Full wwPDB X-ray Structure Validation Report ⓘ

Aug 1, 2016 – 11:06 PM EDT

PDB ID	:	5IB7
Title	:	Structure of T. thermophilus 70S ribosome complex with mRNA, tRNAfMet, near-cognate tRNALys with U-G mismatch in the A-site and antibiotic paromomycin
Authors	:	Rozov, A.; Demeshkina, N.; Yusupov, M.; Yusupova, G.
Deposited on	:	2016-02-22
Resolution	:	2.99 Å(reported)

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

We welcome your comments at *validation@mail.wwpdb.org*
A user guide is available at
http://wwpdb.org/validation/2016/XrayValidationReportHelp
with specific help available everywhere you see the ⓘ symbol.

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

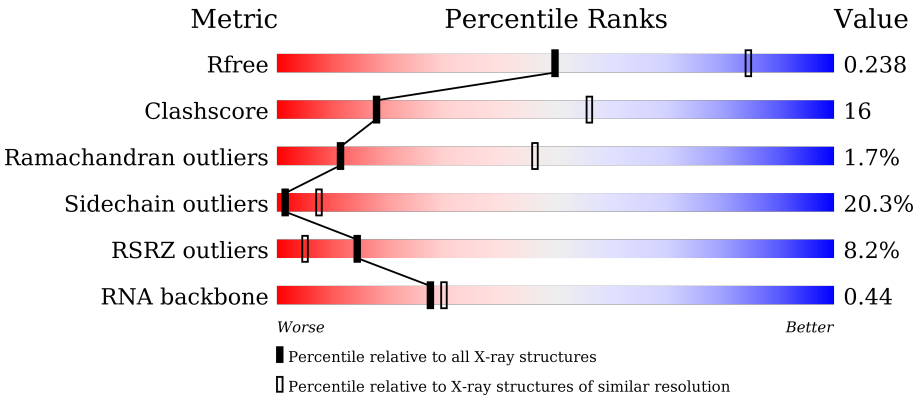
MolProbity	:	4.02b-467
Mogul	:	1.7.1 (RC1), CSD as537be (2016)
Xtriage (Phenix)	:	1.9-1692
EDS	:	rb-20027939
Percentile statistics	:	20151230.v01 (using entries in the PDB archive December 30th 2015)
Refmac	:	5.8.0135
CCP4	:	6.5.0
Ideal geometry (proteins)	:	Engh & Huber (2001)
Ideal geometry (DNA, RNA)	:	Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP)	:	rb-20027939

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 2.99 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}	91344	1578 (3.00-3.00)
Clashscore	102246	1912 (3.00-3.00)
Ramachandran outliers	100387	1853 (3.00-3.00)
Sidechain outliers	100360	1856 (3.00-3.00)
RSRZ outliers	91569	1592 (3.00-3.00)
RNA backbone	2183	1036 (3.40-2.60)

The table below summarises the geometric issues observed across the polymeric chains and their fit to the electron density. The red, orange, yellow and green segments on the lower bar indicate the fraction of residues that contain outliers for >=3, 2, 1 and 0 types of geometric quality criteria. A grey segment represents the fraction of residues that are not modelled. The numeric value for each fraction is indicated below the corresponding segment, with a dot representing fractions <=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	37% 45% 15%
1	1G	1522	% 39% 44% 14%
2	12	256	10% 34% 36% 9% 19%
2	1E	256	5% 39% 38% 12% 10%

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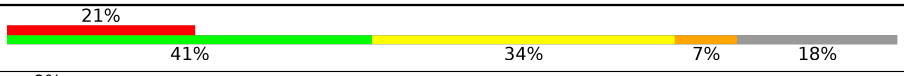
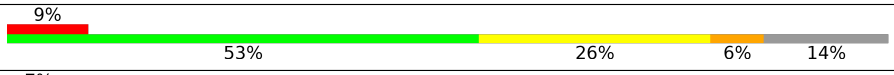
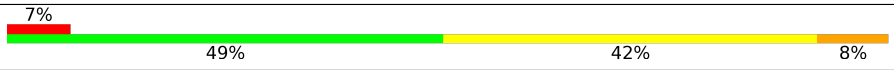
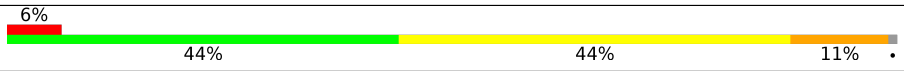
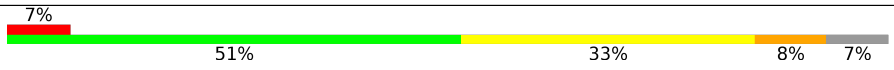
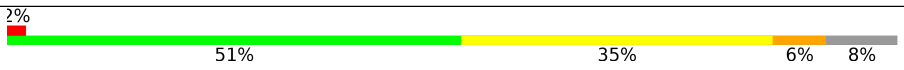
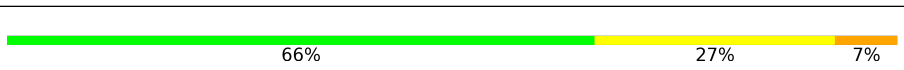
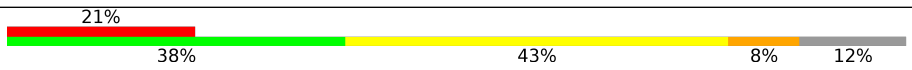
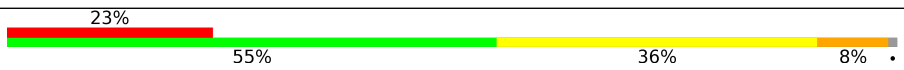
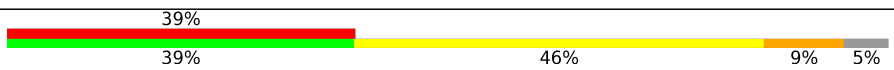
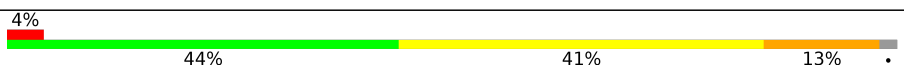
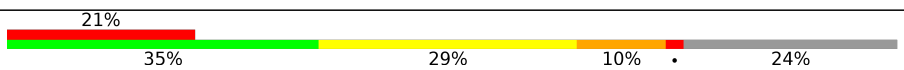
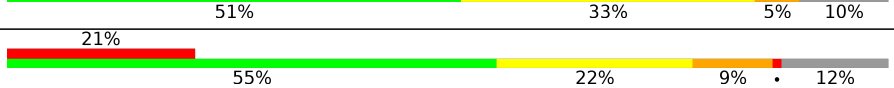
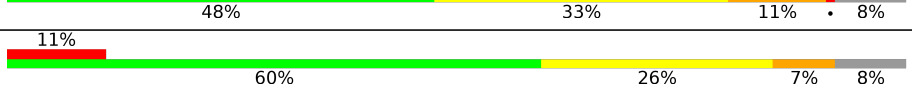
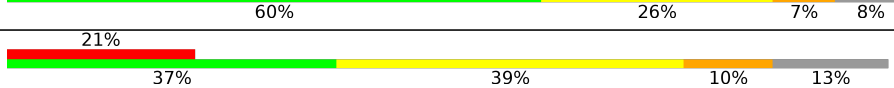
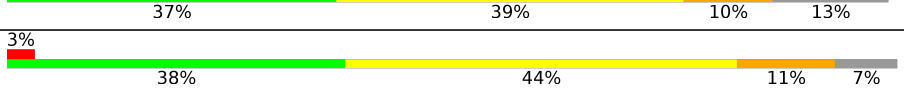
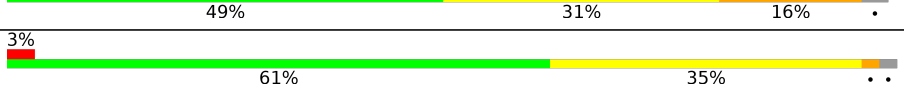
Mol	Chain	Length	Quality of chain
3	22	239	
3	2E	239	
4	32	209	
4	3E	209	
5	42	162	
5	4E	162	
6	52	101	
6	5E	101	
7	62	156	
7	6E	156	
8	72	138	
8	7E	138	
9	82	128	
9	8E	128	
10	1A	105	
10	1I	105	
11	2A	129	
11	2I	129	
12	3A	132	
12	3I	132	
13	4A	126	
13	4I	126	
14	5A	61	
14	5I	61	
15	6A	89	

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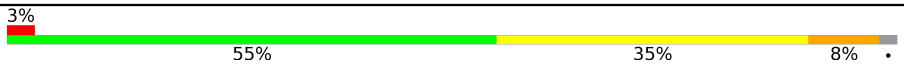
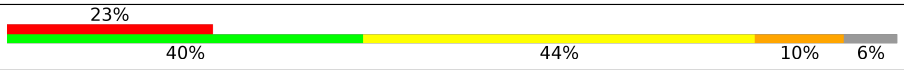
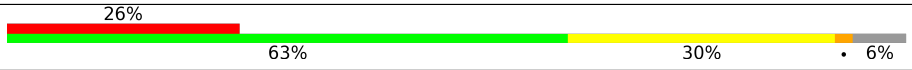
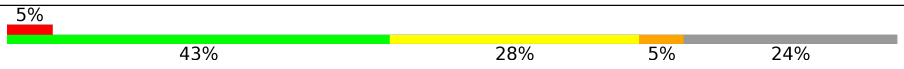
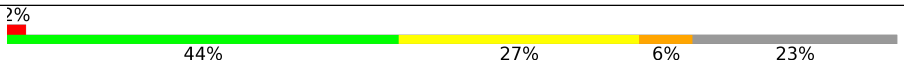
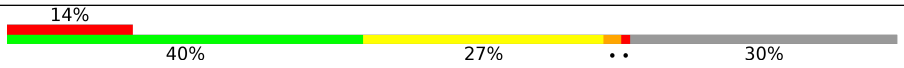
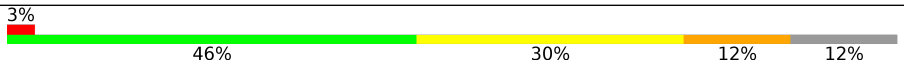
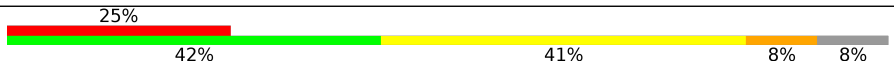
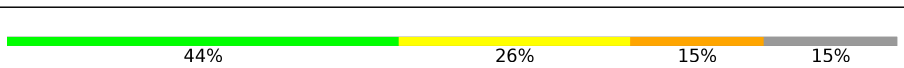
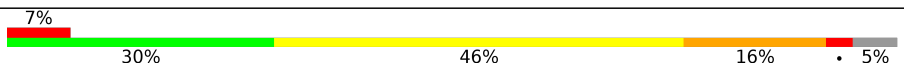
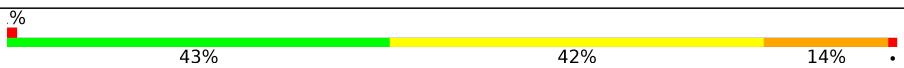
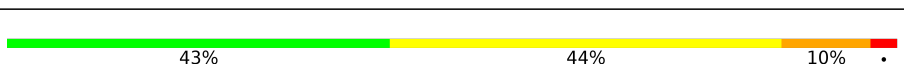
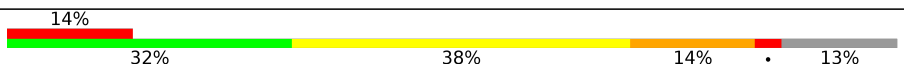
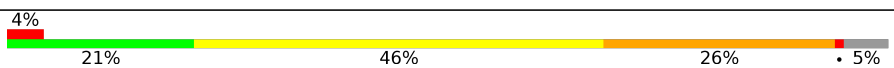
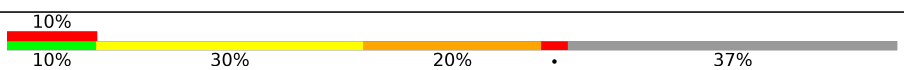
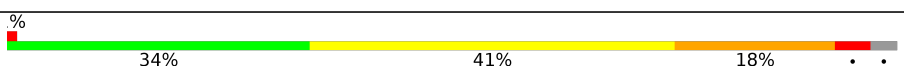
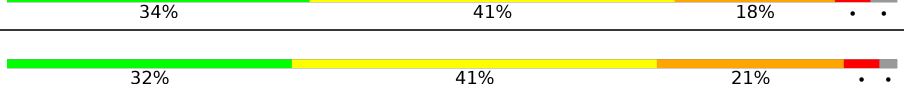
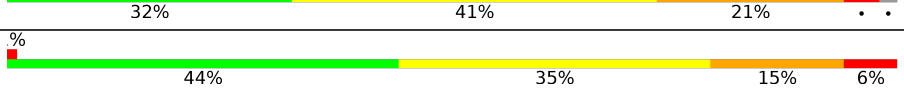
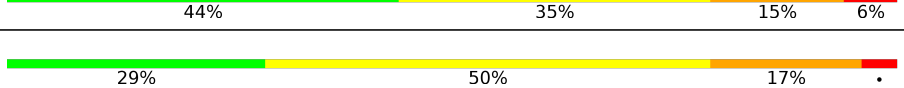
Mol	Chain	Length	Quality of chain
15	6I	89	
16	7A	88	
16	7I	88	
17	8A	105	
17	8I	105	
18	9A	88	
18	9I	88	
19	AA	93	
19	AI	93	
20	BA	106	
20	BI	106	
21	1B	27	
21	1F	27	
22	1K	76	
23	2K	77	
23	2L	77	
24	1L	76	
24	3K	76	
24	3L	76	
25	4K	30	
25	4L	30	
26	14	2917	
26	1H	2917	
27	16	122	
27	1J	122	

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Mol	Chain	Length	Quality of chain
28	71	229	9% 28% 23% 6% 42%
29	11	276	2% 53% 36% 9%
29	19	276	3% 55% 33% 10%
30	21	206	5% 44% 35% 17%
30	29	206	19% 46% 40% 12%
31	31	210	4% 51% 36% 9%
31	39	210	7% 40% 43% 13%
32	41	182	12% 51% 40% 7%
32	49	182	26% 46% 43% 10%
33	51	180	3% 52% 26% 17%
33	59	180	51% 47% 31% 14% 6%
34	61	148	11% 52% 34% 11%
34	69	148	8% 49% 38% 10%
35	15	140	18% 52% 35% 11%
35	58	140	13% 54% 32% 11%
36	25	122	11% 48% 41% 11%
36	68	122	2% 61% 34%
37	35	150	14% 43% 41% 12%
37	78	150	7% 37% 41% 15% 5%
38	45	141	19% 43% 42% 13%
38	88	141	7% 51% 35% 13%
39	55	118	5% 51% 41% 8%
39	98	118	10% 47% 40% 13%
40	65	112	20% 40% 43% 13%
40	A8	112	16% 38% 47% 13%

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Mol	Chain	Length	Quality of chain
41	75	146	5% 41% 38% 11% • 9%
41	B8	146	3% 36% 39% 16% • 8%
42	85	118	3% 56% 31% 11% •
42	C8	118	3% 58% 29% 9% • •
43	95	101	22% 56% 36% 6% ••
43	D8	101	15% 49% 39% 10% ••
44	A5	113	4% 64% 27% 7% •
44	E8	113	7% 58% 31% 8% •
45	B5	96	16% 54% 35% 8% •
45	F8	96	% 68% 23% 8% •
46	C5	110	33% 42% 36% 14% • 5%
46	G8	110	5% 46% 34% 13% • 6%
47	D5	206	34% 34% 40% 11% • 14%
47	H8	206	32% 43% 29% 8% • 17%
48	E5	85	15% 47% 34% 7% • 11%
48	I8	85	6% 49% 31% 11% 9%
49	F5	98	18% 52% 35% 9% •
49	J8	98	9% 49% 37% 9% • •
50	G5	72	7% 43% 43% 7% • •
50	K8	72	% 35% 40% 15% • 6%
51	H5	60	48% 57% 37% • •
51	L8	60	3% 52% 40% 5% •
52	M8	71	23% 35% 32% 14% • 15%
53	J5	60	12% 53% 37% • 7%
53	N8	60	43% 28% 7% • 20%

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Mol	Chain	Length	Quality of chain
54	O8	54	74% 26% 43% 11% 17%
55	L5	49	4% 65% 27%
55	P8	49	2% 53% 39%
56	M5	65	28% 52% 32% 14%
56	Q8	65	43% 35% 18%

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
57	MG	13	1601	-	-	-	X
57	MG	13	1614	-	-	-	X
57	MG	13	1623	-	-	-	X
57	MG	13	1624	-	-	-	X
57	MG	13	1627	-	-	-	X
57	MG	13	1628	-	-	-	X
57	MG	13	1629	-	-	-	X
57	MG	13	1633	-	-	-	X
57	MG	13	1635	-	-	-	X
57	MG	13	1636	-	-	-	X
57	MG	13	1645	-	-	-	X
57	MG	13	1654	-	-	-	X
57	MG	13	1659	-	-	-	X
57	MG	13	1661	-	-	-	X
57	MG	13	1668	-	-	-	X
57	MG	13	1669	-	-	-	X
57	MG	13	1670	-	-	-	X
57	MG	14	3007	-	-	-	X
57	MG	14	3008	-	-	-	X
57	MG	14	3014	-	-	-	X
57	MG	14	3015	-	-	-	X
57	MG	14	3016	-	-	-	X
57	MG	14	3023	-	-	-	X
57	MG	14	3028	-	-	-	X
57	MG	14	3030	-	-	-	X
57	MG	14	3031	-	-	-	X
57	MG	14	3034	-	-	-	X
57	MG	14	3035	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
57	MG	14	3042	-	-	-	X
57	MG	14	3046	-	-	-	X
57	MG	14	3048	-	-	-	X
57	MG	14	3054	-	-	-	X
57	MG	14	3059	-	-	-	X
57	MG	14	3061	-	-	-	X
57	MG	14	3062	-	-	-	X
57	MG	14	3064	-	-	-	X
57	MG	14	3070	-	-	-	X
57	MG	14	3072	-	-	-	X
57	MG	14	3074	-	-	-	X
57	MG	14	3076	-	-	-	X
57	MG	14	3077	-	-	-	X
57	MG	14	3080	-	-	-	X
57	MG	14	3088	-	-	-	X
57	MG	14	3089	-	-	-	X
57	MG	14	3090	-	-	-	X
57	MG	14	3092	-	-	-	X
57	MG	14	3097	-	-	-	X
57	MG	14	3099	-	-	-	X
57	MG	14	3100	-	-	-	X
57	MG	14	3107	-	-	-	X
57	MG	14	3108	-	-	-	X
57	MG	14	3114	-	-	-	X
57	MG	14	3117	-	-	-	X
57	MG	14	3118	-	-	-	X
57	MG	14	3119	-	-	-	X
57	MG	14	3122	-	-	-	X
57	MG	14	3124	-	-	-	X
57	MG	14	3125	-	-	-	X
57	MG	14	3126	-	-	-	X
57	MG	14	3134	-	-	-	X
57	MG	14	3136	-	-	-	X
57	MG	14	3139	-	-	-	X
57	MG	14	3142	-	-	-	X
57	MG	14	3145	-	-	-	X
57	MG	14	3151	-	-	-	X
57	MG	14	3171	-	-	-	X
57	MG	14	3179	-	-	-	X
57	MG	14	3181	-	-	-	X
57	MG	14	3191	-	-	-	X
57	MG	14	3194	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
57	MG	14	3198	-	-	-	X
57	MG	14	3203	-	-	-	X
57	MG	14	3205	-	-	-	X
57	MG	14	3208	-	-	-	X
57	MG	14	3224	-	-	-	X
57	MG	14	3233	-	-	-	X
57	MG	14	3236	-	-	-	X
57	MG	14	3237	-	-	-	X
57	MG	14	3248	-	-	-	X
57	MG	16	202	-	-	-	X
57	MG	16	203	-	-	-	X
57	MG	19	301	-	-	-	X
57	MG	1G	1614	-	-	-	X
57	MG	1G	1636	-	-	-	X
57	MG	1H	3002	-	-	-	X
57	MG	1H	3009	-	-	-	X
57	MG	1H	3011	-	-	-	X
57	MG	1H	3013	-	-	-	X
57	MG	1H	3015	-	-	-	X
57	MG	1H	3029	-	-	-	X
57	MG	1H	3034	-	-	-	X
57	MG	1H	3040	-	-	-	X
57	MG	1H	3041	-	-	-	X
57	MG	1H	3042	-	-	-	X
57	MG	1H	3045	-	-	-	X
57	MG	1H	3056	-	-	-	X
57	MG	1H	3059	-	-	-	X
57	MG	1H	3061	-	-	-	X
57	MG	1H	3066	-	-	-	X
57	MG	1H	3067	-	-	-	X
57	MG	1H	3074	-	-	-	X
57	MG	1H	3076	-	-	-	X
57	MG	1H	3078	-	-	-	X
57	MG	1H	3079	-	-	-	X
57	MG	1H	3082	-	-	-	X
57	MG	1H	3085	-	-	-	X
57	MG	1H	3098	-	-	-	X
57	MG	1H	3102	-	-	-	X
57	MG	1H	3105	-	-	-	X
57	MG	1H	3108	-	-	-	X
57	MG	1H	3121	-	-	-	X
57	MG	1H	3130	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
57	MG	1H	3133	-	-	-	X
57	MG	1H	3140	-	-	-	X
57	MG	1H	3143	-	-	-	X
57	MG	1H	3151	-	-	-	X
57	MG	1H	3165	-	-	-	X
57	MG	1H	3167	-	-	-	X
57	MG	1H	3172	-	-	-	X
57	MG	1H	3175	-	-	-	X
57	MG	1H	3178	-	-	-	X
57	MG	1H	3181	-	-	-	X
57	MG	1H	3186	-	-	-	X
57	MG	1H	3187	-	-	-	X
57	MG	1H	3188	-	-	-	X
57	MG	1H	3200	-	-	-	X
57	MG	1H	3203	-	-	-	X
57	MG	1H	3205	-	-	-	X
57	MG	1H	3212	-	-	-	X
57	MG	1H	3221	-	-	-	X
57	MG	1H	3235	-	-	-	X
57	MG	1H	3264	-	-	-	X
57	MG	1H	3321	-	-	-	X
57	MG	2L	101	-	-	-	X
59	SF4	32	302	-	-	X	-
61	SPE	14	3436	-	-	-	X
61	SPE	14	3437	-	-	-	X

2 Entry composition

There are 62 unique types of molecules in this entry. The entry contains 297444 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	1499	Total	C	N	O	P	0	0	0
			32223	14342	5973	10409	1499			
1	1G	1508	Total	C	N	O	P	0	0	0
			32414	14427	6005	10474	1508			

There are 6 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
13	1542	G	-	expression tag	GB 55771382
13	1543	C	-	expression tag	GB 55771382
13	1544	U	-	expression tag	GB 55771382
1G	1542	G	-	expression tag	GB 55771382
1G	1543	C	-	expression tag	GB 55771382
1G	1544	U	-	expression tag	GB 55771382

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
2	1E	231	Total	C	N	O	S	0	0	0
			1874	1199	334	336	5			
2	12	207	Total	C	N	O	S	0	0	0
			1696	1083	306	303	4			

- 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	195	Total	C	N	O	S	0	0	0
			1537	973	297	266	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
4	3E	207	Total	C	N	O	S	0	0	0
			1698	1064	338	289	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	149	Total	C	N	O	S	0	0	0
			1142	722	216	200	4			
5	42	150	Total	C	N	O	S	0	0	0
			1141	719	217	201	4			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
6	5E	100	Total	C	N	O	S	0	0	0
			837	528	154	152	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	154	Total	C	N	O	S	0	0	0
			1242	770	250	216	6			
7	62	138	Total	C	N	O	S	0	0	0
			1110	689	221	194	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	137	Total	C	N	O	S	0	0	0
			1107	700	214	191	2			

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

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
9	8E	126	Total	C	N	O	0	0	0
			1000	634	196	170			

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Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
9	82	121	Total	C	N	O	0	0	0
			953	605	186	162			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
10	1I	94	Total	C	N	O	S	0	0	0
			749	468	147	133	1			
10	1A	80	Total	C	N	O		0	0	0
			646	403	129	114				

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
11	2I	111	Total	C	N	O	S	0	0	0
			823	512	154	154	3			
11	2A	113	Total	C	N	O	S	0	0	0
			835	520	156	156	3			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
12	3I	122	Total	C	N	O	S	0	0	0
			956	603	193	159	1			
12	3A	121	Total	C	N	O	S	0	0	0
			947	597	191	158	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
13	4I	117	Total	C	N	O	S	0	0	0
			933	577	192	162	2			
13	4A	109	Total	C	N	O	S	0	0	0
			879	544	181	152	2			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
14	5I	59	Total	C	N	O	S	0	0	0
			486	309	103	70	4			
14	5A	59	Total	C	N	O	S	0	0	0
			486	309	103	70	4			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
15	6I	87	Total	C	N	O	S	0	0	0
			729	457	146	124	2			
15	6A	87	Total	C	N	O	S	0	0	0
			729	457	146	124	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	99	Total	C	N	O	S	0	0	0
			823	528	151	142	2			
17	8A	99	Total	C	N	O	S	0	0	0
			823	528	151	142	2			

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

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
18	9I	68	Total	C	N	O	0	0	0
			549	352	105	92			
18	9A	67	Total	C	N	O	0	0	0
			544	349	104	91			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
19	AI	82	Total	C	N	O	S	0	0	0
			661	422	123	114	2			
19	AA	65	Total	C	N	O	S	0	0	0
			510	324	92	92	2			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
20	BI	97	Total	C	N	O	S	0	0	0
			746	461	157	126	2			
20	BA	99	Total	C	N	O	S	0	0	0
			762	470	162	128	2			

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

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
21	1F	23	Total	C	N	O	0	0	0
			199	122	48	29			
21	1B	22	Total	C	N	O	0	0	0
			188	116	44	28			

- Molecule 22 is a RNA chain called tRNALys.

Mol	Chain	Residues	Atoms						ZeroOcc	AltConf	Trace
22	1K	72	Total	C	N	O	P	S	0	0	0
			1542	691	269	509	72	1			

- Molecule 23 is a RNA chain called tRNAfMet.

Mol	Chain	Residues	Atoms						ZeroOcc	AltConf	Trace
23	2K	77	Total	C	N	O	P	S	0	0	0
			1646	735	298	535	77	1			
23	2L	77	Total	C	N	O	P	S	0	0	0
			1646	735	298	535	77	1			

- Molecule 24 is a RNA chain called tRNALys.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
24	3K	70	Total	C	N	O	P	0	0	0
			1483	664	260	490	69			
24	1L	66	Total	C	N	O	P	0	0	0
			1401	626	244	465	66			
24	3L	72	Total	C	N	O	P	0	0	0
			1528	684	270	503	71			

- Molecule 25 is a RNA chain called mRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
25	4K	19	Total	C	N	O	P	0	0	0
			420	188	89	124	19			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
25	4L	19	Total	C	N	O	P	0	0	0
			419	188	89	123	19			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
26	1H	2860	Total	C	N	O	P	0	0	0
			61609	27418	11525	19806	2860			
26	14	2826	Total	C	N	O	P	0	0	0
			60877	27092	11393	19566	2826			

There are 14 discrepancies between the modelled and reference sequences:

Chain	Residue	Modelled	Actual	Comment	Reference
1H	161	U	UNK	conflict	GB 55771382
1H	654A	A	G	conflict	GB 55771382
1H	654E	C	G	conflict	GB 55771382
1H	654P	G	C	conflict	GB 55771382
1H	654T	A	C	conflict	GB 55771382
1H	1059B	U	G	conflict	GB 55771382
1H	1080	A	C	conflict	GB 55771382
14	158	U	UNK	conflict	GB 55771382
14	654A	A	G	conflict	GB 55771382
14	654E	C	G	conflict	GB 55771382
14	654P	G	C	conflict	GB 55771382
14	654T	A	C	conflict	GB 55771382
14	1058	U	G	conflict	GB 55771382
14	1080	A	C	conflict	GB 55771382

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

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

- Molecule 28 is a protein called 50S ribosomal protein L1.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
28	71	132	Total	C	N	O	S	0	0	0
			1027	648	193	185	1			

- Molecule 29 is a protein called 50S ribosomal protein L2.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
29	11	273	Total	C	N	O	S	0	0	0
			2120	1338	421	358	3			
29	19	274	Total	C	N	O	S	0	0	0
			2125	1341	422	359	3			

- Molecule 30 is a protein called 50S ribosomal protein L3.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
30	21	203	Total	C	N	O	S	0	0	0
			1546	978	295	267	6			
30	29	204	Total	C	N	O	S	0	0	0
			1563	988	299	270	6			

- Molecule 31 is a protein called 50S ribosomal protein L4.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
31	31	202	Total	C	N	O	S	0	0	0
			1585	1011	297	275	2			
31	39	204	Total	C	N	O	S	0	0	0
			1602	1022	299	279	2			

- Molecule 32 is a protein called 50S ribosomal protein L5.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
32	41	179	Total	C	N	O	S	0	0	0
			1457	931	265	257	4			
32	49	180	Total	C	N	O	S	0	0	0
			1459	931	266	258	4			

- Molecule 33 is a protein called 50S ribosomal protein L6.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
33	51	174	Total	C	N	O	S	0	0	0
			1328	842	249	236	1			
33	59	169	Total	C	N	O	S	0	0	0
			1295	823	241	230	1			

- Molecule 34 is a protein called 50S ribosomal protein L9.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
34	61	145	Total	C	N	O	S	0	0	0
			1131	723	200	207	1			
34	69	145	Total	C	N	O	S	0	0	0
			1131	723	200	207	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
35	58	137	Total	C	N	O	S	0	0	0
			1096	706	205	181	4			
35	15	137	Total	C	N	O	S	0	0	0
			1096	707	205	181	3			

- 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	147	Total	C	N	O	S	0	0	0
			1122	698	229	192	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
			1117	712	211	187	7			
38	45	138	Total	C	N	O	S	0	0	0
			1099	702	208	183	6			

- 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			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
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	110	Total	C	N	O	0	0	0
			876	553	175	148			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
41	B8	135	Total	C	N	O	S	0	0	0
			1119	697	230	191	1			
41	75	133	Total	C	N	O	S	0	0	0
			1109	691	228	189	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
42	C8	115	Total	C	N	O	S	0	0	0
			950	603	199	147	1			
42	85	116	Total	C	N	O	S	0	0	0
			959	608	201	149	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	100	Total	C	N	O	S	0	0	0
			770	496	140	133	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
44	E8	110	Total	C	N	O	S	0	0	0
			876	552	171	151	2			
44	A5	111	Total	C	N	O	S	0	0	0
			886	558	174	152	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
			750	488	135	126	1			
45	B5	94	Total	C	N	O		0	0	0
			735	477	133	125				

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

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

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
47	H8	170	Total	C	N	O	S	0	0	0
			1365	870	246	246	3			
47	D5	177	Total	C	N	O	S	0	0	0
			1411	901	253	255	2			

- 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			
48	E5	76	Total	C	N	O	S	0	0	0
			603	372	128	102	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
			747	469	148	129	1			
49	F5	94	Total	C	N	O	S	0	0	0
			737	463	146	127	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
50	K8	68	Total	C	N	O	S	0	0	0
			575	358	116	100	1			
50	G5	69	Total	C	N	O	S	0	0	0
			576	358	116	101	1			

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

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
51	L8	58	Total	C	N	O	0	0	0
			459	293	89	77			
51	H5	58	Total	C	N	O	0	0	0
			459	293	89	77			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
52	M8	60	Total	C	N	O	S	0	0	0
			475	300	84	86	5			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
53	N8	48	Total	C	N	O	S	0	0	0
			369	229	75	60	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	45	Total	C	N	O	S	0	0	0
			389	241	79	65	4			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
55	P8	47	Total	C	N	O	S	0	0	0
			401	246	99	54	2			
55	L5	47	Total	C	N	O	S	0	0	0
			401	246	99	54	2			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
56	Q8	64	Total 516	C 331	N 102	O 81	S 2	0	0	0
56	M5	64	Total 516	C 331	N 102	O 81	S 2	0	0	0

- Molecule 57 is MAGNESIUM ION (three-letter code: MG) (formula: Mg).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	45	2	Total 2	Mg 2	0	0
57	P8	1	Total 1	Mg 1	0	0
57	32	1	Total 1	Mg 1	0	0
57	C5	1	Total 1	Mg 1	0	0
57	13	140	Total 140	Mg 140	0	0
57	1J	8	Total 8	Mg 8	0	0
57	5I	1	Total 1	Mg 1	0	0
57	35	1	Total 1	Mg 1	0	0
57	4L	1	Total 1	Mg 1	0	0
57	16	11	Total 11	Mg 11	0	0
57	42	1	Total 1	Mg 1	0	0
57	25	1	Total 1	Mg 1	0	0
57	M5	1	Total 1	Mg 1	0	0
57	21	3	Total 3	Mg 3	0	0
57	31	2	Total 2	Mg 2	0	0
57	Q8	1	Total 1	Mg 1	0	0
57	4I	1	Total 1	Mg 1	0	0
57	3I	1	Total 1	Mg 1	0	0

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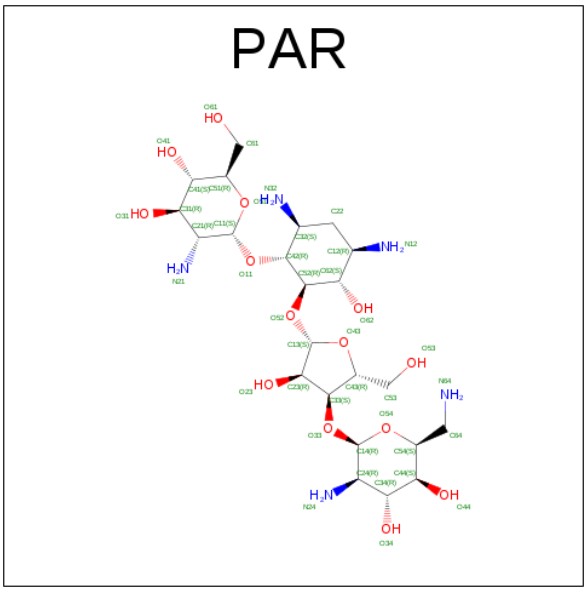
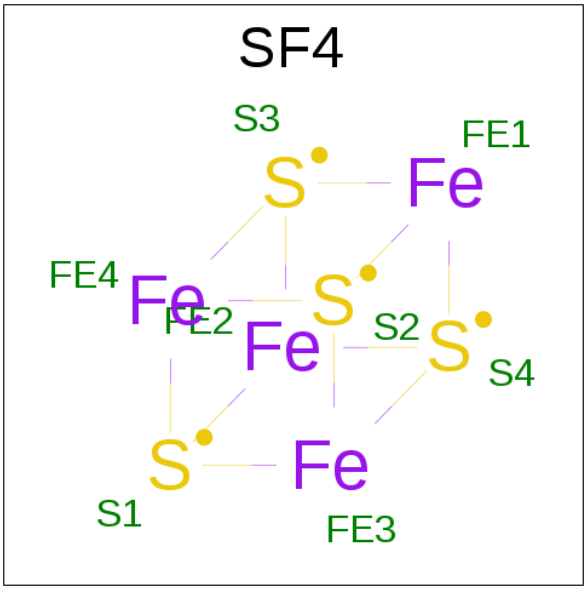
Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
57	I8	2	Total Mg 2 2	0	0
57	52	1	Total Mg 1 1	0	0
57	29	1	Total Mg 1 1	0	0
57	7A	1	Total Mg 1 1	0	0
57	2K	2	Total Mg 2 2	0	0
57	39	2	Total Mg 2 2	0	0
57	1G	102	Total Mg 102 102	0	0
57	1H	525	Total Mg 525 525	0	0
57	E5	1	Total Mg 1 1	0	0
57	88	3	Total Mg 3 3	0	0
57	14	435	Total Mg 435 435	0	0
57	78	1	Total Mg 1 1	0	0
57	19	1	Total Mg 1 1	0	0
57	41	1	Total Mg 1 1	0	0
57	2L	2	Total Mg 2 2	0	0

- Molecule 58 is PAROMOMYCIN (three-letter code: PAR) (formula: $C_{23}H_{45}N_5O_{14}$).



Mol	Chain	Residues	Atoms				ZeroOcc	AltConf
58	13	1	Total	C	N	O	0	0
			42	23	5	14		
58	1G	1	Total	C	N	O	0	0
			42	23	5	14		

- Molecule 59 is IRON/SULFUR CLUSTER (three-letter code: SF4) (formula: Fe_4S_4).

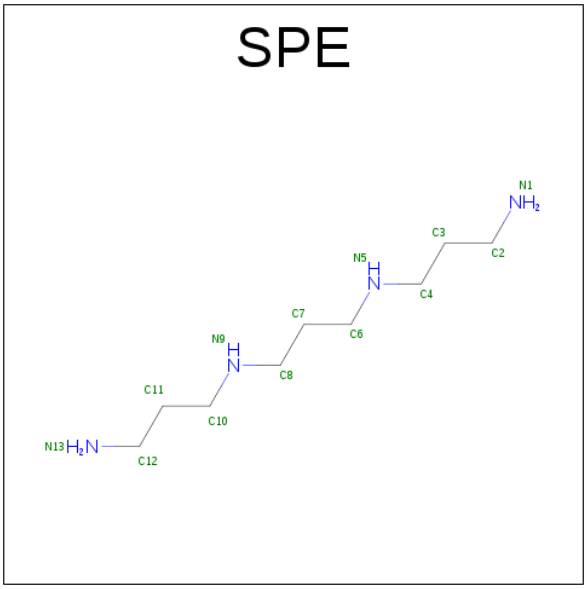


Mol	Chain	Residues	Atoms				ZeroOcc	AltConf
59	3E	1	Total	Fe	S		0	0
			8	4	4			
59	32	1	Total	Fe	S		0	0
			8	4	4			

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

Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
60	C5	1	Total Zn 1 1	0	0
60	5A	1	Total Zn 1 1	0	0
60	G8	1	Total Zn 1 1	0	0
60	5I	1	Total Zn 1 1	0	0

- Molecule 61 is THERMINE (three-letter code: SPE) (formula: $C_9H_{24}N_4$).



Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
61	1G	1	Total C N 13 9 4	0	0
61	14	1	Total C N 13 9 4	0	0
61	14	1	Total C N 13 9 4	0	0

- Molecule 62 is water.

Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
62	13	320	Total O 320 320	0	0
62	4E	3	Total O 3 3	0	0

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Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
62	8E	2	Total O 2 2	0	0
62	1I	2	Total O 2 2	0	0
62	3I	2	Total O 2 2	0	0
62	5I	2	Total O 2 2	0	0
62	6I	3	Total O 3 3	0	0
62	7I	1	Total O 1 1	0	0
62	BI	2	Total O 2 2	0	0
62	1F	1	Total O 1 1	0	0
62	1K	2	Total O 2 2	0	0
62	2K	8	Total O 8 8	0	0
62	4K	5	Total O 5 5	0	0
62	1H	1470	Total O 1470 1470	0	0
62	16	12	Total O 12 12	0	0
62	11	17	Total O 17 17	0	0
62	21	7	Total O 7 7	0	0
62	31	5	Total O 5 5	0	0
62	41	1	Total O 1 1	0	0
62	58	2	Total O 2 2	0	0
62	78	10	Total O 10 10	0	0
62	B8	1	Total O 1 1	0	0
62	C8	3	Total O 3 3	0	0

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Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
62	F8	2	Total O 2 2	0	0
62	G8	2	Total O 2 2	0	0
62	I8	6	Total O 6 6	0	0
62	J8	4	Total O 4 4	0	0
62	L8	3	Total O 3 3	0	0
62	P8	1	Total O 1 1	0	0
62	Q8	5	Total O 5 5	0	0
62	1G	317	Total O 317 317	0	0
62	32	1	Total O 1 1	0	0
62	52	4	Total O 4 4	0	0
62	1A	1	Total O 1 1	0	0
62	6A	2	Total O 2 2	0	0
62	7A	5	Total O 5 5	0	0
62	9A	2	Total O 2 2	0	0
62	BA	2	Total O 2 2	0	0
62	2L	6	Total O 6 6	0	0
62	4L	6	Total O 6 6	0	0
62	14	1144	Total O 1144 1144	0	0
62	1J	12	Total O 12 12	0	0
62	19	15	Total O 15 15	0	0
62	29	4	Total O 4 4	0	0

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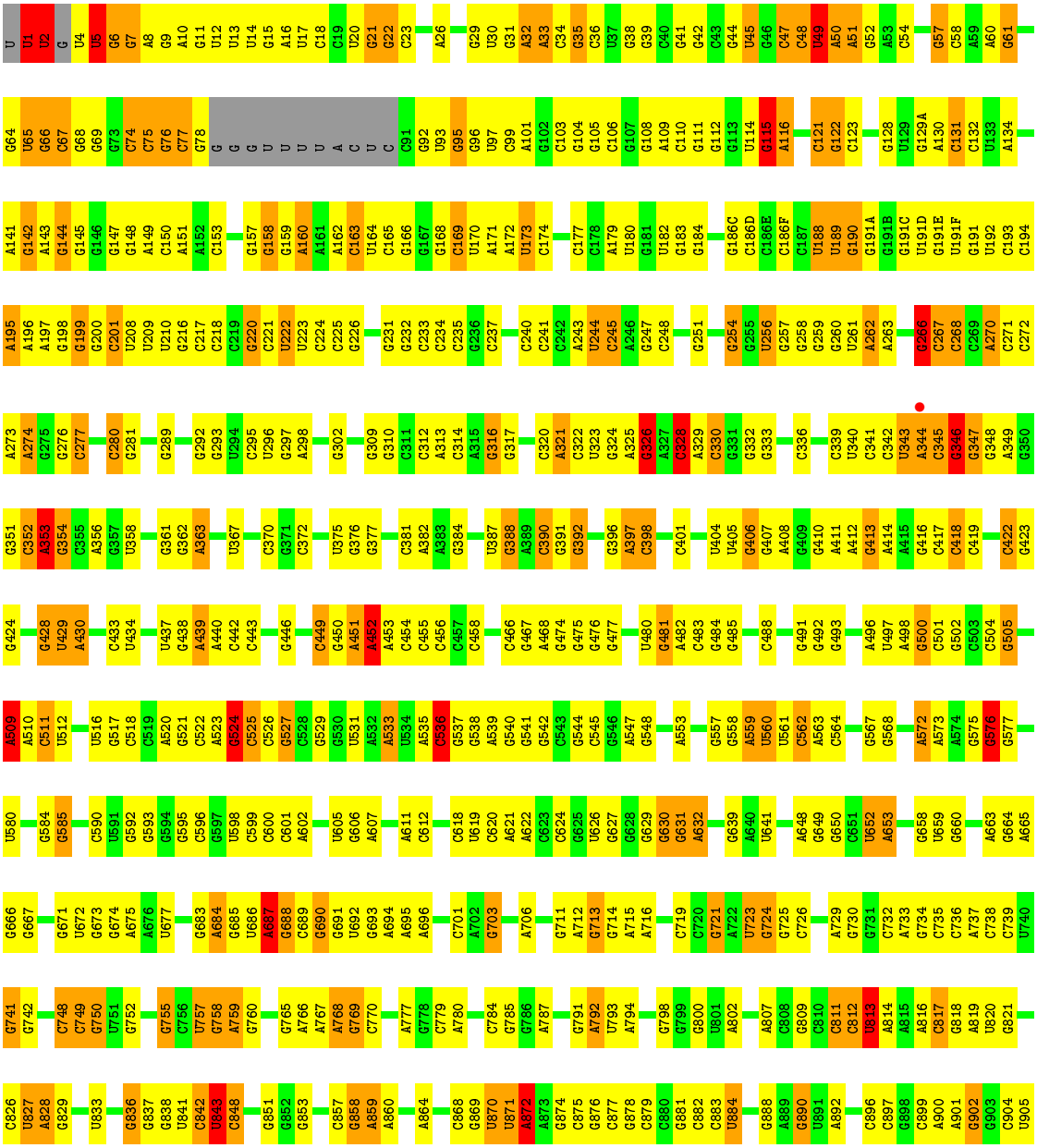
Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
62	39	5	Total O 5 5	0	0
62	15	1	Total O 1 1	0	0
62	25	6	Total O 6 6	0	0
62	35	8	Total O 8 8	0	0
62	55	3	Total O 3 3	0	0
62	85	2	Total O 2 2	0	0
62	A5	1	Total O 1 1	0	0
62	B5	1	Total O 1 1	0	0
62	C5	3	Total O 3 3	0	0
62	F5	1	Total O 1 1	0	0
62	H5	2	Total O 2 2	0	0
62	M5	9	Total O 9 9	0	0

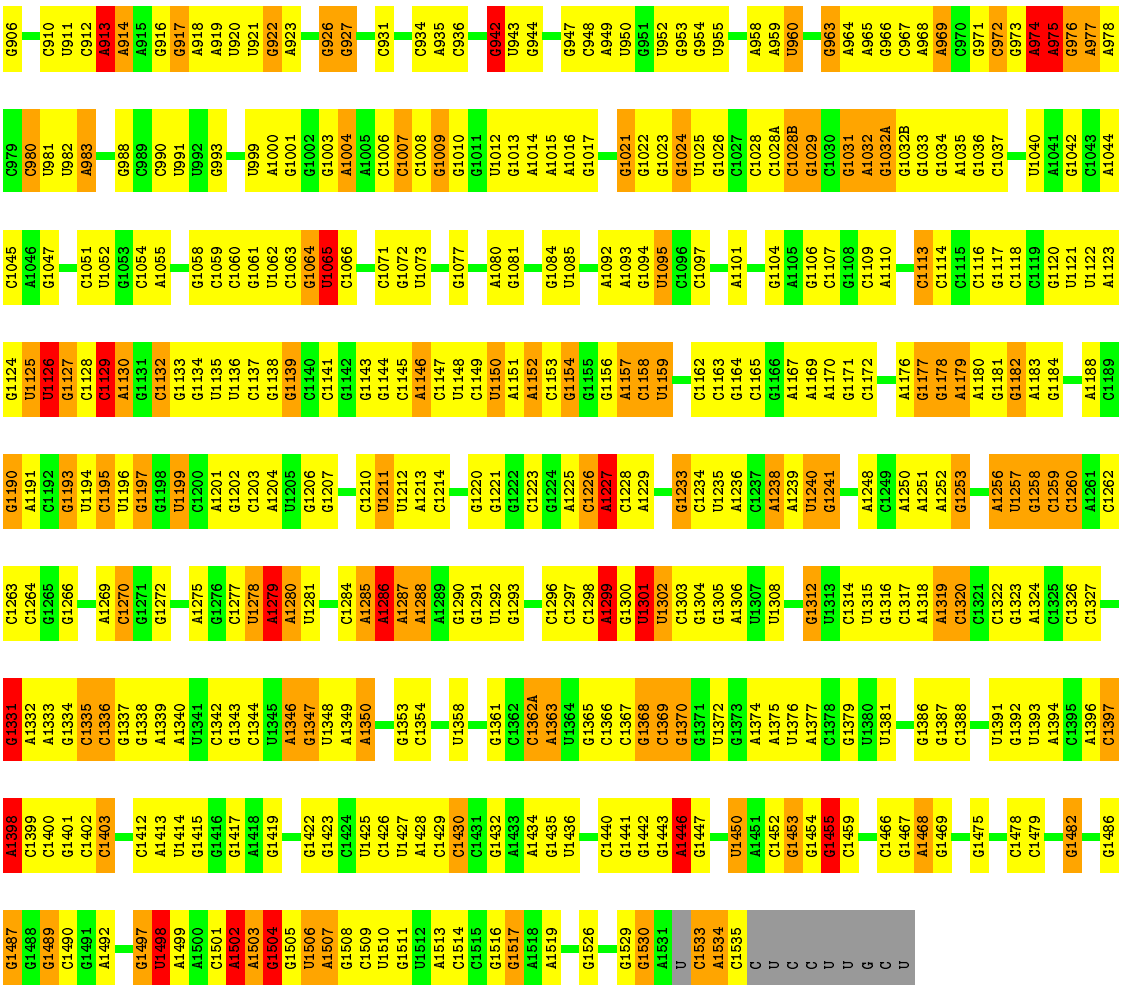
3 Residue-property plots (i)

These plots are drawn for all protein, RNA and DNA chains in the entry. The first graphic for a chain summarises the proportions of errors 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

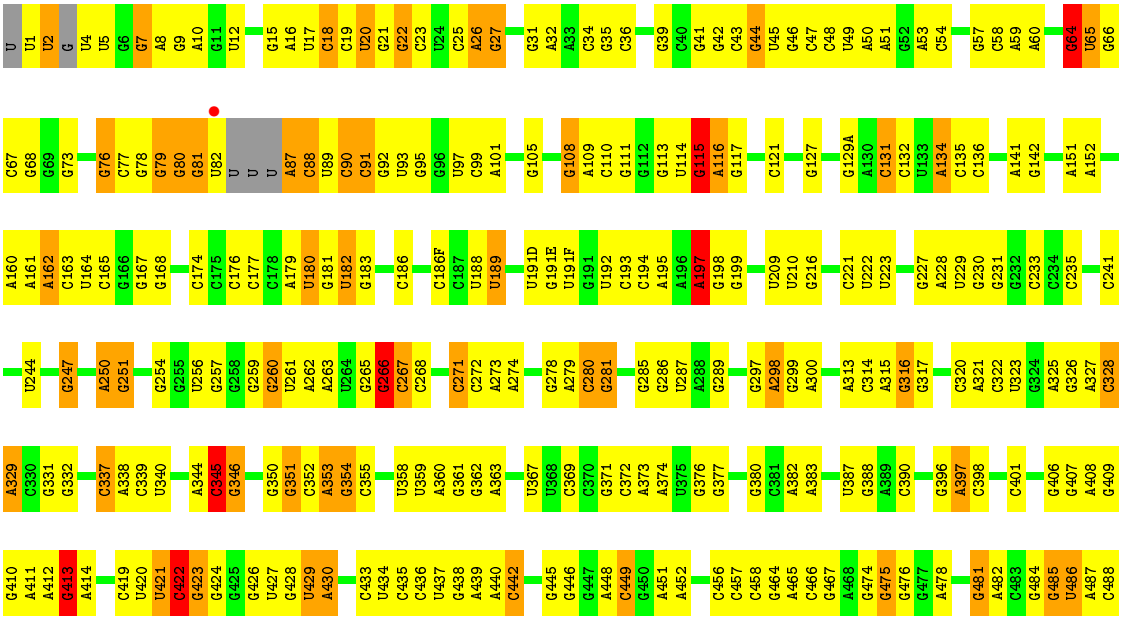
Chain 13: 37% 45% 15%

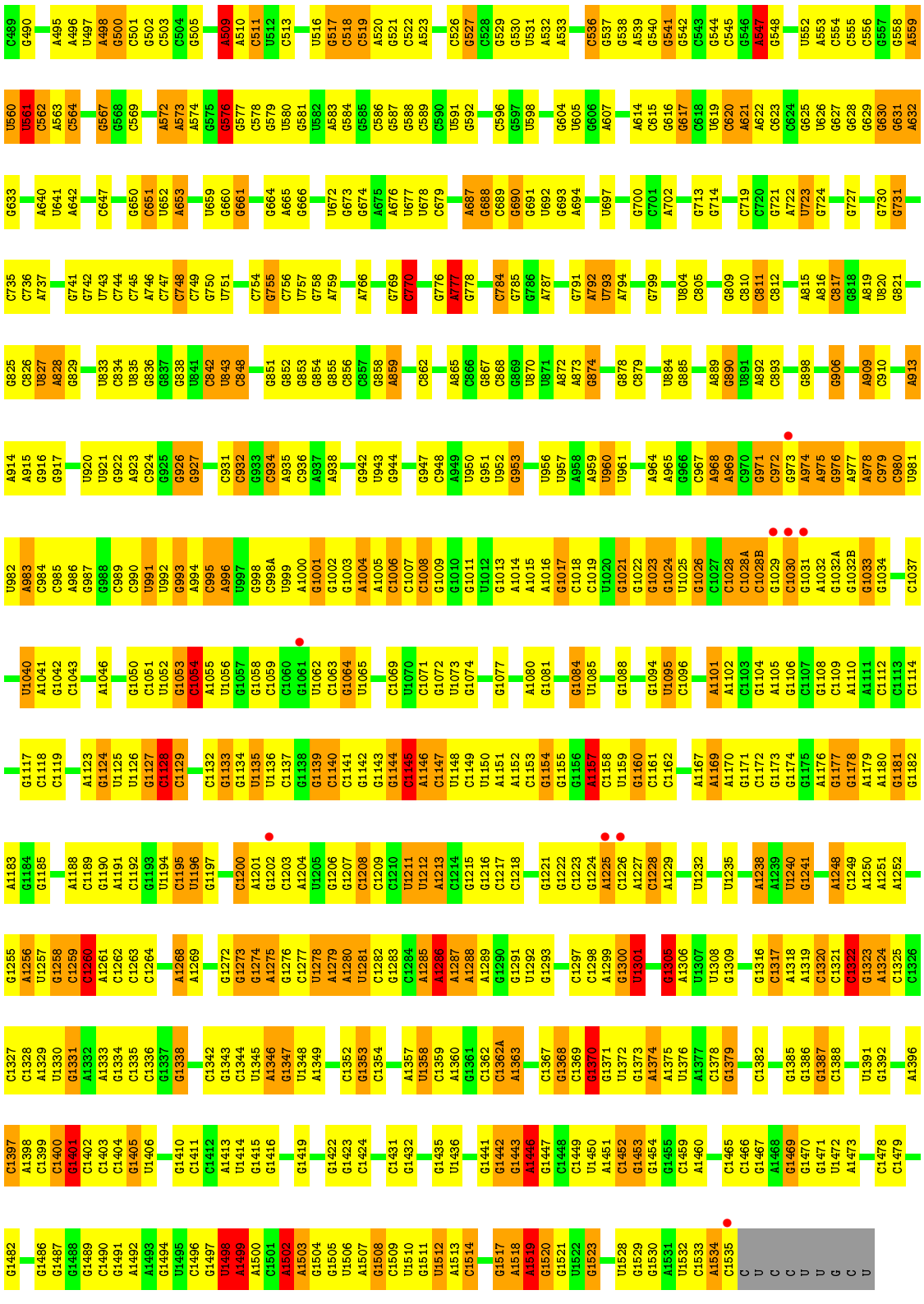




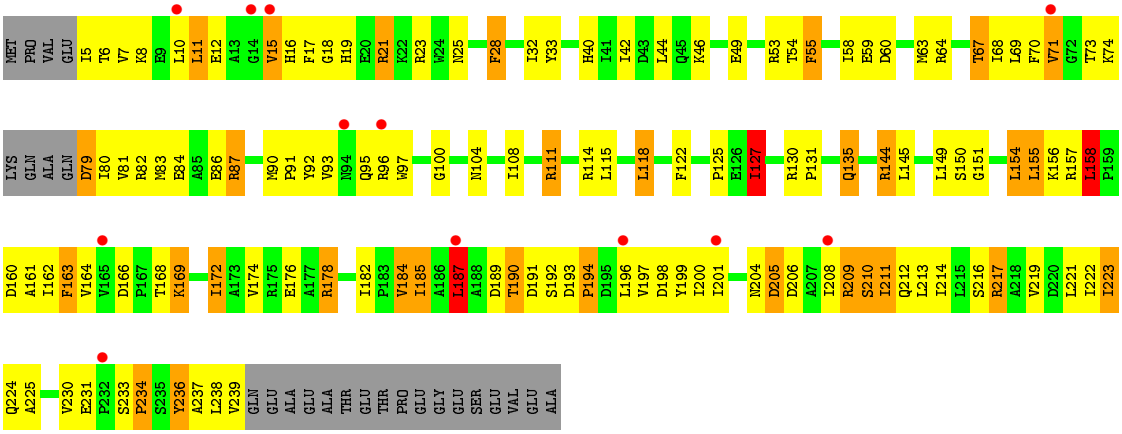
• Molecule 1: 16S ribosomal RNA

Chain 1G:



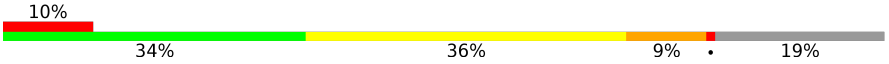


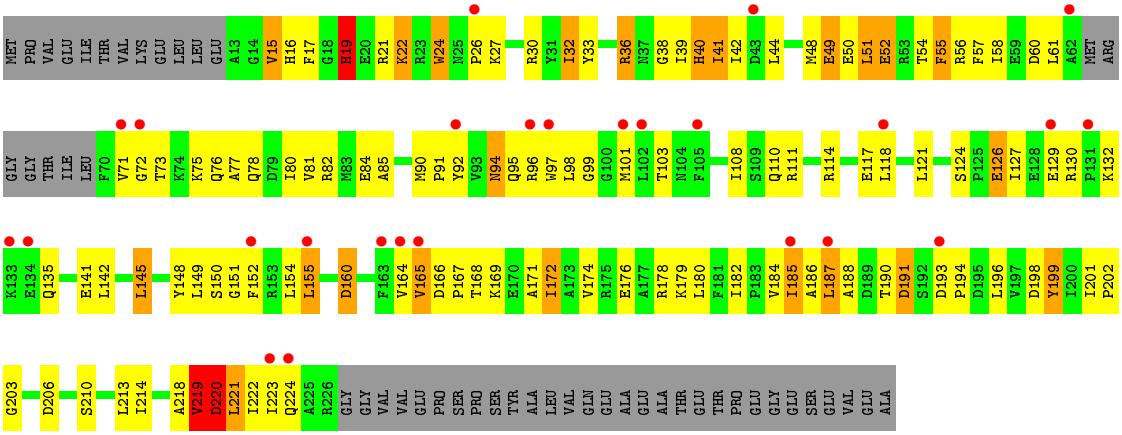
• Molecule 2: 30S ribosomal protein S2



• Molecule 2: 30S ribosomal protein S2

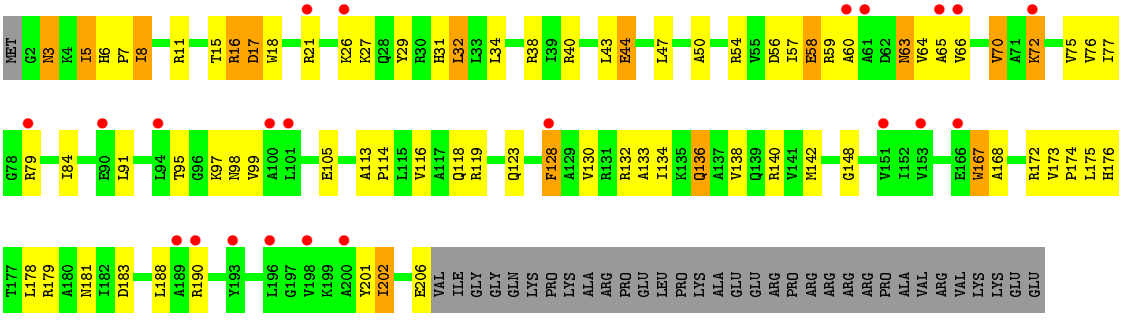
Chain 12:





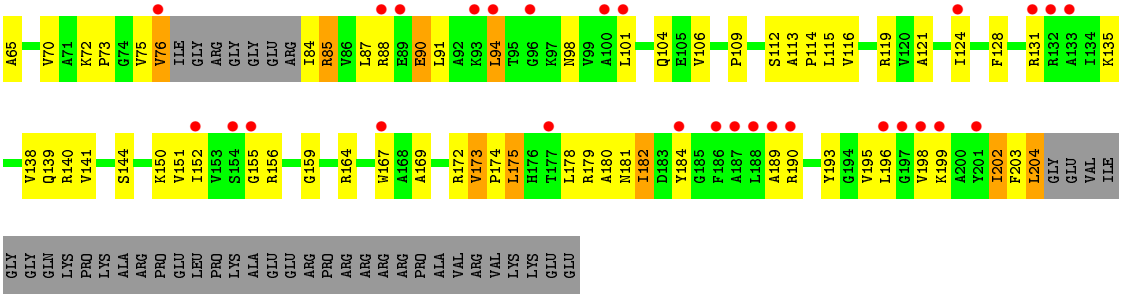
• Molecule 3: 30S ribosomal protein S3

Chain 2E:



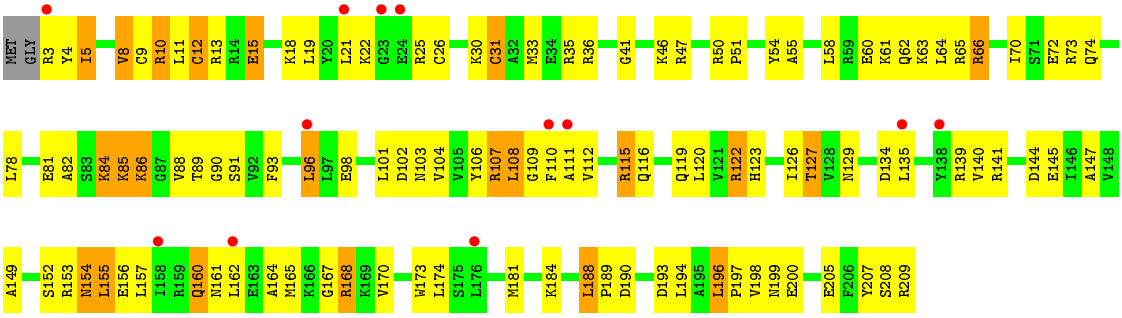
• Molecule 3: 30S ribosomal protein S3

Chain 22:



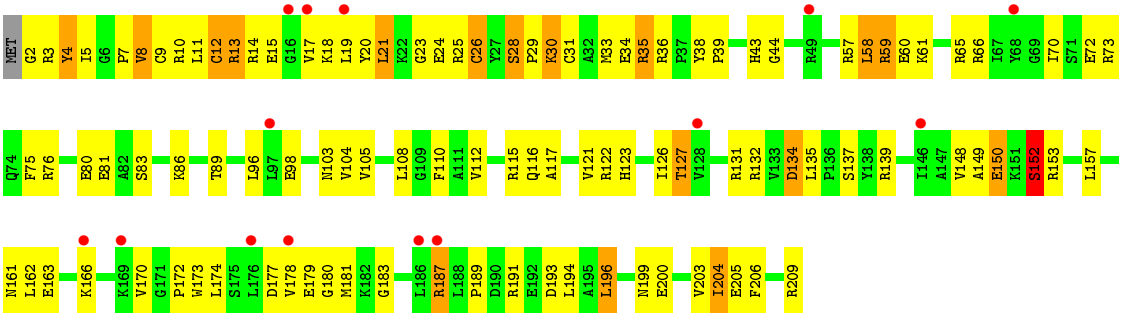
• Molecule 4: 30S ribosomal protein S4

Chain 3E:



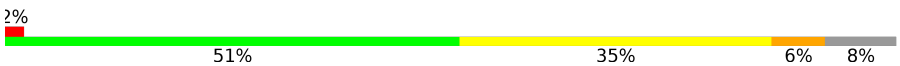
• Molecule 4: 30S ribosomal protein S4

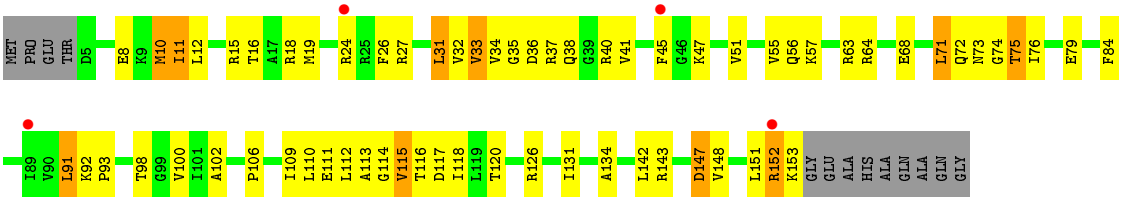
Chain 32:



• Molecule 5: 30S ribosomal protein S5

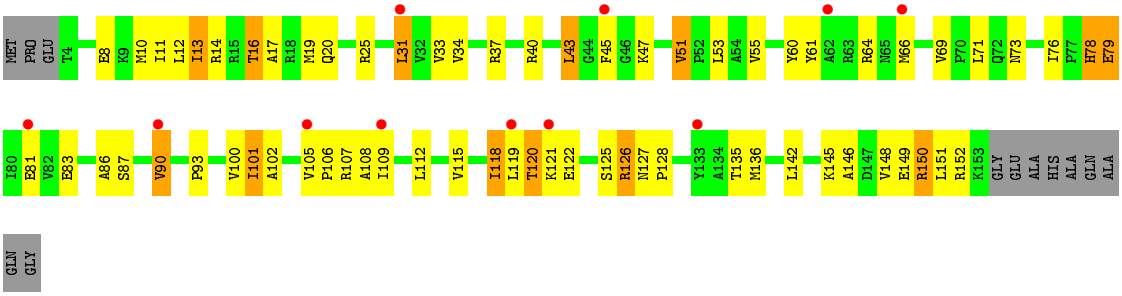
Chain 4E:



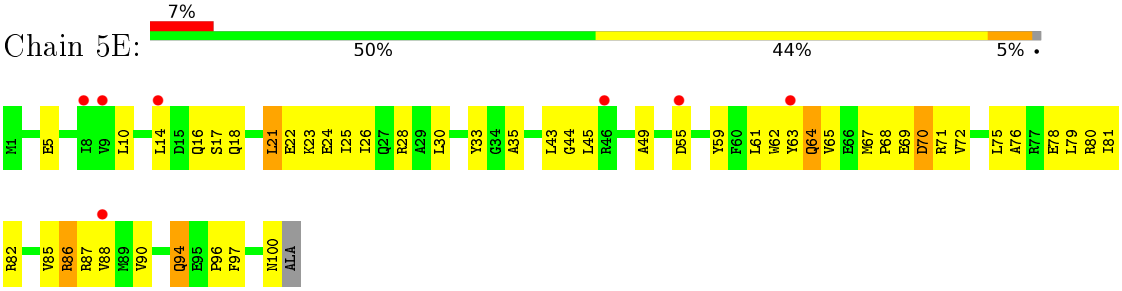


• Molecule 5: 30S ribosomal protein S5

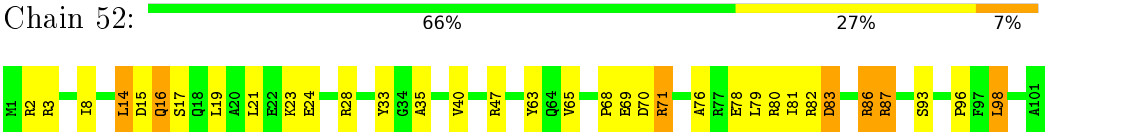
Chain 42:



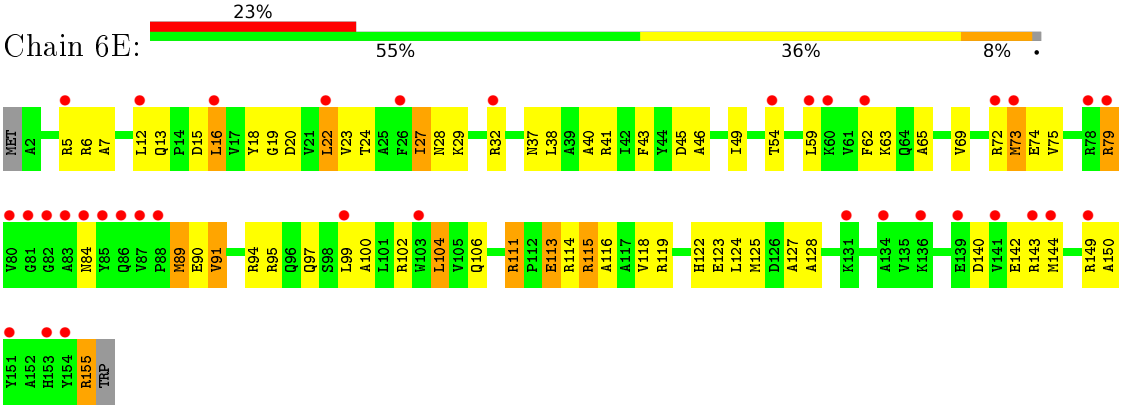
- Molecule 6: 30S ribosomal protein S6



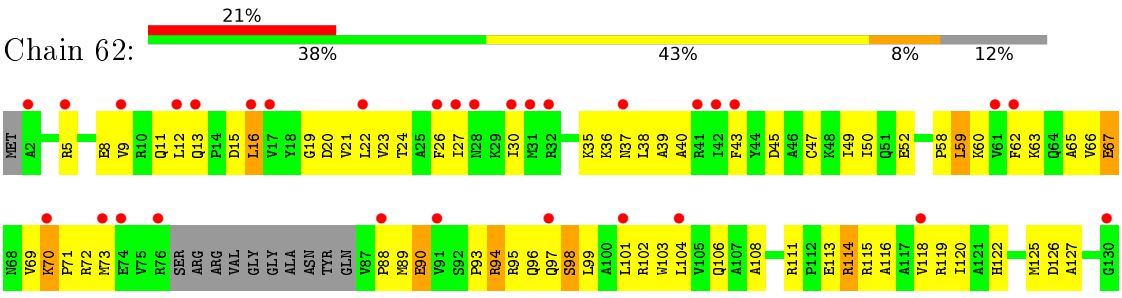
- Molecule 6: 30S ribosomal protein S6

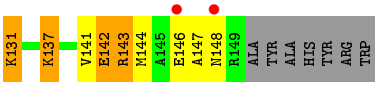


- Molecule 7: 30S ribosomal protein S7

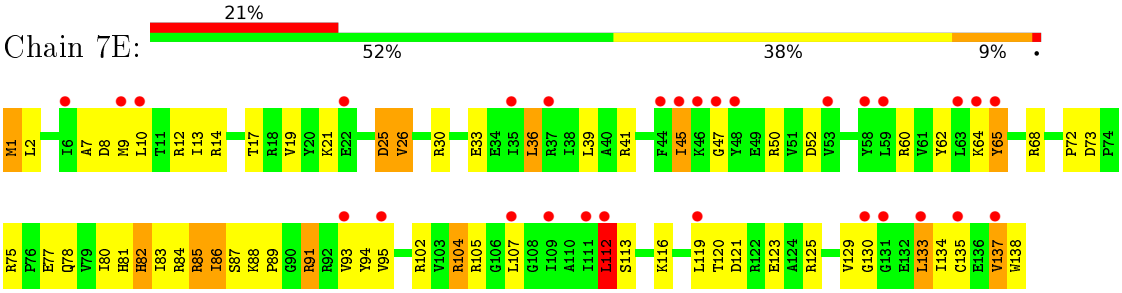


- Molecule 7: 30S ribosomal protein S7

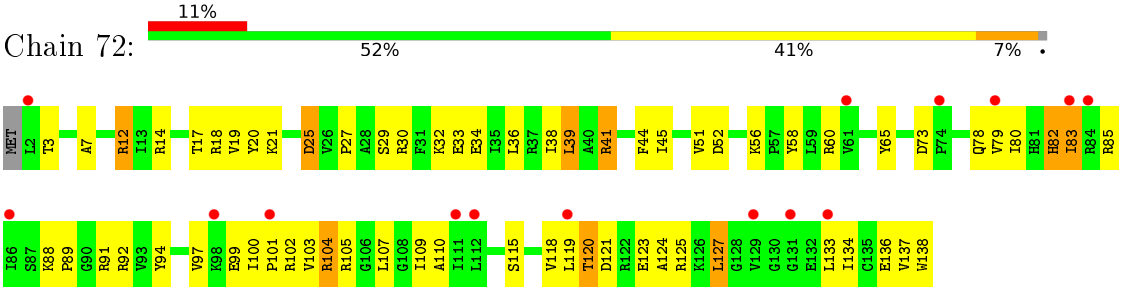




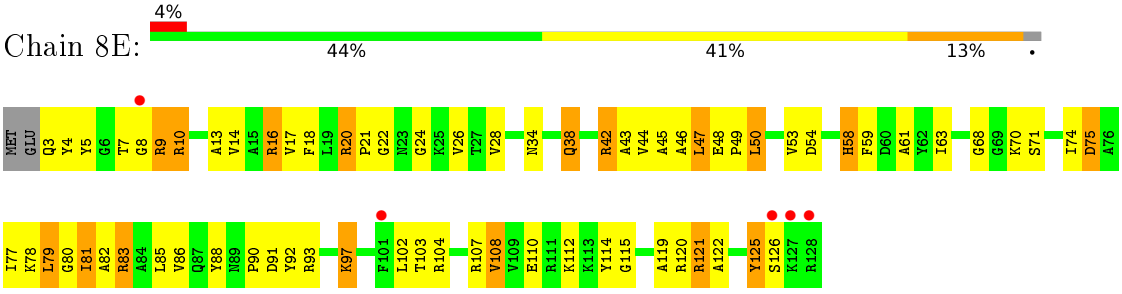
• Molecule 8: 30S ribosomal protein S8



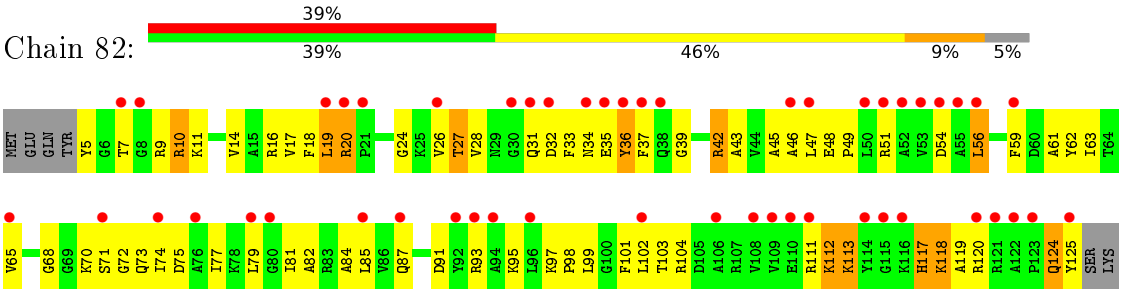
• Molecule 8: 30S ribosomal protein S8



• Molecule 9: 30S ribosomal protein S9

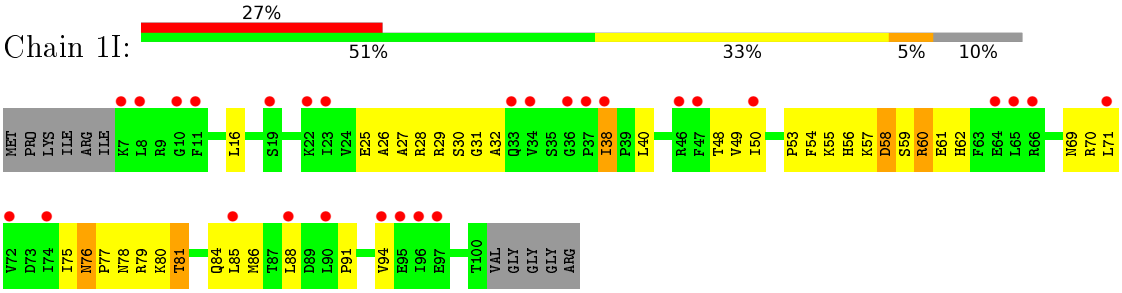


• Molecule 9: 30S ribosomal protein S9



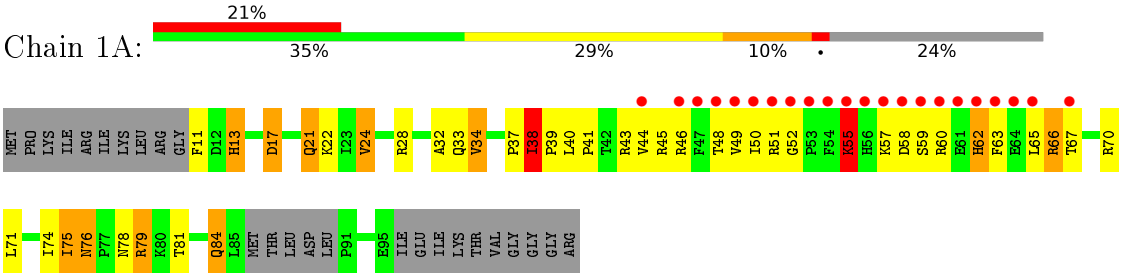
• Molecule 10: 30S ribosomal protein S10

Chain 1I:



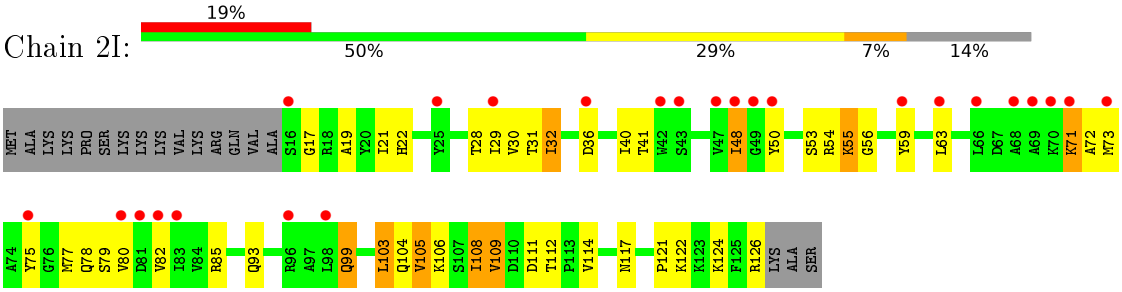
• Molecule 10: 30S ribosomal protein S10

Chain 1A:



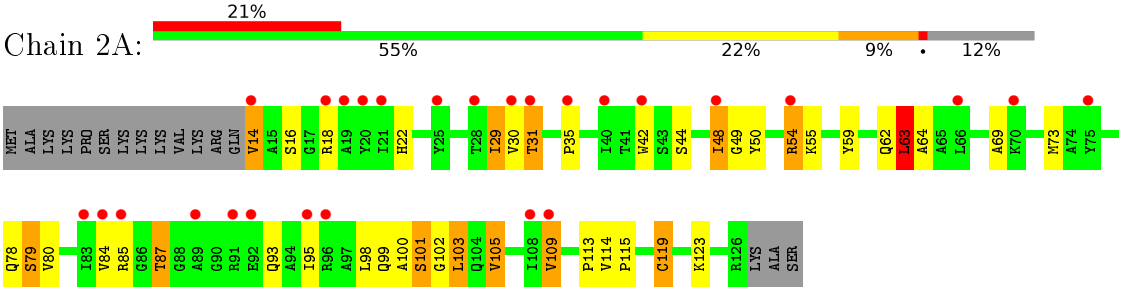
• Molecule 11: 30S ribosomal protein S11

Chain 2I:



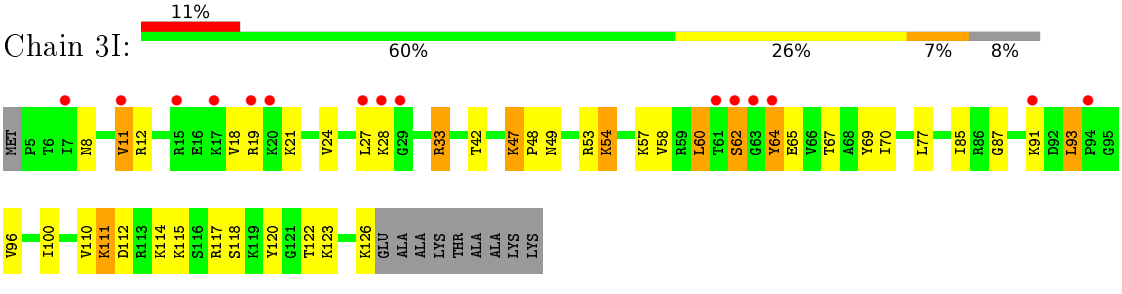
• Molecule 11: 30S ribosomal protein S11

Chain 2A:



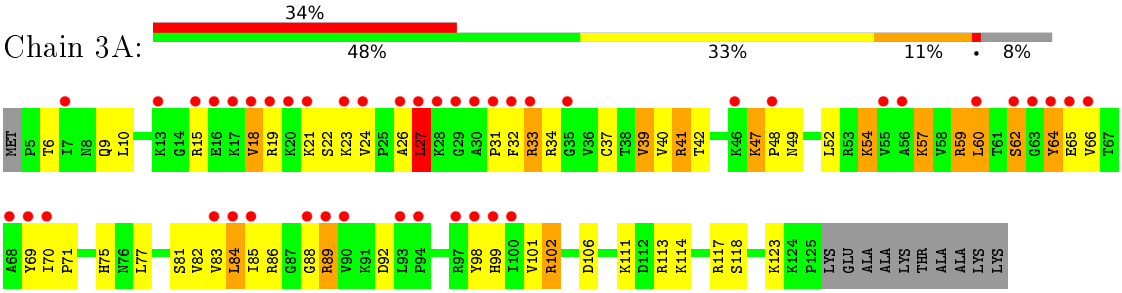
• Molecule 12: 30S ribosomal protein S12

Chain 3I:



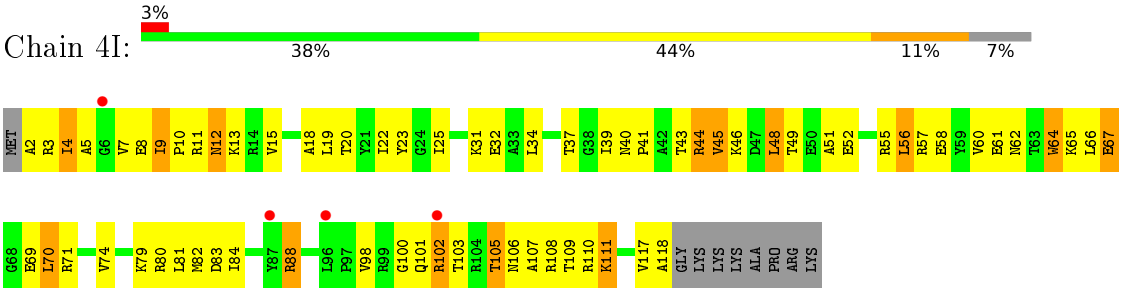
• Molecule 12: 30S ribosomal protein S12

Chain 3A:



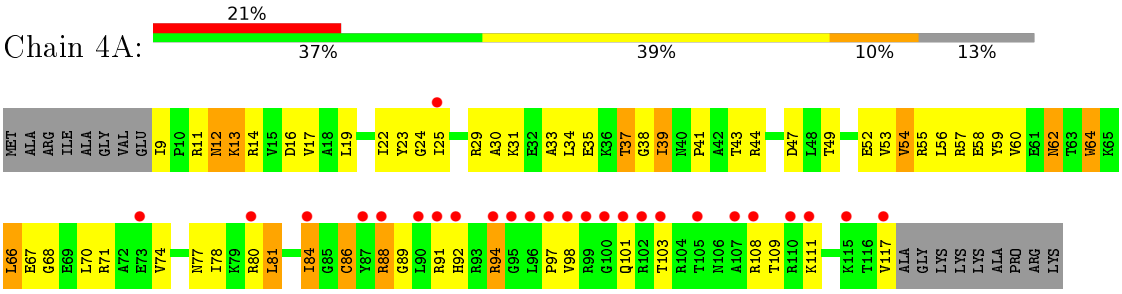
• Molecule 13: 30S ribosomal protein S13

Chain 4I:



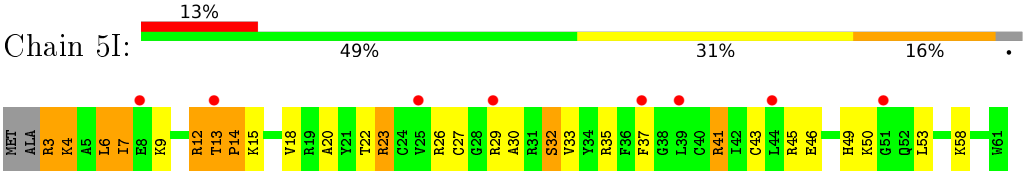
• Molecule 13: 30S ribosomal protein S13

Chain 4A:



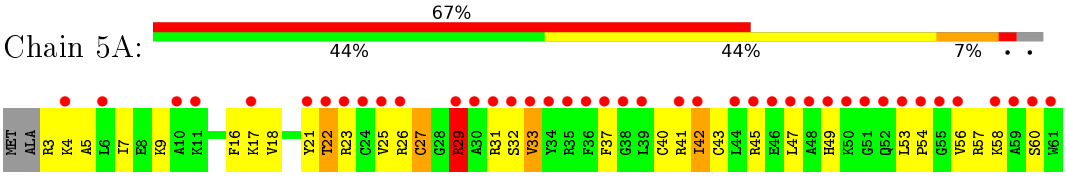
• Molecule 14: 30S ribosomal protein S14 type Z

Chain 5I:



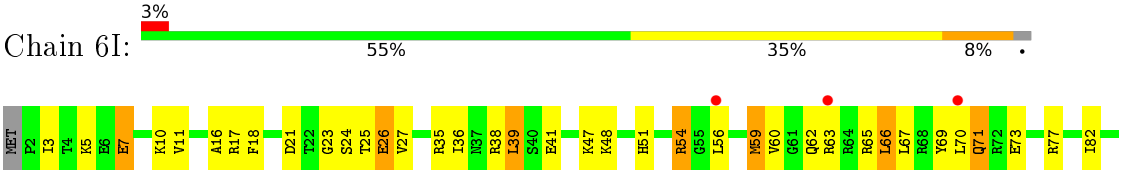
• Molecule 14: 30S ribosomal protein S14 type Z

Chain 5A:



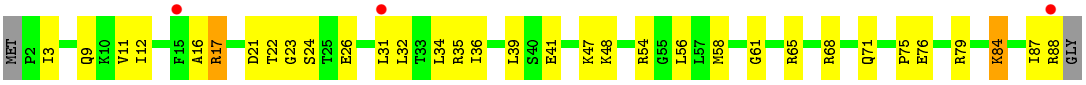
• Molecule 15: 30S ribosomal protein S15

Chain 6I:



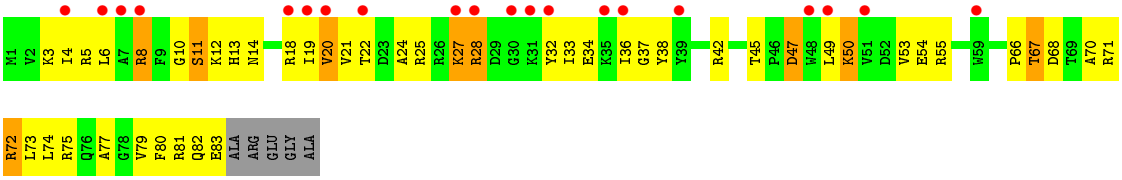
• Molecule 15: 30S ribosomal protein S15

Chain 6A:



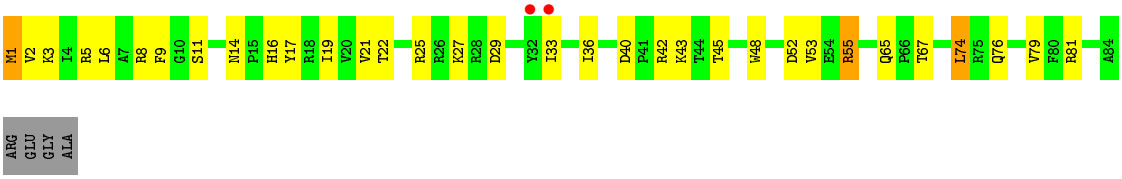
• Molecule 16: 30S ribosomal protein S16

Chain 7I:



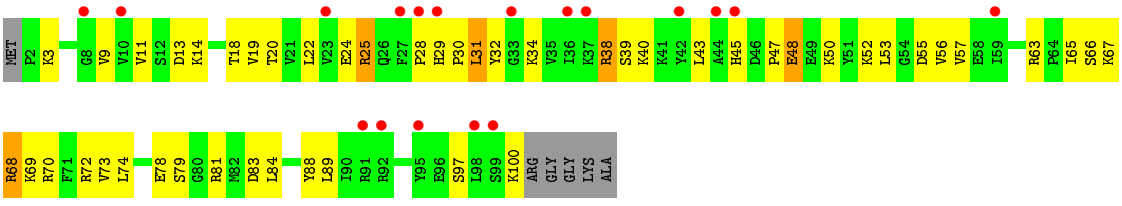
• Molecule 16: 30S ribosomal protein S16

Chain 7A:



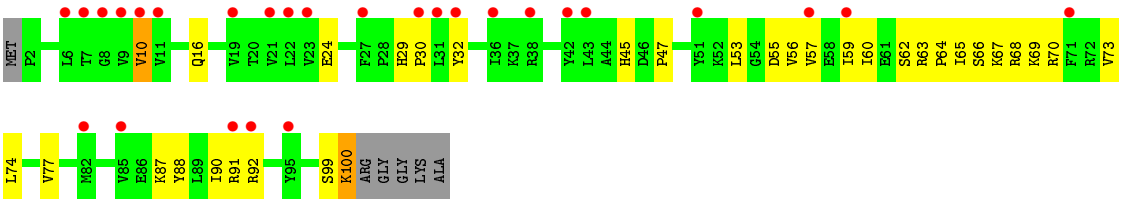
• Molecule 17: 30S ribosomal protein S17

Chain 8I:



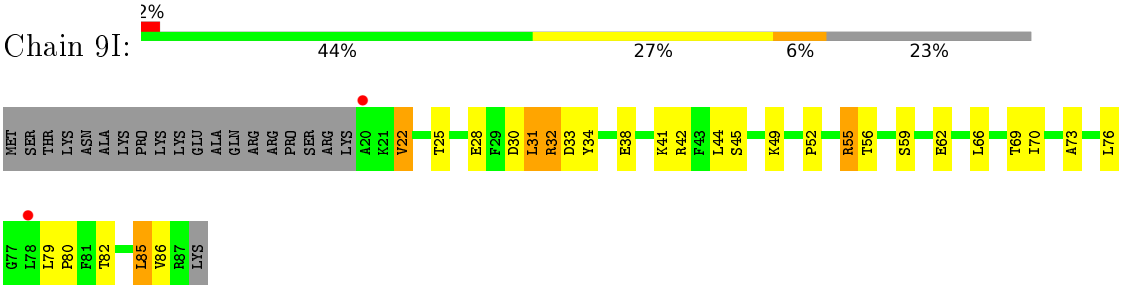
• Molecule 17: 30S ribosomal protein S17

Chain 8A:



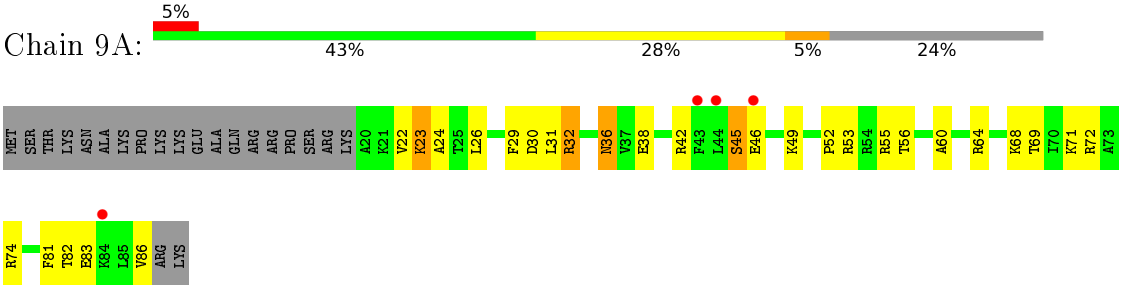
• Molecule 18: 30S ribosomal protein S18

Chain 9I:



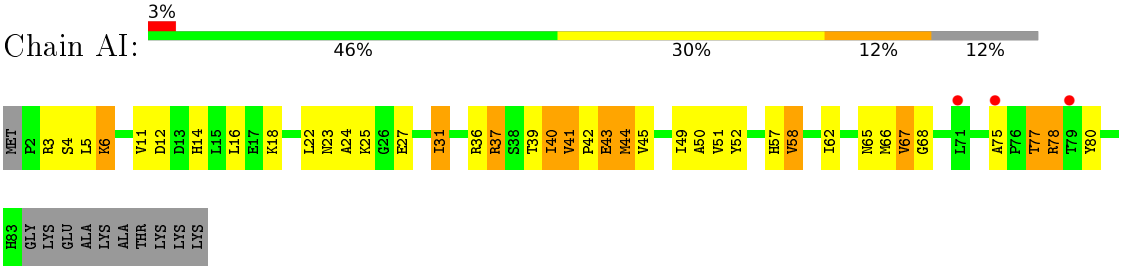
• Molecule 18: 30S ribosomal protein S18

Chain 9A:



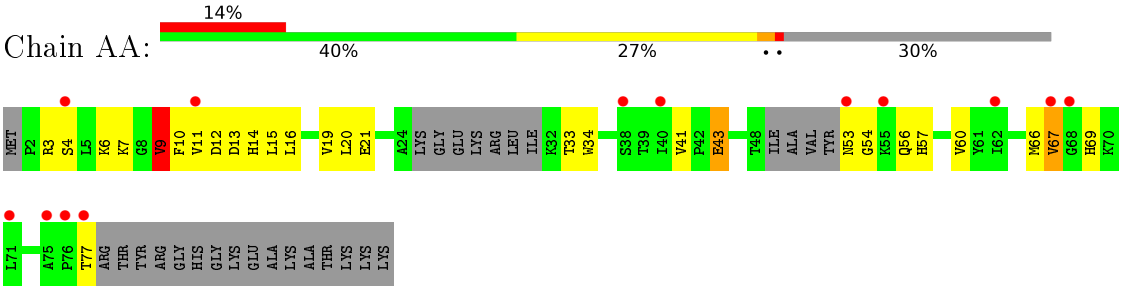
• Molecule 19: 30S ribosomal protein S19

Chain AI:



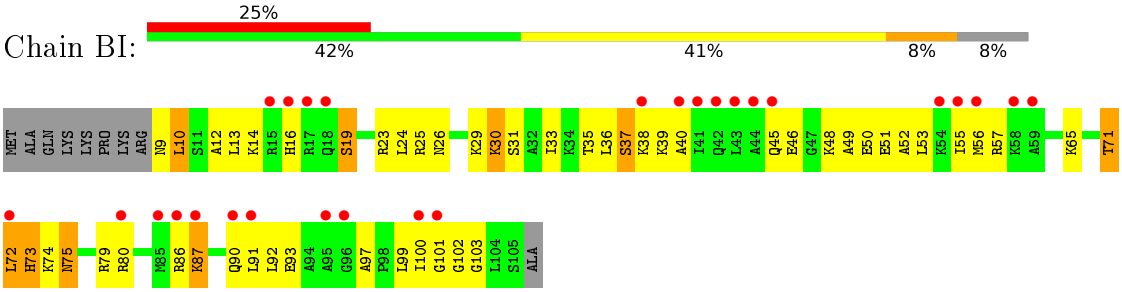
• Molecule 19: 30S ribosomal protein S19

Chain AA:



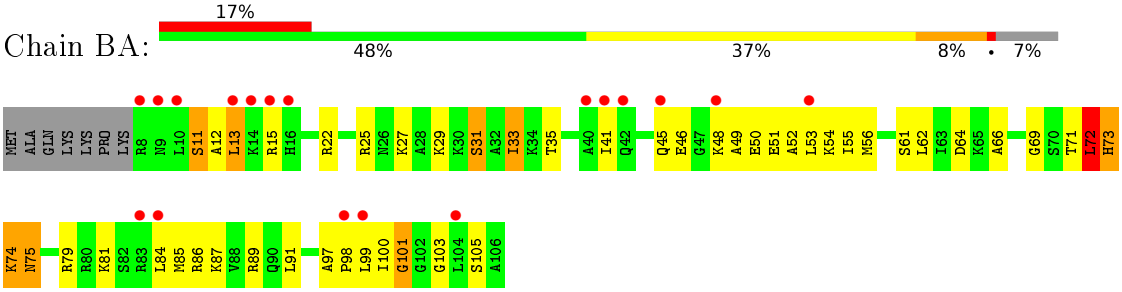
• Molecule 20: 30S ribosomal protein S20

Chain BI:



• Molecule 20: 30S ribosomal protein S20

Chain BA:

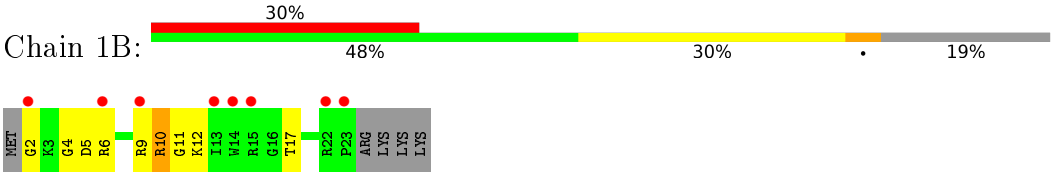


• Molecule 21: 30S ribosomal protein Thx

Chain 1F:

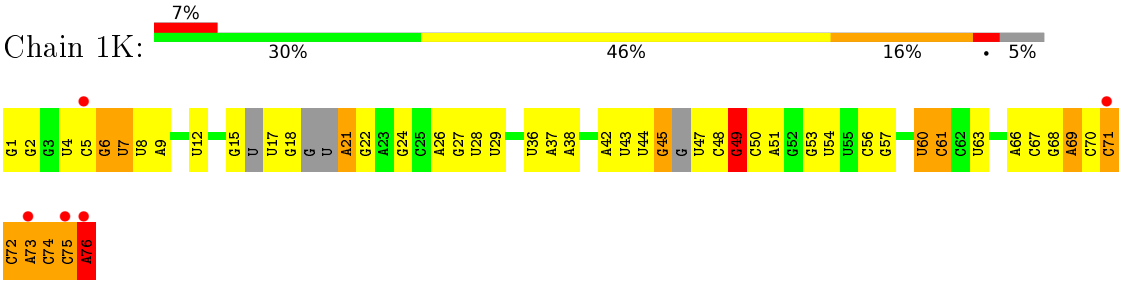
• Molecule 21: 30S ribosomal protein Thx

Chain 1B:



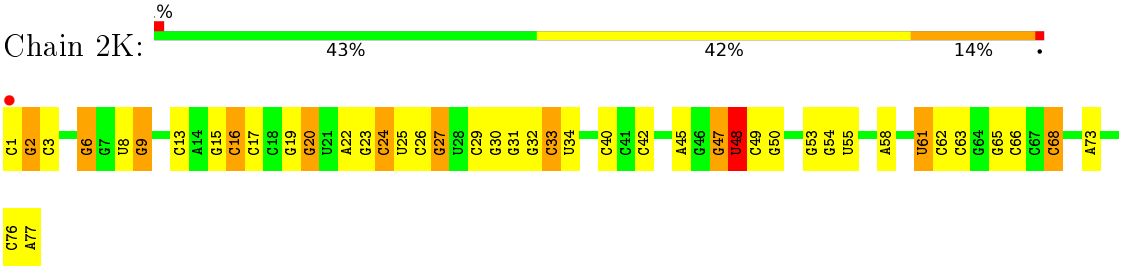
• Molecule 22: tRNALys

Chain 1K:



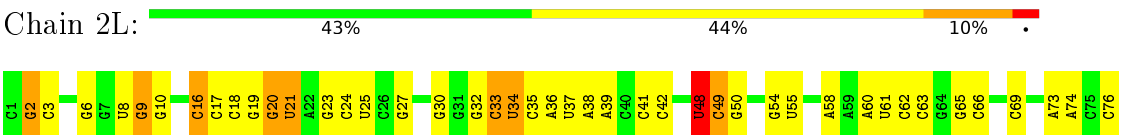
• Molecule 23: tRNAfMet

Chain 2K:



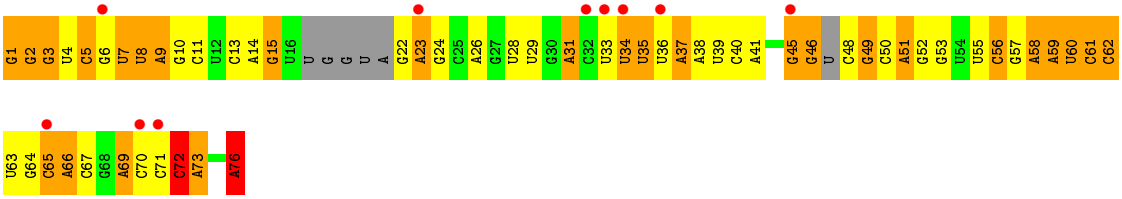
• Molecule 23: tRNAfMet

Chain 2L:



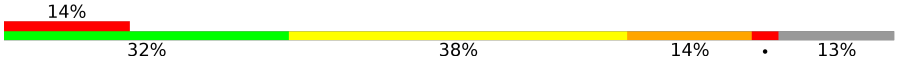
- Molecule 24: tRNALys

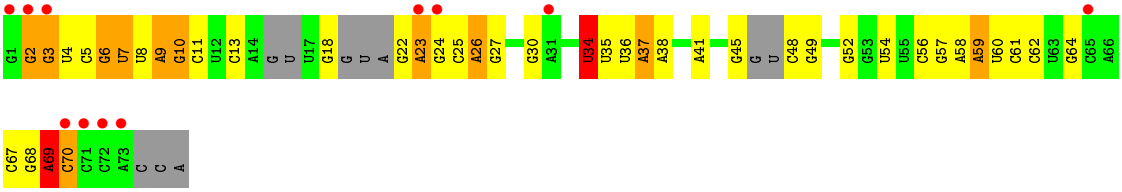
Chain 3K:



- Molecule 24: tRNALys

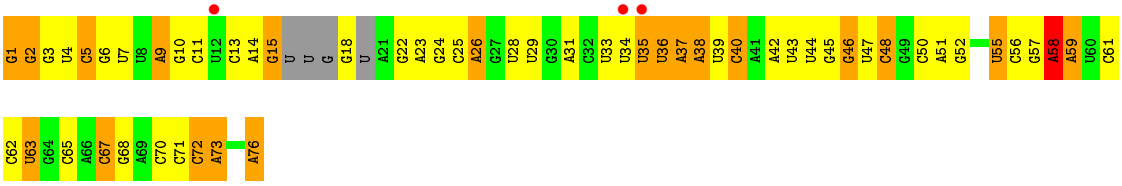
Chain 1L:





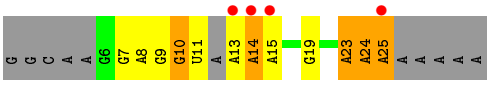
- Molecule 24: tRNALys

Chain 3L:



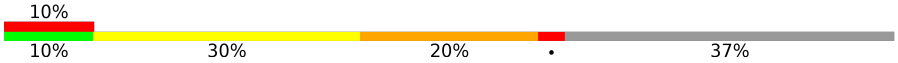
- Molecule 25: mRNA

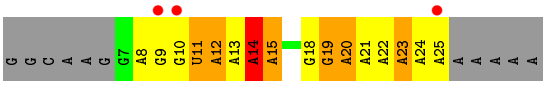
Chain 4K:



- Molecule 25: mRNA

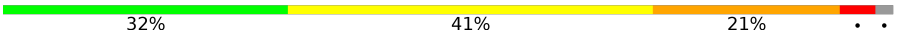
Chain 4L:

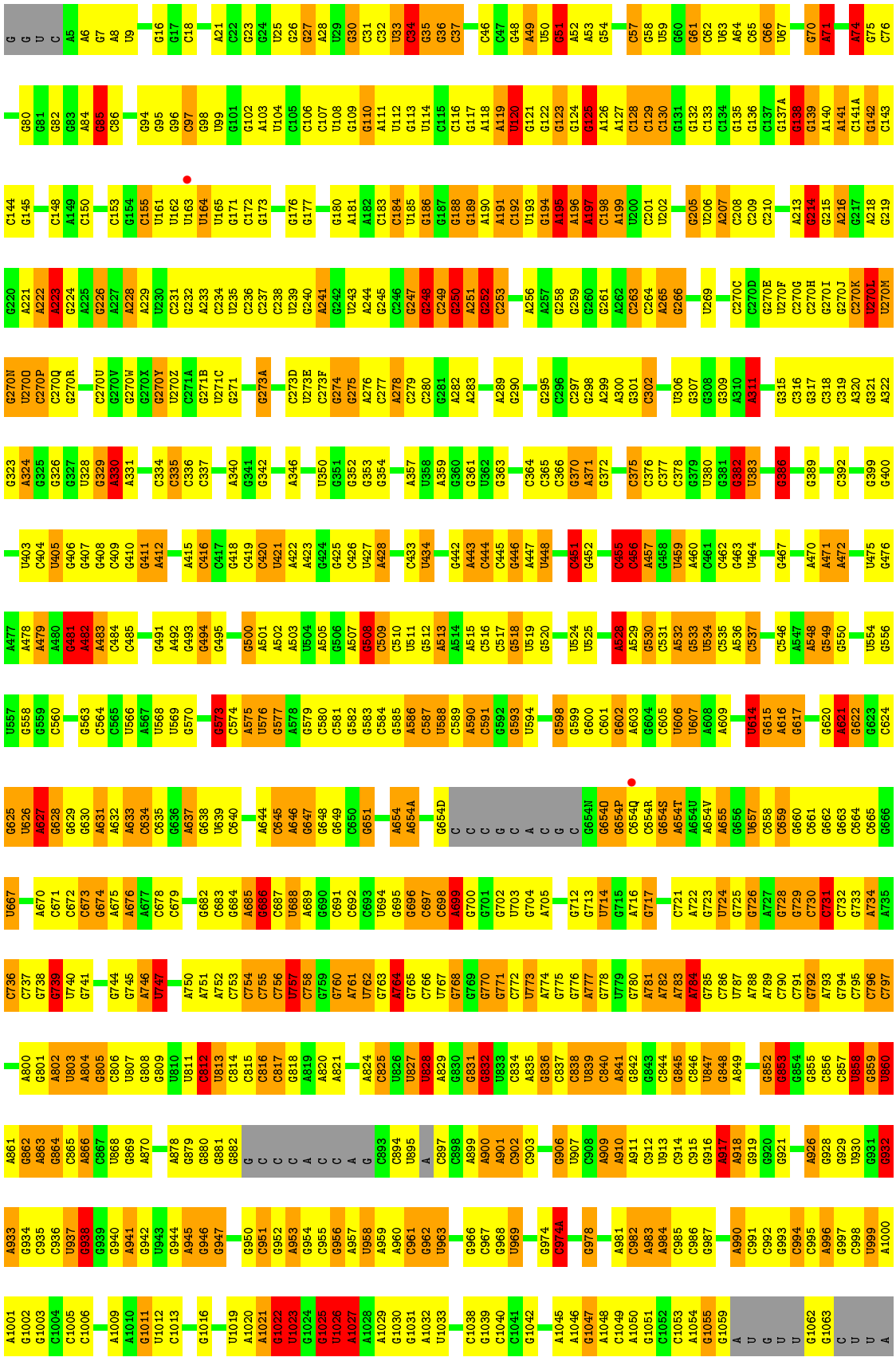


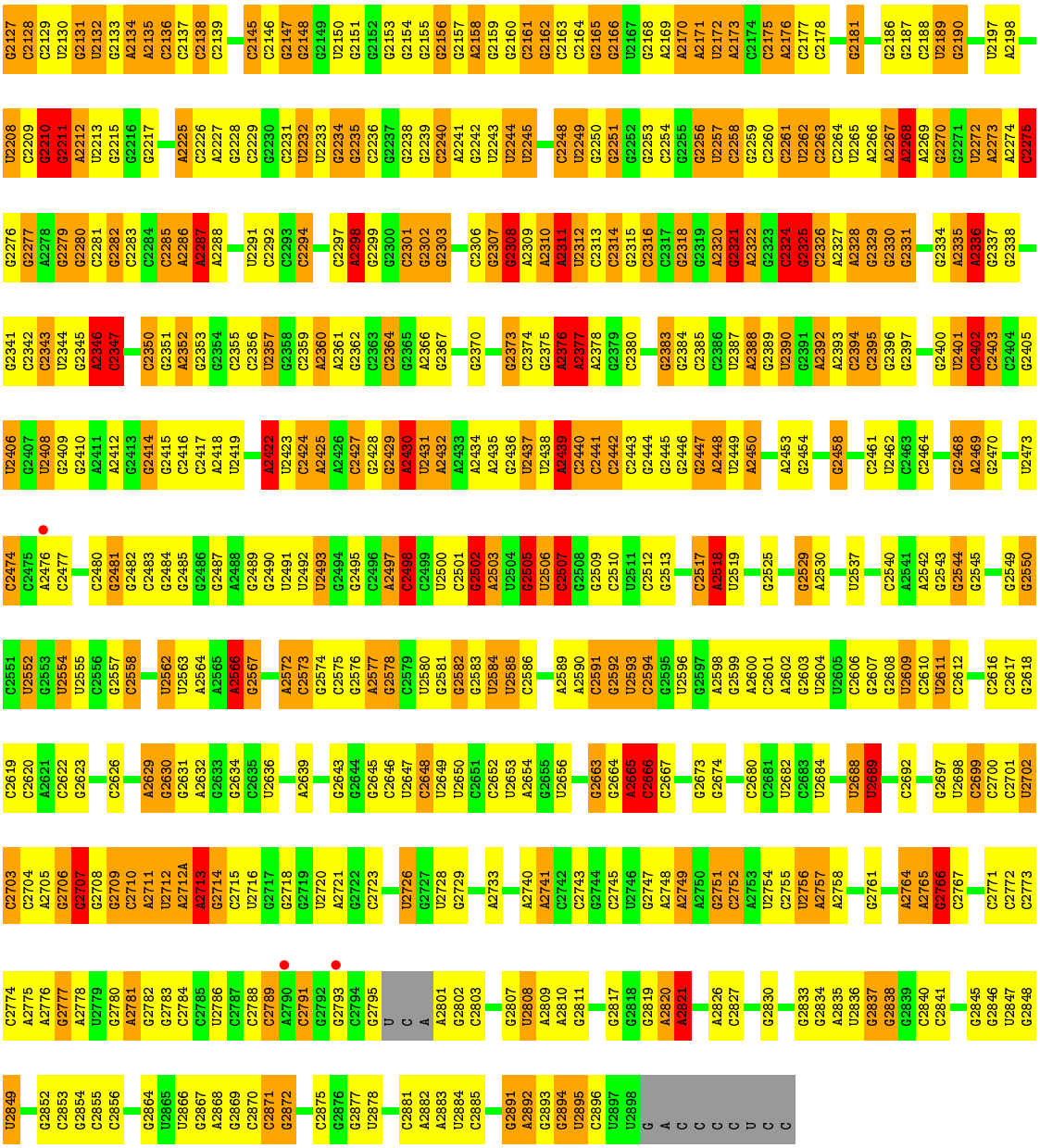


- Molecule 26: 23S ribosomal RNA

Chain 1H:

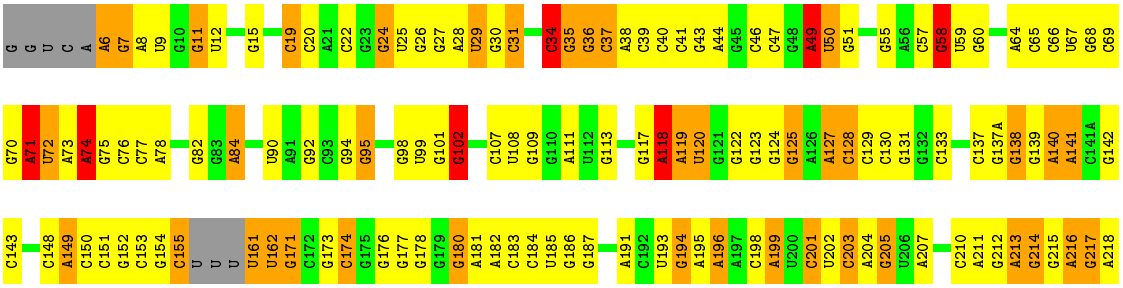


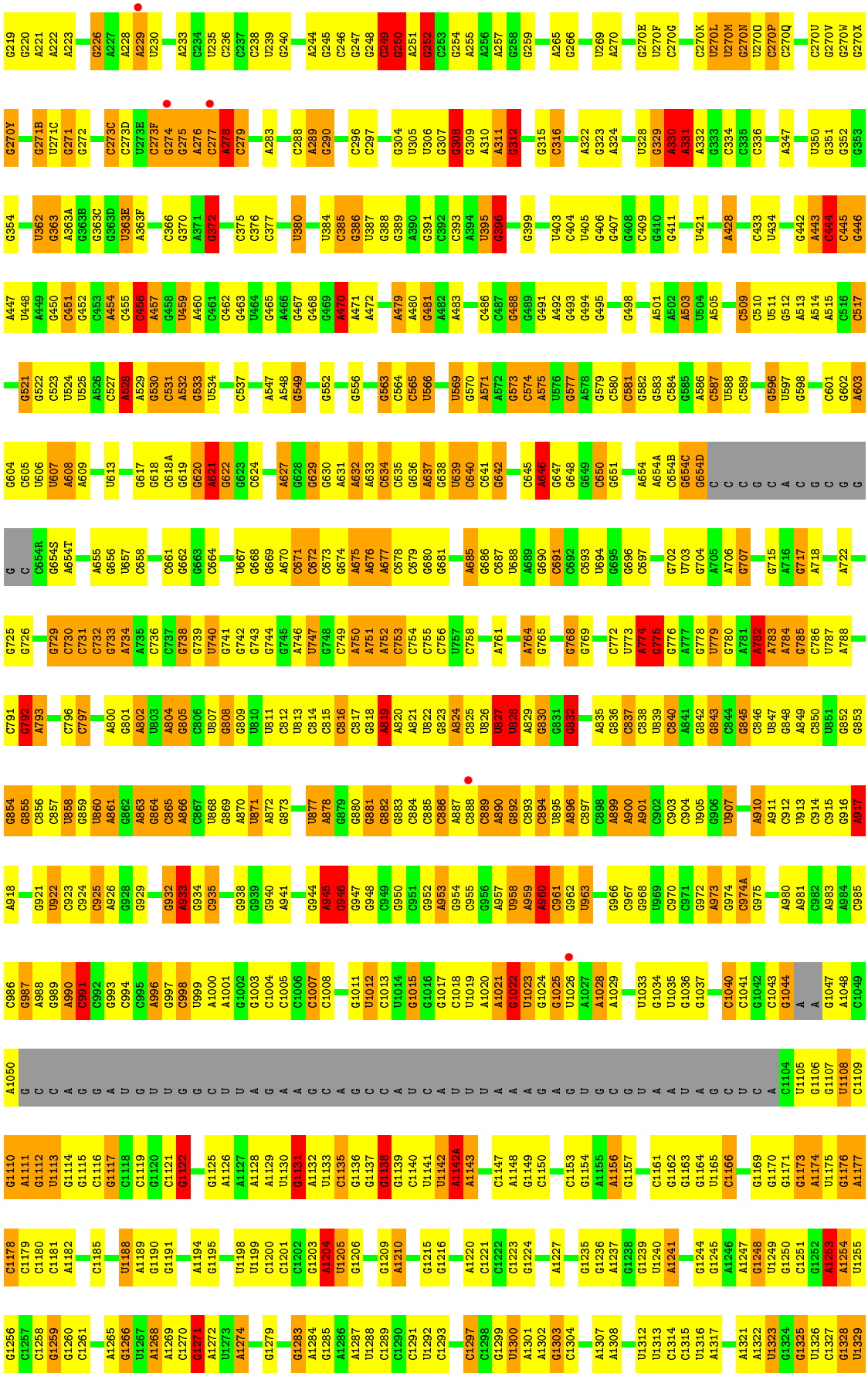




• Molecule 26: 23S ribosomal RNA

Chain 14:

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G270Y G271B U271C G271 G272 C273C C273D U273E C273F G274 G275 A276 C277 A278 C279 A283 C288 A289 G290 C296 C297 G304 U305 U306 G307 G308 G309 A310 A311 G312 G315 C316 A322 G323 A324 U328 G329 A330 A331 A332 G333 C334 C335 C336 A347 U350 G351 G352 G353
G354 U362 G363 A363A G363B G363C G363D U363E A363F C366 G370 A371 G372 C375 C376 C377 U380 U384 C385 G386 U387 G388 G389 A390 G391 C392 C393 A394 U395 G396 G399 U403 C404 U405 G406 G407 G408 C409 G410 G411 U421 A428 C433 U434 G442 A443 C444 C445 G446
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G521 G522 C523 U524 U525 A526 C527 A528 A529 G530 C531 A532 G533 U534 C537 A547 A548 G549 G552 G556 G563 C564 C565 U566 U569 G570 A571 A572 G573 C574 A575 U576 G577 A578 G579 C580 C581 G582 G583 C584 G585 A586 C587 U588 C589 G596 U597 G598 C601 G602 A603
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G G C G654R G654S A654T A655 G656 U657 C658 C661 G662 G663 C664 U667 G668 G669 A670 C671 C672 C673 G674 A675 A676 A677 C678 C679 G680 G681 A685 G686 C687 U688 A689 G690 C691 C692 C693 U694 G695 G696 C697 G702 U703 G704 A705 A706 G707 G715 A716 G717 A718 A722
G725 G726 G729 C730 C731 C732 G733 A734 A735 C736 C737 G738 G739 U740 G741 G742 G743 G744 G745 A746 U747 G748 C749 A750 A751 A752 C753 C754 C755 C756 U757 C758 A761 A764 G765 G768 G769 C772 U773 A774 G775 G776 A777 G778 U779 G780 A781 A782 A783 A784 G785 C786 U787 A788
C791 G792 A793 C796 C797 A800 G801 A802 U803 A804 G805 C806 U807 G808 G809 U810 U811 C812 U813 C814 C815 C816 C817 G818 A819 A820 A821 U822 G823 A824 C825 U826 U827 U828 A829 G830 G831 G832 A835 G836 C837 C838 U839 G840 A841 G842 G843 C844 G845 C846 U847 G848 A849 C850 U851 G852 G853
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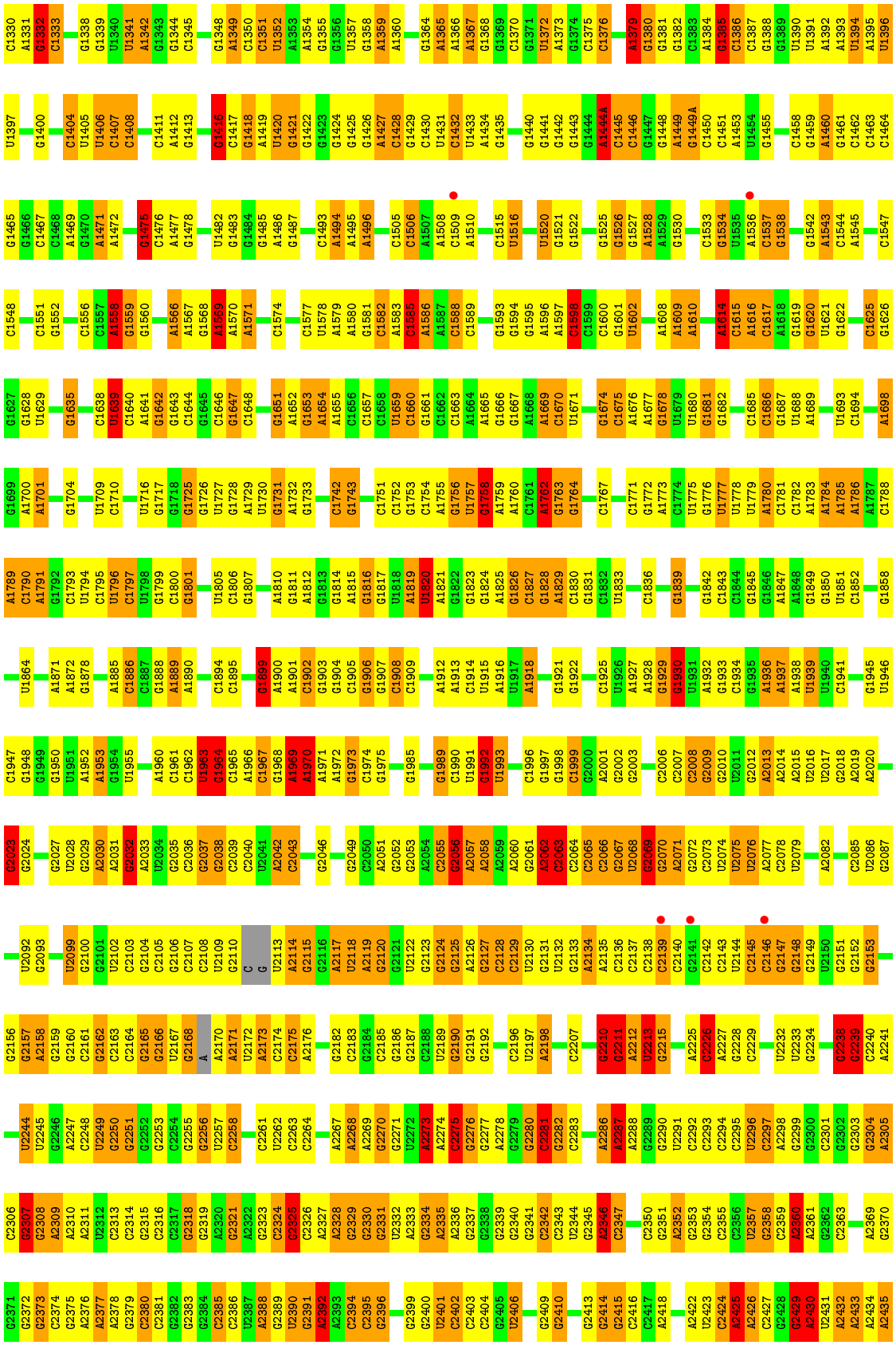
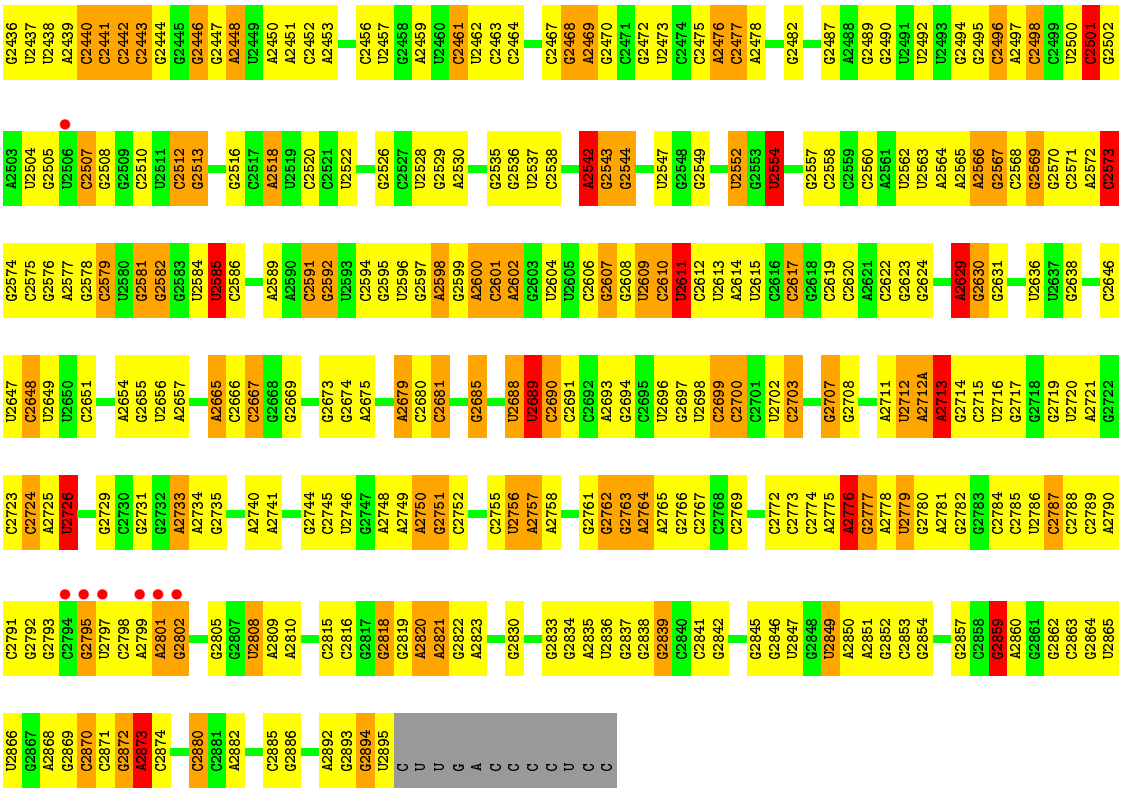
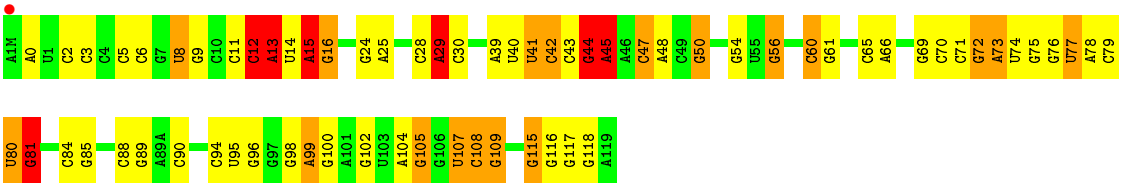
C1178 C1179 C1180 C1181 A1182 C1185 U1188 A1189 G1190 G1191 A1194 G1195 U1198 U1199 C1200 C1201 C1202 G1203 A1204 A1205 G1206 G1209 A1210 G1215 G1216 A1220 C1221 C1222 C1223 G1224 A1227 G1235 G1236 A1237 G1238 G1239 U1240 A1241 G1244 G1245 A1246 A1247 G1248 U1249 G1250 C1251 G1252 A1253 A1254 U1255
G1256 C1257 C1258 G1259 C1260 C1261 A1265 G1266 U1267 A1268 A1269 C1270 G1271 A1272 U1273 A1274 G1279 G1283 A1284 G1285 A1286 A1287 U1288 C1289 C1290 C1291 U1292 C1293 C1297 G1298 G1299 U1300 A1301 A1302 G1303 C1304 A1307 A1308 U1312 U1313 C1314 C1315 U1316 A1317 A1321 A1322 U1323 G1324 G1325 U1326 C1327 G1328 U1329





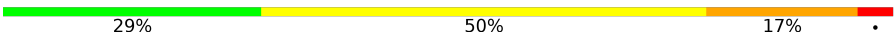
• Molecule 27: 5S ribosomal RNA

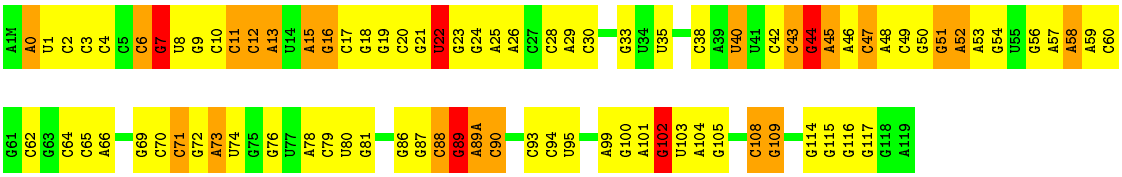
Chain 16:



• Molecule 27: 5S ribosomal RNA

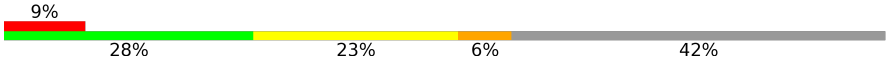
Chain 1J:

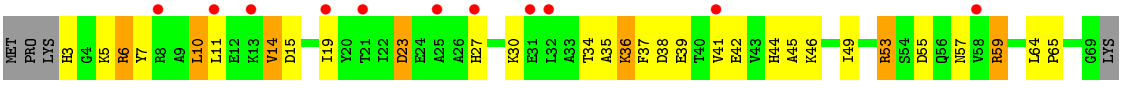


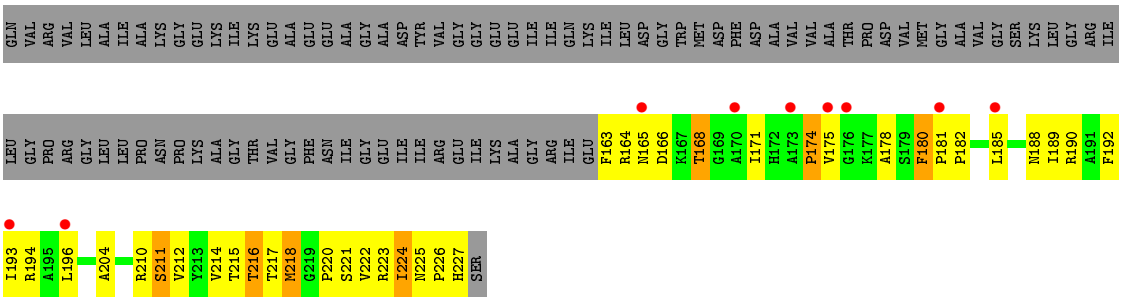


• Molecule 28: 50S ribosomal protein L1

Chain 71:

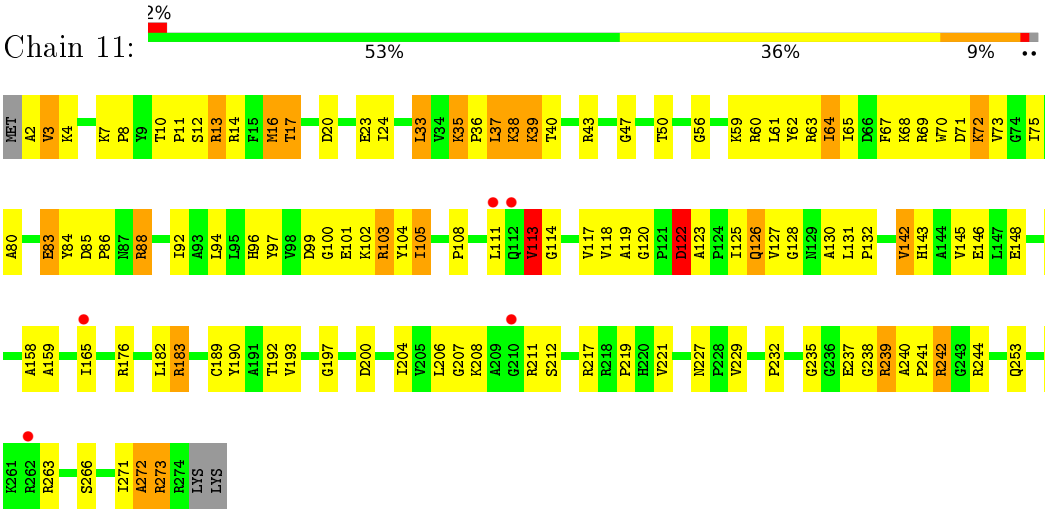






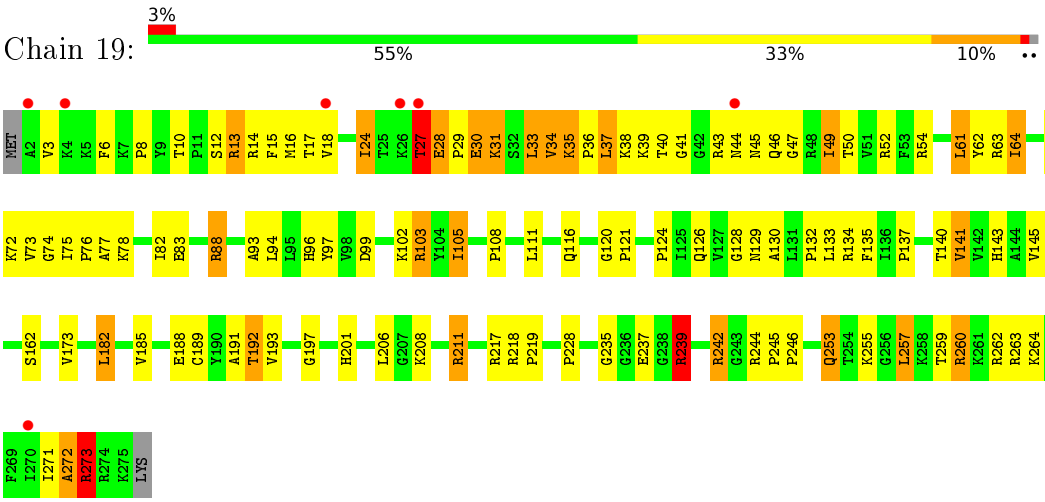
• Molecule 29: 50S ribosomal protein L2

Chain 11:



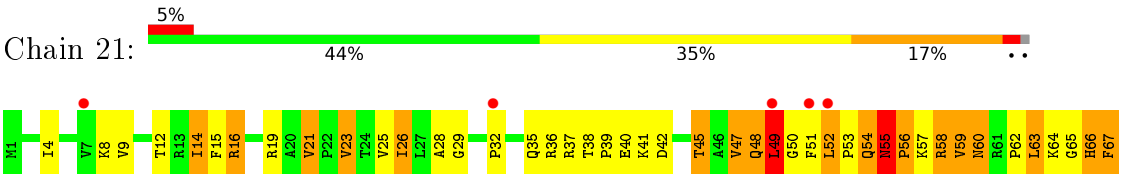
• Molecule 29: 50S ribosomal protein L2

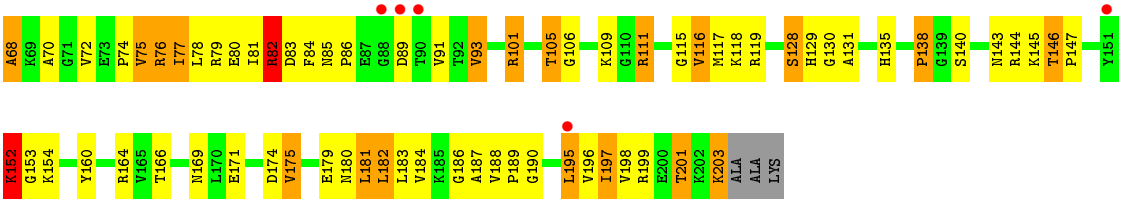
Chain 19:



• Molecule 30: 50S ribosomal protein L3

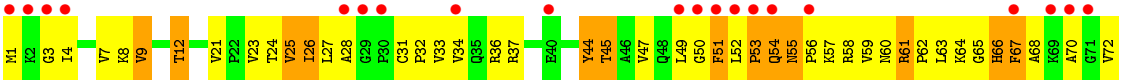
Chain 21:

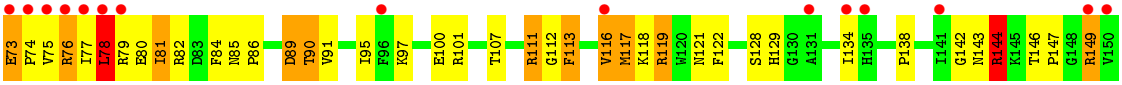


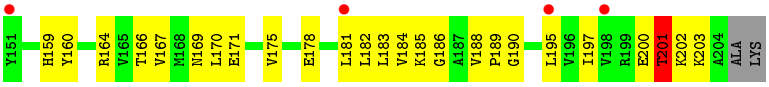


• Molecule 30: 50S ribosomal protein L3

Chain 29:

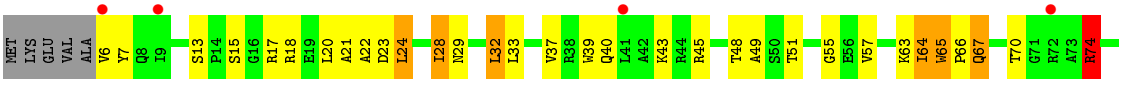


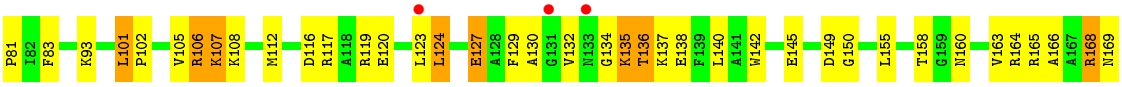


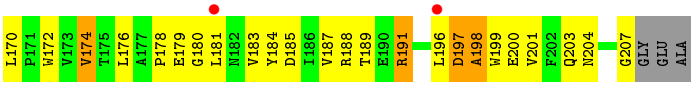


• Molecule 31: 50S ribosomal protein L4

Chain 31:

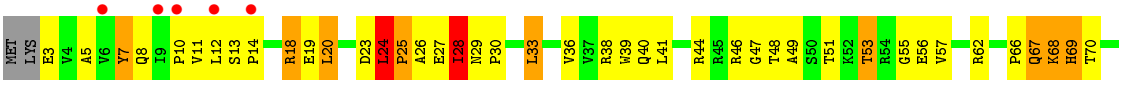


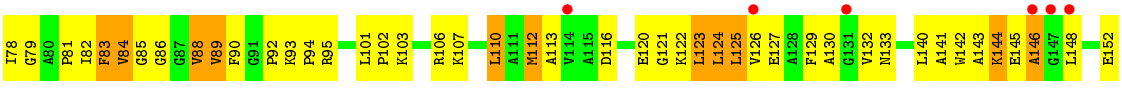


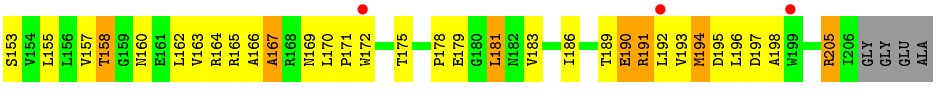


• Molecule 31: 50S ribosomal protein L4

Chain 39:

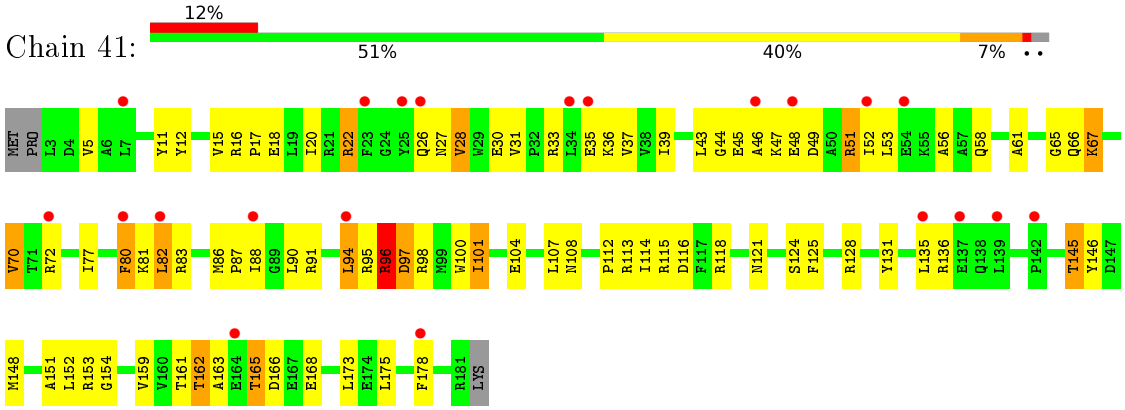






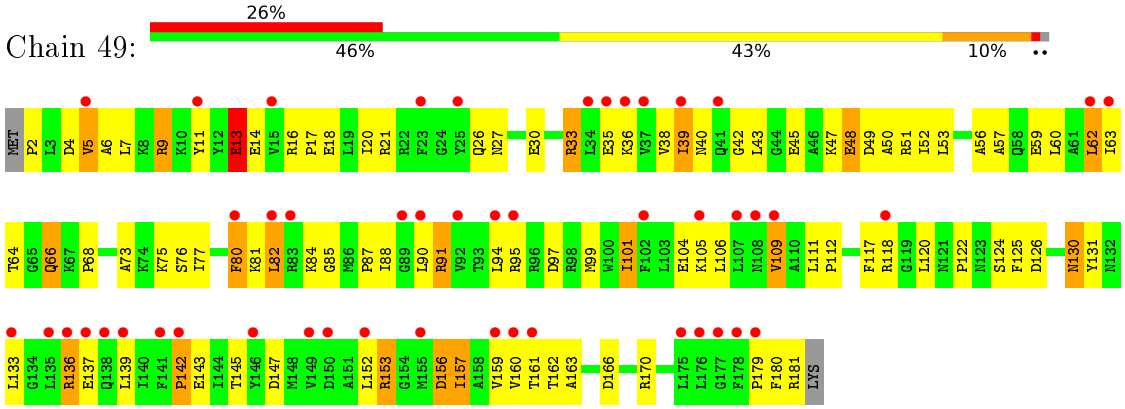
• Molecule 32: 50S ribosomal protein L5

Chain 41:



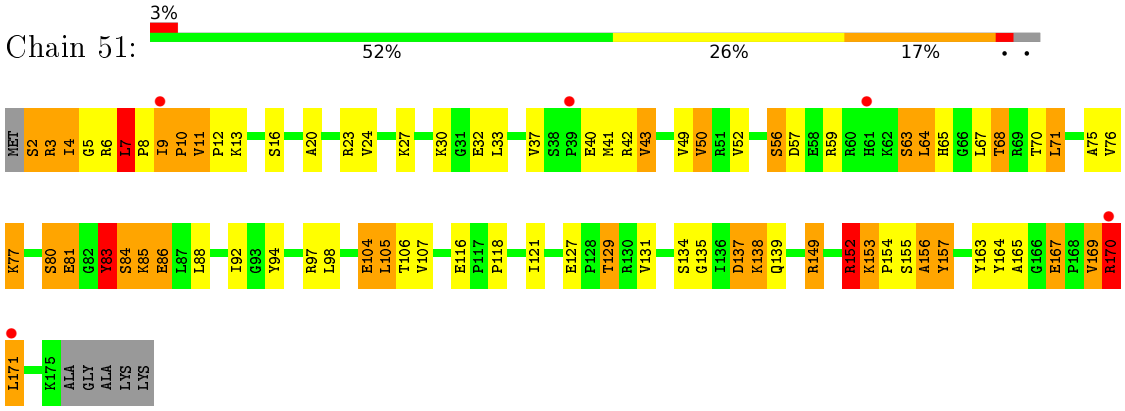
• Molecule 32: 50S ribosomal protein L5

Chain 49:



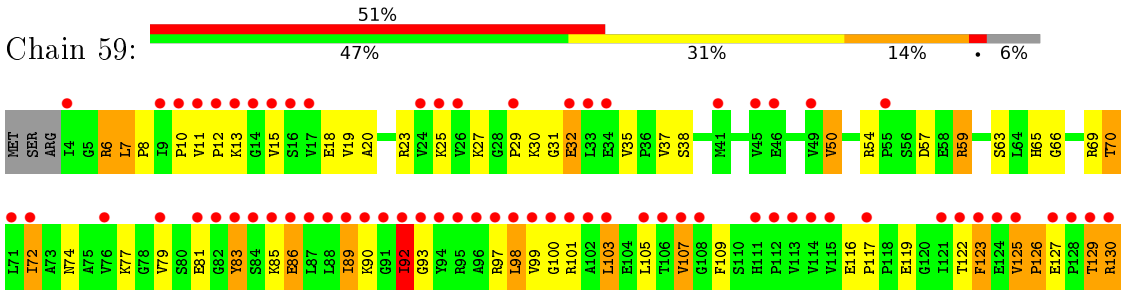
• Molecule 33: 50S ribosomal protein L6

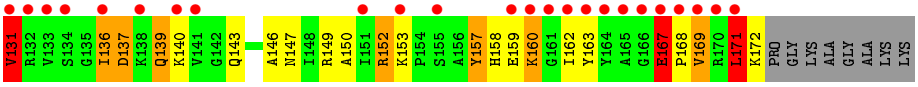
Chain 51:



• Molecule 33: 50S ribosomal protein L6

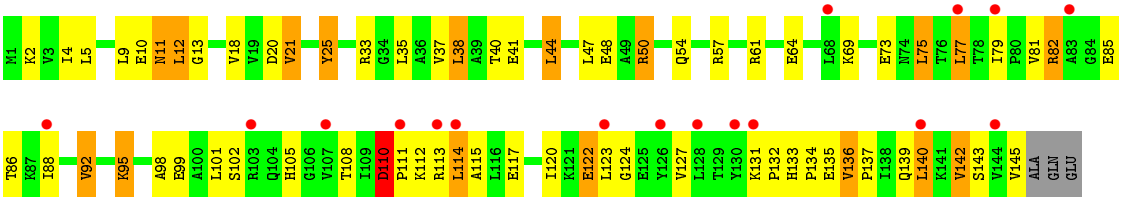
Chain 59:





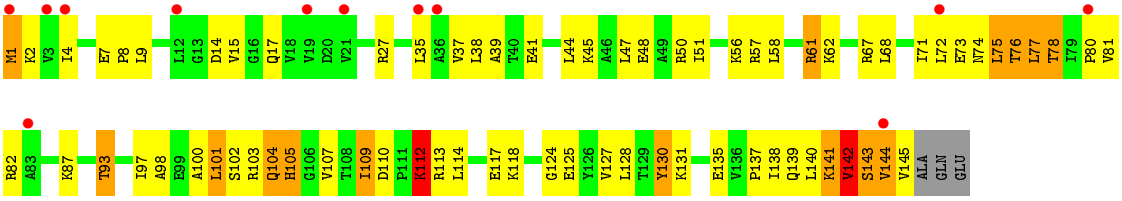
• Molecule 34: 50S ribosomal protein L9

Chain 61:



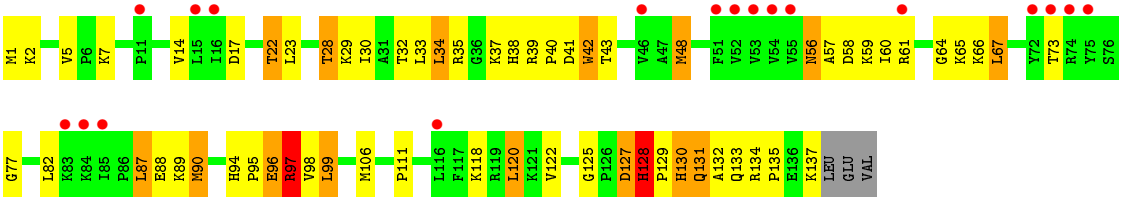
• Molecule 34: 50S ribosomal protein L9

Chain 69:



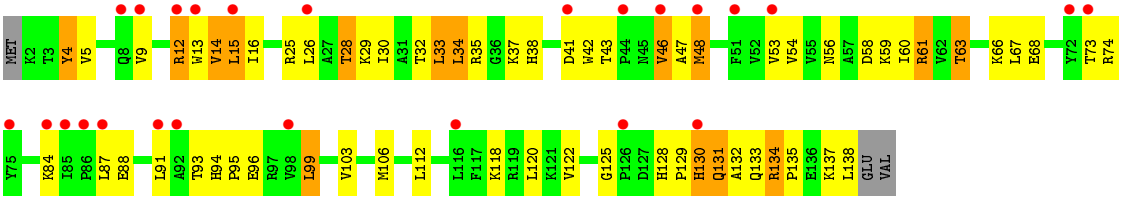
• Molecule 35: 50S ribosomal protein L13

Chain 58:



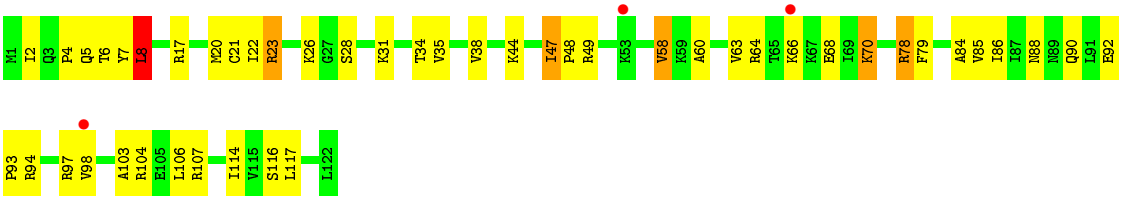
• Molecule 35: 50S ribosomal protein L13

Chain 15:



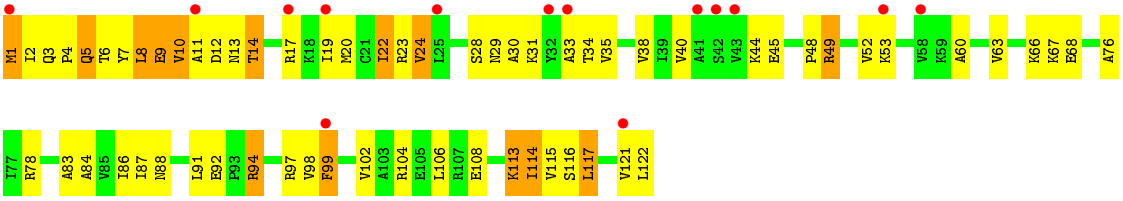
• Molecule 36: 50S ribosomal protein L14

Chain 68:



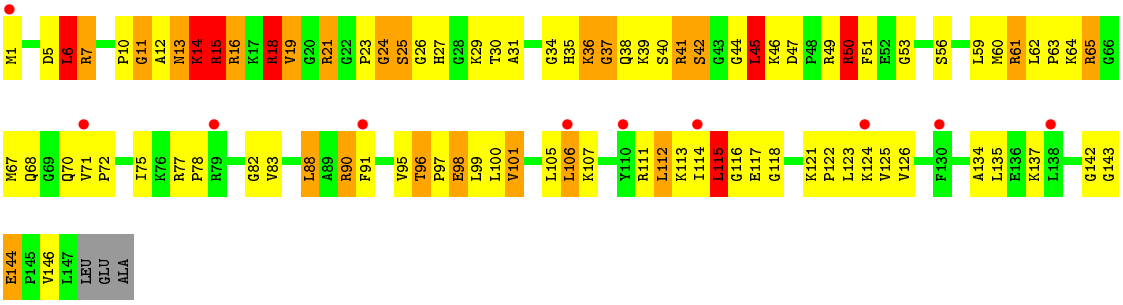
• Molecule 36: 50S ribosomal protein L14

Chain 25:



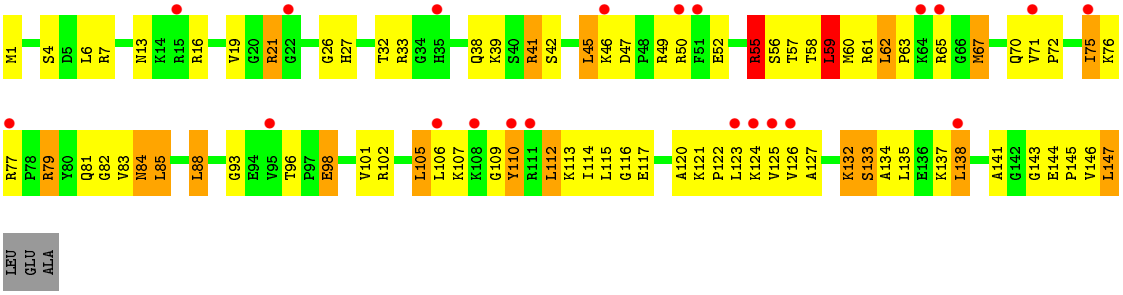
• Molecule 37: 50S ribosomal protein L15

Chain 78:



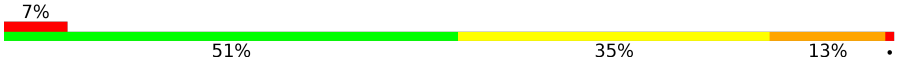
• Molecule 37: 50S ribosomal protein L15

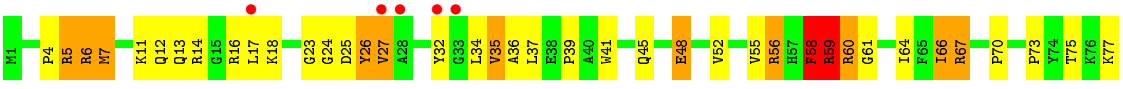
Chain 35:

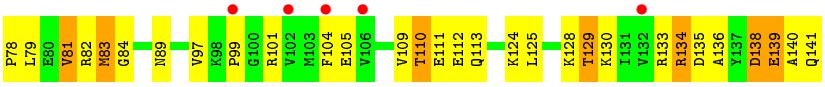


• Molecule 38: 50S ribosomal protein L16

Chain 88:

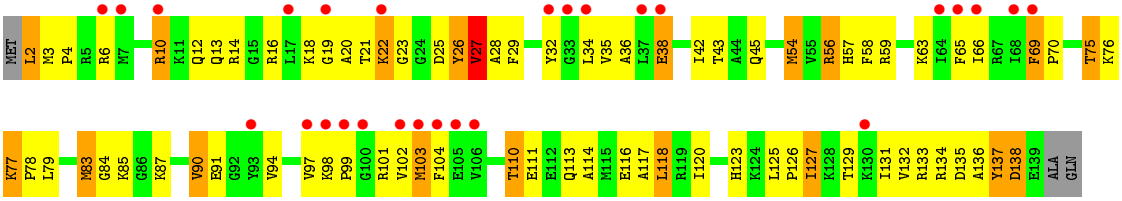






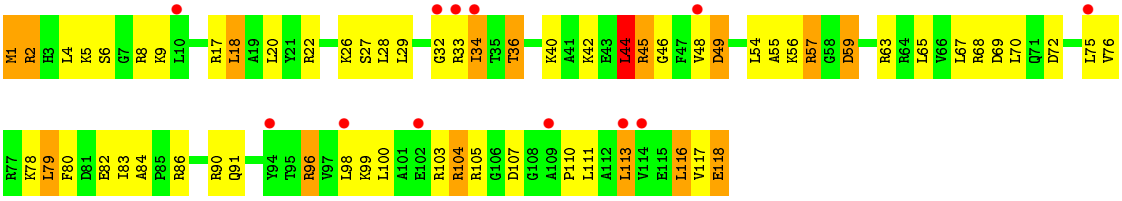
• Molecule 38: 50S ribosomal protein L16

Chain 45:



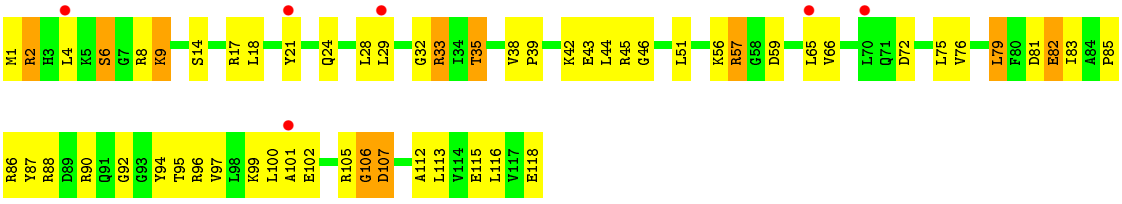
• Molecule 39: 50S ribosomal protein L17

Chain 98:



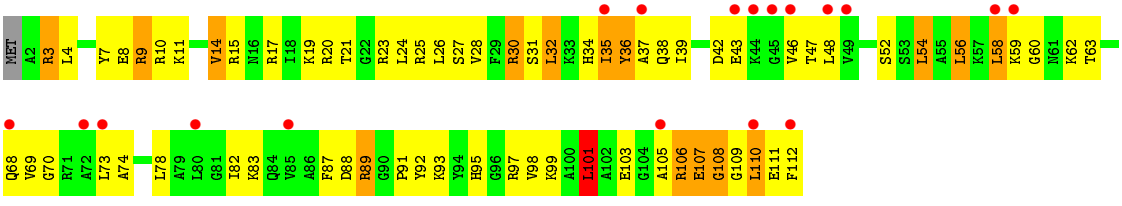
• Molecule 39: 50S ribosomal protein L17

Chain 55:



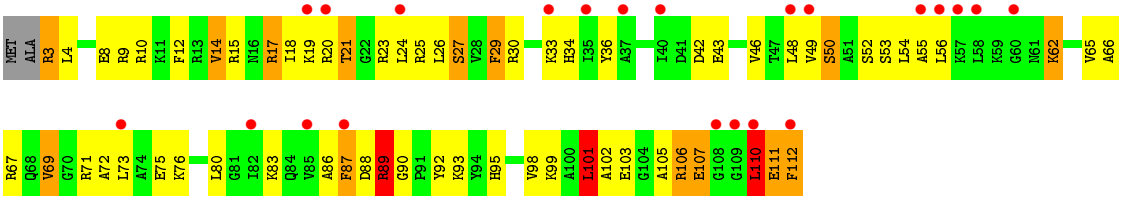
• Molecule 40: 50S ribosomal protein L18

Chain A8:



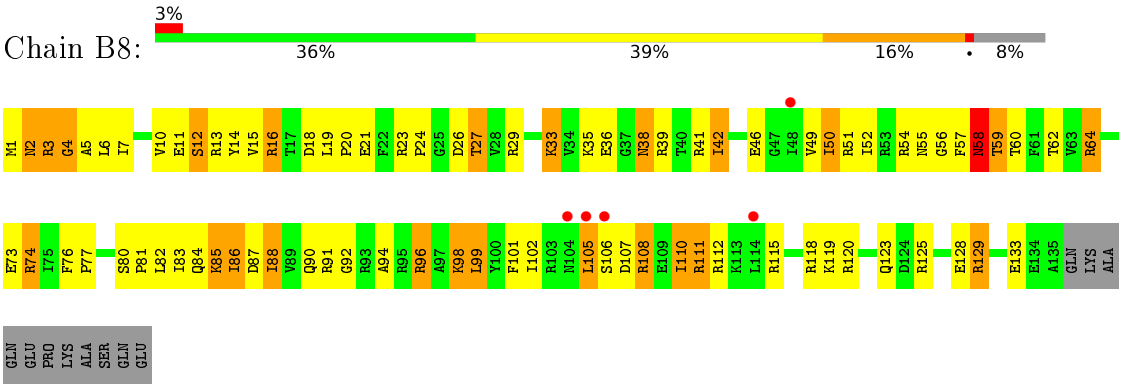
• Molecule 40: 50S ribosomal protein L18

Chain 65:



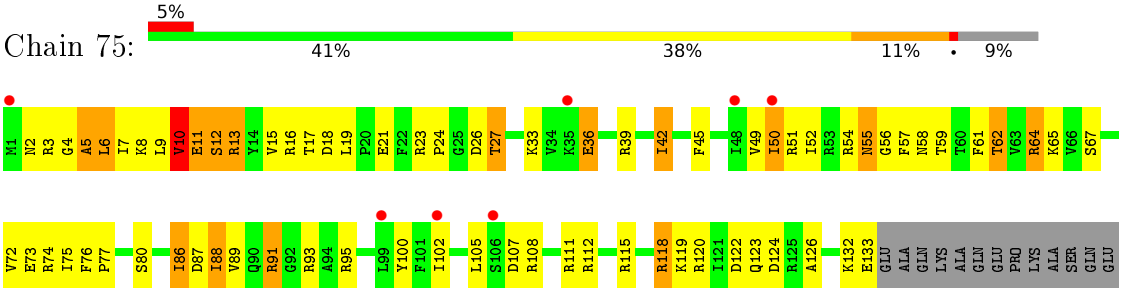
• Molecule 41: 50S ribosomal protein L19

Chain B8:



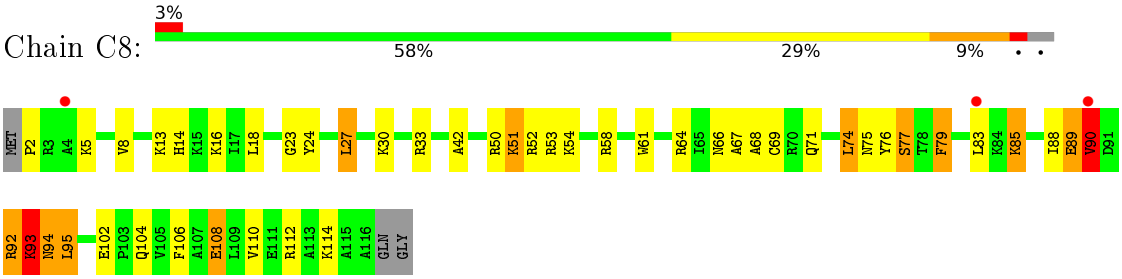
• Molecule 41: 50S ribosomal protein L19

Chain 75:



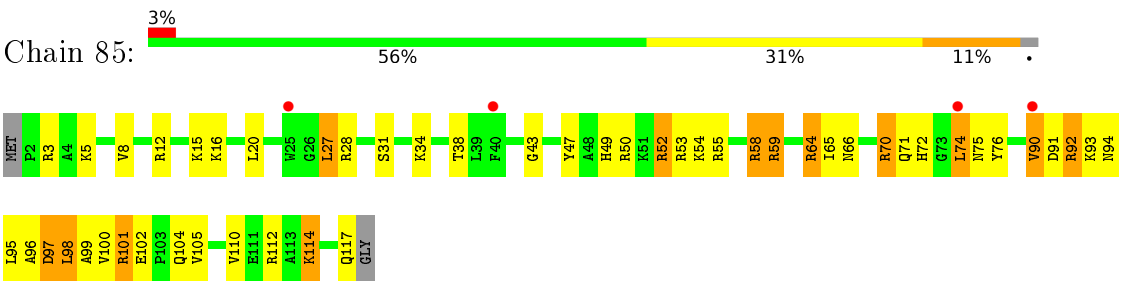
• Molecule 42: 50S ribosomal protein L20

Chain C8:



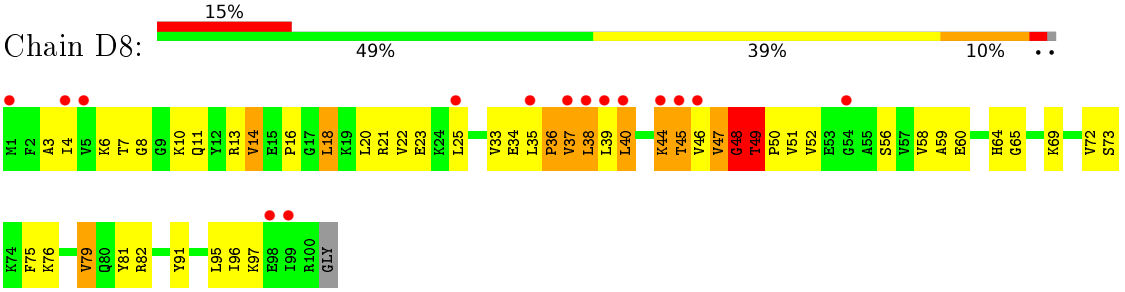
• Molecule 42: 50S ribosomal protein L20

Chain 85:



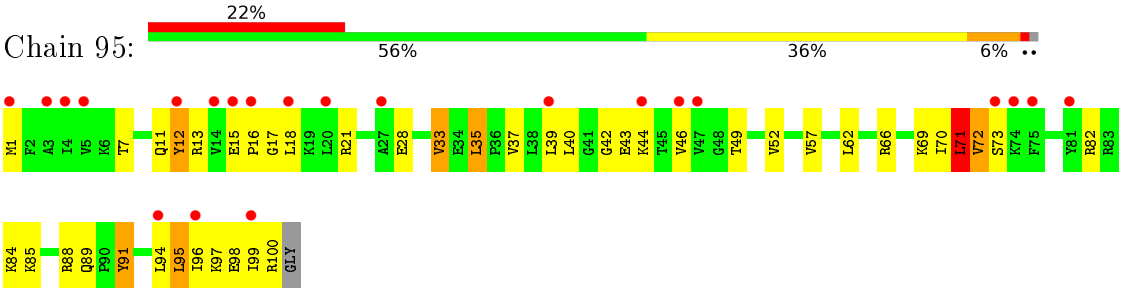
• Molecule 43: 50S ribosomal protein L21

Chain D8:



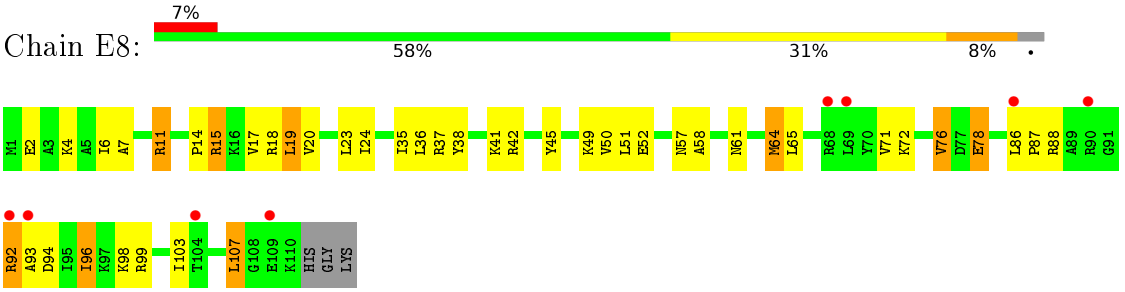
• Molecule 43: 50S ribosomal protein L21

Chain 95:



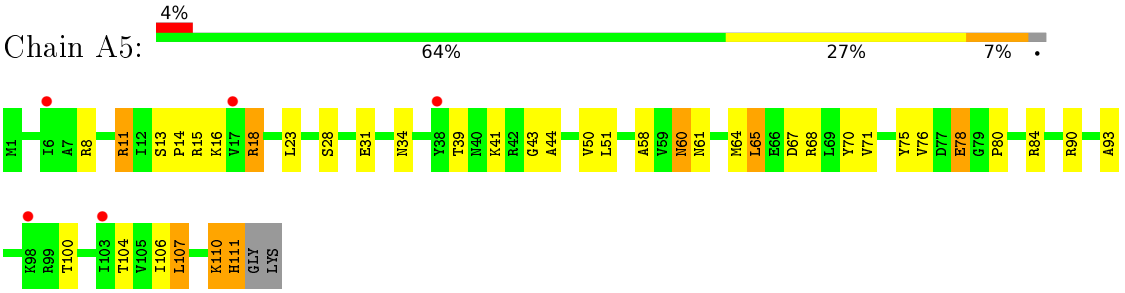
• Molecule 44: 50S ribosomal protein L22

Chain E8:



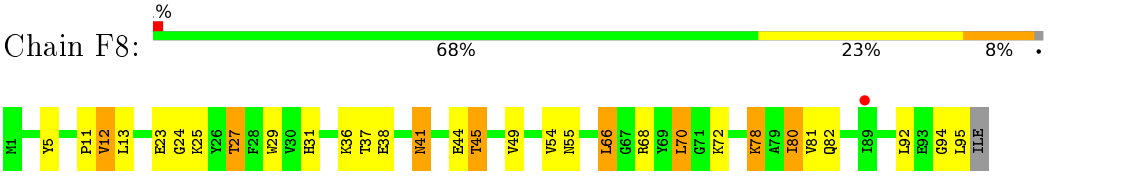
• Molecule 44: 50S ribosomal protein L22

Chain A5:



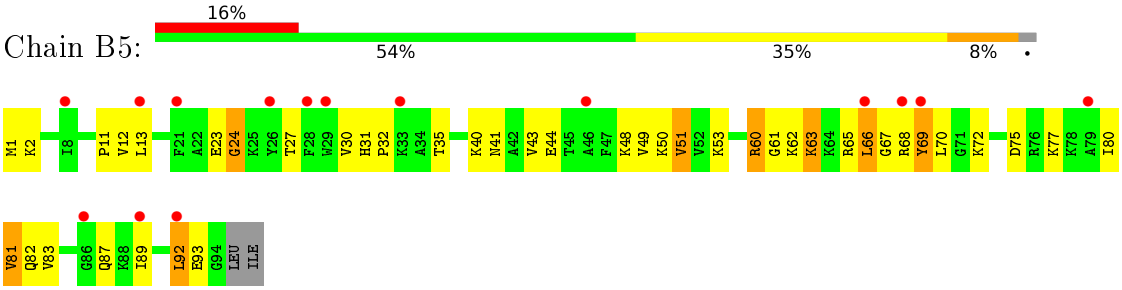
• Molecule 45: 50S ribosomal protein L23

Chain F8:



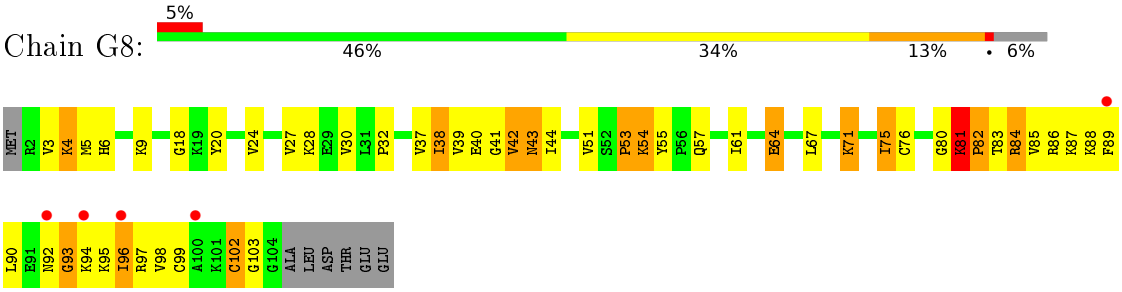
• Molecule 45: 50S ribosomal protein L23

Chain B5:



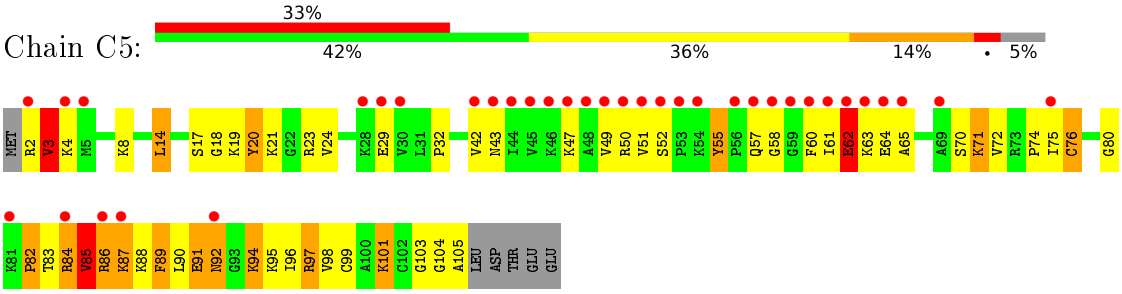
• Molecule 46: 50S ribosomal protein L24

Chain G8:



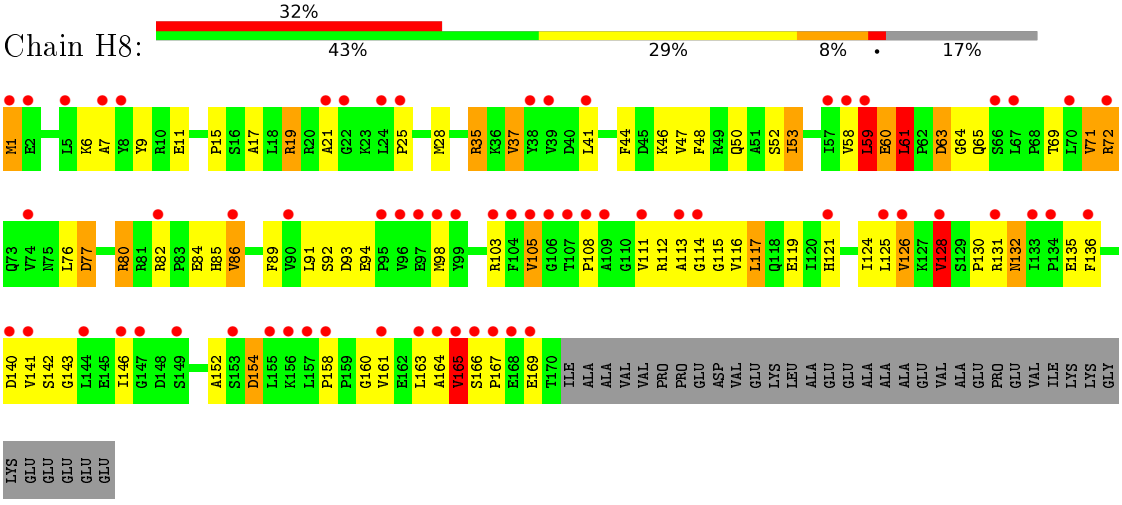
• Molecule 46: 50S ribosomal protein L24

Chain C5:



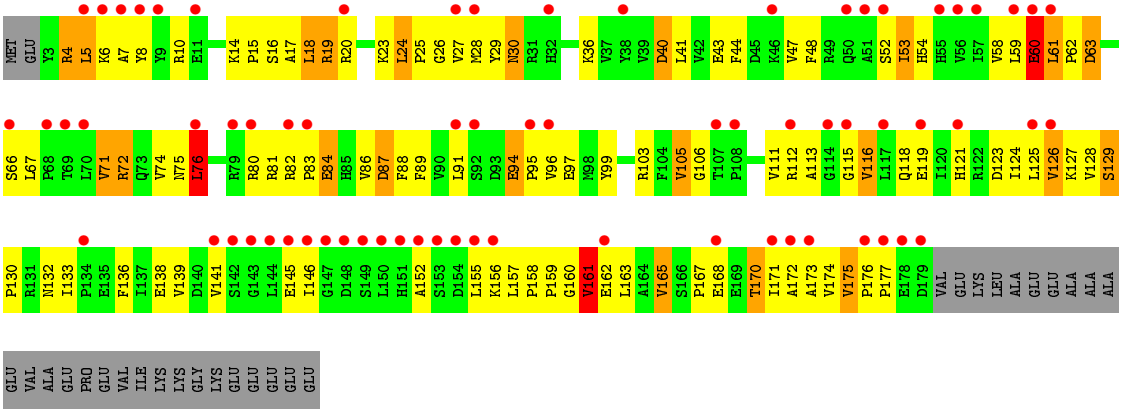
• Molecule 47: 50S ribosomal protein L25

Chain H8:



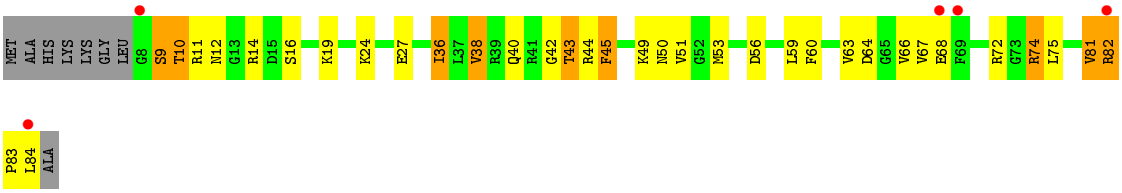
• Molecule 47: 50S ribosomal protein L25

Chain D5:



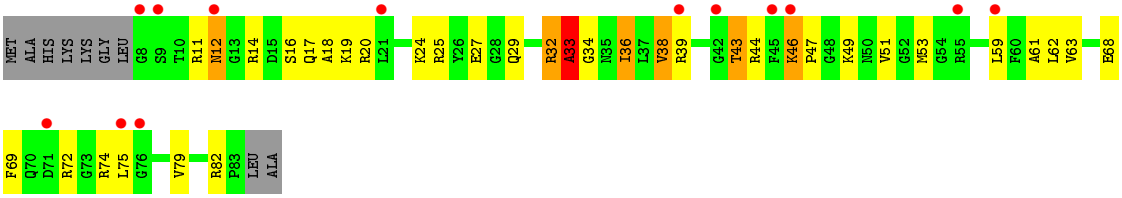
• Molecule 48: 50S ribosomal protein L27

Chain I8:



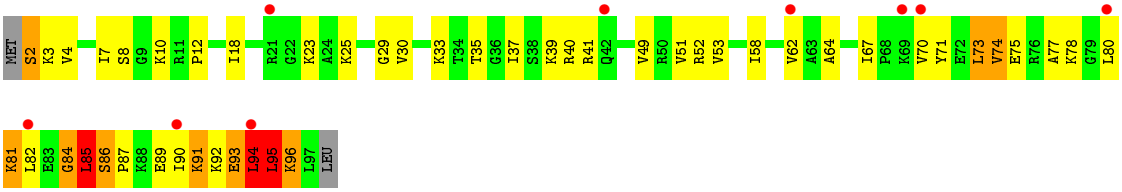
• Molecule 48: 50S ribosomal protein L27

Chain E5:



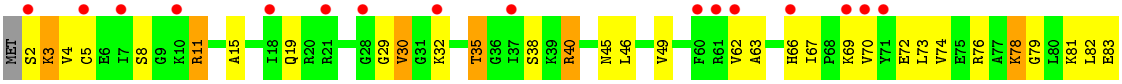
• Molecule 49: 50S ribosomal protein L28

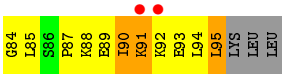
Chain J8:



• Molecule 49: 50S ribosomal protein L28

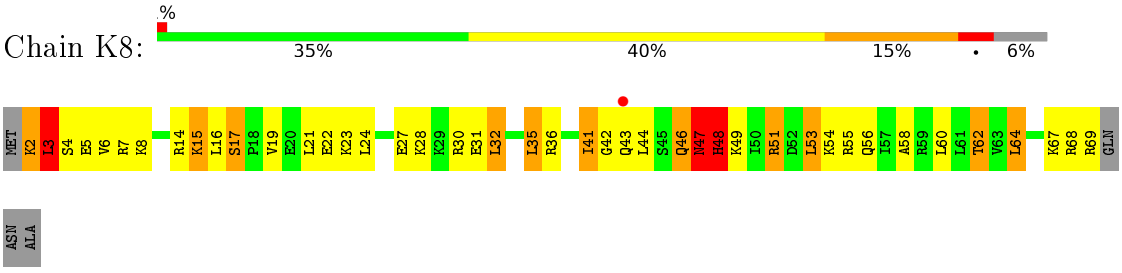
Chain F5:





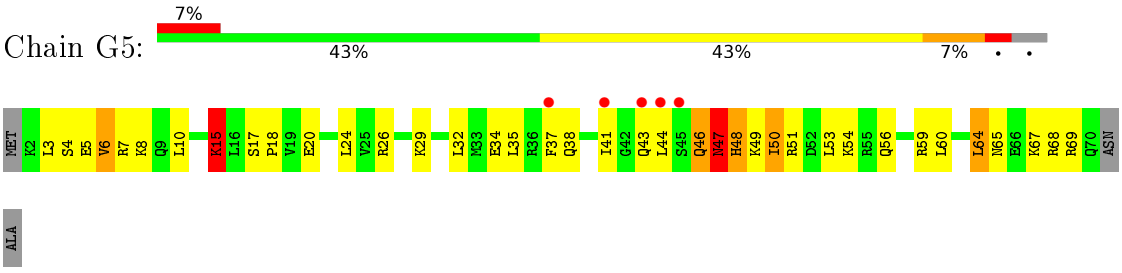
• Molecule 50: 50S ribosomal protein L29

Chain K8:



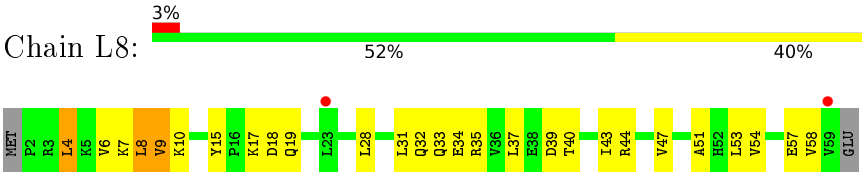
• Molecule 50: 50S ribosomal protein L29

Chain G5:



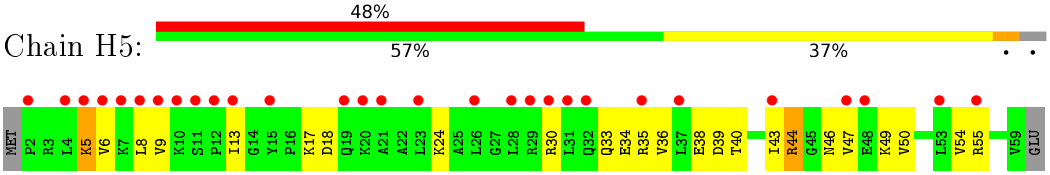
• Molecule 51: 50S ribosomal protein L30

Chain L8:



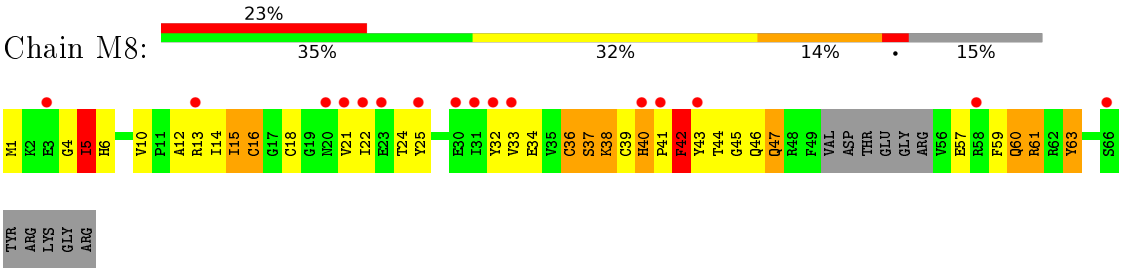
• Molecule 51: 50S ribosomal protein L30

Chain H5:



• Molecule 52: 50S ribosomal protein L31

Chain M8:

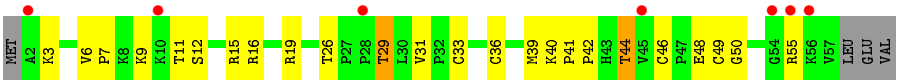


• Molecule 53: 50S ribosomal protein L32

Chain N8:

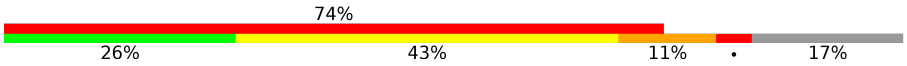
• Molecule 53: 50S ribosomal protein L32

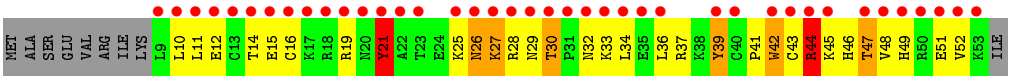
Chain J5:



• Molecule 54: 50S ribosomal protein L33

Chain O8:



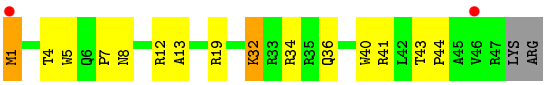


• Molecule 55: 50S ribosomal protein L34

Chain P8:

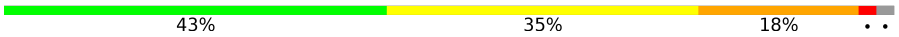
• Molecule 55: 50S ribosomal protein L34

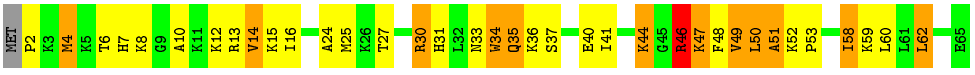
Chain L5:



• Molecule 56: 50S ribosomal protein L35

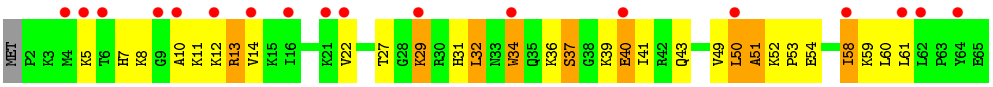
Chain Q8:





• Molecule 56: 50S ribosomal protein L35

Chain M5:



4 Data and refinement statistics (i)

Property	Value	Source
Space group	P 21 21 21	Depositor
Cell constants a, b, c, α, β, γ	209.30Å 448.80Å 620.20Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	147.08 – 2.99 161.81 – 2.99	Depositor EDS
% Data completeness (in resolution range)	99.9 (147.08-2.99) 90.8 (161.81-2.99)	Depositor EDS
R_{merge}	0.34	Depositor
R_{sym}	(Not available)	Depositor
$< I/\sigma(I) >$ [1]	1.76 (at 3.01Å)	Xtriage
Refinement program	PHENIX	Depositor
R, R_{free}	0.202 , 0.241 0.200 , 0.238	Depositor DCC
R_{free} test set	2000 reflections (0.19%)	DCC
Wilson B-factor (Å^2)	82.3	Xtriage
Anisotropy	0.334	Xtriage
Bulk solvent k_{sol}(e/Å^3), B_{sol}(Å^2)	0.28 , 75.7	EDS
L-test for twinning[2]	$< \|L\| > = 0.48$, $< L^2 > = 0.31$	Xtriage
Estimated twinning fraction	No twinning to report.	Xtriage
F_o,F_c correlation	0.95	EDS
Total number of atoms	297444	wwPDB-VP
Average B, all atoms (Å^2)	104.0	wwPDB-VP

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 1.49% of the height of the origin peak. No significant pseudotranslation is detected.*

[1] Intensities estimated from amplitudes.

[2] Theoretical values of $< |L| >$, $< L^2 >$ for acentric reflections are 0.5, 0.333 respectively for untwinned datasets, and 0.375, 0.2 for perfectly twinned datasets.

5 Model quality

5.1 Standard geometry

Bond lengths and bond angles in the following residue types are not validated in this section: 5MU, OMC, PAR, U8U, MG, SF4, ZN, 7MG, 4SU, T6A, SPE, PSU

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

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	#$\|Z\|>5$	RMSZ	#$\|Z\|>5$
1	13	0.70	1/36068 (0.0%)	1.35	292/56287 (0.5%)
1	1G	0.61	0/36282	1.22	170/56623 (0.3%)
2	12	0.37	0/1727	0.61	1/2326 (0.0%)
2	1E	0.40	0/1908	0.63	2/2573 (0.1%)
3	22	0.43	1/1560 (0.1%)	0.56	0/2104
3	2E	0.47	1/1629 (0.1%)	0.62	1/2195 (0.0%)
4	32	0.45	1/1732 (0.1%)	0.64	0/2318
4	3E	0.48	1/1728 (0.1%)	0.62	1/2313 (0.0%)
5	42	0.38	0/1156	0.60	0/1557
5	4E	0.45	0/1158	0.63	0/1559
6	52	0.49	0/855	0.66	1/1154 (0.1%)
6	5E	0.46	0/850	0.61	0/1147
7	62	0.39	0/1122	0.61	0/1500
7	6E	0.39	0/1259	0.54	0/1686
8	72	0.37	0/1127	0.57	0/1517
8	7E	0.41	0/1135	0.64	1/1527 (0.1%)
9	82	0.36	0/971	0.62	0/1304
9	8E	0.39	0/1019	0.61	0/1367
10	1A	1.00	2/658 (0.3%)	0.56	0/885
10	1I	0.40	0/762	0.62	0/1027
11	2A	0.40	0/850	0.61	1/1150 (0.1%)
11	2I	0.47	0/838	0.65	0/1133
12	3A	0.44	0/963	0.66	1/1290 (0.1%)
12	3I	0.63	0/972	0.80	1/1301 (0.1%)
13	4A	0.35	0/889	0.59	0/1192
13	4I	0.50	0/943	0.67	0/1265
14	5A	0.34	0/495	0.65	0/657
14	5I	0.47	0/495	0.69	1/657 (0.2%)
15	6A	0.40	0/740	0.58	0/987
15	6I	0.44	0/740	0.61	0/987
16	7A	0.43	0/721	0.65	0/970
16	7I	0.43	0/716	0.68	0/963

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	#|Z| >5	RMSZ	#|Z| >5
17	8A	0.45	0/836	0.59	0/1117
17	8I	0.51	0/836	0.65	0/1117
18	9A	0.46	0/549	0.67	0/732
18	9I	0.42	0/554	0.63	0/739
19	AA	0.38	0/520	0.67	0/700
19	AI	0.42	0/676	0.72	0/910
20	BA	0.35	0/764	0.67	1/1007 (0.1%)
20	BI	0.50	1/748 (0.1%)	0.63	0/986
21	1B	0.40	0/192	0.61	0/252
21	1F	0.42	0/203	0.71	0/266
22	1K	0.56	0/1589	1.11	6/2464 (0.2%)
23	2K	0.77	0/1721	1.42	16/2682 (0.6%)
23	2L	0.66	1/1721 (0.1%)	1.22	7/2682 (0.3%)
24	1L	0.39	0/1560	0.96	3/2418 (0.1%)
24	3K	0.52	0/1654	1.19	13/2570 (0.5%)
24	3L	0.52	0/1705	1.12	9/2650 (0.3%)
25	4K	0.74	0/473	1.15	0/735
25	4L	0.69	0/473	1.29	3/737 (0.4%)
26	14	0.84	45/68181 (0.1%)	1.54	1291/106432 (1.2%)
26	1H	1.01	117/68997 (0.2%)	1.75	2061/107696 (1.9%)
27	16	0.83	0/2928	1.57	59/4568 (1.3%)
27	1J	0.70	0/2928	1.37	32/4568 (0.7%)
28	71	0.29	0/1049	0.54	0/1417
29	11	0.66	1/2170 (0.0%)	0.90	4/2926 (0.1%)
29	19	0.64	1/2175 (0.0%)	0.85	2/2933 (0.1%)
30	21	0.57	0/1579	0.92	3/2131 (0.1%)
30	29	0.53	0/1596	0.80	3/2153 (0.1%)
31	31	0.63	1/1620 (0.1%)	0.90	3/2194 (0.1%)
31	39	0.51	0/1637	0.80	1/2218 (0.0%)
32	41	0.47	0/1481	0.68	0/1994
32	49	0.38	0/1483	0.63	1/1997 (0.1%)
33	51	0.53	0/1354	0.86	4/1833 (0.2%)
33	59	0.35	0/1320	0.69	3/1787 (0.2%)
34	61	0.41	0/1146	0.71	1/1551 (0.1%)
34	69	0.41	0/1146	0.70	1/1551 (0.1%)
35	15	0.41	0/1123	0.61	0/1515
35	58	0.52	0/1123	0.75	0/1514
36	25	0.48	0/942	0.71	1/1269 (0.1%)
36	68	0.55	0/942	0.76	2/1269 (0.2%)
37	35	0.53	0/1139	0.83	2/1514 (0.1%)
37	78	0.61	0/1139	1.04	8/1514 (0.5%)
38	45	0.61	2/1120 (0.2%)	0.81	0/1498
38	88	0.69	0/1138	0.92	1/1523 (0.1%)

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	#|Z| >5	RMSZ	#|Z| >5
39	55	0.51	0/981	0.83	0/1312
39	98	0.49	0/981	0.81	1/1312 (0.1%)
40	65	0.47	0/886	0.83	3/1180 (0.3%)
40	A8	0.59	0/891	0.83	2/1187 (0.2%)
41	75	0.51	0/1123	0.74	2/1500 (0.1%)
41	B8	0.58	0/1133	0.83	2/1514 (0.1%)
42	85	0.50	0/977	0.67	0/1301
42	C8	0.61	0/968	0.82	2/1289 (0.2%)
43	95	0.47	0/781	0.76	0/1048
43	D8	0.53	0/785	0.74	1/1052 (0.1%)
44	A5	0.55	0/897	0.77	1/1204 (0.1%)
44	E8	0.57	0/886	0.81	0/1189
45	B5	0.57	0/749	0.71	0/1007
45	F8	0.64	0/764	0.80	1/1025 (0.1%)
46	C5	0.52	0/807	0.81	1/1076 (0.1%)
46	G8	0.65	0/796	0.95	2/1062 (0.2%)
47	D5	0.54	1/1443 (0.1%)	0.64	1/1960 (0.1%)
47	H8	0.44	0/1395	0.73	1/1890 (0.1%)
48	E5	0.52	0/611	0.77	0/814
48	I8	0.76	1/619 (0.2%)	0.94	1/825 (0.1%)
49	F5	0.52	0/744	0.90	1/989 (0.1%)
49	J8	0.69	0/754	0.96	4/1003 (0.4%)
50	G5	0.53	0/578	0.73	0/766
50	K8	0.69	0/577	1.02	3/763 (0.4%)
51	H5	0.46	0/464	0.64	0/623
51	L8	0.50	0/464	0.73	0/623
52	M8	0.47	0/485	0.83	0/652
53	J5	0.58	0/448	0.76	0/606
53	N8	0.61	0/381	0.83	1/516 (0.2%)
54	O8	0.63	1/396 (0.3%)	0.90	1/529 (0.2%)
55	L5	0.57	0/409	0.78	0/540
55	P8	0.75	0/409	0.98	2/540 (0.4%)
56	M5	0.65	0/524	0.87	1/691 (0.1%)
56	Q8	0.69	0/524	1.02	3/691 (0.4%)
All	All	0.76	179/317928 (0.1%)	1.36	4051/476129 (0.9%)

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	5

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Mol	Chain	#Chirality outliers	#Planarity outliers
2	1E	0	4
4	32	0	2
9	82	0	1
9	8E	0	2
10	1A	0	1
11	2A	0	1
12	3A	0	1
12	3I	0	2
13	4A	0	1
13	4I	0	1
14	5A	0	1
19	AA	0	2
19	AI	0	1
20	BA	0	2
29	11	0	5
29	19	0	3
30	21	0	8
30	29	0	5
31	39	0	7
32	41	0	1
32	49	0	2
33	51	0	5
33	59	0	1
34	61	0	3
34	69	0	4
35	58	0	1
37	35	0	2
37	78	0	6
38	45	0	2
38	88	0	4
39	55	0	1
39	98	0	2
40	65	0	1
40	A8	0	2
41	75	0	3
41	B8	0	3
42	85	0	4
42	C8	0	2
43	D8	0	3
44	A5	0	1
45	B5	0	2
46	C5	0	2

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Mol	Chain	#Chirality outliers	#Planarity outliers
46	G8	0	5
47	D5	0	3
47	H8	0	6
48	E5	0	2
48	I8	0	1
49	F5	0	1
49	J8	0	1
50	G5	0	2
50	K8	0	2
52	M8	0	3
54	O8	0	1
55	P8	0	1
56	M5	0	2
56	Q8	0	3
All	All	0	145

All (179) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
10	1A	38	ILE	C-N	19.47	1.71	1.34
47	D5	94	GLU	C-N	14.46	1.61	1.34
10	1A	76	ASN	C-N	14.15	1.61	1.34
26	1H	774	A	N9-C4	-13.65	1.29	1.37
26	14	783	A	N9-C4	-12.02	1.30	1.37
26	1H	2430	A	N9-C4	-11.31	1.31	1.37
3	22	173	VAL	C-N	11.03	1.55	1.34
26	1H	783	A	N9-C4	-10.93	1.31	1.37
26	1H	2287	A	N9-C4	-10.75	1.31	1.37
26	1H	783	A	N3-C4	-10.14	1.28	1.34
26	1H	1899	G	N9-C4	-9.96	1.29	1.38
26	1H	676	A	N9-C4	-9.78	1.31	1.37
26	1H	1698	A	N9-C4	-9.63	1.32	1.37
26	1H	945	A	N7-C5	-9.55	1.33	1.39
26	1H	1698	A	N3-C4	-9.36	1.29	1.34
26	14	783	A	N3-C4	-9.25	1.29	1.34
26	1H	2346	A	N3-C4	-9.10	1.29	1.34
20	BI	97	ALA	C-N	9.06	1.51	1.34
26	1H	783	A	C5-C6	-8.99	1.32	1.41
26	1H	1332	G	N9-C4	-8.81	1.30	1.38
26	14	74	A	N9-C4	-8.65	1.32	1.37
26	14	1786	A	N9-C4	-8.52	1.32	1.37
3	2E	173	VAL	C-N	8.40	1.50	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
26	14	528	A	N9-C4	-8.40	1.32	1.37
26	1H	1678	G	N9-C8	8.34	1.43	1.37
26	1H	1786	A	N9-C4	-8.23	1.32	1.37
26	1H	528	A	N9-C4	-8.18	1.32	1.37
26	1H	676	A	N9-C8	8.12	1.44	1.37
26	1H	1899	G	N9-C8	8.06	1.43	1.37
26	1H	192	C	C2-N3	8.02	1.42	1.35
4	3E	36	ARG	C-N	-8.01	1.19	1.34
26	1H	1616	A	C5-C6	-7.95	1.33	1.41
26	1H	2062	A	N9-C4	7.92	1.42	1.37
26	1H	1950	G	N9-C8	7.88	1.43	1.37
26	14	2287	A	N9-C4	-7.86	1.33	1.37
26	1H	2062	A	N7-C5	7.84	1.44	1.39
26	1H	2062	A	N3-C4	7.69	1.39	1.34
26	14	783	A	N7-C5	-7.67	1.34	1.39
26	1H	621	A	N9-C4	-7.61	1.33	1.37
26	1H	1142(A)	A	N9-C4	-7.59	1.33	1.37
26	14	774	A	N9-C4	-7.56	1.33	1.37
26	1H	140	A	N9-C4	-7.47	1.33	1.37
26	1H	945	A	C5-C6	-7.31	1.34	1.41
26	1H	698	C	N1-C6	-7.31	1.32	1.37
26	1H	192	C	N3-C4	7.23	1.39	1.33
26	14	1772	G	N3-C4	7.21	1.40	1.35
26	14	1616	A	N9-C4	-7.17	1.33	1.37
38	45	77	LYS	C-N	-7.17	1.20	1.34
26	1H	71	A	N9-C4	-7.12	1.33	1.37
26	1H	71	A	C5-C6	-7.09	1.34	1.41
26	1H	71	A	C5-C4	7.05	1.43	1.38
26	1H	778	G	P-OP2	7.03	1.60	1.49
26	1H	1966	A	N9-C4	-6.99	1.33	1.37
26	1H	1786	A	C5-C4	6.96	1.43	1.38
26	1H	2392	A	N9-C8	6.88	1.43	1.37
26	1H	2062	A	C5-C6	6.78	1.47	1.41
26	14	783	A	C5-C6	-6.74	1.34	1.41
26	14	2685	G	C6-O6	6.74	1.30	1.24
26	1H	1786	A	N3-C4	-6.73	1.30	1.34
26	1H	138	G	N9-C8	6.72	1.42	1.37
26	1H	676	A	C5-C4	6.70	1.43	1.38
4	32	196	LEU	C-N	6.67	1.47	1.34
26	1H	774	A	N9-C8	6.66	1.43	1.37
26	1H	1786	A	C5-C6	-6.66	1.35	1.41
26	1H	829	A	N9-C4	-6.63	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
26	1H	2490	G	N9-C8	6.60	1.42	1.37
26	1H	1021	A	N9-C4	-6.52	1.33	1.37
26	1H	2713	A	C5-C4	6.52	1.43	1.38
26	1H	74	A	N9-C4	-6.44	1.33	1.37
26	1H	752	A	N9-C4	-6.43	1.33	1.37
26	1H	2051	A	N7-C5	-6.39	1.35	1.39
26	1H	1678	G	N9-C4	-6.36	1.32	1.38
26	1H	1614	A	N9-C4	-6.33	1.34	1.37
26	1H	1332	G	N9-C8	6.30	1.42	1.37
26	14	2346	A	N3-C4	-6.28	1.31	1.34
31	31	65	TRP	CB-CG	-6.21	1.39	1.50
38	45	69	PHE	C-N	6.20	1.46	1.34
26	1H	1332	G	N3-C4	-6.19	1.31	1.35
26	14	945	A	C5-C6	-6.18	1.35	1.41
26	14	2451	A	N9-C4	-6.16	1.34	1.37
26	14	676	A	N9-C4	-6.15	1.34	1.37
26	14	676	A	N9-C8	6.13	1.42	1.37
26	1H	1899	G	C2-N3	-6.09	1.27	1.32
26	1H	71	A	N9-C8	6.07	1.42	1.37
26	1H	1966	A	N3-C4	-6.06	1.31	1.34
26	14	945	A	N7-C5	-6.05	1.35	1.39
26	14	1903	G	N9-C8	-6.04	1.33	1.37
26	14	1786	A	N3-C4	-6.04	1.31	1.34
26	14	774	A	N9-C8	6.03	1.42	1.37
26	14	2062	A	N7-C5	6.01	1.42	1.39
26	14	786	C	N1-C6	5.99	1.40	1.37
26	1H	1776	G	C8-N7	-5.96	1.27	1.30
26	14	1698	A	N9-C4	-5.95	1.34	1.37
26	1H	825	C	N1-C6	-5.92	1.33	1.37
26	1H	2392	A	C5-C4	5.88	1.42	1.38
26	1H	1824	G	N7-C5	-5.87	1.35	1.39
26	1H	141	A	N9-C8	5.85	1.42	1.37
26	1H	530	G	N9-C8	5.83	1.42	1.37
26	1H	1775	U	C2-O2	-5.82	1.17	1.22
26	1H	573	G	P-OP1	5.79	1.58	1.49
26	1H	2518	A	N9-C4	-5.78	1.34	1.37
26	1H	2506	U	N1-C2	5.76	1.43	1.38
26	14	2873	A	N7-C5	-5.74	1.35	1.39
26	1H	2072	G	C8-N7	-5.72	1.27	1.30
26	1H	1204	A	N9-C4	-5.71	1.34	1.37
26	14	71	A	N9-C4	-5.70	1.34	1.37
26	1H	2689	U	C5-C6	-5.69	1.29	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
26	14	1786	A	C5-C6	-5.68	1.35	1.41
26	1H	57	C	N3-C4	-5.68	1.29	1.33
26	1H	471	A	N9-C4	-5.66	1.34	1.37
26	1H	1241	A	N9-C4	-5.65	1.34	1.37
26	14	2430	A	N9-C4	-5.65	1.34	1.37
26	1H	1349	A	C5-C4	5.64	1.42	1.38
26	14	2629	A	N9-C4	5.62	1.41	1.37
26	1H	576	U	N3-C4	-5.61	1.33	1.38
26	1H	621	A	C5-C4	5.58	1.42	1.38
23	2L	77	A	N9-C4	-5.58	1.34	1.37
26	1H	945	A	N3-C4	-5.58	1.31	1.34
26	14	1678	G	N9-C4	-5.57	1.33	1.38
26	1H	774	A	N3-C4	-5.56	1.31	1.34
26	1H	783	A	N7-C5	-5.53	1.35	1.39
26	14	1785	A	N7-C5	-5.52	1.35	1.39
26	1H	1786	A	N7-C5	-5.52	1.35	1.39
26	1H	621	A	C5-C6	-5.50	1.36	1.41
26	1H	1899	G	N3-C4	-5.50	1.31	1.35
26	1H	452	G	N1-C2	-5.47	1.33	1.37
26	14	974(A)	C	N1-C2	5.46	1.45	1.40
26	1H	2062	A	C6-N1	5.46	1.39	1.35
26	14	1899	G	C2-N3	5.46	1.37	1.32
26	14	1678	G	N3-C4	-5.45	1.31	1.35
26	1H	860	U	N1-C2	5.43	1.43	1.38
26	1H	945	A	N1-C2	5.43	1.39	1.34
26	14	788	A	N7-C5	-5.42	1.35	1.39
26	1H	917	A	C5-C6	-5.40	1.36	1.41
26	1H	253	C	N1-C6	5.40	1.40	1.37
26	14	1566	A	C5-C6	-5.40	1.36	1.41
26	1H	2688	U	N3-C4	-5.37	1.33	1.38
1	13	963	G	C6-N1	-5.34	1.35	1.39
26	1H	1616	A	N7-C5	-5.34	1.36	1.39
26	14	1021	A	N9-C4	-5.34	1.34	1.37
26	1H	74	A	N3-C4	-5.34	1.31	1.34
26	1H	2448	A	N7-C5	-5.32	1.36	1.39
26	14	774	A	N1-C2	5.31	1.39	1.34
26	1H	2589	A	C5-C4	-5.29	1.35	1.38
26	1H	472	A	N3-C4	-5.28	1.31	1.34
26	1H	1356	G	C6-O6	5.26	1.28	1.24
26	1H	188	G	C2-N3	5.26	1.36	1.32
26	1H	1349	A	N9-C8	5.25	1.42	1.37
26	14	2581	G	N1-C2	-5.25	1.33	1.37

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
26	14	2873	A	N3-C4	-5.25	1.31	1.34
26	1H	945	A	C2-N3	5.25	1.38	1.33
26	1H	761	A	N3-C4	-5.24	1.31	1.34
26	1H	1966	A	C5-C4	-5.23	1.35	1.38
26	1H	2430	A	N7-C5	-5.21	1.36	1.39
26	1H	1950	G	N3-C4	-5.21	1.31	1.35
26	14	1899	G	C5-C4	5.19	1.42	1.38
26	14	1786	A	C5-C4	5.18	1.42	1.38
26	1H	1950	G	C5-C4	5.18	1.42	1.38
26	14	2873	A	C5-C6	-5.17	1.36	1.41
26	1H	1678	G	C5-C4	5.17	1.42	1.38
26	1H	1032	A	C6-N1	5.17	1.39	1.35
26	1H	2707	G	N7-C5	5.16	1.42	1.39
26	1H	1313	U	C4-C5	-5.14	1.39	1.43
29	11	122	ASP	CB-CG	5.13	1.62	1.51
26	1H	1889	A	N9-C4	-5.12	1.34	1.37
26	1H	2497	A	N3-C4	-5.12	1.31	1.34
48	I8	45	PHE	C-N	5.11	1.45	1.34
54	O8	42	TRP	CB-CG	5.11	1.59	1.50
26	1H	774	A	C6-N1	5.09	1.39	1.35
26	1H	2392	A	N9-C4	-5.09	1.34	1.37
26	1H	2518	A	C5-C6	-5.09	1.36	1.41
26	1H	1621	U	N1-C6	-5.09	1.33	1.38
26	1H	1676	A	N9-C4	-5.08	1.34	1.37
26	14	2392	A	C5-C4	5.07	1.42	1.38
29	19	30	GLU	CG-CD	5.07	1.59	1.51
26	1H	452	G	C6-N1	-5.04	1.36	1.39
26	1H	2287	A	C5-C6	-5.01	1.36	1.41
26	14	2332	U	N1-C2	5.00	1.43	1.38
26	1H	463	G	N1-C2	-5.00	1.33	1.37

All (4051) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	1899	G	N3-C4-N9	-23.89	111.67	126.00
26	1H	945	A	C6-C5-N7	-20.89	117.67	132.30
26	1H	945	A	N1-C6-N6	20.76	131.05	118.60
26	1H	1899	G	N3-C4-C5	20.42	138.81	128.60
26	1H	2430	A	C2-N3-C4	-20.30	100.45	110.60
26	1H	1786	A	C2-N3-C4	-17.93	101.64	110.60
26	1H	71	A	C5-N7-C8	-17.86	94.97	103.90
26	1H	1786	A	C5-N7-C8	-17.71	95.05	103.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	2430	A	O5'-P-OP2	-17.54	89.65	110.70
26	1H	1786	A	N7-C8-N9	17.36	122.48	113.80
26	1H	71	A	N1-C6-N6	17.13	128.88	118.60
26	1H	2287	A	C2-N3-C4	-17.11	102.05	110.60
26	1H	1332	G	C5-N7-C8	-16.98	95.81	104.30
26	1H	1698	A	C2-N3-C4	-16.87	102.17	110.60
26	1H	1678	G	C2-N3-C4	-16.66	103.57	111.90
26	1H	1332	G	C2-N3-C4	-16.29	103.75	111.90
26	1H	945	A	C4-C5-C6	16.19	125.10	117.00
26	1H	576	U	N3-C2-O2	-16.13	110.91	122.20
26	1H	621	A	C2-N3-C4	-16.09	102.55	110.60
26	1H	74	A	C2-N3-C4	-15.98	102.61	110.60
26	1H	2346	A	N1-C2-N3	15.85	137.22	129.30
26	14	74	A	C2-N3-C4	-15.52	102.84	110.60
26	1H	774	A	N3-C4-C5	15.47	137.63	126.80
26	1H	71	A	C4-C5-N7	15.40	118.40	110.70
26	14	1786	A	C5-N7-C8	-15.24	96.28	103.90
26	1H	140	A	C5-N7-C8	-15.22	96.29	103.90
26	1H	774	A	N3-C4-N9	-15.19	115.25	127.40
26	1H	783	A	C5-N7-C8	-15.18	96.31	103.90
26	1H	783	A	C2-N3-C4	-14.98	103.11	110.60
26	1H	676	A	C2-N3-C4	-14.96	103.12	110.60
26	14	963	U	O5'-P-OP1	-14.82	92.36	105.70
26	1H	2689	U	C2-N1-C1'	14.73	135.37	117.70
26	14	1786	A	C2-N3-C4	-14.68	103.26	110.60
26	1H	917	A	C2-N3-C4	-14.56	103.32	110.60
26	1H	2490	G	C4-C5-N7	14.52	116.61	110.80
26	14	945	A	N1-C6-N6	14.51	127.31	118.60
26	14	945	A	C6-C5-N7	-14.49	122.16	132.30
26	14	510	C	O5'-P-OP2	-14.47	92.68	105.70
26	1H	2490	G	C5-N7-C8	-14.46	97.07	104.30
26	1H	945	A	C5-N7-C8	-14.40	96.70	103.90
26	14	783	A	N1-C6-N6	14.30	127.18	118.60
26	1H	1899	G	C2-N3-C4	-14.26	104.77	111.90
26	1H	1678	G	C5-N7-C8	-14.18	97.21	104.30
26	1H	860	U	C4-C5-C6	14.14	128.18	119.70
26	14	1786	A	N7-C8-N9	14.14	120.87	113.80
26	1H	576	U	C5-C4-O4	14.13	134.38	125.90
26	1H	1332	G	N3-C4-C5	14.05	135.62	128.60
26	1H	774	A	C2-N3-C4	-13.97	103.61	110.60
26	1H	1678	G	N3-C4-C5	13.92	135.56	128.60
26	1H	71	A	C6-C5-N7	-13.79	122.64	132.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	1899	G	N1-C2-N2	-13.74	103.83	116.20
26	14	783	A	C2-N3-C4	-13.70	103.75	110.60
26	1H	1616	A	C5-N7-C8	-13.67	97.07	103.90
26	1H	621	A	C5-N7-C8	-13.54	97.13	103.90
26	1H	1678	G	N3-C4-N9	-13.51	117.90	126.00
26	1H	71	A	C5-C6-N6	-13.49	112.91	123.70
26	14	2430	A	C2-N3-C4	-13.43	103.88	110.60
26	1H	71	A	N7-C8-N9	13.37	120.48	113.80
26	1H	676	A	C5-N7-C8	-13.33	97.23	103.90
26	1H	945	A	N7-C8-N9	13.32	120.46	113.80
26	14	1602	U	O5'-P-OP2	13.29	126.64	110.70
26	14	774	A	C2-N3-C4	-13.15	104.03	110.60
26	1H	192	C	C2-N3-C4	-13.08	113.36	119.90
26	1H	987	G	O5'-P-OP2	13.07	126.38	110.70
26	1H	945	A	C4-C5-N7	13.06	117.23	110.70
26	1H	1698	A	N1-C2-N3	12.90	135.75	129.30
26	1H	1332	G	N7-C8-N9	12.90	119.55	113.10
26	1H	1950	G	C5-N7-C8	-12.74	97.93	104.30
26	14	945	A	C2-N3-C4	-12.68	104.26	110.60
26	1H	1786	A	N1-C2-N3	12.68	135.64	129.30
26	1H	71	A	C2-N3-C4	-12.67	104.27	110.60
26	1H	624	C	O5'-P-OP1	-12.66	94.30	105.70
26	1H	140	A	N7-C8-N9	12.65	120.13	113.80
26	1H	1332	G	N3-C4-N9	-12.64	118.42	126.00
26	1H	1616	A	C4-C5-N7	12.59	117.00	110.70
26	1H	1786	A	C6-C5-N7	-12.55	123.52	132.30
26	1H	2582	G	O5'-P-OP2	-12.53	94.42	105.70
26	1H	783	A	N1-C6-N6	12.50	126.10	118.60
26	1H	1786	A	C8-N9-C4	-12.46	100.81	105.80
26	1H	2430	A	N3-C4-C5	12.41	135.49	126.80
26	1H	2430	A	N1-C6-N6	12.38	126.03	118.60
26	1H	2689	U	N3-C4-O4	12.35	128.04	119.40
26	1H	1332	G	C4-C5-N7	12.33	115.73	110.80
26	14	2873	A	C2-N3-C4	-12.32	104.44	110.60
26	1H	2346	A	O4'-C1'-N9	12.28	118.03	108.20
26	1H	49	A	O5'-P-OP2	-12.28	94.65	105.70
26	14	2873	A	C6-C5-N7	-12.27	123.71	132.30
26	1H	1678	G	N7-C8-N9	12.10	119.15	113.10
26	14	528	A	C2-N3-C4	-12.08	104.56	110.60
26	1H	929	G	N1-C6-O6	12.07	127.14	119.90
26	14	783	A	C5-N7-C8	-12.04	97.88	103.90
26	1H	1382	G	C5-C6-O6	-11.91	121.45	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	860	U	C5-C6-N1	-11.90	116.75	122.70
26	1H	1204	A	C2-N3-C4	-11.87	104.66	110.60
26	14	2287	A	C2-N3-C4	-11.87	104.67	110.60
26	14	1816	G	O5'-P-OP1	-11.85	95.04	105.70
26	1H	945	A	O4'-C1'-N9	11.78	117.62	108.20
27	16	13	A	O5'-P-OP2	-11.73	95.15	105.70
26	14	2873	A	N1-C2-N3	11.71	135.15	129.30
26	14	1616	A	C5-N7-C8	-11.70	98.05	103.90
26	14	1678	G	N7-C8-N9	11.70	118.95	113.10
26	14	1899	G	N3-C2-N2	11.67	128.07	119.90
26	1H	74	A	N1-C2-N3	11.62	135.11	129.30
26	14	1678	G	C8-N9-C4	-11.59	101.76	106.40
26	1H	1899	G	N3-C2-N2	-11.57	111.80	119.90
26	14	2873	A	N7-C8-N9	11.52	119.56	113.80
26	14	774	A	N3-C4-C5	11.49	134.84	126.80
26	1H	2406	U	O5'-P-OP1	-11.49	95.36	105.70
26	1H	917	A	C5-N7-C8	-11.48	98.16	103.90
26	1H	1382	G	N1-C6-O6	11.46	126.78	119.90
24	3K	76	A	C5-N7-C8	-11.45	98.17	103.90
26	1H	31	C	O5'-P-OP1	-11.43	95.41	105.70
26	1H	240	G	C5-C6-O6	-11.36	121.79	128.60
1	1G	766	A	O5'-P-OP2	-11.35	95.48	105.70
26	14	676	A	C2-N3-C4	-11.33	104.94	110.60
26	1H	793	A	O5'-P-OP2	-11.32	95.52	105.70
26	1H	783	A	C4-C5-N7	11.31	116.36	110.70
26	1H	2430	A	C5-C6-N1	-11.31	112.05	117.70
26	1H	2430	A	O5'-P-OP1	11.29	124.25	110.70
1	13	690	G	C6-C5-N7	-11.27	123.64	130.40
26	1H	676	A	N3-C4-C5	11.26	134.68	126.80
26	14	1763	G	O5'-P-OP2	-11.23	95.59	105.70
26	1H	1786	A	C4-C5-N7	11.22	116.31	110.70
26	1H	528	A	C6-N1-C2	11.21	125.33	118.60
26	1H	1950	G	N7-C8-N9	11.21	118.70	113.10
26	1H	2688	U	C5-C4-O4	11.19	132.61	125.90
26	14	1496	A	N7-C8-N9	11.19	119.39	113.80
26	1H	2714	G	O5'-P-OP2	-11.17	95.64	105.70
26	1H	2591	C	N1-C2-O2	-11.17	112.20	118.90
26	1H	528	A	N3-C4-C5	11.12	134.59	126.80
26	14	1678	G	C5-N7-C8	-11.06	98.77	104.30
1	13	974	A	N1-C6-N6	11.05	125.23	118.60
26	1H	1624	G	N1-C6-O6	-11.05	113.27	119.90
26	14	945	A	C4-C5-C6	11.04	122.52	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	1616	A	N1-C6-N6	11.03	125.22	118.60
26	1H	945	A	C5-C6-N6	-10.99	114.90	123.70
26	14	783	A	C6-C5-N7	-10.99	124.61	132.30
26	1H	2688	U	N3-C2-O2	-10.97	114.52	122.20
26	1H	945	A	C2-N3-C4	-10.93	105.14	110.60
26	1H	1496	A	N7-C8-N9	10.91	119.25	113.80
26	14	676	A	C5-N7-C8	-10.91	98.45	103.90
26	14	2873	A	N1-C6-N6	10.89	125.13	118.60
26	1H	140	A	C4-C5-N7	10.88	116.14	110.70
26	14	330	A	C2-N3-C4	-10.88	105.16	110.60
26	1H	576	U	N1-C2-N3	10.88	121.42	114.90
26	1H	2490	G	N7-C8-N9	10.80	118.50	113.10
26	1H	929	G	C5-C6-O6	-10.79	122.12	128.60
26	1H	783	A	N7-C8-N9	10.78	119.19	113.80
26	1H	1021	A	C2-N3-C4	-10.74	105.23	110.60
26	1H	1610	A	N9-C4-C5	-10.71	101.51	105.80
26	1H	1142(A)	A	C2-N3-C4	-10.71	105.25	110.60
1	13	690	G	C4-N9-C1'	10.70	140.41	126.50
24	3K	76	A	N7-C8-N9	10.68	119.14	113.80
26	1H	2346	A	C2-N3-C4	-10.68	105.26	110.60
26	1H	1639	U	O5'-P-OP2	-10.67	96.10	105.70
1	13	690	G	O4'-C1'-N9	10.66	116.73	108.20
26	1H	1604	C	O5'-P-OP1	-10.65	96.11	105.70
26	14	1772	G	C2-N3-C4	-10.64	106.58	111.90
26	1H	786	C	N3-C4-N4	-10.58	110.59	118.00
26	14	774	A	N1-C6-N6	10.58	124.95	118.60
26	14	2600	A	O5'-P-OP2	-10.57	96.19	105.70
26	1H	2448	A	N1-C6-N6	10.56	124.94	118.60
26	14	2873	A	C5-N7-C8	-10.54	98.63	103.90
26	1H	621	A	N7-C8-N9	10.51	119.05	113.80
26	14	788	A	N1-C6-N6	10.49	124.89	118.60
26	1H	945	A	C4-N9-C1'	10.48	145.16	126.30
26	1H	2287	A	N3-C4-C5	10.47	134.13	126.80
26	14	71	A	C5-N7-C8	-10.46	98.67	103.90
26	1H	2248	C	O5'-P-OP2	-10.43	96.31	105.70
26	1H	2392	A	C5-N7-C8	-10.43	98.69	103.90
26	14	621	A	C2-N3-C4	-10.42	105.39	110.60
26	1H	621	A	C4-C5-N7	10.42	115.91	110.70
26	14	1698	A	C5-N7-C8	-10.40	98.70	103.90
26	14	31	C	O5'-P-OP1	-10.40	96.34	105.70
26	1H	2331	G	N1-C6-O6	10.39	126.14	119.90
1	13	121	C	N1-C2-O2	10.38	125.13	118.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	2713	A	N7-C8-N9	10.37	118.98	113.80
26	1H	1204	A	O4'-C1'-N9	10.36	116.49	108.20
26	14	778	G	N1-C6-O6	-10.36	113.68	119.90
26	1H	138	G	C4-C5-N7	10.32	114.93	110.80
26	1H	917	A	N1-C6-N6	10.30	124.78	118.60
26	14	2092	U	C5-C4-O4	10.30	132.08	125.90
26	1H	1616	A	N7-C8-N9	10.29	118.94	113.80
26	1H	1698	A	C5-N7-C8	-10.28	98.76	103.90
26	14	2464	C	C6-N1-C2	10.28	124.41	120.30
26	1H	1899	G	N9-C4-C5	10.27	109.51	105.40
26	1H	459	U	O5'-P-OP2	-10.25	96.47	105.70
26	1H	945	A	N1-C2-N3	10.25	134.42	129.30
26	1H	1642	G	O5'-P-OP1	-10.25	96.48	105.70
26	1H	1565	C	C6-N1-C2	10.24	124.39	120.30
26	1H	1678	G	C8-N9-C4	-10.23	102.31	106.40
26	14	2273	A	O5'-P-OP2	-10.22	96.50	105.70
26	1H	1022	G	N9-C4-C5	10.21	109.48	105.40
26	1H	2287	A	C5-C6-N1	-10.21	112.59	117.70
26	1H	1313	U	C5-C6-N1	10.21	127.81	122.70
26	1H	2429	G	OP1-P-OP2	-10.20	104.30	119.60
26	1H	2265	U	O5'-P-OP1	-10.20	96.53	105.70
26	1H	2438	U	C5-C6-N1	-10.17	117.62	122.70
26	1H	1899	G	C8-N9-C1'	10.16	140.21	127.00
26	1H	1950	G	C8-N9-C4	-10.11	102.36	106.40
26	1H	736	C	O5'-P-OP1	-10.10	96.61	105.70
26	14	216	A	O5'-P-OP1	-10.09	96.62	105.70
26	1H	1955	U	C5-C6-N1	-10.09	117.66	122.70
26	14	945	A	N1-C2-N3	10.09	134.34	129.30
26	14	945	A	C5-N7-C8	-10.08	98.86	103.90
26	14	2357	U	O5'-P-OP2	-10.07	96.63	105.70
26	1H	2713	A	C5-N7-C8	-10.06	98.87	103.90
26	14	74	A	N3-C4-C5	10.06	133.84	126.80
26	14	974(A)	C	N1-C2-O2	10.05	124.93	118.90
26	1H	1129	A	O5'-P-OP2	-10.04	96.66	105.70
26	1H	774	A	C5-C6-N1	-10.03	112.69	117.70
31	31	74	ARG	NE-CZ-NH1	10.00	125.30	120.30
26	1H	774	A	C5-N7-C8	-9.99	98.91	103.90
26	1H	2518	A	N1-C6-N6	9.98	124.59	118.60
26	1H	1678	G	C4-C5-N7	9.98	114.79	110.80
26	1H	2700	C	C6-N1-C2	9.96	124.28	120.30
1	13	330	C	N1-C2-O2	9.96	124.88	118.90
26	1H	783	A	C6-C5-N7	-9.94	125.34	132.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	2609	U	C5-C6-N1	-9.92	117.74	122.70
26	1H	34	C	O5'-P-OP1	-9.92	96.78	105.70
26	1H	2689	U	C6-N1-C1'	-9.92	107.31	121.20
26	14	1619	G	O5'-P-OP2	-9.90	96.79	105.70
26	1H	917	A	N1-C2-N3	9.90	134.25	129.30
26	14	1204	A	C2-N3-C4	-9.90	105.65	110.60
26	1H	392	C	O5'-P-OP1	-9.85	96.83	105.70
26	1H	1616	A	C6-C5-N7	-9.85	125.41	132.30
26	14	140	A	N7-C8-N9	9.85	118.72	113.80
26	1H	2688	U	C4-C5-C6	9.84	125.60	119.70
26	1H	787	U	O5'-P-OP1	-9.84	96.85	105.70
26	1H	2584	U	N3-C2-O2	-9.83	115.32	122.20
26	1H	1255	U	N3-C4-O4	9.82	126.27	119.40
26	14	1772	G	N9-C4-C5	-9.82	101.47	105.40
26	14	2346	A	O4'-C1'-N9	9.81	116.05	108.20
26	14	2713	A	C5-N7-C8	-9.81	99.00	103.90
26	14	672	C	O5'-P-OP2	-9.80	96.88	105.70
26	1H	676	A	N3-C4-N9	-9.80	119.56	127.40
26	1H	71	A	N1-C2-N3	9.79	134.20	129.30
26	14	694	U	O5'-P-OP2	-9.79	96.89	105.70
26	14	1332	G	C5-N7-C8	-9.78	99.41	104.30
26	1H	1610	A	N1-C6-N6	9.78	124.47	118.60
26	14	140	A	C5-N7-C8	-9.77	99.02	103.90
26	1H	140	A	N1-C6-N6	9.76	124.46	118.60
26	1H	2432	A	O5'-P-OP2	-9.74	96.93	105.70
26	14	1786	A	C4-C5-N7	9.74	115.57	110.70
26	1H	1496	A	C5-N7-C8	-9.74	99.03	103.90
26	14	2430	A	N1-C6-N6	9.73	124.44	118.60
26	1H	2287	A	N1-C6-N6	9.73	124.44	118.60
26	14	1325	G	O5'-P-OP2	9.72	122.37	110.70
26	1H	528	A	C5-C6-N1	-9.72	112.84	117.70
26	1H	2032	G	C2-N3-C4	-9.71	107.05	111.90
26	14	621	A	C5-C6-N1	-9.69	112.85	117.70
1	13	1195	C	C6-N1-C2	-9.69	116.42	120.30
26	1H	2352	A	O5'-P-OP1	-9.69	96.98	105.70
26	1H	2503	A	N1-C2-N3	-9.69	124.45	129.30
26	1H	2036	C	O5'-P-OP1	-9.69	96.98	105.70
26	1H	210	C	C6-N1-C2	9.68	124.17	120.30
26	1H	1589	C	O5'-P-OP2	9.68	122.32	110.70
26	1H	2688	U	C5-C6-N1	-9.68	117.86	122.70
26	14	1616	A	C4-C5-N7	9.68	115.54	110.70
26	1H	621	A	N1-C6-N6	9.67	124.40	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	125	G	C5-C6-O6	-9.67	122.80	128.60
26	1H	1698	A	C6-C5-N7	-9.66	125.53	132.30
26	1H	1950	G	C4-C5-N7	9.65	114.66	110.80
26	1H	192	C	C5-C4-N4	-9.63	113.46	120.20
26	1H	917	A	C4-C5-N7	9.62	115.51	110.70
26	1H	930	U	C5-C4-O4	9.62	131.67	125.90
26	1H	2062	A	C2-N3-C4	9.62	115.41	110.60
26	1H	1021	A	C5-N7-C8	-9.61	99.10	103.90
26	1H	913	U	O5'-P-OP2	-9.60	97.06	105.70
26	1H	2689	U	N1-C2-O2	9.60	129.52	122.80
26	14	530	G	C4-C5-N7	9.60	114.64	110.80
26	14	963	U	O5'-P-OP2	9.59	122.21	110.70
26	14	1914	C	C6-N1-C2	-9.59	116.47	120.30
26	14	2873	A	C4-C5-C6	9.58	121.79	117.00
26	1H	2689	U	N3-C2-O2	-9.58	115.50	122.20
26	14	2518	A	N1-C6-N6	9.57	124.34	118.60
26	1H	329	G	O5'-P-OP2	-9.57	97.09	105.70
26	1H	752	A	C8-N9-C4	9.56	109.62	105.80
26	1H	141	A	C5-N7-C8	-9.56	99.12	103.90
26	1H	1698	A	N1-C6-N6	9.55	124.33	118.60
26	14	1678	G	N3-C4-N9	-9.55	120.27	126.00
26	1H	2507	C	C6-N1-C2	-9.54	116.48	120.30
26	14	462	C	O5'-P-OP2	-9.54	97.12	105.70
26	1H	676	A	N7-C8-N9	9.52	118.56	113.80
26	1H	224	G	O5'-P-OP2	-9.51	97.14	105.70
26	1H	138	G	C5-N7-C8	-9.50	99.55	104.30
26	1H	1324	G	N1-C6-O6	9.49	125.60	119.90
26	1H	783	A	N3-C4-C5	9.48	133.44	126.80
26	14	1396	U	O5'-P-OP1	-9.47	97.18	105.70
33	59	171	LEU	CA-CB-CG	9.46	137.05	115.30
26	1H	1616	A	C5-C6-N6	-9.45	116.14	123.70
26	1H	207	A	N1-C6-N6	9.44	124.27	118.60
26	14	1496	A	C5-N7-C8	-9.44	99.18	103.90
26	1H	1900	A	O5'-P-OP1	9.44	122.02	110.70
26	14	1678	G	C2-N3-C4	-9.42	107.19	111.90
26	1H	2311	A	C2-N3-C4	-9.41	105.89	110.60
26	14	1332	G	N7-C8-N9	9.41	117.80	113.10
1	13	971	G	O5'-P-OP2	-9.39	97.25	105.70
26	14	205	G	N9-C4-C5	-9.38	101.65	105.40
26	1H	518	G	O5'-P-OP2	-9.38	97.26	105.70
26	14	746	A	O5'-P-OP2	9.38	121.95	110.70
26	14	1496	A	C8-N9-C4	-9.38	102.05	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	1342	A	C2-N3-C4	-9.37	105.92	110.60
26	14	2282	G	O5'-P-OP1	-9.36	97.27	105.70
26	14	783	A	C4-C5-N7	9.36	115.38	110.70
26	1H	528	A	N3-C4-N9	-9.35	119.92	127.40
30	21	152	LYS	C-N-CA	-9.32	102.73	122.30
26	1H	2425	A	O5'-P-OP2	-9.30	97.33	105.70
26	1H	2448	A	C5-C6-N6	-9.30	116.26	123.70
26	1H	1899	G	C6-C5-N7	9.30	135.98	130.40
26	1H	2402	C	C6-N1-C2	-9.29	116.58	120.30
26	14	2490	G	N7-C8-N9	9.29	117.75	113.10
26	1H	216	A	O5'-P-OP1	-9.29	97.34	105.70
26	1H	2518	A	C5-N7-C8	-9.28	99.26	103.90
26	14	778	G	C5-C6-O6	9.27	134.16	128.60
26	14	746	A	O5'-P-OP1	-9.27	97.36	105.70
26	1H	16	G	O5'-P-OP2	-9.26	97.36	105.70
26	14	2297	C	O5'-P-OP1	-9.26	97.37	105.70
26	14	945	A	C4-C5-N7	9.24	115.32	110.70
26	1H	528	A	C2-N3-C4	-9.24	105.98	110.60
26	14	672	C	O5'-P-OP1	9.24	121.79	110.70
26	1H	1297	C	OP2-P-O3'	-9.22	84.91	105.20
26	1H	2430	A	N3-C4-N9	-9.22	120.03	127.40
26	14	2346	A	N1-C2-N3	9.20	133.90	129.30
26	1H	1784	A	O5'-P-OP1	-9.20	97.42	105.70
26	14	929	G	N1-C6-O6	9.20	125.42	119.90
26	14	1786	A	C6-C5-N7	-9.19	125.87	132.30
26	1H	137(A)	G	N1-C6-O6	9.19	125.41	119.90
26	14	1342	A	N1-C2-N3	9.19	133.89	129.30
26	1H	676	A	C4-C5-N7	9.18	115.29	110.70
26	1H	1786	A	N1-C6-N6	9.17	124.10	118.60
26	14	1332	G	C6-C5-N7	-9.17	124.90	130.40
26	1H	399	G	O5'-P-OP2	-9.16	97.45	105.70
26	14	786	C	N3-C4-N4	-9.15	111.59	118.00
1	1G	690	G	C5-N7-C8	-9.14	99.73	104.30
24	3K	76	A	N1-C6-N6	9.14	124.08	118.60
26	1H	1639	U	O5'-P-OP1	9.13	121.66	110.70
26	1H	2689	U	C5-C4-O4	-9.12	120.43	125.90
26	1H	828	U	C5-C4-O4	9.12	131.37	125.90
26	14	1249	U	O5'-P-OP1	-9.12	97.50	105.70
26	1H	1021	A	N7-C8-N9	9.09	118.34	113.80
26	1H	2503	A	C5-C6-N6	-9.09	116.43	123.70
26	1H	1022	G	C8-N9-C4	-9.08	102.77	106.40
26	1H	1950	G	C2-N3-C4	-9.07	107.36	111.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	2331	G	C2-N3-C4	-9.06	107.37	111.90
26	1H	1829	A	O5'-P-OP1	-9.06	97.55	105.70
26	1H	1394	U	O5'-P-OP2	9.06	121.57	110.70
26	1H	252	G	O5'-P-OP2	-9.04	97.56	105.70
26	1H	1931	U	C5-C4-O4	9.04	131.33	125.90
26	1H	51	G	O5'-P-OP1	-9.03	97.57	105.70
26	1H	1394	U	O5'-P-OP1	-9.03	97.58	105.70
1	13	974	A	O4'-C1'-N9	9.02	115.42	108.20
26	1H	2712	U	N3-C4-O4	-9.01	113.09	119.40
26	14	1332	G	C2-N3-C4	-9.00	107.40	111.90
31	31	74	ARG	NE-CZ-NH2	-8.99	115.80	120.30
26	1H	945	A	C8-N9-C4	-8.99	102.20	105.80
26	14	1964	G	O5'-P-OP2	-8.97	97.62	105.70
26	1H	624	C	O5'-P-OP2	8.92	121.41	110.70
26	1H	140	A	C8-N9-C4	-8.92	102.23	105.80
1	13	974	A	C6-C5-N7	-8.91	126.06	132.30
26	14	2296	U	C2-N1-C1'	8.90	128.38	117.70
26	1H	2374	C	C5-C6-N1	-8.89	116.56	121.00
26	14	205	G	C8-N9-C4	8.88	109.95	106.40
26	1H	1210	A	C8-N9-C4	-8.87	102.25	105.80
26	1H	933	A	O5'-P-OP2	-8.87	97.72	105.70
26	1H	1982	C	C6-N1-C2	-8.85	116.76	120.30
26	14	1284	A	O5'-P-OP2	-8.84	97.74	105.70
26	1H	788	A	N1-C6-N6	8.84	123.90	118.60
26	14	2713	A	N7-C8-N9	8.84	118.22	113.80
26	14	687	C	O5'-P-OP1	-8.84	97.75	105.70
26	14	2688	U	N3-C2-O2	-8.83	116.02	122.20
26	14	2430	A	N1-C2-N3	8.82	133.71	129.30
26	1H	2394	C	O5'-P-OP2	-8.80	97.78	105.70
26	1H	955	C	O5'-P-OP2	-8.79	97.79	105.70
26	14	1899	G	C2-N3-C4	-8.79	107.51	111.90
26	1H	2598	A	O5'-P-OP1	-8.77	97.81	105.70
26	1H	783	A	C8-N9-C4	-8.77	102.29	105.80
26	14	676	A	O4'-C1'-N9	8.77	115.21	108.20
26	1H	464	U	C5-C6-N1	-8.76	118.32	122.70
26	14	1899	G	C6-C5-N7	-8.75	125.15	130.40
26	1H	915	C	N1-C2-O2	8.75	124.15	118.90
26	1H	576	U	N3-C4-O4	-8.75	113.28	119.40
26	14	1021	A	C2-N3-C4	-8.75	106.23	110.60
26	14	2296	U	O5'-P-OP1	-8.74	97.83	105.70
26	1H	1899	G	C4-C5-C6	-8.74	113.56	118.80
26	14	1989	G	N3-C2-N2	-8.73	113.79	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	1566	A	N1-C6-N6	8.72	123.83	118.60
26	1H	1822	G	O5'-P-OP2	8.72	121.16	110.70
26	14	2073	C	O5'-P-OP2	8.71	121.16	110.70
26	1H	446	G	N1-C6-O6	8.71	125.12	119.90
26	1H	1931	U	N3-C2-O2	-8.70	116.11	122.20
26	1H	2427	C	O5'-P-OP2	8.69	121.13	110.70
26	1H	1806	C	O5'-P-OP2	-8.69	97.88	105.70
26	1H	829	A	OP1-P-OP2	8.67	132.60	119.60
26	1H	2346	A	O5'-P-OP1	-8.66	97.91	105.70
26	1H	1332	G	C8-N9-C4	-8.65	102.94	106.40
1	13	576	G	N1-C6-O6	8.63	125.08	119.90
26	1H	446	G	O5'-P-OP2	8.63	121.06	110.70
26	14	1842	G	N1-C6-O6	-8.63	114.72	119.90
26	14	1772	G	C8-N9-C4	8.60	109.84	106.40
26	1H	917	A	N7-C8-N9	8.59	118.09	113.80
26	1H	2441	C	N3-C2-O2	-8.59	115.89	121.90
26	14	1783	A	O5'-P-OP1	-8.59	97.97	105.70
1	13	690	G	C8-N9-C1'	-8.59	115.84	127.00
26	1H	1210	A	N7-C8-N9	8.59	118.09	113.80
26	1H	1361	G	N1-C6-O6	-8.58	114.75	119.90
26	1H	2598	A	O5'-P-OP2	8.58	120.99	110.70
26	14	1616	A	C2-N3-C4	-8.57	106.31	110.60
26	14	1265	A	N1-C6-N6	8.56	123.74	118.60
26	1H	679	C	C5-C6-N1	-8.56	116.72	121.00
26	1H	698	C	C4-C5-C6	8.56	121.68	117.40
26	14	774	A	C4-C5-N7	8.55	114.98	110.70
26	14	2360	A	O5'-P-OP2	-8.55	98.00	105.70
26	1H	251	A	O5'-P-OP1	-8.55	98.01	105.70
26	1H	137(A)	G	C5-C6-O6	-8.54	123.48	128.60
26	1H	1528	A	N7-C8-N9	8.52	118.06	113.80
1	1G	690	G	N7-C8-N9	8.52	117.36	113.10
26	1H	1204	A	C5-C6-N1	-8.52	113.44	117.70
26	1H	2623	G	C8-N9-C4	-8.52	102.99	106.40
1	13	963	G	N1-C2-N2	-8.51	108.54	116.20
26	1H	508	G	C8-N9-C4	-8.51	103.00	106.40
26	1H	1950	G	N3-C4-N9	-8.51	120.89	126.00
26	14	1975	G	O5'-P-OP2	-8.51	98.04	105.70
26	1H	945	A	C8-N9-C1'	-8.51	112.39	127.70
26	1H	71	A	N9-C4-C5	-8.50	102.40	105.80
26	1H	120	U	C5-C6-N1	-8.50	118.45	122.70
26	1H	1784	A	O5'-P-OP2	8.50	120.90	110.70
26	1H	917	A	C6-C5-N7	-8.49	126.36	132.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	1698	A	C4-C5-N7	8.49	114.94	110.70
26	14	1963	U	C2-N1-C1'	8.49	127.89	117.70
26	14	2490	G	C5-N7-C8	-8.49	100.06	104.30
26	14	1807	G	C8-N9-C4	8.48	109.79	106.40
26	1H	740	U	O5'-P-OP2	-8.48	98.07	105.70
26	1H	774	A	C8-N9-C1'	8.47	142.95	127.70
1	13	328	C	C2-N1-C1'	8.47	128.12	118.80
26	1H	2392	A	C4-C5-N7	8.46	114.93	110.70
26	14	2429	G	O5'-P-OP1	8.45	120.84	110.70
26	1H	1528	A	C8-N9-C4	-8.45	102.42	105.80
26	1H	2572	A	C8-N9-C4	8.45	109.18	105.80
26	1H	1676	A	C2-N3-C4	-8.45	106.38	110.60
26	1H	1950	G	O4'-C1'-N9	8.45	114.96	108.20
26	1H	1752	C	C6-N1-C2	8.44	123.67	120.30
26	14	1963	U	N1-C2-O2	8.44	128.71	122.80
26	1H	1203	G	O5'-P-OP2	-8.43	98.11	105.70
26	1H	2311	A	N1-C2-N3	8.43	133.51	129.30
26	14	1785	A	OP2-P-O3'	8.43	123.73	105.20
26	14	74	A	C5-C6-N1	-8.42	113.49	117.70
26	1H	1404	C	O5'-P-OP2	-8.40	98.14	105.70
1	13	5	U	N1-C2-O2	8.40	128.68	122.80
26	14	1948	G	O5'-P-OP1	-8.40	98.14	105.70
26	1H	1310	G	N1-C6-O6	8.39	124.93	119.90
26	14	2712	U	C5-C6-N1	-8.39	118.51	122.70
26	1H	778	G	N1-C6-O6	-8.38	114.87	119.90
26	1H	1899	G	C4-N9-C1'	-8.38	115.61	126.50
26	1H	774	A	C6-N1-C2	8.38	123.62	118.60
1	1G	690	G	O4'-C1'-N9	8.37	114.90	108.20
26	1H	124	G	C5-C6-O6	-8.37	123.58	128.60
1	13	1504	G	O5'-P-OP1	-8.36	98.17	105.70
24	3K	76	A	C4-C5-N7	8.35	114.88	110.70
26	1H	125	G	N1-C6-O6	8.34	124.90	119.90
26	14	929	G	C5-C6-O6	-8.34	123.60	128.60
26	14	34	C	N1-C2-O2	8.33	123.90	118.90
26	1H	786	C	C5-C4-N4	8.33	126.03	120.20
26	1H	1379	A	N1-C6-N6	8.32	123.59	118.60
1	13	5	U	N3-C2-O2	-8.32	116.38	122.20
26	14	1614	A	C2-N3-C4	-8.32	106.44	110.60
26	1H	1950	G	N3-C4-C5	8.31	132.76	128.60
26	1H	2346	A	C8-N9-C4	-8.30	102.48	105.80
1	13	980	C	O5'-P-OP1	-8.30	98.23	105.70
26	14	2058	A	O5'-P-OP2	-8.30	98.23	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	530	G	C6-C5-N7	-8.29	125.42	130.40
26	1H	2360	A	O5'-P-OP2	-8.29	98.24	105.70
26	14	2023	G	O5'-P-OP2	-8.29	98.24	105.70
26	1H	966	G	N1-C6-O6	-8.28	114.93	119.90
26	14	676	A	C4-C5-N7	8.29	114.84	110.70
26	1H	783	A	N3-C4-N9	-8.28	120.78	127.40
26	1H	2506	U	N1-C2-O2	8.28	128.60	122.80
26	14	74	A	N1-C6-N6	8.28	123.57	118.60
26	1H	1496	A	C8-N9-C4	-8.28	102.49	105.80
26	1H	2713	A	C8-N9-C4	-8.28	102.49	105.80
26	1H	731	C	O5'-P-OP2	-8.27	98.25	105.70
26	14	991	C	O5'-P-OP1	-8.27	98.26	105.70
26	1H	2380	C	C5-C6-N1	-8.27	116.87	121.00
26	14	2435	A	C8-N9-C4	-8.27	102.49	105.80
1	13	770	C	O5'-P-OP2	8.25	120.60	110.70
26	1H	821	A	OP1-P-OP2	8.25	131.98	119.60
26	1H	140	A	C6-C5-N7	-8.25	126.53	132.30
26	1H	1365	A	C5-C6-N1	-8.25	113.58	117.70
26	1H	766	C	C2-N3-C4	-8.25	115.78	119.90
26	1H	1888	G	N3-C4-N9	8.25	130.95	126.00
27	16	115	G	C5-C6-O6	-8.24	123.66	128.60
1	13	690	G	N7-C8-N9	8.23	117.22	113.10
26	1H	1781	C	C6-N1-C2	8.23	123.59	120.30
26	1H	2331	G	C4-C5-N7	8.22	114.09	110.80
26	14	621	A	N7-C8-N9	8.22	117.91	113.80
26	1H	2688	U	N1-C2-N3	8.22	119.83	114.90
26	14	2685	G	C5-C6-N1	-8.21	107.39	111.50
26	14	1616	A	N7-C8-N9	8.21	117.90	113.80
26	1H	733	G	O5'-P-OP2	-8.21	98.31	105.70
26	1H	74	A	C5-C6-N1	-8.20	113.60	117.70
26	1H	2503	A	C2-N3-C4	8.20	114.70	110.60
26	1H	751	A	O5'-P-OP1	-8.19	98.33	105.70
29	11	56	GLY	C-N-CA	-8.19	105.11	122.30
26	1H	2244	U	N3-C2-O2	-8.19	116.47	122.20
26	1H	2503	A	N1-C6-N6	8.19	123.51	118.60
26	1H	120	U	C4-C5-C6	8.18	124.61	119.70
26	1H	1301	A	N1-C6-N6	8.18	123.51	118.60
26	1H	467	G	O5'-P-OP2	-8.17	98.34	105.70
26	1H	2085	C	O5'-P-OP2	-8.17	98.35	105.70
26	1H	1271	G	O5'-P-OP2	-8.16	98.35	105.70
26	14	2326	C	C6-N1-C2	-8.16	117.03	120.30
26	1H	451	C	N1-C2-O2	-8.15	114.01	118.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	1265	A	O5'-P-OP2	-8.14	98.37	105.70
26	1H	2699	C	C6-N1-C2	8.14	123.56	120.30
26	1H	825	C	N3-C4-N4	8.14	123.70	118.00
1	1G	1281	U	N3-C2-O2	-8.14	116.50	122.20
26	1H	2518	A	C4-C5-N7	8.13	114.77	110.70
26	1H	1616	A	O4'-C1'-N9	8.13	114.70	108.20
26	1H	2380	C	C2-N3-C4	-8.13	115.84	119.90
26	1H	915	C	N3-C2-O2	-8.12	116.21	121.90
26	1H	2424	C	OP1-P-OP2	8.12	131.78	119.60
26	14	1786	A	C8-N9-C4	-8.12	102.55	105.80
26	1H	2623	G	N3-C4-C5	-8.12	124.54	128.60
26	1H	1602	U	N3-C4-O4	-8.12	113.72	119.40
23	2K	61	U	O5'-P-OP2	-8.11	98.40	105.70
26	1H	812	C	N1-C2-O2	-8.11	114.03	118.90
26	1H	205	G	C8-N9-C4	8.11	109.64	106.40
26	1H	2258	C	C4-C5-C6	8.10	121.45	117.40
26	14	774	A	C5-N7-C8	-8.10	99.85	103.90
26	1H	621	A	N3-C4-C5	8.10	132.47	126.80
26	1H	778	G	C5-C6-O6	8.10	133.46	128.60
26	14	2067	G	O5'-P-OP1	-8.10	98.41	105.70
26	14	1388	G	O5'-P-OP2	-8.09	98.42	105.70
26	14	1314	C	N1-C2-O2	8.09	123.75	118.90
26	14	1786	A	C5-C6-N1	-8.09	113.66	117.70
27	16	6	C	N1-C2-O2	-8.09	114.05	118.90
26	14	120	U	O5'-P-OP2	-8.08	98.43	105.70
26	1H	2392	A	N7-C8-N9	8.08	117.84	113.80
46	G8	81	LYS	C-N-CD	-8.08	102.83	120.60
26	14	2452	C	O5'-P-OP2	-8.07	98.44	105.70
1	13	422	C	C6-N1-C2	-8.07	117.07	120.30
1	1G	117	G	N1-C6-O6	8.06	124.74	119.90
26	14	2430	A	C5-C6-N1	-8.06	113.67	117.70
26	1H	2346	A	N7-C8-N9	8.06	117.83	113.80
1	13	1260	C	C6-N1-C2	-8.05	117.08	120.30
26	1H	930	U	N3-C4-O4	-8.05	113.76	119.40
26	1H	2402	C	N3-C2-O2	-8.05	116.26	121.90
26	14	140	A	C4-C5-N7	8.05	114.73	110.70
26	1H	2490	G	N3-C4-C5	8.05	132.63	128.60
26	1H	2446	G	C5-C6-O6	-8.05	123.77	128.60
26	1H	621	A	C6-C5-N7	-8.05	126.67	132.30
26	1H	2712	U	N3-C2-O2	-8.05	116.57	122.20
26	14	1602	U	O5'-P-OP1	-8.04	98.46	105.70
26	14	1785	A	C8-N9-C4	-8.04	102.58	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	2272	U	O5'-P-OP1	8.04	120.34	110.70
26	1H	917	A	O5'-P-OP1	-8.03	98.47	105.70
1	1G	690	G	C2-N3-C4	-8.04	107.88	111.90
26	1H	1899	G	C5-C6-O6	8.03	133.41	128.60
1	13	974	A	C4-C5-N7	8.02	114.71	110.70
23	2K	6	G	C8-N9-C4	8.02	109.61	106.40
26	1H	184	C	C6-N1-C2	8.01	123.50	120.30
26	1H	2495	G	N1-C6-O6	8.01	124.70	119.90
26	14	1396	U	N3-C2-O2	-8.01	116.60	122.20
26	14	2258	C	C4-C5-C6	8.00	121.40	117.40
26	14	1698	A	N1-C6-N6	8.00	123.40	118.60
26	1H	71	A	O4'-C1'-N9	-7.99	101.81	108.20
26	14	2609	U	O5'-P-OP2	-7.98	98.52	105.70
1	1G	1128	C	C6-N1-C2	-7.96	117.11	120.30
26	14	2566	A	O5'-P-OP2	-7.96	98.54	105.70
26	14	252	G	O5'-P-OP2	-7.96	98.54	105.70
26	14	945	A	N7-C8-N9	7.96	117.78	113.80
1	13	800	G	N1-C6-O6	7.96	124.67	119.90
26	14	1950	G	N7-C8-N9	7.96	117.08	113.10
26	1H	1660	C	N3-C2-O2	-7.95	116.33	121.90
26	1H	2715	C	N3-C4-C5	7.95	125.08	121.90
26	1H	560	C	O5'-P-OP1	-7.95	98.54	105.70
26	1H	1764	G	C5-C6-O6	7.95	133.37	128.60
26	14	783	A	N7-C8-N9	7.95	117.77	113.80
26	1H	2331	G	C6-C5-N7	-7.94	125.64	130.40
26	1H	1408	C	N1-C2-O2	-7.94	114.14	118.90
26	1H	2346	A	C6-C5-N7	-7.94	126.75	132.30
26	1H	2346	A	C4-N9-C1'	7.94	140.59	126.30
26	1H	837	C	C6-N1-C2	-7.93	117.13	120.30
26	1H	2741	A	C8-N9-C4	7.93	108.97	105.80
26	1H	856	C	O5'-P-OP1	-7.93	98.56	105.70
26	1H	755	C	C6-N1-C2	-7.93	117.13	120.30
26	1H	1835	G	O5'-P-OP1	-7.93	98.57	105.70
26	1H	1624	G	C5-C6-O6	7.92	133.35	128.60
26	1H	1626	G	N3-C2-N2	-7.92	114.36	119.90
26	14	2241	A	O5'-P-OP2	7.92	120.20	110.70
26	14	37	C	O5'-P-OP2	-7.91	98.58	105.70
26	1H	34	C	O5'-P-OP2	7.91	120.19	110.70
26	1H	210	C	N3-C4-C5	7.91	125.06	121.90
26	1H	676	A	O4'-C1'-N9	7.91	114.53	108.20
26	14	1807	G	N1-C6-O6	7.90	124.64	119.90
26	1H	1698	A	C5-C6-N1	-7.90	113.75	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	2436	G	N1-C6-O6	7.90	124.64	119.90
26	14	71	A	N7-C8-N9	7.90	117.75	113.80
26	1H	2235	G	C5-C6-O6	-7.90	123.86	128.60
26	1H	730	C	OP2-P-O3'	7.90	122.57	105.20
26	1H	132	G	N1-C6-O6	-7.89	115.16	119.90
26	1H	576	U	C4-C5-C6	7.89	124.44	119.70
26	14	1382	G	C4-C5-N7	7.89	113.96	110.80
26	14	2062	A	C8-N9-C4	7.89	108.96	105.80
26	14	1185	C	O5'-P-OP2	-7.88	98.60	105.70
26	1H	446	G	N9-C4-C5	-7.88	102.25	105.40
26	14	620	G	O5'-P-OP2	-7.88	98.61	105.70
26	1H	739	G	O5'-P-OP2	-7.88	98.61	105.70
26	1H	410	G	C5-C6-O6	-7.87	123.88	128.60
26	1H	2253	G	C5-C6-O6	-7.87	123.88	128.60
26	1H	2346	A	C4-C5-C6	7.87	120.93	117.00
26	14	1332	G	C4-C5-N7	7.86	113.94	110.80
26	14	1904	G	O5'-P-OP2	-7.86	98.63	105.70
26	14	917	A	O5'-P-OP1	-7.86	98.63	105.70
26	1H	192	C	N3-C4-C5	7.85	125.04	121.90
26	1H	265	A	C2-N3-C4	-7.85	106.67	110.60
26	1H	1253	A	C8-N9-C4	7.85	108.94	105.80
26	14	2424	C	O5'-P-OP1	-7.85	98.64	105.70
26	14	985	C	OP2-P-O3'	7.84	122.45	105.20
26	1H	2006	C	C6-N1-C2	7.83	123.43	120.30
26	1H	1255	U	C5-C4-O4	-7.83	121.20	125.90
26	1H	2430	A	N1-C2-N3	7.83	133.21	129.30
26	1H	797	C	C4-C5-C6	7.82	121.31	117.40
26	1H	2525	G	N9-C4-C5	-7.82	102.27	105.40
26	1H	2258	C	C2-N3-C4	-7.82	115.99	119.90
26	1H	1203	G	C5-C6-O6	7.82	133.29	128.60
26	1H	2622	C	O5'-P-OP2	-7.81	98.67	105.70
26	14	676	A	N7-C8-N9	7.81	117.70	113.80
26	14	2392	A	C8-N9-C4	-7.81	102.68	105.80
26	1H	140	A	C2-N3-C4	-7.81	106.70	110.60
26	1H	192	C	N1-C2-O2	-7.81	114.22	118.90
26	1H	1382	G	C4-C5-N7	7.80	113.92	110.80
26	1H	821	A	O5'-P-OP2	-7.80	98.68	105.70
26	1H	2490	G	C2-N3-C4	-7.80	108.00	111.90
26	1H	686	G	N9-C4-C5	-7.79	102.28	105.40
26	14	801	G	O5'-P-OP2	-7.79	98.69	105.70
26	1H	1324	G	N3-C2-N2	-7.79	114.45	119.90
26	14	1780	A	C8-N9-C4	-7.79	102.69	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	13	1369	C	O5'-P-OP2	-7.78	98.70	105.70
26	14	1812	A	O5'-P-OP2	-7.78	98.70	105.70
26	14	2776	A	C8-N9-C4	-7.77	102.69	105.80
26	14	71	A	C4-C5-N7	7.77	114.58	110.70
26	1H	2331	G	C5-C6-O6	-7.76	123.94	128.60
26	14	2324	C	N3-C4-C5	7.76	125.01	121.90
26	1H	1604	C	N1-C2-O2	-7.76	114.24	118.90
26	1H	1610	A	C4-C5-N7	7.76	114.58	110.70
26	14	569	U	C5-C6-N1	-7.75	118.82	122.70
26	1H	1376	C	C6-N1-C2	-7.75	117.20	120.30
26	14	1585	C	N1-C2-O2	7.75	123.55	118.90
26	1H	1970	A	O5'-P-OP1	-7.75	98.73	105.70
26	14	1786	A	N1-C6-N6	7.74	123.25	118.60
26	1H	1352	U	O5'-P-OP2	-7.74	98.73	105.70
26	14	1970	A	O5'-P-OP1	7.74	119.99	110.70
26	1H	2751	G	C8-N9-C4	7.74	109.50	106.40
26	14	1950	G	C5-N7-C8	-7.74	100.43	104.30
26	14	2238	G	O4'-C1'-N9	-7.73	102.01	108.20
26	14	2446	G	N1-C6-O6	-7.73	115.26	119.90
1	13	1446	A	O4'-C1'-N9	7.72	114.38	108.20
26	1H	2318	G	O4'-C1'-N9	7.72	114.38	108.20
26	14	2249	U	C6-N1-C2	-7.72	116.37	121.00
26	1H	852	G	O5'-P-OP2	-7.71	98.76	105.70
27	1J	30	C	C6-N1-C2	-7.71	117.22	120.30
26	1H	1357	U	C4-C5-C6	7.71	124.32	119.70
26	1H	256	A	N1-C6-N6	7.70	123.22	118.60
26	14	49	A	P-O3'-C3'	7.70	128.94	119.70
26	1H	2821	A	N1-C6-N6	7.70	123.22	118.60
26	14	2073	C	OP1-P-OP2	-7.70	108.05	119.60
26	1H	2518	A	C6-C5-N7	-7.70	126.91	132.30
26	14	1302	A	OP1-P-OP2	7.70	131.14	119.60
26	14	74	A	N3-C4-N9	-7.69	121.25	127.40
26	14	2518	A	C6-C5-N7	-7.69	126.92	132.30
26	1H	1763	G	O5'-P-OP2	-7.68	98.78	105.70
26	14	1776	G	N3-C4-N9	7.68	130.61	126.00
27	16	30	C	C6-N1-C2	-7.68	117.23	120.30
26	1H	747	U	O5'-P-OP1	-7.67	98.79	105.70
26	1H	1332	G	N1-C6-O6	7.67	124.50	119.90
26	1H	113	G	N3-C4-N9	-7.67	121.40	126.00
27	16	81	G	C4-C5-N7	7.67	113.87	110.80
1	1G	690	G	N3-C4-C5	7.67	132.43	128.60
26	1H	1778	U	O5'-P-OP1	-7.66	98.80	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	795	C	C5-C6-N1	-7.66	117.17	121.00
26	1H	797	C	C5-C6-N1	-7.66	117.17	121.00
26	1H	512	G	O4'-C1'-N9	7.66	114.33	108.20
26	14	2324	C	C6-N1-C2	7.65	123.36	120.30
26	14	1142	U	C2-N1-C1'	7.65	126.88	117.70
26	1H	841	A	O5'-P-OP2	-7.65	98.81	105.70
26	1H	1437	C	N3-C2-O2	-7.65	116.55	121.90
26	14	2436	G	C5-C6-O6	-7.64	124.02	128.60
26	1H	1349	A	O5'-P-OP1	-7.64	98.82	105.70
26	14	676	A	N3-C4-C5	7.64	132.15	126.80
26	1H	869	G	N1-C6-O6	-7.64	115.32	119.90
26	1H	848	G	O5'-P-OP2	-7.64	98.83	105.70
26	14	1162	G	O5'-P-OP1	-7.63	98.83	105.70
26	1H	508	G	N7-C8-N9	7.63	116.92	113.10
26	14	530	G	C5-N7-C8	-7.63	100.48	104.30
26	1H	330	A	C2-N3-C4	-7.63	106.78	110.60
26	1H	774	A	N1-C6-N6	7.63	123.18	118.60
1	13	330	C	N3-C2-O2	-7.63	116.56	121.90
26	14	929	G	C6-C5-N7	-7.62	125.83	130.40
26	14	752	A	P-O3'-C3'	7.62	128.84	119.70
26	1H	1673	U	C5-C6-N1	-7.62	118.89	122.70
26	1H	138	G	N7-C8-N9	7.62	116.91	113.10
1	1G	1286	A	N7-C8-N9	7.62	117.61	113.80
26	14	1616	A	N1-C6-N6	7.62	123.17	118.60
26	14	2066	C	OP1-P-O3'	7.61	121.94	105.20
26	1H	2837	G	C5-N7-C8	-7.61	100.50	104.30
26	14	2490	G	C8-N9-C4	-7.61	103.36	106.40
26	14	1762	A	C5-C6-N1	-7.61	113.90	117.70
26	14	2079	U	O5'-P-OP1	-7.60	98.86	105.70
26	14	2565	A	O5'-P-OP2	7.60	119.82	110.70
26	1H	1314	C	C2-N1-C1'	7.60	127.16	118.80
26	1H	966	G	C5-C6-O6	7.60	133.16	128.60
1	1G	449	C	C6-N1-C2	-7.60	117.26	120.30
26	14	2873	A	C4-C5-N7	7.60	114.50	110.70
27	1J	89	G	O5'-P-OP2	-7.60	98.86	105.70
26	1H	130	C	C5-C4-N4	-7.58	114.89	120.20
26	1H	410	G	N1-C6-O6	7.58	124.45	119.90
26	14	2401	U	C5-C6-N1	7.58	126.49	122.70
26	14	1303	G	C5-C6-O6	7.58	133.15	128.60
26	1H	36	G	O5'-P-OP2	-7.57	98.89	105.70
26	14	1826	G	C4-C5-N7	-7.57	107.77	110.80
26	1H	774	A	C4-N9-C1'	-7.57	112.68	126.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	1998	G	C8-N9-C4	7.57	109.43	106.40
26	1H	2390	U	O5'-P-OP1	-7.56	98.89	105.70
26	1H	1602	U	O5'-P-OP2	7.56	119.77	110.70
26	14	1786	A	N1-C2-N3	7.56	133.08	129.30
24	3K	76	A	C6-C5-N7	-7.55	127.01	132.30
26	1H	2498	C	N3-C4-C5	7.54	124.92	121.90
26	14	71	A	C2-N3-C4	-7.54	106.83	110.60
26	14	1559	G	C6-C5-N7	-7.54	125.88	130.40
1	13	802	A	N1-C6-N6	7.54	123.12	118.60
26	14	1616	A	O4'-C1'-N9	7.53	114.23	108.20
26	14	788	A	C6-C5-N7	-7.53	127.03	132.30
1	13	328	C	N1-C2-O2	7.53	123.42	118.90
26	1H	951	C	N3-C4-N4	-7.53	112.73	118.00
26	1H	2275	C	OP1-P-O3'	7.53	121.76	105.20
26	1H	2287	A	C5-N7-C8	-7.52	100.14	103.90
26	1H	1698	A	O4'-C1'-N9	7.52	114.22	108.20
27	1J	6	C	C6-N1-C2	7.52	123.31	120.30
1	13	509	A	P-O3'-C3'	7.51	128.72	119.70
1	13	853	G	N1-C6-O6	7.51	124.41	119.90
26	14	2287	A	N3-C4-C5	7.51	132.06	126.80
1	13	1279	A	N7-C8-N9	7.50	117.55	113.80
26	14	1304	C	N3-C4-N4	-7.50	112.75	118.00
1	13	108	G	C4-C5-N7	7.50	113.80	110.80
22	1K	76	A	O4'-C1'-N9	7.50	114.20	108.20
26	1H	371	A	O5'-P-OP2	-7.49	98.96	105.70
26	1H	1695	G	OP1-P-OP2	7.49	130.83	119.60
26	1H	451	C	N3-C2-O2	7.49	127.14	121.90
26	14	1993	U	O5'-P-OP1	-7.49	98.96	105.70
26	14	252	G	O5'-P-OP1	7.48	119.68	110.70
26	14	2598	A	C8-N9-C4	7.48	108.79	105.80
26	1H	140	A	O4'-C1'-N9	7.48	114.18	108.20
24	3L	76	A	O4'-C1'-N9	7.47	114.18	108.20
26	14	2838	G	N1-C6-O6	7.47	124.38	119.90
26	1H	929	G	C6-C5-N7	-7.47	125.92	130.40
26	1H	2298	A	O5'-P-OP2	-7.47	98.98	105.70
27	16	81	G	C5-N7-C8	-7.47	100.56	104.30
26	1H	1430	C	OP1-P-O3'	7.46	121.62	105.20
26	14	71	A	N1-C6-N6	7.46	123.08	118.60
26	14	1762	A	C2-N3-C4	-7.46	106.87	110.60
26	1H	451	C	O5'-P-OP2	-7.46	98.99	105.70
26	1H	621	A	N1-C2-N3	7.46	133.03	129.30
26	1H	1142(A)	A	N3-C4-C5	7.46	132.02	126.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	2211	G	C4-N9-C1'	7.46	136.20	126.50
26	1H	2713	A	C2-N3-C4	-7.46	106.87	110.60
26	14	613	U	C5-C4-O4	7.46	130.37	125.90
26	14	140	A	N1-C6-N6	7.45	123.07	118.60
26	14	2873	A	C8-N9-C4	-7.45	102.82	105.80
26	1H	1678	G	C5-C6-N1	-7.45	107.78	111.50
24	3L	5	C	C6-N1-C2	-7.45	117.32	120.30
26	1H	2507	C	N3-C2-O2	-7.44	116.69	121.90
26	1H	226	G	O4'-C1'-N9	7.44	114.15	108.20
26	1H	2689	U	C5-C6-N1	7.44	126.42	122.70
26	14	2244	U	N1-C2-N3	7.43	119.36	114.90
26	14	2591	C	N1-C2-O2	-7.43	114.44	118.90
26	14	2490	G	C4-C5-N7	7.43	113.77	110.80
26	14	2873	A	C5-C6-N1	-7.43	113.99	117.70
26	14	528	A	N3-C4-C5	7.43	132.00	126.80
26	1H	783	A	C5-C6-N1	-7.42	113.99	117.70
26	1H	2264	C	OP1-P-O3'	7.42	121.54	105.20
26	14	786	C	N3-C4-C5	7.42	124.87	121.90
26	1H	2439	A	O5'-P-OP2	-7.42	99.02	105.70
26	1H	1784	A	O4'-C1'-N9	-7.42	102.26	108.20
26	1H	1295	C	N1-C2-O2	-7.42	114.45	118.90
37	78	61	ARG	NE-CZ-NH1	7.42	124.01	120.30
26	1H	673	C	N3-C4-N4	7.42	123.19	118.00
26	1H	1930	G	O5'-P-OP1	-7.42	99.03	105.70
26	1H	1544	C	N1-C2-O2	7.41	123.34	118.90
26	1H	195	A	P-O3'-C3'	7.40	128.58	119.70
26	1H	2700	C	C5-C4-N4	-7.40	115.02	120.20
26	14	2448	A	O5'-P-OP1	-7.39	99.05	105.70
26	14	1348	G	O5'-P-OP2	7.39	119.57	110.70
26	14	1899	G	N7-C8-N9	7.39	116.79	113.10
26	1H	951	C	N1-C2-O2	7.38	123.33	118.90
26	1H	1496	A	C4-C5-N7	7.38	114.39	110.70
26	1H	2441	C	N3-C4-N4	-7.38	112.83	118.00
26	1H	2387	U	OP2-P-O3'	7.38	121.44	105.20
26	14	624	C	O5'-P-OP1	-7.38	99.06	105.70
26	14	1796	U	O5'-P-OP2	7.38	119.56	110.70
26	1H	1790	C	N3-C4-C5	7.38	124.85	121.90
26	1H	2623	G	N1-C6-O6	-7.38	115.47	119.90
26	14	140	A	C8-N9-C4	-7.38	102.85	105.80
26	1H	1261	C	C2-N3-C4	-7.38	116.21	119.90
1	13	892	A	N1-C6-N6	7.37	123.02	118.60
26	14	2335	A	O4'-C1'-N9	7.37	114.09	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	2688	U	C5-C4-O4	7.37	130.32	125.90
1	1G	909	A	N1-C6-N6	7.36	123.02	118.60
26	14	330	A	N1-C6-N6	7.36	123.02	118.60
26	14	621	A	C8-N9-C4	-7.36	102.86	105.80
26	14	102	G	O4'-C1'-N9	7.35	114.08	108.20
26	14	205	G	OP1-P-OP2	7.35	130.62	119.60
26	14	707	G	C5-C6-N1	-7.34	107.83	111.50
26	1H	1681	G	N3-C4-C5	7.34	132.27	128.60
26	1H	1698	A	N7-C8-N9	7.34	117.47	113.80
26	14	569	U	C2-N3-C4	-7.34	122.60	127.00
26	1H	1321	A	C8-N9-C4	7.33	108.73	105.80
26	1H	1559	G	N1-C6-O6	7.33	124.30	119.90
27	16	13	A	OP1-P-OP2	7.33	130.59	119.60
26	1H	945	A	C5-C6-N1	-7.32	114.04	117.70
26	1H	1239	G	C8-N9-C4	7.32	109.33	106.40
26	14	463	G	O5'-P-OP2	-7.32	99.11	105.70
26	1H	2040	C	C6-N1-C2	7.32	123.23	120.30
1	1G	690	G	N3-C4-N9	-7.32	121.61	126.00
26	14	2335	A	N1-C6-N6	-7.32	114.21	118.60
1	13	562	C	O5'-P-OP2	-7.31	99.12	105.70
26	14	1307	A	C2-N3-C4	-7.31	106.94	110.60
1	13	896	C	C5-C6-N1	-7.31	117.35	121.00
1	1G	1502	A	O5'-P-OP2	-7.31	99.12	105.70
26	14	774	A	N3-C4-N9	-7.30	121.56	127.40
26	14	1777	U	OP2-P-O3'	7.30	121.27	105.20
26	1H	733	G	N1-C6-O6	7.30	124.28	119.90
26	1H	796	C	N3-C4-N4	-7.30	112.89	118.00
26	14	1341	U	O5'-P-OP1	-7.30	99.13	105.70
22	1K	76	A	C8-N9-C4	-7.29	102.88	105.80
26	14	1253	A	C8-N9-C4	7.29	108.72	105.80
26	1H	508	G	C4-N9-C1'	7.29	135.97	126.50
26	1H	839	U	O5'-P-OP2	-7.29	99.14	105.70
26	14	1786	A	N9-C1'-C2'	7.29	123.47	114.00
26	1H	324	A	O5'-P-OP1	-7.28	99.15	105.70
1	1G	1286	A	C8-N9-C4	-7.28	102.89	105.80
26	1H	2446	G	C4-C5-N7	7.28	113.71	110.80
26	1H	797	C	N1-C2-O2	-7.28	114.53	118.90
26	1H	864	G	C2-N3-C4	7.27	115.54	111.90
26	1H	1789	A	O5'-P-OP2	-7.26	99.16	105.70
27	16	85	G	O5'-P-OP2	-7.26	99.16	105.70
26	14	2009	G	O5'-P-OP2	-7.26	99.16	105.70
49	J8	95	LEU	CA-CB-CG	7.26	132.00	115.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	1826	G	C5-N7-C8	7.26	107.93	104.30
26	1H	838	C	N1-C2-O2	-7.26	114.55	118.90
26	14	801	G	N1-C6-O6	-7.26	115.55	119.90
26	14	2032	G	N7-C8-N9	-7.25	109.47	113.10
26	1H	462	C	O5'-P-OP2	-7.25	99.18	105.70
26	1H	576	U	OP2-P-O3'	7.25	121.14	105.20
26	1H	594	U	C5-C6-N1	-7.25	119.08	122.70
26	1H	860	U	N3-C2-O2	-7.25	117.13	122.20
26	14	2352	A	O5'-P-OP1	-7.24	99.18	105.70
26	1H	141	A	C4-C5-N7	7.24	114.32	110.70
26	1H	2513	G	C8-N9-C4	-7.24	103.50	106.40
26	14	1566	A	N9-C4-C5	-7.24	102.90	105.80
26	14	2452	C	OP1-P-OP2	7.24	130.46	119.60
26	1H	1021	A	N1-C2-N3	7.24	132.92	129.30
26	14	613	U	N3-C2-O2	-7.24	117.14	122.20
26	1H	844	C	N1-C2-O2	-7.23	114.56	118.90
26	1H	2700	C	N3-C4-C5	7.23	124.79	121.90
26	14	2210	G	C4-N9-C1'	7.23	135.90	126.50
22	1K	76	A	N7-C8-N9	7.23	117.42	113.80
26	14	797	C	N1-C2-O2	-7.23	114.56	118.90
26	1H	2277	G	C4-C5-N7	-7.22	107.91	110.80
26	14	1807	G	C5-C6-O6	-7.22	124.27	128.60
26	1H	1698	A	C4-C5-N7	7.22	114.31	110.70
26	1H	2211	G	O5'-P-OP2	-7.22	99.20	105.70
26	1H	774	A	C4-C5-C6	-7.21	113.39	117.00
26	14	1902	C	N3-C4-C5	7.21	124.79	121.90
26	14	1559	G	N1-C6-O6	7.21	124.23	119.90
26	1H	1614	A	O5'-P-OP1	-7.20	99.22	105.70
26	1H	2438	U	C2-N3-C4	-7.20	122.68	127.00
26	1H	1517	G	OP1-P-O3'	7.20	121.04	105.20
26	1H	2402	C	N1-C2-O2	7.20	123.22	118.90
26	1H	189	G	C4-C5-N7	7.20	113.68	110.80
26	14	2473	U	C2-N1-C1'	7.20	126.33	117.70
26	1H	2001	A	C2-N3-C4	7.19	114.20	110.60
1	1G	519	C	C6-N1-C2	7.19	123.17	120.30
1	1G	690	G	C8-N9-C4	-7.18	103.53	106.40
26	14	1566	A	C5-C6-N6	-7.18	117.95	123.70
26	14	2873	A	C4-N9-C1'	7.18	139.23	126.30
26	1H	74	A	C5-N7-C8	-7.18	100.31	103.90
1	13	1279	A	C8-N9-C4	-7.18	102.93	105.80
26	1H	733	G	C6-C5-N7	-7.18	126.09	130.40
26	14	2392	A	C5-C6-N1	-7.17	114.11	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
24	3K	76	A	C8-N9-C4	-7.17	102.93	105.80
24	3L	76	A	N7-C8-N9	7.17	117.38	113.80
26	1H	2430	A	C5-N7-C8	-7.16	100.32	103.90
26	14	621	A	C5-N7-C8	-7.16	100.32	103.90
26	1H	794	G	N3-C4-C5	-7.16	125.02	128.60
26	1H	2232	U	C5-C4-O4	7.15	130.19	125.90
26	14	2596	U	O5'-P-OP2	-7.15	99.26	105.70
26	14	1950	G	C4-C5-N7	7.15	113.66	110.80
26	14	2498	C	C6-N1-C2	7.15	123.16	120.30
26	14	2681	C	N3-C4-N4	-7.15	113.00	118.00
26	1H	967	C	C5-C6-N1	-7.15	117.42	121.00
26	14	1313	U	C6-N1-C2	-7.15	116.71	121.00
26	1H	750	A	OP2-P-O3'	7.15	120.93	105.20
26	1H	1574	C	C6-N1-C2	7.14	123.16	120.30
26	14	783	A	N3-C4-C5	7.14	131.80	126.80
26	1H	245	G	N1-C6-O6	7.14	124.19	119.90
26	1H	1557	C	O5'-P-OP2	-7.14	99.27	105.70
26	14	330	A	N1-C2-N3	7.13	132.87	129.30
26	14	642	G	N7-C8-N9	7.13	116.67	113.10
26	1H	1324	G	C5-C6-O6	-7.13	124.32	128.60
26	1H	1616	A	C8-N9-C4	-7.13	102.95	105.80
26	14	1394	U	O5'-P-OP2	7.13	119.26	110.70
26	14	2575	C	N3-C4-C5	-7.13	119.05	121.90
26	14	783	A	N1-C2-N3	7.12	132.86	129.30
26	1H	2509	G	N1-C6-O6	-7.12	115.63	119.90
26	14	945	A	C5-C6-N1	-7.12	114.14	117.70
26	14	2330	G	N1-C6-O6	7.11	124.17	119.90
26	14	2267	A	OP1-P-OP2	7.11	130.27	119.60
26	14	783	A	C5-C6-N6	-7.11	118.01	123.70
26	1H	845	G	P-O3'-C3'	7.11	128.23	119.70
26	1H	121	G	C5-C6-N1	7.10	115.05	111.50
26	1H	2245	U	OP1-P-OP2	-7.10	108.95	119.60
26	14	2838	G	C5-C6-O6	-7.10	124.34	128.60
26	1H	728	G	C8-N9-C4	7.10	109.24	106.40
27	16	41	U	C5-C6-N1	-7.10	119.15	122.70
26	1H	2688	U	N3-C4-O4	-7.09	114.43	119.40
26	14	1965	C	N3-C4-C5	7.09	124.74	121.90
26	14	1408	C	N1-C2-O2	-7.09	114.64	118.90
26	1H	1395	A	O5'-P-OP2	-7.09	99.32	105.70
26	14	946	G	N1-C6-O6	7.09	124.16	119.90
26	14	2392	A	N7-C8-N9	7.09	117.35	113.80
26	14	808	G	N1-C2-N2	-7.09	109.82	116.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	13	523	A	N1-C6-N6	7.08	122.85	118.60
26	14	703	U	C5-C4-O4	7.08	130.15	125.90
26	1H	124	G	N1-C2-N2	7.08	122.57	116.20
26	14	2425	A	O4'-C1'-N9	7.08	113.87	108.20
26	14	74	A	C5-N7-C8	-7.08	100.36	103.90
1	13	529	G	N1-C6-O6	7.08	124.15	119.90
26	1H	2254	C	N1-C2-O2	-7.08	114.65	118.90
26	14	34	C	C2-N1-C1'	7.07	126.58	118.80
26	14	2426	A	N9-C4-C5	-7.07	102.97	105.80
1	13	1502	A	C2-N3-C4	-7.07	107.06	110.60
48	I8	81	VAL	C-N-CA	-7.07	104.03	121.70
26	1H	74	A	N7-C8-N9	7.06	117.33	113.80
26	1H	744	G	O5'-P-OP2	-7.06	99.34	105.70
26	1H	865	C	O5'-P-OP2	7.06	119.17	110.70
26	1H	2490	G	O4'-C1'-N9	7.06	113.85	108.20
1	13	974	A	C5-C6-N6	-7.06	118.05	123.70
1	13	1354	C	C6-N1-C2	-7.06	117.48	120.30
26	1H	788	A	N9-C4-C5	-7.06	102.98	105.80
26	1H	2070	G	N1-C2-N2	-7.05	109.85	116.20
26	1H	2270	G	C8-N9-C4	7.05	109.22	106.40
26	1H	2713	A	C6-C5-N7	-7.05	127.37	132.30
26	14	1907	G	O5'-P-OP1	-7.05	99.36	105.70
26	14	632	A	O5'-P-OP2	7.05	119.16	110.70
26	1H	755	C	N3-C4-C5	-7.04	119.08	121.90
1	1G	1322	C	N1-C2-O2	7.04	123.12	118.90
26	14	1315	C	N3-C4-N4	-7.04	113.07	118.00
1	13	975	A	N1-C6-N6	7.04	122.82	118.60
26	1H	860	U	C2-N1-C1'	7.04	126.15	117.70
26	1H	2318	G	N7-C8-N9	7.04	116.62	113.10
26	14	1566	A	C4-C5-N7	7.04	114.22	110.70
26	1H	1786	A	N9-C1'-C2'	7.04	123.15	114.00
27	16	115	G	C4-C5-N7	7.04	113.61	110.80
26	1H	999	U	O5'-P-OP2	7.03	119.14	110.70
26	1H	1241	A	C5-C6-N1	-7.03	114.18	117.70
26	1H	125	G	C4-C5-N7	7.03	113.61	110.80
26	14	1698	A	N7-C8-N9	7.03	117.31	113.80
26	1H	1332	G	C6-C5-N7	-7.03	126.18	130.40
26	1H	825	C	N1-C2-O2	-7.02	114.69	118.90
1	13	422	C	P-O3'-C3'	7.02	128.12	119.70
1	13	892	A	C2-N3-C4	-7.02	107.09	110.60
26	14	843	G	N1-C6-O6	7.02	124.11	119.90
26	1H	1365	A	N9-C4-C5	7.02	108.61	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	853	G	O5'-P-OP2	-7.01	99.39	105.70
26	1H	2256	G	O5'-P-OP1	7.01	119.12	110.70
26	14	2092	U	N3-C4-O4	-7.01	114.49	119.40
26	14	2270	G	C5-C6-O6	-7.01	124.39	128.60
26	1H	2550	G	C8-N9-C4	-7.01	103.60	106.40
26	14	1404	C	O5'-P-OP2	-7.01	99.39	105.70
26	14	312	G	O5'-P-OP1	-7.01	99.39	105.70
26	14	2779	U	N3-C2-O2	-7.00	117.30	122.20
26	1H	1996	C	C6-N1-C2	7.00	123.10	120.30
26	14	137	C	C6-N1-C2	-7.00	117.50	120.30
26	14	1899	G	N1-C2-N3	7.00	128.10	123.90
26	14	1653	G	O5'-P-OP2	-7.00	99.40	105.70
26	1H	1297	C	OP1-P-O3'	7.00	120.60	105.20
26	1H	2708	G	C8-N9-C4	7.00	109.20	106.40
50	K8	35	LEU	CA-CB-CG	7.00	131.39	115.30
26	1H	2712	U	N1-C2-N3	6.99	119.10	114.90
1	1G	1401	G	C4-N9-C1'	6.99	135.59	126.50
26	14	827	U	N3-C2-O2	6.99	127.09	122.20
26	14	1644	C	N3-C2-O2	-6.99	117.01	121.90
26	1H	686	G	C8-N9-C4	6.99	109.19	106.40
26	1H	738	G	C5-C6-O6	-6.99	124.41	128.60
1	1G	784	C	C6-N1-C2	6.99	123.09	120.30
26	14	2281	C	C6-N1-C2	-6.99	117.51	120.30
26	1H	1249	U	O5'-P-OP1	-6.98	99.42	105.70
26	1H	2331	G	N3-C4-C5	6.98	132.09	128.60
26	1H	2712	U	C5-C4-O4	6.98	130.09	125.90
26	1H	1352	U	O5'-P-OP1	6.98	119.07	110.70
1	1G	136	C	O5'-P-OP2	-6.98	99.42	105.70
26	1H	795	C	C2-N3-C4	-6.98	116.41	119.90
26	14	2296	U	C5-C6-N1	6.98	126.19	122.70
1	13	353	A	C8-N9-C4	-6.97	103.01	105.80
26	1H	1257	C	C4-C5-C6	6.97	120.89	117.40
26	14	1762	A	N1-C2-N3	6.97	132.78	129.30
26	14	1678	G	N3-C4-C5	6.97	132.08	128.60
26	1H	1955	U	C2-N3-C4	-6.96	122.82	127.00
26	14	802	A	C5-C6-N1	6.96	121.18	117.70
26	14	1698	A	C2-N3-C4	-6.96	107.12	110.60
26	14	2501	C	C2-N1-C1'	-6.96	111.14	118.80
26	14	2575	C	C5-C4-N4	6.96	125.07	120.20
26	1H	1817	G	N1-C6-O6	-6.96	115.72	119.90
27	16	115	G	C5-C6-N1	6.96	114.98	111.50
26	1H	124	G	N3-C4-C5	6.96	132.08	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	1559	G	N3-C4-C5	6.96	132.08	128.60
26	1H	239	U	C5-C6-N1	-6.96	119.22	122.70
33	51	153	LYS	C-N-CD	-6.96	105.30	120.60
1	1G	413	G	C4-N9-C1'	-6.96	117.46	126.50
26	1H	193	U	N1-C2-O2	-6.95	117.93	122.80
26	14	970	C	N1-C2-O2	-6.95	114.73	118.90
1	13	974	A	C5-N7-C8	-6.95	100.42	103.90
26	1H	400	G	N1-C6-O6	6.95	124.07	119.90
26	14	1824	G	C8-N9-C4	-6.95	103.62	106.40
26	1H	917	A	C5-C6-N1	-6.95	114.22	117.70
26	1H	2084	C	C2-N3-C4	-6.95	116.42	119.90
26	1H	1817	G	C5-C6-O6	6.95	132.77	128.60
1	1G	413	G	O4'-C1'-N9	6.95	113.76	108.20
27	1J	60	C	C6-N1-C2	-6.95	117.52	120.30
1	13	346	G	N7-C8-N9	6.95	116.57	113.10
26	1H	1379	A	C5-N7-C8	-6.95	100.43	103.90
1	1G	380	G	N3-C4-N9	-6.95	121.83	126.00
1	1G	950	U	O5'-P-OP2	6.95	119.04	110.70
26	14	2331	G	C5-C6-O6	-6.95	124.43	128.60
1	13	963	G	N1-C6-O6	-6.94	115.73	119.90
26	1H	207	A	C2-N3-C4	-6.94	107.13	110.60
1	1G	27	G	N1-C6-O6	6.94	124.06	119.90
1	1G	576	G	C4-N9-C1'	6.94	135.52	126.50
1	13	254	G	O5'-P-OP1	-6.94	99.46	105.70
26	1H	576	U	N1-C2-O2	6.94	127.66	122.80
26	1H	2311	A	C5-N7-C8	-6.94	100.43	103.90
1	13	585	G	O5'-P-OP2	-6.93	99.46	105.70
1	13	690	G	C4-C5-N7	6.93	113.57	110.80
26	14	2003	G	C5-C6-O6	-6.93	124.44	128.60
26	14	2443	C	O5'-P-OP2	6.93	119.02	110.70
26	14	2573	C	C2-N1-C1'	6.93	126.43	118.80
26	1H	794	G	C4-C5-N7	-6.93	108.03	110.80
26	1H	1310	G	O5'-P-OP2	6.93	119.02	110.70
26	1H	1621	U	N3-C4-O4	6.93	124.25	119.40
26	14	1332	G	N1-C6-O6	6.93	124.06	119.90
26	14	1968	G	C5-N7-C8	-6.93	100.84	104.30
1	13	843	U	C2-N1-C1'	6.93	126.01	117.70
26	1H	263	C	O5'-P-OP2	-6.93	99.46	105.70
26	1H	955	C	OP1-P-OP2	6.92	129.99	119.60
26	14	207	A	C8-N9-C4	6.92	108.57	105.80
26	1H	1574	C	OP2-P-O3'	6.92	120.43	105.20
26	1H	202	U	N3-C4-C5	6.92	118.75	114.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	860	U	C2-N3-C4	-6.92	122.85	127.00
26	1H	1610	A	C5-C6-N6	-6.92	118.17	123.70
26	1H	1827	C	C4-C5-C6	6.91	120.86	117.40
26	1H	1313	U	C6-N1-C2	-6.91	116.86	121.00
26	14	575	A	O5'-P-OP1	-6.91	99.48	105.70
26	1H	132	G	C5-C6-O6	6.91	132.74	128.60
26	1H	739	G	N1-C6-O6	6.91	124.04	119.90
26	14	528	A	N3-C4-N9	-6.91	121.88	127.40
26	1H	2503	A	N9-C4-C5	-6.90	103.04	105.80
26	14	74	A	N1-C2-N3	6.90	132.75	129.30
26	1H	1543	A	C2-N3-C4	-6.90	107.15	110.60
26	14	2685	G	N3-C2-N2	-6.90	115.07	119.90
26	1H	1021	A	C8-N9-C4	-6.89	103.04	105.80
25	4L	23	A	OP1-P-O3'	6.89	120.37	105.20
26	14	2346	A	C2-N3-C4	-6.89	107.15	110.60
26	1H	1790	C	C2-N3-C4	-6.89	116.45	119.90
26	1H	1241	A	C2-N3-C4	-6.89	107.16	110.60
26	1H	2287	A	N3-C4-N9	-6.89	121.89	127.40
26	1H	2370	G	N1-C6-O6	-6.89	115.77	119.90
26	14	2433	A	N1-C6-N6	6.89	122.73	118.60
26	1H	954	G	N3-C2-N2	-6.88	115.08	119.90
26	1H	946	G	C8-N9-C4	6.88	109.15	106.40
26	1H	2598	A	N9-C4-C5	-6.88	103.05	105.80
26	1H	2609	U	C6-N1-C2	6.88	125.13	121.00
26	14	510	C	OP1-P-OP2	6.88	129.92	119.60
26	1H	1395	A	OP1-P-OP2	6.88	129.91	119.60
26	14	2275	C	P-O3'-C3'	6.87	127.94	119.70
1	1G	547	A	C8-N9-C4	6.87	108.55	105.80
40	65	110	LEU	CA-CB-CG	6.87	131.10	115.30
1	13	49	U	P-O3'-C3'	6.87	127.94	119.70
26	1H	631	A	N7-C8-N9	-6.87	110.37	113.80
26	1H	2618	G	N9-C4-C5	6.86	108.14	105.40
26	14	856	C	C6-N1-C2	-6.86	117.55	120.30
26	14	2066	C	C5-C4-N4	-6.86	115.39	120.20
37	78	50	ARG	NE-CZ-NH2	6.86	123.73	120.30
26	14	148	C	C6-N1-C2	6.86	123.04	120.30
26	1H	148	C	C2-N3-C4	-6.86	116.47	119.90
26	14	1989	G	C5-C6-O6	-6.86	124.48	128.60
26	1H	37	C	C5-C4-N4	6.86	125.00	120.20
26	1H	1185	C	O5'-P-OP2	-6.86	99.53	105.70
26	1H	1842	G	N1-C6-O6	-6.86	115.78	119.90
26	14	2211	G	C8-N9-C1'	-6.86	118.08	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	1373	A	O5'-P-OP1	6.86	118.93	110.70
26	14	750	A	C8-N9-C4	-6.85	103.06	105.80
26	1H	792	G	O4'-C1'-N9	-6.85	102.72	108.20
1	13	703	G	C6-C5-N7	-6.85	126.29	130.40
26	1H	825	C	C5-C4-N4	-6.84	115.41	120.20
26	1H	1428	C	O5'-P-OP1	-6.84	99.54	105.70
26	14	2092	U	N3-C2-O2	-6.84	117.41	122.20
26	14	509	C	C4-C5-C6	6.84	120.82	117.40
26	1H	2713	A	N1-C6-N6	6.84	122.70	118.60
26	14	1342	A	O4'-C1'-N9	6.84	113.67	108.20
1	13	1502	A	C4-C5-N7	6.84	114.12	110.70
26	1H	508	G	N3-C4-C5	-6.84	125.18	128.60
26	1H	1325	G	C5-C6-O6	-6.84	124.50	128.60
26	1H	2586	C	C2-N3-C4	-6.84	116.48	119.90
26	14	1303	G	N1-C6-O6	-6.84	115.80	119.90
26	1H	2598	A	N1-C6-N6	6.83	122.70	118.60
26	1H	2346	A	C5-N7-C8	-6.83	100.48	103.90
26	14	1600	C	O5'-P-OP2	-6.83	99.55	105.70
26	1H	1528	A	O4'-C1'-N9	6.83	113.66	108.20
26	1H	838	C	C2-N3-C4	-6.83	116.49	119.90
26	1H	1574	C	C5-C6-N1	-6.83	117.59	121.00
26	1H	1798	U	N3-C4-C5	6.82	118.69	114.60
26	1H	2648	C	C6-N1-C2	6.82	123.03	120.30
26	1H	862	G	C5-C6-O6	6.82	132.69	128.60
1	1G	413	G	C6-C5-N7	6.82	134.49	130.40
1	1G	481	G	C6-C5-N7	-6.82	126.31	130.40
26	14	2429	G	O5'-P-OP2	-6.82	99.56	105.70
26	1H	1298	C	OP1-P-O3'	6.82	120.20	105.20
26	1H	1300	U	N1-C2-N3	6.82	118.99	114.90
26	1H	1767	C	O5'-P-OP1	-6.82	99.56	105.70
26	1H	695	G	N3-C2-N2	6.82	124.67	119.90
26	1H	1142(A)	A	C5-C6-N1	-6.81	114.29	117.70
27	16	44	G	C8-N9-C1'	6.81	135.86	127.00
26	14	1939	U	OP2-P-O3'	6.81	120.19	105.20
26	14	2032	G	C5-N7-C8	6.81	107.71	104.30
26	14	2352	A	N1-C2-N3	6.81	132.71	129.30
26	1H	576	U	C5-C6-N1	-6.81	119.30	122.70
26	14	855	G	C8-N9-C4	-6.81	103.68	106.40
26	14	2542	A	C8-N9-C4	6.81	108.52	105.80
26	1H	2593	U	N3-C2-O2	-6.80	117.44	122.20
26	1H	240	G	C5-C6-N1	6.80	114.90	111.50
26	1H	672	C	OP2-P-O3'	6.80	120.16	105.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	1022	G	N3-C2-N2	-6.80	115.14	119.90
26	1H	190	A	N1-C2-N3	-6.80	125.90	129.30
1	13	703	G	N1-C6-O6	6.80	123.98	119.90
26	1H	2507	C	N1-C2-O2	6.80	122.98	118.90
26	1H	2037	G	N1-C6-O6	-6.79	115.82	119.90
26	14	2313	C	C6-N1-C2	-6.79	117.58	120.30
1	13	1434	A	C8-N9-C4	6.79	108.52	105.80
26	14	732	C	N1-C2-O2	-6.79	114.83	118.90
1	1G	990	C	C6-N1-C2	-6.78	117.59	120.30
27	16	44	G	P-O3'-C3'	6.78	127.83	119.70
26	1H	1565	C	N3-C4-C5	6.78	124.61	121.90
26	1H	2712	U	C5-C6-N1	-6.78	119.31	122.70
26	1H	1429	G	C5-C6-O6	6.77	132.66	128.60
26	1H	733	G	C4-C5-C6	6.77	122.86	118.80
26	1H	752	A	N9-C4-C5	-6.77	103.09	105.80
26	1H	1404	C	OP1-P-OP2	6.77	129.76	119.60
26	14	694	U	O5'-P-OP1	6.77	118.82	110.70
26	1H	2235	G	N1-C6-O6	6.77	123.96	119.90
26	14	2249	U	N3-C4-C5	-6.77	110.54	114.60
1	13	449	C	C2-N1-C1'	6.77	126.24	118.80
26	14	571	A	N1-C6-N6	6.76	122.66	118.60
26	14	751	A	O5'-P-OP1	-6.76	99.61	105.70
26	1H	2441	C	N1-C2-O2	6.76	122.96	118.90
26	1H	1558	A	N1-C2-N3	6.76	132.68	129.30
46	G8	81	LYS	C-N-CA	6.76	150.37	122.00
26	1H	1598	C	OP1-P-O3'	6.75	120.06	105.20
26	1H	265	A	N7-C8-N9	6.75	117.17	113.80
26	1H	698	C	C5-C6-N1	-6.75	117.62	121.00
26	1H	1023	U	O5'-P-OP1	-6.75	99.63	105.70
26	14	1313	U	C2-N1-C1'	6.75	125.80	117.70
26	1H	1300	U	C2-N3-C4	-6.74	122.95	127.00
26	1H	2544	G	N1-C6-O6	6.74	123.94	119.90
26	14	2453	A	N1-C6-N6	6.74	122.64	118.60
26	1H	140	A	OP2-P-O3'	6.74	120.02	105.20
26	14	1440	G	O5'-P-OP2	-6.74	99.64	105.70
26	14	2688	U	C5-C6-N1	-6.74	119.33	122.70
26	1H	756	C	C6-N1-C2	-6.74	117.61	120.30
23	2K	6	G	N9-C4-C5	-6.73	102.71	105.40
26	1H	1363	C	N3-C4-N4	-6.73	113.29	118.00
26	1H	621	A	C5-C6-N1	-6.73	114.34	117.70
1	1G	266	G	P-O3'-C3'	6.73	127.78	119.70
26	1H	831	G	C8-N9-C4	6.73	109.09	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	13	798	G	N1-C6-O6	6.72	123.94	119.90
26	1H	696	G	C5-C6-N1	6.72	114.86	111.50
27	16	99	A	C8-N9-C4	-6.72	103.11	105.80
26	14	1762	A	C6-C5-N7	-6.72	127.59	132.30
26	14	2248	C	N1-C2-O2	6.72	122.93	118.90
26	1H	1621	U	N1-C2-O2	-6.72	118.09	122.80
26	14	2681	C	C5-C4-N4	6.72	124.91	120.20
37	78	15	ARG	C-N-CA	6.72	138.50	121.70
26	14	933	A	C5-N7-C8	-6.72	100.54	103.90
26	14	2510	C	C5-C6-N1	-6.72	117.64	121.00
26	14	1328	G	C5-C6-O6	-6.72	124.57	128.60
26	1H	2070	G	N3-C2-N2	6.72	124.60	119.90
26	14	1973	G	N3-C2-N2	6.72	124.60	119.90
26	1H	245	G	C6-C5-N7	-6.71	126.37	130.40
26	1H	1558	A	C2-N3-C4	-6.71	107.25	110.60
26	1H	1817	G	N3-C2-N2	6.71	124.60	119.90
26	14	1279	G	O5'-P-OP2	-6.71	99.66	105.70
26	1H	783	A	N9-C1'-C2'	-6.71	104.62	112.00
26	14	467	G	O5'-P-OP2	-6.71	99.67	105.70
26	1H	906	G	N3-C4-N9	-6.70	121.98	126.00
26	1H	1253	A	N7-C8-N9	-6.70	110.45	113.80
26	1H	2540	C	N3-C4-N4	-6.70	113.31	118.00
26	1H	1022	G	N3-C4-N9	-6.70	121.98	126.00
26	1H	593	G	O5'-P-OP2	-6.70	99.67	105.70
1	1G	1054	C	OP1-P-OP2	-6.70	109.56	119.60
26	14	1376	C	O5'-P-OP1	-6.69	99.68	105.70
26	1H	2513	G	N7-C8-N9	6.69	116.45	113.10
26	14	2296	U	C6-N1-C1'	-6.69	111.83	121.20
26	1H	2593	U	N3-C4-O4	-6.69	114.72	119.40
26	1H	2682	U	O5'-P-OP2	-6.69	99.68	105.70
27	16	115	G	C5-N7-C8	-6.69	100.95	104.30
26	14	2286	A	N7-C8-N9	6.69	117.14	113.80
26	1H	386	G	C5-C6-O6	-6.69	124.59	128.60
26	1H	1261	C	C5-C6-N1	-6.69	117.66	121.00
26	1H	2244	U	C4-C5-C6	6.69	123.71	119.70
1	13	328	C	C6-N1-C1'	-6.68	112.78	120.80
26	1H	1914	C	N3-C2-O2	-6.68	117.22	121.90
1	1G	254	G	O5'-P-OP1	-6.68	99.69	105.70
1	13	811	C	O5'-P-OP2	-6.68	99.69	105.70
1	13	1502	A	C5-N7-C8	-6.68	100.56	103.90
26	1H	144	C	C5-C6-N1	-6.68	117.66	121.00
26	1H	481	G	O5'-P-OP2	-6.67	99.69	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	2712	U	C2-N3-C4	-6.67	123.00	127.00
1	13	5	U	C2-N1-C1'	6.67	125.70	117.70
26	1H	1204	A	C5-N7-C8	-6.67	100.56	103.90
26	1H	2609	U	C2-N3-C4	-6.67	123.00	127.00
26	1H	673	C	C5-C4-N4	-6.67	115.53	120.20
26	1H	863	A	O5'-P-OP2	-6.66	99.70	105.70
26	14	2712	U	O4'-C1'-N1	6.66	113.53	108.20
26	1H	197	A	C5-N7-C8	-6.66	100.57	103.90
26	1H	2032	G	N1-C2-N3	6.66	127.89	123.90
26	14	1812	A	OP1-P-OP2	6.66	129.58	119.60
26	14	1992	G	C8-N9-C4	-6.66	103.74	106.40
1	13	990	C	C6-N1-C2	-6.65	117.64	120.30
2	1E	187	LEU	CA-CB-CG	6.65	130.60	115.30
1	13	690	G	C4-C5-C6	6.65	122.79	118.80
26	1H	696	G	C5-C6-O6	-6.65	124.61	128.60
26	1H	141	A	N7-C8-N9	6.65	117.12	113.80
26	14	1762	A	N1-C6-N6	6.65	122.59	118.60
26	1H	1382	G	C6-C5-N7	-6.64	126.41	130.40
26	1H	626	U	N1-C2-N3	6.64	118.89	114.90
26	1H	138	G	C5-C6-O6	-6.64	124.62	128.60
26	14	668	G	C8-N9-C4	6.64	109.06	106.40
26	1H	1310	G	N3-C2-N2	-6.64	115.25	119.90
26	1H	1496	A	N1-C6-N6	6.64	122.58	118.60
26	1H	1940	U	N1-C2-O2	-6.64	118.15	122.80
26	1H	1986	A	C8-N9-C4	-6.64	103.14	105.80
26	1H	2392	A	C2-N3-C4	-6.64	107.28	110.60
26	1H	1528	A	C5-N7-C8	-6.64	100.58	103.90
26	14	140	A	C6-C5-N7	-6.64	127.66	132.30
26	14	1326	U	OP2-P-O3'	6.64	119.80	105.20
1	13	1129	C	C2-N1-C1'	6.63	126.10	118.80
26	1H	1241	A	C6-N1-C2	6.63	122.58	118.60
26	1H	679	C	C4-C5-C6	6.63	120.72	117.40
26	1H	2256	G	N3-C2-N2	6.63	124.54	119.90
26	1H	809	G	C5-C6-N1	6.63	114.81	111.50
26	1H	746	A	O5'-P-OP2	6.63	118.66	110.70
26	1H	1440	G	O5'-P-OP2	-6.63	99.73	105.70
26	1H	1694	C	C6-N1-C2	6.63	122.95	120.30
26	14	249	C	O5'-P-OP1	-6.63	99.73	105.70
1	13	813	U	N3-C4-O4	-6.63	114.76	119.40
26	1H	270(O)	U	C2-N1-C1'	6.63	125.65	117.70
23	2K	9	G	N1-C6-O6	6.62	123.88	119.90
1	13	690	G	C8-N9-C4	-6.62	103.75	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	1G	1502	A	N1-C2-N3	6.62	132.61	129.30
26	1H	207	A	C5-C6-N6	-6.62	118.40	123.70
26	1H	631	A	C8-N9-C4	6.62	108.45	105.80
26	1H	2577	A	O5'-P-OP1	-6.62	99.74	105.70
49	J8	85	LEU	CA-CB-CG	6.62	130.53	115.30
1	13	966	G	C5-C6-O6	-6.62	124.63	128.60
26	1H	150	C	N3-C4-N4	-6.62	113.37	118.00
26	1H	2506	U	C2-N1-C1'	6.62	125.64	117.70
26	14	1779	U	C5-C4-O4	-6.62	121.93	125.90
33	51	105	LEU	CA-CB-CG	6.61	130.51	115.30
26	1H	847	U	C5-C6-N1	-6.61	119.39	122.70
1	13	974	A	N9-C4-C5	-6.61	103.16	105.80
26	14	2053	G	C8-N9-C4	6.61	109.04	106.40
26	1H	676	A	C8-N9-C4	-6.61	103.16	105.80
26	14	783	A	C8-N9-C4	-6.61	103.16	105.80
26	1H	1321	A	N7-C8-N9	-6.60	110.50	113.80
1	13	758	G	N1-C6-O6	6.60	123.86	119.90
26	1H	1313	U	C2-N1-C1'	6.60	125.62	117.70
26	1H	2329	G	OP1-P-OP2	6.60	129.50	119.60
26	1H	2441	C	C5-C4-N4	6.60	124.82	120.20
26	1H	530	G	N1-C6-O6	-6.60	115.94	119.90
26	1H	768	G	O5'-P-OP2	-6.60	99.76	105.70
1	13	963	G	N3-C2-N2	6.60	124.52	119.90
26	1H	773	U	C5-C6-N1	-6.60	119.40	122.70
26	1H	1572	A	C2-N3-C4	-6.60	107.30	110.60
26	1H	947	G	C8-N9-C4	-6.60	103.76	106.40
26	1H	1319	G	O5'-P-OP1	-6.60	99.76	105.70
1	1G	1502	A	C6-C5-N7	-6.60	127.68	132.30
26	14	1204	A	C5-C6-N1	-6.60	114.40	117.70
26	1H	1203	G	N1-C6-O6	-6.59	115.94	119.90
26	1H	2591	C	N3-C2-O2	6.59	126.52	121.90
1	13	687	A	P-O3'-C3'	6.59	127.61	119.70
1	1G	1473	A	C8-N9-C4	6.59	108.44	105.80
26	1H	2401	U	C5-C6-N1	6.59	126.00	122.70
26	14	584	C	C6-N1-C2	6.59	122.94	120.30
26	1H	144	C	C6-N1-C2	6.59	122.94	120.30
26	1H	74	A	C6-C5-N7	-6.59	127.69	132.30
26	1H	2346	A	C6-N1-C2	-6.59	114.65	118.60
26	1H	2490	G	C8-N9-C4	-6.59	103.77	106.40
26	14	1999	C	OP2-P-O3'	6.59	119.69	105.20
26	1H	2272	U	O5'-P-OP2	-6.58	99.77	105.70
26	1H	2249	U	N3-C4-O4	-6.58	114.79	119.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	1G	449	C	N3-C2-O2	-6.58	117.29	121.90
24	3L	76	A	C5-N7-C8	-6.58	100.61	103.90
26	14	2607	G	N9-C4-C5	-6.58	102.77	105.40
26	1H	202	U	N1-C2-N3	-6.58	110.95	114.90
26	1H	222	A	P-O3'-C3'	6.58	127.59	119.70
26	1H	599	G	N3-C4-N9	6.58	129.95	126.00
26	1H	1367	A	N1-C6-N6	6.58	122.55	118.60
26	1H	1617	C	O5'-P-OP2	6.58	118.59	110.70
26	1H	2318	G	C8-N9-C4	-6.58	103.77	106.40
30	29	78	LEU	CA-CB-CG	6.58	130.43	115.30
26	14	2700	C	C6-N1-C2	6.58	122.93	120.30
1	1G	690	G	C4-C5-N7	6.57	113.43	110.80
1	13	576	G	C6-C5-N7	-6.57	126.46	130.40
1	13	1301	U	P-O3'-C3'	6.57	127.58	119.70
1	13	1450	U	N3-C2-O2	-6.57	117.60	122.20
26	1H	85	G	O5'-P-OP1	6.57	118.58	110.70
26	14	2392	A	C5-N7-C8	-6.57	100.62	103.90
1	1G	60	A	C8-N9-C4	6.56	108.42	105.80
26	1H	1355	G	N1-C6-O6	-6.56	115.97	119.90
26	14	642	G	C8-N9-C4	-6.56	103.78	106.40
26	14	2053	G	C5-C6-O6	-6.56	124.67	128.60
26	1H	236	C	C4-C5-C6	6.56	120.68	117.40
26	14	1156	A	O5'-P-OP2	-6.55	99.80	105.70
26	14	2713	A	N1-C6-N6	6.55	122.53	118.60
1	13	827	U	C2-N1-C1'	6.55	125.56	117.70
26	1H	265	A	C8-N9-C4	-6.55	103.18	105.80
26	1H	774	A	C4-C5-N7	6.55	113.97	110.70
26	1H	809	G	C5-C6-O6	-6.55	124.67	128.60
26	14	199	A	C2-N3-C4	6.55	113.88	110.60
26	1H	728	G	N1-C6-O6	6.55	123.83	119.90
26	14	2713	A	C4-C5-N7	6.55	113.97	110.70
26	1H	834	C	N1-C2-O2	-6.54	114.97	118.90
26	14	1332	G	N1-C2-N3	6.54	127.83	123.90
26	1H	1669	A	C5-C6-N6	-6.54	118.46	123.70
26	14	1614	A	N1-C2-N3	6.54	132.57	129.30
26	1H	1939	U	C4-C5-C6	-6.54	115.78	119.70
26	14	2592	G	N3-C4-N9	6.54	129.93	126.00
26	14	1585	C	C6-N1-C2	-6.54	117.69	120.30
26	14	2239	G	N1-C2-N2	-6.54	110.31	116.20
1	13	1233	G	N1-C6-O6	-6.54	115.98	119.90
26	1H	614	U	C5-C6-N1	6.53	125.97	122.70
26	1H	400	G	C5-C6-O6	-6.53	124.68	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	491	G	O5'-P-OP1	-6.53	99.83	105.70
26	1H	728	G	N9-C4-C5	-6.53	102.79	105.40
26	1H	1617	C	O5'-P-OP1	-6.53	99.82	105.70
26	14	2067	G	C6-N1-C2	-6.53	121.18	125.10
26	14	2446	G	N1-C2-N2	-6.53	110.33	116.20
1	13	1344	C	OP1-P-O3'	6.53	119.56	105.20
26	1H	1259	G	OP2-P-O3'	6.53	119.56	105.20
26	14	2498	C	O5'-P-OP2	-6.53	99.83	105.70
26	14	2542	A	N7-C8-N9	-6.52	110.54	113.80
26	14	2248	C	N3-C2-O2	-6.52	117.33	121.90
26	1H	216	A	O5'-P-OP2	6.52	118.53	110.70
26	1H	1571	A	N1-C6-N6	6.52	122.51	118.60
26	1H	699	A	N1-C6-N6	6.52	122.51	118.60
27	16	115	G	C6-N1-C2	-6.52	121.19	125.10
26	1H	2490	G	C6-C5-N7	-6.51	126.49	130.40
1	1G	315	A	N1-C6-N6	6.51	122.51	118.60
26	14	395	U	O4'-C1'-N1	6.51	113.41	108.20
26	1H	2544	G	C5-C6-O6	-6.51	124.69	128.60
1	1G	576	G	C8-N9-C1'	-6.51	118.53	127.00
1	1G	777	A	O5'-P-OP2	-6.51	99.84	105.70
26	14	1786	A	N3-C4-C5	6.51	131.36	126.80
26	1H	683	C	N3-C4-C5	6.51	124.50	121.90
26	14	1964	G	N3-C2-N2	6.51	124.45	119.90
26	14	2057	A	N1-C6-N6	6.51	122.50	118.60
26	1H	528	A	C5-N7-C8	-6.50	100.65	103.90
26	1H	1444	G	N1-C6-O6	-6.50	116.00	119.90
26	1H	2245	U	C5-C4-O4	-6.50	122.00	125.90
26	14	1342	A	C6-C5-N7	-6.50	127.75	132.30
26	1H	1318	C	O5'-P-OP1	-6.50	99.85	105.70
26	14	774	A	C5-C6-N1	-6.50	114.45	117.70
26	1H	209	C	C2-N3-C4	-6.50	116.65	119.90
1	13	1199	U	C5-C4-O4	6.50	129.80	125.90
26	1H	528	A	C8-N9-C1'	6.50	139.40	127.70
26	1H	1786	A	C5-C6-N1	-6.50	114.45	117.70
26	14	1950	G	O4'-C1'-N9	6.50	113.40	108.20
26	14	2821	A	C2-N3-C4	-6.50	107.35	110.60
26	1H	1370	C	N1-C2-O2	-6.50	115.00	118.90
26	14	1391	U	O5'-P-OP1	-6.50	99.85	105.70
26	14	1831	G	C6-C5-N7	-6.50	126.50	130.40
1	13	33	A	O5'-P-OP2	-6.49	99.86	105.70
26	1H	2710	C	C6-N1-C2	6.49	122.90	120.30
26	14	1396	U	C2-N1-C1'	6.49	125.49	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	1591	G	N1-C6-O6	-6.49	116.01	119.90
26	1H	120	U	N3-C2-O2	-6.49	117.66	122.20
26	14	113	G	C5-C6-O6	-6.49	124.71	128.60
26	14	1852	C	C6-N1-C2	-6.49	117.71	120.30
26	14	2818	G	C8-N9-C4	6.49	109.00	106.40
26	14	2726	U	N3-C2-O2	-6.48	117.66	122.20
26	1H	1392	A	O5'-P-OP1	-6.48	99.87	105.70
26	1H	2392	A	N3-C4-C5	6.48	131.34	126.80
26	1H	1761	C	C5-C4-N4	-6.48	115.66	120.20
26	1H	1657	C	OP1-P-O3'	6.48	119.45	105.20
1	1G	503	C	C6-N1-C2	-6.48	117.71	120.30
1	13	1489	G	C8-N9-C4	6.47	108.99	106.40
26	14	1992	G	P-O3'-C3'	6.47	127.47	119.70
26	1H	1974	C	C6-N1-C2	-6.47	117.71	120.30
26	1H	130	C	C6-N1-C2	6.47	122.89	120.30
26	1H	1357	U	N3-C4-C5	-6.47	110.72	114.60
26	1H	929	G	C8-N9-C4	6.47	108.99	106.40
1	1G	337	C	C6-N1-C2	-6.47	117.71	120.30
26	14	1313	U	N3-C4-O4	6.47	123.93	119.40
26	14	1973	G	C5-C6-O6	6.47	132.48	128.60
26	1H	1202	C	N1-C2-O2	-6.47	115.02	118.90
26	1H	23	G	N3-C2-N2	-6.46	115.38	119.90
26	1H	109	G	N1-C6-O6	-6.46	116.02	119.90
26	1H	202	U	C6-N1-C2	6.46	124.88	121.00
26	1H	1306	C	C2-N3-C4	-6.46	116.67	119.90
26	14	704	G	N3-C2-N2	-6.46	115.38	119.90
14	5I	12	ARG	C-N-CA	6.46	137.84	121.70
26	1H	1327	C	N1-C2-O2	-6.46	115.03	118.90
26	1H	1630	G	O5'-P-OP1	-6.46	99.89	105.70
26	14	511	U	C6-N1-C2	-6.46	117.12	121.00
1	13	2	U	O5'-P-OP1	-6.45	99.89	105.70
26	14	646	A	C8-N9-C4	-6.45	103.22	105.80
26	1H	535	C	O5'-P-OP2	-6.45	99.89	105.70
1	1G	1305	G	N3-C2-N2	-6.45	115.38	119.90
1	1G	1401	G	N1-C6-O6	6.45	123.77	119.90
26	14	2430	A	C6-C5-N7	-6.45	127.79	132.30
26	14	2610	C	O5'-P-OP2	6.45	118.44	110.70
26	14	1781	C	C2-N1-C1'	6.45	125.89	118.80
26	1H	2713	A	N1-C2-N3	6.44	132.52	129.30
26	14	1654	A	N1-C6-N6	-6.44	114.73	118.60
26	1H	739	G	C5-C6-O6	-6.44	124.73	128.60
26	1H	1678	G	N1-C2-N3	6.44	127.76	123.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	945	A	C5-C6-N6	-6.44	118.55	123.70
26	14	1209	G	O5'-P-OP2	-6.44	99.91	105.70
26	14	2581	G	C5-C6-O6	6.44	132.46	128.60
26	1H	1969	A	OP1-P-O3'	6.44	119.36	105.20
26	1H	112	U	N3-C2-O2	6.43	126.70	122.20
26	14	2057	A	OP1-P-OP2	6.43	129.25	119.60
26	14	2226	C	N1-C2-O2	6.43	122.76	118.90
26	1H	704	G	N3-C2-N2	-6.42	115.40	119.90
26	1H	2003	G	O5'-P-OP1	-6.42	99.92	105.70
26	14	1807	G	N9-C4-C5	-6.42	102.83	105.40
1	13	1502	A	O5'-P-OP2	-6.42	99.92	105.70
26	14	2386	C	C6-N1-C2	6.42	122.87	120.30
1	13	328	C	N3-C2-O2	-6.42	117.41	121.90
1	13	584	G	C5-C6-O6	6.42	132.45	128.60
27	16	98	G	C4-C5-N7	6.42	113.37	110.80
26	1H	754	C	C2-N3-C4	-6.41	116.69	119.90
26	14	1936	A	O4'-C1'-N9	6.41	113.33	108.20
26	1H	788	A	OP2-P-O3'	6.41	119.30	105.20
26	1H	793	A	OP1-P-OP2	6.41	129.21	119.60
26	1H	1782	C	C4-C5-C6	6.41	120.60	117.40
26	1H	2697	G	OP1-P-OP2	6.41	129.21	119.60
26	1H	1955	U	C4-C5-C6	6.41	123.54	119.70
26	14	199	A	N9-C4-C5	6.41	108.36	105.80
1	13	811	C	C5-C6-N1	-6.40	117.80	121.00
26	14	2592	G	N3-C4-C5	-6.40	125.40	128.60
26	1H	188	G	C6-C5-N7	-6.40	126.56	130.40
26	1H	609	A	N1-C6-N6	6.40	122.44	118.60
26	1H	196	A	O4'-C1'-N9	6.40	113.32	108.20
6	52	14	LEU	CA-CB-CG	6.39	130.01	115.30
26	14	1845	G	N1-C6-O6	6.39	123.74	119.90
26	1H	162	U	C2-N1-C1'	6.39	125.37	117.70
26	1H	580	C	C6-N1-C2	-6.39	117.74	120.30
26	14	2339	G	O5'-P-OP2	-6.39	99.95	105.70
26	14	211	A	N1-C6-N6	6.38	122.43	118.60
26	14	664	C	C5-C6-N1	-6.38	117.81	121.00
26	14	808	G	N3-C4-N9	6.38	129.83	126.00
26	14	1781	C	O4'-C1'-N1	6.38	113.31	108.20
26	1H	412	A	C8-N9-C4	6.38	108.35	105.80
26	1H	686	G	OP1-P-OP2	6.38	129.17	119.60
26	1H	2458	G	N3-C2-N2	-6.38	115.43	119.90
26	1H	194	G	C5-C6-O6	-6.38	124.77	128.60
24	1L	34	U	C5-C6-N1	6.38	125.89	122.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	1950	G	C6-C5-N7	-6.38	126.57	130.40
26	1H	2430	A	C4-C5-N7	6.38	113.89	110.70
26	1H	835	A	O5'-P-OP2	-6.38	99.96	105.70
26	1H	99	U	N3-C2-O2	-6.38	117.74	122.20
26	1H	1800	C	C4-C5-C6	6.37	120.59	117.40
26	1H	1816	G	O5'-P-OP1	-6.37	99.96	105.70
26	1H	2311	A	N7-C8-N9	6.37	116.99	113.80
26	1H	2445	G	N3-C2-N2	6.37	124.36	119.90
1	1G	1469	G	N1-C6-O6	6.37	123.72	119.90
26	1H	2503	A	C4-C5-N7	6.37	113.89	110.70
1	1G	1446	A	O4'-C1'-N9	6.37	113.30	108.20
26	14	528	A	N1-C2-N3	6.37	132.49	129.30
26	1H	528	A	C4-N9-C1'	-6.37	114.83	126.30
26	1H	1225	C	C6-N1-C2	6.37	122.85	120.30
26	1H	150	C	C5-C4-N4	6.37	124.66	120.20
26	1H	1888	G	C8-N9-C1'	-6.37	118.72	127.00
26	1H	2434	A	C2-N3-C4	-6.37	107.42	110.60
26	1H	750	A	OP1-P-O3'	-6.37	91.19	105.20
26	1H	1142(A)	A	C5-N7-C8	-6.37	100.72	103.90
26	1H	192	C	C6-N1-C2	6.36	122.84	120.30
26	1H	2665	A	N1-C2-N3	6.36	132.48	129.30
1	1G	1487	G	N3-C2-N2	-6.36	115.45	119.90
26	14	667	U	N1-C2-O2	-6.36	118.35	122.80
26	14	1382	G	C5-C6-O6	-6.36	124.78	128.60
26	1H	2256	G	N1-C2-N2	-6.36	110.47	116.20
1	1G	108	G	C4-C5-N7	6.36	113.34	110.80
26	14	1265	A	C5-C6-N6	-6.36	118.61	123.70
26	14	2700	C	C5-C4-N4	-6.36	115.75	120.20
26	1H	858	U	O5'-P-OP2	-6.36	99.98	105.70
26	1H	987	G	OP1-P-OP2	-6.36	110.06	119.60
26	1H	2331	G	N9-C4-C5	-6.36	102.86	105.40
26	1H	2518	A	N7-C8-N9	6.36	116.98	113.80
26	1H	416	C	N3-C4-N4	-6.35	113.55	118.00
26	1H	770	G	C5-C6-O6	-6.35	124.79	128.60
26	14	1559	G	C4-C5-N7	6.35	113.34	110.80
26	1H	1210	A	C6-C5-N7	-6.35	127.85	132.30
26	1H	1437	C	C6-N1-C2	-6.35	117.76	120.30
26	14	1776	G	O5'-P-OP2	-6.35	99.99	105.70
26	14	2496	C	C6-N1-C2	-6.35	117.76	120.30
26	1H	94	G	N1-C6-O6	6.34	123.71	119.90
26	1H	2275	C	O4'-C1'-N1	-6.34	103.12	108.20
26	1H	2437	U	C5-C4-O4	6.34	129.71	125.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	1614	A	OP1-P-OP2	6.34	129.11	119.60
26	14	1760	A	O5'-P-OP2	-6.34	99.99	105.70
27	16	81	G	O4'-C1'-N9	6.34	113.27	108.20
26	14	2463	C	C6-N1-C2	6.34	122.84	120.30
24	3L	1	G	C2-N3-C4	6.34	115.07	111.90
26	1H	1210	A	C5-N7-C8	-6.34	100.73	103.90
1	13	899	C	N1-C2-O2	-6.33	115.10	118.90
26	14	1914	C	C2-N1-C1'	6.33	125.77	118.80
26	1H	917	A	N3-C4-C5	6.33	131.23	126.80
26	1H	2267	A	OP1-P-O3'	6.33	119.13	105.20
26	1H	2562	U	C5-C6-N1	-6.33	119.53	122.70
26	1H	94	G	C5-C6-O6	-6.33	124.80	128.60
26	1H	2081	C	N3-C2-O2	-6.33	117.47	121.90
26	14	629	G	O5'-P-OP2	-6.33	100.00	105.70
26	1H	729	G	C8-N9-C4	-6.33	103.87	106.40
26	14	1939	U	C5-C4-O4	6.33	129.70	125.90
26	1H	481	G	N3-C2-N2	-6.33	115.47	119.90
26	14	2056	G	C5-C6-O6	-6.33	124.81	128.60
1	13	1227	A	O5'-P-OP2	-6.32	100.01	105.70
26	14	2838	G	N3-C2-N2	-6.32	115.47	119.90
26	1H	1782	C	P-O3'-C3'	6.32	127.29	119.70
26	14	1639	U	N3-C2-O2	-6.32	117.78	122.20
26	14	1960	A	O5'-P-OP2	-6.32	100.01	105.70
27	16	6	C	N3-C2-O2	6.32	126.32	121.90
27	16	77	U	C5-C6-N1	-6.32	119.54	122.70
26	14	793	A	O5'-P-OP2	-6.32	100.01	105.70
26	1H	409	C	C6-N1-C2	6.31	122.83	120.30
26	1H	2713	A	C4-C5-N7	6.31	113.86	110.70
1	1G	481	G	C4-N9-C1'	6.31	134.71	126.50
26	14	2595	G	O5'-P-OP1	-6.31	100.02	105.70
1	13	892	A	C6-C5-N7	-6.31	127.88	132.30
26	1H	930	U	N3-C2-O2	-6.31	117.78	122.20
26	1H	1786	A	C4-C5-C6	6.31	120.15	117.00
26	1H	781	A	C8-N9-C4	6.31	108.32	105.80
26	1H	103	A	C8-N9-C4	6.30	108.32	105.80
26	1H	129	C	N3-C4-N4	6.30	122.41	118.00
26	1H	703	U	C5-C4-O4	6.30	129.68	125.90
26	1H	816	C	O5'-P-OP1	6.30	118.27	110.70
26	1H	1558	A	P-O3'-C3'	6.30	127.27	119.70
26	1H	1698	A	C4-C5-C6	6.30	120.15	117.00
26	14	199	A	N1-C6-N6	-6.30	114.82	118.60
26	1H	928	G	N1-C6-O6	6.30	123.68	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	732	C	N3-C4-N4	6.30	122.41	118.00
26	14	2355	C	C2-N1-C1'	6.30	125.73	118.80
26	14	2447	G	N1-C6-O6	6.30	123.68	119.90
26	14	2249	U	C5-C6-N1	6.30	125.85	122.70
26	14	2330	G	C5-C6-O6	-6.30	124.82	128.60
26	14	2518	A	C2-N3-C4	-6.30	107.45	110.60
26	14	1142	U	N1-C2-O2	6.30	127.21	122.80
1	13	121	C	N3-C2-O2	-6.30	117.49	121.90
26	1H	1258	C	OP2-P-O3'	6.30	119.05	105.20
26	1H	825	C	C4-C5-C6	6.29	120.55	117.40
26	1H	2513	G	O5'-P-OP2	-6.29	100.04	105.70
26	14	1784	A	C5-N7-C8	-6.29	100.75	103.90
1	13	792	A	O4'-C1'-N9	6.29	113.23	108.20
26	1H	133	C	C6-N1-C2	6.29	122.82	120.30
26	1H	265	A	N1-C2-N3	6.29	132.45	129.30
26	1H	967	C	O5'-P-OP2	-6.29	100.04	105.70
1	1G	413	G	N3-C4-N9	-6.29	122.22	126.00
26	14	912	C	C6-N1-C2	-6.29	117.78	120.30
26	1H	691	C	N1-C2-O2	-6.29	115.13	118.90
26	1H	730	C	N3-C2-O2	-6.29	117.50	121.90
26	1H	2419	U	O5'-P-OP2	6.29	118.24	110.70
26	1H	2712	U	C2-N3-C4	-6.29	123.23	127.00
1	13	346	G	C8-N9-C4	-6.28	103.89	106.40
26	1H	1416	G	O4'-C1'-N9	6.28	113.23	108.20
26	1H	1984	G	O5'-P-OP2	-6.28	100.04	105.70
26	14	2444	G	N3-C2-N2	-6.28	115.50	119.90
27	16	6	C	C5-C4-N4	-6.28	115.80	120.20
1	13	684	A	C8-N9-C4	-6.28	103.29	105.80
26	1H	138	G	C8-N9-C4	-6.28	103.89	106.40
26	14	330	A	C4-C5-N7	6.28	113.84	110.70
26	14	778	G	N1-C2-N2	-6.27	110.56	116.20
26	1H	135	G	N7-C8-N9	-6.27	109.96	113.10
1	13	1519	A	C5-C6-N1	-6.27	114.56	117.70
26	1H	1142(A)	A	N3-C4-N9	-6.27	122.38	127.40
26	1H	1379	A	C5-C6-N6	-6.27	118.69	123.70
26	1H	2442	C	C5-C6-N1	-6.27	117.86	121.00
26	1H	745	G	N3-C4-C5	-6.27	125.47	128.60
26	1H	767	U	C5-C4-O4	6.27	129.66	125.90
26	1H	2287	A	C4-C5-N7	6.26	113.83	110.70
26	1H	311	A	O5'-P-OP2	6.26	118.22	110.70
26	1H	1777	U	OP1-P-O3'	6.26	118.98	105.20
27	16	44	G	C4-N9-C1'	-6.26	118.36	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	732	C	N3-C2-O2	6.26	126.28	121.90
26	1H	906	G	N9-C4-C5	6.26	107.91	105.40
26	1H	1602	U	C5-C6-N1	-6.26	119.57	122.70
26	1H	1888	G	C4-N9-C1'	6.26	134.64	126.50
26	14	2392	A	C2-N3-C4	-6.26	107.47	110.60
26	1H	862	G	N1-C6-O6	-6.26	116.14	119.90
26	1H	1700	A	O5'-P-OP2	-6.26	100.07	105.70
26	1H	788	A	C6-N1-C2	6.26	122.35	118.60
26	14	2610	C	O5'-P-OP1	-6.26	100.07	105.70
26	14	1258	C	C5-C4-N4	-6.25	115.82	120.20
26	1H	1401	G	C8-N9-C4	-6.25	103.90	106.40
26	14	1339	G	O5'-P-OP2	6.25	118.20	110.70
26	1H	2596	U	OP1-P-OP2	6.25	128.97	119.60
26	1H	664	C	C5-C6-N1	-6.25	117.88	121.00
26	1H	1782	C	C2'-C3'-O3'	6.25	123.70	113.70
26	1H	2269	A	C8-N9-C4	6.25	108.30	105.80
26	14	1796	U	O5'-P-OP1	-6.25	100.08	105.70
26	1H	1775	U	O5'-P-OP2	-6.25	100.08	105.70
26	14	584	C	N3-C2-O2	6.25	126.27	121.90
1	13	525	C	C5-C6-N1	6.25	124.12	121.00
26	1H	1773	A	C2-N3-C4	-6.25	107.48	110.60
26	1H	1838	C	C6-N1-C2	6.24	122.80	120.30
26	1H	2500	U	N3-C4-O4	6.24	123.77	119.40
26	14	2326	C	N3-C4-C5	-6.24	119.40	121.90
26	1H	198	C	N3-C4-C5	6.24	124.40	121.90
26	1H	2665	A	O4'-C1'-N9	6.24	113.19	108.20
26	14	783	A	N3-C4-N9	-6.24	122.41	127.40
26	14	1845	G	N3-C2-N2	-6.24	115.53	119.90
26	14	2581	G	N1-C6-O6	-6.24	116.16	119.90
27	1J	7	G	N1-C6-O6	6.24	123.64	119.90
26	1H	205	G	N9-C4-C5	-6.24	102.90	105.40
26	1H	1339	G	O5'-P-OP2	6.24	118.19	110.70
1	1G	1401	G	C8-N9-C1'	-6.24	118.89	127.00
26	14	1616	A	N3-C4-C5	6.24	131.17	126.80
26	14	2003	G	C4-C5-N7	6.24	113.29	110.80
26	1H	201	C	C2-N3-C4	-6.23	116.78	119.90
24	3K	76	A	O4'-C1'-N9	6.23	113.18	108.20
26	1H	1379	A	N7-C8-N9	6.23	116.92	113.80
26	1H	1771	C	C2-N3-C4	-6.23	116.78	119.90
1	1G	1259	C	C6-N1-C2	-6.23	117.81	120.30
1	13	1502	A	C6-C5-N7	-6.23	127.94	132.30
27	16	44	G	N9-C4-C5	6.23	107.89	105.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	830	G	C8-N9-C4	6.23	108.89	106.40
26	14	1963	U	C6-N1-C1'	-6.22	112.49	121.20
26	14	2581	G	OP1-P-OP2	6.22	128.94	119.60
26	14	2239	G	N3-C2-N2	6.22	124.26	119.90
26	14	2518	A	C5-N7-C8	-6.22	100.79	103.90
23	2K	45	A	N1-C6-N6	6.22	122.33	118.60
26	1H	1984	G	N3-C2-N2	6.22	124.25	119.90
26	14	1678	G	C5-C6-N1	-6.22	108.39	111.50
24	3K	76	A	C2-N3-C4	-6.22	107.49	110.60
26	14	133	C	C6-N1-C2	6.22	122.79	120.30
26	14	488	G	N3-C4-N9	6.22	129.73	126.00
26	14	1989	G	N1-C6-O6	6.22	123.63	119.90
27	16	100	G	C8-N9-C4	6.21	108.89	106.40
26	14	2210	G	C8-N9-C1'	-6.21	118.92	127.00
26	1H	930	U	N1-C2-N3	6.21	118.63	114.90
26	1H	1241	A	C5-N7-C8	-6.21	100.80	103.90
26	1H	2053	G	C5-C6-O6	-6.21	124.87	128.60
26	1H	2604	U	N3-C2-O2	-6.21	117.85	122.20
26	14	1698	A	C6-C5-N7	-6.21	127.95	132.30
1	13	817	C	C5-C4-N4	-6.21	115.86	120.20
26	1H	2424	C	N1-C2-O2	6.21	122.62	118.90
27	16	15	A	O4'-C1'-N9	6.21	113.17	108.20
26	1H	1429	G	O5'-P-OP2	-6.20	100.12	105.70
26	1H	1957	C	N3-C2-O2	-6.20	117.56	121.90
26	1H	945	A	N9-C4-C5	-6.20	103.32	105.80
26	1H	1333	C	C5-C4-N4	-6.20	115.86	120.20
26	14	71	A	N1-C2-N3	6.20	132.40	129.30
26	1H	917	A	C8-N9-C4	-6.20	103.32	105.80
26	1H	2346	A	N9-C1'-C2'	6.20	122.05	114.00
26	14	1686	C	C6-N1-C2	6.20	122.78	120.30
26	14	2301	C	C6-N1-C2	-6.20	117.82	120.30
26	1H	752	A	C2-N3-C4	-6.19	107.50	110.60
26	1H	1626	G	O5'-P-OP2	6.19	118.13	110.70
26	1H	2510	C	N3-C4-N4	-6.19	113.67	118.00
26	14	1379	A	C5-N7-C8	-6.19	100.81	103.90
26	14	2473	U	N1-C2-O2	6.19	127.13	122.80
26	1H	96	G	N1-C6-O6	6.19	123.61	119.90
26	1H	1786	A	C4-N9-C1'	6.19	137.44	126.30
26	1H	2397	G	O5'-P-OP2	6.19	118.13	110.70
26	14	676	A	N1-C6-N6	6.19	122.31	118.60
26	1H	845	G	OP1-P-O3'	6.19	118.81	105.20
26	14	845	G	C6-C5-N7	-6.18	126.69	130.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	2417	C	O5'-P-OP2	-6.18	100.14	105.70
26	14	530	G	N1-C6-O6	6.18	123.61	119.90
26	14	1142(A)	A	C2-N3-C4	-6.18	107.51	110.60
26	14	2446	G	N3-C2-N2	6.18	124.22	119.90
1	13	800	G	C5-C6-O6	-6.18	124.89	128.60
26	1H	2035	G	O5'-P-OP1	-6.18	100.14	105.70
26	1H	2593	U	C5-C4-O4	6.18	129.61	125.90
1	13	1398	A	N1-C6-N6	6.18	122.31	118.60
26	1H	1995	U	C5-C6-N1	-6.18	119.61	122.70
1	1G	700	G	C6-C5-N7	6.18	134.11	130.40
26	14	2062	A	C4-C5-C6	-6.18	113.91	117.00
26	14	2427	C	O5'-P-OP2	6.18	118.11	110.70
26	1H	1394	U	C5-C6-N1	6.17	125.79	122.70
26	1H	1611	C	C5-C4-N4	-6.17	115.88	120.20
26	1H	621	A	C8-N9-C4	-6.17	103.33	105.80
27	1J	89	G	C4-N9-C1'	6.17	134.52	126.50
26	1H	1955	U	N1-C2-N3	6.17	118.60	114.90
26	14	933	A	N1-C6-N6	6.17	122.30	118.60
26	14	1241	A	C5-C6-N1	-6.17	114.61	117.70
26	14	2464	C	N3-C4-C5	6.17	124.37	121.90
26	1H	736	C	O5'-P-OP2	6.17	118.10	110.70
23	2L	21	U	N3-C2-O2	-6.17	117.88	122.20
26	14	1797	C	C6-N1-C2	6.17	122.77	120.30
26	1H	760	G	N1-C6-O6	6.17	123.60	119.90
26	14	2245	U	C4-C5-C6	-6.17	116.00	119.70
1	13	326	G	C4-C5-N7	-6.16	108.33	110.80
26	1H	599	G	N3-C4-C5	-6.16	125.52	128.60
26	14	2426	A	N1-C6-N6	6.16	122.30	118.60
26	14	1725	G	C4-N9-C1'	6.16	134.51	126.50
1	13	1220	G	N1-C6-O6	-6.16	116.20	119.90
1	1G	1487	G	N1-C6-O6	6.16	123.59	119.90
1	1G	817	C	C6-N1-C2	6.16	122.76	120.30
26	14	577	G	N1-C6-O6	6.16	123.59	119.90
8	7E	112	LEU	CA-CB-CG	6.16	129.46	115.30
26	14	1314	C	C2-N1-C1'	6.16	125.57	118.80
26	1H	1332	G	N1-C2-N3	6.15	127.59	123.90
26	1H	2572	A	N9-C4-C5	-6.15	103.34	105.80
27	16	69	G	OP2-P-O3'	6.15	118.73	105.20
1	1G	1397	C	C2-N1-C1'	6.15	125.57	118.80
1	1G	1500	A	O5'-P-OP1	-6.15	100.16	105.70
26	14	1241	A	C2-N3-C4	-6.15	107.52	110.60
26	14	2251	G	N1-C6-O6	-6.15	116.21	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	13	266	G	C5-N7-C8	-6.15	101.22	104.30
26	1H	728	G	C5-C6-O6	-6.15	124.91	128.60
26	1H	1611	C	C2-N3-C4	-6.15	116.82	119.90
26	14	1968	G	N7-C8-N9	6.15	116.18	113.10
26	14	2435	A	N7-C8-N9	6.15	116.88	113.80
26	1H	456	C	OP1-P-O3'	6.15	118.73	105.20
26	14	681	G	N1-C2-N2	-6.15	110.67	116.20
26	14	1496	A	C4-C5-N7	6.15	113.78	110.70
26	14	113	G	N3-C4-C5	6.15	131.67	128.60
26	1H	2328	A	N1-C2-N3	6.15	132.37	129.30
26	14	945	A	C4-N9-C1'	6.15	137.36	126.30
26	14	1390	U	OP1-P-O3'	6.15	118.72	105.20
1	13	1158	C	C2-N1-C1'	6.14	125.56	118.80
26	1H	2258	C	C5-C6-N1	-6.14	117.93	121.00
26	1H	2689	U	C6-N1-C2	-6.14	117.31	121.00
26	14	1416	G	C4-N9-C1'	-6.14	118.51	126.50
26	14	2496	C	C5-C6-N1	6.14	124.07	121.00
26	1H	746	A	O4'-C1'-N9	6.14	113.11	108.20
26	1H	2752	C	C6-N1-C2	-6.14	117.84	120.30
26	14	140	A	O4'-C1'-N9	6.14	113.11	108.20
26	14	71	A	P-O3'-C3'	6.14	127.07	119.70
26	1H	651	G	C8-N9-C4	-6.14	103.94	106.40
26	14	1299	G	O5'-P-OP1	-6.13	100.18	105.70
27	16	81	G	N7-C8-N9	6.13	116.17	113.10
26	1H	1599	C	N3-C4-N4	-6.13	113.71	118.00
24	3K	71	C	C6-N1-C2	-6.13	117.85	120.30
26	1H	655	A	N7-C8-N9	6.13	116.86	113.80
26	1H	1797	C	C5-C4-N4	-6.13	115.91	120.20
26	1H	2353	G	OP1-P-OP2	6.13	128.80	119.60
26	14	778	G	N3-C2-N2	6.13	124.19	119.90
26	1H	1365	A	C8-N9-C4	-6.13	103.35	105.80
26	14	72	U	C5-C6-N1	-6.13	119.64	122.70
26	14	2607	G	C6-C5-N7	-6.13	126.72	130.40
26	1H	1396	U	C5-C4-O4	6.12	129.57	125.90
26	1H	1796	U	O5'-P-OP2	6.12	118.05	110.70
26	1H	2377	A	C8-N9-C4	6.12	108.25	105.80
26	14	1333	C	N3-C4-C5	6.12	124.35	121.90
1	13	767	A	C2-N3-C4	-6.12	107.54	110.60
26	1H	983	A	C8-N9-C4	6.12	108.25	105.80
26	1H	1284	A	OP1-P-OP2	6.12	128.78	119.60
26	1H	2518	A	O5'-P-OP2	6.12	118.05	110.70
26	14	1259	G	OP2-P-O3'	6.12	118.67	105.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
36	25	8	LEU	CA-CB-CG	6.12	129.38	115.30
26	1H	2500	U	OP2-P-O3'	6.12	118.66	105.20
26	1H	2578	G	C5-C6-N1	6.12	114.56	111.50
26	1H	632	A	O5'-P-OP2	6.12	118.04	110.70
26	1H	1899	G	N1-C2-N2	6.12	121.70	116.20
26	1H	2500	U	C5-C4-O4	-6.12	122.23	125.90
1	13	690	G	C5-N7-C8	-6.11	101.24	104.30
23	2L	21	U	C6-N1-C2	-6.11	117.33	121.00
26	1H	932	G	C8-N9-C4	-6.11	103.95	106.40
1	1G	536	C	C6-N1-C2	-6.11	117.86	120.30
26	14	330	A	C5-N7-C8	-6.11	100.84	103.90
26	1H	180	G	N9-C4-C5	-6.11	102.96	105.40
26	1H	213	A	C5-N7-C8	-6.11	100.85	103.90
26	1H	777	A	N1-C2-N3	6.11	132.35	129.30
26	1H	189	G	C6-C5-N7	-6.11	126.74	130.40
26	1H	1663	C	C5-C4-N4	-6.11	115.93	120.20
26	14	2270	G	N1-C6-O6	6.11	123.56	119.90
1	13	57	G	N3-C4-C5	-6.10	125.55	128.60
26	1H	1790	C	P-O3'-C3'	6.10	127.02	119.70
26	14	1475	G	N7-C8-N9	6.10	116.15	113.10
26	14	1638	C	OP2-P-O3'	6.10	118.63	105.20
26	14	2594	C	C5-C4-N4	-6.10	115.93	120.20
1	13	963	G	N3-C4-N9	6.10	129.66	126.00
26	1H	520	G	N1-C6-O6	-6.10	116.24	119.90
26	1H	1184	G	OP2-P-O3'	6.10	118.62	105.20
27	1J	47	C	C6-N1-C2	6.10	122.74	120.30
26	1H	1678	G	C6-C5-N7	-6.10	126.74	130.40
1	13	481	G	N1-C6-O6	6.10	123.56	119.90
26	1H	219	G	C5-C6-N1	6.10	114.55	111.50
26	14	2065	C	N3-C2-O2	-6.10	117.63	121.90
1	13	115	G	P-O3'-C3'	6.09	127.01	119.70
26	14	1308	A	C8-N9-C4	-6.09	103.36	105.80
26	14	2069	G	N9-C4-C5	-6.09	102.96	105.40
26	14	2842	G	N1-C6-O6	6.09	123.55	119.90
26	1H	967	C	N3-C4-C5	6.09	124.33	121.90
27	16	28	C	C6-N1-C2	-6.09	117.86	120.30
1	13	813	U	C5-C4-O4	6.09	129.55	125.90
26	1H	2604	U	N1-C2-O2	6.09	127.06	122.80
26	14	1122	G	C5-C6-O6	-6.09	124.95	128.60
26	14	808	G	N3-C2-N2	6.08	124.16	119.90
26	14	2592	G	O5'-P-OP2	-6.08	100.22	105.70
26	1H	2052	G	N1-C6-O6	-6.08	116.25	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	2830	G	C8-N9-C4	-6.08	103.97	106.40
26	14	2307	G	C4-N9-C1'	6.08	134.41	126.50
26	1H	999	U	OP1-P-OP2	-6.08	110.48	119.60
26	1H	1431	U	C5-C6-N1	6.08	125.74	122.70
26	1H	270(L)	U	O4'-C1'-N1	6.08	113.06	108.20
26	1H	1197	G	C8-N9-C4	6.08	108.83	106.40
26	1H	1611	C	N3-C4-C5	6.08	124.33	121.90
26	1H	1764	G	N1-C6-O6	-6.08	116.25	119.90
26	14	396	G	C4-N9-C1'	6.08	134.40	126.50
1	13	1450	U	N1-C2-O2	6.08	127.05	122.80
1	13	576	G	C5-C6-O6	-6.07	124.96	128.60
26	1H	2837	G	C8-N9-C4	-6.07	103.97	106.40
26	14	1391	U	O5'-P-OP2	6.07	117.99	110.70
26	1H	2706	G	C5-N7-C8	6.07	107.34	104.30
1	1G	197	A	N7-C8-N9	6.07	116.84	113.80
26	14	1932	A	O5'-P-OP1	-6.07	100.23	105.70
26	14	2358	G	N3-C2-N2	-6.07	115.65	119.90
26	1H	1157	G	N3-C4-C5	-6.07	125.56	128.60
26	1H	1191	G	C8-N9-C4	6.07	108.83	106.40
26	1H	2082	A	N1-C2-N3	6.07	132.34	129.30
26	14	827	U	N1-C2-O2	-6.07	118.55	122.80
1	13	703	G	C4-C5-N7	6.07	113.23	110.80
26	1H	189	G	N9-C4-C5	-6.07	102.97	105.40
26	1H	944	G	C4-N9-C1'	6.07	134.38	126.50
26	1H	1555	G	O5'-P-OP1	-6.07	100.24	105.70
26	1H	1970	A	O5'-P-OP2	-6.07	100.24	105.70
26	14	729	G	OP2-P-O3'	6.07	118.54	105.20
26	14	1950	G	C6-C5-N7	-6.07	126.76	130.40
26	1H	1496	A	C6-C5-N7	-6.06	128.06	132.30
26	1H	2512	C	C6-N1-C2	6.06	122.72	120.30
26	1H	1142(A)	A	N1-C6-N6	6.06	122.24	118.60
26	1H	1198	U	N3-C2-O2	-6.06	117.96	122.20
27	1J	30	C	N3-C4-C5	-6.06	119.48	121.90
26	1H	1591	G	C5-C6-O6	6.06	132.23	128.60
26	1H	1858	G	C4-N9-C1'	6.06	134.38	126.50
1	1G	906	G	N1-C6-O6	6.06	123.53	119.90
26	14	331	A	C2-N3-C4	-6.06	107.57	110.60
26	1H	820	A	C5-C6-N6	6.06	128.54	123.70
27	16	12	C	C6-N1-C2	-6.06	117.88	120.30
26	1H	2706	G	N7-C8-N9	-6.05	110.07	113.10
1	13	975	A	O4'-C1'-N9	-6.05	103.36	108.20
26	1H	808	G	N1-C2-N2	-6.05	110.75	116.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	1G	1498	U	P-O3'-C3'	6.05	126.96	119.70
26	1H	921	G	C8-N9-C4	-6.05	103.98	106.40
26	1H	121	G	C5-C6-O6	-6.05	124.97	128.60
26	1H	141	A	N3-C4-C5	6.05	131.03	126.80
26	1H	616	A	OP2-P-O3'	6.05	118.51	105.20
26	1H	834	C	OP2-P-O3'	6.05	118.51	105.20
26	1H	2039	C	C6-N1-C2	-6.05	117.88	120.30
26	1H	576	U	C6-N1-C2	-6.05	117.37	121.00
26	14	1934	C	N1-C2-O2	6.05	122.53	118.90
26	14	2842	G	C5-C6-O6	-6.05	124.97	128.60
26	1H	946	G	N7-C8-N9	-6.04	110.08	113.10
26	14	47	C	OP2-P-O3'	6.04	118.50	105.20
26	14	1323	U	OP1-P-O3'	6.04	118.49	105.20
26	14	2068	U	OP1-P-O3'	6.04	118.50	105.20
26	1H	508	G	C6-C5-N7	-6.04	126.78	130.40
26	14	1914	C	N3-C4-C5	-6.04	119.48	121.90
26	1H	2408	U	OP2-P-O3'	6.04	118.49	105.20
1	1G	1401	G	C6-C5-N7	-6.04	126.78	130.40
26	14	2560	C	O5'-P-OP1	-6.04	100.26	105.70
1	13	536	C	N1-C2-O2	-6.04	115.28	118.90
26	1H	2572	A	N1-C6-N6	6.04	122.22	118.60
26	14	1992	G	C2-N3-C4	6.04	114.92	111.90
26	1H	2057	A	OP1-P-O3'	6.03	118.47	105.20
26	14	565	C	C6-N1-C2	6.03	122.71	120.30
26	14	1367	A	N1-C6-N6	6.03	122.22	118.60
26	14	1258	C	OP2-P-O3'	6.03	118.47	105.20
26	1H	824	A	N1-C6-N6	-6.03	114.98	118.60
26	1H	1645	G	OP1-P-O3'	6.03	118.47	105.20
26	14	642	G	C6-C5-N7	-6.03	126.78	130.40
26	14	1671	U	O5'-P-OP1	-6.03	100.27	105.70
26	14	503	A	N1-C6-N6	-6.03	114.98	118.60
26	14	783	A	C4-C5-C6	6.03	120.02	117.00
26	1H	845	G	N3-C4-C5	6.03	131.61	128.60
26	1H	2616	C	N3-C4-C5	6.03	124.31	121.90
1	1G	1128	C	N1-C2-O2	6.03	122.52	118.90
26	1H	590	A	C6-N1-C2	-6.03	114.98	118.60
27	16	98	G	N9-C4-C5	-6.03	102.99	105.40
26	1H	330	A	C5-N7-C8	-6.02	100.89	103.90
26	1H	821	A	OP1-P-O3'	6.02	118.45	105.20
26	1H	860	U	C6-N1-C1'	-6.02	112.77	121.20
26	14	837	C	C6-N1-C2	-6.02	117.89	120.30
26	1H	797	C	C2-N3-C4	-6.02	116.89	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	2706	G	N1-C6-O6	-6.02	116.29	119.90
26	1H	679	C	C2-N3-C4	-6.02	116.89	119.90
26	1H	1776	G	N3-C4-N9	6.02	129.61	126.00
26	14	2304	G	C8-N9-C4	-6.02	103.99	106.40
26	1H	2251	G	C4-C5-N7	-6.02	108.39	110.80
26	14	2708	G	N1-C2-N2	-6.02	110.79	116.20
26	14	252	G	N1-C6-O6	-6.01	116.29	119.90
26	14	1965	C	C6-N1-C2	6.01	122.70	120.30
26	1H	113	G	N3-C4-C5	6.01	131.60	128.60
26	14	2346	A	C4-N9-C1'	6.01	137.12	126.30
26	1H	738	G	C4-C5-N7	6.01	113.20	110.80
26	1H	1445	C	C6-N1-C2	-6.01	117.90	120.30
26	1H	1432	C	N1-C2-O2	-6.01	115.30	118.90
26	1H	2270	G	N9-C4-C5	-6.01	103.00	105.40
26	1H	2485	G	C8-N9-C4	6.01	108.80	106.40
26	14	596	G	N1-C2-N2	6.01	121.61	116.20
26	1H	1858	G	P-O3'-C3'	6.00	126.91	119.70
22	1K	75	C	C5-C6-N1	6.00	124.00	121.00
26	1H	23	G	N1-C2-N2	6.00	121.60	116.20
27	16	47	C	C6-N1-C2	6.00	122.70	120.30
26	1H	585	G	O5'-P-OP2	6.00	117.90	110.70
26	14	2394	C	O5'-P-OP2	-6.00	100.30	105.70
26	1H	245	G	C5-C6-O6	-6.00	125.00	128.60
26	14	1643	G	OP2-P-O3'	6.00	118.40	105.20
26	1H	74	A	N1-C6-N6	6.00	122.20	118.60
26	1H	516	C	C6-N1-C2	-6.00	117.90	120.30
26	1H	2609	U	O5'-P-OP2	-6.00	100.30	105.70
26	1H	142	G	C2-N3-C4	-6.00	108.90	111.90
26	14	1255	U	N1-C2-O2	6.00	127.00	122.80
26	14	2473	U	N3-C2-O2	-6.00	118.00	122.20
26	1H	2084	C	C5-C6-N1	-5.99	118.00	121.00
26	1H	2430	A	C6-C5-N7	-5.99	128.10	132.30
26	1H	832	G	C8-N9-C4	-5.99	104.00	106.40
26	1H	1193	G	C8-N9-C4	5.99	108.80	106.40
26	14	1762	A	C4-N9-C1'	5.99	137.09	126.30
26	1H	113	G	N3-C2-N2	-5.99	115.71	119.90
26	1H	392	C	O5'-P-OP2	5.99	117.89	110.70
26	14	944	G	N3-C4-N9	5.99	129.59	126.00
26	14	621	A	N1-C2-N3	5.99	132.29	129.30
26	14	675	A	N9-C4-C5	-5.99	103.41	105.80
1	1G	687	A	P-O3'-C3'	5.99	126.88	119.70
1	1G	1200	C	C2-N1-C1'	5.99	125.39	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	1905	C	O5'-P-OP2	-5.99	100.31	105.70
26	1H	2507	C	C2-N1-C1'	5.98	125.38	118.80
26	14	1496	A	C6-C5-N7	-5.98	128.11	132.30
26	1H	2346	A	C8-N9-C1'	-5.98	116.93	127.70
26	14	704	G	N1-C6-O6	5.98	123.49	119.90
26	1H	655	A	C5-N7-C8	-5.98	100.91	103.90
26	1H	1893	C	O5'-P-OP2	-5.98	100.32	105.70
26	1H	124	G	N3-C4-N9	-5.98	122.41	126.00
26	1H	240	G	N9-C4-C5	-5.98	103.01	105.40
26	14	2238	G	OP1-P-O3'	5.98	118.35	105.20
26	1H	1399	C	OP2-P-O3'	5.98	118.35	105.20
26	14	1251	C	N3-C4-N4	5.98	122.18	118.00
26	1H	1022	G	C4-C5-N7	-5.97	108.41	110.80
26	1H	1027	A	C2-N3-C4	-5.97	107.61	110.60
26	1H	1314	C	C6-N1-C1'	-5.97	113.63	120.80
45	F8	70	LEU	CA-CB-CG	5.97	129.04	115.30
1	13	505	G	C4-C5-N7	5.97	113.19	110.80
26	1H	917	A	O5'-P-OP2	5.97	117.87	110.70
26	14	459	U	O5'-P-OP2	-5.97	100.33	105.70
26	14	2213	U	C2-N1-C1'	5.97	124.86	117.70
1	13	1158	C	N1-C2-O2	5.97	122.48	118.90
26	14	530	G	N9-C4-C5	-5.97	103.01	105.40
26	14	1657	C	C6-N1-C2	-5.97	117.91	120.30
26	14	2713	A	C8-N9-C4	-5.97	103.41	105.80
26	1H	2235	G	C4-C5-N7	5.96	113.19	110.80
26	1H	2277	G	C5-N7-C8	5.96	107.28	104.30
26	1H	789	A	O5'-P-OP1	-5.96	100.33	105.70
26	1H	1785	A	OP2-P-O3'	5.96	118.31	105.20
27	16	60	C	C6-N1-C2	-5.96	117.92	120.30
26	14	1767	C	N3-C4-N4	-5.96	113.83	118.00
27	1J	7	G	N9-C4-C5	-5.96	103.02	105.40
26	14	829	A	OP1-P-OP2	5.96	128.54	119.60
26	14	944	G	C8-N9-C1'	-5.96	119.25	127.00
26	14	1617	C	C4-C5-C6	5.96	120.38	117.40
26	14	960	A	OP1-P-O3'	5.96	118.31	105.20
1	13	911	U	N3-C2-O2	-5.95	118.03	122.20
26	1H	141(A)	C	OP1-P-O3'	-5.95	92.10	105.20
26	1H	209	C	N3-C4-C5	5.95	124.28	121.90
26	1H	247	G	C5-C6-O6	-5.95	125.03	128.60
26	1H	807	U	OP1-P-OP2	5.95	128.53	119.60
26	1H	2611	U	N3-C4-O4	-5.95	115.23	119.40
1	1G	449	C	C5-C4-N4	5.95	124.37	120.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	445	C	O5'-P-OP1	-5.95	100.34	105.70
26	14	2599	G	OP2-P-O3'	5.95	118.30	105.20
26	1H	337	C	N1-C2-O2	-5.95	115.33	118.90
26	1H	1757	U	OP1-P-O3'	5.95	118.29	105.20
1	13	897	C	C5-C6-N1	-5.95	118.03	121.00
26	1H	1773	A	C5-C6-N1	-5.95	114.72	117.70
26	1H	2031	A	C2-N3-C4	5.95	113.57	110.60
26	14	1616	A	C6-C5-N7	-5.95	128.14	132.30
26	14	2085	C	O5'-P-OP2	-5.95	100.35	105.70
26	14	2256	G	N1-C2-N2	-5.95	110.85	116.20
1	13	1498	U	P-O3'-C3'	5.95	126.83	119.70
23	2K	9	G	C5-C6-O6	-5.95	125.03	128.60
26	14	380	U	O5'-P-OP2	-5.95	100.35	105.70
26	14	2440	C	O5'-P-OP1	-5.95	100.35	105.70
26	1H	247	G	C8-N9-C4	5.94	108.78	106.40
26	1H	273(A)	G	C8-N9-C4	5.94	108.78	106.40
26	1H	1437	C	C2-N1-C1'	5.94	125.34	118.80
26	1H	2554	U	C5-C4-O4	-5.94	122.33	125.90
1	1G	27	G	N3-C2-N2	-5.94	115.74	119.90
1	13	1199	U	N1-C2-N3	5.94	118.46	114.90
26	1H	128	C	N1-C2-O2	5.94	122.47	118.90
26	1H	1625	C	N3-C4-N4	-5.94	113.84	118.00
1	1G	917	G	O5'-P-OP1	-5.94	100.35	105.70
26	14	1639	U	N3-C4-O4	-5.94	115.24	119.40
26	14	1807	G	N9-C1'-C2'	-5.94	105.47	112.00
26	1H	1699	G	O5'-P-OP1	-5.94	100.36	105.70
1	1G	413	G	C8-N9-C1'	5.94	134.72	127.00
26	1H	263	C	O5'-P-OP1	5.94	117.83	110.70
26	1H	815	C	N3-C4-C5	5.94	124.28	121.90
26	1H	2610	C	O5'-P-OP1	-5.94	100.36	105.70
26	14	577	G	C8-N9-C4	5.94	108.78	106.40
26	1H	2754	U	C5-C4-O4	-5.93	122.34	125.90
1	13	917	G	OP1-P-O3'	5.93	118.25	105.20
26	1H	1685	C	O5'-P-OP2	5.93	117.82	110.70
1	1G	422	C	O4'-C1'-N1	5.93	112.94	108.20
1	1G	1157	A	P-O3'-C3'	5.93	126.82	119.70
26	14	2069	G	C8-N9-C4	5.93	108.77	106.40
1	13	827	U	N3-C2-O2	-5.93	118.05	122.20
26	1H	861	A	N9-C4-C5	-5.93	103.43	105.80
26	1H	1604	C	O5'-P-OP2	5.93	117.81	110.70
26	1H	741	G	C5-C6-O6	-5.93	125.04	128.60
26	1H	120	U	O5'-P-OP2	5.93	117.81	110.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	383	U	O5'-P-OP2	5.93	117.81	110.70
26	1H	1931	U	N1-C2-N3	5.93	118.46	114.90
26	1H	376	C	N3-C2-O2	-5.92	117.75	121.90
26	1H	463	G	N3-C2-N2	5.92	124.05	119.90
26	1H	795	C	O5'-P-OP2	-5.92	100.37	105.70
1	13	1199	U	N3-C2-O2	-5.92	118.06	122.20
26	14	2518	A	O4'-C1'-N9	-5.92	103.46	108.20
26	14	1992	G	N3-C4-C5	-5.92	125.64	128.60
26	14	2244	U	C4-C5-C6	5.92	123.25	119.70
26	14	2592	G	N1-C2-N2	-5.92	110.87	116.20
1	13	1129	C	C5-C6-N1	5.92	123.96	121.00
26	1H	1957	C	N3-C4-N4	-5.92	113.86	118.00
26	1H	2388	A	O4'-C1'-N9	5.92	112.93	108.20
23	2L	48	U	P-O3'-C3'	5.92	126.80	119.70
26	14	988	A	N1-C6-N6	5.92	122.15	118.60
26	14	2596	U	OP1-P-OP2	5.92	128.47	119.60
26	1H	859	G	N3-C4-C5	5.91	131.56	128.60
26	1H	2072	G	OP1-P-O3'	5.91	118.21	105.20
27	16	79	C	N3-C2-O2	-5.91	117.76	121.90
41	B8	13	ARG	N-CA-C	5.91	126.97	111.00
26	14	1963	U	C5-C6-N1	5.91	125.66	122.70
27	1J	47	C	OP1-P-O3'	5.91	118.21	105.20
26	1H	816	C	N3-C4-N4	5.91	122.14	118.00
26	1H	1137	G	OP1-P-O3'	5.91	118.20	105.20
26	1H	2837	G	N7-C8-N9	5.91	116.06	113.10
26	14	1496	A	O4'-C1'-N9	5.91	112.93	108.20
4	3E	11	LEU	CA-CB-CG	5.91	128.89	115.30
1	1G	1281	U	N1-C2-O2	5.91	126.94	122.80
26	14	2598	A	OP2-P-O3'	5.91	118.20	105.20
26	14	2607	G	N3-C4-N9	5.91	129.54	126.00
26	1H	2699	C	C5-C4-N4	-5.91	116.06	120.20
26	1H	1825	A	N1-C6-N6	-5.91	115.06	118.60
25	4L	23	A	P-O3'-C3'	5.91	126.79	119.70
26	14	1786	A	C4-N9-C1'	5.91	136.93	126.30
26	14	1815	A	OP1-P-O3'	5.91	118.19	105.20
26	1H	1315	C	N1-C2-O2	5.90	122.44	118.90
26	1H	1520	U	N3-C2-O2	-5.90	118.07	122.20
26	1H	189	G	C5-C6-O6	-5.90	125.06	128.60
1	1G	1301	U	C2-N1-C1'	5.90	124.78	117.70
26	14	1506	C	C6-N1-C2	-5.90	117.94	120.30
26	14	1762	A	N9-C1'-C2'	5.90	121.67	114.00
26	14	512	G	O4'-C1'-N9	5.90	112.92	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	857	C	O5'-P-OP2	-5.90	100.39	105.70
26	1H	1674	G	O4'-C1'-N9	-5.90	103.48	108.20
26	14	1304	C	C5-C4-N4	5.90	124.33	120.20
26	1H	929	G	N9-C4-C5	-5.90	103.04	105.40
26	1H	1376	C	N3-C4-C5	-5.90	119.54	121.90
26	1H	1649	G	N3-C4-C5	-5.90	125.65	128.60
26	14	1304	C	N3-C2-O2	-5.90	117.77	121.90
26	1H	1534	G	C4-N9-C1'	5.89	134.16	126.50
26	1H	2848	G	O5'-P-OP2	-5.89	100.40	105.70
26	1H	835	A	O5'-P-OP1	5.89	117.77	110.70
26	1H	1432	C	N3-C4-N4	5.89	122.12	118.00
1	1G	481	G	N3-C4-N9	5.89	129.53	126.00
26	14	946	G	N9-C4-C5	-5.89	103.04	105.40
26	1H	2450	A	N7-C8-N9	5.89	116.75	113.80
26	1H	2507	C	C5-C6-N1	5.89	123.94	121.00
26	14	750	A	N7-C8-N9	5.89	116.75	113.80
26	1H	598	G	O5'-P-OP2	-5.89	100.40	105.70
26	1H	2518	A	C5-C6-N6	-5.89	118.99	123.70
26	1H	952	G	N3-C2-N2	5.89	124.02	119.90
26	1H	1618	A	C8-N9-C4	-5.89	103.44	105.80
26	1H	1776	G	N9-C4-C5	-5.89	103.05	105.40
26	1H	1899	G	C8-N9-C4	-5.89	104.05	106.40
26	14	2597	G	C6-C5-N7	-5.89	126.87	130.40
26	1H	1585	C	C4-C5-C6	5.88	120.34	117.40
26	14	693	C	OP2-P-O3'	5.88	118.15	105.20
26	14	2413	G	N1-C6-O6	5.88	123.43	119.90
26	1H	112	U	C5-C4-O4	-5.88	122.37	125.90
26	1H	455	C	C6-N1-C2	5.88	122.65	120.30
26	1H	2374	C	C6-N1-C2	5.88	122.65	120.30
26	14	1350	C	O5'-P-OP1	-5.88	100.41	105.70
26	14	2573	C	N1-C2-O2	5.88	122.43	118.90
27	1J	89	G	N3-C4-N9	5.88	129.53	126.00
26	1H	138	G	O4'-C1'-N9	5.88	112.90	108.20
26	14	2724	C	OP2-P-O3'	5.88	118.13	105.20
1	13	905	U	O5'-P-OP2	5.88	117.75	110.70
26	1H	1301	A	C5-C6-N6	-5.87	119.00	123.70
26	1H	1915	U	N3-C2-O2	-5.87	118.09	122.20
1	13	872	A	C6-N1-C2	5.87	122.12	118.60
26	1H	136	G	N1-C6-O6	-5.87	116.38	119.90
26	14	1899	G	C5-C6-O6	5.87	132.12	128.60
27	1J	54	G	C8-N9-C4	-5.87	104.05	106.40
26	1H	99	U	C2-N1-C1'	5.87	124.74	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	378	C	C6-N1-C2	5.87	122.65	120.30
26	1H	1396	U	OP1-P-OP2	5.87	128.40	119.60
26	1H	1672	C	O5'-P-OP2	5.87	117.74	110.70
26	1H	2618	G	C8-N9-C4	-5.87	104.05	106.40
26	14	1939	U	N3-C4-O4	-5.87	115.29	119.40
26	1H	2297	C	OP1-P-OP2	5.87	128.40	119.60
1	1G	817	C	C5-C6-N1	-5.87	118.07	121.00
26	14	953	A	OP1-P-O3'	5.87	118.11	105.20
26	1H	2270	G	C5-C6-O6	-5.87	125.08	128.60
1	13	57	G	N1-C6-O6	-5.86	116.38	119.90
26	1H	66	C	C6-N1-C2	-5.86	117.95	120.30
26	1H	1379	A	C4-C5-N7	5.86	113.63	110.70
26	14	1022	G	C8-N9-C4	-5.86	104.05	106.40
26	14	1902	C	O5'-P-OP2	5.86	117.74	110.70
26	14	2055	C	C2-N1-C1'	-5.86	112.35	118.80
26	1H	226	G	C5-C6-O6	-5.86	125.08	128.60
26	1H	906	G	C8-N9-C1'	5.86	134.62	127.00
26	1H	1940	U	O5'-P-OP2	-5.86	100.42	105.70
26	1H	2574	G	C5-C6-N1	5.86	114.43	111.50
26	14	2270	G	C4-C5-N7	5.86	113.14	110.80
26	1H	2589	A	N7-C8-N9	-5.86	110.87	113.80
26	1H	2752	C	C5-C6-N1	5.86	123.93	121.00
26	1H	674	G	C8-N9-C4	5.86	108.74	106.40
26	1H	2324	C	C5-C4-N4	-5.86	116.10	120.20
26	14	1342	A	N1-C6-N6	5.86	122.11	118.60
47	D5	76	LEU	CA-CB-CG	5.86	128.78	115.30
1	13	703	G	C4-N9-C1'	5.86	134.12	126.50
1	13	1335	C	C6-N1-C2	5.86	122.64	120.30
26	1H	48	G	OP2-P-O3'	5.86	118.09	105.20
26	1H	74	A	O4'-C1'-N9	-5.86	103.51	108.20
26	14	2841	C	C6-N1-C2	5.86	122.64	120.30
26	1H	2582	G	O5'-P-OP1	5.85	117.72	110.70
26	1H	383	U	C5-C6-N1	-5.85	119.78	122.70
26	14	676	A	N3-C4-N9	-5.85	122.72	127.40
26	14	1973	G	N1-C2-N2	-5.85	110.94	116.20
1	13	22	G	N3-C2-N2	-5.85	115.81	119.90
26	14	1520	U	C5-C4-O4	5.85	129.41	125.90
26	14	2415	G	N3-C2-N2	-5.85	115.81	119.90
26	1H	2374	C	C2-N3-C4	-5.85	116.98	119.90
26	1H	2392	A	C6-N1-C2	5.84	122.11	118.60
27	16	98	G	OP1-P-OP2	5.84	128.37	119.60
26	1H	265	A	C5-N7-C8	-5.84	100.98	103.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	808	G	N3-C4-C5	-5.84	125.68	128.60
26	1H	786	C	N3-C2-O2	-5.84	117.81	121.90
26	1H	836	G	C2-N3-C4	5.84	114.82	111.90
26	1H	1601	G	OP1-P-O3'	5.84	118.05	105.20
1	13	942	G	OP1-P-O3'	5.84	118.05	105.20
26	14	1819	A	P-O3'-C3'	5.84	126.71	119.70
26	14	2347	C	N3-C2-O2	-5.84	117.81	121.90
26	14	2464	C	C5-C6-N1	-5.84	118.08	121.00
26	1H	1204	A	C4-C5-N7	5.84	113.62	110.70
27	1J	60	C	C5-C6-N1	5.84	123.92	121.00
26	1H	667	U	N3-C4-O4	5.84	123.48	119.40
26	1H	820	A	N1-C6-N6	-5.84	115.10	118.60
26	1H	1752	C	N3-C2-O2	5.84	125.98	121.90
26	1H	2449	U	OP2-P-O3'	5.84	118.04	105.20
26	14	2755	C	C2-N1-C1'	5.84	125.22	118.80
32	49	2	PRO	N-CA-CB	5.84	110.30	103.30
26	1H	1315	C	N3-C2-O2	-5.83	117.82	121.90
26	1H	1763	G	O5'-P-OP1	5.83	117.70	110.70
26	1H	2086	U	C5-C4-O4	5.83	129.40	125.90
26	1H	860	U	O5'-P-OP1	5.83	117.70	110.70
26	1H	1698	A	N3-C4-C5	5.83	130.88	126.80
1	13	802	A	N9-C4-C5	-5.83	103.47	105.80
1	13	1455	G	C8-N9-C4	5.83	108.73	106.40
26	1H	2069	G	C8-N9-C4	5.83	108.73	106.40
26	1H	2392	A	O4'-C1'-N9	5.83	112.86	108.20
12	3A	27	LEU	CA-CB-CG	5.83	128.71	115.30
26	14	1253	A	N9-C4-C5	-5.83	103.47	105.80
26	14	2067	G	N3-C4-C5	-5.83	125.69	128.60
26	1H	2510	C	C5-C4-N4	5.83	124.28	120.20
1	1G	495	A	N1-C6-N6	-5.83	115.10	118.60
1	1G	1338	G	N3-C4-C5	-5.83	125.69	128.60
26	14	793	A	N1-C6-N6	5.83	122.10	118.60
26	1H	1662	C	C2-N3-C4	-5.82	116.99	119.90
26	1H	1799	G	N1-C6-O6	-5.82	116.41	119.90
26	1H	2352	A	C8-N9-C4	5.82	108.13	105.80
26	1H	702	G	O5'-P-OP2	-5.82	100.46	105.70
26	14	1394	U	O5'-P-OP1	-5.82	100.46	105.70
24	3K	76	A	C5-C6-N6	-5.82	119.04	123.70
26	1H	863	A	O5'-P-OP1	5.82	117.68	110.70
27	16	77	U	C2-N3-C4	-5.82	123.51	127.00
26	14	428	A	C8-N9-C4	-5.82	103.47	105.80
26	1H	186	G	C2-N3-C4	5.82	114.81	111.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	761	A	O5'-P-OP2	5.82	117.68	110.70
26	1H	784	A	N9-C4-C5	5.82	108.13	105.80
26	1H	2377	A	C2-N3-C4	-5.82	107.69	110.60
26	14	330	A	C6-C5-N7	-5.82	128.23	132.30
26	14	2070	G	N1-C6-O6	-5.82	116.41	119.90
26	1H	716	A	C8-N9-C4	-5.82	103.47	105.80
26	1H	906	G	C6-C5-N7	5.82	133.89	130.40
26	1H	1396	U	N3-C2-O2	-5.82	118.13	122.20
55	P8	28	ARG	NE-CZ-NH1	5.82	123.21	120.30
26	1H	1660	C	C2-N3-C4	-5.81	116.99	119.90
31	31	32	LEU	CA-CB-CG	5.81	128.67	115.30
26	14	768	G	OP1-P-OP2	5.81	128.32	119.60
26	14	1266	G	C8-N9-C4	5.81	108.72	106.40
26	1H	62	C	C6-N1-C2	5.81	122.62	120.30
26	1H	1136	G	C5-C6-O6	-5.81	125.11	128.60
26	1H	1954	G	C5-C6-N1	-5.81	108.59	111.50
25	4L	14	A	O4'-C1'-N9	5.81	112.85	108.20
26	14	1762	A	N7-C8-N9	5.81	116.70	113.80
1	13	1065	U	P-O3'-C3'	5.81	126.67	119.70
26	14	213	A	C8-N9-C4	5.81	108.12	105.80
26	1H	687	C	C2-N3-C4	5.81	122.80	119.90
26	14	2685	G	N3-C4-N9	-5.81	122.52	126.00
26	1H	634	C	N3-C4-N4	-5.81	113.94	118.00
26	1H	2060	A	P-O3'-C3'	5.80	126.67	119.70
26	14	470	A	O5'-P-OP1	-5.80	100.48	105.70
26	14	1772	G	C6-C5-N7	-5.80	126.92	130.40
26	14	1970	A	O5'-P-OP2	-5.80	100.48	105.70
26	1H	969	U	N3-C4-O4	5.80	123.46	119.40
26	14	2741	A	C8-N9-C4	5.80	108.12	105.80
1	13	309	G	N3-C2-N2	-5.80	115.84	119.90
26	1H	1668	A	C2-N3-C4	5.80	113.50	110.60
26	1H	1957	C	C5-C4-N4	5.80	124.26	120.20
26	1H	2506	U	C5-C6-N1	5.80	125.60	122.70
1	13	509	A	C2'-C3'-O3'	5.80	122.97	113.70
26	1H	35	G	O5'-P-OP2	-5.80	100.48	105.70
26	1H	518	G	N1-C6-O6	-5.80	116.42	119.90
26	1H	762	U	C2-N1-C1'	5.80	124.66	117.70
1	1G	20	U	O5'-P-OP2	-5.80	100.48	105.70
1	1G	337	C	C5-C6-N1	5.80	123.90	121.00
26	14	1789	A	C6-N1-C2	-5.80	115.12	118.60
3	2E	188	LEU	CA-CB-CG	5.79	128.63	115.30
26	1H	210	C	C2-N3-C4	-5.79	117.00	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	2038	G	O5'-P-OP1	5.79	117.66	110.70
26	1H	1342	A	N1-C6-N6	5.79	122.08	118.60
26	1H	1396	U	O5'-P-OP1	-5.79	100.49	105.70
26	1H	1979	C	C6-N1-C2	-5.79	117.98	120.30
26	1H	2578	G	OP2-P-O3'	5.79	117.95	105.20
26	14	2258	C	C5-C6-N1	-5.79	118.10	121.00
26	1H	1272	A	O4'-C1'-N9	5.79	112.83	108.20
26	1H	1752	C	C2-N1-C1'	-5.79	112.43	118.80
26	14	1655	A	N7-C8-N9	-5.79	110.91	113.80
26	1H	812	C	C6-N1-C2	-5.79	117.98	120.30
26	1H	1379	A	C6-C5-N7	-5.79	128.25	132.30
26	1H	1803	A	C8-N9-C4	-5.79	103.48	105.80
26	1H	2351	G	N1-C6-O6	-5.79	116.43	119.90
26	14	783	A	C5-C6-N1	-5.79	114.81	117.70
27	1J	74	U	C5-C4-O4	5.79	129.37	125.90
26	1H	2041	U	O5'-P-OP1	-5.79	100.49	105.70
26	14	2623	G	C8-N9-C4	-5.79	104.09	106.40
26	1H	2260	C	N3-C4-C5	5.78	124.21	121.90
26	1H	2611	U	C5-C4-O4	5.78	129.37	125.90
1	13	792	A	C8-N9-C4	5.78	108.11	105.80
26	1H	766	C	C5-C6-N1	-5.78	118.11	121.00
26	14	187	G	N1-C6-O6	-5.78	116.43	119.90
26	14	571	A	C5-C6-N6	-5.78	119.08	123.70
26	1H	994	C	N1-C2-O2	-5.78	115.43	118.90
26	1H	1594	G	OP1-P-O3'	5.78	117.92	105.20
26	1H	683	C	C2-N3-C4	-5.78	117.01	119.90
26	14	180	G	C8-N9-C4	5.78	108.71	106.40
26	1H	137(A)	G	N3-C2-N2	-5.78	115.86	119.90
26	1H	1306	C	C5-C6-N1	-5.78	118.11	121.00
37	78	42	SER	C-N-CA	-5.77	110.17	122.30
26	14	769	G	N7-C8-N9	-5.77	110.21	113.10
1	13	437	U	C6-N1-C2	-5.77	117.54	121.00
26	1H	837	C	N3-C4-N4	5.77	122.04	118.00
1	1G	576	G	C4-C5-C6	5.77	122.26	118.80
26	14	1161	C	C5-C6-N1	5.77	123.89	121.00
26	1H	1377	G	O5'-P-OP2	-5.77	100.51	105.70
26	14	1899	G	C8-N9-C4	-5.77	104.09	106.40
26	1H	1660	C	N3-C4-N4	-5.77	113.96	118.00
26	14	444	C	OP2-P-O3'	5.77	117.89	105.20
26	1H	944	G	N7-C8-N9	5.77	115.98	113.10
26	1H	138	G	N9-C1'-C2'	5.76	121.50	114.00
26	1H	631	A	C5-N7-C8	5.76	106.78	103.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	1622	G	N3-C2-N2	-5.76	115.86	119.90
26	14	1694	C	O5'-P-OP1	-5.76	100.51	105.70
23	2K	1	C	C6-N1-C2	-5.76	118.00	120.30
26	1H	2249	U	C4-C5-C6	-5.76	116.24	119.70
26	1H	2287	A	N1-C2-N3	5.76	132.18	129.30
1	1G	18	C	O5'-P-OP1	-5.76	100.51	105.70
26	1H	180	G	C8-N9-C4	5.76	108.70	106.40
26	1H	1021	A	C5-C6-N1	-5.76	114.82	117.70
26	1H	2392	A	C8-N9-C4	-5.76	103.50	105.80
26	1H	2741	A	N1-C6-N6	5.76	122.06	118.60
24	3L	76	A	N1-C6-N6	5.76	122.06	118.60
26	14	2075	U	OP2-P-O3'	5.76	117.88	105.20
26	1H	482	A	C8-N9-C4	-5.76	103.50	105.80
26	14	207	A	N7-C8-N9	-5.76	110.92	113.80
26	1H	1936	A	N7-C8-N9	5.76	116.68	113.80
26	1H	2586	C	N1-C2-O2	-5.76	115.44	118.90
26	1H	2761	G	N1-C2-N3	5.76	127.36	123.90
26	14	740	U	O5'-P-OP1	5.76	117.61	110.70
26	14	2513	G	C8-N9-C4	-5.76	104.10	106.40
26	1H	768	G	OP1-P-OP2	5.76	128.24	119.60
26	1H	865	C	C6-N1-C2	5.76	122.60	120.30
26	1H	1189	A	N1-C6-N6	5.76	122.05	118.60
26	1H	1623	G	N1-C6-O6	-5.76	116.45	119.90
1	1G	1128	C	C5-C6-N1	5.76	123.88	121.00
26	14	752	A	OP2-P-O3'	5.76	117.87	105.20
26	14	2581	G	O4'-C1'-N9	5.75	112.80	108.20
26	1H	189	G	C2-N3-C4	-5.75	109.02	111.90
26	1H	1300	U	N1-C2-O2	-5.75	118.77	122.80
26	1H	2766	G	C8-N9-C4	-5.75	104.10	106.40
29	11	260	ARG	NE-CZ-NH1	5.75	123.18	120.30
26	14	1372	U	N1-C2-N3	5.75	118.35	114.90
26	1H	2253	G	N1-C6-O6	5.75	123.35	119.90
26	14	773	U	C5-C6-N1	-5.75	119.83	122.70
26	14	974(A)	C	N3-C2-O2	-5.75	117.88	121.90
26	14	1241	A	O4'-C1'-N9	5.75	112.80	108.20
26	14	1315	C	C5-C4-N4	5.75	124.23	120.20
26	1H	659	C	C6-N1-C2	5.75	122.60	120.30
26	14	2776	A	P-O3'-C3'	5.75	126.60	119.70
26	1H	127	A	C5-C6-N6	-5.75	119.10	123.70
26	14	511	U	N3-C4-C5	-5.75	111.15	114.60
26	1H	664	C	C2-N3-C4	-5.75	117.03	119.90
26	1H	1382	G	N9-C4-C5	-5.75	103.10	105.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	1930	G	C4-C5-N7	-5.75	108.50	110.80
26	14	2688	U	N3-C4-O4	-5.75	115.38	119.40
26	14	1785	A	N7-C8-N9	5.75	116.67	113.80
26	14	2597	G	O5'-P-OP1	5.75	117.59	110.70
26	1H	145	G	C6-C5-N7	-5.74	126.95	130.40
26	1H	1310	G	N1-C2-N2	5.74	121.37	116.20
26	1H	1681	G	C4-C5-N7	5.74	113.10	110.80
37	78	61	ARG	NE-CZ-NH2	-5.74	117.43	120.30
1	1G	1397	C	C6-N1-C1'	-5.74	113.91	120.80
1	1G	1519	A	C8-N9-C4	-5.74	103.50	105.80
26	14	265	A	C2-N3-C4	-5.74	107.73	110.60
26	14	792	G	N1-C6-O6	-5.74	116.45	119.90
26	14	786	C	C2-N3-C4	-5.74	117.03	119.90
26	1H	1561	G	C8-N9-C4	-5.74	104.10	106.40
26	1H	2052	G	C5-C6-O6	5.74	132.04	128.60
26	1H	2497	A	C4-C5-C6	5.74	119.87	117.00
26	14	1142(A)	A	N1-C2-N3	5.74	132.17	129.30
26	1H	1204	A	N3-C4-C5	5.74	130.82	126.80
26	1H	2244	U	N1-C2-N3	5.74	118.34	114.90
26	1H	2550	G	N3-C4-C5	-5.74	125.73	128.60
27	16	80	U	N3-C2-O2	-5.74	118.18	122.20
26	14	2870	C	C6-N1-C2	-5.74	118.00	120.30
1	13	760	G	N1-C6-O6	5.74	123.34	119.90
1	13	1468	A	C5-C6-N1	5.74	120.57	117.70
26	1H	839	U	C5-C4-O4	5.74	129.34	125.90
26	1H	2743	C	N1-C2-O2	-5.74	115.46	118.90
26	1H	2821	A	C5-C6-N6	-5.74	119.11	123.70
26	14	743	G	C8-N9-C4	5.74	108.69	106.40
26	1H	1191	G	O5'-P-OP1	-5.73	100.54	105.70
26	14	1779	U	O4'-C1'-N1	5.73	112.79	108.20
26	14	2287	A	C8-N9-C4	5.73	108.09	105.80
26	1H	463	G	C8-N9-C4	5.73	108.69	106.40
26	1H	1673	U	C2-N3-C4	-5.73	123.56	127.00
1	1G	1200	C	N1-C2-O2	5.73	122.34	118.90
26	14	1558	A	P-O3'-C3'	5.73	126.58	119.70
26	1H	1310	G	C5-C6-O6	-5.73	125.16	128.60
26	14	2323	G	C8-N9-C4	5.73	108.69	106.40
26	1H	1204	A	N1-C6-N6	5.73	122.04	118.60
26	14	1209	G	OP1-P-OP2	5.73	128.19	119.60
26	1H	130	C	N3-C4-C5	5.72	124.19	121.90
26	1H	244	A	N1-C6-N6	5.72	122.03	118.60
26	1H	1357	U	O5'-P-OP2	-5.72	100.55	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	2688	U	C4-C5-C6	5.72	123.14	119.70
26	1H	59	U	N3-C4-C5	-5.72	111.17	114.60
26	1H	1365	A	C5-C6-N6	5.72	128.28	123.70
26	1H	1665	A	O5'-P-OP1	-5.72	100.55	105.70
26	1H	1777	U	C4-C5-C6	5.72	123.13	119.70
26	14	828	U	N3-C4-C5	-5.72	111.17	114.60
26	14	2582	G	C6-C5-N7	-5.72	126.97	130.40
1	13	505	G	C5-N7-C8	-5.72	101.44	104.30
26	1H	399	G	N1-C6-O6	-5.72	116.47	119.90
26	1H	966	G	N3-C2-N2	5.72	123.90	119.90
26	1H	270(L)	U	C5-C6-N1	5.72	125.56	122.70
26	14	2413	G	C5-C6-O6	-5.72	125.17	128.60
1	13	1486	G	O5'-P-OP2	-5.71	100.56	105.70
26	1H	117	G	N3-C4-N9	5.71	129.43	126.00
26	1H	859	G	C4-N9-C1'	-5.71	119.07	126.50
26	14	1790	C	OP1-P-O3'	5.71	117.77	105.20
26	1H	2067	G	N3-C4-N9	-5.71	122.57	126.00
26	1H	2073	C	C4-C5-C6	5.71	120.26	117.40
1	13	560	U	P-O3'-C3'	5.71	126.55	119.70
26	1H	1307	A	N1-C6-N6	5.71	122.03	118.60
26	14	1527	G	N3-C2-N2	-5.71	115.90	119.90
26	14	1351	C	C5-C6-N1	-5.71	118.15	121.00
26	14	1585	C	N3-C2-O2	-5.71	117.90	121.90
26	14	577	G	C5-C6-O6	-5.71	125.18	128.60
26	14	1427	A	C6-N1-C2	-5.71	115.18	118.60
26	14	503	A	N9-C4-C5	5.70	108.08	105.80
26	14	788	A	C5-C6-N6	-5.70	119.14	123.70
26	14	954	G	N1-C6-O6	-5.70	116.48	119.90
26	14	1297	C	OP2-P-O3'	-5.70	92.65	105.20
26	1H	1888	G	N3-C4-C5	-5.70	125.75	128.60
1	13	169	C	C6-N1-C2	-5.70	118.02	120.30
1	13	703	G	C5-C6-O6	-5.70	125.18	128.60
26	1H	2367	G	N7-C8-N9	5.70	115.95	113.10
1	1G	503	C	N3-C4-C5	-5.70	119.62	121.90
26	14	137	C	N3-C4-C5	-5.70	119.62	121.90
26	14	2544	G	C5-C6-O6	-5.70	125.18	128.60
26	14	2767	C	C6-N1-C2	-5.70	118.02	120.30
26	1H	1324	G	N1-C2-N2	5.69	121.33	116.20
26	1H	758	C	N3-C4-C5	5.69	124.18	121.90
26	1H	1253	A	C5-N7-C8	5.69	106.75	103.90
27	1J	114	G	C8-N9-C4	5.69	108.68	106.40
26	1H	961	C	O4'-C1'-N1	5.69	112.75	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	2572	A	N7-C8-N9	-5.69	110.95	113.80
26	14	1937	A	O4'-C1'-N9	5.69	112.75	108.20
26	14	1985	G	O5'-P-OP2	-5.69	100.58	105.70
1	13	108	G	N9-C4-C5	-5.69	103.12	105.40
26	1H	452	G	N1-C6-O6	-5.69	116.49	119.90
31	39	125	LEU	CA-CB-CG	5.69	128.38	115.30
1	1G	1528	U	C6-N1-C2	5.69	124.41	121.00
26	14	774	A	O5'-P-OP2	-5.69	100.58	105.70
26	14	2057	A	C5-C6-N6	-5.69	119.15	123.70
37	78	45	LEU	CA-CB-CG	5.69	128.38	115.30
26	14	1349	A	C5-N7-C8	-5.69	101.06	103.90
26	14	2063	C	OP2-P-O3'	5.69	117.71	105.20
26	14	2325	G	N3-C2-N2	-5.69	115.92	119.90
26	1H	80	G	C8-N9-C4	-5.68	104.13	106.40
26	1H	110	G	OP1-P-OP2	5.68	128.13	119.60
26	1H	240	G	N1-C6-O6	5.68	123.31	119.90
26	14	1728	G	N3-C4-N9	5.68	129.41	126.00
26	14	1762	A	C4-C5-C6	5.68	119.84	117.00
26	14	1764	G	O5'-P-OP2	-5.68	100.58	105.70
26	1H	513	A	C8-N9-C4	-5.68	103.53	105.80
26	1H	2434	A	OP2-P-O3'	5.68	117.70	105.20
53	N8	16	ARG	NE-CZ-NH1	5.68	123.14	120.30
26	1H	573	G	C2-N3-C4	5.68	114.74	111.90
26	14	769	G	C8-N9-C4	5.68	108.67	106.40
26	14	1313	U	N1-C2-O2	-5.68	118.82	122.80
26	1H	2622	C	OP2-P-O3'	5.68	117.69	105.20
26	1H	2697	G	C5-C6-O6	5.68	132.01	128.60
26	14	2238	G	P-O3'-C3'	5.68	126.52	119.70
26	1H	116	C	OP2-P-O3'	5.68	117.69	105.20
26	1H	1996	C	C5-C6-N1	-5.68	118.16	121.00
26	1H	958	U	C6-N1-C2	-5.68	117.59	121.00
26	14	1781	C	C6-N1-C1'	-5.68	113.99	120.80
26	1H	621	A	O4'-C1'-N9	5.67	112.74	108.20
26	14	1964	G	N1-C6-O6	-5.67	116.50	119.90
26	1H	614	U	C6-N1-C2	-5.67	117.60	121.00
26	1H	937	U	O5'-P-OP1	5.67	117.51	110.70
26	1H	1776	G	OP1-P-O3'	5.67	117.68	105.20
39	98	75	LEU	CA-CB-CG	5.67	128.34	115.30
26	14	581	C	N3-C4-N4	-5.67	114.03	118.00
26	1H	207	A	C4-C5-N7	5.67	113.53	110.70
26	1H	664	C	N3-C4-N4	-5.67	114.03	118.00
26	1H	2458	G	N1-C6-O6	5.67	123.30	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	2367	G	C8-N9-C4	-5.67	104.13	106.40
26	1H	2442	C	C2-N3-C4	-5.67	117.07	119.90
26	14	824	A	C8-N9-C4	5.67	108.07	105.80
26	14	1824	G	O5'-P-OP2	-5.67	100.60	105.70
26	14	2032	G	C8-N9-C4	5.67	108.67	106.40
26	14	2256	G	O5'-P-OP2	-5.67	100.60	105.70
26	14	2598	A	N9-C4-C5	-5.67	103.53	105.80
26	14	2763	G	N3-C4-C5	-5.67	125.77	128.60
26	1H	144	C	C2-N3-C4	-5.67	117.07	119.90
26	1H	399	G	C5-C6-N1	5.67	114.33	111.50
26	1H	953	A	O5'-P-OP1	-5.67	100.60	105.70
26	1H	2392	A	C5-C6-N1	-5.67	114.87	117.70
26	1H	2493	U	C5-C6-N1	-5.67	119.87	122.70
26	14	933	A	N7-C8-N9	5.67	116.63	113.80
26	14	1204	A	O4'-C1'-N9	5.67	112.73	108.20
26	14	675	A	C8-N9-C4	5.67	108.07	105.80
26	14	2629	A	C2-N3-C4	5.67	113.43	110.60
26	1H	2392	A	OP1-P-OP2	-5.66	111.11	119.60
29	19	272	ALA	C-N-CA	5.66	135.86	121.70
26	1H	2266	A	N1-C2-N3	5.66	132.13	129.30
26	1H	560	C	O5'-P-OP2	5.66	117.49	110.70
26	1H	866	A	C4-N9-C1'	5.66	136.49	126.30
26	1H	1936	A	O4'-C1'-N9	5.66	112.73	108.20
27	1J	22	U	C6-N1-C2	-5.66	117.60	121.00
26	1H	180	G	N3-C2-N2	5.66	123.86	119.90
26	1H	770	G	N1-C6-O6	5.66	123.30	119.90
26	1H	1415	U	C5-C4-O4	5.66	129.30	125.90
26	1H	753	C	N1-C2-O2	5.66	122.29	118.90
26	1H	2016	U	C5-C6-N1	-5.66	119.87	122.70
26	14	2030	A	OP1-P-OP2	5.66	128.09	119.60
26	14	2067	G	C4-C5-N7	-5.66	108.54	110.80
26	14	740	U	O5'-P-OP2	-5.66	100.61	105.70
26	14	1271	G	C8-N9-C4	5.66	108.66	106.40
26	1H	1899	G	OP2-P-O3'	5.65	117.64	105.20
40	65	101	LEU	CA-CB-CG	5.65	128.30	115.30
26	1H	1437	C	N1-C2-O2	5.65	122.29	118.90
26	1H	1761	C	N1-C2-O2	-5.65	115.51	118.90
26	1H	1804	C	N3-C4-N4	-5.65	114.04	118.00
26	1H	963	U	O5'-P-OP2	5.65	117.48	110.70
26	1H	2519	U	N1-C2-O2	-5.65	118.84	122.80
26	14	678	C	N3-C4-C5	5.65	124.16	121.90
26	1H	1081	U	P-O3'-C3'	5.65	126.48	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	2572	A	C5-C6-N6	-5.65	119.18	123.70
26	14	1661	G	C8-N9-C4	5.65	108.66	106.40
26	14	1780	A	O5'-P-OP1	5.65	117.48	110.70
26	1H	1564	C	N3-C4-N4	-5.65	114.05	118.00
26	1H	1752	C	N1-C2-O2	-5.65	115.51	118.90
26	1H	2258	C	C5-C4-N4	-5.64	116.25	120.20
26	1H	2497	A	C6-N1-C2	-5.64	115.21	118.60
38	88	24	GLY	N-CA-C	-5.64	98.99	113.10
1	13	527	G	N1-C6-O6	-5.64	116.52	119.90
26	1H	1446	C	C6-N1-C2	-5.64	118.04	120.30
26	1H	2848	G	O4'-C1'-N9	5.64	112.72	108.20
1	1G	115	G	P-O3'-C3'	5.64	126.47	119.70
26	1H	2712	U	P-O3'-C3'	5.64	126.47	119.70
26	1H	1241	A	N3-C4-C5	5.64	130.75	126.80
26	14	2436	G	N3-C2-N2	-5.64	115.95	119.90
26	1H	944	G	C5-C6-N1	-5.64	108.68	111.50
26	14	1142	U	C6-N1-C1'	-5.64	113.31	121.20
1	13	1506	U	C5-C4-O4	-5.64	122.52	125.90
26	1H	2584	U	N1-C2-O2	5.64	126.75	122.80
41	75	13	ARG	N-CA-C	-5.64	95.78	111.00
1	13	11	G	OP1-P-O3'	5.63	117.59	105.20
1	13	326	G	C5-C6-O6	5.63	131.98	128.60
1	13	1319	A	O5'-P-OP2	-5.63	100.63	105.70
26	1H	508	G	N3-C2-N2	5.63	123.84	119.90
26	1H	622	G	C8-N9-C4	5.63	108.65	106.40
26	1H	1636	C	N1-C2-O2	-5.63	115.52	118.90
26	1H	2665	A	C2-N3-C4	-5.63	107.78	110.60
27	16	50	G	OP2-P-O3'	5.63	117.59	105.20
26	14	2779	U	N3-C4-O4	-5.63	115.45	119.40
26	1H	2555	U	O5'-P-OP1	-5.63	100.63	105.70
26	14	1332	G	C4-N9-C1'	5.63	133.82	126.50
1	1G	244	U	C5-C4-O4	-5.63	122.52	125.90
26	1H	1993	U	N1-C2-O2	-5.63	118.86	122.80
26	1H	1804	C	N3-C2-O2	-5.63	117.96	121.90
26	14	74	A	C4-C5-N7	5.63	113.51	110.70
26	14	621	A	N1-C6-N6	5.63	121.98	118.60
26	14	2056	G	OP1-P-O3'	5.63	117.58	105.20
1	13	1286	A	N7-C8-N9	5.62	116.61	113.80
26	1H	951	C	N3-C2-O2	-5.62	117.96	121.90
26	1H	1204	A	C6-C5-N7	-5.62	128.36	132.30
36	68	70	LYS	C-N-CA	5.62	135.76	121.70
1	13	913	A	P-O3'-C3'	5.62	126.45	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	913	U	O5'-P-OP2	-5.62	100.64	105.70
26	1H	141(A)	C	C6-N1-C2	5.62	122.55	120.30
26	1H	537	C	O5'-P-OP1	5.62	117.45	110.70
26	1H	2065	C	O5'-P-OP1	-5.62	100.64	105.70
26	1H	2380	C	C6-N1-C2	5.62	122.55	120.30
1	1G	180	U	C5-C6-N1	5.62	125.51	122.70
26	1H	258	G	O5'-P-OP2	-5.62	100.64	105.70
26	1H	1428	C	C6-N1-C2	5.62	122.55	120.30
26	14	574	C	N3-C4-N4	-5.62	114.07	118.00
26	1H	1367	A	C2-N3-C4	-5.62	107.79	110.60
26	14	2401	U	C6-N1-C2	-5.62	117.63	121.00
26	14	2065	C	N1-C2-O2	5.61	122.27	118.90
26	14	2589	A	N9-C4-C5	-5.61	103.56	105.80
26	1H	1375	C	OP1-P-O3'	5.61	117.55	105.20
26	14	2685	G	C4-C5-N7	-5.61	108.56	110.80
26	14	2707	G	C5-C6-N1	5.61	114.31	111.50
26	1H	240	G	C8-N9-C4	5.61	108.64	106.40
26	1H	330	A	N1-C2-N3	5.61	132.10	129.30
26	1H	1196	C	C6-N1-C2	5.61	122.54	120.30
26	1H	2578	G	C8-N9-C4	5.61	108.64	106.40
26	14	194	G	N1-C6-O6	5.61	123.27	119.90
26	14	1313	U	C5-C6-N1	5.61	125.51	122.70
26	14	1678	G	N9-C4-C5	5.61	107.64	105.40
1	13	652	U	O5'-P-OP1	-5.61	100.65	105.70
26	14	2554	U	O5'-P-OP2	5.61	117.43	110.70
1	13	328	C	O5'-P-OP1	-5.61	100.66	105.70
1	13	759	A	O5'-P-OP2	-5.61	100.66	105.70
26	1H	1800	C	N1-C2-N3	5.61	123.12	119.20
26	1H	2711	A	OP1-P-O3'	5.61	117.53	105.20
56	Q8	47	LYS	N-CA-C	-5.61	95.87	111.00
1	1G	576	G	N3-C4-N9	5.61	129.36	126.00
1	1G	700	G	N1-C6-O6	-5.61	116.54	119.90
26	1H	122	G	C8-N9-C4	5.60	108.64	106.40
26	1H	734	A	C2-N3-C4	-5.60	107.80	110.60
26	1H	1351	C	OP1-P-O3'	5.60	117.53	105.20
26	1H	2623	G	N7-C8-N9	5.60	115.90	113.10
44	A5	8	ARG	NE-CZ-NH1	5.60	123.10	120.30
1	13	1517	G	O5'-P-OP2	-5.60	100.66	105.70
26	1H	688	U	C5-C6-N1	-5.60	119.90	122.70
26	1H	860	U	N1-C2-N3	5.60	118.26	114.90
26	1H	2303	G	OP1-P-O3'	5.60	117.52	105.20
26	14	528	A	C5-N7-C8	-5.60	101.10	103.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	1574	C	OP2-P-O3'	5.60	117.53	105.20
1	13	1158	C	N3-C2-O2	-5.60	117.98	121.90
26	1H	694	U	O5'-P-OP1	5.60	117.42	110.70
26	1H	1825	A	C2-N3-C4	5.60	113.40	110.60
26	1H	2329	G	C8-N9-C4	5.60	108.64	106.40
26	1H	2741	A	N9-C4-C5	-5.60	103.56	105.80
26	14	680	G	O5'-P-OP1	-5.60	100.66	105.70
26	14	2082	A	C5-C6-N6	-5.60	119.22	123.70
26	14	2415	G	OP1-P-O3'	5.60	117.52	105.20
26	1H	1993	U	C5-C6-N1	-5.60	119.90	122.70
26	1H	2438	U	C4-C5-C6	5.60	123.06	119.70
26	1H	2620	C	N3-C4-N4	-5.60	114.08	118.00
1	1G	271	C	C6-N1-C2	-5.60	118.06	120.30
26	1H	599	G	N3-C2-N2	5.59	123.81	119.90
26	1H	1984	G	N1-C6-O6	-5.59	116.54	119.90
26	1H	1997	G	O5'-P-OP2	-5.59	100.67	105.70
26	14	1930	G	C4-N9-C1'	-5.59	119.23	126.50
26	1H	405	U	C2-N1-C1'	5.59	124.41	117.70
26	1H	969	U	C5-C4-O4	-5.59	122.55	125.90
26	1H	1325	G	OP1-P-OP2	-5.59	111.21	119.60
26	1H	1379	A	C8-N9-C4	-5.59	103.56	105.80
26	14	2053	G	N1-C6-O6	5.59	123.25	119.90
26	1H	302	C	N3-C2-O2	-5.59	117.99	121.90
26	1H	335	C	C2-N3-C4	5.59	122.69	119.90
26	1H	1261	C	C6-N1-C2	5.59	122.54	120.30
26	1H	1574	C	C2-N3-C4	-5.59	117.11	119.90
26	1H	2321	G	O5'-P-OP1	5.59	117.41	110.70
26	14	2554	U	O5'-P-OP1	-5.59	100.67	105.70
26	1H	982	C	C6-N1-C2	-5.59	118.06	120.30
26	1H	2645	G	N3-C4-C5	5.59	131.39	128.60
26	14	1941	C	O5'-P-OP1	-5.59	100.67	105.70
26	1H	936	C	C5-C6-N1	-5.59	118.21	121.00
46	C5	103	GLY	N-CA-C	5.59	127.07	113.10
2	1E	158	LEU	CA-CB-CG	5.58	128.15	115.30
26	1H	1605	C	O5'-P-OP1	-5.58	100.67	105.70
26	1H	124	G	N1-C6-O6	5.58	123.25	119.90
26	1H	135	G	C5-N7-C8	5.58	107.09	104.30
26	1H	236	C	C5-C6-N1	-5.58	118.21	121.00
26	1H	687	C	C6-N1-C2	-5.58	118.07	120.30
26	14	2689	U	P-O3'-C3'	5.58	126.40	119.70
26	1H	929	G	C8-N9-C1'	-5.58	119.75	127.00
26	1H	1021	A	C6-C5-N7	-5.58	128.39	132.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	1141	U	O4'-C1'-N1	5.58	112.67	108.20
26	14	2373	G	OP1-P-OP2	5.58	127.97	119.60
1	1G	898	G	N3-C4-C5	5.58	131.39	128.60
27	1J	11	C	N1-C2-O2	5.58	122.25	118.90
1	13	35	G	C5-C6-N1	-5.58	108.71	111.50
1	13	974	A	C4-N9-C1'	5.58	136.34	126.30
26	1H	141	A	O4'-C1'-N9	5.58	112.66	108.20
26	1H	2257	U	C2-N3-C4	-5.58	123.65	127.00
26	1H	238	C	C5-C6-N1	-5.58	118.21	121.00
26	1H	673	C	OP1-P-OP2	-5.58	111.23	119.60
26	14	71	A	C6-C5-N7	-5.58	128.40	132.30
1	13	1299	A	C2-N3-C4	-5.58	107.81	110.60
26	1H	238	C	C4-C5-C6	5.58	120.19	117.40
26	1H	2031	A	C5-C6-N6	-5.58	119.24	123.70
26	14	2581	G	N1-C2-N2	-5.58	111.18	116.20
26	14	2607	G	O5'-P-OP1	5.58	117.39	110.70
1	13	481	G	C6-C5-N7	-5.57	127.06	130.40
26	14	130	C	N3-C4-C5	5.57	124.13	121.90
26	14	797	C	N3-C2-O2	5.57	125.80	121.90
26	1H	1332	G	O4'-C1'-N9	-5.57	103.74	108.20
26	1H	1669	A	C5-C6-N1	5.57	120.49	117.70
26	14	2565	A	C8-N9-C4	5.57	108.03	105.80
1	13	768	A	C6-N1-C2	-5.57	115.26	118.60
26	1H	335	C	C5-C6-N1	5.57	123.79	121.00
26	1H	1610	A	C6-C5-N7	-5.57	128.40	132.30
26	1H	2276	G	N3-C2-N2	-5.57	116.00	119.90
26	1H	2819	G	N1-C6-O6	5.57	123.24	119.90
27	16	6	C	N3-C4-N4	5.57	121.90	118.00
26	14	127	A	C5-C6-N6	-5.57	119.24	123.70
26	14	2003	G	N1-C6-O6	5.57	123.24	119.90
26	1H	621	A	N3-C4-N9	-5.57	122.94	127.40
26	1H	2064	C	N3-C2-O2	-5.57	118.00	121.90
1	13	690	G	N1-C6-O6	5.57	123.24	119.90
26	1H	127	A	N1-C6-N6	5.57	121.94	118.60
26	1H	1413	G	C8-N9-C4	-5.57	104.17	106.40
26	14	1351	C	C2-N3-C4	-5.57	117.12	119.90
26	14	2512	C	N3-C4-C5	5.57	124.13	121.90
26	1H	1573	G	OP2-P-O3'	5.57	117.44	105.20
26	1H	2064	C	OP1-P-O3'	5.57	117.44	105.20
1	1G	481	G	C8-N9-C1'	-5.57	119.77	127.00
26	14	668	G	N3-C4-C5	5.57	131.38	128.60
26	1H	1261	C	N3-C4-C5	5.56	124.12	121.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	420	C	C6-N1-C2	5.56	122.53	120.30
26	1H	1761	C	N3-C4-N4	5.56	121.89	118.00
26	1H	2469	A	N1-C6-N6	5.56	121.94	118.60
26	14	2281	C	O5'-P-OP2	5.56	117.38	110.70
1	13	802	A	C4-C5-N7	5.56	113.48	110.70
26	1H	2743	C	C2-N3-C4	-5.56	117.12	119.90
1	1G	413	G	C4-C5-N7	-5.56	108.58	110.80
26	14	1210	A	C5-N7-C8	-5.56	101.12	103.90
1	13	1279	A	C6-C5-N7	-5.56	128.41	132.30
26	1H	52	A	N1-C2-N3	-5.56	126.52	129.30
26	1H	500	G	O5'-P-OP1	-5.56	100.70	105.70
26	1H	2869	G	C8-N9-C4	-5.56	104.18	106.40
1	1G	906	G	C5-C6-O6	-5.56	125.26	128.60
26	14	2490	G	O4'-C1'-N9	5.56	112.65	108.20
26	1H	223	A	O5'-P-OP2	-5.56	100.70	105.70
27	16	98	G	C8-N9-C4	5.56	108.62	106.40
26	14	2276	G	O5'-P-OP1	-5.56	100.70	105.70
26	1H	1489	U	C5-C4-O4	5.56	129.23	125.90
26	14	214	G	C8-N9-C4	-5.55	104.18	106.40
26	14	1614	A	C5-C6-N1	-5.55	114.92	117.70
26	1H	1840	G	N3-C2-N2	-5.55	116.01	119.90
26	1H	2355	C	N3-C4-C5	5.55	124.12	121.90
26	14	1166	C	C6-N1-C2	-5.55	118.08	120.30
26	1H	463	G	N7-C8-N9	-5.55	110.32	113.10
26	1H	508	G	N3-C4-N9	5.55	129.33	126.00
26	1H	918	A	O5'-P-OP2	5.55	117.36	110.70
26	1H	2234	G	C8-N9-C4	5.55	108.62	106.40
26	1H	2544	G	C8-N9-C4	5.55	108.62	106.40
26	1H	915	C	OP1-P-OP2	-5.55	111.28	119.60
26	1H	1899	G	C5-N7-C8	-5.55	101.53	104.30
26	1H	2412	A	O5'-P-OP2	-5.55	100.70	105.70
26	14	945	A	N9-C1'-C2'	5.55	121.21	114.00
1	13	1158	C	C6-N1-C2	-5.55	118.08	120.30
26	1H	1939	U	N3-C4-C5	5.55	117.93	114.60
26	14	945	A	O4'-C1'-N9	5.55	112.64	108.20
26	1H	674	G	N9-C4-C5	-5.55	103.18	105.40
26	1H	1799	G	P-O3'-C3'	5.55	126.36	119.70
26	14	1918	A	N9-C4-C5	-5.55	103.58	105.80
26	1H	109	G	C5-C6-O6	5.54	131.93	128.60
26	1H	2469	A	C2-N3-C4	-5.54	107.83	110.60
26	14	1644	C	N1-C2-O2	5.54	122.22	118.90
26	14	2688	U	N1-C2-N3	5.54	118.22	114.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	2518	A	N7-C8-N9	5.54	116.57	113.80
26	1H	122	G	N7-C8-N9	-5.54	110.33	113.10
26	1H	129	C	C4-C5-C6	5.54	120.17	117.40
26	1H	683	C	C5-C4-N4	-5.54	116.32	120.20
26	1H	1032	A	N1-C2-N3	-5.54	126.53	129.30
26	1H	2276	G	C4-C5-N7	-5.54	108.58	110.80
37	78	18	ARG	NE-CZ-NH2	5.54	123.07	120.30
26	14	929	G	C4-C5-N7	5.54	113.02	110.80
1	13	1336	C	C5-C6-N1	5.54	123.77	121.00
26	1H	697	C	C5-C4-N4	-5.54	116.32	120.20
26	1H	2210	G	OP2-P-O3'	5.54	117.38	105.20
26	1H	1678	G	N3-C2-N2	-5.54	116.03	119.90
26	14	680	G	C6-C5-N7	-5.54	127.08	130.40
26	14	226	G	O4'-C1'-N9	5.53	112.63	108.20
1	13	67	C	C6-N1-C2	-5.53	118.09	120.30
26	1H	1900	A	OP1-P-OP2	-5.53	111.31	119.60
43	D8	18	LEU	CA-CB-CG	5.53	128.02	115.30
50	K8	3	LEU	C-N-CA	5.53	135.53	121.70
26	14	2576	G	OP1-P-OP2	-5.53	111.31	119.60
26	1H	1520	U	C5-C4-O4	5.53	129.22	125.90
26	14	1007	C	C6-N1-C2	-5.53	118.09	120.30
1	13	47	C	C4-C5-C6	5.53	120.16	117.40
1	13	336	C	N3-C2-O2	5.53	125.77	121.90
26	1H	74	A	N3-C4-C5	5.53	130.67	126.80
26	1H	2331	G	C8-N9-C4	5.53	108.61	106.40
1	1G	1405	G	C8-N9-C4	5.53	108.61	106.40
26	14	2307	G	O4'-C1'-N9	5.53	112.62	108.20
26	1H	1426	G	C5-C6-O6	-5.53	125.28	128.60
26	14	1444(A)	A	N1-C6-N6	-5.53	115.28	118.60
26	1H	209	C	C5-C6-N1	-5.52	118.24	121.00
26	1H	579	G	N3-C2-N2	-5.52	116.03	119.90
26	1H	1299	G	O5'-P-OP2	5.52	117.33	110.70
26	1H	1700	A	OP1-P-OP2	5.52	127.89	119.60
26	1H	1773	A	N9-C1'-C2'	-5.52	105.93	112.00
26	1H	2094	G	O5'-P-OP2	-5.52	100.73	105.70
26	14	774	A	C6-N1-C2	5.52	121.91	118.60
26	1H	2338	G	O5'-P-OP1	-5.52	100.73	105.70
26	14	726	G	O4'-C1'-N9	5.52	112.62	108.20
26	14	2699	C	C6-N1-C2	5.52	122.51	120.30
26	14	2871	C	O5'-P-OP2	-5.52	100.73	105.70
41	75	6	LEU	N-CA-C	-5.52	96.09	111.00
26	1H	716	A	N7-C8-N9	5.52	116.56	113.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	2574	G	C5-C6-O6	-5.52	125.29	128.60
26	14	779	U	N3-C4-O4	5.52	123.26	119.40
26	14	2700	C	N3-C4-C5	5.52	124.11	121.90
26	1H	194	G	N1-C6-O6	5.52	123.21	119.90
26	1H	816	C	N3-C4-C5	-5.52	119.69	121.90
26	1H	1571	A	C5-C6-N6	-5.52	119.28	123.70
26	1H	2376	A	C8-N9-C4	5.52	108.01	105.80
26	14	1780	A	N9-C4-C5	5.52	108.01	105.80
23	2K	24	C	C2-N3-C4	-5.51	117.14	119.90
26	1H	629	G	N1-C6-O6	-5.51	116.59	119.90
26	1H	1616	A	OP1-P-O3'	5.51	117.33	105.20
1	1G	913	A	P-O3'-C3'	5.51	126.31	119.70
1	13	121	C	C2-N1-C1'	5.51	124.86	118.80
26	1H	2269	A	C2-N3-C4	-5.51	107.84	110.60
26	1H	2590	A	C2-N3-C4	-5.51	107.84	110.60
26	14	2328	A	C6-N1-C2	-5.51	115.29	118.60
26	14	2427	C	C5-C4-N4	-5.51	116.34	120.20
26	1H	760	G	N9-C4-C5	-5.51	103.20	105.40
26	14	774	A	N9-C4-C5	-5.51	103.60	105.80
26	1H	1363	C	C5-C6-N1	-5.51	118.25	121.00
26	1H	2262	U	O5'-P-OP1	5.51	117.31	110.70
26	1H	2592	G	OP2-P-O3'	5.51	117.32	105.20
26	1H	226	G	N1-C6-O6	5.51	123.20	119.90
26	1H	698	C	O5'-P-OP2	-5.51	100.74	105.70
26	1H	813	U	OP1-P-OP2	5.51	127.86	119.60
26	1H	1407	C	OP1-P-O3'	5.51	117.31	105.20
26	14	1166	C	N3-C4-C5	-5.51	119.70	121.90
26	14	1365	A	C8-N9-C4	-5.51	103.60	105.80
26	14	2082	A	N1-C6-N6	5.51	121.90	118.60
26	1H	2024	G	O5'-P-OP1	-5.50	100.75	105.70
33	51	153	LYS	C-N-CA	5.50	145.12	122.00
26	14	1245	G	C5-C6-O6	-5.50	125.30	128.60
26	14	2385	C	C2-N3-C4	-5.50	117.15	119.90
26	1H	205	G	N3-C2-N2	5.50	123.75	119.90
26	1H	2308	G	C6-N1-C2	5.50	128.40	125.10
1	1G	320	C	C6-N1-C2	5.50	122.50	120.30
26	14	1475	G	C8-N9-C4	-5.50	104.20	106.40
27	1J	114	G	N3-C4-C5	5.50	131.35	128.60
26	14	479	A	N1-C6-N6	-5.50	115.30	118.60
26	14	1585	C	C5-C6-N1	5.50	123.75	121.00
26	14	2448	A	C6-N1-C2	-5.50	115.30	118.60
1	13	108	G	C4-N9-C1'	5.50	133.65	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	1786	A	OP1-P-O3'	5.50	117.30	105.20
26	1H	2446	G	C5-N7-C8	-5.50	101.55	104.30
26	14	2227	A	C5-N7-C8	-5.50	101.15	103.90
1	1G	59	A	N1-C6-N6	5.50	121.90	118.60
1	1G	567	G	N3-C4-N9	-5.50	122.70	126.00
26	14	308	G	C4-C5-N7	5.50	113.00	110.80
26	14	2429	G	OP2-P-O3'	5.50	117.29	105.20
26	14	2516	G	OP2-P-O3'	5.50	117.29	105.20
26	14	36	G	OP2-P-O3'	5.50	117.29	105.20
26	1H	21	A	OP2-P-O3'	5.49	117.28	105.20
34	69	131	LYS	C-N-CD	-5.49	108.51	120.60
23	2K	6	G	N1-C6-O6	5.49	123.19	119.90
26	1H	2525	G	C8-N9-C4	5.49	108.60	106.40
40	65	26	LEU	CA-CB-CG	5.49	127.93	115.30
26	1H	1779	U	O5'-P-OP2	-5.49	100.76	105.70
1	1G	812	C	N1-C2-O2	-5.49	115.61	118.90
26	14	470	A	C4-C5-N7	5.49	113.45	110.70
1	13	703	G	C8-N9-C1'	-5.49	119.86	127.00
26	1H	682	G	C8-N9-C1'	-5.49	119.87	127.00
26	1H	703	U	N3-C4-O4	-5.49	115.56	119.40
26	1H	1835	G	C8-N9-C4	-5.49	104.20	106.40
26	1H	1136	G	N1-C2-N2	5.49	121.14	116.20
26	1H	1900	A	C5'-C4'-O4'	-5.49	102.52	109.10
26	1H	2419	U	OP1-P-OP2	-5.49	111.37	119.60
26	1H	2751	G	N3-C4-C5	5.49	131.34	128.60
27	16	29	A	C8-N9-C4	-5.49	103.61	105.80
26	1H	1210	A	C2-N3-C4	-5.48	107.86	110.60
1	13	943	U	N3-C2-O2	5.48	126.04	122.20
26	1H	2585	U	N1-C2-O2	5.48	126.64	122.80
1	1G	810	C	N1-C2-O2	5.48	122.19	118.90
1	1G	893	C	C6-N1-C2	5.48	122.49	120.30
26	14	2256	G	N3-C2-N2	5.48	123.74	119.90
1	13	1455	G	N3-C4-C5	5.48	131.34	128.60
26	1H	250	G	N7-C8-N9	5.48	115.84	113.10
26	1H	1801	G	C5-C6-O6	-5.48	125.31	128.60
27	16	44	G	C4-C5-N7	-5.48	108.61	110.80
26	14	118	A	N1-C6-N6	-5.48	115.31	118.60
26	14	563	G	N1-C6-O6	-5.48	116.61	119.90
26	14	2457	U	OP2-P-O3'	5.48	117.26	105.20
1	13	1446	A	O5'-P-OP1	5.48	117.27	110.70
27	16	98	G	C5-C6-O6	-5.48	125.31	128.60
1	1G	108	G	N3-C2-N2	5.48	123.73	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	1986	A	N7-C8-N9	5.48	116.54	113.80
26	14	707	G	N1-C6-O6	5.48	123.19	119.90
26	14	2437	U	C5-C4-O4	5.48	129.19	125.90
26	14	2776	A	N7-C8-N9	5.48	116.54	113.80
26	1H	1568	G	OP1-P-OP2	-5.48	111.39	119.60
26	14	1780	A	O5'-P-OP2	-5.48	100.77	105.70
26	14	1933	G	OP1-P-OP2	5.48	127.81	119.60
26	1H	1624	G	N3-C2-N2	5.47	123.73	119.90
26	14	835	A	O5'-P-OP1	5.47	117.27	110.70
26	14	855	G	N7-C8-N9	5.47	115.84	113.10
1	13	800	G	C6-C5-N7	-5.47	127.12	130.40
2	12	17	PHE	C-N-CA	5.47	133.79	122.30
26	14	742	G	C5-C6-O6	5.47	131.88	128.60
26	14	822	U	N1-C2-N3	5.47	118.18	114.90
26	14	2008	C	OP2-P-O3'	5.47	117.24	105.20
26	14	2607	G	N3-C2-N2	5.47	123.73	119.90
26	14	1556	C	O5'-P-OP1	-5.47	100.78	105.70
26	1H	795	C	C4-C5-C6	5.47	120.14	117.40
26	1H	1322	A	OP2-P-O3'	5.47	117.23	105.20
26	14	758	C	O5'-P-OP2	-5.47	100.78	105.70
26	14	773	U	C2-N3-C4	-5.47	123.72	127.00
1	1G	579	G	C4-N9-C1'	5.47	133.61	126.50
1	1G	1401	G	C4-C5-C6	5.47	122.08	118.80
26	1H	667	U	C5-C4-O4	-5.47	122.62	125.90
26	14	58	G	C6-C5-N7	-5.47	127.12	130.40
26	14	668	G	N7-C8-N9	-5.47	110.37	113.10
26	14	2490	G	C6-C5-N7	-5.47	127.12	130.40
26	1H	755	C	C4-C5-C6	5.46	120.13	117.40
26	1H	1210	A	C4-C5-C6	5.46	119.73	117.00
26	1H	1280	G	OP1-P-OP2	-5.46	111.41	119.60
26	14	1914	C	N3-C2-O2	-5.46	118.08	121.90
50	K8	3	LEU	CA-C-N	5.46	129.22	117.20
1	13	798	G	C5-C6-N1	-5.46	108.77	111.50
26	1H	195	A	OP1-P-O3'	-5.46	93.19	105.20
26	1H	633	A	N1-C6-N6	5.46	121.88	118.60
26	1H	1971	A	O5'-P-OP2	-5.46	100.78	105.70
26	14	1314	C	C6-N1-C1'	-5.46	114.25	120.80
26	14	1385	G	O4'-C1'-N9	5.46	112.57	108.20
26	14	1899	G	C5-C6-N1	-5.46	108.77	111.50
26	14	2582	G	N1-C6-O6	5.46	123.18	119.90
1	1G	890	G	O4'-C1'-N9	5.46	112.57	108.20
24	3L	76	A	C4-C5-N7	5.46	113.43	110.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	1349	A	N1-C6-N6	5.46	121.88	118.60
26	14	681	G	N1-C2-N3	5.46	127.17	123.90
27	1J	102	G	C4-C5-N7	-5.46	108.62	110.80
26	1H	697	C	N3-C4-C5	5.46	124.08	121.90
26	1H	1761	C	C6-N1-C2	5.46	122.48	120.30
26	1H	1796	U	C5-C6-N1	-5.46	119.97	122.70
26	1H	2081	C	N1-C2-O2	5.46	122.17	118.90
26	14	801	G	C6-C5-N7	5.46	133.67	130.40
26	1H	796	C	N3-C2-O2	-5.46	118.08	121.90
27	16	100	G	N3-C4-N9	5.46	129.27	126.00
1	1G	1502	A	C4-N9-C1'	5.46	136.12	126.30
26	1H	141(A)	C	OP2-P-O3'	5.45	117.20	105.20
26	1H	210	C	OP2-P-O3'	5.45	117.20	105.20
26	1H	1432	C	N3-C2-O2	5.45	125.72	121.90
26	14	1558	A	C2-N3-C4	-5.45	107.87	110.60
26	1H	2871	C	O5'-P-OP2	-5.45	100.79	105.70
26	14	672	C	C6-N1-C2	-5.45	118.12	120.30
26	1H	30	G	OP1-P-O3'	5.45	117.19	105.20
26	1H	1356	G	C5-C6-N1	-5.45	108.77	111.50
26	14	2510	C	C4-C5-C6	5.45	120.12	117.40
26	1H	1429	G	OP1-P-OP2	5.45	127.77	119.60
26	1H	2427	C	O5'-P-OP1	-5.45	100.80	105.70
26	1H	2510	C	O5'-P-OP2	-5.45	100.80	105.70
1	13	896	C	C4-C5-C6	5.45	120.12	117.40
24	1L	34	U	C2-N1-C1'	5.45	124.24	117.70
26	1H	97	C	O5'-P-OP1	-5.45	100.80	105.70
26	1H	657	U	N3-C4-O4	-5.45	115.59	119.40
26	14	1407	C	N3-C2-O2	5.45	125.71	121.90
26	14	2592	G	O5'-P-OP1	5.45	117.23	110.70
26	14	1806	C	O5'-P-OP1	-5.44	100.80	105.70
24	3K	71	C	O4'-C1'-N1	5.44	112.55	108.20
1	1G	697	U	C5-C6-N1	-5.44	119.98	122.70
26	14	385	C	OP1-P-OP2	5.44	127.77	119.60
26	14	2873	A	C8-N9-C1'	-5.44	117.90	127.70
26	1H	724	U	C5-C4-O4	5.44	129.16	125.90
26	1H	730	C	N1-C2-O2	5.44	122.16	118.90
26	1H	855	G	C8-N9-C4	-5.44	104.22	106.40
26	14	1396	U	N1-C2-O2	5.44	126.61	122.80
26	14	1585	C	C2-N1-C1'	5.44	124.78	118.80
26	14	1950	G	C8-N9-C4	-5.44	104.22	106.40
1	1G	721	G	C6-C5-N7	-5.44	127.14	130.40
26	14	1680	U	C6-N1-C2	5.44	124.26	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	13	346	G	C4-N9-C1'	5.44	133.57	126.50
1	13	523	A	C2-N3-C4	-5.44	107.88	110.60
26	1H	120	U	C5-C4-O4	5.44	129.16	125.90
26	1H	2373	G	C6-N1-C2	-5.44	121.84	125.10
36	68	8	LEU	CA-CB-CG	5.44	127.81	115.30
1	13	150	C	C6-N1-C2	-5.43	118.13	120.30
26	1H	1571	A	C8-N9-C4	5.43	107.97	105.80
26	1H	2070	G	O5'-P-OP2	-5.43	100.81	105.70
26	1H	2294	C	C6-N1-C2	-5.43	118.13	120.30
1	1G	197	A	C8-N9-C4	-5.43	103.63	105.80
26	14	1674	G	N3-C4-N9	5.43	129.26	126.00
26	1H	2208	U	C5-C6-N1	-5.43	119.98	122.70
49	J8	2	SER	N-CA-C	5.43	125.67	111.00
1	13	15	G	C8-N9-C1'	-5.43	119.94	127.00
26	1H	1970	A	O4'-C1'-N9	-5.43	103.86	108.20
26	14	1304	C	N1-C2-O2	5.43	122.16	118.90
26	14	2518	A	C4-C5-N7	5.43	113.41	110.70
1	13	1498	U	C6-N1-C2	-5.43	117.75	121.00
26	1H	250	G	C8-N9-C4	-5.43	104.23	106.40
27	16	44	G	OP2-P-O3'	5.43	117.14	105.20
26	14	50	U	C6-N1-C2	5.43	124.26	121.00
26	1H	1252	G	N7-C8-N9	-5.42	110.39	113.10
26	1H	1608	A	C8-N9-C4	5.42	107.97	105.80
26	1H	1800	C	C2-N3-C4	-5.42	117.19	119.90
1	13	346	G	C5-N7-C8	-5.42	101.59	104.30
26	1H	956	G	N1-C6-O6	5.42	123.15	119.90
26	1H	1256	G	C8-N9-C4	5.42	108.57	106.40
27	16	45	A	O5'-P-OP1	-5.42	100.82	105.70
1	1G	380	G	N9-C4-C5	5.42	107.57	105.40
26	14	1516	U	N1-C2-O2	5.42	126.60	122.80
1	13	500	G	OP2-P-O3'	5.42	117.13	105.20
26	1H	1026	U	OP1-P-O3'	5.42	117.13	105.20
26	1H	2826	A	C8-N9-C4	5.42	107.97	105.80
26	14	1969	A	OP1-P-OP2	-5.42	111.47	119.60
26	14	2763	G	N3-C4-N9	5.42	129.25	126.00
26	14	2226	C	N3-C2-O2	-5.42	118.11	121.90
26	1H	1982	C	O5'-P-OP2	-5.42	100.82	105.70
26	1H	2838	G	N1-C6-O6	-5.42	116.65	119.90
26	14	1313	U	N1-C2-N3	5.42	118.15	114.90
26	14	1342	A	C5-N7-C8	-5.42	101.19	103.90
26	1H	838	C	N1-C2-N3	5.42	122.99	119.20
26	1H	196	A	C2-N3-C4	-5.42	107.89	110.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	1663	C	C6-N1-C2	5.42	122.47	120.30
26	1H	99	U	N1-C2-O2	5.41	126.59	122.80
1	1G	64	G	P-O3'-C3'	5.41	126.20	119.70
26	14	837	C	C5-C6-N1	5.41	123.71	121.00
49	F5	35	THR	C-N-CA	-5.41	110.93	122.30
1	13	726	C	OP1-P-O3'	5.41	117.11	105.20
26	1H	205	G	N7-C8-N9	-5.41	110.39	113.10
26	1H	1796	U	O5'-P-OP1	-5.41	100.83	105.70
26	1H	2626	C	N3-C4-C5	5.41	124.06	121.90
26	1H	120	U	O5'-P-OP1	-5.41	100.83	105.70
26	1H	270(O)	U	C5-C6-N1	5.41	125.41	122.70
26	1H	686	G	C6-C5-N7	-5.41	127.15	130.40
26	1H	944	G	C6-C5-N7	-5.41	127.15	130.40
26	1H	954	G	N9-C4-C5	5.41	107.56	105.40
26	1H	2286	A	C8-N9-C4	-5.41	103.64	105.80
26	1H	2416	C	C6-N1-C2	-5.41	118.14	120.30
55	P8	23	ARG	C-N-CA	5.41	135.23	121.70
26	14	130	C	C2-N3-C4	-5.41	117.19	119.90
26	14	203	C	N3-C4-N4	-5.41	114.21	118.00
26	1H	2392	A	N1-C6-N6	5.41	121.84	118.60
26	14	2282	G	O5'-P-OP2	5.41	117.19	110.70
26	1H	1365	A	C4-C5-C6	5.41	119.70	117.00
26	1H	2346	A	C1'-O4'-C4'	-5.41	105.58	109.90
26	14	1820	U	C5-C6-N1	-5.41	120.00	122.70
26	1H	2325	G	O5'-P-OP1	-5.40	100.84	105.70
26	14	2304	G	N3-C2-N2	-5.40	116.12	119.90
1	13	1113	C	C6-N1-C2	-5.40	118.14	120.30
1	13	1279	A	C5-N7-C8	-5.40	101.20	103.90
26	1H	606	U	O5'-P-OP2	-5.40	100.84	105.70
56	Q8	50	LEU	CB-CG-CD1	5.40	120.19	111.00
26	14	2334	G	N9-C4-C5	-5.40	103.24	105.40
1	13	858	G	C8-N9-C4	-5.40	104.24	106.40
26	14	1908	C	OP2-P-O3'	5.40	117.08	105.20
1	13	50	A	C2-N3-C4	5.40	113.30	110.60
26	1H	1129	A	OP1-P-OP2	5.40	127.69	119.60
26	14	1906	G	C8-N9-C4	-5.40	104.24	106.40
1	13	354	G	O5'-P-OP2	-5.39	100.85	105.70
26	14	155	C	N1-C2-O2	5.39	122.14	118.90
26	14	1973	G	N1-C6-O6	-5.39	116.66	119.90
26	1H	124	G	N1-C2-N3	-5.39	120.66	123.90
26	1H	1122	G	C5-C6-O6	-5.39	125.36	128.60
26	14	2072	G	OP1-P-OP2	-5.39	111.51	119.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	2430	A	C4-C5-C6	5.39	119.70	117.00
1	13	442	C	C6-N1-C2	-5.39	118.14	120.30
1	13	1498	U	N3-C4-O4	5.39	123.17	119.40
26	14	270(Y)	G	C5-C6-O6	5.39	131.84	128.60
26	1H	127	A	N9-C4-C5	-5.39	103.64	105.80
26	14	223	A	C8-N9-C4	-5.39	103.64	105.80
26	14	646	A	N7-C8-N9	5.39	116.49	113.80
1	13	974	A	C8-N9-C1'	-5.39	118.00	127.70
26	1H	2286	A	N1-C6-N6	5.39	121.83	118.60
27	16	12	C	N3-C2-O2	-5.39	118.13	121.90
26	1H	71	A	C6-N1-C2	-5.39	115.37	118.60
26	1H	382	G	OP1-P-O3'	5.39	117.05	105.20
26	1H	1786	A	N3-C4-C5	5.39	130.57	126.80
26	1H	1992	G	P-O3'-C3'	5.39	126.17	119.70
26	1H	311	A	O5'-P-OP1	-5.38	100.85	105.70
1	1G	117	G	C5-C6-O6	-5.38	125.37	128.60
1	1G	345	C	P-O3'-C3'	5.38	126.16	119.70
26	14	1404	C	OP1-P-OP2	5.38	127.68	119.60
26	14	1597	A	O5'-P-OP2	-5.38	100.86	105.70
26	14	1825	A	O5'-P-OP2	-5.38	100.86	105.70
56	Q8	46	ARG	C-N-CA	5.38	135.16	121.70
26	1H	1302	A	C5-C6-N1	5.38	120.39	117.70
1	13	1354	C	N3-C2-O2	-5.38	118.13	121.90
26	1H	2741	A	N7-C8-N9	-5.38	111.11	113.80
26	14	1779	U	O5'-P-OP2	-5.38	100.86	105.70
26	1H	210	C	C5-C6-N1	-5.38	118.31	121.00
26	1H	1367	A	C8-N9-C4	5.38	107.95	105.80
26	1H	2318	G	C5-N7-C8	-5.38	101.61	104.30
26	1H	2665	A	C8-N9-C4	-5.38	103.65	105.80
26	14	1248	G	OP1-P-OP2	-5.38	111.53	119.60
26	1H	1026	U	C2-N1-C1'	5.38	124.15	117.70
26	1H	1882	C	C2-N1-C1'	5.38	124.72	118.80
26	1H	2040	C	C5-C4-N4	-5.38	116.44	120.20
26	14	2067	G	N9-C4-C5	5.38	107.55	105.40
26	14	2346	A	C5-C6-N1	-5.38	115.01	117.70
26	1H	195	A	OP2-P-O3'	5.38	117.02	105.20
26	1H	2591	C	C5-C4-N4	-5.38	116.44	120.20
26	14	2073	C	N1-C2-O2	-5.38	115.67	118.90
26	1H	861	A	O5'-P-OP2	5.37	117.15	110.70
23	2L	76	C	N3-C4-C5	-5.37	119.75	121.90
26	14	596	G	N3-C2-N2	-5.37	116.14	119.90
26	1H	214	G	C8-N9-C4	-5.37	104.25	106.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	1253	A	N9-C4-C5	-5.37	103.65	105.80
26	14	409	C	C6-N1-C2	5.37	122.45	120.30
26	14	639	U	C5-C4-O4	5.37	129.12	125.90
37	35	59	LEU	CA-CB-CG	5.37	127.65	115.30
1	13	888	G	C5-C6-O6	-5.37	125.38	128.60
26	1H	2489	G	OP2-P-O3'	5.37	117.01	105.20
27	16	30	C	O5'-P-OP1	-5.37	100.87	105.70
26	14	1138	G	C8-N9-C4	-5.37	104.25	106.40
26	14	1406	U	OP1-P-O3'	5.37	117.01	105.20
26	14	465	G	O5'-P-OP2	5.37	117.14	110.70
26	14	2712	U	N1-C2-N3	5.37	118.12	114.90
1	13	800	G	N9-C4-C5	-5.37	103.25	105.40
26	1H	1752	C	C5-C6-N1	-5.37	118.32	121.00
26	14	250	G	C5-C6-O6	-5.37	125.38	128.60
1	13	190	G	P-O3'-C3'	5.36	126.14	119.70
26	1H	380	U	N3-C2-O2	-5.36	118.44	122.20
26	14	624	C	N1-C2-O2	-5.36	115.68	118.90
26	14	2056	G	N9-C4-C5	-5.36	103.25	105.40
26	1H	579	G	N1-C2-N2	5.36	121.03	116.20
26	1H	686	G	C8-N9-C1'	-5.36	120.03	127.00
26	1H	2040	C	N3-C2-O2	5.36	125.65	121.90
26	14	1253	A	O4'-C1'-N9	-5.36	103.91	108.20
26	14	1789	A	O5'-P-OP2	-5.36	100.88	105.70
26	1H	967	C	C2-N3-C4	-5.36	117.22	119.90
26	1H	2464	C	C6-N1-C2	5.36	122.44	120.30
1	1G	561	U	O5'-P-OP1	-5.36	100.88	105.70
26	14	988	A	N7-C8-N9	5.36	116.48	113.80
26	14	1022	G	P-O3'-C3'	5.36	126.13	119.70
26	14	1254	A	N1-C2-N3	5.36	131.98	129.30
26	14	2024	G	C5-C6-O6	-5.36	125.38	128.60
26	14	2245	U	OP1-P-O3'	5.36	116.99	105.20
26	1H	788	A	C5-C6-N1	-5.36	115.02	117.70
26	1H	1301	A	C6-C5-N7	-5.36	128.55	132.30
26	1H	1982	C	N3-C4-C5	-5.36	119.76	121.90
1	1G	1195	C	C6-N1-C2	-5.36	118.16	120.30
26	14	204	A	N1-C6-N6	5.36	121.81	118.60
26	14	2076	U	N1-C2-O2	-5.36	119.05	122.80
26	14	2461	C	O5'-P-OP1	-5.36	100.88	105.70
26	1H	978	G	C2-N3-C4	-5.36	109.22	111.90
26	1H	1798	U	N3-C4-O4	-5.36	115.65	119.40
26	14	1963	U	N3-C2-O2	-5.36	118.45	122.20
26	14	2501	C	C6-N1-C1'	5.36	127.23	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	193	U	N1-C2-N3	5.35	118.11	114.90
26	1H	665	C	C6-N1-C2	5.35	122.44	120.30
26	1H	1543	A	C5-C6-N1	-5.35	115.02	117.70
26	1H	1595	G	C5-C6-O6	-5.35	125.39	128.60
1	1G	1523	G	N1-C6-O6	-5.35	116.69	119.90
26	14	1925	C	N1-C2-O2	-5.35	115.69	118.90
24	3K	1	G	O4'-C1'-N9	5.35	112.48	108.20
24	3K	72	C	C6-N1-C2	-5.35	118.16	120.30
26	1H	1284	A	C5-N7-C8	-5.35	101.22	103.90
26	14	2296	U	N3-C4-O4	5.35	123.15	119.40
26	14	2346	A	N9-C1'-C2'	5.35	120.96	114.00
26	14	2592	G	C4-N9-C1'	5.35	133.46	126.50
26	1H	796	C	C5-C4-N4	5.35	123.94	120.20
26	1H	944	G	C8-N9-C1'	-5.35	120.05	127.00
26	1H	1305	C	C5-C6-N1	-5.35	118.33	121.00
26	1H	1346	G	N1-C6-O6	-5.35	116.69	119.90
26	1H	1374	G	C5-C6-N1	-5.35	108.83	111.50
26	1H	1811	G	N3-C2-N2	-5.35	116.16	119.90
1	1G	1301	U	N1-C2-O2	5.35	126.54	122.80
26	1H	752	A	N3-C4-C5	5.35	130.54	126.80
26	1H	2586	C	OP1-P-O3'	5.35	116.96	105.20
26	1H	2745	C	O5'-P-OP1	-5.35	100.89	105.70
26	14	1831	G	C8-N9-C1'	-5.35	120.05	127.00
26	14	137(A)	G	C5-C6-O6	-5.35	125.39	128.60
26	1H	2056	G	OP1-P-O3'	5.34	116.96	105.20
26	1H	2231	C	N1-C2-O2	-5.34	115.69	118.90
26	14	830	G	N9-C4-C5	-5.34	103.26	105.40
26	1H	698	C	OP1-P-OP2	5.34	127.61	119.60
1	1G	1502	A	N7-C8-N9	5.34	116.47	113.80
26	1H	74	A	C8-N9-C4	-5.34	103.66	105.80
26	1H	1417	C	C4-C5-C6	5.34	120.07	117.40
26	1H	1520	U	OP2-P-O3'	5.34	116.95	105.20
26	1H	2023	G	O5'-P-OP1	-5.34	100.89	105.70
26	1H	2030	A	C5-C6-N6	-5.34	119.43	123.70
26	14	2496	C	OP1-P-O3'	5.34	116.95	105.20
26	14	34	C	C6-N1-C1'	-5.34	114.39	120.80
26	14	2424	C	OP1-P-OP2	5.34	127.61	119.60
26	1H	2703	C	C6-N1-C2	-5.34	118.17	120.30
27	16	79	C	OP2-P-O3'	5.34	116.94	105.20
26	1H	733	G	C5-C6-N1	-5.34	108.83	111.50
26	1H	787	U	N3-C4-O4	-5.34	115.66	119.40
26	1H	2611	U	N3-C2-O2	-5.34	118.47	122.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	2334	G	N3-C4-N9	5.33	129.20	126.00
26	1H	848	G	O5'-P-OP1	5.33	117.10	110.70
26	1H	2666	C	C6-N1-C2	-5.33	118.17	120.30
26	14	1299	G	O5'-P-OP2	5.33	117.10	110.70
26	14	2287	A	N1-C6-N6	5.33	121.80	118.60
1	13	15	G	N3-C4-N9	5.33	129.20	126.00
1	13	904	C	C6-N1-C2	5.33	122.43	120.30
26	1H	2518	A	C2-N3-C4	-5.33	107.93	110.60
26	14	832	G	OP1-P-O3'	5.33	116.93	105.20
26	14	946	G	C8-N9-C4	5.33	108.53	106.40
26	14	2518	A	C4-C5-C6	5.33	119.67	117.00
1	13	972	C	N3-C2-O2	-5.33	118.17	121.90
26	1H	186	G	C5-C6-N1	5.33	114.17	111.50
1	13	703	G	OP1-P-O3'	5.33	116.92	105.20
26	1H	632	A	OP1-P-OP2	-5.33	111.61	119.60
26	1H	791	C	P-O3'-C3'	5.33	126.09	119.70
26	1H	2555	U	N1-C2-O2	-5.33	119.07	122.80
42	C8	74	LEU	CA-CB-CG	5.33	127.56	115.30
1	1G	576	G	C6-C5-N7	-5.33	127.20	130.40
1	1G	1322	C	N3-C2-O2	-5.33	118.17	121.90
26	14	1836	C	O5'-P-OP1	5.33	117.09	110.70
26	14	2609	U	C5-C6-N1	-5.33	120.04	122.70
29	11	272	ALA	C-N-CA	5.33	135.02	121.70
1	1G	811	C	C6-N1-C2	5.33	122.43	120.30
26	1H	2567	G	O5'-P-OP2	5.33	117.09	110.70
26	14	249	C	O5'-P-OP2	5.33	117.09	110.70
26	1H	2507	C	N3-C4-C5	-5.32	119.77	121.90
26	14	1585	C	C2-N3-C4	5.32	122.56	119.90
26	14	1831	G	N1-C2-N3	5.32	127.09	123.90
26	14	1836	C	OP1-P-O3'	5.32	116.91	105.20
26	1H	835	A	C2-N3-C4	5.32	113.26	110.60
26	1H	1291	C	N1-C2-O2	-5.32	115.71	118.90
26	1H	1610	A	C5-N7-C8	-5.32	101.24	103.90
26	1H	1776	G	C5-C6-O6	-5.32	125.41	128.60
26	14	2700	C	C6-N1-C1'	-5.32	114.41	120.80
26	1H	33	U	OP1-P-O3'	5.32	116.91	105.20
26	1H	2440	C	C5-C4-N4	5.32	123.92	120.20
26	1H	2761	G	C2-N3-C4	-5.32	109.24	111.90
26	14	677	A	O5'-P-OP2	-5.32	100.91	105.70
26	14	819	A	OP2-P-O3'	5.32	116.91	105.20
26	14	864	G	N3-C4-C5	-5.32	125.94	128.60
1	13	1301	U	OP1-P-O3'	5.32	116.90	105.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	869	G	C5-C6-O6	5.32	131.79	128.60
26	1H	938	G	O5'-P-OP1	5.32	117.08	110.70
1	1G	909	A	N9-C4-C5	-5.32	103.67	105.80
1	1G	950	U	O5'-P-OP1	-5.32	100.91	105.70
26	14	205	G	C4-C5-N7	5.32	112.93	110.80
26	1H	1941	C	C6-N1-C2	-5.32	118.17	120.30
26	14	205	G	N3-C2-N2	5.32	123.62	119.90
26	1H	201	C	N3-C4-C5	5.32	124.03	121.90
26	1H	534	U	C5-C6-N1	-5.32	120.04	122.70
26	1H	974(A)	C	C5-C6-N1	-5.32	118.34	121.00
26	1H	1669	A	C6-N1-C2	-5.32	115.41	118.60
1	1G	1260	C	C6-N1-C2	-5.32	118.17	120.30
26	14	124	G	N1-C6-O6	5.32	123.09	119.90
26	14	1899	G	C4-C5-C6	5.32	121.99	118.80
26	1H	446	G	C8-N9-C4	5.31	108.53	106.40
26	1H	508	G	P-O3'-C3'	5.31	126.08	119.70
26	1H	1593	G	C8-N9-C4	-5.31	104.28	106.40
26	1H	2032	G	C8-N9-C4	5.31	108.53	106.40
26	1H	2316	C	O5'-P-OP2	5.31	117.08	110.70
26	14	640	C	OP1-P-O3'	5.31	116.89	105.20
26	14	669	G	N3-C2-N2	-5.31	116.18	119.90
26	14	1204	A	N3-C4-C5	5.31	130.52	126.80
26	14	1382	G	N9-C4-C5	-5.31	103.28	105.40
26	14	1950	G	O5'-P-OP2	-5.31	100.92	105.70
26	1H	479	A	C8-N9-C4	5.31	107.92	105.80
1	13	45	U	OP2-P-O3'	5.31	116.88	105.20
26	1H	139	G	N3-C4-C5	-5.31	125.95	128.60
26	1H	726	G	C8-N9-C4	5.31	108.52	106.40
26	1H	1521	G	N3-C4-C5	-5.31	125.94	128.60
26	1H	1610	A	C8-N9-C4	5.31	107.92	105.80
1	1G	799	G	OP2-P-O3'	5.31	116.88	105.20
26	14	922	U	O5'-P-OP1	-5.31	100.92	105.70
26	1H	2239	G	N3-C2-N2	5.31	123.62	119.90
26	1H	2257	U	OP1-P-OP2	-5.31	111.64	119.60
1	1G	770	C	O5'-P-OP2	-5.31	100.92	105.70
26	14	2055	C	OP1-P-O3'	5.31	116.88	105.20
26	14	2334	G	C8-N9-C4	5.31	108.52	106.40
1	13	108	G	C6-C5-N7	-5.31	127.22	130.40
26	1H	1499	C	O5'-P-OP1	-5.31	100.92	105.70
26	1H	1731	G	N1-C6-O6	-5.31	116.72	119.90
26	1H	2357	U	O5'-P-OP2	-5.31	100.92	105.70
26	14	1659	U	O5'-P-OP1	-5.31	100.92	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	1793	C	C6-N1-C2	-5.31	118.18	120.30
27	1J	114	G	C4-N9-C1'	-5.31	119.60	126.50
26	1H	509	C	O5'-P-OP2	-5.30	100.92	105.70
26	1H	673	C	N1-C2-O2	-5.30	115.72	118.90
26	14	788	A	C4-C5-N7	5.30	113.35	110.70
26	14	2681	C	C5-C6-N1	-5.30	118.35	121.00
26	1H	831	G	N7-C8-N9	-5.30	110.45	113.10
26	1H	1502	C	C6-N1-C2	-5.30	118.18	120.30
26	1H	1611	C	C6-N1-C2	5.30	122.42	120.30
26	1H	2318	G	C4-N9-C1'	5.30	133.39	126.50
26	14	2390	U	C6-N1-C2	-5.30	117.82	121.00
27	1J	54	G	N7-C8-N9	5.30	115.75	113.10
1	13	690	G	N3-C4-N9	5.30	129.18	126.00
1	13	1516	G	N3-C4-N9	-5.30	122.82	126.00
26	1H	1136	G	N3-C2-N2	-5.30	116.19	119.90
26	1H	1609	A	C6-N1-C2	-5.30	115.42	118.60
26	1H	2062	A	N3-C4-N9	5.30	131.64	127.40
26	1H	2447	G	C6-N1-C2	-5.30	121.92	125.10
26	1H	501	A	O5'-P-OP2	-5.30	100.93	105.70
26	1H	2005	A	N1-C6-N6	5.30	121.78	118.60
26	14	2425	A	C5'-C4'-O4'	5.30	115.46	109.10
1	13	47	C	C5-C6-N1	-5.30	118.35	121.00
26	1H	628	G	C8-N9-C4	5.30	108.52	106.40
26	1H	84	A	N1-C2-N3	-5.30	126.65	129.30
26	1H	2272	U	OP2-P-O3'	5.30	116.85	105.20
26	14	571	A	N9-C4-C5	-5.30	103.68	105.80
26	14	2374	C	C2-N3-C4	-5.30	117.25	119.90
26	14	2377	A	C8-N9-C4	5.30	107.92	105.80
26	1H	2286	A	N7-C8-N9	5.29	116.45	113.80
26	14	2622	C	O5'-P-OP2	-5.29	100.93	105.70
26	1H	1990	C	C2-N3-C4	-5.29	117.25	119.90
26	14	141	A	OP2-P-O3'	5.29	116.84	105.20
26	14	832	G	N1-C2-N2	5.29	120.96	116.20
26	14	1725	G	C8-N9-C1'	-5.29	120.12	127.00
26	1H	817	C	C6-N1-C2	-5.29	118.18	120.30
26	1H	1254	A	N1-C6-N6	5.29	121.78	118.60
1	13	827	U	C4-C5-C6	5.29	122.87	119.70
26	1H	2586	C	C4-C5-C6	5.29	120.05	117.40
26	14	2595	G	C5-C6-N1	5.29	114.14	111.50
26	1H	1594	G	O5'-P-OP1	-5.29	100.94	105.70
23	2L	17	C	N1-C2-O2	5.29	122.07	118.90
26	14	2610	C	N1-C2-O2	5.29	122.07	118.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	13	721	G	N3-C4-N9	5.29	129.17	126.00
29	11	272	ALA	N-CA-C	5.29	125.27	111.00
26	1H	528	A	O4'-C1'-N9	-5.29	103.97	108.20
1	13	50	A	P-O3'-C3'	5.28	126.04	119.70
26	1H	789	A	C2-N3-C4	-5.28	107.96	110.60
26	1H	2032	G	OP1-P-O3'	5.28	116.82	105.20
1	13	768	A	N7-C8-N9	-5.28	111.16	113.80
26	1H	145	G	C5-C6-O6	-5.28	125.43	128.60
26	1H	1021	A	C4-C5-N7	5.28	113.34	110.70
26	14	1801	G	O5'-P-OP1	-5.28	100.95	105.70
26	1H	1565	C	C5-C6-N1	-5.28	118.36	121.00
26	1H	1957	C	C5-C6-N1	-5.28	118.36	121.00
26	14	779	U	C4-C5-C6	5.28	122.87	119.70
1	13	529	G	C5-C6-O6	-5.28	125.43	128.60
1	13	791	G	OP2-P-O3'	5.28	116.81	105.20
26	1H	231	C	N1-C2-O2	-5.28	115.73	118.90
26	1H	1914	C	C6-N1-C2	-5.28	118.19	120.30
26	14	2307	G	N7-C8-N9	5.28	115.74	113.10
33	59	98	LEU	CA-CB-CG	5.28	127.44	115.30
1	13	452	A	O5'-P-OP1	-5.28	100.95	105.70
26	1H	740	U	O5'-P-OP1	5.28	117.03	110.70
26	14	456	C	N1-C2-O2	-5.28	115.73	118.90
26	1H	2239	G	N1-C6-O6	-5.27	116.74	119.90
26	14	1826	G	N7-C8-N9	-5.27	110.46	113.10
1	13	904	C	N3-C4-C5	5.27	124.01	121.90
12	3I	93	LEU	CA-CB-CG	5.27	127.43	115.30
1	1G	1260	C	C5-C6-N1	5.27	123.64	121.00
26	14	1312	U	O5'-P-OP1	-5.27	100.95	105.70
26	1H	1835	G	N3-C4-C5	-5.27	125.96	128.60
26	1H	2374	C	C4-C5-C6	5.27	120.03	117.40
26	1H	256	A	N9-C4-C5	-5.27	103.69	105.80
1	13	1061	G	N1-C6-O6	5.27	123.06	119.90
1	1G	1259	C	C5-C6-N1	5.27	123.63	121.00
26	14	509	C	C5-C6-N1	-5.27	118.37	121.00
26	14	715	G	C4-C5-N7	5.27	112.91	110.80
26	14	945	A	C8-N9-C1'	-5.27	118.22	127.70
22	1K	49	G	C4-N9-C1'	-5.27	119.65	126.50
26	1H	800	A	C5-N7-C8	5.27	106.53	103.90
26	14	621	A	O4'-C1'-N9	5.27	112.41	108.20
26	14	1842	G	C5-C6-N1	5.27	114.13	111.50
1	13	15	G	C4-N9-C1'	5.26	133.34	126.50
26	1H	213	A	C4-C5-N7	5.26	113.33	110.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	446	G	C6-C5-N7	-5.26	127.24	130.40
26	1H	745	G	N3-C2-N2	-5.26	116.22	119.90
26	1H	1210	A	N1-C6-N6	5.26	121.76	118.60
26	1H	1647	G	O5'-P-OP1	-5.26	100.96	105.70
26	1H	2011	U	C2-N1-C1'	-5.26	111.38	117.70
26	1H	2709	G	C8-N9-C4	-5.26	104.29	106.40
26	14	1216	G	OP1-P-O3'	5.26	116.78	105.20
1	13	388	G	C8-N9-C4	5.26	108.50	106.40
56	M5	50	LEU	CA-CB-CG	-5.26	103.19	115.30
26	1H	1888	G	N9-C4-C5	-5.26	103.30	105.40
26	14	1681	G	N7-C8-N9	5.26	115.73	113.10
26	14	1831	G	C4-N9-C1'	5.26	133.34	126.50
26	14	2056	G	N3-C4-N9	5.26	129.16	126.00
26	14	2598	A	N7-C8-N9	-5.26	111.17	113.80
26	1H	737	C	N1-C2-O2	-5.26	115.75	118.90
26	1H	1224	G	C4-N9-C1'	-5.26	119.66	126.50
26	1H	2505	G	C2-N3-C4	-5.26	109.27	111.90
26	1H	2558	C	N1-C2-O2	-5.26	115.74	118.90
26	1H	2718	G	C4-C5-N7	5.26	112.90	110.80
1	13	807	A	C8-N9-C4	-5.26	103.70	105.80
26	1H	140	A	C5-C6-N6	-5.26	119.49	123.70
1	1G	481	G	C4-C5-C6	5.26	121.95	118.80
26	14	2258	C	OP1-P-O3'	5.26	116.77	105.20
1	13	1403	C	O5'-P-OP2	-5.26	100.97	105.70
26	1H	199	A	N1-C2-N3	-5.26	126.67	129.30
26	1H	481	G	N1-C6-O6	5.26	123.05	119.90
26	1H	2517	C	O4'-C1'-N1	5.26	112.41	108.20
34	61	110	ASP	C-N-CD	-5.26	109.04	120.60
1	13	108	G	C8-N9-C1'	-5.25	120.17	127.00
26	1H	698	C	N3-C4-N4	5.25	121.68	118.00
26	1H	1770	G	N1-C6-O6	-5.25	116.75	119.90
26	1H	2469	A	C4-C5-N7	5.25	113.33	110.70
26	14	774	A	OP1-P-O3'	-5.25	93.64	105.20
26	1H	141	A	C4-C5-C6	-5.25	114.37	117.00
26	1H	2438	U	C6-N1-C2	5.25	124.15	121.00
26	14	22	C	N3-C4-C5	5.25	124.00	121.90
26	14	773	U	N3-C2-O2	-5.25	118.52	122.20
26	14	1635	G	OP1-P-O3'	5.25	116.76	105.20
26	14	1950	G	C4-N9-C1'	5.25	133.33	126.50
1	13	899	C	OP2-P-O3'	5.25	116.75	105.20
26	1H	110	G	N7-C8-N9	-5.25	110.47	113.10
26	1H	512	G	N7-C8-N9	-5.25	110.47	113.10

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	1G	1128	C	N3-C2-O2	-5.25	118.22	121.90
26	14	29	U	C2-N1-C1'	5.25	124.00	117.70
26	14	621	A	C6-C5-N7	-5.25	128.62	132.30
1	13	703	G	N9-C4-C5	-5.25	103.30	105.40
26	1H	1878	G	N9-C4-C5	5.25	107.50	105.40
26	1H	2552	U	N1-C2-O2	-5.25	119.12	122.80
40	A8	101	LEU	CA-CB-CG	5.25	127.38	115.30
26	14	2731	G	C6-C5-N7	-5.25	127.25	130.40
26	1H	1678	G	N1-C6-O6	5.25	123.05	119.90
26	1H	2375	G	OP2-P-O3'	5.25	116.75	105.20
27	16	105	G	N3-C4-C5	-5.25	125.98	128.60
26	14	1819	A	N9-C4-C5	5.25	107.90	105.80
26	14	2873	A	N9-C1'-C2'	5.25	120.82	114.00
29	19	257	LEU	CA-CB-CG	5.25	127.37	115.30
1	13	817	C	N3-C4-N4	5.25	121.67	118.00
26	1H	141(A)	C	C2-N3-C4	-5.25	117.28	119.90
26	14	1757	U	C5-C4-O4	5.25	129.05	125.90
26	14	2329	G	C5-C6-N1	5.25	114.12	111.50
26	1H	248	G	C6-N1-C2	-5.24	121.95	125.10
26	1H	664	C	N3-C2-O2	-5.24	118.23	121.90
26	1H	2040	C	N1-C2-O2	-5.24	115.75	118.90
23	2K	73	A	C8-N9-C4	5.24	107.90	105.80
26	1H	815	C	C6-N1-C2	5.24	122.40	120.30
26	1H	2603	G	O5'-P-OP1	-5.24	100.98	105.70
1	1G	1496	C	O5'-P-OP2	-5.24	100.98	105.70
11	2A	63	LEU	CA-CB-CG	5.24	127.35	115.30
26	14	137(A)	G	N1-C6-O6	5.24	123.05	119.90
26	14	2332	U	C5-C4-O4	5.24	129.04	125.90
1	13	910	C	O5'-P-OP2	-5.24	100.99	105.70
1	13	963	G	N3-C4-C5	-5.24	125.98	128.60
26	1H	835	A	N9-C4-C5	5.24	107.90	105.80
37	78	23	PRO	C-N-CA	-5.24	111.30	122.30
1	1G	924	C	OP1-P-OP2	5.24	127.46	119.60
26	14	1781	C	C5-C4-N4	-5.24	116.53	120.20
26	1H	2458	G	C5-C6-O6	-5.24	125.46	128.60
1	13	1	U	N1-C1'-C2'	5.24	120.81	114.00
1	13	1126	U	C6-N1-C2	-5.24	117.86	121.00
26	1H	1432	C	C5-C4-N4	-5.24	116.53	120.20
26	1H	1496	A	O4'-C1'-N9	5.24	112.39	108.20
26	1H	2347	C	OP2-P-O3'	5.24	116.72	105.20
26	14	552	G	C8-N9-C4	5.24	108.49	106.40
26	14	2013	A	C2-N3-C4	-5.24	107.98	110.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	1G	1502	A	C5-N7-C8	-5.23	101.28	103.90
26	14	773	U	N1-C2-N3	5.23	118.04	114.90
26	14	827	U	O5'-P-OP1	5.23	116.98	110.70
26	14	916	G	O5'-P-OP1	-5.23	100.99	105.70
26	14	1772	G	C4-C5-N7	5.23	112.89	110.80
26	1H	123	G	C5-C6-N1	5.23	114.12	111.50
26	14	2729	G	C4-C5-N7	5.23	112.89	110.80
26	1H	1361	G	C6-C5-N7	5.23	133.54	130.40
26	14	788	A	C4-C5-C6	5.23	119.61	117.00
26	14	972	G	OP1-P-O3'	5.23	116.70	105.20
1	13	952	U	N1-C2-N3	5.23	118.04	114.90
26	1H	350	U	C5-C4-O4	5.23	129.04	125.90
26	1H	655	A	C8-N9-C4	-5.23	103.71	105.80
26	14	133	C	N3-C4-C5	5.23	123.99	121.90
26	14	764	A	N1-C2-N3	-5.23	126.69	129.30
26	1H	36	G	OP2-P-O3'	5.23	116.70	105.20
26	1H	121	G	C6-N1-C2	-5.23	121.97	125.10
26	14	608	A	C4-C5-C6	5.23	119.61	117.00
26	1H	1157	G	C4-N9-C1'	5.22	133.29	126.50
26	14	2438	U	O5'-P-OP2	-5.22	101.00	105.70
1	13	800	G	N3-C4-N9	5.22	129.13	126.00
26	1H	125	G	N9-C4-C5	-5.22	103.31	105.40
26	1H	591	C	C4-C5-C6	5.22	120.01	117.40
26	14	1571	A	C5-C6-N6	-5.22	119.52	123.70
26	14	1762	A	C8-N9-C1'	-5.22	118.30	127.70
26	14	2346	A	C8-N9-C1'	-5.22	118.30	127.70
23	2K	40	C	C5-C6-N1	5.22	123.61	121.00
26	14	845	G	C4-C5-N7	5.22	112.89	110.80
26	14	944	G	C4-N9-C1'	5.22	133.29	126.50
26	14	1767	C	C5-C4-N4	5.22	123.86	120.20
26	1H	428	A	OP1-P-O3'	5.22	116.68	105.20
26	1H	717	G	C6-C5-N7	-5.22	127.27	130.40
26	14	203	C	N3-C4-C5	5.22	123.99	121.90
26	1H	1302	A	C2-N3-C4	5.22	113.21	110.60
26	1H	2286	A	C6-C5-N7	-5.22	128.65	132.30
26	14	2388	A	O5'-P-OP1	5.22	116.96	110.70
26	14	2565	A	N9-C4-C5	-5.22	103.71	105.80
26	1H	1448	G	O5'-P-OP1	-5.22	101.01	105.70
27	1J	22	U	C5-C6-N1	5.22	125.31	122.70
26	1H	1598	C	C5-C6-N1	-5.21	118.39	121.00
26	14	2573	C	C6-N1-C1'	-5.21	114.54	120.80
26	1H	2062	A	C8-N9-C4	5.21	107.89	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	2891	G	C5-C6-O6	-5.21	125.47	128.60
27	16	80	U	N1-C2-O2	5.21	126.45	122.80
26	14	946	G	C2-N3-C4	-5.21	109.30	111.90
26	14	1379	A	C4-C5-N7	5.21	113.31	110.70
37	35	55	ARG	NE-CZ-NH1	-5.21	117.69	120.30
26	14	2287	A	N1-C2-N3	5.21	131.91	129.30
26	1H	1544	C	C2-N1-C1'	5.21	124.53	118.80
26	1H	2699	C	N3-C4-C5	5.21	123.98	121.90
26	1H	2765	A	OP1-P-OP2	5.21	127.41	119.60
49	J8	2	SER	CB-CA-C	-5.21	100.20	110.10
26	14	246	C	C6-N1-C2	5.21	122.38	120.30
26	14	2275	C	C5'-C4'-O4'	-5.21	102.85	109.10
30	29	44	TYR	CA-CB-CG	5.21	123.30	113.40
26	1H	462	C	OP1-P-OP2	5.21	127.41	119.60
26	1H	2281	C	C5-C4-N4	-5.21	116.56	120.20
26	14	679	C	N1-C2-O2	-5.21	115.78	118.90
26	1H	124	G	C4-N9-C1'	-5.21	119.73	126.50
26	1H	1602	U	C5-C4-O4	5.20	129.02	125.90
40	A8	9	ARG	NE-CZ-NH1	-5.20	117.70	120.30
1	1G	810	C	N3-C2-O2	-5.20	118.26	121.90
26	14	970	C	C6-N1-C1'	5.20	127.04	120.80
26	1H	1252	G	C5-N7-C8	5.20	106.90	104.30
26	14	1799	G	C5-C6-O6	5.20	131.72	128.60
26	1H	207	A	N9-C4-C5	-5.20	103.72	105.80
26	1H	1882	C	C6-N1-C2	-5.20	118.22	120.30
26	1H	1934	C	N3-C4-C5	-5.20	119.82	121.90
26	1H	2370	G	N1-C2-N2	-5.20	111.52	116.20
1	13	853	G	C5-C6-N1	-5.20	108.90	111.50
26	1H	1497	U	OP1-P-O3'	5.20	116.64	105.20
1	1G	1305	G	N3-C4-N9	-5.20	122.88	126.00
26	14	641	C	C6-N1-C2	5.20	122.38	120.30
27	16	100	G	N9-C4-C5	-5.20	103.32	105.40
1	13	963	G	C5-C6-O6	5.20	131.72	128.60
1	13	1502	A	N7-C8-N9	5.20	116.40	113.80
26	1H	1275	A	C2-N3-C4	-5.20	108.00	110.60
26	1H	2375	G	N9-C1'-C2'	-5.20	106.28	112.00
26	1H	2623	G	N9-C4-C5	5.20	107.48	105.40
1	1G	1499	A	C8-N9-C4	5.20	107.88	105.80
26	14	141	A	C5-N7-C8	-5.20	101.30	103.90
26	14	933	A	C4-C5-N7	5.20	113.30	110.70
26	14	998	C	N1-C2-O2	5.20	122.02	118.90
26	14	2211	G	O5'-P-OP2	-5.20	101.02	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	2297	C	OP1-P-OP2	5.20	127.39	119.60
1	13	741	G	O5'-P-OP2	-5.19	101.03	105.70
26	14	2446	G	OP2-P-O3'	5.19	116.63	105.20
26	1H	245	G	C4-C5-N7	5.19	112.88	110.80
26	1H	478	A	N1-C6-N6	-5.19	115.48	118.60
26	1H	695	G	N1-C6-O6	-5.19	116.78	119.90
26	1H	2282	G	C5-C6-O6	5.19	131.72	128.60
26	1H	2390	U	C2-N1-C1'	5.19	123.93	117.70
26	14	2331	G	C4-C5-N7	5.19	112.88	110.80
26	14	2581	G	N3-C4-C5	-5.19	126.00	128.60
1	13	533	A	C2-N3-C4	-5.19	108.00	110.60
26	1H	136	G	C5-C6-N1	5.19	114.09	111.50
26	1H	1410	G	N3-C4-C5	5.19	131.20	128.60
26	1H	627	A	OP1-P-O3'	5.19	116.62	105.20
1	13	413	G	O4'-C1'-N9	5.19	112.35	108.20
1	13	522	C	O5'-P-OP2	-5.19	101.03	105.70
26	1H	766	C	N3-C4-C5	5.19	123.97	121.90
26	1H	1777	U	N1-C2-N3	5.19	118.01	114.90
26	1H	2007	C	O5'-P-OP2	-5.19	101.03	105.70
26	1H	2502	G	C8-N9-C4	-5.19	104.33	106.40
26	1H	2872	G	O5'-P-OP2	-5.19	101.03	105.70
26	1H	1344	G	C8-N9-C4	-5.19	104.33	106.40
26	1H	1470	G	N1-C6-O6	5.19	123.01	119.90
26	1H	1799	G	N3-C4-C5	-5.19	126.01	128.60
26	14	201	C	C6-N1-C2	5.19	122.38	120.30
26	14	393	C	C5-C4-N4	5.19	123.83	120.20
26	14	2681	C	N3-C2-O2	-5.19	118.27	121.90
26	1H	1842	G	C5-C6-O6	5.18	131.71	128.60
26	1H	2033	A	C5-C6-N6	-5.18	119.55	123.70
23	2L	6	G	C8-N9-C4	5.18	108.47	106.40
26	14	1678	G	C4-C5-N7	5.18	112.87	110.80
26	14	2304	G	N9-C4-C5	5.18	107.47	105.40
27	1J	71	C	C2-N1-C1'	5.18	124.50	118.80
26	1H	840	C	O5'-P-OP2	-5.18	101.04	105.70
26	1H	1997	G	C2-N3-C4	-5.18	109.31	111.90
26	1H	2821	A	N9-C4-C5	-5.18	103.73	105.80
26	1H	2280	G	OP1-P-OP2	-5.18	111.83	119.60
26	14	1742	C	C6-N1-C2	-5.18	118.23	120.30
26	1H	829	A	O5'-P-OP1	-5.18	101.04	105.70
26	1H	2328	A	C2-N3-C4	-5.18	108.01	110.60
1	1G	25	C	O5'-P-OP2	-5.18	101.04	105.70
26	14	2699	C	C5-C6-N1	-5.18	118.41	121.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	471	A	C6-N1-C2	5.18	121.71	118.60
26	1H	772	C	C6-N1-C2	5.18	122.37	120.30
26	14	2281	C	C5-C6-N1	5.18	123.59	121.00
1	13	1227	A	C2-N3-C4	-5.18	108.01	110.60
26	1H	630	G	C4-N9-C1'	-5.18	119.77	126.50
26	1H	848	G	C4-N9-C1'	5.18	133.23	126.50
26	1H	1307	A	N9-C4-C5	-5.18	103.73	105.80
26	1H	1357	U	OP1-P-OP2	5.18	127.36	119.60
26	1H	1669	A	N3-C4-N9	5.18	131.54	127.40
26	1H	1858	G	C8-N9-C1'	-5.18	120.27	127.00
26	1H	2519	U	N3-C2-O2	5.18	125.82	122.20
26	14	852	G	O5'-P-OP2	-5.18	101.04	105.70
26	14	1667	G	N1-C6-O6	5.18	123.01	119.90
26	1H	733	G	N3-C4-N9	5.17	129.10	126.00
26	1H	1674	G	N1-C6-O6	5.17	123.00	119.90
26	14	1785	A	C4-C5-C6	5.17	119.59	117.00
26	1H	126	A	OP2-P-O3'	5.17	116.58	105.20
26	1H	726	G	OP1-P-OP2	5.17	127.36	119.60
26	14	566	U	O5'-P-OP2	-5.17	101.04	105.70
26	1H	74	A	C4-C5-C6	5.17	119.59	117.00
26	1H	952	G	O5'-P-OP2	5.17	116.91	110.70
26	1H	2509	G	C8-N9-C4	5.17	108.47	106.40
26	14	250	G	C8-N9-C4	-5.17	104.33	106.40
26	14	744	G	O5'-P-OP2	-5.17	101.05	105.70
26	1H	71	A	C8-N9-C4	-5.17	103.73	105.80
26	14	973	A	OP1-P-OP2	-5.17	111.84	119.60
1	13	827	U	C6-N1-C1'	-5.17	113.97	121.20
26	1H	745	G	OP1-P-OP2	-5.17	111.85	119.60
26	1H	2713	A	O4'-C1'-N9	-5.17	104.06	108.20
24	3L	76	A	C6-C5-N7	-5.17	128.68	132.30
26	14	2029	G	O5'-P-OP1	-5.17	101.05	105.70
1	13	757	U	C5-C6-N1	-5.17	120.12	122.70
1	13	1530	G	N1-C6-O6	5.17	123.00	119.90
26	1H	123	G	O5'-P-OP2	-5.17	101.05	105.70
26	1H	1315	C	N3-C4-N4	-5.17	114.38	118.00
26	1H	2446	G	N1-C6-O6	5.17	123.00	119.90
26	1H	2581	G	N3-C2-N2	5.17	123.52	119.90
1	13	966	G	C8-N9-C4	5.17	108.47	106.40
26	1H	1978	A	N1-C6-N6	-5.17	115.50	118.60
26	1H	103	A	N9-C4-C5	-5.16	103.73	105.80
26	1H	140	A	N3-C4-C5	5.16	130.41	126.80
26	1H	528	A	N1-C6-N6	5.16	121.70	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	1210	A	C5-C6-N1	-5.16	115.12	117.70
26	1H	1973	G	N1-C6-O6	-5.16	116.80	119.90
26	1H	1978	A	N9-C4-C5	5.16	107.87	105.80
1	1G	1370	G	C5-C6-N1	-5.16	108.92	111.50
26	14	149	A	C4-C5-C6	5.16	119.58	117.00
26	14	530	G	N7-C8-N9	5.16	115.68	113.10
26	14	2042	A	C8-N9-C4	5.16	107.87	105.80
26	14	2057	A	O5'-P-OP2	-5.16	101.06	105.70
26	14	2703	C	N3-C2-O2	-5.16	118.29	121.90
26	1H	451	C	C6-N1-C2	5.16	122.36	120.30
26	1H	602	G	C6-C5-N7	-5.16	127.30	130.40
26	1H	1473	G	OP1-P-O3'	5.16	116.55	105.20
26	1H	2249	U	N3-C4-C5	5.16	117.69	114.60
26	1H	2706	G	C5-C6-N1	5.16	114.08	111.50
30	21	49	LEU	CA-CB-CG	-5.16	103.44	115.30
26	14	372	G	O4'-C1'-N9	5.16	112.33	108.20
26	14	1831	G	N1-C2-N2	-5.16	111.56	116.20
26	14	2030	A	O5'-P-OP2	-5.16	101.06	105.70
1	13	813	U	N3-C2-O2	-5.16	118.59	122.20
1	13	1516	G	C5-C6-O6	5.16	131.69	128.60
26	14	1899	G	C4-C5-N7	5.16	112.86	110.80
26	1H	139	G	C2-N3-C4	5.16	114.48	111.90
26	1H	1573	G	C4-C5-N7	5.16	112.86	110.80
26	14	678	C	C6-N1-C2	5.16	122.36	120.30
26	14	2263	C	C6-N1-C2	-5.16	118.24	120.30
26	14	2380	C	N1-C2-O2	-5.16	115.81	118.90
26	14	2607	G	N1-C2-N2	-5.16	111.56	116.20
27	1J	44	G	N7-C8-N9	-5.16	110.52	113.10
26	1H	2623	G	N1-C2-N2	-5.15	111.56	116.20
26	1H	2697	G	N1-C6-O6	-5.15	116.81	119.90
27	16	44	G	C6-C5-N7	5.15	133.49	130.40
27	16	108	C	O4'-C1'-N1	5.15	112.32	108.20
26	14	783	A	N9-C1'-C2'	-5.15	106.33	112.00
26	14	1776	G	N9-C4-C5	-5.15	103.34	105.40
26	1H	1888	G	O4'-C1'-N9	5.15	112.32	108.20
26	1H	2623	G	C5-C6-O6	5.15	131.69	128.60
26	14	1610	A	C8-N9-C4	5.15	107.86	105.80
26	14	1975	G	O5'-P-OP1	5.15	116.88	110.70
27	1J	71	C	C6-N1-C2	-5.15	118.24	120.30
1	13	108	G	C5-N7-C8	-5.15	101.72	104.30
1	13	1197	G	OP1-P-O3'	5.15	116.53	105.20
26	1H	1604	C	C2-N3-C4	-5.15	117.33	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	1G	932	C	N1-C2-O2	5.15	121.99	118.90
1	1G	1071	C	C6-N1-C2	-5.15	118.24	120.30
1	1G	898	G	O5'-P-OP2	-5.15	101.07	105.70
26	14	2430	A	N3-C4-C5	5.15	130.40	126.80
26	14	2589	A	C8-N9-C4	5.15	107.86	105.80
1	13	988	G	C8-N9-C4	-5.15	104.34	106.40
1	13	1489	G	N9-C4-C5	-5.15	103.34	105.40
26	1H	241	A	C2-N3-C4	-5.15	108.03	110.60
26	1H	1393	A	O5'-P-OP2	-5.15	101.07	105.70
26	14	2542	A	O5'-P-OP2	-5.15	101.07	105.70
1	13	1519	A	C5-C6-N6	5.15	127.82	123.70
26	1H	591	C	C2-N3-C4	-5.15	117.33	119.90
26	1H	463	G	N1-C2-N2	-5.14	111.57	116.20
26	1H	803	U	O4'-C1'-N1	5.14	112.31	108.20
26	1H	2537	U	N3-C4-O4	-5.14	115.80	119.40
26	14	161	U	C5-C6-N1	5.14	125.27	122.70
26	14	681	G	C8-N9-C4	5.14	108.46	106.40
26	14	772	C	O5'-P-OP1	-5.14	101.07	105.70
26	14	1642	G	OP2-P-O3'	5.14	116.52	105.20
26	14	1799	G	C4-C5-N7	-5.14	108.74	110.80
26	1H	481	G	C5-C6-O6	-5.14	125.51	128.60
26	1H	1308	A	N1-C2-N3	5.14	131.87	129.30
26	1H	1996	C	N3-C4-N4	-5.14	114.40	118.00
26	14	746	A	O4'-C1'-N9	5.14	112.31	108.20
26	14	775	G	N3-C4-N9	5.14	129.09	126.00
26	14	1681	G	C8-N9-C4	-5.14	104.34	106.40
27	1J	35	U	N3-C4-O4	-5.14	115.80	119.40
26	14	2359	C	O5'-P-OP1	-5.14	101.07	105.70
26	1H	265	A	N9-C1'-C2'	5.14	120.68	114.00
26	1H	777	A	OP2-P-O3'	5.14	116.50	105.20
26	1H	1022	G	P-O3'-C3'	5.14	125.87	119.70
26	14	1379	A	N7-C8-N9	5.14	116.37	113.80
1	13	896	C	C6-N1-C2	5.14	122.36	120.30
26	1H	1401	G	C5-C6-O6	5.14	131.68	128.60
26	1H	1673	U	C6-N1-C2	5.14	124.08	121.00
26	1H	2257	U	N1-C2-N3	5.14	117.98	114.90
26	1H	126	A	OP1-P-OP2	5.14	127.31	119.60
26	1H	1778	U	OP2-P-O3'	5.14	116.50	105.20
26	1H	2078	C	N3-C4-N4	5.14	121.59	118.00
1	1G	509	A	P-O3'-C3'	5.14	125.86	119.70
1	1G	576	G	N3-C4-C5	-5.14	126.03	128.60
26	14	514	A	OP1-P-OP2	5.14	127.31	119.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	804	A	O4'-C1'-N9	5.14	112.31	108.20
26	14	1680	U	C5-C4-O4	-5.14	122.82	125.90
26	1H	411	G	OP1-P-OP2	5.13	127.30	119.60
26	1H	868	U	N3-C2-O2	-5.13	118.61	122.20
26	1H	906	G	C5-C6-O6	5.13	131.68	128.60
26	1H	2415	G	N3-C2-N2	-5.13	116.31	119.90
26	14	1300	U	O5'-P-OP1	5.13	116.86	110.70
26	14	1807	G	O5'-P-OP2	-5.13	101.08	105.70
26	1H	963	U	N3-C4-O4	5.13	122.99	119.40
1	1G	1228	C	O5'-P-OP2	-5.13	101.08	105.70
26	14	784	A	P-O3'-C3'	5.13	125.86	119.70
26	14	1349	A	C4-C5-N7	5.13	113.27	110.70
26	14	1379	A	P-O3'-C3'	5.13	125.86	119.70
26	14	1482	U	C6-N1-C1'	5.13	128.38	121.20
26	14	2392	A	N3-C4-N9	-5.13	123.30	127.40
26	14	2518	A	O5'-P-OP2	-5.13	101.08	105.70
26	1H	2261	C	OP2-P-O3'	5.13	116.48	105.20
26	1H	2604	U	C2-N1-C1'	5.13	123.86	117.70
24	1L	69	A	P-O3'-C3'	5.13	125.86	119.70
1	13	266	G	N7-C8-N9	5.13	115.66	113.10
26	1H	1683	C	C2-N3-C4	-5.13	117.34	119.90
26	1H	1801	G	N3-C4-N9	5.13	129.08	126.00
26	1H	2380	C	N1-C2-O2	-5.13	115.82	118.90
1	1G	1518	A	O5'-P-OP1	-5.13	101.08	105.70
1	13	268	C	O5'-P-OP2	5.13	116.85	110.70
26	1H	252	G	C6-C5-N7	-5.13	127.33	130.40
26	1H	535	C	C2-N1-C1'	-5.13	113.16	118.80
26	1H	829	A	C2-N3-C4	-5.13	108.04	110.60
26	1H	2377	A	N3-C4-C5	5.13	130.39	126.80
26	1H	2599	G	N1-C6-O6	-5.13	116.82	119.90
26	1H	2689	U	N3-C4-C5	-5.13	111.52	114.60
26	14	1463	C	C6-N1-C2	-5.13	118.25	120.30
26	1H	1804	C	N3-C4-C5	5.12	123.95	121.90
26	14	1251	C	C5-C4-N4	-5.12	116.61	120.20
26	14	1482	U	C2-N1-C1'	-5.12	111.55	117.70
26	1H	139	G	O5'-P-OP1	-5.12	101.09	105.70
26	1H	767	U	N1-C2-N3	5.12	117.97	114.90
26	14	2251	G	C4-C5-N7	-5.12	108.75	110.80
26	14	2430	A	O4'-C1'-N9	-5.12	104.10	108.20
1	13	572	A	N1-C6-N6	-5.12	115.53	118.60
1	1G	134	A	N1-C6-N6	5.12	121.67	118.60
23	2K	27	G	N1-C6-O6	5.12	122.97	119.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	625	G	N9-C4-C5	5.12	107.45	105.40
26	1H	1032	A	N1-C6-N6	5.12	121.67	118.60
26	1H	1818	U	N3-C4-C5	5.12	117.67	114.60
26	1H	2716	U	OP2-P-O3'	5.12	116.47	105.20
26	1H	602	G	N1-C2-N2	-5.12	111.59	116.20
26	1H	766	C	N1-C2-O2	-5.12	115.83	118.90
26	1H	1162	G	O5'-P-OP1	-5.12	101.09	105.70
26	1H	2070	G	N1-C6-O6	-5.12	116.83	119.90
26	14	238	C	N1-C2-O2	-5.12	115.83	118.90
1	13	524	G	O5'-P-OP1	-5.12	101.09	105.70
26	1H	737	C	OP1-P-O3'	-5.12	93.94	105.20
26	1H	1021	A	N1-C6-N6	5.12	121.67	118.60
26	1H	1535	U	OP1-P-O3'	5.12	116.45	105.20
26	1H	1555	G	O5'-P-OP2	5.12	116.84	110.70
1	1G	541	G	N1-C6-O6	5.12	122.97	119.90
27	1J	89	G	C8-N9-C1'	-5.12	120.35	127.00
26	1H	732	C	C4-C5-C6	5.11	119.96	117.40
26	14	1588	C	C6-N1-C2	-5.11	118.25	120.30
26	14	608	A	C8-N9-C4	-5.11	103.76	105.80
26	14	1372	U	C2-N3-C4	-5.11	123.93	127.00
26	14	2490	G	C4-N9-C1'	5.11	133.14	126.50
1	13	302	G	N1-C6-O6	-5.11	116.83	119.90
26	1H	958	U	N1-C2-N3	5.11	117.97	114.90
1	1G	331	G	N1-C6-O6	5.11	122.97	119.90
26	14	1998	G	C2-N3-C4	-5.11	109.35	111.90
23	2K	68	C	C6-N1-C2	-5.11	118.26	120.30
26	1H	191	A	O5'-P-OP1	5.11	116.83	110.70
26	1H	2566	A	P-O3'-C3'	5.11	125.83	119.70
26	14	113	G	N1-C6-O6	5.11	122.96	119.90
26	14	1682	G	O5'-P-OP2	-5.11	101.10	105.70
1	13	843	U	C5-C6-N1	5.11	125.25	122.70
26	1H	1574	C	N3-C4-C5	5.11	123.94	121.90
1	13	1503	A	C8-N9-C4	5.10	107.84	105.80
26	1H	1423	G	O5'-P-OP2	-5.10	101.11	105.70
47	H8	128	VAL	C-N-CA	-5.10	108.94	121.70
26	1H	682	G	C4-N9-C1'	5.10	133.13	126.50
26	1H	2330	G	C6-C5-N7	-5.10	127.34	130.40
33	51	156	ALA	C-N-CA	5.10	134.46	121.70
26	14	1379	A	C8-N9-C4	-5.10	103.76	105.80
26	1H	754	C	C5-C6-N1	-5.10	118.45	121.00
26	1H	821	A	C8-N9-C4	-5.10	103.76	105.80
42	C8	27	LEU	CA-CB-CG	5.10	127.03	115.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	2279	G	C5-C6-O6	5.10	131.66	128.60
26	14	842	G	N1-C6-O6	-5.10	116.84	119.90
26	14	2286	A	C8-N9-C4	-5.10	103.76	105.80
26	1H	590	A	C5-C6-N6	-5.10	119.62	123.70
26	1H	686	G	N3-C2-N2	5.10	123.47	119.90
26	1H	1603	A	P-O3'-C3'	5.10	125.82	119.70
27	16	54	G	N7-C8-N9	5.10	115.65	113.10
26	1H	909	A	C2-N3-C4	5.10	113.15	110.60
26	1H	1348	G	N9-C1'-C2'	-5.10	106.39	112.00
26	1H	1401	G	N7-C8-N9	5.10	115.65	113.10
26	14	1930	G	C8-N9-C1'	5.10	133.62	127.00
1	13	833	U	C2-N1-C1'	-5.09	111.59	117.70
26	1H	1931	U	C4-C5-C6	5.09	122.76	119.70
26	1H	2336	A	O5'-P-OP1	-5.09	101.12	105.70
26	14	863	A	O5'-P-OP2	-5.09	101.12	105.70
26	14	866	A	N9-C4-C5	-5.09	103.76	105.80
26	14	1617	C	C5-C6-N1	-5.09	118.45	121.00
26	14	2726	U	C5-C4-O4	5.09	128.96	125.90
1	13	978	A	N1-C6-N6	5.09	121.66	118.60
26	1H	2266	A	C6-N1-C2	-5.09	115.54	118.60
26	1H	2326	C	C6-N1-C2	-5.09	118.26	120.30
26	1H	2506	U	N3-C2-O2	-5.09	118.64	122.20
26	14	308	G	O5'-P-OP2	-5.09	101.12	105.70
26	14	1776	G	N3-C4-C5	-5.09	126.05	128.60
26	1H	1298	C	OP2-P-O3'	-5.09	94.00	105.20
26	1H	1634	A	OP1-P-OP2	5.09	127.24	119.60
26	1H	2827	C	C5-C4-N4	-5.09	116.64	120.20
26	14	186	G	C8-N9-C4	5.09	108.44	106.40
26	14	871	U	OP1-P-O3'	5.09	116.40	105.20
1	13	356	A	O4'-C1'-N9	5.09	112.27	108.20
26	1H	838	C	OP2-P-O3'	5.09	116.40	105.20
26	14	90	U	N3-C2-O2	-5.09	118.64	122.20
26	14	1215	G	C8-N9-C4	-5.09	104.36	106.40
26	14	2442	C	C2-N3-C4	-5.09	117.36	119.90
26	14	2552	U	N3-C4-O4	5.09	122.96	119.40
1	13	703	G	N3-C4-N9	5.09	129.05	126.00
26	14	2390	U	N1-C2-N3	5.09	117.95	114.90
26	14	2433	A	N9-C4-C5	-5.09	103.77	105.80
26	14	2733	A	C8-N9-C4	-5.09	103.77	105.80
27	1J	7	G	C6-C5-N7	-5.09	127.35	130.40
26	1H	330	A	O5'-P-OP2	-5.09	101.12	105.70
26	1H	513	A	N1-C6-N6	-5.09	115.55	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	733	G	C2-N3-C4	-5.09	109.36	111.90
26	1H	1496	A	C5-C6-N6	-5.09	119.63	123.70
26	1H	2440	C	OP1-P-O3'	5.09	116.39	105.20
1	1G	1374	A	O4'-C1'-N9	5.09	112.27	108.20
26	1H	2244	U	C6-N1-C2	-5.08	117.95	121.00
1	13	713	G	C8-N9-C4	5.08	108.43	106.40
26	1H	1626	G	N1-C2-N2	5.08	120.78	116.20
26	1H	1993	U	C2-N3-C4	-5.08	123.95	127.00
26	1H	2311	A	N1-C6-N6	5.08	121.65	118.60
26	1H	2586	C	C5-C4-N4	-5.08	116.64	120.20
26	1H	2618	G	O5'-P-OP2	-5.08	101.12	105.70
26	14	1424	G	N1-C6-O6	5.08	122.95	119.90
27	1J	102	G	C5-C6-O6	5.08	131.65	128.60
26	1H	532	A	C5-N7-C8	-5.08	101.36	103.90
26	1H	832	G	C5-C6-N1	-5.08	108.96	111.50
26	1H	983	A	OP2-P-O3'	5.08	116.38	105.20
26	1H	1025	G	C4-C5-N7	-5.08	108.77	110.80
26	1H	1951	U	O5'-P-OP2	5.08	116.80	110.70
26	1H	1998	G	N7-C8-N9	-5.08	110.56	113.10
26	1H	2023	G	C8-N9-C4	-5.08	104.37	106.40
26	1H	2726	U	C5-C6-N1	-5.08	120.16	122.70
26	14	503	A	C5-C6-N6	5.08	127.77	123.70
26	14	521	G	OP1-P-OP2	-5.08	111.98	119.60
1	13	1468	A	C6-N1-C2	-5.08	115.55	118.60
26	1H	837	C	N3-C4-C5	-5.08	119.87	121.90
26	1H	2047	U	N3-C4-O4	-5.08	115.84	119.40
26	1H	2287	A	C6-N1-C2	5.08	121.65	118.60
41	B8	105	LEU	CA-CB-CG	5.08	126.98	115.30
26	14	19	C	C6-N1-C2	5.08	122.33	120.30
26	14	2724	C	N1-C2-O2	-5.08	115.85	118.90
30	29	144	ARG	NE-CZ-NH1	5.08	122.84	120.30
1	13	723	U	C5-C6-N1	5.08	125.24	122.70
23	2K	6	G	C2-N3-C4	-5.08	109.36	111.90
26	1H	634	C	O5'-P-OP2	-5.08	101.13	105.70
26	1H	712	G	N1-C6-O6	5.08	122.95	119.90
26	1H	1609	A	C5-C6-N6	-5.08	119.64	123.70
26	1H	1936	A	C5-N7-C8	-5.08	101.36	103.90
26	1H	2618	G	C4-C5-N7	-5.08	108.77	110.80
26	1H	141(A)	C	N3-C4-C5	5.08	123.93	121.90
26	1H	410	G	N3-C2-N2	-5.08	116.35	119.90
26	1H	114	U	OP1-P-O3'	5.08	116.36	105.20
26	1H	747	U	OP1-P-OP2	5.08	127.21	119.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	1630	G	O5'-P-OP2	5.08	116.79	110.70
26	1H	2350	C	N3-C2-O2	-5.08	118.35	121.90
26	1H	2387	U	OP1-P-O3'	-5.08	94.03	105.20
54	O8	21	TYR	C-N-CA	5.08	134.39	121.70
26	14	24	G	N3-C4-N9	-5.08	122.95	126.00
26	14	1758	G	O5'-P-OP1	-5.08	101.13	105.70
26	14	940	G	O5'-P-OP2	-5.07	101.13	105.70
26	1H	808	G	N1-C6-O6	-5.07	116.86	119.90
26	1H	2870	C	OP2-P-O3'	5.07	116.36	105.20
1	1G	909	A	C5-C6-N6	-5.07	119.64	123.70
26	14	1675	C	OP1-P-O3'	5.07	116.36	105.20
26	1H	49	A	N9-C4-C5	-5.07	103.77	105.80
26	1H	98	G	OP1-P-OP2	5.07	127.20	119.60
26	1H	1770	G	OP1-P-O3'	5.07	116.36	105.20
1	1G	1397	C	N1-C2-O2	5.07	121.94	118.90
26	14	470	A	N9-C4-C5	-5.07	103.77	105.80
26	14	2489	G	OP2-P-O3'	5.07	116.36	105.20
26	14	2729	G	N9-C4-C5	-5.07	103.37	105.40
26	1H	762	U	C6-N1-C1'	-5.07	114.10	121.20
26	14	396	G	C8-N9-C1'	-5.07	120.41	127.00
26	1H	237	C	C4-C5-C6	5.07	119.93	117.40
26	1H	1797	C	C2-N3-C4	-5.07	117.37	119.90
26	1H	1964	G	O4'-C1'-N9	-5.07	104.15	108.20
1	1G	1145	C	P-O3'-C3'	5.07	125.78	119.70
1	1G	1508	G	O5'-P-OP1	-5.07	101.14	105.70
26	14	1258	C	C5-C6-N1	5.07	123.53	121.00
26	14	1806	C	O5'-P-OP2	-5.07	101.14	105.70
1	13	963	G	N1-C2-N3	5.07	126.94	123.90
23	2L	35	C	C2-N1-C1'	5.07	124.37	118.80
26	14	130	C	C6-N1-C2	5.07	122.33	120.30
26	14	1446	C	N1-C2-O2	5.07	121.94	118.90
26	14	2576	G	C2-N3-C4	5.07	114.43	111.90
26	14	140	A	C2-N3-C4	-5.06	108.07	110.60
26	14	1432	C	N1-C2-O2	-5.06	115.86	118.90
26	14	2062	A	N9-C4-C5	-5.06	103.77	105.80
26	14	2281	C	OP1-P-O3'	5.06	116.34	105.20
26	1H	192	C	N3-C2-O2	5.06	125.44	121.90
26	1H	377	C	C6-N1-C2	5.06	122.33	120.30
26	1H	1678	G	O4'-C1'-N9	-5.06	104.15	108.20
26	1H	2227	A	N1-C6-N6	-5.06	115.56	118.60
26	14	782	A	C8-N9-C4	5.06	107.83	105.80
26	14	1283	G	N3-C4-C5	-5.06	126.07	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	1400	G	O5'-P-OP1	5.06	116.78	110.70
26	14	2032	G	O4'-C1'-N9	-5.06	104.15	108.20
26	14	2211	G	C6-C5-N7	-5.06	127.36	130.40
23	2K	42	C	O5'-P-OP2	-5.06	101.14	105.70
20	BA	11	SER	C-N-CA	5.06	134.35	121.70
26	1H	695	G	N1-C2-N2	-5.06	111.65	116.20
1	1G	518	C	O5'-P-OP2	-5.06	101.15	105.70
26	14	858	U	C6-N1-C2	-5.06	117.96	121.00
26	1H	692	C	C5-C6-N1	-5.06	118.47	121.00
26	1H	1325	G	N1-C6-O6	5.06	122.94	119.90
26	1H	2584	U	N1-C2-N3	5.06	117.93	114.90
26	14	24	G	N3-C4-C5	5.06	131.13	128.60
26	14	472	A	N9-C4-C5	5.06	107.82	105.80
26	14	639	U	N3-C2-O2	-5.06	118.66	122.20
26	14	1395	A	O4'-C1'-N9	5.06	112.25	108.20
26	14	1462	C	C6-N1-C2	-5.06	118.28	120.30
26	14	1777	U	P-O3'-C3'	5.06	125.77	119.70
26	14	691	C	C4-C5-C6	5.06	119.93	117.40
26	14	2250	G	O5'-P-OP1	-5.06	101.15	105.70
1	13	1475	G	N7-C8-N9	5.05	115.63	113.10
26	1H	984	A	OP1-P-O3'	5.05	116.32	105.20
1	1G	700	G	C8-N9-C1'	5.05	133.57	127.00
26	14	929	G	N7-C8-N9	5.05	115.63	113.10
1	13	353	A	OP2-P-O3'	5.05	116.32	105.20
23	2K	48	U	P-O3'-C3'	5.05	125.76	119.70
26	1H	698	C	C5-C4-N4	-5.05	116.66	120.20
26	1H	2101	G	C5-C6-N1	-5.05	108.97	111.50
1	13	346	G	C4-C5-N7	5.05	112.82	110.80
26	1H	987	G	O5'-P-OP1	-5.05	101.15	105.70
26	1H	1594	G	N3-C2-N2	-5.05	116.36	119.90
26	14	229	A	O4'-C1'-N9	5.05	112.24	108.20
26	14	1660	C	N3-C4-C5	5.05	123.92	121.90
26	14	2329	G	O5'-P-OP1	5.05	116.76	110.70
26	1H	804	A	O4'-C1'-N9	5.05	112.24	108.20
26	1H	2826	A	N7-C8-N9	-5.05	111.28	113.80
26	14	2307	G	C8-N9-C4	-5.05	104.38	106.40
26	14	2859	G	P-O3'-C3'	5.05	125.76	119.70
26	1H	688	U	C2-N3-C4	-5.05	123.97	127.00
30	21	195	LEU	CA-CB-CG	5.05	126.91	115.30
26	14	1674	G	N3-C4-C5	-5.05	126.08	128.60
26	14	2248	C	C5-C4-N4	5.05	123.73	120.20
33	59	92	ILE	CG1-CB-CG2	-5.05	100.29	111.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
1	13	890	G	O4'-C1'-N9	5.05	112.24	108.20
1	13	1259	C	N1-C2-O2	5.05	121.93	118.90
26	1H	74	A	N3-C4-N9	-5.05	123.36	127.40
26	1H	241	A	N1-C2-N3	5.05	131.82	129.30
26	1H	528	A	C4-C5-C6	-5.05	114.48	117.00
26	1H	866	A	C8-N9-C1'	-5.05	118.62	127.70
26	1H	2257	U	C5-C6-N1	-5.05	120.18	122.70
26	14	468	G	OP1-P-OP2	-5.05	112.03	119.60
26	1H	25	U	C5-C4-O4	-5.04	122.87	125.90
26	1H	1626	G	N3-C4-N9	-5.04	122.97	126.00
26	1H	2422	A	C8-N9-C4	-5.04	103.78	105.80
26	14	2437	U	OP1-P-OP2	5.04	127.17	119.60
1	13	320	C	C6-N1-C2	5.04	122.32	120.30
26	1H	51	G	N1-C6-O6	-5.04	116.87	119.90
26	1H	566	U	OP1-P-O3'	5.04	116.30	105.20
26	1H	684	G	C5-C6-N1	5.04	114.02	111.50
26	1H	732	C	C2-N3-C4	-5.04	117.38	119.90
26	1H	1235	G	C8-N9-C4	-5.04	104.38	106.40
26	1H	2598	A	C5-C6-N6	-5.04	119.67	123.70
1	1G	1514	C	N1-C2-O2	-5.04	115.87	118.90
26	14	961	C	N1-C2-O2	-5.04	115.87	118.90
26	14	2355	C	C6-N1-C1'	-5.04	114.75	120.80
26	14	2685	G	N1-C2-N2	5.04	120.74	116.20
1	13	1430	C	O5'-P-OP1	-5.04	101.16	105.70
1	13	1482	G	O5'-P-OP2	-5.04	101.16	105.70
26	1H	684	G	N3-C4-C5	-5.04	126.08	128.60
26	1H	1153	C	N1-C2-O2	-5.04	115.88	118.90
26	1H	1644	C	N3-C2-O2	-5.04	118.37	121.90
26	1H	2047	U	N1-C2-O2	5.04	126.33	122.80
26	1H	2287	A	C6-C5-N7	-5.04	128.77	132.30
27	16	81	G	C2-N3-C4	-5.04	109.38	111.90
26	14	1827	C	C5-C6-N1	-5.04	118.48	121.00
26	14	1953	A	O5'-P-OP2	5.04	116.75	110.70
26	14	2426	A	C5-N7-C8	-5.04	101.38	103.90
26	1H	586	A	OP1-P-O3'	5.04	116.29	105.20
26	1H	764	A	OP1-P-O3'	5.04	116.29	105.20
26	14	49	A	OP2-P-O3'	5.04	116.29	105.20
26	14	1496	A	N1-C6-N6	5.04	121.62	118.60
26	14	1831	G	C2-N3-C4	-5.04	109.38	111.90
1	13	1434	A	N7-C8-N9	-5.04	111.28	113.80
26	1H	446	G	O5'-P-OP1	-5.04	101.17	105.70
26	1H	476	G	O5'-P-OP2	-5.04	101.17	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	829	A	N1-C6-N6	5.04	121.62	118.60
26	14	1601	G	OP1-P-O3'	5.04	116.28	105.20
26	14	1638	C	OP1-P-O3'	-5.04	94.11	105.20
27	1J	7	G	C4-C5-N7	5.04	112.81	110.80
26	1H	82	G	C5-C6-O6	5.04	131.62	128.60
26	1H	1204	A	N1-C2-N3	5.04	131.82	129.30
26	1H	1616	A	C2-N3-C4	-5.04	108.08	110.60
26	1H	2639	A	C2-N3-C4	-5.04	108.08	110.60
26	14	2277	G	C4-C5-N7	-5.04	108.78	110.80
1	13	405	U	C5-C6-N1	5.04	125.22	122.70
26	1H	244	A	C5-C6-N6	-5.04	119.67	123.70
26	1H	1029	A	N1-C6-N6	5.04	121.62	118.60
26	1H	1198	U	C5-C6-N1	-5.04	120.18	122.70
26	1H	1268	A	C2-N3-C4	-5.04	108.08	110.60
27	16	9	G	OP2-P-O3'	5.04	116.28	105.20
26	14	828	U	C6-N1-C2	-5.04	117.98	121.00
1	13	277	C	OP2-P-O3'	5.03	116.27	105.20
26	1H	757	U	C5-C6-N1	-5.03	120.18	122.70
26	1H	1306	C	C6-N1-C2	5.03	122.31	120.30
26	1H	1761	C	N3-C2-O2	5.03	125.42	121.90
26	1H	2600	A	C6-N1-C2	-5.03	115.58	118.60
1	1G	1512	U	O5'-P-OP2	-5.03	101.17	105.70
26	14	2611	U	OP2-P-O3'	5.03	116.27	105.20
1	13	1331	G	O5'-P-OP2	-5.03	101.17	105.70
27	16	5	C	C5-C4-N4	-5.03	116.68	120.20
26	14	1992	G	N1-C6-O6	-5.03	116.88	119.90
1	13	280	C	C6-N1-C2	5.03	122.31	120.30
26	1H	746	A	O5'-P-OP1	-5.03	101.17	105.70
26	1H	2240	C	OP1-P-O3'	5.03	116.27	105.20
26	14	1344	G	N1-C6-O6	5.03	122.92	119.90
26	14	2037	G	N1-C6-O6	-5.03	116.88	119.90
27	1J	43	C	C6-N1-C2	-5.03	118.29	120.30
26	1H	113	G	C4-N9-C1'	-5.03	119.96	126.50
26	1H	528	A	N1-C2-N3	-5.03	126.79	129.30
26	1H	651	G	OP1-P-OP2	-5.03	112.06	119.60
26	1H	945	A	C6-N1-C2	-5.03	115.58	118.60
26	14	1969	A	O5'-P-OP2	5.03	116.73	110.70
27	1J	6	C	C5-C6-N1	-5.03	118.48	121.00
1	13	1498	U	C2-N1-C1'	5.03	123.73	117.70
26	1H	1156	A	O4'-C1'-N9	-5.03	104.18	108.20
24	3L	58	A	OP1-P-O3'	5.03	116.26	105.20
26	14	786	C	OP2-P-O3'	5.03	116.26	105.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	14	1571	A	N1-C6-N6	5.03	121.62	118.60
26	14	1615	C	N1-C2-O2	5.03	121.92	118.90
1	13	317	G	N3-C4-N9	5.03	129.02	126.00
1	13	960	U	C6-N1-C2	-5.03	117.98	121.00
26	1H	2301	C	C6-N1-C2	-5.03	118.29	120.30
26	14	278	A	OP1-P-O3'	5.03	116.26	105.20
26	14	1968	G	C4-C5-N7	5.03	112.81	110.80
26	1H	2067	G	C5-C6-O6	5.02	131.61	128.60
1	1G	1502	A	C2-N3-C4	-5.02	108.09	110.60
26	14	960	A	N1-C6-N6	5.02	121.61	118.60
26	14	1255	U	N3-C2-O2	-5.02	118.68	122.20
26	1H	599	G	OP1-P-OP2	5.02	127.13	119.60
27	16	6	C	C6-N1-C2	5.02	122.31	120.30
26	14	671	C	C2-N3-C4	-5.02	117.39	119.90
26	14	768	G	O5'-P-OP2	-5.02	101.18	105.70
26	14	1598	C	OP1-P-OP2	-5.02	112.07	119.60
26	14	2211	G	N3-C4-N9	5.02	129.01	126.00
26	1H	781	A	N7-C8-N9	-5.02	111.29	113.80
26	14	801	G	C5-C6-O6	5.02	131.61	128.60
26	14	2595	G	C8-N9-C4	5.02	108.41	106.40
26	1H	113	G	C6-C5-N7	5.02	133.41	130.40
26	1H	205	G	O5'-P-OP2	-5.02	101.18	105.70
26	1H	1605	C	C2-N3-C4	-5.02	117.39	119.90
26	1H	1835	G	C4-N9-C1'	5.02	133.03	126.50
26	1H	2197	U	OP2-P-O3'	5.02	116.24	105.20
1	1G	449	C	N3-C4-N4	-5.02	114.49	118.00
1	1G	889	A	N1-C6-N6	5.02	121.61	118.60
26	14	725	G	N1-C6-O6	5.02	122.91	119.90
26	14	1646	C	OP1-P-O3'	5.02	116.24	105.20
26	14	1918	A	C8-N9-C4	5.02	107.81	105.80
1	13	1214	C	C6-N1-C2	5.02	122.31	120.30
26	1H	66	C	C5-C6-N1	5.02	123.51	121.00
26	1H	648	G	O5'-P-OP2	-5.02	101.18	105.70
26	1H	649	G	C8-N9-C4	-5.02	104.39	106.40
26	1H	1608	A	C2-N3-C4	-5.02	108.09	110.60
26	14	1655	A	C8-N9-C4	5.02	107.81	105.80
26	14	2003	G	C6-C5-N7	-5.02	127.39	130.40
26	1H	247	G	N9-C4-C5	-5.02	103.39	105.40
26	1H	508	G	C8-N9-C1'	-5.02	120.48	127.00
26	1H	1905	C	C2-N3-C4	5.02	122.41	119.90
26	14	1021	A	C5-C6-N1	-5.02	115.19	117.70
1	13	871	U	P-O3'-C3'	5.01	125.72	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	80	G	N7-C8-N9	5.01	115.61	113.10
26	1H	2819	G	C5-C6-O6	-5.01	125.59	128.60
26	1H	2827	C	C2-N3-C4	-5.01	117.39	119.90
26	14	1274	A	N1-C6-N6	5.01	121.61	118.60
1	1G	700	G	N9-C4-C5	5.01	107.41	105.40
26	14	1569	A	C8-N9-C4	-5.01	103.80	105.80
26	1H	1831	G	C8-N9-C4	-5.01	104.39	106.40
26	1H	2265	U	O5'-P-OP2	5.01	116.71	110.70
26	1H	2501	C	OP2-P-O3'	5.01	116.22	105.20
1	1G	197	A	C5-N7-C8	-5.01	101.39	103.90
26	14	102	G	C4-N9-C1'	-5.01	119.98	126.50
26	14	210	C	C6-N1-C2	5.01	122.31	120.30
26	14	733	G	C6-C5-N7	-5.01	127.39	130.40
26	14	747	U	OP1-P-O3'	5.01	116.23	105.20
26	14	2585	U	N1-C2-O2	5.01	126.31	122.80
1	13	346	G	C6-C5-N7	-5.01	127.39	130.40
26	1H	197	A	C2-N3-C4	-5.01	108.09	110.60
26	1H	994	C	O5'-P-OP1	-5.01	101.19	105.70
26	1H	1930	G	C6-N1-C2	5.01	128.10	125.10
26	1H	2269	A	N9-C4-C5	-5.01	103.80	105.80
26	14	801	G	C4-N9-C1'	-5.01	119.99	126.50
26	1H	232	G	N9-C4-C5	-5.01	103.40	105.40
1	1G	317	G	N1-C6-O6	5.01	122.91	119.90
26	14	446	G	N1-C6-O6	5.01	122.91	119.90
26	14	2447	G	C5-C6-O6	-5.01	125.59	128.60
26	1H	575	A	O5'-P-OP2	5.01	116.71	110.70
26	1H	1303	G	N3-C2-N2	5.01	123.41	119.90
26	1H	1772	G	N9-C1'-C2'	-5.01	106.49	112.00
26	1H	2302	G	N1-C6-O6	-5.01	116.90	119.90
26	14	2346	A	C1'-O4'-C4'	-5.01	105.90	109.90
26	1H	1625	C	N1-C2-O2	5.00	121.90	118.90
26	1H	1914	C	N1-C2-O2	5.00	121.90	118.90
1	13	809	G	N1-C6-O6	-5.00	116.90	119.90
22	1K	69	A	P-O3'-C3'	5.00	125.70	119.70
26	1H	1644	C	N3-C4-C5	5.00	123.90	121.90
26	1H	1824	G	C5-C6-O6	-5.00	125.60	128.60
26	1H	2033	A	C5-C6-N1	5.00	120.20	117.70
26	1H	2268	A	N9-C4-C5	-5.00	103.80	105.80
27	16	85	G	C5-C6-O6	-5.00	125.60	128.60
26	14	1131	G	O4'-C1'-N9	5.00	112.20	108.20
26	14	2679	A	C8-N9-C4	5.00	107.80	105.80
1	13	274	A	N1-C6-N6	-5.00	115.60	118.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
26	1H	202	U	C6-N1-C1'	-5.00	114.20	121.20
26	1H	577	G	OP1-P-OP2	-5.00	112.10	119.60
26	1H	1257	C	N1-C2-N3	5.00	122.70	119.20
26	1H	1797	C	C5-C6-N1	-5.00	118.50	121.00
26	1H	2018	G	N1-C6-O6	5.00	122.90	119.90
26	1H	2448	A	N9-C4-C5	-5.00	103.80	105.80
26	14	211	A	C4-C5-N7	5.00	113.20	110.70
26	14	308	G	C5-C6-O6	-5.00	125.60	128.60
26	14	800	A	O5'-P-OP1	-5.00	101.20	105.70
26	14	2038	G	C8-N9-C4	5.00	108.40	106.40
26	14	2433	A	N7-C8-N9	5.00	116.30	113.80

There are no chirality outliers.

All (145) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
29	11	113	VAL	Peptide
29	11	114	GLY	Peptide
29	11	122	ASP	Peptide
29	11	197	GLY	Peptide
29	11	237	GLU	Peptide
2	12	15	VAL	Peptide
2	12	19	HIS	Peptide
2	12	199	TYR	Peptide
2	12	219	VAL	Peptide
2	12	22	LYS	Peptide
29	19	197	GLY	Peptide
29	19	27	THR	Peptide
29	19	28	GLU	Peptide
10	1A	55	LYS	Peptide
2	1E	15	VAL	Peptide
2	1E	194	PRO	Peptide
2	1E	234	PRO	Peptide
2	1E	236	TYR	Peptide
30	21	153	GLY	Peptide
30	21	186	GLY	Peptide
30	21	187	ALA	Peptide
30	21	56	PRO	Peptide
30	21	58	ARG	Peptide
30	21	66	HIS	Peptide
30	21	68	ALA	Peptide
30	21	82	ARG	Peptide

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Mol	Chain	Res	Type	Group
30	29	117	MET	Peptide
30	29	186	GLY	Peptide
30	29	201	THR	Peptide
30	29	53	PRO	Peptide
30	29	61	ARG	Peptide
11	2A	49	GLY	Peptide
4	32	152	SER	Peptide
4	32	179	GLU	Peptide
37	35	110	TYR	Peptide
37	35	70	GLN	Peptide
31	39	127	GLU	Peptide
31	39	146	ALA	Peptide
31	39	166	ALA	Peptide
31	39	20	LEU	Peptide
31	39	24	LEU	Peptide
31	39	26	ALA	Peptide
31	39	89	VAL	Peptide
12	3A	18	VAL	Peptide
12	3I	47	LYS	Peptide
12	3I	87	GLY	Peptide
32	41	95	ARG	Peptide
38	45	58	PHE	Peptide
38	45	78	PRO	Peptide
32	49	13	GLU	Peptide
32	49	142	PRO	Peptide
13	4A	94	ARG	Peptide
13	4I	105	THR	Peptide
33	51	137	ASP	Peptide
33	51	152	ARG	Peptide
33	51	156	ALA	Peptide
33	51	170	ARG	Peptide
33	51	7	LEU	Peptide
39	55	106	GLY	Peptide
35	58	56	ASN	Peptide
33	59	90	LYS	Peptide
14	5A	27	CYS	Peptide
34	61	11	ASN	Peptide
34	61	134	PRO	Peptide
34	61	82	ARG	Peptide
40	65	53	SER	Peptide
34	69	101	LEU	Peptide
34	69	112	LYS	Peptide

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Mol	Chain	Res	Type	Group
34	69	142	VAL	Peptide
34	69	143	SER	Peptide
41	75	10	VAL	Peptide
41	75	12	SER	Peptide
41	75	5	ALA	Peptide
37	78	11	GLY	Peptide
37	78	115	LEU	Peptide
37	78	15	ARG	Peptide
37	78	24	GLY	Peptide
37	78	36	LYS	Peptide
37	78	70	GLN	Peptide
9	82	117	HIS	Peptide
42	85	72	HIS	Peptide
42	85	90	VAL	Peptide
42	85	98	LEU	Peptide
42	85	99	ALA	Peptide
38	88	139	GLU	Peptide
38	88	23	GLY	Peptide
38	88	58	PHE	Peptide
38	88	59	ARG	Peptide
9	8E	110	GLU	Peptide
9	8E	4	TYR	Peptide
39	98	44	LEU	Peptide
39	98	8	ARG	Peptide
44	A5	43	GLY	Peptide
40	A8	107	GLU	Peptide
40	A8	108	GLY	Peptide
19	AA	4	SER	Peptide
19	AA	9	VAL	Peptide
19	AI	24	ALA	Peptide
45	B5	24	GLY	Peptide
45	B5	61	GLY	Peptide
41	B8	12	SER	Peptide
41	B8	4	GLY	Peptide
41	B8	58	ASN	Peptide
20	BA	101	GLY	Peptide
20	BA	72	LEU	Peptide
46	C5	82	PRO	Peptide
46	C5	91	GLU	Peptide
42	C8	90	VAL	Peptide
42	C8	95	LEU	Peptide
47	D5	175	VAL	Peptide

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Mol	Chain	Res	Type	Group
47	D5	60	GLU	Peptide
47	D5	61	LEU	Peptide
43	D8	36	PRO	Peptide
43	D8	44	LYS	Peptide
43	D8	48	GLY	Peptide
48	E5	32	ARG	Peptide
48	E5	33	ALA	Peptide
49	F5	81	LYS	Peptide
50	G5	15	LYS	Peptide
50	G5	43	GLN	Peptide
46	G8	53	PRO	Peptide
46	G8	54	LYS	Peptide
46	G8	80	GLY	Peptide
46	G8	84	ARG	Peptide
46	G8	93	GLY	Peptide
47	H8	117	LEU	Peptide
47	H8	143	GLY	Peptide
47	H8	158	PRO	Peptide
47	H8	165	VAL	Peptide
47	H8	59	LEU	Peptide
47	H8	63	ASP	Peptide
48	I8	44	ARG	Peptide
49	J8	84	GLY	Peptide
50	K8	17	SER	Peptide
50	K8	46	GLN	Peptide
56	M5	40	GLU	Peptide
56	M5	51	ALA	Peptide
52	M8	37	SER	Peptide
52	M8	40	HIS	Peptide
52	M8	42	PHE	Peptide
54	O8	44	ARG	Peptide
55	P8	45	ALA	Peptide
56	Q8	30	ARG	Peptide
56	Q8	49	VAL	Peptide
56	Q8	51	ALA	Peptide

5.2 Too-close contacts (i)

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	32223	0	16267	692	0
1	1G	32414	0	16360	752	0
2	12	1696	0	1730	89	0
2	1E	1874	0	1926	95	0
3	22	1537	0	1603	84	0
3	2E	1605	0	1668	54	0
4	32	1702	0	1765	93	0
4	3E	1698	0	1759	84	0
5	42	1141	0	1199	40	0
5	4E	1142	0	1204	40	0
6	52	842	0	857	19	0
6	5E	837	0	852	34	0
7	62	1110	0	1163	53	0
7	6E	1242	0	1286	51	0
8	72	1107	0	1165	49	0
8	7E	1115	0	1177	46	0
9	82	953	0	983	62	0
9	8E	1000	0	1031	63	0
10	1A	646	0	662	45	0
10	1I	749	0	767	42	0
11	2A	835	0	847	25	0
11	2I	823	0	833	29	0
12	3A	947	0	1033	45	0
12	3I	956	0	1046	35	0
13	4A	879	0	935	46	0
13	4I	933	0	992	57	0
14	5A	486	0	525	35	0
14	5I	486	0	524	29	0
15	6A	729	0	768	26	0
15	6I	729	0	768	23	0
16	7A	705	0	725	23	0
16	7I	700	0	720	45	0
17	8A	823	0	891	22	0
17	8I	823	0	891	34	0
18	9A	544	0	605	19	0
18	9I	549	0	607	21	0
19	AA	510	0	507	21	0
19	AI	661	0	683	43	0
20	BA	762	0	861	43	0
20	BI	746	0	843	44	0
21	1B	188	0	195	8	0
21	1F	199	0	208	9	0
22	1K	1542	0	790	25	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
23	2K	1646	0	845	18	0
23	2L	1646	0	845	25	0
24	1L	1401	0	713	22	0
24	3K	1483	0	756	50	0
24	3L	1528	0	778	42	0
25	4K	420	0	209	7	0
25	4L	419	0	208	17	0
26	14	60877	0	30690	1171	0
26	1H	61609	0	31058	1189	0
27	16	2617	0	1328	50	0
27	1J	2617	0	1328	76	0
28	71	1027	0	1043	57	0
29	11	2120	0	2197	100	0
29	19	2125	0	2199	105	0
30	21	1546	0	1602	86	0
30	29	1563	0	1629	93	0
31	31	1585	0	1632	70	0
31	39	1602	0	1649	88	0
32	41	1457	0	1514	68	0
32	49	1459	0	1507	65	0
33	51	1328	0	1396	64	0
33	59	1295	0	1366	58	0
34	61	1131	0	1218	36	0
34	69	1131	0	1218	49	0
35	15	1096	0	1168	47	0
35	58	1096	0	1169	48	0
36	25	932	0	996	46	0
36	68	932	0	996	41	0
37	35	1122	0	1206	75	0
37	78	1122	0	1206	80	0
38	45	1099	0	1154	73	0
38	88	1117	0	1168	58	0
39	55	967	0	1033	42	0
39	98	967	0	1033	45	0
40	65	876	0	938	63	0
40	A8	881	0	943	54	0
41	75	1109	0	1170	61	0
41	B8	1119	0	1177	72	0
42	85	959	0	1019	40	0
42	C8	950	0	1011	58	0
43	95	770	0	838	32	0
43	D8	774	0	849	40	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
44	A5	886	0	948	23	0
44	E8	876	0	941	30	0
45	B5	735	0	785	30	0
45	F8	750	0	814	19	0
46	C5	794	0	885	52	0
46	G8	783	0	869	49	0
47	D5	1411	0	1436	82	0
47	H8	1365	0	1391	57	0
48	E5	603	0	620	33	0
48	I8	611	0	631	34	0
49	F5	737	0	813	32	0
49	J8	747	0	817	42	0
50	G5	576	0	625	27	0
50	K8	575	0	634	42	0
51	H5	459	0	512	11	0
51	L8	459	0	512	13	0
52	M8	475	0	465	34	0
53	J5	434	0	454	22	0
53	N8	369	0	388	21	0
54	O8	389	0	404	26	0
55	L5	401	0	436	10	0
55	P8	401	0	436	11	0
56	M5	516	0	582	25	0
56	Q8	516	0	582	37	0
57	13	140	0	0	0	0
57	14	435	0	0	0	0
57	16	11	0	0	0	0
57	19	1	0	0	0	0
57	1G	102	0	0	0	0
57	1H	525	0	0	0	0
57	1J	8	0	0	0	0
57	21	3	0	0	0	0
57	25	1	0	0	0	0
57	29	1	0	0	0	0
57	2K	2	0	0	0	0
57	2L	2	0	0	0	0
57	31	2	0	0	0	0
57	32	1	0	0	0	0
57	35	1	0	0	0	0
57	39	2	0	0	0	0
57	3I	1	0	0	0	0
57	41	1	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
57	42	1	0	0	0	0
57	45	2	0	0	0	0
57	4I	1	0	0	0	0
57	4L	1	0	0	0	0
57	52	1	0	0	0	0
57	5I	1	0	0	0	0
57	78	1	0	0	0	0
57	7A	1	0	0	0	0
57	88	3	0	0	0	0
57	C5	1	0	0	0	0
57	E5	1	0	0	0	0
57	I8	2	0	0	0	0
57	M5	1	0	0	0	0
57	P8	1	0	0	0	0
57	Q8	1	0	0	0	0
58	13	42	0	43	0	0
58	1G	42	0	45	1	0
59	32	8	0	0	3	0
59	3E	8	0	0	1	0
60	5A	1	0	0	0	0
60	5I	1	0	0	0	0
60	C5	1	0	0	0	0
60	G8	1	0	0	0	0
61	14	26	0	45	8	0
61	1G	13	0	24	0	0
62	11	17	0	0	4	0
62	13	320	0	0	12	0
62	14	1144	0	0	65	0
62	15	1	0	0	0	0
62	16	12	0	0	1	0
62	19	15	0	0	3	0
62	1A	1	0	0	0	0
62	1F	1	0	0	0	0
62	1G	317	0	0	21	0
62	1H	1470	0	0	90	0
62	1I	2	0	0	0	0
62	1J	12	0	0	1	0
62	1K	2	0	0	0	0
62	21	7	0	0	1	0
62	25	6	0	0	0	0
62	29	4	0	0	0	0
62	2K	8	0	0	1	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
62	2L	6	0	0	0	0
62	31	5	0	0	0	0
62	32	1	0	0	0	0
62	35	8	0	0	1	0
62	39	5	0	0	1	0
62	3I	2	0	0	0	0
62	41	1	0	0	0	0
62	4E	3	0	0	0	0
62	4K	5	0	0	0	0
62	4L	6	0	0	0	0
62	52	4	0	0	0	0
62	55	3	0	0	0	0
62	58	2	0	0	0	0
62	5I	2	0	0	0	0
62	6A	2	0	0	0	0
62	6I	3	0	0	0	0
62	78	10	0	0	0	0
62	7A	5	0	0	0	0
62	7I	1	0	0	0	0
62	85	2	0	0	0	0
62	8E	2	0	0	0	0
62	9A	2	0	0	0	0
62	A5	1	0	0	0	0
62	B5	1	0	0	0	0
62	B8	1	0	0	0	0
62	BA	2	0	0	0	0
62	BI	2	0	0	1	0
62	C5	3	0	0	0	0
62	C8	3	0	0	0	0
62	F5	1	0	0	1	0
62	F8	2	0	0	0	0
62	G8	2	0	0	0	0
62	H5	2	0	0	1	0
62	I8	6	0	0	1	0
62	J8	4	0	0	0	0
62	L8	3	0	0	1	0
62	M5	9	0	0	1	0
62	P8	1	0	0	0	0
62	Q8	5	0	0	2	0
All	All	297444	0	197360	7486	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 16.

All (7486) 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)
10:1A:38:ILE:C	10:1A:39:PRO:N	1.71	1.44
38:45:27:VAL:HB	38:45:28:ALA:HA	1.19	1.13
29:11:182:LEU:H	29:11:272:ALA:HB3	1.23	1.02
37:78:63:PRO:HB2	56:Q8:30:ARG:HH21	1.23	1.01
26:1H:1496:A:H8	26:1H:1577:C:HO2'	1.00	0.99
26:14:330:A:H2	26:14:1210:A:HO2'	1.05	0.98
8:7E:87:SER:HB2	8:7E:93:VAL:HB	1.47	0.97
29:19:39:LYS:HG3	29:19:40:THR:H	1.31	0.96
29:19:44:ASN:HA	29:19:47:GLY:H	1.30	0.95
29:19:44:ASN:HB3	29:19:45:ASN:HA	1.48	0.94
1:13:1502:A:H2	1:13:1505:G:H1	1.14	0.94
19:AI:40:ILE:HG12	19:AI:41:VAL:HG13	1.49	0.94
26:1H:1689:A:H62	26:1H:1698:A:H2	1.08	0.94
31:31:101:LEU:HD23	31:31:102:PRO:HD2	1.51	0.93
26:1H:1359:A:N1	26:1H:1372:U:N3	2.17	0.93
26:1H:1771:C:HO2'	26:1H:1786:A:H8	0.95	0.92
26:14:2032:G:H21	30:29:146:THR:HG23	1.32	0.92
27:1J:80:U:H2'	27:1J:81:G:H21	1.36	0.91
26:1H:1055:G:N2	26:1H:1104:C:N3	2.18	0.91
26:1H:620:G:H4'	26:1H:621:A:H5"	1.54	0.90
26:14:67:U:H3	26:14:74:A:H2	1.19	0.90
30:29:60:ASN:HB2	30:29:62:PRO:HD2	1.53	0.90
47:D5:115:GLY:HA2	47:D5:177:PRO:HG2	1.52	0.90
26:1H:1456:G:OP2	62:1H:3742:HOH:O	1.89	0.90
26:1H:2308:G:H1	26:1H:2311:A:H2	1.11	0.89
4:3E:107:ARG:HH22	4:3E:194:LEU:HD22	1.37	0.89
1:1G:1368:G:H5'	9:82:112:LYS:HB3	1.56	0.88
30:29:54:GLN:HA	30:29:74:PRO:HA	1.55	0.88
26:1H:1653:G:H3'	39:98:2:ARG:HG3	1.52	0.88
47:H8:108:PRO:HB2	47:H8:112:ARG:HA	1.55	0.88
38:45:21:THR:HG22	38:45:23:GLY:HA3	1.56	0.88
13:4I:3:ARG:HB3	13:4I:9:ILE:HG12	1.55	0.88
26:14:676:A:H8	26:14:2069:G:H21	1.18	0.87
26:1H:442:G:H1'	31:31:48:THR:HG21	1.55	0.87
26:14:2873:A:H8	39:55:6:SER:H	1.17	0.87
26:1H:138:G:N2	45:F8:44:GLU:OE2	2.07	0.87
26:1H:607:U:H3	26:1H:621:A:H2	1.19	0.87
9:82:28:VAL:HG22	9:82:63:ILE:HB	1.56	0.87
41:B8:57:PHE:O	41:B8:58:ASN:ND2	2.08	0.87
26:1H:674:G:H1'	31:31:74:ARG:HD3	1.57	0.87

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:3K:7:U:H2'	24:3K:49:G:H5'	1.56	0.86
26:14:2624:G:N7	62:14:3631:HOH:O	2.09	0.86
1:13:975:A:H4'	1:13:976:G:H5"	1.57	0.86
31:39:25:PRO:HB2	31:39:27:GLU:H	1.38	0.86
26:14:123:G:N7	62:14:3630:HOH:O	2.08	0.86
26:1H:780:G:H21	26:1H:783:A:H62	1.18	0.86
1:13:1348:U:H3	1:13:1374:A:H2	1.24	0.86
1:1G:1028:C:N3	1:1G:1033:G:N2	2.22	0.86
31:39:122:LYS:HD2	31:39:191:ARG:HE	1.41	0.86
26:14:607:U:H3	26:14:621:A:H2	1.20	0.85
1:13:601:C:H2'	1:13:602:A:H8	1.42	0.85
1:1G:1502:A:H2	1:1G:1505:G:H1	1.23	0.85
38:45:27:VAL:CB	38:45:28:ALA:HA	2.06	0.85
26:14:1899:G:H21	26:14:1902:C:N4	1.75	0.85
26:1H:2032:G:H21	30:21:146:THR:HG23	1.40	0.84
48:E5:27:GLU:HG3	48:E5:68:GLU:HA	1.58	0.84
2:12:75:LYS:HA	2:12:78:GLN:HB2	1.59	0.84
26:1H:49:A:N7	26:1H:120:U:H5	1.75	0.84
38:45:26:TYR:CD1	38:45:27:VAL:HG23	2.13	0.84
26:14:1689:A:H62	26:14:1698:A:H2	1.24	0.84
26:1H:1678:G:H22	26:1H:1989:G:H22	1.24	0.84
26:14:483:A:H4'	46:C5:49:VAL:HA	1.60	0.84
26:1H:270(K):C:O2'	26:1H:270(N):G:N2	2.09	0.84
26:1H:1210:A:H8	26:1H:1210:A:H5'	1.42	0.84
13:4I:23:TYR:HD2	13:4I:67:GLU:HA	1.42	0.84
40:A8:78:LEU:HD12	40:A8:108:GLY:HA2	1.58	0.84
7:62:93:PRO:HD2	7:62:94:ARG:HH21	1.42	0.84
27:16:80:U:H2'	27:16:81:G:H21	1.43	0.83
26:1H:1049:C:N3	33:51:3:ARG:NH1	2.26	0.83
29:19:182:LEU:H	29:19:272:ALA:HB3	1.43	0.83
1:13:153:C:H42	1:13:168:G:H1	1.24	0.83
26:14:1757:U:H3	26:14:1762:A:H2	1.25	0.83
26:1H:2867:G:OP2	41:B8:119:LYS:NZ	2.09	0.83
3:22:135:LYS:NZ	3:22:135:LYS:O	2.11	0.83
24:3K:76:A:H8	26:1H:2394:C:H42	1.22	0.83
26:1H:2749:A:H5"	33:51:4:ILE:HD11	1.59	0.83
13:4I:10:PRO:HB2	13:4I:18:ALA:HB1	1.60	0.83
26:1H:67:U:H3	26:1H:74:A:H2	1.26	0.83
31:39:53:THR:HG22	31:39:56:GLU:HG3	1.58	0.83
2:12:91:PRO:HG2	2:12:155:LEU:HG	1.57	0.83
34:61:98:ALA:HB2	34:61:111:PRO:HB3	1.57	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:82:112:LYS:HA	9:82:119:ALA:HB2	1.58	0.82
26:14:84:A:N6	26:14:102:G:O2'	2.11	0.82
47:D5:27:VAL:HG12	47:D5:87:ASP:HB3	1.61	0.82
26:1H:2781:A:H5"	26:1H:2782:G:H5'	1.60	0.82
26:1H:847:U:OP2	62:1H:3743:HOH:O	1.96	0.82
10:1I:61:GLU:OE2	14:5I:45:ARG:NH1	2.13	0.82
1:13:1104:G:OP1	2:1E:144:ARG:NH2	2.11	0.82
26:14:2287:A:N6	26:14:2344:U:H3	1.77	0.82
4:32:23:GLY:N	4:32:26:CYS:SG	2.51	0.82
26:1H:1332:G:N2	26:1H:1609:A:O2'	2.13	0.81
34:69:81:VAL:H	34:69:143:SER:HB3	1.42	0.81
39:98:86:ARG:HH21	39:98:118:GLU:HG2	1.42	0.81
32:49:125:PHE:HB3	32:49:166:ASP:HB2	1.60	0.81
1:13:1305:G:H22	1:13:1331:G:H2'	1.43	0.81
26:14:1496:A:H8	26:14:1577:C:HO2'	1.28	0.81
26:14:2292:C:OP1	40:65:17:ARG:NH2	2.13	0.81
3:2E:16:ARG:HD2	3:2E:54:ARG:HH21	1.45	0.81
1:1G:1352:C:H42	1:1G:1370:G:H1	1.24	0.81
47:D5:157:LEU:HA	47:D5:161:VAL:HG11	1.59	0.81
26:1H:860:U:H5	26:1H:917:A:C2	1.97	0.81
26:14:958:U:OP2	38:45:14:ARG:NH1	2.11	0.81
26:14:152:G:H1	26:14:174:C:H42	1.28	0.81
4:3E:167:GLY:HA2	29:19:135:PHE:HE1	1.44	0.81
49:J8:92:LYS:HA	49:J8:95:LEU:HG	1.62	0.81
26:14:780:G:H21	26:14:783:A:H62	1.28	0.81
26:1H:1670:C:OP1	62:1H:3744:HOH:O	1.98	0.81
33:59:15:VAL:HG12	33:59:29:PRO:HD2	1.62	0.81
1:13:659:U:H2'	1:13:660:G:H8	1.45	0.81
26:1H:2418:A:N7	62:1H:3763:HOH:O	2.12	0.81
26:1H:270(I):G:H1	26:1H:270(Q):C:H42	1.29	0.81
30:29:81:ILE:HG22	30:29:82:ARG:H	1.46	0.81
1:13:1178:G:OP2	9:8E:93:ARG:NH2	2.14	0.81
47:H8:9:TYR:HE1	47:H8:35:ARG:HG2	1.45	0.81
48:I8:38:VAL:HG23	48:I8:59:LEU:HB2	1.63	0.81
54:O8:15:GLU:OE2	54:O8:44:ARG:NH2	2.14	0.80
29:11:84:TYR:HE1	29:11:86:PRO:HB3	1.46	0.80
26:14:1814:G:OP1	29:19:40:THR:HG21	1.81	0.80
38:88:138:ASP:N	38:88:138:ASP:OD1	2.13	0.80
26:1H:958:U:OP2	38:88:14:ARG:NH1	2.14	0.80
26:1H:1604:C:OP2	62:1H:3611:HOH:O	1.99	0.80
31:39:123:LEU:O	31:39:125:LEU:N	2.14	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1434:A:H61	26:1H:1558:A:N6	1.80	0.80
4:32:157:LEU:O	4:32:161:ASN:ND2	2.14	0.80
33:51:7:LEU:HD12	33:51:7:LEU:H	1.44	0.80
20:BI:71:THR:HG22	20:BI:72:LEU:H	1.47	0.80
26:14:1138:G:H21	35:15:106:MET:HE3	1.47	0.80
11:2A:29:ILE:HG22	11:2A:44:SER:HB2	1.64	0.80
51:L8:8:LEU:HB2	51:L8:28:LEU:HD22	1.64	0.80
26:1H:1899:G:H22	26:1H:1902:C:H5	1.30	0.80
1:13:812:C:N3	62:13:1840:HOH:O	2.13	0.79
1:1G:976:G:N2	1:1G:1362(A):C:OP2	2.15	0.79
17:8I:67:LYS:HA	17:8I:70:ARG:HH12	1.46	0.79
26:1H:1021:A:H62	26:1H:1141:U:H3	1.30	0.79
2:1E:208:ILE:HG22	2:1E:211:ILE:HD11	1.62	0.79
26:1H:2111:C:N4	26:1H:2147:G:O6	2.14	0.79
26:1H:2176:A:H1'	28:71:215:THR:HG21	1.65	0.79
50:K8:15:LYS:HZ2	50:K8:15:LYS:H	1.28	0.79
26:14:2681:C:H5	26:14:2725:A:H62	1.29	0.79
26:14:275:G:N2	26:14:276:A:N7	2.31	0.79
37:78:71:VAL:HG13	37:78:72:PRO:HD3	1.65	0.79
1:1G:1288:A:O2'	21:1B:10:ARG:NH2	2.15	0.79
26:1H:1900:A:H5'	26:1H:1900:A:H8	1.47	0.79
37:35:79:ARG:HG2	37:35:110:TYR:HB2	1.64	0.79
26:1H:1138:G:H21	35:58:106:MET:HE3	1.45	0.79
11:2I:54:ARG:O	11:2I:56:GLY:N	2.15	0.79
16:7A:53:VAL:HG13	16:7A:79:VAL:HG22	1.65	0.79
26:1H:1778:U:H2'	26:1H:1784:A:N6	1.97	0.78
11:2I:22:HIS:HB3	11:2I:29:ILE:HG23	1.64	0.78
38:45:27:VAL:HG22	38:45:137:TYR:O	1.83	0.78
48:I8:11:ARG:O	48:I8:14:ARG:NH2	2.16	0.78
51:L8:10:LYS:NZ	51:L8:15:TYR:OH	2.16	0.78
26:1H:676:A:H8	26:1H:2069:G:H21	1.30	0.78
13:4I:39:ILE:HD12	13:4I:56:LEU:HD23	1.65	0.78
25:4L:21:A:H2'	25:4L:22:A:H5"	1.63	0.78
24:3L:6:G:N1	24:3L:67:C:O2	2.15	0.78
29:19:69:ARG:NH2	29:19:128:GLY:O	2.17	0.78
20:BI:26:ASN:HB2	20:BI:71:THR:HG23	1.63	0.78
1:1G:1324:A:H4'	1:1G:1362:C:H4'	1.63	0.78
26:1H:1022:G:N2	26:1H:1023:U:O4	2.16	0.78
26:1H:2392:A:H2	26:1H:2424:C:H42	1.30	0.78
9:82:5:TYR:N	9:82:87:GLN:OE1	2.16	0.78
43:D8:44:LYS:O	43:D8:46:VAL:N	2.17	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2720:U:H3	26:14:2873:A:H2	1.30	0.78
26:1H:1382:G:O6	62:1H:3746:HOH:O	2.02	0.78
37:35:26:GLY:O	62:35:301:HOH:O	2.02	0.78
47:D5:74:VAL:HG13	47:D5:86:VAL:HG22	1.64	0.78
47:H8:116:VAL:HG22	47:H8:146:ILE:HG12	1.65	0.78
26:1H:2061:G:H5'	62:1H:4045:HOH:O	1.82	0.78
38:45:138:ASP:N	38:45:138:ASP:OD1	2.16	0.78
26:14:1678:G:N2	26:14:1989:G:H22	1.80	0.78
1:1G:1248:A:N3	9:82:70:LYS:NZ	2.29	0.78
30:29:91:VAL:HB	30:29:95:ILE:HD11	1.66	0.78
26:1H:607:U:OP1	31:31:102:PRO:HA	1.83	0.78
1:1G:523:A:H61	12:3A:92:ASP:HB2	1.48	0.77
23:2K:76:C:OP1	62:2K:202:HOH:O	2.01	0.77
13:4A:16:ASP:HB3	13:4A:34:LEU:HD11	1.66	0.77
1:13:1422:G:H5"	36:68:48:PRO:HB3	1.66	0.77
1:1G:975:A:H4'	1:1G:976:G:H5"	1.66	0.77
26:14:71:A:H2	45:B5:31:HIS:HE2	1.30	0.77
47:H8:165:VAL:HB	47:H8:167:PRO:HD3	1.65	0.77
29:11:235:GLY:O	62:11:301:HOH:O	2.01	0.77
1:13:143:A:H2	1:13:220:G:H1	1.33	0.77
26:1H:2447:G:O5'	62:1H:3745:HOH:O	2.01	0.77
7:62:148:ASN:ND2	7:62:148:ASN:O	2.16	0.77
1:13:1256:A:N6	1:13:1278:U:OP2	2.17	0.77
26:14:2357:U:OP1	48:E5:20:ARG:NH1	2.17	0.77
26:1H:2656:U:H3	26:1H:2665:A:H2	1.29	0.77
1:13:974:A:OP2	14:5I:41:ARG:NH1	2.17	0.77
26:14:2719:G:OP2	62:14:3625:HOH:O	2.01	0.77
10:1A:51:ARG:HB2	10:1A:60:ARG:HA	1.66	0.77
26:1H:1728:G:H8	26:1H:1732:A:H62	1.33	0.77
10:1A:28:ARG:HH21	10:1A:34:VAL:H	1.30	0.77
1:13:1305:G:N2	1:13:1331:G:H2'	1.99	0.77
12:3A:52:LEU:O	12:3A:54:LYS:NZ	2.17	0.77
5:42:43:LEU:HD22	5:42:136:MET:HG3	1.66	0.77
20:BA:53:LEU:HD12	20:BA:100:ILE:HG22	1.66	0.77
26:14:259:G:H21	26:14:621:A:H8	1.31	0.77
26:1H:1009:A:OP2	35:58:37:LYS:NZ	2.17	0.77
3:22:70:VAL:HG12	3:22:72:LYS:H	1.50	0.77
31:31:185:ASP:OD1	31:31:188:ARG:NH1	2.18	0.77
1:1G:1281:U:OP2	1:1G:1282:C:N4	2.17	0.77
42:85:49:HIS:HA	42:85:52:ARG:HB3	1.67	0.77
38:88:109:VAL:HG13	38:88:113:GLN:HB3	1.67	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2456:C:N4	62:14:3643:HOH:O	2.17	0.76
26:1H:993:G:OP1	42:C8:50:ARG:NH2	2.16	0.76
36:68:88:ASN:HD21	36:68:92:GLU:HB2	1.50	0.76
1:13:780:A:OP2	62:13:1837:HOH:O	2.02	0.76
26:1H:2469:A:H2	26:1H:2481:G:H21	1.33	0.76
22:1K:36:U:H3	25:4K:19:G:H1	1.33	0.76
12:3A:60:LEU:HB2	12:3A:64:TYR:HB2	1.67	0.76
24:3K:33:U:H2'	24:3K:34:U:H2'	1.67	0.76
41:B8:56:GLY:O	41:B8:59:THR:HG22	1.84	0.76
2:1E:16:HIS:HE1	2:1E:213:LEU:HD12	1.49	0.76
41:75:56:GLY:O	41:75:59:THR:HG23	1.85	0.76
43:95:98:GLU:OE1	43:95:100:ARG:NH1	2.16	0.76
53:N8:33:CYS:SG	53:N8:40:LYS:NZ	2.54	0.76
26:14:2392:A:H2	26:14:2424:C:H42	1.31	0.76
30:21:38:THR:HG22	30:21:40:GLU:H	1.51	0.76
7:6E:122:HIS:HA	7:6E:125:MET:HE2	1.68	0.76
26:14:761:A:N7	61:14:3437:SPE:H112	2.01	0.76
3:22:106:VAL:HB	3:22:109:PRO:HB3	1.68	0.76
30:29:50:GLY:HA2	30:29:78:LEU:HB3	1.68	0.76
46:G8:30:VAL:HG22	46:G8:37:VAL:HG12	1.67	0.76
26:1H:2334:G:O6	48:I8:74:ARG:NH2	2.18	0.76
31:39:157:VAL:HB	31:39:194:MET:HG3	1.67	0.76
24:3L:3:G:N2	24:3L:70:C:N3	2.32	0.76
6:52:76:ALA:HB1	6:52:80:ARG:HH21	1.49	0.76
46:G8:102:CYS:SG	46:G8:103:GLY:N	2.59	0.76
47:H8:19:ARG:NH1	47:H8:84:GLU:O	2.19	0.76
2:12:40:HIS:HD2	2:12:190:THR:HG21	1.50	0.76
26:14:1141:U:OP2	35:15:63:THR:OG1	2.03	0.76
26:1H:2580:U:H4'	30:21:130:GLY:HA3	1.68	0.76
26:14:1019:U:OP1	26:14:1035:U:O2'	2.02	0.76
26:14:907:U:O2'	38:45:101:ARG:NH2	2.17	0.76
26:1H:780:G:H21	26:1H:783:A:N6	1.83	0.76
41:75:50:ILE:HD11	41:75:102:ILE:HD11	1.68	0.76
46:C5:88:LYS:HG3	46:C5:89:PHE:H	1.48	0.76
1:13:524:G:H2'	1:13:525:C:C6	2.21	0.75
26:1H:2312:U:H5'	32:41:88:ILE:HD11	1.68	0.75
33:51:2:SER:HB2	33:51:3:ARG:HD3	1.67	0.75
26:14:889:C:H2'	26:14:890:A:H4'	1.69	0.75
1:1G:999:U:H3	1:1G:1041:A:H61	1.32	0.75
26:1H:2099:U:N3	26:1H:2190:G:O6	2.17	0.75
27:1J:80:U:H2'	27:1J:81:G:N2	2.01	0.75

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:4E:102:ALA:HB1	5:4E:106:PRO:HG2	1.68	0.75
42:C8:92:ARG:O	42:C8:94:ASN:N	2.19	0.75
1:1G:827:U:H3	1:1G:872:A:H62	1.32	0.75
26:1H:1332:G:OP1	62:1H:3747:HOH:O	2.04	0.75
30:21:53:PRO:HA	30:21:75:VAL:H	1.50	0.75
16:7I:47:ASP:N	16:7I:47:ASP:OD1	2.20	0.75
44:A5:13:SER:HB3	44:A5:16:LYS:HD2	1.67	0.75
10:1A:28:ARG:NH2	10:1A:34:VAL:O	2.18	0.75
24:3L:3:G:H1	24:3L:70:C:H42	1.31	0.75
26:14:6:A:H3'	26:14:7:G:H5'	1.69	0.75
29:19:228:PRO:O	62:19:401:HOH:O	2.04	0.75
30:21:197:ILE:HD11	30:21:199:ARG:HE	1.52	0.75
19:AI:41:VAL:O	52:M8:63:TYR:OH	2.05	0.75
2:1E:80:ILE:HG12	2:1E:212:GLN:HB2	1.69	0.75
1:1G:176:C:OP1	20:BA:29:LYS:NZ	2.17	0.75
26:1H:70:G:H21	26:1H:71:A:N6	1.85	0.75
30:21:74:PRO:HA	30:21:75:VAL:HB	1.67	0.75
48:I8:27:GLU:HG3	48:I8:68:GLU:HA	1.69	0.75
1:13:1007:C:H42	1:13:1022:G:H1	1.34	0.75
1:13:141:A:O2'	1:13:182:U:O2	2.04	0.75
26:14:2789:C:O2	26:14:2894:G:N2	2.20	0.75
26:1H:259:G:O2'	26:1H:621:A:O2'	2.05	0.75
31:39:25:PRO:HB3	31:39:28:ILE:HG23	1.68	0.75
43:D8:65:GLY:HA3	43:D8:91:TYR:CZ	2.21	0.75
26:14:270(W):G:N7	62:14:3651:HOH:O	2.20	0.75
1:1G:542:G:OP1	4:32:10:ARG:NH2	2.19	0.75
1:1G:576:G:N2	1:1G:759:A:OP1	2.20	0.75
8:7E:21:LYS:O	8:7E:65:TYR:OH	2.05	0.75
52:M8:13:ARG:HH12	52:M8:22:ILE:HG23	1.52	0.75
26:14:2226:C:OP2	62:14:3627:HOH:O	2.05	0.74
29:19:30:GLU:HG3	29:19:63:ARG:CZ	2.17	0.74
31:39:181:LEU:HD21	31:39:186:ILE:HD11	1.67	0.74
26:1H:1604:C:OP2	62:1H:3748:HOH:O	2.05	0.74
36:25:14:THR:HG21	36:25:86:ILE:HG13	1.69	0.74
24:3K:49:G:H1'	24:3K:66:A:C6	2.22	0.74
35:58:96:GLU:O	35:58:98:VAL:N	2.19	0.74
1:1G:793:U:OP1	62:1G:1854:HOH:O	2.04	0.74
26:1H:1016:G:N7	62:1H:3777:HOH:O	2.20	0.74
15:6I:26:GLU:OE2	15:6I:77:ARG:NH1	2.20	0.74
56:M5:40:GLU:HA	56:M5:43:GLN:HB2	1.67	0.74
1:1G:54:C:N4	1:1G:353:A:OP2	2.20	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:8E:121:ARG:NH1	9:8E:122:ALA:O	2.19	0.74
50:K8:4:SER:H	50:K8:7:ARG:H	1.34	0.74
40:A8:48:LEU:HD23	40:A8:82:ILE:HD11	1.68	0.74
2:12:184:VAL:HG23	2:12:198:ASP:H	1.53	0.74
26:14:1864:U:OP1	26:14:2410:G:O2'	2.06	0.74
28:71:7:TYR:HA	28:71:10:LEU:HB2	1.68	0.74
53:N8:40:LYS:HE2	53:N8:47:PRO:HD2	1.68	0.74
33:51:10:PRO:HD2	33:51:50:VAL:O	1.87	0.74
28:71:57:ASN:HA	28:71:165:ASN:HD21	1.50	0.74
50:G5:50:ILE:HD12	50:G5:51:ARG:H	1.52	0.74
50:K8:42:GLY:O	50:K8:44:LEU:N	2.20	0.74
26:14:2255:G:OP2	62:14:3626:HOH:O	2.04	0.74
26:1H:1021:A:H8	26:1H:1022:G:H5"	1.50	0.74
47:D5:76:LEU:HA	47:D5:83:PRO:HA	1.69	0.74
2:1E:80:ILE:HG21	2:1E:212:GLN:HA	1.70	0.74
1:13:346:G:OP1	41:B8:41:ARG:NH2	2.20	0.74
49:J8:89:GLU:OE2	49:J8:89:GLU:N	2.20	0.74
1:1G:1028(B):C:O2	1:1G:1030:C:N4	2.20	0.74
42:C8:69:CYS:HG	42:C8:79:PHE:HD2	1.34	0.74
30:29:36:ARG:HH21	30:29:89:ASP:HB3	1.53	0.73
29:19:39:LYS:HG3	29:19:40:THR:N	2.03	0.73
1:1G:1305:G:HO2'	1:1G:1306:A:H8	1.34	0.73
1:1G:558:G:OP1	62:1G:1855:HOH:O	2.05	0.73
26:1H:1597:A:N7	62:1H:3778:HOH:O	2.20	0.73
26:1H:2849:U:O2'	62:1H:3749:HOH:O	2.05	0.73
28:71:23:ASP:HB2	28:71:190:ARG:HH22	1.54	0.73
45:B5:11:PRO:HB3	45:B5:92:LEU:HD11	1.70	0.73
26:14:1022:G:O2'	26:14:1023:U:OP2	2.06	0.73
1:1G:235:C:H5'	17:8A:70:ARG:HG2	1.69	0.73
26:14:815:C:OP1	43:95:85:LYS:NZ	2.21	0.73
32:49:161:THR:HG22	32:49:163:ALA:H	1.54	0.73
1:13:76:G:N1	1:13:93:U:O2	2.19	0.73
26:14:1537:C:H2'	26:14:1538:G:C8	2.23	0.73
26:14:1729:A:H2'	26:14:1731:G:N2	2.04	0.73
26:14:2404:C:OP2	62:14:3628:HOH:O	2.07	0.73
1:1G:1435:G:H2'	1:1G:1436:U:C6	2.23	0.73
1:1G:617:G:N7	62:1G:1865:HOH:O	2.21	0.73
26:1H:568:U:O4	62:1H:3683:HOH:O	2.06	0.73
39:98:100:LEU:HD11	39:98:113:LEU:HD13	1.71	0.73
37:35:122:PRO:HB3	37:35:141:ALA:HB1	1.70	0.73
1:13:455:C:N3	1:13:477:G:N2	2.37	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2329:G:N7	62:1H:3782:HOH:O	2.21	0.73
1:13:692:U:O4	11:2I:53:SER:HB2	1.88	0.73
26:1H:617:G:OP1	31:31:40:GLN:NE2	2.22	0.73
31:39:192:LEU:HD13	31:39:194:MET:HE1	1.70	0.73
31:39:53:THR:HG23	31:39:55:GLY:H	1.52	0.73
12:3A:41:ARG:HB3	12:3A:41:ARG:HH11	1.53	0.73
32:41:67:LYS:HE2	52:M8:6:HIS:CE1	2.23	0.73
26:14:1022:G:H22	26:14:1142(A):A:H2	1.37	0.73
26:1H:1174:A:H1'	26:1H:1178:C:H41	1.54	0.73
24:3L:9:A:H2'	24:3L:11:C:H41	1.54	0.73
1:1G:377:G:OP1	16:7A:3:LYS:NZ	2.22	0.73
1:13:1348:U:H4'	9:8E:120:ARG:HD2	1.70	0.72
27:1J:86:G:N2	27:1J:90:C:O2	2.17	0.72
34:69:78:THR:HG21	34:69:104:GLN:HG3	1.70	0.72
41:B8:26:ASP:HB3	41:B8:92:GLY:H	1.54	0.72
26:14:141:A:H8	26:14:1595:G:H21	1.37	0.72
1:1G:1204:A:OP1	14:5A:3:ARG:NH1	2.23	0.72
3:22:138:VAL:HG23	3:22:151:VAL:HG23	1.70	0.72
13:4I:7:VAL:HB	32:41:115:ARG:HH22	1.54	0.72
19:AA:3:ARG:HD2	19:AA:7:LYS:HG2	1.71	0.72
47:H8:9:TYR:CE1	47:H8:35:ARG:HG2	2.24	0.72
2:12:22:LYS:HB3	2:12:40:HIS:CE1	2.23	0.72
26:14:761:A:C8	61:14:3437:SPE:H112	2.25	0.72
14:5I:6:LEU:HD13	14:5I:23:ARG:HH22	1.55	0.72
41:75:45:PHE:CE2	41:75:74:ARG:HG3	2.24	0.72
46:C5:74:PRO:HG2	46:C5:82:PRO:HG2	1.71	0.72
50:G5:47:ASN:O	50:G5:49:LYS:N	2.19	0.72
50:K8:32:LEU:HA	50:K8:35:LEU:HD23	1.72	0.72
29:11:38:LYS:HG2	29:11:40:THR:HG23	1.70	0.72
26:14:848:G:H2'	26:14:849:A:C8	2.25	0.72
26:1H:763:G:OP1	62:1H:3751:HOH:O	2.07	0.72
35:58:56:ASN:N	35:58:125:GLY:O	2.15	0.72
26:14:1678:G:H22	26:14:1989:G:H22	1.35	0.72
26:14:6:A:H62	35:15:131:GLN:H	1.35	0.72
24:3K:53:G:N2	24:3K:61:C:N3	2.36	0.72
1:13:449:C:H5	16:7I:42:ARG:HH11	1.38	0.72
1:13:501:C:H2'	1:13:502:G:H8	1.53	0.72
1:13:563:A:N6	62:13:1843:HOH:O	2.23	0.72
26:14:2655:G:N2	26:14:2665:A:OP2	2.23	0.72
22:1K:12:U:O2	22:1K:24:G:N2	2.22	0.72
3:22:84:ILE:HG23	3:22:85:ARG:HD2	1.71	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:71:35:ALA:HB2	28:71:218:MET:HG2	1.72	0.72
26:1H:2882:A:OP1	39:98:96:ARG:NH1	2.22	0.72
1:13:501:C:H2'	1:13:502:G:C8	2.24	0.72
26:14:1342:A:H2	26:14:1602:U:H3	1.36	0.72
26:14:1762:A:H2'	62:14:4247:HOH:O	1.89	0.72
1:1G:448:A:OP2	1:1G:485:G:N2	2.19	0.72
26:1H:1055:G:H1	26:1H:1104:C:H42	1.36	0.72
22:1K:6:G:N2	22:1K:67:C:O2'	2.23	0.72
29:19:30:GLU:HG3	29:19:63:ARG:NH2	2.05	0.72
1:1G:1162:C:H42	1:1G:1174:G:H1	1.35	0.72
26:1H:1364:G:N7	49:J8:2:SER:HB3	2.04	0.72
42:C8:69:CYS:SG	42:C8:79:PHE:HD2	2.13	0.72
47:D5:30:ASN:HA	47:D5:89:PHE:HE1	1.54	0.72
26:14:1899:G:H21	26:14:1902:C:H42	1.38	0.72
22:1K:17:U:HO2'	22:1K:57:G:H1	1.36	0.72
26:1H:2313:C:H4'	32:41:91:ARG:HG3	1.70	0.72
1:1G:1535:C:H41	25:4L:10:G:H21	1.38	0.72
1:13:1291:G:OP1	7:6E:37:ASN:ND2	2.23	0.72
26:14:2293:C:H5"	40:65:89:ARG:HH21	1.54	0.72
10:1A:48:THR:HA	10:1A:62:HIS:HB3	1.71	0.72
26:1H:1903:G:OP1	29:11:241:PRO:HB2	1.90	0.72
31:31:6:VAL:N	31:31:24:LEU:O	2.22	0.72
5:4E:110:LEU:HD13	5:4E:118:ILE:HD13	1.72	0.72
40:65:88:ASP:O	40:65:89:ARG:HB3	1.90	0.72
36:68:2:ILE:HD12	36:68:6:THR:HG21	1.71	0.72
56:M5:14:VAL:HG11	56:M5:58:ILE:HD11	1.70	0.72
26:14:2711:A:OP2	62:14:3508:HOH:O	2.08	0.71
1:1G:1347:G:O2'	1:1G:1373:G:O6	2.08	0.71
26:1H:2711:A:OP2	62:1H:3750:HOH:O	2.06	0.71
39:55:33:ARG:HB2	39:55:115:GLU:HB3	1.71	0.71
38:88:59:ARG:HB3	38:88:61:GLY:H	1.53	0.71
24:1L:5:C:H42	24:1L:68:G:H1	1.38	0.71
31:39:12:LEU:HD23	31:39:14:PRO:HD3	1.72	0.71
8:72:29:SER:HB3	8:72:32:LYS:HG3	1.71	0.71
54:O8:28:ARG:HE	54:O8:30:THR:HG23	1.54	0.71
1:13:659:U:H2'	1:13:660:G:C8	2.25	0.71
1:1G:1305:G:H22	1:1G:1331:G:H2'	1.55	0.71
1:1G:576:G:OP1	62:1G:1856:HOH:O	2.06	0.71
33:51:30:LYS:HD2	33:51:81:GLU:H	1.54	0.71
36:25:115:VAL:HG13	36:25:121:VAL:HG21	1.70	0.71
4:3E:85:LYS:HE2	4:3E:89:THR:HA	1.73	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
45:B5:43:VAL:HG23	45:B5:51:VAL:HG21	1.72	0.71
1:13:165:C:H2'	1:13:166:G:H8	1.56	0.71
1:13:235:C:H5'	17:8I:70:ARG:HG2	1.72	0.71
31:31:29:ASN:H	31:31:112:MET:HE1	1.56	0.71
12:3I:117:ARG:HB3	12:3I:122:THR:HB	1.70	0.71
32:49:7:LEU:HD12	32:49:104:GLU:HA	1.72	0.71
38:88:140:ALA:O	38:88:141:GLN:NE2	2.23	0.71
51:H5:44:ARG:HB2	51:H5:44:ARG:HH11	1.54	0.71
26:14:2598:A:OP1	62:14:3629:HOH:O	2.08	0.71
26:14:323:G:HO2'	26:14:1205:U:H3	1.38	0.71
4:3E:129:ASN:ND2	4:3E:144:ASP:OD1	2.24	0.71
16:7I:53:VAL:HG13	16:7I:79:VAL:HG22	1.71	0.71
47:H8:7:ALA:HB2	47:H8:59:LEU:HD22	1.73	0.71
29:19:37:LEU:HD12	29:19:37:LEU:H	1.54	0.71
30:21:101:ARG:HG2	30:21:169:ASN:OD1	1.91	0.71
33:59:89:ILE:HG21	33:59:130:ARG:HA	1.73	0.71
39:98:32:GLY:HA2	39:98:116:LEU:HD12	1.72	0.71
26:14:2331:G:H4'	48:E5:43:THR:H	1.54	0.71
52:M8:57:GLU:O	52:M8:61:ARG:NH1	2.23	0.71
1:13:664:G:H22	1:13:741:G:H1	1.38	0.71
26:14:832:G:H5'	37:35:45:LEU:HD11	1.71	0.71
1:1G:1023:G:C5	1:1G:1024:G:H1'	2.26	0.71
1:1G:987:G:H1	1:1G:1218:C:H42	1.39	0.71
1:1G:1443:G:O2'	41:75:122:ASP:OD2	2.09	0.71
26:1H:2487:G:O6	62:1H:3754:HOH:O	2.09	0.71
26:1H:248:G:OP1	62:1H:3753:HOH:O	2.09	0.71
26:1H:731:C:H5"	62:1H:3885:HOH:O	1.90	0.71
36:25:2:ILE:HD12	36:25:6:THR:HG21	1.70	0.71
37:78:65:ARG:HB3	62:Q8:404:HOH:O	1.90	0.71
1:13:598:U:H4'	8:7E:94:TYR:CD2	2.26	0.71
9:82:112:LYS:HE3	9:82:118:LYS:H	1.55	0.71
30:21:9:VAL:HG13	41:B8:3:ARG:HG3	1.73	0.71
44:E8:86:LEU:HD12	44:E8:87:PRO:HD2	1.73	0.71
1:13:1124:G:N7	1:13:1145:C:O2'	2.23	0.71
1:13:1391:U:H2'	1:13:1392:G:C8	2.26	0.71
1:13:963:G:H21	10:1I:55:LYS:NZ	1.89	0.71
1:1G:1345:U:OP2	62:1G:1857:HOH:O	2.09	0.71
37:35:71:VAL:HG13	37:35:72:PRO:HD3	1.71	0.71
34:69:98:ALA:HA	34:69:109:ILE:HD11	1.73	0.71
26:14:1154:G:OP2	42:85:58:ARG:NH1	2.23	0.70
26:14:2656:U:H3	26:14:2665:A:H2	1.38	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:1A:40:LEU:HD22	10:1A:71:LEU:HD13	1.73	0.70
1:1G:1003:G:N2	1:1G:1005:A:OP1	2.24	0.70
3:2E:136:GLN:OE1	3:2E:140:ARG:NH1	2.24	0.70
34:61:110:ASP:OD1	34:61:110:ASP:N	2.21	0.70
41:75:77:PRO:HG2	41:75:80:SER:HB2	1.73	0.70
16:7A:74:LEU:HD12	16:7A:79:VAL:HG21	1.72	0.70
26:1H:1228:G:OP2	42:C8:16:LYS:NZ	2.24	0.70
1:13:1306:A:H61	1:13:1331:G:H1'	1.55	0.70
29:19:69:ARG:HD3	29:19:105:ILE:HD11	1.73	0.70
3:22:87:LEU:HD12	3:22:88:ARG:HH21	1.55	0.70
18:9I:59:SER:HB3	18:9I:62:GLU:HB2	1.74	0.70
20:BA:69:GLY:O	20:BA:73:HIS:NE2	2.24	0.70
48:I8:14:ARG:NH1	62:I8:201:HOH:O	2.22	0.70
1:1G:1353:G:OP1	21:1B:10:ARG:NH1	2.23	0.70
26:1H:1899:G:N2	26:1H:1902:C:H5	1.88	0.70
31:31:66:PRO:O	31:31:67:GLN:HB3	1.89	0.70
4:3E:65:ARG:NH1	4:3E:70:ILE:O	2.23	0.70
34:61:92:VAL:HG13	34:61:120:ILE:HG23	1.73	0.70
8:7E:41:ARG:NH2	8:7E:123:GLU:OE1	2.22	0.70
1:13:1062:U:H2'	1:13:1063:C:C6	2.26	0.70
1:1G:1256:A:N6	1:1G:1278:U:OP2	2.24	0.70
26:1H:2135:A:N6	26:1H:2156:G:O2'	2.24	0.70
3:22:152:ILE:HB	3:22:199:LYS:HB2	1.72	0.70
4:3E:89:THR:HG23	4:3E:91:SER:H	1.57	0.70
26:14:2882:A:H5'	39:55:96:ARG:HG3	1.74	0.70
29:11:238:GLY:O	62:11:302:HOH:O	2.10	0.70
1:13:262:A:H2'	1:13:263:A:C8	2.27	0.70
26:14:1418:G:N7	62:14:3668:HOH:O	2.24	0.70
26:1H:2789:C:O2	26:1H:2894:G:N2	2.16	0.70
36:68:104:ARG:HH11	41:B8:36:GLU:HG3	1.55	0.70
47:D5:60:GLU:HA	47:D5:66:SER:HA	1.74	0.70
49:J8:91:LYS:HD3	49:J8:91:LYS:N	2.06	0.70
1:13:944:G:OP1	62:13:1838:HOH:O	2.09	0.70
10:1I:49:VAL:HG23	14:5I:41:ARG:HB2	1.72	0.70
33:51:149:ARG:NH1	33:51:167:GLU:OE2	2.25	0.70
20:BI:33:ILE:O	20:BI:37:SER:OG	2.06	0.70
46:G8:9:LYS:HA	46:G8:27:VAL:HG22	1.72	0.70
29:11:10:THR:OG1	29:11:13:ARG:HB2	1.90	0.70
27:1J:15:A:OP2	27:1J:69:G:N2	2.24	0.70
30:29:101:ARG:HG3	30:29:203:LYS:HD3	1.72	0.70
1:13:601:C:H2'	1:13:602:A:C8	2.26	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:816:C:OP2	62:14:3632:HOH:O	2.09	0.70
26:14:932:G:N7	62:14:3670:HOH:O	2.25	0.70
1:1G:1401:G:OP1	62:1G:1858:HOH:O	2.10	0.70
38:45:135:ASP:N	38:45:136:ALA:HA	2.06	0.70
1:1G:1095:U:P	1:1G:1108:G:H1	2.14	0.70
26:1H:1525:G:H2'	26:1H:1526:G:H8	1.55	0.70
26:1H:1607:C:H4'	26:1H:1608:A:O5'	1.92	0.70
8:72:12:ARG:NH2	8:72:27:PRO:HD3	2.07	0.70
27:1J:11:C:OP2	27:1J:12:C:N4	2.20	0.69
3:22:18:TRP:HE3	3:22:18:TRP:H	1.40	0.69
37:78:47:ASP:OD2	37:78:50:ARG:NH2	2.25	0.69
1:1G:1346:A:H5"	9:82:120:ARG:HH12	1.57	0.69
26:1H:1156:A:C8	42:C8:51:LYS:HG2	2.27	0.69
26:14:2306:C:H3'	26:14:2307:G:H5"	1.74	0.69
1:1G:564:C:O2'	8:72:91:ARG:NH2	2.22	0.69
26:1H:1900:A:C8	26:1H:1900:A:H5'	2.26	0.69
1:13:404:U:H5'	4:3E:122:ARG:HD2	1.73	0.69
1:13:1240:U:OP2	7:6E:116:ALA:N	2.25	0.69
15:6I:63:ARG:HG2	15:6I:67:LEU:HD12	1.75	0.69
29:11:8:PRO:HB3	29:11:14:ARG:HB2	1.73	0.69
26:14:2785:C:O2'	30:29:64:LYS:NZ	2.24	0.69
35:15:128:HIS:ND1	35:15:129:PRO:O	2.25	0.69
1:1G:1286:A:C8	1:1G:1287:A:H4'	2.26	0.69
1:1G:501:C:H2'	1:1G:502:G:H8	1.57	0.69
5:42:16:THR:OG1	5:42:17:ALA:N	2.24	0.69
7:62:113:GLU:HB2	7:62:119:ARG:HG2	1.73	0.69
5:4E:45:PHE:CE2	5:4E:47:LYS:HD2	2.28	0.69
26:1H:270(L):U:C2	34:61:50:ARG:HG2	2.27	0.69
2:12:27:LYS:O	2:12:30:ARG:NH1	2.26	0.69
1:1G:1262:C:H42	1:1G:1273:G:H1	1.40	0.69
26:1H:1534:G:N1	26:1H:1539:G:N3	2.40	0.69
26:1H:1843:C:H5'	29:11:253:GLN:OE1	1.92	0.69
26:1H:2503:A:OP2	62:1H:3755:HOH:O	2.10	0.69
26:1H:259:G:H21	26:1H:621:A:H8	1.39	0.69
33:59:8:PRO:HB2	33:59:69:ARG:HH21	1.56	0.69
26:1H:1534:G:H21	26:1H:1538:G:N2	1.91	0.69
26:1H:1932:A:OP2	62:1H:3752:HOH:O	2.09	0.69
26:1H:2217:G:O6	62:1H:3757:HOH:O	2.10	0.69
30:21:105:THR:OG1	30:21:199:ARG:NH2	2.25	0.69
3:22:150:LYS:HG3	3:22:169:ALA:HB2	1.72	0.69
26:14:1952:A:C6	36:25:22:ILE:HD11	2.27	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1226:C:O2'	13:4A:111:LYS:NZ	2.21	0.69
36:68:93:PRO:HG3	36:68:114:ILE:HG12	1.74	0.69
26:14:94:G:H21	50:G5:47:ASN:HD22	1.40	0.69
26:14:1035:U:H2'	26:14:1036:G:C8	2.27	0.69
26:14:1043:C:H42	26:14:1112:G:H1	1.38	0.69
37:78:50:ARG:HH21	37:78:50:ARG:HG3	1.57	0.69
1:1G:1347:G:N7	9:82:10:ARG:NH2	2.41	0.69
52:M8:45:GLY:O	52:M8:47:GLN:NE2	2.25	0.69
1:1G:1028(A):C:H42	1:1G:1032(B):G:H1	1.40	0.69
26:1H:2124:G:O6	26:1H:2173:A:N6	2.25	0.69
26:1H:2210:G:H4'	26:1H:2211:G:OP2	1.92	0.69
26:1H:635:C:O2'	26:1H:639:U:OP1	2.11	0.69
32:41:77:ILE:HG22	32:41:82:LEU:HD12	1.75	0.69
33:51:153:LYS:HB2	33:51:155:SER:H	1.58	0.69
7:6E:111:ARG:NH1	7:6E:113:GLU:OE2	2.26	0.69
24:3K:63:U:H6	28:71:53:ARG:HH22	1.41	0.69
1:13:468:A:H5"	16:7I:80:PHE:HB3	1.74	0.69
19:AI:11:VAL:HG11	19:AI:16:LEU:HD22	1.75	0.69
44:E8:14:PRO:HG2	44:E8:78:GLU:HB2	1.75	0.69
1:13:1455:G:OP1	20:BI:35:THR:OG1	2.04	0.69
1:13:362:G:O2'	12:3I:33:ARG:NH2	2.26	0.69
26:14:674:G:OP2	62:14:3634:HOH:O	2.10	0.69
26:14:910:A:H62	38:45:12:GLN:HA	1.57	0.69
10:1A:50:ILE:HG22	10:1A:52:GLY:H	1.58	0.69
1:1G:1218:C:OP2	14:5A:9:LYS:NZ	2.22	0.69
1:1G:1401:G:N7	62:1G:1867:HOH:O	2.24	0.69
1:13:8:A:N7	4:3E:208:SER:HB3	2.06	0.69
32:41:112:PRO:HB3	52:M8:37:SER:HB2	1.73	0.69
1:13:735:C:H2'	1:13:736:C:H6	1.58	0.69
1:1G:957:U:O2'	1:1G:959:A:N7	2.22	0.69
26:1H:1006:C:OP2	62:1H:3756:HOH:O	2.10	0.69
26:1H:1210:A:C8	26:1H:1210:A:H5'	2.28	0.69
36:25:102:VAL:HB	36:25:106:LEU:HD12	1.74	0.69
26:14:2379:G:O2'	40:65:17:ARG:NH1	2.25	0.69
44:E8:88:ARG:HB2	44:E8:92:ARG:HB3	1.73	0.69
26:1H:2210:G:H5'	26:1H:2211:G:N7	2.08	0.69
26:1H:2314:C:H2'	26:1H:2315:G:H8	1.56	0.69
30:29:134:ILE:O	30:29:134:ILE:HD12	1.93	0.69
24:3L:76:A:H8	26:14:2394:C:H42	1.38	0.68
26:14:483:A:H5'	46:C5:49:VAL:HG22	1.76	0.68
26:1H:2689:U:C6	26:1H:2689:U:H5'	2.28	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:6:A:H4'	35:58:129:PRO:HB3	1.76	0.68
62:1H:3693:HOH:O	30:21:135:HIS:NE2	2.25	0.68
8:7E:9:MET:HG3	8:7E:26:VAL:HG21	1.74	0.68
45:B5:41:ASN:HA	45:B5:44:GLU:HB2	1.75	0.68
44:E8:2:GLU:OE1	44:E8:72:LYS:NZ	2.27	0.68
26:14:1622:G:OP2	62:14:3633:HOH:O	2.09	0.68
29:19:237:GLU:OE2	62:19:402:HOH:O	2.09	0.68
1:1G:1014:A:H2'	1:1G:1015:A:C8	2.29	0.68
1:1G:998:G:H22	1:1G:1043:C:H42	1.38	0.68
1:1G:1154:G:H2'	1:1G:1155:G:H8	1.57	0.68
1:1G:278:G:OP2	17:8A:92:ARG:NH2	2.26	0.68
26:1H:2270:G:OP2	62:1H:3758:HOH:O	2.10	0.68
26:1H:982:C:OP2	62:1H:3759:HOH:O	2.11	0.68
13:4I:80:ARG:NH1	19:AI:65:ASN:O	2.26	0.68
26:14:602:G:HO2'	26:14:604:G:HO2'	1.39	0.68
1:1G:1263:C:N4	1:1G:1272:G:O6	2.20	0.68
1:1G:79:G:H1	1:1G:90:C:H42	1.40	0.68
26:1H:2118:U:O2	26:1H:2148:G:O2'	2.11	0.68
26:1H:2275:C:H5'	26:1H:2275:C:H6	1.57	0.68
41:75:24:PRO:HD3	41:75:52:ILE:HD12	1.76	0.68
1:13:963:G:H1	1:13:972:C:H42	1.38	0.68
26:14:1041:C:H42	26:14:1114:G:H1	1.40	0.68
40:A8:74:ALA:HB1	40:A8:108:GLY:HA3	1.73	0.68
1:13:323:U:H5'	20:BI:23:ARG:HB2	1.75	0.68
26:1H:1165:U:H2'	26:1H:1166:C:C6	2.29	0.68
27:1J:76:G:N7	62:1J:305:HOH:O	2.26	0.68
26:14:1359:A:H62	26:14:1372:U:H3	1.42	0.68
1:1G:1124:G:O2'	1:1G:1145:C:N4	2.27	0.68
1:1G:345:C:OP2	41:75:39:ARG:NH2	2.27	0.68
1:1G:362:G:H4'	12:3A:33:ARG:HH21	1.57	0.68
26:1H:1381:G:N7	62:1H:3795:HOH:O	2.27	0.68
26:1H:1486:A:H2'	26:1H:1487:G:H8	1.59	0.68
26:1H:2352:A:OP2	62:1H:3760:HOH:O	2.12	0.68
27:1J:44:G:H1'	27:1J:47:C:H42	1.58	0.68
11:2I:99:GLN:HB3	11:2I:105:VAL:HG11	1.76	0.68
36:25:68:GLU:OE2	36:25:78:ARG:NH1	2.27	0.68
5:42:102:ALA:HB1	5:42:106:PRO:HG2	1.76	0.68
34:69:77:LEU:HD23	34:69:78:THR:H	1.58	0.68
41:B8:12:SER:HA	41:B8:14:TYR:H	1.58	0.68
45:B5:1:MET:H2	50:G5:29:LYS:HE3	1.59	0.68
26:14:389:G:N1	37:35:71:VAL:HG12	2.09	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:4I:37:THR:O	13:4I:55:ARG:NH1	2.27	0.68
1:1G:1196:U:H5	1:1G:1397:C:H41	1.42	0.68
26:1H:2154:G:H2'	26:1H:2155:G:H8	1.59	0.68
26:1H:860:U:C5	26:1H:917:A:C2	2.81	0.68
3:2E:123:GLN:O	3:2E:128:PHE:HB2	1.94	0.68
32:41:161:THR:HG22	32:41:163:ALA:H	1.57	0.68
39:98:20:LEU:HD21	39:98:40:LYS:HD3	1.75	0.68
26:14:1430:C:H2'	26:14:1431:U:C6	2.29	0.68
26:14:1676:A:N7	62:14:3674:HOH:O	2.26	0.68
8:72:12:ARG:HH21	8:72:27:PRO:HD3	1.59	0.68
18:9I:38:GLU:OE1	18:9I:41:LYS:NZ	2.25	0.68
1:1G:1274:G:H21	1:1G:1275:A:H62	1.41	0.67
33:51:86:GLU:CD	33:51:86:GLU:H	1.97	0.67
35:58:132:ALA:O	35:58:134:ARG:NH2	2.27	0.67
42:C8:68:ALA:O	42:C8:71:GLN:HB2	1.94	0.67
26:1H:2688:U:H5	26:1H:2720:U:OP2	1.77	0.67
32:49:56:ALA:HA	32:49:59:GLU:HB3	1.76	0.67
33:51:6:ARG:HH21	33:51:7:LEU:HD11	1.59	0.67
17:8I:22:LEU:HD11	17:8I:39:SER:HB3	1.76	0.67
19:AA:10:PHE:N	19:AA:11:VAL:HB	2.09	0.67
43:D8:45:THR:O	43:D8:47:VAL:HG23	1.95	0.67
26:14:2331:G:O3'	48:E5:43:THR:HG22	1.94	0.67
1:13:353:A:H5'	1:13:353:A:H8	1.59	0.67
1:13:953:G:OP2	62:13:1839:HOH:O	2.11	0.67
26:14:2247:A:N6	62:14:3676:HOH:O	2.26	0.67
31:31:29:ASN:H	31:31:112:MET:CE	2.05	0.67
32:49:130:ASN:HB3	32:49:160:VAL:HA	1.75	0.67
3:22:6:HIS:HB2	14:5A:49:HIS:HD2	1.59	0.67
45:B5:51:VAL:HG13	45:B5:81:VAL:HG23	1.76	0.67
1:13:1446:A:OP1	1:13:1446:A:H4'	1.95	0.67
1:1G:971:G:N2	1:1G:1363:A:OP2	2.27	0.67
5:42:11:ILE:HD12	5:42:31:LEU:HD12	1.77	0.67
26:1H:2751:G:N7	33:51:3:ARG:CZ	2.57	0.67
38:88:104:PHE:HE2	38:88:125:LEU:HD11	1.57	0.67
1:13:145:G:H1	1:13:177:C:H42	1.40	0.67
26:14:1970:A:H4'	62:14:3695:HOH:O	1.92	0.67
27:16:102:G:N7	62:16:302:HOH:O	2.27	0.67
40:65:50:SER:O	40:65:76:LYS:NZ	2.23	0.67
42:C8:92:ARG:NH1	42:C8:94:ASN:OD1	2.27	0.67
1:13:1015:A:H2'	1:13:1016:A:C8	2.30	0.67
26:14:2836:U:H2'	26:14:2837:G:C8	2.29	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1452:C:H4'	1:1G:1453:G:H5'	1.75	0.67
26:1H:634:C:H2'	26:1H:635:C:C6	2.30	0.67
27:16:42:C:O2'	32:41:67:LYS:HE3	1.94	0.67
5:4E:8:GLU:OE2	5:4E:63:ARG:NH2	2.28	0.67
35:58:96:GLU:C	35:58:98:VAL:H	1.96	0.67
19:AI:36:ARG:NH1	19:AI:52:TYR:O	2.26	0.67
26:14:1364:G:OP2	49:F5:2:SER:N	2.27	0.67
46:G8:82:PRO:HB3	46:G8:99:CYS:HB2	1.77	0.67
26:1H:191:A:N1	62:1H:3798:HOH:O	2.27	0.67
26:1H:2048:G:N7	62:1H:3802:HOH:O	2.28	0.67
26:1H:2233:U:H2'	26:1H:2234:G:C8	2.29	0.67
26:1H:848:G:H2'	26:1H:849:A:C8	2.30	0.67
24:1L:36:U:H3	25:4L:19:G:H1	1.41	0.67
12:3A:37:CYS:SG	12:3A:81:SER:OG	2.47	0.67
1:13:160:A:N6	1:13:346:G:O6	2.28	0.67
26:14:2520:C:H41	26:14:2542:A:H62	1.42	0.67
26:1H:601:C:N4	62:1H:3799:HOH:O	2.28	0.67
40:A8:27:SER:HA	40:A8:88:ASP:HB2	1.77	0.67
41:B8:24:PRO:HD3	41:B8:52:ILE:HD12	1.77	0.67
53:J5:49:CYS:SG	53:J5:50:GLY:N	2.67	0.67
1:13:410:G:OP1	4:3E:30:LYS:NZ	2.26	0.67
26:14:275:G:O2'	26:14:276:A:O4'	2.10	0.67
2:1E:16:HIS:CE1	2:1E:213:LEU:HD12	2.28	0.67
26:1H:1406:U:H2'	26:1H:1407:C:C6	2.30	0.67
9:8E:10:ARG:HG3	9:8E:75:ASP:HB3	1.77	0.67
41:B8:16:ARG:NH2	41:B8:83:ILE:O	2.28	0.67
20:BA:11:SER:HA	20:BA:13:LEU:HD23	1.77	0.67
26:14:271(B):G:N7	26:14:421:U:H2'	2.09	0.67
26:14:6:A:C8	35:15:129:PRO:HB2	2.30	0.67
10:1A:24:VAL:HG21	10:1A:37:PRO:HD3	1.76	0.67
1:1G:1023:G:H5"	1:1G:1024:G:H21	1.59	0.67
1:1G:1192:C:OP2	3:22:4:LYS:NZ	2.28	0.67
14:5A:27:CYS:O	14:5A:29:ARG:NH2	2.25	0.67
42:85:28:ARG:NH1	42:85:38:THR:OG1	2.26	0.67
7:6E:16:LEU:HG	9:8E:42:ARG:HA	1.77	0.67
46:C5:42:VAL:HG13	46:C5:65:ALA:HB3	1.76	0.67
54:O8:19:ARG:HG3	54:O8:21:TYR:HE1	1.60	0.67
26:14:389:G:H1	37:35:71:VAL:HG12	1.59	0.66
29:19:93:ALA:HB3	29:19:105:ILE:HG22	1.77	0.66
26:1H:1380:G:N7	62:1H:3767:HOH:O	2.28	0.66
13:4A:66:LEU:HA	13:4A:70:LEU:HB2	1.77	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:D5:111:VAL:HG12	47:D5:145:GLU:HB2	1.76	0.66
1:13:200:G:N2	1:13:218:C:N3	2.43	0.66
1:1G:1235:U:O2'	1:1G:1305:G:O5'	2.12	0.66
26:1H:2287:A:H62	26:1H:2344:U:H3	1.42	0.66
12:3A:47:LYS:HD2	12:3A:48:PRO:HD2	1.75	0.66
32:49:60:LEU:HD22	32:49:68:PRO:HB3	1.77	0.66
42:85:92:ARG:HG3	42:85:94:ASN:HB3	1.76	0.66
26:14:95:G:H4'	50:G5:46:GLN:HB2	1.76	0.66
35:15:56:ASN:H	35:15:125:GLY:HA3	1.60	0.66
26:1H:862:G:OP2	62:1H:3764:HOH:O	2.13	0.66
10:1I:78:ASN:O	10:1I:81:THR:OG1	2.14	0.66
37:35:39:LYS:HG3	37:35:45:LEU:HD22	1.77	0.66
13:4A:22:ILE:HB	13:4A:25:ILE:HG13	1.77	0.66
14:5A:22:THR:HB	14:5A:33:VAL:HG21	1.77	0.66
14:5I:27:CYS:SG	14:5I:29:ARG:HB2	2.36	0.66
43:D8:37:VAL:HG23	43:D8:51:VAL:HG21	1.78	0.66
1:13:411:A:C4	1:13:413:G:H1'	2.29	0.66
26:14:1952:A:C5	36:25:22:ILE:HD11	2.31	0.66
26:14:2400:G:H2'	26:14:2401:U:C6	2.31	0.66
26:1H:1899:G:H1	26:1H:1902:C:H41	1.44	0.66
3:2E:72:LYS:HD3	3:2E:75:VAL:HG21	1.77	0.66
37:35:93:GLY:H	37:35:123:LEU:HD22	1.59	0.66
17:8A:66:SER:O	17:8A:70:ARG:NH1	2.28	0.66
26:1H:1491:G:O4'	29:11:99:ASP:HB3	1.96	0.66
26:14:1997:G:OP2	62:14:3635:HOH:O	2.13	0.66
1:1G:353:A:H8	1:1G:353:A:H5'	1.60	0.66
1:1G:973:G:O3'	14:5A:41:ARG:NH2	2.28	0.66
26:1H:945:A:N3	62:1H:3796:HOH:O	2.27	0.66
1:13:8:A:N6	4:3E:205:GLU:O	2.27	0.66
12:3I:42:THR:HG22	12:3I:54:LYS:HD2	1.77	0.66
47:D5:5:LEU:HD11	47:D5:44:PHE:HA	1.77	0.66
47:H8:128:VAL:HB	47:H8:161:VAL:HG12	1.76	0.66
26:1H:1292:U:H2'	26:1H:1293:C:C6	2.31	0.66
30:21:64:LYS:HA	30:21:67:PHE:O	1.95	0.66
24:3K:15:G:O6	24:3K:48:C:N4	2.29	0.66
26:1H:637:A:H2'	37:78:117:GLU:OE1	1.96	0.66
26:1H:969:U:OP1	51:L8:17:LYS:HG2	1.96	0.66
26:14:1044:G:O2'	26:14:1047:G:O2'	2.14	0.66
26:14:1727:U:H3	26:14:1733:G:H1	1.44	0.66
26:14:247:G:H4'	26:14:386:G:C5	2.31	0.66
26:14:71:A:OP2	26:14:71:A:H3'	1.96	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1300:G:O2'	1:1G:1301:U:O5'	2.13	0.66
1:1G:1352:C:N4	1:1G:1370:G:H1	1.93	0.66
1:1G:300:A:N1	62:1G:1872:HOH:O	2.29	0.66
26:1H:1412:A:H2'	26:1H:1413:G:C8	2.30	0.66
23:2L:24:C:H2'	23:2L:25:U:C6	2.30	0.66
27:1J:7:G:H4'	40:65:29:PHE:CD2	2.31	0.66
2:12:71:VAL:HB	2:12:165:VAL:HG22	1.77	0.66
26:14:2162:G:O2'	26:14:2173:A:OP1	2.13	0.66
2:1E:166:ASP:HB3	2:1E:169:LYS:HB2	1.77	0.66
4:32:7:PRO:HB2	4:32:10:ARG:HD2	1.78	0.66
37:35:55:ARG:HG2	37:35:56:SER:N	2.11	0.66
41:75:10:VAL:O	41:75:12:SER:N	2.29	0.66
29:11:35:LYS:HD3	29:11:36:PRO:HD2	1.78	0.66
1:13:1346:A:H5"	9:8E:120:ARG:NH1	2.11	0.66
2:1E:178:ARG:NH1	2:1E:196:LEU:O	2.27	0.66
38:88:66:ILE:O	38:88:104:PHE:N	2.28	0.66
26:1H:581:C:OP1	42:C8:33:ARG:HG3	1.96	0.66
47:D5:10:ARG:HH21	47:D5:26:GLY:H	1.40	0.66
2:12:50:GLU:HB2	2:12:199:TYR:HB3	1.77	0.66
1:1G:659:U:OP1	15:6A:9:GLN:NE2	2.29	0.66
1:1G:838:G:N2	1:1G:848:C:N3	2.43	0.66
34:61:110:ASP:HB2	34:61:112:LYS:HG2	1.78	0.66
43:95:1:MET:HA	43:95:42:GLY:H	1.61	0.66
40:A8:59:LYS:HG2	40:A8:60:GLY:H	1.60	0.66
41:B8:26:ASP:HB2	41:B8:91:ARG:HA	1.77	0.66
41:B8:26:ASP:O	41:B8:49:VAL:HG13	1.96	0.66
1:13:1034:G:N2	1:13:1035:A:N7	2.44	0.65
26:14:993:G:OP1	42:85:50:ARG:NH2	2.28	0.65
26:1H:65:C:H2'	26:1H:66:C:H6	1.61	0.65
37:35:98:GLU:HA	37:35:101:VAL:HG12	1.77	0.65
1:13:345:C:N4	36:68:116:SER:O	2.24	0.65
26:1H:71:A:H2	45:F8:31:HIS:HE2	1.44	0.65
56:Q8:33:ASN:HA	56:Q8:36:LYS:HD2	1.78	0.65
1:1G:1274:G:N2	1:1G:1275:A:H62	1.94	0.65
1:1G:261:U:OP2	20:BA:79:ARG:NH2	2.29	0.65
1:1G:474:G:H2'	1:1G:475:G:H8	1.61	0.65
1:1G:548:G:OP1	62:1G:1859:HOH:O	2.12	0.65
30:21:82:ARG:O	30:21:84:PHE:N	2.29	0.65
5:42:31:LEU:HD22	5:42:45:PHE:HB2	1.78	0.65
26:14:1653:G:H3'	39:55:2:ARG:HG2	1.77	0.65
38:88:110:THR:HG23	38:88:113:GLN:OE1	1.96	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:8A:87:LYS:O	17:8A:91:ARG:HG3	1.95	0.65
44:A5:65:LEU:HD13	44:A5:68:ARG:HD3	1.77	0.65
26:14:1015:G:N2	26:14:1147:C:O2	2.26	0.65
26:1H:2324:C:H5"	26:1H:2325:G:H5'	1.79	0.65
26:1H:2409:G:N7	62:1H:3809:HOH:O	2.29	0.65
26:1H:981:A:OP1	62:1H:3765:HOH:O	2.14	0.65
38:45:75:THR:OG1	38:45:87:LYS:NZ	2.30	0.65
1:13:1202:G:N2	14:5I:46:GLU:OE1	2.24	0.65
34:69:104:GLN:OE1	34:69:105:HIS:ND1	2.30	0.65
26:14:1044:G:HO2'	26:14:1047:G:HO2'	1.40	0.65
26:14:945:A:N3	62:14:3683:HOH:O	2.28	0.65
1:1G:1028:C:H42	1:1G:1033:G:H1	1.44	0.65
26:1H:548:A:H2'	26:1H:549:G:H5'	1.79	0.65
4:3E:154:ASN:OD1	4:3E:154:ASN:N	2.30	0.65
4:3E:30:LYS:HA	4:3E:35:ARG:HE	1.62	0.65
1:13:474:G:H5"	16:7I:81:ARG:HE	1.60	0.65
26:1H:881:G:H1	26:1H:895:U:H3	1.44	0.65
42:C8:92:ARG:HD3	42:C8:94:ASN:HB3	1.76	0.65
47:H8:77:ASP:OD1	47:H8:80:ARG:HD2	1.96	0.65
1:13:619:U:H3	4:3E:134:ASP:HB2	1.62	0.65
26:14:1495:A:N7	62:14:3690:HOH:O	2.30	0.65
26:14:2745:C:O2	33:59:139:GLN:NE2	2.30	0.65
26:14:1113:U:OP1	26:14:2751:G:N2	2.30	0.65
10:1A:55:LYS:NZ	10:1A:57:LYS:HG2	2.12	0.65
30:29:54:GLN:HB2	30:29:72:VAL:HB	1.78	0.65
26:1H:910:A:C5	38:88:13:GLN:HG3	2.32	0.65
47:H8:165:VAL:HB	47:H8:166:SER:HA	1.78	0.65
1:13:719:C:O2'	18:9I:49:LYS:HB3	1.97	0.65
26:14:2148:G:H2'	26:14:2149:G:H8	1.60	0.65
1:1G:1316:G:H4'	14:5A:18:VAL:HG11	1.79	0.65
38:45:27:VAL:HB	38:45:28:ALA:CA	2.10	0.65
5:4E:8:GLU:HG2	5:4E:34:VAL:HG22	1.77	0.65
14:5I:23:ARG:HH11	14:5I:30:ALA:HB2	1.61	0.65
40:65:29:PHE:HD1	40:65:30:ARG:N	1.93	0.65
2:12:71:VAL:HG11	2:12:164:VAL:HA	1.77	0.65
1:13:611:A:H61	1:13:629:G:H1	1.44	0.65
1:1G:862:C:H1'	1:1G:874:G:H5"	1.78	0.65
26:1H:860:U:C5	26:1H:917:A:H2	2.15	0.65
31:39:79:GLY:HA2	31:39:86:GLY:HA2	1.77	0.65
38:45:34:LEU:HB2	38:45:118:LEU:HD13	1.78	0.65
2:12:118:LEU:HB3	2:12:142:LEU:HD12	1.78	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:620:G:H4'	26:14:621:A:H5"	1.79	0.65
1:1G:976:G:P	14:5A:32:SER:H	2.20	0.65
26:1H:573:G:O2'	26:1H:574:C:H3'	1.97	0.65
36:25:49:ARG:HA	36:25:53:LYS:HZ2	1.60	0.65
41:75:18:ASP:N	41:75:18:ASP:OD1	2.30	0.65
38:88:66:ILE:HG13	38:88:67:ARG:H	1.61	0.65
9:8E:17:VAL:HG11	9:8E:81:ILE:HD13	1.79	0.65
9:8E:13:ALA:HB2	9:8E:68:GLY:HA3	1.77	0.65
42:C8:90:VAL:HG22	43:D8:39:LEU:HB3	1.79	0.65
49:J8:84:GLY:HA2	49:J8:85:LEU:HB3	1.78	0.65
10:1A:32:ALA:HA	10:1A:76:ASN:HD21	1.62	0.65
1:1G:620:C:H2'	1:1G:621:A:O4'	1.97	0.65
1:1G:587:G:N2	1:1G:754:C:OP2	2.29	0.65
26:1H:1049:C:H2'	26:1H:1050:A:H5'	1.78	0.65
26:1H:832:G:H5'	37:78:45:LEU:HD11	1.78	0.65
23:2L:24:C:H2'	23:2L:25:U:H6	1.61	0.65
39:55:38:VAL:HG22	39:55:112:ALA:HB2	1.79	0.65
27:1J:9:G:P	40:65:25:ARG:HH22	2.20	0.65
37:78:18:ARG:HG3	37:78:18:ARG:HH21	1.60	0.65
10:1A:75:ILE:HG13	10:1A:76:ASN:N	2.11	0.64
1:1G:664:G:H22	1:1G:741:G:H1	1.45	0.64
26:1H:1430:C:H2'	26:1H:1431:U:C6	2.32	0.64
26:1H:1453:A:OP2	62:1H:3766:HOH:O	2.14	0.64
26:1H:2849:U:H4'	26:1H:2868:A:C2	2.33	0.64
42:85:91:ASP:O	42:85:92:ARG:HG2	1.97	0.64
40:A8:38:GLN:HG2	40:A8:47:THR:HG21	1.79	0.64
47:D5:10:ARG:NH2	47:D5:26:GLY:O	2.30	0.64
26:14:1169:G:N2	26:14:1180:C:N3	2.40	0.64
26:14:1329:U:H5"	26:14:1330:C:H5	1.62	0.64
26:14:2210:G:H3'	26:14:2211:G:C8	2.33	0.64
26:14:498:G:H21	46:C5:47:LYS:NZ	1.93	0.64
26:14:6:A:H62	35:15:131:GLN:N	1.95	0.64
26:1H:2105:C:H2'	26:1H:2106:G:H8	1.62	0.64
8:7E:121:ASP:HB2	8:7E:125:ARG:NH2	2.12	0.64
43:95:21:ARG:NH1	43:95:91:TYR:OH	2.29	0.64
41:B8:4:GLY:HA2	41:B8:7:ILE:HG12	1.78	0.64
1:13:1296:C:OP1	13:4I:44:ARG:NH2	2.29	0.64
1:13:67:C:H2'	1:13:68:G:C8	2.32	0.64
26:14:30:G:H2'	26:14:31:C:C6	2.32	0.64
26:14:900:A:H3'	26:14:901:A:H8	1.60	0.64
1:1G:1305:G:H22	1:1G:1331:G:C2'	2.09	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1386:C:H2'	26:1H:1387:C:H6	1.61	0.64
26:14:1670:C:O2	30:29:129:HIS:NE2	2.27	0.64
11:2A:48:ILE:HD11	11:2A:64:ALA:HA	1.79	0.64
37:35:63:PRO:HD3	56:M5:27:THR:HG22	1.79	0.64
25:4K:24:A:H2'	25:4K:25:A:C8	2.32	0.64
1:1G:1319:A:OP2	19:AA:3:ARG:HG3	1.97	0.64
47:D5:30:ASN:HA	47:D5:89:PHE:CE1	2.32	0.64
52:M8:16:CYS:HB3	52:M8:36:CYS:HB3	1.79	0.64
1:13:737:A:H2'	1:13:738:C:C6	2.32	0.64
26:14:2147:G:H2'	26:14:2148:G:H4'	1.78	0.64
35:15:42:TRP:O	42:85:64:ARG:NH2	2.30	0.64
1:1G:972:C:O2'	10:1A:55:LYS:HG3	1.97	0.64
2:1E:53:ARG:NH2	2:1E:198:ASP:O	2.30	0.64
1:1G:1268:A:H2'	1:1G:1269:A:C8	2.33	0.64
1:1G:1401:G:OP1	25:4L:18:G:O2'	2.13	0.64
30:29:116:VAL:HG11	30:29:138:PRO:HB3	1.78	0.64
34:61:123:LEU:HD23	34:61:143:SER:HA	1.77	0.64
1:1G:1147:C:O2	9:82:16:ARG:NH1	2.30	0.64
9:8E:26:VAL:HG13	9:8E:61:ALA:HB3	1.78	0.64
40:A8:28:VAL:HG11	40:A8:98:VAL:HG13	1.80	0.64
36:68:104:ARG:HD3	41:B8:36:GLU:HG3	1.79	0.64
1:13:1286:A:C8	1:13:1287:A:H4'	2.33	0.64
26:14:1040:C:H2'	26:14:1041:C:C6	2.33	0.64
27:16:66:A:H61	27:16:107:U:H2'	1.62	0.64
26:1H:176:G:O2'	26:1H:177:G:H5'	1.98	0.64
14:5A:41:ARG:HG3	14:5A:42:ILE:HG13	1.80	0.64
17:8I:9:VAL:HG21	17:8I:84:LEU:HD12	1.79	0.64
40:A8:106:ARG:NH1	40:A8:107:GLU:HG2	2.13	0.64
48:E5:12:ASN:HA	48:E5:14:ARG:HH21	1.61	0.64
53:N8:36:CYS:HB2	53:N8:49:CYS:SG	2.37	0.64
33:51:157:TYR:H	33:51:171:LEU:HA	1.61	0.64
37:78:18:ARG:O	37:78:19:VAL:HB	1.97	0.64
9:8E:50:LEU:HA	9:8E:53:VAL:HG22	1.80	0.64
47:H8:132:ASN:OD1	47:H8:132:ASN:N	2.31	0.64
26:14:1169:G:H1	26:14:1180:C:H42	1.44	0.64
26:14:990:A:H8	26:14:990:A:H5'	1.63	0.64
26:1H:1521:G:N7	62:1H:3812:HOH:O	2.30	0.64
30:21:116:VAL:O	30:21:117:MET:HB3	1.97	0.64
26:14:1428:C:N4	26:14:1570:A:OP2	2.27	0.64
26:14:2712:U:O2'	62:14:3636:HOH:O	2.14	0.64
26:1H:1364:G:OP2	49:J8:2:SER:OG	2.14	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:1K:17:U:O2'	22:1K:57:G:N1	2.30	0.64
33:59:70:THR:O	33:59:74:ASN:ND2	2.22	0.64
33:59:7:LEU:HD12	33:59:8:PRO:HD3	1.80	0.64
35:15:13:TRP:O	35:15:135:PRO:HD2	1.97	0.64
29:19:108:PRO:HB3	29:19:143:HIS:CE1	2.33	0.64
1:1G:1134:G:C2	1:1G:1135:U:H1'	2.33	0.64
1:1G:1127:G:N2	1:1G:1145:C:N3	2.46	0.64
1:1G:8:A:N6	4:32:209:ARG:HB2	2.11	0.64
26:1H:1021:A:C8	26:1H:1022:G:H5"	2.32	0.64
26:1H:1113:U:OP1	33:51:2:SER:N	2.30	0.64
26:1H:2130:U:OP2	28:71:6:ARG:NH1	2.23	0.64
26:1H:2723:C:H5"	39:98:1:MET:HE2	1.80	0.64
4:3E:107:ARG:NH2	4:3E:194:LEU:HD22	2.11	0.64
1:13:974:A:OP2	14:5I:29:ARG:NH1	2.30	0.64
15:6A:16:ALA:HB1	15:6A:21:ASP:HB3	1.78	0.64
8:72:120:THR:HG23	8:72:123:GLU:H	1.62	0.64
41:75:7:ILE:HG13	41:75:8:LYS:H	1.63	0.64
38:88:35:VAL:HG13	38:88:130:LYS:HB3	1.78	0.64
48:E5:27:GLU:HB2	48:E5:69:PHE:HD1	1.63	0.64
26:14:2773:C:OP1	30:29:166:THR:OG1	2.15	0.64
26:14:34:C:H1'	26:14:35:G:OP1	1.99	0.64
1:1G:448:A:P	1:1G:485:G:H22	2.21	0.64
30:29:97:LYS:N	30:29:100:GLU:OE1	2.28	0.64
1:1G:36:C:OP1	12:3A:123:LYS:NZ	2.31	0.64
32:49:11:TYR:OH	32:49:16:ARG:NH2	2.30	0.64
34:69:45:LYS:HA	34:69:48:GLU:HB3	1.80	0.64
42:85:66:ASN:O	42:85:70:ARG:HB2	1.97	0.64
19:AI:41:VAL:HG12	19:AI:44:MET:HB2	1.80	0.64
1:13:1223:C:P	19:AI:78:ARG:HH12	2.21	0.64
41:B8:42:ILE:HD12	41:B8:42:ILE:H	1.63	0.64
1:13:1368:G:H5"	9:8E:112:LYS:HB3	1.80	0.63
26:14:1614:A:H2	62:14:4025:HOH:O	1.81	0.63
1:1G:1203:C:H2'	1:1G:1204:A:C8	2.33	0.63
1:1G:1251:A:H2'	1:1G:1252:A:C8	2.33	0.63
26:1H:620:G:H4'	26:1H:621:A:C5'	2.28	0.63
26:1H:631:A:H5'	62:1H:4891:HOH:O	1.98	0.63
26:1H:900:A:H3'	26:1H:901:A:H8	1.61	0.63
27:1J:15:A:H1'	27:1J:109:G:C8	2.33	0.63
13:4I:107:ALA:HB3	13:4I:111:LYS:HD3	1.79	0.63
26:14:1771:C:HO2'	26:14:1786:A:H8	1.46	0.63
26:14:2156:G:N7	26:14:2157:G:N2	2.46	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2296:U:OP2	40:65:9:ARG:NH1	2.26	0.63
2:1E:237:ALA:O	2:1E:239:VAL:N	2.31	0.63
1:1G:1104:G:O2'	2:12:111:ARG:NH2	2.31	0.63
1:1G:560:U:O2'	1:1G:561:U:OP2	2.15	0.63
1:1G:973:G:OP1	10:1A:57:LYS:NZ	2.32	0.63
35:58:57:ALA:C	35:58:59:LYS:H	2.00	0.63
1:13:453:A:H4'	16:7I:72:ARG:HB2	1.78	0.63
9:8E:5:TYR:HE1	9:8E:16:ARG:HG2	1.62	0.63
46:C5:86:ARG:HG3	46:C5:87:LYS:N	2.11	0.63
26:14:451:C:OP2	62:14:3637:HOH:O	2.15	0.63
4:3E:167:GLY:HA2	29:19:135:PHE:CE1	2.31	0.63
26:1H:320:A:H2'	31:31:136:THR:HG21	1.80	0.63
26:1H:33:U:H4'	26:1H:34:C:OP1	1.97	0.63
31:39:178:PRO:HB3	31:39:198:ALA:HA	1.80	0.63
14:5I:3:ARG:HB2	14:5I:3:ARG:NH1	2.12	0.63
26:1H:2336:A:H61	48:I8:43:THR:HB	1.64	0.63
26:14:2432:A:H2'	26:14:2433:A:C8	2.34	0.63
26:14:588:U:H2'	26:14:589:C:C6	2.33	0.63
1:1G:1181:G:N2	1:1G:1182:G:O2'	2.32	0.63
26:1H:273(F):C:H3'	26:1H:274:G:H5"	1.80	0.63
3:2E:8:ILE:HG23	3:2E:16:ARG:HG2	1.81	0.63
4:3E:98:GLU:HG2	4:3E:189:PRO:HG2	1.81	0.63
5:4E:33:VAL:HG11	5:4E:109:ILE:HA	1.80	0.63
5:4E:10:MET:HB3	5:4E:32:VAL:HG22	1.80	0.63
19:AI:41:VAL:HA	19:AI:44:MET:HG3	1.81	0.63
26:14:370:G:N7	62:14:3691:HOH:O	2.31	0.63
30:29:25:VAL:HG12	30:29:26:ILE:H	1.63	0.63
28:71:185:LEU:O	28:71:189:ILE:N	2.31	0.63
39:98:55:ALA:HA	39:98:80:PHE:CE1	2.33	0.63
47:D5:23:LYS:HD3	47:D5:40:ASP:HA	1.81	0.63
26:14:1843:C:H5'	29:19:253:GLN:OE1	1.97	0.63
26:14:2520:C:H41	26:14:2542:A:N6	1.97	0.63
26:14:273(C):C:H42	26:14:363(C):G:H1	1.45	0.63
10:1A:44:VAL:HG22	10:1A:66:ARG:HB3	1.80	0.63
1:1G:1127:G:O2'	1:1G:1148:U:O2	2.17	0.63
1:1G:785:G:N7	62:1G:1873:HOH:O	2.30	0.63
26:1H:974(A):C:OP1	62:1H:3768:HOH:O	2.15	0.63
27:1J:103:U:HO2'	47:D5:29:TYR:HH	1.44	0.63
26:14:993:G:N3	43:95:89:GLN:NE2	2.46	0.63
41:B8:84:GLN:HG2	41:B8:85:LYS:HD2	1.81	0.63
26:14:2839:G:H5'	39:55:46:GLY:HA2	1.81	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1E:17:PHE:HD1	2:1E:44:LEU:HD11	1.64	0.63
1:1G:1127:G:H2'	1:1G:1128:C:H6	1.64	0.63
26:1H:2068:U:H3	26:1H:2430:A:H2	1.44	0.63
26:1H:2347:C:OP1	54:O8:39:TYR:OH	2.10	0.63
12:3I:57:LYS:HD3	12:3I:67:THR:HG22	1.81	0.63
33:59:159:GLU:O	33:59:163:TYR:OH	2.15	0.63
26:1H:2124:G:H4'	28:71:174:PRO:HG3	1.80	0.63
1:1G:1349:A:OP2	9:82:118:LYS:NZ	2.32	0.63
39:98:55:ALA:HB2	39:98:79:LEU:HD13	1.80	0.63
26:1H:61:G:OP1	50:K8:51:ARG:NH1	2.32	0.63
1:13:1023:G:H3'	1:13:1024:G:H5"	1.81	0.63
26:1H:1329:U:H5"	26:1H:1330:C:H5	1.63	0.63
11:2I:121:PRO:HG2	11:2I:126:ARG:HG2	1.79	0.63
34:61:10:GLU:OE1	34:61:11:ASN:ND2	2.30	0.63
41:75:54:ARG:HG3	41:75:59:THR:HG21	1.79	0.63
26:1H:534:U:H5'	42:C8:42:ALA:HB1	1.81	0.63
56:Q8:49:VAL:HG12	56:Q8:49:VAL:O	1.98	0.63
1:13:280:C:C2	17:8I:38:ARG:HG3	2.34	0.63
1:1G:1028(A):C:N3	1:1G:1032(B):G:N2	2.41	0.63
1:1G:1521:G:N3	62:1G:1874:HOH:O	2.31	0.63
10:1I:26:ALA:O	10:1I:30:SER:OG	2.09	0.63
13:4I:37:THR:HB	13:4I:55:ARG:HD2	1.79	0.63
20:BI:75:ASN:N	20:BI:75:ASN:OD1	2.27	0.63
50:K8:42:GLY:C	50:K8:44:LEU:H	2.02	0.63
26:1H:2032:G:H21	30:21:146:THR:CG2	2.10	0.62
26:1H:2287:A:C2	26:1H:2346:A:H2	2.17	0.62
22:1K:7:U:H3	22:1K:66:A:H61	1.47	0.62
26:14:587:C:O2	37:35:33:ARG:NH1	2.31	0.62
31:39:157:VAL:HG12	31:39:198:ALA:HB1	1.81	0.62
1:13:590:C:O3'	8:7E:30:ARG:NH1	2.30	0.62
9:82:46:ALA:HB2	9:82:74:ILE:HG23	1.80	0.62
19:AI:42:PRO:HD3	52:M8:63:TYR:OH	1.98	0.62
42:C8:92:ARG:HD2	43:D8:11:GLN:HB2	1.80	0.62
52:M8:37:SER:HB3	52:M8:42:PHE:CZ	2.33	0.62
29:11:2:ALA:HA	29:11:20:ASP:CB	2.29	0.62
1:13:1454:G:OP1	20:BI:39:LYS:NZ	2.29	0.62
26:1H:863:A:H2'	26:1H:864:G:H8	1.63	0.62
1:1G:777:A:H2	11:2A:119:CYS:HB3	1.64	0.62
11:2A:14:VAL:HG11	11:2A:35:PRO:HD3	1.81	0.62
3:2E:15:THR:HG21	3:2E:181:ASN:HA	1.80	0.62
1:1G:363:A:OP1	12:3A:33:ARG:HG3	1.99	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
33:59:30:LYS:HB3	33:59:79:VAL:O	1.99	0.62
39:98:104:ARG:NH1	39:98:107:ASP:OD2	2.30	0.62
18:9A:22:VAL:HG22	18:9A:23:LYS:H	1.64	0.62
26:14:1198:U:H2'	26:14:1199:U:C6	2.34	0.62
26:14:142:G:H5"	26:14:1598:C:O2'	1.99	0.62
26:14:1794:U:H2'	26:14:1795:C:H6	1.64	0.62
26:14:774:A:H2	26:14:787:U:HO2'	1.46	0.62
1:1G:1534:A:N6	25:4L:11:U:O4	2.33	0.62
1:1G:770:C:OP1	62:1G:1860:HOH:O	2.16	0.62
26:1H:1345:C:OP2	62:1H:3769:HOH:O	2.16	0.62
6:5E:94:GLN:HE21	18:9I:32:ARG:HG2	1.64	0.62
1:1G:584:G:H5'	17:8A:91:ARG:HH12	1.64	0.62
26:1H:1178:C:H4'	26:1H:1179:C:OP1	1.98	0.62
26:1H:1786:A:H2	26:1H:2606:C:H1'	1.64	0.62
5:4E:35:GLY:H	5:4E:112:LEU:HD13	1.64	0.62
45:F8:11:PRO:HG2	45:F8:13:LEU:HD21	1.80	0.62
26:1H:76:C:O2'	50:K8:62:THR:HG21	2.00	0.62
1:13:973:G:H3'	1:13:974:A:H5"	1.82	0.62
26:14:193:U:OP2	62:14:3639:HOH:O	2.16	0.62
26:14:71:A:C8	26:14:71:A:H5'	2.35	0.62
29:19:145:VAL:HG13	29:19:191:ALA:HB2	1.79	0.62
2:1E:84:GLU:HB3	2:1E:219:VAL:HG21	1.80	0.62
1:1G:1513:A:H2'	1:1G:1514:C:C6	2.35	0.62
1:1G:181:G:O2'	1:1G:183:G:O6	2.18	0.62
1:1G:589:C:H42	1:1G:650:G:H1	1.48	0.62
26:1H:226:G:H21	26:1H:228:A:H2	1.46	0.62
13:4A:14:ARG:HA	13:4A:43:THR:O	1.99	0.62
1:13:1149:C:H2'	1:13:1150:U:C6	2.34	0.62
1:13:1301:U:O2'	1:13:1302:U:H3'	2.00	0.62
1:13:624:C:O3'	16:7I:10:GLY:HA2	1.99	0.62
26:1H:1525:G:H2'	26:1H:1526:G:C8	2.35	0.62
26:1H:2400:G:H2'	26:1H:2401:U:H6	1.64	0.62
26:1H:2784:C:O2'	30:21:37:ARG:NH1	2.32	0.62
40:65:89:ARG:HG3	40:65:92:TYR:O	1.99	0.62
26:1H:2178:C:O2'	28:71:168:THR:OG1	2.12	0.62
41:75:55:ASN:N	41:75:59:THR:HG22	2.15	0.62
17:8I:45:HIS:HB2	17:8I:65:ILE:HD13	1.81	0.62
1:13:714:G:H2'	1:13:715:A:C8	2.33	0.62
1:1G:673:G:H2'	1:1G:674:G:C8	2.34	0.62
1:13:256:U:H2'	1:13:257:G:C8	2.34	0.62
26:14:270:A:OP2	26:14:270(Y):G:N1	2.32	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:305:U:H2'	26:14:306:U:C6	2.35	0.62
26:1H:639:U:H2'	26:1H:640:C:C6	2.35	0.62
11:2A:100:ALA:O	11:2A:102:GLY:N	2.33	0.62
12:3A:59:ARG:HA	12:3A:65:GLU:H	1.64	0.62
3:22:6:HIS:HB2	14:5A:49:HIS:CD2	2.35	0.62
40:65:67:ARG:CZ	40:65:67:ARG:HB2	2.30	0.62
26:14:1266:G:O5'	44:A5:15:ARG:NH2	2.33	0.62
41:B8:54:ARG:HA	41:B8:59:THR:HB	1.82	0.62
2:12:40:HIS:CD2	2:12:190:THR:HG21	2.35	0.62
1:13:1504:G:OP1	1:13:1507:A:H4'	2.00	0.62
26:14:1048:A:N6	26:14:1111:A:O2'	2.33	0.62
26:14:1771:C:O2'	26:14:1786:A:H8	1.81	0.62
26:14:636:G:N7	37:35:113:LYS:NZ	2.40	0.62
1:1G:114:U:H2'	1:1G:115:G:C8	2.35	0.62
1:1G:1157:A:H61	1:1G:1177:G:H1	1.47	0.62
26:1H:2470:G:H5'	38:88:56:ARG:HH12	1.64	0.62
26:1H:270(E):G:H1	26:1H:270(U):C:H42	1.48	0.62
26:1H:336:C:OP1	46:G8:83:THR:HG23	2.00	0.62
30:29:111:ARG:HD2	30:29:160:TYR:CE2	2.34	0.62
24:3L:50:C:H2'	24:3L:51:A:H8	1.65	0.62
24:3L:72:C:H3'	24:3L:73:A:H5"	1.82	0.62
19:AI:51:VAL:O	19:AI:57:HIS:HA	2.00	0.62
51:H5:13:ILE:O	62:H5:101:HOH:O	2.16	0.62
26:1H:1796:U:H2'	26:1H:1797:C:C6	2.34	0.62
4:3E:64:LEU:HD22	4:3E:198:VAL:HG11	1.82	0.62
53:N8:40:LYS:HG2	53:N8:46:CYS:HA	1.82	0.62
26:14:581:C:H2'	26:14:582:G:H8	1.65	0.61
1:1G:842:C:O2'	1:1G:848:C:N3	2.33	0.61
26:1H:2789:C:H1'	26:1H:2892:A:H2	1.65	0.61
23:2K:54:G:H2'	23:2K:55:5MU:H6	1.65	0.61
34:61:69:LYS:HG3	34:61:136:VAL:HB	1.82	0.61
9:82:27:THR:OG1	9:82:31:GLN:O	2.11	0.61
9:8E:112:LYS:HA	9:8E:119:ALA:HB2	1.81	0.61
20:BI:57:ARG:HH11	20:BI:102:GLY:HA2	1.64	0.61
26:14:987:G:O2'	26:14:1000:A:N3	2.29	0.61
26:14:1012:U:OP1	42:85:75:ASN:ND2	2.33	0.61
26:14:2537:U:H2'	26:14:2538:C:C6	2.34	0.61
10:1A:11:PHE:HE1	10:1A:67:THR:HG22	1.65	0.61
36:25:10:VAL:HG12	36:25:19:ILE:HG12	1.82	0.61
12:3A:41:ARG:HD2	12:3A:42:THR:H	1.65	0.61
15:6I:82:ILE:O	15:6I:86:GLY:N	2.33	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:71:212:VAL:HG21	28:71:226:PRO:HG3	1.81	0.61
9:82:43:ALA:HA	9:82:74:ILE:HD13	1.82	0.61
52:M8:36:CYS:SG	52:M8:37:SER:N	2.72	0.61
26:14:2415:G:H4'	37:35:67:MET:N	2.16	0.61
26:14:38:A:H2'	26:14:39:C:C6	2.36	0.61
26:14:491:G:H2'	26:14:492:A:C8	2.35	0.61
26:14:635:C:O2'	26:14:639:U:OP1	2.18	0.61
26:1H:1729:A:O2'	26:1H:1730:U:O5'	2.18	0.61
26:1H:2101:G:H1	26:1H:2188:C:H42	1.47	0.61
22:1K:53:G:H1	22:1K:61:C:H42	1.48	0.61
7:62:23:VAL:HG13	7:62:43:PHE:HE2	1.66	0.61
1:13:1118:C:H1'	1:13:1179:A:C4	2.35	0.61
1:13:77:C:O2'	1:13:92:G:N2	2.34	0.61
26:14:1021:A:H62	26:14:1141:U:H3	1.48	0.61
26:14:2648:C:H2'	26:14:2649:U:C6	2.36	0.61
33:51:64:LEU:O	33:51:68:THR:OG1	2.18	0.61
34:69:143:SER:OG	34:69:144:VAL:N	2.34	0.61
18:9A:32:ARG:HA	18:9A:69:THR:HG21	1.82	0.61
45:B5:65:ARG:HG3	45:B5:67:GLY:H	1.65	0.61
20:BA:46:GLU:HB2	20:BA:48:LYS:HG2	1.82	0.61
37:35:50:ARG:HD3	56:M5:7:HIS:CD2	2.36	0.61
1:13:153:C:N4	1:13:168:G:H1	1.98	0.61
1:13:595:G:H1	1:13:641:U:HO2'	1.48	0.61
1:13:618:C:H5"	1:13:619:U:H5"	1.83	0.61
1:13:953:G:H5'	1:13:965:A:H61	1.66	0.61
1:1G:1460:A:OP2	20:BA:27:LYS:NZ	2.32	0.61
1:1G:920:U:H2'	1:1G:921:U:C6	2.36	0.61
26:1H:2706:G:O6	62:1H:3761:HOH:O	2.12	0.61
4:32:60:GLU:OE2	4:32:199:ASN:N	2.31	0.61
12:3I:53:ARG:HG3	12:3I:53:ARG:HH11	1.66	0.61
1:1G:564:C:HO2'	8:72:91:ARG:HH22	1.47	0.61
38:88:78:PRO:O	38:88:79:LEU:HB3	2.00	0.61
40:A8:34:HIS:CE1	40:A8:54:LEU:HD23	2.35	0.61
1:13:163:C:O2'	1:13:164:U:O4'	2.19	0.61
1:13:837:G:OP2	1:13:842:C:N4	2.34	0.61
26:14:273(F):C:H3'	26:14:274:G:H5"	1.83	0.61
26:14:2788:C:O2'	26:14:2809:A:N3	2.34	0.61
26:14:34:C:O2'	26:14:35:G:O5'	2.14	0.61
26:1H:587:C:OP2	37:78:21:ARG:NH2	2.34	0.61
3:22:50:ALA:HB2	3:22:75:VAL:HB	1.82	0.61
42:85:92:ARG:NH2	43:95:11:GLN:H	1.98	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
46:C5:88:LYS:O	46:C5:89:PHE:HB3	1.98	0.61
26:14:2611:U:C4	53:J5:3:LYS:HG3	2.36	0.61
50:K8:4:SER:N	50:K8:7:ARG:H	1.98	0.61
26:14:2294:C:P	40:65:89:ARG:HH22	2.24	0.61
35:15:132:ALA:HB1	35:15:133:GLN:HG2	1.83	0.61
29:19:242:ARG:HG2	29:19:246:PRO:HG3	1.82	0.61
2:1E:18:GLY:H	2:1E:42:ILE:HB	1.66	0.61
26:1H:1405:U:H2'	26:1H:1406:U:C6	2.35	0.61
26:1H:2138:C:N3	26:1H:2154:G:N2	2.49	0.61
26:1H:2275:C:H5'	26:1H:2275:C:C6	2.36	0.61
26:1H:71:A:H8	26:1H:71:A:H5'	1.65	0.61
24:3L:5:C:H2'	24:3L:6:G:C8	2.36	0.61
32:49:6:ALA:O	32:49:9:ARG:N	2.34	0.61
35:58:73:THR:HB	35:58:82:LEU:HD11	1.82	0.61
33:59:92:ILE:HG22	33:59:93:GLY:N	2.16	0.61
41:75:26:ASP:O	41:75:49:VAL:HG22	2.00	0.61
2:12:71:VAL:HG11	2:12:164:VAL:HG13	1.82	0.61
26:1H:1449:A:H5'	26:1H:1449(A):G:OP2	2.00	0.61
26:1H:71:A:H5'	26:1H:71:A:C8	2.36	0.61
26:14:872:A:H4'	38:45:66:ILE:HD11	1.83	0.61
41:75:2:ASN:C	41:75:4:GLY:HA3	2.21	0.61
20:BI:49:ALA:HB3	20:BI:99:LEU:HD22	1.81	0.61
49:J8:18:ILE:HG12	49:J8:37:ILE:HG12	1.82	0.61
50:K8:58:ALA:O	50:K8:62:THR:HG22	2.00	0.61
2:12:24:TRP:HE1	2:12:26:PRO:HG3	1.66	0.61
2:12:56:ARG:O	2:12:56:ARG:NH1	2.33	0.61
26:14:1778:U:H2'	26:14:1784:A:N6	2.16	0.61
26:14:2363:C:O2	48:E5:39:ARG:NH2	2.32	0.61
26:14:1786:A:H2	26:14:2606:C:H1'	1.65	0.61
1:1G:1129:C:H5"	1:1G:1139:G:N7	2.16	0.61
1:1G:1273:G:H3'	1:1G:1274:G:H8	1.66	0.61
1:1G:474:G:H2'	1:1G:475:G:C8	2.35	0.61
22:1K:76:A:H8	26:1H:2583:G:H21	1.46	0.61
26:1H:2698:U:H2'	26:1H:2699:C:C6	2.36	0.61
3:22:14:ILE:HG23	3:22:15:THR:HG23	1.81	0.61
30:29:68:ALA:C	30:29:70:ALA:H	2.04	0.61
5:4E:36:ASP:OD2	5:4E:38:GLN:HB2	2.01	0.61
28:71:39:GLU:HG3	28:71:178:ALA:HB2	1.82	0.61
50:K8:2:LYS:HD3	50:K8:6:VAL:HG23	1.83	0.61
52:M8:37:SER:HB3	52:M8:42:PHE:CE2	2.35	0.61
1:13:407:G:OP1	4:3E:115:ARG:NH2	2.34	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1341:U:OP2	26:14:1394:U:O2'	2.16	0.61
26:14:2273:A:H2'	26:14:2274:A:C8	2.36	0.61
1:1G:1007:C:H1'	1:1G:1023:G:N1	2.16	0.61
26:1H:507:A:H5"	26:1H:508:G:H3'	1.83	0.61
26:1H:322:A:P	31:31:168:ARG:HH21	2.24	0.61
41:75:7:ILE:HG13	41:75:8:LYS:N	2.15	0.61
26:14:2232:U:P	49:F5:40:ARG:HH22	2.24	0.61
53:J5:41:PRO:O	53:J5:44:THR:OG1	2.19	0.61
2:12:91:PRO:HG3	2:12:154:LEU:HB2	1.83	0.60
26:14:1149:G:H2'	26:14:1150:C:C6	2.36	0.60
24:3L:71:C:O2'	26:14:1851:U:O2'	2.03	0.60
1:1G:1124:G:HO2'	1:1G:1145:C:N4	1.99	0.60
1:1G:1321:C:N4	1:1G:1322:C:H41	1.99	0.60
26:1H:1639:U:O2'	26:1H:1640:C:H5"	2.01	0.60
36:25:49:ARG:HA	36:25:53:LYS:NZ	2.15	0.60
25:4L:14:A:O2'	25:4L:15:A:O5'	2.19	0.60
26:1H:252:G:OP2	37:78:50:ARG:NH1	2.34	0.60
8:7E:36:LEU:HA	8:7E:39:LEU:HB2	1.83	0.60
20:BA:86:ARG:NH1	20:BA:86:ARG:HB2	2.16	0.60
48:E5:38:VAL:HG13	48:E5:59:LEU:HB2	1.82	0.60
46:G8:42:VAL:HG23	46:G8:43:ASN:H	1.66	0.60
44:E8:38:TYR:OH	53:N8:47:PRO:HG2	2.01	0.60
29:11:2:ALA:HA	29:11:20:ASP:HB2	1.83	0.60
1:13:438:G:H4'	4:3E:123:HIS:CD2	2.35	0.60
1:13:677:U:H3	1:13:713:G:H22	1.50	0.60
26:14:2016:U:O2	53:J5:7:PRO:HG2	2.01	0.60
29:19:182:LEU:N	29:19:272:ALA:HB3	2.14	0.60
26:1H:863:A:H2'	26:1H:864:G:C8	2.36	0.60
30:29:60:ASN:ND2	30:29:62:PRO:O	2.34	0.60
23:2L:62:C:H2'	23:2L:63:C:H6	1.66	0.60
40:65:106:ARG:HA	40:65:110:LEU:HD11	1.82	0.60
8:7E:87:SER:HB2	8:7E:93:VAL:CB	2.27	0.60
26:1H:2591:C:P	29:11:239:ARG:HG3	2.41	0.60
1:13:443:C:H42	1:13:491:G:H1	1.46	0.60
26:14:524:U:H2'	26:14:525:U:C6	2.36	0.60
29:19:264:LYS:HE2	29:19:266:SER:HB3	1.82	0.60
1:1G:1392:G:H21	1:1G:1502:A:H8	1.49	0.60
26:1H:511:U:OP2	62:1H:3770:HOH:O	2.16	0.60
37:35:59:LEU:O	37:35:59:LEU:HD22	2.01	0.60
31:39:24:LEU:HD22	31:39:25:PRO:HD3	1.83	0.60
24:3K:14:A:H2'	24:3K:15:G:C8	2.36	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:F5:90:ILE:HA	49:F5:93:GLU:OE1	2.02	0.60
26:14:528:A:C2	26:14:2043:C:H4'	2.37	0.60
2:1E:5:ILE:HB	2:1E:221:LEU:HD23	1.83	0.60
1:1G:41:G:H2'	1:1G:42:G:C8	2.36	0.60
26:1H:1026:U:H4'	26:1H:1027:A:OP1	2.00	0.60
26:1H:1991:U:H2'	26:1H:1992:G:H5"	1.83	0.60
26:1H:2062:A:OP1	62:1H:3771:HOH:O	2.16	0.60
26:1H:451:C:H5'	62:1H:3709:HOH:O	1.94	0.60
30:29:64:LYS:N	30:29:73:GLU:OE2	2.34	0.60
4:32:31:CYS:HB2	4:32:33:MET:O	2.01	0.60
26:14:2467:C:H4'	38:45:123:HIS:CD2	2.35	0.60
7:62:22:LEU:HD23	7:62:62:PHE:HE2	1.66	0.60
1:13:1347:G:H5"	9:8E:107:ARG:HB3	1.83	0.60
20:BI:30:LYS:HE3	20:BI:80:ARG:HH22	1.65	0.60
26:14:498:G:H21	46:C5:47:LYS:HZ2	1.49	0.60
46:C5:17:SER:HB2	46:C5:71:LYS:HE2	1.83	0.60
1:13:1367:C:H5'	10:1I:60:ARG:NH1	2.15	0.60
1:13:1366:C:H2'	1:13:1367:C:H6	1.66	0.60
26:14:2128:C:H42	26:14:2160:G:H1	1.50	0.60
2:1E:21:ARG:O	2:1E:23:ARG:N	2.33	0.60
1:1G:1021:G:H2'	1:1G:1022:G:C8	2.36	0.60
26:1H:443:A:H1'	26:1H:1201:C:O4'	2.01	0.60
26:1H:2321:G:H5"	62:1H:3776:HOH:O	2.01	0.60
4:3E:31:CYS:SG	4:3E:33:MET:HB2	2.42	0.60
41:B8:58:ASN:C	41:B8:58:ASN:HD22	2.03	0.60
41:B8:26:ASP:CB	41:B8:92:GLY:H	2.14	0.60
26:14:517:C:OP1	53:J5:16:ARG:NH2	2.34	0.60
29:11:37:LEU:HD23	29:11:62:TYR:HB2	1.84	0.60
1:13:1073:U:OP2	5:4E:57:LYS:NZ	2.35	0.60
1:13:1145:C:H4'	1:13:1146:A:H5'	1.83	0.60
26:14:2378:A:H4'	40:65:23:ARG:HH11	1.66	0.60
27:16:15:A:H5'	27:16:16:G:C8	2.37	0.60
37:35:85:LEU:HA	37:35:88:LEU:HB2	1.84	0.60
26:14:38:A:H1'	31:39:48:THR:HB	1.83	0.60
24:3K:62:C:H2'	28:71:53:ARG:HH21	1.65	0.60
13:4I:58:GLU:O	13:4I:62:ASN:ND2	2.31	0.60
6:52:83:ASP:N	6:52:83:ASP:OD1	2.34	0.60
39:55:38:VAL:HG12	39:55:42:LYS:HD2	1.83	0.60
19:AA:10:PHE:H	19:AA:11:VAL:HB	1.65	0.60
26:14:125:G:H5"	55:L5:19:ARG:HD3	1.82	0.60
26:14:1817:G:OP1	29:19:88:ARG:NH2	2.33	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2118:U:O2	26:14:2147:G:N2	2.34	0.60
26:14:2734:A:H2'	26:14:2735:G:O4'	2.01	0.60
26:14:796:C:H2'	26:14:797:C:C6	2.36	0.60
33:51:153:LYS:CB	33:51:155:SER:H	2.14	0.60
33:59:6:ARG:HB3	33:59:66:GLY:HA2	1.84	0.60
6:5E:80:ARG:NH1	6:5E:88:VAL:O	2.34	0.60
37:78:19:VAL:HG13	37:78:31:ALA:HB1	1.82	0.60
26:1H:910:A:N7	38:88:13:GLN:HG3	2.17	0.60
38:88:77:LYS:HE3	38:88:84:GLY:O	2.01	0.60
26:14:582:G:H2'	26:14:583:G:C8	2.37	0.60
26:1H:2315:G:OP1	32:41:36:LYS:NZ	2.33	0.60
26:1H:870:A:OP1	38:88:5:ARG:NH2	2.34	0.60
3:22:70:VAL:HG21	3:22:76:VAL:HG11	1.84	0.60
11:2I:53:SER:HA	11:2I:54:ARG:C	2.22	0.60
31:39:66:PRO:O	31:39:67:GLN:HB3	2.01	0.60
38:45:77:LYS:HE3	38:45:84:GLY:H	1.65	0.60
7:6E:15:ASP:HB3	7:6E:20:ASP:H	1.65	0.60
41:B8:77:PRO:HG2	41:B8:80:SER:HB2	1.83	0.60
35:58:38:HIS:O	42:C8:67:ALA:HB1	2.02	0.60
37:35:50:ARG:HG2	56:M5:61:LEU:HD21	1.83	0.60
29:11:70:TRP:O	29:11:73:VAL:HG23	2.02	0.60
2:12:118:LEU:HD11	2:12:141:GLU:HG2	1.82	0.60
26:14:1430:C:H2'	26:14:1431:U:H6	1.66	0.60
26:1H:1420:U:O2'	26:1H:1421:G:OP1	2.16	0.60
4:32:127:THR:HG21	4:32:149:ALA:HB2	1.83	0.60
5:42:105:VAL:HG21	5:42:128:PRO:HB3	1.84	0.60
32:49:124:SER:HB2	32:49:131:TYR:CE1	2.36	0.60
1:13:1226:C:H2'	13:4I:103:THR:HB	1.84	0.60
34:69:57:ARG:O	34:69:61:ARG:HB2	2.01	0.60
41:B8:64:ARG:HB2	41:B8:73:GLU:HG2	1.84	0.60
50:G5:4:SER:HA	50:G5:7:ARG:H	1.67	0.60
49:J8:64:ALA:HA	49:J8:67:ILE:HG13	1.83	0.60
45:F8:5:TYR:CE1	50:K8:30:ARG:HG3	2.36	0.60
50:K8:4:SER:OG	50:K8:4:SER:O	2.15	0.60
1:13:1129:C:H42	1:13:1143:G:H1	1.49	0.60
1:13:310:G:OP2	16:7I:27:LYS:NZ	2.33	0.60
26:14:1593:G:H2'	26:14:1594:G:C8	2.37	0.60
26:14:2409:G:N7	62:14:3692:HOH:O	2.31	0.60
26:14:2577:A:H5'	53:J5:3:LYS:HD3	1.84	0.60
26:14:270(X):G:OP2	62:14:3641:HOH:O	2.16	0.60
26:1H:674:G:C1'	31:31:74:ARG:HD3	2.29	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:796:C:H2'	26:1H:797:C:C6	2.37	0.60
3:22:180:ALA:HB1	3:22:203:PHE:HE1	1.66	0.60
31:39:129:PHE:HA	31:39:142:TRP:NE1	2.16	0.60
24:3K:57:G:N2	24:3K:60:U:O4	2.35	0.60
41:B8:90:GLN:OE1	41:B8:91:ARG:N	2.32	0.60
20:BA:12:ALA:O	20:BA:15:ARG:N	2.34	0.60
26:1H:1570:A:H2'	26:1H:1571:A:C8	2.36	0.59
10:1I:16:LEU:HD11	10:1I:70:ARG:HB2	1.84	0.59
13:4A:12:ASN:O	13:4A:12:ASN:ND2	2.31	0.59
33:51:107:VAL:HB	33:51:152:ARG:HG2	1.84	0.59
33:59:152:ARG:HG3	33:59:153:LYS:HB2	1.84	0.59
14:5A:26:ARG:HH12	14:5A:47:LEU:HD21	1.65	0.59
9:8E:28:VAL:HA	9:8E:63:ILE:O	2.02	0.59
49:F5:4:VAL:HG12	49:F5:11:ARG:HB3	1.83	0.59
51:L8:9:VAL:HG12	51:L8:53:LEU:O	2.02	0.59
1:13:1129:C:H1'	1:13:1146:A:H61	1.67	0.59
1:1G:979:C:H3'	1:1G:980:C:H5"	1.84	0.59
26:1H:2432:A:C4	49:J8:33:LYS:HG2	2.38	0.59
1:1G:1423:G:OP1	36:25:49:ARG:NH2	2.34	0.59
31:39:116:ASP:OD2	37:35:1:MET:N	2.35	0.59
9:8E:114:TYR:HE2	10:1I:59:SER:HA	1.67	0.59
47:D5:80:ARG:HH11	47:D5:82:ARG:HH21	1.50	0.59
29:11:24:ILE:HG23	29:11:83:GLU:HA	1.84	0.59
26:14:2693:A:H2'	26:14:2694:G:H8	1.67	0.59
26:14:2713:A:OP2	62:14:3636:HOH:O	2.16	0.59
26:14:2745:C:H1'	33:59:143:GLN:HG2	1.83	0.59
29:19:31:LYS:NZ	29:19:33:LEU:HB3	2.17	0.59
26:1H:1332:G:H21	26:1H:1610:A:H8	1.48	0.59
26:1H:243:U:OP1	56:Q8:6:THR:OG1	2.20	0.59
26:1H:270(F):U:H2'	26:1H:270(G):C:C6	2.37	0.59
9:8E:7:THR:O	9:8E:83:ARG:NH1	2.35	0.59
26:14:495:G:N3	44:A5:61:ASN:ND2	2.51	0.59
47:D5:139:VAL:HA	47:D5:156:LYS:HE3	1.84	0.59
26:14:850:C:O3'	51:H5:49:LYS:HE2	2.00	0.59
50:K8:23:LYS:NZ	50:K8:27:GLU:OE2	2.35	0.59
52:M8:13:ARG:HA	52:M8:24:THR:HG21	1.85	0.59
1:13:868:C:H2'	1:13:869:G:O4'	2.03	0.59
26:14:1927:A:H2'	26:14:1928:A:C8	2.37	0.59
29:19:39:LYS:CG	29:19:40:THR:H	2.11	0.59
2:1E:185:ILE:HB	2:1E:199:TYR:HB2	1.85	0.59
1:1G:164:U:H2'	1:1G:165:C:C6	2.38	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1932:A:H2'	26:1H:1933:G:O4'	2.02	0.59
26:1H:577:G:OP2	62:1H:3772:HOH:O	2.17	0.59
30:29:119:ARG:HG2	30:29:160:TYR:HB2	1.84	0.59
3:2E:91:LEU:HB2	3:2E:99:VAL:HG21	1.85	0.59
11:2I:85:ARG:HG2	11:2I:112:THR:H	1.68	0.59
23:2L:41:C:H2'	23:2L:42:C:H6	1.66	0.59
38:45:133:ARG:O	38:45:134:ARG:HG3	2.03	0.59
38:45:35:VAL:HG12	38:45:36:ALA:H	1.67	0.59
15:6I:25:THR:HG21	15:6I:70:LEU:HB2	1.84	0.59
9:8E:47:LEU:HB2	9:8E:50:LEU:HD11	1.83	0.59
44:E8:92:ARG:NH1	44:E8:94:ASP:OD1	2.36	0.59
46:G8:42:VAL:HG23	46:G8:43:ASN:N	2.18	0.59
48:I8:72:ARG:HB3	48:I8:75:LEU:HB2	1.84	0.59
26:14:2577:A:O4'	53:J5:3:LYS:HB2	2.01	0.59
50:K8:3:LEU:H	50:K8:5:GLU:N	2.00	0.59
2:12:114:ARG:NH1	2:12:117:GLU:OE1	2.36	0.59
29:19:37:LEU:H	29:19:37:LEU:CD1	2.06	0.59
2:1E:212:GLN:NE2	2:1E:233:SER:O	2.36	0.59
1:1G:1160:G:H1	1:1G:1176:A:H61	1.49	0.59
26:1H:86:C:H4'	26:1H:104:U:H1'	1.82	0.59
26:1H:2801:A:H5'	26:1H:2895:U:H1'	1.84	0.59
27:1J:89(A):A:H5'	27:1J:90:C:OP2	2.03	0.59
4:3E:84:LYS:HB3	4:3E:86:LYS:HG3	1.84	0.59
7:6E:69:VAL:HG12	7:6E:100:ALA:HA	1.85	0.59
28:71:64:LEU:HD21	28:71:188:ASN:ND2	2.18	0.59
41:75:2:ASN:HB3	41:75:4:GLY:O	2.03	0.59
41:B8:24:PRO:HA	41:B8:49:VAL:HG22	1.83	0.59
1:13:1126:U:H2'	1:13:1127:G:H5'	1.84	0.59
26:14:2062:A:O2'	26:14:2063:C:OP1	2.19	0.59
26:14:570:G:H5"	62:14:4024:HOH:O	2.01	0.59
29:19:37:LEU:N	29:19:37:LEU:HD12	2.18	0.59
1:1G:191(F):U:H3	20:BA:105:SER:HG	1.48	0.59
1:1G:371:G:H1	1:1G:390:C:H42	1.50	0.59
26:1H:2176:A:H5'	28:71:221:SER:HB3	1.83	0.59
26:1H:860:U:H5	26:1H:917:A:N1	2.00	0.59
10:1I:40:LEU:HB2	10:1I:69:ASN:HB2	1.83	0.59
27:1J:15:A:H3'	27:1J:16:G:H5'	1.83	0.59
30:21:116:VAL:HG11	30:21:138:PRO:HB3	1.84	0.59
3:22:182:ILE:HG23	3:22:203:PHE:HD1	1.66	0.59
31:31:197:ASP:N	31:31:197:ASP:OD1	2.33	0.59
33:51:59:ARG:O	33:51:63:SER:OG	2.20	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:52:35:ALA:HB1	6:52:65:VAL:HG11	1.85	0.59
33:59:32:GLU:H	33:59:32:GLU:CD	2.06	0.59
38:88:78:PRO:HB2	38:88:81:VAL:HG11	1.83	0.59
9:8E:28:VAL:HG22	9:8E:63:ILE:HB	1.83	0.59
20:BI:73:HIS:HB3	20:BI:74:LYS:HG2	1.85	0.59
47:H8:111:VAL:HG11	47:H8:146:ILE:HG13	1.85	0.59
54:O8:43:CYS:HB3	54:O8:44:ARG:NH1	2.17	0.59
26:14:782:A:H5'	26:14:783:A:C2	2.38	0.59
35:15:130:HIS:HB3	35:15:134:ARG:HH21	1.67	0.59
1:1G:79:G:N2	1:1G:90:C:N3	2.46	0.59
26:1H:49:A:N7	26:1H:120:U:C5	2.66	0.59
8:7E:73:ASP:OD1	8:7E:75:ARG:NE	2.36	0.59
45:B5:32:PRO:HA	45:B5:77:LYS:HB2	1.85	0.59
54:O8:27:LYS:HZ2	54:O8:27:LYS:H	1.50	0.59
26:14:1449(A):G:H2'	26:14:1450:C:H6	1.67	0.59
26:14:1568:G:P	29:19:63:ARG:HH12	2.26	0.59
26:1H:827:U:H2'	26:1H:2430:A:C2	2.38	0.59
10:1I:86:MET:SD	10:1I:86:MET:N	2.76	0.59
12:3I:93:LEU:O	12:3I:96:VAL:HG13	2.02	0.59
13:4A:54:VAL:HA	13:4A:57:ARG:HB3	1.83	0.59
6:5E:100:ASN:HB2	18:9I:28:GLU:HA	1.85	0.59
37:78:36:LYS:O	37:78:40:SER:HB3	2.03	0.59
1:13:1347:G:C8	9:8E:107:ARG:HB2	2.38	0.59
26:1H:2208:U:H4'	29:11:151:LYS:HG2	1.83	0.59
26:14:1019:U:H3	26:14:1142(A):A:H62	1.50	0.59
1:1G:973:G:H5'	10:1A:55:LYS:HZ2	1.68	0.59
26:1H:1230:C:H2'	26:1H:1231:G:C8	2.38	0.59
26:1H:1762:A:N1	62:1H:3823:HOH:O	2.32	0.59
26:1H:1678:G:N2	26:1H:1989:G:H22	1.97	0.59
26:1H:736:C:H42	26:1H:760:G:H1	1.51	0.59
26:1H:994:C:H3'	42:C8:54:LYS:HE3	1.85	0.59
1:1G:490:G:P	4:32:132:ARG:HH22	2.25	0.59
13:4I:12:ASN:OD1	13:4I:46:LYS:NZ	2.30	0.59
8:7E:7:ALA:HB2	8:7E:85:ARG:HH11	1.68	0.59
46:G8:38:ILE:HD11	46:G8:64:GLU:HG3	1.83	0.59
2:12:33:TYR:HB3	2:12:41:ILE:HG23	1.85	0.59
26:14:1198:U:H2'	26:14:1199:U:H6	1.66	0.59
2:1E:231:GLU:OE1	2:1E:231:GLU:N	2.31	0.59
1:1G:1157:A:H2	1:1G:1180:A:C6	2.21	0.59
1:1G:1469:G:H2'	1:1G:1470:G:C8	2.37	0.59
26:1H:1111:A:N3	26:1H:1112:G:H1'	2.18	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1858:G:O2'	26:1H:1859:A:OP2	2.21	0.59
31:39:3:GLU:HA	31:39:24:LEU:HD12	1.85	0.59
34:61:110:ASP:OD2	34:61:113:ARG:HD3	2.03	0.59
34:69:110:ASP:N	34:69:130:TYR:OH	2.26	0.59
15:6I:16:ALA:HB1	15:6I:21:ASP:HB3	1.85	0.59
9:82:36:TYR:OH	9:82:73:GLN:NE2	2.33	0.59
51:H5:39:ASP:O	51:H5:44:ARG:NH1	2.36	0.59
26:1H:593:G:H1'	56:Q8:4:MET:HE1	1.84	0.59
1:13:1435:G:H2'	1:13:1436:U:C6	2.38	0.58
1:13:735:C:H2'	1:13:736:C:C6	2.37	0.58
26:14:1028:A:N6	26:14:1125:G:H2'	2.18	0.58
3:2E:58:GLU:H	3:2E:65:ALA:HB3	1.69	0.58
56:Q8:52:LYS:H	56:Q8:53:PRO:HD2	1.68	0.58
26:14:2819:G:N7	62:14:3699:HOH:O	2.32	0.58
26:14:548:A:C6	26:14:549:G:H1'	2.39	0.58
26:14:71:A:H2	45:B5:31:HIS:NE2	1.99	0.58
26:14:824:A:H1'	26:14:2358:G:N7	2.18	0.58
1:1G:581:G:OP1	15:6A:61:GLY:HA3	2.03	0.58
26:1H:2306:C:H3'	26:1H:2307:G:C5'	2.33	0.58
26:1H:85:G:OP2	46:G8:9:LYS:HB2	2.03	0.58
30:21:152:LYS:HD3	35:58:77:GLY:HA3	1.85	0.58
40:65:62:LYS:O	40:65:66:ALA:N	2.35	0.58
1:13:1259:C:N4	1:13:1260:C:O2	2.36	0.58
1:13:1292:U:H2'	1:13:1293:G:C8	2.37	0.58
26:14:375:C:H2'	26:14:376:C:C6	2.37	0.58
1:1G:1018:C:H2'	1:1G:1019:C:O4'	2.04	0.58
1:1G:838:G:H1	1:1G:848:C:H42	1.52	0.58
26:1H:614:U:H6	26:1H:614:U:OP2	1.86	0.58
4:32:31:CYS:H	4:32:35:ARG:CZ	2.16	0.58
24:3K:57:G:H2'	24:3K:58:A:H5'	1.85	0.58
24:3K:5:C:H2'	24:3K:6:G:H8	1.68	0.58
5:42:69:VAL:O	5:42:71:LEU:N	2.35	0.58
32:49:118:ARG:HB3	32:49:181:ARG:HD3	1.85	0.58
13:4I:60:VAL:HG13	13:4I:64:TRP:HE1	1.67	0.58
39:98:55:ALA:HB1	39:98:84:ALA:HB2	1.86	0.58
43:D8:65:GLY:HA3	43:D8:91:TYR:CE1	2.38	0.58
50:K8:28:LYS:HB3	50:K8:53:LEU:HD21	1.84	0.58
1:13:1000:A:H2'	1:13:1001:G:C8	2.38	0.58
26:14:1420:U:O2'	26:14:1421:G:OP1	2.20	0.58
1:1G:426:G:OP1	4:32:36:ARG:NH2	2.35	0.58
26:1H:2129:C:OP2	28:71:36:LYS:NZ	2.36	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:25:24:VAL:HB	36:25:33:ALA:HB2	1.85	0.58
11:2I:59:TYR:CE2	11:2I:63:LEU:HD11	2.39	0.58
31:31:101:LEU:O	31:31:106:ARG:NH1	2.37	0.58
31:39:170:LEU:HD22	31:39:172:TRP:HE1	1.68	0.58
33:59:10:PRO:HD2	33:59:50:VAL:HG13	1.85	0.58
14:5A:45:ARG:O	14:5A:49:HIS:ND1	2.33	0.58
7:62:26:PHE:O	7:62:30:ILE:HG13	2.03	0.58
30:29:12:THR:HG22	41:75:58:ASN:OD1	2.02	0.58
48:E5:47:PRO:HG3	48:E5:53:MET:HB2	1.84	0.58
1:13:1117:G:H5"	9:8E:104:ARG:NH1	2.18	0.58
1:13:652:U:O2'	1:13:653:A:O5'	2.21	0.58
26:1H:2776:A:H4'	26:1H:2777:G:H5"	1.85	0.58
26:1H:671:C:OP1	37:78:42:SER:O	2.20	0.58
36:25:4:PRO:O	36:25:5:GLN:HB2	2.03	0.58
4:3E:173:TRP:CD1	4:3E:174:LEU:HG	2.38	0.58
40:65:107:GLU:H	40:65:110:LEU:HD11	1.69	0.58
37:78:46:LYS:O	37:78:47:ASP:HB3	2.03	0.58
16:7I:71:ARG:O	16:7I:75:ARG:N	2.35	0.58
26:1H:1278:A:OP1	39:98:36:THR:HG22	2.04	0.58
39:98:72:ASP:O	39:98:76:VAL:HG23	2.04	0.58
2:12:121:LEU:HG	2:12:126:GLU:HG2	1.85	0.58
1:13:7:G:H5'	1:13:298:A:O4'	2.03	0.58
26:14:2287:A:H62	26:14:2344:U:H3	1.48	0.58
26:14:2779:U:OP1	62:14:3640:HOH:O	2.16	0.58
26:14:67:U:N3	26:14:74:A:H2	1.95	0.58
26:14:821:A:O2'	26:14:946:G:OP2	2.21	0.58
26:1H:1512:G:H2'	26:1H:1513:C:C6	2.39	0.58
26:1H:1605:C:O3'	62:1H:3773:HOH:O	2.17	0.58
36:25:13:ASN:HD21	36:25:97:ARG:H	1.51	0.58
31:31:116:ASP:O	31:31:120:GLU:HG3	2.04	0.58
36:68:68:GLU:OE2	36:68:78:ARG:NH1	2.36	0.58
9:82:119:ALA:O	9:82:120:ARG:HB2	2.03	0.58
39:98:27:SER:HB3	39:98:34:ILE:HD11	1.84	0.58
20:BA:49:ALA:HA	20:BA:52:ALA:HB3	1.86	0.58
2:12:73:THR:HG21	2:12:97:TRP:H	1.68	0.58
1:13:1510:U:H2'	1:13:1511:G:C8	2.39	0.58
1:13:667:G:H4'	15:6I:51:HIS:ND1	2.19	0.58
1:13:685:G:O2'	1:13:686:U:H5'	2.04	0.58
1:13:93:U:H5'	1:13:95:G:OP2	2.04	0.58
26:14:1292:U:H2'	26:14:1293:C:C6	2.38	0.58
26:14:1794:U:H2'	26:14:1795:C:C6	2.38	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2572:A:OP1	26:14:2574:G:O2'	2.22	0.58
26:14:2648:C:H2'	26:14:2649:U:H6	1.66	0.58
26:14:2749:A:N1	26:14:2750:A:N6	2.52	0.58
26:14:2830:G:O6	62:14:3642:HOH:O	2.16	0.58
29:19:17:THR:O	29:19:211:ARG:NH2	2.37	0.58
1:1G:1305:G:O2'	1:1G:1306:A:H8	1.87	0.58
30:21:131:ALA:HB1	62:21:403:HOH:O	2.03	0.58
30:29:53:PRO:HA	30:29:74:PRO:HB3	1.86	0.58
31:39:85:GLY:O	62:39:401:HOH:O	2.17	0.58
32:41:16:ARG:O	32:41:20:ILE:HG13	2.03	0.58
33:51:20:ALA:HB3	33:51:23:ARG:HG3	1.86	0.58
35:58:130:HIS:C	35:58:134:ARG:HH12	2.07	0.58
44:A5:71:VAL:HA	44:A5:107:LEU:HD12	1.86	0.58
41:B8:108:ARG:HA	41:B8:111:ARG:NE	2.19	0.58
20:BI:26:ASN:O	20:BI:30:LYS:HB2	2.03	0.58
1:13:1226:C:OP2	13:4I:103:THR:OG1	2.15	0.58
1:13:376:G:H1	1:13:387:U:H3	1.50	0.58
26:14:2391:G:O6	26:14:2425:A:H8	1.87	0.58
2:1E:73:THR:HG22	2:1E:74:LYS:HG2	1.86	0.58
1:1G:247:G:OP2	17:8A:100:LYS:HA	2.04	0.58
1:1G:727:G:N2	1:1G:730:G:OP2	2.29	0.58
26:1H:1331:A:O2'	26:1H:1332:G:H8	1.87	0.58
26:1H:2211:G:O2'	26:1H:2212:A:OP1	2.20	0.58
26:1H:330:A:HO2'	26:1H:331:A:H8	1.52	0.58
26:1H:960:A:H2'	26:1H:962:G:H5'	1.86	0.58
3:22:121:ALA:HB1	3:22:189:ALA:HB2	1.85	0.58
31:31:51:THR:O	31:31:93:LYS:HE2	2.04	0.58
47:D5:132:ASN:HD22	47:D5:159:PRO:HG2	1.68	0.58
1:1G:19:C:OP1	5:42:125:SER:OG	2.18	0.58
1:1G:572:A:H5'	1:1G:573:A:OP2	2.03	0.58
1:1G:972:C:O3'	10:1A:55:LYS:NZ	2.37	0.58
26:1H:2145:C:H5	26:1H:2148:G:H21	1.50	0.58
26:1H:2401:U:H2'	26:1H:2402:C:O4'	2.04	0.58
1:13:963:G:N3	10:1I:55:LYS:NZ	2.52	0.58
26:14:607:U:OP1	31:39:102:PRO:HA	2.04	0.58
31:39:92:PRO:O	31:39:93:LYS:HD2	2.02	0.58
26:1H:2751:G:N7	33:51:3:ARG:NH2	2.52	0.58
33:59:35:VAL:HG11	33:59:72:ILE:HG12	1.84	0.58
28:71:45:ALA:HB2	28:71:212:VAL:HG22	1.86	0.58
49:F5:88:LYS:O	49:F5:91:LYS:HB3	2.04	0.58
46:G8:89:PHE:HD1	46:G8:90:LEU:N	2.02	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
46:G8:76:CYS:SG	46:G8:97:ARG:HG3	2.43	0.58
2:12:165:VAL:HG23	2:12:166:ASP:H	1.68	0.58
1:13:964:A:N3	1:13:969:A:O2'	2.33	0.58
26:14:2776:A:OP1	26:14:2776:A:H3'	2.04	0.58
1:1G:1:U:H1'	1:1G:2:U:H5'	1.86	0.58
26:1H:1113:U:H5'	33:51:2:SER:OG	2.03	0.58
26:1H:1264:G:H5'	53:N8:11:THR:HG21	1.86	0.58
26:1H:2327:A:H2'	26:1H:2328:A:C8	2.39	0.58
24:1L:8:U:H3'	24:1L:13:C:H42	1.69	0.58
23:2L:48:U:O2'	23:2L:49:C:OP2	2.21	0.58
31:31:160:ASN:OD1	31:31:163:VAL:HG23	2.04	0.58
31:31:66:PRO:HD2	31:31:70:THR:HG21	1.86	0.58
4:32:13:ARG:C	4:32:15:GLU:H	2.07	0.58
32:41:17:PRO:HA	32:41:20:ILE:HD12	1.85	0.58
32:49:106:LEU:O	32:49:111:LEU:HD12	2.04	0.58
25:4L:20:A:H2'	25:4L:21:A:O4'	2.04	0.58
6:52:70:ASP:OD1	6:52:70:ASP:N	2.34	0.58
51:H5:46:ASN:O	51:H5:50:VAL:HG22	2.04	0.58
56:M5:54:GLU:O	56:M5:58:ILE:HG23	2.04	0.58
2:12:78:GLN:O	2:12:94:ASN:ND2	2.36	0.57
1:13:1239:A:H62	1:13:1299:A:H62	1.51	0.57
26:1H:827:U:H2'	26:1H:2430:A:H2	1.69	0.57
26:1H:994:C:OP1	42:C8:53:ARG:NH2	2.37	0.57
31:39:164:ARG:O	31:39:167:ALA:HB3	2.03	0.57
13:4A:34:LEU:HD13	13:4A:41:PRO:HB3	1.86	0.57
38:88:58:PHE:O	38:88:60:ARG:N	2.36	0.57
20:BI:57:ARG:HH22	20:BI:100:ILE:HD13	1.68	0.57
49:J8:73:LEU:HD13	49:J8:93:GLU:HB3	1.86	0.57
26:14:2528:U:O2'	26:14:2530:A:OP1	2.18	0.57
2:1E:189:ASP:CG	2:1E:205:ASP:HB3	2.24	0.57
26:1H:2392:A:H2	26:1H:2424:C:N4	2.02	0.57
30:21:53:PRO:HA	30:21:75:VAL:N	2.16	0.57
13:4A:91:ARG:NH1	13:4A:97:PRO:O	2.37	0.57
1:1G:1346:A:H5"	9:82:120:ARG:NH1	2.18	0.57
42:85:97:ASP:OD1	42:85:98:LEU:N	2.37	0.57
46:C5:17:SER:HB2	46:C5:71:LYS:CE	2.34	0.57
1:1G:1028:C:N4	1:1G:1033:G:H1	2.02	0.57
1:1G:1104:G:H4'	2:12:111:ARG:HE	1.69	0.57
26:1H:309:G:H4'	46:G8:18:GLY:HA2	1.86	0.57
3:22:44:GLU:HG3	3:22:52:LEU:HD21	1.86	0.57
31:31:155:LEU:HD11	31:31:176:LEU:HD13	1.86	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2415:G:H4'	37:35:67:MET:H	1.69	0.57
24:3K:9:A:O2'	24:3K:46:G:O4'	2.19	0.57
20:BI:86:ARG:O	20:BI:90:GLN:NE2	2.37	0.57
47:D5:158:PRO:HB2	47:D5:159:PRO:HD2	1.86	0.57
45:F8:29:TRP:CZ3	45:F8:78:LYS:HG2	2.40	0.57
52:M8:12:ALA:HB3	52:M8:24:THR:HB	1.86	0.57
29:11:71:ASP:N	29:11:71:ASP:OD1	2.36	0.57
1:13:313:A:H2'	1:13:314:C:C6	2.40	0.57
1:1G:222:U:H2'	1:1G:223:U:C6	2.40	0.57
1:1G:458:C:H2'	1:1G:464:G:H8	1.70	0.57
26:1H:2128:C:H2'	26:1H:2129:C:C6	2.39	0.57
36:68:63:VAL:HG12	36:68:106:LEU:HD11	1.87	0.57
8:72:36:LEU:HA	8:72:39:LEU:HB2	1.87	0.57
48:E5:36:ILE:HD11	48:E5:39:ARG:HG2	1.85	0.57
1:13:456:C:H42	1:13:476:G:H1	1.50	0.57
26:14:315:G:H2'	26:14:316:C:C6	2.39	0.57
21:1F:5:ASP:HB3	21:1F:8:THR:HG22	1.87	0.57
1:1G:1203:C:H2'	1:1G:1204:A:H8	1.69	0.57
1:1G:297:G:N2	1:1G:300:A:OP2	2.36	0.57
1:1G:44:G:H2'	1:1G:45:U:O4'	2.04	0.57
26:1H:760:G:H5"	62:1H:3668:HOH:O	2.05	0.57
4:32:76:ARG:NH2	4:32:80:GLU:OE1	2.38	0.57
31:39:28:ILE:HA	31:39:112:MET:HG2	1.85	0.57
27:1J:90:C:P	38:45:16:ARG:HH21	2.27	0.57
33:51:77:LYS:HE2	33:51:138:LYS:HD2	1.84	0.57
34:61:113:ARG:HH21	34:61:132:PRO:HB3	1.70	0.57
16:7A:8:ARG:HD3	16:7A:17:TYR:CE1	2.40	0.57
9:82:77:ILE:O	9:82:81:ILE:HG12	2.04	0.57
42:85:91:ASP:OD1	42:85:96:ALA:HB2	2.03	0.57
9:8E:43:ALA:HA	9:8E:74:ILE:HD13	1.84	0.57
40:A8:35:ILE:HD11	40:A8:101:LEU:HD23	1.87	0.57
26:14:138:G:N2	45:B5:44:GLU:OE2	2.25	0.57
47:D5:113:ALA:O	47:D5:116:VAL:N	2.23	0.57
47:D5:115:GLY:HA3	47:D5:174:VAL:HG13	1.86	0.57
51:L8:7:LYS:HE2	51:L8:32:GLN:O	2.05	0.57
1:13:158:G:H2'	1:13:159:G:H8	1.68	0.57
26:14:2378:A:O2'	40:65:21:THR:HG21	2.05	0.57
1:1G:976:G:OP1	14:5A:32:SER:N	2.36	0.57
26:1H:1042:G:H1	26:1H:1113:U:H3	1.51	0.57
26:1H:125:G:H5'	26:1H:125:G:C8	2.39	0.57
26:1H:2061:G:OP2	26:1H:2502:G:H5'	2.04	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2257:U:O2'	26:1H:2258:C:H5'	2.05	0.57
31:31:129:PHE:HA	31:31:142:TRP:NE1	2.19	0.57
33:51:164:TYR:O	33:51:167:GLU:HB3	2.05	0.57
1:13:750:G:N3	15:6I:23:GLY:HA3	2.20	0.57
1:1G:376:G:H5"	16:7A:5:ARG:HB2	1.85	0.57
19:AI:41:VAL:HG12	19:AI:44:MET:CB	2.34	0.57
45:B5:63:LYS:HA	45:B5:72:LYS:HA	1.87	0.57
42:C8:92:ARG:CZ	43:D8:11:GLN:H	2.16	0.57
46:G8:97:ARG:NH2	46:G8:103:GLY:O	2.34	0.57
29:11:182:LEU:N	29:11:272:ALA:HB3	2.06	0.57
26:14:1381:G:OP2	62:14:3645:HOH:O	2.18	0.57
26:14:2425:A:H5'	26:14:2426:A:H3'	1.87	0.57
1:1G:501:C:H2'	1:1G:502:G:C8	2.37	0.57
26:1H:754:C:H2'	26:1H:755:C:H6	1.70	0.57
23:2L:33:OMC:O5'	23:2L:33:OMC:H6	1.88	0.57
24:3K:15:G:H1	24:3K:48:C:H41	1.52	0.57
24:3K:76:A:H8	26:1H:2394:C:N4	1.99	0.57
13:4A:37:THR:HG22	13:4A:55:ARG:HE	1.70	0.57
17:8I:66:SER:O	17:8I:70:ARG:NH1	2.37	0.57
41:B8:1:MET:N	41:B8:2:ASN:OD1	2.33	0.57
42:C8:88:ILE:O	42:C8:90:VAL:N	2.38	0.57
47:D5:157:LEU:HB3	47:D5:161:VAL:HG21	1.86	0.57
2:12:110:GLN:HB3	2:12:111:ARG:HH12	1.70	0.57
1:13:1266:G:N2	1:13:1270:C:N3	2.52	0.57
26:14:1614:A:OP1	26:14:1617:C:N4	2.31	0.57
2:1E:209:ARG:HD2	2:1E:239:VAL:HG13	1.85	0.57
1:1G:1025:U:H5'	1:1G:1026:G:H5'	1.85	0.57
1:1G:1127:G:H2'	1:1G:1128:C:C6	2.40	0.57
1:1G:1154:G:H2'	1:1G:1155:G:C8	2.38	0.57
1:1G:1345:U:OP1	62:1G:1861:HOH:O	2.17	0.57
1:1G:135:C:O2	16:7A:1:MET:HB3	2.05	0.57
1:1G:458:C:N3	1:1G:474:G:N2	2.49	0.57
1:1G:87:A:H4'	1:1G:88:C:OP1	2.03	0.57
3:22:58:GLU:HB2	3:22:65:ALA:HB3	1.86	0.57
5:4E:142:LEU:O	5:4E:143:ARG:NH1	2.34	0.57
36:68:107:ARG:NH1	41:B8:36:GLU:HG2	2.20	0.57
34:69:1:MET:HE1	34:69:38:LEU:HD23	1.86	0.57
16:7A:22:THR:HA	16:7A:33:ILE:HD12	1.84	0.57
38:88:39:PRO:HA	38:88:97:VAL:O	2.05	0.57
26:1H:1188:U:H4'	43:D8:79:VAL:HG22	1.87	0.57
1:13:1130:A:H5'	9:8E:18:PHE:CE2	2.40	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:446:G:H1	1:13:488:C:H42	1.51	0.57
26:14:1997:G:OP2	62:14:3644:HOH:O	2.18	0.57
26:14:2275:C:H6	26:14:2275:C:H5'	1.70	0.57
27:16:40:U:H1'	27:16:45:A:N6	2.20	0.57
26:1H:223:A:O4'	26:1H:422:A:H5'	2.05	0.57
30:21:63:LEU:O	30:21:66:HIS:HB3	2.05	0.57
24:3L:4:U:H2'	24:3L:5:C:O4'	2.05	0.57
5:42:93:PRO:HG2	8:72:105:ARG:HG3	1.87	0.57
35:58:96:GLU:O	35:58:98:VAL:HG12	2.04	0.57
6:5E:86:ARG:O	6:5E:87:ARG:HG2	2.05	0.57
7:62:27:ILE:HD12	7:62:40:ALA:HA	1.86	0.57
15:6I:39:LEU:HD13	15:6I:56:LEU:HB2	1.87	0.57
44:A5:14:PRO:HG2	44:A5:78:GLU:HB2	1.85	0.57
1:13:201:C:H42	1:13:216:G:H1	1.53	0.57
26:14:827:U:H2'	26:14:2430:A:H2	1.70	0.57
1:1G:1:U:H4'	1:1G:630:G:N2	2.20	0.57
24:1L:3:G:N2	24:1L:4:U:O4	2.37	0.57
1:13:881:G:OP2	12:3I:12:ARG:NH2	2.37	0.57
7:6E:13:GLN:O	7:6E:24:THR:HG21	2.05	0.57
16:7A:14:ASN:OD1	16:7A:42:ARG:NH2	2.38	0.57
49:J8:86:SER:HB2	49:J8:87:PRO:HD2	1.87	0.57
49:J8:90:ILE:HG22	49:J8:94:LEU:HD22	1.87	0.57
1:13:316:G:OP2	1:13:351:G:O2'	2.22	0.56
1:13:769:G:N7	62:13:1852:HOH:O	2.32	0.56
26:14:2324:C:H5"	26:14:2325:G:H5'	1.85	0.56
26:14:395:U:O2'	26:14:396:G:C8	2.52	0.56
1:1G:1342:C:H4'	9:82:125:TYR:HB3	1.86	0.56
26:1H:1257:C:H4'	31:31:83:PHE:CD1	2.40	0.56
26:1H:2298:A:H62	26:1H:2318:G:H8	1.51	0.56
3:2E:16:ARG:NH2	3:2E:183:ASP:OD1	2.38	0.56
3:2E:40:ARG:O	3:2E:44:GLU:HG2	2.05	0.56
4:3E:122:ARG:HG2	4:3E:122:ARG:HH11	1.68	0.56
32:41:151:ALA:O	32:41:153:ARG:NH1	2.38	0.56
30:29:12:THR:HG21	41:75:11:GLU:OE1	2.04	0.56
41:B8:12:SER:CA	41:B8:14:TYR:H	2.19	0.56
41:B8:35:LYS:HE3	41:B8:38:ASN:HA	1.86	0.56
47:D5:19:ARG:NH1	47:D5:84:GLU:O	2.38	0.56
48:E5:34:GLY:HA2	48:E5:61:ALA:O	2.05	0.56
26:1H:1264:G:H5'	53:N8:11:THR:CG2	2.35	0.56
1:13:1003:G:H1	1:13:1037:C:H42	1.52	0.56
1:13:1071:C:H2'	1:13:1072:G:C8	2.40	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:683:G:N7	62:13:1853:HOH:O	2.33	0.56
26:14:2262:U:H4'	26:14:2328:A:C2	2.39	0.56
27:16:42:C:O3'	32:41:67:LYS:NZ	2.38	0.56
1:1G:1238:A:N3	1:1G:1241:G:O2'	2.27	0.56
1:1G:376:G:H1	1:1G:387:U:H3	1.53	0.56
33:59:6:ARG:HB2	33:59:65:HIS:CD2	2.39	0.56
6:5E:45:LEU:HD12	6:5E:59:TYR:HD2	1.69	0.56
17:8I:67:LYS:O	17:8I:68:ARG:HB3	2.05	0.56
26:14:71:A:C2	45:B5:31:HIS:NE2	2.73	0.56
45:F8:11:PRO:HB3	45:F8:92:LEU:HD21	1.87	0.56
48:I8:72:ARG:NH1	48:I8:75:LEU:HD12	2.20	0.56
26:1H:1500:G:O2'	29:11:100:GLY:O	2.18	0.56
26:14:122:G:OP1	26:14:149:A:O2'	2.21	0.56
26:14:1332:G:H21	26:14:1610:A:H8	1.54	0.56
26:14:1786:A:C2	26:14:2606:C:H1'	2.39	0.56
26:14:2331:G:O2'	48:E5:43:THR:HB	2.06	0.56
2:1E:60:ASP:O	2:1E:64:ARG:NE	2.38	0.56
1:1G:1478:C:H2'	1:1G:1479:C:H6	1.70	0.56
26:1H:1593:G:H2'	26:1H:1594:G:C8	2.40	0.56
26:1H:661:C:O2'	37:78:13:ASN:O	2.23	0.56
26:1H:880:G:H1	26:1H:897:C:H42	1.53	0.56
26:1H:991:C:H2'	26:1H:992:C:H6	1.70	0.56
26:1H:444:C:H4'	31:31:49:ALA:HB2	1.87	0.56
26:14:637:A:H2'	37:35:117:GLU:OE2	2.05	0.56
37:35:57:THR:HG22	37:35:60:MET:H	1.71	0.56
7:62:70:LYS:HG2	7:62:96:GLN:HB3	1.88	0.56
34:69:77:LEU:HA	34:69:141:LYS:HB3	1.86	0.56
34:69:81:VAL:H	34:69:143:SER:CB	2.15	0.56
15:6A:39:LEU:HD12	15:6A:56:LEU:HB2	1.87	0.56
28:71:64:LEU:HD21	28:71:188:ASN:HD21	1.69	0.56
39:98:86:ARG:NH2	39:98:118:GLU:HG2	2.16	0.56
19:AI:50:ALA:HB1	19:AI:57:HIS:HB3	1.87	0.56
29:11:33:LEU:O	29:11:64:ILE:HG23	2.04	0.56
2:12:22:LYS:HB3	2:12:40:HIS:HE1	1.70	0.56
27:16:66:A:N6	27:16:107:U:H2'	2.20	0.56
2:1E:12:GLU:HB3	2:1E:44:LEU:HD13	1.86	0.56
1:1G:1213:A:N6	1:1G:1215:G:N3	2.53	0.56
26:1H:1871:A:H2'	26:1H:1872:A:C8	2.40	0.56
26:1H:2584:U:H2'	26:1H:2585:U:H2'	1.87	0.56
11:2I:17:GLY:O	11:2I:80:VAL:HA	2.05	0.56
31:39:192:LEU:O	31:39:193:VAL:HG23	2.04	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:55:97:VAL:HA	39:55:113:LEU:O	2.05	0.56
41:75:108:ARG:HA	41:75:111:ARG:HG2	1.87	0.56
37:78:116:GLY:H	37:78:134:ALA:HB2	1.70	0.56
41:B8:99:LEU:HB3	41:B8:101:PHE:CE1	2.41	0.56
47:D5:29:TYR:CE2	47:D5:87:ASP:HB2	2.41	0.56
50:G5:4:SER:HB3	50:G5:7:ARG:HB2	1.88	0.56
26:1H:189:G:OP2	49:J8:39:LYS:HE3	2.05	0.56
1:13:177:C:OP2	20:BI:65:LYS:NZ	2.36	0.56
26:14:2158:A:H1'	26:14:2159:G:C8	2.40	0.56
26:14:403:U:H4'	26:14:404:C:H5'	1.86	0.56
26:14:491:G:H2'	26:14:492:A:H8	1.70	0.56
1:1G:1052:U:O2'	1:1G:1055:A:OP2	2.13	0.56
1:1G:971:G:OP1	1:1G:972:C:H5"	2.05	0.56
26:1H:1188:U:C4'	43:D8:79:VAL:HG22	2.35	0.56
26:1H:370:G:H4'	26:1H:371:A:OP2	2.04	0.56
27:1J:13:A:H5"	27:1J:15:A:C6	2.40	0.56
33:51:94:TYR:HA	33:51:106:THR:O	2.06	0.56
34:61:79:ILE:HB	34:61:142:VAL:HG12	1.88	0.56
1:1G:1291:G:O3'	9:82:39:GLY:HA3	2.06	0.56
9:82:19:LEU:HD11	9:82:84:ALA:HB1	1.86	0.56
18:9A:74:ARG:NH1	18:9A:81:PHE:HA	2.20	0.56
48:E5:32:ARG:O	48:E5:34:GLY:N	2.36	0.56
49:F5:62:VAL:HG21	49:F5:70:VAL:HG21	1.88	0.56
47:H8:163:LEU:HB3	47:H8:165:VAL:H	1.69	0.56
54:O8:41:PRO:HB2	54:O8:44:ARG:HH12	1.69	0.56
29:11:38:LYS:HG2	29:11:40:THR:CG2	2.36	0.56
2:12:132:LYS:HA	2:12:135:GLN:HB2	1.88	0.56
26:14:270(M):U:H5"	26:14:270(N):G:OP1	2.05	0.56
26:14:29:U:H2'	26:14:30:G:C8	2.39	0.56
10:1A:81:THR:O	10:1A:84:GLN:NE2	2.36	0.56
2:1E:60:ASP:HB3	2:1E:64:ARG:HH21	1.71	0.56
1:1G:991:U:H5	1:1G:1212:U:H1'	1.71	0.56
1:1G:604:G:H2'	1:1G:605:U:O4'	2.05	0.56
26:1H:2473:U:H2'	26:1H:2474:C:H5'	1.87	0.56
26:1H:65:C:H2'	26:1H:66:C:C6	2.41	0.56
11:2A:87:THR:O	11:2A:87:THR:OG1	2.21	0.56
32:41:47:LYS:HD2	32:41:81:LYS:HB2	1.86	0.56
5:42:100:VAL:HG23	5:42:118:ILE:HG22	1.87	0.56
41:75:45:PHE:CZ	41:75:74:ARG:HG3	2.40	0.56
37:78:59:LEU:O	56:Q8:13:ARG:HD2	2.05	0.56
37:78:78:PRO:HB3	37:78:111:ARG:NH2	2.20	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:82:51:ARG:HG2	9:82:56:LEU:HD13	1.87	0.56
39:98:46:GLY:HA2	39:98:49:ASP:HB2	1.88	0.56
40:A8:3:ARG:HG3	40:A8:4:LEU:HB2	1.88	0.56
20:BA:25:ARG:HG3	20:BA:29:LYS:HE3	1.88	0.56
52:M8:14:ILE:HG23	52:M8:21:VAL:HB	1.85	0.56
1:13:838:G:H1	1:13:848:C:H42	1.53	0.56
26:14:198:C:H5'	26:14:2244:U:OP1	2.05	0.56
26:14:2280:G:O2'	26:14:2388:A:N1	2.34	0.56
35:15:58:ASP:N	35:15:58:ASP:OD1	2.37	0.56
2:1E:114:ARG:O	2:1E:118:LEU:HB2	2.05	0.56
2:1E:71:VAL:HG23	2:1E:164:VAL:HA	1.88	0.56
1:1G:1148:U:O3'	9:82:14:VAL:HG11	2.06	0.56
26:1H:1412:A:H2'	26:1H:1413:G:H8	1.71	0.56
26:1H:1508:A:O2'	26:1H:1509:C:O5'	2.23	0.56
26:1H:646:A:H2'	26:1H:647:G:O4'	2.05	0.56
26:1H:754:C:H2'	26:1H:755:C:C6	2.41	0.56
26:1H:762:U:H4'	26:1H:763:G:O5'	2.06	0.56
31:31:183:VAL:O	31:31:187:VAL:HG23	2.05	0.56
26:1H:663:G:OP1	37:78:16:ARG:HB2	2.04	0.56
37:78:63:PRO:CB	56:Q8:30:ARG:HH21	2.09	0.56
26:1H:1800:C:OP2	29:11:183:ARG:NH2	2.38	0.56
2:12:72:GLY:HA3	2:12:81:VAL:HG21	1.87	0.56
1:13:1125:U:HO2'	1:13:1126:U:H6	1.50	0.56
1:13:838:G:O6	1:13:848:C:N4	2.38	0.56
26:14:1018:C:H2'	26:14:1019:U:H6	1.71	0.56
26:14:1693:U:O2'	29:19:14:ARG:NH2	2.39	0.56
26:14:2719:G:O6	62:14:3638:HOH:O	2.15	0.56
35:15:61:ARG:HH11	35:15:61:ARG:HA	1.71	0.56
27:16:2:C:O2	27:16:118:G:N2	2.19	0.56
26:1H:1520:U:H2'	26:1H:1521:G:O4'	2.06	0.56
4:32:108:LEU:HD21	4:32:183:GLY:HA3	1.88	0.56
4:32:18:LYS:HG2	59:32:302:SF4:S1	2.45	0.56
31:39:155:LEU:HD23	31:39:186:ILE:HD13	1.87	0.56
6:5E:78:GLU:O	6:5E:81:ILE:HG22	2.06	0.56
1:1G:376:G:H5"	16:7A:5:ARG:HD2	1.86	0.56
18:9A:71:LYS:HA	18:9A:74:ARG:HD2	1.88	0.56
1:13:963:G:H21	10:1I:55:LYS:CE	2.18	0.56
26:14:1582:C:HO2'	26:14:1586:A:H8	1.51	0.56
26:14:2037:G:H2'	26:14:2038:G:C8	2.40	0.56
26:14:2117:A:H2'	26:14:2118:U:H5	1.71	0.56
26:14:981:A:N1	26:14:2027:G:O2'	2.35	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1E:32:ILE:HD13	2:1E:40:HIS:HB3	1.88	0.56
1:1G:1343:G:H2'	1:1G:1344:C:H6	1.71	0.56
1:1G:323:U:O3'	20:BA:22:ARG:HD3	2.06	0.56
1:1G:973:G:H5"	1:1G:974:A:H5"	1.88	0.56
22:1K:76:A:H1'	26:1H:2583:G:H21	1.70	0.56
4:32:34:GLU:HB2	4:32:35:ARG:HH22	1.71	0.56
32:41:39:ILE:HD12	32:41:94:LEU:HD21	1.88	0.56
13:4I:23:TYR:HB3	13:4I:67:GLU:HB2	1.88	0.56
6:5E:22:GLU:OE1	6:5E:82:ARG:NH2	2.39	0.56
17:8A:99:SER:OG	17:8A:100:LYS:N	2.37	0.56
9:8E:34:ASN:O	9:8E:38:GLN:HB2	2.06	0.56
26:1H:2292:C:P	40:A8:17:ARG:HH22	2.29	0.56
47:D5:111:VAL:HG22	47:D5:112:ARG:HG2	1.88	0.56
47:H8:60:GLU:O	47:H8:61:LEU:HB3	2.05	0.56
1:13:1497:G:H2'	1:13:1498:U:H5'	1.87	0.56
26:14:2142:C:H2'	26:14:2143:C:C6	2.41	0.56
26:14:2418:A:OP2	56:M5:29:LYS:NZ	2.27	0.56
26:14:1775:U:O2'	61:14:3437:SPE:N1	2.39	0.56
26:14:646:A:H2'	26:14:647:G:O4'	2.06	0.56
1:1G:1251:A:O2'	1:1G:1369:C:O2'	2.24	0.56
1:1G:811:C:N4	62:1G:1884:HOH:O	2.38	0.56
26:1H:1221:C:H2'	26:1H:1222:C:H6	1.71	0.56
26:1H:1230:C:H2'	26:1H:1231:G:H8	1.70	0.56
26:1H:2688:U:C5	26:1H:2720:U:OP2	2.59	0.56
26:1H:2789:C:H1'	26:1H:2892:A:C2	2.41	0.56
26:1H:926:A:N7	62:1H:3826:HOH:O	2.33	0.56
31:39:89:VAL:HG12	31:39:90:PHE:H	1.71	0.56
32:49:27:ASN:HB3	32:49:30:GLU:HG3	1.86	0.56
26:14:2822:G:O6	39:55:2:ARG:HD3	2.06	0.56
8:72:51:VAL:HG11	8:72:60:ARG:HE	1.71	0.56
26:1H:625:G:N7	37:78:107:LYS:NZ	2.54	0.56
37:78:13:ASN:O	37:78:15:ARG:N	2.39	0.56
16:7I:4:ILE:HG12	16:7I:21:VAL:HG12	1.87	0.56
26:14:486:C:O2'	44:A5:60:ASN:ND2	2.39	0.56
40:A8:30:ARG:O	40:A8:30:ARG:HG3	2.05	0.56
26:1H:997:G:OP1	42:C8:93:LYS:HD2	2.05	0.56
51:L8:43:ILE:O	51:L8:47:VAL:HG23	2.06	0.56
56:M5:8:LYS:HB3	56:M5:12:LYS:HE3	1.88	0.56
54:O8:25:LYS:HB3	56:Q8:34:TRP:CD1	2.41	0.56
1:13:922:G:C6	1:13:923:A:C6	2.94	0.56
26:14:1416:G:HO2'	26:14:1417:C:H6	1.54	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2117:A:H2'	26:14:2118:U:C5	2.41	0.56
26:14:2327:A:H2'	26:14:2328:A:C8	2.41	0.56
1:1G:1259:C:N4	1:1G:1260:C:O2	2.39	0.56
1:1G:1469:G:H2'	1:1G:1470:G:H8	1.71	0.56
3:22:14:ILE:HG12	3:22:15:THR:H	1.71	0.56
3:22:32:LEU:HD22	3:22:59:ARG:HH22	1.71	0.56
38:45:20:ALA:HA	38:45:99:PRO:HG2	1.87	0.56
13:4A:70:LEU:O	13:4A:74:VAL:HG23	2.06	0.56
6:52:76:ALA:HB1	6:52:80:ARG:NH2	2.19	0.56
33:59:149:ARG:HA	33:59:162:ILE:HG21	1.88	0.56
14:5A:32:SER:O	14:5A:40:CYS:HA	2.06	0.56
16:7I:21:VAL:O	16:7I:33:ILE:N	2.36	0.56
44:E8:58:ALA:HB1	44:E8:64:MET:HB2	1.88	0.56
54:O8:43:CYS:HB3	54:O8:44:ARG:HH11	1.71	0.56
1:13:1060:C:HO2'	10:1I:56:HIS:HD1	1.54	0.55
1:13:592:G:H2'	1:13:593:G:H8	1.71	0.55
26:14:1581:G:H2'	26:14:1582:C:O4'	2.04	0.55
26:14:2275:C:H5'	26:14:2275:C:C6	2.41	0.55
27:16:94:C:H2'	27:16:95:U:C6	2.41	0.55
29:19:141:VAL:HG23	29:19:162:SER:HB2	1.88	0.55
29:19:70:TRP:C	29:19:70:TRP:CD1	2.80	0.55
1:1G:692:U:O2'	1:1G:694:A:N7	2.36	0.55
26:1H:1126:A:H4'	26:1H:1127:A:O5'	2.06	0.55
26:1H:2362:G:OP1	56:Q8:44:LYS:NZ	2.29	0.55
26:1H:271(B):G:N7	26:1H:421:U:H2'	2.20	0.55
23:2K:47:7MG:O2'	23:2K:48:U:OP2	2.21	0.55
4:32:163:GLU:HA	4:32:166:LYS:HE3	1.88	0.55
33:59:54:ARG:NH2	33:59:57:ASP:OD1	2.29	0.55
1:1G:878:G:H5'	8:72:89:PRO:HG2	1.87	0.55
1:13:1248:A:N3	9:8E:70:LYS:HE2	2.22	0.55
40:A8:111:GLU:HB2	40:A8:112:PHE:CE2	2.41	0.55
40:A8:27:SER:HA	40:A8:88:ASP:CB	2.36	0.55
47:D5:59:LEU:O	47:D5:60:GLU:HB3	2.05	0.55
43:D8:16:PRO:HA	43:D8:96:ILE:HG22	1.87	0.55
52:M8:24:THR:OG1	52:M8:25:TYR:N	2.38	0.55
2:12:166:ASP:OD2	2:12:169:LYS:HB2	2.05	0.55
2:12:71:VAL:HG21	2:12:164:VAL:HA	1.88	0.55
1:13:148:G:H2'	1:13:149:A:C8	2.41	0.55
1:13:179:A:H2'	1:13:180:U:C6	2.41	0.55
1:13:673:G:H2'	1:13:674:G:C8	2.41	0.55
26:14:1449(A):G:H2'	26:14:1450:C:C6	2.41	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2113:U:H3'	26:14:2114:A:H4'	1.88	0.55
26:14:2139:C:N3	26:14:2153:G:N2	2.54	0.55
2:1E:17:PHE:HB3	2:1E:44:LEU:HG	1.88	0.55
1:1G:316:G:OP2	1:1G:351:G:O2'	2.22	0.55
26:1H:1315:C:OP2	62:1H:3747:HOH:O	2.17	0.55
26:1H:247:G:H4'	26:1H:386:G:C5	2.41	0.55
27:1J:2:C:H2'	27:1J:3:C:H6	1.69	0.55
30:29:47:VAL:HG21	30:29:86:PRO:HD2	1.87	0.55
37:35:59:LEU:HD21	56:M5:10:ALA:HA	1.88	0.55
1:1G:527:G:O6	12:3A:49:ASN:ND2	2.37	0.55
13:4A:13:LYS:HD3	13:4A:14:ARG:H	1.70	0.55
1:13:339:C:OP2	36:68:97:ARG:NH1	2.39	0.55
41:75:91:ARG:NH1	41:75:124:ASP:OD2	2.30	0.55
37:78:82:GLY:HA2	37:78:113:LYS:O	2.06	0.55
41:B8:74:ARG:HD3	41:B8:76:PHE:CZ	2.41	0.55
47:D5:97:GLU:HB3	47:D5:125:LEU:HD11	1.87	0.55
26:1H:1266:G:O5'	44:E8:15:ARG:NH2	2.40	0.55
1:13:1028(B):C:H41	1:13:1032(A):G:H21	1.52	0.55
1:13:10:A:OP2	5:4E:126:ARG:HD3	2.06	0.55
1:13:1226:C:O3'	13:4I:111:LYS:NZ	2.39	0.55
1:13:1233:G:H2'	1:13:1234:C:C6	2.41	0.55
1:13:580:U:OP1	15:6I:54:ARG:NH2	2.39	0.55
26:14:1041:C:H42	26:14:1114:G:H22	1.54	0.55
26:14:323:G:O2'	26:14:1205:U:N3	2.31	0.55
1:1G:1177:G:O2'	1:1G:1178:G:O4'	2.24	0.55
1:1G:411:A:H62	1:1G:413:G:H21	1.53	0.55
26:1H:1827:C:C2'	26:1H:1828:G:H5'	2.36	0.55
38:45:110:THR:HG23	38:45:113:GLN:HB2	1.88	0.55
8:7E:119:LEU:HB3	8:7E:123:GLU:HG3	1.87	0.55
6:5E:94:GLN:NE2	18:9I:33:ASP:OD1	2.30	0.55
47:H8:77:ASP:N	47:H8:84:GLU:HG2	2.21	0.55
2:12:118:LEU:HD22	2:12:142:LEU:HB2	1.87	0.55
26:14:886:C:H1'	26:14:890:A:H2	1.70	0.55
1:1G:998:G:H22	1:1G:1043:C:N4	2.05	0.55
26:1H:2208:U:O2'	26:1H:2209:C:H5'	2.05	0.55
26:1H:34:C:O2'	26:1H:35:G:OP2	2.18	0.55
26:1H:7:G:H2'	26:1H:8:A:O4'	2.06	0.55
3:2E:7:PRO:O	3:2E:11:ARG:HG2	2.06	0.55
38:45:57:HIS:NE2	38:45:116:GLU:HG2	2.21	0.55
13:4I:4:ILE:HG23	13:4I:57:ARG:HA	1.89	0.55
33:59:18:GLU:HB2	33:59:25:LYS:HB2	1.89	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
18:9A:31:LEU:H	18:9A:31:LEU:HD23	1.71	0.55
41:B8:94:ALA:HB1	41:B8:99:LEU:HD21	1.89	0.55
26:1H:142:G:H1'	45:F8:37:THR:HG21	1.88	0.55
29:11:17:THR:CG2	29:11:204:ILE:HA	2.36	0.55
1:13:1286:A:H8	1:13:1287:A:H4'	1.71	0.55
26:14:2308:G:O2'	26:14:2309:A:OP1	2.20	0.55
26:14:802:A:H4'	62:14:4111:HOH:O	2.05	0.55
29:19:237:GLU:OE1	62:19:403:HOH:O	2.17	0.55
1:1G:1352:C:H2'	1:1G:1353:G:C8	2.41	0.55
26:1H:1586:A:H5"	26:1H:1586:A:N3	2.21	0.55
26:1H:2122:U:H2'	26:1H:2123:G:C8	2.41	0.55
26:1H:218:A:H2	26:1H:235:U:H4'	1.72	0.55
26:1H:479:A:N3	26:1H:481:G:H5"	2.22	0.55
3:2E:167:TRP:CD1	3:2E:168:ALA:N	2.75	0.55
4:32:31:CYS:C	4:32:33:MET:H	2.10	0.55
46:C5:20:TYR:CZ	46:C5:42:VAL:HA	2.42	0.55
46:C5:76:CYS:SG	46:C5:97:ARG:HG3	2.45	0.55
29:11:75:ILE:HG21	29:11:99:ASP:OD2	2.07	0.55
2:12:160:ASP:N	2:12:160:ASP:OD1	2.40	0.55
1:13:652:U:O4	1:13:752:G:O2'	2.17	0.55
1:13:688:G:H2'	1:13:689:C:H6	1.72	0.55
26:14:1359:A:H5'	26:14:1359:A:H8	1.70	0.55
26:14:1939:U:OP1	26:14:2604:U:O2'	2.20	0.55
26:14:2068:U:H3	26:14:2430:A:H2	1.55	0.55
26:14:2123:G:N2	26:14:2176:A:N1	2.54	0.55
1:1G:1308:U:H5"	13:4A:98:VAL:HG22	1.89	0.55
1:1G:1512:U:H2'	1:1G:1513:A:C8	2.42	0.55
1:1G:540:G:H2'	1:1G:541:G:O4'	2.06	0.55
26:1H:500:G:N7	62:1H:3827:HOH:O	2.33	0.55
11:2I:122:LYS:HE3	11:2I:124:LYS:HE3	1.89	0.55
34:69:109:ILE:HB	34:69:130:TYR:CZ	2.42	0.55
1:13:376:G:H5"	16:7I:5:ARG:HB2	1.89	0.55
46:C5:83:THR:HG22	46:C5:84:ARG:H	1.70	0.55
52:M8:4:GLY:C	52:M8:5:ILE:HG13	2.27	0.55
26:14:1166:C:O2'	62:14:3646:HOH:O	2.18	0.55
26:14:2646:C:H2'	26:14:2647:U:O4'	2.07	0.55
10:1I:61:GLU:OE1	14:5I:58:LYS:NZ	2.27	0.55
3:22:112:SER:HB3	3:22:115:LEU:HB2	1.88	0.55
4:32:31:CYS:HB2	4:32:33:MET:H	1.72	0.55
37:35:106:LEU:HD13	37:35:112:LEU:HD23	1.89	0.55
4:3E:61:LYS:NZ	4:3E:72:GLU:OE2	2.24	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:41:124:SER:HB2	32:41:131:TYR:CE2	2.42	0.55
35:58:40:PRO:O	42:C8:64:ARG:HG2	2.07	0.55
17:8I:18:THR:OG1	17:8I:69:LYS:NZ	2.26	0.55
42:C8:95:LEU:HD22	43:D8:4:ILE:HD13	1.89	0.55
45:B5:11:PRO:HD3	50:G5:37:PHE:CD2	2.42	0.55
1:13:134:A:H61	16:7I:25:ARG:NH1	2.05	0.55
26:14:127:A:H5"	26:14:128:C:C6	2.42	0.55
26:14:2118:U:O2'	26:14:2145:C:N3	2.39	0.55
26:14:2406:U:H2'	26:14:2406:U:OP2	2.07	0.55
26:14:270(E):G:H2'	26:14:270(F):U:C6	2.42	0.55
26:14:774:A:O2'	26:14:775:G:O5'	2.23	0.55
26:14:848:G:H2'	26:14:849:A:H8	1.68	0.55
26:1H:919:G:H4'	27:16:81:G:H4'	1.88	0.55
1:1G:1065:U:C5	1:1G:1190:G:H1'	2.42	0.55
1:1G:1466:C:H2'	1:1G:1467:G:O4'	2.06	0.55
1:1G:1499:A:H1'	1:1G:1520:G:H5'	1.89	0.55
1:1G:45:U:H2'	1:1G:46:G:C8	2.42	0.55
26:1H:1021:A:H8	26:1H:1021:A:H3'	1.71	0.55
26:1H:1354:A:H2'	26:1H:1355:G:O4'	2.07	0.55
3:2E:15:THR:HG22	3:2E:16:ARG:N	2.22	0.55
31:39:178:PRO:HG2	31:39:179:GLU:OE1	2.07	0.55
5:42:71:LEU:HD21	5:42:115:VAL:HG22	1.89	0.55
36:68:7:TYR:CZ	36:68:44:LYS:HG3	2.42	0.55
45:B5:63:LYS:O	45:B5:63:LYS:NZ	2.29	0.55
20:BA:41:ILE:HD13	20:BA:87:LYS:HD2	1.89	0.55
46:C5:8:LYS:HZ3	46:C5:95:LYS:HD3	1.70	0.55
47:D5:170:THR:O	47:D5:172:ALA:N	2.39	0.55
50:K8:28:LYS:HE3	50:K8:56:GLN:NE2	2.22	0.55
26:1H:2228:G:OP2	29:11:263:ARG:NH2	2.40	0.55
1:13:157:G:H2'	1:13:158:G:H8	1.72	0.55
1:13:535:A:H5"	62:13:1845:HOH:O	2.06	0.55
26:14:1417:C:H42	26:14:1581:G:H1	1.55	0.55
26:14:2542:A:H5"	26:14:2542:A:N3	2.22	0.55
1:1G:1181:G:C2	1:1G:1182:G:H1'	2.42	0.55
1:1G:1396:A:H4'	1:1G:1397:C:H5"	1.88	0.55
30:21:128:SER:OG	30:21:129:HIS:N	2.40	0.55
3:22:36:ASP:HA	3:22:39:ILE:HD12	1.88	0.55
3:2E:16:ARG:HH22	3:2E:183:ASP:HA	1.72	0.55
4:3E:127:THR:HG22	4:3E:147:ALA:HB3	1.89	0.55
5:42:78:HIS:CD2	8:72:107:LEU:HD22	2.42	0.55
32:49:120:LEU:HB2	32:49:180:PHE:CD1	2.41	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:55:72:ASP:O	39:55:76:VAL:HG23	2.07	0.55
34:69:76:THR:HG23	34:69:139:GLN:O	2.06	0.55
26:1H:870:A:OP1	38:88:6:ARG:NH2	2.39	0.55
19:AA:13:ASP:O	19:AA:16:LEU:HB3	2.07	0.55
43:D8:36:PRO:O	43:D8:38:LEU:N	2.40	0.55
29:11:72:LYS:HE2	29:11:101:GLU:OE2	2.06	0.55
26:14:1486:A:H2'	26:14:1487:G:C8	2.42	0.55
26:14:1810:A:H2'	26:14:1811:G:O4'	2.07	0.55
26:14:2270:G:OP2	62:14:3648:HOH:O	2.18	0.55
26:14:2629:A:N3	26:14:2629:A:H2'	2.21	0.55
26:14:2693:A:H2'	26:14:2694:G:C8	2.43	0.55
1:1G:736:C:H2'	1:1G:737:A:C8	2.42	0.55
1:1G:978:A:H5'	1:1G:979:C:OP2	2.07	0.55
26:1H:1021:A:C8	26:1H:1021:A:H3'	2.42	0.55
26:1H:1639:U:C2'	26:1H:1640:C:H5"	2.37	0.55
26:1H:524:U:H2'	26:1H:525:U:C6	2.43	0.55
30:29:68:ALA:O	30:29:70:ALA:N	2.40	0.55
4:32:4:TYR:CE2	4:32:11:LEU:HD11	2.42	0.55
1:1G:619:U:N3	4:32:134:ASP:OD1	2.38	0.55
25:4L:21:A:C2'	25:4L:22:A:H5"	2.35	0.55
39:98:56:LYS:NZ	39:98:90:ARG:O	2.38	0.55
1:13:1182:G:H4'	1:13:1183:A:H5"	1.89	0.54
26:14:2849:U:H4'	26:14:2868:A:C2	2.42	0.54
2:1E:115:LEU:HD13	2:1E:145:LEU:HB3	1.89	0.54
1:1G:1413:A:H2'	1:1G:1414:U:O4'	2.07	0.54
26:1H:1165:U:H2'	26:1H:1166:C:H6	1.72	0.54
26:1H:1339:G:H21	26:1H:1603:A:H1'	1.72	0.54
22:1K:72:C:H2'	22:1K:73:A:H5"	1.89	0.54
23:2L:54:G:H2'	23:2L:55:5MU:H6	1.71	0.54
12:3A:6:THR:OG1	12:3A:9:GLN:HG3	2.07	0.54
26:14:873:G:O3'	38:45:63:LYS:NZ	2.39	0.54
35:58:17:ASP:O	35:58:56:ASN:HB2	2.07	0.54
14:5A:29:ARG:NH2	14:5A:40:CYS:SG	2.80	0.54
7:6E:22:LEU:HD23	7:6E:62:PHE:CE2	2.42	0.54
42:85:98:LEU:HB2	42:85:102:GLU:HB2	1.89	0.54
43:D8:14:VAL:HB	43:D8:96:ILE:HG13	1.89	0.54
45:F8:41:ASN:O	45:F8:45:THR:HG23	2.07	0.54
26:14:528:A:C2	26:14:2042:A:H2'	2.42	0.54
26:14:586:A:N1	26:14:809:G:O2'	2.32	0.54
26:14:843:G:H1	26:14:935:C:H42	1.54	0.54
2:1E:234:PRO:HB2	2:1E:236:TYR:H	1.71	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:942:G:C2	1:1G:1342:C:C2	2.95	0.54
1:1G:596:C:H3'	62:1G:1801:HOH:O	2.07	0.54
26:1H:1179:C:H2'	26:1H:1180:C:C6	2.41	0.54
26:1H:1590:U:H2'	26:1H:1591:G:C8	2.42	0.54
26:1H:300:A:N6	62:1H:3860:HOH:O	2.39	0.54
3:22:8:ILE:O	3:22:11:ARG:N	2.39	0.54
30:29:60:ASN:OD1	30:29:63:LEU:HD22	2.06	0.54
3:2E:18:TRP:H	3:2E:18:TRP:HE3	1.55	0.54
4:3E:155:LEU:O	4:3E:157:LEU:N	2.40	0.54
36:68:2:ILE:HG13	36:68:8:LEU:HD11	1.90	0.54
41:75:3:ARG:HA	41:75:6:LEU:HB2	1.88	0.54
8:7E:85:ARG:HD3	8:7E:88:LYS:HG2	1.89	0.54
20:BI:10:LEU:HD11	20:BI:12:ALA:HB3	1.88	0.54
47:D5:60:GLU:HG2	47:D5:60:GLU:O	2.06	0.54
50:G5:32:LEU:HD21	50:G5:54:LYS:HG2	1.90	0.54
46:G8:94:LYS:HZ2	46:G8:95:LYS:H	1.53	0.54
1:13:396:G:O2'	1:13:398:C:OP1	2.08	0.54
26:14:754:C:H2'	26:14:755:C:C6	2.42	0.54
1:1G:1028(A):C:N4	1:1G:1032(B):G:H1	2.05	0.54
26:1H:1062:G:H1'	26:1H:1088:A:C5	2.42	0.54
26:1H:1441:G:H2'	26:1H:1442:G:C8	2.42	0.54
10:1I:85:LEU:HB2	10:1I:86:MET:SD	2.47	0.54
27:1J:3:C:H2'	27:1J:4:C:C6	2.42	0.54
26:1H:2572:A:C8	30:21:144:ARG:HD3	2.42	0.54
3:2E:119:ARG:O	3:2E:123:GLN:HG3	2.07	0.54
4:32:28:SER:HB3	4:32:29:PRO:HA	1.89	0.54
31:39:122:LYS:HB3	31:39:191:ARG:HB2	1.89	0.54
13:4I:40:ASN:HB3	13:4I:43:THR:HG23	1.88	0.54
33:51:10:PRO:O	33:51:11:VAL:HG13	2.08	0.54
34:61:4:ILE:HD11	34:61:44:LEU:HD13	1.89	0.54
7:62:113:GLU:O	7:62:119:ARG:HD3	2.07	0.54
26:14:2295:C:OP1	40:65:10:ARG:NH1	2.40	0.54
37:78:63:PRO:HG2	56:Q8:25:MET:HB2	1.89	0.54
17:8I:81:ARG:HB3	17:8I:83:ASP:OD1	2.08	0.54
47:D5:87:ASP:N	47:D5:87:ASP:OD1	2.41	0.54
48:I8:36:ILE:HD13	48:I8:36:ILE:O	2.07	0.54
49:J8:95:LEU:HD12	49:J8:96:LYS:HD2	1.89	0.54
1:13:1414:U:H2'	1:13:1415:G:H8	1.73	0.54
1:13:165:C:H2'	1:13:166:G:C8	2.38	0.54
1:13:224:C:H2'	1:13:225:C:C6	2.42	0.54
1:13:540:G:H2'	1:13:541:G:O4'	2.07	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:588:G:H1	1:1G:651:C:H42	1.56	0.54
26:1H:1050:A:H2'	26:1H:1051:G:O4'	2.07	0.54
26:1H:1697:G:OP2	26:1H:1698:A:O2'	2.18	0.54
26:1H:2000:G:HO2'	26:1H:2689:U:H5	1.52	0.54
26:1H:2256:G:H4'	48:I8:9:SER:HB2	1.90	0.54
26:1H:2817:G:OP1	39:98:99:LYS:NZ	2.32	0.54
26:14:2724:C:OP1	30:29:118:LYS:HE3	2.07	0.54
38:45:38:GLU:HG3	38:45:127:ILE:HG22	1.88	0.54
33:51:56:SER:OG	33:51:57:ASP:N	2.39	0.54
7:6E:89:MET:HE1	7:6E:155:ARG:HD2	1.89	0.54
55:L5:5:TRP:NE1	55:L5:7:PRO:HG3	2.22	0.54
1:13:31:G:O2'	1:13:48:C:N4	2.41	0.54
26:14:2162:G:H2'	26:14:2163:C:H5'	1.90	0.54
26:14:2726:U:H4'	36:25:1:MET:HE3	1.90	0.54
26:1H:918:A:O2'	27:16:96:G:N2	2.41	0.54
1:1G:115:G:H1'	1:1G:116:A:N7	2.23	0.54
1:1G:421:U:O2'	1:1G:423:G:N7	2.39	0.54
1:1G:555:C:H2'	1:1G:556:C:C6	2.42	0.54
1:1G:628:G:H2'	1:1G:629:G:C8	2.43	0.54
26:1H:1332:G:N2	26:1H:1610:A:C8	2.76	0.54
26:1H:768:G:O2'	26:1H:1379:A:N6	2.40	0.54
30:29:12:THR:O	30:29:23:VAL:HG22	2.08	0.54
31:39:3:GLU:O	31:39:19:GLU:HB2	2.07	0.54
4:3E:156:GLU:O	4:3E:160:GLN:HB3	2.08	0.54
24:3K:22:G:N7	24:3K:46:G:N1	2.55	0.54
38:45:19:GLY:O	38:45:98:LYS:HB3	2.08	0.54
15:6A:39:LEU:HD12	15:6A:56:LEU:HD13	1.88	0.54
28:71:7:TYR:CE1	28:71:220:PRO:HB3	2.43	0.54
28:71:214:VAL:HG23	28:71:224:ILE:HG12	1.89	0.54
8:7E:8:ASP:O	8:7E:12:ARG:HG3	2.07	0.54
9:8E:5:TYR:CE1	9:8E:16:ARG:HG2	2.43	0.54
48:E5:18:ALA:HB3	48:E5:20:ARG:HH21	1.73	0.54
48:E5:25:ARG:HD2	48:E5:29:GLN:NE2	2.22	0.54
44:E8:37:ARG:HD3	44:E8:38:TYR:CE2	2.42	0.54
29:11:70:TRP:CD1	29:11:70:TRP:C	2.81	0.54
1:13:1133:G:N2	1:13:1141:C:O2	2.41	0.54
1:13:182:U:H5	1:13:183:G:C4	2.26	0.54
26:14:2318:G:H5'	26:14:2319:G:OP2	2.08	0.54
26:14:252:G:OP2	37:35:50:ARG:NH2	2.41	0.54
26:14:531:C:H4'	26:14:532:A:H5"	1.89	0.54
1:1G:192:U:H2'	1:1G:193:C:H6	1.72	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:983:A:H2	1:1G:984:C:C6	2.25	0.54
26:1H:2315:G:H2'	26:1H:2316:C:C6	2.42	0.54
3:22:113:ALA:HA	3:22:202:ILE:HD11	1.88	0.54
1:13:35:G:O2'	12:3I:118:SER:O	2.18	0.54
13:4I:5:ALA:HB2	13:4I:61:GLU:HG2	1.90	0.54
39:55:106:GLY:O	39:55:107:ASP:HB3	2.07	0.54
35:58:22:THR:OG1	35:58:23:LEU:N	2.41	0.54
35:58:96:GLU:HB2	35:58:122:VAL:HG12	1.90	0.54
26:14:270(L):U:O2	34:69:50:ARG:HD3	2.07	0.54
26:1H:533:G:H5'	42:C8:24:TYR:CD1	2.43	0.54
47:H8:93:ASP:HB3	47:H8:131:ARG:HH21	1.73	0.54
1:13:401:C:OP2	4:3E:73:ARG:HD3	2.08	0.54
1:13:686:U:O4	1:13:703:G:H1'	2.07	0.54
1:1G:1024:G:OP1	1:1G:1024:G:H4'	2.08	0.54
1:1G:1431:C:H42	1:1G:1469:G:H1	1.55	0.54
1:1G:250:A:H4'	1:1G:251:G:O5'	2.08	0.54
26:1H:2017:U:O2	53:N8:10:LYS:HB2	2.07	0.54
26:1H:2518:A:H8	26:1H:2518:A:H5'	1.73	0.54
26:1H:536:A:H2'	26:1H:537:C:C6	2.43	0.54
27:1J:4:C:H42	27:1J:116:G:H1	1.55	0.54
30:21:105:THR:HG21	30:21:164:ARG:NH1	2.22	0.54
26:14:673:C:H5"	31:39:81:PRO:HD2	1.90	0.54
43:95:17:GLY:H	43:95:96:ILE:HB	1.71	0.54
20:BA:86:ARG:HH11	20:BA:86:ARG:HB2	1.72	0.54
46:C5:75:ILE:HG13	46:C5:80:GLY:HA2	1.89	0.54
26:14:1434:A:H61	26:14:1558:A:H62	1.56	0.54
26:14:2233:U:H2'	26:14:2234:G:C8	2.42	0.54
26:14:278:A:OP2	26:14:278:A:H2'	2.08	0.54
26:14:363(E):U:H5'	26:14:363(F):A:OP2	2.07	0.54
26:14:955:C:OP1	38:45:87:LYS:HE2	2.07	0.54
26:1H:2507:C:OP1	62:1H:3774:HOH:O	2.18	0.54
23:2K:24:C:H2'	23:2K:25:U:C6	2.42	0.54
4:32:104:VAL:O	4:32:108:LEU:HB2	2.08	0.54
4:32:173:TRP:CZ3	4:32:193:ASP:HB3	2.43	0.54
24:3K:72:C:H2'	24:3K:73:A:H5"	1.90	0.54
42:85:92:ARG:C	42:85:94:ASN:H	2.10	0.54
52:M8:14:ILE:HG22	52:M8:24:THR:HG22	1.89	0.54
1:13:631:G:O2'	1:13:632:A:O5'	2.21	0.54
26:14:1946:U:H2'	26:14:1947:C:C6	2.43	0.54
26:14:67:U:H2'	26:14:68:G:C8	2.43	0.54
26:14:811:U:O2'	37:35:21:ARG:HG3	2.08	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:15:47:ALA:HB2	35:15:112:LEU:HD21	1.89	0.54
27:16:15:A:H3'	27:16:16:G:H5'	1.88	0.54
2:1E:100:GLY:O	2:1E:104:ASN:N	2.36	0.54
26:1H:1290:C:H2'	26:1H:1291:C:C6	2.43	0.54
26:1H:1420:U:HO2'	26:1H:1421:G:P	2.31	0.54
26:1H:364:C:H3'	62:1H:3873:HOH:O	2.07	0.54
26:1H:569:U:C4	26:1H:570:G:C6	2.96	0.54
26:1H:593:G:C1'	56:Q8:4:MET:HE1	2.38	0.54
30:29:4:ILE:HD13	30:29:28:ALA:HB1	1.89	0.54
13:4A:13:LYS:HD3	13:4A:14:ARG:N	2.23	0.54
35:58:42:TRP:HA	35:58:48:MET:SD	2.48	0.54
26:1H:2820:A:O5'	39:98:4:LEU:HD23	2.07	0.54
26:1H:2232:U:P	49:J8:40:ARG:HH12	2.31	0.54
1:13:244:U:H4'	1:13:245:C:O5'	2.08	0.54
1:13:648:A:N6	1:13:649:G:O6	2.40	0.54
10:1A:11:PHE:CE1	10:1A:67:THR:HG22	2.43	0.54
1:1G:1139:G:H22	1:1G:1143:G:H1	1.55	0.54
1:1G:719:C:O2'	18:9A:49:LYS:HB3	2.08	0.54
26:1H:107:C:H2'	26:1H:108:U:H6	1.73	0.54
26:1H:1259:G:H2'	26:1H:1260:G:C8	2.43	0.54
26:1H:1386:C:H2'	26:1H:1387:C:C6	2.41	0.54
26:1H:1533:C:H3'	26:1H:1534:G:H5"	1.90	0.54
26:1H:259:G:N2	26:1H:621:A:H8	2.05	0.54
10:1I:54:PHE:CZ	10:1I:55:LYS:NZ	2.68	0.54
3:2E:114:PRO:O	3:2E:118:GLN:HG3	2.08	0.54
13:4A:91:ARG:HB2	13:4A:98:VAL:HG12	1.89	0.54
33:59:27:LYS:HD3	33:59:32:GLU:HG3	1.90	0.54
7:62:59:LEU:HD21	7:62:63:LYS:HZ2	1.72	0.54
1:1G:1178:G:H5"	9:82:93:ARG:NH2	2.23	0.54
9:8E:8:GLY:HA3	9:8E:79:LEU:HB3	1.90	0.54
41:B8:58:ASN:O	41:B8:58:ASN:ND2	2.40	0.54
47:D5:128:VAL:HG22	47:D5:129:SER:H	1.73	0.54
26:1H:142:G:H1'	45:F8:37:THR:CG2	2.38	0.54
1:13:1149:C:H2'	1:13:1150:U:H6	1.72	0.53
1:13:271:C:H2'	1:13:272:C:H6	1.73	0.53
1:13:2:U:H5"	1:13:630:G:H21	1.73	0.53
1:13:843:U:H3'	1:13:848:C:C6	2.43	0.53
26:14:579:G:H2'	26:14:580:C:C6	2.43	0.53
26:14:754:C:H2'	26:14:755:C:H6	1.73	0.53
27:16:73:A:C4	27:16:104:A:C2	2.96	0.53
27:16:80:U:H2'	27:16:81:G:N2	2.20	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1019:U:HO2'	26:1H:1021:A:H2	1.56	0.53
26:1H:2518:A:C8	26:1H:2518:A:H5'	2.42	0.53
26:1H:2053:G:H5'	30:21:144:ARG:O	2.08	0.53
26:1H:2572:A:N7	30:21:145:LYS:HB2	2.22	0.53
36:25:63:VAL:HB	36:25:102:VAL:HG12	1.89	0.53
30:29:51:PHE:CG	30:29:52:LEU:N	2.74	0.53
11:2I:72:ALA:HB1	11:2I:77:MET:HE3	1.88	0.53
40:65:23:ARG:HB2	40:65:86:ALA:HB2	1.89	0.53
37:78:63:PRO:HB2	56:Q8:30:ARG:NH2	2.06	0.53
41:B8:50:ILE:HD11	41:B8:102:ILE:HG13	1.88	0.53
49:F5:32:LYS:O	62:F5:101:HOH:O	2.18	0.53
53:N8:41:PRO:O	53:N8:44:THR:OG1	2.24	0.53
37:78:59:LEU:HD11	56:Q8:10:ALA:HA	1.90	0.53
29:11:84:TYR:CE1	29:11:86:PRO:HB3	2.36	0.53
1:13:1126:U:O2	1:13:1280:A:C8	2.62	0.53
26:14:1332:G:N2	26:14:1609:A:O2'	2.42	0.53
26:14:273(C):C:N4	26:14:363(C):G:H1	2.07	0.53
26:14:67:U:H2'	26:14:68:G:H8	1.73	0.53
26:14:1569:A:O2'	29:19:37:LEU:HD23	2.09	0.53
2:1E:11:LEU:HG	2:1E:213:LEU:HD13	1.91	0.53
1:1G:1494:G:N7	58:1G:1702:PAR:N32	2.56	0.53
26:1H:1203:G:H3'	26:1H:1204:A:H5"	1.91	0.53
26:1H:1359:A:C2	26:1H:1372:U:O4	2.61	0.53
26:1H:2788:C:O2'	26:1H:2809:A:N3	2.41	0.53
26:1H:315:G:H2'	26:1H:316:C:C6	2.42	0.53
27:1J:44:G:H1'	27:1J:47:C:N4	2.23	0.53
3:2E:15:THR:CG2	3:2E:181:ASN:HA	2.38	0.53
31:39:146:ALA:HB3	31:39:148:LEU:HG	1.89	0.53
24:3L:35:U:O2	24:3L:36:U:N3	2.41	0.53
32:41:97:ASP:O	32:41:100:TRP:N	2.42	0.53
38:45:57:HIS:ND1	38:45:117:ALA:HB2	2.23	0.53
13:4A:29:ARG:HD3	13:4A:64:TRP:CE2	2.44	0.53
7:62:69:VAL:HG12	7:62:103:TRP:HE3	1.72	0.53
34:69:124:GLY:O	34:69:142:VAL:HG23	2.07	0.53
47:D5:10:ARG:HH21	47:D5:26:GLY:N	2.06	0.53
47:H8:98:MET:O	47:H8:125:LEU:HA	2.08	0.53
50:K8:15:LYS:HZ2	50:K8:15:LYS:N	2.03	0.53
55:P8:12:ARG:NH2	55:P8:44:PRO:HB3	2.23	0.53
2:12:103:THR:HG23	2:12:176:GLU:HB3	1.90	0.53
1:13:1157:A:H61	1:13:1178:G:H21	1.57	0.53
1:13:157:G:H2'	1:13:158:G:C8	2.44	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:536:C:H2'	1:13:537:G:C8	2.42	0.53
1:13:5:U:O2'	1:13:6:G:O5'	2.26	0.53
1:13:920:U:H2'	1:13:921:U:C6	2.42	0.53
26:14:1871:A:H2'	26:14:1872:A:C8	2.42	0.53
26:14:2125:G:H21	26:14:2173:A:N6	2.07	0.53
26:14:235:U:H2'	26:14:236:C:C6	2.44	0.53
26:14:602:G:OP2	26:14:602:G:H8	1.91	0.53
1:1G:1306:A:N6	1:1G:1331:G:O2'	2.41	0.53
26:1H:1486:A:H2'	26:1H:1487:G:C8	2.42	0.53
26:1H:2125:G:H21	26:1H:2173:A:H62	1.57	0.53
30:29:37:ARG:HD2	30:29:44:TYR:OH	2.08	0.53
31:31:198:ALA:O	31:31:201:VAL:N	2.41	0.53
24:3K:35:U:H2'	24:3K:36:U:C6	2.43	0.53
5:42:51:VAL:O	5:42:55:VAL:HG23	2.07	0.53
13:4A:19:LEU:HB3	13:4A:25:ILE:HG21	1.90	0.53
33:51:4:ILE:HG23	33:51:6:ARG:NH2	2.23	0.53
2:1E:178:ARG:HG3	8:7E:72:PRO:HA	1.91	0.53
46:G8:43:ASN:O	46:G8:64:GLU:HA	2.08	0.53
49:J8:77:ALA:HA	49:J8:78:LYS:C	2.28	0.53
29:11:238:GLY:HA2	62:11:311:HOH:O	2.08	0.53
1:13:976:G:N2	1:13:1362(A):C:OP2	2.39	0.53
1:13:345:C:H4'	1:13:346:G:N7	2.23	0.53
26:14:1520:U:H2'	26:14:1521:G:O4'	2.09	0.53
26:14:307:G:N2	26:14:309:G:H3'	2.24	0.53
2:1E:54:THR:HG23	2:1E:185:ILE:HD11	1.90	0.53
1:1G:1228:C:H2'	1:1G:1229:A:H8	1.73	0.53
1:1G:1316:G:H2'	1:1G:1317:C:H5"	1.89	0.53
26:1H:1088:A:H5'	26:1H:1089:G:H5'	1.90	0.53
26:1H:2235:G:H2'	26:1H:2236:C:C6	2.44	0.53
26:1H:2306:C:H3'	26:1H:2307:G:H5'	1.91	0.53
26:1H:2308:G:N1	26:1H:2311:A:H2	1.94	0.53
26:1H:2573:C:H3'	62:1H:3962:HOH:O	2.07	0.53
26:1H:2756:U:H4'	26:1H:2757:A:OP1	2.07	0.53
30:21:101:ARG:CZ	30:21:171:GLU:HB2	2.39	0.53
26:1H:2786:U:O2'	30:21:62:PRO:O	2.20	0.53
26:14:1665:A:H4'	36:25:67:LYS:HB2	1.91	0.53
30:29:101:ARG:CZ	30:29:171:GLU:HB2	2.38	0.53
30:29:31:CYS:SG	30:29:51:PHE:HB2	2.48	0.53
11:2I:21:ILE:HG12	11:2I:30:VAL:HG12	1.89	0.53
12:3A:60:LEU:HD23	12:3A:64:TYR:HB3	1.91	0.53
12:3I:8:ASN:O	12:3I:11:VAL:HG23	2.09	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:5E:96:PRO:HB3	18:9I:30:ASP:OD2	2.08	0.53
14:5I:3:ARG:HB2	14:5I:3:ARG:HH11	1.72	0.53
8:72:103:VAL:HG21	8:72:110:ALA:HB2	1.91	0.53
1:1G:1344:C:H5"	9:82:120:ARG:O	2.09	0.53
38:88:32:TYR:OH	38:88:111:GLU:OE1	2.19	0.53
38:88:37:LEU:HD21	38:88:130:LYS:HE3	1.89	0.53
45:B5:50:LYS:HB2	45:B5:87:GLN:NE2	2.24	0.53
1:13:491:G:H2'	1:13:492:G:O4'	2.08	0.53
1:13:703:G:H8	1:13:703:G:O5'	1.90	0.53
26:14:2849:U:OP1	41:75:95:ARG:NH1	2.42	0.53
26:14:94:G:N3	50:G5:47:ASN:ND2	2.56	0.53
1:1G:1169:A:C6	1:1G:1170:A:C6	2.97	0.53
1:1G:1167:A:H2'	1:1G:1169:A:O4'	2.09	0.53
1:1G:1238:A:H62	1:1G:1301:U:H3	1.57	0.53
1:1G:750:G:N3	15:6A:23:GLY:HA3	2.23	0.53
1:1G:828:A:H2'	1:1G:829:G:O4'	2.08	0.53
1:1G:8:A:C6	4:32:209:ARG:HB2	2.44	0.53
26:1H:2001:A:H2'	26:1H:2002:G:C8	2.44	0.53
3:22:175:LEU:H	3:22:175:LEU:HD12	1.74	0.53
40:65:67:ARG:NH1	40:65:67:ARG:HB2	2.23	0.53
8:72:20:TYR:HA	8:72:65:TYR:CZ	2.43	0.53
41:75:88:ILE:HD11	41:75:91:ARG:HG2	1.89	0.53
16:7A:21:VAL:HG22	16:7A:33:ILE:HB	1.91	0.53
2:12:95:GLN:HB2	2:12:148:TYR:HA	1.90	0.53
2:12:97:TRP:HZ3	2:12:99:GLY:HA2	1.74	0.53
1:13:1376:U:H2'	1:13:1377:A:C8	2.43	0.53
1:13:1402:C:H2'	1:13:1403:C:O4'	2.07	0.53
1:13:1478:C:H2'	1:13:1479:C:C6	2.44	0.53
1:13:417:C:H2'	1:13:418:C:H6	1.73	0.53
1:13:542:G:OP1	4:3E:10:ARG:NH2	2.42	0.53
1:13:690:G:H2'	1:13:691:G:O4'	2.08	0.53
1:13:976:G:OP1	14:5I:32:SER:N	2.41	0.53
26:14:1203:G:H3'	26:14:1204:A:H5"	1.90	0.53
26:14:1321:A:H2'	26:14:1322:A:O4'	2.09	0.53
26:14:2129:C:H3'	26:14:2130:U:C6	2.43	0.53
26:14:2396:G:H4'	49:F5:30:VAL:H	1.74	0.53
26:14:39:C:H2'	26:14:40:C:C6	2.44	0.53
26:14:636:G:O2'	26:14:638:G:O2'	2.19	0.53
1:1G:751:U:H4'	15:6A:24:SER:HA	1.89	0.53
33:51:83:TYR:HB3	33:51:135:GLY:H	1.74	0.53
33:59:167:GLU:HG3	33:59:169:VAL:HG23	1.90	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:5A:17:LYS:HZ3	14:5A:18:VAL:HG13	1.74	0.53
9:82:71:SER:HA	9:82:74:ILE:HD12	1.90	0.53
42:85:110:VAL:HG12	42:85:114:LYS:HD3	1.90	0.53
9:8E:21:PRO:HA	9:8E:59:PHE:HD1	1.73	0.53
27:16:8:U:O3'	40:A8:25:ARG:NH2	2.41	0.53
27:16:44:G:H1'	27:16:47:C:N4	2.24	0.53
29:19:77:ALA:HB2	29:19:97:TYR:CD2	2.44	0.53
10:1A:40:LEU:HG	10:1A:41:PRO:HD2	1.90	0.53
26:1H:2287:A:N1	26:1H:2346:A:H2	2.07	0.53
26:1H:2780:G:OP1	35:58:118:LYS:HE2	2.09	0.53
10:1I:84:GLN:HB3	10:1I:88:LEU:HD23	1.90	0.53
39:55:82:GLU:H	39:55:85:PRO:HG2	1.73	0.53
27:1J:7:G:H4'	40:65:29:PHE:HD2	1.70	0.53
36:68:34:THR:OG1	36:68:35:VAL:N	2.42	0.53
7:6E:102:ARG:O	7:6E:106:GLN:HG3	2.08	0.53
41:75:132:LYS:HB3	41:75:133:GLU:OE1	2.08	0.53
16:7I:5:ARG:HE	16:7I:22:THR:HG21	1.73	0.53
26:1H:910:A:H62	38:88:12:GLN:HA	1.74	0.53
9:8E:46:ALA:HB2	9:8E:74:ILE:HG23	1.90	0.53
20:BI:71:THR:HG22	20:BI:72:LEU:N	2.21	0.53
47:H8:116:VAL:H	47:H8:146:ILE:HG12	1.74	0.53
1:13:1000:A:H2'	1:13:1001:G:H8	1.74	0.53
35:15:30:ILE:HG22	35:15:34:LEU:HD22	1.90	0.53
1:1G:422:C:HO2'	1:1G:423:G:N2	2.06	0.53
1:1G:438:G:H4'	4:32:123:HIS:ND1	2.24	0.53
1:1G:777:A:C2	11:2A:119:CYS:HB3	2.44	0.53
26:1H:1899:G:H22	26:1H:1902:C:H41	1.56	0.53
26:1H:1968:G:OP1	62:1H:3775:HOH:O	2.19	0.53
26:1H:528:A:N1	26:1H:2042:A:H2'	2.23	0.53
27:1J:94:C:H2'	27:1J:95:U:C6	2.43	0.53
11:2A:54:ARG:NH2	24:3L:40:C:OP1	2.41	0.53
37:35:105:LEU:O	37:35:106:LEU:HB3	2.09	0.53
13:4I:34:LEU:HD13	13:4I:41:PRO:HA	1.90	0.53
16:7A:19:ILE:HB	16:7A:36:ILE:O	2.09	0.53
9:8E:86:VAL:O	9:8E:90:PRO:HB3	2.09	0.53
26:14:495:G:H21	44:A5:61:ASN:HD21	1.57	0.53
41:B8:29:ARG:HB2	41:B8:46:GLU:HG3	1.91	0.53
52:M8:42:PHE:CG	52:M8:42:PHE:O	2.61	0.53
1:13:1256:A:O2'	1:13:1257:U:O5'	2.25	0.53
26:14:363:G:H2'	26:14:363(A):A:H8	1.74	0.53
1:1G:345:C:H5'	1:1G:346:G:C5	2.44	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:998(A):C:H2'	1:1G:999:U:C6	2.43	0.53
4:32:12:CYS:SG	4:32:18:LYS:HA	2.49	0.53
37:35:85:LEU:HG	37:35:120:ALA:HA	1.91	0.53
45:B5:89:ILE:O	45:B5:93:GLU:HG2	2.09	0.53
1:13:1190:G:H5"	3:2E:176:HIS:CE1	2.44	0.53
1:13:376:G:H5"	16:7I:5:ARG:HD2	1.90	0.53
26:14:2291:U:H2'	26:14:2292:C:C6	2.44	0.53
26:14:399:G:OP2	62:14:3647:HOH:O	2.18	0.53
2:1E:54:THR:HG21	2:1E:201:ILE:HD11	1.90	0.53
1:1G:867:G:O2'	1:1G:868:C:H5'	2.09	0.53
26:1H:2108:C:H2'	26:1H:2109:U:O4'	2.09	0.53
26:1H:2314:C:H2'	26:1H:2315:G:C8	2.41	0.53
26:1H:950:G:H2'	26:1H:951:C:C6	2.44	0.53
26:1H:95:G:O2'	50:K8:48:HIS:HB3	2.08	0.53
30:21:143:ASN:HD22	30:21:147:PRO:HD2	1.74	0.53
26:14:2823:A:OP1	30:29:113:PHE:HB2	2.09	0.53
4:3E:161:ASN:O	4:3E:165:MET:HB2	2.09	0.53
14:5A:37:PHE:CE1	14:5A:53:LEU:HD13	2.44	0.53
7:62:16:LEU:HD11	9:82:45:ALA:HB2	1.91	0.53
8:7E:129:VAL:HG23	8:7E:130:GLY:H	1.74	0.53
17:8A:88:TYR:CZ	17:8A:92:ARG:HD2	2.44	0.53
44:A5:60:ASN:OD1	44:A5:60:ASN:N	2.41	0.53
29:11:126:GLN:HG2	29:11:127:VAL:N	2.24	0.52
1:13:1396:A:H4'	1:13:1397:C:H5"	1.91	0.52
1:13:411:A:N9	1:13:413:G:H1'	2.24	0.52
26:14:1652:A:OP1	39:55:8:ARG:NH1	2.42	0.52
26:14:1678:G:N2	26:14:1989:G:N2	2.56	0.52
1:1G:1343:G:H2'	1:1G:1344:C:C6	2.44	0.52
1:1G:629:G:H2'	1:1G:630:G:O4'	2.09	0.52
26:1H:1332:G:N2	26:1H:1609:A:HO2'	2.07	0.52
26:1H:1378:A:O2'	62:1H:3767:HOH:O	2.15	0.52
26:1H:2656:U:N3	26:1H:2665:A:H2	2.04	0.52
31:31:107:LYS:HE2	31:31:207:GLY:N	2.24	0.52
32:41:97:ASP:H	32:41:100:TRP:HD1	1.56	0.52
5:4E:10:MET:HA	5:4E:32:VAL:HA	1.91	0.52
33:59:171:LEU:HD13	33:59:172:LYS:H	1.74	0.52
41:75:88:ILE:HG13	41:75:88:ILE:O	2.08	0.52
40:A8:10:ARG:O	40:A8:14:VAL:HG13	2.08	0.52
47:H8:103:ARG:HD3	47:H8:136:PHE:CD1	2.44	0.52
47:H8:61:LEU:O	47:H8:64:GLY:HA2	2.09	0.52
56:Q8:46:ARG:HB2	56:Q8:47:LYS:HB2	1.89	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:429:U:H3'	4:3E:9:CYS:SG	2.49	0.52
1:13:757:U:H2'	1:13:758:G:O4'	2.08	0.52
26:14:273(F):C:H3'	26:14:274:G:C5'	2.39	0.52
27:16:60:C:C2	27:16:61:G:C8	2.97	0.52
1:1G:1062:U:H2'	1:1G:1063:C:C6	2.44	0.52
1:1G:1132:C:H2'	1:1G:1133:G:C8	2.43	0.52
1:1G:256:U:H2'	1:1G:257:G:C8	2.45	0.52
1:1G:736:C:OP1	18:9A:72:ARG:NH2	2.43	0.52
26:1H:2689:U:H6	26:1H:2689:U:H5'	1.74	0.52
26:1H:600:G:N2	26:1H:605:C:O3'	2.42	0.52
3:22:11:ARG:NH2	3:22:182:ILE:HD11	2.24	0.52
31:39:133:ASN:HA	31:39:162:LEU:HD23	1.91	0.52
36:68:4:PRO:HA	36:68:21:CYS:O	2.09	0.52
34:69:14:ASP:O	34:69:17:GLN:HB2	2.09	0.52
34:69:72:LEU:HD21	34:69:107:VAL:HG11	1.90	0.52
34:69:77:LEU:HG	34:69:141:LYS:HE2	1.91	0.52
41:75:3:ARG:N	41:75:4:GLY:O	2.42	0.52
9:8E:45:ALA:O	9:8E:78:LYS:NZ	2.35	0.52
42:85:92:ARG:CZ	43:95:11:GLN:H	2.22	0.52
39:98:32:GLY:HA2	39:98:116:LEU:CD1	2.39	0.52
46:G8:55:TYR:CZ	46:G8:61:ILE:HD11	2.44	0.52
56:Q8:52:LYS:N	56:Q8:53:PRO:HD2	2.23	0.52
1:13:313:A:H2'	1:13:314:C:H6	1.75	0.52
26:14:1239:G:H2'	26:14:1240:U:O4'	2.09	0.52
26:14:2689:U:P	26:14:2719:G:H22	2.32	0.52
26:14:2777:G:OP2	26:14:2781:A:O2'	2.20	0.52
26:14:662:G:H5"	37:35:16:ARG:HG2	1.91	0.52
26:14:7:G:H2'	26:14:8:A:C8	2.45	0.52
1:1G:934:C:O2'	1:1G:1344:C:OP2	2.26	0.52
1:1G:41:G:H2'	1:1G:42:G:H8	1.74	0.52
26:1H:389:G:H1	37:78:71:VAL:HG12	1.74	0.52
24:1L:9:A:N6	24:1L:23:A:N7	2.57	0.52
23:2K:16:C:O2'	23:2K:62:C:OP1	2.25	0.52
24:3L:9:A:H2'	24:3L:11:C:N4	2.24	0.52
1:13:1423:G:OP1	36:68:49:ARG:NH2	2.42	0.52
37:78:118:GLY:O	37:78:137:LYS:NZ	2.42	0.52
17:8I:88:TYR:CD1	17:8I:89:LEU:HD23	2.45	0.52
43:D8:65:GLY:HA3	43:D8:91:TYR:CE2	2.45	0.52
47:H8:113:ALA:N	47:H8:114:GLY:HA2	2.24	0.52
29:11:146:GLU:HB2	29:11:189:CYS:HB3	1.91	0.52
26:14:1133:U:O2	26:14:1137:G:H5"	2.10	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1416:G:O2'	26:14:1417:C:O5'	2.25	0.52
26:14:2129:C:H5"	26:14:2130:U:C5	2.44	0.52
29:19:133:LEU:HB3	29:19:173:VAL:HG11	1.92	0.52
2:1E:162:ILE:O	2:1E:185:ILE:HG23	2.09	0.52
2:1E:93:VAL:HG11	2:1E:97:TRP:HD1	1.75	0.52
1:1G:1142:G:H3'	1:1G:1143:G:C8	2.45	0.52
26:1H:2065:C:H2'	26:1H:2066:C:C6	2.44	0.52
26:1H:2105:C:H2'	26:1H:2106:G:C8	2.41	0.52
26:1H:760:G:H2'	26:1H:761:A:O4'	2.09	0.52
1:13:963:G:N2	10:1I:55:LYS:NZ	2.57	0.52
26:1H:2680:C:H5'	30:21:189:PRO:HA	1.92	0.52
3:22:43:LEU:HD23	3:22:47:LEU:HB2	1.90	0.52
4:32:31:CYS:H	4:32:35:ARG:NH1	2.06	0.52
9:8E:17:VAL:HA	9:8E:63:ILE:HG12	1.90	0.52
47:D5:99:TYR:HD2	47:D5:123:ASP:HB3	1.74	0.52
29:11:79:VAL:O	29:11:113:VAL:HG23	2.09	0.52
1:13:1190:G:H5"	3:2E:176:HIS:NE2	2.23	0.52
1:13:130:A:O2'	1:13:131:C:O5'	2.24	0.52
26:14:2018:G:P	53:J5:9:LYS:HZ1	2.32	0.52
26:14:450:G:N7	62:14:3706:HOH:O	2.34	0.52
26:14:823:G:H2'	26:14:824:A:C8	2.44	0.52
1:1G:1289:A:OP1	21:1B:9:ARG:NH2	2.43	0.52
21:1F:10:ARG:HH11	21:1F:10:ARG:HB3	1.74	0.52
1:1G:134:A:H61	16:7A:25:ARG:NH1	2.07	0.52
1:1G:67:C:H2'	1:1G:68:G:C8	2.45	0.52
1:1G:892:A:O2'	1:1G:1415:G:H4'	2.09	0.52
26:1H:1449(A):G:H2'	26:1H:1450:C:H6	1.74	0.52
4:32:70:ILE:HD11	4:32:75:PHE:HD1	1.75	0.52
13:4A:31:LYS:O	13:4A:35:GLU:HG2	2.09	0.52
25:4K:13:A:N6	25:4K:14:A:N7	2.57	0.52
28:71:27:HIS:HA	28:71:182:PRO:HB3	1.90	0.52
5:4E:148:VAL:HG21	8:7E:107:LEU:HG	1.91	0.52
16:7I:3:LYS:HG2	16:7I:24:ALA:HB2	1.90	0.52
42:85:98:LEU:HA	42:85:100:VAL:O	2.10	0.52
17:8I:45:HIS:HB3	17:8I:72:ARG:HG2	1.89	0.52
40:A8:99:LYS:O	40:A8:103:GLU:HG2	2.09	0.52
42:C8:102:GLU:OE1	43:D8:13:ARG:NH2	2.42	0.52
47:D5:5:LEU:HG	47:D5:47:VAL:HG21	1.92	0.52
44:E8:24:ILE:HD12	44:E8:24:ILE:O	2.10	0.52
49:J8:53:VAL:HB	49:J8:58:ILE:HD13	1.92	0.52
52:M8:15:ILE:HB	52:M8:32:TYR:CD1	2.45	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1015:A:H2'	1:13:1016:A:H8	1.74	0.52
1:13:1126:U:H2'	1:13:1127:G:C5'	2.40	0.52
1:13:539:A:H2'	1:13:540:G:C8	2.44	0.52
1:13:827:U:C5	1:13:872:A:N1	2.77	0.52
26:14:1190:G:H2'	26:14:1191:G:H8	1.75	0.52
26:14:1568:G:OP1	29:19:63:ARG:NH1	2.38	0.52
26:14:1757:U:N3	26:14:1762:A:H2	2.02	0.52
26:14:2282:G:H4'	26:14:2389:G:O2'	2.09	0.52
26:14:2820:A:P	39:55:2:ARG:HH12	2.33	0.52
29:19:35:LYS:HG2	29:19:35:LYS:O	2.10	0.52
29:19:44:ASN:CB	29:19:45:ASN:HA	2.31	0.52
1:1G:411:A:H62	1:1G:413:G:N2	2.07	0.52
1:1G:815:A:N7	1:1G:1509:C:O2'	2.36	0.52
26:1H:1107:G:H2'	26:1H:1108:U:C6	2.45	0.52
26:1H:1314:C:OP1	62:1H:3747:HOH:O	2.19	0.52
26:1H:2068:U:N3	26:1H:2430:A:H2	2.08	0.52
26:1H:2836:U:H2'	26:1H:2837:G:C8	2.45	0.52
26:1H:515:A:H1'	26:1H:581:C:H1'	1.92	0.52
26:1H:945:A:H4'	62:1H:3603:HOH:O	2.09	0.52
3:22:150:LYS:HE3	3:22:152:ILE:HG13	1.91	0.52
26:1H:443:A:N7	31:31:45:ARG:HG2	2.24	0.52
1:1G:411:A:OP1	4:32:30:LYS:NZ	2.43	0.52
1:1G:429:U:H3'	4:32:9:CYS:SG	2.50	0.52
31:39:124:LEU:HG	31:39:126:VAL:HG12	1.91	0.52
31:39:129:PHE:HA	31:39:142:TRP:CD1	2.45	0.52
32:41:114:ILE:HG22	32:41:115:ARG:O	2.10	0.52
38:45:137:TYR:HD1	38:45:137:TYR:C	2.13	0.52
1:1G:1059:C:O3'	14:5A:45:ARG:NH2	2.43	0.52
37:78:121:LYS:O	37:78:123:LEU:N	2.42	0.52
56:M5:22:VAL:O	56:M5:50:LEU:HB3	2.10	0.52
52:M8:42:PHE:CD2	52:M8:42:PHE:O	2.63	0.52
1:13:1315:U:H2'	1:13:1316:G:O4'	2.10	0.52
1:13:1533:C:O2'	1:13:1534:A:OP1	2.23	0.52
26:14:1641:A:H2'	26:14:1642:G:O4'	2.10	0.52
26:14:2105:C:H2'	26:14:2106:G:O4'	2.09	0.52
26:14:691:C:O4'	29:19:43:ARG:NH2	2.43	0.52
1:1G:554:C:H2'	1:1G:555:C:H6	1.75	0.52
1:1G:583:A:H2'	1:1G:584:G:O4'	2.10	0.52
26:1H:1794:U:H2'	26:1H:1795:C:H6	1.75	0.52
26:1H:2131:G:H5'	26:1H:2132:U:H3'	1.92	0.52
26:1H:2389:G:H5"	26:1H:2390:U:O4'	2.10	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:589:C:H2'	26:1H:590:A:C8	2.45	0.52
26:1H:658:C:H2'	26:1H:659:C:C6	2.45	0.52
26:1H:729:G:O5'	29:11:208:LYS:NZ	2.42	0.52
30:21:59:VAL:HG13	30:21:60:ASN:H	1.75	0.52
3:2E:175:LEU:HD21	3:2E:201:TYR:CE2	2.45	0.52
11:2I:79:SER:HB2	11:2I:106:LYS:HD2	1.90	0.52
4:32:61:LYS:NZ	4:32:72:GLU:OE2	2.31	0.52
31:39:130:ALA:H	31:39:142:TRP:HD1	1.58	0.52
33:51:43:VAL:HB	33:51:52:VAL:HG22	1.91	0.52
33:51:33:LEU:HD12	33:51:75:ALA:HA	1.91	0.52
28:71:218:MET:N	28:71:218:MET:SD	2.82	0.52
16:7I:4:ILE:HB	16:7I:66:PRO:HB3	1.91	0.52
39:98:117:VAL:O	39:98:118:GLU:HB2	2.09	0.52
18:9I:52:PRO:O	18:9I:56:THR:HG23	2.09	0.52
40:A8:106:ARG:HA	40:A8:110:LEU:H	1.75	0.52
20:BI:46:GLU:HB2	20:BI:48:LYS:HG3	1.91	0.52
47:D5:53:ILE:HG22	47:D5:71:VAL:HG13	1.90	0.52
50:K8:3:LEU:H	50:K8:4:SER:C	2.13	0.52
50:K8:47:ASN:O	50:K8:49:LYS:HG3	2.10	0.52
2:12:127:ILE:HA	2:12:130:ARG:CZ	2.39	0.52
1:13:1194:U:H2'	1:13:1195:C:C6	2.45	0.52
1:13:1240:U:OP1	7:6E:119:ARG:NH2	2.41	0.52
1:13:129(A):G:C2	1:13:188:U:O2'	2.63	0.52
1:13:17:U:H2'	1:13:18:C:C6	2.45	0.52
26:14:1108:U:H5"	26:14:1109:C:OP2	2.09	0.52
26:14:2329:G:H2'	26:14:2330:G:C8	2.44	0.52
1:1G:113:G:O4'	1:1G:354:G:H4'	2.10	0.52
1:1G:26:A:N6	1:1G:558:G:O2'	2.41	0.52
1:1G:973:G:H1'	10:1A:55:LYS:HG2	1.92	0.52
26:1H:1359:A:H2'	26:1H:1360:A:H5'	1.92	0.52
26:1H:1050:A:C8	26:1H:2751:G:N7	2.78	0.52
27:1J:17:C:H2'	27:1J:18:G:O4'	2.09	0.52
24:3K:8:U:H5'	24:3K:48:C:O2'	2.09	0.52
7:62:115:ARG:O	7:62:118:VAL:HG22	2.09	0.52
1:1G:192:U:C4'	20:BA:103:GLY:HA2	2.40	0.52
47:D5:94:GLU:HB3	47:D5:96:VAL:HG23	1.91	0.52
43:D8:59:ALA:HB2	43:D8:96:ILE:HD13	1.92	0.52
56:M5:14:VAL:CG1	56:M5:22:VAL:HG13	2.39	0.52
54:O8:15:GLU:HA	54:O8:49:HIS:HA	1.92	0.52
26:14:185:U:H4'	26:14:218:A:H4'	1.92	0.52
26:14:827:U:H2'	26:14:2430:A:C2	2.44	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2689:U:OP2	26:14:2719:G:N2	2.42	0.52
26:14:395:U:O2'	26:14:396:G:N7	2.32	0.52
1:1G:1008:C:H42	1:1G:1021:G:H22	1.57	0.52
1:1G:1324:A:H2'	1:1G:1325:C:H6	1.75	0.52
1:1G:1423:G:H2'	1:1G:1424:C:H6	1.75	0.52
1:1G:198:G:H2'	1:1G:199:G:H8	1.75	0.52
1:1G:1:U:O5'	1:1G:630:G:N2	2.42	0.52
1:1G:690:G:H2'	1:1G:691:G:O4'	2.10	0.52
1:1G:742:G:OP2	15:6A:35:ARG:NH2	2.34	0.52
26:1H:2389:G:H5"	26:1H:2390:U:H5'	1.91	0.52
26:1H:2563:U:H1'	26:1H:2566:A:N6	2.25	0.52
30:21:51:PHE:CE2	30:21:52:LEU:HD23	2.44	0.52
36:25:22:ILE:HG22	36:25:40:VAL:HB	1.92	0.52
31:31:130:ALA:H	31:31:132:VAL:HG13	1.73	0.52
32:41:56:ALA:HB2	32:41:153:ARG:HE	1.75	0.52
5:42:33:VAL:HG21	5:42:109:ILE:HG12	1.92	0.52
5:4E:75:THR:OG1	5:4E:76:ILE:N	2.42	0.52
7:62:126:ASP:HB3	7:62:131:LYS:O	2.09	0.52
41:75:23:ARG:HG3	41:75:120:ARG:NH1	2.25	0.52
40:A8:106:ARG:CZ	40:A8:107:GLU:HG2	2.39	0.52
47:D5:15:PRO:HA	47:D5:18:LEU:HB2	1.92	0.52
46:G8:39:VAL:O	46:G8:42:VAL:HG22	2.09	0.52
46:G8:28:LYS:NZ	46:G8:64:GLU:OE2	2.30	0.52
53:N8:31:VAL:HG23	53:N8:42:PRO:HG3	1.92	0.52
26:1H:1797:C:O2'	29:11:259:THR:OG1	2.23	0.52
1:13:1258:G:H2'	1:13:1259:C:C6	2.45	0.52
1:1G:1055:A:N7	1:1G:1200:C:N4	2.58	0.52
1:1G:652:U:H1'	1:1G:653:A:H2	1.74	0.52
26:1H:1025:G:C4	26:1H:1135:C:H1'	2.45	0.52
26:1H:1636:C:H2'	26:1H:1637:A:C8	2.45	0.52
26:1H:1675:C:N3	30:21:128:SER:OG	2.42	0.52
26:1H:459:U:H2'	26:1H:460:A:H8	1.75	0.52
12:3I:53:ARG:HB3	12:3I:69:TYR:HE1	1.75	0.52
38:45:90:VAL:O	38:45:91:GLU:HB2	2.10	0.52
33:51:164:TYR:HB2	33:51:167:GLU:HB2	1.92	0.52
7:6E:45:ASP:O	7:6E:49:ILE:HG13	2.10	0.52
40:A8:105:ALA:O	40:A8:109:GLY:HA3	2.10	0.52
50:K8:28:LYS:HB3	50:K8:53:LEU:CD2	2.40	0.52
52:M8:39:CYS:SG	52:M8:41:PRO:HD2	2.50	0.52
1:13:191(C):G:H2'	1:13:191(D):U:O4'	2.10	0.51
1:13:191(D):U:H2'	1:13:191(E):G:C8	2.44	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1154:G:OP1	42:85:58:ARG:HD3	2.11	0.51
26:14:528:A:O2'	26:14:529:A:H5'	2.09	0.51
29:19:72:LYS:HB3	29:19:75:ILE:HD12	1.92	0.51
1:1G:1105:A:H2'	1:1G:1106:G:H8	1.75	0.51
1:1G:1512:U:H2'	1:1G:1513:A:H8	1.75	0.51
26:1H:155:C:H5'	26:1H:161:U:OP2	2.10	0.51
26:1H:2747:G:O6	26:1H:2755:C:H5"	2.10	0.51
24:1L:59:A:H5"	24:1L:60:U:C5	2.45	0.51
36:25:10:VAL:HG13	36:25:17:ARG:O	2.10	0.51
3:2E:64:VAL:HG12	3:2E:66:VAL:HG23	1.91	0.51
31:31:191:ARG:HB3	31:31:191:ARG:HH11	1.74	0.51
24:3K:56:C:H1'	26:1H:2169:A:H62	1.73	0.51
13:4I:8:GLU:O	13:4I:10:PRO:HD3	2.09	0.51
33:59:107:VAL:HG11	33:59:152:ARG:HG2	1.93	0.51
7:62:26:PHE:CE2	7:62:30:ILE:HD11	2.45	0.51
1:13:1298:C:P	7:6E:114:ARG:HH22	2.33	0.51
8:7E:86:ILE:HG13	8:7E:133:LEU:HD22	1.92	0.51
38:88:52:VAL:O	38:88:56:ARG:HB2	2.10	0.51
40:A8:36:TYR:HB3	40:A8:52:SER:HB3	1.92	0.51
56:Q8:51:ALA:HB1	56:Q8:52:LYS:HA	1.92	0.51
1:13:827:U:H5	1:13:872:A:N1	2.07	0.51
1:13:890:G:O2'	1:13:906:G:O6	2.24	0.51
26:14:1758:G:C2	26:14:2696:U:H5'	2.45	0.51
26:14:2010:G:O6	62:14:3650:HOH:O	2.19	0.51
24:3L:76:A:O2'	26:14:2394:C:N3	2.40	0.51
2:1E:28:PHE:CE2	2:1E:190:THR:HA	2.44	0.51
1:1G:108:G:H5'	1:1G:109:A:H5"	1.92	0.51
1:1G:998(A):C:O2	1:1G:1042:G:N2	2.33	0.51
26:1H:1174:A:H1'	26:1H:1178:C:N4	2.23	0.51
26:1H:1658:C:H2'	26:1H:1659:U:C6	2.44	0.51
26:1H:1728:G:O6	26:1H:1730:U:H5"	2.10	0.51
26:1H:306:U:H2'	26:1H:307:G:O4'	2.10	0.51
26:1H:35:G:H2'	26:1H:36:G:O4'	2.10	0.51
26:1H:50:U:H3'	26:1H:51:G:H5'	1.90	0.51
24:1L:52:G:N2	24:1L:62:C:O2	2.31	0.51
4:3E:112:VAL:HG12	4:3E:116:GLN:OE1	2.10	0.51
32:49:66:GLN:HE21	32:49:94:LEU:HD23	1.74	0.51
1:1G:1535:C:H41	25:4L:10:G:N2	2.05	0.51
14:5I:6:LEU:HB3	14:5I:23:ARG:NH2	2.25	0.51
15:6I:18:PHE:CZ	15:6I:21:ASP:HB2	2.45	0.51
1:1G:1371:G:OP1	9:82:11:LYS:HG2	2.11	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
17:8A:67:LYS:HA	17:8A:70:ARG:HH12	1.75	0.51
46:G8:87:LYS:HD2	46:G8:89:PHE:HD2	1.75	0.51
1:13:128:G:O2'	17:8I:3:LYS:NZ	2.40	0.51
26:14:1028:A:H2'	26:14:1029:A:C8	2.46	0.51
26:14:111:A:H4'	50:G5:69:ARG:HH22	1.76	0.51
26:14:2238:G:N7	62:14:3707:HOH:O	2.34	0.51
26:14:1759:A:H4'	26:14:2715:C:O4'	2.11	0.51
26:14:1012:U:C5	35:15:28:THR:HG21	2.44	0.51
26:1H:1259:G:H2'	26:1H:1260:G:H8	1.74	0.51
26:1H:1678:G:H22	26:1H:1989:G:N2	2.00	0.51
26:1H:2111:C:N3	26:1H:2118:U:O2'	2.39	0.51
26:1H:2321:G:H5"	26:1H:2322:A:OP2	2.09	0.51
26:1H:270(K):C:H1'	26:1H:270(N):G:H1	1.75	0.51
26:1H:818:G:H4'	26:1H:838:C:O3'	2.11	0.51
27:1J:93:C:H2'	27:1J:94:C:H6	1.75	0.51
30:21:68:ALA:HB1	30:21:70:ALA:O	2.11	0.51
30:29:52:LEU:O	30:29:74:PRO:HB2	2.10	0.51
3:2E:142:MET:SD	3:2E:148:GLY:HA2	2.51	0.51
4:3E:50:ARG:HD3	4:3E:51:PRO:HD2	1.91	0.51
6:5E:70:ASP:N	6:5E:70:ASP:OD1	2.41	0.51
9:82:112:LYS:HA	9:82:119:ALA:CB	2.34	0.51
9:82:26:VAL:HG13	9:82:61:ALA:O	2.10	0.51
20:BI:50:GLU:HG2	20:BI:100:ILE:HB	1.92	0.51
48:I8:63:VAL:HG23	48:I8:64:ASP:O	2.10	0.51
51:L8:4:LEU:HD11	51:L8:39:ASP:HA	1.92	0.51
1:13:1331:G:OP2	13:4I:23:TYR:HD1	1.92	0.51
1:13:51:A:OP2	1:13:52:G:H8	1.94	0.51
26:14:1442:G:H2'	26:14:1443:G:C8	2.45	0.51
26:14:2152:G:N3	26:14:2152:G:H2'	2.25	0.51
26:14:2542:A:H4'	26:14:2542:A:OP1	2.09	0.51
26:14:2572:A:C8	30:29:144:ARG:HD2	2.45	0.51
26:14:2651:C:H42	26:14:2669:G:H1	1.57	0.51
26:14:307:G:H21	26:14:330:A:H62	1.59	0.51
26:14:443:A:H1'	26:14:1201:C:O4'	2.10	0.51
26:14:768:G:O2'	26:14:1379:A:N6	2.43	0.51
29:19:35:LYS:HB2	29:19:62:TYR:O	2.10	0.51
2:1E:163:PHE:HA	2:1E:185:ILE:O	2.11	0.51
26:1H:1614:A:H8	26:1H:1614:A:P	2.33	0.51
26:1H:97:C:H5"	50:K8:2:LYS:HB2	1.92	0.51
27:1J:13:A:H5"	27:1J:15:A:N6	2.26	0.51
26:1H:2052:G:H4'	30:21:143:ASN:O	2.11	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:2E:113:ALA:N	3:2E:183:ASP:OD2	2.41	0.51
31:39:3:GLU:N	31:39:3:GLU:OE1	2.44	0.51
33:51:83:TYR:HB2	33:51:134:SER:HA	1.93	0.51
33:59:109:PHE:CZ	33:59:152:ARG:HD3	2.45	0.51
40:65:14:VAL:HG21	40:65:89:ARG:HD3	1.92	0.51
15:6I:67:LEU:O	15:6I:71:GLN:HB2	2.11	0.51
26:1H:2132:U:H3	28:71:5:LYS:HB2	1.76	0.51
37:78:64:LYS:HD2	56:Q8:12:LYS:HB3	1.92	0.51
9:8E:53:VAL:HG21	9:8E:85:LEU:HD22	1.92	0.51
6:52:96:PRO:HB3	18:9A:30:ASP:CG	2.31	0.51
45:B5:1:MET:N	50:G5:29:LYS:HE3	2.24	0.51
55:P8:15:THR:HG22	55:P8:16:HIS:CE1	2.46	0.51
26:1H:1568:G:H5"	29:11:61:LEU:CD2	2.41	0.51
1:13:1412:C:H2'	1:13:1413:A:C8	2.46	0.51
1:13:266:G:H5"	1:13:267:C:C5	2.46	0.51
1:13:77:C:H2'	1:13:78:G:H8	1.74	0.51
1:13:8:A:H62	4:3E:208:SER:HB2	1.75	0.51
26:14:1425:G:H2'	26:14:1426:G:C8	2.46	0.51
26:14:2513:G:N2	30:29:143:ASN:HD21	2.08	0.51
26:14:2579:C:H4'	30:29:134:ILE:HG12	1.93	0.51
26:14:2711:A:OP2	62:14:3507:HOH:O	2.19	0.51
26:14:994:C:OP2	42:85:54:LYS:NZ	2.34	0.51
27:16:88:C:H2'	27:16:89:G:O4'	2.10	0.51
26:1H:1799:G:O2'	26:1H:1800:C:OP2	2.19	0.51
26:1H:1919:A:H5"	26:1H:1920:C:OP2	2.10	0.51
30:29:54:GLN:HG2	30:29:55:ASN:N	2.25	0.51
24:3L:76:A:H8	26:14:2394:C:N4	2.08	0.51
6:52:33:TYR:CE1	6:52:78:GLU:HG3	2.46	0.51
6:5E:97:PHE:HD1	18:9I:31:LEU:HD11	1.75	0.51
7:6E:15:ASP:HB3	7:6E:19:GLY:N	2.26	0.51
41:75:24:PRO:HA	41:75:49:VAL:HG23	1.91	0.51
38:88:4:PRO:HD3	38:88:70:PRO:O	2.10	0.51
9:8E:3:GLN:HB3	9:8E:20:ARG:HD3	1.92	0.51
17:8I:88:TYR:HD1	17:8I:89:LEU:HD23	1.76	0.51
18:9I:34:TYR:HB3	18:9I:69:THR:HG23	1.92	0.51
47:D5:126:VAL:HG12	47:D5:163:LEU:HA	1.92	0.51
47:H8:128:VAL:HG12	47:H8:161:VAL:HB	1.91	0.51
55:P8:5:TRP:HA	55:P8:5:TRP:CE3	2.44	0.51
1:13:1346:A:H5"	9:8E:120:ARG:HH12	1.73	0.51
1:13:1478:C:H2'	1:13:1479:C:H6	1.76	0.51
1:13:963:G:N2	1:13:972:C:N3	2.43	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1007:C:OP1	35:15:37:LYS:NZ	2.37	0.51
26:14:1003:G:N2	26:14:1153:C:C2	2.79	0.51
26:14:1342:A:H2	26:14:1602:U:N3	2.05	0.51
26:14:1464:C:HO2'	26:14:1528:A:H8	1.58	0.51
26:14:270(E):G:N2	26:14:270(U):C:O2	2.35	0.51
35:15:96:GLU:H	35:15:96:GLU:CD	2.13	0.51
29:19:73:VAL:HG13	29:19:120:GLY:HA3	1.92	0.51
1:1G:1133:G:N2	1:1G:1141:C:O2	2.43	0.51
1:1G:1160:G:H2'	1:1G:1161:C:C6	2.45	0.51
1:1G:1216:G:H2'	1:1G:1217:C:C6	2.45	0.51
1:1G:827:U:H3	1:1G:872:A:N6	2.06	0.51
26:1H:1494:A:C2'	26:1H:1495:A:H5'	2.41	0.51
26:1H:234:C:H2'	26:1H:235:U:H6	1.74	0.51
26:1H:984:A:H5"	26:1H:985:C:H5	1.75	0.51
22:1K:76:A:H1'	26:1H:2583:G:N2	2.25	0.51
23:2L:60:A:H2'	23:2L:61:U:H5'	1.93	0.51
4:32:24:GLU:HG2	4:32:25:ARG:H	1.74	0.51
12:3A:70:ILE:HD13	12:3A:77:LEU:HD12	1.92	0.51
5:42:76:ILE:HG23	5:42:142:LEU:HD13	1.93	0.51
26:14:2839:G:H5'	39:55:46:GLY:CA	2.40	0.51
33:59:125:VAL:HG22	33:59:126:PRO:HA	1.91	0.51
33:59:92:ILE:HD13	33:59:160:LYS:HD3	1.92	0.51
36:68:58:VAL:HG21	36:68:86:ILE:HG12	1.93	0.51
8:7E:120:THR:H	8:7E:123:GLU:CG	2.23	0.51
8:7E:33:GLU:HA	8:7E:36:LEU:HD12	1.92	0.51
40:A8:32:LEU:O	40:A8:62:LYS:NZ	2.35	0.51
41:B8:12:SER:CB	41:B8:15:VAL:H	2.23	0.51
47:H8:126:VAL:HG12	47:H8:163:LEU:HA	1.91	0.51
1:13:953:G:H2'	1:13:954:G:O4'	2.11	0.51
26:14:1530:G:O6	26:14:1542:G:N2	2.43	0.51
26:14:2468:G:H3'	26:14:2476:A:N1	2.25	0.51
35:15:73:THR:HG22	35:15:84:LYS:HB3	1.93	0.51
27:16:40:U:H1'	27:16:45:A:H61	1.76	0.51
27:16:78:A:C2	27:16:99:A:C4	2.99	0.51
1:1G:134:A:H61	16:7A:25:ARG:HH12	1.59	0.51
26:1H:1432:C:H2'	26:1H:1433:U:O4'	2.11	0.51
26:1H:1942:C:OP2	26:1H:1943:U:O2'	2.18	0.51
26:1H:7:G:H1	26:1H:2896:C:H42	1.58	0.51
26:1H:581:C:H2'	26:1H:582:G:C8	2.45	0.51
22:1K:43:U:H2'	22:1K:44:U:C6	2.46	0.51
30:29:171:GLU:OE1	30:29:185:LYS:HE2	2.11	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:2I:53:SER:HA	11:2I:55:LYS:HB2	1.93	0.51
4:3E:164:ALA:O	4:3E:168:ARG:NE	2.43	0.51
5:42:101:ILE:HD11	5:42:119:LEU:HD23	1.93	0.51
5:42:78:HIS:HD2	8:72:107:LEU:HD22	1.75	0.51
38:45:137:TYR:CD1	38:45:137:TYR:C	2.84	0.51
5:4E:91:LEU:HD12	5:4E:120:THR:HG22	1.92	0.51
13:4I:46:LYS:HZ2	13:4I:46:LYS:HB2	1.75	0.51
13:4I:60:VAL:HG12	13:4I:66:LEU:HD11	1.92	0.51
35:58:96:GLU:HB2	35:58:122:VAL:CG1	2.40	0.51
33:59:6:ARG:HG2	33:59:7:LEU:H	1.74	0.51
6:5E:44:GLY:HA2	6:5E:59:TYR:CZ	2.46	0.51
40:65:67:ARG:HH21	40:65:103:GLU:HB2	1.75	0.51
42:85:66:ASN:HB2	42:85:76:TYR:HB2	1.93	0.51
39:98:78:LYS:HE2	39:98:83:ILE:HD11	1.91	0.51
40:A8:37:ALA:HB2	40:A8:101:LEU:HD21	1.93	0.51
46:C5:17:SER:O	46:C5:21:LYS:HB2	2.11	0.51
46:C5:62:GLU:CD	46:C5:63:LYS:H	2.14	0.51
26:1H:111:A:H4'	50:K8:69:ARG:NH2	2.26	0.51
1:13:1007:C:N4	1:13:1022:G:H1	2.05	0.51
1:13:1348:U:H2'	1:13:1349:A:H8	1.76	0.51
1:13:626:U:C2	1:13:627:G:C8	2.99	0.51
1:13:748:C:O5'	1:13:748:C:H6	1.94	0.51
26:14:1533:C:H5'	26:14:1534:G:OP2	2.11	0.51
26:14:2107:C:N3	26:14:2182:G:N2	2.56	0.51
26:14:733:G:N7	61:14:3437:SPE:H121	2.26	0.51
26:14:882:G:H22	26:14:894:C:H42	1.59	0.51
1:1G:108:G:H5'	1:1G:109:A:C5'	2.41	0.51
1:1G:1369:C:OP2	9:82:112:LYS:N	2.44	0.51
1:1G:626:U:C2	1:1G:627:G:C8	2.99	0.51
26:1H:1210:A:H5"	26:1H:1212:G:O4'	2.11	0.51
26:1H:2564:A:C2	26:1H:2647:U:H4'	2.46	0.51
26:1H:443:A:OP2	26:1H:615:G:N2	2.33	0.51
37:35:107:LYS:O	37:35:109:GLY:N	2.40	0.51
26:14:1247:A:OP1	31:39:95:ARG:NH2	2.44	0.51
4:3E:149:ALA:O	4:3E:153:ARG:HG2	2.11	0.51
4:3E:85:LYS:HG3	4:3E:88:VAL:O	2.11	0.51
33:51:169:VAL:HG13	33:51:170:ARG:HG3	1.93	0.51
28:71:163:PHE:HB3	28:71:192:PHE:HZ	1.76	0.51
8:72:110:ALA:O	8:72:121:ASP:N	2.44	0.51
26:14:495:G:N2	44:A5:61:ASN:HD21	2.09	0.51
49:F5:84:GLY:HA3	49:F5:87:PRO:HD2	1.93	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:M5:51:ALA:HB1	56:M5:52:LYS:HA	1.91	0.51
1:13:1133:G:H2'	1:13:1134:G:H8	1.75	0.51
1:13:144:G:H2'	1:13:145:G:O4'	2.10	0.51
1:13:649:G:H2'	1:13:650:G:H8	1.74	0.51
1:13:674:G:H2'	1:13:675:A:C8	2.46	0.51
26:14:1291:C:H2'	26:14:1292:U:C6	2.46	0.51
26:14:1357:U:H2'	26:14:1358:G:O4'	2.11	0.51
26:14:1486:A:H2'	26:14:1487:G:H8	1.76	0.51
26:14:2287:A:C2	26:14:2346:A:H2	2.28	0.51
26:14:2461:C:H2'	26:14:2462:U:C6	2.46	0.51
26:14:34:C:HO2'	26:14:35:G:P	2.34	0.51
26:14:890:A:H2'	26:14:892:G:C8	2.46	0.51
1:1G:1080:A:OP1	5:42:14:ARG:NH2	2.44	0.51
1:1G:1423:G:H2'	1:1G:1424:C:C6	2.45	0.51
1:1G:967:C:H3'	1:1G:968:A:H2'	1.93	0.51
26:1H:125:G:H5'	26:1H:125:G:H8	1.76	0.51
26:1H:274:G:H2'	26:1H:275:G:O4'	2.10	0.51
26:1H:353:G:H2'	26:1H:354:G:H8	1.76	0.51
26:1H:721:C:H2'	26:1H:722:A:H8	1.75	0.51
30:21:36:ARG:NH1	30:21:85:ASN:OD1	2.43	0.51
30:29:171:GLU:O	30:29:184:VAL:HA	2.10	0.51
31:31:29:ASN:HB3	31:31:112:MET:HE1	1.92	0.51
1:1G:407:G:P	4:32:115:ARG:HH21	2.33	0.51
4:3E:108:LEU:HD12	4:3E:170:VAL:HG11	1.93	0.51
4:3E:9:CYS:O	4:3E:13:ARG:HG3	2.11	0.51
12:3I:58:VAL:O	12:3I:65:GLU:HA	2.11	0.51
5:42:37:ARG:HG2	5:42:112:LEU:HA	1.92	0.51
39:55:57:ARG:HE	39:55:59:ASP:CG	2.15	0.51
34:61:5:LEU:HD13	34:61:13:GLY:O	2.11	0.51
28:71:64:LEU:HG	28:71:65:PRO:HD2	1.92	0.51
38:88:48:GLU:O	38:88:48:GLU:HG3	2.11	0.51
2:12:16:HIS:CE1	2:12:213:LEU:HD22	2.47	0.51
1:13:292:G:N7	1:13:293:G:H1'	2.26	0.51
1:13:765:G:H5"	1:13:766:A:OP1	2.11	0.51
26:14:308:G:H5"	26:14:309:G:OP2	2.11	0.51
1:1G:1329:A:H2'	1:1G:1330:U:O4'	2.11	0.51
1:1G:757:U:H2'	1:1G:758:G:O4'	2.11	0.51
22:1K:2:G:N2	22:1K:71:C:O2	2.40	0.51
30:21:29:GLY:H	30:21:51:PHE:HE1	1.58	0.51
12:3A:71:PRO:O	12:3A:102:ARG:NH1	2.42	0.51
4:32:57:ARG:HH22	5:42:107:ARG:HD2	1.76	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:71:11:LEU:HD13	28:71:220:PRO:HB2	1.93	0.51
8:72:38:ILE:HD11	8:72:118:VAL:O	2.11	0.51
43:95:62:LEU:HD21	43:95:95:LEU:HB2	1.92	0.51
41:B8:11:GLU:HG2	41:B8:57:PHE:HB3	1.93	0.51
47:D5:130:PRO:HA	47:D5:133:ILE:HD11	1.93	0.51
49:J8:3:LYS:O	49:J8:12:PRO:HD3	2.11	0.51
26:14:180:G:P	55:L5:32:LYS:HD2	2.51	0.51
54:O8:11:LEU:HD11	54:O8:51:GLU:HG3	1.91	0.51
1:13:57:G:C5	1:13:58:C:C4	2.99	0.50
26:14:2314:C:H5'	32:49:38:VAL:HG11	1.94	0.50
26:14:2441:C:OP2	26:14:2586:C:O2'	2.25	0.50
29:19:108:PRO:HB3	29:19:143:HIS:HE1	1.71	0.50
1:1G:1224:G:O2'	1:1G:1322:C:OP2	2.29	0.50
26:1H:1406:U:H2'	26:1H:1407:C:H6	1.76	0.50
26:1H:274:G:N2	26:1H:276:A:H61	2.09	0.50
26:1H:582:G:H2'	26:1H:583:G:C8	2.46	0.50
27:1J:70:C:H2'	27:1J:71:C:H6	1.76	0.50
3:22:155:GLY:HA3	3:22:196:LEU:HD13	1.92	0.50
3:2E:47:LEU:HG	3:2E:76:VAL:HG12	1.93	0.50
1:1G:552:U:O2'	12:3A:86:ARG:O	2.26	0.50
12:3I:54:LYS:N	12:3I:54:LYS:HD3	2.25	0.50
24:3K:1:G:N3	24:3K:1:G:H2'	2.26	0.50
32:49:75:LYS:HA	32:49:84:LYS:HG3	1.93	0.50
39:55:101:ALA:HA	53:J5:44:THR:HG21	1.92	0.50
7:62:111:ARG:NH2	7:62:122:HIS:HB3	2.26	0.50
8:72:123:GLU:O	8:72:127:LEU:HB2	2.11	0.50
43:95:72:VAL:HG13	43:95:72:VAL:O	2.11	0.50
54:O8:44:ARG:H	54:O8:44:ARG:HH11	1.58	0.50
29:11:17:THR:HG22	29:11:204:ILE:HA	1.93	0.50
2:12:48:MET:HA	2:12:51:LEU:HD11	1.93	0.50
26:14:1034:G:H2'	26:14:1035:U:O4'	2.11	0.50
26:14:1533:C:H3'	26:14:1534:G:H4'	1.93	0.50
26:14:1666:G:OP1	36:25:66:LYS:HD3	2.12	0.50
26:14:2185:C:H2'	26:14:2186:G:C8	2.46	0.50
26:14:2655:G:H1'	26:14:2656:U:H5	1.76	0.50
26:14:997:G:O2'	26:14:998:C:H5'	2.12	0.50
29:19:146:GLU:HB2	29:19:189:CYS:HB3	1.93	0.50
29:19:68:LYS:HB3	29:19:70:TRP:CH2	2.47	0.50
1:1G:1134:G:N2	1:1G:1140:C:O2	2.40	0.50
1:1G:1162:C:N4	1:1G:1174:G:H1	2.07	0.50
1:1G:1179:A:H4'	9:82:103:THR:HA	1.94	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1914:C:H2'	26:1H:1915:U:O4'	2.11	0.50
26:1H:2127:G:H2'	26:1H:2128:C:O4'	2.10	0.50
26:1H:2895:U:H2'	26:1H:2896:C:C6	2.46	0.50
27:1J:88:C:H5"	27:1J:89:G:C5	2.46	0.50
1:1G:537:G:H5"	12:3A:113:ARG:NH1	2.27	0.50
1:1G:562:C:H1'	12:3A:15:ARG:HD2	1.92	0.50
1:1G:10:A:OP2	5:42:126:ARG:HG2	2.11	0.50
13:4A:53:VAL:O	13:4A:57:ARG:N	2.25	0.50
4:3E:197:PRO:HD3	6:52:16:GLN:HG3	1.94	0.50
3:22:13:GLY:HA2	14:5A:57:ARG:HD2	1.93	0.50
34:61:21:VAL:HG21	34:61:25:TYR:HD2	1.76	0.50
7:6E:5:ARG:CZ	7:6E:7:ALA:HA	2.41	0.50
43:D8:60:GLU:HB2	43:D8:97:LYS:HE2	1.93	0.50
46:G8:82:PRO:HG3	46:G8:97:ARG:HB3	1.92	0.50
53:J5:33:CYS:SG	53:J5:46:CYS:SG	3.04	0.50
52:M8:13:ARG:NH1	52:M8:22:ILE:HG23	2.23	0.50
2:12:15:VAL:HB	2:12:16:HIS:CE1	2.46	0.50
1:13:57:G:H2'	1:13:58:C:C6	2.46	0.50
1:13:736:C:H2'	1:13:737:A:C8	2.46	0.50
1:1G:1132:C:H2'	1:1G:1133:G:H8	1.77	0.50
1:1G:1151:A:O2'	1:1G:1152:A:O5'	2.30	0.50
1:1G:661:G:H1	1:1G:744:C:H42	1.59	0.50
26:1H:102:G:OP1	50:K8:7:ARG:NH2	2.44	0.50
26:1H:2282:G:H4'	26:1H:2389:G:O2'	2.12	0.50
26:1H:2377:A:H2'	26:1H:2378:A:C8	2.46	0.50
10:1I:38:ILE:HG12	10:1I:71:LEU:O	2.11	0.50
30:21:105:THR:HG1	30:21:199:ARG:NH2	2.09	0.50
31:31:63:LYS:NZ	31:31:67:GLN:HB2	2.27	0.50
37:35:126:VAL:HA	37:35:145:PRO:HD2	1.93	0.50
12:3I:8:ASN:OD1	17:8I:34:LYS:NZ	2.42	0.50
32:41:35:GLU:OE1	32:41:36:LYS:N	2.44	0.50
1:1G:1348:U:H4'	9:82:120:ARG:HD2	1.93	0.50
39:98:2:ARG:CZ	39:98:2:ARG:HB3	2.41	0.50
20:BA:89:ARG:NH1	20:BA:105:SER:O	2.44	0.50
45:F8:27:THR:HB	45:F8:80:ILE:HB	1.93	0.50
47:H8:111:VAL:HB	47:H8:115:GLY:HA3	1.93	0.50
49:J8:93:GLU:O	49:J8:96:LYS:N	2.35	0.50
53:N8:33:CYS:SG	53:N8:40:LYS:HD3	2.52	0.50
1:13:160:A:N6	1:13:343:U:O2'	2.40	0.50
1:13:950:U:OP2	13:4I:102:ARG:HD2	2.10	0.50
26:14:577:G:O2'	26:14:1254:A:OP1	2.27	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1525:G:H2'	26:14:1526:G:H8	1.76	0.50
26:14:2584:U:H2'	26:14:2585:U:H2'	1.93	0.50
26:14:2820:A:C6	39:55:4:LEU:HD11	2.47	0.50
35:15:133:GLN:O	35:15:134:ARG:HG3	2.11	0.50
29:19:24:ILE:HG13	29:19:83:GLU:HA	1.94	0.50
29:19:35:LYS:HA	29:19:64:ILE:HG22	1.92	0.50
1:1G:1004:A:C2	1:1G:1006:C:H1'	2.46	0.50
1:1G:1288:A:H2'	1:1G:1289:A:H8	1.76	0.50
1:1G:192:U:H2'	1:1G:193:C:C6	2.45	0.50
1:1G:408:A:H2'	1:1G:409:G:O4'	2.11	0.50
26:1H:1055:G:O2'	26:1H:1086:A:N6	2.43	0.50
26:1H:1221:C:H2'	26:1H:1222:C:C6	2.47	0.50
26:1H:1204:A:C2	26:1H:1241:A:N1	2.79	0.50
26:1H:1478:G:O6	26:1H:1510:A:N6	2.44	0.50
26:1H:1530:G:O6	26:1H:1542:G:N2	2.44	0.50
26:1H:2170:A:OP2	26:1H:2170:A:H3'	2.11	0.50
26:1H:2400:G:H2'	26:1H:2401:U:C6	2.44	0.50
26:1H:2031:A:C6	26:1H:2498:C:H1'	2.46	0.50
26:1H:860:U:H5	26:1H:917:A:H2	1.46	0.50
26:1H:962:G:H2'	26:1H:963:U:C6	2.46	0.50
27:1J:51:G:C6	27:1J:52:A:C2	2.99	0.50
4:32:33:MET:C	4:32:35:ARG:HH12	2.13	0.50
4:3E:153:ARG:HB3	4:3E:181:MET:SD	2.51	0.50
24:3K:2:G:H2'	24:3K:3:G:C8	2.46	0.50
32:49:53:LEU:HD13	32:49:90:LEU:HD21	1.93	0.50
41:75:4:GLY:N	41:75:5:ALA:C	2.65	0.50
46:C5:14:LEU:HB2	46:C5:75:ILE:HD11	1.93	0.50
37:78:65:ARG:HD3	56:Q8:25:MET:SD	2.52	0.50
1:13:145:G:H1	1:13:177:C:N4	2.09	0.50
26:14:1412:A:H2'	26:14:1413:G:C8	2.47	0.50
26:14:1729:A:H2'	26:14:1731:G:H22	1.74	0.50
10:1A:78:ASN:O	10:1A:81:THR:OG1	2.27	0.50
2:1E:19:HIS:NE2	2:1E:206:ASP:HB2	2.25	0.50
1:1G:1206:G:OP1	3:22:190:ARG:NH2	2.38	0.50
1:1G:1224:G:C6	1:1G:1322:C:H1'	2.46	0.50
1:1G:328:C:H4'	1:1G:329:A:H5"	1.94	0.50
1:1G:539:A:H2'	1:1G:540:G:C8	2.47	0.50
1:1G:664:G:P	18:9A:64:ARG:HH21	2.34	0.50
26:1H:1731:G:H2'	26:1H:1732:A:H8	1.76	0.50
26:1H:2845:G:H2'	26:1H:2846:G:C8	2.47	0.50
27:1J:103:U:O2'	47:D5:72:ARG:HG3	2.10	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
27:1J:13:A:H2'	27:1J:70:C:O2'	2.12	0.50
27:1J:52:A:N6	40:65:33:LYS:HG3	2.27	0.50
27:1J:90:C:OP2	38:45:16:ARG:NH2	2.45	0.50
3:22:9:GLY:HA3	14:5A:49:HIS:HA	1.93	0.50
32:41:28:VAL:O	32:41:31:VAL:HG13	2.12	0.50
32:49:56:ALA:HB2	32:49:153:ARG:CZ	2.41	0.50
13:4I:49:THR:HB	13:4I:52:GLU:HG2	1.94	0.50
13:4I:81:LEU:HD22	13:4I:88:ARG:HB3	1.92	0.50
1:13:667:G:H4'	15:6I:51:HIS:CE1	2.46	0.50
17:8A:45:HIS:HB2	17:8A:65:ILE:HD13	1.92	0.50
39:98:67:LEU:HD22	39:98:76:VAL:HG21	1.93	0.50
18:9I:59:SER:H	18:9I:62:GLU:HB2	1.75	0.50
40:A8:89:ARG:HG3	40:A8:92:TYR:O	2.12	0.50
26:14:372:G:OP2	49:F5:69:LYS:NZ	2.45	0.50
1:13:1003:G:H2'	1:13:1004:A:H4'	1.93	0.50
1:13:1226:C:H4'	19:AI:80:TYR:OH	2.12	0.50
1:13:1459:C:OP1	20:BI:31:SER:OG	2.25	0.50
1:13:259:G:C6	1:13:260:G:C5	2.99	0.50
1:13:323:U:H2'	1:13:324:G:O4'	2.12	0.50
1:13:134:A:H1'	1:13:325:A:C5	2.47	0.50
26:14:1338:G:N3	26:14:1393:A:H2	2.10	0.50
26:14:1366:A:H2'	26:14:1367:A:O4'	2.11	0.50
26:14:2012:G:OP1	44:A5:11:ARG:NH2	2.43	0.50
26:14:2147:G:C5	26:14:2148:G:H1'	2.46	0.50
26:14:2429:G:O6	37:35:61:ARG:NH2	2.44	0.50
26:14:900:A:N3	26:14:900:A:H2'	2.27	0.50
29:19:44:ASN:HA	29:19:47:GLY:N	2.13	0.50
1:1G:1007:C:H2'	1:1G:1008:C:C6	2.47	0.50
1:1G:458:C:H2'	1:1G:464:G:C8	2.45	0.50
26:1H:2756:U:H1'	26:1H:2757:A:H5"	1.93	0.50
26:1H:728:G:H4'	29:11:13:ARG:HD3	1.94	0.50
27:1J:2:C:H2'	27:1J:3:C:C6	2.46	0.50
37:35:120:ALA:O	37:35:121:LYS:HD2	2.12	0.50
5:42:61:TYR:HA	5:42:64:ARG:HG3	1.92	0.50
33:51:169:VAL:HG13	33:51:170:ARG:N	2.25	0.50
40:65:36:TYR:HA	40:65:52:SER:HB3	1.93	0.50
7:6E:20:ASP:HB3	7:6E:23:VAL:H	1.76	0.50
16:7I:4:ILE:HA	16:7I:20:VAL:O	2.12	0.50
40:A8:89:ARG:HG2	40:A8:89:ARG:O	2.12	0.50
48:E5:68:GLU:OE1	48:E5:82:ARG:HG3	2.12	0.50
26:1H:1614:A:C2	44:E8:93:ALA:HB2	2.46	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:170:U:H2'	1:13:171:A:H8	1.77	0.50
26:14:1012:U:H5	35:15:28:THR:HG21	1.76	0.50
26:14:1204:A:C2	26:14:1241:A:N1	2.80	0.50
26:14:2674:G:H2'	26:14:2675:A:C8	2.46	0.50
26:14:597:U:H2'	26:14:598:G:C8	2.47	0.50
26:14:910:A:C5	38:45:13:GLN:HG3	2.46	0.50
1:1G:1129:C:H1'	1:1G:1132:C:H41	1.77	0.50
1:1G:1273:G:H3'	1:1G:1274:G:C8	2.45	0.50
1:1G:1375:A:H2'	1:1G:1376:U:O4'	2.11	0.50
26:1H:2335:A:C8	26:1H:2337:G:C5	3.00	0.50
26:1H:459:U:H2'	26:1H:460:A:C8	2.46	0.50
26:1H:581:C:H2'	26:1H:582:G:H8	1.76	0.50
27:1J:78:A:H2'	27:1J:79:C:O4'	2.12	0.50
30:21:55:ASN:HB3	30:21:58:ARG:H	1.76	0.50
4:3E:60:GLU:OE2	4:3E:199:ASN:N	2.42	0.50
38:45:132:VAL:HG21	47:D5:81:ARG:HE	1.77	0.50
1:1G:974:A:P	14:5A:41:ARG:HH22	2.35	0.50
6:5E:82:ARG:HB2	6:5E:85:VAL:HG23	1.94	0.50
27:1J:7:G:O5'	40:65:29:PHE:HE2	1.94	0.50
15:6I:17:ARG:CZ	15:6I:77:ARG:HD2	2.42	0.50
40:A8:111:GLU:HB2	40:A8:112:PHE:CD2	2.46	0.50
42:C8:88:ILE:C	42:C8:90:VAL:H	2.13	0.50
47:D5:105:VAL:HG13	47:D5:106:GLY:H	1.77	0.50
49:F5:45:ASN:O	49:F5:63:ALA:HA	2.11	0.50
26:1H:2331:G:O3'	48:I8:43:THR:HG22	2.11	0.50
1:13:191(F):U:H2'	1:13:191:G:C8	2.46	0.50
1:13:345:C:H4'	1:13:346:G:C5	2.46	0.50
1:13:407:G:H2'	1:13:408:A:C8	2.46	0.50
29:19:34:VAL:O	29:19:35:LYS:HE2	2.12	0.50
26:1H:1434:A:H61	26:1H:1558:A:H61	1.55	0.50
26:1H:2261:C:H1'	26:1H:2388:A:N3	2.27	0.50
26:1H:26:G:C6	26:1H:27:G:N1	2.80	0.50
26:1H:607:U:N3	26:1H:621:A:C2	2.73	0.50
3:22:40:ARG:HG3	3:22:55:VAL:HG11	1.93	0.50
1:13:1256:A:OP2	3:2E:26:LYS:NZ	2.45	0.50
12:3A:62:SER:OG	12:3A:62:SER:O	2.29	0.50
33:59:37:VAL:HG13	33:59:38:SER:O	2.12	0.50
34:69:8:PRO:N	34:69:15:VAL:HG22	2.26	0.50
15:6A:76:GLU:HA	15:6A:79:ARG:HD2	1.94	0.50
1:13:1291:G:P	7:6E:37:ASN:HD22	2.34	0.50
7:6E:95:ARG:NH2	7:6E:99:LEU:HD11	2.27	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:75:55:ASN:H	41:75:59:THR:HG22	1.75	0.50
19:AI:40:ILE:HG12	19:AI:41:VAL:H	1.76	0.50
47:D5:170:THR:C	47:D5:172:ALA:H	2.15	0.50
26:14:2615:U:C2	53:J5:7:PRO:HA	2.47	0.50
29:11:96:HIS:CE1	29:11:102:LYS:HE2	2.47	0.50
1:13:1287:A:H2'	1:13:1288:A:C8	2.46	0.50
1:13:141:A:H2'	1:13:142:G:H8	1.77	0.50
1:13:429:U:H1'	1:13:430:A:H5"	1.94	0.50
26:14:1165:U:H2'	26:14:1166:C:C6	2.47	0.50
26:14:1434:A:H2'	26:14:1435:G:C8	2.47	0.50
26:14:1686:C:H2'	26:14:1687:G:O4'	2.12	0.50
26:14:395:U:H2'	62:14:4364:HOH:O	2.11	0.50
26:14:64:A:C4	45:B5:66:LEU:HD12	2.47	0.50
26:14:6:A:C3'	26:14:7:G:H5'	2.41	0.50
26:14:729:G:C6	29:19:208:LYS:HB2	2.46	0.50
29:19:137:PRO:O	29:19:140:THR:OG1	2.23	0.50
10:1A:49:VAL:O	10:1A:60:ARG:HB2	2.12	0.50
1:1G:538:G:H2'	1:1G:539:A:C8	2.47	0.50
1:1G:756:C:H2'	1:1G:757:U:O4'	2.11	0.50
26:1H:1509:C:O2'	26:1H:1510:A:OP1	2.23	0.50
26:1H:2593:U:H2'	26:1H:2594:C:C6	2.47	0.50
26:1H:2652:C:H2'	26:1H:2653:U:O4'	2.12	0.50
26:1H:831:G:N7	62:1H:3831:HOH:O	2.34	0.50
22:1K:37:T6A:H2'	22:1K:38:A:O4'	2.12	0.50
30:29:3:GLY:HA3	30:29:81:ILE:HD12	1.93	0.50
30:29:47:VAL:HG21	30:29:85:ASN:HA	1.94	0.50
31:31:134:GLY:CA	31:31:166:ALA:HB2	2.42	0.50
31:39:83:PHE:O	31:39:84:VAL:HB	2.10	0.50
13:4A:78:ILE:HG23	13:4A:92:HIS:ND1	2.27	0.50
8:72:44:PHE:HD1	8:72:80:ILE:HG13	1.76	0.50
37:78:71:VAL:CG1	37:78:72:PRO:HD3	2.39	0.50
42:85:110:VAL:O	42:85:114:LYS:HG2	2.11	0.50
18:9A:36:ASN:N	18:9A:36:ASN:HD22	2.08	0.50
50:G5:15:LYS:H	50:G5:67:LYS:NZ	2.10	0.50
50:G5:47:ASN:N	50:G5:47:ASN:OD1	2.45	0.50
46:G8:89:PHE:CD1	46:G8:90:LEU:N	2.80	0.50
47:H8:152:ALA:HB3	47:H8:167:PRO:O	2.12	0.50
51:L8:35:ARG:HB3	51:L8:37:LEU:HD21	1.94	0.50
2:12:219:VAL:CG2	2:12:221:LEU:H	2.25	0.49
1:13:105:G:H2'	1:13:106:C:C6	2.47	0.49
1:13:1314:C:OP2	19:AI:4:SER:OG	2.29	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:297:G:H4'	1:13:557:G:H4'	1.94	0.49
26:14:1704:G:O6	61:14:3436:SPE:H82	2.11	0.49
26:14:270(F):U:H2'	26:14:270(G):C:C6	2.47	0.49
26:14:309:G:H4'	46:C5:18:GLY:HA3	1.92	0.49
26:14:329:G:P	46:C5:71:LYS:HE3	2.52	0.49
26:14:877:U:H4'	26:14:878:A:OP1	2.10	0.49
29:19:273:ARG:O	29:19:273:ARG:HG2	2.12	0.49
29:19:30:GLU:HG3	29:19:63:ARG:NE	2.26	0.49
2:1E:18:GLY:HA2	2:1E:42:ILE:HG13	1.93	0.49
1:1G:983:A:N1	1:1G:1222:G:N2	2.60	0.49
1:1G:554:C:H2'	1:1G:555:C:C6	2.46	0.49
26:1H:1446:C:H2'	26:1H:1447:G:H8	1.77	0.49
26:1H:1980:G:O2'	26:1H:1982:C:OP2	2.23	0.49
26:1H:2154:G:H2'	26:1H:2155:G:C8	2.43	0.49
26:1H:270(W):G:N7	62:1H:3838:HOH:O	2.35	0.49
26:14:2787:C:O2'	30:29:61:ARG:O	2.16	0.49
26:1H:321:G:O3'	31:31:168:ARG:NH2	2.45	0.49
12:3A:114:LYS:O	12:3A:117:ARG:HG3	2.12	0.49
32:41:33:ARG:O	32:41:162:THR:HG23	2.12	0.49
6:52:2:ARG:NH2	6:52:69:GLU:HG3	2.27	0.49
35:58:133:GLN:C	35:58:134:ARG:HE	2.14	0.49
33:59:86:GLU:H	33:59:86:GLU:CD	2.15	0.49
7:6E:115:ARG:O	7:6E:118:VAL:HG12	2.12	0.49
26:1H:2175:C:H1'	28:71:217:THR:O	2.12	0.49
1:1G:1432:G:OP1	41:75:107:ASP:HB2	2.12	0.49
19:AA:66:MET:HA	19:AA:67:VAL:C	2.32	0.49
47:D5:152:ALA:HB3	47:D5:167:PRO:HA	1.94	0.49
43:D8:3:ALA:HB3	43:D8:14:VAL:HG22	1.93	0.49
50:G5:64:LEU:HD22	50:G5:64:LEU:O	2.12	0.49
50:K8:4:SER:N	50:K8:7:ARG:HG2	2.27	0.49
26:14:1109:C:H2'	26:14:1110:G:H1'	1.93	0.49
26:14:1268:A:H2'	26:14:1269:A:O4'	2.12	0.49
26:14:1274:A:N3	26:14:1297:C:H1'	2.27	0.49
26:14:1828:G:H8	26:14:1828:G:OP2	1.95	0.49
26:14:2103:C:H2'	26:14:2104:G:C8	2.47	0.49
27:16:116:G:H2'	27:16:117:G:O4'	2.11	0.49
1:1G:1324:A:H2'	1:1G:1325:C:C6	2.47	0.49
1:1G:1422:G:H5"	36:25:48:PRO:HB3	1.93	0.49
1:1G:520:A:N1	1:1G:536:C:H1'	2.27	0.49
26:1H:1170:G:N2	26:1H:1180:C:C2	2.80	0.49
26:1H:1864:U:H2'	26:1H:1869:G:H5"	1.93	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2134:A:HO2'	26:1H:2159:G:N2	2.10	0.49
26:1H:330:A:O2'	26:1H:331:A:H8	1.95	0.49
26:1H:375:C:H5"	62:1H:4278:HOH:O	2.12	0.49
26:1H:724:U:H2'	26:1H:725:G:O4'	2.12	0.49
26:1H:840:C:H2'	26:1H:841:A:C8	2.48	0.49
4:32:35:ARG:HH11	4:32:35:ARG:HB2	1.76	0.49
12:3A:86:ARG:HB2	12:3A:101:VAL:HG23	1.94	0.49
4:3E:12:CYS:SG	4:3E:19:LEU:N	2.69	0.49
13:4I:4:ILE:HG22	13:4I:5:ALA:H	1.76	0.49
39:55:56:LYS:NZ	39:55:90:ARG:O	2.44	0.49
1:13:112:G:P	16:7I:27:LYS:HD2	2.53	0.49
1:1G:1117:G:O2'	9:82:104:ARG:HG2	2.12	0.49
43:95:46:VAL:HG22	43:95:52:VAL:HG22	1.92	0.49
18:9A:38:GLU:OE2	18:9A:38:GLU:N	2.42	0.49
41:B8:125:ARG:O	41:B8:129:ARG:N	2.37	0.49
46:G8:89:PHE:HD1	46:G8:90:LEU:H	1.61	0.49
29:11:92:ILE:HD12	29:11:104:TYR:CD1	2.47	0.49
2:12:61:LEU:HG	2:12:160:ASP:CG	2.32	0.49
1:13:1167:A:OP1	1:13:1167:A:H8	1.95	0.49
1:13:1333:A:H2'	1:13:1334:G:O4'	2.12	0.49
1:13:826:C:H2'	1:13:827:U:O2	2.12	0.49
26:14:2119:A:H61	26:14:2170:A:H62	1.60	0.49
26:14:706:A:H2'	26:14:707:G:O4'	2.12	0.49
26:14:958:U:O2	27:1J:89(A):A:O2'	2.24	0.49
26:14:997:G:H2'	26:14:998:C:H6	1.78	0.49
1:1G:362:G:H4'	12:3A:33:ARG:NH2	2.23	0.49
1:1G:397:A:N3	1:1G:397:A:H3'	2.28	0.49
1:1G:957:U:H2'	1:1G:959:A:OP2	2.11	0.49
26:1H:1170:G:N2	26:1H:1180:C:O2	2.45	0.49
26:1H:243:U:OP2	56:Q8:8:LYS:NZ	2.45	0.49
3:22:91:LEU:HD21	3:22:101:LEU:HD21	1.93	0.49
30:29:101:ARG:O	30:29:201:THR:OG1	2.30	0.49
12:3A:88:GLY:O	12:3A:99:HIS:HD2	1.94	0.49
5:42:145:LYS:O	5:42:149:GLU:HG3	2.11	0.49
32:49:42:GLY:O	32:49:43:LEU:HD13	2.12	0.49
7:62:122:HIS:HA	7:62:125:MET:HE2	1.94	0.49
40:65:95:HIS:HA	40:65:99:LYS:HD2	1.94	0.49
15:6I:7:GLU:O	15:6I:11:VAL:HG23	2.13	0.49
9:82:99:LEU:HB3	9:82:101:PHE:CE1	2.47	0.49
17:8A:66:SER:OG	17:8A:69:LYS:HB2	2.13	0.49
26:1H:1754:C:OP1	41:B8:96:ARG:NH1	2.44	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:BA:72:LEU:O	20:BA:73:HIS:HB2	2.11	0.49
49:F5:29:GLY:O	49:F5:30:VAL:HG22	2.11	0.49
46:G8:87:LYS:HD3	46:G8:88:LYS:N	2.28	0.49
53:N8:40:LYS:NZ	53:N8:46:CYS:HB3	2.26	0.49
26:1H:1798:U:H5'	29:11:259:THR:OG1	2.12	0.49
1:13:1413:A:H2	1:13:1487:G:H22	1.60	0.49
1:13:1399:C:C2	1:13:1502:A:N6	2.81	0.49
26:14:1126:A:OP1	26:14:1126:A:H8	1.96	0.49
26:14:1444(A):A:N3	26:14:1444(A):A:H2'	2.28	0.49
26:14:239:U:H2'	26:14:240:G:O4'	2.12	0.49
26:14:2734:A:C8	26:14:2735:G:C8	2.99	0.49
1:1G:1152:A:H5'	10:1A:13:HIS:HD2	1.77	0.49
1:1G:1316:G:N2	1:1G:1318:A:H3'	2.28	0.49
1:1G:313:A:H2'	1:1G:314:C:C6	2.47	0.49
1:1G:910:C:OP2	12:3A:21:LYS:NZ	2.40	0.49
26:1H:141:A:C8	26:1H:1408:C:H1'	2.47	0.49
26:1H:1339:G:N2	26:1H:1603:A:H1'	2.28	0.49
26:1H:1791:A:C8	26:1H:1792:G:C8	3.00	0.49
26:1H:2243:U:H2'	26:1H:2244:U:C6	2.47	0.49
26:1H:198:C:H5'	26:1H:2244:U:OP1	2.12	0.49
30:21:14:ILE:HB	30:21:21:VAL:HG22	1.92	0.49
30:21:15:PHE:HA	30:21:19:ARG:O	2.13	0.49
36:25:98:VAL:HG12	36:25:117:LEU:HB3	1.94	0.49
4:3E:81:GLU:CD	4:3E:139:ARG:HH22	2.16	0.49
13:4I:69:GLU:HG3	32:41:118:ARG:NH2	2.26	0.49
35:58:96:GLU:C	35:58:98:VAL:N	2.64	0.49
26:14:2864:G:OP1	41:75:119:LYS:HD3	2.13	0.49
20:BA:50:GLU:HA	20:BA:100:ILE:HG21	1.93	0.49
42:C8:66:ASN:HB2	42:C8:76:TYR:HB2	1.95	0.49
1:13:963:G:H1	1:13:972:C:N4	2.08	0.49
26:14:2062:A:HO2'	26:14:2063:C:P	2.34	0.49
26:14:2292:C:H4'	26:14:2375:G:H4'	1.93	0.49
26:14:244:A:C2	26:14:255:A:C4	3.00	0.49
26:14:2689:U:H5"	26:14:2713:A:C2	2.47	0.49
26:14:868:U:C4	26:14:869:G:N7	2.80	0.49
35:15:46:VAL:HG12	35:15:48:MET:HG3	1.94	0.49
10:1A:55:LYS:HZ1	10:1A:57:LYS:HG2	1.76	0.49
1:1G:78:G:H1	1:1G:91:C:H42	1.59	0.49
26:1H:107:C:H2'	26:1H:108:U:C6	2.48	0.49
26:1H:2298:A:H2'	26:1H:2299:G:O4'	2.11	0.49
26:1H:2287:A:C2	26:1H:2346:A:C2	2.98	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:1L:34:U:H6	24:1L:34:U:H5"	1.77	0.49
36:25:20:MET:HE3	36:25:44:LYS:HE3	1.94	0.49
32:41:125:PHE:HB3	32:41:166:ASP:OD2	2.13	0.49
33:59:136:ILE:H	33:59:136:ILE:HD12	1.78	0.49
33:59:157:TYR:CD1	33:59:171:LEU:HD23	2.47	0.49
40:65:34:HIS:CE1	40:65:54:LEU:HD12	2.47	0.49
41:75:2:ASN:OD1	41:75:4:GLY:HA3	2.13	0.49
16:7A:9:PHE:HB2	16:7A:16:HIS:O	2.13	0.49
9:82:99:LEU:HB3	9:82:101:PHE:HE1	1.77	0.49
42:85:92:ARG:HD3	42:85:95:LEU:HG	1.94	0.49
48:I8:72:ARG:HH11	48:I8:75:LEU:HD12	1.77	0.49
2:12:219:VAL:HA	2:12:220:ASP:HB3	1.93	0.49
1:13:276:G:O3'	17:8I:68:ARG:NH1	2.45	0.49
1:13:784:C:H2'	1:13:785:G:O4'	2.12	0.49
26:14:1041:C:N4	26:14:1114:G:H22	2.10	0.49
26:14:2784:C:O2	30:29:37:ARG:NH2	2.46	0.49
35:15:15:LEU:HB2	35:15:134:ARG:HB2	1.94	0.49
29:19:70:TRP:CH2	29:19:150:LYS:HA	2.47	0.49
1:1G:1240:U:OP2	7:62:116:ALA:N	2.45	0.49
1:1G:1320:C:H2'	1:1G:1321:C:C6	2.47	0.49
26:1H:814:C:O2'	26:1H:1225:C:N3	2.46	0.49
26:1H:2393:A:H2'	26:1H:2394:C:H6	1.76	0.49
31:39:39:TRP:CH2	31:39:106:ARG:HD3	2.46	0.49
31:39:11:VAL:HG22	31:39:13:SER:CB	2.42	0.49
4:3E:102:ASP:OD1	4:3E:103:ASN:N	2.46	0.49
4:3E:141:ARG:HB2	4:3E:141:ARG:NH1	2.28	0.49
26:1H:558:G:OP1	35:58:111:PRO:HD2	2.11	0.49
34:69:75:LEU:HG	34:69:76:THR:H	1.78	0.49
40:A8:78:LEU:CD1	40:A8:108:GLY:HA2	2.37	0.49
46:G8:20:TYR:CE2	46:G8:43:ASN:HA	2.47	0.49
49:J8:93:GLU:HA	49:J8:96:LYS:HD3	1.92	0.49
2:12:185:ILE:HG23	2:12:199:TYR:HB2	1.95	0.49
1:13:1009:G:C2	1:13:1010:G:C8	3.01	0.49
1:13:1113:C:H2'	1:13:1114:C:H6	1.77	0.49
1:13:1129:C:H3'	1:13:1139:G:N7	2.27	0.49
1:13:671:G:H2'	1:13:672:U:H6	1.76	0.49
26:14:1443:G:H1	26:14:1548:C:H42	1.60	0.49
26:14:2110:G:O2'	26:14:2120:G:H5'	2.13	0.49
26:14:218:A:H2	26:14:235:U:H4'	1.78	0.49
26:14:2520:C:N4	26:14:2542:A:H62	2.08	0.49
26:14:634:C:H2'	26:14:635:C:C6	2.48	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:850:C:O5'	26:14:850:C:H6	1.96	0.49
1:1G:1261:A:H5'	1:1G:1283:G:O3'	2.12	0.49
1:1G:371:G:O2'	1:1G:373:A:N7	2.45	0.49
1:1G:986:A:H1'	19:AA:54:GLY:O	2.12	0.49
26:1H:1103:A:H3'	26:1H:1104:C:C6	2.48	0.49
26:1H:1359:A:N1	26:1H:1372:U:C4	2.79	0.49
26:1H:265:A:H1'	26:1H:266:G:O4'	2.12	0.49
26:1H:2000:G:O2'	26:1H:2689:U:C5	2.60	0.49
26:1H:415:A:H2'	26:1H:416:C:O4'	2.13	0.49
26:1H:606:U:H4'	26:1H:658:C:H4'	1.95	0.49
26:1H:192:C:O2'	26:1H:802:A:N3	2.39	0.49
10:1I:32:ALA:HB3	10:1I:76:ASN:O	2.12	0.49
30:29:112:GLY:O	30:29:159:HIS:HA	2.13	0.49
4:32:105:VAL:HG13	4:32:110:PHE:HB2	1.94	0.49
12:3A:27:LEU:HD23	12:3A:60:LEU:HG	1.93	0.49
33:51:163:TYR:HB3	33:51:167:GLU:HG2	1.94	0.49
37:78:90:ARG:HD3	37:78:91:PHE:CE2	2.47	0.49
48:E5:53:MET:HG3	48:E5:59:LEU:HD23	1.94	0.49
49:J8:85:LEU:O	49:J8:86:SER:OG	2.27	0.49
29:11:132:PRO:HG3	29:11:190:TYR:CE1	2.48	0.49
2:12:180:LEU:HB2	2:12:182:ILE:HD12	1.93	0.49
1:13:1031:G:H2'	1:13:1032:A:H5'	1.95	0.49
1:13:1369:C:H2'	1:13:1370:G:C8	2.48	0.49
1:13:381:C:H2'	1:13:382:A:O4'	2.13	0.49
1:13:60:A:H4'	1:13:61:G:H5'	1.93	0.49
1:13:67:C:H2'	1:13:68:G:H8	1.75	0.49
1:13:955:U:H1'	1:13:1227:A:N6	2.28	0.49
26:14:2557:G:H2'	26:14:2558:C:C6	2.48	0.49
26:14:2816:C:O3'	39:55:99:LYS:NZ	2.44	0.49
26:14:481:G:OP2	46:C5:47:LYS:HB2	2.13	0.49
26:14:853:G:C2'	26:14:854:G:H5'	2.42	0.49
1:1G:179:A:H2'	1:1G:180:U:C6	2.48	0.49
1:1G:428:G:C5	1:1G:430:A:C6	3.00	0.49
26:1H:1022:G:N7	35:58:66:LYS:NZ	2.61	0.49
26:1H:197:A:N6	26:1H:2430:A:H2'	2.27	0.49
26:1H:2544:G:H2'	26:1H:2545:G:H8	1.77	0.49
26:1H:2592:G:C6	26:1H:2593:U:C4	3.01	0.49
26:1H:598:G:H5'	37:78:11:GLY:HA3	1.94	0.49
3:22:195:VAL:O	3:22:196:LEU:HD22	2.13	0.49
3:22:35:GLU:OE1	3:22:59:ARG:NH2	2.46	0.49
11:2I:109:VAL:HA	18:9I:85:LEU:O	2.12	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:32:26:CYS:HB3	59:32:302:SF4:S1	2.53	0.49
31:39:181:LEU:CD2	31:39:186:ILE:HD11	2.41	0.49
32:41:165:THR:OG1	32:41:168:GLU:HG3	2.13	0.49
32:41:67:LYS:HE2	32:41:67:LYS:H	1.78	0.49
32:49:50:ALA:HA	32:49:53:LEU:HD23	1.93	0.49
13:4A:23:TYR:HB3	13:4A:67:GLU:HA	1.94	0.49
13:4I:23:TYR:CE2	13:4I:71:ARG:HG3	2.48	0.49
33:51:170:ARG:HA	33:51:171:LEU:HB2	1.95	0.49
34:61:37:VAL:HG12	34:61:38:LEU:HD12	1.93	0.49
26:14:2685:G:P	41:75:51:ARG:HH22	2.36	0.49
37:78:114:ILE:HD13	37:78:125:VAL:HG11	1.93	0.49
17:8I:56:VAL:HB	17:8I:78:GLU:HB2	1.93	0.49
20:BI:100:ILE:HG12	20:BI:101:GLY:H	1.77	0.49
46:C5:88:LYS:CG	46:C5:89:PHE:H	2.21	0.49
47:D5:158:PRO:O	47:D5:161:VAL:HG22	2.13	0.49
42:C8:92:ARG:NH1	43:D8:11:GLN:O	2.46	0.49
48:I8:51:VAL:HG23	48:I8:81:VAL:HG23	1.95	0.49
55:P8:10:ARG:O	55:P8:14:LYS:HB2	2.13	0.49
26:1H:459:U:H5"	55:P8:40:TRP:CD2	2.48	0.49
2:12:98:LEU:O	2:12:101:MET:HG2	2.13	0.49
1:13:1159:U:O4'	1:13:1182:G:N2	2.46	0.49
1:13:345:C:O2'	1:13:346:G:N3	2.40	0.49
1:13:450:G:N7	1:13:481:G:C6	2.80	0.49
1:13:4:U:O4	8:7E:105:ARG:HG3	2.13	0.49
26:14:1204:A:H2	26:14:1241:A:N1	2.10	0.49
26:14:2115:G:N1	26:14:2117:A:N7	2.61	0.49
26:14:2394:C:H1'	62:14:3815:HOH:O	2.12	0.49
26:14:249:C:H4'	26:14:250:G:O5'	2.13	0.49
26:14:2869:G:H2'	26:14:2870:C:O4'	2.13	0.49
26:14:581:C:H2'	26:14:582:G:C8	2.45	0.49
1:1G:1353:G:P	21:1B:10:ARG:HH12	2.35	0.49
1:1G:110:C:H2'	1:1G:111:G:O4'	2.12	0.49
1:1G:1300:G:HO2'	1:1G:1301:U:P	2.36	0.49
1:1G:1459:C:OP1	20:BA:31:SER:OG	2.20	0.49
1:1G:20:U:H2'	1:1G:21:G:O4'	2.13	0.49
1:1G:973:G:H5'	10:1A:55:LYS:NZ	2.26	0.49
26:1H:1206:G:C6	26:1H:1207:C:C4	3.01	0.49
26:1H:1434:A:H61	26:1H:1558:A:H62	1.57	0.49
26:1H:2505:G:O6	26:1H:2576:G:H2'	2.12	0.49
27:1J:18:G:H1	27:1J:65:C:H42	1.61	0.49
3:22:119:ARG:NH2	3:22:140:ARG:HD2	2.28	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:35:127:ALA:O	37:35:147:LEU:HB2	2.13	0.49
24:3L:10:G:H22	24:3L:26:A:H1'	1.78	0.49
24:3L:48:C:C6	24:3L:59:A:H1'	2.48	0.49
26:1H:1140:C:OP1	35:58:23:LEU:HB3	2.13	0.49
40:65:101:LEU:O	40:65:105:ALA:N	2.43	0.49
9:82:24:GLY:HA2	9:82:59:PHE:O	2.13	0.49
44:A5:75:TYR:CZ	44:A5:104:THR:HG21	2.48	0.49
26:14:2012:G:OP2	44:A5:16:LYS:NZ	2.45	0.49
27:1J:103:U:O2'	47:D5:29:TYR:OH	2.18	0.49
29:11:142:VAL:HG23	29:11:193:VAL:HA	1.94	0.49
2:12:178:ARG:HD2	2:12:196:LEU:O	2.12	0.49
1:13:1126:U:C4	1:13:1127:G:C4	3.01	0.49
1:13:1469:G:O6	62:13:1841:HOH:O	2.20	0.49
1:13:2:U:H5"	1:13:630:G:N2	2.28	0.49
1:13:813:U:OP2	1:13:816:A:N6	2.42	0.49
26:14:1204:A:O2'	26:14:1205:U:OP2	2.29	0.49
26:14:304:G:H2'	26:14:305:U:C6	2.48	0.49
26:14:311:A:C6	26:14:328:U:C4	3.01	0.49
26:14:77:C:OP1	50:G5:59:ARG:HD3	2.13	0.49
2:1E:158:LEU:HD22	2:1E:182:ILE:HD11	1.95	0.49
1:1G:1011:G:N2	1:1G:1019:C:H1'	2.28	0.49
1:1G:426:G:OP1	4:32:38:TYR:OH	2.26	0.49
1:1G:730:G:C5	1:1G:731:G:H1'	2.48	0.49
1:1G:979:C:H3'	1:1G:980:C:C5'	2.42	0.49
26:1H:2308:G:N3	26:1H:2308:G:H2'	2.28	0.49
26:1H:493:G:H2'	26:1H:494:G:O4'	2.12	0.49
26:1H:721:C:H2'	26:1H:722:A:C8	2.47	0.49
27:1J:44:G:H5"	27:1J:45:A:OP1	2.13	0.49
30:29:147:PRO:HB2	30:29:149:ARG:HD3	1.95	0.49
30:29:81:ILE:HG22	30:29:82:ARG:N	2.23	0.49
24:3L:22:G:C8	24:3L:46:G:N2	2.79	0.49
38:45:66:ILE:HG22	38:45:104:PHE:HE1	1.78	0.49
33:59:149:ARG:NH1	33:59:162:ILE:O	2.46	0.49
7:62:116:ALA:O	7:62:120:ILE:HG12	2.13	0.49
34:69:102:SER:OG	34:69:103:ARG:N	2.45	0.49
15:6A:11:VAL:HG21	15:6A:34:LEU:HD12	1.94	0.49
39:98:78:LYS:O	39:98:83:ILE:HG13	2.12	0.49
40:A8:70:GLY:HA2	40:A8:101:LEU:HD13	1.95	0.49
47:D5:40:ASP:OD1	47:D5:43:GLU:N	2.33	0.49
47:D5:6:LYS:HG3	47:D5:7:ALA:H	1.77	0.49
47:H8:52:SER:O	47:H8:53:ILE:HG12	2.13	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:O8:41:PRO:HB2	54:O8:44:ARG:NH1	2.27	0.49
56:Q8:15:LYS:HG3	56:Q8:16:ILE:N	2.28	0.49
26:1H:1491:G:O2'	29:11:101:GLU:HB2	2.13	0.48
1:13:1071:C:H2'	1:13:1072:G:H8	1.77	0.48
1:13:1308:U:OP1	13:4I:98:VAL:HG22	2.14	0.48
1:13:35:G:H2'	1:13:36:C:C6	2.48	0.48
1:13:411:A:C8	1:13:413:G:H1'	2.48	0.48
1:13:953:G:C2	1:13:954:G:H1'	2.48	0.48
26:14:1181:C:H2'	26:14:1182:A:H8	1.78	0.48
26:14:2280:G:C2'	26:14:2281:C:H5'	2.43	0.48
26:14:959:A:N6	26:14:960:A:N1	2.61	0.48
26:14:975:G:H1'	26:14:990:A:C2	2.47	0.48
1:1G:1143:G:H2'	1:1G:1144:G:C8	2.48	0.48
1:1G:976:G:N7	1:1G:1358:U:C5	2.80	0.48
1:1G:227:G:H2'	1:1G:228:A:C8	2.48	0.48
1:1G:266:G:H5'	1:1G:268:C:H41	1.78	0.48
1:1G:433:C:H2'	1:1G:434:U:H6	1.78	0.48
1:1G:592:G:H1	1:1G:647:C:H42	1.61	0.48
1:1G:828:A:H5"	1:1G:859:A:C2	2.48	0.48
26:1H:1243:G:O2'	37:78:7:ARG:NH2	2.46	0.48
26:1H:1358:G:N2	26:1H:1372:U:C5	2.81	0.48
26:1H:1705:G:C6	26:1H:1706:U:C4	3.00	0.48
26:1H:2056:G:C2	26:1H:2057:A:C8	3.01	0.48
26:1H:2287:A:N6	26:1H:2344:U:H3	2.10	0.48
26:1H:234:C:H2'	26:1H:235:U:C6	2.47	0.48
26:1H:2854:G:H2'	26:1H:2855:C:C6	2.48	0.48
26:1H:847:U:C5	26:1H:933:A:N1	2.81	0.48
31:39:68:LYS:HB3	31:39:69:HIS:CD2	2.48	0.48
38:45:32:TYR:CE1	38:45:133:ARG:HG3	2.48	0.48
13:4I:79:LYS:O	13:4I:83:ASP:HB2	2.13	0.48
34:61:110:ASP:OD1	34:61:111:PRO:HA	2.12	0.48
44:A5:110:LYS:HG3	44:A5:111:HIS:ND1	2.27	0.48
20:BA:33:ILE:HD11	20:BA:62:LEU:O	2.13	0.48
43:D8:37:VAL:O	43:D8:38:LEU:HG	2.13	0.48
51:H5:6:VAL:O	51:H5:34:GLU:HA	2.13	0.48
26:1H:2261:C:C5	48:I8:16:SER:HB3	2.48	0.48
1:13:1016:A:H2'	1:13:1017:G:O4'	2.13	0.48
1:13:347:G:H2'	1:13:348:G:O4'	2.12	0.48
1:13:814:A:N7	1:13:816:A:C4	2.81	0.48
26:14:152:G:H1	26:14:174:C:N4	2.04	0.48
26:14:1585:C:H2'	26:14:1585:C:O2	2.12	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1973:G:H2'	26:14:1974:C:C6	2.47	0.48
26:14:1999:C:H4'	26:14:2723:C:O2	2.13	0.48
26:14:2315:G:H2'	26:14:2316:C:C6	2.48	0.48
26:14:569:U:C4	26:14:570:G:C6	3.01	0.48
26:14:627:A:H62	37:35:84:ASN:ND2	2.11	0.48
1:1G:1316:G:HO2'	1:1G:1318:A:H62	1.60	0.48
1:1G:1372:U:OP1	9:82:72:GLY:N	2.46	0.48
1:1G:141:A:H1'	1:1G:182:U:O2	2.13	0.48
1:1G:995:C:O2'	1:1G:996:A:H5'	2.13	0.48
26:1H:141:A:H8	26:1H:1595:G:H21	1.60	0.48
26:1H:2320:A:N3	26:1H:2320:A:H2'	2.29	0.48
26:1H:2462:U:H1'	26:1H:2491:U:O4	2.13	0.48
26:1H:558:G:P	35:58:111:PRO:HD2	2.53	0.48
11:2I:41:THR:HG21	11:2I:71:LYS:HD2	1.95	0.48
31:31:134:GLY:HA2	31:31:166:ALA:HB2	1.95	0.48
1:1G:407:G:O2'	4:32:116:GLN:HG3	2.13	0.48
26:14:322:A:OP2	31:39:169:ASN:HB2	2.13	0.48
5:42:107:ARG:NH2	5:42:108:ALA:HB2	2.28	0.48
32:49:120:LEU:HG	32:49:179:PRO:O	2.12	0.48
6:52:68:PRO:HG2	6:52:71:ARG:HG3	1.94	0.48
34:61:81:VAL:HG11	34:61:88:ILE:HD13	1.94	0.48
16:7A:45:THR:O	16:7A:48:TRP:HD1	1.97	0.48
8:7E:10:LEU:HD22	8:7E:83:ILE:HD11	1.95	0.48
9:8E:114:TYR:CE2	10:1I:59:SER:HA	2.48	0.48
30:21:111:ARG:HA	39:98:1:MET:HE3	1.95	0.48
42:C8:75:ASN:HB3	42:C8:77:SER:H	1.78	0.48
42:C8:90:VAL:O	42:C8:92:ARG:N	2.46	0.48
50:G5:4:SER:N	50:G5:5:GLU:HB2	2.28	0.48
54:O8:47:THR:HG22	54:O8:48:VAL:HG23	1.95	0.48
54:O8:47:THR:HG22	54:O8:48:VAL:H	1.77	0.48
2:12:95:GLN:HB3	2:12:148:TYR:HD1	1.78	0.48
1:13:1210:C:H2'	1:13:1211:U:H5'	1.94	0.48
1:13:1417:G:N2	1:13:1482:G:H2'	2.28	0.48
1:13:148:G:H2'	1:13:149:A:H8	1.76	0.48
26:14:1332:G:H5'	26:14:1332:G:C8	2.47	0.48
26:14:141:A:C8	26:14:1408:C:H1'	2.48	0.48
26:14:1622:G:OP1	62:14:3652:HOH:O	2.20	0.48
26:14:2126:A:H2	26:14:2162:G:H22	1.60	0.48
26:14:376:C:H2'	26:14:377:C:C6	2.48	0.48
2:1E:25:ASN:ND2	2:1E:193:ASP:HB3	2.28	0.48
1:1G:1084:G:H5'	1:1G:1102:A:OP2	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:956:U:H1'	1:1G:1225:A:H2	1.76	0.48
1:1G:407:G:H2'	1:1G:408:A:C8	2.48	0.48
1:1G:457:C:H2'	1:1G:458:C:C6	2.48	0.48
26:1H:1441:G:H2'	26:1H:1442:G:H8	1.78	0.48
26:1H:2259:G:H1'	26:1H:2427:C:H2'	1.95	0.48
26:1H:2502:G:N7	62:1H:3840:HOH:O	2.35	0.48
26:1H:2895:U:H2'	26:1H:2896:C:H6	1.78	0.48
26:1H:723:G:H2'	26:1H:724:U:O4'	2.13	0.48
26:1H:811:U:H2'	37:78:21:ARG:HA	1.94	0.48
26:1H:991:C:H2'	26:1H:992:C:C6	2.48	0.48
3:22:70:VAL:HG12	3:22:72:LYS:N	2.25	0.48
23:2L:2:G:H2'	23:2L:3:C:C6	2.48	0.48
31:31:197:ASP:O	31:31:199:TRP:N	2.46	0.48
5:42:146:ALA:HB1	5:42:150:ARG:HH21	1.78	0.48
32:49:101:ILE:HD12	32:49:105:LYS:HD2	1.95	0.48
32:49:97:ASP:O	32:49:101:ILE:HG23	2.14	0.48
13:4I:11:ARG:HG2	13:4I:12:ASN:H	1.79	0.48
1:1G:1535:C:N4	25:4L:10:G:H21	2.08	0.48
33:51:118:PRO:HD2	33:51:121:ILE:HG21	1.95	0.48
34:61:86:THR:HA	34:61:123:LEU:HD13	1.94	0.48
26:1H:2469:A:O2'	38:88:56:ARG:NH1	2.46	0.48
1:13:1342:C:H4'	9:8E:125:TYR:HB3	1.94	0.48
17:8I:29:HIS:CD2	17:8I:30:PRO:HD2	2.48	0.48
44:A5:34:ASN:ND2	53:J5:39:MET:HG3	2.28	0.48
19:AI:23:ASN:HD21	19:AI:43:GLU:HB2	1.79	0.48
41:B8:60:THR:HG22	41:B8:77:PRO:HA	1.95	0.48
42:C8:88:ILE:C	42:C8:90:VAL:N	2.67	0.48
51:H5:6:VAL:HB	51:H5:54:VAL:HG21	1.93	0.48
45:F8:5:TYR:O	50:K8:36:ARG:NH2	2.46	0.48
51:L8:18:ASP:HB3	62:L8:102:HOH:O	2.13	0.48
1:13:198:G:N7	1:13:220:G:N2	2.61	0.48
1:13:260:G:H2'	1:13:261:U:C6	2.48	0.48
26:14:1019:U:H2'	26:14:1020:A:C8	2.48	0.48
26:14:1181:C:H2'	26:14:1182:A:C8	2.49	0.48
26:14:1224:G:N2	26:14:1227:A:OP2	2.34	0.48
26:14:1270:C:H5"	26:14:1271:G:O5'	2.12	0.48
26:14:140:A:C8	26:14:1408:C:O2'	2.61	0.48
26:14:2238:G:N3	26:14:2238:G:H2'	2.28	0.48
26:14:774:A:H2	26:14:787:U:O2'	1.96	0.48
26:14:881:G:H3'	26:14:882:G:C8	2.48	0.48
1:1G:1054:C:H4'	1:1G:1055:A:H5"	1.96	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1255:G:OP1	10:1A:45:ARG:NH1	2.47	0.48
1:1G:922:G:N3	1:1G:1398:A:H2	2.10	0.48
1:1G:142:G:H1	1:1G:221:C:H42	1.60	0.48
1:1G:622:A:C8	1:1G:623:C:C6	3.01	0.48
26:1H:2630:G:H2'	26:1H:2631:G:O4'	2.13	0.48
26:1H:265:A:C8	26:1H:266:G:H1'	2.48	0.48
26:1H:2772:C:H2'	26:1H:2773:C:C6	2.48	0.48
26:1H:500:G:N2	26:1H:502:A:H3'	2.28	0.48
3:2E:113:ALA:HB2	3:2E:202:ILE:HG13	1.94	0.48
4:32:61:LYS:HB2	4:32:203:VAL:HG13	1.95	0.48
4:32:89:THR:HG21	4:32:204:ILE:HD11	1.95	0.48
24:3K:3:G:H1	24:3K:70:C:H42	1.61	0.48
28:71:53:ARG:HA	28:71:53:ARG:HD3	1.54	0.48
16:7A:43:LYS:HG2	16:7A:48:TRP:CD2	2.47	0.48
16:7A:40:ASP:HB3	16:7A:48:TRP:HB2	1.95	0.48
42:85:90:VAL:O	42:85:92:ARG:N	2.47	0.48
27:16:90:C:H5'	38:88:18:LYS:HA	1.95	0.48
43:95:39:LEU:HD23	43:95:40:LEU:N	2.28	0.48
18:9A:22:VAL:HG12	18:9A:55:ARG:O	2.14	0.48
20:BI:87:LYS:HA	20:BI:87:LYS:HD2	1.53	0.48
46:C5:8:LYS:NZ	46:C5:95:LYS:HD3	2.28	0.48
42:C8:79:PHE:O	42:C8:79:PHE:HD1	1.96	0.48
47:D5:174:VAL:O	47:D5:175:VAL:HB	2.12	0.48
46:G8:88:LYS:HD2	46:G8:92:ASN:OD1	2.13	0.48
50:K8:64:LEU:HD11	50:K8:68:ARG:HH11	1.79	0.48
2:12:210:SER:O	2:12:214:ILE:HG12	2.13	0.48
1:13:872:A:C4	1:13:874:G:N7	2.81	0.48
26:14:1375:C:H2'	26:14:1376:C:H6	1.77	0.48
26:14:483:A:H1'	46:C5:60:PHE:HE1	1.77	0.48
26:14:780:G:H21	26:14:783:A:N6	2.03	0.48
1:1G:1006:C:H2'	1:1G:1007:C:C6	2.48	0.48
1:1G:1342:C:H1'	9:82:124:GLN:NE2	2.29	0.48
1:1G:938:A:N3	1:1G:1376:U:O2'	2.33	0.48
26:1H:1001:A:H2'	26:1H:1002:G:O4'	2.13	0.48
26:1H:1126:A:H8	26:1H:1126:A:O5'	1.96	0.48
26:1H:1614:A:N1	44:E8:93:ALA:HB2	2.28	0.48
26:1H:2159:G:H2'	26:1H:2160:G:H8	1.78	0.48
26:1H:2172:U:H5'	26:1H:2173:A:OP2	2.13	0.48
26:1H:2186:G:H2'	26:1H:2187:G:H8	1.78	0.48
26:1H:2414:G:H21	37:78:67:MET:HE1	1.78	0.48
26:1H:2846:G:H2'	26:1H:2847:U:O4'	2.13	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:698:C:O2'	26:1H:734:A:N6	2.47	0.48
26:1H:916:G:C2'	26:1H:917:A:H5"	2.43	0.48
1:13:963:G:H21	10:1I:55:LYS:HZ3	1.60	0.48
27:1J:44:G:O2'	27:1J:47:C:N4	2.46	0.48
30:29:60:ASN:OD1	30:29:61:ARG:N	2.47	0.48
23:2L:41:C:H2'	23:2L:42:C:C6	2.48	0.48
37:35:132:LYS:HD2	37:35:132:LYS:HA	1.61	0.48
12:3I:24:VAL:HB	12:3I:27:LEU:HD12	1.96	0.48
32:49:73:ALA:HB3	32:49:85:GLY:H	1.78	0.48
13:4A:13:LYS:HA	13:4A:44:ARG:HH11	1.79	0.48
34:61:10:GLU:O	34:61:10:GLU:HG3	2.12	0.48
36:68:64:ARG:HD3	36:68:79:PHE:CD2	2.49	0.48
15:6A:87:ILE:HG22	15:6A:88:ARG:N	2.29	0.48
1:1G:640:A:N3	8:72:115:SER:HB2	2.28	0.48
8:72:14:ARG:O	8:72:18:ARG:HG2	2.13	0.48
26:14:994:C:OP1	42:85:53:ARG:NH2	2.46	0.48
45:F8:12:VAL:HG13	45:F8:27:THR:O	2.13	0.48
51:L8:7:LYS:HB2	51:L8:34:GLU:HG2	1.94	0.48
1:13:1171:G:H2'	1:13:1172:C:C6	2.49	0.48
1:13:1297:C:OP1	13:4I:13:LYS:NZ	2.39	0.48
1:13:666:G:H5"	1:13:732:C:O2	2.14	0.48
1:13:901:A:C5	1:13:902:G:H1'	2.48	0.48
26:14:1432:C:H2'	26:14:1433:U:O4'	2.13	0.48
26:14:1639:U:O2'	26:14:1640:C:H5'	2.14	0.48
26:14:2138:C:N4	26:14:2153:G:H22	2.12	0.48
26:14:2232:U:OP1	49:F5:40:ARG:NH2	2.47	0.48
26:14:2286:A:H4'	26:14:2287:A:O4'	2.12	0.48
26:14:528:A:N1	26:14:2042:A:H2'	2.28	0.48
26:14:618:G:H2'	26:14:618(A):C:O4'	2.12	0.48
26:14:71:A:H5'	26:14:71:A:H8	1.76	0.48
26:14:755:C:H2'	26:14:756:C:C6	2.48	0.48
29:19:228:PRO:HD3	29:19:235:GLY:N	2.28	0.48
1:1G:1223:C:H5"	1:1G:1224:G:H5"	1.94	0.48
1:1G:1327:C:H2'	1:1G:1328:C:C6	2.47	0.48
1:1G:1329:A:H4'	13:4A:24:GLY:HA2	1.95	0.48
26:1H:1464:C:HO2'	26:1H:1528:A:H8	1.59	0.48
26:1H:1683:C:H2'	26:1H:1684:C:H6	1.78	0.48
26:1H:1683:C:H2'	26:1H:1684:C:C6	2.48	0.48
26:1H:2383:G:O2'	26:1H:2384:G:H5'	2.13	0.48
26:1H:2801:A:H5'	26:1H:2895:U:O2'	2.13	0.48
26:1H:783:A:C8	26:1H:783:A:H3'	2.48	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:1I:50:ILE:HA	10:1I:60:ARG:HB3	1.96	0.48
23:2L:24:C:C2	23:2L:25:U:C5	3.01	0.48
12:3A:47:LYS:CD	12:3A:48:PRO:HD2	2.43	0.48
5:4E:71:LEU:HD22	5:4E:114:GLY:HA3	1.96	0.48
14:5I:9:LYS:HA	14:5I:12:ARG:HG2	1.95	0.48
40:65:87:PHE:CZ	40:65:102:ALA:HB2	2.49	0.48
26:1H:2483:C:N3	38:88:124:LYS:HE3	2.28	0.48
43:95:33:VAL:HG22	43:95:35:LEU:HD23	1.96	0.48
19:AI:40:ILE:HD11	19:AI:62:ILE:HG23	1.95	0.48
30:21:181:LEU:HD21	41:B8:6:LEU:HG	1.95	0.48
1:1G:192:U:O4'	20:BA:103:GLY:HA2	2.13	0.48
44:E8:19:LEU:HB3	53:N8:25:LEU:HD11	1.96	0.48
1:13:1453:G:H2'	20:BI:39:LYS:HE2	1.96	0.48
1:13:674:G:H2'	1:13:675:A:H8	1.79	0.48
26:14:251:A:C5	26:14:252:G:H1'	2.49	0.48
26:14:2820:A:C5	39:55:4:LEU:HD11	2.48	0.48
2:1E:155:LEU:HA	2:1E:155:LEU:HD23	1.72	0.48
1:1G:1287:A:H2'	1:1G:1288:A:C8	2.48	0.48
1:1G:197:A:C6	1:1G:221:C:H4'	2.48	0.48
1:1G:23:C:OP2	1:1G:561:U:N3	2.44	0.48
1:1G:407:G:OP1	4:32:115:ARG:NE	2.43	0.48
26:1H:1142(A):A:C4	26:1H:1144:G:C8	3.00	0.48
26:1H:1164:G:H2'	26:1H:1165:U:C6	2.48	0.48
26:1H:2291:U:H2'	26:1H:2292:C:C6	2.48	0.48
26:1H:389:G:N1	37:78:71:VAL:HG12	2.28	0.48
1:13:1190:G:OP1	3:2E:5:ILE:HG23	2.13	0.48
11:2I:19:ALA:HA	11:2I:32:ILE:HG22	1.95	0.48
4:32:12:CYS:SG	4:32:19:LEU:N	2.70	0.48
4:3E:10:ARG:HB2	4:3E:10:ARG:NH1	2.29	0.48
24:3K:5:C:H2'	24:3K:6:G:C8	2.47	0.48
5:4E:15:ARG:HE	5:4E:26:PHE:HE2	1.59	0.48
6:5E:69:GLU:O	6:5E:72:VAL:HG12	2.14	0.48
14:5I:13:THR:HG23	14:5I:20:ALA:HB2	1.96	0.48
27:1J:116:G:H5"	40:65:55:ALA:HB2	1.95	0.48
41:75:5:ALA:HB3	41:75:9:LEU:HB2	1.94	0.48
26:1H:2292:C:OP2	40:A8:17:ARG:NH2	2.47	0.48
45:B5:67:GLY:C	45:B5:69:TYR:H	2.17	0.48
41:B8:20:PRO:HG2	41:B8:86:ILE:O	2.13	0.48
48:E5:51:VAL:N	48:E5:62:LEU:HD12	2.29	0.48
47:H8:98:MET:O	47:H8:125:LEU:HD12	2.13	0.48
1:13:1122:U:O4	1:13:1123:A:N6	2.47	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:857:C:H2'	26:14:858:U:C6	2.49	0.48
29:19:133:LEU:HD13	29:19:173:VAL:HG11	1.95	0.48
29:19:31:LYS:HZ1	29:19:33:LEU:HB3	1.77	0.48
2:1E:111:ARG:HD2	2:1E:145:LEU:HD21	1.94	0.48
1:1G:1104:G:H4'	2:12:111:ARG:NE	2.29	0.48
1:1G:1151:A:HO2'	1:1G:1152:A:C5'	2.26	0.48
1:1G:164:U:H2'	1:1G:165:C:H6	1.79	0.48
1:1G:322:C:H5	1:1G:328:C:C5	2.32	0.48
1:1G:952:U:H4'	1:1G:964:A:N1	2.29	0.48
1:1G:972:C:C2'	10:1A:55:LYS:HG3	2.44	0.48
26:1H:1108:U:C2'	26:1H:1109:C:H5'	2.44	0.48
26:1H:1171:G:N2	26:1H:1178:C:N3	2.55	0.48
26:1H:1791:A:H5'	29:11:206:LEU:HD12	1.96	0.48
26:1H:2118:U:O4'	26:1H:2147:G:N1	2.43	0.48
26:1H:2126:A:H8	26:1H:2163:C:H1'	1.79	0.48
26:1H:2740:A:H2'	26:1H:2741:A:C8	2.48	0.48
26:1H:518:G:H2'	26:1H:519:U:C6	2.49	0.48
27:1J:88:C:H5"	27:1J:89:G:C6	2.49	0.48
3:2E:57:ILE:HG12	3:2E:66:VAL:HG22	1.96	0.48
11:2I:73:MET:HG2	11:2I:103:LEU:HD13	1.96	0.48
23:2L:54:G:H2'	23:2L:55:5MU:C6	2.49	0.48
31:31:101:LEU:HD23	31:31:102:PRO:CD	2.33	0.48
37:35:101:VAL:HA	37:35:105:LEU:O	2.14	0.48
24:3L:2:G:H1	24:3L:71:C:H42	1.62	0.48
32:41:81:LYS:HZ2	32:41:81:LYS:H	1.61	0.48
13:4A:81:LEU:CD2	13:4A:88:ARG:HH21	2.26	0.48
5:4E:33:VAL:HB	5:4E:112:LEU:HD12	1.95	0.48
33:59:77:LYS:HA	33:59:77:LYS:HD2	1.60	0.48
34:69:39:ALA:O	34:69:44:LEU:HG	2.14	0.48
28:71:45:ALA:H	28:71:171:ILE:HG22	1.79	0.48
8:72:101:PRO:HG2	8:72:133:LEU:HD11	1.95	0.48
19:AI:41:VAL:HB	19:AI:42:PRO:C	2.34	0.48
44:E8:18:ARG:HD3	44:E8:76:VAL:HG13	1.96	0.48
49:J8:51:VAL:HG21	49:J8:74:VAL:HG11	1.95	0.48
50:K8:2:LYS:O	50:K8:3:LEU:HD23	2.14	0.48
2:12:73:THR:HG21	2:12:97:TRP:N	2.29	0.48
1:13:1151:A:O2'	1:13:1152:A:H8	1.96	0.48
1:13:1179:A:H2'	1:13:1180:A:O4'	2.14	0.48
26:14:228:A:H2'	26:14:230:U:O4'	2.14	0.48
26:14:642:G:H3'	26:14:642:G:C8	2.49	0.48
35:15:33:LEU:HD12	35:15:38:HIS:CE1	2.49	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
29:19:27:THR:HG22	29:19:29:PRO:O	2.13	0.48
29:19:49:ILE:HD11	29:19:52:ARG:HA	1.96	0.48
1:1G:1008:C:N4	1:1G:1021:G:H22	2.11	0.48
1:1G:921:U:O2	5:42:19:MET:HB3	2.14	0.48
26:1H:1021:A:H3'	26:1H:1022:G:H5"	1.95	0.48
26:1H:1899:G:H1	26:1H:1902:C:N4	2.10	0.48
26:1H:2062:A:H2'	26:1H:2062:A:N3	2.29	0.48
26:1H:2322:A:OP2	62:1H:3776:HOH:O	2.20	0.48
26:1H:275:G:N2	26:1H:278:A:H61	2.11	0.48
26:1H:297:C:H2'	26:1H:298:G:O4'	2.14	0.48
30:21:8:LYS:NZ	30:21:190:GLY:O	2.39	0.48
30:29:169:ASN:OD1	30:29:201:THR:HG21	2.14	0.48
26:14:2680:C:H5'	30:29:189:PRO:HA	1.95	0.48
30:29:76:ARG:HG2	30:29:195:LEU:HD13	1.95	0.48
37:35:82:GLY:HA2	37:35:113:LYS:O	2.12	0.48
4:3E:122:ARG:HG2	4:3E:122:ARG:NH1	2.29	0.48
32:41:96:ARG:HB2	32:41:96:ARG:HH11	1.79	0.48
38:45:136:ALA:N	38:45:137:TYR:HA	2.29	0.48
5:4E:11:ILE:HG12	5:4E:31:LEU:HB3	1.96	0.48
33:51:2:SER:C	33:51:3:ARG:HE	2.16	0.48
1:13:974:A:P	14:5I:41:ARG:HH12	2.37	0.48
17:8I:45:HIS:O	17:8I:73:VAL:HG23	2.14	0.48
18:9I:73:ALA:HB3	18:9I:79:LEU:HD12	1.95	0.48
20:BI:90:GLN:HA	20:BI:93:GLU:HB2	1.94	0.48
47:D5:157:LEU:CB	47:D5:161:VAL:HG21	2.43	0.48
47:D5:75:ASN:O	47:D5:84:GLU:HG3	2.13	0.48
45:B5:11:PRO:HD3	50:G5:37:PHE:CE2	2.49	0.48
48:I8:82:ARG:NH2	48:I8:83:PRO:O	2.47	0.48
1:13:1022:G:H2'	1:13:1023:G:H8	1.79	0.48
1:13:1156:G:H2'	1:13:1157:A:H5"	1.96	0.48
1:13:1306:A:N6	1:13:1331:G:H1'	2.26	0.48
1:13:346:G:H21	1:13:347:G:H1'	1.79	0.48
1:13:351:G:H4'	1:13:352:C:OP1	2.13	0.48
1:13:688:G:H2'	1:13:689:C:C6	2.49	0.48
26:14:1048:A:OP2	26:14:1110:G:N2	2.46	0.48
26:14:194:G:H2'	26:14:195:A:O4'	2.14	0.48
27:16:70:C:H2'	27:16:71:C:H6	1.79	0.48
2:1E:70:PHE:HE1	2:1E:90:MET:HB3	1.79	0.48
1:1G:1104:G:H4'	2:12:111:ARG:HH21	1.79	0.48
1:1G:1273:G:C4	1:1G:1274:G:C8	3.02	0.48
1:1G:45:U:H2'	1:1G:46:G:H8	1.79	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:689:C:C2'	1:1G:690:G:H5'	2.44	0.48
1:1G:980:C:H3'	1:1G:981:U:C6	2.49	0.48
26:1H:1794:U:H2'	26:1H:1795:C:C6	2.49	0.48
26:1H:2210:G:H3'	26:1H:2211:G:C4	2.48	0.48
26:1H:2352:A:C4	26:1H:2366:A:C2	3.02	0.48
26:1H:2443:C:O2'	26:1H:2444:G:H5'	2.14	0.48
26:1H:2543:G:H2'	26:1H:2544:G:C8	2.48	0.48
26:1H:322:A:OP2	31:31:169:ASN:HB2	2.12	0.48
27:1J:49:C:H2'	27:1J:50:G:C8	2.49	0.48
30:21:16:ARG:O	30:21:16:ARG:HG3	2.14	0.48
3:22:135:LYS:HZ1	3:22:139:GLN:HB2	1.79	0.48
1:13:1193:G:P	3:2E:167:TRP:HZ3	2.37	0.48
23:2K:26:C:H2'	23:2K:27:G:O4'	2.14	0.48
23:2L:32:G:C6	23:2L:33:OMC:N4	2.82	0.48
4:32:173:TRP:CD1	4:32:174:LEU:HG	2.49	0.48
31:39:27:GLU:O	31:39:28:ILE:HG12	2.13	0.48
24:3K:6:G:N2	24:3K:67:C:O2	2.47	0.48
38:45:29:PHE:HB3	38:45:65:PHE:CE2	2.48	0.48
40:65:27:SER:HA	40:65:88:ASP:HB2	1.96	0.48
36:68:7:TYR:HE1	36:68:20:MET:HE3	1.79	0.48
8:7E:88:LYS:HB3	8:7E:89:PRO:HD2	1.95	0.48
18:9I:31:LEU:HD21	18:9I:62:GLU:HG2	1.96	0.48
26:14:1323:U:OP1	44:A5:84:ARG:HD2	2.13	0.48
41:B8:16:ARG:HB2	41:B8:18:ASP:OD1	2.14	0.48
42:C8:79:PHE:HE2	42:C8:106:PHE:CZ	2.32	0.48
49:J8:71:TYR:HA	49:J8:74:VAL:HG13	1.96	0.48
56:M5:52:LYS:N	56:M5:53:PRO:HD2	2.29	0.48
1:13:221:C:H2'	1:13:222:U:H6	1.79	0.47
1:13:256:U:H3	1:13:270:A:H61	1.62	0.47
1:13:321:A:H62	1:13:328:C:H1'	1.79	0.47
1:13:981:U:H5	1:13:982:U:HO2'	1.60	0.47
26:14:1827:C:C2'	26:14:1828:G:H5'	2.43	0.47
26:14:1889:A:N1	26:14:2234:G:H1'	2.29	0.47
26:14:76:C:O3'	50:G5:59:ARG:HG3	2.13	0.47
29:19:12:SER:HB2	29:19:208:LYS:HB3	1.95	0.47
2:1E:46:LYS:HA	2:1E:49:GLU:OE1	2.13	0.47
1:1G:1288:A:H2'	1:1G:1289:A:C8	2.49	0.47
26:1H:1171:G:C5	26:1H:1174:A:N6	2.82	0.47
26:1H:1332:G:C8	26:1H:1332:G:H5'	2.49	0.47
26:1H:2840:C:H2'	26:1H:2841:C:C6	2.48	0.47
26:1H:699:A:H2'	26:1H:700:G:O4'	2.13	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:722:A:H2'	26:1H:723:G:C8	2.49	0.47
26:1H:963:U:H5"	62:1H:4635:HOH:O	2.13	0.47
30:21:105:THR:HG21	30:21:164:ARG:CZ	2.44	0.47
30:29:117:MET:HA	30:29:122:PHE:N	2.28	0.47
4:32:126:ILE:HG22	4:32:127:THR:N	2.29	0.47
31:39:51:THR:HG23	31:39:92:PRO:HG2	1.95	0.47
12:3A:41:ARG:NH1	12:3A:41:ARG:HB3	2.25	0.47
24:3L:15:G:H22	24:3L:48:C:H41	1.61	0.47
32:41:161:THR:CG2	32:41:163:ALA:H	2.24	0.47
32:41:11:TYR:OH	32:41:16:ARG:NH1	2.46	0.47
32:49:49:ASP:O	32:49:52:ILE:HG22	2.14	0.47
32:49:63:ILE:HG22	32:49:143:GLU:HB2	1.96	0.47
6:52:81:ILE:HG23	6:52:82:ARG:HG3	1.95	0.47
6:5E:5:GLU:HA	6:5E:63:TYR:O	2.14	0.47
7:62:23:VAL:HG13	7:62:43:PHE:CE2	2.48	0.47
27:1J:52:A:H62	40:65:33:LYS:HG3	1.78	0.47
7:6E:150:ALA:HB2	11:2I:50:TYR:HE2	1.77	0.47
8:72:7:ALA:HB2	8:72:85:ARG:HD3	1.95	0.47
41:75:26:ASP:OD1	41:75:120:ARG:NH2	2.41	0.47
18:9A:22:VAL:C	18:9A:24:ALA:H	2.17	0.47
20:BI:49:ALA:O	20:BI:52:ALA:N	2.47	0.47
46:C5:75:ILE:HG22	46:C5:76:CYS:N	2.28	0.47
47:H8:58:VAL:O	47:H8:60:GLU:N	2.47	0.47
54:O8:44:ARG:O	54:O8:45:LYS:HG2	2.14	0.47
26:1H:764:A:H2	29:11:219:PRO:HG3	1.79	0.47
26:1H:1826:G:H4'	29:11:242:ARG:CZ	2.45	0.47
1:13:157:G:N2	1:13:165:C:O2	2.47	0.47
1:13:295:C:H2'	1:13:296:U:O4'	2.14	0.47
1:13:375:U:O3'	16:7I:6:LEU:HB2	2.13	0.47
26:14:1022:G:C6	26:14:1140:C:C4	3.03	0.47
26:14:1147:C:H2'	26:14:1148:A:C8	2.50	0.47
26:14:1188:U:O2'	26:14:1189:A:H5'	2.14	0.47
26:14:1235:G:C6	26:14:1236:G:N1	2.82	0.47
26:14:1342:A:C2	26:14:1397:U:C2	3.02	0.47
26:14:1494:A:C2	26:14:1495:A:C4	3.02	0.47
26:14:1889:A:H2'	26:14:1890:A:O4'	2.14	0.47
26:14:702:G:C2	26:14:731:C:C2	3.02	0.47
26:14:882:G:H22	26:14:894:C:N4	2.12	0.47
21:1F:3:LYS:HB3	21:1F:14:TRP:CD1	2.49	0.47
1:1G:978:A:H1'	1:1G:1322:C:C2	2.49	0.47
1:1G:1422:G:OP1	36:25:48:PRO:HA	2.14	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:485:G:O2'	1:1G:486:U:O5'	2.33	0.47
26:1H:1535:U:O4	26:1H:1538:G:O2'	2.15	0.47
26:1H:2837:G:H2'	26:1H:2838:G:H8	1.80	0.47
26:1H:320:A:OP1	31:31:135:LYS:NZ	2.47	0.47
26:1H:456:C:O2'	26:1H:457:A:OP2	2.31	0.47
3:22:141:VAL:HA	3:22:144:SER:HB3	1.96	0.47
36:25:60:ALA:HB1	36:25:84:ALA:HB1	1.96	0.47
31:39:101:LEU:O	31:39:106:ARG:NH1	2.46	0.47
31:39:107:LYS:HZ1	31:39:205:ARG:HD2	1.78	0.47
32:41:107:LEU:HD21	32:41:178:PHE:CE1	2.50	0.47
13:4I:49:THR:HG22	13:4I:51:ALA:H	1.79	0.47
26:14:1651:G:H5'	39:55:39:PRO:HG2	1.96	0.47
7:6E:94:ARG:O	7:6E:97:GLN:HB3	2.14	0.47
37:78:15:ARG:HA	37:78:15:ARG:HD2	1.62	0.47
37:78:29:LYS:HG2	37:78:30:THR:N	2.29	0.47
37:78:97:PRO:HB3	37:78:112:LEU:HB2	1.96	0.47
37:78:98:GLU:O	37:78:101:VAL:HG13	2.13	0.47
9:8E:77:ILE:O	9:8E:81:ILE:HG12	2.14	0.47
37:78:50:ARG:HD3	56:Q8:7:HIS:CD2	2.49	0.47
2:12:77:ALA:O	2:12:81:VAL:HG23	2.14	0.47
1:13:1125:U:C2	1:13:1126:U:C5	3.02	0.47
26:14:1239:G:H5"	62:14:4324:HOH:O	2.14	0.47
26:14:1331:A:O2'	26:14:1332:G:H8	1.96	0.47
26:14:1351:C:H2'	26:14:1352:U:C6	2.50	0.47
26:14:2115:G:O2'	26:14:2171:A:N6	2.47	0.47
26:14:2592:G:N7	62:14:3715:HOH:O	2.35	0.47
26:14:43:G:H2'	26:14:44:A:O4'	2.13	0.47
26:14:529:A:H4'	26:14:530:G:H5'	1.96	0.47
26:14:774:A:HO2'	26:14:775:G:P	2.37	0.47
26:14:805:G:OP2	37:35:41:ARG:HG2	2.14	0.47
2:1E:84:GLU:OE2	2:1E:216:SER:HA	2.14	0.47
1:1G:451:A:OP1	1:1G:481:G:N2	2.36	0.47
1:1G:660:G:H1	1:1G:745:C:H42	1.60	0.47
1:1G:976:G:H5'	1:1G:1358:U:O2'	2.14	0.47
26:1H:1187:G:H5"	43:D8:81:TYR:CE1	2.49	0.47
26:1H:1541:U:H2'	26:1H:1542:G:O4'	2.14	0.47
26:1H:1686:C:H2'	26:1H:1687:G:O4'	2.15	0.47
26:1H:1639:U:H4'	26:1H:2699:C:H4'	1.96	0.47
31:31:184:TYR:CE2	31:31:188:ARG:HD2	2.49	0.47
31:39:7:TYR:HD1	31:39:18:ARG:H	1.61	0.47
5:42:8:GLU:HA	5:42:34:VAL:HA	1.96	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:4I:15:VAL:O	13:4I:19:LEU:HD23	2.14	0.47
7:62:97:GLN:O	7:62:101:LEU:HG	2.14	0.47
7:62:20:ASP:HB3	7:62:23:VAL:HB	1.95	0.47
7:62:88:PRO:O	7:62:89:MET:HG2	2.14	0.47
36:68:47:ILE:HG13	36:68:48:PRO:HD2	1.97	0.47
7:6E:140:ASP:O	7:6E:144:MET:HG2	2.14	0.47
7:6E:150:ALA:HB2	11:2I:50:TYR:CE2	2.49	0.47
15:6I:56:LEU:O	15:6I:60:VAL:HG23	2.14	0.47
1:1G:826:C:H5'	8:72:12:ARG:NH1	2.28	0.47
41:75:91:ARG:HD2	41:75:124:ASP:OD2	2.14	0.47
1:13:345:C:H4'	1:13:346:G:C8	2.50	0.47
1:13:963:G:N7	62:13:1854:HOH:O	2.35	0.47
26:14:1199:U:O2'	62:14:3649:HOH:O	2.19	0.47
26:14:1536:A:H8	26:14:1537:C:H1'	1.78	0.47
26:14:1784:A:H4'	26:14:1785:A:O5'	2.15	0.47
26:14:2328:A:H2'	26:14:2329:G:O4'	2.15	0.47
26:14:2402:C:H5	26:14:2415:G:H22	1.63	0.47
26:14:49:A:H5"	26:14:51:G:O4'	2.14	0.47
26:14:819:A:H2'	26:14:820:A:H5'	1.95	0.47
26:14:959:A:C6	26:14:960:A:N1	2.82	0.47
29:19:8:PRO:HB3	29:19:14:ARG:HB2	1.96	0.47
1:1G:999:U:H2'	1:1G:1000:A:C8	2.50	0.47
1:1G:382:A:H2'	1:1G:383:A:C8	2.50	0.47
1:1G:487:A:H2'	1:1G:488:C:O4'	2.15	0.47
1:1G:858:G:H8	1:1G:858:G:OP2	1.97	0.47
26:1H:2330:G:H2'	26:1H:2331:G:O4'	2.14	0.47
26:1H:444:C:C4'	31:31:49:ALA:HB2	2.44	0.47
26:1H:782:A:H5'	26:1H:783:A:C2	2.49	0.47
22:1K:7:U:O2'	22:1K:8:U:H5'	2.15	0.47
3:22:113:ALA:HB3	3:22:114:PRO:HD3	1.96	0.47
31:39:121:GLY:O	31:39:122:LYS:HD3	2.15	0.47
24:3L:18:G:O2'	24:3L:57:G:H2'	2.14	0.47
32:49:95:ARG:O	32:49:99:MET:HG2	2.14	0.47
33:59:59:ARG:O	33:59:63:SER:OG	2.20	0.47
15:6A:39:LEU:CD1	15:6A:56:LEU:HB2	2.43	0.47
28:71:6:ARG:CZ	28:71:6:ARG:HB3	2.43	0.47
8:7E:104:ARG:HG3	8:7E:138:TRP:CD1	2.50	0.47
9:8E:97:LYS:HB2	9:8E:102:LEU:HD12	1.97	0.47
39:98:26:LYS:HE2	39:98:70:LEU:O	2.14	0.47
41:B8:88:ILE:O	41:B8:88:ILE:HG13	2.13	0.47
47:D5:4:ARG:NH1	47:D5:60:GLU:OE2	2.46	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:E5:72:ARG:HH21	48:E5:75:LEU:CD1	2.28	0.47
45:F8:41:ASN:OD1	45:F8:41:ASN:N	2.47	0.47
50:G5:17:SER:HB2	50:G5:20:GLU:HG3	1.95	0.47
2:12:188:ALA:O	2:12:203:GLY:N	2.48	0.47
1:13:131:C:H2'	1:13:132:C:C6	2.49	0.47
1:13:1375:A:H4'	7:6E:29:LYS:HD3	1.97	0.47
1:13:1466:C:H2'	1:13:1467:G:O4'	2.15	0.47
1:13:475:G:H2'	1:13:476:G:O4'	2.14	0.47
1:13:947:G:O3'	13:4I:109:THR:OG1	2.32	0.47
26:14:1033:U:H3'	26:14:1033:U:H6	1.79	0.47
26:14:2865:U:C4	26:14:2866:U:C4	3.02	0.47
26:14:332:A:O2'	26:14:334:C:OP2	2.23	0.47
26:14:853:G:O2'	26:14:854:G:H5'	2.15	0.47
27:16:29:A:OP2	40:A8:31:SER:HB2	2.13	0.47
21:1B:2:GLY:O	21:1B:4:GLY:N	2.47	0.47
1:1G:991:U:C5	1:1G:1212:U:H1'	2.48	0.47
1:1G:1281:U:H3'	1:1G:1282:C:C5	2.49	0.47
1:1G:281:G:H8	1:1G:281:G:OP2	1.96	0.47
26:1H:941:A:H4'	62:1H:4465:HOH:O	2.13	0.47
3:22:5:ILE:HG21	10:1A:51:ARG:HH21	1.80	0.47
30:29:23:VAL:HG11	30:29:183:LEU:HD23	1.95	0.47
4:32:65:ARG:HD2	4:32:72:GLU:HA	1.96	0.47
24:3K:9:A:H1'	24:3K:46:G:C8	2.49	0.47
5:4E:27:ARG:HE	5:4E:27:ARG:HB2	1.39	0.47
33:59:19:VAL:HG12	33:59:20:ALA:H	1.79	0.47
34:69:128:LEU:O	34:69:137:PRO:HA	2.14	0.47
8:72:119:LEU:HD12	8:72:124:ALA:HA	1.95	0.47
26:1H:1250:G:OP2	37:78:21:ARG:HD3	2.15	0.47
37:78:59:LEU:HB2	56:Q8:58:ILE:CD1	2.44	0.47
37:78:59:LEU:HB2	56:Q8:58:ILE:HD11	1.96	0.47
8:7E:83:ILE:HB	8:7E:137:VAL:HG13	1.96	0.47
1:13:392:G:H5"	16:7I:12:LYS:HE3	1.96	0.47
9:8E:17:VAL:HG21	9:8E:80:GLY:HA3	1.96	0.47
9:8E:9:ARG:HE	9:8E:14:VAL:HG13	1.79	0.47
17:8I:28:PRO:HA	17:8I:34:LYS:O	2.14	0.47
40:A8:26:LEU:HD22	40:A8:87:PHE:CD1	2.50	0.47
46:C5:97:ARG:NH1	46:C5:104:GLY:O	2.47	0.47
47:D5:105:VAL:HG13	47:D5:106:GLY:N	2.30	0.47
1:13:1028(A):C:N4	1:13:1029:G:N7	2.62	0.47
1:13:1162:C:H2'	1:13:1163:C:C6	2.49	0.47
1:13:1322:C:H5"	13:4I:100:GLY:HA2	1.95	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:142:G:H2'	26:14:143:C:C6	2.50	0.47
26:14:2124:G:H2'	26:14:2124:G:N3	2.29	0.47
26:14:2688:U:H5	26:14:2720:U:OP2	1.97	0.47
1:1G:411:A:C5	1:1G:413:G:H1'	2.49	0.47
1:1G:804:U:H5"	1:1G:805:C:OP2	2.15	0.47
26:1H:1678:G:N2	26:1H:1989:G:N2	2.60	0.47
26:1H:185:U:H4'	26:1H:218:A:H4'	1.97	0.47
26:1H:2436:G:C5	26:1H:2437:U:C5	3.03	0.47
26:1H:2775:A:N6	62:1H:3724:HOH:O	2.48	0.47
26:1H:2877:G:H2'	26:1H:2878:U:O4'	2.14	0.47
26:1H:775:G:O5'	26:1H:777:A:H1'	2.14	0.47
26:1H:847:U:H5'	62:1H:3743:HOH:O	2.15	0.47
27:1J:21:G:H2'	27:1J:22:U:O4'	2.15	0.47
23:2K:65:G:C2	23:2K:66:C:C2	3.03	0.47
4:32:24:GLU:OE2	4:32:24:GLU:N	2.48	0.47
37:35:6:LEU:HA	37:35:6:LEU:HD12	1.52	0.47
1:13:881:G:P	12:3I:12:ARG:HH22	2.37	0.47
24:3L:44:U:H2'	24:3L:45:G:O4'	2.14	0.47
13:4I:70:LEU:O	13:4I:74:VAL:HG23	2.15	0.47
33:59:137:ASP:CB	33:59:140:LYS:HB2	2.45	0.47
6:5E:96:PRO:HB3	18:9I:30:ASP:CG	2.35	0.47
36:68:64:ARG:HB2	36:68:79:PHE:CG	2.49	0.47
7:6E:111:ARG:HE	7:6E:123:GLU:HB2	1.79	0.47
9:8E:112:LYS:CA	9:8E:119:ALA:HB2	2.43	0.47
40:A8:58:LEU:HD12	40:A8:68:GLN:OE1	2.15	0.47
19:AA:53:ASN:HB2	19:AA:77:THR:HG22	1.97	0.47
29:11:79:VAL:HG21	29:11:111:LEU:HD11	1.97	0.47
29:11:77:ALA:HB2	29:11:97:TYR:CD2	2.49	0.47
1:13:1164:G:C6	1:13:1165:C:C4	3.03	0.47
1:13:1427:U:H2'	1:13:1428:A:C8	2.48	0.47
1:13:41:G:H2'	1:13:42:G:C8	2.49	0.47
1:13:452:A:H2'	1:13:453:A:C8	2.50	0.47
1:13:631:G:HO2'	1:13:632:A:P	2.37	0.47
26:14:1621:U:OP1	62:14:3652:HOH:O	2.20	0.47
26:14:631:A:H2'	26:14:632:A:O4'	2.15	0.47
27:16:15:A:H5'	27:16:16:G:H8	1.80	0.47
1:1G:855:G:C2	1:1G:856:C:C2	3.02	0.47
26:1H:1053:C:N4	26:1H:1106:G:H1	2.13	0.47
26:1H:1688:U:O2	26:1H:1700:A:H5"	2.14	0.47
26:1H:1971:A:C4	29:11:241:PRO:HD3	2.49	0.47
26:1H:2061:G:H5"	26:1H:2503:A:C2	2.50	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2648:C:H2'	26:1H:2649:U:C6	2.49	0.47
26:1H:32:C:O2'	26:1H:33:U:H5'	2.15	0.47
1:1G:1112:C:C4	3:22:178:LEU:HD23	2.49	0.47
30:29:76:ARG:HD3	30:29:76:ARG:HA	1.59	0.47
30:29:9:VAL:HA	41:75:3:ARG:HD2	1.97	0.47
31:39:30:PRO:O	31:39:33:LEU:N	2.47	0.47
5:4E:98:THR:HB	5:4E:117:ASP:HB3	1.97	0.47
25:4L:19:G:N3	25:4L:20:A:C8	2.83	0.47
33:51:97:ARG:NH2	33:51:104:GLU:OE2	2.42	0.47
33:51:5:GLY:HA2	33:51:8:PRO:HD3	1.97	0.47
26:14:1653:G:H3'	39:55:2:ARG:CG	2.43	0.47
34:61:33:ARG:HB3	34:61:35:LEU:HD13	1.97	0.47
34:69:143:SER:O	34:69:144:VAL:HG22	2.14	0.47
28:71:23:ASP:HB2	28:71:190:ARG:NH2	2.26	0.47
16:7A:52:ASP:OD2	16:7A:55:ARG:HG3	2.15	0.47
16:7I:20:VAL:HG21	16:7I:32:TYR:CD2	2.50	0.47
9:82:81:ILE:HG22	9:82:85:LEU:HD23	1.96	0.47
40:A8:58:LEU:HD23	40:A8:58:LEU:H	1.80	0.47
20:BA:87:LYS:O	20:BA:91:LEU:HG	2.15	0.47
20:BI:40:ALA:HB2	20:BI:55:ILE:HG22	1.97	0.47
26:14:2261:C:C6	48:E5:16:SER:HB3	2.49	0.47
46:G8:96:ILE:HA	46:G8:102:CYS:O	2.14	0.47
47:H8:116:VAL:HG13	47:H8:146:ILE:HD13	1.97	0.47
27:16:43:C:H5"	52:M8:1:MET:HG2	1.96	0.47
29:11:92:ILE:HD12	29:11:104:TYR:CE1	2.49	0.47
1:13:1120:G:H2'	1:13:1121:U:C6	2.50	0.47
1:13:1391:U:H2'	1:13:1392:G:H8	1.72	0.47
1:13:553:A:H5"	12:3I:24:VAL:HG21	1.97	0.47
1:13:737:A:H2'	1:13:738:C:H6	1.74	0.47
26:14:1445:C:H2'	26:14:1446:C:C6	2.49	0.47
26:14:1551:C:H2'	26:14:1552:G:O4'	2.15	0.47
26:14:1579:A:H2'	26:14:1580:A:C8	2.49	0.47
26:14:1826:G:H4'	29:19:242:ARG:CZ	2.45	0.47
26:14:1833:U:O2'	26:14:1969:A:N1	2.38	0.47
26:14:202:U:H2'	26:14:203:C:O4'	2.14	0.47
26:14:2182:G:H2'	26:14:2183:C:C6	2.50	0.47
26:14:2104:G:C2	26:14:2186:G:C2	3.03	0.47
26:14:2400:G:H2'	26:14:2401:U:H6	1.79	0.47
26:14:2591:C:H2'	26:14:2592:G:C8	2.50	0.47
26:14:582:G:H2'	26:14:583:G:H8	1.79	0.47
35:15:61:ARG:NH1	35:15:61:ARG:HA	2.29	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1152:A:H2'	1:1G:1153:C:C6	2.49	0.47
1:1G:690:G:H22	11:2A:55:LYS:HE2	1.79	0.47
1:1G:843:U:H3'	1:1G:848:C:O4'	2.13	0.47
26:1H:139:G:N3	26:1H:141:A:N1	2.63	0.47
26:1H:1771:C:H1'	26:1H:1786:A:C8	2.50	0.47
26:1H:1799:G:H5'	26:1H:1819:A:N6	2.29	0.47
26:1H:2439:A:C8	26:1H:2439:A:H5'	2.49	0.47
26:1H:270(J):G:N2	26:1H:270(P):C:O2	2.30	0.47
26:1H:482:A:H5"	26:1H:483:A:OP1	2.15	0.47
26:1H:806:C:OP2	37:78:41:ARG:HD3	2.15	0.47
4:32:112:VAL:HG12	4:32:116:GLN:OE1	2.14	0.47
37:35:113:LYS:HD3	37:35:115:LEU:HD21	1.96	0.47
37:35:27:HIS:HB3	37:35:32:THR:HG23	1.96	0.47
4:3E:108:LEU:HB3	4:3E:110:PHE:CD1	2.50	0.47
6:52:33:TYR:HE1	6:52:78:GLU:HG3	1.80	0.47
39:55:51:LEU:HD22	39:55:66:VAL:HG13	1.97	0.47
26:14:2839:G:H21	39:55:92:GLY:CA	2.28	0.47
3:22:18:TRP:HE1	14:5A:56:VAL:H	1.62	0.47
8:72:121:ASP:OD1	8:72:125:ARG:NH2	2.47	0.47
26:1H:249:C:O2'	37:78:64:LYS:HE2	2.15	0.47
37:78:96:THR:C	37:78:98:GLU:H	2.18	0.47
9:82:37:PHE:HB3	9:82:43:ALA:CB	2.44	0.47
38:88:34:LEU:HD11	38:88:129:THR:OG1	2.14	0.47
39:98:2:ARG:O	39:98:5:LYS:HB2	2.15	0.47
41:B8:12:SER:HA	41:B8:14:TYR:N	2.26	0.47
29:11:69:ARG:NH2	29:11:128:GLY:O	2.28	0.47
1:13:1210:C:C2'	1:13:1211:U:H5'	2.44	0.47
1:13:1422:G:H5"	36:68:48:PRO:CB	2.41	0.47
1:13:1:U:C6	1:13:630:G:H2'	2.48	0.47
26:14:1570:A:H2'	26:14:1571:A:C8	2.49	0.47
26:14:2019:A:OP2	53:J5:9:LYS:NZ	2.44	0.47
26:14:276:A:H2'	26:14:277:C:C5	2.50	0.47
26:14:278:A:HO2'	26:14:279:C:H5	1.63	0.47
26:14:384:U:H2'	26:14:385:C:H6	1.79	0.47
1:1G:1028:C:N4	1:1G:1034:G:H21	2.12	0.47
1:1G:1001:G:N2	1:1G:1040:U:O2	2.48	0.47
1:1G:1095:U:OP1	1:1G:1108:G:N1	2.47	0.47
26:1H:106:C:H2'	26:1H:107:C:C6	2.50	0.47
26:1H:1534:G:N2	26:1H:1535:U:H5	2.13	0.47
26:1H:214:G:OP1	26:1H:214:G:H4'	2.14	0.47
26:1H:2285:C:H5	54:O8:27:LYS:HD2	1.80	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2453:A:H2'	26:1H:2454:G:O4'	2.15	0.47
26:1H:2728:U:H2'	26:1H:2729:G:C8	2.49	0.47
26:1H:818:G:H5'	26:1H:839:U:OP1	2.15	0.47
22:1K:7:U:O2'	22:1K:49:G:N2	2.48	0.47
31:31:155:LEU:HB2	31:31:189:THR:HG21	1.97	0.47
4:32:96:LEU:HD22	4:32:139:ARG:NH1	2.30	0.47
32:41:11:TYR:HA	32:41:15:VAL:HB	1.97	0.47
32:41:18:GLU:O	32:41:22:ARG:HB2	2.14	0.47
32:49:99:MET:HE2	32:49:99:MET:HB2	1.66	0.47
13:4I:67:GLU:HG2	13:4I:71:ARG:NH2	2.30	0.47
7:62:47:CYS:O	7:62:50:ILE:HB	2.15	0.47
28:71:171:ILE:HA	28:71:171:ILE:HD12	1.81	0.47
37:78:68:GLN:HG3	56:Q8:12:LYS:HD3	1.95	0.47
37:78:83:VAL:O	37:78:114:ILE:HA	2.15	0.47
8:7E:113:SER:HB3	8:7E:134:ILE:HD11	1.96	0.47
17:8A:56:VAL:O	17:8A:77:VAL:N	2.43	0.47
19:AI:5:LEU:HA	19:AI:5:LEU:HD12	1.80	0.47
20:BI:38:LYS:HE2	20:BI:38:LYS:HB3	1.73	0.47
44:E8:71:VAL:HA	44:E8:107:LEU:HD12	1.95	0.47
44:E8:96:ILE:HD13	44:E8:96:ILE:H	1.79	0.47
49:J8:7:ILE:HG13	49:J8:62:VAL:HG12	1.97	0.47
37:35:59:LEU:HD22	56:M5:13:ARG:HD2	1.96	0.47
52:M8:16:CYS:HB3	52:M8:36:CYS:H	1.80	0.47
1:13:342:C:C2	1:13:348:G:C2	3.03	0.47
1:13:439:A:H3'	1:13:440:A:H8	1.79	0.47
26:14:1259:G:H2'	26:14:1260:G:C8	2.50	0.47
26:14:2074:U:H2'	26:14:2075:U:C6	2.50	0.47
26:14:2352:A:C2	48:E5:33:ALA:HB1	2.50	0.47
26:14:2690:C:OP2	39:55:14:SER:HB2	2.14	0.47
26:14:276:A:N3	26:14:277:C:N4	2.63	0.47
26:14:2795:G:N3	26:14:2795:G:H2'	2.30	0.47
26:14:2850:A:C2	26:14:2851:A:C4	3.03	0.47
26:14:68:G:H2'	26:14:69:C:O4'	2.15	0.47
26:14:870:A:H5"	38:45:6:ARG:HB3	1.97	0.47
2:1E:100:GLY:N	2:1E:176:GLU:OE2	2.47	0.47
2:1E:68:ILE:O	2:1E:91:PRO:HD2	2.14	0.47
1:1G:134:A:H1'	1:1G:325:A:C5	2.50	0.47
1:1G:1410:G:H2'	1:1G:1411:C:C6	2.50	0.47
1:1G:1482:G:O6	62:1G:1863:HOH:O	2.20	0.47
1:1G:298:A:H5"	1:1G:299:G:OP2	2.14	0.47
26:1H:1668:A:C8	26:1H:1674:G:C6	3.03	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1268:A:C2	26:1H:2013:A:C4	3.03	0.47
26:1H:2130:U:H2'	26:1H:2131:G:N7	2.30	0.47
26:1H:2210:G:H3'	26:1H:2211:G:C5	2.50	0.47
26:1H:2436:G:C6	26:1H:2437:U:C4	3.03	0.47
26:1H:275:G:N2	26:1H:276:A:N7	2.42	0.47
26:1H:602:G:N2	26:1H:655:A:C8	2.70	0.47
22:1K:45:G:O2'	22:1K:47:U:H5'	2.15	0.47
36:25:10:VAL:HG23	36:25:12:ASP:OD1	2.15	0.47
30:29:9:VAL:HG23	30:29:26:ILE:O	2.15	0.47
12:3I:70:ILE:HD13	12:3I:77:LEU:HD12	1.97	0.47
32:49:33:ARG:NH2	32:49:162:THR:HG21	2.30	0.47
13:4A:39:ILE:HG23	13:4A:52:GLU:HG2	1.97	0.47
13:4A:59:TYR:CD2	13:4A:60:VAL:HG22	2.50	0.47
39:55:79:LEU:HA	39:55:83:ILE:HB	1.95	0.47
35:58:35:ARG:O	35:58:42:TRP:HZ3	1.97	0.47
33:59:107:VAL:CG1	33:59:152:ARG:HG2	2.45	0.47
33:59:152:ARG:HD2	33:59:153:LYS:HG3	1.96	0.47
7:62:22:LEU:HD23	7:62:62:PHE:CE2	2.48	0.47
42:85:92:ARG:O	42:85:94:ASN:N	2.48	0.47
9:8E:86:VAL:HG11	9:8E:93:ARG:HG3	1.97	0.47
17:8I:31:LEU:HD22	17:8I:32:TYR:CE1	2.50	0.47
49:F5:92:LYS:O	49:F5:94:LEU:N	2.48	0.47
2:12:42:ILE:HG21	2:12:202:PRO:HB2	1.97	0.47
1:13:1318:A:H1'	19:AI:37:ARG:HH21	1.79	0.47
1:13:1432:G:OP1	41:B8:107:ASP:HB2	2.14	0.47
26:14:1639:U:C2'	26:14:1640:C:H5'	2.45	0.47
26:14:1665:A:H2'	26:14:1666:G:O4'	2.14	0.47
26:14:195:A:H4'	26:14:251:A:O2'	2.15	0.47
26:14:2846:G:H2'	26:14:2847:U:O4'	2.14	0.47
26:14:480:A:N3	26:14:480:A:H2'	2.30	0.47
26:14:57:C:H2'	26:14:58:G:O4'	2.15	0.47
35:15:35:ARG:HB3	35:15:42:TRP:CZ3	2.50	0.47
1:1G:1072:G:H2'	1:1G:1073:U:O4'	2.15	0.47
1:1G:1069:C:O2'	1:1G:1192:C:H1'	2.15	0.47
1:1G:1352:C:N3	1:1G:1370:G:N2	2.51	0.47
1:1G:577:G:H2'	1:1G:578:C:H6	1.80	0.47
1:1G:922:G:O5'	5:42:20:GLN:NE2	2.43	0.47
26:1H:270(H):C:H2'	26:1H:270(I):G:C8	2.50	0.47
26:1H:627:A:H4'	26:1H:628:G:OP1	2.15	0.47
26:1H:739:G:H8	26:1H:739:G:OP2	1.97	0.47
4:3E:101:LEU:HD11	4:3E:126:ILE:HG21	1.97	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:3L:72:C:C3'	24:3L:73:A:H5"	2.45	0.47
33:59:97:ARG:HG2	33:59:98:LEU:H	1.79	0.47
36:68:35:VAL:HG21	36:68:103:ALA:HB3	1.97	0.47
26:1H:1952:A:C2	36:68:22:ILE:HG23	2.50	0.47
1:1G:750:G:N2	15:6A:23:GLY:HA3	2.30	0.47
28:71:64:LEU:HD22	28:71:175:VAL:O	2.15	0.47
1:13:377:G:H5'	16:7I:5:ARG:HH12	1.80	0.47
16:7I:73:LEU:O	16:7I:77:ALA:HB2	2.15	0.47
38:88:27:VAL:HA	38:88:105:GLU:OE1	2.15	0.47
19:AA:56:GLN:HG2	19:AA:57:HIS:H	1.79	0.47
20:BA:51:GLU:HA	20:BA:54:LYS:HE3	1.96	0.47
26:1H:1816:G:H8	29:11:62:TYR:CZ	2.34	0.46
1:13:44:G:C2	1:13:45:U:H1'	2.50	0.46
1:13:520:A:N1	1:13:536:C:H1'	2.30	0.46
1:13:963:G:N2	10:1I:55:LYS:HZ3	2.13	0.46
26:14:1379:A:H1'	26:14:1380:G:OP1	2.15	0.46
26:14:1790:C:H2'	26:14:1791:A:C5	2.50	0.46
26:14:2859:G:H3'	26:14:2859:G:C8	2.51	0.46
26:14:817:C:H3'	26:14:818:G:H8	1.80	0.46
29:19:173:VAL:HG12	29:19:185:VAL:O	2.15	0.46
1:1G:973:G:C5'	10:1A:55:LYS:HZ2	2.27	0.46
2:1E:59:GLU:HB2	2:1E:221:LEU:HD11	1.97	0.46
1:1G:1286:A:H3'	1:1G:1286:A:C8	2.50	0.46
26:1H:1379:A:H1'	26:1H:1380:G:OP1	2.14	0.46
26:1H:353:G:H2'	26:1H:354:G:C8	2.49	0.46
27:1J:117:G:H8	27:1J:117:G:O5'	1.97	0.46
27:1J:21:G:H1	27:1J:62:C:H42	1.63	0.46
24:1L:8:U:H3'	24:1L:13:C:N4	2.30	0.46
23:2L:9:G:O2'	23:2L:10:G:N7	2.40	0.46
31:31:150:GLY:HA2	31:31:172:TRP:CD2	2.50	0.46
31:31:20:LEU:HD12	31:31:21:ALA:H	1.79	0.46
4:32:162:LEU:HD23	4:32:162:LEU:HA	1.57	0.46
26:14:588:U:H1'	31:39:90:PHE:CG	2.49	0.46
1:1G:529:G:O6	12:3A:49:ASN:HA	2.15	0.46
13:4I:117:VAL:HG13	13:4I:118:ALA:H	1.79	0.46
1:1G:750:G:C2	15:6A:23:GLY:HA3	2.50	0.46
8:7E:77:GLU:HG2	8:7E:78:GLN:H	1.79	0.46
16:7I:68:ASP:O	16:7I:71:ARG:HG2	2.15	0.46
38:88:16:ARG:HB3	38:88:16:ARG:HE	1.46	0.46
19:AA:9:VAL:CB	19:AA:10:PHE:HA	2.45	0.46
19:AI:40:ILE:HG21	19:AI:66:MET:O	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
43:D8:48:GLY:O	43:D8:49:THR:O	2.33	0.46
48:E5:49:LYS:HB2	48:E5:82:ARG:HH21	1.79	0.46
49:F5:3:LYS:HA	49:F5:46:LEU:HD22	1.96	0.46
48:I8:84:LEU:HD12	48:I8:84:LEU:HA	1.79	0.46
56:Q8:6:THR:HG22	56:Q8:62:LEU:HA	1.97	0.46
1:13:1316:G:N2	1:13:1318:A:H3'	2.30	0.46
1:13:1349:A:H2'	1:13:1350:A:C8	2.49	0.46
1:13:232:G:H1'	1:13:262:A:N1	2.29	0.46
1:13:729:A:H2'	1:13:730:G:H8	1.79	0.46
26:14:1771:C:H1'	26:14:1786:A:C8	2.51	0.46
26:14:2745:C:H2'	26:14:2746:U:O4'	2.16	0.46
26:14:329:G:OP2	46:C5:71:LYS:HE3	2.16	0.46
26:14:854:G:H2'	26:14:855:G:H8	1.81	0.46
27:16:41:U:C5	32:41:70:VAL:HG13	2.50	0.46
29:19:49:ILE:CD1	29:19:52:ARG:HA	2.45	0.46
1:1G:1023:G:H3'	1:1G:1024:G:O4'	2.15	0.46
1:1G:1095:U:OP1	1:1G:1108:G:N2	2.48	0.46
1:1G:1145:C:H5"	1:1G:1146:A:OP1	2.15	0.46
1:1G:131:C:H2'	1:1G:132:C:C6	2.51	0.46
1:1G:401:C:N4	62:1G:1898:HOH:O	2.48	0.46
1:1G:420:U:O2'	1:1G:423:G:O6	2.26	0.46
1:1G:538:G:H2'	1:1G:539:A:H8	1.80	0.46
1:1G:560:U:OP2	62:1G:1864:HOH:O	2.20	0.46
26:1H:1899:G:H22	26:1H:1902:C:N4	2.12	0.46
26:1H:2273:A:H2'	26:1H:2274:A:C8	2.49	0.46
26:1H:2402:C:H1'	26:1H:2403:C:H5	1.80	0.46
26:1H:662:G:O2'	26:1H:663:G:H5'	2.16	0.46
30:21:111:ARG:HG3	30:21:160:TYR:CD2	2.51	0.46
31:31:23:ASP:CG	31:31:24:LEU:H	2.18	0.46
24:3K:65:C:H2'	24:3K:66:A:H5"	1.97	0.46
38:45:42:ILE:HD13	38:45:97:VAL:CG2	2.45	0.46
32:49:47:LYS:HG2	32:49:48:GLU:H	1.80	0.46
32:49:66:GLN:NE2	32:49:94:LEU:HD23	2.31	0.46
35:58:134:ARG:HB2	35:58:134:ARG:HE	1.49	0.46
33:59:137:ASP:HB3	33:59:140:LYS:HB2	1.96	0.46
34:61:18:VAL:HG21	34:61:44:LEU:HD11	1.96	0.46
7:62:95:ARG:HH21	7:62:99:LEU:HD11	1.80	0.46
40:65:15:ARG:HD2	40:65:25:ARG:HH21	1.80	0.46
15:6A:12:ILE:HG12	15:6A:31:LEU:HD11	1.97	0.46
37:78:14:LYS:O	37:78:15:ARG:HB2	2.15	0.46
8:7E:116:LYS:HG3	8:7E:129:VAL:HG11	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:112:G:OP2	16:7I:27:LYS:HD2	2.15	0.46
9:8E:93:ARG:NH2	9:8E:97:LYS:HD2	2.30	0.46
46:G8:94:LYS:HZ2	46:G8:95:LYS:N	2.12	0.46
47:H8:132:ASN:ND2	47:H8:160:GLY:HA3	2.31	0.46
54:O8:27:LYS:NZ	54:O8:27:LYS:H	2.12	0.46
2:12:75:LYS:H	2:12:78:GLN:HG3	1.81	0.46
2:12:82:ARG:HB2	2:12:94:ASN:OD1	2.15	0.46
1:13:1058:G:C6	1:13:1059:C:N3	2.84	0.46
1:13:1092:A:C6	1:13:1093:A:C6	3.03	0.46
1:13:1203:C:H2'	1:13:1204:A:O4'	2.14	0.46
1:13:575:G:H4'	1:13:576:G:O5'	2.15	0.46
1:13:948:C:O2'	1:13:949:A:H5'	2.15	0.46
26:14:1485:G:H2'	26:14:1486:A:C8	2.51	0.46
26:14:1754:C:H2'	26:14:1755:A:C8	2.49	0.46
26:14:196:A:H2'	26:14:196:A:N3	2.31	0.46
26:14:270(P):C:H2'	26:14:270(Q):C:C6	2.50	0.46
27:16:13:A:O2'	27:16:14:U:H3'	2.16	0.46
1:1G:1348:U:H3	1:1G:1374:A:H2	1.56	0.46
1:1G:735:C:H2'	1:1G:736:C:H6	1.79	0.46
26:1H:1453:A:O2'	26:1H:1454:U:H2'	2.14	0.46
26:1H:1478:G:H2'	26:1H:1479:G:H8	1.80	0.46
26:1H:1680:U:H2'	26:1H:1681:G:O4'	2.15	0.46
26:1H:2343:C:HO2'	26:1H:2373:G:HO2'	1.57	0.46
26:1H:2503:A:OP2	26:1H:2503:A:H3'	2.15	0.46
26:1H:2575:C:H5'	30:21:144:ARG:HB2	1.97	0.46
26:1H:445:C:O2'	26:1H:446:G:H5'	2.15	0.46
27:1J:73:A:C4	27:1J:104:A:C2	3.04	0.46
31:39:146:ALA:CB	31:39:148:LEU:HG	2.44	0.46
31:39:78:ILE:HA	31:39:83:PHE:CD2	2.50	0.46
31:39:49:ALA:O	31:39:92:PRO:HB2	2.15	0.46
4:3E:15:GLU:HG3	4:3E:63:LYS:HB2	1.97	0.46
40:65:110:LEU:HD13	40:65:112:PHE:CE1	2.50	0.46
40:65:27:SER:HA	40:65:88:ASP:CB	2.45	0.46
15:6A:84:LYS:HA	15:6A:84:LYS:HD3	1.63	0.46
7:6E:79:ARG:HE	7:6E:84:ASN:HD22	1.64	0.46
26:1H:2414:G:H21	37:78:67:MET:CE	2.29	0.46
8:7E:25:ASP:OD2	8:7E:60:ARG:HG3	2.15	0.46
16:7I:12:LYS:O	16:7I:13:HIS:HB2	2.14	0.46
42:85:50:ARG:NH1	43:95:72:VAL:HG23	2.31	0.46
1:13:237:C:H5"	17:8I:25:ARG:CZ	2.46	0.46
45:B5:1:MET:HA	45:B5:2:LYS:HA	1.50	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
46:C5:52:SER:HA	46:C5:55:TYR:O	2.16	0.46
26:1H:533:G:H5'	42:C8:24:TYR:CE1	2.50	0.46
42:C8:88:ILE:O	42:C8:88:ILE:HG22	2.15	0.46
26:14:180:G:OP2	55:L5:32:LYS:HD2	2.15	0.46
52:M8:42:PHE:CD1	52:M8:43:TYR:HB3	2.50	0.46
56:Q8:4:MET:HB2	56:Q8:4:MET:HE2	1.59	0.46
29:11:85:ASP:OD2	29:11:88:ARG:NH1	2.39	0.46
2:12:92:TYR:HH	2:12:150:SER:HG	1.64	0.46
1:13:1044:A:C5	1:13:1045:C:H1'	2.51	0.46
1:13:1113:C:H2'	1:13:1114:C:C6	2.51	0.46
1:13:1132:C:H2'	1:13:1133:G:C8	2.50	0.46
1:13:1263:C:H2'	1:13:1264:C:H6	1.80	0.46
1:13:1292:U:H2'	1:13:1293:G:H8	1.77	0.46
1:13:258:G:H2'	1:13:259:G:H8	1.79	0.46
1:13:443:C:N4	1:13:491:G:H1	2.12	0.46
26:14:2212:A:O2'	26:14:2213:U:O5'	2.34	0.46
26:14:2557:G:H2'	26:14:2558:C:H6	1.81	0.46
26:14:2720:U:N3	26:14:2873:A:H2	2.08	0.46
27:16:11:C:H3'	27:16:12:C:H6	1.80	0.46
1:1G:1502:A:H2	1:1G:1505:G:N1	2.02	0.46
1:1G:337:C:H2'	1:1G:338:A:C8	2.51	0.46
26:1H:1991:U:C2'	26:1H:1992:G:H5"	2.46	0.46
26:1H:2126:A:OP1	28:71:38:ASP:HB3	2.14	0.46
26:1H:2150:U:H2'	26:1H:2151:G:C8	2.50	0.46
26:1H:2171:A:O2'	26:1H:2172:U:O4'	2.23	0.46
26:1H:2186:G:H2'	26:1H:2187:G:C8	2.50	0.46
26:1H:2400:G:O2'	26:1H:2401:U:H5'	2.16	0.46
26:1H:270(I):G:H1	26:1H:270(Q):C:N4	2.05	0.46
26:1H:403:U:H3'	62:1H:4080:HOH:O	2.15	0.46
26:1H:530:G:C5	26:1H:2022:U:H5"	2.50	0.46
26:14:2562:U:H1'	36:25:23:ARG:NE	2.31	0.46
3:2E:16:ARG:HD2	3:2E:54:ARG:NH2	2.23	0.46
4:3E:104:VAL:HG21	4:3E:140:VAL:HG21	1.97	0.46
5:42:148:VAL:O	5:42:152:ARG:HG3	2.15	0.46
32:49:111:LEU:HB2	32:49:112:PRO:HD3	1.98	0.46
5:4E:76:ILE:HG13	5:4E:93:PRO:HG3	1.98	0.46
7:62:102:ARG:HG2	7:62:106:GLN:NE2	2.30	0.46
1:13:1292:U:P	7:6E:41:ARG:HH22	2.38	0.46
37:78:124:LYS:HA	37:78:143:GLY:O	2.14	0.46
1:13:878:G:H5'	8:7E:89:PRO:HG2	1.98	0.46
9:82:18:PHE:HD2	9:82:62:TYR:HD2	1.63	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2250:G:C5	38:88:83:MET:HB3	2.51	0.46
19:AA:66:MET:N	19:AA:67:VAL:HB	2.29	0.46
41:B8:107:ASP:O	41:B8:110:ILE:HG23	2.16	0.46
46:G8:5:MET:HE1	46:G8:32:PRO:HA	1.98	0.46
51:H5:5:LYS:HG2	51:H5:36:VAL:HG22	1.96	0.46
26:14:2046:G:H5'	53:J5:19:ARG:HG3	1.98	0.46
50:K8:3:LEU:O	50:K8:6:VAL:HB	2.15	0.46
53:N8:16:ARG:HG3	53:N8:17:ASP:N	2.30	0.46
54:O8:27:LYS:HB2	54:O8:27:LYS:HZ2	1.80	0.46
1:13:1238:A:N3	1:13:1241:G:O2'	2.39	0.46
1:13:1262:C:H2'	1:13:1263:C:C6	2.49	0.46
1:13:1386:G:O2'	1:13:1387:G:H5'	2.16	0.46
26:14:1901:A:OP2	29:19:255:LYS:HE2	2.14	0.46
26:14:2104:G:H2'	26:14:2105:C:C6	2.51	0.46
26:14:2168:G:H3'	26:14:2168:G:N3	2.31	0.46
35:15:15:LEU:HD22	35:15:53:VAL:HB	1.96	0.46
27:16:44:G:C2	27:16:48:A:C2	3.04	0.46
10:1A:63:PHE:CD1	14:5A:58:LYS:HA	2.50	0.46
2:1E:174:VAL:HG13	2:1E:184:VAL:HG11	1.97	0.46
1:1G:1305:G:N2	1:1G:1331:G:O2'	2.48	0.46
1:1G:21:G:H2'	1:1G:22:G:C8	2.51	0.46
1:1G:974:A:P	14:5A:41:ARG:HH12	2.39	0.46
26:1H:1727:U:H2'	26:1H:1728:G:O4'	2.16	0.46
26:1H:2146:C:H4'	26:1H:2147:G:N7	2.30	0.46
26:1H:2629:A:OP1	26:1H:2629:A:H4'	2.16	0.46
26:1H:492:A:H2'	26:1H:493:G:O4'	2.15	0.46
24:3K:37:A:H3'	24:3K:38:A:C8	2.50	0.46
38:45:134:ARG:HG2	38:45:136:ALA:CB	2.46	0.46
38:45:69:PHE:CD1	38:45:70:PRO:HD2	2.51	0.46
5:4E:74:GLY:O	5:4E:115:VAL:HA	2.16	0.46
26:14:2761:G:H1'	33:59:143:GLN:HE22	1.80	0.46
28:71:45:ALA:HA	28:71:211:SER:O	2.15	0.46
28:71:30:LYS:HG3	28:71:182:PRO:HD3	1.96	0.46
17:8I:22:LEU:HD12	17:8I:40:LYS:O	2.16	0.46
45:B5:67:GLY:O	45:B5:69:TYR:N	2.47	0.46
20:BA:12:ALA:O	20:BA:15:ARG:HB2	2.15	0.46
47:H8:92:SER:O	47:H8:130:PRO:HG2	2.15	0.46
47:H8:52:SER:O	47:H8:52:SER:OG	2.21	0.46
29:11:38:LYS:HB3	29:11:39:LYS:HA	1.97	0.46
1:13:1278:U:H5'	1:13:1279:A:C5'	2.45	0.46
1:13:1260:C:O5'	1:13:1284:C:H4'	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1298:C:C5	7:6E:114:ARG:HD3	2.50	0.46
1:13:280:C:H4'	1:13:281:G:OP2	2.16	0.46
1:13:353:A:C8	1:13:353:A:H5'	2.47	0.46
1:13:377:G:H5'	16:7I:5:ARG:NH1	2.30	0.46
26:14:1826:G:H2'	26:14:1827:C:O4'	2.16	0.46
26:14:1839:G:C8	26:14:1927:A:H1'	2.50	0.46
26:14:384:U:H2'	26:14:385:C:C6	2.51	0.46
2:1E:166:ASP:C	2:1E:168:THR:H	2.19	0.46
1:1G:509:A:C8	1:1G:509:A:H3'	2.50	0.46
1:1G:757:U:O2'	1:1G:879:C:O2	2.31	0.46
26:1H:1105:U:H2'	26:1H:1106:G:C8	2.51	0.46
26:1H:1163:G:C2	26:1H:1164:G:C8	3.03	0.46
26:1H:1396:U:H2'	26:1H:1396:U:O2	2.15	0.46
26:1H:780:G:N2	26:1H:783:A:N6	2.59	0.46
3:22:73:PRO:O	3:22:76:VAL:HG22	2.16	0.46
30:29:8:LYS:HE2	30:29:188:VAL:HG22	1.98	0.46
3:2E:77:ILE:O	3:2E:84:ILE:HG22	2.15	0.46
4:32:139:ARG:HG3	4:32:139:ARG:HH11	1.81	0.46
31:39:66:PRO:O	31:39:67:GLN:CB	2.64	0.46
38:45:43:THR:HG22	38:45:94:VAL:HG12	1.98	0.46
25:4K:9:G:H3'	25:4K:10:G:O4'	2.16	0.46
39:55:33:ARG:NH2	39:55:115:GLU:OE2	2.41	0.46
26:14:2839:G:H21	39:55:92:GLY:HA2	1.81	0.46
1:13:1047:G:O3'	14:5I:4:LYS:HB2	2.15	0.46
34:61:124:GLY:H	34:61:142:VAL:HG23	1.80	0.46
7:62:65:ALA:HB1	7:62:127:ALA:HB3	1.96	0.46
7:6E:143:ARG:NH1	24:3K:41:A:O2'	2.49	0.46
28:71:180:PHE:HA	28:71:181:PRO:HD3	1.80	0.46
38:88:32:TYR:O	38:88:105:GLU:HA	2.16	0.46
39:98:59:ASP:OD1	39:98:59:ASP:N	2.37	0.46
19:AI:40:ILE:HG23	19:AI:41:VAL:N	2.31	0.46
41:B8:7:ILE:O	41:B8:11:GLU:HB2	2.15	0.46
20:BA:87:LYS:HG2	20:BA:91:LEU:HD11	1.98	0.46
47:D5:132:ASN:N	47:D5:132:ASN:OD1	2.47	0.46
43:D8:36:PRO:C	43:D8:38:LEU:H	2.19	0.46
45:F8:94:GLY:O	45:F8:95:LEU:HB2	2.15	0.46
46:G8:95:LYS:HE2	46:G8:97:ARG:HH12	1.80	0.46
48:I8:53:MET:HB2	48:I8:59:LEU:CD2	2.45	0.46
2:12:71:VAL:HG23	2:12:165:VAL:HG13	1.98	0.46
26:14:1036:G:H1	26:14:1119:C:H42	1.62	0.46
26:14:991:C:O2	26:14:1164:G:C2	2.69	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2291:U:H5"	26:14:2380:C:O2'	2.15	0.46
26:14:451:C:H41	26:14:454:A:H5'	1.81	0.46
26:14:521:G:H2'	26:14:522:G:H8	1.80	0.46
26:14:533:G:H2'	26:14:534:U:O4'	2.16	0.46
26:14:65:C:H2'	26:14:66:C:H6	1.80	0.46
26:14:588:U:O4	26:14:670:A:H1'	2.15	0.46
35:15:67:LEU:HG	35:15:88:GLU:HG2	1.97	0.46
10:1A:79:ARG:HH11	10:1A:79:ARG:HB3	1.80	0.46
1:1G:1142:G:H3'	1:1G:1143:G:H8	1.79	0.46
1:1G:501:C:OP1	12:3A:117:ARG:NH2	2.48	0.46
26:1H:1869:G:H5"	26:1H:1869:G:H8	1.80	0.46
26:1H:207:A:H2'	26:1H:208:C:O4'	2.16	0.46
26:1H:2262:U:H4'	26:1H:2328:A:C2	2.51	0.46
26:1H:2492:U:H2'	26:1H:2493:U:C6	2.50	0.46
26:1H:2544:G:H2'	26:1H:2545:G:C8	2.51	0.46
26:1H:705:A:O3'	29:11:7:LYS:HD2	2.16	0.46
27:1J:104:A:H2'	27:1J:105:G:O4'	2.16	0.46
30:21:4:ILE:HD13	30:21:28:ALA:HB1	1.98	0.46
1:1G:1206:G:O2'	3:22:193:TYR:HA	2.16	0.46
23:2K:20:G:C2	23:2K:58:A:N3	2.84	0.46
37:35:126:VAL:HG12	37:35:147:LEU:HD22	1.97	0.46
37:35:59:LEU:HA	37:35:62:LEU:HD22	1.97	0.46
13:4I:12:ASN:HD22	13:4I:13:LYS:N	2.13	0.46
7:62:141:VAL:HA	7:62:142:GLU:HB2	1.97	0.46
7:62:94:ARG:HA	7:62:97:GLN:HB3	1.97	0.46
40:65:101:LEU:HD12	40:65:105:ALA:HB2	1.97	0.46
27:1J:116:G:C5'	40:65:55:ALA:HB2	2.46	0.46
1:13:1240:U:P	7:6E:116:ALA:HB2	2.55	0.46
41:75:61:PHE:CE1	41:75:76:PHE:HB2	2.51	0.46
17:8A:59:ILE:HD13	17:8A:73:VAL:HA	1.98	0.46
20:BI:45:GLN:HA	20:BI:91:LEU:HB3	1.98	0.46
42:C8:50:ARG:NH2	43:D8:72:VAL:HG22	2.30	0.46
26:14:459:U:H5"	55:L5:40:TRP:CD2	2.51	0.46
2:12:24:TRP:NE1	2:12:26:PRO:HG3	2.31	0.46
1:13:1129:C:H4'	1:13:1130:A:OP1	2.15	0.46
1:13:1468:A:P	62:13:1850:HOH:O	2.73	0.46
1:13:342:C:C2'	1:13:343:U:H5'	2.46	0.46
1:13:724:G:C2	1:13:725:G:C8	3.03	0.46
26:14:1178:C:H2'	26:14:1179:C:C6	2.51	0.46
26:14:734:A:O2'	26:14:1635:G:H5'	2.16	0.46
26:14:1921:G:H2'	26:14:1922:G:H8	1.81	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2360:A:H2'	26:14:2361:A:O4'	2.15	0.46
2:1E:16:HIS:CE1	2:1E:213:LEU:HB2	2.51	0.46
1:1G:1256:A:N6	1:1G:1277:C:H3'	2.31	0.46
1:1G:631:G:H1'	1:1G:632:A:H5'	1.98	0.46
1:1G:953:G:H5"	1:1G:965:A:H61	1.80	0.46
26:1H:1470:G:N7	62:1H:3844:HOH:O	2.36	0.46
26:1H:194:G:H2'	26:1H:195:A:O4'	2.15	0.46
26:1H:2109:U:H1'	26:1H:2181:G:N2	2.31	0.46
26:1H:2711:A:P	62:1H:3750:HOH:O	2.74	0.46
26:1H:2766:G:H2'	26:1H:2766:G:N3	2.31	0.46
26:1H:674:G:O2'	31:31:74:ARG:HG3	2.16	0.46
26:1H:910:A:N1	26:1H:2277:G:H1'	2.31	0.46
26:1H:934:G:H2'	26:1H:935:C:C6	2.51	0.46
10:1I:80:LYS:O	10:1I:84:GLN:HG2	2.15	0.46
22:1K:27:G:H1	22:1K:43:U:H3	1.62	0.46
30:29:128:SER:OG	30:29:129:HIS:N	2.49	0.46
23:2K:54:G:H2'	23:2K:55:5MU:C6	2.49	0.46
31:31:108:LYS:O	31:31:112:MET:HG3	2.15	0.46
37:35:50:ARG:HD3	56:M5:7:HIS:HD2	1.79	0.46
12:3A:32:PHE:HB3	12:3A:84:LEU:HD22	1.97	0.46
13:4I:57:ARG:O	13:4I:61:GLU:HG3	2.15	0.46
26:1H:270(M):U:OP2	34:61:57:ARG:NH2	2.49	0.46
43:95:71:LEU:O	43:95:85:LYS:O	2.33	0.46
39:98:42:LYS:O	39:98:45:ARG:HD3	2.16	0.46
45:B5:63:LYS:H	45:B5:63:LYS:CE	2.29	0.46
47:D5:48:PHE:O	47:D5:52:SER:HB3	2.16	0.46
26:14:1364:G:OP1	49:F5:3:LYS:HD2	2.15	0.46
26:14:94:G:N2	50:G5:47:ASN:HD22	2.11	0.46
47:H8:59:LEU:HD23	47:H8:59:LEU:HA	1.82	0.46
52:M8:38:LYS:HE2	52:M8:44:THR:HG21	1.98	0.46
1:13:1171:G:O2'	1:13:1172:C:H5'	2.16	0.46
1:13:1:U:H6	1:13:630:G:H2'	1.80	0.46
26:14:1109:C:H2'	26:14:1110:G:C1'	2.46	0.46
26:14:1111:A:O3'	26:14:1112:G:H4'	2.15	0.46
26:14:1114:G:H2'	26:14:1115:G:C8	2.51	0.46
26:14:1448:G:H1'	26:14:1528:A:H62	1.81	0.46
26:14:2542:A:O2'	26:14:2543:G:OP2	2.24	0.46
23:2L:77:A:O2'	26:14:2602:A:N7	2.49	0.46
26:14:640:C:H42	26:14:648:G:H1	1.64	0.46
26:14:886:C:H1'	26:14:890:A:C2	2.49	0.46
1:1G:433:C:H2'	1:1G:434:U:C6	2.51	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:437:U:C4	1:1G:438:G:C6	3.04	0.46
26:1H:1329:U:H5"	26:1H:1330:C:C5	2.48	0.46
26:1H:1878:G:H2'	26:1H:1879:C:C6	2.51	0.46
26:1H:2126:A:C8	26:1H:2163:C:H1'	2.51	0.46
26:1H:2164:C:OP2	26:1H:2166:G:N2	2.48	0.46
26:1H:2251:G:OP1	38:88:82:ARG:NH1	2.48	0.46
26:1H:638:G:C5	26:1H:651:G:C2	3.03	0.46
3:22:61:ALA:C	3:22:63:ASN:H	2.19	0.46
3:22:94:LEU:H	3:22:94:LEU:HG	1.36	0.46
30:29:81:ILE:O	30:29:82:ARG:HB2	2.15	0.46
23:2L:73:A:C6	23:2L:74:A:C6	3.04	0.46
4:32:57:ARG:NH2	5:42:107:ARG:HD2	2.31	0.46
13:4A:92:HIS:CD2	13:4A:98:VAL:HG11	2.51	0.46
33:59:103:LEU:HD22	33:59:123:PHE:CZ	2.51	0.46
8:72:88:LYS:O	8:72:92:ARG:HD3	2.16	0.46
31:31:37:VAL:HG21	37:78:6:LEU:HD21	1.97	0.46
40:A8:25:ARG:NH1	40:A8:42:ASP:OD1	2.47	0.46
49:J8:23:LYS:HB3	49:J8:29:GLY:HA3	1.97	0.46
50:K8:4:SER:H	50:K8:7:ARG:HG2	1.79	0.46
55:L5:12:ARG:NH2	55:L5:44:PRO:HB3	2.31	0.46
1:13:1171:G:H2'	1:13:1172:C:H6	1.81	0.46
1:13:223:U:H2'	1:13:224:C:H6	1.81	0.46
1:13:272:C:H2'	1:13:273:A:C8	2.51	0.46
1:13:458:C:H42	1:13:474:G:H1	1.64	0.46
1:13:652:U:H1'	1:13:653:A:C2	2.51	0.46
26:14:1141:U:OP1	35:15:25:ARG:NE	2.39	0.46
26:14:2152:G:C6	26:14:2153:G:H1'	2.51	0.46
26:14:2186:G:H2'	26:14:2187:G:H8	1.81	0.46
26:14:2772:C:H2'	26:14:2773:C:C6	2.51	0.46
35:15:133:GLN:C	35:15:134:ARG:HG3	2.37	0.46
2:1E:11:LEU:HD11	2:1E:209:ARG:NH2	2.31	0.46
1:1G:1148:U:H2'	1:1G:1149:C:O4'	2.16	0.46
1:1G:1211:U:O2	1:1G:1213:A:H2	1.99	0.46
1:1G:1258:G:H8	1:1G:1258:G:OP2	1.99	0.46
1:1G:80:G:O2'	1:1G:81:G:OP1	2.32	0.46
26:1H:1049:C:C2'	26:1H:1050:A:H5'	2.44	0.46
26:1H:1337:G:H2'	26:1H:1338:G:H8	1.81	0.46
26:1H:1731:G:H2'	26:1H:1732:A:C8	2.50	0.46
26:1H:2074:U:H2'	26:1H:2075:U:C6	2.51	0.46
26:1H:2126:A:H1'	26:1H:2162:G:H21	1.80	0.46
26:1H:817:C:H4'	26:1H:932:G:C5	2.51	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:847:U:C5'	62:1H:3743:HOH:O	2.64	0.46
26:1H:918:A:H8	26:1H:918:A:O5'	1.99	0.46
30:21:81:ILE:HG22	30:21:81:ILE:O	2.16	0.46
30:29:111:ARG:HD2	30:29:160:TYR:CD2	2.50	0.46
31:31:64:ILE:HG23	31:31:65:TRP:CD1	2.51	0.46
12:3I:85:ILE:HD13	12:3I:85:ILE:HA	1.67	0.46
32:41:81:LYS:NZ	32:41:81:LYS:H	2.14	0.46
38:45:36:ALA:HB2	38:45:103:MET:SD	2.55	0.46
38:45:117:ALA:HA	38:45:120:ILE:HB	1.96	0.46
5:4E:37:ARG:HH12	5:4E:111:GLU:HG2	1.80	0.46
5:4E:147:ASP:O	5:4E:151:LEU:HB2	2.16	0.46
26:1H:6:A:H1'	35:58:131:GLN:HG3	1.98	0.46
14:5A:29:ARG:HD3	14:5A:31:ARG:O	2.16	0.46
7:62:12:LEU:HB2	7:62:21:VAL:HB	1.98	0.46
37:78:122:PRO:HA	37:78:142:GLY:CA	2.46	0.46
1:1G:235:C:C5'	17:8A:70:ARG:HG2	2.43	0.46
26:14:1614:A:H62	44:A5:93:ALA:HB2	1.81	0.46
20:BI:26:ASN:HB2	20:BI:71:THR:CG2	2.39	0.46
47:H8:53:ILE:HG22	47:H8:71:VAL:HG22	1.97	0.46
26:14:2815:C:H5'	53:J5:29:THR:HG21	1.98	0.46
50:K8:47:ASN:C	50:K8:49:LYS:H	2.19	0.46
29:11:68:LYS:HB3	29:11:70:TRP:CH2	2.52	0.45
2:12:51:LEU:H	2:12:51:LEU:HG	1.41	0.45
1:13:1084:G:C5	1:13:1085:U:C4	3.03	0.45
1:13:1126:U:C5	1:13:1127:G:N7	2.84	0.45
1:13:1425:U:H2'	1:13:1426:C:H6	1.81	0.45
26:14:1047:G:N3	26:14:1047:G:H2'	2.31	0.45
26:14:1442:G:H2'	26:14:1443:G:H8	1.81	0.45
26:14:1496:A:H8	26:14:1577:C:O2'	1.95	0.45
26:14:2006:C:H2'	26:14:2007:C:H6	1.81	0.45
26:14:2099:U:H3	26:14:2190:G:H1	1.65	0.45
26:14:2343:C:O2'	26:14:2373:G:O2'	2.14	0.45
26:14:2416:C:OP1	37:35:65:ARG:O	2.33	0.45
26:14:2773:C:H2'	26:14:2774:C:H6	1.82	0.45
26:14:2801:A:H2'	26:14:2802:G:O4'	2.16	0.45
26:14:289:A:H3'	26:14:290:G:H8	1.80	0.45
26:14:492:A:H2'	26:14:493:G:O4'	2.16	0.45
26:14:603:A:H8	26:14:604:G:H1'	1.81	0.45
26:14:654(C):G:H2'	26:14:654(D):G:H5'	1.99	0.45
21:1F:9:ARG:HD3	21:1F:13:ILE:HD11	1.97	0.45
26:1H:1217:C:H2'	26:1H:1218:C:H5"	1.97	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1728:G:H2'	26:1H:1731:G:O6	2.16	0.45
26:1H:1820:U:H4'	26:1H:1821:A:OP2	2.16	0.45
3:22:90:GLU:H	3:22:90:GLU:CD	2.20	0.45
3:2E:50:ALA:O	3:2E:70:VAL:HG22	2.16	0.45
31:39:7:TYR:HE2	31:39:10:PRO:HG3	1.81	0.45
31:39:155:LEU:HB2	31:39:189:THR:HG21	1.98	0.45
4:3E:90:GLY:O	4:3E:93:PHE:HB3	2.16	0.45
24:3K:2:G:O2'	24:3K:3:G:OP1	2.28	0.45
24:3K:50:C:H2'	24:3K:51:A:O4'	2.17	0.45
38:45:34:LEU:HD11	38:45:129:THR:HB	1.98	0.45
13:4I:45:VAL:HA	13:4I:48:LEU:HD22	1.96	0.45
6:5E:18:GLN:HA	6:5E:21:LEU:HB2	1.97	0.45
7:62:137:LYS:HE2	7:62:137:LYS:HB3	1.81	0.45
7:62:146:GLU:HG3	11:2A:50:TYR:OH	2.16	0.45
7:62:16:LEU:HD21	9:82:45:ALA:HB2	1.98	0.45
38:88:26:TYR:CD2	38:88:141:GLN:HG2	2.51	0.45
1:13:967:C:O2'	9:8E:125:TYR:OH	2.21	0.45
1:13:1372:U:H5"	9:8E:71:SER:HB3	1.98	0.45
43:95:37:VAL:O	43:95:39:LEU:N	2.45	0.45
40:A8:108:GLY:O	40:A8:110:LEU:HD12	2.16	0.45
26:1H:2334:G:H5'	40:A8:9:ARG:HG2	1.96	0.45
41:B8:51:ARG:HG3	41:B8:98:LYS:HD3	1.98	0.45
26:1H:18:C:O3'	42:C8:23:GLY:HA2	2.16	0.45
42:C8:85:LYS:HD2	42:C8:85:LYS:HA	1.50	0.45
44:E8:11:ARG:HH21	44:E8:99:ARG:N	2.14	0.45
52:M8:15:ILE:HB	52:M8:32:TYR:HD1	1.80	0.45
29:11:182:LEU:H	29:11:272:ALA:CB	2.10	0.45
2:12:124:SER:O	2:12:126:GLU:N	2.45	0.45
1:13:1129:C:N4	1:13:1143:G:H1	2.14	0.45
1:13:755:G:OP2	15:6I:65:ARG:HD2	2.16	0.45
1:13:881:G:P	12:3I:12:ARG:NH2	2.89	0.45
26:14:1525:G:H2'	26:14:1526:G:C8	2.51	0.45
26:14:1628:G:H2'	26:14:1629:U:C6	2.50	0.45
26:14:2086:U:H2'	26:14:2087:G:C8	2.51	0.45
26:14:2286:A:H8	26:14:2287:A:N6	2.14	0.45
26:14:2394:C:H2'	26:14:2395:C:H5'	1.98	0.45
26:14:259:G:HO2'	26:14:621:A:HO2'	1.63	0.45
26:14:307:G:H21	26:14:330:A:N6	2.13	0.45
26:14:607:U:N3	26:14:621:A:C2	2.74	0.45
26:14:882:G:H1	26:14:894:C:H42	1.64	0.45
26:14:952:G:C6	26:14:966:G:C6	3.04	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
27:16:15:A:H1'	27:16:109:G:C8	2.52	0.45
29:19:36:PRO:HB2	29:19:37:LEU:HD12	1.99	0.45
2:1E:231:GLU:CD	2:1E:231:GLU:H	2.18	0.45
2:1E:97:TRP:CZ3	2:1E:172:ILE:HB	2.51	0.45
1:1G:1194:U:H2'	1:1G:1195:C:C6	2.52	0.45
1:1G:280:C:H3'	1:1G:281:G:H5'	1.98	0.45
26:1H:1019:U:H3	26:1H:1142(A):A:H62	1.64	0.45
26:1H:1316:U:H2'	26:1H:1317:A:C8	2.51	0.45
26:1H:2210:G:H3'	26:1H:2211:G:C8	2.51	0.45
26:1H:2689:U:H5"	26:1H:2713:A:C2	2.51	0.45
26:1H:433:C:H2'	26:1H:434:U:C6	2.50	0.45
26:1H:900:A:H5'	26:1H:901:A:OP2	2.17	0.45
26:1H:940:G:H2'	26:1H:941:A:O4'	2.16	0.45
10:1I:61:GLU:OE1	14:5I:49:HIS:HE1	1.99	0.45
10:1I:91:PRO:HB3	10:1I:94:VAL:HB	1.98	0.45
22:1K:28:U:H3	22:1K:42:A:H2	1.62	0.45
36:25:113:LYS:O	36:25:117:LEU:HD22	2.16	0.45
30:29:34:VAL:HG12	30:29:64:LYS:HE3	1.98	0.45
31:31:28:ILE:HG12	31:31:119:ARG:HH21	1.82	0.45
1:1G:430:A:OP2	4:32:8:VAL:HG23	2.16	0.45
24:3K:1:G:C2	24:3K:73:A:C6	3.04	0.45
13:4I:66:LEU:O	13:4I:70:LEU:HB2	2.17	0.45
40:65:10:ARG:O	40:65:14:VAL:HG12	2.16	0.45
34:69:102:SER:HA	34:69:107:VAL:O	2.16	0.45
15:6A:17:ARG:HD3	15:6A:26:GLU:HG3	1.98	0.45
41:75:123:GLN:HA	41:75:126:ALA:HB3	1.96	0.45
9:82:32:ASP:HB3	9:82:35:GLU:HB3	1.98	0.45
39:98:117:VAL:HG22	39:98:118:GLU:H	1.81	0.45
44:A5:28:SER:OG	44:A5:31:GLU:HB2	2.16	0.45
47:D5:99:TYR:CD2	47:D5:123:ASP:HB3	2.51	0.45
49:F5:32:LYS:HB3	49:F5:32:LYS:HE3	1.82	0.45
46:G8:40:GLU:HA	46:G8:41:GLY:HA2	1.80	0.45
50:K8:21:LEU:HD13	50:K8:64:LEU:HA	1.98	0.45
26:1H:729:G:C6	29:11:208:LYS:HB2	2.51	0.45
2:12:141:GLU:O	2:12:145:LEU:HB2	2.17	0.45
1:13:1014:A:H4'	19:AI:14:HIS:NE2	2.31	0.45
1:13:1013:G:N2	1:13:1016:A:OP2	2.48	0.45
1:13:1399:C:C2	1:13:1401:G:C5	3.05	0.45
1:13:397:A:C6	1:13:548:G:N7	2.84	0.45
1:13:74:C:H2'	1:13:75:C:C5	2.51	0.45
1:13:663:A:H5'	1:13:836:G:OP1	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:28:A:C2	26:14:513:A:C8	3.05	0.45
10:1A:32:ALA:HA	10:1A:76:ASN:ND2	2.31	0.45
2:1E:156:LYS:HA	2:1E:156:LYS:HD2	1.68	0.45
21:1F:5:ASP:O	21:1F:11:GLY:HA3	2.15	0.45
1:1G:1171:G:H2'	1:1G:1172:C:C6	2.52	0.45
1:1G:129(A):G:C6	1:1G:188:U:H4'	2.51	0.45
1:1G:688:G:H2'	1:1G:689:C:H6	1.81	0.45
1:1G:872:A:C4	1:1G:874:G:N7	2.84	0.45
26:1H:1049:C:C4	26:1H:1050:A:C2	3.04	0.45
26:1H:2022:U:O2'	26:1H:2617:C:H5'	2.16	0.45
26:1H:2051:A:H5'	26:1H:2578:G:O4'	2.17	0.45
26:1H:2364:C:H4'	48:I8:56:ASP:OD1	2.17	0.45
26:1H:2461:C:H2'	26:1H:2462:U:C6	2.51	0.45
26:1H:942:G:H5'	62:1H:3621:HOH:O	2.16	0.45
27:1J:46:A:H2'	27:1J:47:C:C6	2.51	0.45
30:21:60:ASN:OD1	30:21:63:LEU:HB2	2.16	0.45
3:22:76:VAL:O	3:22:84:ILE:HA	2.15	0.45
36:25:114:ILE:H	36:25:114:ILE:HG12	1.50	0.45
4:32:149:ALA:O	4:32:153:ARG:NE	2.49	0.45
4:3E:108:LEU:HB3	4:3E:110:PHE:CE1	2.50	0.45
4:3E:18:LYS:HG2	59:3E:301:SF4:S1	2.57	0.45
32:49:125:PHE:HB3	32:49:166:ASP:CB	2.38	0.45
5:4E:19:MET:HE1	5:4E:24:ARG:HH21	1.82	0.45
13:4I:2:ALA:O	13:4I:10:PRO:HD2	2.17	0.45
33:51:97:ARG:HH21	33:51:104:GLU:CD	2.19	0.45
35:58:96:GLU:HG2	35:58:97:ARG:N	2.31	0.45
40:65:36:TYR:HE2	40:65:54:LEU:HD22	1.81	0.45
8:72:30:ARG:O	8:72:34:GLU:HG2	2.17	0.45
1:1G:1446:A:N3	41:75:118:ARG:HD2	2.30	0.45
36:25:104:ARG:HD3	41:75:36:GLU:HB2	1.99	0.45
16:7I:21:VAL:HG23	16:7I:34:GLU:H	1.82	0.45
38:88:14:ARG:HG2	38:88:41:TRP:HH2	1.81	0.45
17:8A:29:HIS:CG	17:8A:30:PRO:HD2	2.51	0.45
6:5E:97:PHE:CD1	18:9I:31:LEU:HD11	2.51	0.45
45:B5:62:LYS:HB3	45:B5:63:LYS:HZ1	1.81	0.45
41:B8:120:ARG:HA	41:B8:123:GLN:HG2	1.98	0.45
45:F8:25:LYS:HA	45:F8:81:VAL:O	2.16	0.45
47:H8:53:ILE:HA	47:H8:71:VAL:HG13	1.97	0.45
29:11:145:VAL:HG12	29:11:146:GLU:O	2.16	0.45
29:11:59:LYS:HG2	29:11:60:ARG:N	2.31	0.45
1:13:188:U:H2'	1:13:189:U:H5"	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:695:A:H2'	1:13:696:A:C8	2.51	0.45
26:14:2014:A:H2'	26:14:2015:A:C8	2.51	0.45
26:14:1701:A:OP2	61:14:3436:SPE:N1	2.49	0.45
26:14:627:A:H62	37:35:84:ASN:HD21	1.63	0.45
26:14:639:U:H2'	26:14:640:C:C6	2.51	0.45
27:16:94:C:H2'	27:16:95:U:H6	1.82	0.45
2:1E:97:TRP:HZ3	2:1E:172:ILE:HB	1.81	0.45
2:1E:6:THR:OG1	2:1E:7:VAL:N	2.49	0.45
1:1G:1058:G:H2'	1:1G:1059:C:O4'	2.16	0.45
1:1G:1241:G:OP1	7:62:35:LYS:NZ	2.48	0.45
1:1G:1277:C:O2'	1:1G:1279:A:H8	1.99	0.45
1:1G:49:U:C2	1:1G:361:G:N2	2.85	0.45
1:1G:959:A:HO2'	1:1G:984:C:HO2'	1.64	0.45
26:1H:1030:G:OP2	38:88:128:LYS:NZ	2.43	0.45
26:1H:1425:G:N2	26:1H:1573:G:N7	2.65	0.45
10:1I:31:GLY:HA2	10:1I:78:ASN:HB2	1.98	0.45
3:22:37:GLN:O	3:22:40:ARG:N	2.48	0.45
23:2K:62:C:H2'	23:2K:63:C:H6	1.81	0.45
31:39:11:VAL:HG23	31:39:12:LEU:N	2.31	0.45
31:39:141:ALA:O	31:39:144:LYS:HB3	2.17	0.45
31:39:89:VAL:HG12	31:39:90:PHE:N	2.32	0.45
24:3K:48:C:C5	24:3K:59:A:H1'	2.51	0.45
32:41:80:PHE:O	32:41:82:LEU:HB2	2.17	0.45
5:42:60:TYR:HB3	5:42:64:ARG:CZ	2.46	0.45
38:45:38:GLU:HG3	38:45:127:ILE:CG2	2.47	0.45
35:58:57:ALA:C	35:58:59:LYS:N	2.68	0.45
40:65:106:ARG:NH1	40:65:107:GLU:OE2	2.49	0.45
34:69:74:ASN:O	34:69:75:LEU:HB2	2.16	0.45
1:13:1374:A:O2'	7:6E:28:ASN:HB3	2.17	0.45
1:13:277:C:P	17:8I:68:ARG:HH12	2.39	0.45
46:C5:52:SER:H	46:C5:57:GLN:N	2.15	0.45
46:G8:95:LYS:O	46:G8:103:GLY:HA2	2.15	0.45
49:J8:81:LYS:HG3	49:J8:82:LEU:H	1.81	0.45
54:O8:41:PRO:HD2	54:O8:46:HIS:N	2.31	0.45
2:12:57:PHE:HZ	2:12:199:TYR:CZ	2.35	0.45
1:13:1125:U:O2'	1:13:1126:U:H6	1.97	0.45
26:14:1754:C:N3	26:14:2716:U:O2'	2.48	0.45
26:14:2027:G:H2'	26:14:2028:U:O4'	2.16	0.45
26:14:336:C:OP1	46:C5:83:THR:HG23	2.16	0.45
26:14:573:G:O2'	26:14:574:C:H3'	2.16	0.45
26:14:730:C:O2'	26:14:731:C:H5'	2.16	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:910:A:N3	26:14:2264:C:O2'	2.43	0.45
26:14:922:U:H2'	26:14:923:C:C6	2.51	0.45
26:14:999:U:H5"	26:14:1154:G:O6	2.16	0.45
29:19:13:ARG:HD2	29:19:13:ARG:HA	1.75	0.45
1:1G:1286:A:H3'	1:1G:1286:A:H8	1.81	0.45
1:1G:1523:G:OP1	11:2A:123:LYS:HD2	2.17	0.45
1:1G:407:G:C2	1:1G:436:C:C2	3.05	0.45
1:1G:422:C:O2'	1:1G:423:G:N2	2.50	0.45
26:1H:142:G:H2'	26:1H:143:C:C6	2.50	0.45
26:1H:1828:G:H8	26:1H:1828:G:OP2	1.99	0.45
26:1H:2356:C:H2'	26:1H:2357:U:O4'	2.17	0.45
26:1H:2557:G:H2'	26:1H:2558:C:C6	2.51	0.45
26:1H:2619:C:OP1	30:21:152:LYS:HE3	2.17	0.45
26:1H:801:G:OP2	31:31:55:GLY:HA2	2.17	0.45
22:1K:1:G:H2'	22:1K:2:G:O4'	2.16	0.45
36:25:34:THR:OG1	36:25:35:VAL:N	2.50	0.45
11:2A:103:LEU:HD12	11:2A:103:LEU:HA	1.71	0.45
23:2L:50:G:H1	23:2L:66:C:H42	1.63	0.45
31:39:103:LYS:HA	31:39:106:ARG:HG3	1.98	0.45
12:3A:39:VAL:HG23	12:3A:57:LYS:HD3	1.97	0.45
24:3K:3:G:O6	24:3K:69:A:N6	2.49	0.45
32:49:80:PHE:O	32:49:82:LEU:HB2	2.17	0.45
6:52:98:LEU:H	6:52:98:LEU:HD12	1.82	0.45
35:58:67:LEU:O	35:58:88:GLU:HG2	2.17	0.45
6:5E:67:MET:HB2	6:5E:68:PRO:HD2	1.98	0.45
40:65:62:LYS:HA	40:65:65:VAL:HB	1.99	0.45
34:69:109:ILE:HB	34:69:130:TYR:OH	2.17	0.45
7:6E:65:ALA:HB1	7:6E:127:ALA:HB3	1.99	0.45
28:71:225:ASN:HB3	28:71:227:HIS:ND1	2.31	0.45
41:75:2:ASN:HB3	41:75:4:GLY:C	2.37	0.45
16:7I:36:ILE:HG13	16:7I:36:ILE:O	2.16	0.45
9:82:63:ILE:HD11	9:82:81:ILE:HD11	1.98	0.45
38:88:138:ASP:HA	38:88:139:GLU:HA	1.69	0.45
38:88:5:ARG:H	38:88:5:ARG:HG3	1.61	0.45
18:9A:52:PRO:O	18:9A:56:THR:HG23	2.17	0.45
20:BA:50:GLU:CA	20:BA:100:ILE:HG12	2.46	0.45
20:BA:64:ASP:OD2	20:BA:81:LYS:NZ	2.40	0.45
35:58:42:TRP:O	42:C8:64:ARG:HD2	2.16	0.45
47:D5:94:GLU:O	47:D5:129:SER:HA	2.15	0.45
47:H8:63:ASP:OD1	47:H8:65:GLN:NE2	2.36	0.45
2:12:152:PHE:O	2:12:155:LEU:HD12	2.17	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1131:G:O6	26:14:2040:C:H1'	2.16	0.45
26:14:1005:C:C2	26:14:1143:A:C5	3.05	0.45
26:14:1248:G:C5	42:85:3:ARG:HB2	2.52	0.45
26:14:1253:A:OP1	62:14:3654:HOH:O	2.21	0.45
26:14:176:G:O2'	26:14:177:G:H5'	2.16	0.45
26:14:2038:G:H2'	26:14:2039:C:C6	2.52	0.45
26:14:2129:C:H3'	26:14:2130:U:H6	1.80	0.45
26:14:860:U:C2	26:14:2268:A:C8	3.05	0.45
26:14:2262:U:H4'	26:14:2328:A:H2	1.79	0.45
26:14:2472:G:H1	26:14:2477:C:P	2.39	0.45
26:14:2023:G:OP2	26:14:2617:C:H4'	2.17	0.45
26:14:2740:A:C6	26:14:2764:A:C8	3.05	0.45
26:14:839:U:H2'	26:14:840:C:C6	2.52	0.45
2:1E:11:LEU:HD12	2:1E:15:VAL:HG23	1.99	0.45
1:1G:109:A:H5'	1:1G:110:C:C5	2.52	0.45
1:1G:17:U:H2'	1:1G:18:C:C6	2.51	0.45
1:1G:500:G:H2'	1:1G:501:C:C6	2.52	0.45
26:1H:1429:G:O2'	26:1H:1430:C:H5'	2.16	0.45
26:1H:2562:U:H1'	36:68:23:ARG:HD3	1.98	0.45
26:1H:2023:G:H5'	26:1H:2617:C:H4'	1.98	0.45
10:1I:27:ALA:HA	10:1I:81:THR:HG22	1.97	0.45
27:1J:72:G:O2'	27:1J:104:A:N6	2.44	0.45
30:21:75:VAL:HG12	30:21:75:VAL:O	2.16	0.45
12:3I:60:LEU:HD12	12:3I:62:SER:OG	2.17	0.45
24:3K:48:C:OP2	24:3K:59:A:O2'	2.31	0.45
5:42:122:GLU:O	5:42:126:ARG:NH1	2.50	0.45
33:51:11:VAL:HG23	33:51:76:VAL:HG11	1.98	0.45
3:22:18:TRP:HD1	14:5A:54:PRO:HA	1.81	0.45
6:5E:16:GLN:HG2	6:5E:17:SER:N	2.31	0.45
40:65:54:LEU:O	40:65:56:LEU:N	2.44	0.45
7:6E:27:ILE:HD13	7:6E:40:ALA:HA	1.99	0.45
7:6E:73:MET:HG3	7:6E:89:MET:O	2.16	0.45
8:72:17:THR:HG23	8:72:65:TYR:HE2	1.80	0.45
17:8A:88:TYR:CE1	17:8A:92:ARG:HD2	2.52	0.45
9:8E:21:PRO:HA	9:8E:59:PHE:HA	1.99	0.45
44:A5:13:SER:O	44:A5:16:LYS:HB2	2.16	0.45
46:C5:89:PHE:O	46:C5:90:LEU:HB3	2.16	0.45
1:13:1153:C:H2'	1:13:1154:G:O4'	2.16	0.45
1:13:347:G:C2	1:13:348:G:H1'	2.52	0.45
1:13:509:A:H5"	4:3E:55:ALA:HB2	1.99	0.45
1:13:648:A:C6	1:13:649:G:C6	3.05	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:195:A:H61	26:14:198:C:H3'	1.81	0.45
26:14:2126:A:O2'	26:14:2127:G:H5"	2.17	0.45
26:14:2261:C:H1'	26:14:2388:A:N3	2.31	0.45
2:1E:187:LEU:HD11	2:1E:204:ASN:O	2.16	0.45
2:1E:11:LEU:HG	2:1E:213:LEU:HD22	1.98	0.45
1:1G:262:A:C6	1:1G:263:A:C6	3.05	0.45
1:1G:325:A:H2'	1:1G:326:G:O4'	2.16	0.45
1:1G:358:U:H2'	1:1G:359:U:H6	1.81	0.45
26:1H:129:C:H2'	26:1H:130:C:C6	2.52	0.45
26:1H:2136:C:H2'	26:1H:2137:C:O4'	2.17	0.45
26:1H:2607:G:H2'	26:1H:2608:G:O4'	2.17	0.45
26:1H:270(Y):G:C2	26:1H:270(Z):U:O4	2.70	0.45
26:1H:475:U:C4	26:1H:481:G:O6	2.70	0.45
26:1H:638:G:H2'	26:1H:639:U:C6	2.52	0.45
26:1H:755:C:H2'	26:1H:756:C:C6	2.52	0.45
30:29:182:LEU:HA	30:29:182:LEU:HD12	1.63	0.45
30:29:52:LEU:HA	30:29:52:LEU:HD12	1.56	0.45
11:2I:79:SER:HA	11:2I:104:GLN:O	2.16	0.45
11:2I:40:ILE:CG2	11:2I:75:TYR:HD2	2.30	0.45
23:2L:20:G:C4	23:2L:58:A:C2	3.04	0.45
31:31:149:ASP:OD1	31:31:149:ASP:N	2.41	0.45
32:49:18:GLU:OE2	32:49:21:ARG:NH2	2.43	0.45
32:49:50:ALA:HB2	32:49:87:PRO:HG3	1.99	0.45
13:4A:81:LEU:HD11	13:4A:86:CYS:SG	2.57	0.45
7:62:15:ASP:HB3	7:62:19:GLY:N	2.31	0.45
34:69:128:LEU:O	34:69:138:ILE:HG22	2.17	0.45
15:6A:11:VAL:HG21	15:6A:34:LEU:CD1	2.47	0.45
41:75:10:VAL:C	41:75:12:SER:H	2.20	0.45
37:78:97:PRO:HD3	37:78:126:VAL:O	2.16	0.45
37:78:88:LEU:HD12	37:78:88:LEU:HA	1.72	0.45
1:1G:1179:A:O3'	9:82:103:THR:HG23	2.17	0.45
9:82:42:ARG:NH1	9:82:75:ASP:OD1	2.37	0.45
18:9A:36:ASN:HB2	18:9A:38:GLU:OE1	2.16	0.45
46:C5:104:GLY:HA2	46:C5:105:ALA:HA	1.74	0.45
48:E5:27:GLU:HB2	48:E5:69:PHE:CD1	2.48	0.45
47:H8:105:VAL:HG22	47:H8:140:ASP:HA	1.99	0.45
1:13:1127:G:H2'	1:13:1128:C:O4'	2.16	0.45
1:13:1239:A:C4	1:13:1298:C:N4	2.85	0.45
1:13:1326:C:H2'	1:13:1327:C:C6	2.51	0.45
1:13:1349:A:H2'	1:13:1350:A:H8	1.81	0.45
1:13:272:C:H2'	1:13:273:A:H8	1.81	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:108:U:H4'	26:14:347:A:H2	1.82	0.45
26:14:1116:C:H2'	26:14:1117:G:C8	2.52	0.45
26:14:1204:A:N1	26:14:1241:A:C2	2.84	0.45
26:14:1805:U:O2	29:19:50:THR:HB	2.17	0.45
26:14:183:C:H1'	26:14:433:C:H1'	1.98	0.45
26:14:2261:C:O2'	26:14:2262:U:H5'	2.17	0.45
26:14:2342:C:OP2	26:14:2342:C:H6	1.98	0.45
26:14:2389:G:H5"	26:14:2390:U:O4'	2.15	0.45
26:14:2399:G:H2'	26:14:2400:G:O4'	2.17	0.45
26:14:2845:G:H5"	41:75:55:ASN:HA	1.99	0.45
26:14:2591:C:OP1	29:19:239:ARG:HG2	2.16	0.45
1:1G:1073:U:H2'	1:1G:1074:G:C8	2.52	0.45
1:1G:1221:G:OP1	1:1G:1321:C:N4	2.47	0.45
26:1H:1553:A:HO2'	26:1H:1554:A:H8	1.62	0.45
26:1H:1931:U:O4'	26:1H:1931:U:O2	2.35	0.45
26:1H:2138:C:H42	26:1H:2153:G:H1	1.64	0.45
26:1H:250:G:C6	26:1H:251:A:C6	3.04	0.45
26:1H:2562:U:H1'	36:68:23:ARG:NH1	2.32	0.45
26:1H:2820:A:OP2	26:1H:2821:A:N6	2.45	0.45
26:1H:576:U:OP1	62:1H:3779:HOH:O	2.21	0.45
10:1I:50:ILE:HD11	10:1I:57:LYS:HD2	1.98	0.45
3:22:18:TRP:HE3	3:22:18:TRP:N	2.13	0.45
30:29:23:VAL:HA	30:29:184:VAL:O	2.17	0.45
11:2A:48:ILE:HG21	11:2A:63:LEU:HD12	1.98	0.45
31:31:197:ASP:O	31:31:198:ALA:HB3	2.16	0.45
4:32:34:GLU:HB2	4:32:35:ARG:NH2	2.31	0.45
38:45:79:LEU:HD23	38:45:79:LEU:HA	1.73	0.45
32:49:145:THR:C	32:49:147:ASP:H	2.19	0.45
14:5A:17:LYS:NZ	14:5A:18:VAL:HG13	2.31	0.45
14:5I:6:LEU:CD1	14:5I:23:ARG:HH22	2.26	0.45
34:61:86:THR:HG22	34:61:122:GLU:HG2	1.99	0.45
34:61:95:LYS:HB3	34:61:95:LYS:HE3	1.87	0.45
34:69:14:ASP:N	34:69:17:GLN:OE1	2.41	0.45
7:6E:65:ALA:HB2	7:6E:128:ALA:HB2	1.99	0.45
5:4E:152:ARG:HA	8:7E:64:LYS:HZ3	1.80	0.45
38:88:133:ARG:O	38:88:134:ARG:HB2	2.17	0.45
38:88:36:ALA:O	38:88:99:PRO:HA	2.16	0.45
17:8I:78:GLU:OE2	17:8I:81:ARG:HD2	2.16	0.45
26:1H:2820:A:C6	39:98:4:LEU:HD11	2.52	0.45
47:H8:15:PRO:HB2	47:H8:19:ARG:NH2	2.32	0.45
49:J8:84:GLY:HA2	49:J8:85:LEU:CB	2.46	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
50:K8:41:ILE:HD13	50:K8:44:LEU:HG	1.99	0.45
56:M5:14:VAL:HG13	56:M5:22:VAL:HG13	1.97	0.45
1:13:1147:C:O2	9:8E:16:ARG:NH1	2.50	0.45
1:13:1234:C:H2'	1:13:1235:U:C6	2.51	0.45
1:13:612:C:O2	1:13:629:G:N2	2.50	0.45
26:14:1244:G:OP1	37:35:7:ARG:HD3	2.17	0.45
26:14:1796:U:H2'	26:14:1797:C:C6	2.52	0.45
26:14:2212:A:H1'	26:14:2215:G:C5	2.51	0.45
26:14:2512:C:H2'	26:14:2513:G:O4'	2.17	0.45
26:14:524:U:H2'	26:14:525:U:H6	1.80	0.45
26:14:923:C:H2'	26:14:924:C:C6	2.52	0.45
35:15:99:LEU:O	35:15:103:VAL:HG23	2.17	0.45
1:1G:1004:A:H8	1:1G:1026:G:C8	2.35	0.45
1:1G:1224:G:O6	1:1G:1322:C:H1'	2.17	0.45
1:1G:1322:C:O2'	1:1G:1323:G:H5'	2.16	0.45
1:1G:362:G:O2'	12:3A:33:ARG:NH2	2.49	0.45
26:1H:1108:U:H2'	26:1H:1109:C:H5'	1.99	0.45
26:1H:1805:U:O2	29:11:50:THR:HB	2.17	0.45
26:1H:205:G:HO2'	26:1H:206:U:P	2.39	0.45
26:1H:270(L):U:O2	34:61:50:ARG:HG2	2.16	0.45
26:1H:747:U:O2	26:1H:2014:A:H1'	2.17	0.45
26:1H:781:A:C8	29:11:219:PRO:HG2	2.52	0.45
27:1J:38:C:O2'	40:65:93:LYS:NZ	2.50	0.45
30:21:59:VAL:HG22	30:21:60:ASN:N	2.32	0.45
3:2E:134:ILE:HG22	3:2E:168:ALA:HB3	1.98	0.45
31:31:101:LEU:HD22	31:31:105:VAL:HB	1.99	0.45
4:32:33:MET:O	4:32:34:GLU:HB2	2.17	0.45
12:3A:60:LEU:HD13	12:3A:60:LEU:HA	1.75	0.45
24:3K:45:G:H4'	24:3K:46:G:OP1	2.16	0.45
1:1G:947:G:O3'	13:4A:109:THR:OG1	2.35	0.45
5:4E:36:ASP:OD2	5:4E:40:ARG:NH1	2.50	0.45
5:4E:51:VAL:O	5:4E:55:VAL:HG23	2.17	0.45
8:72:41:ARG:H	8:72:41:ARG:HG3	1.49	0.45
16:7I:74:LEU:HD22	16:7I:79:VAL:HG21	1.99	0.45
43:95:70:ILE:O	43:95:70:ILE:HG22	2.17	0.45
19:AA:12:ASP:OD1	19:AA:13:ASP:N	2.48	0.45
19:AI:51:VAL:HG12	19:AI:52:TYR:H	1.81	0.45
41:B8:50:ILE:HG13	41:B8:99:LEU:O	2.17	0.45
46:C5:89:PHE:O	46:C5:89:PHE:CG	2.70	0.45
42:C8:58:ARG:HA	42:C8:61:TRP:CE3	2.51	0.45
47:D5:128:VAL:HG23	47:D5:160:GLY:O	2.17	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:E8:88:ARG:HB2	44:E8:92:ARG:CB	2.44	0.45
49:F5:84:GLY:HA2	49:F5:85:LEU:HB3	1.99	0.45
46:G8:93:GLY:O	46:G8:94:LYS:HB2	2.17	0.45
52:M8:43:TYR:O	52:M8:46:GLN:HA	2.16	0.45
26:14:1138:G:H21	35:15:106:MET:CE	2.25	0.45
26:14:11:G:H2'	26:14:12:U:H5'	1.98	0.45
26:14:1316:U:H2'	26:14:1317:A:C8	2.51	0.45
26:14:1784:A:H4'	26:14:1785:A:C5'	2.47	0.45
26:14:387:U:H4'	26:14:388:G:O5'	2.17	0.45
26:14:921:G:C6	26:14:922:U:C4	3.06	0.45
29:19:245:PRO:HG2	29:19:253:GLN:NE2	2.32	0.45
2:1E:187:LEU:HA	2:1E:201:ILE:O	2.17	0.45
1:1G:1052:U:H5"	1:1G:1053:G:OP2	2.17	0.45
1:1G:746:A:H2'	1:1G:747:C:C6	2.52	0.45
1:1G:957:U:H1'	1:1G:960:U:C5	2.52	0.45
1:1G:991:U:O2	1:1G:993:G:H8	2.00	0.45
26:1H:1048:A:H5'	26:1H:1049:C:OP2	2.17	0.45
26:1H:1359:A:H2	26:1H:1372:U:O4	1.99	0.45
26:1H:1470:G:H5"	26:1H:1471:A:OP1	2.17	0.45
26:1H:1557:C:OP2	26:1H:1558:A:O2'	2.28	0.45
26:1H:1614:A:H8	26:1H:1614:A:O5'	2.00	0.45
26:1H:2855:C:H2'	26:1H:2856:C:H6	1.81	0.45
26:1H:524:U:H4'	26:1H:554:U:H4'	1.98	0.45
26:1H:616:A:C4	31:31:180:GLY:HA3	2.52	0.45
27:1J:101:A:H3'	27:1J:102:G:H8	1.82	0.45
3:22:152:ILE:O	3:22:198:VAL:HA	2.17	0.45
3:2E:17:ASP:O	3:2E:54:ARG:NH2	2.40	0.45
3:2E:72:LYS:HD3	3:2E:75:VAL:CG2	2.47	0.45
37:35:57:THR:O	37:35:61:ARG:HG3	2.17	0.45
31:39:69:HIS:CD2	31:39:69:HIS:N	2.84	0.45
24:3K:56:C:H2'	24:3K:57:G:O4'	2.16	0.45
24:3L:28:U:H2'	24:3L:29:U:C6	2.52	0.45
38:45:98:LYS:HB3	38:45:99:PRO:HD2	1.99	0.45
13:4A:80:ARG:O	13:4A:84:ILE:HB	2.17	0.45
5:4E:110:LEU:HD13	5:4E:118:ILE:HG21	1.99	0.45
33:51:4:ILE:HG23	33:51:6:ARG:CZ	2.47	0.45
34:61:38:LEU:HD11	49:J8:75:GLU:HG3	1.99	0.45
34:69:73:GLU:OE2	34:69:137:PRO:HD2	2.16	0.45
34:69:9:LEU:HD11	34:69:35:LEU:HB3	1.99	0.45
34:69:93:THR:O	34:69:97:ILE:HG13	2.17	0.45
8:72:83:ILE:HB	8:72:137:VAL:HG13	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:82:5:TYR:HA	9:82:17:VAL:O	2.16	0.45
42:85:102:GLU:HB3	42:85:105:VAL:HG13	1.99	0.45
43:95:1:MET:HA	43:95:42:GLY:N	2.30	0.45
40:A8:93:LYS:HG2	40:A8:95:HIS:HB2	1.99	0.45
44:E8:7:ALA:HB2	44:E8:50:VAL:HG22	1.99	0.45
53:J5:11:THR:HG23	53:J5:15:ARG:HB3	1.99	0.45
52:M8:38:LYS:HE3	52:M8:38:LYS:HA	1.99	0.45
53:N8:33:CYS:SG	53:N8:46:CYS:HB3	2.56	0.45
1:13:363:A:OP1	12:3I:33:ARG:HG3	2.17	0.44
1:13:828:A:H2'	1:13:829:G:O4'	2.16	0.44
26:14:1283:G:N2	26:14:1285:G:H3'	2.32	0.44
26:14:2062:A:O2'	26:14:2063:C:P	2.75	0.44
26:14:470:A:H8	26:14:470:A:C5'	2.30	0.44
26:14:657:U:H2'	26:14:658:C:C6	2.52	0.44
26:14:65:C:H2'	26:14:66:C:C6	2.52	0.44
2:1E:55:PHE:CD1	2:1E:58:ILE:HD12	2.52	0.44
21:1F:6:ARG:HH11	21:1F:15:ARG:NE	2.16	0.44
1:1G:987:G:N2	1:1G:1218:C:N3	2.46	0.44
1:1G:1497:G:H2'	1:1G:1498:U:H5'	1.99	0.44
1:1G:322:C:H5	1:1G:328:C:H5	1.65	0.44
1:1G:589:C:N3	1:1G:650:G:N2	2.46	0.44
1:1G:947:G:H2'	1:1G:948:C:C6	2.53	0.44
26:1H:141:A:H8	26:1H:1408:C:H1'	1.82	0.44
26:1H:1550:C:H2'	26:1H:1551:C:H6	1.83	0.44
26:1H:2845:G:H2'	26:1H:2846:G:H8	1.82	0.44
26:1H:517:C:OP1	53:N8:16:ARG:NH2	2.39	0.44
27:1J:56:G:H4'	27:1J:57:A:C8	2.51	0.44
30:21:49:LEU:HD12	30:21:49:LEU:HA	1.40	0.44
3:22:51:GLY:O	3:22:70:VAL:HG13	2.17	0.44
37:35:122:PRO:CB	37:35:141:ALA:HB1	2.45	0.44
31:39:25:PRO:C	31:39:27:GLU:N	2.70	0.44
24:3L:13:C:H2'	24:3L:14:A:H8	1.82	0.44
24:3L:55:U:H2'	24:3L:57:G:P	2.56	0.44
32:41:145:THR:O	32:41:146:TYR:HB3	2.18	0.44
6:5E:22:GLU:O	6:5E:26:ILE:HG13	2.16	0.44
34:61:73:GLU:OE2	34:61:137:PRO:HD2	2.17	0.44
16:7A:11:SER:H	16:7A:14:ASN:HB3	1.82	0.44
18:9I:44:LEU:HD11	18:9I:70:ILE:HG21	1.99	0.44
47:D5:62:PRO:O	47:D5:63:ASP:HB3	2.17	0.44
43:D8:49:THR:HG23	43:D8:51:VAL:H	1.81	0.44
51:H5:5:LYS:HB3	51:H5:5:LYS:HE3	1.72	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:H8:164:ALA:O	47:H8:165:VAL:HG22	2.18	0.44
32:41:67:LYS:CE	52:M8:6:HIS:CE1	2.98	0.44
54:O8:14:THR:OG1	54:O8:15:GLU:N	2.50	0.44
6:5E:81:ILE:HD11	29:11:125:ILE:HB	2.00	0.44
1:13:158:G:C4	1:13:159:G:C8	3.05	0.44
26:14:142:G:H2'	26:14:143:C:H6	1.81	0.44
26:14:656:G:H2'	26:14:657:U:O4'	2.16	0.44
26:14:671:C:OP1	37:35:42:SER:O	2.34	0.44
26:14:817:C:H2'	26:14:818:G:O4'	2.17	0.44
35:15:26:LEU:HD23	35:15:60:ILE:HD11	1.98	0.44
27:16:75:G:H21	47:H8:85:HIS:CE1	2.34	0.44
1:1G:1441:G:H8	1:1G:1441:G:O5'	2.00	0.44
1:1G:922:G:H2'	1:1G:923:A:C8	2.52	0.44
1:1G:931:C:H2'	1:1G:932:C:H6	1.82	0.44
1:1G:951:G:O4'	1:1G:971:G:H5'	2.17	0.44
26:1H:1019:U:O2'	26:1H:1021:A:H2	1.99	0.44
26:1H:330:A:H2	26:1H:1210:A:O2'	2.01	0.44
26:1H:1484:G:C2	26:1H:1506:C:C2	3.05	0.44
26:1H:184:C:H2'	26:1H:185:U:C6	2.52	0.44
26:1H:2244:U:H2'	26:1H:2245:U:O4'	2.17	0.44
26:1H:2774:C:H2'	26:1H:2775:A:O4'	2.16	0.44
26:1H:761:A:OP1	62:1H:3780:HOH:O	2.21	0.44
26:1H:770:G:C8	62:1H:4083:HOH:O	2.56	0.44
26:1H:813:U:H2'	26:1H:814:C:C6	2.52	0.44
30:21:26:ILE:HD11	30:21:198:VAL:HG21	1.99	0.44
36:25:3:GLN:HB2	36:25:4:PRO:HD2	2.00	0.44
36:25:13:ASN:ND2	36:25:97:ARG:H	2.13	0.44
31:31:155:LEU:HD13	31:31:174:VAL:HG22	1.98	0.44
4:32:173:TRP:HZ3	4:32:193:ASP:HB3	1.82	0.44
4:32:81:GLU:OE2	4:32:139:ARG:NH2	2.51	0.44
12:3A:47:LYS:HB3	12:3A:48:PRO:HD2	1.99	0.44
4:3E:98:GLU:OE2	4:3E:107:ARG:NH1	2.50	0.44
4:3E:109:GLY:HA3	4:3E:165:MET:SD	2.57	0.44
4:3E:86:LYS:H	4:3E:86:LYS:HD3	1.83	0.44
32:41:49:ASP:OD2	32:41:51:ARG:NH2	2.50	0.44
32:41:46:ALA:HB2	32:41:52:ILE:HB	2.00	0.44
38:45:126:PRO:O	38:45:127:ILE:HG23	2.17	0.44
38:45:29:PHE:HB3	38:45:65:PHE:CZ	2.52	0.44
38:45:4:PRO:HD3	38:45:70:PRO:O	2.18	0.44
6:52:35:ALA:HB1	6:52:65:VAL:CG1	2.47	0.44
34:69:44:LEU:HD23	34:69:44:LEU:HA	1.74	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:82:79:LEU:HD22	9:82:82:ALA:HB3	1.99	0.44
42:85:47:TYR:HA	42:85:50:ARG:NH2	2.32	0.44
17:8I:11:VAL:HG23	17:8I:20:THR:HB	1.99	0.44
39:98:18:LEU:HD23	39:98:18:LEU:HA	1.73	0.44
26:1H:2378:A:O2'	40:A8:21:THR:HG21	2.17	0.44
19:AI:5:LEU:O	19:AI:6:LYS:HB3	2.17	0.44
46:C5:101:LYS:HE3	46:C5:101:LYS:H	1.81	0.44
46:C5:63:LYS:HA	46:C5:63:LYS:NZ	2.32	0.44
26:14:297:C:OP1	46:C5:84:ARG:NH2	2.51	0.44
43:D8:34:GLU:HG2	43:D8:56:SER:OG	2.17	0.44
44:E8:11:ARG:CZ	44:E8:98:LYS:HB3	2.46	0.44
26:1H:495:G:H1'	44:E8:57:ASN:OD1	2.18	0.44
46:G8:81:LYS:HD2	46:G8:99:CYS:SG	2.58	0.44
50:K8:3:LEU:HA	50:K8:3:LEU:HD23	1.50	0.44
29:11:17:THR:HG21	29:11:204:ILE:HA	2.00	0.44
2:12:91:PRO:HB3	2:12:151:GLY:O	2.18	0.44
1:13:1226:C:H4'	1:13:1227:A:OP1	2.17	0.44
1:13:321:A:C2	1:13:333:G:C2	3.05	0.44
26:14:1929:G:H4'	26:14:1930:G:OP1	2.17	0.44
26:14:2271:G:H5"	48:E5:20:ARG:NE	2.32	0.44
26:14:957:A:N6	26:14:2459:A:C8	2.86	0.44
26:14:2619:C:H2'	26:14:2620:C:C6	2.53	0.44
26:14:396:G:H8	26:14:396:G:O5'	2.00	0.44
2:1E:155:LEU:HD13	2:1E:157:ARG:O	2.17	0.44
2:1E:237:ALA:O	2:1E:239:VAL:HG23	2.18	0.44
1:1G:545:C:OP1	4:32:61:LYS:NZ	2.50	0.44
26:1H:1011:G:OP1	42:C8:77:SER:OG	2.19	0.44
26:1H:1204:A:N1	26:1H:1241:A:C2	2.85	0.44
26:1H:1317:A:H2'	26:1H:1318:C:C6	2.52	0.44
26:1H:1338:G:O2'	26:1H:1393:A:N1	2.40	0.44
26:1H:2160:G:C5	26:1H:2161:C:H1'	2.52	0.44
26:1H:270(Y):G:H5"	26:1H:273(A):G:OP1	2.17	0.44
26:1H:660:G:H21	37:78:12:ALA:HA	1.82	0.44
26:14:861:A:N3	27:1J:79:C:O2'	2.51	0.44
3:22:106:VAL:O	3:22:109:PRO:HD3	2.18	0.44
3:22:113:ALA:O	3:22:116:VAL:HG12	2.16	0.44
3:2E:175:LEU:HD11	3:2E:201:TYR:CD2	2.52	0.44
4:32:148:VAL:O	4:32:152:SER:OG	2.28	0.44
31:39:5:ALA:HB1	31:39:125:LEU:HD21	1.98	0.44
31:39:164:ARG:HG2	31:39:175:THR:OG1	2.18	0.44
12:3I:21:LYS:HB3	12:3I:21:LYS:HE2	1.52	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:41:37:VAL:HG22	32:41:159:VAL:HG13	1.99	0.44
7:6E:5:ARG:HG3	7:6E:7:ALA:H	1.81	0.44
28:71:59:ARG:HG3	28:71:164:ARG:HD2	1.98	0.44
8:72:44:PHE:HA	8:72:79:VAL:HG11	1.99	0.44
8:7E:82:HIS:HE1	8:7E:84:ARG:HD3	1.81	0.44
9:82:5:TYR:CE1	9:82:16:ARG:HG2	2.51	0.44
18:9I:66:LEU:O	18:9I:70:ILE:HG13	2.17	0.44
19:AA:3:ARG:NH2	19:AA:11:VAL:HG12	2.32	0.44
47:D5:95:PRO:HA	47:D5:128:VAL:O	2.16	0.44
47:D5:27:VAL:HG23	47:D5:36:LYS:HA	2.00	0.44
26:1H:2261:C:C6	48:I8:16:SER:HB3	2.52	0.44
53:J5:31:VAL:HG13	53:J5:42:PRO:HG3	2.00	0.44
50:K8:31:GLU:HB3	50:K8:53:LEU:HD11	1.98	0.44
51:L8:18:ASP:OD1	51:L8:19:GLN:N	2.50	0.44
51:L8:51:ALA:HA	51:L8:54:VAL:HG12	1.99	0.44
56:M5:40:GLU:HG3	56:M5:43:GLN:NE2	2.32	0.44
19:AI:68:GLY:HA3	52:M8:59:PHE:HD2	1.83	0.44
1:13:110:C:H2'	1:13:111:G:O4'	2.17	0.44
1:13:1392:G:O2'	1:13:1393:U:H5'	2.18	0.44
1:13:820:U:H4'	1:13:821:G:OP2	2.18	0.44
26:14:2076:U:O5'	26:14:2076:U:H6	1.99	0.44
26:14:2151:G:C2	26:14:2152:G:H1'	2.51	0.44
26:14:235:U:H2'	26:14:236:C:H6	1.80	0.44
26:14:2823:A:OP1	30:29:159:HIS:NE2	2.48	0.44
26:14:2872:G:C4	26:14:2873:A:C2	3.05	0.44
26:14:312:G:H4'	26:14:331:A:N3	2.33	0.44
26:14:41:C:H2'	26:14:43:G:H8	1.82	0.44
26:14:839:U:H2'	26:14:840:C:H6	1.82	0.44
35:15:63:THR:O	35:15:66:LYS:HE2	2.17	0.44
27:16:11:C:H3'	27:16:12:C:C6	2.53	0.44
29:19:268:ARG:HG3	29:19:268:ARG:O	2.17	0.44
10:1A:55:LYS:HZ3	10:1A:57:LYS:HG2	1.81	0.44
1:1G:1519:A:H5"	1:1G:1520:G:OP2	2.17	0.44
1:1G:266:G:H2'	1:1G:266:G:N3	2.32	0.44
26:1H:1510:A:N3	26:1H:1510:A:H2'	2.32	0.44
26:1H:1535:U:OP2	26:1H:1538:G:N2	2.50	0.44
26:1H:2025:C:H2'	26:1H:2026:C:C6	2.52	0.44
26:1H:2310:A:H5"	26:1H:2311:A:OP2	2.17	0.44
26:1H:2542:A:H4'	26:1H:2543:G:C8	2.53	0.44
26:1H:273(E):U:H2'	26:1H:273(F):C:H5'	1.99	0.44
27:1J:66:A:N6	27:1J:108:C:O5'	2.50	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
22:1K:21:A:C5	22:1K:47:U:C4	3.06	0.44
24:1L:22:G:OP1	24:1L:48:C:N4	2.50	0.44
3:22:18:TRP:CE3	3:22:18:TRP:N	2.85	0.44
3:22:151:VAL:HA	3:22:199:LYS:O	2.17	0.44
3:22:44:GLU:HA	3:22:52:LEU:HD11	1.99	0.44
11:2A:69:ALA:O	11:2A:73:MET:HG3	2.18	0.44
11:2A:85:ARG:HD3	11:2A:113:PRO:HD3	1.98	0.44
4:32:177:ASP:O	4:32:180:GLY:N	2.46	0.44
4:32:189:PRO:HB2	4:32:194:LEU:HD21	1.99	0.44
12:3I:110:VAL:CG2	12:3I:120:TYR:HB3	2.47	0.44
38:45:26:TYR:HD1	38:45:27:VAL:HG23	1.73	0.44
13:4A:37:THR:HG22	13:4A:55:ARG:NE	2.33	0.44
5:4E:113:ALA:O	5:4E:115:VAL:HG23	2.18	0.44
13:4I:22:ILE:HB	13:4I:25:ILE:HD12	1.99	0.44
13:4I:82:MET:C	13:4I:84:ILE:H	2.21	0.44
25:4L:21:A:H8	25:4L:21:A:O5'	2.01	0.44
6:52:8:ILE:HD11	6:52:79:LEU:HD13	1.98	0.44
39:55:33:ARG:HB2	39:55:115:GLU:CB	2.45	0.44
35:58:127:ASP:O	35:58:128:HIS:HB3	2.17	0.44
28:71:226:PRO:HD2	28:71:227:HIS:CE1	2.53	0.44
18:9A:22:VAL:CG1	18:9A:56:THR:HA	2.47	0.44
19:AA:66:MET:O	19:AA:69:HIS:HB2	2.18	0.44
41:B8:11:GLU:O	41:B8:12:SER:HB2	2.18	0.44
47:D5:6:LYS:HD2	47:D5:40:ASP:CB	2.47	0.44
47:D5:4:ARG:NH2	47:D5:58:VAL:HG11	2.33	0.44
47:H8:48:PHE:HE1	47:H8:71:VAL:HG11	1.81	0.44
54:O8:26:ASN:ND2	54:O8:28:ARG:O	2.51	0.44
1:13:1394:A:C6	1:13:1501:C:H4'	2.53	0.44
1:13:1513:A:H2'	1:13:1514:C:C6	2.52	0.44
1:13:741:G:H2'	1:13:742:G:O4'	2.17	0.44
26:14:1000:A:C6	26:14:1001:A:N1	2.85	0.44
26:14:1820:U:H4'	26:14:1821:A:OP2	2.17	0.44
26:14:244:A:H2'	26:14:245:G:O4'	2.18	0.44
26:14:459:U:H4'	55:L5:40:TRP:CZ3	2.52	0.44
26:14:924:C:H2'	26:14:925:C:C6	2.52	0.44
29:19:245:PRO:HG2	29:19:253:GLN:HE21	1.83	0.44
2:1E:194:PRO:O	2:1E:196:LEU:N	2.44	0.44
1:1G:1051:C:H2'	1:1G:1052:U:C6	2.53	0.44
1:1G:373:A:C2	1:1G:374:A:C8	3.05	0.44
1:1G:588:G:H1	1:1G:651:C:N4	2.14	0.44
1:1G:659:U:H2'	1:1G:660:G:O4'	2.16	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:784:C:H2'	1:1G:785:G:O4'	2.17	0.44
26:1H:2028:U:H2'	26:1H:2029:G:O4'	2.18	0.44
26:1H:270(G):C:H2'	26:1H:270(H):C:O4'	2.18	0.44
26:1H:370:G:H5"	26:1H:423:A:N6	2.33	0.44
26:1H:937:U:H2'	26:1H:938:G:O4'	2.18	0.44
26:1H:992:C:H2'	26:1H:993:G:H8	1.82	0.44
26:1H:2032:G:N2	30:21:146:THR:HG23	2.20	0.44
30:21:67:PHE:HA	30:21:68:ALA:HA	1.71	0.44
30:21:65:GLY:HA2	30:21:67:PHE:O	2.18	0.44
3:2E:3:ASN:OD1	3:2E:3:ASN:N	2.50	0.44
4:32:31:CYS:CB	4:32:33:MET:H	2.29	0.44
4:3E:10:ARG:HB2	4:3E:10:ARG:HH11	1.82	0.44
12:3I:8:ASN:O	12:3I:12:ARG:HG3	2.18	0.44
24:3K:9:A:O2'	24:3K:46:G:O5'	2.36	0.44
24:3L:67:C:H2'	24:3L:68:G:C8	2.52	0.44
32:41:61:ALA:HB2	32:41:67:LYS:HA	1.98	0.44
32:41:83:ARG:H	32:41:86:MET:CE	2.30	0.44
39:55:86:ARG:HB3	39:55:118:GLU:OE1	2.17	0.44
33:59:31:GLY:O	33:59:79:VAL:HB	2.18	0.44
1:13:737:A:H5'	6:5E:90:VAL:O	2.18	0.44
7:62:146:GLU:OE2	11:2A:54:ARG:NE	2.48	0.44
36:68:22:ILE:HG21	36:68:22:ILE:HD13	1.69	0.44
1:1G:755:G:OP2	15:6A:65:ARG:HD3	2.17	0.44
7:6E:74:GLU:HG2	7:6E:91:VAL:HG13	1.99	0.44
28:71:189:ILE:O	28:71:193:ILE:HD13	2.18	0.44
28:71:49:ILE:HG22	28:71:204:ALA:HB1	2.00	0.44
8:72:19:VAL:CG2	8:72:21:LYS:HB3	2.48	0.44
19:AA:3:ARG:HD2	19:AA:7:LYS:CG	2.44	0.44
41:B8:23:ARG:HG3	41:B8:120:ARG:NH1	2.31	0.44
42:C8:79:PHE:C	42:C8:79:PHE:CD1	2.91	0.44
43:D8:50:PRO:HB2	43:D8:51:VAL:HG12	2.00	0.44
48:E5:72:ARG:HB3	48:E5:75:LEU:HB2	1.99	0.44
2:12:111:ARG:HA	2:12:111:ARG:HD3	1.84	0.44
1:13:183:G:H2'	1:13:184:G:H8	1.82	0.44
1:13:599:C:H2'	1:13:600:C:H6	1.82	0.44
1:13:827:U:C5	1:13:870:U:C4	3.06	0.44
1:13:977:A:H1'	1:13:982:U:O4	2.18	0.44
26:14:1287:A:C5	26:14:1288:U:C4	3.06	0.44
26:14:1915:U:H2'	26:14:1916:A:H5'	1.99	0.44
26:14:2353:G:N7	62:14:3712:HOH:O	2.36	0.44
26:14:2494:G:H2'	26:14:2495:G:H8	1.82	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:868:U:C2	26:14:869:G:C8	3.05	0.44
26:14:1567:A:O2'	29:19:63:ARG:NH2	2.51	0.44
29:19:68:LYS:HB3	29:19:70:TRP:CZ3	2.53	0.44
2:1E:87:ARG:HH11	2:1E:223:ILE:HD11	1.83	0.44
2:1E:63:MET:HB2	2:1E:225:ALA:HB1	1.98	0.44
1:1G:1119:C:H42	1:1G:1154:G:H1	1.65	0.44
1:1G:1309:G:C6	1:1G:1329:A:N1	2.86	0.44
1:1G:1347:G:N2	1:1G:1373:G:H2'	2.33	0.44
1:1G:511:C:H4'	4:32:43:HIS:CD2	2.52	0.44
1:1G:652:U:O2'	1:1G:653:A:N3	2.46	0.44
1:1G:652:U:O2'	1:1G:653:A:O5'	2.35	0.44
1:1G:748:C:O5'	1:1G:748:C:H6	2.01	0.44
1:1G:836:G:C6	1:1G:851:G:C6	3.05	0.44
26:1H:1408:C:C2	26:1H:1595:G:N2	2.86	0.44
26:1H:1900:A:N1	26:1H:1970:A:C6	2.86	0.44
26:1H:713:G:H2'	26:1H:714:U:C6	2.53	0.44
26:1H:828:U:H4'	26:1H:831:G:N1	2.32	0.44
26:14:1996:C:OP1	36:25:31:LYS:HE2	2.17	0.44
31:31:29:ASN:N	31:31:112:MET:HE1	2.30	0.44
31:31:119:ARG:HB3	31:31:119:ARG:CZ	2.46	0.44
1:1G:547:A:OP2	4:32:2:GLY:N	2.51	0.44
37:35:55:ARG:HG2	37:35:56:SER:H	1.82	0.44
31:39:41:LEU:O	31:39:44:ARG:HG2	2.16	0.44
24:3K:38:A:H2'	24:3K:39:U:O4'	2.18	0.44
32:41:107:LEU:HD11	32:41:178:PHE:CE1	2.53	0.44
35:58:95:PRO:O	35:58:96:GLU:CD	2.56	0.44
33:59:10:PRO:HD2	33:59:50:VAL:O	2.17	0.44
10:1I:49:VAL:CG2	14:5I:41:ARG:HB2	2.46	0.44
8:72:17:THR:HG23	8:72:65:TYR:CE2	2.52	0.44
9:82:5:TYR:HE1	9:82:16:ARG:HG2	1.83	0.44
41:B8:108:ARG:HA	41:B8:111:ARG:CZ	2.48	0.44
1:1G:1454:G:H5"	20:BA:35:THR:HG21	2.00	0.44
20:BI:25:ARG:O	20:BI:29:LYS:HG3	2.17	0.44
26:1H:2331:G:H4'	48:I8:42:GLY:HA3	1.98	0.44
50:K8:42:GLY:C	50:K8:44:LEU:N	2.70	0.44
55:P8:5:TRP:NE1	55:P8:7:PRO:HG3	2.33	0.44
37:78:62:LEU:O	56:Q8:13:ARG:HD3	2.17	0.44
1:13:1064:G:H4'	1:13:1065:U:OP1	2.17	0.44
1:13:20:U:H2'	1:13:21:G:O4'	2.18	0.44
1:13:22:G:H2'	1:13:23:C:C6	2.53	0.44
1:13:857:C:H2'	1:13:858:G:O4'	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1536:A:C8	26:14:1537:C:H1'	2.53	0.44
26:14:1669:A:H5"	26:14:1670:C:OP2	2.18	0.44
26:14:19:C:H2'	26:14:20:C:H6	1.83	0.44
26:14:2138:C:H42	26:14:2153:G:H22	1.65	0.44
26:14:2064:C:H1'	26:14:2450:A:C2	2.53	0.44
26:14:24:G:H2'	26:14:25:U:O4'	2.17	0.44
26:14:873:G:C2	26:14:905:U:O2	2.71	0.44
29:19:34:VAL:C	29:19:35:LYS:HE2	2.39	0.44
21:1B:5:ASP:O	21:1B:11:GLY:HA3	2.17	0.44
2:1E:28:PHE:CD2	2:1E:190:THR:HA	2.53	0.44
2:1E:87:ARG:NH1	2:1E:223:ILE:HD11	2.33	0.44
1:1G:1277:C:O2'	1:1G:1279:A:H1'	2.18	0.44
1:1G:1399:C:C2	1:1G:1502:A:N6	2.86	0.44
1:1G:513:C:H42	1:1G:538:G:H1	1.66	0.44
1:1G:713:G:H2'	1:1G:714:G:C8	2.53	0.44
1:1G:722:A:H5"	1:1G:723:U:OP2	2.17	0.44
1:1G:865:A:H8	1:1G:865:A:O5'	1.99	0.44
26:1H:1000:A:H62	26:1H:1154:G:H2'	1.83	0.44
26:1H:1446:C:H2'	26:1H:1447:G:C8	2.53	0.44
26:1H:1729:A:C8	26:1H:1731:G:C8	3.05	0.44
26:1H:1786:A:H1'	26:1H:1938:A:N6	2.33	0.44
26:1H:382:G:H5"	26:1H:383:U:OP2	2.17	0.44
26:1H:978:G:C2	26:1H:986:C:C2	3.06	0.44
10:1I:77:PRO:HG2	10:1I:79:ARG:HH12	1.82	0.44
4:32:19:LEU:HB2	4:32:21:LEU:HD22	2.00	0.44
4:32:31:CYS:HB2	4:32:35:ARG:HH12	1.82	0.44
37:35:46:LYS:HE2	37:35:46:LYS:HB3	1.62	0.44
4:3E:22:LYS:HB2	4:3E:26:CYS:SG	2.57	0.44
12:3I:123:LYS:H	12:3I:123:LYS:HG2	1.65	0.44
24:3K:35:U:H2'	24:3K:36:U:H6	1.82	0.44
32:41:27:ASN:HB3	32:41:30:GLU:HG3	1.99	0.44
32:41:83:ARG:H	32:41:86:MET:HE1	1.82	0.44
33:59:81:GLU:HG3	33:59:83:TYR:HB2	1.98	0.44
36:68:4:PRO:O	36:68:5:GLN:HB2	2.18	0.44
8:72:41:ARG:HE	8:72:41:ARG:HB2	1.27	0.44
41:75:54:ARG:HG3	41:75:59:THR:CG2	2.47	0.44
1:13:624:C:H4'	16:7I:11:SER:N	2.32	0.44
16:7I:3:LYS:O	16:7I:21:VAL:HA	2.17	0.44
16:7I:74:LEU:HA	16:7I:77:ALA:HB2	1.99	0.44
42:85:43:GLY:HA3	43:95:73:SER:OG	2.18	0.44
42:C8:30:LYS:HD3	42:C8:30:LYS:HA	1.85	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
44:E8:17:VAL:HG13	44:E8:76:VAL:HG11	1.99	0.44
49:J8:87:PRO:HA	49:J8:90:ILE:HD12	1.99	0.44
29:11:126:GLN:HG2	29:11:127:VAL:H	1.83	0.44
2:12:145:LEU:O	2:12:149:LEU:HB2	2.18	0.44
2:12:36:ARG:N	2:12:38:GLY:O	2.50	0.44
1:13:1342:C:H2'	1:13:1343:G:C8	2.53	0.44
1:13:192:U:C1'	20:BI:103:GLY:HA2	2.47	0.44
1:13:34:C:H2'	1:13:35:G:C8	2.53	0.44
26:14:1115:G:H2'	26:14:1116:C:C6	2.53	0.44
26:14:1163:G:H2'	26:14:1164:G:H8	1.82	0.44
26:14:1260:G:H2'	26:14:1261:C:C6	2.53	0.44
26:14:1659:U:C4	26:14:1660:C:C5	3.05	0.44
26:14:2115:G:H1'	26:14:2171:A:H61	1.82	0.44
26:14:2394:C:C2'	26:14:2395:C:H5'	2.48	0.44
26:14:270(M):U:H4'	26:14:270(N):G:O5'	2.18	0.44
26:14:442:G:C6	26:14:444:C:N4	2.86	0.44
26:14:259:G:N2	26:14:621:A:H8	2.07	0.44
26:14:634:C:H2'	26:14:635:C:H6	1.82	0.44
26:14:736:C:H5"	62:14:3926:HOH:O	2.17	0.44
26:14:868:U:N3	26:14:869:G:N7	2.65	0.44
35:15:120:LEU:HG	35:15:122:VAL:HG23	1.98	0.44
29:19:6:PHE:HE1	29:19:18:VAL:HG23	1.82	0.44
1:1G:1013:G:O2'	1:1G:1014:A:N7	2.35	0.44
1:1G:1109:C:H2'	1:1G:1110:A:O4'	2.18	0.44
1:1G:129(A):G:O2'	1:1G:189:U:OP1	2.24	0.44
1:1G:373:A:N3	1:1G:374:A:C8	2.86	0.44
1:1G:434:U:H2'	1:1G:435:C:C6	2.53	0.44
1:1G:440:A:H3'	1:1G:442:C:C6	2.53	0.44
1:1G:516:U:O2'	1:1G:519:C:N3	2.51	0.44
1:1G:573:A:H5'	1:1G:573:A:H8	1.83	0.44
26:1H:1055:G:H1	26:1H:1104:C:N4	2.10	0.44
26:1H:1425:G:H2'	26:1H:1426:G:O4'	2.17	0.44
26:1H:1482:U:O4	26:1H:1510:A:H1'	2.17	0.44
26:1H:1590:U:H2'	26:1H:1591:G:H8	1.82	0.44
26:1H:1675:C:H2'	26:1H:1676:A:O4'	2.18	0.44
26:1H:1858:G:HO2'	26:1H:1859:A:P	2.41	0.44
26:1H:2135:A:C6	26:1H:2136:C:C2	3.06	0.44
26:1H:2111:C:C4	26:1H:2145:C:C2	3.06	0.44
26:1H:2396:G:H5"	49:J8:25:LYS:NZ	2.33	0.44
26:1H:484:C:H2'	26:1H:485:C:C6	2.53	0.44
26:1H:784:A:O4'	29:11:227:ASN:ND2	2.51	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:2K:20:G:OP1	23:2K:61:U:N3	2.48	0.44
1:13:1060:C:OP1	14:5I:45:ARG:NH2	2.50	0.44
34:61:40:THR:O	34:61:44:LEU:HB2	2.17	0.44
40:65:95:HIS:N	40:65:99:LYS:HB2	2.33	0.44
28:71:194:ARG:NH2	28:71:226:PRO:O	2.47	0.44
8:72:19:VAL:HG23	8:72:21:LYS:HB3	2.00	0.44
16:7I:12:LYS:C	16:7I:14:ASN:H	2.21	0.44
40:A8:7:TYR:HA	40:A8:10:ARG:NH2	2.32	0.44
19:AI:37:ARG:HG3	19:AI:37:ARG:H	1.35	0.44
47:D5:139:VAL:HG21	47:D5:155:LEU:HD22	2.00	0.44
47:D5:24:LEU:HD12	47:D5:25:PRO:O	2.18	0.44
42:C8:92:ARG:NE	43:D8:11:GLN:H	2.16	0.44
2:12:27:LYS:HE3	2:12:194:PRO:HD2	2.00	0.44
1:13:1144:G:H21	1:13:1146:A:H62	1.64	0.44
1:13:141:A:H2'	1:13:142:G:C8	2.51	0.44
1:13:516:U:C4	1:13:517:G:C6	3.06	0.44
26:14:150:C:H2'	26:14:151:C:C6	2.53	0.44
26:14:196:A:OP2	37:35:46:LYS:NZ	2.50	0.44
26:14:747:U:O2	26:14:2014:A:H1'	2.17	0.44
26:14:2165:G:H3'	26:14:2166:G:H5'	2.00	0.44
26:14:2543:G:H2'	26:14:2544:G:C8	2.53	0.44
26:14:877:U:O4	26:14:899:A:N6	2.51	0.44
26:14:947:G:H2'	26:14:948:G:C8	2.52	0.44
29:19:245:PRO:HA	29:19:246:PRO:HD3	1.92	0.44
26:14:1797:C:O2'	29:19:259:THR:OG1	2.28	0.44
2:1E:125:PRO:HA	2:1E:127:ILE:HG12	2.00	0.44
1:1G:135:C:C2	16:7A:1:MET:HB3	2.52	0.44
1:1G:1251:A:HO2'	1:1G:1369:C:HO2'	1.63	0.44
1:1G:1378:C:H3'	1:1G:1379:G:H5"	2.00	0.44
1:1G:1404:C:H2'	1:1G:1405:G:C8	2.53	0.44
1:1G:561:U:O2'	1:1G:562:C:OP2	2.35	0.44
1:1G:567:G:H1'	62:1G:1897:HOH:O	2.18	0.44
1:1G:580:U:H2'	1:1G:581:G:O4'	2.17	0.44
1:1G:964:A:N3	1:1G:969:A:O2'	2.41	0.44
26:1H:1639:U:H5"	62:1H:4619:HOH:O	2.17	0.44
26:1H:1693:U:O2'	29:11:14:ARG:NH2	2.51	0.44
26:1H:2134:A:O2'	26:1H:2159:G:N2	2.50	0.44
10:1I:57:LYS:O	10:1I:60:ARG:NH2	2.48	0.44
27:1J:87:G:N2	27:1J:89:G:H3'	2.33	0.44
3:22:180:ALA:O	3:22:181:ASN:HB3	2.18	0.44
3:22:88:ARG:HA	3:22:91:LEU:HD13	1.98	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:3L:13:C:H2'	24:3L:14:A:C8	2.53	0.44
38:45:25:ASP:HA	38:45:102:VAL:HG23	2.00	0.44
38:45:97:VAL:HG21	38:45:103:MET:HE2	2.00	0.44
38:45:114:ALA:O	38:45:118:LEU:HB2	2.18	0.44
26:14:2469:A:O2'	38:45:56:ARG:HG2	2.18	0.44
33:51:10:PRO:C	33:51:11:VAL:HG22	2.39	0.44
7:62:15:ASP:OD1	7:62:16:LEU:N	2.51	0.44
7:62:45:ASP:O	7:62:49:ILE:HG13	2.18	0.44
40:65:102:ALA:HA	40:65:105:ALA:HB3	1.99	0.44
28:71:196:LEU:HD12	28:71:196:LEU:HA	1.66	0.44
16:7A:1:MET:HE3	16:7A:1:MET:HB2	1.51	0.44
9:82:111:ARG:HB2	9:82:113:LYS:HE2	1.99	0.44
19:AA:41:VAL:HG23	19:AA:43:GLU:H	1.82	0.44
20:BA:73:HIS:CB	20:BA:74:LYS:HZ2	2.31	0.44
26:14:2354:G:O2'	48:E5:36:ILE:HG23	2.18	0.44
48:E5:43:THR:HG23	48:E5:46:LYS:HE2	2.00	0.44
45:F8:66:LEU:HD22	45:F8:66:LEU:HA	1.80	0.44
47:H8:44:PHE:CE2	47:H8:86:VAL:HG11	2.53	0.44
32:41:101:ILE:HG13	52:M8:25:TYR:O	2.17	0.44
1:13:1095:U:H5'	1:13:1109:C:O2	2.17	0.43
1:13:1256:A:HO2'	1:13:1257:U:P	2.40	0.43
1:13:1505:G:OP1	62:13:1842:HOH:O	2.21	0.43
1:13:67:C:O2'	1:13:171:A:N3	2.45	0.43
1:13:605:U:H2'	1:13:606:G:O4'	2.18	0.43
1:13:652:U:C4	1:13:752:G:N3	2.86	0.43
26:14:108:U:H2'	26:14:109:G:C8	2.53	0.43
26:14:1154:G:P	42:85:58:ARG:HH11	2.41	0.43
26:14:350:U:H2'	26:14:351:G:O4'	2.18	0.43
26:14:452:G:N3	26:14:457:A:H2	2.15	0.43
26:14:493:G:H2'	26:14:494:G:O4'	2.18	0.43
26:14:51:G:N3	26:14:119:A:C2	2.86	0.43
26:14:70:G:H21	26:14:71:A:H62	1.66	0.43
29:19:71:ASP:OD2	29:19:103:ARG:NH2	2.49	0.43
1:1G:42:G:H2'	1:1G:43:C:O4'	2.17	0.43
26:1H:129:C:H2'	26:1H:130:C:H6	1.83	0.43
26:1H:1515:C:H2'	26:1H:1516:U:H6	1.82	0.43
26:1H:164:U:H5'	26:1H:165:U:OP2	2.18	0.43
26:1H:2058:A:H5"	26:1H:2059:A:OP2	2.18	0.43
26:1H:2211:G:O2'	26:1H:2212:A:P	2.76	0.43
26:1H:2704:C:H2'	26:1H:2705:A:C8	2.53	0.43
26:1H:2771:C:H2'	26:1H:2772:C:C6	2.53	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2810:A:H2'	26:1H:2811:G:O4'	2.18	0.43
26:1H:317:G:C2	26:1H:318:C:C2	3.06	0.43
26:1H:53:A:H2'	26:1H:54:G:O4'	2.18	0.43
27:1J:65:C:H41	27:1J:108:C:C2'	2.31	0.43
30:21:111:ARG:HG2	30:21:111:ARG:H	1.58	0.43
30:21:32:PRO:HD2	30:21:50:GLY:O	2.17	0.43
3:22:181:ASN:ND2	3:22:204:LEU:HB2	2.33	0.43
4:32:13:ARG:HG2	4:32:13:ARG:H	1.53	0.43
4:3E:85:LYS:CE	4:3E:89:THR:HA	2.46	0.43
5:42:118:ILE:HG12	5:42:119:LEU:H	1.82	0.43
5:42:60:TYR:HB3	5:42:64:ARG:NH2	2.33	0.43
38:45:57:HIS:CG	38:45:117:ALA:HB2	2.52	0.43
5:4E:100:VAL:HG22	5:4E:115:VAL:HG12	2.00	0.43
1:1G:1503:A:H1'	25:4L:12:A:H61	1.83	0.43
6:52:15:ASP:O	6:52:19:LEU:HB2	2.17	0.43
33:59:11:VAL:HB	33:59:13:LYS:HD2	2.00	0.43
27:1J:48:A:O2'	40:65:95:HIS:HE1	2.01	0.43
8:72:82:HIS:HE1	8:72:136:GLU:HG3	1.83	0.43
41:75:54:ARG:HA	41:75:59:THR:HB	2.00	0.43
26:1H:805:G:O5'	37:78:41:ARG:HG2	2.18	0.43
37:78:39:LYS:HB2	37:78:45:LEU:HD21	1.99	0.43
9:8E:22:GLY:N	9:8E:58:HIS:O	2.37	0.43
40:A8:110:LEU:O	40:A8:111:GLU:HB2	2.17	0.43
40:A8:88:ASP:O	40:A8:89:ARG:HB3	2.17	0.43
19:AI:65:ASN:OD1	19:AI:65:ASN:N	2.51	0.43
20:BI:16:HIS:O	20:BI:19:SER:HB2	2.18	0.43
44:E8:64:MET:HE3	44:E8:64:MET:HB3	1.83	0.43
1:13:1022:G:H2'	1:13:1023:G:C8	2.54	0.43
1:13:1132:C:H2'	1:13:1133:G:H8	1.82	0.43
1:13:706:A:N3	11:2I:31:THR:HG21	2.33	0.43
1:13:721:G:C6	1:13:733:A:C2	3.07	0.43
26:14:151:C:H2'	26:14:152:G:C8	2.54	0.43
26:14:1839:G:N3	26:14:1839:G:H2'	2.33	0.43
26:14:2299:G:N1	26:14:2318:G:C8	2.86	0.43
26:14:2657:A:O3'	33:59:160:LYS:NZ	2.48	0.43
26:14:2862:G:H2'	26:14:2863:C:C6	2.53	0.43
27:16:75:G:H21	47:H8:85:HIS:HE1	1.64	0.43
1:1G:1391:U:H2'	1:1G:1392:G:C8	2.54	0.43
1:1G:1478:C:H2'	1:1G:1479:C:C6	2.52	0.43
1:1G:265:G:H5'	17:8A:64:PRO:O	2.18	0.43
1:1G:536:C:H2'	1:1G:537:G:C8	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:980:C:H5'	1:1G:981:U:C5	2.53	0.43
26:1H:1131:G:O6	26:1H:2040:C:H1'	2.18	0.43
26:1H:1226:G:OP1	43:D8:69:LYS:NZ	2.31	0.43
26:1H:1296:G:OP1	26:1H:2709:G:O2'	2.30	0.43
26:1H:1337:G:C4	26:1H:1338:G:C8	3.07	0.43
26:1H:1538:G:H2'	26:1H:1539:G:C8	2.52	0.43
26:1H:2062:A:C2'	26:1H:2062:A:N3	2.80	0.43
1:1G:1191:A:H5"	3:22:4:LYS:NZ	2.33	0.43
1:1G:676:A:H1'	11:2A:115:PRO:HB3	1.99	0.43
4:32:15:GLU:HG2	4:32:66:ARG:HH11	1.83	0.43
4:3E:165:MET:O	4:3E:168:ARG:HD2	2.18	0.43
4:3E:93:PHE:HA	4:3E:96:LEU:HD22	2.00	0.43
24:3L:15:G:H1	24:3L:48:C:N4	2.16	0.43
32:41:145:THR:HG1	32:41:148:MET:CG	2.31	0.43
32:49:39:ILE:HG23	32:49:157:ILE:HG23	2.00	0.43
13:4A:81:LEU:HD13	13:4A:81:LEU:HA	1.69	0.43
3:2E:6:HIS:CD2	14:5I:49:HIS:HB3	2.52	0.43
7:62:116:ALA:HA	7:62:119:ARG:NE	2.33	0.43
40:65:29:PHE:CD1	40:65:30:ARG:N	2.81	0.43
40:65:49:VAL:HG22	40:65:80:LEU:HD12	2.01	0.43
36:68:98:VAL:HG11	36:68:114:ILE:HG23	2.00	0.43
8:72:29:SER:HB3	8:72:32:LYS:CG	2.45	0.43
41:75:5:ALA:N	41:75:6:LEU:CA	2.81	0.43
41:75:62:THR:HG22	41:75:75:ILE:HG12	1.99	0.43
37:78:18:ARG:HA	37:78:18:ARG:HD2	1.71	0.43
20:BA:74:LYS:HG3	20:BA:75:ASN:H	1.83	0.43
46:G8:41:GLY:HA2	46:G8:64:GLU:OE1	2.18	0.43
46:G8:4:LYS:HD3	46:G8:4:LYS:HA	1.71	0.43
26:1H:2336:A:N6	48:I8:43:THR:HB	2.29	0.43
56:M5:37:SER:OG	56:M5:39:LYS:O	2.32	0.43
1:13:192:U:H1'	20:BI:103:GLY:HA2	1.99	0.43
1:13:343:U:H2'	1:13:344:A:H5"	2.01	0.43
26:14:1176:G:H8	26:14:1177:A:H2	1.66	0.43
26:14:1259:G:H2'	26:14:1260:G:H8	1.83	0.43
26:14:2146:C:H4'	26:14:2147:G:C8	2.53	0.43
26:14:322:A:H3'	31:39:169:ASN:OD1	2.18	0.43
26:14:606:U:H4'	26:14:658:C:H4'	2.00	0.43
26:14:685:A:H1'	26:14:688:U:O4	2.18	0.43
26:14:740:U:H2'	26:14:741:G:C8	2.53	0.43
29:19:70:TRP:O	29:19:73:VAL:HG23	2.18	0.43
2:1E:16:HIS:CE1	2:1E:210:SER:O	2.71	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1E:7:VAL:HG21	2:1E:217:ARG:HH11	1.84	0.43
1:1G:34:C:H2'	1:1G:35:G:C8	2.53	0.43
1:1G:64:G:H4'	1:1G:65:U:O5'	2.17	0.43
1:1G:66:G:C2	1:1G:67:C:C6	3.06	0.43
1:1G:965:A:C2	1:1G:969:A:C2	3.06	0.43
26:1H:1266:G:O2'	26:1H:2012:G:O6	2.28	0.43
26:1H:2345:G:H4'	26:1H:2346:A:O5'	2.17	0.43
26:1H:2663:G:C6	26:1H:2664:G:C4	3.07	0.43
26:1H:654(A):A:C2	26:1H:654(T):A:N1	2.85	0.43
9:8E:115:GLY:HA2	10:1I:58:ASP:OD2	2.17	0.43
22:1K:17:U:H4'	22:1K:60:U:C4	2.53	0.43
30:21:50:GLY:HA2	30:21:77:ILE:O	2.18	0.43
3:22:32:LEU:HB3	3:22:59:ARG:HH12	1.83	0.43
3:2E:16:ARG:HB2	3:2E:16:ARG:HH11	1.84	0.43
11:2I:78:GLN:O	11:2I:103:LEU:HA	2.18	0.43
32:49:109:VAL:HG12	32:49:142:PRO:HG3	2.01	0.43
35:58:94:HIS:C	35:58:95:PRO:O	2.56	0.43
14:5A:21:TYR:HE1	14:5A:23:ARG:HB2	1.83	0.43
1:1G:994:A:C2	14:5A:5:ALA:HB2	2.53	0.43
34:61:50:ARG:HA	34:61:50:ARG:HD3	1.40	0.43
16:7I:49:LEU:HD22	16:7I:73:LEU:HD22	2.01	0.43
38:88:75:THR:HG22	38:88:89:ASN:C	2.38	0.43
17:8I:48:GLU:O	17:8I:50:LYS:HG2	2.18	0.43
43:95:43:GLU:HA	43:95:44:LYS:HA	1.81	0.43
40:A8:110:LEU:HB3	40:A8:111:GLU:H	1.66	0.43
1:13:1320:C:N4	19:AI:36:ARG:HG3	2.33	0.43
47:D5:43:GLU:O	47:D5:47:VAL:HG23	2.18	0.43
44:E8:6:ILE:HA	44:E8:103:ILE:O	2.18	0.43
46:G8:81:LYS:HB2	46:G8:99:CYS:SG	2.58	0.43
26:1H:2361:A:O5'	56:Q8:27:THR:OG1	2.36	0.43
29:11:11:PRO:O	29:11:12:SER:OG	2.33	0.43
29:11:67:PHE:HB3	29:11:153:ALA:H	1.83	0.43
29:11:232:PRO:HB3	29:11:244:ARG:CZ	2.48	0.43
2:12:171:ALA:HA	2:12:174:VAL:HG23	2.01	0.43
2:12:19:HIS:CE1	2:12:206:ASP:HB2	2.54	0.43
1:13:103:C:C2	1:13:104:G:C8	3.07	0.43
26:14:1025:G:C4	26:14:1135:C:H1'	2.54	0.43
26:14:1142:U:O2	26:14:1142:U:H2'	2.18	0.43
26:14:2290:G:O2'	26:14:2381:C:H1'	2.18	0.43
26:14:2494:G:C4	26:14:2495:G:C8	3.06	0.43
26:14:2880:C:H1'	39:55:92:GLY:HA3	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:565:C:H4'	26:14:1253:A:C6	2.53	0.43
1:1G:973:G:C1'	10:1A:55:LYS:HG2	2.48	0.43
1:1G:1007:C:H1'	1:1G:1023:G:H1	1.81	0.43
1:1G:1017:G:H2'	1:1G:1018:C:O4'	2.19	0.43
1:1G:1149:C:H2'	1:1G:1150:U:O4'	2.19	0.43
1:1G:429:U:H1'	1:1G:430:A:H5"	2.01	0.43
1:1G:560:U:H4'	1:1G:561:U:O5'	2.17	0.43
1:1G:744:C:O2'	1:1G:851:G:N2	2.50	0.43
1:1G:927:G:N2	1:1G:1391:U:H1'	2.33	0.43
26:1H:1062:G:H2'	26:1H:1063:G:O4'	2.19	0.43
26:1H:2078:C:H2'	26:1H:2079:U:O4'	2.19	0.43
26:1H:2164:C:H41	26:1H:2165:G:N2	2.15	0.43
26:1H:273(D):C:H2'	26:1H:273(E):U:C6	2.53	0.43
26:1H:389:G:H8	26:1H:389:G:O5'	2.01	0.43
26:1H:644:A:H4'	26:1H:645:C:C5	2.53	0.43
27:1J:19:G:H2'	27:1J:20:C:O4'	2.18	0.43
22:1K:66:A:H5"	22:1K:67:C:C5	2.54	0.43
30:21:64:LYS:HD2	30:21:65:GLY:HA2	2.00	0.43
30:29:81:ILE:HG21	30:29:84:PHE:HD2	1.84	0.43
3:2E:175:LEU:HD21	3:2E:201:TYR:HE2	1.82	0.43
23:2K:33:OMC:HM22	23:2K:34:U:H5'	2.00	0.43
31:31:123:LEU:HD12	31:31:124:LEU:H	1.83	0.43
12:3A:32:PHE:HD1	12:3A:86:ARG:HA	1.84	0.43
1:1G:19:C:H5"	5:42:86:ALA:HB3	2.00	0.43
32:49:33:ARG:HE	32:49:33:ARG:HB2	1.54	0.43
13:4A:62:ASN:N	13:4A:62:ASN:OD1	2.52	0.43
13:4A:77:ASN:O	13:4A:81:LEU:HD23	2.18	0.43
14:5I:3:ARG:HD3	14:5I:3:ARG:O	2.19	0.43
40:65:46:VAL:HG12	40:65:48:LEU:HD12	1.99	0.43
40:65:18:ILE:HD13	40:65:87:PHE:O	2.19	0.43
34:69:76:THR:HA	34:69:105:HIS:NE2	2.34	0.43
15:6I:26:GLU:H	15:6I:26:GLU:HG2	1.44	0.43
1:13:660:G:OP1	15:6I:5:LYS:HD3	2.18	0.43
1:1G:825:G:O2'	8:72:12:ARG:NH1	2.51	0.43
8:7E:121:ASP:OD1	8:7E:121:ASP:N	2.48	0.43
1:13:875:C:O2'	8:7E:14:ARG:NH1	2.50	0.43
1:13:1130:A:O2'	9:8E:3:GLN:OE1	2.22	0.43
47:D5:8:TYR:HA	47:D5:62:PRO:CD	2.48	0.43
45:F8:36:LYS:HE2	45:F8:54:VAL:O	2.18	0.43
50:K8:64:LEU:O	50:K8:68:ARG:HG3	2.18	0.43
26:1H:1158:C:H4'	51:L8:32:GLN:HB2	2.01	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
56:M5:50:LEU:HA	56:M5:50:LEU:HD12	1.52	0.43
2:12:49:GLU:O	2:12:52:GLU:HG3	2.18	0.43
2:12:98:LEU:HD23	2:12:98:LEU:HA	1.66	0.43
1:13:1228:C:H2'	1:13:1229:A:H8	1.83	0.43
1:13:38:G:C2	1:13:397:A:C2	3.06	0.43
1:13:433:C:H2'	1:13:434:U:C6	2.53	0.43
1:13:926:G:H5'	1:13:927:G:O5'	2.17	0.43
26:14:818:G:C2	26:14:1190:G:O6	2.70	0.43
26:14:125:G:H1'	55:L5:13:ALA:HB1	2.01	0.43
26:14:1460:A:H3'	26:14:1461:G:H5'	2.00	0.43
26:14:162:U:H4'	26:14:171:G:C4	2.52	0.43
26:14:1894:C:O2'	26:14:1895:C:H5'	2.19	0.43
26:14:2175:C:N4	26:14:2176:A:N7	2.66	0.43
26:14:648:G:O2'	26:14:2351:G:OP1	2.30	0.43
26:14:738:G:H3'	26:14:739:G:C8	2.53	0.43
26:14:932:G:H4'	26:14:933:A:O5'	2.19	0.43
35:15:96:GLU:OE1	35:15:96:GLU:N	2.37	0.43
21:1B:12:LYS:HB3	21:1B:17:THR:O	2.18	0.43
1:1G:1372:U:H2'	1:1G:1373:G:O4'	2.19	0.43
1:1G:1497:G:C2'	1:1G:1498:U:H5'	2.49	0.43
1:1G:562:C:H4'	1:1G:563:A:O5'	2.18	0.43
26:1H:1059:G:H1	26:1H:1088:A:H8	1.67	0.43
26:1H:1498:C:O4'	26:1H:1577:C:H4'	2.18	0.43
26:1H:528:A:C2	26:1H:2043:C:H4'	2.53	0.43
26:1H:2079:U:O3'	49:J8:35:THR:HB	2.18	0.43
26:1H:2259:G:C2	26:1H:2282:G:N1	2.87	0.43
26:1H:2272:U:H5"	26:1H:2273:A:OP1	2.17	0.43
26:1H:2287:A:N1	26:1H:2346:A:C2	2.87	0.43
26:1H:600:G:H2'	26:1H:601:C:C6	2.54	0.43
26:1H:757:U:H2'	26:1H:758:C:O4'	2.18	0.43
30:21:47:VAL:O	30:21:80:GLU:HA	2.18	0.43
3:22:9:GLY:HA2	3:22:12:LEU:HG	2.00	0.43
36:25:1:MET:O	36:25:2:ILE:HD13	2.19	0.43
11:2A:109:VAL:HG13	18:9A:86:VAL:HG22	2.00	0.43
3:2E:178:LEU:HD13	3:2E:178:LEU:HA	1.78	0.43
24:3K:13:C:H42	24:3K:22:G:N2	2.17	0.43
38:45:102:VAL:O	38:45:102:VAL:HG12	2.19	0.43
32:49:136:ARG:H	32:49:136:ARG:HG3	1.57	0.43
5:4E:35:GLY:HA3	5:4E:112:LEU:HB3	1.99	0.43
26:14:1653:G:C6	39:55:9:LYS:HB3	2.53	0.43
1:1G:1202:G:O2'	14:5A:27:CYS:HB2	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:62:63:LYS:O	7:62:67:GLU:N	2.47	0.43
36:68:60:ALA:HB1	36:68:84:ALA:HB1	2.01	0.43
7:6E:15:ASP:OD2	7:6E:18:TYR:N	2.48	0.43
15:6I:69:TYR:CE1	15:6I:73:GLU:HG3	2.53	0.43
9:82:118:LYS:HG2	9:82:118:LYS:O	2.19	0.43
42:85:100:VAL:O	42:85:101:ARG:HG2	2.18	0.43
41:B8:4:GLY:HA2	41:B8:7:ILE:CG1	2.46	0.43
20:BA:84:LEU:HD23	20:BA:84:LEU:HA	1.85	0.43
20:BI:57:ARG:NH2	20:BI:100:ILE:HD13	2.33	0.43
20:BI:36:LEU:HD12	20:BI:55:ILE:HD12	2.01	0.43
42:C8:50:ARG:HH22	43:D8:72:VAL:HG22	1.83	0.43
47:D5:99:TYR:HA	47:D5:124:ILE:O	2.18	0.43
44:E8:57:ASN:O	44:E8:61:ASN:HB2	2.17	0.43
49:F5:84:GLY:CA	49:F5:85:LEU:HB3	2.49	0.43
50:G5:18:PRO:HB2	50:G5:68:ARG:NH2	2.34	0.43
52:M8:60:GLN:HG3	52:M8:61:ARG:NH1	2.33	0.43
53:N8:42:PRO:HB2	53:N8:43:HIS:ND1	2.33	0.43
2:12:54:THR:HA	2:12:57:PHE:CD2	2.53	0.43
1:13:256:U:H2'	1:13:257:G:H8	1.81	0.43
1:13:49:U:C2	1:13:361:G:N2	2.87	0.43
1:13:942:G:C2	1:13:1342:C:C2	3.06	0.43
26:14:1716:U:H2'	26:14:1717:G:H8	1.82	0.43
26:14:1885:A:H3'	26:14:1886:C:C6	2.54	0.43
26:14:2129:C:H5"	26:14:2130:U:H5	1.84	0.43
26:14:2129:C:H5'	26:14:2130:U:OP2	2.19	0.43
26:14:2197:U:H1'	26:14:2198:A:C8	2.53	0.43
26:14:2255:G:C6	26:14:2256:G:C5	3.06	0.43
26:14:2461:C:H2'	26:14:2462:U:H6	1.84	0.43
26:14:2552:U:H2'	26:14:2554:U:H5"	2.00	0.43
26:14:2781:A:H5"	26:14:2782:G:H5'	1.99	0.43
26:14:433:C:C4	26:14:434:U:O4	2.72	0.43
35:15:91:LEU:O	35:15:95:PRO:HB3	2.19	0.43
29:19:134:ARG:NH1	29:19:188:GLU:OE2	2.46	0.43
2:1E:187:LEU:HD23	2:1E:201:ILE:HG22	2.01	0.43
1:1G:241:C:H42	1:1G:285:G:H1	1.67	0.43
1:1G:973:G:H5"	1:1G:974:A:C5'	2.48	0.43
26:1H:130:C:O3'	26:1H:1349:A:H1'	2.19	0.43
26:1H:1465:G:C4	26:1H:1466:G:C8	3.05	0.43
26:1H:1825:A:H2'	26:1H:1826:G:C8	2.53	0.43
26:1H:189:G:H2'	26:1H:205:G:N2	2.33	0.43
26:1H:2171:A:O2'	26:1H:2172:U:O5'	2.36	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:654(O):G:H5"	26:1H:654(P):G:C2	2.54	0.43
26:1H:900:A:H3'	26:1H:901:A:C8	2.49	0.43
24:1L:68:G:N3	24:1L:69:A:N6	2.66	0.43
30:21:101:ARG:HB3	30:21:201:THR:OG1	2.18	0.43
30:21:37:ARG:HD2	30:21:42:ASP:CG	2.39	0.43
30:29:202:LYS:N	30:29:202:LYS:HD2	2.34	0.43
11:2A:59:TYR:O	11:2A:62:GLN:HB3	2.18	0.43
3:2E:43:LEU:O	3:2E:47:LEU:HB2	2.18	0.43
3:2E:50:ALA:HB2	3:2E:75:VAL:HB	2.01	0.43
23:2K:65:G:H2'	23:2K:66:C:O4'	2.18	0.43
4:32:57:ARG:NH2	4:32:205:GLU:OE1	2.51	0.43
32:41:173:LEU:HD13	32:41:173:LEU:HA	1.82	0.43
32:41:66:GLN:OE1	32:41:98:ARG:NH1	2.51	0.43
32:49:88:ILE:HD12	32:49:88:ILE:HA	1.82	0.43
13:4A:67:GLU:HG3	13:4A:68:GLY:H	1.83	0.43
6:52:23:LYS:HB3	6:52:23:LYS:HE2	1.76	0.43
35:58:67:LEU:HA	35:58:87:LEU:HD13	2.01	0.43
40:65:89:ARG:O	40:65:90:GLY:C	2.57	0.43
34:69:51:ILE:HD13	34:69:51:ILE:HA	1.90	0.43
1:1G:742:G:P	15:6A:35:ARG:HH22	2.41	0.43
7:6E:104:LEU:HD13	7:6E:104:LEU:HA	1.80	0.43
8:7E:86:ILE:HG12	8:7E:135:CYS:HA	2.00	0.43
8:7E:45:ILE:HD12	8:7E:47:GLY:HA2	1.99	0.43
42:85:74:LEU:HD11	42:85:110:VAL:HG13	1.99	0.43
1:13:1320:C:H42	19:AI:36:ARG:HG3	1.83	0.43
42:C8:14:HIS:O	42:C8:18:LEU:HD12	2.18	0.43
42:C8:58:ARG:HH11	42:C8:93:LYS:HE2	1.84	0.43
47:D5:15:PRO:HB2	47:D5:19:ARG:NH2	2.33	0.43
47:D5:52:SER:O	47:D5:54:HIS:N	2.51	0.43
48:E5:11:ARG:HB2	48:E5:11:ARG:HE	1.68	0.43
40:A8:20:ARG:NH2	48:I8:51:VAL:O	2.52	0.43
48:I8:45:PHE:O	48:I8:59:LEU:HD11	2.18	0.43
49:J8:87:PRO:HB2	49:J8:91:LYS:HZ3	1.83	0.43
54:O8:19:ARG:HG3	54:O8:21:TYR:CE1	2.47	0.43
29:11:120:GLY:O	29:11:123:ALA:HB3	2.19	0.43
2:12:220:ASP:H	2:12:222:ILE:HG13	1.83	0.43
1:13:1120:G:H2'	1:13:1121:U:H6	1.83	0.43
1:13:1124:G:O2'	1:13:1145:C:C4	2.71	0.43
1:13:198:G:C2	1:13:199:G:C8	3.07	0.43
1:13:266:G:H5"	1:13:267:C:H5	1.83	0.43
1:13:428:G:O4'	1:13:430:A:C8	2.72	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:509:A:H5'	4:3E:54:TYR:CD2	2.54	0.43
1:13:600:C:H2'	1:13:601:C:C6	2.54	0.43
1:13:859:A:H2'	1:13:860:A:O4'	2.19	0.43
1:13:947:G:H2'	1:13:948:C:C6	2.53	0.43
26:14:1420:U:HO2'	26:14:1421:G:P	2.41	0.43
26:14:1677:A:H2'	26:14:1678:G:C8	2.54	0.43
26:14:1885:A:H3'	26:14:1886:C:H6	1.84	0.43
26:14:191:A:N1	62:14:3718:HOH:O	2.36	0.43
26:14:2056:G:C2	26:14:2057:A:C8	3.06	0.43
26:14:2067:G:O2'	26:14:2069:G:H5"	2.19	0.43
26:14:2629:A:H4'	26:14:2630:G:H5'	2.01	0.43
26:14:479:A:N3	26:14:481:G:H5"	2.34	0.43
26:14:630:G:N2	26:14:633:A:OP2	2.45	0.43
26:14:675:A:C4	26:14:804:A:C2	3.07	0.43
26:14:962:G:H2'	26:14:963:U:C6	2.54	0.43
27:16:24:G:N7	27:16:56:G:H2'	2.33	0.43
29:19:74:GLY:O	29:19:76:PRO:HD3	2.18	0.43
1:1G:109:A:H2'	1:1G:326:G:N2	2.33	0.43
1:1G:1324:A:C4'	1:1G:1362:C:H4'	2.40	0.43
1:1G:1402:C:H2'	1:1G:1403:C:O4'	2.18	0.43
1:1G:1:U:H4'	1:1G:630:G:H21	1.84	0.43
1:1G:519:C:H2'	1:1G:520:A:O4'	2.18	0.43
1:1G:756:C:N4	62:1G:1906:HOH:O	2.52	0.43
26:1H:1101:U:H2'	26:1H:1102:C:C6	2.54	0.43
26:1H:999:U:H5"	26:1H:1154:G:O6	2.19	0.43
26:1H:1728:G:C6	26:1H:1730:U:H5"	2.53	0.43
26:1H:2301:C:H2'	26:1H:2302:G:H8	1.84	0.43
26:1H:241:A:H5'	26:1H:243:U:O4'	2.18	0.43
26:1H:2468:G:N3	26:1H:2468:G:O4'	2.51	0.43
26:1H:2801:A:H2'	26:1H:2802:G:H8	1.84	0.43
26:1H:280:C:C2	26:1H:361:G:C2	3.07	0.43
26:1H:500:G:N1	26:1H:503:A:OP2	2.48	0.43
26:1H:906:G:OP1	38:88:26:TYR:OH	2.27	0.43
27:1J:7:G:H4'	40:65:29:PHE:CE2	2.54	0.43
30:21:38:THR:O	30:21:42:ASP:N	2.52	0.43
30:21:59:VAL:HG22	30:21:60:ASN:H	1.83	0.43
36:25:88:ASN:HB3	36:25:94:ARG:HD3	2.00	0.43
30:29:65:GLY:O	30:29:68:ALA:HB2	2.17	0.43
23:2L:16:C:O2'	23:2L:62:C:OP1	2.29	0.43
31:39:113:ALA:HB1	31:39:186:ILE:HG21	2.01	0.43
24:3L:50:C:H2'	24:3L:51:A:C8	2.51	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
24:3L:59:A:N3	24:3L:59:A:H2'	2.32	0.43
32:41:173:LEU:HD12	32:41:178:PHE:CE2	2.54	0.43
39:55:35:THR:HG21	39:55:100:LEU:HD11	1.99	0.43
7:6E:15:ASP:OD1	7:6E:16:LEU:N	2.51	0.43
7:6E:79:ARG:NH2	24:3K:33:U:O2'	2.51	0.43
41:75:51:ARG:HD3	41:75:100:TYR:OH	2.19	0.43
16:7I:37:GLY:HA3	16:7I:50:LYS:O	2.19	0.43
43:95:15:GLU:HG3	43:95:16:PRO:HD2	2.00	0.43
43:95:21:ARG:HB3	43:95:91:TYR:HD2	1.84	0.43
26:14:751:A:H5'	44:A5:90:ARG:HA	1.99	0.43
45:B5:24:GLY:O	45:B5:83:VAL:HG12	2.19	0.43
20:BI:55:ILE:HD13	20:BI:55:ILE:HA	1.91	0.43
47:D5:59:LEU:HB2	47:D5:67:LEU:O	2.19	0.43
49:F5:78:LYS:HG2	49:F5:79:GLY:N	2.32	0.43
46:G8:87:LYS:HE3	46:G8:89:PHE:HB3	1.99	0.43
47:H8:69:THR:HA	47:H8:89:PHE:O	2.18	0.43
56:M5:34:TRP:CE3	56:M5:34:TRP:HA	2.54	0.43
1:13:1004:A:H5"	1:13:1024:G:H2'	2.01	0.43
1:13:1117:G:H5"	9:8E:104:ARG:CZ	2.49	0.43
1:13:1126:U:C4	1:13:1127:G:C5	3.07	0.43
1:13:345:C:H41	36:68:116:SER:C	2.17	0.43
1:13:468:A:O2'	16:7I:82:GLN:HG2	2.18	0.43
1:13:542:G:H5'	4:3E:41:GLY:HA3	2.01	0.43
1:13:64:G:H4'	1:13:65:U:H5'	2.00	0.43
1:13:958:A:C6	1:13:959:A:N1	2.87	0.43
26:14:118:A:N3	26:14:178:G:H1'	2.33	0.43
26:14:128:C:H3'	26:14:128:C:C6	2.54	0.43
26:14:1471:A:C2	26:14:1472:A:C4	3.07	0.43
26:14:1753:G:N1	26:14:1756:G:C2	2.86	0.43
26:14:184:C:H2'	26:14:185:U:C6	2.53	0.43
26:14:629:G:H5"	26:14:650:C:O2'	2.19	0.43
26:14:828:U:C5	26:14:2247:A:H4'	2.54	0.43
1:1G:1016:A:O2'	1:1G:1217:C:O2'	2.34	0.43
1:1G:1084:G:H2'	1:1G:1085:U:C6	2.54	0.43
1:1G:983:A:N3	1:1G:983:A:H3'	2.33	0.43
26:1H:2086:U:H2'	26:1H:2087:G:C8	2.54	0.43
26:1H:2213:U:O4'	49:J8:52:ARG:NH2	2.52	0.43
26:1H:2309:A:N6	26:1H:2310:A:N6	2.67	0.43
26:1H:2327:A:H2'	26:1H:2328:A:H8	1.81	0.43
26:1H:2064:C:H1'	26:1H:2450:A:C2	2.53	0.43
26:1H:270(I):G:C2	26:1H:270(R):G:C2	3.06	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:2820:A:O2'	26:1H:2821:A:OP1	2.31	0.43
26:1H:442:G:C4	26:1H:444:C:C5	3.06	0.43
24:1L:37:A:C6	25:4L:19:G:C6	3.07	0.43
3:22:135:LYS:NZ	3:22:139:GLN:HB2	2.34	0.43
3:2E:15:THR:HG22	3:2E:16:ARG:H	1.84	0.43
31:39:160:ASN:HB3	31:39:163:VAL:HB	2.00	0.43
32:41:11:TYR:O	32:41:15:VAL:HB	2.18	0.43
13:4I:11:ARG:HG2	13:4I:12:ASN:N	2.34	0.43
13:4I:65:LYS:HD2	13:4I:69:GLU:CD	2.39	0.43
33:51:129:THR:OG1	33:51:129:THR:O	2.37	0.43
40:65:24:LEU:O	40:65:86:ALA:N	2.51	0.43
28:71:64:LEU:HD11	28:71:188:ASN:HD21	1.84	0.43
8:7E:13:ILE:O	8:7E:17:THR:HG23	2.19	0.43
8:7E:81:HIS:HB2	8:7E:138:TRP:CE3	2.54	0.43
43:95:37:VAL:C	43:95:39:LEU:H	2.22	0.43
40:A8:32:LEU:N	40:A8:32:LEU:HD23	2.34	0.43
40:A8:11:LYS:HD3	40:A8:91:PRO:HD3	1.99	0.43
19:AA:66:MET:HA	19:AA:67:VAL:O	2.19	0.43
45:B5:40:LYS:HA	45:B5:51:VAL:HG11	2.01	0.43
41:B8:16:ARG:NH2	41:B8:19:LEU:HD21	2.34	0.43
46:C5:91:GLU:HG3	46:C5:92:ASN:N	2.34	0.43
47:D5:118:GLN:HB2	47:D5:173:ALA:O	2.19	0.43
47:H8:1:MET:CE	47:H8:135:GLU:HG2	2.49	0.43
55:P8:24:THR:HG23	55:P8:27:GLY:H	1.83	0.43
29:11:105:ILE:HA	29:11:105:ILE:HD12	1.78	0.43
26:1H:1818:U:O4	29:11:154:LYS:HE3	2.19	0.43
1:13:1051:C:H2'	1:13:1052:U:C6	2.54	0.43
1:13:116:A:H61	1:13:313:A:H1'	1.84	0.43
1:13:122:G:O5'	1:13:122:G:H8	2.00	0.43
1:13:411:A:C5	1:13:413:G:H1'	2.54	0.43
35:15:137:LYS:O	35:15:138:LEU:HD23	2.18	0.43
35:15:28:THR:HG22	35:15:29:LYS:N	2.34	0.43
2:1E:150:SER:OG	2:1E:151:GLY:N	2.50	0.43
1:1G:1250:A:H4'	9:82:68:GLY:N	2.33	0.43
1:1G:12:U:H4'	1:1G:526:C:H4'	2.00	0.43
1:1G:1333:A:H2'	1:1G:1334:G:O4'	2.18	0.43
1:1G:1369:C:H2'	1:1G:1370:G:O4'	2.19	0.43
1:1G:192:U:H4'	20:BA:103:GLY:HA2	1.99	0.43
1:1G:616:G:H1'	1:1G:625:G:N2	2.33	0.43
1:1G:678:U:H2'	1:1G:679:C:C6	2.54	0.43
26:1H:1181:C:O2'	26:1H:1182:A:H5'	2.18	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1405:U:H2'	26:1H:1406:U:H6	1.81	0.43
26:1H:183:C:H1'	26:1H:433:C:H1'	2.00	0.43
26:1H:2241:A:H2'	26:1H:2242:G:C8	2.53	0.43
26:1H:2341:G:H2'	26:1H:2342:C:C6	2.54	0.43
26:1H:2712(A):A:OP2	62:1H:3750:HOH:O	2.21	0.43
26:1H:2728:U:H2'	26:1H:2729:G:H8	1.82	0.43
26:1H:633:A:H2'	26:1H:634:C:H5'	2.01	0.43
26:1H:686:G:H8	55:P8:6:GLN:O	2.01	0.43
24:1L:6:G:O2'	24:1L:7:U:OP1	2.33	0.43
3:22:172:ARG:HH21	3:22:174:PRO:HG2	1.84	0.43
30:29:167:VAL:HG11	30:29:189:PRO:HD3	2.01	0.43
37:35:82:GLY:HA3	37:35:115:LEU:HD11	2.01	0.43
12:3A:27:LEU:HG	12:3A:33:ARG:HG2	2.01	0.43
1:1G:362:G:C4'	12:3A:33:ARG:HH21	2.29	0.43
24:3K:69:A:H2'	24:3K:70:C:C6	2.53	0.43
6:5E:24:GLU:HG2	6:5E:28:ARG:NH2	2.33	0.43
34:69:77:LEU:CD2	34:69:78:THR:H	2.30	0.43
28:71:10:LEU:HA	28:71:10:LEU:HD12	1.74	0.43
28:71:46:LYS:HB3	28:71:210:ARG:HB2	2.01	0.43
17:8I:14:LYS:HD2	17:8I:14:LYS:N	2.33	0.43
19:AI:41:VAL:HG21	19:AI:45:VAL:HG23	2.01	0.43
41:B8:16:ARG:NH1	41:B8:18:ASP:OD2	2.52	0.43
20:BA:29:LYS:O	20:BA:33:ILE:HD12	2.18	0.43
42:C8:92:ARG:HD3	42:C8:94:ASN:CB	2.45	0.43
46:G8:71:LYS:HE3	46:G8:71:LYS:HB3	1.71	0.43
47:H8:161:VAL:HG23	47:H8:161:VAL:O	2.19	0.43
49:J8:49:VAL:HG11	49:J8:70:VAL:HG11	2.00	0.43
56:Q8:24:ALA:O	56:Q8:48:PHE:N	2.50	0.43
56:Q8:49:VAL:CG1	56:Q8:49:VAL:O	2.65	0.43
29:11:238:GLY:CA	62:11:311:HOH:O	2.66	0.43
1:13:1126:U:C5	1:13:1127:G:C8	3.07	0.43
1:13:141:A:C2	1:13:142:G:C5	3.07	0.43
1:13:1427:U:H2'	1:13:1428:A:H8	1.84	0.43
1:13:390:C:H2'	1:13:391:G:C8	2.54	0.43
26:14:1156:A:O5'	26:14:1156:A:H8	2.02	0.43
26:14:1199:U:H2'	26:14:1200:C:C6	2.54	0.43
26:14:1475:G:H5'	26:14:1476:C:OP2	2.19	0.43
26:14:1515:C:H2'	26:14:1516:U:H6	1.83	0.43
26:14:1582:C:O2'	26:14:1586:A:C8	2.72	0.43
26:14:2057:A:H2'	26:14:2058:A:C8	2.54	0.43
26:14:2068:U:N3	26:14:2430:A:H2	2.17	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2287:A:C2	26:14:2346:A:C2	3.06	0.43
26:14:2401:U:H2'	26:14:2402:C:H5"	2.00	0.43
26:14:2600:A:H2'	26:14:2601:C:C6	2.54	0.43
26:14:675:A:N6	26:14:676:A:N6	2.67	0.43
26:14:753:C:O2'	26:14:754:C:H5'	2.19	0.43
27:16:44:G:H1'	27:16:47:C:H42	1.84	0.43
2:1E:135:GLN:HG3	2:1E:135:GLN:H	1.65	0.43
2:1E:91:PRO:HB3	2:1E:154:LEU:HB2	2.00	0.43
1:1G:151:A:H2'	1:1G:152:A:O4'	2.18	0.43
1:1G:853:G:H2'	1:1G:854:G:H8	1.83	0.43
1:1G:92:G:H2'	1:1G:93:U:O4'	2.18	0.43
26:1H:1514:U:H2'	26:1H:1515:C:C6	2.53	0.43
26:1H:1800:C:OP1	29:11:266:SER:OG	2.33	0.43
26:1H:2137:C:O2	26:1H:2155:G:N1	2.52	0.43
26:1H:218:A:C2	26:1H:235:U:H4'	2.53	0.43
26:1H:2577:A:H5"	26:1H:2578:G:H5'	2.01	0.43
26:1H:841:A:H2'	26:1H:842:G:C8	2.54	0.43
26:1H:998:C:OP2	42:C8:58:ARG:NH1	2.52	0.43
1:13:1059:C:O2'	10:1I:53:PRO:HD3	2.19	0.43
1:13:973:G:H4'	10:1I:54:PHE:O	2.19	0.43
27:1J:40:U:O2	27:1J:43:C:H5"	2.19	0.43
24:1L:59:A:H5"	24:1L:60:U:H5	1.83	0.43
24:1L:69:A:H1'	24:1L:70:C:O5'	2.18	0.43
3:22:135:LYS:HA	3:22:135:LYS:HD2	1.73	0.43
3:22:7:PRO:O	3:22:11:ARG:HB2	2.19	0.43
30:29:33:VAL:HG13	30:29:47:VAL:HG13	2.00	0.43
30:29:54:GLN:O	30:29:55:ASN:ND2	2.52	0.43
3:2E:32:LEU:HD13	3:2E:59:ARG:HD3	2.01	0.43
23:2L:10:G:N2	23:2L:27:G:H1'	2.34	0.43
24:3L:37:A:H2'	24:3L:38:A:O4'	2.18	0.43
38:45:19:GLY:O	38:45:99:PRO:HD2	2.19	0.43
32:49:56:ALA:HB2	32:49:153:ARG:NE	2.34	0.43
13:4A:68:GLY:HA2	13:4A:71:ARG:NE	2.33	0.43
13:4I:3:ARG:HB2	13:4I:7:VAL:O	2.18	0.43
26:14:2749:A:O4'	33:59:63:SER:HA	2.19	0.43
14:5A:53:LEU:HD23	14:5A:53:LEU:HA	1.67	0.43
15:6I:24:SER:HB3	15:6I:27:VAL:HG23	2.00	0.43
17:8I:13:ASP:HA	17:8I:19:VAL:HG12	2.01	0.43
17:8I:43:LEU:HD12	17:8I:68:ARG:HG2	2.00	0.43
26:14:1223:C:OP2	43:95:88:ARG:NH2	2.52	0.43
40:A8:36:TYR:N	40:A8:36:TYR:CD1	2.87	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:AI:41:VAL:HG13	19:AI:41:VAL:H	1.55	0.43
20:BI:57:ARG:NH1	20:BI:102:GLY:HA2	2.32	0.43
46:C5:50:ARG:HB2	46:C5:58:GLY:O	2.19	0.43
47:D5:10:ARG:HD2	47:D5:36:LYS:HD2	2.00	0.43
38:45:137:TYR:CE2	47:D5:76:LEU:HD22	2.53	0.43
49:F5:88:LYS:HA	49:F5:90:ILE:HG22	2.00	0.43
50:G5:4:SER:H	50:G5:6:VAL:HG13	1.84	0.43
49:J8:25:LYS:HB3	49:J8:25:LYS:HE3	1.72	0.43
1:13:390:C:O3'	16:7I:28:ARG:NH2	2.52	0.42
1:13:406:G:H21	4:3E:119:GLN:HE22	1.67	0.42
1:13:416:G:C5	1:13:417:C:C4	3.06	0.42
1:13:430:A:OP2	4:3E:8:VAL:HG23	2.19	0.42
1:13:715:A:H2'	1:13:716:A:C8	2.54	0.42
26:14:1647:G:P	26:14:1647:G:H3'	2.59	0.42
26:14:2128:C:N3	26:14:2160:G:N2	2.66	0.42
26:14:226:G:H21	26:14:228:A:H62	1.67	0.42
26:14:2508:G:HO2'	26:14:2554:U:HO2'	1.66	0.42
26:14:733:G:O6	26:14:761:A:C8	2.72	0.42
26:14:792:G:H5"	26:14:793:A:H5'	1.99	0.42
29:19:228:PRO:HD3	29:19:235:GLY:CA	2.49	0.42
2:1E:74:LYS:C	2:1E:208:ILE:HG21	2.40	0.42
1:1G:265:G:O3'	17:8A:66:SER:HA	2.18	0.42
1:1G:27:G:H8	1:1G:27:G:O5'	2.02	0.42
1:1G:498:A:H4'	1:1G:500:G:OP1	2.19	0.42
1:1G:591:U:H2'	1:1G:592:G:C8	2.54	0.42
1:1G:890:G:O2'	1:1G:906:G:O6	2.32	0.42
26:1H:1448:G:O2'	26:1H:1529:A:N1	2.42	0.42
26:1H:1581:G:H2'	26:1H:1582:C:C6	2.54	0.42
26:1H:1655:A:H3'	26:1H:1656:C:H6	1.84	0.42
26:1H:1899:G:N2	26:1H:1902:C:H41	2.17	0.42
26:1H:2159:G:H2'	26:1H:2160:G:C8	2.54	0.42
26:1H:2701:C:H2'	26:1H:2702:U:H2'	2.00	0.42
26:1H:2820:A:C8	30:21:109:LYS:HE2	2.54	0.42
26:1H:322:A:H5'	26:1H:340:A:H1'	2.00	0.42
26:1H:839:U:H2'	26:1H:840:C:C6	2.54	0.42
30:21:105:THR:HG22	30:21:106:GLY:H	1.83	0.42
30:21:169:ASN:OD1	30:21:201:THR:HG21	2.19	0.42
30:21:38:THR:HB	30:21:41:LYS:H	1.83	0.42
30:21:79:ARG:HD3	30:21:79:ARG:HA	1.73	0.42
1:1G:1256:A:H3'	3:22:27:LYS:NZ	2.34	0.42
30:29:56:PRO:HD2	30:29:58:ARG:NH2	2.34	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:2K:47:7MG:O2'	23:2K:48:U:C6	2.72	0.42
31:31:127:GLU:OE2	31:31:127:GLU:HA	2.18	0.42
31:31:28:ILE:HA	31:31:112:MET:HE3	2.00	0.42
4:32:126:ILE:HG22	4:32:127:THR:H	1.84	0.42
4:32:98:GLU:OE2	4:32:103:ASN:ND2	2.50	0.42
26:14:587:C:C2	37:35:33:ARG:NH1	2.86	0.42
31:39:46:ARG:O	31:39:48:THR:HG23	2.19	0.42
4:3E:25:ARG:NH1	4:3E:30:LYS:O	2.52	0.42
24:3L:1:G:N3	24:3L:1:G:H2'	2.34	0.42
32:41:113:ARG:HD2	52:M8:33:VAL:HG13	2.01	0.42
32:49:122:PRO:O	32:49:125:PHE:HD2	2.02	0.42
26:1H:2751:G:C6	33:51:3:ARG:CG	3.02	0.42
33:51:9:ILE:HB	33:51:49:VAL:HB	2.00	0.42
34:61:75:LEU:HG	34:61:105:HIS:HD1	1.82	0.42
7:62:114:ARG:H	7:62:114:ARG:HG2	1.30	0.42
34:69:112:LYS:HE2	34:69:112:LYS:HB3	1.87	0.42
7:6E:72:ARG:HG3	7:6E:142:GLU:OE1	2.19	0.42
28:71:19:ILE:HG12	28:71:223:ARG:HD3	2.01	0.42
41:75:64:ARG:HB2	41:75:73:GLU:HG2	2.01	0.42
9:82:97:LYS:HB3	9:82:98:PRO:HD3	2.00	0.42
38:88:37:LEU:HA	38:88:37:LEU:HD23	1.64	0.42
1:13:1116:C:O2'	9:8E:108:VAL:HG21	2.19	0.42
44:A5:58:ALA:HB1	44:A5:64:MET:HB2	2.01	0.42
45:B5:11:PRO:HG2	45:B5:13:LEU:HD21	2.01	0.42
46:C5:19:LYS:HB3	46:C5:20:TYR:H	1.61	0.42
46:C5:17:SER:HB3	46:C5:71:LYS:HB3	2.00	0.42
49:J8:87:PRO:HB2	49:J8:91:LYS:NZ	2.34	0.42
56:Q8:41:ILE:HG13	56:Q8:41:ILE:H	1.62	0.42
29:11:148:GLU:HB2	29:11:151:LYS:HD2	2.01	0.42
1:13:1510:U:H1'	1:13:1526:G:N2	2.34	0.42
1:13:199:G:O6	1:13:218:C:N4	2.52	0.42
1:13:29:G:O2'	1:13:30:U:H5'	2.19	0.42
1:13:658:G:C6	1:13:659:U:C4	3.07	0.42
1:13:687:A:H2'	1:13:701:C:H41	1.85	0.42
1:13:836:G:C6	1:13:851:G:C6	3.07	0.42
1:13:883:C:C2'	1:13:884:U:H5'	2.49	0.42
26:14:1048:A:N6	26:14:1112:G:O2'	2.36	0.42
26:14:1198:U:C2	26:14:1199:U:C5	3.07	0.42
26:14:1354:A:H2'	26:14:1355:G:O4'	2.19	0.42
26:14:1404:C:O2'	26:14:1405:U:H5'	2.19	0.42
26:14:1912:A:C8	26:14:1918:A:C2	3.07	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2345:G:N3	26:14:2381:C:H2'	2.35	0.42
26:14:2611:U:OP2	26:14:2611:U:H3'	2.18	0.42
26:14:221:A:C4	26:14:266:G:N7	2.87	0.42
26:14:2697:G:H2'	26:14:2698:U:O4'	2.18	0.42
26:14:2756:U:H1'	26:14:2757:A:H5"	2.01	0.42
26:14:2780:G:OP2	35:15:118:LYS:HD3	2.19	0.42
26:14:35:G:H2'	26:14:36:G:O4'	2.20	0.42
26:14:483:A:H5"	46:C5:49:VAL:HG13	2.01	0.42
26:14:596:G:H2'	26:14:597:U:O4'	2.19	0.42
26:14:805:G:O4'	37:35:38:GLN:NE2	2.52	0.42
26:14:914:C:N4	26:14:915:C:C2	2.88	0.42
29:19:124:PRO:HG2	29:19:129:ASN:ND2	2.34	0.42
2:1E:79:ASP:N	2:1E:81:VAL:HG22	2.34	0.42
1:1G:1022:G:C6	1:1G:1023:G:C8	3.07	0.42
1:1G:1206:G:H2'	1:1G:1207:G:C8	2.54	0.42
1:1G:1510:U:H2'	1:1G:1511:G:C8	2.55	0.42
1:1G:229:U:H2'	1:1G:230:G:O4'	2.18	0.42
1:1G:539:A:H2'	1:1G:540:G:H8	1.84	0.42
1:1G:854:G:C2	1:1G:855:G:C8	3.07	0.42
26:1H:1316:U:H2'	26:1H:1317:A:H8	1.83	0.42
26:1H:1430:C:H2'	26:1H:1431:U:H6	1.83	0.42
26:1H:1475:G:C4	26:1H:1519:G:N2	2.86	0.42
26:1H:2093:G:C6	26:1H:2225:A:C8	3.08	0.42
26:1H:2302:G:C4	26:1H:2303:G:C8	3.07	0.42
26:1H:273(E):U:C2'	26:1H:273(F):C:H5'	2.49	0.42
26:1H:36:G:C5	26:1H:37:C:C5	3.07	0.42
26:1H:455:C:N3	26:1H:472:A:H2'	2.34	0.42
26:1H:590:A:H2'	26:1H:591:C:C6	2.54	0.42
26:1H:760:G:H4'	26:1H:1776:G:OP1	2.19	0.42
26:1H:840:C:H2'	26:1H:841:A:H8	1.84	0.42
1:13:1199:U:H4'	10:1I:54:PHE:CE2	2.54	0.42
27:1J:0:A:H2'	27:1J:1:U:C6	2.54	0.42
3:22:63:ASN:HA	3:22:98:ASN:HB2	2.01	0.42
11:2A:84:VAL:HG11	11:2A:95:ILE:HD11	2.00	0.42
4:32:150:GLU:C	4:32:152:SER:H	2.21	0.42
4:32:61:LYS:HD2	4:32:206:PHE:CE2	2.54	0.42
31:39:186:ILE:HD12	31:39:192:LEU:HD11	2.01	0.42
5:4E:84:PHE:HB3	5:4E:134:ALA:HB2	2.01	0.42
33:51:27:LYS:HD2	33:51:32:GLU:OE1	2.19	0.42
33:51:84:SER:O	33:51:85:LYS:HB2	2.19	0.42
39:55:32:GLY:HA2	39:55:116:LEU:HD12	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
39:55:45:ARG:HA	39:55:95:THR:HG21	2.00	0.42
34:61:75:LEU:HG	34:61:105:HIS:ND1	2.34	0.42
28:71:14:VAL:HG11	28:71:222:VAL:HG22	2.02	0.42
28:71:44:HIS:O	28:71:212:VAL:HA	2.20	0.42
8:72:39:LEU:HB3	8:72:45:ILE:HG12	2.00	0.42
9:82:63:ILE:HD13	9:82:77:ILE:HG23	2.01	0.42
43:95:21:ARG:HB3	43:95:91:TYR:CD2	2.54	0.42
40:A8:56:LEU:CB	40:A8:58:LEU:HD22	2.49	0.42
41:B8:2:ASN:HD21	41:B8:6:LEU:HD13	1.84	0.42
41:B8:2:ASN:O	41:B8:6:LEU:N	2.48	0.42
41:B8:1:MET:HA	41:B8:3:ARG:H	1.83	0.42
26:1H:2845:G:H5"	41:B8:54:ARG:O	2.19	0.42
41:B8:81:PRO:HG2	41:B8:82:LEU:HD12	2.00	0.42
46:C5:82:PRO:HB3	46:C5:97:ARG:HB3	2.01	0.42
44:E8:107:LEU:HA	44:E8:107:LEU:HD12	1.82	0.42
46:G8:75:ILE:HD12	46:G8:75:ILE:HA	1.61	0.42
49:J8:87:PRO:O	49:J8:91:LYS:HE2	2.19	0.42
2:12:222:ILE:H	2:12:222:ILE:HG13	1.52	0.42
1:13:1240:U:C4	7:6E:32:ARG:HD2	2.54	0.42
1:13:1366:C:H2'	1:13:1367:C:C6	2.52	0.42
1:13:342:C:N4	1:13:343:U:O4	2.52	0.42
1:13:418:C:H2'	1:13:419:C:C6	2.54	0.42
1:13:509:A:H5'	4:3E:54:TYR:HD2	1.83	0.42
1:13:963:G:H21	10:1I:55:LYS:HE2	1.84	0.42
26:14:1022:G:N2	26:14:1142(A):A:H2	2.11	0.42
26:14:1543:A:H1'	26:14:1545:A:H1'	2.00	0.42
26:14:2352:A:H2	48:E5:33:ALA:O	2.02	0.42
26:14:2392:A:H2	26:14:2424:C:N4	2.06	0.42
26:14:2853:C:H2'	26:14:2854:G:C8	2.53	0.42
26:14:2862:G:H2'	26:14:2863:C:H6	1.84	0.42
26:14:601:C:O2	26:14:605:C:H4'	2.19	0.42
26:14:901:A:H2'	26:14:901:A:N3	2.34	0.42
26:14:975:G:C2	26:14:990:A:C8	3.06	0.42
26:14:1007:C:P	35:15:37:LYS:HZ2	2.41	0.42
1:1G:1104:G:C2	1:1G:1105:A:C4	3.07	0.42
26:1H:1062:G:H1'	26:1H:1088:A:C4	2.54	0.42
26:1H:110:G:N7	62:1H:3849:HOH:O	2.37	0.42
26:1H:1213:A:H1'	26:1H:1238:G:N3	2.33	0.42
26:1H:172:C:H2'	26:1H:173:G:H8	1.84	0.42
26:1H:1788:C:H2'	26:1H:1789:A:H8	1.83	0.42
26:1H:1826:G:H2'	26:1H:1827:C:O4'	2.18	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1857:G:C6	26:1H:1858:G:N1	2.88	0.42
26:1H:2394:C:H2'	26:1H:2395:C:H6	1.84	0.42
22:1K:74:C:N4	26:1H:2507:C:O3'	2.53	0.42
26:1H:252:G:O2'	26:1H:253:C:H5'	2.19	0.42
26:1H:263:C:H2'	26:1H:264:C:O4'	2.19	0.42
26:1H:1637:A:H4'	26:1H:2711:A:O2'	2.19	0.42
26:1H:319:C:OP1	31:31:137:LYS:NZ	2.36	0.42
26:1H:902:C:O2'	26:1H:903:C:H5'	2.19	0.42
3:22:152:ILE:HD12	3:22:199:LYS:HD2	2.00	0.42
36:25:122:LEU:HD13	41:75:72:VAL:HG11	2.00	0.42
30:29:52:LEU:O	30:29:75:VAL:N	2.50	0.42
4:32:148:VAL:HG12	4:32:152:SER:CB	2.49	0.42
4:32:172:PRO:HB2	4:32:187:ARG:NH1	2.34	0.42
37:35:133:SER:O	37:35:137:LYS:HG3	2.19	0.42
12:3A:102:ARG:HB3	12:3A:102:ARG:HE	1.63	0.42
12:3I:42:THR:HA	12:3I:53:ARG:O	2.19	0.42
32:41:35:GLU:HG3	32:41:36:LYS:HB2	2.00	0.42
38:45:57:HIS:CE1	38:45:116:GLU:HG2	2.55	0.42
38:45:2:LEU:O	38:45:70:PRO:HG2	2.20	0.42
13:4A:30:ALA:O	13:4A:34:LEU:N	2.39	0.42
13:4A:92:HIS:HD2	13:4A:98:VAL:HG11	1.84	0.42
5:4E:131:ILE:HA	5:4E:131:ILE:HD13	1.89	0.42
33:51:4:ILE:O	33:51:6:ARG:NE	2.51	0.42
14:5I:37:PHE:CE1	14:5I:53:LEU:HD13	2.54	0.42
34:61:114:LEU:HB2	34:61:115:ALA:H	1.73	0.42
34:69:71:ILE:HG22	34:69:72:LEU:HD23	2.00	0.42
15:6A:54:ARG:O	15:6A:58:MET:HG3	2.20	0.42
41:75:12:SER:HA	41:75:15:VAL:HG22	2.00	0.42
8:7E:1:MET:HB3	8:7E:2:LEU:H	1.46	0.42
27:16:90:C:P	38:88:16:ARG:HH21	2.42	0.42
19:AI:41:VAL:HG11	19:AI:45:VAL:HG23	2.01	0.42
1:13:1221:G:O3'	19:AI:77:THR:HG21	2.19	0.42
41:B8:2:ASN:O	41:B8:5:ALA:HB3	2.18	0.42
46:C5:3:VAL:HG11	46:C5:32:PRO:O	2.19	0.42
26:1H:18:C:H4'	42:C8:23:GLY:O	2.20	0.42
47:D5:127:LYS:O	47:D5:162:GLU:HB2	2.19	0.42
47:D5:72:ARG:HD2	47:D5:72:ARG:HA	1.31	0.42
50:G5:38:GLN:O	50:G5:41:ILE:HG12	2.19	0.42
29:11:102:LYS:C	29:11:103:ARG:HG2	2.39	0.42
29:11:119:ALA:CB	29:11:130:ALA:HB3	2.49	0.42
1:13:1121:U:H2'	1:13:1122:U:C6	2.54	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:1397:C:H4'	1:13:1398:A:O5'	2.18	0.42
1:13:1401:G:C2	1:13:1402:C:H1'	2.53	0.42
1:13:439:A:OP2	1:13:493:G:N2	2.46	0.42
1:13:538:G:O3'	12:3I:114:LYS:HD3	2.20	0.42
1:13:711:G:H2'	1:13:712:A:H8	1.84	0.42
1:13:967:C:H4'	9:8E:125:TYR:HE1	1.84	0.42
26:14:1244:G:P	37:35:7:ARG:HD3	2.60	0.42
26:14:1268:A:OP1	62:14:3656:HOH:O	2.21	0.42
26:14:1441:G:H2'	26:14:1442:G:H8	1.84	0.42
26:14:1515:C:H2'	26:14:1516:U:C6	2.54	0.42
26:14:2607:G:O3'	62:14:3658:HOH:O	2.22	0.42
26:14:2543:G:H21	26:14:2646:C:H5"	1.84	0.42
26:14:1999:C:H5"	26:14:2723:C:O2'	2.20	0.42
26:14:813:U:C2	26:14:1195:G:N2	2.87	0.42
29:19:71:ASP:OD1	29:19:71:ASP:N	2.52	0.42
1:1G:1367:C:H5'	10:1A:60:ARG:NH2	2.34	0.42
10:1A:66:ARG:HE	10:1A:66:ARG:HB3	1.63	0.42
2:1E:67:THR:HG21	2:1E:155:LEU:HG	2.01	0.42
1:1G:1173:G:OP1	7:62:5:ARG:NH2	2.47	0.42
1:1G:1279:A:H5"	1:1G:1280:A:OP2	2.20	0.42
1:1G:1442:G:C6	1:1G:1446:A:N6	2.87	0.42
1:1G:834:C:H2'	1:1G:835:U:C6	2.54	0.42
26:1H:1086:A:H1'	26:1H:1103:A:C2	2.54	0.42
26:1H:1386:C:OP2	26:1H:1396:U:H5	2.03	0.42
26:1H:1799:G:H5'	26:1H:1819:A:H61	1.83	0.42
26:1H:2431:U:P	62:1H:3605:HOH:O	2.77	0.42
26:1H:2646:C:H2'	26:1H:2647:U:O4'	2.19	0.42
26:1H:289:A:H2'	26:1H:290:G:O4'	2.20	0.42
1:13:963:G:C2	10:1I:55:LYS:NZ	2.88	0.42
3:2E:130:VAL:HG12	3:2E:134:ILE:HD11	2.02	0.42
3:2E:95:THR:HB	3:2E:97:LYS:HG3	2.01	0.42
12:3I:28:LYS:HD3	12:3I:28:LYS:HA	1.72	0.42
12:3I:47:LYS:HA	12:3I:49:ASN:H	1.84	0.42
32:49:48:GLU:H	32:49:48:GLU:HG2	1.40	0.42
13:4I:15:VAL:HG23	13:4I:43:THR:O	2.19	0.42
33:51:80:SER:O	33:51:81:GLU:HG3	2.19	0.42
35:58:131:GLN:HG2	35:58:131:GLN:H	1.59	0.42
7:62:71:PRO:HG3	7:62:103:TRP:CH2	2.55	0.42
36:68:117:LEU:HA	36:68:117:LEU:HD23	1.84	0.42
34:69:128:LEU:O	34:69:138:ILE:N	2.41	0.42
15:6A:32:LEU:O	15:6A:36:ILE:HG13	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:88:135:ASP:N	38:88:135:ASP:OD1	2.52	0.42
38:88:12:GLN:HG2	38:88:73:PRO:HD2	2.01	0.42
26:1H:996:A:H4'	42:C8:92:ARG:NE	2.34	0.42
26:14:98:G:OP1	50:G5:3:LEU:HB3	2.20	0.42
26:14:2056:G:H1	53:J5:3:LYS:HB3	1.83	0.42
29:11:123:ALA:HB3	29:11:131:LEU:HG	2.02	0.42
1:13:1162:C:O5'	1:13:1162:C:H6	2.02	0.42
1:13:1399:C:H4'	1:13:1400:C:O5'	2.19	0.42
1:13:222:U:H2'	1:13:223:U:C6	2.54	0.42
1:13:376:G:O3'	16:7I:5:ARG:NH1	2.45	0.42
1:13:872:A:C5	1:13:874:G:C8	3.06	0.42
26:14:1121:C:H2'	26:14:1122:G:O4'	2.20	0.42
26:14:1131:G:C2	26:14:1132:A:C4	3.08	0.42
26:14:201:C:H4'	26:14:386:G:C2	2.53	0.42
26:14:2552:U:C2	26:14:2554:U:H5'	2.54	0.42
26:14:2674:G:H4'	36:25:30:ALA:HB2	2.01	0.42
26:14:26:G:C6	26:14:27:G:N1	2.87	0.42
26:14:270(V):G:H2'	26:14:270(W):G:H8	1.83	0.42
26:14:2857:G:N2	26:14:2859:G:H3'	2.35	0.42
26:14:785:G:OP2	62:14:3657:HOH:O	2.22	0.42
26:14:895:U:H4'	26:14:896:A:C2	2.55	0.42
26:14:987:G:OP2	62:14:3655:HOH:O	2.21	0.42
21:1F:9:ARG:O	21:1F:13:ILE:HG13	2.19	0.42
1:1G:1279:A:H5"	1:1G:1280:A:P	2.58	0.42
1:1G:191(E):G:H2'	1:1G:191(F):U:C6	2.55	0.42
1:1G:160:A:H1'	1:1G:344:A:N7	2.35	0.42
1:1G:456:C:H42	1:1G:476:G:H1	1.67	0.42
1:1G:79:G:H1	1:1G:90:C:N4	2.10	0.42
26:1H:1449(A):G:H2'	26:1H:1450:C:C6	2.53	0.42
26:1H:1478:G:H2'	26:1H:1479:G:C8	2.55	0.42
26:1H:2376:A:H2'	26:1H:2377:A:O4'	2.19	0.42
26:1H:2740:A:C6	26:1H:2764:A:C8	3.07	0.42
27:1J:15:A:H1'	27:1J:109:G:C5	2.54	0.42
27:1J:18:G:H2'	27:1J:19:G:C8	2.54	0.42
27:1J:6:C:H2'	27:1J:7:G:O4'	2.19	0.42
11:2A:31:THR:HA	11:2A:42:TRP:HA	2.02	0.42
24:3L:3:G:H2'	24:3L:4:U:O4'	2.18	0.42
38:45:54:MET:O	38:45:57:HIS:N	2.52	0.42
32:49:4:ASP:OD2	32:49:9:ARG:NH1	2.52	0.42
39:55:35:THR:HG23	39:55:113:LEU:HD12	2.00	0.42
6:5E:10:LEU:HD13	6:5E:61:LEU:HD13	2.00	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:62:15:ASP:HB3	7:62:20:ASP:H	1.83	0.42
7:62:97:GLN:HG3	7:62:98:SER:N	2.35	0.42
40:65:102:ALA:O	40:65:105:ALA:N	2.53	0.42
7:6E:15:ASP:CB	7:6E:20:ASP:H	2.32	0.42
41:75:33:LYS:HA	41:75:42:ILE:HD12	2.01	0.42
37:78:113:LYS:HG2	37:78:115:LEU:HD23	2.02	0.42
18:9A:60:ALA:O	18:9A:64:ARG:HG3	2.20	0.42
18:9I:55:ARG:HD2	18:9I:55:ARG:HA	1.91	0.42
26:1H:2378:A:H4'	40:A8:23:ARG:NH1	2.35	0.42
19:AI:25:LYS:HG2	19:AI:27:GLU:OE1	2.20	0.42
20:BA:56:MET:HG3	20:BA:84:LEU:HD13	2.02	0.42
42:C8:83:LEU:HG	42:C8:88:ILE:HB	2.01	0.42
47:D5:17:ALA:O	47:D5:20:ARG:HB2	2.19	0.42
48:E5:69:PHE:CE2	48:E5:79:VAL:HG22	2.55	0.42
49:F5:85:LEU:O	49:F5:88:LYS:N	2.32	0.42
48:I8:36:ILE:C	48:I8:36:ILE:HD13	2.40	0.42
53:J5:40:LYS:HE3	53:J5:44:THR:O	2.19	0.42
56:M5:32:LEU:HA	56:M5:32:LEU:HD12	1.82	0.42
56:M5:32:LEU:O	56:M5:36:LYS:HE3	2.20	0.42
1:13:1284:C:H2'	1:13:1285:A:N7	2.34	0.42
1:13:129(A):G:H4'	1:13:130:A:H5"	2.01	0.42
1:13:1304:G:N2	1:13:1332:A:OP2	2.49	0.42
1:13:186(C):G:H2'	1:13:186(D):C:C6	2.55	0.42
1:13:585:G:O2'	1:13:879:C:H5"	2.19	0.42
1:13:980:C:H2'	1:13:981:U:O4'	2.19	0.42
26:14:1041:C:H42	26:14:1114:G:N2	2.17	0.42
26:14:1385:G:C4	26:14:1386:C:C5	3.08	0.42
26:14:1441:G:H2'	26:14:1442:G:C8	2.55	0.42
26:14:1990:C:H2'	26:14:1991:U:C6	2.55	0.42
26:14:459:U:H2'	26:14:460:A:H8	1.84	0.42
26:14:637:A:OP2	37:35:116:GLY:N	2.50	0.42
26:14:59:U:O2'	26:14:73:A:H2'	2.20	0.42
1:1G:64:G:OP1	1:1G:64:G:H3'	2.19	0.42
1:1G:750:G:H21	15:6A:23:GLY:HA3	1.84	0.42
26:1H:1207:C:H2'	26:1H:1208:C:H6	1.84	0.42
26:1H:1604:C:P	62:1H:3611:HOH:O	2.72	0.42
26:1H:165:U:H2'	26:1H:165:U:H6	1.62	0.42
26:1H:1916:A:H2'	26:1H:1917:U:O4'	2.20	0.42
26:1H:858:U:O2	26:1H:2268:A:H2'	2.19	0.42
26:1H:2529:G:OP2	26:1H:2530:A:H5"	2.20	0.42
26:1H:508:G:N3	26:1H:508:G:H5"	2.35	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:588:U:O4	26:1H:670:A:H1'	2.19	0.42
22:1K:53:G:O2'	22:1K:54:5MU:H5"	2.20	0.42
30:29:117:MET:HA	30:29:122:PHE:H	1.84	0.42
30:29:107:THR:O	30:29:190:GLY:HA2	2.19	0.42
3:2E:34:LEU:HD22	3:2E:38:ARG:NH1	2.34	0.42
23:2L:62:C:H2'	23:2L:63:C:C6	2.50	0.42
4:32:162:LEU:HD12	4:32:181:MET:HG2	2.02	0.42
4:32:58:LEU:HA	4:32:58:LEU:HD23	1.89	0.42
4:3E:196:LEU:H	4:3E:196:LEU:HD12	1.85	0.42
12:3I:70:ILE:HG12	12:3I:100:ILE:HD12	2.01	0.42
24:3L:22:G:N2	24:3L:23:A:C5	2.87	0.42
24:3L:28:U:H3	24:3L:43:U:H3	1.68	0.42
32:49:106:LEU:HG	32:49:111:LEU:HD12	2.01	0.42
35:58:128:HIS:HB2	35:58:129:PRO:HD2	2.01	0.42
33:59:146:ALA:O	33:59:150:ALA:N	2.49	0.42
14:5I:4:LYS:O	14:5I:7:ILE:HG23	2.19	0.42
40:65:83:LYS:HE3	40:65:83:LYS:HB3	1.96	0.42
40:65:89:ARG:O	40:65:92:TYR:N	2.45	0.42
26:14:270(Q):C:H5"	34:69:45:LYS:HE3	2.01	0.42
37:78:106:LEU:O	37:78:107:LYS:C	2.58	0.42
37:78:38:GLN:O	37:78:41:ARG:HB2	2.20	0.42
1:13:112:G:OP1	16:7I:27:LYS:HD2	2.20	0.42
42:85:8:VAL:HB	42:85:12:ARG:HE	1.85	0.42
38:88:112:GLU:CD	38:88:112:GLU:H	2.22	0.42
38:88:35:VAL:HA	38:88:101:ARG:O	2.18	0.42
40:A8:15:ARG:O	40:A8:19:LYS:HD2	2.20	0.42
41:B8:29:ARG:HD3	41:B8:46:GLU:OE2	2.20	0.42
20:BA:66:ALA:HB1	20:BA:71:THR:HB	2.02	0.42
46:C5:2:ARG:HD2	46:C5:2:ARG:HA	1.90	0.42
46:G8:87:LYS:HD2	46:G8:89:PHE:CD2	2.54	0.42
50:K8:22:GLU:OE2	50:K8:68:ARG:NH1	2.51	0.42
53:N8:36:CYS:SG	53:N8:37:LYS:N	2.92	0.42
29:11:70:TRP:CD1	29:11:71:ASP:N	2.87	0.42
29:11:68:LYS:HB3	29:11:70:TRP:CZ3	2.55	0.42
1:13:1126:U:O4	1:13:1127:G:C4	2.72	0.42
1:13:1319:A:O2'	1:13:1323:G:N7	2.47	0.42
1:13:14:U:H2'	1:13:16:A:OP2	2.20	0.42
1:13:233:C:H2'	1:13:234:C:H6	1.84	0.42
1:13:652:U:HO2'	1:13:653:A:P	2.42	0.42
26:14:111:A:H4'	50:G5:69:ARG:NH2	2.33	0.42
26:14:1477:A:H2'	26:14:1478:G:O4'	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2257:U:O2'	26:14:2258:C:H5'	2.19	0.42
26:14:330:A:H2	26:14:1210:A:O2'	1.83	0.42
26:14:717:G:H2'	26:14:718:A:O4'	2.18	0.42
26:14:818:G:O2'	26:14:838:C:O2'	2.28	0.42
1:1G:1181:G:N1	1:1G:1182:G:H1'	2.34	0.42
1:1G:1255:G:H5'	1:1G:1256:A:OP2	2.18	0.42
1:1G:1509:C:H2'	1:1G:1510:U:O4'	2.20	0.42
1:1G:18:C:H6	1:1G:18:C:O5'	2.03	0.42
1:1G:979:C:H5"	1:1G:980:C:OP2	2.18	0.42
26:1H:1021:A:C3'	26:1H:1021:A:C8	3.02	0.42
26:1H:1655:A:H4'	30:21:115:GLY:N	2.34	0.42
26:1H:1988:C:H2'	26:1H:1989:G:O4'	2.19	0.42
26:1H:270(E):G:H1	26:1H:270(U):C:N4	2.16	0.42
26:1H:300:A:H2'	26:1H:334:C:O2'	2.19	0.42
26:1H:484:C:OP1	46:G8:51:VAL:HG22	2.19	0.42
26:1H:28:A:O2'	26:1H:583:G:H5'	2.20	0.42
26:1H:675:A:C8	26:1H:804:A:C6	3.07	0.42
26:1H:960:A:C8	26:1H:962:G:C8	3.07	0.42
10:1I:26:ALA:HA	10:1I:29:ARG:CZ	2.49	0.42
24:1L:35:U:H2'	24:1L:36:U:O4'	2.20	0.42
3:22:7:PRO:O	3:22:11:ARG:NH1	2.53	0.42
36:25:22:ILE:HD13	36:25:22:ILE:HA	1.41	0.42
23:2L:38:A:H2'	23:2L:39:A:O4'	2.18	0.42
37:35:75:ILE:HG13	37:35:77:ARG:NH2	2.35	0.42
31:39:158:THR:HA	31:39:195:ASP:HB2	2.01	0.42
4:3E:196:LEU:HB3	4:3E:197:PRO:HD2	2.02	0.42
32:49:170:ARG:HA	32:49:170:ARG:HD2	1.80	0.42
32:49:36:LYS:HD2	32:49:95:ARG:HH22	1.85	0.42
32:49:62:LEU:HD12	32:49:63:ILE:HG23	2.02	0.42
33:51:169:VAL:HG22	33:51:170:ARG:H	1.84	0.42
6:5E:26:ILE:O	6:5E:30:LEU:HD12	2.20	0.42
36:68:85:VAL:HG11	36:68:114:ILE:HD13	2.02	0.42
28:71:185:LEU:HD23	28:71:185:LEU:HA	1.87	0.42
8:7E:112:LEU:HD12	8:7E:113:SER:N	2.35	0.42
16:7I:8:ARG:HB3	16:7I:28:ARG:NH1	2.34	0.42
26:1H:2483:C:O2	38:88:124:LYS:HE3	2.19	0.42
39:98:113:LEU:HD12	39:98:113:LEU:HA	1.84	0.42
19:AA:3:ARG:HB3	19:AA:7:LYS:CB	2.49	0.42
41:B8:105:LEU:O	41:B8:107:ASP:OD1	2.38	0.42
48:E5:24:LYS:O	48:E5:25:ARG:HD3	2.20	0.42
46:G8:88:LYS:HA	46:G8:88:LYS:HD3	1.78	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:J8:90:ILE:O	49:J8:94:LEU:HB2	2.18	0.42
56:M5:14:VAL:HG11	56:M5:22:VAL:HG13	2.02	0.42
29:11:101:GLU:OE1	29:11:103:ARG:HD3	2.20	0.42
2:12:55:PHE:CD1	2:12:221:LEU:HG	2.55	0.42
2:12:52:GLU:O	2:12:55:PHE:HB2	2.19	0.42
1:13:1009:G:C2	1:13:1021:G:C6	3.08	0.42
1:13:114:U:O2'	1:13:115:G:H5'	2.20	0.42
1:13:1170:A:H2'	1:13:1171:G:O4'	2.20	0.42
1:13:1425:U:H2'	1:13:1426:C:C6	2.55	0.42
1:13:123:C:OP1	1:13:312:C:H5'	2.20	0.42
1:13:345:C:O2'	1:13:346:G:C4	2.69	0.42
1:13:734:G:C6	1:13:735:C:C4	3.08	0.42
1:13:13:U:O2	1:13:914:A:H3'	2.19	0.42
26:14:1024:G:C8	26:14:1025:G:H2'	2.55	0.42
26:14:1022:G:N2	26:14:1142(A):A:C2	2.86	0.42
26:14:1542:G:H3'	26:14:1543:A:H5"	2.01	0.42
26:14:579:G:C8	26:14:2017:U:C4	3.08	0.42
26:14:2051:A:H5'	26:14:2578:G:O4'	2.19	0.42
26:14:219:G:C6	26:14:220:G:C6	3.08	0.42
26:14:2228:G:OP2	29:19:263:ARG:NH2	2.53	0.42
26:14:240:G:O2'	26:14:257:A:N6	2.41	0.42
26:14:2414:G:H21	37:35:67:MET:HE1	1.85	0.42
26:14:2577:A:H2'	26:14:2614:A:N6	2.35	0.42
26:14:2638:G:OP2	30:29:82:ARG:NH2	2.50	0.42
29:19:96:HIS:CD2	29:19:102:LYS:HE2	2.55	0.42
29:19:260:ARG:HH12	29:19:267:SER:HB3	1.84	0.42
21:1B:6:ARG:O	21:1B:12:LYS:HE2	2.20	0.42
1:1G:1208:C:H2'	1:1G:1209:C:H6	1.84	0.42
1:1G:1228:C:H2'	1:1G:1229:A:C8	2.54	0.42
1:1G:1224:G:C2	1:1G:1322:C:H4'	2.55	0.42
1:1G:176:C:H2'	1:1G:177:C:C6	2.54	0.42
1:1G:259:G:H2'	1:1G:260:G:O4'	2.19	0.42
1:1G:689:C:H2'	1:1G:690:G:H5'	2.02	0.42
26:1H:1038:C:H2'	26:1H:1039:G:O4'	2.19	0.42
26:1H:1260:G:C6	26:1H:1261:C:C4	3.08	0.42
26:1H:1263:U:O2'	53:N8:11:THR:HG23	2.20	0.42
26:1H:1543:A:C8	26:1H:1545:A:H5"	2.55	0.42
26:1H:1299:G:H3'	26:1H:1639:U:O4	2.20	0.42
26:1H:270(E):G:C5	26:1H:270(F):U:C4	3.08	0.42
26:1H:852:G:O2'	26:1H:853:G:H5'	2.19	0.42
11:2A:98:LEU:O	11:2A:101:SER:OG	2.24	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:32:25:ARG:HG2	4:32:30:LYS:O	2.20	0.42
37:35:65:ARG:HB2	62:M5:204:HOH:O	2.20	0.42
4:3E:62:GLN:O	4:3E:66:ARG:HB2	2.20	0.42
38:45:103:MET:HE1	38:45:125:LEU:HD13	2.02	0.42
32:49:13:GLU:H	32:49:13:GLU:HG3	1.76	0.42
32:49:5:VAL:O	32:49:5:VAL:HG12	2.20	0.42
5:4E:147:ASP:OD1	5:4E:147:ASP:N	2.43	0.42
35:58:57:ALA:O	35:58:59:LYS:N	2.52	0.42
33:59:37:VAL:HG22	33:59:38:SER:H	1.84	0.42
6:5E:76:ALA:HA	6:5E:79:LEU:HD12	2.00	0.42
14:5I:13:THR:N	14:5I:14:PRO:HD2	2.35	0.42
7:6E:124:LEU:HD23	7:6E:124:LEU:HA	1.72	0.42
41:75:8:LYS:O	41:75:11:GLU:HB3	2.20	0.42
1:1G:943:U:H1'	9:82:124:GLN:OE1	2.20	0.42
19:AI:31:ILE:HG23	19:AI:49:ILE:HG12	2.02	0.42
41:B8:27:THR:HG23	41:B8:90:GLN:HB3	2.02	0.42
46:G8:94:LYS:NZ	46:G8:95:LYS:H	2.18	0.42
53:N8:49:CYS:O	53:N8:49:CYS:SG	2.78	0.42
29:11:75:ILE:HD13	29:11:99:ASP:OD2	2.20	0.42
1:13:1023:G:C3'	1:13:1024:G:H5"	2.49	0.42
1:13:1106:G:C6	1:13:1107:C:C4	3.07	0.42
1:13:779:C:H2'	1:13:780:A:O4'	2.20	0.42
1:13:811:C:H4'	1:13:900:A:N6	2.35	0.42
26:14:1625:C:H2'	26:14:1626:G:O4'	2.20	0.42
26:14:2196:C:O2'	26:14:2197:U:H5'	2.19	0.42
26:14:690:G:H2'	26:14:691:C:C6	2.54	0.42
26:14:819:A:C2'	26:14:820:A:H5'	2.50	0.42
27:16:15:A:H1'	27:16:109:G:C4	2.55	0.42
26:14:1820:U:O2	29:19:201:HIS:HB3	2.20	0.42
29:19:206:LEU:HD23	29:19:206:LEU:HA	1.80	0.42
2:1E:192:SER:OG	2:1E:193:ASP:N	2.51	0.42
2:1E:223:ILE:HG12	2:1E:223:ILE:H	1.52	0.42
1:1G:677:U:H3	1:1G:713:G:H22	1.68	0.42
1:1G:994:A:N7	1:1G:1216:G:H4'	2.35	0.42
26:1H:1168:G:C2	26:1H:1182:A:C2	3.08	0.42
26:1H:1569:A:O5'	29:11:61:LEU:HD21	2.19	0.42
26:1H:2159:G:C4	26:1H:2160:G:C8	3.08	0.42
26:1H:2262:U:C2'	26:1H:2263:C:H5'	2.49	0.42
26:1H:2517:C:C2	26:1H:2542:A:N6	2.87	0.42
26:1H:2552:U:O5'	26:1H:2552:U:H6	2.03	0.42
26:1H:2881:C:C2	26:1H:2882:A:C8	3.07	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:589:C:H2'	26:1H:590:A:H8	1.85	0.42
26:1H:58:G:N2	26:1H:70:G:C4	2.88	0.42
26:1H:825:C:H5"	62:1H:4527:HOH:O	2.19	0.42
27:1J:23:G:C2	27:1J:24:G:O6	2.73	0.42
30:21:182:LEU:HD12	30:21:183:LEU:N	2.34	0.42
31:31:165:ARG:HA	31:31:168:ARG:HD3	2.02	0.42
4:32:162:LEU:HD22	4:32:178:VAL:HG13	2.02	0.42
37:35:81:GLN:OE1	37:35:107:LYS:HB2	2.20	0.42
4:3E:74:GLN:O	4:3E:78:LEU:HG	2.20	0.42
12:3I:111:LYS:HD3	12:3I:112:ASP:N	2.35	0.42
1:13:363:A:N7	12:3I:33:ARG:CZ	2.83	0.42
25:4L:21:A:H61	25:4L:22:A:N6	2.18	0.42
33:59:6:ARG:HG2	33:59:7:LEU:HG	2.02	0.42
14:5A:3:ARG:HA	14:5A:4:LYS:HA	1.76	0.42
26:1H:2562:U:H1'	36:68:23:ARG:HH11	1.85	0.42
34:69:144:VAL:O	34:69:144:VAL:HG23	2.19	0.42
15:6A:75:PRO:HB2	15:6A:79:ARG:NH2	2.35	0.42
37:78:50:ARG:NH2	37:78:50:ARG:HG3	2.24	0.42
8:7E:39:LEU:HB3	8:7E:45:ILE:HG12	2.00	0.42
8:7E:60:ARG:HD3	8:7E:62:TYR:OH	2.20	0.42
9:8E:26:VAL:HA	9:8E:61:ALA:HB3	2.01	0.42
26:14:456:C:C2	45:B5:69:TYR:CE2	3.08	0.42
41:B8:105:LEU:O	41:B8:107:ASP:N	2.52	0.42
41:B8:1:MET:H2	41:B8:2:ASN:HB3	1.85	0.42
46:C5:85:VAL:HG23	46:C5:96:ILE:O	2.19	0.42
42:C8:110:VAL:O	42:C8:114:LYS:HG3	2.20	0.42
47:D5:146:ILE:HD12	47:D5:146:ILE:HA	1.88	0.42
50:G5:5:GLU:O	50:G5:8:LYS:N	2.51	0.42
46:G8:9:LYS:HA	46:G8:27:VAL:CG2	2.47	0.42
29:11:108:PRO:HG3	29:11:143:HIS:CE1	2.54	0.42
29:11:176:ARG:HH11	29:11:176:ARG:HG2	1.85	0.42
1:13:1036:G:N7	1:13:1037:C:C4	2.88	0.42
1:13:406:G:H2'	1:13:407:G:H8	1.83	0.42
1:13:417:C:H2'	1:13:418:C:C6	2.52	0.42
1:13:454:C:H3'	1:13:455:C:C6	2.55	0.42
1:13:451:A:N6	1:13:480:U:H2'	2.35	0.42
1:13:954:G:C2	1:13:955:U:C2	3.08	0.42
26:14:1384:A:N3	26:14:1405:U:H1'	2.35	0.42
26:14:1709:U:H2'	26:14:1710:C:C6	2.55	0.42
26:14:1899:G:N2	26:14:1902:C:N4	2.56	0.42
26:14:2119:A:C6	26:14:2171:A:H2	2.38	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2250:G:OP2	26:14:2275:C:H2'	2.19	0.42
26:14:2467:C:H4'	38:45:123:HIS:CG	2.55	0.42
26:14:2697:G:C6	26:14:2698:U:C4	3.08	0.42
26:14:304:G:H2'	26:14:305:U:H6	1.85	0.42
29:19:71:ASP:CG	29:19:103:ARG:HH22	2.23	0.42
1:1G:559:A:H4'	1:1G:560:U:H5"	2.02	0.42
1:1G:673:G:H5"	6:52:87:ARG:NH1	2.35	0.42
26:1H:51:G:N3	26:1H:119:A:C2	2.88	0.42
26:1H:1836:C:H2'	26:1H:1837:C:H6	1.84	0.42
26:1H:185:U:H2'	26:1H:186:G:H8	1.85	0.42
26:1H:1917:U:H2'	26:1H:1918:A:O4'	2.19	0.42
26:1H:1834:U:H4'	26:1H:1969:A:C6	2.55	0.42
26:1H:2119:A:C2	26:1H:2171:A:H2	2.38	0.42
26:1H:2248:C:H2'	26:1H:2249:U:O4'	2.20	0.42
26:1H:2864:G:OP1	41:B8:119:LYS:HD2	2.20	0.42
26:1H:282:A:C4	26:1H:359:A:C2	3.08	0.42
24:1L:37:A:N6	24:1L:38:A:C2	2.88	0.42
30:21:54:GLN:HB3	30:21:76:ARG:HH21	1.84	0.42
3:22:6:HIS:CE1	3:22:184:TYR:CE2	3.08	0.42
30:29:68:ALA:C	30:29:70:ALA:N	2.71	0.42
3:2E:60:ALA:N	3:2E:63:ASN:OD1	2.53	0.42
11:2I:48:ILE:HA	11:2I:48:ILE:HD12	1.73	0.42
23:2K:17:C:H5'	23:2K:62:C:OP1	2.20	0.42
31:31:178:PRO:HG2	31:31:179:GLU:CD	2.40	0.42
31:39:116:ASP:O	31:39:120:GLU:HG2	2.20	0.42
31:39:3:GLU:CA	31:39:24:LEU:HD12	2.50	0.42
31:39:51:THR:HB	31:39:88:VAL:HG11	2.02	0.42
4:3E:162:LEU:HA	4:3E:162:LEU:HD23	1.80	0.42
4:3E:18:LYS:HD3	4:3E:31:CYS:SG	2.60	0.42
4:3E:82:ALA:O	4:3E:85:LYS:HE3	2.20	0.42
24:3K:31:A:OP2	24:3K:31:A:H8	2.02	0.42
24:3L:48:C:H6	24:3L:59:A:H1'	1.84	0.42
13:4A:35:GLU:O	13:4A:38:GLY:N	2.52	0.42
13:4A:56:LEU:O	13:4A:60:VAL:HG23	2.20	0.42
13:4A:88:ARG:HG3	13:4A:88:ARG:H	1.41	0.42
33:51:71:LEU:HD12	33:51:71:LEU:HA	1.77	0.42
35:58:34:LEU:HD21	35:58:120:LEU:HB2	2.02	0.42
6:5E:44:GLY:HA2	6:5E:59:TYR:CE2	2.55	0.42
7:62:36:LYS:O	7:62:39:ALA:N	2.53	0.42
15:6I:66:LEU:HA	15:6I:66:LEU:HD12	1.78	0.42
8:72:120:THR:OG1	8:72:121:ASP:N	2.53	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
42:85:65:ILE:HD11	42:85:96:ALA:HB3	2.02	0.42
38:88:17:LEU:HA	38:88:17:LEU:HD23	1.66	0.42
9:8E:125:TYR:HD1	9:8E:126:SER:H	1.67	0.42
42:85:92:ARG:HD2	43:95:11:GLN:HB2	2.01	0.42
39:98:44:LEU:HA	39:98:44:LEU:HD23	1.87	0.42
19:AI:39:THR:HG22	19:AI:40:ILE:H	1.84	0.42
45:B5:60:ARG:HB2	45:B5:60:ARG:HE	1.56	0.42
20:BA:73:HIS:C	20:BA:74:LYS:HG2	2.40	0.42
46:C5:87:LYS:H	46:C5:94:LYS:HG2	1.85	0.42
47:D5:52:SER:C	47:D5:54:HIS:H	2.23	0.42
49:F5:15:ALA:O	49:F5:40:ARG:HG3	2.20	0.42
27:16:77:U:OP1	47:H8:19:ARG:NH2	2.53	0.42
48:I8:60:PHE:CD1	48:I8:60:PHE:N	2.88	0.42
1:13:1036:G:H5'	1:13:1037:C:OP2	2.20	0.41
1:13:1176:A:H2'	1:13:1177:G:O4'	2.20	0.41
1:13:1277:C:O2'	1:13:1279:A:H8	2.03	0.41
1:13:1323:G:H2'	1:13:1324:A:C8	2.54	0.41
1:13:1338:G:C6	1:13:1339:A:C6	3.08	0.41
1:13:1508:G:H2'	1:13:1509:C:O4'	2.19	0.41
1:13:339:C:H2'	1:13:340:U:H6	1.85	0.41
1:13:502:G:OP1	12:3I:118:SER:HB2	2.19	0.41
1:13:917:G:H2'	1:13:918:A:C8	2.55	0.41
26:14:1180:C:H2'	26:14:1181:C:C6	2.54	0.41
26:14:973:A:O4'	26:14:1188:U:C6	2.73	0.41
26:14:1448:G:H2'	26:14:1449:A:C8	2.55	0.41
26:14:1582:C:O2'	26:14:1586:A:H8	2.01	0.41
26:14:1615:C:C5	26:14:1617:C:C4	3.08	0.41
26:14:1689:A:N6	26:14:1698:A:H2	2.03	0.41
26:14:1716:U:H2'	26:14:1717:G:C8	2.54	0.41
26:14:1742:C:H5'	26:14:1743:G:OP2	2.19	0.41
26:14:296:C:OP2	46:C5:4:LYS:NZ	2.44	0.41
26:14:483:A:H1'	46:C5:60:PHE:CE1	2.54	0.41
26:14:620:G:H8	26:14:622:G:O6	2.03	0.41
26:14:805:G:C8	37:35:38:GLN:NE2	2.88	0.41
26:14:811:U:H2'	37:35:21:ARG:HA	2.01	0.41
26:14:825:C:H2'	26:14:826:U:O4'	2.20	0.41
10:1A:44:VAL:HG21	10:1A:66:ARG:HH21	1.85	0.41
1:1G:1063:C:H3'	1:1G:1064:G:H2'	2.02	0.41
1:1G:1258:G:H2'	1:1G:1259:C:H6	1.85	0.41
1:1G:4:U:O4	8:72:105:ARG:HD3	2.20	0.41
1:1G:743:U:H2'	1:1G:744:C:C6	2.55	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:791:G:C6	1:1G:792:A:N7	2.87	0.41
1:1G:834:C:H2'	1:1G:835:U:H6	1.84	0.41
1:1G:97:U:H2'	1:1G:99:C:C6	2.55	0.41
26:1H:1110:G:O2'	26:1H:1111:A:O5'	2.34	0.41
26:1H:1118:C:H2'	26:1H:1119:C:C6	2.55	0.41
26:1H:1280:G:N2	26:1H:1291:C:C2	2.88	0.41
26:1H:1628:G:H2'	26:1H:1629:U:C6	2.55	0.41
26:1H:1692:U:H2'	26:1H:1694:C:C5	2.54	0.41
26:1H:2422:A:N7	56:Q8:31:HIS:HE1	2.16	0.41
26:1H:249:C:H4'	26:1H:250:G:O5'	2.20	0.41
26:1H:2649:U:H2'	26:1H:2650:U:C6	2.54	0.41
26:1H:448:U:C4	26:1H:583:G:H1'	2.55	0.41
26:1H:909:A:H2'	26:1H:912:C:C5	2.55	0.41
4:32:13:ARG:C	4:32:15:GLU:N	2.73	0.41
4:32:86:LYS:HD2	4:32:86:LYS:HA	1.75	0.41
37:35:122:PRO:O	37:35:123:LEU:HD23	2.20	0.41
12:3A:69:TYR:CE2	12:3A:71:PRO:HA	2.55	0.41
12:3A:89:ARG:HE	12:3A:89:ARG:HB3	1.51	0.41
4:3E:107:ARG:HH22	4:3E:194:LEU:CD2	2.19	0.41
32:41:115:ARG:HB3	32:41:115:ARG:NH1	2.35	0.41
32:41:125:PHE:CD1	32:41:131:TYR:HB2	2.55	0.41
32:41:12:TYR:HA	32:41:16:ARG:HG3	2.02	0.41
5:4E:100:VAL:HA	5:4E:118:ILE:HG22	2.02	0.41
33:51:86:GLU:CD	33:51:165:ALA:HB3	2.40	0.41
33:59:99:VAL:HG13	33:59:100:GLY:H	1.84	0.41
33:59:130:ARG:O	33:59:131:VAL:HB	2.20	0.41
9:8E:46:ALA:O	9:8E:78:LYS:HA	2.20	0.41
43:95:97:LYS:HD3	43:95:97:LYS:HA	1.81	0.41
49:F5:49:VAL:HG21	49:F5:67:ILE:HD12	2.02	0.41
47:H8:154:ASP:OD1	47:H8:154:ASP:N	2.43	0.41
48:I8:10:THR:C	48:I8:12:ASN:H	2.24	0.41
55:P8:37:LYS:HG2	55:P8:37:LYS:O	2.20	0.41
2:12:149:LEU:HA	2:12:149:LEU:HD23	1.78	0.41
1:13:1151:A:C2	1:13:1152:A:C5	3.08	0.41
1:13:1206:G:C6	1:13:1207:G:C5	3.08	0.41
1:13:1440:C:H2'	1:13:1441:G:O4'	2.20	0.41
1:13:492:G:C6	1:13:493:G:C4	3.07	0.41
1:13:598:U:H4'	8:7E:94:TYR:CG	2.55	0.41
1:13:621:A:H2'	1:13:622:A:O4'	2.20	0.41
1:13:693:G:H2'	1:13:694:A:C8	2.55	0.41
1:13:953:G:O5'	1:13:953:G:H8	2.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1288:U:H4'	26:14:1289:C:OP2	2.20	0.41
26:14:1996:C:P	36:25:31:LYS:HE2	2.60	0.41
26:14:2065:C:H2'	26:14:2066:C:C6	2.55	0.41
26:14:2070:G:H2'	26:14:2071:A:C8	2.55	0.41
26:14:2130:U:O2'	26:14:2134:A:H1'	2.20	0.41
26:14:254:G:O6	56:M5:5:LYS:HG2	2.20	0.41
35:15:12:ARG:HG2	35:15:14:VAL:HG22	2.02	0.41
1:1G:1262:C:N4	1:1G:1273:G:H1	2.13	0.41
1:1G:183:G:O5'	1:1G:183:G:H8	2.03	0.41
1:1G:641:U:O3'	1:1G:642:A:H8	2.03	0.41
1:1G:973:G:H4'	10:1A:55:LYS:HD3	2.02	0.41
26:1H:1222:C:H2'	26:1H:1223:C:C6	2.55	0.41
26:1H:1321:A:H2'	26:1H:1322:A:O4'	2.21	0.41
26:1H:1424:G:H2'	26:1H:1425:G:O4'	2.21	0.41
26:1H:1591:G:H2'	26:1H:1592:C:C6	2.55	0.41
26:1H:2188:C:N4	26:1H:2189:U:O2	2.53	0.41
26:1H:2360:A:H2'	26:1H:2361:A:O4'	2.20	0.41
26:1H:30:G:OP2	42:C8:5:LYS:NZ	2.52	0.41
26:1H:771:G:OP1	55:P8:10:ARG:NH1	2.54	0.41
27:1J:16:G:H2'	27:1J:17:C:C6	2.54	0.41
30:21:14:ILE:O	30:21:15:PHE:HB2	2.20	0.41
23:2L:30:G:N2	23:2L:42:C:O2	2.49	0.41
31:31:39:TRP:O	31:31:43:LYS:HG2	2.20	0.41
4:32:20:TYR:HD1	4:32:26:CYS:HB2	1.84	0.41
31:39:36:VAL:HG11	31:39:183:VAL:HG21	2.01	0.41
4:3E:111:ALA:HB2	4:3E:120:LEU:HD12	2.01	0.41
38:45:22:LYS:N	38:45:23:GLY:HA3	2.35	0.41
26:1H:2748:A:P	33:51:70:THR:HG21	2.60	0.41
33:59:10:PRO:O	33:59:12:PRO:HD3	2.20	0.41
33:59:97:ARG:O	33:59:99:VAL:HG12	2.20	0.41
6:5E:63:TYR:HB3	6:5E:65:VAL:HG12	2.01	0.41
14:5I:23:ARG:NH1	14:5I:30:ALA:HB2	2.31	0.41
40:65:86:ALA:O	40:65:87:PHE:HB2	2.20	0.41
26:1H:2563:U:O2'	36:68:28:SER:HB3	2.20	0.41
26:1H:2132:U:N3	28:71:5:LYS:HD2	2.35	0.41
39:98:103:ARG:NH2	39:98:110:PRO:HD3	2.34	0.41
39:98:2:ARG:NH1	39:98:2:ARG:HB3	2.35	0.41
20:BI:35:THR:HG23	62:BI:202:HOH:O	2.20	0.41
48:E5:17:GLN:O	48:E5:19:LYS:NZ	2.46	0.41
44:E8:86:LEU:HD12	44:E8:87:PRO:CD	2.46	0.41
47:H8:17:ALA:O	47:H8:21:ALA:N	2.50	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
47:H8:28:MET:HE2	47:H8:37:VAL:CG1	2.50	0.41
2:12:164:VAL:HB	2:12:186:ALA:HB2	2.02	0.41
1:13:1021:G:H2'	1:13:1022:G:O4'	2.21	0.41
1:13:1363:A:H1'	1:13:1365:G:N7	2.35	0.41
1:13:567:G:H2'	1:13:568:G:O4'	2.20	0.41
1:13:749:C:H2'	1:13:750:G:H8	1.85	0.41
1:13:843:U:H5"	1:13:848:C:C5	2.55	0.41
1:13:912:C:O2'	1:13:913:A:H5'	2.20	0.41
26:14:1171:G:O2'	26:14:1173:G:OP2	2.37	0.41
26:14:1174:A:H2'	26:14:1176:G:OP1	2.19	0.41
26:14:1429:G:H2'	26:14:1430:C:C6	2.55	0.41
26:14:2001:A:H2'	26:14:2002:G:C8	2.55	0.41
26:14:2008:C:H2'	26:14:2009:G:H8	1.84	0.41
26:14:2228:G:C5	26:14:2229:C:C4	3.08	0.41
26:14:2070:G:C2	26:14:2442:C:C2	3.08	0.41
26:14:288:C:H2'	26:14:289:A:C8	2.55	0.41
26:14:870:A:H2'	26:14:871:U:O4'	2.19	0.41
26:14:986:C:C2'	26:14:987:G:H5'	2.50	0.41
26:14:6:A:N9	35:15:129:PRO:HB2	2.34	0.41
27:16:42:C:H4'	32:41:67:LYS:HD2	2.02	0.41
27:16:50:G:OP1	40:A8:63:THR:OG1	2.29	0.41
29:19:218:ARG:HB3	29:19:219:PRO:HD2	2.02	0.41
29:19:44:ASN:HB3	29:19:45:ASN:CA	2.34	0.41
29:19:36:PRO:O	29:19:61:LEU:HD12	2.20	0.41
10:1A:40:LEU:HD13	10:1A:71:LEU:HD22	2.00	0.41
1:1G:1368:G:H4'	10:1A:46:ARG:HH22	1.85	0.41
10:1A:75:ILE:HG13	10:1A:76:ASN:H	1.82	0.41
2:1E:212:GLN:OE1	2:1E:216:SER:OG	2.35	0.41
1:1G:1073:U:H2'	1:1G:1074:G:H8	1.84	0.41
1:1G:1357:A:C5	1:1G:1358:U:N3	2.83	0.41
1:1G:407:G:OP1	4:32:115:ARG:NH2	2.51	0.41
1:1G:409:G:H2'	1:1G:410:G:O4'	2.20	0.41
26:1H:1086:A:H1'	26:1H:1103:A:N1	2.35	0.41
26:1H:1198:U:H2'	26:1H:1199:U:C6	2.55	0.41
26:1H:1533:C:C6	26:1H:1534:G:H5"	2.54	0.41
26:1H:1539:G:C2	26:1H:1540:G:C5	3.08	0.41
26:1H:1591:G:H2'	26:1H:1592:C:H6	1.86	0.41
26:1H:2712:U:H1'	26:1H:2712(A):A:C8	2.55	0.41
26:1H:445:C:OP1	42:C8:2:PRO:HA	2.20	0.41
26:1H:685:A:H1'	26:1H:688:U:O4	2.20	0.41
26:1H:881:G:H3'	26:1H:881:G:N3	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:21:26:ILE:O	30:21:26:ILE:HG12	2.20	0.41
30:21:39:PRO:HD3	30:21:45:THR:HG22	2.03	0.41
30:21:93:VAL:HG21	30:21:180:ASN:HA	2.01	0.41
1:1G:1191:A:H5"	3:22:4:LYS:HZ2	1.84	0.41
36:25:7:TYR:CE1	36:25:20:MET:HB2	2.56	0.41
30:29:37:ARG:NH1	30:29:80:GLU:OE2	2.50	0.41
23:2L:33:OMC:O2'	23:2L:34:U:H6	2.03	0.41
31:39:11:VAL:CG2	31:39:12:LEU:N	2.83	0.41
4:3E:4:TYR:CG	4:3E:5:ILE:N	2.88	0.41
12:3I:64:TYR:O	12:3I:65:GLU:HB3	2.20	0.41
32:41:44:GLY:O	32:41:47:LYS:HB2	2.20	0.41
5:42:13:ILE:HG13	5:42:13:ILE:O	2.20	0.41
38:45:10:ARG:CZ	38:45:10:ARG:HA	2.50	0.41
32:49:161:THR:CG2	32:49:163:ALA:H	2.30	0.41
35:58:90:MET:O	35:58:94:HIS:N	2.53	0.41
33:59:129:THR:C	33:59:130:ARG:HG3	2.39	0.41
6:5E:14:LEU:HD22	6:5E:18:GLN:HB3	2.01	0.41
14:5I:26:ARG:HB3	14:5I:43:CYS:SG	2.60	0.41
26:1H:2674:G:H5'	36:68:26:LYS:HE2	2.02	0.41
36:68:2:ILE:N	36:68:2:ILE:HD13	2.35	0.41
36:68:88:ASN:OD1	36:68:90:GLN:HB2	2.21	0.41
34:69:97:ILE:O	34:69:100:ALA:HB3	2.20	0.41
7:6E:90:GLU:H	7:6E:90:GLU:HG2	1.64	0.41
15:6I:36:ILE:HG12	15:6I:59:MET:HE3	2.02	0.41
41:75:55:ASN:ND2	41:75:55:ASN:O	2.51	0.41
62:1H:4624:HOH:O	37:78:26:GLY:HA3	2.20	0.41
8:7E:75:ARG:HE	8:7E:75:ARG:HB2	1.73	0.41
39:98:44:LEU:O	39:98:45:ARG:C	2.58	0.41
27:16:48:A:P	40:A8:30:ARG:HH22	2.43	0.41
41:B8:110:ILE:HG13	41:B8:111:ARG:N	2.35	0.41
43:D8:38:LEU:HD12	43:D8:40:LEU:H	1.86	0.41
49:F5:91:LYS:HB2	49:F5:91:LYS:HE2	1.67	0.41
49:F5:95:LEU:HA	49:F5:95:LEU:HD13	1.76	0.41
47:H8:124:ILE:HD12	47:H8:125:LEU:H	1.84	0.41
47:H8:132:ASN:HD22	47:H8:160:GLY:HA3	1.85	0.41
47:H8:53:ILE:HG13	47:H8:53:ILE:O	2.21	0.41
56:M5:40:GLU:HG3	56:M5:43:GLN:HB2	2.03	0.41
1:13:1180:A:OP1	9:8E:103:THR:OG1	2.34	0.41
1:13:1223:C:P	19:AI:78:ARG:NH1	2.92	0.41
1:13:407:G:O2'	4:3E:116:GLN:HG3	2.21	0.41
1:13:738:C:H2'	1:13:739:C:C6	2.55	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:107:C:H2'	26:14:108:U:C6	2.55	0.41
26:14:1204:A:N1	26:14:1241:A:H2	2.18	0.41
26:14:1849:G:H2'	26:14:1850:G:H8	1.86	0.41
26:14:2238:G:H4'	26:14:2239:G:N7	2.35	0.41
26:14:2273:A:O2'	26:14:2274:A:H5'	2.20	0.41
26:14:2298:A:N6	26:14:2318:G:C8	2.86	0.41
26:14:2570:G:H2'	26:14:2571:C:O4'	2.21	0.41
26:14:2775:A:N7	62:14:3613:HOH:O	2.52	0.41
26:14:362:U:H5'	26:14:363:G:OP2	2.21	0.41
26:14:577:G:H8	26:14:577:G:O5'	2.04	0.41
26:14:753:C:H2'	26:14:754:C:H6	1.84	0.41
35:15:4:TYR:O	42:85:64:ARG:NH1	2.53	0.41
29:19:130:ALA:HA	29:19:192:THR:HA	2.01	0.41
1:1G:1207:G:H2'	1:1G:1208:C:C6	2.55	0.41
1:1G:598:U:H4'	8:72:94:TYR:CG	2.55	0.41
1:1G:947:G:H2'	1:1G:948:C:H6	1.85	0.41
26:1H:1182:A:H2'	26:1H:1183:G:O4'	2.21	0.41
26:1H:1191:G:P	62:1H:3785:HOH:O	2.79	0.41
26:1H:1401:G:H2'	26:1H:1402:C:C6	2.56	0.41
26:1H:2692:C:OP1	26:1H:2871:C:H5'	2.20	0.41
26:1H:270(E):G:C6	26:1H:270(F):U:C4	3.09	0.41
26:1H:2849:U:H1'	26:1H:2866:U:O2	2.20	0.41
26:1H:2883:A:H5'	26:1H:2884:U:H5'	2.02	0.41
26:1H:320:A:H2'	31:31:136:THR:CG2	2.49	0.41
26:1H:259:G:N2	26:1H:621:A:C8	2.86	0.41
26:1H:657:U:H2'	26:1H:658:C:C6	2.56	0.41
26:1H:725:G:C6	26:1H:726:G:N1	2.88	0.41
27:1J:10:C:C4	27:1J:11:C:C5	3.08	0.41
27:1J:88:C:H4'	27:1J:89:G:OP2	2.21	0.41
30:21:201:THR:HB	30:21:203:LYS:HA	2.02	0.41
36:25:9:GLU:O	36:25:83:ALA:HA	2.20	0.41
30:29:101:ARG:NH2	30:29:171:GLU:HB2	2.34	0.41
11:2I:79:SER:HB2	11:2I:106:LYS:CD	2.50	0.41
23:2L:36:A:H2'	23:2L:37:U:C6	2.55	0.41
31:31:196:LEU:C	31:31:197:ASP:O	2.58	0.41
26:14:589:C:H5"	31:39:95:ARG:HH12	1.85	0.41
24:3K:53:G:H1	24:3K:61:C:N4	2.18	0.41
26:14:960:A:H61	38:45:83:MET:HE2	1.86	0.41
39:55:35:THR:CG2	39:55:100:LEU:HD11	2.51	0.41
35:58:30:ILE:HG22	35:58:34:LEU:HD22	2.02	0.41
34:61:77:LEU:H	34:61:77:LEU:HD12	1.84	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:62:141:VAL:O	7:62:143:ARG:HG3	2.20	0.41
36:68:106:LEU:HD23	36:68:106:LEU:HA	1.84	0.41
15:6I:3:ILE:HA	15:6I:7:GLU:OE1	2.19	0.41
28:71:39:GLU:O	28:71:178:ALA:HB2	2.19	0.41
37:78:114:ILE:HD13	37:78:114:ILE:HG21	1.87	0.41
17:8I:25:ARG:O	17:8I:25:ARG:HG2	2.19	0.41
19:AI:18:LYS:O	19:AI:22:LEU:HD13	2.19	0.41
41:B8:12:SER:CB	41:B8:14:TYR:H	2.33	0.41
43:D8:22:VAL:HG12	43:D8:23:GLU:O	2.20	0.41
43:D8:76:LYS:O	43:D8:79:VAL:HG12	2.20	0.41
51:H5:18:ASP:OD1	51:H5:18:ASP:N	2.53	0.41
47:H8:47:VAL:O	47:H8:50:GLN:HB2	2.21	0.41
48:I8:19:LYS:HD3	48:I8:19:LYS:HA	1.67	0.41
50:K8:4:SER:H	50:K8:7:ARG:N	2.10	0.41
50:K8:64:LEU:HD21	50:K8:68:ARG:NH1	2.35	0.41
26:1H:2347:C:P	54:O8:39:TYR:HH	2.36	0.41
2:12:16:HIS:ND1	2:12:213:LEU:HD22	2.35	0.41
1:13:109:A:N7	1:13:326:G:H2'	2.35	0.41
1:13:192:U:O4'	20:BI:103:GLY:HA2	2.21	0.41
1:13:254:G:O3'	17:8I:69:LYS:NZ	2.43	0.41
1:13:498:A:H4'	1:13:500:G:OP1	2.21	0.41
1:13:66:G:O4'	1:13:173:U:C4	2.74	0.41
1:13:77:C:H2'	1:13:78:G:C8	2.54	0.41
26:14:1392:A:N6	26:14:1393:A:N6	2.68	0.41
26:14:1505:C:H2'	26:14:1506:C:H6	1.85	0.41
26:14:1568:G:OP2	29:19:63:ARG:NH2	2.53	0.41
26:14:1751:C:H2'	26:14:1752:C:C6	2.56	0.41
26:14:1963:U:H4'	26:14:1964:G:OP2	2.20	0.41
26:14:571:A:H5'	26:14:2030:A:N7	2.35	0.41
26:14:212:G:H2'	26:14:213:A:O4'	2.21	0.41
26:14:2569:G:C2	26:14:2570:G:C8	3.09	0.41
26:14:2607:G:H2'	26:14:2608:G:O4'	2.20	0.41
26:14:2749:A:O5'	26:14:2749:A:H8	2.04	0.41
26:14:696:G:H2'	26:14:697:C:H6	1.84	0.41
26:14:864:G:C2'	26:14:865:C:H5'	2.50	0.41
27:16:40:U:C1'	27:16:45:A:H61	2.33	0.41
2:1E:131:PRO:O	2:1E:135:GLN:HG3	2.21	0.41
2:1E:178:ARG:HD3	2:1E:178:ARG:HA	1.82	0.41
2:1E:213:LEU:HG	2:1E:213:LEU:H	1.58	0.41
1:1G:1077:G:N2	1:1G:1080:A:OP2	2.53	0.41
1:1G:1123:A:H4'	10:1A:37:PRO:HD2	2.01	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:1216:G:H2'	1:1G:1217:C:H6	1.85	0.41
1:1G:1277:C:H1'	1:1G:1282:C:O2	2.21	0.41
1:1G:1321:C:N4	1:1G:1322:C:N4	2.66	0.41
1:1G:161:A:C6	1:1G:162:A:C6	3.08	0.41
1:1G:631:G:OP2	1:1G:631:G:H8	2.03	0.41
1:1G:833:U:O2'	1:1G:834:C:H5'	2.21	0.41
26:1H:1222:C:H2'	26:1H:1223:C:H6	1.86	0.41
26:1H:1380:G:N2	26:1H:1570:A:C2	2.88	0.41
26:1H:1475:G:H2'	26:1H:1476:C:C6	2.56	0.41
26:1H:311:A:C6	26:1H:328:U:C4	3.09	0.41
26:1H:639:U:O2'	26:1H:640:C:H5'	2.21	0.41
26:1H:729:G:C4	26:1H:1775:U:O2	2.74	0.41
26:1H:910:A:H2'	26:1H:911:A:C8	2.55	0.41
26:1H:991:C:C4	26:1H:1185:C:N4	2.89	0.41
30:21:47:VAL:HG11	30:21:86:PRO:HD2	2.01	0.41
26:14:2572:A:N7	30:29:144:ARG:HD2	2.36	0.41
3:2E:27:LYS:HA	3:2E:27:LYS:HD2	1.77	0.41
31:39:11:VAL:HG22	31:39:13:SER:HB2	2.01	0.41
4:3E:119:GLN:HG2	4:3E:119:GLN:O	2.21	0.41
4:3E:188:LEU:HA	4:3E:188:LEU:HD22	1.81	0.41
32:41:173:LEU:HD12	32:41:178:PHE:CD2	2.56	0.41
5:42:101:ILE:HG12	5:42:101:ILE:H	1.74	0.41
5:42:90:VAL:HG23	5:42:121:LYS:HB3	2.02	0.41
32:49:106:LEU:HG	32:49:111:LEU:CD1	2.50	0.41
32:49:64:THR:HB	32:49:94:LEU:HD21	2.02	0.41
13:4I:12:ASN:HD22	13:4I:13:LYS:H	1.69	0.41
13:4I:13:LYS:HZ3	13:4I:13:LYS:HA	1.85	0.41
33:59:117:PRO:HB3	33:59:123:PHE:HZ	1.84	0.41
26:1H:2128:C:H3'	28:71:36:LYS:NZ	2.35	0.41
41:75:16:ARG:NH2	41:75:19:LEU:HD21	2.35	0.41
41:75:27:THR:OG1	41:75:89:VAL:HG22	2.20	0.41
37:78:116:GLY:N	37:78:134:ALA:HB2	2.35	0.41
37:78:38:GLN:O	37:78:44:GLY:HA2	2.19	0.41
16:7A:43:LYS:HG2	16:7A:48:TRP:CG	2.55	0.41
9:8E:48:GLU:N	9:8E:49:PRO:HD2	2.35	0.41
26:1H:2707:G:O3'	39:98:68:ARG:HG2	2.21	0.41
19:AA:41:VAL:HG23	19:AA:43:GLU:N	2.36	0.41
41:B8:33:LYS:HG2	41:B8:82:LEU:O	2.20	0.41
20:BA:100:ILE:HG23	20:BA:101:GLY:N	2.35	0.41
20:BI:35:THR:O	20:BI:38:LYS:HB2	2.21	0.41
42:C8:92:ARG:HB2	43:D8:11:GLN:NE2	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
43:D8:75:PHE:HD1	43:D8:82:ARG:HG3	1.86	0.41
47:H8:72:ARG:HA	47:H8:72:ARG:HD3	1.90	0.41
48:I8:24:LYS:HG3	48:I8:36:ILE:HD11	2.02	0.41
54:O8:29:ASN:O	54:O8:32:ASN:HB3	2.21	0.41
2:12:187:LEU:HA	2:12:201:ILE:O	2.20	0.41
2:12:219:VAL:HG23	2:12:221:LEU:H	1.86	0.41
1:13:1252:A:H2'	1:13:1253:G:O4'	2.20	0.41
1:13:263:A:OP2	20:BI:79:ARG:NH1	2.54	0.41
1:13:322:C:OP2	1:13:328:C:N4	2.53	0.41
1:13:358:U:OP1	34:69:87:LYS:NZ	2.50	0.41
1:13:26:A:N6	1:13:558:G:O2'	2.46	0.41
1:13:768:A:H2'	1:13:769:G:O4'	2.21	0.41
26:14:1952:A:C6	26:14:1953:A:C6	3.09	0.41
26:14:2077:A:O2'	26:14:2078:C:H5'	2.21	0.41
26:14:2303:G:C2'	26:14:2304:G:H5'	2.50	0.41
26:14:26:G:OP1	44:A5:80:PRO:HB3	2.21	0.41
26:14:329:G:H1	46:C5:19:LYS:NZ	2.18	0.41
27:16:60:C:N3	27:16:61:G:N7	2.68	0.41
1:1G:114:U:O2'	1:1G:115:G:H5'	2.20	0.41
1:1G:1386:G:C2	1:1G:1387:G:C8	3.09	0.41
1:1G:517:G:N2	1:1G:530:G:OP1	2.44	0.41
26:1H:1176:G:H5'	26:1H:1177:A:P	2.61	0.41
26:1H:2154:G:O5'	26:1H:2154:G:H8	2.03	0.41
26:1H:2070:G:C2	26:1H:2442:C:C2	3.08	0.41
26:1H:2666:C:H3'	26:1H:2667:C:H6	1.86	0.41
26:1H:2688:U:H1'	26:1H:2721:A:N6	2.35	0.41
26:1H:2807:G:H3'	26:1H:2808:U:H5"	2.02	0.41
26:1H:330:A:O2'	26:1H:331:A:C8	2.72	0.41
26:1H:70:G:H21	26:1H:71:A:H62	1.59	0.41
26:1H:803:U:C4	26:1H:804:A:N7	2.88	0.41
24:1L:2:G:N3	24:1L:2:G:H2'	2.36	0.41
3:22:156:ARG:NH2	3:22:159:GLY:O	2.40	0.41
11:2I:59:TYR:O	11:2I:63:LEU:HD12	2.21	0.41
23:2K:29:C:H2'	23:2K:30:G:H8	1.86	0.41
37:35:134:ALA:O	37:35:138:LEU:HB2	2.20	0.41
24:3K:28:U:H2'	24:3K:29:U:C6	2.56	0.41
32:41:135:LEU:O	32:41:154:GLY:HA3	2.20	0.41
13:4A:33:ALA:HA	13:4A:59:TYR:CE2	2.55	0.41
33:51:13:LYS:HD3	33:51:13:LYS:HA	1.83	0.41
39:55:87:TYR:HD1	39:55:90:ARG:HE	1.68	0.41
35:58:99:LEU:HD23	35:58:99:LEU:HA	1.77	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
28:71:41:VAL:HA	28:71:216:THR:HG23	2.02	0.41
16:7I:45:THR:HB	16:7I:47:ASP:OD1	2.21	0.41
16:7I:67:THR:H	16:7I:70:ALA:HB3	1.85	0.41
17:8A:45:HIS:CD2	17:8A:47:PRO:HD3	2.55	0.41
43:95:84:LYS:O	43:95:85:LYS:HG3	2.20	0.41
39:98:118:GLU:OE1	39:98:118:GLU:HA	2.21	0.41
20:BA:50:GLU:HA	20:BA:100:ILE:CG2	2.50	0.41
1:13:103:C:OP2	20:BI:14:LYS:HD2	2.20	0.41
46:C5:64:GLU:HG3	46:C5:64:GLU:H	1.70	0.41
44:E8:24:ILE:HG12	44:E8:36:LEU:HD21	2.02	0.41
49:F5:73:LEU:HA	49:F5:73:LEU:HD23	1.87	0.41
47:H8:25:PRO:O	47:H8:85:HIS:HA	2.20	0.41
55:L5:34:ARG:NH1	55:L5:41:ARG:O	2.54	0.41
1:13:1223:C:OP1	19:AI:78:ARG:NH1	2.54	0.41
1:13:1256:A:O2'	1:13:1257:U:P	2.79	0.41
1:13:658:G:H2'	1:13:659:U:C6	2.56	0.41
26:14:117:G:C6	26:14:119:A:C6	3.08	0.41
26:14:1322:A:N1	26:14:1333:C:O2'	2.35	0.41
26:14:1542:G:O5'	26:14:1543:A:H5"	2.20	0.41
26:14:1966:A:H4'	26:14:1967:C:OP1	2.20	0.41
26:14:2535:G:N3	26:14:2536:G:C8	2.89	0.41
26:14:2591:C:H2'	26:14:2592:G:H8	1.85	0.41
26:14:2522:U:H4'	26:14:2648:C:OP1	2.21	0.41
26:14:2698:U:H2'	26:14:2699:C:C6	2.56	0.41
26:14:2852:G:H2'	26:14:2853:C:C6	2.56	0.41
26:14:732:C:C5	61:14:3437:SPE:H102	2.56	0.41
27:16:66:A:C5	27:16:108:C:C5	3.09	0.41
27:16:71:C:C2	27:16:72:G:C8	3.08	0.41
2:1E:149:LEU:HD23	2:1E:149:LEU:HA	1.71	0.41
1:1G:1328:C:O2'	13:4A:29:ARG:NH2	2.47	0.41
1:1G:167:G:O2'	1:1G:168:G:H5'	2.20	0.41
1:1G:271:C:H2'	1:1G:272:C:H6	1.86	0.41
1:1G:345:C:H4'	1:1G:346:G:O5'	2.20	0.41
1:1G:427:U:H4'	1:1G:541:G:H5"	2.02	0.41
1:1G:672:U:H2'	1:1G:673:G:C8	2.55	0.41
1:1G:741:G:H2'	1:1G:742:G:O4'	2.21	0.41
26:1H:1087:G:N7	26:1H:1089:G:H1'	2.35	0.41
26:1H:1091:G:C6	26:1H:1101:U:C2	3.09	0.41
26:1H:1164:G:C6	26:1H:1165:U:C4	3.09	0.41
26:1H:1389:G:C2	26:1H:1390:U:C2	3.09	0.41
26:1H:1677:A:H2'	26:1H:1678:G:C8	2.56	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:1H:1894:C:C2'	26:1H:1895:C:H5'	2.51	0.41
26:1H:1931:U:H5	26:1H:1969:A:N7	2.18	0.41
26:1H:2000:G:OP1	39:98:5:LYS:NZ	2.51	0.41
26:1H:2347:C:H4'	54:O8:39:TYR:HE1	1.86	0.41
26:1H:2359:C:H2'	26:1H:2360:A:O4'	2.21	0.41
26:1H:2395:C:H5"	26:1H:2396:G:OP2	2.20	0.41
26:1H:2688:U:H1'	26:1H:2721:A:H61	1.86	0.41
26:1H:654:A:N3	26:1H:654(A):A:H5"	2.35	0.41
26:1H:934:G:H2'	26:1H:935:C:H6	1.86	0.41
26:1H:956:G:H5"	26:1H:957:A:OP2	2.21	0.41
4:32:132:ARG:HB3	4:32:132:ARG:HE	1.41	0.41
4:32:15:GLU:OE1	4:32:59:ARG:NH2	2.43	0.41
4:32:31:CYS:HA	59:32:302:SF4:S2	2.60	0.41
4:32:39:PRO:O	4:32:44:GLY:HA3	2.21	0.41
26:14:661:C:H4'	37:35:13:ASN:OD1	2.20	0.41
31:39:110:LEU:HD21	31:39:181:LEU:HD22	2.03	0.41
12:3A:10:LEU:HB3	17:8A:32:TYR:CE2	2.56	0.41
1:1G:363:A:C5	12:3A:31:PRO:HD2	2.55	0.41
12:3A:40:VAL:HG11	12:3A:75:HIS:HE1	1.86	0.41
24:3K:76:A:H5"	49:J8:30:VAL:HG11	2.02	0.41
32:49:53:LEU:O	32:49:57:ALA:N	2.53	0.41
32:49:81:LYS:HB3	32:49:82:LEU:H	1.52	0.41
13:4I:117:VAL:O	13:4I:118:ALA:HB2	2.21	0.41
25:4K:23:A:O2'	25:4K:24:A:N7	2.37	0.41
33:51:80:SER:C	33:51:81:GLU:HG3	2.40	0.41
39:55:21:TYR:OH	39:55:43:GLU:HG2	2.21	0.41
6:5E:35:ALA:HA	6:5E:67:MET:HB3	2.02	0.41
6:5E:33:TYR:HD1	6:5E:71:ARG:HB3	1.86	0.41
40:65:80:LEU:HD23	40:65:80:LEU:HA	1.86	0.41
15:6A:87:ILE:HG22	15:6A:88:ARG:H	1.85	0.41
8:7E:77:GLU:HG2	8:7E:78:GLN:N	2.36	0.41
42:85:16:LYS:HE3	42:85:16:LYS:HB3	1.91	0.41
38:88:32:TYR:CE2	38:88:133:ARG:HG3	2.56	0.41
18:9I:22:VAL:HA	18:9I:25:THR:OG1	2.21	0.41
44:A5:18:ARG:HA	44:A5:76:VAL:HG11	2.03	0.41
29:11:108:PRO:HD2	29:11:111:LEU:HG	2.02	0.41
29:11:118:VAL:HG22	29:11:119:ALA:N	2.36	0.41
29:11:127:VAL:HA	29:11:193:VAL:HG22	2.02	0.41
26:1H:1568:G:P	29:11:63:ARG:HH12	2.44	0.41
29:11:80:ALA:HB2	29:11:96:HIS:CD2	2.56	0.41
2:12:168:THR:HG21	2:12:191:ASP:O	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:13:339:C:H2'	1:13:340:U:C6	2.56	0.41
1:13:664:G:N2	1:13:741:G:H1	2.11	0.41
26:14:1681:G:N2	62:14:3653:HOH:O	2.20	0.41
26:14:1829:A:N3	29:19:15:PHE:HE2	2.19	0.41
26:14:2134:A:H2'	26:14:2134:A:N3	2.36	0.41
26:14:807:U:H4'	26:14:2446:G:OP1	2.20	0.41
26:14:2500:U:H5"	26:14:2501:C:OP2	2.21	0.41
26:14:2688:U:H1'	26:14:2721:A:N6	2.35	0.41
26:14:547:A:H2'	26:14:548:A:C8	2.56	0.41
26:14:670:A:H4'	26:14:671:C:O5'	2.20	0.41
26:14:807:U:C2	26:14:808:G:C8	3.08	0.41
29:19:132:PRO:HA	29:19:189:CYS:O	2.20	0.41
1:1G:1152:A:H5'	10:1A:13:HIS:CD2	2.55	0.41
1:1G:1264:C:H1'	1:1G:1272:G:N2	2.36	0.41
1:1G:1300:G:O2'	1:1G:1301:U:P	2.78	0.41
1:1G:1465:C:H2'	1:1G:1466:C:O4'	2.21	0.41
1:1G:339:C:H2'	1:1G:340:U:C6	2.55	0.41
1:1G:909:A:H2'	1:1G:910:C:O4'	2.20	0.41
26:1H:1176:G:H5'	26:1H:1177:A:OP2	2.20	0.41
26:1H:2129:C:P	28:71:6:ARG:HH11	2.44	0.41
26:1H:2131:G:H1'	26:1H:2158:A:C6	2.55	0.41
26:1H:1889:A:N1	26:1H:2234:G:H1'	2.35	0.41
26:1H:2820:A:HO2'	26:1H:2821:A:P	2.43	0.41
26:1H:957:A:N1	26:1H:2458:G:H4'	2.35	0.41
27:1J:15:A:H1'	27:1J:109:G:C4	2.56	0.41
30:21:181:LEU:HD12	30:21:181:LEU:HA	1.85	0.41
30:21:23:VAL:HA	30:21:184:VAL:O	2.20	0.41
30:29:32:PRO:HA	30:29:90:THR:HA	2.03	0.41
37:35:71:VAL:CG1	37:35:72:PRO:HD3	2.46	0.41
31:39:143:ALA:O	31:39:148:LEU:HB2	2.20	0.41
4:3E:207:TYR:O	4:3E:209:ARG:HG3	2.21	0.41
24:3K:1:G:N2	24:3K:2:G:N7	2.69	0.41
13:4A:81:LEU:HD12	13:4A:89:GLY:HA3	2.03	0.41
6:52:86:ARG:O	6:52:87:ARG:HG2	2.20	0.41
35:58:127:ASP:OD1	35:58:127:ASP:N	2.53	0.41
33:59:20:ALA:HB3	33:59:23:ARG:O	2.21	0.41
40:65:3:ARG:HD2	40:65:4:LEU:N	2.36	0.41
7:6E:15:ASP:O	7:6E:19:GLY:HA2	2.20	0.41
37:78:60:MET:HE3	37:78:60:MET:HB2	1.94	0.41
9:82:20:ARG:O	9:82:20:ARG:HG3	2.21	0.41
9:82:48:GLU:N	9:82:49:PRO:HD2	2.35	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:88:17:LEU:HB3	38:88:39:PRO:HB2	2.03	0.41
9:8E:82:ALA:O	9:8E:86:VAL:HG23	2.21	0.41
43:95:12:TYR:CD1	43:95:12:TYR:N	2.88	0.41
19:AI:51:VAL:HB	19:AI:75:ALA:HB2	2.03	0.41
45:B5:66:LEU:HD23	45:B5:66:LEU:HA	1.82	0.41
41:B8:1:MET:HA	41:B8:3:ARG:N	2.35	0.41
20:BI:49:ALA:HA	20:BI:92:LEU:HD11	2.03	0.41
29:11:33:LEU:HD12	29:11:33:LEU:HA	1.73	0.41
1:1G:1104:G:H4'	2:12:111:ARG:NH2	2.35	0.41
2:12:167:PRO:O	2:12:171:ALA:N	2.54	0.41
2:12:218:ALA:O	2:12:219:VAL:HG22	2.21	0.41
1:13:1134:G:N1	1:13:1135:U:H1'	2.35	0.41
1:13:1277:C:O2'	1:13:1279:A:H1'	2.20	0.41
1:13:1286:A:C2	21:1F:18:TYR:OH	2.74	0.41
1:13:1326:C:H2'	1:13:1327:C:H6	1.86	0.41
1:13:200:G:N2	1:13:218:C:C2	2.88	0.41
1:13:199:G:H2'	1:13:200:G:O4'	2.21	0.41
1:13:416:G:C6	1:13:417:C:C4	3.08	0.41
26:14:597:U:H2'	26:14:598:G:H8	1.83	0.41
26:14:863:A:H2'	26:14:864:G:C8	2.55	0.41
26:14:861:A:C2	26:14:917:A:C4	3.09	0.41
26:14:933:A:C5	26:14:934:G:C8	3.09	0.41
26:14:950:G:C2	26:14:968:G:C2	3.09	0.41
29:19:126:GLN:O	29:19:193:VAL:HG22	2.21	0.41
29:19:16:MET:HG3	29:19:206:LEU:O	2.21	0.41
2:1E:130:ARG:HA	2:1E:131:PRO:HD3	1.92	0.41
2:1E:68:ILE:HG13	2:1E:161:ALA:HB3	2.02	0.41
1:1G:1385:G:C6	1:1G:1386:G:N7	2.88	0.41
1:1G:1517:G:C6	1:1G:1518:A:C5	3.08	0.41
1:1G:15:G:H2'	1:1G:16:A:C8	2.55	0.41
1:1G:690:G:H2'	1:1G:691:G:C8	2.56	0.41
26:1H:1047:G:H2'	26:1H:1110:G:N1	2.35	0.41
26:1H:1141:U:C5	35:58:64:GLY:HA3	2.56	0.41
26:1H:1583:A:O5'	26:1H:1585:C:H5	2.04	0.41
26:1H:1657:C:H2'	26:1H:1658:C:C6	2.56	0.41
26:1H:2331:G:C4'	48:I8:42:GLY:HA3	2.51	0.41
26:1H:2346:A:H5'	26:1H:2383:G:O4'	2.21	0.41
26:1H:2507:C:H5'	26:1H:2573:C:N4	2.36	0.41
27:1J:13:A:N1	27:1J:69:G:O2'	2.43	0.41
27:1J:51:G:C6	27:1J:52:A:H2	2.38	0.41
30:21:35:GLN:HB3	30:21:48:GLN:HB2	2.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:22:172:ARG:HE	3:22:172:ARG:HB3	1.68	0.41
36:25:11:ALA:HB1	36:25:99:PHE:O	2.21	0.41
31:31:135:LYS:HB3	31:31:138:GLU:HG3	2.03	0.41
31:39:170:LEU:HA	31:39:171:PRO:HD3	1.90	0.41
31:39:194:MET:HB2	31:39:194:MET:HE3	1.83	0.41
24:3K:72:C:N3	24:3K:73:A:C8	2.89	0.41
24:3L:55:U:H2'	24:3L:57:G:OP1	2.21	0.41
5:42:90:VAL:O	5:42:120:THR:HA	2.20	0.41
32:49:136:ARG:HD3	32:49:137:GLU:HG3	2.03	0.41
32:49:14:GLU:O	32:49:17:PRO:HG2	2.21	0.41
32:49:84:LYS:HE2	32:49:84:LYS:HB3	1.92	0.41
33:51:92:ILE:HD12	33:51:92:ILE:N	2.36	0.41
35:58:39:ARG:NH2	35:58:41:ASP:OD2	2.54	0.41
14:5I:3:ARG:HG3	14:5I:4:LYS:N	2.35	0.41
7:62:108:ALA:O	7:62:111:ARG:HG3	2.21	0.41
34:69:140:LEU:HD12	34:69:140:LEU:HA	1.74	0.41
34:69:41:GLU:HA	34:69:44:LEU:HB2	2.03	0.41
15:6A:48:LYS:HB2	15:6A:48:LYS:HE3	1.87	0.41
1:1G:878:G:H1'	8:72:3:THR:HG21	2.03	0.41
9:8E:70:LYS:O	9:8E:74:ILE:HG13	2.20	0.41
39:98:22:ARG:HG2	39:98:69:ASP:HB3	2.03	0.41
39:98:57:ARG:HB3	39:98:59:ASP:OD1	2.20	0.41
39:98:98:LEU:HA	39:98:98:LEU:HD23	1.86	0.41
26:1H:1152:C:H4'	42:C8:77:SER:HA	2.03	0.41
42:C8:95:LEU:HD22	43:D8:4:ILE:CD1	2.49	0.41
26:1H:302:C:OP1	46:G8:81:LYS:HD3	2.21	0.41
48:I8:66:VAL:O	48:I8:82:ARG:N	2.53	0.41
27:16:12:C:O2'	48:I8:74:ARG:HG2	2.20	0.41
53:J5:12:SER:OG	53:J5:15:ARG:HB2	2.20	0.41
29:11:16:MET:HG3	29:11:207:GLY:HA3	2.03	0.41
1:13:1179:A:H4'	9:8E:103:THR:HA	2.03	0.41
1:13:1157:A:C6	1:13:1180:A:C5	3.09	0.41
1:13:1260:C:H6	1:13:1260:C:H3'	1.86	0.41
1:13:160:A:H2'	1:13:160:A:N3	2.35	0.41
1:13:240:C:H2'	1:13:241:C:H6	1.85	0.41
1:13:564:C:O2'	8:7E:91:ARG:NH2	2.53	0.41
1:13:703:G:O5'	1:13:703:G:C8	2.71	0.41
26:14:1019:U:H2'	26:14:1020:A:H8	1.86	0.41
26:14:1628:G:H2'	26:14:1629:U:H6	1.85	0.41
26:14:2287:A:H61	26:14:2344:U:H3	1.60	0.41
26:14:2432:A:C2	49:F5:35:THR:HG22	2.56	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2579:C:C4'	30:29:134:ILE:HG12	2.50	0.41
26:14:2716:U:O2'	26:14:2717:G:H5'	2.21	0.41
26:14:271:G:H2'	26:14:272:G:C8	2.56	0.41
26:14:488:G:N2	26:14:492:A:OP2	2.54	0.41
2:1E:55:PHE:HD1	2:1E:58:ILE:HD12	1.83	0.41
1:1G:109:A:C6	1:1G:326:G:C6	3.08	0.41
1:1G:1285:A:OP1	1:1G:1285:A:H8	2.03	0.41
1:1G:1343:G:O2'	1:1G:1344:C:H5'	2.21	0.41
1:1G:1415:G:C6	1:1G:1486:G:C6	3.08	0.41
1:1G:445:G:H2'	1:1G:446:G:H8	1.86	0.41
1:1G:629:G:H3'	1:1G:630:G:H5"	2.02	0.41
1:1G:693:G:H2'	1:1G:694:A:C8	2.56	0.41
26:1H:1970:A:H4'	26:1H:1971:A:OP1	2.21	0.41
26:1H:2145:C:H5	26:1H:2148:G:N2	2.17	0.41
26:1H:2783:G:H2'	26:1H:2784:C:C6	2.55	0.41
26:1H:2791:C:N4	26:1H:2807:G:H1	2.19	0.41
30:21:174:ASP:OD1	30:21:175:VAL:N	2.54	0.41
26:14:2052:G:O4'	30:29:142:GLY:HA3	2.21	0.41
11:2I:59:TYR:CZ	11:2I:63:LEU:HD11	2.56	0.41
4:32:162:LEU:HD12	4:32:181:MET:CE	2.50	0.41
31:39:152:GLU:HA	31:39:190:GLU:OE2	2.20	0.41
31:39:47:GLY:O	31:39:94:PRO:HA	2.21	0.41
24:3K:22:G:N2	24:3K:23:A:N7	2.68	0.41
24:3L:55:U:N3	24:3L:58:A:OP1	2.36	0.41
5:4E:126:ARG:HH11	5:4E:126:ARG:HG3	1.85	0.41
13:4I:7:VAL:H	32:41:115:ARG:HH12	1.69	0.41
26:1H:2758:A:C4	33:51:67:LEU:HD21	2.56	0.41
1:1G:1114:C:O2'	14:5A:60:SER:O	2.33	0.41
7:6E:113:GLU:HG2	7:6E:119:ARG:HG2	2.03	0.41
8:72:85:ARG:NH1	8:72:134:ILE:HG23	2.35	0.41
16:7A:11:SER:HB2	16:7A:14:ASN:HB3	2.03	0.41
20:BA:48:LYS:O	20:BA:50:GLU:N	2.54	0.41
46:G8:28:LYS:NZ	46:G8:40:GLU:HG3	2.36	0.41
56:Q8:14:VAL:HG11	56:Q8:58:ILE:HG21	2.02	0.41
1:13:1023:G:H3'	1:13:1024:G:C5'	2.48	0.41
1:13:439:A:C8	1:13:440:A:C8	3.09	0.41
1:13:12:U:O2'	1:13:526:C:H4'	2.21	0.41
1:13:626:U:H5"	16:7I:38:TYR:CD2	2.56	0.41
26:14:1328:G:H2'	26:14:1330:C:C4	2.56	0.41
26:14:1685:C:H2'	26:14:1686:C:C6	2.56	0.41
26:14:1827:C:O2'	26:14:1828:G:H5'	2.21	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2273:A:H2'	26:14:2274:A:H8	1.85	0.41
26:14:2305:A:H8	32:49:156:ASP:OD1	2.03	0.41
26:14:2369:A:H2'	26:14:2370:G:H8	1.86	0.41
26:14:2430:A:OP1	62:14:3659:HOH:O	2.22	0.41
26:14:2711:A:P	62:14:3507:HOH:O	2.78	0.41
26:14:442:G:HO2'	26:14:444:C:H6	1.67	0.41
26:14:522:G:H2'	26:14:523:C:C6	2.56	0.41
26:14:564:C:H2'	26:14:565:C:O4'	2.21	0.41
35:15:56:ASN:N	35:15:125:GLY:HA3	2.32	0.41
1:1G:1292:U:H2'	1:1G:1293:G:C8	2.56	0.41
1:1G:1410:G:H2'	1:1G:1411:C:H6	1.86	0.41
1:1G:1414:U:H2'	1:1G:1415:G:C8	2.56	0.41
1:1G:926:G:C6	1:1G:1505:G:C6	3.08	0.41
1:1G:448:A:H2'	1:1G:449:C:O2	2.21	0.41
1:1G:7:G:H5'	1:1G:298:A:O4'	2.20	0.41
1:1G:984:C:H2'	1:1G:985:C:C6	2.56	0.41
26:1H:1204:A:H2	26:1H:1241:A:N1	2.18	0.41
26:1H:125:G:C6	55:P8:10:ARG:HG3	2.56	0.41
26:1H:1533:C:H2'	26:1H:1534:G:C8	2.56	0.41
26:1H:1580:A:OP2	26:1H:1580:A:H8	2.04	0.41
26:1H:2430:A:H8	26:1H:2431:U:C5	2.38	0.41
26:1H:2483:C:C2	38:88:124:LYS:HE3	2.56	0.41
26:1H:654(R):C:C2'	26:1H:654(S):G:H5'	2.51	0.41
27:1J:57:A:C2'	27:1J:58:A:H5'	2.51	0.41
3:2E:58:GLU:N	3:2E:65:ALA:HB3	2.35	0.41
23:2K:2:G:H2'	23:2K:3:C:C6	2.56	0.41
4:32:117:ALA:O	4:32:121:VAL:HG23	2.20	0.41
24:3L:22:G:H8	24:3L:46:G:N2	2.19	0.41
25:4K:13:A:N3	25:4K:13:A:H2'	2.36	0.41
1:1G:1400:C:H4'	25:4L:18:G:C5	2.56	0.41
33:51:137:ASP:OD1	33:51:138:LYS:N	2.44	0.41
26:1H:2748:A:C2	33:51:63:SER:HB3	2.56	0.41
33:51:92:ILE:C	33:51:94:TYR:H	2.24	0.41
6:5E:62:TRP:HH2	6:5E:64:GLN:HG3	1.86	0.41
7:62:36:LYS:O	7:62:40:ALA:N	2.54	0.41
40:65:21:THR:HG22	40:65:21:THR:H	1.56	0.41
34:69:80:PRO:HA	34:69:143:SER:HB2	2.03	0.41
1:1G:1128:C:H4'	9:82:16:ARG:HH22	1.85	0.41
38:88:37:LEU:HD21	38:88:130:LYS:CE	2.50	0.41
38:88:136:ALA:CB	47:H8:52:SER:HB2	2.51	0.41
26:14:565:C:OP1	43:95:82:ARG:NH2	2.54	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:AA:3:ARG:HH22	19:AA:9:VAL:C	2.24	0.41
42:C8:79:PHE:C	42:C8:79:PHE:HD1	2.24	0.41
49:F5:73:LEU:HB3	49:F5:90:ILE:HD11	2.02	0.41
49:F5:8:SER:HB3	49:F5:66:HIS:CD2	2.56	0.41
46:G8:30:VAL:HG12	46:G8:32:PRO:HD3	2.03	0.41
51:H5:43:ILE:O	51:H5:47:VAL:HG23	2.21	0.41
53:J5:41:PRO:HG2	53:J5:44:THR:OG1	2.21	0.41
26:1H:2432:A:C5	49:J8:33:LYS:HG2	2.56	0.41
50:K8:15:LYS:HD3	50:K8:15:LYS:HA	1.76	0.41
44:E8:35:ILE:HG23	53:N8:28:PRO:HD2	2.02	0.41
29:11:61:LEU:HA	29:11:61:LEU:HD13	1.95	0.40
2:12:85:ALA:HB1	2:12:92:TYR:HB3	2.02	0.40
1:13:1097:C:O2'	1:13:1169:A:N3	2.40	0.40
1:13:1429:C:H2'	1:13:1430:C:H6	1.86	0.40
1:13:1489:G:H2'	1:13:1490:C:O4'	2.20	0.40
1:13:21:G:H2'	1:13:22:G:C8	2.56	0.40
1:13:544:G:C6	1:13:545:C:C4	3.09	0.40
26:14:1132:A:H2'	26:14:1133:U:C6	2.56	0.40
26:14:1788:C:C2	26:14:1789:A:C8	3.09	0.40
26:14:2335:A:C8	26:14:2337:G:C5	3.09	0.40
26:14:2340:G:O2'	26:14:2341:G:H5'	2.21	0.40
26:14:2567:G:H2'	26:14:2568:C:C6	2.56	0.40
26:14:26:G:H1'	26:14:515:A:H61	1.86	0.40
26:14:2808:U:O2'	26:14:2809:A:H5'	2.21	0.40
26:14:445:C:O2'	26:14:446:G:H5'	2.21	0.40
26:14:761:A:OP2	61:14:3437:SPE:H111	2.21	0.40
26:14:996:A:C2	26:14:997:G:C8	3.09	0.40
35:15:16:ILE:HB	35:15:54:VAL:HG22	2.01	0.40
29:19:40:THR:OG1	29:19:41:GLY:N	2.53	0.40
10:1A:17:ASP:O	10:1A:21:GLN:HB2	2.21	0.40
1:1G:1248:A:C6	1:1G:1249:C:N4	2.89	0.40
1:1G:1387:G:H2'	1:1G:1388:C:C6	2.57	0.40
1:1G:1471:G:H2'	1:1G:1472:U:C6	2.56	0.40
26:1H:1049:C:H1'	26:1H:1113:U:O2'	2.21	0.40
26:1H:1593:G:H2'	26:1H:1594:G:H8	1.83	0.40
26:1H:2248:C:C5	26:1H:2249:U:C4	3.09	0.40
26:1H:2684:U:H1'	36:68:70:LYS:HD2	2.03	0.40
26:1H:2772:C:H2'	26:1H:2773:C:H6	1.85	0.40
26:1H:696:G:O2'	26:1H:697:C:H5'	2.21	0.40
10:1I:48:THR:OG1	10:1I:62:HIS:CE1	2.74	0.40
30:29:143:ASN:HB2	30:29:147:PRO:HD2	2.03	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
30:29:70:ALA:O	30:29:72:VAL:N	2.53	0.40
11:2A:105:VAL:HG22	11:2A:105:VAL:O	2.20	0.40
11:2A:22:HIS:HB3	11:2A:29:ILE:HG12	2.03	0.40
11:2A:16:SER:OG	11:2A:79:SER:HB3	2.22	0.40
23:2K:54:G:O2'	23:2K:55:5MU:H5"	2.21	0.40
37:35:124:LYS:HE2	37:35:143:GLY:O	2.22	0.40
37:35:47:ASP:HB3	37:35:49:ARG:N	2.37	0.40
24:3L:15:G:C4	24:3L:59:A:C2	3.09	0.40
32:41:72:ARG:NH1	32:41:87:PRO:HD3	2.35	0.40
32:49:76:SER:O	32:49:77:ILE:HG13	2.21	0.40
33:51:2:SER:HB2	33:51:3:ARG:CD	2.43	0.40
33:51:6:ARG:HB3	33:51:65:HIS:CG	2.56	0.40
39:55:107:ASP:C	39:55:107:ASP:OD1	2.60	0.40
35:58:28:THR:HG22	35:58:29:LYS:N	2.36	0.40
34:61:77:LEU:HD13	34:61:140:LEU:HB3	2.03	0.40
8:72:25:ASP:OD1	8:72:25:ASP:N	2.53	0.40
26:1H:637:A:O5'	37:78:116:GLY:HA3	2.21	0.40
37:78:39:LYS:HB2	37:78:45:LEU:CD2	2.50	0.40
42:85:27:LEU:O	42:85:31:SER:HB3	2.22	0.40
43:95:94:LEU:HA	43:95:94:LEU:HD23	1.81	0.40
43:95:18:LEU:O	43:95:96:ILE:HG12	2.22	0.40
40:A8:23:ARG:NH1	40:A8:111:GLU:HG2	2.36	0.40
43:D8:8:GLY:O	43:D8:10:LYS:HE3	2.21	0.40
43:D8:96:ILE:HD13	43:D8:96:ILE:HA	1.86	0.40
44:E8:45:TYR:CE2	44:E8:49:LYS:HD2	2.57	0.40
45:F8:24:GLY:CA	45:F8:82:GLN:HE22	2.33	0.40
56:Q8:14:VAL:C	62:Q8:401:HOH:O	2.59	0.40
26:1H:667:U:O2	56:Q8:2:PRO:HD2	2.21	0.40
1:13:1269:A:H2	1:13:1312:G:N3	2.20	0.40
1:13:1347:G:H22	1:13:1374:A:P	2.44	0.40
1:13:32:A:H2'	1:13:33:A:C8	2.56	0.40
1:13:511:C:C2	1:13:512:U:C5	3.09	0.40
1:13:982:U:H4'	1:13:983:A:O5'	2.21	0.40
26:14:1034:G:H8	26:14:1034:G:OP1	2.03	0.40
26:14:819:A:C4	26:14:1189:A:C2	3.09	0.40
26:14:1372:U:H2'	26:14:1373:A:O4'	2.21	0.40
26:14:1992:G:C8	26:14:1992:G:O5'	2.74	0.40
26:14:217:G:H2'	26:14:218:A:O4'	2.22	0.40
26:14:2075:U:H2'	26:14:2238:G:N2	2.36	0.40
26:14:2666:C:H3'	26:14:2667:C:H6	1.85	0.40
26:14:270(L):U:H1'	34:69:50:ARG:HD3	2.03	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2762:G:H5'	26:14:2763:G:OP2	2.22	0.40
26:14:565:C:H2'	26:14:566:U:O4'	2.21	0.40
26:14:671:C:H2'	26:14:672:C:C6	2.56	0.40
26:14:836:G:C5	26:14:837:C:C4	3.09	0.40
35:15:137:LYS:HE3	35:15:137:LYS:HB3	1.97	0.40
1:1G:266:G:H5"	1:1G:267:C:C5	2.56	0.40
1:1G:272:C:H2'	1:1G:273:A:C8	2.56	0.40
1:1G:522:C:H2'	1:1G:523:A:O4'	2.21	0.40
1:1G:57:G:C6	1:1G:58:C:C4	3.09	0.40
1:1G:652:U:H1'	1:1G:653:A:C2	2.55	0.40
1:1G:852:G:C6	1:1G:853:G:N7	2.90	0.40
26:1H:2189:U:H2'	26:1H:2190:G:C8	2.56	0.40
26:1H:2542:A:H4'	26:1H:2543:G:H8	1.86	0.40
26:1H:2710:C:P	62:1H:4023:HOH:O	2.79	0.40
26:1H:2766:G:H5"	26:1H:2767:C:OP2	2.22	0.40
26:1H:301:G:C4	26:1H:302:C:C5	3.09	0.40
26:1H:425:G:H2'	26:1H:426:C:H6	1.86	0.40
26:1H:57:C:H2'	26:1H:58:G:O4'	2.21	0.40
24:1L:57:G:H2'	24:1L:58:A:H5'	2.03	0.40
3:22:173:VAL:HG12	3:22:175:LEU:HG	2.02	0.40
26:14:1665:A:C4'	36:25:67:LYS:HB2	2.50	0.40
3:2E:133:ALA:O	3:2E:136:GLN:HG3	2.21	0.40
26:14:389:G:H22	37:35:72:PRO:HD3	1.86	0.40
32:41:61:ALA:O	32:41:65:GLY:N	2.52	0.40
32:41:82:LEU:HD22	32:41:82:LEU:HA	1.97	0.40
38:45:132:VAL:HG21	47:D5:81:ARG:NE	2.35	0.40
6:5E:49:ALA:HB1	18:9I:80:PRO:HB3	2.03	0.40
40:65:72:ALA:O	40:65:76:LYS:HG3	2.21	0.40
7:6E:15:ASP:HB3	7:6E:19:GLY:H	1.86	0.40
28:71:59:ARG:HG2	28:71:163:PHE:O	2.21	0.40
9:82:18:PHE:HD2	9:82:62:TYR:CD2	2.38	0.40
40:A8:25:ARG:O	40:A8:39:ILE:HA	2.21	0.40
20:BI:26:ASN:HD22	20:BI:71:THR:HA	1.86	0.40
46:G8:94:LYS:CE	46:G8:95:LYS:H	2.34	0.40
48:I8:38:VAL:HG21	48:I8:59:LEU:HD12	2.04	0.40
48:I8:50:ASN:ND2	48:I8:83:PRO:HD3	2.36	0.40
26:1H:773:U:H5'	29:11:47:GLY:HA3	2.03	0.40
1:13:1125:U:C4	1:13:1126:U:C4	3.09	0.40
1:13:1468:A:H8	1:13:1468:A:O5'	2.04	0.40
1:13:406:G:H2'	1:13:407:G:C8	2.56	0.40
1:13:684:A:N6	1:13:685:G:C6	2.89	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:1392:A:C6	26:14:1393:A:C6	3.09	0.40
26:14:1827:C:H2'	26:14:1828:G:O4'	2.21	0.40
26:14:19:C:H2'	26:14:20:C:C6	2.56	0.40
26:14:2321:G:H2'	26:14:2321:G:N3	2.37	0.40
26:14:2376:A:H2'	26:14:2377:A:O4'	2.21	0.40
26:14:2688:U:C5	26:14:2720:U:OP2	2.75	0.40
26:14:37:C:H4'	26:14:451:C:OP1	2.21	0.40
26:14:608:A:H2'	26:14:609:A:C8	2.57	0.40
26:14:675:A:N3	26:14:2443:C:O2'	2.48	0.40
26:14:838:C:O2'	26:14:839:U:H5'	2.22	0.40
27:16:24:G:C2	27:16:56:G:C2	3.09	0.40
29:19:121:PRO:HB3	29:19:135:PHE:CD2	2.56	0.40
1:1G:1095:U:H2'	1:1G:1096:C:O4'	2.21	0.40
1:1G:1286:A:C8	1:1G:1286:A:C3'	3.05	0.40
1:1G:194:C:H3'	62:1G:1806:HOH:O	2.18	0.40
1:1G:197:A:N1	1:1G:221:C:H4'	2.37	0.40
26:1H:123:G:O6	26:1H:128:C:N3	2.55	0.40
26:1H:1568:G:H5'	29:11:60:ARG:HA	2.02	0.40
26:1H:2663:G:H3'	26:1H:2664:G:H8	1.86	0.40
26:1H:2820:A:OP1	39:98:2:ARG:NH2	2.45	0.40
26:1H:2852:G:H2'	26:1H:2853:C:O4'	2.21	0.40
26:1H:365:C:H2'	26:1H:366:C:O4'	2.22	0.40
26:1H:990:A:H5"	26:1H:991:C:OP1	2.21	0.40
26:1H:990:A:N6	26:1H:1186:G:H1'	2.37	0.40
27:1J:64:C:H2'	27:1J:65:C:C6	2.57	0.40
26:14:2786:U:H5"	30:29:66:HIS:HB3	2.02	0.40
30:29:66:HIS:CG	30:29:67:PHE:N	2.89	0.40
30:29:54:GLN:HG3	30:29:72:VAL:O	2.21	0.40
4:32:13:ARG:O	4:32:14:ARG:HB3	2.21	0.40
12:3I:111:LYS:HD3	12:3I:112:ASP:H	1.86	0.40
32:41:11:TYR:O	32:41:16:ARG:HG3	2.21	0.40
32:41:15:VAL:HG13	32:41:175:LEU:HB2	2.03	0.40
38:45:84:GLY:HA2	38:45:85:LYS:HB2	2.03	0.40
13:4A:89:GLY:HA2	13:4A:92:HIS:HB2	2.03	0.40
5:4E:35:GLY:HA3	5:4E:112:LEU:O	2.22	0.40
39:55:90:ARG:HD2	39:55:94:TYR:HD1	1.86	0.40
3:22:6:HIS:CG	14:5A:49:HIS:HB3	2.57	0.40
34:61:61:ARG:HA	34:61:61:ARG:HD3	1.74	0.40
7:62:47:CYS:HB3	7:62:58:PRO:CB	2.51	0.40
7:62:59:LEU:HD21	7:62:63:LYS:NZ	2.36	0.40
7:62:90:GLU:HG2	7:62:90:GLU:H	1.61	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
40:65:66:ALA:O	40:65:69:VAL:HG13	2.21	0.40
36:68:64:ARG:HB2	36:68:79:PHE:CD1	2.56	0.40
34:69:130:TYR:HD1	34:69:130:TYR:HA	1.74	0.40
5:42:79:GLU:OE1	8:72:104:ARG:HA	2.21	0.40
36:25:76:ALA:HB3	41:75:75:ILE:HB	2.04	0.40
37:78:122:PRO:HA	37:78:142:GLY:HA3	2.03	0.40
37:78:37:GLY:HA2	37:78:41:ARG:NH2	2.36	0.40
37:78:51:PHE:CE2	37:78:53:GLY:HA2	2.56	0.40
1:1G:1149:C:OP2	9:82:9:ARG:NH1	2.55	0.40
20:BA:50:GLU:HA	20:BA:100:ILE:HG12	2.03	0.40
47:D5:60:GLU:HB2	47:D5:66:SER:OG	2.22	0.40
49:J8:58:ILE:HG23	49:J8:90:ILE:HG13	2.03	0.40
29:11:158:ALA:O	29:11:159:ALA:C	2.59	0.40
1:13:1148:U:H2'	1:13:1149:C:O4'	2.21	0.40
1:13:1366:C:O2'	10:1I:60:ARG:NH1	2.36	0.40
1:13:1064:G:OP1	1:13:1386:G:H4'	2.21	0.40
1:13:183:G:H2'	1:13:184:G:C8	2.56	0.40
1:13:193:C:O2'	1:13:194:C:H5'	2.21	0.40
1:13:195:A:C5	1:13:196:A:N1	2.89	0.40
1:13:818:G:O2'	1:13:819:A:H5'	2.21	0.40
1:13:848:C:H6	1:13:848:C:O5'	2.05	0.40
26:14:1176:G:H5'	26:14:1177:A:OP1	2.22	0.40
26:14:1288:U:C2	26:14:1327:C:O2	2.75	0.40
26:14:1434:A:H61	26:14:1558:A:N6	2.18	0.40
26:14:1407:C:C2	26:14:1596:A:C2	3.09	0.40
26:14:2303:G:O4'	32:49:126:ASP:HB3	2.21	0.40
26:14:2526:G:C2	26:14:2538:C:O2	2.75	0.40
26:14:2547:U:O2	36:25:23:ARG:NH2	2.54	0.40
26:14:2699:C:H2'	26:14:2700:C:O4'	2.21	0.40
26:14:527:C:H5"	62:14:3502:HOH:O	2.21	0.40
26:14:602:G:N2	26:14:655:A:C8	2.89	0.40
29:19:30:GLU:HB2	29:19:35:LYS:NZ	2.37	0.40
2:1E:189:ASP:CG	2:1E:191:ASP:HB2	2.41	0.40
2:1E:21:ARG:C	2:1E:23:ARG:H	2.20	0.40
2:1E:82:ARG:HG3	2:1E:92:TYR:CE2	2.56	0.40
1:1G:1101:A:C8	2:12:172:ILE:HD11	2.56	0.40
1:1G:1154:G:N3	1:1G:1155:G:C8	2.89	0.40
1:1G:1331:G:OP1	1:1G:1331:G:H4'	2.22	0.40
1:1G:1489:G:H2'	1:1G:1490:C:O4'	2.21	0.40
1:1G:247:G:OP1	1:1G:247:G:H4'	2.21	0.40
1:1G:359:U:H2'	1:1G:360:A:C8	2.56	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1G:569:C:H1'	1:1G:574:A:C4	2.57	0.40
1:1G:617:G:H1	1:1G:623:C:H42	1.70	0.40
1:1G:76:G:C6	1:1G:77:C:C4	3.10	0.40
26:1H:1368:G:C2	26:1H:1369:G:C8	3.09	0.40
26:1H:1516:U:H2'	26:1H:1517:G:H8	1.86	0.40
26:1H:1568:G:H5"	29:11:61:LEU:HD22	2.02	0.40
26:1H:1578:U:H5	62:1H:4883:HOH:O	2.03	0.40
26:1H:1668:A:N6	26:1H:1676:A:H61	2.19	0.40
26:1H:1669:A:H5"	26:1H:2550:G:OP1	2.21	0.40
26:1H:172:C:H2'	26:1H:173:G:C8	2.57	0.40
26:1H:2228:G:H2'	26:1H:2229:C:C6	2.56	0.40
26:1H:2593:U:O2'	26:1H:2594:C:H5'	2.21	0.40
26:1H:419:C:H2'	26:1H:420:C:O4'	2.22	0.40
26:1H:443:A:H5"	26:1H:444:C:OP1	2.21	0.40
26:1H:617:G:OP2	31:31:43:LYS:HE2	2.22	0.40
26:1H:962:G:H2'	26:1H:963:U:H6	1.85	0.40
1:1G:1112:C:N3	3:22:178:LEU:HD23	2.37	0.40
3:2E:172:ARG:HH21	3:2E:174:PRO:HG2	1.86	0.40
31:31:22:ALA:HB1	31:31:24:LEU:HD13	2.02	0.40
31:39:107:LYS:HA	31:39:107:LYS:HD3	1.61	0.40
12:3A:82:VAL:N	12:3A:106:ASP:OD2	2.33	0.40
24:3L:52:G:C6	24:3L:63:U:C4	3.10	0.40
38:45:32:TYR:OH	38:45:111:GLU:HB2	2.21	0.40
38:45:77:LYS:HE3	38:45:84:GLY:N	2.34	0.40
32:49:117:PHE:CG	32:49:117:PHE:O	2.74	0.40
13:4A:88:ARG:HB2	13:4A:88:ARG:CZ	2.51	0.40
25:4K:14:A:OP2	25:4K:14:A:H3'	2.21	0.40
1:1G:1216:G:H5"	14:5A:5:ALA:HB3	2.02	0.40
8:72:109:ILE:HG22	8:72:137:VAL:HB	2.02	0.40
26:14:1008:C:H4'	42:85:59:ARG:HH12	1.87	0.40
38:88:135:ASP:O	38:88:138:ASP:N	2.44	0.40
17:8A:10:VAL:HG12	17:8A:55:ASP:O	2.22	0.40
17:8I:45:HIS:NE2	17:8I:47:PRO:HG3	2.37	0.40
17:8I:55:ASP:HB3	17:8I:57:VAL:CG1	2.51	0.40
19:AI:41:VAL:CA	19:AI:44:MET:HG3	2.50	0.40
19:AI:50:ALA:HA	19:AI:58:VAL:O	2.21	0.40
46:C5:19:LYS:C	46:C5:21:LYS:H	2.24	0.40
47:D5:127:LYS:HB3	47:D5:127:LYS:HE2	1.81	0.40
47:D5:19:ARG:NH1	47:D5:84:GLU:HB2	2.37	0.40
49:F5:91:LYS:HZ2	49:F5:92:LYS:H	1.69	0.40
26:1H:298:G:OP2	46:G8:84:ARG:HD2	2.22	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:J8:8:SER:OG	49:J8:10:LYS:HG3	2.22	0.40
1:13:1077:G:N2	1:13:1080:A:OP2	2.53	0.40
1:13:1250:A:H2'	1:13:1251:A:C8	2.57	0.40
1:13:267:C:H2'	1:13:268:C:C6	2.57	0.40
1:13:370:C:C2	1:13:392:G:N2	2.90	0.40
1:13:559:A:H2'	1:13:559:A:N3	2.37	0.40
1:13:881:G:H2'	1:13:882:C:O4'	2.22	0.40
1:13:919:A:O2'	1:13:920:U:H5'	2.20	0.40
1:13:955:U:H1'	1:13:1227:A:H61	1.86	0.40
26:14:1268:A:C2	26:14:2013:A:C4	3.10	0.40
26:14:1329:U:H5"	26:14:1330:C:C5	2.49	0.40
26:14:1359:A:N7	26:14:1372:U:O4	2.54	0.40
26:14:1386:C:H2'	26:14:1387:C:C6	2.56	0.40
26:14:1421:G:C2	26:14:1422:G:C8	3.10	0.40
26:14:1620:G:O4'	55:L5:1:MET:N	2.50	0.40
26:14:1688:U:H2'	26:14:1698:A:N6	2.36	0.40
26:14:1945:G:H2'	26:14:1946:U:H6	1.87	0.40
26:14:1961:C:O2'	26:14:1962:C:H5'	2.21	0.40
26:14:2507:C:H5"	26:14:2573:C:N4	2.36	0.40
26:14:2712:U:H1'	26:14:2712(A):A:C8	2.56	0.40
26:14:521:G:H2'	26:14:522:G:C8	2.56	0.40
26:14:635:C:H2'	26:14:636:G:O4'	2.22	0.40
29:19:206:LEU:HD22	29:19:211:ARG:HG2	2.02	0.40
21:1F:8:THR:OG1	21:1F:9:ARG:N	2.54	0.40
1:1G:1072:G:C5	1:1G:1073:U:C4	3.10	0.40
1:1G:1275:A:H2'	1:1G:1276:G:O4'	2.21	0.40
1:1G:198:G:H2'	1:1G:199:G:C8	2.54	0.40
1:1G:286:G:C6	1:1G:287:U:C4	3.10	0.40
26:1H:1728:G:C2	26:1H:1730:U:OP2	2.75	0.40
26:1H:1915:U:H2'	26:1H:1916:A:O4'	2.21	0.40
26:1H:2123:G:H2'	26:1H:2124:G:O4'	2.21	0.40
26:1H:2503:A:P	62:1H:3755:HOH:O	2.78	0.40
26:1H:34:C:C6	26:1H:34:C:OP2	2.75	0.40
26:1H:812:C:H5"	26:1H:1250:G:O2'	2.22	0.40
27:1J:12:C:O2'	48:E5:74:ARG:HG2	2.21	0.40
24:1L:25:C:C2	24:1L:26:A:H1'	2.57	0.40
24:1L:59:A:C8	24:1L:60:U:C4	3.10	0.40
24:1L:9:A:H3'	24:1L:10:G:C8	2.56	0.40
30:21:37:ARG:O	30:21:45:THR:HA	2.21	0.40
3:22:182:ILE:HG22	3:22:203:PHE:HA	2.03	0.40
30:29:24:THR:HG21	30:29:188:VAL:HG22	2.03	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
26:14:2786:U:H5"	30:29:66:HIS:CB	2.52	0.40
11:2I:32:ILE:HG12	11:2I:41:THR:O	2.21	0.40
26:1H:673:C:H5"	31:31:81:PRO:HD2	2.04	0.40
38:45:118:LEU:HD12	38:45:131:ILE:HG23	2.02	0.40
32:49:91:ARG:HB3	32:49:91:ARG:HE	1.43	0.40
25:4L:19:G:O2'	25:4L:20:A:OP2	2.38	0.40
33:51:4:ILE:HD13	33:51:4:ILE:HG21	1.88	0.40
7:6E:43:PHE:O	7:6E:46:ALA:HB3	2.21	0.40
8:72:82:HIS:CE1	8:72:136:GLU:HG3	2.56	0.40
8:72:51:VAL:HG22	8:72:58:TYR:O	2.22	0.40
41:75:86:ILE:HG21	41:75:86:ILE:HD13	1.87	0.40
37:78:125:VAL:O	37:78:144:GLU:HB2	2.22	0.40
9:82:14:VAL:O	9:82:65:VAL:HG23	2.22	0.40
17:8A:45:HIS:HB2	17:8A:65:ILE:CD1	2.51	0.40
9:8E:24:GLY:HA2	9:8E:59:PHE:O	2.21	0.40
18:9A:45:SER:OG	18:9A:46:GLU:N	2.55	0.40
19:AA:16:LEU:HD12	19:AA:19:VAL:HB	2.03	0.40
20:BA:55:ILE:HD13	20:BA:55:ILE:HA	1.97	0.40
20:BA:97:ALA:HA	20:BA:98:PRO:HD3	1.92	0.40
42:C8:108:GLU:HG3	43:D8:44:LYS:CE	2.52	0.40
49:F5:92:LYS:HA	49:F5:95:LEU:HB2	2.03	0.40
45:F8:55:ASN:HB2	45:F8:80:ILE:HG23	2.03	0.40
26:14:2019:A:N7	53:J5:9:LYS:HD2	2.36	0.40
50:K8:60:LEU:HA	50:K8:60:LEU:HD23	1.90	0.40
56:Q8:37:SER:O	56:Q8:40:GLU:N	2.54	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	203/256 (79%)	173 (85%)	25 (12%)	5 (2%)	7 34

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles
2	1E	227/256 (89%)	185 (82%)	39 (17%)	3 (1%)	15 53
3	22	191/239 (80%)	171 (90%)	20 (10%)	0	100 100
3	2E	203/239 (85%)	186 (92%)	17 (8%)	0	100 100
4	32	206/209 (99%)	184 (89%)	21 (10%)	1 (0%)	34 76
4	3E	205/209 (98%)	192 (94%)	12 (6%)	1 (0%)	34 76
5	42	148/162 (91%)	141 (95%)	7 (5%)	0	100 100
5	4E	147/162 (91%)	141 (96%)	5 (3%)	1 (1%)	26 70
6	52	99/101 (98%)	96 (97%)	3 (3%)	0	100 100
6	5E	98/101 (97%)	94 (96%)	4 (4%)	0	100 100
7	62	134/156 (86%)	123 (92%)	10 (8%)	1 (1%)	26 70
7	6E	152/156 (97%)	144 (95%)	8 (5%)	0	100 100
8	72	135/138 (98%)	125 (93%)	8 (6%)	2 (2%)	13 50
8	7E	136/138 (99%)	125 (92%)	10 (7%)	1 (1%)	26 70
9	82	119/128 (93%)	109 (92%)	9 (8%)	1 (1%)	24 66
9	8E	124/128 (97%)	108 (87%)	16 (13%)	0	100 100
10	1A	76/105 (72%)	71 (93%)	5 (7%)	0	100 100
10	1I	92/105 (88%)	85 (92%)	7 (8%)	0	100 100
11	2A	111/129 (86%)	101 (91%)	8 (7%)	2 (2%)	11 45
11	2I	109/129 (84%)	94 (86%)	12 (11%)	3 (3%)	6 30
12	3A	119/132 (90%)	100 (84%)	15 (13%)	4 (3%)	5 25
12	3I	120/132 (91%)	107 (89%)	12 (10%)	1 (1%)	24 66
13	4A	107/126 (85%)	88 (82%)	18 (17%)	1 (1%)	21 64
13	4I	115/126 (91%)	96 (84%)	18 (16%)	1 (1%)	21 64
14	5A	57/61 (93%)	48 (84%)	8 (14%)	1 (2%)	11 45
14	5I	57/61 (93%)	48 (84%)	7 (12%)	2 (4%)	4 24
15	6A	85/89 (96%)	81 (95%)	4 (5%)	0	100 100
15	6I	85/89 (96%)	77 (91%)	8 (9%)	0	100 100
16	7A	82/88 (93%)	76 (93%)	6 (7%)	0	100 100
16	7I	81/88 (92%)	77 (95%)	4 (5%)	0	100 100
17	8A	97/105 (92%)	90 (93%)	7 (7%)	0	100 100
17	8I	97/105 (92%)	92 (95%)	4 (4%)	1 (1%)	19 61

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles
18	9A	65/88 (74%)	62 (95%)	3 (5%)	0	100 100
18	9I	66/88 (75%)	64 (97%)	1 (2%)	1 (2%)	13 50
19	AA	59/93 (63%)	49 (83%)	8 (14%)	2 (3%)	5 25
19	AI	80/93 (86%)	68 (85%)	9 (11%)	3 (4%)	4 22
20	BA	97/106 (92%)	83 (86%)	13 (13%)	1 (1%)	19 61
20	BI	95/106 (90%)	84 (88%)	10 (10%)	1 (1%)	17 58
21	1B	20/27 (74%)	19 (95%)	1 (5%)	0	100 100
21	1F	21/27 (78%)	20 (95%)	1 (5%)	0	100 100
28	71	128/229 (56%)	120 (94%)	8 (6%)	0	100 100
29	11	271/276 (98%)	249 (92%)	17 (6%)	5 (2%)	11 45
29	19	272/276 (99%)	246 (90%)	23 (8%)	3 (1%)	17 58
30	21	201/206 (98%)	158 (79%)	33 (16%)	10 (5%)	3 15
30	29	202/206 (98%)	149 (74%)	42 (21%)	11 (5%)	2 14
31	31	200/210 (95%)	181 (90%)	17 (8%)	2 (1%)	19 61
31	39	202/210 (96%)	162 (80%)	34 (17%)	6 (3%)	5 29
32	41	177/182 (97%)	156 (88%)	18 (10%)	3 (2%)	11 46
32	49	178/182 (98%)	155 (87%)	22 (12%)	1 (1%)	30 72
33	51	172/180 (96%)	141 (82%)	20 (12%)	11 (6%)	2 9
33	59	167/180 (93%)	129 (77%)	32 (19%)	6 (4%)	4 24
34	61	143/148 (97%)	122 (85%)	19 (13%)	2 (1%)	14 51
34	69	143/148 (97%)	111 (78%)	29 (20%)	3 (2%)	9 40
35	15	135/140 (96%)	122 (90%)	13 (10%)	0	100 100
35	58	135/140 (96%)	114 (84%)	17 (13%)	4 (3%)	5 29
36	25	120/122 (98%)	112 (93%)	8 (7%)	0	100 100
36	68	120/122 (98%)	115 (96%)	5 (4%)	0	100 100
37	35	145/150 (97%)	119 (82%)	26 (18%)	0	100 100
37	78	145/150 (97%)	114 (79%)	22 (15%)	9 (6%)	2 10
38	45	136/141 (96%)	115 (85%)	19 (14%)	2 (2%)	13 50
38	88	139/141 (99%)	116 (84%)	18 (13%)	5 (4%)	4 24
39	55	116/118 (98%)	110 (95%)	5 (4%)	1 (1%)	21 64
39	98	116/118 (98%)	104 (90%)	11 (10%)	1 (1%)	21 64

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles
40	65	108/112 (96%)	89 (82%)	16 (15%)	3 (3%)	6 30
40	A8	109/112 (97%)	90 (83%)	19 (17%)	0	100 100
41	75	131/146 (90%)	117 (89%)	12 (9%)	2 (2%)	13 50
41	B8	133/146 (91%)	118 (89%)	15 (11%)	0	100 100
42	85	114/118 (97%)	104 (91%)	9 (8%)	1 (1%)	21 64
42	C8	113/118 (96%)	104 (92%)	6 (5%)	3 (3%)	6 32
43	95	98/101 (97%)	80 (82%)	15 (15%)	3 (3%)	5 28
43	D8	98/101 (97%)	88 (90%)	6 (6%)	4 (4%)	3 20
44	A5	109/113 (96%)	101 (93%)	7 (6%)	1 (1%)	21 64
44	E8	108/113 (96%)	100 (93%)	8 (7%)	0	100 100
45	B5	92/96 (96%)	84 (91%)	6 (6%)	2 (2%)	8 38
45	F8	93/96 (97%)	88 (95%)	5 (5%)	0	100 100
46	C5	102/110 (93%)	74 (72%)	21 (21%)	7 (7%)	1 7
46	G8	101/110 (92%)	81 (80%)	16 (16%)	4 (4%)	4 21
47	D5	175/206 (85%)	136 (78%)	31 (18%)	8 (5%)	3 18
47	H8	168/206 (82%)	136 (81%)	25 (15%)	7 (4%)	3 20
48	E5	74/85 (87%)	66 (89%)	6 (8%)	2 (3%)	6 32
48	I8	75/85 (88%)	67 (89%)	7 (9%)	1 (1%)	15 53
49	F5	92/98 (94%)	79 (86%)	12 (13%)	1 (1%)	17 58
49	J8	94/98 (96%)	83 (88%)	9 (10%)	2 (2%)	9 40
50	G5	67/72 (93%)	61 (91%)	4 (6%)	2 (3%)	5 29
50	K8	66/72 (92%)	60 (91%)	3 (4%)	3 (4%)	3 18
51	H5	56/60 (93%)	53 (95%)	3 (5%)	0	100 100
51	L8	56/60 (93%)	54 (96%)	2 (4%)	0	100 100
52	M8	56/71 (79%)	40 (71%)	15 (27%)	1 (2%)	11 45
53	J5	54/60 (90%)	50 (93%)	4 (7%)	0	100 100
53	N8	46/60 (77%)	43 (94%)	3 (6%)	0	100 100
54	O8	43/54 (80%)	30 (70%)	12 (28%)	1 (2%)	8 36
55	L5	45/49 (92%)	42 (93%)	3 (7%)	0	100 100
55	P8	45/49 (92%)	42 (93%)	3 (7%)	0	100 100
56	M5	62/65 (95%)	52 (84%)	9 (14%)	1 (2%)	12 48

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles
56	Q8	62/65 (95%)	53 (86%)	7 (11%)	2 (3%)	5 27
All	All	11128/12158 (92%)	9762 (88%)	1180 (11%)	186 (2%)	11 46

All (186) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
11	2I	55	LYS
12	3I	48	PRO
18	9I	22	VAL
19	AI	41	VAL
30	21	83	ASP
37	78	37	GLY
42	C8	89	GLU
47	H8	165	VAL
2	12	219	VAL
9	82	118	LYS
20	BA	73	HIS
30	29	25	VAL
30	29	54	GLN
31	39	28	ILE
31	39	84	VAL
32	49	5	VAL
39	55	107	ASP
41	75	10	VAL
41	75	11	GLU
47	D5	53	ILE
47	D5	171	ILE
49	F5	30	VAL
8	7E	86	ILE
29	11	239	ARG
30	21	59	VAL
30	21	60	ASN
30	21	77	ILE
30	21	118	LYS
33	51	10	PRO
33	51	157	TYR
33	51	171	LEU
37	78	25	SER
37	78	35	HIS
38	88	6	ARG
38	88	66	ILE
42	C8	93	LYS

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Mol	Chain	Res	Type
43	D8	45	THR
46	G8	54	LYS
46	G8	81	LYS
47	H8	6	LYS
47	H8	60	GLU
49	J8	94	LEU
56	Q8	50	LEU
11	2A	48	ILE
12	3A	18	VAL
12	3A	26	ALA
14	5A	29	ARG
19	AA	9	VAL
29	19	273	ARG
30	29	59	VAL
30	29	81	ILE
33	59	131	VAL
34	69	113	ARG
40	65	87	PHE
40	65	89	ARG
46	C5	20	TYR
46	C5	92	ASN
47	D5	105	VAL
47	D5	165	VAL
50	G5	48	HIS
14	5I	13	THR
29	11	273	ARG
30	21	72	VAL
33	51	83	TYR
33	51	84	SER
33	51	138	LYS
35	58	97	ARG
35	58	128	HIS
37	78	6	LEU
38	88	7	MET
38	88	134	ARG
50	K8	43	GLN
50	K8	48	HIS
11	2A	101	SER
30	29	9	VAL
30	29	51	PHE
30	29	55	ASN
31	39	25	PRO

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Mol	Chain	Res	Type
31	39	124	LEU
31	39	132	VAL
31	39	167	ALA
33	59	92	ILE
38	45	27	VAL
38	45	90	VAL
48	E5	33	ALA
50	G5	47	ASN
2	1E	238	LEU
4	3E	155	LEU
14	5I	14	PRO
17	8I	79	SER
29	11	3	VAL
29	11	122	ASP
30	21	21	VAL
32	41	96	ARG
32	41	97	ASP
33	51	154	PRO
35	58	22	THR
42	C8	90	VAL
43	D8	49	THR
47	H8	59	LEU
49	J8	86	SER
50	K8	47	ASN
56	Q8	35	GLN
2	12	220	ASP
12	3A	19	ARG
30	29	90	THR
33	59	168	PRO
34	69	112	LYS
40	65	111	GLU
43	95	71	LEU
45	B5	68	ARG
46	C5	29	GLU
47	D5	116	VAL
47	D5	176	PRO
56	M5	34	TRP
2	1E	10	LEU
19	AI	67	VAL
20	BI	71	THR
29	11	240	ALA
30	21	56	PRO

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Mol	Chain	Res	Type
31	31	198	ALA
32	41	5	VAL
33	51	12	PRO
33	51	169	VAL
34	61	12	LEU
34	61	133	HIS
35	58	135	PRO
37	78	14	LYS
37	78	19	VAL
37	78	95	VAL
46	G8	42	VAL
48	I8	9	SER
7	62	147	ALA
8	72	73	ASP
30	29	26	ILE
30	29	45	THR
42	85	93	LYS
44	A5	44	ALA
46	C5	99	CYS
47	D5	161	VAL
2	1E	127	ILE
11	2I	82	VAL
33	51	85	LYS
33	51	170	ARG
37	78	34	GLY
39	98	45	ARG
46	G8	53	PRO
4	32	28	SER
29	19	239	ARG
33	59	167	GLU
46	C5	62	GLU
48	E5	44	ARG
19	AI	40	ILE
30	21	55	ASN
30	21	75	VAL
47	H8	61	LEU
47	H8	141	VAL
12	3A	47	LYS
29	19	3	VAL
13	4I	4	ILE
43	D8	47	VAL
43	95	72	VAL

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Mol	Chain	Res	Type
45	B5	51	VAL
46	C5	85	VAL
5	4E	115	VAL
38	88	27	VAL
43	D8	48	GLY
47	H8	53	ILE
54	O8	52	VAL
2	12	39	ILE
33	59	169	VAL
34	69	144	VAL
43	95	99	ILE
46	C5	3	VAL
11	2I	108	ILE
31	31	24	LEU
37	78	24	GLY
52	M8	5	ILE
2	12	32	ILE
2	12	223	ILE
8	72	100	ILE
13	4A	84	ILE
30	29	77	ILE
47	D5	141	VAL
19	AA	67	VAL
33	59	126	PRO

5.3.2 Protein sidechains (i)

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	179/220 (81%)	143 (80%)	36 (20%)	1 8
2	1E	200/220 (91%)	154 (77%)	46 (23%)	1 5
3	22	154/188 (82%)	128 (83%)	26 (17%)	2 13
3	2E	159/188 (85%)	131 (82%)	28 (18%)	2 12
4	32	180/181 (99%)	151 (84%)	29 (16%)	3 14

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles
4	3E	180/181 (99%)	146 (81%)	34 (19%)	2 10
5	42	114/123 (93%)	87 (76%)	27 (24%)	1 4
5	4E	115/123 (94%)	93 (81%)	22 (19%)	2 10
6	52	90/90 (100%)	74 (82%)	16 (18%)	2 11
6	5E	90/90 (100%)	80 (89%)	10 (11%)	8 29
7	62	114/127 (90%)	88 (77%)	26 (23%)	1 5
7	6E	125/127 (98%)	105 (84%)	20 (16%)	3 15
8	72	118/119 (99%)	101 (86%)	17 (14%)	4 18
8	7E	119/119 (100%)	99 (83%)	20 (17%)	2 13
9	82	92/99 (93%)	73 (79%)	19 (21%)	1 7
9	8E	97/99 (98%)	75 (77%)	22 (23%)	1 5
10	1A	71/92 (77%)	51 (72%)	20 (28%)	0 2
10	1I	81/92 (88%)	73 (90%)	8 (10%)	10 35
11	2A	85/99 (86%)	67 (79%)	18 (21%)	1 6
11	2I	84/99 (85%)	70 (83%)	14 (17%)	3 13
12	3A	102/109 (94%)	79 (78%)	23 (22%)	1 5
12	3I	103/109 (94%)	91 (88%)	12 (12%)	7 27
13	4A	90/101 (89%)	68 (76%)	22 (24%)	1 4
13	4I	94/101 (93%)	74 (79%)	20 (21%)	1 6
14	5A	49/50 (98%)	41 (84%)	8 (16%)	3 14
14	5I	49/50 (98%)	36 (74%)	13 (26%)	0 3
15	6A	79/80 (99%)	71 (90%)	8 (10%)	9 34
15	6I	79/80 (99%)	65 (82%)	14 (18%)	2 11
16	7A	72/74 (97%)	61 (85%)	11 (15%)	3 16
16	7I	72/74 (97%)	58 (81%)	14 (19%)	2 9
17	8A	94/97 (97%)	82 (87%)	12 (13%)	5 23
17	8I	94/97 (97%)	82 (87%)	12 (13%)	5 23
18	9A	58/77 (75%)	47 (81%)	11 (19%)	2 10
18	9I	58/77 (75%)	49 (84%)	9 (16%)	3 16
19	AA	56/80 (70%)	47 (84%)	9 (16%)	3 14
19	AI	72/80 (90%)	61 (85%)	11 (15%)	3 16

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles
20	BA	76/82 (93%)	66 (87%)	10 (13%)	5 22
20	BI	75/82 (92%)	61 (81%)	14 (19%)	2 10
21	1B	17/22 (77%)	16 (94%)	1 (6%)	24 63
21	1F	18/22 (82%)	14 (78%)	4 (22%)	1 5
28	71	108/181 (60%)	87 (81%)	21 (19%)	2 9
29	11	214/218 (98%)	173 (81%)	41 (19%)	2 10
29	19	214/218 (98%)	171 (80%)	43 (20%)	1 8
30	21	162/166 (98%)	118 (73%)	44 (27%)	0 2
30	29	165/166 (99%)	135 (82%)	30 (18%)	2 11
31	31	161/166 (97%)	128 (80%)	33 (20%)	1 7
31	39	163/166 (98%)	126 (77%)	37 (23%)	1 5
32	41	153/156 (98%)	126 (82%)	27 (18%)	2 12
32	49	152/156 (97%)	125 (82%)	27 (18%)	2 11
33	51	143/148 (97%)	106 (74%)	37 (26%)	0 3
33	59	140/148 (95%)	106 (76%)	34 (24%)	1 4
34	61	122/124 (98%)	86 (70%)	36 (30%)	0 2
34	69	122/124 (98%)	89 (73%)	33 (27%)	0 3
35	15	116/119 (98%)	90 (78%)	26 (22%)	1 5
35	58	116/119 (98%)	87 (75%)	29 (25%)	1 3
36	25	100/100 (100%)	76 (76%)	24 (24%)	1 4
36	68	100/100 (100%)	90 (90%)	10 (10%)	9 34
37	35	114/116 (98%)	82 (72%)	32 (28%)	0 2
37	78	114/116 (98%)	78 (68%)	36 (32%)	0 1
38	45	109/111 (98%)	88 (81%)	21 (19%)	2 9
38	88	110/111 (99%)	90 (82%)	20 (18%)	2 11
39	55	101/101 (100%)	80 (79%)	21 (21%)	1 7
39	98	101/101 (100%)	72 (71%)	29 (29%)	0 2
40	65	87/88 (99%)	61 (70%)	26 (30%)	0 2
40	A8	87/88 (99%)	66 (76%)	21 (24%)	1 4
41	75	117/127 (92%)	95 (81%)	22 (19%)	2 10
41	B8	117/127 (92%)	83 (71%)	34 (29%)	0 2

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles
42	85	93/94 (99%)	73 (78%)	20 (22%)	1 6
42	C8	92/94 (98%)	76 (83%)	16 (17%)	2 12
43	95	81/82 (99%)	68 (84%)	13 (16%)	3 15
43	D8	82/82 (100%)	63 (77%)	19 (23%)	1 5
44	A5	91/92 (99%)	74 (81%)	17 (19%)	2 10
44	E8	90/92 (98%)	73 (81%)	17 (19%)	2 10
45	B5	74/78 (95%)	56 (76%)	18 (24%)	1 4
45	F8	77/78 (99%)	64 (83%)	13 (17%)	2 13
46	C5	85/91 (93%)	63 (74%)	22 (26%)	0 3
46	G8	84/91 (92%)	66 (79%)	18 (21%)	1 6
47	D5	156/179 (87%)	124 (80%)	32 (20%)	1 7
47	H8	151/179 (84%)	124 (82%)	27 (18%)	2 11
48	E5	61/67 (91%)	55 (90%)	6 (10%)	10 36
48	I8	62/67 (92%)	53 (86%)	9 (14%)	4 18
49	F5	79/83 (95%)	63 (80%)	16 (20%)	1 7
49	J8	79/83 (95%)	67 (85%)	12 (15%)	3 16
50	G5	63/67 (94%)	46 (73%)	17 (27%)	0 3
50	K8	64/67 (96%)	43 (67%)	21 (33%)	0 1
51	H5	50/52 (96%)	38 (76%)	12 (24%)	1 4
51	L8	50/52 (96%)	40 (80%)	10 (20%)	1 8
52	M8	52/63 (82%)	38 (73%)	14 (27%)	0 3
53	J5	48/52 (92%)	41 (85%)	7 (15%)	4 18
53	N8	43/52 (83%)	34 (79%)	9 (21%)	1 7
54	O8	44/52 (85%)	29 (66%)	15 (34%)	0 1
55	L5	38/42 (90%)	32 (84%)	6 (16%)	3 15
55	P8	38/42 (90%)	31 (82%)	7 (18%)	2 10
56	M5	54/55 (98%)	43 (80%)	11 (20%)	1 7
56	Q8	54/55 (98%)	44 (82%)	10 (18%)	2 10
All	All	9397/10064 (93%)	7493 (80%)	1904 (20%)	1 7

All (1904) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
2	1E	8	LYS
2	1E	11	LEU
2	1E	21	ARG
2	1E	28	PHE
2	1E	33	TYR
2	1E	55	PHE
2	1E	67	THR
2	1E	69	LEU
2	1E	71	VAL
2	1E	79	ASP
2	1E	83	MET
2	1E	86	GLU
2	1E	87	ARG
2	1E	95	GLN
2	1E	96	ARG
2	1E	108	ILE
2	1E	111	ARG
2	1E	118	LEU
2	1E	122	PHE
2	1E	127	ILE
2	1E	135	GLN
2	1E	144	ARG
2	1E	154	LEU
2	1E	155	LEU
2	1E	158	LEU
2	1E	160	ASP
2	1E	163	PHE
2	1E	169	LYS
2	1E	172	ILE
2	1E	178	ARG
2	1E	184	VAL
2	1E	185	ILE
2	1E	187	LEU
2	1E	190	THR
2	1E	197	VAL
2	1E	200	ILE
2	1E	205	ASP
2	1E	209	ARG
2	1E	210	SER
2	1E	211	ILE
2	1E	214	ILE
2	1E	217	ARG
2	1E	222	ILE

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Mol	Chain	Res	Type
2	1E	223	ILE
2	1E	224	GLN
2	1E	230	VAL
3	2E	3	ASN
3	2E	5	ILE
3	2E	8	ILE
3	2E	16	ARG
3	2E	17	ASP
3	2E	21	ARG
3	2E	29	TYR
3	2E	31	HIS
3	2E	32	LEU
3	2E	44	GLU
3	2E	56	ASP
3	2E	58	GLU
3	2E	63	ASN
3	2E	70	VAL
3	2E	72	LYS
3	2E	79	ARG
3	2E	98	ASN
3	2E	105	GLU
3	2E	116	VAL
3	2E	128	PHE
3	2E	132	ARG
3	2E	136	GLN
3	2E	138	VAL
3	2E	167	TRP
3	2E	179	ARG
3	2E	190	ARG
3	2E	202	ILE
3	2E	206	GLU
4	3E	3	ARG
4	3E	5	ILE
4	3E	8	VAL
4	3E	10	ARG
4	3E	12	CYS
4	3E	15	GLU
4	3E	21	LEU
4	3E	31	CYS
4	3E	46	LYS
4	3E	47	ARG
4	3E	58	LEU

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Mol	Chain	Res	Type
4	3E	66	ARG
4	3E	84	LYS
4	3E	85	LYS
4	3E	86	LYS
4	3E	96	LEU
4	3E	106	TYR
4	3E	107	ARG
4	3E	108	LEU
4	3E	115	ARG
4	3E	122	ARG
4	3E	127	THR
4	3E	135	LEU
4	3E	145	GLU
4	3E	152	SER
4	3E	154	ASN
4	3E	160	GLN
4	3E	168	ARG
4	3E	184	LYS
4	3E	188	LEU
4	3E	190	ASP
4	3E	193	ASP
4	3E	196	LEU
4	3E	200	GLU
5	4E	10	MET
5	4E	11	ILE
5	4E	12	LEU
5	4E	16	THR
5	4E	18	ARG
5	4E	31	LEU
5	4E	33	VAL
5	4E	41	VAL
5	4E	56	GLN
5	4E	64	ARG
5	4E	68	GLU
5	4E	71	LEU
5	4E	72	GLN
5	4E	73	ASN
5	4E	75	THR
5	4E	79	GLU
5	4E	91	LEU
5	4E	92	LYS
5	4E	116	THR

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Mol	Chain	Res	Type
5	4E	147	ASP
5	4E	152	ARG
5	4E	153	LYS
6	5E	21	LEU
6	5E	23	LYS
6	5E	25	ILE
6	5E	43	LEU
6	5E	55	ASP
6	5E	64	GLN
6	5E	70	ASP
6	5E	75	LEU
6	5E	86	ARG
6	5E	94	GLN
7	6E	6	ARG
7	6E	12	LEU
7	6E	16	LEU
7	6E	22	LEU
7	6E	27	ILE
7	6E	38	LEU
7	6E	54	THR
7	6E	59	LEU
7	6E	63	LYS
7	6E	73	MET
7	6E	75	VAL
7	6E	79	ARG
7	6E	89	MET
7	6E	91	VAL
7	6E	104	LEU
7	6E	111	ARG
7	6E	113	GLU
7	6E	115	ARG
7	6E	149	ARG
7	6E	155	ARG
8	7E	1	MET
8	7E	19	VAL
8	7E	25	ASP
8	7E	26	VAL
8	7E	36	LEU
8	7E	45	ILE
8	7E	50	ARG
8	7E	52	ASP
8	7E	65	TYR

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Mol	Chain	Res	Type
8	7E	68	ARG
8	7E	80	ILE
8	7E	82	HIS
8	7E	85	ARG
8	7E	91	ARG
8	7E	95	VAL
8	7E	102	ARG
8	7E	104	ARG
8	7E	112	LEU
8	7E	133	LEU
8	7E	137	VAL
9	8E	9	ARG
9	8E	10	ARG
9	8E	16	ARG
9	8E	20	ARG
9	8E	38	GLN
9	8E	42	ARG
9	8E	44	VAL
9	8E	47	LEU
9	8E	50	LEU
9	8E	54	ASP
9	8E	58	HIS
9	8E	75	ASP
9	8E	79	LEU
9	8E	81	ILE
9	8E	83	ARG
9	8E	88	TYR
9	8E	91	ASP
9	8E	92	TYR
9	8E	97	LYS
9	8E	108	VAL
9	8E	121	ARG
9	8E	125	TYR
10	1I	25	GLU
10	1I	28	ARG
10	1I	38	ILE
10	1I	58	ASP
10	1I	60	ARG
10	1I	75	ILE
10	1I	76	ASN
10	1I	81	THR
11	2I	28	THR

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Mol	Chain	Res	Type
11	2I	32	ILE
11	2I	36	ASP
11	2I	48	ILE
11	2I	71	LYS
11	2I	93	GLN
11	2I	99	GLN
11	2I	103	LEU
11	2I	105	VAL
11	2I	108	ILE
11	2I	109	VAL
11	2I	111	ASP
11	2I	114	VAL
11	2I	117	ASN
12	3I	11	VAL
12	3I	18	VAL
12	3I	19	ARG
12	3I	33	ARG
12	3I	54	LYS
12	3I	60	LEU
12	3I	62	SER
12	3I	64	TYR
12	3I	91	LYS
12	3I	111	LYS
12	3I	115	LYS
12	3I	126	LYS
13	4I	9	ILE
13	4I	12	ASN
13	4I	20	THR
13	4I	31	LYS
13	4I	32	GLU
13	4I	44	ARG
13	4I	45	VAL
13	4I	48	LEU
13	4I	56	LEU
13	4I	64	TRP
13	4I	67	GLU
13	4I	70	LEU
13	4I	88	ARG
13	4I	101	GLN
13	4I	102	ARG
13	4I	105	THR
13	4I	106	ASN

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Mol	Chain	Res	Type
13	4I	108	ARG
13	4I	110	ARG
13	4I	111	LYS
14	5I	3	ARG
14	5I	4	LYS
14	5I	6	LEU
14	5I	7	ILE
14	5I	15	LYS
14	5I	18	VAL
14	5I	22	THR
14	5I	23	ARG
14	5I	32	SER
14	5I	33	VAL
14	5I	35	ARG
14	5I	41	ARG
14	5I	50	LYS
15	6I	7	GLU
15	6I	10	LYS
15	6I	26	GLU
15	6I	35	ARG
15	6I	38	ARG
15	6I	39	LEU
15	6I	41	GLU
15	6I	47	LYS
15	6I	48	LYS
15	6I	54	ARG
15	6I	59	MET
15	6I	62	GLN
15	6I	66	LEU
15	6I	71	GLN
16	7I	8	ARG
16	7I	11	SER
16	7I	18	ARG
16	7I	19	ILE
16	7I	20	VAL
16	7I	27	LYS
16	7I	28	ARG
16	7I	47	ASP
16	7I	50	LYS
16	7I	54	GLU
16	7I	55	ARG
16	7I	67	THR

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Mol	Chain	Res	Type
16	7I	72	ARG
16	7I	83	GLU
17	8I	24	GLU
17	8I	25	ARG
17	8I	31	LEU
17	8I	38	ARG
17	8I	48	GLU
17	8I	52	LYS
17	8I	53	LEU
17	8I	63	ARG
17	8I	68	ARG
17	8I	74	LEU
17	8I	97	SER
17	8I	100	LYS
18	9I	31	LEU
18	9I	32	ARG
18	9I	42	ARG
18	9I	45	SER
18	9I	55	ARG
18	9I	76	LEU
18	9I	82	THR
18	9I	85	LEU
18	9I	86	VAL
19	AI	3	ARG
19	AI	6	LYS
19	AI	12	ASP
19	AI	31	ILE
19	AI	37	ARG
19	AI	43	GLU
19	AI	44	MET
19	AI	58	VAL
19	AI	67	VAL
19	AI	77	THR
19	AI	78	ARG
20	BI	9	ASN
20	BI	10	LEU
20	BI	13	LEU
20	BI	19	SER
20	BI	24	LEU
20	BI	30	LYS
20	BI	37	SER
20	BI	51	GLU

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Mol	Chain	Res	Type
20	BI	53	LEU
20	BI	56	MET
20	BI	72	LEU
20	BI	73	HIS
20	BI	75	ASN
20	BI	87	LYS
21	1F	6	ARG
21	1F	8	THR
21	1F	9	ARG
21	1F	10	ARG
28	71	3	HIS
28	71	6	ARG
28	71	10	LEU
28	71	14	VAL
28	71	15	ASP
28	71	23	ASP
28	71	34	THR
28	71	36	LYS
28	71	37	PHE
28	71	42	GLU
28	71	53	ARG
28	71	55	ASP
28	71	59	ARG
28	71	166	ASP
28	71	168	THR
28	71	174	PRO
28	71	180	PHE
28	71	211	SER
28	71	216	THR
28	71	218	MET
28	71	224	ILE
29	11	3	VAL
29	11	4	LYS
29	11	13	ARG
29	11	16	MET
29	11	17	THR
29	11	23	GLU
29	11	33	LEU
29	11	35	LYS
29	11	37	LEU
29	11	38	LYS
29	11	39	LYS

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Mol	Chain	Res	Type
29	11	43	ARG
29	11	64	ILE
29	11	65	ILE
29	11	72	LYS
29	11	78	LYS
29	11	83	GLU
29	11	88	ARG
29	11	94	LEU
29	11	103	ARG
29	11	105	ILE
29	11	113	VAL
29	11	117	VAL
29	11	126	GLN
29	11	142	VAL
29	11	154	LYS
29	11	165	ILE
29	11	183	ARG
29	11	192	THR
29	11	200	ASP
29	11	211	ARG
29	11	212	SER
29	11	217	ARG
29	11	221	VAL
29	11	229	VAL
29	11	242	ARG
29	11	257	LEU
29	11	259	THR
29	11	260	ARG
29	11	271	ILE
29	11	273	ARG
30	21	12	THR
30	21	14	ILE
30	21	16	ARG
30	21	23	VAL
30	21	25	VAL
30	21	26	ILE
30	21	45	THR
30	21	47	VAL
30	21	48	GLN
30	21	49	LEU
30	21	52	LEU
30	21	54	GLN

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Mol	Chain	Res	Type
30	21	55	ASN
30	21	57	LYS
30	21	63	LEU
30	21	67	PHE
30	21	76	ARG
30	21	78	LEU
30	21	82	ARG
30	21	89	ASP
30	21	91	VAL
30	21	93	VAL
30	21	101	ARG
30	21	105	THR
30	21	111	ARG
30	21	116	VAL
30	21	119	ARG
30	21	128	SER
30	21	138	PRO
30	21	140	SER
30	21	146	THR
30	21	152	LYS
30	21	154	LYS
30	21	166	THR
30	21	175	VAL
30	21	179	GLU
30	21	181	LEU
30	21	182	LEU
30	21	188	VAL
30	21	195	LEU
30	21	196	VAL
30	21	197	ILE
30	21	201	THR
30	21	203	LYS
31	31	7	TYR
31	31	13	SER
31	31	15	SER
31	31	17	ARG
31	31	18	ARG
31	31	28	ILE
31	31	32	LEU
31	31	33	LEU
31	31	57	VAL
31	31	64	ILE

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Mol	Chain	Res	Type
31	31	67	GLN
31	31	74	ARG
31	31	101	LEU
31	31	106	ARG
31	31	107	LYS
31	31	117	ARG
31	31	124	LEU
31	31	127	GLU
31	31	135	LYS
31	31	136	THR
31	31	140	LEU
31	31	145	GLU
31	31	158	THR
31	31	164	ARG
31	31	168	ARG
31	31	170	LEU
31	31	174	VAL
31	31	181	LEU
31	31	191	ARG
31	31	197	ASP
31	31	200	GLU
31	31	203	GLN
31	31	204	ASN
32	41	22	ARG
32	41	26	GLN
32	41	28	VAL
32	41	43	LEU
32	41	45	GLU
32	41	48	GLU
32	41	51	ARG
32	41	53	LEU
32	41	58	GLN
32	41	67	LYS
32	41	70	VAL
32	41	80	PHE
32	41	82	LEU
32	41	90	LEU
32	41	94	LEU
32	41	96	ARG
32	41	101	ILE
32	41	104	GLU
32	41	108	ASN

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Mol	Chain	Res	Type
32	41	116	ASP
32	41	121	ASN
32	41	128	ARG
32	41	136	ARG
32	41	145	THR
32	41	152	LEU
32	41	162	THR
32	41	165	THR
33	51	2	SER
33	51	3	ARG
33	51	4	ILE
33	51	7	LEU
33	51	9	ILE
33	51	11	VAL
33	51	16	SER
33	51	24	VAL
33	51	37	VAL
33	51	40	GLU
33	51	41	MET
33	51	42	ARG
33	51	43	VAL
33	51	50	VAL
33	51	56	SER
33	51	63	SER
33	51	64	LEU
33	51	68	THR
33	51	71	LEU
33	51	77	LYS
33	51	80	SER
33	51	81	GLU
33	51	83	TYR
33	51	86	GLU
33	51	88	LEU
33	51	98	LEU
33	51	104	GLU
33	51	105	LEU
33	51	116	GLU
33	51	127	GLU
33	51	129	THR
33	51	131	VAL
33	51	139	GLN
33	51	149	ARG

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Mol	Chain	Res	Type
33	51	152	ARG
33	51	167	GLU
33	51	170	ARG
34	61	2	LYS
34	61	9	LEU
34	61	12	LEU
34	61	20	ASP
34	61	21	VAL
34	61	25	TYR
34	61	38	LEU
34	61	41	GLU
34	61	44	LEU
34	61	47	LEU
34	61	48	GLU
34	61	50	ARG
34	61	54	GLN
34	61	64	GLU
34	61	75	LEU
34	61	77	LEU
34	61	82	ARG
34	61	85	GLU
34	61	92	VAL
34	61	95	LYS
34	61	99	GLU
34	61	101	LEU
34	61	102	SER
34	61	108	THR
34	61	110	ASP
34	61	114	LEU
34	61	117	GLU
34	61	122	GLU
34	61	127	VAL
34	61	131	LYS
34	61	135	GLU
34	61	136	VAL
34	61	139	GLN
34	61	140	LEU
34	61	142	VAL
34	61	145	VAL
35	58	1	MET
35	58	2	LYS
35	58	5	VAL

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Mol	Chain	Res	Type
35	58	7	LYS
35	58	14	VAL
35	58	28	THR
35	58	32	THR
35	58	33	LEU
35	58	34	LEU
35	58	42	TRP
35	58	43	THR
35	58	48	MET
35	58	58	ASP
35	58	60	ILE
35	58	61	ARG
35	58	65	LYS
35	58	67	LEU
35	58	87	LEU
35	58	89	LYS
35	58	90	MET
35	58	96	GLU
35	58	97	ARG
35	58	99	LEU
35	58	120	LEU
35	58	127	ASP
35	58	128	HIS
35	58	130	HIS
35	58	131	GLN
35	58	137	LYS
36	68	8	LEU
36	68	17	ARG
36	68	23	ARG
36	68	31	LYS
36	68	38	VAL
36	68	47	ILE
36	68	58	VAL
36	68	66	LYS
36	68	78	ARG
36	68	94	ARG
37	78	1	MET
37	78	5	ASP
37	78	6	LEU
37	78	7	ARG
37	78	10	PRO
37	78	13	ASN

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Mol	Chain	Res	Type
37	78	14	LYS
37	78	15	ARG
37	78	16	ARG
37	78	18	ARG
37	78	21	ARG
37	78	25	SER
37	78	27	HIS
37	78	41	ARG
37	78	45	LEU
37	78	49	ARG
37	78	50	ARG
37	78	56	SER
37	78	61	ARG
37	78	65	ARG
37	78	75	ILE
37	78	77	ARG
37	78	88	LEU
37	78	90	ARG
37	78	96	THR
37	78	98	GLU
37	78	99	LEU
37	78	100	LEU
37	78	101	VAL
37	78	105	LEU
37	78	106	LEU
37	78	112	LEU
37	78	115	LEU
37	78	135	LEU
37	78	144	GLU
37	78	146	VAL
38	88	5	ARG
38	88	7	MET
38	88	11	LYS
38	88	25	ASP
38	88	26	TYR
38	88	35	VAL
38	88	45	GLN
38	88	48	GLU
38	88	55	VAL
38	88	56	ARG
38	88	58	PHE
38	88	59	ARG

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Mol	Chain	Res	Type
38	88	60	ARG
38	88	64	ILE
38	88	67	ARG
38	88	81	VAL
38	88	83	MET
38	88	110	THR
38	88	129	THR
38	88	138	ASP
39	98	1	MET
39	98	2	ARG
39	98	6	SER
39	98	9	LYS
39	98	17	ARG
39	98	18	LEU
39	98	28	LEU
39	98	29	LEU
39	98	33	ARG
39	98	34	ILE
39	98	36	THR
39	98	44	LEU
39	98	48	VAL
39	98	49	ASP
39	98	54	LEU
39	98	57	ARG
39	98	59	ASP
39	98	63	ARG
39	98	65	LEU
39	98	79	LEU
39	98	82	GLU
39	98	91	GLN
39	98	96	ARG
39	98	104	ARG
39	98	105	ARG
39	98	111	LEU
39	98	113	LEU
39	98	116	LEU
39	98	118	GLU
40	A8	3	ARG
40	A8	8	GLU
40	A8	14	VAL
40	A8	24	LEU
40	A8	30	ARG

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Mol	Chain	Res	Type
40	A8	32	LEU
40	A8	35	ILE
40	A8	36	TYR
40	A8	43	GLU
40	A8	46	VAL
40	A8	54	LEU
40	A8	56	LEU
40	A8	58	LEU
40	A8	69	VAL
40	A8	73	LEU
40	A8	83	LYS
40	A8	89	ARG
40	A8	97	ARG
40	A8	101	LEU
40	A8	106	ARG
40	A8	110	LEU
41	B8	2	ASN
41	B8	3	ARG
41	B8	10	VAL
41	B8	16	ARG
41	B8	21	GLU
41	B8	27	THR
41	B8	33	LYS
41	B8	38	ASN
41	B8	39	ARG
41	B8	42	ILE
41	B8	50	ILE
41	B8	55	ASN
41	B8	58	ASN
41	B8	59	THR
41	B8	62	THR
41	B8	64	ARG
41	B8	74	ARG
41	B8	85	LYS
41	B8	86	ILE
41	B8	87	ASP
41	B8	88	ILE
41	B8	96	ARG
41	B8	98	LYS
41	B8	99	LEU
41	B8	106	SER
41	B8	108	ARG

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Mol	Chain	Res	Type
41	B8	110	ILE
41	B8	111	ARG
41	B8	112	ARG
41	B8	115	ARG
41	B8	118	ARG
41	B8	128	GLU
41	B8	129	ARG
41	B8	133	GLU
42	C8	8	VAL
42	C8	13	LYS
42	C8	27	LEU
42	C8	51	LYS
42	C8	52	ARG
42	C8	74	LEU
42	C8	77	SER
42	C8	79	PHE
42	C8	85	LYS
42	C8	89	GLU
42	C8	92	ARG
42	C8	93	LYS
42	C8	94	ASN
42	C8	104	GLN
42	C8	108	GLU
42	C8	112	ARG
43	D8	6	LYS
43	D8	7	THR
43	D8	14	VAL
43	D8	18	LEU
43	D8	20	LEU
43	D8	21	ARG
43	D8	25	LEU
43	D8	33	VAL
43	D8	35	LEU
43	D8	37	VAL
43	D8	38	LEU
43	D8	40	LEU
43	D8	49	THR
43	D8	52	VAL
43	D8	58	VAL
43	D8	64	HIS
43	D8	73	SER
43	D8	79	VAL

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Mol	Chain	Res	Type
43	D8	95	LEU
44	E8	4	LYS
44	E8	11	ARG
44	E8	15	ARG
44	E8	19	LEU
44	E8	20	VAL
44	E8	23	LEU
44	E8	41	LYS
44	E8	42	ARG
44	E8	51	LEU
44	E8	52	GLU
44	E8	64	MET
44	E8	65	LEU
44	E8	76	VAL
44	E8	78	GLU
44	E8	92	ARG
44	E8	96	ILE
44	E8	107	LEU
45	F8	12	VAL
45	F8	23	GLU
45	F8	27	THR
45	F8	38	GLU
45	F8	41	ASN
45	F8	45	THR
45	F8	49	VAL
45	F8	66	LEU
45	F8	68	ARG
45	F8	70	LEU
45	F8	72	LYS
45	F8	78	LYS
45	F8	80	ILE
46	G8	3	VAL
46	G8	4	LYS
46	G8	6	HIS
46	G8	24	VAL
46	G8	38	ILE
46	G8	43	ASN
46	G8	44	ILE
46	G8	57	GLN
46	G8	64	GLU
46	G8	67	LEU
46	G8	71	LYS

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Mol	Chain	Res	Type
46	G8	75	ILE
46	G8	82	PRO
46	G8	85	VAL
46	G8	86	ARG
46	G8	96	ILE
46	G8	98	VAL
46	G8	102	CYS
47	H8	1	MET
47	H8	11	GLU
47	H8	19	ARG
47	H8	35	ARG
47	H8	37	VAL
47	H8	41	LEU
47	H8	46	LYS
47	H8	61	LEU
47	H8	71	VAL
47	H8	72	ARG
47	H8	76	LEU
47	H8	77	ASP
47	H8	80	ARG
47	H8	82	ARG
47	H8	86	VAL
47	H8	91	LEU
47	H8	94	GLU
47	H8	105	VAL
47	H8	117	LEU
47	H8	119	GLU
47	H8	121	HIS
47	H8	126	VAL
47	H8	128	VAL
47	H8	132	ASN
47	H8	142	SER
47	H8	154	ASP
47	H8	169	GLU
48	I8	10	THR
48	I8	36	ILE
48	I8	38	VAL
48	I8	40	GLN
48	I8	43	THR
48	I8	49	LYS
48	I8	67	VAL
48	I8	74	ARG

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Mol	Chain	Res	Type
48	I8	82	ARG
49	J8	4	VAL
49	J8	41	ARG
49	J8	73	LEU
49	J8	74	VAL
49	J8	80	LEU
49	J8	81	LYS
49	J8	85	LEU
49	J8	91	LYS
49	J8	93	GLU
49	J8	94	LEU
49	J8	95	LEU
49	J8	96	LYS
50	K8	2	LYS
50	K8	3	LEU
50	K8	8	LYS
50	K8	14	ARG
50	K8	15	LYS
50	K8	16	LEU
50	K8	17	SER
50	K8	19	VAL
50	K8	24	LEU
50	K8	32	LEU
50	K8	41	ILE
50	K8	46	GLN
50	K8	47	ASN
50	K8	48	HIS
50	K8	51	ARG
50	K8	53	LEU
50	K8	54	LYS
50	K8	55	ARG
50	K8	62	THR
50	K8	64	LEU
50	K8	67	LYS
51	L8	4	LEU
51	L8	6	VAL
51	L8	8	LEU
51	L8	9	VAL
51	L8	31	LEU
51	L8	33	GLN
51	L8	40	THR
51	L8	44	ARG

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Mol	Chain	Res	Type
51	L8	57	GLU
51	L8	58	VAL
52	M8	5	ILE
52	M8	10	VAL
52	M8	15	ILE
52	M8	16	CYS
52	M8	18	CYS
52	M8	34	GLU
52	M8	36	CYS
52	M8	38	LYS
52	M8	40	HIS
52	M8	42	PHE
52	M8	47	GLN
52	M8	60	GLN
52	M8	61	ARG
52	M8	63	TYR
53	N8	3	LYS
53	N8	6	VAL
53	N8	11	THR
53	N8	15	ARG
53	N8	16	ARG
53	N8	29	THR
53	N8	37	LYS
53	N8	40	LYS
53	N8	46	CYS
54	O8	10	LEU
54	O8	12	GLU
54	O8	16	CYS
54	O8	21	TYR
54	O8	26	ASN
54	O8	27	LYS
54	O8	30	THR
54	O8	33	LYS
54	O8	34	LEU
54	O8	36	LEU
54	O8	37	ARG
54	O8	39	TYR
54	O8	42	TRP
54	O8	44	ARG
54	O8	47	THR
55	P8	4	THR
55	P8	8	ASN

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Mol	Chain	Res	Type
55	P8	14	LYS
55	P8	24	THR
55	P8	29	LYS
55	P8	32	LYS
55	P8	43	THR
56	Q8	4	MET
56	Q8	14	VAL
56	Q8	34	TRP
56	Q8	35	GLN
56	Q8	44	LYS
56	Q8	46	ARG
56	Q8	58	ILE
56	Q8	59	LYS
56	Q8	60	LEU
56	Q8	62	LEU
2	12	19	HIS
2	12	21	ARG
2	12	24	TRP
2	12	32	ILE
2	12	36	ARG
2	12	40	HIS
2	12	41	ILE
2	12	44	LEU
2	12	49	GLU
2	12	51	LEU
2	12	52	GLU
2	12	55	PHE
2	12	58	ILE
2	12	60	ASP
2	12	76	GLN
2	12	80	ILE
2	12	84	GLU
2	12	90	MET
2	12	94	ASN
2	12	96	ARG
2	12	108	ILE
2	12	126	GLU
2	12	129	GLU
2	12	145	LEU
2	12	155	LEU
2	12	160	ASP
2	12	165	VAL

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Mol	Chain	Res	Type
2	12	172	ILE
2	12	179	LYS
2	12	185	ILE
2	12	187	LEU
2	12	191	ASP
2	12	193	ASP
2	12	220	ASP
2	12	221	LEU
2	12	224	GLN
3	22	4	LYS
3	22	6	HIS
3	22	11	ARG
3	22	16	ARG
3	22	18	TRP
3	22	22	TRP
3	22	29	TYR
3	22	34	LEU
3	22	47	LEU
3	22	52	LEU
3	22	59	ARG
3	22	76	VAL
3	22	85	ARG
3	22	90	GLU
3	22	94	LEU
3	22	104	GLN
3	22	124	ILE
3	22	128	PHE
3	22	131	ARG
3	22	164	ARG
3	22	167	TRP
3	22	175	LEU
3	22	179	ARG
3	22	182	ILE
3	22	202	ILE
3	22	204	LEU
4	32	3	ARG
4	32	4	TYR
4	32	5	ILE
4	32	8	VAL
4	32	12	CYS
4	32	13	ARG
4	32	17	VAL

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Mol	Chain	Res	Type
4	32	21	LEU
4	32	26	CYS
4	32	30	LYS
4	32	35	ARG
4	32	58	LEU
4	32	59	ARG
4	32	73	ARG
4	32	83	SER
4	32	122	ARG
4	32	127	THR
4	32	131	ARG
4	32	134	ASP
4	32	135	LEU
4	32	137	SER
4	32	150	GLU
4	32	152	SER
4	32	170	VAL
4	32	187	ARG
4	32	191	ARG
4	32	196	LEU
4	32	200	GLU
4	32	204	ILE
5	42	10	MET
5	42	12	LEU
5	42	13	ILE
5	42	16	THR
5	42	25	ARG
5	42	31	LEU
5	42	40	ARG
5	42	43	LEU
5	42	47	LYS
5	42	51	VAL
5	42	53	LEU
5	42	66	MET
5	42	73	ASN
5	42	78	HIS
5	42	79	GLU
5	42	81	GLU
5	42	83	GLU
5	42	87	SER
5	42	90	VAL
5	42	101	ILE

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Mol	Chain	Res	Type
5	42	118	ILE
5	42	120	THR
5	42	126	ARG
5	42	127	ASN
5	42	135	THR
5	42	150	ARG
5	42	151	LEU
6	52	3	ARG
6	52	14	LEU
6	52	16	GLN
6	52	17	SER
6	52	21	LEU
6	52	24	GLU
6	52	28	ARG
6	52	40	VAL
6	52	47	ARG
6	52	63	TYR
6	52	71	ARG
6	52	83	ASP
6	52	86	ARG
6	52	87	ARG
6	52	93	SER
6	52	98	LEU
7	62	8	GLU
7	62	9	VAL
7	62	11	GLN
7	62	13	GLN
7	62	16	LEU
7	62	24	THR
7	62	37	ASN
7	62	38	LEU
7	62	52	GLU
7	62	59	LEU
7	62	60	LYS
7	62	66	VAL
7	62	67	GLU
7	62	70	LYS
7	62	72	ARG
7	62	73	MET
7	62	90	GLU
7	62	94	ARG
7	62	98	SER

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Mol	Chain	Res	Type
7	62	104	LEU
7	62	114	ARG
7	62	131	LYS
7	62	137	LYS
7	62	142	GLU
7	62	143	ARG
7	62	144	MET
8	72	12	ARG
8	72	25	ASP
8	72	33	GLU
8	72	39	LEU
8	72	41	ARG
8	72	52	ASP
8	72	56	LYS
8	72	78	GLN
8	72	82	HIS
8	72	83	ILE
8	72	97	VAL
8	72	99	GLU
8	72	102	ARG
8	72	104	ARG
8	72	120	THR
8	72	127	LEU
8	72	138	TRP
9	82	7	THR
9	82	10	ARG
9	82	19	LEU
9	82	20	ARG
9	82	27	THR
9	82	33	PHE
9	82	34	ASN
9	82	36	TYR
9	82	42	ARG
9	82	47	LEU
9	82	54	ASP
9	82	56	LEU
9	82	91	ASP
9	82	95	LYS
9	82	102	LEU
9	82	112	LYS
9	82	113	LYS
9	82	117	HIS

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Mol	Chain	Res	Type
9	82	124	GLN
10	1A	13	HIS
10	1A	17	ASP
10	1A	21	GLN
10	1A	22	LYS
10	1A	24	VAL
10	1A	33	GLN
10	1A	34	VAL
10	1A	38	ILE
10	1A	43	ARG
10	1A	55	LYS
10	1A	58	ASP
10	1A	59	SER
10	1A	62	HIS
10	1A	65	LEU
10	1A	66	ARG
10	1A	70	ARG
10	1A	74	ILE
10	1A	75	ILE
10	1A	79	ARG
10	1A	84	GLN
11	2A	14	VAL
11	2A	18	ARG
11	2A	29	ILE
11	2A	30	VAL
11	2A	31	THR
11	2A	54	ARG
11	2A	63	LEU
11	2A	78	GLN
11	2A	79	SER
11	2A	80	VAL
11	2A	87	THR
11	2A	93	GLN
11	2A	99	GLN
11	2A	103	LEU
11	2A	105	VAL
11	2A	109	VAL
11	2A	114	VAL
11	2A	119	CYS
12	3A	22	SER
12	3A	23	LYS
12	3A	24	VAL

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Mol	Chain	Res	Type
12	3A	27	LEU
12	3A	33	ARG
12	3A	34	ARG
12	3A	39	VAL
12	3A	41	ARG
12	3A	54	LYS
12	3A	57	LYS
12	3A	59	ARG
12	3A	60	LEU
12	3A	62	SER
12	3A	64	TYR
12	3A	66	VAL
12	3A	83	VAL
12	3A	84	LEU
12	3A	85	ILE
12	3A	89	ARG
12	3A	98	TYR
12	3A	102	ARG
12	3A	111	LYS
12	3A	118	SER
13	4A	9	ILE
13	4A	11	ARG
13	4A	12	ASN
13	4A	13	LYS
13	4A	17	VAL
13	4A	37	THR
13	4A	39	ILE
13	4A	47	ASP
13	4A	49	THR
13	4A	54	VAL
13	4A	58	GLU
13	4A	62	ASN
13	4A	64	TRP
13	4A	66	LEU
13	4A	81	LEU
13	4A	86	CYS
13	4A	88	ARG
13	4A	94	ARG
13	4A	101	GLN
13	4A	103	THR
13	4A	108	ARG
13	4A	117	VAL

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Mol	Chain	Res	Type
14	5A	7	ILE
14	5A	16	PHE
14	5A	22	THR
14	5A	25	VAL
14	5A	29	ARG
14	5A	33	VAL
14	5A	42	ILE
14	5A	43	CYS
15	6A	3	ILE
15	6A	17	ARG
15	6A	22	THR
15	6A	41	GLU
15	6A	47	LYS
15	6A	68	ARG
15	6A	71	GLN
15	6A	84	LYS
16	7A	1	MET
16	7A	2	VAL
16	7A	6	LEU
16	7A	27	LYS
16	7A	29	ASP
16	7A	55	ARG
16	7A	65	GLN
16	7A	67	THR
16	7A	74	LEU
16	7A	76	GLN
16	7A	81	ARG
17	8A	10	VAL
17	8A	16	GLN
17	8A	24	GLU
17	8A	53	LEU
17	8A	57	VAL
17	8A	60	ILE
17	8A	62	SER
17	8A	63	ARG
17	8A	68	ARG
17	8A	74	LEU
17	8A	90	ILE
17	8A	100	LYS
18	9A	23	LYS
18	9A	26	LEU
18	9A	29	PHE

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Mol	Chain	Res	Type
18	9A	32	ARG
18	9A	36	ASN
18	9A	42	ARG
18	9A	45	SER
18	9A	53	ARG
18	9A	68	LYS
18	9A	82	THR
18	9A	83	GLU
19	AA	6	LYS
19	AA	14	HIS
19	AA	15	LEU
19	AA	20	LEU
19	AA	21	GLU
19	AA	33	THR
19	AA	34	TRP
19	AA	43	GLU
19	AA	60	VAL
20	BA	13	LEU
20	BA	31	SER
20	BA	33	ILE
20	BA	45	GLN
20	BA	61	SER
20	BA	72	LEU
20	BA	74	LYS
20	BA	75	ASN
20	BA	85	MET
20	BA	99	LEU
21	1B	10	ARG
29	19	10	THR
29	19	13	ARG
29	19	24	ILE
29	19	27	THR
29	19	28	GLU
29	19	31	LYS
29	19	33	LEU
29	19	34	VAL
29	19	35	LYS
29	19	37	LEU
29	19	38	LYS
29	19	46	GLN
29	19	49	ILE
29	19	54	ARG

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Mol	Chain	Res	Type
29	19	61	LEU
29	19	64	ILE
29	19	69	ARG
29	19	78	LYS
29	19	82	ILE
29	19	88	ARG
29	19	94	LEU
29	19	99	ASP
29	19	103	ARG
29	19	105	ILE
29	19	111	LEU
29	19	116	GLN
29	19	141	VAL
29	19	147	LEU
29	19	182	LEU
29	19	192	THR
29	19	211	ARG
29	19	217	ARG
29	19	239	ARG
29	19	242	ARG
29	19	244	ARG
29	19	253	GLN
29	19	257	LEU
29	19	260	ARG
29	19	262	ARG
29	19	266	SER
29	19	268	ARG
29	19	271	ILE
29	19	273	ARG
30	29	1	MET
30	29	7	VAL
30	29	12	THR
30	29	21	VAL
30	29	27	LEU
30	29	45	THR
30	29	49	LEU
30	29	57	LYS
30	29	66	HIS
30	29	67	PHE
30	29	73	GLU
30	29	76	ARG
30	29	78	LEU

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Mol	Chain	Res	Type
30	29	79	ARG
30	29	89	ASP
30	29	111	ARG
30	29	113	PHE
30	29	116	VAL
30	29	119	ARG
30	29	121	ASN
30	29	144	ARG
30	29	149	ARG
30	29	164	ARG
30	29	170	LEU
30	29	175	VAL
30	29	178	GLU
30	29	181	LEU
30	29	197	ILE
30	29	200	GLU
30	29	201	THR
31	39	7	TYR
31	39	8	GLN
31	39	18	ARG
31	39	20	LEU
31	39	23	ASP
31	39	24	LEU
31	39	28	ILE
31	39	29	ASN
31	39	33	LEU
31	39	38	ARG
31	39	40	GLN
31	39	53	THR
31	39	57	VAL
31	39	62	ARG
31	39	67	GLN
31	39	68	LYS
31	39	69	HIS
31	39	70	THR
31	39	82	ILE
31	39	83	PHE
31	39	88	VAL
31	39	110	LEU
31	39	112	MET
31	39	123	LEU
31	39	140	LEU

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Mol	Chain	Res	Type
31	39	144	LYS
31	39	145	GLU
31	39	153	SER
31	39	158	THR
31	39	165	ARG
31	39	181	LEU
31	39	190	GLU
31	39	191	ARG
31	39	194	MET
31	39	196	LEU
31	39	197	ASP
31	39	205	ARG
32	49	9	ARG
32	49	13	GLU
32	49	20	ILE
32	49	26	GLN
32	49	33	ARG
32	49	35	GLU
32	49	39	ILE
32	49	40	ASN
32	49	45	GLU
32	49	48	GLU
32	49	51	ARG
32	49	62	LEU
32	49	66	GLN
32	49	80	PHE
32	49	82	LEU
32	49	91	ARG
32	49	101	ILE
32	49	109	VAL
32	49	130	ASN
32	49	133	LEU
32	49	136	ARG
32	49	139	LEU
32	49	152	LEU
32	49	153	ARG
32	49	156	ASP
32	49	157	ILE
32	49	159	VAL
33	59	6	ARG
33	59	7	LEU
33	59	32	GLU

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Mol	Chain	Res	Type
33	59	50	VAL
33	59	59	ARG
33	59	70	THR
33	59	72	ILE
33	59	83	TYR
33	59	85	LYS
33	59	86	GLU
33	59	89	ILE
33	59	101	ARG
33	59	103	LEU
33	59	105	LEU
33	59	107	VAL
33	59	116	GLU
33	59	119	GLU
33	59	122	THR
33	59	123	PHE
33	59	125	VAL
33	59	127	GLU
33	59	129	THR
33	59	130	ARG
33	59	131	VAL
33	59	136	ILE
33	59	137	ASP
33	59	139	GLN
33	59	147	ASN
33	59	152	ARG
33	59	157	TYR
33	59	158	HIS
33	59	160	LYS
33	59	167	GLU
33	59	171	LEU
34	69	1	MET
34	69	2	LYS
34	69	4	ILE
34	69	7	GLU
34	69	27	ARG
34	69	37	VAL
34	69	47	LEU
34	69	56	LYS
34	69	58	LEU
34	69	61	ARG
34	69	62	LYS

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Mol	Chain	Res	Type
34	69	67	ARG
34	69	68	LEU
34	69	75	LEU
34	69	76	THR
34	69	77	LEU
34	69	78	THR
34	69	82	ARG
34	69	93	THR
34	69	101	LEU
34	69	104	GLN
34	69	105	HIS
34	69	109	ILE
34	69	114	LEU
34	69	117	GLU
34	69	118	LYS
34	69	125	GLU
34	69	127	VAL
34	69	130	TYR
34	69	135	GLU
34	69	141	LYS
34	69	142	VAL
34	69	145	VAL
35	15	4	TYR
35	15	5	VAL
35	15	9	VAL
35	15	12	ARG
35	15	14	VAL
35	15	15	LEU
35	15	28	THR
35	15	32	THR
35	15	33	LEU
35	15	34	LEU
35	15	41	ASP
35	15	43	THR
35	15	46	VAL
35	15	48	MET
35	15	59	LYS
35	15	61	ARG
35	15	63	THR
35	15	68	GLU
35	15	74	ARG
35	15	87	LEU

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Mol	Chain	Res	Type
35	15	93	THR
35	15	94	HIS
35	15	99	LEU
35	15	130	HIS
35	15	131	GLN
35	15	134	ARG
36	25	1	MET
36	25	5	GLN
36	25	8	LEU
36	25	9	GLU
36	25	10	VAL
36	25	14	THR
36	25	22	ILE
36	25	24	VAL
36	25	28	SER
36	25	29	ASN
36	25	38	VAL
36	25	45	GLU
36	25	49	ARG
36	25	52	VAL
36	25	87	ILE
36	25	91	LEU
36	25	92	GLU
36	25	94	ARG
36	25	99	PHE
36	25	108	GLU
36	25	113	LYS
36	25	114	ILE
36	25	116	SER
36	25	117	LEU
37	35	4	SER
37	35	19	VAL
37	35	21	ARG
37	35	41	ARG
37	35	45	LEU
37	35	52	GLU
37	35	55	ARG
37	35	58	THR
37	35	59	LEU
37	35	62	LEU
37	35	67	MET
37	35	75	ILE

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Mol	Chain	Res	Type
37	35	76	LYS
37	35	79	ARG
37	35	83	VAL
37	35	84	ASN
37	35	85	LEU
37	35	88	LEU
37	35	96	THR
37	35	98	GLU
37	35	102	ARG
37	35	105	LEU
37	35	112	LEU
37	35	114	ILE
37	35	125	VAL
37	35	132	LYS
37	35	133	SER
37	35	135	LEU
37	35	138	LEU
37	35	144	GLU
37	35	146	VAL
37	35	147	LEU
38	45	2	LEU
38	45	3	MET
38	45	10	ARG
38	45	18	LYS
38	45	22	LYS
38	45	26	TYR
38	45	27	VAL
38	45	38	GLU
38	45	45	GLN
38	45	54	MET
38	45	56	ARG
38	45	59	ARG
38	45	75	THR
38	45	76	LYS
38	45	83	MET
38	45	103	MET
38	45	110	THR
38	45	118	LEU
38	45	127	ILE
38	45	137	TYR
38	45	138	ASP
39	55	1	MET

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Mol	Chain	Res	Type
39	55	2	ARG
39	55	6	SER
39	55	9	LYS
39	55	17	ARG
39	55	18	LEU
39	55	24	GLN
39	55	28	LEU
39	55	29	LEU
39	55	33	ARG
39	55	35	THR
39	55	44	LEU
39	55	57	ARG
39	55	65	LEU
39	55	75	LEU
39	55	79	LEU
39	55	81	ASP
39	55	82	GLU
39	55	88	ARG
39	55	102	GLU
39	55	105	ARG
40	65	3	ARG
40	65	8	GLU
40	65	12	PHE
40	65	14	VAL
40	65	17	ARG
40	65	19	LYS
40	65	20	ARG
40	65	21	THR
40	65	27	SER
40	65	29	PHE
40	65	42	ASP
40	65	43	GLU
40	65	50	SER
40	65	62	LYS
40	65	69	VAL
40	65	71	ARG
40	65	73	LEU
40	65	75	GLU
40	65	89	ARG
40	65	98	VAL
40	65	101	LEU
40	65	106	ARG

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Mol	Chain	Res	Type
40	65	107	GLU
40	65	110	LEU
40	65	111	GLU
40	65	112	PHE
41	75	13	ARG
41	75	17	THR
41	75	21	GLU
41	75	27	THR
41	75	36	GLU
41	75	42	ILE
41	75	50	ILE
41	75	55	ASN
41	75	57	PHE
41	75	62	THR
41	75	64	ARG
41	75	65	LYS
41	75	67	SER
41	75	86	ILE
41	75	87	ASP
41	75	88	ILE
41	75	91	ARG
41	75	93	ARG
41	75	105	LEU
41	75	112	ARG
41	75	115	ARG
41	75	118	ARG
42	85	5	LYS
42	85	15	LYS
42	85	20	LEU
42	85	27	LEU
42	85	34	LYS
42	85	52	ARG
42	85	55	ARG
42	85	58	ARG
42	85	59	ARG
42	85	64	ARG
42	85	70	ARG
42	85	71	GLN
42	85	74	LEU
42	85	92	ARG
42	85	97	ASP
42	85	101	ARG

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Mol	Chain	Res	Type
42	85	104	GLN
42	85	112	ARG
42	85	114	LYS
42	85	117	GLN
43	95	7	THR
43	95	12	TYR
43	95	13	ARG
43	95	28	GLU
43	95	33	VAL
43	95	35	LEU
43	95	49	THR
43	95	57	VAL
43	95	66	ARG
43	95	69	LYS
43	95	71	LEU
43	95	91	TYR
43	95	95	LEU
44	A5	11	ARG
44	A5	18	ARG
44	A5	23	LEU
44	A5	39	THR
44	A5	41	LYS
44	A5	50	VAL
44	A5	51	LEU
44	A5	60	ASN
44	A5	65	LEU
44	A5	67	ASP
44	A5	70	TYR
44	A5	78	GLU
44	A5	100	THR
44	A5	106	ILE
44	A5	107	LEU
44	A5	110	LYS
44	A5	111	HIS
45	B5	12	VAL
45	B5	23	GLU
45	B5	27	THR
45	B5	30	VAL
45	B5	35	THR
45	B5	48	LYS
45	B5	49	VAL
45	B5	53	LYS

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Mol	Chain	Res	Type
45	B5	60	ARG
45	B5	63	LYS
45	B5	66	LEU
45	B5	69	TYR
45	B5	70	LEU
45	B5	75	ASP
45	B5	80	ILE
45	B5	81	VAL
45	B5	82	GLN
45	B5	92	LEU
46	C5	3	VAL
46	C5	14	LEU
46	C5	23	ARG
46	C5	24	VAL
46	C5	43	ASN
46	C5	51	VAL
46	C5	55	TYR
46	C5	61	ILE
46	C5	62	GLU
46	C5	70	SER
46	C5	71	LYS
46	C5	72	VAL
46	C5	76	CYS
46	C5	84	ARG
46	C5	85	VAL
46	C5	86	ARG
46	C5	87	LYS
46	C5	89	PHE
46	C5	94	LYS
46	C5	97	ARG
46	C5	98	VAL
46	C5	101	LYS
47	D5	4	ARG
47	D5	5	LEU
47	D5	14	LYS
47	D5	16	SER
47	D5	18	LEU
47	D5	19	ARG
47	D5	24	LEU
47	D5	28	MET
47	D5	30	ASN
47	D5	40	ASP

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Mol	Chain	Res	Type
47	D5	41	LEU
47	D5	60	GLU
47	D5	61	LEU
47	D5	63	ASP
47	D5	71	VAL
47	D5	72	ARG
47	D5	76	LEU
47	D5	84	GLU
47	D5	87	ASP
47	D5	88	PHE
47	D5	91	LEU
47	D5	103	ARG
47	D5	119	GLU
47	D5	121	HIS
47	D5	126	VAL
47	D5	129	SER
47	D5	136	PHE
47	D5	138	GLU
47	D5	161	VAL
47	D5	165	VAL
47	D5	168	GLU
47	D5	170	THR
48	E5	12	ASN
48	E5	36	ILE
48	E5	38	VAL
48	E5	43	THR
48	E5	46	LYS
48	E5	63	VAL
49	F5	3	LYS
49	F5	5	CYS
49	F5	11	ARG
49	F5	19	GLN
49	F5	38	SER
49	F5	40	ARG
49	F5	72	GLU
49	F5	74	VAL
49	F5	76	ARG
49	F5	78	LYS
49	F5	82	LEU
49	F5	83	GLU
49	F5	89	GLU
49	F5	90	ILE

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Mol	Chain	Res	Type
49	F5	91	LYS
49	F5	95	LEU
50	G5	6	VAL
50	G5	10	LEU
50	G5	15	LYS
50	G5	24	LEU
50	G5	26	ARG
50	G5	34	GLU
50	G5	35	LEU
50	G5	44	LEU
50	G5	46	GLN
50	G5	47	ASN
50	G5	48	HIS
50	G5	50	ILE
50	G5	53	LEU
50	G5	56	GLN
50	G5	60	LEU
50	G5	64	LEU
50	G5	65	ASN
51	H5	5	LYS
51	H5	8	LEU
51	H5	9	VAL
51	H5	17	LYS
51	H5	24	LYS
51	H5	30	ARG
51	H5	33	GLN
51	H5	35	ARG
51	H5	38	GLU
51	H5	40	THR
51	H5	44	ARG
51	H5	55	ARG
53	J5	6	VAL
53	J5	26	THR
53	J5	29	THR
53	J5	36	CYS
53	J5	44	THR
53	J5	48	GLU
53	J5	55	ARG
55	L5	1	MET
55	L5	4	THR
55	L5	8	ASN
55	L5	32	LYS

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Mol	Chain	Res	Type
55	L5	36	GLN
55	L5	43	THR
56	M5	11	LYS
56	M5	13	ARG
56	M5	29	LYS
56	M5	31	HIS
56	M5	32	LEU
56	M5	37	SER
56	M5	41	ILE
56	M5	49	VAL
56	M5	58	ILE
56	M5	59	LYS
56	M5	60	LEU

Some sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (13) such sidechains are listed below:

Mol	Chain	Res	Type
7	6E	84	ASN
19	AI	23	ASN
28	71	188	ASN
2	12	16	HIS
2	12	19	HIS
2	12	40	HIS
5	42	78	HIS
18	9A	36	ASN
29	19	227	ASN
29	19	253	GLN
40	65	95	HIS
50	G5	47	ASN
51	H5	33	GLN

5.3.3 RNA

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	13	1497/1522 (98%)	331 (22%)	33 (2%)
1	1G	1506/1522 (98%)	352 (23%)	33 (2%)
22	1K	68/76 (89%)	27 (39%)	4 (5%)
23	2K	76/77 (98%)	21 (27%)	2 (2%)
23	2L	76/77 (98%)	15 (19%)	2 (2%)
24	1L	61/76 (80%)	24 (39%)	2 (3%)

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Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
24	3K	67/76 (88%)	37 (55%)	2 (2%)
24	3L	69/76 (90%)	30 (43%)	2 (2%)
25	4K	17/30 (56%)	9 (52%)	1 (5%)
25	4L	18/30 (60%)	12 (66%)	1 (5%)
26	14	2821/2917 (96%)	675 (23%)	42 (1%)
26	1H	2850/2917 (97%)	630 (22%)	52 (1%)
27	16	121/122 (99%)	24 (19%)	1 (0%)
27	1J	121/122 (99%)	32 (26%)	3 (2%)
All	All	9368/9640 (97%)	2219 (23%)	180 (1%)

All (2219) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	13	2	U
1	13	5	U
1	13	6	G
1	13	7	G
1	13	9	G
1	13	21	G
1	13	32	A
1	13	39	G
1	13	47	C
1	13	48	C
1	13	49	U
1	13	50	A
1	13	51	A
1	13	54	C
1	13	61	G
1	13	65	U
1	13	66	G
1	13	69	G
1	13	74	C
1	13	75	C
1	13	76	G
1	13	77	C
1	13	95	G
1	13	96	G
1	13	97	U
1	13	99	C
1	13	101	A
1	13	116	A
1	13	121	C

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Mol	Chain	Res	Type
1	13	122	G
1	13	131	C
1	13	142	G
1	13	144	G
1	13	147	G
1	13	151	A
1	13	158	G
1	13	160	A
1	13	162	A
1	13	163	C
1	13	169	C
1	13	172	A
1	13	173	U
1	13	174	C
1	13	186(F)	C
1	13	188	U
1	13	189	U
1	13	191(A)	G
1	13	195	A
1	13	197	A
1	13	199	G
1	13	201	C
1	13	208	U
1	13	209	U
1	13	210	U
1	13	217	C
1	13	220	G
1	13	222	U
1	13	226	G
1	13	231	G
1	13	243	A
1	13	244	U
1	13	245	C
1	13	247	G
1	13	248	C
1	13	251	G
1	13	256	U
1	13	262	A
1	13	266	G
1	13	267	C
1	13	270	A
1	13	274	A

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Mol	Chain	Res	Type
1	13	289	G
1	13	316	G
1	13	321	A
1	13	326	G
1	13	328	C
1	13	329	A
1	13	330	C
1	13	332	G
1	13	341	C
1	13	343	U
1	13	344	A
1	13	345	C
1	13	346	G
1	13	347	G
1	13	349	A
1	13	352	C
1	13	353	A
1	13	354	G
1	13	363	A
1	13	367	U
1	13	372	C
1	13	384	G
1	13	388	G
1	13	390	C
1	13	392	G
1	13	397	A
1	13	398	C
1	13	406	G
1	13	412	A
1	13	414	A
1	13	418	C
1	13	423	G
1	13	424	G
1	13	429	U
1	13	430	A
1	13	439	A
1	13	451	A
1	13	452	A
1	13	466	C
1	13	467	G
1	13	482	A
1	13	483	C

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Mol	Chain	Res	Type
1	13	485	G
1	13	496	A
1	13	497	U
1	13	504	C
1	13	505	G
1	13	509	A
1	13	510	A
1	13	511	C
1	13	518	C
1	13	521	G
1	13	524	G
1	13	527	G
1	13	531	U
1	13	533	A
1	13	536	C
1	13	547	A
1	13	559	A
1	13	561	U
1	13	562	C
1	13	572	A
1	13	573	A
1	13	576	G
1	13	577	G
1	13	596	C
1	13	607	A
1	13	620	C
1	13	630	G
1	13	631	G
1	13	632	A
1	13	639	G
1	13	653	A
1	13	665	A
1	13	687	A
1	13	688	G
1	13	723	U
1	13	724	G
1	13	749	C
1	13	750	G
1	13	755	G
1	13	759	A
1	13	769	G
1	13	777	A

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Mol	Chain	Res	Type
1	13	787	A
1	13	792	A
1	13	793	U
1	13	794	A
1	13	812	C
1	13	813	U
1	13	817	C
1	13	828	A
1	13	836	G
1	13	841	U
1	13	842	C
1	13	843	U
1	13	848	C
1	13	859	A
1	13	864	A
1	13	870	U
1	13	872	A
1	13	876	G
1	13	877	C
1	13	884	U
1	13	902	G
1	13	914	A
1	13	916	G
1	13	922	G
1	13	926	G
1	13	927	G
1	13	931	C
1	13	934	C
1	13	935	A
1	13	936	C
1	13	942	G
1	13	960	U
1	13	968	A
1	13	969	A
1	13	974	A
1	13	975	A
1	13	976	G
1	13	977	A
1	13	983	A
1	13	991	U
1	13	993	G
1	13	999	U

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Mol	Chain	Res	Type
1	13	1004	A
1	13	1006	C
1	13	1007	C
1	13	1008	C
1	13	1009	G
1	13	1012	U
1	13	1021	G
1	13	1024	G
1	13	1025	U
1	13	1026	G
1	13	1028	C
1	13	1028(B)	C
1	13	1029	G
1	13	1031	G
1	13	1032	A
1	13	1032(A)	G
1	13	1032(B)	G
1	13	1033	G
1	13	1040	U
1	13	1042	G
1	13	1054	C
1	13	1055	A
1	13	1064	G
1	13	1065	U
1	13	1066	C
1	13	1081	G
1	13	1094	G
1	13	1095	U
1	13	1101	A
1	13	1110	A
1	13	1125	U
1	13	1126	U
1	13	1127	G
1	13	1129	C
1	13	1130	A
1	13	1132	C
1	13	1136	U
1	13	1137	C
1	13	1138	G
1	13	1139	G
1	13	1146	A
1	13	1150	U

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Mol	Chain	Res	Type
1	13	1152	A
1	13	1154	G
1	13	1157	A
1	13	1158	C
1	13	1159	U
1	13	1177	G
1	13	1178	G
1	13	1179	A
1	13	1181	G
1	13	1182	G
1	13	1184	G
1	13	1188	A
1	13	1190	G
1	13	1191	A
1	13	1193	G
1	13	1196	U
1	13	1197	G
1	13	1201	A
1	13	1211	U
1	13	1212	U
1	13	1213	A
1	13	1225	A
1	13	1226	C
1	13	1227	A
1	13	1236	A
1	13	1238	A
1	13	1240	U
1	13	1241	G
1	13	1253	G
1	13	1256	A
1	13	1257	U
1	13	1258	G
1	13	1270	C
1	13	1272	G
1	13	1275	A
1	13	1278	U
1	13	1279	A
1	13	1280	A
1	13	1281	U
1	13	1286	A
1	13	1287	A
1	13	1288	A

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Mol	Chain	Res	Type
1	13	1290	G
1	13	1299	A
1	13	1300	G
1	13	1301	U
1	13	1302	U
1	13	1303	C
1	13	1312	G
1	13	1317	C
1	13	1320	C
1	13	1331	G
1	13	1335	C
1	13	1336	C
1	13	1337	G
1	13	1340	A
1	13	1346	A
1	13	1347	G
1	13	1350	A
1	13	1353	G
1	13	1358	U
1	13	1361	G
1	13	1362(A)	C
1	13	1363	A
1	13	1368	G
1	13	1370	G
1	13	1379	G
1	13	1381	U
1	13	1388	C
1	13	1398	A
1	13	1419	G
1	13	1442	G
1	13	1443	G
1	13	1446	A
1	13	1447	G
1	13	1450	U
1	13	1452	C
1	13	1453	G
1	13	1455	G
1	13	1487	G
1	13	1492	A
1	13	1497	G
1	13	1499	A
1	13	1502	A

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Mol	Chain	Res	Type
1	13	1504	G
1	13	1506	U
1	13	1507	A
1	13	1517	G
1	13	1529	G
1	13	1530	G
1	13	1534	A
1	13	1535	C
22	1K	4	U
22	1K	5	C
22	1K	6	G
22	1K	7	U
22	1K	9	A
22	1K	15	G
22	1K	18	G
22	1K	22	G
22	1K	26	A
22	1K	29	U
22	1K	45	G
22	1K	48	C
22	1K	50	C
22	1K	51	A
22	1K	56	C
22	1K	60	U
22	1K	61	C
22	1K	63	U
22	1K	68	G
22	1K	69	A
22	1K	70	C
22	1K	71	C
22	1K	72	C
22	1K	73	A
22	1K	74	C
22	1K	75	C
22	1K	76	A
23	2K	2	G
23	2K	6	G
23	2K	8	4SU
23	2K	9	G
23	2K	13	C
23	2K	15	G
23	2K	16	C

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Mol	Chain	Res	Type
23	2K	19	G
23	2K	20	G
23	2K	22	A
23	2K	23	G
23	2K	31	G
23	2K	32	G
23	2K	33	OMC
23	2K	47	7MG
23	2K	48	U
23	2K	49	C
23	2K	50	G
23	2K	53	G
23	2K	68	C
23	2K	77	A
24	3K	2	G
24	3K	3	G
24	3K	4	U
24	3K	5	C
24	3K	7	U
24	3K	8	U
24	3K	9	A
24	3K	10	G
24	3K	11	C
24	3K	15	G
24	3K	23	A
24	3K	24	G
24	3K	26	A
24	3K	31	A
24	3K	34	U
24	3K	35	U
24	3K	37	A
24	3K	40	C
24	3K	45	G
24	3K	46	G
24	3K	49	G
24	3K	51	A
24	3K	52	G
24	3K	55	U
24	3K	56	C
24	3K	58	A
24	3K	59	A
24	3K	60	U

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Mol	Chain	Res	Type
24	3K	61	C
24	3K	62	C
24	3K	64	G
24	3K	65	C
24	3K	66	A
24	3K	69	A
24	3K	72	C
24	3K	73	A
24	3K	76	A
25	4K	7	G
25	4K	8	A
25	4K	10	G
25	4K	11	U
25	4K	14	A
25	4K	15	A
25	4K	23	A
25	4K	24	A
25	4K	25	A
26	1H	9	U
26	1H	27	G
26	1H	34	C
26	1H	46	C
26	1H	51	G
26	1H	61	G
26	1H	63	U
26	1H	64	A
26	1H	70	G
26	1H	71	A
26	1H	74	A
26	1H	75	G
26	1H	85	G
26	1H	118	A
26	1H	119	A
26	1H	120	U
26	1H	125	G
26	1H	138	G
26	1H	153	C
26	1H	155	C
26	1H	163	U
26	1H	164	U
26	1H	171	G
26	1H	181	A

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Mol	Chain	Res	Type
26	1H	188	G
26	1H	196	A
26	1H	197	A
26	1H	199	A
26	1H	214	G
26	1H	215	G
26	1H	216	A
26	1H	221	A
26	1H	222	A
26	1H	223	A
26	1H	228	A
26	1H	229	A
26	1H	233	A
26	1H	248	G
26	1H	250	G
26	1H	252	G
26	1H	261	G
26	1H	266	G
26	1H	269	U
26	1H	270(C)	C
26	1H	270(K)	C
26	1H	270(L)	U
26	1H	270(M)	U
26	1H	270(N)	G
26	1H	270(O)	U
26	1H	270(P)	C
26	1H	270(Y)	G
26	1H	271(C)	U
26	1H	271	G
26	1H	274	G
26	1H	275	G
26	1H	277	C
26	1H	278	A
26	1H	279	C
26	1H	283	A
26	1H	295	G
26	1H	299	A
26	1H	311	A
26	1H	323	G
26	1H	324	A
26	1H	326	G
26	1H	329	G

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Mol	Chain	Res	Type
26	1H	330	A
26	1H	335	C
26	1H	342	G
26	1H	346	A
26	1H	352	G
26	1H	357	A
26	1H	363	G
26	1H	370	G
26	1H	372	G
26	1H	375	C
26	1H	382	G
26	1H	386	G
26	1H	405	U
26	1H	406	G
26	1H	407	G
26	1H	408	G
26	1H	411	G
26	1H	412	A
26	1H	418	G
26	1H	421	U
26	1H	427	U
26	1H	428	A
26	1H	434	U
26	1H	443	A
26	1H	444	C
26	1H	447	A
26	1H	448	U
26	1H	451	C
26	1H	455	C
26	1H	456	C
26	1H	457	A
26	1H	470	A
26	1H	471	A
26	1H	481	G
26	1H	482	A
26	1H	483	A
26	1H	494	G
26	1H	505	A
26	1H	508	G
26	1H	509	C
26	1H	510	C
26	1H	513	A

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Mol	Chain	Res	Type
26	1H	528	A
26	1H	529	A
26	1H	531	C
26	1H	532	A
26	1H	533	G
26	1H	546	C
26	1H	548	A
26	1H	549	G
26	1H	550	G
26	1H	556	G
26	1H	563	G
26	1H	564	C
26	1H	573	G
26	1H	575	A
26	1H	584	C
26	1H	586	A
26	1H	588	U
26	1H	603	A
26	1H	607	U
26	1H	614	U
26	1H	615	G
26	1H	617	G
26	1H	621	A
26	1H	622	G
26	1H	626	U
26	1H	627	A
26	1H	637	A
26	1H	645	C
26	1H	646	A
26	1H	647	G
26	1H	654	A
26	1H	654(A)	A
26	1H	654(D)	G
26	1H	654(O)	G
26	1H	654(P)	G
26	1H	654(Q)	C
26	1H	654(S)	G
26	1H	654(T)	A
26	1H	654(V)	A
26	1H	678	C
26	1H	686	G
26	1H	689	A

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Mol	Chain	Res	Type
26	1H	699	A
26	1H	714	U
26	1H	717	G
26	1H	730	C
26	1H	731	C
26	1H	739	G
26	1H	747	U
26	1H	757	U
26	1H	764	A
26	1H	765	G
26	1H	771	G
26	1H	776	G
26	1H	782	A
26	1H	784	A
26	1H	785	G
26	1H	790	C
26	1H	792	G
26	1H	802	A
26	1H	805	G
26	1H	812	C
26	1H	816	C
26	1H	827	U
26	1H	828	U
26	1H	832	G
26	1H	836	G
26	1H	845	G
26	1H	846	C
26	1H	853	G
26	1H	858	U
26	1H	859	G
26	1H	860	U
26	1H	866	A
26	1H	878	A
26	1H	879	G
26	1H	882	G
26	1H	894	C
26	1H	899	A
26	1H	900	A
26	1H	901	A
26	1H	902	C
26	1H	907	U
26	1H	910	A

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Mol	Chain	Res	Type
26	1H	914	C
26	1H	917	A
26	1H	926	A
26	1H	932	G
26	1H	938	G
26	1H	941	A
26	1H	946	G
26	1H	947	G
26	1H	953	A
26	1H	959	A
26	1H	961	C
26	1H	962	G
26	1H	968	G
26	1H	974	G
26	1H	974(A)	C
26	1H	983	A
26	1H	990	A
26	1H	995	C
26	1H	996	A
26	1H	1003	G
26	1H	1005	C
26	1H	1011	G
26	1H	1012	U
26	1H	1013	C
26	1H	1020	A
26	1H	1022	G
26	1H	1023	U
26	1H	1025	G
26	1H	1026	U
26	1H	1027	A
26	1H	1031	G
26	1H	1033	U
26	1H	1040	C
26	1H	1045	A
26	1H	1046	A
26	1H	1047	G
26	1H	1054	A
26	1H	1055	G
26	1H	1071	G
26	1H	1072	C
26	1H	1079	C
26	1H	1080	A

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Mol	Chain	Res	Type
26	1H	1081	U
26	1H	1082	U
26	1H	1085	A
26	1H	1086	A
26	1H	1087	G
26	1H	1088	A
26	1H	1090	U
26	1H	1093	G
26	1H	1101	U
26	1H	1106	G
26	1H	1109	C
26	1H	1110	G
26	1H	1111	A
26	1H	1112	G
26	1H	1126	A
26	1H	1129	A
26	1H	1130	U
26	1H	1135	C
26	1H	1136	G
26	1H	1138	G
26	1H	1139	G
26	1H	1142	U
26	1H	1142(A)	A
26	1H	1144	G
26	1H	1149	G
26	1H	1170	G
26	1H	1177	A
26	1H	1178	C
26	1H	1179	C
26	1H	1194	A
26	1H	1195	G
26	1H	1204	A
26	1H	1205	U
26	1H	1210	A
26	1H	1211	U
26	1H	1218	C
26	1H	1220	A
26	1H	1225	C
26	1H	1237	A
26	1H	1241	A
26	1H	1244	G
26	1H	1250	G

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Mol	Chain	Res	Type
26	1H	1253	A
26	1H	1256	G
26	1H	1267	U
26	1H	1268	A
26	1H	1271	G
26	1H	1272	A
26	1H	1273	U
26	1H	1275	A
26	1H	1276	A
26	1H	1282	U
26	1H	1300	U
26	1H	1301	A
26	1H	1303	G
26	1H	1306	C
26	1H	1319	G
26	1H	1329	U
26	1H	1332	G
26	1H	1349	A
26	1H	1359	A
26	1H	1360	A
26	1H	1365	A
26	1H	1379	A
26	1H	1380	G
26	1H	1384	A
26	1H	1385	G
26	1H	1386	C
26	1H	1388	G
26	1H	1389	G
26	1H	1395	A
26	1H	1407	C
26	1H	1416	G
26	1H	1417	C
26	1H	1420	U
26	1H	1421	G
26	1H	1428	C
26	1H	1444(A)	A
26	1H	1449	A
26	1H	1449(A)	G
26	1H	1453	A
26	1H	1455	G
26	1H	1458	C
26	1H	1459	G

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Mol	Chain	Res	Type
26	1H	1460	A
26	1H	1461	G
26	1H	1467	C
26	1H	1471	A
26	1H	1483	G
26	1H	1493	C
26	1H	1494	A
26	1H	1495	A
26	1H	1497	U
26	1H	1500	G
26	1H	1505	C
26	1H	1508	A
26	1H	1509	C
26	1H	1510	A
26	1H	1511	A
26	1H	1517	G
26	1H	1522	G
26	1H	1526	G
26	1H	1534	G
26	1H	1535	U
26	1H	1537	C
26	1H	1538	G
26	1H	1539	G
26	1H	1540	G
26	1H	1543	A
26	1H	1545	A
26	1H	1548	C
26	1H	1554	A
26	1H	1558	A
26	1H	1559	G
26	1H	1560	G
26	1H	1566	A
26	1H	1569	A
26	1H	1578	U
26	1H	1580	A
26	1H	1585	C
26	1H	1586	A
26	1H	1587	A
26	1H	1594	G
26	1H	1604	C
26	1H	1608	A
26	1H	1609	A

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Mol	Chain	Res	Type
26	1H	1610	A
26	1H	1617	C
26	1H	1619	G
26	1H	1625	C
26	1H	1634	A
26	1H	1640	C
26	1H	1647	G
26	1H	1648	C
26	1H	1653	G
26	1H	1656	C
26	1H	1660	C
26	1H	1661	G
26	1H	1664	A
26	1H	1674	G
26	1H	1695	G
26	1H	1706	U
26	1H	1728	G
26	1H	1730	U
26	1H	1731	G
26	1H	1733	G
26	1H	1750	G
26	1H	1756	G
26	1H	1758	G
26	1H	1762	A
26	1H	1763	G
26	1H	1764	G
26	1H	1773	A
26	1H	1782	C
26	1H	1783	A
26	1H	1791	A
26	1H	1799	G
26	1H	1800	C
26	1H	1801	G
26	1H	1816	G
26	1H	1828	G
26	1H	1829	A
26	1H	1836	C
26	1H	1847	A
26	1H	1859	A
26	1H	1869	G
26	1H	1870	C
26	1H	1878	G

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Mol	Chain	Res	Type
26	1H	1889	A
26	1H	1895	C
26	1H	1900	A
26	1H	1901	A
26	1H	1906	G
26	1H	1913	A
26	1H	1914	C
26	1H	1919	A
26	1H	1926	U
26	1H	1929	G
26	1H	1930	G
26	1H	1936	A
26	1H	1938	A
26	1H	1945	G
26	1H	1955	U
26	1H	1960	A
26	1H	1963	U
26	1H	1965	C
26	1H	1967	C
26	1H	1969	A
26	1H	1970	A
26	1H	1971	A
26	1H	1972	A
26	1H	1982	C
26	1H	1992	G
26	1H	1993	U
26	1H	1994	C
26	1H	2020	A
26	1H	2021	C
26	1H	2023	G
26	1H	2031	A
26	1H	2033	A
26	1H	2043	C
26	1H	2049	G
26	1H	2055	C
26	1H	2056	G
26	1H	2060	A
26	1H	2061	G
26	1H	2062	A
26	1H	2069	G
26	1H	2072	G
26	1H	2093	G

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Mol	Chain	Res	Type
26	1H	2096	U
26	1H	2099	U
26	1H	2108	C
26	1H	2111	C
26	1H	2113	U
26	1H	2114	A
26	1H	2115	G
26	1H	2116	G
26	1H	2117	A
26	1H	2119	A
26	1H	2123	G
26	1H	2126	A
26	1H	2127	G
26	1H	2128	C
26	1H	2131	G
26	1H	2132	U
26	1H	2133	G
26	1H	2134	A
26	1H	2135	A
26	1H	2136	C
26	1H	2138	C
26	1H	2139	C
26	1H	2145	C
26	1H	2147	G
26	1H	2148	G
26	1H	2156	G
26	1H	2157	G
26	1H	2158	A
26	1H	2161	C
26	1H	2162	G
26	1H	2165	G
26	1H	2166	G
26	1H	2168	G
26	1H	2170	A
26	1H	2171	A
26	1H	2172	U
26	1H	2173	A
26	1H	2175	C
26	1H	2176	A
26	1H	2177	C
26	1H	2181	G
26	1H	2189	U

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Mol	Chain	Res	Type
26	1H	2190	G
26	1H	2198	A
26	1H	2210	G
26	1H	2211	G
26	1H	2212	A
26	1H	2215	G
26	1H	2225	A
26	1H	2226	C
26	1H	2238	G
26	1H	2240	C
26	1H	2263	C
26	1H	2267	A
26	1H	2268	A
26	1H	2273	A
26	1H	2275	C
26	1H	2279	G
26	1H	2280	G
26	1H	2283	C
26	1H	2285	C
26	1H	2286	A
26	1H	2287	A
26	1H	2288	A
26	1H	2294	C
26	1H	2298	A
26	1H	2307	G
26	1H	2308	G
26	1H	2310	A
26	1H	2311	A
26	1H	2312	U
26	1H	2314	C
26	1H	2320	A
26	1H	2321	G
26	1H	2322	A
26	1H	2324	C
26	1H	2325	G
26	1H	2326	C
26	1H	2335	A
26	1H	2336	A
26	1H	2343	C
26	1H	2346	A
26	1H	2347	C
26	1H	2350	C

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Mol	Chain	Res	Type
26	1H	2364	C
26	1H	2376	A
26	1H	2377	A
26	1H	2383	G
26	1H	2385	C
26	1H	2395	C
26	1H	2402	C
26	1H	2403	C
26	1H	2405	G
26	1H	2406	U
26	1H	2408	U
26	1H	2410	G
26	1H	2414	G
26	1H	2422	A
26	1H	2423	U
26	1H	2425	A
26	1H	2428	G
26	1H	2429	G
26	1H	2430	A
26	1H	2431	U
26	1H	2435	A
26	1H	2439	A
26	1H	2440	C
26	1H	2441	C
26	1H	2448	A
26	1H	2468	G
26	1H	2474	C
26	1H	2476	A
26	1H	2477	C
26	1H	2480	C
26	1H	2482	G
26	1H	2484	G
26	1H	2497	A
26	1H	2498	C
26	1H	2502	G
26	1H	2505	G
26	1H	2506	U
26	1H	2507	C
26	1H	2518	A
26	1H	2529	G
26	1H	2549	G
26	1H	2554	U

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Mol	Chain	Res	Type
26	1H	2566	A
26	1H	2567	G
26	1H	2573	C
26	1H	2582	G
26	1H	2594	C
26	1H	2601	C
26	1H	2602	A
26	1H	2609	U
26	1H	2611	U
26	1H	2612	C
26	1H	2629	A
26	1H	2630	G
26	1H	2632	A
26	1H	2634	G
26	1H	2636	U
26	1H	2643	G
26	1H	2654	A
26	1H	2663	G
26	1H	2665	A
26	1H	2666	C
26	1H	2673	G
26	1H	2689	U
26	1H	2702	U
26	1H	2703	C
26	1H	2707	G
26	1H	2712(A)	A
26	1H	2713	A
26	1H	2714	G
26	1H	2726	U
26	1H	2733	A
26	1H	2749	A
26	1H	2752	C
26	1H	2756	U
26	1H	2757	A
26	1H	2764	A
26	1H	2765	A
26	1H	2766	G
26	1H	2777	G
26	1H	2778	A
26	1H	2781	A
26	1H	2789	C
26	1H	2791	C

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Mol	Chain	Res	Type
26	1H	2793	G
26	1H	2795	G
26	1H	2803	C
26	1H	2808	U
26	1H	2820	A
26	1H	2821	A
26	1H	2833	G
26	1H	2834	G
26	1H	2835	A
26	1H	2849	U
26	1H	2872	G
26	1H	2875	C
26	1H	2885	C
26	1H	2891	G
26	1H	2892	A
26	1H	2893	G
26	1H	2894	G
26	1H	2895	U
27	16	0	A
27	16	3	C
27	16	8	U
27	16	12	C
27	16	13	A
27	16	15	A
27	16	16	G
27	16	25	A
27	16	29	A
27	16	39	A
27	16	42	C
27	16	45	A
27	16	56	G
27	16	65	C
27	16	72	G
27	16	73	A
27	16	74	U
27	16	76	G
27	16	81	G
27	16	84	C
27	16	105	G
27	16	107	U
27	16	109	G
27	16	115	G

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Mol	Chain	Res	Type
1	1G	2	U
1	1G	5	U
1	1G	7	G
1	1G	9	G
1	1G	22	G
1	1G	26	A
1	1G	31	G
1	1G	32	A
1	1G	39	G
1	1G	44	G
1	1G	47	C
1	1G	48	C
1	1G	50	A
1	1G	51	A
1	1G	53	A
1	1G	65	U
1	1G	73	G
1	1G	76	G
1	1G	79	G
1	1G	81	G
1	1G	82	U
1	1G	88	C
1	1G	90	C
1	1G	91	C
1	1G	95	G
1	1G	101	A
1	1G	105	G
1	1G	115	G
1	1G	116	A
1	1G	121	C
1	1G	127	G
1	1G	131	C
1	1G	162	A
1	1G	163	C
1	1G	174	C
1	1G	182	U
1	1G	186	C
1	1G	186(F)	C
1	1G	189	U
1	1G	191(D)	U
1	1G	195	A
1	1G	197	A

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Mol	Chain	Res	Type
1	1G	209	U
1	1G	210	U
1	1G	216	G
1	1G	231	G
1	1G	233	C
1	1G	247	G
1	1G	251	G
1	1G	260	G
1	1G	266	G
1	1G	267	C
1	1G	274	A
1	1G	279	A
1	1G	280	C
1	1G	281	G
1	1G	289	G
1	1G	298	A
1	1G	316	G
1	1G	321	A
1	1G	328	C
1	1G	329	A
1	1G	332	G
1	1G	346	G
1	1G	350	G
1	1G	351	G
1	1G	352	C
1	1G	353	A
1	1G	354	G
1	1G	355	C
1	1G	367	U
1	1G	369	C
1	1G	372	C
1	1G	388	G
1	1G	396	G
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

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Mol	Chain	Res	Type
1	1G	423	G
1	1G	424	G
1	1G	429	U
1	1G	430	A
1	1G	439	A
1	1G	442	C
1	1G	452	A
1	1G	465	A
1	1G	466	C
1	1G	467	G
1	1G	475	G
1	1G	478	A
1	1G	482	A
1	1G	484	G
1	1G	485	G
1	1G	486	U
1	1G	496	A
1	1G	497	U
1	1G	498	A
1	1G	500	G
1	1G	505	G
1	1G	509	A
1	1G	510	A
1	1G	511	C
1	1G	517	G
1	1G	518	C
1	1G	521	G
1	1G	527	G
1	1G	531	U
1	1G	532	A
1	1G	533	A
1	1G	544	G
1	1G	547	A
1	1G	553	A
1	1G	559	A
1	1G	561	U
1	1G	562	C
1	1G	564	C
1	1G	572	A
1	1G	573	A
1	1G	576	G
1	1G	586	C

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Mol	Chain	Res	Type
1	1G	607	A
1	1G	614	A
1	1G	615	C
1	1G	617	G
1	1G	620	C
1	1G	621	A
1	1G	630	G
1	1G	631	G
1	1G	632	A
1	1G	633	G
1	1G	651	C
1	1G	653	A
1	1G	661	G
1	1G	665	A
1	1G	666	G
1	1G	687	A
1	1G	688	G
1	1G	702	A
1	1G	723	U
1	1G	724	G
1	1G	731	G
1	1G	749	C
1	1G	755	G
1	1G	769	G
1	1G	770	C
1	1G	776	G
1	1G	777	A
1	1G	778	G
1	1G	787	A
1	1G	792	A
1	1G	793	U
1	1G	794	A
1	1G	809	G
1	1G	816	A
1	1G	817	C
1	1G	819	A
1	1G	820	U
1	1G	821	G
1	1G	827	U
1	1G	828	A
1	1G	842	C
1	1G	843	U

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Mol	Chain	Res	Type
1	1G	848	C
1	1G	859	A
1	1G	870	U
1	1G	873	A
1	1G	874	G
1	1G	885	G
1	1G	914	A
1	1G	915	A
1	1G	916	G
1	1G	926	G
1	1G	927	G
1	1G	934	C
1	1G	935	A
1	1G	936	C
1	1G	944	G
1	1G	953	G
1	1G	960	U
1	1G	961	U
1	1G	968	A
1	1G	969	A
1	1G	971	G
1	1G	972	C
1	1G	974	A
1	1G	975	A
1	1G	976	G
1	1G	977	A
1	1G	978	A
1	1G	979	C
1	1G	980	C
1	1G	982	U
1	1G	983	A
1	1G	989	C
1	1G	991	U
1	1G	992	U
1	1G	993	G
1	1G	995	C
1	1G	996	A
1	1G	1001	G
1	1G	1002	G
1	1G	1004	A
1	1G	1006	C
1	1G	1008	C

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Mol	Chain	Res	Type
1	1G	1009	G
1	1G	1017	G
1	1G	1021	G
1	1G	1023	G
1	1G	1024	G
1	1G	1026	G
1	1G	1028	C
1	1G	1028(A)	C
1	1G	1028(B)	C
1	1G	1029	G
1	1G	1030	C
1	1G	1031	G
1	1G	1032	A
1	1G	1032(A)	G
1	1G	1033	G
1	1G	1037	C
1	1G	1040	U
1	1G	1046	A
1	1G	1050	G
1	1G	1054	C
1	1G	1056	U
1	1G	1064	G
1	1G	1081	G
1	1G	1084	G
1	1G	1088	G
1	1G	1094	G
1	1G	1095	U
1	1G	1101	A
1	1G	1118	C
1	1G	1124	G
1	1G	1125	U
1	1G	1127	G
1	1G	1128	C
1	1G	1129	C
1	1G	1133	G
1	1G	1135	U
1	1G	1136	U
1	1G	1137	C
1	1G	1139	G
1	1G	1140	C
1	1G	1144	G
1	1G	1146	A

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Mol	Chain	Res	Type
1	1G	1147	C
1	1G	1154	G
1	1G	1157	A
1	1G	1158	C
1	1G	1159	U
1	1G	1160	G
1	1G	1169	A
1	1G	1177	G
1	1G	1178	G
1	1G	1181	G
1	1G	1183	A
1	1G	1185	G
1	1G	1188	A
1	1G	1189	C
1	1G	1196	U
1	1G	1197	G
1	1G	1201	A
1	1G	1208	C
1	1G	1211	U
1	1G	1212	U
1	1G	1213	A
1	1G	1225	A
1	1G	1227	A
1	1G	1232	U
1	1G	1238	A
1	1G	1240	U
1	1G	1241	G
1	1G	1248	A
1	1G	1256	A
1	1G	1257	U
1	1G	1258	G
1	1G	1260	C
1	1G	1268	A
1	1G	1273	G
1	1G	1274	G
1	1G	1275	A
1	1G	1278	U
1	1G	1279	A
1	1G	1280	A
1	1G	1286	A
1	1G	1287	A
1	1G	1288	A

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Mol	Chain	Res	Type
1	1G	1297	C
1	1G	1298	C
1	1G	1299	A
1	1G	1301	U
1	1G	1305	G
1	1G	1317	C
1	1G	1320	C
1	1G	1322	C
1	1G	1323	G
1	1G	1324	A
1	1G	1331	G
1	1G	1335	C
1	1G	1336	C
1	1G	1338	G
1	1G	1346	A
1	1G	1347	G
1	1G	1353	G
1	1G	1354	C
1	1G	1358	U
1	1G	1359	C
1	1G	1360	A
1	1G	1362(A)	C
1	1G	1363	A
1	1G	1368	G
1	1G	1370	G
1	1G	1379	G
1	1G	1382	C
1	1G	1387	G
1	1G	1400	C
1	1G	1401	G
1	1G	1406	U
1	1G	1416	G
1	1G	1419	G
1	1G	1442	G
1	1G	1443	G
1	1G	1446	A
1	1G	1447	G
1	1G	1450	U
1	1G	1451	A
1	1G	1452	C
1	1G	1453	G
1	1G	1491	G

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Mol	Chain	Res	Type
1	1G	1492	A
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	1517	G
1	1G	1519	A
1	1G	1520	G
1	1G	1529	G
1	1G	1530	G
1	1G	1532	U
1	1G	1533	C
1	1G	1534	A
24	1L	2	G
24	1L	3	G
24	1L	7	U
24	1L	9	A
24	1L	10	G
24	1L	11	C
24	1L	18	G
24	1L	23	A
24	1L	24	G
24	1L	26	A
24	1L	27	G
24	1L	30	G
24	1L	34	U
24	1L	37	A
24	1L	41	A
24	1L	45	G
24	1L	49	G
24	1L	54	U
24	1L	56	C
24	1L	59	A
24	1L	61	C
24	1L	64	G
24	1L	67	C
24	1L	70	C
23	2L	2	G
23	2L	8	4SU

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Mol	Chain	Res	Type
23	2L	9	G
23	2L	16	C
23	2L	18	C
23	2L	19	G
23	2L	20	G
23	2L	21	U
23	2L	23	G
23	2L	34	U
23	2L	48	U
23	2L	49	C
23	2L	65	G
23	2L	69	C
23	2L	77	A
24	3L	2	G
24	3L	7	U
24	3L	9	A
24	3L	15	G
24	3L	24	G
24	3L	25	C
24	3L	26	A
24	3L	31	A
24	3L	33	U
24	3L	34	U
24	3L	35	U
24	3L	37	A
24	3L	38	A
24	3L	39	U
24	3L	40	C
24	3L	42	A
24	3L	46	G
24	3L	47	U
24	3L	48	C
24	3L	55	U
24	3L	56	C
24	3L	58	A
24	3L	59	A
24	3L	61	C
24	3L	62	C
24	3L	63	U
24	3L	65	C
24	3L	67	C
24	3L	72	C

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Mol	Chain	Res	Type
24	3L	73	A
25	4L	8	A
25	4L	9	G
25	4L	11	U
25	4L	12	A
25	4L	13	A
25	4L	14	A
25	4L	15	A
25	4L	19	G
25	4L	20	A
25	4L	23	A
25	4L	24	A
25	4L	25	A
26	14	7	G
26	14	9	U
26	14	11	G
26	14	15	G
26	14	35	G
26	14	46	C
26	14	49	A
26	14	50	U
26	14	55	G
26	14	58	G
26	14	60	G
26	14	71	A
26	14	72	U
26	14	74	A
26	14	75	G
26	14	78	A
26	14	82	G
26	14	84	A
26	14	92	G
26	14	95	G
26	14	99	U
26	14	101	G
26	14	102	G
26	14	118	A
26	14	119	A
26	14	120	U
26	14	125	G
26	14	129	C
26	14	131	G

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Mol	Chain	Res	Type
26	14	138	G
26	14	139	G
26	14	153	C
26	14	154	G
26	14	155	C
26	14	161	U
26	14	162	U
26	14	171	G
26	14	173	G
26	14	174	C
26	14	181	A
26	14	182	A
26	14	196	A
26	14	199	A
26	14	205	G
26	14	214	G
26	14	215	G
26	14	216	A
26	14	217	G
26	14	222	A
26	14	229	A
26	14	233	A
26	14	248	G
26	14	249	C
26	14	250	G
26	14	252	G
26	14	269	U
26	14	270(K)	C
26	14	270(L)	U
26	14	270(N)	G
26	14	270(O)	U
26	14	270(P)	C
26	14	271(B)	G
26	14	271(C)	U
26	14	271	G
26	14	273(C)	C
26	14	273(D)	C
26	14	273(F)	C
26	14	274	G
26	14	275	G
26	14	276	A
26	14	277	C

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Mol	Chain	Res	Type
26	14	278	A
26	14	279	C
26	14	283	A
26	14	289	A
26	14	290	G
26	14	308	G
26	14	311	A
26	14	312	G
26	14	316	C
26	14	324	A
26	14	329	G
26	14	330	A
26	14	331	A
26	14	352	G
26	14	354	G
26	14	362	U
26	14	363	G
26	14	363(E)	U
26	14	366	C
26	14	372	G
26	14	380	U
26	14	386	G
26	14	391	G
26	14	396	G
26	14	405	U
26	14	406	G
26	14	407	G
26	14	411	G
26	14	428	A
26	14	443	A
26	14	444	C
26	14	447	A
26	14	448	U
26	14	451	C
26	14	454	A
26	14	455	C
26	14	456	C
26	14	457	A
26	14	470	A
26	14	471	A
26	14	481	G
26	14	501	A

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Mol	Chain	Res	Type
26	14	505	A
26	14	509	C
26	14	517	C
26	14	528	A
26	14	531	C
26	14	532	A
26	14	533	G
26	14	537	C
26	14	549	G
26	14	556	G
26	14	563	G
26	14	573	G
26	14	575	A
26	14	587	C
26	14	603	A
26	14	607	U
26	14	617	G
26	14	619	G
26	14	621	A
26	14	622	G
26	14	627	A
26	14	634	C
26	14	637	A
26	14	645	C
26	14	646	A
26	14	650	C
26	14	651	G
26	14	654	A
26	14	654(A)	A
26	14	654(B)	C
26	14	654(C)	G
26	14	654(D)	G
26	14	654(S)	G
26	14	654(T)	A
26	14	677	A
26	14	686	G
26	14	717	G
26	14	722	A
26	14	730	C
26	14	731	C
26	14	734	A
26	14	738	G

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Mol	Chain	Res	Type
26	14	749	C
26	14	750	A
26	14	752	A
26	14	753	C
26	14	764	A
26	14	765	G
26	14	775	G
26	14	776	G
26	14	779	U
26	14	782	A
26	14	784	A
26	14	785	G
26	14	791	C
26	14	792	G
26	14	805	G
26	14	812	C
26	14	814	C
26	14	816	C
26	14	819	A
26	14	827	U
26	14	828	U
26	14	830	G
26	14	832	G
26	14	840	C
26	14	845	G
26	14	846	C
26	14	847	U
26	14	854	G
26	14	859	G
26	14	860	U
26	14	861	A
26	14	865	C
26	14	866	A
26	14	878	A
26	14	880	G
26	14	881	G
26	14	882	G
26	14	883	G
26	14	884	C
26	14	885	C
26	14	886	C
26	14	887	A

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Mol	Chain	Res	Type
26	14	888	C
26	14	889	C
26	14	890	A
26	14	892	G
26	14	894	C
26	14	896	A
26	14	897	C
26	14	899	A
26	14	900	A
26	14	901	A
26	14	903	C
26	14	904	C
26	14	907	U
26	14	910	A
26	14	911	A
26	14	917	A
26	14	918	A
26	14	925	C
26	14	926	A
26	14	932	G
26	14	933	A
26	14	935	C
26	14	938	G
26	14	941	A
26	14	945	A
26	14	946	G
26	14	953	A
26	14	958	U
26	14	959	A
26	14	961	C
26	14	967	C
26	14	974	G
26	14	974(A)	C
26	14	980	A
26	14	983	A
26	14	987	G
26	14	989	G
26	14	990	A
26	14	991	C
26	14	996	A
26	14	1004	C
26	14	1012	U

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Mol	Chain	Res	Type
26	14	1013	C
26	14	1015	G
26	14	1017	G
26	14	1022	G
26	14	1023	U
26	14	1025	G
26	14	1026	U
26	14	1028	A
26	14	1037	G
26	14	1040	C
26	14	1044	G
26	14	1050	A
26	14	1105	U
26	14	1106	G
26	14	1107	G
26	14	1108	U
26	14	1110	G
26	14	1111	A
26	14	1112	G
26	14	1113	U
26	14	1117	G
26	14	1122	G
26	14	1128	A
26	14	1129	A
26	14	1130	U
26	14	1131	G
26	14	1135	C
26	14	1136	G
26	14	1138	G
26	14	1139	G
26	14	1142(A)	A
26	14	1143	A
26	14	1157	G
26	14	1170	G
26	14	1173	G
26	14	1174	A
26	14	1175	U
26	14	1176	G
26	14	1177	A
26	14	1178	C
26	14	1188	U
26	14	1194	A

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Mol	Chain	Res	Type
26	14	1204	A
26	14	1205	U
26	14	1206	G
26	14	1220	A
26	14	1221	C
26	14	1237	A
26	14	1250	G
26	14	1253	A
26	14	1256	G
26	14	1268	A
26	14	1271	G
26	14	1272	A
26	14	1300	U
26	14	1301	A
26	14	1303	G
26	14	1325	G
26	14	1329	U
26	14	1332	G
26	14	1345	C
26	14	1349	A
26	14	1352	U
26	14	1359	A
26	14	1360	A
26	14	1365	A
26	14	1368	G
26	14	1370	C
26	14	1379	A
26	14	1380	G
26	14	1385	G
26	14	1386	C
26	14	1406	U
26	14	1411	C
26	14	1416	G
26	14	1418	G
26	14	1419	A
26	14	1421	G
26	14	1427	A
26	14	1428	C
26	14	1444(A)	A
26	14	1445	C
26	14	1449	A
26	14	1449(A)	G

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Mol	Chain	Res	Type
26	14	1451	C
26	14	1453	A
26	14	1455	G
26	14	1458	C
26	14	1459	G
26	14	1460	A
26	14	1465	G
26	14	1467	C
26	14	1469	A
26	14	1471	A
26	14	1475	G
26	14	1483	G
26	14	1493	C
26	14	1494	A
26	14	1508	A
26	14	1509	C
26	14	1510	A
26	14	1522	G
26	14	1526	G
26	14	1528	A
26	14	1534	G
26	14	1537	C
26	14	1538	G
26	14	1543	A
26	14	1544	C
26	14	1547	C
26	14	1558	A
26	14	1559	G
26	14	1560	G
26	14	1566	A
26	14	1569	A
26	14	1578	U
26	14	1582	C
26	14	1583	A
26	14	1585	C
26	14	1586	A
26	14	1588	C
26	14	1589	C
26	14	1598	C
26	14	1608	A
26	14	1609	A
26	14	1614	A

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Mol	Chain	Res	Type
26	14	1616	A
26	14	1620	G
26	14	1625	C
26	14	1639	U
26	14	1647	G
26	14	1648	C
26	14	1651	G
26	14	1654	A
26	14	1669	A
26	14	1670	C
26	14	1674	G
26	14	1675	C
26	14	1700	A
26	14	1701	A
26	14	1725	G
26	14	1726	G
26	14	1730	U
26	14	1731	G
26	14	1732	A
26	14	1743	G
26	14	1756	G
26	14	1758	G
26	14	1762	A
26	14	1763	G
26	14	1764	G
26	14	1773	A
26	14	1777	U
26	14	1780	A
26	14	1782	C
26	14	1791	A
26	14	1800	C
26	14	1801	G
26	14	1816	G
26	14	1819	A
26	14	1820	U
26	14	1823	G
26	14	1828	G
26	14	1829	A
26	14	1830	C
26	14	1839	G
26	14	1847	A
26	14	1858	G

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Mol	Chain	Res	Type
26	14	1878	G
26	14	1886	C
26	14	1888	G
26	14	1889	A
26	14	1899	G
26	14	1900	A
26	14	1906	G
26	14	1908	C
26	14	1909	C
26	14	1913	A
26	14	1929	G
26	14	1930	G
26	14	1936	A
26	14	1937	A
26	14	1938	A
26	14	1955	U
26	14	1963	U
26	14	1964	G
26	14	1967	C
26	14	1969	A
26	14	1970	A
26	14	1971	A
26	14	1972	A
26	14	1993	U
26	14	2020	A
26	14	2023	G
26	14	2031	A
26	14	2032	G
26	14	2033	A
26	14	2036	C
26	14	2043	C
26	14	2049	G
26	14	2055	C
26	14	2056	G
26	14	2060	A
26	14	2061	G
26	14	2062	A
26	14	2063	C
26	14	2069	G
26	14	2071	A
26	14	2093	G
26	14	2099	U

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Mol	Chain	Res	Type
26	14	2100	G
26	14	2102	U
26	14	2108	C
26	14	2109	U
26	14	2114	A
26	14	2115	G
26	14	2117	A
26	14	2118	U
26	14	2119	A
26	14	2120	G
26	14	2122	U
26	14	2124	G
26	14	2125	G
26	14	2127	G
26	14	2128	C
26	14	2129	C
26	14	2131	G
26	14	2132	U
26	14	2133	G
26	14	2134	A
26	14	2135	A
26	14	2136	C
26	14	2137	C
26	14	2139	C
26	14	2140	C
26	14	2144	U
26	14	2145	C
26	14	2146	C
26	14	2147	G
26	14	2148	G
26	14	2153	G
26	14	2157	G
26	14	2158	A
26	14	2161	C
26	14	2162	G
26	14	2164	C
26	14	2165	G
26	14	2166	G
26	14	2167	U
26	14	2168	G
26	14	2171	A
26	14	2172	U

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Mol	Chain	Res	Type
26	14	2173	A
26	14	2174	C
26	14	2175	C
26	14	2189	U
26	14	2190	G
26	14	2191	G
26	14	2192	G
26	14	2198	A
26	14	2207	C
26	14	2210	G
26	14	2211	G
26	14	2212	A
26	14	2213	U
26	14	2215	G
26	14	2225	A
26	14	2226	C
26	14	2238	G
26	14	2239	G
26	14	2240	C
26	14	2249	U
26	14	2251	G
26	14	2253	G
26	14	2268	A
26	14	2269	A
26	14	2273	A
26	14	2275	C
26	14	2276	G
26	14	2278	A
26	14	2280	G
26	14	2281	C
26	14	2283	C
26	14	2287	A
26	14	2288	A
26	14	2297	C
26	14	2305	A
26	14	2307	G
26	14	2309	A
26	14	2310	A
26	14	2311	A
26	14	2318	G
26	14	2321	G
26	14	2325	G

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Mol	Chain	Res	Type
26	14	2333	A
26	14	2334	G
26	14	2336	A
26	14	2342	C
26	14	2346	A
26	14	2347	C
26	14	2350	C
26	14	2360	A
26	14	2372	G
26	14	2383	G
26	14	2385	C
26	14	2391	G
26	14	2392	A
26	14	2395	C
26	14	2396	G
26	14	2402	C
26	14	2403	C
26	14	2406	U
26	14	2410	G
26	14	2414	G
26	14	2422	A
26	14	2423	U
26	14	2425	A
26	14	2429	G
26	14	2430	A
26	14	2431	U
26	14	2432	A
26	14	2434	A
26	14	2435	A
26	14	2439	A
26	14	2440	C
26	14	2441	C
26	14	2448	A
26	14	2468	G
26	14	2469	A
26	14	2470	G
26	14	2475	C
26	14	2476	A
26	14	2477	C
26	14	2478	A
26	14	2482	G
26	14	2487	G

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Mol	Chain	Res	Type
26	14	2492	U
26	14	2496	C
26	14	2497	A
26	14	2498	C
26	14	2501	C
26	14	2502	G
26	14	2504	U
26	14	2505	G
26	14	2507	C
26	14	2518	A
26	14	2529	G
26	14	2542	A
26	14	2543	G
26	14	2549	G
26	14	2554	U
26	14	2563	U
26	14	2564	A
26	14	2566	A
26	14	2567	G
26	14	2569	G
26	14	2573	C
26	14	2579	C
26	14	2581	G
26	14	2582	G
26	14	2585	U
26	14	2601	C
26	14	2602	A
26	14	2609	U
26	14	2610	C
26	14	2611	U
26	14	2612	C
26	14	2613	U
26	14	2617	C
26	14	2630	G
26	14	2631	G
26	14	2636	U
26	14	2648	C
26	14	2654	A
26	14	2665	A
26	14	2667	C
26	14	2673	G
26	14	2679	A

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Mol	Chain	Res	Type
26	14	2689	U
26	14	2690	C
26	14	2691	C
26	14	2702	U
26	14	2703	C
26	14	2707	G
26	14	2712(A)	A
26	14	2713	A
26	14	2714	G
26	14	2726	U
26	14	2733	A
26	14	2744	G
26	14	2748	A
26	14	2750	A
26	14	2751	G
26	14	2752	C
26	14	2757	A
26	14	2758	A
26	14	2762	G
26	14	2764	A
26	14	2765	A
26	14	2766	G
26	14	2769	C
26	14	2777	G
26	14	2778	A
26	14	2787	C
26	14	2790	A
26	14	2791	C
26	14	2792	G
26	14	2793	G
26	14	2795	G
26	14	2797	U
26	14	2798	C
26	14	2799	A
26	14	2801	A
26	14	2802	G
26	14	2805	G
26	14	2808	U
26	14	2810	A
26	14	2818	G
26	14	2820	A
26	14	2821	A

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Mol	Chain	Res	Type
26	14	2833	G
26	14	2834	G
26	14	2835	A
26	14	2839	G
26	14	2849	U
26	14	2860	A
26	14	2872	G
26	14	2873	A
26	14	2874	C
26	14	2880	C
26	14	2885	C
26	14	2886	G
26	14	2892	A
26	14	2893	G
26	14	2894	G
26	14	2895	U
27	1J	0	A
27	1J	7	G
27	1J	8	U
27	1J	12	C
27	1J	13	A
27	1J	15	A
27	1J	16	G
27	1J	22	U
27	1J	25	A
27	1J	26	A
27	1J	28	C
27	1J	29	A
27	1J	33	G
27	1J	40	U
27	1J	42	C
27	1J	44	G
27	1J	45	A
27	1J	51	G
27	1J	53	A
27	1J	58	A
27	1J	59	A
27	1J	73	A
27	1J	88	C
27	1J	89	G
27	1J	89(A)	A
27	1J	90	C

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Mol	Chain	Res	Type
27	1J	99	A
27	1J	100	G
27	1J	102	G
27	1J	108	C
27	1J	109	G
27	1J	115	G

All (180) RNA pucker outliers are listed below:

Mol	Chain	Res	Type
1	13	1	U
1	13	5	U
1	13	49	U
1	13	50	A
1	13	115	G
1	13	190	G
1	13	244	U
1	13	266	G
1	13	353	A
1	13	422	C
1	13	428	G
1	13	429	U
1	13	484	G
1	13	509	A
1	13	560	U
1	13	687	A
1	13	748	C
1	13	793	U
1	13	871	U
1	13	913	A
1	13	1025	U
1	13	1064	G
1	13	1065	U
1	13	1129	C
1	13	1256	A
1	13	1285	A
1	13	1301	U
1	13	1336	C
1	13	1397	C
1	13	1443	G
1	13	1498	U
1	13	1503	A

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Mol	Chain	Res	Type
1	13	1533	C
22	1K	6	G
22	1K	21	A
22	1K	49	G
22	1K	69	A
23	2K	47	7MG
23	2K	48	U
24	3K	2	G
24	3K	34	U
25	4K	24	A
26	1H	125	G
26	1H	195	A
26	1H	196	A
26	1H	199	A
26	1H	222	A
26	1H	249	C
26	1H	404	C
26	1H	508	G
26	1H	587	C
26	1H	685	A
26	1H	746	A
26	1H	764	A
26	1H	776	G
26	1H	827	U
26	1H	845	G
26	1H	859	G
26	1H	961	C
26	1H	1022	G
26	1H	1026	U
26	1H	1081	U
26	1H	1110	G
26	1H	1178	C
26	1H	1210	A
26	1H	1275	A
26	1H	1378	A
26	1H	1379	A
26	1H	1394	U
26	1H	1396	U
26	1H	1420	U
26	1H	1451	C
26	1H	1508	A
26	1H	1509	C

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Mol	Chain	Res	Type
26	1H	1558	A
26	1H	1608	A
26	1H	1609	A
26	1H	1617	C
26	1H	1757	U
26	1H	1782	C
26	1H	1799	G
26	1H	1858	G
26	1H	1900	A
26	1H	1992	G
26	1H	2060	A
26	1H	2171	A
26	1H	2172	U
26	1H	2210	G
26	1H	2211	G
26	1H	2422	A
26	1H	2439	A
26	1H	2481	G
26	1H	2566	A
26	1H	2756	U
27	16	44	G
1	1G	64	G
1	1G	80	G
1	1G	87	A
1	1G	89	U
1	1G	115	G
1	1G	250	A
1	1G	266	G
1	1G	327	A
1	1G	345	C
1	1G	412	A
1	1G	429	U
1	1G	466	C
1	1G	485	G
1	1G	509	A
1	1G	560	U
1	1G	687	A
1	1G	748	C
1	1G	793	U
1	1G	884	U
1	1G	913	A
1	1G	992	U

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Mol	Chain	Res	Type
1	1G	1053	G
1	1G	1064	G
1	1G	1126	U
1	1G	1145	C
1	1G	1157	A
1	1G	1285	A
1	1G	1298	C
1	1G	1300	G
1	1G	1442	G
1	1G	1449	C
1	1G	1498	U
1	1G	1533	C
24	1L	6	G
24	1L	69	A
23	2L	33	OMC
23	2L	48	U
24	3L	36	U
24	3L	58	A
25	4L	23	A
26	14	6	A
26	14	34	C
26	14	49	A
26	14	71	A
26	14	128	C
26	14	270(M)	U
26	14	278	A
26	14	310	A
26	14	503	A
26	14	685	A
26	14	752	A
26	14	764	A
26	14	774	A
26	14	877	U
26	14	893	C
26	14	960	A
26	14	1011	G
26	14	1022	G
26	14	1325	G
26	14	1379	A
26	14	1396	U
26	14	1420	U
26	14	1558	A

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Mol	Chain	Res	Type
26	14	1608	A
26	14	1762	A
26	14	1819	A
26	14	1992	G
26	14	2035	G
26	14	2062	A
26	14	2212	A
26	14	2275	C
26	14	2308	G
26	14	2406	U
26	14	2425	A
26	14	2439	A
26	14	2477	C
26	14	2629	A
26	14	2689	U
26	14	2756	U
26	14	2776	A
26	14	2791	C
26	14	2859	G
27	1J	52	A
27	1J	88	C
27	1J	89	G

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

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

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

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\#\|Z\| > 2$	Counts	RMSZ	$\#\|Z\| > 2$
22	U8U	1K	34	25,22	15,24,25	2.57	4 (26%)	17,34,37	1.85	2 (11%)
22	T6A	1K	37	22	23,34,35	2.47	4 (17%)	26,49,52	3.42	7 (26%)
22	PSU	1K	39	22	15,21,22	0.96	1 (6%)	16,30,33	2.02	4 (25%)
22	5MU	1K	54	22	13,22,23	1.73	2 (15%)	16,32,35	1.78	2 (12%)
22	PSU	1K	55	22	15,21,22	1.13	1 (6%)	16,30,33	2.29	3 (18%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\#\|Z\|>2$	Counts	RMSZ	$\#\|Z\|>2$
23	OMC	2K	33	23	15,22,23	2.22	4 (26%)	20,31,34	1.82	3 (15%)
23	7MG	2K	47	23	20,26,27	3.34	6 (30%)	23,39,42	2.29	5 (21%)
23	5MU	2K	55	23	13,22,23	1.69	2 (15%)	16,32,35	1.53	2 (12%)
23	PSU	2K	56	23	15,21,22	1.04	1 (6%)	16,30,33	1.87	3 (18%)
23	4SU	2K	8	23	12,21,22	3.01	2 (16%)	15,30,33	0.76	1 (6%)
23	OMC	2L	33	23	15,22,23	2.29	4 (26%)	20,31,34	1.11	2 (10%)
23	7MG	2L	47	23	20,26,27	3.42	5 (25%)	23,39,42	2.27	6 (26%)
23	5MU	2L	55	23	13,22,23	1.71	2 (15%)	16,32,35	1.50	1 (6%)
23	PSU	2L	56	23	15,21,22	1.10	1 (6%)	16,30,33	1.97	3 (18%)
23	4SU	2L	8	23	12,21,22	3.40	2 (16%)	15,30,33	0.92	1 (6%)

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

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
22	U8U	1K	34	25,22	-	0/5/28/29	0/2/2/2
22	T6A	1K	37	22	-	0/15/41/42	0/3/3/3
22	PSU	1K	39	22	-	0/7/25/26	0/2/2/2
22	5MU	1K	54	22	-	0/3/25/26	0/2/2/2
22	PSU	1K	55	22	-	0/7/25/26	0/2/2/2
23	OMC	2K	33	23	-	0/5/27/28	0/2/2/2
23	7MG	2K	47	23	-	0/7/37/38	0/3/3/3
23	5MU	2K	55	23	-	0/3/25/26	0/2/2/2
23	PSU	2K	56	23	-	0/7/25/26	0/2/2/2
23	4SU	2K	8	23	-	0/3/25/26	0/2/2/2
23	OMC	2L	33	23	-	0/5/27/28	0/2/2/2
23	7MG	2L	47	23	-	0/7/37/38	0/3/3/3
23	5MU	2L	55	23	-	0/3/25/26	0/2/2/2
23	PSU	2L	56	23	-	0/7/25/26	0/2/2/2
23	4SU	2L	8	23	-	0/3/25/26	0/2/2/2

All (41) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
23	2K	47	7MG	C5-C4	-5.57	1.24	1.39
23	2L	47	7MG	C5-C4	-5.28	1.25	1.39
22	1K	34	U8U	C2-S2	-3.90	1.58	1.66
23	2K	55	5MU	C4-N3	-2.85	1.27	1.33

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
22	1K	37	T6A	C5-C4	-2.82	1.34	1.40
23	2L	55	5MU	C4-N3	-2.80	1.28	1.33
22	1K	54	5MU	C4-N3	-2.65	1.28	1.33
22	1K	34	U8U	O4-C4	-2.01	1.19	1.24
23	2K	47	7MG	C2-N1	2.08	1.39	1.35
23	2K	56	PSU	C4-N3	2.53	1.37	1.33
23	2L	33	OMC	C4-N4	2.88	1.43	1.35
23	2K	33	OMC	C4-N4	3.05	1.43	1.35
22	1K	39	PSU	C4-N3	3.15	1.38	1.33
23	2L	56	PSU	C4-N3	3.55	1.39	1.33
22	1K	55	PSU	C4-N3	3.56	1.39	1.33
23	2L	47	7MG	C2-N2	3.56	1.41	1.34
23	2K	47	7MG	C2-N2	3.65	1.41	1.34
23	2L	33	OMC	C2-N3	3.70	1.45	1.38
23	2K	33	OMC	C2-N3	3.87	1.46	1.38
23	2L	33	OMC	C5-C4	4.04	1.50	1.41
23	2K	33	OMC	C5-C4	4.40	1.51	1.41
23	2L	47	7MG	C8-N7	4.51	1.64	1.43
23	2K	47	7MG	C8-N7	4.59	1.64	1.43
22	1K	37	T6A	C10-N11	4.93	1.49	1.35
23	2K	55	5MU	C2-N3	5.03	1.48	1.38
23	2K	33	OMC	C6-N1	5.06	1.42	1.35
23	2K	47	7MG	C6-C5	5.18	1.48	1.41
23	2L	55	5MU	C2-N3	5.23	1.49	1.38
22	1K	54	5MU	C2-N3	5.31	1.49	1.38
22	1K	34	U8U	C6-C5	5.40	1.49	1.36
23	2L	47	7MG	C6-C5	5.50	1.49	1.41
23	2L	33	OMC	C6-N1	5.78	1.43	1.35
23	2K	8	4SU	C6-N1	6.44	1.44	1.35
22	1K	37	T6A	C10-N6	6.48	1.49	1.37
22	1K	37	T6A	C6-N6	6.92	1.49	1.36
22	1K	34	U8U	C4-N3	7.02	1.45	1.33
23	2L	8	4SU	C6-N1	7.28	1.45	1.35
23	2K	8	4SU	C5-C4	8.00	1.48	1.38
23	2L	8	4SU	C5-C4	9.04	1.50	1.38
23	2K	47	7MG	C4-N3	10.88	1.48	1.34
23	2L	47	7MG	C4-N3	11.48	1.49	1.34

All (45) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
22	1K	37	T6A	N3-C2-N1	-11.74	119.65	128.87

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
23	2K	47	7MG	C5-C4-N3	-7.75	118.84	126.74
23	2L	47	7MG	C5-C4-N3	-7.35	119.25	126.74
22	1K	34	U8U	C5-C4-N3	-6.24	119.59	125.19
22	1K	37	T6A	C13-C12-N11	-5.21	103.39	113.40
22	1K	37	T6A	C6-N6-C10	-4.91	123.70	130.33
22	1K	55	PSU	C5-C1'-C2'	-3.97	108.69	115.44
22	1K	37	T6A	C1'-N9-C4	-3.37	123.04	126.81
23	2L	47	7MG	N1-C2-N3	-3.22	120.25	125.51
23	2L	56	PSU	C5-C6-N1	-2.89	120.35	124.38
23	2L	8	4SU	C5-C4-N3	-2.84	120.55	123.56
23	2K	47	7MG	N1-C2-N3	-2.82	120.90	125.51
22	1K	39	PSU	C5-C1'-C2'	-2.79	110.69	115.44
23	2K	47	7MG	C5-C6-N1	-2.54	119.61	123.39
23	2L	47	7MG	C5-C6-N1	-2.50	119.67	123.39
22	1K	54	5MU	C5-C4-N3	-2.38	123.34	125.35
22	1K	37	T6A	O10-C10-N6	-2.26	120.13	123.59
23	2K	33	OMC	C5-C4-N4	-2.25	117.58	121.19
23	2K	56	PSU	C5-C6-N1	-2.25	121.25	124.38
22	1K	39	PSU	C5-C6-N1	-2.24	121.25	124.38
23	2K	8	4SU	C5-C4-N3	-2.14	121.29	123.56
23	2K	55	5MU	C5-C4-N3	-2.10	123.58	125.35
23	2L	47	7MG	C2-N3-C4	2.02	120.26	114.50
22	1K	55	PSU	O4'-C1'-C2'	2.32	107.20	104.69
23	2K	56	PSU	O4'-C1'-C2'	2.36	107.24	104.69
23	2L	33	OMC	C6-C5-C4	2.38	118.37	117.44
22	1K	39	PSU	O4'-C1'-C2'	2.41	107.30	104.69
23	2L	33	OMC	N4-C4-N3	2.42	120.72	116.50
23	2L	56	PSU	O4'-C1'-C2'	2.43	107.31	104.69
23	2K	47	7MG	C6-N1-C2	2.68	119.02	115.88
23	2K	33	OMC	N4-C4-N3	3.18	122.06	116.50
22	1K	34	U8U	C2-N3-C4	3.29	119.55	115.89
23	2L	47	7MG	C6-N1-C2	3.39	119.85	115.88
23	2L	47	7MG	C5-C4-N9	4.61	113.69	106.25
23	2K	47	7MG	C5-C4-N9	5.05	114.40	106.25
23	2L	55	5MU	C4-N3-C2	5.24	119.53	115.16
23	2K	55	5MU	C4-N3-C2	5.26	119.55	115.16
23	2K	56	PSU	C4-N3-C2	5.97	120.14	115.16
23	2L	56	PSU	C4-N3-C2	6.18	120.32	115.16
22	1K	54	5MU	C4-N3-C2	6.33	120.44	115.16
22	1K	39	PSU	C4-N3-C2	6.45	120.54	115.16
22	1K	37	T6A	C2-N1-C6	6.68	121.27	116.47
22	1K	37	T6A	C12-N11-C10	6.72	134.61	120.82

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Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
23	2K	33	OMC	C6-C5-C4	6.81	120.11	117.44
22	1K	55	PSU	C4-N3-C2	7.30	121.25	115.16

There are no chirality outliers.

There are no torsion outliers.

There are no ring outliers.

7 monomers are involved in 13 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
22	1K	37	T6A	1	0
22	1K	54	5MU	1	0
23	2K	33	OMC	1	0
23	2K	47	7MG	2	0
23	2K	55	5MU	3	0
23	2L	33	OMC	3	0
23	2L	55	5MU	2	0

5.5 Carbohydrates (i)

There are no carbohydrates in this entry.

5.6 Ligand geometry (i)

Of 1269 ligands modelled in this entry, 1262 are monoatomic - leaving 7 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	$\#\|Z\| > 2$
58	PAR	13	1741	1	45,45,45	0.67	0	60,67,67	1.42	9 (15%)
61	SPE	14	3436	-	12,12,12	0.44	0	11,11,11	0.64	0
61	SPE	14	3437	26	12,12,12	0.82	0	11,11,11	1.16	1 (9%)
58	PAR	1G	1702	-	45,45,45	0.66	1 (2%)	60,67,67	1.36	7 (11%)
61	SPE	1G	1703	-	12,12,12	0.37	0	11,11,11	0.73	0

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\#\|Z\|>2$	Counts	RMSZ	$\#\|Z\|>2$
59	SF4	32	302	4	0,12,12	0.00	-	0,24,24	0.00	-
59	SF4	3E	301	4	0,12,12	0.00	-	0,24,24	0.00	-

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

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
58	PAR	13	1741	1	-	0/18/94/94	0/4/4/4
61	SPE	14	3436	-	-	0/10/10/10	0/0/0/0
61	SPE	14	3437	26	-	0/10/10/10	0/0/0/0
58	PAR	1G	1702	-	-	0/18/94/94	0/4/4/4
61	SPE	1G	1703	-	-	0/10/10/10	0/0/0/0
59	SF4	32	302	4	-	0/0/48/48	0/6/5/5
59	SF4	3E	301	4	-	0/0/48/48	0/6/5/5

All (1) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
58	1G	1702	PAR	C24-N24	-2.19	1.43	1.47

All (17) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(^{o})	Ideal(^{o})
58	1G	1702	PAR	C14-O33-C33	-3.33	109.14	118.00
58	1G	1702	PAR	C13-O52-C52	-3.07	109.84	118.00
58	13	1741	PAR	O33-C14-O54	-3.01	102.86	110.69
61	14	3437	SPE	C6-N5-C4	-2.79	103.68	113.35
58	13	1741	PAR	C14-O33-C33	-2.70	110.83	118.00
58	13	1741	PAR	O34-C34-C24	-2.69	105.46	110.28
58	1G	1702	PAR	O54-C14-C24	-2.58	103.99	109.88
58	1G	1702	PAR	C62-C12-N12	-2.15	106.92	110.66
58	13	1741	PAR	C22-C32-C42	2.14	114.65	109.31
58	1G	1702	PAR	C52-C62-C12	2.16	115.20	109.99
58	13	1741	PAR	O51-C51-C41	2.17	113.81	109.67
58	1G	1702	PAR	O51-C11-C21	2.50	115.61	109.88
58	13	1741	PAR	O51-C11-C21	2.85	116.39	109.88
58	13	1741	PAR	C52-C42-C32	2.95	117.01	111.09
58	1G	1702	PAR	C11-O51-C51	3.08	119.79	113.74
58	13	1741	PAR	C22-C12-C62	3.13	115.00	110.14
58	13	1741	PAR	C11-O51-C51	3.62	120.85	113.74

There are no chirality outliers.

There are no torsion outliers.

There are no ring outliers.

5 monomers are involved in 13 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
61	14	3436	SPE	2	0
61	14	3437	SPE	6	0
58	1G	1702	PAR	1	0
59	32	302	SF4	3	0
59	3E	301	SF4	1	0

5.7 Other polymers

There are no such residues in this entry.

5.8 Polymer linkage issues

The following chains have linkage breaks:

Mol	Chain	Number of breaks
10	1A	2
26	1H	1
4	3E	1
47	D5	1
24	1L	1

All chain breaks are listed below:

Model	Chain	Residue-1	Atom-1	Residue-2	Atom-2	Distance (Å)
1	1H	1055:G	O3'	1059:G	P	14.13
1	1L	72:C	O3'	73:A	P	3.25
1	1A	38:ILE	C	39:PRO	N	1.71
1	1A	76:ASN	C	77:PRO	N	1.61
1	D5	94:GLU	C	95:PRO	N	1.61
1	3E	36:ARG	C	37:PRO	N	1.19

6 Fit of model and data

6.1 Protein, DNA and RNA chains

In the following table, the column labelled '#RSRZ> 2' contains the number (and percentage) of RSRZ outliers, followed by percent RSRZ outliers for the chain as percentile scores relative to all X-ray entries and entries of similar resolution. The OWAB column contains the minimum, median, 95^{th} 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	1499/1522 (98%)	-0.23	1 (0%) 95 90	57, 100, 176, 247	0
1	1G	1508/1522 (99%)	-0.17	10 (0%) 89 70	71, 119, 192, 267	0
2	12	207/256 (80%)	0.76	26 (12%) 5 2	131, 164, 184, 192	0
2	1E	231/256 (90%)	0.24	12 (5%) 31 12	112, 143, 169, 174	0
3	22	195/239 (81%)	1.21	50 (25%) 1 1	135, 160, 173, 178	0
3	2E	205/239 (85%)	0.67	22 (10%) 8 3	88, 111, 143, 151	0
4	32	208/209 (99%)	0.65	14 (6%) 21 7	92, 113, 136, 142	0
4	3E	207/209 (99%)	0.31	12 (5%) 26 10	84, 108, 132, 144	0
5	42	150/162 (92%)	0.49	11 (7%) 18 6	108, 128, 148, 160	0
5	4E	149/162 (91%)	0.30	4 (2%) 58 28	82, 103, 121, 131	0
6	52	101/101 (100%)	0.02	0 100 100	85, 104, 121, 132	0
6	5E	100/101 (99%)	0.52	7 (7%) 19 7	85, 105, 120, 132	0
7	62	138/156 (88%)	1.18	33 (23%) 1 1	116, 130, 141, 147	0
7	6E	154/156 (98%)	1.23	36 (23%) 1 1	102, 120, 150, 171	0
8	72	137/138 (99%)	0.73	15 (10%) 7 3	100, 131, 142, 149	0
8	7E	138/138 (100%)	1.07	29 (21%) 1 1	93, 110, 122, 134	0
9	82	121/128 (94%)	1.87	50 (41%) 0 0	116, 163, 174, 182	0
9	8E	126/128 (98%)	0.24	5 (3%) 42 17	87, 138, 157, 163	0
10	1A	80/105 (76%)	1.02	22 (27%) 1 0	133, 156, 169, 174	0
10	1I	94/105 (89%)	1.42	28 (29%) 1 0	81, 131, 168, 173	0
11	2A	113/129 (87%)	1.33	27 (23%) 1 1	84, 109, 124, 135	0
11	2I	111/129 (86%)	1.14	25 (22%) 1 1	72, 107, 123, 133	0
12	3A	121/132 (91%)	1.72	45 (37%) 0 0	90, 112, 135, 151	0
12	3I	122/132 (92%)	0.72	15 (12%) 5 2	66, 76, 102, 135	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å^2)	Q<0.9
13	4A	109/126 (86%)	0.85	26 (23%) 1 1	123, 150, 180, 194	0
13	4I	117/126 (92%)	0.31	4 (3%) 49 21	88, 116, 131, 138	0
14	5A	59/61 (96%)	3.89	41 (69%) 0 0	137, 158, 180, 183	0
14	5I	59/61 (96%)	0.83	8 (13%) 4 1	81, 96, 114, 122	0
15	6A	87/89 (97%)	0.13	3 (3%) 49 21	84, 113, 128, 131	0
15	6I	87/89 (97%)	0.35	3 (3%) 49 21	80, 98, 113, 123	0
16	7A	84/88 (95%)	0.26	2 (2%) 62 32	92, 105, 129, 155	0
16	7I	83/88 (94%)	1.13	20 (24%) 1 1	96, 111, 137, 155	0
17	8A	99/105 (94%)	1.30	27 (27%) 1 0	97, 113, 130, 135	0
17	8I	99/105 (94%)	1.10	18 (18%) 2 1	89, 106, 115, 124	0
18	9A	67/88 (76%)	0.35	4 (5%) 25 9	92, 112, 132, 138	0
18	9I	68/88 (77%)	0.20	2 (2%) 55 26	88, 107, 131, 135	0
19	AA	65/93 (69%)	1.17	13 (20%) 1 1	162, 178, 187, 192	0
19	AI	82/93 (88%)	0.24	3 (3%) 45 19	96, 117, 138, 151	0
20	BA	99/106 (93%)	1.15	18 (18%) 2 1	90, 114, 140, 152	0
20	BI	97/106 (91%)	1.36	27 (27%) 1 0	106, 119, 145, 151	0
21	1B	22/27 (81%)	1.72	8 (36%) 0 0	127, 141, 145, 148	0
21	1F	23/27 (85%)	0.24	0 100 100	92, 99, 107, 115	0
22	1K	67/76 (88%)	0.54	5 (7%) 17 6	90, 193, 226, 233	0
23	2K	72/77 (93%)	0.03	1 (1%) 78 51	67, 92, 124, 140	0
23	2L	72/77 (93%)	0.07	0 100 100	80, 116, 152, 163	0
24	1L	66/76 (86%)	1.06	11 (16%) 2 1	145, 224, 245, 249	0
24	3K	70/76 (92%)	0.91	10 (14%) 4 1	76, 229, 251, 253	0
24	3L	72/76 (94%)	0.16	3 (4%) 40 16	85, 220, 238, 240	0
25	4K	19/30 (63%)	0.97	4 (21%) 1 1	71, 140, 219, 219	0
25	4L	19/30 (63%)	1.35	3 (15%) 3 1	98, 158, 230, 230	0
26	14	2826/2917 (96%)	-0.06	17 (0%) 90 73	50, 85, 199, 264	0
26	1H	2860/2917 (98%)	-0.10	7 (0%) 95 87	40, 69, 186, 257	0
27	16	122/122 (100%)	-0.35	1 (0%) 87 67	61, 88, 107, 197	0
27	1J	122/122 (100%)	-0.45	0 100 100	85, 119, 145, 205	0
28	71	132/229 (57%)	0.80	20 (15%) 3 1	146, 210, 235, 243	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å^2)	Q<0.9
29	11	273/276 (98%)	0.46	5 (1%) 71 43	41, 63, 79, 93	0
29	19	274/276 (99%)	0.59	7 (2%) 59 29	47, 73, 88, 107	0
30	21	203/206 (98%)	0.63	10 (4%) 33 13	48, 84, 120, 130	0
30	29	204/206 (99%)	1.03	39 (19%) 2 1	60, 95, 134, 149	0
31	31	202/210 (96%)	0.62	9 (4%) 37 15	46, 76, 112, 127	0
31	39	204/210 (97%)	0.56	14 (6%) 20 7	57, 101, 148, 177	0
32	41	179/182 (98%)	0.77	21 (11%) 6 2	79, 100, 134, 146	0
32	49	180/182 (98%)	1.22	48 (26%) 1 0	117, 138, 156, 174	0
33	51	174/180 (96%)	0.30	5 (2%) 55 26	77, 102, 119, 130	0
33	59	169/180 (93%)	2.48	92 (54%) 0 0	152, 196, 220, 233	0
34	61	145/148 (97%)	0.52	17 (11%) 6 2	76, 131, 149, 157	0
34	69	145/148 (97%)	0.46	12 (8%) 14 5	86, 122, 149, 156	0
35	15	137/140 (97%)	1.16	25 (18%) 2 1	81, 108, 135, 151	0
35	58	137/140 (97%)	0.86	18 (13%) 5 2	63, 85, 121, 138	0
36	25	122/122 (100%)	0.91	14 (11%) 6 2	69, 88, 105, 117	0
36	68	122/122 (100%)	0.39	3 (2%) 61 30	56, 72, 90, 99	0
37	35	147/150 (98%)	0.86	21 (14%) 4 1	58, 102, 135, 152	0
37	78	147/150 (98%)	0.57	10 (6%) 20 7	47, 80, 104, 115	0
38	45	138/141 (97%)	0.95	27 (19%) 1 1	76, 105, 124, 155	0
38	88	141/141 (100%)	0.71	10 (7%) 19 7	54, 75, 95, 123	0
39	55	118/118 (100%)	0.59	6 (5%) 32 12	65, 80, 97, 112	0
39	98	118/118 (100%)	0.81	12 (10%) 9 3	60, 78, 94, 106	0
40	65	110/112 (98%)	1.06	22 (20%) 1 1	89, 112, 132, 137	0
40	A8	111/112 (99%)	1.09	18 (16%) 3 1	72, 85, 105, 116	0
41	75	133/146 (91%)	0.40	7 (5%) 30 12	80, 96, 128, 151	0
41	B8	135/146 (92%)	0.14	5 (3%) 45 19	67, 84, 128, 139	0
42	85	116/118 (98%)	0.54	4 (3%) 49 21	63, 96, 127, 134	0
42	C8	115/118 (97%)	0.31	3 (2%) 59 29	53, 77, 102, 108	0
43	95	100/101 (99%)	1.08	22 (22%) 1 1	65, 116, 135, 142	0
43	D8	100/101 (99%)	0.89	15 (15%) 3 1	53, 101, 119, 126	0
44	A5	111/113 (98%)	0.79	5 (4%) 37 15	57, 75, 101, 137	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å^2)	Q<0.9
44	E8	110/113 (97%)	0.49	8 (7%) 18 6	53, 70, 94, 106	0
45	B5	94/96 (97%)	1.09	15 (15%) 3 1	68, 83, 104, 118	0
45	F8	95/96 (98%)	0.21	1 (1%) 82 58	49, 66, 97, 110	0
46	C5	104/110 (94%)	2.12	36 (34%) 0 0	92, 118, 156, 164	0
46	G8	103/110 (93%)	0.26	5 (4%) 33 13	73, 95, 122, 130	0
47	D5	177/206 (85%)	1.81	70 (39%) 0 0	108, 148, 226, 236	0
47	H8	170/206 (82%)	1.81	65 (38%) 0 0	80, 116, 202, 210	0
48	E5	76/85 (89%)	1.15	13 (17%) 2 1	66, 88, 102, 115	0
48	I8	77/85 (90%)	0.58	5 (6%) 22 8	54, 69, 91, 102	0
49	F5	94/98 (95%)	1.20	18 (19%) 2 1	60, 83, 122, 136	0
49	J8	96/98 (97%)	0.85	9 (9%) 11 4	51, 71, 126, 138	0
50	G5	69/72 (95%)	0.44	5 (7%) 18 7	81, 104, 124, 138	0
50	K8	68/72 (94%)	0.12	1 (1%) 76 49	59, 79, 96, 120	0
51	H5	58/60 (96%)	2.04	29 (50%) 0 0	77, 99, 122, 135	0
51	L8	58/60 (96%)	0.54	2 (3%) 49 21	60, 77, 103, 110	0
52	M8	60/71 (84%)	1.54	16 (26%) 1 0	102, 143, 168, 171	0
53	J5	56/60 (93%)	0.72	7 (12%) 5 2	58, 85, 131, 141	0
53	N8	48/60 (80%)	0.35	0 100 100	46, 78, 119, 127	0
54	O8	45/54 (83%)	4.56	40 (88%) 0 0	117, 153, 171, 177	0
55	L5	47/49 (95%)	0.47	2 (4%) 39 16	47, 60, 87, 97	0
55	P8	47/49 (95%)	0.09	1 (2%) 67 36	42, 50, 72, 87	0
56	M5	64/65 (98%)	1.47	18 (28%) 1 0	68, 80, 97, 117	0
56	Q8	64/65 (98%)	0.36	0 100 100	51, 65, 80, 98	0
All	All	20730/21798 (95%)	0.42	1700 (8%) 14 5	40, 99, 184, 267	0

All (1700) RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
46	C5	59	GLY	14.7
14	5A	38	GLY	12.9
33	59	96	ALA	12.6
46	C5	49	VAL	12.1
52	M8	40	HIS	11.7
43	D8	37	VAL	11.6

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Mol	Chain	Res	Type	RSRZ
14	5A	39	LEU	11.3
12	3A	64	TYR	10.8
33	59	17	VAL	10.7
9	82	115	GLY	9.9
46	C5	58	GLY	9.9
46	C5	50	ARG	9.8
19	AA	68	GLY	9.3
54	O8	13	CYS	9.3
14	5A	26	ARG	9.2
14	5A	25	VAL	9.2
54	O8	19	ARG	9.0
12	3A	28	LYS	8.8
14	5A	34	TYR	8.8
54	O8	14	THR	8.3
37	35	110	TYR	8.3
40	A8	110	LEU	8.2
54	O8	52	VAL	8.2
54	O8	20	ASN	8.1
46	C5	60	PHE	7.9
54	O8	34	LEU	7.9
14	5A	30	ALA	7.8
12	3A	27	LEU	7.6
54	O8	53	LYS	7.6
14	5A	37	PHE	7.5
10	1A	47	PHE	7.4
33	59	170	ARG	7.4
47	D5	142	SER	7.4
47	H8	113	ALA	7.3
50	G5	44	LEU	7.2
54	O8	26	ASN	7.2
13	4A	102	ARG	7.2
33	59	83	TYR	7.1
46	C5	29	GLU	7.1
33	59	168	PRO	7.1
54	O8	50	ARG	7.1
54	O8	42	TRP	7.0
14	5A	53	LEU	7.0
54	O8	43	CYS	7.0
33	59	99	VAL	7.0
46	C5	47	LYS	7.0
33	59	107	VAL	6.9
7	6E	81	GLY	6.9

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Mol	Chain	Res	Type	RSRZ
45	B5	69	TYR	6.9
47	D5	9	TYR	6.9
14	5A	41	ARG	6.8
54	O8	18	ARG	6.8
14	5A	31	ARG	6.7
12	3A	19	ARG	6.6
9	82	109	VAL	6.6
3	22	6	HIS	6.6
33	59	87	LEU	6.6
14	5A	51	GLY	6.5
10	1A	54	PHE	6.5
33	59	33	LEU	6.5
46	C5	63	LYS	6.5
46	C5	61	ILE	6.4
26	14	888	C	6.4
10	1A	62	HIS	6.4
25	4L	25	A	6.3
33	59	128	PRO	6.3
14	5A	35	ARG	6.3
14	5A	42	ILE	6.3
41	B8	106	SER	6.3
24	1L	1	G	6.2
24	1L	71	C	6.2
2	12	62	ALA	6.2
19	AA	67	VAL	6.2
33	59	169	VAL	6.2
12	3A	63	GLY	6.2
22	1K	76	A	6.2
7	62	16	LEU	6.1
33	59	103	LEU	6.1
7	6E	80	VAL	6.1
10	1A	65	LEU	6.1
7	6E	85	TYR	6.0
33	59	95	ARG	6.0
7	6E	84	ASN	6.0
33	59	105	LEU	5.9
10	1I	10	GLY	5.9
52	M8	41	PRO	5.9
54	O8	49	HIS	5.9
33	59	164	TYR	5.8
12	3A	62	SER	5.8
7	6E	83	ALA	5.8

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Mol	Chain	Res	Type	RSRZ
2	12	72	GLY	5.8
7	6E	78	ARG	5.7
33	59	76	VAL	5.7
32	49	139	LEU	5.7
3	22	39	ILE	5.7
10	1I	22	LYS	5.7
16	7I	32	TYR	5.7
45	B5	68	ARG	5.7
46	C5	46	LYS	5.6
19	AA	40	ILE	5.6
12	3A	20	LYS	5.6
12	3I	64	TYR	5.6
12	3A	68	ALA	5.6
47	D5	155	LEU	5.6
6	5E	46	ARG	5.6
45	B5	79	ALA	5.6
47	D5	146	ILE	5.6
12	3A	21	LYS	5.6
32	49	138	GLN	5.5
51	H5	6	VAL	5.5
14	5A	6	LEU	5.5
52	M8	31	ILE	5.5
32	49	90	LEU	5.5
33	59	115	VAL	5.4
34	69	1	MET	5.4
10	1A	61	GLU	5.4
2	1E	10	LEU	5.4
33	59	25	LYS	5.4
54	O8	32	ASN	5.4
14	5A	33	VAL	5.3
12	3A	32	PHE	5.3
9	82	116	LYS	5.3
33	59	26	VAL	5.3
3	22	198	VAL	5.3
33	59	106	THR	5.3
14	5A	58	LYS	5.2
10	1A	59	SER	5.2
20	BA	9	ASN	5.2
31	39	10	PRO	5.2
28	71	175	VAL	5.2
46	C5	48	ALA	5.2
47	H8	165	VAL	5.2

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Mol	Chain	Res	Type	RSRZ
47	D5	112	ARG	5.2
29	19	26	LYS	5.2
38	88	104	PHE	5.2
47	D5	121	HIS	5.2
47	D5	69	THR	5.2
33	59	130	ARG	5.2
12	3I	61	THR	5.2
33	59	90	LYS	5.1
19	AI	71	LEU	5.1
54	O8	12	GLU	5.1
3	22	177	THR	5.1
33	59	162	ILE	5.1
24	3K	6	G	5.1
54	O8	25	LYS	5.1
47	H8	96	VAL	5.1
7	6E	79	ARG	5.1
2	12	152	PHE	5.0
47	D5	68	PRO	5.0
13	4A	111	LYS	5.0
33	59	114	VAL	5.0
47	D5	179	ASP	5.0
13	4A	95	GLY	5.0
46	C5	44	ILE	5.0
51	H5	26	LEU	5.0
10	1A	48	THR	5.0
14	5A	50	LYS	5.0
14	5A	44	LEU	5.0
33	59	171	LEU	5.0
32	49	133	LEU	5.0
3	2E	193	TYR	4.9
47	H8	147	GLY	4.9
10	1A	49	VAL	4.9
14	5A	23	ARG	4.9
33	59	94	TYR	4.9
19	AA	53	ASN	4.9
30	29	77	ILE	4.9
52	M8	22	ILE	4.9
25	4K	25	A	4.9
30	29	76	ARG	4.9
54	O8	23	THR	4.9
46	C5	53	PRO	4.9
9	82	110	GLU	4.9

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Mol	Chain	Res	Type	RSRZ
9	82	123	PRO	4.9
11	2A	75	TYR	4.9
52	M8	66	SER	4.9
35	15	9	VAL	4.8
33	59	89	ILE	4.8
38	88	33	GLY	4.8
10	1A	63	PHE	4.8
9	82	102	LEU	4.8
48	I8	8	GLY	4.8
32	41	80	PHE	4.8
14	5A	29	ARG	4.8
14	5A	36	PHE	4.8
54	O8	22	ALA	4.8
7	6E	154	TYR	4.8
47	D5	153	SER	4.7
17	8I	36	ILE	4.7
47	H8	166	SER	4.7
43	D8	38	LEU	4.7
9	82	20	ARG	4.7
12	3I	20	LYS	4.7
49	F5	91	LYS	4.7
47	D5	5	LEU	4.7
47	D5	150	LEU	4.7
33	59	16	SER	4.7
40	A8	49	VAL	4.7
3	22	28	GLN	4.7
33	59	153	LYS	4.7
47	D5	70	LEU	4.6
35	58	72	TYR	4.6
12	3I	19	ARG	4.6
40	A8	48	LEU	4.6
37	35	71	VAL	4.6
24	3L	34	U	4.6
7	6E	82	GLY	4.6
7	6E	16	LEU	4.6
13	4A	103	THR	4.5
47	D5	171	ILE	4.5
54	O8	15	GLU	4.5
14	5A	22	THR	4.5
32	41	25	TYR	4.5
40	65	60	GLY	4.5
47	H8	70	LEU	4.5

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Mol	Chain	Res	Type	RSRZ
3	22	186	PHE	4.5
2	12	165	VAL	4.5
29	19	27	THR	4.5
32	49	62	LEU	4.5
3	22	35	GLU	4.5
10	1I	94	VAL	4.5
14	5A	10	ALA	4.4
46	C5	56	PRO	4.4
7	62	61	VAL	4.4
14	5A	47	LEU	4.4
51	H5	30	ARG	4.4
54	O8	31	PRO	4.4
54	O8	10	LEU	4.4
47	H8	146	ILE	4.4
50	G5	43	GLN	4.4
20	BI	55	ILE	4.4
33	59	88	LEU	4.4
45	B5	92	LEU	4.4
33	59	49	VAL	4.4
54	O8	16	CYS	4.4
3	22	53	ALA	4.4
51	H5	15	TYR	4.4
43	D8	1	MET	4.4
30	21	88	GLY	4.4
47	D5	144	LEU	4.3
47	H8	104	PHE	4.3
2	12	133	LYS	4.3
8	7E	63	LEU	4.3
33	59	155	SER	4.3
10	1A	51	ARG	4.3
3	22	101	LEU	4.3
30	29	78	LEU	4.3
56	M5	12	LYS	4.3
53	J5	56	LYS	4.3
12	3A	23	LYS	4.3
44	E8	92	ARG	4.3
16	7I	35	LYS	4.3
20	BI	72	LEU	4.3
32	49	34	LEU	4.3
28	71	176	GLY	4.3
16	7I	48	TRP	4.3
17	8A	7	THR	4.2

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Mol	Chain	Res	Type	RSRZ
19	AA	62	ILE	4.2
28	71	13	LYS	4.2
33	59	123	PHE	4.2
13	4A	98	VAL	4.2
38	45	104	PHE	4.2
2	12	224	GLN	4.2
47	D5	51	ALA	4.2
33	59	132	ARG	4.2
12	3A	69	TYR	4.2
14	5A	46	GLU	4.2
47	H8	153	SER	4.2
54	O8	29	ASN	4.2
30	29	71	GLY	4.2
47	H8	38	TYR	4.2
40	65	55	ALA	4.2
47	D5	96	VAL	4.2
10	1I	90	LEU	4.2
51	H5	28	LEU	4.2
41	75	106	SER	4.2
43	95	1	MET	4.2
12	3A	85	ILE	4.1
9	82	36	TYR	4.1
49	J8	70	VAL	4.1
30	29	69	LYS	4.1
47	H8	98	MET	4.1
26	1H	1536	A	4.1
47	D5	125	LEU	4.1
32	49	152	LEU	4.1
47	H8	155	LEU	4.1
47	H8	8	TYR	4.1
9	82	53	VAL	4.1
9	82	50	LEU	4.1
11	2I	42	TRP	4.1
13	4I	96	LEU	4.1
40	65	58	LEU	4.1
47	H8	144	LEU	4.1
13	4A	101	GLN	4.1
10	1A	55	LYS	4.1
19	AA	71	LEU	4.1
7	6E	149	ARG	4.1
54	O8	21	TYR	4.1
31	31	6	VAL	4.1

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Mol	Chain	Res	Type	RSRZ
7	62	42	ILE	4.1
33	59	72	ILE	4.1
35	58	15	LEU	4.1
47	H8	161	VAL	4.0
30	29	3	GLY	4.0
32	49	142	PRO	4.0
54	O8	9	LEU	4.0
28	71	27	HIS	4.0
48	E5	8	GLY	4.0
46	C5	45	VAL	4.0
3	2E	200	ALA	4.0
25	4L	10	G	4.0
36	25	1	MET	4.0
33	59	10	PRO	4.0
32	49	92	VAL	4.0
47	D5	168	GLU	4.0
47	D5	173	ALA	4.0
2	12	163	PHE	4.0
33	59	34	GLU	4.0
17	8A	23	VAL	4.0
54	O8	33	LYS	3.9
38	45	102	VAL	3.9
32	49	39	ILE	3.9
48	E5	12	ASN	3.9
32	49	177	GLY	3.9
10	1I	95	GLU	3.9
10	1I	65	LEU	3.9
10	1I	72	VAL	3.9
20	BA	10	LEU	3.9
40	65	56	LEU	3.9
39	55	101	ALA	3.9
51	H5	5	LYS	3.9
8	7E	95	VAL	3.9
24	3L	35	U	3.9
2	12	134	GLU	3.9
8	7E	109	ILE	3.9
17	8I	98	LEU	3.9
14	5A	52	GLN	3.9
40	65	108	GLY	3.9
47	H8	1	MET	3.9
26	14	229	A	3.9
26	14	2799	A	3.9

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Mol	Chain	Res	Type	RSRZ
7	62	41	ARG	3.9
33	59	98	LEU	3.9
53	J5	55	ARG	3.9
3	22	8	ILE	3.8
33	59	125	VAL	3.8
52	M8	58	ARG	3.8
2	1E	232	PRO	3.8
10	1I	88	LEU	3.8
3	22	155	GLY	3.8
28	71	181	PRO	3.8
24	3K	36	U	3.8
9	82	114	TYR	3.8
44	A5	6	ILE	3.8
3	22	60	ALA	3.8
47	H8	164	ALA	3.8
35	15	8	GLN	3.8
38	45	103	MET	3.8
37	35	106	LEU	3.8
11	2A	14	VAL	3.8
12	3A	15	ARG	3.8
14	5A	24	CYS	3.8
49	F5	5	CYS	3.8
21	1B	13	ILE	3.8
30	29	67	PHE	3.8
30	29	73	GLU	3.8
10	1A	60	ARG	3.8
47	H8	157	LEU	3.8
14	5A	54	PRO	3.8
30	29	28	ALA	3.8
10	1I	96	ILE	3.8
33	59	131	VAL	3.8
47	H8	111	VAL	3.8
4	3E	96	LEU	3.8
26	14	2802	G	3.8
11	2A	25	TYR	3.8
30	29	70	ALA	3.8
33	59	97	ARG	3.8
33	59	124	GLU	3.8
49	F5	62	VAL	3.8
14	5A	49	HIS	3.8
26	14	2795	G	3.8
27	16	1(M)	A	3.8

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Mol	Chain	Res	Type	RSRZ
35	15	84	LYS	3.8
33	59	45	VAL	3.8
24	1L	3	G	3.7
39	55	70	LEU	3.7
55	L5	1	MET	3.7
9	8E	126	SER	3.7
47	D5	117	LEU	3.7
54	O8	36	LEU	3.7
56	M5	40	GLU	3.7
32	41	88	ILE	3.7
46	C5	86	ARG	3.7
17	8A	9	VAL	3.7
47	H8	5	LEU	3.7
56	M5	22	VAL	3.7
12	3A	65	GLU	3.7
38	45	100	GLY	3.7
47	D5	50	GLN	3.7
2	12	164	VAL	3.7
32	49	175	LEU	3.7
40	A8	112	PHE	3.7
17	8A	22	LEU	3.7
37	35	123	LEU	3.7
48	E5	71	ASP	3.7
49	F5	28	GLY	3.7
7	6E	86	GLN	3.7
10	1A	57	LYS	3.7
18	9A	84	LYS	3.7
12	3A	30	ALA	3.7
11	2A	83	ILE	3.7
47	H8	106	GLY	3.7
20	BA	42	GLN	3.7
3	22	23	TYR	3.7
11	2I	48	ILE	3.7
34	69	35	LEU	3.7
14	5A	59	ALA	3.6
33	59	111	HIS	3.6
28	71	32	LEU	3.6
32	49	82	LEU	3.6
45	B5	89	ILE	3.6
12	3A	33	ARG	3.6
20	BI	41	ILE	3.6
3	22	10	PHE	3.6

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Mol	Chain	Res	Type	RSRZ
30	29	29	GLY	3.6
10	1A	64	GLU	3.6
3	22	190	ARG	3.6
3	2E	94	LEU	3.6
12	3A	55	VAL	3.6
17	8A	6	LEU	3.6
47	H8	168	GLU	3.6
1	1G	82	U	3.6
40	A8	68	GLN	3.6
54	O8	40	CYS	3.6
46	C5	65	ALA	3.6
32	41	34	LEU	3.6
33	59	32	GLU	3.6
56	M5	16	ILE	3.6
33	59	29	PRO	3.6
21	1B	14	TRP	3.6
8	72	133	LEU	3.6
29	11	111	LEU	3.6
33	59	4	ILE	3.6
47	D5	141	VAL	3.6
13	4A	94	ARG	3.6
40	65	20	ARG	3.6
3	22	184	TYR	3.5
17	8A	32	TYR	3.5
38	45	17	LEU	3.5
9	82	120	ARG	3.5
34	61	107	VAL	3.5
46	G8	89	PHE	3.5
10	1A	46	ARG	3.5
47	H8	134	PRO	3.5
56	M5	64	TYR	3.5
20	BI	40	ALA	3.5
33	59	11	VAL	3.5
35	15	85	ILE	3.5
51	H5	9	VAL	3.5
17	8A	27	PHE	3.5
32	49	178	PHE	3.5
30	29	181	LEU	3.5
5	42	90	VAL	3.5
20	BI	100	ILE	3.5
56	M5	34	TRP	3.5
3	22	96	GLY	3.5

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Mol	Chain	Res	Type	RSRZ
24	1L	70	C	3.5
14	5A	55	GLY	3.5
38	45	33	GLY	3.5
11	2I	50	TYR	3.5
46	C5	30	VAL	3.5
46	C5	51	VAL	3.5
24	3K	45	G	3.5
47	D5	152	ALA	3.5
52	M8	13	ARG	3.5
10	1A	58	ASP	3.5
38	45	99	PRO	3.4
33	59	71	LEU	3.4
24	3K	70	C	3.4
3	22	7	PRO	3.4
47	D5	95	PRO	3.4
13	4A	87	TYR	3.4
47	H8	169	GLU	3.4
17	8I	95	TYR	3.4
13	4A	110	ARG	3.4
10	1A	56	HIS	3.4
30	29	4	ILE	3.4
35	58	85	ILE	3.4
35	15	73	THR	3.4
30	29	51	PHE	3.4
10	1A	53	PRO	3.4
45	B5	66	LEU	3.4
5	42	109	ILE	3.4
43	95	4	ILE	3.4
35	15	98	VAL	3.4
32	49	23	PHE	3.4
33	59	81	GLU	3.4
54	O8	30	THR	3.4
40	A8	59	LYS	3.4
40	65	57	LYS	3.4
8	7E	59	LEU	3.4
12	3A	24	VAL	3.4
37	78	71	VAL	3.4
40	65	109	GLY	3.4
47	D5	8	TYR	3.4
14	5A	17	LYS	3.4
45	B5	13	LEU	3.4
4	3E	3	ARG	3.4

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Mol	Chain	Res	Type	RSRZ
8	7E	46	LYS	3.4
35	15	51	PHE	3.4
46	C5	2	ARG	3.4
49	F5	61	ARG	3.4
1	1G	1202	G	3.4
12	3I	28	LYS	3.4
9	82	79	LEU	3.3
13	4A	96	LEU	3.3
12	3A	29	GLY	3.3
8	7E	58	TYR	3.3
35	15	12	ARG	3.3
33	59	122	THR	3.3
11	2A	21	ILE	3.3
20	BI	44	ALA	3.3
25	4L	9	G	3.3
3	22	94	LEU	3.3
31	31	123	LEU	3.3
12	3A	31	PRO	3.3
28	71	19	ILE	3.3
43	D8	99	ILE	3.3
38	45	65	PHE	3.3
38	45	22	LYS	3.3
37	78	106	LEU	3.3
26	1H	163	U	3.3
35	58	16	ILE	3.3
17	8A	21	VAL	3.3
32	41	23	PHE	3.3
32	49	160	VAL	3.3
24	1L	72	C	3.3
35	15	72	TYR	3.3
3	22	187	ALA	3.3
16	7I	19	ILE	3.3
33	59	100	GLY	3.3
32	49	179	PRO	3.3
47	H8	74	VAL	3.3
52	M8	3	GLU	3.3
40	A8	44	LYS	3.3
47	D5	114	GLY	3.3
40	65	40	ILE	3.3
33	59	159	GLU	3.3
7	62	5	ARG	3.3
6	5E	63	TYR	3.3

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Mol	Chain	Res	Type	RSRZ
35	15	87	LEU	3.3
43	D8	54	GLY	3.3
47	D5	143	GLY	3.3
13	4A	92	HIS	3.3
32	49	176	LEU	3.3
3	22	197	GLY	3.3
21	1B	2	GLY	3.3
26	1H	2476	A	3.3
33	51	9	ILE	3.3
43	95	14	VAL	3.3
35	15	48	MET	3.3
10	1I	33	GLN	3.3
12	3A	98	TYR	3.2
33	59	151	ILE	3.2
32	49	109	VAL	3.2
47	H8	141	VAL	3.2
13	4A	88	ARG	3.2
21	1B	22	ARG	3.2
30	29	49	LEU	3.2
51	H5	53	LEU	3.2
51	H5	48	GLU	3.2
38	88	132	VAL	3.2
13	4A	90	LEU	3.2
20	BI	17	ARG	3.2
17	8I	23	VAL	3.2
43	D8	45	THR	3.2
20	BA	8	ARG	3.2
37	35	125	VAL	3.2
11	2I	71	LYS	3.2
28	71	31	GLU	3.2
51	H5	19	GLN	3.2
17	8A	8	GLY	3.2
2	12	92	TYR	3.2
30	29	151	TYR	3.2
33	59	15	VAL	3.2
46	C5	52	SER	3.2
9	82	31	GLN	3.2
12	3I	63	GLY	3.2
40	A8	58	LEU	3.2
7	62	62	PHE	3.2
7	62	28	ASN	3.2
17	8A	92	ARG	3.2

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Mol	Chain	Res	Type	RSRZ
47	D5	134	PRO	3.2
47	D5	176	PRO	3.2
26	14	2141	G	3.2
13	4A	117	VAL	3.2
35	15	92	ALA	3.2
9	82	92	TYR	3.2
11	2I	98	LEU	3.2
14	5I	39	LEU	3.2
51	H5	2	PRO	3.2
54	O8	11	LEU	3.2
32	41	52	ILE	3.2
46	C5	62	GLU	3.2
12	3A	97	ARG	3.2
20	BI	59	ALA	3.2
2	1E	14	GLY	3.2
11	2I	25	TYR	3.2
13	4A	115	LYS	3.2
43	95	16	PRO	3.2
17	8A	71	PHE	3.2
46	C5	42	VAL	3.1
7	6E	99	LEU	3.1
7	62	12	LEU	3.1
9	82	59	PHE	3.1
34	61	111	PRO	3.1
11	2I	68	ALA	3.1
17	8I	27	PHE	3.1
32	49	155	MET	3.1
40	65	112	PHE	3.1
43	D8	98	GLU	3.1
41	75	50	ILE	3.1
3	2E	189	ALA	3.1
9	82	30	GLY	3.1
20	BA	99	LEU	3.1
47	H8	99	TYR	3.1
31	39	14	PRO	3.1
12	3A	16	GLU	3.1
16	7I	6	LEU	3.1
32	41	137	GLU	3.1
14	5A	45	ARG	3.1
16	7I	28	ARG	3.1
47	H8	2	GLU	3.1
7	6E	12	LEU	3.1

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Mol	Chain	Res	Type	RSRZ
8	72	2	LEU	3.1
20	BI	43	LEU	3.1
34	61	130	TYR	3.1
9	82	21	PRO	3.1
16	7I	36	ILE	3.1
32	49	89	GLY	3.1
54	O8	48	VAL	3.1
8	72	112	LEU	3.1
37	78	110	TYR	3.1
33	59	93	GLY	3.1
33	59	79	VAL	3.1
3	22	32	LEU	3.1
4	32	186	LEU	3.1
8	7E	133	LEU	3.1
17	8I	45	HIS	3.1
47	H8	156	LYS	3.1
47	D5	156	LYS	3.1
2	1E	96	ARG	3.1
36	25	42	SER	3.1
30	29	150	VAL	3.1
7	62	101	LEU	3.1
12	3I	17	LYS	3.1
12	3A	93	LEU	3.1
38	45	130	LYS	3.1
21	1B	6	ARG	3.1
47	H8	149	SER	3.0
47	D5	7	ALA	3.0
12	3A	46	LYS	3.0
32	49	135	LEU	3.0
56	M5	61	LEU	3.0
7	6E	151	TYR	3.0
24	3K	71	C	3.0
33	59	9	ILE	3.0
29	11	112	GLN	3.0
32	49	36	LYS	3.0
16	7I	49	LEU	3.0
7	62	27	ILE	3.0
16	7A	32	TYR	3.0
3	22	64	VAL	3.0
14	5A	56	VAL	3.0
12	3A	56	ALA	3.0
47	H8	66	SER	3.0

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Mol	Chain	Res	Type	RSRZ
7	62	76	ARG	3.0
34	61	128	LEU	3.0
3	22	31	HIS	3.0
37	35	108	LYS	3.0
41	75	1	MET	3.0
43	95	99	ILE	3.0
35	58	55	VAL	3.0
49	F5	70	VAL	3.0
9	82	106	ALA	3.0
40	A8	37	ALA	3.0
43	95	94	LEU	3.0
43	95	96	ILE	3.0
16	7I	7	ALA	3.0
33	59	86	GLU	3.0
35	58	73	THR	3.0
3	22	199	LYS	3.0
47	H8	67	LEU	3.0
11	2A	91	ARG	3.0
11	2A	84	VAL	3.0
7	62	37	ASN	3.0
18	9I	78	LEU	3.0
24	3K	34	U	3.0
47	D5	162	GLU	3.0
2	12	96	ARG	3.0
3	2E	79	ARG	3.0
3	2E	190	ARG	3.0
10	1I	34	VAL	3.0
33	59	13	LYS	3.0
1	1G	1029	G	3.0
47	H8	133	ILE	3.0
7	62	73	MET	3.0
4	32	128	VAL	3.0
6	5E	9	VAL	3.0
9	82	65	VAL	3.0
42	85	90	VAL	3.0
11	2I	66	LEU	3.0
18	9A	43	PHE	3.0
43	95	12	TYR	3.0
29	19	44	ASN	3.0
16	7I	30	GLY	3.0
8	72	79	VAL	3.0
12	3A	18	VAL	3.0

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Mol	Chain	Res	Type	RSRZ
47	D5	126	VAL	3.0
47	D5	151	HIS	3.0
33	59	165	ALA	2.9
34	69	12	LEU	2.9
47	D5	108	PRO	2.9
32	41	48	GLU	2.9
46	C5	64	GLU	2.9
32	49	149	VAL	2.9
29	19	2	ALA	2.9
8	7E	112	LEU	2.9
11	2I	63	LEU	2.9
13	4I	6	GLY	2.9
17	8A	95	TYR	2.9
10	1A	44	VAL	2.9
11	2I	70	LYS	2.9
38	45	6	ARG	2.9
3	22	42	LEU	2.9
13	4A	107	ALA	2.9
4	32	187	ARG	2.9
20	BI	80	ARG	2.9
31	39	199	TRP	2.9
54	O8	47	THR	2.9
12	3A	60	LEU	2.9
47	D5	172	ALA	2.9
24	1L	31	A	2.9
7	6E	72	ARG	2.9
56	M5	6	THR	2.9
2	12	155	LEU	2.9
14	5A	48	ALA	2.9
32	41	26	GLN	2.9
12	3I	7	ILE	2.9
11	2I	43	SER	2.9
32	49	159	VAL	2.9
42	85	40	PHE	2.9
6	5E	14	LEU	2.9
32	49	137	GLU	2.9
38	45	37	LEU	2.9
8	7E	131	GLY	2.9
12	3I	91	LYS	2.9
3	22	30	ARG	2.9
8	72	74	PRO	2.9
11	2A	35	PRO	2.9

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Mol	Chain	Res	Type	RSRZ
54	O8	44	ARG	2.9
50	K8	43	GLN	2.9
2	12	129	GLU	2.9
3	2E	151	VAL	2.9
4	32	17	VAL	2.9
2	12	102	LEU	2.9
10	1I	46	ARG	2.9
47	H8	109	ALA	2.9
3	22	201	TYR	2.9
17	8A	11	VAL	2.9
33	59	141	VAL	2.9
3	2E	60	ALA	2.9
3	22	188	LEU	2.9
47	D5	28	MET	2.9
51	H5	11	SER	2.8
3	22	189	ALA	2.8
25	4K	13	A	2.8
34	61	123	LEU	2.8
14	5A	32	SER	2.8
34	61	103	ARG	2.8
39	98	34	ILE	2.8
46	C5	84	ARG	2.8
30	29	131	ALA	2.8
43	95	44	LYS	2.8
14	5A	60	SER	2.8
33	59	134	SER	2.8
1	1G	1226	C	2.8
37	78	91	PHE	2.8
33	59	85	LYS	2.8
33	59	113	VAL	2.8
33	59	133	VAL	2.8
47	D5	91	LEU	2.8
48	I8	84	LEU	2.8
56	M5	50	LEU	2.8
9	82	122	ALA	2.8
47	H8	103	ARG	2.8
2	12	223	ILE	2.8
10	1I	36	GLY	2.8
12	3A	17	LYS	2.8
46	C5	54	LYS	2.8
47	H8	114	GLY	2.8
8	7E	10	LEU	2.8

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Mol	Chain	Res	Type	RSRZ
46	C5	43	ASN	2.8
47	D5	61	LEU	2.8
5	4E	152	ARG	2.8
29	11	262	ARG	2.8
30	29	79	ARG	2.8
12	3A	88	GLY	2.8
30	21	195	LEU	2.8
45	B5	26	TYR	2.8
47	H8	163	LEU	2.8
47	D5	59	LEU	2.8
49	J8	80	LEU	2.8
52	M8	32	TYR	2.8
51	H5	55	ARG	2.8
26	14	2797	U	2.8
48	E5	46	LYS	2.8
4	32	146	ILE	2.8
17	8A	59	ILE	2.8
30	29	56	PRO	2.8
48	E5	42	GLY	2.8
7	6E	153	HIS	2.8
9	82	37	PHE	2.8
35	15	41	ASP	2.8
9	82	19	LEU	2.8
38	45	34	LEU	2.8
52	M8	20	ASN	2.8
8	7E	6	ILE	2.8
40	65	35	ILE	2.8
48	E5	45	PHE	2.8
2	1E	187	LEU	2.8
18	9A	46	GLU	2.8
33	59	101	ARG	2.8
7	6E	60	LYS	2.8
11	2A	19	ALA	2.8
20	BI	45	GLN	2.8
8	72	83	ILE	2.8
17	8I	99	SER	2.8
32	49	35	GLU	2.8
33	59	46	GLU	2.8
56	M5	29	LYS	2.8
51	H5	37	LEU	2.8
33	59	129	THR	2.8
23	2K	1	C	2.8

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Mol	Chain	Res	Type	RSRZ
46	C5	5	MET	2.7
10	1I	64	GLU	2.7
33	59	12	PRO	2.7
35	15	86	PRO	2.7
9	82	32	ASP	2.7
8	72	129	VAL	2.7
11	2I	82	VAL	2.7
17	8A	10	VAL	2.7
35	58	52	VAL	2.7
38	45	97	VAL	2.7
39	98	113	LEU	2.7
22	1K	5	C	2.7
41	B8	104	ASN	2.7
17	8A	36	ILE	2.7
45	F8	89	ILE	2.7
47	D5	83	PRO	2.7
29	19	18	VAL	2.7
17	8I	44	ALA	2.7
46	C5	69	ALA	2.7
4	32	169	LYS	2.7
14	5A	11	LYS	2.7
17	8I	37	LYS	2.7
46	C5	81	LYS	2.7
9	82	34	ASN	2.7
47	D5	119	GLU	2.7
10	1I	23	ILE	2.7
11	2A	108	ILE	2.7
20	BA	41	ILE	2.7
30	29	74	PRO	2.7
3	22	43	LEU	2.7
12	3A	84	LEU	2.7
20	BA	45	GLN	2.7
28	71	11	LEU	2.7
32	49	41	GLN	2.7
43	D8	39	LEU	2.7
8	72	98	LYS	2.7
47	H8	21	ALA	2.7
47	H8	131	ARG	2.7
49	J8	21	ARG	2.7
11	2A	95	ILE	2.7
15	6A	15	PHE	2.7
7	62	9	VAL	2.7

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Mol	Chain	Res	Type	RSRZ
53	J5	45	VAL	2.7
3	22	100	ALA	2.7
39	98	109	ALA	2.7
43	95	27	ALA	2.7
14	5I	8	GLU	2.7
11	2A	20	TYR	2.7
52	M8	25	TYR	2.7
56	M5	9	GLY	2.7
16	7I	4	ILE	2.7
56	M5	58	ILE	2.7
51	H5	20	LYS	2.7
7	62	104	LEU	2.7
37	35	77	ARG	2.7
38	45	105	GLU	2.7
49	F5	69	LYS	2.7
51	H5	10	LYS	2.7
48	I8	82	ARG	2.7
2	12	43	ASP	2.7
3	2E	66	VAL	2.7
20	BA	84	LEU	2.7
32	49	150	ASP	2.7
39	98	114	VAL	2.7
47	H8	105	VAL	2.7
47	D5	76	LEU	2.7
9	82	52	ALA	2.7
8	72	86	ILE	2.7
11	2A	18	ARG	2.7
32	41	135	LEU	2.7
32	49	107	LEU	2.7
51	H5	31	LEU	2.7
19	AA	38	SER	2.7
11	2A	40	ILE	2.7
37	35	75	ILE	2.7
32	49	161	THR	2.7
34	61	114	LEU	2.7
43	95	5	VAL	2.7
17	8A	91	ARG	2.7
52	M8	43	TYR	2.6
6	5E	55	ASP	2.6
20	BA	104	LEU	2.6
28	71	21	THR	2.6
51	H5	4	LEU	2.6

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Mol	Chain	Res	Type	RSRZ
12	3A	94	PRO	2.6
47	H8	25	PRO	2.6
32	49	136	ARG	2.6
11	2I	73	MET	2.6
8	7E	48	TYR	2.6
33	59	136	ILE	2.6
33	59	167	GLU	2.6
47	D5	57	ILE	2.6
34	61	77	LEU	2.6
19	AA	76	PRO	2.6
35	58	61	ARG	2.6
20	BI	95	ALA	2.6
7	62	26	PHE	2.6
40	65	19	LYS	2.6
41	B8	48	ILE	2.6
20	BI	18	GLN	2.6
20	BA	53	LEU	2.6
35	15	26	LEU	2.6
11	2I	47	VAL	2.6
31	39	114	VAL	2.6
36	25	58	VAL	2.6
16	7I	22	THR	2.6
44	E8	104	THR	2.6
12	3A	26	ALA	2.6
20	BI	87	LYS	2.6
34	69	83	ALA	2.6
24	3K	65	C	2.6
33	59	92	ILE	2.6
37	78	114	ILE	2.6
9	82	96	LEU	2.6
12	3A	89	ARG	2.6
14	5A	21	TYR	2.6
38	88	17	LEU	2.6
43	D8	25	LEU	2.6
43	95	73	SER	2.6
30	29	141	ILE	2.6
38	45	64	ILE	2.6
12	3A	90	VAL	2.6
30	21	151	TYR	2.6
31	39	131	GLY	2.6
32	49	15	VAL	2.6
34	69	3	VAL	2.6

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Mol	Chain	Res	Type	RSRZ
47	D5	27	VAL	2.6
12	3I	94	PRO	2.6
33	59	55	PRO	2.6
33	59	127	GLU	2.6
22	1K	73	A	2.6
16	7I	18	ARG	2.6
49	F5	21	ARG	2.6
33	59	41	MET	2.6
38	45	7	MET	2.6
49	F5	66	HIS	2.6
32	49	94	LEU	2.6
39	98	98	LEU	2.6
2	1E	15	VAL	2.6
17	8A	57	VAL	2.6
28	71	41	VAL	2.6
40	65	49	VAL	2.6
49	F5	2	SER	2.6
7	62	43	PHE	2.6
17	8I	28	PRO	2.6
46	C5	92	ASN	2.6
50	G5	37	PHE	2.6
7	62	97	GLN	2.6
10	1A	50	ILE	2.6
24	1L	2	G	2.6
37	35	124	LYS	2.6
47	D5	46	LYS	2.6
7	6E	59	LEU	2.6
10	1I	71	LEU	2.6
33	59	161	GLY	2.6
40	65	24	LEU	2.6
43	95	18	LEU	2.6
30	21	89	ASP	2.6
30	29	75	VAL	2.6
47	H8	140	ASP	2.6
47	D5	56	VAL	2.6
9	8E	101	PHE	2.6
5	42	121	LYS	2.6
12	3A	48	PRO	2.6
49	F5	7	ILE	2.6
17	8A	43	LEU	2.6
7	62	146	GLU	2.6
26	14	2146	C	2.6

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Mol	Chain	Res	Type	RSRZ
30	29	34	VAL	2.6
38	88	27	VAL	2.6
35	58	74	ARG	2.6
38	45	32	TYR	2.6
9	8E	127	LYS	2.6
3	22	167	TRP	2.6
31	39	172	TRP	2.6
35	15	126	PRO	2.6
2	1E	208	ILE	2.5
9	82	38	GLN	2.5
30	29	52	LEU	2.5
47	H8	125	LEU	2.5
47	D5	60	GLU	2.5
2	12	193	ASP	2.5
30	29	116	VAL	2.5
38	88	102	VAL	2.5
5	4E	45	PHE	2.5
10	1I	11	PHE	2.5
28	71	170	ALA	2.5
37	35	51	PHE	2.5
7	62	88	PRO	2.5
13	4A	97	PRO	2.5
7	62	74	GLU	2.5
43	D8	40	LEU	2.5
51	H5	8	LEU	2.5
51	H5	23	LEU	2.5
20	BI	56	MET	2.5
24	3L	12	U	2.5
49	F5	10	LYS	2.5
56	M5	21	LYS	2.5
3	2E	153	VAL	2.5
8	7E	137	VAL	2.5
11	2I	75	TYR	2.5
40	65	87	PHE	2.5
53	J5	2	ALA	2.5
47	H8	158	PRO	2.5
7	62	30	ILE	2.5
8	7E	47	GLY	2.5
11	2I	29	ILE	2.5
33	59	14	GLY	2.5
36	25	19	ILE	2.5
51	H5	43	ILE	2.5

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Mol	Chain	Res	Type	RSRZ
52	M8	30	GLU	2.5
20	BA	13	LEU	2.5
34	69	72	LEU	2.5
35	15	116	LEU	2.5
42	85	25	TRP	2.5
2	12	101	MET	2.5
30	29	198	VAL	2.5
34	69	21	VAL	2.5
3	22	133	ALA	2.5
47	D5	79	ARG	2.5
41	75	99	LEU	2.5
33	59	160	LYS	2.5
40	65	33	LYS	2.5
30	29	50	GLY	2.5
50	G5	45	SER	2.5
38	45	66	ILE	2.5
4	3E	135	LEU	2.5
9	82	85	LEU	2.5
35	15	46	VAL	2.5
9	82	54	ASP	2.5
14	5I	37	PHE	2.5
30	21	51	PHE	2.5
35	15	130	HIS	2.5
47	D5	6	LYS	2.5
49	J8	69	LYS	2.5
44	E8	90	ARG	2.5
48	E5	76	GLY	2.5
2	1E	201	ILE	2.5
12	3A	100	ILE	2.5
30	29	30	PRO	2.5
34	69	4	ILE	2.5
34	69	80	PRO	2.5
40	A8	35	ILE	2.5
17	8A	31	LEU	2.5
30	21	49	LEU	2.5
31	31	196	LEU	2.5
32	41	139	LEU	2.5
12	3A	83	VAL	2.5
33	59	24	VAL	2.5
34	61	131	LYS	2.5
37	35	46	LYS	2.5
3	22	132	ARG	2.5

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Mol	Chain	Res	Type	RSRZ
11	2I	96	ARG	2.5
14	5I	13	THR	2.5
21	1B	9	ARG	2.5
36	25	33	ALA	2.5
12	3A	35	GLY	2.5
38	88	32	TYR	2.5
47	H8	57	ILE	2.5
51	H5	13	ILE	2.5
10	1I	7	LYS	2.5
17	8I	10	VAL	2.5
47	H8	39	VAL	2.5
9	82	93	ARG	2.5
14	5A	61	TRP	2.5
36	25	11	ALA	2.5
49	J8	42	GLN	2.5
16	7I	39	TYR	2.5
3	22	154	SER	2.5
2	12	71	VAL	2.5
6	5E	88	VAL	2.5
20	BI	15	ARG	2.5
30	21	7	VAL	2.5
47	H8	97	GLU	2.5
1	1G	1225	A	2.5
9	82	8	GLY	2.5
33	59	102	ALA	2.5
35	15	13	TRP	2.5
36	68	53	LYS	2.5
41	B8	114	LEU	2.5
48	E5	75	LEU	2.5
54	O8	39	TYR	2.5
2	12	131	PRO	2.4
22	1K	71	C	2.4
26	14	2794	C	2.4
48	E5	9	SER	2.4
14	5A	4	LYS	2.4
20	BI	42	GLN	2.4
3	2E	196	LEU	2.4
31	39	9	ILE	2.4
48	E5	21	LEU	2.4
47	H8	95	PRO	2.4
47	D5	52	SER	2.4
3	2E	72	LYS	2.4

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Mol	Chain	Res	Type	RSRZ
8	72	61	VAL	2.4
9	82	108	VAL	2.4
12	3I	11	VAL	2.4
35	15	53	VAL	2.4
39	98	48	VAL	2.4
56	M5	14	VAL	2.4
30	29	1	MET	2.4
42	C8	4	ALA	2.4
14	5I	29	ARG	2.4
32	41	164	GLU	2.4
32	49	63	ILE	2.4
47	H8	107	THR	2.4
11	2A	70	LYS	2.4
53	J5	10	LYS	2.4
4	3E	110	PHE	2.4
8	7E	44	PHE	2.4
17	8A	19	VAL	2.4
34	69	19	VAL	2.4
35	58	51	PHE	2.4
7	6E	5	ARG	2.4
7	62	32	ARG	2.4
34	69	36	ALA	2.4
37	35	111	ARG	2.4
7	6E	136	LYS	2.4
11	2A	66	LEU	2.4
37	78	138	LEU	2.4
7	62	17	VAL	2.4
38	45	106	VAL	2.4
47	H8	90	VAL	2.4
31	31	133	ASN	2.4
8	7E	64	LYS	2.4
10	1I	97	GLU	2.4
20	BA	48	LYS	2.4
31	39	148	LEU	2.4
43	95	20	LEU	2.4
11	2A	31	THR	2.4
24	1L	23	A	2.4
7	62	91	VAL	2.4
17	8A	85	VAL	2.4
11	2I	49	GLY	2.4
51	H5	29	ARG	2.4
33	59	140	LYS	2.4

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Mol	Chain	Res	Type	RSRZ
54	O8	35	GLU	2.4
43	95	3	ALA	2.4
51	H5	21	ALA	2.4
3	2E	101	LEU	2.4
8	7E	45	ILE	2.4
44	A5	103	ILE	2.4
47	H8	59	LEU	2.4
9	82	26	VAL	2.4
11	2A	28	THR	2.4
15	6A	88	ARG	2.4
32	49	141	PHE	2.4
39	55	21	TYR	2.4
47	H8	167	PRO	2.4
11	2A	42	TRP	2.4
33	59	91	GLY	2.4
49	F5	92	LYS	2.4
51	H5	7	LYS	2.4
54	O8	17	LYS	2.4
36	25	41	ALA	2.4
2	12	185	ILE	2.4
10	1I	8	LEU	2.4
17	8I	59	ILE	2.4
29	19	270	ILE	2.4
30	29	195	LEU	2.4
36	25	25	LEU	2.4
41	75	48	ILE	2.4
43	D8	35	LEU	2.4
1	1G	973	G	2.4
4	32	166	LYS	2.4
7	62	118	VAL	2.4
14	5I	51	GLY	2.4
40	A8	45	GLY	2.4
47	D5	147	GLY	2.4
2	12	97	TRP	2.4
17	8A	82	MET	2.4
4	3E	21	LEU	2.4
8	7E	107	LEU	2.4
9	82	56	LEU	2.4
31	31	181	LEU	2.4
35	15	91	LEU	2.4
47	H8	41	LEU	2.4
38	45	10	ARG	2.4

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Mol	Chain	Res	Type	RSRZ
38	45	98	LYS	2.4
51	H5	35	ARG	2.4
30	29	96	PHE	2.4
48	I8	69	PHE	2.4
34	61	126	TYR	2.4
10	1A	67	THR	2.4
47	D5	107	THR	2.4
9	82	46	ALA	2.4
38	88	28	ALA	2.4
7	62	22	LEU	2.4
9	8E	128	ARG	2.4
11	2A	48	ILE	2.4
11	2A	96	ARG	2.4
31	31	41	LEU	2.4
32	41	94	LEU	2.4
3	2E	166	GLU	2.3
8	7E	53	VAL	2.3
34	69	144	VAL	2.3
9	82	7	THR	2.3
30	21	90	THR	2.3
37	78	1	MET	2.3
4	3E	111	ALA	2.3
13	4I	102	ARG	2.3
16	7I	8	ARG	2.3
13	4A	25	ILE	2.3
39	55	4	LEU	2.3
40	65	48	LEU	2.3
40	65	82	ILE	2.3
46	G8	92	ASN	2.3
13	4A	73	GLU	2.3
17	8I	29	HIS	2.3
47	H8	128	VAL	2.3
13	4I	87	TYR	2.3
32	49	146	TYR	2.3
41	75	35	LYS	2.3
46	C5	87	LYS	2.3
7	6E	144	MET	2.3
37	35	15	ARG	2.3
7	62	2	ALA	2.3
3	22	152	ILE	2.3
9	82	71	SER	2.3
12	3I	27	LEU	2.3

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Mol	Chain	Res	Type	RSRZ
14	5I	44	LEU	2.3
26	1H	2062	A	2.3
26	14	277	C	2.3
43	95	46	VAL	2.3
52	M8	33	VAL	2.3
53	J5	28	PRO	2.3
33	59	163	TYR	2.3
2	12	118	LEU	2.3
8	7E	22	GLU	2.3
8	7E	35	ILE	2.3
15	6I	56	LEU	2.3
29	11	165	ILE	2.3
33	51	171	LEU	2.3
40	A8	43	GLU	2.3
49	J8	90	ILE	2.3
3	22	20	SER	2.3
49	F5	60	PHE	2.3
12	3A	99	HIS	2.3
28	71	58	VAL	2.3
36	25	121	VAL	2.3
38	88	106	VAL	2.3
47	D5	32	HIS	2.3
7	6E	73	MET	2.3
39	98	94	TYR	2.3
2	12	187	LEU	2.3
7	6E	22	LEU	2.3
10	1I	38	ILE	2.3
16	7A	33	ILE	2.3
49	J8	82	LEU	2.3
20	BA	14	LYS	2.3
46	C5	4	LYS	2.3
12	3I	62	SER	2.3
39	98	32	GLY	2.3
9	82	111	ARG	2.3
13	4A	80	ARG	2.3
20	BA	83	ARG	2.3
47	H8	82	ARG	2.3
40	A8	85	VAL	2.3
1	13	344	A	2.3
24	3K	23	A	2.3
17	8A	51	TYR	2.3
37	35	138	LEU	2.3

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Mol	Chain	Res	Type	RSRZ
49	F5	32	LYS	2.3
4	3E	23	GLY	2.3
7	6E	143	ARG	2.3
8	72	131	GLY	2.3
10	1A	52	GLY	2.3
39	98	33	ARG	2.3
48	I8	68	GLU	2.3
24	3K	32	C	2.3
7	62	31	MET	2.3
35	15	75	TYR	2.3
49	F5	71	TYR	2.3
9	82	74	ILE	2.3
7	6E	26	PHE	2.3
13	4A	100	GLY	2.3
20	BI	101	GLY	2.3
46	C5	75	ILE	2.3
45	B5	28	PHE	2.3
32	49	108	ASN	2.3
7	6E	87	VAL	2.3
12	3A	66	VAL	2.3
40	A8	46	VAL	2.3
46	C5	28	LYS	2.3
54	O8	27	LYS	2.3
11	2I	81	ASP	2.3
47	D5	148	ASP	2.3
2	1E	196	LEU	2.3
4	32	176	LEU	2.3
22	1K	75	C	2.3
32	41	7	LEU	2.3
45	B5	46	ALA	2.3
8	72	111	ILE	2.3
11	2I	83	ILE	2.3
26	1H	2790	A	2.3
32	49	80	PHE	2.3
37	78	130	PHE	2.3
2	1E	165	VAL	2.3
14	5I	25	VAL	2.3
32	41	35	GLU	2.3
2	12	26	PRO	2.3
32	49	83	ARG	2.3
8	7E	119	LEU	2.3
3	22	124	ILE	2.3

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Mol	Chain	Res	Type	RSRZ
4	3E	138	TYR	2.3
8	7E	130	GLY	2.3
41	75	102	ILE	2.3
7	62	70	LYS	2.3
35	58	84	LYS	2.3
37	35	64	LYS	2.3
9	82	35	GLU	2.3
43	95	15	GLU	2.3
3	22	15	THR	2.2
31	39	126	VAL	2.2
37	35	126	VAL	2.2
3	22	59	ARG	2.2
34	61	113	ARG	2.2
36	25	17	ARG	2.2
9	82	80	GLY	2.2
9	82	94	ALA	2.2
28	71	185	LEU	2.2
39	98	75	LEU	2.2
40	A8	105	ALA	2.2
32	49	11	TYR	2.2
32	49	5	VAL	2.2
37	35	95	VAL	2.2
24	1L	24	G	2.2
33	59	84	SER	2.2
15	6I	70	LEU	2.2
33	59	108	GLY	2.2
34	61	140	LEU	2.2
39	55	29	LEU	2.2
47	D5	55	HIS	2.2
7	6E	103	TRP	2.2
38	45	69	PHE	2.2
50	G5	41	ILE	2.2
3	2E	198	VAL	2.2
4	32	178	VAL	2.2
5	4E	24	ARG	2.2
32	49	37	VAL	2.2
32	49	95	ARG	2.2
37	78	79	ARG	2.2
29	19	4	LYS	2.2
36	68	66	LYS	2.2
44	A5	98	LYS	2.2
5	42	31	LEU	2.2

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Mol	Chain	Res	Type	RSRZ
10	1I	85	LEU	2.2
5	42	81	GLU	2.2
33	59	117	PRO	2.2
43	D8	4	ILE	2.2
44	E8	93	ALA	2.2
9	82	51	ARG	2.2
9	82	125	TYR	2.2
17	8I	92	ARG	2.2
24	1L	65	C	2.2
26	1H	654(Q)	C	2.2
52	M8	21	VAL	2.2
33	59	166	GLY	2.2
3	2E	65	ALA	2.2
7	6E	134	ALA	2.2
40	A8	80	LEU	2.2
20	BA	16	HIS	2.2
32	49	102	PHE	2.2
47	H8	7	ALA	2.2
1	1G	1031	G	2.2
3	22	131	ARG	2.2
5	42	133	TYR	2.2
26	14	1509	C	2.2
31	39	147	GLY	2.2
39	98	102	GLU	2.2
49	J8	94	LEU	2.2
54	O8	51	GLU	2.2
7	6E	62	PHE	2.2
11	2A	85	ARG	2.2
11	2A	89	ALA	2.2
19	AA	4	SER	2.2
47	D5	92	SER	2.2
56	M5	62	LEU	2.2
20	BA	40	ALA	2.2
20	BI	38	LYS	2.2
28	71	193	ILE	2.2
30	29	149	ARG	2.2
32	41	142	PRO	2.2
37	78	124	LYS	2.2
43	95	81	TYR	2.2
19	AA	11	VAL	2.2
45	B5	29	TRP	2.2
7	62	148	ASN	2.2

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Mol	Chain	Res	Type	RSRZ
8	72	119	LEU	2.2
12	3A	13	LYS	2.2
20	BI	91	LEU	2.2
5	4E	89	ILE	2.2
9	82	76	ALA	2.2
10	1I	47	PHE	2.2
19	AI	75	ALA	2.2
32	41	178	PHE	2.2
46	G8	94	LYS	2.2
56	M5	5	LYS	2.2
38	45	68	ILE	2.2
47	D5	149	SER	2.2
8	72	101	PRO	2.2
26	14	2801	A	2.2
32	49	25	TYR	2.2
35	58	54	VAL	2.2
36	25	32	TYR	2.2
47	H8	58	VAL	2.2
47	H8	86	VAL	2.2
7	62	130	GLY	2.2
8	72	84	ARG	2.2
26	14	274	G	2.2
32	41	54	GLU	2.2
38	45	19	GLY	2.2
45	B5	86	GLY	2.2
53	J5	54	GLY	2.2
2	1E	94	ASN	2.2
4	3E	176	LEU	2.2
28	71	196	LEU	2.2
39	55	65	LEU	2.2
46	G8	100	ALA	2.2
56	M5	10	ALA	2.2
33	51	61	HIS	2.2
35	58	11	PRO	2.2
37	35	35	HIS	2.2
46	C5	57	GLN	2.2
24	1L	73	A	2.2
3	22	26	LYS	2.2
16	7I	51	VAL	2.2
17	8A	42	TYR	2.2
31	39	6	VAL	2.2
7	6E	32	ARG	2.2

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Mol	Chain	Res	Type	RSRZ
45	B5	33	LYS	2.2
47	H8	72	ARG	2.2
54	O8	28	ARG	2.2
9	82	47	LEU	2.2
24	3K	33	U	2.2
10	1I	74	ILE	2.2
40	65	37	ALA	2.2
17	8I	91	ARG	2.2
35	58	53	VAL	2.2
37	35	65	ARG	2.2
48	E5	55	ARG	2.2
4	32	68	TYR	2.2
11	2I	59	TYR	2.2
25	4K	15	A	2.2
44	A5	38	TYR	2.2
49	J8	62	VAL	2.2
31	31	131	GLY	2.1
34	61	68	LEU	2.1
43	95	39	LEU	2.1
43	95	75	PHE	2.1
44	E8	69	LEU	2.1
4	3E	158	ILE	2.1
10	1I	50	ILE	2.1
19	AA	75	ALA	2.1
3	22	89	GLU	2.1
7	6E	88	PRO	2.1
19	AA	55	LYS	2.1
54	O8	45	LYS	2.1
13	4A	91	ARG	2.1
3	22	33	LEU	2.1
4	3E	162	LEU	2.1
4	32	97	LEU	2.1
30	21	52	LEU	2.1
44	E8	86	LEU	2.1
45	B5	21	PHE	2.1
3	22	93	LYS	2.1
6	5E	8	ILE	2.1
8	7E	111	ILE	2.1
11	2I	69	ALA	2.1
20	BI	54	LYS	2.1
1	1G	1030	C	2.1
32	49	118	ARG	2.1

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Mol	Chain	Res	Type	RSRZ
51	H5	12	PRO	2.1
56	M5	4	MET	2.1
35	58	46	VAL	2.1
3	22	196	LEU	2.1
4	32	19	LEU	2.1
43	95	74	LYS	2.1
3	2E	21	ARG	2.1
12	3A	7	ILE	2.1
13	4A	84	ILE	2.1
15	6I	63	ARG	2.1
20	BA	15	ARG	2.1
45	B5	8	ILE	2.1
47	D5	82	ARG	2.1
5	42	66	MET	2.1
36	68	98	VAL	2.1
47	H8	121	HIS	2.1
51	L8	59	VAL	2.1
55	L5	46	VAL	2.1
32	49	105	LYS	2.1
15	6A	31	LEU	2.1
31	39	12	LEU	2.1
32	41	82	LEU	2.1
47	D5	38	TYR	2.1
51	H5	32	GLN	2.1
52	M8	23	GLU	2.1
8	7E	93	VAL	2.1
11	2I	80	VAL	2.1
47	D5	66	SER	2.1
3	2E	128	PHE	2.1
41	B8	105	LEU	2.1
42	C8	83	LEU	2.1
42	85	74	LEU	2.1
9	82	87	GLN	2.1
20	BI	90	GLN	2.1
28	71	173	ALA	2.1
7	6E	131	LYS	2.1
30	21	32	PRO	2.1
33	59	82	GLY	2.1
37	35	22	GLY	2.1
47	H8	108	PRO	2.1
3	22	76	VAL	2.1
34	61	144	VAL	2.1

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Mol	Chain	Res	Type	RSRZ
10	1I	19	SER	2.1
17	8A	38	ARG	2.1
30	29	135	HIS	2.1
5	42	45	PHE	2.1
5	42	119	LEU	2.1
17	8I	42	TYR	2.1
19	AA	77	THR	2.1
44	E8	109	GLU	2.1
48	E5	59	LEU	2.1
47	D5	154	ASP	2.1
5	42	62	ALA	2.1
9	82	55	ALA	2.1
31	39	146	ALA	2.1
49	F5	18	ILE	2.1
26	1H	2793	G	2.1
10	1I	66	ARG	2.1
11	2A	54	ARG	2.1
11	2A	109	VAL	2.1
43	D8	5	VAL	2.1
43	D8	46	VAL	2.1
47	H8	126	VAL	2.1
47	D5	80	ARG	2.1
3	2E	90	GLU	2.1
7	6E	139	GLU	2.1
16	7I	59	TRP	2.1
20	BI	16	HIS	2.1
26	14	2139	C	2.1
30	29	40	GLU	2.1
47	D5	11	GLU	2.1
35	15	15	LEU	2.1
40	65	73	LEU	2.1
40	65	110	LEU	2.1
35	58	83	LYS	2.1
36	25	53	LYS	2.1
38	45	93	TYR	2.1
3	2E	100	ALA	2.1
18	9I	20	ALA	2.1
26	14	1026	U	2.1
3	22	88	ARG	2.1
8	7E	9	MET	2.1
8	7E	37	ARG	2.1
32	41	72	ARG	2.1

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Mol	Chain	Res	Type	RSRZ
48	E5	39	ARG	2.1
4	3E	24	GLU	2.1
42	C8	90	VAL	2.1
43	95	47	VAL	2.1
16	7I	27	LYS	2.1
30	29	2	LYS	2.1
40	A8	73	LEU	2.1
8	7E	135	CYS	2.1
7	6E	54	THR	2.1
8	7E	65	TYR	2.1
11	2I	36	ASP	2.1
19	AI	79	THR	2.1
28	71	25	ALA	2.1
33	59	121	ILE	2.1
13	4A	99	ARG	2.1
17	8I	33	GLY	2.1
29	11	210	GLY	2.1
44	E8	68	ARG	2.1
44	A5	17	VAL	2.1
47	D5	178	GLU	2.1
20	BI	58	LYS	2.1
20	BA	98	PRO	2.1
33	51	39	PRO	2.1
2	12	105	PHE	2.1
36	25	99	PHE	2.1
7	62	13	GLN	2.0
3	22	22	TRP	2.0
3	2E	61	ALA	2.0
12	3I	15	ARG	2.0
21	1B	15	ARG	2.0
31	31	9	ILE	2.0
32	41	46	ALA	2.0
34	61	88	ILE	2.0
17	8I	8	GLY	2.0
26	14	2506	U	2.0
35	58	75	TYR	2.0
40	A8	72	ALA	2.0
47	D5	20	ARG	2.0
47	H8	22	GLY	2.0
47	D5	115	GLY	2.0
47	D5	145	GLU	2.0
21	1B	23	PRO	2.0

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Mol	Chain	Res	Type	RSRZ
55	P8	1	MET	2.0
28	71	165	ASN	2.0
30	29	54	GLN	2.0
1	1G	1061	G	2.0
1	1G	1535	C	2.0
4	32	49	ARG	2.0
12	3A	70	ILE	2.0
9	8E	8	GLY	2.0
20	BI	96	GLY	2.0
26	14	1536	A	2.0
16	7I	20	VAL	2.0
36	25	43	VAL	2.0
40	65	85	VAL	2.0
10	1I	37	PRO	2.0
35	15	44	PRO	2.0
38	88	99	PRO	2.0
35	58	116	LEU	2.0
47	H8	136	PHE	2.0
51	L8	23	LEU	2.0
9	82	121	ARG	2.0
4	32	16	GLY	2.0
11	2I	16	SER	2.0
33	59	138	LYS	2.0
46	G8	96	ILE	2.0
34	61	83	ALA	2.0
25	4K	14	A	2.0
13	4A	105	THR	2.0
2	1E	71	VAL	2.0
11	2A	30	VAL	2.0
20	BI	85	MET	2.0
51	H5	47	VAL	2.0
13	4A	108	ARG	2.0
30	29	53	PRO	2.0
31	39	192	LEU	2.0
39	98	10	LEU	2.0
47	D5	177	PRO	2.0
16	7I	31	LYS	2.0
11	2A	92	GLU	2.0
12	3I	29	GLY	2.0
30	29	134	ILE	2.0
34	61	79	ILE	2.0
38	45	38	GLU	2.0

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Mol	Chain	Res	Type	RSRZ
49	F5	37	ILE	2.0
5	42	105	VAL	2.0
7	6E	141	VAL	2.0
20	BI	86	ARG	2.0
28	71	8	ARG	2.0
31	31	72	ARG	2.0
33	51	170	ARG	2.0
37	35	50	ARG	2.0
3	2E	26	LYS	2.0
17	8A	30	PRO	2.0
18	9A	44	LEU	2.0
33	59	112	PRO	2.0
43	D8	44	LYS	2.0
47	H8	24	LEU	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. LLDF column lists the quality of electron density of the group with respect to its neighbouring residues in protein, DNA or RNA chains. The B-factors column lists the minimum, median, 95^{th} 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	LLDF	B-factors(Å^2)	Q<0.9
22	PSU	1K	55	20/21	0.87	0.16	-	107,127,140,145	0
23	OMC	2K	33	21/22	0.96	0.21	-	73,79,82,94	0
23	4SU	2K	8	20/21	0.94	0.17	-	89,95,101,103	0
23	7MG	2L	47	24/25	0.92	0.14	-	118,128,137,141	0
23	OMC	2L	33	21/22	0.90	0.19	-	94,103,107,109	0
22	5MU	1K	54	21/22	0.92	0.16	-	104,125,133,143	0
23	5MU	2K	55	21/22	0.95	0.14	-	98,105,112,114	0
23	4SU	2L	8	20/21	0.88	0.16	-	107,118,124,125	0
23	7MG	2K	47	24/25	0.94	0.15	-	96,101,113,116	0
22	U8U	1K	34	23/24	0.96	0.17	-	83,98,108,110	0
22	PSU	1K	39	20/21	0.93	0.21	-	94,112,118,120	0
23	PSU	2L	56	20/21	0.91	0.10	-	109,121,126,126	0
22	T6A	1K	37	32/33	0.94	0.20	-	81,92,114,116	0
23	PSU	2K	56	20/21	0.92	0.12	-	91,101,110,117	0
23	5MU	2L	55	21/22	0.94	0.14	-	115,122,127,130	0

6.3 Carbohydrates (i)

There are no carbohydrates in this entry.

6.4 Ligands (i)

In the following table, the Atoms column lists the number of modelled atoms in the group and the number defined in the chemical component dictionary. LLDF column lists the quality of electron density of the group with respect to its neighbouring residues in protein, DNA or RNA chains. The B-factors column lists the minimum, median, 95^{th} 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	LLDF	B-factors(Å^2)	Q<0.9
57	MG	1H	3205	1/1	0.62	0.60	117.16	79,79,79,79	0
57	MG	1H	3151	1/1	0.92	0.55	58.39	80,80,80,80	0
57	MG	13	1627	1/1	0.84	0.48	53.46	71,71,71,71	0
57	MG	14	3208	1/1	0.61	0.95	39.77	88,88,88,88	0
57	MG	1H	3009	1/1	0.94	0.54	36.07	71,71,71,71	0
57	MG	1G	1614	1/1	0.90	0.60	34.55	88,88,88,88	0
57	MG	14	3117	1/1	0.85	0.47	32.77	64,64,64,64	0
57	MG	1H	3130	1/1	0.70	0.47	31.50	63,63,63,63	0
57	MG	14	3136	1/1	0.95	0.46	30.42	92,92,92,92	0
57	MG	14	3064	1/1	0.65	0.48	29.88	63,63,63,63	0
57	MG	14	3016	1/1	0.96	0.38	25.60	74,74,74,74	0
57	MG	14	3203	1/1	0.89	0.53	23.30	79,79,79,79	0
57	MG	1H	3188	1/1	0.96	0.41	22.29	49,49,49,49	0
57	MG	14	3048	1/1	0.97	0.47	20.80	51,51,51,51	0
57	MG	1H	3061	1/1	0.85	0.28	19.51	53,53,53,53	0
57	MG	14	3035	1/1	0.94	0.41	19.15	58,58,58,58	0
57	MG	13	1601	1/1	0.98	0.42	18.90	80,80,80,80	0
57	MG	14	3088	1/1	0.87	0.49	18.86	87,87,87,87	0
57	MG	14	3090	1/1	0.91	0.35	17.94	86,86,86,86	0
57	MG	14	3126	1/1	0.95	0.28	17.89	72,72,72,72	0
57	MG	1H	3187	1/1	0.98	0.42	17.89	44,44,44,44	0
57	MG	1H	3041	1/1	0.94	0.28	17.49	51,51,51,51	0
57	MG	14	3046	1/1	0.94	0.45	17.39	79,79,79,79	0
57	MG	14	3142	1/1	0.80	0.29	15.82	80,80,80,80	0
57	MG	1H	3098	1/1	0.89	0.41	15.63	46,46,46,46	0
57	MG	14	3198	1/1	0.83	0.43	15.02	81,81,81,81	0
57	MG	13	1669	1/1	0.73	0.53	14.98	95,95,95,95	0
57	MG	1H	3143	1/1	0.95	0.37	14.68	50,50,50,50	0
57	MG	1H	3172	1/1	0.73	0.30	14.52	83,83,83,83	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	1H	3079	1/1	0.66	0.34	14.30	63,63,63,63	0
57	MG	1H	3015	1/1	0.98	0.31	13.81	51,51,51,51	0
57	MG	13	1654	1/1	0.85	0.41	13.77	101,101,101,101	0
57	MG	1H	3165	1/1	0.88	0.29	13.72	64,64,64,64	0
57	MG	1H	3078	1/1	0.92	0.28	13.54	58,58,58,58	0
57	MG	14	3072	1/1	0.82	0.27	13.49	94,94,94,94	0
57	MG	1H	3074	1/1	0.81	0.38	13.21	64,64,64,64	0
57	MG	13	1628	1/1	0.82	0.32	12.83	91,91,91,91	0
57	MG	13	1624	1/1	0.95	0.32	12.09	59,59,59,59	0
57	MG	14	3205	1/1	0.94	0.45	12.03	59,59,59,59	0
57	MG	14	3034	1/1	0.98	0.49	11.19	40,40,40,40	0
57	MG	13	1614	1/1	0.96	0.34	10.95	84,84,84,84	0
57	MG	14	3118	1/1	0.74	0.44	10.73	61,61,61,61	0
57	MG	1H	3076	1/1	0.93	0.24	10.25	72,72,72,72	0
57	MG	1G	1636	1/1	0.80	0.33	10.22	90,90,90,90	0
57	MG	1H	3066	1/1	0.83	0.26	10.06	60,60,60,60	0
57	MG	1H	3029	1/1	0.96	0.32	10.02	77,77,77,77	0
57	MG	1H	3056	1/1	0.97	0.31	9.95	49,49,49,49	0
57	MG	14	3122	1/1	0.67	0.37	9.89	75,75,75,75	0
57	MG	14	3030	1/1	0.95	0.36	9.78	57,57,57,57	0
57	MG	1H	3321	1/1	0.90	0.46	9.62	86,86,86,86	0
57	MG	1H	3178	1/1	0.84	0.43	9.48	63,63,63,63	0
57	MG	1H	3181	1/1	0.89	0.24	9.42	65,65,65,65	0
57	MG	19	301	1/1	0.79	0.42	9.18	48,48,48,48	0
57	MG	1H	3105	1/1	0.78	0.23	8.86	72,72,72,72	0
57	MG	1H	3186	1/1	0.97	0.26	8.62	62,62,62,62	0
57	MG	14	3134	1/1	0.78	0.33	8.45	85,85,85,85	0
57	MG	14	3179	1/1	0.95	0.32	8.43	77,77,77,77	0
57	MG	13	1668	1/1	0.70	0.29	8.26	66,66,66,66	0
57	MG	1H	3175	1/1	0.94	0.29	8.15	64,64,64,64	0
61	SPE	14	3437	13/13	0.82	0.29	8.02	59,68,73,74	0
57	MG	1H	3040	1/1	0.97	0.31	7.97	44,44,44,44	0
57	MG	14	3114	1/1	0.64	0.22	7.95	70,70,70,70	0
57	MG	1H	3108	1/1	0.87	0.24	7.76	64,64,64,64	0
57	MG	13	1629	1/1	0.89	0.21	7.61	72,72,72,72	0
57	MG	14	3181	1/1	0.73	0.30	7.46	75,75,75,75	0
57	MG	16	202	1/1	0.89	0.23	7.44	82,82,82,82	0
57	MG	13	1670	1/1	0.83	0.30	7.44	82,82,82,82	0
57	MG	14	3124	1/1	0.93	0.18	7.39	97,97,97,97	0
57	MG	14	3089	1/1	0.96	0.24	7.31	54,54,54,54	0
57	MG	1H	3045	1/1	0.98	0.28	6.97	46,46,46,46	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	14	3107	1/1	0.84	0.36	6.81	54,54,54,54	0
57	MG	13	1645	1/1	0.91	0.28	6.80	92,92,92,92	0
57	MG	1H	3203	1/1	0.78	0.24	6.70	74,74,74,74	0
57	MG	14	3108	1/1	0.80	0.28	6.49	72,72,72,72	0
57	MG	14	3171	1/1	0.97	0.34	6.48	56,56,56,56	0
57	MG	16	203	1/1	0.75	0.21	6.41	82,82,82,82	0
57	MG	2L	101	1/1	0.93	0.40	6.35	91,91,91,91	0
57	MG	14	3248	1/1	0.95	0.29	6.28	47,47,47,47	0
57	MG	1H	3085	1/1	0.69	0.28	6.26	70,70,70,70	0
57	MG	14	3031	1/1	0.96	0.28	6.17	73,73,73,73	0
57	MG	1H	3200	1/1	0.95	0.26	6.13	71,71,71,71	0
57	MG	1H	3212	1/1	0.88	0.27	6.11	65,65,65,65	0
57	MG	14	3151	1/1	0.74	0.29	6.10	70,70,70,70	0
57	MG	1H	3121	1/1	0.99	0.24	5.84	39,39,39,39	0
57	MG	14	3099	1/1	0.84	0.18	5.82	78,78,78,78	0
57	MG	14	3097	1/1	0.94	0.26	5.65	97,97,97,97	0
57	MG	14	3023	1/1	0.73	0.27	5.52	76,76,76,76	0
57	MG	14	3028	1/1	0.95	0.33	5.41	45,45,45,45	0
57	MG	1H	3167	1/1	0.92	0.53	5.31	98,98,98,98	0
57	MG	13	1636	1/1	0.95	0.28	5.29	74,74,74,74	0
57	MG	14	3054	1/1	0.94	0.25	5.13	79,79,79,79	0
57	MG	1H	3011	1/1	0.97	0.30	5.08	56,56,56,56	0
57	MG	14	3125	1/1	0.87	0.21	5.02	81,81,81,81	0
57	MG	14	3092	1/1	0.95	0.26	5.01	66,66,66,66	0
57	MG	1H	3102	1/1	0.92	0.21	4.89	60,60,60,60	0
57	MG	1H	3042	1/1	0.96	0.26	4.75	65,65,65,65	0
61	SPE	14	3436	13/13	0.75	0.25	4.68	83,88,93,94	0
57	MG	1H	3221	1/1	0.89	0.24	4.55	73,73,73,73	0
57	MG	14	3042	1/1	0.90	0.19	4.49	65,65,65,65	0
57	MG	1H	3082	1/1	0.88	0.28	4.48	71,71,71,71	0
57	MG	14	3145	1/1	0.94	0.21	4.44	90,90,90,90	0
57	MG	1H	3002	1/1	0.86	0.21	4.41	48,48,48,48	0
57	MG	14	3059	1/1	0.96	0.23	4.36	89,89,89,89	0
57	MG	14	3007	1/1	0.80	0.26	4.20	60,60,60,60	0
57	MG	13	1633	1/1	0.94	0.22	4.15	73,73,73,73	0
57	MG	14	3100	1/1	0.76	0.27	4.08	71,71,71,71	0
57	MG	14	3062	1/1	0.95	0.25	4.04	59,59,59,59	0
57	MG	14	3070	1/1	0.89	0.22	4.03	52,52,52,52	0
57	MG	1H	3264	1/1	0.97	0.25	3.98	50,50,50,50	0
57	MG	1H	3133	1/1	0.86	0.23	3.81	55,55,55,55	0
57	MG	1H	3034	1/1	0.92	0.21	3.56	46,46,46,46	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	14	3139	1/1	0.78	0.31	3.47	68,68,68,68	0
57	MG	14	3074	1/1	0.81	0.27	3.39	69,69,69,69	0
57	MG	14	3015	1/1	0.91	0.29	3.35	63,63,63,63	0
57	MG	1H	3235	1/1	0.86	0.20	3.32	71,71,71,71	0
57	MG	14	3224	1/1	0.92	0.24	3.12	45,45,45,45	0
57	MG	1H	3140	1/1	0.84	0.20	3.10	54,54,54,54	0
57	MG	14	3077	1/1	0.86	0.22	3.09	52,52,52,52	0
57	MG	14	3076	1/1	0.87	0.45	3.09	81,81,81,81	0
57	MG	14	3233	1/1	0.93	0.26	2.99	65,65,65,65	0
57	MG	14	3236	1/1	0.88	0.28	2.99	44,44,44,44	0
57	MG	14	3008	1/1	0.60	0.21	2.91	77,77,77,77	0
57	MG	14	3061	1/1	0.85	0.31	2.91	61,61,61,61	0
57	MG	13	1635	1/1	0.88	0.24	2.90	81,81,81,81	0
57	MG	13	1659	1/1	0.85	0.15	2.84	126,126,126,126	0
57	MG	13	1661	1/1	0.78	0.19	2.69	90,90,90,90	0
57	MG	14	3014	1/1	0.97	0.25	2.67	59,59,59,59	0
57	MG	14	3080	1/1	0.76	0.25	2.58	68,68,68,68	0
57	MG	1H	3013	1/1	0.90	0.27	2.50	40,40,40,40	0
57	MG	1H	3067	1/1	0.93	0.21	2.50	58,58,58,58	0
57	MG	14	3194	1/1	0.96	0.22	2.43	80,80,80,80	0
57	MG	14	3191	1/1	0.98	0.27	2.28	67,67,67,67	0
57	MG	14	3119	1/1	0.90	0.20	2.24	81,81,81,81	0
57	MG	1H	3059	1/1	0.99	0.27	2.20	56,56,56,56	0
57	MG	13	1623	1/1	0.96	0.22	2.14	76,76,76,76	0
57	MG	14	3237	1/1	0.91	0.26	2.08	51,51,51,51	0
57	MG	1H	3193	1/1	0.91	0.26	1.97	61,61,61,61	0
57	MG	1H	3275	1/1	0.96	0.18	1.95	57,57,57,57	0
57	MG	1H	3080	1/1	0.86	0.17	1.83	80,80,80,80	0
58	PAR	13	1741	42/42	0.95	0.24	1.83	64,71,75,82	0
57	MG	13	1607	1/1	0.99	0.20	1.81	72,72,72,72	0
57	MG	13	1646	1/1	0.95	0.17	1.58	71,71,71,71	0
57	MG	4I	200	1/1	0.70	0.23	1.54	95,95,95,95	0
57	MG	13	1641	1/1	0.90	0.19	1.50	81,81,81,81	0
57	MG	14	3086	1/1	0.97	0.23	1.45	56,56,56,56	0
57	MG	1H	3022	1/1	0.96	0.19	1.43	49,49,49,49	0
57	MG	14	3231	1/1	0.97	0.23	1.41	55,55,55,55	0
57	MG	14	3040	1/1	0.98	0.25	1.24	68,68,68,68	0
57	MG	14	3235	1/1	0.98	0.24	1.16	48,48,48,48	0
57	MG	1H	3090	1/1	0.97	0.21	1.08	42,42,42,42	0
57	MG	14	3041	1/1	0.92	0.20	0.98	55,55,55,55	0
57	MG	14	3290	1/1	0.95	0.22	0.97	59,59,59,59	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	1H	3092	1/1	0.92	0.24	0.93	61,61,61,61	0
57	MG	1H	3134	1/1	0.83	0.14	0.89	57,57,57,57	0
57	MG	13	1692	1/1	0.96	0.18	0.85	72,72,72,72	0
58	PAR	1G	1702	42/42	0.94	0.22	0.77	83,92,100,103	0
57	MG	14	3288	1/1	0.95	0.20	0.73	49,49,49,49	0
57	MG	1H	3417	1/1	0.92	0.18	0.72	83,83,83,83	0
57	MG	1G	1622	1/1	0.98	0.20	0.71	125,125,125,125	0
57	MG	1H	3117	1/1	0.91	0.18	0.70	60,60,60,60	0
57	MG	14	3087	1/1	0.95	0.27	0.66	76,76,76,76	0
57	MG	1G	1633	1/1	0.98	0.21	0.56	118,118,118,118	0
57	MG	14	3005	1/1	0.84	0.26	0.54	54,54,54,54	0
57	MG	1H	3170	1/1	0.79	0.34	0.46	80,80,80,80	0
57	MG	13	1606	1/1	0.97	0.25	0.42	72,72,72,72	0
57	MG	1G	1644	1/1	0.74	0.20	0.42	106,106,106,106	0
57	MG	14	3216	1/1	0.98	0.21	0.36	52,52,52,52	0
57	MG	42	201	1/1	0.90	0.21	0.35	120,120,120,120	0
57	MG	14	3045	1/1	0.97	0.20	0.22	53,53,53,53	0
57	MG	14	3111	1/1	0.85	0.19	0.22	58,58,58,58	0
57	MG	1H	3300	1/1	0.93	0.19	0.21	62,62,62,62	0
57	MG	13	1642	1/1	0.94	0.13	0.19	103,103,103,103	0
57	MG	1G	1649	1/1	0.88	0.14	0.19	113,113,113,113	0
57	MG	1G	1607	1/1	0.98	0.20	0.18	86,86,86,86	0
57	MG	14	3217	1/1	0.96	0.20	0.10	56,56,56,56	0
57	MG	1H	3208	1/1	0.96	0.15	0.10	72,72,72,72	0
57	MG	1H	3294	1/1	0.90	0.16	0.06	54,54,54,54	0
57	MG	1H	3114	1/1	0.94	0.15	0.00	71,71,71,71	0
57	MG	1H	3039	1/1	0.95	0.20	-0.04	33,33,33,33	0
57	MG	1H	3027	1/1	0.97	0.19	-0.05	52,52,52,52	0
57	MG	1H	3525	1/1	0.84	0.22	-0.06	68,68,68,68	0
57	MG	1H	3397	1/1	0.92	0.19	-0.11	51,51,51,51	0
57	MG	1H	3365	1/1	0.94	0.16	-0.13	67,67,67,67	0
57	MG	13	1648	1/1	0.98	0.16	-0.14	67,67,67,67	0
57	MG	1H	3021	1/1	0.94	0.17	-0.16	49,49,49,49	0
57	MG	1H	3323	1/1	0.96	0.17	-0.19	57,57,57,57	0
57	MG	1G	1615	1/1	0.87	0.21	-0.21	86,86,86,86	0
57	MG	14	3029	1/1	0.94	0.21	-0.22	57,57,57,57	0
57	MG	52	201	1/1	0.79	0.16	-0.23	129,129,129,129	0
57	MG	14	3155	1/1	0.82	0.16	-0.24	78,78,78,78	0
57	MG	1H	3259	1/1	0.94	0.17	-0.28	54,54,54,54	0
57	MG	1H	3313	1/1	0.94	0.17	-0.30	47,47,47,47	0
57	MG	1G	1685	1/1	0.93	0.14	-0.31	110,110,110,110	0
59	SF4	3E	301	8/8	0.99	0.21	-0.33	82,91,95,98	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	1H	3433	1/1	0.85	0.14	-0.34	72,72,72,72	0
57	MG	1G	1704	1/1	0.80	0.20	-0.35	84,84,84,84	0
57	MG	1G	1604	1/1	0.91	0.17	-0.35	127,127,127,127	0
57	MG	1H	3263	1/1	0.96	0.19	-0.36	46,46,46,46	0
57	MG	14	3390	1/1	0.87	0.22	-0.41	94,94,94,94	0
57	MG	88	201	1/1	0.93	0.19	-0.45	76,76,76,76	0
57	MG	13	1672	1/1	0.91	0.15	-0.47	88,88,88,88	0
57	MG	13	1657	1/1	0.90	0.15	-0.47	68,68,68,68	0
57	MG	13	1602	1/1	0.97	0.14	-0.52	120,120,120,120	0
57	MG	16	206	1/1	0.85	0.12	-0.52	77,77,77,77	0
57	MG	39	301	1/1	0.91	0.18	-0.61	89,89,89,89	0
57	MG	13	1700	1/1	0.88	0.12	-0.63	102,102,102,102	0
57	MG	1H	3215	1/1	0.89	0.13	-0.63	63,63,63,63	0
57	MG	14	3081	1/1	0.97	0.15	-0.64	56,56,56,56	0
59	SF4	32	302	8/8	0.99	0.18	-0.72	100,109,118,124	0
57	MG	14	3105	1/1	0.94	0.14	-0.75	73,73,73,73	0
57	MG	1H	3415	1/1	0.75	0.14	-0.77	72,72,72,72	0
57	MG	1H	3285	1/1	0.94	0.14	-0.80	55,55,55,55	0
57	MG	14	3395	1/1	0.67	0.10	-0.82	118,118,118,118	0
57	MG	14	3261	1/1	0.96	0.17	-0.84	59,59,59,59	0
57	MG	14	3027	1/1	0.96	0.14	-0.86	66,66,66,66	0
57	MG	1H	3020	1/1	0.82	0.14	-0.93	73,73,73,73	0
57	MG	1G	1678	1/1	0.58	0.11	-0.96	111,111,111,111	0
57	MG	13	1663	1/1	0.97	0.10	-0.99	107,107,107,107	0
57	MG	14	3113	1/1	0.92	0.18	-1.00	52,52,52,52	0
57	MG	1H	3273	1/1	0.97	0.14	-1.02	78,78,78,78	0
60	ZN	5I	102	1/1	0.99	0.13	-1.02	86,86,86,86	0
57	MG	1H	3297	1/1	0.96	0.14	-1.02	49,49,49,49	0
57	MG	13	1612	1/1	0.92	0.12	-1.05	86,86,86,86	0
57	MG	1H	3210	1/1	0.98	0.18	-1.05	56,56,56,56	0
57	MG	1H	3303	1/1	0.90	0.17	-1.05	48,48,48,48	0
57	MG	14	3268	1/1	0.87	0.15	-1.07	82,82,82,82	0
57	MG	1G	1638	1/1	0.95	0.14	-1.11	127,127,127,127	0
57	MG	13	1618	1/1	0.90	0.20	-1.13	48,48,48,48	0
57	MG	1G	1684	1/1	0.94	0.06	-1.16	96,96,96,96	0
57	MG	1G	1603	1/1	0.96	0.14	-1.16	88,88,88,88	0
57	MG	1H	3279	1/1	0.94	0.15	-1.17	48,48,48,48	0
57	MG	14	3249	1/1	0.93	0.15	-1.18	52,52,52,52	0
57	MG	1H	3262	1/1	0.97	0.18	-1.20	43,43,43,43	0
57	MG	1G	1666	1/1	0.94	0.14	-1.20	100,100,100,100	0
57	MG	14	3310	1/1	0.96	0.15	-1.22	54,54,54,54	0
57	MG	1H	3055	1/1	0.81	0.18	-1.25	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	1H	3023	1/1	0.92	0.17	-1.26	50,50,50,50	0
57	MG	45	201	1/1	0.95	0.13	-1.28	101,101,101,101	0
57	MG	13	1691	1/1	0.94	0.12	-1.31	92,92,92,92	0
57	MG	1H	3492	1/1	0.86	0.15	-1.32	78,78,78,78	0
57	MG	14	3083	1/1	0.91	0.15	-1.33	72,72,72,72	0
57	MG	41	201	1/1	0.81	0.12	-1.35	69,69,69,69	0
57	MG	1H	3047	1/1	0.95	0.15	-1.37	55,55,55,55	0
57	MG	21	301	1/1	0.94	0.14	-1.38	59,59,59,59	0
57	MG	1H	3386	1/1	0.81	0.18	-1.42	42,42,42,42	0
60	ZN	5A	101	1/1	0.98	0.09	-1.43	146,146,146,146	0
57	MG	13	1680	1/1	0.96	0.16	-1.45	64,64,64,64	0
57	MG	1H	3116	1/1	0.91	0.16	-1.46	53,53,53,53	0
57	MG	13	1733	1/1	0.85	0.12	-1.46	101,101,101,101	0
57	MG	14	3378	1/1	0.93	0.13	-1.49	92,92,92,92	0
57	MG	14	3238	1/1	0.94	0.20	-1.50	50,50,50,50	0
57	MG	1H	3335	1/1	0.96	0.14	-1.51	48,48,48,48	0
57	MG	1H	3036	1/1	0.97	0.16	-1.56	37,37,37,37	0
57	MG	1H	3312	1/1	0.92	0.15	-1.57	44,44,44,44	0
57	MG	13	1682	1/1	0.92	0.11	-1.58	93,93,93,93	0
57	MG	14	3349	1/1	0.90	0.13	-1.59	83,83,83,83	0
57	MG	13	1631	1/1	0.90	0.09	-1.66	78,78,78,78	0
57	MG	14	3156	1/1	0.77	0.17	-1.67	87,87,87,87	0
57	MG	1H	3509	1/1	0.89	0.12	-1.69	47,47,47,47	0
57	MG	1H	3281	1/1	0.98	0.15	-1.71	61,61,61,61	0
57	MG	14	3230	1/1	0.83	0.17	-1.72	63,63,63,63	0
57	MG	14	3025	1/1	0.86	0.12	-1.74	80,80,80,80	0
57	MG	1H	3298	1/1	0.83	0.11	-1.75	51,51,51,51	0
57	MG	1H	3305	1/1	0.96	0.14	-1.76	63,63,63,63	0
57	MG	1H	3336	1/1	0.90	0.14	-1.81	62,62,62,62	0
57	MG	1H	3228	1/1	0.91	0.15	-1.81	60,60,60,60	0
57	MG	1G	1619	1/1	0.91	0.10	-1.82	102,102,102,102	0
57	MG	1G	1617	1/1	0.95	0.12	-1.85	116,116,116,116	0
57	MG	14	3287	1/1	0.96	0.11	-1.86	88,88,88,88	0
57	MG	14	3254	1/1	0.86	0.10	-1.87	77,77,77,77	0
57	MG	5I	101	1/1	0.83	0.10	-1.89	79,79,79,79	0
57	MG	14	3245	1/1	0.90	0.14	-1.97	63,63,63,63	0
57	MG	14	3121	1/1	0.90	0.12	-2.00	61,61,61,61	0
57	MG	14	3270	1/1	0.97	0.13	-2.01	67,67,67,67	0
57	MG	14	3272	1/1	0.95	0.09	-2.03	71,71,71,71	0
57	MG	14	3020	1/1	0.92	0.12	-2.06	75,75,75,75	0
57	MG	1H	3155	1/1	0.93	0.15	-2.12	55,55,55,55	0
57	MG	14	3295	1/1	0.82	0.10	-2.13	60,60,60,60	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	14	3262	1/1	0.82	0.14	-2.15	83,83,83,83	0
57	MG	1H	3222	1/1	0.82	0.14	-2.19	69,69,69,69	0
61	SPE	1G	1703	13/13	0.88	0.09	-2.22	90,107,113,114	0
57	MG	14	3387	1/1	0.67	0.14	-2.22	83,83,83,83	0
60	ZN	C5	202	1/1	0.85	0.06	-2.22	165,165,165,165	0
57	MG	M5	101	1/1	0.93	0.18	-2.23	88,88,88,88	0
57	MG	Q8	300	1/1	0.92	0.12	-2.24	80,80,80,80	0
57	MG	14	3018	1/1	0.94	0.14	-2.28	44,44,44,44	0
57	MG	14	3150	1/1	0.94	0.12	-2.36	63,63,63,63	0
57	MG	1G	1667	1/1	0.86	0.11	-2.39	103,103,103,103	0
57	MG	1H	3325	1/1	0.99	0.15	-2.40	46,46,46,46	0
57	MG	1H	3491	1/1	0.94	0.10	-2.42	70,70,70,70	0
57	MG	13	1683	1/1	0.87	0.05	-2.43	93,93,93,93	0
57	MG	1H	3267	1/1	0.73	0.10	-2.44	56,56,56,56	0
57	MG	13	1714	1/1	0.74	0.09	-2.44	122,122,122,122	0
57	MG	78	201	1/1	0.86	0.10	-2.45	66,66,66,66	0
57	MG	14	3321	1/1	0.94	0.08	-2.48	69,69,69,69	0
57	MG	14	3309	1/1	0.91	0.14	-2.49	55,55,55,55	0
57	MG	1H	3194	1/1	0.85	0.10	-2.52	63,63,63,63	0
57	MG	1H	3257	1/1	0.95	0.14	-2.56	43,43,43,43	0
57	MG	14	3373	1/1	0.86	0.12	-2.57	65,65,65,65	0
57	MG	14	3293	1/1	0.97	0.10	-2.60	63,63,63,63	0
57	MG	1H	3469	1/1	0.94	0.15	-2.61	58,58,58,58	0
57	MG	1H	3094	1/1	0.92	0.15	-2.64	33,33,33,33	0
57	MG	14	3322	1/1	0.98	0.14	-2.67	58,58,58,58	0
57	MG	14	3211	1/1	0.74	0.09	-2.69	81,81,81,81	0
57	MG	14	3410	1/1	0.93	0.06	-2.71	85,85,85,85	0
57	MG	14	3296	1/1	0.90	0.12	-2.72	72,72,72,72	0
57	MG	14	3056	1/1	0.99	0.11	-2.73	54,54,54,54	0
57	MG	1H	3107	1/1	0.94	0.12	-2.73	60,60,60,60	0
57	MG	1H	3400	1/1	0.96	0.12	-2.77	53,53,53,53	0
57	MG	1G	1647	1/1	0.85	0.10	-2.79	97,97,97,97	0
57	MG	14	3159	1/1	0.88	0.11	-2.81	79,79,79,79	0
57	MG	1H	3138	1/1	0.91	0.14	-2.84	54,54,54,54	0
57	MG	14	3427	1/1	0.84	0.08	-2.88	65,65,65,65	0
57	MG	14	3116	1/1	0.96	0.13	-3.03	88,88,88,88	0
57	MG	1G	1632	1/1	0.79	0.09	-3.04	91,91,91,91	0
57	MG	1H	3051	1/1	0.92	0.11	-3.04	60,60,60,60	0
57	MG	13	1698	1/1	0.92	0.11	-3.08	83,83,83,83	0
57	MG	14	3354	1/1	0.90	0.14	-3.09	60,60,60,60	0
57	MG	1H	3381	1/1	0.95	0.11	-3.11	74,74,74,74	0
57	MG	1H	3046	1/1	0.86	0.13	-3.12	33,33,33,33	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	1G	1672	1/1	0.97	0.12	-3.17	110,110,110,110	0
57	MG	1H	3361	1/1	0.97	0.07	-3.17	52,52,52,52	0
57	MG	1H	3278	1/1	0.98	0.09	-3.18	75,75,75,75	0
57	MG	1H	3115	1/1	0.90	0.10	-3.19	61,61,61,61	0
57	MG	1H	3331	1/1	0.90	0.12	-3.19	70,70,70,70	0
57	MG	14	3302	1/1	0.92	0.11	-3.29	58,58,58,58	0
57	MG	14	3283	1/1	0.95	0.07	-3.31	68,68,68,68	0
57	MG	1H	3431	1/1	0.82	0.09	-3.33	98,98,98,98	0
57	MG	1H	3307	1/1	0.96	0.14	-3.37	44,44,44,44	0
57	MG	13	1703	1/1	0.95	0.13	-3.44	70,70,70,70	0
57	MG	1H	3269	1/1	0.95	0.16	-3.46	41,41,41,41	0
57	MG	14	3324	1/1	0.96	0.15	-3.46	48,48,48,48	0
57	MG	14	3282	1/1	0.89	0.06	-3.54	90,90,90,90	0
57	MG	1H	3369	1/1	0.96	0.08	-3.67	44,44,44,44	0
57	MG	1H	3031	1/1	0.99	0.14	-3.67	51,51,51,51	0
57	MG	1H	3028	1/1	0.93	0.10	-3.69	67,67,67,67	0
57	MG	1H	3280	1/1	0.98	0.15	-3.72	36,36,36,36	0
57	MG	14	3251	1/1	0.91	0.09	-3.72	80,80,80,80	0
57	MG	1H	3371	1/1	0.89	0.10	-3.74	56,56,56,56	0
57	MG	14	3280	1/1	0.96	0.14	-3.89	61,61,61,61	0
57	MG	13	1718	1/1	0.94	0.09	-3.90	90,90,90,90	0
57	MG	14	3229	1/1	0.98	0.13	-3.93	59,59,59,59	0
57	MG	1H	3355	1/1	0.95	0.09	-4.11	59,59,59,59	0
57	MG	1H	3364	1/1	0.97	0.09	-4.12	44,44,44,44	0
57	MG	14	3220	1/1	0.93	0.12	-4.17	54,54,54,54	0
57	MG	14	3307	1/1	0.90	0.15	-4.21	66,66,66,66	0
57	MG	1H	3356	1/1	0.94	0.11	-4.21	54,54,54,54	0
57	MG	14	3275	1/1	0.87	0.10	-4.31	65,65,65,65	0
57	MG	14	3091	1/1	0.81	0.09	-4.33	78,78,78,78	0
57	MG	13	1617	1/1	0.88	0.14	-4.45	51,51,51,51	0
57	MG	14	3315	1/1	0.95	0.12	-4.48	68,68,68,68	0
57	MG	1H	3252	1/1	0.98	0.09	-4.48	65,65,65,65	0
57	MG	1H	3199	1/1	0.96	0.12	-4.48	61,61,61,61	0
57	MG	1H	3427	1/1	0.91	0.10	-4.49	67,67,67,67	0
57	MG	1H	3330	1/1	0.95	0.10	-4.51	71,71,71,71	0
57	MG	1G	1646	1/1	0.88	0.10	-4.54	86,86,86,86	0
57	MG	14	3253	1/1	0.97	0.07	-4.55	58,58,58,58	0
57	MG	1H	3299	1/1	0.92	0.13	-4.55	48,48,48,48	0
57	MG	16	204	1/1	0.88	0.06	-4.64	59,59,59,59	0
57	MG	1H	3304	1/1	0.97	0.10	-4.72	58,58,58,58	0
57	MG	14	3244	1/1	0.94	0.15	-4.76	54,54,54,54	0
57	MG	1H	3093	1/1	0.91	0.09	-4.77	56,56,56,56	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	1H	3477	1/1	0.87	0.14	-4.78	83,83,83,83	0
57	MG	1H	3104	1/1	0.91	0.13	-4.85	43,43,43,43	0
57	MG	14	3289	1/1	0.92	0.13	-4.88	74,74,74,74	0
57	MG	1H	3239	1/1	0.94	0.10	-4.99	67,67,67,67	0
57	MG	1H	3302	1/1	0.98	0.13	-5.00	51,51,51,51	0
57	MG	1H	3437	1/1	0.97	0.09	-5.01	92,92,92,92	0
57	MG	14	3305	1/1	0.98	0.12	-5.04	49,49,49,49	0
57	MG	1G	1657	1/1	0.84	0.06	-5.17	121,121,121,121	0
57	MG	1G	1652	1/1	0.91	0.15	-5.19	81,81,81,81	0
57	MG	14	3242	1/1	0.99	0.10	-5.21	62,62,62,62	0
57	MG	1G	1651	1/1	0.97	0.11	-5.22	76,76,76,76	0
57	MG	1H	3392	1/1	0.86	0.15	-5.23	48,48,48,48	0
57	MG	1G	1606	1/1	0.85	0.13	-5.28	73,73,73,73	0
57	MG	1H	3470	1/1	0.81	0.08	-5.31	83,83,83,83	0
57	MG	1G	1645	1/1	0.93	0.14	-5.32	81,81,81,81	0
57	MG	1H	3048	1/1	0.93	0.13	-5.40	45,45,45,45	0
57	MG	1H	3272	1/1	0.96	0.09	-5.47	63,63,63,63	0
57	MG	14	3250	1/1	0.93	0.08	-5.56	79,79,79,79	0
57	MG	14	3388	1/1	0.89	0.07	-5.60	85,85,85,85	0
57	MG	1H	3248	1/1	0.96	0.11	-5.67	44,44,44,44	0
57	MG	1H	3465	1/1	0.96	0.09	-5.76	67,67,67,67	0
57	MG	14	3247	1/1	0.95	0.09	-5.89	52,52,52,52	0
57	MG	13	1621	1/1	0.98	0.09	-6.11	85,85,85,85	0
57	MG	14	3392	1/1	0.91	0.07	-6.22	63,63,63,63	0
57	MG	1H	3290	1/1	0.96	0.12	-6.25	55,55,55,55	0
57	MG	14	3389	1/1	0.84	0.07	-6.44	93,93,93,93	0
57	MG	14	3240	1/1	0.96	0.11	-6.57	73,73,73,73	0
57	MG	1H	3334	1/1	0.90	0.10	-6.58	77,77,77,77	0
57	MG	14	3218	1/1	0.94	0.09	-6.61	65,65,65,65	0
57	MG	14	3024	1/1	0.89	0.07	-6.63	81,81,81,81	0
57	MG	14	3317	1/1	0.93	0.08	-6.66	82,82,82,82	0
57	MG	14	3380	1/1	0.85	0.08	-6.77	78,78,78,78	0
57	MG	14	3400	1/1	0.92	0.05	-6.92	108,108,108,108	0
57	MG	1H	3265	1/1	0.97	0.08	-7.32	46,46,46,46	0
57	MG	13	1689	1/1	0.96	0.07	-7.41	88,88,88,88	0
57	MG	1H	3246	1/1	0.89	0.09	-7.44	55,55,55,55	0
57	MG	1H	3398	1/1	0.98	0.07	-7.55	78,78,78,78	0
57	MG	1H	3387	1/1	0.96	0.10	-7.68	33,33,33,33	0
57	MG	1J	204	1/1	0.89	0.09	-7.69	89,89,89,89	0
57	MG	14	3375	1/1	0.97	0.07	-7.70	93,93,93,93	0
57	MG	1H	3450	1/1	1.00	0.06	-7.73	44,44,44,44	0
57	MG	1H	3474	1/1	0.94	0.09	-8.12	93,93,93,93	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	1H	3245	1/1	0.94	0.11	-8.20	47,47,47,47	0
57	MG	14	3368	1/1	0.83	0.09	-8.36	88,88,88,88	0
57	MG	14	3267	1/1	0.84	0.06	-8.46	82,82,82,82	0
57	MG	1H	3466	1/1	0.97	0.05	-8.55	72,72,72,72	0
57	MG	1H	3375	1/1	0.87	0.10	-8.87	79,79,79,79	0
57	MG	13	1715	1/1	0.94	0.06	-9.54	76,76,76,76	0
57	MG	1H	3289	1/1	0.95	0.12	-9.55	59,59,59,59	0
57	MG	1H	3393	1/1	0.93	0.11	-9.98	50,50,50,50	0
57	MG	1H	3378	1/1	0.93	0.09	-10.02	50,50,50,50	0
57	MG	1H	3461	1/1	0.96	0.06	-10.21	45,45,45,45	0
57	MG	1H	3284	1/1	0.95	0.07	-10.66	83,83,83,83	0
57	MG	14	3065	1/1	0.95	0.10	-10.76	47,47,47,47	0
57	MG	1H	3421	1/1	0.80	0.06	-12.06	82,82,82,82	0
57	MG	13	1638	1/1	0.92	0.07	-12.51	85,85,85,85	0
57	MG	1H	3418	1/1	0.92	0.10	-12.61	46,46,46,46	0
57	MG	13	1732	1/1	0.84	0.08	-12.70	114,114,114,114	0
57	MG	14	3225	1/1	0.83	0.07	-12.89	73,73,73,73	0
57	MG	1H	3064	1/1	0.92	0.07	-12.92	51,51,51,51	0
57	MG	1H	3485	1/1	0.83	0.06	-15.92	77,77,77,77	0
57	MG	13	1701	1/1	0.91	0.12	-	108,108,108,108	0
57	MG	14	3022	1/1	0.90	0.13	-	81,81,81,81	0
57	MG	1G	1689	1/1	0.91	0.06	-	116,116,116,116	0
57	MG	1J	202	1/1	0.86	0.28	-	92,92,92,92	0
57	MG	1H	3156	1/1	0.83	0.15	-	83,83,83,83	0
57	MG	1G	1668	1/1	0.74	0.09	-	124,124,124,124	0
57	MG	1H	3318	1/1	0.96	0.15	-	59,59,59,59	0
57	MG	1H	3254	1/1	0.98	0.08	-	46,46,46,46	0
57	MG	1H	3128	1/1	0.85	0.42	-	85,85,85,85	0
57	MG	1H	3410	1/1	0.88	0.10	-	83,83,83,83	0
57	MG	14	3279	1/1	0.90	0.08	-	58,58,58,58	0
57	MG	14	3102	1/1	0.96	0.56	-	85,85,85,85	0
57	MG	14	3415	1/1	0.93	0.15	-	103,103,103,103	0
57	MG	1H	3301	1/1	0.91	0.17	-	45,45,45,45	0
57	MG	14	3271	1/1	0.80	0.06	-	94,94,94,94	0
57	MG	1H	3489	1/1	0.85	0.32	-	73,73,73,73	0
57	MG	14	3146	1/1	0.80	0.28	-	75,75,75,75	0
57	MG	14	3342	1/1	0.78	0.09	-	113,113,113,113	0
57	MG	1H	3024	1/1	0.96	0.24	-	83,83,83,83	0
57	MG	1H	3050	1/1	0.59	0.38	-	72,72,72,72	0
57	MG	1H	3166	1/1	0.96	0.43	-	78,78,78,78	0
57	MG	1H	3348	1/1	0.87	0.10	-	111,111,111,111	0
57	MG	14	3206	1/1	0.98	0.27	-	85,85,85,85	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	14	3384	1/1	0.63	0.14	-	78,78,78,78	0
57	MG	14	3053	1/1	0.95	0.07	-	95,95,95,95	0
57	MG	14	3416	1/1	0.87	0.13	-	102,102,102,102	0
57	MG	1H	3004	1/1	0.67	0.50	-	89,89,89,89	0
57	MG	14	3311	1/1	0.93	0.11	-	67,67,67,67	0
57	MG	1H	3270	1/1	0.91	0.15	-	50,50,50,50	0
57	MG	1H	3380	1/1	0.98	0.18	-	55,55,55,55	0
57	MG	13	1603	1/1	0.98	0.24	-	74,74,74,74	0
57	MG	1H	3382	1/1	0.97	0.09	-	49,49,49,49	0
57	MG	1H	3484	1/1	0.90	0.09	-	105,105,105,105	0
57	MG	14	3137	1/1	0.86	0.53	-	99,99,99,99	0
57	MG	1G	1605	1/1	0.81	0.39	-	104,104,104,104	0
57	MG	1H	3332	1/1	0.80	0.16	-	102,102,102,102	0
57	MG	1H	3315	1/1	0.97	0.10	-	63,63,63,63	0
57	MG	1G	1620	1/1	0.93	0.19	-	91,91,91,91	0
57	MG	14	3379	1/1	0.77	0.14	-	84,84,84,84	0
57	MG	14	3210	1/1	0.88	0.29	-	91,91,91,91	0
57	MG	1H	3184	1/1	0.94	0.24	-	70,70,70,70	0
57	MG	14	3209	1/1	0.87	0.17	-	83,83,83,83	0
57	MG	1H	3373	1/1	0.84	0.09	-	80,80,80,80	0
57	MG	13	1679	1/1	0.57	0.30	-	99,99,99,99	0
57	MG	1H	3512	1/1	0.86	0.13	-	87,87,87,87	0
57	MG	13	1637	1/1	0.74	0.32	-	74,74,74,74	0
57	MG	14	3371	1/1	0.93	0.10	-	107,107,107,107	0
57	MG	14	3075	1/1	0.82	0.16	-	73,73,73,73	0
57	MG	13	1613	1/1	0.88	0.13	-	87,87,87,87	0
57	MG	1H	3457	1/1	0.86	0.16	-	84,84,84,84	0
57	MG	1H	3120	1/1	0.60	0.18	-	88,88,88,88	0
57	MG	2L	102	1/1	0.88	0.18	-	132,132,132,132	0
57	MG	14	3286	1/1	0.85	0.11	-	73,73,73,73	0
57	MG	1G	1686	1/1	0.57	0.13	-	106,106,106,106	0
57	MG	14	3133	1/1	0.81	1.00	-	88,88,88,88	0
57	MG	1H	3359	1/1	0.98	0.17	-	44,44,44,44	0
57	MG	14	3278	1/1	0.83	0.09	-	102,102,102,102	0
57	MG	14	3012	1/1	0.96	0.32	-	57,57,57,57	0
57	MG	1H	3196	1/1	0.76	0.38	-	77,77,77,77	0
57	MG	14	3068	1/1	0.93	0.41	-	53,53,53,53	0
57	MG	1H	3122	1/1	0.91	0.09	-	52,52,52,52	0
57	MG	1H	3456	1/1	0.82	0.06	-	94,94,94,94	0
57	MG	1G	1637	1/1	0.80	0.16	-	103,103,103,103	0
57	MG	1H	3319	1/1	0.98	0.18	-	57,57,57,57	0
57	MG	14	3325	1/1	0.98	0.04	-	79,79,79,79	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	1G	1612	1/1	0.90	0.15	-	106,106,106,106	0
57	MG	14	3033	1/1	0.94	0.33	-	80,80,80,80	0
57	MG	1H	3225	1/1	0.72	0.43	-	71,71,71,71	0
57	MG	13	1686	1/1	0.92	0.17	-	76,76,76,76	0
57	MG	1H	3463	1/1	0.96	0.09	-	81,81,81,81	0
57	MG	14	3393	1/1	0.32	0.08	-	165,165,165,165	0
57	MG	1H	3032	1/1	0.99	0.25	-	47,47,47,47	0
57	MG	1H	3467	1/1	0.95	0.03	-	84,84,84,84	0
57	MG	14	3346	1/1	0.89	0.08	-	99,99,99,99	0
57	MG	14	3433	1/1	0.64	0.26	-	114,114,114,114	0
57	MG	13	1647	1/1	0.79	0.47	-	88,88,88,88	0
57	MG	13	1704	1/1	0.32	0.13	-	106,106,106,106	0
57	MG	14	3197	1/1	0.76	0.61	-	77,77,77,77	0
57	MG	14	3098	1/1	0.76	0.35	-	77,77,77,77	0
57	MG	1H	3189	1/1	0.92	0.35	-	68,68,68,68	0
57	MG	14	3228	1/1	0.98	0.14	-	56,56,56,56	0
57	MG	1H	3110	1/1	0.95	0.23	-	79,79,79,79	0
57	MG	14	3320	1/1	0.95	0.09	-	53,53,53,53	0
57	MG	14	3110	1/1	0.70	0.24	-	69,69,69,69	0
57	MG	14	3147	1/1	0.98	0.40	-	69,69,69,69	0
57	MG	1H	3423	1/1	0.67	0.18	-	84,84,84,84	0
57	MG	1H	3197	1/1	0.82	0.51	-	77,77,77,77	0
57	MG	13	1729	1/1	0.68	0.11	-	110,110,110,110	0
57	MG	13	1666	1/1	0.94	0.20	-	85,85,85,85	0
57	MG	14	3148	1/1	0.89	0.39	-	85,85,85,85	0
57	MG	1H	3434	1/1	0.79	0.17	-	95,95,95,95	0
57	MG	13	1609	1/1	0.97	0.15	-	70,70,70,70	0
57	MG	14	3421	1/1	0.79	0.10	-	97,97,97,97	0
57	MG	14	3093	1/1	0.97	0.42	-	84,84,84,84	0
57	MG	14	3312	1/1	0.86	0.16	-	92,92,92,92	0
57	MG	14	3085	1/1	0.94	0.25	-	88,88,88,88	0
57	MG	1H	3035	1/1	0.94	0.11	-	40,40,40,40	0
57	MG	1H	3202	1/1	0.91	0.45	-	73,73,73,73	0
57	MG	13	1710	1/1	0.86	0.07	-	96,96,96,96	0
57	MG	14	3327	1/1	0.90	0.09	-	82,82,82,82	0
57	MG	1H	3328	1/1	0.97	0.13	-	47,47,47,47	0
57	MG	1H	3081	1/1	0.19	0.45	-	83,83,83,83	0
57	MG	14	3345	1/1	0.82	0.16	-	99,99,99,99	0
57	MG	14	3339	1/1	0.80	0.10	-	107,107,107,107	0
57	MG	1H	3180	1/1	0.34	0.42	-	90,90,90,90	0
57	MG	1H	3148	1/1	0.85	0.37	-	83,83,83,83	0
57	MG	14	3017	1/1	0.68	0.34	-	119,119,119,119	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	1H	3216	1/1	0.69	0.27	-	82,82,82,82	0
57	MG	14	3376	1/1	0.65	0.23	-	102,102,102,102	0
57	MG	13	1719	1/1	0.89	0.04	-	81,81,81,81	0
57	MG	14	3187	1/1	0.79	0.25	-	117,117,117,117	0
57	MG	1H	3389	1/1	0.77	0.14	-	88,88,88,88	0
57	MG	14	3260	1/1	0.90	0.11	-	106,106,106,106	0
57	MG	13	1656	1/1	0.90	0.38	-	74,74,74,74	0
57	MG	13	1608	1/1	0.96	0.08	-	76,76,76,76	0
57	MG	1H	3514	1/1	0.95	0.05	-	98,98,98,98	0
57	MG	14	3409	1/1	0.72	0.22	-	88,88,88,88	0
57	MG	1H	3075	1/1	0.73	0.29	-	76,76,76,76	0
57	MG	1H	3404	1/1	0.90	0.13	-	71,71,71,71	0
57	MG	1G	1663	1/1	0.80	0.23	-	98,98,98,98	0
57	MG	1G	1624	1/1	0.91	0.13	-	95,95,95,95	0
57	MG	1H	3411	1/1	0.97	0.07	-	63,63,63,63	0
57	MG	1H	3296	1/1	0.88	0.08	-	91,91,91,91	0
57	MG	14	3258	1/1	0.77	0.07	-	79,79,79,79	0
57	MG	14	3274	1/1	0.77	0.05	-	85,85,85,85	0
57	MG	1H	3100	1/1	0.92	0.54	-	68,68,68,68	0
57	MG	13	1722	1/1	0.90	0.09	-	87,87,87,87	0
57	MG	14	3213	1/1	0.84	0.70	-	87,87,87,87	0
57	MG	1H	3125	1/1	0.81	0.20	-	56,56,56,56	0
57	MG	1G	1693	1/1	0.80	0.07	-	129,129,129,129	0
57	MG	14	3109	1/1	0.71	0.43	-	67,67,67,67	0
57	MG	1H	3096	1/1	0.95	0.14	-	64,64,64,64	0
57	MG	14	3177	1/1	0.87	0.45	-	101,101,101,101	0
57	MG	14	3383	1/1	0.91	0.07	-	90,90,90,90	0
57	MG	1G	1654	1/1	0.86	0.30	-	108,108,108,108	0
57	MG	1H	3065	1/1	0.68	0.53	-	65,65,65,65	0
57	MG	14	3140	1/1	0.96	0.13	-	51,51,51,51	0
57	MG	1G	1631	1/1	0.96	0.56	-	92,92,92,92	0
57	MG	13	1693	1/1	0.96	0.07	-	85,85,85,85	0
57	MG	1G	1630	1/1	0.91	0.37	-	72,72,72,72	0
57	MG	1H	3142	1/1	0.88	0.29	-	68,68,68,68	0
57	MG	14	3004	1/1	0.97	0.28	-	63,63,63,63	0
57	MG	13	1725	1/1	0.90	0.04	-	115,115,115,115	0
57	MG	13	1734	1/1	0.97	0.10	-	94,94,94,94	0
57	MG	1H	3471	1/1	0.96	0.12	-	90,90,90,90	0
57	MG	1H	3340	1/1	0.92	0.05	-	76,76,76,76	0
57	MG	14	3172	1/1	0.88	0.39	-	77,77,77,77	0
57	MG	14	3246	1/1	0.95	0.13	-	61,61,61,61	0
57	MG	14	3164	1/1	0.71	0.33	-	78,78,78,78	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	14	3372	1/1	0.96	0.12	-	86,86,86,86	0
57	MG	1H	3311	1/1	0.98	0.08	-	53,53,53,53	0
57	MG	14	3407	1/1	0.69	0.12	-	123,123,123,123	0
57	MG	14	3047	1/1	0.98	0.34	-	65,65,65,65	0
57	MG	1G	1626	1/1	0.94	0.19	-	83,83,83,83	0
57	MG	1H	3072	1/1	0.95	0.20	-	58,58,58,58	0
57	MG	1G	1613	1/1	0.93	0.76	-	89,89,89,89	0
57	MG	14	3168	1/1	0.78	0.24	-	89,89,89,89	0
57	MG	14	3170	1/1	0.85	0.59	-	81,81,81,81	0
57	MG	14	3397	1/1	0.90	0.17	-	96,96,96,96	0
57	MG	1H	3220	1/1	0.82	0.45	-	89,89,89,89	0
57	MG	1H	3445	1/1	0.18	0.12	-	92,92,92,92	0
57	MG	1H	3480	1/1	0.89	0.07	-	87,87,87,87	0
57	MG	1H	3038	1/1	0.82	0.32	-	50,50,50,50	0
57	MG	14	3352	1/1	0.93	0.07	-	82,82,82,82	0
57	MG	1H	3448	1/1	0.97	0.07	-	80,80,80,80	0
57	MG	1H	3077	1/1	0.92	0.42	-	80,80,80,80	0
57	MG	14	3291	1/1	0.72	0.21	-	85,85,85,85	0
57	MG	1H	3498	1/1	0.86	0.20	-	98,98,98,98	0
57	MG	14	3323	1/1	0.85	0.11	-	81,81,81,81	0
57	MG	14	3006	1/1	0.85	0.32	-	75,75,75,75	0
57	MG	14	3396	1/1	0.91	0.03	-	126,126,126,126	0
57	MG	14	3408	1/1	0.86	0.13	-	92,92,92,92	0
57	MG	1H	3037	1/1	0.90	0.54	-	75,75,75,75	0
57	MG	13	1687	1/1	0.94	0.06	-	103,103,103,103	0
57	MG	13	1658	1/1	0.89	0.25	-	82,82,82,82	0
57	MG	1H	3360	1/1	0.99	0.09	-	78,78,78,78	0
57	MG	1H	3493	1/1	0.95	0.04	-	101,101,101,101	0
57	MG	14	3180	1/1	0.91	0.41	-	91,91,91,91	0
57	MG	14	3021	1/1	0.95	0.35	-	38,38,38,38	0
57	MG	1H	3422	1/1	0.69	0.33	-	65,65,65,65	0
57	MG	14	3169	1/1	0.90	0.31	-	85,85,85,85	0
57	MG	14	3195	1/1	0.95	0.25	-	78,78,78,78	0
57	MG	14	3239	1/1	0.97	0.17	-	77,77,77,77	0
57	MG	1H	3227	1/1	0.72	0.28	-	72,72,72,72	0
57	MG	13	1675	1/1	0.81	0.47	-	101,101,101,101	0
57	MG	1H	3234	1/1	0.68	0.20	-	79,79,79,79	0
57	MG	1H	3174	1/1	0.86	0.23	-	55,55,55,55	0
57	MG	14	3204	1/1	0.83	0.22	-	72,72,72,72	0
57	MG	1H	3306	1/1	0.98	0.09	-	75,75,75,75	0
57	MG	14	3026	1/1	0.97	0.29	-	56,56,56,56	0
57	MG	1H	3501	1/1	0.92	0.06	-	90,90,90,90	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	14	3183	1/1	0.81	0.30	-	103,103,103,103	0
57	MG	1G	1642	1/1	0.55	0.28	-	99,99,99,99	0
57	MG	13	1717	1/1	0.92	0.11	-	99,99,99,99	0
57	MG	1H	3508	1/1	0.90	0.10	-	117,117,117,117	0
57	MG	1H	3129	1/1	0.78	0.20	-	76,76,76,76	0
57	MG	2K	101	1/1	0.91	0.11	-	100,100,100,100	0
57	MG	1H	3062	1/1	0.90	0.13	-	38,38,38,38	0
57	MG	14	3192	1/1	0.95	0.28	-	74,74,74,74	0
57	MG	1H	3157	1/1	0.54	0.38	-	89,89,89,89	0
57	MG	14	3049	1/1	0.86	1.18	-	83,83,83,83	0
57	MG	1G	1623	1/1	0.94	0.21	-	115,115,115,115	0
57	MG	14	3009	1/1	0.90	0.34	-	57,57,57,57	0
57	MG	1H	3443	1/1	0.86	0.10	-	75,75,75,75	0
57	MG	13	1632	1/1	0.96	0.22	-	78,78,78,78	0
57	MG	13	1655	1/1	0.89	0.50	-	82,82,82,82	0
57	MG	14	3357	1/1	0.92	0.08	-	105,105,105,105	0
57	MG	1H	3345	1/1	0.92	0.06	-	102,102,102,102	0
57	MG	13	1738	1/1	0.64	0.09	-	133,133,133,133	0
57	MG	13	1676	1/1	0.77	0.47	-	93,93,93,93	0
57	MG	1H	3049	1/1	0.83	0.25	-	62,62,62,62	0
57	MG	13	1713	1/1	0.88	0.08	-	86,86,86,86	0
57	MG	14	3165	1/1	0.86	0.66	-	77,77,77,77	0
57	MG	14	3332	1/1	0.65	0.09	-	102,102,102,102	0
57	MG	14	3106	1/1	0.94	0.41	-	57,57,57,57	0
57	MG	16	207	1/1	0.92	0.33	-	74,74,74,74	0
57	MG	1H	3266	1/1	0.94	0.08	-	54,54,54,54	0
57	MG	1H	3123	1/1	0.77	0.37	-	65,65,65,65	0
57	MG	14	3223	1/1	0.89	0.10	-	87,87,87,87	0
57	MG	1G	1653	1/1	0.95	0.11	-	97,97,97,97	0
57	MG	1H	3482	1/1	0.85	0.09	-	94,94,94,94	0
57	MG	1H	3366	1/1	0.94	0.17	-	68,68,68,68	0
57	MG	1H	3060	1/1	0.99	0.16	-	35,35,35,35	0
57	MG	14	3010	1/1	0.94	0.24	-	57,57,57,57	0
57	MG	14	3423	1/1	0.90	0.07	-	120,120,120,120	0
57	MG	1J	201	1/1	0.90	0.29	-	73,73,73,73	0
57	MG	14	3037	1/1	0.97	0.26	-	65,65,65,65	0
57	MG	13	1667	1/1	0.93	0.67	-	89,89,89,89	0
57	MG	1H	3261	1/1	0.93	0.08	-	63,63,63,63	0
57	MG	1H	3344	1/1	0.95	0.08	-	103,103,103,103	0
57	MG	14	3273	1/1	0.88	0.09	-	86,86,86,86	0
57	MG	13	1706	1/1	0.77	0.08	-	117,117,117,117	0
57	MG	14	3414	1/1	0.84	0.12	-	97,97,97,97	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	1H	3012	1/1	0.97	0.41	-	58,58,58,58	0
57	MG	14	3094	1/1	0.93	0.65	-	73,73,73,73	0
57	MG	1H	3163	1/1	0.78	0.44	-	91,91,91,91	0
57	MG	1G	1635	1/1	0.71	0.20	-	105,105,105,105	0
57	MG	1H	3154	1/1	0.53	0.23	-	58,58,58,58	0
57	MG	1G	1687	1/1	0.90	0.08	-	111,111,111,111	0
57	MG	16	201	1/1	0.84	0.25	-	62,62,62,62	0
57	MG	14	3178	1/1	0.92	0.73	-	78,78,78,78	0
57	MG	14	3330	1/1	0.71	0.14	-	94,94,94,94	0
57	MG	1H	3468	1/1	0.93	0.24	-	82,82,82,82	0
57	MG	E5	101	1/1	0.60	0.58	-	89,89,89,89	0
57	MG	1H	3226	1/1	0.74	0.31	-	71,71,71,71	0
57	MG	1H	3083	1/1	0.81	0.84	-	83,83,83,83	0
57	MG	1H	3523	1/1	0.83	0.11	-	82,82,82,82	0
57	MG	14	3363	1/1	0.95	0.09	-	107,107,107,107	0
57	MG	13	1716	1/1	0.94	0.10	-	71,71,71,71	0
57	MG	14	3153	1/1	0.84	0.59	-	90,90,90,90	0
57	MG	1G	1629	1/1	0.88	0.52	-	85,85,85,85	0
57	MG	13	1684	1/1	0.89	0.08	-	85,85,85,85	0
57	MG	1H	3444	1/1	0.74	0.11	-	74,74,74,74	0
57	MG	1G	1655	1/1	0.93	0.12	-	110,110,110,110	0
57	MG	14	3308	1/1	0.97	0.10	-	94,94,94,94	0
57	MG	14	3411	1/1	0.87	0.04	-	105,105,105,105	0
57	MG	14	3292	1/1	0.94	0.12	-	74,74,74,74	0
57	MG	1G	1694	1/1	0.79	0.12	-	113,113,113,113	0
57	MG	1H	3233	1/1	0.95	0.07	-	93,93,93,93	0
57	MG	13	1696	1/1	0.88	0.05	-	93,93,93,93	0
57	MG	14	3207	1/1	0.71	0.37	-	70,70,70,70	0
57	MG	1H	3006	1/1	0.72	0.28	-	74,74,74,74	0
60	ZN	G8	201	1/1	0.92	0.14	-	139,139,139,139	0
57	MG	1H	3118	1/1	0.75	0.15	-	69,69,69,69	0
57	MG	14	3422	1/1	0.52	0.24	-	108,108,108,108	0
57	MG	14	3430	1/1	0.61	0.18	-	107,107,107,107	0
57	MG	1G	1634	1/1	0.79	0.72	-	85,85,85,85	0
57	MG	1H	3005	1/1	0.86	0.21	-	68,68,68,68	0
57	MG	14	3078	1/1	0.96	0.29	-	56,56,56,56	0
57	MG	14	3394	1/1	0.96	0.05	-	91,91,91,91	0
57	MG	14	3173	1/1	0.70	0.45	-	79,79,79,79	0
57	MG	1H	3231	1/1	0.91	0.49	-	66,66,66,66	0
57	MG	1H	3016	1/1	0.98	0.19	-	44,44,44,44	0
57	MG	1H	3101	1/1	0.86	0.41	-	78,78,78,78	0
57	MG	1H	3250	1/1	0.95	0.18	-	41,41,41,41	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	1G	1661	1/1	0.97	0.08	-	95,95,95,95	0
57	MG	1H	3368	1/1	0.80	0.12	-	69,69,69,69	0
57	MG	1H	3291	1/1	0.99	0.13	-	59,59,59,59	0
57	MG	13	1625	1/1	0.94	0.16	-	82,82,82,82	0
57	MG	1H	3391	1/1	0.80	0.13	-	89,89,89,89	0
57	MG	1G	1700	1/1	0.93	0.06	-	109,109,109,109	0
57	MG	13	1727	1/1	0.76	0.13	-	90,90,90,90	0
57	MG	13	1711	1/1	0.80	0.16	-	74,74,74,74	0
57	MG	13	1673	1/1	0.68	0.40	-	84,84,84,84	0
57	MG	14	3319	1/1	0.90	0.07	-	95,95,95,95	0
57	MG	1H	3363	1/1	0.77	0.12	-	96,96,96,96	0
57	MG	1H	3310	1/1	0.91	0.18	-	70,70,70,70	0
57	MG	14	3050	1/1	0.94	0.20	-	57,57,57,57	0
57	MG	1H	3145	1/1	0.56	0.53	-	80,80,80,80	0
57	MG	13	1651	1/1	0.87	0.29	-	98,98,98,98	0
57	MG	14	3052	1/1	0.97	0.23	-	54,54,54,54	0
57	MG	14	3297	1/1	0.91	0.17	-	71,71,71,71	0
57	MG	1J	207	1/1	0.79	0.15	-	97,97,97,97	0
57	MG	1H	3113	1/1	0.48	0.26	-	81,81,81,81	0
57	MG	1H	3438	1/1	0.93	0.04	-	62,62,62,62	0
57	MG	1H	3058	1/1	0.85	0.17	-	60,60,60,60	0
57	MG	13	1615	1/1	0.88	0.24	-	88,88,88,88	0
57	MG	14	3424	1/1	0.88	0.09	-	109,109,109,109	0
57	MG	14	3039	1/1	0.91	0.71	-	89,89,89,89	0
57	MG	14	3182	1/1	0.96	0.35	-	74,74,74,74	0
57	MG	1G	1695	1/1	0.86	0.23	-	107,107,107,107	0
57	MG	1G	1671	1/1	0.72	0.17	-	112,112,112,112	0
57	MG	1H	3146	1/1	0.92	0.16	-	52,52,52,52	0
57	MG	1H	3001	1/1	0.87	0.39	-	84,84,84,84	0
57	MG	1H	3183	1/1	0.96	0.21	-	70,70,70,70	0
57	MG	3I	201	1/1	0.92	0.24	-	60,60,60,60	0
57	MG	14	3043	1/1	0.93	0.29	-	71,71,71,71	0
57	MG	1H	3008	1/1	0.87	0.39	-	82,82,82,82	0
57	MG	14	3386	1/1	0.82	0.09	-	74,74,74,74	0
57	MG	1H	3238	1/1	0.94	0.34	-	80,80,80,80	0
57	MG	1H	3274	1/1	0.84	0.14	-	97,97,97,97	0
57	MG	1H	3160	1/1	0.93	0.18	-	71,71,71,71	0
57	MG	1H	3333	1/1	0.98	0.13	-	77,77,77,77	0
57	MG	1H	3276	1/1	0.94	0.10	-	78,78,78,78	0
57	MG	14	3264	1/1	0.71	0.14	-	93,93,93,93	0
57	MG	1G	1669	1/1	0.81	0.09	-	105,105,105,105	0
57	MG	1H	3019	1/1	0.95	0.23	-	32,32,32,32	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	1H	3505	1/1	0.90	0.14	-	100,100,100,100	0
57	MG	13	1740	1/1	0.86	0.06	-	98,98,98,98	0
57	MG	1H	3292	1/1	0.98	0.21	-	43,43,43,43	0
57	MG	1H	3449	1/1	0.94	0.10	-	80,80,80,80	0
57	MG	32	301	1/1	0.85	0.47	-	113,113,113,113	0
57	MG	1G	1625	1/1	0.89	0.08	-	99,99,99,99	0
57	MG	1H	3214	1/1	0.96	0.47	-	77,77,77,77	0
57	MG	1H	3379	1/1	0.88	0.09	-	73,73,73,73	0
57	MG	35	201	1/1	0.69	0.22	-	75,75,75,75	0
57	MG	1H	3401	1/1	0.81	0.09	-	64,64,64,64	0
57	MG	1H	3244	1/1	0.90	0.42	-	74,74,74,74	0
57	MG	1G	1602	1/1	0.92	0.33	-	106,106,106,106	0
57	MG	1H	3374	1/1	0.76	0.11	-	83,83,83,83	0
57	MG	1H	3412	1/1	0.87	0.10	-	75,75,75,75	0
57	MG	1H	3518	1/1	0.78	0.11	-	96,96,96,96	0
57	MG	14	3284	1/1	0.93	0.14	-	72,72,72,72	0
57	MG	1H	3152	1/1	0.98	0.41	-	70,70,70,70	0
57	MG	14	3135	1/1	0.66	0.32	-	79,79,79,79	0
57	MG	14	3252	1/1	0.72	0.18	-	90,90,90,90	0
57	MG	13	1616	1/1	0.87	0.48	-	74,74,74,74	0
57	MG	14	3333	1/1	0.92	0.17	-	50,50,50,50	0
57	MG	1G	1688	1/1	0.79	0.24	-	113,113,113,113	0
57	MG	1H	3095	1/1	0.92	0.17	-	70,70,70,70	0
57	MG	1H	3502	1/1	0.85	0.18	-	141,141,141,141	0
57	MG	1H	3453	1/1	0.83	0.12	-	93,93,93,93	0
57	MG	14	3115	1/1	0.40	0.78	-	87,87,87,87	0
57	MG	14	3360	1/1	0.92	0.07	-	96,96,96,96	0
57	MG	14	3167	1/1	0.88	0.32	-	78,78,78,78	0
57	MG	14	3266	1/1	0.65	0.08	-	107,107,107,107	0
57	MG	14	3011	1/1	0.91	0.18	-	44,44,44,44	0
57	MG	14	3435	1/1	0.70	0.20	-	92,92,92,92	0
57	MG	1H	3426	1/1	0.96	0.05	-	74,74,74,74	0
57	MG	14	3001	1/1	0.89	0.17	-	57,57,57,57	0
57	MG	1G	1696	1/1	0.91	0.07	-	128,128,128,128	0
57	MG	13	1736	1/1	0.96	0.12	-	101,101,101,101	0
57	MG	1H	3451	1/1	0.65	0.08	-	102,102,102,102	0
57	MG	1H	3516	1/1	0.77	0.14	-	102,102,102,102	0
57	MG	1H	3342	1/1	0.96	0.06	-	86,86,86,86	0
57	MG	14	3129	1/1	0.88	0.42	-	118,118,118,118	0
57	MG	1H	3161	1/1	0.83	0.38	-	67,67,67,67	0
57	MG	14	3269	1/1	0.94	0.05	-	78,78,78,78	0
57	MG	1H	3068	1/1	0.95	0.40	-	72,72,72,72	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	1H	3124	1/1	0.92	0.21	-	61,61,61,61	0
57	MG	1H	3464	1/1	0.96	0.08	-	63,63,63,63	0
57	MG	13	1685	1/1	0.64	0.16	-	90,90,90,90	0
57	MG	1H	3191	1/1	0.82	0.27	-	58,58,58,58	0
57	MG	1H	3256	1/1	0.89	0.13	-	47,47,47,47	0
57	MG	1H	3223	1/1	0.82	0.43	-	85,85,85,85	0
57	MG	C5	201	1/1	0.65	0.10	-	113,113,113,113	0
57	MG	14	3158	1/1	0.90	0.24	-	66,66,66,66	0
57	MG	14	3084	1/1	0.87	0.27	-	79,79,79,79	0
57	MG	1G	1679	1/1	0.88	0.09	-	94,94,94,94	0
57	MG	13	1728	1/1	0.83	0.05	-	117,117,117,117	0
57	MG	14	3347	1/1	0.64	0.10	-	88,88,88,88	0
57	MG	1H	3413	1/1	0.97	0.10	-	78,78,78,78	0
57	MG	1H	3258	1/1	0.96	0.11	-	47,47,47,47	0
57	MG	1H	3159	1/1	0.95	0.13	-	80,80,80,80	0
57	MG	13	1678	1/1	0.01	0.16	-	112,112,112,112	0
57	MG	1G	1608	1/1	0.92	0.37	-	93,93,93,93	0
57	MG	1H	3017	1/1	0.95	0.29	-	62,62,62,62	0
57	MG	1H	3271	1/1	0.86	0.19	-	51,51,51,51	0
57	MG	1H	3481	1/1	0.85	0.11	-	103,103,103,103	0
57	MG	13	1702	1/1	0.98	0.13	-	85,85,85,85	0
57	MG	1H	3007	1/1	0.59	0.31	-	81,81,81,81	0
57	MG	14	3328	1/1	0.85	0.08	-	81,81,81,81	0
57	MG	1G	1662	1/1	0.92	0.13	-	91,91,91,91	0
57	MG	1H	3211	1/1	0.66	0.46	-	78,78,78,78	0
57	MG	1H	3206	1/1	0.78	0.56	-	93,93,93,93	0
57	MG	1H	3287	1/1	0.96	0.08	-	71,71,71,71	0
57	MG	13	1735	1/1	0.91	0.07	-	100,100,100,100	0
57	MG	1H	3018	1/1	0.97	0.18	-	41,41,41,41	0
57	MG	1H	3277	1/1	0.77	0.16	-	71,71,71,71	0
57	MG	1H	3144	1/1	0.80	0.60	-	102,102,102,102	0
57	MG	14	3374	1/1	0.89	0.12	-	92,92,92,92	0
57	MG	13	1626	1/1	0.90	0.33	-	84,84,84,84	0
57	MG	1H	3500	1/1	0.62	0.14	-	94,94,94,94	0
57	MG	1H	3179	1/1	0.95	0.14	-	80,80,80,80	0
57	MG	14	3299	1/1	0.89	0.11	-	79,79,79,79	0
57	MG	13	1721	1/1	0.95	0.17	-	106,106,106,106	0
57	MG	21	302	1/1	0.86	0.25	-	72,72,72,72	0
57	MG	1H	3370	1/1	0.92	0.08	-	68,68,68,68	0
57	MG	14	3428	1/1	0.77	0.29	-	119,119,119,119	0
57	MG	14	3127	1/1	0.56	1.03	-	87,87,87,87	0
57	MG	1G	1698	1/1	0.96	0.10	-	123,123,123,123	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	13	1652	1/1	0.96	0.62	-	71,71,71,71	0
57	MG	39	302	1/1	0.82	0.20	-	74,74,74,74	0
57	MG	1H	3137	1/1	0.86	0.43	-	70,70,70,70	0
57	MG	13	1730	1/1	0.88	0.09	-	94,94,94,94	0
57	MG	13	1660	1/1	0.74	0.23	-	88,88,88,88	0
57	MG	14	3051	1/1	0.93	0.27	-	61,61,61,61	0
57	MG	1H	3424	1/1	0.89	0.04	-	101,101,101,101	0
57	MG	1H	3406	1/1	0.84	0.08	-	85,85,85,85	0
57	MG	1H	3503	1/1	0.64	0.12	-	91,91,91,91	0
57	MG	1G	1628	1/1	0.90	0.31	-	113,113,113,113	0
57	MG	14	3101	1/1	0.96	0.33	-	87,87,87,87	0
57	MG	1H	3218	1/1	0.94	0.21	-	37,37,37,37	0
57	MG	14	3256	1/1	0.84	0.13	-	83,83,83,83	0
57	MG	1H	3164	1/1	0.85	0.41	-	69,69,69,69	0
57	MG	1H	3003	1/1	0.77	0.37	-	65,65,65,65	0
57	MG	14	3355	1/1	0.56	0.10	-	104,104,104,104	0
57	MG	14	3227	1/1	0.90	0.15	-	58,58,58,58	0
57	MG	14	3193	1/1	0.90	0.13	-	45,45,45,45	0
57	MG	1H	3376	1/1	0.83	0.15	-	66,66,66,66	0
57	MG	14	3277	1/1	0.68	0.16	-	84,84,84,84	0
57	MG	14	3382	1/1	0.78	0.12	-	66,66,66,66	0
57	MG	14	3334	1/1	0.95	0.10	-	60,60,60,60	0
57	MG	1H	3241	1/1	0.78	0.22	-	58,58,58,58	0
57	MG	1H	3475	1/1	0.94	0.12	-	97,97,97,97	0
57	MG	14	3103	1/1	0.76	0.43	-	96,96,96,96	0
57	MG	1H	3053	1/1	0.76	0.46	-	75,75,75,75	0
57	MG	1G	1621	1/1	0.85	0.42	-	82,82,82,82	0
57	MG	14	3335	1/1	0.93	0.08	-	81,81,81,81	0
57	MG	1H	3462	1/1	0.93	0.10	-	94,94,94,94	0
57	MG	1H	3317	1/1	0.96	0.07	-	66,66,66,66	0
57	MG	1H	3192	1/1	0.87	0.34	-	75,75,75,75	0
57	MG	1H	3416	1/1	0.90	0.06	-	89,89,89,89	0
57	MG	13	1634	1/1	0.94	0.36	-	75,75,75,75	0
57	MG	14	3337	1/1	0.92	0.14	-	104,104,104,104	0
57	MG	1H	3385	1/1	0.98	0.15	-	25,25,25,25	0
57	MG	1H	3182	1/1	0.95	0.57	-	60,60,60,60	0
57	MG	1H	3479	1/1	0.76	0.10	-	83,83,83,83	0
57	MG	1H	3513	1/1	0.86	0.15	-	53,53,53,53	0
57	MG	1H	3507	1/1	0.94	0.08	-	57,57,57,57	0
57	MG	1H	3497	1/1	0.79	0.10	-	107,107,107,107	0
57	MG	1H	3436	1/1	0.93	0.10	-	70,70,70,70	0
57	MG	14	3138	1/1	0.71	0.67	-	78,78,78,78	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	14	3316	1/1	0.90	0.13	-	69,69,69,69	0
57	MG	1G	1690	1/1	0.80	0.06	-	117,117,117,117	0
57	MG	1H	3139	1/1	0.95	0.35	-	71,71,71,71	0
57	MG	14	3234	1/1	0.96	0.18	-	57,57,57,57	0
57	MG	14	3257	1/1	0.78	0.16	-	69,69,69,69	0
57	MG	13	1650	1/1	0.85	0.40	-	75,75,75,75	0
57	MG	1H	3253	1/1	0.92	0.08	-	42,42,42,42	0
57	MG	1H	3033	1/1	0.96	0.25	-	70,70,70,70	0
57	MG	14	3358	1/1	0.90	0.15	-	86,86,86,86	0
57	MG	31	301	1/1	0.69	0.13	-	63,63,63,63	0
57	MG	1H	3504	1/1	0.83	0.07	-	109,109,109,109	0
57	MG	1H	3520	1/1	0.54	0.10	-	105,105,105,105	0
57	MG	1H	3521	1/1	0.87	0.12	-	82,82,82,82	0
57	MG	13	1665	1/1	0.85	0.69	-	82,82,82,82	0
57	MG	1H	3026	1/1	0.94	0.39	-	55,55,55,55	0
57	MG	1H	3476	1/1	0.74	0.21	-	82,82,82,82	0
57	MG	1H	3487	1/1	0.68	0.11	-	85,85,85,85	0
57	MG	1H	3106	1/1	0.81	0.23	-	87,87,87,87	0
57	MG	14	3343	1/1	0.84	0.06	-	80,80,80,80	0
57	MG	1H	3044	1/1	0.98	0.36	-	67,67,67,67	0
57	MG	14	3255	1/1	0.86	0.13	-	79,79,79,79	0
57	MG	1H	3324	1/1	0.96	0.08	-	70,70,70,70	0
57	MG	1H	3405	1/1	0.80	0.09	-	66,66,66,66	0
57	MG	14	3032	1/1	0.97	0.41	-	64,64,64,64	0
57	MG	1J	203	1/1	0.95	0.26	-	73,73,73,73	0
57	MG	1H	3428	1/1	0.70	0.09	-	89,89,89,89	0
57	MG	1G	1639	1/1	0.90	0.54	-	78,78,78,78	0
57	MG	14	3298	1/1	0.82	0.11	-	86,86,86,86	0
57	MG	1H	3293	1/1	0.93	0.14	-	73,73,73,73	0
57	MG	14	3412	1/1	0.82	0.10	-	97,97,97,97	0
57	MG	1H	3446	1/1	0.93	0.06	-	98,98,98,98	0
57	MG	1H	3111	1/1	0.79	0.61	-	79,79,79,79	0
57	MG	14	3221	1/1	0.82	0.20	-	62,62,62,62	0
57	MG	14	3176	1/1	0.67	0.20	-	82,82,82,82	0
57	MG	14	3367	1/1	0.94	0.08	-	106,106,106,106	0
57	MG	14	3301	1/1	0.98	0.04	-	85,85,85,85	0
57	MG	1H	3442	1/1	0.65	0.09	-	103,103,103,103	0
57	MG	13	1622	1/1	0.86	0.26	-	63,63,63,63	0
57	MG	14	3079	1/1	0.93	0.30	-	48,48,48,48	0
57	MG	14	3069	1/1	0.80	0.46	-	86,86,86,86	0
57	MG	14	3338	1/1	0.93	0.05	-	107,107,107,107	0
57	MG	1H	3511	1/1	0.68	0.16	-	101,101,101,101	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	1H	3329	1/1	0.80	0.08	-	85,85,85,85	0
57	MG	14	3265	1/1	0.94	0.07	-	91,91,91,91	0
57	MG	13	1697	1/1	0.91	0.07	-	84,84,84,84	0
57	MG	1H	3224	1/1	0.80	0.33	-	85,85,85,85	0
57	MG	14	3401	1/1	0.86	0.15	-	110,110,110,110	0
57	MG	1H	3394	1/1	0.85	0.11	-	49,49,49,49	0
57	MG	14	3163	1/1	0.51	0.38	-	93,93,93,93	0
57	MG	1G	1616	1/1	0.95	0.13	-	94,94,94,94	0
57	MG	1H	3384	1/1	0.87	0.08	-	67,67,67,67	0
57	MG	1H	3337	1/1	0.94	0.05	-	86,86,86,86	0
57	MG	1G	1658	1/1	0.95	0.06	-	104,104,104,104	0
57	MG	14	3128	1/1	0.74	0.41	-	89,89,89,89	0
57	MG	1H	3063	1/1	0.76	0.27	-	100,100,100,100	0
57	MG	14	3362	1/1	0.94	0.06	-	114,114,114,114	0
57	MG	14	3353	1/1	0.95	0.08	-	83,83,83,83	0
57	MG	1H	3236	1/1	0.83	0.31	-	67,67,67,67	0
57	MG	14	3157	1/1	0.85	0.32	-	92,92,92,92	0
57	MG	1H	3213	1/1	0.87	0.29	-	71,71,71,71	0
57	MG	1H	3362	1/1	0.88	0.13	-	100,100,100,100	0
57	MG	1G	1683	1/1	0.82	0.13	-	121,121,121,121	0
57	MG	1H	3515	1/1	0.88	0.19	-	58,58,58,58	0
57	MG	P8	101	1/1	0.73	0.56	-	71,71,71,71	0
57	MG	1H	3171	1/1	0.73	0.38	-	73,73,73,73	0
57	MG	1H	3488	1/1	0.77	0.28	-	105,105,105,105	0
57	MG	1H	3052	1/1	0.95	0.18	-	59,59,59,59	0
57	MG	1H	3103	1/1	0.90	0.32	-	61,61,61,61	0
57	MG	14	3226	1/1	0.97	0.16	-	51,51,51,51	0
57	MG	1H	3494	1/1	0.92	0.08	-	72,72,72,72	0
57	MG	1H	3354	1/1	0.97	0.17	-	60,60,60,60	0
57	MG	14	3403	1/1	0.87	0.12	-	84,84,84,84	0
57	MG	14	3417	1/1	0.73	0.12	-	108,108,108,108	0
57	MG	1H	3388	1/1	0.66	0.13	-	55,55,55,55	0
57	MG	1H	3247	1/1	0.97	0.14	-	36,36,36,36	0
57	MG	16	209	1/1	0.91	0.47	-	76,76,76,76	0
57	MG	1H	3069	1/1	0.88	0.41	-	69,69,69,69	0
57	MG	1H	3286	1/1	0.96	0.12	-	40,40,40,40	0
57	MG	1J	206	1/1	0.86	0.24	-	94,94,94,94	0
57	MG	1H	3414	1/1	0.76	0.14	-	105,105,105,105	0
57	MG	14	3067	1/1	0.78	0.23	-	78,78,78,78	0
57	MG	1G	1670	1/1	0.81	0.06	-	134,134,134,134	0
57	MG	1H	3341	1/1	0.81	0.12	-	78,78,78,78	0
57	MG	14	3044	1/1	0.96	0.18	-	57,57,57,57	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	14	3419	1/1	0.85	0.06	-	110,110,110,110	0
57	MG	14	3359	1/1	0.91	0.08	-	93,93,93,93	0
57	MG	25	201	1/1	0.91	0.05	-	115,115,115,115	0
57	MG	14	3214	1/1	0.83	0.70	-	80,80,80,80	0
57	MG	13	1694	1/1	0.88	0.14	-	82,82,82,82	0
57	MG	1H	3496	1/1	0.67	0.16	-	89,89,89,89	0
57	MG	13	1695	1/1	0.90	0.10	-	106,106,106,106	0
57	MG	14	3434	1/1	0.66	0.12	-	110,110,110,110	0
57	MG	1H	3185	1/1	0.91	0.48	-	64,64,64,64	0
57	MG	1G	1659	1/1	0.87	0.04	-	115,115,115,115	0
57	MG	14	3161	1/1	0.71	0.42	-	78,78,78,78	0
57	MG	1H	3372	1/1	0.94	0.08	-	89,89,89,89	0
57	MG	14	3144	1/1	0.89	0.32	-	102,102,102,102	0
57	MG	1H	3054	1/1	0.95	0.35	-	52,52,52,52	0
57	MG	1H	3229	1/1	0.83	0.50	-	90,90,90,90	0
57	MG	14	3344	1/1	0.66	0.09	-	93,93,93,93	0
57	MG	1H	3030	1/1	0.94	0.10	-	39,39,39,39	0
57	MG	16	205	1/1	0.73	0.18	-	69,69,69,69	0
57	MG	1H	3168	1/1	0.72	0.49	-	80,80,80,80	0
57	MG	13	1640	1/1	0.90	0.28	-	92,92,92,92	0
57	MG	14	3073	1/1	0.74	0.31	-	78,78,78,78	0
57	MG	1H	3407	1/1	0.83	0.09	-	72,72,72,72	0
57	MG	13	1644	1/1	0.94	0.45	-	85,85,85,85	0
57	MG	14	3241	1/1	0.89	0.05	-	78,78,78,78	0
57	MG	14	3071	1/1	0.93	0.19	-	74,74,74,74	0
57	MG	13	1707	1/1	0.92	0.10	-	56,56,56,56	0
57	MG	1H	3308	1/1	0.81	0.14	-	50,50,50,50	0
57	MG	14	3196	1/1	0.85	0.51	-	71,71,71,71	0
57	MG	16	210	1/1	0.93	0.11	-	73,73,73,73	0
57	MG	14	3385	1/1	0.70	0.16	-	67,67,67,67	0
57	MG	14	3399	1/1	0.85	0.09	-	123,123,123,123	0
57	MG	1H	3322	1/1	0.98	0.06	-	58,58,58,58	0
57	MG	14	3381	1/1	0.79	0.31	-	79,79,79,79	0
57	MG	14	3130	1/1	0.85	0.24	-	66,66,66,66	0
57	MG	14	3058	1/1	0.95	0.34	-	84,84,84,84	0
57	MG	1H	3510	1/1	0.94	0.14	-	39,39,39,39	0
57	MG	14	3404	1/1	0.82	0.16	-	97,97,97,97	0
57	MG	14	3038	1/1	0.94	0.21	-	52,52,52,52	0
57	MG	14	3285	1/1	0.81	0.06	-	108,108,108,108	0
57	MG	1H	3314	1/1	0.87	0.08	-	68,68,68,68	0
57	MG	14	3013	1/1	0.98	0.23	-	47,47,47,47	0
57	MG	1H	3351	1/1	0.91	0.09	-	68,68,68,68	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	13	1726	1/1	0.87	0.08	-	74,74,74,74	0
57	MG	1G	1692	1/1	0.90	0.12	-	101,101,101,101	0
57	MG	14	3188	1/1	0.91	0.50	-	72,72,72,72	0
57	MG	1G	1691	1/1	0.84	0.05	-	133,133,133,133	0
57	MG	16	208	1/1	0.70	0.39	-	85,85,85,85	0
57	MG	1H	3087	1/1	0.92	0.20	-	71,71,71,71	0
57	MG	1H	3349	1/1	0.98	0.12	-	50,50,50,50	0
57	MG	1H	3435	1/1	0.90	0.06	-	62,62,62,62	0
57	MG	14	3131	1/1	0.81	0.28	-	64,64,64,64	0
57	MG	1H	3316	1/1	0.95	0.07	-	78,78,78,78	0
57	MG	14	3189	1/1	0.95	0.33	-	51,51,51,51	0
57	MG	14	3215	1/1	0.84	1.26	-	86,86,86,86	0
57	MG	1G	1610	1/1	0.84	0.23	-	91,91,91,91	0
57	MG	14	3212	1/1	0.70	0.32	-	75,75,75,75	0
57	MG	1G	1641	1/1	0.85	0.59	-	103,103,103,103	0
57	MG	14	3418	1/1	0.84	0.07	-	90,90,90,90	0
57	MG	1H	3169	1/1	0.97	0.45	-	91,91,91,91	0
57	MG	1H	3399	1/1	0.96	0.06	-	73,73,73,73	0
57	MG	1H	3409	1/1	0.90	0.07	-	85,85,85,85	0
57	MG	1G	1697	1/1	0.81	0.09	-	128,128,128,128	0
57	MG	14	3402	1/1	0.89	0.08	-	78,78,78,78	0
57	MG	1G	1674	1/1	0.83	0.08	-	107,107,107,107	0
57	MG	1H	3367	1/1	0.82	0.08	-	66,66,66,66	0
57	MG	1H	3396	1/1	0.74	0.22	-	59,59,59,59	0
57	MG	1H	3429	1/1	0.90	0.10	-	88,88,88,88	0
57	MG	1G	1643	1/1	0.90	0.54	-	83,83,83,83	0
57	MG	14	3003	1/1	0.83	0.27	-	78,78,78,78	0
57	MG	1J	205	1/1	0.82	0.16	-	73,73,73,73	0
57	MG	14	3340	1/1	0.94	0.09	-	107,107,107,107	0
57	MG	1H	3352	1/1	0.77	0.12	-	57,57,57,57	0
57	MG	14	3331	1/1	0.93	0.20	-	90,90,90,90	0
57	MG	14	3391	1/1	0.83	0.23	-	99,99,99,99	0
57	MG	14	3219	1/1	0.96	0.10	-	57,57,57,57	0
57	MG	13	1619	1/1	0.98	0.21	-	52,52,52,52	0
57	MG	13	1639	1/1	0.93	0.58	-	70,70,70,70	0
57	MG	1H	3237	1/1	0.57	0.46	-	81,81,81,81	0
57	MG	14	3123	1/1	0.94	0.28	-	83,83,83,83	0
57	MG	1H	3153	1/1	0.77	0.44	-	77,77,77,77	0
57	MG	1H	3499	1/1	0.68	0.09	-	112,112,112,112	0
57	MG	14	3066	1/1	0.98	0.57	-	61,61,61,61	0
57	MG	14	3300	1/1	0.95	0.09	-	89,89,89,89	0
57	MG	14	3160	1/1	0.89	0.18	-	65,65,65,65	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	1H	3204	1/1	0.66	0.20	-	91,91,91,91	0
57	MG	14	3425	1/1	0.85	0.12	-	91,91,91,91	0
57	MG	1H	3135	1/1	0.71	0.49	-	79,79,79,79	0
57	MG	1H	3440	1/1	0.96	0.09	-	92,92,92,92	0
57	MG	1H	3010	1/1	0.89	0.37	-	39,39,39,39	0
57	MG	1H	3402	1/1	0.96	0.05	-	85,85,85,85	0
57	MG	1H	3243	1/1	0.76	0.17	-	82,82,82,82	0
57	MG	14	3406	1/1	0.72	0.12	-	101,101,101,101	0
57	MG	14	3175	1/1	0.79	0.21	-	88,88,88,88	0
57	MG	1H	3158	1/1	0.92	0.17	-	71,71,71,71	0
57	MG	1H	3209	1/1	0.92	0.60	-	76,76,76,76	0
57	MG	1H	3338	1/1	0.96	0.11	-	66,66,66,66	0
57	MG	1H	3014	1/1	0.96	0.17	-	53,53,53,53	0
57	MG	1H	3472	1/1	0.95	0.04	-	83,83,83,83	0
57	MG	1H	3132	1/1	0.91	0.23	-	65,65,65,65	0
57	MG	1G	1673	1/1	0.34	0.14	-	115,115,115,115	0
57	MG	13	1705	1/1	0.96	0.17	-	97,97,97,97	0
57	MG	1H	3295	1/1	0.94	0.09	-	89,89,89,89	0
57	MG	1H	3089	1/1	0.78	0.20	-	38,38,38,38	0
57	MG	14	3002	1/1	0.97	0.37	-	57,57,57,57	0
57	MG	13	1690	1/1	0.96	0.05	-	70,70,70,70	0
57	MG	1H	3455	1/1	0.87	0.15	-	76,76,76,76	0
57	MG	1G	1648	1/1	0.89	0.16	-	115,115,115,115	0
57	MG	1H	3232	1/1	0.93	0.22	-	48,48,48,48	0
57	MG	1H	3126	1/1	0.88	0.43	-	86,86,86,86	0
57	MG	1H	3326	1/1	0.95	0.10	-	62,62,62,62	0
57	MG	1G	1665	1/1	0.95	0.05	-	91,91,91,91	0
57	MG	14	3259	1/1	0.92	0.10	-	71,71,71,71	0
57	MG	1H	3357	1/1	0.94	0.12	-	52,52,52,52	0
57	MG	1H	3255	1/1	0.98	0.08	-	49,49,49,49	0
57	MG	13	1604	1/1	0.96	0.15	-	62,62,62,62	0
57	MG	14	3143	1/1	0.88	0.17	-	57,57,57,57	0
57	MG	14	3199	1/1	0.97	0.38	-	76,76,76,76	0
57	MG	13	1664	1/1	0.80	0.46	-	78,78,78,78	0
57	MG	1H	3136	1/1	0.84	0.41	-	116,116,116,116	0
57	MG	1H	3420	1/1	0.84	0.12	-	69,69,69,69	0
57	MG	14	3365	1/1	0.93	0.13	-	78,78,78,78	0
57	MG	1G	1609	1/1	0.49	0.87	-	92,92,92,92	0
57	MG	1G	1681	1/1	0.93	0.06	-	96,96,96,96	0
57	MG	14	3364	1/1	0.96	0.07	-	92,92,92,92	0
57	MG	13	1708	1/1	0.90	0.16	-	65,65,65,65	0
57	MG	13	1671	1/1	0.66	0.51	-	79,79,79,79	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	1H	3230	1/1	0.60	0.36	-	82,82,82,82	0
57	MG	1H	3043	1/1	0.81	0.40	-	75,75,75,75	0
57	MG	1H	3071	1/1	0.83	0.22	-	79,79,79,79	0
57	MG	1H	3217	1/1	0.75	0.64	-	82,82,82,82	0
57	MG	1H	3219	1/1	0.92	0.20	-	71,71,71,71	0
57	MG	1H	3478	1/1	0.93	0.05	-	101,101,101,101	0
57	MG	14	3431	1/1	0.88	0.25	-	116,116,116,116	0
57	MG	13	1720	1/1	0.91	0.15	-	109,109,109,109	0
57	MG	14	3201	1/1	0.91	0.20	-	88,88,88,88	0
57	MG	1H	3309	1/1	0.78	0.23	-	54,54,54,54	0
57	MG	14	3185	1/1	0.73	0.41	-	86,86,86,86	0
57	MG	14	3019	1/1	0.98	0.35	-	57,57,57,57	0
57	MG	14	3096	1/1	0.96	0.20	-	65,65,65,65	0
57	MG	14	3377	1/1	0.47	0.10	-	131,131,131,131	0
57	MG	14	3329	1/1	0.95	0.06	-	105,105,105,105	0
57	MG	1H	3408	1/1	0.82	0.12	-	67,67,67,67	0
57	MG	14	3154	1/1	0.58	0.40	-	95,95,95,95	0
57	MG	1H	3403	1/1	0.93	0.07	-	99,99,99,99	0
57	MG	14	3162	1/1	0.94	0.40	-	74,74,74,74	0
57	MG	14	3351	1/1	0.95	0.11	-	78,78,78,78	0
57	MG	88	203	1/1	0.83	0.33	-	79,79,79,79	0
57	MG	1H	3441	1/1	0.94	0.06	-	105,105,105,105	0
57	MG	14	3082	1/1	0.96	0.49	-	79,79,79,79	0
57	MG	14	3366	1/1	0.73	0.23	-	82,82,82,82	0
57	MG	13	1737	1/1	0.73	0.11	-	99,99,99,99	0
57	MG	14	3370	1/1	0.95	0.06	-	97,97,97,97	0
57	MG	1H	3086	1/1	0.89	0.18	-	53,53,53,53	0
57	MG	14	3243	1/1	0.83	0.15	-	58,58,58,58	0
57	MG	1H	3522	1/1	0.75	0.14	-	110,110,110,110	0
57	MG	13	1681	1/1	0.94	0.14	-	59,59,59,59	0
57	MG	1H	3119	1/1	0.82	0.39	-	83,83,83,83	0
57	MG	13	1677	1/1	0.91	0.47	-	77,77,77,77	0
57	MG	1H	3486	1/1	0.81	0.07	-	95,95,95,95	0
57	MG	1H	3177	1/1	0.70	0.90	-	82,82,82,82	0
57	MG	1H	3524	1/1	0.92	0.18	-	81,81,81,81	0
57	MG	21	303	1/1	0.97	0.09	-	46,46,46,46	0
57	MG	1H	3447	1/1	0.91	0.24	-	70,70,70,70	0
57	MG	1H	3439	1/1	0.97	0.07	-	67,67,67,67	0
57	MG	14	3200	1/1	0.87	0.15	-	89,89,89,89	0
57	MG	1G	1627	1/1	0.98	0.41	-	87,87,87,87	0
57	MG	1G	1682	1/1	0.72	0.05	-	123,123,123,123	0
57	MG	14	3174	1/1	0.95	0.08	-	89,89,89,89	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	14	3055	1/1	0.99	0.30	-	60,60,60,60	0
57	MG	1H	3395	1/1	0.75	0.14	-	68,68,68,68	0
57	MG	13	1605	1/1	0.97	0.20	-	71,71,71,71	0
57	MG	1H	3084	1/1	0.84	0.20	-	62,62,62,62	0
57	MG	14	3104	1/1	0.94	0.18	-	78,78,78,78	0
57	MG	14	3341	1/1	0.98	0.12	-	66,66,66,66	0
57	MG	1H	3483	1/1	0.84	0.17	-	91,91,91,91	0
57	MG	14	3190	1/1	0.92	0.37	-	59,59,59,59	0
57	MG	1G	1680	1/1	0.79	0.07	-	123,123,123,123	0
57	MG	29	301	1/1	0.98	0.35	-	58,58,58,58	0
57	MG	14	3348	1/1	0.78	0.11	-	96,96,96,96	0
57	MG	1H	3097	1/1	0.98	0.06	-	65,65,65,65	0
57	MG	1H	3149	1/1	0.89	0.24	-	55,55,55,55	0
57	MG	88	202	1/1	0.90	0.32	-	66,66,66,66	0
57	MG	13	1611	1/1	0.95	0.24	-	70,70,70,70	0
57	MG	1H	3473	1/1	0.94	0.14	-	95,95,95,95	0
57	MG	1H	3347	1/1	0.83	0.10	-	67,67,67,67	0
57	MG	14	3306	1/1	0.92	0.08	-	58,58,58,58	0
57	MG	1H	3346	1/1	0.85	0.10	-	92,92,92,92	0
57	MG	1H	3195	1/1	0.72	0.31	-	55,55,55,55	0
57	MG	1H	3150	1/1	0.61	0.70	-	98,98,98,98	0
57	MG	1H	3242	1/1	0.85	0.47	-	90,90,90,90	0
57	MG	14	3326	1/1	0.78	0.09	-	77,77,77,77	0
57	MG	1H	3490	1/1	0.90	0.07	-	89,89,89,89	0
57	MG	14	3036	1/1	0.95	0.27	-	85,85,85,85	0
57	MG	I8	102	1/1	0.95	0.05	-	62,62,62,62	0
57	MG	14	3263	1/1	0.90	0.08	-	71,71,71,71	0
57	MG	45	202	1/1	0.91	0.44	-	64,64,64,64	0
57	MG	14	3314	1/1	0.96	0.16	-	69,69,69,69	0
57	MG	1H	3430	1/1	0.93	0.08	-	73,73,73,73	0
57	MG	13	1649	1/1	0.82	0.42	-	75,75,75,75	0
57	MG	14	3281	1/1	0.95	0.16	-	69,69,69,69	0
57	MG	14	3057	1/1	0.90	0.25	-	49,49,49,49	0
57	MG	14	3095	1/1	0.91	0.18	-	62,62,62,62	0
57	MG	14	3112	1/1	0.93	0.20	-	83,83,83,83	0
57	MG	1H	3283	1/1	0.94	0.15	-	71,71,71,71	0
57	MG	1H	3091	1/1	0.99	0.15	-	41,41,41,41	0
57	MG	1H	3454	1/1	0.54	0.07	-	111,111,111,111	0
57	MG	14	3413	1/1	0.88	0.15	-	107,107,107,107	0
57	MG	14	3313	1/1	0.91	0.05	-	105,105,105,105	0
57	MG	14	3152	1/1	0.66	0.17	-	78,78,78,78	0
57	MG	16	211	1/1	0.97	0.11	-	73,73,73,73	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	14	3120	1/1	0.90	0.18	-	78,78,78,78	0
57	MG	1H	3358	1/1	0.94	0.18	-	45,45,45,45	0
57	MG	14	3294	1/1	0.98	0.05	-	75,75,75,75	0
57	MG	14	3063	1/1	0.84	0.24	-	67,67,67,67	0
57	MG	13	1712	1/1	0.88	0.07	-	89,89,89,89	0
57	MG	1H	3327	1/1	0.98	0.08	-	50,50,50,50	0
57	MG	1H	3506	1/1	0.84	0.08	-	106,106,106,106	0
57	MG	1H	3268	1/1	0.90	0.17	-	59,59,59,59	0
57	MG	1H	3459	1/1	0.95	0.13	-	75,75,75,75	0
57	MG	I8	101	1/1	0.86	0.09	-	85,85,85,85	0
57	MG	1H	3057	1/1	0.88	0.38	-	75,75,75,75	0
57	MG	1H	3519	1/1	0.92	0.18	-	116,116,116,116	0
57	MG	13	1630	1/1	0.99	0.34	-	96,96,96,96	0
57	MG	1H	3339	1/1	0.61	0.12	-	106,106,106,106	0
57	MG	1H	3141	1/1	0.59	0.29	-	83,83,83,83	0
57	MG	14	3276	1/1	0.79	0.10	-	120,120,120,120	0
57	MG	1H	3460	1/1	0.89	0.06	-	96,96,96,96	0
57	MG	1G	1650	1/1	0.93	0.16	-	101,101,101,101	0
57	MG	1H	3173	1/1	0.94	0.22	-	59,59,59,59	0
57	MG	14	3426	1/1	0.65	0.16	-	104,104,104,104	0
57	MG	1G	1676	1/1	0.89	0.09	-	93,93,93,93	0
57	MG	1H	3260	1/1	0.95	0.19	-	42,42,42,42	0
57	MG	1J	208	1/1	0.65	0.10	-	103,103,103,103	0
57	MG	1H	3383	1/1	0.86	0.10	-	68,68,68,68	0
57	MG	13	1620	1/1	0.88	0.52	-	86,86,86,86	0
57	MG	1G	1611	1/1	0.87	0.65	-	76,76,76,76	0
57	MG	1H	3432	1/1	0.92	0.11	-	93,93,93,93	0
57	MG	1H	3390	1/1	0.92	0.20	-	64,64,64,64	0
57	MG	1H	3353	1/1	0.97	0.14	-	48,48,48,48	0
57	MG	1H	3201	1/1	0.82	0.74	-	104,104,104,104	0
57	MG	14	3184	1/1	0.83	0.26	-	78,78,78,78	0
57	MG	1H	3249	1/1	0.96	0.13	-	63,63,63,63	0
57	MG	14	3303	1/1	0.92	0.07	-	74,74,74,74	0
57	MG	1H	3088	1/1	0.90	0.42	-	80,80,80,80	0
57	MG	1H	3070	1/1	0.96	0.34	-	75,75,75,75	0
57	MG	1H	3320	1/1	0.86	0.08	-	58,58,58,58	0
57	MG	14	3356	1/1	0.98	0.06	-	75,75,75,75	0
57	MG	14	3336	1/1	0.87	0.09	-	94,94,94,94	0
57	MG	1G	1664	1/1	0.89	0.05	-	80,80,80,80	0
57	MG	1H	3112	1/1	0.96	0.26	-	55,55,55,55	0
57	MG	1H	3377	1/1	0.95	0.09	-	76,76,76,76	0
57	MG	14	3432	1/1	0.76	0.13	-	116,116,116,116	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	14	3232	1/1	0.94	0.14	-	66,66,66,66	0
57	MG	13	1674	1/1	0.74	0.36	-	92,92,92,92	0
57	MG	1H	3425	1/1	0.92	0.07	-	64,64,64,64	0
57	MG	1H	3288	1/1	0.94	0.08	-	54,54,54,54	0
57	MG	14	3186	1/1	0.90	0.16	-	108,108,108,108	0
57	MG	13	1610	1/1	0.87	0.38	-	82,82,82,82	0
57	MG	1G	1699	1/1	0.75	0.07	-	133,133,133,133	0
57	MG	1H	3240	1/1	0.86	0.38	-	73,73,73,73	0
57	MG	1H	3198	1/1	0.79	0.23	-	79,79,79,79	0
57	MG	1H	3350	1/1	0.80	0.11	-	74,74,74,74	0
57	MG	13	1688	1/1	0.95	0.10	-	109,109,109,109	0
57	MG	14	3202	1/1	0.81	0.49	-	84,84,84,84	0
57	MG	1H	3517	1/1	0.83	0.07	-	91,91,91,91	0
57	MG	1G	1701	1/1	0.92	0.03	-	111,111,111,111	0
57	MG	1G	1660	1/1	0.92	0.14	-	74,74,74,74	0
57	MG	14	3132	1/1	0.61	0.44	-	82,82,82,82	0
57	MG	14	3318	1/1	0.85	0.12	-	89,89,89,89	0
57	MG	1H	3099	1/1	0.94	0.44	-	56,56,56,56	0
57	MG	13	1723	1/1	0.66	0.07	-	128,128,128,128	0
57	MG	14	3149	1/1	0.92	0.28	-	72,72,72,72	0
57	MG	1G	1640	1/1	0.92	0.39	-	108,108,108,108	0
57	MG	1H	3190	1/1	0.91	0.38	-	91,91,91,91	0
57	MG	1H	3343	1/1	0.93	0.07	-	78,78,78,78	0
57	MG	1H	3109	1/1	0.90	0.31	-	59,59,59,59	0
57	MG	1H	3073	1/1	0.92	0.21	-	66,66,66,66	0
57	MG	1H	3282	1/1	0.95	0.04	-	71,71,71,71	0
57	MG	1H	3025	1/1	0.96	0.35	-	75,75,75,75	0
57	MG	31	302	1/1	0.95	0.16	-	77,77,77,77	0
57	MG	14	3398	1/1	0.76	0.14	-	76,76,76,76	0
57	MG	2K	102	1/1	0.95	0.17	-	78,78,78,78	0
57	MG	1H	3131	1/1	0.94	0.50	-	83,83,83,83	0
57	MG	1H	3495	1/1	0.96	0.05	-	97,97,97,97	0
57	MG	13	1643	1/1	0.59	0.23	-	101,101,101,101	0
57	MG	1H	3162	1/1	0.83	0.29	-	60,60,60,60	0
57	MG	1G	1601	1/1	0.97	0.20	-	77,77,77,77	0
57	MG	14	3304	1/1	0.94	0.10	-	74,74,74,74	0
57	MG	1H	3127	1/1	0.90	0.12	-	91,91,91,91	0
57	MG	1G	1656	1/1	0.87	0.08	-	100,100,100,100	0
57	MG	1H	3176	1/1	0.83	0.14	-	83,83,83,83	0
57	MG	13	1724	1/1	0.84	0.11	-	117,117,117,117	0
57	MG	13	1662	1/1	0.69	0.50	-	89,89,89,89	0
57	MG	1H	3147	1/1	0.84	0.39	-	84,84,84,84	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(Å^2)	Q<0.9
57	MG	14	3350	1/1	0.39	0.16	-	101,101,101,101	0
57	MG	1H	3207	1/1	0.96	0.24	-	92,92,92,92	0
57	MG	14	3420	1/1	0.90	0.07	-	107,107,107,107	0
57	MG	1G	1618	1/1	0.77	0.69	-	78,78,78,78	0
57	MG	13	1709	1/1	0.54	0.26	-	99,99,99,99	0
57	MG	14	3141	1/1	0.92	0.28	-	95,95,95,95	0
57	MG	14	3369	1/1	0.91	0.10	-	91,91,91,91	0
57	MG	14	3361	1/1	0.92	0.05	-	79,79,79,79	0
57	MG	1H	3419	1/1	0.70	0.15	-	86,86,86,86	0
57	MG	13	1653	1/1	0.94	0.33	-	73,73,73,73	0
57	MG	14	3166	1/1	0.65	0.46	-	79,79,79,79	0
57	MG	13	1739	1/1	0.59	0.06	-	121,121,121,121	0
57	MG	13	1699	1/1	0.96	0.09	-	83,83,83,83	0
57	MG	4L	101	1/1	0.74	0.32	-	131,131,131,131	0
57	MG	1H	3452	1/1	0.88	0.10	-	72,72,72,72	0
57	MG	7A	101	1/1	0.73	0.26	-	114,114,114,114	0
57	MG	1H	3251	1/1	0.91	0.16	-	48,48,48,48	0
57	MG	14	3405	1/1	0.90	0.07	-	108,108,108,108	0
57	MG	1G	1675	1/1	0.76	0.08	-	103,103,103,103	0
57	MG	1H	3458	1/1	0.73	0.10	-	82,82,82,82	0
57	MG	14	3222	1/1	0.98	0.20	-	59,59,59,59	0
57	MG	14	3060	1/1	0.77	0.30	-	98,98,98,98	0
57	MG	1G	1677	1/1	0.93	0.17	-	92,92,92,92	0
57	MG	13	1731	1/1	0.88	0.05	-	112,112,112,112	0
57	MG	14	3429	1/1	0.67	0.30	-	95,95,95,95	0

6.5 Other polymers (i)

There are no such residues in this entry.