
25:1H:989:G:OP2	51:L8:11:SER:OG	2.26	0.46
25:1H:1085:A:O2'	25:1H:1086:A:OP1	2.27	0.46
25:1H:1173:G:H5''	25:1H:1174:A:OP1	2.15	0.46
25:1H:2122:U:H2'	25:1H:2123:G:C8	2.50	0.46
25:1H:2556:C:H2'	25:1H:2557:G:O4'	2.14	0.46
25:1H:2688:U:H1'	25:1H:2721:A:N6	2.30	0.46
1:1G:1443:G:C6	41:75:118:ARG:HB2	2.51	0.46
25:14:485:C:H42	25:14:495:G:H1	1.63	0.46
25:14:774:A:OP1	28:19:48:ARG:NH2	2.47	0.46
25:14:856:C:O2'	25:14:857:C:OP1	2.31	0.46
25:14:1029:A:N6	25:14:1125:G:O2'	2.48	0.46
25:14:1639:U:H2'	25:14:1640:C:H5''	1.97	0.46
25:14:2247:A:H2'	25:14:2248:C:H6	1.80	0.46
25:14:2247:A:H2'	25:14:2248:C:C6	2.50	0.46
25:14:2712:U:O2'	25:14:2713:A:H5'	2.15	0.46
28:19:246:PRO:HD2	28:19:255:LYS:HD3	1.96	0.46
32:59:157:TYR:CE2	32:59:172:LYS:HG3	2.50	0.46
37:35:1:MET:HE2	37:35:5:ASP:HB3	1.96	0.46
37:35:57:THR:HG21	37:35:60:MET:N	2.31	0.46
41:75:50:ILE:HD11	41:75:102:ILE:HD11	1.96	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
43:95:24:LYS:HB2	43:95:90:PRO:HB2	1.96	0.46
1:13:407:G:H2'	1:13:408:A:C8	2.50	0.46
1:13:465:A:N6	1:13:467:G:O6	2.44	0.46
3:2E:22:TRP:HB3	3:2E:59:ARG:HB2	1.98	0.46
5:4E:6:PHE:CE1	5:4E:36:ASP:HB3	2.50	0.46
7:6E:78:ARG:HG3	7:6E:79:ARG:N	2.30	0.46
9:8E:65:VAL:HG21	9:8E:73:GLN:HB3	1.97	0.46
14:5I:24:CYS:SG	14:5I:26:ARG:N	2.88	0.46
16:7I:22:THR:HA	16:7I:33:ILE:HG12	1.97	0.46
20:BI:43:LEU:HA	20:BI:46:GLU:HB2	1.97	0.46
23:2K:69:C:H2'	23:2K:70:C:H6	1.80	0.46
25:1H:270(H):C:H42	25:1H:270(R):G:H1	1.62	0.46
25:1H:873:G:H1	25:1H:904:C:H42	1.64	0.46
25:1H:1020:A:N6	25:1H:1141:U:O2'	2.49	0.46
25:1H:1213:A:N3	25:1H:1238:G:O2'	2.43	0.46
25:1H:2185:C:H2'	25:1H:2186:G:C8	2.50	0.46
27:7I:190:ARG:HG3	27:7I:193:ILE:HD12	1.96	0.46
28:11:17:THR:HG22	28:11:204:ILE:HA	1.97	0.46
29:21:17:ASP:N	29:21:17:ASP:OD1	2.46	0.46
41:B8:97:ALA:O	41:B8:98:LYS:HD2	2.16	0.46
47:H8:45:ASP:OD1	47:H8:49:ARG:NE	2.47	0.46
48:I8:68:GLU:HG3	48:I8:80:HIS:HD2	1.79	0.46
54:O8:52:VAL:HG22	54:O8:53:LYS:HG3	1.97	0.46
1:1G:1269:A:OP2	1:1G:1269:A:H8	1.97	0.46
1:1G:1279:A:OP2	10:1A:9:ARG:HD3	2.15	0.46
1:1G:1325:C:H5''	21:1B:6:ARG:HH21	1.80	0.46
1:1G:1395:C:H5'	1:1G:1402:4OC:O2'	2.16	0.46
4:32:32:ALA:N	60:32:303:SF4:S1	2.88	0.46
8:72:36:LEU:O	8:72:40:ALA:N	2.41	0.46
12:3A:25:LYS:HE3	12:3A:25:LYS:HB3	1.84	0.46
57:2L:7:G:O2'	57:2L:8:4SU:H5''	2.14	0.46
25:14:139:G:H1'	25:14:140:A:H2	1.79	0.46
25:14:618:G:OP2	30:39:106:ARG:NH2	2.36	0.46
25:14:629:G:OP2	56:M5:15:LYS:NZ	2.48	0.46
25:14:1297:C:H2'	25:14:1298:C:C6	2.50	0.46
25:14:2128:C:H2'	25:14:2129:C:O4'	2.14	0.46
25:14:2734:A:N6	25:14:2770:G:O2'	2.48	0.46
31:49:135:LEU:HD22	31:49:155:MET:HE2	1.96	0.46
43:95:35:LEU:HB2	43:95:37:VAL:HG13	1.97	0.46
50:G5:44:LEU:C	50:G5:46:GLN:H	2.23	0.46
1:13:663:A:H5''	18:9I:61:LYS:HE3	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1222:G:H5'	19:AI:78:ARG:NH1	2.31	0.46
2:1E:82:ARG:HB3	2:1E:82:ARG:HH11	1.80	0.46
16:7I:74:LEU:HB3	16:7I:79:VAL:HB	1.98	0.46
22:1K:16:C:O3'	22:1K:61:U:O2'	2.33	0.46
25:1H:2805:G:H2'	25:1H:2807:G:C8	2.50	0.46
25:1H:2892:A:H2'	25:1H:2893:G:O4'	2.16	0.46
26:16:111:U:H2'	26:16:112:G:C8	2.50	0.46
28:11:147:LEU:HG	28:11:155:LEU:HD21	1.97	0.46
29:21:11:MET:HE2	29:21:11:MET:HB3	1.67	0.46
34:38:113:GLN:O	34:38:113:GLN:NE2	2.48	0.46
35:58:15:LEU:O	35:58:136:GLU:HA	2.14	0.46
1:1G:707:C:H4'	11:2A:20:TYR:CD2	2.50	0.46
1:1G:936:C:H2'	1:1G:937:A:O4'	2.16	0.46
1:1G:1064:G:O2'	1:1G:1065:U:O5'	2.27	0.46
6:52:16:GLN:H	6:52:16:GLN:CD	2.22	0.46
8:72:86:ILE:HD11	8:72:136:GLU:HG2	1.96	0.46
14:5A:50:LYS:HB3	14:5A:52:GLN:HG3	1.97	0.46
25:14:700:G:H2'	25:14:701:G:O4'	2.14	0.46
25:14:1636:C:H2'	25:14:1637:A:H8	1.79	0.46
25:14:2018:G:H2'	25:14:2019:A:O4'	2.15	0.46
25:14:2065:C:H2'	25:14:2066:C:O4'	2.15	0.46
25:14:2458:G:O2'	25:14:2489:G:O6	2.34	0.46
25:14:2557:G:H2'	25:14:2558:C:H6	1.79	0.46
25:14:2734:A:H3'	25:14:2735:G:H8	1.79	0.46
25:14:2838:G:H1	25:14:2880:C:H42	1.63	0.46
26:1J:114:G:O2'	40:65:50:SER:HB3	2.16	0.46
28:19:234:GLY:N	62:19:405:HOH:O	2.48	0.46
38:45:40:ALA:HB3	38:45:127:ILE:HD11	1.97	0.46
48:E5:24:LYS:HB2	48:E5:37:LEU:HA	1.97	0.46
48:E5:36:ILE:HA	48:E5:60:PHE:HA	1.96	0.46
53:J5:49:CYS:HA	53:J5:56:LYS:HD2	1.97	0.46
1:13:143:A:H2	1:13:220:G:H1	1.60	0.46
1:13:633:G:H5'	1:13:634:C:OP2	2.15	0.46
1:13:1020:U:H2'	1:13:1021:G:C8	2.50	0.46
1:13:1059:C:O2'	10:1I:53:PRO:HD3	2.15	0.46
1:13:1187:G:H4'	9:8E:111:ARG:HH11	1.79	0.46
1:13:1254:C:OP1	10:1I:45:ARG:NE	2.47	0.46
5:4E:10:MET:HA	5:4E:32:VAL:HA	1.97	0.46
11:2I:99:GLN:HG2	11:2I:105:VAL:HG21	1.97	0.46
19:AI:40:ILE:HG21	19:AI:66:MET:O	2.15	0.46
25:1H:586:A:H5'	30:31:89:VAL:HG21	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:931:G:O3'	51:L8:24:LYS:NZ	2.49	0.46
25:1H:972:G:OP2	25:1H:973:A:O2'	2.32	0.46
25:1H:1007:C:O3'	35:58:108:PRO:HB3	2.15	0.46
25:1H:1107:G:H2'	25:1H:1108:U:C6	2.50	0.46
25:1H:1380:G:N2	25:1H:1570:A:N1	2.62	0.46
25:1H:2581:G:H3'	62:1H:4038:HOH:O	2.14	0.46
30:31:106:ARG:H	30:31:106:ARG:HG2	1.47	0.46
33:61:21:VAL:HG22	33:61:22:LYS:H	1.79	0.46
1:1G:222:U:H2'	1:1G:223:U:C6	2.50	0.46
1:1G:1320:C:H5''	19:AA:36:ARG:HH21	1.80	0.46
1:1G:1466:C:H2'	1:1G:1467:G:O4'	2.16	0.46
15:6A:74:ASP:HB2	15:6A:77:ARG:HH21	1.81	0.46
22:3L:73:A:H2'	22:3L:74:A:O4'	2.15	0.46
25:14:2117:A:H2'	25:14:2118:U:C5	2.51	0.46
25:14:2439:A:O2'	25:14:2440:C:OP2	2.29	0.46
25:14:2737:G:H2'	25:14:2738:A:C8	2.50	0.46
25:14:2776:A:H4'	25:14:2777:G:O5'	2.15	0.46
31:49:145:THR:C	31:49:147:ASP:H	2.23	0.46
33:69:76:THR:HG23	33:69:140:LEU:HD13	1.98	0.46
41:75:105:LEU:C	41:75:107:ASP:H	2.23	0.46
46:C5:27:VAL:HA	46:C5:39:VAL:HA	1.98	0.46
47:D5:148:ASP:O	47:D5:172:ALA:HB3	2.15	0.46
48:E5:15:ASP:OD1	48:E5:16:SER:N	2.48	0.46
48:E5:23:VAL:HG22	48:E5:38:VAL:HG22	1.97	0.46
49:F5:51:VAL:HG11	49:F5:74:VAL:HG21	1.98	0.46
1:13:643:C:H5'	8:7E:31:PHE:CD1	2.51	0.46
1:13:1235:U:H5''	21:1F:3:LYS:HD2	1.96	0.46
10:1I:49:VAL:HG13	14:5I:41:ARG:HG3	1.98	0.46
12:3I:88:LYS:HG3	12:3I:89:OTD:H5	1.98	0.46
14:5I:50:LYS:HE2	14:5I:50:LYS:HB3	1.79	0.46
22:1K:44:A:H2'	22:1K:45:A:C8	2.50	0.46
25:1H:1068:G:O2'	25:1H:1096:A:N3	2.46	0.46
25:1H:1140:C:OP2	35:58:66:LYS:NZ	2.47	0.46
25:1H:1252:G:O2'	25:1H:1253:A:O5'	2.29	0.46
25:1H:1430:C:H2'	25:1H:1431:U:C6	2.51	0.46
25:1H:1497:U:H3'	25:1H:1498:C:H6	1.80	0.46
25:1H:2099:U:H2'	25:1H:2100:G:C8	2.50	0.46
25:1H:2292:C:P	40:A8:17:ARG:HH22	2.38	0.46
25:1H:2294:C:N4	25:1H:2338:G:H1	2.14	0.46
30:31:126:VAL:H	30:31:195:ASP:HA	1.79	0.46
31:41:17:PRO:HA	31:41:20:ILE:HG13	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:51:42:ARG:HG2	32:51:43:VAL:N	2.30	0.46
37:78:97:PRO:C	37:78:99:LEU:H	2.23	0.46
1:1G:956:U:H1'	1:1G:1225:A:H2	1.81	0.46
1:1G:1190:G:H4'	3:22:176:HIS:CE1	2.50	0.46
1:1G:1353:G:H1	1:1G:1369:C:H42	1.64	0.46
7:62:47:CYS:HB3	7:62:58:PRO:HG3	1.98	0.46
9:82:28:VAL:HG13	9:82:29:ASN:O	2.16	0.46
25:14:848:G:OP2	25:14:929:G:N2	2.39	0.46
25:14:1331:A:O2'	25:14:1332:G:H8	1.99	0.46
25:14:1801:G:OP2	28:19:154:LYS:NZ	2.38	0.46
25:14:1815:A:P	28:19:54:ARG:HH22	2.39	0.46
25:14:1902:C:H5'	28:19:246:PRO:HD3	1.97	0.46
25:14:2734:A:H5'	25:14:2735:G:OP2	2.15	0.46
29:29:11:MET:HG2	29:29:24:THR:HB	1.98	0.46
35:15:99:LEU:O	35:15:103:VAL:HG23	2.15	0.46
44:A5:4:LYS:HE2	44:A5:4:LYS:HB3	1.84	0.46
46:C5:42:VAL:HG23	46:C5:43:ASN:H	1.80	0.46
49:F5:85:LEU:HD13	49:F5:88:LYS:HB2	1.97	0.46
1:13:177:C:H2'	1:13:178:C:C6	2.51	0.46
1:13:496:A:N3	1:13:496:A:H2'	2.29	0.46
1:13:603:U:H2'	1:13:604:G:C8	2.50	0.46
1:13:690:G:H2'	1:13:691:G:O4'	2.16	0.46
1:13:956:U:H2'	1:13:957:U:O4'	2.15	0.46
1:13:973:G:O4'	10:1I:55:LYS:HG2	2.16	0.46
1:13:1262:C:H2'	1:13:1263:C:C6	2.50	0.46
1:13:1298:C:P	7:6E:114:ARG:HH22	2.39	0.46
12:3I:102:TYR:HD1	12:3I:102:TYR:HA	1.70	0.46
19:AI:10:PHE:HE2	19:AI:12:ASP:HB3	1.80	0.46
25:1H:268:C:H42	25:1H:424:G:H1	1.62	0.46
25:1H:566:U:OP1	37:78:29:LYS:HE2	2.15	0.46
25:1H:1176:G:O2'	25:1H:1178:C:N4	2.49	0.46
29:21:77:ILE:HD13	29:21:77:ILE:HA	1.67	0.46
42:C8:66:ASN:HB2	42:C8:76:TYR:HB2	1.97	0.46
44:E8:48:ALA:O	44:E8:52:GLU:HG2	2.16	0.46
1:1G:1001:G:H22	1:1G:1039:C:N4	2.12	0.46
1:1G:1046:A:H3'	1:1G:1047:G:H8	1.81	0.46
7:62:29:LYS:HE2	7:62:101:LEU:HD12	1.97	0.46
9:82:47:LEU:HD12	9:82:50:LEU:HD12	1.98	0.46
20:BA:82:SER:HB2	20:BA:86:ARG:HD2	1.97	0.46
25:14:70:G:OP1	25:14:112:U:N3	2.40	0.46
25:14:247:G:N1	25:14:247:G:C5	2.79	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:489:G:N2	25:14:1321:A:OP1	2.44	0.46
25:14:806:C:OP2	37:35:41:ARG:NH2	2.36	0.46
25:14:877:U:H2'	25:14:878:A:H5''	1.97	0.46
25:14:972:G:OP2	25:14:974:G:H5''	2.14	0.46
25:14:1434:A:H61	25:14:1558:A:N6	2.13	0.46
25:14:1936:A:H4'	25:14:1937:A:O5'	2.15	0.46
28:19:95:LEU:HD11	28:19:103:ARG:HB2	1.97	0.46
31:49:67:LYS:NZ	52:I5:5:ILE:HG21	2.31	0.46
42:85:16:LYS:O	42:85:20:LEU:HD23	2.16	0.46
43:95:35:LEU:HD23	43:95:37:VAL:HG22	1.98	0.46
43:95:55:ALA:HB1	43:95:101:GLY:HA2	1.98	0.46
44:A5:88:ARG:HA	44:A5:88:ARG:HD2	1.73	0.46
47:D5:117:LEU:H	47:D5:117:LEU:HG	1.57	0.46
1:13:260:G:H2'	1:13:261:U:C6	2.50	0.46
1:13:762:C:H2'	1:13:763:G:H8	1.80	0.46
1:13:1532:U:O2	24:4K:9:G:H8	1.99	0.46
2:1E:87:ARG:CZ	2:1E:219:VAL:HB	2.46	0.46
15:6I:39:LEU:HD13	15:6I:56:LEU:HD12	1.97	0.46
18:9I:22:VAL:HG13	18:9I:42:ARG:NH1	2.30	0.46
19:AI:3:ARG:HA	19:AI:3:ARG:HD2	1.72	0.46
22:1K:54:G:O2'	38:88:51:ARG:NH2	2.49	0.46
22:1K:59:A:HO2'	22:1K:61:U:H6	1.63	0.46
25:1H:99:U:H1'	25:1H:101:G:OP2	2.16	0.46
25:1H:144:C:H4'	45:F8:2:LYS:HG3	1.96	0.46
25:1H:2143:C:H2'	25:1H:2144:U:O4'	2.16	0.46
25:1H:2314:C:H2'	25:1H:2315:G:C8	2.45	0.46
25:1H:2822:G:O2'	25:1H:2824:C:OP2	2.27	0.46
34:38:98:LYS:HD2	34:38:98:LYS:HA	1.74	0.46
35:58:42:TRP:HE3	35:58:48:MET:HE1	1.80	0.46
35:58:130:HIS:HB3	35:58:134:ARG:HH22	1.81	0.46
36:68:68:GLU:OE2	36:68:78:ARG:NH1	2.49	0.46
41:B8:13:ARG:CZ	41:B8:13:ARG:HB3	2.46	0.46
53:N8:40:LYS:NZ	53:N8:46:CYS:HB3	2.31	0.46
1:1G:362:G:N2	1:1G:365:U:OP2	2.48	0.46
1:1G:975:A:H5'	1:1G:1363:A:H62	1.79	0.46
1:1G:1004:A:O2'	1:1G:1005:A:OP2	2.28	0.46
1:1G:1226:C:O2'	13:4A:111:LYS:NZ	2.44	0.46
2:12:32:ILE:O	2:12:33:TYR:HB2	2.14	0.46
6:52:54:LYS:N	6:52:54:LYS:HD3	2.31	0.46
8:72:104:ARG:HB3	8:72:108:GLY:H	1.80	0.46
10:1A:22:LYS:HD2	10:1A:26:ALA:HB2	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:3A:15:VAL:HG23	12:3A:16:ARG:H	1.80	0.46
15:6A:5:LYS:O	15:6A:9:GLN:HG2	2.16	0.46
26:1J:89(A):A:C5	26:1J:90:C:H1'	2.50	0.46
30:39:29:ASN:OD1	30:39:32:LEU:HB2	2.16	0.46
31:49:118:ARG:HB3	31:49:181:ARG:HE	1.81	0.46
1:13:122:G:H2'	1:13:123:C:O4'	2.15	0.46
1:13:190:G:HO2'	1:13:191(A):G:P	2.38	0.46
1:13:960:U:H4'	1:13:961:U:O5'	2.15	0.46
1:13:1235:U:H5''	21:1F:3:LYS:HB2	1.98	0.46
1:13:1368:G:H5''	9:8E:112:LYS:HB3	1.97	0.46
2:1E:16:HIS:CE1	2:1E:209:ARG:HH21	2.33	0.46
4:3E:13:ARG:HB3	4:3E:33:MET:HE3	1.98	0.46
5:4E:139:LEU:O	5:4E:142:LEU:HB2	2.16	0.46
8:7E:104:ARG:HB3	8:7E:107:LEU:HB2	1.98	0.46
8:7E:112:LEU:HA	8:7E:134:ILE:HG12	1.97	0.46
13:4I:12:ASN:O	13:4I:13:LYS:HB2	2.15	0.46
25:1H:481:G:H4'	25:1H:482:A:O5'	2.14	0.46
25:1H:957:A:N1	25:1H:2458:G:H4'	2.31	0.46
25:1H:2687:U:C4	25:1H:2688:U:C5	3.04	0.46
25:1H:2831:G:O2'	25:1H:2884:U:OP1	2.23	0.46
27:71:6:ARG:HH12	27:71:7:TYR:HB2	1.81	0.46
28:11:76:PRO:HB2	28:11:116:GLN:HE21	1.81	0.46
30:31:155:LEU:HD13	30:31:174:VAL:CG1	2.46	0.46
31:41:163:ALA:HB1	31:41:168:GLU:HB2	1.98	0.46
33:61:130:TYR:C	33:61:131:LYS:HD2	2.41	0.46
34:38:30:GLN:H	34:38:30:GLN:CD	2.23	0.46
39:98:2:ARG:HG3	39:98:5:LYS:HB2	1.98	0.46
41:B8:112:ARG:HA	41:B8:115:ARG:HH11	1.81	0.46
44:E8:37:ARG:NH2	53:N8:48:GLU:OE2	2.48	0.46
1:1G:10:A:OP2	5:42:126:ARG:HG2	2.15	0.46
1:1G:334:C:H2'	1:1G:335:C:C6	2.51	0.46
1:1G:1000:A:C8	1:1G:1001:G:H1'	2.50	0.46
1:1G:1028(B):C:H3'	1:1G:1029:G:H4'	1.97	0.46
1:1G:1516:G:N2	1:1G:1519:MA6:OP2	2.47	0.46
2:12:121:LEU:HD23	2:12:126:GLU:HG3	1.98	0.46
3:22:136:GLN:HA	3:22:139:GLN:HB3	1.98	0.46
5:42:72:GLN:O	5:42:75:THR:HG22	2.16	0.46
10:1A:49:VAL:O	10:1A:60:ARG:HB2	2.15	0.46
19:AA:9:VAL:HG11	52:I5:63:TYR:HD1	1.81	0.46
19:AA:65:ASN:HA	52:I5:55:ARG:NH1	2.31	0.46
25:14:818:G:H5'	25:14:819:A:OP1	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:846:C:H42	25:14:931:G:H1	1.63	0.46
25:14:1210:A:H5''	25:14:1211:U:H3'	1.98	0.46
26:1J:105:G:H5''	47:D5:31:ARG:HG3	1.97	0.46
28:19:43:ARG:HG3	28:19:47:GLY:O	2.16	0.46
29:29:32:PRO:HA	29:29:90:THR:HA	1.98	0.46
38:45:139:GLU:HG2	47:D5:74:VAL:HB	1.97	0.46
2:1E:104:ASN:OD1	2:1E:107:THR:OG1	2.32	0.46
2:1E:163:PHE:CD1	2:1E:185:ILE:HG13	2.50	0.46
25:1H:154:G:H2'	25:1H:155:C:O4'	2.16	0.46
25:1H:172:C:H2'	25:1H:173:G:H8	1.80	0.46
25:1H:1140:C:OP1	35:58:23:LEU:HB3	2.16	0.46
25:1H:1248:G:P	30:31:92:PRO:HG3	2.56	0.46
25:1H:2153:G:H2'	25:1H:2154:G:O4'	2.15	0.46
25:1H:2212:A:H1'	25:1H:2215:G:C4	2.51	0.46
25:1H:2520:C:H2'	25:1H:2521:C:H6	1.81	0.46
25:1H:2731:G:H8	25:1H:2731:G:O5'	1.99	0.46
32:51:8:PRO:HD2	32:51:69:ARG:NE	2.31	0.46
34:38:87:VAL:HG21	34:38:91:LYS:HD3	1.98	0.46
43:D8:35:LEU:HB3	43:D8:57:VAL:HG13	1.97	0.46
47:H8:111:VAL:HG23	47:H8:117:LEU:HB2	1.98	0.46
47:H8:156:LYS:N	47:H8:156:LYS:HE2	2.30	0.46
1:1G:93:U:H2'	1:1G:95:G:H8	1.81	0.46
1:1G:1115:C:H42	1:1G:1185:G:H1	1.64	0.46
1:1G:1157:A:H1'	1:1G:1158:C:C2	2.50	0.46
1:1G:1304:G:N1	1:1G:1332:A:OP2	2.44	0.46
1:1G:1490:C:H2'	1:1G:1491:G:O4'	2.16	0.46
9:82:32:ASP:HB3	9:82:34:ASN:H	1.81	0.46
15:6A:29:VAL:HG11	15:6A:67:LEU:HD21	1.97	0.46
20:BA:67:ALA:HB2	20:BA:77:ALA:HB2	1.97	0.46
25:14:270(I):G:H2'	25:14:270(J):G:H8	1.81	0.46
25:14:633:A:H1'	25:14:2403:C:H4'	1.98	0.46
25:14:673:C:H5''	30:39:81:PRO:HD2	1.97	0.46
25:14:767:U:H2'	25:14:768:G:H8	1.80	0.46
25:14:1794:U:H2'	25:14:1795:C:H6	1.80	0.46
25:14:2105:C:H2'	25:14:2106:G:O4'	2.16	0.46
25:14:2683:C:P	41:75:53:ARG:HH22	2.38	0.46
25:14:2847:U:OP1	41:75:98:LYS:HD3	2.16	0.46
26:1J:28:C:OP2	40:65:33:LYS:HE3	2.16	0.46
33:69:120:ILE:HG22	33:69:122:GLU:H	1.80	0.46
37:35:64:LYS:H	37:35:64:LYS:HG3	1.60	0.46
47:D5:48:PHE:HE1	47:D5:71:VAL:HG11	1.81	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:D5:156:LYS:O	47:D5:157:LEU:HD22	2.16	0.46
1:13:833:U:H2'	1:13:834:C:H6	1.81	0.46
1:13:885:G:H2'	1:13:886:G:C8	2.51	0.46
1:13:1216:G:P	14:5I:3:ARG:HH21	2.39	0.46
3:2E:18:TRP:CD1	14:5I:54:PRO:HA	2.51	0.46
3:2E:73:PRO:HG3	3:2E:105:GLU:HG3	1.98	0.46
13:4I:94:ARG:HH22	25:1H:887:A:P	2.39	0.46
22:1K:15:G:H21	22:1K:22:A:H1'	1.80	0.46
22:3K:22:A:N6	22:3K:47:G:H2'	2.27	0.46
25:1H:639:U:H2'	25:1H:640:C:C6	2.51	0.46
25:1H:730:C:H3'	62:1H:3622:HOH:O	2.15	0.46
25:1H:1265:A:H8	25:1H:1265:A:OP1	1.99	0.46
25:1H:1510:A:H2'	25:1H:1511:A:C8	2.51	0.46
25:1H:1819:A:H4'	25:1H:1820:U:O5'	2.16	0.46
25:1H:2028:U:O4	62:1H:3636:HOH:O	2.20	0.46
25:1H:2154:G:C2	25:1H:2155:G:C5	3.04	0.46
25:1H:2228:G:C5	25:1H:2229:C:C4	3.04	0.46
25:1H:2572:A:C8	29:21:144:ARG:HB3	2.51	0.46
30:31:7:TYR:HD2	30:31:21:ALA:HB1	1.81	0.46
42:C8:91:ASP:OD1	42:C8:96:ALA:HB2	2.16	0.46
44:E8:29:LEU:HD21	44:E8:33:ARG:CZ	2.45	0.46
45:F8:63:LYS:O	45:F8:64:LYS:HD2	2.16	0.46
52:M8:63:TYR:HA	52:M8:66:SER:OG	2.16	0.46
1:1G:3:G:H5''	1:1G:4:U:OP2	2.16	0.46
1:1G:29:G:O2'	1:1G:295:C:H4'	2.16	0.46
1:1G:109:A:C6	1:1G:326:G:C6	3.03	0.46
1:1G:1082:G:OP1	5:42:18:ARG:HD3	2.15	0.46
6:52:7:ASN:HD21	18:9A:34:TYR:HE2	1.62	0.46
22:3L:48:U:H3'	22:3L:49:C:H4'	1.98	0.46
25:14:910:A:C5	38:45:13:GLN:HG3	2.50	0.46
25:14:1166:C:H42	25:14:1182:A:H61	1.63	0.46
25:14:2130:U:O4	25:14:2158:A:O2'	2.30	0.46
32:59:43:VAL:HG22	32:59:44:VAL:H	1.80	0.46
36:25:88:ASN:O	36:25:90:GLN:N	2.49	0.46
37:35:11:GLY:C	37:35:13:ASN:H	2.23	0.46
42:85:88:ILE:HG13	42:85:88:ILE:O	2.15	0.46
47:D5:146:ILE:HD12	47:D5:175:VAL:HA	1.98	0.46
51:H5:8:LEU:HD12	51:H5:30:ARG:O	2.16	0.46
1:13:191:G:C1'	20:BI:105:SER:HB3	2.46	0.45
2:1E:136:VAL:HA	2:1E:139:LYS:HB2	1.97	0.45
3:2E:150:LYS:HB3	3:2E:201:TYR:HB2	1.97	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:3E:9:CYS:HB3	4:3E:32:ALA:HB2	1.99	0.45
7:6E:57:GLU:HB2	7:6E:60:LYS:HD2	1.98	0.45
20:BI:26:ASN:HB3	20:BI:71:THR:HG23	1.97	0.45
22:3K:65:G:H2'	22:3K:66:C:O4'	2.16	0.45
25:1H:675:A:H4'	30:31:67:GLN:OE1	2.15	0.45
25:1H:1297:C:OP1	25:1H:2710:C:H4'	2.16	0.45
25:1H:1795:C:H2'	25:1H:1796:U:O4'	2.15	0.45
25:1H:2114:A:O2'	25:1H:2168:G:OP2	2.33	0.45
31:41:75:LYS:HA	31:41:84:LYS:HG3	1.98	0.45
1:1G:324:G:N2	1:1G:326:G:H3'	2.30	0.45
1:1G:419:C:H42	1:1G:424:G:H1	1.63	0.45
1:1G:1376:U:OP1	7:62:98:SER:OG	2.25	0.45
1:1G:1414:U:H2'	1:1G:1415:G:H8	1.81	0.45
4:32:61:LYS:HD3	4:32:206:PHE:CE2	2.51	0.45
5:42:9:LYS:HD2	5:42:9:LYS:HA	1.81	0.45
9:82:18:PHE:O	9:82:19:LEU:HB2	2.15	0.45
25:14:30:G:H2'	25:14:31:C:C6	2.51	0.45
25:14:188:G:H1	25:14:208:C:H42	1.64	0.45
25:14:978:G:H1	25:14:985:C:N4	2.14	0.45
25:14:1151:G:H5''	42:85:81:HIS:CD2	2.50	0.45
25:14:1153:C:H2'	25:14:1154:G:O4'	2.16	0.45
25:14:2883:A:H5'	25:14:2884:U:H5'	1.98	0.45
30:39:164:ARG:O	30:39:168:ARG:HB2	2.15	0.45
37:35:85:LEU:HD23	37:35:85:LEU:H	1.80	0.45
48:E5:27:GLU:OE1	48:E5:69:PHE:N	2.44	0.45
1:13:376:G:H5''	16:7I:5:ARG:HB2	1.98	0.45
1:13:437:U:OP1	4:3E:155:LEU:HD21	2.17	0.45
1:13:1305:G:N2	1:13:1331:G:H2'	2.30	0.45
1:13:1389:C:H2'	1:13:1390:U:O4'	2.16	0.45
1:13:1434:A:H2'	1:13:1435:G:O4'	2.15	0.45
5:4E:94:ALA:HB2	5:4E:119:LEU:HG	1.97	0.45
6:5E:10:LEU:HD13	6:5E:61:LEU:HD13	1.98	0.45
8:7E:39:LEU:HB3	8:7E:45:ILE:HG12	1.97	0.45
19:AI:44:MET:H	19:AI:44:MET:HG3	1.49	0.45
23:2K:24:C:H2'	23:2K:25:U:C6	2.52	0.45
24:4K:18:G:H4'	24:4K:19:A:H5'	1.98	0.45
25:1H:1313:U:OP1	62:1H:3697:HOH:O	2.21	0.45
25:1H:1568:G:P	28:11:63:ARG:HH12	2.37	0.45
25:1H:1782:C:H1'	25:1H:2609:U:H5''	1.98	0.45
25:1H:2064:C:H2'	25:1H:2065:C:C6	2.51	0.45
25:1H:2564:A:OP1	25:1H:2648:C:H4'	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:31:12:LEU:HD11	30:31:17:ARG:HE	1.81	0.45
34:38:7:VAL:C	34:38:11:ALA:HB2	2.41	0.45
34:38:23:SER:HB2	34:38:68:LEU:HB2	1.98	0.45
46:G8:5:MET:HE1	46:G8:32:PRO:HB3	1.98	0.45
1:1G:1219:U:O2'	19:AA:34:TRP:HB3	2.16	0.45
2:12:22:LYS:O	2:12:24:TRP:N	2.49	0.45
11:2A:83:ILE:HG12	11:2A:109:VAL:HB	1.99	0.45
11:2A:98:LEU:O	11:2A:101:SER:OG	2.22	0.45
13:4A:3:ARG:HD3	31:49:113:ARG:HH21	1.80	0.45
22:3L:77:A:O2'	25:14:2394:C:N3	2.40	0.45
25:14:395:U:H2'	25:14:396:G:N7	2.32	0.45
25:14:539:G:H2'	25:14:540:G:C8	2.51	0.45
25:14:602:G:OP2	25:14:602:G:H8	1.98	0.45
25:14:661:C:H1'	37:35:12:ALA:HA	1.98	0.45
25:14:818:G:H3'	25:14:1187:G:H22	1.81	0.45
25:14:1024:G:C3'	25:14:1025:G:H5''	2.45	0.45
25:14:1469:A:H2'	25:14:1470:G:O4'	2.16	0.45
25:14:1612:C:O3'	55:L5:5:TRP:HD1	2.00	0.45
25:14:1805:U:O2	28:19:50:THR:HB	2.16	0.45
25:14:2142:C:H2'	25:14:2143:C:C6	2.51	0.45
25:14:2364:C:H4'	48:E5:56:ASP:OD1	2.16	0.45
25:14:2438:U:O3'	25:14:2439:A:H3'	2.16	0.45
26:1J:24:G:H4'	26:1J:25:A:O5'	2.16	0.45
28:19:225:ALA:O	62:19:403:HOH:O	2.21	0.45
28:19:253:GLN:HB3	28:19:255:LYS:NZ	2.31	0.45
41:75:123:GLN:HG3	41:75:124:ASP:OD1	2.16	0.45
42:85:98:LEU:O	42:85:99:ALA:HB3	2.16	0.45
1:13:262:A:H2'	1:13:263:A:C8	2.52	0.45
1:13:292:G:H21	1:13:608:A:H61	1.64	0.45
1:13:407:G:OP1	4:3E:115:ARG:NH1	2.50	0.45
1:13:736:C:H2'	1:13:737:A:C8	2.51	0.45
1:13:812:C:O2'	1:13:813:U:H6	2.00	0.45
1:13:1298:C:N4	7:6E:114:ARG:HB3	2.32	0.45
2:1E:16:HIS:HA	2:1E:210:SER:HB2	1.98	0.45
7:6E:15:ASP:OD1	7:6E:16:LEU:N	2.47	0.45
7:6E:86:GLN:HE21	7:6E:86:GLN:HB2	1.57	0.45
25:1H:2152:G:C2	25:1H:2153:G:N7	2.85	0.45
25:1H:2378:A:H4'	40:A8:23:ARG:CZ	2.46	0.45
25:1H:2444:G:OP2	30:31:68:LYS:NZ	2.45	0.45
25:1H:2845:G:H5''	41:B8:55:ASN:HA	1.98	0.45
31:41:61:ALA:HB2	31:41:68:PRO:HD3	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:68:73:ASP:OD1	41:B8:32:TYR:OH	2.31	0.45
38:88:47:ILE:HD12	38:88:70:PRO:HG3	1.99	0.45
41:B8:49:VAL:HG21	41:B8:52:ILE:HD11	1.98	0.45
49:J8:64:ALA:HA	49:J8:67:ILE:HG13	1.97	0.45
1:1G:35:G:O2'	12:3A:115:SER:O	2.27	0.45
1:1G:560:U:H4'	1:1G:561:U:O5'	2.16	0.45
1:1G:818:G:O3'	1:1G:819:A:H4'	2.16	0.45
1:1G:1119:C:P	9:82:9:ARG:HH21	2.37	0.45
1:1G:1143:G:H2'	1:1G:1144:G:C8	2.52	0.45
9:82:58:HIS:HB3	9:82:59:PHE:HD1	1.81	0.45
10:1A:6:ILE:HG22	10:1A:98:ILE:HG12	1.98	0.45
18:9A:88:LYS:HE3	18:9A:88:LYS:HB2	1.72	0.45
25:14:276:A:O2'	25:14:277:C:O4'	2.19	0.45
25:14:755:C:H2'	25:14:756:C:C6	2.51	0.45
25:14:1015:G:H2'	25:14:1016:G:H8	1.81	0.45
26:1J:56:G:H4'	26:1J:57:A:C8	2.51	0.45
30:39:174:VAL:HG11	30:39:188:ARG:HH22	1.81	0.45
31:49:94:LEU:HD12	31:49:99:MET:HA	1.98	0.45
32:59:3:ARG:HH21	32:59:7:LEU:HD11	1.81	0.45
41:75:12:SER:O	41:75:15:VAL:HG22	2.16	0.45
41:75:21:GLU:O	41:75:91:ARG:NH1	2.41	0.45
41:75:23:ARG:HG2	41:75:26:ASP:OD2	2.17	0.45
1:13:429:U:H3'	4:3E:9:CYS:SG	2.56	0.45
1:13:692:U:H2'	1:13:694:A:OP2	2.17	0.45
1:13:977:A:O2'	1:13:979:C:OP2	2.27	0.45
2:1E:63:MET:HB3	2:1E:64:ARG:HH21	1.81	0.45
6:5E:20:ALA:O	6:5E:24:GLU:N	2.35	0.45
19:AI:33:THR:OG1	19:AI:34:TRP:N	2.46	0.45
25:1H:7:G:H2'	25:1H:8:A:C8	2.51	0.45
25:1H:510:C:H5''	62:1H:3631:HOH:O	2.16	0.45
25:1H:787:U:H5''	25:1H:788:A:H5'	1.98	0.45
25:1H:1139:G:O2'	25:1H:1143:A:N1	2.38	0.45
25:1H:2037:G:H2'	25:1H:2038:G:C8	2.52	0.45
25:1H:2126:A:H2	25:1H:2162:G:N2	2.11	0.45
25:1H:2679:A:OP2	29:21:160:TYR:OH	2.30	0.45
29:21:52:LEU:O	29:21:76:ARG:HG2	2.17	0.45
29:21:78:LEU:HD13	29:21:78:LEU:HA	1.82	0.45
29:21:104:VAL:HG11	29:21:188:VAL:HG13	1.99	0.45
36:68:47:ILE:HD12	36:68:48:PRO:HD2	1.98	0.45
41:B8:124:ASP:H	41:B8:127:ALA:CB	2.26	0.45
42:C8:85:LYS:HD3	42:C8:85:LYS:HA	1.72	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:H8:19:ARG:HD3	47:H8:19:ARG:N	2.31	0.45
47:H8:100:VAL:HG11	47:H8:134:PRO:HG2	1.99	0.45
52:M8:42:PHE:CZ	52:M8:43:TYR:HD1	2.34	0.45
1:1G:334:C:H2'	1:1G:335:C:H6	1.81	0.45
1:1G:614:A:H5'	1:1G:615:C:OP2	2.17	0.45
1:1G:1357:A:H2	1:1G:1365:G:H22	1.63	0.45
2:12:17:PHE:CD1	2:12:42:ILE:HG12	2.51	0.45
7:62:41:ARG:O	7:62:45:ASP:HB2	2.17	0.45
25:14:265:A:H2'	25:14:266:G:C4'	2.46	0.45
25:14:580:C:H2'	25:14:581:C:C6	2.52	0.45
25:14:1153:C:OP1	42:85:76:TYR:OH	2.34	0.45
25:14:1416:G:H2'	25:14:1417:C:C6	2.52	0.45
25:14:1536:A:C8	25:14:1537:C:H1'	2.51	0.45
25:14:1657:C:H2'	25:14:1658:C:C6	2.51	0.45
25:14:1709:U:H2'	25:14:1710:C:C6	2.51	0.45
25:14:1942:5MC:OP2	25:14:1943:U:O2'	2.25	0.45
35:15:4:TYR:CG	42:85:100:VAL:HG11	2.51	0.45
36:25:68:GLU:HB3	36:25:78:ARG:HB2	1.97	0.45
38:45:135:ASP:OD1	38:45:135:ASP:N	2.41	0.45
40:65:36:TYR:HA	40:65:52:SER:HB3	1.98	0.45
45:B5:12:VAL:HG12	45:B5:29:TRP:CD1	2.52	0.45
1:13:955:U:H1'	1:13:1227:A:H61	1.82	0.45
1:13:1055:A:H62	1:13:1200:C:H42	1.63	0.45
1:13:1086:U:H2'	1:13:1087:G:O4'	2.17	0.45
1:13:1314:C:C5	19:AI:4:SER:HB2	2.50	0.45
21:1F:3:LYS:HB3	21:1F:14:TRP:CD1	2.51	0.45
25:1H:988:A:H8	25:1H:988:A:O5'	1.99	0.45
25:1H:1403:C:H5''	25:1H:1471:A:H1'	1.97	0.45
25:1H:1667:G:O2'	25:1H:1991:U:O4	2.24	0.45
25:1H:2439:A:H3'	25:1H:2439:A:P	2.57	0.45
30:31:64:ILE:HG23	30:31:65:TRP:CD1	2.52	0.45
33:61:122:GLU:O	33:61:126:TYR:OH	2.27	0.45
38:88:79:LEU:HD23	38:88:79:LEU:HA	1.63	0.45
48:I8:7:LEU:HD23	48:I8:11:ARG:HG2	1.98	0.45
1:1G:225:C:H2'	1:1G:226:G:C8	2.52	0.45
1:1G:1011:G:N2	1:1G:1019:C:O2	2.50	0.45
1:1G:1075:C:H2'	1:1G:1076:C:C6	2.51	0.45
1:1G:1352:C:N4	1:1G:1370:G:H1	2.06	0.45
1:1G:1434:A:H2'	1:1G:1435:G:O4'	2.16	0.45
1:1G:1446:A:OP1	1:1G:1446:A:H4'	2.16	0.45
8:72:46:LYS:HB2	8:72:62:TYR:HB2	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:82:34:ASN:HB3	9:82:35:GLU:H	1.68	0.45
16:7A:7:ALA:HB2	16:7A:20:VAL:HG11	1.98	0.45
19:AA:15:LEU:HD13	19:AA:19:VAL:HG21	1.99	0.45
19:AA:84:GLY:C	19:AA:86:GLU:H	2.25	0.45
25:14:270:A:OP2	25:14:270(Y):G:N2	2.28	0.45
25:14:300:A:O2'	25:14:318:C:O2	2.35	0.45
25:14:320:A:H4'	25:14:322:A:C8	2.51	0.45
25:14:574:C:N3	29:29:145:LYS:NZ	2.60	0.45
25:14:807:U:O2'	25:14:2060:A:N1	2.48	0.45
25:14:1032:A:O2'	25:14:1033:U:H5'	2.17	0.45
25:14:1386:C:H2'	25:14:1387:C:H6	1.81	0.45
25:14:1537:C:O2'	25:14:1538:G:O4'	2.14	0.45
25:14:2078:C:C4	25:14:2079:U:C4	3.04	0.45
25:14:2130:U:O3'	25:14:2131:G:H4'	2.16	0.45
25:14:2876:G:OP1	41:75:3:ARG:HG2	2.17	0.45
26:1J:56:G:OP1	31:49:27:ASN:ND2	2.49	0.45
28:19:260:ARG:CZ	28:19:264:LYS:HD3	2.46	0.45
30:39:29:ASN:O	30:39:112:MET:HE1	2.16	0.45
32:59:72:ILE:H	32:59:72:ILE:HD12	1.82	0.45
32:59:96:ALA:O	32:59:125:VAL:HG11	2.17	0.45
37:35:56:SER:O	37:35:57:THR:HB	2.16	0.45
39:55:87:TYR:CE2	39:55:117:VAL:HG22	2.51	0.45
40:65:35:ILE:HG21	40:65:69:VAL:HG11	1.99	0.45
42:85:92:ARG:NH2	43:95:10:LYS:HA	2.23	0.45
50:G5:70:GLN:HG2	50:G5:71:ASN:H	1.81	0.45
52:I5:2:LYS:HB2	52:I5:5:ILE:CG2	2.46	0.45
1:13:491:G:H2'	1:13:492:G:O4'	2.16	0.45
1:13:625:G:H2'	1:13:626:U:C6	2.52	0.45
1:13:712:A:H2'	1:13:713:G:C8	2.52	0.45
1:13:1218:C:H2'	1:13:1219:U:C6	2.51	0.45
2:1E:86:GLU:C	2:1E:89:GLY:H	2.25	0.45
4:3E:153:ARG:HE	4:3E:153:ARG:HB3	1.64	0.45
25:1H:67:U:H2'	25:1H:68:G:H8	1.81	0.45
25:1H:1729:A:H2'	25:1H:1730:U:H5''	1.98	0.45
25:1H:2052:G:H4'	29:21:143:ASN:O	2.17	0.45
25:1H:2336:A:H61	48:I8:43:THR:CG2	2.30	0.45
31:41:7:LEU:HB2	31:41:104:GLU:HG3	1.99	0.45
52:M8:34:GLU:HG3	52:M8:35:VAL:H	1.81	0.45
1:1G:615:C:H2'	1:1G:616:G:O4'	2.16	0.45
1:1G:1034:G:H2'	1:1G:1035:A:O4'	2.17	0.45
1:1G:1475:G:OP1	25:14:1689:A:H1'	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1540:U:H5'	18:9A:55:ARG:NH1	2.31	0.45
7:62:16:LEU:HG	9:82:41:VAL:HG12	1.98	0.45
12:3A:57:LEU:HB3	12:3A:59:SER:H	1.82	0.45
19:AA:16:LEU:HD12	19:AA:20:LEU:HG	1.98	0.45
22:3L:6:G:H4'	22:3L:7:G:OP1	2.16	0.45
25:14:83:G:H2'	46:C5:95:LYS:HD2	1.99	0.45
25:14:1050:A:H2'	25:14:1051:G:O4'	2.16	0.45
25:14:2615:U:H2'	25:14:2616:C:H6	1.82	0.45
28:19:26:LYS:HB3	28:19:83:GLU:HG2	1.98	0.45
28:19:165:ILE:HD13	28:19:175:LEU:HD21	1.98	0.45
29:29:111:ARG:HG3	29:29:160:TYR:CD2	2.52	0.45
30:39:135:LYS:HA	30:39:135:LYS:HD2	1.72	0.45
30:39:149:ASP:OD1	30:39:149:ASP:N	2.43	0.45
33:69:50:ARG:HA	33:69:53:ALA:HB3	1.98	0.45
44:A5:86:LEU:HD12	44:A5:87:PRO:HD2	1.98	0.45
1:13:236:G:H2'	1:13:237:C:C6	2.52	0.45
1:13:291:C:H42	1:13:309:G:H1	1.65	0.45
1:13:438:G:N2	1:13:495:A:C8	2.84	0.45
1:13:731:G:H2'	1:13:732:C:C6	2.51	0.45
1:13:821:G:H4'	62:13:1912:HOH:O	2.17	0.45
1:13:963:G:H1	1:13:972:C:H42	1.65	0.45
1:13:1014:A:H2	1:13:1219:U:O2	2.00	0.45
25:1H:71:A:H2	45:F8:31:HIS:HE2	1.65	0.45
25:1H:592:G:H1	25:1H:665:C:H42	1.63	0.45
25:1H:1021:A:C8	25:1H:1021:A:C3'	3.00	0.45
25:1H:1270:C:H5''	25:1H:1271:G:O5'	2.17	0.45
25:1H:1908:C:H2'	25:1H:1909:C:H6	1.82	0.45
25:1H:2402:C:H2'	25:1H:2403:C:H5'	1.98	0.45
27:71:6:ARG:NH1	27:71:7:TYR:HB2	2.32	0.45
32:51:26:VAL:HG12	32:51:32:GLU:HA	1.99	0.45
38:88:24:GLY:C	38:88:26:TYR:H	2.24	0.45
54:O8:9:LEU:HD22	54:O8:9:LEU:HA	1.71	0.45
54:O8:35:GLU:O	54:O8:51:GLU:HB2	2.16	0.45
1:1G:406:G:H5'	4:32:5:ILE:HD13	1.98	0.45
5:42:33:VAL:HG11	5:42:109:ILE:HA	1.98	0.45
6:52:98:LEU:HD22	18:9A:28:GLU:HG3	1.97	0.45
7:62:69:VAL:HG12	7:62:103:TRP:HE3	1.82	0.45
9:82:82:ALA:HB1	9:82:96:LEU:HD21	1.98	0.45
22:3L:19:G:N2	22:3L:58:A:H5'	2.32	0.45
25:14:196:A:OP2	37:35:46:LYS:HE2	2.15	0.45
25:14:328:U:H4'	46:C5:68:HIS:ND1	2.32	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:389:G:N2	37:35:72:PRO:HD3	2.30	0.45
25:14:1686:C:H2'	25:14:1687:G:O4'	2.16	0.45
25:14:1972:A:H2'	25:14:1973:G:H8	1.82	0.45
25:14:2057:A:H2'	25:14:2058:A:O4'	2.17	0.45
37:35:52:GLU:OE1	37:35:54:GLY:N	2.49	0.45
41:75:11:GLU:O	41:75:15:VAL:HG13	2.16	0.45
1:13:269:C:H2'	1:13:270:A:C8	2.51	0.45
1:13:880:C:OP1	12:3I:5:ASN:ND2	2.49	0.45
1:13:914:A:H2'	1:13:915:A:C8	2.41	0.45
1:13:983:A:H5''	1:13:984:C:OP2	2.17	0.45
3:2E:6:HIS:HD2	3:2E:7:PRO:HD2	1.81	0.45
3:2E:52:LEU:HD23	3:2E:52:LEU:H	1.81	0.45
10:1I:39:PRO:HB3	10:1I:70:ARG:NH1	2.32	0.45
11:2I:20:TYR:N	11:2I:31:THR:O	2.48	0.45
25:1H:95:G:O2'	50:K8:48:HIS:HB3	2.17	0.45
25:1H:898:C:H2'	25:1H:899:A:H5'	1.99	0.45
25:1H:1301:A:O2'	25:1H:1302:A:P	2.75	0.45
26:16:48:A:H2'	26:16:49:C:C6	2.52	0.45
29:21:67:PHE:CE1	29:21:74:PRO:HA	2.52	0.45
32:51:102:ALA:HA	32:51:117:PRO:HD3	1.99	0.45
35:58:55:VAL:HB	35:58:126:PRO:HA	1.98	0.45
40:A8:26:LEU:HD22	40:A8:87:PHE:CD1	2.51	0.45
41:B8:126:ALA:HA	41:B8:129:ARG:HB2	1.98	0.45
43:D8:17:GLY:N	43:D8:96:ILE:O	2.38	0.45
54:O8:6:ARG:NH1	54:O8:6:ARG:HA	2.31	0.45
1:1G:581:G:OP1	15:6A:61:GLY:HA3	2.15	0.45
1:1G:1206:G:H2'	1:1G:1207:2MG:C8	2.51	0.45
3:22:131:ARG:HE	3:22:166:GLU:CD	2.24	0.45
4:32:13:ARG:NH1	4:32:38:TYR:O	2.39	0.45
9:82:1:MET:N	9:82:20:ARG:HH12	2.15	0.45
21:1B:15:ARG:HB2	21:1B:15:ARG:HH11	1.81	0.45
25:14:446:G:OP1	42:85:3:ARG:NH1	2.47	0.45
25:14:464:U:H4'	55:L5:5:TRP:CZ3	2.51	0.45
25:14:1111:A:H5'	32:59:3:ARG:NH1	2.31	0.45
25:14:1869:G:H5'	25:14:1870:C:OP2	2.16	0.45
25:14:2495:G:H5''	38:45:81:VAL:HG22	1.99	0.45
25:14:2517:C:H42	25:14:2567:G:H1	1.65	0.45
26:1J:45:A:O4'	31:49:95:ARG:NH1	2.50	0.45
28:19:215:LEU:HD23	28:19:215:LEU:HA	1.79	0.45
29:29:96:PHE:HD2	29:29:182:LEU:HD21	1.81	0.45
30:39:154:VAL:HB	30:39:173:VAL:HG22	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:35:90:ARG:HG3	37:35:91:PHE:CE2	2.51	0.45
41:75:110:ILE:HD12	41:75:110:ILE:HG23	1.72	0.45
47:D5:91:LEU:HD23	47:D5:91:LEU:H	1.82	0.45
50:G5:66:GLU:O	50:G5:70:GLN:HB2	2.17	0.45
1:13:155:C:H2'	1:13:156:G:C8	2.52	0.45
1:13:952:U:H4'	1:13:964:A:H61	1.82	0.45
1:13:1316:G:N2	1:13:1318:A:H3'	2.31	0.45
14:5I:3:ARG:HG2	14:5I:6:LEU:HD12	1.98	0.45
25:1H:264:C:C2'	25:1H:265:A:H5''	2.46	0.45
25:1H:274:G:N3	25:1H:275:G:H5''	2.32	0.45
25:1H:1308:A:H2'	25:1H:1309:G:O4'	2.17	0.45
25:1H:1796:U:H2'	25:1H:1797:C:C6	2.52	0.45
25:1H:2107:C:H5''	27:7I:2:LYS:HE3	1.99	0.45
28:11:132:PRO:HG3	28:11:190:TYR:CE1	2.52	0.45
31:41:114:ILE:HG22	31:41:115:ARG:N	2.32	0.45
35:58:35:ARG:O	35:58:37:LYS:N	2.40	0.45
37:78:101:VAL:HG22	37:78:106:LEU:HB3	1.99	0.45
40:A8:24:LEU:H	40:A8:24:LEU:HD22	1.81	0.45
1:1G:294:U:OP1	1:1G:610:G:O2'	2.33	0.45
1:1G:370:C:N4	1:1G:391:G:H1	2.11	0.45
1:1G:458:C:N3	1:1G:474:G:N2	2.55	0.45
2:12:133:LYS:O	2:12:137:ARG:NH1	2.50	0.45
7:62:65:ALA:O	7:62:69:VAL:HG23	2.17	0.45
20:BA:39:LYS:HA	20:BA:42:GLN:HB3	1.98	0.45
20:BA:84:LEU:HD22	20:BA:84:LEU:HA	1.82	0.45
25:14:616:A:H4'	25:14:617:G:OP1	2.16	0.45
25:14:981:A:H8	25:14:982:C:C5	2.35	0.45
25:14:1025:G:H8	25:14:1025:G:OP1	2.00	0.45
25:14:2577:A:H2'	25:14:2614:A:N6	2.32	0.45
25:14:2679:A:OP2	29:29:160:TYR:OH	2.34	0.45
30:39:2:LYS:HG3	30:39:24:LEU:HD12	1.99	0.45
32:59:144:VAL:O	32:59:148:ILE:HG12	2.17	0.45
33:69:125:GLU:HA	33:69:141:LYS:CB	2.47	0.45
35:15:56:ASN:H	35:15:125:GLY:N	2.14	0.45
39:55:87:TYR:HD2	39:55:90:ARG:HD2	1.80	0.45
47:D5:30:ASN:CG	47:D5:32:HIS:H	2.25	0.45
48:E5:49:LYS:HD3	48:E5:49:LYS:HA	1.30	0.45
53:J5:40:LYS:HG3	53:J5:46:CYS:HB3	1.99	0.45
4:3E:99:SER:O	4:3E:140:VAL:HG22	2.17	0.45
9:8E:32:ASP:O	9:8E:36:TYR:N	2.46	0.45
9:8E:47:LEU:H	9:8E:47:LEU:HD22	1.82	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:7I:26:ARG:NH1	62:7I:201:HOH:O	2.50	0.45
21:1F:12:LYS:HG2	21:1F:22:ARG:HB3	1.99	0.45
25:1H:300:A:H1'	25:1H:319:C:H1'	1.99	0.45
25:1H:300:A:H8	46:G8:84:ARG:HH22	1.65	0.45
25:1H:432:A:H2'	25:1H:433:C:C6	2.51	0.45
25:1H:577:G:O2'	25:1H:1254:A:OP1	2.33	0.45
25:1H:613:U:H5'	25:1H:616:A:N6	2.32	0.45
25:1H:1641:A:H2'	25:1H:1642:G:O4'	2.17	0.45
25:1H:2115:G:N1	25:1H:2116:G:O6	2.50	0.45
31:41:60:LEU:HD22	31:41:68:PRO:HB3	1.99	0.45
31:41:107:LEU:HD21	31:41:178:PHE:CE1	2.52	0.45
39:98:36:THR:OG1	39:98:37:THR:N	2.50	0.45
41:B8:26:ASP:O	41:B8:49:VAL:HG12	2.16	0.45
46:G8:30:VAL:HG22	46:G8:37:VAL:HG12	1.98	0.45
46:G8:55:TYR:CE2	46:G8:61:ILE:HD11	2.52	0.45
1:1G:619:U:N3	4:32:134:ASP:OD1	2.40	0.45
1:1G:946:A:H2'	1:1G:947:G:C8	2.51	0.45
1:1G:1134:G:C2	1:1G:1135:U:H1'	2.52	0.45
1:1G:1149:C:OP1	9:82:9:ARG:HD3	2.17	0.45
4:32:57:ARG:HG3	4:32:202:LEU:HB3	1.98	0.45
9:82:79:LEU:O	9:82:83:ARG:N	2.50	0.45
11:2A:85:ARG:HA	11:2A:112:THR:HG23	1.99	0.45
25:14:2585:U:HO2'	25:14:2586:C:P	2.40	0.45
26:1J:80:U:H2'	26:1J:81:G:N2	2.31	0.45
31:49:18:GLU:OE1	31:49:22:ARG:NH1	2.50	0.45
31:49:67:LYS:HD3	52:I5:5:ILE:CG1	2.47	0.45
37:35:41:ARG:H	37:35:41:ARG:HG2	1.45	0.45
40:65:19:LYS:O	40:65:20:ARG:HD3	2.17	0.45
41:75:52:ILE:O	41:75:98:LYS:NZ	2.39	0.45
1:13:977:A:O2'	1:13:981:U:N3	2.36	0.44
1:13:1189:C:H5''	3:2E:5:ILE:HD12	1.97	0.44
1:13:1296:C:H5'	13:4I:14:ARG:HD3	1.99	0.44
3:2E:36:ASP:HA	3:2E:39:ILE:HD12	1.99	0.44
4:3E:9:CYS:HB3	4:3E:32:ALA:CB	2.47	0.44
16:7I:13:HIS:C	16:7I:15:PRO:HD3	2.43	0.44
16:7I:75:ARG:HA	16:7I:80:PHE:HB2	1.99	0.44
18:9I:30:ASP:OD2	18:9I:33:ASP:HB2	2.18	0.44
25:1H:270(Q):C:H2'	25:1H:270(R):G:O4'	2.16	0.44
25:1H:1226:G:OP1	43:D8:69:LYS:NZ	2.37	0.44
25:1H:1231:G:H2'	25:1H:1232:G:C8	2.52	0.44
25:1H:2208:U:H1'	28:11:151:LYS:HE2	1.98	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:16:74:U:O4	26:16:102:G:N2	2.47	0.44
40:A8:57:LYS:H	40:A8:57:LYS:HD3	1.82	0.44
45:F8:15:GLU:CD	45:F8:15:GLU:H	2.23	0.44
45:F8:53:LYS:H	45:F8:82:GLN:HB3	1.83	0.44
47:H8:72:ARG:NH2	47:H8:97:GLU:O	2.50	0.44
50:K8:6:VAL:O	50:K8:10:LEU:HB2	2.17	0.44
4:32:154:ASN:OD1	4:32:154:ASN:N	2.50	0.44
15:6A:61:GLY:O	15:6A:65:ARG:HG3	2.17	0.44
17:8A:55:ASP:HB3	17:8A:76:LEU:HD13	1.98	0.44
20:BA:33:ILE:HD11	20:BA:62:LEU:C	2.43	0.44
25:14:107:C:H2'	25:14:108:U:H6	1.82	0.44
25:14:363(B):G:H2'	25:14:363(C):G:C8	2.51	0.44
25:14:817:C:O2'	25:14:839:U:H5''	2.17	0.44
25:14:2249:U:N3	25:14:2253:G:OP2	2.36	0.44
28:19:223:GLY:HA2	28:19:226:MET:HG3	1.98	0.44
31:49:104:GLU:O	31:49:108:ASN:ND2	2.41	0.44
33:69:102:SER:HB3	33:69:107:VAL:O	2.17	0.44
33:69:113:ARG:HB3	33:69:131:LYS:HD3	1.99	0.44
40:65:84:GLN:HG3	40:65:109:GLY:HA2	1.99	0.44
41:75:111:ARG:C	41:75:113:LYS:H	2.23	0.44
47:D5:10:ARG:NH1	47:D5:26:GLY:O	2.37	0.44
1:13:373:A:H2'	1:13:374:A:H8	1.81	0.44
1:13:436:C:H2'	1:13:437:U:O4'	2.17	0.44
1:13:789:U:O2'	1:13:791:G:N7	2.46	0.44
1:13:827:U:C2	1:13:874:G:N2	2.85	0.44
6:5E:4:TYR:HD1	6:5E:92:LYS:HA	1.82	0.44
8:7E:20:TYR:CE2	8:7E:75:ARG:HB3	2.51	0.44
8:7E:73:ASP:OD1	8:7E:75:ARG:NH2	2.50	0.44
12:3I:21:VAL:HG13	12:3I:95:TYR:CE1	2.52	0.44
13:4I:117:VAL:HG22	13:4I:118:ALA:H	1.81	0.44
22:1K:38:A:H2'	22:1K:39:A:O4'	2.17	0.44
25:1H:902:C:H2'	25:1H:903:C:H6	1.81	0.44
25:1H:924:C:H2'	25:1H:925:C:C6	2.52	0.44
25:1H:1059:G:H3'	25:1H:1060:U:C5'	2.47	0.44
25:1H:2530:A:O2'	25:1H:2532:G:OP2	2.28	0.44
25:1H:2846:G:H2'	25:1H:2847:U:O4'	2.17	0.44
28:11:131:LEU:O	28:11:190:TYR:HA	2.17	0.44
29:21:52:LEU:HD23	29:21:76:ARG:HD3	1.98	0.44
39:98:74:LYS:C	39:98:76:VAL:H	2.25	0.44
50:K8:6:VAL:HA	50:K8:9:GLN:HB3	1.98	0.44
1:1G:690:G:H2'	1:1G:691:G:O4'	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:885:G:H1	1:1G:912:C:H42	1.65	0.44
1:1G:1235:U:H5''	21:1B:3:LYS:HE2	1.99	0.44
1:1G:1298:C:O5'	7:62:114:ARG:NH2	2.50	0.44
1:1G:1372:U:P	9:82:71:SER:HG	2.40	0.44
3:22:157:ILE:HD12	3:22:164:ARG:HB3	1.99	0.44
4:32:176:LEU:HG	4:32:178:VAL:HG22	1.99	0.44
9:82:112:LYS:HD2	9:82:113:LYS:N	2.32	0.44
13:4A:48:LEU:O	13:4A:49:THR:C	2.58	0.44
25:14:1639:U:O2'	25:14:2699:C:H4'	2.17	0.44
25:14:2185:C:H2'	25:14:2186:G:C8	2.52	0.44
25:14:2298:A:H2'	25:14:2299:G:O4'	2.17	0.44
25:14:2340:G:H2'	25:14:2341:G:H8	1.82	0.44
25:14:2345:G:N3	25:14:2381:C:H2'	2.32	0.44
25:14:2765:A:H2	25:14:2766:G:O4'	1.99	0.44
38:45:35:VAL:N	38:45:130:LYS:O	2.46	0.44
1:13:667:G:H4'	15:6I:51:HIS:CE1	2.52	0.44
1:13:895:G:H2'	1:13:896:C:H6	1.82	0.44
1:13:938:A:HO2'	1:13:1376:U:HO2'	1.60	0.44
1:13:1328:C:H2'	1:13:1329:A:O4'	2.17	0.44
9:8E:83:ARG:O	9:8E:86:VAL:HG12	2.18	0.44
9:8E:113:LYS:H	9:8E:119:ALA:HA	1.83	0.44
13:4I:84:ILE:HD11	19:AI:65:ASN:ND2	2.32	0.44
25:1H:451:C:H4'	30:31:52:LYS:HZ2	1.82	0.44
25:1H:1778:U:H2'	25:1H:1784:A:N6	2.32	0.44
25:1H:2481:G:O2'	25:1H:2482:G:OP2	2.28	0.44
25:1H:2646:C:H2'	25:1H:2647:U:O4'	2.16	0.44
35:58:21:LYS:HB2	35:58:26:LEU:HD22	2.00	0.44
42:C8:34:LYS:HE2	42:C8:34:LYS:HA	1.99	0.44
42:C8:89:GLU:HG2	43:D8:50:PRO:HB3	1.99	0.44
45:F8:56:THR:HG22	45:F8:79:ALA:HB2	2.00	0.44
47:H8:156:LYS:HG2	47:H8:157:LEU:N	2.32	0.44
48:I8:49:LYS:HB2	48:I8:80:HIS:HB3	1.99	0.44
1:1G:429:U:H1'	1:1G:430:A:H5''	1.99	0.44
1:1G:1118:C:N4	1:1G:1156:G:H21	2.15	0.44
1:1G:1207:2MG:H2'	1:1G:1208:C:C6	2.53	0.44
5:42:78:HIS:HB3	8:72:107:LEU:HD13	1.99	0.44
7:62:150:ALA:HB1	11:2A:57:THR:HG21	2.00	0.44
8:72:84:ARG:HG2	8:72:86:ILE:HD13	1.99	0.44
9:82:40:LEU:HB3	9:82:43:ALA:HB2	2.00	0.44
13:4A:111:LYS:H	13:4A:111:LYS:HG3	1.61	0.44
20:BA:34:LYS:O	20:BA:38:LYS:HB2	2.17	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:717:G:H2'	25:14:718:A:O4'	2.16	0.44
25:14:1418:G:H8	25:14:1418:G:O5'	1.99	0.44
25:14:2108:C:H2'	25:14:2109:U:O4'	2.18	0.44
25:14:2696:U:H2'	25:14:2697:G:C8	2.52	0.44
25:14:2882:A:O5'	39:55:96:ARG:HG2	2.18	0.44
26:1J:32:C:H2'	26:1J:33:G:C8	2.52	0.44
31:49:174:GLU:HG2	31:49:180:PHE:CD2	2.53	0.44
35:15:56:ASN:O	35:15:56:ASN:ND2	2.45	0.44
52:I5:12:ALA:HB1	52:I5:24:THR:H	1.83	0.44
1:13:170:U:H2'	1:13:171:A:H8	1.83	0.44
1:13:391:G:O3'	16:7I:8:ARG:NH2	2.49	0.44
1:13:451:A:H4'	1:13:452:A:O4'	2.17	0.44
1:13:803:G:H2'	1:13:804:U:O4'	2.18	0.44
1:13:857:C:H2'	1:13:858:G:O4'	2.18	0.44
1:13:984:C:H2'	1:13:985:C:C6	2.52	0.44
1:13:1055:A:H62	1:13:1200:C:N4	2.16	0.44
1:13:1329:A:H5'	13:4I:29:ARG:HE	1.81	0.44
4:3E:12:CYS:SG	4:3E:19:LEU:HB2	2.57	0.44
5:4E:100:VAL:HG22	5:4E:118:ILE:HG22	1.98	0.44
13:4I:68:GLY:HA3	31:41:116:ASP:OD2	2.17	0.44
23:2K:24:C:H2'	23:2K:25:U:H6	1.81	0.44
25:1H:296:C:H2'	25:1H:297:C:C6	2.53	0.44
25:1H:519:U:H2'	25:1H:520:G:H8	1.82	0.44
25:1H:768:G:O2'	25:1H:1379:A:N6	2.50	0.44
25:1H:973:A:H8	25:1H:973:A:OP1	2.00	0.44
25:1H:1783:A:H5'	25:1H:2608:G:H4'	2.00	0.44
25:1H:1872:A:H5'	25:1H:1878:G:OP2	2.18	0.44
25:1H:1945:G:O6	25:1H:1960:A:N6	2.50	0.44
25:1H:2078:C:H2'	25:1H:2079:U:C6	2.52	0.44
25:1H:2138:C:O2	25:1H:2153:G:N1	2.43	0.44
29:21:18:ASP:N	29:21:18:ASP:OD1	2.49	0.44
29:21:188:VAL:HG13	29:21:189:PRO:HD2	1.98	0.44
31:41:111:LEU:O	31:41:114:ILE:HG12	2.17	0.44
31:41:145:THR:O	31:41:146:TYR:HB3	2.16	0.44
54:O8:25:LYS:HB3	56:Q8:35:GLN:HG3	1.99	0.44
1:1G:823:G:H2'	1:1G:824:C:C6	2.52	0.44
3:22:12:LEU:HD12	14:5A:56:VAL:O	2.17	0.44
13:4A:65:LYS:HE2	52:I5:52:THR:HG22	1.99	0.44
20:BA:10:LEU:HA	20:BA:13:LEU:HB2	2.00	0.44
25:14:896:A:H3'	25:14:897:C:H5'	1.99	0.44
25:14:981:A:N1	25:14:2027:G:O2'	2.39	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:1071:G:N3	25:14:1089:G:O2'	2.42	0.44
25:14:1170:G:O6	25:14:1178:C:N4	2.50	0.44
25:14:2322:A:H2'	25:14:2323:G:O4'	2.17	0.44
25:14:2686:G:H5'	62:14:3654:HOH:O	2.16	0.44
25:14:2688:U:H1'	25:14:2721:A:H61	1.81	0.44
25:14:2867:G:O2'	25:14:2868:A:C8	2.71	0.44
40:65:108:GLY:H	40:65:110:LEU:HG	1.82	0.44
46:C5:102:CYS:SG	46:C5:105:ALA:HB2	2.56	0.44
47:D5:30:ASN:HB3	47:D5:90:VAL:HB	1.98	0.44
49:F5:46:LEU:HA	49:F5:46:LEU:HD12	1.67	0.44
54:K5:25:LYS:HD3	56:M5:34:TRP:HE1	1.82	0.44
1:13:938:A:O2'	1:13:1376:U:O2'	2.24	0.44
1:13:1004:A:C2	1:13:1024:G:C8	3.04	0.44
1:13:1129:C:C5'	1:13:1130:A:H5'	2.35	0.44
1:13:1327:C:H2'	1:13:1328:C:C6	2.53	0.44
2:1E:115:LEU:HB2	2:1E:145:LEU:HD12	1.99	0.44
2:1E:142:LEU:O	2:1E:146:GLN:N	2.29	0.44
3:2E:16:ARG:HE	3:2E:54:ARG:HH21	1.65	0.44
8:7E:41:ARG:HB3	8:7E:41:ARG:HH11	1.82	0.44
10:1I:6:ILE:HG22	10:1I:98:ILE:HG23	1.98	0.44
19:AI:80:TYR:O	19:AI:81:ARG:C	2.59	0.44
20:BI:56:MET:SD	20:BI:88:VAL:HG11	2.58	0.44
25:1H:270(L):U:H2'	33:61:50:ARG:HD3	1.99	0.44
25:1H:792:G:H5''	25:1H:793:A:H5'	1.99	0.44
25:1H:970:C:H2'	25:1H:971:C:C6	2.52	0.44
25:1H:1055:G:H22	25:1H:1104:C:H42	1.64	0.44
25:1H:1404:C:O2'	25:1H:1405:U:H5'	2.18	0.44
25:1H:1711:C:H2'	25:1H:1712:C:H6	1.82	0.44
25:1H:1812:A:O4'	28:11:45:ASN:ND2	2.50	0.44
25:1H:1919:A:H5''	25:1H:1920:OMC:OP2	2.18	0.44
25:1H:2210:G:N3	25:1H:2210:G:H2'	2.32	0.44
25:1H:2396:G:O5'	49:J8:25:LYS:HE2	2.17	0.44
25:1H:2454:G:H1'	62:1H:3876:HOH:O	2.17	0.44
31:4I:118:ARG:NH2	52:M8:43:TYR:HB2	2.32	0.44
40:A8:65:VAL:O	40:A8:69:VAL:HG12	2.16	0.44
48:I8:77:ARG:HB2	48:I8:77:ARG:HH11	1.82	0.44
1:1G:1004:A:C4	1:1G:1025:U:N3	2.85	0.44
3:22:130:VAL:O	3:22:134:ILE:HG12	2.16	0.44
4:32:173:TRP:CD2	4:32:189:PRO:HB3	2.53	0.44
9:82:15:ALA:HB2	9:82:65:VAL:HG23	1.99	0.44
9:82:95:LYS:HE2	9:82:95:LYS:HB2	1.90	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
57:2L:76:C:H3'	57:2L:77:A:C5'	2.45	0.44
25:14:41:C:H2'	25:14:43:G:C8	2.52	0.44
25:14:1142(A):A:C5	25:14:1144:G:C5	3.06	0.44
25:14:1308:A:H2'	25:14:1309:G:O4'	2.18	0.44
25:14:1813:G:H4'	28:19:43:ARG:O	2.18	0.44
25:14:2544:G:H2'	25:14:2545:G:C8	2.53	0.44
26:1J:46:A:H2'	26:1J:47:C:O4'	2.16	0.44
31:49:43:LEU:O	31:49:88:ILE:HG13	2.18	0.44
31:49:64:THR:HG23	31:49:66:GLN:N	2.30	0.44
36:25:113:LYS:H	36:25:113:LYS:HD2	1.82	0.44
43:95:70:ILE:HG22	43:95:72:VAL:HG23	1.98	0.44
46:C5:91:GLU:O	46:C5:93:GLY:N	2.50	0.44
47:D5:152:ALA:C	47:D5:154:ASP:N	2.76	0.44
1:13:9:G:H2'	1:13:10:A:C8	2.53	0.44
1:13:936:C:O2	1:13:1382:C:N4	2.50	0.44
1:13:1007:C:H2'	1:13:1008:C:O4'	2.17	0.44
1:13:1133:G:H2'	1:13:1134:G:H8	1.82	0.44
1:13:1243:C:P	21:1F:10:ARG:HH22	2.41	0.44
1:13:1352:C:OP1	21:1F:3:LYS:NZ	2.28	0.44
2:1E:7:VAL:HG21	2:1E:217:ARG:NH1	2.33	0.44
9:8E:49:PRO:HB3	9:8E:96:LEU:HD11	1.99	0.44
17:8I:101:ARG:HA	17:8I:101:ARG:HE	1.82	0.44
21:1F:6:ARG:CZ	21:1F:15:ARG:HE	2.30	0.44
22:3K:31:G:H1	22:3K:41:C:H42	1.65	0.44
25:1H:161:U:H3'	25:1H:162:U:C5'	2.48	0.44
25:1H:270(E):G:H2'	25:1H:270(F):U:O4'	2.18	0.44
25:1H:533:G:H5'	42:C8:24:TYR:CE1	2.53	0.44
25:1H:732:C:OP2	62:1H:3693:HOH:O	2.20	0.44
25:1H:1932:A:H2'	25:1H:1933:G:O4'	2.17	0.44
25:1H:2046:G:O5'	53:N8:19:ARG:HA	2.17	0.44
25:1H:2208:U:O2'	28:11:151:LYS:HG2	2.17	0.44
25:1H:2567:G:H2'	25:1H:2568:C:H6	1.81	0.44
25:1H:2667:C:H1'	32:51:109:PHE:HD2	1.82	0.44
31:41:55:LYS:O	31:41:59:GLU:HB2	2.16	0.44
37:78:57:THR:HG22	37:78:58:THR:N	2.32	0.44
40:A8:25:ARG:HB3	40:A8:40:ILE:HB	1.99	0.44
41:B8:51:ARG:HG2	41:B8:52:ILE:N	2.33	0.44
43:D8:1:MET:HE3	43:D8:1:MET:HB3	1.76	0.44
43:D8:6:LYS:HE3	43:D8:6:LYS:HB2	1.77	0.44
43:D8:37:VAL:HG22	43:D8:56:SER:HA	2.00	0.44
1:1G:56:U:H2'	1:1G:57:G:C8	2.52	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:280:C:N3	17:8A:39:SER:OG	2.45	0.44
1:1G:446:G:O6	62:1G:1813:HOH:O	2.20	0.44
1:1G:767:A:H2'	1:1G:768:A:C8	2.53	0.44
1:1G:1092:A:O3'	7:62:4:ARG:NH1	2.50	0.44
1:1G:1165:C:N4	1:1G:1166:G:O6	2.51	0.44
1:1G:1353:G:H1	1:1G:1369:C:N4	2.14	0.44
1:1G:1410:G:H2'	1:1G:1411:C:C6	2.53	0.44
3:22:56:ASP:CG	3:22:69:HIS:HE1	2.25	0.44
3:22:155:GLY:HA2	3:22:163:ALA:HB1	1.99	0.44
4:32:155:LEU:O	4:32:158:ILE:HG22	2.18	0.44
5:42:11:ILE:HG22	5:42:12:LEU:N	2.31	0.44
8:72:83:ILE:HB	8:72:137:VAL:HG12	2.00	0.44
9:82:81:ILE:HG22	9:82:85:LEU:HD23	1.98	0.44
13:4A:7:VAL:HG23	13:4A:9:ILE:H	1.81	0.44
25:14:796:C:H2'	25:14:797:C:C6	2.52	0.44
29:29:24:THR:HG21	29:29:188:VAL:HG23	2.00	0.44
31:49:120:LEU:HD11	31:49:133:LEU:HD13	1.99	0.44
35:15:91:LEU:O	35:15:95:PRO:HB3	2.18	0.44
36:25:63:VAL:HG22	36:25:84:ALA:HA	1.99	0.44
41:75:4:GLY:O	41:75:7:ILE:HD13	2.18	0.44
43:95:40:LEU:HD22	43:95:40:LEU:HA	1.85	0.44
47:D5:161:VAL:HG22	47:D5:163:LEU:HD22	1.99	0.44
1:13:390:C:O2'	16:7I:28:ARG:NH2	2.51	0.44
1:13:890:G:O2'	1:13:906:G:O6	2.24	0.44
1:13:1015:A:H2'	1:13:1016:A:O4'	2.17	0.44
2:1E:47:THR:O	2:1E:51:LEU:N	2.50	0.44
19:AI:3:ARG:NH1	19:AI:10:PHE:HA	2.33	0.44
25:1H:935:C:H2'	25:1H:936:C:C6	2.52	0.44
25:1H:1410:G:H1	25:1H:1592:C:H42	1.66	0.44
25:1H:1680:U:O2'	25:1H:1763:G:N7	2.41	0.44
25:1H:2528:U:O2'	25:1H:2530:A:OP1	2.35	0.44
25:1H:2721:A:H2'	25:1H:2722:G:O4'	2.17	0.44
34:38:112:LEU:HD11	34:38:118:THR:OG1	2.18	0.44
37:78:112:LEU:HD13	37:78:127:ALA:HB2	2.00	0.44
40:A8:76:LYS:HE2	40:A8:76:LYS:HB3	1.81	0.44
46:G8:39:VAL:O	46:G8:42:VAL:HG22	2.17	0.44
52:M8:40:HIS:CE1	52:M8:44:THR:HG22	2.53	0.44
1:1G:464:G:O6	1:1G:466:C:H5'	2.17	0.44
1:1G:755:G:H2'	1:1G:756:C:C6	2.53	0.44
2:12:35:GLU:HG2	2:12:36:ARG:N	2.32	0.44
3:22:47:LEU:HD23	3:22:52:LEU:HD13	2.00	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:22:120:VAL:O	3:22:123:GLN:N	2.51	0.44
5:42:92:LYS:HB3	5:42:119:LEU:HB2	1.98	0.44
12:3A:55:VAL:O	12:3A:62:GLU:HA	2.18	0.44
13:4A:53:VAL:HG12	13:4A:57:ARG:HH11	1.81	0.44
16:7A:69:THR:O	16:7A:73:LEU:HG	2.17	0.44
22:3L:23:G:HO2'	22:3L:24:C:P	2.40	0.44
25:14:27:G:N2	25:14:512:G:O2'	2.51	0.44
25:14:902:C:H2'	25:14:903:C:C6	2.52	0.44
25:14:1015:G:H2'	25:14:1016:G:C8	2.53	0.44
25:14:1024:G:H8	25:14:1024:G:O5'	2.00	0.44
25:14:1144:G:C6	25:14:1145:C:C4	3.05	0.44
25:14:1171:G:H8	25:14:1173:G:H21	1.66	0.44
25:14:1266:G:O5'	44:A5:15:ARG:NH2	2.51	0.44
25:14:1401:G:H2'	25:14:1402:C:O4'	2.17	0.44
25:14:1582:C:O2'	25:14:1586:A:H8	2.00	0.44
25:14:2113:U:H3'	25:14:2114:A:H4'	1.99	0.44
25:14:2490:G:H2'	25:14:2490:G:N3	2.32	0.44
32:59:4:ILE:HD13	32:59:4:ILE:H	1.82	0.44
35:15:71:ILE:H	35:15:71:ILE:HG13	1.68	0.44
35:15:111:PRO:HA	35:15:114:ARG:NH1	2.33	0.44
40:65:72:ALA:O	40:65:76:LYS:HG3	2.17	0.44
1:13:693:G:H2'	1:13:694:A:C8	2.52	0.44
1:13:1321:C:C4	1:13:1322:C:C4	3.06	0.44
2:1E:27:LYS:NZ	2:1E:193:ASP:OD2	2.43	0.44
3:2E:19:GLU:OE2	3:2E:40:ARG:NE	2.46	0.44
13:4I:4:ILE:HG22	13:4I:5:ALA:H	1.83	0.44
18:9I:38:GLU:H	18:9I:38:GLU:HG2	1.52	0.44
25:1H:248:G:H5''	25:1H:386:G:N2	2.33	0.44
25:1H:306:U:H2'	25:1H:307:G:O4'	2.17	0.44
25:1H:1040:C:H2'	25:1H:1041:C:C6	2.52	0.44
25:1H:1062:G:H2'	25:1H:1063:G:H8	1.83	0.44
25:1H:1178:C:O2'	25:1H:1179:C:O4'	2.36	0.44
25:1H:2061:G:H2'	25:1H:2501:C:O2'	2.17	0.44
25:1H:2062:A:H2'	25:1H:2062:A:N3	2.32	0.44
27:71:202:GLU:C	27:71:204:ALA:H	2.25	0.44
29:21:116:VAL:HG13	29:21:122:PHE:HB2	1.99	0.44
32:51:51:ARG:HG2	32:51:52:VAL:H	1.83	0.44
42:C8:95:LEU:HD13	43:D8:4:ILE:CG2	2.42	0.44
47:H8:101:PRO:HA	47:H8:123:ASP:HB3	1.98	0.44
1:1G:127:G:O2'	17:8A:2:PRO:O	2.36	0.44
1:1G:192:U:H2'	1:1G:193:C:C6	2.53	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:254:G:N2	17:8A:16:GLN:OE1	2.26	0.44
1:1G:401:C:O2'	1:1G:621:A:N3	2.42	0.44
1:1G:517:G:N2	1:1G:533:A:OP2	2.39	0.44
1:1G:806:C:H2'	1:1G:807:A:C8	2.53	0.44
1:1G:1127:G:H21	1:1G:1147:C:H41	1.66	0.44
1:1G:1305:G:N2	1:1G:1331:G:H2'	2.33	0.44
3:22:150:LYS:HB3	3:22:201:TYR:HB2	2.00	0.44
21:1B:6:ARG:HG2	21:1B:15:ARG:HH22	1.83	0.44
22:3L:20:G:N1	25:14:2112:G:O4'	2.50	0.44
22:3L:23:G:O2'	22:3L:24:C:OP1	2.30	0.44
25:14:422:A:H2'	25:14:423:A:C8	2.53	0.44
25:14:438:G:H2'	25:14:439:G:C8	2.53	0.44
25:14:463:G:N2	25:14:466:A:OP2	2.30	0.44
25:14:556:G:H8	25:14:556:G:O5'	2.00	0.44
25:14:632:A:H2'	25:14:633:A:C8	2.52	0.44
25:14:1043:C:N4	25:14:1112:G:H22	2.15	0.44
25:14:1197:G:H22	25:14:1249:U:H2'	1.82	0.44
25:14:1759:A:H5''	25:14:2715:C:O2'	2.17	0.44
25:14:2082:A:H2'	25:14:2083:G:O4'	2.17	0.44
25:14:2392:A:OP2	25:14:2422:A:N6	2.51	0.44
31:49:64:THR:HA	31:49:102:PHE:HD2	1.83	0.44
32:59:102:ALA:HB1	32:59:115:VAL:N	2.33	0.44
36:25:69:ILE:HD12	36:25:77:ILE:O	2.18	0.44
40:65:27:SER:HA	40:65:88:ASP:HB3	2.00	0.44
40:65:87:PHE:HB2	40:65:112:PHE:CE2	2.53	0.44
48:E5:50:ASN:HB3	48:E5:81:VAL:HB	1.99	0.44
1:13:372:C:N4	1:13:389:A:H62	2.14	0.44
1:13:795:C:H1'	1:13:1506:U:C5	2.53	0.44
1:13:1145:C:H4'	1:13:1146:A:H8	1.83	0.44
1:13:1223:C:P	19:AI:78:ARG:HH12	2.41	0.44
1:13:1376:U:H2'	1:13:1377:A:H8	1.82	0.44
2:1E:24:TRP:HA	2:1E:190:THR:O	2.18	0.44
22:1K:11:A:H2'	22:1K:12:G:O4'	2.18	0.44
25:1H:195:A:H61	25:1H:198:C:H3'	1.83	0.44
25:1H:330:A:O2'	25:1H:331:A:H8	2.01	0.44
25:1H:528:A:H8	25:1H:528:A:H3'	1.82	0.44
25:1H:1019:U:OP1	25:1H:1035:U:O2'	2.30	0.44
25:1H:1038:C:H2'	25:1H:1039:G:O4'	2.18	0.44
25:1H:1054:A:H5'	25:1H:1055:G:OP2	2.17	0.44
25:1H:1101:U:H2'	25:1H:1102:C:H6	1.83	0.44
25:1H:1510:A:H2'	25:1H:1511:A:O4'	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:2295:C:OP2	40:A8:10:ARG:HG2	2.18	0.44
25:1H:2766:G:N3	25:1H:2766:G:H2'	2.32	0.44
29:21:144:ARG:O	29:21:146:THR:N	2.51	0.44
34:38:131:MET:HG3	34:38:132:ASP:N	2.33	0.44
37:78:125:VAL:O	37:78:145:PRO:HD2	2.18	0.44
38:88:132:VAL:HG11	47:H8:81:ARG:HH11	1.83	0.44
40:A8:24:LEU:HD12	40:A8:41:ASP:HB2	2.00	0.44
54:O8:9:LEU:HB3	54:O8:26:ASN:O	2.17	0.44
1:1G:67:C:H2'	1:1G:68:G:H8	1.82	0.44
1:1G:736:C:H2'	1:1G:737:A:H8	1.79	0.44
1:1G:832:C:H42	1:1G:854:G:H1	1.66	0.44
4:32:36:ARG:HD2	4:32:38:TYR:CZ	2.53	0.44
24:4L:12:A:OP1	24:4L:12:A:H4'	2.16	0.44
25:14:671:C:H2'	25:14:672:C:H6	1.82	0.44
25:14:947:G:H2'	25:14:948:G:C8	2.53	0.44
25:14:1030:G:H2'	25:14:1031:G:C8	2.53	0.44
25:14:1171:G:H1	25:14:1178:C:H42	1.64	0.44
25:14:1180:C:H5'	25:14:1181:C:OP2	2.18	0.44
25:14:1375:C:H2'	25:14:1376:C:C6	2.53	0.44
25:14:2022:U:O2'	25:14:2023:G:OP2	2.31	0.44
25:14:2079:U:O3'	49:F5:35:THR:HB	2.18	0.44
25:14:2098:U:H2'	25:14:2099:U:O4'	2.18	0.44
25:14:2385:C:OP2	25:14:2385:C:H4'	2.18	0.44
25:14:2467:C:H2'	25:14:2468:G:O4'	2.17	0.44
25:14:2844:G:H3'	25:14:2845:G:C8	2.49	0.44
30:39:64:ILE:H	30:39:64:ILE:HG13	1.48	0.44
31:49:121:ASN:HB2	31:49:181:ARG:HH22	1.83	0.44
31:49:129:GLY:O	31:49:161:THR:N	2.51	0.44
36:25:71:ARG:HB2	36:25:73:ASP:OD1	2.18	0.44
38:45:48:GLU:HA	38:45:51:ARG:HB3	1.99	0.44
41:75:39:ARG:HG2	41:75:40:THR:N	2.31	0.44
42:85:99:ALA:HB2	42:85:106:PHE:CG	2.53	0.44
46:C5:75:ILE:HD13	46:C5:75:ILE:HA	1.81	0.44
46:C5:79:CYS:HB3	46:C5:80:GLY:O	2.18	0.44
47:D5:5:LEU:HD21	47:D5:44:PHE:HA	2.00	0.44
54:K5:10:LEU:HD12	54:K5:24:GLU:HG2	1.99	0.44
1:13:21:G:H2'	1:13:22:G:C8	2.53	0.43
1:13:551:U:H5'	12:3I:116:LYS:HE2	2.00	0.43
1:13:825:G:H2'	1:13:826:C:C6	2.53	0.43
1:13:1303:C:H2'	1:13:1304:G:O4'	2.18	0.43
4:3E:15:GLU:OE1	4:3E:59:ARG:NE	2.42	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:2K:21:H2U:H2'	23:2K:22:A:H5'	1.99	0.43
25:1H:528:A:H3'	25:1H:528:A:C8	2.53	0.43
25:1H:817:C:O2'	25:1H:839:U:H5''	2.18	0.43
25:1H:1482:U:H5'	25:1H:1483:G:OP2	2.16	0.43
25:1H:1871:A:H8	25:1H:1871:A:OP2	2.00	0.43
25:1H:2307:G:H1'	25:1H:2308:G:C2	2.52	0.43
25:1H:2494:G:H2'	25:1H:2495:G:C8	2.50	0.43
26:16:9:G:OP1	40:A8:15:ARG:NH1	2.51	0.43
32:51:73:ALA:O	32:51:77:LYS:N	2.37	0.43
40:A8:71:ARG:HH12	40:A8:106:ARG:NH2	2.16	0.43
43:D8:1:MET:HG3	43:D8:43:GLU:HG2	2.00	0.43
1:1G:688:G:H2'	1:1G:689:C:H6	1.82	0.43
1:1G:1137:C:H5'	1:1G:1138:G:N3	2.33	0.43
1:1G:1229:A:OP2	13:4A:114:ARG:HD3	2.17	0.43
1:1G:1299:A:C6	1:1G:1301:U:C2	3.06	0.43
1:1G:1327:C:OP1	21:1B:20:LYS:HB3	2.18	0.43
3:22:20:SER:HB2	3:22:40:ARG:HH21	1.83	0.43
3:22:36:ASP:HA	3:22:39:ILE:HD12	1.99	0.43
4:32:119:GLN:HG2	4:32:123:HIS:CE1	2.53	0.43
4:32:201:GLN:NE2	5:42:99:GLY:HA2	2.33	0.43
8:72:30:ARG:HG3	8:72:31:PHE:N	2.32	0.43
19:AA:52:TYR:HB2	19:AA:57:HIS:HE1	1.83	0.43
25:14:78:A:H2'	25:14:79:G:C8	2.53	0.43
25:14:589:C:H2'	25:14:590:A:H8	1.82	0.43
25:14:779:U:OP1	28:19:49:ILE:HG13	2.18	0.43
25:14:858:U:O2	25:14:2268:A:H2'	2.18	0.43
25:14:1652:A:HO2'	25:14:1653:G:P	2.41	0.43
25:14:2261:C:OP1	48:E5:17:GLN:HB2	2.18	0.43
25:14:2361:A:OP1	56:M5:26:LYS:HD3	2.18	0.43
26:1J:109:G:H2'	26:1J:110:G:C8	2.52	0.43
31:49:173:LEU:HB3	31:49:178:PHE:CD2	2.53	0.43
33:69:9:LEU:HD13	33:69:9:LEU:H	1.81	0.43
35:15:54:VAL:HB	35:15:122:VAL:HG22	1.99	0.43
47:D5:29:TYR:CE2	47:D5:87:ASP:HB3	2.53	0.43
47:D5:138:GLU:H	47:D5:138:GLU:HG2	1.46	0.43
47:D5:153:SER:C	47:D5:155:LEU:H	2.25	0.43
1:13:160:A:N6	1:13:347:G:H1'	2.33	0.43
1:13:190:G:O2'	1:13:191(A):G:OP2	2.32	0.43
1:13:503:C:O2	1:13:542:G:N2	2.40	0.43
1:13:792:A:H1'	1:13:793:U:OP2	2.18	0.43
1:13:1118:C:H1'	1:13:1179:A:C8	2.53	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1E:114:ARG:O	2:1E:118:LEU:N	2.43	0.43
2:1E:130:ARG:O	2:1E:135:GLN:HG3	2.18	0.43
5:4E:129:ILE:O	5:4E:133:TYR:HB2	2.18	0.43
7:6E:84:ASN:N	7:6E:84:ASN:OD1	2.51	0.43
8:7E:103:VAL:HG21	8:7E:110:ALA:HB2	1.99	0.43
9:8E:7:THR:H	9:8E:83:ARG:HD2	1.83	0.43
15:6I:68:ARG:O	15:6I:72:ARG:HB2	2.18	0.43
20:BI:25:ARG:HG2	20:BI:29:LYS:HE3	2.00	0.43
22:3K:68:C:H2'	22:3K:69:C:C6	2.53	0.43
25:1H:41:C:H42	25:1H:438:G:H1	1.67	0.43
25:1H:407:G:H2'	25:1H:408:G:C8	2.53	0.43
25:1H:571:A:O5'	25:1H:2030:A:N6	2.47	0.43
25:1H:956:G:H2'	25:1H:957:A:H2'	2.00	0.43
25:1H:1639:U:C2'	25:1H:1640:C:H5''	2.48	0.43
25:1H:2776:A:H3'	25:1H:2776:A:OP1	2.17	0.43
62:1H:3995:HOH:O	37:78:37:GLY:HA3	2.17	0.43
33:61:85:GLU:OE1	33:61:86:THR:OG1	2.32	0.43
34:38:64:LYS:H	34:38:64:LYS:HG2	1.58	0.43
37:78:124:LYS:HA	37:78:143:GLY:O	2.17	0.43
45:F8:36:LYS:HG2	45:F8:54:VAL:HG12	2.01	0.43
46:G8:5:MET:HE1	46:G8:32:PRO:HA	1.99	0.43
1:1G:717:C:O2'	11:2A:117:ASN:HB2	2.18	0.43
1:1G:973:G:H1'	10:1A:55:LYS:HE2	1.99	0.43
1:1G:1243:C:H5''	21:1B:8:THR:HG21	1.99	0.43
1:1G:1282:C:H2'	1:1G:1283:G:O4'	2.18	0.43
1:1G:1346:A:H5''	9:82:120:ARG:HH12	1.82	0.43
1:1G:1481:U:H2'	1:1G:1482:G:C8	2.53	0.43
2:12:15:VAL:HG11	2:12:213:LEU:HD22	2.00	0.43
12:3A:81:LEU:HD22	12:3A:82:ILE:H	1.82	0.43
15:6A:87:ILE:HG22	15:6A:88:ARG:H	1.83	0.43
19:AA:29:ARG:CZ	19:AA:48:THR:HG21	2.48	0.43
57:2L:76:C:C2	25:14:2251:OMG:N2	2.81	0.43
25:14:578:A:OP1	25:14:1255:U:O2'	2.24	0.43
25:14:738:G:C6	25:14:739:G:C2	3.06	0.43
25:14:1085:A:HO2'	25:14:1086:A:P	2.40	0.43
25:14:1120:G:H2'	25:14:1121:C:C6	2.53	0.43
25:14:2111:C:H1'	25:14:2118:U:O2'	2.17	0.43
25:14:2165:G:C5	25:14:2166:G:N2	2.86	0.43
25:14:2216:G:H2'	25:14:2217:G:C8	2.53	0.43
25:14:2562:U:H4'	36:25:25:LEU:HD21	2.00	0.43
26:1J:60:C:H2'	26:1J:61:G:C8	2.53	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1J:111:U:H2'	26:1J:112:G:H8	1.83	0.43
28:19:83:GLU:OE1	28:19:104:TYR:OH	2.20	0.43
30:39:154:VAL:HA	30:39:191:ARG:O	2.18	0.43
1:13:191(E):G:H2'	1:13:191(F):U:C6	2.53	0.43
1:13:433:C:H2'	1:13:434:U:C6	2.51	0.43
1:13:538:G:H2'	1:13:539:A:C8	2.51	0.43
1:13:620:C:H2'	1:13:621:A:O4'	2.19	0.43
1:13:950:U:H1'	1:13:971:G:C5	2.53	0.43
1:13:1347:G:N2	1:13:1373:G:H2'	2.33	0.43
1:13:1376:U:OP1	7:6E:98:SER:HB3	2.18	0.43
1:13:1392:G:N2	1:13:1502:A:H8	2.15	0.43
4:3E:135:LEU:C	4:3E:137:SER:H	2.27	0.43
5:4E:39:GLY:C	5:4E:69:VAL:HB	2.43	0.43
10:1I:49:VAL:O	10:1I:60:ARG:HB3	2.18	0.43
22:3K:64:G:H2'	22:3K:65:G:C8	2.53	0.43
25:1H:273(F):C:H3'	25:1H:274:G:C5'	2.47	0.43
25:1H:589:C:H2'	25:1H:590:A:C8	2.53	0.43
25:1H:612:G:N2	25:1H:616:A:O2'	2.51	0.43
25:1H:828:U:H2'	25:1H:829:A:C8	2.53	0.43
25:1H:1019:U:O2'	25:1H:1021:A:H2	2.00	0.43
25:1H:2210:G:H5'	25:1H:2211:G:C5	2.53	0.43
25:1H:2591:C:P	28:11:239:ARG:HG3	2.59	0.43
27:71:5:LYS:HZ2	27:71:8:ARG:HH12	1.66	0.43
32:51:169:VAL:O	32:51:170:ARG:NE	2.24	0.43
33:61:8:PRO:HA	33:61:14:ASP:HA	2.00	0.43
36:68:119:PRO:HB2	41:B8:68:TYR:CE2	2.54	0.43
38:88:39:PRO:HB3	38:88:99:PRO:HD3	2.00	0.43
1:1G:122:G:H1	1:1G:239:U:H3	1.65	0.43
1:1G:489:C:H2'	1:1G:490:G:C8	2.48	0.43
1:1G:595:G:HO2'	1:1G:596:C:P	2.34	0.43
1:1G:607:A:H2'	1:1G:608:A:O4'	2.18	0.43
1:1G:653:A:C8	8:72:56:LYS:HE2	2.53	0.43
1:1G:748:C:H4'	1:1G:749:C:O5'	2.18	0.43
1:1G:998(A):C:H2'	1:1G:999:U:O4'	2.18	0.43
1:1G:1087:G:H2'	1:1G:1088:G:C8	2.53	0.43
3:22:79:ARG:HE	3:22:84:ILE:HG21	1.82	0.43
4:32:15:GLU:HG3	4:32:63:LYS:NZ	2.34	0.43
7:62:136:LYS:O	7:62:140:ASP:HB2	2.18	0.43
9:82:17:VAL:HG13	9:82:63:ILE:HG13	2.01	0.43
9:82:95:LYS:C	9:82:98:PRO:HD2	2.44	0.43
10:1A:50:ILE:HA	10:1A:60:ARG:HD3	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:AA:32:LYS:HE2	19:AA:32:LYS:HB2	1.89	0.43
57:2L:59:A:O2'	57:2L:61:U:OP2	2.28	0.43
25:14:1226:G:OP1	43:95:85:LYS:NZ	2.30	0.43
25:14:1309:G:H4'	55:L5:7:PRO:HB2	2.00	0.43
25:14:2255:G:H1	25:14:2275:C:H42	1.65	0.43
25:14:2363:C:O2	48:E5:39:ARG:NH2	2.52	0.43
25:14:2881:C:O2'	39:55:96:ARG:HB2	2.18	0.43
28:19:129:ASN:O	28:19:193:VAL:HG12	2.17	0.43
35:15:18:ALA:HA	35:15:21:LYS:HD2	2.00	0.43
36:25:87:ILE:HD13	36:25:87:ILE:HA	1.87	0.43
37:35:6:LEU:HA	37:35:6:LEU:HD13	1.76	0.43
46:C5:54:LYS:HD2	46:C5:55:TYR:H	1.83	0.43
1:13:8:A:N6	4:3E:205:GLU:O	2.51	0.43
1:13:271:C:H2'	1:13:272:C:C6	2.54	0.43
1:13:407:G:H5''	4:3E:115:ARG:HB3	1.99	0.43
1:13:624:C:H2'	1:13:625:G:C8	2.53	0.43
2:1E:62:ALA:HB1	2:1E:225:ALA:HB3	1.99	0.43
2:1E:215:LEU:HD22	2:1E:215:LEU:HA	1.77	0.43
4:3E:61:LYS:HA	4:3E:203:VAL:HG22	2.00	0.43
4:3E:94:LEU:H	4:3E:94:LEU:HG	1.55	0.43
9:8E:97:LYS:HB3	9:8E:98:PRO:HD3	2.01	0.43
11:2I:103:LEU:HD12	11:2I:103:LEU:HA	1.86	0.43
13:4I:11:ARG:HG3	13:4I:12:ASN:CG	2.44	0.43
13:4I:23:TYR:HB3	13:4I:67:GLU:HA	1.99	0.43
13:4I:49:THR:O	13:4I:53:VAL:N	2.33	0.43
19:AI:63:THR:OG1	19:AI:64:GLU:N	2.51	0.43
25:1H:251:A:C5	25:1H:252:G:H1'	2.54	0.43
25:1H:898:C:H3'	25:1H:899:A:H8	1.83	0.43
25:1H:1203:G:H3'	25:1H:1204:A:H5''	1.99	0.43
25:1H:1332:G:H8	25:1H:1332:G:H2'	1.65	0.43
25:1H:2030:A:H4'	25:1H:2031:A:C8	2.49	0.43
25:1H:2309:A:N6	25:1H:2310:A:H2	2.16	0.43
25:1H:2405:G:O2'	25:1H:2406:U:OP2	2.33	0.43
25:1H:2688:U:O2	25:1H:2688:U:H3'	2.18	0.43
25:1H:2758:A:C2	25:1H:2759:G:H1'	2.53	0.43
31:4I:66:GLN:HA	52:M8:6:HIS:HE1	1.81	0.43
33:6I:10:GLU:O	33:6I:11:ASN:HB3	2.17	0.43
36:68:105:GLU:HA	36:68:108:GLU:HG3	2.01	0.43
41:B8:105:LEU:C	41:B8:107:ASP:N	2.77	0.43
1:1G:740:U:H2'	1:1G:741:G:C8	2.54	0.43
1:1G:791:G:C6	1:1G:792:A:N7	2.86	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:828:A:N6	1:1G:858:G:O2'	2.48	0.43
1:1G:895:G:H1	1:1G:904:C:N4	2.09	0.43
1:1G:980:C:H5'	1:1G:981:U:OP2	2.19	0.43
1:1G:1161:C:H2'	1:1G:1162:C:H6	1.83	0.43
1:1G:1224:G:O2'	1:1G:1225:A:OP1	2.32	0.43
1:1G:1320:C:C4	19:AA:36:ARG:HD3	2.54	0.43
1:1G:1498:UR3:HO2'	1:1G:1499:A:P	2.36	0.43
3:22:174:PRO:HB2	3:22:177:THR:HG22	2.00	0.43
18:9A:19:LYS:H	18:9A:19:LYS:HE2	1.82	0.43
19:AA:39:THR:HA	19:AA:70:LYS:HB3	2.01	0.43
20:BA:36:LEU:HB3	20:BA:59:ALA:HB2	2.00	0.43
25:14:139:G:H1'	25:14:140:A:C2	2.52	0.43
25:14:329:G:N7	46:C5:71:LYS:NZ	2.66	0.43
25:14:668:G:H2'	25:14:670:A:H62	1.83	0.43
25:14:920:G:H2'	25:14:921:G:H8	1.81	0.43
25:14:1056:G:N2	25:14:1102:C:H5	2.17	0.43
25:14:1789:A:H2'	25:14:1790:C:O4'	2.19	0.43
25:14:1885:A:H3'	25:14:1886:C:C6	2.52	0.43
25:14:2233:U:H2'	25:14:2234:G:C8	2.53	0.43
25:14:2688:U:H1'	25:14:2721:A:N6	2.33	0.43
26:1J:66:A:O2'	26:1J:67:G:OP2	2.35	0.43
28:19:144:ALA:HB3	28:19:192:THR:HG23	1.99	0.43
30:39:162:LEU:HA	30:39:165:ARG:HB2	2.00	0.43
38:45:2:LEU:O	38:45:70:PRO:HG2	2.18	0.43
38:45:81:VAL:HB	38:45:82:ARG:HG2	2.00	0.43
43:95:1:MET:HE3	43:95:1:MET:HB2	1.93	0.43
50:G5:31:GLU:O	50:G5:35:LEU:N	2.50	0.43
1:13:135:C:H2'	1:13:136:C:H5'	1.99	0.43
1:13:172:A:H8	1:13:172:A:OP2	2.01	0.43
1:13:191(D):U:H2'	1:13:191(E):G:C8	2.54	0.43
1:13:339:C:H42	1:13:350:G:H1	1.66	0.43
1:13:611:A:H61	1:13:629:G:H1	1.66	0.43
1:13:685:G:N2	1:13:686:U:O4	2.50	0.43
1:13:1301:U:O2	1:13:1301:U:H2'	2.17	0.43
1:13:1312:G:P	52:M8:58:ARG:HH22	2.41	0.43
1:13:1360:A:OP2	14:5I:35:ARG:NH2	2.46	0.43
15:6I:17:ARG:HD2	15:6I:17:ARG:HA	1.85	0.43
23:2K:37:U:H2'	23:2K:38:A:H8	1.84	0.43
25:1H:402:A:H61	25:1H:423:A:H61	1.66	0.43
25:1H:460:A:H2'	25:1H:461:C:O4'	2.17	0.43
25:1H:663:G:C6	25:1H:664:C:C4	3.06	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:821:A:H5'	25:1H:822:U:C6	2.50	0.43
25:1H:1341:U:OP2	25:1H:1394:U:O2'	2.23	0.43
25:1H:1351:C:O2'	25:1H:1571:A:N3	2.49	0.43
25:1H:2025:C:OP1	29:21:149:ARG:HD3	2.19	0.43
34:38:21:GLN:HB3	34:38:24:PHE:HE2	1.84	0.43
40:A8:78:LEU:HD12	40:A8:108:GLY:HA3	2.00	0.43
43:D8:24:LYS:HB3	43:D8:24:LYS:NZ	2.33	0.43
46:G8:89:PHE:HD1	46:G8:94:LYS:HB3	1.83	0.43
51:L8:18:ASP:OD1	51:L8:18:ASP:N	2.52	0.43
1:1G:201:C:C4	1:1G:209:U:H1'	2.53	0.43
1:1G:297:G:O2'	1:1G:299:G:N7	2.49	0.43
1:1G:1129:C:C5'	1:1G:1130:A:H5'	2.48	0.43
1:1G:1326:C:H2'	1:1G:1327:C:C6	2.53	0.43
15:6A:56:LEU:O	15:6A:60:VAL:HG23	2.18	0.43
25:14:171:G:O2'	25:14:172:C:O4'	2.32	0.43
25:14:363(B):G:H2'	25:14:363(C):G:H8	1.83	0.43
25:14:550:G:O2'	25:14:1220:A:N3	2.36	0.43
25:14:1027:A:N6	25:14:1126:A:C4	2.87	0.43
25:14:1637:A:H4'	25:14:2711:A:O2'	2.18	0.43
25:14:1815:A:C5	25:14:1817:G:C6	3.07	0.43
25:14:2873:A:C8	39:55:5:LYS:HA	2.54	0.43
28:19:93:ALA:HB3	28:19:105:ILE:HG22	2.00	0.43
28:19:175:LEU:HA	28:19:175:LEU:HD23	1.69	0.43
44:A5:33:ARG:HD3	44:A5:52:GLU:OE1	2.19	0.43
48:E5:18:ALA:HB3	48:E5:20:ARG:NH2	2.34	0.43
1:13:162:A:N7	1:13:163:C:H1'	2.33	0.43
1:13:474:G:H2'	1:13:475:G:H8	1.84	0.43
1:13:972:C:P	10:1I:57:LYS:HZ3	2.41	0.43
2:1E:68:ILE:O	2:1E:91:PRO:HD2	2.18	0.43
4:3E:65:ARG:HB3	4:3E:75:PHE:CD1	2.54	0.43
4:3E:121:VAL:HG12	4:3E:133:VAL:O	2.19	0.43
6:5E:100:ASN:HB2	18:9I:28:GLU:HA	2.00	0.43
7:6E:108:ALA:HA	7:6E:111:ARG:HD2	2.01	0.43
12:3I:20:LYS:HD3	12:3I:21:VAL:HG23	2.01	0.43
17:8I:12:SER:O	17:8I:19:VAL:HA	2.18	0.43
17:8I:68:ARG:HG3	17:8I:68:ARG:O	2.19	0.43
22:1K:52:C:H2'	22:1K:53:G:C8	2.54	0.43
25:1H:141:A:H8	25:1H:1595:G:H21	1.66	0.43
25:1H:1151:G:C2	25:1H:1152:C:C2	3.06	0.43
25:1H:1681:G:H4'	25:1H:1682:G:OP2	2.18	0.43
25:1H:1751:C:H2'	25:1H:1752:C:C6	2.53	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:1784:A:H5''	62:1H:3899:HOH:O	2.17	0.43
25:1H:2400:G:H2'	25:1H:2401:U:C6	2.54	0.43
25:1H:2726:U:O2'	25:1H:2727:G:H8	2.01	0.43
32:51:137:ASP:OD1	32:51:138:LYS:N	2.52	0.43
35:58:15:LEU:HA	35:58:53:VAL:O	2.18	0.43
48:I8:26:TYR:O	48:I8:29:GLN:HB2	2.19	0.43
54:O8:41:PRO:HG2	54:O8:44:ARG:HB2	2.00	0.43
1:1G:74:C:N3	1:1G:96:G:N2	2.45	0.43
1:1G:544:G:OP1	4:32:59:ARG:NH2	2.51	0.43
1:1G:1216:G:H2'	1:1G:1217:C:C6	2.53	0.43
1:1G:1309:G:P	13:4A:88:ARG:HH12	2.41	0.43
3:22:136:GLN:H	3:22:136:GLN:HG3	1.50	0.43
9:82:96:LEU:HD23	9:82:102:LEU:HD12	1.99	0.43
10:1A:76:ASN:HA	10:1A:77:PRO:HD3	1.90	0.43
15:6A:53:HIS:O	15:6A:56:LEU:HB3	2.19	0.43
18:9A:22:VAL:HG13	18:9A:56:THR:HA	2.00	0.43
22:3L:23:G:H8	22:3L:23:G:OP2	2.01	0.43
25:14:442:G:O4'	30:39:46:ARG:HD3	2.19	0.43
25:14:465:G:C6	25:14:466:A:N6	2.86	0.43
25:14:1311:G:N7	55:L5:47:ARG:NH1	2.56	0.43
25:14:2123:G:H2'	25:14:2124:G:C8	2.53	0.43
25:14:2299:G:C2	25:14:2300:G:C5	3.07	0.43
25:14:2299:G:N1	25:14:2318:G:C8	2.86	0.43
25:14:2607:G:H2'	25:14:2608:G:O4'	2.18	0.43
26:1J:8:U:O3'	40:65:25:ARG:NH2	2.51	0.43
26:1J:111:U:H2'	26:1J:112:G:C8	2.53	0.43
32:59:22:GLY:C	32:59:37:VAL:HG12	2.44	0.43
41:75:51:ARG:HE	41:75:62:THR:CG2	2.32	0.43
42:85:95:LEU:HD13	43:95:4:ILE:HG23	2.00	0.43
44:A5:5:ALA:HB2	44:A5:57:ASN:HD22	1.83	0.43
46:C5:42:VAL:HG23	46:C5:43:ASN:N	2.34	0.43
1:13:130:A:C8	17:8I:63:ARG:HG3	2.53	0.43
1:13:1128:C:H4'	9:8E:16:ARG:NH1	2.34	0.43
8:7E:96:GLY:O	8:7E:100:ILE:HG13	2.18	0.43
14:5I:9:LYS:HG2	14:5I:12:ARG:NH1	2.32	0.43
25:1H:32:C:OP2	62:1H:3699:HOH:O	2.21	0.43
25:1H:194:G:H2'	25:1H:195:A:O4'	2.19	0.43
25:1H:553:U:H2'	25:1H:554:U:O4'	2.19	0.43
25:1H:862:G:H5'	26:16:79:C:H4'	2.01	0.43
25:1H:2243:U:O4	62:1H:3694:HOH:O	2.20	0.43
25:1H:2404:C:O3'	37:78:77:ARG:NH2	2.51	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:2712:U:O2'	25:1H:2712(A):A:P	2.76	0.43
28:11:206:LEU:HD23	28:11:206:LEU:HA	1.86	0.43
30:31:59:TYR:CD2	30:31:78:ILE:HG13	2.53	0.43
31:41:118:ARG:HA	31:41:118:ARG:HD2	1.57	0.43
33:61:133:HIS:CG	33:61:134:PRO:HD2	2.54	0.43
35:58:132:ALA:O	35:58:134:ARG:HD3	2.18	0.43
41:B8:123:GLN:H	41:B8:123:GLN:HG3	1.60	0.43
56:Q8:8:LYS:O	56:Q8:12:LYS:HE3	2.19	0.43
1:1G:674:G:H2'	1:1G:675:A:H8	1.84	0.43
1:1G:986:A:H1'	19:AA:54:GLY:O	2.19	0.43
1:1G:1033:G:O2'	1:1G:1034:G:O5'	2.31	0.43
1:1G:1129:C:C2	1:1G:1139:G:N2	2.87	0.43
1:1G:1132:C:H2'	1:1G:1133:G:C8	2.53	0.43
1:1G:1216:G:H2'	1:1G:1217:C:H6	1.83	0.43
1:1G:1320:C:O4'	19:AA:36:ARG:NH2	2.52	0.43
9:82:27:THR:CG2	9:82:33:PHE:HB2	2.48	0.43
13:4A:80:ARG:HH12	52:I5:55:ARG:HD3	1.82	0.43
14:5A:17:LYS:HG3	14:5A:18:VAL:N	2.34	0.43
57:2L:45:A:H2'	57:2L:46:G:O4'	2.19	0.43
25:14:26:G:H1'	25:14:515:A:H61	1.84	0.43
25:14:77:C:OP2	62:14:3559:HOH:O	2.21	0.43
25:14:380:U:H2'	25:14:381:G:H8	1.83	0.43
25:14:447:A:O2'	25:14:473:G:N7	2.44	0.43
25:14:513:A:H2'	25:14:514:A:H8	1.83	0.43
25:14:863:A:H2'	25:14:864:G:C8	2.53	0.43
25:14:876:C:H2'	25:14:877:U:O4'	2.19	0.43
25:14:942:G:O2'	25:14:1189:A:N3	2.49	0.43
26:1J:104:A:H2'	26:1J:105:G:O4'	2.19	0.43
28:19:262:ARG:O	28:19:264:LYS:N	2.51	0.43
33:69:101:LEU:HD12	33:69:107:VAL:HB	1.99	0.43
36:25:8:LEU:HB2	36:25:83:ALA:HA	2.01	0.43
38:45:68:ILE:HG22	38:45:101:ARG:HE	1.83	0.43
41:75:102:ILE:HB	41:75:110:ILE:CD1	2.49	0.43
47:D5:82:ARG:NH2	47:D5:83:PRO:O	2.51	0.43
54:K5:12:GLU:HG3	54:K5:22:ALA:HB3	2.00	0.43
54:K5:23:THR:O	54:K5:24:GLU:HB2	2.19	0.43
1:13:413:G:H2'	1:13:428:G:H22	1.83	0.43
1:13:474:G:H5''	16:7I:81:ARG:CZ	2.49	0.43
1:13:719:C:H3'	1:13:720:C:C6	2.54	0.43
1:13:765:G:H5''	1:13:766:A:OP1	2.19	0.43
1:13:872:A:C8	1:13:874:G:C8	3.07	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1347:G:H2'	1:13:1373:G:H1	1.83	0.43
8:7E:86:ILE:HG12	8:7E:135:CYS:HA	2.01	0.43
9:8E:69:GLY:O	9:8E:73:GLN:HG3	2.17	0.43
13:4I:69:GLU:O	13:4I:73:GLU:HB3	2.18	0.43
19:AI:36:ARG:H	19:AI:36:ARG:HG2	1.59	0.43
20:BI:52:ALA:O	20:BI:56:MET:N	2.43	0.43
25:1H:773:U:C4'	28:11:47:GLY:HA3	2.48	0.43
25:1H:859:G:H5'	25:1H:2268:A:O2'	2.18	0.43
25:1H:1417:C:H42	25:1H:1581:G:H1	1.66	0.43
25:1H:1820:U:H5''	25:1H:1821:A:C8	2.54	0.43
25:1H:2445:G:OP1	30:31:74:ARG:NH2	2.42	0.43
25:1H:2747:G:OP1	32:51:138:LYS:NZ	2.51	0.43
30:31:154:VAL:HB	30:31:173:VAL:HG22	2.00	0.43
35:58:114:ARG:C	35:58:116:LEU:H	2.27	0.43
37:78:65:ARG:HG3	37:78:65:ARG:O	2.17	0.43
41:B8:54:ARG:HA	41:B8:59:THR:OG1	2.19	0.43
42:C8:61:TRP:HB3	42:C8:93:LYS:O	2.19	0.43
46:G8:75:ILE:HD13	46:G8:75:ILE:HA	1.68	0.43
1:1G:682:G:C6	1:1G:709:G:C6	3.07	0.43
1:1G:767:A:H2'	1:1G:768:A:H8	1.83	0.43
1:1G:922:G:N3	1:1G:1398:A:H2	2.17	0.43
1:1G:1011:G:H1	1:1G:1018:C:N4	2.17	0.43
1:1G:1090:U:H2'	1:1G:1091:U:H6	1.83	0.43
1:1G:1513:A:H2'	1:1G:1514:C:C6	2.54	0.43
2:12:187:LEU:HD22	2:12:201:ILE:O	2.19	0.43
5:42:131:ILE:HD13	5:42:131:ILE:HA	1.89	0.43
14:5A:24:CYS:SG	14:5A:25:VAL:N	2.92	0.43
19:AA:16:LEU:HD22	19:AA:16:LEU:H	1.83	0.43
19:AA:66:MET:HG2	19:AA:74:PHE:CZ	2.53	0.43
25:14:204:A:O3'	25:14:205:G:H4'	2.18	0.43
25:14:724:U:H2'	25:14:725:G:O4'	2.18	0.43
25:14:826:U:H5''	25:14:2428:G:O3'	2.19	0.43
25:14:1190:G:OP1	37:35:30:THR:OG1	2.36	0.43
25:14:1332:G:N2	25:14:1609:A:O2'	2.52	0.43
25:14:2141:G:H2'	25:14:2142:C:O4'	2.19	0.43
25:14:2313:C:H2'	25:14:2314:C:C6	2.54	0.43
46:C5:39:VAL:O	46:C5:42:VAL:HG22	2.18	0.43
49:F5:83:GLU:H	49:F5:83:GLU:CD	2.27	0.43
1:13:343:U:HO2'	1:13:346:G:H1	1.67	0.43
1:13:590:C:H2'	1:13:591:U:H6	1.84	0.43
1:13:748:C:H4'	1:13:749:C:O5'	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:797:C:OP1	11:2I:124:LYS:NZ	2.43	0.43
1:13:887:G:H2'	1:13:888:G:O4'	2.18	0.43
1:13:1004:A:C8	1:13:1026:G:C5	3.03	0.43
1:13:1124:G:OP2	1:13:1124:G:H8	2.01	0.43
1:13:1316:G:N1	1:13:1319:A:OP2	2.49	0.43
1:13:1384:C:H2'	1:13:1385:G:C8	2.54	0.43
2:1E:10:LEU:HD23	2:1E:10:LEU:HA	1.89	0.43
2:1E:130:ARG:HB3	2:1E:134:GLU:HB2	2.00	0.43
4:3E:14:ARG:HG2	4:3E:40:PRO:HD2	2.00	0.43
5:4E:15:ARG:HD2	5:4E:26:PHE:CD2	2.54	0.43
17:8I:26:GLN:HB3	17:8I:37:LYS:HG2	2.00	0.43
22:1K:20:G:H4'	22:1K:21:U:OP2	2.15	0.43
25:1H:469:G:O6	55:P8:37:LYS:HE2	2.19	0.43
25:1H:1278:A:H2'	25:1H:1279:G:H8	1.83	0.43
25:1H:1337:G:H2'	25:1H:1338:G:O4'	2.19	0.43
25:1H:1628:G:H1	25:1H:1638:C:H42	1.67	0.43
25:1H:2074:U:H2'	25:1H:2075:U:C6	2.54	0.43
25:1H:2251:OMG:H1'	25:1H:2251:OMG:HM23	1.63	0.43
25:1H:2600:A:C6	25:1H:2601:C:N4	2.87	0.43
27:71:30:LYS:HD2	27:71:182:PRO:HB3	2.00	0.43
30:31:12:LEU:HA	30:31:12:LEU:HD12	1.86	0.43
30:31:184:TYR:O	30:31:188:ARG:HG3	2.18	0.43
31:41:99:MET:HB2	31:41:99:MET:HE3	1.58	0.43
32:51:86:GLU:O	32:51:87:LEU:HB2	2.19	0.43
38:88:66:ILE:HA	38:88:104:PHE:HA	2.00	0.43
45:F8:31:HIS:HE1	45:F8:33:LYS:HB2	1.83	0.43
47:H8:10:ARG:HH21	47:H8:26:GLY:H	1.67	0.43
54:O8:33:LYS:HG2	54:O8:34:LEU:HD22	2.00	0.43
1:1G:186(F):C:H2'	1:1G:187:C:O4'	2.18	0.43
1:1G:474:G:H5'	16:7A:81:ARG:HD3	2.00	0.43
1:1G:1015:A:H8	1:1G:1015:A:O5'	2.02	0.43
1:1G:1028:C:H2'	1:1G:1028(A):C:O4'	2.19	0.43
1:1G:1515:C:H2'	1:1G:1516:G:C8	2.54	0.43
2:12:167:PRO:HG2	2:12:192:SER:CB	2.49	0.43
3:22:37:GLN:O	3:22:41:GLY:N	2.43	0.43
6:52:70:ASP:OD1	6:52:71:ARG:HG2	2.18	0.43
6:52:77:ARG:HH11	6:52:77:ARG:HB3	1.84	0.43
8:72:20:TYR:CE2	8:72:75:ARG:HD2	2.52	0.43
9:82:4:TYR:HE1	9:82:88:TYR:CD1	2.37	0.43
19:AA:70:LYS:O	19:AA:71:LEU:HD23	2.19	0.43
25:14:29:U:H2'	25:14:30:G:H8	1.83	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:572:A:H5''	25:14:573:G:OP2	2.18	0.43
25:14:839:U:H2'	25:14:840:C:C6	2.54	0.43
25:14:872:A:C6	25:14:906:G:C2	3.07	0.43
25:14:1032:A:H2	25:14:1122:G:H1	1.67	0.43
25:14:1177:A:H5'	25:14:1178:C:OP1	2.19	0.43
25:14:1614:A:P	25:14:1614:A:H8	2.41	0.43
25:14:2127:G:C2	25:14:2128:C:H1'	2.54	0.43
25:14:2528:U:H2'	25:14:2530:A:O5'	2.19	0.43
25:14:2689:U:P	25:14:2719:G:H22	2.41	0.43
32:59:20:ALA:HB1	32:59:21:PRO:HD2	2.00	0.43
36:25:79:PHE:CD2	41:75:72:VAL:HG22	2.54	0.43
38:45:66:ILE:HG12	38:45:67:ARG:N	2.32	0.43
39:55:52:ILE:HD13	39:55:79:LEU:HD21	1.99	0.43
47:D5:30:ASN:OD1	47:D5:33:LEU:N	2.48	0.43
49:F5:40:ARG:NH2	49:F5:42:GLN:HG2	2.32	0.43
1:13:295:C:H2'	1:13:296:U:O4'	2.18	0.43
1:13:341:C:O2'	1:13:342:C:H5'	2.19	0.43
1:13:375:U:O3'	16:7I:6:LEU:HB2	2.19	0.43
1:13:642:A:H5''	8:7E:30:ARG:HH21	1.83	0.43
1:13:1425:U:H2'	1:13:1426:C:C6	2.54	0.43
2:1E:21:ARG:HA	2:1E:39:ILE:HA	2.00	0.43
2:1E:22:LYS:HG3	2:1E:35:GLU:OE1	2.18	0.43
2:1E:189:ASP:HB3	2:1E:203:GLY:O	2.19	0.43
4:3E:121:VAL:HG13	4:3E:126:ILE:HD12	2.01	0.43
13:4I:57:ARG:HG3	13:4I:61:GLU:HG3	2.00	0.43
21:1F:4:GLY:HA2	21:1F:15:ARG:HH12	1.83	0.43
22:1K:5:G:N3	22:1K:5:G:H2'	2.34	0.43
25:1H:48:G:C2	25:1H:178:G:C6	3.07	0.43
25:1H:725:G:C6	25:1H:726:G:N1	2.87	0.43
25:1H:1363:C:H2'	25:1H:1364:G:H8	1.83	0.43
25:1H:1364:G:C8	49:J8:2:SER:HB3	2.54	0.43
25:1H:1567:A:OP2	28:11:84:TYR:OH	2.21	0.43
25:1H:1957:C:H2'	25:1H:1958:C:C6	2.54	0.43
25:1H:2131:G:H4'	25:1H:2132:U:C4'	2.43	0.43
25:1H:2164:C:H2'	25:1H:2165:G:O4'	2.19	0.43
25:1H:2637:U:H2'	25:1H:2638:G:O4'	2.19	0.43
25:1H:2849:U:H4'	25:1H:2868:A:N1	2.34	0.43
27:71:3:HIS:HB2	27:71:7:TYR:CD1	2.54	0.43
32:51:121:ILE:HD13	32:51:121:ILE:HA	1.88	0.43
37:78:6:LEU:HD12	37:78:6:LEU:HA	1.69	0.43
41:B8:102:ILE:HB	41:B8:110:ILE:HD11	2.01	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:E8:73:ALA:HB3	44:E8:106:ILE:HG12	2.01	0.43
49:J8:81:LYS:HB3	49:J8:82:LEU:H	1.54	0.43
52:M8:42:PHE:CD1	52:M8:42:PHE:C	2.97	0.43
53:N8:20:ARG:HA	53:N8:23:HIS:ND1	2.33	0.43
54:O8:17:LYS:C	54:O8:19:ARG:H	2.25	0.43
1:1G:468:A:H2'	1:1G:474:G:O4'	2.19	0.43
1:1G:928:G:H2'	1:1G:929:G:H8	1.83	0.43
1:1G:994:A:H8	1:1G:994:A:OP1	2.02	0.43
1:1G:1092:A:H61	1:1G:1110:A:H4'	1.83	0.43
1:1G:1324:A:P	13:4A:99:ARG:HH22	2.42	0.43
2:12:42:ILE:CD1	2:12:202:PRO:HB2	2.49	0.43
4:32:168:ARG:HB3	4:32:168:ARG:NH1	2.31	0.43
9:82:5:TYR:H	9:82:87:GLN:NE2	2.16	0.43
25:14:196:A:N3	25:14:196:A:H2'	2.34	0.43
25:14:671:C:H2'	25:14:672:C:C6	2.54	0.43
25:14:1349:A:N6	25:14:1598:C:H42	2.17	0.43
29:29:101:ARG:HA	29:29:170:LEU:O	2.19	0.43
33:69:123:LEU:HD22	33:69:143:SER:CB	2.49	0.43
39:55:90:ARG:NH2	39:55:117:VAL:HG21	2.33	0.43
46:C5:106:LEU:C	46:C5:108:THR:N	2.76	0.43
47:D5:33:LEU:HD13	47:D5:33:LEU:HA	1.82	0.43
3:2E:106:VAL:HG11	3:2E:115:LEU:HD11	2.01	0.42
3:2E:130:VAL:O	3:2E:134:ILE:HG12	2.19	0.42
6:5E:86:ARG:O	6:5E:87:ARG:HG2	2.19	0.42
11:2I:99:GLN:HG2	11:2I:105:VAL:HG11	2.01	0.42
12:3I:74:LEU:HD21	12:3I:104:ALA:HB2	2.01	0.42
16:7I:71:ARG:HE	16:7I:80:PHE:HZ	1.66	0.42
22:3K:52:C:H2'	22:3K:53:G:C8	2.53	0.42
25:1H:1178:C:H4'	25:1H:1179:C:OP1	2.19	0.42
25:1H:1244:G:O5'	37:78:7:ARG:HG3	2.19	0.42
25:1H:1566:A:P	28:11:211:ARG:HH21	2.41	0.42
25:1H:1753:G:N1	25:1H:1756:G:OP2	2.51	0.42
25:1H:1858:G:H1'	25:1H:1884:A:N6	2.34	0.42
25:1H:2282:G:H4'	25:1H:2389:G:O2'	2.19	0.42
25:1H:2542:A:H4'	25:1H:2543:G:C8	2.54	0.42
25:1H:2561:A:H2'	25:1H:2562:U:O4'	2.19	0.42
27:71:225:ASN:ND2	27:71:228:SER:H	2.17	0.42
28:11:4:LYS:NZ	28:11:20:ASP:OD1	2.52	0.42
29:21:67:PHE:O	29:21:71:GLY:HA2	2.19	0.42
31:41:112:PRO:HB3	52:M8:37:SER:H	1.82	0.42
32:51:88:LEU:H	32:51:88:LEU:HG	1.66	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:78:61:ARG:HB3	37:78:61:ARG:HH11	1.83	0.42
40:A8:26:LEU:HD22	40:A8:87:PHE:HD1	1.84	0.42
40:A8:30:ARG:NH2	40:A8:92:TYR:HD1	2.16	0.42
42:C8:98:LEU:HD23	42:C8:99:ALA:N	2.34	0.42
48:I8:12:ASN:HA	48:I8:14:ARG:NH2	2.34	0.42
48:I8:29:GLN:O	48:I8:31:VAL:HG13	2.19	0.42
54:O8:36:LEU:HD22	54:O8:50:ARG:HH12	1.83	0.42
1:1G:596:C:N4	1:1G:644:G:H1	2.17	0.42
1:1G:689:C:P	11:2A:46:GLY:HA3	2.59	0.42
1:1G:752:G:H1'	1:1G:754:C:N4	2.34	0.42
1:1G:860:A:H4'	8:72:75:ARG:NH1	2.34	0.42
1:1G:948:C:OP2	13:4A:106:ASN:HB2	2.19	0.42
1:1G:1035:A:H2'	1:1G:1036:G:H5''	2.01	0.42
1:1G:1437:C:N4	1:1G:1464:G:H1	2.08	0.42
9:82:54:ASP:OD1	9:82:54:ASP:N	2.51	0.42
10:1A:33:GLN:O	10:1A:75:ILE:HG12	2.19	0.42
12:3A:9:ARG:O	12:3A:11:GLY:N	2.52	0.42
17:8A:58:GLU:O	17:8A:74:LEU:N	2.50	0.42
25:14:389:G:N1	37:35:71:VAL:HG12	2.34	0.42
25:14:565:C:H4'	25:14:1253:A:C6	2.53	0.42
25:14:597:U:H3	25:14:660:G:H1	1.67	0.42
25:14:1141:U:H4'	25:14:1142(A):A:O4'	2.19	0.42
25:14:1492:G:H3'	25:14:1493:C:C5'	2.49	0.42
25:14:1776:G:O6	62:14:3548:HOH:O	2.19	0.42
25:14:1814:G:H2'	25:14:1815:A:C8	2.53	0.42
25:14:2481:G:O2'	25:14:2482:G:H8	2.01	0.42
25:14:2875:C:H2'	25:14:2876:G:O4'	2.19	0.42
26:1J:40:U:H3	26:1J:43:C:H5''	1.84	0.42
28:19:136:ILE:O	28:19:168:ARG:NH2	2.52	0.42
33:69:117:GLU:OE1	33:69:118:LYS:HG2	2.19	0.42
41:75:26:ASP:HB3	41:75:88:ILE:HD13	2.01	0.42
45:B5:61:GLY:N	45:B5:75:ASP:HB3	2.34	0.42
1:13:167:G:H2'	1:13:168:G:H8	1.83	0.42
1:13:270:A:H2'	1:13:271:C:C6	2.55	0.42
1:13:538:G:O3'	12:3I:111:LYS:HD3	2.19	0.42
1:13:601:C:H2'	1:13:602:A:C8	2.55	0.42
1:13:991:U:O2	1:13:993:G:H8	2.02	0.42
4:3E:108:LEU:HD12	4:3E:174:LEU:HD13	2.00	0.42
9:8E:7:THR:O	9:8E:83:ARG:HD2	2.19	0.42
25:1H:27:G:O2'	25:1H:28:A:H8	2.02	0.42
25:1H:71:A:H4'	25:1H:72:U:H5''	1.99	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:738:G:C6	25:1H:739:G:C2	3.06	0.42
25:1H:979:G:H2'	25:1H:982:C:H42	1.84	0.42
25:1H:1786:A:H1'	25:1H:1938:A:N6	2.34	0.42
25:1H:1846:G:H5'	25:1H:1847:A:OP2	2.19	0.42
25:1H:2505:G:O6	25:1H:2576:G:H2'	2.19	0.42
25:1H:2712:U:OP1	25:1H:2714:G:H4'	2.19	0.42
29:21:9:VAL:HG13	29:21:25:VAL:HB	2.00	0.42
30:31:53:THR:HG23	30:31:56:GLU:OE1	2.19	0.42
34:38:36:GLU:C	34:38:102:LYS:HZ1	2.27	0.42
35:58:128:HIS:ND1	35:58:129:PRO:O	2.52	0.42
39:98:44:LEU:O	39:98:48:VAL:HG22	2.19	0.42
42:C8:29:SER:OG	42:C8:30:LYS:HE2	2.19	0.42
46:G8:40:GLU:HA	46:G8:41:GLY:HA2	1.57	0.42
1:1G:183:G:H1	1:1G:194:C:H42	1.67	0.42
1:1G:225:C:H2'	1:1G:226:G:H8	1.83	0.42
1:1G:1279:A:O2'	1:1G:1282:C:N4	2.53	0.42
1:1G:1292:U:H2'	1:1G:1293:G:C8	2.55	0.42
11:2A:27:ASN:OD1	11:2A:28:THR:N	2.52	0.42
12:3A:10:LYS:H	12:3A:10:LYS:HG3	1.59	0.42
18:9A:65:ILE:H	18:9A:65:ILE:HG13	1.60	0.42
25:14:1044:G:N2	25:14:1112:G:H1'	2.34	0.42
25:14:1186:G:H2'	25:14:1187:G:O4'	2.19	0.42
25:14:1331:A:OP2	62:14:3561:HOH:O	2.21	0.42
25:14:1360:A:H5'	25:14:1361:G:OP2	2.18	0.42
25:14:1910:G:H2'	25:14:1911:PSU:H6	1.84	0.42
25:14:2679:A:H4'	29:29:165:VAL:HG11	2.02	0.42
36:25:12:ASP:N	36:25:12:ASP:OD1	2.51	0.42
42:85:82:GLY:O	42:85:86:ALA:N	2.51	0.42
43:95:12:TYR:CZ	43:95:22:VAL:HG23	2.54	0.42
45:B5:57:LEU:HD21	45:B5:78:LYS:HG3	2.01	0.42
47:D5:144:LEU:H	47:D5:144:LEU:HD12	1.84	0.42
1:13:828:A:N6	1:13:858:G:O2'	2.47	0.42
1:13:1057:G:H2'	1:13:1058:G:O4'	2.19	0.42
1:13:1347:G:H2'	1:13:1373:G:N1	2.34	0.42
1:13:1510:U:H2'	1:13:1511:G:C8	2.54	0.42
5:4E:118:ILE:HG12	5:4E:120:THR:HG23	2.01	0.42
15:6I:58:MET:HE2	15:6I:58:MET:HB3	1.90	0.42
16:7I:4:ILE:HG12	16:7I:21:VAL:HG12	2.02	0.42
20:BI:55:ILE:HD13	20:BI:55:ILE:HA	1.88	0.42
22:3K:20:G:C4	25:1H:2112:G:N7	2.87	0.42
22:3K:59:A:O2'	22:3K:61:U:H5	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:27:G:N2	25:1H:512:G:H2'	2.31	0.42
25:1H:120:U:OP2	62:1H:3679:HOH:O	2.21	0.42
25:1H:570:G:H2'	25:1H:2030:A:C5	2.54	0.42
25:1H:582:G:H2'	25:1H:583:G:C8	2.54	0.42
25:1H:938:G:OP2	56:Q8:52:LYS:NZ	2.50	0.42
25:1H:1279:G:H4'	39:98:31:HIS:CD2	2.54	0.42
25:1H:2010:G:H5''	44:E8:42:ARG:HB2	2.01	0.42
25:1H:2157:G:H4'	25:1H:2158:A:O5'	2.20	0.42
25:1H:2271:G:OP1	48:I8:18:ALA:HB1	2.19	0.42
25:1H:2423:U:H6	25:1H:2423:U:H2'	1.72	0.42
25:1H:2751:G:HO2'	25:1H:2752:C:P	2.42	0.42
32:51:89:ILE:HB	32:51:129:THR:O	2.18	0.42
37:78:49:ARG:HG3	56:Q8:61:LEU:HD11	2.01	0.42
37:78:128:HIS:HA	37:78:147:LEU:HB2	2.00	0.42
38:88:125:LEU:HD23	38:88:125:LEU:HA	1.88	0.42
1:1G:452:A:HO2'	1:1G:453:A:C5'	2.33	0.42
1:1G:1112:C:O2	3:22:179:ARG:HG2	2.18	0.42
1:1G:1369:C:H2'	1:1G:1370:G:O4'	2.19	0.42
2:12:111:ARG:HD3	2:12:111:ARG:HA	1.50	0.42
3:22:35:GLU:O	3:22:39:ILE:HG13	2.19	0.42
4:32:61:LYS:HD3	4:32:206:PHE:HE2	1.83	0.42
4:32:127:THR:HG22	4:32:147:ALA:HB3	2.02	0.42
6:52:45:LEU:HD21	6:52:57:GLN:HB3	2.01	0.42
25:14:182:A:H8	25:14:182:A:OP2	2.02	0.42
25:14:693:C:H42	25:14:769:G:H1	1.68	0.42
25:14:1087:G:P	25:14:1087:G:H8	2.42	0.42
25:14:1526:G:H3'	25:14:1527:G:C8	2.53	0.42
25:14:1567:A:H5'	28:19:86:PRO:CB	2.48	0.42
25:14:1578:U:H6	25:14:1578:U:OP2	2.02	0.42
25:14:1858:G:C6	25:14:1883:G:C6	3.07	0.42
25:14:2094:G:H5'	33:69:25:TYR:CD1	2.54	0.42
25:14:2166:G:H4'	25:14:2167:U:OP1	2.20	0.42
25:14:2712(A):A:H5''	25:14:2713:A:OP2	2.18	0.42
26:1J:31:C:H42	26:1J:51:G:H1	1.67	0.42
26:1J:37:C:N3	26:1J:48:A:O2'	2.50	0.42
30:39:25:PRO:C	30:39:27:GLU:H	2.27	0.42
32:59:41:MET:HE2	32:59:55:PRO:HD2	2.01	0.42
38:45:47:ILE:O	38:45:51:ARG:N	2.43	0.42
41:75:51:ARG:HE	41:75:62:THR:HG23	1.84	0.42
1:13:197:A:H4'	1:13:198:G:O5'	2.18	0.42
1:13:198:G:H2'	1:13:199:G:C8	2.54	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:608:A:HO2'	16:7I:32:TYR:HH	1.65	0.42
1:13:623:C:H2'	1:13:624:C:O4'	2.19	0.42
1:13:1149:C:H2'	1:13:1150:U:O4'	2.18	0.42
2:1E:224:GLN:HA	2:1E:229:VAL:HG22	2.02	0.42
3:2E:11:ARG:NH2	3:2E:180:ALA:HB3	2.34	0.42
6:5E:67:MET:HE3	6:5E:67:MET:HB2	1.86	0.42
7:6E:45:ASP:O	7:6E:48:LYS:HB3	2.20	0.42
7:6E:66:VAL:O	7:6E:70:LYS:HG3	2.19	0.42
11:2I:69:ALA:O	11:2I:73:MET:HG3	2.19	0.42
13:4I:65:LYS:HE2	13:4I:65:LYS:HB3	1.83	0.42
25:1H:1587:A:H2'	25:1H:1588:C:C6	2.54	0.42
25:1H:1590:U:H2'	25:1H:1591:G:C8	2.55	0.42
25:1H:2163:C:H2'	25:1H:2164:C:C6	2.53	0.42
26:16:95:U:H2'	26:16:96:G:H8	1.82	0.42
28:11:38:LYS:HE2	28:11:39:LYS:O	2.19	0.42
28:11:223:GLY:HA3	28:11:231:HIS:CE1	2.54	0.42
30:31:68:LYS:HB3	30:31:69:HIS:CD2	2.55	0.42
30:31:136:THR:HA	30:31:166:ALA:HB1	2.02	0.42
32:51:26:VAL:HG11	32:51:33:LEU:HD22	2.01	0.42
33:61:113:ARG:HD3	33:61:113:ARG:N	2.34	0.42
42:C8:66:ASN:O	42:C8:70:ARG:HB2	2.19	0.42
42:C8:92:ARG:C	42:C8:94:ASN:N	2.78	0.42
1:1G:34:C:H2'	1:1G:35:G:C8	2.54	0.42
1:1G:44:G:H1	1:1G:398:C:H42	1.67	0.42
1:1G:345:C:H4'	1:1G:346:G:C4	2.55	0.42
1:1G:686:U:H1'	11:2A:42:TRP:NE1	2.33	0.42
1:1G:849:C:H2'	1:1G:850:U:O4'	2.20	0.42
1:1G:960:U:HO2'	1:1G:961:U:P	2.43	0.42
1:1G:1403:C:H2'	1:1G:1404:5MC:HM53	2.00	0.42
1:1G:1465:C:H2'	1:1G:1466:C:O4'	2.20	0.42
3:22:54:ARG:HB2	3:22:69:HIS:CG	2.54	0.42
3:22:95:THR:HG22	3:22:97:LYS:H	1.85	0.42
4:32:63:LYS:HB2	4:32:63:LYS:HE2	1.79	0.42
7:62:71:PRO:HG2	7:62:91:VAL:HG21	2.01	0.42
11:2A:85:ARG:HG2	11:2A:111:ASP:O	2.19	0.42
12:3A:83:ARG:HB2	12:3A:98:VAL:HG22	2.01	0.42
17:8A:85:VAL:O	17:8A:89:LEU:HG	2.20	0.42
18:9A:50:ILE:HD12	18:9A:70:ILE:HD13	2.01	0.42
19:AA:36:ARG:CG	19:AA:72:GLY:H	2.30	0.42
25:14:640:C:N4	25:14:648:G:H1	2.11	0.42
25:14:1138:G:O2'	35:15:102:ALA:O	2.37	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:1542:G:H5''	25:14:1543:A:OP2	2.19	0.42
25:14:2867:G:O2'	25:14:2868:A:H8	2.01	0.42
26:1J:6:C:H2'	26:1J:7:G:H8	1.84	0.42
29:29:92:THR:O	29:29:95:ILE:HG12	2.19	0.42
35:15:90:MET:HE2	35:15:90:MET:HA	2.01	0.42
36:25:104:ARG:O	36:25:108:GLU:HG2	2.20	0.42
41:75:36:GLU:HB3	41:75:39:ARG:NH2	2.32	0.42
47:D5:123:ASP:OD1	47:D5:123:ASP:N	2.52	0.42
1:13:607:A:H2'	1:13:608:A:H8	1.84	0.42
1:13:1014:A:P	1:13:1014:A:H8	2.43	0.42
1:13:1239:A:H62	1:13:1299:A:H62	1.68	0.42
1:13:1281:U:OP2	1:13:1282:C:N4	2.41	0.42
1:13:1320:C:OP2	19:AI:70:LYS:HE3	2.19	0.42
2:1E:24:TRP:CE3	2:1E:26:PRO:HA	2.54	0.42
25:1H:26:G:C6	25:1H:27:G:N1	2.87	0.42
25:1H:141(A):C:H2'	25:1H:142:G:O4'	2.19	0.42
25:1H:414:C:H1'	25:1H:1864:U:O2'	2.19	0.42
25:1H:673:C:H5''	30:31:81:PRO:HD2	2.00	0.42
25:1H:711:G:H1	25:1H:720:C:H42	1.68	0.42
25:1H:1287:A:C8	39:98:107:ASP:HB2	2.54	0.42
25:1H:1291:C:H2'	25:1H:1292:U:H6	1.85	0.42
25:1H:2810:A:O3'	29:21:61:ARG:HD3	2.19	0.42
25:1H:2815:C:H2'	25:1H:2816:C:H6	1.83	0.42
26:16:80:U:H2'	26:16:81:G:H21	1.84	0.42
28:11:17:THR:CG2	28:11:204:ILE:HA	2.50	0.42
30:31:65:TRP:CZ3	30:31:72:ARG:HB3	2.54	0.42
33:61:133:HIS:ND1	33:61:133:HIS:N	2.67	0.42
34:38:7:VAL:O	34:38:11:ALA:HB2	2.19	0.42
35:58:53:VAL:HA	35:58:121:LYS:O	2.19	0.42
35:58:70:LYS:O	35:58:70:LYS:HG3	2.15	0.42
35:58:76:SER:O	35:58:78:TYR:N	2.53	0.42
37:78:55:ARG:O	37:78:56:SER:OG	2.31	0.42
39:98:75:LEU:HA	39:98:78:LYS:HB3	2.00	0.42
44:E8:97:LYS:H	44:E8:97:LYS:HG2	1.64	0.42
45:F8:51:VAL:HG13	45:F8:81:VAL:HG23	2.01	0.42
50:K8:50:ILE:H	50:K8:50:ILE:HG13	1.51	0.42
1:1G:327:A:O2'	1:1G:328:C:O4'	2.28	0.42
1:1G:765:G:H5''	1:1G:766:A:OP1	2.19	0.42
1:1G:1198:G:H2'	1:1G:1199:U:C6	2.55	0.42
1:1G:1250:A:H4'	9:82:68:GLY:N	2.34	0.42
1:1G:1263:C:N3	1:1G:1273:G:N2	2.68	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:72:11:THR:HG22	8:72:15:ASN:HD21	1.84	0.42
13:4A:70:LEU:HD13	13:4A:71:ARG:N	2.35	0.42
18:9A:56:THR:HB	18:9A:58:LEU:HG	2.00	0.42
20:BA:33:ILE:O	20:BA:37:SER:OG	2.30	0.42
25:14:298:G:OP2	62:14:3558:HOH:O	2.21	0.42
25:14:1879:C:H2'	25:14:1880:C:O4'	2.19	0.42
25:14:1999:C:H4'	25:14:2723:C:O2	2.19	0.42
37:35:25:SER:C	37:35:27:HIS:H	2.27	0.42
37:35:57:THR:HG21	37:35:60:MET:CB	2.43	0.42
43:95:71:LEU:CA	43:95:86:GLY:H	2.30	0.42
46:C5:31:LEU:O	46:C5:35:TYR:N	2.51	0.42
47:D5:133:ILE:H	47:D5:133:ILE:HG13	1.68	0.42
47:D5:150:LEU:HB2	47:D5:169:GLU:O	2.19	0.42
56:M5:36:LYS:O	56:M5:37:SER:C	2.62	0.42
1:13:437:U:H5''	4:3E:155:LEU:HD11	2.01	0.42
1:13:565:U:H5''	1:13:566:G:C2'	2.48	0.42
1:13:936:C:N4	1:13:1379:G:H1	2.15	0.42
1:13:944:G:N2	1:13:1338:G:C8	2.88	0.42
1:13:1020:U:H2'	1:13:1021:G:H8	1.85	0.42
1:13:1410:G:H2'	1:13:1411:C:H6	1.84	0.42
1:13:1502:A:H2	1:13:1505:G:N1	2.05	0.42
2:1E:80:ILE:HD13	2:1E:212:GLN:HB2	2.02	0.42
7:6E:5:ARG:HB3	7:6E:7:ALA:N	2.33	0.42
12:3I:50:ARG:HB2	12:3I:90:LEU:HD11	2.02	0.42
18:9I:72:ARG:O	18:9I:76:LEU:HD22	2.20	0.42
25:1H:443:A:C3'	30:31:45:ARG:HH12	2.31	0.42
25:1H:869:G:H2'	25:1H:870:A:H8	1.85	0.42
25:1H:978:G:H1	25:1H:985:C:H42	1.67	0.42
25:1H:1011:G:OP1	42:C8:75:ASN:HB3	2.19	0.42
25:1H:1628:G:H2'	25:1H:1629:U:C6	2.54	0.42
25:1H:2107:C:H42	25:1H:2182:G:H1	1.67	0.42
25:1H:2447:G:OP2	62:1H:3696:HOH:O	2.21	0.42
25:1H:2643:G:H2'	25:1H:2644:G:O4'	2.19	0.42
27:71:46:LYS:HG2	27:71:210:ARG:HB2	2.01	0.42
28:11:62:TYR:HA	28:11:87:ASN:OD1	2.19	0.42
28:11:164:GLN:CD	28:11:176:ARG:HH21	2.27	0.42
31:41:150:ASP:OD1	31:41:150:ASP:N	2.53	0.42
34:38:75:GLN:O	34:38:111:LEU:HB3	2.19	0.42
36:68:88:ASN:O	36:68:91:LEU:N	2.47	0.42
47:H8:138:GLU:C	47:H8:156:LYS:HZ3	2.27	0.42
47:H8:170:THR:O	47:H8:172:ALA:N	2.52	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:I8:49:LYS:H	48:I8:80:HIS:HD1	1.67	0.42
53:N8:9:LYS:HD3	53:N8:9:LYS:HA	1.94	0.42
1:1G:329:A:O2'	1:1G:332:G:N7	2.51	0.42
1:1G:735:C:H2'	1:1G:736:C:C6	2.48	0.42
1:1G:1055:A:O2'	3:22:156:ARG:NH1	2.52	0.42
1:1G:1144:G:N2	1:1G:1146:A:H62	2.18	0.42
7:62:9:VAL:HG22	7:62:94:ARG:NH1	2.35	0.42
14:5A:23:ARG:NH1	14:5A:28:GLY:O	2.48	0.42
18:9A:59:SER:OG	18:9A:60:ALA:N	2.53	0.42
25:14:300:A:C8	46:C5:84:ARG:NH2	2.87	0.42
25:14:528:A:H2	25:14:2042:A:H2'	1.81	0.42
25:14:1255:U:C5	30:39:73:ALA:HA	2.55	0.42
25:14:1412:A:H2'	25:14:1413:G:C8	2.54	0.42
25:14:1525:G:H2'	25:14:1526:G:C8	2.53	0.42
25:14:1567:A:H5'	28:19:86:PRO:HG3	2.01	0.42
25:14:2625:G:H2'	25:14:2626:C:O4'	2.20	0.42
25:14:2808:U:H5'	25:14:2891:G:O6	2.19	0.42
29:29:63:LEU:HD23	29:29:63:LEU:HA	1.93	0.42
31:49:23:PHE:HB2	31:49:25:TYR:CE1	2.55	0.42
31:49:75:LYS:HG3	31:49:76:SER:H	1.85	0.42
31:49:126:ASP:OD2	31:49:130:ASN:ND2	2.42	0.42
32:59:23:ARG:HA	32:59:37:VAL:H	1.84	0.42
32:59:87:LEU:HA	32:59:163:TYR:O	2.19	0.42
37:35:52:GLU:CD	37:35:54:GLY:H	2.28	0.42
47:D5:152:ALA:HA	47:D5:155:LEU:HG	2.01	0.42
47:D5:173:ALA:C	47:D5:175:VAL:H	2.28	0.42
1:13:637:G:H2'	1:13:638:G:C8	2.54	0.42
1:13:1359:C:H3'	14:5I:35:ARG:HH12	1.85	0.42
1:13:1367:C:H5'	10:1I:60:ARG:NH1	2.34	0.42
1:13:1387:G:H2'	1:13:1388:C:C6	2.55	0.42
4:3E:88:VAL:HG12	4:3E:92:VAL:HG22	2.01	0.42
8:7E:1:MET:HE2	8:7E:1:MET:N	2.34	0.42
10:1I:81:THR:HA	10:1I:84:GLN:HE22	1.84	0.42
17:8I:21:VAL:HG11	17:8I:59:ILE:HD12	2.01	0.42
18:9I:29:PHE:N	18:9I:29:PHE:HD1	2.16	0.42
25:1H:341:G:H2'	25:1H:342:G:O4'	2.20	0.42
25:1H:767:U:H2'	25:1H:768:G:H8	1.84	0.42
25:1H:1113:U:OP1	32:51:2:SER:N	2.52	0.42
25:1H:1166:C:H2'	25:1H:1167:U:C6	2.55	0.42
25:1H:1204:A:HO2'	25:1H:1205:U:P	2.41	0.42
25:1H:1341:U:O4	45:F8:16:LYS:NZ	2.47	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:1396:U:O2	25:1H:1396:U:H2'	2.19	0.42
25:1H:1468:C:H2'	25:1H:1469:A:H8	1.85	0.42
25:1H:1913:A:H4'	25:1H:1914:C:H5''	2.01	0.42
25:1H:2128:C:OP2	25:1H:2128:C:H6	2.02	0.42
25:1H:2373:G:H2'	25:1H:2374:C:C6	2.54	0.42
25:1H:2392:A:OP2	25:1H:2422:A:N6	2.51	0.42
62:1H:3862:HOH:O	49:J8:17:SER:HA	2.20	0.42
27:71:29:VAL:HG21	27:71:41:VAL:HG22	2.02	0.42
41:B8:65:LYS:NZ	41:B8:65:LYS:HB2	2.35	0.42
44:E8:38:TYR:OH	53:N8:40:LYS:HG3	2.20	0.42
46:G8:63:LYS:HD2	46:G8:63:LYS:HA	1.91	0.42
1:1G:17:U:H2'	1:1G:18:C:C6	2.55	0.42
1:1G:129(A):G:C6	1:1G:191(A):G:H1'	2.55	0.42
1:1G:686:U:H1'	11:2A:42:TRP:HE1	1.84	0.42
1:1G:928:G:H2'	1:1G:929:G:C8	2.55	0.42
1:1G:1157:A:O2'	1:1G:1158:C:P	2.78	0.42
2:12:205:ASP:OD1	2:12:206:ASP:N	2.52	0.42
9:82:27:THR:OG1	9:82:28:VAL:N	2.52	0.42
10:1A:15:THR:HA	10:1A:18:ALA:HB3	2.02	0.42
20:BA:28:ALA:HA	20:BA:31:SER:HB2	2.02	0.42
22:3L:34:U:H2'	22:3L:35:C:H3'	2.02	0.42
25:14:34:C:O2'	25:14:35:G:C8	2.71	0.42
25:14:270(D):C:H2'	25:14:270(E):G:C8	2.55	0.42
25:14:625:G:O6	37:35:107:LYS:HD3	2.20	0.42
25:14:833:U:O2'	37:35:55:ARG:NH1	2.53	0.42
25:14:935:C:H2'	25:14:936:C:H6	1.85	0.42
30:39:34:TRP:HD1	37:35:6:LEU:HB3	1.85	0.42
31:49:111:LEU:HD23	31:49:111:LEU:HA	1.86	0.42
35:15:14:VAL:HG12	35:15:51:PHE:O	2.20	0.42
41:75:57:PHE:O	41:75:58:ASN:CG	2.63	0.42
42:85:92:ARG:O	42:85:93:LYS:C	2.63	0.42
47:D5:100:VAL:N	47:D5:124:ILE:O	2.52	0.42
1:13:191(C):G:H2'	1:13:191(D):U:O4'	2.20	0.42
1:13:484:G:O2'	1:13:485:G:OP2	2.28	0.42
1:13:687:A:N1	1:13:700:G:O2'	2.49	0.42
1:13:1461:G:H2'	1:13:1462:G:H8	1.84	0.42
2:1E:16:HIS:HE1	2:1E:209:ARG:HH21	1.67	0.42
3:2E:39:ILE:HG22	3:2E:43:LEU:HD11	2.02	0.42
4:3E:121:VAL:HG22	4:3E:126:ILE:HD12	2.00	0.42
17:8I:18:THR:OG1	17:8I:69:LYS:NZ	2.26	0.42
25:1H:933:A:OP1	51:L8:24:LYS:NZ	2.36	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:1021:A:C8	25:1H:1022:G:H5''	2.48	0.42
25:1H:1035:U:H2'	25:1H:1036:G:C8	2.55	0.42
25:1H:1278:A:H2'	25:1H:1279:G:C8	2.54	0.42
25:1H:1333:C:H2'	25:1H:1334:G:H8	1.85	0.42
25:1H:1368:G:C2	25:1H:1369:G:C8	3.08	0.42
25:1H:1469:A:H2'	25:1H:1470:G:O4'	2.19	0.42
25:1H:1511:A:H2'	25:1H:1512:G:O4'	2.19	0.42
25:1H:1869:G:N2	25:1H:1872:A:OP2	2.47	0.42
25:1H:2119:A:N6	25:1H:2170:A:C6	2.88	0.42
25:1H:2688:U:H1'	25:1H:2721:A:H61	1.85	0.42
25:1H:2867:G:O2'	25:1H:2868:A:H8	2.03	0.42
27:71:190:ARG:HA	27:71:193:ILE:HB	2.01	0.42
39:98:41:ALA:O	39:98:44:LEU:N	2.53	0.42
47:H8:137:ILE:HG22	47:H8:158:PRO:HG2	2.00	0.42
50:K8:52:ASP:HA	50:K8:55:ARG:HB2	2.02	0.42
52:M8:47:GLN:C	52:M8:49:PHE:H	2.28	0.42
53:N8:16:ARG:HG3	53:N8:17:ASP:N	2.35	0.42
1:1G:974:A:P	14:5A:41:ARG:HH12	2.43	0.42
1:1G:1037:C:H2'	1:1G:1038:C:C6	2.53	0.42
1:1G:1142:G:H3'	1:1G:1143:G:H8	1.85	0.42
1:1G:1376:U:O4	7:62:10:ARG:NH1	2.52	0.42
1:1G:1483:A:H1'	25:14:1948:G:H1'	2.01	0.42
2:12:52:GLU:O	2:12:56:ARG:HG3	2.20	0.42
10:1A:8:LEU:HD22	10:1A:20:ALA:HB2	2.02	0.42
11:2A:56:GLY:O	11:2A:89:ALA:HB3	2.20	0.42
18:9A:23:LYS:O	18:9A:23:LYS:HG3	2.19	0.42
22:3L:61:U:H3'	22:3L:62:C:C5	2.55	0.42
24:4L:19:A:H2'	24:4L:20:A:C8	2.55	0.42
25:14:84:A:H61	25:14:102:G:H1'	1.85	0.42
25:14:506:G:O3'	25:14:507:A:H8	2.03	0.42
25:14:848:G:H2'	25:14:849:A:H8	1.81	0.42
25:14:1535:U:C5	25:14:1536:A:H1'	2.55	0.42
25:14:1853:A:H2'	25:14:1854:A:H8	1.83	0.42
25:14:2615:U:C2	53:J5:7:PRO:HA	2.54	0.42
25:14:2820:A:C4	39:55:4:LEU:HD11	2.55	0.42
26:1J:21:G:H2'	26:1J:22:U:O4'	2.20	0.42
28:19:72:LYS:HD3	28:19:97:TYR:CE1	2.54	0.42
30:39:103:LYS:HA	30:39:106:ARG:HG3	2.01	0.42
32:59:74:ASN:HB3	32:59:138:LYS:HB2	2.02	0.42
39:55:107:ASP:O	39:55:107:ASP:CG	2.63	0.42
40:65:51:ALA:HB2	40:65:76:LYS:HD3	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:85:40:PHE:HB3	43:95:75:PHE:CD2	2.55	0.42
47:D5:24:LEU:HD23	47:D5:41:LEU:HG	2.02	0.42
47:D5:81:ARG:HE	47:D5:81:ARG:HB2	1.71	0.42
47:D5:99:TYR:HA	47:D5:124:ILE:O	2.20	0.42
49:F5:80:LEU:HD22	49:F5:80:LEU:H	1.85	0.42
1:13:781:A:C8	1:13:782:A:C8	3.08	0.42
1:13:901:A:C5	1:13:902:G:H1'	2.55	0.42
1:13:947:G:H2'	1:13:948:C:O4'	2.19	0.42
1:13:991:U:O2'	1:13:992:U:P	2.78	0.42
1:13:1035:A:N6	1:13:1036:G:N3	2.68	0.42
1:13:1274:G:N2	1:13:1275:A:N7	2.58	0.42
1:13:1312:G:H1	1:13:1325:C:N4	2.13	0.42
2:1E:74:LYS:HG3	2:1E:77:ALA:HB3	2.02	0.42
3:2E:47:LEU:HG	3:2E:52:LEU:HD22	2.02	0.42
6:5E:47:ARG:NH1	6:5E:57:GLN:HB3	2.35	0.42
11:2I:32:ILE:HG12	11:2I:41:THR:O	2.20	0.42
20:BI:22:ARG:O	20:BI:26:ASN:ND2	2.53	0.42
25:1H:270(G):C:H42	25:1H:270(S):G:H1	1.67	0.42
25:1H:746:A:HO2'	25:1H:747:U:P	2.38	0.42
25:1H:1068:G:O2'	25:1H:1096:A:H1'	2.20	0.42
25:1H:1448:G:H2'	25:1H:1449:A:C8	2.55	0.42
25:1H:1750:G:O2'	25:1H:2860:A:N1	2.40	0.42
25:1H:2163:C:H2'	25:1H:2164:C:H6	1.85	0.42
29:21:3:GLY:HA2	29:21:199:ARG:HA	2.00	0.42
30:31:159:GLY:HA2	30:31:164:ARG:HH21	1.85	0.42
42:C8:90:VAL:HG11	43:D8:39:LEU:HD13	2.02	0.42
46:G8:81:LYS:HD3	46:G8:81:LYS:HA	1.76	0.42
52:M8:39:CYS:C	52:M8:41:PRO:HD3	2.45	0.42
1:1G:270:A:C5	1:1G:271:C:C4	3.08	0.42
1:1G:1025:U:H2'	1:1G:1026:G:C8	2.54	0.42
1:1G:1038:C:O2'	1:1G:1039:C:P	2.77	0.42
1:1G:1289:A:H5''	21:1B:10:ARG:HH21	1.85	0.42
1:1G:1315:U:H2'	1:1G:1316:G:O4'	2.19	0.42
7:62:146:GLU:C	7:62:148:ASN:H	2.28	0.42
19:AA:20:LEU:HD23	19:AA:44:MET:HE1	2.01	0.42
25:14:124:G:H5'	25:14:1376:C:O2'	2.19	0.42
25:14:186:G:H2'	25:14:187:G:H8	1.84	0.42
25:14:748:G:C8	44:A5:89:ALA:HB1	2.55	0.42
25:14:829:A:C8	25:14:2248:C:H5'	2.55	0.42
25:14:840:C:H2'	25:14:841:A:H8	1.83	0.42
25:14:923:C:H2'	25:14:924:C:C6	2.55	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:1203:G:N1	25:14:1241:A:OP2	2.50	0.42
25:14:1839:G:C8	25:14:1927:A:H1'	2.54	0.42
25:14:2267:A:H5''	25:14:2268:A:H5'	2.01	0.42
25:14:2554:U:H2'	25:14:2555:U:C6	2.55	0.42
28:19:249:PRO:HD2	28:19:250:TRP:CZ3	2.55	0.42
29:29:33:VAL:HG12	29:29:89:ASP:O	2.19	0.42
30:39:21:ALA:O	30:39:23:ASP:N	2.53	0.42
32:59:131:VAL:HG12	32:59:132:ARG:H	1.84	0.42
38:45:17:LEU:HD22	38:45:96:VAL:HG13	2.01	0.42
39:55:73:VAL:O	39:55:76:VAL:HG12	2.20	0.42
43:95:35:LEU:HB2	43:95:37:VAL:CG1	2.50	0.42
48:E5:29:GLN:O	48:E5:67:VAL:HG23	2.20	0.42
1:13:1074:G:C2	1:13:1075:C:C2	3.08	0.42
1:13:1137:C:O2'	1:13:1138:G:H5''	2.19	0.42
6:5E:19:LEU:HD11	6:5E:59:TYR:CD2	2.54	0.42
9:8E:48:GLU:N	9:8E:49:PRO:HD2	2.35	0.42
25:1H:870:A:OP1	38:88:5:ARG:NH2	2.53	0.42
25:1H:1087:G:H2'	25:1H:1089:G:O4'	2.20	0.42
25:1H:1675:C:H2'	25:1H:1676:A:O4'	2.20	0.42
25:1H:2817:G:H1	25:1H:2829:C:H42	1.67	0.42
29:21:8:LYS:HB2	29:21:196:VAL:HG21	2.02	0.42
41:B8:86:ILE:HD12	41:B8:87:ASP:H	1.84	0.42
47:H8:48:PHE:HA	47:H8:51:ALA:CB	2.43	0.42
48:I8:70:GLN:O	48:I8:77:ARG:HG3	2.20	0.42
1:1G:520:A:N1	1:1G:536:C:H1'	2.35	0.42
1:1G:559:A:H4'	1:1G:560:U:H3'	2.01	0.42
1:1G:598:U:H2'	1:1G:599:C:C6	2.54	0.42
1:1G:729:A:H2'	1:1G:730:G:C8	2.51	0.42
1:1G:816:A:OP1	1:1G:1526:G:O2'	2.38	0.42
3:22:78:GLY:O	3:22:80:GLY:N	2.52	0.42
4:32:88:VAL:O	4:32:90:GLY:N	2.53	0.42
13:4A:14:ARG:HB2	13:4A:17:VAL:HG23	2.02	0.42
13:4A:60:VAL:HA	13:4A:64:TRP:HE1	1.83	0.42
18:9A:23:LYS:HA	18:9A:26:LEU:HD13	2.02	0.42
25:14:28:A:H2'	25:14:29:U:O4'	2.20	0.42
25:14:372:G:O2'	25:14:373:U:OP2	2.38	0.42
25:14:465:G:H2'	25:14:466:A:C8	2.55	0.42
25:14:1149:G:H2'	25:14:1150:C:C6	2.55	0.42
25:14:1491:G:H2'	25:14:1492:G:H8	1.85	0.42
25:14:2642:G:H1	25:14:2772:C:H42	1.67	0.42
28:19:208:LYS:HG3	28:19:211:ARG:H	1.84	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:29:181:LEU:HD12	29:29:181:LEU:HA	1.83	0.42
30:39:25:PRO:CD	30:39:115:ALA:HB1	2.50	0.42
30:39:125:LEU:HD13	30:39:199:TRP:CD1	2.55	0.42
41:75:36:GLU:HB3	41:75:39:ARG:HH12	1.84	0.42
42:85:96:ALA:C	42:85:98:LEU:N	2.78	0.42
1:13:177:C:H2'	1:13:178:C:H6	1.85	0.41
1:13:464:G:C6	1:13:466:C:H5'	2.55	0.41
1:13:501:C:H1'	1:13:549:C:H1'	2.02	0.41
1:13:748:C:H1'	1:13:749:C:OP2	2.19	0.41
1:13:919:A:O5'	1:13:919:A:H8	2.03	0.41
1:13:991:U:C4	1:13:1212:U:H1'	2.54	0.41
1:13:1095:U:P	1:13:1108:G:H1	2.42	0.41
1:13:1316:G:H5''	14:51:17:LYS:HE3	2.01	0.41
4:3E:148:VAL:HG23	4:3E:181:MET:HB3	2.01	0.41
15:6I:29:VAL:HB	15:6I:81:LEU:HD21	2.02	0.41
18:9I:29:PHE:HD1	18:9I:29:PHE:H	1.68	0.41
25:1H:51:G:H1'	25:1H:119:A:N1	2.34	0.41
25:1H:322:A:P	30:31:168:ARG:HH21	2.43	0.41
25:1H:669:G:H2'	25:1H:669:G:N3	2.34	0.41
25:1H:680:G:H2'	25:1H:681:G:C8	2.55	0.41
25:1H:719:C:H2'	25:1H:720:C:C6	2.54	0.41
25:1H:770:G:OP2	62:1H:3700:HOH:O	2.21	0.41
25:1H:1444(A):A:O2'	25:1H:1445:C:OP1	2.35	0.41
25:1H:1507:A:H2'	25:1H:1508:A:O4'	2.20	0.41
25:1H:1510:A:H2'	25:1H:1511:A:H8	1.85	0.41
28:11:111:LEU:HD22	28:11:115:GLN:CD	2.45	0.41
30:31:34:TRP:CD1	37:78:8:PRO:HD3	2.55	0.41
35:58:115:ARG:HA	35:58:118:LYS:HB2	2.02	0.41
44:E8:38:TYR:HE2	53:N8:41:PRO:HD3	1.82	0.41
45:F8:27:THR:HB	45:F8:80:ILE:HG22	2.02	0.41
47:H8:56:VAL:HA	47:H8:70:LEU:HD12	2.02	0.41
47:H8:98:MET:HB2	47:H8:98:MET:HE3	1.78	0.41
50:K8:2:LYS:HE3	50:K8:6:VAL:HB	2.02	0.41
51:L8:7:LYS:NZ	51:L8:32:GLN:HG3	2.35	0.41
56:Q8:40:GLU:OE2	56:Q8:44:LYS:HE2	2.19	0.41
1:1G:458:C:N4	1:1G:474:G:H1	2.16	0.41
1:1G:1351:U:H2'	1:1G:1352:C:C6	2.56	0.41
2:12:167:PRO:O	2:12:170:GLU:N	2.53	0.41
3:22:180:ALA:HB1	3:22:182:ILE:HG13	2.02	0.41
4:32:31:CYS:O	4:32:35:ARG:HG3	2.20	0.41
4:32:36:ARG:HD2	4:32:38:TYR:OH	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:3A:31:ARG:HG2	12:3A:32:GLY:N	2.35	0.41
13:4A:80:ARG:NH1	52:I5:55:ARG:HD3	2.35	0.41
13:4A:83:ASP:OD1	13:4A:83:ASP:N	2.49	0.41
25:14:67:U:H2'	25:14:68:G:C8	2.55	0.41
25:14:185:U:H4'	25:14:218:A:H4'	2.02	0.41
25:14:710:G:H2'	25:14:711:G:H8	1.84	0.41
25:14:778:G:H5''	28:19:48:ARG:HD2	2.02	0.41
25:14:1055:G:N2	25:14:1086:A:O4'	2.53	0.41
25:14:1220:A:OP2	42:85:19:LYS:NZ	2.49	0.41
25:14:1287:A:C6	25:14:1288:U:N3	2.88	0.41
25:14:1678:G:N2	25:14:1989:G:N2	2.68	0.41
25:14:2578:G:C5	29:29:140:SER:HB2	2.55	0.41
25:14:2695:C:H2'	25:14:2696:U:C6	2.55	0.41
25:14:2859:G:H2'	25:14:2860:A:C8	2.54	0.41
26:1J:82:G:H8	26:1J:82:G:O5'	2.03	0.41
28:19:245:PRO:HA	28:19:246:PRO:HD3	1.96	0.41
29:29:5:LEU:HD23	29:29:197:ILE:HG12	2.01	0.41
29:29:39:PRO:HG3	29:29:45:THR:OG1	2.20	0.41
31:49:120:LEU:HB2	31:49:179:PRO:O	2.20	0.41
37:35:21:ARG:O	37:35:28:GLY:HA2	2.20	0.41
39:55:87:TYR:HE2	39:55:117:VAL:HG13	1.85	0.41
40:65:83:LYS:C	40:65:83:LYS:HD3	2.45	0.41
56:M5:37:SER:O	56:M5:39:LYS:N	2.53	0.41
1:13:35:G:OP1	12:3I:101:VAL:HG11	2.20	0.41
1:13:104:G:OP1	20:BI:21:LYS:NZ	2.49	0.41
1:13:280:C:C2	17:8I:38:ARG:HG3	2.55	0.41
1:13:876:G:H1'	8:7E:11:THR:HG21	2.02	0.41
1:13:939:G:H5''	7:6E:102:ARG:CZ	2.50	0.41
1:13:999:U:O2'	25:14:2137:C:H5''	2.20	0.41
1:13:1003:G:H1	1:13:1037:C:H42	1.68	0.41
1:13:1112:C:H1'	3:2E:179:ARG:HH11	1.85	0.41
1:13:1159:U:H1'	1:13:1182:G:N2	2.34	0.41
1:13:1429:C:H2'	1:13:1430:C:C6	2.55	0.41
4:3E:100:ARG:NH2	4:3E:137:SER:HB3	2.35	0.41
4:3E:173:TRP:CE3	4:3E:189:PRO:HB3	2.55	0.41
6:5E:61:LEU:HD23	6:5E:63:TYR:HE1	1.84	0.41
9:8E:91:ASP:C	9:8E:93:ARG:H	2.28	0.41
25:1H:270(I):G:H22	25:1H:270(R):G:H1'	1.85	0.41
25:1H:708:C:H5'	25:1H:709:U:OP2	2.20	0.41
25:1H:828:U:H4'	25:1H:831:G:N1	2.35	0.41
25:1H:2227:A:H2'	25:1H:2228:G:O4'	2.19	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:2405:G:H8	25:1H:2405:G:O5'	2.03	0.41
25:1H:2846:G:OP2	41:B8:54:ARG:HB2	2.20	0.41
26:16:81:G:C2	26:16:82:G:N7	2.88	0.41
28:11:61:LEU:HD13	28:11:61:LEU:HA	1.84	0.41
28:11:112:GLN:H	28:11:115:GLN:HE22	1.68	0.41
30:31:81:PRO:CB	30:31:89:VAL:HG22	2.49	0.41
32:51:28:GLY:HA3	32:51:79:VAL:HB	2.02	0.41
35:58:34:LEU:HD13	35:58:34:LEU:HA	1.85	0.41
41:B8:108:ARG:HA	41:B8:111:ARG:CZ	2.50	0.41
41:B8:117:ASP:O	41:B8:121:ILE:HG13	2.19	0.41
42:C8:59:ARG:O	42:C8:63:VAL:HG23	2.20	0.41
47:H8:10:ARG:NH2	47:H8:37:VAL:O	2.53	0.41
47:H8:53:ILE:HG22	47:H8:71:VAL:HG13	2.02	0.41
48:I8:40:GLN:NE2	48:I8:43:THR:HA	2.35	0.41
51:L8:52:HIS:CD2	51:L8:53:LEU:HG	2.55	0.41
55:P8:30:VAL:O	55:P8:34:ARG:HG3	2.20	0.41
1:1G:201:C:O2'	1:1G:208:U:P	2.78	0.41
1:1G:210:U:O2'	1:1G:216:G:O5'	2.38	0.41
1:1G:374:A:C6	1:1G:375:U:C4	3.08	0.41
1:1G:559:A:H4'	1:1G:560:U:C5'	2.50	0.41
1:1G:1211:U:H4'	1:1G:1212:U:OP1	2.19	0.41
1:1G:1298:C:H5	7:62:114:ARG:HD2	1.85	0.41
1:1G:1502:A:H5''	1:1G:1504:G:C8	2.55	0.41
3:22:4:LYS:HZ2	3:22:4:LYS:HG3	1.64	0.41
4:32:21:LEU:HD23	4:32:21:LEU:HA	1.89	0.41
6:52:47:ARG:HG3	6:52:57:GLN:HG2	2.02	0.41
6:52:50:TYR:OH	18:9A:74:ARG:O	2.32	0.41
15:6A:82:ILE:HD11	15:6A:88:ARG:HB2	2.01	0.41
25:14:863:A:O2'	25:14:864:G:H5'	2.20	0.41
25:14:1106:G:H3'	25:14:1107:G:C8	2.55	0.41
25:14:1442:G:H5'	25:14:1628:G:H5''	2.02	0.41
25:14:2403:C:H2'	25:14:2404:C:C6	2.55	0.41
25:14:2774:C:H2'	25:14:2775:A:O4'	2.19	0.41
33:69:3:VAL:O	33:69:18:VAL:HA	2.20	0.41
38:45:98:LYS:HB3	38:45:99:PRO:HD2	2.02	0.41
38:45:134:ARG:HH21	47:D5:122:ARG:CZ	2.33	0.41
39:55:54:LEU:HG	39:55:62:ALA:HB1	2.02	0.41
45:B5:12:VAL:HG12	45:B5:29:TRP:NE1	2.35	0.41
47:D5:148:ASP:C	47:D5:172:ALA:HB3	2.45	0.41
51:H5:19:GLN:O	51:H5:23:LEU:HD12	2.20	0.41
52:I5:40:HIS:HA	52:I5:44:THR:HB	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:K5:27:LYS:O	54:K5:28:ARG:HB2	2.20	0.41
1:13:17:U:H1'	1:13:1080:A:H1'	2.02	0.41
1:13:91:C:H2'	1:13:92:G:O4'	2.20	0.41
1:13:854:G:O6	62:13:1809:HOH:O	2.22	0.41
1:13:977:A:H4'	1:13:981:U:O2	2.20	0.41
1:13:1409:C:H2'	1:13:1410:G:C8	2.55	0.41
25:1H:275:G:H4'	25:1H:275:G:OP1	2.19	0.41
25:1H:319:C:H2'	25:1H:320:A:O4'	2.21	0.41
25:1H:580:C:H2'	25:1H:581:C:C6	2.54	0.41
25:1H:631:A:O2'	37:78:67:MET:HB3	2.19	0.41
25:1H:767:U:H2'	25:1H:768:G:C8	2.56	0.41
25:1H:869:G:H1	25:1H:908:C:H42	1.67	0.41
25:1H:1025:G:C4	25:1H:1135:C:H1'	2.56	0.41
25:1H:1497:U:H5''	25:1H:1498:C:C5	2.55	0.41
25:1H:1978:A:H2'	25:1H:1979:C:O4'	2.21	0.41
25:1H:2663:G:H3'	25:1H:2664:G:H8	1.85	0.41
25:1H:2849:U:H4'	25:1H:2868:A:C2	2.55	0.41
25:1H:2867:G:O2'	25:1H:2868:A:C8	2.73	0.41
26:16:103:U:O3'	47:H8:72:ARG:HD3	2.21	0.41
31:41:138:GLN:C	31:41:140:ILE:H	2.28	0.41
31:41:161:THR:HG21	31:41:172:LEU:HD22	2.01	0.41
31:41:174:GLU:HG2	31:41:180:PHE:CD1	2.54	0.41
32:51:24:VAL:HG23	32:51:35:VAL:HB	2.02	0.41
44:E8:35:ILE:HG23	53:N8:28:PRO:HD2	2.03	0.41
45:F8:41:ASN:OD1	45:F8:41:ASN:N	2.53	0.41
46:G8:4:LYS:HD3	46:G8:4:LYS:HA	1.71	0.41
47:H8:58:VAL:C	47:H8:60:GLU:H	2.29	0.41
53:N8:40:LYS:HD3	53:N8:46:CYS:SG	2.60	0.41
54:O8:6:ARG:HA	54:O8:6:ARG:HH11	1.85	0.41
1:1G:1126:U:H3'	1:1G:1280:A:C8	2.55	0.41
1:1G:1249:C:O2'	9:82:73:GLN:OE1	2.37	0.41
1:1G:1301:U:O2	1:1G:1301:U:H2'	2.19	0.41
4:32:96:LEU:HG	4:32:139:ARG:NH1	2.34	0.41
6:52:7:ASN:OD1	6:52:62:TRP:HD1	2.04	0.41
15:6A:26:GLU:OE2	15:6A:77:ARG:HD2	2.20	0.41
16:7A:45:THR:HG23	16:7A:48:TRP:HB3	2.02	0.41
19:AA:25:LYS:HD2	19:AA:25:LYS:C	2.46	0.41
25:14:274:G:N2	25:14:363:G:C2	2.88	0.41
25:14:321:G:H5'	30:39:134:GLY:O	2.20	0.41
25:14:776:G:N7	25:14:793:A:O2'	2.53	0.41
25:14:957:A:H5''	38:45:14:ARG:HH22	1.86	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:1729:A:N7	25:14:1731:G:C2	2.89	0.41
25:14:2577:A:O4'	53:J5:3:LYS:HB2	2.20	0.41
25:14:2875:C:O2'	41:75:2:ASN:OD1	2.23	0.41
29:29:41:LYS:HE3	29:29:41:LYS:HB2	1.91	0.41
29:29:108:SER:HB3	29:29:165:VAL:HG21	2.02	0.41
30:39:122:LYS:HD2	30:39:191:ARG:HG2	2.01	0.41
30:39:170:LEU:HD23	30:39:170:LEU:HA	1.95	0.41
33:69:26:ALA:HB1	33:69:31:LEU:HD13	2.01	0.41
33:69:125:GLU:HA	33:69:141:LYS:HB3	2.02	0.41
37:35:77:ARG:HB2	37:35:78:PRO:HD2	2.03	0.41
39:55:65:LEU:O	39:55:68:ARG:HB2	2.20	0.41
43:95:17:GLY:N	43:95:96:ILE:O	2.48	0.41
46:C5:54:LYS:HD2	46:C5:55:TYR:CD1	2.56	0.41
47:D5:7:ALA:HB3	47:D5:59:LEU:HD12	2.02	0.41
47:D5:102:LEU:HB3	47:D5:139:VAL:HB	2.01	0.41
1:13:468:A:H4'	16:7I:80:PHE:O	2.21	0.41
1:13:687:A:H4'	1:13:688:G:O5'	2.20	0.41
1:13:792:A:H2'	1:13:792:A:N3	2.35	0.41
1:13:859:A:OP2	1:13:869:G:N2	2.47	0.41
1:13:1187:G:H4'	9:8E:111:ARG:NH1	2.35	0.41
4:3E:13:ARG:HB2	4:3E:38:TYR:O	2.20	0.41
16:7I:6:LEU:HB3	16:7I:17:TYR:CD1	2.54	0.41
17:8I:27:PHE:CZ	17:8I:36:ILE:HD11	2.55	0.41
25:1H:270(R):G:H2'	25:1H:270(S):G:C8	2.56	0.41
25:1H:384:U:H2'	25:1H:385:C:H6	1.85	0.41
25:1H:476:G:N2	25:1H:478:A:H3'	2.35	0.41
25:1H:518:G:H4'	44:E8:18:ARG:NH1	2.35	0.41
25:1H:668:G:H2'	25:1H:670:A:H62	1.85	0.41
25:1H:1065:U:H1'	25:1H:1074:G:N2	2.35	0.41
25:1H:1331:A:C6	25:1H:1333:C:C2	3.09	0.41
25:1H:1512:G:H2'	25:1H:1513:C:H6	1.84	0.41
25:1H:1841:U:H2'	25:1H:1842:G:C8	2.56	0.41
30:31:51:THR:HB	30:31:88:VAL:HG21	2.02	0.41
31:41:54:GLU:CD	31:41:54:GLU:H	2.27	0.41
34:38:18:GLU:OE1	34:38:66:LEU:HD13	2.20	0.41
34:38:27:VAL:HG22	34:38:111:LEU:N	2.34	0.41
41:B8:1:MET:HB3	41:B8:1:MET:HE2	1.64	0.41
43:D8:49:THR:O	43:D8:50:PRO:C	2.63	0.41
1:1G:59:A:H1'	1:1G:354:G:N2	2.34	0.41
1:1G:664:G:OP1	18:9A:64:ARG:NH2	2.37	0.41
1:1G:714:G:H2'	1:1G:715:A:C8	2.55	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:757:U:H2'	1:1G:758:G:O4'	2.20	0.41
1:1G:981:U:H5'	14:5A:21:TYR:CE2	2.55	0.41
1:1G:1075:C:H4'	2:12:175:ARG:HH12	1.86	0.41
1:1G:1418:A:H2	25:14:1948:G:N3	2.18	0.41
8:72:124:ALA:O	8:72:128:GLY:N	2.52	0.41
13:4A:105:THR:OG1	13:4A:106:ASN:N	2.52	0.41
19:AA:7:LYS:HE3	19:AA:7:LYS:H	1.86	0.41
19:AA:36:ARG:HG3	19:AA:72:GLY:N	2.31	0.41
25:14:273(F):C:H2'	25:14:274:G:H5''	2.02	0.41
25:14:293:U:O2	25:14:348:G:N2	2.53	0.41
25:14:527:C:O5'	25:14:2779:U:H5	2.03	0.41
25:14:957:A:H4'	38:45:74:TYR:OH	2.20	0.41
25:14:1405:U:H2'	25:14:1406:U:C6	2.54	0.41
25:14:1449:A:H8	25:14:1449:A:OP2	2.03	0.41
25:14:2611:U:C5	53:J5:3:LYS:HG2	2.54	0.41
28:19:39:LYS:HE2	28:19:39:LYS:HB3	1.78	0.41
30:39:107:LYS:HA	30:39:107:LYS:HD3	1.76	0.41
37:35:60:MET:HE3	37:35:60:MET:HB3	1.70	0.41
39:55:63:ARG:HA	39:55:80:PHE:CZ	2.55	0.41
40:65:15:ARG:NH2	40:65:90:GLY:HA2	2.36	0.41
42:85:98:LEU:C	42:85:100:VAL:N	2.73	0.41
43:95:21:ARG:NE	43:95:91:TYR:HB3	2.35	0.41
52:I5:22:ILE:O	52:I5:23:GLU:HB2	2.20	0.41
1:13:186(E):C:H2'	1:13:186(F):C:O4'	2.20	0.41
1:13:1073:U:O2'	2:1E:104:ASN:ND2	2.53	0.41
1:13:1250:A:H2	1:13:1370:G:H1'	1.86	0.41
2:1E:103:THR:N	2:1E:176:GLU:OE1	2.40	0.41
4:3E:105:VAL:HG13	4:3E:110:PHE:HB2	2.01	0.41
6:5E:89:MET:HG3	18:9I:76:LEU:HD11	2.02	0.41
8:7E:7:ALA:HB2	8:7E:85:ARG:HD3	2.02	0.41
12:3I:19:SER:OG	12:3I:20:LYS:N	2.54	0.41
19:AI:71:LEU:HD23	19:AI:71:LEU:HA	1.91	0.41
25:1H:71:A:H2	45:F8:31:HIS:NE2	2.19	0.41
25:1H:272:G:H2'	25:1H:273:G:O4'	2.20	0.41
25:1H:1206:G:C6	25:1H:1207:C:C4	3.09	0.41
25:1H:1432:C:H2'	25:1H:1433:U:O4'	2.20	0.41
25:1H:1825:A:H2'	25:1H:1826:G:C8	2.55	0.41
25:1H:1885:A:H2'	25:1H:1886:C:O4'	2.20	0.41
25:1H:2195:C:H3'	62:1H:3660:HOH:O	2.19	0.41
25:1H:2285:C:H5	54:O8:27:LYS:HD2	1.85	0.41
25:1H:2466:C:N4	25:1H:2484:G:H1	2.17	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
27:71:163:PHE:HB3	27:71:192:PHE:HE1	1.86	0.41
28:11:79:VAL:HB	28:11:114:GLY:H	1.85	0.41
28:11:242:ARG:HE	28:11:242:ARG:H	1.69	0.41
29:21:117:MET:O	29:21:118:LYS:HB3	2.19	0.41
38:88:97:VAL:HG11	38:88:103:MET:HE3	2.02	0.41
41:B8:67:SER:OG	41:B8:67:SER:O	2.38	0.41
41:B8:136:GLN:H	41:B8:136:GLN:HG2	1.65	0.41
42:C8:92:ARG:HB3	43:D8:11:GLN:CD	2.46	0.41
43:D8:44:LYS:HE2	43:D8:44:LYS:HB3	1.93	0.41
49:J8:82:LEU:HB2	49:J8:83:GLU:H	1.69	0.41
1:1G:164:U:H2'	1:1G:165:C:C6	2.56	0.41
1:1G:310:G:OP1	16:7A:27:LYS:HD3	2.20	0.41
1:1G:482:A:H2'	1:1G:482:A:N3	2.35	0.41
1:1G:661:G:H1	1:1G:744:C:H42	1.69	0.41
1:1G:688:G:H2'	1:1G:689:C:C6	2.55	0.41
1:1G:773:G:H1	1:1G:806:C:H42	1.68	0.41
1:1G:878:G:H5'	8:72:89:PRO:HG2	2.03	0.41
1:1G:1442:G:C6	1:1G:1446:A:C5	3.08	0.41
2:12:121:LEU:HD22	2:12:127:ILE:HG12	2.02	0.41
5:42:118:ILE:HD12	5:42:118:ILE:HA	1.95	0.41
7:62:102:ARG:HG2	7:62:106:GLN:NE2	2.36	0.41
19:AA:53:ASN:HB2	19:AA:77:THR:HG22	2.03	0.41
25:14:78:A:H2'	25:14:79:G:H8	1.85	0.41
25:14:219:G:N3	25:14:234:C:O2'	2.49	0.41
25:14:747:U:O2	25:14:2014:A:H1'	2.21	0.41
25:14:997:G:OP1	42:85:93:LYS:N	2.52	0.41
25:14:1352:U:O2	25:14:1570:A:H2	2.04	0.41
25:14:2298:A:C2	25:14:2299:G:H1'	2.55	0.41
25:14:2320:A:H61	25:14:2333:A:H2'	1.85	0.41
25:14:2694:G:H2'	25:14:2695:C:H6	1.85	0.41
26:1J:6:C:H2'	26:1J:7:G:C8	2.55	0.41
26:1J:11:C:OP2	26:1J:12:C:N4	2.42	0.41
30:39:153:SER:N	30:39:190:GLU:OE2	2.35	0.41
30:39:172:TRP:H	30:39:172:TRP:CD1	2.39	0.41
37:35:8:PRO:HG2	37:35:13:ASN:OD1	2.21	0.41
44:A5:36:LEU:HD13	44:A5:48:ALA:HA	2.02	0.41
45:B5:72:LYS:HG2	45:B5:73:ARG:O	2.21	0.41
52:I5:14:ILE:HA	52:I5:30:GLU:HA	2.01	0.41
1:13:35:G:C6	1:13:36:C:N4	2.89	0.41
1:13:438:G:H3'	1:13:493:G:H1	1.85	0.41
1:13:1077:G:N2	1:13:1079:G:H3'	2.36	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:8E:40:LEU:HD23	9:8E:40:LEU:HA	1.86	0.41
19:AI:64:GLU:H	19:AI:64:GLU:HG2	1.60	0.41
25:1H:48:G:H22	25:1H:177:G:H5'	1.85	0.41
25:1H:372:G:O2'	25:1H:373:U:P	2.79	0.41
25:1H:873:G:H1	25:1H:904:C:N4	2.17	0.41
25:1H:2035:G:H4'	25:1H:2036:C:OP2	2.20	0.41
25:1H:2286:A:P	54:O8:28:ARG:HH21	2.44	0.41
25:1H:2684:U:H3	25:1H:2725:A:H61	1.69	0.41
25:1H:2762:G:H5'	25:1H:2763:G:OP2	2.21	0.41
25:1H:2853:C:H42	25:1H:2864:G:H1	1.67	0.41
27:71:5:LYS:NZ	27:71:8:ARG:HH12	2.18	0.41
27:71:15:ASP:HB3	27:71:18:LYS:CB	2.50	0.41
31:41:173:LEU:O	31:41:178:PHE:HB2	2.21	0.41
38:88:54:MET:HE2	38:88:54:MET:HB2	1.82	0.41
41:B8:78:LEU:HD22	41:B8:78:LEU:HA	1.85	0.41
42:C8:82:GLY:HA3	42:C8:113:ALA:HB1	2.02	0.41
50:K8:8:LYS:O	50:K8:12:GLU:HB2	2.20	0.41
50:K8:13:ALA:O	50:K8:16:LEU:HB3	2.19	0.41
54:O8:27:LYS:NZ	54:O8:27:LYS:HB2	2.35	0.41
56:Q8:6:THR:HG23	56:Q8:63:PRO:HD2	2.03	0.41
1:1G:562:C:H4'	1:1G:563:A:O5'	2.21	0.41
1:1G:663:A:H2'	1:1G:664:G:O4'	2.20	0.41
1:1G:718:G:H5'	11:2A:117:ASN:ND2	2.35	0.41
1:1G:779:C:O5'	62:1G:1816:HOH:O	2.22	0.41
1:1G:980:C:O2	14:5A:19:ARG:HA	2.21	0.41
1:1G:1190:G:OP1	3:22:5:ILE:HG23	2.21	0.41
1:1G:1502:A:H2	1:1G:1505:G:N1	2.19	0.41
3:22:35:GLU:CD	3:22:59:ARG:HH22	2.28	0.41
4:32:3:ARG:H	4:32:3:ARG:HG3	1.64	0.41
4:32:172:PRO:HB2	4:32:187:ARG:HH22	1.85	0.41
8:72:29:SER:HB3	8:72:32:LYS:CG	2.50	0.41
20:BA:9:ASN:O	20:BA:12:ALA:N	2.51	0.41
25:14:568:U:O2'	25:14:570:G:N7	2.45	0.41
25:14:883:G:O2'	25:14:884:C:OP1	2.27	0.41
25:14:1359:A:O2'	25:14:1360:A:OP1	2.38	0.41
25:14:1536:A:H8	25:14:1537:C:H1'	1.86	0.41
25:14:1536:A:H3'	25:14:1537:C:O4'	2.21	0.41
25:14:1983:C:OP1	62:14:3563:HOH:O	2.22	0.41
25:14:2101:G:N2	25:14:2189:U:O2	2.53	0.41
25:14:2119:A:C2	25:14:2172:U:H1'	2.55	0.41
25:14:2478:A:H3'	25:14:2479:G:C8	2.56	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:2602:A:OP2	25:14:2603:G:H5''	2.21	0.41
25:14:2688:U:H6	25:14:2721:A:H62	1.68	0.41
28:19:228:PRO:HD3	28:19:235:GLY:CA	2.51	0.41
33:69:66:GLU:HA	33:69:69:LYS:HB2	2.03	0.41
33:69:121:LYS:HB3	33:69:121:LYS:HE3	1.79	0.41
33:69:131:LYS:HB3	33:69:132:PRO:HA	2.02	0.41
37:35:97:PRO:HD3	37:35:126:VAL:O	2.20	0.41
38:45:19:GLY:O	38:45:98:LYS:HB3	2.21	0.41
41:75:55:ASN:ND2	41:75:58:ASN:HD21	2.18	0.41
43:95:61:VAL:HA	43:95:94:LEU:HD23	2.01	0.41
45:B5:50:LYS:HB3	45:B5:84:ALA:HB2	2.03	0.41
45:B5:55:ASN:HB2	45:B5:80:ILE:HG13	2.02	0.41
46:C5:37:VAL:N	46:C5:67:LEU:O	2.44	0.41
1:13:17:U:H2'	1:13:18:C:C6	2.55	0.41
1:13:244:U:H4'	1:13:245:C:C5'	2.50	0.41
1:13:545:C:OP1	4:3E:61:LYS:NZ	2.44	0.41
1:13:694:A:H5''	11:2I:53:SER:CB	2.50	0.41
1:13:701:C:O2'	1:13:702:A:OP2	2.36	0.41
3:2E:3:ASN:OD1	3:2E:3:ASN:N	2.53	0.41
6:5E:19:LEU:HD11	6:5E:59:TYR:CE2	2.55	0.41
8:7E:100:ILE:HA	8:7E:101:PRO:HD3	1.89	0.41
9:8E:95:LYS:H	9:8E:95:LYS:HD3	1.84	0.41
10:1I:22:LYS:HB3	10:1I:22:LYS:HE3	1.71	0.41
12:3I:55:VAL:O	12:3I:62:GLU:HA	2.21	0.41
13:4I:49:THR:HG22	13:4I:51:ALA:H	1.85	0.41
25:1H:338:G:N2	25:1H:339:U:H1'	2.35	0.41
25:1H:707:G:H1	25:1H:724:U:H3	1.67	0.41
25:1H:849:A:H61	25:1H:929:G:H1'	1.85	0.41
25:1H:977:G:C4	25:1H:978:G:C8	3.09	0.41
25:1H:2592:G:H2'	25:1H:2593:U:O4'	2.20	0.41
28:11:72:LYS:HE3	28:11:72:LYS:HB3	1.77	0.41
28:11:95:LEU:HD13	28:11:97:TYR:CE1	2.54	0.41
34:38:4:LYS:O	34:38:7:VAL:HG12	2.21	0.41
34:38:6:ASN:O	34:38:8:GLU:N	2.54	0.41
35:58:115:ARG:HA	35:58:118:LYS:HD2	2.03	0.41
38:88:72:LYS:HA	38:88:73:PRO:HD3	1.94	0.41
44:E8:24:ILE:O	44:E8:27:LYS:HG3	2.21	0.41
47:H8:91:LEU:HD13	47:H8:91:LEU:HA	1.90	0.41
49:J8:91:LYS:O	49:J8:94:LEU:N	2.45	0.41
50:K8:20:GLU:HA	50:K8:23:LYS:HB3	2.03	0.41
54:O8:13:CYS:O	54:O8:21:TYR:HA	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:552:U:H5'	12:3A:83:ARG:HH21	1.85	0.41
1:1G:627:G:H2'	1:1G:628:G:H8	1.86	0.41
1:1G:1240:U:H1'	7:62:42:ILE:HD11	2.02	0.41
1:1G:1254:C:OP1	10:1A:45:ARG:HA	2.19	0.41
1:1G:1509:C:H2'	1:1G:1510:U:O4'	2.21	0.41
1:1G:1517:G:C3'	1:1G:1518:MA6:H8	2.51	0.41
6:52:62:TRP:CZ2	6:52:64:GLN:HB2	2.54	0.41
13:4A:93:ARG:HA	13:4A:93:ARG:HD3	1.95	0.41
25:14:394:A:H5'	25:14:395:U:OP2	2.21	0.41
25:14:448:U:H3'	25:14:449:A:H8	1.85	0.41
25:14:845:G:H21	25:14:933:A:H61	1.69	0.41
25:14:999:U:H3'	25:14:1154:G:O6	2.20	0.41
25:14:1296:G:OP1	25:14:2709:G:O2'	2.23	0.41
25:14:1816:G:O6	28:19:35:LYS:NZ	2.52	0.41
28:19:264:LYS:HE2	28:19:266:SER:HB3	2.03	0.41
29:29:107:THR:O	29:29:190:GLY:HA2	2.21	0.41
29:29:182:LEU:HD12	29:29:182:LEU:HA	1.83	0.41
30:39:22:ALA:C	30:39:24:LEU:H	2.28	0.41
41:75:107:ASP:O	41:75:111:ARG:NH2	2.51	0.41
43:95:57:VAL:HG13	43:95:99:ILE:HG12	2.02	0.41
44:A5:67:ASP:OD1	44:A5:67:ASP:N	2.42	0.41
47:D5:119:GLU:H	47:D5:119:GLU:HG3	1.56	0.41
1:13:106:C:O2	1:13:379:C:H4'	2.21	0.41
1:13:439:A:OP2	1:13:493:G:N2	2.53	0.41
1:13:1095:U:OP1	1:13:1108:G:N2	2.53	0.41
1:13:1137:C:H1'	1:13:1138:G:N2	2.36	0.41
1:13:1159:U:H1'	1:13:1182:G:H21	1.86	0.41
4:3E:31:CYS:SG	4:3E:33:MET:N	2.94	0.41
6:5E:33:TYR:HB2	6:5E:75:LEU:HD12	2.01	0.41
7:6E:20:ASP:O	7:6E:24:THR:OG1	2.39	0.41
22:1K:35:C:H2'	22:1K:36:A:C8	2.55	0.41
22:3K:69:C:C4	22:3K:70:C:C4	3.08	0.41
24:4K:16:A:H2'	24:4K:17:U:O4'	2.21	0.41
25:1H:218:A:H8	25:1H:218:A:O5'	2.04	0.41
25:1H:273(D):C:H2'	25:1H:273(E):U:C6	2.56	0.41
25:1H:273(F):C:C3'	25:1H:274:G:H5''	2.46	0.41
25:1H:622:G:OP2	37:78:108:LYS:HD3	2.21	0.41
25:1H:1918:A:O2'	25:1H:1920:OMC:N4	2.53	0.41
25:1H:2751:G:C6	32:51:3:ARG:HB2	2.56	0.41
27:71:10:LEU:HD23	27:71:219:GLY:HA2	2.02	0.41
29:21:51:PHE:O	29:21:75:VAL:HB	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:31:195:ASP:OD1	30:31:195:ASP:N	2.54	0.41
31:41:95:ARG:C	31:41:99:MET:HG2	2.45	0.41
34:38:4:LYS:HG2	34:38:7:VAL:HB	2.03	0.41
36:68:25:LEU:HB2	36:68:38:VAL:O	2.21	0.41
39:98:56:LYS:C	39:98:58:GLY:H	2.28	0.41
39:98:100:LEU:HA	39:98:100:LEU:HD13	1.86	0.41
41:B8:12:SER:O	41:B8:15:VAL:HG22	2.20	0.41
41:B8:123:GLN:C	41:B8:125:ARG:H	2.29	0.41
46:G8:54:LYS:NZ	46:G8:54:LYS:HB3	2.35	0.41
47:H8:58:VAL:HA	47:H8:68:PRO:HA	2.02	0.41
53:N8:25:LEU:HD23	53:N8:25:LEU:HA	1.96	0.41
1:1G:468:A:O2'	16:7A:82:GLN:HG2	2.20	0.41
1:1G:482:A:H5'	1:1G:483:C:OP2	2.21	0.41
1:1G:882:C:H41	12:3A:6:GLN:HE22	1.68	0.41
1:1G:929:G:H5'	1:1G:1534:A:OP2	2.20	0.41
1:1G:1035:A:O5'	1:1G:1035:A:H8	2.04	0.41
1:1G:1176:A:H62	1:1G:1177:G:N2	2.18	0.41
5:42:48:ALA:HB1	5:42:49:PRO:HD2	2.03	0.41
7:62:70:LYS:O	7:62:138:LYS:HE3	2.21	0.41
9:82:77:ILE:O	9:82:81:ILE:HG12	2.21	0.41
10:1A:12:ASP:O	10:1A:16:LEU:HB2	2.21	0.41
13:4A:56:LEU:HD22	13:4A:56:LEU:HA	1.88	0.41
17:8A:10:VAL:HG13	17:8A:19:VAL:HB	2.01	0.41
25:14:459:U:OP2	55:L5:39:ARG:NH1	2.54	0.41
25:14:522:G:C6	25:14:523:C:C4	3.09	0.41
25:14:656:G:H2'	25:14:657:U:O4'	2.21	0.41
25:14:883:G:H1	25:14:893:C:N4	2.19	0.41
25:14:932:G:H4'	25:14:933:A:O5'	2.20	0.41
25:14:1348:G:H2'	25:14:1349:A:H5''	2.02	0.41
25:14:1791:A:H3'	25:14:1792:G:H8	1.85	0.41
25:14:2291:U:OP1	25:14:2381:C:H5'	2.21	0.41
25:14:2299:G:N1	25:14:2318:G:H8	2.18	0.41
25:14:2699:C:H2'	25:14:2700:C:O4'	2.21	0.41
26:1J:42:C:O2	31:49:93:THR:HB	2.21	0.41
26:1J:65:C:N4	26:1J:108:C:H2'	2.36	0.41
28:19:176:ARG:HA	28:19:182:LEU:HD22	2.01	0.41
29:29:107:THR:HA	29:29:163:GLU:O	2.21	0.41
29:29:163:GLU:O	29:29:165:VAL:HG23	2.21	0.41
30:39:176:LEU:HD23	30:39:181:LEU:HA	2.02	0.41
30:39:183:VAL:O	30:39:187:VAL:HG23	2.20	0.41
31:49:133:LEU:HD23	31:49:157:ILE:HB	2.02	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:15:19:GLU:CD	35:15:59:LYS:HD2	2.46	0.41
40:65:11:LYS:HD3	40:65:91:PRO:HG3	2.02	0.41
41:75:19:LEU:H	41:75:19:LEU:HG	1.67	0.41
41:75:96:ARG:HG3	41:75:97:ALA:H	1.86	0.41
41:75:120:ARG:HG2	41:75:123:GLN:NE2	2.36	0.41
45:B5:54:VAL:HG13	45:B5:81:VAL:HG12	2.02	0.41
47:D5:106:GLY:HA2	47:D5:109:ALA:HB3	2.03	0.41
47:D5:145:GLU:HG2	47:D5:146:ILE:N	2.35	0.41
56:M5:37:SER:O	56:M5:38:GLY:C	2.63	0.41
1:13:175:C:H2'	1:13:176:C:H6	1.85	0.41
1:13:309:G:H2'	1:13:310:G:C8	2.54	0.41
1:13:548:G:OP1	4:3E:73:ARG:NH2	2.54	0.41
1:13:563:A:H2	12:3I:12:ARG:NH1	2.19	0.41
1:13:719:C:O2'	18:9I:49:LYS:HB3	2.21	0.41
1:13:765:G:H22	1:13:812:C:H2'	1.85	0.41
1:13:954:G:H2'	1:13:955:U:C6	2.56	0.41
1:13:1003:G:H2'	1:13:1004:A:H5'	2.03	0.41
1:13:1296:C:H5''	1:13:1297:C:OP2	2.21	0.41
1:13:1448:C:N3	1:13:1455:G:N2	2.62	0.41
3:2E:29:TYR:O	3:2E:33:LEU:HB3	2.20	0.41
3:2E:76:VAL:HG21	3:2E:103:VAL:HG11	2.01	0.41
3:2E:188:LEU:HD13	3:2E:188:LEU:HA	1.92	0.41
5:4E:6:PHE:HE1	5:4E:36:ASP:HB3	1.84	0.41
6:5E:41:GLU:O	6:5E:43:LEU:HG	2.20	0.41
7:6E:24:THR:HA	7:6E:27:ILE:HG22	2.03	0.41
8:7E:14:ARG:NE	8:7E:83:ILE:O	2.39	0.41
8:7E:32:LYS:O	8:7E:35:ILE:HG13	2.21	0.41
9:8E:8:GLY:HA3	9:8E:76:ALA:O	2.21	0.41
10:1I:44:VAL:HG22	10:1I:66:ARG:HG2	2.02	0.41
12:3I:114:ARG:HB3	12:3I:119:THR:HB	2.03	0.41
13:4I:37:THR:HG22	13:4I:55:ARG:NE	2.36	0.41
13:4I:81:LEU:HD21	13:4I:88:ARG:HH11	1.86	0.41
17:8I:48:GLU:HB2	17:8I:50:LYS:HD3	2.03	0.41
22:1K:8:U:H2'	22:1K:13:C:N4	2.36	0.41
22:1K:64:G:H2'	22:1K:65:G:C8	2.55	0.41
24:4K:10:G:O2'	24:4K:11:U:OP1	2.32	0.41
25:1H:173:G:H2'	25:1H:174:C:C6	2.56	0.41
25:1H:270(I):G:N2	25:1H:270(R):G:H1'	2.36	0.41
25:1H:442:G:H1'	30:3I:48:THR:HG21	2.02	0.41
25:1H:546:C:H5	25:1H:547:A:C2	2.39	0.41
25:1H:644:A:H8	25:1H:644:A:OP2	2.04	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:855:G:C6	25:1H:856:C:N4	2.89	0.41
25:1H:1062:G:H1'	25:1H:1088:A:C5	2.55	0.41
25:1H:1608:A:H1'	25:1H:1610:A:OP2	2.20	0.41
25:1H:1835:G:N3	25:1H:1835:G:H2'	2.35	0.41
25:1H:1889:A:N1	25:1H:2234:G:H1'	2.36	0.41
25:1H:1922:G:H2'	25:1H:1923:U:C6	2.56	0.41
25:1H:2082:A:H2'	25:1H:2083:G:O4'	2.20	0.41
25:1H:2168:G:N3	25:1H:2168:G:H2'	2.35	0.41
25:1H:2252:G:H2'	25:1H:2253:G:O4'	2.21	0.41
25:1H:2319:G:H4'	25:1H:2320:A:OP1	2.20	0.41
25:1H:2402:C:H4'	25:1H:2402:C:OP1	2.21	0.41
25:1H:2512:C:H2'	25:1H:2513:G:O4'	2.21	0.41
25:1H:2523:G:H2'	25:1H:2524:G:H5''	2.02	0.41
25:1H:2728:U:H2'	25:1H:2729:G:C8	2.56	0.41
26:16:24:G:H4'	26:16:25:A:H8	1.86	0.41
34:38:27:VAL:HG22	34:38:111:LEU:H	1.86	0.41
34:38:32:LEU:HB2	34:38:33:PRO:HD3	2.03	0.41
34:38:40:LEU:H	34:38:40:LEU:HD12	1.86	0.41
39:98:73:VAL:O	39:98:76:VAL:HG12	2.21	0.41
41:B8:29:ARG:HH12	41:B8:89:VAL:HG11	1.86	0.41
42:C8:58:ARG:HA	42:C8:61:TRP:CE3	2.56	0.41
47:H8:140:ASP:HB3	47:H8:141:VAL:H	1.65	0.41
54:O8:18:ARG:HB2	54:O8:44:ARG:HH12	1.85	0.41
1:1G:15:G:H2'	1:1G:16:A:C8	2.56	0.41
1:1G:110:C:H2'	1:1G:111:G:O4'	2.20	0.41
1:1G:302:G:O2'	1:1G:556:C:H5''	2.21	0.41
1:1G:656:C:O2	15:6A:28:GLN:NE2	2.51	0.41
1:1G:806:C:H2'	1:1G:807:A:H8	1.86	0.41
1:1G:960:U:O2'	1:1G:961:U:P	2.79	0.41
1:1G:1053:G:O2'	1:1G:1054:C:P	2.78	0.41
1:1G:1116:C:H42	1:1G:1184:G:H1	1.69	0.41
1:1G:1330:U:H4'	13:4A:23:TYR:CD1	2.56	0.41
1:1G:1402:4OC:O5'	1:1G:1402:4OC:H6	2.21	0.41
1:1G:1404:5MC:H2'	1:1G:1405:G:C8	2.56	0.41
1:1G:1411:C:H2'	1:1G:1412:C:C6	2.56	0.41
2:12:29:ALA:O	2:12:32:ILE:HG22	2.21	0.41
2:12:178:ARG:NH1	8:72:68:ARG:HH22	2.19	0.41
3:22:129:ALA:HB3	3:22:132:ARG:HB3	2.03	0.41
4:32:20:TYR:HA	4:32:26:CYS:SG	2.61	0.41
4:32:121:VAL:O	4:32:134:ASP:HA	2.20	0.41
5:42:36:ASP:CG	5:42:38:GLN:HB3	2.46	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:62:27:ILE:HD11	7:62:43:PHE:CE2	2.56	0.41
7:62:113:GLU:H	7:62:113:GLU:HG2	1.69	0.41
8:72:2:LEU:HD23	8:72:2:LEU:HA	1.90	0.41
9:82:9:ARG:HB3	9:82:104:ARG:NH1	2.36	0.41
13:4A:118:ALA:HB1	57:2L:30:G:H5'	2.03	0.41
25:14:68:G:H3'	25:14:69:C:C6	2.56	0.41
25:14:242:G:O2'	25:14:254:G:O6	2.37	0.41
25:14:302:C:H42	25:14:315:G:H1	1.68	0.41
25:14:307:G:N2	25:14:310:A:OP2	2.53	0.41
25:14:428:A:H5'	25:14:429:A:OP2	2.21	0.41
25:14:1063:G:C6	25:14:1076:C:N3	2.89	0.41
25:14:1085:A:N3	25:14:1085:A:H2'	2.35	0.41
25:14:1210:A:H4'	25:14:1211:U:O5'	2.17	0.41
25:14:1363:C:O2'	25:14:1809:A:N3	2.51	0.41
25:14:1491:G:H2'	25:14:1492:G:C8	2.56	0.41
25:14:1814:G:O3'	28:19:54:ARG:NH2	2.54	0.41
25:14:1825:A:H2'	25:14:1826:G:C8	2.56	0.41
25:14:1826:G:OP1	28:19:224:ALA:N	2.52	0.41
25:14:2075:U:H2'	25:14:2238:G:N2	2.36	0.41
25:14:2090:G:C6	25:14:2091:U:C4	3.09	0.41
25:14:2538:C:H2'	25:14:2539:C:C6	2.55	0.41
25:14:2585:U:O2'	25:14:2586:C:P	2.78	0.41
25:14:2611:U:C4	53:J5:3:LYS:HG2	2.55	0.41
25:14:2617:C:C2	25:14:2618:G:C8	3.09	0.41
25:14:2704:C:H2'	25:14:2705:A:O4'	2.21	0.41
25:14:2837:G:N2	39:55:96:ARG:HH12	2.18	0.41
26:1J:21:G:H1	26:1J:62:C:H42	1.67	0.41
26:1J:86:G:H1	26:1J:90:C:H42	1.68	0.41
26:1J:87:G:C2	26:1J:89:G:H5''	2.55	0.41
28:19:71:ASP:CG	28:19:103:ARG:HH22	2.28	0.41
37:35:22:GLY:O	37:35:25:SER:OG	2.35	0.41
37:35:36:LYS:HZ2	37:35:36:LYS:HG3	1.63	0.41
40:65:20:ARG:O	40:65:20:ARG:HG2	2.20	0.41
40:65:34:HIS:CD2	40:65:54:LEU:HG	2.56	0.41
40:65:93:LYS:HD3	40:65:95:HIS:HB2	2.03	0.41
47:D5:156:LYS:C	47:D5:157:LEU:HD13	2.46	0.41
53:J5:42:PRO:HB2	53:J5:43:HIS:ND1	2.36	0.41
1:13:75:C:H2'	1:13:76:G:O4'	2.21	0.41
1:13:834:C:OP1	18:9I:60:ALA:HB2	2.20	0.41
1:13:949:A:N6	1:13:1232:U:H3	2.19	0.41
1:13:962:C:H2'	1:13:963:G:O4'	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1356:G:H2'	1:13:1357:A:C8	2.56	0.41
1:13:1397:C:N4	24:4K:22:A:H3'	2.35	0.41
2:1E:187:LEU:HD22	2:1E:201:ILE:O	2.21	0.41
5:4E:48:ALA:HB3	5:4E:54:ALA:HB2	2.02	0.41
9:8E:59:PHE:HZ	9:8E:88:TYR:CE2	2.39	0.41
16:7I:49:LEU:HD12	16:7I:50:LYS:H	1.86	0.41
25:1H:588:U:H2'	25:1H:589:C:C6	2.56	0.41
25:1H:625:G:O6	37:78:107:LYS:NZ	2.41	0.41
25:1H:1020:A:N1	25:1H:1141:U:H2'	2.36	0.41
25:1H:1048:A:P	25:1H:1110:G:H22	2.44	0.41
25:1H:1324:G:C4	25:1H:1328:G:O6	2.74	0.41
25:1H:2008:C:H2'	25:1H:2009:G:H8	1.85	0.41
25:1H:2124:G:O5'	25:1H:2124:G:H8	2.04	0.41
25:1H:2205:C:H2'	25:1H:2206:C:H6	1.86	0.41
25:1H:2238:G:H2'	25:1H:2238:G:N3	2.36	0.41
25:1H:2293:C:H2'	25:1H:2294:C:O4'	2.20	0.41
25:1H:2382:G:H8	25:1H:2382:G:OP2	2.04	0.41
25:1H:2471:C:H2'	25:1H:2472:G:O4'	2.20	0.41
25:1H:2485:G:OP1	38:88:46:GLN:NE2	2.37	0.41
29:21:105:THR:HG23	29:21:166:THR:OG1	2.20	0.41
31:41:99:MET:O	31:41:103:LEU:HB2	2.21	0.41
33:61:14:ASP:CG	33:61:15:VAL:H	2.29	0.41
35:58:48:MET:SD	35:58:48:MET:N	2.90	0.41
37:78:138:LEU:HD21	37:78:144:GLU:HG2	2.03	0.41
53:N8:30:LEU:HD23	53:N8:41:PRO:HB3	2.02	0.41
54:O8:30:THR:HA	54:O8:31:PRO:HA	1.90	0.41
55:P8:13:ALA:O	55:P8:17:GLY:HA3	2.21	0.41
1:1G:184:G:H2'	1:1G:185:A:C8	2.55	0.41
1:1G:232:G:H1'	1:1G:262:A:N1	2.35	0.41
1:1G:627:G:H2'	1:1G:628:G:C8	2.55	0.41
1:1G:920:U:H2'	1:1G:921:U:H6	1.83	0.41
1:1G:1128:C:C4	1:1G:1139:G:N2	2.89	0.41
1:1G:1512:U:H2'	1:1G:1513:A:H8	1.84	0.41
6:52:1:MET:HE3	6:52:67:MET:HA	2.03	0.41
6:52:20:ALA:O	6:52:24:GLU:N	2.53	0.41
18:9A:31:LEU:CD1	18:9A:65:ILE:HD13	2.50	0.41
21:1B:15:ARG:HB2	21:1B:15:ARG:NH1	2.35	0.41
25:14:84:A:N6	25:14:102:G:O2'	2.53	0.41
25:14:152:G:H2'	25:14:153:C:O4'	2.21	0.41
25:14:1164:G:H2'	25:14:1165:U:O4'	2.21	0.41
25:14:1332:G:H8	25:14:1332:G:H2'	1.66	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:1728:G:H5''	25:14:1728:G:N3	2.35	0.41
25:14:2313:C:O4'	31:49:40:ASN:ND2	2.54	0.41
25:14:2366:A:H2'	25:14:2367:G:O4'	2.21	0.41
25:14:2543:G:H21	25:14:2646:C:H5''	1.85	0.41
25:14:2735:G:H2'	25:14:2736:G:C8	2.56	0.41
31:49:75:LYS:O	31:49:76:SER:HB2	2.21	0.41
32:59:6:ARG:O	32:59:65:HIS:NE2	2.54	0.41
38:45:78:PRO:HG3	38:45:87:LYS:HD3	2.03	0.41
47:D5:110:GLY:H	47:D5:142:SER:HB2	1.86	0.41
53:J5:37:LYS:HA	53:J5:37:LYS:HD2	1.86	0.41
1:13:328:C:H4'	1:13:329:A:C5'	2.50	0.40
1:13:438:G:H4'	4:3E:123:HIS:CG	2.56	0.40
1:13:665:A:N3	1:13:732:C:H2'	2.36	0.40
1:13:848:C:H6	1:13:848:C:H3'	1.86	0.40
1:13:1084:G:C5	1:13:1085:U:C4	3.09	0.40
1:13:1309:G:OP2	13:4I:99:ARG:NH2	2.53	0.40
1:13:1309:G:N1	1:13:1328:C:N3	2.52	0.40
1:13:1346:A:H2	1:13:1347:G:H21	1.70	0.40
8:7E:87:SER:HB2	8:7E:93:VAL:HB	2.03	0.40
9:8E:4:TYR:CE1	9:8E:87:GLN:HB2	2.57	0.40
19:AI:19:VAL:HA	19:AI:22:LEU:HD23	2.02	0.40
25:1H:300:A:H2'	25:1H:334:C:H1'	2.02	0.40
25:1H:362:U:O2'	25:1H:363:G:H5'	2.21	0.40
25:1H:734:A:O2'	25:1H:1635:G:H5'	2.21	0.40
25:1H:909:A:C6	25:1H:912:C:C2	3.09	0.40
25:1H:1690:A:H2'	25:1H:1691:C:O4'	2.21	0.40
25:1H:1773:A:N7	25:1H:1829:A:H1'	2.36	0.40
25:1H:2577:A:H5''	25:1H:2578:G:H5'	2.04	0.40
28:11:215:LEU:H	28:11:215:LEU:HG	1.70	0.40
34:38:39:ALA:HB3	34:38:96:PHE:HE1	1.86	0.40
37:78:95:VAL:HG23	37:78:123:LEU:HD11	2.01	0.40
40:A8:42:ASP:O	40:A8:43:GLU:HB3	2.21	0.40
46:G8:79:CYS:HB2	46:G8:102:CYS:HB3	2.01	0.40
47:H8:6:LYS:HB2	47:H8:6:LYS:HE3	1.88	0.40
1:1G:507:C:H2'	1:1G:508:C:C5	2.56	0.40
1:1G:522:C:OP2	12:3A:66:TYR:OH	2.36	0.40
1:1G:582:U:H2'	1:1G:583:A:C8	2.56	0.40
1:1G:926:G:H3'	1:1G:1505:G:H21	1.85	0.40
1:1G:936:C:H42	1:1G:1379:G:H1	1.68	0.40
1:1G:1095:U:H2'	1:1G:1096:C:O4'	2.21	0.40
1:1G:1290:G:H2'	1:1G:1290:G:N3	2.37	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1347:G:O2'	1:1G:1348:U:P	2.79	0.40
1:1G:1370:G:N7	9:82:109:VAL:HG11	2.35	0.40
5:42:141:GLN:O	5:42:143:ARG:HG2	2.21	0.40
8:72:49:GLU:HB3	8:72:62:TYR:OH	2.20	0.40
57:2L:19:G:H22	57:2L:58:A:H2'	1.86	0.40
57:2L:20:G:N2	57:2L:58:A:N3	2.69	0.40
25:14:300:A:H8	46:C5:84:ARG:HH22	1.69	0.40
25:14:767:U:H2'	25:14:768:G:C8	2.56	0.40
25:14:857:C:H2'	25:14:858:U:C6	2.56	0.40
25:14:1173:G:C8	25:14:1175:U:H5''	2.56	0.40
25:14:1410:G:H2'	25:14:1411:C:C6	2.55	0.40
25:14:1475:G:C2	25:14:1519:G:C2	3.09	0.40
25:14:1496:A:H8	25:14:1577:C:O2'	2.03	0.40
25:14:2030:A:H4'	25:14:2031:A:H8	1.85	0.40
25:14:2113:U:H3'	25:14:2114:A:C4'	2.51	0.40
25:14:2473:U:O2	25:14:2473:U:H2'	2.20	0.40
25:14:2882:A:O4'	39:55:96:ARG:CZ	2.69	0.40
28:19:132:PRO:HG3	28:19:190:TYR:CE1	2.57	0.40
30:39:66:PRO:HD2	30:39:70:THR:HG23	2.04	0.40
32:59:95:ARG:HB3	32:59:95:ARG:NH1	2.36	0.40
32:59:175:LYS:HE2	32:59:175:LYS:HB3	1.90	0.40
36:25:76:ALA:HB3	41:75:75:ILE:HB	2.02	0.40
37:35:15:ARG:O	37:35:16:ARG:C	2.64	0.40
39:55:107:ASP:O	39:55:108:GLY:C	2.65	0.40
45:B5:29:TRP:CE3	45:B5:78:LYS:HB3	2.56	0.40
1:13:41:G:H2'	1:13:42:G:C8	2.56	0.40
1:13:603:U:H2'	1:13:604:G:H8	1.87	0.40
1:13:703:G:O2'	1:13:704:A:P	2.80	0.40
1:13:812:C:N3	62:13:1832:HOH:O	2.37	0.40
1:13:983:A:H2	1:13:984:C:C6	2.39	0.40
1:13:1033:G:H2'	1:13:1034:G:O4'	2.21	0.40
1:13:1348:U:H5	1:13:1373:G:N2	2.19	0.40
1:13:1364:U:O2'	1:13:1365:G:OP1	2.34	0.40
5:4E:81:GLU:HB3	5:4E:90:VAL:HG22	2.03	0.40
25:1H:524:U:H2'	25:1H:525:U:C6	2.56	0.40
25:1H:598:G:H1	25:1H:659:C:N4	2.18	0.40
25:1H:797:C:H2'	25:1H:798:G:O4'	2.21	0.40
25:1H:892:G:H2'	25:1H:892:G:N3	2.37	0.40
25:1H:1805:U:O2	28:11:50:THR:HB	2.22	0.40
25:1H:2138:C:C4	25:1H:2154:G:N2	2.89	0.40
25:1H:2176:A:H4'	27:71:215:THR:HG21	2.04	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:16:74:U:H2'	26:16:75:G:O4'	2.20	0.40
26:16:86:G:H1	26:16:90:C:H42	1.68	0.40
28:11:34:VAL:HG22	28:11:61:LEU:HD12	2.03	0.40
29:21:105:THR:HG21	29:21:164:ARG:CZ	2.51	0.40
30:31:168:ARG:H	30:31:168:ARG:HG3	1.65	0.40
35:58:26:LEU:HG	35:58:30:ILE:HD11	2.03	0.40
36:68:70:LYS:HB3	36:68:70:LYS:HE3	1.85	0.40
38:88:27:VAL:HG21	38:88:134:ARG:HA	2.02	0.40
50:K8:55:ARG:O	50:K8:59:ARG:HB2	2.21	0.40
1:1G:457:C:H2'	1:1G:458:C:H6	1.84	0.40
1:1G:1366:C:H2'	1:1G:1367:C:C6	2.56	0.40
5:42:79:GLU:HG3	5:42:93:PRO:HD2	2.02	0.40
5:42:100:VAL:HG12	5:42:118:ILE:HG22	2.04	0.40
7:62:69:VAL:HG21	7:62:104:LEU:HD21	2.02	0.40
9:82:10:ARG:HB2	9:82:76:ALA:HB2	2.03	0.40
14:5A:23:ARG:NH1	14:5A:30:ALA:HB2	2.36	0.40
17:8A:68:ARG:O	17:8A:68:ARG:HG3	2.20	0.40
57:2L:20:G:H3'	57:2L:21:U:C6	2.57	0.40
25:14:674:G:C1'	30:39:74:ARG:HD3	2.50	0.40
25:14:863:A:O3'	26:1J:100:G:N2	2.42	0.40
25:14:890:A:H3'	25:14:892:G:C8	2.56	0.40
25:14:1114:G:N1	25:14:1115:G:O6	2.55	0.40
25:14:1419:A:H61	25:14:1494:A:N6	2.18	0.40
25:14:1614:A:C2	44:A5:93:ALA:HB2	2.56	0.40
25:14:2313:C:H2'	25:14:2314:C:C5	2.56	0.40
25:14:2692:C:H2'	25:14:2693:A:C8	2.56	0.40
25:14:2844:G:H5'	25:14:2845:G:OP2	2.21	0.40
29:29:134:ILE:HA	29:29:137:HIS:CD2	2.56	0.40
30:39:21:ALA:C	30:39:23:ASP:H	2.29	0.40
30:39:179:GLU:H	30:39:179:GLU:CD	2.29	0.40
32:59:53:GLU:HA	32:59:65:HIS:CE1	2.56	0.40
37:35:14:LYS:O	37:35:15:ARG:C	2.64	0.40
38:45:8:LYS:C	38:45:10:ARG:HH21	2.29	0.40
1:13:614:A:N6	1:13:626:U:H3	2.19	0.40
1:13:728:A:C6	15:6I:54:ARG:HD2	2.56	0.40
1:13:926:G:C6	1:13:1505:G:C6	3.10	0.40
1:13:1073:U:H2'	1:13:1074:G:C8	2.52	0.40
1:13:1195:C:C4	1:13:1197:G:C8	3.10	0.40
1:13:1326:C:H2'	1:13:1327:C:C6	2.56	0.40
1:13:1328:C:O2'	13:4I:29:ARG:NH2	2.49	0.40
1:13:1436:U:H2'	1:13:1437:C:O4'	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1E:100:GLY:N	2:1E:176:GLU:OE2	2.55	0.40
6:5E:50:TYR:CE1	18:9I:77:GLY:HA2	2.57	0.40
7:6E:111:ARG:HD2	7:6E:123:GLU:HB2	2.03	0.40
7:6E:136:LYS:HB3	7:6E:136:LYS:HE3	1.83	0.40
12:3I:102:TYR:C	12:3I:104:ALA:N	2.79	0.40
13:4I:3:ARG:HH22	31:4I:139:LEU:HD13	1.85	0.40
17:8I:52:LYS:HD2	17:8I:52:LYS:H	1.87	0.40
20:BI:30:LYS:HD2	20:BI:30:LYS:HA	1.92	0.40
25:1H:250:G:C6	25:1H:251:A:C6	3.09	0.40
25:1H:665:C:H2'	25:1H:666:G:H8	1.86	0.40
25:1H:994:C:O2	43:D8:10:LYS:HE2	2.22	0.40
25:1H:1087:G:O5'	25:1H:1103:A:N6	2.54	0.40
25:1H:1512:G:H2'	25:1H:1513:C:C6	2.55	0.40
25:1H:1820:U:C2	28:1I:202:LYS:HB3	2.57	0.40
25:1H:1910:G:H2'	25:1H:1911:PSU:H6	1.86	0.40
25:1H:2030:A:H5''	25:1H:2031:A:OP1	2.20	0.40
25:1H:2075:U:OP2	25:1H:2238:G:O2'	2.37	0.40
25:1H:2702:U:H1'	25:1H:2703:C:C5	2.57	0.40
29:21:50:GLY:HA2	29:21:78:LEU:HD13	2.04	0.40
30:31:168:ARG:HG2	30:31:175:THR:HG21	2.03	0.40
38:88:17:LEU:HB3	38:88:39:PRO:HB2	2.02	0.40
46:G8:93:GLY:C	46:G8:94:LYS:HG3	2.47	0.40
46:G8:102:CYS:SG	46:G8:103:GLY:N	2.94	0.40
1:1G:145:G:H1	1:1G:177:C:H42	1.67	0.40
1:1G:245:C:C2	1:1G:284:G:C2	3.10	0.40
1:1G:247:G:C6	1:1G:278:G:C2	3.09	0.40
1:1G:1111:A:H2'	1:1G:1112:C:C6	2.56	0.40
1:1G:1273:G:C4	1:1G:1274:G:C8	3.09	0.40
1:1G:1354:C:H2'	1:1G:1355:G:C8	2.56	0.40
2:12:196:LEU:HD12	2:12:197:VAL:HG23	2.01	0.40
3:22:71:ALA:HB1	3:22:109:PRO:HG3	2.04	0.40
3:22:149:ALA:HA	3:22:201:TYR:O	2.21	0.40
5:42:15:ARG:HG3	5:42:26:PHE:HB3	2.03	0.40
7:62:38:LEU:O	7:62:42:ILE:HG13	2.21	0.40
11:2A:24:SER:N	11:2A:27:ASN:O	2.54	0.40
11:2A:85:ARG:HD3	11:2A:113:PRO:HD3	2.03	0.40
13:4A:14:ARG:HD2	13:4A:42:ALA:HA	2.04	0.40
20:BA:41:ILE:H	20:BA:41:ILE:HG12	1.61	0.40
22:3L:23:G:O2'	22:3L:24:C:P	2.79	0.40
25:14:68:G:H3'	25:14:69:C:H6	1.86	0.40
25:14:270(V):G:H2'	25:14:270(W):G:H8	1.86	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:407:G:H2'	25:14:408:G:H8	1.87	0.40
25:14:959:A:N6	25:14:960:A:N1	2.69	0.40
25:14:1257:C:H4'	30:39:83:PHE:CE1	2.56	0.40
25:14:1387:C:C2	25:14:1388:G:C8	3.09	0.40
25:14:1904:G:H2'	25:14:1905:C:O4'	2.22	0.40
25:14:2093:G:H2'	25:14:2094:G:H8	1.85	0.40
25:14:2584:U:C6	25:14:2585:U:C5	3.09	0.40
25:14:2791:C:H5	25:14:2893:G:H3'	1.87	0.40
25:14:2887:U:H2'	25:14:2888:C:H6	1.86	0.40
26:1J:53:A:H2'	26:1J:54:G:O4'	2.21	0.40
26:1J:86:G:H1	26:1J:90:C:N4	2.20	0.40
26:1J:95:U:C2	26:1J:96:G:N7	2.89	0.40
29:29:55:ASN:HA	29:29:56:PRO:HD3	1.96	0.40
29:29:96:PHE:CD2	29:29:182:LEU:HD21	2.56	0.40
31:49:143:GLU:OE1	31:49:143:GLU:N	2.53	0.40
35:15:16:ILE:HD13	35:15:16:ILE:HA	1.86	0.40
43:95:38:LEU:O	43:95:52:VAL:HB	2.21	0.40
47:D5:111:VAL:C	47:D5:113:ALA:H	2.29	0.40
53:J5:41:PRO:HD2	53:J5:44:THR:CG2	2.51	0.40
1:13:440:A:H8	1:13:440:A:OP2	2.04	0.40
1:13:547:A:N3	1:13:548:G:H1'	2.37	0.40
1:13:701:C:H1'	1:13:703:G:C5	2.56	0.40
1:13:960:U:O2	1:13:960:U:H2'	2.22	0.40
1:13:1195:C:N4	1:13:1197:G:N7	2.68	0.40
1:13:1240:U:OP2	7:6E:116:ALA:N	2.54	0.40
1:13:1267:C:O2	21:1F:20:LYS:HD2	2.22	0.40
4:3E:148:VAL:HG12	4:3E:149:ALA:N	2.37	0.40
10:1I:25:GLU:O	10:1I:29:ARG:HB3	2.21	0.40
15:6I:5:LYS:O	15:6I:8:LYS:HB3	2.21	0.40
17:8I:90:ILE:HG22	17:8I:94:ASN:ND2	2.36	0.40
18:9I:40:LEU:H	18:9I:40:LEU:HG	1.59	0.40
20:BI:57:ARG:NH1	20:BI:102:GLY:HA3	2.36	0.40
25:1H:270(C):C:H42	25:1H:270(W):G:H1	1.69	0.40
25:1H:307:G:H8	25:1H:307:G:O5'	2.05	0.40
25:1H:955:C:OP1	38:88:87:LYS:HE2	2.21	0.40
25:1H:978:G:C2	25:1H:986:C:N3	2.89	0.40
25:1H:1045:A:H4'	25:1H:1046:A:O5'	2.21	0.40
25:1H:1141:U:C5	35:58:64:GLY:HA3	2.56	0.40
25:1H:1728:G:H2'	25:1H:1731:G:O6	2.21	0.40
25:1H:1728:G:H5''	25:1H:1728:G:N3	2.37	0.40
25:1H:2031:A:C6	25:1H:2498:C:H1'	2.57	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:1H:2320:A:H2'	25:1H:2320:A:N3	2.37	0.40
25:1H:2520:C:H2'	25:1H:2521:C:C6	2.57	0.40
25:1H:2790:A:H2'	25:1H:2791:C:H5''	2.02	0.40
25:1H:2847:U:P	41:B8:98:LYS:HZ3	2.44	0.40
26:16:87:G:C2	26:16:89:G:H5''	2.57	0.40
29:21:112:GLY:O	29:21:159:HIS:HA	2.22	0.40
30:31:179:GLU:CD	30:31:179:GLU:H	2.28	0.40
40:A8:59:LYS:HD3	40:A8:60:GLY:N	2.32	0.40
42:C8:5:LYS:H	42:C8:5:LYS:HG3	1.60	0.40
44:E8:18:ARG:NH1	44:E8:76:VAL:O	2.55	0.40
54:O8:7:ILE:O	54:O8:8:LYS:HE3	2.22	0.40
1:1G:109:A:C6	1:1G:327:A:C6	3.10	0.40
1:1G:123:C:OP1	1:1G:312:C:H5'	2.22	0.40
1:1G:1534:A:H2'	1:1G:1535:C:C6	2.57	0.40
2:12:33:TYR:HB2	2:12:42:ILE:C	2.46	0.40
2:12:87:ARG:HE	2:12:233:SER:HB2	1.85	0.40
2:12:132:LYS:HD3	2:12:132:LYS:HA	1.87	0.40
4:32:108:LEU:HD13	4:32:174:LEU:HD22	2.04	0.40
18:9A:37:VAL:HG11	18:9A:78:LEU:HB3	2.03	0.40
19:AA:65:ASN:C	19:AA:66:MET:HG3	2.47	0.40
25:14:336:C:H4'	46:C5:6:HIS:CD2	2.56	0.40
25:14:1540:G:H2'	25:14:1541:U:O4'	2.21	0.40
25:14:1541:U:H2'	25:14:1542:G:O4'	2.22	0.40
25:14:1795:C:C4	25:14:1796:U:C4	3.10	0.40
25:14:2361:A:O5'	56:M5:27:THR:OG1	2.40	0.40
28:19:36:PRO:HA	28:19:61:LEU:HD12	2.04	0.40
28:19:169:GLU:HG2	28:19:174:ILE:HD11	2.04	0.40
31:49:166:ASP:OD1	31:49:166:ASP:N	2.53	0.40
35:15:19:GLU:O	35:15:21:LYS:HE3	2.22	0.40
38:45:42:ILE:HD13	38:45:97:VAL:HB	2.02	0.40
39:55:78:LYS:O	39:55:83:ILE:HG13	2.21	0.40
45:B5:48:LYS:N	45:B5:48:LYS:HD3	2.37	0.40
47:D5:126:VAL:HG12	47:D5:163:LEU:HA	2.03	0.40
47:D5:129:SER:HB2	47:D5:132:ASN:H	1.87	0.40
53:J5:16:ARG:HG3	53:J5:17:ASP:N	2.36	0.40
56:M5:32:LEU:HD23	56:M5:32:LEU:HA	1.74	0.40
1:13:46:G:O2'	1:13:365:U:H1'	2.22	0.40
1:13:537:G:H2'	1:13:538:G:C8	2.56	0.40
1:13:652:U:C4	1:13:752:G:N3	2.90	0.40
1:13:763:G:H2'	1:13:764:C:H6	1.86	0.40
1:13:1013:G:N2	1:13:1016:A:OP2	2.55	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1187:G:O5'	9:8E:113:LYS:HE2	2.21	0.40
1:13:1369:C:OP1	14:5I:61:TRP:NE1	2.55	0.40
1:13:1479:C:H2'	1:13:1480:G:C8	2.56	0.40
3:2E:105:GLU:HG2	3:2E:106:VAL:N	2.36	0.40
12:3I:32:GLY:HA2	12:3I:57:LEU:HA	2.03	0.40
15:6I:3:ILE:HG21	15:6I:34:LEU:HG	2.03	0.40
16:7I:23:ASP:O	16:7I:26:ARG:HB2	2.22	0.40
22:1K:66:C:H2'	22:1K:67:C:O4'	2.21	0.40
23:2K:41:C:H2'	23:2K:42:C:H6	1.86	0.40
24:4K:10:G:HO2'	24:4K:11:U:P	2.44	0.40
25:1H:556:G:H8	25:1H:556:G:O5'	2.04	0.40
25:1H:733:G:C8	25:1H:761:A:N6	2.90	0.40
25:1H:1698:A:H4'	25:1H:1699:G:OP1	2.21	0.40
25:1H:2028:U:H2'	25:1H:2029:G:O4'	2.21	0.40
25:1H:2302:G:H1	25:1H:2314:C:H42	1.69	0.40
25:1H:2472:G:N2	25:1H:2477:C:H5'	2.35	0.40
26:16:41:U:C4	31:41:70:VAL:HG23	2.56	0.40
30:31:45:ARG:HH11	30:31:45:ARG:HG2	1.87	0.40
33:61:1:MET:HG3	33:61:23:PRO:HA	2.03	0.40
36:68:7:TYR:OH	36:68:44:LYS:HG3	2.21	0.40
40:A8:109:GLY:C	40:A8:110:LEU:HD13	2.47	0.40
42:C8:90:VAL:HG12	42:C8:91:ASP:H	1.85	0.40
44:E8:64:MET:HE2	44:E8:69:LEU:HD23	2.04	0.40
45:F8:12:VAL:HG22	45:F8:29:TRP:CE2	2.57	0.40
46:G8:11:ASP:O	46:G8:26:LYS:HG3	2.22	0.40
49:J8:13:ILE:HD11	49:J8:42:GLN:OE1	2.21	0.40
56:Q8:50:LEU:HD22	56:Q8:50:LEU:HA	1.85	0.40
1:1G:181:G:H1'	1:1G:182:U:H5	1.86	0.40
1:1G:376:G:OP2	16:7A:67:THR:HG21	2.21	0.40
1:1G:377:G:O6	62:1G:1814:HOH:O	2.21	0.40
1:1G:533:A:H4'	1:1G:534:U:OP1	2.20	0.40
1:1G:668:G:H1	1:1G:738:C:H42	1.69	0.40
1:1G:709:G:H2'	1:1G:710:G:C8	2.55	0.40
1:1G:779:C:H2'	1:1G:780:A:O4'	2.22	0.40
1:1G:932:C:H5''	7:62:2:ALA:N	2.36	0.40
1:1G:1118:C:H2'	1:1G:1119:C:C6	2.56	0.40
1:1G:1363:A:H4'	1:1G:1364:U:H5''	2.04	0.40
1:1G:1367:C:H5'	10:1A:60:ARG:NH2	2.36	0.40
4:32:64:LEU:HB2	4:32:198:VAL:HG11	2.03	0.40
9:82:25:LYS:N	9:82:25:LYS:HD3	2.36	0.40
15:6A:17:ARG:HH12	15:6A:77:ARG:HH11	1.70	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:14:96:G:H4'	50:G5:48:HIS:CD2	2.56	0.40
25:14:861:A:H2'	25:14:862:G:O4'	2.22	0.40
25:14:998:C:N4	62:14:3710:HOH:O	2.53	0.40
25:14:1116:C:H2'	25:14:1117:G:C8	2.57	0.40
25:14:1607:C:H4'	25:14:1608:A:O5'	2.20	0.40
25:14:2889:C:H2'	25:14:2891:G:O4'	2.21	0.40
28:19:27:THR:HG22	28:19:28:GLU:HG2	2.03	0.40
29:29:6:GLY:HA2	29:29:28:ALA:HA	2.04	0.40
30:39:28:ILE:H	30:39:28:ILE:HG12	1.59	0.40
37:35:57:THR:HG21	37:35:60:MET:H	1.87	0.40
42:85:15:LYS:HE2	42:85:15:LYS:HB2	1.87	0.40
54:K5:9:LEU:O	54:K5:10:LEU:HD13	2.22	0.40

There are no symmetry-related clashes.

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	12	234/256 (91%)	193 (82%)	36 (15%)	5 (2%)	5	31
2	1E	235/256 (92%)	200 (85%)	35 (15%)	0	100	100
3	22	204/239 (85%)	161 (79%)	41 (20%)	2 (1%)	12	44
3	2E	203/239 (85%)	172 (85%)	31 (15%)	0	100	100
4	32	206/209 (99%)	188 (91%)	18 (9%)	0	100	100
4	3E	206/209 (99%)	190 (92%)	16 (8%)	0	100	100
5	42	152/162 (94%)	134 (88%)	18 (12%)	0	100	100
5	4E	149/162 (92%)	140 (94%)	9 (6%)	0	100	100
6	52	99/101 (98%)	94 (95%)	5 (5%)	0	100	100
6	5E	99/101 (98%)	91 (92%)	8 (8%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
7	62	145/156 (93%)	122 (84%)	21 (14%)	2 (1%)	9	38
7	6E	153/156 (98%)	140 (92%)	13 (8%)	0	100	100
8	72	136/138 (99%)	125 (92%)	11 (8%)	0	100	100
8	7E	136/138 (99%)	123 (90%)	13 (10%)	0	100	100
9	82	125/128 (98%)	90 (72%)	29 (23%)	6 (5%)	2	16
9	8E	122/128 (95%)	107 (88%)	15 (12%)	0	100	100
10	1A	97/105 (92%)	79 (81%)	18 (19%)	0	100	100
10	1I	97/105 (92%)	84 (87%)	13 (13%)	0	100	100
11	2A	117/129 (91%)	100 (86%)	16 (14%)	1 (1%)	14	47
11	2I	115/129 (89%)	108 (94%)	7 (6%)	0	100	100
12	3A	122/132 (92%)	101 (83%)	18 (15%)	3 (2%)	4	28
12	3I	122/132 (92%)	102 (84%)	19 (16%)	1 (1%)	16	49
13	4A	116/126 (92%)	79 (68%)	31 (27%)	6 (5%)	1	14
13	4I	117/126 (93%)	99 (85%)	16 (14%)	2 (2%)	7	35
14	5A	58/61 (95%)	49 (84%)	8 (14%)	1 (2%)	7	35
14	5I	58/61 (95%)	48 (83%)	9 (16%)	1 (2%)	7	35
15	6A	86/89 (97%)	81 (94%)	5 (6%)	0	100	100
15	6I	86/89 (97%)	77 (90%)	9 (10%)	0	100	100
16	7A	82/88 (93%)	76 (93%)	6 (7%)	0	100	100
16	7I	81/88 (92%)	73 (90%)	8 (10%)	0	100	100
17	8A	97/105 (92%)	91 (94%)	6 (6%)	0	100	100
17	8I	98/105 (93%)	92 (94%)	6 (6%)	0	100	100
18	9A	69/88 (78%)	62 (90%)	7 (10%)	0	100	100
18	9I	68/88 (77%)	61 (90%)	7 (10%)	0	100	100
19	AA	81/93 (87%)	56 (69%)	25 (31%)	0	100	100
19	AI	84/93 (90%)	68 (81%)	15 (18%)	1 (1%)	10	41
20	BA	101/106 (95%)	89 (88%)	12 (12%)	0	100	100
20	BI	99/106 (93%)	85 (86%)	13 (13%)	1 (1%)	12	44
21	1B	23/27 (85%)	19 (83%)	4 (17%)	0	100	100
21	1F	23/27 (85%)	19 (83%)	4 (17%)	0	100	100
27	71	131/229 (57%)	118 (90%)	13 (10%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
28	11	272/276 (99%)	247 (91%)	24 (9%)	1 (0%)	30	62
28	19	270/276 (98%)	244 (90%)	26 (10%)	0	100	100
29	21	202/206 (98%)	177 (88%)	23 (11%)	2 (1%)	12	44
29	29	202/206 (98%)	183 (91%)	18 (9%)	1 (0%)	24	57
30	31	200/210 (95%)	174 (87%)	25 (12%)	1 (0%)	24	57
30	39	204/210 (97%)	166 (81%)	34 (17%)	4 (2%)	6	32
31	41	179/182 (98%)	144 (80%)	33 (18%)	2 (1%)	11	43
31	49	179/182 (98%)	141 (79%)	35 (20%)	3 (2%)	7	35
32	51	172/180 (96%)	139 (81%)	31 (18%)	2 (1%)	10	41
32	59	172/180 (96%)	131 (76%)	39 (23%)	2 (1%)	10	41
33	61	144/148 (97%)	112 (78%)	30 (21%)	2 (1%)	9	38
33	69	144/148 (97%)	113 (78%)	30 (21%)	1 (1%)	18	51
34	38	135/173 (78%)	75 (56%)	57 (42%)	3 (2%)	5	30
35	15	136/140 (97%)	121 (89%)	15 (11%)	0	100	100
35	58	136/140 (97%)	110 (81%)	23 (17%)	3 (2%)	5	30
36	25	120/122 (98%)	105 (88%)	15 (12%)	0	100	100
36	68	120/122 (98%)	110 (92%)	10 (8%)	0	100	100
37	35	145/150 (97%)	109 (75%)	35 (24%)	1 (1%)	18	51
37	78	146/150 (97%)	110 (75%)	33 (23%)	3 (2%)	5	31
38	45	138/141 (98%)	111 (80%)	26 (19%)	1 (1%)	18	51
38	88	139/141 (99%)	105 (76%)	32 (23%)	2 (1%)	9	38
39	55	116/118 (98%)	107 (92%)	8 (7%)	1 (1%)	14	47
39	98	116/118 (98%)	105 (90%)	9 (8%)	2 (2%)	7	35
40	65	109/112 (97%)	88 (81%)	21 (19%)	0	100	100
40	A8	109/112 (97%)	84 (77%)	24 (22%)	1 (1%)	14	47
41	75	135/146 (92%)	107 (79%)	27 (20%)	1 (1%)	18	51
41	B8	135/146 (92%)	113 (84%)	22 (16%)	0	100	100
42	85	115/118 (98%)	99 (86%)	15 (13%)	1 (1%)	14	47
42	C8	115/118 (98%)	106 (92%)	8 (7%)	1 (1%)	14	47
43	95	99/101 (98%)	78 (79%)	20 (20%)	1 (1%)	12	44
43	D8	98/101 (97%)	83 (85%)	15 (15%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
44	A5	110/113 (97%)	104 (94%)	6 (6%)	0	100	100
44	E8	111/113 (98%)	99 (89%)	12 (11%)	0	100	100
45	B5	91/96 (95%)	77 (85%)	14 (15%)	0	100	100
45	F8	93/96 (97%)	85 (91%)	8 (9%)	0	100	100
46	C5	107/110 (97%)	82 (77%)	22 (21%)	3 (3%)	4	26
46	G8	105/110 (96%)	79 (75%)	24 (23%)	2 (2%)	6	33
47	D5	174/206 (84%)	114 (66%)	55 (32%)	5 (3%)	3	26
47	H8	172/206 (84%)	123 (72%)	45 (26%)	4 (2%)	5	30
48	E5	77/85 (91%)	65 (84%)	12 (16%)	0	100	100
48	I8	75/85 (88%)	65 (87%)	10 (13%)	0	100	100
49	F5	90/96 (94%)	75 (83%)	15 (17%)	0	100	100
49	J8	94/96 (98%)	75 (80%)	16 (17%)	3 (3%)	3	24
50	G5	67/72 (93%)	58 (87%)	9 (13%)	0	100	100
50	K8	69/72 (96%)	56 (81%)	11 (16%)	2 (3%)	3	26
51	H5	57/60 (95%)	53 (93%)	4 (7%)	0	100	100
51	L8	57/60 (95%)	52 (91%)	5 (9%)	0	100	100
52	I5	69/71 (97%)	40 (58%)	28 (41%)	1 (1%)	9	38
52	M8	64/71 (90%)	32 (50%)	30 (47%)	2 (3%)	3	25
53	J5	54/60 (90%)	42 (78%)	10 (18%)	2 (4%)	2	21
53	N8	54/60 (90%)	45 (83%)	7 (13%)	2 (4%)	2	21
54	K5	46/54 (85%)	21 (46%)	22 (48%)	3 (6%)	1	11
54	O8	47/54 (87%)	27 (57%)	17 (36%)	3 (6%)	1	11
55	L5	46/49 (94%)	46 (100%)	0	0	100	100
55	P8	46/49 (94%)	45 (98%)	1 (2%)	0	100	100
56	M5	62/65 (95%)	53 (86%)	7 (11%)	2 (3%)	3	24
56	Q8	62/65 (95%)	58 (94%)	4 (6%)	0	100	100
All	All	11633/12452 (93%)	9754 (84%)	1771 (15%)	108 (1%)	14	47

All (108) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
14	5I	16	PHE
35	58	96	GLU

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Mol	Chain	Res	Type
39	98	3	HIS
47	H8	53	ILE
49	J8	30	VAL
49	J8	82	LEU
50	K8	48	HIS
9	82	30	GLY
9	82	35	GLU
13	4A	68	GLY
13	4A	106	ASN
30	39	22	ALA
31	49	96	ARG
31	49	97	ASP
47	D5	53	ILE
47	D5	153	SER
53	J5	41	PRO
54	K5	31	PRO
33	61	15	VAL
34	38	30	GLN
35	58	22	THR
37	78	57	THR
38	88	80	GLU
39	98	58	GLY
42	C8	92	ARG
46	G8	54	LYS
53	N8	41	PRO
7	62	21	VAL
9	82	29	ASN
11	2A	15	ALA
13	4A	107	ALA
29	29	71	GLY
30	39	90	PHE
37	35	57	THR
42	85	97	ASP
46	C5	54	LYS
47	D5	112	ARG
47	D5	140	ASP
47	D5	152	ALA
53	J5	42	PRO
13	4I	14	ARG
28	11	273	ARG
29	21	71	GLY
29	21	145	LYS

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Mol	Chain	Res	Type
31	41	14	GLU
32	51	168	PRO
34	38	22	GLY
35	58	95	PRO
37	78	106	LEU
40	A8	62	LYS
47	H8	155	LEU
50	K8	47	ASN
52	M8	24	THR
52	M8	49	PHE
53	N8	42	PRO
54	O8	15	GLU
54	O8	21	TYR
2	12	23	ARG
9	82	19	LEU
12	3A	10	LYS
13	4A	67	GLU
39	55	107	ASP
54	K5	24	GLU
31	41	15	VAL
33	61	12	LEU
37	78	107	LYS
38	88	104	PHE
2	12	22	LYS
2	12	33	TYR
3	22	16	ARG
9	82	34	ASN
12	3A	16	ARG
13	4A	44	ARG
33	69	78	THR
41	75	106	SER
46	C5	53	PRO
12	3I	102	TYR
19	AI	81	ARG
20	BI	74	LYS
30	31	130	ALA
46	G8	53	PRO
49	J8	83	GLU
30	39	25	PRO
46	C5	56	PRO
32	51	12	PRO
34	38	44	LEU

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Mol	Chain	Res	Type
2	12	21	ARG
2	12	30	ARG
3	22	160	ALA
7	62	17	VAL
12	3A	15	VAL
14	5A	15	LYS
32	59	128	PRO
56	M5	37	SER
47	H8	110	GLY
13	4A	100	GLY
52	I5	6	HIS
30	39	89	VAL
32	59	168	PRO
38	45	78	PRO
54	O8	7	ILE
9	82	28	VAL
43	95	50	PRO
54	K5	41	PRO
47	H8	141	VAL
13	4I	15	VAL
31	49	44	GLY
56	M5	38	GLY

5.3.2 Protein sidechains ⓘ

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
2	12	204/220 (93%)	148 (72%)	56 (28%)	0	3
2	1E	205/220 (93%)	162 (79%)	43 (21%)	1	6
3	22	160/188 (85%)	115 (72%)	45 (28%)	0	3
3	2E	159/188 (85%)	133 (84%)	26 (16%)	2	14
4	32	180/181 (99%)	156 (87%)	24 (13%)	4	20
4	3E	180/181 (99%)	148 (82%)	32 (18%)	2	10
5	42	119/123 (97%)	88 (74%)	31 (26%)	0	3

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
5	4E	116/123 (94%)	92 (79%)	24 (21%)	1	6
6	52	90/90 (100%)	83 (92%)	7 (8%)	11	36
6	5E	90/90 (100%)	73 (81%)	17 (19%)	1	9
7	62	123/127 (97%)	102 (83%)	21 (17%)	2	12
7	6E	126/127 (99%)	102 (81%)	24 (19%)	1	8
8	72	119/119 (100%)	99 (83%)	20 (17%)	2	13
8	7E	119/119 (100%)	93 (78%)	26 (22%)	1	6
9	82	97/99 (98%)	76 (78%)	21 (22%)	1	6
9	8E	95/99 (96%)	79 (83%)	16 (17%)	2	13
10	1A	89/92 (97%)	70 (79%)	19 (21%)	1	6
10	1I	89/92 (97%)	73 (82%)	16 (18%)	2	10
11	2A	91/99 (92%)	73 (80%)	18 (20%)	1	7
11	2I	88/99 (89%)	70 (80%)	18 (20%)	1	7
12	3A	103/108 (95%)	82 (80%)	21 (20%)	1	7
12	3I	103/108 (95%)	86 (84%)	17 (16%)	2	13
13	4A	94/101 (93%)	66 (70%)	28 (30%)	0	2
13	4I	95/101 (94%)	67 (70%)	28 (30%)	0	2
14	5A	48/50 (96%)	46 (96%)	2 (4%)	26	52
14	5I	49/50 (98%)	41 (84%)	8 (16%)	2	14
15	6A	79/80 (99%)	66 (84%)	13 (16%)	2	13
15	6I	79/80 (99%)	66 (84%)	13 (16%)	2	13
16	7A	72/74 (97%)	61 (85%)	11 (15%)	3	16
16	7I	72/74 (97%)	59 (82%)	13 (18%)	2	10
17	8A	93/97 (96%)	80 (86%)	13 (14%)	3	19
17	8I	95/97 (98%)	83 (87%)	12 (13%)	4	21
18	9A	62/77 (80%)	48 (77%)	14 (23%)	1	5
18	9I	61/77 (79%)	49 (80%)	12 (20%)	1	8
19	AA	72/80 (90%)	56 (78%)	16 (22%)	1	5
19	AI	74/80 (92%)	55 (74%)	19 (26%)	0	3
20	BA	75/82 (92%)	63 (84%)	12 (16%)	2	14
20	BI	75/82 (92%)	62 (83%)	13 (17%)	2	11

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
21	1B	20/22 (91%)	14 (70%)	6 (30%)	0	2
21	1F	20/22 (91%)	19 (95%)	1 (5%)	22	48
27	71	111/181 (61%)	95 (86%)	16 (14%)	3	18
28	11	214/218 (98%)	166 (78%)	48 (22%)	1	5
28	19	213/218 (98%)	181 (85%)	32 (15%)	3	17
29	21	164/166 (99%)	134 (82%)	30 (18%)	2	10
29	29	164/166 (99%)	134 (82%)	30 (18%)	2	10
30	31	161/166 (97%)	135 (84%)	26 (16%)	2	14
30	39	165/166 (99%)	135 (82%)	30 (18%)	2	10
31	41	155/156 (99%)	130 (84%)	25 (16%)	2	14
31	49	155/156 (99%)	126 (81%)	29 (19%)	1	9
32	51	145/148 (98%)	114 (79%)	31 (21%)	1	6
32	59	145/148 (98%)	114 (79%)	31 (21%)	1	6
33	61	122/124 (98%)	80 (66%)	42 (34%)	0	2
33	69	122/124 (98%)	91 (75%)	31 (25%)	0	3
34	38	112/135 (83%)	77 (69%)	35 (31%)	0	2
35	15	117/119 (98%)	98 (84%)	19 (16%)	2	14
35	58	117/119 (98%)	88 (75%)	29 (25%)	1	4
36	25	100/100 (100%)	80 (80%)	20 (20%)	1	7
36	68	100/100 (100%)	84 (84%)	16 (16%)	2	14
37	35	114/116 (98%)	82 (72%)	32 (28%)	0	3
37	78	115/116 (99%)	88 (76%)	27 (24%)	1	4
38	45	110/111 (99%)	89 (81%)	21 (19%)	1	8
38	88	111/111 (100%)	88 (79%)	23 (21%)	1	6
39	55	101/101 (100%)	83 (82%)	18 (18%)	2	10
39	98	101/101 (100%)	78 (77%)	23 (23%)	1	5
40	65	87/88 (99%)	73 (84%)	14 (16%)	2	14
40	A8	87/88 (99%)	61 (70%)	26 (30%)	0	2
41	75	120/127 (94%)	90 (75%)	30 (25%)	0	4
41	B8	120/127 (94%)	75 (62%)	45 (38%)	0	1
42	85	93/94 (99%)	77 (83%)	16 (17%)	2	12

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
42	C8	93/94 (99%)	73 (78%)	20 (22%)	1	6
43	95	82/82 (100%)	59 (72%)	23 (28%)	0	3
43	D8	82/82 (100%)	54 (66%)	28 (34%)	0	2
44	A5	91/92 (99%)	71 (78%)	20 (22%)	1	5
44	E8	92/92 (100%)	69 (75%)	23 (25%)	0	4
45	B5	74/78 (95%)	61 (82%)	13 (18%)	2	11
45	F8	76/78 (97%)	60 (79%)	16 (21%)	1	6
46	C5	88/91 (97%)	58 (66%)	30 (34%)	0	2
46	G8	85/91 (93%)	63 (74%)	22 (26%)	0	3
47	D5	155/179 (87%)	112 (72%)	43 (28%)	0	3
47	H8	153/179 (86%)	116 (76%)	37 (24%)	1	4
48	E5	63/67 (94%)	51 (81%)	12 (19%)	1	8
48	I8	62/67 (92%)	54 (87%)	8 (13%)	4	21
49	F5	77/81 (95%)	56 (73%)	21 (27%)	0	3
49	J8	81/81 (100%)	71 (88%)	10 (12%)	4	22
50	G5	64/67 (96%)	48 (75%)	16 (25%)	0	4
50	K8	64/67 (96%)	42 (66%)	22 (34%)	0	2
51	H5	51/52 (98%)	45 (88%)	6 (12%)	5	23
51	L8	51/52 (98%)	39 (76%)	12 (24%)	1	4
52	I5	63/63 (100%)	43 (68%)	20 (32%)	0	2
52	M8	59/63 (94%)	43 (73%)	16 (27%)	0	3
53	J5	48/52 (92%)	38 (79%)	10 (21%)	1	6
53	N8	48/52 (92%)	37 (77%)	11 (23%)	1	5
54	K5	47/52 (90%)	34 (72%)	13 (28%)	0	3
54	O8	48/52 (92%)	28 (58%)	20 (42%)	0	0
55	L5	41/42 (98%)	33 (80%)	8 (20%)	1	8
55	P8	41/42 (98%)	36 (88%)	5 (12%)	5	22
56	M5	52/55 (94%)	46 (88%)	6 (12%)	5	24
56	Q8	52/55 (94%)	46 (88%)	6 (12%)	5	24
All	All	9804/10308 (95%)	7736 (79%)	2068 (21%)	1	6

All (2068) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
2	1E	4	GLU
2	1E	8	LYS
2	1E	9	GLU
2	1E	15	VAL
2	1E	23	ARG
2	1E	28	PHE
2	1E	32	ILE
2	1E	36	ARG
2	1E	47	THR
2	1E	49	GLU
2	1E	64	ARG
2	1E	69	LEU
2	1E	71	VAL
2	1E	74	LYS
2	1E	79	ASP
2	1E	82	ARG
2	1E	96	ARG
2	1E	107	THR
2	1E	136	VAL
2	1E	145	LEU
2	1E	149	LEU
2	1E	155	LEU
2	1E	158	LEU
2	1E	160	ASP
2	1E	162	ILE
2	1E	164	VAL
2	1E	168	THR
2	1E	172	ILE
2	1E	174	VAL
2	1E	190	THR
2	1E	196	LEU
2	1E	197	VAL
2	1E	200	ILE
2	1E	204	ASN
2	1E	208	ILE
2	1E	215	LEU
2	1E	219	VAL
2	1E	221	LEU
2	1E	222	ILE
2	1E	223	ILE
2	1E	229	VAL
2	1E	230	VAL
2	1E	239	VAL

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Mol	Chain	Res	Type
3	2E	3	ASN
3	2E	5	ILE
3	2E	6	HIS
3	2E	12	LEU
3	2E	21	ARG
3	2E	31	HIS
3	2E	34	LEU
3	2E	43	LEU
3	2E	45	LYS
3	2E	47	LEU
3	2E	48	TYR
3	2E	75	VAL
3	2E	89	GLU
3	2E	94	LEU
3	2E	95	THR
3	2E	102	ASN
3	2E	104	GLN
3	2E	107	GLN
3	2E	127	ARG
3	2E	138	VAL
3	2E	140	ARG
3	2E	164	ARG
3	2E	167	TRP
3	2E	190	ARG
3	2E	195	VAL
3	2E	198	VAL
4	3E	10	ARG
4	3E	14	ARG
4	3E	15	GLU
4	3E	24	GLU
4	3E	30	LYS
4	3E	31	CYS
4	3E	33	MET
4	3E	38	TYR
4	3E	57	ARG
4	3E	66	ARG
4	3E	83	SER
4	3E	86	LYS
4	3E	94	LEU
4	3E	101	LEU
4	3E	114	ARG
4	3E	120	LEU

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Mol	Chain	Res	Type
4	3E	121	VAL
4	3E	122	ARG
4	3E	132	ARG
4	3E	134	ASP
4	3E	137	SER
4	3E	150	GLU
4	3E	155	LEU
4	3E	159	ARG
4	3E	165	MET
4	3E	168	ARG
4	3E	176	LEU
4	3E	184	LYS
4	3E	193	ASP
4	3E	194	LEU
4	3E	196	LEU
4	3E	209	ARG
5	4E	6	PHE
5	4E	8	GLU
5	4E	10	MET
5	4E	13	ILE
5	4E	16	THR
5	4E	31	LEU
5	4E	33	VAL
5	4E	41	VAL
5	4E	47	LYS
5	4E	51	VAL
5	4E	55	VAL
5	4E	64	ARG
5	4E	71	LEU
5	4E	73	ASN
5	4E	79	GLU
5	4E	90	VAL
5	4E	101	ILE
5	4E	112	LEU
5	4E	116	THR
5	4E	120	THR
5	4E	133	TYR
5	4E	142	LEU
5	4E	147	ASP
5	4E	153	LYS
6	5E	1	MET
6	5E	7	ASN

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Mol	Chain	Res	Type
6	5E	11	ASN
6	5E	14	LEU
6	5E	21	LEU
6	5E	25	ILE
6	5E	27	GLN
6	5E	30	LEU
6	5E	36	ARG
6	5E	69	GLU
6	5E	70	ASP
6	5E	74	ASP
6	5E	77	ARG
6	5E	81	ILE
6	5E	89	MET
6	5E	92	LYS
6	5E	98	LEU
7	6E	8	GLU
7	6E	24	THR
7	6E	30	ILE
7	6E	38	LEU
7	6E	52	GLU
7	6E	54	THR
7	6E	56	GLN
7	6E	73	MET
7	6E	74	GLU
7	6E	75	VAL
7	6E	76	ARG
7	6E	78	ARG
7	6E	79	ARG
7	6E	80	VAL
7	6E	84	ASN
7	6E	89	MET
7	6E	90	GLU
7	6E	92	SER
7	6E	97	GLN
7	6E	98	SER
7	6E	113	GLU
7	6E	120	ILE
7	6E	124	LEU
7	6E	135	VAL
8	7E	1	MET
8	7E	10	LEU
8	7E	12	ARG

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Mol	Chain	Res	Type
8	7E	19	VAL
8	7E	22	GLU
8	7E	24	THR
8	7E	26	VAL
8	7E	32	LYS
8	7E	35	ILE
8	7E	38	ILE
8	7E	39	LEU
8	7E	41	ARG
8	7E	50	ARG
8	7E	51	VAL
8	7E	53	VAL
8	7E	56	LYS
8	7E	63	LEU
8	7E	82	HIS
8	7E	83	ILE
8	7E	95	VAL
8	7E	98	LYS
8	7E	102	ARG
8	7E	109	ILE
8	7E	112	LEU
8	7E	129	VAL
8	7E	133	LEU
9	8E	14	VAL
9	8E	25	LYS
9	8E	35	GLU
9	8E	44	VAL
9	8E	47	LEU
9	8E	48	GLU
9	8E	56	LEU
9	8E	64	THR
9	8E	71	SER
9	8E	79	LEU
9	8E	86	VAL
9	8E	95	LYS
9	8E	99	LEU
9	8E	104	ARG
9	8E	117	HIS
9	8E	118	LYS
10	1I	6	ILE
10	1I	13	HIS
10	1I	16	LEU

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Mol	Chain	Res	Type
10	1I	28	ARG
10	1I	38	ILE
10	1I	49	VAL
10	1I	62	HIS
10	1I	68	HIS
10	1I	72	VAL
10	1I	73	ASP
10	1I	74	ILE
10	1I	75	ILE
10	1I	76	ASN
10	1I	84	GLN
10	1I	92	THR
10	1I	96	ILE
11	2I	24	SER
11	2I	29	ILE
11	2I	31	THR
11	2I	32	ILE
11	2I	36	ASP
11	2I	41	THR
11	2I	47	VAL
11	2I	55	LYS
11	2I	57	THR
11	2I	63	LEU
11	2I	87	THR
11	2I	92	GLU
11	2I	103	LEU
11	2I	105	VAL
11	2I	109	VAL
11	2I	114	VAL
11	2I	124	LYS
11	2I	126	ARG
12	3I	4	ILE
12	3I	9	ARG
12	3I	14	LYS
12	3I	17	LYS
12	3I	20	LYS
12	3I	24	LEU
12	3I	30	ARG
12	3I	33	VAL
12	3I	51	LYS
12	3I	52	VAL
12	3I	64	THR

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Mol	Chain	Res	Type
12	3I	67	ILE
12	3I	80	VAL
12	3I	101	VAL
12	3I	114	ARG
12	3I	123	LYS
12	3I	124	GLU
13	4I	4	ILE
13	4I	9	ILE
13	4I	13	LYS
13	4I	17	VAL
13	4I	19	LEU
13	4I	32	GLU
13	4I	36	LYS
13	4I	37	THR
13	4I	45	VAL
13	4I	48	LEU
13	4I	56	LEU
13	4I	61	GLU
13	4I	64	TRP
13	4I	65	LYS
13	4I	66	LEU
13	4I	71	ARG
13	4I	73	GLU
13	4I	77	ASN
13	4I	82	MET
13	4I	84	ILE
13	4I	88	ARG
13	4I	96	LEU
13	4I	98	VAL
13	4I	105	THR
13	4I	108	ARG
13	4I	109	THR
13	4I	115	LYS
13	4I	116	THR
14	5I	7	ILE
14	5I	8	GLU
14	5I	16	PHE
14	5I	18	VAL
14	5I	41	ARG
14	5I	43	CYS
14	5I	44	LEU
14	5I	50	LYS

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Mol	Chain	Res	Type
15	6I	3	ILE
15	6I	4	THR
15	6I	13	GLN
15	6I	17	ARG
15	6I	21	ASP
15	6I	26	GLU
15	6I	31	LEU
15	6I	39	LEU
15	6I	41	GLU
15	6I	66	LEU
15	6I	82	ILE
15	6I	84	LYS
15	6I	87	ILE
16	7I	2	VAL
16	7I	8	ARG
16	7I	19	ILE
16	7I	20	VAL
16	7I	28	ARG
16	7I	44	THR
16	7I	45	THR
16	7I	67	THR
16	7I	68	ASP
16	7I	69	THR
16	7I	72	ARG
16	7I	80	PHE
16	7I	83	GLU
17	8I	5	VAL
17	8I	9	VAL
17	8I	25	ARG
17	8I	37	LYS
17	8I	38	ARG
17	8I	52	LYS
17	8I	59	ILE
17	8I	60	ILE
17	8I	74	LEU
17	8I	86	GLU
17	8I	97	SER
17	8I	101	ARG
18	9I	29	PHE
18	9I	32	ARG
18	9I	36	ASN
18	9I	38	GLU

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Mol	Chain	Res	Type
18	9I	40	LEU
18	9I	42	ARG
18	9I	46	GLU
18	9I	54	ARG
18	9I	76	LEU
18	9I	83	GLU
18	9I	84	LYS
18	9I	88	LYS
19	AI	5	LEU
19	AI	22	LEU
19	AI	27	GLU
19	AI	30	LEU
19	AI	31	ILE
19	AI	33	THR
19	AI	39	THR
19	AI	41	VAL
19	AI	43	GLU
19	AI	44	MET
19	AI	45	VAL
19	AI	47	HIS
19	AI	48	THR
19	AI	58	VAL
19	AI	63	THR
19	AI	67	VAL
19	AI	77	THR
19	AI	78	ARG
19	AI	83	HIS
20	BI	11	SER
20	BI	13	LEU
20	BI	16	HIS
20	BI	36	LEU
20	BI	55	ILE
20	BI	62	LEU
20	BI	73	HIS
20	BI	82	SER
20	BI	86	ARG
20	BI	88	VAL
20	BI	93	GLU
20	BI	99	LEU
20	BI	100	ILE
21	1F	25	LYS
27	7I	22	ILE

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Mol	Chain	Res	Type
27	71	30	LYS
27	71	34	THR
27	71	49	ILE
27	71	57	ASN
27	71	68	LEU
27	71	172	HIS
27	71	180	PHE
27	71	184	LYS
27	71	187	ASP
27	71	199	HIS
27	71	202	GLU
27	71	208	PHE
27	71	215	THR
27	71	216	THR
27	71	224	ILE
28	11	3	VAL
28	11	17	THR
28	11	18	VAL
28	11	27	THR
28	11	34	VAL
28	11	38	LYS
28	11	39	LYS
28	11	40	THR
28	11	52	ARG
28	11	61	LEU
28	11	65	ILE
28	11	72	LYS
28	11	78	LYS
28	11	94	LEU
28	11	95	LEU
28	11	101	GLU
28	11	103	ARG
28	11	105	ILE
28	11	106	ILE
28	11	111	LEU
28	11	112	GLN
28	11	113	VAL
28	11	122	ASP
28	11	127	VAL
28	11	136	ILE
28	11	141	VAL
28	11	147	LEU

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Mol	Chain	Res	Type
28	11	155	LEU
28	11	157	ARG
28	11	162	SER
28	11	165	ILE
28	11	171	ASP
28	11	173	VAL
28	11	175	LEU
28	11	183	ARG
28	11	192	THR
28	11	193	VAL
28	11	215	LEU
28	11	221	VAL
28	11	226	MET
28	11	237	GLU
28	11	242	ARG
28	11	257	LEU
28	11	260	ARG
28	11	261	LYS
28	11	270	ILE
28	11	271	ILE
28	11	273	ARG
29	21	7	VAL
29	21	9	VAL
29	21	14	ILE
29	21	17	ASP
29	21	18	ASP
29	21	21	VAL
29	21	23	VAL
29	21	24	THR
29	21	38	THR
29	21	41	LYS
29	21	45	THR
29	21	73	GLU
29	21	77	ILE
29	21	78	LEU
29	21	89	ASP
29	21	90	THR
29	21	111	ARG
29	21	116	VAL
29	21	117	MET
29	21	118	LYS
29	21	119	ARG

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Mol	Chain	Res	Type
29	21	134	ILE
29	21	143	ASN
29	21	152	LYS
29	21	154	LYS
29	21	167	VAL
29	21	175	VAL
29	21	179	GLU
29	21	184	VAL
29	21	198	VAL
30	31	6	VAL
30	31	8	GLN
30	31	17	ARG
30	31	24	LEU
30	31	28	ILE
30	31	33	LEU
30	31	53	THR
30	31	57	VAL
30	31	60	SER
30	31	65	TRP
30	31	67	GLN
30	31	68	LYS
30	31	70	THR
30	31	72	ARG
30	31	74	ARG
30	31	82	ILE
30	31	89	VAL
30	31	106	ARG
30	31	108	LYS
30	31	149	ASP
30	31	170	LEU
30	31	174	VAL
30	31	179	GLU
30	31	183	VAL
30	31	193	VAL
30	31	206	ILE
31	41	10	LYS
31	41	26	GLN
31	41	28	VAL
31	41	34	LEU
31	41	35	GLU
31	41	45	GLU
31	41	53	LEU

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Mol	Chain	Res	Type
31	41	54	GLU
31	41	63	ILE
31	41	67	LYS
31	41	70	VAL
31	41	82	LEU
31	41	88	ILE
31	41	96	ARG
31	41	99	MET
31	41	103	LEU
31	41	118	ARG
31	41	137	GLU
31	41	145	THR
31	41	147	ASP
31	41	159	VAL
31	41	161	THR
31	41	162	THR
31	41	166	ASP
31	41	174	GLU
32	51	4	ILE
32	51	6	ARG
32	51	7	LEU
32	51	24	VAL
32	51	26	VAL
32	51	33	LEU
32	51	37	VAL
32	51	40	GLU
32	51	50	VAL
32	51	56	SER
32	51	57	ASP
32	51	72	ILE
32	51	77	LYS
32	51	79	VAL
32	51	83	TYR
32	51	86	GLU
32	51	88	LEU
32	51	97	ARG
32	51	103	LEU
32	51	104	GLU
32	51	113	VAL
32	51	119	GLU
32	51	125	VAL
32	51	129	THR

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Mol	Chain	Res	Type
32	51	136	ILE
32	51	139	GLN
32	51	149	ARG
32	51	151	ILE
32	51	153	LYS
32	51	169	VAL
32	51	171	LEU
33	61	1	MET
33	61	4	ILE
33	61	7	GLU
33	61	10	GLU
33	61	12	LEU
33	61	14	ASP
33	61	18	VAL
33	61	25	TYR
33	61	27	ARG
33	61	33	ARG
33	61	35	LEU
33	61	38	LEU
33	61	41	GLU
33	61	43	ASN
33	61	56	LYS
33	61	57	ARG
33	61	64	GLU
33	61	67	ARG
33	61	68	LEU
33	61	70	GLU
33	61	71	ILE
33	61	75	LEU
33	61	78	THR
33	61	81	VAL
33	61	82	ARG
33	61	85	GLU
33	61	86	THR
33	61	92	VAL
33	61	93	THR
33	61	96	ASP
33	61	104	GLN
33	61	112	LYS
33	61	113	ARG
33	61	129	THR
33	61	133	HIS

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Mol	Chain	Res	Type
33	61	134	PRO
33	61	135	GLU
33	61	138	ILE
33	61	139	GLN
33	61	140	LEU
33	61	142	VAL
33	61	144	VAL
34	38	5	ARG
34	38	7	VAL
34	38	9	LEU
34	38	10	LEU
34	38	12	THR
34	38	13	LEU
34	38	14	LYS
34	38	18	GLU
34	38	21	GLN
34	38	25	PHE
34	38	27	VAL
34	38	29	TYR
34	38	37	THR
34	38	38	HIS
34	38	40	LEU
34	38	41	ARG
34	38	42	GLN
34	38	44	LEU
34	38	45	LYS
34	38	52	PHE
34	38	60	ARG
34	38	61	LEU
34	38	66	LEU
34	38	71	LEU
34	38	85	ASP
34	38	87	VAL
34	38	94	VAL
34	38	100	ASN
34	38	111	LEU
34	38	112	LEU
34	38	113	GLN
34	38	120	LYS
34	38	122	VAL
34	38	131	MET
34	38	142	LEU

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Mol	Chain	Res	Type
35	58	7	LYS
35	58	14	VAL
35	58	16	ILE
35	58	28	THR
35	58	34	LEU
35	58	35	ARG
35	58	43	THR
35	58	46	VAL
35	58	48	MET
35	58	58	ASP
35	58	60	ILE
35	58	61	ARG
35	58	62	VAL
35	58	67	LEU
35	58	70	LYS
35	58	73	THR
35	58	82	LEU
35	58	87	LEU
35	58	90	MET
35	58	91	LEU
35	58	96	GLU
35	58	98	VAL
35	58	99	LEU
35	58	109	LYS
35	58	112	LEU
35	58	114	ARG
35	58	120	LEU
35	58	134	ARG
35	58	136	GLU
36	68	5	GLN
36	68	22	ILE
36	68	24	VAL
36	68	39	ILE
36	68	47	ILE
36	68	53	LYS
36	68	57	VAL
36	68	61	VAL
36	68	66	LYS
36	68	68	GLU
36	68	89	ASN
36	68	96	THR
36	68	97	ARG

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Mol	Chain	Res	Type
36	68	98	VAL
36	68	108	GLU
36	68	115	VAL
37	78	1	MET
37	78	5	ASP
37	78	6	LEU
37	78	21	ARG
37	78	29	LYS
37	78	32	THR
37	78	40	SER
37	78	61	ARG
37	78	65	ARG
37	78	71	VAL
37	78	84	ASN
37	78	88	LEU
37	78	94	GLU
37	78	98	GLU
37	78	100	LEU
37	78	108	LYS
37	78	112	LEU
37	78	123	LEU
37	78	125	VAL
37	78	126	VAL
37	78	133	SER
37	78	135	LEU
37	78	138	LEU
37	78	139	LYS
37	78	144	GLU
37	78	146	VAL
37	78	148	LEU
38	88	5	ARG
38	88	18	LYS
38	88	21	THR
38	88	25	ASP
38	88	27	VAL
38	88	43	THR
38	88	45	GLN
38	88	54	MET
38	88	56	ARG
38	88	59	ARG
38	88	66	ILE
38	88	79	LEU

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Mol	Chain	Res	Type
38	88	80	GLU
38	88	90	VAL
38	88	91	GLU
38	88	101	ARG
38	88	109	VAL
38	88	110	THR
38	88	111	GLU
38	88	112	GLU
38	88	127	ILE
38	88	129	THR
38	88	130	LYS
39	98	1	MET
39	98	2	ARG
39	98	4	LEU
39	98	18	LEU
39	98	28	LEU
39	98	29	LEU
39	98	30	THR
39	98	36	THR
39	98	37	THR
39	98	44	LEU
39	98	51	LEU
39	98	54	LEU
39	98	65	LEU
39	98	67	LEU
39	98	73	VAL
39	98	75	LEU
39	98	79	LEU
39	98	98	LEU
39	98	100	LEU
39	98	104	ARG
39	98	105	ARG
39	98	113	LEU
39	98	116	LEU
40	A8	10	ARG
40	A8	11	LYS
40	A8	17	ARG
40	A8	20	ARG
40	A8	23	ARG
40	A8	25	ARG
40	A8	27	SER
40	A8	35	ILE

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Mol	Chain	Res	Type
40	A8	44	LYS
40	A8	46	VAL
40	A8	49	VAL
40	A8	52	SER
40	A8	54	LEU
40	A8	56	LEU
40	A8	57	LYS
40	A8	63	THR
40	A8	69	VAL
40	A8	73	LEU
40	A8	82	ILE
40	A8	88	ASP
40	A8	89	ARG
40	A8	98	VAL
40	A8	101	LEU
40	A8	106	ARG
40	A8	110	LEU
40	A8	111	GLU
41	B8	1	MET
41	B8	2	ASN
41	B8	6	LEU
41	B8	7	ILE
41	B8	9	LEU
41	B8	10	VAL
41	B8	21	GLU
41	B8	23	ARG
41	B8	27	THR
41	B8	28	VAL
41	B8	34	VAL
41	B8	35	LYS
41	B8	38	ASN
41	B8	42	ILE
41	B8	43	GLN
41	B8	44	ASP
41	B8	46	GLU
41	B8	51	ARG
41	B8	54	ARG
41	B8	55	ASN
41	B8	58	ASN
41	B8	62	THR
41	B8	65	LYS
41	B8	66	VAL

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Mol	Chain	Res	Type
41	B8	67	SER
41	B8	73	GLU
41	B8	74	ARG
41	B8	78	LEU
41	B8	84	GLN
41	B8	86	ILE
41	B8	87	ASP
41	B8	88	ILE
41	B8	89	VAL
41	B8	96	ARG
41	B8	99	LEU
41	B8	104	ASN
41	B8	107	ASP
41	B8	108	ARG
41	B8	110	ILE
41	B8	111	ARG
41	B8	112	ARG
41	B8	122	ASP
41	B8	125	ARG
41	B8	128	GLU
41	B8	134	GLU
42	C8	5	LYS
42	C8	9	VAL
42	C8	31	SER
42	C8	34	LYS
42	C8	36	ARG
42	C8	59	ARG
42	C8	64	ARG
42	C8	70	ARG
42	C8	74	LEU
42	C8	83	LEU
42	C8	91	ASP
42	C8	93	LYS
42	C8	98	LEU
42	C8	100	VAL
42	C8	101	ARG
42	C8	104	GLN
42	C8	108	GLU
42	C8	111	GLU
42	C8	114	LYS
42	C8	117	GLN
43	D8	1	MET

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Mol	Chain	Res	Type
43	D8	5	VAL
43	D8	6	LYS
43	D8	7	THR
43	D8	11	GLN
43	D8	18	LEU
43	D8	19	LYS
43	D8	21	ARG
43	D8	24	LYS
43	D8	33	VAL
43	D8	39	LEU
43	D8	40	LEU
43	D8	47	VAL
43	D8	51	VAL
43	D8	52	VAL
43	D8	57	VAL
43	D8	58	VAL
43	D8	61	VAL
43	D8	66	ARG
43	D8	70	ILE
43	D8	71	LEU
43	D8	72	VAL
43	D8	73	SER
43	D8	79	VAL
43	D8	82	ARG
43	D8	92	THR
43	D8	95	LEU
43	D8	99	ILE
44	E8	11	ARG
44	E8	16	LYS
44	E8	19	LEU
44	E8	20	VAL
44	E8	28	SER
44	E8	29	LEU
44	E8	37	ARG
44	E8	41	LYS
44	E8	50	VAL
44	E8	51	LEU
44	E8	63	ASP
44	E8	65	LEU
44	E8	69	LEU
44	E8	71	VAL
44	E8	76	VAL

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Mol	Chain	Res	Type
44	E8	78	GLU
44	E8	88	ARG
44	E8	94	ASP
44	E8	95	ILE
44	E8	97	LYS
44	E8	103	ILE
44	E8	106	ILE
44	E8	107	LEU
45	F8	1	MET
45	F8	3	THR
45	F8	15	GLU
45	F8	27	THR
45	F8	28	PHE
45	F8	48	LYS
45	F8	49	VAL
45	F8	57	LEU
45	F8	65	ARG
45	F8	70	LEU
45	F8	78	LYS
45	F8	80	ILE
45	F8	81	VAL
45	F8	83	VAL
45	F8	88	LYS
45	F8	92	LEU
46	G8	3	VAL
46	G8	5	MET
46	G8	7	VAL
46	G8	14	LEU
46	G8	21	LYS
46	G8	27	VAL
46	G8	38	ILE
46	G8	44	ILE
46	G8	54	LYS
46	G8	57	GLN
46	G8	67	LEU
46	G8	72	VAL
46	G8	75	ILE
46	G8	81	LYS
46	G8	84	ARG
46	G8	85	VAL
46	G8	90	LEU
46	G8	91	GLU

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Mol	Chain	Res	Type
46	G8	92	ASN
46	G8	94	LYS
46	G8	95	LYS
46	G8	98	VAL
47	H8	2	GLU
47	H8	5	LEU
47	H8	19	ARG
47	H8	32	HIS
47	H8	35	ARG
47	H8	42	VAL
47	H8	50	GLN
47	H8	53	ILE
47	H8	58	VAL
47	H8	61	LEU
47	H8	67	LEU
47	H8	70	LEU
47	H8	71	VAL
47	H8	72	ARG
47	H8	76	LEU
47	H8	77	ASP
47	H8	80	ARG
47	H8	86	VAL
47	H8	91	LEU
47	H8	93	ASP
47	H8	97	GLU
47	H8	117	LEU
47	H8	119	GLU
47	H8	121	HIS
47	H8	122	ARG
47	H8	126	VAL
47	H8	132	ASN
47	H8	135	GLU
47	H8	137	ILE
47	H8	140	ASP
47	H8	154	ASP
47	H8	155	LEU
47	H8	156	LYS
47	H8	157	LEU
47	H8	162	GLU
47	H8	170	THR
47	H8	171	ILE
48	I8	12	ASN

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Mol	Chain	Res	Type
48	I8	36	ILE
48	I8	38	VAL
48	I8	40	GLN
48	I8	41	ARG
48	I8	58	THR
48	I8	77	ARG
48	I8	80	HIS
49	J8	26	ARG
49	J8	41	ARG
49	J8	46	LEU
49	J8	51	VAL
49	J8	56	GLN
49	J8	69	LYS
49	J8	81	LYS
49	J8	83	GLU
49	J8	85	LEU
49	J8	91	LYS
50	K8	3	LEU
50	K8	4	SER
50	K8	5	GLU
50	K8	8	LYS
50	K8	10	LEU
50	K8	11	GLU
50	K8	12	GLU
50	K8	16	LEU
50	K8	19	VAL
50	K8	32	LEU
50	K8	33	MET
50	K8	35	LEU
50	K8	45	SER
50	K8	47	ASN
50	K8	48	HIS
50	K8	50	ILE
50	K8	53	LEU
50	K8	55	ARG
50	K8	59	ARG
50	K8	62	THR
50	K8	63	VAL
50	K8	69	ARG
51	L8	8	LEU
51	L8	9	VAL
51	L8	13	ILE

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Mol	Chain	Res	Type
51	L8	28	LEU
51	L8	29	ARG
51	L8	31	LEU
51	L8	33	GLN
51	L8	38	GLU
51	L8	40	THR
51	L8	44	ARG
51	L8	54	VAL
51	L8	60	GLU
52	M8	5	ILE
52	M8	9	LEU
52	M8	10	VAL
52	M8	14	ILE
52	M8	15	ILE
52	M8	18	CYS
52	M8	21	VAL
52	M8	27	THR
52	M8	35	VAL
52	M8	38	LYS
52	M8	42	PHE
52	M8	43	TYR
52	M8	48	ARG
52	M8	57	GLU
52	M8	59	PHE
52	M8	61	ARG
53	N8	6	VAL
53	N8	13	LYS
53	N8	29	THR
53	N8	31	VAL
53	N8	37	LYS
53	N8	40	LYS
53	N8	44	THR
53	N8	49	CYS
53	N8	51	TYR
53	N8	56	LYS
53	N8	57	VAL
54	O8	5	VAL
54	O8	6	ARG
54	O8	7	ILE
54	O8	8	LYS
54	O8	9	LEU
54	O8	10	LEU

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Mol	Chain	Res	Type
54	O8	11	LEU
54	O8	13	CYS
54	O8	14	THR
54	O8	16	CYS
54	O8	18	ARG
54	O8	20	ASN
54	O8	27	LYS
54	O8	30	THR
54	O8	34	LEU
54	O8	35	GLU
54	O8	37	ARG
54	O8	48	VAL
54	O8	49	HIS
54	O8	50	ARG
55	P8	1	MET
55	P8	4	THR
55	P8	11	LYS
55	P8	46	VAL
55	P8	48	LYS
56	Q8	6	THR
56	Q8	12	LYS
56	Q8	14	VAL
56	Q8	34	TRP
56	Q8	35	GLN
56	Q8	50	LEU
2	12	7	VAL
2	12	10	LEU
2	12	11	LEU
2	12	12	GLU
2	12	17	PHE
2	12	21	ARG
2	12	27	LYS
2	12	30	ARG
2	12	31	TYR
2	12	32	ILE
2	12	33	TYR
2	12	35	GLU
2	12	42	ILE
2	12	44	LEU
2	12	73	THR
2	12	75	LYS
2	12	76	GLN

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Mol	Chain	Res	Type
2	12	80	ILE
2	12	81	VAL
2	12	84	GLU
2	12	101	MET
2	12	103	THR
2	12	108	ILE
2	12	111	ARG
2	12	114	ARG
2	12	115	LEU
2	12	116	GLU
2	12	117	GLU
2	12	127	ILE
2	12	128	GLU
2	12	134	GLU
2	12	136	VAL
2	12	139	LYS
2	12	142	LEU
2	12	146	GLN
2	12	149	LEU
2	12	154	LEU
2	12	158	LEU
2	12	163	PHE
2	12	165	VAL
2	12	168	THR
2	12	178	ARG
2	12	185	ILE
2	12	189	ASP
2	12	192	SER
2	12	196	LEU
2	12	198	ASP
2	12	201	ILE
2	12	208	ILE
2	12	209	ARG
2	12	219	VAL
2	12	226	ARG
2	12	229	VAL
2	12	235	SER
2	12	238	LEU
2	12	240	GLN
3	22	5	ILE
3	22	12	LEU
3	22	14	ILE

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Mol	Chain	Res	Type
3	22	16	ARG
3	22	17	ASP
3	22	23	TYR
3	22	26	LYS
3	22	28	GLN
3	22	34	LEU
3	22	43	LEU
3	22	44	GLU
3	22	62	ASP
3	22	64	VAL
3	22	66	VAL
3	22	67	THR
3	22	72	LYS
3	22	75	VAL
3	22	77	ILE
3	22	82	GLU
3	22	83	ARG
3	22	86	VAL
3	22	94	LEU
3	22	101	LEU
3	22	102	ASN
3	22	103	VAL
3	22	116	VAL
3	22	119	ARG
3	22	122	GLU
3	22	127	ARG
3	22	128	PHE
3	22	136	GLN
3	22	140	ARG
3	22	143	GLU
3	22	151	VAL
3	22	154	SER
3	22	161	GLU
3	22	165	THR
3	22	166	GLU
3	22	173	VAL
3	22	178	LEU
3	22	186	PHE
3	22	190	ARG
3	22	196	LEU
3	22	206	GLU
3	22	207	VAL

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Mol	Chain	Res	Type
4	32	3	ARG
4	32	8	VAL
4	32	12	CYS
4	32	13	ARG
4	32	14	ARG
4	32	15	GLU
4	32	19	LEU
4	32	30	LYS
4	32	33	MET
4	32	58	LEU
4	32	60	GLU
4	32	77	ASN
4	32	78	LEU
4	32	107	ARG
4	32	115	ARG
4	32	127	THR
4	32	134	ASP
4	32	135	LEU
4	32	150	GLU
4	32	151	LYS
4	32	155	LEU
4	32	162	LEU
4	32	178	VAL
4	32	186	LEU
5	42	3	GLU
5	42	6	PHE
5	42	8	GLU
5	42	10	MET
5	42	12	LEU
5	42	13	ILE
5	42	25	ARG
5	42	47	LYS
5	42	53	LEU
5	42	63	ARG
5	42	64	ARG
5	42	66	MET
5	42	72	GLN
5	42	73	ASN
5	42	75	THR
5	42	78	HIS
5	42	79	GLU
5	42	80	ILE

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Mol	Chain	Res	Type
5	42	82	VAL
5	42	83	GLU
5	42	89	ILE
5	42	90	VAL
5	42	100	VAL
5	42	116	THR
5	42	118	ILE
5	42	126	ARG
5	42	131	ILE
5	42	135	THR
5	42	144	THR
5	42	150	ARG
5	42	152	ARG
6	52	16	GLN
6	52	28	ARG
6	52	40	VAL
6	52	54	LYS
6	52	74	ASP
6	52	77	ARG
6	52	85	VAL
7	62	8	GLU
7	62	9	VAL
7	62	12	LEU
7	62	22	LEU
7	62	24	THR
7	62	28	ASN
7	62	30	ILE
7	62	32	ARG
7	62	54	THR
7	62	63	LYS
7	62	67	GLU
7	62	72	ARG
7	62	78	ARG
7	62	101	LEU
7	62	113	GLU
7	62	118	VAL
7	62	124	LEU
7	62	131	LYS
7	62	135	VAL
7	62	137	LYS
7	62	151	TYR
8	72	1	MET

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Mol	Chain	Res	Type
8	72	17	THR
8	72	18	ARG
8	72	21	LYS
8	72	22	GLU
8	72	38	ILE
8	72	52	ASP
8	72	56	LYS
8	72	80	ILE
8	72	99	GLU
8	72	102	ARG
8	72	103	VAL
8	72	105	ARG
8	72	109	ILE
8	72	114	THR
8	72	115	SER
8	72	119	LEU
8	72	122	ARG
8	72	133	LEU
8	72	137	VAL
9	82	1	MET
9	82	3	GLN
9	82	10	ARG
9	82	19	LEU
9	82	25	LYS
9	82	32	ASP
9	82	38	GLN
9	82	40	LEU
9	82	47	LEU
9	82	56	LEU
9	82	58	HIS
9	82	78	LYS
9	82	83	ARG
9	82	87	GLN
9	82	95	LYS
9	82	96	LEU
9	82	99	LEU
9	82	102	LEU
9	82	110	GLU
9	82	112	LYS
9	82	125	TYR
10	1A	16	LEU
10	1A	47	PHE

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Mol	Chain	Res	Type
10	1A	49	VAL
10	1A	50	ILE
10	1A	57	LYS
10	1A	62	HIS
10	1A	65	LEU
10	1A	68	HIS
10	1A	70	ARG
10	1A	74	ILE
10	1A	75	ILE
10	1A	76	ASN
10	1A	80	LYS
10	1A	82	ILE
10	1A	87	THR
10	1A	89	ASP
10	1A	90	LEU
10	1A	96	ILE
10	1A	97	GLU
11	2A	9	LYS
11	2A	13	GLN
11	2A	29	ILE
11	2A	31	THR
11	2A	47	VAL
11	2A	63	LEU
11	2A	66	LEU
11	2A	81	ASP
11	2A	82	VAL
11	2A	87	THR
11	2A	96	ARG
11	2A	103	LEU
11	2A	105	VAL
11	2A	112	THR
11	2A	114	VAL
11	2A	120	ARG
11	2A	123	LYS
11	2A	127	LYS
12	3A	8	VAL
12	3A	15	VAL
12	3A	20	LYS
12	3A	21	VAL
12	3A	24	LEU
12	3A	30	ARG
12	3A	34	CYS

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Mol	Chain	Res	Type
12	3A	39	THR
12	3A	41	THR
12	3A	43	LYS
12	3A	51	LYS
12	3A	52	VAL
12	3A	70	GLU
12	3A	79	VAL
12	3A	81	LEU
12	3A	82	ILE
12	3A	83	ARG
12	3A	86	ARG
12	3A	101	VAL
12	3A	119	THR
12	3A	123	LYS
13	4A	4	ILE
13	4A	7	VAL
13	4A	8	GLU
13	4A	11	ARG
13	4A	15	VAL
13	4A	17	VAL
13	4A	22	ILE
13	4A	35	GLU
13	4A	37	THR
13	4A	39	ILE
13	4A	44	ARG
13	4A	45	VAL
13	4A	46	LYS
13	4A	47	ASP
13	4A	48	LEU
13	4A	49	THR
13	4A	50	GLU
13	4A	56	LEU
13	4A	67	GLU
13	4A	70	LEU
13	4A	73	GLU
13	4A	74	VAL
13	4A	79	LYS
13	4A	81	LEU
13	4A	105	THR
13	4A	108	ARG
13	4A	111	LYS
13	4A	116	THR

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Mol	Chain	Res	Type
14	5A	15	LYS
14	5A	50	LYS
15	6A	3	ILE
15	6A	6	GLU
15	6A	10	LYS
15	6A	24	SER
15	6A	27	VAL
15	6A	35	ARG
15	6A	39	LEU
15	6A	41	GLU
15	6A	48	LYS
15	6A	56	LEU
15	6A	58	MET
15	6A	64	ARG
15	6A	87	ILE
16	7A	2	VAL
16	7A	5	ARG
16	7A	20	VAL
16	7A	27	LYS
16	7A	29	ASP
16	7A	45	THR
16	7A	53	VAL
16	7A	67	THR
16	7A	69	THR
16	7A	72	ARG
16	7A	79	VAL
17	8A	6	LEU
17	8A	7	THR
17	8A	19	VAL
17	8A	21	VAL
17	8A	25	ARG
17	8A	35	VAL
17	8A	52	LYS
17	8A	53	LEU
17	8A	55	ASP
17	8A	59	ILE
17	8A	73	VAL
17	8A	74	LEU
17	8A	78	GLU
18	9A	18	ARG
18	9A	19	LYS
18	9A	23	LYS

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Mol	Chain	Res	Type
18	9A	25	THR
18	9A	31	LEU
18	9A	32	ARG
18	9A	36	ASN
18	9A	42	ARG
18	9A	58	LEU
18	9A	65	ILE
18	9A	68	LYS
18	9A	74	ARG
18	9A	84	LYS
18	9A	88	LYS
19	AA	5	LEU
19	AA	6	LYS
19	AA	7	LYS
19	AA	9	VAL
19	AA	15	LEU
19	AA	16	LEU
19	AA	22	LEU
19	AA	27	GLU
19	AA	31	ILE
19	AA	33	THR
19	AA	40	ILE
19	AA	60	VAL
19	AA	64	GLU
19	AA	66	MET
19	AA	67	VAL
19	AA	69	HIS
20	BA	10	LEU
20	BA	23	ARG
20	BA	24	LEU
20	BA	36	LEU
20	BA	41	ILE
20	BA	56	MET
20	BA	74	LYS
20	BA	84	LEU
20	BA	86	ARG
20	BA	87	LYS
20	BA	88	VAL
20	BA	104	LEU
21	1B	6	ARG
21	1B	8	THR
21	1B	13	ILE

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Mol	Chain	Res	Type
21	1B	15	ARG
21	1B	18	TYR
21	1B	25	LYS
28	19	3	VAL
28	19	4	LYS
28	19	10	THR
28	19	17	THR
28	19	18	VAL
28	19	54	ARG
28	19	61	LEU
28	19	64	ILE
28	19	65	ILE
28	19	72	LYS
28	19	79	VAL
28	19	94	LEU
28	19	95	LEU
28	19	101	GLU
28	19	105	ILE
28	19	106	ILE
28	19	111	LEU
28	19	136	ILE
28	19	138	VAL
28	19	157	ARG
28	19	161	THR
28	19	166	GLN
28	19	173	VAL
28	19	182	LEU
28	19	185	VAL
28	19	192	THR
28	19	200	ASP
28	19	211	ARG
28	19	255	LYS
28	19	257	LEU
28	19	261	LYS
28	19	271	ILE
29	29	9	VAL
29	29	14	ILE
29	29	21	VAL
29	29	24	THR
29	29	34	VAL
29	29	38	THR
29	29	40	GLU

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Mol	Chain	Res	Type
29	29	42	ASP
29	29	45	THR
29	29	52	LEU
29	29	72	VAL
29	29	77	ILE
29	29	78	LEU
29	29	87	GLU
29	29	89	ASP
29	29	92	THR
29	29	116	VAL
29	29	119	ARG
29	29	134	ILE
29	29	143	ASN
29	29	144	ARG
29	29	152	LYS
29	29	154	LYS
29	29	167	VAL
29	29	174	ASP
29	29	175	VAL
29	29	181	LEU
29	29	182	LEU
29	29	184	VAL
29	29	196	VAL
30	39	3	GLU
30	39	8	GLN
30	39	20	LEU
30	39	24	LEU
30	39	28	ILE
30	39	32	LEU
30	39	33	LEU
30	39	43	LYS
30	39	52	LYS
30	39	53	THR
30	39	64	ILE
30	39	68	LYS
30	39	70	THR
30	39	72	ARG
30	39	89	VAL
30	39	106	ARG
30	39	123	LEU
30	39	124	LEU
30	39	125	LEU

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Mol	Chain	Res	Type
30	39	138	GLU
30	39	158	THR
30	39	165	ARG
30	39	174	VAL
30	39	183	VAL
30	39	186	ILE
30	39	190	GLU
30	39	191	ARG
30	39	193	VAL
30	39	199	TRP
30	39	206	ILE
31	49	5	VAL
31	49	7	LEU
31	49	10	LYS
31	49	21	ARG
31	49	22	ARG
31	49	28	VAL
31	49	31	VAL
31	49	34	LEU
31	49	35	GLU
31	49	43	LEU
31	49	47	LYS
31	49	54	GLU
31	49	60	LEU
31	49	72	ARG
31	49	81	LYS
31	49	86	MET
31	49	88	ILE
31	49	106	LEU
31	49	114	ILE
31	49	115	ARG
31	49	118	ARG
31	49	133	LEU
31	49	138	GLN
31	49	139	LEU
31	49	147	ASP
31	49	148	MET
31	49	155	MET
31	49	159	VAL
31	49	174	GLU
32	59	4	ILE
32	59	7	LEU

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Mol	Chain	Res	Type
32	59	9	ILE
32	59	13	LYS
32	59	23	ARG
32	59	37	VAL
32	59	41	MET
32	59	42	ARG
32	59	45	VAL
32	59	46	GLU
32	59	50	VAL
32	59	51	ARG
32	59	56	SER
32	59	70	THR
32	59	72	ILE
32	59	81	GLU
32	59	83	TYR
32	59	88	LEU
32	59	95	ARG
32	59	98	LEU
32	59	104	GLU
32	59	105	LEU
32	59	107	VAL
32	59	114	VAL
32	59	122	THR
32	59	131	VAL
32	59	132	ARG
32	59	138	LYS
32	59	153	LYS
32	59	164	TYR
32	59	169	VAL
33	69	1	MET
33	69	3	VAL
33	69	4	ILE
33	69	5	LEU
33	69	6	LEU
33	69	7	GLU
33	69	9	LEU
33	69	10	GLU
33	69	18	VAL
33	69	48	GLU
33	69	52	ARG
33	69	56	LYS
33	69	60	GLU

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Mol	Chain	Res	Type
33	69	64	GLU
33	69	75	LEU
33	69	78	THR
33	69	81	VAL
33	69	86	THR
33	69	87	LYS
33	69	101	LEU
33	69	105	HIS
33	69	109	ILE
33	69	110	ASP
33	69	114	LEU
33	69	117	GLU
33	69	122	GLU
33	69	135	GLU
33	69	136	VAL
33	69	138	ILE
33	69	144	VAL
33	69	145	VAL
35	15	9	VAL
35	15	16	ILE
35	15	17	ASP
35	15	28	THR
35	15	29	LYS
35	15	32	THR
35	15	48	MET
35	15	62	VAL
35	15	71	ILE
35	15	85	ILE
35	15	94	HIS
35	15	98	VAL
35	15	99	LEU
35	15	109	LYS
35	15	112	LEU
35	15	120	LEU
35	15	131	GLN
35	15	136	GLU
35	15	137	LYS
36	25	3	GLN
36	25	5	GLN
36	25	8	LEU
36	25	10	VAL
36	25	12	ASP

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Mol	Chain	Res	Type
36	25	14	THR
36	25	23	ARG
36	25	28	SER
36	25	43	VAL
36	25	58	VAL
36	25	65	THR
36	25	69	ILE
36	25	82	ASN
36	25	87	ILE
36	25	89	ASN
36	25	96	THR
36	25	97	ARG
36	25	98	VAL
36	25	108	GLU
36	25	114	ILE
37	35	2	LYS
37	35	5	ASP
37	35	6	LEU
37	35	13	ASN
37	35	14	LYS
37	35	19	VAL
37	35	21	ARG
37	35	41	ARG
37	35	52	GLU
37	35	58	THR
37	35	59	LEU
37	35	64	LYS
37	35	68	GLN
37	35	75	ILE
37	35	79	ARG
37	35	81	GLN
37	35	87	ASP
37	35	88	LEU
37	35	91	PHE
37	35	101	VAL
37	35	110	TYR
37	35	111	ARG
37	35	112	LEU
37	35	114	ILE
37	35	117	GLU
37	35	123	LEU
37	35	125	VAL

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Mol	Chain	Res	Type
37	35	135	LEU
37	35	136	GLU
37	35	138	LEU
37	35	144	GLU
37	35	147	LEU
38	45	1	MET
38	45	3	MET
38	45	10	ARG
38	45	22	LYS
38	45	45	GLN
38	45	51	ARG
38	45	54	MET
38	45	55	VAL
38	45	56	ARG
38	45	58	PHE
38	45	59	ARG
38	45	64	ILE
38	45	66	ILE
38	45	80	GLU
38	45	81	VAL
38	45	89	ASN
38	45	109	VAL
38	45	115	MET
38	45	128	LYS
38	45	138	ASP
38	45	139	GLU
39	55	8	ARG
39	55	15	SER
39	55	24	GLN
39	55	35	THR
39	55	54	LEU
39	55	65	LEU
39	55	67	LEU
39	55	74	LYS
39	55	75	LEU
39	55	76	VAL
39	55	79	LEU
39	55	86	ARG
39	55	105	ARG
39	55	107	ASP
39	55	114	VAL
39	55	115	GLU

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Mol	Chain	Res	Type
39	55	117	VAL
39	55	118	GLU
40	65	8	GLU
40	65	14	VAL
40	65	20	ARG
40	65	24	LEU
40	65	34	HIS
40	65	40	ILE
40	65	58	LEU
40	65	59	LYS
40	65	73	LEU
40	65	84	GLN
40	65	85	VAL
40	65	93	LYS
40	65	98	VAL
40	65	106	ARG
41	75	1	MET
41	75	7	ILE
41	75	10	VAL
41	75	13	ARG
41	75	17	THR
41	75	19	LEU
41	75	23	ARG
41	75	27	THR
41	75	30	VAL
41	75	38	ASN
41	75	39	ARG
41	75	42	ILE
41	75	50	ILE
41	75	55	ASN
41	75	58	ASN
41	75	62	THR
41	75	64	ARG
41	75	65	LYS
41	75	78	LEU
41	75	82	LEU
41	75	86	ILE
41	75	88	ILE
41	75	98	LYS
41	75	102	ILE
41	75	105	LEU
41	75	111	ARG

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Mol	Chain	Res	Type
41	75	112	ARG
41	75	113	LYS
41	75	125	ARG
41	75	132	LYS
42	85	5	LYS
42	85	8	VAL
42	85	17	ILE
42	85	27	LEU
42	85	52	ARG
42	85	54	LYS
42	85	55	ARG
42	85	63	VAL
42	85	65	ILE
42	85	74	LEU
42	85	75	ASN
42	85	83	LEU
42	85	85	LYS
42	85	98	LEU
42	85	101	ARG
42	85	104	GLN
43	95	5	VAL
43	95	10	LYS
43	95	15	GLU
43	95	19	LYS
43	95	28	GLU
43	95	35	LEU
43	95	39	LEU
43	95	40	LEU
43	95	47	VAL
43	95	51	VAL
43	95	57	VAL
43	95	58	VAL
43	95	60	GLU
43	95	61	VAL
43	95	64	HIS
43	95	66	ARG
43	95	71	LEU
43	95	79	VAL
43	95	80	GLN
43	95	81	TYR
43	95	85	LYS
43	95	91	TYR

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Mol	Chain	Res	Type
43	95	96	ILE
44	A5	1	MET
44	A5	11	ARG
44	A5	18	ARG
44	A5	19	LEU
44	A5	20	VAL
44	A5	24	ILE
44	A5	31	GLU
44	A5	39	THR
44	A5	52	GLU
44	A5	67	ASP
44	A5	69	LEU
44	A5	70	TYR
44	A5	76	VAL
44	A5	88	ARG
44	A5	94	ASP
44	A5	96	ILE
44	A5	97	LYS
44	A5	105	VAL
44	A5	106	ILE
44	A5	107	LEU
45	B5	6	ASP
45	B5	8	ILE
45	B5	27	THR
45	B5	28	PHE
45	B5	30	VAL
45	B5	43	VAL
45	B5	48	LYS
45	B5	51	VAL
45	B5	54	VAL
45	B5	63	LYS
45	B5	66	LEU
45	B5	80	ILE
45	B5	81	VAL
46	C5	3	VAL
46	C5	6	HIS
46	C5	9	LYS
46	C5	21	LYS
46	C5	24	VAL
46	C5	27	VAL
46	C5	38	ILE
46	C5	39	VAL

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Mol	Chain	Res	Type
46	C5	49	VAL
46	C5	50	ARG
46	C5	52	SER
46	C5	54	LYS
46	C5	55	TYR
46	C5	56	PRO
46	C5	61	ILE
46	C5	72	VAL
46	C5	75	ILE
46	C5	81	LYS
46	C5	84	ARG
46	C5	85	VAL
46	C5	89	PHE
46	C5	90	LEU
46	C5	92	ASN
46	C5	94	LYS
46	C5	95	LYS
46	C5	96	ILE
46	C5	102	CYS
46	C5	106	LEU
46	C5	109	GLU
46	C5	110	GLU
47	D5	4	ARG
47	D5	5	LEU
47	D5	9	TYR
47	D5	10	ARG
47	D5	19	ARG
47	D5	24	LEU
47	D5	28	MET
47	D5	33	LEU
47	D5	40	ASP
47	D5	53	ILE
47	D5	55	HIS
47	D5	56	VAL
47	D5	59	LEU
47	D5	61	LEU
47	D5	70	LEU
47	D5	71	VAL
47	D5	76	LEU
47	D5	80	ARG
47	D5	81	ARG
47	D5	87	ASP

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Mol	Chain	Res	Type
47	D5	93	ASP
47	D5	96	VAL
47	D5	98	MET
47	D5	103	ARG
47	D5	104	PHE
47	D5	112	ARG
47	D5	117	LEU
47	D5	119	GLU
47	D5	120	ILE
47	D5	123	ASP
47	D5	128	VAL
47	D5	131	ARG
47	D5	132	ASN
47	D5	133	ILE
47	D5	135	GLU
47	D5	138	GLU
47	D5	142	SER
47	D5	149	SER
47	D5	157	LEU
47	D5	165	VAL
47	D5	171	ILE
47	D5	174	VAL
47	D5	175	VAL
48	E5	7	LEU
48	E5	9	SER
48	E5	20	ARG
48	E5	27	GLU
48	E5	31	VAL
48	E5	36	ILE
48	E5	37	LEU
48	E5	49	LYS
48	E5	53	MET
48	E5	58	THR
48	E5	64	ASP
48	E5	71	ASP
49	F5	4	VAL
49	F5	7	ILE
49	F5	13	ILE
49	F5	17	SER
49	F5	41	ARG
49	F5	46	LEU
49	F5	51	VAL

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Mol	Chain	Res	Type
49	F5	53	VAL
49	F5	59	THR
49	F5	67	ILE
49	F5	72	GLU
49	F5	76	ARG
49	F5	78	LYS
49	F5	80	LEU
49	F5	82	LEU
49	F5	83	GLU
49	F5	85	LEU
49	F5	88	LYS
49	F5	90	ILE
49	F5	91	LYS
49	F5	93	GLU
50	G5	4	SER
50	G5	5	GLU
50	G5	6	VAL
50	G5	9	GLN
50	G5	15	LYS
50	G5	20	GLU
50	G5	34	GLU
50	G5	35	LEU
50	G5	41	ILE
50	G5	44	LEU
50	G5	49	LYS
50	G5	50	ILE
50	G5	52	ASP
50	G5	55	ARG
50	G5	60	LEU
50	G5	64	LEU
51	H5	9	VAL
51	H5	24	LYS
51	H5	32	GLN
51	H5	40	THR
51	H5	44	ARG
51	H5	56	VAL
52	I5	2	LYS
52	I5	6	HIS
52	I5	13	ARG
52	I5	14	ILE
52	I5	16	CYS
52	I5	20	ASN

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Mol	Chain	Res	Type
52	I5	21	VAL
52	I5	34	GLU
52	I5	35	VAL
52	I5	40	HIS
52	I5	42	PHE
52	I5	44	THR
52	I5	46	GLN
52	I5	53	GLU
52	I5	55	ARG
52	I5	56	VAL
52	I5	58	ARG
52	I5	59	PHE
52	I5	61	ARG
52	I5	69	LYS
53	J5	6	VAL
53	J5	25	LEU
53	J5	36	CYS
53	J5	39	MET
53	J5	40	LYS
53	J5	44	THR
53	J5	48	GLU
53	J5	49	CYS
53	J5	51	TYR
53	J5	52	TYR
54	K5	7	ILE
54	K5	8	LYS
54	K5	9	LEU
54	K5	17	LYS
54	K5	23	THR
54	K5	28	ARG
54	K5	37	ARG
54	K5	38	LYS
54	K5	39	TYR
54	K5	43	CYS
54	K5	44	ARG
54	K5	45	LYS
54	K5	47	THR
55	L5	1	MET
55	L5	4	THR
55	L5	9	ARG
55	L5	24	THR
55	L5	41	ARG

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Mol	Chain	Res	Type
55	L5	42	LEU
55	L5	43	THR
55	L5	46	VAL
56	M5	6	THR
56	M5	16	ILE
56	M5	34	TRP
56	M5	35	GLN
56	M5	59	LYS
56	M5	62	LEU

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (69) such sidechains are listed below:

Mol	Chain	Res	Type
2	1E	146	GLN
3	2E	98	ASN
4	3E	42	GLN
5	4E	38	GLN
5	4E	73	ASN
5	4E	78	HIS
7	6E	13	GLN
7	6E	37	ASN
7	6E	64	GLN
7	6E	68	ASN
7	6E	86	GLN
7	6E	106	GLN
9	8E	3	GLN
9	8E	23	ASN
10	1I	56	HIS
11	2I	99	GLN
13	4I	12	ASN
13	4I	40	ASN
13	4I	77	ASN
15	6I	50	HIS
17	8I	26	GLN
27	7I	17	ASN
28	11	115	GLN
28	11	220	HIS
28	11	227	ASN
29	21	35	GLN
31	41	27	ASN
34	38	113	GLN
34	38	143	GLN

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Mol	Chain	Res	Type
35	58	69	GLN
39	98	31	HIS
41	B8	58	ASN
42	C8	75	ASN
42	C8	117	GLN
43	D8	11	GLN
43	D8	80	GLN
44	E8	40	ASN
47	H8	54	HIS
50	K8	38	GLN
51	L8	46	ASN
3	22	69	HIS
4	32	123	HIS
4	32	161	ASN
6	52	7	ASN
7	62	109	ASN
8	72	81	HIS
9	82	31	GLN
9	82	87	GLN
11	2A	26	ASN
11	2A	38	ASN
12	3A	6	GLN
14	5A	52	GLN
15	6A	37	ASN
19	AA	53	ASN
20	BA	42	GLN
29	29	35	GLN
29	29	48	GLN
29	29	132	HIS
29	29	169	ASN
30	39	40	GLN
31	49	66	GLN
31	49	121	ASN
35	15	69	GLN
38	45	45	GLN
41	75	58	ASN
47	D5	151	HIS
49	F5	66	HIS
50	G5	46	GLN
51	H5	32	GLN

5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	13	1504/1522 (98%)	361 (24%)	47 (3%)
1	1G	1508/1522 (99%)	398 (26%)	53 (3%)
22	1K	64/77 (83%)	20 (31%)	2 (3%)
22	3K	76/77 (98%)	23 (30%)	0
22	3L	76/77 (98%)	31 (40%)	2 (2%)
23	2K	76/77 (98%)	9 (11%)	1 (1%)
24	4K	16/27 (59%)	10 (62%)	2 (12%)
24	4L	20/27 (74%)	8 (40%)	1 (5%)
25	14	2876/2917 (98%)	759 (26%)	50 (1%)
25	1H	2874/2917 (98%)	676 (23%)	63 (2%)
26	16	119/122 (97%)	28 (23%)	2 (1%)
26	1J	121/122 (99%)	40 (33%)	3 (2%)
57	2L	76/77 (98%)	11 (14%)	0
All	All	9406/9561 (98%)	2374 (25%)	226 (2%)

All (2374) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	13	5	U
1	13	6	G
1	13	7	G
1	13	8	A
1	13	32	A
1	13	35	G
1	13	39	G
1	13	44	G
1	13	47	C
1	13	48	C
1	13	50	A
1	13	51	A
1	13	61	G
1	13	64	G
1	13	65	U
1	13	66	G
1	13	69	G
1	13	77	C
1	13	78	G
1	13	81	G
1	13	89	U
1	13	90	C
1	13	91	C
1	13	92	G
1	13	95	G

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Mol	Chain	Res	Type
1	13	101	A
1	13	115	G
1	13	120	A
1	13	121	C
1	13	122	G
1	13	129(A)	G
1	13	131	C
1	13	135	C
1	13	144	G
1	13	147	G
1	13	151	A
1	13	163	C
1	13	172	A
1	13	173	U
1	13	186(F)	C
1	13	188	U
1	13	190	G
1	13	191(A)	G
1	13	191(E)	G
1	13	195	A
1	13	196	A
1	13	197	A
1	13	201	C
1	13	209	U
1	13	210	U
1	13	217	C
1	13	222	U
1	13	226	G
1	13	243	A
1	13	244	U
1	13	245	C
1	13	247	G
1	13	248	C
1	13	250	A
1	13	251	G
1	13	264	U
1	13	266	G
1	13	267	C
1	13	270	A
1	13	281	G
1	13	289	G
1	13	314	C

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Mol	Chain	Res	Type
1	13	321	A
1	13	328	C
1	13	329	A
1	13	332	G
1	13	342	C
1	13	344	A
1	13	345	C
1	13	346	G
1	13	351	G
1	13	352	C
1	13	353	A
1	13	354	G
1	13	367	U
1	13	372	C
1	13	382	A
1	13	384	G
1	13	388	G
1	13	390	C
1	13	397	A
1	13	398	C
1	13	406	G
1	13	409	G
1	13	411	A
1	13	412	A
1	13	413	G
1	13	414	A
1	13	422	C
1	13	423	G
1	13	424	G
1	13	429	U
1	13	430	A
1	13	435	C
1	13	440	A
1	13	442	C
1	13	448	A
1	13	466	C
1	13	467	G
1	13	482	A
1	13	485	G
1	13	486	U
1	13	494	U
1	13	496	A

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Mol	Chain	Res	Type
1	13	497	U
1	13	509	A
1	13	510	A
1	13	511	C
1	13	512	U
1	13	513	C
1	13	518	C
1	13	521	G
1	13	528	C
1	13	531	U
1	13	532	A
1	13	533	A
1	13	534	U
1	13	545	C
1	13	547	A
1	13	550	G
1	13	559	A
1	13	560	U
1	13	561	U
1	13	562	C
1	13	569	C
1	13	572	A
1	13	573	A
1	13	576	G
1	13	577	G
1	13	595	G
1	13	596	C
1	13	614	A
1	13	618	C
1	13	623	C
1	13	630	G
1	13	632	A
1	13	633	G
1	13	653	A
1	13	665	A
1	13	671	G
1	13	683	G
1	13	687	A
1	13	688	G
1	13	700	G
1	13	701	C
1	13	702	A

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Mol	Chain	Res	Type
1	13	703	G
1	13	704	A
1	13	720	C
1	13	724	G
1	13	731	G
1	13	747	C
1	13	749	C
1	13	753	A
1	13	754	C
1	13	755	G
1	13	777	A
1	13	792	A
1	13	793	U
1	13	794	A
1	13	813	U
1	13	815	A
1	13	816	A
1	13	817	C
1	13	828	A
1	13	836	G
1	13	841	U
1	13	843	U
1	13	848	C
1	13	859	A
1	13	870	U
1	13	871	U
1	13	872	A
1	13	875	C
1	13	880	C
1	13	902	G
1	13	913	A
1	13	914	A
1	13	922	G
1	13	926	G
1	13	927	G
1	13	933	G
1	13	934	C
1	13	935	A
1	13	936	C
1	13	940	C
1	13	944	G
1	13	950	U

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Mol	Chain	Res	Type
1	13	958	A
1	13	960	U
1	13	961	U
1	13	966	M2G
1	13	968	A
1	13	969	A
1	13	971	G
1	13	972	C
1	13	974	A
1	13	975	A
1	13	976	G
1	13	977	A
1	13	980	C
1	13	981	U
1	13	982	U
1	13	983	A
1	13	991	U
1	13	992	U
1	13	993	G
1	13	994	A
1	13	1001	G
1	13	1002	G
1	13	1004	A
1	13	1005	A
1	13	1006	C
1	13	1008	C
1	13	1009	G
1	13	1017	G
1	13	1021	G
1	13	1024	G
1	13	1025	U
1	13	1028	C
1	13	1028(A)	C
1	13	1029	G
1	13	1032(A)	G
1	13	1037	C
1	13	1040	U
1	13	1054	C
1	13	1064	G
1	13	1065	U
1	13	1066	C
1	13	1082	G

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Mol	Chain	Res	Type
1	13	1090	U
1	13	1094	G
1	13	1095	U
1	13	1101	A
1	13	1113	C
1	13	1117	G
1	13	1123	A
1	13	1124	G
1	13	1125	U
1	13	1126	U
1	13	1127	G
1	13	1129	C
1	13	1130	A
1	13	1131	G
1	13	1133	G
1	13	1136	U
1	13	1137	C
1	13	1138	G
1	13	1139	G
1	13	1144	G
1	13	1146	A
1	13	1151	A
1	13	1157	A
1	13	1158	C
1	13	1159	U
1	13	1160	G
1	13	1171	G
1	13	1174	G
1	13	1178	G
1	13	1181	G
1	13	1182	G
1	13	1183	A
1	13	1187	G
1	13	1188	A
1	13	1191	A
1	13	1193	G
1	13	1196	U
1	13	1200	C
1	13	1201	A
1	13	1212	U
1	13	1213	A
1	13	1218	C

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Mol	Chain	Res	Type
1	13	1225	A
1	13	1226	C
1	13	1231	G
1	13	1236	A
1	13	1238	A
1	13	1240	U
1	13	1241	G
1	13	1256	A
1	13	1257	U
1	13	1258	G
1	13	1261	A
1	13	1263	C
1	13	1270	C
1	13	1272	G
1	13	1273	G
1	13	1275	A
1	13	1277	C
1	13	1279	A
1	13	1280	A
1	13	1281	U
1	13	1282	C
1	13	1286	A
1	13	1287	A
1	13	1288	A
1	13	1290	G
1	13	1292	U
1	13	1296	C
1	13	1300	G
1	13	1301	U
1	13	1302	U
1	13	1303	C
1	13	1305	G
1	13	1320	C
1	13	1321	C
1	13	1322	C
1	13	1323	G
1	13	1327	C
1	13	1331	G
1	13	1332	A
1	13	1333	A
1	13	1335	C
1	13	1336	C

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Mol	Chain	Res	Type
1	13	1337	G
1	13	1338	G
1	13	1347	G
1	13	1348	U
1	13	1353	G
1	13	1360	A
1	13	1362(A)	C
1	13	1364	U
1	13	1365	G
1	13	1370	G
1	13	1380	U
1	13	1398	A
1	13	1400	5MC
1	13	1402	4OC
1	13	1406	U
1	13	1419	G
1	13	1442	G
1	13	1443	G
1	13	1446	A
1	13	1447	G
1	13	1450	U
1	13	1452	C
1	13	1453	G
1	13	1454	G
1	13	1482	G
1	13	1487	G
1	13	1492	A
1	13	1494	G
1	13	1499	A
1	13	1503	A
1	13	1504	G
1	13	1505	G
1	13	1506	U
1	13	1507	A
1	13	1517	G
1	13	1529	G
1	13	1530	G
1	13	1531	A
1	13	1532	U
1	13	1533	C
1	13	1534	A
1	13	1535	C

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Mol	Chain	Res	Type
22	1K	6	G
22	1K	8	U
22	1K	9	G
22	1K	10	G
22	1K	16	C
22	1K	17	C
22	1K	20	G
22	1K	21	U
22	1K	22	A
22	1K	24	C
22	1K	37	U
22	1K	43	G
22	1K	48	U
22	1K	49	C
22	1K	50	G
22	1K	55	U
22	1K	57	C
22	1K	60	A
22	1K	61	U
22	1K	62	C
23	2K	7	G
23	2K	8	4SU
23	2K	13	C
23	2K	18	U
23	2K	21	H2U
23	2K	48	U
23	2K	49	C
23	2K	69	C
23	2K	77	A
22	3K	3	C
22	3K	5	G
22	3K	8	U
22	3K	9	G
22	3K	13	C
22	3K	14	A
22	3K	15	G
22	3K	17	C
22	3K	19	G
22	3K	21	U
22	3K	22	A
22	3K	23	G
22	3K	36	A

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Mol	Chain	Res	Type
22	3K	38	A
22	3K	45	A
22	3K	46	G
22	3K	49	C
22	3K	50	G
22	3K	56	U
22	3K	57	C
22	3K	58	A
22	3K	62	C
22	3K	73	A
24	4K	10	G
24	4K	11	U
24	4K	12	A
24	4K	13	A
24	4K	14	A
24	4K	19	A
24	4K	20	A
24	4K	23	A
24	4K	24	A
24	4K	25	A
25	1H	15	G
25	1H	26	G
25	1H	34	C
25	1H	35	G
25	1H	46	C
25	1H	49	A
25	1H	55	G
25	1H	61	G
25	1H	71	A
25	1H	72	U
25	1H	74	A
25	1H	75	G
25	1H	82	G
25	1H	83	G
25	1H	88	G
25	1H	99	U
25	1H	101	G
25	1H	102	G
25	1H	116	C
25	1H	118	A
25	1H	119	A
25	1H	120	U

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Mol	Chain	Res	Type
25	1H	125	G
25	1H	137	C
25	1H	140	A
25	1H	155	C
25	1H	161	U
25	1H	162	U
25	1H	163	U
25	1H	165	U
25	1H	177	G
25	1H	181	A
25	1H	188	G
25	1H	196	A
25	1H	197	A
25	1H	199	A
25	1H	206	U
25	1H	214	G
25	1H	215	G
25	1H	216	A
25	1H	221	A
25	1H	222	A
25	1H	223	A
25	1H	225	A
25	1H	227	A
25	1H	228	A
25	1H	229	A
25	1H	230	U
25	1H	232	G
25	1H	238	C
25	1H	239	U
25	1H	242	G
25	1H	248	G
25	1H	249	C
25	1H	252	G
25	1H	253	C
25	1H	265	A
25	1H	266	G
25	1H	269	U
25	1H	270(G)	C
25	1H	270(I)	G
25	1H	270(L)	U
25	1H	270(M)	U
25	1H	270(N)	G

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Mol	Chain	Res	Type
25	1H	270(P)	C
25	1H	270(Y)	G
25	1H	271(C)	U
25	1H	271	G
25	1H	273(C)	C
25	1H	274	G
25	1H	275	G
25	1H	276	A
25	1H	277	C
25	1H	278	A
25	1H	279	C
25	1H	284	U
25	1H	285	C
25	1H	299	A
25	1H	310	A
25	1H	317	G
25	1H	324	A
25	1H	329	G
25	1H	330	A
25	1H	332	A
25	1H	343	C
25	1H	352	G
25	1H	357	A
25	1H	363	G
25	1H	363(E)	U
25	1H	364	C
25	1H	371	A
25	1H	372	G
25	1H	373	U
25	1H	386	G
25	1H	388	G
25	1H	395	U
25	1H	396	G
25	1H	405	U
25	1H	411	G
25	1H	412	A
25	1H	421	U
25	1H	428	A
25	1H	444	C
25	1H	448	U
25	1H	457	A
25	1H	470	A

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Mol	Chain	Res	Type
25	1H	471	A
25	1H	479	A
25	1H	481	G
25	1H	482	A
25	1H	496	G
25	1H	503	A
25	1H	504	U
25	1H	505	A
25	1H	508	G
25	1H	509	C
25	1H	512	G
25	1H	513	A
25	1H	529	A
25	1H	530	G
25	1H	531	C
25	1H	532	A
25	1H	535	C
25	1H	537	C
25	1H	539	G
25	1H	540	G
25	1H	544	C
25	1H	545	G
25	1H	546	C
25	1H	547	A
25	1H	556	G
25	1H	563	G
25	1H	564	C
25	1H	568	U
25	1H	573	G
25	1H	574	C
25	1H	575	A
25	1H	588	U
25	1H	603	A
25	1H	607	U
25	1H	609	A
25	1H	614	U
25	1H	615	G
25	1H	616	A
25	1H	617	G
25	1H	620	G
25	1H	621	A
25	1H	622	G

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Mol	Chain	Res	Type
25	1H	624	C
25	1H	627	A
25	1H	629	G
25	1H	631	A
25	1H	634	C
25	1H	637	A
25	1H	638	G
25	1H	644	A
25	1H	645	C
25	1H	646	A
25	1H	654	A
25	1H	654(A)	A
25	1H	654(B)	C
25	1H	654(U)	A
25	1H	660	G
25	1H	667	U
25	1H	668	G
25	1H	669	G
25	1H	670	A
25	1H	671	C
25	1H	686	G
25	1H	714	U
25	1H	717	G
25	1H	730	C
25	1H	738	G
25	1H	747	U
25	1H	760	G
25	1H	762	U
25	1H	765	G
25	1H	776	G
25	1H	782	A
25	1H	784	A
25	1H	785	G
25	1H	792	G
25	1H	802	A
25	1H	805	G
25	1H	812	C
25	1H	819	A
25	1H	821	A
25	1H	827	U
25	1H	828	U
25	1H	845	G

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Mol	Chain	Res	Type
25	1H	846	C
25	1H	847	U
25	1H	856	C
25	1H	857	C
25	1H	859	G
25	1H	860	U
25	1H	866	A
25	1H	869	G
25	1H	871	U
25	1H	880	G
25	1H	882	G
25	1H	884	C
25	1H	885	C
25	1H	886	C
25	1H	889	C
25	1H	893	C
25	1H	898	C
25	1H	899	A
25	1H	900	A
25	1H	901	A
25	1H	904	C
25	1H	907	U
25	1H	910	A
25	1H	914	C
25	1H	915	C
25	1H	917	A
25	1H	918	A
25	1H	932	G
25	1H	940	G
25	1H	941	A
25	1H	946	G
25	1H	953	A
25	1H	959	A
25	1H	961	C
25	1H	968	G
25	1H	974	G
25	1H	974(A)	C
25	1H	980	A
25	1H	983	A
25	1H	990	A
25	1H	991	C
25	1H	996	A

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Mol	Chain	Res	Type
25	1H	999	U
25	1H	1002	G
25	1H	1003	G
25	1H	1005	C
25	1H	1010	A
25	1H	1011	G
25	1H	1012	U
25	1H	1013	C
25	1H	1015	G
25	1H	1016	G
25	1H	1023	U
25	1H	1025	G
25	1H	1026	U
25	1H	1027	A
25	1H	1032	A
25	1H	1033	U
25	1H	1045	A
25	1H	1046	A
25	1H	1050	A
25	1H	1054	A
25	1H	1055	G
25	1H	1057	A
25	1H	1058	U
25	1H	1060	U
25	1H	1061	U
25	1H	1062	G
25	1H	1065	U
25	1H	1067	A
25	1H	1068	G
25	1H	1070	A
25	1H	1071	G
25	1H	1076	C
25	1H	1077	A
25	1H	1078	U
25	1H	1079	C
25	1H	1080	A
25	1H	1082	U
25	1H	1083	U
25	1H	1084	A
25	1H	1085	A
25	1H	1086	A
25	1H	1087	G

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Mol	Chain	Res	Type
25	1H	1088	A
25	1H	1089	G
25	1H	1090	U
25	1H	1095	A
25	1H	1096	A
25	1H	1103	A
25	1H	1104	C
25	1H	1105	U
25	1H	1110	G
25	1H	1111	A
25	1H	1112	G
25	1H	1122	G
25	1H	1130	U
25	1H	1135	C
25	1H	1136	G
25	1H	1139	G
25	1H	1142	U
25	1H	1142(A)	A
25	1H	1151	G
25	1H	1155	A
25	1H	1156	A
25	1H	1165	U
25	1H	1169	G
25	1H	1173	G
25	1H	1174	A
25	1H	1175	U
25	1H	1176	G
25	1H	1178	C
25	1H	1179	C
25	1H	1180	C
25	1H	1195	G
25	1H	1204	A
25	1H	1205	U
25	1H	1210	A
25	1H	1211	U
25	1H	1218	C
25	1H	1220	A
25	1H	1237	A
25	1H	1251	C
25	1H	1256	G
25	1H	1267	U
25	1H	1269	A

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Mol	Chain	Res	Type
25	1H	1271	G
25	1H	1272	A
25	1H	1273	U
25	1H	1283	G
25	1H	1300	U
25	1H	1302	A
25	1H	1303	G
25	1H	1319	G
25	1H	1321	A
25	1H	1329	U
25	1H	1345	C
25	1H	1349	A
25	1H	1352	U
25	1H	1359	A
25	1H	1365	A
25	1H	1368	G
25	1H	1370	C
25	1H	1380	G
25	1H	1385	G
25	1H	1388	G
25	1H	1395	A
25	1H	1407	C
25	1H	1411	C
25	1H	1412	A
25	1H	1416	G
25	1H	1417	C
25	1H	1419	A
25	1H	1420	U
25	1H	1421	G
25	1H	1427	A
25	1H	1428	C
25	1H	1444(A)	A
25	1H	1445	C
25	1H	1449	A
25	1H	1449(A)	G
25	1H	1451	C
25	1H	1455	G
25	1H	1456	G
25	1H	1458	C
25	1H	1459	G
25	1H	1461	G
25	1H	1467	C

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Mol	Chain	Res	Type
25	1H	1471	A
25	1H	1478	G
25	1H	1479	G
25	1H	1482	U
25	1H	1483	G
25	1H	1486	A
25	1H	1493	C
25	1H	1494	A
25	1H	1495	A
25	1H	1497	U
25	1H	1504	C
25	1H	1509	C
25	1H	1510	A
25	1H	1522	G
25	1H	1528	A
25	1H	1540	G
25	1H	1543	A
25	1H	1545	A
25	1H	1546	C
25	1H	1558	A
25	1H	1559	G
25	1H	1560	G
25	1H	1566	A
25	1H	1569	A
25	1H	1578	U
25	1H	1579	A
25	1H	1581	G
25	1H	1585	C
25	1H	1586	A
25	1H	1593	G
25	1H	1598	C
25	1H	1608	A
25	1H	1617	C
25	1H	1618	A
25	1H	1622	G
25	1H	1639	U
25	1H	1640	C
25	1H	1647	G
25	1H	1648	C
25	1H	1674	G
25	1H	1678	G
25	1H	1681	G

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Mol	Chain	Res	Type
25	1H	1682	G
25	1H	1694	C
25	1H	1695	G
25	1H	1699	G
25	1H	1725	G
25	1H	1728	G
25	1H	1729	A
25	1H	1730	U
25	1H	1731	G
25	1H	1732	A
25	1H	1742	C
25	1H	1743	G
25	1H	1750	G
25	1H	1753	G
25	1H	1756	G
25	1H	1762	A
25	1H	1763	G
25	1H	1764	G
25	1H	1767	C
25	1H	1773	A
25	1H	1780	A
25	1H	1781	C
25	1H	1782	C
25	1H	1787	A
25	1H	1791	A
25	1H	1798	U
25	1H	1799	G
25	1H	1800	C
25	1H	1801	G
25	1H	1816	G
25	1H	1820	U
25	1H	1821	A
25	1H	1829	A
25	1H	1835	G
25	1H	1847	A
25	1H	1858	G
25	1H	1869	G
25	1H	1870	C
25	1H	1878	G
25	1H	1882	C
25	1H	1885	A
25	1H	1889	A

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Mol	Chain	Res	Type
25	1H	1906	G
25	1H	1908	C
25	1H	1910	G
25	1H	1913	A
25	1H	1919	A
25	1H	1929	G
25	1H	1930	G
25	1H	1935	G
25	1H	1936	A
25	1H	1937	A
25	1H	1938	A
25	1H	1955	U
25	1H	1963	U
25	1H	1964	G
25	1H	1967	C
25	1H	1969	A
25	1H	1970	A
25	1H	1971	A
25	1H	1972	A
25	1H	1981	A
25	1H	1982	C
25	1H	1991	U
25	1H	1992	G
25	1H	1993	U
25	1H	2023	G
25	1H	2031	A
25	1H	2033	A
25	1H	2043	C
25	1H	2049	G
25	1H	2055	C
25	1H	2056	G
25	1H	2059	A
25	1H	2060	A
25	1H	2061	G
25	1H	2062	A
25	1H	2063	C
25	1H	2069	G
25	1H	2086	U
25	1H	2092	U
25	1H	2093	G
25	1H	2099	U
25	1H	2108	C

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Mol	Chain	Res	Type
25	1H	2111	C
25	1H	2113	U
25	1H	2114	A
25	1H	2115	G
25	1H	2116	G
25	1H	2117	A
25	1H	2119	A
25	1H	2120	G
25	1H	2121	G
25	1H	2126	A
25	1H	2127	G
25	1H	2128	C
25	1H	2129	C
25	1H	2131	G
25	1H	2132	U
25	1H	2133	G
25	1H	2144	U
25	1H	2145	C
25	1H	2146	C
25	1H	2147	G
25	1H	2148	G
25	1H	2153	G
25	1H	2156	G
25	1H	2157	G
25	1H	2158	A
25	1H	2159	G
25	1H	2166	G
25	1H	2168	G
25	1H	2169	A
25	1H	2171	A
25	1H	2173	A
25	1H	2189	U
25	1H	2190	G
25	1H	2193	G
25	1H	2198	A
25	1H	2199	A
25	1H	2209	C
25	1H	2210	G
25	1H	2211	G
25	1H	2212	A
25	1H	2213	U
25	1H	2215	G

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Mol	Chain	Res	Type
25	1H	2225	A
25	1H	2238	G
25	1H	2248	C
25	1H	2266	A
25	1H	2269	A
25	1H	2275	C
25	1H	2280	G
25	1H	2283	C
25	1H	2286	A
25	1H	2287	A
25	1H	2294	C
25	1H	2298	A
25	1H	2302	G
25	1H	2305	A
25	1H	2307	G
25	1H	2308	G
25	1H	2310	A
25	1H	2311	A
25	1H	2319	G
25	1H	2320	A
25	1H	2321	G
25	1H	2325	G
25	1H	2334	G
25	1H	2335	A
25	1H	2336	A
25	1H	2342	C
25	1H	2345	G
25	1H	2346	A
25	1H	2347	C
25	1H	2350	C
25	1H	2352	A
25	1H	2382	G
25	1H	2383	G
25	1H	2385	C
25	1H	2392	A
25	1H	2393	A
25	1H	2396	G
25	1H	2399	G
25	1H	2402	C
25	1H	2403	C
25	1H	2406	U
25	1H	2410	G

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Mol	Chain	Res	Type
25	1H	2423	U
25	1H	2425	A
25	1H	2426	A
25	1H	2428	G
25	1H	2429	G
25	1H	2430	A
25	1H	2435	A
25	1H	2439	A
25	1H	2440	C
25	1H	2441	C
25	1H	2448	A
25	1H	2450	A
25	1H	2468	G
25	1H	2469	A
25	1H	2472	G
25	1H	2474	C
25	1H	2475	C
25	1H	2476	A
25	1H	2477	C
25	1H	2478	A
25	1H	2482	G
25	1H	2483	C
25	1H	2484	G
25	1H	2497	A
25	1H	2498	C
25	1H	2502	G
25	1H	2503	2MA
25	1H	2505	G
25	1H	2518	A
25	1H	2519	U
25	1H	2520	C
25	1H	2524	G
25	1H	2529	G
25	1H	2554	U
25	1H	2558	C
25	1H	2562	U
25	1H	2566	A
25	1H	2567	G
25	1H	2574	G
25	1H	2583	G
25	1H	2602	A
25	1H	2609	U

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Mol	Chain	Res	Type
25	1H	2611	U
25	1H	2612	C
25	1H	2621	A
25	1H	2630	G
25	1H	2641	G
25	1H	2649	U
25	1H	2654	A
25	1H	2661	G
25	1H	2665	A
25	1H	2673	G
25	1H	2682	U
25	1H	2689	U
25	1H	2690	C
25	1H	2691	C
25	1H	2697	G
25	1H	2702	U
25	1H	2703	C
25	1H	2707	G
25	1H	2712	U
25	1H	2712(A)	A
25	1H	2713	A
25	1H	2714	G
25	1H	2717	G
25	1H	2724	C
25	1H	2726	U
25	1H	2727	G
25	1H	2733	A
25	1H	2734	A
25	1H	2739	U
25	1H	2744	G
25	1H	2752	C
25	1H	2757	A
25	1H	2758	A
25	1H	2762	G
25	1H	2764	A
25	1H	2765	A
25	1H	2766	G
25	1H	2777	G
25	1H	2778	A
25	1H	2779	U
25	1H	2780	G
25	1H	2789	C

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Mol	Chain	Res	Type
25	1H	2790	A
25	1H	2791	C
25	1H	2792	G
25	1H	2793	G
25	1H	2802	G
25	1H	2803	C
25	1H	2807	G
25	1H	2808	U
25	1H	2820	A
25	1H	2821	A
25	1H	2823	A
25	1H	2833	G
25	1H	2835	A
25	1H	2861	G
25	1H	2864	G
25	1H	2867	G
25	1H	2872	G
25	1H	2874	C
25	1H	2879	C
25	1H	2880	C
25	1H	2891	G
25	1H	2892	A
25	1H	2894	G
25	1H	2895	U
26	16	2	C
26	16	8	U
26	16	10	C
26	16	12	C
26	16	13	A
26	16	15	A
26	16	16	G
26	16	19	G
26	16	21	G
26	16	22	U
26	16	24	G
26	16	25	A
26	16	27	C
26	16	33	G
26	16	38	C
26	16	40	U
26	16	41	U
26	16	42	C

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Mol	Chain	Res	Type
26	16	45	A
26	16	52	A
26	16	65	C
26	16	67	G
26	16	73	A
26	16	81	G
26	16	89	G
26	16	108	C
26	16	109	G
26	16	116	G
1	1G	4	U
1	1G	5	U
1	1G	6	G
1	1G	9	G
1	1G	13	U
1	1G	14	U
1	1G	15	G
1	1G	22	G
1	1G	32	A
1	1G	39	G
1	1G	47	C
1	1G	48	C
1	1G	51	A
1	1G	65	U
1	1G	66	G
1	1G	76	G
1	1G	78	G
1	1G	79	G
1	1G	80	G
1	1G	89	U
1	1G	90	C
1	1G	91	C
1	1G	95	G
1	1G	99	C
1	1G	108	G
1	1G	116	A
1	1G	121	C
1	1G	129(A)	G
1	1G	137	C
1	1G	138	G
1	1G	151	A
1	1G	163	C

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Mol	Chain	Res	Type
1	1G	182	U
1	1G	186	C
1	1G	187	C
1	1G	188	U
1	1G	189	U
1	1G	190	G
1	1G	191(A)	G
1	1G	191(D)	U
1	1G	195	A
1	1G	196	A
1	1G	208	U
1	1G	216	G
1	1G	217	C
1	1G	247	G
1	1G	251	G
1	1G	266	G
1	1G	267	C
1	1G	270	A
1	1G	279	A
1	1G	280	C
1	1G	281	G
1	1G	289	G
1	1G	298	A
1	1G	306	G
1	1G	314	C
1	1G	316	G
1	1G	321	A
1	1G	328	C
1	1G	329	A
1	1G	332	G
1	1G	341	C
1	1G	344	A
1	1G	345	C
1	1G	346	G
1	1G	347	G
1	1G	351	G
1	1G	352	C
1	1G	353	A
1	1G	354	G
1	1G	356	A
1	1G	367	U
1	1G	369	C

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Mol	Chain	Res	Type
1	1G	372	C
1	1G	373	A
1	1G	384	G
1	1G	388	G
1	1G	389	A
1	1G	397	A
1	1G	398	C
1	1G	406	G
1	1G	412	A
1	1G	413	G
1	1G	414	A
1	1G	419	C
1	1G	421	U
1	1G	422	C
1	1G	424	G
1	1G	429	U
1	1G	430	A
1	1G	435	C
1	1G	439	A
1	1G	440	A
1	1G	442	C
1	1G	448	A
1	1G	452	A
1	1G	465	A
1	1G	466	C
1	1G	467	G
1	1G	476	G
1	1G	478	A
1	1G	482	A
1	1G	484	G
1	1G	485	G
1	1G	486	U
1	1G	494	U
1	1G	495	A
1	1G	496	A
1	1G	497	U
1	1G	505	G
1	1G	510	A
1	1G	511	C
1	1G	518	C
1	1G	519	C
1	1G	521	G

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Mol	Chain	Res	Type
1	1G	527	7MG
1	1G	532	A
1	1G	533	A
1	1G	534	U
1	1G	535	A
1	1G	545	C
1	1G	547	A
1	1G	548	G
1	1G	559	A
1	1G	561	U
1	1G	568	G
1	1G	572	A
1	1G	573	A
1	1G	575	G
1	1G	576	G
1	1G	577	G
1	1G	581	G
1	1G	595	G
1	1G	596	C
1	1G	614	A
1	1G	618	C
1	1G	630	G
1	1G	652	U
1	1G	653	A
1	1G	656	C
1	1G	665	A
1	1G	688	G
1	1G	702	A
1	1G	703	G
1	1G	704	A
1	1G	721	G
1	1G	722	A
1	1G	724	G
1	1G	731	G
1	1G	749	C
1	1G	755	G
1	1G	770	C
1	1G	773	G
1	1G	777	A
1	1G	786	G
1	1G	792	A
1	1G	801	U

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Mol	Chain	Res	Type
1	1G	810	C
1	1G	813	U
1	1G	815	A
1	1G	816	A
1	1G	817	C
1	1G	818	G
1	1G	819	A
1	1G	820	U
1	1G	821	G
1	1G	828	A
1	1G	835	U
1	1G	838	G
1	1G	841	U
1	1G	842	C
1	1G	843	U
1	1G	848	C
1	1G	859	A
1	1G	870	U
1	1G	871	U
1	1G	873	A
1	1G	874	G
1	1G	884	U
1	1G	885	G
1	1G	889	A
1	1G	914	A
1	1G	915	A
1	1G	926	G
1	1G	927	G
1	1G	934	C
1	1G	935	A
1	1G	936	C
1	1G	942	G
1	1G	943	U
1	1G	947	G
1	1G	954	G
1	1G	960	U
1	1G	961	U
1	1G	966	M2G
1	1G	968	A
1	1G	969	A
1	1G	971	G
1	1G	972	C

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Mol	Chain	Res	Type
1	1G	974	A
1	1G	975	A
1	1G	976	G
1	1G	977	A
1	1G	978	A
1	1G	980	C
1	1G	989	C
1	1G	991	U
1	1G	992	U
1	1G	993	G
1	1G	999	U
1	1G	1000	A
1	1G	1001	G
1	1G	1003	G
1	1G	1005	A
1	1G	1007	C
1	1G	1008	C
1	1G	1009	G
1	1G	1014	A
1	1G	1020	U
1	1G	1024	G
1	1G	1025	U
1	1G	1026	G
1	1G	1027	C
1	1G	1028(A)	C
1	1G	1029	G
1	1G	1030	C
1	1G	1031	G
1	1G	1032	A
1	1G	1032(A)	G
1	1G	1032(B)	G
1	1G	1033	G
1	1G	1034	G
1	1G	1036	G
1	1G	1038	C
1	1G	1039	C
1	1G	1042	G
1	1G	1046	A
1	1G	1050	G
1	1G	1054	C
1	1G	1064	G
1	1G	1065	U

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Mol	Chain	Res	Type
1	1G	1066	C
1	1G	1067	A
1	1G	1068	G
1	1G	1081	G
1	1G	1082	G
1	1G	1084	G
1	1G	1086	U
1	1G	1087	G
1	1G	1094	G
1	1G	1095	U
1	1G	1096	C
1	1G	1099	G
1	1G	1100	C
1	1G	1101	A
1	1G	1109	C
1	1G	1113	C
1	1G	1117	G
1	1G	1118	C
1	1G	1124	G
1	1G	1125	U
1	1G	1127	G
1	1G	1130	A
1	1G	1131	G
1	1G	1134	G
1	1G	1135	U
1	1G	1136	U
1	1G	1137	C
1	1G	1138	G
1	1G	1139	G
1	1G	1140	C
1	1G	1141	C
1	1G	1144	G
1	1G	1145	C
1	1G	1146	A
1	1G	1147	C
1	1G	1149	C
1	1G	1151	A
1	1G	1157	A
1	1G	1158	C
1	1G	1159	U
1	1G	1160	G
1	1G	1161	C

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Mol	Chain	Res	Type
1	1G	1164	G
1	1G	1171	G
1	1G	1175	G
1	1G	1176	A
1	1G	1177	G
1	1G	1178	G
1	1G	1181	G
1	1G	1182	G
1	1G	1183	A
1	1G	1186	G
1	1G	1188	A
1	1G	1190	G
1	1G	1196	U
1	1G	1198	G
1	1G	1200	C
1	1G	1201	A
1	1G	1211	U
1	1G	1212	U
1	1G	1213	A
1	1G	1215	G
1	1G	1220	G
1	1G	1225	A
1	1G	1227	A
1	1G	1236	A
1	1G	1238	A
1	1G	1240	U
1	1G	1241	G
1	1G	1247	U
1	1G	1257	U
1	1G	1258	G
1	1G	1260	C
1	1G	1269	A
1	1G	1273	G
1	1G	1278	U
1	1G	1279	A
1	1G	1280	A
1	1G	1281	U
1	1G	1285	A
1	1G	1286	A
1	1G	1287	A
1	1G	1288	A
1	1G	1291	G

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Mol	Chain	Res	Type
1	1G	1294	G
1	1G	1296	C
1	1G	1297	C
1	1G	1299	A
1	1G	1300	G
1	1G	1302	U
1	1G	1303	C
1	1G	1305	G
1	1G	1306	A
1	1G	1317	C
1	1G	1320	C
1	1G	1321	C
1	1G	1322	C
1	1G	1323	G
1	1G	1325	C
1	1G	1331	G
1	1G	1334	G
1	1G	1335	C
1	1G	1336	C
1	1G	1337	G
1	1G	1344	C
1	1G	1346	A
1	1G	1347	G
1	1G	1348	U
1	1G	1358	U
1	1G	1362	C
1	1G	1362(A)	C
1	1G	1363	A
1	1G	1364	U
1	1G	1365	G
1	1G	1368	G
1	1G	1370	G
1	1G	1379	G
1	1G	1381	U
1	1G	1389	C
1	1G	1393	U
1	1G	1397	C
1	1G	1398	A
1	1G	1400	5MC
1	1G	1402	4OC
1	1G	1406	U
1	1G	1419	G

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Mol	Chain	Res	Type
1	1G	1442	G
1	1G	1443	G
1	1G	1446	A
1	1G	1447	G
1	1G	1450	U
1	1G	1452	C
1	1G	1453	G
1	1G	1454	G
1	1G	1475	G
1	1G	1492	A
1	1G	1498	UR3
1	1G	1499	A
1	1G	1502	A
1	1G	1503	A
1	1G	1504	G
1	1G	1506	U
1	1G	1507	A
1	1G	1508	G
1	1G	1510	U
1	1G	1517	G
1	1G	1519	MA6
1	1G	1520	G
1	1G	1529	G
1	1G	1530	G
1	1G	1531	A
1	1G	1532	U
1	1G	1533	C
1	1G	1534	A
1	1G	1536	C
1	1G	1540	U
57	2L	8	4SU
57	2L	19	G
57	2L	20	G
57	2L	21	U
57	2L	43	G
57	2L	48	U
57	2L	50	G
57	2L	53	G
57	2L	64	G
57	2L	70	C
57	2L	77	A
22	3L	2	G

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Mol	Chain	Res	Type
22	3L	5	G
22	3L	7	G
22	3L	8	U
22	3L	13	C
22	3L	15	G
22	3L	16	C
22	3L	17	C
22	3L	18	U
22	3L	19	G
22	3L	20	G
22	3L	21	U
22	3L	22	A
22	3L	23	G
22	3L	24	C
22	3L	35	C
22	3L	45	A
22	3L	48	U
22	3L	49	C
22	3L	54	G
22	3L	55	U
22	3L	58	A
22	3L	59	A
22	3L	60	A
22	3L	61	U
22	3L	64	G
22	3L	66	C
22	3L	68	C
22	3L	70	C
22	3L	72	C
22	3L	74	A
24	4L	7	G
24	4L	8	A
24	4L	10	G
24	4L	13	A
24	4L	14	A
24	4L	19	A
24	4L	23	A
24	4L	25	A
25	14	5	A
25	14	6	A
25	14	8	A
25	14	15	G

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Mol	Chain	Res	Type
25	14	34	C
25	14	35	G
25	14	36	G
25	14	46	C
25	14	50	U
25	14	54	G
25	14	55	G
25	14	58	G
25	14	60	G
25	14	63	U
25	14	68	G
25	14	69	C
25	14	71	A
25	14	72	U
25	14	74	A
25	14	75	G
25	14	82	G
25	14	83	G
25	14	85	G
25	14	90	U
25	14	91	A
25	14	96	G
25	14	99	U
25	14	101	G
25	14	103	A
25	14	118	A
25	14	120	U
25	14	124	G
25	14	129	C
25	14	131	G
25	14	137	C
25	14	138	G
25	14	146	G
25	14	153	C
25	14	155	C
25	14	161	U
25	14	172	C
25	14	173	G
25	14	174	C
25	14	175	G
25	14	196	A
25	14	199	A

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Mol	Chain	Res	Type
25	14	201	C
25	14	214	G
25	14	215	G
25	14	216	A
25	14	221	A
25	14	222	A
25	14	225	A
25	14	226	G
25	14	228	A
25	14	229	A
25	14	248	G
25	14	249	C
25	14	252	G
25	14	265	A
25	14	266	G
25	14	269	U
25	14	270(B)	A
25	14	270(C)	C
25	14	270(G)	C
25	14	270(J)	G
25	14	270(K)	C
25	14	270(L)	U
25	14	270(M)	U
25	14	270(O)	U
25	14	270(Q)	C
25	14	270(U)	C
25	14	270(Z)	U
25	14	271(C)	U
25	14	271	G
25	14	274	G
25	14	275	G
25	14	276	A
25	14	277	C
25	14	278	A
25	14	279	C
25	14	281	G
25	14	286	C
25	14	289	A
25	14	311	A
25	14	324	A
25	14	329	G
25	14	330	A

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Mol	Chain	Res	Type
25	14	332	A
25	14	333	G
25	14	344	G
25	14	345	A
25	14	352	G
25	14	354	G
25	14	356	G
25	14	363(E)	U
25	14	371	A
25	14	372	G
25	14	386	G
25	14	387	U
25	14	388	G
25	14	395	U
25	14	396	G
25	14	399	G
25	14	405	U
25	14	406	G
25	14	407	G
25	14	411	G
25	14	412	A
25	14	416	C
25	14	422	A
25	14	428	A
25	14	444	C
25	14	447	A
25	14	448	U
25	14	449	A
25	14	453	C
25	14	455	C
25	14	456	C
25	14	457	A
25	14	470	A
25	14	481	G
25	14	486	C
25	14	494	G
25	14	504	U
25	14	505	A
25	14	507	A
25	14	508	G
25	14	509	C
25	14	510	C

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Mol	Chain	Res	Type
25	14	512	G
25	14	513	A
25	14	527	C
25	14	529	A
25	14	531	C
25	14	532	A
25	14	533	G
25	14	548	A
25	14	549	G
25	14	554	U
25	14	561	G
25	14	563	G
25	14	571	A
25	14	573	G
25	14	574	C
25	14	575	A
25	14	588	U
25	14	603	A
25	14	604	G
25	14	607	U
25	14	614	U
25	14	615	G
25	14	617	G
25	14	620	G
25	14	621	A
25	14	622	G
25	14	624	C
25	14	626	U
25	14	627	A
25	14	631	A
25	14	637	A
25	14	645	C
25	14	646	A
25	14	647	G
25	14	650	C
25	14	651	G
25	14	654(U)	A
25	14	677	A
25	14	686	G
25	14	694	U
25	14	699	A
25	14	708	C

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Mol	Chain	Res	Type
25	14	717	G
25	14	721	C
25	14	722	A
25	14	730	C
25	14	735	A
25	14	749	C
25	14	752	A
25	14	753	C
25	14	762	U
25	14	776	G
25	14	777	A
25	14	779	U
25	14	782	A
25	14	784	A
25	14	785	G
25	14	790	C
25	14	791	C
25	14	792	G
25	14	793	A
25	14	805	G
25	14	812	C
25	14	819	A
25	14	827	U
25	14	828	U
25	14	832	G
25	14	846	C
25	14	847	U
25	14	854	G
25	14	856	C
25	14	857	C
25	14	859	G
25	14	860	U
25	14	865	C
25	14	866	A
25	14	869	G
25	14	875	G
25	14	878	A
25	14	879	G
25	14	881	G
25	14	882	G
25	14	883	G
25	14	884	C

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Mol	Chain	Res	Type
25	14	885	C
25	14	886	C
25	14	887	A
25	14	888	C
25	14	889	C
25	14	890	A
25	14	894	C
25	14	895	U
25	14	897	C
25	14	901	A
25	14	905	U
25	14	906	G
25	14	907	U
25	14	910	A
25	14	914	C
25	14	915	C
25	14	917	A
25	14	919	G
25	14	928	G
25	14	932	G
25	14	938	G
25	14	941	A
25	14	943	U
25	14	944	G
25	14	946	G
25	14	958	U
25	14	959	A
25	14	961	C
25	14	968	G
25	14	973	A
25	14	974	G
25	14	974(A)	C
25	14	980	A
25	14	983	A
25	14	990	A
25	14	996	A
25	14	997	G
25	14	1004	C
25	14	1009	A
25	14	1010	A
25	14	1011	G
25	14	1012	U

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Mol	Chain	Res	Type
25	14	1013	C
25	14	1022	G
25	14	1023	U
25	14	1024	G
25	14	1025	G
25	14	1026	U
25	14	1027	A
25	14	1031	G
25	14	1033	U
25	14	1034	G
25	14	1044	G
25	14	1045	A
25	14	1046	A
25	14	1047	G
25	14	1050	A
25	14	1051	G
25	14	1053	C
25	14	1056	G
25	14	1057	A
25	14	1060	U
25	14	1061	U
25	14	1062	G
25	14	1065	U
25	14	1067	A
25	14	1070	A
25	14	1071	G
25	14	1074	G
25	14	1075	C
25	14	1076	C
25	14	1077	A
25	14	1080	A
25	14	1083	U
25	14	1084	A
25	14	1086	A
25	14	1087	G
25	14	1088	A
25	14	1089	G
25	14	1090	U
25	14	1092	C
25	14	1095	A
25	14	1096	A
25	14	1099	G

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Mol	Chain	Res	Type
25	14	1100	C
25	14	1101	U
25	14	1105	U
25	14	1108	U
25	14	1111	A
25	14	1112	G
25	14	1113	U
25	14	1115	G
25	14	1122	G
25	14	1126	A
25	14	1128	A
25	14	1129	A
25	14	1130	U
25	14	1135	C
25	14	1136	G
25	14	1139	G
25	14	1141	U
25	14	1142(A)	A
25	14	1143	A
25	14	1148	A
25	14	1150	C
25	14	1155	A
25	14	1160	G
25	14	1170	G
25	14	1171	G
25	14	1173	G
25	14	1174	A
25	14	1175	U
25	14	1176	G
25	14	1177	A
25	14	1178	C
25	14	1180	C
25	14	1183	G
25	14	1195	G
25	14	1196	C
25	14	1205	U
25	14	1206	G
25	14	1210	A
25	14	1211	U
25	14	1212	G
25	14	1219	G
25	14	1220	A

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Mol	Chain	Res	Type
25	14	1222	C
25	14	1224	G
25	14	1230	C
25	14	1233	C
25	14	1244	G
25	14	1247	A
25	14	1249	U
25	14	1250	G
25	14	1253	A
25	14	1255	U
25	14	1256	G
25	14	1271	G
25	14	1272	A
25	14	1281	G
25	14	1300	U
25	14	1301	A
25	14	1312	U
25	14	1313	U
25	14	1314	C
25	14	1319	G
25	14	1321	A
25	14	1329	U
25	14	1332	G
25	14	1337	G
25	14	1345	C
25	14	1349	A
25	14	1352	U
25	14	1359	A
25	14	1360	A
25	14	1365	A
25	14	1368	G
25	14	1379	A
25	14	1380	G
25	14	1383	C
25	14	1384	A
25	14	1385	G
25	14	1386	C
25	14	1390	U
25	14	1391	U
25	14	1395	A
25	14	1406	U
25	14	1416	G

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Mol	Chain	Res	Type
25	14	1420	U
25	14	1421	G
25	14	1428	C
25	14	1429	G
25	14	1436	G
25	14	1437	C
25	14	1444	G
25	14	1444(A)	A
25	14	1445	C
25	14	1449	A
25	14	1450	C
25	14	1455	G
25	14	1458	C
25	14	1459	G
25	14	1460	A
25	14	1461	G
25	14	1467	C
25	14	1471	A
25	14	1475	G
25	14	1478	G
25	14	1482	U
25	14	1483	G
25	14	1486	A
25	14	1488	G
25	14	1490	A
25	14	1493	C
25	14	1494	A
25	14	1502	C
25	14	1505	C
25	14	1508	A
25	14	1509	C
25	14	1510	A
25	14	1514	U
25	14	1515	C
25	14	1521	G
25	14	1522	G
25	14	1527	G
25	14	1534	G
25	14	1535	U
25	14	1536	A
25	14	1537	C
25	14	1540	G

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Mol	Chain	Res	Type
25	14	1543	A
25	14	1544	C
25	14	1545	A
25	14	1547	C
25	14	1549	C
25	14	1558	A
25	14	1559	G
25	14	1560	G
25	14	1566	A
25	14	1567	A
25	14	1569	A
25	14	1578	U
25	14	1579	A
25	14	1580	A
25	14	1581	G
25	14	1585	C
25	14	1586	A
25	14	1588	C
25	14	1591	G
25	14	1592	C
25	14	1593	G
25	14	1595	G
25	14	1598	C
25	14	1608	A
25	14	1610	A
25	14	1618	A
25	14	1634	A
25	14	1635	G
25	14	1639	U
25	14	1640	C
25	14	1647	G
25	14	1648	C
25	14	1653	G
25	14	1654	A
25	14	1665	A
25	14	1667	G
25	14	1668	A
25	14	1674	G
25	14	1688	U
25	14	1694	C
25	14	1695	G
25	14	1696	G

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Mol	Chain	Res	Type
25	14	1699	G
25	14	1700	A
25	14	1701	A
25	14	1725	G
25	14	1728	G
25	14	1729	A
25	14	1730	U
25	14	1731	G
25	14	1733	G
25	14	1735	C
25	14	1743	G
25	14	1750	G
25	14	1753	G
25	14	1754	C
25	14	1756	G
25	14	1759	A
25	14	1762	A
25	14	1763	G
25	14	1764	G
25	14	1773	A
25	14	1780	A
25	14	1782	C
25	14	1783	A
25	14	1787	A
25	14	1791	A
25	14	1800	C
25	14	1801	G
25	14	1815	A
25	14	1816	G
25	14	1820	U
25	14	1823	G
25	14	1829	A
25	14	1835	G
25	14	1847	A
25	14	1858	G
25	14	1860	G
25	14	1869	G
25	14	1878	G
25	14	1881	C
25	14	1882	C
25	14	1888	G
25	14	1889	A

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Mol	Chain	Res	Type
25	14	1900	A
25	14	1903	G
25	14	1905	C
25	14	1906	G
25	14	1909	C
25	14	1910	G
25	14	1913	A
25	14	1914	C
25	14	1915	5MU
25	14	1919	A
25	14	1920	OMC
25	14	1929	G
25	14	1930	G
25	14	1937	A
25	14	1938	A
25	14	1948	G
25	14	1949	G
25	14	1955	U
25	14	1956	U
25	14	1963	U
25	14	1966	A
25	14	1967	C
25	14	1970	A
25	14	1971	A
25	14	1972	A
25	14	1982	C
25	14	1993	U
25	14	2019	A
25	14	2022	U
25	14	2023	G
25	14	2026	C
25	14	2027	G
25	14	2031	A
25	14	2033	A
25	14	2039	C
25	14	2043	C
25	14	2050	C
25	14	2052	G
25	14	2055	C
25	14	2056	G
25	14	2059	A
25	14	2060	A

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Mol	Chain	Res	Type
25	14	2061	G
25	14	2062	A
25	14	2063	C
25	14	2069	G
25	14	2071	A
25	14	2092	U
25	14	2099	U
25	14	2100	G
25	14	2101	G
25	14	2102	U
25	14	2107	C
25	14	2111	C
25	14	2112	G
25	14	2113	U
25	14	2114	A
25	14	2115	G
25	14	2117	A
25	14	2119	A
25	14	2120	G
25	14	2128	C
25	14	2130	U
25	14	2131	G
25	14	2132	U
25	14	2136	C
25	14	2137	C
25	14	2139	C
25	14	2140	C
25	14	2144	U
25	14	2145	C
25	14	2146	C
25	14	2147	G
25	14	2148	G
25	14	2154	G
25	14	2161	C
25	14	2167	U
25	14	2169	A
25	14	2171	A
25	14	2172	U
25	14	2173	A
25	14	2174	C
25	14	2175	C
25	14	2182	G

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Mol	Chain	Res	Type
25	14	2189	U
25	14	2190	G
25	14	2191	G
25	14	2192	G
25	14	2198	A
25	14	2199	A
25	14	2210	G
25	14	2211	G
25	14	2212	A
25	14	2213	U
25	14	2215	G
25	14	2225	A
25	14	2226	C
25	14	2238	G
25	14	2239	G
25	14	2261	C
25	14	2275	C
25	14	2276	G
25	14	2278	A
25	14	2280	G
25	14	2283	C
25	14	2286	A
25	14	2287	A
25	14	2288	A
25	14	2289	G
25	14	2290	G
25	14	2304	G
25	14	2305	A
25	14	2306	C
25	14	2307	G
25	14	2308	G
25	14	2310	A
25	14	2314	C
25	14	2316	C
25	14	2319	G
25	14	2320	A
25	14	2321	G
25	14	2322	A
25	14	2325	G
25	14	2334	G
25	14	2342	C
25	14	2345	G

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Mol	Chain	Res	Type
25	14	2346	A
25	14	2347	C
25	14	2350	C
25	14	2372	G
25	14	2382	G
25	14	2383	G
25	14	2384	G
25	14	2385	C
25	14	2388	A
25	14	2396	G
25	14	2402	C
25	14	2403	C
25	14	2406	U
25	14	2409	G
25	14	2425	A
25	14	2429	G
25	14	2430	A
25	14	2435	A
25	14	2439	A
25	14	2440	C
25	14	2441	C
25	14	2445	G
25	14	2446	G
25	14	2448	A
25	14	2450	A
25	14	2469	A
25	14	2470	G
25	14	2472	G
25	14	2473	U
25	14	2476	A
25	14	2478	A
25	14	2479	G
25	14	2482	G
25	14	2497	A
25	14	2502	G
25	14	2504	U
25	14	2505	G
25	14	2517	C
25	14	2518	A
25	14	2519	U
25	14	2520	C
25	14	2527	C

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Mol	Chain	Res	Type
25	14	2529	G
25	14	2542	A
25	14	2543	G
25	14	2546	U
25	14	2554	U
25	14	2556	C
25	14	2566	A
25	14	2567	G
25	14	2576	G
25	14	2582	G
25	14	2586	C
25	14	2602	A
25	14	2603	G
25	14	2609	U
25	14	2610	C
25	14	2611	U
25	14	2612	C
25	14	2615	U
25	14	2629	A
25	14	2630	G
25	14	2638	G
25	14	2655	G
25	14	2656	U
25	14	2665	A
25	14	2673	G
25	14	2674	G
25	14	2675	A
25	14	2683	C
25	14	2689	U
25	14	2691	C
25	14	2702	U
25	14	2703	C
25	14	2712	U
25	14	2712(A)	A
25	14	2713	A
25	14	2718	G
25	14	2719	G
25	14	2725	A
25	14	2733	A
25	14	2734	A
25	14	2739	U
25	14	2748	A

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Mol	Chain	Res	Type
25	14	2749	A
25	14	2750	A
25	14	2751	G
25	14	2752	C
25	14	2754	U
25	14	2756	U
25	14	2757	A
25	14	2758	A
25	14	2762	G
25	14	2763	G
25	14	2764	A
25	14	2766	G
25	14	2769	C
25	14	2777	G
25	14	2778	A
25	14	2779	U
25	14	2780	G
25	14	2790	A
25	14	2791	C
25	14	2792	G
25	14	2794	C
25	14	2802	G
25	14	2808	U
25	14	2818	G
25	14	2820	A
25	14	2821	A
25	14	2827	C
25	14	2833	G
25	14	2834	G
25	14	2860	A
25	14	2861	G
25	14	2867	G
25	14	2868	A
25	14	2872	G
25	14	2876	G
25	14	2879	C
25	14	2882	A
25	14	2883	A
25	14	2893	G
25	14	2895	U
25	14	2896	C
26	1J	0	A

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Mol	Chain	Res	Type
26	1J	3	C
26	1J	5	C
26	1J	6	C
26	1J	8	U
26	1J	13	A
26	1J	14	U
26	1J	15	A
26	1J	16	G
26	1J	19	G
26	1J	22	U
26	1J	24	G
26	1J	25	A
26	1J	27	C
26	1J	30	C
26	1J	31	C
26	1J	32	C
26	1J	34	U
26	1J	35	U
26	1J	40	U
26	1J	41	U
26	1J	42	C
26	1J	44	G
26	1J	45	A
26	1J	47	C
26	1J	52	A
26	1J	53	A
26	1J	63	G
26	1J	66	A
26	1J	67	G
26	1J	73	A
26	1J	81	G
26	1J	88	C
26	1J	89	G
26	1J	96	G
26	1J	108	C
26	1J	109	G
26	1J	112	G
26	1J	115	G
26	1J	119	A

All (226) RNA pucker outliers are listed below:

Mol	Chain	Res	Type
1	13	5	U
1	13	60	A
1	13	89	U
1	13	190	G
1	13	243	A
1	13	244	U
1	13	266	G
1	13	328	C
1	13	352	C
1	13	410	G
1	13	412	A
1	13	423	G
1	13	429	U
1	13	481	G
1	13	484	G
1	13	485	G
1	13	495	A
1	13	509	A
1	13	560	U
1	13	595	G
1	13	686	U
1	13	687	A
1	13	701	C
1	13	703	G
1	13	748	C
1	13	753	A
1	13	792	A
1	13	812	C
1	13	815	A
1	13	871	U
1	13	913	A
1	13	960	U
1	13	965	A
1	13	991	U
1	13	992	U
1	13	1065	U
1	13	1126	U
1	13	1182	G
1	13	1200	C
1	13	1280	A
1	13	1285	A
1	13	1302	U
1	13	1336	C

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Mol	Chain	Res	Type
1	13	1347	G
1	13	1397	C
1	13	1503	A
1	13	1504	G
22	1K	20	G
22	1K	49	C
23	2K	48	U
24	4K	10	G
24	4K	18	G
25	1H	99	U
25	1H	221	A
25	1H	222	A
25	1H	229	A
25	1H	249	C
25	1H	271(B)	G
25	1H	275	G
25	1H	278	A
25	1H	372	G
25	1H	404	C
25	1H	481	G
25	1H	503	A
25	1H	512	G
25	1H	587	C
25	1H	614	U
25	1H	637	A
25	1H	653	A
25	1H	746	A
25	1H	845	G
25	1H	856	C
25	1H	859	G
25	1H	1012	U
25	1H	1022	G
25	1H	1026	U
25	1H	1045	A
25	1H	1078	U
25	1H	1085	A
25	1H	1109	C
25	1H	1177	A
25	1H	1178	C
25	1H	1204	A
25	1H	1210	A
25	1H	1266	G

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Mol	Chain	Res	Type
25	1H	1301	A
25	1H	1302	A
25	1H	1379	A
25	1H	1444(A)	A
25	1H	1558	A
25	1H	1608	A
25	1H	1681	G
25	1H	1694	C
25	1H	1698	A
25	1H	1799	G
25	1H	1819	A
25	1H	1980	G
25	1H	1992	G
25	1H	2022	U
25	1H	2126	A
25	1H	2146	C
25	1H	2157	G
25	1H	2351	G
25	1H	2405	G
25	1H	2439	A
25	1H	2481	G
25	1H	2518	A
25	1H	2566	A
25	1H	2610	C
25	1H	2689	U
25	1H	2712	U
25	1H	2751	G
25	1H	2756	U
25	1H	2776	A
25	1H	2832	U
26	16	24	G
26	16	66	A
1	1G	5	U
1	1G	31	G
1	1G	115	G
1	1G	181	G
1	1G	201	C
1	1G	250	A
1	1G	266	G
1	1G	279	A
1	1G	328	C
1	1G	353	A

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Mol	Chain	Res	Type
1	1G	412	A
1	1G	413	G
1	1G	429	U
1	1G	485	G
1	1G	496	A
1	1G	518	C
1	1G	547	A
1	1G	560	U
1	1G	595	G
1	1G	687	A
1	1G	701	C
1	1G	748	C
1	1G	812	C
1	1G	815	A
1	1G	818	G
1	1G	870	U
1	1G	872	A
1	1G	913	A
1	1G	960	U
1	1G	976	G
1	1G	991	U
1	1G	992	U
1	1G	1000	A
1	1G	1004	A
1	1G	1008	C
1	1G	1029	G
1	1G	1030	C
1	1G	1033	G
1	1G	1038	C
1	1G	1049	U
1	1G	1053	G
1	1G	1065	U
1	1G	1067	A
1	1G	1137	C
1	1G	1139	G
1	1G	1157	A
1	1G	1200	C
1	1G	1211	U
1	1G	1240	U
1	1G	1346	A
1	1G	1347	G
1	1G	1452	C

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Mol	Chain	Res	Type
1	1G	1532	U
22	3L	23	G
22	3L	58	A
24	4L	7	G
25	14	34	C
25	14	49	A
25	14	71	A
25	14	119	A
25	14	221	A
25	14	275	G
25	14	387	U
25	14	587	C
25	14	616	A
25	14	752	A
25	14	790	C
25	14	846	C
25	14	856	C
25	14	877	U
25	14	878	A
25	14	883	G
25	14	1033	U
25	14	1085	A
25	14	1171	G
25	14	1210	A
25	14	1300	U
25	14	1359	A
25	14	1427	A
25	14	1460	A
25	14	1558	A
25	14	1634	A
25	14	1647	G
25	14	1652	A
25	14	1694	C
25	14	1698	A
25	14	1762	A
25	14	1782	C
25	14	1819	A
25	14	1936	A
25	14	1955	U
25	14	1992	G
25	14	2144	U
25	14	2166	G

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Mol	Chain	Res	Type
25	14	2191	G
25	14	2211	G
25	14	2439	A
25	14	2481	G
25	14	2518	A
25	14	2585	U
25	14	2602	A
25	14	2610	C
25	14	2776	A
25	14	2789	C
25	14	2859	G
25	14	2867	G
26	1J	24	G
26	1J	34	U
26	1J	66	A

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

59 non-standard protein/DNA/RNA residues are modelled in this entry.

In the following table, the Counts columns list the number of bonds (or angles) for which Mogul statistics could be retrieved, the number of bonds (or angles) that are observed in the model and the number of bonds (or angles) that are defined in the Chemical Component Dictionary. The Link column lists molecule types, if any, to which the group is linked. The Z score for a bond length (or angle) is the number of standard deviations the observed value is removed from the expected value. A bond length (or angle) with $|Z| > 2$ is considered an outlier worth inspection. RMSZ is the root-mean-square of all Z scores of the bond lengths (or angles).

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
25	OMC	1H	1920	25	19,22,23	1.76	4 (21%)	25,31,34	1.07	4 (16%)
1	M2G	1G	966	1	24,27,28	2.90	8 (33%)	33,40,43	1.71	7 (21%)
25	PSU	1H	2605	25	18,21,22	1.17	2 (11%)	21,30,33	2.13	5 (23%)
25	OMG	14	2251	57,25,58	23,26,27	5.12	11 (47%)	32,38,41	4.73	18 (56%)
25	2MA	1H	2503	59,25,58	22,25,26	3.57	7 (31%)	32,37,40	2.35	9 (28%)
23	H2U	2K	21	23	18,21,22	2.17	4 (22%)	19,30,33	1.64	3 (15%)
57	OMC	2L	33	57	19,22,23	1.82	4 (21%)	25,31,34	1.09	3 (12%)
25	5MC	1H	1942	25	19,22,23	3.83	8 (42%)	26,32,35	1.04	2 (7%)
57	5MU	2L	55	57	19,22,23	3.98	5 (26%)	27,32,35	3.24	10 (37%)
25	OMU	1H	2552	59,25	19,22,23	2.96	8 (42%)	25,31,34	1.88	5 (20%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
1	MA6	1G	1519	1	23,26,27	1.43	5 (21%)	33,38,41	2.71	12 (36%)
25	PSU	1H	1917	25	18,21,22	1.19	1 (5%)	21,30,33	1.97	4 (19%)
1	4OC	13	1402	1	20,23,24	3.12	8 (40%)	25,32,35	0.87	1 (4%)
1	5MC	13	1407	1	19,22,23	4.09	8 (42%)	26,32,35	1.04	1 (3%)
1	PSU	13	516	59,1	18,21,22	1.19	1 (5%)	21,30,33	1.75	4 (19%)
25	5MU	1H	1939	25,58	19,22,23	3.81	5 (26%)	27,32,35	3.70	8 (29%)
1	5MC	1G	1407	1	19,22,23	4.09	8 (42%)	26,32,35	1.10	2 (7%)
1	PSU	1G	516	59,1	18,21,22	1.19	1 (5%)	21,30,33	1.89	4 (19%)
23	PSU	2K	56	23	18,21,22	1.17	1 (5%)	21,30,33	1.95	5 (23%)
25	5MU	14	1939	25,58	19,22,23	3.78	5 (26%)	27,32,35	3.42	9 (33%)
25	5MU	14	1915	25	19,22,23	4.05	5 (26%)	27,32,35	3.28	9 (33%)
1	MA6	13	1518	1	23,26,27	1.49	5 (21%)	33,38,41	2.71	12 (36%)
25	5MC	14	1962	25	19,22,23	4.13	8 (42%)	26,32,35	1.41	5 (19%)
25	PSU	14	2605	25	18,21,22	1.18	1 (5%)	21,30,33	1.82	3 (14%)
12	0TD	3A	89	12	8,9,10	1.43	0	6,11,13	3.19	3 (50%)
1	4OC	1G	1402	1	20,23,24	2.97	8 (40%)	25,32,35	1.15	1 (4%)
1	MA6	1G	1518	1	23,26,27	1.50	5 (21%)	33,38,41	2.61	11 (33%)
25	PSU	14	1911	25	18,21,22	1.14	1 (5%)	21,30,33	1.73	4 (19%)
57	4SU	2L	8	57	18,21,22	1.85	3 (16%)	25,30,33	1.98	4 (16%)
1	5MC	13	1400	1	19,22,23	4.28	8 (42%)	26,32,35	1.08	2 (7%)
23	7MG	2K	47	23	23,26,27	3.14	8 (34%)	27,39,42	2.77	9 (33%)
25	OMC	14	1920	25	19,22,23	1.78	4 (21%)	25,31,34	0.84	1 (4%)
1	7MG	13	527	58,1	23,26,27	3.12	8 (34%)	27,39,42	2.83	9 (33%)
57	PSU	2L	56	57	18,21,22	1.21	1 (5%)	21,30,33	1.92	4 (19%)
1	7MG	1G	527	58,1	23,26,27	3.10	8 (34%)	27,39,42	2.83	9 (33%)
25	2MA	14	2503	59,25,58	22,25,26	3.68	8 (36%)	32,37,40	2.26	11 (34%)
25	OMG	1H	2251	25,58,23	23,26,27	5.07	11 (47%)	32,38,41	4.75	16 (50%)
1	5MC	1G	1400	1	19,22,23	4.14	8 (42%)	26,32,35	1.08	2 (7%)
1	UR3	13	1498	1	19,22,23	2.84	7 (36%)	26,32,35	1.72	4 (15%)
57	7MG	2L	47	57	23,26,27	3.09	8 (34%)	27,39,42	2.81	9 (33%)
1	UR3	1G	1498	1	19,22,23	2.84	7 (36%)	26,32,35	1.83	4 (15%)
1	5MC	13	1404	1	19,22,23	4.12	8 (42%)	26,32,35	1.12	2 (7%)
1	5MC	1G	1404	1	19,22,23	4.17	8 (42%)	26,32,35	1.00	2 (7%)
25	OMU	14	2552	59,25	19,22,23	3.05	8 (42%)	25,31,34	2.02	5 (20%)
25	5MC	14	1942	25	19,22,23	4.09	8 (42%)	26,32,35	1.07	2 (7%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
23	5MU	2K	55	23	19,22,23	3.99	5 (26%)	27,32,35	3.19	8 (29%)
23	OMC	2K	33	23	19,22,23	1.80	3 (15%)	25,31,34	1.03	2 (8%)
25	5MU	1H	1915	25	19,22,23	4.02	5 (26%)	27,32,35	3.30	10 (37%)
12	0TD	3I	89	12	8,9,10	1.28	0	6,11,13	2.92	3 (50%)
1	2MG	1G	1207	1	23,26,27	3.13	8 (34%)	33,38,41	2.17	8 (24%)
1	MA6	13	1519	1	23,26,27	1.49	5 (21%)	33,38,41	2.66	11 (33%)
23	4SU	2K	8	23	18,21,22	1.87	3 (16%)	25,30,33	2.45	5 (20%)
25	PSU	14	1917	25	18,21,22	1.10	1 (5%)	21,30,33	1.95	5 (23%)
1	2MG	13	1207	1	23,26,27	3.04	8 (34%)	33,38,41	2.29	13 (39%)
25	5MC	1H	1962	25,58	19,22,23	4.14	8 (42%)	26,32,35	1.06	1 (3%)
1	5MC	1G	967	1	19,22,23	4.17	8 (42%)	26,32,35	1.04	2 (7%)
1	M2G	13	966	1	24,27,28	2.99	8 (33%)	33,40,43	1.68	6 (18%)
25	PSU	1H	1911	25	18,21,22	1.14	1 (5%)	21,30,33	1.84	4 (19%)
1	5MC	13	967	1	19,22,23	4.13	8 (42%)	26,32,35	1.02	2 (7%)

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
25	OMC	1H	1920	25	-	2/9/27/28	0/2/2/2
1	M2G	1G	966	1	-	0/11/29/30	0/3/3/3
25	PSU	1H	2605	25	-	2/7/25/26	0/2/2/2
25	OMG	14	2251	57,25,58	-	0/9/27/28	0/3/3/3
25	2MA	1H	2503	59,25,58	-	3/7/25/26	0/3/3/3
23	H2U	2K	21	23	-	4/7/38/39	0/2/2/2
57	OMC	2L	33	57	-	3/9/27/28	0/2/2/2
25	5MC	1H	1942	25	-	0/7/25/26	0/2/2/2
57	5MU	2L	55	57	-	0/7/25/26	0/2/2/2
25	OMU	1H	2552	59,25	-	0/9/27/28	0/2/2/2
1	MA6	1G	1519	1	-	3/11/29/30	0/3/3/3
25	PSU	1H	1917	25	-	1/7/25/26	0/2/2/2
1	4OC	13	1402	1	-	2/9/29/30	0/2/2/2
1	5MC	13	1407	1	-	0/7/25/26	0/2/2/2
1	PSU	13	516	59,1	-	0/7/25/26	0/2/2/2
25	5MU	1H	1939	25,58	-	0/7/25/26	0/2/2/2
1	5MC	1G	1407	1	-	0/7/25/26	0/2/2/2

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
1	PSU	1G	516	59,1	-	0/7/25/26	0/2/2/2
23	PSU	2K	56	23	-	2/7/25/26	0/2/2/2
25	5MU	14	1939	25,58	-	0/7/25/26	0/2/2/2
25	5MU	14	1915	25	-	3/7/25/26	0/2/2/2
1	MA6	13	1518	1	-	0/11/29/30	0/3/3/3
25	5MC	14	1962	25	-	2/7/25/26	0/2/2/2
25	PSU	14	2605	25	-	0/7/25/26	0/2/2/2
12	0TD	3A	89	12	-	5/7/12/14	-
1	4OC	1G	1402	1	-	2/9/29/30	0/2/2/2
1	MA6	1G	1518	1	-	2/11/29/30	0/3/3/3
25	PSU	14	1911	25	-	0/7/25/26	0/2/2/2
57	4SU	2L	8	57	-	3/7/25/26	0/2/2/2
1	5MC	13	1400	1	-	2/7/25/26	0/2/2/2
23	7MG	2K	47	23	-	2/7/37/38	0/3/3/3
25	OMC	14	1920	25	-	2/9/27/28	0/2/2/2
1	7MG	13	527	58,1	-	2/7/37/38	0/3/3/3
57	PSU	2L	56	57	-	0/7/25/26	0/2/2/2
1	7MG	1G	527	58,1	-	3/7/37/38	0/3/3/3
25	2MA	14	2503	59,25,58	-	0/7/25/26	0/3/3/3
25	OMG	1H	2251	25,58,23	-	1/9/27/28	0/3/3/3
1	5MC	1G	1400	1	-	2/7/25/26	0/2/2/2
1	UR3	13	1498	1	-	0/7/25/26	0/2/2/2
57	7MG	2L	47	57	-	3/7/37/38	0/3/3/3
1	UR3	1G	1498	1	-	0/7/25/26	0/2/2/2
1	5MC	13	1404	1	-	0/7/25/26	0/2/2/2
1	5MC	1G	1404	1	-	0/7/25/26	0/2/2/2
25	OMU	14	2552	59,25	-	1/9/27/28	0/2/2/2
25	5MC	14	1942	25	-	0/7/25/26	0/2/2/2
23	5MU	2K	55	23	-	0/7/25/26	0/2/2/2
23	OMC	2K	33	23	-	1/9/27/28	0/2/2/2
25	5MU	1H	1915	25	-	0/7/25/26	0/2/2/2
12	0TD	3I	89	12	-	3/7/12/14	-
1	2MG	1G	1207	1	-	0/9/27/28	0/3/3/3
1	MA6	13	1519	1	-	4/11/29/30	0/3/3/3
23	4SU	2K	8	23	-	0/7/25/26	0/2/2/2
25	PSU	14	1917	25	-	0/7/25/26	0/2/2/2
1	2MG	13	1207	1	-	1/9/27/28	0/3/3/3
25	5MC	1H	1962	25,58	-	2/7/25/26	0/2/2/2

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
1	5MC	1G	967	1	-	0/7/25/26	0/2/2/2
1	M2G	13	966	1	-	1/11/29/30	0/3/3/3
25	PSU	1H	1911	25	-	0/7/25/26	0/2/2/2
1	5MC	13	967	1	-	2/7/25/26	0/2/2/2

All (329) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
25	14	1915	5MU	C2-N1	13.36	1.59	1.38
25	1H	1915	5MU	C2-N1	13.12	1.59	1.38
57	2L	55	5MU	C2-N1	13.00	1.58	1.38
23	2K	55	5MU	C2-N1	12.97	1.58	1.38
25	1H	1939	5MU	C2-N1	12.34	1.57	1.38
25	14	1939	5MU	C2-N1	12.07	1.57	1.38
25	14	2251	OMG	C4-N3	11.71	1.61	1.34
25	14	2251	OMG	C5-N7	-11.70	1.15	1.39
25	1H	2251	OMG	C4-N3	11.44	1.60	1.34
25	1H	2251	OMG	C5-N7	-11.43	1.16	1.39
25	1H	2503	2MA	C4-N3	11.32	1.48	1.34
25	14	2503	2MA	C4-N3	11.29	1.48	1.34
1	1G	1407	5MC	C6-C5	9.82	1.50	1.34
25	1H	1962	5MC	C6-C5	9.72	1.50	1.34
1	13	1400	5MC	C6-C5	9.69	1.50	1.34
1	1G	967	5MC	C6-C5	9.62	1.50	1.34
1	1G	1400	5MC	C6-C5	9.62	1.50	1.34
1	13	1407	5MC	C6-C5	9.56	1.50	1.34
1	1G	1404	5MC	C6-C5	9.52	1.50	1.34
25	14	1942	5MC	C6-C5	9.44	1.50	1.34
1	13	967	5MC	C6-C5	9.42	1.50	1.34
25	14	1962	5MC	C6-C5	9.28	1.49	1.34
1	13	1404	5MC	C6-C5	9.27	1.49	1.34
25	1H	1942	5MC	C6-C5	8.97	1.49	1.34
25	1H	2251	OMG	C4-N9	8.89	1.60	1.38
25	14	2251	OMG	C4-N9	8.82	1.60	1.38
25	14	2251	OMG	C8-N7	-8.50	1.08	1.32
23	2K	47	7MG	C5-N7	8.41	1.46	1.35
25	1H	2251	OMG	C8-N7	-8.31	1.09	1.32
25	1H	2251	OMG	C5-C4	8.28	1.61	1.38
57	2L	47	7MG	C5-N7	8.17	1.46	1.35
25	14	2251	OMG	C5-C4	8.12	1.61	1.38
1	13	527	7MG	C5-N7	8.03	1.45	1.35
1	1G	1207	2MG	C2-N3	8.02	1.47	1.32

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	1G	527	7MG	C5-N7	7.98	1.45	1.35
1	13	1400	5MC	C4-N3	7.95	1.46	1.34
1	13	1400	5MC	C5-C4	7.92	1.50	1.44
25	1H	2251	OMG	C6-N1	-7.91	1.24	1.38
1	13	966	M2G	C2-N2	7.91	1.49	1.35
25	14	1962	5MC	C4-N3	7.76	1.46	1.34
1	1G	967	5MC	C4-N3	7.70	1.46	1.34
25	14	2251	OMG	C6-N1	-7.70	1.24	1.38
1	13	527	7MG	C4-N9	-7.69	1.28	1.37
1	13	1404	5MC	C5-C4	7.66	1.49	1.44
1	13	967	5MC	C4-N3	7.65	1.46	1.34
1	13	1207	2MG	C2-N3	7.63	1.46	1.32
1	1G	1498	UR3	C2-N1	7.63	1.49	1.38
25	14	1942	5MC	C4-N3	7.61	1.46	1.34
1	1G	1404	5MC	C4-N3	7.57	1.46	1.34
1	1G	1404	5MC	C5-C4	7.56	1.49	1.44
1	13	1404	5MC	C4-N3	7.53	1.46	1.34
1	13	1498	UR3	C2-N1	7.52	1.48	1.38
1	1G	1400	5MC	C4-N3	7.51	1.46	1.34
25	1H	1962	5MC	C4-N3	7.46	1.46	1.34
1	1G	966	M2G	C2-N2	7.43	1.48	1.35
1	1G	966	M2G	C4-N3	7.42	1.51	1.34
1	13	967	5MC	C5-C4	7.41	1.49	1.44
1	13	966	M2G	C4-N3	7.38	1.51	1.34
1	1G	967	5MC	C5-C4	7.34	1.49	1.44
1	1G	1407	5MC	C4-N3	7.29	1.45	1.34
1	1G	527	7MG	C4-N9	-7.29	1.28	1.37
1	13	1402	4OC	C4-N3	7.27	1.45	1.32
1	13	1407	5MC	C4-N3	7.27	1.45	1.34
23	2K	47	7MG	C4-N9	-7.26	1.28	1.37
25	1H	1942	5MC	C4-N3	7.25	1.45	1.34
57	2L	47	7MG	C4-N9	-7.23	1.29	1.37
1	13	1407	5MC	C5-C4	7.20	1.49	1.44
1	1G	1400	5MC	C5-C4	7.18	1.49	1.44
25	14	1962	5MC	C5-C4	7.09	1.49	1.44
1	1G	1402	4OC	C4-N3	7.08	1.44	1.32
25	1H	2552	OMU	C2-N3	7.07	1.50	1.38
1	1G	1207	2MG	C2-N2	7.05	1.48	1.33
1	1G	1207	2MG	C4-N3	7.02	1.50	1.34
25	1H	1962	5MC	C5-C4	7.02	1.49	1.44
25	14	2552	OMU	C2-N3	7.00	1.50	1.38
25	14	1942	5MC	C5-C4	6.99	1.49	1.44

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
25	14	2552	OMU	C2-N1	6.88	1.49	1.38
25	1H	1939	5MU	C4-N3	-6.87	1.26	1.38
1	13	1207	2MG	C4-N3	6.86	1.50	1.34
1	13	1207	2MG	C2-N2	6.85	1.47	1.33
1	13	1400	5MC	C2-N3	6.81	1.49	1.36
25	14	2503	2MA	C2-N1	6.81	1.45	1.34
25	14	1939	5MU	C4-N3	-6.77	1.26	1.38
25	14	1942	5MC	C2-N3	6.72	1.49	1.36
25	1H	1962	5MC	C2-N3	6.69	1.49	1.36
1	1G	1404	5MC	C2-N3	6.66	1.49	1.36
1	1G	1407	5MC	C2-N3	6.65	1.49	1.36
1	1G	967	5MC	C2-N3	6.64	1.49	1.36
1	1G	1400	5MC	C2-N3	6.57	1.49	1.36
1	13	1404	5MC	C2-N3	6.56	1.49	1.36
1	13	967	5MC	C2-N3	6.56	1.49	1.36
1	1G	1407	5MC	C5-C4	6.55	1.49	1.44
25	14	1962	5MC	C2-N3	6.54	1.49	1.36
1	13	1407	5MC	C2-N3	6.48	1.49	1.36
23	2K	21	H2U	C2-N1	6.48	1.44	1.35
25	1H	1942	5MC	C2-N3	6.43	1.49	1.36
25	1H	1915	5MU	C4-N3	-6.43	1.26	1.38
25	1H	2503	2MA	C2-N1	6.31	1.44	1.34
25	14	2503	2MA	C6-N1	6.29	1.43	1.35
25	14	2503	2MA	C2-N3	6.28	1.44	1.34
25	14	1915	5MU	C4-N3	-6.22	1.27	1.38
1	13	1498	UR3	C6-C5	6.21	1.49	1.35
57	2L	55	5MU	C4-N3	-6.18	1.27	1.38
1	1G	1498	UR3	C6-C5	6.16	1.49	1.35
23	2K	47	7MG	C4-N3	6.14	1.48	1.34
1	13	1402	4OC	C6-C5	6.14	1.49	1.35
1	13	1402	4OC	C2-N3	6.13	1.48	1.36
57	2L	8	4SU	C5-C4	6.11	1.49	1.42
57	2L	47	7MG	C4-N3	6.10	1.48	1.34
1	1G	527	7MG	C4-N3	6.09	1.48	1.34
23	2K	55	5MU	C4-N3	-6.07	1.27	1.38
25	1H	2552	OMU	C2-N1	6.04	1.47	1.38
1	13	966	M2G	C2-N3	5.98	1.46	1.32
25	1H	2503	2MA	C2-N3	5.96	1.44	1.34
1	13	527	7MG	C4-N3	5.93	1.47	1.34
23	2K	55	5MU	C2-N3	5.93	1.48	1.38
1	1G	1402	4OC	C2-N3	5.88	1.48	1.36
25	1H	2503	2MA	C6-N1	5.85	1.42	1.35

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	1G	966	M2G	C2-N3	5.82	1.46	1.32
25	14	1915	5MU	C2-N3	5.80	1.48	1.38
57	2L	55	5MU	C2-N3	5.78	1.48	1.38
25	14	2552	OMU	C6-C5	5.77	1.48	1.35
1	1G	1402	4OC	C6-C5	5.76	1.48	1.35
25	1H	2552	OMU	C6-C5	5.75	1.48	1.35
25	14	1915	5MU	C6-N1	5.70	1.47	1.38
23	2K	8	4SU	C5-C4	5.68	1.49	1.42
23	2K	55	5MU	C6-N1	5.65	1.47	1.38
25	1H	1942	5MC	C5-C4	5.64	1.48	1.44
25	1H	1915	5MU	C6-N1	5.60	1.47	1.38
57	2L	55	5MU	C6-N1	5.58	1.47	1.38
25	1H	1915	5MU	C2-N3	5.56	1.47	1.38
25	14	1939	5MU	C6-N1	5.34	1.47	1.38
1	13	1498	UR3	C2-N3	5.24	1.49	1.39
1	13	1400	5MC	C4-N4	5.11	1.47	1.34
25	14	1962	5MC	C4-N4	5.11	1.47	1.34
25	14	1939	5MU	C2-N3	5.10	1.46	1.38
25	1H	1915	5MU	C4-C5	5.10	1.53	1.44
23	2K	55	5MU	C4-C5	5.09	1.53	1.44
57	2L	55	5MU	C4-C5	5.07	1.53	1.44
1	13	967	5MC	C4-N4	5.07	1.47	1.34
1	1G	1498	UR3	C2-N3	5.05	1.49	1.39
25	1H	1962	5MC	C4-N4	5.04	1.47	1.34
1	1G	1404	5MC	C4-N4	5.02	1.46	1.34
25	14	1942	5MC	C4-N4	5.01	1.46	1.34
1	1G	1400	5MC	C4-N4	5.01	1.46	1.34
1	13	1404	5MC	C4-N4	5.00	1.46	1.34
1	1G	967	5MC	C4-N4	5.00	1.46	1.34
25	14	1962	5MC	C2-N1	4.97	1.50	1.40
25	1H	1939	5MU	C6-N1	4.95	1.46	1.38
25	14	1915	5MU	C4-C5	4.95	1.52	1.44
1	13	1407	5MC	C4-N4	4.93	1.46	1.34
25	1H	1939	5MU	C2-N3	4.91	1.46	1.38
25	1H	1942	5MC	C4-N4	4.89	1.46	1.34
1	1G	1407	5MC	C4-N4	4.88	1.46	1.34
1	13	1407	5MC	C6-N1	4.88	1.46	1.38
1	1G	1207	2MG	C2-N1	4.87	1.44	1.36
1	1G	1404	5MC	C6-N1	4.87	1.46	1.38
1	1G	967	5MC	C6-N1	4.85	1.46	1.38
25	1H	1962	5MC	C6-N1	4.85	1.46	1.38
25	1H	1939	5MU	C4-C5	4.79	1.52	1.44

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	1G	1400	5MC	C6-N1	4.77	1.46	1.38
1	13	1400	5MC	C6-N1	4.76	1.46	1.38
1	13	967	5MC	C6-N1	4.73	1.46	1.38
1	1G	1407	5MC	C6-N1	4.73	1.46	1.38
1	13	1207	2MG	C2-N1	4.71	1.44	1.36
1	1G	1400	5MC	C2-N1	4.71	1.49	1.40
23	2K	33	OMC	C2-N3	4.69	1.45	1.36
1	1G	1407	5MC	C2-N1	4.68	1.49	1.40
1	13	1400	5MC	C2-N1	4.68	1.49	1.40
23	2K	47	7MG	C8-N7	4.66	1.64	1.42
57	2L	33	OMC	C2-N3	4.65	1.45	1.36
25	14	1942	5MC	C6-N1	4.62	1.45	1.38
57	2L	47	7MG	C8-N7	4.61	1.64	1.42
1	13	527	7MG	C5-C4	-4.61	1.23	1.37
1	13	966	M2G	C2-N1	4.60	1.47	1.36
25	14	1962	5MC	C6-N1	4.59	1.45	1.38
1	1G	527	7MG	C8-N7	4.59	1.64	1.42
23	2K	47	7MG	C5-C4	-4.58	1.23	1.37
1	13	527	7MG	C8-N7	4.57	1.64	1.42
1	1G	527	7MG	C5-C4	-4.56	1.24	1.37
25	1H	1962	5MC	C2-N1	4.55	1.49	1.40
57	2L	47	7MG	C5-C4	-4.55	1.24	1.37
1	1G	1404	5MC	C2-N1	4.53	1.49	1.40
1	13	1404	5MC	C6-N1	4.53	1.45	1.38
25	14	2251	OMG	C8-N9	-4.52	1.27	1.37
1	1G	966	M2G	C2-N1	4.47	1.47	1.36
25	14	1920	OMC	C2-N3	4.47	1.45	1.36
25	14	1939	5MU	C4-C5	4.46	1.52	1.44
25	14	1942	5MC	C2-N1	4.46	1.49	1.40
25	1H	1920	OMC	C2-N3	4.46	1.45	1.36
25	1H	1942	5MC	C2-N1	4.43	1.49	1.40
1	1G	967	5MC	C2-N1	4.43	1.49	1.40
1	13	1407	5MC	C2-N1	4.42	1.49	1.40
1	13	1404	5MC	C2-N1	4.39	1.49	1.40
25	1H	2251	OMG	C8-N9	-4.36	1.27	1.37
25	1H	1942	5MC	C6-N1	4.35	1.45	1.38
1	13	967	5MC	C2-N1	4.27	1.49	1.40
25	14	1920	OMC	C4-N4	4.27	1.44	1.33
57	2L	33	OMC	C4-N4	4.26	1.44	1.33
25	1H	2552	OMU	C4-N3	4.26	1.45	1.38
23	2K	21	H2U	C2-N3	4.24	1.45	1.38
23	2K	33	OMC	C4-N4	4.21	1.44	1.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
25	1H	1920	OMC	C4-N4	4.19	1.44	1.33
1	13	1402	4OC	C4-N4	4.18	1.44	1.36
1	13	1402	4OC	C2-N1	4.16	1.48	1.40
1	1G	1402	4OC	C2-N1	4.11	1.48	1.40
25	14	2552	OMU	C4-N3	4.11	1.45	1.38
57	2L	56	PSU	C6-C5	4.10	1.39	1.35
23	2K	8	4SU	C2-N1	3.96	1.44	1.38
1	13	516	PSU	C6-C5	3.95	1.39	1.35
25	1H	1917	PSU	C6-C5	3.94	1.39	1.35
23	2K	56	PSU	C6-C5	3.94	1.39	1.35
1	1G	516	PSU	C6-C5	3.94	1.39	1.35
25	14	1911	PSU	C6-C5	3.90	1.39	1.35
25	14	2605	PSU	C6-C5	3.85	1.39	1.35
25	1H	1911	PSU	C6-C5	3.84	1.39	1.35
25	14	1920	OMC	C5-C4	3.82	1.51	1.42
1	1G	966	M2G	C5-N7	-3.74	1.31	1.39
1	13	1402	4OC	C5-C4	3.72	1.49	1.41
1	1G	1519	MA6	C6-N6	3.70	1.47	1.36
25	14	1917	PSU	C6-C5	3.69	1.39	1.35
25	1H	2605	PSU	C6-C5	3.69	1.39	1.35
57	2L	33	OMC	C5-C4	3.66	1.51	1.42
23	2K	33	OMC	C5-C4	3.65	1.51	1.42
25	14	2503	2MA	C5-N7	-3.65	1.32	1.39
1	1G	1402	4OC	C4-N4	3.60	1.43	1.36
1	13	966	M2G	C5-N7	-3.59	1.31	1.39
25	1H	2503	2MA	C5-N7	-3.58	1.32	1.39
1	13	1518	MA6	C6-N6	3.55	1.46	1.36
25	1H	1920	OMC	C5-C4	3.54	1.51	1.42
1	1G	1402	4OC	C5-C4	3.54	1.48	1.41
1	13	1519	MA6	C6-N6	3.45	1.46	1.36
23	2K	21	H2U	C4-N3	3.44	1.43	1.37
1	1G	1518	MA6	C6-N6	3.44	1.46	1.36
25	1H	2503	2MA	C5-C6	3.43	1.50	1.41
25	14	2503	2MA	C5-C6	3.39	1.50	1.41
1	1G	1207	2MG	C5-N7	-3.32	1.32	1.39
57	2L	8	4SU	C2-N1	3.25	1.43	1.38
25	14	1962	5MC	O2-C2	-3.17	1.17	1.23
1	1G	1407	5MC	O2-C2	-3.16	1.17	1.23
1	13	1207	2MG	O6-C6	-3.16	1.17	1.23
1	1G	1518	MA6	C8-N9	-3.13	1.32	1.37
1	13	1402	4OC	C6-N1	3.12	1.45	1.38
1	1G	527	7MG	C2-N2	3.12	1.41	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	13	1518	MA6	C8-N9	-3.11	1.32	1.37
1	13	1404	5MC	O2-C2	-3.11	1.17	1.23
1	13	527	7MG	C2-N2	3.10	1.41	1.34
1	1G	1207	2MG	O6-C6	-3.10	1.17	1.23
1	13	1519	MA6	C5-N7	-3.09	1.33	1.39
23	2K	47	7MG	C2-N2	3.07	1.41	1.34
1	13	967	5MC	O2-C2	-3.07	1.18	1.23
25	1H	2251	OMG	O6-C6	-3.06	1.17	1.23
57	2L	47	7MG	C2-N2	3.06	1.41	1.34
1	1G	1518	MA6	C5-N7	-3.04	1.33	1.39
1	1G	1404	5MC	O2-C2	-3.04	1.18	1.23
1	1G	967	5MC	O2-C2	-3.02	1.18	1.23
1	1G	1498	UR3	C6-N1	3.02	1.45	1.38
25	1H	1962	5MC	O2-C2	-3.02	1.18	1.23
1	13	1207	2MG	C5-N7	-3.01	1.33	1.39
25	14	2251	OMG	O6-C6	-3.00	1.17	1.23
1	13	1498	UR3	C6-N1	2.97	1.45	1.38
25	1H	1942	5MC	O2-C2	-2.97	1.18	1.23
1	1G	1400	5MC	O2-C2	-2.94	1.18	1.23
25	14	2251	OMG	C2-N3	2.94	1.40	1.33
25	14	2552	OMU	O4-C4	-2.93	1.18	1.24
1	13	1407	5MC	O2-C2	-2.92	1.18	1.23
1	1G	1402	4OC	C6-N1	2.90	1.45	1.38
1	13	1518	MA6	C5-N7	-2.89	1.33	1.39
25	14	1942	5MC	O2-C2	-2.88	1.18	1.23
1	13	1400	5MC	O2-C2	-2.87	1.18	1.23
1	13	966	M2G	C5-C6	2.84	1.55	1.44
1	13	1519	MA6	C8-N9	-2.83	1.32	1.37
1	1G	966	M2G	C5-C6	2.82	1.54	1.44
57	2L	8	4SU	C6-N1	2.81	1.44	1.38
25	14	2552	OMU	C6-N1	2.80	1.44	1.38
1	13	1518	MA6	C5-C4	-2.79	1.34	1.39
1	1G	1519	MA6	C8-N9	-2.77	1.32	1.37
25	1H	2251	OMG	C2-N3	2.76	1.39	1.33
25	14	2251	OMG	C2-N2	2.74	1.40	1.34
25	1H	2552	OMU	O4-C4	-2.73	1.19	1.24
25	1H	2552	OMU	C6-N1	2.70	1.44	1.38
1	1G	1519	MA6	C5-N7	-2.69	1.34	1.39
1	13	1402	4OC	O2-C2	-2.68	1.18	1.23
1	13	966	M2G	C4-N9	-2.66	1.31	1.38
1	1G	1518	MA6	C5-C4	-2.66	1.34	1.39
1	1G	1402	4OC	O2-C2	-2.61	1.18	1.23

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	13	1519	MA6	C5-C4	-2.61	1.34	1.39
1	13	1207	2MG	C5-C6	2.59	1.54	1.44
23	2K	21	H2U	C6-N1	-2.59	1.42	1.47
1	1G	1498	UR3	C4-N3	2.53	1.45	1.40
1	13	966	M2G	C6-N1	2.51	1.43	1.38
25	1H	2251	OMG	C2-N2	2.49	1.40	1.34
1	13	1498	UR3	C4-N3	2.49	1.45	1.40
1	1G	1207	2MG	C5-C6	2.48	1.53	1.44
25	1H	2251	OMG	C2-N1	-2.48	1.31	1.37
25	14	2552	OMU	C5-C4	2.47	1.49	1.43
25	1H	2552	OMU	O2-C2	-2.46	1.18	1.23
23	2K	8	4SU	C6-N1	2.44	1.43	1.38
25	1H	2552	OMU	C5-C4	2.40	1.48	1.43
25	14	2552	OMU	O2-C2	-2.39	1.18	1.23
25	14	2503	2MA	C8-N7	2.39	1.36	1.31
1	1G	1519	MA6	C8-N7	2.38	1.36	1.31
25	14	2251	OMG	C2-N1	-2.37	1.31	1.37
1	1G	966	M2G	C6-N1	2.36	1.43	1.38
1	13	1207	2MG	C6-N1	2.33	1.43	1.38
1	1G	1498	UR3	O4-C4	-2.29	1.18	1.23
1	13	1498	UR3	C5-C4	2.28	1.49	1.43
1	1G	1519	MA6	C5-C4	-2.27	1.35	1.39
1	1G	966	M2G	C4-N9	-2.25	1.32	1.38
23	2K	47	7MG	C5-C6	2.24	1.49	1.43
25	14	2503	2MA	CM2-C2	2.23	1.55	1.49
25	1H	2503	2MA	C8-N7	2.21	1.35	1.31
1	1G	1498	UR3	C5-C4	2.21	1.49	1.43
1	13	1498	UR3	O4-C4	-2.18	1.18	1.23
25	1H	2605	PSU	C4-C5	-2.18	1.38	1.44
1	13	1518	MA6	C8-N7	2.17	1.35	1.31
57	2L	33	OMC	C6-N1	2.17	1.43	1.38
1	1G	1207	2MG	C6-N1	2.15	1.42	1.38
25	1H	1920	OMC	C6-N1	2.13	1.43	1.38
1	13	1519	MA6	C8-N7	2.12	1.35	1.31
1	1G	527	7MG	C5-C6	2.10	1.48	1.43
1	1G	1518	MA6	C8-N7	2.10	1.35	1.31
23	2K	47	7MG	C6-N1	-2.09	1.34	1.38
1	13	527	7MG	C6-N1	-2.05	1.35	1.38
57	2L	47	7MG	C5-C6	2.04	1.48	1.43
25	14	1920	OMC	C6-N1	2.03	1.42	1.38
1	1G	527	7MG	C8-N9	2.03	1.47	1.45
57	2L	47	7MG	C6-N1	-2.02	1.35	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	13	527	7MG	C5-C6	2.01	1.48	1.43

All (339) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
25	1H	1939	5MU	C5-C4-N3	11.63	125.43	115.32
25	1H	2251	OMG	C6-C5-N7	11.24	150.75	130.29
25	14	1939	5MU	C5-C4-N3	11.05	124.93	115.32
25	14	2251	OMG	C6-C5-N7	10.94	150.20	130.29
25	1H	1915	5MU	C5-C4-N3	10.50	124.45	115.32
25	14	1915	5MU	C5-C4-N3	10.48	124.43	115.32
57	2L	55	5MU	C5-C4-N3	10.28	124.26	115.32
23	2K	55	5MU	C5-C4-N3	10.06	124.07	115.32
25	1H	2251	OMG	C8-N7-C5	10.03	122.12	104.26
25	14	2251	OMG	N9-C4-N3	9.97	145.89	125.95
25	14	2251	OMG	C8-N7-C5	9.94	121.96	104.26
25	1H	2251	OMG	N9-C4-N3	9.72	145.39	125.95
1	1G	1519	MA6	N1-C6-N6	-8.51	106.48	116.86
25	1H	2251	OMG	C1'-N9-C4	8.43	151.39	126.49
25	14	2251	OMG	C1'-N9-C4	8.13	150.52	126.49
1	13	527	7MG	C4-C5-N7	8.06	114.89	105.38
1	13	1518	MA6	N1-C6-N6	-8.01	107.10	116.86
57	2L	47	7MG	C4-C5-N7	7.96	114.79	105.38
23	2K	8	4SU	C4-N3-C2	-7.91	119.73	127.31
1	1G	527	7MG	C4-C5-N7	7.83	114.63	105.38
25	1H	2503	2MA	C5-C4-N3	-7.77	118.99	127.18
25	1H	1939	5MU	C5-C6-N1	-7.71	114.93	123.31
23	2K	47	7MG	C4-C5-N7	7.60	114.36	105.38
25	1H	1939	5MU	C4-N3-C2	-7.57	117.41	127.34
1	1G	1518	MA6	N1-C6-N6	-7.50	107.72	116.86
25	14	2251	OMG	C6-C5-C4	-7.50	107.55	118.83
25	1H	2251	OMG	C6-C5-C4	-7.43	107.65	118.83
1	13	1519	MA6	N1-C6-N6	-7.33	107.93	116.86
1	13	1207	2MG	C2-N3-C4	6.82	120.53	112.00
25	14	1939	5MU	C4-N3-C2	-6.82	118.40	127.34
25	1H	1939	5MU	C6-C5-C4	6.74	123.58	118.02
25	14	1939	5MU	C5-C6-N1	-6.64	116.10	123.31
25	14	2251	OMG	C5-C6-N1	6.64	130.15	113.25
25	1H	2251	OMG	C5-C6-N1	6.58	130.01	113.25
1	1G	1207	2MG	C2-N3-C4	6.49	120.12	112.00
25	1H	1915	5MU	C4-N3-C2	-6.41	118.93	127.34
25	1H	2251	OMG	C5-C4-N3	-6.40	118.21	128.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
57	2L	8	4SU	C4-N3-C2	-6.39	121.19	127.31
25	14	2552	OMU	C4-N3-C2	-6.32	118.76	126.61
25	14	2251	OMG	C5-C4-N3	-6.27	118.42	128.39
57	2L	55	5MU	C4-N3-C2	-6.22	119.19	127.34
25	14	1915	5MU	C4-N3-C2	-6.18	119.23	127.34
23	2K	55	5MU	C4-N3-C2	-6.09	119.35	127.34
25	14	1939	5MU	C6-C5-C4	6.04	123.00	118.02
1	1G	527	7MG	C5-C4-N9	5.99	114.00	106.33
23	2K	47	7MG	C5-C4-N9	5.93	113.93	106.33
1	1G	1207	2MG	C5-C4-N3	-5.88	119.03	128.39
1	13	1519	MA6	N1-C2-N3	-5.88	119.69	128.58
1	13	527	7MG	C5-C4-N9	5.83	113.80	106.33
25	14	2503	2MA	C5-C4-N3	-5.82	121.05	127.18
25	1H	2251	OMG	C4-C5-N7	-5.73	101.59	110.67
25	1H	2552	OMU	C4-N3-C2	-5.68	119.57	126.61
1	13	1207	2MG	C5-C4-N3	-5.63	119.43	128.39
1	13	1518	MA6	N1-C2-N3	-5.62	120.07	128.58
57	2L	55	5MU	C6-C5-C4	5.61	122.65	118.02
23	2K	55	5MU	C6-C5-C4	5.60	122.64	118.02
57	2L	47	7MG	C5-C4-N9	5.60	113.51	106.33
1	1G	1518	MA6	N1-C2-N3	-5.59	120.12	128.58
25	14	2251	OMG	C5-C4-N9	-5.57	95.69	105.66
25	1H	1915	5MU	C6-C5-C4	5.56	122.60	118.02
1	1G	1519	MA6	N1-C2-N3	-5.54	120.19	128.58
25	14	1915	5MU	C6-C5-C4	5.52	122.57	118.02
25	1H	2503	2MA	N3-C4-N9	5.51	133.98	126.99
23	2K	8	4SU	C5-C4-N3	5.48	119.85	114.75
1	1G	966	M2G	C5-C4-N3	-5.38	119.83	128.39
1	13	1498	UR3	C4-N3-C2	-5.37	120.25	124.58
57	2L	55	5MU	C5-C6-N1	-5.35	117.50	123.31
25	14	2251	OMG	C4-C5-N7	-5.32	102.25	110.67
1	1G	1498	UR3	C4-N3-C2	-5.30	120.32	124.58
25	1H	1915	5MU	C5-C6-N1	-5.29	117.56	123.31
25	1H	2251	OMG	C1'-N9-C8	-5.29	111.70	126.73
1	13	1519	MA6	C5-C4-N3	-5.28	119.45	126.72
12	3A	89	0TD	OD2-CG-CB	5.24	124.47	113.15
23	2K	55	5MU	C5-C6-N1	-5.24	117.63	123.31
25	14	1915	5MU	C5-C6-N1	-5.21	117.65	123.31
1	1G	1519	MA6	C5-C6-N6	5.17	133.52	125.33
25	1H	2251	OMG	N9-C8-N7	5.17	122.99	113.40
25	1H	2251	OMG	C5-C4-N9	-5.17	96.40	105.66
25	14	2251	OMG	C1'-N9-C8	-5.16	112.08	126.73

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
25	1H	2605	PSU	C4-N3-C2	-5.13	119.30	126.37
23	2K	47	7MG	C5-C6-N1	5.11	119.94	110.94
25	1H	1939	5MU	O4-C4-C5	-5.08	119.10	124.92
25	14	1939	5MU	O4-C4-C5	-5.05	119.14	124.92
25	1H	2605	PSU	N1-C2-N3	5.05	120.49	115.17
1	13	527	7MG	C5-C6-N1	5.03	119.79	110.94
25	1H	1917	PSU	C4-N3-C2	-5.03	119.45	126.37
23	2K	55	5MU	O4-C4-C5	-5.03	119.17	124.92
25	14	2251	OMG	N9-C8-N7	5.02	122.72	113.40
25	14	1915	5MU	O4-C4-C5	-5.00	119.19	124.92
1	13	966	M2G	C5-C4-N3	-5.00	120.43	128.39
1	1G	516	PSU	C4-N3-C2	-5.00	119.49	126.37
25	1H	1917	PSU	N1-C2-N3	4.99	120.43	115.17
23	2K	56	PSU	C4-N3-C2	-4.98	119.52	126.37
57	2L	47	7MG	C5-C6-N1	4.97	119.69	110.94
25	14	1917	PSU	C4-N3-C2	-4.95	119.55	126.37
25	14	1917	PSU	N1-C2-N3	4.91	120.35	115.17
1	1G	527	7MG	C5-C6-N1	4.89	119.55	110.94
1	1G	1519	MA6	C5-C4-N3	-4.89	119.98	126.72
57	2L	56	PSU	C4-N3-C2	-4.87	119.66	126.37
25	1H	2251	OMG	C8-N9-C4	-4.87	96.89	106.03
1	13	1518	MA6	C5-C6-N6	4.85	133.01	125.33
23	2K	56	PSU	N1-C2-N3	4.81	120.24	115.17
57	2L	56	PSU	N1-C2-N3	4.74	120.17	115.17
1	1G	1518	MA6	C5-C4-N3	-4.74	120.19	126.72
25	1H	1911	PSU	N1-C2-N3	4.73	120.16	115.17
25	14	2605	PSU	C4-N3-C2	-4.72	119.88	126.37
1	13	1518	MA6	C5-C4-N3	-4.71	120.23	126.72
1	13	1518	MA6	N9-C8-N7	-4.69	107.28	113.94
57	2L	8	4SU	C5-C4-N3	4.69	119.11	114.75
57	2L	55	5MU	C5M-C5-C6	-4.66	116.55	122.85
57	2L	55	5MU	O4-C4-C5	-4.65	119.60	124.92
23	2K	8	4SU	C5-C4-S4	-4.65	119.00	124.31
25	14	2251	OMG	C8-N9-C4	-4.61	97.38	106.03
25	1H	1915	5MU	C5M-C5-C6	-4.56	116.68	122.85
25	14	2552	OMU	N3-C2-N1	4.53	120.79	114.89
1	1G	516	PSU	N1-C2-N3	4.52	119.94	115.17
1	13	516	PSU	C4-N3-C2	-4.52	120.14	126.37
1	1G	527	7MG	C6-C5-N7	-4.48	124.99	131.93
1	1G	1518	MA6	C5-C6-N6	4.46	132.39	125.33
25	1H	1911	PSU	C4-N3-C2	-4.45	120.24	126.37
25	1H	1915	5MU	O4-C4-C5	-4.45	119.83	124.92

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	13	527	7MG	C6-C5-N7	-4.44	125.06	131.93
25	14	2503	2MA	C4-N9-C1'	-4.42	116.30	126.63
25	14	1911	PSU	N1-C2-N3	4.42	119.83	115.17
57	2L	47	7MG	C2-N3-C4	4.41	119.89	112.30
23	2K	47	7MG	C2-N3-C4	4.39	119.86	112.30
23	2K	55	5MU	C5M-C5-C6	-4.36	116.96	122.85
1	1G	527	7MG	C2-N3-C4	4.35	119.79	112.30
25	14	2251	OMG	O6-C6-C5	-4.33	115.10	126.53
1	13	527	7MG	C2-N3-C4	4.31	119.72	112.30
57	2L	47	7MG	C6-C5-N7	-4.30	125.26	131.93
1	13	1519	MA6	N9-C8-N7	-4.29	107.85	113.94
25	14	1915	5MU	C5M-C5-C6	-4.28	117.06	122.85
12	3I	89	0TD	CSB-SB-CB	4.28	110.05	102.36
25	14	2605	PSU	N1-C2-N3	4.26	119.67	115.17
1	13	1207	2MG	C2-N1-C6	-4.26	119.40	124.55
12	3I	89	0TD	OD2-CG-CB	4.26	122.35	113.15
1	1G	527	7MG	C5-C4-N3	-4.21	120.23	128.13
1	1G	1518	MA6	N9-C8-N7	-4.20	107.98	113.94
25	14	1911	PSU	C4-N3-C2	-4.16	120.64	126.37
1	13	516	PSU	N1-C2-N3	4.14	119.54	115.17
25	1H	1939	5MU	C5M-C5-C6	-4.10	117.30	122.85
57	2L	47	7MG	C5-C4-N3	-4.07	120.48	128.13
23	2K	8	4SU	N3-C2-N1	4.06	120.18	114.89
12	3A	89	0TD	CSB-SB-CB	4.06	109.66	102.36
1	13	966	M2G	C2-N3-C4	4.06	120.01	112.51
1	13	1519	MA6	C4-C5-C6	4.05	120.10	115.91
1	1G	1519	MA6	N9-C8-N7	-4.05	108.19	113.94
1	13	527	7MG	C5-C4-N3	-4.05	120.53	128.13
23	2K	47	7MG	C5-C4-N3	-4.03	120.56	128.13
1	1G	1207	2MG	C2-N1-C6	-4.03	119.68	124.55
25	1H	2552	OMU	N3-C2-N1	4.01	120.12	114.89
1	1G	1518	MA6	C4-C5-C6	4.00	120.05	115.91
1	1G	1207	2MG	N9-C4-N3	3.98	133.92	125.95
25	14	2503	2MA	N3-C4-N9	3.97	132.03	126.99
1	13	1519	MA6	C5-C6-N6	3.97	131.61	125.33
57	2L	8	4SU	N3-C2-N1	3.96	120.04	114.89
25	1H	2251	OMG	O6-C6-C5	-3.92	116.17	126.53
25	1H	2503	2MA	N9-C8-N7	-3.92	108.38	113.94
23	2K	21	H2U	C5-C4-N3	3.90	120.84	116.69
25	14	2503	2MA	N9-C8-N7	-3.87	108.45	113.94
1	1G	966	M2G	C2-N3-C4	3.86	119.64	112.51
25	1H	1915	5MU	C6-N1-C2	-3.85	117.47	121.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
23	2K	47	7MG	C6-C5-N7	-3.85	125.97	131.93
1	1G	1498	UR3	C5-C4-N3	3.79	120.03	115.04
25	14	2503	2MA	C1'-N9-C8	3.74	135.41	127.09
25	14	2552	OMU	C5-C4-N3	3.73	120.02	114.80
1	13	1518	MA6	C4-C5-C6	3.71	119.75	115.91
1	1G	1519	MA6	C4-C5-C6	3.68	119.71	115.91
25	14	2503	2MA	N6-C6-N1	3.66	121.96	117.03
1	1G	1498	UR3	C1'-N1-C2	3.65	123.02	117.04
25	14	1915	5MU	C6-N1-C2	-3.63	117.69	121.30
1	13	967	5MC	C5-C6-N1	-3.60	119.41	123.31
1	13	1519	MA6	N3-C4-N9	3.56	133.22	127.17
25	1H	2552	OMU	O2-C2-N1	-3.54	118.19	122.80
1	13	1207	2MG	N9-C4-N3	3.53	133.02	125.95
23	2K	21	H2U	C5-C6-N1	3.52	122.18	111.52
1	13	1519	MA6	C2-N3-C4	3.52	120.42	111.83
1	13	1498	UR3	C5-C4-N3	3.48	119.62	115.04
23	2K	55	5MU	C6-N1-C2	-3.47	117.85	121.30
1	13	1407	5MC	C5-C6-N1	-3.46	119.55	123.31
23	2K	21	H2U	N3-C2-N1	3.43	120.09	116.65
1	13	1518	MA6	C2-N1-C6	3.42	120.19	111.83
25	1H	1942	5MC	C5-C6-N1	-3.42	119.60	123.31
1	1G	1518	MA6	C2-N1-C6	3.40	120.13	111.83
1	13	1404	5MC	C5-C6-N1	-3.40	119.62	123.31
1	13	1400	5MC	C5-C6-N1	-3.39	119.63	123.31
1	1G	1519	MA6	C2-N1-C6	3.37	120.07	111.83
57	2L	55	5MU	C6-N1-C2	-3.37	117.94	121.30
1	1G	1402	4OC	CM4-N4-C4	-3.37	115.87	122.45
1	1G	1519	MA6	C2-N3-C4	3.37	120.05	111.83
25	1H	2251	OMG	O6-C6-N1	-3.35	113.81	120.11
1	1G	966	M2G	N9-C4-N3	3.34	132.63	125.95
25	1H	2552	OMU	C5-C4-N3	3.33	119.47	114.80
25	14	1962	5MC	C1'-N1-C6	-3.31	115.70	121.15
1	13	1519	MA6	C2-N1-C6	3.30	119.89	111.83
1	13	1518	MA6	C2-N3-C4	3.29	119.88	111.83
1	1G	1400	5MC	C5-C6-N1	-3.29	119.75	123.31
25	14	1939	5MU	C5M-C5-C6	-3.28	118.41	122.85
25	1H	1939	5MU	N3-C2-N1	3.28	119.16	114.89
25	14	2503	2MA	N3-C2-N1	-3.28	119.99	125.77
25	1H	1911	PSU	O2-C2-N1	-3.24	119.45	122.79
23	2K	47	7MG	N9-C8-N7	-3.22	98.81	103.37
25	1H	1962	5MC	C5-C6-N1	-3.21	119.82	123.31
1	1G	1518	MA6	C2-N3-C4	3.19	119.62	111.83

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	13	1207	2MG	N1-C2-N2	3.18	119.81	116.56
1	1G	1518	MA6	N3-C4-N9	3.18	132.57	127.17
12	3A	89	0TD	OD2-CG-OD1	-3.18	116.87	124.08
25	1H	2503	2MA	C6-C5-C4	3.16	121.49	117.18
1	13	527	7MG	O6-C6-C5	-3.16	119.86	127.62
1	1G	1407	5MC	C5-C6-N1	-3.15	119.89	123.31
25	1H	1915	5MU	N3-C2-N1	3.15	118.99	114.89
25	1H	2605	PSU	O2-C2-N1	-3.15	119.54	122.79
1	13	1518	MA6	C4-N9-C8	3.12	109.02	105.74
25	1H	2503	2MA	N6-C6-N1	3.10	121.22	117.03
57	2L	47	7MG	N9-C8-N7	-3.10	98.98	103.37
1	13	1518	MA6	C5-N7-C8	3.10	108.32	103.45
25	14	1939	5MU	N3-C2-N1	3.10	118.92	114.89
25	14	1942	5MC	C5-C6-N1	-3.09	119.96	123.31
1	13	966	M2G	N9-C8-N7	-3.08	107.69	113.40
1	1G	1404	5MC	C5-C6-N1	-3.07	119.98	123.31
57	2L	47	7MG	O6-C6-C5	-3.06	120.10	127.62
1	13	1518	MA6	N3-C4-N9	3.05	132.36	127.17
1	13	527	7MG	C2-N1-C6	-3.04	119.59	125.11
1	1G	527	7MG	N9-C8-N7	-3.01	99.10	103.37
1	1G	527	7MG	O6-C6-C5	-3.01	120.24	127.62
25	14	1915	5MU	C1'-N1-C2	3.00	122.99	117.59
25	1H	2605	PSU	C6-C5-C4	3.00	120.20	118.17
1	1G	527	7MG	C2-N1-C6	-2.98	119.71	125.11
1	13	1207	2MG	N9-C8-N7	-2.98	107.88	113.40
25	14	2552	OMU	O2-C2-N1	-2.96	118.94	122.80
23	2K	47	7MG	C2-N1-C6	-2.95	119.75	125.11
1	13	1519	MA6	C5-N7-C8	2.95	108.09	103.45
1	13	1404	5MC	CM5-C5-C6	-2.95	118.86	122.85
1	1G	967	5MC	C5-C6-N1	-2.94	120.11	123.31
1	13	527	7MG	N9-C8-N7	-2.91	99.25	103.37
25	1H	2503	2MA	N3-C2-N1	-2.91	120.63	125.77
1	1G	1519	MA6	N3-C4-N9	2.90	132.10	127.17
1	1G	1519	MA6	C5-N7-C8	2.89	108.00	103.45
25	14	1911	PSU	O2-C2-N1	-2.88	119.81	122.79
1	13	966	M2G	N9-C4-N3	2.88	131.71	125.95
23	2K	47	7MG	O6-C6-C5	-2.86	120.60	127.62
25	14	2251	OMG	O6-C6-N1	-2.85	114.76	120.11
25	14	1911	PSU	C6-N1-C2	-2.84	120.05	122.69
57	2L	47	7MG	C2-N1-C6	-2.84	119.96	125.11
25	14	2552	OMU	O4-C4-C5	-2.83	120.28	125.16
25	1H	2552	OMU	O4-C4-C5	-2.83	120.29	125.16

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
25	1H	2503	2MA	C5-N7-C8	2.82	107.88	103.45
25	1H	1911	PSU	C6-N1-C2	-2.81	120.08	122.69
1	1G	1518	MA6	C5-N7-C8	2.81	107.86	103.45
25	14	1962	5MC	C1'-N1-C2	2.78	124.59	118.44
23	2K	55	5MU	N3-C2-N1	2.75	118.47	114.89
57	2L	55	5MU	N3-C2-N1	2.74	118.46	114.89
1	13	966	M2G	C5-C6-N1	2.74	120.22	113.25
1	13	1498	UR3	C1'-N1-C2	2.73	121.51	117.04
25	1H	2251	OMG	N2-C2-N1	2.73	122.51	116.76
25	14	1915	5MU	N3-C2-N1	2.72	118.43	114.89
1	13	1207	2MG	C5-C6-N1	2.70	120.13	113.25
1	13	1498	UR3	C6-N1-C2	-2.67	119.61	121.80
25	14	1939	5MU	C6-N1-C2	-2.67	118.64	121.30
1	1G	966	M2G	N9-C8-N7	-2.66	108.46	113.40
1	1G	1518	MA6	C4-N9-C8	2.65	108.53	105.74
25	14	1962	5MC	C5-C6-N1	-2.59	120.50	123.31
57	2L	33	OMC	C1'-N1-C2	2.58	124.14	118.44
23	2K	8	4SU	C1'-N1-C2	2.57	122.21	117.59
25	14	2251	OMG	C2-N1-C6	2.56	129.76	125.11
1	1G	1207	2MG	C5-C6-N1	2.54	119.73	113.25
25	1H	1920	OMC	C1'-N1-C2	2.54	124.06	118.44
57	2L	33	OMC	O2-C2-N3	-2.54	118.33	122.33
25	14	2251	OMG	N2-C2-N1	2.54	122.11	116.76
1	13	1207	2MG	C1'-N9-C4	-2.53	119.02	126.49
25	14	2503	2MA	C6-C5-C4	2.51	120.60	117.18
25	1H	1917	PSU	O2-C2-N1	-2.50	120.21	122.79
1	13	1519	MA6	C4-N9-C8	2.50	108.36	105.74
1	13	966	M2G	O6-C6-C5	-2.49	119.95	126.53
25	14	1942	5MC	CM5-C5-C6	-2.49	119.48	122.85
25	14	2251	OMG	N1-C2-N3	-2.49	118.77	123.32
1	1G	1404	5MC	CM5-C5-C6	-2.48	119.49	122.85
25	1H	1920	OMC	O2-C2-N3	-2.48	118.42	122.33
25	14	1962	5MC	O2-C2-N3	-2.48	118.42	122.33
25	14	1917	PSU	O2-C2-N1	-2.48	120.23	122.79
25	1H	1939	5MU	O4-C4-N3	-2.47	115.47	120.11
1	1G	966	M2G	O6-C6-C5	-2.47	120.02	126.53
25	1H	2605	PSU	C6-N1-C2	-2.46	120.41	122.69
1	1G	1207	2MG	N9-C8-N7	-2.45	108.85	113.40
25	14	2503	2MA	CM2-C2-N3	2.45	120.80	117.13
25	1H	1917	PSU	C6-N1-C2	-2.44	120.43	122.69
25	14	2503	2MA	C4-N9-C8	2.44	108.30	105.74
23	2K	56	PSU	O2-C2-N1	-2.42	120.29	122.79

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	1G	1519	MA6	C4-C5-N7	-2.42	107.82	110.58
1	1G	516	PSU	O2-C2-N1	-2.40	120.31	122.79
1	1G	966	M2G	C5-C6-N1	2.40	119.37	113.25
25	14	1917	PSU	C6-C5-C4	2.39	119.79	118.17
1	13	1400	5MC	CM5-C5-C6	-2.39	119.61	122.85
57	2L	56	PSU	O2-C2-N1	-2.37	120.35	122.79
25	1H	1915	5MU	O4-C4-N3	-2.34	115.72	120.11
57	2L	56	PSU	C6-N1-C2	-2.32	120.53	122.69
1	1G	1207	2MG	O6-C6-C5	-2.30	120.47	126.53
25	14	1962	5MC	CM5-C5-C6	-2.29	119.75	122.85
57	2L	55	5MU	C1'-N1-C2	2.29	121.70	117.59
1	13	1518	MA6	C4-C5-N7	-2.28	107.97	110.58
25	14	2503	2MA	C5-N7-C8	2.28	107.03	103.45
1	13	1207	2MG	O6-C6-C5	-2.27	120.55	126.53
25	1H	2503	2MA	C4-N9-C1'	-2.26	121.34	126.63
1	13	1402	4OC	C6-C5-C4	2.26	119.72	117.00
1	1G	1498	UR3	C6-N1-C2	-2.25	119.96	121.80
25	1H	2251	OMG	C2-N1-C6	2.24	129.18	125.11
25	1H	1915	5MU	C1'-N1-C2	2.24	121.61	117.59
1	1G	516	PSU	O4'-C1'-C2'	2.23	108.24	105.15
23	2K	33	OMC	C1'-N1-C2	2.23	123.37	118.44
1	1G	1519	MA6	C4-N9-C8	2.23	108.08	105.74
25	14	1939	5MU	O4-C4-N3	-2.23	115.93	120.11
25	14	2605	PSU	C6-C5-C4	2.22	119.67	118.17
23	2K	56	PSU	C6-C5-C4	2.22	119.67	118.17
23	2K	56	PSU	C6-N1-C2	-2.21	120.64	122.69
1	13	1207	2MG	C8-N7-C5	2.21	108.19	104.26
1	1G	966	M2G	C1'-N9-C8	-2.19	120.51	126.73
1	13	516	PSU	O4'-C1'-C2'	2.18	108.17	105.15
1	1G	1400	5MC	CM5-C5-C6	-2.18	119.91	122.85
23	2K	33	OMC	O2-C2-N3	-2.17	118.91	122.33
25	1H	2503	2MA	C4-N9-C8	2.17	108.01	105.74
1	1G	967	5MC	CM5-C5-C6	-2.15	119.94	122.85
1	13	1207	2MG	N1-C2-N3	-2.14	120.08	123.68
12	3I	89	0TD	OD1-CG-CB	-2.13	117.99	122.44
57	2L	55	5MU	O4-C4-N3	-2.11	116.14	120.11
25	14	1917	PSU	C6-N1-C2	-2.11	120.74	122.69
1	13	1207	2MG	C1'-N9-C8	2.09	132.67	126.73
25	1H	1942	5MC	CM5-C5-C6	-2.08	120.04	122.85
1	1G	1207	2MG	N1-C2-N3	-2.06	120.21	123.68
1	13	1207	2MG	CM2-N2-C2	-2.04	119.27	123.65
25	1H	1920	OMC	O2-C2-N1	2.04	122.89	118.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	13	516	PSU	O2-C2-N1	-2.03	120.69	122.79
25	1H	1920	OMC	C1'-N1-C6	-2.03	116.44	120.78
57	2L	8	4SU	S4-C4-N3	-2.03	118.08	120.20
25	14	1920	OMC	C2'-C1'-N1	-2.03	110.39	114.24
25	14	2251	OMG	C2'-C1'-N9	-2.02	110.41	114.24
57	2L	33	OMC	C1'-N1-C6	-2.01	116.49	120.78
1	13	967	5MC	CM5-C5-C6	-2.00	120.14	122.85
1	1G	1407	5MC	O2-C2-N3	-2.00	119.17	122.33

There are no chirality outliers.

All (71) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
12	3I	89	0TD	SB-CB-CG-OD2
12	3I	89	0TD	SB-CB-CG-OD1
12	3A	89	0TD	CA-CB-SB-CSB
1	13	1402	4OC	O4'-C4'-C5'-O5'
1	1G	1402	4OC	O4'-C4'-C5'-O5'
1	1G	1518	MA6	C5-C6-N6-C10
1	13	1519	MA6	C5-C6-N6-C10
1	1G	1519	MA6	O4'-C4'-C5'-O5'
23	2K	21	H2U	O4'-C1'-N1-C2
23	2K	21	H2U	O4'-C1'-N1-C6
25	1H	2251	OMG	C1'-C2'-O2'-CM2
57	2L	47	7MG	C2'-C1'-N9-C8
1	13	1400	5MC	O4'-C4'-C5'-O5'
1	1G	1400	5MC	O4'-C4'-C5'-O5'
1	1G	1402	4OC	C3'-C4'-C5'-O5'
1	13	1519	MA6	O4'-C4'-C5'-O5'
1	1G	527	7MG	C3'-C4'-C5'-O5'
23	2K	21	H2U	C3'-C4'-C5'-O5'
23	2K	47	7MG	O4'-C4'-C5'-O5'
25	14	1920	OMC	C3'-C4'-C5'-O5'
25	14	1920	OMC	O4'-C4'-C5'-O5'
25	14	1962	5MC	O4'-C4'-C5'-O5'
1	13	1402	4OC	C3'-C4'-C5'-O5'
1	1G	1519	MA6	C3'-C4'-C5'-O5'
1	13	527	7MG	C3'-C4'-C5'-O5'
1	1G	1400	5MC	C3'-C4'-C5'-O5'
23	2K	21	H2U	O4'-C4'-C5'-O5'
1	1G	527	7MG	O4'-C4'-C5'-O5'
1	13	967	5MC	O4'-C4'-C5'-O5'

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Mol	Chain	Res	Type	Atoms
1	13	1400	5MC	C3'-C4'-C5'-O5'
1	13	1519	MA6	C3'-C4'-C5'-O5'
25	1H	2503	2MA	C3'-C4'-C5'-O5'
25	14	1962	5MC	C3'-C4'-C5'-O5'
25	1H	2503	2MA	O4'-C4'-C5'-O5'
57	2L	47	7MG	O4'-C4'-C5'-O5'
25	14	1915	5MU	C4'-C5'-O5'-P
12	3I	89	0TD	CG-CB-SB-CSB
25	1H	1962	5MC	O4'-C4'-C5'-O5'
25	1H	2503	2MA	C4'-C5'-O5'-P
23	2K	56	PSU	O4'-C1'-C5-C4
25	1H	1917	PSU	O4'-C1'-C5-C4
25	1H	2605	PSU	O4'-C1'-C5-C4
1	13	527	7MG	O4'-C4'-C5'-O5'
1	13	1207	2MG	C3'-C4'-C5'-O5'
1	1G	1518	MA6	N1-C6-N6-C10
1	13	1519	MA6	N1-C6-N6-C10
23	2K	47	7MG	C3'-C4'-C5'-O5'
1	1G	527	7MG	C4'-C5'-O5'-P
1	13	966	M2G	C4'-C5'-O5'-P
1	1G	1519	MA6	C4'-C5'-O5'-P
57	2L	8	4SU	C4'-C5'-O5'-P
57	2L	8	4SU	C3'-C4'-C5'-O5'
12	3A	89	0TD	SB-CB-CG-OD2
25	1H	1920	OMC	C2'-C1'-N1-C6
57	2L	47	7MG	C2'-C1'-N9-C4
23	2K	56	PSU	O4'-C1'-C5-C6
25	1H	2605	PSU	O4'-C1'-C5-C6
25	1H	1920	OMC	C2'-C1'-N1-C2
25	14	1915	5MU	C2'-C1'-N1-C6
57	2L	33	OMC	C2'-C1'-N1-C6
12	3A	89	0TD	CA-CB-CG-OD2
12	3A	89	0TD	CA-CB-CG-OD1
12	3A	89	0TD	CG-CB-SB-CSB
57	2L	8	4SU	O4'-C4'-C5'-O5'
57	2L	33	OMC	C3'-C2'-O2'-CM2
25	14	2552	OMU	O4'-C4'-C5'-O5'
23	2K	33	OMC	C2'-C1'-N1-C2
1	13	967	5MC	C3'-C4'-C5'-O5'
25	1H	1962	5MC	C3'-C4'-C5'-O5'
25	14	1915	5MU	C2'-C1'-N1-C2
57	2L	33	OMC	C2'-C1'-N1-C2

There are no ring outliers.

31 monomers are involved in 51 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
25	1H	1920	OMC	3	0
25	14	2251	OMG	3	0
23	2K	21	H2U	2	0
57	2L	33	OMC	1	0
25	1H	2552	OMU	1	0
1	1G	1519	MA6	1	0
1	13	1402	4OC	1	0
1	13	1407	5MC	1	0
25	1H	1939	5MU	1	0
25	14	1939	5MU	1	0
25	14	1962	5MC	1	0
12	3A	89	0TD	1	0
1	1G	1402	4OC	2	0
1	1G	1518	MA6	2	0
25	14	1911	PSU	1	0
57	2L	8	4SU	3	0
25	14	1920	OMC	1	0
25	14	2503	2MA	4	0
25	1H	2251	OMG	3	0
1	1G	1400	5MC	1	0
1	13	1498	UR3	1	0
1	1G	1498	UR3	2	0
1	1G	1404	5MC	2	0
25	14	1942	5MC	1	0
23	2K	55	5MU	2	0
23	2K	33	OMC	2	0
12	3I	89	0TD	1	0
1	1G	1207	2MG	3	0
1	13	1519	MA6	1	0
1	13	966	M2G	1	0
25	1H	1911	PSU	1	0

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

Of 1272 ligands modelled in this entry, 1270 are monoatomic - leaving 2 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	$\# Z > 2$
60	SF4	3E	302	-	0,12,12	-	-	-		
60	SF4	32	303	-	0,12,12	-	-	-		

In the following table, the Chirals column lists the number of chiral outliers, the number of chiral centers analysed, the number of these observed in the model and the number defined in the Chemical Component Dictionary. Similar counts are reported in the Torsion and Rings columns. '-' means no outliers of that kind were identified.

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
60	SF4	3E	302	-	-	-	0/6/5/5
60	SF4	32	303	-	-	-	0/6/5/5

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

There are no torsion outliers.

There are no ring outliers.

2 monomers are involved in 3 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
60	3E	302	SF4	1	0
60	32	303	SF4	2	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 ⓘ

In the following table, the column labelled ‘#RSRZ > 2’ contains the number (and percentage) of RSRZ outliers, followed by percent RSRZ outliers for the chain as percentile scores relative to all X-ray entries and entries of similar resolution. The OWAB column contains the minimum, median, 95th percentile and maximum values of the occupancy-weighted average B-factor per residue. The column labelled ‘Q < 0.9’ lists the number of (and percentage) of residues with an average occupancy less than 0.9.

Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
1	13	1494/1522 (98%)	-0.49	4 (0%) 90 76	40, 102, 169, 228	0
1	1G	1498/1522 (98%)	-0.53	1 (0%) 92 86	53, 104, 169, 213	0
2	12	236/256 (92%)	-0.20	0 100 100	114, 155, 171, 178	0
2	1E	237/256 (92%)	-0.26	0 100 100	108, 143, 167, 181	0
3	22	206/239 (86%)	0.09	2 (0%) 79 56	110, 131, 163, 182	0
3	2E	205/239 (85%)	-0.33	0 100 100	95, 124, 157, 172	0
4	32	208/209 (99%)	0.11	4 (1%) 66 41	78, 99, 120, 133	0
4	3E	208/209 (99%)	-0.03	2 (0%) 79 56	83, 112, 134, 144	0
5	42	154/162 (95%)	-0.09	0 100 100	86, 105, 134, 174	0
5	4E	151/162 (93%)	-0.20	0 100 100	77, 103, 130, 152	0
6	52	101/101 (100%)	-0.29	0 100 100	78, 103, 117, 125	0
6	5E	101/101 (100%)	0.16	2 (1%) 65 39	72, 104, 117, 140	0
7	62	149/156 (95%)	-0.22	0 100 100	110, 132, 151, 159	0
7	6E	155/156 (99%)	-0.14	0 100 100	99, 115, 145, 161	0
8	72	138/138 (100%)	-0.32	0 100 100	83, 107, 123, 128	0
8	7E	138/138 (100%)	-0.27	0 100 100	84, 106, 120, 126	0
9	82	127/128 (99%)	-0.31	0 100 100	104, 154, 173, 185	0
9	8E	124/128 (96%)	-0.38	0 100 100	84, 139, 155, 167	0
10	1A	99/105 (94%)	-0.05	0 100 100	111, 147, 169, 180	0
10	1I	99/105 (94%)	-0.26	0 100 100	93, 151, 166, 170	0
11	2A	119/129 (92%)	-0.28	0 100 100	87, 109, 142, 171	0
11	2I	117/129 (90%)	0.04	2 (1%) 69 43	63, 94, 129, 162	0
12	3A	124/132 (93%)	-0.13	0 100 100	71, 88, 113, 157	0
12	3I	124/132 (93%)	-0.21	0 100 100	62, 81, 111, 156	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
13	4A	118/126 (93%)	-0.01	0 100 100	108, 146, 166, 175	0
13	4I	119/126 (94%)	0.17	2 (1%) 69 43	81, 123, 139, 150	0
14	5A	60/61 (98%)	0.01	0 100 100	114, 131, 142, 149	0
14	5I	60/61 (98%)	-0.22	0 100 100	101, 118, 129, 135	0
15	6A	88/89 (98%)	-0.31	0 100 100	72, 98, 121, 128	0
15	6I	88/89 (98%)	-0.24	1 (1%) 78 54	65, 98, 120, 127	0
16	7A	84/88 (95%)	-0.30	0 100 100	75, 88, 106, 142	0
16	7I	83/88 (94%)	-0.16	0 100 100	95, 107, 130, 157	0
17	8A	99/105 (94%)	-0.23	0 100 100	77, 92, 108, 114	0
17	8I	100/105 (95%)	0.01	2 (2%) 65 39	75, 98, 113, 120	0
18	9A	71/88 (80%)	-0.30	0 100 100	86, 107, 135, 146	0
18	9I	70/88 (79%)	0.04	1 (1%) 73 48	78, 100, 127, 152	0
19	AA	83/93 (89%)	0.26	2 (2%) 59 35	120, 151, 166, 171	0
19	AI	86/93 (92%)	0.32	1 (1%) 76 52	102, 130, 148, 155	0
20	BA	103/106 (97%)	-0.22	0 100 100	82, 98, 125, 137	0
20	BI	101/106 (95%)	-0.21	0 100 100	92, 110, 130, 137	0
21	1B	25/27 (92%)	-0.22	0 100 100	123, 140, 157, 165	0
21	1F	25/27 (92%)	-0.43	0 100 100	104, 117, 136, 147	0
22	1K	65/77 (84%)	0.61	1 (1%) 72 47	88, 119, 149, 174	59 (90%)
22	3K	77/77 (100%)	-0.12	0 100 100	71, 198, 220, 236	0
22	3L	77/77 (100%)	-0.41	0 100 100	87, 216, 245, 255	0
23	2K	71/77 (92%)	-0.51	0 100 100	54, 91, 130, 147	0
24	4K	17/27 (62%)	-0.03	0 100 100	75, 132, 206, 207	0
24	4L	21/27 (77%)	-0.22	0 100 100	88, 171, 227, 232	0
25	14	2868/2917 (98%)	-0.59	2 (0%) 92 86	40, 79, 193, 238	0
25	1H	2868/2917 (98%)	-0.70	5 (0%) 91 82	27, 61, 173, 226	0
26	16	120/122 (98%)	-0.71	0 100 100	66, 93, 111, 146	0
26	1J	122/122 (100%)	-0.56	0 100 100	94, 132, 160, 193	0
27	7I	135/229 (58%)	-0.00	0 100 100	117, 180, 198, 211	0
28	11	274/276 (99%)	-0.26	0 100 100	29, 47, 70, 96	0
28	19	274/276 (99%)	-0.04	1 (0%) 88 72	39, 66, 86, 100	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
29	21	204/206 (99%)	-0.35	0 100 100	34, 74, 110, 127	0
29	29	204/206 (99%)	-0.21	0 100 100	47, 80, 111, 120	0
30	31	202/210 (96%)	-0.32	2 (0%) 79 56	31, 68, 110, 127	0
30	39	206/210 (98%)	-0.09	0 100 100	46, 91, 152, 177	0
31	41	181/182 (99%)	0.07	0 100 100	82, 103, 138, 149	0
31	49	181/182 (99%)	0.08	2 (1%) 78 54	118, 146, 161, 184	0
32	51	174/180 (96%)	-0.11	0 100 100	72, 98, 118, 136	0
32	59	174/180 (96%)	-0.02	0 100 100	115, 155, 176, 187	0
33	61	146/148 (98%)	0.15	1 (0%) 84 63	60, 109, 130, 140	0
33	69	146/148 (98%)	0.07	0 100 100	79, 120, 142, 156	0
34	38	139/173 (80%)	0.39	2 (1%) 73 48	156, 193, 210, 232	0
35	15	138/140 (98%)	-0.09	0 100 100	68, 102, 129, 140	0
35	58	138/140 (98%)	-0.18	0 100 100	56, 76, 118, 127	0
36	25	122/122 (100%)	0.07	3 (2%) 58 35	53, 75, 93, 103	0
36	68	122/122 (100%)	-0.33	0 100 100	45, 65, 85, 105	0
37	35	147/150 (98%)	-0.35	1 (0%) 84 63	51, 96, 127, 150	0
37	78	148/150 (98%)	-0.41	0 100 100	32, 72, 102, 123	0
38	45	140/141 (99%)	-0.14	0 100 100	72, 99, 127, 138	0
38	88	141/141 (100%)	-0.13	0 100 100	46, 73, 106, 128	0
39	55	118/118 (100%)	-0.26	0 100 100	48, 71, 99, 122	0
39	98	118/118 (100%)	-0.14	1 (0%) 82 60	40, 69, 94, 116	0
40	65	111/112 (99%)	-0.41	0 100 100	103, 125, 139, 148	0
40	A8	111/112 (99%)	-0.14	0 100 100	75, 93, 115, 127	0
41	75	137/146 (93%)	-0.12	2 (1%) 72 47	68, 84, 137, 157	0
41	B8	137/146 (93%)	-0.26	0 100 100	61, 81, 134, 164	0
42	85	117/118 (99%)	-0.25	0 100 100	57, 90, 147, 158	0
42	C8	117/118 (99%)	-0.36	0 100 100	37, 63, 101, 120	0
43	95	101/101 (100%)	0.05	0 100 100	53, 123, 143, 159	0
43	D8	100/101 (99%)	-0.10	0 100 100	41, 90, 118, 134	0
44	A5	112/113 (99%)	-0.28	1 (0%) 81 58	52, 66, 100, 144	0
44	E8	113/113 (100%)	-0.44	0 100 100	40, 55, 97, 140	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
45	B5	93/96 (96%)	-0.09	0 100 100	58, 77, 106, 124	0
45	F8	95/96 (98%)	-0.40	0 100 100	42, 56, 85, 99	0
46	C5	109/110 (99%)	0.30	3 (2%) 55 32	73, 102, 120, 128	0
46	G8	107/110 (97%)	-0.19	1 (0%) 81 58	62, 87, 135, 148	0
47	D5	176/206 (85%)	-0.11	0 100 100	117, 145, 178, 186	0
47	H8	174/206 (84%)	0.29	3 (1%) 69 43	79, 118, 177, 192	0
48	E5	79/85 (92%)	-0.66	0 100 100	70, 90, 113, 160	0
48	I8	77/85 (90%)	-0.41	1 (1%) 75 50	49, 67, 93, 114	0
49	F5	92/96 (95%)	0.06	2 (2%) 62 37	54, 79, 115, 130	0
49	J8	96/96 (100%)	0.02	2 (2%) 63 38	41, 66, 117, 150	0
50	G5	69/72 (95%)	-0.11	0 100 100	69, 95, 121, 145	0
50	K8	71/72 (98%)	-0.14	2 (2%) 55 32	45, 69, 89, 110	0
51	H5	59/60 (98%)	-0.24	0 100 100	70, 97, 146, 169	0
51	L8	59/60 (98%)	-0.34	0 100 100	49, 71, 105, 114	0
52	I5	71/71 (100%)	0.23	1 (1%) 73 48	147, 178, 194, 204	0
52	M8	66/71 (92%)	0.49	3 (4%) 38 20	114, 141, 171, 192	0
53	J5	56/60 (93%)	0.04	0 100 100	50, 76, 143, 153	0
53	N8	56/60 (93%)	-0.19	1 (1%) 67 42	35, 75, 145, 151	0
54	K5	48/54 (88%)	0.01	0 100 100	128, 165, 181, 188	0
54	O8	49/54 (90%)	0.39	2 (4%) 41 22	106, 140, 158, 162	0
55	L5	48/49 (97%)	-0.38	0 100 100	41, 51, 82, 94	0
55	P8	48/49 (97%)	-0.62	0 100 100	26, 35, 71, 76	0
56	M5	64/65 (98%)	-0.64	0 100 100	59, 73, 98, 119	0
56	Q8	64/65 (98%)	-0.59	0 100 100	38, 53, 74, 95	0
57	2L	72/77 (93%)	-0.55	1 (1%) 73 48	67, 113, 154, 178	0
All	All	21207/22013 (96%)	-0.33	72 (0%) 90 76	26, 95, 171, 255	59 (0%)

All (72) RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
54	O8	7	ILE	3.7
13	4I	119	GLY	3.6
30	31	131	GLY	3.5

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Mol	Chain	Res	Type	RSRZ
25	1H	1538	G	3.4
17	8I	98	LEU	3.3
11	2I	129	SER	3.0
48	I8	7	LEU	2.9
41	75	36	GLU	2.8
34	38	116	ILE	2.7
46	C5	55	TYR	2.7
33	61	122	GLU	2.7
36	25	12	ASP	2.6
49	J8	1	MET	2.6
52	M8	3	GLU	2.5
25	1H	2805	G	2.5
50	K8	72	ALA	2.5
49	J8	84	GLY	2.5
31	49	20	ILE	2.5
47	H8	116	VAL	2.5
3	22	80	GLY	2.4
22	1K	5	G	2.4
19	AA	49	ILE	2.4
37	35	52	GLU	2.4
25	1H	1830	C	2.4
36	25	57	VAL	2.4
4	32	37	PRO	2.4
47	H8	120	ILE	2.4
4	32	161	ASN	2.3
25	14	546	C	2.3
52	M8	64	GLY	2.3
4	32	164	ALA	2.3
4	3E	24	GLU	2.3
6	5E	55	ASP	2.3
15	6I	88	ARG	2.3
47	H8	60	GLU	2.3
34	38	63	LEU	2.3
49	F5	79	GLY	2.3
52	M8	21	VAL	2.3
1	13	1511	G	2.2
25	14	1176	G	2.2
50	K8	70	GLN	2.2
46	C5	29	GLU	2.2
1	13	344	A	2.2
25	1H	2402	C	2.2
30	31	72	ARG	2.2

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Mol	Chain	Res	Type	RSRZ
28	19	5	LYS	2.2
11	2I	81	ASP	2.2
31	49	48	GLU	2.2
18	9I	88	LYS	2.2
13	4I	102	ARG	2.1
1	13	1394	A	2.1
19	AI	43	GLU	2.1
25	1H	1963	U	2.1
36	25	91	LEU	2.1
57	2L	1	C	2.1
4	32	166	LYS	2.1
44	A5	94	ASP	2.1
39	98	7	GLY	2.1
1	1G	693	G	2.1
4	3E	90	GLY	2.1
46	G8	108	THR	2.1
41	75	106	SER	2.1
19	AA	41	VAL	2.1
3	22	81	GLY	2.0
17	8I	96	GLU	2.0
49	F5	91	LYS	2.0
53	N8	53	ALA	2.0
54	O8	8	LYS	2.0
1	13	913	A	2.0
46	C5	58	GLY	2.0
6	5E	50	TYR	2.0
52	I5	40	HIS	2.0

6.2 Non-standard residues in protein, DNA, RNA chains [i](#)

In the following table, the Atoms column lists the number of modelled atoms in the group and the number defined in the chemical component dictionary. The B-factors column lists the minimum, median, 95th percentile and maximum values of B factors of atoms in the group. The column labelled 'Q< 0.9' lists the number of atoms with occupancy less than 0.9.

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(Å ²)	Q<0.9
25	5MU	14	1915	21/22	0.80	0.08	116,128,139,153	0
23	5MU	2K	55	21/22	0.83	0.08	103,113,125,145	0
57	5MU	2L	55	21/22	0.84	0.06	119,129,134,137	0
57	4SU	2L	8	20/21	0.85	0.07	120,131,140,141	0
23	H2U	2K	21	20/21	0.85	0.11	117,140,147,150	0
23	PSU	2K	56	20/21	0.86	0.07	92,104,120,124	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	PSU	2L	56	20/21	0.87	0.06	123,128,134,135	0
1	5MC	1G	967	21/22	0.88	0.07	92,107,118,120	0
57	7MG	2L	47	24/25	0.89	0.06	114,141,151,162	0
25	PSU	14	1911	20/21	0.89	0.07	86,94,118,121	0
25	5MU	1H	1915	21/22	0.89	0.08	75,91,102,116	0
1	2MG	13	1207	24/25	0.90	0.11	100,112,118,119	0
1	7MG	13	527	24/25	0.90	0.10	71,80,86,92	0
23	7MG	2K	47	24/25	0.90	0.08	96,105,114,126	0
1	PSU	1G	516	20/21	0.91	0.06	98,102,108,110	0
12	0TD	3I	89	10/11	0.91	0.10	81,87,92,107	0
1	2MG	1G	1207	24/25	0.91	0.06	115,120,125,127	0
1	M2G	1G	966	25/26	0.91	0.07	87,107,115,124	0
23	OMC	2K	33	21/22	0.91	0.08	77,82,89,101	0
1	5MC	1G	1404	21/22	0.92	0.10	77,85,89,91	0
57	OMC	2L	33	21/22	0.92	0.08	91,100,107,111	0
1	5MC	13	967	21/22	0.92	0.09	71,82,94,99	0
1	5MC	1G	1400	21/22	0.92	0.07	81,92,98,102	0
25	PSU	14	1917	20/21	0.92	0.08	87,104,109,110	0
25	OMC	14	1920	21/22	0.93	0.08	73,91,99,102	0
1	5MC	1G	1407	21/22	0.93	0.10	75,80,85,98	0
12	0TD	3A	89	10/11	0.93	0.11	89,95,100,105	0
25	PSU	1H	1917	20/21	0.94	0.08	65,76,90,91	0
1	M2G	13	966	25/26	0.94	0.08	68,77,90,94	0
1	4OC	1G	1402	22/23	0.94	0.07	71,84,91,99	0
25	5MC	14	1962	21/22	0.94	0.08	59,67,76,83	0
1	5MC	13	1404	21/22	0.94	0.10	54,59,69,79	0
1	PSU	13	516	20/21	0.94	0.05	84,95,98,98	0
1	7MG	1G	527	24/25	0.94	0.09	88,99,101,102	0
1	UR3	1G	1498	21/22	0.94	0.10	65,76,83,86	0
1	5MC	13	1400	21/22	0.94	0.08	58,74,79,86	0
25	5MC	1H	1942	21/22	0.95	0.09	39,50,56,61	0
25	5MC	14	1942	21/22	0.95	0.09	62,71,82,90	0
1	MA6	1G	1519	24/25	0.95	0.12	64,79,84,86	0
25	2MA	14	2503	23/24	0.95	0.10	39,50,63,79	0
1	4OC	13	1402	22/23	0.95	0.09	61,71,79,84	0
1	UR3	13	1498	21/22	0.95	0.11	44,62,69,73	0
25	OMC	1H	1920	21/22	0.95	0.08	60,66,70,72	0
1	5MC	13	1407	21/22	0.95	0.09	53,59,65,67	0
25	5MU	1H	1939	21/22	0.95	0.12	36,41,48,50	0
25	OMG	14	2251	24/25	0.96	0.07	51,57,64,69	0
25	2MA	1H	2503	23/24	0.96	0.09	30,38,48,59	0
1	MA6	13	1518	24/25	0.96	0.10	50,57,61,62	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
25	OMU	1H	2552	21/22	0.96	0.08	37,45,51,53	0
25	PSU	1H	2605	20/21	0.96	0.09	35,40,48,53	0
25	PSU	14	2605	20/21	0.96	0.08	42,52,74,77	0
23	4SU	2K	8	20/21	0.96	0.06	85,91,97,101	0
1	MA6	1G	1518	24/25	0.96	0.09	68,78,82,84	0
25	PSU	1H	1911	20/21	0.96	0.08	50,63,72,74	0
25	5MC	1H	1962	21/22	0.96	0.08	50,57,61,66	0
1	MA6	13	1519	24/25	0.96	0.12	51,56,62,64	0
25	OMU	14	2552	21/22	0.97	0.08	48,55,62,67	0
25	5MU	14	1939	21/22	0.97	0.08	44,52,57,62	0
25	OMG	1H	2251	24/25	0.98	0.05	36,43,49,53	0

6.3 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

6.4 Ligands [i](#)

In the following table, the Atoms column lists the number of modelled atoms in the group and the number defined in the chemical component dictionary. The B-factors column lists the minimum, median, 95th percentile and maximum values of B factors of atoms in the group. The column labelled 'Q< 0.9' lists the number of atoms with occupancy less than 0.9.

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
58	K	1H	3072	1/1	0.40	0.12	100,100,100,100	0
59	MG	1H	3326[A]	1/1	0.47	0.37	58,58,58,58	1
59	MG	1H	3326[B]	1/1	0.47	0.37	55,55,55,55	1
59	MG	14	3174	1/1	0.51	0.24	78,78,78,78	0
58	K	16	202	1/1	0.56	0.12	116,116,116,116	0
59	MG	1G	1656	1/1	0.57	0.25	100,100,100,100	0
58	K	1H	3010	1/1	0.57	0.11	93,93,93,93	0
59	MG	1H	3239	1/1	0.59	0.26	83,83,83,83	0
59	MG	13	1682	1/1	0.61	0.19	101,101,101,101	0
59	MG	1G	1665	1/1	0.62	0.23	84,84,84,84	0
59	MG	1H	3140	1/1	0.63	0.26	82,82,82,82	0
59	MG	14	3211	1/1	0.63	0.23	64,64,64,64	0
58	K	13	1635	1/1	0.64	0.18	104,104,104,104	0
59	MG	13	1710	1/1	0.64	0.19	97,97,97,97	0
58	K	14	3088	1/1	0.64	0.13	101,101,101,101	0
59	MG	1G	1648	1/1	0.64	0.30	83,83,83,83	0
59	MG	1H	3231	1/1	0.66	0.21	67,67,67,67	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	1G	1628	1/1	0.67	0.23	92,92,92,92	0
59	MG	13	1688	1/1	0.67	0.20	83,83,83,83	0
58	K	1H	3016	1/1	0.68	0.09	114,114,114,114	0
58	K	13	1636	1/1	0.69	0.09	136,136,136,136	0
59	MG	16	205	1/1	0.69	0.22	84,84,84,84	0
59	MG	14	3254	1/1	0.69	0.24	71,71,71,71	0
58	K	5E	201	1/1	0.70	0.17	113,113,113,113	0
58	K	1H	3014	1/1	0.70	0.20	84,84,84,84	0
58	K	1G	1604	1/1	0.71	0.11	111,111,111,111	0
59	MG	14	3246	1/1	0.71	0.15	76,76,76,76	0
59	MG	1G	1661	1/1	0.71	0.17	114,114,114,114	0
58	K	1H	3052	1/1	0.72	0.12	127,127,127,127	0
59	MG	1G	1635	1/1	0.72	0.21	80,80,80,80	0
58	K	3A	201	1/1	0.73	0.12	93,93,93,93	0
59	MG	13	1693	1/1	0.73	0.18	78,78,78,78	0
59	MG	14	3196	1/1	0.73	0.20	90,90,90,90	0
59	MG	14	3202	1/1	0.73	0.13	84,84,84,84	0
58	K	41	202	1/1	0.73	0.09	128,128,128,128	0
58	K	1H	3044	1/1	0.73	0.07	82,82,82,82	0
59	MG	1H	3158	1/1	0.73	0.18	89,89,89,89	0
58	K	14	3042	1/1	0.74	0.09	83,83,83,83	0
58	K	14	3069	1/1	0.74	0.08	89,89,89,89	0
58	K	1G	1605	1/1	0.74	0.10	92,92,92,92	0
59	MG	1H	3221	1/1	0.74	0.28	77,77,77,77	0
59	MG	1G	1662	1/1	0.74	0.17	74,74,74,74	0
59	MG	13	1695	1/1	0.74	0.23	76,76,76,76	0
59	MG	13	1696	1/1	0.75	0.16	80,80,80,80	0
58	K	13	1626	1/1	0.75	0.12	99,99,99,99	0
58	K	14	3005	1/1	0.75	0.12	91,91,91,91	0
59	MG	1H	3310	1/1	0.75	0.23	78,78,78,78	0
58	K	1G	1606	1/1	0.75	0.08	102,102,102,102	0
59	MG	1H	3174	1/1	0.76	0.10	71,71,71,71	0
59	MG	1G	1649	1/1	0.76	0.17	85,85,85,85	0
59	MG	14	3197	1/1	0.76	0.17	60,60,60,60	0
58	K	8I	201	1/1	0.76	0.09	104,104,104,104	0
59	MG	13	1686	1/1	0.76	0.22	95,95,95,95	0
59	MG	14	3218	1/1	0.76	0.17	66,66,66,66	0
58	K	1H	3079	1/1	0.76	0.08	114,114,114,114	0
59	MG	1H	3159	1/1	0.76	0.18	68,68,68,68	0
59	MG	1H	3284	1/1	0.77	0.23	66,66,66,66	0
58	K	52	201	1/1	0.77	0.12	116,116,116,116	0
58	K	1G	1601	1/1	0.77	0.11	102,102,102,102	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
58	K	13	1613	1/1	0.78	0.08	100,100,100,100	0
59	MG	1G	1627	1/1	0.78	0.21	88,88,88,88	0
59	MG	13	1652	1/1	0.78	0.23	95,95,95,95	0
58	K	14	3055	1/1	0.78	0.07	116,116,116,116	0
58	K	13	1610	1/1	0.78	0.09	114,114,114,114	0
58	K	14	3075	1/1	0.78	0.09	116,116,116,116	0
59	MG	14	3137	1/1	0.79	0.11	83,83,83,83	0
58	K	14	3049	1/1	0.79	0.11	103,103,103,103	0
59	MG	1H	3244	1/1	0.79	0.12	52,52,52,52	0
59	MG	1G	1652	1/1	0.79	0.14	94,94,94,94	0
59	MG	16	207	1/1	0.79	0.23	88,88,88,88	0
59	MG	1H	3213	1/1	0.79	0.11	73,73,73,73	0
59	MG	3E	301	1/1	0.79	0.23	84,84,84,84	0
59	MG	1H	3139	1/1	0.79	0.25	76,76,76,76	0
59	MG	1G	1666	1/1	0.79	0.19	77,77,77,77	0
59	MG	1H	3207	1/1	0.80	0.09	56,56,56,56	0
59	MG	13	1709	1/1	0.80	0.19	85,85,85,85	0
58	K	14	3058	1/1	0.80	0.20	84,84,84,84	0
59	MG	1G	1663	1/1	0.80	0.24	73,73,73,73	0
59	MG	1G	1664	1/1	0.80	0.25	73,73,73,73	0
59	MG	13	1687	1/1	0.80	0.17	75,75,75,75	0
58	K	13	1606	1/1	0.80	0.09	87,87,87,87	0
59	MG	1H	3188	1/1	0.80	0.20	84,84,84,84	0
59	MG	14	3267	1/1	0.80	0.21	92,92,92,92	0
59	MG	1H	3285	1/1	0.81	0.27	66,66,66,66	0
59	MG	1H	3304	1/1	0.81	0.22	69,69,69,69	0
58	K	1G	1615	1/1	0.81	0.14	106,106,106,106	0
59	MG	13	1685	1/1	0.81	0.26	85,85,85,85	0
58	K	1H	3099	1/1	0.81	0.11	81,81,81,81	0
59	MG	1J	202	1/1	0.81	0.15	69,69,69,69	0
59	MG	13	1645	1/1	0.82	0.10	71,71,71,71	0
58	K	13	1618	1/1	0.82	0.11	90,90,90,90	0
59	MG	13	1678	1/1	0.82	0.18	79,79,79,79	0
59	MG	13	1698	1/1	0.82	0.25	67,67,67,67	0
59	MG	13	1701	1/1	0.82	0.25	72,72,72,72	0
58	K	14	3008	1/1	0.82	0.08	88,88,88,88	0
58	K	14	3033	1/1	0.82	0.14	84,84,84,84	0
59	MG	1H	3238	1/1	0.82	0.27	82,82,82,82	0
58	K	13	1620	1/1	0.82	0.06	115,115,115,115	0
59	MG	BI	201	1/1	0.82	0.11	108,108,108,108	0
59	MG	14	3220	1/1	0.82	0.07	53,53,53,53	0
59	MG	1H	3133	1/1	0.82	0.15	55,55,55,55	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
58	K	14	3076	1/1	0.82	0.07	89,89,89,89	0
58	K	13	1607	1/1	0.82	0.10	90,90,90,90	0
59	MG	13	1691	1/1	0.82	0.18	88,88,88,88	0
58	K	1H	3038	1/1	0.83	0.07	86,86,86,86	0
59	MG	1H	3325	1/1	0.83	0.26	65,65,65,65	0
59	MG	13	1663	1/1	0.83	0.24	68,68,68,68	0
59	MG	13	1676	1/1	0.83	0.14	53,53,53,53	0
58	K	14	3026	1/1	0.83	0.10	109,109,109,109	0
59	MG	14	3209	1/1	0.83	0.20	72,72,72,72	0
59	MG	13	1699	1/1	0.83	0.23	68,68,68,68	0
58	K	13	1633	1/1	0.83	0.06	108,108,108,108	0
59	MG	13	1707	1/1	0.83	0.15	65,65,65,65	0
59	MG	14	3233	1/1	0.83	0.14	80,80,80,80	0
59	MG	1G	1630	1/1	0.83	0.18	64,64,64,64	0
59	MG	1H	3309	1/1	0.83	0.13	56,56,56,56	0
59	MG	14	3265	1/1	0.83	0.26	67,67,67,67	0
59	MG	14	3096	1/1	0.83	0.12	59,59,59,59	0
59	MG	14	3131	1/1	0.83	0.24	66,66,66,66	0
59	MG	14	3136	1/1	0.84	0.16	77,77,77,77	0
59	MG	13	1656	1/1	0.84	0.21	73,73,73,73	0
59	MG	14	3142	1/1	0.84	0.18	62,62,62,62	0
58	K	14	3081	1/1	0.84	0.10	89,89,89,89	0
59	MG	14	3185	1/1	0.84	0.17	76,76,76,76	0
59	MG	14	3192	1/1	0.84	0.14	63,63,63,63	0
59	MG	1H	3316	1/1	0.84	0.21	59,59,59,59	0
59	MG	7I	101	1/1	0.84	0.16	124,124,124,124	0
59	MG	1G	1660	1/1	0.84	0.22	79,79,79,79	0
58	K	14	3083	1/1	0.84	0.08	119,119,119,119	0
58	K	13	1637	1/1	0.84	0.08	123,123,123,123	0
58	K	14	3092	1/1	0.84	0.08	83,83,83,83	0
58	K	29	302	1/1	0.84	0.07	89,89,89,89	0
58	K	14	3039	1/1	0.84	0.09	94,94,94,94	0
58	K	13	1634	1/1	0.84	0.08	111,111,111,111	0
59	MG	14	3250	1/1	0.84	0.25	84,84,84,84	0
59	MG	14	3252	1/1	0.84	0.23	75,75,75,75	0
59	MG	13	1655	1/1	0.84	0.12	80,80,80,80	0
59	MG	14	3103	1/1	0.84	0.17	82,82,82,82	0
59	MG	14	3124	1/1	0.84	0.06	63,63,63,63	0
59	MG	14	3268	1/1	0.84	0.12	79,79,79,79	0
59	MG	1H	3180	1/1	0.84	0.17	72,72,72,72	0
59	MG	1G	1676	1/1	0.85	0.19	81,81,81,81	0
59	MG	1H	3317	1/1	0.85	0.25	63,63,63,63	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	1H	3323	1/1	0.85	0.29	73,73,73,73	0
59	MG	14	3115	1/1	0.85	0.19	46,46,46,46	0
59	MG	13	1674	1/1	0.85	0.11	92,92,92,92	0
58	K	BA	201	1/1	0.85	0.12	96,96,96,96	0
58	K	1H	3117	1/1	0.85	0.11	85,85,85,85	0
58	K	1H	3009	1/1	0.85	0.13	97,97,97,97	0
59	MG	1H	3223	1/1	0.85	0.12	64,64,64,64	0
59	MG	1H	3229	1/1	0.85	0.08	51,51,51,51	0
58	K	14	3019	1/1	0.85	0.16	81,81,81,81	0
58	K	13	1627	1/1	0.85	0.07	91,91,91,91	0
58	K	13	1630	1/1	0.85	0.08	82,82,82,82	0
59	MG	1G	1646	1/1	0.85	0.16	107,107,107,107	0
58	K	13	1605	1/1	0.85	0.07	94,94,94,94	0
59	MG	1H	3252	1/1	0.85	0.18	72,72,72,72	0
59	MG	1G	1651	1/1	0.85	0.24	63,63,63,63	0
59	MG	1H	3276	1/1	0.85	0.23	63,63,63,63	0
58	K	1H	3020	1/1	0.85	0.12	105,105,105,105	0
59	MG	1G	1658	1/1	0.85	0.21	76,76,76,76	0
59	MG	14	3243	1/1	0.85	0.26	74,74,74,74	0
58	K	1H	3082	1/1	0.85	0.05	80,80,80,80	0
59	MG	1H	3289	1/1	0.85	0.16	75,75,75,75	0
58	K	1H	3033	1/1	0.85	0.10	94,94,94,94	0
59	MG	14	3253	1/1	0.85	0.21	52,52,52,52	0
58	K	1H	3110	1/1	0.85	0.10	111,111,111,111	0
58	K	1H	3112	1/1	0.85	0.09	87,87,87,87	0
59	MG	14	3266	1/1	0.85	0.18	74,74,74,74	0
59	MG	1H	3312	1/1	0.85	0.14	70,70,70,70	0
59	MG	13	1671	1/1	0.85	0.17	65,65,65,65	0
59	MG	1G	1672	1/1	0.85	0.20	63,63,63,63	0
59	MG	1G	1643	1/1	0.86	0.08	56,56,56,56	0
59	MG	13	1697	1/1	0.86	0.14	70,70,70,70	0
58	K	1G	1617	1/1	0.86	0.07	94,94,94,94	0
58	K	1H	3091	1/1	0.86	0.14	62,62,62,62	0
59	MG	1H	3308	1/1	0.86	0.17	55,55,55,55	0
59	MG	13	1679	1/1	0.86	0.12	89,89,89,89	0
58	K	14	3027	1/1	0.86	0.10	97,97,97,97	0
59	MG	14	3199	1/1	0.86	0.10	105,105,105,105	0
59	MG	14	3201	1/1	0.86	0.18	68,68,68,68	0
58	K	13	1628	1/1	0.86	0.07	103,103,103,103	0
59	MG	1H	3215	1/1	0.86	0.28	65,65,65,65	0
59	MG	13	1653	1/1	0.86	0.16	74,74,74,74	0
59	MG	1H	3321	1/1	0.86	0.16	55,55,55,55	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
58	K	1H	3101	1/1	0.86	0.08	60,60,60,60	0
59	MG	1H	3227	1/1	0.86	0.20	58,58,58,58	0
59	MG	8E	201	1/1	0.86	0.16	49,49,49,49	0
59	MG	4I	201	1/1	0.86	0.25	84,84,84,84	0
59	MG	1H	3327	1/1	0.86	0.28	72,72,72,72	0
58	K	4I	201	1/1	0.86	0.11	114,114,114,114	0
59	MG	14	3095	1/1	0.86	0.19	57,57,57,57	0
59	MG	13	1690	1/1	0.86	0.20	77,77,77,77	0
58	K	14	3047	1/1	0.86	0.08	97,97,97,97	0
59	MG	13	1665	1/1	0.86	0.15	50,50,50,50	0
58	K	1H	3018	1/1	0.86	0.07	93,93,93,93	0
59	MG	1G	1634	1/1	0.86	0.17	110,110,110,110	0
59	MG	14	3269	1/1	0.86	0.17	79,79,79,79	0
58	K	14	3091	1/1	0.86	0.10	103,103,103,103	0
59	MG	14	3190	1/1	0.87	0.14	81,81,81,81	0
59	MG	1H	3236	1/1	0.87	0.12	54,54,54,54	0
58	K	1H	3113[B]	1/1	0.87	0.17	39,39,39,39	1
58	K	14	3036	1/1	0.87	0.08	85,85,85,85	0
59	MG	1H	3243	1/1	0.87	0.22	94,94,94,94	0
59	MG	1H	3336	1/1	0.87	0.15	54,54,54,54	0
58	K	13	1614	1/1	0.87	0.12	103,103,103,103	0
59	MG	14	3203	1/1	0.87	0.13	85,85,85,85	0
58	K	1H	3120	1/1	0.87	0.11	79,79,79,79	0
59	MG	1G	1671	1/1	0.87	0.26	79,79,79,79	0
59	MG	1H	3170	1/1	0.87	0.13	47,47,47,47	0
58	K	1H	3096	1/1	0.87	0.14	75,75,75,75	0
58	K	1H	3069	1/1	0.87	0.06	74,74,74,74	0
58	K	14	3052	1/1	0.87	0.06	105,105,105,105	0
59	MG	1H	3203	1/1	0.87	0.12	52,52,52,52	0
58	K	1H	3019	1/1	0.87	0.08	78,78,78,78	0
58	K	1H	3102	1/1	0.87	0.06	94,94,94,94	0
58	K	1H	3040	1/1	0.87	0.18	111,111,111,111	0
58	K	14	3070	1/1	0.87	0.08	95,95,95,95	0
58	K	13	1604	1/1	0.87	0.07	90,90,90,90	0
58	K	1H	3113[A]	1/1	0.87	0.17	45,45,45,45	1
59	MG	14	3155	1/1	0.87	0.15	57,57,57,57	0
59	MG	14	3162	1/1	0.87	0.20	58,58,58,58	0
59	MG	13	1670	1/1	0.87	0.17	72,72,72,72	0
59	MG	14	3270	1/1	0.87	0.12	57,57,57,57	0
59	MG	1H	3138	1/1	0.87	0.22	66,66,66,66	0
59	MG	1H	3278	1/1	0.88	0.22	68,68,68,68	0
59	MG	1H	3282	1/1	0.88	0.12	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	1H	3184	1/1	0.88	0.19	65,65,65,65	0
59	MG	1G	1637	1/1	0.88	0.17	77,77,77,77	0
59	MG	14	3150	1/1	0.88	0.17	63,63,63,63	0
59	MG	14	3154	1/1	0.88	0.19	72,72,72,72	0
59	MG	1G	1639	1/1	0.88	0.19	55,55,55,55	0
58	K	1G	1622	1/1	0.88	0.07	118,118,118,118	0
59	MG	14	3167	1/1	0.88	0.19	60,60,60,60	0
58	K	39	301	1/1	0.88	0.08	107,107,107,107	0
59	MG	14	3179	1/1	0.88	0.20	66,66,66,66	0
59	MG	14	3183	1/1	0.88	0.17	63,63,63,63	0
59	MG	14	3184	1/1	0.88	0.19	73,73,73,73	0
58	K	1H	3034	1/1	0.88	0.07	83,83,83,83	0
59	MG	1H	3211	1/1	0.88	0.21	47,47,47,47	0
59	MG	1G	1650	1/1	0.88	0.17	65,65,65,65	0
58	K	31	302	1/1	0.88	0.06	90,90,90,90	0
58	K	1H	3092	1/1	0.88	0.07	70,70,70,70	0
59	MG	14	3198	1/1	0.88	0.18	63,63,63,63	0
59	MG	1G	1655	1/1	0.88	0.20	67,67,67,67	0
58	K	14	3074	1/1	0.88	0.08	94,94,94,94	0
59	MG	1H	3315	1/1	0.88	0.30	70,70,70,70	0
59	MG	13	1689	1/1	0.88	0.09	80,80,80,80	0
58	K	14	3001	1/1	0.88	0.10	87,87,87,87	0
59	MG	1H	3318	1/1	0.88	0.19	85,85,85,85	0
58	K	14	3041	1/1	0.88	0.12	54,54,54,54	0
58	K	1G	1609	1/1	0.88	0.12	113,113,113,113	0
59	MG	14	3225	1/1	0.88	0.09	62,62,62,62	0
59	MG	14	3226	1/1	0.88	0.10	49,49,49,49	0
59	MG	1H	3235	1/1	0.88	0.14	62,62,62,62	0
59	MG	14	3235	1/1	0.88	0.16	61,61,61,61	0
58	K	14	3082	1/1	0.88	0.14	76,76,76,76	0
59	MG	1G	1667	1/1	0.88	0.16	60,60,60,60	0
59	MG	1G	1670	1/1	0.88	0.19	60,60,60,60	0
58	K	1H	3025	1/1	0.88	0.05	100,100,100,100	0
58	K	14	3017	1/1	0.88	0.12	107,107,107,107	0
59	MG	1G	1674	1/1	0.88	0.18	50,50,50,50	0
59	MG	14	3263	1/1	0.88	0.23	66,66,66,66	0
58	K	14	3051	1/1	0.88	0.08	93,93,93,93	0
59	MG	2L	102	1/1	0.88	0.11	132,132,132,132	0
59	MG	1H	3389	1/1	0.88	0.12	38,38,38,38	0
59	MG	13	1677	1/1	0.88	0.11	63,63,63,63	0
58	K	1H	3012	1/1	0.88	0.09	80,80,80,80	0
59	MG	1H	3259	1/1	0.88	0.14	59,59,59,59	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	13	1702	1/1	0.88	0.22	62,62,62,62	0
59	MG	14	3160	1/1	0.89	0.13	73,73,73,73	0
59	MG	13	1661	1/1	0.89	0.12	42,42,42,42	0
59	MG	13	1662	1/1	0.89	0.17	67,67,67,67	0
59	MG	14	3171	1/1	0.89	0.07	71,71,71,71	0
58	K	14	3079	1/1	0.89	0.09	112,112,112,112	0
58	K	14	3003	1/1	0.89	0.08	86,86,86,86	0
59	MG	13	1669	1/1	0.89	0.16	53,53,53,53	0
58	K	1H	3001	1/1	0.89	0.06	88,88,88,88	0
59	MG	13	1704	1/1	0.89	0.13	53,53,53,53	0
58	K	1H	3097	1/1	0.89	0.08	69,69,69,69	0
58	K	14	3084	1/1	0.89	0.08	75,75,75,75	0
59	MG	1H	3319	1/1	0.89	0.14	55,55,55,55	0
58	K	14	3085	1/1	0.89	0.05	94,94,94,94	0
58	K	14	3086	1/1	0.89	0.14	97,97,97,97	0
59	MG	1H	3324	1/1	0.89	0.20	59,59,59,59	0
58	K	1H	3063	1/1	0.89	0.07	55,55,55,55	0
58	K	14	3089	1/1	0.89	0.07	97,97,97,97	0
59	MG	1G	1668	1/1	0.89	0.24	55,55,55,55	0
58	K	13	1608	1/1	0.89	0.08	96,96,96,96	0
59	MG	13	1683	1/1	0.89	0.10	55,55,55,55	0
59	MG	14	3214	1/1	0.89	0.25	57,57,57,57	0
58	K	14	3022	1/1	0.89	0.07	92,92,92,92	0
59	MG	1H	3134	1/1	0.89	0.23	59,59,59,59	0
58	K	1G	1619	1/1	0.89	0.09	79,79,79,79	0
59	MG	1G	1690	1/1	0.89	0.08	109,109,109,109	0
58	K	14	3067	1/1	0.89	0.07	90,90,90,90	0
59	MG	D8	202	1/1	0.89	0.08	52,52,52,52	0
59	MG	D8	203	1/1	0.89	0.12	55,55,55,55	0
59	MG	1H	3253	1/1	0.89	0.15	53,53,53,53	0
59	MG	13	1642	1/1	0.89	0.19	59,59,59,59	0
59	MG	14	3251	1/1	0.89	0.21	52,52,52,52	0
59	MG	1H	3263	1/1	0.89	0.18	33,33,33,33	0
58	K	1H	3037	1/1	0.89	0.07	77,77,77,77	0
58	K	13	1629	1/1	0.89	0.06	92,92,92,92	0
59	MG	14	3262	1/1	0.89	0.17	45,45,45,45	0
59	MG	1H	3280	1/1	0.89	0.15	64,64,64,64	0
59	MG	1H	3161	1/1	0.89	0.15	32,32,32,32	0
59	MG	14	3145	1/1	0.89	0.17	86,86,86,86	0
59	MG	14	3149	1/1	0.89	0.18	46,46,46,46	0
58	K	1H	3017	1/1	0.89	0.06	94,94,94,94	0
59	MG	14	3153	1/1	0.89	0.12	68,68,68,68	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
58	K	1H	3043	1/1	0.89	0.07	82,82,82,82	0
59	MG	14	3315	1/1	0.89	0.07	84,84,84,84	0
58	K	1H	3031	1/1	0.89	0.07	96,96,96,96	0
59	MG	E5	101	1/1	0.89	0.05	36,36,36,36	0
58	K	14	3066	1/1	0.90	0.13	80,80,80,80	0
59	MG	1G	1654	1/1	0.90	0.18	58,58,58,58	0
59	MG	14	3176	1/1	0.90	0.23	67,67,67,67	0
58	K	1H	3027	1/1	0.90	0.13	91,91,91,91	0
58	K	31	301	1/1	0.90	0.08	60,60,60,60	0
59	MG	1H	3217	1/1	0.90	0.14	52,52,52,52	0
58	K	1G	1620	1/1	0.90	0.07	101,101,101,101	0
58	K	1H	3029	1/1	0.90	0.06	104,104,104,104	0
58	K	1G	1625	1/1	0.90	0.10	106,106,106,106	0
58	K	13	1625	1/1	0.90	0.07	94,94,94,94	0
59	MG	1H	3126	1/1	0.90	0.23	53,53,53,53	0
59	MG	1H	3131	1/1	0.90	0.16	42,42,42,42	0
58	K	14	3078	1/1	0.90	0.20	98,98,98,98	0
58	K	1H	3032	1/1	0.90	0.09	58,58,58,58	0
58	K	14	3040	1/1	0.90	0.13	72,72,72,72	0
58	K	13	1609	1/1	0.90	0.10	91,91,91,91	0
59	MG	1H	3332	1/1	0.90	0.29	59,59,59,59	0
59	MG	1H	3335	1/1	0.90	0.13	50,50,50,50	0
59	MG	1G	1673	1/1	0.90	0.22	73,73,73,73	0
58	K	1H	3084	1/1	0.90	0.07	77,77,77,77	0
59	MG	1H	3248	1/1	0.90	0.12	47,47,47,47	0
59	MG	1H	3142	1/1	0.90	0.13	71,71,71,71	0
59	MG	13	1666	1/1	0.90	0.07	83,83,83,83	0
58	K	1H	3088	1/1	0.90	0.07	74,74,74,74	0
58	K	14	3048	1/1	0.90	0.10	101,101,101,101	0
59	MG	14	3239	1/1	0.90	0.21	66,66,66,66	0
58	K	14	3004	1/1	0.90	0.08	55,55,55,55	0
59	MG	14	3105	1/1	0.90	0.20	36,36,36,36	0
59	MG	1H	3172	1/1	0.90	0.15	48,48,48,48	0
59	MG	1G	1629	1/1	0.90	0.09	69,69,69,69	0
58	K	13	1615	1/1	0.90	0.05	91,91,91,91	0
58	K	1H	3054	1/1	0.90	0.10	68,68,68,68	0
59	MG	1H	3181	1/1	0.90	0.19	41,41,41,41	0
59	MG	14	3141	1/1	0.90	0.17	60,60,60,60	0
59	MG	13	1703	1/1	0.90	0.25	69,69,69,69	0
58	K	14	3014	1/1	0.90	0.06	82,82,82,82	0
59	MG	1G	1640	1/1	0.90	0.20	53,53,53,53	0
59	MG	1H	3291	1/1	0.90	0.13	62,62,62,62	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	1G	1645	1/1	0.90	0.17	40,40,40,40	0
59	MG	1H	3295	1/1	0.90	0.09	46,46,46,46	0
59	MG	1H	3297	1/1	0.90	0.19	52,52,52,52	0
59	MG	1H	3191	1/1	0.90	0.27	72,72,72,72	0
59	MG	14	3375	1/1	0.90	0.09	57,57,57,57	0
59	MG	14	3377	1/1	0.90	0.08	70,70,70,70	0
59	MG	13	1705	1/1	0.90	0.20	60,60,60,60	0
59	MG	29	304	1/1	0.90	0.19	62,62,62,62	0
58	K	1H	3060	1/1	0.90	0.15	55,55,55,55	0
59	MG	1H	3484	1/1	0.91	0.08	47,47,47,47	0
59	MG	14	3188	1/1	0.91	0.12	54,54,54,54	0
59	MG	16	203	1/1	0.91	0.12	57,57,57,57	0
59	MG	1H	3199	1/1	0.91	0.11	60,60,60,60	0
59	MG	14	3193	1/1	0.91	0.11	63,63,63,63	0
59	MG	1H	3132	1/1	0.91	0.20	58,58,58,58	0
59	MG	D8	201	1/1	0.91	0.14	71,71,71,71	0
59	MG	1H	3204	1/1	0.91	0.20	51,51,51,51	0
59	MG	1H	3283	1/1	0.91	0.17	58,58,58,58	0
59	MG	13	1643	1/1	0.91	0.11	76,76,76,76	0
59	MG	1H	3208	1/1	0.91	0.27	52,52,52,52	0
59	MG	1G	1696	1/1	0.91	0.07	129,129,129,129	0
59	MG	14	3206	1/1	0.91	0.16	67,67,67,67	0
58	K	1H	3081	1/1	0.91	0.13	66,66,66,66	0
59	MG	1H	3290	1/1	0.91	0.21	51,51,51,51	0
59	MG	14	3213	1/1	0.91	0.30	58,58,58,58	0
59	MG	1G	1633	1/1	0.91	0.14	65,65,65,65	0
59	MG	14	3217	1/1	0.91	0.17	59,59,59,59	0
59	MG	1H	3212	1/1	0.91	0.07	57,57,57,57	0
58	K	32	301	1/1	0.91	0.10	133,133,133,133	0
59	MG	14	3113	1/1	0.91	0.24	51,51,51,51	0
58	K	13	1602	1/1	0.91	0.07	113,113,113,113	0
59	MG	14	3231	1/1	0.91	0.08	48,48,48,48	0
59	MG	14	3118	1/1	0.91	0.13	53,53,53,53	0
58	K	14	3023	1/1	0.91	0.11	93,93,93,93	0
59	MG	1H	3218	1/1	0.91	0.11	53,53,53,53	0
58	K	14	3025	1/1	0.91	0.07	97,97,97,97	0
59	MG	14	3244	1/1	0.91	0.18	46,46,46,46	0
59	MG	1H	3150	1/1	0.91	0.11	76,76,76,76	0
59	MG	14	3247	1/1	0.91	0.17	60,60,60,60	0
59	MG	14	3139	1/1	0.91	0.19	46,46,46,46	0
59	MG	1H	3151	1/1	0.91	0.26	54,54,54,54	0
59	MG	1H	3156	1/1	0.91	0.15	55,55,55,55	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
58	K	1G	1610	1/1	0.91	0.07	95,95,95,95	0
58	K	1G	1613	1/1	0.91	0.06	96,96,96,96	0
59	MG	14	3260	1/1	0.91	0.19	56,56,56,56	0
59	MG	14	3261	1/1	0.91	0.14	75,75,75,75	0
58	K	1H	3003	1/1	0.91	0.05	76,76,76,76	0
58	K	1H	3109	1/1	0.91	0.08	54,54,54,54	0
58	K	1G	1618	1/1	0.91	0.10	102,102,102,102	0
58	K	1H	3118	1/1	0.91	0.18	89,89,89,89	0
59	MG	13	1692	1/1	0.91	0.15	70,70,70,70	0
59	MG	1H	3246	1/1	0.91	0.13	58,58,58,58	0
58	K	14	3093	1/1	0.91	0.09	72,72,72,72	0
59	MG	14	3169	1/1	0.91	0.12	57,57,57,57	0
59	MG	1H	3183	1/1	0.91	0.18	49,49,49,49	0
59	MG	14	3350	1/1	0.91	0.20	69,69,69,69	0
58	K	1H	3035	1/1	0.91	0.09	98,98,98,98	0
59	MG	1H	3257	1/1	0.91	0.09	49,49,49,49	0
59	MG	14	3381	1/1	0.91	0.10	77,77,77,77	0
59	MG	14	3387	1/1	0.91	0.07	45,45,45,45	0
59	MG	14	3389	1/1	0.91	0.09	69,69,69,69	0
59	MG	1J	201	1/1	0.91	0.07	131,131,131,131	0
59	MG	1H	3186	1/1	0.91	0.23	50,50,50,50	0
59	MG	29	303	1/1	0.91	0.14	37,37,37,37	0
58	K	14	3013	1/1	0.91	0.06	98,98,98,98	0
58	K	1H	3111	1/1	0.91	0.11	75,75,75,75	0
58	K	1H	3089	1/1	0.92	0.09	50,50,50,50	0
59	MG	1H	3273	1/1	0.92	0.06	49,49,49,49	0
59	MG	1H	3274	1/1	0.92	0.10	56,56,56,56	0
59	MG	13	1711	1/1	0.92	0.15	48,48,48,48	0
59	MG	16	206	1/1	0.92	0.12	62,62,62,62	0
59	MG	14	3200	1/1	0.92	0.09	102,102,102,102	0
59	MG	13	1712	1/1	0.92	0.12	50,50,50,50	0
59	MG	1G	1691	1/1	0.92	0.08	84,84,84,84	0
59	MG	13	1741	1/1	0.92	0.06	106,106,106,106	0
59	MG	13	1680	1/1	0.92	0.18	71,71,71,71	0
59	MG	14	3208	1/1	0.92	0.07	61,61,61,61	0
58	K	29	301	1/1	0.92	0.09	56,56,56,56	0
58	K	14	3061	1/1	0.92	0.06	103,103,103,103	0
59	MG	14	3101	1/1	0.92	0.09	61,61,61,61	0
59	MG	14	3102	1/1	0.92	0.08	57,57,57,57	0
59	MG	13	1684	1/1	0.92	0.14	56,56,56,56	0
59	MG	1H	3288	1/1	0.92	0.09	51,51,51,51	0
59	MG	14	3106	1/1	0.92	0.18	34,34,34,34	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	14	3222	1/1	0.92	0.07	57,57,57,57	0
58	K	14	3064	1/1	0.92	0.15	70,70,70,70	0
59	MG	13	1640	1/1	0.92	0.13	72,72,72,72	0
59	MG	14	3228	1/1	0.92	0.16	88,88,88,88	0
58	K	1H	3119	1/1	0.92	0.08	92,92,92,92	0
59	MG	14	3120	1/1	0.92	0.08	44,44,44,44	0
58	K	13	1624	1/1	0.92	0.11	83,83,83,83	0
59	MG	14	3129	1/1	0.92	0.08	44,44,44,44	0
59	MG	14	3240	1/1	0.92	0.12	59,59,59,59	0
58	K	1H	3053	1/1	0.92	0.12	63,63,63,63	0
58	K	1G	1624	1/1	0.92	0.07	116,116,116,116	0
59	MG	14	3245	1/1	0.92	0.22	64,64,64,64	0
59	MG	1H	3305	1/1	0.92	0.18	57,57,57,57	0
59	MG	1G	1642	1/1	0.92	0.19	59,59,59,59	0
59	MG	14	3140	1/1	0.92	0.18	45,45,45,45	0
58	K	1H	3093	1/1	0.92	0.06	79,79,79,79	0
58	K	13	1631	1/1	0.92	0.08	93,93,93,93	0
59	MG	14	3143	1/1	0.92	0.16	72,72,72,72	0
58	K	1H	3015	1/1	0.92	0.07	80,80,80,80	0
59	MG	14	3256	1/1	0.92	0.24	64,64,64,64	0
59	MG	14	3258	1/1	0.92	0.12	37,37,37,37	0
58	K	1H	3026	1/1	0.92	0.13	62,62,62,62	0
59	MG	1H	3313	1/1	0.92	0.12	52,52,52,52	0
59	MG	1H	3145	1/1	0.92	0.13	32,32,32,32	0
58	K	13	1612[B]	1/1	0.92	0.25	23,23,23,23	1
58	K	1H	3028	1/1	0.92	0.07	50,50,50,50	0
59	MG	14	3159	1/1	0.92	0.17	53,53,53,53	0
58	K	1H	3073	1/1	0.92	0.10	47,47,47,47	0
58	K	14	3046	1/1	0.92	0.14	68,68,68,68	0
58	K	1H	3075	1/1	0.92	0.07	81,81,81,81	0
59	MG	1H	3322	1/1	0.92	0.16	60,60,60,60	0
59	MG	1H	3242	1/1	0.92	0.15	57,57,57,57	0
58	K	13	1612[A]	1/1	0.92	0.25	35,35,35,35	1
59	MG	1H	3168	1/1	0.92	0.22	46,46,46,46	0
59	MG	14	3177	1/1	0.92	0.14	64,64,64,64	0
59	MG	14	3378	1/1	0.92	0.09	71,71,71,71	0
59	MG	14	3178	1/1	0.92	0.10	36,36,36,36	0
58	K	1H	3041	1/1	0.92	0.10	66,66,66,66	0
58	K	14	3050	1/1	0.92	0.09	81,81,81,81	0
58	K	1H	3030	1/1	0.92	0.10	86,86,86,86	0
59	MG	13	1706	1/1	0.92	0.20	63,63,63,63	0
58	K	5I	101	1/1	0.92	0.09	129,129,129,129	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
58	K	1H	3045	1/1	0.92	0.06	72,72,72,72	0
59	MG	1H	3352	1/1	0.92	0.07	59,59,59,59	0
59	MG	1H	3179	1/1	0.93	0.09	51,51,51,51	0
58	K	14	3010	1/1	0.93	0.06	53,53,53,53	0
59	MG	14	3191	1/1	0.93	0.20	45,45,45,45	0
58	K	1H	3057	1/1	0.93	0.05	43,43,43,43	0
58	K	1H	3013	1/1	0.93	0.06	83,83,83,83	0
59	MG	1H	3261	1/1	0.93	0.22	49,49,49,49	0
59	MG	1H	3262	1/1	0.93	0.12	49,49,49,49	0
58	K	1G	1608	1/1	0.93	0.07	113,113,113,113	0
59	MG	1H	3268	1/1	0.93	0.16	50,50,50,50	0
59	MG	BI	202	1/1	0.93	0.07	94,94,94,94	0
59	MG	1H	3122	1/1	0.93	0.18	29,29,29,29	0
59	MG	1G	1711	1/1	0.93	0.06	104,104,104,104	0
59	MG	1G	1716	1/1	0.93	0.07	107,107,107,107	0
59	MG	1H	3125	1/1	0.93	0.15	41,41,41,41	0
58	K	13	1616	1/1	0.93	0.04	96,96,96,96	0
59	MG	41	203	1/1	0.93	0.13	40,40,40,40	0
59	MG	14	3100	1/1	0.93	0.07	47,47,47,47	0
59	MG	1H	3279	1/1	0.93	0.20	74,74,74,74	0
59	MG	1H	3129	1/1	0.93	0.16	34,34,34,34	0
58	K	14	3021	1/1	0.93	0.07	47,47,47,47	0
59	MG	N8	101	1/1	0.93	0.10	56,56,56,56	0
59	MG	14	3219	1/1	0.93	0.12	45,45,45,45	0
58	K	1H	3011	1/1	0.93	0.04	61,61,61,61	0
59	MG	14	3110	1/1	0.93	0.19	48,48,48,48	0
59	MG	14	3112	1/1	0.93	0.06	55,55,55,55	0
59	MG	13	1675	1/1	0.93	0.10	91,91,91,91	0
59	MG	14	3114	1/1	0.93	0.15	57,57,57,57	0
58	K	1H	3090	1/1	0.93	0.06	65,65,65,65	0
59	MG	1H	3137	1/1	0.93	0.14	42,42,42,42	0
59	MG	14	3234	1/1	0.93	0.34	67,67,67,67	0
59	MG	13	1700	1/1	0.93	0.11	63,63,63,63	0
58	K	14	3024	1/1	0.93	0.07	57,57,57,57	0
59	MG	14	3126	1/1	0.93	0.12	69,69,69,69	0
59	MG	14	3242	1/1	0.93	0.18	44,44,44,44	0
58	K	1H	3048	1/1	0.93	0.05	67,67,67,67	0
59	MG	1H	3293	1/1	0.93	0.10	55,55,55,55	0
59	MG	14	3135	1/1	0.93	0.17	42,42,42,42	0
59	MG	13	1647	1/1	0.93	0.15	59,59,59,59	0
59	MG	1H	3219	1/1	0.93	0.15	50,50,50,50	0
59	MG	1H	3301	1/1	0.93	0.12	59,59,59,59	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(Å ²)	Q<0.9
59	MG	13	1650	1/1	0.93	0.06	73,73,73,73	0
59	MG	1H	3222	1/1	0.93	0.18	67,67,67,67	0
59	MG	1H	3306	1/1	0.93	0.14	44,44,44,44	0
59	MG	1G	1647	1/1	0.93	0.10	80,80,80,80	0
59	MG	1H	3307	1/1	0.93	0.09	39,39,39,39	0
59	MG	14	3257	1/1	0.93	0.15	48,48,48,48	0
59	MG	13	1681	1/1	0.93	0.13	76,76,76,76	0
58	K	14	3054	1/1	0.93	0.04	77,77,77,77	0
58	K	1G	1616	1/1	0.93	0.05	89,89,89,89	0
59	MG	1H	3157	1/1	0.93	0.12	62,62,62,62	0
59	MG	1G	1653	1/1	0.93	0.19	52,52,52,52	0
59	MG	14	3264	1/1	0.93	0.16	55,55,55,55	0
59	MG	1H	3233	1/1	0.93	0.14	59,59,59,59	0
59	MG	13	1708	1/1	0.93	0.18	59,59,59,59	0
58	K	1H	3039	1/1	0.93	0.05	81,81,81,81	0
59	MG	14	3165	1/1	0.93	0.14	70,70,70,70	0
59	MG	1G	1657	1/1	0.93	0.12	74,74,74,74	0
59	MG	1H	3160	1/1	0.93	0.16	29,29,29,29	0
59	MG	14	3170	1/1	0.93	0.12	52,52,52,52	0
59	MG	14	3341	1/1	0.93	0.10	52,52,52,52	0
59	MG	1G	1659	1/1	0.93	0.13	72,72,72,72	0
59	MG	14	3172	1/1	0.93	0.15	72,72,72,72	0
59	MG	14	3173	1/1	0.93	0.28	64,64,64,64	0
58	K	13	1601	1/1	0.93	0.07	87,87,87,87	0
59	MG	1H	3241	1/1	0.93	0.12	39,39,39,39	0
59	MG	14	3386	1/1	0.93	0.10	82,82,82,82	0
59	MG	1H	3167	1/1	0.93	0.05	33,33,33,33	0
59	MG	13	1658	1/1	0.93	0.13	35,35,35,35	0
59	MG	13	1660	1/1	0.93	0.15	60,60,60,60	0
59	MG	14	3182	1/1	0.93	0.12	42,42,42,42	0
58	K	1H	3023	1/1	0.93	0.07	68,68,68,68	0
58	K	1H	3055	1/1	0.93	0.04	40,40,40,40	0
59	MG	55	201	1/1	0.93	0.15	59,59,59,59	0
59	MG	1H	3249	1/1	0.93	0.15	53,53,53,53	0
59	MG	1H	3196	1/1	0.94	0.06	42,42,42,42	0
59	MG	14	3210	1/1	0.94	0.19	45,45,45,45	0
58	K	1G	1603	1/1	0.94	0.16	79,79,79,79	0
58	K	13	1623	1/1	0.94	0.10	121,121,121,121	0
59	MG	1H	3320	1/1	0.94	0.19	43,43,43,43	0
59	MG	14	3215	1/1	0.94	0.13	45,45,45,45	0
58	K	1H	3021	1/1	0.94	0.06	78,78,78,78	0
59	MG	14	3134	1/1	0.94	0.08	67,67,67,67	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	1H	3143	1/1	0.94	0.18	63,63,63,63	0
59	MG	13	1648	1/1	0.94	0.10	61,61,61,61	0
59	MG	1H	3209	1/1	0.94	0.16	33,33,33,33	0
59	MG	14	3224	1/1	0.94	0.04	53,53,53,53	0
59	MG	1H	3269	1/1	0.94	0.17	54,54,54,54	0
59	MG	1H	3210	1/1	0.94	0.09	50,50,50,50	0
59	MG	1H	3147	1/1	0.94	0.09	19,19,19,19	0
58	K	1H	3103	1/1	0.94	0.05	65,65,65,65	0
58	K	1H	3107	1/1	0.94	0.05	82,82,82,82	0
59	MG	1H	3334	1/1	0.94	0.16	55,55,55,55	0
59	MG	14	3148	1/1	0.94	0.11	58,58,58,58	0
59	MG	14	3237	1/1	0.94	0.11	58,58,58,58	0
59	MG	1H	3214	1/1	0.94	0.10	26,26,26,26	0
58	K	14	3030	1/1	0.94	0.07	56,56,56,56	0
59	MG	14	3151	1/1	0.94	0.14	57,57,57,57	0
59	MG	1H	3281	1/1	0.94	0.08	93,93,93,93	0
58	K	14	3032	1/1	0.94	0.07	61,61,61,61	0
59	MG	1H	3393	1/1	0.94	0.08	44,44,44,44	0
59	MG	1H	3415	1/1	0.94	0.07	94,94,94,94	0
59	MG	1H	3474	1/1	0.94	0.07	47,47,47,47	0
59	MG	1H	3477	1/1	0.94	0.08	43,43,43,43	0
58	K	14	3012	1/1	0.94	0.05	91,91,91,91	0
59	MG	1H	3518	1/1	0.94	0.07	40,40,40,40	0
58	K	14	3056	1/1	0.94	0.04	57,57,57,57	0
59	MG	16	204	1/1	0.94	0.13	80,80,80,80	0
58	K	14	3057	1/1	0.94	0.05	68,68,68,68	0
58	K	13	1603	1/1	0.94	0.07	117,117,117,117	0
59	MG	1H	3165	1/1	0.94	0.14	30,30,30,30	0
58	K	14	3060	1/1	0.94	0.04	67,67,67,67	0
59	MG	1G	1683	1/1	0.94	0.12	90,90,90,90	0
59	MG	1G	1685	1/1	0.94	0.09	87,87,87,87	0
58	K	1H	3047	1/1	0.94	0.14	37,37,37,37	0
59	MG	13	1664	1/1	0.94	0.17	53,53,53,53	0
59	MG	14	3181	1/1	0.94	0.07	46,46,46,46	0
58	K	14	3063	1/1	0.94	0.06	84,84,84,84	0
59	MG	1H	3173	1/1	0.94	0.09	57,57,57,57	0
58	K	1H	3024	1/1	0.94	0.08	56,56,56,56	0
59	MG	1H	3177	1/1	0.94	0.14	54,54,54,54	0
58	K	13	1622	1/1	0.94	0.05	73,73,73,73	0
59	MG	14	3289	1/1	0.94	0.08	53,53,53,53	0
59	MG	1H	3240	1/1	0.94	0.21	52,52,52,52	0
59	MG	14	3325	1/1	0.94	0.09	81,81,81,81	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	14	3099	1/1	0.94	0.10	32,32,32,32	0
59	MG	14	3347	1/1	0.94	0.09	57,57,57,57	0
59	MG	1G	1632	1/1	0.94	0.09	109,109,109,109	0
59	MG	14	3357	1/1	0.94	0.17	95,95,95,95	0
58	K	14	3094	1/1	0.94	0.05	111,111,111,111	0
59	MG	14	3195	1/1	0.94	0.11	64,64,64,64	0
58	K	1H	3086	1/1	0.94	0.09	79,79,79,79	0
59	MG	14	3379	1/1	0.94	0.12	42,42,42,42	0
58	K	14	3068	1/1	0.94	0.10	48,48,48,48	0
59	MG	1G	1636	1/1	0.94	0.10	89,89,89,89	0
58	K	14	3044	1/1	0.94	0.08	95,95,95,95	0
59	MG	1G	1638	1/1	0.94	0.15	43,43,43,43	0
59	MG	14	3398	1/1	0.94	0.09	63,63,63,63	0
59	MG	14	3406	1/1	0.94	0.08	55,55,55,55	0
59	MG	13	1638	1/1	0.94	0.12	58,58,58,58	0
58	K	1H	3042	1/1	0.94	0.14	62,62,62,62	0
59	MG	1H	3190	1/1	0.94	0.18	49,49,49,49	0
59	MG	14	3205	1/1	0.94	0.09	37,37,37,37	0
58	K	14	3073	1/1	0.94	0.10	81,81,81,81	0
59	MG	1G	1644	1/1	0.94	0.15	55,55,55,55	0
59	MG	1G	1725	1/1	0.95	0.06	97,97,97,97	0
59	MG	1G	1727	1/1	0.95	0.05	93,93,93,93	0
59	MG	2L	101	1/1	0.95	0.11	55,55,55,55	0
58	K	14	3077	1/1	0.95	0.12	72,72,72,72	0
59	MG	13	1733	1/1	0.95	0.09	117,117,117,117	0
59	MG	13	1738	1/1	0.95	0.09	91,91,91,91	0
59	MG	14	3207	1/1	0.95	0.17	58,58,58,58	0
59	MG	1H	3287	1/1	0.95	0.21	43,43,43,43	0
59	MG	16	210	1/1	0.95	0.08	98,98,98,98	0
58	K	14	3009	1/1	0.95	0.06	69,69,69,69	0
59	MG	78	201	1/1	0.95	0.06	32,32,32,32	0
59	MG	1H	3164	1/1	0.95	0.09	45,45,45,45	0
59	MG	13	1651	1/1	0.95	0.12	51,51,51,51	0
59	MG	1H	3225	1/1	0.95	0.21	58,58,58,58	0
58	K	14	3053	1/1	0.95	0.06	63,63,63,63	0
59	MG	1H	3294	1/1	0.95	0.13	35,35,35,35	0
58	K	14	3031	1/1	0.95	0.05	78,78,78,78	0
59	MG	1H	3296	1/1	0.95	0.10	50,50,50,50	0
58	K	1H	3108	1/1	0.95	0.09	86,86,86,86	0
59	MG	14	3223	1/1	0.95	0.08	53,53,53,53	0
58	K	13	1621	1/1	0.95	0.05	59,59,59,59	0
59	MG	14	3119	1/1	0.95	0.10	51,51,51,51	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
58	K	14	3034	1/1	0.95	0.06	49,49,49,49	0
59	MG	14	3121	1/1	0.95	0.06	43,43,43,43	0
59	MG	2K	101	1/1	0.95	0.16	42,42,42,42	0
59	MG	14	3232	1/1	0.95	0.23	69,69,69,69	0
58	K	14	3035	1/1	0.95	0.12	62,62,62,62	0
59	MG	1H	3178	1/1	0.95	0.18	34,34,34,34	0
58	K	14	3059	1/1	0.95	0.10	73,73,73,73	0
58	K	1H	3064	1/1	0.95	0.07	49,49,49,49	0
58	K	14	3037	1/1	0.95	0.06	60,60,60,60	0
58	K	14	3090	1/1	0.95	0.06	65,65,65,65	0
58	K	1H	3066	1/1	0.95	0.06	80,80,80,80	0
59	MG	1H	3314	1/1	0.95	0.14	49,49,49,49	0
59	MG	1H	3245	1/1	0.95	0.14	41,41,41,41	0
58	K	1G	1612	1/1	0.95	0.09	101,101,101,101	0
59	MG	1H	3247	1/1	0.95	0.18	44,44,44,44	0
59	MG	13	1668	1/1	0.95	0.09	41,41,41,41	0
59	MG	14	3248	1/1	0.95	0.05	41,41,41,41	0
59	MG	14	3249	1/1	0.95	0.08	54,54,54,54	0
59	MG	1H	3136	1/1	0.95	0.19	37,37,37,37	0
58	K	14	3065	1/1	0.95	0.07	92,92,92,92	0
59	MG	1H	3193	1/1	0.95	0.09	43,43,43,43	0
59	MG	1H	3254	1/1	0.95	0.13	50,50,50,50	0
58	K	1H	3100	1/1	0.95	0.10	78,78,78,78	0
59	MG	1H	3258	1/1	0.95	0.11	48,48,48,48	0
58	K	1H	3002	1/1	0.95	0.04	71,71,71,71	0
59	MG	1H	3260	1/1	0.95	0.12	43,43,43,43	0
59	MG	1H	3200	1/1	0.95	0.10	67,67,67,67	0
58	K	1H	3050	1/1	0.95	0.09	70,70,70,70	0
59	MG	1H	3328	1/1	0.95	0.19	61,61,61,61	0
59	MG	1H	3141	1/1	0.95	0.11	37,37,37,37	0
59	MG	1H	3265	1/1	0.95	0.23	63,63,63,63	0
58	K	14	3002	1/1	0.95	0.06	90,90,90,90	0
58	K	1G	1602	1/1	0.95	0.18	70,70,70,70	0
59	MG	1H	3347	1/1	0.95	0.09	50,50,50,50	0
58	K	14	3072	1/1	0.95	0.04	89,89,89,89	0
59	MG	13	1641	1/1	0.95	0.12	90,90,90,90	0
59	MG	1H	3275	1/1	0.95	0.23	60,60,60,60	0
59	MG	14	3175	1/1	0.95	0.23	59,59,59,59	0
58	K	1H	3061	1/1	0.95	0.09	47,47,47,47	0
59	MG	1H	3419	1/1	0.95	0.09	43,43,43,43	0
59	MG	14	3340	1/1	0.95	0.07	74,74,74,74	0
59	MG	1G	1669	1/1	0.95	0.05	72,72,72,72	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	1H	3426	1/1	0.95	0.11	68,68,68,68	0
59	MG	14	3348	1/1	0.95	0.09	102,102,102,102	0
59	MG	1H	3427	1/1	0.95	0.08	41,41,41,41	0
59	MG	1H	3461	1/1	0.95	0.08	33,33,33,33	0
59	MG	1H	3466	1/1	0.95	0.07	34,34,34,34	0
58	K	1H	3106	1/1	0.95	0.10	92,92,92,92	0
59	MG	1H	3153	1/1	0.95	0.11	44,44,44,44	0
59	MG	14	3186	1/1	0.95	0.09	48,48,48,48	0
59	MG	14	3380	1/1	0.95	0.09	70,70,70,70	0
59	MG	1G	1677	1/1	0.95	0.04	76,76,76,76	0
59	MG	14	3189	1/1	0.95	0.17	26,26,26,26	0
58	K	1H	3095	1/1	0.95	0.06	44,44,44,44	0
59	MG	1H	3493	1/1	0.95	0.13	63,63,63,63	0
59	MG	14	3394	1/1	0.95	0.08	82,82,82,82	0
59	MG	14	3395	1/1	0.95	0.08	50,50,50,50	0
59	MG	1G	1686	1/1	0.95	0.06	94,94,94,94	0
59	MG	14	3399	1/1	0.95	0.07	55,55,55,55	0
59	MG	14	3401	1/1	0.95	0.13	89,89,89,89	0
59	MG	1H	3514	1/1	0.95	0.07	88,88,88,88	0
59	MG	14	3194	1/1	0.95	0.05	64,64,64,64	0
59	MG	1H	3515	1/1	0.95	0.13	90,90,90,90	0
58	K	14	3028	1/1	0.95	0.06	56,56,56,56	0
59	MG	1H	3522	1/1	0.95	0.11	102,102,102,102	0
59	MG	1H	3216	1/1	0.95	0.12	45,45,45,45	0
59	MG	1G	1722	1/1	0.95	0.05	97,97,97,97	0
59	MG	1H	3388	1/1	0.96	0.08	38,38,38,38	0
58	K	1H	3046	1/1	0.96	0.10	50,50,50,50	0
58	K	1H	3058	1/1	0.96	0.03	56,56,56,56	0
59	MG	1H	3394	1/1	0.96	0.06	38,38,38,38	0
59	MG	1H	3399	1/1	0.96	0.08	54,54,54,54	0
59	MG	13	1739	1/1	0.96	0.05	97,97,97,97	0
58	K	14	3006	1/1	0.96	0.12	62,62,62,62	0
59	MG	1G	1675	1/1	0.96	0.11	54,54,54,54	0
59	MG	13	1654	1/1	0.96	0.12	36,36,36,36	0
58	K	1H	3076	1/1	0.96	0.05	86,86,86,86	0
59	MG	1G	1678	1/1	0.96	0.06	71,71,71,71	0
59	MG	1H	3437	1/1	0.96	0.16	49,49,49,49	0
59	MG	1H	3450	1/1	0.96	0.05	91,91,91,91	0
59	MG	1H	3163	1/1	0.96	0.14	40,40,40,40	0
59	MG	1H	3220	1/1	0.96	0.07	45,45,45,45	0
58	K	1H	3005	1/1	0.96	0.05	67,67,67,67	0
59	MG	1H	3475	1/1	0.96	0.08	91,91,91,91	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	1G	1702	1/1	0.96	0.11	87,87,87,87	0
59	MG	1G	1704	1/1	0.96	0.04	110,110,110,110	0
59	MG	1G	1710	1/1	0.96	0.08	70,70,70,70	0
59	MG	1H	3476	1/1	0.96	0.08	42,42,42,42	0
58	K	14	3087	1/1	0.96	0.03	84,84,84,84	0
59	MG	1G	1718	1/1	0.96	0.05	91,91,91,91	0
59	MG	14	3212	1/1	0.96	0.19	34,34,34,34	0
59	MG	1H	3481	1/1	0.96	0.07	43,43,43,43	0
59	MG	1H	3483	1/1	0.96	0.12	75,75,75,75	0
58	K	1H	3006	1/1	0.96	0.03	59,59,59,59	0
59	MG	1H	3492	1/1	0.96	0.06	67,67,67,67	0
59	MG	1H	3286	1/1	0.96	0.20	52,52,52,52	0
59	MG	1H	3498	1/1	0.96	0.08	37,37,37,37	0
59	MG	1H	3508	1/1	0.96	0.19	78,78,78,78	0
59	MG	14	3221	1/1	0.96	0.19	52,52,52,52	0
59	MG	14	3097	1/1	0.96	0.14	38,38,38,38	0
59	MG	14	3098	1/1	0.96	0.12	38,38,38,38	0
59	MG	1H	3224	1/1	0.96	0.17	51,51,51,51	0
58	K	14	3011	1/1	0.96	0.05	56,56,56,56	0
58	K	1H	3022	1/1	0.96	0.08	51,51,51,51	0
59	MG	1H	3228	1/1	0.96	0.11	48,48,48,48	0
59	MG	14	3229	1/1	0.96	0.17	65,65,65,65	0
58	K	1G	1621	1/1	0.96	0.04	74,74,74,74	0
59	MG	14	3104	1/1	0.96	0.14	29,29,29,29	0
59	MG	1H	3292	1/1	0.96	0.14	54,54,54,54	0
59	MG	1H	3230	1/1	0.96	0.09	38,38,38,38	0
59	MG	14	3107	1/1	0.96	0.11	40,40,40,40	0
59	MG	14	3108	1/1	0.96	0.11	62,62,62,62	0
58	K	1H	3007	1/1	0.96	0.12	54,54,54,54	0
58	K	14	3015	1/1	0.96	0.09	52,52,52,52	0
59	MG	14	3241	1/1	0.96	0.14	34,34,34,34	0
59	MG	1H	3234	1/1	0.96	0.14	33,33,33,33	0
59	MG	21	303	1/1	0.96	0.14	30,30,30,30	0
59	MG	1H	3176	1/1	0.96	0.14	32,32,32,32	0
59	MG	1H	3298	1/1	0.96	0.13	23,23,23,23	0
59	MG	1H	3300	1/1	0.96	0.10	31,31,31,31	0
59	MG	1H	3127	1/1	0.96	0.22	54,54,54,54	0
59	MG	1H	3302	1/1	0.96	0.16	41,41,41,41	0
59	MG	1H	3237	1/1	0.96	0.15	51,51,51,51	0
59	MG	Q8	101	1/1	0.96	0.08	32,32,32,32	0
59	MG	1G	1626	1/1	0.96	0.12	72,72,72,72	0
58	K	1G	1623	1/1	0.96	0.04	126,126,126,126	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	14	3132	1/1	0.96	0.05	47,47,47,47	0
59	MG	13	1667	1/1	0.96	0.15	39,39,39,39	0
59	MG	14	3255	1/1	0.96	0.14	52,52,52,52	0
58	K	14	3018	1/1	0.96	0.06	71,71,71,71	0
58	K	1H	3085	1/1	0.96	0.04	64,64,64,64	0
58	K	1H	3116	1/1	0.96	0.04	80,80,80,80	0
59	MG	14	3138	1/1	0.96	0.13	40,40,40,40	0
59	MG	1H	3135	1/1	0.96	0.05	50,50,50,50	0
59	MG	1H	3311	1/1	0.96	0.21	57,57,57,57	0
58	K	1H	3036	1/1	0.96	0.05	55,55,55,55	0
59	MG	13	1673	1/1	0.96	0.12	56,56,56,56	0
58	K	1H	3067	1/1	0.96	0.05	49,49,49,49	0
59	MG	14	3144	1/1	0.96	0.06	50,50,50,50	0
58	K	1H	3008	1/1	0.96	0.12	61,61,61,61	0
59	MG	14	3146	1/1	0.96	0.05	30,30,30,30	0
59	MG	14	3147	1/1	0.96	0.08	54,54,54,54	0
59	MG	1H	3192	1/1	0.96	0.08	49,49,49,49	0
59	MG	14	3277	1/1	0.96	0.08	53,53,53,53	0
59	MG	14	3278	1/1	0.96	0.08	43,43,43,43	0
58	K	1H	3105	1/1	0.96	0.05	60,60,60,60	0
59	MG	14	3293	1/1	0.96	0.06	91,91,91,91	0
59	MG	14	3308	1/1	0.96	0.06	80,80,80,80	0
59	MG	1G	1641	1/1	0.96	0.14	53,53,53,53	0
59	MG	14	3317	1/1	0.96	0.05	94,94,94,94	0
59	MG	14	3324	1/1	0.96	0.06	62,62,62,62	0
59	MG	1H	3194	1/1	0.96	0.17	45,45,45,45	0
59	MG	14	3332	1/1	0.96	0.04	110,110,110,110	0
59	MG	14	3338	1/1	0.96	0.06	73,73,73,73	0
58	K	1G	1611	1/1	0.96	0.06	94,94,94,94	0
58	K	16	201	1/1	0.96	0.06	102,102,102,102	0
59	MG	14	3344	1/1	0.96	0.13	89,89,89,89	0
59	MG	13	1646	1/1	0.96	0.07	59,59,59,59	0
59	MG	14	3157	1/1	0.96	0.10	57,57,57,57	0
59	MG	1H	3201	1/1	0.96	0.11	48,48,48,48	0
59	MG	1H	3202	1/1	0.96	0.13	46,46,46,46	0
59	MG	14	3366	1/1	0.96	0.06	77,77,77,77	0
58	K	13	1632	1/1	0.96	0.06	83,83,83,83	0
59	MG	14	3376	1/1	0.96	0.09	61,61,61,61	0
58	K	14	3080	1/1	0.96	0.03	94,94,94,94	0
59	MG	1H	3148	1/1	0.96	0.10	39,39,39,39	0
59	MG	1H	3149	1/1	0.96	0.14	29,29,29,29	0
59	MG	1H	3264	1/1	0.96	0.15	47,47,47,47	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
58	K	14	3029	1/1	0.96	0.05	68,68,68,68	0
59	MG	14	3385	1/1	0.96	0.05	63,63,63,63	0
59	MG	1H	3267	1/1	0.96	0.06	60,60,60,60	0
59	MG	1H	3333	1/1	0.96	0.13	50,50,50,50	0
59	MG	13	1714	1/1	0.96	0.04	106,106,106,106	0
59	MG	14	3391	1/1	0.96	0.06	62,62,62,62	0
59	MG	14	3392	1/1	0.96	0.08	69,69,69,69	0
59	MG	13	1732	1/1	0.96	0.08	81,81,81,81	0
59	MG	1H	3271	1/1	0.96	0.17	45,45,45,45	0
59	MG	1H	3341	1/1	0.96	0.09	31,31,31,31	0
59	MG	1H	3272	1/1	0.96	0.09	60,60,60,60	0
59	MG	1H	3350	1/1	0.96	0.07	26,26,26,26	0
59	MG	14	3402	1/1	0.96	0.04	123,123,123,123	0
59	MG	1H	3154	1/1	0.96	0.17	31,31,31,31	0
59	MG	1H	3359	1/1	0.96	0.07	55,55,55,55	0
59	MG	1H	3363	1/1	0.96	0.07	32,32,32,32	0
59	MG	1H	3369	1/1	0.96	0.06	52,52,52,52	0
59	MG	1H	3373	1/1	0.96	0.07	39,39,39,39	0
59	MG	1H	3377	1/1	0.96	0.06	44,44,44,44	0
59	MG	14	3187	1/1	0.96	0.12	41,41,41,41	0
58	K	1H	3114	1/1	0.97	0.04	58,58,58,58	0
59	MG	14	3116	1/1	0.97	0.11	41,41,41,41	0
58	K	1H	3071	1/1	0.97	0.08	65,65,65,65	0
59	MG	1H	3404	1/1	0.97	0.05	65,65,65,65	0
58	K	1H	3004	1/1	0.97	0.13	66,66,66,66	0
59	MG	1H	3175	1/1	0.97	0.10	34,34,34,34	0
59	MG	1H	3421	1/1	0.97	0.05	80,80,80,80	0
59	MG	14	3125	1/1	0.97	0.10	45,45,45,45	0
59	MG	14	3227	1/1	0.97	0.12	41,41,41,41	0
59	MG	1H	3424	1/1	0.97	0.06	84,84,84,84	0
58	K	1G	1614	1/1	0.97	0.06	100,100,100,100	0
59	MG	14	3230	1/1	0.97	0.13	44,44,44,44	0
58	K	1H	3094	1/1	0.97	0.05	65,65,65,65	0
59	MG	1H	3432	1/1	0.97	0.11	73,73,73,73	0
59	MG	13	1725	1/1	0.97	0.04	116,116,116,116	0
59	MG	1H	3438	1/1	0.97	0.07	36,36,36,36	0
59	MG	1H	3447	1/1	0.97	0.04	98,98,98,98	0
59	MG	14	3236	1/1	0.97	0.11	41,41,41,41	0
59	MG	13	1728	1/1	0.97	0.04	106,106,106,106	0
59	MG	1H	3454	1/1	0.97	0.06	42,42,42,42	0
59	MG	1H	3266	1/1	0.97	0.05	40,40,40,40	0
58	K	1H	3083	1/1	0.97	0.05	43,43,43,43	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	1H	3473	1/1	0.97	0.08	27,27,27,27	0
58	K	1H	3077	1/1	0.97	0.05	49,49,49,49	0
59	MG	13	1736	1/1	0.97	0.05	100,100,100,100	0
59	MG	1H	3226	1/1	0.97	0.04	42,42,42,42	0
58	K	14	3071	1/1	0.97	0.09	75,75,75,75	0
59	MG	1H	3479	1/1	0.97	0.07	47,47,47,47	0
59	MG	1H	3480	1/1	0.97	0.06	58,58,58,58	0
59	MG	13	1694	1/1	0.97	0.09	43,43,43,43	0
59	MG	1H	3144	1/1	0.97	0.08	38,38,38,38	0
59	MG	13	1672	1/1	0.97	0.11	24,24,24,24	0
59	MG	1H	3487	1/1	0.97	0.04	85,85,85,85	0
59	MG	14	3152	1/1	0.97	0.10	69,69,69,69	0
59	MG	1H	3488	1/1	0.97	0.05	83,83,83,83	0
59	MG	1H	3490	1/1	0.97	0.06	53,53,53,53	0
59	MG	13	1742	1/1	0.97	0.05	94,94,94,94	0
59	MG	14	3156	1/1	0.97	0.12	55,55,55,55	0
59	MG	1H	3232	1/1	0.97	0.21	40,40,40,40	0
59	MG	14	3259	1/1	0.97	0.05	45,45,45,45	0
59	MG	14	3158	1/1	0.97	0.12	35,35,35,35	0
59	MG	1H	3495	1/1	0.97	0.12	80,80,80,80	0
59	MG	13	1744	1/1	0.97	0.04	107,107,107,107	0
59	MG	1H	3500	1/1	0.97	0.08	45,45,45,45	0
59	MG	14	3163	1/1	0.97	0.06	48,48,48,48	0
59	MG	14	3164	1/1	0.97	0.08	70,70,70,70	0
58	K	1H	3121	1/1	0.97	0.03	66,66,66,66	0
59	MG	14	3166	1/1	0.97	0.14	37,37,37,37	0
59	MG	1H	3509	1/1	0.97	0.05	101,101,101,101	0
59	MG	1H	3511	1/1	0.97	0.05	93,93,93,93	0
59	MG	1H	3512	1/1	0.97	0.07	61,61,61,61	0
59	MG	14	3273	1/1	0.97	0.06	46,46,46,46	0
59	MG	14	3274	1/1	0.97	0.07	30,30,30,30	0
59	MG	14	3276	1/1	0.97	0.09	44,44,44,44	0
59	MG	1H	3513	1/1	0.97	0.05	105,105,105,105	0
58	K	14	3020	1/1	0.97	0.04	72,72,72,72	0
59	MG	14	3288	1/1	0.97	0.15	79,79,79,79	0
59	MG	1H	3195	1/1	0.97	0.12	51,51,51,51	0
59	MG	14	3291	1/1	0.97	0.16	64,64,64,64	0
59	MG	1G	1684	1/1	0.97	0.07	75,75,75,75	0
59	MG	14	3297	1/1	0.97	0.05	44,44,44,44	0
59	MG	14	3299	1/1	0.97	0.06	57,57,57,57	0
59	MG	1H	3516	1/1	0.97	0.10	60,60,60,60	0
59	MG	14	3310	1/1	0.97	0.07	27,27,27,27	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(Å ²)	Q<0.9
59	MG	14	3312	1/1	0.97	0.09	87,87,87,87	0
59	MG	1H	3517	1/1	0.97	0.05	55,55,55,55	0
58	K	1H	3104	1/1	0.97	0.04	53,53,53,53	0
59	MG	14	3320	1/1	0.97	0.09	90,90,90,90	0
59	MG	14	3322	1/1	0.97	0.04	93,93,93,93	0
59	MG	14	3323	1/1	0.97	0.06	47,47,47,47	0
59	MG	1H	3521	1/1	0.97	0.05	113,113,113,113	0
59	MG	1G	1695	1/1	0.97	0.04	97,97,97,97	0
59	MG	14	3327	1/1	0.97	0.06	66,66,66,66	0
59	MG	14	3328	1/1	0.97	0.05	66,66,66,66	0
59	MG	1H	3198	1/1	0.97	0.10	82,82,82,82	0
59	MG	1G	1697	1/1	0.97	0.04	81,81,81,81	0
59	MG	1G	1699	1/1	0.97	0.07	101,101,101,101	0
59	MG	1H	3524	1/1	0.97	0.06	35,35,35,35	0
58	K	1H	3070	1/1	0.97	0.04	54,54,54,54	0
59	MG	1G	1706	1/1	0.97	0.04	96,96,96,96	0
59	MG	1G	1707	1/1	0.97	0.05	120,120,120,120	0
59	MG	1H	3329	1/1	0.97	0.14	73,73,73,73	0
58	K	1G	1607	1/1	0.97	0.03	93,93,93,93	0
59	MG	14	3359	1/1	0.97	0.10	94,94,94,94	0
59	MG	14	3361	1/1	0.97	0.05	71,71,71,71	0
58	K	14	3007	1/1	0.97	0.21	52,52,52,52	0
59	MG	14	3371	1/1	0.97	0.08	61,61,61,61	0
59	MG	14	3372	1/1	0.97	0.05	73,73,73,73	0
59	MG	13	1657	1/1	0.97	0.13	26,26,26,26	0
59	MG	1G	1720	1/1	0.97	0.05	106,106,106,106	0
58	K	11	301	1/1	0.97	0.04	47,47,47,47	0
59	MG	1G	1724	1/1	0.97	0.05	109,109,109,109	0
59	MG	21	301	1/1	0.97	0.05	34,34,34,34	0
59	MG	1H	3123	1/1	0.97	0.15	20,20,20,20	0
59	MG	1H	3337	1/1	0.97	0.07	37,37,37,37	0
59	MG	14	3383	1/1	0.97	0.07	71,71,71,71	0
59	MG	1H	3339	1/1	0.97	0.06	33,33,33,33	0
59	MG	1H	3340	1/1	0.97	0.07	29,29,29,29	0
59	MG	1H	3206	1/1	0.97	0.09	35,35,35,35	0
59	MG	1H	3344	1/1	0.97	0.06	21,21,21,21	0
59	MG	G8	201	1/1	0.97	0.06	89,89,89,89	0
59	MG	1H	3124	1/1	0.97	0.12	28,28,28,28	0
59	MG	14	3393	1/1	0.97	0.22	81,81,81,81	0
59	MG	13	1659	1/1	0.97	0.07	54,54,54,54	0
59	MG	1H	3351	1/1	0.97	0.06	33,33,33,33	0
59	MG	14	3397	1/1	0.97	0.05	46,46,46,46	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
58	K	21	302	1/1	0.97	0.06	66,66,66,66	0
58	K	14	3062	1/1	0.97	0.07	65,65,65,65	0
59	MG	1H	3250	1/1	0.97	0.05	47,47,47,47	0
59	MG	1H	3251	1/1	0.97	0.09	52,52,52,52	0
59	MG	1G	1631	1/1	0.97	0.10	49,49,49,49	0
59	MG	1H	3128	1/1	0.97	0.14	40,40,40,40	0
59	MG	13	1639	1/1	0.97	0.07	62,62,62,62	0
59	MG	1J	205	1/1	0.97	0.05	104,104,104,104	0
59	MG	1H	3130	1/1	0.97	0.14	28,28,28,28	0
58	K	1H	3098	1/1	0.97	0.07	67,67,67,67	0
59	MG	1H	3390	1/1	0.97	0.07	39,39,39,39	0
59	MG	1H	3303	1/1	0.97	0.15	41,41,41,41	0
59	MG	1H	3395	1/1	0.98	0.13	56,56,56,56	0
59	MG	16	208	1/1	0.98	0.04	63,63,63,63	0
59	MG	1H	3396	1/1	0.98	0.05	35,35,35,35	0
59	MG	1H	3397	1/1	0.98	0.06	29,29,29,29	0
59	MG	1H	3166	1/1	0.98	0.10	17,17,17,17	0
58	K	1H	3062	1/1	0.98	0.07	42,42,42,42	0
59	MG	1H	3406	1/1	0.98	0.04	95,95,95,95	0
59	MG	1H	3408	1/1	0.98	0.05	60,60,60,60	0
59	MG	1H	3410	1/1	0.98	0.04	86,86,86,86	0
59	MG	1H	3414	1/1	0.98	0.05	89,89,89,89	0
58	K	1H	3049	1/1	0.98	0.05	66,66,66,66	0
59	MG	I8	102	1/1	0.98	0.06	72,72,72,72	0
59	MG	J8	101	1/1	0.98	0.06	49,49,49,49	0
59	MG	1H	3416	1/1	0.98	0.08	59,59,59,59	0
59	MG	1H	3169	1/1	0.98	0.07	43,43,43,43	0
59	MG	1H	3420	1/1	0.98	0.05	95,95,95,95	0
59	MG	13	1740	1/1	0.98	0.04	105,105,105,105	0
58	K	1H	3087	1/1	0.98	0.03	76,76,76,76	0
58	K	1H	3078	1/1	0.98	0.09	61,61,61,61	0
59	MG	13	1743	1/1	0.98	0.04	105,105,105,105	0
59	MG	1H	3431	1/1	0.98	0.05	85,85,85,85	0
59	MG	14	3117	1/1	0.98	0.07	42,42,42,42	0
58	K	14	3038	1/1	0.98	0.11	63,63,63,63	0
59	MG	1H	3434	1/1	0.98	0.15	68,68,68,68	0
59	MG	13	1745	1/1	0.98	0.09	78,78,78,78	0
59	MG	13	1746	1/1	0.98	0.05	93,93,93,93	0
59	MG	14	3122	1/1	0.98	0.07	49,49,49,49	0
59	MG	1H	3439	1/1	0.98	0.06	55,55,55,55	0
59	MG	1H	3440	1/1	0.98	0.11	63,63,63,63	0
59	MG	1H	3441	1/1	0.98	0.10	51,51,51,51	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	14	3128	1/1	0.98	0.08	35,35,35,35	0
59	MG	1H	3444	1/1	0.98	0.04	94,94,94,94	0
59	MG	14	3130	1/1	0.98	0.12	45,45,45,45	0
59	MG	1H	3446	1/1	0.98	0.04	92,92,92,92	0
59	MG	13	1747	1/1	0.98	0.03	91,91,91,91	0
59	MG	1H	3449	1/1	0.98	0.06	57,57,57,57	0
59	MG	13	1748	1/1	0.98	0.05	104,104,104,104	0
59	MG	1H	3330	1/1	0.98	0.04	45,45,45,45	0
59	MG	1H	3455	1/1	0.98	0.20	55,55,55,55	0
59	MG	1H	3457	1/1	0.98	0.04	35,35,35,35	0
59	MG	1H	3459	1/1	0.98	0.08	50,50,50,50	0
59	MG	1H	3460	1/1	0.98	0.04	62,62,62,62	0
59	MG	13	1644	1/1	0.98	0.04	49,49,49,49	0
59	MG	1H	3462	1/1	0.98	0.04	46,46,46,46	0
59	MG	1H	3465	1/1	0.98	0.06	39,39,39,39	0
59	MG	1H	3256	1/1	0.98	0.08	47,47,47,47	0
59	MG	1H	3467	1/1	0.98	0.05	67,67,67,67	0
59	MG	1H	3468	1/1	0.98	0.04	50,50,50,50	0
59	MG	1H	3470	1/1	0.98	0.10	45,45,45,45	0
59	MG	1H	3472	1/1	0.98	0.11	71,71,71,71	0
58	K	1H	3115	1/1	0.98	0.04	50,50,50,50	0
59	MG	14	3272	1/1	0.98	0.09	26,26,26,26	0
58	K	13	1619	1/1	0.98	0.03	77,77,77,77	0
58	K	1H	3065	1/1	0.98	0.06	43,43,43,43	0
59	MG	1H	3185	1/1	0.98	0.06	41,41,41,41	0
59	MG	1H	3338	1/1	0.98	0.06	35,35,35,35	0
59	MG	1H	3478	1/1	0.98	0.04	38,38,38,38	0
59	MG	14	3280	1/1	0.98	0.05	48,48,48,48	0
59	MG	14	3281	1/1	0.98	0.06	45,45,45,45	0
59	MG	14	3283	1/1	0.98	0.07	36,36,36,36	0
59	MG	14	3286	1/1	0.98	0.04	47,47,47,47	0
59	MG	14	3287	1/1	0.98	0.07	50,50,50,50	0
58	K	13	1611	1/1	0.98	0.03	79,79,79,79	0
59	MG	13	1713	1/1	0.98	0.04	83,83,83,83	0
59	MG	13	1649	1/1	0.98	0.10	101,101,101,101	0
59	MG	1H	3343	1/1	0.98	0.05	35,35,35,35	0
59	MG	14	3294	1/1	0.98	0.04	77,77,77,77	0
59	MG	13	1718	1/1	0.98	0.07	89,89,89,89	0
59	MG	1H	3486	1/1	0.98	0.05	54,54,54,54	0
59	MG	14	3302	1/1	0.98	0.04	52,52,52,52	0
59	MG	14	3303	1/1	0.98	0.05	73,73,73,73	0
59	MG	14	3304	1/1	0.98	0.05	38,38,38,38	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	14	3305	1/1	0.98	0.08	69,69,69,69	0
59	MG	14	3306	1/1	0.98	0.06	49,49,49,49	0
59	MG	13	1719	1/1	0.98	0.05	112,112,112,112	0
59	MG	14	3309	1/1	0.98	0.04	85,85,85,85	0
59	MG	1H	3348	1/1	0.98	0.05	34,34,34,34	0
59	MG	1H	3349	1/1	0.98	0.05	28,28,28,28	0
59	MG	14	3313	1/1	0.98	0.05	89,89,89,89	0
59	MG	14	3314	1/1	0.98	0.05	91,91,91,91	0
59	MG	1H	3155	1/1	0.98	0.15	35,35,35,35	0
59	MG	13	1723	1/1	0.98	0.03	109,109,109,109	0
59	MG	14	3318	1/1	0.98	0.12	68,68,68,68	0
59	MG	1H	3494	1/1	0.98	0.04	78,78,78,78	0
59	MG	14	3168	1/1	0.98	0.04	37,37,37,37	0
58	K	19	301	1/1	0.98	0.15	61,61,61,61	0
58	K	14	3043	1/1	0.98	0.03	71,71,71,71	0
59	MG	1H	3499	1/1	0.98	0.12	68,68,68,68	0
59	MG	1H	3197	1/1	0.98	0.09	50,50,50,50	0
59	MG	1G	1679	1/1	0.98	0.07	64,64,64,64	0
59	MG	14	3331	1/1	0.98	0.07	58,58,58,58	0
59	MG	1G	1681	1/1	0.98	0.04	49,49,49,49	0
59	MG	14	3333	1/1	0.98	0.06	39,39,39,39	0
59	MG	14	3335	1/1	0.98	0.11	50,50,50,50	0
59	MG	1H	3501	1/1	0.98	0.03	78,78,78,78	0
59	MG	14	3339	1/1	0.98	0.04	77,77,77,77	0
59	MG	1H	3503	1/1	0.98	0.06	63,63,63,63	0
59	MG	1H	3504	1/1	0.98	0.13	85,85,85,85	0
59	MG	1H	3506	1/1	0.98	0.04	65,65,65,65	0
59	MG	1G	1688	1/1	0.98	0.04	82,82,82,82	0
59	MG	14	3180	1/1	0.98	0.15	40,40,40,40	0
59	MG	1G	1689	1/1	0.98	0.04	90,90,90,90	0
59	MG	14	3351	1/1	0.98	0.04	61,61,61,61	0
59	MG	14	3354	1/1	0.98	0.08	70,70,70,70	0
59	MG	14	3356	1/1	0.98	0.13	84,84,84,84	0
59	MG	13	1729	1/1	0.98	0.05	68,68,68,68	0
59	MG	1H	3370	1/1	0.98	0.06	38,38,38,38	0
59	MG	1G	1693	1/1	0.98	0.04	89,89,89,89	0
59	MG	14	3362	1/1	0.98	0.04	69,69,69,69	0
59	MG	14	3364	1/1	0.98	0.15	71,71,71,71	0
59	MG	1H	3371	1/1	0.98	0.04	35,35,35,35	0
59	MG	14	3368	1/1	0.98	0.10	40,40,40,40	0
59	MG	1H	3372	1/1	0.98	0.09	77,77,77,77	0
58	K	1H	3074	1/1	0.98	0.02	53,53,53,53	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	14	3373	1/1	0.98	0.06	84,84,84,84	0
59	MG	14	3374	1/1	0.98	0.08	96,96,96,96	0
59	MG	1G	1698	1/1	0.98	0.08	103,103,103,103	0
59	MG	1H	3374	1/1	0.98	0.07	70,70,70,70	0
59	MG	1G	1700	1/1	0.98	0.07	108,108,108,108	0
59	MG	1H	3375	1/1	0.98	0.04	69,69,69,69	0
59	MG	1G	1703	1/1	0.98	0.04	93,93,93,93	0
58	K	14	3045	1/1	0.98	0.07	71,71,71,71	0
59	MG	1H	3380	1/1	0.98	0.05	61,61,61,61	0
59	MG	14	3382	1/1	0.98	0.07	77,77,77,77	0
59	MG	1H	3385	1/1	0.98	0.07	69,69,69,69	0
59	MG	14	3384	1/1	0.98	0.05	108,108,108,108	0
59	MG	1G	1709	1/1	0.98	0.05	102,102,102,102	0
59	MG	1H	3519	1/1	0.98	0.08	56,56,56,56	0
59	MG	1H	3520	1/1	0.98	0.05	70,70,70,70	0
59	MG	1G	1712	1/1	0.98	0.04	106,106,106,106	0
59	MG	14	3390	1/1	0.98	0.05	78,78,78,78	0
59	MG	1G	1713	1/1	0.98	0.05	80,80,80,80	0
59	MG	1G	1714	1/1	0.98	0.07	82,82,82,82	0
59	MG	1G	1715	1/1	0.98	0.06	92,92,92,92	0
59	MG	1H	3386	1/1	0.98	0.09	67,67,67,67	0
59	MG	1G	1717	1/1	0.98	0.05	88,88,88,88	0
59	MG	14	3396	1/1	0.98	0.05	51,51,51,51	0
59	MG	1H	3162	1/1	0.98	0.07	21,21,21,21	0
59	MG	1G	1719	1/1	0.98	0.05	101,101,101,101	0
59	MG	1H	3523	1/1	0.98	0.04	34,34,34,34	0
59	MG	14	3400	1/1	0.98	0.04	92,92,92,92	0
59	MG	1G	1721	1/1	0.98	0.04	88,88,88,88	0
59	MG	13	1735	1/1	0.98	0.04	57,57,57,57	0
59	MG	14	3405	1/1	0.98	0.09	45,45,45,45	0
59	MG	1G	1723	1/1	0.98	0.03	85,85,85,85	0
59	MG	1H	3277	1/1	0.98	0.11	18,18,18,18	0
59	MG	1H	3392	1/1	0.98	0.06	32,32,32,32	0
59	MG	1J	204	1/1	0.98	0.06	115,115,115,115	0
59	MG	1G	1726	1/1	0.98	0.04	98,98,98,98	0
58	K	1H	3056	1/1	0.98	0.04	43,43,43,43	0
59	MG	14	3216	1/1	0.98	0.07	42,42,42,42	0
59	MG	32	302	1/1	0.98	0.05	92,92,92,92	0
59	MG	13	1737	1/1	0.98	0.04	90,90,90,90	0
59	MG	1H	3270	1/1	0.99	0.09	37,37,37,37	0
59	MG	14	3271	1/1	0.99	0.04	52,52,52,52	0
59	MG	1H	3442	1/1	0.99	0.05	80,80,80,80	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	16	209	1/1	0.99	0.07	75,75,75,75	0
59	MG	1G	1705	1/1	0.99	0.02	105,105,105,105	0
59	MG	14	3275	1/1	0.99	0.04	58,58,58,58	0
58	K	1H	3059	1/1	0.99	0.05	38,38,38,38	0
59	MG	1H	3445	1/1	0.99	0.08	42,42,42,42	0
59	MG	1G	1708	1/1	0.99	0.03	69,69,69,69	0
59	MG	14	3279	1/1	0.99	0.03	46,46,46,46	0
59	MG	1H	3299	1/1	0.99	0.10	34,34,34,34	0
59	MG	1H	3171	1/1	0.99	0.10	28,28,28,28	0
59	MG	14	3282	1/1	0.99	0.04	50,50,50,50	0
59	MG	1H	3448	1/1	0.99	0.04	61,61,61,61	0
59	MG	14	3285	1/1	0.99	0.04	43,43,43,43	0
59	MG	13	1730	1/1	0.99	0.05	79,79,79,79	0
59	MG	1H	3152	1/1	0.99	0.07	37,37,37,37	0
59	MG	1H	3451	1/1	0.99	0.10	63,63,63,63	0
59	MG	1H	3453	1/1	0.99	0.05	67,67,67,67	0
59	MG	14	3290	1/1	0.99	0.04	48,48,48,48	0
59	MG	I8	101	1/1	0.99	0.11	48,48,48,48	0
59	MG	14	3292	1/1	0.99	0.04	49,49,49,49	0
59	MG	1H	3376	1/1	0.99	0.04	29,29,29,29	0
59	MG	13	1731	1/1	0.99	0.06	92,92,92,92	0
59	MG	14	3295	1/1	0.99	0.08	51,51,51,51	0
59	MG	1H	3456	1/1	0.99	0.03	38,38,38,38	0
59	MG	14	3298	1/1	0.99	0.06	54,54,54,54	0
59	MG	1H	3379	1/1	0.99	0.03	67,67,67,67	0
59	MG	14	3300	1/1	0.99	0.05	41,41,41,41	0
59	MG	1H	3458	1/1	0.99	0.05	89,89,89,89	0
59	MG	1H	3331	1/1	0.99	0.24	55,55,55,55	0
59	MG	1H	3381	1/1	0.99	0.03	59,59,59,59	0
59	MG	1H	3382	1/1	0.99	0.06	36,36,36,36	0
59	MG	1H	3384	1/1	0.99	0.06	60,60,60,60	0
59	MG	14	3307	1/1	0.99	0.04	51,51,51,51	0
59	MG	1H	3463	1/1	0.99	0.03	40,40,40,40	0
59	MG	1H	3464	1/1	0.99	0.04	19,19,19,19	0
59	MG	13	1715	1/1	0.99	0.06	75,75,75,75	0
59	MG	14	3311	1/1	0.99	0.07	56,56,56,56	0
59	MG	13	1716	1/1	0.99	0.04	66,66,66,66	0
59	MG	13	1734	1/1	0.99	0.07	61,61,61,61	0
59	MG	13	1717	1/1	0.99	0.08	66,66,66,66	0
59	MG	1H	3469	1/1	0.99	0.06	87,87,87,87	0
59	MG	14	3316	1/1	0.99	0.09	53,53,53,53	0
58	K	1H	3051	1/1	0.99	0.03	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	1H	3471	1/1	0.99	0.05	30,30,30,30	0
59	MG	14	3319	1/1	0.99	0.04	59,59,59,59	0
59	MG	1H	3391	1/1	0.99	0.07	31,31,31,31	0
59	MG	14	3321	1/1	0.99	0.05	51,51,51,51	0
58	K	13	1617	1/1	0.99	0.09	72,72,72,72	0
59	MG	1H	3255	1/1	0.99	0.05	36,36,36,36	0
59	MG	1H	3205	1/1	0.99	0.06	34,34,34,34	0
59	MG	14	3204	1/1	0.99	0.06	44,44,44,44	0
59	MG	14	3326	1/1	0.99	0.03	45,45,45,45	0
59	MG	13	1720	1/1	0.99	0.05	73,73,73,73	0
59	MG	1H	3182	1/1	0.99	0.02	47,47,47,47	0
59	MG	14	3329	1/1	0.99	0.13	70,70,70,70	0
59	MG	1H	3342	1/1	0.99	0.04	50,50,50,50	0
59	MG	1H	3398	1/1	0.99	0.08	74,74,74,74	0
59	MG	13	1721	1/1	0.99	0.07	68,68,68,68	0
59	MG	14	3334	1/1	0.99	0.05	48,48,48,48	0
59	MG	1H	3400	1/1	0.99	0.08	46,46,46,46	0
59	MG	14	3336	1/1	0.99	0.08	47,47,47,47	0
59	MG	14	3337	1/1	0.99	0.10	66,66,66,66	0
59	MG	14	3109	1/1	0.99	0.05	36,36,36,36	0
59	MG	1H	3402	1/1	0.99	0.04	63,63,63,63	0
59	MG	14	3111	1/1	0.99	0.02	44,44,44,44	0
59	MG	1H	3403	1/1	0.99	0.07	59,59,59,59	0
59	MG	14	3342	1/1	0.99	0.07	54,54,54,54	0
59	MG	14	3343	1/1	0.99	0.03	76,76,76,76	0
59	MG	13	1722	1/1	0.99	0.12	77,77,77,77	0
59	MG	14	3345	1/1	0.99	0.09	89,89,89,89	0
59	MG	14	3346	1/1	0.99	0.04	76,76,76,76	0
59	MG	1H	3345	1/1	0.99	0.04	24,24,24,24	0
59	MG	1H	3407	1/1	0.99	0.06	65,65,65,65	0
59	MG	14	3349	1/1	0.99	0.03	77,77,77,77	0
59	MG	1H	3489	1/1	0.99	0.08	76,76,76,76	0
59	MG	1H	3346	1/1	0.99	0.05	27,27,27,27	0
59	MG	14	3352	1/1	0.99	0.06	64,64,64,64	0
59	MG	14	3353	1/1	0.99	0.06	65,65,65,65	0
59	MG	1H	3491	1/1	0.99	0.10	51,51,51,51	0
59	MG	14	3355	1/1	0.99	0.08	64,64,64,64	0
59	MG	1H	3409	1/1	0.99	0.10	64,64,64,64	0
58	K	14	3016	1/1	0.99	0.04	72,72,72,72	0
59	MG	14	3358	1/1	0.99	0.04	35,35,35,35	0
59	MG	1H	3412	1/1	0.99	0.12	85,85,85,85	0
59	MG	14	3360	1/1	0.99	0.03	64,64,64,64	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	1H	3413	1/1	0.99	0.08	42,42,42,42	0
59	MG	14	3123	1/1	0.99	0.02	47,47,47,47	0
59	MG	14	3363	1/1	0.99	0.03	85,85,85,85	0
59	MG	1H	3496	1/1	0.99	0.03	64,64,64,64	0
59	MG	14	3365	1/1	0.99	0.06	80,80,80,80	0
59	MG	1H	3497	1/1	0.99	0.04	83,83,83,83	0
59	MG	14	3367	1/1	0.99	0.05	52,52,52,52	0
59	MG	13	1724	1/1	0.99	0.05	80,80,80,80	0
59	MG	14	3369	1/1	0.99	0.12	43,43,43,43	0
59	MG	14	3370	1/1	0.99	0.04	56,56,56,56	0
59	MG	14	3127	1/1	0.99	0.05	41,41,41,41	0
59	MG	1H	3187	1/1	0.99	0.06	37,37,37,37	0
58	K	1H	3080	1/1	0.99	0.02	51,51,51,51	0
59	MG	1H	3417	1/1	0.99	0.05	44,44,44,44	0
59	MG	1H	3502	1/1	0.99	0.03	94,94,94,94	0
59	MG	1H	3418	1/1	0.99	0.04	58,58,58,58	0
59	MG	14	3133	1/1	0.99	0.10	50,50,50,50	0
59	MG	1H	3189	1/1	0.99	0.09	46,46,46,46	0
59	MG	1H	3505	1/1	0.99	0.03	34,34,34,34	0
59	MG	14	3238	1/1	0.99	0.04	42,42,42,42	0
59	MG	1H	3146	1/1	0.99	0.12	24,24,24,24	0
59	MG	1H	3507	1/1	0.99	0.06	45,45,45,45	0
59	MG	1H	3354	1/1	0.99	0.02	70,70,70,70	0
59	MG	1H	3422	1/1	0.99	0.05	43,43,43,43	0
59	MG	1H	3510	1/1	0.99	0.08	80,80,80,80	0
59	MG	1H	3423	1/1	0.99	0.04	33,33,33,33	0
59	MG	1H	3355	1/1	0.99	0.03	32,32,32,32	0
59	MG	1H	3425	1/1	0.99	0.06	42,42,42,42	0
59	MG	1G	1680	1/1	0.99	0.03	90,90,90,90	0
59	MG	1H	3358	1/1	0.99	0.04	43,43,43,43	0
59	MG	1G	1682	1/1	0.99	0.04	60,60,60,60	0
59	MG	13	1726	1/1	0.99	0.02	122,122,122,122	0
59	MG	1H	3428	1/1	0.99	0.02	55,55,55,55	0
59	MG	1H	3429	1/1	0.99	0.05	63,63,63,63	0
59	MG	1H	3430	1/1	0.99	0.04	79,79,79,79	0
59	MG	1G	1687	1/1	0.99	0.04	93,93,93,93	0
59	MG	1H	3360	1/1	0.99	0.04	39,39,39,39	0
59	MG	1H	3361	1/1	0.99	0.07	32,32,32,32	0
59	MG	1H	3433	1/1	0.99	0.13	44,44,44,44	0
59	MG	1H	3362	1/1	0.99	0.03	33,33,33,33	0
59	MG	1G	1692	1/1	0.99	0.06	97,97,97,97	0
59	MG	14	3403	1/1	0.99	0.03	75,75,75,75	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
59	MG	14	3404	1/1	0.99	0.04	34,34,34,34	0
59	MG	1H	3435	1/1	0.99	0.09	72,72,72,72	0
59	MG	1G	1694	1/1	0.99	0.04	62,62,62,62	0
59	MG	1H	3436	1/1	0.99	0.04	33,33,33,33	0
59	MG	1H	3525	1/1	0.99	0.06	46,46,46,46	0
59	MG	1J	203	1/1	0.99	0.04	105,105,105,105	0
59	MG	14	3161	1/1	0.99	0.13	44,44,44,44	0
59	MG	13	1727	1/1	0.99	0.03	47,47,47,47	0
59	MG	1H	3364	1/1	0.99	0.04	28,28,28,28	0
59	MG	1H	3367	1/1	0.99	0.07	61,61,61,61	0
58	K	1H	3068	1/1	0.99	0.08	37,37,37,37	0
59	MG	1G	1701	1/1	0.99	0.04	108,108,108,108	0
60	SF4	3E	302	8/8	0.99	0.05	78,87,103,104	0
60	SF4	32	303	8/8	0.99	0.07	73,95,102,116	0
61	ZN	5I	102	1/1	0.99	0.04	110,110,110,110	0
61	ZN	5A	101	1/1	0.99	0.04	115,115,115,115	0
59	MG	1H	3378	1/1	1.00	0.02	30,30,30,30	0
59	MG	1H	3443	1/1	1.00	0.08	61,61,61,61	0
59	MG	14	3301	1/1	1.00	0.06	53,53,53,53	0
59	MG	1H	3387	1/1	1.00	0.03	37,37,37,37	0
59	MG	14	3284	1/1	1.00	0.02	39,39,39,39	0
59	MG	1H	3366	1/1	1.00	0.01	44,44,44,44	0
59	MG	1H	3356	1/1	1.00	0.04	31,31,31,31	0
59	MG	1H	3368	1/1	1.00	0.01	20,20,20,20	0
59	MG	1H	3411	1/1	1.00	0.02	28,28,28,28	0
59	MG	1H	3357	1/1	1.00	0.02	34,34,34,34	0
59	MG	14	3388	1/1	1.00	0.07	53,53,53,53	0
59	MG	1H	3401	1/1	1.00	0.02	73,73,73,73	0
59	MG	1H	3383	1/1	1.00	0.04	54,54,54,54	0
59	MG	14	3330	1/1	1.00	0.01	48,48,48,48	0
59	MG	1H	3452	1/1	1.00	0.03	48,48,48,48	0
59	MG	1H	3353	1/1	1.00	0.07	53,53,53,53	0
59	MG	1H	3482	1/1	1.00	0.03	22,22,22,22	0
59	MG	1H	3365	1/1	1.00	0.03	40,40,40,40	0
59	MG	14	3296	1/1	1.00	0.03	44,44,44,44	0
59	MG	1H	3405	1/1	1.00	0.07	50,50,50,50	0
59	MG	1H	3485	1/1	1.00	0.03	27,27,27,27	0

6.5 Other polymers ⓘ

There are no such residues in this entry.