



Full wwPDB X-ray Structure Validation Report ⓘ

Dec 16, 2024 – 09:31 AM EST

PDB ID : 6OSI
Title : Unmodified tRNA(Pro) bound to Thermus thermophilus 70S (near cognate)
Authors : Hoffer, E.D.; Subaramanian, S.; Hong, S.; Maehigashi, T.; Dunham, C.M.
Deposited on : 2019-05-01
Resolution : 4.14 Å(reported)

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

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

A user guide is available at

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

with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

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

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

MolProbity : 4.02b-467
Mogul : 2022.3.0, CSD as543be (2022)
Xtriage (Phenix) : 1.21
EDS : 3.0
Percentile statistics : 20231227.v01 (using entries in the PDB archive December 27th 2023)
CCP4 : 9.0.004 (Gargrove)
Density-Fitness : 1.0.11
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.40

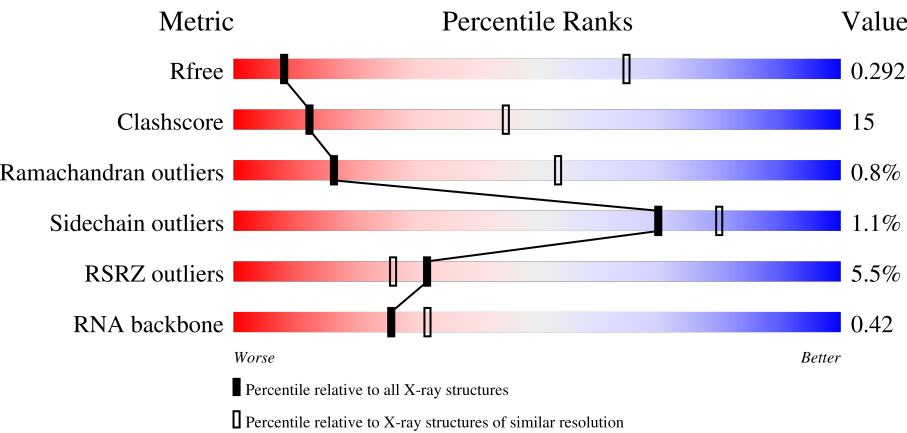
1 Overall quality at a glance i

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

The reported resolution of this entry is 4.14 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}	164625	1166 (4.48-3.80)
Clashscore	180529	1233 (4.48-3.80)
Ramachandran outliers	177936	1158 (4.48-3.80)
Sidechain outliers	177891	1145 (4.48-3.80)
RSRZ outliers	164620	1164 (4.48-3.80)
RNA backbone	3690	1145 (5.04-3.00)

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

Mol	Chain	Length	Quality of chain
1	QA	1521	<div><div>24%48%23%</div><div></div></div>
1	XA	1521	<div><div>28%44%21%7%</div><div></div></div>
2	QB	256	<div><div>4%64%28%8%</div><div></div></div>
2	XB	256	<div><div>7%69%23%8%</div><div></div></div>

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Mol	Chain	Length	Quality of chain
3	QC	239	
3	XC	239	
4	QD	209	
4	XD	209	
5	QE	162	
5	XE	162	
6	QF	101	
6	XF	101	
7	QG	156	
7	XG	156	
8	QH	138	
8	XH	138	
9	QI	128	
9	XI	128	
10	QJ	105	
10	XJ	105	
11	QK	129	
11	XK	129	
12	QL	132	
12	XL	132	
13	QM	126	
13	XM	126	
14	QN	61	
14	XN	61	
15	QO	89	

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Mol	Chain	Length	Quality of chain
15	XO	89	
16	QP	88	
16	XP	88	
17	QQ	105	
17	XQ	105	
18	QR	88	
18	XR	88	
19	QS	93	
19	XS	93	
20	QT	106	
20	XT	106	
21	QU	27	
21	XU	27	
22	QV	77	
22	XV	77	
23	QX	19	
23	XX	19	
24	R0	85	
24	Y0	85	
25	R1	98	
25	Y1	98	
26	R2	72	
26	Y2	72	
27	R3	60	
27	Y3	60	

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Mol	Chain	Length	Quality of chain
28	R4	71	
28	Y4	71	
29	R5	60	
29	Y5	60	
30	R6	54	
30	Y6	54	
31	R7	49	
31	Y7	49	
32	R8	65	
32	Y8	65	
33	R9	37	
33	Y9	37	
34	RA	2905	
34	YA	2905	
35	RB	122	
35	YB	122	
36	RD	276	
36	YD	276	
37	RE	206	
37	YE	206	
38	RF	210	
38	YF	210	
39	RG	182	
39	YG	182	
40	RH	180	

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Mol	Chain	Length	Quality of chain
40	YH	180	
41	RI	148	
41	YI	148	
42	RN	140	
42	YN	140	
43	RO	122	
43	YO	122	
44	RP	150	
44	YP	150	
45	RQ	141	
45	YQ	141	
46	RR	118	
46	YR	118	
47	RS	112	
47	YS	112	
48	RT	146	
48	YT	146	
49	RU	118	
49	YU	118	
50	RV	101	
50	YV	101	
51	RW	113	
51	YW	113	
52	RX	96	
52	YX	96	

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Mol	Chain	Length	Quality of chain
53	RY	110	
53	YY	110	
54	RZ	206	
54	YZ	206	

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

Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
55	MG	QA	1622	-	-	-	X
55	MG	QE	201	-	-	-	X
55	MG	RA	2912	-	-	-	X
55	MG	RA	2921	-	-	-	X
55	MG	RA	2954	-	-	-	X
55	MG	RA	2970	-	-	-	X
55	MG	RA	2975	-	-	-	X
55	MG	RA	2976	-	-	-	X
55	MG	RA	3013	-	-	-	X
55	MG	RA	3051	-	-	-	X
55	MG	RA	3057	-	-	-	X
55	MG	RA	3064	-	-	-	X
55	MG	RA	3071	-	-	-	X
55	MG	RA	3075	-	-	-	X
55	MG	RA	3078	-	-	-	X
55	MG	RA	3083	-	-	-	X
55	MG	RA	3101	-	-	-	X
55	MG	RA	3103	-	-	-	X
55	MG	RA	3133	-	-	-	X
55	MG	RA	3139	-	-	-	X
55	MG	RA	3144	-	-	-	X
55	MG	RA	3147	-	-	-	X
55	MG	RA	3152	-	-	-	X
55	MG	RA	3162	-	-	-	X
55	MG	RA	3195	-	-	-	X
55	MG	RA	3220	-	-	-	X
55	MG	RA	3221	-	-	-	X
55	MG	RA	3223	-	-	-	X
55	MG	RA	3260	-	-	-	X
55	MG	RA	3286	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
55	MG	RA	3307	-	-	-	X
55	MG	RA	3310	-	-	-	X
55	MG	RA	3318	-	-	-	X
55	MG	RE	304	-	-	-	X
55	MG	XA	1605	-	-	-	X
55	MG	XA	1610	-	-	-	X
55	MG	XA	1612	-	-	-	X
55	MG	XA	1613	-	-	-	X
55	MG	XA	1621	-	-	-	X
55	MG	XA	1657	-	-	-	X
55	MG	XA	1665	-	-	-	X
55	MG	XA	1688	-	-	X	-
55	MG	Y8	101	-	-	-	X
55	MG	YA	2904	-	-	-	X
55	MG	YA	2905	-	-	-	X
55	MG	YA	2907	-	-	-	X
55	MG	YA	2921	-	-	-	X
55	MG	YA	2930	-	-	-	X
55	MG	YA	2931	-	-	-	X
55	MG	YA	2932	-	-	-	X
55	MG	YA	2933	-	-	-	X
55	MG	YA	2934	-	-	-	X
55	MG	YA	2936	-	-	-	X
55	MG	YA	2952	-	-	-	X
55	MG	YA	2953	-	-	-	X
55	MG	YA	2954	-	-	-	X
55	MG	YA	2955	-	-	-	X
55	MG	YA	2958	-	-	-	X
55	MG	YA	2962	-	-	-	X
55	MG	YA	2974	-	-	-	X
55	MG	YA	2983	-	-	-	X
55	MG	YA	2984	-	-	-	X
55	MG	YA	3004	-	-	-	X
55	MG	YA	3010	-	-	-	X
55	MG	YA	3017	-	-	-	X
55	MG	YA	3018	-	-	-	X
55	MG	YA	3087	-	-	-	X
55	MG	YA	3097	-	-	-	X
55	MG	YA	3101	-	-	-	X
55	MG	YA	3117	-	-	-	X
55	MG	YA	3120	-	-	-	X
55	MG	YA	3122	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
55	MG	YA	3126	-	-	-	X
55	MG	YA	3153	-	-	-	X
55	MG	YA	3168	-	-	-	X
55	MG	YA	3175	-	-	-	X
55	MG	YA	3192	-	-	X	-
55	MG	YA	3194	-	-	-	X
55	MG	YA	3205	-	-	-	X
55	MG	YA	3210	-	-	-	X
55	MG	YA	3211	-	-	-	X
55	MG	YA	3237	-	-	-	X
55	MG	YA	3239	-	-	-	X
55	MG	YA	3245	-	-	-	X
55	MG	YA	3262	-	-	-	X
55	MG	YA	3276	-	-	-	X
55	MG	YA	3281	-	-	-	X
55	MG	YA	3288	-	-	-	X
55	MG	YA	3290	-	-	-	X
56	SF4	QD	301	-	-	X	-

2 Entry composition

There are 57 unique types of molecules in this entry. The entry contains 291185 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 rRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
1	QA	1511	Total	C	N	O	P	0	0	0
			32469	14453	6011	10495	1510			
1	XA	1515	Total	C	N	O	P	0	0	0
			32551	14490	6022	10525	1514			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
2	QB	235	Total	C	N	O	S	0	0	0
			1907	1217	342	343	5			
2	XB	236	Total	C	N	O	S	0	0	0
			1915	1223	343	344	5			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
3	QC	205	Total	C	N	O	S	0	0	0
			1605	1011	313	280	1			
3	XC	205	Total	C	N	O	S	0	0	0
			1605	1011	313	280	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
4	QD	208	Total	C	N	O	S	0	0	0
			1703	1066	339	291	7			
4	XD	208	Total	C	N	O	S	0	0	0
			1703	1066	339	291	7			

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

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

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
6	QF	101	Total	C	N	O	S	0	0	0
			843	531	155	154	3			
6	XF	101	Total	C	N	O	S	0	0	0
			843	531	155	154	3			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
7	QG	155	Total	C	N	O	S	0	0	0
			1257	781	252	218	6			
7	XG	155	Total	C	N	O	S	0	0	0
			1257	781	252	218	6			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
8	QH	137	Total	C	N	O	S	0	0	0
			1108	700	214	192	2			
8	XH	137	Total	C	N	O	S	0	0	0
			1108	700	214	192	2			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
9	QI	105	Total	C	N	O		0	0	0
			816	519	152	145				
9	XI	107	Total	C	N	O		0	0	0
			834	530	157	147				

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

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

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
10	XJ	96	Total	C	N	O	S	0	0	0
			777	487	153	136	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
11	QK	119	Total	C	N	O	S	0	0	0
			885	549	168	165	3			
11	XK	116	Total	C	N	O	S	0	0	0
			864	537	164	160	3			

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

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

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
13	QM	115	Total	C	N	O	S	0	0	0
			921	569	190	160	2			
13	XM	114	Total	C	N	O	S	0	0	0
			914	565	189	158	2			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
14	QN	60	Total	C	N	O	S	0	0	0
			492	312	104	72	4			
14	XN	60	Total	C	N	O	S	0	0	0
			492	312	104	72	4			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
15	QO	88	Total	C	N	O	S	0	0	0
			734	459	147	126	2			
15	XO	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	QP	84	Total	C	N	O	S	0	0	0
			705	446	140	118	1			
16	XP	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	QQ	100	Total	C	N	O	S	0	0	0
			834	534	155	143	2			
17	XQ	100	Total	C	N	O	S	0	0	0
			834	534	155	143	2			

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

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
18	QR	70	Total	C	N	O	0	0	0
			574	367	112	95			
18	XR	70	Total	C	N	O	0	0	0
			574	367	112	95			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
19	QS	83	Total	C	N	O	S	0	0	0
			665	424	124	115	2			
19	XS	84	Total	C	N	O	S	0	0	0
			674	430	126	116	2			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
20	QT	99	Total	C	N	O	S	0	0	0
			763	470	162	129	2			
20	XT	99	Total	C	N	O	S	0	0	0
			763	470	162	129	2			

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

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
21	QU	25	Total	C	N	O	0	0	0
			217	134	52	31			
21	XU	25	Total	C	N	O	0	0	0
			217	134	52	31			

- Molecule 22 is a RNA chain called tRNA(Pro).

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
22	QV	68	Total	C	N	O	P	0	0	0
			1452	647	260	477	68			
22	XV	68	Total	C	N	O	P	0	0	0
			1452	647	260	477	68			

- Molecule 23 is a RNA chain called mRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
23	QX	19	Total	C	N	O	P	0	0	0
			409	184	81	126	18			
23	XX	19	Total	C	N	O	P	0	0	0
			409	184	81	126	18			

- Molecule 24 is a protein called 50S ribosomal protein L27.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
24	R0	81	Total	C	N	O	S	0	0	0
			643	398	137	107	1			
24	Y0	82	Total	C	N	O	S	0	0	0
			648	401	138	108	1			

- Molecule 25 is a protein called 50S ribosomal protein L28.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
25	R1	95	Total	C	N	O	S	0	0	0
			746	469	148	128	1			
25	Y1	93	Total	C	N	O	S	0	0	0
			729	457	145	126	1			

- Molecule 26 is a protein called 50S ribosomal protein L29.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
26	R2	69	Total	C	N	O	S	0	0	0
			581	358	118	104	1			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
26	Y2	68	Total	C	N	O	S	0	0	0
			575	355	117	102	1			

- Molecule 27 is a protein called 50S ribosomal protein L30.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
27	R3	59	Total	C	N	O		0	0	0
			469	298	90	81				
27	Y3	59	Total	C	N	O		0	0	0
			469	298	90	81				

- Molecule 28 is a protein called 50S ribosomal protein L31.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
28	R4	45	Total	C	N	O	S	0	0	0
			348	224	57	62	5			
28	Y4	46	Total	C	N	O	S	0	0	0
			357	229	59	64	5			

- Molecule 29 is a protein called 50S ribosomal protein L32.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
29	R5	59	Total	C	N	O	S	0	0	0
			459	288	90	76	5			
29	Y5	59	Total	C	N	O	S	0	0	0
			459	288	90	76	5			

- Molecule 30 is a protein called 50S ribosomal protein L33.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
30	R6	53	Total	C	N	O	S	0	0	0
			453	281	91	77	4			
30	Y6	53	Total	C	N	O	S	0	0	0
			453	281	91	77	4			

- Molecule 31 is a protein called 50S ribosomal protein L34.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
31	R7	47	Total	C	N	O	S	0	0	0
			409	251	102	54	2			
31	Y7	48	Total	C	N	O	S	0	0	0
			418	257	104	55	2			

- Molecule 32 is a protein called 50S ribosomal protein L35.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
32	R8	64	Total	C	N	O	S	0	0	0
			517	331	102	82	2			
32	Y8	64	Total	C	N	O	S	0	0	0
			517	331	102	82	2			

- Molecule 33 is a protein called 50S ribosomal protein L36.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
33	R9	37	Total	C	N	O	S	0	0	0
			307	188	68	47	4			
33	Y9	37	Total	C	N	O	S	0	0	0
			307	188	68	47	4			

- Molecule 34 is a RNA chain called 23S rRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
34	RA	2882	Total	C	N	O	P	0	0	0
			62070	27627	11611	19951	2881			
34	YA	2883	Total	C	N	O	P	0	0	0
			62091	27636	11613	19960	2882			

- Molecule 35 is a RNA chain called 5S rRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
35	RB	120	Total	C	N	O	P	0	0	0
			2573	1146	476	832	119			
35	YB	120	Total	C	N	O	P	0	0	0
			2573	1146	476	832	119			

- Molecule 36 is a protein called 50S ribosomal protein L2.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
36	RD	272	Total	C	N	O	S	0	0	0
			2115	1335	420	357	3			
36	YD	272	Total	C	N	O	S	0	0	0
			2115	1335	420	357	3			

- Molecule 37 is a protein called 50S ribosomal protein L3.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
37	RE	205	Total	C	N	O	S	0	0	0
			1568	991	300	271	6			
37	YE	205	Total	C	N	O	S	0	0	0
			1568	991	300	271	6			

- Molecule 38 is a protein called 50S ribosomal protein L4.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
38	RF	202	Total	C	N	O	S	0	0	0
			1585	1011	297	275	2			
38	YF	202	Total	C	N	O	S	0	0	0
			1585	1011	297	275	2			

- Molecule 39 is a protein called 50S ribosomal protein L5.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
39	RG	181	Total	C	N	O	S	0	0	0
			1474	942	268	260	4			
39	YG	181	Total	C	N	O	S	0	0	0
			1474	942	268	260	4			

- Molecule 40 is a protein called 50S ribosomal protein L6.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
40	RH	174	Total	C	N	O	S	0	0	0
			1336	848	251	236	1			
40	YH	174	Total	C	N	O	S	0	0	0
			1336	848	251	236	1			

- Molecule 41 is a protein called 50S ribosomal protein L9.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
41	RI	146	Total	C	N	O	S	0	0	0
			1136	726	201	208	1			
41	YI	146	Total	C	N	O	S	0	0	0
			1136	726	201	208	1			

- Molecule 42 is a protein called 50S ribosomal protein L13.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
42	RN	138	Total	C	N	O	S	0	0	0
			1104	712	206	182	4			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
42	YN	138	Total	C	N	O	S	0	0	0
			1104	712	206	182	4			

- Molecule 43 is a protein called 50S ribosomal protein L14.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
43	RO	122	Total	C	N	O	S	0	0	0
			933	588	171	170	4			
43	YO	122	Total	C	N	O	S	0	0	0
			933	588	171	170	4			

- Molecule 44 is a protein called 50S ribosomal protein L15.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
44	RP	150	Total	C	N	O	S	0	0	0
			1145	712	232	198	3			
44	YP	147	Total	C	N	O	S	0	0	0
			1122	698	229	192	3			

- Molecule 45 is a protein called 50S ribosomal protein L16.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
45	RQ	141	Total	C	N	O	S	0	0	0
			1122	715	212	188	7			
45	YQ	141	Total	C	N	O	S	0	0	0
			1122	715	212	188	7			

- Molecule 46 is a protein called 50S ribosomal protein L17.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
46	RR	117	Total	C	N	O	0	0	0
			960	599	202	159			
46	YR	117	Total	C	N	O	0	0	0
			960	599	202	159			

- Molecule 47 is a protein called 50S ribosomal protein L18.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
47	RS	111	Total	C	N	O	0	0	0
			882	556	176	150			
47	YS	111	Total	C	N	O	0	0	0
			882	556	176	150			

- Molecule 48 is a protein called 50S ribosomal protein L19.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
48	RT	137	Total	C	N	O	S	0	0	0
			1141	710	234	196	1			
48	YT	137	Total	C	N	O	S	0	0	0
			1141	710	234	196	1			

- Molecule 49 is a protein called 50S ribosomal protein L20.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
49	RU	117	Total	C	N	O	S	0	0	0
			964	610	202	151	1			
49	YU	117	Total	C	N	O	S	0	0	0
			964	610	202	151	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
50	RV	101	Total	C	N	O	S	0	0	0
			779	501	142	135	1			
50	YV	101	Total	C	N	O	S	0	0	0
			779	501	142	135	1			

- Molecule 51 is a protein called 50S ribosomal protein L22.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
51	RW	113	Total	C	N	O	S	0	0	0
			900	566	177	155	2			
51	YW	113	Total	C	N	O	S	0	0	0
			900	566	177	155	2			

- Molecule 52 is a protein called 50S ribosomal protein L23.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
52	RX	92	Total	C	N	O	0	0	0
			725	471	131	123			
52	YX	92	Total	C	N	O	0	0	0
			725	471	131	123			

- Molecule 53 is a protein called 50S ribosomal protein L24.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
53	RY	107	Total	C	N	O	S	0	0	0
			818	525	155	132	6			
53	YY	107	Total	C	N	O	S	0	0	0
			818	525	155	132	6			

- Molecule 54 is a protein called 50S ribosomal protein L25.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
54	RZ	183	Total	C	N	O	S	0	0	0
			1461	933	260	265	3			
54	YZ	193	Total	C	N	O	S	0	0	0
			1529	973	270	283	3			

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

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
55	QA	70	Total	Mg	0	0
			70	70		
55	QE	1	Total	Mg	0	0
			1	1		
55	QF	1	Total	Mg	0	0
			1	1		
55	QH	2	Total	Mg	0	0
			2	2		
55	QL	2	Total	Mg	0	0
			2	2		
55	R0	2	Total	Mg	0	0
			2	2		
55	R3	1	Total	Mg	0	0
			1	1		
55	R8	2	Total	Mg	0	0
			2	2		
55	RA	432	Total	Mg	0	0
			432	432		
55	RD	1	Total	Mg	0	0
			1	1		
55	RE	4	Total	Mg	0	0
			4	4		
55	RF	2	Total	Mg	0	0
			2	2		
55	RN	1	Total	Mg	0	0
			1	1		
55	RO	1	Total	Mg	0	0
			1	1		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
55	XA	88	Total 88	Mg 88	0	0
55	XE	1	Total 1	Mg 1	0	0
55	XO	1	Total 1	Mg 1	0	0
55	Y1	1	Total 1	Mg 1	0	0
55	Y2	1	Total 1	Mg 1	0	0
55	Y5	1	Total 1	Mg 1	0	0
55	Y7	1	Total 1	Mg 1	0	0
55	Y8	2	Total 2	Mg 2	0	0
55	YA	394	Total 394	Mg 394	0	0
55	YB	1	Total 1	Mg 1	0	0
55	YD	2	Total 2	Mg 2	0	0
55	YE	4	Total 4	Mg 4	0	0
55	YF	1	Total 1	Mg 1	0	0
55	YP	1	Total 1	Mg 1	0	0
55	YQ	1	Total 1	Mg 1	0	0
55	YR	2	Total 2	Mg 2	0	0
55	YU	1	Total 1	Mg 1	0	0
55	YX	1	Total 1	Mg 1	0	0

- Molecule 56 is IRON/SULFUR CLUSTER (three-letter code: SF4) (formula: Fe₄S₄).



Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
56	QD	1	Total	Fe	S	0	0
			8	4	4		
56	XD	1	Total	Fe	S	0	0
			8	4	4		

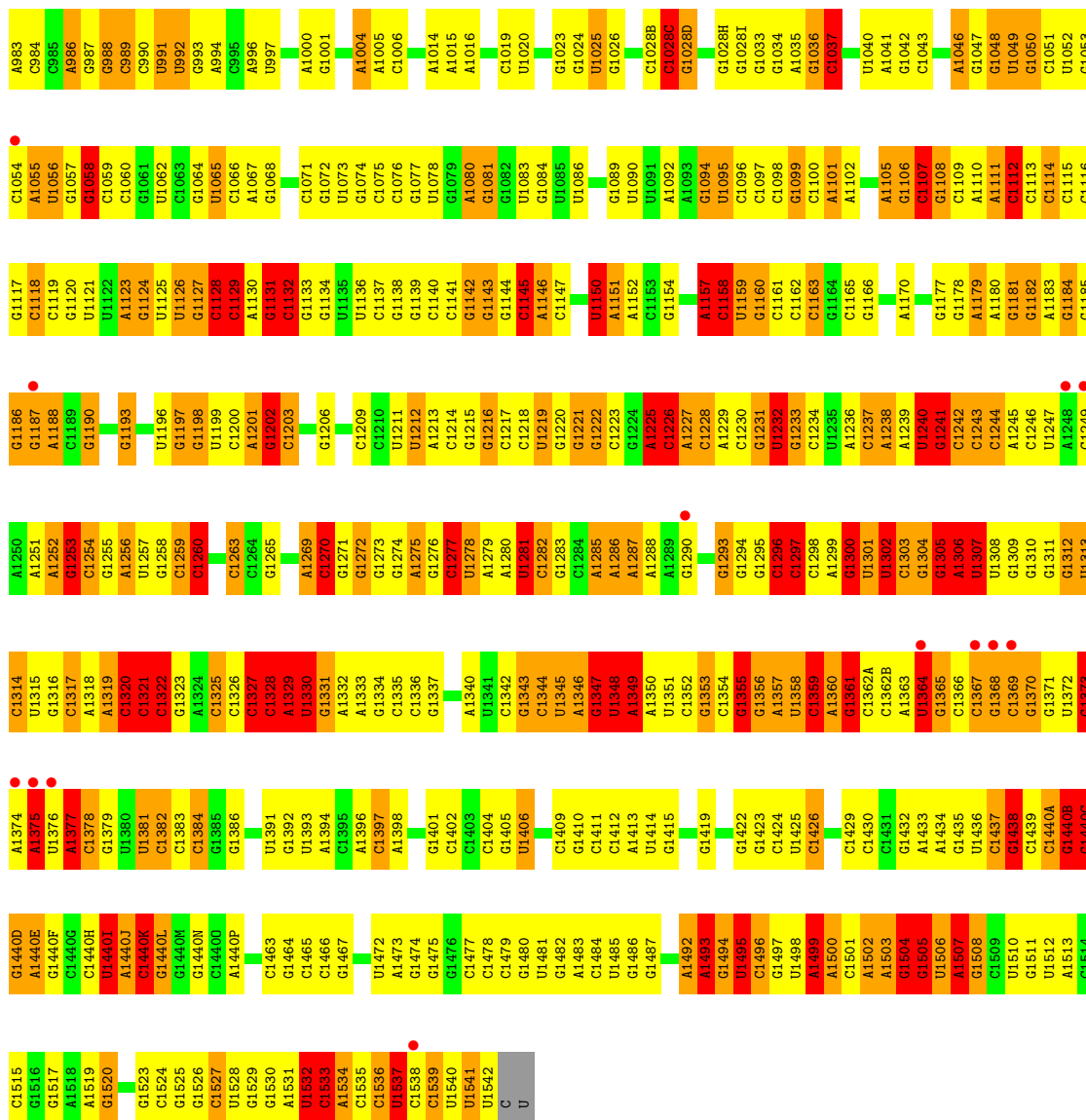
- Molecule 57 is ZINC ION (three-letter code: ZN) (formula: Zn).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	QN	1	Total	Zn	0	0
			1	1		
57	R5	1	Total	Zn	0	0
			1	1		
57	R6	1	Total	Zn	0	0
			1	1		
57	R9	1	Total	Zn	0	0
			1	1		
57	RY	1	Total	Zn	0	0
			1	1		
57	XN	1	Total	Zn	0	0
			1	1		
57	Y5	1	Total	Zn	0	0
			1	1		
57	Y6	1	Total	Zn	0	0
			1	1		
57	Y9	1	Total	Zn	0	0
			1	1		
57	YY	1	Total	Zn	0	0
			1	1		

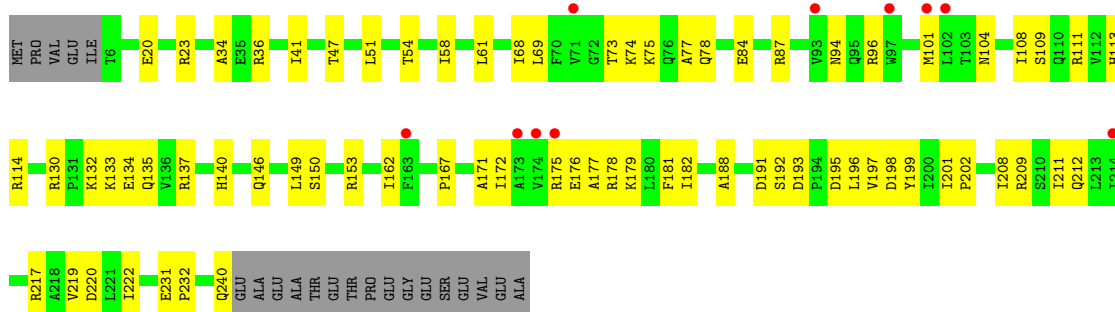
C1533	A1534	A1472	C1395	A1333	G1273	A1213	G1142	U1078	U1012	C980	C810	C748	A684
A1535	A1473	A1474	A1396	G1334	G1274	C1214	G1143	G1079	G1013	C883	C811	C749	G685
C1536	A1475	A1476	C1397	C1335	A1275	G1215	G1144	A1014	A1015	C884	C812	C750	U686
U1537	G1476	G1477	C1398	C1336	G1276	G1216	C1145	G1081	A1016	G885	A814	G751	A687
C1538	G1478	G1479	G1400	G1337	G1277	C1217	C1146	U1085	G1017	A889	A815	G752	G688
C1539	C1477	C1478	G1401	G1338	G1278	G1218	C1147	U1086	U956	A890	A816	G753	C689
U1540	C1479	C1480	A1340	A1339	A1279	U1219	U1148	U1087	G1023	G890	C817	G754	G690
U1541	C1478	C1479	U1341	A1340	A1280	G1220	C1149	U1088	G1024	G891	A818	G755	G691
U1542	G1480	G1481	C1403	U1341	U1281	G1221	U1150	A958	U957	G892	C819	G756	U692
C	C1342	C1343	C1404	C1342	G1282	G1222	A1151	U959	U958	A892	A819	G757	G693
U	U1481	G1343	G1405	G1343	G1283	C1223	A1152	U960	U960	A893	U820	G758	A694
	A1482	U1482	U1406	A1346	G1284	G1224	C1153	U961	C1027	G894	G821	G759	A695
	A1483	C1407	C1407	A1347	A1285	A1225	A1092	U962	G1028A	U897	G822	G760	A696
	C1484	A1408	A1408	G1347	A1286	C1226	G1094	U963	C1028B	C897	G823	G761	U697
	U1485	C1409	C1409	U1348	A1287	A1227	U1095	U964	C1028C	G898	G824	G762	G698
	G1487	G1410	G1410	A1349	C1288	G1228	C1158	U965	G1028D	C899	G825	G763	C699
		C1411	C1411	A1350	A1289	A1229	U1159	C967	G1028E	U905	C826	G764	G700
		C1412	U1351	U1351	G1290	C1230	G1160	U968	G1028F	U906	U827	G765	C701
		A1413	C1352	C1352	G1291	G1231	C1161	A969	A1028G	G906	A828	G766	A702
		U1414	C1353	U1292	U1292	U1232	C1162	U970	G1028H	A907	G829	G767	G703
		G1415	C1354	G1293	G1293	G1233	C1163	U971	U1028I	A908	G830	A768	A704
		G1416	G1355	G1294	G1294	C1234	A1169	C972	C1037	A909	U831	G769	U705
		G1417	G1356	G1295	U1235	U1235	U1170	A974	C1038	C912	U832	G770	A706
		A1418	A1357	C1296	U1236	U1236	G1104	A975	C1039	C913	U833	G771	C707
		G1419	U1358	C1297	C1237	C1237	U1105	G976	U1040	A912	U834	G772	C708
			C1359	C1298	A1238	A1238	G1176	U977	A1041	A913	C834	G773	G709
			G1360	A1299	A1239	A1239	U1177	A978	A1042	A915	G837	G774	G710
			G1361	G1300	U1240	U1240	G1178	A979	G1043	G916	U838	G775	A711
			A1427	U1301	G1241	G1241	C1179	C980	C1044	A918	U839	G776	A712
			C1362B	U1302	G1242	A1179	A1179	C981	A1044	A919	U838B	G777	A713
			C1363	G1303	C1243	A1180	A1111	U981	C1045	U920	U839D	G778	G714
			U1364	G1304	C1244	G1181	C1112	U982	A1046	U921	C848	A715	A715
			G1365	G1305	A1245	G1182	C1113	A983	G1047	G922	C849	A716	A716
			C1367	A1306	C1246	A1183	C1114	C984	U1048	U925	U850	C717	C717
			A1432	U1307	U1247	G1184	C1115	C985	U1049	G926	G851	G718	G718
			G1368	U1308	A1248	G1185	C1116	A986	G1050	G927	G852	C719	C719
			C1369	G1309	C1249	G1186	C1051	G987	G1051	G928	G853	G720	G720
			U1370	G1310	A1250	G1187	U1052	U988	U1052	G929	G854	G721	G721
			G1371	G1311	A1251	A1188	G1053	C989	G1053	G929	C855	A722	A722
			U1372	G1312	A1252	C1189	C1054	U990	U1054	C930	C856	U723	U723
			G1373	U1313	G1253	G1190	A1055	U991	U1055	C931	C857	G724	G724
			A1374	C1314	C1254	A1191	U1056	U992	U1056	G932	G861	G725	G725
			U1375	U1315	G1255	C1195	C1060	A993	C933	G933	C862	C726	C726
			U1376	G1316	A1256	U1196	G1061	A994	C934	C934	U863	G727	G727
			A1377	C1317	U1257	U1197	G1062	C995	A996	A935	U864	A728	A728
			C1378	A1318	G1258	G1198	U1063	A997	U997	A936	A865	A729	A729
			G1379	C1319	C1259	C1199	C1064	G998A	G998A	A937	C866	G730	G730
			U1380	C1320	C1260	U1199	U1065	C998B	A938	C937	G867	G731	G731
			U1381	C1321	A1261	C1200	U1065	U999	G939	G939	G867	C732	C732
			C1382	G1322	C1262	A1201	C1066	U999	C940	C940	U870	C736	C736
			C1383	G1323	C1263	G1202	C1069	A1000	C941	C941	U871	A737	A737
			C1384	A1324	C1264	C1203	U1070	G1001	G942	G942	A872	G800	G800
			C1388	C1325	G1265	A1204	U1071	G1002	G1002	A873	U801	C738	C738
			U1389	G1326	G1266	U1205	C1071	G1003	G1003	A874	A802	C739	C739
			C1390	C1327	C1267	G1206	U1072	A1004	G944	G944	C874	G803	G803
			U1391	C1328	A1268	G1207	U1073	A1005	G945	G945	C875	U740	U740
			U1391	A1269	A1269	C1207	G1074	A946	A946	G947	G876	G804	G804
			C1463	G1330	C1270	C1271	C1075	G947	C948	C948	C877	C805	C805
			G1464	U1331	U1271	U1211	G1076	G1010	G1010	C878	C878	A806	A806
			C1465	U1332	U1272	U1212	G1077	G1011	G1011	A949	C879	U743	U743
			G1467	A1332	G1272	U1212	G1077	G1011	G1011	A949	C879	U744	U744

- Molecule 1: 16S rRNA



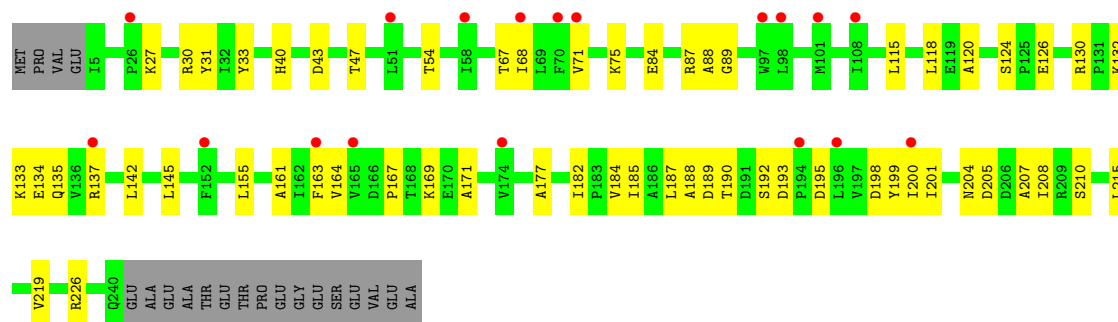


- Molecule 2: 30S ribosomal protein S2



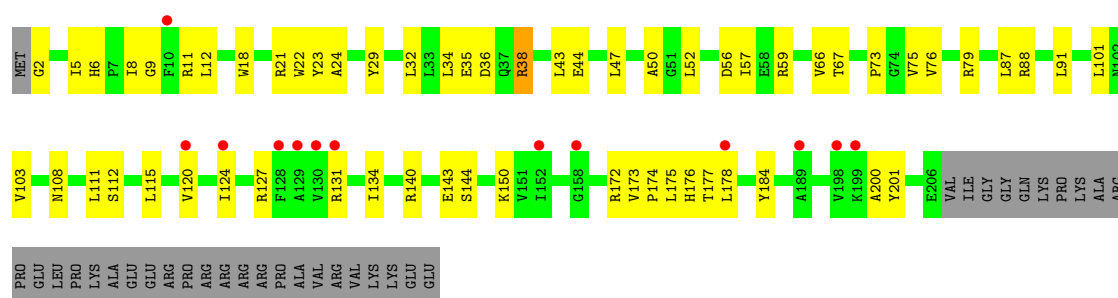
- Molecule 2: 30S ribosomal protein S2

Chain XB: 



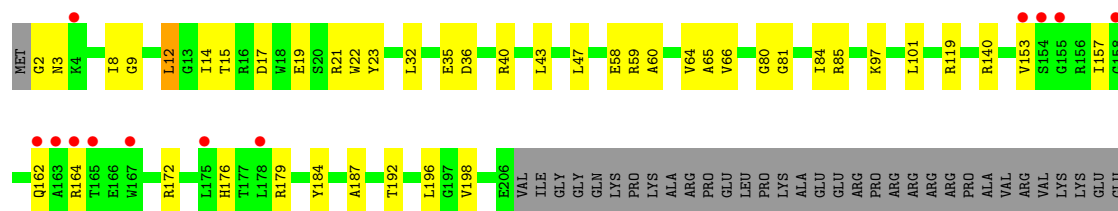
- Molecule 3: 30S ribosomal protein S3

Chain QC: 




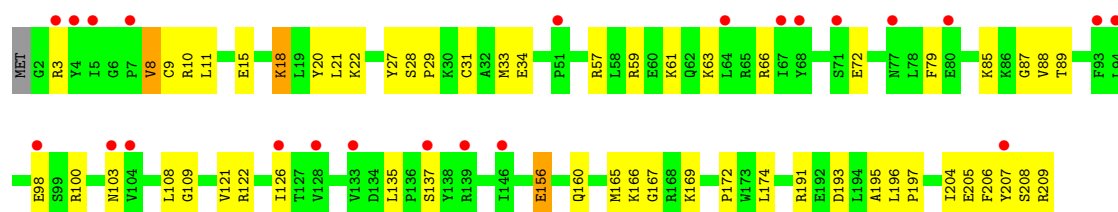
- Molecule 3: 30S ribosomal protein S3

Chain XC: 

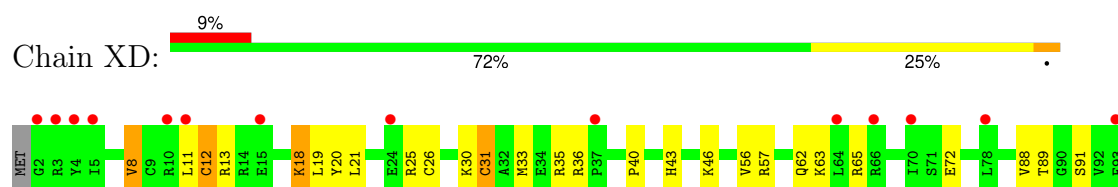


- Molecule 4: 30S ribosomal protein S4

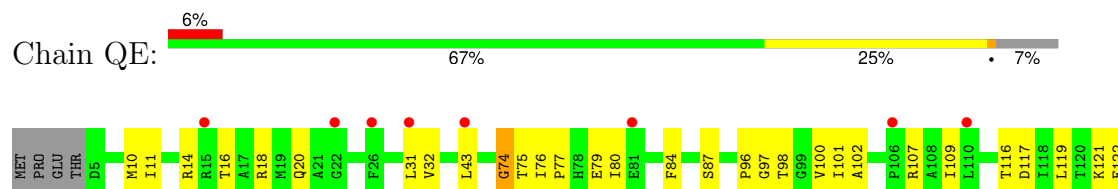
Chain QD: 



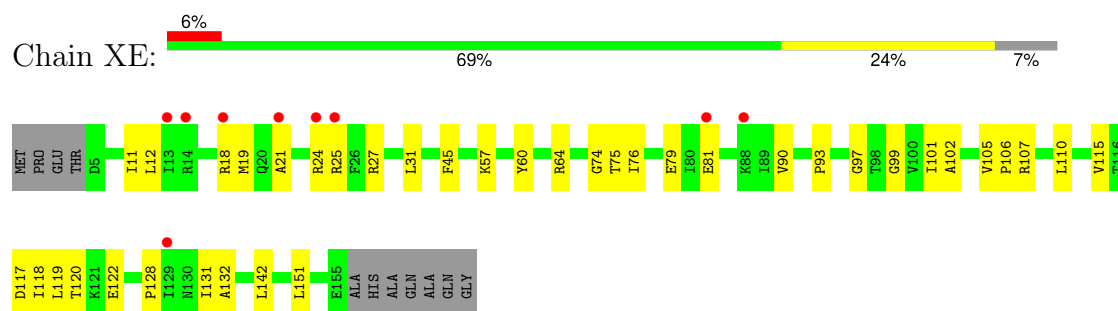
- Molecule 4: 30S ribosomal protein S4



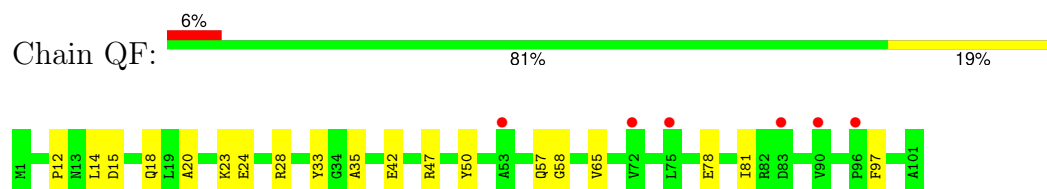
- Molecule 5: 30S ribosomal protein S5



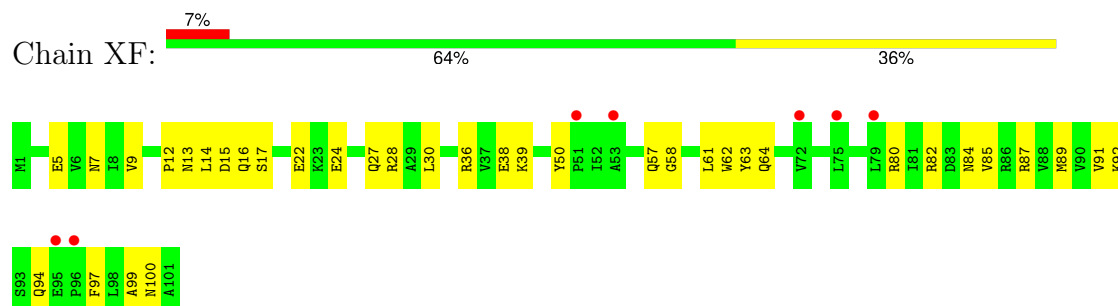
- Molecule 5: 30S ribosomal protein S5



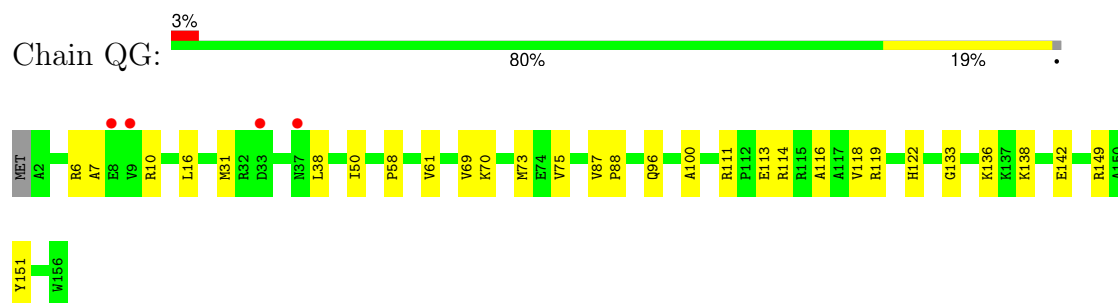
- Molecule 6: 30S ribosomal protein S6



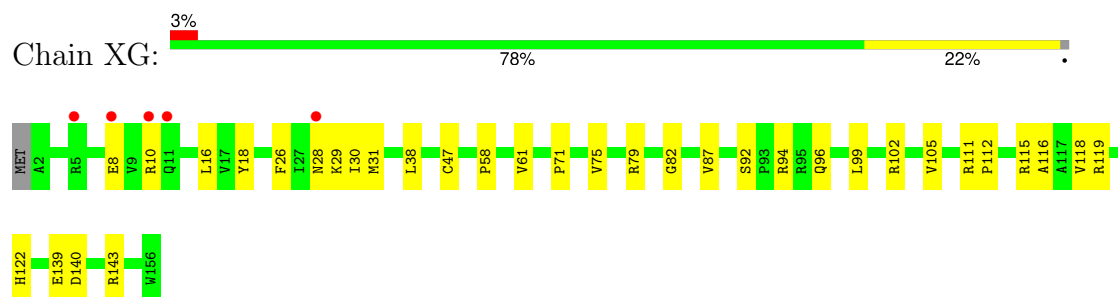
- Molecule 6: 30S ribosomal protein S6



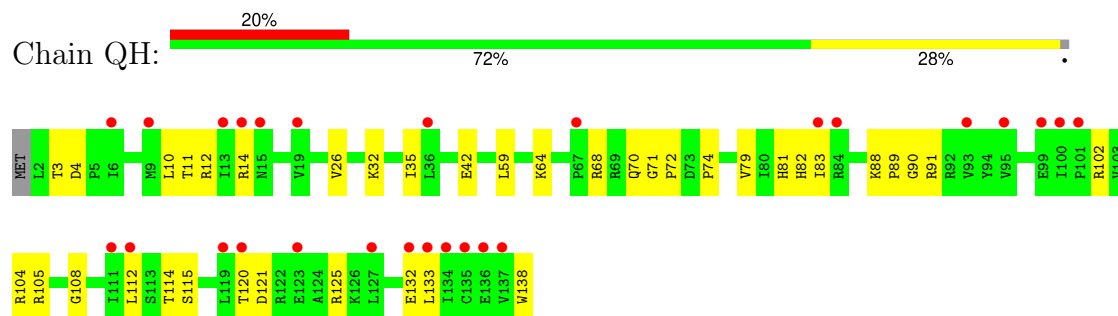
- Molecule 7: 30S ribosomal protein S7



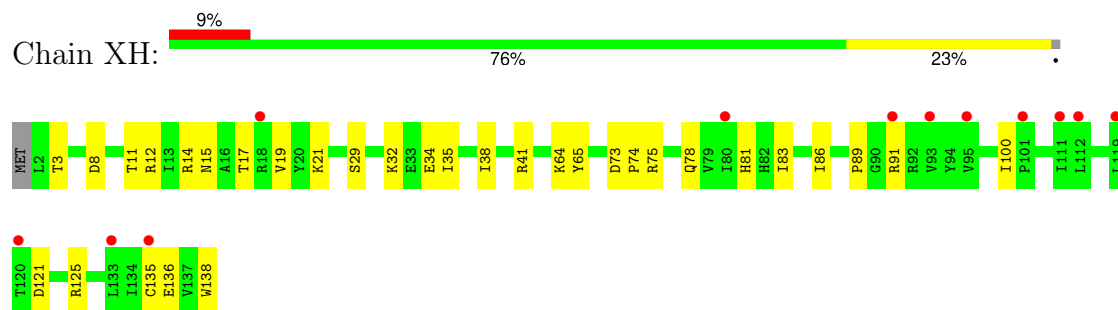
- Molecule 7: 30S ribosomal protein S7



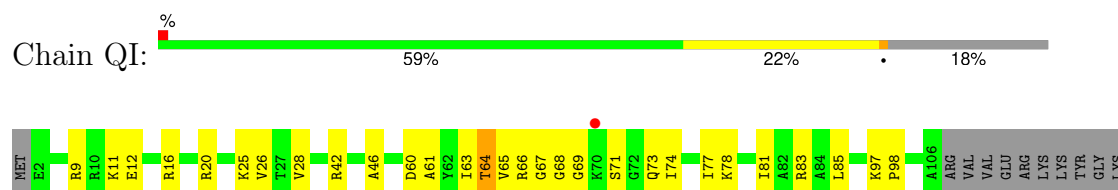
- Molecule 8: 30S ribosomal protein S8



- Molecule 8: 30S ribosomal protein S8



- Molecule 9: 30S ribosomal protein S9



HIS
LYS
ALA
ARG
ARG
ALA
PRO
GLN
TYR
SER
LYS
ARG

• Molecule 9: 30S ribosomal protein S9



MET E2 T7 R10 K11 E12 A13 V17 F18 L19 R20 P21 G22 T27 V28 V44 A45 H58 I63 G68 G69 K70 S71 G72 Q73 I74 I77 D105 A106 R107 V108 VAL GLU ARG LYS LYS TYR GLY LYS HIS LYS ALA ARG ARG ALA PRO GLN TYR SER LYS

ARG

• Molecule 10: 30S ribosomal protein S10



MET PRO K3 I4 R5 I6 R9 G10 F11 D12 H13 K14 T15 L16 D17 V24 R28 Q33 V34 S35 G36 P37 I38 P39 L40 P41 T42 R43 R44 R45 R46 F47 T48 V49 I50 R51 G52 P53 P54 F55 H56 K57 D58 S59 H62 L65 R66 T67 N68 N69 R70 V72 D73

N76 R79 G83 V94 E95 I98 K99 T100 V101 GLY GLY GLY ARG

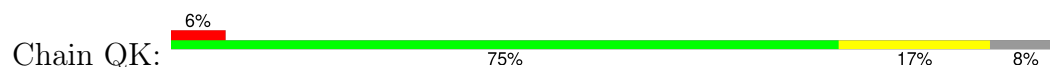
• Molecule 10: 30S ribosomal protein S10



MET PRO LYS ILE R5 I6 K7 L8 R9 G10 H13 K14 T15 L16 V24 E25 A26 G36 P37 I38 P39 L40 P41 T42 R43 V44 R45 R46 F47 T48 V49 I50 P53 F54 K55 D58 S59 R60 E61 H62 E63 E64 L65 R66 T67 H68 N69 R70 L71 V72 N78 R79 R80 T81

Q84 L90 P91 T92 G93 V94 E95 I96 E97 I98 K99 T100 VAL GLY GLY GLY ARG

• Molecule 11: 30S ribosomal protein S11



MET ALA LYS LYS PRO SER LYS LYS VAL K11 R12 Q13 G17 N27 T28 T32 P35 P39 T40 T41 G52 S53 R54 K55 G56 T57 A60 K70 K71 V82 I83 V84 S101 I108 V109 D110 P115 H116 N117 G118 C119 G119 F125 R126 K127 A128 S129

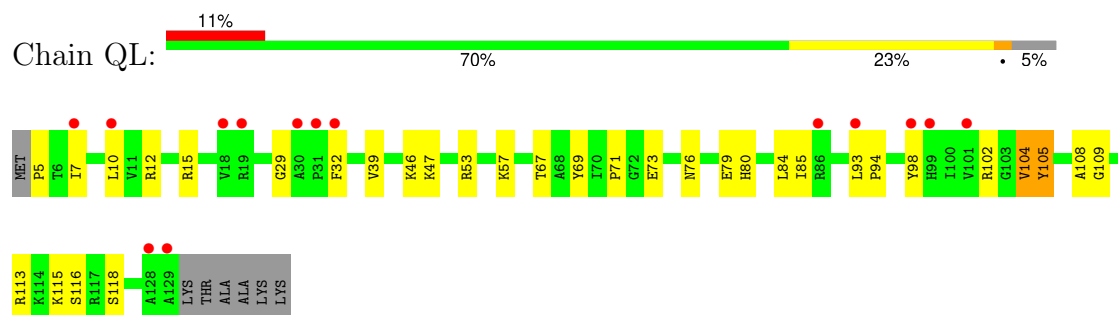
• Molecule 11: 30S ribosomal protein S11



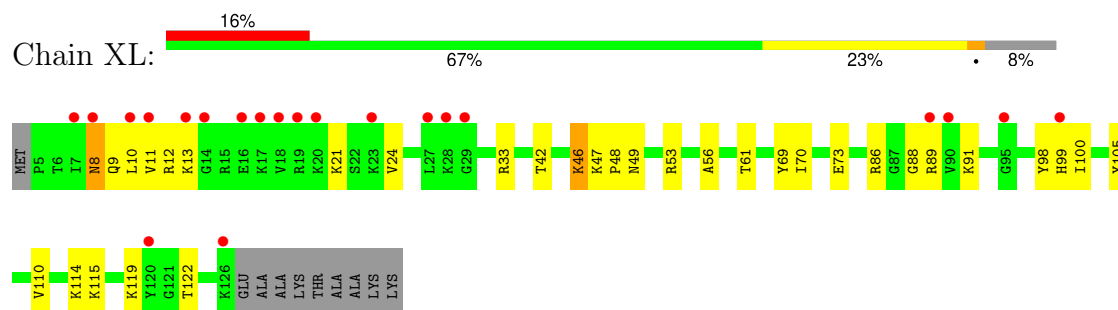
MET ALA LYS LYS PRO SER LYS LYS VAL K11 A15 S16 G17 T31 T35 W42 K51 R54 T57 M77 V82 I83 V84 R85 R91 Q104 V105 K106 S107 I108 V109 D110 P113 V114 P115 H116 N117 G118 C119 R120 P121 K122 K123 K124 F125 R126 LYS

ALA
SER

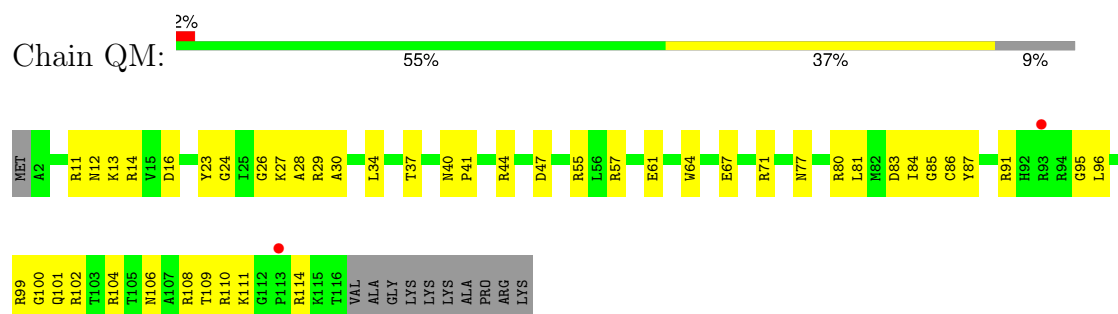
- Molecule 12: 30S ribosomal protein S12



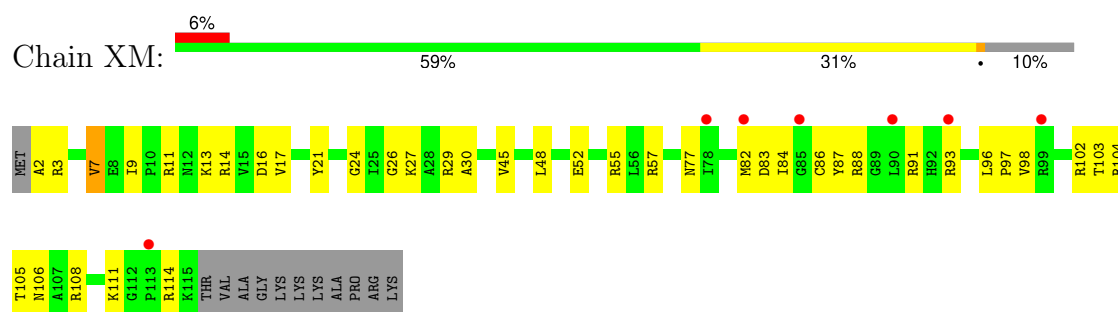
- Molecule 12: 30S ribosomal protein S12



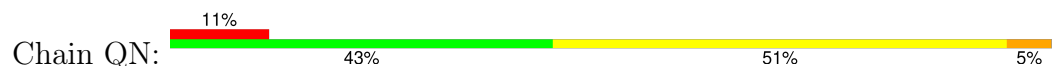
- Molecule 13: 30S ribosomal protein S13



- Molecule 13: 30S ribosomal protein S13



- Molecule 14: 30S ribosomal protein S14 type Z

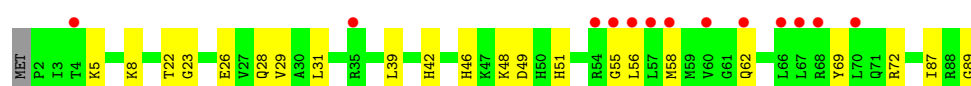
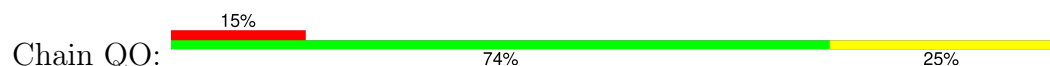




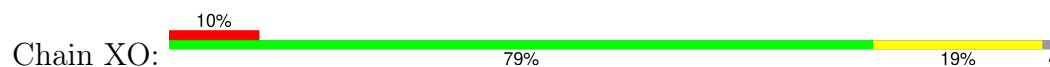
- Molecule 14: 30S ribosomal protein S14 type Z



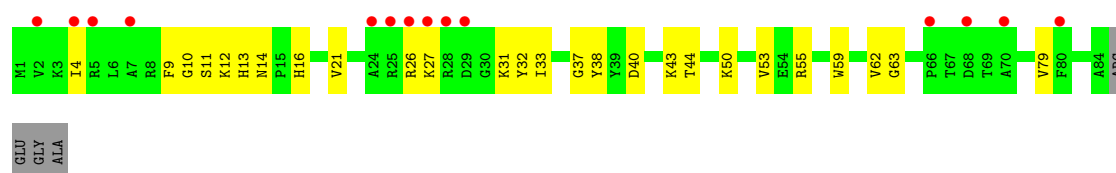
- Molecule 15: 30S ribosomal protein S15



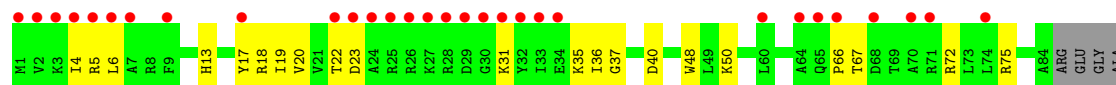
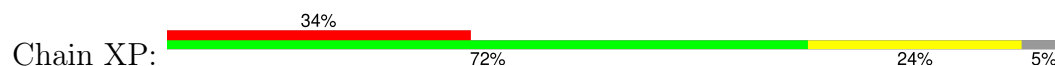
- Molecule 15: 30S ribosomal protein S15



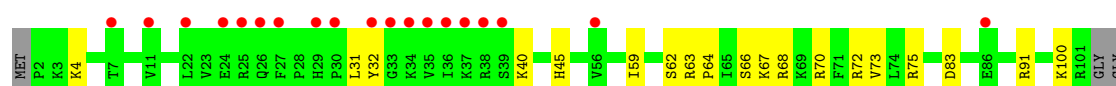
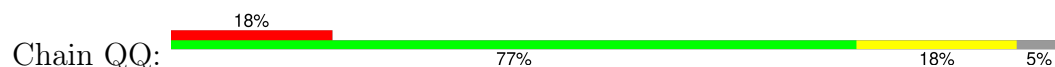
- Molecule 16: 30S ribosomal protein S16



- Molecule 16: 30S ribosomal protein S16




- Molecule 17: 30S ribosomal protein S17



LYS
ALA

- Molecule 17: 30S ribosomal protein S17

Chain XQ: 



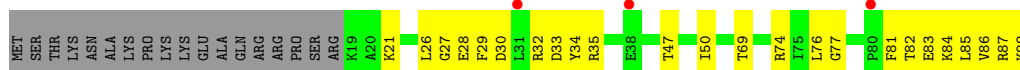
- Molecule 18: 30S ribosomal protein S18

Chain QR: 



- Molecule 18: 30S ribosomal protein S18

Chain XR: 



- Molecule 19: 30S ribosomal protein S19

Chain QS: 



LYS
LYS
LYS

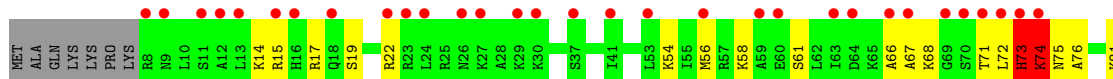
- Molecule 19: 30S ribosomal protein S19

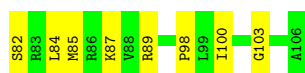
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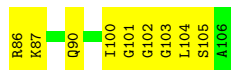
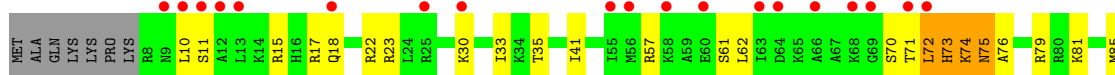
- Molecule 20: 30S ribosomal protein S20

Chain QT: 

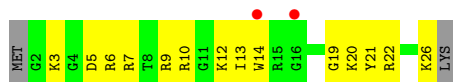
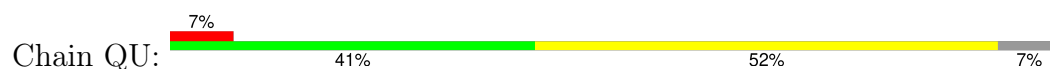




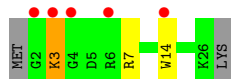
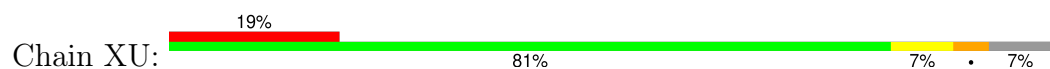
- Molecule 20: 30S ribosomal protein S20



- Molecule 21: 30S ribosomal protein Thx



- Molecule 21: 30S ribosomal protein Thx



- Molecule 22: tRNA(Pro)



- Molecule 22: tRNA(Pro)



- Molecule 23: mRNA




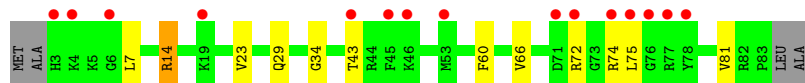
- Molecule 23: mRNA

Chain XX: 




- Molecule 24: 50S ribosomal protein L27

Chain R0: 




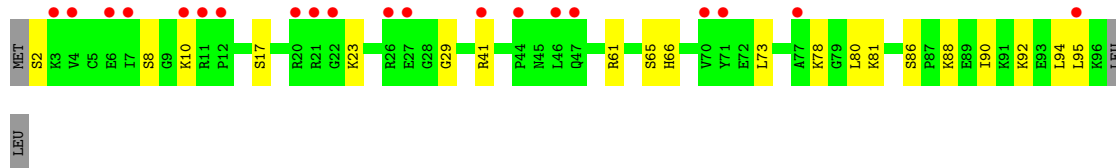
- Molecule 24: 50S ribosomal protein L27

Chain Y0: 




- Molecule 25: 50S ribosomal protein L28

Chain R1: 




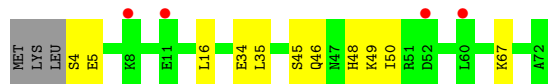
- Molecule 25: 50S ribosomal protein L28

Chain Y1: 




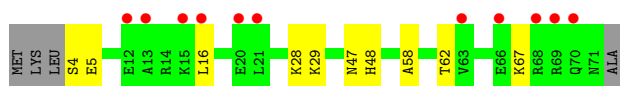
- Molecule 26: 50S ribosomal protein L29

Chain R2: 

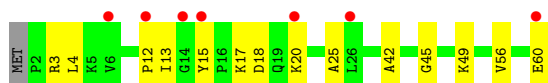
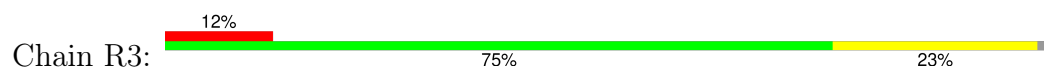


- Molecule 26: 50S ribosomal protein L29

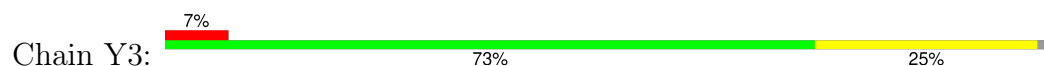
Chain Y2: 



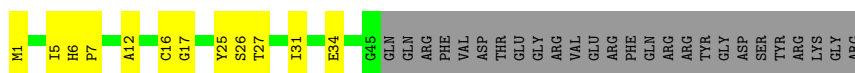
- Molecule 27: 50S ribosomal protein L30



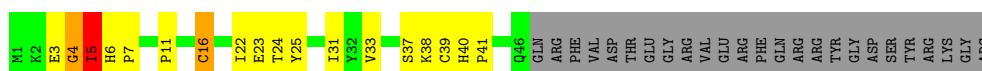
- Molecule 27: 50S ribosomal protein L30



- Molecule 28: 50S ribosomal protein L31



- Molecule 28: 50S ribosomal protein L31



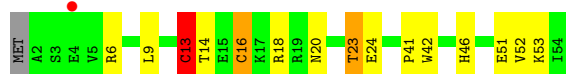
- Molecule 29: 50S ribosomal protein L32



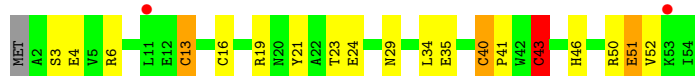
- Molecule 29: 50S ribosomal protein L32



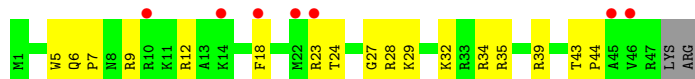
- Molecule 30: 50S ribosomal protein L33



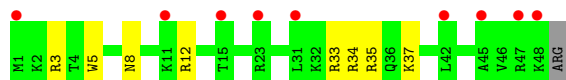
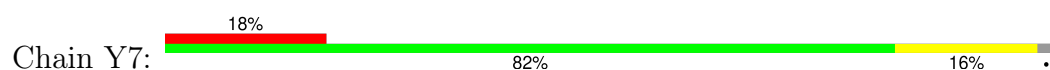
- Molecule 30: 50S ribosomal protein L33



- Molecule 31: 50S ribosomal protein L34



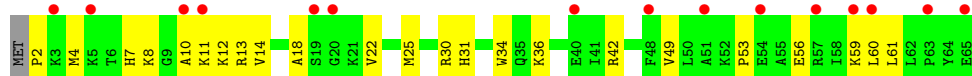
- Molecule 31: 50S ribosomal protein L34



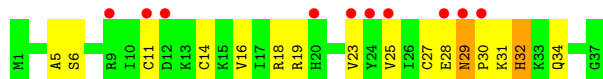
- Molecule 32: 50S ribosomal protein L35



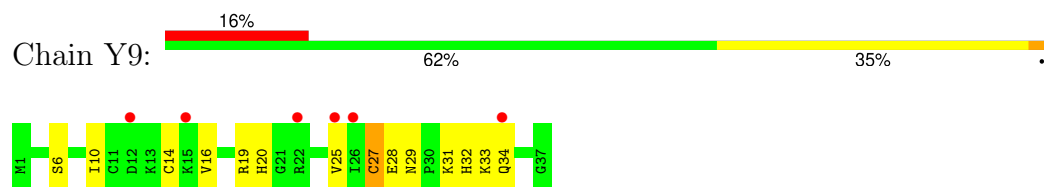
- Molecule 32: 50S ribosomal protein L35



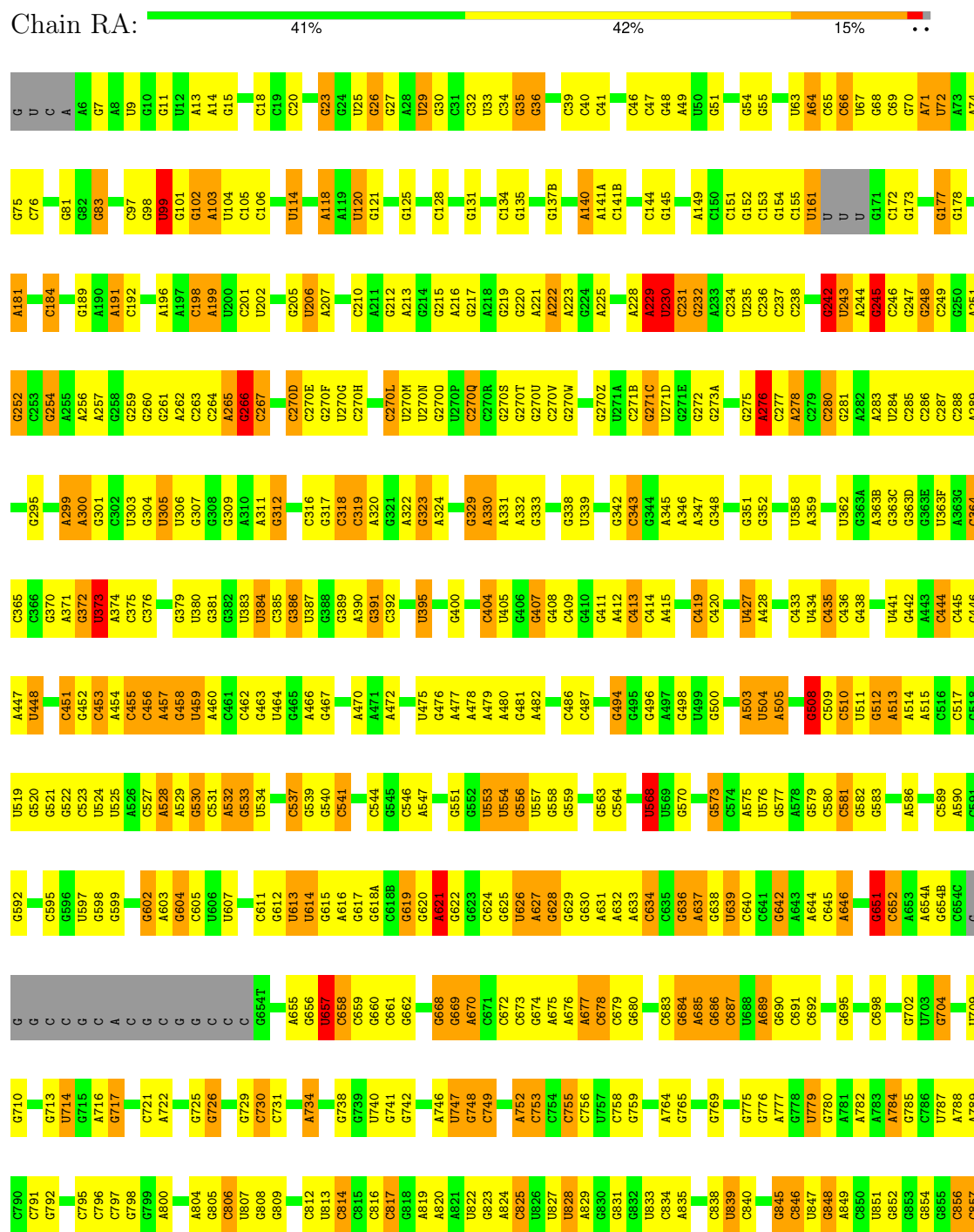
- Molecule 33: 50S ribosomal protein L36



• Molecule 33: 50S ribosomal protein L36



• Molecule 34: 23S rRNA



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U1955	C1961	A1871	U1693	A1614	C1544	G1473	G1400	C1314	U1240	A1156	A1086	G1015	U943	G859
U1956	C1962	A1872	C1694	C1615	A1545A	C1474	G1401	C1315	A1241	A1157	A1087	G1016	G944	U860
G1959	C1963	A1878	G1695	A1616	C1545B	G1475	C1402	U1316	G1244	C1161	A1088	U1019	A945	A863
A1960	C1964	C1879	C1617	C1617	C1546	A1476	C1403	A1317	G1244	G1162	G1089	A1020	G946	G864
A1961	C1965	A1880	A1618	A1618	C1547	A1477	C1404	C1318	G1247	U1165	U1090	A1021	C949	C865
C1961	C1966	G1792	U1621	U1621	C1548	G1480	U1405	G1319	A1247	C1166	A1096	G1022	G950	A866
C1962	C1967	C1793	U1629	U1629	C1549	U1482	U1406	U1326	U1249	U1167	U1097	G1023	G951	C867
U1963	C1968	C1794	G1630A	G1630A	C1557	G1483	C1408	C1327	U1250	G1168	A1098	G1024	U944	U868
G1964	C1969	U1796	A1885A	A1885A	C1558	G1484	C1409	G1328	C1251	G1169	G1099	G1025	A953	G869
C1965	C1970	C1797	G1708	G1708	C1559	G1485	G1410	U1329	G1252	C1170	C1100	U1026	C954	G869
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C1967	C1972	G1799	C1710	C1710	G1561	A1491	C1412	A1331	U1254	G1172	C1102	A1028	G956	U877
A1968	C1973	C1800	C1638	C1638	C1562	U1496	C1413	G1332	A1255	A1173	A1103	A1029	A957	U877
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A1971	C1976	A1803	A1641	A1641	C1565	A1499	C1416	C1335	C1258	A1176	U1106	A1032	A960	G883
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G1973	C1978	U1808	G1643	G1643	C1567	C1498	G1421	G1337	G1260	C1178	U1108	G1034	C965	A887
G1989	C1979	A1809	G1644	G1644	A1567	C1501	A1427	G1338	G1265	C1179	C1109	C1041	G966	C888
A1900	C1980	A1810	U1730	U1730	C1568	C1502	C1428	G1339	U1267	C1180	G1110	G1042	C967	C889
C1903	C1981	G1811	C1646	C1646	A1569	C1503	G1429	U1340	G1267	C1181	A1111	C1043	G968	A890
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G1906	C1984	A1815	C1734	C1734	G1572	C1506	U1434	C1341	U1272	A1188	A1047	A1036	C971	U895
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C1908	C1986	G1817	C1736	C1736	C1574	A1508	G1435	U1352	A1274	G1190	C1116	A1048	A973	C903
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G1910	C1988	C1819	C1657	C1657	C1577	C1509	C1437	A1354	G1278	G1195	C1121	A1050	C974B	C904
A1913	C1989	U1820	G1658	G1658	U1578	A1510	G1441	C1362	G1280	C1200	G1122	G1052	G975	U905
C1914	C1990	A1822	U1659	U1659	A1579	A1511	G1442	C1363	U1281	C1201	C1123	C1053	A980	U907
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C1929	C1995	A1829	C1665	C1665	C1587	C1518	G1447	G1368	U1287	C1207	G1130	G1059	G987	C915
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C1943	C2004	A1847	C1673	C1673	C1594	G1527	U1454	A1378	G1296	G1216	U1141	A1067	C994	C923
A2013	C2005	C1846	G1674	G1674	G1595	A1528	G1455	C1379	C1297	C1217	U1142A	G1068	C995	C924
A2014	C2006	A1847	C1675	C1675	C1596	A1529	A1460	G1380	C1298	G1218	A1143B	A1069	A996	C925
C2019	C2007	C1848	A1676	A1676	C1598	G1529	G1461	G1381	G1299	G1219	A1143	A1070	G997	A926
A2020	C2008	G1849	U1676	U1676	A1603	C1533	G1462	C1382	U1300	A1220	C1144	G1071	C998	G928
C2021	C2009	C1850	C1773	C1773	C1604	C1534	C1462	G1383	A1301	C1221	C1145	C1072	U999	G929
G2022	C2010	A1936	C1774	C1774	C1605	G1535	C1463	A1384	U1302	C1222	C1146	A1073	U999	G930
C2023	C2011	U1775	U1775	U1775	C1606	U1535	C1464	A1385	G1303	C1222	C1147	G1074	A1000	U930
C2024	C2012	C1853	G1776	G1776	G1606	A1536	G1465	G1385	C1304	G1225	A1148	A1074	C1075	G931
C2025	C2013	A1854	U1777	U1777	C1607	C1537	G1466	U1391	C1304	G1226	C1149	C1075	C1005	G932
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A2028	C2015	G1857	G1777	G1777	A1608	C1538	C1467	U1391	A1308	G1226	C1149	C1076	C1008	G934
C2029	C2016	C1858	C1778	C1778	A1609	G1539	C1468	U1394	G1309	G1235	G1151	A1077	C1008	A934
A2030	C2017	A1689	U1688	U1688	C1609	G1540	C1469	U1394	G1310	G1236	C1152	U1082	G1011	C936
C2031	C2018	C1782	C1782	C1782	A1610	G1540	A1469	U1394	G1310	G1236	C1152	U1082	G1011	C936
G2032	C2019	A1783	A1690	A1690	C1611	U1541	G1470	U1398	C1311	G1237	C1153	U1083	U1012	A941
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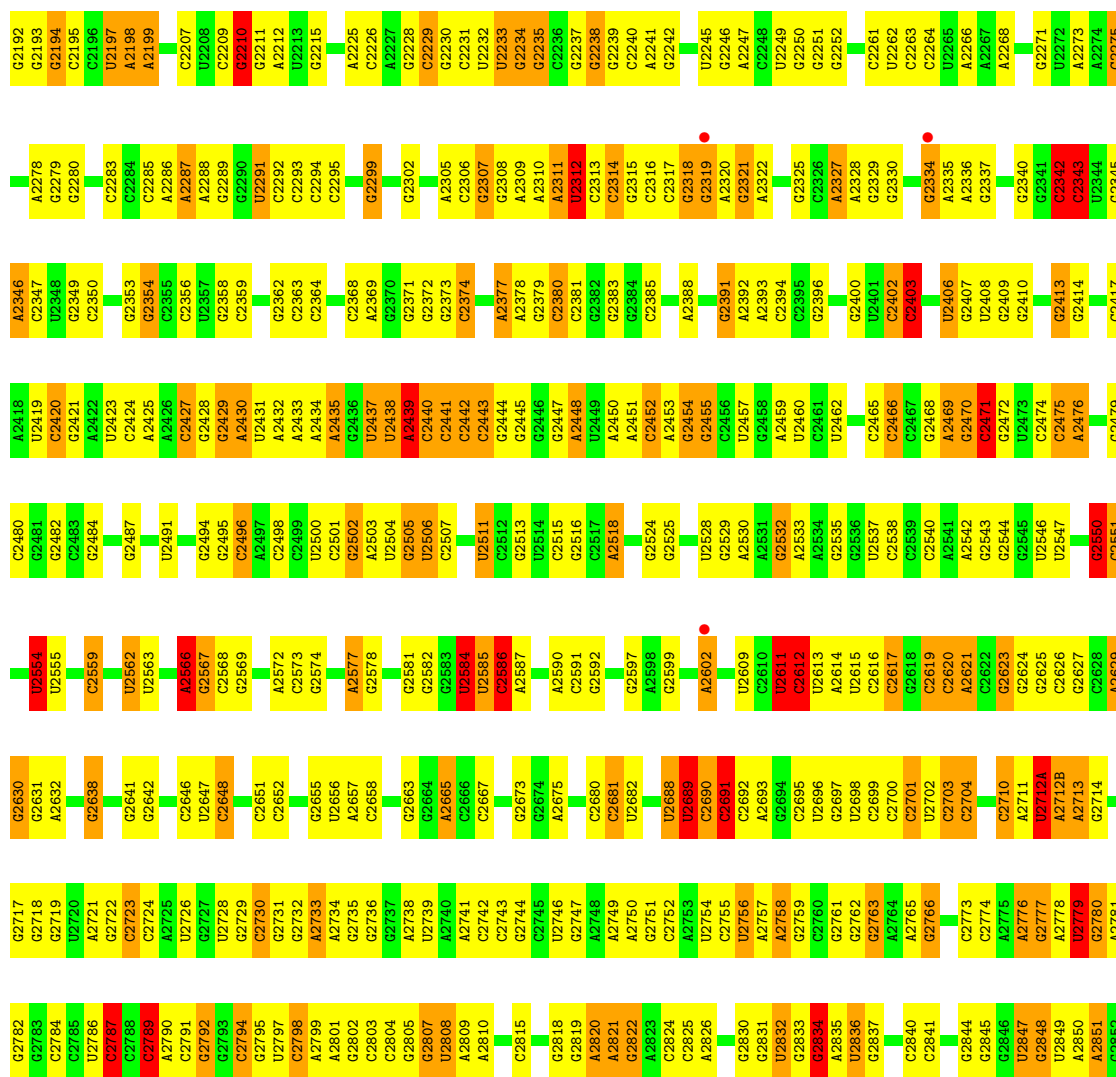


Response	Percentage
Yes, the U.S. should take action to protect the environment	38%
No, the U.S. should focus on the economy	45%
It's not the U.S.'s responsibility	15%
Don't know	2%



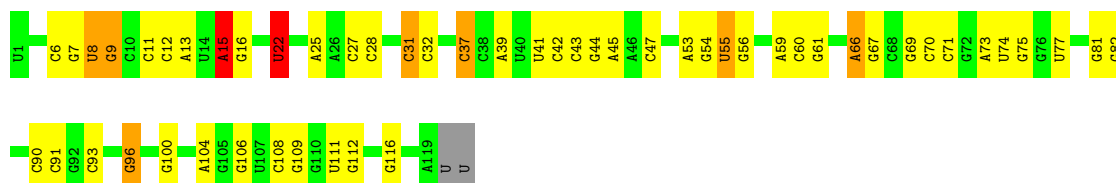
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A1057	C992	G842	G775	C687	A647	G583	C509	G438	A363G	C292	G261	A190	G96
G1058	G993	G843	G776	U688	G648	C584	C510	G439	C364	C293	A262	A191	G97
G1059	G994	G844	G777	A689	G649	G585	U511	G442	C365	C296	C263	C192	G98
U1061	C995	G845	G778	G690	G649	G586	G512	A443	C366	C297	C264	C98	
G1062	C996	C846	U779	C691	C650	A586	A513	G444	G370	G298	A265	A195	U99
	A996	U847	G780	C692	G651	C589	A514	C445	A371	A299	A266	A196	G101
U1065	G997	G848	A781	G695	C652	A590	A515	G446	G372	A197	C267	A197	G102
U1066	C998	A849	A782	G696	A653	C591	G592	A447	U373	A103	C268	A198	A104
G1067	A1000	U851	A783	A699	A654A	G593	G594	U448	A374	C375	A270A	A199	C105
A1068	G1005	G852	G786	G701	C654C	C595	G521	C452	C376	U306	A270B	C106	C107
A1070	C1008	G854	U787	G704	G	G596	G522	C453	U380	G307	C270D	A204	U109
G1071	A1009	G855	A788	G705	G	U597	C523	A454	G381	G311	C270E	G205	U114
C1072	A1010	C856	A789	U709	C	G598	U524	C455	G382	G312	G270F	A206	
G1073	G1011	C857	G790	G710	C	G599	U525	C456	U383	C208	C270G	C208	A118
G1074	U1012	G858	G792	G713	C	C601	A527	C458	U384	C209	C270H	C210	U120
C1075	G1013	U860	A793	G714	A	G602	A528	U459	C385	G316	C270I	C211	G125
A1077	U1014	A861	G794	U714	C	A603	A529	A460	G386	G317	G270J	A216	
U1078	G1015	G862	C795	G715	C	G604	G530	C461	U387	C318	G270K	G215	
G1079	G1016	A863	C796	C721	C	C531	C462	G463	G388	C319	C270L	G216	
C1080	G1017	G864	C797	A722	C	A532	A532	G464	G389	A320	U270M	G217	
U1081	G1018	C865	G798	A723	G	G607	A533	U464	A390	G321	U270N	G217	C128
U1082	U1019	A866	G799	G726	C	A608	G534	G465	G391	G322	G270O	G220	C129
U1083	A1020	G869	A800	A727	C	A609A	U534	A466	C392	G323	C270P	A221	G130
A1084	G1021	A870	A804	C730	C	C910	A536	A467	G395	G329	C270Q	A222	G131
A1085	U1022	U871	G805	C731	G654T	G612	C537	G468	G398	A331	G270R	A223	C134
A1086	G1023	A872	C806	C732	C654U	U613	G539	G469	G400	A332	G270T	G135	
G1087	G1024		U807	G738	A654V	U614	C540	A470	G399		G270U	G226	G137B
A1088	G1025		G808	G739	A655	G615	C541	A471	G400	C335	C270V	A227	G138
U1089	U1026		U811	U740	G656	A616	C544	A472	G400	C336	G270W	A228	G139
U1090	A1027		U812	G741	U657	G617	G545	G473	C404	G337	G270X	U230	A140
	A953		U813	G742	C658	G619	G546	U475	U405	C338	U271A	C231	A141A
G1093	G956		U814	G745	C659	G620	A547		G406	G232	C271B	G232	G141B
U1094	A957		C815	A746	G661	A621	U553	A479	G407	A233	G271C	G233	G142
A1095	U958		C816	U747	G662	G622	U554	A480	G408	C234	U271D	C234	C143
U1096	A959		G663	G748	G663	G623	G556	G481	C409	G343	G271E	U235	C144
A1098	G960		C664	C749	C665	G624	U557	A482	G410	C236		C236	G145
G1099	C961		G818		C665	G625	G558	A483	G411	C237	C273E	C237	
U1101	G962		A819	A752	C666	U626	G559	C484	A412	C238	U273F	C238	C151
C1098	U963		A820	C753	G667	G627	G564	C486	C413	A347	C273G		G152
G1102	C964		G823	C754	G669	G628	G563	C487	C414	G348	G274	A241	C155
A1103	C971		C824	C755	A670	G629	C564	C415	C416	G352	G275	G242	C156
C1104	A972		A824	G756	C671	G632	C565	C417	C277	G245	A276	A244	U161
U1105	A973		U827	U757	C672	A633	U568	G418	A278	C246	C277	G245	U
U1108	G974A		G758	C758	C673	A634	C564	C419	C279	G247	G279	G247	U
G1109	C974B		G759	G759	C674	C634	U569	G494	C280	G248		C172	G171
G1110	G975		A676	G764	A675	G635	G573	G495	A283	G249	A284	G173	G173
A1111	C976		A677	A764	A676	G636	C574	A497	G360	G250	U284	G250	G177
G1112	A900		G765	G765	A677	G637	U575	G498	G361	A251	C285	G252	G178
U1113	A901		C766	C766	C678	G638	U576		G362	G253	C286	G253	G178
C1049	A953		A834		C679	U639	G577	A428	G363A	C287	C287	G254	C184
G1114	A954		A835		C679	C540	A578	A429	A363B				
G1115	C985		C838	G771	G680	G579	A505	G430	G363C				
C1116	G1051		U839	C772	G684	C580		G506	G363D				
G1117	G987												





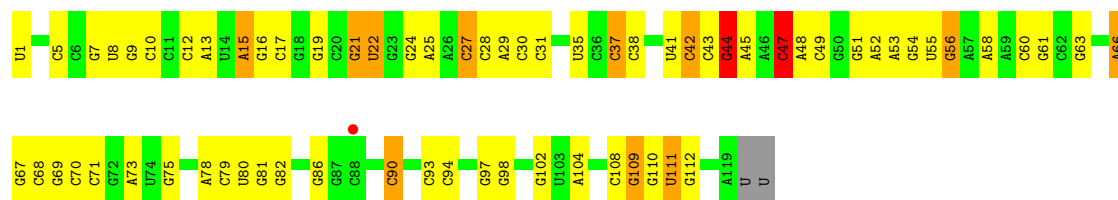
• Molecule 35: 5S rRNA

Chain RB: 55% 36% 6% ..

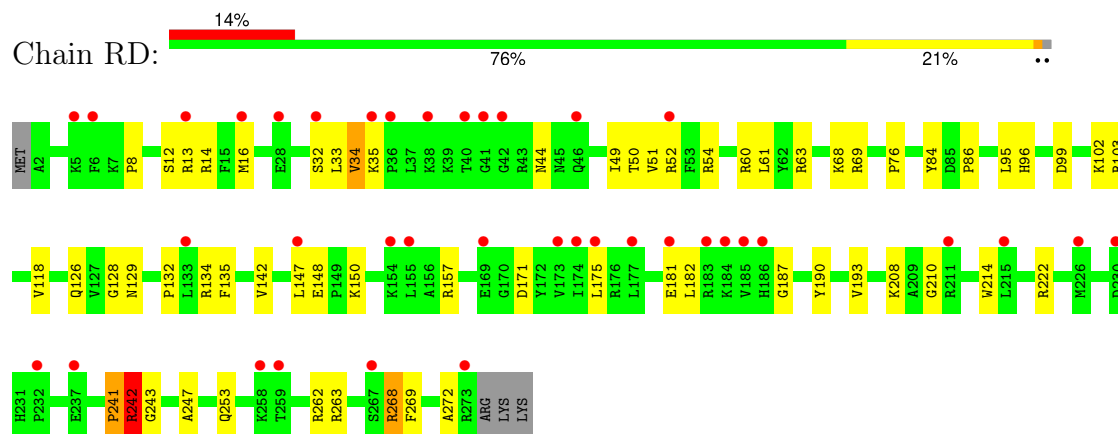


• Molecule 35: 5S rRNA

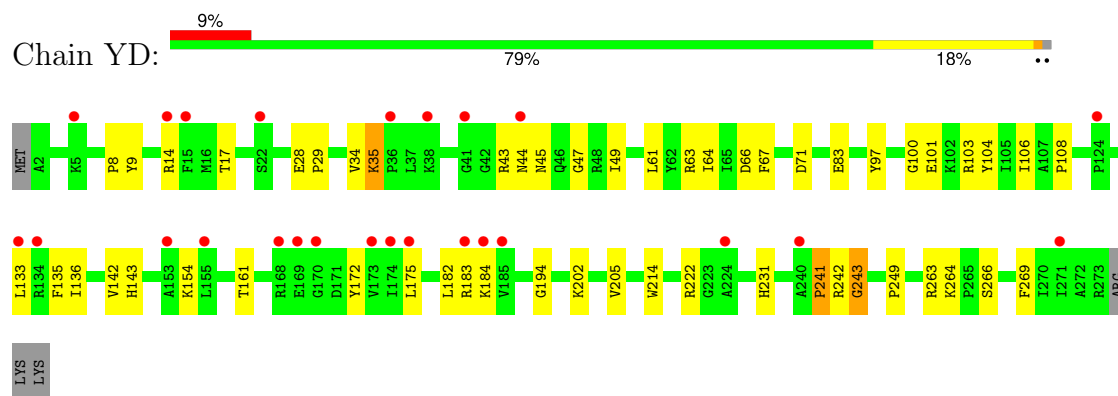
Chain YB: 43% 45% 9% ..



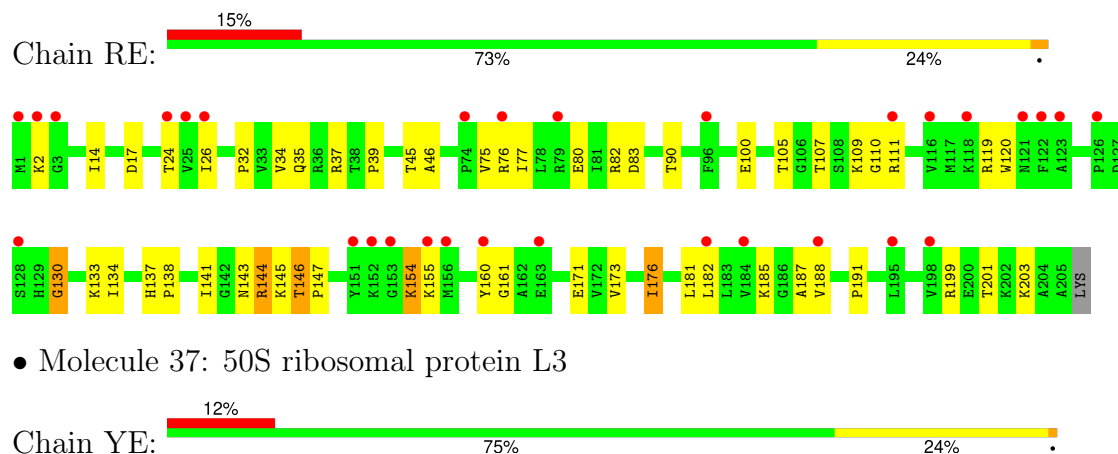
• Molecule 36: 50S ribosomal protein L2

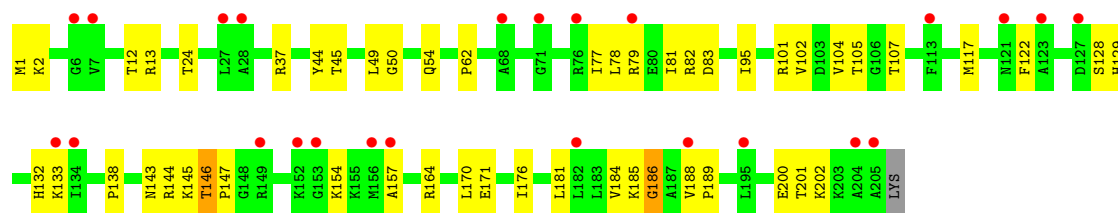


• Molecule 37: 50S ribosomal protein L3

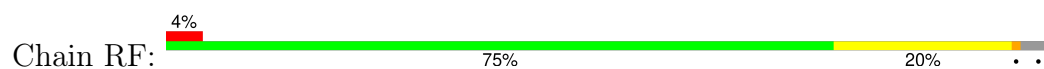


• Molecule 37: 50S ribosomal protein L3

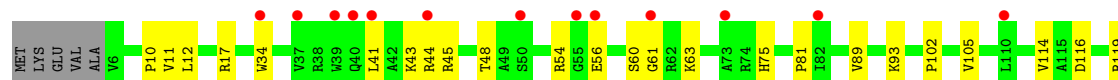
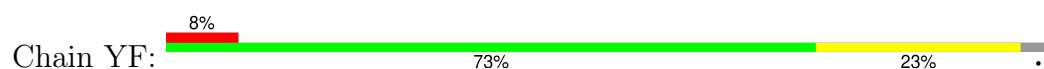




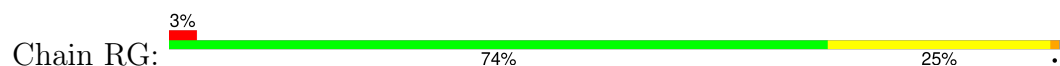
- Molecule 38: 50S ribosomal protein L4



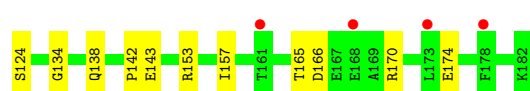
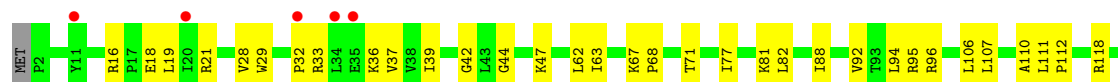
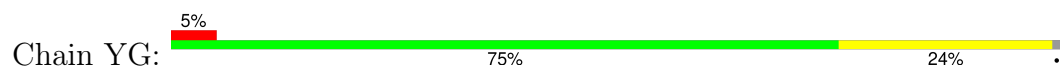
- Molecule 38: 50S ribosomal protein L4



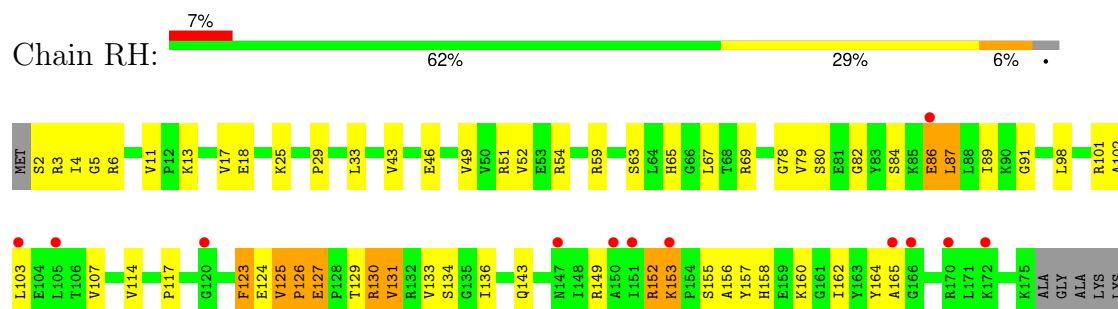
- Molecule 39: 50S ribosomal protein L5



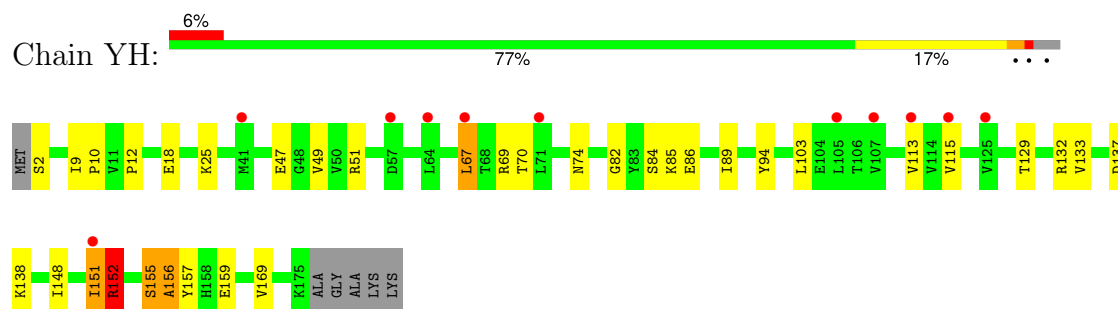
- Molecule 39: 50S ribosomal protein L5



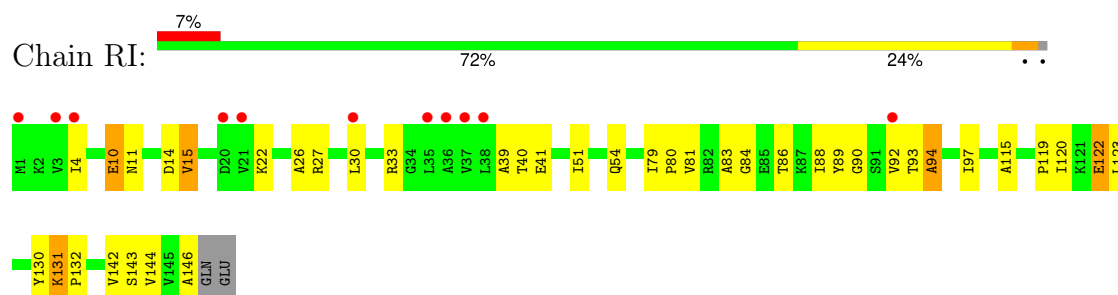
- Molecule 40: 50S ribosomal protein L6



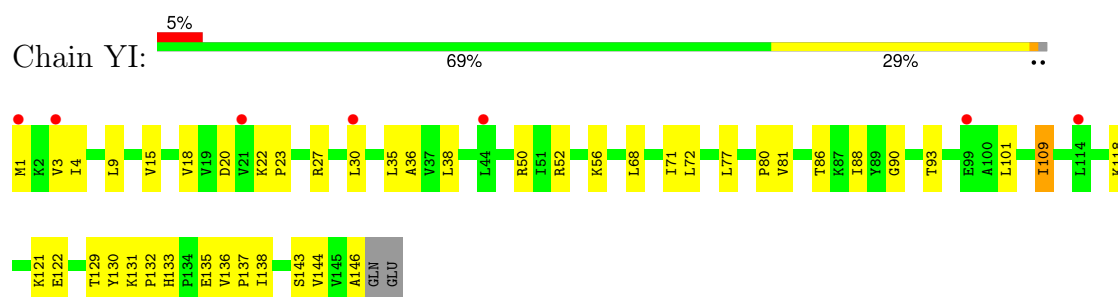
- Molecule 40: 50S ribosomal protein L6



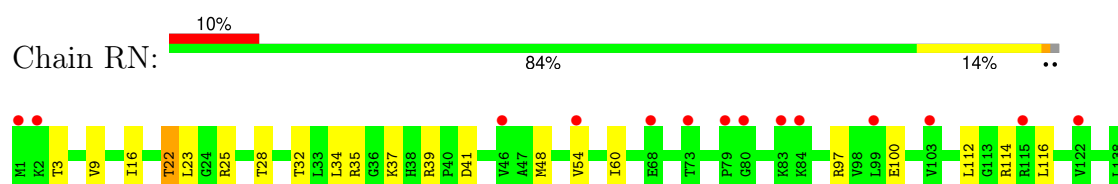
- Molecule 41: 50S ribosomal protein L9



- Molecule 41: 50S ribosomal protein L9




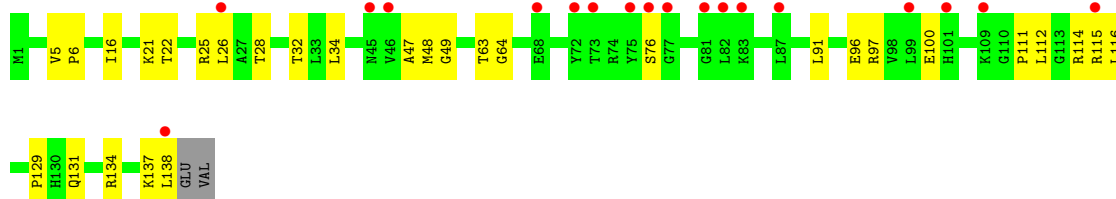
- Molecule 42: 50S ribosomal protein L13




GLU
VAL

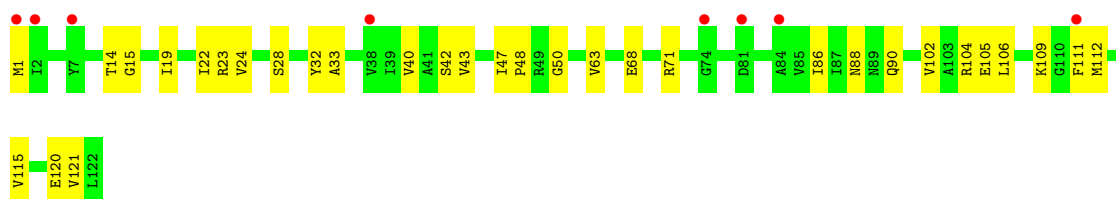
- Molecule 42: 50S ribosomal protein L13

Chain YN:  13% 77% 21%




- Molecule 43: 50S ribosomal protein L14

Chain RO:  7% 74% 26%




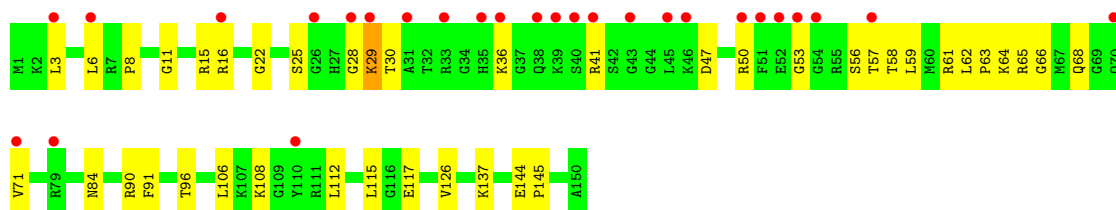
- Molecule 43: 50S ribosomal protein L14

Chain YO:  82% 18%




- Molecule 44: 50S ribosomal protein L15

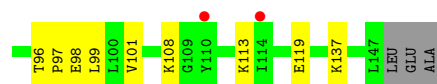
Chain RP:  18% 73% 27%



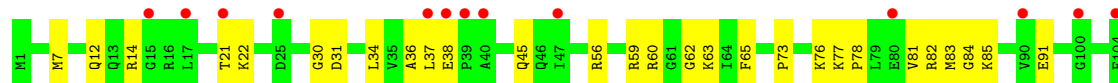
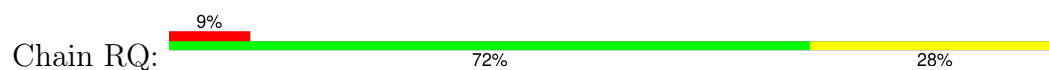
- Molecule 44: 50S ribosomal protein L15

Chain YP:  11% 71% 27%

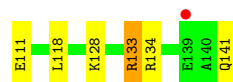
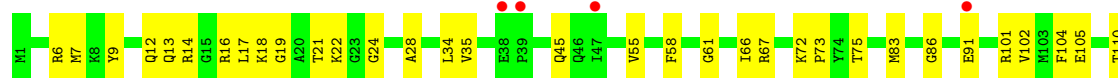
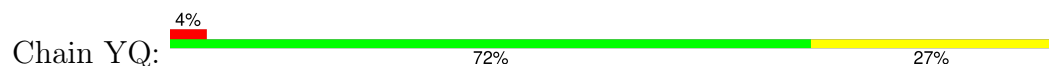




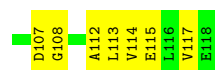
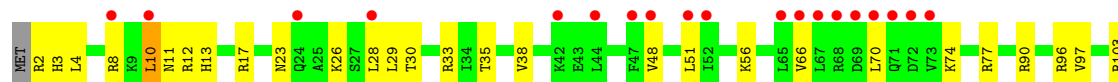
- Molecule 45: 50S ribosomal protein L16



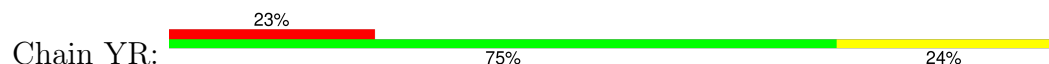
- Molecule 45: 50S ribosomal protein L16



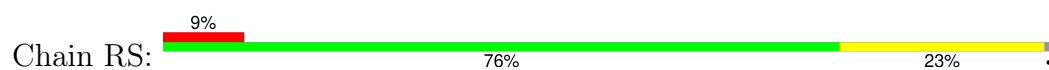
- Molecule 46: 50S ribosomal protein L17



- Molecule 46: 50S ribosomal protein L17

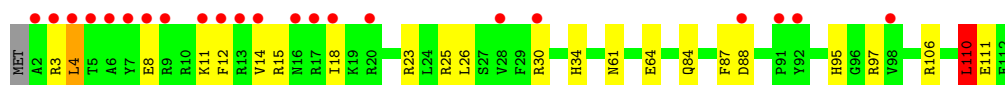
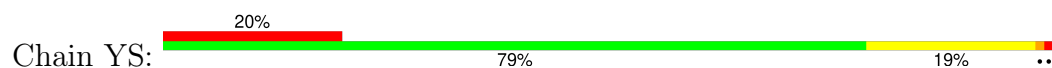


- Molecule 47: 50S ribosomal protein L18

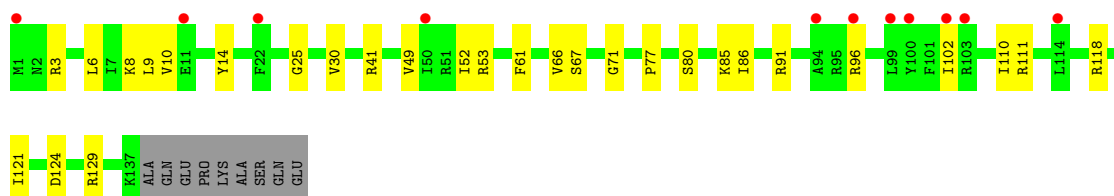
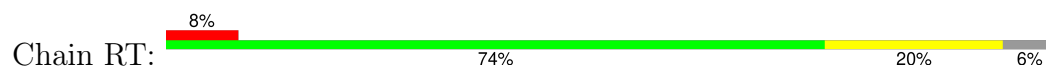




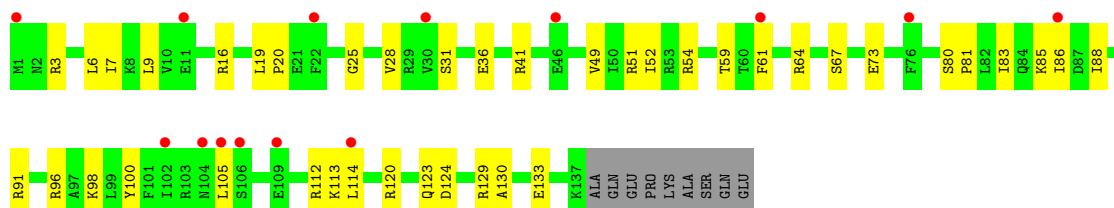
- Molecule 47: 50S ribosomal protein L18



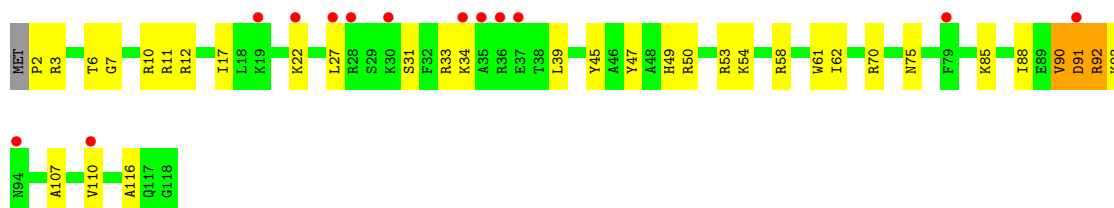
- Molecule 48: 50S ribosomal protein L19



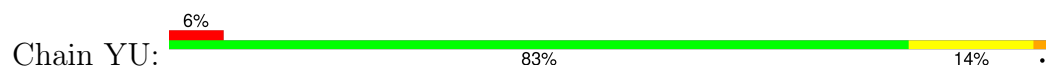
- Molecule 48: 50S ribosomal protein L19



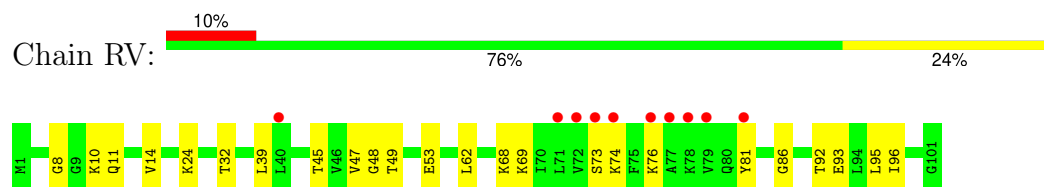
- Molecule 49: 50S ribosomal protein L20



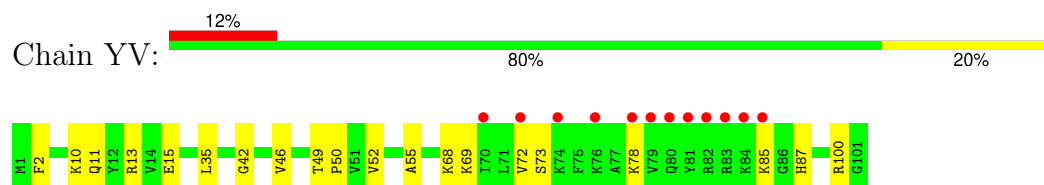
- Molecule 49: 50S ribosomal protein L20



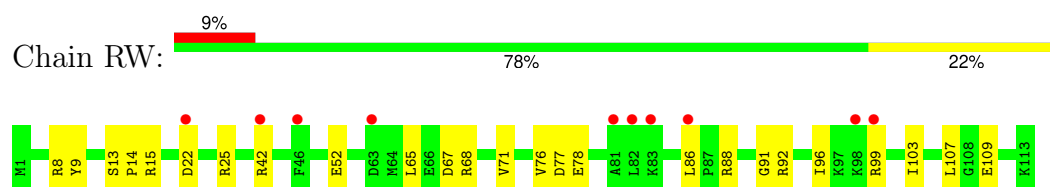
- Molecule 50: 50S ribosomal protein L21



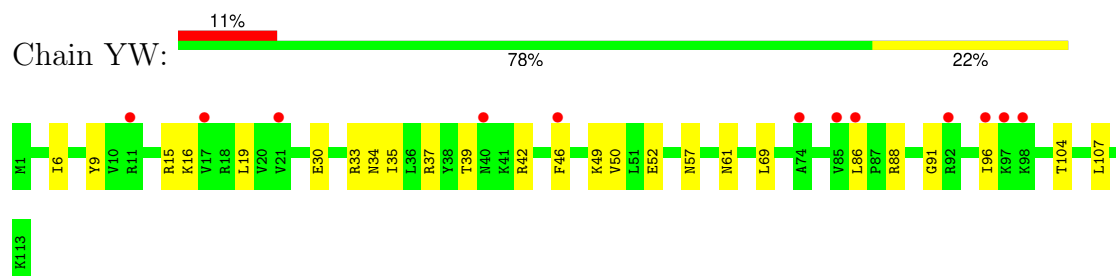
- Molecule 50: 50S ribosomal protein L21



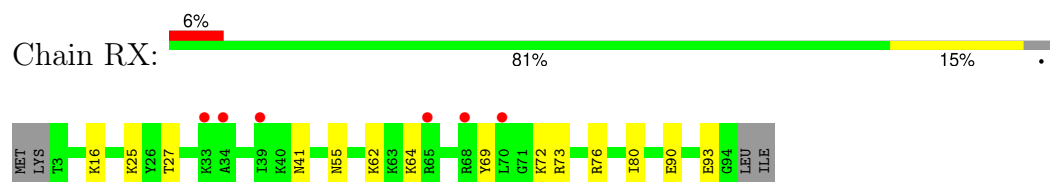
- Molecule 51: 50S ribosomal protein L22



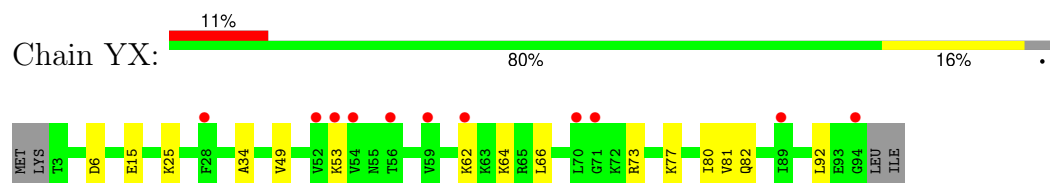
- Molecule 51: 50S ribosomal protein L22



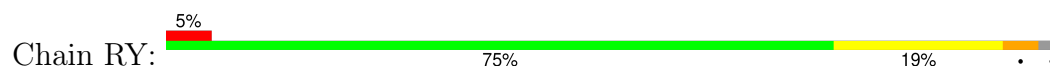
- Molecule 52: 50S ribosomal protein L23



- Molecule 52: 50S ribosomal protein L23

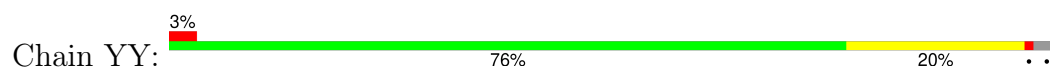


- Molecule 53: 50S ribosomal protein L24

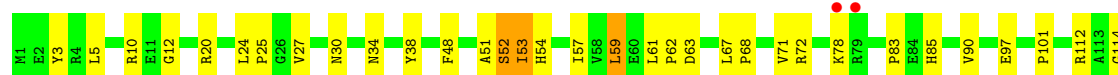




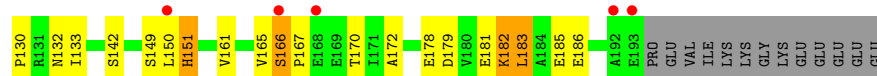
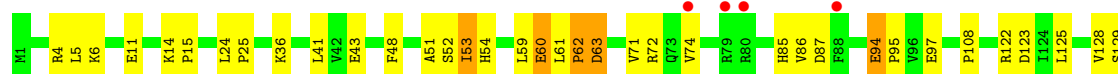
- Molecule 53: 50S ribosomal protein L24



- Molecule 54: 50S ribosomal protein L25



- Molecule 54: 50S ribosomal protein L25



4 Data and refinement statistics

Property	Value	Source
Space group	P 21 21 21	Depositor
Cell constants a, b, c, α , β , γ	210.09Å 450.32Å 622.89Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	49.85 – 4.14 49.85 – 4.14	Depositor EDS
% Data completeness (in resolution range)	98.5 (49.85-4.14) 98.4 (49.85-4.14)	Depositor EDS
R_{merge}	0.24	Depositor
R_{sym}	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ ¹	1.35 (at 4.14Å)	Xtriage
Refinement program	PHENIX 1.15.2_3472	Depositor
R, R_{free}	0.248 , 0.294 0.246 , 0.292	Depositor DCC
R_{free} test set	427005 reflections (4.34%)	wwPDB-VP
Wilson B-factor (Å ²)	171.1	Xtriage
Anisotropy	0.389	Xtriage
Bulk solvent k_{sol} (e/Å ³), B_{sol} (Å ²)	0.19 , 97.5	EDS
L-test for twinning ²	$\langle L \rangle = 0.39$, $\langle L^2 \rangle = 0.22$	Xtriage
Estimated twinning fraction	No twinning to report.	Xtriage
F_o, F_c correlation	0.92	EDS
Total number of atoms	291185	wwPDB-VP
Average B, all atoms (Å ²)	247.0	wwPDB-VP

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 1.91% of the height of the origin peak. No significant pseudotranslation is detected.*

¹Intensities estimated from amplitudes.

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

5 Model quality

5.1 Standard geometry

Bond lengths and bond angles in the following residue types are not validated in this section: SF4, MG, ZN

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	QA	0.31	0/36343	1.10	485/56720 (0.9%)
1	XA	0.44	28/36435 (0.1%)	1.31	648/56865 (1.1%)
2	QB	0.36	0/1942	0.67	0/2619
2	XB	0.37	0/1950	0.64	1/2630 (0.0%)
3	QC	0.36	0/1629	0.66	0/2195
3	XC	0.37	0/1629	0.61	0/2195
4	QD	0.45	1/1733 (0.1%)	0.65	0/2318
4	XD	0.52	2/1733 (0.1%)	0.70	2/2318 (0.1%)
5	QE	0.37	0/1171	0.67	0/1576
5	XE	0.43	0/1171	0.62	0/1576
6	QF	0.39	0/856	0.68	0/1154
6	XF	0.41	0/856	0.62	0/1154
7	QG	0.35	0/1276	0.63	1/1709 (0.1%)
7	XG	0.36	0/1276	0.61	0/1709
8	QH	0.40	0/1128	0.62	0/1517
8	XH	0.42	0/1128	0.66	0/1517
9	QI	0.42	0/831	0.74	0/1120
9	XI	0.36	0/849	0.72	0/1144
10	QJ	0.35	0/814	0.67	0/1095
10	XJ	0.68	1/790 (0.1%)	0.80	1/1063 (0.1%)
11	QK	0.36	0/900	0.57	0/1213
11	XK	0.39	0/879	0.59	0/1187
12	QL	0.41	0/991	0.70	1/1327 (0.1%)
12	XL	0.45	0/972	0.76	2/1301 (0.2%)
13	QM	0.35	0/931	0.75	0/1248
13	XM	0.37	0/924	0.66	0/1238
14	QN	0.67	1/501 (0.2%)	0.84	3/664 (0.5%)
14	XN	0.69	1/501 (0.2%)	0.89	2/664 (0.3%)
15	QO	0.38	0/745	0.57	0/992
15	XO	0.40	0/740	0.56	0/987
16	QP	0.40	0/721	0.64	0/970
16	XP	0.38	0/721	0.66	0/970

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
17	QQ	0.38	0/847	0.62	0/1131
17	XQ	0.47	0/847	0.64	0/1131
18	QR	0.38	0/579	0.56	0/768
18	XR	0.39	0/579	0.58	0/768
19	QS	0.35	0/680	0.72	1/915 (0.1%)
19	XS	0.36	0/689	0.70	0/926
20	QT	0.77	2/765 (0.3%)	1.14	8/1007 (0.8%)
20	XT	0.37	0/765	0.75	2/1007 (0.2%)
21	QU	0.34	0/221	0.58	0/288
21	XU	0.52	0/221	0.61	0/288
22	QV	0.28	0/1621	0.84	5/2523 (0.2%)
22	XV	0.44	0/1621	1.24	15/2523 (0.6%)
23	QX	0.41	0/459	1.04	0/715
23	XX	0.60	0/459	1.31	2/715 (0.3%)
24	R0	0.40	0/652	0.63	0/867
24	Y0	0.59	0/657	0.60	0/874
25	R1	0.54	0/753	0.68	0/1000
25	Y1	0.59	0/736	0.74	0/978
26	R2	0.37	0/583	0.62	0/771
26	Y2	0.47	0/577	0.62	0/764
27	R3	0.39	0/474	0.59	0/635
27	Y3	0.62	0/474	0.59	0/635
28	R4	0.33	0/357	0.60	0/483
28	Y4	1.56	2/366 (0.5%)	1.47	9/495 (1.8%)
29	R5	0.87	3/473 (0.6%)	0.79	2/639 (0.3%)
29	Y5	0.94	2/473 (0.4%)	0.77	1/639 (0.2%)
30	R6	0.95	3/460 (0.7%)	0.81	2/613 (0.3%)
30	Y6	1.33	6/460 (1.3%)	1.01	3/613 (0.5%)
31	R7	0.53	0/417	0.62	0/550
31	Y7	0.63	0/426	0.66	0/561
32	R8	0.43	0/525	0.88	4/691 (0.6%)
32	Y8	0.59	0/525	0.84	0/691
33	R9	0.62	1/310 (0.3%)	0.72	1/407 (0.2%)
33	Y9	0.64	0/310	0.73	0/407
34	RA	0.26	0/69520	1.00	605/108527 (0.6%)
34	YA	0.29	2/69543 (0.0%)	1.02	662/108563 (0.6%)
35	RB	0.57	0/2878	1.40	38/4490 (0.8%)
35	YB	0.72	0/2878	1.67	85/4490 (1.9%)
36	RD	0.52	0/2165	0.71	3/2919 (0.1%)
36	YD	0.64	0/2165	0.74	4/2919 (0.1%)
37	RE	0.50	0/1601	0.83	3/2160 (0.1%)
37	YE	0.66	0/1601	0.85	3/2160 (0.1%)
38	RF	0.49	0/1620	0.70	1/2194 (0.0%)

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
38	YF	0.67	0/1620	0.65	1/2194 (0.0%)
39	RG	0.41	0/1499	0.69	0/2016
39	YG	0.43	0/1499	0.68	0/2016
40	RH	0.39	0/1362	0.83	5/1841 (0.3%)
40	YH	0.68	1/1362 (0.1%)	0.86	6/1841 (0.3%)
41	RI	0.45	1/1151 (0.1%)	0.81	3/1558 (0.2%)
41	YI	0.45	1/1151 (0.1%)	0.79	0/1558
42	RN	0.45	0/1131	0.68	1/1525 (0.1%)
42	YN	0.63	0/1131	0.71	2/1525 (0.1%)
43	RO	0.51	0/943	0.65	0/1269
43	YO	0.60	0/943	0.63	0/1269
44	RP	0.44	0/1162	0.76	1/1544 (0.1%)
44	YP	0.54	0/1139	0.83	1/1514 (0.1%)
45	RQ	0.45	0/1143	0.73	0/1527
45	YQ	0.61	0/1143	0.77	2/1527 (0.1%)
46	RR	0.48	0/974	0.68	0/1302
46	YR	0.57	0/974	0.70	0/1302
47	RS	0.40	0/892	0.66	0/1187
47	YS	0.52	0/892	0.67	0/1187
48	RT	0.43	0/1155	0.69	0/1542
48	YT	0.54	0/1155	0.72	1/1542 (0.1%)
49	RU	0.49	0/982	0.62	0/1306
49	YU	0.70	0/982	0.62	0/1306
50	RV	0.48	0/790	0.74	1/1057 (0.1%)
50	YV	0.63	0/790	0.76	1/1057 (0.1%)
51	RW	0.52	0/911	0.63	0/1220
51	YW	0.68	0/911	0.64	0/1220
52	RX	0.52	0/739	0.60	0/993
52	YX	0.66	0/739	0.68	0/993
53	RY	0.72	4/831 (0.5%)	0.67	2/1108 (0.2%)
53	YY	0.73	1/831 (0.1%)	0.72	1/1108 (0.1%)
54	RZ	0.43	0/1493	0.89	6/2026 (0.3%)
54	YZ	0.51	0/1561	0.85	5/2119 (0.2%)
All	All	0.40	63/315379 (0.0%)	1.01	2644/471694 (0.6%)

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

Mol	Chain	#Chirality outliers	#Planarity outliers
1	XA	1	16

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Mol	Chain	#Chirality outliers	#Planarity outliers
28	Y4	1	1
34	RA	0	1
34	YA	0	6
37	RE	0	1
37	YE	0	1
50	RV	0	2
54	RZ	0	1
54	YZ	0	1
All	All	2	30

All (63) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
28	Y4	5	ILE	CA-CB	-21.62	1.05	1.54
28	Y4	4	GLY	N-CA	-18.71	1.18	1.46
1	XA	309	G	C3'-C2'	-15.74	1.35	1.52
30	R6	16	CYS	CB-SG	14.21	2.06	1.82
30	Y6	16	CYS	CB-SG	-14.07	1.58	1.82
20	QT	74	LYS	CA-CB	-13.70	1.23	1.53
29	Y5	32	PRO	N-CA	13.21	1.69	1.47
40	YH	12	PRO	N-CD	-13.08	1.29	1.47
30	Y6	13	CYS	CB-SG	-12.97	1.60	1.82
20	QT	73	HIS	N-CA	-12.33	1.21	1.46
1	XA	617	G	C1'-N9	12.04	1.66	1.48
14	QN	43	CYS	CB-SG	11.16	2.01	1.82
14	XN	43	CYS	CB-SG	10.98	2.00	1.82
4	XD	12	CYS	CB-SG	10.97	2.00	1.82
53	RY	102	CYS	CB-SG	-10.92	1.63	1.82
29	R5	34	PRO	N-CD	10.48	1.62	1.47
53	YY	79	CYS	CB-SG	-10.34	1.64	1.82
1	XA	309	G	C4'-O4'	-9.97	1.32	1.45
1	XA	309	G	C4'-C3'	9.41	1.63	1.53
29	R5	33	CYS	C-N	8.94	1.51	1.34
30	Y6	40	CYS	CB-SG	8.90	1.97	1.82
30	Y6	40	CYS	C-N	8.78	1.50	1.34
1	XA	116	A	P-O5'	8.69	1.68	1.59
1	XA	1358	U	P-O5'	8.49	1.68	1.59
1	XA	608	A	C1'-N9	-8.48	1.34	1.46
33	R9	29	ASN	C-N	8.38	1.50	1.34
53	RY	79	CYS	CB-SG	8.21	1.96	1.82
1	XA	1320	C	C2'-C1'	7.96	1.62	1.53
30	R6	13	CYS	CB-SG	-7.91	1.68	1.82

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
30	R6	41	PRO	N-CD	7.44	1.58	1.47
34	YA	745	G	O3'-P	-7.19	1.52	1.61
1	XA	608	A	C2'-C1'	7.09	1.61	1.53
41	RI	94	ALA	C-N	7.05	1.50	1.34
30	Y6	41	PRO	N-CD	7.05	1.57	1.47
1	XA	309	G	O4'-C1'	-6.81	1.32	1.41
4	QD	8	VAL	CB-CG1	6.79	1.67	1.52
1	XA	1347	G	C4'-C3'	6.59	1.60	1.53
1	XA	617	G	O4'-C1'	6.57	1.50	1.41
1	XA	636	U	P-O5'	6.49	1.66	1.59
1	XA	1320	C	C1'-N1	6.33	1.58	1.48
1	XA	1160	G	C1'-N9	6.18	1.58	1.48
53	RY	99	CYS	CB-SG	-6.10	1.71	1.82
1	XA	1158	C	C1'-N1	6.03	1.57	1.48
1	XA	1253	G	O3'-P	6.00	1.68	1.61
29	Y5	31	VAL	C-N	5.97	1.45	1.34
1	XA	1158	C	C2'-C1'	5.95	1.59	1.53
53	RY	76	CYS	CB-SG	-5.84	1.72	1.81
10	XJ	39	PRO	N-CD	-5.78	1.39	1.47
1	XA	1349	A	C1'-N9	5.77	1.57	1.48
1	XA	116	A	P-OP2	-5.67	1.39	1.49
4	XD	31	CYS	CB-SG	5.57	1.91	1.82
1	XA	1359	C	P-O5'	-5.46	1.54	1.59
1	XA	1158	C	C3'-C2'	5.37	1.58	1.52
30	Y6	51	GLU	CG-CD	-5.32	1.44	1.51
34	YA	745	G	C3'-O3'	-5.32	1.34	1.42
41	YI	109	ILE	C-N	5.26	1.46	1.34
1	XA	1440(B)	G	C5'-C4'	5.25	1.57	1.51
1	XA	635	G	O3'-P	5.13	1.67	1.61
1	XA	1225	A	C1'-N9	-5.13	1.39	1.46
29	R5	33	CYS	CB-SG	-5.13	1.73	1.81
1	XA	608	A	C3'-C2'	5.07	1.58	1.52
1	XA	309	G	C5'-C4'	5.06	1.57	1.51
1	XA	308	C	C2'-C1'	-5.01	1.47	1.53

All (2644) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	617	G	C4-N9-C1'	38.95	177.13	126.50
1	XA	617	G	C8-N9-C1'	-38.89	76.44	127.00
1	XA	1505	G	C8-N9-C1'	-27.87	90.77	127.00
1	XA	625	G	C8-N9-C1'	-27.83	90.82	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	1505	G	C4-N9-C1'	27.65	162.44	126.50
1	XA	877	C	C2-N1-C1'	25.01	146.31	118.80
1	XA	877	C	C6-N1-C1'	-23.72	92.33	120.80
1	XA	309	G	C4-N9-C1'	-23.02	96.57	126.50
1	XA	309	G	C8-N9-C1'	22.84	156.70	127.00
1	XA	1234	C	C2-N1-C1'	22.21	143.24	118.80
20	QT	74	LYS	N-CA-CB	21.24	148.84	110.60
1	XA	1440(B)	G	C8-N9-C1'	-21.03	99.66	127.00
1	XA	1440(B)	G	C4-N9-C1'	20.48	153.12	126.50
1	XA	1234	C	C6-N1-C1'	-20.33	96.41	120.80
34	YA	2111	C	C2-N1-C1'	20.05	140.86	118.80
34	YA	2111	C	C6-N1-C1'	-19.88	96.95	120.80
1	XA	618	C	C6-N1-C1'	-19.79	97.06	120.80
1	XA	1112	C	C6-N1-C1'	-19.56	97.33	120.80
1	XA	1348	U	C6-N1-C1'	-19.54	93.84	121.20
1	XA	186(B)	C	C2-N1-C1'	18.68	139.35	118.80
34	YA	2014	A	O5'-P-OP2	-18.66	88.31	110.70
1	XA	1112	C	C2-N1-C1'	18.30	138.93	118.80
1	XA	618	C	C2-N1-C1'	18.11	138.72	118.80
1	XA	972	C	C6-N1-C1'	-18.08	99.10	120.80
1	XA	972	C	C2-N1-C1'	18.07	138.68	118.80
1	XA	310	G	O5'-P-OP1	-17.82	89.31	110.70
1	XA	186(B)	C	C6-N1-C1'	-17.76	99.49	120.80
1	XA	625	G	C4-N9-C1'	17.56	149.32	126.50
1	XA	1358	U	O5'-P-OP1	17.50	131.70	110.70
28	Y4	5	ILE	CB-CA-C	17.14	145.87	111.60
1	XA	608	A	C8-N9-C1'	-16.92	97.25	127.70
1	XA	957	U	C2-N1-C1'	16.57	137.59	117.70
1	XA	877	C	O4'-C1'-N1	16.55	121.44	108.20
1	XA	878	G	C8-N9-C1'	-16.48	105.57	127.00
1	XA	309	G	O4'-C1'-N9	-16.30	95.16	108.20
1	XA	860	A	C8-N9-C1'	-15.99	98.92	127.70
1	XA	1320	C	N1-C1'-C2'	15.78	134.51	114.00
1	XA	1369	C	O5'-P-OP1	-15.53	91.73	105.70
1	XA	980	C	C2-N1-C1'	15.51	135.86	118.80
1	XA	635	G	P-O3'-C3'	15.38	138.15	119.70
37	RE	146	THR	C-N-CD	-15.28	86.99	120.60
1	XA	957	U	C6-N1-C1'	-15.26	99.83	121.20
1	XA	860	A	C4-N9-C1'	15.23	153.71	126.30
1	QA	194	C	C2-N1-C1'	14.87	135.16	118.80
1	XA	1348	U	C2-N1-C1'	14.68	135.32	117.70
1	XA	1320	C	C6-N1-C1'	-14.48	103.42	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	YA	887	A	O5'-P-OP2	-14.38	92.76	105.70
1	XA	309	G	O5'-P-OP1	14.31	127.87	110.70
1	XA	980	C	C6-N1-C1'	-14.23	103.73	120.80
54	RZ	166	SER	C-N-CD	-14.13	89.51	120.60
37	YE	146	THR	C-N-CD	-14.12	89.53	120.60
1	XA	674	G	C8-N9-C1'	-14.03	108.77	127.00
1	XA	1358	U	OP1-P-OP2	-13.99	98.61	119.60
1	XA	878	G	C4-N9-C1'	13.99	144.68	126.50
1	XA	674	G	C4-N9-C1'	13.92	144.60	126.50
1	QA	194	C	C6-N1-C1'	-13.60	104.48	120.80
35	YB	30	C	C6-N1-C2	-13.53	114.89	120.30
1	XA	625	G	O4'-C1'-N9	13.51	119.00	108.20
1	XA	1349	A	O4'-C1'-N9	13.16	118.73	108.20
34	YA	2442	C	C2-N1-C1'	13.03	133.13	118.80
40	RH	86	GLU	CB-CA-C	-12.83	84.74	110.40
1	XA	1302	U	C2-N1-C1'	12.81	133.07	117.70
1	XA	608	A	N9-C1'-C2'	12.73	130.55	114.00
34	YA	1076	C	C2-N1-C1'	12.72	132.79	118.80
54	YZ	166	SER	C-N-CD	-12.57	92.94	120.60
34	YA	270(L)	C	C2-N1-C1'	12.42	132.47	118.80
1	XA	1244	C	O5'-P-OP2	-12.26	94.67	105.70
1	XA	1320	C	C2-N1-C1'	12.11	132.12	118.80
34	YA	2442	C	C6-N1-C1'	-12.10	106.28	120.80
1	XA	1301	U	C2-N1-C1'	12.03	132.13	117.70
1	XA	686	U	C2-N1-C1'	12.01	132.11	117.70
1	XA	723	U	C2-N1-C1'	11.76	131.82	117.70
34	YA	1076	C	C6-N1-C1'	-11.63	106.84	120.80
1	XA	132	C	OP1-P-O3'	11.60	130.72	105.20
34	RA	2789	C	C2-N1-C1'	11.56	131.52	118.80
1	XA	116	A	OP1-P-OP2	-11.50	102.34	119.60
1	QA	856	C	C2-N1-C1'	11.47	131.42	118.80
34	YA	270(L)	C	C6-N1-C1'	-11.38	107.15	120.80
1	XA	1232	U	O5'-P-OP1	11.36	124.33	110.70
1	XA	618	C	O4'-C1'-N1	11.31	117.25	108.20
1	XA	1507	A	C4-N9-C1'	-11.26	106.03	126.30
34	RA	2836	U	C2-N1-C1'	11.24	131.19	117.70
1	XA	1527	C	C2-N1-C1'	11.13	131.04	118.80
1	XA	607	A	C4-N9-C1'	11.10	146.29	126.30
1	XA	607	A	C8-N9-C1'	-11.08	107.76	127.70
1	XA	1302	U	C6-N1-C1'	-11.00	105.80	121.20
1	XA	1347	G	C4-N9-C1'	-10.94	112.28	126.50
1	XA	1359	C	O5'-P-OP1	-10.93	95.86	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	1307	U	O5'-P-OP1	-10.88	95.91	105.70
34	YA	2394	C	C2-N1-C1'	10.86	130.75	118.80
1	XA	1344	C	C2-N1-C1'	10.85	130.74	118.80
1	XA	68(T)	C	C2-N1-C1'	10.83	130.72	118.80
34	YA	2064	C	C2-N1-C1'	10.83	130.71	118.80
34	YA	1535	U	C2-N1-C1'	10.82	130.68	117.70
1	XA	948	C	C2-N1-C1'	-10.81	106.91	118.80
1	XA	1347	G	C8-N9-C1'	10.77	141.00	127.00
34	YA	2394	C	OP1-P-OP2	-10.76	103.46	119.60
34	RA	2789	C	C6-N1-C1'	-10.67	108.00	120.80
1	XA	1507	A	O4'-C1'-N9	10.58	116.67	108.20
1	XA	1507	A	C8-N9-C1'	10.58	146.74	127.70
34	RA	2431	U	C2-N1-C1'	10.57	130.38	117.70
1	XA	259	G	O5'-P-OP2	-10.53	96.22	105.70
1	XA	1375	A	C4-N9-C1'	10.53	145.25	126.30
1	XA	1375	A	C8-N9-C1'	-10.52	108.77	127.70
1	XA	677	U	C2-N1-C1'	10.51	130.31	117.70
1	XA	686	U	C6-N1-C1'	-10.50	106.50	121.20
1	QA	856	C	C6-N1-C1'	-10.49	108.22	120.80
1	XA	1343	G	C4-N9-C1'	10.48	140.13	126.50
34	RA	1534	G	C4-N9-C1'	10.47	140.12	126.50
34	YA	2443	C	C2-N1-C1'	10.47	130.32	118.80
1	XA	1440(C)	G	C8-N9-C1'	-10.44	113.43	127.00
1	XA	1259	C	C2-N1-C1'	10.44	130.28	118.80
1	XA	1328	C	O5'-P-OP1	-10.43	96.31	105.70
1	QA	677	U	C2-N1-C1'	10.42	130.21	117.70
1	QA	1234	C	C2-N1-C1'	10.42	130.26	118.80
1	XA	636	U	C2-N1-C1'	10.41	130.19	117.70
1	XA	1343	G	C8-N9-C1'	-10.39	113.49	127.00
34	YA	2098	U	C2-N1-C1'	10.39	130.16	117.70
34	RA	1534	G	C8-N9-C1'	-10.38	113.50	127.00
1	XA	956	U	C2-N1-C1'	10.37	130.15	117.70
1	XA	1438	G	O5'-P-OP1	10.33	123.10	110.70
1	QA	980	C	C2-N1-C1'	10.32	130.15	118.80
1	XA	1301	U	C6-N1-C1'	-10.29	106.79	121.20
1	XA	1112	C	O4'-C1'-N1	10.28	116.42	108.20
1	XA	617	G	O5'-P-OP2	-10.27	96.45	105.70
1	XA	608	A	O4'-C1'-N9	10.26	116.40	108.20
1	XA	981	U	O5'-P-OP2	-10.25	96.48	105.70
1	XA	1527	C	C6-N1-C1'	-10.24	108.51	120.80
1	XA	1111	A	C4-N9-C1'	10.23	144.71	126.30
1	XA	1111	A	C8-N9-C1'	-10.16	109.41	127.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	1260	C	C2-N1-C1'	10.13	129.94	118.80
1	XA	1328	C	O5'-P-OP2	10.12	122.84	110.70
34	YA	1094	U	C2-N1-C1'	10.11	129.83	117.70
34	YA	2064	C	C6-N1-C1'	-10.10	108.68	120.80
34	YA	2822	G	O5'-P-OP2	-10.10	96.61	105.70
34	RA	419	C	C2-N1-C1'	10.08	129.89	118.80
1	QA	88	C	C2-N1-C1'	10.04	129.85	118.80
1	QA	1037	C	C2-N1-C1'	10.04	129.84	118.80
1	XA	1440(C)	G	C4-N9-C1'	10.03	139.54	126.50
34	RA	270(L)	C	C2-N1-C1'	10.02	129.82	118.80
1	XA	723	U	C6-N1-C1'	-10.01	107.19	121.20
1	XA	948	C	C6-N1-C1'	9.96	132.75	120.80
34	YA	2394	C	C6-N1-C1'	-9.96	108.85	120.80
1	XA	1344	C	C6-N1-C1'	-9.93	108.89	120.80
1	XA	186(A)	C	C2-N1-C1'	9.92	129.71	118.80
34	YA	2789	C	C2-N1-C1'	9.91	129.70	118.80
1	XA	385	C	C2-N1-C1'	9.90	129.70	118.80
1	XA	1355	G	C4-N9-C1'	9.90	139.37	126.50
1	XA	1355	G	C8-N9-C1'	-9.90	114.13	127.00
34	YA	2474	C	C2-N1-C1'	9.87	129.66	118.80
34	RA	2189	U	C2-N1-C1'	9.87	129.54	117.70
1	XA	68(T)	C	C6-N1-C1'	-9.87	108.95	120.80
30	Y6	43	CYS	N-CA-CB	-9.83	92.90	110.60
1	XA	1328	C	OP1-P-OP2	-9.83	104.86	119.60
34	YA	1078	U	C2-N1-C1'	9.83	129.49	117.70
34	RA	1742	C	C2-N1-C1'	9.79	129.57	118.80
1	XA	1297	C	C2-N1-C1'	9.74	129.51	118.80
34	RA	1644	C	C2-N1-C1'	9.70	129.47	118.80
1	XA	1356	G	C4-N9-C1'	9.65	139.04	126.50
34	YA	2443	C	C6-N1-C1'	-9.64	109.23	120.80
1	XA	308	C	P-O3'-C3'	-9.64	108.14	119.70
1	XA	623	C	C2-N1-C1'	9.59	129.34	118.80
34	YA	1644	C	C2-N1-C1'	9.59	129.34	118.80
34	RA	2836	U	C6-N1-C1'	-9.57	107.80	121.20
1	XA	1356	G	C8-N9-C1'	-9.57	114.55	127.00
1	XA	1259	C	C6-N1-C1'	-9.55	109.34	120.80
1	QA	536	C	C2-N1-C1'	9.54	129.29	118.80
1	QA	1234	C	C6-N1-C1'	-9.53	109.37	120.80
37	YE	146	THR	C-N-CA	9.52	161.97	122.00
34	YA	2620	C	C2-N1-C1'	9.51	129.26	118.80
1	XA	635	G	N9-C1'-C2'	-9.49	101.56	112.00
34	YA	2098	U	C6-N1-C1'	-9.48	107.93	121.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	YA	2871	C	C2-N1-C1'	9.48	129.23	118.80
1	XA	1342	C	C2-N1-C1'	9.45	129.20	118.80
34	RA	373	U	C2-N1-C1'	9.43	129.01	117.70
1	XA	1297	C	C6-N1-C1'	-9.38	109.54	120.80
1	XA	692	U	C2-N1-C1'	9.38	128.96	117.70
1	QA	980	C	C6-N1-C1'	-9.34	109.59	120.80
1	XA	1505	G	N9-C1'-C2'	9.31	126.11	114.00
1	XA	1343	G	N9-C1'-C2'	-9.27	101.81	112.00
1	QA	1260	C	C6-N1-C1'	-9.26	109.68	120.80
1	XA	310	G	OP1-P-OP2	-9.25	105.72	119.60
1	QA	995	C	C2-N1-C1'	9.22	128.95	118.80
1	XA	309	G	C5'-C4'-O4'	-9.20	98.06	109.10
1	QA	88	C	C6-N1-C1'	-9.20	109.76	120.80
1	XA	617	G	O4'-C1'-N9	9.19	115.55	108.20
34	RA	270(L)	C	C6-N1-C1'	-9.18	109.78	120.80
1	XA	986	A	C4-N9-C1'	9.18	142.82	126.30
1	XA	68(S)	C	C2-N1-C1'	9.18	128.90	118.80
1	QA	1037	C	C6-N1-C1'	-9.16	109.81	120.80
1	XA	669	U	C2-N1-C1'	9.15	128.68	117.70
34	RA	419	C	C6-N1-C1'	-9.15	109.82	120.80
54	RZ	166	SER	C-N-CA	9.14	160.40	122.00
1	XA	677	U	C6-N1-C1'	-9.14	108.40	121.20
34	YA	888	C	O5'-P-OP2	-9.13	97.48	105.70
34	YA	2342	C	C2-N1-C1'	9.12	128.84	118.80
1	XA	986	A	C8-N9-C1'	-9.12	111.29	127.70
34	YA	1535	U	C6-N1-C1'	-9.10	108.47	121.20
30	R6	16	CYS	CA-CB-SG	9.09	130.37	114.00
1	XA	186(A)	C	C6-N1-C1'	-9.09	109.89	120.80
20	QT	73	HIS	N-CA-C	9.09	135.53	111.00
20	QT	74	LYS	N-CA-C	-9.08	86.47	111.00
34	YA	2787	C	C2-N1-C1'	9.08	128.78	118.80
34	YA	1941	C	C2-N1-C1'	9.07	128.78	118.80
34	YA	373	U	C2-N1-C1'	9.06	128.58	117.70
1	XA	1342	C	C6-N1-C1'	-9.06	109.93	120.80
34	YA	2794	C	C2-N1-C1'	9.05	128.75	118.80
1	XA	385	C	C6-N1-C1'	-9.04	109.95	120.80
1	XA	739	C	O5'-P-OP1	9.03	121.54	110.70
1	XA	115	G	C4-N9-C1'	9.02	138.22	126.50
1	XA	860	A	O5'-P-OP2	-9.02	97.58	105.70
1	XA	498	U	C2-N1-C1'	9.01	128.52	117.70
1	XA	1157	A	O4'-C1'-N9	9.01	115.41	108.20
34	RA	1742	C	C6-N1-C1'	-9.00	110.00	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	YA	2789	C	C6-N1-C1'	-9.00	110.00	120.80
1	QA	714	G	C4-N9-C1'	8.99	138.18	126.50
34	YA	2474	C	C6-N1-C1'	-8.99	110.02	120.80
34	RA	2431	U	C6-N1-C1'	-8.98	108.62	121.20
1	XA	1129	C	C2-N1-C1'	8.98	128.68	118.80
1	QA	1114	C	C2-N1-C1'	8.96	128.65	118.80
34	RA	2691	C	C2-N1-C1'	8.96	128.65	118.80
1	XA	115	G	C8-N9-C1'	-8.94	115.37	127.00
1	QA	1028(C)	C	C2-N1-C1'	8.93	128.62	118.80
1	QA	1131	G	C4-N9-C1'	8.92	138.10	126.50
1	XA	115	G	P-O3'-C3'	8.92	130.41	119.70
1	QA	714	G	C8-N9-C1'	-8.92	115.41	127.00
34	YA	2164	C	C2-N1-C1'	8.91	128.60	118.80
1	QA	677	U	C6-N1-C1'	-8.91	108.73	121.20
1	XA	956	U	C6-N1-C1'	-8.90	108.74	121.20
1	XA	636	U	C6-N1-C1'	-8.89	108.75	121.20
28	Y4	39	CYS	C-N-CA	8.88	143.91	121.70
1	QA	1325	C	C2-N1-C1'	8.88	128.56	118.80
1	XA	1329	A	OP1-P-OP2	-8.87	106.29	119.60
35	RB	22	U	C5-C6-N1	8.86	127.13	122.70
1	QA	1131	G	C8-N9-C1'	-8.85	115.49	127.00
1	XA	856	C	C2-N1-C1'	8.85	128.53	118.80
34	YA	2111	C	O4'-C1'-N1	8.85	115.28	108.20
34	RA	2114	A	C4-N9-C1'	8.83	142.20	126.30
34	YA	2871	C	C6-N1-C1'	-8.83	110.20	120.80
1	XA	981	U	OP1-P-OP2	8.82	132.84	119.60
1	XA	1347	G	C5'-C4'-C3'	8.82	130.12	116.00
1	QA	1440(K)	C	C2-N1-C1'	8.82	128.50	118.80
34	RA	1644	C	C6-N1-C1'	-8.81	110.22	120.80
34	RA	2874	C	C2-N1-C1'	8.80	128.48	118.80
1	QA	1235	U	C2-N1-C1'	8.78	128.23	117.70
1	QA	400	C	C2-N1-C1'	8.77	128.45	118.80
1	QA	1277	C	C2-N1-C1'	8.77	128.44	118.80
1	XA	623	C	C6-N1-C1'	-8.75	110.30	120.80
34	YA	1644	C	C6-N1-C1'	-8.75	110.30	120.80
1	XA	136	C	C2-N1-C1'	8.73	128.41	118.80
34	YA	2502	G	C8-N9-C1'	-8.73	115.65	127.00
34	RA	2114	A	C8-N9-C1'	-8.72	112.00	127.70
34	YA	1534	G	C8-N9-C1'	-8.72	115.66	127.00
34	YA	12	U	C2-N1-C1'	8.70	128.15	117.70
34	RA	1882	C	C2-N1-C1'	8.70	128.37	118.80
34	YA	2620	C	C6-N1-C1'	-8.70	110.37	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	536	C	C6-N1-C1'	-8.69	110.37	120.80
1	XA	309	G	OP2-P-O3'	8.69	124.31	105.20
1	QA	379	C	C2-N1-C1'	8.65	128.32	118.80
34	YA	413	C	C2-N1-C1'	8.65	128.32	118.80
34	YA	1790	C	C2-N1-C1'	8.64	128.31	118.80
34	RA	1391	U	C2-N1-C1'	8.60	128.01	117.70
1	XA	826	C	C2-N1-C1'	8.59	128.25	118.80
34	YA	1094	U	C6-N1-C1'	-8.59	109.17	121.20
1	QA	43	C	C2-N1-C1'	8.59	128.25	118.80
35	YB	60	C	C6-N1-C2	-8.59	116.87	120.30
34	RA	338	G	C8-N9-C1'	-8.58	115.85	127.00
1	QA	723	U	C2-N1-C1'	8.57	127.99	117.70
1	XA	1198	G	C8-N9-C1'	-8.57	115.85	127.00
1	XA	267	C	O5'-P-OP1	-8.57	97.98	105.70
1	QA	1301	U	C2-N1-C1'	8.57	127.98	117.70
34	RA	338	G	C4-N9-C1'	8.57	137.64	126.50
34	YA	2619	C	C2-N1-C1'	8.57	128.22	118.80
1	XA	1277	C	C2-N1-C1'	8.56	128.22	118.80
1	XA	1150	U	C2-N1-C1'	8.55	127.97	117.70
34	YA	1611	C	C2-N1-C1'	8.53	128.19	118.80
34	RA	2098	U	C2-N1-C1'	8.53	127.94	117.70
1	XA	1198	G	C4-N9-C1'	8.53	137.59	126.50
35	YB	54	G	N1-C6-O6	8.53	125.02	119.90
1	QA	498	U	C2-N1-C1'	8.52	127.92	117.70
1	XA	1327	C	C2-N1-C1'	8.51	128.17	118.80
34	YA	1403	C	C2-N1-C1'	8.49	128.14	118.80
34	YA	2312	U	C2-N1-C1'	8.49	127.89	117.70
2	XB	89	GLY	C-N-CA	8.48	142.91	121.70
54	YZ	166	SER	C-N-CA	8.48	157.62	122.00
34	YA	1534	G	C4-N9-C1'	8.48	137.52	126.50
1	QA	385	C	C2-N1-C1'	8.47	128.12	118.80
1	XA	201(C)	U	C2-N1-C1'	8.47	127.87	117.70
30	Y6	43	CYS	CB-CA-C	8.47	127.34	110.40
34	RA	904	C	C2-N1-C1'	8.47	128.11	118.80
1	QA	995	C	C6-N1-C1'	-8.46	110.64	120.80
34	RA	2559	C	C2-N1-C1'	8.46	128.11	118.80
1	QA	193	C	C2-N1-C1'	8.46	128.10	118.80
37	RE	146	THR	C-N-CA	8.44	157.45	122.00
1	XA	68(Q)	C	C2-N1-C1'	8.44	128.09	118.80
34	RA	2506	U	C2-N1-C1'	8.44	127.83	117.70
34	RA	508	G	C4-N9-C1'	8.43	137.46	126.50
1	XA	68(S)	C	C6-N1-C1'	-8.42	110.69	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	RA	384	U	C2-N1-C1'	8.41	127.80	117.70
1	XA	1158	C	C2-N1-C1'	-8.41	109.55	118.80
34	RA	2189	U	C6-N1-C1'	-8.40	109.44	121.20
34	RA	2006	C	C2-N1-C1'	8.40	128.04	118.80
34	RA	1506	C	C2-N1-C1'	8.39	128.03	118.80
34	YA	1078	U	C6-N1-C1'	-8.38	109.47	121.20
34	RA	280	C	C2-N1-C1'	8.38	128.01	118.80
34	RA	35	G	C8-N9-C1'	-8.37	116.12	127.00
1	XA	68(V)	U	C2-N1-C1'	8.36	127.73	117.70
40	YH	12	PRO	CA-N-CD	8.35	123.40	111.70
1	QA	1313	U	C2-N1-C1'	8.35	127.72	117.70
34	RA	508	G	C8-N9-C1'	-8.34	116.15	127.00
34	YA	2342	C	C6-N1-C1'	-8.34	110.79	120.80
34	RA	714	U	C2-N1-C1'	8.33	127.70	117.70
34	YA	1882	C	C2-N1-C1'	8.33	127.96	118.80
34	YA	2787	C	C6-N1-C1'	-8.32	110.81	120.80
34	RA	1437	C	C2-N1-C1'	8.31	127.94	118.80
34	RA	35	G	C4-N9-C1'	8.31	137.31	126.50
1	XA	1129	C	C6-N1-C1'	-8.31	110.83	120.80
34	YA	2502	G	C4-N9-C1'	8.30	137.30	126.50
1	XA	1345	U	C2-N1-C1'	8.30	127.66	117.70
34	RA	2803	C	C2-N1-C1'	8.30	127.93	118.80
4	XD	18	LYS	CD-CE-NZ	8.30	130.79	111.70
34	YA	2794	C	C6-N1-C1'	-8.29	110.85	120.80
34	YA	1941	C	C6-N1-C1'	-8.29	110.85	120.80
34	YA	2501	C	C2-N1-C1'	8.28	127.91	118.80
34	YA	2889	C	C2-N1-C1'	8.28	127.91	118.80
1	XA	754	C	N1-C1'-C2'	8.27	124.75	114.00
1	QA	1145	C	C2-N1-C1'	8.27	127.89	118.80
1	XA	1226	C	C2-N1-C1'	-8.26	109.71	118.80
1	XA	323	U	C2-N1-C1'	8.25	127.61	117.70
1	QA	1440(K)	C	C6-N1-C1'	-8.24	110.91	120.80
34	RA	2787	C	C2-N1-C1'	8.24	127.86	118.80
1	XA	1260	C	C2-N1-C1'	8.24	127.86	118.80
1	QA	624	C	C2-N1-C1'	8.22	127.84	118.80
1	XA	116	A	O5'-P-OP2	8.20	120.54	110.70
34	YA	509	C	C2-N1-C1'	8.20	127.82	118.80
34	RA	2691	C	C6-N1-C1'	-8.18	110.98	120.80
1	QA	1114	C	C6-N1-C1'	-8.18	110.98	120.80
36	YD	34	VAL	N-CA-C	-8.16	88.97	111.00
1	XA	1145	C	C2-N1-C1'	8.16	127.77	118.80
34	YA	2815	C	C2-N1-C1'	8.15	127.77	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	YA	2164	C	C6-N1-C1'	-8.13	111.05	120.80
1	QA	1325	C	C6-N1-C1'	-8.12	111.06	120.80
28	Y4	5	ILE	N-CA-C	-8.11	89.11	111.00
34	RA	2554	U	C2-N1-C1'	8.11	127.43	117.70
1	QA	163	C	C2-N1-C1'	8.10	127.71	118.80
34	YA	856	C	C2-N1-C1'	8.09	127.70	118.80
1	QA	1028(C)	C	C6-N1-C1'	-8.08	111.10	120.80
1	QA	1219	U	C2-N1-C1'	8.08	127.40	117.70
34	RA	1870	C	C2-N1-C1'	8.08	127.69	118.80
1	QA	443	C	C2-N1-C1'	8.07	127.68	118.80
34	YA	1742	C	C2-N1-C1'	8.06	127.67	118.80
35	YB	30	C	C5-C6-N1	8.06	125.03	121.00
34	RA	1905	C	C2-N1-C1'	8.06	127.66	118.80
34	YA	2619	C	C6-N1-C1'	-8.05	111.13	120.80
1	XA	692	U	C6-N1-C1'	-8.05	109.93	121.20
1	QA	400	C	C6-N1-C1'	-8.04	111.15	120.80
34	RA	1881	C	C2-N1-C1'	8.04	127.64	118.80
34	RA	2168	G	C8-N9-C1'	-8.04	116.55	127.00
1	XA	1253	G	P-O3'-C3'	-8.04	110.05	119.70
34	YA	1880	C	C2-N1-C1'	8.04	127.64	118.80
34	RA	2874	C	C6-N1-C1'	-8.03	111.17	120.80
1	XA	856	C	C6-N1-C1'	-8.03	111.17	120.80
29	Y5	32	PRO	CA-N-CD	-8.02	100.27	111.50
1	XA	1197	G	C4-N9-C1'	8.02	136.93	126.50
1	XA	1277	C	C6-N1-C1'	-8.02	111.18	120.80
40	RH	152	ARG	C-N-CA	8.01	141.73	121.70
1	XA	669	U	C6-N1-C1'	-8.01	109.99	121.20
1	QA	1277	C	C6-N1-C1'	-8.01	111.19	120.80
1	XA	133	U	OP1-P-OP2	-8.00	107.60	119.60
34	RA	1649	G	C8-N9-C1'	-7.99	116.61	127.00
34	RA	373	U	C6-N1-C1'	-7.99	110.02	121.20
34	YA	270(Q)	C	C2-N1-C1'	7.99	127.59	118.80
40	YH	151	ILE	N-CA-C	-7.99	89.43	111.00
34	RA	1882	C	C6-N1-C1'	-7.99	111.22	120.80
34	RA	2168	G	C4-N9-C1'	7.97	136.86	126.50
34	RA	2456	C	C2-N1-C1'	7.97	127.56	118.80
1	XA	1202	G	C4-N9-C1'	7.97	136.86	126.50
1	XA	943	U	C2-N1-C1'	7.97	127.26	117.70
34	YA	2127	G	C4-N9-C1'	7.97	136.86	126.50
34	YA	420	C	C2-N1-C1'	7.96	127.56	118.80
1	QA	1367	C	O4'-C1'-N1	7.96	114.57	108.20
1	XA	1270	C	C2-N1-C1'	7.96	127.55	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	RA	1879	C	C2-N1-C1'	7.95	127.55	118.80
1	XA	1197	G	C8-N9-C1'	-7.95	116.67	127.00
1	XA	136	C	C6-N1-C1'	-7.94	111.27	120.80
1	QA	623	C	C2-N1-C1'	7.93	127.53	118.80
34	YA	721	C	C2-N1-C1'	7.93	127.53	118.80
34	YA	860	U	O4'-C1'-N1	7.93	114.54	108.20
34	RA	2291	U	C2-N1-C1'	7.92	127.21	117.70
1	XA	1347	G	P-O3'-C3'	7.92	129.20	119.70
34	YA	541	C	C2-N1-C1'	7.92	127.51	118.80
34	YA	1790	C	C6-N1-C1'	-7.92	111.30	120.80
1	QA	110	C	C2-N1-C1'	7.92	127.51	118.80
1	XA	1401	G	C4-N9-C1'	7.91	136.78	126.50
1	QA	1369	C	C2-N1-C1'	7.91	127.50	118.80
1	QA	43	C	C6-N1-C1'	-7.90	111.32	120.80
1	XA	1202	G	C8-N9-C1'	-7.90	116.73	127.00
35	YB	70	C	C6-N1-C2	-7.89	117.14	120.30
34	YA	2127	G	C8-N9-C1'	-7.88	116.75	127.00
34	YA	413	C	C6-N1-C1'	-7.88	111.34	120.80
34	YA	1533	C	C2-N1-C1'	7.88	127.46	118.80
1	QA	379	C	C6-N1-C1'	-7.87	111.36	120.80
1	XA	36	C	C2-N1-C1'	7.87	127.46	118.80
1	QA	984	C	C2-N1-C1'	7.86	127.44	118.80
34	RA	2343	C	C2-N1-C1'	7.85	127.43	118.80
1	QA	201(A)	C	C2-N1-C1'	7.84	127.43	118.80
34	YA	1951	U	C2-N1-C1'	7.84	127.11	117.70
34	YA	2087	G	C8-N9-C1'	-7.83	116.81	127.00
34	RA	2658	C	C2-N1-C1'	7.83	127.41	118.80
1	XA	1401	G	C8-N9-C1'	-7.82	116.83	127.00
1	XA	381	C	C2-N1-C1'	7.82	127.40	118.80
34	YA	2656	U	C2-N1-C1'	7.82	127.08	117.70
34	RA	141(B)	C	C2-N1-C1'	7.82	127.40	118.80
34	RA	2178	C	C2-N1-C1'	7.81	127.39	118.80
1	XA	836	G	C8-N9-C1'	-7.81	116.84	127.00
1	QA	186(H)	C	C2-N1-C1'	7.81	127.39	118.80
1	QA	1235	U	C6-N1-C1'	-7.81	110.27	121.20
34	YA	2087	G	C4-N9-C1'	7.81	136.65	126.50
34	YA	1611	C	C6-N1-C1'	-7.80	111.44	120.80
1	XA	826	C	C6-N1-C1'	-7.80	111.44	120.80
1	XA	1505	G	O5'-P-OP2	7.79	120.05	110.70
34	RA	2695	C	C2-N1-C1'	7.79	127.37	118.80
35	RB	11	C	C6-N1-C2	-7.79	117.18	120.30
1	QA	838(B)	U	C2-N1-C1'	7.79	127.05	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	YA	2251	G	C4-N9-C1'	7.78	136.62	126.50
35	YB	37	C	N3-C2-O2	-7.78	116.45	121.90
1	XA	836	G	C4-N9-C1'	7.78	136.61	126.50
1	XA	1426	C	C2-N1-C1'	7.77	127.35	118.80
34	RA	1087	G	O4'-C1'-N9	-7.77	101.98	108.20
34	RA	2559	C	C6-N1-C1'	-7.76	111.49	120.80
1	XA	1158	C	C6-N1-C1'	7.76	130.11	120.80
34	YA	1403	C	C6-N1-C1'	-7.76	111.49	120.80
34	YA	114	U	C2-N1-C1'	7.75	127.00	117.70
1	QA	193	C	C6-N1-C1'	-7.75	111.50	120.80
34	YA	2836	U	C2-N1-C1'	7.75	126.99	117.70
35	YB	111	U	C5-C4-O4	7.74	130.54	125.90
1	QA	320	C	C2-N1-C1'	7.74	127.31	118.80
1	QA	385	C	C6-N1-C1'	-7.74	111.52	120.80
1	XA	1327	C	C6-N1-C1'	-7.73	111.52	120.80
28	Y4	3	GLU	C-N-CA	-7.73	106.06	122.30
34	RA	2130	U	C2-N1-C1'	7.73	126.97	117.70
1	XA	1058	G	O5'-P-OP1	7.72	119.97	110.70
22	XV	15	G	C4-N9-C1'	7.72	136.54	126.50
1	XA	68(Q)	C	C6-N1-C1'	-7.72	111.54	120.80
34	RA	904	C	C6-N1-C1'	-7.72	111.54	120.80
1	XA	1325	C	C2-N1-C1'	7.72	127.29	118.80
1	XA	1058	G	C4-N9-C1'	7.71	136.53	126.50
34	YA	2251	G	C8-N9-C1'	-7.71	116.97	127.00
1	XA	1049	U	C2-N1-C1'	7.71	126.95	117.70
1	XA	322	C	C2-N1-C1'	7.70	127.27	118.80
34	YA	373	U	C6-N1-C1'	-7.69	110.43	121.20
34	RA	1533	C	C2-N1-C1'	7.69	127.26	118.80
34	YA	2235	G	C4-N9-C1'	7.69	136.49	126.50
1	XA	1140	C	C2-N1-C1'	7.67	127.24	118.80
1	XA	310	G	O5'-P-OP2	7.67	119.90	110.70
34	RA	2006	C	C6-N1-C1'	-7.67	111.60	120.80
34	RA	2656	U	C2-N1-C1'	7.66	126.89	117.70
1	QA	705	U	C2-N1-C1'	7.66	126.89	117.70
34	RA	280	C	C6-N1-C1'	-7.66	111.61	120.80
1	XA	1381	U	C2'-C3'-O3'	7.66	126.35	109.50
34	RA	544	C	C2-N1-C1'	7.65	127.21	118.80
34	YA	414	C	C2-N1-C1'	7.64	127.21	118.80
1	QA	186(B)	C	C2-N1-C1'	7.64	127.20	118.80
1	XA	498	U	C6-N1-C1'	-7.64	110.51	121.20
14	QN	24	CYS	CA-CB-SG	7.64	127.75	114.00
1	XA	1260	C	C6-N1-C1'	-7.64	111.64	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
22	XV	15	G	N3-C4-C5	-7.63	124.78	128.60
34	RA	63	U	N1-C1'-C2'	7.63	123.92	114.00
35	YB	10	C	C6-N1-C2	-7.63	117.25	120.30
34	YA	2501	C	C6-N1-C1'	-7.62	111.65	120.80
34	RA	1437	C	C6-N1-C1'	-7.62	111.65	120.80
34	RA	1225	C	C2-N1-C1'	7.62	127.19	118.80
34	YA	2235	G	C8-N9-C1'	-7.62	117.10	127.00
34	RA	114	U	C2-N1-C1'	7.61	126.83	117.70
34	RA	1506	C	C6-N1-C1'	-7.60	111.67	120.80
1	XA	1058	G	C8-N9-C1'	-7.60	117.11	127.00
1	QA	957	U	C2-N1-C1'	7.60	126.82	117.70
34	RA	613	U	C2-N1-C1'	7.60	126.82	117.70
1	XA	1233	G	C4-N9-C1'	7.60	136.38	126.50
34	YA	2889	C	C6-N1-C1'	-7.60	111.68	120.80
1	XA	1358	U	O5'-C5'-C4'	7.59	126.13	111.70
1	QA	524	G	C4-N9-C1'	7.59	136.37	126.50
34	RA	1790	C	C2-N1-C1'	7.59	127.14	118.80
34	YA	1990	C	C2-N1-C1'	7.59	127.14	118.80
34	YA	2431	U	C2-N1-C1'	7.59	126.80	117.70
34	RA	2803	C	C6-N1-C1'	-7.58	111.70	120.80
1	QA	624	C	C6-N1-C1'	-7.58	111.70	120.80
34	RA	2787	C	C6-N1-C1'	-7.58	111.70	120.80
1	QA	989	C	C2-N1-C1'	7.58	127.14	118.80
1	XA	323	U	OP1-P-OP2	-7.58	108.23	119.60
1	XA	68(R)	U	C2-N1-C1'	7.57	126.78	117.70
34	RA	2832	U	P-O3'-C3'	7.56	128.78	119.70
34	YA	1882	C	C6-N1-C1'	-7.56	111.72	120.80
34	RA	2666	C	C2-N1-C1'	7.55	127.11	118.80
34	RA	413	C	C2-N1-C1'	7.55	127.10	118.80
1	QA	524	G	C8-N9-C1'	-7.54	117.19	127.00
1	QA	692	U	C2-N1-C1'	7.54	126.75	117.70
34	RA	721	C	C2-N1-C1'	7.54	127.09	118.80
34	YA	2132	U	C2-N1-C1'	7.54	126.74	117.70
34	YA	509	C	C6-N1-C1'	-7.53	111.76	120.80
34	RA	2262	U	C2-N1-C1'	7.53	126.74	117.70
1	QA	1145	C	C6-N1-C1'	-7.52	111.77	120.80
34	YA	2130	U	C2-N1-C1'	7.52	126.72	117.70
20	QT	73	HIS	N-CA-CB	7.52	124.13	110.60
1	XA	309	G	OP1-P-OP2	-7.52	108.33	119.60
34	RA	1535	U	C2-N1-C1'	7.51	126.71	117.70
35	YB	31	C	N3-C2-O2	-7.51	116.64	121.90
1	QA	1440(B)	G	C8-N9-C1'	-7.51	117.24	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	1128	C	C2-N1-C1'	7.50	127.06	118.80
1	XA	1233	G	C8-N9-C1'	-7.50	117.25	127.00
1	XA	309	G	C3'-C2'-C1'	-7.50	95.50	101.50
34	YA	1404	C	C2-N1-C1'	7.49	127.04	118.80
1	QA	20	U	C2-N1-C1'	7.49	126.68	117.70
1	QA	1028(D)	G	C4-N9-C1'	7.48	136.22	126.50
1	QA	848	C	C2-N1-C1'	7.48	127.03	118.80
1	QA	1326	C	C2-N1-C1'	7.48	127.03	118.80
1	XA	1062	U	C2-N1-C1'	7.48	126.67	117.70
34	YA	2190	G	C8-N9-C1'	-7.47	117.28	127.00
1	QA	1361	G	C4-N9-C1'	7.47	136.21	126.50
1	XA	1495	U	N1-C1'-C2'	-7.47	103.78	112.00
1	XA	690	G	C4-N9-C1'	7.47	136.21	126.50
1	QA	757	U	C2-N1-C1'	7.46	126.66	117.70
1	QA	1440(B)	G	C4-N9-C1'	7.46	136.20	126.50
1	XA	309	G	P-O5'-C5'	7.46	132.84	120.90
1	QA	24	U	C2-N1-C1'	7.46	126.65	117.70
34	YA	269	U	C2-N1-C1'	7.46	126.65	117.70
1	XA	618	C	O5'-P-OP1	7.45	119.64	110.70
1	XA	1107	C	O5'-P-OP1	7.45	119.64	110.70
1	XA	1145	C	C6-N1-C1'	-7.45	111.87	120.80
35	YB	82	G	C8-N9-C4	-7.45	103.42	106.40
1	QA	581	G	C4-N9-C1'	7.44	136.18	126.50
34	YA	12	U	C6-N1-C1'	-7.44	110.79	121.20
1	XA	1373	G	O5'-P-OP1	7.43	119.62	110.70
34	RA	1672	C	C2-N1-C1'	7.43	126.98	118.80
34	YA	856	C	C6-N1-C1'	-7.43	111.88	120.80
35	YB	10	C	C5-C6-N1	7.43	124.71	121.00
1	QA	1361	G	C8-N9-C1'	-7.43	117.35	127.00
1	XA	1329	A	OP2-P-O3'	7.42	121.53	105.20
1	XA	1226	C	C6-N1-C1'	7.42	129.71	120.80
34	RA	1558	A	P-O3'-C3'	7.41	128.59	119.70
34	YA	2815	C	C6-N1-C1'	-7.41	111.91	120.80
34	RA	1549	C	C2-N1-C1'	7.41	126.94	118.80
34	RA	1870	C	C6-N1-C1'	-7.40	111.92	120.80
1	QA	1028(D)	G	C8-N9-C1'	-7.40	117.38	127.00
1	XA	1314	C	C2-N1-C1'	7.40	126.94	118.80
34	RA	1075	C	C2-N1-C1'	7.40	126.94	118.80
1	XA	68(M)	U	OP1-P-OP2	-7.39	108.51	119.60
1	XA	690	G	C8-N9-C1'	-7.39	117.39	127.00
36	RD	34	VAL	N-CA-C	-7.39	91.04	111.00
1	QA	163	C	C6-N1-C1'	-7.39	111.93	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	YA	1742	C	C6-N1-C1'	-7.38	111.94	120.80
34	RA	602	G	C8-N9-C1'	-7.38	117.41	127.00
34	RA	1577	C	C2-N1-C1'	7.38	126.91	118.80
34	YA	2034	U	C2-N1-C1'	7.38	126.55	117.70
34	RA	1881	C	C6-N1-C1'	-7.37	111.95	120.80
34	YA	1880	C	C6-N1-C1'	-7.37	111.95	120.80
1	QA	1315	U	C2-N1-C1'	7.37	126.54	117.70
34	RA	1905	C	C6-N1-C1'	-7.37	111.96	120.80
1	QA	1056	U	C2-N1-C1'	7.37	126.54	117.70
34	YA	2190	G	C4-N9-C1'	7.37	136.08	126.50
34	RA	1734	C	C2-N1-C1'	7.37	126.90	118.80
35	YB	1	U	N1-C2-O2	7.37	127.95	122.80
35	RB	11	C	N1-C2-O2	7.36	123.32	118.90
1	XA	838(B)	U	C2-N1-C1'	7.36	126.53	117.70
1	XA	726	C	C2-N1-C1'	7.35	126.89	118.80
34	YA	2161	C	C2-N1-C1'	7.35	126.89	118.80
1	QA	581	G	C8-N9-C1'	-7.35	117.45	127.00
34	YA	1005	C	C2-N1-C1'	7.35	126.89	118.80
35	YB	70	C	C5-C6-N1	7.35	124.67	121.00
1	QA	1150	U	C2-N1-C1'	7.34	126.51	117.70
35	YB	37	C	C6-N1-C2	-7.34	117.36	120.30
1	QA	154	C	C2-N1-C1'	7.34	126.87	118.80
1	QA	54	C	C2-N1-C1'	7.34	126.87	118.80
1	QA	443	C	C6-N1-C1'	-7.33	112.01	120.80
1	QA	862	C	C2-N1-C1'	7.33	126.86	118.80
34	RA	1433	U	C2-N1-C1'	7.32	126.48	117.70
34	RA	2374	C	C2-N1-C1'	7.32	126.85	118.80
34	RA	602	G	C4-N9-C1'	7.31	136.01	126.50
1	XA	1367	C	O4'-C1'-N1	7.31	114.05	108.20
34	RA	9	U	C2-N1-C1'	7.31	126.47	117.70
1	XA	115	G	OP2-P-O3'	7.31	121.28	105.20
34	YA	1558	A	P-O3'-C3'	7.31	128.47	119.70
1	QA	322	C	C2-N1-C1'	7.31	126.84	118.80
1	XA	1270	C	C6-N1-C1'	-7.31	112.03	120.80
1	QA	137	C	C2-N1-C1'	7.30	126.83	118.80
1	QA	1113	C	C2-N1-C1'	7.30	126.83	118.80
34	RA	231	C	C2-N1-C1'	7.30	126.83	118.80
1	QA	723	U	C6-N1-C1'	-7.30	110.98	121.20
1	QA	1376	U	C2-N1-C1'	7.30	126.46	117.70
34	RA	1788	C	C2-N1-C1'	7.30	126.83	118.80
1	QA	1230	C	C2-N1-C1'	7.29	126.83	118.80
34	RA	2766	G	C4-N9-C1'	7.29	135.98	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	1301	U	C6-N1-C1'	-7.29	110.99	121.20
34	RA	1391	U	C6-N1-C1'	-7.29	110.99	121.20
1	XA	1113	C	C2-N1-C1'	7.29	126.82	118.80
1	QA	201(A)	C	C6-N1-C1'	-7.29	112.05	120.80
35	YB	94	C	C6-N1-C2	-7.29	117.38	120.30
34	YA	270(Q)	C	C6-N1-C1'	-7.29	112.06	120.80
34	YA	1407	C	C2-N1-C1'	7.29	126.81	118.80
35	YB	102	G	N1-C6-O6	-7.28	115.53	119.90
1	XA	1327	C	OP1-P-O3'	7.28	121.21	105.20
1	XA	368	U	C2-N1-C1'	7.28	126.43	117.70
34	RA	2456	C	C6-N1-C1'	-7.27	112.08	120.80
1	QA	198	G	C4-N9-C1'	7.27	135.94	126.50
34	RA	657	U	C2-N1-C1'	7.26	126.42	117.70
34	RA	1879	C	C6-N1-C1'	-7.26	112.08	120.80
1	XA	443	C	C2-N1-C1'	7.26	126.79	118.80
34	YA	420	C	C6-N1-C1'	-7.26	112.08	120.80
1	QA	198	G	C8-N9-C1'	-7.26	117.56	127.00
34	RA	2821	A	O5'-P-OP1	-7.26	99.16	105.70
1	XA	590	C	C2-N1-C1'	7.26	126.78	118.80
34	YA	2312	U	C6-N1-C1'	-7.25	111.04	121.20
34	RA	384	U	C6-N1-C1'	-7.25	111.05	121.20
35	YB	47	C	N1-C2-O2	7.25	123.25	118.90
34	RA	161	U	C2-N1-C1'	7.25	126.40	117.70
35	RB	22	U	C6-N1-C2	-7.25	116.65	121.00
1	QA	623	C	C6-N1-C1'	-7.24	112.11	120.80
34	RA	2098	U	C6-N1-C1'	-7.24	111.06	121.20
1	QA	1369	C	C6-N1-C1'	-7.24	112.12	120.80
34	YA	721	C	C6-N1-C1'	-7.23	112.12	120.80
34	RA	779	U	C2-N1-C1'	7.23	126.38	117.70
1	XA	201(C)	U	C6-N1-C1'	-7.23	111.08	121.20
34	YA	1533	C	C6-N1-C1'	-7.23	112.12	120.80
1	QA	110	C	C6-N1-C1'	-7.23	112.12	120.80
34	RA	2343	C	C6-N1-C1'	-7.23	112.12	120.80
34	RA	2766	G	C8-N9-C1'	-7.23	117.60	127.00
34	YA	2063	C	C2-N1-C1'	7.23	126.75	118.80
34	YA	541	C	C6-N1-C1'	-7.21	112.15	120.80
1	QA	1097	C	C2-N1-C1'	7.21	126.73	118.80
1	QA	437	U	C2-N1-C1'	7.20	126.34	117.70
1	QA	498	U	C6-N1-C1'	-7.20	111.12	121.20
34	RA	1213	A	C8-N9-C1'	-7.20	114.74	127.70
1	QA	320	C	C6-N1-C1'	-7.19	112.17	120.80
34	RA	1213	A	C4-N9-C1'	7.19	139.24	126.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	1131	G	C4-N9-C1'	7.19	135.85	126.50
1	XA	1343	G	C3'-C2'-C1'	7.19	107.25	101.50
1	XA	1225	A	C5'-C4'-O4'	7.19	117.72	109.10
1	XA	1131	G	C8-N9-C1'	-7.19	117.66	127.00
1	XA	1277	C	P-O3'-C3'	7.18	128.32	119.70
34	YA	2559	C	C2-N1-C1'	7.18	126.70	118.80
34	YA	2858	C	C2-N1-C1'	7.18	126.70	118.80
1	XA	1346	A	C8-N9-C1'	-7.18	114.77	127.70
34	RA	2658	C	C6-N1-C1'	-7.18	112.18	120.80
34	RA	714	U	C6-N1-C1'	-7.18	111.15	121.20
34	RA	1533	C	C6-N1-C1'	-7.18	112.19	120.80
1	QA	79	G	C4-N9-C1'	7.17	135.83	126.50
1	QA	984	C	C6-N1-C1'	-7.17	112.20	120.80
34	RA	453	C	C2-N1-C1'	7.17	126.69	118.80
34	YA	1271	G	C4-N9-C1'	7.17	135.82	126.50
1	XA	1140	C	C6-N1-C1'	-7.17	112.20	120.80
1	QA	1328	C	OP1-P-OP2	-7.16	108.86	119.60
1	XA	946	A	O5'-P-OP1	7.16	119.29	110.70
34	YA	105	C	C2-N1-C1'	7.16	126.68	118.80
1	XA	381	C	C6-N1-C1'	-7.16	112.21	120.80
1	XA	36	C	C6-N1-C1'	-7.16	112.21	120.80
36	YD	35	LYS	CA-CB-CG	7.16	129.14	113.40
1	XA	1361	G	C4-N9-C1'	7.15	135.80	126.50
34	RA	1649	G	C4-N9-C1'	7.15	135.80	126.50
1	QA	283	C	C2-N1-C1'	7.15	126.67	118.80
1	XA	68(V)	U	C6-N1-C1'	-7.15	111.19	121.20
1	XA	955	U	C2-N1-C1'	7.15	126.28	117.70
34	RA	2178	C	C6-N1-C1'	-7.14	112.23	120.80
1	QA	643	C	C2-N1-C1'	7.14	126.66	118.80
1	QA	719	C	C2-N1-C1'	7.14	126.66	118.80
1	QA	223	U	C2-N1-C1'	7.13	126.26	117.70
34	RA	2506	U	C6-N1-C1'	-7.13	111.22	121.20
1	QA	789	U	C2-N1-C1'	7.13	126.25	117.70
1	QA	186(H)	C	C6-N1-C1'	-7.12	112.26	120.80
1	QA	1313	U	C6-N1-C1'	-7.12	111.24	121.20
1	XA	1361	G	C8-N9-C1'	-7.12	117.75	127.00
1	XA	1150	U	C6-N1-C1'	-7.11	111.24	121.20
1	QA	79	G	C8-N9-C1'	-7.11	117.76	127.00
1	XA	946	A	O5'-P-OP2	-7.11	99.30	105.70
34	YA	1271	G	C8-N9-C1'	-7.11	117.76	127.00
34	YA	1022	G	P-O3'-C3'	7.11	128.23	119.70
1	QA	1406	U	C2-N1-C1'	7.10	126.22	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	323	U	C6-N1-C1'	-7.10	111.25	121.20
34	RA	2584	U	O4'-C1'-N1	7.10	113.88	108.20
1	XA	658	G	C4-N9-C1'	7.09	135.72	126.50
34	YA	629	G	C4-N9-C1'	7.09	135.72	126.50
1	QA	826	C	C2-N1-C1'	7.09	126.60	118.80
1	XA	531	U	C2-N1-C1'	7.09	126.21	117.70
34	RA	141(B)	C	C6-N1-C1'	-7.09	112.29	120.80
35	YB	1	U	N3-C2-O2	-7.09	117.24	122.20
34	YA	652	C	C2-N1-C1'	7.08	126.59	118.80
35	YB	80	U	N3-C2-O2	-7.08	117.24	122.20
1	XA	1348	U	O4'-C1'-N1	7.08	113.86	108.20
34	YA	2784	C	C2-N1-C1'	7.08	126.59	118.80
34	RA	2889	C	C2-N1-C1'	7.08	126.58	118.80
34	RA	2695	C	C6-N1-C1'	-7.07	112.31	120.80
34	RA	1178	C	C2-N1-C1'	7.07	126.58	118.80
34	RA	1370	C	C2-N1-C1'	7.07	126.58	118.80
1	QA	748	C	P-O3'-C3'	7.07	128.18	119.70
34	RA	537	C	C2-N1-C1'	7.06	126.57	118.80
34	YA	1314	C	C2-N1-C1'	7.06	126.57	118.80
34	YA	1045	A	P-O3'-C3'	7.06	128.17	119.70
35	YB	27	C	C6-N1-C2	-7.06	117.48	120.30
34	RA	846	C	P-O3'-C3'	7.05	128.16	119.70
1	QA	186(B)	C	C6-N1-C1'	-7.05	112.34	120.80
34	YA	2550	G	N9-C1'-C2'	7.05	123.17	114.00
1	XA	678	U	C2-N1-C1'	7.05	126.16	117.70
1	XA	1426	C	C6-N1-C1'	-7.05	112.34	120.80
34	YA	846	C	P-O3'-C3'	7.05	128.16	119.70
1	XA	658	G	C8-N9-C1'	-7.04	117.85	127.00
1	XA	1325	C	C6-N1-C1'	-7.04	112.35	120.80
34	YA	629	G	C8-N9-C1'	-7.03	117.86	127.00
35	YB	30	C	N3-C2-O2	-7.03	116.98	121.90
1	QA	312	C	C2-N1-C1'	7.03	126.53	118.80
1	XA	309	G	C5'-C4'-C3'	7.03	127.25	116.00
34	YA	1070	A	C4-N9-C1'	7.03	138.95	126.30
1	QA	634	C	C2-N1-C1'	7.03	126.53	118.80
1	XA	1345	U	C6-N1-C1'	-7.03	111.36	121.20
35	YB	68	C	C6-N1-C2	-7.03	117.49	120.30
34	YA	2380	C	C2-N1-C1'	7.02	126.52	118.80
1	XA	811	C	C2-N1-C1'	7.02	126.52	118.80
1	QA	618	C	C2-N1-C1'	7.02	126.52	118.80
34	YA	2584	U	O4'-C1'-N1	7.02	113.81	108.20
35	RB	27	C	C6-N1-C2	-7.01	117.49	120.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	1384	C	C2-N1-C1'	7.01	126.52	118.80
34	RA	544	C	C6-N1-C1'	-7.01	112.39	120.80
1	QA	1086	U	C2-N1-C1'	7.01	126.11	117.70
34	RA	2698	U	C2-N1-C1'	7.01	126.11	117.70
1	XA	1358	U	OP1-P-O3'	7.01	120.62	105.20
34	RA	1141	U	C2-N1-C1'	7.01	126.11	117.70
35	RB	11	C	N3-C2-O2	-7.01	117.00	121.90
34	YA	1741	C	C2-N1-C1'	7.00	126.50	118.80
35	YB	54	G	C4-C5-N7	7.00	113.60	110.80
1	XA	1232	U	OP1-P-OP2	-7.00	109.10	119.60
1	XA	322	C	C6-N1-C1'	-7.00	112.40	120.80
34	RA	1790	C	C6-N1-C1'	-6.99	112.41	120.80
34	RA	184	C	C2-N1-C1'	6.99	126.49	118.80
34	RA	2321	G	C4-N9-C1'	6.99	135.59	126.50
34	YA	2723	C	C2-N1-C1'	6.99	126.49	118.80
35	YB	31	C	N1-C2-O2	6.98	123.09	118.90
34	YA	613	U	C2-N1-C1'	6.98	126.07	117.70
34	YA	1728	G	O4'-C1'-N9	6.98	113.78	108.20
1	QA	1128	C	C6-N1-C1'	-6.97	112.43	120.80
34	YA	1433	U	C2-N1-C1'	6.97	126.07	117.70
34	YA	271(C)	G	P-O3'-C3'	6.97	128.06	119.70
1	QA	195	A	C4-N9-C1'	6.96	138.84	126.30
1	QA	1219	U	C6-N1-C1'	-6.96	111.45	121.20
1	XA	1037	C	C2-N1-C1'	6.96	126.46	118.80
1	XA	131	C	C2-N1-C1'	6.96	126.45	118.80
34	YA	1070	A	C8-N9-C1'	-6.95	115.19	127.70
1	XA	1346	A	C4-N9-C1'	6.95	138.81	126.30
1	XA	1366	C	C2-N1-C1'	6.95	126.44	118.80
34	RA	1022	G	P-O3'-C3'	6.95	128.03	119.70
34	RA	2554	U	C6-N1-C1'	-6.95	111.48	121.20
34	YA	503	A	P-O3'-C3'	6.94	128.03	119.70
34	RA	721	C	C6-N1-C1'	-6.94	112.47	120.80
20	QT	73	HIS	CB-CA-C	-6.94	96.52	110.40
34	YA	1956	U	C2-N1-C1'	6.94	126.02	117.70
34	RA	1330	C	C2-N1-C1'	6.93	126.42	118.80
1	XA	115	G	O4'-C1'-N9	6.93	113.75	108.20
34	RA	2483	C	C2-N1-C1'	6.93	126.42	118.80
1	XA	323	U	O5'-P-OP1	-6.93	99.47	105.70
1	XA	619	U	C2-N1-C1'	-6.93	109.39	117.70
1	XA	879	C	C2-N1-C1'	6.93	126.42	118.80
1	XA	310	G	C8-N9-C1'	6.92	136.00	127.00
34	YA	1097	U	C2-N1-C1'	6.92	126.01	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	YA	1990	C	C6-N1-C1'	-6.92	112.49	120.80
1	QA	989	C	C6-N1-C1'	-6.92	112.49	120.80
34	RA	2321	G	C8-N9-C1'	-6.92	118.01	127.00
1	QA	590	C	C2-N1-C1'	6.92	126.41	118.80
34	YA	414	C	C6-N1-C1'	-6.90	112.52	120.80
34	RA	2666	C	C6-N1-C1'	-6.90	112.52	120.80
35	YB	54	G	C5-N7-C8	-6.90	100.85	104.30
1	QA	1371	G	C4-N9-C1'	6.89	135.46	126.50
34	RA	1225	C	C6-N1-C1'	-6.89	112.53	120.80
1	XA	1242	C	C2-N1-C1'	6.89	126.38	118.80
34	RA	413	C	C6-N1-C1'	-6.89	112.53	120.80
34	RA	2471	C	C2-N1-C1'	6.89	126.38	118.80
41	RI	131	LYS	N-CA-C	6.89	129.60	111.00
34	RA	1577	C	C6-N1-C1'	-6.89	112.53	120.80
34	YA	2465	C	C2-N1-C1'	6.89	126.38	118.80
1	XA	931	C	C2-N1-C1'	6.89	126.38	118.80
34	YA	1788	C	C2-N1-C1'	6.88	126.37	118.80
1	QA	195	A	C8-N9-C1'	-6.87	115.33	127.70
1	QA	848	C	C6-N1-C1'	-6.87	112.55	120.80
34	RA	2075	U	C2-N1-C1'	6.87	125.95	117.70
35	YB	1	U	C5-C6-N1	6.87	126.14	122.70
34	RA	2703	C	C2-N1-C1'	6.87	126.36	118.80
34	RA	1675	C	C2-N1-C1'	6.87	126.35	118.80
34	YA	904	C	C2-N1-C1'	6.87	126.35	118.80
34	RA	2871	C	C2-N1-C1'	6.86	126.34	118.80
1	XA	948	C	C5'-C4'-O4'	-6.86	100.87	109.10
1	QA	952	U	C2-N1-C1'	6.85	125.92	117.70
1	QA	328	C	P-O3'-C3'	6.85	127.92	119.70
1	XA	186(O)	U	C2-N1-C1'	6.85	125.92	117.70
1	QA	1259	C	C2-N1-C1'	6.84	126.33	118.80
34	YA	1506	C	C2-N1-C1'	6.84	126.33	118.80
34	RA	2403	C	C2-N1-C1'	6.84	126.32	118.80
1	XA	525	C	C2-N1-C1'	6.84	126.32	118.80
34	YA	404	C	P-O3'-C3'	6.84	127.91	119.70
34	YA	859	G	P-O3'-C3'	6.84	127.91	119.70
1	QA	1326	C	C6-N1-C1'	-6.84	112.59	120.80
1	XA	310	G	C4-N9-C1'	-6.84	117.61	126.50
34	YA	270(Z)	G	C4-N9-C1'	6.84	135.39	126.50
34	YA	2343	C	C2-N1-C1'	6.83	126.32	118.80
34	YA	1653	G	P-O3'-C3'	6.83	127.90	119.70
34	YA	2321	G	C4-N9-C1'	6.83	135.38	126.50
1	QA	1371	G	C8-N9-C1'	-6.83	118.12	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	1307	U	OP1-P-OP2	6.83	129.84	119.60
1	QA	1263	C	C2-N1-C1'	6.83	126.31	118.80
34	YA	537	C	C2-N1-C1'	6.83	126.31	118.80
34	RA	1066	U	C2-N1-C1'	6.82	125.88	117.70
34	RA	1404	C	C2-N1-C1'	6.82	126.30	118.80
35	RB	71	C	N1-C2-O2	6.82	122.99	118.90
34	YA	591	C	C2-N1-C1'	6.82	126.30	118.80
1	QA	620	C	C2-N1-C1'	6.82	126.30	118.80
34	RA	642	G	C8-N9-C1'	-6.81	118.14	127.00
34	YA	1372	U	C2-N1-C1'	6.81	125.87	117.70
34	YA	384	U	C2-N1-C1'	6.80	125.86	117.70
34	YA	1404	C	C6-N1-C1'	-6.80	112.64	120.80
34	RA	404	C	P-O3'-C3'	6.80	127.86	119.70
34	RA	1130	U	P-O3'-C3'	6.80	127.86	119.70
34	RA	1686	C	C2-N1-C1'	6.80	126.28	118.80
34	RA	1777	U	C2-N1-C1'	6.80	125.86	117.70
34	RA	1516	U	C2-N1-C1'	6.80	125.86	117.70
1	QA	595	G	C4-N9-C1'	6.80	135.33	126.50
1	QA	1242	C	C2-N1-C1'	6.80	126.28	118.80
34	RA	343	C	C2-N1-C1'	6.79	126.28	118.80
34	RA	642	G	C4-N9-C1'	6.79	135.33	126.50
1	XA	1128	C	C2-N1-C1'	6.79	126.27	118.80
1	XA	943	U	C6-N1-C1'	-6.79	111.70	121.20
34	YA	1870	C	C2-N1-C1'	6.79	126.27	118.80
34	YA	2895	U	C2-N1-C1'	6.79	125.84	117.70
34	RA	1105	U	C2-N1-C1'	6.78	125.84	117.70
34	RA	1672	C	C6-N1-C1'	-6.78	112.66	120.80
34	YA	270(Z)	G	C8-N9-C1'	-6.78	118.18	127.00
34	YA	2554	U	C2-N1-C1'	6.78	125.83	117.70
34	YA	1687	G	C4-N9-C1'	6.77	135.31	126.50
22	XV	56	C	N1-C2-O2	6.77	122.96	118.90
1	QA	344	A	O4'-C1'-N9	-6.77	102.79	108.20
34	YA	1914	C	O4'-C1'-N1	6.77	113.61	108.20
34	RA	1734	C	C6-N1-C1'	-6.76	112.68	120.80
1	XA	948	C	O5'-P-OP1	-6.76	99.61	105.70
1	QA	137	C	C6-N1-C1'	-6.76	112.69	120.80
1	XA	1314	C	C6-N1-C1'	-6.76	112.69	120.80
1	XA	757	U	C2-N1-C1'	6.75	125.81	117.70
42	YN	114	ARG	N-CA-C	-6.75	92.77	111.00
1	QA	1482	G	C4-N9-C1'	6.75	135.28	126.50
1	QA	19	C	C2-N1-C1'	6.74	126.22	118.80
1	XA	1358	U	P-O5'-C5'	6.73	131.67	120.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	RA	1075	C	C6-N1-C1'	-6.73	112.72	120.80
1	QA	595	G	C8-N9-C1'	-6.73	118.25	127.00
34	RA	1549	C	C6-N1-C1'	-6.73	112.72	120.80
1	QA	862	C	C6-N1-C1'	-6.73	112.73	120.80
34	YA	2321	G	C8-N9-C1'	-6.73	118.26	127.00
1	QA	1140	C	C2-N1-C1'	6.72	126.20	118.80
1	QA	1366	C	C2-N1-C1'	6.72	126.19	118.80
34	RA	568	U	C2-N1-C1'	6.72	125.76	117.70
34	RA	2291	U	C6-N1-C1'	-6.71	111.80	121.20
1	QA	580	U	C2-N1-C1'	6.71	125.76	117.70
1	XA	1241	G	C4-N9-C1'	6.71	135.23	126.50
34	YA	1005	C	C6-N1-C1'	-6.71	112.75	120.80
1	XA	1313	U	C2-N1-C1'	6.71	125.75	117.70
34	RA	2752	C	C2-N1-C1'	6.71	126.18	118.80
34	YA	1951	U	C6-N1-C1'	-6.71	111.81	121.20
1	XA	1225	A	C8-N9-C1'	-6.71	115.63	127.70
34	RA	1188	U	C2-N1-C1'	6.70	125.74	117.70
1	XA	714	G	C4-N9-C1'	6.70	135.22	126.50
35	RB	22	U	C2-N1-C1'	6.70	125.74	117.70
34	RA	420	C	C2-N1-C1'	6.70	126.17	118.80
1	XA	714	G	C8-N9-C1'	-6.70	118.30	127.00
34	YA	2374	C	C2-N1-C1'	6.70	126.16	118.80
34	YA	1391	U	C2-N1-C1'	6.69	125.73	117.70
1	XA	1537	U	P-O3'-C3'	6.69	127.73	119.70
1	QA	578	C	C2-N1-C1'	6.69	126.16	118.80
34	RA	1653	G	P-O3'-C3'	6.69	127.72	119.70
20	QT	74	LYS	CA-C-O	6.68	134.14	120.10
34	YA	464	U	C2-N1-C1'	6.68	125.72	117.70
1	XA	944	G	C4-N9-C1'	6.68	135.19	126.50
34	YA	1687	G	C8-N9-C1'	-6.68	118.31	127.00
1	XA	590	C	C6-N1-C1'	-6.68	112.79	120.80
34	YA	2173	A	O5'-P-OP1	6.68	118.71	110.70
40	YH	152	ARG	C-N-CA	6.68	138.39	121.70
1	QA	1157	A	O4'-C1'-N9	6.67	113.54	108.20
34	RA	2667	C	C2-N1-C1'	6.67	126.14	118.80
34	RA	1788	C	C6-N1-C1'	-6.67	112.79	120.80
34	YA	2681	C	P-O3'-C3'	6.67	127.71	119.70
1	XA	726	C	C6-N1-C1'	-6.67	112.80	120.80
1	QA	154	C	C6-N1-C1'	-6.66	112.80	120.80
1	QA	992	U	P-O3'-C3'	6.66	127.70	119.70
1	QA	1482	G	C8-N9-C1'	-6.66	118.34	127.00
34	RA	2374	C	C6-N1-C1'	-6.66	112.80	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	343	U	C2-N1-C1'	6.66	125.70	117.70
34	RA	1407	C	C2-N1-C1'	6.66	126.13	118.80
34	RA	231	C	C6-N1-C1'	-6.66	112.81	120.80
34	RA	1240	U	C2-N1-C1'	6.66	125.69	117.70
34	YA	1754	C	C2-N1-C1'	6.65	126.11	118.80
34	YA	2161	C	C6-N1-C1'	-6.65	112.82	120.80
1	QA	1351	U	C2-N1-C1'	6.65	125.68	117.70
1	XA	62	U	C2-N1-C1'	6.65	125.68	117.70
1	QA	54	C	C6-N1-C1'	-6.65	112.82	120.80
1	QA	322	C	C6-N1-C1'	-6.65	112.82	120.80
34	YA	1407	C	C6-N1-C1'	-6.64	112.83	120.80
1	QA	122	G	C4-N9-C1'	6.64	135.13	126.50
1	XA	307	C	C2-N1-C1'	6.63	126.10	118.80
1	XA	1113	C	C6-N1-C1'	-6.63	112.84	120.80
1	XA	1241	G	C8-N9-C1'	-6.63	118.38	127.00
35	RB	27	C	N1-C2-O2	6.63	122.88	118.90
1	QA	36	C	C2-N1-C1'	6.63	126.09	118.80
1	QA	355	C	C2-N1-C1'	6.63	126.09	118.80
1	QA	1113	C	C6-N1-C1'	-6.63	112.85	120.80
1	QA	1230	C	C6-N1-C1'	-6.62	112.85	120.80
1	XA	718	G	C8-N9-C1'	-6.62	118.39	127.00
34	YA	2656	U	C6-N1-C1'	-6.62	111.93	121.20
34	RA	1046	A	C4-N9-C1'	6.62	138.21	126.30
1	QA	1129	C	C2-N1-C1'	6.62	126.08	118.80
34	RA	229	A	P-O3'-C3'	6.62	127.64	119.70
34	RA	1611	C	C2-N1-C1'	6.61	126.08	118.80
34	YA	897	C	C2-N1-C1'	6.61	126.07	118.80
1	XA	443	C	C6-N1-C1'	-6.61	112.86	120.80
19	QS	41	VAL	N-CA-C	6.61	128.85	111.00
1	QA	1206	G	C4-N9-C1'	6.60	135.09	126.50
34	RA	503	A	P-O3'-C3'	6.60	127.62	119.70
34	RA	636	G	C8-N9-C1'	-6.60	118.42	127.00
1	XA	944	G	C8-N9-C1'	-6.60	118.42	127.00
34	RA	2871	C	C6-N1-C1'	-6.60	112.88	120.80
34	YA	1774	C	C2-N1-C1'	6.60	126.06	118.80
1	QA	838(B)	U	C6-N1-C1'	-6.59	111.97	121.20
34	YA	1857	G	C4-N9-C1'	6.59	135.07	126.50
34	YA	2559	C	C6-N1-C1'	-6.59	112.89	120.80
34	YA	2612	C	O5'-P-OP2	-6.59	99.77	105.70
34	YA	2836	U	C6-N1-C1'	-6.59	111.98	121.20
1	QA	343	U	C2-N1-C1'	6.58	125.60	117.70
1	XA	277	C	C2-N1-C1'	6.58	126.04	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	617	G	O4'-C1'-C2'	-6.58	99.22	105.80
1	QA	1097	C	C6-N1-C1'	-6.58	112.91	120.80
34	RA	1920	C	C2-N1-C1'	6.58	126.03	118.80
34	YA	2063	C	C6-N1-C1'	-6.58	112.91	120.80
34	RA	271(C)	G	P-O3'-C3'	6.57	127.59	119.70
34	RA	1045	A	P-O3'-C3'	6.57	127.59	119.70
34	RA	2040	C	C2-N1-C1'	6.57	126.03	118.80
1	QA	122	G	C8-N9-C1'	-6.57	118.46	127.00
35	YB	54	G	C2-N3-C4	-6.57	108.62	111.90
34	YA	1049	C	C2-N1-C1'	6.57	126.02	118.80
1	QA	1206	G	C8-N9-C1'	-6.56	118.48	127.00
34	RA	2636	U	C2-N1-C1'	6.56	125.57	117.70
14	XN	28	GLY	N-CA-C	6.55	129.49	113.10
35	YB	71	C	C6-N1-C2	-6.55	117.68	120.30
34	RA	2130	U	C6-N1-C1'	-6.55	112.03	121.20
34	YA	637	A	P-O3'-C3'	6.55	127.56	119.70
34	YA	1857	G	C8-N9-C1'	-6.55	118.48	127.00
1	QA	56	U	C2-N1-C1'	6.55	125.56	117.70
1	QA	957	U	C6-N1-C1'	-6.55	112.03	121.20
1	XA	578	C	C2-N1-C1'	6.55	126.00	118.80
22	XV	76	A	C8-N9-C4	-6.55	103.18	105.80
34	YA	114	U	C6-N1-C1'	-6.55	112.03	121.20
35	YB	27	C	N3-C2-O2	-6.55	117.32	121.90
1	XA	718	G	C4-N9-C1'	6.54	135.01	126.50
34	YA	1881	C	C2-N1-C1'	6.54	126.00	118.80
20	QT	74	LYS	CA-C-N	-6.54	102.81	117.20
1	XA	12	U	C2-N1-C1'	6.54	125.55	117.70
34	YA	2710	C	C2-N1-C1'	6.54	125.99	118.80
1	QA	390	C	C2-N1-C1'	6.54	125.99	118.80
35	YB	63	G	N1-C6-O6	-6.54	115.98	119.90
34	RA	1662	C	C2-N1-C1'	6.53	125.99	118.80
34	YA	2858	C	C6-N1-C1'	-6.53	112.96	120.80
34	RA	541	C	C2-N1-C1'	6.53	125.98	118.80
1	XA	1049	U	C6-N1-C1'	-6.53	112.06	121.20
1	QA	283	C	C6-N1-C1'	-6.53	112.97	120.80
1	QA	1148	U	C2-N1-C1'	6.53	125.53	117.70
34	YA	2617	C	C2-N1-C1'	6.53	125.98	118.80
1	XA	964	A	C8-N9-C1'	-6.52	115.96	127.70
34	YA	512	G	O4'-C1'-N9	6.52	113.42	108.20
34	YA	1982	C	C2-N1-C1'	6.52	125.97	118.80
1	QA	643	C	C6-N1-C1'	-6.52	112.97	120.80
34	RA	453	C	C6-N1-C1'	-6.52	112.97	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	RA	856	C	C2-N1-C1'	6.52	125.97	118.80
34	RA	2824	C	C2-N1-C1'	6.52	125.97	118.80
1	QA	521	G	C4-N9-C1'	6.52	134.97	126.50
1	XA	266	G	O4'-C1'-N9	-6.51	102.99	108.20
34	RA	1046	A	C8-N9-C1'	-6.51	115.98	127.70
1	QA	719	C	C6-N1-C1'	-6.50	112.99	120.80
1	XA	636	U	O5'-P-OP1	-6.50	99.84	105.70
1	XA	1074	G	C4-N9-C1'	6.50	134.95	126.50
1	QA	822	C	C2-N1-C1'	6.50	125.95	118.80
34	RA	2342	C	C2-N1-C1'	6.50	125.95	118.80
34	YA	2784	C	C6-N1-C1'	-6.50	113.00	120.80
34	RA	363(F)	U	C2-N1-C1'	6.50	125.50	117.70
34	RA	2656	U	C6-N1-C1'	-6.50	112.11	121.20
1	XA	241	C	C2-N1-C1'	6.50	125.95	118.80
1	QA	136	C	C2-N1-C1'	6.49	125.94	118.80
34	YA	105	C	C6-N1-C1'	-6.49	113.01	120.80
34	RA	636	G	C4-N9-C1'	6.49	134.93	126.50
34	YA	1026	U	P-O3'-C3'	6.49	127.48	119.70
34	YA	1686	C	C2-N1-C1'	6.49	125.93	118.80
34	RA	637	A	P-O3'-C3'	6.48	127.48	119.70
34	RA	2096	U	C2-N1-C1'	6.48	125.48	117.70
1	XA	748	C	P-O3'-C3'	6.48	127.48	119.70
35	YB	60	C	C5-C6-N1	6.48	124.24	121.00
1	XA	131	C	C6-N1-C1'	-6.48	113.02	120.80
1	QA	826	C	C6-N1-C1'	-6.48	113.03	120.80
1	XA	201(A)	C	C2-N1-C1'	6.48	125.93	118.80
34	RA	537	C	C6-N1-C1'	-6.47	113.03	120.80
34	RA	2043	C	C2-N1-C1'	6.47	125.92	118.80
1	QA	705	U	C6-N1-C1'	-6.47	112.14	121.20
34	YA	652	C	C6-N1-C1'	-6.47	113.04	120.80
34	RA	2889	C	C6-N1-C1'	-6.47	113.04	120.80
34	YA	2431	U	C6-N1-C1'	-6.47	112.15	121.20
1	QA	312	C	C6-N1-C1'	-6.46	113.04	120.80
1	QA	1017	G	C8-N9-C1'	-6.46	118.60	127.00
34	RA	270(Q)	C	C2-N1-C1'	6.46	125.90	118.80
1	QA	1127	G	C4-N9-C1'	6.45	134.89	126.50
34	RA	114	U	C6-N1-C1'	-6.45	112.16	121.20
1	QA	1327	C	O4'-C1'-N1	6.45	113.36	108.20
1	XA	20	U	C2-N1-C1'	6.45	125.44	117.70
1	QA	6	G	C4-N9-C1'	6.45	134.88	126.50
1	QA	775	G	C4-N9-C1'	6.45	134.88	126.50
34	RA	2095	C	C2-N1-C1'	6.45	125.89	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	RA	2262	U	C6-N1-C1'	-6.45	112.17	121.20
34	YA	745	G	P-O3'-C3'	-6.45	111.97	119.70
34	YA	753	C	C2-N1-C1'	6.44	125.89	118.80
54	RZ	62	PRO	C-N-CA	6.44	137.81	121.70
1	QA	775	G	C8-N9-C1'	-6.44	118.63	127.00
34	YA	1415	U	C2-N1-C1'	6.44	125.43	117.70
34	RA	1178	C	C6-N1-C1'	-6.44	113.07	120.80
35	YB	27	C	N1-C2-O2	6.44	122.76	118.90
34	RA	753	C	C2-N1-C1'	6.44	125.88	118.80
34	RA	1728	G	O4'-C1'-N9	6.44	113.35	108.20
34	YA	2132	U	C6-N1-C1'	-6.44	112.19	121.20
1	XA	834	C	C2-N1-C1'	6.44	125.88	118.80
1	XA	1384	C	C6-N1-C1'	-6.43	113.08	120.80
34	RA	1370	C	C6-N1-C1'	-6.43	113.08	120.80
1	QA	521	G	C8-N9-C1'	-6.43	118.64	127.00
1	XA	1074	G	C8-N9-C1'	-6.43	118.65	127.00
34	YA	565	C	C2-N1-C1'	6.42	125.87	118.80
34	YA	2494	G	O5'-P-OP1	-6.42	99.92	105.70
34	RA	669	G	C4-N9-C1'	6.42	134.85	126.50
1	QA	692	U	C6-N1-C1'	-6.42	112.21	121.20
22	QV	30	C	C2-N1-C1'	6.42	125.86	118.80
1	XA	796	C	C2-N1-C1'	6.42	125.86	118.80
34	YA	974(B)	C	C2-N1-C1'	6.42	125.86	118.80
34	RA	639	U	C2-N1-C1'	6.42	125.40	117.70
34	RA	1892	C	C2-N1-C1'	6.42	125.86	118.80
42	RN	114	ARG	N-CA-C	-6.42	93.68	111.00
30	R6	13	CYS	CA-CB-SG	-6.41	102.46	114.00
34	RA	613	U	C6-N1-C1'	-6.41	112.23	121.20
34	RA	1528	A	O4'-C1'-N9	6.41	113.33	108.20
34	YA	2097	C	C2-N1-C1'	6.41	125.85	118.80
34	YA	2393	A	OP1-P-O3'	6.41	119.30	105.20
1	QA	6	G	C8-N9-C1'	-6.41	118.67	127.00
1	QA	1017	G	C4-N9-C1'	6.41	134.83	126.50
34	RA	184	C	C6-N1-C1'	-6.41	113.11	120.80
1	QA	687	A	P-O3'-C3'	6.41	127.39	119.70
1	XA	972	C	N1-C1'-C2'	6.41	122.33	114.00
1	XA	1349	A	C8-N9-C1'	6.40	139.22	127.70
34	YA	1314	C	C6-N1-C1'	-6.40	113.12	120.80
34	YA	1640	C	C2-N1-C1'	6.40	125.84	118.80
34	YA	2089	U	C2-N1-C1'	6.39	125.37	117.70
1	XA	68(R)	U	C6-N1-C1'	-6.39	112.26	121.20
1	XA	232	G	C4-N9-C1'	6.39	134.80	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	YA	2040	C	C2-N1-C1'	6.39	125.82	118.80
34	RA	1774	C	C2-N1-C1'	6.38	125.82	118.80
1	QA	1045	C	C2-N1-C1'	6.38	125.82	118.80
53	RY	99	CYS	CA-CB-SG	6.38	125.48	114.00
1	QA	1440(H)	C	C2-N1-C1'	6.38	125.82	118.80
1	QA	1538	C	P-O3'-C3'	6.38	127.35	119.70
1	XA	598	U	C2-N1-C1'	6.38	125.35	117.70
34	YA	650	C	C2-N1-C1'	6.38	125.81	118.80
34	YA	2130	U	C6-N1-C1'	-6.38	112.28	121.20
1	QA	672	U	C2-N1-C1'	6.37	125.34	117.70
34	RA	1509	C	C2-N1-C1'	6.37	125.81	118.80
1	XA	811	C	C6-N1-C1'	-6.37	113.15	120.80
34	RA	266	G	C4-N9-C1'	6.37	134.78	126.50
1	QA	1075	C	C2-N1-C1'	6.37	125.81	118.80
1	QA	1127	G	C8-N9-C1'	-6.37	118.72	127.00
34	YA	2380	C	C6-N1-C1'	-6.37	113.16	120.80
1	QA	618	C	C6-N1-C1'	-6.37	113.16	120.80
22	XV	15	G	C8-N9-C4	-6.37	103.85	106.40
34	YA	35	G	C4-N9-C1'	6.36	134.77	126.50
35	YB	38	C	C6-N1-C2	-6.36	117.76	120.30
1	XA	1366	C	C6-N1-C1'	-6.36	113.17	120.80
1	QA	1384	C	C2-N1-C1'	6.36	125.79	118.80
34	RA	669	G	C8-N9-C1'	-6.36	118.74	127.00
34	RA	2028	U	C2-N1-C1'	6.35	125.33	117.70
1	XA	1465	C	C2-N1-C1'	6.35	125.79	118.80
1	QA	634	C	C6-N1-C1'	-6.35	113.18	120.80
34	YA	1741	C	C6-N1-C1'	-6.35	113.18	120.80
1	QA	24	U	C6-N1-C1'	-6.35	112.31	121.20
34	YA	1598	C	C2-N1-C1'	6.35	125.79	118.80
34	RA	266	G	C8-N9-C1'	-6.35	118.75	127.00
34	YA	2723	C	C6-N1-C1'	-6.35	113.18	120.80
1	XA	972	C	O4'-C1'-N1	6.35	113.28	108.20
1	XA	989	C	C2-N1-C1'	6.35	125.78	118.80
35	YB	86	G	C8-N9-C4	6.35	108.94	106.40
1	XA	1062	U	C6-N1-C1'	-6.35	112.32	121.20
1	XA	322	C	P-O3'-C3'	6.34	127.31	119.70
1	XA	805	C	C2-N1-C1'	6.34	125.78	118.80
1	XA	1377	A	P-O3'-C3'	6.34	127.31	119.70
34	YA	2195	C	C2-N1-C1'	6.34	125.78	118.80
34	YA	2658	C	C2-N1-C1'	6.34	125.78	118.80
1	XA	283	C	C2-N1-C1'	6.34	125.78	118.80
34	YA	2146	C	C2-N1-C1'	6.34	125.78	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	YA	2466	C	C2-N1-C1'	6.34	125.78	118.80
34	RA	2179	C	C2-N1-C1'	6.34	125.77	118.80
34	RA	2483	C	C6-N1-C1'	-6.34	113.20	120.80
1	QA	20	U	C6-N1-C1'	-6.33	112.33	121.20
1	XA	618	C	O5'-P-OP2	-6.33	100.00	105.70
1	XA	1219	U	C2-N1-C1'	-6.33	110.10	117.70
34	YA	242	G	P-O3'-C3'	6.33	127.30	119.70
35	YB	28	C	C6-N1-C2	-6.33	117.77	120.30
34	RA	2582	G	C4-N9-C1'	6.33	134.73	126.50
34	RA	1330	C	C6-N1-C1'	-6.33	113.20	120.80
1	XA	1349	A	C4-N9-C1'	-6.33	114.91	126.30
34	YA	2343	C	C6-N1-C1'	-6.33	113.20	120.80
34	RA	339	U	C2-N1-C1'	6.33	125.29	117.70
34	RA	18	C	C2-N1-C1'	6.33	125.76	118.80
1	QA	757	U	C6-N1-C1'	-6.32	112.35	121.20
34	RA	2403	C	C6-N1-C1'	-6.32	113.21	120.80
34	RA	242	G	P-O3'-C3'	6.32	127.29	119.70
34	YA	269	U	C6-N1-C1'	-6.32	112.35	121.20
34	YA	1914	C	C2-N1-C1'	6.32	125.75	118.80
1	QA	955	U	C2-N1-C1'	6.32	125.28	117.70
34	RA	1535	U	C6-N1-C1'	-6.32	112.35	121.20
1	QA	1403	C	C2-N1-C1'	6.32	125.75	118.80
34	RA	1857	G	C4-N9-C1'	6.32	134.71	126.50
34	RA	2712(A)	U	P-O3'-C3'	6.32	127.28	119.70
34	RA	1154	G	C4-N9-C1'	6.32	134.71	126.50
1	XA	232	G	C8-N9-C1'	-6.32	118.79	127.00
34	YA	76	C	C2-N1-C1'	6.31	125.74	118.80
34	RA	1951	U	C2-N1-C1'	6.31	125.27	117.70
1	XA	879	C	C6-N1-C1'	-6.31	113.23	120.80
1	XA	595	G	C4-N9-C1'	6.31	134.70	126.50
34	RA	1990	C	C2-N1-C1'	6.31	125.74	118.80
34	RA	683	C	C2-N1-C1'	6.30	125.73	118.80
40	YH	82	GLY	N-CA-C	6.30	128.86	113.10
42	YN	48	MET	CG-SD-CE	-6.30	90.11	100.20
1	QA	1382	C	C2-N1-C1'	6.30	125.73	118.80
34	YA	229	A	P-O3'-C3'	6.30	127.26	119.70
1	QA	1150	U	C6-N1-C1'	-6.30	112.38	121.20
12	QL	104	VAL	C-N-CA	6.30	137.45	121.70
1	XA	1505	G	O5'-P-OP1	-6.30	100.03	105.70
1	QA	117	G	C4-N9-C1'	6.30	134.69	126.50
34	RA	1510	A	C4-N9-C1'	6.30	137.63	126.30
1	XA	656	C	C2-N1-C1'	6.30	125.73	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	1201	A	O4'-C1'-N9	-6.29	103.17	108.20
34	YA	2126	A	P-O3'-C3'	6.29	127.25	119.70
1	QA	1315	U	C6-N1-C1'	-6.29	112.39	121.20
34	YA	2145	C	C2-N1-C1'	6.29	125.72	118.80
1	XA	501	C	C2-N1-C1'	6.29	125.72	118.80
1	XA	931	C	C6-N1-C1'	-6.29	113.25	120.80
34	RA	2471	C	C6-N1-C1'	-6.29	113.25	120.80
34	YA	35	G	C8-N9-C1'	-6.29	118.83	127.00
35	YB	30	C	C2-N1-C1'	6.29	125.72	118.80
1	XA	950	U	C2-N1-C1'	6.29	125.24	117.70
1	QA	590	C	C6-N1-C1'	-6.28	113.26	120.80
1	XA	1504	G	P-O3'-C3'	6.28	127.24	119.70
34	YA	904	C	C6-N1-C1'	-6.28	113.26	120.80
1	QA	570	G	C4-N9-C1'	6.28	134.67	126.50
34	RA	687	C	C2-N1-C1'	6.28	125.71	118.80
34	RA	1049	C	C2-N1-C1'	6.28	125.71	118.80
34	YA	2034	U	C6-N1-C1'	-6.28	112.41	121.20
1	XA	1440(I)	U	C2-N1-C1'	6.28	125.23	117.70
1	XA	1297	C	N1-C1'-C2'	6.28	122.16	114.00
1	XA	1499	A	C4-N9-C1'	6.27	137.59	126.30
1	QA	1242	C	C6-N1-C1'	-6.27	113.28	120.80
34	YA	1799	G	P-O3'-C3'	6.27	127.22	119.70
1	QA	381	C	C2-N1-C1'	6.27	125.70	118.80
1	QA	1056	U	C6-N1-C1'	-6.27	112.43	121.20
34	RA	2582	G	C8-N9-C1'	-6.27	118.85	127.00
34	YA	2465	C	C6-N1-C1'	-6.27	113.28	120.80
1	QA	981	U	C2-N1-C1'	6.26	125.22	117.70
34	RA	1327	C	C2-N1-C1'	6.26	125.69	118.80
34	RA	9	U	C6-N1-C1'	-6.26	112.43	121.20
34	YA	343	C	C2-N1-C1'	6.26	125.69	118.80
34	RA	2703	C	C6-N1-C1'	-6.26	113.29	120.80
1	XA	920	U	C2-N1-C1'	6.26	125.21	117.70
34	YA	1788	C	C6-N1-C1'	-6.26	113.29	120.80
34	RA	2379	G	C8-N9-C1'	-6.26	118.86	127.00
1	QA	249	U	C2-N1-C1'	6.26	125.21	117.70
34	RA	427	U	C2-N1-C1'	6.26	125.21	117.70
10	XJ	39	PRO	CA-N-CD	6.26	120.46	111.70
34	RA	1857	G	C8-N9-C1'	-6.25	118.87	127.00
1	XA	525	C	C6-N1-C1'	-6.25	113.30	120.80
1	XA	838(B)	U	C6-N1-C1'	-6.25	112.45	121.20
1	QA	1129	C	C6-N1-C1'	-6.25	113.30	120.80
1	QA	963	G	C4-N9-C1'	6.25	134.62	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	RB	11	C	C2-N1-C1'	6.25	125.67	118.80
34	YA	2891	G	C4-N9-C1'	6.25	134.62	126.50
34	RA	2379	G	C4-N9-C1'	6.25	134.62	126.50
1	XA	608	A	C5'-C4'-C3'	6.25	125.99	116.00
34	RA	222	A	P-O3'-C3'	6.24	127.19	119.70
34	RA	2174	C	C2-N1-C1'	6.24	125.67	118.80
1	QA	620	C	C6-N1-C1'	-6.24	113.31	120.80
1	XA	1037	C	C6-N1-C1'	-6.24	113.31	120.80
1	XA	595	G	C8-N9-C1'	-6.24	118.89	127.00
34	YA	537	C	C6-N1-C1'	-6.24	113.32	120.80
1	XA	252	U	C2-N1-C1'	6.23	125.18	117.70
1	QA	19	C	C6-N1-C1'	-6.23	113.32	120.80
34	RA	2126	A	P-O3'-C3'	6.23	127.18	119.70
1	QA	1259	C	C6-N1-C1'	-6.23	113.32	120.80
1	XA	1242	C	C6-N1-C1'	-6.23	113.32	120.80
1	QA	117	G	C8-N9-C1'	-6.23	118.90	127.00
34	RA	1154	G	C8-N9-C1'	-6.23	118.90	127.00
1	QA	1263	C	C6-N1-C1'	-6.23	113.33	120.80
34	RA	343	C	C6-N1-C1'	-6.22	113.33	120.80
1	XA	186(H)	C	C2-N1-C1'	6.22	125.65	118.80
34	YA	591	C	C6-N1-C1'	-6.22	113.33	120.80
1	XA	974	A	C8-N9-C1'	6.22	138.90	127.70
34	YA	1201	C	C2-N1-C1'	6.22	125.64	118.80
34	YA	1506	C	C6-N1-C1'	-6.22	113.33	120.80
34	RA	749	C	C2-N1-C1'	6.22	125.64	118.80
34	RA	1404	C	C6-N1-C1'	-6.22	113.34	120.80
34	RA	1675	C	C6-N1-C1'	-6.22	113.34	120.80
34	RA	229	A	OP2-P-O3'	6.22	118.88	105.20
34	YA	619	G	C4-N9-C1'	6.22	134.58	126.50
1	XA	1128	C	C6-N1-C1'	-6.21	113.34	120.80
1	XA	1326	C	C2-N1-C1'	6.21	125.64	118.80
1	QA	261	U	C2-N1-C1'	6.21	125.15	117.70
34	RA	2891	G	C4-N9-C1'	6.21	134.57	126.50
38	RF	197	ASP	N-CA-C	-6.21	94.24	111.00
1	XA	1321	C	C2-N1-C1'	6.21	125.63	118.80
1	QA	1336	C	C2-N1-C1'	6.21	125.63	118.80
34	RA	1433	U	C6-N1-C1'	-6.20	112.52	121.20
34	YA	1870	C	C6-N1-C1'	-6.20	113.36	120.80
1	QA	1003	G	C4-N9-C1'	6.20	134.56	126.50
34	RA	1686	C	C6-N1-C1'	-6.20	113.36	120.80
34	YA	2471	C	C2-N1-C1'	6.20	125.62	118.80
34	RA	1510	A	C8-N9-C1'	-6.20	116.55	127.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	1499	A	C8-N9-C1'	-6.20	116.55	127.70
34	RA	2891	G	C8-N9-C1'	-6.20	118.94	127.00
1	XA	222	U	C2-N1-C1'	6.20	125.13	117.70
34	YA	2648	C	C2-N1-C1'	6.20	125.61	118.80
1	QA	570	G	C8-N9-C1'	-6.19	118.95	127.00
1	XA	868	C	C2-N1-C1'	6.19	125.61	118.80
1	XA	962	C	C2-N1-C1'	6.19	125.61	118.80
1	XA	974	A	C4-N9-C1'	-6.19	115.16	126.30
1	QA	963	G	C8-N9-C1'	-6.19	118.95	127.00
1	QA	358	U	C2-N1-C1'	6.18	125.12	117.70
34	RA	161	U	C6-N1-C1'	-6.18	112.54	121.20
34	YA	1542	G	C4-N9-C1'	6.18	134.54	126.50
34	RA	657	U	C6-N1-C1'	-6.18	112.55	121.20
1	QA	1140	C	C6-N1-C1'	-6.18	113.38	120.80
1	XA	68(I)	G	C4-N9-C1'	6.18	134.53	126.50
34	YA	619	G	C8-N9-C1'	-6.18	118.97	127.00
34	YA	2891	G	C8-N9-C1'	-6.17	118.97	127.00
34	RA	1741	C	C2-N1-C1'	6.17	125.59	118.80
34	RA	779	U	C6-N1-C1'	-6.17	112.56	121.20
1	QA	1429	C	C2-N1-C1'	6.17	125.58	118.80
1	QA	67	C	C2-N1-C1'	6.17	125.58	118.80
35	RB	70	C	C5-C6-N1	6.17	124.08	121.00
1	QA	578	C	C6-N1-C1'	-6.16	113.41	120.80
7	QG	73	MET	N-CA-CB	-6.16	99.50	110.60
1	QA	1414	U	C2-N1-C1'	6.16	125.09	117.70
34	YA	1327	C	C2-N1-C1'	6.16	125.58	118.80
34	RA	2439	A	P-O3'-C3'	6.16	127.09	119.70
22	XV	15	G	N3-C4-N9	6.16	129.69	126.00
34	RA	420	C	C6-N1-C1'	-6.16	113.41	120.80
34	YA	2374	C	C6-N1-C1'	-6.16	113.41	120.80
34	YA	669	G	C4-N9-C1'	6.15	134.50	126.50
1	XA	368	U	C6-N1-C1'	-6.15	112.59	121.20
34	YA	2437	U	C2-N1-C1'	6.15	125.08	117.70
34	YA	2231	C	C2-N1-C1'	6.15	125.57	118.80
34	RA	1819	A	P-O3'-C3'	6.15	127.08	119.70
34	RA	1218	C	C2-N1-C1'	6.15	125.56	118.80
1	XA	1109	C	C2-N1-C1'	6.15	125.56	118.80
1	QA	1003	G	C8-N9-C1'	-6.14	119.01	127.00
34	YA	2420	C	C2-N1-C1'	6.14	125.56	118.80
34	RA	2532	G	C4-N9-C1'	6.14	134.48	126.50
1	QA	1376	U	C6-N1-C1'	-6.14	112.61	121.20
34	YA	202	U	C2-N1-C1'	6.14	125.07	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	9	G	O5'-P-OP1	-6.14	100.18	105.70
34	RA	1411	C	C2-N1-C1'	6.14	125.55	118.80
22	XV	15	G	C2-N3-C4	6.14	114.97	111.90
34	RA	1930	G	C4-N9-C1'	6.13	134.47	126.50
34	YA	2207	C	C2-N1-C1'	6.13	125.55	118.80
1	XA	328	C	O4'-C1'-N1	6.13	113.11	108.20
34	RA	2752	C	C6-N1-C1'	-6.13	113.45	120.80
34	YA	1514	U	C2-N1-C1'	6.13	125.05	117.70
1	QA	651	C	C2-N1-C1'	6.12	125.54	118.80
1	XA	938	A	C4-N9-C1'	6.12	137.31	126.30
35	YB	86	G	N7-C8-N9	-6.12	110.04	113.10
34	RA	2689	U	P-O3'-C3'	6.12	127.04	119.70
1	XA	1253	G	C5'-C4'-O4'	-6.12	101.76	109.10
34	YA	2803	C	C2-N1-C1'	6.12	125.53	118.80
1	XA	1225	A	O4'-C1'-N9	6.12	113.09	108.20
34	RA	1328	G	C4-N9-C1'	6.12	134.45	126.50
34	RA	1312	U	P-O3'-C3'	6.11	127.04	119.70
1	XA	1437	C	C2-N1-C1'	-6.11	112.07	118.80
35	YB	29	A	C4-C5-N7	6.11	113.75	110.70
34	RA	1992	G	P-O3'-C3'	6.11	127.03	119.70
1	XA	68(I)	G	C8-N9-C1'	-6.11	119.06	127.00
34	YA	97	C	C2-N1-C1'	6.10	125.51	118.80
34	YA	1542	G	C8-N9-C1'	-6.10	119.07	127.00
1	XA	688	G	C4-N9-C1'	6.10	134.43	126.50
1	XA	797	C	C2-N1-C1'	6.10	125.51	118.80
34	RA	1694	C	P-O3'-C3'	6.10	127.02	119.70
34	RA	2234	G	C8-N9-C1'	-6.10	119.07	127.00
35	RB	54	G	C8-N9-C4	-6.10	103.96	106.40
41	RI	10	GLU	C-N-CA	6.10	136.94	121.70
34	RA	1542	G	C4-N9-C1'	6.09	134.42	126.50
36	RD	33	LEU	CA-CB-CG	6.09	129.32	115.30
1	XA	323	U	O5'-P-OP2	6.09	118.02	110.70
34	RA	2532	G	C8-N9-C1'	-6.09	119.08	127.00
34	YA	642	G	C4-N9-C1'	6.09	134.42	126.50
34	YA	1694	C	P-O3'-C3'	6.09	127.01	119.70
34	RA	2356	C	C2-N1-C1'	6.09	125.50	118.80
34	RA	2667	C	C6-N1-C1'	-6.09	113.49	120.80
34	YA	1315	C	C2-N1-C1'	6.09	125.50	118.80
1	QA	1366	C	C6-N1-C1'	-6.09	113.50	120.80
34	YA	395	U	C2-N1-C1'	6.09	125.00	117.70
1	QA	223	U	C6-N1-C1'	-6.08	112.68	121.20
1	XA	1203	C	C2-N1-C1'	6.08	125.49	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	1270	C	C2-N1-C1'	6.08	125.49	118.80
34	YA	221	A	P-O3'-C3'	6.08	127.00	119.70
1	XA	688	G	C8-N9-C1'	-6.08	119.10	127.00
1	XA	955	U	C6-N1-C1'	-6.08	112.69	121.20
35	YB	29	A	C5-N7-C8	-6.08	100.86	103.90
34	YA	1964	G	O4'-C1'-N9	-6.07	103.34	108.20
34	RA	1930	G	C8-N9-C1'	-6.07	119.12	127.00
1	XA	1507	A	N9-C1'-C2'	6.07	121.89	114.00
34	YA	669	G	C8-N9-C1'	-6.06	119.12	127.00
34	YA	1754	C	C6-N1-C1'	-6.06	113.52	120.80
1	QA	789	U	C6-N1-C1'	-6.06	112.72	121.20
34	YA	484	C	C2-N1-C1'	6.06	125.47	118.80
1	XA	1263	C	C2-N1-C1'	6.05	125.46	118.80
1	QA	97	U	C2-N1-C1'	6.05	124.96	117.70
1	QA	437	U	C6-N1-C1'	-6.05	112.73	121.20
1	QA	36	C	C6-N1-C1'	-6.05	113.54	120.80
1	XA	578	C	C6-N1-C1'	-6.05	113.54	120.80
1	XA	617	G	C3'-C2'-O2'	6.05	130.85	113.30
34	YA	184	C	C2-N1-C1'	6.05	125.45	118.80
34	YA	2884	U	C2-N1-C1'	6.05	124.96	117.70
34	RA	1542	G	C8-N9-C1'	-6.05	119.14	127.00
1	QA	1137	C	C2-N1-C1'	6.04	125.45	118.80
34	RA	1427	A	P-O3'-C3'	6.04	126.95	119.70
1	QA	355	C	C6-N1-C1'	-6.04	113.55	120.80
1	XA	307	C	C6-N1-C1'	-6.04	113.55	120.80
1	QA	1406	U	C6-N1-C1'	-6.04	112.75	121.20
34	RA	1920	C	C6-N1-C1'	-6.04	113.56	120.80
34	YA	595	C	C2-N1-C1'	6.04	125.44	118.80
34	YA	657	U	C2-N1-C1'	6.04	124.95	117.70
34	RA	1328	G	C8-N9-C1'	-6.04	119.15	127.00
28	Y4	5	ILE	N-CA-CB	6.04	124.68	110.80
34	YA	897	C	C6-N1-C1'	-6.04	113.56	120.80
34	RA	2359	C	C2-N1-C1'	6.03	125.44	118.80
1	QA	75	C	C2-N1-C1'	6.03	125.43	118.80
40	RH	87	LEU	N-CA-CB	-6.03	98.34	110.40
1	XA	1330	U	OP1-P-OP2	-6.03	110.56	119.60
34	RA	1314	C	C2-N1-C1'	6.03	125.43	118.80
34	YA	1108	U	C2-N1-C1'	6.03	124.93	117.70
34	YA	1240	U	C2-N1-C1'	6.03	124.93	117.70
1	XA	531	U	C6-N1-C1'	-6.02	112.77	121.20
34	YA	155	C	C2-N1-C1'	6.02	125.42	118.80
34	YA	544	C	C2-N1-C1'	6.02	125.42	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	252	U	C2-N1-C1'	6.02	124.92	117.70
34	RA	1407	C	C6-N1-C1'	-6.02	113.58	120.80
35	RB	37	C	C6-N1-C2	-6.02	117.89	120.30
1	XA	938	A	C8-N9-C1'	-6.02	116.87	127.70
34	RA	2574	G	C4-N9-C1'	6.02	134.32	126.50
1	XA	1285	A	P-O3'-C3'	6.02	126.92	119.70
34	RA	1611	C	C6-N1-C1'	-6.01	113.58	120.80
1	XA	864	A	C4-N9-C1'	6.01	137.13	126.30
34	YA	1879	C	C2-N1-C1'	6.01	125.42	118.80
34	YA	1047	G	C4-N9-C1'	6.01	134.31	126.50
34	YA	1774	C	C6-N1-C1'	-6.01	113.59	120.80
1	XA	277	C	C6-N1-C1'	-6.01	113.59	120.80
34	YA	2712(A)	U	P-O3'-C3'	6.01	126.91	119.70
34	RA	1101	U	C2-N1-C1'	6.00	124.91	117.70
1	XA	436	C	C2-N1-C1'	6.00	125.40	118.80
1	XA	1478	C	C2-N1-C1'	6.00	125.40	118.80
34	YA	642	G	C8-N9-C1'	-6.00	119.20	127.00
1	QA	1362(A)	C	C2-N1-C1'	6.00	125.40	118.80
34	YA	2439	A	P-O3'-C3'	6.00	126.90	119.70
34	RA	2040	C	C6-N1-C1'	-6.00	113.61	120.80
1	XA	587	G	C4-N9-C1'	6.00	134.29	126.50
1	XA	678	U	C6-N1-C1'	-6.00	112.81	121.20
34	RA	1662	C	C6-N1-C1'	-5.99	113.61	120.80
34	YA	1930	G	C4-N9-C1'	5.99	134.29	126.50
41	RI	130	TYR	C-N-CA	5.99	136.67	121.70
34	RA	541	C	C6-N1-C1'	-5.99	113.61	120.80
34	YA	267	C	C2-N1-C1'	5.99	125.39	118.80
34	YA	1049	C	C6-N1-C1'	-5.99	113.61	120.80
34	YA	1549	C	C2-N1-C1'	5.99	125.39	118.80
35	YB	31	C	C2-N1-C1'	5.99	125.39	118.80
35	YB	54	G	C6-C5-N7	-5.99	126.81	130.40
1	QA	390	C	C6-N1-C1'	-5.98	113.62	120.80
34	RA	1994	C	C2-N1-C1'	5.98	125.38	118.80
1	QA	536	C	O5'-P-OP2	-5.98	100.32	105.70
1	XA	587	G	C8-N9-C1'	-5.98	119.23	127.00
34	YA	613	U	C6-N1-C1'	-5.97	112.84	121.20
34	YA	2060	A	O4'-C1'-N9	-5.97	103.42	108.20
34	YA	635	C	C2-N1-C1'	5.97	125.37	118.80
1	XA	201(A)	C	C6-N1-C1'	-5.97	113.64	120.80
1	XA	241	C	C6-N1-C1'	-5.97	113.64	120.80
1	XA	1100	C	C2-N1-C1'	5.96	125.36	118.80
34	YA	1427	A	P-O3'-C3'	5.96	126.86	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	YA	1992	G	P-O3'-C3'	5.96	126.86	119.70
40	YH	12	PRO	N-CA-CB	-5.96	96.04	102.60
1	QA	456	C	C2-N1-C1'	5.96	125.36	118.80
1	QA	678	U	C2-N1-C1'	5.96	124.85	117.70
1	QA	1285	A	P-O3'-C3'	5.96	126.85	119.70
34	RA	2698	U	C6-N1-C1'	-5.96	112.85	121.20
34	RA	2824	C	C6-N1-C1'	-5.96	113.65	120.80
35	RB	70	C	C6-N1-C2	-5.96	117.92	120.30
34	YA	1675	C	C2-N1-C1'	5.96	125.36	118.80
34	YA	1047	G	C8-N9-C1'	-5.96	119.25	127.00
34	YA	1433	U	C6-N1-C1'	-5.96	112.86	121.20
34	YA	1235	G	C4-N9-C1'	5.95	134.24	126.50
34	YA	2582	G	C4-N9-C1'	5.95	134.24	126.50
34	YA	2710	C	C6-N1-C1'	-5.95	113.66	120.80
35	YB	22	U	C5-C6-N1	5.95	125.67	122.70
34	RA	2234	G	C4-N9-C1'	5.95	134.23	126.50
1	XA	169	C	C2-N1-C1'	5.95	125.34	118.80
1	XA	635	G	C4'-C3'-C2'	-5.95	96.65	102.60
1	XA	795	C	C2-N1-C1'	5.95	125.34	118.80
34	YA	2175	C	C2-N1-C1'	5.95	125.34	118.80
34	YA	2617	C	C6-N1-C1'	-5.95	113.66	120.80
1	QA	1343	G	C4-N9-C1'	5.95	134.23	126.50
1	QA	1359	C	C2-N1-C1'	5.95	125.34	118.80
34	RA	2574	G	C8-N9-C1'	-5.95	119.27	127.00
1	QA	905	U	C2-N1-C1'	5.94	124.83	117.70
1	QA	920	U	C2-N1-C1'	5.94	124.83	117.70
34	YA	419	C	C2-N1-C1'	5.94	125.34	118.80
34	RA	2095	C	C6-N1-C1'	-5.94	113.67	120.80
1	QA	177	C	C2-N1-C1'	5.94	125.33	118.80
34	RA	2739	U	C2-N1-C1'	5.94	124.83	117.70
34	YA	391	G	C4-N9-C1'	5.94	134.22	126.50
34	YA	2780	G	C8-N9-C1'	5.93	134.72	127.00
34	RA	2463	C	C2-N1-C1'	5.93	125.33	118.80
1	XA	1440(H)	C	C2-N1-C1'	5.93	125.33	118.80
34	YA	1672	C	C2-N1-C1'	5.93	125.33	118.80
34	YA	1881	C	C6-N1-C1'	-5.93	113.68	120.80
1	QA	822	C	C6-N1-C1'	-5.93	113.69	120.80
1	XA	619	U	C6-N1-C1'	5.93	129.50	121.20
1	XA	862	C	C2-N1-C1'	5.93	125.32	118.80
1	XA	1143	G	C8-N9-C1'	-5.93	119.29	127.00
34	YA	1516	U	C2-N1-C1'	5.93	124.82	117.70
34	YA	2394	C	O5'-P-OP2	5.93	117.82	110.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	RB	27	C	C5-C6-N1	5.93	123.96	121.00
1	XA	1493	A	C4-N9-C1'	5.93	136.97	126.30
34	YA	1930	G	C8-N9-C1'	-5.93	119.29	127.00
1	QA	930	C	C2-N1-C1'	5.93	125.32	118.80
1	QA	1327	C	OP1-P-O3'	5.93	118.24	105.20
34	YA	1982	C	C6-N1-C1'	-5.93	113.69	120.80
1	QA	1086	U	C6-N1-C1'	-5.92	112.91	121.20
1	XA	56	U	C2-N1-C1'	5.92	124.81	117.70
34	YA	838	C	C2-N1-C1'	5.92	125.31	118.80
1	QA	559	A	OP2-P-O3'	5.92	118.22	105.20
34	YA	1093	G	C4-N9-C1'	5.92	134.19	126.50
35	YB	75	G	C8-N9-C4	5.92	108.77	106.40
34	YA	568	U	C2-N1-C1'	5.92	124.80	117.70
34	YA	1093	G	C8-N9-C1'	-5.92	119.31	127.00
1	QA	1151	A	C4-N9-C1'	5.92	136.95	126.30
34	YA	1154	G	C4-N9-C1'	5.92	134.19	126.50
34	RA	856	C	C6-N1-C1'	-5.91	113.70	120.80
34	YA	1686	C	C6-N1-C1'	-5.91	113.70	120.80
1	QA	192	U	C2-N1-C1'	5.91	124.79	117.70
34	RA	581	C	C2-N1-C1'	5.91	125.30	118.80
1	QA	1211	U	C2-N1-C1'	5.91	124.79	117.70
34	RA	2144	U	C2-N1-C1'	5.91	124.79	117.70
35	RB	54	G	N7-C8-N9	5.91	116.05	113.10
34	YA	974(B)	C	C6-N1-C1'	-5.91	113.71	120.80
1	XA	864	A	C8-N9-C1'	-5.90	117.07	127.70
34	RA	1226	G	O5'-P-OP1	-5.90	100.39	105.70
34	RA	1930	G	P-O3'-C3'	5.90	126.78	119.70
34	RA	2342	C	C6-N1-C1'	-5.90	113.72	120.80
1	XA	121	C	C2-N1-C1'	5.90	125.29	118.80
34	YA	1097	U	C6-N1-C1'	-5.90	112.94	121.20
34	YA	2780	G	C4-N9-C1'	-5.89	118.84	126.50
34	RA	2034	U	C2-N1-C1'	5.89	124.77	117.70
34	YA	26	G	C4-N9-C1'	5.89	134.16	126.50
34	RA	254	G	C4-N9-C1'	5.89	134.16	126.50
1	XA	34	C	C2-N1-C1'	5.89	125.28	118.80
1	XA	335	C	C2-N1-C1'	5.89	125.28	118.80
1	QA	1282	C	C2-N1-C1'	5.88	125.27	118.80
1	QA	1343	G	C8-N9-C1'	-5.88	119.35	127.00
34	YA	1920	C	C2-N1-C1'	5.88	125.27	118.80
34	YA	2582	G	C8-N9-C1'	-5.88	119.35	127.00
34	RA	553	U	C2-N1-C1'	5.88	124.76	117.70
34	RA	1141	U	C6-N1-C1'	-5.88	112.96	121.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	YA	753	C	C6-N1-C1'	-5.88	113.74	120.80
1	QA	545	C	C2-N1-C1'	5.88	125.27	118.80
34	RA	2043	C	C6-N1-C1'	-5.88	113.74	120.80
1	QA	447	G	C4-N9-C1'	5.88	134.14	126.50
1	XA	656	C	C6-N1-C1'	-5.88	113.75	120.80
1	XA	1143	G	C4-N9-C1'	5.88	134.14	126.50
34	RA	245	G	C8-N9-C1'	-5.88	119.36	127.00
34	RA	1964	G	O4'-C1'-N9	-5.87	103.50	108.20
1	XA	796	C	C6-N1-C1'	-5.87	113.75	120.80
1	QA	136	C	C6-N1-C1'	-5.87	113.76	120.80
34	RA	1105	U	C6-N1-C1'	-5.87	112.98	121.20
1	XA	685	G	C4-N9-C1'	5.87	134.13	126.50
28	Y4	5	ILE	CA-CB-CG1	5.87	122.15	111.00
34	YA	1956	U	C6-N1-C1'	-5.87	112.98	121.20
34	YA	2776	A	P-O3'-C3'	5.87	126.74	119.70
1	XA	1056	U	C2-N1-C1'	5.87	124.74	117.70
1	QA	1517	G	C4-N9-C1'	5.87	134.13	126.50
1	XA	132	C	C2-N1-C1'	5.87	125.25	118.80
34	YA	364	C	C2-N1-C1'	5.86	125.25	118.80
34	YA	391	G	C8-N9-C1'	-5.86	119.38	127.00
34	YA	2691	C	C2-N1-C1'	5.86	125.25	118.80
1	QA	952	U	C6-N1-C1'	-5.86	112.99	121.20
34	YA	565	C	C6-N1-C1'	-5.86	113.77	120.80
1	XA	163	C	C2-N1-C1'	5.86	125.24	118.80
28	Y4	5	ILE	CA-CB-CG2	-5.86	99.19	110.90
34	YA	1537	C	C2-N1-C1'	5.86	125.24	118.80
34	RA	1649	G	N9-C1'-C2'	-5.86	105.56	112.00
34	RA	1864	U	C2-N1-C1'	5.86	124.73	117.70
4	XD	8	VAL	CG1-CB-CG2	5.86	120.27	110.90
34	YA	796	C	C2-N1-C1'	5.86	125.24	118.80
35	YB	55	U	N3-C4-O4	5.85	123.50	119.40
54	RZ	59	LEU	CA-CB-CG	5.85	128.76	115.30
34	YA	1235	G	C8-N9-C1'	-5.85	119.39	127.00
34	YA	1437	C	C2-N1-C1'	5.85	125.24	118.80
34	RA	245	G	C4-N9-C1'	5.85	134.11	126.50
1	XA	857	C	C2-N1-C1'	5.85	125.24	118.80
34	RA	1688	U	C2-N1-C1'	5.84	124.72	117.70
34	RA	365	C	C2-N1-C1'	5.84	125.23	118.80
34	RA	1892	C	C6-N1-C1'	-5.84	113.79	120.80
34	RA	254	G	C8-N9-C1'	-5.84	119.41	127.00
1	XA	718	G	O4'-C1'-N9	-5.84	103.53	108.20
1	XA	1493	A	C8-N9-C1'	-5.84	117.19	127.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	YA	650	C	C6-N1-C1'	-5.84	113.79	120.80
34	RA	270(Q)	C	C6-N1-C1'	-5.84	113.80	120.80
34	YA	1927	A	C4-N9-C1'	5.84	136.81	126.30
1	XA	980	C	OP1-P-O3'	5.84	118.04	105.20
1	XA	1075	C	C2-N1-C1'	5.84	125.22	118.80
34	YA	2097	C	C6-N1-C1'	-5.84	113.80	120.80
34	YA	2895	U	C6-N1-C1'	-5.84	113.03	121.20
1	QA	825	G	C4-N9-C1'	5.83	134.08	126.50
34	RA	1990	C	C6-N1-C1'	-5.83	113.80	120.80
1	XA	626	U	O5'-P-OP1	-5.83	100.45	105.70
1	QA	1151	A	C8-N9-C1'	-5.83	117.20	127.70
1	QA	1045	C	C6-N1-C1'	-5.83	113.80	120.80
1	QA	1440(H)	C	C6-N1-C1'	-5.83	113.81	120.80
34	RA	2776	A	P-O3'-C3'	5.83	126.69	119.70
1	XA	834	C	C6-N1-C1'	-5.83	113.81	120.80
34	RA	1474	C	C2-N1-C1'	5.83	125.21	118.80
35	YB	93	C	N1-C2-O2	5.83	122.39	118.90
34	YA	2150	U	C2-N1-C1'	5.82	124.69	117.70
34	RA	276	A	O5'-P-OP1	5.82	117.69	110.70
34	RA	943	U	C2-N1-C1'	5.82	124.69	117.70
34	RA	2093	G	C4-N9-C1'	5.82	134.07	126.50
34	YA	1085	A	P-O3'-C3'	5.82	126.68	119.70
34	YA	1154	G	C8-N9-C1'	-5.82	119.44	127.00
34	YA	2040	C	C6-N1-C1'	-5.82	113.82	120.80
1	QA	500	G	C4-N9-C1'	5.82	134.06	126.50
35	RB	15	A	C8-N9-C4	5.82	108.13	105.80
1	QA	500	G	C8-N9-C1'	-5.82	119.44	127.00
1	QA	1075	C	C6-N1-C1'	-5.82	113.82	120.80
34	RA	753	C	C6-N1-C1'	-5.82	113.82	120.80
35	RB	6	C	C6-N1-C2	-5.82	117.97	120.30
34	YA	2146	C	C6-N1-C1'	-5.81	113.83	120.80
1	QA	1517	G	C8-N9-C1'	-5.81	119.44	127.00
34	YA	26	G	C8-N9-C1'	-5.81	119.45	127.00
34	YA	779	U	C2-N1-C1'	5.81	124.67	117.70
1	QA	564	C	C2-N1-C1'	5.81	125.19	118.80
32	R8	62	LEU	CA-CB-CG	5.81	128.66	115.30
34	RA	1670	C	C2-N1-C1'	5.81	125.19	118.80
1	XA	283	C	C6-N1-C1'	-5.81	113.83	120.80
34	YA	2195	C	C6-N1-C1'	-5.81	113.83	120.80
34	YA	678	C	C2-N1-C1'	5.81	125.19	118.80
1	XA	685	G	C8-N9-C1'	-5.80	119.45	127.00
1	XA	805	C	C6-N1-C1'	-5.80	113.83	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	RA	1799	G	P-O3'-C3'	5.80	126.66	119.70
1	XA	186(O)	U	C6-N1-C1'	-5.80	113.08	121.20
1	QA	447	G	C8-N9-C1'	-5.80	119.46	127.00
34	RA	284	U	C2-N1-C1'	5.80	124.66	117.70
34	YA	2688	U	O4'-C1'-N1	5.80	112.84	108.20
1	QA	1384	C	C6-N1-C1'	-5.80	113.84	120.80
34	RA	1707	G	C4-N9-C1'	5.80	134.04	126.50
1	XA	1494	G	C4-N9-C1'	5.80	134.04	126.50
34	YA	1598	C	C6-N1-C1'	-5.80	113.84	120.80
34	YA	1640	C	C6-N1-C1'	-5.80	113.84	120.80
34	YA	2482	G	C4-N9-C1'	5.80	134.04	126.50
1	QA	1074	G	C4-N9-C1'	5.80	134.04	126.50
34	YA	1510	A	O5'-P-OP2	-5.80	100.48	105.70
34	RA	1186	G	C4-N9-C1'	5.79	134.03	126.50
12	XL	47	LYS	N-CA-C	5.79	126.64	111.00
34	YA	1019	U	O5'-P-OP1	-5.79	100.49	105.70
34	RA	2075	U	C6-N1-C1'	-5.79	113.09	121.20
1	QA	1237	C	C2-N1-C1'	5.79	125.17	118.80
34	RA	825	C	C2-N1-C1'	5.79	125.17	118.80
34	RA	2401	U	C2-N1-C1'	5.79	124.64	117.70
34	YA	2554	U	C6-N1-C1'	-5.79	113.10	121.20
34	RA	725	G	C4-N9-C1'	5.79	134.02	126.50
1	XA	1465	C	C6-N1-C1'	-5.79	113.86	120.80
1	QA	825	G	C8-N9-C1'	-5.78	119.48	127.00
34	RA	856	C	P-O3'-C3'	5.78	126.64	119.70
34	RA	1186	G	C8-N9-C1'	-5.78	119.48	127.00
1	XA	309	G	O5'-P-OP2	-5.78	100.50	105.70
34	RA	2179	C	C6-N1-C1'	-5.78	113.86	120.80
34	YA	2586	C	C2-N1-C1'	5.78	125.16	118.80
34	RA	1777	U	C6-N1-C1'	-5.78	113.11	121.20
1	XA	266	G	OP2-P-O3'	5.78	117.91	105.20
34	YA	2460	U	C2-N1-C1'	5.78	124.64	117.70
34	YA	2689	U	P-O3'-C3'	5.78	126.64	119.70
34	YA	284	U	C2-N1-C1'	5.78	124.63	117.70
1	QA	133	U	C2-N1-C1'	5.78	124.63	117.70
34	RA	1774	C	C6-N1-C1'	-5.78	113.87	120.80
1	XA	1354	C	C2-N1-C1'	5.78	125.15	118.80
34	YA	2466	C	C6-N1-C1'	-5.78	113.87	120.80
34	RA	1516	U	C6-N1-C1'	-5.77	113.12	121.20
34	YA	1405	U	C2-N1-C1'	5.77	124.63	117.70
34	YA	2658	C	C6-N1-C1'	-5.77	113.87	120.80
34	YA	1372	U	C6-N1-C1'	-5.77	113.12	121.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	YA	1649	G	C4-N9-C1'	5.77	134.00	126.50
1	QA	516	U	C2-N1-C1'	5.77	124.62	117.70
1	QA	1213	A	C4-N9-C1'	5.77	136.68	126.30
1	XA	1364	U	C2-N1-C1'	5.77	124.62	117.70
34	RA	18	C	C6-N1-C1'	-5.77	113.88	120.80
34	RA	140	A	O4'-C1'-N9	5.76	112.81	108.20
1	XA	458(B)	G	C4-N9-C1'	5.76	134.00	126.50
34	YA	318	C	C2-N1-C1'	5.76	125.14	118.80
1	QA	1215	G	C4-N9-C1'	5.76	133.99	126.50
1	XA	989	C	C6-N1-C1'	-5.76	113.88	120.80
1	QA	67	C	C6-N1-C1'	-5.76	113.89	120.80
1	QA	1377	A	O4'-C1'-N9	5.76	112.81	108.20
34	YA	76	C	C6-N1-C1'	-5.76	113.89	120.80
34	RA	687	C	C6-N1-C1'	-5.76	113.89	120.80
34	YA	1927	A	C8-N9-C1'	-5.76	117.33	127.70
34	RA	1509	C	C6-N1-C1'	-5.76	113.89	120.80
1	XA	1090	U	C2-N1-C1'	5.76	124.61	117.70
34	YA	1577	C	C2-N1-C1'	5.76	125.13	118.80
1	QA	824	C	C2-N1-C1'	5.75	125.13	118.80
34	RA	1180	C	C2-N1-C1'	5.75	125.13	118.80
1	XA	501	C	C6-N1-C1'	-5.75	113.89	120.80
34	RA	725	G	C8-N9-C1'	-5.75	119.52	127.00
1	QA	353	A	C4-N9-C1'	5.75	136.65	126.30
34	RA	512	G	P-O3'-C3'	5.75	126.60	119.70
34	YA	1649	G	C8-N9-C1'	-5.75	119.52	127.00
34	YA	1819	A	P-O3'-C3'	5.75	126.60	119.70
1	QA	176	C	C2-N1-C1'	5.75	125.12	118.80
34	RA	1066	U	C6-N1-C1'	-5.75	113.15	121.20
35	YB	97	G	N1-C2-N3	5.75	127.35	123.90
34	RA	26	G	C4-N9-C1'	5.75	133.97	126.50
1	XA	236	G	C4-N9-C1'	5.75	133.97	126.50
34	RA	2093	G	C8-N9-C1'	-5.75	119.53	127.00
34	YA	2233	U	C2-N1-C1'	5.74	124.59	117.70
35	YB	66	A	P-O3'-C3'	5.74	126.59	119.70
34	RA	2064	C	C2-N1-C1'	5.74	125.11	118.80
1	XA	1317	C	C2-N1-C1'	5.74	125.11	118.80
22	XV	15	G	C8-N9-C1'	-5.74	119.54	127.00
34	YA	343	C	C6-N1-C1'	-5.74	113.91	120.80
34	YA	384	U	C6-N1-C1'	-5.74	113.17	121.20
1	XA	1437	C	P-O3'-C3'	-5.74	112.82	119.70
1	QA	943	U	C2-N1-C1'	5.74	124.58	117.70
1	QA	1401	G	C4-N9-C1'	5.74	133.96	126.50

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	RA	678	C	C2-N1-C1'	5.74	125.11	118.80
34	RA	1982	C	C2-N1-C1'	5.74	125.11	118.80
1	XA	117	G	C4-N9-C1'	5.74	133.96	126.50
1	XA	1326	C	C6-N1-C1'	-5.73	113.92	120.80
34	YA	1289	C	C2-N1-C1'	5.73	125.11	118.80
1	QA	1403	C	C6-N1-C1'	-5.73	113.92	120.80
35	RB	66	A	P-O3'-C3'	5.73	126.58	119.70
1	QA	370	C	C2-N1-C1'	5.73	125.10	118.80
1	QA	1247	U	C2-N1-C1'	5.73	124.58	117.70
34	RA	25	U	C2-N1-C1'	5.73	124.58	117.70
34	YA	271(C)	G	OP2-P-O3'	5.73	117.81	105.20
35	YB	42	C	C6-N1-C2	-5.73	118.01	120.30
23	XX	19	U	C2-N1-C1'	5.73	124.57	117.70
34	YA	871	U	C2-N1-C1'	5.73	124.57	117.70
34	YA	2145	C	C6-N1-C1'	-5.73	113.93	120.80
34	YA	2452	C	C2-N1-C1'	5.73	125.10	118.80
1	QA	1382	C	C6-N1-C1'	-5.72	113.93	120.80
34	RA	1707	G	C8-N9-C1'	-5.72	119.56	127.00
1	QA	1074	G	C8-N9-C1'	-5.72	119.56	127.00
22	QV	31	G	C4-N9-C1'	5.72	133.94	126.50
34	YA	1101	U	C2-N1-C1'	5.72	124.57	117.70
34	YA	2291	U	C2-N1-C1'	5.72	124.57	117.70
1	QA	556	C	C2-N1-C1'	5.72	125.09	118.80
34	RA	1049	C	C6-N1-C1'	-5.72	113.94	120.80
34	YA	1914	C	C6-N1-C1'	-5.72	113.94	120.80
34	RA	1121	C	C2-N1-C1'	5.72	125.09	118.80
1	XA	757	U	C6-N1-C1'	-5.72	113.20	121.20
1	XA	1537	U	OP1-P-O3'	5.71	117.77	105.20
22	XV	76	A	N7-C8-N9	5.71	116.66	113.80
34	RA	556	G	C4-N9-C1'	5.71	133.93	126.50
34	RA	839	U	C2-N1-C1'	5.71	124.56	117.70
1	XA	1313	U	C6-N1-C1'	-5.71	113.20	121.20
34	YA	1411	C	C2-N1-C1'	5.71	125.08	118.80
1	XA	513	C	C2-N1-C1'	5.71	125.08	118.80
1	XA	1494	G	C8-N9-C1'	-5.71	119.58	127.00
34	YA	2482	G	C8-N9-C1'	-5.71	119.58	127.00
35	YB	79	C	N3-C2-O2	-5.71	117.90	121.90
35	YB	82	G	N7-C8-N9	5.71	115.95	113.10
34	RA	683	C	C6-N1-C1'	-5.71	113.95	120.80
34	RA	568	U	C6-N1-C1'	-5.70	113.22	121.20
1	QA	381	C	C6-N1-C1'	-5.70	113.96	120.80
1	QA	580	U	C6-N1-C1'	-5.70	113.22	121.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	RA	1327	C	C6-N1-C1'	-5.70	113.96	120.80
34	YA	1556	C	C2-N1-C1'	5.70	125.07	118.80
34	YA	1734	C	C2-N1-C1'	5.70	125.07	118.80
35	YB	43	C	C6-N1-C2	-5.70	118.02	120.30
1	XA	607	A	O4'-C1'-N9	5.70	112.76	108.20
34	RA	66	C	C2-N1-C1'	5.69	125.06	118.80
34	RA	2312	U	C2-N1-C1'	5.69	124.53	117.70
1	XA	117	G	C8-N9-C1'	-5.69	119.60	127.00
1	QA	1215	G	C8-N9-C1'	-5.69	119.60	127.00
34	RA	395	U	C2-N1-C1'	5.69	124.53	117.70
34	RA	2174	C	C6-N1-C1'	-5.69	113.97	120.80
35	YB	80	U	C5-C4-O4	5.69	129.31	125.90
1	XA	832	C	C2-N1-C1'	5.69	125.06	118.80
34	RA	634	C	C2-N1-C1'	5.69	125.06	118.80
34	RA	1188	U	C6-N1-C1'	-5.69	113.24	121.20
34	RA	2636	U	C6-N1-C1'	-5.69	113.24	121.20
1	XA	236	G	C8-N9-C1'	-5.69	119.61	127.00
1	XA	458(B)	G	C8-N9-C1'	-5.69	119.61	127.00
34	YA	2663	G	C4-N9-C1'	5.68	133.89	126.50
1	XA	352	C	C2-N1-C1'	5.68	125.05	118.80
34	YA	208	C	C2-N1-C1'	5.68	125.05	118.80
14	QN	43	CYS	CB-CA-C	5.68	121.76	110.40
1	XA	34	C	C6-N1-C1'	-5.68	113.98	120.80
1	XA	1240	U	C2-N1-C1'	-5.68	110.88	117.70
35	RB	91	C	C6-N1-C2	-5.68	118.03	120.30
34	YA	1391	U	C6-N1-C1'	-5.68	113.25	121.20
45	YQ	19	GLY	N-CA-C	-5.68	98.91	113.10
1	XA	134	A	C4-N9-C1'	5.67	136.51	126.30
34	YA	1448	G	C4-N9-C1'	5.67	133.88	126.50
1	QA	1322	C	C2-N1-C1'	5.67	125.04	118.80
34	RA	1218	C	C6-N1-C1'	-5.67	113.99	120.80
34	YA	1588	C	C2-N1-C1'	5.67	125.04	118.80
34	RA	2210	G	C4-N9-C1'	5.67	133.87	126.50
34	RA	2664	G	C4-N9-C1'	5.66	133.86	126.50
34	YA	1201	C	C6-N1-C1'	-5.66	114.01	120.80
34	YA	2084	C	C2-N1-C1'	5.66	125.03	118.80
1	QA	1401	G	C8-N9-C1'	-5.66	119.64	127.00
1	XA	343	U	C6-N1-C1'	-5.66	113.28	121.20
34	YA	923	C	C2-N1-C1'	5.66	125.02	118.80
34	RA	26	G	C8-N9-C1'	-5.66	119.65	127.00
1	XA	186(H)	C	C6-N1-C1'	-5.66	114.01	120.80
1	XA	1440(A)	C	C2-N1-C1'	5.66	125.02	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	YA	18	C	C2-N1-C1'	5.66	125.02	118.80
1	QA	1213	A	C8-N9-C1'	-5.66	117.52	127.70
34	RA	556	G	C8-N9-C1'	-5.66	119.65	127.00
34	YA	271(B)	C	C2-N1-C1'	5.65	125.02	118.80
35	YB	1	U	C6-N1-C2	-5.65	117.61	121.00
1	QA	295	C	C2-N1-C1'	5.65	125.02	118.80
34	YA	464	U	C6-N1-C1'	-5.65	113.28	121.20
34	YA	2763	G	C4-N9-C1'	5.65	133.85	126.50
1	QA	1351	U	C6-N1-C1'	-5.65	113.29	121.20
1	XA	1357	A	C4-N9-C1'	5.65	136.47	126.30
34	RA	1240	U	C6-N1-C1'	-5.65	113.29	121.20
1	QA	353	A	C8-N9-C1'	-5.65	117.53	127.70
34	RA	749	C	C6-N1-C1'	-5.65	114.02	120.80
34	YA	1417	C	C2-N1-C1'	5.65	125.01	118.80
34	YA	2471	C	C6-N1-C1'	-5.65	114.02	120.80
34	RA	568	U	O5'-P-OP1	-5.65	100.62	105.70
1	QA	1336	C	C6-N1-C1'	-5.64	114.03	120.80
34	RA	2496	C	O5'-P-OP1	-5.64	100.62	105.70
35	YB	58	A	N1-C6-N6	-5.64	115.21	118.60
1	XA	62	U	C6-N1-C1'	-5.64	113.30	121.20
34	YA	229	A	OP2-P-O3'	5.64	117.61	105.20
34	YA	2663	G	C8-N9-C1'	-5.64	119.67	127.00
37	YE	186	GLY	N-CA-C	5.64	127.19	113.10
1	XA	537	G	C8-N9-C1'	5.64	134.33	127.00
34	YA	2179	C	C2-N1-C1'	5.64	125.00	118.80
1	XA	1321	C	C6-N1-C1'	-5.63	114.04	120.80
34	YA	2756	U	OP1-P-O3'	5.63	117.59	105.20
1	XA	309	G	P-O3'-C3'	5.63	126.46	119.70
34	RA	2318	G	O4'-C1'-N9	5.63	112.70	108.20
1	QA	1359	C	C6-N1-C1'	-5.63	114.05	120.80
1	XA	68(J)	G	C4-N9-C1'	5.63	133.81	126.50
34	RA	2210	G	C8-N9-C1'	-5.62	119.69	127.00
34	RA	848	G	C4-N9-C1'	5.62	133.81	126.50
1	XA	1243	C	O4'-C1'-N1	5.62	112.70	108.20
36	YD	241	PRO	C-N-CA	5.62	135.76	121.70
1	XA	868	C	C6-N1-C1'	-5.62	114.06	120.80
1	QA	343	U	C6-N1-C1'	-5.62	113.33	121.20
34	RA	1005	C	C2-N1-C1'	5.62	124.98	118.80
1	XA	1328	C	OP2-P-O3'	5.62	117.56	105.20
1	QA	1538	C	OP1-P-O3'	5.62	117.56	105.20
34	YA	2752	C	C2-N1-C1'	5.62	124.98	118.80
34	RA	1992	G	OP2-P-O3'	5.62	117.55	105.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	RA	2867	G	C4-N9-C1'	5.62	133.80	126.50
34	YA	270(D)	C	C2-N1-C1'	5.62	124.98	118.80
1	XA	797	C	C6-N1-C1'	-5.61	114.07	120.80
1	XA	980	C	C3'-C2'-C1'	-5.61	97.01	101.50
1	QA	1429	C	C6-N1-C1'	-5.61	114.07	120.80
34	RA	2785	C	C2-N1-C1'	5.61	124.97	118.80
1	XA	312	C	C2-N1-C1'	5.61	124.97	118.80
1	XA	905	U	C2-N1-C1'	5.61	124.43	117.70
1	XA	1231	G	OP1-P-O3'	-5.61	92.87	105.20
20	QT	73	HIS	CA-CB-CG	5.60	123.13	113.60
34	RA	102	G	P-O3'-C3'	5.60	126.42	119.70
34	RA	1741	C	C6-N1-C1'	-5.60	114.08	120.80
34	RA	2466	C	C2-N1-C1'	5.60	124.96	118.80
34	YA	1327	C	C6-N1-C1'	-5.60	114.08	120.80
34	RA	752	A	P-O3'-C3'	5.60	126.42	119.70
1	XA	636	U	P-O5'-C5'	5.60	129.86	120.90
34	RA	1411	C	C6-N1-C1'	-5.60	114.08	120.80
34	YA	155	C	C6-N1-C1'	-5.60	114.08	120.80
34	YA	2207	C	C6-N1-C1'	-5.60	114.08	120.80
34	YA	2566	A	P-O3'-C3'	5.60	126.42	119.70
34	YA	2763	G	C8-N9-C1'	-5.60	119.72	127.00
1	XA	848	C	C2-N1-C1'	5.59	124.95	118.80
34	YA	887	A	OP1-P-OP2	5.59	127.99	119.60
34	YA	2648	C	C6-N1-C1'	-5.59	114.09	120.80
1	QA	56	U	C6-N1-C1'	-5.59	113.37	121.20
35	RB	60	C	C5-C6-N1	5.59	123.80	121.00
34	YA	1448	G	C8-N9-C1'	-5.59	119.73	127.00
1	QA	1229	A	C4-N9-C1'	5.59	136.37	126.30
34	RA	2044	C	C2-N1-C1'	5.59	124.95	118.80
35	RB	108	C	C6-N1-C2	5.59	122.53	120.30
1	XA	134	A	C8-N9-C1'	-5.59	117.64	127.70
35	YB	54	G	N3-C4-C5	5.59	131.39	128.60
34	RA	848	G	C8-N9-C1'	-5.59	119.73	127.00
34	YA	2231	C	C6-N1-C1'	-5.59	114.09	120.80
1	XA	749	C	C2-N1-C1'	5.59	124.94	118.80
1	XA	962	C	C6-N1-C1'	-5.59	114.09	120.80
34	YA	862	G	C4-N9-C1'	5.59	133.76	126.50
34	YA	1777	U	C2-N1-C1'	5.59	124.40	117.70
1	QA	428	G	C4-N9-C1'	5.58	133.76	126.50
1	QA	1417	G	C4-N9-C1'	5.58	133.76	126.50
34	RA	1298	C	C2-N1-C1'	5.58	124.94	118.80
34	RA	1776	G	C8-N9-C1'	-5.58	119.74	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	1358	U	C2-N1-C1'	-5.58	111.00	117.70
1	QA	651	C	C6-N1-C1'	-5.58	114.11	120.80
34	YA	97	C	C6-N1-C1'	-5.58	114.11	120.80
44	RP	59	LEU	CA-CB-CG	5.58	128.12	115.30
1	XA	1158	C	N1-C1'-C2'	5.58	121.25	114.00
34	YA	2420	C	C6-N1-C1'	-5.58	114.11	120.80
1	QA	962	C	C2-N1-C1'	5.57	124.93	118.80
1	QA	1303	C	C2-N1-C1'	5.57	124.93	118.80
1	XA	537	G	C4-N9-C1'	-5.57	119.26	126.50
1	QA	647	C	C2-N1-C1'	5.57	124.93	118.80
34	RA	1776	G	C4-N9-C1'	5.57	133.74	126.50
1	XA	400	C	C2-N1-C1'	5.57	124.93	118.80
34	YA	484	C	C6-N1-C1'	-5.57	114.12	120.80
34	YA	636	G	C8-N9-C1'	-5.57	119.76	127.00
1	QA	1270	C	C6-N1-C1'	-5.57	114.12	120.80
29	R5	34	PRO	CA-N-CD	-5.57	103.71	111.50
1	XA	1109	C	C6-N1-C1'	-5.57	114.12	120.80
34	YA	602	G	C4-N9-C1'	5.57	133.74	126.50
1	XA	12	U	C6-N1-C1'	-5.56	113.41	121.20
34	YA	636	G	C4-N9-C1'	5.56	133.73	126.50
34	YA	2210	G	N9-C1'-C2'	5.56	121.23	114.00
34	YA	2704	C	C2-N1-C1'	5.56	124.92	118.80
34	RA	2664	G	C8-N9-C1'	-5.56	119.77	127.00
1	QA	1197	G	C8-N9-C1'	-5.56	119.77	127.00
1	QA	849	C	C2-N1-C1'	5.56	124.91	118.80
1	XA	68(J)	G	C8-N9-C1'	-5.56	119.77	127.00
1	QA	1148	U	C6-N1-C1'	-5.56	113.42	121.20
34	RA	2663	G	C4-N9-C1'	5.55	133.72	126.50
34	YA	595	C	C6-N1-C1'	-5.55	114.13	120.80
34	YA	2134	A	C8-N9-C1'	-5.55	117.70	127.70
34	YA	2730	C	C2-N1-C1'	5.55	124.91	118.80
1	QA	1137	C	C6-N1-C1'	-5.55	114.14	120.80
1	XA	980	C	OP2-P-O3'	-5.55	92.99	105.20
1	XA	1357	A	C8-N9-C1'	-5.55	117.71	127.70
34	RA	1304	C	C2-N1-C1'	5.55	124.91	118.80
34	YA	448	U	C2-N1-C1'	5.55	124.36	117.70
1	QA	1197	G	C4-N9-C1'	5.55	133.71	126.50
1	XA	1098	C	C2-N1-C1'	5.55	124.90	118.80
34	RA	1914	C	O4'-C1'-N1	5.54	112.64	108.20
1	QA	1417	G	C8-N9-C1'	-5.54	119.79	127.00
34	RA	2356	C	C6-N1-C1'	-5.54	114.15	120.80
34	RA	2420	C	C2-N1-C1'	5.54	124.90	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	1437	C	C6-N1-C1'	5.54	127.45	120.80
34	RA	2663	G	C8-N9-C1'	-5.54	119.80	127.00
1	XA	479	C	C2-N1-C1'	5.54	124.89	118.80
28	Y4	39	CYS	N-CA-C	-5.54	96.04	111.00
32	R8	61	LEU	C-N-CA	5.54	135.55	121.70
34	YA	1315	C	C6-N1-C1'	-5.54	114.15	120.80
34	RA	1312	U	C2-N1-C1'	5.54	124.35	117.70
34	RA	1313	U	C2-N1-C1'	5.54	124.35	117.70
54	YZ	62	PRO	C-N-CA	5.54	135.54	121.70
34	YA	544	C	C6-N1-C1'	-5.54	114.16	120.80
34	YA	2803	C	C6-N1-C1'	-5.54	114.16	120.80
34	RA	2566	A	P-O3'-C3'	5.53	126.34	119.70
34	RA	2580	U	C2-N1-C1'	5.53	124.34	117.70
34	YA	184	C	C6-N1-C1'	-5.53	114.16	120.80
34	YA	576	U	C2-N1-C1'	5.53	124.34	117.70
1	QA	794	A	C4-N9-C1'	5.53	136.25	126.30
34	RA	202	U	C2-N1-C1'	5.53	124.33	117.70
34	RA	1483	G	C4-N9-C1'	5.53	133.69	126.50
40	RH	153	LYS	N-CA-C	5.53	125.92	111.00
34	RA	2359	C	C6-N1-C1'	-5.53	114.17	120.80
34	YA	1507	A	O4'-C1'-N9	5.53	112.62	108.20
1	QA	1388	C	C2-N1-C1'	5.52	124.88	118.80
1	XA	1203	C	C6-N1-C1'	-5.52	114.17	120.80
1	XA	1353	G	C8-N9-C1'	-5.52	119.82	127.00
34	YA	1515	C	C2-N1-C1'	5.52	124.88	118.80
34	RA	2575	C	C2-N1-C1'	5.52	124.87	118.80
1	XA	420	U	C2-N1-C1'	5.52	124.33	117.70
1	QA	1229	A	C8-N9-C1'	-5.52	117.76	127.70
35	YB	61	G	C8-N9-C4	-5.52	104.19	106.40
34	RA	509	C	C2-N1-C1'	5.52	124.87	118.80
1	XA	1105	A	N9-C1'-C2'	-5.52	105.93	112.00
34	RA	2867	G	C8-N9-C1'	-5.51	119.83	127.00
34	YA	2133	G	C4-N9-C1'	5.51	133.67	126.50
34	YA	9	U	C2-N1-C1'	5.51	124.31	117.70
34	YA	270(N)	U	C2-N1-C1'	5.51	124.31	117.70
1	QA	956	U	C2-N1-C1'	5.51	124.31	117.70
1	XA	25	C	C2-N1-C1'	5.51	124.86	118.80
1	XA	526	C	C2-N1-C1'	5.51	124.86	118.80
34	YA	602	G	C8-N9-C1'	-5.51	119.84	127.00
1	QA	1352	C	C2-N1-C1'	5.51	124.86	118.80
34	YA	2334	G	O4'-C1'-N9	-5.51	103.79	108.20
34	YA	2475	C	C2-N1-C1'	5.51	124.86	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	RA	619	G	C4-N9-C1'	5.51	133.66	126.50
34	YA	2808	U	C2-N1-C1'	5.51	124.31	117.70
1	XA	1353	G	C4-N9-C1'	5.50	133.65	126.50
34	YA	848	G	C4-N9-C1'	5.50	133.66	126.50
34	YA	2532	G	C4-N9-C1'	5.50	133.66	126.50
34	RA	817	C	C2-N1-C1'	5.50	124.85	118.80
1	XA	37	U	C2-N1-C1'	5.50	124.30	117.70
34	YA	365	C	C2-N1-C1'	5.50	124.85	118.80
34	YA	1879	C	C6-N1-C1'	-5.50	114.20	120.80
44	YP	59	LEU	CA-CB-CG	5.50	127.95	115.30
34	RA	2205	C	C2-N1-C1'	5.50	124.85	118.80
1	XA	858	G	C4-N9-C1'	5.50	133.64	126.50
35	YB	37	C	N1-C2-O2	5.50	122.20	118.90
1	QA	75	C	C6-N1-C1'	-5.49	114.21	120.80
1	QA	428	G	C8-N9-C1'	-5.49	119.86	127.00
34	YA	862	G	C8-N9-C1'	-5.49	119.86	127.00
1	QA	1362(A)	C	C6-N1-C1'	-5.49	114.21	120.80
34	YA	206	U	C2-N1-C1'	5.49	124.29	117.70
34	YA	546	C	C2-N1-C1'	5.49	124.84	118.80
34	YA	2511	U	C2-N1-C1'	5.49	124.29	117.70
1	XA	1440(A)	C	C6-N1-C1'	-5.49	114.22	120.80
34	YA	1498	C	C2-N1-C1'	5.49	124.84	118.80
34	RA	201	C	C2-N1-C1'	5.49	124.83	118.80
34	RA	2404	C	C2-N1-C1'	5.49	124.83	118.80
34	YA	1599	C	C2-N1-C1'	5.49	124.83	118.80
34	RA	806	C	C2-N1-C1'	5.48	124.83	118.80
35	RB	31	C	C6-N1-C2	-5.48	118.11	120.30
1	XA	1263	C	C6-N1-C1'	-5.48	114.22	120.80
1	QA	707	C	C2-N1-C1'	5.48	124.83	118.80
34	RA	1994	C	C6-N1-C1'	-5.48	114.22	120.80
34	RA	2233	U	C2-N1-C1'	5.48	124.28	117.70
34	YA	838	C	C6-N1-C1'	-5.48	114.22	120.80
34	YA	2824	C	C2-N1-C1'	5.48	124.83	118.80
1	XA	266	G	P-O3'-C3'	5.48	126.27	119.70
34	YA	898	C	C2-N1-C1'	5.48	124.83	118.80
1	XA	1219	U	C6-N1-C1'	5.48	128.87	121.20
34	YA	2496	C	O5'-P-OP1	-5.48	100.77	105.70
1	QA	1409	C	C2-N1-C1'	5.47	124.82	118.80
34	RA	435	C	C2-N1-C1'	5.47	124.82	118.80
34	RA	2096	U	C6-N1-C1'	-5.47	113.54	121.20
1	XA	1302	U	P-O3'-C3'	5.47	126.27	119.70
35	YB	90	C	N1-C2-O2	5.47	122.19	118.90

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	YA	1415	U	C6-N1-C1'	-5.47	113.54	121.20
1	QA	241	C	C2-N1-C1'	5.47	124.81	118.80
1	QA	788	U	C2-N1-C1'	5.47	124.26	117.70
1	XA	1478	C	C6-N1-C1'	-5.47	114.24	120.80
34	YA	752	A	P-O3'-C3'	5.47	126.26	119.70
34	RA	1483	G	C8-N9-C1'	-5.46	119.90	127.00
34	RA	1687	G	C4-N9-C1'	5.46	133.60	126.50
34	RA	2563	U	C2-N1-C1'	5.46	124.25	117.70
35	YB	94	C	N3-C4-C5	-5.46	119.72	121.90
34	RA	363(F)	U	C6-N1-C1'	-5.46	113.56	121.20
1	XA	651	C	C2-N1-C1'	5.46	124.80	118.80
1	XA	992	U	P-O3'-C3'	5.46	126.25	119.70
34	RA	619	G	C8-N9-C1'	-5.46	119.91	127.00
1	XA	947	G	C4-N9-C1'	5.46	133.59	126.50
34	RA	1314	C	C6-N1-C1'	-5.46	114.25	120.80
34	RA	1498	C	C2-N1-C1'	5.45	124.80	118.80
34	YA	435	C	C2-N1-C1'	5.45	124.80	118.80
1	QA	1440(L)	G	C4-N9-C1'	5.45	133.59	126.50
34	YA	2028	U	C2-N1-C1'	5.45	124.24	117.70
1	XA	795	C	C6-N1-C1'	-5.45	114.26	120.80
34	YA	267	C	C6-N1-C1'	-5.45	114.26	120.80
34	YA	848	G	C8-N9-C1'	-5.45	119.91	127.00
34	YA	2792	G	C4-N9-C1'	5.45	133.59	126.50
1	XA	436	C	C6-N1-C1'	-5.45	114.26	120.80
1	XA	916	G	C4-N9-C1'	5.45	133.58	126.50
1	XA	1373	G	OP1-P-OP2	-5.45	111.43	119.60
34	YA	1218	C	C2-N1-C1'	5.45	124.79	118.80
1	QA	456	C	C6-N1-C1'	-5.45	114.26	120.80
1	QA	1435	G	C4-N9-C1'	5.45	133.58	126.50
34	RA	219	G	C4-N9-C1'	5.45	133.58	126.50
34	RA	930	U	C2-N1-C1'	5.45	124.24	117.70
1	XA	153	C	C2-N1-C1'	5.45	124.79	118.80
22	QV	31	G	C8-N9-C1'	-5.44	119.92	127.00
34	YA	419	C	C6-N1-C1'	-5.44	114.27	120.80
34	YA	964	C	C2-N1-C1'	5.44	124.79	118.80
1	QA	1327	C	C2-N1-C1'	5.44	124.79	118.80
34	YA	919	G	C4-N9-C1'	5.44	133.58	126.50
1	QA	794	A	C8-N9-C1'	-5.44	117.91	127.70
50	RV	48	GLY	C-N-CA	5.44	135.30	121.70
1	XA	5	U	C2-N1-C1'	5.44	124.23	117.70
1	XA	1440(H)	C	C6-N1-C1'	-5.44	114.27	120.80
1	QA	656	C	C2-N1-C1'	5.44	124.78	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	YA	254	G	C4-N9-C1'	5.44	133.57	126.50
34	YA	1343	G	C4-N9-C1'	5.44	133.57	126.50
34	YA	2133	G	C8-N9-C1'	-5.44	119.93	127.00
34	RA	639	U	C6-N1-C1'	-5.43	113.59	121.20
34	RA	825	C	C6-N1-C1'	-5.43	114.28	120.80
34	YA	1930	G	P-O3'-C3'	5.43	126.22	119.70
34	RA	41	C	C2-N1-C1'	5.43	124.78	118.80
35	RB	43	C	C6-N1-C2	-5.43	118.13	120.30
1	XA	1100	C	C6-N1-C1'	-5.43	114.28	120.80
34	RA	1504	C	C2-N1-C1'	5.43	124.78	118.80
34	YA	1102	C	C2-N1-C1'	5.43	124.77	118.80
34	YA	2134	A	C4-N9-C1'	5.43	136.08	126.30
1	XA	20	U	C6-N1-C1'	-5.43	113.60	121.20
1	XA	137	C	C2-N1-C1'	5.43	124.77	118.80
34	YA	635	C	C6-N1-C1'	-5.43	114.28	120.80
35	YB	68	C	C5-C6-N1	5.43	123.71	121.00
34	YA	922	U	C2-N1-C1'	5.43	124.21	117.70
34	YA	2273	A	C4-N9-C1'	5.43	136.07	126.30
1	QA	709	G	C4-N9-C1'	5.42	133.55	126.50
12	XL	46	LYS	C-N-CA	-5.42	108.14	121.70
1	QA	906	G	C4-N9-C1'	5.42	133.55	126.50
34	YA	234	C	C2-N1-C1'	5.42	124.77	118.80
34	RA	2173	A	O5'-P-OP1	5.42	117.21	110.70
1	XA	1515	C	C2-N1-C1'	5.42	124.76	118.80
1	XA	68(E)	C	C2-N1-C1'	5.42	124.76	118.80
1	QA	372	C	C2-N1-C1'	5.42	124.76	118.80
34	RA	838	C	C2-N1-C1'	5.42	124.76	118.80
34	RA	1687	G	C8-N9-C1'	-5.42	119.95	127.00
1	XA	328	C	P-O3'-C3'	5.42	126.20	119.70
34	RA	191	A	C4-N9-C1'	5.42	136.05	126.30
35	YB	98	G	C8-N9-C4	-5.42	104.23	106.40
34	YA	2089	U	C6-N1-C1'	-5.42	113.62	121.20
34	RA	2648	C	C2-N1-C1'	5.41	124.75	118.80
1	XA	916	G	C8-N9-C1'	-5.41	119.96	127.00
1	QA	177	C	C6-N1-C1'	-5.41	114.31	120.80
34	RA	621	A	O4'-C1'-N9	5.41	112.53	108.20
34	RA	2299	G	C8-N9-C1'	-5.41	119.97	127.00
34	YA	75	G	C4-N9-C1'	5.41	133.54	126.50
34	YA	1549	C	C6-N1-C1'	-5.41	114.31	120.80
34	RA	76	C	C2-N1-C1'	5.41	124.75	118.80
1	XA	945	G	P-O3'-C3'	-5.41	113.21	119.70
34	YA	2175	C	C6-N1-C1'	-5.41	114.31	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	972	C	C2-N1-C1'	5.41	124.75	118.80
1	XA	524	G	C4-N9-C1'	5.41	133.53	126.50
34	YA	1418	G	C4-N9-C1'	5.41	133.53	126.50
34	YA	2532	G	C8-N9-C1'	-5.41	119.97	127.00
34	YA	1672	C	C6-N1-C1'	-5.41	114.31	120.80
34	RA	391	G	C4-N9-C1'	5.41	133.53	126.50
34	RA	628	G	C4-N9-C1'	5.41	133.53	126.50
34	RA	2299	G	C4-N9-C1'	5.41	133.53	126.50
1	XA	549	C	C2-N1-C1'	5.41	124.75	118.80
34	YA	407	G	C4-N9-C1'	5.41	133.53	126.50
34	YA	523	C	C2-N1-C1'	5.41	124.75	118.80
1	XA	858	G	C8-N9-C1'	-5.40	119.97	127.00
1	QA	1435	G	C8-N9-C1'	-5.40	119.98	127.00
1	XA	598	U	C6-N1-C1'	-5.40	113.64	121.20
1	XA	947	G	C8-N9-C1'	-5.40	119.98	127.00
34	RA	1951	U	C6-N1-C1'	-5.40	113.64	121.20
1	XA	1243	C	OP1-P-O3'	5.40	117.07	105.20
34	YA	2834	G	C4-N9-C1'	5.40	133.51	126.50
34	RA	1178	C	P-O3'-C3'	5.39	126.17	119.70
34	YA	994	C	C2-N1-C1'	5.39	124.73	118.80
34	YA	2072	G	C4-N9-C1'	5.39	133.51	126.50
1	XA	234	C	C2-N1-C1'	5.39	124.73	118.80
1	XA	1440(I)	U	C6-N1-C1'	-5.39	113.65	121.20
34	YA	1178	C	P-O3'-C3'	5.39	126.17	119.70
34	YA	1819	A	OP2-P-O3'	5.39	117.06	105.20
1	XA	169	C	C6-N1-C1'	-5.39	114.33	120.80
34	YA	1920	C	C6-N1-C1'	-5.39	114.33	120.80
48	YT	114	LEU	CA-CB-CG	5.39	127.70	115.30
1	QA	401	C	C2-N1-C1'	5.38	124.72	118.80
1	QA	834	C	C2-N1-C1'	5.38	124.72	118.80
34	YA	1992	G	OP2-P-O3'	5.38	117.05	105.20
34	RA	581	C	C6-N1-C1'	-5.38	114.34	120.80
34	YA	2667	C	C2-N1-C1'	5.38	124.72	118.80
34	RA	281	G	C4-N9-C1'	5.38	133.50	126.50
34	RA	2028	U	C6-N1-C1'	-5.38	113.67	121.20
34	RA	2745	C	C2-N1-C1'	5.38	124.72	118.80
34	YA	254	G	C8-N9-C1'	-5.38	120.00	127.00
1	QA	591	U	C2-N1-C1'	5.38	124.16	117.70
34	RA	1961	C	C2-N1-C1'	5.38	124.72	118.80
1	XA	862	C	C6-N1-C1'	-5.38	114.34	120.80
40	YH	155	SER	N-CA-C	5.38	125.53	111.00
1	XA	132	C	C6-N1-C1'	-5.38	114.35	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	524	G	C8-N9-C1'	-5.38	120.01	127.00
34	YA	887	A	O4'-C1'-N9	5.38	112.50	108.20
34	RA	628	G	C8-N9-C1'	-5.38	120.01	127.00
1	XA	739	C	C2-N1-C1'	5.38	124.71	118.80
34	YA	1188	U	C2-N1-C1'	5.38	124.15	117.70
1	QA	672	U	C6-N1-C1'	-5.38	113.67	121.20
1	QA	764	C	C2-N1-C1'	5.38	124.71	118.80
34	RA	219	G	C8-N9-C1'	-5.38	120.01	127.00
1	QA	805	C	C2-N1-C1'	5.37	124.71	118.80
1	XA	1075	C	C6-N1-C1'	-5.37	114.35	120.80
34	YA	102	G	P-O3'-C3'	5.37	126.15	119.70
34	YA	919	G	C8-N9-C1'	-5.37	120.02	127.00
1	QA	1225	A	O4'-C1'-N9	5.37	112.50	108.20
34	RA	1405	U	C2-N1-C1'	5.37	124.14	117.70
34	RA	2463	C	C6-N1-C1'	-5.37	114.36	120.80
34	YA	971	C	C2-N1-C1'	5.37	124.71	118.80
34	YA	1370	C	C2-N1-C1'	5.37	124.71	118.80
34	YA	1437	C	C6-N1-C1'	-5.37	114.36	120.80
34	YA	1675	C	C6-N1-C1'	-5.37	114.36	120.80
1	QA	930	C	C6-N1-C1'	-5.37	114.36	120.80
34	YA	1508	A	P-O3'-C3'	5.37	126.14	119.70
34	YA	2056	G	C4-N9-C1'	5.37	133.48	126.50
35	YB	75	G	C5-C6-O6	5.37	131.82	128.60
34	YA	1396	U	C2-N1-C1'	5.37	124.14	117.70
14	QN	43	CYS	CA-CB-SG	-5.37	104.34	114.00
34	RA	365	C	C6-N1-C1'	-5.37	114.36	120.80
34	RA	370	G	O4'-C1'-N9	-5.37	103.91	108.20
1	XA	335	C	C6-N1-C1'	-5.37	114.36	120.80
34	YA	858	U	C2-N1-C1'	5.36	124.14	117.70
34	YA	2792	G	C8-N9-C1'	-5.36	120.03	127.00
1	QA	709	G	C8-N9-C1'	-5.36	120.03	127.00
1	QA	906	G	C8-N9-C1'	-5.36	120.03	127.00
1	QA	1282	C	C6-N1-C1'	-5.36	114.36	120.80
14	XN	58	LYS	N-CA-C	-5.36	96.52	111.00
34	RA	833	U	C2-N1-C1'	5.36	124.13	117.70
34	YA	1343	G	C8-N9-C1'	-5.36	120.03	127.00
1	QA	955	U	C6-N1-C1'	-5.36	113.70	121.20
1	XA	1188	A	C4-N9-C1'	5.36	135.94	126.30
33	R9	32	HIS	CB-CA-C	5.36	121.11	110.40
34	RA	339	U	C6-N1-C1'	-5.36	113.70	121.20
1	XA	980	C	O3'-P-O5'	5.36	114.17	104.00
1	XA	1296	C	O4'-C1'-N1	5.36	112.48	108.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	1440(L)	G	C8-N9-C1'	-5.35	120.04	127.00
34	RA	2089	U	C2-N1-C1'	5.35	124.12	117.70
34	RA	191	A	C8-N9-C1'	-5.35	118.07	127.70
1	QA	945	G	C4-N9-C1'	5.35	133.45	126.50
22	QV	30	C	C6-N1-C1'	-5.35	114.38	120.80
34	RA	1636	C	C2-N1-C1'	5.35	124.68	118.80
1	XA	121	C	C6-N1-C1'	-5.35	114.38	120.80
1	XA	920	U	C6-N1-C1'	-5.35	113.71	121.20
1	XA	1160	G	C5'-C4'-O4'	5.35	115.52	109.10
1	QA	981	U	C6-N1-C1'	-5.35	113.71	121.20
34	RA	2441	C	C2-N1-C1'	5.35	124.68	118.80
34	YA	1537	C	C6-N1-C1'	-5.35	114.38	120.80
1	QA	1187	G	C4-N9-C1'	5.35	133.45	126.50
34	RA	427	U	C6-N1-C1'	-5.35	113.72	121.20
1	QA	8	A	OP1-P-O3'	5.34	116.96	105.20
1	QA	545	C	C6-N1-C1'	-5.34	114.39	120.80
1	QA	1515	C	C2-N1-C1'	5.34	124.68	118.80
1	XA	290	C	C2-N1-C1'	5.34	124.68	118.80
20	XT	10	LEU	CA-CB-CG	5.34	127.59	115.30
1	QA	479	C	C2-N1-C1'	5.34	124.68	118.80
1	QA	1028(G)	A	C4-N9-C1'	5.34	135.91	126.30
1	XA	564	C	C6-N1-C1'	-5.34	114.39	120.80
34	YA	1104	C	C2-N1-C1'	5.34	124.67	118.80
34	YA	2273	A	C8-N9-C1'	-5.34	118.09	127.70
34	YA	75	G	C8-N9-C1'	-5.34	120.06	127.00
34	RA	2166	G	C4-N9-C1'	5.33	133.44	126.50
34	RA	2392	A	O4'-C1'-N9	5.33	112.47	108.20
1	QA	593	G	C4-N9-C1'	5.33	133.43	126.50
34	RA	595	C	C2-N1-C1'	5.33	124.67	118.80
34	RA	658	C	C2-N1-C1'	5.33	124.67	118.80
35	RB	27	C	C2-N1-C1'	5.33	124.67	118.80
1	XA	1423	G	C4-N9-C1'	5.33	133.43	126.50
1	XA	1536	C	C2-N1-C1'	5.33	124.67	118.80
1	QA	251	G	OP1-P-O3'	5.33	116.93	105.20
1	QA	201(C)	U	C2-N1-C1'	5.33	124.09	117.70
34	RA	629	G	C4-N9-C1'	5.33	133.43	126.50
34	RA	2157	G	C4-N9-C1'	5.33	133.43	126.50
1	XA	950	U	C6-N1-C1'	-5.33	113.74	121.20
35	YB	110	G	N1-C6-O6	5.33	123.10	119.90
1	QA	519	C	C2-N1-C1'	5.33	124.66	118.80
34	RA	1271	G	C4-N9-C1'	5.33	133.42	126.50
1	XA	163	C	C6-N1-C1'	-5.33	114.41	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	789	U	C2-N1-C1'	5.33	124.09	117.70
34	YA	108	U	C2-N1-C1'	5.33	124.09	117.70
1	QA	1210	C	C2-N1-C1'	5.32	124.66	118.80
34	RA	2060	A	P-O3'-C3'	5.32	126.09	119.70
22	XV	56	C	C2-N1-C1'	5.32	124.66	118.80
22	XV	56	C	C6-N1-C2	-5.32	118.17	120.30
34	YA	2691	C	C6-N1-C1'	-5.32	114.41	120.80
1	QA	93	U	C2-N1-C1'	5.32	124.09	117.70
34	RA	1581	G	C4-N9-C1'	5.32	133.42	126.50
1	QA	21	G	C4-N9-C1'	5.32	133.42	126.50
1	QA	89	U	C2-N1-C1'	5.32	124.08	117.70
34	RA	1068	G	C4-N9-C1'	5.32	133.41	126.50
34	YA	796	C	C6-N1-C1'	-5.32	114.42	120.80
34	YA	1264	G	OP1-P-O3'	5.32	116.90	105.20
34	YA	1688	U	C2-N1-C1'	5.32	124.08	117.70
1	QA	17	U	C2-N1-C1'	5.32	124.08	117.70
34	YA	1418	G	C8-N9-C1'	-5.32	120.09	127.00
45	YQ	17	LEU	CB-CA-C	-5.32	100.10	110.20
1	XA	1354	C	C6-N1-C1'	-5.31	114.42	120.80
34	RA	1474	C	C6-N1-C1'	-5.31	114.42	120.80
1	XA	964	A	C4-N9-C1'	5.31	135.86	126.30
34	YA	2798	C	C2-N1-C1'	5.31	124.64	118.80
34	YA	534	U	C2-N1-C1'	5.31	124.07	117.70
34	RA	684	G	C4-N9-C1'	5.31	133.40	126.50
1	XA	1404	C	C2-N1-C1'	5.31	124.64	118.80
34	YA	407	G	C8-N9-C1'	-5.31	120.10	127.00
35	YB	49	C	N3-C4-C5	-5.31	119.78	121.90
1	QA	564	C	C6-N1-C1'	-5.31	114.43	120.80
1	QA	744	C	C2-N1-C1'	5.31	124.64	118.80
1	QA	174	C	C2-N1-C1'	5.30	124.64	118.80
34	RA	1670	C	C6-N1-C1'	-5.30	114.43	120.80
1	XA	631	G	C4-N9-C1'	5.30	133.40	126.50
1	XA	1105	A	O4'-C1'-N9	5.30	112.44	108.20
1	QA	786	G	C4-N9-C1'	5.30	133.39	126.50
1	XA	857	C	C6-N1-C1'	-5.30	114.44	120.80
34	YA	364	C	C6-N1-C1'	-5.30	114.44	120.80
34	YA	398	G	OP1-P-O3'	5.30	116.86	105.20
34	YA	553	U	C2-N1-C1'	5.30	124.06	117.70
34	YA	2081	C	C2-N1-C1'	5.30	124.63	118.80
34	YA	2403	C	C2-N1-C1'	5.30	124.63	118.80
34	YA	2099	U	C2-N1-C1'	5.30	124.06	117.70
34	RA	391	G	C8-N9-C1'	-5.30	120.11	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	RA	651	G	C8-N9-C1'	-5.30	120.11	127.00
1	QA	674	G	C4-N9-C1'	5.30	133.39	126.50
34	RA	1100	C	C2-N1-C1'	5.30	124.62	118.80
34	RA	1430	C	C2-N1-C1'	5.30	124.63	118.80
35	RB	47	C	N1-C2-O2	5.30	122.08	118.90
1	XA	878	G	O4'-C1'-N9	5.30	112.44	108.20
34	YA	2834	G	C8-N9-C1'	-5.30	120.11	127.00
1	QA	249	U	C6-N1-C1'	-5.29	113.79	121.20
1	QA	1187	G	C8-N9-C1'	-5.29	120.12	127.00
1	XA	1188	A	C8-N9-C1'	-5.29	118.17	127.70
1	XA	1423	G	C8-N9-C1'	-5.29	120.12	127.00
34	RA	155	C	C2-N1-C1'	5.29	124.62	118.80
34	RA	281	G	C8-N9-C1'	-5.29	120.12	127.00
34	RA	1121	C	C6-N1-C1'	-5.29	114.45	120.80
1	XA	1327	C	O4'-C1'-N1	5.29	112.43	108.20
34	YA	2072	G	C8-N9-C1'	-5.29	120.12	127.00
34	YA	2110	G	O4'-C1'-N9	-5.29	103.97	108.20
34	YA	2354	G	C4-N9-C1'	5.29	133.38	126.50
28	Y4	4	GLY	N-CA-C	5.29	126.33	113.10
1	QA	112	G	C4-N9-C1'	5.29	133.38	126.50
1	QA	1012	U	C2-N1-C1'	5.29	124.05	117.70
34	RA	1588	C	C2-N1-C1'	5.29	124.62	118.80
1	XA	252	U	C6-N1-C1'	-5.29	113.80	121.20
34	YA	1508	A	OP2-P-O3'	5.29	116.83	105.20
1	XA	831	U	C2-N1-C1'	5.28	124.04	117.70
34	YA	2056	G	C8-N9-C1'	-5.28	120.13	127.00
34	YA	2599	G	C4-N9-C1'	5.28	133.37	126.50
1	QA	593	G	C8-N9-C1'	-5.28	120.13	127.00
1	QA	754	C	N1-C1'-C2'	5.28	120.86	114.00
1	QA	1203	C	C2-N1-C1'	5.28	124.61	118.80
34	RA	1068	G	C8-N9-C1'	-5.28	120.14	127.00
35	RB	96	G	C8-N9-C4	-5.28	104.29	106.40
34	RA	2166	G	C8-N9-C1'	-5.28	120.14	127.00
34	YA	604	G	C4-N9-C1'	5.28	133.36	126.50
35	YB	29	A	N7-C8-N9	5.28	116.44	113.80
1	QA	217	C	C2-N1-C1'	5.28	124.60	118.80
34	RA	1113	U	C2-N1-C1'	5.28	124.03	117.70
34	RA	2157	G	C8-N9-C1'	-5.28	120.14	127.00
1	XA	1281	U	C2-N1-C1'	5.28	124.03	117.70
34	YA	2162	G	C4-N9-C1'	5.27	133.36	126.50
34	YA	2586	C	C6-N1-C1'	-5.27	114.47	120.80
35	YB	78	A	C8-N9-C4	-5.27	103.69	105.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	1404	C	C2-N1-C1'	5.27	124.60	118.80
34	RA	1201	C	C2-N1-C1'	5.27	124.60	118.80
1	XA	1347	G	N9-C1'-C2'	-5.27	106.20	112.00
34	YA	1105	U	C2-N1-C1'	5.27	124.03	117.70
1	QA	221	C	C2-N1-C1'	5.27	124.60	118.80
1	QA	455	C	C2-N1-C1'	5.27	124.60	118.80
34	RA	267	C	C2-N1-C1'	5.27	124.60	118.80
1	QA	824	C	C6-N1-C1'	-5.27	114.48	120.80
1	QA	1237	C	C6-N1-C1'	-5.27	114.48	120.80
34	RA	1581	G	C8-N9-C1'	-5.27	120.15	127.00
22	QV	4	U	C2-N1-C1'	5.26	124.02	117.70
34	YA	678	C	C6-N1-C1'	-5.26	114.48	120.80
1	QA	786	G	C8-N9-C1'	-5.26	120.16	127.00
1	QA	945	G	C8-N9-C1'	-5.26	120.16	127.00
1	XA	1108	G	C8-N9-C1'	-5.26	120.16	127.00
35	YB	38	C	N3-C2-O2	-5.26	118.22	121.90
54	YZ	151	HIS	N-CA-C	5.26	125.21	111.00
34	YA	271(E)	G	C4-N9-C1'	5.26	133.34	126.50
1	XA	222	U	C6-N1-C1'	-5.26	113.84	121.20
34	YA	604	G	C8-N9-C1'	-5.26	120.17	127.00
34	YA	2359	C	C2-N1-C1'	5.26	124.58	118.80
1	XA	1440(K)	C	C2-N1-C1'	5.26	124.58	118.80
34	YA	2779	U	C2-N1-C1'	5.26	124.01	117.70
1	QA	556	C	C6-N1-C1'	-5.25	114.49	120.80
35	RB	60	C	C6-N1-C2	-5.25	118.20	120.30
34	YA	2354	G	C8-N9-C1'	-5.25	120.17	127.00
34	RA	684	G	C8-N9-C1'	-5.25	120.17	127.00
34	RA	2061	G	O5'-P-OP2	-5.25	100.97	105.70
1	XA	849	C	C2-N1-C1'	5.25	124.58	118.80
34	YA	1577	C	C6-N1-C1'	-5.25	114.50	120.80
34	YA	2128	C	C2-N1-C1'	5.25	124.58	118.80
1	QA	1402	C	C2-N1-C1'	5.25	124.58	118.80
1	QA	1514	C	C2-N1-C1'	5.25	124.58	118.80
34	RA	104	U	C2-N1-C1'	5.25	124.00	117.70
34	RA	1180	C	C6-N1-C1'	-5.25	114.50	120.80
34	RA	2610	C	P-O3'-C3'	5.25	126.00	119.70
34	RA	2832	U	C2-N1-C1'	5.25	123.99	117.70
1	XA	186(F)	C	C2-N1-C1'	5.24	124.57	118.80
34	YA	318	C	C6-N1-C1'	-5.24	114.51	120.80
1	QA	370	C	C6-N1-C1'	-5.24	114.51	120.80
1	XA	744	C	C2-N1-C1'	5.24	124.57	118.80
1	XA	1364	U	C6-N1-C1'	-5.24	113.86	121.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	RB	28	C	C5-C6-N1	5.24	123.62	121.00
34	YA	2438	U	C2-N1-C1'	5.24	123.99	117.70
34	RA	613	U	O4'-C1'-N1	5.24	112.39	108.20
34	RA	1271	G	C8-N9-C1'	-5.24	120.19	127.00
1	XA	617	G	P-O3'-C3'	5.24	125.98	119.70
1	XA	631	G	C8-N9-C1'	-5.24	120.19	127.00
1	XA	1186	G	O4'-C1'-N9	-5.24	104.01	108.20
34	RA	198	C	C2-N1-C1'	5.23	124.56	118.80
34	RA	2665	A	O4'-C1'-N9	5.23	112.39	108.20
1	XA	342	C	C2-N1-C1'	5.23	124.56	118.80
34	YA	794	G	C4-N9-C1'	5.23	133.30	126.50
34	YA	2437	U	C6-N1-C1'	-5.23	113.88	121.20
1	XA	1108	G	C4-N9-C1'	5.23	133.30	126.50
22	XV	15	G	N7-C8-N9	5.23	115.72	113.10
34	YA	1556	C	C6-N1-C1'	-5.23	114.53	120.80
1	QA	501	C	C2-N1-C1'	5.23	124.55	118.80
1	XA	1317	C	C6-N1-C1'	-5.23	114.53	120.80
34	YA	1782	C	O5'-P-OP2	5.23	116.97	110.70
35	RB	82	G	N1-C6-O6	5.22	123.03	119.90
34	YA	2455	G	C4-N9-C1'	5.22	133.29	126.50
1	QA	261	U	C6-N1-C1'	-5.22	113.89	121.20
34	RA	270(D)	C	C2-N1-C1'	5.22	124.55	118.80
34	RA	2064	C	C6-N1-C1'	-5.22	114.53	120.80
1	XA	1300	G	O4'-C1'-N9	5.22	112.38	108.20
1	XA	1533	C	C2-N1-C1'	5.22	124.55	118.80
34	YA	1544	C	C2-N1-C1'	5.22	124.55	118.80
1	QA	112	G	C8-N9-C1'	-5.22	120.21	127.00
34	RA	2264	C	C2-N1-C1'	5.22	124.54	118.80
1	QA	358	U	C6-N1-C1'	-5.22	113.89	121.20
34	RA	1153	C	C2-N1-C1'	5.22	124.54	118.80
1	QA	21	G	C8-N9-C1'	-5.22	120.22	127.00
1	QA	1028(G)	A	C8-N9-C1'	-5.22	118.31	127.70
1	QA	1381	U	C2-N1-C1'	5.21	123.96	117.70
34	RA	629	G	C8-N9-C1'	-5.21	120.22	127.00
34	RA	2477	C	O5'-P-OP1	-5.21	101.01	105.70
34	YA	684	G	C4-N9-C1'	5.21	133.28	126.50
34	YA	976	C	C2-N1-C1'	5.21	124.54	118.80
34	YA	235	U	C2-N1-C1'	5.21	123.96	117.70
34	YA	2189	U	C2-N1-C1'	5.21	123.96	117.70
34	YA	1588	C	C6-N1-C1'	-5.21	114.55	120.80
35	YB	108	C	C6-N1-C2	5.21	122.38	120.30
34	RA	678	C	C6-N1-C1'	-5.21	114.55	120.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
54	RZ	178	GLU	C-N-CA	5.21	134.72	121.70
1	QA	1028(I)	G	C4-N9-C1'	5.21	133.27	126.50
34	RA	1556	C	C2-N1-C1'	5.21	124.53	118.80
1	XA	513	C	C6-N1-C1'	-5.21	114.55	120.80
1	XA	1508	G	C8-N9-C1'	-5.21	120.23	127.00
34	YA	653	A	P-O3'-C3'	5.21	125.95	119.70
34	RA	1982	C	C6-N1-C1'	-5.21	114.55	120.80
1	XA	1322	C	C2-N1-C1'	5.21	124.53	118.80
22	XV	31	G	C4-N9-C1'	5.21	133.27	126.50
1	QA	1414	U	C6-N1-C1'	-5.21	113.91	121.20
34	RA	884	C	C2-N1-C1'	5.21	124.53	118.80
1	XA	1078	U	C2-N1-C1'	5.20	123.94	117.70
34	YA	1026	U	OP1-P-O3'	5.20	116.65	105.20
34	YA	1411	C	C6-N1-C1'	-5.20	114.56	120.80
34	RA	1349	A	O4'-C1'-N9	5.20	112.36	108.20
34	YA	312	G	C4-N9-C1'	5.20	133.26	126.50
34	YA	1514	U	C6-N1-C1'	-5.20	113.92	121.20
34	YA	2452	C	C6-N1-C1'	-5.20	114.56	120.80
35	YB	17	C	N1-C2-O2	5.20	122.02	118.90
35	YB	49	C	C6-N1-C2	-5.20	118.22	120.30
1	QA	546	G	P-O3'-C3'	5.20	125.94	119.70
34	RA	965	C	C2-N1-C1'	5.20	124.52	118.80
53	YY	79	CYS	N-CA-CB	5.20	119.95	110.60
34	RA	66	C	C6-N1-C1'	-5.20	114.56	120.80
1	XA	1406	U	C2-N1-C1'	5.20	123.93	117.70
34	YA	2599	G	C8-N9-C1'	-5.20	120.25	127.00
34	YA	2611	U	C2-N1-C1'	5.20	123.94	117.70
34	YA	2495	G	OP1-P-O3'	5.19	116.62	105.20
34	YA	2874	C	C2-N1-C1'	5.19	124.51	118.80
35	YB	54	G	N7-C8-N9	5.19	115.70	113.10
1	XA	1132	C	C2-N1-C1'	5.19	124.51	118.80
34	YA	639	U	C2-N1-C1'	5.19	123.93	117.70
34	RA	230	U	C2-N1-C1'	5.19	123.92	117.70
34	YA	2137	C	C2-N1-C1'	5.19	124.51	118.80
1	QA	674	G	C8-N9-C1'	-5.19	120.26	127.00
1	XA	352	C	C6-N1-C1'	-5.18	114.58	120.80
1	XA	1142	G	C4-N9-C1'	5.18	133.24	126.50
34	YA	271(E)	G	C8-N9-C1'	-5.18	120.26	127.00
34	YA	1707	G	C4-N9-C1'	5.18	133.24	126.50
34	YA	2085	C	C2-N1-C1'	5.18	124.50	118.80
1	XA	981	U	O5'-P-OP1	-5.18	101.04	105.70
1	QA	8	A	P-O3'-C3'	5.18	125.92	119.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	176	C	C6-N1-C1'	-5.18	114.58	120.80
34	RA	2452	C	C2-N1-C1'	5.18	124.50	118.80
34	RA	2805	G	C4-N9-C1'	5.18	133.23	126.50
1	XA	1232	U	C2-N1-C1'	5.18	123.92	117.70
1	QA	1261	A	C4-N9-C1'	5.18	135.62	126.30
1	XA	724	G	C4-N9-C1'	5.17	133.23	126.50
34	RA	1110	G	C4-N9-C1'	5.17	133.22	126.50
1	XA	1508	G	C4-N9-C1'	5.17	133.22	126.50
34	YA	1289	C	C6-N1-C1'	-5.17	114.59	120.80
34	YA	2162	G	C8-N9-C1'	-5.17	120.28	127.00
34	RA	1544	C	C2-N1-C1'	5.17	124.49	118.80
34	YA	624	C	C2-N1-C1'	5.17	124.49	118.80
1	XA	1206	G	C4-N9-C1'	5.17	133.22	126.50
34	YA	1406	U	C2-N1-C1'	5.17	123.90	117.70
34	YA	1734	C	C6-N1-C1'	-5.17	114.60	120.80
34	RA	1956	U	C2-N1-C1'	5.17	123.90	117.70
34	RA	2195	C	C2-N1-C1'	5.17	124.48	118.80
34	YA	480	A	C4-N9-C1'	5.17	135.60	126.30
35	RB	59	A	N1-C2-N3	-5.16	126.72	129.30
1	QA	851	G	C4-N9-C1'	5.16	133.21	126.50
1	QA	1157	A	OP2-P-O3'	5.16	116.56	105.20
1	XA	1157	A	C3'-C2'-C1'	5.16	105.63	101.50
1	XA	1348	U	O5'-C5'-C4'	5.16	121.51	111.70
34	YA	1313	U	C2-N1-C1'	5.16	123.89	117.70
34	RA	2099	U	C2-N1-C1'	5.16	123.89	117.70
1	XA	545	C	C2-N1-C1'	5.16	124.48	118.80
1	XA	617	G	C1'-C2'-O2'	5.16	126.08	110.60
34	YA	2179	C	C6-N1-C1'	-5.16	114.61	120.80
36	YD	243	GLY	N-CA-C	5.16	126.00	113.10
34	RA	364	C	C2-N1-C1'	5.16	124.47	118.80
34	RA	651	G	C4-N9-C1'	5.16	133.21	126.50
34	RA	1298	C	C6-N1-C1'	-5.16	114.61	120.80
1	XA	851	G	C4-N9-C1'	5.16	133.21	126.50
1	XA	1219	U	OP1-P-O3'	-5.16	93.85	105.20
1	QA	1027	C	C2-N1-C1'	5.16	124.47	118.80
34	YA	657	U	C6-N1-C1'	-5.16	113.98	121.20
1	XA	1359	C	O4'-C1'-N1	5.16	112.32	108.20
34	YA	202	U	C6-N1-C1'	-5.16	113.98	121.20
34	YA	684	G	C8-N9-C1'	-5.16	120.30	127.00
1	XA	1157	A	OP2-P-O3'	5.15	116.53	105.20
34	YA	208	C	C6-N1-C1'	-5.15	114.62	120.80
34	YA	395	U	C6-N1-C1'	-5.15	113.99	121.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	YA	508	G	C4-N9-C1'	5.15	133.20	126.50
34	YA	2455	G	C8-N9-C1'	-5.15	120.30	127.00
53	RY	99	CYS	CB-CA-C	-5.15	100.10	110.40
34	YA	312	G	C8-N9-C1'	-5.15	120.31	127.00
1	QA	178	C	C2-N1-C1'	5.15	124.46	118.80
1	QA	1440(I)	U	C2-N1-C1'	5.15	123.88	117.70
34	RA	634	C	C6-N1-C1'	-5.15	114.62	120.80
34	YA	94	G	C4-N9-C1'	5.15	133.19	126.50
34	YA	1303	G	C4-N9-C1'	5.15	133.19	126.50
35	YB	69	G	C8-N9-C4	-5.15	104.34	106.40
1	QA	252	U	C6-N1-C1'	-5.15	114.00	121.20
1	QA	262	A	C4-N9-C1'	5.15	135.56	126.30
34	RA	2466	C	C6-N1-C1'	-5.14	114.63	120.80
36	RD	241	PRO	C-N-CA	5.14	134.55	121.70
34	YA	2752	C	C6-N1-C1'	-5.14	114.63	120.80
35	YB	86	G	C5-N7-C8	5.14	106.87	104.30
1	QA	851	G	C8-N9-C1'	-5.14	120.32	127.00
34	RA	319	C	C2-N1-C1'	5.14	124.45	118.80
34	RA	1510	A	P-O5'-C5'	5.14	129.12	120.90
34	RA	1674	G	C4-N9-C1'	5.14	133.18	126.50
1	XA	1142	G	C8-N9-C1'	-5.14	120.32	127.00
1	XA	1320	C	O4'-C4'-C3'	-5.14	98.86	104.00
34	YA	923	C	C6-N1-C1'	-5.14	114.63	120.80
1	QA	295	C	C6-N1-C1'	-5.14	114.64	120.80
34	RA	755	C	C2-N1-C1'	5.14	124.45	118.80
1	XA	1349	A	O5'-P-OP1	-5.14	101.08	105.70
30	Y6	13	CYS	CA-CB-SG	-5.14	104.75	114.00
34	YA	2084	C	C6-N1-C1'	-5.14	114.64	120.80
50	YV	35	LEU	CA-CB-CG	5.14	127.11	115.30
1	QA	1322	C	C6-N1-C1'	-5.13	114.64	120.80
34	RA	929	G	C4-N9-C1'	5.13	133.17	126.50
1	XA	564	C	C2-N1-C1'	5.13	124.45	118.80
1	XA	832	C	C6-N1-C1'	-5.13	114.64	120.80
34	YA	271(B)	C	C6-N1-C1'	-5.13	114.64	120.80
34	YA	1417	C	C6-N1-C1'	-5.13	114.64	120.80
34	YA	1707	G	C8-N9-C1'	-5.13	120.33	127.00
32	R8	28	GLY	N-CA-C	5.13	125.92	113.10
1	XA	1347	G	O5'-C5'-C4'	5.13	121.44	111.70
34	YA	658	C	C2-N1-C1'	5.13	124.44	118.80
34	YA	794	G	C8-N9-C1'	-5.13	120.33	127.00
34	YA	866	A	C4-N9-C1'	5.13	135.53	126.30
34	YA	2013	A	OP1-P-O3'	5.13	116.48	105.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
35	YB	27	C	C2-N1-C1'	5.13	124.44	118.80
34	RA	1005	C	C6-N1-C1'	-5.13	114.65	120.80
34	RA	1674	G	C8-N9-C1'	-5.13	120.34	127.00
35	RB	90	C	C5-C6-N1	5.13	123.56	121.00
1	XA	1306	A	P-O3'-C3'	-5.13	113.55	119.70
34	RA	1135	C	C2-N1-C1'	5.12	124.44	118.80
1	XA	724	G	C8-N9-C1'	-5.12	120.34	127.00
1	QA	97	U	C6-N1-C1'	-5.12	114.03	121.20
35	RB	15	A	N9-C4-C5	-5.12	103.75	105.80
1	XA	288	A	C8-N9-C1'	-5.12	118.48	127.70
34	YA	18	C	C6-N1-C1'	-5.12	114.65	120.80
1	QA	234	C	C2-N1-C1'	5.12	124.43	118.80
1	QA	1028(I)	G	C8-N9-C1'	-5.12	120.34	127.00
34	RA	2805	G	C8-N9-C1'	-5.12	120.34	127.00
35	RB	55	U	C5-C6-N1	5.12	125.26	122.70
1	XA	988	G	C4-N9-C1'	5.12	133.16	126.50
20	XT	72	LEU	CA-CB-CG	5.12	127.08	115.30
35	YB	44	G	C5-C6-N1	5.12	114.06	111.50
1	QA	28	G	C4-N9-C1'	5.12	133.16	126.50
34	YA	94	G	C8-N9-C1'	-5.12	120.34	127.00
1	XA	1306	A	N9-C1'-C2'	5.12	120.66	114.00
34	YA	2540	C	C2-N1-C1'	5.12	124.43	118.80
1	QA	1261	A	C8-N9-C1'	-5.12	118.49	127.70
34	RA	1112	G	C4-N9-C1'	5.12	133.15	126.50
1	XA	288	A	C4-N9-C1'	5.12	135.51	126.30
34	RA	814	C	C2-N1-C1'	5.12	124.43	118.80
34	YA	270(D)	C	C6-N1-C1'	-5.12	114.66	120.80
35	YB	1	U	C2-N1-C1'	5.12	123.84	117.70
34	RA	2443	C	C2-N1-C1'	5.11	124.42	118.80
35	RB	106	G	C8-N9-C4	-5.11	104.36	106.40
1	XA	312	C	C6-N1-C1'	-5.11	114.66	120.80
34	YA	1479	G	C4-N9-C1'	5.11	133.15	126.50
1	QA	797	C	C2-N1-C1'	5.11	124.42	118.80
1	XA	1028(C)	C	C2-N1-C1'	5.11	124.42	118.80
1	QA	262	A	C8-N9-C1'	-5.11	118.50	127.70
1	XA	1532	U	C2-N1-C1'	5.11	123.83	117.70
34	RA	1579	A	C4-N9-C1'	5.11	135.50	126.30
1	XA	1098	C	C6-N1-C1'	-5.11	114.67	120.80
35	YB	30	C	N3-C4-C5	-5.11	119.86	121.90
29	R5	34	PRO	N-CA-CB	5.10	109.42	103.30
34	RA	1101	U	C6-N1-C1'	-5.10	114.06	121.20
34	YA	1579	A	C4-N9-C1'	5.10	135.49	126.30

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	RA	305	U	C2-N1-C1'	5.10	123.82	117.70
34	RA	351	G	C8-N9-C1'	-5.10	120.37	127.00
34	RA	1026	U	OP1-P-O3'	5.10	116.42	105.20
34	RA	2895	U	C2-N1-C1'	5.10	123.82	117.70
35	RB	27	C	N3-C2-O2	-5.10	118.33	121.90
1	XA	749	C	C6-N1-C1'	-5.10	114.68	120.80
35	YB	102	G	C5-C6-N1	5.10	114.05	111.50
1	QA	186(C)	C	C2-N1-C1'	5.10	124.41	118.80
34	YA	851	U	C2-N1-C1'	5.10	123.82	117.70
1	QA	856	C	O4'-C1'-N1	5.10	112.28	108.20
1	QA	962	C	C6-N1-C1'	-5.10	114.68	120.80
34	RA	2785	C	C6-N1-C1'	-5.10	114.68	120.80
34	YA	1072	C	C2-N1-C1'	-5.10	113.19	118.80
35	YB	61	G	N7-C8-N9	5.10	115.65	113.10
1	XA	1206	G	C8-N9-C1'	-5.10	120.37	127.00
34	YA	1240	U	C6-N1-C1'	-5.10	114.06	121.20
38	YF	133	ASN	N-CA-C	-5.10	97.24	111.00
34	YA	2229	C	C2-N1-C1'	5.09	124.40	118.80
1	QA	186(G)	C	C2-N1-C1'	5.09	124.40	118.80
1	QA	1256	A	O4'-C1'-N9	5.09	112.27	108.20
34	RA	29	U	C2-N1-C1'	5.09	123.81	117.70
35	RB	28	C	C6-N1-C2	-5.09	118.26	120.30
34	RA	915	C	C2-N1-C1'	5.09	124.40	118.80
34	RA	1265	A	OP1-P-OP2	-5.09	111.97	119.60
34	YA	1303	G	C8-N9-C1'	-5.09	120.38	127.00
34	YA	1869	G	C4-N9-C1'	5.09	133.12	126.50
35	YB	44	G	C4-N9-C1'	-5.09	119.88	126.50
34	YA	2462	U	C2-N1-C1'	5.09	123.80	117.70
35	YB	51	G	N1-C2-N3	5.09	126.95	123.90
1	QA	311	C	C2-N1-C1'	5.08	124.39	118.80
1	XA	1305	G	C1'-C2'-O2'	-5.08	95.34	110.60
34	YA	189	G	C4-N9-C1'	5.08	133.11	126.50
1	QA	1244	C	C2-N1-C1'	5.08	124.39	118.80
34	RA	2217	G	C4-N9-C1'	5.08	133.11	126.50
34	RA	2351	G	C8-N9-C1'	-5.08	120.39	127.00
34	YA	2884	U	C6-N1-C1'	-5.08	114.08	121.20
34	RA	509	C	C6-N1-C1'	-5.08	114.70	120.80
1	XA	848	C	C6-N1-C1'	-5.08	114.70	120.80
34	YA	1050	A	O4'-C1'-N9	5.08	112.27	108.20
34	YA	1108	U	C6-N1-C1'	-5.08	114.09	121.20
1	QA	1316	G	C8-N9-C1'	-5.08	120.40	127.00
34	RA	318	C	C2-N1-C1'	5.08	124.39	118.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
34	RA	2351	G	C4-N9-C1'	5.08	133.10	126.50
35	YB	31	C	C6-N1-C1'	-5.08	114.70	120.80
34	RA	99	U	P-O3'-C3'	5.08	125.80	119.70
34	RA	1110	G	C8-N9-C1'	-5.08	120.40	127.00
34	RA	1174	A	C4-N9-C1'	5.08	135.44	126.30
34	YA	1516	U	C6-N1-C1'	-5.08	114.09	121.20
34	RA	120	U	C2-N1-C1'	5.08	123.79	117.70
54	RZ	12	GLY	N-CA-C	-5.08	100.41	113.10
34	YA	508	G	C8-N9-C1'	-5.08	120.40	127.00
34	YA	2093	G	C4-N9-C1'	5.07	133.09	126.50
1	QA	1115	C	C2-N1-C1'	5.07	124.38	118.80
1	QA	1352	C	C6-N1-C1'	-5.07	114.72	120.80
1	XA	851	G	C8-N9-C1'	-5.07	120.41	127.00
34	YA	690	G	C4-N9-C1'	5.07	133.09	126.50
1	XA	456	C	C2-N1-C1'	5.07	124.38	118.80
22	XV	56	C	N3-C2-O2	-5.07	118.35	121.90
35	YB	21	G	C8-N9-C4	-5.07	104.37	106.40
1	QA	192	U	C6-N1-C1'	-5.07	114.11	121.20
34	YA	1178	C	C2-N1-C1'	5.07	124.37	118.80
1	XA	1220	G	O5'-P-OP1	-5.07	101.14	105.70
34	RA	929	G	C8-N9-C1'	-5.06	120.42	127.00
34	RA	2044	C	C6-N1-C1'	-5.06	114.73	120.80
34	RA	2420	C	C6-N1-C1'	-5.06	114.72	120.80
1	XA	526	C	C6-N1-C1'	-5.06	114.72	120.80
1	QA	1158	C	O4'-C1'-N1	5.06	112.25	108.20
37	RE	146	THR	N-CA-C	5.06	124.66	111.00
1	XA	400	C	C6-N1-C1'	-5.06	114.73	120.80
1	XA	1247	U	C2-N1-C1'	5.06	123.77	117.70
34	YA	1252	G	O4'-C1'-N9	-5.06	104.15	108.20
34	YA	2730	C	C6-N1-C1'	-5.06	114.73	120.80
1	QA	14	U	C2-N1-C1'	5.06	123.77	117.70
1	QA	905	U	C6-N1-C1'	-5.06	114.12	121.20
34	RA	1257	C	C2-N1-C1'	5.06	124.36	118.80
34	RA	1289	C	C2-N1-C1'	5.06	124.36	118.80
34	YA	480	A	C8-N9-C1'	-5.06	118.59	127.70
1	QA	281	G	C4-N9-C1'	5.06	133.07	126.50
34	RA	1927	A	C4-N9-C1'	5.06	135.40	126.30
34	RA	2441	C	C6-N1-C1'	-5.06	114.73	120.80
34	YA	1432	C	C2-N1-C1'	5.06	124.36	118.80
34	YA	1967	C	C2-N1-C1'	5.06	124.36	118.80
1	QA	849	C	C6-N1-C1'	-5.06	114.73	120.80
34	YA	813	U	C2-N1-C1'	5.06	123.77	117.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	QA	920	U	C6-N1-C1'	-5.05	114.12	121.20
34	RA	1578	U	C2-N1-C1'	5.05	123.77	117.70
34	RA	1680	U	C2-N1-C1'	5.05	123.77	117.70
40	RH	155	SER	N-CA-C	5.05	124.65	111.00
1	QA	28	G	C8-N9-C1'	-5.05	120.43	127.00
1	XA	581	G	C4-N9-C1'	5.05	133.07	126.50
1	QA	12	U	C2-N1-C1'	5.05	123.76	117.70
34	RA	704	G	C4-N9-C1'	5.05	133.07	126.50
22	XV	28	U	C5-C6-N1	5.05	125.23	122.70
34	YA	1633	G	C4-N9-C1'	5.05	133.06	126.50
34	YA	2562	U	C2-N1-C1'	5.05	123.76	117.70
34	YA	1869	G	C8-N9-C1'	-5.05	120.44	127.00
34	RA	2739	U	C6-N1-C1'	-5.04	114.14	121.20
1	XA	988	G	C8-N9-C1'	-5.04	120.44	127.00
34	YA	2101	G	C4-N9-C1'	5.04	133.06	126.50
35	YB	22	U	C6-N1-C2	-5.04	117.97	121.00
1	QA	1303	C	C6-N1-C1'	-5.04	114.75	120.80
34	RA	1112	G	C8-N9-C1'	-5.04	120.45	127.00
34	YA	74	A	O4'-C1'-N9	-5.04	104.17	108.20
34	YA	1515	C	C6-N1-C1'	-5.04	114.75	120.80
1	QA	647	C	C6-N1-C1'	-5.04	114.75	120.80
1	QA	678	U	C6-N1-C1'	-5.04	114.15	121.20
34	RA	2144	U	C6-N1-C1'	-5.04	114.15	121.20
1	XA	25	C	C6-N1-C1'	-5.04	114.76	120.80
1	QA	1388	C	C6-N1-C1'	-5.03	114.76	120.80
34	RA	2226	C	C2-N1-C1'	5.03	124.33	118.80
34	YA	866	A	C8-N9-C1'	-5.03	118.64	127.70
1	XA	479	C	C6-N1-C1'	-5.03	114.77	120.80
1	XA	1349	A	N9-C1'-C2'	5.03	120.54	114.00
34	YA	2093	G	C8-N9-C1'	-5.03	120.46	127.00
1	QA	855	G	C4-N9-C1'	5.03	133.04	126.50
34	RA	435	C	C6-N1-C1'	-5.03	114.77	120.80
34	RA	1579	A	C8-N9-C1'	-5.03	118.65	127.70
1	QA	1316	G	C4-N9-C1'	5.03	133.03	126.50
1	XA	1440(K)	C	C6-N1-C1'	-5.03	114.77	120.80
34	YA	568	U	C6-N1-C1'	-5.03	114.16	121.20
34	YA	690	G	C8-N9-C1'	-5.03	120.47	127.00
34	YA	1479	G	C8-N9-C1'	-5.03	120.47	127.00
34	YA	1579	A	C8-N9-C1'	-5.03	118.65	127.70
34	RA	652	C	C2-N1-C1'	5.02	124.33	118.80
34	YA	649	G	C4-N9-C1'	5.02	133.03	126.50
34	RA	2217	G	C8-N9-C1'	-5.02	120.47	127.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	XA	1243	C	C2-N1-C1'	5.02	124.32	118.80
54	YZ	63	ASP	CB-CG-OD1	5.02	122.82	118.30
34	RA	351	G	C4-N9-C1'	5.02	133.02	126.50
1	XA	1170	A	C4-N9-C1'	5.02	135.34	126.30
34	YA	189	G	C8-N9-C1'	-5.02	120.47	127.00
1	QA	998(A)	G	C4-N9-C1'	5.02	133.02	126.50
1	XA	1402	C	C2-N1-C1'	5.02	124.32	118.80
34	YA	539	G	C4-N9-C1'	5.02	133.02	126.50
34	YA	1924	C	C2-N1-C1'	5.02	124.32	118.80
34	RA	838	C	C6-N1-C1'	-5.02	114.78	120.80
34	YA	1218	C	C6-N1-C1'	-5.02	114.78	120.80
34	RA	201	C	C6-N1-C1'	-5.01	114.78	120.80
34	RA	2292	C	C2-N1-C1'	5.01	124.32	118.80
1	XA	860	A	O4'-C1'-N9	5.01	112.21	108.20
1	QA	883	C	C2-N1-C1'	5.01	124.31	118.80
1	XA	1371	G	C4-N9-C1'	5.01	133.01	126.50
1	QA	372	C	C6-N1-C1'	-5.01	114.79	120.80
34	RA	1136	G	C4-N9-C1'	5.01	133.01	126.50
1	XA	1056	U	C6-N1-C1'	-5.01	114.19	121.20
34	YA	1633	G	C8-N9-C1'	-5.01	120.49	127.00
34	YA	2101	G	C8-N9-C1'	-5.01	120.49	127.00
32	R8	62	LEU	N-CA-C	5.01	124.52	111.00
34	RA	1304	C	C6-N1-C1'	-5.01	114.79	120.80
35	YB	78	A	N9-C4-C5	5.01	107.80	105.80
34	RA	856	C	C2'-C3'-O3'	5.00	121.71	113.70
34	RA	1174	A	C8-N9-C1'	-5.00	118.69	127.70
34	RA	2086	U	C2-N1-C1'	5.00	123.70	117.70
1	XA	1347	G	C2'-C3'-O3'	-5.00	98.49	109.50
34	YA	365	C	C6-N1-C1'	-5.00	114.80	120.80
1	QA	352	C	C2-N1-C1'	5.00	124.30	118.80
34	RA	806	C	C6-N1-C1'	-5.00	114.80	120.80
35	RB	91	C	N3-C4-C5	-5.00	119.90	121.90
1	XA	930	C	C2-N1-C1'	5.00	124.30	118.80
23	XX	15	A	N9-C4-C5	-5.00	103.80	105.80
34	YA	1926	U	C2-N1-C1'	5.00	123.70	117.70
34	YA	2271	G	C4-N9-C1'	5.00	133.00	126.50

All (2) chirality outliers are listed below:

Mol	Chain	Res	Type	Atom
1	XA	617	G	C2'
28	Y4	5	ILE	CA

All (30) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
34	RA	1649	G	Sidechain
37	RE	146	THR	Peptide
50	RV	49	THR	Mainchain,Peptide
54	RZ	166	SER	Peptide
1	XA	1112	C	Sidechain
1	XA	1225	A	Sidechain
1	XA	1320	C	Sidechain
1	XA	1348	U	Sidechain
1	XA	1507	A	Sidechain
1	XA	308	C	Sidechain
1	XA	564	C	Sidechain
1	XA	608	A	Sidechain
1	XA	618	C	Sidechain
1	XA	625	G	Sidechain
1	XA	635	G	Sidechain
1	XA	860	A	Sidechain
1	XA	878	G	Sidechain
1	XA	957	U	Sidechain
1	XA	964	A	Sidechain
1	XA	972	C	Sidechain
28	Y4	5	ILE	Mainchain
34	YA	2060	A	Sidechain
34	YA	2098	U	Sidechain
34	YA	2111	C	Sidechain
34	YA	2454	G	Sidechain
34	YA	2459	A	Sidechain
34	YA	2550	G	Sidechain
37	YE	146	THR	Peptide
54	YZ	166	SER	Peptide

5.2 Too-close contacts

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	QA	32469	0	16385	1571	0
1	XA	32551	0	16418	1288	0
2	QB	1907	0	1958	99	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
2	XB	1915	0	1965	40	0
3	QC	1605	0	1667	135	0
3	XC	1605	0	1668	92	18
4	QD	1703	0	1762	108	0
4	XD	1703	0	1763	57	7
5	QE	1155	0	1213	76	0
5	XE	1155	0	1213	33	0
6	QF	843	0	857	10	5
6	XF	843	0	855	89	0
7	QG	1257	0	1296	53	18
7	XG	1257	0	1295	62	0
8	QH	1108	0	1165	52	0
8	XH	1108	0	1165	26	0
9	QI	816	0	822	51	6
9	XI	834	0	847	22	0
10	QJ	801	0	843	199	0
10	XJ	777	0	816	114	6
11	QK	885	0	904	35	0
11	XK	864	0	880	70	0
12	QL	975	0	1062	46	0
12	XL	956	0	1046	30	0
13	QM	921	0	974	137	0
13	XM	914	0	966	128	0
14	QN	492	0	528	288	0
14	XN	492	0	521	172	0
15	QO	734	0	770	57	0
15	XO	729	0	767	44	0
16	QP	705	0	725	71	0
16	XP	705	0	725	16	0
17	QQ	834	0	904	43	0
17	XQ	834	0	902	23	0
18	QR	574	0	644	8	0
18	XR	574	0	643	102	0
19	QS	665	0	678	223	0
19	XS	674	0	695	110	0
20	QT	763	0	861	26	0
20	XT	763	0	861	40	0
21	QU	217	0	223	44	0
21	XU	217	0	234	10	0
22	QV	1452	0	736	3	0
22	XV	1452	0	736	16	0
23	QX	409	0	209	5	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
23	XX	409	0	209	23	0
24	R0	643	0	667	12	0
24	Y0	648	0	672	11	0
25	R1	746	0	826	26	0
25	Y1	729	0	802	11	0
26	R2	581	0	629	6	1
26	Y2	575	0	624	6	0
27	R3	469	0	518	13	2
27	Y3	469	0	518	12	0
28	R4	348	0	354	25	0
28	Y4	357	0	362	9	0
29	R5	459	0	477	20	0
29	Y5	459	0	476	25	1
30	R6	453	0	474	11	0
30	Y6	453	0	473	13	0
31	R7	409	0	454	12	0
31	Y7	418	0	467	18	0
32	R8	517	0	582	31	0
32	Y8	517	0	582	27	0
33	R9	307	0	335	24	0
33	Y9	307	0	336	19	0
34	RA	62070	0	31282	990	0
34	YA	62091	0	31289	1253	1
35	RB	2573	0	1306	21	0
35	YB	2573	0	1306	13	0
36	RD	2115	0	2195	53	2
36	YD	2115	0	2195	64	0
37	RE	1568	0	1634	34	0
37	YE	1568	0	1633	33	0
38	RF	1585	0	1632	33	0
38	YF	1585	0	1632	31	0
39	RG	1474	0	1535	49	0
39	YG	1474	0	1535	27	0
40	RH	1336	0	1418	80	0
40	YH	1336	0	1418	25	0
41	RI	1136	0	1223	41	0
41	YI	1136	0	1223	30	0
42	RN	1104	0	1180	13	0
42	YN	1104	0	1180	17	0
43	RO	933	0	996	23	0
43	YO	933	0	996	19	0
44	RP	1145	0	1228	37	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
44	YP	1122	0	1206	44	2
45	RQ	1122	0	1179	41	0
45	YQ	1122	0	1179	33	0
46	RR	960	0	1021	22	0
46	YR	960	0	1021	24	0
47	RS	882	0	943	19	0
47	YS	882	0	943	16	0
48	RT	1141	0	1202	19	0
48	YT	1141	0	1202	29	0
49	RU	964	0	1022	38	0
49	YU	964	0	1022	22	0
50	RV	779	0	852	17	0
50	YV	779	0	852	11	1
51	RW	900	0	964	20	0
51	YW	900	0	964	19	0
52	RX	725	0	778	11	0
52	YX	725	0	778	9	0
53	RY	818	0	911	25	0
53	YY	818	0	910	19	0
54	RZ	1461	0	1493	36	0
54	YZ	1529	0	1551	33	0
55	QA	70	0	0	0	0
55	QE	1	0	0	0	0
55	QF	1	0	0	0	0
55	QH	2	0	0	0	0
55	QL	2	0	0	0	0
55	R0	2	0	0	0	0
55	R3	1	0	0	0	0
55	R8	2	0	0	0	0
55	RA	432	0	0	0	0
55	RD	1	0	0	0	0
55	RE	4	0	0	0	0
55	RF	2	0	0	0	0
55	RN	1	0	0	0	0
55	RO	1	0	0	0	0
55	XA	88	0	0	2	0
55	XE	1	0	0	0	0
55	XO	1	0	0	0	0
55	Y1	1	0	0	0	0
55	Y2	1	0	0	0	0
55	Y5	1	0	0	0	0
55	Y7	1	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
55	Y8	2	0	0	0	0
55	YA	394	0	0	3	0
55	YB	1	0	0	0	0
55	YD	2	0	0	0	0
55	YE	4	0	0	0	0
55	YF	1	0	0	0	0
55	YP	1	0	0	0	0
55	YQ	1	0	0	0	0
55	YR	2	0	0	0	0
55	YU	1	0	0	0	0
55	YX	1	0	0	0	0
56	QD	8	0	0	2	0
56	XD	8	0	0	0	0
57	QN	1	0	0	0	0
57	R5	1	0	0	0	0
57	R6	1	0	0	0	0
57	R9	1	0	0	0	0
57	RY	1	0	0	0	0
57	XN	1	0	0	0	0
57	Y5	1	0	0	0	0
57	Y6	1	0	0	0	0
57	Y9	1	0	0	0	0
57	YY	1	0	0	1	0
All	All	291185	0	197033	6993	35

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 15.

All (6993) 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:QJ:47:PHE:CZ	14:QN:36:PHE:HB3	1.21	1.72
14:QN:24:CYS:SG	14:QN:40:CYS:HB2	1.24	1.68
1:QA:980:C:C1'	14:QN:19:ARG:HG2	1.23	1.68
1:QA:1049:U:C5	14:QN:3:ARG:HB3	1.26	1.66
1:XA:1190:G:H5'	3:XC:176:HIS:CE1	1.30	1.64
34:YA:1127:A:C2	34:YA:2518:A:C5	1.83	1.64
1:QA:1360:A:C1'	14:QN:17:LYS:HE3	1.23	1.62
1:XA:1318:A:C5	14:XN:16:PHE:CE1	1.79	1.60
7:QG:16:LEU:CD2	9:QI:42:ARG:HG2	1.27	1.59
1:QA:1106:G:C4'	3:QC:172:ARG:HG3	1.25	1.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1106:G:C1'	3:QC:172:ARG:HE	1.06	1.58
1:QA:1106:G:H4'	3:QC:172:ARG:CG	1.21	1.57
34:YA:919:G:C6	34:YA:2268:A:N1	1.69	1.57
10:QJ:47:PHE:HZ	14:QN:36:PHE:CB	1.17	1.56
7:QG:16:LEU:HD21	9:QI:42:ARG:CG	1.21	1.55
1:XA:1240:U:C1'	7:XG:38:LEU:HD11	1.16	1.55
1:XA:1190:G:C5'	3:XC:176:HIS:HE1	1.13	1.54
1:XA:1318:A:C2'	19:XS:11:VAL:HG21	1.37	1.53
1:QA:1221:G:C4'	19:QS:36:ARG:NH2	1.68	1.53
1:XA:1318:A:N6	14:XN:16:PHE:CG	1.76	1.52
34:YA:2070:G:N2	34:YA:2442:C:C2	1.77	1.51
6:XF:100:ASN:HD22	18:XR:27:GLY:CA	1.17	1.51
1:XA:1190:G:C5'	3:XC:176:HIS:CE1	1.88	1.49
1:QA:980:C:H1'	14:QN:19:ARG:CG	1.05	1.49
1:QA:1302:U:C1'	13:QM:27:LYS:HE3	1.02	1.49
1:QA:1236:A:C4'	21:QU:10:ARG:NH1	1.70	1.48
1:XA:1228:C:C5	13:XM:104:ARG:HA	1.45	1.48
1:XA:1240:U:H1'	7:XG:38:LEU:CD1	1.40	1.48
1:QA:1360:A:H1'	14:QN:17:LYS:CE	1.44	1.47
11:XK:91:ARG:NE	18:XR:88:LYS:NZ	1.61	1.46
1:XA:1320:C:N4	19:XS:37:ARG:HB3	1.27	1.46
1:QA:1123:A:H4'	10:QJ:37:PRO:CD	1.44	1.45
40:RH:98:LEU:CD2	40:RH:125:VAL:HG11	1.44	1.45
34:YA:2090:G:N1	34:YA:2230:G:C6	1.84	1.44
34:RA:2751:G:C4	40:RH:3:ARG:HB3	1.52	1.44
29:Y5:32:PRO:N	29:Y5:32:PRO:CA	1.69	1.44
30:R6:16:CYS:SG	30:R6:16:CYS:CB	2.06	1.43
1:QA:1158:C:H4'	2:QB:133:LYS:NZ	1.30	1.43
4:QD:57:ARG:NH2	5:QE:107:ARG:HD3	1.17	1.43
1:QA:1100:C:N4	2:QB:96:ARG:HH22	1.14	1.42
1:QA:1221:G:O2'	19:QS:77:THR:CG2	1.65	1.42
1:QA:974:A:C8	14:QN:31:ARG:NH1	1.81	1.42
1:XA:1318:A:O2'	19:XS:11:VAL:CG2	1.65	1.42
3:QC:23:TYR:CD2	10:QJ:95:GLU:HB2	1.55	1.41
34:YA:1127:A:N3	34:YA:2518:A:N7	1.67	1.41
1:XA:1318:A:C6	14:XN:16:PHE:CD1	2.08	1.41
1:QA:1179:A:C5'	9:QI:83:ARG:HH22	1.32	1.41
1:QA:1188:A:H4'	14:QN:58:LYS:NZ	1.30	1.41
1:QA:657:G:O2'	15:QO:28:GLN:CG	1.68	1.40
1:QA:981:U:C4'	14:QN:21:TYR:OH	1.66	1.40
1:QA:1236:A:H4'	21:QU:10:ARG:NH1	1.10	1.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1187:G:C1'	14:XN:61:TRP:O	1.68	1.40
1:XA:1318:A:C3'	19:XS:11:VAL:HG21	1.50	1.40
34:YA:919:G:C6	34:YA:2268:A:C6	2.09	1.40
6:XF:100:ASN:ND2	18:XR:27:GLY:HA2	1.15	1.39
1:QA:1309:G:H5''	13:QM:77:ASN:ND2	1.38	1.38
34:YA:2082:A:H62	34:YA:2237:G:N2	1.20	1.38
1:QA:980:C:C2	14:QN:19:ARG:O	1.76	1.37
10:QJ:50:ILE:CG1	14:QN:41:ARG:CD	2.01	1.37
1:QA:1106:G:C1'	3:QC:172:ARG:NE	1.84	1.37
4:QD:197:PRO:CD	6:XF:16:GLN:CB	1.94	1.36
1:XA:675:A:C2	11:XK:118:GLY:HA2	1.60	1.36
1:XA:1542:U:O3'	2:XB:169:LYS:NZ	1.58	1.36
13:QM:84:ILE:O	19:QS:74:PHE:CE2	1.76	1.35
1:QA:1360:A:C1'	14:QN:17:LYS:CE	2.01	1.35
14:QN:24:CYS:SG	14:QN:40:CYS:CB	2.14	1.35
1:QA:948:C:H5''	13:QM:101:GLN:CG	1.55	1.35
1:QA:1313:U:H3'	19:QS:6:LYS:NZ	1.04	1.34
7:QG:16:LEU:CD2	9:QI:42:ARG:CG	1.87	1.34
1:QA:980:C:C1'	14:QN:19:ARG:CG	1.85	1.34
1:XA:1316:G:H4'	14:XN:17:LYS:CG	1.49	1.34
1:QA:957:U:C4'	19:QS:79:THR:O	1.74	1.34
1:QA:1253:G:P	10:QJ:44:VAL:HB	1.67	1.34
1:XA:1318:A:C5	14:XN:16:PHE:CD1	2.15	1.33
10:QJ:50:ILE:CG1	14:QN:41:ARG:HD3	1.58	1.33
34:YA:919:G:N1	34:YA:2268:A:C6	1.96	1.33
1:QA:1372:U:OP1	9:QI:68:GLY:CA	1.74	1.32
1:QA:1378:C:OP2	7:QG:7:ALA:CB	1.75	1.32
1:XA:675:A:C2	11:XK:118:GLY:CA	2.10	1.32
1:QA:1187:G:N3	14:QN:60:SER:OG	1.61	1.32
1:XA:1360:A:C8	14:XN:17:LYS:O	1.82	1.32
1:QA:1179:A:H5''	9:QI:83:ARG:NH2	1.41	1.31
4:QD:197:PRO:HD3	6:XF:16:GLN:CB	1.05	1.31
10:QJ:50:ILE:HG12	14:QN:41:ARG:CD	1.59	1.31
1:QA:1320:C:N4	19:QS:37:ARG:HG2	1.42	1.31
1:XA:1305:G:C5	1:XA:1331:G:C6	2.18	1.31
1:QA:1330:U:O4	21:QU:7:ARG:NH1	1.60	1.31
1:QA:1376:U:O4	7:QG:10:ARG:NH2	1.61	1.30
1:XA:1187:G:C2'	14:XN:61:TRP:O	1.71	1.30
1:QA:1313:U:C3'	19:QS:6:LYS:NZ	1.95	1.30
34:RA:2751:G:N3	40:RH:3:ARG:HG2	1.44	1.30
6:XF:100:ASN:HD22	18:XR:27:GLY:C	1.35	1.30

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:972:C:P	10:QJ:57:LYS:HD2	1.70	1.29
1:QA:310:G:P	16:QP:27:LYS:HD3	1.72	1.29
1:XA:1320:C:C1'	19:XS:70:LYS:CD	2.04	1.29
1:QA:979:C:N3	14:QN:19:ARG:HD2	1.47	1.29
4:QD:205:GLU:OE2	5:QE:100:VAL:HG12	1.24	1.29
1:QA:979:C:N4	14:QN:19:ARG:HB2	1.46	1.29
1:QA:1378:C:OP2	7:QG:7:ALA:HB1	1.16	1.29
1:XA:1318:A:H4'	19:XS:11:VAL:CG2	1.63	1.29
1:XA:675:A:H2	11:XK:118:GLY:CA	1.44	1.29
1:XA:1320:C:C1'	19:XS:70:LYS:HD2	1.59	1.29
21:XU:3:LYS:HD3	21:XU:14:TRP:CE3	1.67	1.29
1:QA:986:A:H1'	19:QS:54:GLY:O	1.33	1.28
1:XA:1318:A:C4'	19:XS:11:VAL:CG2	2.12	1.28
34:YA:1127:A:C2	34:YA:2518:A:N7	1.95	1.28
1:XA:1228:C:N4	13:XM:104:ARG:HG2	1.49	1.28
1:QA:1223:C:P	19:QS:78:ARG:NH2	2.07	1.28
1:XA:1226:C:H2'	13:XM:103:THR:OG1	1.12	1.28
1:QA:974:A:P	14:QN:29:ARG:NE	2.06	1.28
34:YA:1566:A:C2	36:YD:214:TRP:CD2	2.21	1.28
34:YA:2052:G:N2	34:YA:2617:C:O2	1.61	1.28
6:XF:94:GLN:OE1	18:XR:32:ARG:CD	1.81	1.27
1:XA:1316:G:C4'	14:XM:17:LYS:HG2	1.60	1.27
1:QA:1106:G:H1'	3:QC:172:ARG:NE	1.45	1.27
1:XA:1229:A:N6	13:XM:104:ARG:HE	1.32	1.27
34:YA:2046:G:C2	34:YA:2623:G:C2	2.22	1.27
1:QA:1307:U:P	13:QM:99:ARG:HG3	1.75	1.26
1:QA:974:A:C5	14:QN:31:ARG:NH1	2.02	1.26
34:RA:2751:G:C5	40:RH:3:ARG:HB3	1.67	1.26
1:XA:1318:A:C6	14:XM:16:PHE:CG	2.18	1.26
1:QA:1221:G:H4'	19:QS:36:ARG:NH2	0.93	1.26
1:QA:1317:C:OP1	14:QN:16:PHE:HD2	1.13	1.26
1:XA:1228:C:C4	13:XM:104:ARG:HG2	1.69	1.26
1:QA:986:A:O2'	19:QS:55:LYS:O	1.52	1.26
1:QA:1320:C:H42	19:QS:37:ARG:CG	1.46	1.26
1:QA:957:U:H4'	19:QS:79:THR:C	1.56	1.25
6:XF:50:TYR:CE1	18:XR:77:GLY:O	1.90	1.25
34:YA:2070:G:C2	34:YA:2442:C:C2	2.23	1.25
1:QA:1100:C:H41	2:QB:96:ARG:NH2	1.32	1.25
1:QA:1106:G:O2'	3:QC:172:ARG:CD	1.85	1.25
1:XA:1230:C:H41	13:XM:105:THR:CB	1.47	1.25
6:XF:94:GLN:OE1	18:XR:32:ARG:HD3	1.08	1.25

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:978:A:N6	14:QN:18:VAL:HG21	1.52	1.25
1:QA:1309:G:C5'	13:QM:77:ASN:ND2	1.98	1.25
34:YA:2090:G:C2	34:YA:2230:G:C6	2.23	1.25
1:QA:980:C:O2	14:QN:19:ARG:O	1.52	1.24
1:XA:1320:C:C5	19:XS:37:ARG:HA	1.71	1.24
13:QM:86:CYS:SG	19:QS:73:GLU:OE2	1.95	1.24
34:YA:2049:G:N2	34:YA:2620:C:C2	2.05	1.24
1:QA:607:A:C2	16:QP:31:LYS:HA	1.73	1.24
1:QA:972:C:OP1	10:QJ:57:LYS:CD	1.85	1.24
1:QA:1317:C:OP1	14:QN:16:PHE:CD2	1.91	1.24
40:RH:98:LEU:HD21	40:RH:125:VAL:CG1	1.66	1.24
12:QL:104:VAL:O	12:QL:105:TYR:CD2	1.90	1.23
3:QC:23:TYR:HB2	10:QJ:93:GLY:O	1.39	1.23
1:XA:1318:A:C4	14:YN:16:PHE:CE1	2.27	1.23
40:RH:103:LEU:HD11	40:RH:123:PHE:CZ	1.72	1.23
1:XA:1312:G:C3'	19:XS:6:LYS:NZ	1.98	1.23
6:XF:99:ALA:O	18:XR:28:GLU:HA	1.10	1.23
34:YA:919:G:O6	34:YA:2268:A:N1	1.59	1.22
3:QC:23:TYR:HB2	10:QJ:93:GLY:C	1.59	1.22
1:QA:669:U:H1'	15:QO:46:HIS:CE1	1.73	1.22
1:QA:1240:U:C6	7:QG:38:LEU:HD13	1.71	1.22
1:QA:1347:G:O6	9:QI:11:LYS:HD3	1.35	1.21
1:XA:1240:U:C1'	7:XG:38:LEU:CD1	2.06	1.21
12:QL:7:ILE:CD1	17:QQ:32:TYR:CD1	2.22	1.21
1:XA:1318:A:C4'	19:XS:11:VAL:HG21	1.68	1.21
1:XA:1377:A:N6	7:XG:10:ARG:CG	2.03	1.21
3:QC:23:TYR:CE2	10:QJ:95:GLU:HB2	1.74	1.21
1:XA:1305:G:O2'	1:XA:1332:A:N6	1.72	1.21
4:QD:57:ARG:CZ	5:QE:107:ARG:HD3	1.70	1.21
34:RA:2751:G:N7	40:RH:2:SER:O	1.73	1.21
1:XA:1190:G:H5'	3:XC:176:HIS:ND1	1.56	1.21
1:QA:959:A:N6	19:QS:79:THR:N	1.89	1.20
1:QA:973:G:O3'	14:QN:29:ARG:NH2	1.75	1.20
1:QA:1049:U:C5	14:QN:3:ARG:CB	2.23	1.20
1:XA:376:G:OP1	16:XP:67:THR:HG21	1.39	1.20
31:Y7:37:LYS:NZ	34:YA:469:G:O6	1.72	1.20
34:YA:1566:A:C2	36:YD:214:TRP:CE2	2.30	1.20
34:YA:270(P):U:O4	41:YI:52:ARG:NE	1.74	1.20
1:XA:1377:A:N6	7:XG:10:ARG:HG2	1.56	1.20
1:XA:1230:C:H41	13:XM:105:THR:CG2	1.53	1.20
1:QA:974:A:N7	14:QN:31:ARG:NH1	1.87	1.19

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1108:G:OP1	3:QC:175:LEU:N	1.74	1.19
1:XA:1320:C:C4	19:XS:37:ARG:CA	2.24	1.19
34:YA:2090:G:N1	34:YA:2230:G:O6	1.70	1.19
4:QD:197:PRO:CD	6:XF:16:GLN:HB2	1.63	1.19
34:YA:2081:C:O2	34:YA:2239:G:N2	1.75	1.19
1:XA:421:U:O4'	3:XC:192:THR:HG21	1.38	1.19
1:XA:1059:C:OP2	3:XC:2:GLY:CA	1.91	1.19
1:XA:1320:C:C4	19:XS:37:ARG:HA	1.77	1.18
34:YA:2045:C:C2	34:YA:2624:G:N2	2.11	1.18
1:QA:959:A:H61	19:QS:78:ARG:HA	1.03	1.18
1:QA:1106:G:O2'	3:QC:172:ARG:HD2	1.05	1.18
1:QA:1179:A:C5'	9:QI:83:ARG:NH2	1.97	1.18
1:XA:1318:A:C6	14:YN:16:PHE:CE1	2.27	1.18
1:XA:1312:G:H3'	19:XS:6:LYS:NZ	1.19	1.17
1:XA:1305:G:C6	1:XA:1331:G:C6	2.31	1.17
1:QA:1123:A:C4'	10:QJ:37:PRO:CD	2.23	1.17
34:YA:2082:A:N6	34:YA:2237:G:N3	1.93	1.17
10:QJ:50:ILE:HG12	14:QN:41:ARG:NE	1.60	1.16
1:XA:1226:C:C2'	13:XM:103:THR:OG1	1.91	1.16
4:QD:88:VAL:HG22	5:QE:96:PRO:O	1.45	1.16
34:YA:1127:A:H2	34:YA:2518:A:C5	1.37	1.16
1:QA:979:C:C4	14:QN:19:ARG:HD2	1.81	1.16
1:QA:1152:A:OP1	10:QJ:13:HIS:NE2	1.76	1.16
1:XA:1305:G:C6	1:XA:1331:G:O6	1.99	1.16
34:YA:919:G:N1	34:YA:2268:A:N1	1.87	1.16
1:XA:1230:C:H41	13:XM:105:THR:HB	1.07	1.15
1:XA:1230:C:N4	13:XM:105:THR:CG2	2.07	1.15
1:XA:1312:G:C3'	19:XS:6:LYS:HZ2	1.55	1.15
1:XA:1320:C:N3	19:XS:36:ARG:HG3	1.59	1.15
1:QA:956:U:O3'	19:QS:80:TYR:O	1.65	1.15
1:QA:1320:C:N3	19:QS:37:ARG:HA	1.59	1.15
4:QD:205:GLU:OE2	5:QE:100:VAL:CG1	1.95	1.15
1:QA:669:U:C1'	15:QO:46:HIS:HE1	1.58	1.14
1:QA:1186:G:N2	14:QN:61:TRP:O	1.79	1.14
1:XA:107:G:O6	20:XT:15:ARG:NE	1.80	1.14
1:XA:1229:A:OP2	13:XM:105:THR:HA	1.47	1.14
1:QA:959:A:N6	19:QS:79:THR:H	1.42	1.14
1:QA:1541:U:P	2:QB:23:ARG:HH21	1.70	1.14
1:XA:1228:C:C6	13:XM:104:ARG:HA	1.81	1.14
34:YA:1050:A:C8	34:YA:2751:G:C4	2.34	1.14
1:QA:974:A:C4	14:QN:31:ARG:NH1	2.13	1.14

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:QD:57:ARG:NH2	5:QE:107:ARG:CD	2.10	1.14
34:YA:1456:G:H1	34:YA:2703:C:N4	1.46	1.14
34:YA:2048:G:C2	34:YA:2621:A:C2	2.37	1.14
1:QA:1229:A:H61	13:QM:104:ARG:CZ	1.60	1.13
1:QA:1188:A:C4'	14:QN:58:LYS:NZ	2.10	1.13
1:QA:959:A:H61	19:QS:78:ARG:CA	1.59	1.13
1:XA:1318:A:N1	14:YN:16:PHE:CE2	2.16	1.13
1:XA:1359:C:P	14:YN:22:THR:OG1	2.06	1.13
1:XA:1229:A:N6	13:XM:104:ARG:NE	1.94	1.12
1:XA:1230:C:H42	13:XM:102:ARG:NH1	1.43	1.13
34:YA:2090:G:N2	34:YA:2230:G:C5	2.15	1.13
1:QA:624:C:O3'	16:QP:10:GLY:HA2	1.44	1.12
1:QA:1367:C:C4'	10:QJ:48:THR:HG21	1.79	1.12
11:QK:71:LYS:NZ	34:RA:2146:C:H41	1.46	1.12
1:XA:1377:A:N6	7:YG:10:ARG:CD	1.97	1.12
4:QD:89:THR:OG1	5:QE:97:GLY:O	1.67	1.12
1:XA:1359:C:OP1	14:YN:22:THR:OG1	1.66	1.12
1:XA:1440(N):G:OP1	20:XT:35:THR:HG21	1.50	1.12
1:QA:974:A:C5'	14:QN:31:ARG:HB3	1.79	1.12
1:QA:1240:U:H2'	7:QG:38:LEU:HD11	1.28	1.12
1:XA:1318:A:N6	14:YN:16:PHE:CD1	2.12	1.12
34:YA:1127:A:C2	34:YA:2518:A:C6	2.36	1.12
1:QA:972:C:OP1	10:QJ:57:LYS:HD2	0.94	1.12
6:XF:100:ASN:ND2	18:XR:27:GLY:CA	1.83	1.12
1:QA:954:G:C2	19:QS:83:HIS:HE1	1.67	1.11
1:QA:1229:A:OP1	13:QM:108:ARG:NH2	1.80	1.11
1:QA:1360:A:N7	14:QN:18:VAL:HA	1.49	1.11
3:QC:79:ARG:NE	11:XK:104:GLN:HG3	1.65	1.11
1:XA:974:A:P	14:YN:31:ARG:HB2	1.90	1.11
3:QC:79:ARG:NH2	11:XK:104:GLN:HA	1.64	1.11
34:RA:2751:G:C4	40:RH:3:ARG:CB	2.32	1.11
1:XA:1187:G:H1'	14:YN:61:TRP:O	1.33	1.11
1:XA:1318:A:O2'	19:XS:11:VAL:HG21	1.33	1.11
1:QA:1123:A:C4'	10:QJ:37:PRO:HD3	1.81	1.11
34:YA:2082:A:N6	34:YA:2237:G:H21	1.48	1.11
1:QA:957:U:H4'	19:QS:79:THR:O	1.41	1.10
1:QA:1320:C:C4	19:QS:37:ARG:HA	1.85	1.10
1:XA:1320:C:N4	19:XS:37:ARG:CB	2.14	1.10
1:QA:538:G:H3'	12:QL:115:LYS:HZ2	1.12	1.10
1:QA:1253:G:OP1	10:QJ:44:VAL:CB	1.98	1.10
1:QA:980:C:C6	14:QN:19:ARG:HG3	1.85	1.10

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
25:R1:90:ILE:HG22	25:R1:94:LEU:HD11	1.19	1.10
7:QG:16:LEU:CD2	9:QI:42:ARG:HG3	1.80	1.10
1:XA:1188:A:H1'	14:XN:60:SER:HA	1.26	1.10
1:QA:981:U:C5'	14:QN:21:TYR:OH	1.98	1.09
45:RQ:59:ARG:HA	54:RZ:179:ASP:OD2	1.51	1.09
1:XA:1228:C:C5	13:XM:104:ARG:CA	2.35	1.09
1:XA:1318:A:C4	14:XN:16:PHE:CZ	2.39	1.09
34:YA:2080:G:C2	34:YA:2241:A:C2	2.39	1.09
3:QC:23:TYR:N	10:QJ:93:GLY:HA2	1.65	1.09
34:YA:2046:G:C4	34:YA:2623:G:N2	2.20	1.09
1:XA:1320:C:C1'	19:XS:70:LYS:HD3	1.83	1.09
1:QA:949:A:OP1	13:QM:101:GLN:HA	1.52	1.09
12:QL:7:ILE:HD11	17:QQ:32:TYR:HB3	1.28	1.09
14:XN:56:VAL:HG13	14:XN:57:ARG:H	1.01	1.09
34:YA:919:G:C5	34:YA:2268:A:N6	2.20	1.09
1:QA:974:A:N9	14:QN:31:ARG:NH1	2.01	1.08
1:QA:983:A:H5'	14:QN:2:ALA:HB2	1.15	1.08
1:QA:1150:U:H1'	10:QJ:39:PRO:HG2	1.10	1.08
34:RA:2751:G:N3	40:RH:3:ARG:CG	2.16	1.08
1:XA:1112:C:O4'	3:XC:179:ARG:NH2	1.86	1.08
1:XA:1377:A:P	7:XG:94:ARG:NH2	2.25	1.08
34:YA:2043:C:N4	34:YA:2777:G:N1	2.01	1.08
34:YA:2090:G:C6	34:YA:2230:G:O6	2.05	1.08
1:QA:948:C:H5''	13:QM:101:GLN:HG3	1.14	1.08
3:QC:23:TYR:H	10:QJ:93:GLY:HA2	1.03	1.08
12:QL:7:ILE:HD11	17:QQ:32:TYR:CB	1.83	1.08
1:QA:1219:U:O2'	19:QS:34:TRP:CE3	2.04	1.08
11:XK:91:ARG:HH21	18:XR:88:LYS:HD2	1.13	1.08
3:QC:23:TYR:CE2	10:QJ:95:GLU:CB	2.35	1.08
10:QJ:50:ILE:HD11	14:QN:41:ARG:HD2	1.31	1.08
1:XA:1318:A:N6	14:XN:16:PHE:CB	2.17	1.08
1:XA:1318:A:C2	14:XN:16:PHE:CZ	2.42	1.08
1:XA:1320:C:H1'	19:XS:70:LYS:HD2	1.26	1.08
10:QJ:49:VAL:CG2	14:QN:41:ARG:HB2	1.84	1.08
1:XA:186(B):C:O2	20:XT:105:SER:HB2	1.49	1.08
10:XJ:40:LEU:HB3	10:XJ:41:PRO:CD	1.84	1.08
10:QJ:50:ILE:HD11	14:QN:41:ARG:NH1	1.67	1.07
1:QA:1320:C:O2	19:QS:36:ARG:O	1.71	1.07
1:QA:1358:U:O3'	14:QN:22:THR:HG21	1.53	1.07
6:XF:99:ALA:O	18:XR:28:GLU:CA	2.02	1.07
34:YA:918:A:N7	34:YA:2268:A:N6	2.01	1.07

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:2048:G:N1	34:YA:2621:A:C2	2.20	1.07
1:QA:1100:C:C5	2:QB:96:ARG:NH2	2.21	1.07
1:QA:974:A:O5'	14:QN:31:ARG:HB2	1.53	1.07
1:QA:978:A:H62	14:QN:18:VAL:CG2	1.66	1.07
1:XA:702:A:N6	34:YA:1848:A:C2	2.22	1.07
1:QA:1240:U:N3	7:QG:38:LEU:HB3	1.40	1.07
1:XA:1318:A:C6	14:YN:16:PHE:CD2	2.43	1.07
13:XM:84:ILE:HD11	19:XS:65:ASN:CG	1.73	1.07
34:YA:945:A:N6	34:YA:2448:A:N6	1.61	1.07
34:YA:1127:A:H2	34:YA:2518:A:C4	1.72	1.07
1:QA:972:C:P	10:QJ:57:LYS:CD	2.42	1.06
1:XA:741:G:P	15:XO:35:ARG:NH2	2.26	1.06
1:XA:1377:A:C6	7:XG:10:ARG:CG	2.38	1.06
4:XD:18:LYS:NZ	4:XD:31:CYS:SG	2.26	1.06
1:QA:1280:A:H5''	10:QJ:40:LEU:HD23	1.30	1.06
14:YN:27:CYS:SG	14:YN:28:GLY:N	2.28	1.06
1:QA:1253:G:P	10:QJ:44:VAL:CB	2.43	1.06
1:QA:1253:G:OP1	10:QJ:44:VAL:HB	1.54	1.06
1:QA:1318:A:N3	19:QS:37:ARG:CZ	2.17	1.06
4:QD:197:PRO:HD3	6:XF:16:GLN:HB3	1.06	1.06
34:YA:2046:G:C6	34:YA:2623:G:N1	2.24	1.06
1:QA:657:G:O2'	15:QO:28:GLN:HG2	0.88	1.06
1:QA:1160:G:O4'	2:QB:132:LYS:HE3	1.54	1.06
1:QA:1229:A:N6	13:QM:104:ARG:HG3	1.69	1.06
1:QA:1367:C:H4'	10:QJ:48:THR:HG21	1.06	1.06
1:XA:1318:A:C4'	19:XS:11:VAL:HG23	1.82	1.06
1:QA:1152:A:OP1	10:QJ:13:HIS:CE1	2.09	1.06
34:YA:2044:C:C4	34:YA:2625:G:N2	2.23	1.06
1:QA:538:G:H3'	12:QL:115:LYS:NZ	1.71	1.05
10:QJ:45:ARG:HG2	14:QN:36:PHE:CZ	1.69	1.05
1:XA:1317:C:C5	14:YN:16:PHE:CD1	2.27	1.05
10:XJ:10:GLY:HA3	10:XJ:16:LEU:HD21	1.36	1.05
10:QJ:50:ILE:HG13	14:QN:41:ARG:HD3	1.34	1.05
1:XA:737:A:OP1	6:XF:92:LYS:NZ	1.87	1.05
1:QA:1253:G:O5'	10:QJ:44:VAL:HB	1.57	1.05
12:QL:7:ILE:HD11	17:QQ:32:TYR:CG	1.90	1.05
1:XA:1319:A:OP2	19:XS:3:ARG:NE	1.88	1.05
1:QA:974:A:H5'	14:QN:31:ARG:HB3	1.33	1.05
1:QA:1179:A:H5'	9:QI:83:ARG:HH22	1.17	1.05
40:RH:87:LEU:O	40:RH:131:VAL:HG23	1.54	1.05
1:XA:1304:G:H5''	1:XA:1304:G:H8	1.16	1.05

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:2070:G:N2	34:YA:2442:C:N1	2.04	1.05
1:QA:1320:C:C2	19:QS:36:ARG:O	2.10	1.05
2:QB:196:LEU:HA	8:QH:74:PRO:HG3	1.37	1.05
1:XA:1309:G:C4'	13:XM:77:ASN:ND2	2.20	1.05
10:XJ:16:LEU:HB3	10:XJ:70:ARG:NH1	1.71	1.05
34:YA:1392:A:N6	52:YX:15:GLU:OE2	1.89	1.05
1:QA:669:U:H1'	15:QO:46:HIS:HE1	0.89	1.04
1:QA:876:G:H1'	8:QH:11:THR:HG21	1.40	1.04
1:QA:1221:G:C5'	19:QS:36:ARG:NH2	2.19	1.04
1:QA:607:A:N1	16:QP:31:LYS:HA	1.72	1.04
1:QA:1320:C:N4	19:QS:37:ARG:CG	2.12	1.04
1:QA:1357:A:C5'	10:QJ:45:ARG:HH22	1.70	1.04
34:YA:1566:A:N1	36:YD:214:TRP:CE2	2.26	1.04
34:YA:2099:U:O2	34:YA:2190:G:N2	1.91	1.04
1:XA:280:C:N3	17:XQ:39:SER:N	2.03	1.04
1:XA:1320:C:H1'	19:XS:70:LYS:CD	1.77	1.04
1:QA:973:G:O3'	14:QN:29:ARG:CZ	2.05	1.04
1:XA:974:A:O5'	14:XN:31:ARG:HB3	1.55	1.04
34:YA:2108:C:H42	34:YA:2182:G:N2	1.55	1.04
1:QA:948:C:H5''	13:QM:101:GLN:HG2	1.40	1.03
1:QA:972:C:OP2	10:QJ:57:LYS:HG3	1.58	1.03
1:QA:1357:A:H5'	10:QJ:45:ARG:NH2	1.73	1.03
1:XA:191:G:O2'	20:XT:101:GLY:O	1.74	1.03
1:XA:1190:G:H5''	3:XC:176:HIS:HE1	0.88	1.03
1:XA:1318:A:H4'	19:XS:11:VAL:HG23	1.08	1.03
1:XA:1225:A:H1'	19:XS:78:ARG:HE	1.18	1.03
6:XF:91:VAL:HG21	18:XR:34:TYR:CE1	1.92	1.03
1:QA:981:U:H4'	14:QN:21:TYR:OH	1.22	1.03
1:XA:1227:A:OP1	13:XM:96:LEU:HD21	1.59	1.03
1:XA:1318:A:C2	14:XN:16:PHE:CE2	2.46	1.03
11:XK:91:ARG:HE	18:XR:88:LYS:NZ	1.31	1.03
34:YA:2082:A:N6	34:YA:2237:G:N2	2.03	1.03
1:QA:1313:U:C5	19:QS:6:LYS:HE2	1.92	1.03
1:XA:186(B):C:O2	20:XT:105:SER:CB	2.07	1.03
3:XC:23:TYR:CE2	10:XJ:95:GLU:HB2	1.94	1.03
13:XM:84:ILE:HG13	19:XS:65:ASN:O	1.58	1.03
34:YA:1127:A:N3	34:YA:2518:A:C8	2.27	1.03
1:QA:979:C:C2	14:QN:19:ARG:HD2	1.93	1.02
1:QA:1106:G:O4'	3:QC:172:ARG:NE	1.92	1.02
1:QA:1226:C:OP1	13:QM:87:TYR:OH	1.75	1.02
3:QC:23:TYR:CD2	10:QJ:95:GLU:CB	2.42	1.02

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1372:U:OP1	9:XI:68:GLY:HA2	1.58	1.02
1:QA:1100:C:N4	2:QB:96:ARG:NH2	1.97	1.02
1:XA:1318:A:N7	14:XN:16:PHE:CE1	2.27	1.02
1:XA:1374:A:O2'	7:XG:31:MET:HE2	1.60	1.02
1:QA:1240:U:C6	7:QG:38:LEU:CD1	2.42	1.02
1:QA:1280:A:H5''	10:QJ:40:LEU:CD2	1.90	1.02
1:XA:1309:G:C5'	13:XM:77:ASN:ND2	2.23	1.02
1:XA:1318:A:C5	14:XN:16:PHE:CZ	2.46	1.02
1:XA:1318:A:O2'	19:XS:11:VAL:HG22	1.58	1.02
13:XM:84:ILE:HG22	19:XS:74:PHE:CZ	1.94	1.02
1:QA:309:G:H5''	16:QP:27:LYS:HE2	1.41	1.02
7:QG:16:LEU:HD22	9:QI:42:ARG:CG	1.90	1.02
1:QA:657:G:C2'	15:QO:28:GLN:HG2	1.86	1.01
1:QA:668:G:H4'	15:QO:48:LYS:O	1.58	1.01
1:XA:1305:G:H2'	1:XA:1331:G:N2	1.74	1.01
34:YA:197:A:C4	34:YA:2430:A:C2	2.47	1.01
1:QA:953:G:C5	13:QM:104:ARG:NH2	2.27	1.01
34:YA:2051:A:C2	34:YA:2614:A:C2	2.47	1.01
1:XA:1377:A:OP1	7:XG:94:ARG:NH2	1.93	1.01
6:XF:99:ALA:HB3	18:XR:29:PHE:CD1	1.96	1.01
34:YA:2056:G:C6	34:YA:2577:A:C4	2.48	1.01
34:YA:2082:A:N6	34:YA:2237:G:C2	2.26	1.01
1:XA:412:A:C2	4:XD:35:ARG:HD3	1.96	1.01
1:XA:974:A:C5'	14:XN:31:ARG:HB3	1.89	1.01
34:YA:2070:G:N1	34:YA:2442:C:N3	2.09	1.01
1:QA:974:A:P	14:QN:29:ARG:CZ	2.48	1.01
1:QA:1360:A:O4'	14:QN:17:LYS:CE	2.06	1.01
13:QM:84:ILE:HG21	19:QS:69:HIS:CE1	1.95	1.01
26:R2:48:HIS:HE2	26:R2:49:LYS:HE2	1.21	1.01
1:XA:974:A:C8	14:XN:31:ARG:CD	2.44	1.01
34:YA:1050:A:C8	34:YA:2751:G:C5	2.49	1.00
34:YA:2046:G:N3	34:YA:2623:G:C2	2.29	1.00
34:YA:2311:A:H1'	39:YG:88:ILE:HD12	1.41	1.00
1:QA:980:C:H1'	14:QN:19:ARG:CD	1.90	1.00
1:QA:1123:A:C4'	10:QJ:37:PRO:HD2	1.91	1.00
10:QJ:47:PHE:CZ	14:QN:36:PHE:CB	2.06	1.00
1:XA:719:C:O2	18:XR:50:ILE:HD12	1.60	1.00
1:QA:986:A:C1'	19:QS:55:LYS:HA	1.91	1.00
3:QC:23:TYR:CE1	10:QJ:9:ARG:O	2.14	1.00
26:R2:48:HIS:NE2	26:R2:49:LYS:HE2	1.74	1.00
1:QA:1367:C:H4'	10:QJ:48:THR:CG2	1.92	1.00

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1301:U:O2'	13:XM:17:VAL:HG21	1.60	1.00
1:XA:1318:A:C3'	19:XS:11:VAL:CG2	2.36	1.00
22:XV:8:U:H3	22:XV:14:A:H62	1.01	1.00
34:YA:2046:G:N3	34:YA:2623:G:N2	2.10	1.00
1:QA:740:U:H4'	15:QO:39:LEU:HG	1.44	1.00
1:QA:1186:G:H21	14:QN:61:TRP:C	1.65	1.00
1:QA:1229:A:N6	13:QM:104:ARG:CG	2.25	1.00
13:QM:84:ILE:O	19:QS:74:PHE:CZ	2.15	0.99
1:XA:1160:G:O4'	2:XB:132:LYS:HE3	1.62	0.99
1:QA:980:C:C1'	14:QN:19:ARG:HG3	1.89	0.99
1:QA:1360:A:N7	14:QN:18:VAL:CA	2.12	0.99
34:YA:2090:G:C2	34:YA:2230:G:C5	2.51	0.99
1:XA:1059:C:OP2	3:XC:2:GLY:HA2	1.60	0.99
6:XF:50:TYR:CE1	18:XR:77:GLY:C	2.22	0.99
1:QA:959:A:H62	19:QS:79:THR:N	1.53	0.99
1:QA:1221:G:OP1	19:QS:36:ARG:CZ	2.10	0.99
1:XA:728:A:N7	15:XO:54:ARG:CZ	2.26	0.99
6:XF:91:VAL:HG21	18:XR:34:TYR:CZ	1.97	0.99
1:QA:974:A:OP2	14:QN:29:ARG:NE	1.96	0.99
1:XA:185:A:O2'	20:XT:81:LYS:NZ	1.96	0.99
34:YA:2051:A:H2	34:YA:2614:A:C2	1.78	0.99
1:XA:1305:G:C5	1:XA:1331:G:O6	2.13	0.99
34:YA:919:G:C6	34:YA:2268:A:N6	2.29	0.99
34:YA:1782:C:N4	34:YA:2587:A:C2	2.31	0.99
34:YA:2059:A:N1	34:YA:2503:A:C6	2.31	0.99
1:QA:1158:C:C4'	2:QB:133:LYS:NZ	2.25	0.99
10:QJ:50:ILE:CD1	14:QN:41:ARG:CD	2.41	0.99
10:QJ:50:ILE:CD1	14:QN:41:ARG:HD2	1.92	0.99
1:QA:1313:U:H3'	19:QS:6:LYS:HZ3	1.20	0.98
1:XA:1228:C:O5'	13:XM:108:ARG:NH2	1.95	0.98
1:XA:1230:C:N4	13:XM:102:ARG:HH12	1.60	0.98
1:QA:1302:U:H1'	13:QM:27:LYS:HE3	1.38	0.98
1:XA:1320:C:C4	19:XS:37:ARG:CB	2.44	0.98
1:QA:667:G:H4'	15:QO:51:HIS:CE1	1.96	0.98
1:QA:1188:A:C4'	14:QN:58:LYS:HZ1	1.73	0.98
34:YA:2051:A:N1	34:YA:2614:A:C4	2.31	0.98
28:R4:7:PRO:HG3	39:RG:61:ALA:HB1	1.45	0.98
1:XA:1533:C:N4	23:XX:12:A:C2	2.32	0.98
34:YA:199:A:N6	34:YA:2434:A:C6	2.31	0.98
34:YA:1566:A:N1	36:YD:214:TRP:CZ2	2.31	0.98
1:QA:1221:G:OP1	19:QS:36:ARG:NH1	1.96	0.98

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1229:A:N6	13:QM:104:ARG:NE	2.10	0.98
1:QA:1318:A:N3	19:QS:37:ARG:NH1	2.10	0.98
3:QC:23:TYR:H	10:QJ:93:GLY:CA	1.75	0.98
34:YA:197:A:C5	34:YA:2430:A:C2	2.52	0.98
34:YA:2108:C:H42	34:YA:2182:G:H22	1.09	0.98
1:XA:1318:A:N3	19:XS:37:ARG:NH1	2.11	0.98
1:QA:1372:U:OP1	9:QL:68:GLY:HA2	0.80	0.98
10:QJ:50:ILE:HG12	14:QN:41:ARG:HD3	1.23	0.98
34:YA:197:A:C4	34:YA:2430:A:H2	1.82	0.98
1:QA:1240:U:C5	7:QG:38:LEU:HD13	1.99	0.97
3:QC:23:TYR:CB	10:QJ:93:GLY:O	2.11	0.97
1:XA:1309:G:C4'	13:XM:77:ASN:HD21	1.77	0.97
1:QA:1226:C:C5'	13:QM:91:ARG:HH12	1.76	0.97
4:QD:204:ILE:CG2	5:QE:98:THR:O	2.12	0.97
34:YA:2048:G:C2	34:YA:2621:A:N3	2.32	0.97
1:XA:1304:G:H5''	1:XA:1304:G:C8	1.99	0.97
1:QA:956:U:C2'	19:QS:80:TYR:O	2.11	0.97
25:R1:95:LEU:O	25:R1:95:LEU:HD23	1.64	0.97
29:R5:3:LYS:NZ	34:RA:2611:U:O4	1.96	0.97
34:YA:2059:A:C2	34:YA:2503:A:N6	2.33	0.97
1:QA:974:A:OP2	14:QN:29:ARG:CG	2.12	0.97
1:QA:1150:U:C1'	10:QJ:39:PRO:HG2	1.95	0.97
1:QA:1158:C:C4'	2:QB:133:LYS:HZ2	1.78	0.97
1:QA:1226:C:H5'	13:QM:91:ARG:HH12	1.26	0.96
1:XA:1228:C:H5	13:XM:104:ARG:HA	1.23	0.96
1:XA:1112:C:H1'	3:XC:179:ARG:NE	1.80	0.96
11:XK:110:ASP:HB3	18:XR:85:LEU:O	1.65	0.96
1:QA:1152:A:P	10:QJ:13:HIS:CE1	2.51	0.96
33:R9:19:ARG:NE	34:RA:2756:U:OP2	1.97	0.96
1:XA:1124:G:H4'	10:XJ:36:GLY:H	1.30	0.96
11:XK:116:HIS:NE2	18:XR:82:THR:HB	1.80	0.96
34:YA:1174:A:N6	55:YA:3192:MG:MG	1.21	0.96
1:QA:957:U:O2'	19:QS:79:THR:O	1.83	0.96
30:R6:13:CYS:SG	30:R6:14:THR:N	2.39	0.96
1:XA:1305:G:O2'	1:XA:1332:A:C6	2.18	0.96
1:XA:1318:A:O2'	19:XS:11:VAL:CG1	2.14	0.96
29:Y5:43:HIS:CD2	34:YA:2884:U:H5	1.83	0.96
34:YA:747:U:O4	34:YA:2613:U:N3	1.98	0.96
34:YA:2054:A:H62	34:YA:2577:A:H61	1.06	0.96
1:QA:761:G:H5'	17:QQ:100:LYS:NZ	1.81	0.96
1:QA:1099:G:OP2	2:QB:96:ARG:CD	2.13	0.96

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:974:A:O5'	14:XN:31:ARG:CB	2.13	0.96
1:XA:974:A:C1'	14:XN:31:ARG:HH11	1.77	0.96
1:XA:1360:A:C8	14:XN:17:LYS:C	2.35	0.96
11:XK:108:ILE:O	18:XR:87:ARG:HD2	1.66	0.96
1:QA:742:G:H5''	15:QO:58:MET:SD	2.06	0.96
1:QA:1049:U:C4	14:QN:3:ARG:HB3	2.00	0.95
1:XA:675:A:C2	11:XK:118:GLY:HA3	1.98	0.95
6:XF:87:ARG:NH1	18:XR:76:LEU:O	1.99	0.95
34:YA:1252:G:N3	49:YU:33:ARG:NH1	2.13	0.95
1:QA:675:A:H2	11:QK:118:GLY:HA2	1.29	0.95
1:QA:880:C:H5''	12:QL:12:ARG:NH2	1.80	0.95
13:QM:86:CYS:HB2	19:QS:69:HIS:CE1	1.99	0.95
1:XA:1318:A:C6	14:XN:16:PHE:CZ	2.52	0.95
34:YA:2044:C:N3	34:YA:2625:G:C2	2.33	0.95
1:QA:657:G:HO2'	15:QO:28:GLN:HG2	1.14	0.95
1:QA:1221:G:O2'	19:QS:77:THR:HG21	0.78	0.95
1:QA:982:U:O5'	14:QN:6:LEU:HD21	1.67	0.95
1:XA:974:A:C4	14:XN:31:ARG:NH1	2.35	0.95
1:QA:957:U:C2'	19:QS:79:THR:O	2.15	0.95
1:QA:982:U:H5''	14:QN:6:LEU:CG	1.96	0.95
1:QA:1347:G:O6	9:QI:11:LYS:CD	2.15	0.95
1:QA:1099:G:P	2:QB:96:ARG:HD2	2.05	0.95
1:XA:1226:C:H2'	13:XM:103:THR:HG1	1.30	0.95
1:XA:1318:A:N6	14:XN:16:PHE:HB3	1.80	0.95
10:XJ:37:PRO:HB3	10:XJ:72:VAL:HG22	1.46	0.95
34:YA:1050:A:H8	34:YA:2751:G:C4	1.84	0.95
1:QA:1360:A:O4'	14:QN:17:LYS:NZ	2.00	0.94
1:XA:1230:C:H42	13:XM:102:ARG:HH12	1.00	0.94
1:QA:974:A:C5'	14:QN:31:ARG:CB	2.44	0.94
1:QA:986:A:C1'	19:QS:54:GLY:O	2.15	0.94
34:YA:2046:G:C6	34:YA:2623:G:C6	2.55	0.94
1:XA:1320:C:C5	19:XS:37:ARG:CA	2.44	0.94
1:XA:1320:C:C4	19:XS:37:ARG:HB3	2.00	0.94
1:QA:986:A:H1'	19:QS:55:LYS:HA	1.46	0.94
1:QA:1160:G:H5'	2:QB:132:LYS:HE2	1.48	0.94
1:QA:1223:C:OP1	19:QS:78:ARG:NH2	1.98	0.94
1:XA:1112:C:C1'	3:XC:179:ARG:HH21	1.80	0.94
1:QA:1160:G:O4'	2:QB:132:LYS:CE	2.15	0.94
34:YA:1566:A:C2	36:YD:214:TRP:CG	2.54	0.94
34:YA:2071:A:H2	34:YA:2441:C:N3	1.65	0.94
1:QA:186(K):G:N7	17:QQ:63:ARG:NH2	2.14	0.94

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1188:A:C1'	14:XN:60:SER:HA	1.98	0.94
1:XA:1230:C:N4	13:XM:105:THR:HB	1.82	0.94
1:XA:1318:A:N1	14:XN:16:PHE:CD2	2.36	0.94
1:XA:1377:A:C6	7:XG:10:ARG:HG2	2.02	0.94
34:YA:2046:G:N1	34:YA:2623:G:C6	2.35	0.94
1:XA:1228:C:N4	13:XM:104:ARG:CG	2.29	0.94
10:XJ:40:LEU:HB3	10:XJ:41:PRO:HD2	1.49	0.94
1:QA:626:U:H4'	16:QP:38:TYR:CD2	2.02	0.94
1:QA:1099:G:OP2	2:QB:96:ARG:HD2	1.68	0.94
1:QA:1150:U:O2	10:QJ:39:PRO:HG3	1.68	0.94
10:QJ:50:ILE:CD1	14:QN:41:ARG:CZ	2.45	0.94
7:XG:16:LEU:CD2	9:XI:44:VAL:HG23	1.98	0.94
1:QA:1229:A:C6	13:QM:104:ARG:NE	2.35	0.94
1:QA:1229:A:H62	13:QM:104:ARG:HG3	1.28	0.94
28:R4:26:SER:OG	39:RG:143:GLU:OE2	1.84	0.94
22:XV:8:U:H3	22:XV:14:A:N6	1.65	0.94
1:QA:1236:A:C5'	21:QU:10:ARG:NH1	2.30	0.94
4:QD:197:PRO:CD	6:XF:16:GLN:HB3	1.78	0.94
34:RA:2680:C:OP2	37:RE:111:ARG:NH2	2.01	0.94
1:QA:1313:U:O4	19:QS:4:SER:HB2	1.68	0.93
34:YA:2087:G:O6	34:YA:2233:U:C4	2.21	0.93
3:XC:60:ALA:HB1	10:XJ:91:PRO:HD2	1.50	0.93
6:XF:100:ASN:HB2	18:XR:27:GLY:C	1.89	0.93
1:QA:1229:A:N6	13:QM:104:ARG:CZ	2.31	0.93
1:XA:1360:A:H1'	14:XN:17:LYS:HD2	1.50	0.93
1:QA:1229:A:P	13:QM:108:ARG:NH2	2.40	0.93
1:XA:1309:G:H5'	13:XM:77:ASN:HD22	1.33	0.93
1:QA:1313:U:C3'	19:QS:6:LYS:HZ2	1.69	0.93
1:XA:974:A:O4'	14:XN:31:ARG:HD3	1.69	0.93
1:QA:1158:C:H4'	2:QB:133:LYS:HZ1	1.22	0.93
34:RA:2751:G:C2	40:RH:3:ARG:HG2	2.04	0.93
1:XA:974:A:C4	14:XN:31:ARG:CZ	2.52	0.93
1:QA:956:U:O2'	19:QS:80:TYR:O	1.87	0.93
10:QJ:50:ILE:HD11	14:QN:41:ARG:HH11	1.23	0.93
34:YA:919:G:N1	34:YA:2268:A:C2	2.35	0.93
34:YA:1493:C:N4	34:YA:2210:G:C8	2.37	0.93
1:QA:958:A:N6	19:QS:77:THR:O	2.01	0.93
1:QA:1014:A:O4'	19:QS:34:TRP:CD1	2.22	0.93
1:QA:1360:A:O4'	14:QN:17:LYS:HE3	1.66	0.93
33:R9:31:LYS:HE2	34:RA:2478:A:H5'	1.47	0.93
4:QD:205:GLU:HG2	5:QE:100:VAL:O	1.68	0.92

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1217:C:OP1	14:XN:5:ALA:HB1	1.68	0.92
34:YA:2070:G:C2	34:YA:2442:C:O2	2.20	0.92
1:QA:957:U:O4'	19:QS:79:THR:O	1.85	0.92
1:XA:253:U:OP1	17:XQ:67:LYS:HD3	1.69	0.92
1:QA:1378:C:P	7:QG:7:ALA:HB1	2.07	0.92
1:QA:948:C:C5'	13:QM:101:GLN:CG	2.47	0.92
1:QA:959:A:N6	19:QS:78:ARG:HA	1.83	0.92
10:QJ:50:ILE:CD1	14:QN:41:ARG:NH1	2.31	0.92
34:YA:2054:A:H62	34:YA:2577:A:N6	1.67	0.92
6:XF:100:ASN:ND2	18:XR:27:GLY:C	2.14	0.92
1:QA:1309:G:C5'	13:QM:77:ASN:HD21	1.74	0.92
1:XA:1123:A:O2'	10:XJ:38:ILE:HG23	1.69	0.92
1:QA:946:A:OP1	13:QM:114:ARG:NH2	2.02	0.92
11:XK:91:ARG:CD	18:XR:88:LYS:NZ	2.33	0.92
1:QA:979:C:H2'	14:QN:19:ARG:CZ	1.99	0.92
1:QA:980:C:O4'	14:QN:19:ARG:NE	2.01	0.92
1:QA:1221:G:H4'	19:QS:36:ARG:HH21	1.29	0.92
1:QA:1541:U:P	2:QB:23:ARG:NH2	2.43	0.91
1:XA:974:A:C8	14:XN:31:ARG:HD3	2.05	0.91
1:QA:137:C:H1'	16:QP:63:GLY:HA3	1.50	0.91
11:QK:71:LYS:HZ2	34:RA:2146:C:H41	1.16	0.91
1:QA:1188:A:H4'	14:QN:58:LYS:HZ3	1.14	0.91
1:QA:1357:A:H5'	10:QJ:45:ARG:CZ	1.99	0.91
3:QC:23:TYR:N	10:QJ:93:GLY:CA	2.32	0.91
34:YA:2069:G:N2	34:YA:2443:C:C2	2.38	0.91
1:QA:954:G:C2	19:QS:83:HIS:CE1	2.58	0.91
1:QA:954:G:N2	19:QS:83:HIS:CE1	2.38	0.91
34:YA:491:G:O6	51:YW:49:LYS:NZ	2.02	0.91
1:QA:983:A:C5'	14:QN:2:ALA:HB2	1.99	0.91
1:QA:975:A:N6	10:QJ:48:THR:OG1	2.02	0.91
10:QJ:50:ILE:HD11	14:QN:41:ARG:CD	2.01	0.91
1:XA:1309:G:H4'	13:XM:77:ASN:HD21	1.35	0.91
34:YA:2045:C:C2	34:YA:2624:G:C2	2.58	0.91
1:QA:1313:U:C6	19:QS:6:LYS:HE2	2.05	0.91
4:QD:196:LEU:O	6:XF:16:GLN:NE2	1.81	0.91
14:XN:56:VAL:HG13	14:XN:57:ARG:N	1.85	0.91
1:XA:675:A:N3	11:XK:118:GLY:CA	2.33	0.91
1:XA:1309:G:H5'	13:XM:77:ASN:ND2	1.86	0.91
11:XK:109:VAL:HG22	18:XR:86:VAL:HG23	1.52	0.91
1:QA:1150:U:O2'	10:QJ:39:PRO:HB2	1.70	0.91
1:XA:51:A:C4	1:XA:353:A:C6	2.59	0.91

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:2056:G:C5	34:YA:2577:A:C5	2.58	0.91
1:QA:538:G:H5''	12:QL:115:LYS:HG2	1.53	0.90
1:QA:609:A:H5'	16:QP:9:PHE:CE1	2.06	0.90
1:QA:1100:C:C4	2:QB:96:ARG:NH2	2.39	0.90
1:XA:974:A:P	14:XN:31:ARG:CB	2.59	0.90
34:YA:2059:A:C2	34:YA:2503:A:C6	2.59	0.90
34:YA:2081:C:N3	34:YA:2239:G:N1	2.19	0.90
1:QA:1014:A:P	19:QS:32:LYS:HE2	2.11	0.90
34:YA:1456:G:H1	34:YA:2703:C:H42	0.98	0.90
34:YA:2093:G:H21	34:YA:2198:A:N6	1.68	0.90
1:QA:972:C:P	10:QJ:57:LYS:CG	2.58	0.90
1:QA:1235:U:O3'	21:QU:10:ARG:HD2	1.70	0.90
1:XA:528:C:N4	12:XL:49:ASN:OD1	2.05	0.90
4:QD:57:ARG:HH22	5:QE:107:ARG:CD	1.79	0.90
33:R9:23:VAL:HG11	34:RA:1032:A:H4'	1.51	0.90
1:XA:978:A:N7	14:XN:18:VAL:HG21	1.85	0.90
1:QA:1372:U:P	9:QL:68:GLY:HA2	2.11	0.90
25:R1:90:ILE:HA	25:R1:94:LEU:CD1	2.00	0.90
1:XA:1230:C:N4	13:XM:105:THR:HG22	1.87	0.90
34:YA:2046:G:N1	34:YA:2623:G:N1	2.20	0.90
34:YA:2043:C:N4	34:YA:2777:G:C2	2.25	0.90
1:QA:583:A:H4'	17:QQ:91:ARG:CG	2.01	0.90
1:XA:1305:G:N1	1:XA:1331:G:N7	2.19	0.90
1:XA:1318:A:N7	14:XN:16:PHE:CD1	2.40	0.90
34:YA:919:G:C2	34:YA:2268:A:C6	2.59	0.90
1:QA:1111:A:N1	3:QC:177:THR:HA	1.87	0.90
1:QA:1160:G:H5'	2:QB:132:LYS:CE	2.01	0.90
1:QA:1253:G:P	10:QJ:44:VAL:CG2	2.60	0.90
34:YA:2045:C:N3	34:YA:2624:G:C2	2.39	0.90
10:QJ:49:VAL:HG22	14:QN:41:ARG:HB2	1.52	0.89
1:XA:675:A:N3	11:XK:118:GLY:HA3	1.87	0.89
1:XA:1059:C:OP2	3:XC:2:GLY:C	2.10	0.89
1:XA:1318:A:C6	14:XN:16:PHE:CE2	2.60	0.89
1:QA:1320:C:N3	19:QS:36:ARG:O	2.05	0.89
33:R9:23:VAL:HG21	34:RA:1032:A:H1'	1.51	0.89
1:XA:708:C:OP1	11:XK:85:ARG:NH1	2.03	0.89
1:XA:1123:A:O3'	10:XJ:36:GLY:HA3	1.71	0.89
1:XA:1320:C:C5	19:XS:37:ARG:CB	2.55	0.89
1:XA:403:C:N4	55:XA:1688:MG:MG	1.30	0.89
6:XF:50:TYR:HE1	18:XR:77:GLY:C	1.73	0.89
1:QA:1188:A:H4'	14:QN:58:LYS:HZ1	1.27	0.89

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1357:A:H5'	10:QJ:45:ARG:HH22	1.31	0.89
7:QG:16:LEU:HD22	9:QI:42:ARG:HG3	1.45	0.89
1:QA:676:A:H4'	11:QK:115:PRO:HB3	1.53	0.89
1:QA:960:U:O4	19:QS:78:ARG:HB3	1.71	0.89
1:QA:1123:A:H4'	10:QJ:37:PRO:HD3	0.90	0.89
1:QA:1358:U:H5''	14:QN:35:ARG:N	1.86	0.89
1:QA:253:U:OP1	17:QQ:67:LYS:NZ	2.06	0.89
1:QA:979:C:N3	14:QN:19:ARG:CD	2.33	0.89
1:XA:1225:A:H1'	19:XS:78:ARG:NE	1.87	0.89
28:R4:34:GLU:OE1	39:RG:113:ARG:CZ	2.21	0.89
41:RI:83:ALA:O	41:RI:89:TYR:CZ	2.26	0.89
1:XA:1188:A:H1'	14:YN:60:SER:CA	2.01	0.89
25:R1:90:ILE:CG2	25:R1:94:LEU:HD11	2.03	0.89
27:R3:42:ALA:O	34:RA:851:U:O2'	1.90	0.89
1:QA:1100:C:H5	2:QB:96:ARG:NH2	1.67	0.89
1:XA:1112:C:C1'	3:XC:179:ARG:NH2	2.35	0.89
1:XA:1190:G:H5''	3:XC:176:HIS:CE1	1.76	0.89
43:YO:34:THR:HG22	43:YO:35:VAL:H	1.33	0.89
1:QA:982:U:H5''	14:QN:6:LEU:HG	1.55	0.88
1:XA:728:A:H62	15:XO:54:ARG:HD3	1.34	0.88
1:QA:668:G:C4'	15:QO:48:LYS:O	2.20	0.88
1:QA:1253:G:OP1	10:QJ:44:VAL:CG2	2.20	0.88
1:QA:1280:A:C4	10:QJ:41:PRO:HD3	2.08	0.88
1:QA:1359:C:OP1	14:QN:22:THR:HG22	1.73	0.88
1:QA:1360:A:C1'	14:QN:17:LYS:NZ	2.37	0.88
1:QA:1357:A:H5'	10:QJ:45:ARG:NH1	1.89	0.88
2:QB:197:VAL:O	8:QH:68:ARG:NH2	2.06	0.88
3:XC:21:ARG:HG3	10:XJ:92:THR:O	1.74	0.88
1:XA:1254:C:OP1	10:XJ:45:ARG:HD3	1.73	0.88
1:XA:1318:A:C1'	19:XS:11:VAL:HG21	2.04	0.88
10:XJ:47:PHE:HZ	14:YN:36:PHE:CD2	1.90	0.88
1:XA:1228:C:OP1	13:XM:108:ARG:CZ	2.22	0.88
1:QA:1309:G:H5''	13:QM:77:ASN:HD21	1.20	0.88
4:QD:88:VAL:HG13	5:QE:97:GLY:CA	2.03	0.88
1:XA:728:A:N7	15:XO:54:ARG:NH1	2.22	0.88
6:XF:94:GLN:CD	18:XR:32:ARG:HD3	1.95	0.88
34:YA:2085:C:C2	34:YA:2235:G:N2	2.42	0.88
1:QA:625:G:H4'	16:QP:16:HIS:HB2	1.55	0.87
10:QJ:50:ILE:CG1	14:QN:41:ARG:NE	2.31	0.87
11:XK:116:HIS:CE1	18:XR:82:THR:HB	2.08	0.87
1:QA:741:G:P	15:QO:39:LEU:HD12	2.13	0.87

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:956:U:C3'	19:QS:80:TYR:O	2.21	0.87
1:QA:980:C:H4'	14:QN:19:ARG:HH21	1.38	0.87
1:QA:986:A:H4'	19:QS:55:LYS:HG3	1.56	0.87
34:YA:2099:U:O4	34:YA:2190:G:O6	1.92	0.87
1:QA:975:A:C3'	14:QN:32:SER:HA	2.03	0.87
34:YA:2090:G:C6	34:YA:2230:G:C6	2.57	0.87
1:QA:948:C:C5'	13:QM:101:GLN:HG2	2.04	0.87
1:QA:1131:G:H1	1:QA:1143:G:H21	1.22	0.87
1:XA:1318:A:C2'	19:XS:11:VAL:CG2	2.28	0.87
1:QA:310:G:OP1	16:QP:27:LYS:HD3	1.74	0.87
1:QA:957:U:C1'	19:QS:79:THR:O	2.22	0.87
1:QA:980:C:N1	14:QN:19:ARG:CG	2.36	0.87
34:YA:2053:G:N2	34:YA:2617:C:C2	2.43	0.87
34:YA:2126:A:H61	34:YA:2163:C:H1'	1.37	0.87
1:QA:1307:U:P	13:QM:99:ARG:CG	2.63	0.87
1:QA:1309:G:H5'	13:QM:77:ASN:ND2	1.87	0.87
1:XA:539:A:OP1	12:XL:114:LYS:HD2	1.75	0.87
11:XK:91:ARG:NH2	18:XR:88:LYS:HD2	1.88	0.87
1:QA:958:A:OP1	19:QS:79:THR:CG2	2.18	0.87
1:QA:1374:A:O2'	7:QG:31:MET:SD	2.31	0.87
1:XA:1228:C:H3'	13:XM:104:ARG:O	1.74	0.87
34:YA:389:G:N1	44:YP:71:VAL:HG12	1.90	0.87
1:QA:1358:U:C5'	14:QN:35:ARG:N	2.37	0.86
3:QC:79:ARG:CZ	11:XK:104:GLN:HA	2.04	0.86
1:XA:881:G:OP2	12:XL:9:GLN:NE2	2.08	0.86
21:XU:3:LYS:CD	21:XU:14:TRP:CE3	2.57	0.86
1:QA:880:C:H5''	12:QL:12:ARG:HH21	1.35	0.86
7:XG:16:LEU:HD23	9:XI:44:VAL:CG2	2.05	0.86
1:QA:676:A:H1'	11:QK:118:GLY:HA3	1.55	0.86
10:QJ:49:VAL:HG23	14:QN:41:ARG:HB2	1.55	0.86
34:RA:1411:C:H42	34:RA:1591:G:H1	1.20	0.86
41:RI:83:ALA:C	41:RI:89:TYR:CE2	2.49	0.86
1:XA:1131:G:H1	1:XA:1143:G:H21	1.22	0.86
34:YA:2043:C:N4	34:YA:2777:G:C6	2.43	0.86
1:QA:1106:G:O3'	3:QC:172:ARG:HB3	1.75	0.86
3:QC:79:ARG:HE	11:XK:104:GLN:HG3	1.39	0.86
1:QA:44:G:P	16:QP:12:LYS:HD3	2.15	0.86
1:QA:974:A:O5'	14:QN:31:ARG:CB	2.24	0.86
15:QO:89:GLY:OXT	34:RA:716:A:OP1	1.92	0.86
34:YA:918:A:N6	34:YA:2268:A:H62	1.71	0.86
14:YN:56:VAL:CG1	14:YN:57:ARG:H	1.85	0.86

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:1493:C:C4	34:YA:2210:G:C8	2.64	0.86
1:QA:1150:U:O2	10:QJ:39:PRO:CG	2.23	0.86
1:QA:1220:G:H21	19:QS:54:GLY:CA	1.87	0.86
1:QA:1313:U:C3'	19:QS:6:LYS:HZ3	1.74	0.86
1:XA:1187:G:H2'	14:XN:61:TRP:O	1.75	0.86
12:QL:7:ILE:HD11	17:QQ:32:TYR:CD1	1.98	0.86
1:XA:1309:G:H4'	13:XM:77:ASN:ND2	1.87	0.86
1:XA:1312:G:C3'	19:XS:6:LYS:HZ3	1.72	0.86
1:XA:1360:A:C4	14:XN:17:LYS:HG3	2.11	0.85
28:R4:5:ILE:HB	39:RG:67:LYS:HD2	1.59	0.85
40:RH:103:LEU:HD11	40:RH:123:PHE:CE1	2.09	0.85
1:XA:1230:C:N4	13:XM:102:ARG:NH1	2.21	0.85
1:QA:1160:G:C5'	2:QB:132:LYS:CE	2.53	0.85
1:QA:1229:A:N1	13:QM:104:ARG:NH2	2.23	0.85
1:QA:1313:U:H3'	19:QS:6:LYS:HZ1	1.35	0.85
1:QA:1541:U:OP2	2:QB:23:ARG:NE	2.10	0.85
35:RB:75:G:HO2'	54:RZ:85:HIS:HE2	1.22	0.85
1:QA:954:G:N2	19:QS:83:HIS:HE1	1.72	0.85
1:QA:959:A:H61	19:QS:78:ARG:C	1.80	0.85
1:QA:1307:U:OP2	13:QM:99:ARG:HG3	1.76	0.85
10:QJ:45:ARG:CG	14:QN:36:PHE:CZ	2.55	0.85
1:QA:1230:C:N4	13:QM:104:ARG:HD3	1.90	0.85
1:XA:1374:A:O2'	7:XG:31:MET:CE	2.23	0.85
1:QA:625:G:H4'	16:QP:16:HIS:CB	2.06	0.85
4:QD:20:TYR:OH	6:XF:14:LEU:CA	2.25	0.85
4:QD:61:LYS:HE2	4:QD:206:PHE:CE2	2.11	0.85
34:YA:443:A:N7	38:YF:45:ARG:HD3	1.90	0.85
1:QA:617:G:O2'	16:QP:44:THR:HG21	1.77	0.85
1:QA:975:A:C2	14:QN:34:TYR:CD1	2.64	0.85
1:XA:309:G:O2'	1:XA:607:A:N1	2.10	0.85
34:YA:389:G:N2	44:YP:71:VAL:HG12	1.90	0.85
1:QA:1081:G:H5''	5:QE:18:ARG:HD2	1.59	0.85
1:QA:1179:A:H5'	9:QI:83:ARG:NH2	1.79	0.85
1:QA:1360:A:H1'	14:QN:17:LYS:HE3	0.85	0.85
1:XA:393:A:OP1	16:XP:13:HIS:NE2	2.08	0.85
1:XA:1228:C:C5	13:XM:104:ARG:HG2	2.12	0.84
34:YA:199:A:N6	34:YA:2434:A:C5	2.45	0.84
16:QP:37:GLY:HA3	16:QP:50:LYS:O	1.77	0.84
1:XA:1317:C:H5	14:XN:16:PHE:CD1	1.91	0.84
1:QA:656:C:H4'	15:QO:62:GLN:NE2	1.90	0.84
1:QA:979:C:H42	14:QN:19:ARG:HB2	1.42	0.84

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1235:U:O2'	21:QU:3:LYS:HG3	1.77	0.84
1:QA:1307:U:O5'	13:QM:99:ARG:HG3	1.77	0.84
4:XD:205:GLU:OE1	5:XE:107:ARG:NH1	2.09	0.84
32:Y8:12:LYS:NZ	34:YA:247:G:O6	2.10	0.84
1:QA:972:C:P	10:QJ:57:LYS:HG3	2.17	0.84
11:XK:91:ARG:HE	18:XR:88:LYS:CE	1.89	0.84
34:YA:2048:G:N2	34:YA:2621:A:N3	2.26	0.84
1:QA:1106:G:C2'	3:QC:172:ARG:HE	1.89	0.84
34:RA:2751:G:C2	40:RH:3:ARG:CB	2.59	0.84
1:XA:1377:A:OP2	7:XG:94:ARG:NH2	2.09	0.84
1:XA:1305:G:C6	1:XA:1331:G:C5	2.65	0.84
11:QK:13:GLN:HG3	34:RA:2141:G:H5''	1.60	0.84
1:XA:974:A:C1'	14:XN:31:ARG:NH1	2.41	0.84
1:XA:1230:C:N4	13:XM:105:THR:HG21	1.93	0.84
1:QA:1150:U:H1'	10:QJ:39:PRO:CG	2.02	0.84
5:QE:77:PRO:O	8:QH:105:ARG:HD3	1.77	0.84
40:RH:103:LEU:CD1	40:RH:123:PHE:CZ	2.61	0.84
1:XA:675:A:H2	11:XK:118:GLY:C	1.80	0.84
34:YA:2690:C:OP1	46:YR:17:ARG:NH2	2.09	0.84
4:QD:197:PRO:HD3	6:XF:16:GLN:HB2	0.84	0.84
34:RA:534:U:HO2'	49:RU:49:HIS:HD1	1.25	0.84
34:YA:2046:G:C2	34:YA:2623:G:N3	2.46	0.84
40:YH:9:ILE:HG21	40:YH:49:VAL:HB	1.56	0.84
1:QA:958:A:OP1	19:QS:79:THR:HG21	1.76	0.84
1:QA:1307:U:C5'	13:QM:99:ARG:HG3	2.08	0.84
3:QC:22:TRP:HA	10:QJ:93:GLY:CA	2.07	0.84
28:R4:31:ILE:HG21	39:RG:142:PRO:HB2	1.58	0.84
1:XA:186(A):C:O2'	20:XT:85:MET:SD	2.35	0.84
1:XA:1217:C:P	14:XN:5:ALA:HB1	2.18	0.84
34:YA:2082:A:H62	34:YA:2237:G:H21	0.86	0.84
1:QA:718:G:O5'	11:QK:117:ASN:OD1	1.96	0.83
34:YA:389:G:H1	44:YP:71:VAL:HG12	1.42	0.83
1:QA:974:A:OP2	14:QN:29:ARG:CD	2.25	0.83
1:XA:1376:U:C4	7:XG:10:ARG:NE	2.46	0.83
34:YA:1127:A:C2	34:YA:2518:A:C8	2.65	0.83
3:QC:23:TYR:CB	10:QJ:93:GLY:C	2.46	0.83
10:XJ:39:PRO:HA	10:XJ:70:ARG:HG2	1.57	0.83
34:YA:747:U:C5	34:YA:2613:U:C4	2.65	0.83
34:YA:1127:A:N1	34:YA:2518:A:C6	2.45	0.83
34:YA:1456:G:N2	34:YA:2703:C:N3	2.24	0.83
1:QA:1358:U:H5'	14:QN:35:ARG:H	1.41	0.83

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1318:A:N3	14:XN:16:PHE:CZ	2.46	0.83
21:XU:3:LYS:HD3	21:XU:14:TRP:CZ3	2.14	0.83
34:YA:2049:G:C2	34:YA:2620:C:C2	2.66	0.83
1:QA:675:A:C2	11:QK:118:GLY:HA2	2.12	0.83
1:XA:1377:A:N7	7:XG:10:ARG:HD2	1.92	0.83
1:QA:626:U:H4'	16:QP:38:TYR:CE2	2.12	0.83
1:QA:1123:A:O4'	10:QJ:37:PRO:HD2	1.78	0.83
1:XA:107:G:N7	20:XT:15:ARG:NH2	2.27	0.83
1:XA:618:C:H5'	1:XA:619:U:H5''	1.58	0.83
10:XJ:53:PRO:HB3	14:XN:42:ILE:HG12	1.58	0.83
34:YA:747:U:C5	34:YA:2613:U:O4	2.32	0.83
34:YA:2044:C:C4	34:YA:2625:G:C2	2.66	0.83
34:YA:2048:G:C6	34:YA:2621:A:N1	2.46	0.83
1:QA:948:C:P	13:QM:106:ASN:O	2.36	0.83
1:QA:1106:G:H1'	3:QC:172:ARG:HE	0.68	0.83
1:QA:1108:G:P	3:QC:174:PRO:HA	2.19	0.83
1:QA:1253:G:OP1	10:QJ:44:VAL:HG21	1.79	0.83
1:QA:520:A:O2'	12:QL:73:GLU:OE2	1.97	0.83
1:QA:669:U:C1'	15:QO:46:HIS:CE1	2.45	0.83
1:QA:1248:A:H62	21:QU:26:LYS:HD2	1.44	0.83
4:QD:205:GLU:CD	5:QE:100:VAL:HG13	1.99	0.83
11:QK:71:LYS:HZ3	34:RA:2146:C:H41	1.23	0.83
34:RA:1971:A:C8	36:RD:241:PRO:HB3	2.14	0.83
1:XA:1254:C:P	10:XJ:45:ARG:HD3	2.18	0.83
34:RA:956:G:OP2	45:RQ:14:ARG:NH2	2.11	0.82
34:RA:2312:U:O2	39:RG:40:ASN:ND2	2.12	0.82
34:YA:2094:G:P	41:YI:22:LYS:HD2	2.18	0.82
1:QA:309:G:H5''	16:QP:27:LYS:CE	2.09	0.82
1:QA:980:C:H4'	14:QN:19:ARG:NH2	1.94	0.82
1:QA:1243:C:H5''	21:QU:9:ARG:HG3	1.59	0.82
34:YA:2045:C:N3	34:YA:2624:G:N2	2.26	0.82
34:YA:2078:C:N4	34:YA:2238:G:O6	2.11	0.82
1:QA:1187:G:C2	14:QN:60:SER:OG	2.32	0.82
4:QD:204:ILE:HG21	5:QE:98:THR:C	2.00	0.82
12:QL:7:ILE:HD12	17:QQ:32:TYR:CD1	2.11	0.82
1:XA:974:A:N9	14:XN:31:ARG:HD3	1.92	0.82
1:XA:1377:A:P	7:XG:94:ARG:HH22	1.97	0.82
1:QA:44:G:OP2	16:QP:12:LYS:NZ	2.13	0.82
1:QA:986:A:H4'	19:QS:55:LYS:CG	2.10	0.82
34:RA:2094:G:OP1	41:RI:22:LYS:HE3	1.78	0.82
1:XA:1305:G:N1	1:XA:1331:G:C5	2.47	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:980:C:N1	14:QN:19:ARG:HG2	1.94	0.82
1:QA:1114:C:H1'	14:QN:60:SER:O	1.78	0.82
1:QA:1307:U:C5'	13:QM:99:ARG:CG	2.57	0.82
34:YA:2046:G:C4	34:YA:2623:G:C2	2.65	0.82
34:YA:2070:G:C2	34:YA:2442:C:N3	2.46	0.82
34:YA:2097:C:N3	34:YA:2192:G:O6	2.11	0.82
1:QA:980:C:C4'	14:QN:19:ARG:HE	1.93	0.82
1:XA:619:U:H5'	4:XD:131:ARG:HH21	1.44	0.82
4:QD:57:ARG:HH22	5:QE:107:ARG:HD3	1.01	0.82
1:QA:1049:U:H5	14:QN:3:ARG:CB	1.71	0.82
1:QA:1223:C:OP2	19:QS:78:ARG:NH2	2.12	0.82
1:QA:1313:U:C4	19:QS:4:SER:HB2	2.15	0.82
10:QJ:47:PHE:CE2	14:QN:37:PHE:HD2	1.98	0.82
1:QA:310:G:OP2	16:QP:27:LYS:HD3	1.78	0.82
1:QA:1221:G:C4'	19:QS:36:ARG:HH21	1.87	0.82
1:XA:986:A:H1'	19:XS:52:TYR:OH	1.80	0.81
1:QA:49:U:H3	1:QA:362:G:H1'	1.44	0.81
4:QD:204:ILE:HG21	5:QE:98:THR:O	1.80	0.81
11:QK:71:LYS:NZ	34:RA:2146:C:N4	2.28	0.81
34:RA:2751:G:C8	40:RH:2:SER:O	2.33	0.81
1:XA:1305:G:C2	1:XA:1331:G:C5	2.68	0.81
3:XC:23:TYR:HE2	10:XJ:95:GLU:HB2	1.45	0.81
34:YA:918:A:C5	34:YA:2268:A:N6	2.47	0.81
1:QA:9:G:H5''	5:QE:122:GLU:OE2	1.81	0.81
29:R5:9:LYS:NZ	34:RA:2019:A:N7	2.29	0.81
34:YA:2090:G:N2	34:YA:2230:G:C4	2.48	0.81
1:XA:1320:C:C6	19:XS:37:ARG:HA	2.16	0.81
1:QA:1243:C:C5'	21:QU:9:ARG:HG3	2.09	0.81
34:YA:389:G:C2	44:YP:71:VAL:HG12	2.15	0.81
3:QC:23:TYR:CE2	10:QJ:95:GLU:HB3	2.15	0.81
34:RA:1294:U:O2'	46:RR:23:ASN:OD1	1.98	0.81
34:YA:199:A:C6	34:YA:2434:A:N1	2.49	0.81
34:YA:919:G:C2	34:YA:2268:A:C5	2.69	0.81
1:XA:6:G:N2	5:XE:119:LEU:HD11	1.96	0.81
40:YH:9:ILE:CG2	40:YH:49:VAL:HB	2.11	0.81
1:QA:1188:A:C3'	14:QN:58:LYS:NZ	2.44	0.81
25:R1:90:ILE:HG22	25:R1:94:LEU:CD1	2.06	0.81
1:XA:1112:C:H1'	3:XC:179:ARG:HE	1.45	0.81
29:Y5:45:VAL:HG21	29:Y5:58:LEU:HD21	1.63	0.81
34:YA:1493:C:C5	34:YA:2210:G:C5	2.69	0.81
1:QA:676:A:C4'	11:QK:115:PRO:HB3	2.11	0.80

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1358:U:O3'	14:QN:22:THR:CG2	2.27	0.80
4:QD:204:ILE:HG22	5:QE:98:THR:O	1.81	0.80
1:QA:957:U:H5'	19:QS:81:ARG:HG3	1.60	0.80
1:QA:1188:A:O3'	14:QN:58:LYS:NZ	2.13	0.80
1:QA:1248:A:H62	21:QU:26:LYS:CD	1.94	0.80
4:QD:61:LYS:HE2	4:QD:206:PHE:HE2	1.45	0.80
40:RH:98:LEU:HD21	40:RH:125:VAL:HG11	0.81	0.80
15:XO:88:ARG:NH1	34:YA:713:G:OP2	2.13	0.80
34:YA:389:G:H22	44:YP:71:VAL:HG12	1.44	0.80
1:QA:277:C:OP1	17:QQ:68:ARG:NH2	2.15	0.80
1:QA:1106:G:C5'	3:QC:172:ARG:HG3	2.11	0.80
34:YA:814:C:H41	44:YP:25:SER:HA	1.46	0.80
1:QA:1160:G:C5'	2:QB:132:LYS:HE2	2.10	0.80
1:XA:1318:A:O2'	19:XS:11:VAL:CB	2.30	0.80
34:YA:2090:G:C2	34:YA:2230:G:N1	2.48	0.80
1:XA:1315:U:O3'	14:YN:17:LYS:CE	2.28	0.80
13:XM:84:ILE:HG22	19:XS:74:PHE:CE1	2.14	0.80
1:QA:1357:A:H5''	10:QJ:45:ARG:HH22	1.45	0.80
1:QA:1360:A:C8	14:QN:18:VAL:HA	2.16	0.80
34:YA:2044:C:N3	34:YA:2625:G:N2	2.30	0.80
34:YA:2048:G:C6	34:YA:2621:A:C2	2.69	0.80
40:RH:89:ILE:CD1	40:RH:131:VAL:HG22	2.12	0.80
41:RI:84:GLY:HA3	41:RI:89:TYR:OH	1.82	0.80
34:YA:2051:A:C2	34:YA:2614:A:N1	2.50	0.80
1:QA:1307:U:H5''	13:QM:99:ARG:CG	2.11	0.80
34:YA:1614:A:N6	51:YW:91:GLY:HA2	1.97	0.80
34:YA:2051:A:C2	34:YA:2614:A:C6	2.70	0.80
1:QA:656:C:H4'	15:QO:62:GLN:HE22	1.45	0.80
1:QA:981:U:C5'	14:QN:6:LEU:CD2	2.60	0.80
1:QA:1179:A:H5''	9:QI:83:ARG:HH22	1.06	0.80
4:QD:205:GLU:CD	5:QE:100:VAL:CG1	2.50	0.80
10:QJ:50:ILE:CD1	14:QN:41:ARG:NE	2.44	0.80
34:YA:2087:G:C6	34:YA:2233:U:C2	2.70	0.80
1:QA:1320:C:C4'	19:QS:70:LYS:HG3	2.13	0.79
13:QM:84:ILE:CG2	19:QS:69:HIS:CE1	2.65	0.79
6:XF:100:ASN:CG	18:XR:27:GLY:HA2	2.01	0.79
34:RA:2751:G:C5	40:RH:3:ARG:CB	2.56	0.79
1:XA:609:A:OP1	16:XP:18:ARG:NH2	2.16	0.79
1:XA:1320:C:H41	19:XS:37:ARG:HB3	0.94	0.79
1:QA:617:G:H4'	16:QP:44:THR:HB	1.64	0.79
1:QA:958:A:N9	19:QS:55:LYS:HD2	1.97	0.79

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1106:G:H4'	3:QC:172:ARG:CD	2.13	0.79
1:QA:1204:A:OP1	14:QN:3:ARG:NH2	2.13	0.79
1:QA:1321:C:N3	19:QS:36:ARG:NH1	2.30	0.79
1:XA:1188:A:O2'	14:XN:58:LYS:HE3	1.82	0.79
1:XA:1240:U:C2'	7:XG:38:LEU:HD11	2.12	0.79
6:XF:100:ASN:ND2	18:XR:27:GLY:O	2.15	0.79
34:YA:1782:C:N4	34:YA:2587:A:H2	1.80	0.79
1:QA:974:A:P	14:QN:29:ARG:CD	2.70	0.79
3:QC:23:TYR:HD2	10:QJ:95:GLU:HB2	1.43	0.79
13:XM:84:ILE:HD11	19:XS:65:ASN:OD1	1.83	0.79
1:QA:1280:A:C4	10:QJ:41:PRO:CD	2.65	0.79
4:QD:208:SER:HB2	5:QE:101:ILE:HD12	1.65	0.79
34:RA:1754:C:H5	48:RT:96:ARG:HH22	1.28	0.79
1:XA:339:C:OP2	43:YO:97:ARG:NH1	2.15	0.79
1:QA:986:A:N3	19:QS:52:TYR:OH	2.15	0.79
29:R5:9:LYS:NZ	34:RA:2019:A:C8	2.49	0.79
11:XK:91:ARG:CZ	18:XR:88:LYS:NZ	2.46	0.79
34:YA:2071:A:C2	34:YA:2441:C:N3	2.51	0.79
1:QA:981:U:H5'	14:QN:21:TYR:OH	1.80	0.79
1:QA:1221:G:C2'	19:QS:77:THR:HG21	2.10	0.79
1:QA:1223:C:P	19:QS:78:ARG:HH21	1.91	0.79
3:QC:22:TRP:HA	10:QJ:93:GLY:HA2	1.65	0.79
34:RA:508:G:O6	51:RW:9:TYR:CE1	2.35	0.79
1:XA:1318:A:O3'	19:XS:11:VAL:CB	2.31	0.79
1:QA:742:G:C5'	15:QO:58:MET:SD	2.71	0.79
1:QA:1355:G:H1	1:QA:1367:C:H5	1.29	0.79
2:QB:178:ARG:NE	8:QH:71:GLY:O	2.16	0.79
4:QD:18:LYS:NZ	56:QD:301:SF4:S4	2.55	0.79
34:RA:2751:G:N3	40:RH:3:ARG:CB	2.42	0.79
1:XA:1124:G:H4'	10:XJ:36:GLY:N	1.97	0.79
34:YA:2088:G:N1	34:YA:2232:U:O2	2.15	0.79
1:QA:875:C:O2'	8:QH:14:ARG:NH1	2.15	0.79
1:XA:1254:C:H5''	10:XJ:45:ARG:CZ	2.13	0.79
6:XF:100:ASN:HB2	18:XR:28:GLU:N	1.97	0.79
22:XV:19:G:N7	34:YA:2112:G:N7	2.31	0.79
29:Y5:9:LYS:NZ	34:YA:2019:A:OP2	2.15	0.79
1:QA:957:U:C3'	19:QS:79:THR:O	2.31	0.78
4:QD:57:ARG:CZ	5:QE:107:ARG:CD	2.55	0.78
10:QJ:49:VAL:HG23	14:QN:41:ARG:CB	2.14	0.78
1:XA:1313:U:C4	19:XS:4:SER:OG	2.36	0.78
1:XA:1320:C:N1	19:XS:70:LYS:HD3	1.99	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:XK:108:ILE:O	18:XR:87:ARG:N	2.15	0.78
34:YA:1818:U:O4	36:YD:154:LYS:NZ	2.14	0.78
1:QA:668:G:H4'	15:QO:48:LYS:HB3	1.63	0.78
1:QA:972:C:OP2	10:QJ:57:LYS:CG	2.30	0.78
1:QA:978:A:N6	14:QN:18:VAL:CG2	2.37	0.78
1:QA:1309:G:H5'	13:QM:77:ASN:CG	2.02	0.78
34:YA:199:A:C6	34:YA:2434:A:C6	2.71	0.78
13:QM:84:ILE:HG13	19:QS:74:PHE:HZ	1.49	0.78
1:XA:1377:A:P	7:XG:94:ARG:HH21	2.03	0.78
1:QA:624:C:C3'	16:QP:10:GLY:HA2	2.13	0.78
19:QS:61:TYR:CE1	34:RA:888:C:OP2	2.36	0.78
34:RA:2250:G:C4	45:RQ:82:ARG:HG3	2.17	0.78
1:XA:1237:C:O2'	1:XA:1300:G:N2	2.16	0.78
1:XA:1318:A:N1	14:YN:16:PHE:CZ	2.50	0.78
1:XA:1377:A:C6	7:XG:10:ARG:CD	2.65	0.78
13:XM:84:ILE:CG2	19:XS:74:PHE:CZ	2.66	0.78
34:YA:747:U:C4	34:YA:2613:U:N3	2.52	0.78
1:QA:1109:C:P	3:QC:176:HIS:CD2	2.77	0.78
34:RA:602:G:HO2'	34:RA:604:G:HO2'	1.28	0.78
34:RA:675:A:O2'	38:RF:67:GLN:NE2	2.16	0.78
40:RH:98:LEU:CG	40:RH:125:VAL:HG11	2.13	0.78
1:XA:520:A:O2'	12:XL:73:GLU:OE1	2.02	0.78
34:YA:1566:A:C6	36:YD:214:TRP:CH2	2.72	0.78
34:YA:2108:C:N4	34:YA:2182:G:N2	2.29	0.78
1:QA:1124:G:O5'	10:QJ:36:GLY:N	2.16	0.78
28:R4:5:ILE:O	39:RG:67:LYS:HG3	1.83	0.78
1:XA:728:A:N6	15:XO:54:ARG:CD	2.47	0.78
34:YA:1050:A:N7	34:YA:2751:G:C2	2.52	0.78
1:QA:1109:C:P	3:QC:176:HIS:HD2	2.07	0.78
1:XA:675:A:N3	11:XK:118:GLY:HA2	1.98	0.78
1:XA:1321:C:H5'	1:XA:1322:C:H5''	1.66	0.78
1:QA:957:U:H5'	19:QS:81:ARG:CG	2.14	0.78
40:RH:98:LEU:CD2	40:RH:125:VAL:CG1	2.41	0.78
1:XA:1229:A:H62	13:XM:104:ARG:HE	1.32	0.78
1:QA:1179:A:H5'	9:QI:83:ARG:HH12	1.49	0.78
1:QA:1236:A:H4'	21:QU:10:ARG:CZ	2.09	0.78
1:QA:1360:A:C4'	14:QN:17:LYS:HZ2	1.97	0.78
19:QS:61:TYR:CZ	34:RA:888:C:OP2	2.37	0.78
29:R5:3:LYS:HG2	34:RA:2611:U:C4	2.19	0.78
1:XA:728:A:C8	15:XO:54:ARG:CZ	2.67	0.78
1:XA:1112:C:H1'	3:XC:179:ARG:CZ	2.13	0.78

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:608:A:H1'	16:QP:32:TYR:HE1	1.48	0.78
10:QJ:50:ILE:HD13	14:QN:41:ARG:CZ	2.13	0.78
1:XA:8:A:C6	4:XD:209:ARG:HA	2.19	0.78
1:XA:740:U:OP2	15:XO:2:PRO:HB3	1.83	0.78
1:QA:979:C:H2'	14:QN:19:ARG:NH2	1.98	0.77
1:QA:1226:C:H5'	13:QM:91:ARG:NH1	1.98	0.77
19:QS:65:ASN:O	19:QS:66:MET:HE2	1.84	0.77
27:R3:25:ALA:HB2	34:RA:849:A:C2	2.19	0.77
28:R4:34:GLU:OE1	39:RG:113:ARG:NH1	2.17	0.77
1:XA:1253:G:C2	1:XA:1254:C:C5	2.72	0.77
3:XC:23:TYR:CD2	10:XJ:95:GLU:HB2	2.19	0.77
1:QA:9:G:OP2	5:QE:122:GLU:HG2	1.84	0.77
3:XC:60:ALA:HB1	10:XJ:91:PRO:CD	2.13	0.77
1:QA:958:A:C1'	19:QS:55:LYS:CD	2.61	0.77
11:XK:91:ARG:HG2	18:XR:88:LYS:HZ1	1.47	0.77
34:YA:1493:C:C5	34:YA:2210:G:N7	2.52	0.77
40:RH:87:LEU:O	40:RH:131:VAL:CG2	2.32	0.77
1:QA:44:G:OP1	16:QP:12:LYS:HD3	1.84	0.77
1:QA:752:G:H4'	15:QO:69:TYR:OH	1.84	0.77
1:QA:958:A:C8	19:QS:55:LYS:HD2	2.20	0.77
1:QA:1240:U:C2'	7:QG:38:LEU:HD11	2.14	0.77
3:QC:23:TYR:HB2	10:QJ:94:VAL:N	1.98	0.77
34:RA:99:U:O4	53:RY:8:LYS:NZ	2.16	0.77
1:XA:1152:A:H5''	10:XJ:13:HIS:CD2	2.19	0.77
34:YA:1566:A:H2	36:YD:214:TRP:CD1	2.02	0.77
1:QA:979:C:C4	14:QN:19:ARG:CD	2.67	0.77
1:QA:1236:A:O5'	21:QU:10:ARG:NH1	2.11	0.77
1:QA:1253:G:H5'	10:QJ:44:VAL:H	1.50	0.77
1:QA:1302:U:OP2	13:QM:27:LYS:HE2	1.85	0.77
28:R4:31:ILE:HG23	39:RG:142:PRO:O	1.85	0.77
34:RA:1534:G:H2'	34:RA:1535:U:H4'	1.66	0.77
1:QA:740:U:C4'	15:QO:39:LEU:HG	2.15	0.77
1:QA:1331:G:N1	21:QU:5:ASP:OD1	2.18	0.77
1:XA:957:U:H4'	19:XS:79:THR:OG1	1.85	0.77
34:YA:270(P):U:C4	41:YI:52:ARG:NH2	2.52	0.77
34:YA:919:G:N1	34:YA:2268:A:C5	2.53	0.77
12:QL:7:ILE:CD1	17:QQ:32:TYR:CG	2.59	0.77
13:QM:84:ILE:HG13	19:QS:74:PHE:CZ	2.19	0.77
1:XA:367:U:H5'	1:XA:394:G:H21	1.50	0.77
1:XA:1152:A:H5''	10:XJ:13:HIS:NE2	2.00	0.77
1:QA:1125:U:N3	10:QJ:73:ASP:OD1	2.18	0.77

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1305:G:H4'	1:QA:1332:A:H62	1.50	0.77
1:QA:1313:U:C6	19:QS:6:LYS:CE	2.68	0.77
34:RA:586:A:H5'	38:RF:89:VAL:HG21	1.67	0.77
34:RA:2294:C:OP2	47:RS:13:ARG:NH2	2.17	0.77
1:XA:1357:A:OP1	14:XN:35:ARG:NH1	2.08	0.77
1:XA:1377:A:C5	7:XG:10:ARG:HD2	2.20	0.77
29:Y5:43:HIS:CD2	34:YA:2884:U:C5	2.72	0.77
1:QA:978:A:H62	14:QN:18:VAL:HG21	0.72	0.77
1:QA:1378:C:OP2	7:QG:7:ALA:HB2	1.80	0.77
1:XA:1058:G:O5'	3:XC:2:GLY:N	2.18	0.77
1:QA:974:A:OP1	14:QN:29:ARG:CZ	2.34	0.76
1:QA:980:C:C1'	14:QN:19:ARG:NE	2.46	0.76
1:QA:1330:U:C4	21:QU:7:ARG:NH1	2.53	0.76
3:QC:79:ARG:NH2	11:XK:105:VAL:H	1.82	0.76
10:QJ:50:ILE:CG1	14:QN:41:ARG:HD2	2.04	0.76
34:RA:1789:A:OP1	36:RD:222:ARG:HG3	1.84	0.76
1:XA:1318:A:O3'	19:XS:11:VAL:HB	1.84	0.76
43:YO:34:THR:HG22	43:YO:35:VAL:N	1.99	0.76
34:RA:2483:C:O2	45:RQ:124:LYS:NZ	2.14	0.76
1:XA:1305:G:C4	1:XA:1331:G:C6	2.72	0.76
1:QA:1240:U:H3	7:QG:38:LEU:HB3	1.49	0.76
1:QA:1360:A:N6	14:QN:18:VAL:CG2	2.29	0.76
50:RV:45:THR:O	50:RV:45:THR:HG22	1.83	0.76
34:YA:442:G:H1'	38:YF:48:THR:HG21	1.67	0.76
34:YA:2095:C:O2	34:YA:2194:G:N2	2.15	0.76
1:QA:1236:A:O4'	21:QU:10:ARG:NH1	2.17	0.76
41:RI:92:VAL:O	41:RI:120:ILE:HB	1.85	0.76
1:XA:634:C:H2'	1:XA:635:G:H8	1.50	0.76
34:YA:2069:G:C2	34:YA:2443:C:C2	2.73	0.76
54:YZ:183:LEU:HD23	54:YZ:183:LEU:O	1.85	0.76
25:R1:90:ILE:HA	25:R1:94:LEU:HD12	1.67	0.76
1:QA:761:G:H5'	17:QQ:100:LYS:HZ3	1.49	0.76
1:QA:1240:U:N3	7:QG:38:LEU:CB	2.30	0.76
1:QA:1358:U:H4'	14:QN:33:VAL:O	1.86	0.76
1:XA:1503:A:O2'	23:XX:15:A:N6	2.19	0.76
11:XK:91:ARG:CG	18:XR:88:LYS:NZ	2.48	0.76
1:QA:1152:A:OP1	10:QJ:13:HIS:CD2	2.38	0.76
34:RA:2108:C:HO2'	34:RA:2142:C:HO2'	1.34	0.76
34:RA:2751:G:C2	40:RH:3:ARG:CG	2.66	0.76
1:XA:1240:U:C2'	7:XG:38:LEU:CD1	2.63	0.76
34:YA:2070:G:N2	34:YA:2442:C:O2	2.19	0.76

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:974:A:C4	14:QN:31:ARG:CZ	2.68	0.76
1:QA:1179:A:H5'	9:QI:83:ARG:NH1	2.01	0.76
1:QA:1320:C:O4'	19:QS:70:LYS:HG3	1.84	0.76
2:QB:178:ARG:O	8:QH:72:PRO:HD3	1.86	0.76
1:QA:624:C:O3'	16:QP:10:GLY:CA	2.30	0.76
29:R5:15:ARG:NH2	34:RA:2022:U:OP2	2.17	0.76
7:XG:16:LEU:HD23	9:XI:44:VAL:HG22	1.68	0.76
1:XA:1160:G:C1'	2:XB:132:LYS:HE3	2.15	0.75
1:XA:1230:C:N4	13:XM:105:THR:CB	2.33	0.75
13:XM:84:ILE:CG1	19:XS:65:ASN:O	2.32	0.75
4:QD:208:SER:CB	5:QE:101:ILE:HD12	2.16	0.75
1:XA:421:U:C4'	3:XC:192:THR:HG21	2.15	0.75
34:YA:2054:A:C2	34:YA:2616:C:N3	2.54	0.75
34:YA:2088:G:C2	34:YA:2232:U:O2	2.39	0.75
34:RA:1049:C:N3	40:RH:2:SER:HB3	2.01	0.75
1:QA:1203:C:H5'	14:QN:3:ARG:NH1	2.01	0.75
10:QJ:79:ARG:NH1	10:QJ:79:ARG:O	2.20	0.75
34:RA:2751:G:C4	40:RH:3:ARG:CG	2.68	0.75
1:XA:1318:A:N6	14:YN:16:PHE:CD2	2.48	0.75
1:XA:1320:C:C4	19:XS:37:ARG:N	2.55	0.75
1:QA:739:C:O2'	15:QO:42:HIS:CG	2.40	0.75
10:QJ:50:ILE:HD11	14:QN:41:ARG:CZ	2.12	0.75
41:RI:92:VAL:HB	41:RI:120:ILE:CG2	2.16	0.75
1:XA:728:A:N6	15:XO:54:ARG:HD3	2.00	0.75
3:XC:23:TYR:CD2	10:XJ:95:GLU:HG3	2.22	0.75
34:YA:2053:G:O6	34:YA:2614:A:C2	2.40	0.75
34:YA:2131:G:H4'	34:YA:2132:U:H4'	1.69	0.75
34:RA:2471:C:N4	34:RA:2476:A:O2'	2.20	0.75
1:XA:1360:A:H1'	14:YN:17:LYS:CD	2.16	0.75
33:Y9:27:CYS:SG	33:Y9:29:ASN:N	2.58	0.75
33:Y9:29:ASN:ND2	33:Y9:32:HIS:NE2	2.35	0.75
32:R8:35:GLN:NE2	32:R8:36:LYS:HE2	2.02	0.75
1:XA:1312:G:C2'	19:XS:6:LYS:HZ2	1.98	0.75
34:YA:919:G:C6	34:YA:2268:A:C2	2.69	0.75
34:YA:2087:G:O6	34:YA:2233:U:N3	2.20	0.75
1:QA:959:A:N6	19:QS:78:ARG:C	2.38	0.75
4:QD:205:GLU:OE1	5:QE:107:ARG:NH1	2.20	0.75
1:XA:501:C:H1'	1:XA:549:C:H1'	1.69	0.75
1:XA:974:A:OP2	14:YN:29:ARG:NE	2.20	0.75
1:QA:986:A:O2'	19:QS:55:LYS:C	2.25	0.74
1:QA:1188:A:H4'	14:QN:58:LYS:HZ2	1.52	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:831:G:O2'	44:YP:38:GLN:OE1	2.04	0.74
34:YA:918:A:N6	34:YA:2268:A:N6	2.33	0.74
1:QA:9:G:P	5:QE:122:GLU:HG2	2.26	0.74
34:YA:727:A:H2	36:YD:9:TYR:CD2	2.05	0.74
1:QA:981:U:H5'	14:QN:6:LEU:CD2	2.16	0.74
1:QA:1222:G:H5'	19:QS:77:THR:OG1	1.86	0.74
1:QA:1229:A:N1	13:QM:104:ARG:NE	2.34	0.74
1:QA:1357:A:H5'	10:QJ:45:ARG:HH12	1.51	0.74
1:QA:1360:A:N9	14:QN:17:LYS:HE3	2.01	0.74
31:Y7:37:LYS:NZ	34:YA:469:G:C6	2.55	0.74
34:YA:2584:U:H2'	34:YA:2585:U:H2'	1.70	0.74
1:QA:564:C:H5'	17:QQ:32:TYR:HE1	1.50	0.74
1:QA:608:A:H1'	16:QP:32:TYR:CE1	2.22	0.74
1:XA:1226:C:C3'	13:XM:103:THR:OG1	2.36	0.74
1:QA:1179:A:C5'	9:QI:83:ARG:CZ	2.64	0.74
34:RA:1566:A:C2	36:RD:214:TRP:CD2	2.75	0.74
1:XA:538:G:OP1	12:XL:115:LYS:HB2	1.86	0.74
1:XA:754:C:H6	15:XO:69:TYR:CE2	2.05	0.74
1:QA:1158:C:H4'	2:QB:133:LYS:HZ2	0.92	0.74
1:QA:954:G:N3	19:QS:83:HIS:HE1	1.84	0.74
1:QA:1229:A:H62	13:QM:104:ARG:CG	1.96	0.74
1:QA:1357:A:C5'	10:QJ:45:ARG:NH2	2.39	0.74
4:QD:88:VAL:HG13	5:QE:97:GLY:HA3	1.70	0.74
1:XA:8:A:C5	4:XD:209:ARG:HA	2.23	0.74
1:XA:1240:U:H1'	7:XG:38:LEU:HD11	0.74	0.74
1:XA:1377:A:N6	7:XG:10:ARG:HD2	2.03	0.74
34:YA:1338:G:N7	52:YX:62:LYS:NZ	2.35	0.74
1:QA:642:A:C8	8:QH:115:SER:HA	2.21	0.74
1:QA:922:G:H4'	5:QE:20:GLN:HA	1.70	0.74
10:QJ:45:ARG:HG2	14:QN:36:PHE:CE2	2.21	0.74
1:XA:1252:A:H2	1:XA:1355:G:H1'	1.53	0.74
1:XA:1318:A:C5	14:YN:16:PHE:HE1	1.95	0.74
34:YA:2056:G:C5	34:YA:2577:A:C6	2.76	0.74
1:QA:609:A:H5''	16:QP:9:PHE:CD1	2.23	0.74
11:XK:91:ARG:HG2	18:XR:88:LYS:NZ	2.03	0.74
1:XA:8:A:H62	4:XD:209:ARG:N	1.86	0.74
11:XK:110:ASP:N	18:XR:85:LEU:O	2.20	0.74
1:QA:959:A:N6	19:QS:78:ARG:CA	2.45	0.73
1:QA:980:C:N1	14:QN:19:ARG:HG3	1.99	0.73
1:QA:982:U:O5'	14:QN:6:LEU:CD2	2.30	0.73
1:QA:981:U:O3'	14:QN:23:ARG:NH2	2.21	0.73

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1376:U:C4	7:QG:10:ARG:NH2	2.34	0.73
11:XK:109:VAL:HG22	18:XR:86:VAL:CG2	2.18	0.73
1:QA:862:C:H1'	1:QA:874:G:H5''	1.68	0.73
4:QD:20:TYR:OH	6:XF:14:LEU:HA	1.87	0.73
1:XA:595:G:H1'	1:XA:596:C:H5	1.52	0.73
10:XJ:47:PHE:CZ	14:XN:36:PHE:CD2	2.75	0.73
34:YA:2046:G:C5	34:YA:2623:G:N1	2.55	0.73
1:QA:979:C:C4	14:QN:19:ARG:HB2	2.22	0.73
1:QA:1179:A:H5'	9:QL:83:ARG:CZ	2.18	0.73
28:R4:27:THR:HG21	39:RG:62:LEU:O	1.88	0.73
1:XA:1440(N):G:OP1	20:XT:35:THR:CG2	2.35	0.73
1:QA:982:U:C5'	14:QN:6:LEU:CG	2.62	0.73
1:QA:1219:U:O2'	19:QS:34:TRP:HE3	1.62	0.73
1:XA:1537:U:H3	23:XX:9:G:H1	1.37	0.73
7:XG:16:LEU:HD21	9:XI:44:VAL:HG23	1.69	0.73
34:YA:34:C:H41	34:YA:447:A:H61	1.37	0.73
34:YA:1061:U:H6	34:YA:1062:G:H5''	1.54	0.73
1:QA:609:A:C5'	16:QP:9:PHE:CE1	2.72	0.73
1:QA:642:A:H1'	8:QH:114:THR:O	1.88	0.73
1:QA:979:C:C2	14:QN:19:ARG:CD	2.70	0.73
11:XK:110:ASP:CB	18:XR:85:LEU:O	2.34	0.73
34:YA:919:G:O6	34:YA:2268:A:C2	2.41	0.73
34:YA:1456:G:N2	34:YA:2704:C:C2	2.55	0.73
1:QA:1106:G:C2'	3:QC:172:ARG:NE	2.48	0.73
3:XC:14:ILE:HD11	14:XN:57:ARG:NH2	2.02	0.73
13:XM:86:CYS:HB2	19:XS:73:GLU:OE1	1.89	0.73
1:QA:959:A:H62	19:QS:79:THR:H	1.17	0.73
1:QA:986:A:C4'	19:QS:55:LYS:HG3	2.18	0.73
1:QA:1106:G:H1'	3:QC:172:ARG:CZ	2.18	0.73
34:RA:1510:A:O2'	34:RA:1511:A:N7	2.22	0.73
34:RA:2751:G:C2	40:RH:3:ARG:HB2	2.24	0.73
32:Y8:8:LYS:NZ	34:YA:243:U:OP1	2.22	0.73
1:QA:959:A:H62	19:QS:79:THR:HG23	1.53	0.73
1:QA:1360:A:H1'	14:QN:17:LYS:HE2	1.65	0.73
34:RA:1049:C:N3	40:RH:2:SER:CB	2.52	0.73
1:XA:1217:C:H5''	14:XN:12:ARG:HH12	1.53	0.73
1:XA:1315:U:O3'	14:XN:17:LYS:NZ	2.21	0.73
34:YA:2053:G:O6	34:YA:2614:A:H2	1.71	0.73
1:QA:975:A:C2'	14:QN:32:SER:HA	2.19	0.72
1:QA:1106:G:C2'	3:QC:172:ARG:CD	2.67	0.72
34:YA:2046:G:C2	34:YA:2623:G:N1	2.57	0.72

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:XL:11:VAL:HG23	17:XQ:29:HIS:CD2	2.24	0.72
1:QA:948:C:C5'	13:QM:101:GLN:HG3	2.07	0.72
1:QA:958:A:C1'	19:QS:55:LYS:HD2	2.19	0.72
2:QB:172:ILE:O	2:QB:176:GLU:HB2	1.88	0.72
1:XA:1238:A:H2'	1:XA:1239:A:H8	1.55	0.72
1:XA:1305:G:O6	1:XA:1331:G:O6	2.05	0.72
10:XJ:40:LEU:HB3	10:XJ:41:PRO:HD3	1.71	0.72
27:R3:25:ALA:HB2	34:RA:849:A:H2	1.55	0.72
34:RA:2483:C:C2	45:RQ:124:LYS:NZ	2.57	0.72
1:XA:948:C:OP2	13:XM:106:ASN:CG	2.28	0.72
1:XA:1318:A:O2'	19:XS:11:VAL:HG11	1.90	0.72
1:QA:1049:U:H5	14:QN:3:ARG:HB3	0.92	0.72
1:QA:1307:U:O5'	13:QM:99:ARG:NH1	2.21	0.72
3:QC:79:ARG:NH2	11:XK:104:GLN:CA	2.48	0.72
13:XM:84:ILE:CD1	19:XS:65:ASN:OD1	2.37	0.72
1:QA:657:G:O2'	15:QO:28:GLN:HG3	1.86	0.72
1:QA:667:G:C4'	15:QO:51:HIS:CE1	2.72	0.72
10:QJ:49:VAL:CG2	14:QN:41:ARG:CB	2.66	0.72
12:QL:7:ILE:CD1	17:QQ:32:TYR:HD1	1.93	0.72
1:XA:1107:C:OP1	3:XC:172:ARG:NE	2.22	0.72
1:XA:1226:C:O2'	1:XA:1227:A:N7	2.23	0.72
34:YA:2048:G:N1	34:YA:2621:A:N1	2.38	0.72
4:QD:197:PRO:CG	6:XF:16:GLN:HB2	2.18	0.72
30:R6:23:THR:HG21	34:RA:2286:A:H61	1.55	0.72
1:XA:1187:G:N2	14:XN:60:SER:CB	2.52	0.72
1:XA:1375:A:H4'	7:XG:28:ASN:CG	2.09	0.72
34:YA:919:G:C4	34:YA:2268:A:N6	2.58	0.72
1:QA:959:A:C6	19:QS:77:THR:O	2.42	0.72
1:QA:1235:U:O3'	21:QU:10:ARG:CD	2.38	0.72
34:RA:11:G:H22	34:RA:2627:G:H5''	1.53	0.72
1:XA:1228:C:H41	13:XM:104:ARG:CG	2.02	0.72
1:XA:1320:C:H41	19:XS:37:ARG:CB	1.87	0.72
31:Y7:37:LYS:HE2	34:YA:458:G:C8	2.25	0.72
31:Y7:37:LYS:HE2	34:YA:458:G:N9	2.05	0.72
34:YA:2056:G:O6	34:YA:2577:A:N9	2.22	0.72
1:QA:761:G:C5'	17:QQ:100:LYS:NZ	2.52	0.72
1:QA:955:U:O2'	19:QS:83:HIS:CB	2.38	0.72
35:YB:48:A:OP2	47:YS:30:ARG:NH2	2.23	0.72
1:QA:974:A:OP2	14:QN:29:ARG:HG2	1.90	0.71
3:QC:23:TYR:HE2	10:QJ:95:GLU:OE1	1.73	0.71
1:XA:981:U:H5''	1:XA:982:U:H2'	1.71	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1329:A:N7	21:QU:7:ARG:NH2	2.39	0.71
34:RA:2751:G:C5	40:RH:2:SER:O	2.42	0.71
1:XA:1360:A:H8	14:XN:17:LYS:O	1.63	0.71
1:QA:405:U:H5''	1:QA:495:A:H2	1.56	0.71
1:QA:1240:U:C4	7:QG:38:LEU:HB3	2.24	0.71
36:RD:96:HIS:CE1	36:RD:102:LYS:HE2	2.24	0.71
1:XA:322:C:H4'	20:XT:23:ARG:HG2	1.72	0.71
1:XA:1228:C:C4	13:XM:104:ARG:CG	2.63	0.71
3:XC:14:ILE:HD11	14:XN:57:ARG:HH22	1.52	0.71
1:QA:1128:C:O2'	1:QA:1146:A:N6	2.23	0.71
1:XA:1228:C:C5	13:XM:104:ARG:CG	2.73	0.71
1:QA:617:G:H4'	16:QP:44:THR:CB	2.20	0.71
1:QA:983:A:H5'	14:QN:2:ALA:CB	2.10	0.71
1:QA:1188:A:C3'	14:QN:58:LYS:HZ1	2.02	0.71
28:R4:6:HIS:CE1	39:RG:66:GLN:OE1	2.43	0.71
1:XA:1229:A:H61	13:XM:104:ARG:NE	1.87	0.71
4:XD:89:THR:OG1	5:XE:97:GLY:O	2.03	0.71
1:QA:642:A:N9	8:QH:115:SER:HA	2.05	0.71
1:QA:1329:A:H62	21:QU:7:ARG:NH2	1.88	0.71
13:QM:86:CYS:CB	19:QS:69:HIS:CE1	2.72	0.71
1:XA:1228:C:H41	13:XM:104:ARG:HG2	1.51	0.71
34:YA:2080:G:N2	34:YA:2241:A:N3	2.39	0.71
34:YA:2717:G:O2'	48:YT:96:ARG:NH2	2.23	0.71
1:QA:583:A:H4'	17:QQ:91:ARG:HG2	1.72	0.71
1:QA:624:C:H4'	16:QP:10:GLY:C	2.11	0.71
1:QA:980:C:C1'	14:QN:19:ARG:CD	2.61	0.71
1:XA:1375:A:H4'	7:XG:28:ASN:OD1	1.90	0.71
34:YA:911:A:C2	45:YQ:9:TYR:CG	2.79	0.71
34:YA:2054:A:H2	34:YA:2616:C:C2	2.07	0.71
34:YA:2056:G:C6	34:YA:2577:A:C5	2.79	0.71
1:QA:642:A:H1'	8:QH:114:THR:C	2.11	0.71
1:QA:948:C:OP1	13:QM:101:GLN:HB3	1.88	0.71
1:QA:975:A:O2'	14:QN:32:SER:HA	1.91	0.71
1:QA:1280:A:N3	10:QJ:41:PRO:HD3	2.06	0.71
45:RQ:65:PHE:HB2	45:RQ:105:GLU:HB2	1.72	0.71
1:XA:974:A:C1'	14:XN:31:ARG:HD3	2.19	0.71
1:XA:1229:A:H62	13:XM:104:ARG:NE	1.88	0.71
6:XF:97:PHE:O	18:XR:30:ASP:HA	1.91	0.71
34:YA:2620:C:O2'	37:YE:157:ALA:O	2.08	0.71
1:QA:978:A:H61	1:QA:1316:G:H1'	1.54	0.71
1:QA:1313:U:C5	19:QS:4:SER:HB2	2.26	0.71

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:QI:16:ARG:HB3	9:QI:64:THR:HG23	1.73	0.71
1:XA:1111:A:H2'	3:XC:179:ARG:NH2	2.06	0.71
3:XC:22:TRP:HA	10:XJ:93:GLY:HA2	1.71	0.71
1:QA:186(K):G:N1	1:QA:264:U:H5''	2.05	0.70
1:QA:980:C:C1'	14:QN:19:ARG:HE	2.04	0.70
34:RA:614:U:O4	38:RF:175:THR:OG1	2.08	0.70
1:XA:1535:C:H41	23:XX:10:G:N2	1.88	0.70
34:YA:2043:C:C4	34:YA:2777:G:N3	2.39	0.70
1:QA:761:G:H5'	17:QQ:100:LYS:HZ1	1.55	0.70
1:QA:974:A:P	14:QN:29:ARG:HE	2.07	0.70
1:QA:1307:U:OP2	13:QM:99:ARG:CG	2.38	0.70
1:QA:1320:C:N4	19:QS:37:ARG:HA	2.04	0.70
1:XA:974:A:H1'	14:YN:31:ARG:NH1	2.04	0.70
1:XA:974:A:N3	14:YN:31:ARG:NH1	2.39	0.70
1:XA:1377:A:C6	7:XG:10:ARG:HG3	2.25	0.70
3:XC:23:TYR:HD2	10:XJ:95:GLU:HG3	1.56	0.70
1:QA:583:A:H4'	17:QQ:91:ARG:HD3	1.73	0.70
1:QA:666:G:H21	15:QO:51:HIS:HB2	1.56	0.70
34:RA:839:U:H1'	34:RA:1191:G:H1'	1.73	0.70
1:XA:403:C:H42	55:XA:1688:MG:MG	0.96	0.70
1:XA:974:A:N9	14:YN:31:ARG:NH1	2.39	0.70
1:XA:1059:C:OP2	3:XC:2:GLY:N	2.24	0.70
34:YA:2087:G:C6	34:YA:2233:U:N3	2.59	0.70
1:QA:956:U:C4'	19:QS:83:HIS:HA	2.20	0.70
1:QA:974:A:C4'	14:QN:31:ARG:HB3	2.22	0.70
10:QJ:47:PHE:HE2	14:QN:34:TYR:HB3	1.55	0.70
12:QL:7:ILE:HD13	17:QQ:32:TYR:CD1	2.22	0.70
34:YA:508:G:O6	51:YW:9:TYR:CD1	2.45	0.70
1:QA:186(K):G:N7	17:QQ:63:ARG:CZ	2.54	0.70
41:RI:83:ALA:O	41:RI:89:TYR:CE1	2.45	0.70
2:XB:120:ALA:O	2:XB:124:SER:HB2	1.92	0.70
10:XJ:37:PRO:HB3	10:XJ:72:VAL:CG2	2.21	0.70
1:QA:1253:G:H4'	10:QJ:44:VAL:O	1.89	0.70
1:QA:1290:G:O6	21:QU:26:LYS:HE3	1.91	0.70
34:RA:998:C:OP2	49:RU:58:ARG:NH1	2.25	0.70
4:XD:57:ARG:HB3	4:XD:206:PHE:HB2	1.72	0.70
31:Y7:33:ARG:NH1	34:YA:467:G:OP1	2.24	0.70
1:QA:813:U:H2'	1:QA:814:A:H8	1.57	0.70
1:QA:1080:A:OP1	5:QE:14:ARG:NH2	2.24	0.70
1:QA:1106:G:C4'	3:QC:172:ARG:CD	2.68	0.70
1:QA:1106:G:C1'	3:QC:172:ARG:CZ	2.67	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:2056:G:C8	34:YA:2577:A:N6	2.60	0.70
34:YA:2440:C:H5''	34:YA:2587:A:H4'	1.73	0.70
34:YA:2623:G:HO2'	34:YA:2825:C:HO2'	1.39	0.70
3:QC:22:TRP:CA	10:QJ:93:GLY:HA2	2.22	0.70
1:XA:910:C:OP2	12:XL:21:LYS:NZ	2.20	0.70
1:XA:1111:A:H2'	3:XC:179:ARG:HH22	1.56	0.70
10:XJ:24:VAL:HG21	10:XJ:37:PRO:HG3	1.74	0.70
30:Y6:6:ARG:NH1	34:YA:2285:C:OP2	2.24	0.70
1:QA:186(K):G:H1	1:QA:264:U:H5''	1.56	0.70
1:QA:949:A:OP1	13:QM:101:GLN:CA	2.37	0.70
1:QA:975:A:H2	14:QN:34:TYR:CD1	2.08	0.70
1:QA:1186:G:N2	14:QN:61:TRP:C	2.35	0.70
1:QA:1309:G:H5''	13:QM:77:ASN:HD22	1.51	0.70
1:XA:1330:U:H5''	13:XM:24:GLY:C	2.11	0.70
2:QB:178:ARG:HB3	8:QH:71:GLY:C	2.13	0.70
34:RA:1140:C:O3'	42:RN:25:ARG:NH2	2.24	0.70
6:XF:91:VAL:CG2	18:XR:34:TYR:OH	2.40	0.70
34:YA:685:A:H5''	34:YA:788:A:H62	1.56	0.70
34:YA:2045:C:N3	34:YA:2624:G:N1	2.39	0.70
1:QA:1106:G:O4'	3:QC:172:ARG:CZ	2.40	0.69
1:QA:1307:U:H5''	13:QM:99:ARG:HG2	1.74	0.69
1:QA:1360:A:N6	14:QN:18:VAL:HG21	2.07	0.69
34:RA:685:A:H5''	34:RA:788:A:H62	1.56	0.69
1:XA:1059:C:O2'	10:XJ:53:PRO:HD3	1.91	0.69
3:XC:23:TYR:CD2	10:XJ:95:GLU:CB	2.75	0.69
3:XC:23:TYR:N	10:XJ:93:GLY:HA2	2.06	0.69
34:YA:2046:G:N2	34:YA:2623:G:C4	2.60	0.69
1:QA:583:A:H4'	17:QQ:91:ARG:CD	2.22	0.69
1:QA:1320:C:N3	19:QS:37:ARG:CA	2.48	0.69
35:RB:37:C:O2	47:RS:95:HIS:NE2	2.25	0.69
1:QA:1320:C:H42	19:QS:37:ARG:HG2	0.62	0.69
1:QA:1376:U:C5	7:QG:10:ARG:NH1	2.61	0.69
2:QB:54:THR:HG22	2:QB:199:TYR:HB3	1.74	0.69
4:QD:57:ARG:NH2	5:QE:107:ARG:HH11	1.91	0.69
4:QD:195:ALA:O	6:XF:17:SER:HA	1.93	0.69
1:XA:991:U:H3'	1:XA:1212:U:H3	1.58	0.69
1:XA:1059:C:C5	3:XC:2:GLY:CA	2.75	0.69
1:XA:1318:A:C8	14:XN:16:PHE:CE1	2.79	0.69
34:YA:2080:G:N2	34:YA:2241:A:C4	2.61	0.69
1:QA:107:G:H3'	1:QA:108:G:H21	1.57	0.69
1:QA:186(B):C:O2'	20:QT:89:ARG:HG3	1.92	0.69

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1318:A:H61	14:XN:16:PHE:HB3	1.57	0.69
1:QA:664:G:H22	1:QA:741:G:H1	1.38	0.69
1:QA:1152:A:H1'	10:QJ:17:ASP:OD2	1.92	0.69
27:R3:42:ALA:HB1	34:RA:851:U:O2	1.92	0.69
1:XA:8:A:N1	4:XD:209:ARG:NH1	2.40	0.69
3:XC:22:TRP:CA	10:XJ:93:GLY:HA2	2.23	0.69
4:XD:12:CYS:SG	4:XD:19:LEU:HB2	2.33	0.69
1:QA:936:C:O2'	1:QA:1382:C:N4	2.25	0.69
1:QA:1106:G:O2'	3:QC:172:ARG:NE	2.26	0.69
32:R8:8:LYS:NZ	34:RA:243:U:OP1	2.26	0.69
40:RH:103:LEU:CD1	40:RH:123:PHE:CE1	2.75	0.69
34:YA:1566:A:C2	36:YD:214:TRP:CD1	2.78	0.69
1:QA:413:G:H21	1:QA:428:G:H1'	1.57	0.69
1:QA:1014:A:OP1	19:QS:32:LYS:HE2	1.92	0.69
1:QA:1229:A:N1	13:QM:104:ARG:CZ	2.55	0.69
12:QL:104:VAL:O	12:QL:105:TYR:HD2	1.68	0.69
24:R0:74:ARG:NH2	34:RA:2334:G:O6	2.26	0.69
1:QA:838(B):U:H4'	1:QA:838(C):C:C5	2.28	0.69
1:QA:956:U:O2'	19:QS:80:TYR:CD1	2.46	0.69
1:QA:1160:G:C4'	2:QB:132:LYS:HE2	2.22	0.69
1:QA:1313:U:C6	19:QS:6:LYS:NZ	2.60	0.69
1:QA:1329:A:N6	21:QU:7:ARG:NH2	2.41	0.69
4:QD:61:LYS:CE	4:QD:206:PHE:CE2	2.75	0.69
10:QJ:50:ILE:HG13	14:QN:41:ARG:CD	1.97	0.69
13:QM:86:CYS:HB2	19:QS:69:HIS:ND1	2.07	0.69
1:XA:51:A:C6	1:XA:353:A:C2	2.81	0.69
1:XA:1152:A:C5'	10:XJ:13:HIS:CD2	2.76	0.69
34:YA:747:U:O4	34:YA:2613:U:C2	2.46	0.69
1:QA:186(K):G:N7	17:QQ:63:ARG:NH1	2.41	0.69
1:QA:972:C:O5'	10:QJ:57:LYS:HB2	1.92	0.69
1:QA:1229:A:P	13:QM:108:ARG:HH22	2.10	0.69
34:RA:2453:A:H2'	34:RA:2454:G:H8	1.57	0.69
34:YA:2098:U:H1'	34:YA:2192:G:N2	2.08	0.69
36:YD:35:LYS:HB2	36:YD:63:ARG:HA	1.76	0.69
1:QA:1236:A:C5'	21:QU:10:ARG:HH11	2.03	0.68
4:QD:196:LEU:HA	6:XF:16:GLN:HB3	1.73	0.68
10:QJ:47:PHE:CE2	14:QN:37:PHE:CD2	2.82	0.68
36:RD:8:PRO:HB3	36:RD:14:ARG:HB3	1.75	0.68
1:XA:1228:C:C6	13:XM:104:ARG:CA	2.71	0.68
34:RA:859:G:H21	34:RA:2268:A:H2	1.41	0.68
3:XC:43:LEU:O	3:XC:47:LEU:HB2	1.94	0.68

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:XF:99:ALA:O	18:XR:28:GLU:HG2	1.93	0.68
34:YA:1456:G:N2	34:YA:2704:C:N3	2.41	0.68
3:QC:23:TYR:CE1	10:QJ:10:GLY:HA2	2.28	0.68
1:XA:815:A:H1'	1:XA:1527:C:H1'	1.74	0.68
1:XA:1302:U:H4'	13:XM:27:LYS:HB2	1.75	0.68
34:YA:506:G:H5''	34:YA:509:C:H1'	1.76	0.68
53:RY:99:CYS:HB2	53:RY:103:GLY:H	1.58	0.68
1:XA:664:G:H22	1:XA:741:G:H1	1.38	0.68
1:XA:1318:A:H61	14:XN:16:PHE:CB	2.00	0.68
34:YA:2046:G:N2	34:YA:2623:G:N3	2.41	0.68
42:RN:16:ILE:HB	42:RN:54:VAL:HG12	1.75	0.68
1:QA:186(B):C:N1	20:QT:85:MET:HE2	2.09	0.68
1:QA:754:C:OP1	15:QO:72:ARG:NH2	2.23	0.68
1:QA:959:A:N6	19:QS:79:THR:HG23	2.07	0.68
10:QJ:47:PHE:CE1	14:QN:37:PHE:HE2	2.12	0.68
1:XA:421:U:C4'	3:XC:192:THR:CG2	2.72	0.68
13:XM:84:ILE:O	19:XS:74:PHE:HE1	1.77	0.68
34:YA:1252:G:H21	49:YU:33:ARG:HH11	1.40	0.68
1:QA:950:U:H3	1:QA:1231:G:H1	1.40	0.68
1:QA:956:U:O4'	19:QS:83:HIS:HA	1.92	0.68
4:QD:57:ARG:HH21	5:QE:107:ARG:HH11	1.40	0.68
6:XF:94:GLN:OE1	18:XR:32:ARG:HD2	1.90	0.68
1:QA:1359:C:C5	14:QN:35:ARG:NE	2.61	0.68
45:RQ:38:GLU:HG2	45:RQ:127:ILE:HG23	1.74	0.68
1:QA:975:A:H3'	14:QN:32:SER:HA	1.76	0.68
1:QA:976:G:P	14:QN:32:SER:H	2.17	0.68
1:QA:1359:C:P	14:QN:22:THR:CG2	2.82	0.68
10:QJ:47:PHE:CZ	14:QN:36:PHE:HB2	2.24	0.68
34:YA:1204:A:H2	34:YA:1241:A:H61	1.42	0.68
1:QA:958:A:H1'	19:QS:55:LYS:CD	2.10	0.68
7:QG:16:LEU:HD23	9:QI:42:ARG:HG2	1.66	0.68
34:YA:2097:C:O2	34:YA:2192:G:N1	2.24	0.68
34:YA:2229:C:H2'	34:YA:2230:G:H8	1.58	0.68
1:QA:947:G:OP1	13:QM:108:ARG:HB2	1.94	0.67
1:QA:1359:C:C5	14:QN:35:ARG:CZ	2.77	0.67
38:RF:154:VAL:HG12	38:RF:191:ARG:HB2	1.77	0.67
1:XA:62:U:H1'	1:XA:379:C:H1'	1.76	0.67
1:XA:254:G:H5'	17:XQ:66:SER:OG	1.94	0.67
1:XA:539:A:P	12:XL:114:LYS:HD2	2.34	0.67
1:XA:1316:G:O4'	14:XN:17:LYS:NZ	2.20	0.67
1:XA:1358:U:O3'	14:XN:22:THR:OG1	2.12	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:860:U:H2'	34:YA:861:A:H8	1.58	0.67
1:XA:1128:C:O2'	1:XA:1146:A:N6	2.27	0.67
1:QA:1373:G:P	9:QI:11:LYS:HZ1	2.18	0.67
32:R8:31:HIS:CE1	34:RA:2421:G:N7	2.63	0.67
1:XA:754:C:H6	15:XO:69:TYR:CZ	2.12	0.67
1:XA:979:C:N4	14:XN:18:VAL:HG23	2.08	0.67
34:YA:270(P):U:O4	41:YI:52:ARG:CZ	2.42	0.67
34:YA:1566:A:N3	36:YD:214:TRP:CD2	2.60	0.67
1:QA:675:A:O2'	11:QK:116:HIS:N	2.27	0.67
1:QA:1253:G:OP2	10:QJ:44:VAL:HG23	1.94	0.67
10:QJ:47:PHE:CE2	14:QN:34:TYR:HB3	2.29	0.67
12:QL:7:ILE:HD13	17:QQ:32:TYR:HD1	1.59	0.67
1:XA:107:G:H3'	1:XA:108:G:H21	1.59	0.67
1:XA:1318:A:C8	14:XN:16:PHE:HE1	2.11	0.67
31:Y7:37:LYS:HE2	34:YA:458:G:C4	2.28	0.67
1:QA:1203:C:OP2	14:QN:3:ARG:CD	2.42	0.67
1:QA:1253:G:OP1	10:QJ:44:VAL:CG1	2.42	0.67
4:QD:3:ARG:HH22	4:QD:100:ARG:HH22	1.42	0.67
34:RA:1614:A:N6	51:RW:91:GLY:HA2	2.09	0.67
34:RA:1700:A:H3'	34:RA:1701:A:H8	1.60	0.67
6:XF:89:MET:CE	18:XR:76:LEU:HD13	2.25	0.67
10:XJ:50:ILE:HD11	14:XN:41:ARG:CZ	2.24	0.67
21:XU:3:LYS:HG2	21:XU:14:TRP:HB2	1.76	0.67
42:YN:131:GLN:OE1	42:YN:134:ARG:NH2	2.27	0.67
34:RA:1649:G:O2'	46:RR:107:ASP:OD1	2.04	0.67
34:RA:2429:G:N7	44:RP:56:SER:OG	2.27	0.67
1:XA:974:A:H5'	14:XN:31:ARG:HB3	1.73	0.67
1:XA:1059:C:C5	3:XC:2:GLY:HA3	2.30	0.67
1:XA:1255:G:H1	1:XA:1282:C:H42	1.43	0.67
1:QA:186(K):G:C8	17:QQ:63:ARG:NH2	2.57	0.67
1:QA:1314:C:H5	19:QS:6:LYS:HE3	1.58	0.67
2:QB:178:ARG:O	8:QH:71:GLY:HA2	1.95	0.67
3:QC:23:TYR:CA	10:QJ:93:GLY:O	2.43	0.67
5:QE:139:LEU:HA	5:QE:142:LEU:HD12	1.77	0.67
10:QJ:47:PHE:CZ	14:QN:37:PHE:CD2	2.82	0.67
41:RI:92:VAL:HB	41:RI:120:ILE:HG21	1.74	0.67
54:RZ:52:SER:O	54:RZ:54:HIS:N	2.28	0.67
1:XA:373:A:O2'	1:XA:451:A:N6	2.26	0.67
1:XA:936:C:H2'	1:XA:937:A:H8	1.59	0.67
1:XA:1319:A:OP2	19:XS:3:ARG:CZ	2.42	0.67
6:XF:99:ALA:H	18:XR:29:PHE:H	1.41	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
45:YQ:16:ARG:HH21	45:YQ:18:LYS:HD3	1.59	0.67
1:QA:1014:A:C8	19:QS:34:TRP:NE1	2.62	0.67
1:QA:1376:U:H2'	1:QA:1377:A:H8	1.59	0.67
34:YA:1493:C:C4	34:YA:2210:G:N9	2.63	0.67
34:YA:2107:C:O5'	34:YA:2107:C:H6	1.78	0.67
34:YA:2315:G:OP1	39:YG:36:LYS:NZ	2.25	0.67
1:QA:974:A:OP1	14:QN:29:ARG:HD2	1.95	0.67
1:QA:1230:C:H42	13:QM:104:ARG:HD3	1.55	0.67
1:QA:1541:U:OP2	2:QB:23:ARG:NH2	2.28	0.67
40:RH:101:ARG:HH12	40:RH:123:PHE:H	1.43	0.67
1:XA:1360:A:H1'	14:YN:17:LYS:CG	2.25	0.67
34:YA:1248:G:C2	49:YU:3:ARG:HD2	2.30	0.67
1:XA:421:U:O4'	3:XC:192:THR:CG2	2.30	0.67
1:XA:974:A:OP1	14:YN:31:ARG:HD2	1.95	0.67
1:XA:1160:G:O4'	2:XB:132:LYS:CE	2.41	0.67
7:XG:16:LEU:HD22	9:XI:45:ALA:HB2	1.75	0.67
1:QA:1106:G:C4'	3:QC:172:ARG:NE	2.57	0.66
1:QA:1158:C:C4'	2:QB:133:LYS:HZ1	1.98	0.66
1:QA:1373:G:P	9:QI:11:LYS:NZ	2.68	0.66
4:QD:31:CYS:SG	4:QD:33:MET:HB2	2.35	0.66
14:QN:24:CYS:SG	14:QN:40:CYS:CA	2.81	0.66
1:XA:1320:C:H1'	19:XS:70:LYS:HD3	1.60	0.66
6:XF:94:GLN:CD	18:XR:32:ARG:HH11	1.99	0.66
28:Y4:7:PRO:HG3	39:YG:62:LEU:HA	1.77	0.66
34:YA:94:G:O6	55:YA:3185:MG:MG	1.37	0.66
53:YY:79:CYS:SG	57:YY:201:ZN:ZN	1.84	0.66
1:QA:1160:G:C4'	2:QB:132:LYS:CE	2.72	0.66
1:QA:1484:C:HO2'	34:RA:1960:A:HO2'	1.42	0.66
25:R1:78:LYS:HZ3	34:RA:270(T):G:H1'	1.60	0.66
32:R8:39:LYS:NZ	34:RA:2365:G:O6	2.20	0.66
10:XJ:50:ILE:HD11	14:YN:41:ARG:NH1	2.10	0.66
34:YA:410:G:N2	34:YA:2407:G:C5	2.63	0.66
1:QA:625:G:O2'	16:QP:16:HIS:ND1	2.27	0.66
1:QA:981:U:H5''	14:QN:6:LEU:HD21	1.77	0.66
1:QA:1373:G:O5'	9:QI:11:LYS:NZ	2.28	0.66
34:RA:955:C:OP2	45:RQ:14:ARG:NH1	2.27	0.66
54:RZ:112:ARG:HG3	54:RZ:114:GLY:H	1.59	0.66
1:XA:728:A:N6	15:XO:54:ARG:HD2	2.10	0.66
1:QA:59:A:H5''	1:QA:387:U:H5''	1.78	0.66
1:QA:959:A:N1	19:QS:77:THR:O	2.29	0.66
34:RA:515:A:H1'	34:RA:581:C:H1'	1.76	0.66

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:186(B):C:C2	20:XT:105:SER:HB2	2.30	0.66
1:XA:664:G:N2	1:XA:726:C:O2'	2.25	0.66
24:Y0:33:ALA:O	34:YA:2353:G:O2'	2.12	0.66
1:QA:1360:A:C4'	14:QN:17:LYS:NZ	2.56	0.66
34:RA:379:G:H1	34:RA:395:U:H3	1.44	0.66
34:RA:2440:C:H5''	34:RA:2587:A:H4'	1.77	0.66
35:RB:116:G:H4'	47:RS:54:LEU:HD22	1.76	0.66
1:XA:1059:C:H5	3:XC:2:GLY:CA	2.08	0.66
10:XJ:37:PRO:HG2	10:XJ:37:PRO:O	1.96	0.66
34:YA:1044:G:O2'	34:YA:1047:G:O2'	2.13	0.66
1:XA:1253:G:C6	1:XA:1254:C:N4	2.64	0.66
3:XC:60:ALA:HB1	10:XJ:91:PRO:HG2	1.78	0.66
1:QA:310:G:P	16:QP:27:LYS:CD	2.67	0.66
1:QA:972:C:OP2	10:QJ:57:LYS:CD	2.44	0.66
1:QA:980:C:C6	14:QN:19:ARG:CG	2.68	0.66
1:XA:702:A:C6	34:YA:1848:A:C2	2.83	0.66
1:XA:749:C:H2'	1:XA:750:G:H8	1.59	0.66
1:XA:1301:U:O3'	13:XM:17:VAL:HG23	1.96	0.66
2:XB:185:ILE:HG22	2:XB:199:TYR:HB2	1.78	0.66
9:XI:18:PHE:HB3	9:XI:20:ARG:HH12	1.60	0.66
10:XJ:16:LEU:HB3	10:XJ:70:ARG:HH12	1.60	0.66
32:Y8:42:ARG:NH1	34:YA:2349:G:OP2	2.28	0.66
34:YA:886:C:O2'	34:YA:889:C:N4	2.26	0.66
1:QA:1010:G:H2'	1:QA:1011:G:H8	1.59	0.66
1:XA:1059:C:C5	3:XC:2:GLY:HA2	2.31	0.66
3:XC:23:TYR:CE2	10:XJ:95:GLU:CB	2.77	0.66
34:YA:2056:G:O6	34:YA:2577:A:C8	2.49	0.66
34:YA:2343:C:O2'	34:YA:2373:G:O2'	2.14	0.66
1:QA:1240:U:O2	7:QG:38:LEU:HA	1.95	0.66
34:RA:248:G:C4	34:RA:2431:U:H4'	2.31	0.66
6:XF:99:ALA:C	18:XR:28:GLU:HA	2.10	0.66
13:QM:14:ARG:HG2	13:QM:44:ARG:HD3	1.78	0.66
40:RH:18:GLU:HB2	40:RH:25:LYS:HB2	1.78	0.66
1:XA:51:A:N3	1:XA:353:A:C6	2.64	0.66
1:XA:974:A:C8	14:XN:31:ARG:HD2	2.30	0.66
6:XF:100:ASN:CB	18:XR:27:GLY:C	2.64	0.66
34:YA:199:A:C6	34:YA:2434:A:C2	2.83	0.66
34:YA:2051:A:C2	34:YA:2614:A:C4	2.84	0.66
40:YH:84:SER:HB2	40:YH:132:ARG:HD2	1.78	0.66
1:QA:1060:C:H4'	10:QJ:52:GLY:HA2	1.77	0.65
1:QA:1099:G:OP2	2:QB:96:ARG:HD3	1.93	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1317:C:OP1	14:QN:16:PHE:CE2	2.47	0.65
1:QA:1541:U:OP2	2:QB:23:ARG:CZ	2.44	0.65
2:QB:167:PRO:O	2:QB:171:ALA:HB2	1.96	0.65
2:QB:181:PHE:CD2	8:QH:70:GLN:HG2	2.31	0.65
34:RA:1333:C:H2'	34:RA:1334:G:H8	1.61	0.65
51:RW:88:ARG:HB2	51:RW:92:ARG:HB3	1.77	0.65
1:XA:6:G:H4'	1:XA:298:A:H4'	1.78	0.65
1:XA:186(B):C:O2	20:XT:105:SER:N	2.29	0.65
1:XA:1227:A:OP1	13:XM:96:LEU:CD2	2.40	0.65
3:XC:60:ALA:CB	10:XJ:91:PRO:HD2	2.26	0.65
1:QA:1307:U:C5'	13:QM:99:ARG:HG2	2.26	0.65
41:RI:83:ALA:O	41:RI:89:TYR:CE2	2.48	0.65
1:XA:658:G:H1'	15:XO:22:THR:CG2	2.26	0.65
34:YA:184:C:H1'	34:YA:217:G:H1'	1.78	0.65
1:QA:192:U:H2'	1:QA:193:C:H6	1.62	0.65
1:QA:740:U:O3'	15:QO:39:LEU:HD12	1.96	0.65
1:QA:1377:A:O2'	1:QA:1379:G:O6	2.12	0.65
4:QD:57:ARG:NH1	5:QE:107:ARG:HD3	2.11	0.65
7:QG:16:LEU:CD2	9:QI:42:ARG:HA	2.26	0.65
13:XM:84:ILE:HG21	19:XS:66:MET:HB3	1.77	0.65
14:XN:3:ARG:HB3	14:XN:3:ARG:CZ	2.26	0.65
34:YA:1216:G:OP1	49:YU:11:ARG:NH2	2.26	0.65
34:YA:2150:U:H2'	34:YA:2151:G:C8	2.31	0.65
3:XC:23:TYR:HB2	10:XJ:93:GLY:O	1.97	0.65
35:YB:90:C:OP2	45:YQ:16:ARG:NH1	2.30	0.65
1:QA:607:A:N1	16:QP:31:LYS:CA	2.54	0.65
29:R5:3:LYS:O	34:RA:2056:G:N2	2.19	0.65
38:RF:147:GLY:O	38:RF:191:ARG:NH1	2.29	0.65
1:XA:1305:G:H21	1:XA:1331:G:H3'	1.59	0.65
1:XA:1329:A:N7	21:XU:7:ARG:NH2	2.45	0.65
1:XA:1375:A:H3'	1:XA:1376:U:H6	1.60	0.65
1:XA:1422:G:H5'	43:YO:48:PRO:HG3	1.78	0.65
10:XJ:62:HIS:CD2	14:XN:59:ALA:HB1	2.31	0.65
34:YA:1088:A:H4'	34:YA:1089:G:H8	1.61	0.65
34:RA:445:C:OP1	49:RU:2:PRO:HA	1.96	0.65
34:RA:1044:G:O2'	34:RA:1047:G:O2'	2.15	0.65
34:RA:2483:C:N3	45:RQ:124:LYS:NZ	2.44	0.65
40:RH:89:ILE:HD11	40:RH:131:VAL:HG22	1.76	0.65
1:XA:1202:G:O2'	14:XN:28:GLY:O	2.14	0.65
1:XA:1278:U:H4'	1:XA:1279:A:C5	2.32	0.65
8:XH:14:ARG:HB3	8:XH:83:ILE:HD11	1.78	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:XT:86:ARG:O	20:XT:90:GLN:NE2	2.30	0.65
34:YA:1457:A:C2	34:YA:2703:C:C4	2.85	0.65
1:QA:634:C:H2'	1:QA:635:G:C8	2.32	0.65
1:QA:1314:C:OP2	19:QS:6:LYS:NZ	2.24	0.65
1:QA:1318:A:N6	14:QN:16:PHE:HB3	2.11	0.65
34:RA:1152:C:H2'	34:RA:1153:C:H6	1.61	0.65
50:RV:62:LEU:HD11	50:RV:95:LEU:HB2	1.77	0.65
1:XA:636:U:H3'	1:XA:637:G:H8	1.62	0.65
1:XA:1238:A:H2'	1:XA:1239:A:C8	2.32	0.65
1:XA:1463:C:H4'	48:YT:112:ARG:HH21	1.62	0.65
29:Y5:52:TYR:OH	34:YA:2883:A:OP1	2.06	0.65
34:YA:2080:G:C2	34:YA:2241:A:N3	2.64	0.65
1:QA:1014:A:C8	19:QS:34:TRP:CE2	2.76	0.65
1:QA:1220:G:H21	19:QS:54:GLY:HA3	1.61	0.65
1:XA:376:G:H4'	16:XP:5:ARG:HH11	1.61	0.65
1:XA:1150:U:H2'	1:XA:1151:A:C8	2.32	0.65
1:XA:1377:A:C6	7:XG:10:ARG:HD2	2.32	0.65
34:YA:2238:G:OP1	34:YA:2238:G:N2	2.30	0.65
47:YS:106:ARG:HB2	47:YS:110:LEU:HD23	1.78	0.65
1:QA:1307:U:H3'	13:QM:99:ARG:NH2	2.11	0.65
3:QC:79:ARG:HH21	11:XK:105:VAL:H	1.44	0.65
54:RZ:10:ARG:HD2	54:RZ:38:TYR:HB3	1.79	0.65
1:XA:1253:G:C2	1:XA:1254:C:C4	2.85	0.65
1:XA:1253:G:N1	1:XA:1254:C:C4	2.65	0.65
6:XF:99:ALA:CB	18:XR:29:PHE:CD1	2.75	0.65
10:XJ:62:HIS:HD2	14:XN:59:ALA:HB1	1.62	0.65
11:XK:91:ARG:CG	18:XR:88:LYS:HZ1	2.09	0.65
34:YA:2049:G:N2	34:YA:2620:C:N3	2.44	0.65
1:QA:876:G:C1'	8:QH:11:THR:HG21	2.24	0.65
1:QA:1221:G:H5'	19:QS:36:ARG:NH2	2.08	0.65
1:QA:1299:A:H2'	1:QA:1301:U:H1'	1.79	0.65
1:QA:1541:U:OP1	2:QB:23:ARG:NH2	2.29	0.65
4:QD:121:VAL:HG22	4:QD:126:ILE:HG13	1.79	0.65
37:RE:75:VAL:HG23	37:RE:76:ARG:HG2	1.79	0.65
1:XA:1123:A:H4'	10:XJ:37:PRO:HD2	1.79	0.65
34:YA:1566:A:C2	36:YD:214:TRP:NE1	2.65	0.65
1:QA:981:U:H5''	1:QA:982:U:H2'	1.77	0.64
1:QA:1203:C:OP2	14:QN:3:ARG:HD2	1.96	0.64
2:QB:177:ALA:HB1	2:QB:182:ILE:HB	1.78	0.64
7:QG:16:LEU:HD22	9:QI:42:ARG:HA	1.78	0.64
1:XA:412:A:N3	4:XD:35:ARG:HD3	2.11	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1124:G:C4'	10:XJ:36:GLY:H	2.05	0.64
4:XD:20:TYR:HA	4:XD:26:CYS:SG	2.37	0.64
11:XK:108:ILE:H	18:XR:87:ARG:CD	2.10	0.64
34:YA:805:G:H22	34:YA:828:U:H5''	1.61	0.64
1:QA:980:C:OP1	14:QN:19:ARG:NH2	2.30	0.64
31:R7:5:TRP:CZ3	34:RA:686:G:N7	2.65	0.64
29:Y5:16:ARG:NH1	29:Y5:17:ASP:OD1	2.30	0.64
25:Y1:87:PRO:HA	25:Y1:90:ILE:HG22	1.80	0.64
34:YA:1789:A:OP2	36:YD:222:ARG:NH1	2.30	0.64
1:QA:973:G:O3'	14:QN:29:ARG:NE	2.23	0.64
1:XA:107:G:O6	20:XT:15:ARG:CZ	2.43	0.64
1:XA:1359:C:H5	14:XN:35:ARG:HD3	1.62	0.64
1:XA:1541:U:C2	23:XX:5:A:C2	2.85	0.64
34:YA:2048:G:N2	34:YA:2621:A:C4	2.65	0.64
1:QA:495:A:H1'	1:QA:497:A:H2'	1.78	0.64
2:QB:132:LYS:HA	2:QB:135:GLN:HB2	1.78	0.64
34:RA:414:C:H1'	34:RA:1864:U:H1'	1.78	0.64
35:RB:55:U:O2'	39:RG:27:ASN:ND2	2.31	0.64
1:XA:585:G:OP1	17:XQ:37:LYS:HE2	1.98	0.64
1:XA:1286:A:N6	1:XA:1355:G:OP1	2.30	0.64
1:XA:1319:A:OP2	19:XS:3:ARG:CD	2.46	0.64
34:YA:788:A:OP1	34:YA:791:C:N4	2.28	0.64
34:YA:2402:C:H1'	34:YA:2403:C:H5	1.63	0.64
1:QA:237:C:H2'	1:QA:238:G:H8	1.62	0.64
1:QA:979:C:O2	14:QN:19:ARG:NH1	2.30	0.64
1:QA:1378:C:P	7:QG:7:ALA:CB	2.75	0.64
2:QB:178:ARG:O	8:QH:71:GLY:CA	2.45	0.64
7:QG:16:LEU:CD2	9:QI:42:ARG:CB	2.75	0.64
41:RI:92:VAL:HB	41:RI:120:ILE:HB	1.80	0.64
1:XA:406:G:H2'	1:XA:407:G:H8	1.63	0.64
1:XA:1059:C:OP2	3:XC:3:ASN:N	2.29	0.64
1:XA:1239:A:N1	1:XA:1297:C:H1'	2.12	0.64
1:XA:1359:C:C5	14:XN:35:ARG:HD3	2.33	0.64
9:XI:17:VAL:HG12	9:XI:63:ILE:HD12	1.80	0.64
34:YA:1448:G:O2'	34:YA:1528:A:N6	2.31	0.64
1:QA:1249:C:N4	1:QA:1288:A:OP2	2.30	0.64
1:QA:1368:G:O3'	10:QJ:46:ARG:NH2	2.30	0.64
6:XF:91:VAL:HG21	18:XR:34:TYR:OH	1.97	0.64
10:XJ:50:ILE:HA	10:XJ:60:ARG:HB2	1.78	0.64
29:Y5:3:LYS:HB2	34:YA:2577:A:O4'	1.97	0.64
1:QA:1203:C:P	14:QN:3:ARG:HD2	2.37	0.64

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:657:G:H21	15:XO:22:THR:HG23	1.63	0.64
7:XG:16:LEU:CD2	9:XI:44:VAL:CG2	2.65	0.64
34:YA:2291:U:H1'	34:YA:2374:C:H1'	1.80	0.64
1:QA:1048:G:OP1	14:QN:4:LYS:HE3	1.98	0.64
1:QA:1221:G:O2'	19:QS:77:THR:CB	2.44	0.64
1:QA:1229:A:N6	13:QM:104:ARG:CD	2.61	0.64
4:QD:20:TYR:CG	6:XF:15:ASP:HB3	2.33	0.64
10:XJ:62:HIS:CE1	14:XN:61:TRP:HE3	2.16	0.64
22:XV:8:U:O4	22:XV:14:A:N7	2.31	0.64
22:XV:65:U:H2'	22:XV:66:A:H8	1.62	0.64
1:QA:666:G:H5'	1:QA:726:C:H1'	1.80	0.64
1:QA:1236:A:P	21:QU:10:ARG:HH11	2.20	0.64
1:QA:1320:C:N4	19:QS:37:ARG:CB	2.61	0.64
1:QA:1320:C:C5'	19:QS:70:LYS:HG3	2.27	0.64
1:XA:824:C:H2'	1:XA:825:G:C8	2.33	0.64
34:YA:1226:G:OP1	50:YV:69:LYS:NZ	2.26	0.64
34:YA:2377:A:O2'	47:YS:111:GLU:O	2.14	0.64
1:QA:1125:U:C4	10:QJ:73:ASP:OD1	2.51	0.63
4:QD:18:LYS:HG3	4:QD:33:MET:HG3	1.78	0.63
13:QM:86:CYS:HB2	19:QS:69:HIS:HE1	1.61	0.63
34:RA:2789:C:H1'	34:RA:2892:A:C2	2.33	0.63
36:RD:60:ARG:HD3	36:RD:86:PRO:HB2	1.80	0.63
40:RH:6:ARG:HH22	40:RH:54:ARG:HD3	1.62	0.63
1:XA:1320:C:C5	19:XS:37:ARG:HB2	2.32	0.63
6:XF:50:TYR:CD1	18:XR:77:GLY:O	2.48	0.63
28:Y4:22:ILE:HG22	28:Y4:23:GLU:HG3	1.80	0.63
34:YA:1493:C:C5	34:YA:2210:G:C8	2.85	0.63
1:QA:675:A:H1'	11:QK:116:HIS:HB2	1.80	0.63
1:QA:986:A:H1'	19:QS:54:GLY:C	2.17	0.63
1:QA:1141:C:H2'	1:QA:1142:G:H8	1.63	0.63
1:QA:1229:A:C6	13:QM:104:ARG:CZ	2.79	0.63
1:QA:1251:A:O2'	1:QA:1369:C:O3'	2.16	0.63
1:QA:1342:C:H2'	1:QA:1343:G:H8	1.63	0.63
33:R9:23:VAL:HG21	34:RA:1032:A:C1'	2.24	0.63
34:RA:2250:G:C6	45:RQ:82:ARG:HD2	2.32	0.63
40:RH:78:GLY:HA2	40:RH:82:GLY:HA3	1.79	0.63
41:RI:84:GLY:HA3	41:RI:89:TYR:CZ	2.33	0.63
45:RQ:81:VAL:HG12	45:RQ:82:ARG:HG2	1.80	0.63
1:XA:718:G:O4'	11:XK:117:ASN:HB2	1.98	0.63
2:XB:54:THR:HG22	2:XB:199:TYR:HB3	1.81	0.63
3:XC:60:ALA:HB1	10:XJ:91:PRO:CG	2.29	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
11:XK:109:VAL:CG2	18:XR:86:VAL:HG23	2.27	0.63
32:Y8:12:LYS:NZ	34:YA:249:C:O2	2.31	0.63
34:YA:1447:G:H1'	34:YA:1545(B):A:H1'	1.81	0.63
1:QA:186(B):C:C1'	20:QT:85:MET:HE2	2.29	0.63
1:QA:564:C:H5'	17:QQ:32:TYR:CE1	2.33	0.63
1:QA:975:A:H3'	14:QN:32:SER:CA	2.28	0.63
32:R8:35:GLN:HE22	32:R8:36:LYS:HE2	1.62	0.63
54:RZ:52:SER:O	54:RZ:54:HIS:ND1	2.32	0.63
1:XA:1228:C:P	13:XM:108:ARG:NH2	2.72	0.63
1:XA:1532:U:H2'	1:XA:1534:A:H2	1.64	0.63
6:XF:91:VAL:HG11	18:XR:34:TYR:HE1	1.63	0.63
34:YA:2250:G:O2'	34:YA:2496:C:OP1	2.16	0.63
1:QA:1005:A:H4'	1:QA:1037:C:H1'	1.79	0.63
5:QE:140:ARG:O	5:QE:143:ARG:NH1	2.32	0.63
34:RA:23:G:N2	51:RW:77:ASP:OD1	2.30	0.63
34:RA:1566:A:C2	36:RD:214:TRP:CE2	2.86	0.63
1:XA:8:A:N1	4:XD:209:ARG:CZ	2.61	0.63
1:XA:662:G:H2'	1:XA:663:A:C8	2.33	0.63
1:XA:728:A:C6	15:XO:54:ARG:HD2	2.34	0.63
1:XA:1004:A:N1	1:XA:1025:U:H4'	2.12	0.63
1:XA:1318:A:H62	14:XN:16:PHE:HB3	1.62	0.63
32:Y8:22:VAL:HB	32:Y8:53:PRO:HB3	1.80	0.63
34:YA:784:A:N6	34:YA:2072:G:O2'	2.32	0.63
34:YA:1853:A:N3	34:YA:2233:U:O2'	2.29	0.63
1:QA:109:A:H62	1:QA:324:G:H21	1.45	0.63
1:QA:741:G:O5'	15:QO:39:LEU:CD1	2.46	0.63
1:QA:766:A:H61	1:QA:1511:G:H1'	1.62	0.63
1:QA:948:C:OP1	13:QM:106:ASN:O	2.15	0.63
1:QA:974:A:OP1	14:QN:29:ARG:NE	2.30	0.63
21:QU:12:LYS:HB3	21:QU:22:ARG:HD2	1.81	0.63
1:XA:1005:A:H4'	1:XA:1037:C:H1'	1.81	0.63
17:XQ:29:HIS:HB3	17:XQ:33:GLY:H	1.64	0.63
44:YP:4:SER:O	44:YP:7:ARG:NH2	2.32	0.63
34:RA:508:G:O6	51:RW:9:TYR:CD1	2.52	0.63
34:RA:581:C:H2'	34:RA:582:G:C8	2.34	0.63
1:XA:865:A:N3	1:XA:918:A:O2'	2.30	0.63
1:XA:1071:C:OP1	5:XE:27:ARG:NH2	2.31	0.63
1:XA:1106:G:H4'	3:XC:172:ARG:HG2	1.80	0.63
1:XA:1240:U:H1'	7:XG:38:LEU:CG	2.22	0.63
1:QA:54:C:H42	1:QA:357:G:H1	1.46	0.63
1:QA:948:C:OP2	13:QM:106:ASN:HB2	1.98	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1313:U:H3'	19:QS:6:LYS:HZ2	0.80	0.63
3:QC:79:ARG:NH2	11:XK:105:VAL:N	2.46	0.63
28:R4:26:SER:HG	39:RG:143:GLU:CD	1.97	0.63
33:R9:18:ARG:CZ	34:RA:1034:G:O4'	2.47	0.63
41:RI:83:ALA:HB1	41:RI:88:ILE:HA	1.81	0.63
1:XA:1329:A:H4'	13:XM:29:ARG:NH2	2.13	0.63
34:YA:392:C:H5''	34:YA:409:C:H5''	1.80	0.63
1:QA:1186:G:N2	14:QN:61:TRP:HA	2.13	0.63
1:QA:1318:A:C6	14:QN:16:PHE:CD1	2.87	0.63
3:QC:23:TYR:HB2	10:QJ:94:VAL:CA	2.28	0.63
37:RE:109:LYS:HE3	37:RE:191:PRO:HA	1.79	0.63
11:XK:110:ASP:CA	18:XR:85:LEU:O	2.46	0.63
34:YA:197:A:N9	34:YA:2430:A:H2	1.96	0.63
1:QA:194:C:H4'	20:QT:68:LYS:HE2	1.79	0.63
1:QA:953:G:O6	13:QM:104:ARG:NH1	2.32	0.63
1:QA:1126:U:O2	10:QJ:38:ILE:HD12	1.98	0.63
15:QO:89:GLY:C	34:RA:716:A:OP1	2.36	0.63
34:RA:2250:G:N3	45:RQ:82:ARG:HG3	2.14	0.63
1:XA:373:A:O2'	1:XA:451:A:N7	2.32	0.63
1:XA:1226:C:OP1	13:XM:91:ARG:NH1	2.31	0.63
34:YA:2099:U:N3	34:YA:2190:G:N1	1.97	0.63
34:YA:2118:U:H3	34:YA:2148:G:H4'	1.64	0.63
1:QA:186(B):C:H5'	20:QT:82:SER:HA	1.81	0.62
1:QA:667:G:N3	15:QO:49:ASP:OD1	2.32	0.62
1:QA:677:U:H3	1:QA:713:G:H1	1.47	0.62
1:QA:953:G:C6	13:QM:104:ARG:NH1	2.67	0.62
2:QB:195:ASP:O	8:QH:68:ARG:NH2	2.32	0.62
34:RA:137(B):G:N3	52:RX:41:ASN:ND2	2.47	0.62
1:XA:1198:G:H2'	1:XA:1199:U:C6	2.34	0.62
1:XA:1305:G:C2	1:XA:1331:G:N7	2.67	0.62
4:XD:8:VAL:HA	4:XD:11:LEU:HD13	1.81	0.62
34:YA:2054:A:N6	34:YA:2577:A:H61	1.87	0.62
1:QA:9:G:C5'	5:QE:122:GLU:OE2	2.47	0.62
1:QA:62:U:H1'	1:QA:379:C:H1'	1.80	0.62
1:QA:1229:A:P	13:QM:108:ARG:HH21	2.21	0.62
1:QA:1320:C:H5'	19:QS:70:LYS:HG3	1.80	0.62
10:QJ:53:PRO:HA	14:QN:42:ILE:HD11	1.81	0.62
1:XA:51:A:N3	1:XA:353:A:N6	2.48	0.62
34:YA:67:U:H3	34:YA:74:A:H2	1.46	0.62
1:QA:740:U:O3'	15:QO:39:LEU:CD1	2.46	0.62
34:RA:229:A:H4'	34:RA:230:U:H5'	1.81	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
37:RE:119:ARG:HG3	37:RE:160:TYR:CD1	2.34	0.62
45:RQ:37:LEU:HD11	45:RQ:130:LYS:HG2	1.81	0.62
1:XA:346:G:OP1	48:YT:41:ARG:NH2	2.31	0.62
1:XA:1229:A:N6	13:XM:104:ARG:CD	2.62	0.62
6:XF:99:ALA:N	18:XR:29:PHE:O	2.32	0.62
1:QA:1246:C:N4	21:QU:26:LYS:HB2	2.12	0.62
1:QA:1313:U:C5	19:QS:4:SER:CB	2.81	0.62
1:XA:740:U:OP1	15:XO:2:PRO:HA	2.00	0.62
11:XK:116:HIS:CD2	18:XR:81:PHE:HB3	2.34	0.62
24:Y0:72:ARG:HE	24:Y0:75:LEU:HD12	1.64	0.62
54:YZ:52:SER:O	54:YZ:54:HIS:N	2.32	0.62
1:QA:1188:A:C3'	14:QN:58:LYS:HZ2	2.12	0.62
3:QC:79:ARG:HH21	11:XK:105:VAL:N	1.98	0.62
11:QK:71:LYS:HZ2	34:RA:2146:C:N4	1.94	0.62
29:R5:51:TYR:CE1	29:R5:56:LYS:HB3	2.35	0.62
1:XA:1240:U:H1'	7:XG:38:LEU:HD12	1.67	0.62
25:Y1:45:ASN:HA	34:YA:2230:G:H1'	1.80	0.62
1:QA:359:U:H2'	1:QA:360:A:H8	1.64	0.62
53:RY:99:CYS:CB	53:RY:103:GLY:H	2.12	0.62
1:XA:619:U:C5'	4:XD:131:ARG:HH21	2.12	0.62
1:XA:979:C:N3	14:XN:19:ARG:NE	2.26	0.62
1:XA:1124:G:O5'	10:XJ:36:GLY:N	2.33	0.62
34:YA:2072:G:C2	34:YA:2438:U:O2	2.53	0.62
34:YA:2787:C:H1'	37:YE:62:PRO:HG3	1.80	0.62
34:YA:2848:G:O2'	34:YA:2867:G:N2	2.32	0.62
41:YI:80:PRO:HB2	41:YI:146:ALA:HB2	1.81	0.62
25:R1:95:LEU:O	25:R1:95:LEU:CD2	2.45	0.62
34:RA:475:U:H4'	34:RA:510:C:H5'	1.80	0.62
1:XA:1372:U:OP1	9:XI:68:GLY:CA	2.43	0.62
10:XJ:62:HIS:CE1	14:XN:61:TRP:CE3	2.87	0.62
34:YA:629:G:N3	34:YA:639:U:O2'	2.30	0.62
34:YA:2051:A:C2	34:YA:2614:A:C5	2.88	0.62
1:QA:658:G:H2'	1:QA:659:U:C6	2.35	0.62
1:QA:989:C:H2'	1:QA:990:C:C6	2.35	0.62
1:QA:1114:C:O2	14:QN:60:SER:O	2.18	0.62
1:QA:1364:U:O4'	21:QU:14:TRP:HZ2	1.83	0.62
34:RA:71:A:H5''	34:RA:72:U:H3'	1.81	0.62
1:XA:1228:C:OP1	13:XM:108:ARG:NH1	2.32	0.62
32:Y8:11:LYS:HB3	32:Y8:60:LEU:HD11	1.81	0.62
32:R8:42:ARG:HD2	34:RA:2350:C:O5'	2.00	0.62
34:RA:820:A:H1'	34:RA:943:U:H1'	1.81	0.62

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:855:G:N2	1:XA:1539:C:OP1	2.33	0.62
10:XJ:39:PRO:CA	10:XJ:70:ARG:HG2	2.28	0.62
1:QA:838(B):U:H4'	1:QA:838(C):C:C6	2.34	0.62
34:RA:2630:G:H21	34:RA:2892:A:H1'	1.65	0.62
34:RA:2751:G:N1	40:RH:3:ARG:HB2	2.15	0.62
32:Y8:25:MET:HG2	44:YP:65:ARG:HH12	1.64	0.62
34:YA:2054:A:H2	34:YA:2616:C:O2	1.82	0.62
36:YD:264:LYS:HG2	36:YD:266:SER:H	1.64	0.62
34:RA:749:C:H5'	34:RA:1271:G:H1'	1.81	0.61
47:RS:83:LYS:HG3	47:RS:84:GLN:HG3	1.81	0.61
54:RZ:72:ARG:NH2	54:RZ:97:GLU:O	2.33	0.61
6:XF:99:ALA:CB	18:XR:29:PHE:CE1	2.83	0.61
1:QA:625:G:O2'	16:QP:16:HIS:CG	2.53	0.61
28:R4:16:CYS:SG	28:R4:17:GLY:N	2.73	0.61
33:R9:16:VAL:HG11	34:RA:1033:U:OP1	2.00	0.61
34:RA:1828:G:O6	36:RD:222:ARG:HD3	1.99	0.61
8:XH:17:THR:O	8:XH:78:GLN:NE2	2.32	0.61
34:YA:2053:G:C2	34:YA:2617:C:C2	2.88	0.61
47:YS:23:ARG:NH2	47:YS:84:GLN:OE1	2.33	0.61
1:QA:973:G:H4'	14:QN:29:ARG:HH21	1.65	0.61
1:QA:1221:G:H5'	19:QS:36:ARG:NH1	2.14	0.61
16:QP:53:VAL:HG12	16:QP:79:VAL:HG12	1.82	0.61
27:R3:12:PRO:HB2	27:R3:20:LYS:HD3	1.82	0.61
37:RE:141:ILE:O	37:RE:154:LYS:NZ	2.33	0.61
11:XK:110:ASP:O	18:XR:84:LYS:HB2	2.01	0.61
34:YA:300:A:OP1	53:YY:86:ARG:NH2	2.33	0.61
34:YA:2056:G:N7	34:YA:2577:A:C5	2.67	0.61
38:YF:160:ASN:HB3	38:YF:163:VAL:HG12	1.81	0.61
1:QA:69:G:H1'	1:QA:152:A:H2	1.66	0.61
1:QA:976:G:H21	1:QA:1362(B):C:H2'	1.66	0.61
34:RA:1041:C:H2'	34:RA:1042:G:H8	1.64	0.61
44:RP:47:ASP:OD2	44:RP:50:ARG:NH2	2.33	0.61
1:XA:1187:G:N2	14:YN:60:SER:OG	2.33	0.61
21:XU:3:LYS:HG2	21:XU:14:TRP:CG	2.36	0.61
34:YA:17:G:H21	34:YA:554:U:H5'	1.64	0.61
34:YA:2822:G:O2'	34:YA:2825:C:N4	2.34	0.61
50:YV:72:VAL:HB	50:YV:85:LYS:HB3	1.82	0.61
1:QA:538:G:C5'	12:QL:115:LYS:HG2	2.27	0.61
1:QA:856:C:H2'	1:QA:857:C:H6	1.64	0.61
7:QG:118:VAL:O	7:QG:122:HIS:ND1	2.32	0.61
10:QJ:47:PHE:CE1	14:QN:37:PHE:CE2	2.88	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:2751:G:C6	40:RH:3:ARG:CB	2.83	0.61
1:XA:675:A:O2'	11:XK:115:PRO:HA	2.00	0.61
1:XA:745:C:H2'	1:XA:746:A:H8	1.63	0.61
34:YA:2059:A:H2	34:YA:2503:A:N6	1.95	0.61
1:QA:264:U:O2'	17:QQ:64:PRO:HD2	2.01	0.61
1:QA:1198:G:H1'	10:QJ:54:PHE:CZ	2.35	0.61
34:RA:2490:G:N2	34:RA:2490:G:OP2	2.32	0.61
1:XA:355:C:O5'	1:XA:355:C:H6	1.84	0.61
10:XJ:7:LYS:HE2	10:XJ:9:ARG:HG3	1.83	0.61
16:XP:6:LEU:HB2	16:XP:17:TYR:HB3	1.82	0.61
34:YA:2820:A:H1'	46:YR:3:HIS:ND1	2.15	0.61
1:QA:311:C:OP1	16:QP:26:ARG:NH1	2.33	0.61
1:QA:609:A:H5'	16:QP:9:PHE:HE1	1.64	0.61
10:QJ:49:VAL:HA	14:QN:34:TYR:OH	2.01	0.61
34:RA:2313:C:H5''	39:RG:91:ARG:HH21	1.65	0.61
36:RD:96:HIS:HE1	36:RD:102:LYS:HE2	1.64	0.61
11:QK:71:LYS:HZ3	34:RA:2146:C:N4	1.92	0.61
34:RA:81:G:HO2'	34:RA:295:G:HO2'	1.48	0.61
1:XA:666:G:H5'	1:XA:726:C:H1'	1.82	0.61
1:XA:1302:U:C2	13:XM:27:LYS:HE2	2.36	0.61
25:Y1:43:TYR:HD2	34:YA:2230:G:H5''	1.66	0.61
34:YA:199:A:N1	34:YA:2434:A:C2	2.68	0.61
34:YA:335:C:OP2	53:YY:84:ARG:NH2	2.33	0.61
34:YA:383:U:H2'	34:YA:385:C:H5	1.65	0.61
34:YA:1333:C:H2'	34:YA:1334:G:H8	1.65	0.61
42:YN:112:LEU:O	42:YN:116:LEU:HB2	1.99	0.61
1:QA:953:G:C4	13:QM:104:ARG:NH2	2.69	0.61
1:XA:413:G:H1'	1:XA:428:G:H21	1.66	0.61
1:XA:702:A:H3'	1:XA:703:G:H8	1.66	0.61
1:XA:1124:G:C5'	10:XJ:36:GLY:H	2.14	0.61
1:XA:1330:U:H5''	13:XM:24:GLY:HA2	1.81	0.61
1:XA:1358:U:H5	14:XN:35:ARG:HE	1.48	0.61
34:YA:554:U:H2'	34:YA:556:G:C8	2.36	0.61
34:YA:2046:G:C2	34:YA:2623:G:C4	2.89	0.61
34:YA:2471:C:N4	34:YA:2476:A:O2'	2.34	0.61
41:YI:30:LEU:HB3	41:YI:36:ALA:HB3	1.80	0.61
54:YZ:181:GLU:O	54:YZ:182:LYS:O	2.19	0.61
1:QA:1358:U:C3'	14:QN:22:THR:HG21	2.30	0.61
1:QA:1491:G:H5''	12:QL:46:LYS:HG2	1.82	0.61
13:QM:57:ARG:O	13:QM:61:GLU:HB2	2.01	0.61
34:RA:1490:A:O2'	36:RD:99:ASP:OD1	2.19	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
53:RY:76:CYS:SG	53:RY:80:GLY:N	2.74	0.61
54:RZ:3:TYR:HB2	54:RZ:57:ILE:HG22	1.81	0.61
1:XA:263:A:P	20:XT:79:ARG:HE	2.24	0.61
10:XJ:10:GLY:HA3	10:XJ:16:LEU:CD2	2.22	0.61
1:QA:948:C:OP2	13:QM:106:ASN:CB	2.49	0.60
1:QA:1107:C:H5''	3:QC:173:VAL:H	1.65	0.60
1:QA:1150:U:O2	10:QJ:39:PRO:HG2	2.00	0.60
1:QA:1321:C:C4	19:QS:36:ARG:NH1	2.69	0.60
34:RA:626:U:H5'	34:RA:627:A:H5''	1.81	0.60
34:RA:1019:U:H2'	34:RA:1020:A:H8	1.66	0.60
35:RB:22:U:H3	35:RB:61:G:H1	1.49	0.60
1:XA:728:A:N7	15:XO:54:ARG:NE	2.49	0.60
34:YA:2046:G:N1	34:YA:2623:G:C2	2.62	0.60
1:QA:667:G:N2	15:QO:49:ASP:OD1	2.34	0.60
1:QA:1318:A:C6	14:QN:16:PHE:CG	2.89	0.60
34:RA:2296:U:OP2	47:RS:9:ARG:NH1	2.34	0.60
39:RG:72:ARG:HA	39:RG:87:PRO:HA	1.82	0.60
46:RR:74:LYS:HD3	46:RR:77:ARG:HH21	1.66	0.60
48:RT:77:PRO:HG2	48:RT:80:SER:HB3	1.82	0.60
1:XA:528:C:H41	12:XL:49:ASN:CG	2.04	0.60
1:XA:538:G:P	12:XL:115:LYS:HB2	2.41	0.60
1:XA:901:A:O5'	1:XA:901:A:H8	1.84	0.60
1:XA:1320:C:C6	19:XS:70:LYS:HD3	2.35	0.60
11:XK:109:VAL:HA	18:XR:86:VAL:HG23	1.83	0.60
12:XL:10:LEU:HB3	17:XQ:32:TYR:CE2	2.36	0.60
1:QA:595:G:H1'	1:QA:596:C:H5	1.65	0.60
1:QA:1320:C:H5'	19:QS:70:LYS:HE3	1.83	0.60
1:QA:1359:C:H5	14:QN:35:ARG:CZ	2.14	0.60
1:QA:1376:U:C4	7:QG:10:ARG:NH1	2.68	0.60
5:QE:80:ILE:CD1	8:QH:104:ARG:HH12	2.14	0.60
33:R9:18:ARG:HD2	34:RA:1034:G:H5'	1.83	0.60
34:RA:1700:A:H3'	34:RA:1701:A:C8	2.36	0.60
34:RA:2144:U:H4'	34:RA:2145:C:H5	1.66	0.60
1:XA:975:A:C2	14:XN:34:TYR:HE1	2.19	0.60
1:XA:1059:C:H5	3:XC:2:GLY:HA3	1.64	0.60
34:YA:508:G:O6	51:YW:9:TYR:CE1	2.55	0.60
34:YA:2069:G:C2	34:YA:2443:C:N3	2.69	0.60
39:YG:16:ARG:NH2	39:YG:28:VAL:O	2.35	0.60
1:QA:740:U:C3'	15:QO:39:LEU:HG	2.32	0.60
1:QA:1221:G:H5'	19:QS:36:ARG:HH12	1.65	0.60
11:QK:108:ILE:HG21	18:QR:88:LYS:OXT	2.02	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:642:G:H4'	34:RA:2349:G:H4'	1.83	0.60
34:RA:1604:C:H2'	34:RA:1605:C:H6	1.66	0.60
3:XC:23:TYR:CD2	10:XJ:95:GLU:CG	2.84	0.60
11:XK:17:GLY:HA2	11:XK:35:PRO:HD3	1.83	0.60
34:YA:1252:G:N2	49:YU:33:ARG:HH11	1.99	0.60
1:QA:667:G:H4'	15:QO:51:HIS:HE1	1.62	0.60
1:QA:982:U:C5'	14:QN:6:LEU:HD21	2.31	0.60
1:QA:1110:A:N1	3:QC:177:THR:HG22	2.16	0.60
34:RA:331:A:N6	34:RA:1210:A:OP2	2.34	0.60
34:RA:758:C:H2'	34:RA:759:G:H8	1.65	0.60
41:RI:92:VAL:HB	41:RI:120:ILE:CB	2.30	0.60
4:XD:25:ARG:NE	4:XD:30:LYS:O	2.33	0.60
37:YE:1:MET:HG3	37:YE:200:GLU:HG2	1.82	0.60
44:YP:58:THR:O	44:YP:61:ARG:NH2	2.34	0.60
1:QA:662:G:H2'	1:QA:663:A:C8	2.37	0.60
1:QA:810:C:H1'	1:QA:899:C:H41	1.66	0.60
1:QA:980:C:C2	14:QN:19:ARG:C	2.70	0.60
1:XA:595:G:H1	1:XA:641:U:HO2'	1.49	0.60
34:YA:458:G:N2	34:YA:470:A:OP2	2.34	0.60
34:YA:2051:A:N1	34:YA:2614:A:C5	2.69	0.60
51:YW:6:ILE:HG12	51:YW:104:THR:HG23	1.83	0.60
1:QA:1106:G:HO2'	3:QC:172:ARG:HD2	1.56	0.60
7:QG:138:LYS:HE2	7:QG:142:GLU:HG3	1.83	0.60
34:RA:299:A:N1	34:RA:322:A:O2'	2.29	0.60
34:RA:1295:C:H2'	34:RA:1296:G:H8	1.67	0.60
34:RA:1999:C:H2'	34:RA:2000:G:H8	1.67	0.60
1:XA:1360:A:N9	14:XN:17:LYS:HG3	2.17	0.60
3:XC:9:GLY:HA2	3:XC:12:LEU:HD13	1.83	0.60
34:YA:690:G:O2'	36:YD:43:ARG:NH1	2.30	0.60
34:YA:1094:U:H1'	34:YA:1097:U:H5	1.67	0.60
34:YA:2140:C:H2'	34:YA:2141:G:H8	1.66	0.60
51:YW:30:GLU:O	51:YW:34:ASN:ND2	2.35	0.60
1:QA:7:G:O2'	5:QE:121:LYS:HB2	2.01	0.60
1:QA:980:C:C4'	14:QN:19:ARG:NE	2.62	0.60
1:QA:1312:G:OP1	19:QS:5:LEU:O	2.20	0.60
1:QA:1329:A:N6	21:QU:7:ARG:HH22	1.99	0.60
2:XB:126:GLU:OE2	2:XB:130:ARG:NH1	2.34	0.60
4:XD:13:ARG:HG3	4:XD:40:PRO:HD3	1.84	0.60
34:YA:2080:G:N3	34:YA:2241:A:C2	2.69	0.60
34:YA:2515:C:H2'	34:YA:2516:G:H8	1.66	0.60
1:QA:609:A:C5'	16:QP:9:PHE:CD1	2.85	0.60

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1359:C:P	14:QN:22:THR:HG22	2.42	0.60
1:QA:1359:C:H5	14:QN:35:ARG:NE	1.98	0.60
2:QB:197:VAL:C	8:QH:68:ARG:HH22	2.03	0.60
41:RI:83:ALA:CB	41:RI:88:ILE:HA	2.31	0.60
46:RR:97:VAL:HG22	46:RR:114:VAL:HG12	1.84	0.60
1:XA:68(Q):C:H2'	1:XA:68(R):U:C6	2.37	0.60
1:XA:983:A:H5'	14:XN:2:ALA:HB3	1.84	0.60
2:XB:67:THR:HG21	2:XB:155:LEU:HG	1.84	0.60
2:XB:163:PHE:HA	2:XB:185:ILE:O	2.01	0.60
34:YA:662:G:OP1	44:YP:15:ARG:NH1	2.34	0.60
47:YS:4:LEU:HD11	47:YS:12:PHE:HE2	1.67	0.60
4:QD:88:VAL:CG2	5:QE:96:PRO:O	2.35	0.60
10:QJ:6:ILE:HG22	10:QJ:98:ILE:HG22	1.84	0.60
28:R4:5:ILE:O	39:RG:67:LYS:CG	2.50	0.60
34:RA:2094:G:H1'	34:RA:2198:A:H61	1.67	0.60
45:RQ:21:THR:HB	54:RZ:78:LYS:HE3	1.83	0.60
1:XA:757:U:H1'	1:XA:879:C:H1'	1.83	0.60
1:XA:900:A:O5'	1:XA:900:A:H8	1.85	0.60
34:YA:2090:G:O6	34:YA:2230:G:O6	2.20	0.60
38:YF:11:VAL:HG22	38:YF:125:LEU:HB2	1.84	0.60
40:YH:89:ILE:O	40:YH:129:THR:OG1	2.19	0.60
1:QA:676:A:N3	11:QK:119:CYS:SG	2.72	0.59
1:QA:947:G:OP2	13:QM:106:ASN:OD1	2.19	0.59
1:QA:980:C:H5''	1:QA:981:U:H5	1.67	0.59
1:QA:1307:U:OP2	13:QM:99:ARG:CB	2.50	0.59
7:QG:133:GLY:HA2	7:QG:136:LYS:HE2	1.83	0.59
32:R8:31:HIS:HD2	34:RA:2422:A:N6	2.00	0.59
41:RI:88:ILE:HG22	41:RI:90:GLY:H	1.66	0.59
10:XJ:16:LEU:HB3	10:XJ:70:ARG:HH11	1.64	0.59
10:XJ:47:PHE:HZ	14:XN:36:PHE:CG	2.20	0.59
30:Y6:6:ARG:HH21	30:Y6:24:GLU:HG3	1.65	0.59
34:YA:1566:A:C6	36:YD:214:TRP:CZ2	2.90	0.59
51:YW:33:ARG:NH2	51:YW:52:GLU:OE1	2.34	0.59
1:QA:1103:C:O2'	2:QB:111:ARG:NE	2.30	0.59
1:QA:1123:A:O2'	10:QJ:36:GLY:C	2.41	0.59
1:QA:1372:U:H5''	9:QL:69:GLY:HA3	1.84	0.59
34:RA:698:C:O2'	34:RA:734:A:N6	2.33	0.59
34:RA:1370:C:HO2'	34:RA:1811:G:HO2'	1.47	0.59
34:RA:2751:G:C6	40:RH:3:ARG:HB3	2.31	0.59
34:YA:1151:G:O2'	49:YU:77:SER:O	2.21	0.59
34:YA:2069:G:N1	34:YA:2443:C:N3	2.51	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:126:G:H4'	1:QA:634:C:H1'	1.84	0.59
1:QA:677:U:C2	11:QK:119:CYS:SG	2.95	0.59
34:RA:206:U:H2'	34:RA:207:A:H8	1.68	0.59
34:RA:1076:C:OP1	45:RQ:60:ARG:NH2	2.35	0.59
1:XA:972:C:O2'	10:XJ:55:LYS:HB2	2.02	0.59
1:XA:979:C:H42	14:XN:18:VAL:HG23	1.66	0.59
1:XA:983:A:H5'	14:XN:2:ALA:CB	2.31	0.59
34:YA:1152:C:H2'	34:YA:1153:C:H6	1.67	0.59
53:YY:79:CYS:HB2	53:YY:81:LYS:HG2	1.82	0.59
1:QA:564:C:C6	17:QQ:31:LEU:HD21	2.37	0.59
1:QA:1106:G:C3'	3:QC:172:ARG:HG3	2.24	0.59
1:QA:1220:G:H21	19:QS:54:GLY:HA2	1.66	0.59
3:QC:91:LEU:HD12	3:QC:101:LEU:HD11	1.85	0.59
32:R8:56:GLU:HA	32:R8:59:LYS:HE2	1.83	0.59
34:RA:788:A:OP1	34:RA:791:C:N4	2.34	0.59
34:RA:955:C:OP1	45:RQ:85:LYS:NZ	2.32	0.59
34:RA:1141:U:P	42:RN:25:ARG:HH21	2.25	0.59
37:RE:201:THR:HG22	37:RE:203:LYS:H	1.66	0.59
41:RI:83:ALA:HA	41:RI:89:TYR:H	1.67	0.59
1:XA:974:A:OP1	14:XN:31:ARG:HB2	2.02	0.59
1:XA:1068:G:H22	1:XA:1108:G:H1'	1.68	0.59
1:XA:1117:G:N2	1:XA:1180:A:N3	2.50	0.59
5:XE:79:GLU:HG3	5:XE:93:PRO:HD2	1.84	0.59
34:YA:1546:C:H5'	34:YA:1547:C:H5'	1.85	0.59
34:YA:2063:C:O2	34:YA:2450:A:N1	2.35	0.59
1:QA:1118:C:C5'	9:QI:9:ARG:NH1	2.66	0.59
1:QA:1187:G:O2'	14:QN:60:SER:HA	2.03	0.59
40:RH:89:ILE:HD11	40:RH:131:VAL:CG2	2.32	0.59
1:XA:51:A:C2	1:XA:353:A:N1	2.70	0.59
1:XA:398:C:H2'	1:XA:399:G:H8	1.67	0.59
1:XA:662:G:H2'	1:XA:663:A:H8	1.66	0.59
1:XA:1049:U:HO2'	14:XN:2:ALA:N	2.00	0.59
13:XM:3:ARG:O	13:XM:57:ARG:NH2	2.35	0.59
34:YA:2712(A):U:H2'	34:YA:2712(B):A:H2'	1.83	0.59
37:YE:143:ASN:OD1	37:YE:143:ASN:N	2.35	0.59
1:QA:1170:A:H5'	2:QB:140:HIS:CE1	2.37	0.59
1:QA:1186:G:N2	14:QN:61:TRP:CA	2.66	0.59
3:QC:23:TYR:N	10:QJ:93:GLY:C	2.55	0.59
29:R5:33:CYS:CB	29:R5:46:CYS:SG	2.90	0.59
34:RA:2086:U:OP2	36:RD:263:ARG:NH1	2.36	0.59
40:RH:107:VAL:O	40:RH:152:ARG:NH2	2.35	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1305:G:C5	1:XA:1331:G:N1	2.69	0.59
1:XA:1319:A:P	19:XS:3:ARG:CZ	2.91	0.59
8:XH:100:ILE:HD11	8:XH:125:ARG:HB3	1.84	0.59
34:YA:2080:G:N1	34:YA:2241:A:N1	2.50	0.59
1:QA:68:G:H22	1:QA:101:A:H2	1.49	0.59
1:QA:643:C:H2'	1:QA:644:G:C8	2.38	0.59
10:QJ:45:ARG:HB3	10:QJ:65:LEU:HB3	1.85	0.59
10:QJ:47:PHE:CZ	14:QN:36:PHE:CG	2.90	0.59
34:RA:184:C:H1'	34:RA:217:G:H1'	1.84	0.59
34:RA:2744:G:N2	40:RH:143:GLN:OE1	2.36	0.59
1:XA:165:C:H2'	1:XA:166:G:C8	2.38	0.59
1:XA:838(B):U:H4'	1:XA:838(C):C:C6	2.37	0.59
1:XA:1305:G:H2'	1:XA:1331:G:C2	2.36	0.59
1:XA:1318:A:C2	19:XS:37:ARG:NH1	2.71	0.59
32:Y8:31:HIS:CD2	34:YA:2421:G:O6	2.56	0.59
34:YA:956:G:H2'	34:YA:957:A:H2'	1.82	0.59
34:YA:2647:U:H2'	34:YA:2648:C:H6	1.67	0.59
36:YD:61:LEU:O	36:YD:63:ARG:NH1	2.36	0.59
41:YI:9:LEU:HD21	41:YI:35:LEU:HD12	1.84	0.59
1:QA:974:A:OP1	14:QN:29:ARG:CD	2.51	0.59
1:QA:1248:A:N6	21:QU:26:LYS:HD2	2.17	0.59
1:QA:1251:A:H2'	1:QA:1252:A:C8	2.37	0.59
1:QA:1304:G:H1'	1:QA:1334:G:H1	1.67	0.59
1:QA:1318:A:N6	14:QN:16:PHE:CB	2.65	0.59
4:QD:98:GLU:OE1	4:QD:103:ASN:ND2	2.36	0.59
34:RA:674:G:O2'	38:RF:67:GLN:OE1	2.18	0.59
34:RA:1996:C:H5	43:RO:32:TYR:HH	1.50	0.59
41:RI:93:THR:O	41:RI:97:ILE:HG13	2.03	0.59
46:RR:51:LEU:HG	46:RR:66:VAL:HG23	1.85	0.59
32:Y8:2:PRO:HA	34:YA:591:C:H1'	1.84	0.59
34:YA:2867:G:HO2'	34:YA:2868:A:H8	1.51	0.59
54:YZ:151:HIS:HB3	54:YZ:170:THR:HA	1.85	0.59
1:QA:986:A:H1'	19:QS:55:LYS:CA	2.27	0.59
1:QA:1097:C:H5'	2:QB:140:HIS:NE2	2.18	0.59
1:QA:1228:C:C5'	13:QM:111:LYS:NZ	2.42	0.59
1:QA:1320:C:H41	19:QS:37:ARG:HD2	1.68	0.59
28:R4:27:THR:CG2	39:RG:62:LEU:O	2.51	0.59
34:RA:746:A:O2'	34:RA:2611:U:O2'	2.18	0.59
34:RA:2250:G:C5	45:RQ:82:ARG:HD2	2.37	0.59
43:RO:88:ASN:ND2	43:RO:90:GLN:OE1	2.36	0.59
33:Y9:14:CYS:HA	33:Y9:27:CYS:HB2	1.84	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:727:A:C2	36:YD:9:TYR:CD2	2.89	0.59
1:QA:740:U:O3'	15:QO:39:LEU:HG	2.03	0.59
1:QA:1376:U:H2'	1:QA:1377:A:C8	2.38	0.59
4:QD:88:VAL:HG13	5:QE:97:GLY:HA2	1.85	0.59
1:XA:960:U:H5	19:XS:78:ARG:HG2	1.67	0.59
1:XA:979:C:OP1	1:XA:1223:C:N4	2.35	0.59
38:YF:168:ARG:HG2	38:YF:175:THR:HG21	1.84	0.59
1:QA:642:A:C1'	8:QH:115:SER:HA	2.33	0.58
1:QA:730:G:C5	1:QA:731:G:H1'	2.38	0.58
25:R1:78:LYS:HZ2	34:RA:270(S):G:H21	1.50	0.58
1:XA:571:U:H5''	1:XA:819:A:C4	2.38	0.58
1:XA:702:A:N1	34:YA:1848:A:C6	2.71	0.58
1:XA:1004:A:N6	1:XA:1025:U:O3'	2.36	0.58
1:XA:1318:A:H62	14:YN:16:PHE:CB	2.09	0.58
1:XA:1440(K):C:O2'	1:XA:1440(L):G:N2	2.33	0.58
1:XA:1541:U:N3	23:XX:5:A:C2	2.71	0.58
25:Y1:2:SER:N	34:YA:1364:G:N7	2.50	0.58
31:Y7:34:ARG:NH1	34:YA:466:A:OP1	2.36	0.58
34:YA:1582:C:H2'	34:YA:1583:A:H8	1.68	0.58
34:YA:2275:C:O2	45:YQ:83:MET:HG2	2.02	0.58
1:QA:123:C:H2'	1:QA:124:G:H8	1.67	0.58
1:QA:280:C:O2	17:QQ:40:LYS:NZ	2.30	0.58
1:QA:1306:A:OP2	21:QU:5:ASP:HA	2.03	0.58
10:QJ:47:PHE:CE1	14:QN:36:PHE:CB	2.81	0.58
34:RA:1996:C:H5	43:RO:32:TYR:OH	1.86	0.58
40:RH:152:ARG:HG2	40:RH:153:LYS:HG2	1.84	0.58
1:XA:1345:U:C4	1:XA:1377:A:H1'	2.38	0.58
1:QA:107:G:OP1	1:QA:325:A:N6	2.36	0.58
1:QA:936:C:H2'	1:QA:937:A:C8	2.39	0.58
2:QB:181:PHE:CE2	8:QH:70:GLN:HG2	2.38	0.58
4:QD:20:TYR:CZ	6:XF:14:LEU:HA	2.38	0.58
1:XA:59:A:H3'	1:XA:331:G:H22	1.67	0.58
1:XA:974:A:C8	14:YN:31:ARG:NE	2.71	0.58
1:XA:1105:A:H2'	1:XA:1106:G:C8	2.37	0.58
1:QA:44:G:OP2	16:QP:12:LYS:CE	2.52	0.58
1:QA:954:G:N3	19:QS:83:HIS:CE1	2.67	0.58
3:QC:108:ASN:ND2	3:QC:144:SER:OG	2.36	0.58
7:QG:16:LEU:HD22	9:QI:42:ARG:CA	2.33	0.58
10:QJ:53:PRO:CA	14:QN:42:ILE:HD11	2.33	0.58
13:QM:86:CYS:CB	19:QS:69:HIS:HE1	2.14	0.58
27:R3:17:LYS:HG3	34:RA:969:U:OP1	2.02	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:379:C:H2'	1:XA:380:G:H8	1.69	0.58
1:XA:974:A:H1'	14:XN:31:ARG:HH11	1.59	0.58
1:XA:1193:G:O2'	5:XE:21:ALA:O	2.21	0.58
34:YA:747:U:C4	34:YA:2613:U:C4	2.92	0.58
34:YA:1418:G:H2'	34:YA:1579:A:H61	1.66	0.58
37:YE:128:SER:OG	37:YE:129:HIS:N	2.35	0.58
1:QA:739:C:O2'	15:QO:42:HIS:ND1	2.30	0.58
1:QA:974:A:H5'	14:QN:31:ARG:CB	2.15	0.58
1:QA:1253:G:O5'	10:QJ:44:VAL:CB	2.42	0.58
1:QA:1353:G:OP1	21:QU:13:ILE:HG21	2.03	0.58
2:QB:78:GLN:O	2:QB:94:ASN:ND2	2.37	0.58
7:QG:16:LEU:HD21	9:QI:42:ARG:CD	2.23	0.58
34:RA:191:A:H1'	34:RA:679:C:H1'	1.86	0.58
34:RA:987:G:O2'	34:RA:1000:A:N3	2.33	0.58
34:RA:1024:G:O2'	34:RA:1144:G:O2'	2.22	0.58
34:RA:1076:C:H2'	34:RA:1077:A:H4'	1.86	0.58
34:RA:1226:G:OP1	50:RV:69:LYS:NZ	2.24	0.58
38:RF:148:LEU:HD13	38:RF:191:ARG:HH11	1.68	0.58
1:XA:1318:A:C1'	19:XS:11:VAL:CG2	2.71	0.58
1:XA:1535:C:N4	23:XX:10:G:C2	2.69	0.58
5:XE:18:ARG:NH1	5:XE:25:ARG:O	2.36	0.58
34:YA:1061:U:C6	34:YA:1062:G:H5''	2.35	0.58
36:YD:35:LYS:H	36:YD:64:ILE:HG12	1.68	0.58
1:QA:980:C:C2'	14:QN:19:ARG:HG2	2.23	0.58
4:QD:34:GLU:OE1	6:XF:57:GLN:NE2	2.36	0.58
4:QD:59:ARG:HH12	4:QD:66:ARG:HH22	1.51	0.58
5:QE:75:THR:OG1	5:QE:76:ILE:N	2.35	0.58
34:RA:1297:C:H2'	34:RA:1298:C:H6	1.68	0.58
34:RA:1566:A:C2	36:RD:214:TRP:CG	2.92	0.58
1:XA:1000:A:H2'	1:XA:1001:G:C8	2.38	0.58
3:XC:36:ASP:OD1	3:XC:59:ARG:NH2	2.34	0.58
34:YA:1457:A:C2	34:YA:2703:C:N4	2.71	0.58
1:QA:1097:C:O2'	1:QA:1169:A:N3	2.34	0.58
1:QA:1278:U:H5''	1:QA:1279:A:C8	2.38	0.58
34:RA:2684:U:O2'	43:RO:68:GLU:OE2	2.17	0.58
50:RV:24:LYS:HA	50:RV:92:THR:HG23	1.85	0.58
52:RX:25:LYS:HD3	52:RX:80:ILE:HD11	1.86	0.58
1:XA:501:C:H2'	1:XA:502:G:H8	1.68	0.58
1:XA:781:A:H3'	1:XA:782:A:H8	1.68	0.58
1:XA:1330:U:H5''	13:XM:24:GLY:CA	2.33	0.58
34:YA:530:G:C5	34:YA:2022:U:H5''	2.38	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:581:C:H2'	34:YA:582:G:C8	2.38	0.58
34:YA:863:A:H2'	34:YA:864:G:C8	2.39	0.58
37:YE:50:GLY:HA2	37:YE:77:ILE:HA	1.85	0.58
1:QA:974:A:N9	14:QN:31:ARG:CZ	2.67	0.58
1:QA:1186:G:H22	14:QN:61:TRP:HA	1.69	0.58
1:QA:1238:A:H2'	1:QA:1239:A:C8	2.38	0.58
1:QA:1484:C:O2'	34:RA:1960:A:O2'	2.20	0.58
34:RA:605:C:H1'	34:RA:657:U:H1'	1.85	0.58
34:RA:1011:G:HO2'	34:RA:1012:U:HO2'	1.51	0.58
34:RA:1930:G:H2'	34:RA:1968:G:H1	1.68	0.58
1:XA:15:G:H1	1:XA:920:U:H3	1.50	0.58
1:XA:412:A:C2	4:XD:35:ARG:CD	2.81	0.58
1:XA:1286:A:H2'	1:XA:1287:A:H4'	1.85	0.58
1:XA:1296:C:C5'	13:XM:14:ARG:NH2	2.63	0.58
11:XK:15:ALA:HA	11:XK:77:MET:HA	1.86	0.58
1:QA:6:G:C2	5:QE:119:LEU:HD11	2.39	0.58
1:QA:974:A:C8	14:QN:31:ARG:HD2	2.35	0.58
1:QA:1313:U:H5	19:QS:4:SER:CB	2.17	0.58
3:QC:22:TRP:C	10:QJ:93:GLY:HA2	2.23	0.58
1:XA:1270:C:H2'	1:XA:1271:G:C8	2.39	0.58
1:XA:1318:A:O3'	19:XS:11:VAL:HG21	2.00	0.58
1:XA:1343:G:H2'	1:XA:1344:C:C6	2.39	0.58
1:XA:1376:U:O4	7:XG:10:ARG:NE	2.37	0.58
17:XQ:99:SER:OG	17:XQ:100:LYS:N	2.37	0.58
19:XS:63:THR:OG1	19:XS:65:ASN:OD1	2.21	0.58
34:YA:577:G:O2'	34:YA:1254:A:OP1	2.21	0.58
34:YA:727:A:C2	36:YD:9:TYR:CE2	2.92	0.58
1:QA:19:C:OP1	5:QE:130:ASN:ND2	2.36	0.58
1:QA:25:C:H5'	1:QA:524:G:H1'	1.86	0.58
1:QA:689:C:OP2	11:QK:55:LYS:NZ	2.35	0.58
1:QA:741:G:P	15:QO:39:LEU:CD1	2.90	0.58
1:QA:1028(B):C:H2'	1:QA:1028(C):C:H5	1.69	0.58
1:QA:1371:G:OP1	9:QI:12:GLU:HG2	2.04	0.58
10:QJ:40:LEU:HD11	10:QJ:71:LEU:HB2	1.86	0.58
30:R6:16:CYS:SG	30:R6:42:TRP:HB2	2.44	0.58
34:RA:259:G:H21	34:RA:621:A:H8	1.52	0.58
41:RI:83:ALA:CA	41:RI:89:TYR:CD2	2.86	0.58
43:RO:1:MET:HB2	43:RO:32:TYR:HB3	1.84	0.58
53:RY:99:CYS:HB2	53:RY:103:GLY:N	2.18	0.58
1:XA:1306:A:H1'	1:XA:1332:A:C6	2.39	0.58
1:XA:1375:A:H3'	1:XA:1376:U:C6	2.37	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:XT:71:THR:OG1	20:XT:72:LEU:N	2.36	0.58
39:YG:29:TRP:O	39:YG:33:ARG:NH1	2.37	0.58
1:QA:1080:A:H4'	5:QE:16:THR:HG21	1.86	0.57
1:QA:1110:A:N6	3:QC:176:HIS:HB2	2.19	0.57
6:QF:23:LYS:NZ	6:QF:42:GLU:OE1	2.36	0.57
7:QG:50:ILE:HG12	7:QG:61:VAL:HG11	1.86	0.57
34:RA:992:C:OP1	49:RU:47:TYR:OH	2.18	0.57
39:RG:15:VAL:HG22	39:RG:175:LEU:HD22	1.86	0.57
41:RI:115:ALA:HB2	41:RI:131:LYS:HE3	1.85	0.57
54:RZ:30:ASN:HB3	54:RZ:90:VAL:HG22	1.86	0.57
1:XA:719:C:O2	18:XR:50:ILE:CD1	2.45	0.57
40:YH:9:ILE:HD12	40:YH:51:ARG:HG2	1.85	0.57
41:YI:3:VAL:HG12	41:YI:38:LEU:HA	1.85	0.57
34:RA:1689:A:OP2	34:RA:1698:A:N6	2.37	0.57
37:RE:26:ILE:HG23	37:RE:182:LEU:HB3	1.86	0.57
1:XA:766:A:H61	1:XA:1511:G:H1'	1.67	0.57
8:XH:11:THR:O	8:XH:15:ASN:ND2	2.38	0.57
20:XT:30:LYS:HA	20:XT:33:ILE:HD12	1.86	0.57
27:Y3:42:ALA:O	34:YA:851:U:O2'	2.20	0.57
34:YA:656:G:H2'	34:YA:657:U:C6	2.40	0.57
34:YA:2710:C:H2'	34:YA:2711:A:C8	2.38	0.57
46:YR:56:LYS:O	46:YR:88:ARG:NH2	2.36	0.57
48:YT:36:GLU:OE1	48:YT:41:ARG:NH1	2.37	0.57
1:QA:608:A:H3'	1:QA:609:A:H8	1.69	0.57
1:QA:937:A:H2'	1:QA:1379:G:H21	1.68	0.57
1:QA:1108:G:O3'	3:QC:176:HIS:HD2	1.87	0.57
1:QA:1114:C:C1'	14:QN:60:SER:O	2.51	0.57
1:QA:1364:U:O4'	21:QU:14:TRP:CZ2	2.57	0.57
12:QL:113:ARG:HH21	12:QL:116:SER:HB2	1.68	0.57
19:QS:61:TYR:OH	34:RA:888:C:OP2	2.21	0.57
33:R9:6:SER:HA	34:RA:1031:G:H4'	1.86	0.57
34:RA:1247:A:OP2	44:RP:15:ARG:NH2	2.38	0.57
49:RU:6:THR:OG1	49:RU:7:GLY:N	2.37	0.57
1:XA:1112:C:C1'	3:XC:179:ARG:CZ	2.79	0.57
1:XA:1313:U:O4	19:XS:4:SER:OG	2.22	0.57
1:XA:1539:C:H2'	1:XA:1540:U:C6	2.39	0.57
34:YA:307:G:H21	34:YA:330:A:H62	1.51	0.57
34:YA:664:C:OP1	44:YP:18:ARG:NH1	2.37	0.57
34:YA:1728:G:H8	34:YA:1732:A:H62	1.50	0.57
1:QA:968:A:N7	1:QA:1062:U:O2'	2.31	0.57
1:QA:1158:C:C5'	2:QB:133:LYS:HZ1	2.17	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1176:A:OP1	9:QI:97:LYS:HE2	2.04	0.57
1:QA:1320:C:N4	19:QS:37:ARG:CA	2.68	0.57
3:QC:23:TYR:CG	10:QJ:95:GLU:N	2.72	0.57
33:R9:19:ARG:NH2	34:RA:2754:U:O3'	2.36	0.57
34:RA:511:U:H4'	34:RA:1235:G:H4'	1.86	0.57
34:RA:1216:G:OP1	49:RU:11:ARG:NH2	2.31	0.57
34:RA:2394:C:OP1	44:RP:63:PRO:HD2	2.03	0.57
40:RH:149:ARG:HA	40:RH:162:ILE:HD11	1.85	0.57
1:XA:111:G:H8	1:XA:111:G:O5'	1.87	0.57
5:XE:11:ILE:HG21	5:XE:105:VAL:HG22	1.86	0.57
13:XM:3:ARG:HH12	13:XM:11:ARG:HH21	1.51	0.57
13:XM:84:ILE:HG23	19:XS:66:MET:HE1	1.85	0.57
29:Y5:43:HIS:N	34:YA:2884:U:O4	2.32	0.57
34:YA:911:A:C2	45:YQ:9:TYR:CD2	2.92	0.57
34:YA:996:A:O3'	49:YU:92:ARG:NH2	2.37	0.57
36:YD:28:GLU:HG2	36:YD:29:PRO:HD3	1.86	0.57
1:QA:1219:U:C2'	19:QS:34:TRP:CE3	2.87	0.57
1:QA:1242:C:H4'	1:QA:1303:C:H4'	1.86	0.57
10:QJ:47:PHE:CZ	14:QN:37:PHE:HD2	2.21	0.57
32:R8:46:ARG:NH1	34:RA:630:G:OP1	2.37	0.57
34:RA:2006:C:O2'	34:RA:2823:A:N3	2.35	0.57
34:RA:2419:U:H2'	34:RA:2420:C:H6	1.69	0.57
1:XA:6:G:N3	1:XA:6:G:H2'	2.20	0.57
1:XA:375:U:H5''	1:XA:375:U:H6	1.68	0.57
1:XA:669:U:C1'	15:XO:46:HIS:HE1	2.18	0.57
1:XA:730:G:C5	1:XA:731:G:H1'	2.40	0.57
3:XC:17:ASP:OD1	3:XC:21:ARG:NH2	2.37	0.57
6:XF:62:TRP:CD1	18:XR:35:ARG:NH1	2.72	0.57
34:YA:2044:C:N4	34:YA:2625:G:N1	2.51	0.57
1:QA:1073:U:O2'	2:QB:104:ASN:OD1	2.16	0.57
1:QA:1300:G:H1'	1:QA:1303:C:H42	1.70	0.57
1:QA:1320:C:N4	19:QS:37:ARG:CD	2.66	0.57
3:QC:23:TYR:HE1	10:QJ:10:GLY:HA2	1.66	0.57
13:QM:91:ARG:HD2	13:QM:96:LEU:HD22	1.87	0.57
34:RA:662:G:OP1	44:RP:15:ARG:NH1	2.37	0.57
34:RA:1204:A:H1'	34:RA:1206:G:C8	2.39	0.57
45:RQ:12:GLN:HB2	45:RQ:73:PRO:HD2	1.87	0.57
1:XA:401:C:H2'	1:XA:402:G:C8	2.40	0.57
1:XA:1320:C:N3	19:XS:36:ARG:C	2.58	0.57
1:XA:1320:C:C2	19:XS:36:ARG:O	2.57	0.57
1:XA:1320:C:C2'	19:XS:70:LYS:HD2	2.33	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
8:XH:8:ASP:OD2	8:XH:12:ARG:NH2	2.37	0.57
9:XI:10:ARG:HG3	9:XI:11:LYS:HG2	1.86	0.57
34:YA:184:C:O2'	34:YA:217:G:N3	2.35	0.57
34:YA:828:U:H4'	34:YA:831:G:N1	2.20	0.57
34:YA:1566:A:C2	36:YD:214:TRP:CE3	2.88	0.57
34:YA:2051:A:N1	34:YA:2614:A:N3	2.52	0.57
34:YA:2070:G:N2	34:YA:2442:C:N3	2.40	0.57
34:YA:2085:C:C2	34:YA:2235:G:C2	2.91	0.57
1:QA:1160:G:C4'	2:QB:132:LYS:HE3	2.35	0.57
3:QC:5:ILE:HG21	14:QN:45:ARG:NH2	2.19	0.57
4:QD:167:GLY:O	4:QD:169:LYS:NZ	2.36	0.57
4:QD:193:ASP:HA	6:XF:24:GLU:OE1	2.05	0.57
26:R2:4:SER:OG	26:R2:5:GLU:N	2.37	0.57
34:RA:1216:G:OP2	49:RU:12:ARG:NH2	2.36	0.57
34:RA:1854:A:H62	34:RA:1888:G:H8	1.51	0.57
1:XA:292:G:C5	1:XA:293:G:H1'	2.40	0.57
1:XA:590:C:OP1	8:XH:29:SER:HA	2.03	0.57
1:XA:728:A:H62	15:XO:54:ARG:CD	2.05	0.57
1:XA:741:G:OP1	15:XO:35:ARG:NE	2.38	0.57
1:XA:1251:A:O2'	1:XA:1369:C:O3'	2.21	0.57
1:XA:1296:C:H5'	13:XM:14:ARG:NH2	2.20	0.57
1:XA:1318:A:O3'	19:XS:11:VAL:CG2	2.50	0.57
2:XB:115:LEU:HD12	2:XB:145:LEU:HB3	1.86	0.57
3:XC:22:TRP:C	10:XJ:93:GLY:HA2	2.25	0.57
34:YA:86:C:OP1	53:YY:33:LYS:NZ	2.37	0.57
34:YA:270(P):U:C4	41:YI:52:ARG:NE	2.69	0.57
34:YA:617:G:OP2	38:YF:43:LYS:NZ	2.31	0.57
34:YA:639:U:H2'	34:YA:640:C:C6	2.40	0.57
34:YA:2049:G:N2	34:YA:2620:C:O2	2.36	0.57
1:QA:1306:A:OP2	21:QU:5:ASP:CA	2.53	0.57
2:QB:74:LYS:HG3	2:QB:77:ALA:HB3	1.87	0.57
2:QB:219:VAL:HA	2:QB:222:ILE:HD12	1.84	0.57
4:QD:9:CYS:SG	4:QD:22:LYS:NZ	2.71	0.57
34:RA:1889:A:N1	34:RA:2234:G:H1'	2.19	0.57
34:RA:2094:G:P	41:RI:22:LYS:HE3	2.44	0.57
39:RG:144:ILE:HG22	39:RG:146:TYR:H	1.69	0.57
1:XA:486:U:H2'	1:XA:487:A:H8	1.69	0.57
1:XA:922:G:H1'	5:XE:19:MET:HB2	1.85	0.57
1:XA:1318:A:O4'	19:XS:11:VAL:CG2	2.53	0.57
30:Y6:19:ARG:HH21	30:Y6:52:VAL:HG21	1.68	0.57
39:YG:37:VAL:HG13	39:YG:94:LEU:HB2	1.85	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
46:YR:57:ARG:NH1	46:YR:59:ASP:OD2	2.38	0.57
53:YY:30:VAL:HG22	53:YY:37:VAL:HG12	1.87	0.57
1:QA:248:C:H2'	1:QA:249:U:C6	2.40	0.57
1:QA:1226:C:C5'	13:QM:91:ARG:NH1	2.59	0.57
1:QA:1253:G:OP1	10:QJ:44:VAL:HG11	2.05	0.57
34:RA:2130:U:O2'	34:RA:2133:G:O2'	2.22	0.57
34:RA:2291:U:H2'	34:RA:2292:C:C6	2.40	0.57
1:XA:269:C:H2'	1:XA:270:A:C8	2.40	0.57
1:XA:401:C:H2'	1:XA:402:G:H8	1.69	0.57
1:XA:1495:U:HO2'	34:YA:1919:A:H2	1.53	0.57
12:XL:11:VAL:HG23	17:XQ:29:HIS:NE2	2.19	0.57
13:XM:11:ARG:O	13:XM:13:LYS:NZ	2.35	0.57
34:YA:1566:A:N3	36:YD:214:TRP:CG	2.73	0.57
36:YD:184:LYS:HB3	36:YD:269:PHE:HB3	1.86	0.57
1:QA:1119:C:H2'	1:QA:1120:G:H8	1.69	0.57
1:QA:1253:G:O2'	1:QA:1356:G:O2'	2.20	0.57
4:QD:166:LYS:O	36:YD:135:PHE:CE1	2.58	0.57
34:RA:2039:C:H2'	34:RA:2040:C:H6	1.70	0.57
1:XA:1300:G:C2	1:XA:1334:G:C6	2.93	0.57
10:XJ:53:PRO:C	14:YN:41:ARG:HH21	2.08	0.57
17:XQ:3:LYS:HB2	17:XQ:60:ILE:HD11	1.86	0.57
34:YA:2049:G:C2	34:YA:2620:C:O2	2.57	0.57
34:YA:2054:A:C2	34:YA:2616:C:C2	2.90	0.57
40:YH:155:SER:OG	40:YH:156:ALA:N	2.36	0.57
1:QA:69:G:H1'	1:QA:152:A:C2	2.40	0.56
1:QA:1414:U:H2'	1:QA:1415:G:C8	2.40	0.56
2:QB:87:ARG:NH1	2:QB:220:ASP:OD2	2.38	0.56
4:QD:166:LYS:O	36:YD:135:PHE:HE1	1.87	0.56
10:QJ:47:PHE:CD1	14:QN:37:PHE:HE2	2.22	0.56
31:R7:9:ARG:NE	34:RA:1310:G:OP2	2.38	0.56
34:RA:2751:G:N7	40:RH:2:SER:OG	2.37	0.56
40:RH:29:PRO:HD2	40:RH:79:VAL:HB	1.86	0.56
44:RP:58:THR:O	44:RP:61:ARG:NH2	2.38	0.56
54:RZ:127:LYS:HB3	54:RZ:162:GLU:HB2	1.86	0.56
1:XA:978:A:C5	14:YN:18:VAL:HG21	2.40	0.56
1:XA:1123:A:H1'	10:XJ:38:ILE:CG2	2.35	0.56
1:XA:1278:U:H5''	1:XA:1279:A:C8	2.39	0.56
1:XA:1537:U:O2	23:XX:9:G:N2	2.38	0.56
30:Y6:3:SER:OG	30:Y6:4:GLU:N	2.37	0.56
34:YA:1088:A:H4'	34:YA:1089:G:C8	2.40	0.56
34:YA:2056:G:C8	34:YA:2577:A:C6	2.93	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:689:C:H4'	1:QA:705:U:H1'	1.86	0.56
1:QA:1047:G:O2'	1:QA:1215:G:O2'	2.22	0.56
1:QA:1243:C:H5'	21:QU:9:ARG:HG3	1.87	0.56
9:QI:67:GLY:O	9:QI:73:GLN:NE2	2.38	0.56
33:R9:19:ARG:NH2	34:RA:2754:U:O2'	2.33	0.56
34:RA:1338:G:O6	52:RX:62:LYS:NZ	2.36	0.56
1:XA:51:A:O4'	1:XA:353:A:N7	2.38	0.56
1:XA:1228:C:H5	13:XM:103:THR:O	1.87	0.56
34:YA:917:A:H3'	34:YA:918:A:H8	1.70	0.56
1:QA:975:A:C2	14:QN:34:TYR:CE1	2.93	0.56
1:QA:1014:A:O4'	19:QS:34:TRP:CG	2.58	0.56
1:QA:1041:A:H2'	1:QA:1042:G:C8	2.40	0.56
1:QA:1280:A:N3	10:QJ:41:PRO:CD	2.68	0.56
1:QA:1331:G:O6	21:QU:7:ARG:HB2	2.05	0.56
1:QA:1342:C:H2'	1:QA:1343:G:C8	2.39	0.56
9:QI:64:THR:OG1	9:QI:66:ARG:NH1	2.37	0.56
34:RA:644:A:H2	34:RA:2369:A:H1'	1.71	0.56
34:RA:824:A:H1'	34:RA:2358:G:N7	2.20	0.56
34:RA:1248:G:C2	49:RU:3:ARG:HD2	2.40	0.56
53:RY:30:VAL:HG12	53:RY:37:VAL:HG23	1.87	0.56
1:XA:899:C:O2	1:XA:899:C:H2'	2.05	0.56
1:XA:1028(H):G:H2'	1:XA:1028(I):G:C8	2.39	0.56
1:XA:1105:A:H2'	1:XA:1106:G:H8	1.71	0.56
1:XA:1126:U:H3'	1:XA:1127:G:H8	1.69	0.56
1:XA:1184:G:H2'	1:XA:1185:G:H8	1.70	0.56
8:XH:73:ASP:OD1	8:XH:75:ARG:NH1	2.38	0.56
29:Y5:57:VAL:O	46:YR:33:ARG:NH2	2.30	0.56
32:Y8:31:HIS:HD2	34:YA:2421:G:O6	1.88	0.56
34:YA:270(C):A:H61	34:YA:270(Z):G:H1'	1.69	0.56
34:YA:1566:A:C6	36:YD:214:TRP:CZ3	2.92	0.56
34:YA:2096:U:H3	34:YA:2193:G:H22	1.53	0.56
34:YA:2511:U:O2'	37:YE:138:PRO:O	2.22	0.56
34:YA:2647:U:H2'	34:YA:2648:C:C6	2.39	0.56
37:YE:49:LEU:HD22	37:YE:81:ILE:HD11	1.85	0.56
45:YQ:24:GLY:H	45:YQ:101:ARG:HD2	1.70	0.56
46:YR:14:SER:OG	46:YR:15:SER:N	2.38	0.56
53:YY:76:CYS:HB3	53:YY:79:CYS:SG	2.45	0.56
33:R9:23:VAL:HG11	34:RA:1032:A:C4'	2.31	0.56
34:RA:383:U:H2'	34:RA:385:C:H5	1.71	0.56
34:RA:887:A:H1'	34:RA:889:C:C5	2.40	0.56
45:RQ:138:ASP:O	45:RQ:141:GLN:NE2	2.39	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:962:C:H2'	1:XA:963:G:H8	1.71	0.56
1:XA:1028(H):G:H2'	1:XA:1028(I):G:H8	1.70	0.56
1:XA:1077:G:N2	1:XA:1080:A:OP2	2.37	0.56
11:XK:91:ARG:HH21	18:XR:88:LYS:CD	2.03	0.56
34:YA:515:A:H1'	34:YA:581:C:H1'	1.88	0.56
1:QA:1189:C:O2'	3:QC:176:HIS:ND1	2.39	0.56
30:R6:52:VAL:HG22	30:R6:53:LYS:H	1.70	0.56
34:RA:2114:A:H3'	34:RA:2114:A:N3	2.20	0.56
34:RA:2140:C:H2'	34:RA:2141:G:H8	1.70	0.56
37:RE:39:PRO:HD3	37:RE:45:THR:HG22	1.87	0.56
1:XA:674:G:H2'	1:XA:675:A:C8	2.40	0.56
8:XH:32:LYS:HA	8:XH:35:ILE:HD12	1.86	0.56
34:YA:1802:A:H2'	34:YA:1803:A:C8	2.41	0.56
34:YA:2049:G:C2	34:YA:2620:C:N3	2.72	0.56
34:YA:2250:G:H21	34:YA:2250:G:P	2.27	0.56
1:QA:788:U:H3	1:QA:792:A:H2'	1.68	0.56
1:QA:865:A:N3	1:QA:918:A:O2'	2.35	0.56
1:QA:1358:U:H5'	14:QN:35:ARG:N	2.06	0.56
10:QJ:47:PHE:CE1	14:QN:36:PHE:HB3	2.19	0.56
31:R7:29:LYS:HA	31:R7:32:LYS:HG3	1.88	0.56
34:RA:83:G:N2	34:RA:103:A:OP2	2.39	0.56
37:RE:34:VAL:HG21	37:RE:77:ILE:HD11	1.88	0.56
1:XA:323:U:H3'	1:XA:324:G:C8	2.40	0.56
1:XA:1141:C:H2'	1:XA:1142:G:H8	1.71	0.56
1:XA:1305:G:C5	1:XA:1331:G:C5	2.92	0.56
34:YA:554:U:H2'	34:YA:556:G:H8	1.71	0.56
34:YA:958:U:OP2	45:YQ:14:ARG:NH1	2.39	0.56
33:R9:23:VAL:CG2	34:RA:1032:A:O2'	2.54	0.56
34:RA:1565:C:H1'	34:RA:1566:A:H8	1.70	0.56
38:RF:117:ARG:NH1	38:RF:120:GLU:OE2	2.38	0.56
38:RF:195:ASP:N	38:RF:195:ASP:OD1	2.37	0.56
1:XA:114:U:H1'	1:XA:353:A:H1'	1.86	0.56
1:XA:1270:C:O2'	1:XA:1314:C:OP1	2.24	0.56
34:YA:1204:A:H1'	34:YA:1206:G:C8	2.40	0.56
1:QA:1440(B):G:H4'	1:QA:1440(C):G:C4	2.40	0.56
3:QC:23:TYR:H	10:QJ:93:GLY:C	2.09	0.56
10:QJ:47:PHE:HE2	14:QN:34:TYR:CB	2.17	0.56
34:RA:1576:U:H2'	34:RA:1577:C:H6	1.71	0.56
41:RI:26:ALA:HA	41:RI:30:LEU:HB2	1.88	0.56
49:RU:50:ARG:O	49:RU:54:LYS:NZ	2.39	0.56
1:XA:728:A:C6	15:XO:54:ARG:CD	2.88	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1123:A:O3'	10:XJ:36:GLY:CA	2.49	0.56
34:YA:1493:C:C6	34:YA:2210:G:C5	2.94	0.56
34:YA:2097:C:C2	34:YA:2193:G:N2	2.73	0.56
34:YA:2134:A:H3'	34:YA:2135:A:H8	1.71	0.56
42:YN:63:THR:OG1	42:YN:64:GLY:N	2.39	0.56
54:YZ:52:SER:O	54:YZ:54:HIS:ND1	2.38	0.56
1:QA:377:G:H2'	1:QA:378:G:C8	2.41	0.56
1:QA:1313:U:C1'	19:QS:6:LYS:HZ3	2.18	0.56
32:R8:25:MET:HG3	44:RP:64:LYS:HB3	1.87	0.56
1:XA:501:C:H2'	1:XA:502:G:C8	2.41	0.56
1:XA:943:U:O2'	1:XA:1232:U:OP2	2.23	0.56
1:XA:989:C:H2'	1:XA:990:C:C6	2.40	0.56
1:XA:1312:G:H3'	19:XS:6:LYS:HZ3	0.75	0.56
1:XA:1377:A:C2	7:XG:8:GLU:O	2.59	0.56
1:XA:1503:A:N6	1:XA:1532:U:H1'	2.21	0.56
31:Y7:8:ASN:ND2	34:YA:770:G:OP1	2.38	0.56
34:YA:2051:A:C2	34:YA:2614:A:N3	2.73	0.56
43:YO:34:THR:CG2	43:YO:35:VAL:H	2.12	0.56
7:QG:16:LEU:HD11	9:QI:42:ARG:HH11	1.69	0.56
8:QH:12:ARG:HD2	8:QH:26:VAL:HG12	1.88	0.56
10:QJ:24:VAL:HG21	10:QJ:37:PRO:HG3	1.87	0.56
12:QL:71:PRO:O	12:QL:102:ARG:NH1	2.39	0.56
13:QM:85:GLY:O	19:QS:74:PHE:HD2	1.89	0.56
25:R1:90:ILE:CA	25:R1:94:LEU:HD12	2.34	0.56
34:RA:659:C:H2'	34:RA:660:G:C8	2.41	0.56
34:RA:1834:U:H1'	34:RA:1969:A:H2'	1.87	0.56
1:XA:12:U:H4'	1:XA:526:C:H4'	1.87	0.56
1:XA:458(B):G:H1'	1:XA:458(F):A:H61	1.71	0.56
24:Y0:19:LYS:NZ	34:YA:2261:C:OP1	2.31	0.56
28:Y4:6:HIS:CE1	39:YG:67:LYS:H	2.24	0.56
1:QA:129(B):G:H4'	1:QA:130:A:H5''	1.88	0.55
1:QA:1086:U:H2'	1:QA:1087:G:H8	1.71	0.55
1:QA:1369:C:H2'	1:QA:1370:G:C8	2.42	0.55
3:QC:5:ILE:HD12	14:QN:49:HIS:NE2	2.21	0.55
3:QC:22:TRP:HA	10:QJ:93:GLY:HA3	1.84	0.55
23:QX:8:A:H2'	23:QX:9:G:H8	1.71	0.55
34:RA:319:C:H2'	34:RA:320:A:C8	2.42	0.55
34:RA:2328:A:H2'	34:RA:2329:G:C8	2.41	0.55
42:RN:22:THR:OG1	42:RN:23:LEU:N	2.39	0.55
1:XA:686:U:O4	1:XA:703:G:O2'	2.20	0.55
29:Y5:48:GLU:OE2	51:YW:37:ARG:NH1	2.24	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:597:U:H2'	34:YA:598:G:C8	2.41	0.55
34:YA:2059:A:N1	34:YA:2503:A:C5	2.74	0.55
1:QA:265:G:H5'	17:QQ:64:PRO:O	2.06	0.55
1:QA:741:G:O5'	15:QO:39:LEU:HD11	2.06	0.55
34:RA:372:G:HO2'	34:RA:400:G:H1	1.55	0.55
34:RA:1380:G:O2'	34:RA:1569:A:N6	2.39	0.55
34:RA:2747:G:O6	34:RA:2755:C:H5''	2.06	0.55
43:RO:14:THR:HG21	43:RO:86:ILE:HD12	1.87	0.55
43:RO:106:LEU:HB3	43:RO:111:PHE:HB2	1.87	0.55
44:RP:90:ARG:HG3	44:RP:91:PHE:HD1	1.71	0.55
53:RY:47:LYS:NZ	53:RY:48:ALA:O	2.37	0.55
1:XA:107:G:H4'	1:XA:378:G:H5''	1.88	0.55
1:XA:192:U:H4'	20:XT:57:ARG:HD3	1.87	0.55
1:XA:686:U:O2'	1:XA:703:G:N2	2.40	0.55
1:XA:1228:C:C5	13:XM:103:THR:O	2.59	0.55
1:XA:1228:C:H5	13:XM:104:ARG:CA	1.98	0.55
1:XA:1318:A:H1'	19:XS:37:ARG:NH2	2.21	0.55
34:YA:2108:C:N4	34:YA:2182:G:H22	1.91	0.55
34:YA:2882:A:OP1	46:YR:96:ARG:NH1	2.39	0.55
1:QA:66:G:P	1:QA:66:G:H8	2.29	0.55
1:QA:310:G:OP2	16:QP:27:LYS:CD	2.52	0.55
4:QD:11:LEU:HD13	4:QD:66:ARG:HD2	1.88	0.55
34:RA:1380:G:H1'	34:RA:1569:A:H61	1.71	0.55
1:XA:8:A:N6	4:XD:209:ARG:HA	2.21	0.55
1:XA:261:U:C5	20:XT:79:ARG:CZ	2.88	0.55
1:XA:1358:U:O5'	1:XA:1358:U:H6	1.89	0.55
31:Y7:5:TRP:CZ3	34:YA:686:G:N7	2.74	0.55
34:YA:1030:G:OP2	45:YQ:128:LYS:NZ	2.27	0.55
39:YG:77:ILE:HG22	39:YG:82:LEU:HB2	1.88	0.55
41:YI:131:LYS:HG2	41:YI:135:GLU:HG3	1.87	0.55
48:YT:51:ARG:HD2	48:YT:100:TYR:HE1	1.72	0.55
1:QA:1160:G:C5'	2:QB:132:LYS:HE3	2.33	0.55
1:QA:1280:A:C4	10:QJ:41:PRO:HD2	2.40	0.55
33:R9:5:ALA:O	34:RA:1031:G:H1'	2.06	0.55
34:RA:784:A:N6	34:RA:2072:G:O2'	2.35	0.55
34:RA:2105:C:H2'	34:RA:2106:G:C8	2.42	0.55
35:RB:74:U:H1'	54:RZ:34:ASN:HD21	1.71	0.55
40:RH:33:LEU:HD11	40:RH:136:ILE:HG13	1.87	0.55
41:RI:86:THR:O	41:RI:122:GLU:HG2	2.05	0.55
1:XA:974:A:N9	14:XN:31:ARG:CZ	2.70	0.55
1:XA:1048:G:H2'	1:XA:1050:G:C8	2.42	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1305:G:C2	1:XA:1331:G:C8	2.95	0.55
24:Y0:16:SER:HB3	34:YA:2262:U:H5	1.70	0.55
34:YA:192:C:H1'	34:YA:800:A:H62	1.71	0.55
34:YA:468:G:H5''	38:YF:60:SER:HB2	1.88	0.55
1:QA:546:G:H4'	1:QA:548:G:H4'	1.89	0.55
1:QA:791:G:N1	1:QA:1498:U:OP1	2.38	0.55
1:QA:925:G:H1'	1:QA:1502:A:C4	2.42	0.55
1:QA:1124:G:C5'	10:QJ:36:GLY:H	2.20	0.55
4:QD:57:ARG:NH1	5:QE:107:ARG:CD	2.70	0.55
13:QM:11:ARG:HG3	13:QM:12:ASN:H	1.71	0.55
33:R9:14:CYS:HA	33:R9:27:CYS:HB2	1.88	0.55
34:RA:1571:A:H2'	34:RA:1572:A:C8	2.42	0.55
39:RG:126:ASP:OD2	39:RG:130:ASN:ND2	2.36	0.55
47:RS:25:ARG:HH21	47:RS:40:ILE:HG13	1.70	0.55
1:XA:1301:U:H4'	13:XM:17:VAL:CG2	2.36	0.55
1:XA:1316:G:C4'	14:YN:17:LYS:CG	2.31	0.55
1:XA:1473:A:H2'	1:XA:1474:G:C8	2.42	0.55
1:XA:1485:U:H2'	1:XA:1486:G:C8	2.41	0.55
3:XC:23:TYR:HD2	10:XJ:95:GLU:CB	2.19	0.55
34:YA:270(P):U:C4	41:YI:52:ARG:CZ	2.90	0.55
34:YA:746:A:H3'	34:YA:2612:C:H5	1.71	0.55
34:YA:828:U:O2'	34:YA:831:G:O6	2.17	0.55
34:YA:1791:A:N6	34:YA:1828:G:O2'	2.36	0.55
42:YN:129:PRO:O	42:YN:134:ARG:NH1	2.33	0.55
43:YO:87:ILE:HD12	43:YO:91:LEU:HA	1.89	0.55
48:YT:3:ARG:HG3	48:YT:6:LEU:HB2	1.89	0.55
1:QA:1000:A:H2'	1:QA:1001:G:C8	2.41	0.55
1:QA:1204:A:P	14:QN:3:ARG:NH2	2.79	0.55
1:QA:1239:A:H2	1:QA:1296:C:H5	1.53	0.55
13:QM:67:GLU:OE1	13:QM:71:ARG:NH1	2.40	0.55
26:R2:35:LEU:HD23	26:R2:50:ILE:HG12	1.88	0.55
31:R7:23:ARG:O	31:R7:28:ARG:NH1	2.40	0.55
43:RO:112:MET:HA	43:RO:115:VAL:HG22	1.87	0.55
47:RS:106:ARG:HA	47:RS:110:LEU:HD21	1.87	0.55
1:XA:1123:A:H2'	1:XA:1124:G:C4	2.41	0.55
10:XJ:47:PHE:CE2	14:YN:37:PHE:CZ	2.94	0.55
10:XJ:68:HIS:CD2	10:XJ:68:HIS:N	2.75	0.55
34:YA:34:C:N4	34:YA:454:A:O2'	2.38	0.55
34:YA:1566:A:H2	36:YD:214:TRP:NE1	2.03	0.55
34:YA:2377:A:H2'	34:YA:2378:A:C8	2.42	0.55
41:YI:130:TYR:HB3	41:YI:136:VAL:HG13	1.86	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1014:A:C1'	19:QS:34:TRP:CD2	2.90	0.55
1:QA:1218:C:H2'	1:QA:1219:U:H6	1.72	0.55
34:RA:318:C:H2'	34:RA:319:C:H6	1.71	0.55
38:RF:117:ARG:NH2	38:RF:189:THR:O	2.40	0.55
1:XA:68(Q):C:H2'	1:XA:68(R):U:H6	1.72	0.55
1:XA:186(L):G:H2'	1:XA:186(M):G:C8	2.42	0.55
1:XA:288:A:H2'	1:XA:289:G:H4'	1.87	0.55
1:XA:1162:C:H2'	1:XA:1163:C:C6	2.41	0.55
1:XA:1228:C:C5	13:XM:104:ARG:CB	2.89	0.55
34:YA:2229:C:H2'	34:YA:2230:G:C8	2.41	0.55
1:QA:1320:C:H42	19:QS:37:ARG:CB	2.18	0.55
1:QA:1376:U:C4	7:QG:10:ARG:CZ	2.90	0.55
4:QD:20:TYR:CZ	6:XF:14:LEU:CA	2.90	0.55
34:RA:644:A:C2	34:RA:2369:A:H1'	2.42	0.55
34:RA:1462:C:H4'	34:RA:2703:C:H5'	1.89	0.55
40:RH:86:GLU:HG3	40:RH:165:ALA:HB3	1.89	0.55
43:RO:19:ILE:HG22	43:RO:43:VAL:HG12	1.88	0.55
1:XA:405:U:H5''	1:XA:495:A:H2	1.71	0.55
1:XA:481:G:O2'	1:XA:483:C:N4	2.40	0.55
1:XA:1300:G:H1'	1:XA:1301:U:C6	2.42	0.55
1:XA:1359:C:N4	14:YN:35:ARG:NE	2.54	0.55
7:XG:29:LYS:HE2	7:XG:102:ARG:HB3	1.89	0.55
34:YA:236:C:H2'	34:YA:237:C:C6	2.42	0.55
34:YA:1872:A:C5	34:YA:1878:G:H1'	2.41	0.55
34:YA:2056:G:C6	34:YA:2577:A:N9	2.75	0.55
34:YA:2692:C:H2'	34:YA:2693:A:H8	1.72	0.55
1:QA:856:C:H2'	1:QA:857:C:C6	2.42	0.55
1:QA:1049:U:C4	14:QN:3:ARG:CB	2.77	0.55
1:QA:1221:G:H5'	19:QS:36:ARG:CZ	2.37	0.55
1:QA:1314:C:C5	19:QS:6:LYS:HE3	2.39	0.55
34:RA:1528:A:H2'	34:RA:1529:A:C8	2.42	0.55
1:XA:1229:A:H62	13:XM:104:ARG:CD	2.20	0.55
21:XU:3:LYS:HG2	21:XU:14:TRP:CB	2.36	0.55
43:YO:104:ARG:NH2	48:YT:36:GLU:OE2	2.36	0.55
1:QA:625:G:H5'	16:QP:10:GLY:N	2.21	0.55
14:QN:24:CYS:SG	14:QN:40:CYS:N	2.80	0.55
34:RA:831:G:N2	44:RP:53:GLY:O	2.40	0.55
34:RA:1299:G:H22	34:RA:1640:C:H5'	1.71	0.55
34:RA:1826:G:H4'	36:RD:242:ARG:HH21	1.72	0.55
34:RA:2476:A:H2'	34:RA:2477:C:C6	2.42	0.55
53:RY:6:HIS:O	53:RY:97:ARG:NH2	2.39	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1440(K):C:HO2'	1:XA:1440(L):G:H21	1.54	0.55
3:XC:23:TYR:HB2	10:XJ:93:GLY:C	2.27	0.55
10:XJ:69:ASN:N	10:XJ:69:ASN:OD1	2.39	0.55
11:XK:91:ARG:CD	18:XR:88:LYS:HZ1	2.15	0.55
34:YA:2080:G:N1	34:YA:2241:A:C2	2.74	0.55
34:YA:2134:A:H1'	34:YA:2158:A:N3	2.22	0.55
34:YA:2506:U:H2'	34:YA:2507:C:H6	1.72	0.55
38:YF:56:GLU:OE2	38:YF:93:LYS:NZ	2.36	0.55
53:YY:83:THR:HG21	53:YY:99:CYS:SG	2.48	0.55
1:QA:29:G:HO2'	1:QA:295:C:HO2'	1.47	0.54
1:QA:677:U:H2'	1:QA:678:U:C6	2.43	0.54
1:QA:947:G:H5'	13:QM:109:THR:HG23	1.90	0.54
1:QA:1253:G:P	10:QJ:44:VAL:HG23	2.46	0.54
2:QB:130:ARG:O	2:QB:135:GLN:NE2	2.37	0.54
34:RA:459:U:H2'	34:RA:460:A:H8	1.72	0.54
34:RA:1999:C:H1'	34:RA:2687:U:H1'	1.89	0.54
43:RO:104:ARG:NH2	43:RO:121:VAL:O	2.40	0.54
45:RQ:45:GLN:NE2	45:RQ:91:GLU:O	2.40	0.54
45:RQ:62:GLY:O	54:RZ:178:GLU:OE1	2.25	0.54
1:XA:745:C:H5''	1:XA:851:G:H1'	1.87	0.54
1:XA:941:G:O2'	1:XA:1350:A:OP1	2.24	0.54
1:XA:1541:U:O2	23:XX:5:A:C2	2.60	0.54
30:Y6:40:CYS:HB3	30:Y6:43:CYS:HB3	1.88	0.54
34:YA:482:A:H1'	34:YA:498:G:N2	2.22	0.54
34:YA:1456:G:N1	34:YA:2703:C:N4	2.18	0.54
34:YA:1493:C:C4	34:YA:2210:G:C4	2.95	0.54
34:YA:2056:G:C4	34:YA:2577:A:C6	2.95	0.54
41:YI:129:THR:HA	41:YI:137:PRO:HA	1.89	0.54
45:YQ:134:ARG:NH2	54:YZ:122:ARG:HD2	2.21	0.54
1:QA:954:G:H21	19:QS:83:HIS:CE1	2.23	0.54
1:QA:1221:G:P	19:QS:36:ARG:CZ	2.95	0.54
3:QC:5:ILE:HD12	14:QN:49:HIS:CD2	2.42	0.54
8:QH:64:LYS:HG2	8:QH:79:VAL:HG11	1.90	0.54
34:RA:504:U:H5''	34:RA:505:A:H5'	1.88	0.54
34:RA:747:U:H3'	34:RA:2612:C:H41	1.71	0.54
47:RS:34:HIS:ND1	47:RS:53:SER:OG	2.40	0.54
1:XA:1505:G:H5'	1:XA:1506:U:C6	2.43	0.54
2:XB:187:LEU:HA	2:XB:201:ILE:HB	1.90	0.54
14:XN:23:ARG:NH1	14:XN:24:CYS:O	2.40	0.54
34:YA:390:A:N1	44:YP:71:VAL:HG21	2.22	0.54
1:QA:309:G:C5'	16:QP:27:LYS:HE2	2.27	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:974:A:OP2	14:QN:29:ARG:HG3	2.03	0.54
3:QC:21:ARG:NH2	10:QJ:15:THR:HG21	2.22	0.54
34:RA:2312:U:O2'	39:RG:40:ASN:OD1	2.20	0.54
34:RA:2789:C:H1'	34:RA:2892:A:H2	1.72	0.54
43:RO:22:ILE:HB	43:RO:40:VAL:HG13	1.89	0.54
45:RQ:138:ASP:HB2	54:RZ:122:ARG:NH2	2.23	0.54
1:XA:1221:G:OP1	19:XS:36:ARG:HD3	2.07	0.54
33:Y9:16:VAL:HG12	33:Y9:25:VAL:HG12	1.88	0.54
34:YA:1275:A:O2'	34:YA:1645:G:N3	2.41	0.54
34:YA:1826:G:H4'	36:YD:242:ARG:NH2	2.22	0.54
34:YA:2275:C:O2	45:YQ:83:MET:CG	2.55	0.54
40:YH:2:SER:O	40:YH:2:SER:OG	2.24	0.54
1:QA:625:G:C5'	16:QP:9:PHE:HB3	2.36	0.54
1:QA:957:U:H5'	19:QS:81:ARG:CB	2.37	0.54
1:QA:1004:A:H2	1:QA:1024:G:H2'	1.71	0.54
1:QA:1060:C:C5	3:QC:2:GLY:HA3	2.42	0.54
5:QE:80:ILE:HD13	8:QH:104:ARG:HH12	1.73	0.54
12:QL:53:ARG:HB3	12:QL:69:TYR:HE1	1.73	0.54
1:XA:335:C:H2'	1:XA:336:C:C6	2.41	0.54
1:XA:379:C:H2'	1:XA:380:G:C8	2.43	0.54
1:XA:1112:C:C1'	3:XC:179:ARG:HE	2.18	0.54
1:XA:1396:A:H2	5:XE:19:MET:HG3	1.72	0.54
3:XC:23:TYR:HD2	10:XJ:95:GLU:CG	2.19	0.54
22:XV:19:G:C5	34:YA:2112:G:N7	2.75	0.54
34:YA:2836:U:H2'	34:YA:2837:G:C8	2.43	0.54
54:YZ:97:GLU:HB3	54:YZ:125:LEU:HD11	1.90	0.54
1:QA:111:G:O2'	1:QA:389:A:O2'	2.23	0.54
1:QA:624:C:H4'	16:QP:10:GLY:O	2.07	0.54
1:QA:981:U:H5''	14:QN:6:LEU:CD2	2.35	0.54
1:QA:1229:A:H61	13:QM:104:ARG:NH1	2.02	0.54
1:QA:1279:A:OP2	10:QJ:9:ARG:NH1	2.37	0.54
6:QF:97:PHE:CD2	18:QR:31:LEU:HD13	2.43	0.54
10:QJ:40:LEU:HD13	10:QJ:69:ASN:HB3	1.90	0.54
34:RA:1225:C:O2'	50:RV:86:GLY:N	2.34	0.54
34:RA:1445:C:H2'	34:RA:1446:C:H6	1.72	0.54
34:RA:2185:C:H2'	34:RA:2186:G:C8	2.42	0.54
34:RA:2312:U:H2'	34:RA:2313:C:H6	1.72	0.54
34:RA:2520:C:H2'	34:RA:2521:C:H6	1.72	0.54
37:RE:2:LYS:NZ	37:RE:100:GLU:OE2	2.40	0.54
51:RW:67:ASP:N	51:RW:67:ASP:OD1	2.39	0.54
54:RZ:128:VAL:HG23	54:RZ:161:VAL:HG12	1.88	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:123:C:H2'	1:XA:124:G:C8	2.43	0.54
1:XA:981:U:H3'	1:XA:982:U:C6	2.43	0.54
1:XA:1226:C:H3'	13:XM:103:THR:OG1	2.07	0.54
4:XD:57:ARG:HH12	5:XE:107:ARG:NH1	2.06	0.54
8:XH:86:ILE:HD11	8:XH:136:GLU:HG2	1.89	0.54
34:YA:319:C:H2'	34:YA:320:A:C8	2.43	0.54
39:YG:19:LEU:HD23	39:YG:32:PRO:HD2	1.89	0.54
54:YZ:149:SER:OG	54:YZ:150:LEU:N	2.40	0.54
1:QA:980:C:C4'	14:QN:19:ARG:HH21	2.16	0.54
1:QA:1014:A:OP1	19:QS:32:LYS:CE	2.56	0.54
1:QA:1414:U:H2'	1:QA:1415:G:H8	1.71	0.54
9:QI:46:ALA:HA	9:QI:78:LYS:HB3	1.88	0.54
28:R4:34:GLU:OE1	39:RG:113:ARG:NE	2.40	0.54
34:RA:1638:C:H1'	34:RA:2698:U:H1'	1.90	0.54
34:RA:2212:A:H1'	34:RA:2215:G:C4	2.43	0.54
50:RV:62:LEU:HB2	50:RV:93:GLU:HG3	1.88	0.54
1:XA:421:U:H4'	3:XC:192:THR:HG22	1.88	0.54
1:XA:1315:U:O3'	14:XN:17:LYS:HE2	1.68	0.54
3:XC:14:ILE:HG22	3:XC:15:THR:HG23	1.88	0.54
3:XC:22:TRP:HA	10:XJ:93:GLY:CA	2.38	0.54
34:YA:2108:C:C4	34:YA:2182:G:N2	2.76	0.54
51:YW:69:LEU:HD13	51:YW:107:LEU:HD23	1.89	0.54
1:QA:1198:G:H2'	1:QA:1199:U:C6	2.42	0.54
1:QA:1434:A:H61	1:QA:1467:G:H1'	1.72	0.54
34:RA:1221:C:H2'	34:RA:1222:C:C6	2.43	0.54
34:RA:2030:A:H4'	34:RA:2031:A:H8	1.71	0.54
38:RF:63:LYS:NZ	38:RF:75:HIS:O	2.33	0.54
49:RU:90:VAL:O	49:RU:92:ARG:N	2.35	0.54
1:XA:1041:A:H2'	1:XA:1042:G:C8	2.43	0.54
30:Y6:46:HIS:ND1	34:YA:2371:G:O2'	2.37	0.54
34:YA:1050:A:H8	34:YA:2751:G:N9	2.06	0.54
34:YA:2045:C:O2	34:YA:2624:G:C2	2.59	0.54
46:YR:104:ARG:NH1	46:YR:107:ASP:OD1	2.40	0.54
54:YZ:53:ILE:HG22	54:YZ:71:VAL:HG13	1.90	0.54
1:QA:243:A:H2	1:QA:282:A:H62	1.56	0.54
1:QA:413:G:H1'	1:QA:428:G:H21	1.72	0.54
1:QA:769:G:H4'	1:QA:1513:A:H4'	1.89	0.54
1:QA:979:C:OP1	1:QA:1223:C:N4	2.41	0.54
30:R6:46:HIS:ND1	34:RA:2371:G:O2'	2.28	0.54
33:R9:25:VAL:HB	33:R9:34:GLN:HB2	1.90	0.54
46:RR:33:ARG:NH2	46:RR:115:GLU:OE1	2.41	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:151:A:H3'	1:XA:152:A:H8	1.73	0.54
1:XA:571:U:H4'	1:XA:819:A:C6	2.42	0.54
1:XA:736:C:H2'	1:XA:737:A:C8	2.43	0.54
1:XA:994:A:H1'	1:XA:1216:G:H4'	1.88	0.54
2:XB:192:SER:OG	2:XB:193:ASP:N	2.40	0.54
11:XK:107:SER:HA	18:XR:87:ARG:NH1	2.23	0.54
27:Y3:8:LEU:HD13	27:Y3:23:LEU:HD11	1.90	0.54
34:YA:2646:C:OP2	34:YA:2732:G:O2'	2.24	0.54
54:YZ:11:GLU:O	54:YZ:36:LYS:NZ	2.36	0.54
1:QA:1004:A:C6	1:QA:1025:U:H4'	2.43	0.54
3:QC:108:ASN:HD22	3:QC:111:LEU:HD23	1.72	0.54
27:R3:13:ILE:HG21	34:RA:988:A:N6	2.23	0.54
34:RA:30:G:O2'	34:RA:1214:A:N3	2.38	0.54
34:RA:1397:U:OP2	34:RA:1398:C:N4	2.37	0.54
34:RA:2210:G:OP1	36:RD:68:LYS:NZ	2.41	0.54
1:XA:1181:G:C2	1:XA:1182:G:H1'	2.43	0.54
6:XF:89:MET:HE3	18:XR:76:LEU:HD13	1.90	0.54
22:XV:75:C:OP1	25:Y1:30:VAL:HG22	2.08	0.54
34:YA:358:U:H2'	34:YA:359:A:H8	1.72	0.54
34:YA:451:C:N4	34:YA:454:A:OP2	2.35	0.54
34:YA:2240:C:H2'	34:YA:2241:A:C8	2.43	0.54
45:YQ:45:GLN:NE2	45:YQ:91:GLU:O	2.40	0.54
1:QA:1360:A:C1'	14:QN:17:LYS:HZ1	2.19	0.54
3:QC:11:ARG:NH2	3:QC:177:THR:O	2.40	0.54
34:RA:247:G:H4'	34:RA:386:G:C4	2.43	0.54
34:RA:1275:A:O2'	34:RA:1645:G:N3	2.41	0.54
34:RA:1936:A:OP2	34:RA:1961:C:N4	2.41	0.54
1:XA:702:A:C6	34:YA:1848:A:C4	2.96	0.54
1:XA:1253:G:N3	1:XA:1254:C:C5	2.76	0.54
34:YA:197:A:C5	34:YA:2430:A:H2	2.09	0.54
34:YA:1086:A:O2'	34:YA:1103:A:N6	2.36	0.54
34:YA:1363:C:H2'	34:YA:1364:G:H8	1.73	0.54
54:YZ:108:PRO:HA	54:YZ:142:SER:HA	1.88	0.54
1:QA:1010:G:H2'	1:QA:1011:G:C8	2.42	0.53
1:QA:1280:A:N9	10:QJ:41:PRO:HD2	2.22	0.53
1:QA:1286:A:N6	1:QA:1355:G:OP1	2.41	0.53
1:QA:1305:G:O2'	1:QA:1332:A:N7	2.36	0.53
3:QC:23:TYR:CE1	10:QJ:9:ARG:C	2.82	0.53
9:QI:71:SER:HA	9:QI:74:ILE:HG12	1.90	0.53
34:RA:685:A:C2	34:RA:787:U:H1'	2.43	0.53
34:RA:2750:A:P	40:RH:59:ARG:HH12	2.31	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
41:RI:83:ALA:HA	41:RI:89:TYR:CD2	2.43	0.53
53:RY:19:LYS:HZ1	53:RY:20:TYR:HE2	1.47	0.53
1:XA:51:A:C4	1:XA:353:A:C5	2.96	0.53
1:XA:1230:C:N3	13:XM:102:ARG:NH1	2.55	0.53
1:XA:1304:G:H8	1:XA:1304:G:C5'	2.06	0.53
1:XA:1320:C:N3	19:XS:37:ARG:N	2.56	0.53
34:YA:911:A:C2	45:YQ:9:TYR:CD1	2.96	0.53
1:QA:137:C:H1'	16:QP:63:GLY:CA	2.29	0.53
1:QA:445:G:H2'	1:QA:446:G:H8	1.74	0.53
1:QA:699:C:H2'	1:QA:700:G:C8	2.43	0.53
1:QA:1080:A:P	5:QE:14:ARG:HH22	2.31	0.53
1:QA:1126:U:H5	1:QA:1148:U:H3	1.56	0.53
1:QA:1522:U:H2'	1:QA:1523:G:H8	1.72	0.53
3:QC:23:TYR:CZ	10:QJ:9:ARG:O	2.61	0.53
4:QD:98:GLU:HA	4:QD:103:ASN:HD22	1.73	0.53
27:R3:49:LYS:NZ	34:RA:851:U:OP1	2.38	0.53
34:RA:863:A:O3'	35:RB:100:G:N2	2.39	0.53
1:XA:68(B):G:H1	1:XA:68(Z):C:H42	1.56	0.53
1:XA:130:A:H4'	1:XA:186(K):G:C4	2.42	0.53
1:XA:592:G:H2'	1:XA:593:G:H8	1.73	0.53
1:XA:1502:A:H5'	1:XA:1504:G:N7	2.24	0.53
23:XX:6:G:H2'	23:XX:7:G:C8	2.44	0.53
34:YA:345:A:O2'	34:YA:347:A:N6	2.40	0.53
34:YA:746:A:O2'	34:YA:2611:U:O2'	2.18	0.53
34:YA:998:C:OP2	49:YU:58:ARG:NH1	2.41	0.53
34:YA:2074:U:HO2'	34:YA:2597:G:HO2'	1.55	0.53
34:YA:2319:G:O6	47:YS:3:ARG:O	2.25	0.53
34:YA:2391:G:H1'	34:YA:2429:G:H21	1.72	0.53
1:QA:237:C:H2'	1:QA:238:G:C8	2.43	0.53
1:QA:372:C:N4	1:QA:389:A:N7	2.56	0.53
1:QA:1048:G:OP1	14:QN:4:LYS:HB3	2.07	0.53
1:QA:1293:G:H2'	1:QA:1294:G:C8	2.43	0.53
28:R4:1:MET:N	35:RB:39:A:N1	2.57	0.53
29:R5:36:CYS:SG	29:R5:49:CYS:HB3	2.49	0.53
34:RA:551:G:H5'	34:RA:1220:A:H1'	1.90	0.53
36:RD:44:ASN:N	36:RD:44:ASN:OD1	2.41	0.53
47:RS:18:ILE:HD13	47:RS:88:ASP:HA	1.90	0.53
1:XA:8:A:N6	4:XD:209:ARG:CA	2.72	0.53
1:XA:34:C:H2'	1:XA:35:G:C8	2.43	0.53
1:XA:107:G:O6	20:XT:15:ARG:NH2	2.41	0.53
1:XA:232:G:H1'	1:XA:262:A:N1	2.24	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:948:C:P	13:XM:106:ASN:HB3	2.48	0.53
1:XA:1360:A:H1'	14:XN:17:LYS:HG3	1.89	0.53
4:XD:57:ARG:NH2	5:XE:107:ARG:HD3	2.23	0.53
6:XF:99:ALA:O	18:XR:28:GLU:CG	2.56	0.53
30:Y6:35:GLU:OE2	30:Y6:50:ARG:NH1	2.42	0.53
34:YA:820:A:H1'	34:YA:943:U:H1'	1.90	0.53
34:YA:2092:U:O4	34:YA:2228:G:O6	2.26	0.53
54:YZ:129:SER:OG	54:YZ:132:ASN:OD1	2.24	0.53
3:QC:23:TYR:CD1	10:QJ:9:ARG:O	2.60	0.53
3:QC:88:ARG:HG2	3:QC:101:LEU:HD13	1.90	0.53
4:QD:196:LEU:CA	6:XF:16:GLN:HG2	2.27	0.53
4:QD:205:GLU:HG2	5:QE:100:VAL:C	2.27	0.53
27:R3:45:GLY:HA3	34:RA:852:G:H5'	1.89	0.53
32:R8:25:MET:CE	44:RP:64:LYS:HD2	2.38	0.53
34:RA:631:A:H5'	44:RP:65:ARG:HG2	1.90	0.53
34:RA:668:G:H2'	34:RA:670:A:H62	1.74	0.53
1:XA:261:U:C5	20:XT:79:ARG:NH2	2.76	0.53
1:XA:652:U:O4	1:XA:752:G:O2'	2.26	0.53
1:XA:796:C:H2'	1:XA:797:C:H6	1.73	0.53
7:XG:71:PRO:O	7:XG:96:GLN:NE2	2.41	0.53
16:XP:37:GLY:HA3	16:XP:50:LYS:O	2.08	0.53
33:Y9:6:SER:HB3	34:YA:2466:C:H5''	1.89	0.53
33:Y9:29:ASN:ND2	33:Y9:32:HIS:CD2	2.77	0.53
34:YA:2228:G:OP1	36:YD:263:ARG:NH2	2.41	0.53
45:YQ:28:ALA:N	45:YQ:105:GLU:OE2	2.41	0.53
1:QA:691:G:O6	11:QK:52:GLY:HA2	2.08	0.53
1:QA:986:A:O4'	19:QS:55:LYS:HA	2.08	0.53
1:QA:1302:U:C1'	13:QM:27:LYS:CE	1.95	0.53
1:QA:1306:A:OP2	21:QU:5:ASP:HB2	2.08	0.53
4:QD:20:TYR:CE2	6:XF:14:LEU:HA	2.43	0.53
46:RR:56:LYS:NZ	46:RR:90:ARG:O	2.41	0.53
51:RW:14:PRO:HG2	51:RW:78:GLU:HG3	1.90	0.53
1:XA:901:A:C5	1:XA:902:G:H1'	2.44	0.53
1:XA:976:G:H21	1:XA:1362(B):C:H2'	1.73	0.53
1:XA:1305:G:N7	1:XA:1331:G:O6	2.41	0.53
34:YA:222:A:HO2'	34:YA:420:C:HO2'	1.56	0.53
34:YA:685:A:H5''	34:YA:788:A:N6	2.23	0.53
47:YS:26:LEU:HB3	47:YS:87:PHE:HA	1.90	0.53
1:QA:740:U:O3'	15:QO:39:LEU:CG	2.56	0.53
1:QA:1187:G:N2	14:QN:60:SER:OG	2.42	0.53
3:QC:29:TYR:CE1	14:QN:54:PRO:HG2	2.39	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:QN:6:LEU:HB3	14:QN:23:ARG:HH12	1.72	0.53
34:RA:1782:C:H1'	34:RA:2609:U:H5''	1.91	0.53
34:RA:2849:U:C2	34:RA:2867:G:H1'	2.44	0.53
38:RF:107:LYS:HE3	38:RF:207:GLY:H	1.73	0.53
38:RF:185:ASP:OD1	38:RF:188:ARG:NH1	2.42	0.53
45:RQ:81:VAL:O	45:RQ:82:ARG:NE	2.34	0.53
34:YA:389:G:H1	44:YP:71:VAL:H	1.57	0.53
34:YA:749:C:H4'	34:YA:1271:G:N3	2.23	0.53
34:YA:1457:A:N1	34:YA:2703:C:N4	2.57	0.53
34:YA:2071:A:H2'	34:YA:2072:G:C8	2.44	0.53
41:YI:68:LEU:HA	41:YI:71:ILE:HG22	1.90	0.53
1:QA:539:A:OP2	12:QL:115:LYS:HD3	2.08	0.53
1:QA:878:G:H5''	8:QH:90:GLY:N	2.23	0.53
1:QA:973:G:C4'	14:QN:29:ARG:NH2	2.53	0.53
1:QA:1048:G:OP1	14:QN:4:LYS:CE	2.56	0.53
4:QD:205:GLU:CG	5:QE:100:VAL:O	2.48	0.53
27:R3:12:PRO:HA	27:R3:15:TYR:HD2	1.73	0.53
30:R6:9:LEU:HD13	30:R6:51:GLU:HB2	1.90	0.53
34:RA:642:G:H21	34:RA:646:A:H2	1.56	0.53
34:RA:1815:A:OP2	36:RD:54:ARG:NH2	2.42	0.53
34:RA:2345:G:H1'	34:RA:2382:G:H5'	1.91	0.53
1:XA:481:G:N3	1:XA:482:A:N6	2.56	0.53
1:XA:539:A:OP1	12:XL:114:LYS:CD	2.52	0.53
1:XA:1106:G:H5'	3:XC:172:ARG:CG	2.38	0.53
31:Y7:37:LYS:HE3	34:YA:458:G:C5	2.44	0.53
34:YA:288:C:H2'	34:YA:289:A:H8	1.73	0.53
34:YA:882:G:H2'	34:YA:883:G:H8	1.74	0.53
34:YA:1061:U:H4'	34:YA:1070:A:H1'	1.90	0.53
40:YH:103:LEU:HB3	40:YH:115:VAL:HG22	1.91	0.53
1:QA:797:C:H2'	1:QA:798:G:H8	1.74	0.53
1:QA:1318:A:H62	14:QN:16:PHE:HB3	1.74	0.53
7:QG:114:ARG:O	7:QG:119:ARG:NH2	2.42	0.53
14:QN:24:CYS:HB2	14:QN:28:GLY:H	1.73	0.53
25:R1:41:ARG:NH2	34:RA:1365:A:O4'	2.40	0.53
34:RA:2107:C:H2'	34:RA:2108:C:C6	2.44	0.53
34:RA:2698:U:H2'	34:RA:2699:C:C6	2.43	0.53
37:RE:171:GLU:HB3	37:RE:185:LYS:HE2	1.90	0.53
1:XA:745:C:H2'	1:XA:746:A:C8	2.43	0.53
2:XB:132:LYS:HA	2:XB:135:GLN:HB2	1.89	0.53
10:XJ:37:PRO:O	10:XJ:37:PRO:CG	2.52	0.53
34:YA:191:A:H2'	34:YA:192:C:C6	2.44	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:2680:C:H5'	37:YE:189:PRO:HA	1.90	0.53
49:YU:28:ARG:NH1	49:YU:38:THR:OG1	2.41	0.53
1:QA:229:U:H2'	1:QA:230:G:C8	2.44	0.53
1:QA:980:C:O2	14:QN:21:TYR:CE1	2.62	0.53
2:QB:109:SER:O	2:QB:113:HIS:ND1	2.41	0.53
3:QC:108:ASN:HB3	3:QC:111:LEU:HB2	1.89	0.53
29:R5:49:CYS:SG	29:R5:50:GLY:N	2.82	0.53
34:RA:392:C:H5''	34:RA:409:C:H5''	1.91	0.53
35:RB:7:G:H21	47:RS:38:GLN:HE22	1.57	0.53
1:XA:996:A:N1	1:XA:1046:A:H1'	2.24	0.53
1:XA:1260:C:O2'	1:XA:1283:G:O2'	2.23	0.53
1:XA:1279:A:O2'	1:XA:1282:C:N4	2.42	0.53
34:YA:699:A:O3'	34:YA:1554:A:N6	2.42	0.53
34:YA:793:A:OP2	34:YA:2071:A:O2'	2.27	0.53
34:YA:1050:A:C8	34:YA:2751:G:N3	2.76	0.53
34:YA:1266:G:C8	51:YW:15:ARG:NH2	2.76	0.53
34:YA:1999:C:H2'	34:YA:2000:G:H8	1.74	0.53
34:YA:2317:C:H3'	34:YA:2318:G:H21	1.74	0.53
1:QA:160:A:H1'	1:QA:344:A:C5	2.45	0.53
1:QA:861:G:HO2'	1:QA:874:G:HO2'	1.57	0.53
1:QA:1314:C:C5	19:QS:6:LYS:CE	2.92	0.53
1:QA:1331:G:P	13:QM:24:GLY:H	2.32	0.53
7:QG:16:LEU:CD2	9:QI:42:ARG:CA	2.87	0.53
34:RA:477:A:N6	34:RA:500:G:O2'	2.42	0.53
34:RA:828:U:H4'	34:RA:831:G:N1	2.24	0.53
34:RA:1935:G:H1'	34:RA:1964:G:N2	2.24	0.53
34:RA:2094:G:OP1	41:RI:22:LYS:CE	2.54	0.53
42:RN:39:ARG:NH1	42:RN:48:MET:SD	2.82	0.53
26:Y2:29:LYS:NZ	52:YX:6:ASP:OD2	2.32	0.53
28:Y4:11:PRO:HA	28:Y4:25:TYR:HA	1.90	0.53
34:YA:758:C:H2'	34:YA:759:G:H8	1.73	0.53
34:YA:1958:C:H2'	34:YA:1959:G:H8	1.74	0.53
34:YA:2630:G:N3	34:YA:2892:A:O2'	2.42	0.53
43:YO:120:GLU:OE1	48:YT:67:SER:OG	2.28	0.53
1:QA:643:C:O2'	8:QH:132:GLU:OE1	2.24	0.52
1:QA:675:A:O2'	11:QK:115:PRO:HA	2.09	0.52
1:QA:1223:C:P	19:QS:78:ARG:CZ	2.93	0.52
34:RA:589:C:H2'	34:RA:590:A:C8	2.44	0.52
34:RA:1028:A:H2'	34:RA:1029:A:C8	2.44	0.52
34:RA:1668:A:H2	34:RA:1675:C:H41	1.56	0.52
34:RA:2133:G:N2	34:RA:2157:G:H2'	2.24	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:2291:U:H1'	34:RA:2374:C:H1'	1.91	0.52
46:RR:103:ARG:NH1	46:RR:108:GLY:O	2.40	0.52
50:RV:76:LYS:HB2	50:RV:81:TYR:HB3	1.90	0.52
1:XA:1282:C:H2'	1:XA:1283:G:O4'	2.09	0.52
1:XA:1378:C:O2	1:XA:1378:C:H2'	2.09	0.52
1:XA:1505:G:H5'	1:XA:1506:U:H6	1.74	0.52
1:XA:1510:U:H2'	1:XA:1511:G:C8	2.44	0.52
10:XJ:64:GLU:O	14:XN:56:VAL:CG2	2.57	0.52
34:YA:1500:G:O2'	36:YD:100:GLY:O	2.27	0.52
34:YA:2453:A:H2'	34:YA:2454:G:H8	1.73	0.52
34:YA:2626:C:H2'	34:YA:2627:G:C8	2.44	0.52
36:YD:17:THR:HB	36:YD:205:VAL:H	1.74	0.52
1:QA:677:U:H1'	11:QK:119:CYS:SG	2.49	0.52
1:QA:762:C:H2'	1:QA:763:G:H8	1.75	0.52
1:QA:985:C:H2'	1:QA:986:A:C8	2.44	0.52
1:QA:1305:G:H2'	1:QA:1331:G:C2	2.43	0.52
10:QJ:33:GLN:O	10:QJ:76:ASN:ND2	2.43	0.52
19:QS:4:SER:OG	19:QS:5:LEU:N	2.41	0.52
34:RA:390:A:H1'	34:RA:391:G:C8	2.44	0.52
34:RA:2006:C:H2'	34:RA:2007:C:C6	2.44	0.52
34:RA:2377:A:H2'	34:RA:2378:A:C8	2.44	0.52
34:RA:2729:G:H1'	37:RE:187:ALA:HB2	1.91	0.52
1:XA:129(B):G:H1'	1:XA:186(J):U:H2'	1.92	0.52
1:XA:422:C:H1'	1:XA:423:G:N1	2.25	0.52
1:XA:520:A:C2	1:XA:536:C:H1'	2.44	0.52
1:XA:1216:G:C5'	14:XN:5:ALA:HB2	2.40	0.52
1:XA:1440(F):G:N2	1:XA:1440(P):A:H1'	2.24	0.52
31:Y7:37:LYS:CG	34:YA:458:G:C8	2.93	0.52
34:YA:1094:U:H1'	34:YA:1097:U:C5	2.44	0.52
34:YA:1728:G:H2'	34:YA:1731:G:O6	2.09	0.52
34:YA:2479:G:OP1	34:YA:2537:U:O2'	2.25	0.52
1:QA:217:C:H2'	1:QA:218:C:C6	2.45	0.52
1:QA:825:G:H2'	1:QA:826:C:C6	2.45	0.52
1:QA:855:G:H21	1:QA:1539:C:H5''	1.74	0.52
1:QA:1028(H):G:H2'	1:QA:1028(I):G:C8	2.45	0.52
1:QA:1048:G:OP1	14:QN:4:LYS:CB	2.56	0.52
2:QB:209:ARG:NH1	2:QB:240:GLN:OE1	2.43	0.52
29:R5:46:CYS:HB3	29:R5:49:CYS:SG	2.50	0.52
34:RA:1252:G:H21	49:RU:33:ARG:CZ	2.22	0.52
34:RA:1252:G:H21	49:RU:33:ARG:NH2	2.07	0.52
34:RA:2245:U:H5'	34:RA:2246:G:H5'	1.91	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:2695:C:H2'	34:RA:2696:U:H6	1.74	0.52
40:RH:125:VAL:HG22	40:RH:125:VAL:O	2.08	0.52
54:RZ:27:VAL:HG22	54:RZ:85:HIS:HE1	1.74	0.52
1:XA:608:A:C3'	1:XA:609:A:H8	2.22	0.52
1:XA:860:A:H3'	1:XA:861:G:H8	1.74	0.52
1:XA:974:A:OP2	14:XN:31:ARG:HB2	2.08	0.52
1:XA:1500:A:H5''	1:XA:1508:G:H5'	1.92	0.52
3:XC:32:LEU:O	3:XC:59:ARG:NH2	2.42	0.52
6:XF:50:TYR:HE1	18:XR:77:GLY:O	1.67	0.52
34:YA:691:C:H4'	36:YD:43:ARG:HD3	1.91	0.52
36:YD:67:PHE:HE1	36:YD:106:ILE:HD11	1.75	0.52
39:YG:170:ARG:NH1	39:YG:174:GLU:OE1	2.43	0.52
1:QA:561:U:H5''	1:QA:563:A:N7	2.25	0.52
1:QA:981:U:C5'	14:QN:6:LEU:HD22	2.37	0.52
13:QM:84:ILE:CG1	19:QS:74:PHE:HZ	2.20	0.52
34:RA:2619:C:H2'	34:RA:2620:C:H6	1.73	0.52
39:RG:166:ASP:OD2	39:RG:166:ASP:N	2.42	0.52
44:RP:29:LYS:HD3	44:RP:30:THR:HG23	1.92	0.52
1:XA:960:U:C5	19:XS:78:ARG:HG2	2.45	0.52
1:XA:1301:U:HO2'	13:XM:17:VAL:HG21	1.67	0.52
11:XK:83:ILE:HD13	11:XK:109:VAL:HB	1.90	0.52
11:XK:91:ARG:NH2	18:XR:88:LYS:CD	2.69	0.52
13:XM:84:ILE:CD1	19:XS:65:ASN:CG	2.61	0.52
18:XR:47:THR:HG22	18:XR:85:LEU:HD13	1.91	0.52
34:YA:2789:C:H1'	34:YA:2892:A:H2	1.75	0.52
37:YE:176:ILE:HG13	37:YE:181:LEU:HB2	1.92	0.52
1:QA:1218:C:H2'	1:QA:1219:U:C6	2.45	0.52
1:QA:1327:C:OP1	21:QU:19:GLY:HA2	2.09	0.52
24:R0:7:LEU:HD12	45:RQ:83:MET:SD	2.50	0.52
34:RA:137(B):G:H21	52:RX:41:ASN:HD21	1.57	0.52
34:RA:787:U:H5''	34:RA:788:A:H5'	1.91	0.52
34:RA:1567:A:C8	36:RD:84:TYR:HE2	2.27	0.52
1:XA:908:A:H2'	1:XA:909:A:C8	2.45	0.52
1:XA:1300:G:N1	1:XA:1334:G:C5	2.78	0.52
34:YA:2757:A:N1	40:YH:67:LEU:HD13	2.24	0.52
1:QA:28:G:N3	1:QA:296:U:H4'	2.24	0.52
1:QA:186(Q):U:H2'	1:QA:191:G:C8	2.45	0.52
1:QA:543:C:OP2	4:QD:10:ARG:NH2	2.41	0.52
1:QA:637:G:H2'	1:QA:638:G:C8	2.44	0.52
1:QA:983:A:O4'	14:QN:2:ALA:HB3	2.10	0.52
29:R5:33:CYS:N	29:R5:38:ALA:O	2.28	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:309:G:N3	34:RA:329:G:O2'	2.42	0.52
34:RA:996:A:H2'	34:RA:997:G:H8	1.75	0.52
39:RG:41:GLN:HB2	39:RG:90:LEU:HB2	1.92	0.52
43:RO:24:VAL:HG13	43:RO:33:ALA:HB2	1.91	0.52
54:RZ:24:LEU:HD11	54:RZ:83:PRO:HB2	1.90	0.52
1:XA:21:G:H2'	1:XA:22:G:C8	2.45	0.52
1:XA:608:A:H3'	1:XA:609:A:H8	1.75	0.52
1:XA:975:A:C2	14:XN:34:TYR:CE1	2.97	0.52
1:XA:1540:U:C2	23:XX:6:G:N2	2.78	0.52
7:XG:18:TYR:OH	7:XG:47:CYS:SG	2.64	0.52
34:YA:2085:C:N3	34:YA:2235:G:C2	2.78	0.52
47:YS:61:ASN:ND2	47:YS:64:GLU:OE1	2.42	0.52
1:QA:123:C:H2'	1:QA:124:G:C8	2.44	0.52
1:QA:718:G:C5'	11:QK:117:ASN:OD1	2.58	0.52
1:QA:973:G:H4'	14:QN:29:ARG:NH2	2.22	0.52
7:QG:111:ARG:HH11	7:QG:119:ARG:HA	1.73	0.52
12:QL:85:ILE:HD11	12:QL:98:TYR:HB3	1.92	0.52
34:RA:144:C:H2'	34:RA:145:G:C8	2.45	0.52
34:RA:225:A:N6	34:RA:419:C:O2'	2.43	0.52
48:RT:30:VAL:HG12	48:RT:86:ILE:HG23	1.92	0.52
1:XA:1253:G:O3'	10:XJ:45:ARG:CD	2.58	0.52
5:XE:76:ILE:HB	5:XE:142:LEU:HD21	1.92	0.52
22:XV:1:C:H2'	22:XV:2:G:H8	1.75	0.52
26:Y2:48:HIS:CG	34:YA:96:G:H4'	2.45	0.52
34:YA:410:G:C2	34:YA:2407:G:N7	2.78	0.52
34:YA:2168:G:N2	34:YA:2170:A:H62	2.07	0.52
34:YA:2311:A:N7	39:YG:44:GLY:HA3	2.25	0.52
1:QA:310:G:OP1	16:QP:27:LYS:CD	2.52	0.52
1:QA:377:G:H2'	1:QA:378:G:H8	1.75	0.52
1:QA:797:C:H2'	1:QA:798:G:C8	2.45	0.52
1:QA:946:A:P	13:QM:114:ARG:HH21	2.30	0.52
1:QA:994:A:C8	1:QA:1216:G:H1'	2.45	0.52
1:QA:1108:G:O3'	3:QC:176:HIS:CD2	2.63	0.52
1:QA:1162:C:H2'	1:QA:1163:C:C6	2.44	0.52
1:QA:1279:A:H2	10:QJ:43:ARG:HH12	1.57	0.52
1:QA:1304:G:H5''	21:QU:10:ARG:NH2	2.24	0.52
10:QJ:47:PHE:HD2	14:QN:34:TYR:CD2	2.28	0.52
34:RA:323:G:C2	34:RA:333:G:H1'	2.45	0.52
34:RA:1752:C:H2'	34:RA:1753:G:C8	2.45	0.52
34:RA:1938:A:C6	34:RA:2590:A:H1'	2.45	0.52
34:RA:2730:C:H2'	34:RA:2731:G:C8	2.44	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
35:RB:55:U:O3'	39:RG:27:ASN:ND2	2.42	0.52
38:RF:65:TRP:NE1	38:RF:73:ALA:O	2.42	0.52
48:RT:6:LEU:HA	48:RT:9:LEU:HB2	1.91	0.52
1:XA:770:C:H2'	1:XA:771:G:H8	1.75	0.52
1:XA:899:C:P	1:XA:899:C:C6	3.03	0.52
1:XA:974:A:P	14:XN:29:ARG:CZ	2.98	0.52
1:XA:1124:G:P	10:XJ:36:GLY:HA3	2.50	0.52
5:XE:110:LEU:HD13	5:XE:118:ILE:HG21	1.90	0.52
7:XG:92:SER:O	7:XG:96:GLN:HB2	2.10	0.52
34:YA:28:A:N6	34:YA:512:G:H1'	2.25	0.52
34:YA:586:A:H5'	38:YF:89:VAL:HG21	1.91	0.52
34:YA:1608:A:O2'	34:YA:1611:C:N4	2.43	0.52
34:YA:2054:A:C2	34:YA:2616:C:O2	2.62	0.52
34:YA:2185:C:H2'	34:YA:2186:G:C8	2.45	0.52
34:YA:2577:A:H2'	34:YA:2614:A:H62	1.75	0.52
1:QA:675:A:H61	1:QA:715:A:H61	1.57	0.52
1:QA:1251:A:N1	1:QA:1354:C:O2'	2.35	0.52
1:QA:1357:A:C5'	10:QJ:45:ARG:HH12	2.22	0.52
4:QD:122:ARG:NH1	4:QD:122:ARG:O	2.43	0.52
28:R4:7:PRO:CG	39:RG:61:ALA:HB1	2.30	0.52
34:RA:1567:A:C8	36:RD:84:TYR:CE2	2.98	0.52
34:RA:2306:C:H2'	34:RA:2307:G:N2	2.25	0.52
35:RB:104:A:OP1	54:RZ:72:ARG:NH1	2.43	0.52
1:XA:8:A:H62	4:XD:208:SER:C	2.13	0.52
1:XA:191:G:O2'	20:XT:103:GLY:N	2.43	0.52
1:XA:1440(K):C:HO2'	1:XA:1440(L):G:N2	2.07	0.52
1:XA:1440(N):G:P	20:XT:35:THR:HG21	2.46	0.52
2:XB:43:ASP:O	2:XB:47:THR:OG1	2.23	0.52
4:XD:187:ARG:NH1	4:XD:188:LEU:O	2.43	0.52
32:Y8:49:VAL:HG23	32:Y8:53:PRO:HD3	1.91	0.52
34:YA:10:G:N2	34:YA:2629:A:C2	2.78	0.52
34:YA:1801:G:OP1	34:YA:1801:G:N2	2.32	0.52
1:QA:520:A:C2'	12:QL:73:GLU:OE2	2.57	0.52
1:QA:959:A:H62	19:QS:79:THR:CG2	2.23	0.52
1:QA:986:A:O2'	19:QS:55:LYS:HA	2.10	0.52
1:QA:1160:G:H5'	2:QB:132:LYS:NZ	2.25	0.52
1:QA:1203:C:H5'	14:QN:3:ARG:HH11	1.74	0.52
2:QB:61:LEU:HD21	2:QB:68:ILE:HD11	1.91	0.52
24:R0:66:VAL:O	24:R0:81:VAL:HA	2.10	0.52
35:RB:116:G:H4'	47:RS:54:LEU:CD2	2.39	0.52
44:RP:84:ASN:ND2	44:RP:117:GLU:OE2	2.43	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:718:G:C1'	11:XK:117:ASN:HB2	2.40	0.52
1:XA:1033:G:H2'	1:XA:1034:G:C8	2.44	0.52
1:XA:1123:A:H1'	10:XJ:38:ILE:HG22	1.92	0.52
1:XA:1244:C:H2'	1:XA:1245:A:C8	2.45	0.52
34:YA:197:A:C8	34:YA:2430:A:H2	2.27	0.52
34:YA:389:G:H22	44:YP:72:PRO:HD3	1.75	0.52
41:YI:20:ASP:OD2	41:YI:20:ASP:N	2.43	0.52
45:YQ:21:THR:OG1	45:YQ:22:LYS:N	2.42	0.52
1:QA:164:U:H2'	1:QA:165:C:C6	2.45	0.51
1:QA:186(C):C:H1'	20:QT:89:ARG:HH22	1.75	0.51
1:QA:1240:U:H2'	7:QG:38:LEU:CD1	2.16	0.51
1:QA:1304:G:H1'	1:QA:1334:G:N1	2.25	0.51
33:R9:27:CYS:SG	33:R9:28:GLU:N	2.83	0.51
34:RA:990:A:H1'	34:RA:1156:A:N3	2.25	0.51
34:RA:2102:U:H2'	34:RA:2103:C:H6	1.75	0.51
34:RA:2836:U:H2'	34:RA:2837:G:C8	2.44	0.51
42:RN:35:ARG:HG3	42:RN:37:LYS:HG2	1.91	0.51
1:XA:322:C:H5''	20:XT:23:ARG:HE	1.75	0.51
1:XA:974:A:N9	14:XN:31:ARG:CD	2.63	0.51
1:XA:1382:C:O5'	1:XA:1382:C:H6	1.93	0.51
8:XH:121:ASP:OD1	8:XH:121:ASP:N	2.42	0.51
34:YA:199:A:C5	34:YA:2434:A:N1	2.77	0.51
34:YA:564:C:N4	34:YA:573:G:OP1	2.37	0.51
34:YA:1849:G:H2'	34:YA:1850:G:H8	1.75	0.51
34:YA:2070:G:H2'	34:YA:2071:A:C8	2.44	0.51
34:YA:2417:C:OP1	44:YP:64:LYS:NZ	2.43	0.51
1:QA:749:C:H2'	1:QA:750:G:H8	1.74	0.51
1:QA:960:U:O4	19:QS:78:ARG:CB	2.52	0.51
1:QA:974:A:C4'	14:QN:31:ARG:CB	2.87	0.51
1:QA:1315:U:H2'	1:QA:1316:G:H8	1.75	0.51
1:QA:1360:A:H4'	14:QN:17:LYS:HZ2	1.74	0.51
4:QD:15:GLU:OE2	4:QD:66:ARG:NH2	2.42	0.51
34:RA:1768:U:H2'	34:RA:1769:G:C8	2.45	0.51
34:RA:1992:G:N2	34:RA:1996:C:O2'	2.43	0.51
34:RA:2086:U:H2'	34:RA:2087:G:C8	2.45	0.51
43:RO:23:ARG:NH2	43:RO:28:SER:O	2.43	0.51
1:XA:186(B):C:O2'	20:XT:104:LEU:HD11	2.09	0.51
1:XA:619:U:O4'	4:XD:131:ARG:NH2	2.42	0.51
1:XA:1252:A:C2	1:XA:1355:G:H1'	2.41	0.51
1:XA:1294:G:H2'	1:XA:1295:G:H8	1.76	0.51
29:Y5:48:GLU:CD	51:YW:37:ARG:HH12	2.09	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:414:C:O3'	34:YA:1878:G:N2	2.44	0.51
34:YA:1418:G:H2'	34:YA:1579:A:N6	2.25	0.51
34:YA:1999:C:H2'	34:YA:2000:G:C8	2.45	0.51
34:YA:2044:C:N4	34:YA:2625:G:C2	2.77	0.51
34:YA:2847:U:OP1	48:YT:98:LYS:NZ	2.41	0.51
36:YD:8:PRO:HB3	36:YD:14:ARG:HG2	1.92	0.51
1:QA:617:G:H21	16:QP:14:ASN:ND2	2.08	0.51
1:QA:741:G:O5'	15:QO:39:LEU:HD12	2.07	0.51
3:QC:79:ARG:CD	11:XK:104:GLN:HG3	2.40	0.51
5:QE:147:ASP:HA	5:QE:150:ARG:HD2	1.92	0.51
7:QG:69:VAL:HG13	7:QG:100:ALA:HB1	1.93	0.51
32:R8:6:THR:HG23	34:RA:242:G:H3'	1.93	0.51
34:RA:1674:G:H1'	34:RA:1676:A:N6	2.26	0.51
37:RE:14:ILE:HB	48:RT:14:TYR:HE2	1.76	0.51
41:RI:83:ALA:C	41:RI:89:TYR:CD2	2.83	0.51
53:RY:76:CYS:HB2	53:RY:99:CYS:SG	2.51	0.51
1:XA:406:G:H2'	1:XA:407:G:C8	2.45	0.51
1:XA:1129:C:H1'	1:XA:1132:C:C5	2.45	0.51
1:XA:1259:C:O2'	1:XA:1283:G:N2	2.42	0.51
10:XJ:64:GLU:O	14:XN:56:VAL:HG22	2.11	0.51
12:XL:10:LEU:HD23	17:XQ:32:TYR:CZ	2.44	0.51
16:XP:4:ILE:HB	16:XP:66:PRO:HB3	1.92	0.51
30:Y6:34:LEU:N	30:Y6:51:GLU:OE1	2.44	0.51
34:YA:1297:C:H2'	34:YA:1298:C:H6	1.74	0.51
34:YA:1971:A:C4	36:YD:241:PRO:HB3	2.45	0.51
34:YA:2049:G:N2	34:YA:2620:C:N1	2.53	0.51
34:YA:2641:G:H5''	42:YN:76:SER:HB3	1.93	0.51
35:YB:104:A:OP1	54:YZ:72:ARG:NH1	2.42	0.51
37:YE:78:LEU:HG	37:YE:79:ARG:HD2	1.91	0.51
45:YQ:67:ARG:O	45:YQ:101:ARG:NH2	2.44	0.51
48:YT:19:LEU:HD22	48:YT:86:ILE:HD12	1.92	0.51
1:QA:45:U:H3	1:QA:396:G:H1	1.58	0.51
1:QA:269:C:H2'	1:QA:270:A:C8	2.45	0.51
1:QA:981:U:H5'	14:QN:6:LEU:HD23	1.92	0.51
1:QA:1137:C:H4'	1:QA:1138:G:C2	2.46	0.51
1:QA:1309:G:H2'	1:QA:1310:G:C8	2.45	0.51
2:QB:84:GLU:OE2	2:QB:87:ARG:NH2	2.43	0.51
18:QR:30:ASP:HB3	18:QR:33:ASP:HB2	1.92	0.51
34:RA:220:G:H22	34:RA:427:U:H2'	1.75	0.51
34:RA:242:G:N2	34:RA:254:G:H2'	2.25	0.51
34:RA:307:G:H21	34:RA:330:A:N6	2.09	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:633:A:H1'	34:RA:2403:C:H4'	1.91	0.51
34:RA:679:C:H2'	34:RA:680:G:C8	2.46	0.51
34:RA:1546:C:H5'	34:RA:1547:C:H5'	1.91	0.51
34:RA:1660:C:H2'	34:RA:1661:G:H8	1.75	0.51
41:RI:86:THR:HA	41:RI:123:LEU:CB	2.40	0.51
1:XA:375:U:H6	1:XA:375:U:C5'	2.23	0.51
34:YA:414:C:H2'	34:YA:415:A:C8	2.45	0.51
34:YA:882:G:H2'	34:YA:883:G:C8	2.45	0.51
34:YA:1252:G:H21	49:YU:33:ARG:HD3	1.76	0.51
34:YA:2443:C:H2'	34:YA:2444:G:H8	1.76	0.51
34:YA:2690:C:N4	34:YA:2713:A:N3	2.58	0.51
38:YF:198:ALA:HA	38:YF:201:VAL:HG12	1.91	0.51
1:QA:608:A:O2'	16:QP:9:PHE:CE1	2.62	0.51
1:QA:986:A:H2'	1:QA:987:G:C8	2.46	0.51
25:R1:10:LYS:NZ	25:R1:65:SER:OG	2.41	0.51
28:R4:26:SER:OG	28:R4:27:THR:N	2.43	0.51
34:RA:414:C:H2'	34:RA:415:A:H8	1.75	0.51
34:RA:2575:C:H5'	37:RE:144:ARG:HG3	1.91	0.51
34:RA:2647:U:H2'	34:RA:2648:C:H6	1.76	0.51
40:RH:103:LEU:CG	40:RH:123:PHE:CE1	2.94	0.51
41:RI:86:THR:HA	41:RI:123:LEU:HB2	1.92	0.51
47:RS:26:LEU:O	47:RS:88:ASP:HB3	2.10	0.51
1:XA:123:C:H2'	1:XA:124:G:H8	1.76	0.51
1:XA:784:C:H2'	1:XA:785:G:C8	2.46	0.51
1:XA:1279:A:HO2'	1:XA:1281:U:H5	1.59	0.51
3:XC:58:GLU:HB2	3:XC:65:ALA:HB3	1.92	0.51
34:YA:544:C:H3'	34:YA:545:G:H8	1.75	0.51
34:YA:863:A:H2'	34:YA:864:G:H8	1.75	0.51
34:YA:2105:C:H2'	34:YA:2106:G:C8	2.46	0.51
1:QA:637:G:H2'	1:QA:638:G:H8	1.76	0.51
1:QA:908:A:H2'	1:QA:909:A:C8	2.45	0.51
1:QA:1142:G:C2	1:QA:1143:G:H1'	2.46	0.51
1:QA:1203:C:OP2	14:QN:3:ARG:HD3	2.11	0.51
34:RA:1057:A:H62	34:RA:1087:G:P	2.32	0.51
40:RH:103:LEU:HG	40:RH:123:PHE:CE1	2.46	0.51
41:RI:94:ALA:HA	41:RI:97:ILE:HD12	1.93	0.51
1:XA:1512:U:H2'	1:XA:1513:A:H8	1.76	0.51
5:XE:102:ALA:HB1	5:XE:106:PRO:HG2	1.93	0.51
7:XG:58:PRO:HA	7:XG:61:VAL:HG12	1.92	0.51
34:YA:992:C:H2'	34:YA:993:G:H8	1.76	0.51
34:YA:1576:U:H2'	34:YA:1577:C:H6	1.76	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:2090:G:C6	34:YA:2230:G:N1	2.78	0.51
38:YF:63:LYS:NZ	38:YF:75:HIS:O	2.33	0.51
53:YY:76:CYS:SG	53:YY:79:CYS:SG	3.08	0.51
1:QA:770:C:H2'	1:QA:771:G:C8	2.46	0.51
1:QA:1112:C:O2	3:QC:178:LEU:HB2	2.10	0.51
1:QA:1292:U:H2'	1:QA:1293:G:C8	2.46	0.51
4:QD:57:ARG:HH22	5:QE:107:ARG:CG	2.23	0.51
5:QE:10:MET:HA	5:QE:32:VAL:HG12	1.93	0.51
30:R6:6:ARG:NH1	30:R6:24:GLU:OE2	2.43	0.51
34:RA:191:A:H2'	34:RA:192:C:C6	2.45	0.51
34:RA:1914:C:H2'	34:RA:1915:U:O4'	2.10	0.51
45:RQ:21:THR:OG1	45:RQ:22:LYS:N	2.43	0.51
53:RY:67:LEU:HD22	53:RY:71:LYS:HD2	1.91	0.51
1:XA:554:C:H2'	1:XA:555:C:C6	2.46	0.51
1:XA:636:U:H3'	1:XA:637:G:C8	2.45	0.51
1:XA:838(B):U:H4'	1:XA:838(C):C:C5	2.45	0.51
1:XA:977:A:O3'	1:XA:980:C:N4	2.41	0.51
1:XA:1305:G:N2	1:XA:1331:G:C8	2.79	0.51
1:XA:1422:G:H4'	43:YO:48:PRO:HB3	1.91	0.51
11:XK:109:VAL:HA	18:XR:86:VAL:HA	1.93	0.51
34:YA:36:G:H4'	34:YA:451:C:C2	2.45	0.51
34:YA:197:A:C5	34:YA:2430:A:N3	2.79	0.51
34:YA:2046:G:O6	34:YA:2623:G:O6	2.29	0.51
36:YD:108:PRO:HB3	36:YD:143:HIS:CE1	2.46	0.51
52:YX:25:LYS:HA	52:YX:81:VAL:O	2.11	0.51
1:QA:186(A):C:H1'	20:QT:81:LYS:HE3	1.92	0.51
1:QA:770:C:H2'	1:QA:771:G:H8	1.76	0.51
1:QA:1391:U:H2'	1:QA:1392:G:C8	2.46	0.51
28:R4:7:PRO:HG3	39:RG:61:ALA:CB	2.27	0.51
34:RA:36:G:H4'	34:RA:451:C:C2	2.45	0.51
34:RA:414:C:H2'	34:RA:415:A:C8	2.46	0.51
34:RA:1019:U:H2'	34:RA:1020:A:C8	2.43	0.51
34:RA:2289:G:H1'	34:RA:2346:A:H2	1.75	0.51
37:RE:110:GLY:HA2	37:RE:161:GLY:HA3	1.93	0.51
47:RS:6:ALA:HA	47:RS:9:ARG:HG2	1.92	0.51
48:RT:3:ARG:HG3	48:RT:6:LEU:HB2	1.92	0.51
50:RV:8:GLY:O	50:RV:10:LYS:NZ	2.42	0.51
1:XA:398:C:H2'	1:XA:399:G:C8	2.45	0.51
1:XA:1303:C:O2	1:XA:1303:C:H2'	2.11	0.51
1:XA:1305:G:N7	1:XA:1331:G:C6	2.75	0.51
1:XA:1313:U:H5	19:XS:6:LYS:HZ1	1.57	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1347:G:H22	1:XA:1373:G:H2'	1.76	0.51
8:XH:38:ILE:HD12	8:XH:41:ARG:HH21	1.76	0.51
34:YA:1454:U:H5	46:YR:73:VAL:CG1	2.24	0.51
34:YA:2071:A:H2	34:YA:2441:C:C2	2.28	0.51
34:YA:2085:C:N3	34:YA:2235:G:N2	2.58	0.51
34:YA:2099:U:O4	34:YA:2190:G:C6	2.64	0.51
38:YF:34:TRP:CD2	44:YP:8:PRO:HB3	2.46	0.51
42:YN:97:ARG:HA	42:YN:100:GLU:HB2	1.92	0.51
1:QA:65:U:H2'	1:QA:381:C:C5	2.45	0.51
1:QA:126:G:O2'	1:QA:634:C:O2'	2.21	0.51
1:QA:1131:G:H2'	1:QA:1132:C:C6	2.46	0.51
3:QC:34:LEU:O	3:QC:38:ARG:NE	2.43	0.51
8:QH:32:LYS:HA	8:QH:35:ILE:HD12	1.92	0.51
12:QL:10:LEU:HB3	17:QQ:32:TYR:OH	2.10	0.51
12:QL:46:LYS:HD2	12:QL:47:LYS:HB2	1.93	0.51
34:RA:1794:U:H2'	34:RA:1795:C:C6	2.45	0.51
1:XA:186(K):G:O6	1:XA:264:U:H5''	2.11	0.51
1:XA:401:C:H1'	1:XA:622:A:H1'	1.93	0.51
1:XA:435:C:H2'	1:XA:436:C:H6	1.76	0.51
1:XA:757:U:O2'	1:XA:879:C:O2	2.27	0.51
1:XA:1190:G:C4'	3:XC:176:HIS:CE1	2.83	0.51
1:XA:1216:G:H5''	14:XN:5:ALA:HB2	1.92	0.51
1:XA:1230:C:C4	13:XM:102:ARG:NH1	2.77	0.51
6:XF:9:VAL:HB	6:XF:87:ARG:HB2	1.92	0.51
29:Y5:3:LYS:HB2	34:YA:2577:A:C1'	2.40	0.51
34:YA:1416:G:H2'	34:YA:1417:C:C6	2.46	0.51
34:YA:1820:U:C2	36:YD:202:LYS:HB3	2.46	0.51
1:QA:34:C:H2'	1:QA:35:G:C8	2.45	0.51
1:QA:231:G:H2'	1:QA:232:G:H8	1.75	0.51
1:QA:1131:G:H1	1:QA:1143:G:N2	2.01	0.51
1:QA:1203:C:C5'	14:QN:3:ARG:HD2	2.41	0.51
3:QC:150:LYS:HB3	3:QC:201:TYR:HB2	1.93	0.51
32:R8:21:LYS:HE3	34:RA:651:G:OP2	2.11	0.51
42:RN:39:ARG:NH2	42:RN:41:ASP:OD1	2.44	0.51
44:RP:57:THR:OG1	44:RP:58:THR:N	2.44	0.51
1:XA:148:G:H2'	1:XA:149:A:C8	2.45	0.51
1:XA:878:G:H2'	1:XA:879:C:C6	2.46	0.51
10:XJ:26:ALA:O	10:XJ:84:GLN:NE2	2.43	0.51
34:YA:140:A:H1'	34:YA:1409:C:H5'	1.93	0.51
34:YA:291:C:H2'	34:YA:292:C:H6	1.76	0.51
34:YA:389:G:H1	44:YP:71:VAL:CG1	2.16	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:659:C:H2'	34:YA:660:G:H8	1.76	0.51
34:YA:1700:A:H3'	34:YA:1701:A:H8	1.76	0.51
34:YA:2250:G:OP2	34:YA:2250:G:N2	2.35	0.51
38:YF:154:VAL:HG22	38:YF:191:ARG:HB2	1.93	0.51
41:YI:1:MET:HG2	41:YI:23:PRO:HB3	1.92	0.51
1:QA:447:G:H2'	1:QA:485:G:N2	2.26	0.50
1:QA:617:G:H4'	16:QP:44:THR:OG1	2.11	0.50
1:QA:702:A:H3'	1:QA:703:G:H8	1.75	0.50
1:QA:815:A:H1'	1:QA:1527:C:H1'	1.93	0.50
1:QA:920:U:H2'	1:QA:921:U:C6	2.46	0.50
1:QA:1038:C:H2'	1:QA:1039:C:H6	1.76	0.50
1:QA:1318:A:C4	19:QS:37:ARG:CZ	2.93	0.50
2:QB:178:ARG:CD	8:QH:71:GLY:O	2.58	0.50
34:RA:2853:C:H2'	34:RA:2854:G:C8	2.46	0.50
36:RD:147:LEU:HD12	36:RD:148:GLU:HG3	1.93	0.50
36:RD:175:LEU:O	36:RD:182:LEU:HA	2.12	0.50
39:RG:52:ILE:HG22	39:RG:55:LYS:HD2	1.92	0.50
47:RS:35:ILE:HD11	47:RS:97:ARG:HD2	1.92	0.50
1:XA:140:A:H2'	1:XA:141:A:C8	2.45	0.50
1:XA:708:C:P	11:XK:85:ARG:HH12	2.28	0.50
1:XA:1068:G:N2	1:XA:1108:G:H1'	2.25	0.50
1:XA:1505:G:H1'	23:XX:15:A:C2	2.46	0.50
9:XI:22:GLY:N	9:XI:58:HIS:O	2.37	0.50
34:YA:86:C:H2'	34:YA:87:C:C6	2.46	0.50
34:YA:137(B):G:H2'	34:YA:139:G:N7	2.26	0.50
34:YA:824:A:O2'	34:YA:2358:G:O6	2.17	0.50
34:YA:1316:U:H2'	34:YA:1317:A:C8	2.46	0.50
34:YA:1999:C:H5''	34:YA:2723:C:O2'	2.11	0.50
34:YA:2071:A:H2'	34:YA:2072:G:H8	1.76	0.50
34:YA:2312:U:H2'	34:YA:2313:C:H6	1.76	0.50
34:YA:2532:G:O2'	34:YA:2657:A:N1	2.36	0.50
1:QA:112:G:H22	1:QA:315:A:H2	1.59	0.50
1:QA:266:G:O2'	1:QA:268:C:OP2	2.24	0.50
2:QB:208:ILE:O	2:QB:212:GLN:HB2	2.12	0.50
20:QT:71:THR:OG1	20:QT:72:LEU:N	2.43	0.50
32:R8:8:LYS:HB3	32:R8:12:LYS:HE3	1.94	0.50
32:R8:30:ARG:O	32:R8:30:ARG:HG2	2.12	0.50
34:RA:65:C:H2'	34:RA:66:C:H6	1.76	0.50
34:RA:380:U:H2'	34:RA:381:G:C8	2.47	0.50
34:RA:639:U:H2'	34:RA:640:C:C6	2.46	0.50
34:RA:1150:C:H2'	34:RA:1151:G:C8	2.46	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:24:U:H2'	1:XA:25:C:H6	1.76	0.50
1:XA:1533:C:C4	23:XX:13:A:N1	2.79	0.50
2:XB:167:PRO:O	2:XB:171:ALA:HB2	2.11	0.50
4:XD:19:LEU:HB3	4:XD:21:LEU:HD23	1.92	0.50
4:XD:107:ARG:HB3	4:XD:174:LEU:HD11	1.93	0.50
34:YA:659:C:H2'	34:YA:660:G:C8	2.47	0.50
34:YA:1113:U:H5'	40:YH:2:SER:HB3	1.93	0.50
34:YA:1479:G:N7	34:YA:1510:A:N6	2.59	0.50
34:YA:1582:C:H2'	34:YA:1583:A:C8	2.46	0.50
34:YA:2056:G:N7	34:YA:2577:A:C6	2.79	0.50
34:YA:2691:C:H2'	34:YA:2692:C:C6	2.46	0.50
34:YA:2747:G:O6	34:YA:2755:C:H5''	2.12	0.50
1:QA:877:C:H2'	1:QA:878:G:C8	2.47	0.50
1:QA:979:C:N4	14:QN:19:ARG:CB	2.42	0.50
1:QA:1145:C:O2'	1:QA:1146:A:N7	2.43	0.50
7:QG:111:ARG:NH1	7:QG:113:GLU:OE1	2.43	0.50
11:QK:108:ILE:CG2	18:QR:88:LYS:OXT	2.60	0.50
12:QL:39:VAL:HG12	12:QL:57:LYS:HG2	1.93	0.50
20:QT:67:ALA:O	20:QT:73:HIS:ND1	2.39	0.50
34:RA:597:U:H2'	34:RA:598:G:C8	2.45	0.50
34:RA:686:G:N2	34:RA:788:A:H61	2.10	0.50
34:RA:997:G:H3'	49:RU:58:ARG:HH12	1.76	0.50
34:RA:1116:C:H2'	34:RA:1117:G:C8	2.46	0.50
39:RG:121:ASN:O	39:RG:131:TYR:OH	2.26	0.50
1:XA:702:A:N1	34:YA:1848:A:C5	2.79	0.50
1:XA:1539:C:C2	23:XX:7:G:N1	2.79	0.50
7:XG:111:ARG:HD3	7:XG:112:PRO:HD2	1.93	0.50
34:YA:389:G:N2	44:YP:71:VAL:CG1	2.69	0.50
34:YA:589:C:H2'	34:YA:590:A:C8	2.46	0.50
34:YA:1093:G:H21	34:YA:1098:A:H62	1.59	0.50
34:YA:1827:C:H5'	34:YA:1971:A:H4'	1.94	0.50
1:QA:10:A:H2'	1:QA:11:G:C8	2.46	0.50
1:QA:107:G:H3'	1:QA:108:G:N2	2.24	0.50
1:QA:757:U:H1'	1:QA:879:C:H1'	1.93	0.50
1:QA:1302:U:P	13:QM:27:LYS:HE2	2.51	0.50
4:QD:205:GLU:HB3	5:QE:107:ARG:HH12	1.77	0.50
32:R8:64:TYR:HB3	34:RA:625:G:P	2.51	0.50
34:RA:1796:U:H2'	34:RA:1797:C:C6	2.47	0.50
34:RA:2144:U:H2'	34:RA:2146:C:C5	2.47	0.50
38:RF:100:THR:OG1	38:RF:100:THR:O	2.30	0.50
49:RU:90:VAL:HG13	50:RV:39:LEU:HD22	1.92	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1145:C:O2'	1:XA:1146:A:N7	2.43	0.50
6:XF:91:VAL:CG2	18:XR:34:TYR:CZ	2.82	0.50
18:XR:32:ARG:HA	18:XR:69:THR:HG21	1.92	0.50
28:Y4:16:CYS:HB3	28:Y4:33:VAL:HB	1.93	0.50
34:YA:390:A:H1'	34:YA:391:G:C8	2.47	0.50
34:YA:1853:A:N1	34:YA:2087:G:H1'	2.27	0.50
34:YA:2807:G:H1	34:YA:2892:A:H62	1.60	0.50
40:YH:85:LYS:HB3	40:YH:133:VAL:HG13	1.92	0.50
1:QA:6:G:N3	5:QE:119:LEU:HD11	2.27	0.50
1:QA:37:U:H2'	1:QA:38:G:H8	1.76	0.50
1:QA:760:G:H3'	1:QA:761:G:H8	1.76	0.50
1:QA:1222:G:O3'	19:QS:78:ARG:CZ	2.60	0.50
2:QB:111:ARG:HH11	2:QB:114:ARG:HH12	1.60	0.50
20:QT:66:ALA:O	20:QT:71:THR:OG1	2.28	0.50
34:RA:301:G:OP2	53:RY:84:ARG:NH1	2.35	0.50
34:RA:373:U:H2'	34:RA:374:A:H8	1.77	0.50
42:RN:112:LEU:O	42:RN:116:LEU:HB2	2.10	0.50
53:RY:19:LYS:NZ	53:RY:20:TYR:CE2	2.71	0.50
1:XA:767:A:H2'	1:XA:768:A:C8	2.46	0.50
1:XA:981:U:O5'	1:XA:981:U:H6	1.94	0.50
1:XA:986:A:C1'	19:XS:52:TYR:OH	2.58	0.50
1:XA:1305:G:N2	1:XA:1331:G:H3'	2.26	0.50
1:XA:1305:G:C1'	1:XA:1332:A:H62	2.24	0.50
1:XA:1306:A:OP2	1:XA:1331:G:N2	2.33	0.50
3:XC:157:ILE:HD12	3:XC:164:ARG:HG3	1.93	0.50
10:XJ:42:THR:HG22	10:XJ:44:VAL:HG22	1.92	0.50
19:XS:12:ASP:HB2	19:XS:37:ARG:HE	1.77	0.50
22:XV:19:G:C5	34:YA:2111:C:H4'	2.46	0.50
29:Y5:9:LYS:HE2	34:YA:2019:A:N7	2.26	0.50
30:Y6:21:TYR:OH	32:Y8:36:LYS:O	2.28	0.50
31:Y7:37:LYS:CE	34:YA:458:G:C5	2.95	0.50
34:YA:1295:C:H2'	34:YA:1296:G:H8	1.76	0.50
34:YA:1570:A:H2'	34:YA:1571:A:C8	2.47	0.50
49:YU:43:GLY:HA3	50:YV:73:SER:HB3	1.93	0.50
53:YY:28:LYS:NZ	53:YY:64:GLU:OE2	2.35	0.50
1:QA:22:G:H4'	1:QA:885:G:C8	2.47	0.50
1:QA:1253:G:C4'	10:QJ:44:VAL:O	2.58	0.50
1:QA:1323:G:N2	1:QA:1361:G:O2'	2.39	0.50
3:QC:57:ILE:HG22	3:QC:66:VAL:HG22	1.93	0.50
10:QJ:47:PHE:CE1	14:QN:36:PHE:CG	2.99	0.50
34:RA:990:A:N6	34:RA:1186:G:H1'	2.26	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:1205:U:C4	38:RF:171:PRO:HA	2.46	0.50
34:RA:2148:G:H2'	34:RA:2149:G:C8	2.47	0.50
34:RA:2159:G:H2'	34:RA:2160:G:C8	2.46	0.50
34:RA:2314:C:H2'	34:RA:2315:G:C8	2.46	0.50
34:RA:2317:C:H3'	34:RA:2318:G:H21	1.75	0.50
1:XA:656:C:O2'	15:XO:28:GLN:NE2	2.45	0.50
1:XA:728:A:C5	15:XO:54:ARG:CD	2.95	0.50
1:XA:1347:G:C8	9:XI:108:VAL:C	2.85	0.50
1:XA:1347:G:N2	1:XA:1373:G:H2'	2.26	0.50
20:XT:74:LYS:O	20:XT:76:ALA:N	2.44	0.50
38:YF:10:PRO:HB3	38:YF:17:ARG:HH21	1.76	0.50
40:YH:113:VAL:HG11	40:YH:151:ILE:HD12	1.94	0.50
52:YX:53:LYS:HG2	52:YX:82:GLN:HB3	1.93	0.50
1:QA:17:U:H2'	1:QA:18:C:C6	2.46	0.50
1:QA:1368:G:OP1	10:QJ:62:HIS:CD2	2.64	0.50
1:QA:1510:U:H3	1:QA:1525:G:H1	1.59	0.50
32:R8:39:LYS:NZ	34:RA:2365:G:C6	2.77	0.50
34:RA:276:A:O2'	34:RA:278:A:OP2	2.28	0.50
34:RA:2025:C:H2'	34:RA:2026:C:C6	2.47	0.50
36:RD:69:ARG:NH1	36:RD:128:GLY:O	2.42	0.50
43:RO:71:ARG:NE	43:RO:105:GLU:OE2	2.36	0.50
1:XA:45:U:H3	1:XA:396:G:H1	1.59	0.50
1:XA:754:C:C6	15:XO:69:TYR:CE2	2.95	0.50
3:XC:21:ARG:HH11	10:XJ:15:THR:HG21	1.75	0.50
10:XJ:40:LEU:CB	10:XJ:41:PRO:CD	2.68	0.50
10:XJ:62:HIS:HD2	14:XN:59:ALA:CB	2.24	0.50
13:XM:82:MET:O	13:XM:93:ARG:NH2	2.43	0.50
20:XT:75:ASN:N	20:XT:75:ASN:OD1	2.44	0.50
33:Y9:27:CYS:HB3	33:Y9:32:HIS:HB2	1.94	0.50
34:YA:1041:C:H2'	34:YA:1042:G:H8	1.77	0.50
34:YA:1359:A:N6	34:YA:1372:U:H3	2.10	0.50
34:YA:1604:C:H2'	34:YA:1605:C:H6	1.76	0.50
34:YA:2125:G:N1	34:YA:2172:U:OP1	2.29	0.50
37:YE:101:ARG:NE	37:YE:171:GLU:OE2	2.41	0.50
1:QA:1170:A:H5'	2:QB:140:HIS:HE1	1.73	0.50
1:QA:1253:G:OP2	10:QJ:44:VAL:CG2	2.58	0.50
1:QA:1315:U:H2'	1:QA:1316:G:C8	2.47	0.50
34:RA:288:C:H2'	34:RA:289:A:H8	1.76	0.50
34:RA:2030:A:H4'	34:RA:2031:A:C8	2.46	0.50
34:RA:2153:G:H2'	34:RA:2154:G:C8	2.46	0.50
36:RD:126:GLN:O	36:RD:129:ASN:ND2	2.41	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:895:G:O2'	17:XQ:100:LYS:HD2	2.11	0.50
1:XA:979:C:H5''	1:XA:1221:G:C8	2.47	0.50
2:XB:118:LEU:HD23	2:XB:142:LEU:HB2	1.94	0.50
4:XD:88:VAL:HG13	5:XE:97:GLY:HA3	1.94	0.50
34:YA:959:A:N3	34:YA:2457:U:O2'	2.44	0.50
34:YA:1150:C:H2'	34:YA:1151:G:C8	2.47	0.50
34:YA:1176:G:H3'	34:YA:1177:A:C8	2.47	0.50
34:YA:1918:A:O2'	34:YA:1920:C:N4	2.44	0.50
34:YA:2563:U:H1'	34:YA:2566:A:N6	2.27	0.50
51:YW:86:LEU:HD22	51:YW:96:ILE:HD11	1.93	0.50
1:QA:982:U:C5'	14:QN:6:LEU:CD2	2.89	0.50
1:QA:1491:G:C5'	12:QL:46:LYS:HG2	2.41	0.50
2:QB:146:GLN:HG3	2:QB:153:ARG:HH22	1.77	0.50
12:QL:104:VAL:O	12:QL:105:TYR:CG	2.57	0.50
34:RA:828:U:O2'	34:RA:831:G:O6	2.26	0.50
34:RA:1057:A:N6	34:RA:1087:G:OP1	2.45	0.50
34:RA:1408:C:H2'	34:RA:1409:C:C6	2.47	0.50
34:RA:1638:C:O2	34:RA:2698:U:O2'	2.30	0.50
34:RA:1771:C:H2'	34:RA:1772:G:C8	2.47	0.50
34:RA:2695:C:H2'	34:RA:2696:U:C6	2.47	0.50
42:RN:9:VAL:HG11	42:RN:39:ARG:HH12	1.77	0.50
1:XA:879:C:H2'	1:XA:880:C:C6	2.46	0.50
1:XA:1040:U:H2'	1:XA:1041:A:C8	2.47	0.50
1:XA:1358:U:H5'	14:XN:35:ARG:N	2.27	0.50
8:XH:21:LYS:O	8:XH:65:TYR:OH	2.28	0.50
30:Y6:23:THR:OG1	30:Y6:24:GLU:N	2.42	0.50
34:YA:69:C:H2'	34:YA:70:G:H8	1.77	0.50
34:YA:2097:C:C2	34:YA:2193:G:C2	3.00	0.50
34:YA:2100:G:C6	34:YA:2190:G:C4	3.00	0.50
34:YA:2124:G:C2	34:YA:2125:G:H1'	2.46	0.50
34:YA:2275:C:O2	45:YQ:83:MET:SD	2.69	0.50
34:YA:2698:U:H2'	34:YA:2699:C:C6	2.47	0.50
43:YO:63:VAL:HG12	43:YO:106:LEU:HD21	1.93	0.50
1:QA:625:G:H4'	16:QP:16:HIS:HB3	1.93	0.49
1:QA:1094:G:H4'	1:QA:1095:U:H5	1.76	0.49
1:QA:1100:C:H5	2:QB:96:ARG:HH21	1.46	0.49
4:QD:57:ARG:NH2	5:QE:107:ARG:NH1	2.59	0.49
24:R0:23:VAL:HG21	34:RA:857:C:H4'	1.93	0.49
24:R0:72:ARG:HE	24:R0:75:LEU:HD12	1.76	0.49
34:RA:659:C:H2'	34:RA:660:G:H8	1.77	0.49
34:RA:1494:A:H2	34:RA:1579:A:H1'	1.76	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:1818:U:H5	36:RD:157:ARG:HH21	1.60	0.49
34:RA:2273:A:H2'	34:RA:2274:A:C8	2.46	0.49
1:XA:413:G:H1'	1:XA:428:G:N2	2.27	0.49
1:XA:948:C:OP2	13:XM:106:ASN:CB	2.60	0.49
1:XA:1304:G:H2'	1:XA:1333:A:H61	1.78	0.49
5:XE:81:GLU:HG2	5:XE:90:VAL:HG23	1.94	0.49
34:YA:922:U:H2'	34:YA:923:C:C6	2.47	0.49
34:YA:1127:A:N1	34:YA:2518:A:N6	2.60	0.49
34:YA:1870:C:H2'	34:YA:1871:A:O4'	2.12	0.49
1:QA:62:U:H2'	1:QA:63:C:C6	2.47	0.49
1:QA:248:C:H2'	1:QA:249:U:H6	1.77	0.49
1:QA:584:G:H2'	1:QA:585:G:C8	2.46	0.49
1:QA:980:C:H4'	14:QN:19:ARG:CZ	2.42	0.49
1:QA:1181:G:C2	1:QA:1182:G:H1'	2.47	0.49
1:QA:1221:G:C5'	19:QS:36:ARG:CZ	2.89	0.49
34:RA:345:A:N3	34:RA:346:A:N6	2.61	0.49
34:RA:458:G:H1'	34:RA:459:U:H5	1.77	0.49
1:XA:8:A:N6	4:XD:209:ARG:N	2.57	0.49
1:XA:186(C):C:H2'	1:XA:186(D):G:C8	2.47	0.49
6:XF:89:MET:HE1	18:XR:76:LEU:HB2	1.94	0.49
7:XG:16:LEU:HD21	9:XI:45:ALA:N	2.27	0.49
8:XH:19:VAL:HG23	8:XH:21:LYS:HG3	1.95	0.49
27:Y3:24:LYS:NZ	34:YA:933:A:OP1	2.38	0.49
27:Y3:42:ALA:HB1	34:YA:851:U:O2	2.13	0.49
34:YA:579:G:O2'	34:YA:2019:A:OP1	2.23	0.49
34:YA:2090:G:C2	34:YA:2230:G:C4	2.97	0.49
54:YZ:6:LYS:NZ	54:YZ:43:GLU:OE1	2.39	0.49
1:QA:538:G:C3'	12:QL:115:LYS:NZ	2.60	0.49
1:QA:608:A:C1'	16:QP:32:TYR:HE1	2.22	0.49
1:QA:958:A:C8	19:QS:79:THR:HG21	2.46	0.49
1:QA:1153:C:H2'	1:QA:1154:G:C8	2.47	0.49
5:QE:77:PRO:O	8:QH:105:ARG:CD	2.56	0.49
34:RA:265:A:H2'	34:RA:266:G:H4'	1.93	0.49
34:RA:358:U:H2'	34:RA:359:A:C8	2.48	0.49
34:RA:380:U:H2'	34:RA:381:G:H8	1.76	0.49
34:RA:1476:C:H2'	34:RA:1477:A:C8	2.47	0.49
34:RA:2006:C:H2'	34:RA:2007:C:H6	1.77	0.49
34:RA:2183:C:H2'	34:RA:2184:G:C8	2.48	0.49
34:RA:2735:G:N2	34:RA:2770:G:H1'	2.28	0.49
1:XA:581:G:N2	1:XA:759:A:OP2	2.38	0.49
1:XA:658:G:H1'	15:XO:22:THR:HG21	1.93	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1249:C:N4	1:XA:1288:A:OP2	2.45	0.49
1:XA:1382:C:H2'	1:XA:1383:C:C6	2.47	0.49
1:XA:1481:U:H2'	1:XA:1482:G:C8	2.47	0.49
7:XG:115:ARG:HB2	7:XG:118:VAL:HG12	1.94	0.49
7:XG:140:ASP:OD2	7:XG:143:ARG:NH1	2.45	0.49
13:XM:88:ARG:HD2	13:XM:98:VAL:HB	1.95	0.49
33:Y9:27:CYS:SG	33:Y9:28:GLU:N	2.84	0.49
34:YA:270(P):U:N3	41:YI:52:ARG:NH2	2.60	0.49
34:YA:2261:C:H1'	34:YA:2388:A:H1'	1.94	0.49
35:YB:44:G:O2'	35:YB:47:C:N4	2.44	0.49
1:QA:186(E):C:H2'	1:QA:186(F):C:C6	2.46	0.49
1:QA:676:A:H2	1:QA:714:G:H1	1.60	0.49
1:QA:953:G:N7	13:QM:104:ARG:NH2	2.40	0.49
4:QD:61:LYS:HD3	4:QD:206:PHE:CD2	2.46	0.49
4:QD:205:GLU:HB3	5:QE:107:ARG:NH1	2.27	0.49
5:QE:87:SER:OG	5:QE:125:SER:OG	2.27	0.49
19:QS:10:PHE:HZ	19:QS:15:LEU:HD11	1.76	0.49
25:R1:90:ILE:HA	25:R1:94:LEU:CG	2.42	0.49
34:RA:922:U:H2'	34:RA:923:C:C6	2.47	0.49
34:RA:975:G:N2	34:RA:1156:A:O2'	2.45	0.49
40:RH:89:ILE:HD12	40:RH:131:VAL:HG22	1.94	0.49
45:RQ:36:ALA:HB1	45:RQ:127:ILE:HG21	1.94	0.49
1:XA:412:A:C2	4:XD:35:ARG:HB3	2.47	0.49
1:XA:1233:G:H21	1:XA:1364:U:H3	1.60	0.49
3:XC:19:GLU:O	3:XC:40:ARG:NH2	2.44	0.49
34:YA:1062:G:N2	34:YA:1077:A:N1	2.61	0.49
34:YA:2008:C:H2'	34:YA:2009:G:C8	2.48	0.49
1:QA:5:U:O4	4:QD:87:GLY:N	2.42	0.49
1:QA:192:U:H2'	1:QA:193:C:C6	2.46	0.49
1:QA:422:C:H1'	1:QA:423:G:N1	2.28	0.49
1:QA:891:U:H2'	1:QA:892:A:H8	1.78	0.49
1:QA:893:C:H2'	1:QA:894:G:C8	2.48	0.49
1:QA:922:G:N3	1:QA:1398:A:H2	2.10	0.49
31:R7:35:ARG:NH1	34:RA:54:G:O2'	2.39	0.49
34:RA:2103:C:H2'	34:RA:2104:G:C8	2.48	0.49
50:RV:45:THR:O	50:RV:45:THR:CG2	2.55	0.49
1:XA:25:C:H41	1:XA:559:A:H61	1.58	0.49
1:XA:407:G:C1'	4:XD:119:GLN:HE22	2.26	0.49
1:XA:692:U:H5'	1:XA:797:C:H5'	1.94	0.49
1:XA:891:U:H2'	1:XA:892:A:H8	1.76	0.49
1:XA:974:A:O5'	14:YN:31:ARG:CD	2.60	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:974:A:C5	14:XN:31:ARG:CZ	2.95	0.49
32:Y8:10:ALA:O	32:Y8:14:VAL:HB	2.13	0.49
34:YA:971:C:O2'	34:YA:983:A:N3	2.38	0.49
34:YA:1566:A:C4	36:YD:214:TRP:CE3	3.00	0.49
34:YA:2090:G:C2	34:YA:2230:G:C2	3.00	0.49
37:YE:1:MET:N	37:YE:83:ASP:O	2.36	0.49
1:QA:137:C:C1'	16:QP:63:GLY:HA3	2.32	0.49
1:QA:959:A:N6	19:QS:77:THR:O	2.45	0.49
1:QA:975:A:N6	10:QJ:48:THR:HG1	2.05	0.49
1:QA:1255:G:H1	1:QA:1282:C:H42	1.61	0.49
3:QC:35:GLU:HA	3:QC:38:ARG:HH21	1.77	0.49
4:QD:20:TYR:CZ	6:XF:14:LEU:C	2.85	0.49
10:QJ:47:PHE:CZ	14:QN:37:PHE:CE2	3.00	0.49
25:R1:61:ARG:NH2	34:RA:1364:G:OP2	2.41	0.49
29:R5:9:LYS:NZ	34:RA:2019:A:OP2	2.31	0.49
34:RA:444:C:H2'	34:RA:445:C:C6	2.48	0.49
34:RA:1161:C:H2'	34:RA:1162:G:C8	2.47	0.49
34:RA:2085:C:H4'	36:RD:262:ARG:HH21	1.76	0.49
34:RA:2692:C:H2'	34:RA:2693:A:H8	1.77	0.49
39:RG:135:LEU:O	39:RG:154:GLY:HA3	2.13	0.49
51:RW:22:ASP:OD1	51:RW:25:ARG:NH1	2.45	0.49
1:XA:51:A:C5	1:XA:353:A:C2	3.01	0.49
1:XA:60:A:H2	1:XA:378:G:H1'	1.77	0.49
1:XA:178:C:H2'	1:XA:179:A:H8	1.76	0.49
1:XA:536:C:H2'	1:XA:537:G:C8	2.48	0.49
6:XF:91:VAL:HB	18:XR:34:TYR:OH	2.12	0.49
29:Y5:9:LYS:NZ	34:YA:2019:A:N7	2.59	0.49
34:YA:1028:A:H2'	34:YA:1029:A:C8	2.48	0.49
34:YA:1923:U:H2'	34:YA:1924:C:C6	2.48	0.49
34:YA:2081:C:C2	34:YA:2239:G:N2	2.58	0.49
34:YA:2186:G:H2'	34:YA:2187:G:C8	2.47	0.49
34:YA:2751:G:N7	40:YH:2:SER:OG	2.45	0.49
36:YD:97:TYR:HB2	36:YD:101:GLU:O	2.13	0.49
1:QA:162:A:C5	1:QA:163:C:H1'	2.48	0.49
1:QA:165:C:H2'	1:QA:166:G:C8	2.47	0.49
1:QA:986:A:C2	19:QS:52:TYR:HE1	2.31	0.49
1:QA:1096:C:H2'	1:QA:1097:C:C6	2.47	0.49
1:QA:1306:A:OP2	21:QU:5:ASP:CB	2.60	0.49
1:QA:1313:U:C2'	19:QS:6:LYS:NZ	2.74	0.49
5:QE:151:LEU:HD12	8:QH:79:VAL:HG12	1.94	0.49
33:R9:11:CYS:SG	33:R9:14:CYS:N	2.85	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:896:A:H5''	54:RZ:146:ILE:HG22	1.94	0.49
34:RA:1283:G:H1'	34:RA:1329:U:O2	2.13	0.49
1:XA:657:G:O4'	15:XO:28:GLN:NE2	2.45	0.49
1:XA:974:A:O4'	14:XN:31:ARG:CD	2.53	0.49
1:XA:1106:G:H5'	3:XC:172:ARG:HD2	1.95	0.49
1:XA:1305:G:C4	1:XA:1331:G:N1	2.81	0.49
1:XA:1536:C:O2	23:XX:10:G:N2	2.44	0.49
6:XF:61:LEU:HD23	6:XF:63:TYR:HE2	1.77	0.49
19:XS:40:ILE:HD13	19:XS:71:LEU:HD21	1.95	0.49
26:Y2:16:LEU:O	26:Y2:67:LYS:NZ	2.45	0.49
34:YA:1771:C:H2'	34:YA:1772:G:C8	2.48	0.49
34:YA:2092:U:OP2	41:YI:27:ARG:NH2	2.43	0.49
34:YA:2153:G:H2'	34:YA:2154:G:C8	2.47	0.49
34:YA:2781:A:H5''	34:YA:2782:G:H5'	1.95	0.49
44:YP:52:GLU:OE1	44:YP:55:ARG:NH1	2.45	0.49
1:QA:739:C:O2	15:QO:42:HIS:CE1	2.66	0.49
1:QA:940:C:H2'	1:QA:941:G:C8	2.48	0.49
1:QA:1238:A:H2'	1:QA:1239:A:H8	1.77	0.49
1:QA:1253:G:O5'	10:QJ:44:VAL:O	2.30	0.49
1:QA:1313:U:O4	19:QS:4:SER:CB	2.52	0.49
1:QA:1481:U:H2'	1:QA:1482:G:C8	2.48	0.49
2:QB:178:ARG:NH2	2:QB:198:ASP:OD1	2.39	0.49
20:QT:56:MET:HG3	20:QT:84:LEU:HD21	1.94	0.49
34:RA:318:C:H2'	34:RA:319:C:C6	2.48	0.49
34:RA:1566:A:N1	36:RD:214:TRP:CE2	2.81	0.49
34:RA:2291:U:O3'	34:RA:2379:G:N2	2.46	0.49
34:RA:2532:G:O2'	34:RA:2657:A:N1	2.44	0.49
34:RA:2581:G:N2	34:RA:2581:G:OP2	2.46	0.49
1:XA:132:C:OP1	20:XT:75:ASN:ND2	2.46	0.49
1:XA:308:C:H2'	1:XA:309:G:H5'	1.94	0.49
1:XA:1187:G:N3	14:XN:61:TRP:O	2.46	0.49
7:XG:118:VAL:HG22	7:XG:122:HIS:CE1	2.48	0.49
13:XM:91:ARG:NE	13:XM:97:PRO:O	2.46	0.49
20:XT:41:ILE:HD13	20:XT:87:LYS:HG2	1.95	0.49
31:Y7:37:LYS:CE	34:YA:458:G:C4	2.95	0.49
34:YA:662:G:H5''	44:YP:17:LYS:HG2	1.93	0.49
34:YA:911:A:N1	45:YQ:9:TYR:CG	2.80	0.49
34:YA:1028:A:N6	34:YA:1125:G:H2'	2.28	0.49
1:QA:1292:U:H2'	1:QA:1293:G:H8	1.78	0.49
6:QF:35:ALA:HB1	6:QF:65:VAL:HG21	1.95	0.49
30:R6:18:ARG:O	30:R6:20:ASN:ND2	2.45	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:270(Z):G:H4'	34:RA:273(A):G:H4'	1.95	0.49
34:RA:839:U:H2'	34:RA:840:C:H6	1.78	0.49
34:RA:1788:C:OP1	36:RD:222:ARG:NH2	2.45	0.49
43:RO:88:ASN:HD21	43:RO:90:GLN:HB2	1.78	0.49
1:XA:131:C:OP2	1:XA:186(K):G:O2'	2.30	0.49
1:XA:669:U:H1'	15:XO:46:HIS:HE1	1.77	0.49
1:XA:861:G:HO2'	1:XA:874:G:HO2'	1.49	0.49
1:XA:1540:U:O2	23:XX:6:G:N2	2.46	0.49
24:Y0:11:ARG:O	24:Y0:14:ARG:NH2	2.43	0.49
28:Y4:38:LYS:HE2	39:YG:112:PRO:HG3	1.94	0.49
34:YA:1128:A:H1'	34:YA:1129:A:C4	2.47	0.49
34:YA:1657:C:H2'	34:YA:1658:C:C6	2.47	0.49
34:YA:1794:U:H2'	34:YA:1795:C:C6	2.48	0.49
34:YA:2119:A:H61	34:YA:2168:G:H1	1.61	0.49
34:YA:2391:G:C6	34:YA:2427:C:H1'	2.48	0.49
45:YQ:66:ILE:HA	45:YQ:104:PHE:HA	1.95	0.49
48:YT:28:VAL:HG12	48:YT:88:ILE:HA	1.95	0.49
1:QA:1049:U:H5	14:QN:3:ARG:CG	2.23	0.49
1:QA:1160:G:H5'	2:QB:132:LYS:HZ1	1.78	0.49
34:RA:500:G:H1'	34:RA:505:A:H61	1.78	0.49
34:RA:534:U:HO2'	49:RU:49:HIS:CG	2.30	0.49
34:RA:1070:A:O2'	34:RA:1097:U:O3'	2.30	0.49
34:RA:1802:A:N1	34:RA:1822:G:H1'	2.28	0.49
34:RA:2563:U:H1'	34:RA:2566:A:N6	2.28	0.49
1:XA:68(J):G:H1	1:XA:68(R):U:H3	1.60	0.49
1:XA:107:G:H3'	1:XA:108:G:N2	2.26	0.49
1:XA:1047:G:O2'	1:XA:1215:G:O2'	2.16	0.49
1:XA:1226:C:H3'	13:XM:103:THR:CB	2.43	0.49
1:XA:1227:A:H5''	13:XM:111:LYS:HE2	1.95	0.49
1:XA:1414:U:H2'	1:XA:1415:G:H8	1.78	0.49
4:XD:56:VAL:HG13	4:XD:57:ARG:HD2	1.94	0.49
6:XF:100:ASN:ND2	18:XR:26:LEU:O	2.44	0.49
34:YA:448:U:C4	34:YA:583:G:H1'	2.48	0.49
34:YA:1174:A:H2'	34:YA:1175:U:H4'	1.95	0.49
34:YA:1221:C:H2'	34:YA:1222:C:H6	1.78	0.49
34:YA:2002:G:H2'	34:YA:2003:G:H8	1.78	0.49
1:QA:1108:G:P	3:QC:175:LEU:H	2.29	0.48
1:QA:1244:C:H2'	1:QA:1245:A:C8	2.48	0.48
1:QA:1248:A:N6	21:QU:26:LYS:CD	2.56	0.48
1:QA:1372:U:H5''	9:QI:69:GLY:CA	2.43	0.48
3:QC:47:LEU:HD11	3:QC:87:LEU:HD21	1.95	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
20:QT:54:LYS:HE3	20:QT:100:ILE:HG21	1.95	0.48
34:RA:554:U:H2'	34:RA:556:G:C8	2.47	0.48
53:RY:15:VAL:HA	53:RY:72:VAL:HA	1.94	0.48
54:RZ:163:LEU:HD22	54:RZ:167:PRO:HG3	1.95	0.48
1:XA:518:C:H2'	1:XA:530:G:N3	2.27	0.48
1:XA:1118:C:H1'	1:XA:1179:A:C5	2.48	0.48
1:XA:1223:C:OP1	19:XS:78:ARG:NH2	2.46	0.48
1:XA:1251:A:H2'	1:XA:1252:A:C8	2.48	0.48
10:XJ:48:THR:O	14:XN:34:TYR:OH	2.29	0.48
32:Y8:2:PRO:N	34:YA:591:C:O2	2.45	0.48
32:Y8:13:ARG:HD2	44:YP:61:ARG:HE	1.78	0.48
33:Y9:14:CYS:HB3	33:Y9:27:CYS:HB2	1.95	0.48
33:Y9:19:ARG:HG3	34:YA:2756:U:H5''	1.94	0.48
34:YA:1141:U:P	42:YN:25:ARG:HH21	2.36	0.48
34:YA:1667:G:O2'	34:YA:1991:U:O4	2.30	0.48
34:YA:2691:C:O3'	34:YA:2871:C:H4'	2.12	0.48
54:YZ:24:LEU:HD23	54:YZ:41:LEU:HG	1.94	0.48
1:QA:447:G:H2'	1:QA:485:G:H22	1.78	0.48
1:QA:936:C:H2'	1:QA:937:A:H8	1.78	0.48
1:QA:973:G:H5''	1:QA:974:A:H3'	1.95	0.48
1:QA:1179:A:C5'	9:QI:83:ARG:NH1	2.72	0.48
1:QA:1219:U:H1'	19:QS:34:TRP:CZ3	2.48	0.48
17:QQ:66:SER:O	17:QQ:70:ARG:NH1	2.46	0.48
33:R9:23:VAL:HG21	34:RA:1032:A:O2'	2.13	0.48
34:RA:363(B):A:H2'	34:RA:363(C):G:C8	2.49	0.48
34:RA:441:U:O2	38:RF:46:ARG:NH2	2.46	0.48
34:RA:685:A:H5''	34:RA:788:A:N6	2.25	0.48
34:RA:687:C:N3	34:RA:787:U:H4'	2.28	0.48
34:RA:1394:U:H4'	34:RA:1603:A:H4'	1.93	0.48
34:RA:1482:U:H3	34:RA:1512:G:H1	1.61	0.48
34:RA:1759:A:H1'	34:RA:2711:A:C2	2.47	0.48
43:RO:120:GLU:OE1	48:RT:67:SER:OG	2.24	0.48
1:XA:24:U:H2'	1:XA:25:C:C6	2.49	0.48
1:XA:590:C:OP1	8:XH:29:SER:HB2	2.14	0.48
1:XA:1533:C:H5	23:XX:13:A:H61	1.61	0.48
4:XD:72:GLU:OE2	4:XD:207:TYR:OH	2.19	0.48
6:XF:82:ARG:HB3	6:XF:85:VAL:HG12	1.95	0.48
21:XU:3:LYS:CD	21:XU:14:TRP:CZ3	2.93	0.48
34:YA:236:C:H2'	34:YA:237:C:H6	1.78	0.48
34:YA:910:A:N3	34:YA:2264:C:O2'	2.42	0.48
34:YA:1479:G:H1	34:YA:1514:U:H3	1.61	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:2832:U:H1'	34:YA:2834:G:C4	2.48	0.48
36:YD:108:PRO:HB3	36:YD:143:HIS:HE1	1.77	0.48
41:YI:129:THR:HG22	41:YI:137:PRO:HB3	1.95	0.48
1:QA:875:C:O2'	8:QH:14:ARG:HD2	2.13	0.48
1:QA:1014:A:N3	1:QA:1219:U:H1'	2.27	0.48
1:QA:1141:C:H2'	1:QA:1142:G:C8	2.47	0.48
1:QA:1226:C:O5'	13:QM:91:ARG:NH1	2.44	0.48
1:QA:1280:A:C5'	10:QJ:40:LEU:CD2	2.80	0.48
1:QA:1314:C:P	19:QS:6:LYS:HZ2	2.34	0.48
2:QB:69:LEU:HB3	2:QB:162:ILE:HG22	1.94	0.48
5:QE:152:ARG:HG2	8:QH:42:GLU:O	2.14	0.48
24:R0:43:THR:CG2	34:RA:2336:A:H61	2.26	0.48
34:RA:27:G:N2	34:RA:513:A:OP2	2.44	0.48
34:RA:436:C:H2'	34:RA:438:G:C8	2.48	0.48
34:RA:523:C:O2	34:RA:553:U:O2'	2.30	0.48
34:RA:568:U:OP1	44:RP:36:LYS:HD2	2.13	0.48
34:RA:1173:G:O2'	34:RA:1175:U:O4'	2.30	0.48
34:RA:2594:C:H2'	34:RA:2595:G:C8	2.47	0.48
34:RA:2831:G:H1'	34:RA:2883:A:H2'	1.95	0.48
35:RB:15:A:H5'	35:RB:16:G:C8	2.47	0.48
40:RH:3:ARG:HG3	40:RH:3:ARG:O	2.13	0.48
40:RH:3:ARG:HH12	40:RH:5:GLY:HA2	1.77	0.48
1:XA:201(C):U:H4'	1:XA:216:G:N2	2.28	0.48
1:XA:447:G:H2'	1:XA:485:G:N2	2.28	0.48
1:XA:673:G:H2'	1:XA:674:G:C8	2.48	0.48
4:XD:63:LYS:HD2	4:XD:198:VAL:HG12	1.95	0.48
6:XF:89:MET:HE1	18:XR:76:LEU:HD13	1.92	0.48
6:XF:94:GLN:NE2	18:XR:32:ARG:HH11	2.11	0.48
15:XO:10:LYS:HA	15:XO:13:GLN:HG2	1.95	0.48
34:YA:2037:G:H2'	34:YA:2038:G:C8	2.49	0.48
34:YA:2140:C:H2'	34:YA:2141:G:C8	2.48	0.48
34:YA:2152:G:H2'	34:YA:2153:G:C8	2.48	0.48
34:YA:2590:A:H2'	34:YA:2591:C:C6	2.48	0.48
35:YB:12:C:O4'	35:YB:15:A:N6	2.46	0.48
47:YS:4:LEU:HD22	47:YS:8:GLU:CD	2.34	0.48
48:YT:31:SER:OG	48:YT:85:LYS:NZ	2.44	0.48
1:QA:22:G:H2'	1:QA:23:C:C6	2.48	0.48
1:QA:957:U:H5'	19:QS:81:ARG:HB2	1.93	0.48
1:QA:1187:G:H1'	14:QN:61:TRP:N	2.28	0.48
1:QA:1229:A:OP2	13:QM:108:ARG:NH2	2.40	0.48
1:QA:1286:A:H2'	1:QA:1287:A:H4'	1.94	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
4:QD:79:PHE:HE1	4:QD:204:ILE:HD13	1.77	0.48
13:QM:83:ASP:O	19:QS:66:MET:HE3	2.13	0.48
34:RA:29:U:H2'	34:RA:30:G:C8	2.49	0.48
34:RA:118:A:N3	34:RA:178:G:H1'	2.29	0.48
34:RA:243:U:H2'	34:RA:244:A:H8	1.78	0.48
34:RA:270(S):G:H2'	34:RA:270(T):G:C8	2.49	0.48
34:RA:834:C:H2'	34:RA:835:A:H8	1.78	0.48
34:RA:2142:C:H2'	34:RA:2143:C:C6	2.48	0.48
34:RA:2211:G:N2	34:RA:2212:A:H2	2.11	0.48
43:RO:15:GLY:O	43:RO:47:ILE:N	2.45	0.48
53:RY:76:CYS:SG	53:RY:79:CYS:CA	2.94	0.48
1:XA:222:U:H2'	1:XA:223:U:C6	2.49	0.48
1:XA:522:C:H1'	1:XA:536:C:H5''	1.94	0.48
1:XA:689:C:H4'	1:XA:705:U:H1'	1.94	0.48
1:XA:988:G:N2	1:XA:1016:A:O2'	2.47	0.48
5:XE:60:TYR:OH	5:XE:64:ARG:NH2	2.47	0.48
16:XP:40:ASP:HB3	16:XP:48:TRP:HB2	1.94	0.48
34:YA:65:C:H2'	34:YA:66:C:H6	1.78	0.48
34:YA:1010:A:H1'	34:YA:1153:C:H1'	1.95	0.48
41:YI:77:LEU:HD13	41:YI:101:LEU:HB3	1.96	0.48
54:YZ:130:PRO:HA	54:YZ:133:ILE:HD11	1.96	0.48
1:QA:18:C:H4'	1:QA:1078:U:O2	2.14	0.48
1:QA:620:C:H1'	4:QD:135:LEU:HG	1.96	0.48
1:QA:947:G:OP1	13:QM:108:ARG:CB	2.61	0.48
1:QA:974:A:OP1	14:QN:29:ARG:NH1	2.46	0.48
1:QA:1522:U:H2'	1:QA:1523:G:C8	2.48	0.48
5:QE:102:ALA:O	5:QE:107:ARG:NH2	2.45	0.48
34:RA:141(A):A:H8	34:RA:1595:G:H21	1.61	0.48
34:RA:414:C:O2	34:RA:1864:U:O2'	2.21	0.48
34:RA:589:C:H2'	34:RA:590:A:H8	1.79	0.48
34:RA:1266:G:N7	51:RW:15:ARG:NH1	2.62	0.48
1:XA:156:G:H2'	1:XA:157:G:C8	2.48	0.48
1:XA:859:A:H2'	1:XA:860:A:O4'	2.13	0.48
1:XA:1004:A:C2	1:XA:1024:G:H2'	2.49	0.48
1:XA:1310:G:H1	1:XA:1327:C:H5	1.61	0.48
1:XA:1319:A:N6	1:XA:1361:G:H1'	2.29	0.48
1:XA:1472:U:H2'	1:XA:1473:A:C8	2.49	0.48
3:XC:153:VAL:HB	3:XC:196:LEU:HD21	1.95	0.48
4:XD:20:TYR:HD1	4:XD:26:CYS:HB3	1.78	0.48
6:XF:5:GLU:HA	6:XF:63:TYR:O	2.14	0.48
10:XJ:9:ARG:HG2	10:XJ:9:ARG:HH11	1.78	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:XX:6:G:H2'	23:XX:7:G:H8	1.77	0.48
29:Y5:9:LYS:CE	34:YA:2019:A:N7	2.75	0.48
34:YA:956:G:OP2	45:YQ:14:ARG:NH2	2.46	0.48
34:YA:1657:C:H4'	37:YE:133:LYS:HB3	1.95	0.48
34:YA:2084:C:H2'	34:YA:2085:C:C6	2.48	0.48
34:YA:2094:G:OP1	41:YI:22:LYS:HD2	2.12	0.48
48:YT:52:ILE:HG13	48:YT:61:PHE:HB3	1.96	0.48
1:QA:626:U:O3'	16:QP:38:TYR:CD1	2.66	0.48
1:QA:1151:A:O2'	10:QJ:13:HIS:HB2	2.13	0.48
1:QA:1270:C:O2'	1:QA:1313:U:O3'	2.30	0.48
34:RA:1073:A:H3'	34:RA:1074:G:H8	1.79	0.48
34:RA:1518:C:H2'	34:RA:1519:G:C8	2.49	0.48
38:RF:48:THR:O	38:RF:48:THR:OG1	2.32	0.48
38:RF:159:GLY:O	38:RF:164:ARG:NH2	2.41	0.48
44:RP:106:LEU:HD13	44:RP:112:LEU:HD13	1.95	0.48
46:RR:26:LYS:O	46:RR:30:THR:OG1	2.24	0.48
1:XA:186(Q):U:H2'	1:XA:191:G:C8	2.48	0.48
1:XA:1159:U:C6	1:XA:1182:G:H2'	2.49	0.48
1:XA:1316:G:H4'	14:YN:17:LYS:HG2	0.64	0.48
1:XA:1351:U:H2'	1:XA:1352:C:H6	1.78	0.48
17:XQ:66:SER:H	17:XQ:69:LYS:HB3	1.78	0.48
19:XS:36:ARG:HB2	19:XS:72:GLY:HA3	1.94	0.48
34:YA:629:G:H1'	34:YA:639:U:H1'	1.96	0.48
34:YA:2113:U:C5	34:YA:2114:A:H1'	2.48	0.48
1:QA:155:C:H2'	1:QA:156:G:C8	2.48	0.48
1:QA:490:G:H2'	1:QA:491:G:C8	2.48	0.48
1:QA:608:A:O2'	16:QP:9:PHE:HE1	1.95	0.48
1:QA:986:A:O2'	19:QS:55:LYS:CA	2.60	0.48
1:QA:1106:G:O3'	3:QC:172:ARG:CB	2.56	0.48
1:QA:1278:U:H4'	1:QA:1279:A:C4	2.49	0.48
1:QA:1318:A:N3	19:QS:37:ARG:NH2	2.59	0.48
34:RA:813:U:C5	44:RP:25:SER:HB3	2.49	0.48
34:RA:851:U:H2'	34:RA:852:G:C8	2.48	0.48
34:RA:1732:A:H3'	34:RA:1733:G:H8	1.77	0.48
34:RA:2751:G:C6	40:RH:3:ARG:HB2	2.47	0.48
53:RY:13:VAL:HA	53:RY:74:PRO:HA	1.96	0.48
54:RZ:61:LEU:HD23	54:RZ:67:LEU:HD23	1.94	0.48
1:XA:68(H):G:C5	1:XA:68(I):G:H1'	2.49	0.48
1:XA:112:G:H22	1:XA:315:A:H2	1.60	0.48
1:XA:186(B):C:H2'	1:XA:186(C):C:C6	2.48	0.48
1:XA:856:C:H2'	1:XA:857:C:C6	2.49	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1391:U:H2'	1:XA:1392:G:C8	2.49	0.48
4:XD:91:SER:HB2	4:XD:191:ARG:HD2	1.95	0.48
13:XM:3:ARG:HD2	13:XM:7:VAL:HG12	1.96	0.48
31:Y7:37:LYS:HG2	34:YA:458:G:C8	2.49	0.48
34:YA:305:U:H2'	34:YA:306:U:C6	2.49	0.48
34:YA:1753:G:H2'	34:YA:1755:A:N7	2.28	0.48
34:YA:2053:G:C2	34:YA:2617:C:N3	2.82	0.48
34:YA:2314:C:H2'	34:YA:2315:G:C8	2.49	0.48
34:YA:2688:U:H1'	34:YA:2721:A:H61	1.78	0.48
46:YR:3:HIS:O	46:YR:5:LYS:N	2.47	0.48
1:QA:308:C:H2'	1:QA:309:G:C8	2.49	0.48
1:QA:624:C:H4'	16:QP:11:SER:HB3	1.96	0.48
1:QA:667:G:H2'	1:QA:668:G:C8	2.49	0.48
1:QA:695:A:OP2	11:QK:53:SER:HB3	2.14	0.48
1:QA:1318:A:C4	19:QS:37:ARG:NH2	2.78	0.48
1:QA:1530:G:H2'	1:QA:1531:A:H8	1.78	0.48
13:QM:26:GLY:O	13:QM:30:ALA:CB	2.62	0.48
34:RA:985:C:H2'	34:RA:986:C:C6	2.49	0.48
39:RG:126:ASP:OD1	39:RG:130:ASN:N	2.43	0.48
40:RH:43:VAL:HG23	40:RH:52:VAL:HG12	1.95	0.48
40:RH:46:GLU:HB2	40:RH:49:VAL:HG23	1.95	0.48
1:XA:263:A:H2'	1:XA:264:U:C5	2.49	0.48
1:XA:647:C:H2'	1:XA:648:A:C8	2.49	0.48
1:XA:784:C:H2'	1:XA:785:G:H8	1.79	0.48
1:XA:967:C:H5'	1:XA:968:A:H2'	1.96	0.48
1:XA:1312:G:H2'	19:XS:6:LYS:HZ2	1.77	0.48
1:XA:1409:C:H2'	1:XA:1410:G:C8	2.49	0.48
4:XD:57:ARG:HH12	5:XE:107:ARG:CZ	2.27	0.48
13:XM:84:ILE:HD12	19:XS:65:ASN:OD1	2.13	0.48
16:XP:6:LEU:HD13	16:XP:17:TYR:CG	2.49	0.48
17:XQ:83:ASP:N	17:XQ:83:ASP:OD1	2.40	0.48
20:XT:73:HIS:HB3	20:XT:74:LYS:H	1.47	0.48
34:YA:389:G:H22	44:YP:71:VAL:CG1	2.19	0.48
34:YA:860:U:H2'	34:YA:861:A:C8	2.43	0.48
1:QA:43:C:OP1	16:QP:13:HIS:HD2	1.97	0.48
1:QA:643:C:H2'	1:QA:644:G:H8	1.77	0.48
2:QB:47:THR:HG23	2:QB:202:PRO:HG2	1.96	0.48
2:QB:179:LYS:HA	8:QH:72:PRO:HG3	1.96	0.48
2:QB:197:VAL:O	8:QH:68:ARG:CZ	2.60	0.48
10:QJ:4:ILE:HG12	10:QJ:100:THR:HG22	1.95	0.48
34:RA:270(V):C:H2'	34:RA:270(W):G:C8	2.49	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:558:G:H2'	34:RA:559:G:C8	2.49	0.48
34:RA:1403:C:H5''	34:RA:1471:A:H1'	1.96	0.48
34:RA:1454:U:OP1	46:RR:77:ARG:NH1	2.45	0.48
34:RA:1607:C:H41	34:RA:1621:U:H3'	1.79	0.48
34:RA:1662:C:H2'	34:RA:1663:C:C6	2.49	0.48
34:RA:2210:G:H5'	34:RA:2211:G:C5	2.49	0.48
34:RA:2241:A:H2'	34:RA:2242:G:C8	2.49	0.48
34:RA:2553:G:C2	34:RA:2583:G:H1'	2.49	0.48
34:RA:2657:A:O3'	40:RH:160:LYS:NZ	2.46	0.48
1:XA:106:C:H2'	1:XA:107:G:C8	2.49	0.48
1:XA:669:U:O4'	15:XO:46:HIS:HE1	1.97	0.48
1:XA:1229:A:H62	13:XM:104:ARG:HB3	1.79	0.48
1:XA:1397:C:OP2	5:XE:24:ARG:NH2	2.40	0.48
10:XJ:78:ASN:HB2	10:XJ:81:THR:HG23	1.96	0.48
25:Y1:83:GLU:HG3	25:Y1:85:LEU:H	1.78	0.48
34:YA:29:U:H2'	34:YA:30:G:C8	2.49	0.48
34:YA:356:G:H2'	34:YA:357:A:C8	2.49	0.48
34:YA:410:G:N2	34:YA:2407:G:N7	2.62	0.48
34:YA:828:U:H4'	34:YA:831:G:C6	2.49	0.48
34:YA:920:G:H2'	34:YA:921:G:H8	1.78	0.48
34:YA:1162:G:H2'	34:YA:1163:G:H8	1.79	0.48
34:YA:1223:C:H2'	34:YA:1224:G:C8	2.49	0.48
34:YA:1952:A:P	43:YO:44:LYS:HZ3	2.35	0.48
34:YA:2097:C:H2'	34:YA:2098:U:H6	1.79	0.48
37:YE:37:ARG:O	37:YE:45:THR:HA	2.13	0.48
37:YE:117:MET:HA	37:YE:122:PHE:H	1.79	0.48
39:YG:68:PRO:HB3	39:YG:92:VAL:HB	1.95	0.48
45:YQ:35:VAL:HG12	45:YQ:102:VAL:HG22	1.94	0.48
53:YY:11:ASP:OD1	53:YY:11:ASP:N	2.43	0.48
1:QA:625:G:HO2'	16:QP:16:HIS:CE1	2.30	0.48
1:QA:695:A:H61	1:QA:786:G:H21	1.61	0.48
1:QA:986:A:N9	19:QS:54:GLY:O	2.47	0.48
1:QA:1069:C:H1'	1:QA:1191:A:H2	1.77	0.48
1:QA:1318:A:N6	14:QN:16:PHE:CG	2.82	0.48
1:QA:1407:C:H2'	1:QA:1408:A:H8	1.79	0.48
18:QR:59:SER:OG	18:QR:60:ALA:N	2.46	0.48
34:RA:153:C:H2'	34:RA:154:G:C8	2.49	0.48
34:RA:433:C:H2'	34:RA:434:U:C6	2.49	0.48
34:RA:1213:A:H1'	34:RA:1238:G:N3	2.28	0.48
34:RA:2343:C:H2'	34:RA:2344:U:C6	2.49	0.48
34:RA:2710:C:H2'	34:RA:2711:A:C8	2.48	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:62:U:H2'	1:XA:63:C:C6	2.49	0.48
1:XA:372:C:O2	1:XA:372:C:O2'	2.23	0.48
1:XA:495:A:H1'	1:XA:497:A:H2'	1.96	0.48
1:XA:1256:A:N7	1:XA:1278:U:H5'	2.29	0.48
1:XA:1434:A:H61	1:XA:1467:G:H1'	1.79	0.48
2:XB:30:ARG:NH1	2:XB:31:TYR:OH	2.46	0.48
9:XI:73:GLN:O	9:XI:77:ILE:HG12	2.14	0.48
34:YA:917:A:H3'	34:YA:918:A:C8	2.49	0.48
34:YA:1050:A:N7	34:YA:2751:G:C4	2.80	0.48
34:YA:2097:C:N3	34:YA:2192:G:C6	2.82	0.48
34:YA:2546:U:H4'	34:YA:2566:A:H2	1.79	0.48
34:YA:2695:C:H2'	34:YA:2696:U:C6	2.49	0.48
40:YH:70:THR:O	40:YH:74:ASN:ND2	2.47	0.48
1:QA:1253:G:C5'	10:QJ:44:VAL:H	2.24	0.47
1:QA:1300:G:H1'	1:QA:1303:C:N4	2.29	0.47
1:QA:1309:G:H4'	13:QM:77:ASN:HD21	1.77	0.47
5:QE:74:GLY:O	5:QE:116:THR:OG1	2.31	0.47
5:QE:84:PHE:N	5:QE:87:SER:O	2.45	0.47
11:QK:82:VAL:HG13	11:QK:108:ILE:HA	1.95	0.47
25:R1:17:SER:O	25:R1:17:SER:OG	2.29	0.47
25:R1:78:LYS:NZ	34:RA:270(S):G:H21	2.10	0.47
31:R7:24:THR:HG23	31:R7:27:GLY:H	1.79	0.47
34:RA:65:C:H2'	34:RA:66:C:C6	2.49	0.47
34:RA:137(B):G:N2	52:RX:41:ASN:HD21	2.12	0.47
34:RA:514:A:H2'	34:RA:515:A:C8	2.48	0.47
34:RA:657:U:H2'	34:RA:658:C:C6	2.48	0.47
34:RA:1367:A:C5	34:RA:1368:G:H1'	2.49	0.47
35:RB:8:U:H3	35:RB:112:G:H1	1.61	0.47
41:RI:51:ILE:HA	41:RI:54:GLN:HG2	1.95	0.47
42:RN:60:ILE:HD12	42:RN:60:ILE:HA	1.81	0.47
1:XA:1106:G:C5'	3:XC:172:ARG:HG2	2.44	0.47
1:XA:1376:U:H2'	1:XA:1377:A:C8	2.48	0.47
13:XM:14:ARG:HG2	13:XM:16:ASP:H	1.79	0.47
20:XT:61:SER:OG	20:XT:62:LEU:N	2.47	0.47
34:YA:685:A:O2'	34:YA:773:U:O4	2.25	0.47
34:YA:1930:G:H2'	34:YA:1968:G:H1	1.78	0.47
34:YA:2199:A:N1	34:YA:2226:C:N4	2.57	0.47
34:YA:2641:G:H2'	34:YA:2642:G:H8	1.79	0.47
1:QA:401:C:H2'	1:QA:402:G:C8	2.50	0.47
1:QA:562:C:H1'	12:QL:15:ARG:HD2	1.95	0.47
1:QA:949:A:OP1	13:QM:100:GLY:O	2.32	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:953:G:C6	13:QM:104:ARG:CZ	2.97	0.47
1:QA:1129:C:H1'	1:QA:1132:C:C5	2.49	0.47
1:QA:1253:G:C5'	10:QJ:44:VAL:O	2.62	0.47
5:QE:79:GLU:CD	8:QH:102:ARG:HH12	2.18	0.47
29:R5:12:SER:O	29:R5:16:ARG:HB2	2.14	0.47
34:RA:558:G:H2'	34:RA:559:G:H8	1.79	0.47
34:RA:1681:G:HO2'	34:RA:1762:A:HO2'	1.57	0.47
34:RA:2197:U:H1'	34:RA:2198:A:C8	2.49	0.47
34:RA:2233:U:H2'	34:RA:2234:G:C8	2.48	0.47
47:RS:26:LEU:HB3	47:RS:87:PHE:HA	1.96	0.47
48:RT:91:ARG:NH2	48:RT:124:ASP:OD2	2.47	0.47
49:RU:17:ILE:HG13	49:RU:39:LEU:HD12	1.95	0.47
1:XA:822:C:H2'	1:XA:823:G:C8	2.49	0.47
1:XA:1187:G:N9	14:YN:61:TRP:O	2.42	0.47
2:XB:71:VAL:HB	2:XB:164:VAL:HG12	1.96	0.47
13:XM:84:ILE:CG2	19:XS:74:PHE:HZ	2.21	0.47
34:YA:270(A):A:C2	34:YA:366:C:H4'	2.50	0.47
34:YA:2115:G:H4'	34:YA:2166:G:H4'	1.96	0.47
34:YA:2150:U:H2'	34:YA:2151:G:H8	1.76	0.47
37:YE:102:VAL:O	37:YE:170:LEU:N	2.45	0.47
39:YG:107:LEU:HA	39:YG:111:LEU:HD12	1.95	0.47
1:QA:186(B):C:H2'	1:QA:186(C):C:O4'	2.15	0.47
1:QA:258:G:OP1	20:QT:87:LYS:NZ	2.48	0.47
1:QA:345:C:H5'	48:RT:41:ARG:NH1	2.29	0.47
1:QA:658:G:H5''	15:QO:31:LEU:CD1	2.44	0.47
1:QA:1320:C:N3	19:QS:36:ARG:C	2.67	0.47
1:QA:1534:A:N3	23:QX:12:A:N6	2.55	0.47
4:QD:108:LEU:HD22	4:QD:174:LEU:HD13	1.96	0.47
4:QD:169:LYS:HE2	6:XF:82:ARG:HH22	1.79	0.47
9:QI:20:ARG:HG3	9:QI:60:ASP:HB2	1.96	0.47
34:RA:678:C:H2'	34:RA:679:C:H6	1.79	0.47
34:RA:1568:G:OP1	36:RD:63:ARG:NH1	2.34	0.47
34:RA:2081:C:H2'	34:RA:2082:A:H8	1.80	0.47
34:RA:2373:G:H2'	34:RA:2374:C:C6	2.49	0.47
37:RE:105:THR:OG1	37:RE:199:ARG:NH1	2.47	0.47
1:XA:708:C:H2'	1:XA:709:G:C8	2.49	0.47
1:XA:1319:A:H61	1:XA:1361:G:H1'	1.79	0.47
4:XD:175:SER:O	4:XD:183:GLY:HA2	2.15	0.47
7:XG:99:LEU:HD12	7:XG:102:ARG:HD2	1.96	0.47
25:Y1:39:LYS:NZ	34:YA:205:G:O6	2.44	0.47
26:Y2:4:SER:OG	26:Y2:5:GLU:N	2.39	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:389:G:N1	44:YP:71:VAL:CG1	2.69	0.47
34:YA:996:A:O4'	50:YV:10:LYS:HG2	2.14	0.47
34:YA:2046:G:N1	34:YA:2623:G:C5	2.80	0.47
34:YA:2110:G:OP1	34:YA:2111:C:N4	2.47	0.47
34:YA:2795:G:O2'	34:YA:2799:A:N6	2.47	0.47
1:QA:233:C:O2'	1:QA:264:U:N3	2.47	0.47
1:QA:1123:A:O2'	10:QJ:37:PRO:N	2.47	0.47
1:QA:1307:U:P	13:QM:99:ARG:CB	3.03	0.47
1:QA:1307:U:H5''	13:QM:99:ARG:NE	2.29	0.47
16:QP:40:ASP:OD1	16:QP:43:LYS:N	2.46	0.47
31:R7:39:ARG:NH1	34:RA:459:U:OP2	2.47	0.47
34:RA:521:G:H2'	34:RA:522:G:C8	2.49	0.47
34:RA:521:G:H2'	34:RA:522:G:H8	1.79	0.47
34:RA:679:C:H2'	34:RA:680:G:H8	1.78	0.47
34:RA:990:A:C6	34:RA:1186:G:H1'	2.50	0.47
34:RA:1022:G:N2	34:RA:1023:U:O4	2.47	0.47
38:RF:102:PRO:HB2	38:RF:105:VAL:HG23	1.95	0.47
1:XA:296:U:H1'	1:XA:556:C:H1'	1.95	0.47
1:XA:372:C:N4	1:XA:389:A:N7	2.61	0.47
1:XA:608:A:C2'	1:XA:609:A:H8	2.27	0.47
1:XA:718:G:H1	18:XR:74:ARG:HH12	1.62	0.47
1:XA:1217:C:H5''	14:XN:12:ARG:NH1	2.26	0.47
1:XA:1253:G:O3'	10:XJ:45:ARG:HD2	2.14	0.47
1:XA:1318:A:O3'	19:XS:11:VAL:CG1	2.62	0.47
3:XC:23:TYR:CD2	10:XJ:95:GLU:N	2.82	0.47
11:XK:109:VAL:CA	18:XR:86:VAL:HG23	2.43	0.47
13:XM:84:ILE:O	19:XS:74:PHE:CE1	2.62	0.47
34:YA:223:A:H1'	34:YA:407:G:H21	1.79	0.47
34:YA:270(Q):C:H1'	41:YI:50:ARG:HH22	1.79	0.47
34:YA:685:A:C2	34:YA:787:U:H1'	2.49	0.47
34:YA:1050:A:H1'	34:YA:2751:G:C8	2.49	0.47
34:YA:1068:G:N2	34:YA:1096:A:O5'	2.42	0.47
34:YA:2054:A:N1	34:YA:2616:C:N3	2.62	0.47
34:YA:2312:U:H4'	39:YG:71:THR:OG1	2.14	0.47
34:YA:2577:A:H2'	34:YA:2614:A:N6	2.30	0.47
50:YV:13:ARG:NH1	50:YV:15:GLU:OE2	2.45	0.47
50:YV:52:VAL:HG21	50:YV:55:ALA:HB3	1.95	0.47
1:QA:297:G:H4'	1:QA:557:G:O2'	2.14	0.47
1:QA:956:U:O2'	19:QS:80:TYR:C	2.52	0.47
1:QA:985:C:H2'	1:QA:986:A:H8	1.78	0.47
1:QA:1106:G:C3'	3:QC:172:ARG:CG	2.87	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1112:C:N3	3:QC:178:LEU:CD1	2.78	0.47
11:QK:17:GLY:HA2	11:QK:35:PRO:HD3	1.97	0.47
13:QM:16:ASP:N	13:QM:16:ASP:OD1	2.44	0.47
32:R8:42:ARG:HD2	34:RA:2350:C:C5'	2.44	0.47
34:RA:1339:G:N2	34:RA:1603:A:N3	2.62	0.47
34:RA:1592:C:H2'	34:RA:1593:G:C8	2.50	0.47
1:XA:186(K):G:C6	1:XA:264:U:H5''	2.49	0.47
1:XA:600:C:H2'	1:XA:601:C:C6	2.49	0.47
1:XA:613:C:H2'	1:XA:614:A:C8	2.50	0.47
1:XA:813:U:H2'	1:XA:814:A:H8	1.80	0.47
1:XA:1132:C:H2'	1:XA:1133:G:C8	2.50	0.47
1:XA:1463:C:H2'	1:XA:1464:G:H8	1.79	0.47
2:XB:204:ASN:OD1	2:XB:205:ASP:N	2.48	0.47
27:Y3:18:ASP:OD1	27:Y3:18:ASP:N	2.46	0.47
34:YA:624:C:H2'	34:YA:625:G:H8	1.79	0.47
34:YA:747:U:H5	34:YA:2613:U:C4	2.30	0.47
34:YA:1035:U:H2'	34:YA:1036:G:C8	2.50	0.47
34:YA:1043:C:H2'	34:YA:1044:G:H8	1.79	0.47
34:YA:1445:C:H2'	34:YA:1446:C:C6	2.49	0.47
34:YA:1476:C:H2'	34:YA:1477:A:C8	2.49	0.47
34:YA:2181:G:H2'	34:YA:2182:G:C8	2.49	0.47
34:YA:2722:G:H5''	34:YA:2820:A:N7	2.30	0.47
34:YA:2893:G:O2'	34:YA:2894:G:N2	2.47	0.47
1:QA:9:G:H2'	1:QA:10:A:C8	2.49	0.47
1:QA:711:G:H2'	1:QA:712:A:C8	2.50	0.47
8:QH:120:THR:OG1	8:QH:121:ASP:N	2.47	0.47
12:QL:5:PRO:HG2	12:QL:10:LEU:HD21	1.96	0.47
23:QX:3:C:H2'	23:QX:4:A:C8	2.49	0.47
34:RA:270(D):C:H2'	34:RA:270(E):C:C6	2.50	0.47
34:RA:436:C:H2'	34:RA:438:G:H8	1.79	0.47
34:RA:602:G:H1'	34:RA:656:G:N2	2.30	0.47
34:RA:619:G:H3'	34:RA:620:G:H21	1.79	0.47
34:RA:863:A:H2'	34:RA:864:G:C8	2.49	0.47
34:RA:863:A:H2'	34:RA:864:G:H8	1.80	0.47
1:XA:376:G:H4'	16:XP:5:ARG:NH1	2.28	0.47
1:XA:708:C:H2'	1:XA:709:G:H8	1.80	0.47
1:XA:908:A:H2'	1:XA:909:A:H8	1.80	0.47
1:XA:1112:C:C1'	3:XC:179:ARG:NE	2.67	0.47
1:XA:1112:C:H6	3:XC:179:ARG:HH21	1.51	0.47
1:XA:1143:G:H2'	1:XA:1144:G:C8	2.49	0.47
13:XM:83:ASP:OD2	13:XM:84:ILE:N	2.47	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:XN:23:ARG:NH1	14:XN:28:GLY:HA2	2.29	0.47
15:XO:39:LEU:HG	15:XO:56:LEU:HD12	1.96	0.47
34:YA:363(B):A:H2'	34:YA:363(C):G:C8	2.50	0.47
34:YA:1400:G:H2'	34:YA:1401:G:C8	2.49	0.47
34:YA:1947:C:H2'	34:YA:1948:G:H8	1.79	0.47
34:YA:2008:C:H2'	34:YA:2009:G:H8	1.78	0.47
34:YA:2245:U:H5'	34:YA:2246:G:H5''	1.97	0.47
34:YA:2443:C:H2'	34:YA:2444:G:C8	2.49	0.47
36:YD:35:LYS:HG3	36:YD:63:ARG:HG3	1.96	0.47
37:YE:105:THR:HG21	37:YE:164:ARG:HH21	1.78	0.47
47:YS:25:ARG:HG3	47:YS:88:ASP:HB2	1.96	0.47
1:QA:35:G:O2'	12:QL:118:SER:O	2.20	0.47
1:QA:66:G:H1'	1:QA:173:U:H2'	1.97	0.47
1:QA:269:C:H2'	1:QA:270:A:H8	1.80	0.47
1:QA:667:G:C2	15:QO:49:ASP:OD1	2.68	0.47
1:QA:833:U:H2'	1:QA:834:C:C6	2.49	0.47
1:QA:926:G:N2	23:QX:18:C:OP2	2.45	0.47
1:QA:932:C:H2'	1:QA:933:G:C8	2.49	0.47
1:QA:960:U:H2'	1:QA:1225:A:N6	2.30	0.47
1:QA:1253:G:H5'	10:QJ:44:VAL:N	2.23	0.47
1:QA:1278:U:H4'	1:QA:1279:A:C5	2.49	0.47
1:QA:1280:A:C2	10:QJ:41:PRO:HD3	2.49	0.47
1:QA:1397:C:H4'	1:QA:1398:A:C8	2.50	0.47
10:QJ:50:ILE:HD13	14:QN:41:ARG:NH1	2.18	0.47
17:QQ:62:SER:OG	17:QQ:72:ARG:NE	2.47	0.47
17:QQ:83:ASP:N	17:QQ:83:ASP:OD1	2.47	0.47
34:RA:508:G:C6	51:RW:9:TYR:CE1	3.03	0.47
34:RA:2008:C:H2'	34:RA:2009:G:C8	2.49	0.47
34:RA:2102:U:H2'	34:RA:2103:C:C6	2.50	0.47
34:RA:2319:G:N1	34:RA:2334:G:OP2	2.45	0.47
34:RA:2392:A:H2	34:RA:2424:C:H42	1.61	0.47
34:RA:2468:G:OP2	34:RA:2468:G:N2	2.41	0.47
34:RA:2503:A:O2'	34:RA:2505:G:OP2	2.29	0.47
34:RA:2658:C:H5''	40:RH:158:HIS:CD2	2.49	0.47
52:RX:64:LYS:HD2	52:RX:73:ARG:HH12	1.77	0.47
54:RZ:182:LYS:HA	54:RZ:182:LYS:HD2	1.75	0.47
1:XA:413:G:H4'	1:XA:414:A:H5''	1.97	0.47
1:XA:539:A:H2'	1:XA:540:G:H8	1.80	0.47
1:XA:595:G:N1	1:XA:641:U:O2'	2.47	0.47
1:XA:608:A:H2'	1:XA:609:A:O4'	2.15	0.47
1:XA:656:C:H2'	1:XA:657:G:H8	1.79	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:893:C:H2'	1:XA:894:G:C8	2.50	0.47
1:XA:1084:G:H21	1:XA:1102:A:N6	2.13	0.47
1:XA:1141:C:H2'	1:XA:1142:G:C8	2.50	0.47
1:XA:1269:A:N1	1:XA:1312:G:O2'	2.41	0.47
4:XD:62:GLN:HE22	4:XD:65:ARG:HH21	1.62	0.47
4:XD:175:SER:HB3	4:XD:184:LYS:HB3	1.96	0.47
4:XD:208:SER:OG	5:XE:101:ILE:HD12	2.14	0.47
29:Y5:3:LYS:HD3	34:YA:2577:A:H5'	1.97	0.47
29:Y5:58:LEU:HD23	46:YR:113:LEU:HD11	1.96	0.47
34:YA:151:C:H2'	34:YA:152:G:H8	1.79	0.47
34:YA:320:A:N3	38:YF:169:ASN:ND2	2.62	0.47
34:YA:413:C:H2'	34:YA:414:C:C6	2.49	0.47
34:YA:536:A:OP1	49:YU:53:ARG:NH1	2.47	0.47
34:YA:603:A:H5''	34:YA:655:A:H61	1.80	0.47
34:YA:624:C:H2'	34:YA:625:G:C8	2.50	0.47
34:YA:816:C:H2'	34:YA:817:C:C6	2.50	0.47
34:YA:998:C:H2'	34:YA:999:U:O4'	2.14	0.47
34:YA:1332:G:N2	34:YA:1609:A:O2'	2.48	0.47
34:YA:1529:A:H61	34:YA:1542:G:H1'	1.78	0.47
34:YA:1796:U:H2'	34:YA:1797:C:C6	2.49	0.47
34:YA:2115:G:H22	34:YA:2163:C:H3'	1.78	0.47
34:YA:2316:C:H2'	34:YA:2317:C:H6	1.79	0.47
39:YG:63:ILE:HG22	39:YG:143:GLU:HB2	1.95	0.47
45:YQ:75:THR:HB	45:YQ:86:GLY:HA3	1.96	0.47
1:QA:8:A:C4	4:QD:209:ARG:O	2.68	0.47
1:QA:229:U:H2'	1:QA:230:G:H8	1.78	0.47
1:QA:626:U:O3'	16:QP:38:TYR:CE1	2.68	0.47
1:QA:1099:G:OP2	2:QB:96:ARG:NH1	2.48	0.47
34:RA:251:A:C4	34:RA:252:G:H1'	2.50	0.47
34:RA:730:C:H2'	34:RA:731:C:H6	1.80	0.47
34:RA:1141:U:H1'	34:RA:1142(B):A:C5	2.49	0.47
34:RA:1189:A:C2	34:RA:1190:G:H1'	2.50	0.47
34:RA:1375:C:H2'	34:RA:1376:C:H6	1.78	0.47
34:RA:2084:C:H2'	34:RA:2085:C:H6	1.79	0.47
34:RA:2208:U:O2'	36:RD:150:LYS:O	2.32	0.47
34:RA:2244:U:H2'	34:RA:2245:U:C6	2.50	0.47
34:RA:2262:U:H2'	34:RA:2263:C:C6	2.50	0.47
40:RH:6:ARG:HE	40:RH:65:HIS:HB3	1.80	0.47
1:XA:106:C:H2'	1:XA:107:G:H8	1.80	0.47
1:XA:702:A:C6	34:YA:1848:A:N3	2.83	0.47
1:XA:789:U:H2'	1:XA:791:G:N7	2.30	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:826:C:H4'	8:XH:12:ARG:HG3	1.97	0.47
1:XA:961:U:O2	1:XA:1201:A:N6	2.47	0.47
1:XA:1187:G:N2	14:XN:60:SER:HB2	2.26	0.47
1:XA:1512:U:H3	1:XA:1523:G:H1	1.63	0.47
2:XB:84:GLU:HG3	2:XB:215:LEU:HB3	1.96	0.47
2:XB:184:VAL:HG23	2:XB:198:ASP:H	1.79	0.47
7:XG:16:LEU:HD21	9:XI:44:VAL:C	2.34	0.47
27:Y3:51:ALA:HA	27:Y3:54:VAL:HG12	1.95	0.47
33:Y9:14:CYS:CA	33:Y9:27:CYS:HB2	2.44	0.47
34:YA:671:C:H2'	34:YA:672:C:H6	1.79	0.47
34:YA:749:C:H5'	34:YA:1271:G:H1'	1.96	0.47
34:YA:860:U:C5	34:YA:2268:A:N7	2.83	0.47
34:YA:1838:C:N4	34:YA:1899:G:O4'	2.48	0.47
34:YA:1853:A:H2'	34:YA:1854:A:C8	2.49	0.47
34:YA:2197:U:H1'	34:YA:2198:A:C8	2.50	0.47
35:YB:24:G:O6	35:YB:56:G:O2'	2.26	0.47
40:YH:18:GLU:HB3	40:YH:25:LYS:HG2	1.97	0.47
45:YQ:55:VAL:HG22	54:YZ:178:GLU:HG2	1.97	0.47
47:YS:11:LYS:HG3	47:YS:15:ARG:HE	1.80	0.47
1:QA:106:C:C5	20:QT:15:ARG:NH1	2.83	0.47
1:QA:337:C:H2'	1:QA:338:A:C8	2.50	0.47
1:QA:986:A:C5'	19:QS:55:LYS:HG3	2.44	0.47
1:QA:1065:U:C4	1:QA:1190:G:H1'	2.49	0.47
1:QA:1109:C:H3'	1:QA:1110:A:H8	1.79	0.47
1:QA:1222:G:O3'	19:QS:78:ARG:NH2	2.48	0.47
1:QA:1236:A:P	21:QU:10:ARG:HD3	2.55	0.47
1:QA:1427:U:H2'	1:QA:1428:A:C8	2.49	0.47
3:QC:50:ALA:HB2	3:QC:75:VAL:HB	1.96	0.47
13:QM:26:GLY:O	13:QM:30:ALA:HB2	2.15	0.47
15:QO:26:GLU:HA	15:QO:29:VAL:HG12	1.97	0.47
34:RA:305:U:H2'	34:RA:306:U:C6	2.50	0.47
34:RA:442:G:H1'	38:RF:48:THR:HG21	1.96	0.47
34:RA:478:A:N6	34:RA:500:G:O2'	2.47	0.47
34:RA:482:A:H1'	34:RA:498:G:N2	2.30	0.47
34:RA:1011:G:O2'	34:RA:1012:U:O2'	2.28	0.47
34:RA:1416:G:H2'	34:RA:1417:C:C6	2.49	0.47
34:RA:1768:U:H2'	34:RA:1769:G:H8	1.79	0.47
34:RA:2152:G:H2'	34:RA:2153:G:C8	2.49	0.47
34:RA:2346:A:H5''	34:RA:2346:A:N3	2.30	0.47
34:RA:2691:C:H2'	34:RA:2692:C:C6	2.50	0.47
49:RU:6:THR:HG21	49:RU:10:ARG:HH12	1.80	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
51:RW:68:ARG:NH2	51:RW:109:GLU:OE2	2.48	0.47
1:XA:831:U:H2'	1:XA:832:C:C6	2.50	0.47
1:XA:1096:C:H2'	1:XA:1097:C:C6	2.49	0.47
1:XA:1294:G:H2'	1:XA:1295:G:C8	2.49	0.47
1:XA:1305:G:C4	1:XA:1331:G:C5	3.03	0.47
28:Y4:23:GLU:O	28:Y4:25:TYR:N	2.46	0.47
34:YA:619:G:H3'	34:YA:620:G:H21	1.79	0.47
34:YA:1050:A:C8	34:YA:2751:G:C6	3.02	0.47
34:YA:1050:A:N7	34:YA:2751:G:N1	2.63	0.47
34:YA:1417:C:H2'	34:YA:1418:G:O4'	2.15	0.47
34:YA:2047:U:H2'	34:YA:2048:G:C8	2.49	0.47
34:YA:2406:U:C4	44:YP:75:ILE:HD11	2.50	0.47
34:YA:2804:C:H2'	34:YA:2805:G:C8	2.50	0.47
41:YI:88:ILE:HG22	41:YI:90:GLY:H	1.80	0.47
1:QA:115:G:H21	1:QA:117:G:H1	1.62	0.47
1:QA:625:G:H5'	16:QP:9:PHE:HB3	1.97	0.47
1:QA:673:G:H2'	1:QA:674:G:C8	2.49	0.47
1:QA:742:G:H4'	15:QO:58:MET:SD	2.55	0.47
1:QA:830:G:H2'	1:QA:831:U:C6	2.50	0.47
1:QA:1108:G:OP1	3:QC:174:PRO:HA	2.14	0.47
1:QA:1179:A:H2'	1:QA:1180:A:C8	2.50	0.47
1:QA:1352:C:H2'	1:QA:1353:G:C8	2.50	0.47
2:QB:134:GLU:HA	2:QB:137:ARG:HG2	1.96	0.47
4:QD:156:GLU:O	4:QD:160:GLN:N	2.48	0.47
34:RA:181:A:H1'	34:RA:435:C:H5'	1.96	0.47
34:RA:303:U:H2'	34:RA:304:G:C8	2.50	0.47
34:RA:577:G:O2'	34:RA:1254:A:OP1	2.30	0.47
34:RA:611:C:H2'	34:RA:612:G:H8	1.80	0.47
34:RA:1728:G:H8	34:RA:1732:A:H62	1.61	0.47
34:RA:2036:C:H2'	34:RA:2037:G:H8	1.79	0.47
34:RA:2710:C:H2'	34:RA:2711:A:H8	1.79	0.47
34:RA:2751:G:P	34:RA:2751:G:H8	2.37	0.47
1:XA:34:C:H2'	1:XA:35:G:H8	1.80	0.47
1:XA:324:G:P	20:XT:70:SER:HG	2.38	0.47
1:XA:444:C:H2'	1:XA:445:G:C8	2.50	0.47
1:XA:649:G:H2'	1:XA:650:G:C8	2.49	0.47
1:XA:657:G:H21	15:XO:22:THR:CG2	2.27	0.47
1:XA:1253:G:N3	1:XA:1355:G:O2'	2.46	0.47
1:XA:1318:A:O2'	19:XS:11:VAL:HG13	2.09	0.47
11:XK:31:THR:HA	11:XK:42:TRP:HA	1.97	0.47
19:XS:4:SER:HB2	19:XS:7:LYS:HG2	1.95	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
31:Y7:12:ARG:NH1	34:YA:686:G:O6	2.47	0.47
33:Y9:31:LYS:HE2	34:YA:2528:U:H5'	1.97	0.47
34:YA:1154:G:OP2	49:YU:58:ARG:NH2	2.48	0.47
34:YA:2059:A:C2	34:YA:2503:A:N1	2.83	0.47
34:YA:2738:A:H2	34:YA:2766:G:H22	1.62	0.47
38:YF:157:VAL:HG13	38:YF:194:MET:HB3	1.97	0.47
1:QA:97:U:H2'	1:QA:99:C:C2	2.50	0.46
1:QA:233:C:H2'	1:QA:234:C:H6	1.80	0.46
1:QA:806:C:H2'	1:QA:807:A:C8	2.50	0.46
1:QA:815:A:N6	1:QA:1509:C:H1'	2.30	0.46
1:QA:1195:C:H6	1:QA:1196:U:H4'	1.79	0.46
1:QA:1309:G:C4'	13:QM:77:ASN:HD21	2.24	0.46
1:QA:1422:G:H5'	43:RO:48:PRO:HG3	1.97	0.46
6:QF:24:GLU:OE1	6:QF:28:ARG:NH1	2.48	0.46
8:QH:108:GLY:HA3	8:QH:138:TRP:HB3	1.97	0.46
9:QI:26:VAL:HG12	9:QI:61:ALA:HB3	1.97	0.46
13:QM:108:ARG:HA	13:QM:111:LYS:HB2	1.97	0.46
25:R1:2:SER:O	25:R1:61:ARG:NH1	2.47	0.46
25:R1:90:ILE:HA	25:R1:94:LEU:HD11	1.92	0.46
34:RA:191:A:H2'	34:RA:192:C:H6	1.78	0.46
34:RA:1266:G:H1'	34:RA:1267:U:H5	1.80	0.46
34:RA:1569:A:H5'	36:RD:61:LEU:HD11	1.97	0.46
34:RA:2100:G:H1	34:RA:2189:U:H3	1.64	0.46
34:RA:2198:A:H5'	41:RI:33:ARG:HH12	1.79	0.46
34:RA:2260:C:HO2'	34:RA:2388:A:HO2'	1.58	0.46
34:RA:2308:G:H22	34:RA:2311:A:H2	1.63	0.46
34:RA:2351:G:H1'	34:RA:2367:G:H22	1.80	0.46
34:RA:2393:A:H4'	44:RP:61:ARG:O	2.15	0.46
34:RA:2841:C:H2'	34:RA:2842:G:C8	2.51	0.46
35:RB:93:C:H5''	54:RZ:20:ARG:HH21	1.80	0.46
37:RE:24:THR:HG21	37:RE:188:VAL:HG22	1.97	0.46
40:RH:11:VAL:HG12	40:RH:13:LYS:HG2	1.97	0.46
1:XA:186(B):C:O2	20:XT:105:SER:CA	2.62	0.46
1:XA:421:U:H4'	3:XC:192:THR:CG2	2.43	0.46
1:XA:978:A:H62	14:XN:18:VAL:CG2	2.28	0.46
1:XA:1057:G:H3'	1:XA:1058:G:H8	1.80	0.46
1:XA:1309:G:H2'	1:XA:1310:G:C8	2.50	0.46
4:XD:18:LYS:HZ3	4:XD:33:MET:HG3	1.80	0.46
8:XH:91:ARG:HG3	17:XQ:33:GLY:O	2.15	0.46
12:XL:12:ARG:HH21	12:XL:13:LYS:HE3	1.79	0.46
34:YA:144:C:H2'	34:YA:145:G:C8	2.50	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:504:U:H5''	34:YA:505:A:H5'	1.98	0.46
34:YA:2011:U:OP2	51:YW:16:LYS:NZ	2.35	0.46
34:YA:2863:C:H2'	34:YA:2864:G:H8	1.80	0.46
38:YF:12:LEU:HB3	38:YF:126:VAL:HG12	1.96	0.46
1:QA:625:G:H5''	16:QP:9:PHE:HB3	1.97	0.46
1:QA:705:U:H3'	1:QA:706:A:C8	2.50	0.46
1:QA:757:U:H5''	1:QA:822:C:O2'	2.15	0.46
1:QA:758:G:H8	1:QA:758:G:O5'	1.99	0.46
1:QA:955:U:H2'	1:QA:956:U:O4'	2.15	0.46
1:QA:1313:U:C2'	19:QS:6:LYS:HZ3	2.25	0.46
9:QL:28:VAL:HA	9:QL:63:ILE:HB	1.97	0.46
10:QJ:12:ASP:O	10:QJ:16:LEU:HB3	2.16	0.46
34:RA:301:G:P	53:RY:84:ARG:HH22	2.38	0.46
34:RA:358:U:H2'	34:RA:359:A:H8	1.80	0.46
34:RA:1853:A:H2'	34:RA:1854:A:C8	2.49	0.46
34:RA:2417:C:OP1	44:RP:64:LYS:NZ	2.49	0.46
35:RB:74:U:H1'	54:RZ:34:ASN:ND2	2.29	0.46
1:XA:60:A:C2	1:XA:378:G:H1'	2.50	0.46
1:XA:191:G:N2	20:XT:103:GLY:O	2.27	0.46
1:XA:376:G:OP1	16:XP:67:THR:CG2	2.34	0.46
1:XA:1142:G:H3'	1:XA:1143:G:H8	1.80	0.46
1:XA:1303:C:H3'	1:XA:1304:G:C8	2.51	0.46
1:XA:1359:C:H41	14:XN:35:ARG:NE	2.13	0.46
1:XA:1495:U:O2'	1:XA:1496:C:H5'	2.16	0.46
34:YA:380:U:H2'	34:YA:381:G:C8	2.50	0.46
34:YA:475:U:H4'	34:YA:510:C:H5'	1.97	0.46
34:YA:671:C:H2'	34:YA:672:C:C6	2.49	0.46
34:YA:1466:G:H3'	34:YA:1547:C:H41	1.80	0.46
34:YA:2045:C:H2'	34:YA:2046:G:H8	1.81	0.46
34:YA:2142:C:H2'	34:YA:2143:C:C6	2.51	0.46
42:YN:16:ILE:HG21	42:YN:26:LEU:HD11	1.97	0.46
44:YP:98:GLU:HA	44:YP:101:VAL:HG12	1.96	0.46
1:QA:417:C:H2'	1:QA:418:C:C6	2.50	0.46
1:QA:866:C:H4'	1:QA:919:A:H5'	1.97	0.46
1:QA:980:C:H6	14:QN:19:ARG:HG3	1.63	0.46
1:QA:1150:U:C2	10:QJ:39:PRO:HG2	2.50	0.46
1:QA:1188:A:C5'	14:QN:58:LYS:HZ1	2.25	0.46
1:QA:1203:C:H5'	14:QN:3:ARG:HD2	1.97	0.46
1:QA:1524:C:H2'	1:QA:1525:G:C8	2.50	0.46
3:QC:66:VAL:HB	3:QC:101:LEU:HG	1.98	0.46
8:QH:121:ASP:OD2	8:QH:125:ARG:NH1	2.48	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:QP:21:VAL:HG23	16:QP:33:ILE:HB	1.96	0.46
32:R8:46:ARG:NH1	34:RA:631:A:OP2	2.48	0.46
34:RA:456:C:C5	52:RX:69:TYR:CZ	3.03	0.46
34:RA:462:C:H2'	34:RA:463:G:C8	2.51	0.46
34:RA:464:U:N3	34:RA:684:G:O2'	2.46	0.46
34:RA:1827:C:OP2	36:RD:222:ARG:NH1	2.49	0.46
34:RA:2031:A:N3	34:RA:2455:G:O2'	2.43	0.46
34:RA:2447:G:H1	34:RA:2451:A:H62	1.63	0.46
40:RH:123:PHE:HE2	40:RH:133:VAL:HG22	1.80	0.46
40:RH:126:PRO:HG2	40:RH:130:ARG:HG3	1.96	0.46
46:RR:28:LEU:HD23	46:RR:48:VAL:HG21	1.98	0.46
1:XA:891:U:H2'	1:XA:892:A:C8	2.49	0.46
1:XA:926:G:N2	23:XX:18:C:OP2	2.48	0.46
1:XA:1351:U:H2'	1:XA:1352:C:C6	2.50	0.46
1:XA:1392:G:H2'	1:XA:1393:U:C6	2.50	0.46
12:XL:8:ASN:O	12:XL:12:ARG:HB2	2.15	0.46
16:XP:20:VAL:HG12	16:XP:35:LYS:HA	1.97	0.46
22:XV:56:C:N4	34:YA:2112:G:O6	2.48	0.46
34:YA:192:C:H1'	34:YA:800:A:N6	2.31	0.46
34:YA:383:U:H2'	34:YA:385:C:C5	2.50	0.46
34:YA:553:U:H2'	34:YA:554:U:C6	2.51	0.46
34:YA:920:G:H2'	34:YA:921:G:C8	2.51	0.46
34:YA:2102:U:H2'	34:YA:2103:C:C6	2.49	0.46
34:YA:2688:U:H2'	34:YA:2719:G:N2	2.30	0.46
34:YA:2831:G:H1'	34:YA:2883:A:H2'	1.98	0.46
38:YF:60:SER:OG	38:YF:61:GLY:N	2.49	0.46
1:QA:341:C:H2'	1:QA:342:C:C6	2.50	0.46
1:QA:404:U:H5'	4:QD:122:ARG:HG2	1.97	0.46
1:QA:657:G:H21	15:QO:23:GLY:HA3	1.80	0.46
1:QA:668:G:H1'	15:QO:49:ASP:HB2	1.97	0.46
1:QA:1330:U:H5''	13:QM:24:GLY:HA2	1.97	0.46
2:QB:197:VAL:O	8:QH:68:ARG:NH1	2.49	0.46
32:R8:8:LYS:HG3	34:RA:246:C:N4	2.31	0.46
32:R8:13:ARG:HD2	44:RP:61:ARG:HE	1.80	0.46
34:RA:184:C:O2'	34:RA:217:G:N3	2.40	0.46
34:RA:413:C:H2'	34:RA:414:C:C6	2.49	0.46
34:RA:839:U:H2'	34:RA:840:C:C6	2.50	0.46
34:RA:1058:G:H2'	34:RA:1059:G:C8	2.50	0.46
34:RA:1802:A:H2'	34:RA:1803:A:C8	2.50	0.46
34:RA:1940:U:OP1	34:RA:1965:C:N4	2.48	0.46
34:RA:2291:U:H2'	34:RA:2292:C:H6	1.79	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:2443:C:H2'	34:RA:2444:G:C8	2.50	0.46
34:RA:2647:U:H2'	34:RA:2648:C:C6	2.50	0.46
34:RA:2819:G:H2'	34:RA:2821:A:N7	2.30	0.46
38:RF:24:LEU:HD12	38:RF:115:ALA:HB2	1.96	0.46
46:RR:35:THR:O	46:RR:35:THR:OG1	2.33	0.46
1:XA:665:A:N3	1:XA:732:C:H2'	2.30	0.46
1:XA:768:A:H1'	1:XA:1512:U:H1'	1.98	0.46
1:XA:951:G:H2'	1:XA:952:U:C6	2.51	0.46
1:XA:1131:G:H2'	1:XA:1132:C:C6	2.49	0.46
1:XA:1330:U:H5''	13:XM:24:GLY:O	2.14	0.46
1:XA:1479:C:H2'	1:XA:1480:G:C8	2.50	0.46
2:XB:188:ALA:HB3	2:XB:200:ILE:HD11	1.97	0.46
34:YA:270(E):C:H2'	34:YA:270(F):G:C8	2.51	0.46
34:YA:514:A:H2'	34:YA:515:A:C8	2.50	0.46
34:YA:839:U:H1'	34:YA:1191:G:H1'	1.96	0.46
34:YA:1454:U:H5	46:YR:73:VAL:HG12	1.80	0.46
41:YI:72:LEU:HD12	41:YI:138:ILE:HD12	1.97	0.46
1:QA:44:G:OP2	16:QP:12:LYS:HD3	2.15	0.46
1:QA:186(B):C:N1	20:QT:85:MET:CE	2.76	0.46
1:QA:309:G:C5'	16:QP:27:LYS:HZ3	2.29	0.46
1:QA:617:G:H21	16:QP:14:ASN:HD22	1.63	0.46
1:QA:877:C:H5''	8:QH:88:LYS:CD	2.46	0.46
1:QA:958:A:C4	19:QS:55:LYS:HB2	2.51	0.46
1:QA:958:A:H61	19:QS:77:THR:HB	1.80	0.46
1:QA:986:A:C4	19:QS:54:GLY:O	2.68	0.46
1:QA:1347:G:O2'	1:QA:1373:G:N1	2.44	0.46
1:QA:1514:C:H2'	1:QA:1515:C:C6	2.51	0.46
3:QC:9:GLY:HA3	14:QN:49:HIS:O	2.16	0.46
5:QE:98:THR:N	5:QE:117:ASP:OD1	2.40	0.46
34:RA:532:A:O2'	34:RA:2021:C:N4	2.47	0.46
34:RA:675:A:C2'	38:RF:67:GLN:HE22	2.28	0.46
34:RA:917:A:H3'	34:RA:918:A:H8	1.79	0.46
34:RA:1662:C:H1'	34:RA:2687:U:H5''	1.97	0.46
34:RA:2037:G:H2'	34:RA:2038:G:C8	2.50	0.46
34:RA:2398:U:H2'	34:RA:2399:G:C8	2.51	0.46
34:RA:2844:G:H3'	34:RA:2845:G:H8	1.81	0.46
38:RF:34:TRP:HB2	44:RP:6:LEU:HB3	1.98	0.46
41:RI:88:ILE:HG22	41:RI:90:GLY:N	2.31	0.46
1:XA:115:G:H1'	1:XA:116:A:OP2	2.15	0.46
1:XA:163:C:H2'	1:XA:164:U:C6	2.51	0.46
1:XA:320:C:H2'	1:XA:321:A:C8	2.51	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:324:G:H2'	1:XA:326:G:N7	2.29	0.46
1:XA:417:C:H2'	1:XA:418:C:C6	2.50	0.46
1:XA:754:C:C6	15:XO:69:TYR:CZ	2.98	0.46
1:XA:777:A:H2'	1:XA:778:G:O4'	2.15	0.46
1:XA:1016:A:H1'	1:XA:1218:C:O2	2.16	0.46
1:XA:1301:U:O3'	13:XM:17:VAL:CG2	2.64	0.46
7:XG:75:VAL:HA	7:XG:87:VAL:O	2.16	0.46
31:Y7:3:ARG:NE	34:YA:1613:G:O2'	2.45	0.46
32:Y8:4:MET:HG2	34:YA:592:G:O2'	2.16	0.46
34:YA:247:G:H4'	34:YA:386:G:C4	2.50	0.46
34:YA:2232:U:H2'	34:YA:2233:U:C6	2.50	0.46
34:YA:2855:C:H2'	34:YA:2856:C:H6	1.80	0.46
52:YX:64:LYS:HZ2	52:YX:73:ARG:HE	1.63	0.46
1:QA:757:U:H2'	1:QA:758:G:O4'	2.16	0.46
1:QA:761:G:C5'	17:QQ:100:LYS:HZ3	2.20	0.46
1:QA:975:A:C2	14:QN:34:TYR:HD1	2.29	0.46
1:QA:1241:G:H2'	1:QA:1242:C:C6	2.50	0.46
1:QA:1440(O):C:H2'	1:QA:1440(P):A:C8	2.51	0.46
16:QP:59:TRP:HA	16:QP:62:VAL:HG22	1.97	0.46
33:R9:18:ARG:HD3	34:RA:1034:G:H4'	1.97	0.46
34:RA:910:A:H2'	34:RA:911:A:C8	2.50	0.46
34:RA:1026:U:H1'	34:RA:1027:A:H5''	1.98	0.46
34:RA:1689:A:H62	34:RA:1698:A:H2	1.64	0.46
37:RE:14:ILE:HD11	37:RE:173:VAL:HG11	1.97	0.46
52:RX:72:LYS:NZ	52:RX:73:ARG:O	2.38	0.46
1:XA:59:A:N3	1:XA:59:A:H2'	2.31	0.46
1:XA:113:G:H2'	1:XA:114:U:C6	2.50	0.46
1:XA:713:G:H2'	1:XA:714:G:C8	2.51	0.46
1:XA:1270:C:H4'	1:XA:1314:C:H5'	1.98	0.46
1:XA:1304:G:C8	1:XA:1304:G:C5'	2.85	0.46
1:XA:1327:C:H2'	1:XA:1328:C:C6	2.51	0.46
1:XA:1367:C:H4'	10:XJ:48:THR:HG21	1.98	0.46
1:XA:1375:A:C4'	7:XG:28:ASN:OD1	2.62	0.46
2:XB:118:LEU:HB3	2:XB:142:LEU:HD13	1.98	0.46
34:YA:523:C:O2	34:YA:553:U:O2'	2.34	0.46
34:YA:851:U:H2'	34:YA:852:G:C8	2.50	0.46
34:YA:1454:U:OP1	46:YR:77:ARG:NE	2.35	0.46
34:YA:1566:A:N6	36:YD:214:TRP:CH2	2.83	0.46
34:YA:1772:G:N2	34:YA:1774:C:H5'	2.31	0.46
34:YA:2069:G:C2	34:YA:2443:C:O2	2.68	0.46
34:YA:2439:A:H1'	34:YA:2587:A:H5'	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:YF:41:LEU:HA	38:YF:44:ARG:HG2	1.98	0.46
40:YH:89:ILE:HD11	40:YH:94:TYR:HB3	1.98	0.46
1:QA:955:U:H2'	19:QS:83:HIS:HA	1.97	0.46
1:QA:1004:A:N6	1:QA:1025:U:O3'	2.49	0.46
1:QA:1260:C:O2'	1:QA:1283:G:O2'	2.34	0.46
1:QA:1359:C:OP2	14:QN:35:ARG:NH1	2.49	0.46
10:QJ:50:ILE:HG12	14:QN:41:ARG:HE	1.71	0.46
34:RA:212:G:H2'	34:RA:213:A:C8	2.50	0.46
34:RA:244:A:C2	34:RA:245:G:H1'	2.51	0.46
34:RA:303:U:H2'	34:RA:304:G:H8	1.81	0.46
34:RA:598:G:H5'	44:RP:11:GLY:HA3	1.97	0.46
34:RA:817:C:HO2'	34:RA:932:G:N2	2.13	0.46
34:RA:1791:A:N6	34:RA:1828:G:O2'	2.41	0.46
42:RN:3:THR:HG21	49:RU:61:TRP:HE1	1.79	0.46
52:RX:90:GLU:HA	52:RX:93:GLU:HG2	1.97	0.46
54:RZ:67:LEU:HD13	54:RZ:68:PRO:HD2	1.96	0.46
1:XA:160:A:H2'	1:XA:161:A:O4'	2.15	0.46
1:XA:186(O):U:H2'	1:XA:186(P):G:C8	2.51	0.46
1:XA:620:C:C1'	4:XD:135:LEU:HD13	2.46	0.46
1:XA:674:G:H2'	1:XA:675:A:H8	1.81	0.46
1:XA:702:A:H3'	1:XA:703:G:C8	2.48	0.46
1:XA:896:C:H2'	1:XA:897:C:H6	1.81	0.46
1:XA:1124:G:H2'	1:XA:1145:C:N3	2.30	0.46
1:XA:1228:C:OP1	13:XM:108:ARG:NH2	2.49	0.46
8:XH:86:ILE:HD12	8:XH:135:CYS:HA	1.97	0.46
10:XJ:40:LEU:HD13	10:XJ:70:ARG:HA	1.98	0.46
26:Y2:28:LYS:HD3	26:Y2:28:LYS:HA	1.77	0.46
34:YA:259:G:H21	34:YA:621:A:H1'	1.81	0.46
34:YA:2064:C:H1'	34:YA:2450:A:C2	2.51	0.46
34:YA:2730:C:H2'	34:YA:2731:G:C8	2.50	0.46
34:YA:2820:A:C6	46:YR:4:LEU:HD11	2.50	0.46
42:YN:47:ALA:HB2	42:YN:112:LEU:HD11	1.98	0.46
43:YO:80:ASP:OD2	48:YT:64:ARG:NH2	2.48	0.46
1:QA:140:A:H2'	1:QA:141:A:C8	2.51	0.46
1:QA:260:G:H2'	1:QA:261:U:C6	2.51	0.46
1:QA:414:A:H3'	1:QA:415:A:H8	1.80	0.46
1:QA:687:A:O2'	1:QA:701:C:N4	2.48	0.46
1:QA:775:G:N2	1:QA:804:U:O4	2.47	0.46
1:QA:947:G:H2'	1:QA:948:C:C6	2.51	0.46
1:QA:977:A:O2'	1:QA:981:U:O4	2.30	0.46
1:QA:1382:C:H2'	1:QA:1383:C:O4'	2.16	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:236:C:H2'	34:RA:237:C:C6	2.51	0.46
34:RA:270(G):U:H2'	34:RA:270(H):C:C6	2.51	0.46
34:RA:1147:C:H2'	34:RA:1148:A:C8	2.51	0.46
34:RA:1332:G:N2	34:RA:1609:A:O2'	2.49	0.46
34:RA:1853:A:H2'	34:RA:1854:A:H8	1.81	0.46
34:RA:1927:A:H2'	34:RA:1928:A:C8	2.51	0.46
34:RA:2574:G:H1'	37:RE:143:ASN:HB3	1.97	0.46
38:RF:56:GLU:OE2	38:RF:93:LYS:NZ	2.49	0.46
40:RH:84:SER:HA	40:RH:134:SER:HA	1.97	0.46
1:XA:68(G):C:H2'	1:XA:68(H):G:C8	2.50	0.46
1:XA:370:C:H2'	1:XA:371:G:C8	2.50	0.46
1:XA:445:G:H2'	1:XA:446:G:C8	2.51	0.46
1:XA:718:G:H3'	1:XA:719:C:C6	2.51	0.46
1:XA:728:A:N7	15:XO:54:ARG:CD	2.78	0.46
1:XA:745:C:H1'	1:XA:836:G:H1'	1.98	0.46
1:XA:948:C:OP1	13:XM:106:ASN:HB3	2.16	0.46
11:XK:108:ILE:O	18:XR:87:ARG:CA	2.64	0.46
33:Y9:20:HIS:CE1	34:YA:2756:U:H3'	2.51	0.46
34:YA:842:G:H2'	34:YA:843:G:H8	1.80	0.46
34:YA:1468:C:H2'	34:YA:1469:A:C8	2.51	0.46
34:YA:2096:U:H3	34:YA:2193:G:N2	2.13	0.46
34:YA:2232:U:H2'	34:YA:2233:U:H6	1.81	0.46
34:YA:2533:A:OP1	34:YA:2665:A:O2'	2.33	0.46
1:QA:106:C:H2'	1:QA:107:G:H8	1.81	0.46
1:QA:346:G:OP1	48:RT:41:ARG:NH2	2.48	0.46
1:QA:500:G:H1'	1:QA:547:A:N1	2.31	0.46
1:QA:568:G:N7	12:QL:5:PRO:HD3	2.31	0.46
1:QA:730:G:N3	1:QA:765:G:H4'	2.31	0.46
1:QA:1106:G:C4'	3:QC:172:ARG:CG	2.15	0.46
1:QA:1109:C:H3'	1:QA:1110:A:C8	2.51	0.46
1:QA:1131:G:H2'	1:QA:1132:C:H6	1.81	0.46
1:QA:1307:U:H5''	13:QM:99:ARG:CZ	2.46	0.46
3:QC:22:TRP:CA	10:QJ:93:GLY:CA	2.84	0.46
4:QD:22:LYS:HG3	56:QD:301:SF4:S1	2.56	0.46
11:QK:27:ASN:OD1	11:QK:28:THR:N	2.47	0.46
33:R9:11:CYS:N	33:R9:14:CYS:SG	2.83	0.46
34:RA:1351:C:H2'	34:RA:1352:U:C6	2.51	0.46
34:RA:1657:C:O3'	37:RE:133:LYS:HG2	2.16	0.46
34:RA:1952:A:OP1	43:RO:42:SER:OG	2.32	0.46
34:RA:2036:C:H2'	34:RA:2037:G:C8	2.51	0.46
34:RA:2260:C:O2'	34:RA:2388:A:O2'	2.28	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:2757:A:N1	40:RH:67:LEU:HD13	2.31	0.46
35:RB:31:C:H4'	39:RG:29:TRP:CH2	2.51	0.46
51:RW:86:LEU:HD22	51:RW:96:ILE:HD11	1.98	0.46
1:XA:450:G:H5''	1:XA:451:A:H3'	1.98	0.46
1:XA:996:A:H2'	1:XA:997:U:C6	2.51	0.46
1:XA:1157:A:C2	1:XA:1181:G:H1'	2.51	0.46
1:XA:1312:G:C2'	19:XS:6:LYS:NZ	2.68	0.46
5:XE:151:LEU:O	8:XH:64:LYS:NZ	2.49	0.46
13:XM:45:VAL:HG23	13:XM:48:LEU:HD12	1.98	0.46
20:XT:11:SER:O	20:XT:11:SER:OG	2.28	0.46
31:Y7:35:ARG:NH1	34:YA:54:G:O2'	2.38	0.46
34:YA:98:G:H1'	34:YA:103:A:H1'	1.98	0.46
34:YA:442:G:N2	38:YF:48:THR:OG1	2.48	0.46
34:YA:2688:U:H1'	34:YA:2721:A:N6	2.31	0.46
40:YH:137:ASP:OD2	40:YH:138:LYS:N	2.48	0.46
43:YO:64:ARG:HB2	43:YO:83:ALA:HB3	1.98	0.46
48:YT:91:ARG:NH2	48:YT:124:ASP:OD2	2.48	0.46
1:QA:18:C:H42	1:QA:917:G:H1	1.64	0.46
1:QA:521:G:OP1	12:QL:73:GLU:HA	2.16	0.46
2:QB:101:MET:HA	2:QB:108:ILE:HG13	1.97	0.46
2:QB:178:ARG:HH21	8:QH:70:GLN:HA	1.81	0.46
3:QC:184:TYR:HA	3:QC:200:ALA:O	2.15	0.46
4:QD:57:ARG:HB3	4:QD:206:PHE:HB2	1.97	0.46
4:QD:72:GLU:OE2	4:QD:207:TYR:OH	2.29	0.46
6:QF:33:TYR:OH	6:QF:78:GLU:OE1	2.30	0.46
16:QP:4:ILE:HD13	16:QP:21:VAL:HG12	1.97	0.46
18:QR:74:ARG:HD3	18:QR:81:PHE:HA	1.98	0.46
34:RA:1539:G:H2'	34:RA:1540:G:C8	2.51	0.46
34:RA:1689:A:H2'	34:RA:1690:A:C8	2.51	0.46
34:RA:2318:G:OP2	34:RA:2318:G:N2	2.48	0.46
34:RA:2530:A:O2'	34:RA:2534:A:N6	2.48	0.46
34:RA:2554:U:H2'	34:RA:2555:U:C6	2.51	0.46
38:RF:182:ASN:OD1	38:RF:182:ASN:N	2.46	0.46
49:RU:45:TYR:O	49:RU:49:HIS:ND1	2.49	0.46
1:XA:129(B):G:H4'	1:XA:130:A:H5''	1.98	0.46
1:XA:663:A:H2'	1:XA:664:G:C8	2.51	0.46
1:XA:669:U:H1'	15:XO:46:HIS:CE1	2.50	0.46
1:XA:697:U:H1'	1:XA:786:G:H1'	1.97	0.46
1:XA:739:C:P	15:XO:2:PRO:HD3	2.56	0.46
1:XA:1410:G:H2'	1:XA:1411:C:C6	2.50	0.46
2:XB:27:LYS:HD2	2:XB:193:ASP:HB2	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:XI:7:THR:O	9:XI:7:THR:OG1	2.31	0.46
10:XJ:50:ILE:CD1	14:XN:41:ARG:NH1	2.79	0.46
12:XL:53:ARG:HB3	12:XL:69:TYR:HE1	1.79	0.46
22:XV:50:G:H2'	22:XV:51:A:H8	1.81	0.46
32:Y8:34:TRP:HD1	34:YA:2420:C:OP1	1.98	0.46
34:YA:691:C:H2'	34:YA:692:C:C6	2.50	0.46
34:YA:1380:G:O2'	34:YA:1569:A:N6	2.49	0.46
34:YA:1493:C:N4	34:YA:2210:G:N9	2.61	0.46
34:YA:1493:C:C6	34:YA:2210:G:C6	3.04	0.46
34:YA:1645:G:H5''	34:YA:1646:C:H5'	1.98	0.46
34:YA:2043:C:N4	34:YA:2625:G:H1	2.13	0.46
34:YA:2291:U:H2'	34:YA:2292:C:C6	2.51	0.46
34:YA:2801:A:C5	34:YA:2802:G:H1'	2.51	0.46
45:YQ:133:ARG:HG3	45:YQ:134:ARG:H	1.80	0.46
54:YZ:74:VAL:HG22	54:YZ:86:VAL:HG23	1.97	0.46
1:QA:538:G:H2'	1:QA:539:A:C8	2.51	0.45
1:QA:612:C:H2'	1:QA:613:C:C6	2.51	0.45
1:QA:784:C:H2'	1:QA:785:G:C8	2.51	0.45
1:QA:956:U:C5'	19:QS:83:HIS:HA	2.46	0.45
1:QA:957:U:O2'	19:QS:79:THR:C	2.53	0.45
1:QA:1294:G:H2'	1:QA:1295:G:H8	1.81	0.45
1:QA:1359:C:H4'	1:QA:1362(A):C:N4	2.31	0.45
2:QB:73:THR:O	2:QB:75:LYS:NZ	2.50	0.45
3:QC:5:ILE:CG2	14:QN:45:ARG:NH2	2.78	0.45
7:QG:70:LYS:HB2	7:QG:96:GLN:HB3	1.98	0.45
32:R8:13:ARG:HG2	44:RP:63:PRO:HB3	1.98	0.45
32:R8:29:LYS:O	32:R8:31:HIS:N	2.46	0.45
34:RA:689:A:H2'	34:RA:690:G:C8	2.51	0.45
34:RA:748:G:OP1	51:RW:88:ARG:NH2	2.40	0.45
34:RA:956:G:H2'	34:RA:957:A:H2'	1.98	0.45
34:RA:1102:C:H2'	34:RA:1103:A:H8	1.81	0.45
34:RA:2081:C:H2'	34:RA:2082:A:C8	2.51	0.45
37:RE:46:ALA:HB1	37:RE:80:GLU:HG2	1.99	0.45
46:RR:38:VAL:HG12	46:RR:112:ALA:HB2	1.96	0.45
49:RU:27:LEU:HD22	49:RU:31:SER:HB2	1.97	0.45
1:XA:376:G:H1	1:XA:387:U:H3	1.64	0.45
1:XA:445:G:H2'	1:XA:446:G:H8	1.80	0.45
1:XA:858:G:H3'	1:XA:869:G:H1	1.81	0.45
1:XA:974:A:O5'	14:XN:31:ARG:HD3	2.16	0.45
1:XA:983:A:H2	1:XA:984:C:C6	2.33	0.45
1:XA:1242:C:H4'	1:XA:1303:C:H4'	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1318:A:O3'	19:XS:11:VAL:HG11	2.15	0.45
1:XA:1439:C:H2'	1:XA:1440(A):C:H6	1.81	0.45
6:XF:94:GLN:NE2	18:XR:32:ARG:NH1	2.65	0.45
34:YA:270(G):U:H2'	34:YA:270(H):C:C6	2.51	0.45
34:YA:443:A:C5	38:YF:45:ARG:HD3	2.51	0.45
34:YA:919:G:C5	34:YA:2268:A:C6	2.74	0.45
34:YA:2408:U:H2'	34:YA:2409:G:C8	2.51	0.45
48:YT:16:ARG:HH21	48:YT:81:PRO:HA	1.80	0.45
1:QA:64:G:H1'	1:QA:67:C:H41	1.81	0.45
1:QA:235:C:O2'	17:QQ:4:LYS:HE3	2.17	0.45
1:QA:659:U:OP1	15:QO:8:LYS:HD3	2.16	0.45
1:QA:1065:U:H5''	1:QA:1190:G:H21	1.82	0.45
1:QA:1375:A:H3'	1:QA:1376:U:C6	2.51	0.45
8:QH:91:ARG:NE	17:QQ:32:TYR:O	2.41	0.45
10:QJ:50:ILE:HD11	14:QN:41:ARG:NE	2.22	0.45
15:QO:39:LEU:HD22	15:QO:56:LEU:HD13	1.97	0.45
34:RA:2320:A:N6	34:RA:2333:A:H2'	2.31	0.45
1:XA:109:A:C8	1:XA:326:G:H2'	2.51	0.45
1:XA:320:C:H2'	1:XA:321:A:H8	1.82	0.45
1:XA:427:U:OP2	4:XD:36:ARG:NH1	2.49	0.45
1:XA:876:G:H1'	8:XH:11:THR:HG21	1.98	0.45
1:XA:1185:G:H2'	1:XA:1186:G:O4'	2.16	0.45
1:XA:1305:G:C2'	1:XA:1331:G:N2	2.64	0.45
6:XF:100:ASN:O	18:XR:28:GLU:HG3	2.15	0.45
9:XI:28:VAL:HG12	9:XI:63:ILE:HB	1.98	0.45
29:Y5:4:HIS:O	34:YA:2056:G:N2	2.50	0.45
34:YA:1461:G:P	34:YA:1461:G:H8	2.39	0.45
34:YA:1925:C:H2'	34:YA:1926:U:C6	2.51	0.45
34:YA:2157:G:O2'	34:YA:2158:A:O4'	2.34	0.45
46:YR:33:ARG:HA	46:YR:114:VAL:O	2.16	0.45
1:QA:345:C:H1'	1:QA:346:G:C2	2.52	0.45
1:QA:1105:A:H2'	1:QA:1106:G:H8	1.82	0.45
1:QA:1157:A:N6	1:QA:1178:G:N3	2.65	0.45
1:QA:1306:A:H1'	1:QA:1332:A:C5	2.52	0.45
13:QM:87:TYR:OH	13:QM:91:ARG:NH2	2.47	0.45
15:QO:89:GLY:OXT	34:RA:716:A:P	2.73	0.45
34:RA:581:C:H2'	34:RA:582:G:H8	1.79	0.45
34:RA:995:C:H5''	49:RU:54:LYS:HG2	1.98	0.45
34:RA:1173:G:H1'	34:RA:1175:U:O2	2.16	0.45
34:RA:1278:A:H2'	34:RA:1279:G:C8	2.51	0.45
34:RA:1336:A:H2'	34:RA:1337:G:C8	2.50	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:715:A:H2'	1:XA:716:A:C8	2.51	0.45
1:XA:1307:U:H2'	1:XA:1308:U:C6	2.51	0.45
2:XB:68:ILE:HG22	2:XB:161:ALA:HB3	1.98	0.45
4:XD:21:LEU:N	4:XD:26:CYS:SG	2.89	0.45
5:XE:101:ILE:O	5:XE:120:THR:OG1	2.30	0.45
12:XL:46:LYS:HG3	12:XL:48:PRO:HD2	1.98	0.45
34:YA:191:A:H1'	34:YA:679:C:H1'	1.97	0.45
34:YA:195:A:H2'	34:YA:198:C:H42	1.80	0.45
34:YA:376:C:H2'	34:YA:377:C:C6	2.51	0.45
34:YA:1173:G:H4'	34:YA:1174:A:C8	2.51	0.45
34:YA:1399:C:H2'	34:YA:1400:G:C8	2.52	0.45
34:YA:2010:G:H5''	51:YW:42:ARG:HB2	1.98	0.45
34:YA:2077:A:C5	34:YA:2435:A:C5	3.04	0.45
34:YA:2439:A:N7	34:YA:2586:C:H4'	2.31	0.45
34:YA:2630:G:H2'	34:YA:2631:G:C8	2.52	0.45
34:YA:2773:C:H2'	34:YA:2774:C:H6	1.81	0.45
34:YA:2809:A:H2'	34:YA:2810:A:C8	2.52	0.45
34:YA:2844:G:H3'	34:YA:2845:G:H8	1.81	0.45
44:YP:96:THR:HG22	44:YP:99:LEU:HD22	1.98	0.45
54:YZ:149:SER:OG	54:YZ:172:ALA:O	2.27	0.45
1:QA:730:G:N2	1:QA:765:G:H5''	2.32	0.45
1:QA:967:C:H5'	1:QA:968:A:C4	2.51	0.45
1:QA:1038:C:H2'	1:QA:1039:C:C6	2.52	0.45
1:QA:1071:C:H2'	1:QA:1072:G:C8	2.51	0.45
1:QA:1307:U:OP1	13:QM:99:ARG:HB2	2.16	0.45
1:QA:1359:C:C6	14:QN:35:ARG:CZ	3.00	0.45
2:QB:178:ARG:NH1	2:QB:196:LEU:O	2.49	0.45
4:QD:208:SER:HB2	5:QE:101:ILE:CD1	2.42	0.45
12:QL:7:ILE:CD1	17:QQ:32:TYR:HB3	2.20	0.45
24:R0:34:GLY:HA3	34:RA:2353:G:H1'	1.98	0.45
34:RA:20:C:OP1	49:RU:22:LYS:NZ	2.28	0.45
34:RA:347:A:H2'	34:RA:348:G:C8	2.52	0.45
34:RA:2692:C:O2	34:RA:2847:U:O2'	2.29	0.45
34:RA:2754:U:HO2'	34:RA:2756:U:P	2.39	0.45
36:RD:247:ALA:HA	36:RD:253:GLN:HA	1.98	0.45
39:RG:68:PRO:HB3	39:RG:92:VAL:HB	1.98	0.45
41:RI:123:LEU:HD12	41:RI:142:VAL:HG13	1.98	0.45
51:RW:76:VAL:HG22	51:RW:103:ILE:HG23	1.98	0.45
53:RY:28:LYS:HG3	53:RY:40:GLU:HG2	1.99	0.45
1:XA:68(I):G:N2	1:XA:68(S):C:N3	2.65	0.45
1:XA:309:G:O4'	1:XA:608:A:C2	2.70	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1142:G:C2	1:XA:1143:G:H1'	2.51	0.45
1:XA:1411:C:H2'	1:XA:1412:C:C6	2.52	0.45
2:XB:207:ALA:O	2:XB:210:SER:OG	2.28	0.45
7:XG:139:GLU:OE1	7:XG:143:ARG:NH2	2.49	0.45
10:XJ:16:LEU:CB	10:XJ:70:ARG:HH12	2.27	0.45
17:XQ:34:LYS:NZ	17:XQ:35:VAL:O	2.46	0.45
34:YA:106:C:H2'	34:YA:107:C:C6	2.52	0.45
34:YA:1040:C:H2'	34:YA:1041:C:C6	2.51	0.45
34:YA:1153:C:H5'	49:YU:76:TYR:HE2	1.82	0.45
34:YA:1535:U:N3	34:YA:1537:C:H1'	2.32	0.45
34:YA:2025:C:H2'	34:YA:2026:C:C6	2.51	0.45
34:YA:2044:C:N3	34:YA:2625:G:N3	2.62	0.45
34:YA:2368:C:H2'	34:YA:2369:A:C8	2.52	0.45
34:YA:2641:G:H2'	34:YA:2642:G:C8	2.51	0.45
34:YA:2692:C:H2'	34:YA:2693:A:C8	2.51	0.45
41:YI:4:ILE:HG22	41:YI:18:VAL:HB	1.98	0.45
1:QA:186(A):C:O2'	20:QT:85:MET:SD	2.59	0.45
1:QA:235:C:H2'	1:QA:236:G:C8	2.52	0.45
1:QA:403:C:H4'	4:QD:122:ARG:HD3	1.97	0.45
1:QA:958:A:N7	19:QS:79:THR:CG2	2.80	0.45
10:QJ:48:THR:HG22	10:QJ:62:HIS:HB3	1.98	0.45
10:QJ:67:THR:O	10:QJ:67:THR:OG1	2.32	0.45
28:R4:1:MET:HE2	39:RG:98:ARG:NH1	2.32	0.45
34:RA:1930:G:H2'	34:RA:1968:G:N1	2.30	0.45
34:RA:2071:A:H2'	34:RA:2072:G:H8	1.81	0.45
36:RD:13:ARG:NH1	36:RD:16:MET:SD	2.90	0.45
40:RH:98:LEU:CG	40:RH:125:VAL:CG1	2.91	0.45
40:RH:124:GLU:HG3	40:RH:124:GLU:O	2.17	0.45
43:RO:63:VAL:HB	43:RO:102:VAL:HG13	1.98	0.45
1:XA:68(N):U:H2'	1:XA:68(O):U:O4'	2.16	0.45
1:XA:341:C:H2'	1:XA:342:C:C6	2.52	0.45
1:XA:601:C:H2'	1:XA:602:A:C8	2.52	0.45
1:XA:718:G:H3'	1:XA:719:C:H6	1.81	0.45
1:XA:1028(B):C:H2'	1:XA:1028(C):C:H5	1.81	0.45
1:XA:1081:G:OP1	5:XE:18:ARG:HB3	2.17	0.45
1:XA:1124:G:O5'	10:XJ:36:GLY:CA	2.65	0.45
1:XA:1360:A:C1'	14:XN:17:LYS:HG3	2.46	0.45
1:XA:1405:G:H2'	1:XA:1406:U:H6	1.81	0.45
1:XA:1505:G:C1'	23:XX:15:A:H2	2.29	0.45
4:XD:162:LEU:HD12	4:XD:178:VAL:HG13	1.98	0.45
6:XF:36:ARG:NH2	6:XF:38:GLU:OE2	2.49	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:XL:110:VAL:H	12:XL:122:THR:HG22	1.82	0.45
25:Y1:40:ARG:HG2	25:Y1:41:ARG:H	1.81	0.45
29:Y5:32:PRO:N	29:Y5:32:PRO:C	2.61	0.45
34:YA:286:C:H2'	34:YA:287:C:C6	2.52	0.45
34:YA:834:C:H2'	34:YA:835:A:C8	2.51	0.45
34:YA:1165:U:H2'	34:YA:1166:C:C6	2.52	0.45
34:YA:2031:A:N3	34:YA:2455:G:O2'	2.43	0.45
34:YA:2696:U:H2'	34:YA:2697:G:C8	2.51	0.45
36:YD:133:LEU:HD23	36:YD:136:ILE:HD12	1.98	0.45
39:YG:138:GLN:OE1	39:YG:153:ARG:N	2.43	0.45
40:YH:10:PRO:O	40:YH:49:VAL:HA	2.17	0.45
43:YO:104:ARG:N	43:YO:122:LEU:O	2.49	0.45
49:YU:17:ILE:HG13	49:YU:32:PHE:HE1	1.82	0.45
1:QA:194:C:H4'	20:QT:68:LYS:CE	2.47	0.45
1:QA:451:A:N6	1:QA:480:U:H2'	2.32	0.45
1:QA:677:U:N1	11:QK:119:CYS:SG	2.90	0.45
1:QA:684:A:H1'	11:QK:39:PRO:HD2	1.99	0.45
1:QA:950:U:OP2	13:QM:102:ARG:HD2	2.17	0.45
1:QA:1094:G:H4'	1:QA:1095:U:C5	2.51	0.45
1:QA:1331:G:OP2	13:QM:24:GLY:N	2.34	0.45
2:QB:193:ASP:OD2	2:QB:193:ASP:N	2.50	0.45
16:QP:55:ARG:HD2	16:QP:55:ARG:HA	1.79	0.45
25:R1:90:ILE:CA	25:R1:94:LEU:CD1	2.84	0.45
31:R7:18:PHE:HB2	31:R7:43:THR:HG21	1.99	0.45
34:RA:996:A:H2'	34:RA:997:G:C8	2.51	0.45
34:RA:1048:A:H2	34:RA:1112:G:H21	1.63	0.45
34:RA:2071:A:H2'	34:RA:2072:G:C8	2.52	0.45
36:RD:132:PRO:HA	36:RD:190:TYR:HA	1.99	0.45
48:RT:102:ILE:HD12	48:RT:110:ILE:HD12	1.99	0.45
50:RV:68:LYS:HD2	50:RV:68:LYS:HA	1.69	0.45
1:XA:116:A:C8	1:XA:116:A:O5'	2.70	0.45
1:XA:156:G:H2'	1:XA:157:G:H8	1.81	0.45
1:XA:321:A:H4'	1:XA:1436:U:H5'	1.97	0.45
1:XA:634:C:H2'	1:XA:635:G:C8	2.40	0.45
1:XA:722:A:H4'	1:XA:723:U:C4	2.52	0.45
1:XA:730:G:N2	1:XA:765:G:H5''	2.32	0.45
1:XA:1117:G:H21	1:XA:1180:A:H1'	1.81	0.45
4:XD:30:LYS:HB3	4:XD:35:ARG:HH22	1.81	0.45
7:XG:79:ARG:HH21	7:XG:82:GLY:HA2	1.82	0.45
10:XJ:53:PRO:HB3	14:XN:42:ILE:CG1	2.39	0.45
34:YA:415:A:C2	34:YA:2409:G:C2	3.04	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:834:C:H2'	34:YA:835:A:H8	1.82	0.45
34:YA:2019:A:H2	34:YA:2035:G:H22	1.65	0.45
34:YA:2044:C:C5	34:YA:2625:G:N2	2.83	0.45
34:YA:2638:G:OP1	37:YE:82:ARG:NH2	2.49	0.45
34:YA:2700:C:H2'	34:YA:2701:C:C6	2.51	0.45
36:YD:71:ASP:HB2	36:YD:103:ARG:HH12	1.82	0.45
51:YW:46:PHE:O	51:YW:50:VAL:HG23	2.16	0.45
1:QA:158:G:H2'	1:QA:159:G:C8	2.52	0.45
1:QA:646:U:H2'	1:QA:647:C:C6	2.52	0.45
1:QA:950:U:H2'	1:QA:951:G:C8	2.51	0.45
1:QA:1110:A:H62	3:QC:176:HIS:HB2	1.81	0.45
10:QJ:99:LYS:HA	10:QJ:99:LYS:HD3	1.79	0.45
13:QM:91:ARG:HD2	13:QM:96:LEU:HB3	1.98	0.45
20:QT:58:LYS:O	20:QT:61:SER:OG	2.26	0.45
27:R3:18:ASP:OD1	27:R3:18:ASP:N	2.48	0.45
29:R5:41:PRO:O	29:R5:44:THR:OG1	2.27	0.45
32:R8:31:HIS:CD2	34:RA:2422:A:N6	2.83	0.45
34:RA:64:A:H2'	34:RA:65:C:O4'	2.17	0.45
34:RA:270(E):C:H2'	34:RA:270(F):G:C8	2.52	0.45
34:RA:2572:A:OP1	34:RA:2574:G:O2'	2.31	0.45
34:RA:2795:G:H2'	34:RA:2798:C:H5	1.82	0.45
50:RV:14:VAL:HB	50:RV:96:ILE:HG12	1.98	0.45
1:XA:102:G:N3	1:XA:151:A:H2	2.15	0.45
1:XA:364:A:H2'	1:XA:365:U:C2	2.51	0.45
1:XA:824:C:H2'	1:XA:825:G:H8	1.79	0.45
3:XC:21:ARG:NH1	10:XJ:15:THR:HG21	2.31	0.45
7:XG:26:PHE:O	7:XG:30:ILE:HG12	2.17	0.45
29:Y5:9:LYS:NZ	34:YA:2019:A:C8	2.83	0.45
34:YA:321:G:C4	34:YA:341:G:H4'	2.51	0.45
34:YA:839:U:H2'	34:YA:840:C:C6	2.52	0.45
34:YA:1083:U:O2	34:YA:1085:A:H3'	2.16	0.45
34:YA:1221:C:H2'	34:YA:1222:C:C6	2.52	0.45
34:YA:1445:C:H2'	34:YA:1446:C:H6	1.81	0.45
34:YA:2080:G:C6	34:YA:2241:A:N1	2.85	0.45
34:YA:2184:G:H2'	34:YA:2185:C:C6	2.51	0.45
34:YA:2263:C:H2'	34:YA:2264:C:C6	2.52	0.45
34:YA:2345:G:N3	34:YA:2381:C:H2'	2.31	0.45
34:YA:2880:C:H1'	46:YR:92:GLY:O	2.17	0.45
35:YB:90:C:H5'	45:YQ:18:LYS:HA	1.99	0.45
38:YF:155:LEU:HB2	38:YF:189:THR:HG21	1.99	0.45
42:YN:137:LYS:HD3	42:YN:138:LEU:HG	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
48:YT:54:ARG:HA	48:YT:59:THR:HG23	1.99	0.45
1:QA:435:C:H2'	1:QA:436:C:C6	2.51	0.45
1:QA:565:U:H3'	1:QA:566:G:H2'	1.99	0.45
1:QA:762:C:H2'	1:QA:763:G:C8	2.52	0.45
1:QA:987:G:H2'	1:QA:988:G:H8	1.82	0.45
1:QA:996:A:H2'	1:QA:997:U:C6	2.52	0.45
1:QA:1310:G:H2'	1:QA:1311:G:C8	2.51	0.45
3:QC:23:TYR:OH	10:QJ:9:ARG:HB3	2.17	0.45
4:QD:85:LYS:HA	4:QD:85:LYS:HD2	1.73	0.45
10:QJ:47:PHE:CE2	14:QN:34:TYR:CB	2.97	0.45
31:R7:6:GLN:O	34:RA:686:G:H8	2.00	0.45
34:RA:33:U:O4	34:RA:446:G:O2'	2.31	0.45
34:RA:270(T):G:H2'	34:RA:270(U):G:H8	1.82	0.45
34:RA:863:A:OP1	45:RQ:21:THR:OG1	2.14	0.45
34:RA:994:C:OP1	49:RU:53:ARG:NH2	2.50	0.45
34:RA:1463:C:H2'	34:RA:1464:C:H6	1.81	0.45
34:RA:1947:C:H2'	34:RA:1948:G:C8	2.52	0.45
34:RA:2010:G:H5''	51:RW:42:ARG:HB2	1.98	0.45
34:RA:2419:U:H2'	34:RA:2420:C:C6	2.49	0.45
34:RA:2514:U:H2'	34:RA:2515:C:C6	2.52	0.45
36:RD:208:LYS:HG3	36:RD:210:GLY:H	1.82	0.45
37:RE:111:ARG:HD3	37:RE:160:TYR:CE2	2.52	0.45
39:RG:161:THR:HG22	39:RG:163:ALA:H	1.82	0.45
48:RT:118:ARG:HH11	48:RT:121:ILE:HG21	1.81	0.45
1:XA:167:G:H2'	1:XA:168:G:C8	2.52	0.45
1:XA:694:A:H2'	1:XA:695:A:O4'	2.17	0.45
3:XC:60:ALA:CB	10:XJ:91:PRO:HG2	2.45	0.45
4:XD:43:HIS:HB3	4:XD:46:LYS:HD2	1.98	0.45
10:XJ:47:PHE:CZ	14:XN:36:PHE:HB3	2.51	0.45
33:Y9:25:VAL:HG22	33:Y9:34:GLN:HB3	1.98	0.45
34:YA:574:C:C5	34:YA:2054:A:H4'	2.52	0.45
34:YA:686:G:N2	34:YA:788:A:H61	2.15	0.45
34:YA:987:G:O2'	34:YA:1000:A:N3	2.42	0.45
34:YA:1071:G:H1'	34:YA:1089:G:H3'	1.98	0.45
34:YA:2098:U:O2	34:YA:2191:G:N2	2.49	0.45
34:YA:2581:G:N2	34:YA:2581:G:OP2	2.50	0.45
37:YE:171:GLU:HB2	37:YE:185:LYS:HG3	1.98	0.45
38:YF:54:ARG:HD2	38:YF:81:PRO:HD3	1.98	0.45
1:QA:186(F):C:H2'	1:QA:186(G):C:C6	2.52	0.45
1:QA:647:C:H2'	1:QA:648:A:C8	2.51	0.45
1:QA:953:G:C5	13:QM:104:ARG:CZ	2.96	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1360:A:C2'	14:QN:17:LYS:HZ1	2.30	0.45
11:QK:57:THR:HG23	11:QK:60:ALA:H	1.81	0.45
17:QQ:59:ILE:HG22	17:QQ:73:VAL:HA	1.98	0.45
22:QV:74:C:H5''	25:R1:23:LYS:HG3	1.98	0.45
34:RA:459:U:H2'	34:RA:460:A:C8	2.50	0.45
34:RA:1056:G:H5''	34:RA:1057:A:O4'	2.17	0.45
34:RA:1525:G:H2'	34:RA:1526:G:C8	2.52	0.45
34:RA:1838:C:N4	34:RA:1899:G:O4'	2.50	0.45
34:RA:2591:C:H2'	34:RA:2592:G:C8	2.52	0.45
34:RA:2688:U:OP1	34:RA:2713:A:N6	2.50	0.45
38:RF:187:VAL:HG23	44:RP:3:LEU:HD22	1.97	0.45
42:RN:28:THR:O	42:RN:32:THR:OG1	2.24	0.45
43:RO:106:LEU:HA	43:RO:109:LYS:HB2	1.99	0.45
54:RZ:54:HIS:HB3	54:RZ:101:PRO:HD3	1.99	0.45
1:XA:110:C:O5'	1:XA:110:C:C6	2.70	0.45
1:XA:186(B):C:O2	20:XT:105:SER:OG	2.34	0.45
1:XA:240:C:H2'	1:XA:241:C:H6	1.81	0.45
1:XA:302:G:N3	1:XA:556:C:H4'	2.32	0.45
3:XC:60:ALA:CA	10:XJ:91:PRO:HG2	2.47	0.45
10:XJ:47:PHE:CE1	14:XN:36:PHE:HB3	2.51	0.45
27:Y3:23:LEU:HD12	27:Y3:28:LEU:HB2	1.98	0.45
34:YA:662:G:H2'	34:YA:663:G:C8	2.51	0.45
34:YA:691:C:H2'	34:YA:692:C:H6	1.81	0.45
34:YA:1523:U:H2'	34:YA:1524:G:C8	2.52	0.45
34:YA:1571:A:H2'	34:YA:1572:A:C8	2.52	0.45
34:YA:1812:A:H2'	34:YA:1813:G:H8	1.81	0.45
34:YA:2373:G:H2'	34:YA:2374:C:C6	2.52	0.45
34:YA:2741:A:H61	34:YA:2763:G:H1'	1.82	0.45
34:YA:2849:U:C2	34:YA:2867:G:H1'	2.52	0.45
36:YD:44:ASN:HB3	36:YD:49:ILE:HG22	1.98	0.45
49:YU:8:VAL:HG22	49:YU:12:ARG:HE	1.82	0.45
1:QA:40:C:H2'	1:QA:41:G:C8	2.52	0.45
1:QA:115:G:H1'	1:QA:116:A:N7	2.32	0.45
1:QA:302:G:N3	1:QA:556:C:H4'	2.32	0.45
1:QA:505:G:H2'	1:QA:506:G:C8	2.52	0.45
7:QG:16:LEU:HD21	9:QI:42:ARG:HG2	0.48	0.45
13:QM:83:ASP:O	19:QS:66:MET:CE	2.65	0.45
19:QS:79:THR:O	19:QS:79:THR:OG1	2.35	0.45
34:RA:242:G:N2	34:RA:243:U:O4	2.50	0.45
34:RA:363(C):G:H2'	34:RA:363(D):G:H8	1.82	0.45
34:RA:1012:U:OP1	49:RU:75:ASN:ND2	2.50	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:1254:A:H5''	34:RA:1255:U:H5''	1.99	0.45
34:RA:1470:G:O2'	34:RA:1522:G:O6	2.34	0.45
34:RA:1849:G:H2'	34:RA:1850:G:H8	1.82	0.45
34:RA:1925:C:H2'	34:RA:1926:U:C6	2.52	0.45
34:RA:1939:U:OP1	34:RA:2604:U:O2'	2.35	0.45
34:RA:2250:G:O2'	34:RA:2496:C:OP1	2.23	0.45
40:RH:86:GLU:O	40:RH:164:TYR:HB2	2.17	0.45
40:RH:126:PRO:HG2	40:RH:130:ARG:CG	2.46	0.45
41:RI:79:ILE:HG23	41:RI:142:VAL:HA	1.99	0.45
41:RI:80:PRO:HB2	41:RI:146:ALA:HB2	1.97	0.45
47:RS:23:ARG:NH2	47:RS:111:GLU:OE1	2.49	0.45
1:XA:539:A:H2'	1:XA:540:G:C8	2.52	0.45
1:XA:796:C:H2'	1:XA:797:C:C6	2.52	0.45
1:XA:876:G:C1'	8:XH:11:THR:HG21	2.47	0.45
1:XA:1305:G:C2	1:XA:1331:G:C4	3.04	0.45
6:XF:91:VAL:CB	18:XR:34:TYR:OH	2.64	0.45
15:XO:64:ARG:HH12	15:XO:68:ARG:HH22	1.63	0.45
31:Y7:37:LYS:CE	34:YA:458:G:C8	2.99	0.45
33:Y9:27:CYS:SG	33:Y9:29:ASN:HB3	2.57	0.45
34:YA:285:C:H2'	34:YA:286:C:C6	2.52	0.45
34:YA:347:A:H2'	34:YA:348:G:H8	1.82	0.45
34:YA:779:U:H2'	34:YA:780:G:H8	1.81	0.45
34:YA:1458:C:H1'	34:YA:1459:G:C6	2.52	0.45
34:YA:1476:C:H2'	34:YA:1477:A:H8	1.82	0.45
34:YA:2036:C:H2'	34:YA:2037:G:H8	1.82	0.45
34:YA:2306:C:H2'	34:YA:2307:G:N2	2.32	0.45
34:YA:2863:C:H2'	34:YA:2864:G:C8	2.52	0.45
43:YO:78:ARG:NE	48:YT:73:GLU:OE1	2.47	0.45
1:QA:738:C:H2'	1:QA:739:C:C6	2.52	0.44
1:QA:973:G:C2'	14:QN:29:ARG:NH1	2.68	0.44
1:QA:1206:G:H2'	1:QA:1207:G:O4'	2.17	0.44
1:QA:1277:C:H4'	1:QA:1281:U:O4	2.17	0.44
6:QF:12:PRO:HD3	6:QF:58:GLY:HA2	1.99	0.44
8:QH:112:LEU:HD11	8:QH:133:LEU:HD12	1.99	0.44
34:RA:13:A:N1	34:RA:525:U:H2'	2.31	0.44
34:RA:823:G:H2'	34:RA:824:A:H8	1.82	0.44
34:RA:1399:C:H2'	34:RA:1400:G:C8	2.53	0.44
34:RA:1708:C:H2'	34:RA:1709:U:H6	1.82	0.44
34:RA:2229:C:H2'	34:RA:2230:G:H8	1.82	0.44
34:RA:2511:U:O2'	37:RE:138:PRO:O	2.30	0.44
34:RA:2567:G:H2'	34:RA:2568:C:C6	2.52	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:RD:61:LEU:HD23	36:RD:61:LEU:HA	1.83	0.44
37:RE:32:PRO:HA	37:RE:90:THR:HA	1.99	0.44
1:XA:165:C:H2'	1:XA:166:G:H8	1.78	0.44
1:XA:457:C:H2'	1:XA:458(A):C:C6	2.52	0.44
1:XA:696:A:H61	1:XA:797:C:HO2'	1.65	0.44
1:XA:897:C:H2'	1:XA:898:G:C8	2.52	0.44
1:XA:899:C:C6	1:XA:899:C:OP1	2.70	0.44
1:XA:1119:C:H2'	1:XA:1120:G:C8	2.52	0.44
1:XA:1126:U:H3'	1:XA:1127:G:C8	2.49	0.44
1:XA:1360:A:N3	14:XA:17:LYS:HG3	2.29	0.44
1:XA:1369:C:H2'	1:XA:1370:G:C8	2.53	0.44
1:XA:1378:C:C5	1:XA:1378:C:OP1	2.70	0.44
8:XH:121:ASP:HB2	8:XH:125:ARG:HH21	1.82	0.44
12:XL:88:GLY:H	12:XL:98:TYR:HA	1.82	0.44
34:YA:318:C:H2'	34:YA:319:C:C6	2.52	0.44
34:YA:459:U:H2'	34:YA:460:A:C8	2.52	0.44
34:YA:1394:U:H4'	34:YA:1603:A:H4'	1.99	0.44
34:YA:1397:U:OP2	34:YA:1398:C:N4	2.40	0.44
34:YA:1408:C:H2'	34:YA:1409:C:C6	2.52	0.44
34:YA:1497:U:H5''	34:YA:1498:C:H5	1.82	0.44
34:YA:2059:A:C6	34:YA:2503:A:C2	3.05	0.44
34:YA:2453:A:H2'	34:YA:2454:G:C8	2.52	0.44
34:YA:2779:U:H5'	34:YA:2779:U:O2	2.16	0.44
1:QA:511:C:C6	1:QA:534:U:H1'	2.52	0.44
1:QA:981:U:C4'	14:QN:6:LEU:HD22	2.48	0.44
1:QA:1190:G:H5'	3:QC:176:HIS:HE1	1.82	0.44
1:QA:1229:A:H2'	1:QA:1230:C:H6	1.82	0.44
1:QA:1320:C:N4	19:QS:37:ARG:HD2	2.27	0.44
28:R4:34:GLU:OE1	39:RG:113:ARG:HD3	2.17	0.44
30:R6:23:THR:HG21	34:RA:2286:A:N6	2.27	0.44
34:RA:676:A:H8	34:RA:2069:G:H21	1.64	0.44
34:RA:709:U:H2'	34:RA:710:G:C8	2.53	0.44
34:RA:845:G:H21	34:RA:933:A:H61	1.65	0.44
36:RD:95:LEU:HB2	36:RD:103:ARG:O	2.17	0.44
48:RT:25:GLY:H	48:RT:49:VAL:HG13	1.82	0.44
54:RZ:97:GLU:HB3	54:RZ:125:LEU:HD11	2.00	0.44
1:XA:62:U:H2'	1:XA:63:C:H6	1.82	0.44
1:XA:388:G:C8	1:XA:388:G:O5'	2.70	0.44
1:XA:765:G:N1	1:XA:812:C:O2'	2.45	0.44
1:XA:899:C:OP1	1:XA:899:C:C5	2.70	0.44
1:XA:932:C:H2'	1:XA:933:G:C8	2.53	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:947:G:H2'	1:XA:948:C:C6	2.52	0.44
1:XA:975:A:O4'	1:XA:1365:G:N2	2.49	0.44
1:XA:1305:G:H4'	1:XA:1332:A:H61	1.83	0.44
1:XA:1382:C:O5'	1:XA:1382:C:C6	2.70	0.44
1:XA:1483:A:H2'	1:XA:1484:C:O4'	2.17	0.44
12:XL:24:VAL:HG13	12:XL:98:TYR:HE1	1.82	0.44
27:Y3:40:THR:HB	27:Y3:43:ILE:HG12	1.98	0.44
32:Y8:2:PRO:O	34:YA:666:G:N2	2.49	0.44
34:YA:575:A:OP2	34:YA:2055:C:N4	2.40	0.44
34:YA:787:U:H5''	34:YA:788:A:H5'	1.99	0.44
34:YA:1362:C:H2'	34:YA:1363:C:C6	2.53	0.44
42:YN:21:LYS:HD2	42:YN:26:LEU:HD13	1.98	0.44
42:YN:91:LEU:HD23	42:YN:91:LEU:HA	1.81	0.44
45:YQ:58:PHE:HD2	45:YQ:61:GLY:HA3	1.82	0.44
48:YT:25:GLY:N	48:YT:49:VAL:O	2.42	0.44
49:YU:92:ARG:HH11	50:YV:11:GLN:HB2	1.82	0.44
1:QA:186(B):C:OP1	20:QT:82:SER:OG	2.23	0.44
1:QA:489:C:H2'	1:QA:490:G:C8	2.52	0.44
1:QA:877:C:O3'	8:QH:88:LYS:HD2	2.17	0.44
1:QA:1085:U:H3'	1:QA:1086:U:C5	2.53	0.44
1:QA:1100:C:H41	2:QB:96:ARG:HH22	0.52	0.44
3:QC:23:TYR:CD2	10:QJ:95:GLU:N	2.85	0.44
13:QM:37:THR:HG1	13:QM:55:ARG:HE	1.62	0.44
13:QM:80:ARG:HD3	19:QS:67:VAL:CG1	2.47	0.44
14:QN:6:LEU:HD23	14:QN:9:LYS:HD3	2.00	0.44
19:QS:50:ALA:HA	19:QS:58:VAL:O	2.17	0.44
34:RA:407:G:H2'	34:RA:408:G:H8	1.82	0.44
34:RA:579:G:O2'	34:RA:2019:A:OP1	2.28	0.44
34:RA:883:G:H2'	34:RA:884:C:H6	1.82	0.44
34:RA:1257:C:O2'	38:RF:84:VAL:HG23	2.17	0.44
34:RA:1728:G:H2'	34:RA:1731:G:O6	2.18	0.44
54:RZ:115:GLY:H	54:RZ:177:PRO:HG3	1.83	0.44
1:XA:728:A:H2'	1:XA:729:A:C8	2.53	0.44
1:XA:729:A:H2'	1:XA:730:G:C8	2.52	0.44
1:XA:899:C:C6	1:XA:899:C:OP2	2.70	0.44
1:XA:940:C:H2'	1:XA:941:G:C8	2.52	0.44
1:XA:1014:A:H1'	1:XA:1219:U:O2'	2.18	0.44
1:XA:1065:U:O2	1:XA:1067:A:N6	2.51	0.44
1:XA:1374:A:O2'	7:XG:31:MET:HE3	2.13	0.44
1:XA:1378:C:OP1	1:XA:1378:C:C6	2.70	0.44
34:YA:244:A:C2	34:YA:245:G:H1'	2.52	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:251:A:C4	34:YA:252:G:H1'	2.52	0.44
34:YA:971:C:H2'	34:YA:972:G:O4'	2.17	0.44
34:YA:1149:G:H2'	34:YA:1150:C:C6	2.52	0.44
34:YA:1367:A:C5	34:YA:1368:G:H1'	2.52	0.44
34:YA:2096:U:H3	34:YA:2193:G:H1	1.63	0.44
34:YA:2133:G:H1'	34:YA:2158:A:N6	2.32	0.44
34:YA:2316:C:H2'	34:YA:2317:C:C6	2.52	0.44
34:YA:2546:U:H5''	34:YA:2547:U:H5'	2.00	0.44
45:YQ:12:GLN:HB2	45:YQ:73:PRO:HD2	2.00	0.44
53:YY:13:VAL:HG12	53:YY:74:PRO:HA	1.99	0.44
54:YZ:5:LEU:HB2	54:YZ:59:LEU:HD12	1.98	0.44
1:QA:24:U:O3'	1:QA:524:G:O2'	2.32	0.44
1:QA:335:C:H2'	1:QA:336:C:C6	2.53	0.44
1:QA:767:A:H2'	1:QA:768:A:C8	2.53	0.44
1:QA:1114:C:O2	14:QN:61:TRP:HA	2.18	0.44
1:QA:1318:A:C2	19:QS:37:ARG:NH1	2.84	0.44
1:QA:1331:G:OP1	13:QM:23:TYR:CD2	2.70	0.44
34:RA:312:G:H4'	34:RA:331:A:N3	2.33	0.44
34:RA:632:A:H2'	34:RA:633:A:C8	2.52	0.44
34:RA:685:A:H1'	34:RA:689:A:N6	2.33	0.44
34:RA:1441:G:H2'	34:RA:1442:G:H8	1.83	0.44
34:RA:1570:A:H2'	34:RA:1571:A:C8	2.51	0.44
34:RA:1657:C:H2'	34:RA:1658:C:C6	2.52	0.44
34:RA:1825:A:H2'	34:RA:1826:G:C8	2.52	0.44
34:RA:1972:A:H2'	34:RA:1973:G:H8	1.83	0.44
34:RA:2263:C:H2'	34:RA:2264:C:H6	1.82	0.44
34:RA:2368:C:H2'	34:RA:2369:A:C8	2.53	0.44
34:RA:2730:C:H2'	34:RA:2731:G:H8	1.83	0.44
39:RG:81:LYS:HD2	39:RG:81:LYS:HA	1.74	0.44
39:RG:81:LYS:HB3	39:RG:82:LEU:H	1.59	0.44
41:RI:4:ILE:HG23	41:RI:39:ALA:HB2	2.00	0.44
1:XA:8:A:C2	4:XD:209:ARG:NH1	2.85	0.44
1:XA:592:G:H2'	1:XA:593:G:C8	2.52	0.44
1:XA:1050:G:H2'	1:XA:1051:C:C6	2.52	0.44
1:XA:1092:A:C2	1:XA:1110:A:H5'	2.52	0.44
1:XA:1142:G:H3'	1:XA:1143:G:C8	2.52	0.44
1:XA:1157:A:N3	1:XA:1157:A:H2'	2.32	0.44
1:XA:1309:G:O4'	13:XM:77:ASN:ND2	2.49	0.44
1:XA:1405:G:H2'	1:XA:1406:U:C6	2.52	0.44
1:XA:1541:U:C2	23:XX:5:A:H2	2.34	0.44
4:XD:57:ARG:NH2	4:XD:205:GLU:OE1	2.51	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
32:Y8:30:ARG:HE	44:YP:62:LEU:HD12	1.81	0.44
34:YA:222:A:H61	34:YA:232:G:H1'	1.82	0.44
34:YA:1028:A:H61	34:YA:1125:G:H2'	1.83	0.44
34:YA:1178:C:H2'	34:YA:1179:C:C6	2.53	0.44
34:YA:2327:A:H2'	34:YA:2328:A:C8	2.52	0.44
38:YF:167:ALA:HB1	38:YF:173:VAL:HG11	1.99	0.44
50:YV:2:PHE:H	50:YV:42:GLY:HA3	1.82	0.44
1:QA:24:U:H2'	1:QA:25:C:C6	2.53	0.44
1:QA:607:A:H3'	1:QA:608:A:H8	1.82	0.44
1:QA:737:A:H2'	1:QA:738:C:C6	2.52	0.44
1:QA:856:C:H4'	23:QX:7:G:H21	1.83	0.44
1:QA:1290:G:O6	21:QU:26:LYS:CE	2.63	0.44
1:QA:1515:C:H2'	1:QA:1516:G:H8	1.82	0.44
2:QB:51:LEU:HD23	2:QB:201:ILE:HD12	1.99	0.44
25:R1:29:GLY:O	34:RA:2396:G:O2'	2.36	0.44
32:R8:17:THR:OG1	32:R8:21:LYS:O	2.31	0.44
34:RA:807:U:H2'	34:RA:808:G:H8	1.83	0.44
34:RA:883:G:H2'	34:RA:884:C:C6	2.52	0.44
34:RA:894:C:H2'	34:RA:895:U:C6	2.53	0.44
34:RA:1564:C:H2'	34:RA:1565:C:C6	2.52	0.44
34:RA:1660:C:H2'	34:RA:1661:G:C8	2.51	0.44
34:RA:1947:C:H2'	34:RA:1948:G:H8	1.82	0.44
34:RA:1995:U:H3'	34:RA:1996:C:H2'	1.99	0.44
34:RA:2331:G:H2'	34:RA:2332:U:C6	2.52	0.44
34:RA:2692:C:H2'	34:RA:2693:A:C8	2.52	0.44
36:RD:12:SER:HB2	36:RD:208:LYS:HB3	1.99	0.44
51:RW:65:LEU:HD12	51:RW:68:ARG:HH21	1.83	0.44
1:XA:186(F):C:O2	1:XA:186(M):G:N2	2.50	0.44
1:XA:218:C:H2'	1:XA:219:C:C6	2.53	0.44
1:XA:341:C:H2'	1:XA:342:C:H6	1.82	0.44
1:XA:486:U:H2'	1:XA:487:A:C8	2.52	0.44
1:XA:681:C:H2'	1:XA:682:G:C8	2.53	0.44
1:XA:978:A:H62	14:YN:18:VAL:HG21	1.83	0.44
2:XB:189:ASP:OD1	2:XB:189:ASP:N	2.49	0.44
26:Y2:58:ALA:O	26:Y2:62:THR:OG1	2.30	0.44
34:YA:303:U:H2'	34:YA:304:G:C8	2.51	0.44
34:YA:414:C:O2	34:YA:1864:U:O2'	2.33	0.44
34:YA:887:A:N3	34:YA:889:C:H6	2.14	0.44
34:YA:1048:A:OP2	34:YA:1110:G:N2	2.51	0.44
34:YA:1128:A:H1'	34:YA:1129:A:C5	2.53	0.44
34:YA:2108:C:C6	34:YA:2108:C:O5'	2.70	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:2153:G:H2'	34:YA:2154:G:H8	1.82	0.44
34:YA:2294:C:H2'	34:YA:2295:C:H6	1.83	0.44
34:YA:2691:C:H5''	34:YA:2872:G:H5''	1.98	0.44
45:YQ:13:GLN:O	45:YQ:72:LYS:NZ	2.38	0.44
46:YR:100:LEU:HD11	46:YR:113:LEU:HG	2.00	0.44
1:QA:68:G:H1'	1:QA:151:A:H61	1.81	0.44
1:QA:437:U:H3	1:QA:495:A:H62	1.66	0.44
1:QA:641:U:H4'	8:QH:115:SER:HB2	2.00	0.44
1:QA:713:G:H2'	1:QA:714:G:C8	2.52	0.44
1:QA:956:U:H5'	19:QS:83:HIS:HA	2.00	0.44
1:QA:999:U:H2'	1:QA:1000:A:C8	2.53	0.44
1:QA:1270:C:H2'	1:QA:1271:G:C8	2.53	0.44
1:QA:1383:C:H2'	1:QA:1384:C:C6	2.52	0.44
6:QF:47:ARG:HD2	6:QF:57:GLN:HB3	2.00	0.44
11:QK:70:LYS:NZ	34:RA:2146:C:H3'	2.33	0.44
22:QV:71:C:H5''	34:RA:1892:C:O2'	2.17	0.44
30:R6:46:HIS:CE1	34:RA:2372:G:H1'	2.53	0.44
34:RA:553:U:H2'	34:RA:554:U:C6	2.53	0.44
34:RA:813:U:H2'	34:RA:814:C:C6	2.53	0.44
34:RA:848:G:H2'	34:RA:849:A:C8	2.53	0.44
34:RA:1152:C:H2'	34:RA:1153:C:C6	2.48	0.44
34:RA:1353:A:H2'	34:RA:1354:A:C8	2.53	0.44
34:RA:1380:G:H2'	34:RA:1381:G:H8	1.82	0.44
34:RA:1565:C:O2'	34:RA:1567:A:N7	2.41	0.44
40:RH:4:ILE:HD12	40:RH:6:ARG:HB2	1.99	0.44
1:XA:563:A:O2'	1:XA:566:G:O3'	2.34	0.44
1:XA:657:G:H2'	1:XA:658:G:H8	1.82	0.44
1:XA:896:C:H2'	1:XA:897:C:C6	2.52	0.44
1:XA:936:C:H3'	1:XA:936:C:H6	1.82	0.44
1:XA:951:G:O3'	1:XA:972:C:N4	2.50	0.44
1:XA:1065:U:C5	1:XA:1190:G:H1'	2.53	0.44
1:XA:1279:A:OP2	10:XJ:9:ARG:NH2	2.49	0.44
1:XA:1317:C:H3'	1:XA:1318:A:H8	1.83	0.44
2:XB:33:TYR:HB2	2:XB:43:ASP:HB2	1.99	0.44
11:XK:109:VAL:CB	18:XR:86:VAL:HG23	2.47	0.44
16:XP:19:ILE:HG13	16:XP:36:ILE:HD11	2.00	0.44
27:Y3:31:LEU:HD12	34:YA:1157:G:O2'	2.16	0.44
34:YA:807:U:H2'	34:YA:808:G:H8	1.82	0.44
34:YA:887:A:H1'	34:YA:889:C:C6	2.53	0.44
34:YA:1494:A:H2	34:YA:1579:A:H1'	1.83	0.44
34:YA:2108:C:O5'	34:YA:2108:C:H6	1.99	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:2306:C:H3'	34:YA:2307:G:H5''	2.00	0.44
34:YA:2515:C:H2'	34:YA:2516:G:C8	2.51	0.44
34:YA:2587:A:H8	34:YA:2587:A:O5'	2.00	0.44
53:YY:28:LYS:N	53:YY:38:ILE:O	2.45	0.44
1:QA:403:C:H4'	4:QD:122:ARG:NE	2.33	0.44
1:QA:413:G:H4'	1:QA:414:A:H5''	1.99	0.44
1:QA:669:U:O4'	15:QO:46:HIS:CE1	2.69	0.44
1:QA:1265:G:H2'	1:QA:1266:G:C8	2.53	0.44
1:QA:1307:U:C5'	13:QM:99:ARG:CZ	2.96	0.44
1:QA:1411:C:H2'	1:QA:1412:C:C6	2.52	0.44
8:QH:82:HIS:N	8:QH:138:TRP:OXT	2.41	0.44
34:RA:144:C:H2'	34:RA:145:G:H8	1.83	0.44
34:RA:624:C:H2'	34:RA:625:G:C8	2.53	0.44
34:RA:822:U:C5	34:RA:944:G:H1'	2.52	0.44
34:RA:1251:C:OP2	49:RU:10:ARG:NH1	2.50	0.44
34:RA:1445:C:H2'	34:RA:1446:C:C6	2.51	0.44
34:RA:2290:G:N2	34:RA:2343:C:H1'	2.33	0.44
34:RA:2320:A:H1'	34:RA:2321:G:C6	2.52	0.44
34:RA:2327:A:H2'	34:RA:2328:A:C8	2.53	0.44
34:RA:2351:G:H1'	34:RA:2367:G:N2	2.32	0.44
34:RA:2619:C:H2'	34:RA:2620:C:C6	2.51	0.44
39:RG:135:LEU:HD11	39:RG:140:ILE:HD11	2.00	0.44
1:XA:130:A:H1'	1:XA:263:A:O2'	2.17	0.44
1:XA:193:C:H2'	1:XA:194:C:C6	2.52	0.44
1:XA:833:U:H2'	1:XA:834:C:C6	2.53	0.44
1:XA:907:A:H2'	1:XA:908:A:O4'	2.18	0.44
1:XA:1028(D):G:N2	1:XA:1028(H):G:N7	2.65	0.44
1:XA:1094:G:H4'	1:XA:1095:U:H5	1.83	0.44
1:XA:1240:U:H3	7:XG:38:LEU:HD22	1.60	0.44
1:XA:1318:A:H1'	19:XS:37:ARG:HH22	1.83	0.44
1:XA:1329:A:H4'	13:XM:29:ARG:HH21	1.82	0.44
1:XA:1429:C:H2'	1:XA:1430:C:C6	2.53	0.44
29:Y5:17:ASP:HB3	34:YA:16:G:OP1	2.18	0.44
34:YA:210:C:H4'	34:YA:1367:A:H1'	2.00	0.44
34:YA:524:U:H2'	34:YA:525:U:C6	2.53	0.44
34:YA:1162:G:H2'	34:YA:1163:G:C8	2.53	0.44
34:YA:1468:C:H2'	34:YA:1469:A:H8	1.82	0.44
34:YA:1574:C:H2'	34:YA:1575:C:C6	2.53	0.44
34:YA:1991:U:H2'	34:YA:1992:G:H5''	2.00	0.44
34:YA:2099:U:C4	34:YA:2190:G:N1	2.82	0.44
34:YA:2124:G:H3'	34:YA:2125:G:H8	1.82	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:2246:G:H2'	34:YA:2247:A:C8	2.53	0.44
34:YA:2286:A:H1'	34:YA:2287:A:C6	2.52	0.44
34:YA:2342:C:H2'	34:YA:2374:C:H5''	2.00	0.44
38:YF:102:PRO:HB2	38:YF:105:VAL:HG23	1.99	0.44
46:YR:29:LEU:HD13	46:YR:79:LEU:HD22	2.00	0.44
1:QA:191:G:O2'	20:QT:103:GLY:N	2.45	0.44
1:QA:245:C:H1'	1:QA:284:G:C2	2.53	0.44
1:QA:986:A:H4'	19:QS:55:LYS:HG2	1.94	0.44
1:QA:1092:A:C2	1:QA:1110:A:H5'	2.52	0.44
1:QA:1108:G:OP2	3:QC:174:PRO:HA	2.18	0.44
1:QA:1114:C:H2'	1:QA:1115:C:H6	1.83	0.44
1:QA:1332:A:H3'	1:QA:1333:A:H8	1.83	0.44
1:QA:1463:C:H2'	1:QA:1464:G:H8	1.83	0.44
1:QA:1479:C:H2'	1:QA:1480:G:H8	1.82	0.44
1:QA:1515:C:H2'	1:QA:1516:G:C8	2.53	0.44
3:QC:23:TYR:N	10:QJ:93:GLY:O	2.51	0.44
6:QF:14:LEU:HD12	6:QF:18:GLN:CD	2.38	0.44
7:QG:116:ALA:HA	7:QG:119:ARG:HE	1.82	0.44
31:R7:34:ARG:NH1	34:RA:466:A:OP1	2.50	0.44
34:RA:1104:C:H2'	34:RA:1105:U:H6	1.82	0.44
34:RA:1923:U:H2'	34:RA:1924:C:C6	2.52	0.44
34:RA:2195:C:H2'	34:RA:2196:C:C6	2.53	0.44
51:RW:71:VAL:HA	51:RW:107:LEU:HD23	2.00	0.44
54:RZ:156:LYS:HE3	54:RZ:156:LYS:HB3	1.74	0.44
1:XA:186(L):G:H2'	1:XA:186(M):G:H8	1.82	0.44
1:XA:407:G:H1'	4:XD:119:GLN:HE22	1.82	0.44
1:XA:643:C:H2'	1:XA:644:G:C8	2.52	0.44
1:XA:687:A:O2'	1:XA:701:C:N4	2.50	0.44
1:XA:962:C:H2'	1:XA:963:G:C8	2.51	0.44
1:XA:974:A:P	14:XN:29:ARG:NE	2.90	0.44
1:XA:986:A:H1'	19:XS:52:TYR:HH	1.80	0.44
1:XA:1240:U:C2'	7:XG:38:LEU:HD12	2.45	0.44
14:XN:40:CYS:SG	14:XN:41:ARG:N	2.91	0.44
16:XP:22:THR:OG1	16:XP:23:ASP:N	2.50	0.44
18:XR:47:THR:O	18:XR:83:GLU:N	2.51	0.44
25:Y1:2:SER:HB3	34:YA:1366:A:OP1	2.18	0.44
34:YA:417:C:O2	34:YA:2407:G:C6	2.71	0.44
34:YA:625:G:H2'	34:YA:626:U:C6	2.53	0.44
34:YA:1242:A:H3'	34:YA:1243:G:H8	1.83	0.44
34:YA:1304:C:H2'	34:YA:1305:C:H6	1.83	0.44
34:YA:1353:A:H2'	34:YA:1354:A:C8	2.53	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:1362:C:H2'	34:YA:1363:C:H6	1.83	0.44
34:YA:1834:U:H1'	34:YA:1969:A:H2'	1.99	0.44
34:YA:2307:G:O6	39:YG:42:GLY:O	2.35	0.44
34:YA:2356:C:H1'	34:YA:2362:G:N2	2.33	0.44
34:YA:2447:G:H1	34:YA:2451:A:N6	2.16	0.44
34:YA:2506:U:H2'	34:YA:2507:C:C6	2.51	0.44
34:YA:2619:C:H2'	34:YA:2620:C:C6	2.53	0.44
34:YA:2747:G:H1	34:YA:2754:U:H2'	1.83	0.44
44:YP:21:ARG:HB3	44:YP:22:GLY:H	1.67	0.44
1:QA:181:G:H4'	1:QA:182:U:H5'	1.99	0.44
1:QA:373:A:N1	1:QA:391:G:O2'	2.48	0.44
1:QA:624:C:C4'	16:QP:11:SER:HB3	2.48	0.44
1:QA:697:U:H3'	1:QA:698:G:H8	1.83	0.44
1:QA:863:U:HO2'	1:QA:865:A:H62	1.62	0.44
1:QA:878:G:H5'	8:QH:89:PRO:HG2	1.99	0.44
1:QA:981:U:H5'	14:QN:6:LEU:HD22	1.99	0.44
1:QA:1124:G:H2'	1:QA:1145:C:N3	2.32	0.44
1:QA:1306:A:H1'	1:QA:1332:A:C8	2.53	0.44
1:QA:1358:U:H3'	14:QN:35:ARG:HD2	1.99	0.44
1:QA:1500:A:H5''	1:QA:1508:G:H5''	1.98	0.44
1:QA:1512:U:H2'	1:QA:1513:A:C8	2.53	0.44
34:RA:448:U:C4	34:RA:583:G:H1'	2.52	0.44
34:RA:661:C:H2'	34:RA:662:G:C8	2.53	0.44
34:RA:1316:U:H2'	34:RA:1317:A:C8	2.53	0.44
34:RA:1794:U:H2'	34:RA:1795:C:H6	1.83	0.44
34:RA:2246:G:H2'	34:RA:2247:A:C8	2.53	0.44
36:RD:181:GLU:HA	36:RD:272:ALA:HB3	1.99	0.44
49:RU:91:ASP:O	49:RU:93:LYS:N	2.50	0.44
52:RX:55:ASN:HB2	52:RX:80:ILE:HG23	2.00	0.44
1:XA:150:C:H2'	1:XA:151:A:O4'	2.18	0.44
1:XA:152:A:N6	1:XA:169:C:N3	2.66	0.44
1:XA:385:C:H2'	1:XA:386:C:C6	2.52	0.44
1:XA:817:C:O2'	1:XA:1527:C:O3'	2.35	0.44
1:XA:986:A:H2'	1:XA:987:G:C8	2.53	0.44
1:XA:1106:G:C4'	3:XC:172:ARG:HG2	2.46	0.44
1:XA:1300:G:C6	1:XA:1334:G:N7	2.86	0.44
1:XA:1440(F):G:H22	1:XA:1440(P):A:H1'	1.83	0.44
10:XJ:16:LEU:CB	10:XJ:70:ARG:NH1	2.61	0.44
13:XM:2:ALA:HB3	13:XM:9:ILE:HG21	2.00	0.44
24:Y0:38:VAL:HG22	24:Y0:59:LEU:HB2	1.98	0.44
32:Y8:7:HIS:HD2	44:YP:50:ARG:HH21	1.64	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:662:G:H2'	34:YA:663:G:H8	1.81	0.44
34:YA:864:G:H1'	34:YA:914:C:H42	1.82	0.44
34:YA:2090:G:C4	34:YA:2230:G:N1	2.86	0.44
34:YA:2550:G:H2'	34:YA:2551:C:H6	1.83	0.44
34:YA:2630:G:H2'	34:YA:2631:G:H8	1.83	0.44
43:YO:44:LYS:HA	43:YO:44:LYS:HD3	1.72	0.44
51:YW:35:ILE:O	51:YW:39:THR:OG1	2.27	0.44
1:QA:231:G:H2'	1:QA:232:G:C8	2.52	0.43
1:QA:292:G:C5	1:QA:293:G:H1'	2.53	0.43
1:QA:324:G:OP1	20:QT:22:ARG:HD2	2.18	0.43
1:QA:765:G:N1	1:QA:812:C:O2'	2.44	0.43
1:QA:975:A:O4'	1:QA:1365:G:N2	2.51	0.43
1:QA:1321:C:H5''	1:QA:1322:C:H5''	1.99	0.43
27:R3:45:GLY:HA3	34:RA:851:U:O3'	2.18	0.43
33:R9:23:VAL:HG22	34:RA:1032:A:O2'	2.17	0.43
34:RA:97:C:H2'	34:RA:98:G:C8	2.53	0.43
34:RA:714:U:H1'	34:RA:717:G:N7	2.33	0.43
34:RA:1400:G:H2'	34:RA:1401:G:C8	2.53	0.43
34:RA:2771:C:H2'	34:RA:2772:C:C6	2.52	0.43
34:RA:2810:A:H62	34:RA:2891:G:H21	1.66	0.43
35:RB:9:G:OP1	47:RS:15:ARG:NH1	2.51	0.43
41:RI:79:ILE:HD12	41:RI:80:PRO:HD2	2.00	0.43
1:XA:51:A:N9	1:XA:353:A:C5	2.86	0.43
1:XA:57:G:H2'	1:XA:58:C:C6	2.53	0.43
1:XA:411:A:H2'	1:XA:413:G:C8	2.53	0.43
1:XA:643:C:H2'	1:XA:644:G:H8	1.82	0.43
1:XA:658:G:H1'	15:XO:22:THR:HG23	1.99	0.43
1:XA:878:G:H5'	8:XH:89:PRO:HG2	2.00	0.43
1:XA:1120:G:H2'	1:XA:1121:U:C6	2.53	0.43
1:XA:1230:C:C4	13:XM:105:THR:HB	2.49	0.43
11:XK:110:ASP:O	18:XR:85:LEU:N	2.51	0.43
20:XT:100:ILE:HG23	20:XT:102:GLY:H	1.81	0.43
30:Y6:3:SER:HB3	30:Y6:6:ARG:HB3	2.00	0.43
30:Y6:29:ASN:ND2	34:YA:2286:A:OP1	2.51	0.43
34:YA:274:G:H2'	34:YA:275:G:H8	1.83	0.43
34:YA:519:U:H2'	34:YA:520:G:C8	2.53	0.43
34:YA:1467:C:C5	34:YA:1546:C:H2'	2.53	0.43
34:YA:1710:C:H2'	34:YA:1711:C:C6	2.53	0.43
34:YA:1908:C:H2'	34:YA:1909:C:H6	1.83	0.43
34:YA:1953:A:C2	34:YA:2550:G:C1'	3.01	0.43
34:YA:2853:C:H2'	34:YA:2854:G:C8	2.53	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
36:YD:142:VAL:HA	36:YD:194:GLY:H	1.83	0.43
37:YE:102:VAL:N	37:YE:170:LEU:O	2.46	0.43
1:QA:444:C:H2'	1:QA:445:G:C8	2.53	0.43
1:QA:608:A:C1'	16:QP:32:TYR:CE1	2.99	0.43
1:QA:1327:C:H4'	21:QU:20:LYS:NZ	2.33	0.43
3:QC:112:SER:HB3	3:QC:115:LEU:HD12	2.00	0.43
3:QC:131:ARG:HH11	3:QC:134:ILE:HG21	1.83	0.43
4:QD:61:LYS:HD2	4:QD:207:TYR:CZ	2.52	0.43
25:R1:90:ILE:HA	25:R1:94:LEU:HG	1.99	0.43
28:R4:12:ALA:H	28:R4:25:TYR:HA	1.83	0.43
34:RA:452:G:N2	34:RA:457:A:O2'	2.51	0.43
34:RA:655:A:H8	34:RA:656:G:C8	2.36	0.43
34:RA:1046:A:H5''	34:RA:1046:A:N3	2.34	0.43
34:RA:1295:C:H2'	34:RA:1296:G:C8	2.49	0.43
34:RA:1826:G:H4'	36:RD:242:ARG:NH2	2.32	0.43
34:RA:2092:U:OP2	41:RI:27:ARG:NH2	2.50	0.43
34:RA:2182:G:H2'	34:RA:2183:C:C6	2.53	0.43
34:RA:2463:C:H2'	34:RA:2464:C:H6	1.82	0.43
36:RD:50:THR:OG1	36:RD:51:VAL:N	2.50	0.43
45:RQ:34:LEU:HB2	45:RQ:118:LEU:HD12	2.00	0.43
1:XA:9:G:OP1	5:XE:122:GLU:HB2	2.17	0.43
1:XA:158:G:H2'	1:XA:159:G:C8	2.53	0.43
1:XA:936:C:H3'	1:XA:936:C:C6	2.52	0.43
1:XA:1059:C:C6	3:XC:2:GLY:HA2	2.51	0.43
1:XA:1271:G:H2'	1:XA:1272:G:C8	2.53	0.43
7:XG:116:ALA:HA	7:XG:119:ARG:HE	1.83	0.43
10:XJ:7:LYS:HD2	10:XJ:97:GLU:HB2	1.98	0.43
12:XL:33:ARG:NH2	12:XL:61:THR:OG1	2.51	0.43
34:YA:462:C:H2'	34:YA:463:G:C8	2.52	0.43
34:YA:1509:C:H3'	34:YA:1510:A:H5''	2.01	0.43
34:YA:1927:A:H2'	34:YA:1928:A:C8	2.53	0.43
34:YA:1958:C:H2'	34:YA:1959:G:C8	2.53	0.43
34:YA:2002:G:H2'	34:YA:2003:G:C8	2.54	0.43
34:YA:2131:G:H5''	34:YA:2131:G:H8	1.82	0.43
34:YA:2692:C:O2	34:YA:2847:U:O2'	2.26	0.43
34:YA:2751:G:O6	40:YH:2:SER:OG	2.35	0.43
41:YI:101:LEU:HD11	41:YI:109:ILE:HD13	2.00	0.43
1:QA:222:U:H2'	1:QA:223:U:C6	2.53	0.43
1:QA:311:C:H2'	1:QA:312:C:H6	1.83	0.43
1:QA:435:C:H2'	1:QA:436:C:H6	1.83	0.43
1:QA:563:A:HO2'	1:QA:566:G:HO2'	1.59	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:718:G:H1'	11:QK:116:HIS:HA	2.00	0.43
1:QA:897:C:H2'	1:QA:898:G:C8	2.53	0.43
3:QC:79:ARG:CZ	11:XK:104:GLN:HG3	2.41	0.43
5:QE:43:LEU:HD13	5:QE:109:ILE:HD11	1.99	0.43
5:QE:100:VAL:O	5:QE:107:ARG:NH1	2.48	0.43
29:R5:12:SER:O	29:R5:16:ARG:CB	2.67	0.43
34:RA:619:G:H3'	34:RA:620:G:N2	2.33	0.43
34:RA:865:C:O2	34:RA:867:C:N4	2.51	0.43
34:RA:971:C:H2'	34:RA:972:G:O4'	2.18	0.43
34:RA:1141:U:H1'	34:RA:1142(B):A:C6	2.53	0.43
34:RA:1211:U:H5''	34:RA:1212:G:N7	2.34	0.43
34:RA:1992:G:H5'	34:RA:1994:C:H41	1.83	0.43
34:RA:2250:G:C2	45:RQ:82:ARG:HB3	2.52	0.43
34:RA:2263:C:H2'	34:RA:2264:C:C6	2.54	0.43
49:RU:92:ARG:HD2	50:RV:11:GLN:HG3	1.99	0.43
1:XA:736:C:H2'	1:XA:737:A:H8	1.83	0.43
1:XA:743:U:H2'	1:XA:744:C:C6	2.53	0.43
1:XA:960:U:H1'	1:XA:1222:G:O2'	2.18	0.43
1:XA:1414:U:H2'	1:XA:1415:G:C8	2.53	0.43
14:YN:13:THR:O	14:YN:13:THR:OG1	2.36	0.43
34:YA:143:C:H2'	34:YA:144:C:C6	2.53	0.43
34:YA:242:G:N2	34:YA:254:G:H2'	2.33	0.43
34:YA:1230:C:H2'	34:YA:1231:G:C8	2.53	0.43
34:YA:1311:G:H21	34:YA:1603:A:H62	1.67	0.43
34:YA:1375:C:H2'	34:YA:1376:C:H6	1.83	0.43
34:YA:1504:C:H2'	34:YA:1505:C:C6	2.53	0.43
34:YA:1564:C:H2'	34:YA:1565:C:C6	2.53	0.43
34:YA:1930:G:H2'	34:YA:1968:G:N1	2.34	0.43
34:YA:1992:G:N2	34:YA:1996:C:O2'	2.52	0.43
34:YA:2471:C:H3'	34:YA:2472:G:H8	1.83	0.43
34:YA:2751:G:C5	40:YH:2:SER:O	2.71	0.43
37:YE:37:ARG:NH1	37:YE:44:TYR:OH	2.48	0.43
39:YG:124:SER:O	39:YG:124:SER:OG	2.35	0.43
39:YG:165:THR:OG1	39:YG:166:ASP:N	2.52	0.43
41:YI:81:VAL:HG11	41:YI:88:ILE:HD13	1.99	0.43
1:QA:24:U:H2'	1:QA:25:C:H6	1.83	0.43
1:QA:36:C:H2'	1:QA:37:U:O4'	2.18	0.43
1:QA:645:C:H2'	1:QA:646:U:C6	2.53	0.43
1:QA:1109:C:OP2	3:QC:176:HIS:CD2	2.71	0.43
1:QA:1247:U:H2'	1:QA:1248:A:C8	2.53	0.43
1:QA:1357:A:H2	1:QA:1365:G:H22	1.67	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1383:C:H2'	1:QA:1384:C:H6	1.82	0.43
3:QC:6:HIS:HE1	3:QC:8:ILE:HD12	1.82	0.43
34:RA:81:G:O2'	34:RA:295:G:O2'	2.23	0.43
34:RA:300:A:OP1	53:RY:86:ARG:NH2	2.47	0.43
34:RA:532:A:H4'	34:RA:533:G:C8	2.52	0.43
34:RA:779:U:H2'	34:RA:780:G:C8	2.53	0.43
34:RA:816:C:H2'	34:RA:817:C:C6	2.54	0.43
34:RA:1130:U:N3	34:RA:2025:C:H5''	2.33	0.43
34:RA:1145:C:H2'	34:RA:1146:C:C6	2.54	0.43
34:RA:1818:U:OP2	36:RD:157:ARG:NH1	2.51	0.43
34:RA:2122:U:H2'	34:RA:2123:G:C8	2.53	0.43
34:RA:2341:G:H2'	34:RA:2342:C:C6	2.53	0.43
35:RB:111:U:H2'	35:RB:112:G:H8	1.82	0.43
37:RE:171:GLU:H	37:RE:185:LYS:HB2	1.84	0.43
37:RE:176:ILE:HG13	37:RE:181:LEU:HB2	2.00	0.43
39:RG:43:LEU:HD21	39:RG:153:ARG:HB2	1.99	0.43
41:RI:84:GLY:CA	41:RI:89:TYR:OH	2.61	0.43
43:RO:15:GLY:HA3	43:RO:50:GLY:HA3	1.99	0.43
46:RR:29:LEU:HD12	46:RR:70:LEU:HD21	1.99	0.43
1:XA:355:C:H1'	1:XA:388:G:H1'	2.00	0.43
1:XA:520:A:N1	1:XA:533:A:N6	2.55	0.43
1:XA:726:C:H2'	1:XA:727:G:C8	2.53	0.43
1:XA:1308:U:H2'	1:XA:1309:G:C8	2.53	0.43
1:XA:1347:G:H4'	1:XA:1348:U:C5	2.53	0.43
22:XV:20:G:N2	22:XV:60:U:H3	2.17	0.43
33:Y9:33:LYS:NZ	34:YA:2743:C:OP1	2.43	0.43
34:YA:208:C:H2'	34:YA:209:C:C6	2.53	0.43
34:YA:270(I):C:H2'	34:YA:270(J):G:C8	2.53	0.43
34:YA:270(J):G:H2'	34:YA:270(K):G:C8	2.54	0.43
34:YA:1097:U:H3'	34:YA:1098:A:H8	1.83	0.43
34:YA:1116:C:H2'	34:YA:1117:G:C8	2.54	0.43
34:YA:1539:G:H2'	34:YA:1540:G:H8	1.84	0.43
34:YA:1782:C:H41	34:YA:2587:A:H2	1.62	0.43
34:YA:2179:C:H2'	34:YA:2180:U:C6	2.53	0.43
36:YD:143:HIS:ND1	36:YD:194:GLY:O	2.40	0.43
39:YG:106:LEU:HD12	39:YG:110:ALA:HB3	1.99	0.43
48:YT:19:LEU:HD21	48:YT:83:ILE:HD11	2.00	0.43
49:YU:55:ARG:O	49:YU:59:ARG:HG2	2.18	0.43
1:QA:218:C:H2'	1:QA:219:C:C6	2.54	0.43
1:QA:378:G:H2'	1:QA:379:C:C6	2.52	0.43
1:QA:702:A:H3'	1:QA:703:G:C8	2.53	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:831:U:H2'	1:QA:832:C:C6	2.54	0.43
1:QA:1014:A:H1'	19:QS:34:TRP:CD2	2.54	0.43
1:QA:1112:C:N3	3:QC:178:LEU:HD12	2.33	0.43
1:QA:1255:G:O6	1:QA:1279:A:H2'	2.18	0.43
1:QA:1404:C:H2'	1:QA:1405:G:C8	2.54	0.43
10:QJ:28:ARG:HE	10:QJ:34:VAL:HG12	1.83	0.43
34:RA:32:C:N4	34:RA:447:A:OP2	2.51	0.43
34:RA:873:G:H4'	45:RQ:63:LYS:NZ	2.32	0.43
34:RA:1382:G:H4'	34:RA:1573:G:C2	2.54	0.43
34:RA:1476:C:H2'	34:RA:1477:A:H8	1.83	0.43
34:RA:1796:U:H2'	34:RA:1797:C:H6	1.82	0.43
34:RA:2737:G:H2'	34:RA:2738:A:C8	2.53	0.43
34:RA:2747:G:H21	34:RA:2757:A:H62	1.65	0.43
40:RH:80:SER:O	40:RH:80:SER:OG	2.31	0.43
46:RR:35:THR:HG22	46:RR:113:LEU:HD13	2.01	0.43
1:XA:337:C:H2'	1:XA:338:A:C8	2.53	0.43
1:XA:559:A:H4'	1:XA:560:U:H5''	1.99	0.43
1:XA:1301:U:C3'	13:XM:17:VAL:HG23	2.49	0.43
1:XA:1302:U:H1'	13:XM:27:LYS:HG3	1.83	0.43
1:XA:1524:C:H2'	1:XA:1525:G:C8	2.53	0.43
5:XE:99:GLY:N	5:XE:117:ASP:OD1	2.47	0.43
9:XI:70:LYS:O	9:XI:74:ILE:HG13	2.19	0.43
27:Y3:18:ASP:HB2	27:Y3:49:LYS:HE2	1.99	0.43
28:Y4:37:SER:HA	28:Y4:41:PRO:HD2	1.99	0.43
32:Y8:18:ALA:HB2	34:YA:628:G:H5''	2.01	0.43
34:YA:358:U:H2'	34:YA:359:A:C8	2.51	0.43
34:YA:361:G:H2'	34:YA:362:U:C6	2.54	0.43
34:YA:557:U:H2'	34:YA:558:G:C8	2.53	0.43
34:YA:786:C:H2'	34:YA:787:U:H6	1.83	0.43
34:YA:994:C:OP1	49:YU:53:ARG:NH2	2.51	0.43
34:YA:1161:C:H2'	34:YA:1162:G:C8	2.53	0.43
34:YA:1189:A:C2	34:YA:1190:G:H1'	2.54	0.43
34:YA:2046:G:C6	34:YA:2623:G:O6	2.68	0.43
34:YA:2170:A:H2'	34:YA:2171:A:O4'	2.19	0.43
34:YA:2432:A:H2'	34:YA:2433:A:C8	2.53	0.43
34:YA:2758:A:C2	34:YA:2759:G:H1'	2.53	0.43
44:YP:97:PRO:O	44:YP:98:GLU:HG3	2.18	0.43
1:QA:130:A:H4'	1:QA:186(K):G:C4	2.53	0.43
1:QA:1004:A:C2	1:QA:1024:G:H2'	2.51	0.43
1:QA:1440(I):U:H4'	1:QA:1440(J):A:C2	2.53	0.43
1:QA:1497:G:H1'	1:QA:1518:A:C2	2.54	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:QB:188:ALA:HB1	2:QB:192:SER:HB2	2.01	0.43
3:QC:24:ALA:HB1	3:QC:32:LEU:HD21	1.99	0.43
24:R0:14:ARG:O	34:RA:2278:A:N6	2.52	0.43
29:R5:16:ARG:NH2	34:RA:517:C:OP1	2.51	0.43
34:RA:300:A:OP2	53:RY:84:ARG:NH1	2.52	0.43
34:RA:1310:G:H1'	34:RA:1611:C:H5''	2.00	0.43
34:RA:1732:A:H3'	34:RA:1733:G:C8	2.53	0.43
36:RD:171:ASP:OD1	36:RD:171:ASP:N	2.48	0.43
38:RF:34:TRP:CE3	44:RP:8:PRO:HB3	2.53	0.43
44:RP:96:THR:HA	44:RP:126:VAL:HB	2.01	0.43
48:RT:6:LEU:O	48:RT:10:VAL:HG23	2.19	0.43
1:XA:404:U:H2'	1:XA:405:U:C6	2.53	0.43
1:XA:1440(D):G:H5'	1:XA:1440(E):A:H2	1.83	0.43
1:XA:1463:C:H2'	1:XA:1464:G:C8	2.53	0.43
5:XE:31:LEU:HD13	5:XE:45:PHE:HD1	1.82	0.43
12:XL:86:ARG:HH21	12:XL:99:HIS:CD2	2.37	0.43
22:XV:76:A:C6	34:YA:2421:G:H2'	2.54	0.43
34:YA:273(E):C:H2'	34:YA:273(F):U:C6	2.54	0.43
34:YA:296:C:O3'	53:YY:95:LYS:NZ	2.51	0.43
34:YA:416:C:N3	34:YA:2407:G:O6	2.51	0.43
34:YA:1141:U:H1'	34:YA:1142(B):A:C6	2.54	0.43
34:YA:1339:G:N2	34:YA:1603:A:N3	2.66	0.43
34:YA:1689:A:OP2	34:YA:1698:A:N6	2.52	0.43
34:YA:1796:U:H2'	34:YA:1797:C:H6	1.84	0.43
34:YA:1947:C:H2'	34:YA:1948:G:C8	2.54	0.43
34:YA:2110:G:H3'	34:YA:2111:C:C6	2.53	0.43
44:YP:64:LYS:O	44:YP:66:GLY:N	2.51	0.43
1:QA:25:C:H2'	1:QA:26:A:C8	2.54	0.43
1:QA:60:A:H2	1:QA:378:G:H1'	1.84	0.43
1:QA:296:U:H2'	1:QA:297:G:C8	2.54	0.43
1:QA:662:G:H2'	1:QA:663:A:H8	1.82	0.43
1:QA:689:C:P	11:QK:55:LYS:NZ	2.92	0.43
1:QA:740:U:O2'	15:QO:39:LEU:HD11	2.18	0.43
1:QA:918:A:H2'	1:QA:919:A:C8	2.54	0.43
1:QA:948:C:OP2	13:QM:106:ASN:O	2.36	0.43
1:QA:986:A:C2'	19:QS:55:LYS:HA	2.46	0.43
1:QA:1190:G:H4'	3:QC:176:HIS:CE1	2.54	0.43
2:QB:54:THR:O	2:QB:58:ILE:HG12	2.18	0.43
4:QD:191:ARG:HD2	4:QD:191:ARG:HA	1.72	0.43
7:QG:6:ARG:HA	7:QG:6:ARG:HD3	1.75	0.43
15:QO:5:LYS:HE2	15:QO:5:LYS:HB2	1.90	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
19:QS:39:THR:HG22	19:QS:41:VAL:HG22	2.01	0.43
32:R8:31:HIS:HE1	34:RA:2421:G:N7	2.14	0.43
34:RA:39:C:H2'	34:RA:40:C:C6	2.54	0.43
34:RA:134:C:H2'	34:RA:135:G:C8	2.54	0.43
34:RA:206:U:H2'	34:RA:207:A:C8	2.50	0.43
34:RA:807:U:O2'	34:RA:2060:A:N1	2.48	0.43
34:RA:864:G:H1'	34:RA:914:C:H42	1.83	0.43
34:RA:967:C:H2'	34:RA:968:G:C8	2.54	0.43
34:RA:1467:C:C5	34:RA:1546:C:H2'	2.54	0.43
34:RA:1952:A:N3	34:RA:2560:C:O2'	2.41	0.43
34:RA:2736:G:H2'	34:RA:2737:G:C8	2.54	0.43
49:RU:88:ILE:HD12	50:RV:47:VAL:HG22	2.01	0.43
53:RY:28:LYS:NZ	53:RY:40:GLU:OE2	2.38	0.43
54:RZ:48:PHE:HA	54:RZ:51:ALA:HB3	2.01	0.43
1:XA:186(F):C:H2'	1:XA:186(G):C:C6	2.53	0.43
1:XA:272:C:H2'	1:XA:273:A:C8	2.53	0.43
1:XA:1028(C):C:N3	1:XA:1028(H):G:N1	2.56	0.43
1:XA:1071:C:H2'	1:XA:1072:G:C8	2.54	0.43
21:XU:3:LYS:HG2	21:XU:14:TRP:CD2	2.53	0.43
27:Y3:5:LYS:HE3	27:Y3:57:GLU:HB3	1.99	0.43
34:YA:263:C:H1'	34:YA:430:G:H1'	2.00	0.43
34:YA:270(N):U:H1'	34:YA:270(O):G:C5	2.53	0.43
34:YA:296:C:H2'	34:YA:297:C:H6	1.83	0.43
34:YA:363(A):G:H2'	34:YA:363(B):A:H8	1.83	0.43
34:YA:1501:C:H2'	34:YA:1502:C:C6	2.54	0.43
34:YA:1539:G:H2'	34:YA:1540:G:C8	2.53	0.43
34:YA:1674:G:H1'	34:YA:1676:A:N6	2.33	0.43
34:YA:1800:C:OP2	36:YD:183:ARG:NH1	2.52	0.43
34:YA:2023:G:P	34:YA:2023:G:H8	2.42	0.43
34:YA:2064:C:H2'	34:YA:2065:C:C6	2.54	0.43
34:YA:2391:G:H1'	34:YA:2429:G:N2	2.33	0.43
40:YH:152:ARG:HA	40:YH:152:ARG:HD3	1.80	0.43
53:YY:76:CYS:CB	53:YY:79:CYS:SG	3.07	0.43
1:QA:285:G:H2'	1:QA:286:G:C8	2.54	0.43
1:QA:831:U:H2'	1:QA:832:C:H6	1.84	0.43
1:QA:975:A:H3'	14:QN:32:SER:HB3	2.01	0.43
1:QA:1107:C:O2'	1:QA:1191:A:O2'	2.29	0.43
1:QA:1123:A:H2'	1:QA:1124:G:C4	2.53	0.43
1:QA:1260:C:HO2'	1:QA:1283:G:HO2'	1.63	0.43
2:QB:150:SER:HA	2:QB:153:ARG:HH21	1.83	0.43
5:QE:80:ILE:HD12	8:QH:104:ARG:HH12	1.82	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
13:QM:34:LEU:HA	13:QM:37:THR:HG22	2.01	0.43
13:QM:84:ILE:HG13	19:QS:74:PHE:CE2	2.54	0.43
13:QM:95:GLY:HA2	13:QM:110:ARG:HH21	1.83	0.43
26:R2:16:LEU:H	26:R2:67:LYS:NZ	2.16	0.43
34:RA:29:U:H5''	49:RU:7:GLY:HA2	2.00	0.43
34:RA:210:C:H4'	34:RA:1367:A:H1'	2.00	0.43
34:RA:231:C:H3'	34:RA:232:G:C8	2.53	0.43
34:RA:259:G:H2'	34:RA:260:G:H8	1.84	0.43
34:RA:285:C:H2'	34:RA:286:C:C6	2.52	0.43
34:RA:624:C:H2'	34:RA:625:G:H8	1.83	0.43
34:RA:1258:C:H2'	34:RA:1259:G:C8	2.54	0.43
34:RA:1518:C:H2'	34:RA:1519:G:H8	1.83	0.43
34:RA:1604:C:H2'	34:RA:1605:C:C6	2.50	0.43
34:RA:1629:U:H2'	34:RA:1630(A):G:H8	1.83	0.43
34:RA:2391:G:C6	34:RA:2427:C:H1'	2.54	0.43
34:RA:2517:C:N3	34:RA:2542:A:N6	2.65	0.43
34:RA:2607:G:H2'	34:RA:2608:G:C8	2.54	0.43
34:RA:2737:G:H2'	34:RA:2738:A:H8	1.84	0.43
38:RF:152:GLU:OE1	38:RF:191:ARG:NE	2.52	0.43
53:RY:79:CYS:N	53:RY:102:CYS:SG	2.92	0.43
1:XA:186(A):C:H2'	1:XA:186(B):C:C6	2.54	0.43
1:XA:256:U:H2'	1:XA:257:G:C8	2.54	0.43
1:XA:555:C:H2'	1:XA:556:C:C6	2.53	0.43
1:XA:921:U:H2'	1:XA:922:G:C8	2.53	0.43
1:XA:936:C:C6	1:XA:936:C:C3'	3.01	0.43
1:XA:950:U:H3	1:XA:1231:G:H1	1.65	0.43
1:XA:1240:U:H5''	1:XA:1241:G:C8	2.54	0.43
1:XA:1256:A:N6	1:XA:1277:C:H5''	2.33	0.43
1:XA:1485:U:H2'	1:XA:1486:G:H8	1.82	0.43
6:XF:62:TRP:CD1	18:XR:35:ARG:NH2	2.87	0.43
11:XK:82:VAL:HB	11:XK:108:ILE:HA	2.01	0.43
12:XL:11:VAL:CG2	17:XQ:29:HIS:CD2	3.00	0.43
34:YA:141(A):A:H1'	34:YA:1408:C:H1'	2.01	0.43
34:YA:161:U:H3'	34:YA:162:U:H5''	2.00	0.43
34:YA:679:C:H2'	34:YA:680:G:C8	2.53	0.43
34:YA:1181:C:H2'	34:YA:1182:A:C8	2.54	0.43
34:YA:1662:C:H2'	34:YA:1663:C:C6	2.53	0.43
34:YA:2044:C:C2	34:YA:2625:G:N2	2.87	0.43
34:YA:2346:A:N3	34:YA:2346:A:H5''	2.34	0.43
34:YA:2735:G:H2'	34:YA:2736:G:C8	2.54	0.43
35:YB:37:C:O2	47:YS:95:HIS:NE2	2.41	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
38:YF:158:THR:OG1	38:YF:159:GLY:N	2.51	0.43
47:YS:14:VAL:O	47:YS:18:ILE:HG12	2.18	0.43
48:YT:120:ARG:HA	48:YT:123:GLN:HB2	2.01	0.43
1:QA:1099:G:O5'	2:QB:96:ARG:HD2	2.19	0.43
1:QA:1221:G:O4'	19:QS:36:ARG:NH2	2.43	0.43
1:QA:1377:A:H2'	1:QA:1377:A:N3	2.34	0.43
4:QD:100:ARG:HG3	4:QD:137:SER:HA	2.00	0.43
12:QL:76:ASN:HD21	12:QL:108:ALA:HB3	1.84	0.43
34:RA:69:C:H2'	34:RA:70:G:H8	1.83	0.43
34:RA:121:G:H4'	34:RA:149:A:H5'	1.99	0.43
34:RA:453:C:H4'	34:RA:472:A:H62	1.84	0.43
34:RA:453:C:H4'	34:RA:472:A:N6	2.34	0.43
34:RA:1044:G:H1'	34:RA:1048:A:C4	2.54	0.43
34:RA:2068:U:H3	34:RA:2430:A:H2	1.65	0.43
34:RA:2134:A:H1'	34:RA:2158:A:C2	2.54	0.43
34:RA:2250:G:C6	45:RQ:82:ARG:CD	3.00	0.43
34:RA:2260:C:H2'	34:RA:2261:C:H6	1.82	0.43
37:RE:35:GLN:HB2	37:RE:37:ARG:HH21	1.84	0.43
40:RH:46:GLU:OE1	40:RH:51:ARG:NH1	2.52	0.43
43:RO:15:GLY:HA2	43:RO:47:ILE:HG22	2.00	0.43
1:XA:164:U:H2'	1:XA:165:C:C6	2.53	0.43
1:XA:290:C:H2'	1:XA:291:C:H6	1.83	0.43
1:XA:766:A:H2'	1:XA:767:A:O4'	2.18	0.43
1:XA:964:A:O2'	10:XJ:55:LYS:NZ	2.47	0.43
5:XE:105:VAL:HG11	5:XE:132:ALA:HB2	1.99	0.43
14:XN:53:LEU:HD12	14:XN:54:PRO:HD2	2.01	0.43
17:XQ:21:VAL:N	17:XQ:42:TYR:O	2.40	0.43
18:XR:21:LYS:HE2	18:XR:21:LYS:HB3	1.81	0.43
24:Y0:72:ARG:CB	24:Y0:75:LEU:HB2	2.48	0.43
34:YA:363(A):G:H2'	34:YA:363(B):A:C8	2.54	0.43
34:YA:511:U:H4'	34:YA:1235:G:H4'	2.01	0.43
34:YA:1204:A:H1'	34:YA:1206:G:N9	2.34	0.43
34:YA:1825:A:H2'	34:YA:1826:G:C8	2.53	0.43
34:YA:1849:G:H2'	34:YA:1850:G:C8	2.52	0.43
34:YA:2077:A:C6	34:YA:2435:A:C6	3.07	0.43
34:YA:2406:U:C4	44:YP:75:ILE:CD1	3.02	0.43
34:YA:2437:U:H2'	34:YA:2438:U:C6	2.53	0.43
34:YA:2469:A:H5''	34:YA:2470:G:H8	1.84	0.43
34:YA:2550:G:H2'	34:YA:2551:C:C6	2.54	0.43
34:YA:2567:G:H2'	34:YA:2568:C:C6	2.53	0.43
48:YT:16:ARG:NH2	48:YT:83:ILE:O	2.51	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
54:YZ:48:PHE:HA	54:YZ:51:ALA:HB3	2.00	0.43
1:QA:137:C:H2'	1:QA:138:G:C8	2.54	0.43
1:QA:359:U:H2'	1:QA:360:A:C8	2.49	0.43
1:QA:1143:G:H2'	1:QA:1144:G:C8	2.54	0.43
1:QA:1288:A:N1	1:QA:1371:G:H1'	2.33	0.43
1:QA:1379:G:H2'	1:QA:1380:U:C6	2.54	0.43
1:QA:1440(J):A:H2'	1:QA:1440(L):G:O6	2.19	0.43
1:QA:1440(O):C:H2'	1:QA:1440(P):A:H8	1.84	0.43
6:QF:78:GLU:HA	6:QF:81:ILE:HG12	2.01	0.43
11:QK:101:SER:O	11:QK:101:SER:OG	2.30	0.43
13:QM:29:ARG:HH21	13:QM:64:TRP:HE1	1.67	0.43
34:RA:288:C:H2'	34:RA:289:A:C8	2.53	0.43
34:RA:494:G:OP1	51:RW:8:ARG:NH1	2.52	0.43
34:RA:530:G:N3	34:RA:2021:C:H1'	2.34	0.43
34:RA:740:U:H2'	34:RA:741:G:C8	2.54	0.43
34:RA:746:A:H3'	34:RA:2612:C:C5	2.53	0.43
34:RA:1065:U:H3	34:RA:1069:A:H2'	1.82	0.43
34:RA:1112:G:H2'	34:RA:1113:U:O4'	2.19	0.43
34:RA:1297:C:O2'	34:RA:1302:A:N1	2.44	0.43
34:RA:2179:C:H2'	34:RA:2180:U:C6	2.54	0.43
34:RA:2374:C:H2'	34:RA:2375:G:C8	2.54	0.43
45:RQ:77:LYS:NZ	45:RQ:84:GLY:O	2.40	0.43
1:XA:458(B):G:H2'	1:XA:458(C):A:H2'	2.01	0.43
1:XA:556:C:H2'	1:XA:557:G:H8	1.83	0.43
1:XA:577:G:H1'	1:XA:816:A:H2'	2.00	0.43
1:XA:787:A:H2'	1:XA:788:U:C6	2.54	0.43
1:XA:996:A:H2'	1:XA:997:U:H6	1.84	0.43
1:XA:1253:G:C4	1:XA:1254:C:C5	3.07	0.43
1:XA:1505:G:H1'	23:XX:15:A:H2	1.83	0.43
1:XA:1525:G:H2'	1:XA:1526:G:C8	2.54	0.43
3:XC:8:ILE:HD11	3:XC:184:TYR:H	1.84	0.43
13:XM:52:GLU:HA	13:XM:55:ARG:HG2	2.00	0.43
20:XT:17:ARG:HE	20:XT:17:ARG:HB3	1.73	0.43
21:XU:3:LYS:CD	21:XU:14:TRP:CD2	3.00	0.43
24:Y0:26:TYR:N	24:Y0:29:GLN:OE1	2.52	0.43
32:Y8:56:GLU:OE1	32:Y8:59:LYS:NZ	2.36	0.43
34:YA:436:C:H2'	34:YA:438:G:C8	2.53	0.43
34:YA:635:C:H2'	34:YA:636:G:H8	1.84	0.43
34:YA:675:A:N3	34:YA:2443:C:O2'	2.47	0.43
34:YA:1660:C:H2'	34:YA:1661:G:H8	1.83	0.43
34:YA:1772:G:H21	34:YA:1774:C:H5'	1.84	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:2131:G:H1'	34:YA:2158:A:H61	1.83	0.43
34:YA:2543:G:H2'	34:YA:2544:G:C8	2.53	0.43
38:YF:48:THR:OG1	38:YF:48:THR:O	2.36	0.43
47:YS:4:LEU:HD11	47:YS:12:PHE:CE2	2.51	0.43
1:QA:186(H):C:H1'	1:QA:186(L):G:N2	2.34	0.42
1:QA:324:G:H2'	1:QA:326:G:N7	2.34	0.42
1:QA:418:C:H2'	1:QA:419:C:C6	2.53	0.42
1:QA:457:C:H2'	1:QA:458(A):C:C6	2.54	0.42
1:QA:518:C:H2'	1:QA:530:G:N3	2.34	0.42
1:QA:563:A:O2'	1:QA:566:G:O2'	2.29	0.42
1:QA:718:G:H21	18:QR:49:LYS:HE3	1.83	0.42
1:QA:979:C:C4	14:QN:19:ARG:CB	3.00	0.42
1:QA:1437:C:H2'	1:QA:1438:G:H8	1.83	0.42
3:QC:12:LEU:HG	3:QC:18:TRP:HE1	1.83	0.42
8:QH:10:LEU:HG	8:QH:83:ILE:HD11	1.99	0.42
13:QM:29:ARG:HD2	13:QM:29:ARG:HA	1.79	0.42
28:R4:34:GLU:OE1	39:RG:113:ARG:CD	2.66	0.42
34:RA:236:C:H2'	34:RA:237:C:H6	1.84	0.42
34:RA:272:G:H2'	34:RA:273(A):G:C8	2.54	0.42
34:RA:286:C:H2'	34:RA:287:C:C6	2.54	0.42
34:RA:1870:C:H2'	34:RA:1871:A:O4'	2.19	0.42
34:RA:1908:C:H2'	34:RA:1909:C:H6	1.84	0.42
34:RA:2161:C:H2'	34:RA:2162:G:C8	2.54	0.42
37:RE:120:TRP:CD2	37:RE:155:LYS:HB3	2.54	0.42
44:RP:65:ARG:O	44:RP:68:GLN:NE2	2.52	0.42
54:RZ:53:ILE:HG22	54:RZ:71:VAL:HG13	1.99	0.42
1:XA:7:G:H5'	1:XA:298:A:H5'	2.00	0.42
1:XA:107:G:C2	1:XA:108:G:H1'	2.54	0.42
1:XA:310:G:H5''	16:XP:31:LYS:HB2	2.00	0.42
1:XA:503:C:OP1	12:XL:119:LYS:HE2	2.19	0.42
1:XA:585:G:N3	1:XA:879:C:H4'	2.34	0.42
1:XA:639:G:H2'	1:XA:640:A:H8	1.84	0.42
1:XA:657:G:H4'	15:XO:28:GLN:HG2	2.01	0.42
1:XA:679:C:H2'	1:XA:680:C:C6	2.53	0.42
1:XA:857:C:H2'	1:XA:858:G:O4'	2.20	0.42
1:XA:1049:U:OP1	14:YN:3:ARG:NH2	2.52	0.42
1:XA:1217:C:P	14:YN:5:ALA:CB	2.98	0.42
1:XA:1361:G:C4	14:YN:18:VAL:HG12	2.54	0.42
11:XK:54:ARG:O	11:XK:57:THR:OG1	2.31	0.42
33:Y9:10:ILE:N	33:Y9:14:CYS:SG	2.84	0.42
34:YA:151:C:H2'	34:YA:152:G:C8	2.52	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:270(V):C:H2'	34:YA:270(W):G:C8	2.53	0.42
34:YA:687:C:N3	34:YA:787:U:H4'	2.34	0.42
34:YA:741:G:H2'	34:YA:742:G:C8	2.54	0.42
34:YA:1069:A:H2'	34:YA:1073:A:N7	2.34	0.42
34:YA:1278:A:H2'	34:YA:1279:G:C8	2.54	0.42
34:YA:1535:U:C4	34:YA:1537:C:H1'	2.54	0.42
34:YA:2786:U:O2'	37:YE:62:PRO:HA	2.18	0.42
34:YA:2840:C:H2'	34:YA:2841:C:C6	2.54	0.42
54:YZ:85:HIS:NE2	54:YZ:87:ASP:OD1	2.52	0.42
1:QA:157:G:H2'	1:QA:158:G:C8	2.54	0.42
1:QA:323:U:H4'	20:QT:19:SER:HA	2.01	0.42
1:QA:781:A:H4'	1:QA:1522:U:O2'	2.19	0.42
1:QA:1260:C:H2'	1:QA:1275:A:H61	1.84	0.42
1:QA:1410:G:H2'	1:QA:1411:C:C6	2.55	0.42
13:QM:28:ALA:CB	21:QU:21:TYR:OH	2.68	0.42
34:RA:807:U:H2'	34:RA:808:G:C8	2.54	0.42
34:RA:825:C:H4'	34:RA:2428:G:N7	2.34	0.42
34:RA:1711:C:H2'	34:RA:1712:C:C6	2.54	0.42
34:RA:2547:U:H2'	34:RA:2548:G:C8	2.54	0.42
34:RA:2580:U:H5'	37:RE:130:GLY:O	2.19	0.42
46:RR:11:ASN:O	46:RR:12:ARG:HG3	2.19	0.42
54:RZ:130:PRO:HA	54:RZ:133:ILE:HD11	2.01	0.42
1:XA:68(P):A:C5	1:XA:68(Q):C:H1'	2.54	0.42
1:XA:518:C:H5	1:XA:529:G:H3'	1.83	0.42
1:XA:1305:G:C6	1:XA:1331:G:N7	2.81	0.42
10:XJ:62:HIS:NE2	14:XN:61:TRP:HE3	2.16	0.42
32:Y8:4:MET:HG3	32:Y8:61:LEU:HD11	2.01	0.42
34:YA:67:U:H2'	34:YA:68:G:H8	1.83	0.42
34:YA:189:G:H2'	34:YA:205:G:N2	2.34	0.42
34:YA:335:C:H2'	34:YA:336:C:H6	1.84	0.42
34:YA:779:U:H2'	34:YA:780:G:C8	2.53	0.42
37:YE:144:ARG:HG3	37:YE:145:LYS:H	1.84	0.42
40:YH:9:ILE:HG22	40:YH:49:VAL:HB	1.95	0.42
45:YQ:141:GLN:HG3	54:YZ:123:ASP:OD1	2.19	0.42
1:QA:45:U:H4'	1:QA:306:G:N2	2.34	0.42
1:QA:125:U:H2'	1:QA:126:G:C8	2.55	0.42
1:QA:320:C:H2'	1:QA:321:A:C8	2.53	0.42
1:QA:522:C:H2'	1:QA:523:A:H8	1.84	0.42
1:QA:624:C:O2'	16:QP:10:GLY:CA	2.68	0.42
1:QA:832:C:H2'	1:QA:833:U:C6	2.54	0.42
1:QA:1086:U:H3	1:QA:1099:G:H1	1.65	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1186:G:C2	14:QN:61:TRP:O	2.66	0.42
1:QA:1530:G:H2'	1:QA:1531:A:C8	2.54	0.42
4:QD:122:ARG:HA	4:QD:122:ARG:HD2	1.90	0.42
13:QM:37:THR:O	13:QM:55:ARG:NH2	2.53	0.42
13:QM:84:ILE:O	19:QS:74:PHE:HE2	1.77	0.42
17:QQ:45:HIS:H	17:QQ:72:ARG:HA	1.84	0.42
34:RA:237:C:H2'	34:RA:238:C:C6	2.54	0.42
34:RA:636:G:C4	44:RP:115:LEU:HD11	2.54	0.42
34:RA:828:U:H4'	34:RA:831:G:C2	2.54	0.42
34:RA:1629:U:H2'	34:RA:1630(A):G:C8	2.55	0.42
34:RA:1857:G:H21	34:RA:1886:C:H42	1.67	0.42
34:RA:1909:C:H2'	34:RA:1910:G:C8	2.53	0.42
34:RA:2208:U:H2'	34:RA:2209:C:C6	2.54	0.42
34:RA:2363:C:H2'	34:RA:2364:C:H6	1.84	0.42
34:RA:2658:C:H5''	40:RH:158:HIS:NE2	2.34	0.42
37:RE:109:LYS:HG2	37:RE:191:PRO:HB3	2.00	0.42
1:XA:181:G:H4'	1:XA:182:U:H5'	2.01	0.42
1:XA:280:C:C4	17:XQ:39:SER:N	2.70	0.42
1:XA:538:G:H2'	1:XA:539:A:C8	2.54	0.42
1:XA:677:U:H2'	1:XA:678:U:C6	2.53	0.42
1:XA:684:A:H2'	1:XA:685:G:C8	2.54	0.42
1:XA:712:A:H2'	1:XA:713:G:C8	2.53	0.42
1:XA:787:A:H2'	1:XA:788:U:H6	1.84	0.42
1:XA:951:G:H1'	1:XA:971:G:H5'	2.00	0.42
1:XA:1376:U:C5	7:XG:10:ARG:HD3	2.53	0.42
1:XA:1439:C:H2'	1:XA:1440(A):C:C6	2.54	0.42
3:XC:66:VAL:HB	3:XC:101:LEU:HD13	2.00	0.42
6:XF:62:TRP:NE1	18:XR:35:ARG:NH2	2.63	0.42
6:XF:89:MET:SD	18:XR:76:LEU:HD22	2.59	0.42
7:XG:16:LEU:CD2	9:XI:45:ALA:HB2	2.47	0.42
12:XL:42:THR:HA	12:XL:53:ARG:O	2.18	0.42
14:XN:23:ARG:NH1	14:XN:24:CYS:HB3	2.34	0.42
34:YA:521:G:H2'	34:YA:522:G:H8	1.84	0.42
34:YA:532:A:N1	34:YA:2020:A:H1'	2.33	0.42
34:YA:935:C:H2'	34:YA:936:C:C6	2.54	0.42
34:YA:945:A:H62	34:YA:2448:A:N6	1.74	0.42
34:YA:1198:U:H2'	34:YA:1199:U:C6	2.54	0.42
34:YA:2014:A:H2'	34:YA:2015:A:C4	2.54	0.42
34:YA:2136:C:H2'	34:YA:2137:C:C6	2.54	0.42
34:YA:2182:G:H2'	34:YA:2183:C:C6	2.54	0.42
40:YH:159:GLU:HG2	40:YH:169:VAL:HG21	2.01	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:445:G:H2'	1:QA:446:G:C8	2.52	0.42
1:QA:972:C:OP2	10:QJ:57:LYS:HE3	2.20	0.42
5:QE:102:ALA:H	5:QE:107:ARG:NH2	2.18	0.42
7:QG:58:PRO:HA	7:QG:61:VAL:HG12	2.01	0.42
22:QV:50:G:H2'	22:QV:51:A:H8	1.85	0.42
24:R0:29:GLN:HG3	34:RA:923:C:H4'	2.01	0.42
25:R1:2:SER:N	34:RA:1364:G:OP2	2.52	0.42
34:RA:245:G:N2	34:RA:384:U:O2'	2.37	0.42
34:RA:949:C:H2'	34:RA:950:G:C8	2.54	0.42
34:RA:1162:G:H4'	50:RV:24:LYS:HB3	2.01	0.42
34:RA:1292:U:H2'	34:RA:1293:C:C6	2.54	0.42
34:RA:1999:C:H2'	34:RA:2000:G:C8	2.50	0.42
34:RA:2513:G:N2	37:RE:143:ASN:OD1	2.52	0.42
34:RA:2863:C:H2'	34:RA:2864:G:C8	2.55	0.42
36:RD:13:ARG:HD2	36:RD:13:ARG:HA	1.80	0.42
36:RD:49:ILE:HD11	36:RD:52:ARG:HA	2.00	0.42
46:RR:8:ARG:HD2	46:RR:10:LEU:HD21	2.01	0.42
1:XA:133:U:O5'	1:XA:133:U:H6	2.03	0.42
1:XA:741:G:P	15:XO:35:ARG:HH21	1.54	0.42
1:XA:1055:A:H2'	1:XA:1056:U:O4'	2.18	0.42
33:Y9:6:SER:O	33:Y9:6:SER:OG	2.35	0.42
34:YA:11:G:H22	34:YA:2627:G:H5''	1.83	0.42
34:YA:13:A:N1	34:YA:525:U:H2'	2.34	0.42
34:YA:654(B):G:N2	34:YA:654(V):A:H1'	2.35	0.42
34:YA:665:C:H2'	34:YA:666:G:H8	1.83	0.42
34:YA:741:G:H2'	34:YA:742:G:H8	1.84	0.42
34:YA:1102:C:H2'	34:YA:1103:A:C8	2.55	0.42
34:YA:1399:C:H2'	34:YA:1400:G:H8	1.85	0.42
34:YA:1430:C:H2'	34:YA:1431:U:C6	2.55	0.42
34:YA:1820:U:O2	36:YD:202:LYS:HB3	2.19	0.42
34:YA:2014:A:H8	34:YA:2014:A:OP2	2.03	0.42
34:YA:2452:C:H2'	34:YA:2453:A:C8	2.54	0.42
44:YP:46:LYS:HB3	44:YP:46:LYS:HE3	1.78	0.42
51:YW:57:ASN:O	51:YW:61:ASN:HB2	2.19	0.42
53:YY:79:CYS:SG	53:YY:81:LYS:HE2	2.60	0.42
1:QA:129(B):G:H5'	1:QA:186(L):G:H5'	2.01	0.42
1:QA:410:G:H2'	1:QA:429:U:C4	2.55	0.42
1:QA:630:G:H2'	1:QA:631:G:C8	2.55	0.42
1:QA:953:G:H3'	1:QA:954:G:C8	2.54	0.42
1:QA:976:G:N2	1:QA:1362(A):C:H2'	2.35	0.42
1:QA:979:C:N4	1:QA:1318:A:H61	2.18	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:1014:A:N9	19:QS:34:TRP:NE1	2.50	0.42
1:QA:1271:G:H2'	1:QA:1272:G:C8	2.54	0.42
1:QA:1472:U:H2'	1:QA:1473:A:C8	2.54	0.42
2:QB:34:ALA:H	2:QB:41:ILE:HB	1.84	0.42
3:QC:43:LEU:HG	3:QC:47:LEU:HD22	2.00	0.42
4:QD:59:ARG:HH12	4:QD:66:ARG:NH2	2.16	0.42
4:QD:172:PRO:HG3	6:XF:28:ARG:HH12	1.84	0.42
5:QE:126:ARG:HA	5:QE:131:ILE:HD11	2.01	0.42
12:QL:79:GLU:OE2	12:QL:80:HIS:NE2	2.53	0.42
13:QM:47:ASP:OD1	13:QM:47:ASP:N	2.48	0.42
34:RA:528:A:C2	34:RA:2042:A:H2'	2.55	0.42
34:RA:941:A:O2'	34:RA:1190:G:O3'	2.30	0.42
34:RA:1123:C:H2'	34:RA:1124:C:H6	1.85	0.42
34:RA:2159:G:H2'	34:RA:2160:G:H8	1.82	0.42
37:RE:134:ILE:HD12	37:RE:134:ILE:HA	1.85	0.42
51:RW:13:SER:HA	51:RW:99:ARG:HB2	2.01	0.42
54:RZ:67:LEU:HD12	54:RZ:90:VAL:HB	2.02	0.42
1:XA:262:A:H2'	1:XA:263:A:C8	2.55	0.42
1:XA:400:C:H2'	1:XA:401:C:C6	2.55	0.42
1:XA:714:G:H1'	1:XA:777:A:C8	2.55	0.42
1:XA:757:U:H2'	1:XA:758:G:O4'	2.18	0.42
1:XA:821:G:H2'	1:XA:822:C:O4'	2.19	0.42
1:XA:948:C:OP2	13:XM:106:ASN:OD1	2.36	0.42
1:XA:1084:G:H21	1:XA:1102:A:H62	1.67	0.42
1:XA:1259:C:HO2'	1:XA:1283:G:H21	1.65	0.42
1:XA:1305:G:H4'	1:XA:1332:A:N6	2.34	0.42
3:XC:187:ALA:HB3	3:XC:198:VAL:HG13	2.00	0.42
6:XF:27:GLN:HA	6:XF:30:LEU:HD12	2.02	0.42
10:XJ:62:HIS:CD2	14:YN:59:ALA:CB	2.99	0.42
32:Y8:13:ARG:NH2	34:YA:250:G:OP2	2.52	0.42
34:YA:576:U:H2'	34:YA:577:G:C8	2.55	0.42
34:YA:962:G:H2'	34:YA:963:U:C6	2.55	0.42
34:YA:1819:A:H5''	36:YD:161:THR:HG21	2.02	0.42
34:YA:2154:G:H2'	34:YA:2155:G:C8	2.54	0.42
35:YB:44:G:H1'	35:YB:47:C:H42	1.83	0.42
46:YR:35:THR:HA	46:YR:112:ALA:O	2.19	0.42
47:YS:34:HIS:O	47:YS:97:ARG:NH2	2.52	0.42
1:QA:21:G:H2'	1:QA:22:G:C8	2.55	0.42
1:QA:287:U:H2'	1:QA:288:A:H8	1.85	0.42
1:QA:358:U:H2'	1:QA:359:U:C6	2.55	0.42
1:QA:408:A:H3'	1:QA:409:G:H8	1.83	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:766:A:H2'	1:QA:767:A:O4'	2.19	0.42
1:QA:926:G:H2'	1:QA:1505:G:N3	2.34	0.42
1:QA:987:G:H2'	1:QA:988:G:C8	2.55	0.42
1:QA:1109:C:OP2	3:QC:176:HIS:HD2	2.02	0.42
1:QA:1126:U:H3'	1:QA:1127:G:H8	1.83	0.42
1:QA:1134:G:H3'	1:QA:1135:U:C6	2.54	0.42
1:QA:1390:U:H2'	1:QA:1391:U:C6	2.54	0.42
4:QD:15:GLU:HG2	4:QD:63:LYS:HB3	2.01	0.42
7:QG:75:VAL:HA	7:QG:88:PRO:HA	2.00	0.42
34:RA:13:A:N3	34:RA:14:A:N6	2.66	0.42
34:RA:262:A:H2'	34:RA:263:C:O4'	2.19	0.42
34:RA:375:C:H2'	34:RA:376:C:C6	2.54	0.42
34:RA:672:C:H2'	34:RA:673:C:C6	2.55	0.42
34:RA:704:G:N2	34:RA:726:G:O2'	2.52	0.42
34:RA:1509:C:H3'	34:RA:1510:A:H4'	2.00	0.42
34:RA:2533:A:OP1	34:RA:2665:A:O2'	2.29	0.42
34:RA:2841:C:H2'	34:RA:2842:G:H8	1.84	0.42
36:RD:32:SER:C	36:RD:34:VAL:H	2.23	0.42
36:RD:142:VAL:HG23	36:RD:193:VAL:HA	2.02	0.42
45:RQ:7:MET:SD	45:RQ:7:MET:N	2.90	0.42
1:XA:186(Q):U:O2	20:XT:105:SER:OG	2.37	0.42
1:XA:972:C:C2'	10:XJ:55:LYS:HB2	2.49	0.42
1:XA:1086:U:H3	1:XA:1099:G:H22	1.67	0.42
1:XA:1300:G:O3'	13:XM:21:TYR:HE1	2.03	0.42
1:XA:1492:A:H1'	1:XA:1493:A:C6	2.54	0.42
2:XB:87:ARG:HH21	2:XB:219:VAL:HG13	1.84	0.42
2:XB:195:ASP:O	8:XH:74:PRO:HG2	2.20	0.42
3:XC:64:VAL:HG13	3:XC:97:LYS:HD2	2.01	0.42
7:XG:102:ARG:HA	7:XG:105:VAL:HG22	2.01	0.42
11:XK:108:ILE:O	18:XR:87:ARG:HA	2.19	0.42
18:XR:85:LEU:HD12	18:XR:85:LEU:HA	1.83	0.42
34:YA:65:C:H2'	34:YA:66:C:C6	2.55	0.42
34:YA:128:C:H2'	34:YA:129:C:C6	2.54	0.42
34:YA:373:U:H1'	34:YA:423:A:N3	2.35	0.42
34:YA:654(B):G:H22	34:YA:654(V):A:H1'	1.84	0.42
34:YA:676:A:C8	34:YA:2070:G:H1'	2.55	0.42
34:YA:689:A:H2'	34:YA:690:G:C8	2.54	0.42
34:YA:852:G:H2'	34:YA:853:G:H8	1.85	0.42
34:YA:985:C:H2'	34:YA:986:C:C6	2.55	0.42
34:YA:1012:U:O4	42:YN:25:ARG:HD3	2.19	0.42
34:YA:1303:G:H1'	34:YA:1641:A:C2	2.55	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:1556:C:H2'	34:YA:1557:C:C6	2.54	0.42
34:YA:2299:G:N2	34:YA:2318:G:H1'	2.34	0.42
34:YA:2305:A:O5'	39:YG:134:GLY:HA3	2.20	0.42
34:YA:2584:U:O3'	34:YA:2602:A:H2	2.02	0.42
34:YA:2689:U:OP2	34:YA:2872:G:N2	2.41	0.42
38:YF:114:VAL:HG11	38:YF:202:PHE:CZ	2.55	0.42
44:YP:82:GLY:HA2	44:YP:113:LYS:O	2.18	0.42
1:QA:21:G:H1'	1:QA:915:A:N1	2.34	0.42
1:QA:34:C:H2'	1:QA:35:G:H8	1.83	0.42
1:QA:320:C:N4	1:QA:329:A:OP2	2.52	0.42
1:QA:740:U:O4'	15:QO:42:HIS:CD2	2.73	0.42
1:QA:821:G:H8	1:QA:821:G:P	2.43	0.42
1:QA:943:U:O2'	1:QA:1232:U:OP2	2.36	0.42
1:QA:953:G:C8	13:QM:104:ARG:NH2	2.79	0.42
1:QA:1099:G:C5'	2:QB:96:ARG:HD2	2.49	0.42
1:QA:1466:C:H2'	1:QA:1467:G:O4'	2.19	0.42
12:QL:57:LYS:HA	12:QL:67:THR:HA	2.00	0.42
34:RA:105:C:H2'	34:RA:106:C:C6	2.54	0.42
34:RA:189:G:H2'	34:RA:205:G:N2	2.34	0.42
34:RA:263:C:H2'	34:RA:264:C:O4'	2.20	0.42
34:RA:270(V):C:H2'	34:RA:270(W):G:H8	1.85	0.42
34:RA:524:U:H2'	34:RA:525:U:C6	2.54	0.42
34:RA:1041:C:H2'	34:RA:1042:G:C8	2.51	0.42
34:RA:1096:A:H2'	34:RA:1097:U:O4'	2.20	0.42
34:RA:1107:G:H2'	34:RA:1108:U:C6	2.54	0.42
34:RA:1165:U:H2'	34:RA:1166:C:C6	2.55	0.42
34:RA:1375:C:H2'	34:RA:1376:C:C6	2.55	0.42
34:RA:2065:C:H1'	34:RA:2449:U:H3	1.84	0.42
38:RF:70:THR:HG23	38:RF:72:ARG:H	1.85	0.42
1:XA:105:G:H2'	1:XA:106:C:C6	2.55	0.42
1:XA:111:G:O5'	1:XA:111:G:C8	2.70	0.42
1:XA:373:A:HO2'	1:XA:374:A:H5'	1.84	0.42
1:XA:474:G:H2'	1:XA:475:G:C8	2.55	0.42
1:XA:920:U:O2'	1:XA:1081:G:O2'	2.36	0.42
1:XA:926:G:H2'	1:XA:1505:G:N2	2.35	0.42
1:XA:1330:U:C5'	13:XM:24:GLY:O	2.68	0.42
2:XB:177:ALA:HB1	2:XB:182:ILE:HB	2.01	0.42
3:XC:119:ARG:HG2	3:XC:140:ARG:HH22	1.85	0.42
33:Y9:32:HIS:CD2	33:Y9:32:HIS:N	2.87	0.42
34:YA:118:A:N3	34:YA:178:G:H1'	2.35	0.42
34:YA:134:C:H2'	34:YA:135:G:C8	2.54	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:270(S):G:H2'	34:YA:270(T):G:H8	1.85	0.42
34:YA:584:C:OP2	49:YU:10:ARG:NH2	2.53	0.42
34:YA:603:A:O4'	34:YA:655:A:N6	2.52	0.42
34:YA:654(B):G:H2'	34:YA:654(C):C:C6	2.55	0.42
34:YA:673:C:H2'	34:YA:674:G:C8	2.55	0.42
34:YA:1869:G:H2'	34:YA:1871:A:N7	2.35	0.42
34:YA:2072:G:N1	34:YA:2438:U:C2	2.88	0.42
34:YA:2109:U:H2'	34:YA:2110:G:C8	2.55	0.42
34:YA:2291:U:H5''	34:YA:2380:C:H1'	2.02	0.42
34:YA:2419:U:H2'	34:YA:2420:C:C6	2.55	0.42
34:YA:2855:C:H2'	34:YA:2856:C:C6	2.54	0.42
39:YG:18:GLU:OE1	39:YG:21:ARG:NH2	2.50	0.42
47:YS:18:ILE:HG21	47:YS:88:ASP:HA	2.02	0.42
52:YX:25:LYS:HB3	52:YX:80:ILE:HD11	2.00	0.42
1:QA:8:A:C8	4:QD:209:ARG:O	2.72	0.42
1:QA:344:A:H3'	1:QA:345:C:C6	2.55	0.42
1:QA:345:C:H5'	48:RT:41:ARG:HD2	2.02	0.42
1:QA:581:G:N1	1:QA:759:A:OP2	2.37	0.42
1:QA:822:C:H2'	1:QA:823:G:C8	2.54	0.42
1:QA:1070:U:H2'	1:QA:1071:C:C6	2.55	0.42
1:QA:1294:G:H2'	1:QA:1295:G:C8	2.55	0.42
1:QA:1437:C:H2'	1:QA:1438:G:C8	2.55	0.42
2:QB:175:ARG:O	2:QB:179:LYS:HG3	2.20	0.42
4:QD:205:GLU:CG	5:QE:107:ARG:HH12	2.33	0.42
9:QI:65:VAL:HG11	9:QI:73:GLN:HG3	2.01	0.42
12:QL:102:ARG:HB3	12:QL:109:GLY:HA2	2.01	0.42
14:QN:41:ARG:HG3	14:QN:42:ILE:HG13	2.01	0.42
32:R8:61:LEU:HD12	32:R8:62:LEU:HG	2.02	0.42
34:RA:191:A:N6	34:RA:206:U:O2	2.53	0.42
34:RA:476:G:H1'	34:RA:480:A:H61	1.84	0.42
34:RA:755:C:H2'	34:RA:756:C:C6	2.55	0.42
34:RA:1572:A:H2'	34:RA:1573:G:C8	2.55	0.42
34:RA:2333:A:H1'	34:RA:2335:A:C5	2.55	0.42
34:RA:2683:C:OP1	48:RT:53:ARG:NH1	2.42	0.42
34:RA:2735:G:H2'	34:RA:2736:G:C8	2.55	0.42
1:XA:373:A:H1'	1:XA:481:G:O4'	2.20	0.42
1:XA:477:G:H2'	1:XA:478:A:C8	2.55	0.42
1:XA:714:G:H2'	1:XA:715:A:C8	2.55	0.42
1:XA:729:A:H2'	1:XA:730:G:H8	1.84	0.42
1:XA:901:A:O5'	1:XA:901:A:C8	2.70	0.42
1:XA:946:A:OP1	13:XM:114:ARG:NH2	2.53	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1253:G:N1	1:XA:1254:C:N4	2.67	0.42
1:XA:1311:G:H3'	19:XS:5:LEU:HD11	1.31	0.42
1:XA:1313:U:C5	19:XS:4:SER:OG	2.72	0.42
1:XA:1329:A:H4'	13:XM:29:ARG:CZ	2.50	0.42
1:XA:1512:U:H2'	1:XA:1513:A:C8	2.54	0.42
30:Y6:40:CYS:HB3	30:Y6:43:CYS:CB	2.50	0.42
34:YA:610:C:H2'	34:YA:611:C:C6	2.55	0.42
34:YA:740:U:H2'	34:YA:741:G:C8	2.54	0.42
34:YA:765:G:H2'	34:YA:766:C:C6	2.55	0.42
34:YA:1119:C:H2'	34:YA:1120:G:H8	1.85	0.42
34:YA:1174:A:H2	34:YA:1176:G:H4'	1.85	0.42
34:YA:1292:U:H2'	34:YA:1293:C:C6	2.55	0.42
34:YA:1534:G:H3'	34:YA:1534:G:N3	2.34	0.42
34:YA:1812:A:H2'	34:YA:1813:G:C8	2.55	0.42
34:YA:2053:G:H2'	34:YA:2054:A:C8	2.55	0.42
34:YA:2249:U:C4	34:YA:2252:G:H5''	2.55	0.42
34:YA:2468:G:OP2	34:YA:2468:G:N2	2.51	0.42
34:YA:2554:U:H2'	34:YA:2555:U:C6	2.55	0.42
37:YE:102:VAL:HG23	37:YE:200:GLU:HA	2.02	0.42
45:YQ:110:THR:OG1	45:YQ:111:GLU:N	2.53	0.42
46:YR:87:TYR:OH	46:YR:117:VAL:O	2.31	0.42
48:YT:6:LEU:HA	48:YT:9:LEU:HB2	2.02	0.42
1:QA:129(A):U:H4'	1:QA:130:A:OP1	2.18	0.42
1:QA:163:C:H2'	1:QA:164:U:O4'	2.20	0.42
1:QA:830:G:H21	1:QA:1540:U:H5'	1.84	0.42
1:QA:1313:U:C4'	19:QS:6:LYS:HZ3	2.29	0.42
1:QA:1373:G:P	9:QI:11:LYS:HZ2	2.41	0.42
1:QA:1509:C:H2'	1:QA:1510:U:H6	1.85	0.42
2:QB:197:VAL:C	8:QH:68:ARG:NH2	2.70	0.42
4:QD:18:LYS:HZ2	4:QD:18:LYS:HG2	1.69	0.42
33:R9:29:ASN:HB3	33:R9:32:HIS:CD2	2.55	0.42
34:RA:26:G:H1'	34:RA:515:A:H61	1.85	0.42
34:RA:27:G:H1'	34:RA:513:A:H61	1.84	0.42
34:RA:389:G:H1	44:RP:71:VAL:HG12	1.84	0.42
34:RA:564:C:N4	34:RA:573:G:OP1	2.49	0.42
34:RA:611:C:H2'	34:RA:612:G:C8	2.55	0.42
34:RA:730:C:H2'	34:RA:731:C:C6	2.55	0.42
34:RA:804:A:H2'	34:RA:806:C:C4	2.55	0.42
34:RA:1444(B):A:H1'	34:RA:1460:A:N3	2.35	0.42
34:RA:1629:U:O2	34:RA:2698:U:H4'	2.20	0.42
34:RA:1654:A:OP1	46:RR:2:ARG:HD3	2.20	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:2651:C:H2'	34:RA:2652:C:H6	1.84	0.42
34:RA:2784:C:H2'	34:RA:2785:C:C6	2.55	0.42
34:RA:2859:G:H2'	34:RA:2860:A:C8	2.54	0.42
38:RF:8:GLN:NE2	38:RF:19:GLU:OE1	2.51	0.42
40:RH:126:PRO:HB2	40:RH:127:GLU:H	1.68	0.42
1:XA:321:A:N6	1:XA:328:C:H1'	2.35	0.42
1:XA:599:C:H2'	1:XA:600:C:C6	2.55	0.42
1:XA:1359:C:H4'	1:XA:1362(A):C:N4	2.35	0.42
19:XS:36:ARG:NH2	19:XS:72:GLY:O	2.52	0.42
22:XV:50:G:H2'	22:XV:51:A:C8	2.54	0.42
34:YA:417:C:O2	34:YA:2407:G:N1	2.52	0.42
34:YA:811:U:O4	44:YP:21:ARG:NH2	2.53	0.42
34:YA:1576:U:H2'	34:YA:1577:C:C6	2.54	0.42
34:YA:2043:C:N4	34:YA:2777:G:C5	2.87	0.42
34:YA:2108:C:H5'	34:YA:2150:U:O2'	2.19	0.42
34:YA:2233:U:H2'	34:YA:2234:G:C8	2.54	0.42
36:YD:45:ASN:O	36:YD:47:GLY:N	2.52	0.42
49:YU:94:ASN:HD22	49:YU:94:ASN:C	2.22	0.42
1:QA:5:U:H4'	1:QA:6:G:C8	2.55	0.42
1:QA:124:G:H2'	1:QA:125:U:O4'	2.20	0.42
1:QA:339:C:H2'	1:QA:340:U:C6	2.55	0.42
1:QA:976:G:OP1	14:QN:31:ARG:HG2	2.19	0.42
1:QA:1014:A:C1'	19:QS:34:TRP:CG	3.03	0.42
1:QA:1085:U:H3'	1:QA:1086:U:C6	2.55	0.42
6:QF:50:TYR:CZ	18:QR:77:GLY:HA2	2.55	0.42
21:QU:3:LYS:HD3	21:QU:14:TRP:CG	2.55	0.42
31:R7:12:ARG:HH21	31:R7:44:PRO:HB3	1.84	0.42
32:R8:49:VAL:HG23	32:R8:53:PRO:HD3	2.02	0.42
34:RA:1148:A:H2'	34:RA:1149:G:C8	2.55	0.42
34:RA:1200:C:H2'	34:RA:1201:C:C6	2.55	0.42
34:RA:1333:C:H2'	34:RA:1334:G:C8	2.47	0.42
34:RA:1688:U:H1'	34:RA:1701:A:C6	2.54	0.42
34:RA:2746:U:O4	34:RA:2755:C:H4'	2.20	0.42
36:RD:76:PRO:HB3	36:RD:118:VAL:HG22	2.02	0.42
1:XA:56:U:H2'	1:XA:57:G:C8	2.55	0.42
1:XA:390:C:O5'	1:XA:390:C:H6	2.03	0.42
1:XA:728:A:C8	15:XO:54:ARG:NH1	2.84	0.42
1:XA:1187:G:C4	14:YN:61:TRP:O	2.73	0.42
1:XA:1253:G:C2	1:XA:1254:C:C6	3.08	0.42
1:XA:1256:A:C8	1:XA:1278:U:H5'	2.54	0.42
1:XA:1414:U:H3	1:XA:1486:G:H1	1.68	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:1510:U:H3	1:XA:1525:G:H1	1.68	0.42
4:XD:105:VAL:HG21	4:XD:126:ILE:HD13	2.02	0.42
10:XJ:7:LYS:HE2	10:XJ:9:ARG:CG	2.49	0.42
13:XM:84:ILE:HD11	19:XS:65:ASN:CB	2.45	0.42
34:YA:279:C:H2'	34:YA:280:C:H6	1.85	0.42
34:YA:363(C):G:H2'	34:YA:363(D):G:C8	2.54	0.42
34:YA:453:C:H4'	34:YA:472:A:H62	1.84	0.42
34:YA:635:C:H2'	34:YA:636:G:C8	2.55	0.42
34:YA:892:G:H2'	34:YA:893:C:C6	2.54	0.42
34:YA:2086:U:OP2	36:YD:263:ARG:NH1	2.52	0.42
34:YA:2138:C:H2'	34:YA:2139:C:C6	2.55	0.42
37:YE:2:LYS:HD2	37:YE:95:ILE:HG22	2.02	0.42
42:YN:28:THR:O	42:YN:32:THR:OG1	2.32	0.42
44:YP:7:ARG:HA	44:YP:8:PRO:HD2	1.97	0.42
45:YQ:34:LEU:HB2	45:YQ:118:LEU:HD22	2.01	0.42
1:QA:10:A:OP2	5:QE:126:ARG:HD3	2.20	0.41
1:QA:669:U:H2'	1:QA:670:G:C8	2.54	0.41
1:QA:801:U:H2'	1:QA:802:A:H8	1.85	0.41
1:QA:893:C:H2'	1:QA:894:G:H8	1.85	0.41
1:QA:1124:G:C4'	10:QJ:36:GLY:H	2.33	0.41
1:QA:1130:A:H62	1:QA:1144:G:H21	1.67	0.41
1:QA:1176:A:H62	1:QA:1182:G:H21	1.67	0.41
1:QA:1220:G:N2	19:QS:54:GLY:HA2	2.33	0.41
1:QA:1483:A:H1'	34:RA:1948:G:H1'	2.02	0.41
3:QC:22:TRP:HH2	3:QC:32:LEU:HD13	1.85	0.41
4:QD:109:GLY:HA3	4:QD:165:MET:HG2	2.02	0.41
7:QG:75:VAL:HG12	7:QG:88:PRO:HB3	2.01	0.41
19:QS:70:LYS:N	19:QS:73:GLU:OE1	2.52	0.41
20:QT:14:LYS:HA	20:QT:17:ARG:HG2	2.02	0.41
34:RA:557:U:H2'	34:RA:558:G:C8	2.55	0.41
34:RA:1062:G:N7	34:RA:1088:A:O2'	2.44	0.41
34:RA:1501:C:H2'	34:RA:1502:C:C6	2.55	0.41
34:RA:1574:C:H2'	34:RA:1575:C:C6	2.55	0.41
34:RA:2538:C:H2'	34:RA:2539:C:C6	2.55	0.41
38:RF:178:PRO:HB3	38:RF:198:ALA:HB2	2.02	0.41
39:RG:34:LEU:HD21	39:RG:172:LEU:HD21	2.02	0.41
39:RG:39:ILE:HB	39:RG:92:VAL:HG13	2.01	0.41
44:RP:84:ASN:HA	44:RP:115:LEU:O	2.20	0.41
49:RU:85:LYS:HE2	49:RU:116:ALA:HA	2.01	0.41
1:XA:51:A:H61	1:XA:314:C:H1'	1.85	0.41
1:XA:68(Y):U:H2'	1:XA:68(Z):C:C6	2.55	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:167:G:H2'	1:XA:168:G:H8	1.85	0.41
1:XA:664:G:N3	1:XA:726:C:H4'	2.35	0.41
1:XA:1073:U:OP1	5:XE:57:LYS:HE3	2.19	0.41
1:XA:1184:G:H2'	1:XA:1185:G:C8	2.52	0.41
1:XA:1188:A:H4'	14:YN:59:ALA:O	2.19	0.41
1:XA:1229:A:OP2	13:XM:104:ARG:O	2.37	0.41
1:XA:1256:A:OP2	1:XA:1279:A:N6	2.52	0.41
1:XA:1500:A:H2'	1:XA:1501:C:C6	2.55	0.41
3:XC:8:ILE:HG12	3:XC:184:TYR:HB3	2.02	0.41
9:XI:13:ALA:HB1	9:XI:73:GLN:HG3	2.01	0.41
19:XS:30:LEU:HD13	19:XS:48:THR:HG23	2.01	0.41
32:Y8:60:LEU:HD12	32:Y8:60:LEU:HA	1.82	0.41
34:YA:220:G:N2	34:YA:427:U:H2'	2.35	0.41
34:YA:852:G:H2'	34:YA:853:G:C8	2.55	0.41
34:YA:1510:A:H2'	34:YA:1510:A:N3	2.33	0.41
34:YA:1675:C:O2	37:YE:128:SER:OG	2.37	0.41
34:YA:1709:U:H2'	34:YA:1710:C:C6	2.55	0.41
34:YA:2054:A:H2	34:YA:2616:C:N3	2.01	0.41
34:YA:2651:C:H2'	34:YA:2652:C:H6	1.85	0.41
36:YD:66:ASP:OD1	36:YD:103:ARG:NH1	2.53	0.41
36:YD:83:GLU:OE1	36:YD:104:TYR:OH	2.31	0.41
1:QA:151:A:H3'	1:QA:152:A:H8	1.85	0.41
1:QA:538:G:O5'	12:QL:115:LYS:HG3	2.19	0.41
1:QA:694:A:H2'	1:QA:695:A:O4'	2.20	0.41
1:QA:715:A:OP1	1:QA:805:C:O2'	2.31	0.41
1:QA:1105:A:H2'	1:QA:1106:G:C8	2.54	0.41
1:QA:1160:G:O4'	2:QB:132:LYS:HE2	2.10	0.41
1:QA:1175:G:H2'	1:QA:1176:A:C8	2.55	0.41
1:QA:1310:G:H2'	1:QA:1311:G:H8	1.84	0.41
9:QL:77:ILE:O	9:QL:81:ILE:HG12	2.20	0.41
26:R2:45:SER:OG	26:R2:46:GLN:N	2.53	0.41
34:RA:172:C:H2'	34:RA:173:G:C8	2.55	0.41
34:RA:242:G:H22	34:RA:254:G:H2'	1.84	0.41
34:RA:676:A:C2	34:RA:677:A:C8	3.09	0.41
34:RA:1212:G:H1'	34:RA:1236:G:N2	2.35	0.41
34:RA:1464:C:H2'	34:RA:1465:G:H8	1.85	0.41
34:RA:1511:A:H3'	34:RA:1512:G:H8	1.85	0.41
34:RA:1812:A:H2'	34:RA:1813:G:H8	1.85	0.41
34:RA:2427:C:H5'	34:RA:2429:G:H5'	2.00	0.41
34:RA:2751:G:N1	40:RH:3:ARG:CB	2.77	0.41
34:RA:2773:C:H2'	34:RA:2774:C:C6	2.55	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
49:RU:107:ALA:HA	49:RU:110:VAL:HG12	2.01	0.41
1:XA:260:G:H2'	1:XA:261:U:C6	2.56	0.41
1:XA:355:C:O5'	1:XA:355:C:C6	2.70	0.41
1:XA:532:A:O5'	1:XA:532:A:H8	2.03	0.41
1:XA:576:G:N2	1:XA:760:G:OP2	2.53	0.41
1:XA:580:U:H2'	1:XA:581:G:O4'	2.19	0.41
1:XA:584:G:H1	1:XA:757:U:H3	1.68	0.41
1:XA:730:G:N3	1:XA:765:G:H4'	2.34	0.41
1:XA:960:U:H4'	1:XA:961:U:H5''	2.02	0.41
1:XA:1376:U:C4	7:XG:10:ARG:CD	3.02	0.41
4:XD:101:LEU:HD22	4:XD:138:TYR:HB3	2.01	0.41
5:XE:75:THR:HG22	5:XE:117:ASP:O	2.20	0.41
5:XE:128:PRO:HA	5:XE:131:ILE:HG12	2.02	0.41
6:XF:39:LYS:HB2	6:XF:64:GLN:HB3	2.01	0.41
6:XF:89:MET:SD	18:XR:34:TYR:CE2	3.13	0.41
6:XF:99:ALA:HB3	18:XR:29:PHE:HD1	1.71	0.41
34:YA:279:C:H2'	34:YA:280:C:C6	2.55	0.41
34:YA:486:C:H2'	34:YA:487:C:C6	2.55	0.41
34:YA:870:A:OP1	45:YQ:6:ARG:NH2	2.25	0.41
34:YA:1995:U:H2'	34:YA:1996:C:C5	2.56	0.41
34:YA:2241:A:H2'	34:YA:2242:G:C8	2.54	0.41
34:YA:2732:G:H3'	34:YA:2733:A:O4'	2.20	0.41
34:YA:2749:A:H3'	34:YA:2750:A:H2'	2.02	0.41
34:YA:2819:G:H2'	34:YA:2821:A:N7	2.35	0.41
37:YE:24:THR:HG22	37:YE:186:GLY:H	1.84	0.41
37:YE:201:THR:OG1	37:YE:202:LYS:N	2.53	0.41
44:YP:15:ARG:HD3	44:YP:15:ARG:HA	1.85	0.41
44:YP:47:ASP:OD2	44:YP:50:ARG:NH1	2.53	0.41
1:QA:99:C:H2'	1:QA:101:A:C8	2.55	0.41
1:QA:120:A:H2'	1:QA:122:G:C8	2.54	0.41
1:QA:138:G:H2'	1:QA:139:G:C8	2.54	0.41
1:QA:418:C:H2'	1:QA:419:C:H6	1.85	0.41
1:QA:736:C:H2'	1:QA:737:A:C8	2.56	0.41
1:QA:1418:A:H1'	34:RA:1959:G:H1'	2.02	0.41
8:QH:81:HIS:ND1	8:QH:138:TRP:O	2.39	0.41
27:R3:4:LEU:HD21	27:R3:56:VAL:HB	2.03	0.41
34:RA:49:A:H61	34:RA:177:G:H2'	1.84	0.41
34:RA:523:C:H5''	34:RA:541:C:O2'	2.19	0.41
34:RA:628:G:O2'	34:RA:651:G:O2'	2.34	0.41
34:RA:1069:A:H4'	34:RA:1070:A:H5''	2.02	0.41
34:RA:2084:C:H2'	34:RA:2085:C:C6	2.55	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:2524:G:H21	34:RA:2741:A:H1'	1.84	0.41
34:RA:2881:C:H2'	34:RA:2882:A:H8	1.85	0.41
45:RQ:116:GLU:O	45:RQ:120:ILE:HG12	2.20	0.41
1:XA:59:A:H1'	1:XA:354:G:N2	2.35	0.41
1:XA:196:A:N3	1:XA:222:U:H1'	2.35	0.41
1:XA:435:C:H2'	1:XA:436:C:C6	2.53	0.41
1:XA:571:U:H5''	1:XA:819:A:C5	2.56	0.41
1:XA:861:G:O2'	1:XA:874:G:O2'	2.24	0.41
1:XA:913:A:OP1	12:XL:91:LYS:HE2	2.21	0.41
1:XA:1230:C:C5	13:XM:105:THR:HB	2.55	0.41
1:XA:1301:U:H4'	13:XM:17:VAL:HG23	2.02	0.41
1:XA:1347:G:HO2'	1:XA:1373:G:H1	1.67	0.41
2:XB:115:LEU:HB2	2:XB:145:LEU:HD23	2.02	0.41
3:XC:35:GLU:HB3	3:XC:59:ARG:HH22	1.85	0.41
13:XM:3:ARG:HA	13:XM:3:ARG:HD3	1.96	0.41
14:YN:24:CYS:SG	14:YN:40:CYS:N	2.93	0.41
29:Y5:25:LEU:HG	51:YW:19:LEU:HD12	2.03	0.41
34:YA:222:A:N6	34:YA:232:G:H1'	2.34	0.41
34:YA:303:U:H2'	34:YA:304:G:H8	1.85	0.41
34:YA:534:U:H2'	34:YA:535:C:C6	2.55	0.41
34:YA:629:G:O2'	34:YA:649:G:N2	2.54	0.41
34:YA:911:A:N1	45:YQ:9:TYR:HB3	2.35	0.41
34:YA:1187:G:N2	34:YA:1188:U:O4	2.51	0.41
34:YA:1566:A:N3	36:YD:214:TRP:CE3	2.88	0.41
41:YI:50:ARG:HA	41:YI:50:ARG:HD3	1.80	0.41
41:YI:121:LYS:HA	41:YI:121:LYS:HD3	1.67	0.41
46:YR:104:ARG:HD2	46:YR:109:ALA:HB3	2.02	0.41
53:YY:102:CYS:SG	53:YY:103:GLY:N	2.93	0.41
54:YZ:185:GLU:O	54:YZ:186:GLU:C	2.58	0.41
1:QA:108:G:N2	1:QA:108:G:OP2	2.49	0.41
1:QA:329:A:H2'	1:QA:332:G:N7	2.35	0.41
1:QA:583:A:H2'	1:QA:584:G:O4'	2.21	0.41
1:QA:777:A:H2'	1:QA:778:G:O4'	2.21	0.41
1:QA:974:A:P	14:QN:31:ARG:HB2	2.56	0.41
1:QA:1065:U:C5	1:QA:1190:G:H1'	2.56	0.41
1:QA:1332:A:H3'	1:QA:1333:A:C8	2.55	0.41
1:QA:1477:C:H2'	1:QA:1478:C:C6	2.56	0.41
1:QA:1517:G:H2'	1:QA:1518:A:O4'	2.21	0.41
3:QC:140:ARG:HA	3:QC:143:GLU:HG2	2.02	0.41
8:QH:3:THR:OG1	8:QH:4:ASP:N	2.54	0.41
29:R5:33:CYS:SG	29:R5:46:CYS:HB2	2.60	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:199:A:N1	34:RA:2433:A:H2'	2.36	0.41
34:RA:475:U:H2'	34:RA:476:G:O4'	2.21	0.41
34:RA:1046:A:H5'	34:RA:1047:G:H8	1.85	0.41
34:RA:1053:C:H2'	34:RA:1054:A:C8	2.55	0.41
34:RA:1129:A:H1'	34:RA:2516:G:H1'	2.01	0.41
34:RA:1221:C:H2'	34:RA:1222:C:H6	1.85	0.41
34:RA:2250:G:C2	45:RQ:82:ARG:HG3	2.55	0.41
34:RA:2748:A:O2'	40:RH:63:SER:O	2.20	0.41
36:RD:268:ARG:HG3	36:RD:269:PHE:CD2	2.55	0.41
46:RR:13:HIS:O	46:RR:17:ARG:HB2	2.20	0.41
50:RV:32:THR:O	50:RV:32:THR:OG1	2.37	0.41
1:XA:657:G:O2'	15:XO:23:GLY:HA2	2.20	0.41
1:XA:762:C:H2'	1:XA:763:G:C8	2.55	0.41
1:XA:1209:C:O2'	1:XA:1214:C:N4	2.53	0.41
1:XA:1499:A:H1'	1:XA:1520:G:H4'	2.03	0.41
6:XF:12:PRO:HD3	6:XF:58:GLY:HA2	2.02	0.41
22:XV:70:C:O2'	34:YA:1893:C:O2'	2.30	0.41
34:YA:409:C:H2'	34:YA:410:G:H8	1.85	0.41
34:YA:822:U:C5	34:YA:944:G:H1'	2.55	0.41
34:YA:840:C:H2'	34:YA:841:A:C8	2.55	0.41
34:YA:919:G:N1	34:YA:2268:A:C4	2.88	0.41
34:YA:1688:U:H1'	34:YA:1701:A:C6	2.55	0.41
34:YA:2103:C:H2'	34:YA:2104:G:C8	2.56	0.41
34:YA:2134:A:H3'	34:YA:2135:A:C8	2.53	0.41
34:YA:2286:A:H1'	34:YA:2287:A:C5	2.55	0.41
34:YA:2728:U:H2'	34:YA:2729:G:C8	2.55	0.41
37:YE:132:HIS:CD2	37:YE:132:HIS:O	2.73	0.41
1:QA:343:U:H2'	1:QA:345:C:C4	2.55	0.41
1:QA:782:A:H3'	1:QA:783:C:C6	2.55	0.41
1:QA:801:U:H2'	1:QA:802:A:C8	2.55	0.41
1:QA:1187:G:N2	14:QN:60:SER:HG	2.16	0.41
1:QA:1314:C:C5	19:QS:6:LYS:NZ	2.89	0.41
4:QD:28:SER:HA	4:QD:29:PRO:HD3	1.96	0.41
4:QD:169:LYS:CG	6:XF:82:ARG:HH22	2.33	0.41
13:QM:77:ASN:O	13:QM:81:LEU:HG	2.21	0.41
24:R0:43:THR:HG21	34:RA:2336:A:N6	2.36	0.41
32:R8:39:LYS:NZ	34:RA:2351:G:O6	2.49	0.41
34:RA:270(T):G:H2'	34:RA:270(U):G:C8	2.54	0.41
34:RA:797:C:H2'	34:RA:798:G:C8	2.56	0.41
34:RA:957:A:OP1	45:RQ:76:LYS:HG2	2.21	0.41
34:RA:1362:C:H2'	34:RA:1363:C:H6	1.85	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:1435:G:H2'	34:RA:1436:G:C8	2.56	0.41
34:RA:1589:C:H2'	34:RA:1590:U:C6	2.55	0.41
34:RA:1692:U:H3'	34:RA:1694:C:H41	1.86	0.41
34:RA:2544:G:H1'	34:RA:2646:C:H4'	2.02	0.41
44:RP:137:LYS:HA	44:RP:137:LYS:HD3	1.85	0.41
45:RQ:34:LEU:HD13	45:RQ:118:LEU:HD12	2.03	0.41
53:RY:99:CYS:HB3	53:RY:102:CYS:H	1.85	0.41
1:XA:40:C:H2'	1:XA:41:G:H8	1.86	0.41
1:XA:1466:C:H2'	1:XA:1467:G:O4'	2.21	0.41
5:XE:110:LEU:HB3	5:XE:115:VAL:HB	2.03	0.41
13:XM:87:TYR:OH	13:XM:91:ARG:NH1	2.53	0.41
32:Y8:4:MET:SD	34:YA:592:G:N2	2.89	0.41
34:YA:172:C:H2'	34:YA:173:G:C8	2.55	0.41
34:YA:270(D):C:H2'	34:YA:270(E):C:C6	2.55	0.41
34:YA:372:G:HO2'	34:YA:400:G:H1	1.59	0.41
34:YA:804:A:H2'	34:YA:806:C:C4	2.55	0.41
34:YA:1038:C:H2'	34:YA:1039:G:C8	2.56	0.41
34:YA:1361:G:N2	34:YA:1371:G:H1'	2.35	0.41
34:YA:1561:G:H2'	34:YA:1562:A:C8	2.55	0.41
34:YA:1790:C:H2'	34:YA:1791:A:C5	2.56	0.41
34:YA:2742:C:H2'	34:YA:2743:C:C6	2.56	0.41
39:YG:39:ILE:HG12	39:YG:157:ILE:HD12	2.02	0.41
54:YZ:6:LYS:O	54:YZ:62:PRO:HD3	2.20	0.41
1:QA:80:G:N2	1:QA:89:U:H1'	2.35	0.41
1:QA:186(A):C:C1'	20:QT:81:LYS:HE3	2.50	0.41
1:QA:553:A:O2'	12:QL:29:GLY:O	2.35	0.41
1:QA:870:U:H4'	1:QA:871:U:H3'	2.03	0.41
1:QA:1015:A:H2'	1:QA:1016:A:C8	2.56	0.41
1:QA:1203:C:H5'	14:QN:3:ARG:CZ	2.50	0.41
1:QA:1463:C:H2'	1:QA:1464:G:C8	2.56	0.41
1:QA:1479:C:H2'	1:QA:1480:G:C8	2.55	0.41
1:QA:1494:G:H4'	34:RA:1913:A:N7	2.35	0.41
3:QC:44:GLU:HA	3:QC:52:LEU:HD21	2.03	0.41
3:QC:76:VAL:HG21	3:QC:103:VAL:HG11	2.03	0.41
34:RA:576:U:H2'	34:RA:577:G:C8	2.56	0.41
34:RA:580:C:H2'	34:RA:581:C:C6	2.56	0.41
34:RA:822:U:H2'	34:RA:823:G:C8	2.56	0.41
34:RA:1588:C:H2'	34:RA:1589:C:C6	2.56	0.41
34:RA:1927:A:H2'	34:RA:1928:A:H8	1.86	0.41
34:RA:2014:A:O3'	51:RW:92:ARG:NH2	2.53	0.41
34:RA:2109:U:H2'	34:RA:2110:G:C8	2.56	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:2226:C:H2'	34:RA:2227:A:O4'	2.20	0.41
34:RA:2537:U:H2'	34:RA:2538:C:C6	2.56	0.41
37:RE:134:ILE:HD12	37:RE:137:HIS:HB2	2.02	0.41
39:RG:63:ILE:HD13	39:RG:141:PHE:HB3	2.02	0.41
44:RP:41:ARG:HD3	44:RP:41:ARG:HA	1.82	0.41
45:RQ:31:ASP:OD2	45:RQ:134:ARG:NH2	2.54	0.41
48:RT:66:VAL:HA	48:RT:71:GLY:HA2	2.02	0.41
49:RU:27:LEU:O	49:RU:31:SER:HB3	2.20	0.41
54:RZ:5:LEU:H	54:RZ:59:LEU:HA	1.84	0.41
1:XA:254:G:OP2	17:XQ:67:LYS:HB3	2.20	0.41
1:XA:451:A:C6	1:XA:480:U:H2'	2.55	0.41
1:XA:646:U:H2'	1:XA:647:C:C6	2.55	0.41
1:XA:649:G:H2'	1:XA:650:G:H8	1.84	0.41
1:XA:769:G:N2	1:XA:811:C:H1'	2.35	0.41
1:XA:1073:U:H3	1:XA:1102:A:H61	1.69	0.41
1:XA:1158:C:O3'	2:XB:133:LYS:HD3	2.19	0.41
4:XD:161:ASN:O	4:XD:165:MET:HB2	2.21	0.41
14:XN:47:LEU:HD22	14:XN:52:GLN:HB2	2.02	0.41
32:Y8:34:TRP:CD1	34:YA:2420:C:OP1	2.74	0.41
34:YA:6:A:H2'	34:YA:7:G:O4'	2.20	0.41
34:YA:69:C:H2'	34:YA:70:G:C8	2.55	0.41
34:YA:807:U:H2'	34:YA:808:G:C8	2.56	0.41
34:YA:1329:U:H5''	34:YA:1330:C:H5	1.86	0.41
34:YA:1336:A:H2'	34:YA:1337:G:C8	2.56	0.41
34:YA:1882:C:H3'	34:YA:1883:G:H8	1.86	0.41
35:YB:15:A:H1'	35:YB:109:G:C5	2.56	0.41
35:YB:37:C:N3	35:YB:48:A:O2'	2.50	0.41
42:YN:5:VAL:HA	42:YN:6:PRO:HD3	1.93	0.41
1:QA:250:A:H1'	1:QA:252:U:C6	2.56	0.41
1:QA:581:G:H2'	1:QA:582:U:C6	2.55	0.41
1:QA:739:C:HO2'	15:QO:42:HIS:CE1	2.37	0.41
1:QA:1013:G:N2	1:QA:1017:G:O6	2.54	0.41
1:QA:1052:U:H2'	1:QA:1200:C:N4	2.36	0.41
1:QA:1061:G:H2'	1:QA:1062:U:O4'	2.20	0.41
1:QA:1117:G:N3	1:QA:1180:A:H1'	2.35	0.41
1:QA:1202:G:H2'	1:QA:1202:G:N3	2.36	0.41
1:QA:1409:C:H2'	1:QA:1410:G:C8	2.55	0.41
4:QD:8:VAL:HG13	4:QD:21:LEU:HD12	2.02	0.41
25:R1:8:SER:HB3	25:R1:66:HIS:CD2	2.55	0.41
29:R5:43:HIS:NE2	34:RA:2884:U:OP2	2.50	0.41
29:R5:51:TYR:HD1	29:R5:55:ARG:O	2.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:98:G:H1'	34:RA:103:A:H1'	2.03	0.41
34:RA:272:G:H2'	34:RA:273(A):G:H8	1.86	0.41
34:RA:390:A:H4'	34:RA:391:G:H5'	2.02	0.41
34:RA:678:C:H2'	34:RA:679:C:C6	2.56	0.41
34:RA:691:C:H2'	34:RA:692:C:C6	2.56	0.41
34:RA:935:C:H2'	34:RA:936:C:C6	2.55	0.41
39:RG:33:ARG:H	39:RG:162:THR:HG22	1.85	0.41
40:RH:91:GLY:HA2	40:RH:160:LYS:HG2	2.03	0.41
42:RN:97:ARG:HA	42:RN:100:GLU:HB2	2.02	0.41
48:RT:8:LYS:HB2	48:RT:8:LYS:HE2	1.86	0.41
49:RU:58:ARG:HH11	49:RU:93:LYS:NZ	2.18	0.41
54:RZ:151:HIS:HB2	54:RZ:170:THR:HA	2.02	0.41
1:XA:130:A:OP2	17:XQ:63:ARG:HD3	2.20	0.41
1:XA:587:G:O2'	8:XH:3:THR:HA	2.20	0.41
1:XA:658:G:C1'	15:XO:22:THR:OG1	2.68	0.41
1:XA:990:C:H2'	1:XA:991:U:O4'	2.20	0.41
1:XA:1004:A:H2	1:XA:1024:G:H2'	1.85	0.41
1:XA:1259:C:HO2'	1:XA:1283:G:N2	2.17	0.41
1:XA:1349:A:H2'	1:XA:1350:A:O4'	2.20	0.41
1:XA:1367:C:H2'	1:XA:1368:G:H8	1.86	0.41
1:XA:1440(B):G:O2'	1:XA:1440(C):G:C6	2.72	0.41
6:XF:22:GLU:OE2	6:XF:84:ASN:ND2	2.40	0.41
10:XJ:16:LEU:HD22	10:XJ:94:VAL:HG22	2.02	0.41
29:Y5:11:THR:HG23	29:Y5:15:ARG:HD2	2.02	0.41
34:YA:318:C:H2'	34:YA:319:C:H6	1.84	0.41
34:YA:444:C:H2'	34:YA:445:C:C6	2.56	0.41
34:YA:755:C:H2'	34:YA:756:C:C6	2.55	0.41
34:YA:1130:U:C2	34:YA:2025:C:H5''	2.55	0.41
34:YA:1843:C:H2'	34:YA:1844:C:C6	2.56	0.41
34:YA:1909:C:H2'	34:YA:1910:G:C8	2.56	0.41
34:YA:2036:C:H2'	34:YA:2037:G:C8	2.55	0.41
34:YA:2073:C:H2'	34:YA:2074:U:H6	1.86	0.41
36:YD:231:HIS:CD2	36:YD:249:PRO:HG3	2.56	0.41
37:YE:104:VAL:HG11	37:YE:188:VAL:HG12	2.03	0.41
44:YP:37:GLY:O	44:YP:40:SER:OG	2.36	0.41
48:YT:3:ARG:O	48:YT:7:ILE:HG12	2.21	0.41
50:YV:68:LYS:HD3	50:YV:68:LYS:HA	1.74	0.41
1:QA:55:A:H61	1:QA:357:G:H2'	1.86	0.41
1:QA:663:A:H2'	1:QA:664:G:C8	2.56	0.41
1:QA:958:A:N6	19:QS:77:THR:C	2.71	0.41
3:QC:73:PRO:HA	3:QC:76:VAL:HG22	2.01	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
3:QC:120:VAL:O	3:QC:124:ILE:HG12	2.20	0.41
4:QD:27:TYR:O	6:XF:13:ASN:ND2	2.54	0.41
4:QD:166:LYS:HG2	36:YD:135:PHE:CZ	2.55	0.41
7:QG:87:VAL:HG23	7:QG:151:TYR:HB3	2.02	0.41
10:QJ:53:PRO:CB	14:QN:42:ILE:CD1	2.99	0.41
28:R4:26:SER:OG	39:RG:143:GLU:CD	2.52	0.41
34:RA:445:C:H2'	34:RA:446:G:C8	2.55	0.41
34:RA:568:U:H2'	34:RA:570:G:N7	2.36	0.41
34:RA:740:U:H2'	34:RA:741:G:H8	1.84	0.41
34:RA:903:C:H2'	34:RA:904:C:H6	1.85	0.41
34:RA:1076:C:C4	34:RA:1077:A:H1'	2.56	0.41
34:RA:1381:G:H1'	34:RA:1571:A:N1	2.36	0.41
34:RA:1695:G:H1'	36:RD:8:PRO:O	2.20	0.41
34:RA:1908:C:H2'	34:RA:1909:C:C6	2.56	0.41
34:RA:2185:C:H2'	34:RA:2186:G:H8	1.84	0.41
46:RR:96:ARG:HB2	46:RR:117:VAL:HG12	2.02	0.41
1:XA:125:U:H2'	1:XA:126:G:C8	2.56	0.41
1:XA:309:G:H2'	1:XA:310:G:C8	2.56	0.41
1:XA:421:U:C4'	3:XC:192:THR:HG22	2.46	0.41
1:XA:954:G:H22	1:XA:1228:C:H42	1.67	0.41
1:XA:1124:G:H4'	10:XJ:36:GLY:CA	2.51	0.41
1:XA:1440(I):U:H4'	1:XA:1440(J):A:C6	2.56	0.41
2:XB:75:LYS:HA	2:XB:75:LYS:HD3	1.86	0.41
2:XB:88:ALA:O	2:XB:226:ARG:NH1	2.46	0.41
2:XB:193:ASP:N	2:XB:193:ASP:OD2	2.41	0.41
3:XC:23:TYR:CE2	10:XJ:95:GLU:CG	3.04	0.41
9:XI:10:ARG:CZ	9:XI:105:ASP:HB2	2.51	0.41
10:XJ:40:LEU:N	10:XJ:40:LEU:CD1	2.84	0.41
11:XK:108:ILE:HD13	18:XR:87:ARG:NH2	2.36	0.41
24:Y0:72:ARG:HB3	24:Y0:75:LEU:HB2	2.02	0.41
25:Y1:3:LYS:HB3	34:YA:1364:G:OP1	2.20	0.41
34:YA:7:G:H2'	34:YA:8:A:C8	2.55	0.41
34:YA:312:G:H4'	34:YA:331:A:N3	2.35	0.41
34:YA:2825:C:H2'	34:YA:2826:A:O4'	2.21	0.41
34:YA:2832:U:H1'	34:YA:2834:G:C5	2.55	0.41
36:YD:44:ASN:OD1	36:YD:44:ASN:N	2.53	0.41
43:YO:70:LYS:HE2	43:YO:70:LYS:HB3	1.81	0.41
1:QA:103:C:OP2	20:QT:14:LYS:HD2	2.21	0.41
1:QA:488:C:H2'	1:QA:489:C:C6	2.56	0.41
1:QA:573:A:H2'	1:QA:574:A:C8	2.55	0.41
1:QA:619:U:H2'	1:QA:620:C:C6	2.56	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:648:A:H2'	1:QA:649:G:H8	1.85	0.41
1:QA:684:A:H2'	1:QA:685:G:C8	2.56	0.41
1:QA:955:U:C2'	19:QS:83:HIS:CA	2.95	0.41
1:QA:1238:A:OP1	1:QA:1335:C:O2'	2.35	0.41
1:QA:1240:U:C2'	7:QG:38:LEU:CD1	2.87	0.41
1:QA:1364:U:C1'	21:QU:14:TRP:HZ2	2.34	0.41
1:QA:1407:C:H2'	1:QA:1408:A:C8	2.56	0.41
2:QB:20:GLU:HG3	2:QB:191:ASP:HB2	2.02	0.41
2:QB:231:GLU:HA	2:QB:232:PRO:HD3	1.87	0.41
3:QC:36:ASP:OD1	3:QC:59:ARG:NH2	2.53	0.41
3:QC:79:ARG:CZ	11:XK:104:GLN:CA	2.88	0.41
5:QE:11:ILE:HG21	5:QE:31:LEU:HD23	2.02	0.41
13:QM:40:ASN:HA	13:QM:41:PRO:HD3	1.92	0.41
14:QN:12:ARG:HE	14:QN:14:PRO:HD2	1.86	0.41
25:R1:80:LEU:HD12	25:R1:81:LYS:HB2	2.03	0.41
28:R4:27:THR:HG21	39:RG:62:LEU:HD13	2.03	0.41
32:R8:3:LYS:H	32:R8:3:LYS:HG2	1.71	0.41
34:RA:47:C:H2'	34:RA:48:G:H8	1.86	0.41
34:RA:256:A:H2'	34:RA:257:A:C8	2.56	0.41
34:RA:455:C:N3	34:RA:472:A:H2'	2.36	0.41
34:RA:456:C:C5	52:RX:69:TYR:CE1	3.09	0.41
34:RA:863:A:O2'	35:RB:100:G:N3	2.49	0.41
34:RA:923:C:H2'	34:RA:924:C:C6	2.55	0.41
34:RA:1303:G:H5'	34:RA:1642:G:H21	1.85	0.41
34:RA:1579:A:H2'	34:RA:1580:A:C8	2.55	0.41
34:RA:1793:C:H2'	34:RA:1794:U:C6	2.56	0.41
34:RA:1991:U:H2'	34:RA:1992:G:H5''	2.03	0.41
34:RA:2085:C:H4'	36:RD:262:ARG:NH2	2.35	0.41
34:RA:2134:A:H1'	34:RA:2158:A:N3	2.35	0.41
34:RA:2205:C:O2	34:RA:2226:C:N4	2.54	0.41
34:RA:2385:C:H2'	34:RA:2386:C:C6	2.56	0.41
34:RA:2428:G:O2'	44:RP:56:SER:OG	2.36	0.41
34:RA:2440:C:H2'	34:RA:2441:C:H4'	2.03	0.41
34:RA:2470:G:OP1	45:RQ:56:ARG:NH2	2.48	0.41
34:RA:2470:G:P	45:RQ:56:ARG:HH21	2.43	0.41
34:RA:2497:A:H1'	34:RA:2498:C:H5	1.86	0.41
34:RA:2764:A:H2'	34:RA:2766:G:C8	2.56	0.41
39:RG:141:PHE:HB2	39:RG:144:ILE:HD13	2.03	0.41
40:RH:123:PHE:HD2	40:RH:123:PHE:HA	1.77	0.41
41:RI:14:ASP:HB3	41:RI:15:VAL:H	1.65	0.41
49:RU:39:LEU:HD23	49:RU:39:LEU:HA	1.93	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:XA:240:C:H2'	1:XA:241:C:C6	2.56	0.41
1:XA:354:G:N3	1:XA:354:G:H2'	2.35	0.41
1:XA:668:G:O2'	15:XO:46:HIS:ND1	2.47	0.41
1:XA:735:C:H2'	1:XA:736:C:C6	2.55	0.41
1:XA:802:A:H3'	1:XA:803:G:H8	1.86	0.41
1:XA:1293:G:H2'	1:XA:1294:G:C8	2.56	0.41
1:XA:1320:C:N4	19:XS:37:ARG:N	2.69	0.41
2:XB:40:HIS:HB2	2:XB:190:THR:HG21	2.01	0.41
12:XL:56:ALA:HB2	12:XL:70:ILE:HD11	2.03	0.41
14:YN:56:VAL:CG1	14:YN:57:ARG:N	2.59	0.41
15:XO:26:GLU:HG3	15:XO:81:LEU:HD22	2.03	0.41
18:XR:30:ASP:OD2	18:XR:33:ASP:N	2.54	0.41
20:XT:18:GLN:HE21	20:XT:22:ARG:NH2	2.19	0.41
28:Y4:31:ILE:HG21	39:YG:142:PRO:HB2	2.03	0.41
34:YA:375:C:H2'	34:YA:376:C:C6	2.56	0.41
34:YA:459:U:H2'	34:YA:460:A:H8	1.85	0.41
34:YA:558:G:H2'	34:YA:559:G:H8	1.85	0.41
34:YA:558:G:P	42:YN:111:PRO:HD2	2.61	0.41
34:YA:632:A:H2'	34:YA:633:A:C8	2.56	0.41
34:YA:665:C:H2'	34:YA:666:G:C8	2.56	0.41
34:YA:748:G:OP1	51:YW:88:ARG:NH2	2.40	0.41
34:YA:923:C:H2'	34:YA:924:C:C6	2.56	0.41
34:YA:1050:A:N7	34:YA:2751:G:N3	2.67	0.41
34:YA:1056:G:H5''	34:YA:1057:A:H5'	2.02	0.41
34:YA:1502:C:H2'	34:YA:1503:U:C6	2.56	0.41
34:YA:1503:U:H2'	34:YA:1504:C:C6	2.56	0.41
34:YA:1510:A:O2'	34:YA:1512:G:N7	2.50	0.41
34:YA:1914:C:H2'	34:YA:1915:U:O4'	2.21	0.41
34:YA:2059:A:C6	34:YA:2503:A:C6	3.05	0.41
34:YA:2134:A:H1'	34:YA:2158:A:C2	2.56	0.41
34:YA:2363:C:H2'	34:YA:2364:C:H6	1.86	0.41
34:YA:2402:C:H1'	34:YA:2403:C:C5	2.50	0.41
34:YA:2413:G:O2'	44:YP:70:GLN:NE2	2.48	0.41
34:YA:2503:A:O2'	34:YA:2505:G:OP2	2.23	0.41
34:YA:2820:A:H1'	46:YR:3:HIS:CE1	2.56	0.41
35:YB:111:U:H2'	35:YB:112:G:H8	1.85	0.41
36:YD:61:LEU:HD23	36:YD:61:LEU:HA	1.94	0.41
37:YE:154:LYS:HD2	37:YE:154:LYS:HA	1.86	0.41
38:YF:157:VAL:HG21	38:YF:181:LEU:HD13	2.03	0.41
40:YH:86:GLU:H	40:YH:86:GLU:HG2	1.51	0.41
44:YP:6:LEU:HD23	44:YP:6:LEU:HA	1.90	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
45:YQ:55:VAL:HG13	54:YZ:178:GLU:HB3	2.03	0.41
48:YT:19:LEU:HA	48:YT:20:PRO:HD3	1.91	0.41
52:YX:34:ALA:O	52:YX:77:LYS:NZ	2.54	0.41
52:YX:92:LEU:HD23	52:YX:92:LEU:HA	1.86	0.41
54:YZ:4:ARG:NH1	54:YZ:60:GLU:OE2	2.54	0.41
54:YZ:5:LEU:HB3	54:YZ:6:LYS:H	1.72	0.41
54:YZ:94:GLU:HA	54:YZ:95:PRO:HD2	1.75	0.41
1:QA:10:A:H2'	1:QA:11:G:H8	1.85	0.41
1:QA:65:U:O4'	1:QA:199:G:O2'	2.31	0.41
1:QA:254:G:OP1	17:QQ:67:LYS:O	2.39	0.41
1:QA:411:A:H2'	1:QA:413:G:C8	2.55	0.41
1:QA:692:U:H5'	1:QA:797:C:H5''	2.02	0.41
1:QA:818:G:H1'	1:QA:820:U:C4	2.55	0.41
1:QA:958:A:N7	19:QS:79:THR:HG21	2.36	0.41
1:QA:1026:G:H2'	1:QA:1027:C:O4'	2.20	0.41
1:QA:1080:A:H5'	5:QE:14:ARG:NH2	2.36	0.41
1:QA:1298:C:H6	1:QA:1298:C:H2'	1.55	0.41
3:QC:23:TYR:CD1	10:QJ:10:GLY:HA2	2.56	0.41
12:QL:93:LEU:HA	12:QL:94:PRO:HD3	1.96	0.41
25:R1:86:SER:O	25:R1:88:LYS:N	2.53	0.41
34:RA:713:G:H2'	34:RA:714:U:C6	2.56	0.41
34:RA:795:C:H2'	34:RA:796:C:C6	2.56	0.41
34:RA:1011:G:OP2	49:RU:70:ARG:NH2	2.54	0.41
34:RA:1468:C:H2'	34:RA:1469:A:C8	2.55	0.41
34:RA:1775:U:N3	34:RA:1776:G:H1'	2.36	0.41
34:RA:2345:G:H4'	34:RA:2346:A:H5'	2.03	0.41
36:RD:35:LYS:HB3	36:RD:63:ARG:HA	2.03	0.41
41:RI:143:SER:HB2	41:RI:144:VAL:H	1.64	0.41
44:RP:28:GLY:C	44:RP:30:THR:H	2.24	0.41
47:RS:50:SER:O	47:RS:76:LYS:NZ	2.41	0.41
49:RU:58:ARG:O	49:RU:62:ILE:HG12	2.20	0.41
50:RV:73:SER:OG	50:RV:74:LYS:N	2.53	0.41
54:RZ:24:LEU:HA	54:RZ:25:PRO:HD3	1.92	0.41
1:XA:22:G:H4'	1:XA:885:G:C8	2.56	0.41
1:XA:171:A:H2'	1:XA:172:A:C8	2.56	0.41
1:XA:299:G:H2'	1:XA:300:A:C8	2.56	0.41
1:XA:323:U:H3'	1:XA:324:G:H8	1.84	0.41
1:XA:528:C:N4	12:XL:49:ASN:CG	2.70	0.41
1:XA:1165:C:H2'	1:XA:1166:G:C8	2.55	0.41
1:XA:1314:C:H6	1:XA:1314:C:O5'	2.04	0.41
1:XA:1391:U:H2'	1:XA:1392:G:H8	1.85	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:XJ:90:LEU:HD12	10:XJ:90:LEU:HA	1.90	0.41
19:XS:19:VAL:HA	19:XS:22:LEU:HB2	2.01	0.41
34:YA:709:U:H2'	34:YA:710:G:C8	2.56	0.41
34:YA:881:G:H2'	34:YA:882:G:C8	2.56	0.41
34:YA:1123:C:H2'	34:YA:1124:C:C6	2.56	0.41
34:YA:1252:G:N2	49:YU:33:ARG:HD3	2.36	0.41
34:YA:1836:C:H2'	34:YA:1837:C:C6	2.56	0.41
34:YA:1949:G:H3'	34:YA:1950:G:H21	1.86	0.41
34:YA:2850:A:H3'	34:YA:2851:A:H8	1.86	0.41
46:YR:56:LYS:NZ	46:YR:90:ARG:O	2.53	0.41
48:YT:130:ALA:HA	48:YT:133:GLU:HG2	2.03	0.41
1:QA:296:U:O2'	1:QA:556:C:O2	2.27	0.40
1:QA:542:G:P	4:QD:10:ARG:HH12	2.44	0.40
1:QA:709:G:H2'	1:QA:710:G:O4'	2.22	0.40
1:QA:756:C:H2'	1:QA:757:U:C6	2.56	0.40
1:QA:947:G:H5'	13:QM:109:THR:CG2	2.50	0.40
1:QA:956:U:H5'	19:QS:83:HIS:CA	2.51	0.40
1:QA:986:A:H2'	1:QA:987:G:H8	1.85	0.40
1:QA:996:A:N1	1:QA:1046:A:H1'	2.36	0.40
1:QA:1152:A:H3'	1:QA:1153:C:C6	2.56	0.40
1:QA:1278:U:H4'	1:QA:1279:A:C8	2.56	0.40
1:QA:1409:C:H2'	1:QA:1410:G:H8	1.86	0.40
1:QA:1498:U:H1'	1:QA:1499:A:N7	2.36	0.40
2:QB:167:PRO:O	2:QB:171:ALA:CB	2.67	0.40
13:QM:84:ILE:O	19:QS:74:PHE:CD2	2.57	0.40
20:QT:74:LYS:O	20:QT:76:ALA:N	2.55	0.40
34:RA:271(B):C:O2	34:RA:272:G:H1'	2.20	0.40
34:RA:2469:A:H2	34:RA:2481:G:H21	1.69	0.40
34:RA:2790:A:H2	34:RA:2791:C:H2'	1.86	0.40
34:RA:2820:A:C5	46:RR:4:LEU:HD12	2.56	0.40
35:RB:15:A:OP2	35:RB:69:G:N2	2.54	0.40
37:RE:144:ARG:HB3	37:RE:145:LYS:H	1.66	0.40
40:RH:126:PRO:HD2	40:RH:131:VAL:HA	2.03	0.40
44:RP:144:GLU:HA	44:RP:145:PRO:HD3	1.83	0.40
49:RU:34:LYS:HA	49:RU:34:LYS:HD3	1.72	0.40
1:XA:17:U:H2'	1:XA:18:C:C6	2.56	0.40
1:XA:864:A:H2	1:XA:918:A:H1'	1.86	0.40
1:XA:1241:G:H2'	1:XA:1242:C:C6	2.56	0.40
1:XA:1305:G:H2'	1:XA:1331:G:H21	1.74	0.40
10:XJ:5:ARG:N	10:XJ:99:LYS:O	2.54	0.40
11:XK:109:VAL:HG13	18:XR:86:VAL:HB	2.03	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
16:XP:72:ARG:HA	16:XP:75:ARG:HG2	2.03	0.40
34:YA:71:A:H5''	34:YA:72:U:H3'	2.02	0.40
34:YA:363(D):G:H2'	34:YA:363(E):G:H8	1.85	0.40
34:YA:1128:A:N6	34:YA:2518:A:C6	2.89	0.40
34:YA:1166:C:H2'	34:YA:1167:U:C6	2.56	0.40
34:YA:1174:A:H61	55:YA:3192:MG:MG	1.10	0.40
34:YA:1781:C:H4'	34:YA:1782:C:C5'	2.51	0.40
34:YA:2090:G:N3	34:YA:2230:G:C2	2.89	0.40
34:YA:2859:G:H2'	34:YA:2860:A:C8	2.56	0.40
36:YD:172:TYR:HB3	36:YD:184:LYS:HG2	2.04	0.40
37:YE:12:THR:HG23	37:YE:13:ARG:N	2.36	0.40
38:YF:116:ASP:OD1	38:YF:119:ARG:NH2	2.52	0.40
41:YI:143:SER:HB2	41:YI:144:VAL:H	1.60	0.40
43:YO:71:ARG:HH21	43:YO:77:ILE:HG21	1.86	0.40
1:QA:28:G:H2'	1:QA:296:U:H5''	2.02	0.40
1:QA:385:C:H2'	1:QA:386:C:C6	2.55	0.40
1:QA:665:A:N3	1:QA:732:C:H2'	2.36	0.40
1:QA:767:A:H1'	1:QA:1525:G:H1'	2.02	0.40
1:QA:968:A:N3	1:QA:1197:G:H1'	2.37	0.40
1:QA:980:C:H1'	14:QN:19:ARG:HG2	0.41	0.40
1:QA:1051:C:H2'	1:QA:1052:U:C2	2.57	0.40
1:QA:1107:C:H5''	3:QC:173:VAL:N	2.32	0.40
1:QA:1221:G:OP1	1:QA:1321:C:N4	2.54	0.40
1:QA:1431:C:H2'	1:QA:1432:G:O4'	2.21	0.40
4:QD:205:GLU:CD	5:QE:107:ARG:NH1	2.74	0.40
12:QL:32:PHE:HB3	12:QL:84:LEU:HD21	2.03	0.40
17:QQ:75:ARG:HA	17:QQ:75:ARG:HD2	1.88	0.40
24:R0:60:PHE:CZ	34:RA:2365:G:H4'	2.56	0.40
34:RA:234:C:H2'	34:RA:235:U:H6	1.86	0.40
34:RA:997:G:O5'	49:RU:58:ARG:NH1	2.55	0.40
34:RA:1050:A:H2'	34:RA:1051:G:O4'	2.21	0.40
34:RA:2011:U:H2'	34:RA:2012:G:O4'	2.20	0.40
36:RD:134:ARG:N	36:RD:187:GLY:O	2.54	0.40
40:RH:102:ALA:HA	40:RH:117:PRO:HD3	2.03	0.40
40:RH:103:LEU:O	40:RH:114:VAL:HA	2.22	0.40
1:XA:136:C:H42	1:XA:227:G:H1	1.68	0.40
1:XA:373:A:C2'	1:XA:374:A:H5'	2.52	0.40
1:XA:1019:C:H2'	1:XA:1020:U:C6	2.56	0.40
1:XA:1051:C:H2'	1:XA:1052:U:C6	2.56	0.40
1:XA:1300:G:N1	1:XA:1334:G:C6	2.89	0.40
2:XB:134:GLU:HG3	2:XB:137:ARG:NE	2.37	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:XB:188:ALA:HB1	2:XB:192:SER:HB3	2.03	0.40
4:XD:170:VAL:HB	4:XD:174:LEU:HD12	2.02	0.40
13:XM:26:GLY:O	13:XM:30:ALA:HB2	2.20	0.40
23:XX:8:A:H2'	23:XX:9:G:C8	2.55	0.40
24:Y0:23:VAL:HA	24:Y0:38:VAL:HG12	2.03	0.40
27:Y3:17:LYS:HB2	27:Y3:17:LYS:HE3	1.88	0.40
34:YA:452:G:N2	34:YA:457:A:O2'	2.55	0.40
34:YA:601:C:H2'	34:YA:602:G:O4'	2.21	0.40
34:YA:620:G:H4'	34:YA:621:A:H5'	2.03	0.40
34:YA:797:C:H2'	34:YA:798:G:C8	2.57	0.40
34:YA:839:U:H2'	34:YA:840:C:H6	1.85	0.40
34:YA:844:C:H3'	34:YA:845:G:H8	1.87	0.40
34:YA:849:A:H61	34:YA:929:G:H1'	1.86	0.40
34:YA:1080:C:H2'	34:YA:1081:U:H6	1.86	0.40
34:YA:1087:G:C8	34:YA:1089:G:H1'	2.56	0.40
34:YA:1158:C:H2'	34:YA:1159:U:C6	2.57	0.40
34:YA:1169:G:H2'	34:YA:1170:G:O4'	2.21	0.40
34:YA:2091:U:H2'	34:YA:2092:U:C5	2.57	0.40
34:YA:2537:U:H2'	34:YA:2538:C:C6	2.55	0.40
35:YB:5:C:O2'	35:YB:27:C:O2	2.39	0.40
40:YH:115:VAL:HG21	40:YH:148:ILE:HD11	2.02	0.40
48:YT:80:SER:HA	48:YT:81:PRO:HD3	1.95	0.40
48:YT:96:ARG:HE	48:YT:96:ARG:HB3	1.72	0.40
53:YY:12:THR:HA	53:YY:26:LYS:HA	2.03	0.40
54:YZ:14:LYS:HA	54:YZ:15:PRO:HD3	1.93	0.40
1:QA:95:G:H2'	1:QA:96:G:C8	2.56	0.40
1:QA:728:A:H2'	1:QA:729:A:C8	2.57	0.40
1:QA:741:G:H4'	15:QO:55:GLY:HA3	2.02	0.40
1:QA:955:U:C2'	19:QS:83:HIS:HA	2.52	0.40
1:QA:1049:U:H5''	1:QA:1050:G:C8	2.55	0.40
1:QA:1123:A:C1'	10:QJ:37:PRO:HD2	2.50	0.40
3:QC:56:ASP:HB3	3:QC:67:THR:HG23	2.02	0.40
9:QI:25:LYS:O	9:QI:60:ASP:HA	2.21	0.40
24:R0:43:THR:HG21	34:RA:2336:A:H61	1.85	0.40
34:RA:151:C:H2'	34:RA:152:G:H8	1.86	0.40
34:RA:486:C:H2'	34:RA:487:C:C6	2.56	0.40
34:RA:519:U:H2'	34:RA:520:G:C8	2.56	0.40
34:RA:729:G:P	36:RD:208:LYS:HZ3	2.45	0.40
34:RA:1186:G:H2'	34:RA:1187:G:O4'	2.21	0.40
34:RA:1561:G:H2'	34:RA:1562:A:C8	2.56	0.40
34:RA:1808:U:H2'	34:RA:1809:A:O4'	2.22	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:2348:U:H2'	34:RA:2349:G:C8	2.56	0.40
47:RS:59:LYS:HA	47:RS:59:LYS:HD2	1.94	0.40
53:RY:47:LYS:HA	53:RY:47:LYS:HD2	1.94	0.40
1:XA:68(A):G:H22	1:XA:101:A:H2	1.68	0.40
1:XA:225:C:H2'	1:XA:226:G:C8	2.56	0.40
1:XA:418:C:H2'	1:XA:419:C:H6	1.85	0.40
1:XA:707:C:H2'	1:XA:708:C:C6	2.56	0.40
1:XA:922:G:H1'	5:XE:19:MET:CB	2.50	0.40
1:XA:1035:A:H2'	1:XA:1036:G:H8	1.87	0.40
1:XA:1101:A:N3	1:XA:1102:A:H1'	2.37	0.40
1:XA:1118:C:H2'	1:XA:1119:C:C6	2.56	0.40
1:XA:1302:U:O2	13:XM:27:LYS:HE2	2.21	0.40
1:XA:1320:C:N4	19:XS:37:ARG:CA	2.60	0.40
1:XA:1359:C:H4'	1:XA:1362(A):C:H42	1.86	0.40
8:XH:34:GLU:O	8:XH:38:ILE:HG12	2.22	0.40
11:XK:51:LYS:HE2	11:XK:51:LYS:HB2	1.93	0.40
12:XL:70:ILE:HG13	12:XL:100:ILE:HG13	2.03	0.40
34:YA:38:A:H2'	34:YA:39:C:C6	2.57	0.40
34:YA:238:C:H1'	34:YA:609(A):A:H1'	2.03	0.40
34:YA:797:C:H2'	34:YA:798:G:H8	1.86	0.40
34:YA:887:A:H1'	34:YA:889:C:C5	2.56	0.40
34:YA:2292:C:H2'	34:YA:2293:C:C6	2.56	0.40
39:YG:95:ARG:HB3	39:YG:96:ARG:H	1.70	0.40
43:YO:67:LYS:HD2	43:YO:67:LYS:HA	1.87	0.40
50:YV:69:LYS:HA	50:YV:87:HIS:O	2.21	0.40
54:YZ:178:GLU:HB3	54:YZ:179:ASP:H	1.60	0.40
1:QA:44:G:OP1	16:QP:12:LYS:CD	2.64	0.40
1:QA:107:G:H4'	1:QA:378:G:H4'	2.02	0.40
1:QA:477:G:H2'	1:QA:478:A:C8	2.56	0.40
1:QA:668:G:O4'	15:QO:48:LYS:O	2.39	0.40
1:QA:1002:G:H3'	1:QA:1003:G:H8	1.85	0.40
1:QA:1179:A:H2'	1:QA:1180:A:H8	1.87	0.40
1:QA:1281:U:H5''	1:QA:1282:C:H5	1.87	0.40
1:QA:1285:A:N1	1:QA:1354:C:O2'	2.54	0.40
1:QA:1485:U:H2'	1:QA:1486:G:C8	2.55	0.40
2:QB:208:ILE:HA	2:QB:211:ILE:HG12	2.03	0.40
3:QC:12:LEU:HG	3:QC:18:TRP:NE1	2.36	0.40
10:QJ:14:LYS:HD3	10:QJ:15:THR:HG23	2.02	0.40
16:QP:14:ASN:OD1	16:QP:16:HIS:NE2	2.54	0.40
34:RA:67:U:H2'	34:RA:68:G:H8	1.87	0.40
34:RA:413:C:H2'	34:RA:414:C:H6	1.86	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:RA:741:G:H2'	34:RA:742:G:C8	2.56	0.40
34:RA:817:C:O2'	34:RA:932:G:N2	2.55	0.40
34:RA:1031:G:H2'	34:RA:1032:A:C8	2.57	0.40
34:RA:1556:C:H2'	34:RA:1557:C:C6	2.57	0.40
34:RA:1853:A:N1	34:RA:2087:G:H1'	2.36	0.40
34:RA:2019:A:H2	34:RA:2035:G:H22	1.68	0.40
34:RA:2453:A:H2'	34:RA:2454:G:C8	2.45	0.40
34:RA:2463:C:H2'	34:RA:2464:C:C6	2.56	0.40
34:RA:2767:C:H2'	34:RA:2768:C:C6	2.57	0.40
39:RG:132:ASN:HD22	39:RG:158:ALA:HA	1.86	0.40
46:RR:3:HIS:HB3	46:RR:4:LEU:H	1.64	0.40
53:RY:67:LEU:HD23	53:RY:67:LEU:HA	1.96	0.40
1:XA:162:A:N7	1:XA:163:C:H1'	2.36	0.40
1:XA:200:G:H2'	1:XA:201(A):C:O4'	2.22	0.40
1:XA:254:G:H5'	17:XQ:66:SER:CB	2.51	0.40
1:XA:280:C:N4	17:XQ:39:SER:H	2.15	0.40
1:XA:1424:C:H2'	1:XA:1425:U:H6	1.86	0.40
1:XA:1437:C:H2'	1:XA:1438:G:C8	2.57	0.40
10:XJ:48:THR:HG22	10:XJ:60:ARG:HD2	2.02	0.40
14:YN:7:ILE:HG22	14:YN:23:ARG:HE	1.87	0.40
14:YN:47:LEU:HA	14:YN:47:LEU:HD23	1.86	0.40
24:Y0:70:GLN:HG2	24:Y0:72:ARG:HG2	2.03	0.40
29:Y5:46:CYS:HB3	29:Y5:49:CYS:HB3	2.03	0.40
34:YA:390:A:C6	44:YP:71:VAL:HG21	2.57	0.40
34:YA:642:G:H21	34:YA:646:A:H2	1.69	0.40
34:YA:862:G:H2'	34:YA:863:A:O4'	2.20	0.40
34:YA:1689:A:H2'	34:YA:1690:A:C8	2.56	0.40
34:YA:1754:C:H5''	48:YT:113:LYS:HD3	2.03	0.40
34:YA:1820:U:N3	36:YD:202:LYS:HD2	2.37	0.40
34:YA:1830:C:H2'	34:YA:1831:G:H8	1.87	0.40
34:YA:1836:C:H2'	34:YA:1837:C:H6	1.86	0.40
34:YA:2043:C:H42	34:YA:2625:G:H1	1.69	0.40
34:YA:2096:U:C2	34:YA:2194:G:C2	3.10	0.40
34:YA:2122:U:H2'	34:YA:2123:G:C8	2.57	0.40
38:YF:195:ASP:OD1	38:YF:195:ASP:N	2.47	0.40
50:YV:100:ARG:HE	50:YV:100:ARG:HB2	1.70	0.40
53:YY:39:VAL:HG23	53:YY:42:VAL:HB	2.02	0.40
54:YZ:24:LEU:HA	54:YZ:25:PRO:HD3	1.87	0.40
54:YZ:128:VAL:HG22	54:YZ:161:VAL:HG22	2.03	0.40
1:QA:277:C:P	17:QQ:68:ARG:HH21	2.44	0.40
1:QA:343:U:H1'	1:QA:347:G:N2	2.36	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:QA:390:C:H2'	1:QA:391:G:C8	2.57	0.40
1:QA:450:G:H5''	1:QA:451:A:H3'	2.04	0.40
1:QA:912:C:H2'	1:QA:913:A:C8	2.56	0.40
1:QA:957:U:P	19:QS:80:TYR:O	2.77	0.40
1:QA:1229:A:H2'	1:QA:1230:C:C6	2.56	0.40
1:QA:1329:A:OP1	13:QM:29:ARG:HD3	2.22	0.40
2:QB:75:LYS:HA	2:QB:75:LYS:HD3	1.83	0.40
4:QD:88:VAL:CG1	5:QE:97:GLY:HA3	2.47	0.40
31:R7:7:PRO:HB2	34:RA:1309:G:H4'	2.04	0.40
32:R8:27:THR:HA	44:RP:62:LEU:HD22	2.04	0.40
33:R9:29:ASN:HA	33:R9:30:PRO:HD3	1.93	0.40
34:RA:151:C:H2'	34:RA:152:G:C8	2.57	0.40
34:RA:969:U:H2'	34:RA:970:C:C6	2.56	0.40
34:RA:1464:C:H2'	34:RA:1465:G:C8	2.57	0.40
34:RA:1509:C:H2'	34:RA:1511:A:C8	2.57	0.40
34:RA:1853:A:N3	34:RA:2233:U:O2'	2.49	0.40
34:RA:2145:C:H2'	34:RA:2147:G:C2	2.56	0.40
34:RA:2405:G:H1'	34:RA:2412:A:H61	1.87	0.40
41:RI:40:THR:OG1	41:RI:41:GLU:N	2.55	0.40
44:RP:64:LYS:O	44:RP:66:GLY:N	2.55	0.40
45:RQ:30:GLY:HA2	45:RQ:107:ALA:HB2	2.03	0.40
48:RT:52:ILE:HG13	48:RT:61:PHE:HB3	2.02	0.40
1:XA:59:A:H3'	1:XA:331:G:N2	2.33	0.40
1:XA:293:G:H5'	1:XA:610:G:H21	1.86	0.40
1:XA:690:G:H1'	1:XA:698:G:N2	2.36	0.40
1:XA:767:A:H1'	1:XA:1525:G:H1'	2.02	0.40
1:XA:860:A:H3'	1:XA:861:G:C8	2.54	0.40
1:XA:969:A:O2'	10:XJ:55:LYS:NZ	2.44	0.40
1:XA:978:A:HO2'	1:XA:1322:C:N4	2.19	0.40
1:XA:1114:C:H2'	1:XA:1115:C:C6	2.56	0.40
1:XA:1275:A:H2'	1:XA:1276:G:O4'	2.22	0.40
1:XA:1302:U:H1'	13:XM:27:LYS:HE2	0.96	0.40
1:XA:1332:A:H3'	1:XA:1333:A:H8	1.87	0.40
4:XD:202:LEU:HA	4:XD:205:GLU:HB2	2.03	0.40
6:XF:7:ASN:HD21	18:XR:35:ARG:CZ	2.34	0.40
8:XH:81:HIS:ND1	8:XH:138:TRP:O	2.33	0.40
22:XV:19:G:C8	34:YA:2112:G:N7	2.88	0.40
25:Y1:41:ARG:NH2	34:YA:1365:A:O5'	2.40	0.40
34:YA:197:A:C6	34:YA:2430:A:C2	3.08	0.40
34:YA:270(S):G:H2'	34:YA:270(T):G:C8	2.56	0.40
34:YA:363(F):U:H3'	34:YA:363(G):A:C8	2.56	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
34:YA:594:U:H2'	34:YA:595:C:C6	2.56	0.40
34:YA:1456:G:C2	34:YA:2704:C:N3	2.89	0.40
34:YA:1656:C:H2'	34:YA:1657:C:C6	2.55	0.40
34:YA:1869:G:H5'	34:YA:1870:C:OP2	2.22	0.40
34:YA:2053:G:N2	34:YA:2617:C:N3	2.69	0.40
34:YA:2065:C:H2'	34:YA:2066:C:C6	2.55	0.40
34:YA:2429:G:N7	44:YP:56:SER:OG	2.35	0.40
34:YA:2591:C:H2'	34:YA:2592:G:C8	2.56	0.40
34:YA:2746:U:O4	34:YA:2755:C:H4'	2.21	0.40
34:YA:2820:A:C6	46:YR:4:LEU:CD1	3.04	0.40
36:YD:175:LEU:O	36:YD:182:LEU:HA	2.22	0.40
39:YG:47:LYS:HD3	39:YG:81:LYS:HB2	2.04	0.40
42:YN:34:LEU:O	42:YN:49:GLY:HA3	2.21	0.40

All (35) symmetry-related close contacts are listed below. The label for Atom-2 includes the symmetry operator and encoded unit-cell translations to be applied.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:QG:149:ARG:NE	3:XC:81:GLY:O[4_555]	0.51	1.69
7:QG:149:ARG:NH1	3:XC:85:ARG:N[4_555]	1.02	1.18
7:QG:149:ARG:NH1	3:XC:85:ARG:CA[4_555]	1.16	1.04
7:QG:149:ARG:CZ	3:XC:85:ARG:N[4_555]	1.35	0.85
7:QG:149:ARG:NH1	3:XC:85:ARG:CB[4_555]	1.52	0.68
7:QG:149:ARG:NH2	3:XC:84:ILE:CA[4_555]	1.62	0.58
7:QG:149:ARG:NE	3:XC:81:GLY:C[4_555]	1.67	0.53
7:QG:149:ARG:CZ	3:XC:81:GLY:O[4_555]	1.67	0.53
7:QG:149:ARG:CD	3:XC:81:GLY:O[4_555]	1.70	0.50
7:QG:149:ARG:NH2	3:XC:85:ARG:N[4_555]	1.70	0.50
29:Y5:60:VAL:O	50:YV:49:THR:OG1[4_545]	1.72	0.48
7:QG:149:ARG:O	3:XC:80:GLY:O[4_555]	1.75	0.45
7:QG:149:ARG:NH2	3:XC:84:ILE:C[4_555]	1.77	0.43
9:QI:98:PRO:CB	10:XJ:79:ARG:NH1[4_555]	1.78	0.42
9:QI:98:PRO:CB	10:XJ:79:ARG:CZ[4_555]	1.81	0.39
7:QG:149:ARG:NH2	3:XC:84:ILE:N[4_555]	1.82	0.38
7:QG:149:ARG:NH2	3:XC:84:ILE:CG2[4_555]	1.83	0.37
36:RD:134:ARG:NH1	4:XD:163:GLU:O[4_555]	1.89	0.31
6:QF:20:ALA:CB	4:XD:195:ALA:CB[4_555]	1.92	0.28
9:QI:98:PRO:CA	10:XJ:79:ARG:NH1[4_555]	1.96	0.24
27:R3:3:ARG:NH2	44:YP:137:LYS:NZ[3_455]	1.96	0.24
6:QF:14:LEU:C	4:XD:20:TYR:OH[4_555]	1.98	0.22

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:QF:14:LEU:O	4:XD:20:TYR:OH[4_555]	1.98	0.22
9:QI:98:PRO:O	10:XJ:79:ARG:NH1[4_555]	1.99	0.21
7:QG:149:ARG:CD	3:XC:85:ARG:CG[4_555]	2.01	0.19
7:QG:149:ARG:NH2	3:XC:84:ILE:CB[4_555]	2.01	0.19
9:QI:98:PRO:C	10:XJ:79:ARG:NH1[4_555]	2.05	0.15
36:RD:135:PHE:CE1	4:XD:167:GLY:CA[4_555]	2.07	0.13
26:R2:34:GLU:OE1	34:YA:277:C:O2[3_555]	2.14	0.06
6:QF:15:ASP:CB	4:XD:20:TYR:CD2[4_555]	2.17	0.03
27:R3:60:GLU:O	44:YP:119:GLU:OE2[3_455]	2.17	0.03
9:QI:98:PRO:CB	10:XJ:79:ARG:NH2[4_555]	2.18	0.02
6:QF:14:LEU:CA	4:XD:20:TYR:OH[4_555]	2.19	0.01
7:QG:149:ARG:NH1	3:XC:85:ARG:CG[4_555]	2.19	0.01
7:QG:149:ARG:CD	3:XC:81:GLY:C[4_555]	2.19	0.01

5.3 Torsion angles ⓘ

5.3.1 Protein backbone ⓘ

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
2	QB	233/256 (91%)	206 (88%)	27 (12%)	0	100	100
2	XB	234/256 (91%)	202 (86%)	31 (13%)	1 (0%)	30	67
3	QC	203/239 (85%)	180 (89%)	23 (11%)	0	100	100
3	XC	203/239 (85%)	182 (90%)	21 (10%)	0	100	100
4	QD	206/209 (99%)	195 (95%)	10 (5%)	1 (0%)	25	62
4	XD	206/209 (99%)	194 (94%)	11 (5%)	1 (0%)	25	62
5	QE	149/162 (92%)	134 (90%)	14 (9%)	1 (1%)	19	56
5	XE	149/162 (92%)	140 (94%)	8 (5%)	1 (1%)	19	56
6	QF	99/101 (98%)	96 (97%)	3 (3%)	0	100	100
6	XF	99/101 (98%)	98 (99%)	1 (1%)	0	100	100
7	QG	153/156 (98%)	145 (95%)	8 (5%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
7	XG	153/156 (98%)	142 (93%)	11 (7%)	0	100	100
8	QH	135/138 (98%)	127 (94%)	8 (6%)	0	100	100
8	XH	135/138 (98%)	126 (93%)	9 (7%)	0	100	100
9	QI	103/128 (80%)	91 (88%)	12 (12%)	0	100	100
9	XI	105/128 (82%)	97 (92%)	8 (8%)	0	100	100
10	QJ	97/105 (92%)	88 (91%)	8 (8%)	1 (1%)	13	47
10	XJ	94/105 (90%)	87 (93%)	5 (5%)	2 (2%)	5	32
11	QK	117/129 (91%)	108 (92%)	9 (8%)	0	100	100
11	XK	114/129 (88%)	104 (91%)	10 (9%)	0	100	100
12	QL	123/132 (93%)	98 (80%)	24 (20%)	1 (1%)	16	53
12	XL	120/132 (91%)	99 (82%)	21 (18%)	0	100	100
13	QM	113/126 (90%)	96 (85%)	16 (14%)	1 (1%)	14	50
13	XM	112/126 (89%)	100 (89%)	11 (10%)	1 (1%)	14	50
14	QN	58/61 (95%)	50 (86%)	7 (12%)	1 (2%)	7	37
14	XN	58/61 (95%)	49 (84%)	7 (12%)	2 (3%)	3	24
15	QO	86/89 (97%)	80 (93%)	6 (7%)	0	100	100
15	XO	85/89 (96%)	81 (95%)	4 (5%)	0	100	100
16	QP	82/88 (93%)	76 (93%)	6 (7%)	0	100	100
16	XP	82/88 (93%)	78 (95%)	4 (5%)	0	100	100
17	QQ	98/105 (93%)	91 (93%)	7 (7%)	0	100	100
17	XQ	98/105 (93%)	94 (96%)	4 (4%)	0	100	100
18	QR	68/88 (77%)	66 (97%)	2 (3%)	0	100	100
18	XR	68/88 (77%)	66 (97%)	2 (3%)	0	100	100
19	QS	81/93 (87%)	66 (82%)	15 (18%)	0	100	100
19	XS	82/93 (88%)	65 (79%)	17 (21%)	0	100	100
20	QT	97/106 (92%)	86 (89%)	8 (8%)	3 (3%)	3	25
20	XT	97/106 (92%)	84 (87%)	10 (10%)	3 (3%)	3	25
21	QU	23/27 (85%)	21 (91%)	2 (9%)	0	100	100
21	XU	23/27 (85%)	23 (100%)	0	0	100	100
24	R0	79/85 (93%)	71 (90%)	8 (10%)	0	100	100
24	Y0	80/85 (94%)	75 (94%)	5 (6%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
25	R1	93/98 (95%)	76 (82%)	17 (18%)	0	100	100
25	Y1	91/98 (93%)	78 (86%)	12 (13%)	1 (1%)	12	45
26	R2	67/72 (93%)	63 (94%)	4 (6%)	0	100	100
26	Y2	66/72 (92%)	64 (97%)	2 (3%)	0	100	100
27	R3	57/60 (95%)	56 (98%)	1 (2%)	0	100	100
27	Y3	57/60 (95%)	55 (96%)	2 (4%)	0	100	100
28	R4	43/71 (61%)	41 (95%)	2 (5%)	0	100	100
28	Y4	44/71 (62%)	28 (64%)	12 (27%)	4 (9%)	0	10
29	R5	57/60 (95%)	49 (86%)	7 (12%)	1 (2%)	7	35
29	Y5	57/60 (95%)	49 (86%)	7 (12%)	1 (2%)	7	35
30	R6	51/54 (94%)	46 (90%)	5 (10%)	0	100	100
30	Y6	51/54 (94%)	49 (96%)	2 (4%)	0	100	100
31	R7	45/49 (92%)	43 (96%)	2 (4%)	0	100	100
31	Y7	46/49 (94%)	45 (98%)	1 (2%)	0	100	100
32	R8	62/65 (95%)	51 (82%)	9 (14%)	2 (3%)	3	25
32	Y8	62/65 (95%)	48 (77%)	14 (23%)	0	100	100
33	R9	35/37 (95%)	33 (94%)	2 (6%)	0	100	100
33	Y9	35/37 (95%)	33 (94%)	2 (6%)	0	100	100
36	RD	270/276 (98%)	244 (90%)	24 (9%)	2 (1%)	19	56
36	YD	270/276 (98%)	241 (89%)	28 (10%)	1 (0%)	30	67
37	RE	203/206 (98%)	159 (78%)	39 (19%)	5 (2%)	4	29
37	YE	203/206 (98%)	162 (80%)	39 (19%)	2 (1%)	13	47
38	RF	200/210 (95%)	183 (92%)	15 (8%)	2 (1%)	13	47
38	YF	200/210 (95%)	183 (92%)	16 (8%)	1 (0%)	25	62
39	RG	179/182 (98%)	150 (84%)	28 (16%)	1 (1%)	22	59
39	YG	179/182 (98%)	152 (85%)	27 (15%)	0	100	100
40	RH	172/180 (96%)	145 (84%)	24 (14%)	3 (2%)	7	37
40	YH	172/180 (96%)	147 (86%)	21 (12%)	4 (2%)	5	31
41	RI	144/148 (97%)	114 (79%)	24 (17%)	6 (4%)	2	21
41	YI	144/148 (97%)	118 (82%)	22 (15%)	4 (3%)	4	27
42	RN	136/140 (97%)	122 (90%)	13 (10%)	1 (1%)	19	56

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
42	YN	136/140 (97%)	123 (90%)	12 (9%)	1 (1%)	19	56
43	RO	120/122 (98%)	109 (91%)	11 (9%)	0	100	100
43	YO	120/122 (98%)	112 (93%)	8 (7%)	0	100	100
44	RP	148/150 (99%)	114 (77%)	31 (21%)	3 (2%)	6	34
44	YP	145/150 (97%)	116 (80%)	28 (19%)	1 (1%)	19	56
45	RQ	139/141 (99%)	120 (86%)	18 (13%)	1 (1%)	19	56
45	YQ	139/141 (99%)	111 (80%)	27 (19%)	1 (1%)	19	56
46	RR	115/118 (98%)	103 (90%)	12 (10%)	0	100	100
46	YR	115/118 (98%)	104 (90%)	10 (9%)	1 (1%)	14	50
47	RS	109/112 (97%)	95 (87%)	14 (13%)	0	100	100
47	YS	109/112 (97%)	95 (87%)	13 (12%)	1 (1%)	14	50
48	RT	135/146 (92%)	116 (86%)	19 (14%)	0	100	100
48	YT	135/146 (92%)	121 (90%)	14 (10%)	0	100	100
49	RU	115/118 (98%)	106 (92%)	6 (5%)	3 (3%)	4	29
49	YU	115/118 (98%)	109 (95%)	6 (5%)	0	100	100
50	RV	99/101 (98%)	87 (88%)	11 (11%)	1 (1%)	13	47
50	YV	99/101 (98%)	90 (91%)	8 (8%)	1 (1%)	13	47
51	RW	111/113 (98%)	104 (94%)	7 (6%)	0	100	100
51	YW	111/113 (98%)	107 (96%)	4 (4%)	0	100	100
52	RX	90/96 (94%)	85 (94%)	5 (6%)	0	100	100
52	YX	90/96 (94%)	84 (93%)	6 (7%)	0	100	100
53	RY	105/110 (96%)	102 (97%)	3 (3%)	0	100	100
53	YY	105/110 (96%)	99 (94%)	6 (6%)	0	100	100
54	RZ	181/206 (88%)	139 (77%)	38 (21%)	4 (2%)	5	32
54	YZ	191/206 (93%)	145 (76%)	39 (20%)	7 (4%)	2	23
All	All	11368/12128 (94%)	10080 (89%)	1202 (11%)	86 (1%)	16	53

All (86) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
12	QL	105	TYR
20	QT	75	ASN
32	R8	30	ARG

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Mol	Chain	Res	Type
37	RE	147	PRO
40	RH	157	TYR
41	RI	11	ASN
44	RP	108	LYS
49	RU	91	ASP
49	RU	92	ARG
54	RZ	53	ILE
4	XD	156	GLU
20	XT	74	LYS
20	XT	75	ASN
28	Y4	24	THR
37	YE	147	PRO
40	YH	157	TYR
44	YP	108	LYS
50	YV	50	PRO
54	YZ	53	ILE
54	YZ	182	LYS
14	QN	17	LYS
32	R8	29	LYS
36	RD	243	GLY
39	RG	81	LYS
40	RH	126	PRO
41	RI	132	PRO
42	RN	22	THR
44	RP	22	GLY
49	RU	90	VAL
54	RZ	167	PRO
14	XN	57	ARG
36	YD	243	GLY
40	YH	47	GLU
41	YI	122	GLU
41	YI	132	PRO
47	YS	110	LEU
54	YZ	60	GLU
54	YZ	167	PRO
20	QT	74	LYS
36	RD	242	ARG
37	RE	130	GLY
38	RF	67	GLN
38	RF	129	PHE
10	XJ	37	PRO
14	XN	56	VAL

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Mol	Chain	Res	Type
28	Y4	5	ILE
38	YF	129	PHE
46	YR	4	LEU
10	QJ	55	LYS
29	R5	49	CYS
37	RE	17	ASP
37	RE	83	ASP
41	RI	10	GLU
41	RI	122	GLU
44	RP	29	LYS
54	RZ	52	SER
54	RZ	63	ASP
10	XJ	40	LEU
13	XM	7	VAL
20	XT	73	HIS
28	Y4	40	HIS
40	YH	152	ARG
41	YI	15	VAL
54	YZ	183	LEU
20	QT	98	PRO
37	RE	82	ARG
40	RH	156	ALA
50	RV	53	GLU
5	XE	74	GLY
40	YH	156	ALA
42	YN	22	THR
54	YZ	61	LEU
4	QD	156	GLU
13	QM	13	LYS
41	RI	15	VAL
25	Y1	54	ALA
29	Y5	49	CYS
37	YE	54	GLN
45	YQ	7	MET
54	YZ	94	GLU
5	QE	74	GLY
2	XB	208	ILE
28	Y4	4	GLY
45	RQ	78	PRO
41	YI	133	HIS
41	RI	119	PRO

5.3.2 Protein sidechains ⓘ

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
2	QB	203/220 (92%)	200 (98%)	3 (2%)	60	75
2	XB	204/220 (93%)	204 (100%)	0	100	100
3	QC	159/188 (85%)	157 (99%)	2 (1%)	65	77
3	XC	159/188 (85%)	157 (99%)	2 (1%)	65	77
4	QD	180/181 (99%)	179 (99%)	1 (1%)	84	88
4	XD	180/181 (99%)	178 (99%)	2 (1%)	70	80
5	QE	116/123 (94%)	116 (100%)	0	100	100
5	XE	116/123 (94%)	115 (99%)	1 (1%)	75	83
6	QF	90/90 (100%)	90 (100%)	0	100	100
6	XF	90/90 (100%)	89 (99%)	1 (1%)	70	80
7	QG	126/127 (99%)	126 (100%)	0	100	100
7	XG	126/127 (99%)	126 (100%)	0	100	100
8	QH	118/119 (99%)	117 (99%)	1 (1%)	79	84
8	XH	118/119 (99%)	118 (100%)	0	100	100
9	QI	79/99 (80%)	77 (98%)	2 (2%)	42	62
9	XI	81/99 (82%)	80 (99%)	1 (1%)	67	79
10	QJ	89/92 (97%)	89 (100%)	0	100	100
10	XJ	86/92 (94%)	84 (98%)	2 (2%)	45	64
11	QK	90/99 (91%)	89 (99%)	1 (1%)	70	80
11	XK	88/99 (89%)	87 (99%)	1 (1%)	70	80
12	QL	104/109 (95%)	104 (100%)	0	100	100
12	XL	103/109 (94%)	100 (97%)	3 (3%)	37	58
13	QM	93/101 (92%)	93 (100%)	0	100	100
13	XM	92/101 (91%)	92 (100%)	0	100	100
14	QN	49/50 (98%)	48 (98%)	1 (2%)	50	68
14	XN	49/50 (98%)	47 (96%)	2 (4%)	26	49

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
15	QO	79/80 (99%)	77 (98%)	2 (2%)	42	62
15	XO	79/80 (99%)	79 (100%)	0	100	100
16	QP	72/74 (97%)	72 (100%)	0	100	100
16	XP	72/74 (97%)	72 (100%)	0	100	100
17	QQ	95/97 (98%)	95 (100%)	0	100	100
17	XQ	95/97 (98%)	93 (98%)	2 (2%)	48	66
18	QR	61/77 (79%)	61 (100%)	0	100	100
18	XR	61/77 (79%)	61 (100%)	0	100	100
19	QS	72/80 (90%)	72 (100%)	0	100	100
19	XS	73/80 (91%)	73 (100%)	0	100	100
20	QT	76/82 (93%)	75 (99%)	1 (1%)	65	77
20	XT	76/82 (93%)	76 (100%)	0	100	100
21	QU	20/22 (91%)	19 (95%)	1 (5%)	20	44
21	XU	20/22 (91%)	19 (95%)	1 (5%)	20	44
24	R0	65/67 (97%)	64 (98%)	1 (2%)	60	75
24	Y0	65/67 (97%)	65 (100%)	0	100	100
25	R1	80/83 (96%)	78 (98%)	2 (2%)	42	62
25	Y1	78/83 (94%)	78 (100%)	0	100	100
26	R2	64/67 (96%)	64 (100%)	0	100	100
26	Y2	64/67 (96%)	63 (98%)	1 (2%)	58	74
27	R3	51/52 (98%)	51 (100%)	0	100	100
27	Y3	51/52 (98%)	51 (100%)	0	100	100
28	R4	40/63 (64%)	40 (100%)	0	100	100
28	Y4	41/63 (65%)	40 (98%)	1 (2%)	44	63
29	R5	51/52 (98%)	50 (98%)	1 (2%)	50	68
29	Y5	51/52 (98%)	48 (94%)	3 (6%)	16	40
30	R6	51/52 (98%)	49 (96%)	2 (4%)	27	50
30	Y6	51/52 (98%)	49 (96%)	2 (4%)	27	50
31	R7	40/42 (95%)	40 (100%)	0	100	100
31	Y7	41/42 (98%)	41 (100%)	0	100	100
32	R8	54/55 (98%)	54 (100%)	0	100	100

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
32	Y8	54/55 (98%)	54 (100%)	0	100	100
33	R9	34/34 (100%)	34 (100%)	0	100	100
33	Y9	34/34 (100%)	33 (97%)	1 (3%)	37	58
36	RD	214/218 (98%)	212 (99%)	2 (1%)	75	83
36	YD	214/218 (98%)	214 (100%)	0	100	100
37	RE	165/166 (99%)	161 (98%)	4 (2%)	44	63
37	YE	165/166 (99%)	163 (99%)	2 (1%)	67	79
38	RF	161/166 (97%)	158 (98%)	3 (2%)	52	69
38	YF	161/166 (97%)	161 (100%)	0	100	100
39	RG	155/156 (99%)	155 (100%)	0	100	100
39	YG	155/156 (99%)	154 (99%)	1 (1%)	84	88
40	RH	145/148 (98%)	137 (94%)	8 (6%)	18	42
40	YH	145/148 (98%)	143 (99%)	2 (1%)	62	75
41	RI	122/124 (98%)	121 (99%)	1 (1%)	79	84
41	YI	122/124 (98%)	118 (97%)	4 (3%)	33	55
42	RN	117/119 (98%)	116 (99%)	1 (1%)	75	83
42	YN	117/119 (98%)	115 (98%)	2 (2%)	56	72
43	RO	100/100 (100%)	100 (100%)	0	100	100
43	YO	100/100 (100%)	98 (98%)	2 (2%)	50	68
44	RP	116/116 (100%)	115 (99%)	1 (1%)	75	83
44	YP	114/116 (98%)	114 (100%)	0	100	100
45	RQ	111/111 (100%)	111 (100%)	0	100	100
45	YQ	111/111 (100%)	110 (99%)	1 (1%)	75	83
46	RR	100/101 (99%)	99 (99%)	1 (1%)	73	81
46	YR	100/101 (99%)	99 (99%)	1 (1%)	73	81
47	RS	87/88 (99%)	87 (100%)	0	100	100
47	YS	87/88 (99%)	85 (98%)	2 (2%)	45	64
48	RT	120/127 (94%)	117 (98%)	3 (2%)	42	62
48	YT	120/127 (94%)	118 (98%)	2 (2%)	56	72
49	RU	93/94 (99%)	93 (100%)	0	100	100
49	YU	93/94 (99%)	91 (98%)	2 (2%)	47	65

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
50	RV	82/82 (100%)	82 (100%)	0	100	100
50	YV	82/82 (100%)	80 (98%)	2 (2%)	44	63
51	RW	92/92 (100%)	91 (99%)	1 (1%)	70	80
51	YW	92/92 (100%)	92 (100%)	0	100	100
52	RX	74/78 (95%)	71 (96%)	3 (4%)	26	49
52	YX	74/78 (95%)	72 (97%)	2 (3%)	40	60
53	RY	88/91 (97%)	88 (100%)	0	100	100
53	YY	88/91 (97%)	87 (99%)	1 (1%)	70	80
54	RZ	162/179 (90%)	162 (100%)	0	100	100
54	YZ	167/179 (93%)	165 (99%)	2 (1%)	67	79
All	All	9610/10066 (96%)	9507 (99%)	103 (1%)	70	80

All (103) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
2	QB	36	ARG
2	QB	149	LEU
2	QB	217	ARG
3	QC	38	ARG
3	QC	127	ARG
4	QD	18	LYS
8	QH	59	LEU
9	QI	64	THR
9	QI	85	LEU
11	QK	41	THR
14	QN	43	CYS
15	QO	22	THR
15	QO	87	ILE
20	QT	73	HIS
21	QU	6	ARG
24	R0	14	ARG
25	R1	73	LEU
25	R1	92	LYS
29	R5	36	CYS
30	R6	13	CYS
30	R6	23	THR
36	RD	242	ARG
36	RD	268	ARG

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Mol	Chain	Res	Type
37	RE	107	THR
37	RE	144	ARG
37	RE	154	LYS
37	RE	176	ILE
38	RF	89	VAL
38	RF	144	LYS
38	RF	195	ASP
40	RH	17	VAL
40	RH	69	ARG
40	RH	123	PHE
40	RH	125	VAL
40	RH	127	GLU
40	RH	129	THR
40	RH	130	ARG
40	RH	131	VAL
41	RI	81	VAL
42	RN	34	LEU
44	RP	16	ARG
46	RR	10	LEU
48	RT	85	LYS
48	RT	111	ARG
48	RT	129	ARG
51	RW	52	GLU
52	RX	16	LYS
52	RX	27	THR
52	RX	76	ARG
3	XC	12	LEU
3	XC	162	GLN
4	XD	135	LEU
4	XD	191	ARG
5	XE	12	LEU
6	XF	80	ARG
9	XI	27	THR
10	XJ	8	LEU
10	XJ	69	ASN
11	XK	117	ASN
12	XL	8	ASN
12	XL	89	ARG
12	XL	105	TYR
14	XN	3	ARG
14	XN	27	CYS
17	XQ	50	LYS

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Mol	Chain	Res	Type
17	XQ	74	LEU
21	XU	3	LYS
26	Y2	47	ASN
28	Y4	16	CYS
29	Y5	37	LYS
29	Y5	40	LYS
29	Y5	45	VAL
30	Y6	13	CYS
30	Y6	43	CYS
33	Y9	27	CYS
37	YE	107	THR
37	YE	184	VAL
39	YG	118	ARG
40	YH	67	LEU
40	YH	69	ARG
41	YI	56	LYS
41	YI	86	THR
41	YI	93	THR
41	YI	118	LYS
42	YN	96	GLU
42	YN	115	ARG
43	YO	24	VAL
43	YO	49	ARG
45	YQ	133	ARG
46	YR	2	ARG
47	YS	4	LEU
47	YS	110	LEU
48	YT	105	LEU
48	YT	129	ARG
49	YU	92	ARG
49	YU	94	ASN
50	YV	46	VAL
50	YV	78	LYS
52	YX	49	VAL
52	YX	66	LEU
53	YY	79	CYS
54	YZ	63	ASP
54	YZ	165	VAL

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (40) such sidechains are listed below:

Mol	Chain	Res	Type
3	QC	6	HIS
3	QC	176	HIS
4	QD	103	ASN
4	QD	123	HIS
7	QG	64	GLN
12	QL	9	GLN
13	QM	77	ASN
15	QO	62	GLN
16	QP	13	HIS
16	QP	14	ASN
19	QS	69	HIS
24	R0	12	ASN
25	R1	47	GLN
32	R8	31	HIS
32	R8	35	GLN
36	RD	253	GLN
38	RF	67	GLN
39	RG	132	ASN
42	RN	101	HIS
47	RS	38	GLN
52	RX	41	ASN
3	XC	162	GLN
3	XC	176	HIS
4	XD	116	GLN
4	XD	119	GLN
4	XD	123	HIS
6	XF	7	ASN
6	XF	57	GLN
6	XF	100	ASN
10	XJ	62	HIS
11	XK	117	ASN
12	XL	8	ASN
13	XM	77	ASN
14	XN	49	HIS
15	XO	28	GLN
28	Y4	6	HIS
29	Y5	22	HIS
37	YE	132	HIS
44	YP	70	GLN
50	YV	11	GLN

5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	QA	1509/1521 (99%)	531 (35%)	14 (0%)
1	XA	1514/1521 (99%)	469 (30%)	27 (1%)
22	QV	66/77 (85%)	15 (22%)	1 (1%)
22	XV	66/77 (85%)	16 (24%)	1 (1%)
23	QX	18/19 (94%)	5 (27%)	0
23	XX	18/19 (94%)	5 (27%)	0
34	RA	2878/2905 (99%)	706 (24%)	40 (1%)
34	YA	2880/2905 (99%)	754 (26%)	40 (1%)
35	RB	119/122 (97%)	20 (16%)	1 (0%)
35	YB	119/122 (97%)	23 (19%)	1 (0%)
All	All	9187/9288 (98%)	2544 (27%)	125 (1%)

All (2544) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	QA	6	G
1	QA	7	G
1	QA	8	A
1	QA	9	G
1	QA	15	G
1	QA	16	A
1	QA	25	C
1	QA	27	G
1	QA	29	G
1	QA	32	A
1	QA	39	G
1	QA	41	G
1	QA	42	G
1	QA	44	G
1	QA	47	C
1	QA	48	C
1	QA	49	U
1	QA	50	A
1	QA	51	A
1	QA	53	A
1	QA	54	C
1	QA	58	C
1	QA	62	U
1	QA	65	U
1	QA	68	G
1	QA	69	G
1	QA	79	G
1	QA	80	G

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Mol	Chain	Res	Type
1	QA	82	U
1	QA	95	G
1	QA	102	G
1	QA	109	A
1	QA	113	G
1	QA	121	C
1	QA	129(B)	G
1	QA	130	A
1	QA	131	C
1	QA	132	C
1	QA	134	A
1	QA	136	C
1	QA	137	C
1	QA	138	G
1	QA	142	G
1	QA	144	G
1	QA	146	G
1	QA	147	G
1	QA	151	A
1	QA	153	C
1	QA	156	G
1	QA	157	G
1	QA	161	A
1	QA	163	C
1	QA	169	C
1	QA	171	A
1	QA	178	C
1	QA	181	G
1	QA	182	U
1	QA	183	G
1	QA	186(C)	C
1	QA	186(D)	G
1	QA	186(I)	U
1	QA	186(J)	U
1	QA	186(L)	G
1	QA	186(M)	G
1	QA	195	A
1	QA	196	A
1	QA	197	A
1	QA	201(A)	C
1	QA	201(B)	U
1	QA	201(C)	U

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Mol	Chain	Res	Type
1	QA	201(D)	U
1	QA	216	G
1	QA	228	A
1	QA	229	U
1	QA	235	C
1	QA	240	C
1	QA	241	C
1	QA	243	A
1	QA	244	U
1	QA	245	C
1	QA	247	G
1	QA	250	A
1	QA	252	U
1	QA	254	G
1	QA	258	G
1	QA	264	U
1	QA	266	G
1	QA	267	C
1	QA	274	A
1	QA	275	G
1	QA	281	G
1	QA	289	G
1	QA	296	U
1	QA	297	G
1	QA	299	G
1	QA	301	G
1	QA	305	G
1	QA	306	G
1	QA	315	A
1	QA	316	G
1	QA	321	A
1	QA	322	C
1	QA	324	G
1	QA	327	A
1	QA	328	C
1	QA	329	A
1	QA	330	C
1	QA	332	G
1	QA	344	A
1	QA	345	C
1	QA	346	G
1	QA	347	G

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Mol	Chain	Res	Type
1	QA	348	G
1	QA	351	G
1	QA	352	C
1	QA	353	A
1	QA	354	G
1	QA	356	A
1	QA	367	U
1	QA	368	U
1	QA	372	C
1	QA	373	A
1	QA	381	C
1	QA	382	A
1	QA	384	G
1	QA	389	A
1	QA	390	C
1	QA	392	G
1	QA	393	A
1	QA	394	G
1	QA	397	A
1	QA	398	C
1	QA	399	G
1	QA	401	C
1	QA	406	G
1	QA	408	A
1	QA	412	A
1	QA	413	G
1	QA	414	A
1	QA	422	C
1	QA	424	G
1	QA	426	G
1	QA	428	G
1	QA	429	U
1	QA	437	U
1	QA	438	G
1	QA	440	A
1	QA	443	C
1	QA	452	A
1	QA	453	A
1	QA	458(B)	G
1	QA	458(C)	A
1	QA	458(D)	C
1	QA	474	G

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Mol	Chain	Res	Type
1	QA	476	G
1	QA	481	G
1	QA	484	G
1	QA	485	G
1	QA	486	U
1	QA	489	C
1	QA	497	A
1	QA	498	U
1	QA	501	C
1	QA	508	C
1	QA	509	A
1	QA	510	A
1	QA	511	C
1	QA	514	C
1	QA	515	G
1	QA	517	G
1	QA	518	C
1	QA	519	C
1	QA	521	G
1	QA	522	C
1	QA	527	G
1	QA	531	U
1	QA	533	A
1	QA	534	U
1	QA	536	C
1	QA	537	G
1	QA	538	G
1	QA	541	G
1	QA	542	G
1	QA	546	G
1	QA	547	A
1	QA	550	G
1	QA	559	A
1	QA	562	C
1	QA	564	C
1	QA	572	A
1	QA	573	A
1	QA	574	A
1	QA	576	G
1	QA	577	G
1	QA	580	U
1	QA	582	U

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Mol	Chain	Res	Type
1	QA	587	G
1	QA	588	G
1	QA	590	C
1	QA	592	G
1	QA	596	C
1	QA	597	G
1	QA	599	C
1	QA	603	U
1	QA	607	A
1	QA	609	A
1	QA	614	A
1	QA	616	G
1	QA	617	G
1	QA	618	C
1	QA	619	U
1	QA	620	C
1	QA	624	C
1	QA	635	G
1	QA	641	U
1	QA	644	G
1	QA	647	C
1	QA	652	U
1	QA	653	A
1	QA	655	A
1	QA	661	G
1	QA	664	G
1	QA	665	A
1	QA	670	G
1	QA	683	G
1	QA	684	A
1	QA	686	U
1	QA	687	A
1	QA	688	G
1	QA	695	A
1	QA	700	G
1	QA	702	A
1	QA	703	G
1	QA	708	C
1	QA	710	G
1	QA	716	A
1	QA	718	G
1	QA	721	G

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Mol	Chain	Res	Type
1	QA	723	U
1	QA	724	G
1	QA	729	A
1	QA	749	C
1	QA	753	A
1	QA	754	C
1	QA	755	G
1	QA	773	G
1	QA	776	G
1	QA	777	A
1	QA	785	G
1	QA	787	A
1	QA	793	U
1	QA	794	A
1	QA	796	C
1	QA	799	G
1	QA	811	C
1	QA	812	C
1	QA	815	A
1	QA	816	A
1	QA	817	C
1	QA	818	G
1	QA	820	U
1	QA	821	G
1	QA	828	A
1	QA	829	G
1	QA	837	G
1	QA	838(B)	U
1	QA	838(C)	C
1	QA	838(D)	U
1	QA	848	C
1	QA	853	G
1	QA	855	G
1	QA	867	G
1	QA	871	U
1	QA	872	A
1	QA	873	A
1	QA	874	G
1	QA	876	G
1	QA	880	C
1	QA	885	G
1	QA	889	A

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Mol	Chain	Res	Type
1	QA	890	G
1	QA	914	A
1	QA	916	G
1	QA	918	A
1	QA	919	A
1	QA	926	G
1	QA	927	G
1	QA	928	G
1	QA	934	C
1	QA	935	A
1	QA	938	A
1	QA	941	G
1	QA	942	G
1	QA	946	A
1	QA	952	U
1	QA	957	U
1	QA	960	U
1	QA	961	U
1	QA	966	G
1	QA	967	C
1	QA	968	A
1	QA	969	A
1	QA	971	G
1	QA	972	C
1	QA	973	G
1	QA	974	A
1	QA	975	A
1	QA	976	G
1	QA	977	A
1	QA	979	C
1	QA	980	C
1	QA	981	U
1	QA	986	A
1	QA	988	G
1	QA	992	U
1	QA	993	G
1	QA	998(A)	G
1	QA	999	U
1	QA	1004	A
1	QA	1006	C
1	QA	1010	G
1	QA	1015	A

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Mol	Chain	Res	Type
1	QA	1016	A
1	QA	1023	G
1	QA	1024	G
1	QA	1025	U
1	QA	1026	G
1	QA	1027	C
1	QA	1028(A)	C
1	QA	1028(C)	C
1	QA	1028(D)	G
1	QA	1028(F)	G
1	QA	1037	C
1	QA	1042	G
1	QA	1044	A
1	QA	1046	A
1	QA	1050	G
1	QA	1054	C
1	QA	1056	U
1	QA	1060	C
1	QA	1064	G
1	QA	1065	U
1	QA	1066	C
1	QA	1076	C
1	QA	1088	G
1	QA	1089	G
1	QA	1094	G
1	QA	1095	U
1	QA	1099	G
1	QA	1101	A
1	QA	1104	G
1	QA	1107	C
1	QA	1117	G
1	QA	1123	A
1	QA	1124	G
1	QA	1125	U
1	QA	1126	U
1	QA	1127	G
1	QA	1128	C
1	QA	1129	C
1	QA	1130	A
1	QA	1131	G
1	QA	1132	C
1	QA	1133	G

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Mol	Chain	Res	Type
1	QA	1136	U
1	QA	1137	C
1	QA	1138	G
1	QA	1139	G
1	QA	1140	C
1	QA	1142	G
1	QA	1143	G
1	QA	1145	C
1	QA	1146	A
1	QA	1147	C
1	QA	1151	A
1	QA	1154	G
1	QA	1157	A
1	QA	1158	C
1	QA	1159	U
1	QA	1178	G
1	QA	1179	A
1	QA	1181	G
1	QA	1183	A
1	QA	1184	G
1	QA	1190	G
1	QA	1196	U
1	QA	1197	G
1	QA	1200	C
1	QA	1201	A
1	QA	1202	G
1	QA	1203	C
1	QA	1206	G
1	QA	1211	U
1	QA	1212	U
1	QA	1213	A
1	QA	1216	G
1	QA	1218	C
1	QA	1221	G
1	QA	1222	G
1	QA	1224	G
1	QA	1225	A
1	QA	1226	C
1	QA	1228	C
1	QA	1235	U
1	QA	1236	A
1	QA	1238	A

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Mol	Chain	Res	Type
1	QA	1240	U
1	QA	1244	C
1	QA	1246	C
1	QA	1250	A
1	QA	1251	A
1	QA	1252	A
1	QA	1253	G
1	QA	1254	C
1	QA	1256	A
1	QA	1257	U
1	QA	1258	G
1	QA	1260	C
1	QA	1263	C
1	QA	1265	G
1	QA	1268	A
1	QA	1269	A
1	QA	1270	C
1	QA	1272	G
1	QA	1273	G
1	QA	1274	G
1	QA	1275	A
1	QA	1277	C
1	QA	1278	U
1	QA	1279	A
1	QA	1280	A
1	QA	1281	U
1	QA	1282	C
1	QA	1285	A
1	QA	1286	A
1	QA	1287	A
1	QA	1290	G
1	QA	1293	G
1	QA	1296	C
1	QA	1297	C
1	QA	1299	A
1	QA	1300	G
1	QA	1301	U
1	QA	1303	C
1	QA	1307	U
1	QA	1312	G
1	QA	1318	A
1	QA	1319	A

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Mol	Chain	Res	Type
1	QA	1320	C
1	QA	1321	C
1	QA	1322	C
1	QA	1323	G
1	QA	1324	A
1	QA	1325	C
1	QA	1329	A
1	QA	1331	G
1	QA	1332	A
1	QA	1335	C
1	QA	1336	C
1	QA	1338	G
1	QA	1340	A
1	QA	1346	A
1	QA	1347	G
1	QA	1348	U
1	QA	1350	A
1	QA	1353	G
1	QA	1355	G
1	QA	1356	G
1	QA	1358	U
1	QA	1359	C
1	QA	1361	G
1	QA	1363	A
1	QA	1364	U
1	QA	1365	G
1	QA	1367	C
1	QA	1370	G
1	QA	1373	G
1	QA	1375	A
1	QA	1377	A
1	QA	1378	C
1	QA	1379	G
1	QA	1380	U
1	QA	1381	U
1	QA	1382	C
1	QA	1384	C
1	QA	1394	A
1	QA	1395	C
1	QA	1397	C
1	QA	1398	A
1	QA	1399	C

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Mol	Chain	Res	Type
1	QA	1401	G
1	QA	1413	A
1	QA	1419	G
1	QA	1422	G
1	QA	1432	G
1	QA	1435	G
1	QA	1436	U
1	QA	1440(C)	G
1	QA	1440(D)	G
1	QA	1440(E)	A
1	QA	1440(H)	C
1	QA	1440(J)	A
1	QA	1440(K)	C
1	QA	1440(L)	G
1	QA	1475	G
1	QA	1487	G
1	QA	1490	C
1	QA	1492	A
1	QA	1493	A
1	QA	1494	G
1	QA	1497	G
1	QA	1499	A
1	QA	1502	A
1	QA	1503	A
1	QA	1504	G
1	QA	1505	G
1	QA	1506	U
1	QA	1507	A
1	QA	1517	G
1	QA	1519	A
1	QA	1520	G
1	QA	1529	G
1	QA	1530	G
1	QA	1532	U
1	QA	1533	C
1	QA	1534	A
1	QA	1535	C
1	QA	1537	U
1	QA	1538	C
1	QA	1539	C
1	QA	1541	U
22	QV	8	U

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Mol	Chain	Res	Type
22	QV	10	G
22	QV	11	C
22	QV	15	G
22	QV	18	U
22	QV	19	G
22	QV	21(B)	A
22	QV	22	G
22	QV	31	G
22	QV	46	G
22	QV	47	U
22	QV	48	C
22	QV	59	A
22	QV	61	C
22	QV	67	U
23	QX	10	G
23	QX	12	A
23	QX	13	A
23	QX	16	C
23	QX	19	U
34	RA	7	G
34	RA	15	G
34	RA	23	G
34	RA	34	C
34	RA	35	G
34	RA	36	G
34	RA	46	C
34	RA	51	G
34	RA	55	G
34	RA	64	A
34	RA	71	A
34	RA	72	U
34	RA	74	A
34	RA	75	G
34	RA	83	G
34	RA	101	G
34	RA	102	G
34	RA	103	A
34	RA	114	U
34	RA	118	A
34	RA	120	U
34	RA	125	G
34	RA	128	C

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Mol	Chain	Res	Type
34	RA	131	G
34	RA	140	A
34	RA	161	U
34	RA	177	G
34	RA	181	A
34	RA	196	A
34	RA	198	C
34	RA	199	A
34	RA	206	U
34	RA	215	G
34	RA	216	A
34	RA	221	A
34	RA	222	A
34	RA	223	A
34	RA	228	A
34	RA	229	A
34	RA	230	U
34	RA	232	G
34	RA	242	G
34	RA	243	U
34	RA	245	G
34	RA	248	G
34	RA	249	C
34	RA	252	G
34	RA	261	G
34	RA	265	A
34	RA	266	G
34	RA	267	C
34	RA	270(L)	C
34	RA	270(M)	U
34	RA	270(N)	U
34	RA	270(O)	G
34	RA	270(Q)	C
34	RA	271(D)	U
34	RA	275	G
34	RA	276	A
34	RA	277	C
34	RA	278	A
34	RA	280	C
34	RA	283	A
34	RA	299	A
34	RA	300	A

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Mol	Chain	Res	Type
34	RA	311	A
34	RA	312	G
34	RA	316	C
34	RA	317	G
34	RA	323	G
34	RA	324	A
34	RA	329	G
34	RA	330	A
34	RA	332	A
34	RA	342	G
34	RA	343	C
34	RA	352	G
34	RA	362	U
34	RA	364	C
34	RA	371	A
34	RA	372	G
34	RA	373	U
34	RA	386	G
34	RA	387	U
34	RA	405	U
34	RA	407	G
34	RA	411	G
34	RA	412	A
34	RA	428	A
34	RA	444	C
34	RA	448	U
34	RA	451	C
34	RA	454	A
34	RA	455	C
34	RA	456	C
34	RA	457	A
34	RA	458	G
34	RA	459	U
34	RA	467	G
34	RA	470	A
34	RA	479	A
34	RA	481	G
34	RA	494	G
34	RA	496	G
34	RA	504	U
34	RA	505	A
34	RA	508	G

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Mol	Chain	Res	Type
34	RA	510	C
34	RA	512	G
34	RA	513	A
34	RA	527	C
34	RA	528	A
34	RA	529	A
34	RA	530	G
34	RA	531	C
34	RA	532	A
34	RA	533	G
34	RA	537	C
34	RA	539	G
34	RA	540	G
34	RA	546	C
34	RA	547	A
34	RA	554	U
34	RA	563	G
34	RA	568	U
34	RA	573	G
34	RA	575	A
34	RA	592	G
34	RA	599	G
34	RA	603	A
34	RA	604	G
34	RA	607	U
34	RA	613	U
34	RA	614	U
34	RA	615	G
34	RA	616	A
34	RA	617	G
34	RA	618(A)	G
34	RA	621	A
34	RA	622	G
34	RA	626	U
34	RA	627	A
34	RA	634	C
34	RA	637	A
34	RA	638	G
34	RA	645	C
34	RA	646	A
34	RA	651	G
34	RA	652	C

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Mol	Chain	Res	Type
34	RA	654(A)	A
34	RA	654(B)	G
34	RA	657	U
34	RA	668	G
34	RA	669	G
34	RA	670	A
34	RA	677	A
34	RA	685	A
34	RA	686	G
34	RA	689	A
34	RA	695	G
34	RA	702	G
34	RA	717	G
34	RA	722	A
34	RA	726	G
34	RA	730	C
34	RA	734	A
34	RA	738	G
34	RA	747	U
34	RA	748	G
34	RA	753	C
34	RA	764	A
34	RA	765	G
34	RA	769	G
34	RA	775	G
34	RA	776	G
34	RA	777	A
34	RA	782	A
34	RA	784	A
34	RA	785	G
34	RA	789	A
34	RA	792	G
34	RA	800	A
34	RA	805	G
34	RA	809	G
34	RA	812	C
34	RA	819	A
34	RA	827	U
34	RA	828	U
34	RA	829	A
34	RA	845	G
34	RA	847	U

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Mol	Chain	Res	Type
34	RA	854	G
34	RA	856	C
34	RA	857	C
34	RA	859	G
34	RA	860	U
34	RA	865	C
34	RA	867	C
34	RA	869	G
34	RA	877	U
34	RA	887	A
34	RA	889	C
34	RA	890	A
34	RA	896	A
34	RA	905	U
34	RA	907	U
34	RA	910	A
34	RA	915	C
34	RA	917	A
34	RA	919	G
34	RA	926	A
34	RA	932	G
34	RA	941	A
34	RA	945	A
34	RA	946	G
34	RA	953	A
34	RA	957	A
34	RA	959	A
34	RA	960	A
34	RA	961	C
34	RA	974(A)	G
34	RA	974(B)	C
34	RA	975	G
34	RA	980	A
34	RA	983	A
34	RA	996	A
34	RA	1005	C
34	RA	1008	C
34	RA	1011	G
34	RA	1012	U
34	RA	1013	C
34	RA	1015	G
34	RA	1022	G

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Mol	Chain	Res	Type
34	RA	1023	U
34	RA	1025	G
34	RA	1026	U
34	RA	1027	A
34	RA	1033	U
34	RA	1044	G
34	RA	1045	A
34	RA	1046	A
34	RA	1050	A
34	RA	1058	G
34	RA	1060	U
34	RA	1067	A
34	RA	1070	A
34	RA	1071	G
34	RA	1072	C
34	RA	1073	A
34	RA	1077	A
34	RA	1082	U
34	RA	1083	U
34	RA	1084	A
34	RA	1085	A
34	RA	1088	A
34	RA	1089	G
34	RA	1090	U
34	RA	1097	U
34	RA	1099	G
34	RA	1102	C
34	RA	1103	A
34	RA	1106	G
34	RA	1110	G
34	RA	1112	G
34	RA	1113	U
34	RA	1114	G
34	RA	1122	G
34	RA	1128	A
34	RA	1129	A
34	RA	1130	U
34	RA	1131	G
34	RA	1135	C
34	RA	1136	G
34	RA	1139	G
34	RA	1141	U

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Mol	Chain	Res	Type
34	RA	1142(A)	U
34	RA	1142(B)	A
34	RA	1155	A
34	RA	1156	A
34	RA	1168	G
34	RA	1170	G
34	RA	1173	G
34	RA	1174	A
34	RA	1175	U
34	RA	1176	G
34	RA	1179	C
34	RA	1181	C
34	RA	1190	G
34	RA	1195	G
34	RA	1204	A
34	RA	1205	U
34	RA	1206	G
34	RA	1210	A
34	RA	1212	G
34	RA	1213	A
34	RA	1220	A
34	RA	1236	G
34	RA	1238	G
34	RA	1240	U
34	RA	1241	A
34	RA	1244	G
34	RA	1247	A
34	RA	1250	G
34	RA	1253	A
34	RA	1255	U
34	RA	1256	G
34	RA	1265	A
34	RA	1271	G
34	RA	1272	A
34	RA	1273	U
34	RA	1281	G
34	RA	1282	U
34	RA	1286	A
34	RA	1300	U
34	RA	1301	A
34	RA	1308	A
34	RA	1313	U

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Mol	Chain	Res	Type
34	RA	1314	C
34	RA	1319	G
34	RA	1326	U
34	RA	1329	U
34	RA	1332	G
34	RA	1341	U
34	RA	1349	A
34	RA	1350	C
34	RA	1352	U
34	RA	1365	A
34	RA	1368	G
34	RA	1370	C
34	RA	1378	A
34	RA	1384	A
34	RA	1385	G
34	RA	1394	U
34	RA	1407	C
34	RA	1408	C
34	RA	1410	G
34	RA	1411	C
34	RA	1416	G
34	RA	1419	A
34	RA	1420	U
34	RA	1421	G
34	RA	1427	A
34	RA	1428	C
34	RA	1434	A
34	RA	1444(B)	A
34	RA	1445	C
34	RA	1448	G
34	RA	1449(A)	A
34	RA	1449(B)	G
34	RA	1451	C
34	RA	1455	G
34	RA	1461	G
34	RA	1466	G
34	RA	1467	C
34	RA	1471	A
34	RA	1472	A
34	RA	1474	C
34	RA	1480	G
34	RA	1482	U

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Mol	Chain	Res	Type
34	RA	1483	G
34	RA	1485	G
34	RA	1490	A
34	RA	1493	C
34	RA	1494	A
34	RA	1496	A
34	RA	1497	U
34	RA	1506	C
34	RA	1507	A
34	RA	1508	A
34	RA	1510	A
34	RA	1515	C
34	RA	1521	G
34	RA	1533	C
34	RA	1534	G
34	RA	1535	U
34	RA	1536	A
34	RA	1537	C
34	RA	1538	G
34	RA	1543	A
34	RA	1544	C
34	RA	1545(A)	A
34	RA	1558	A
34	RA	1559	G
34	RA	1560	G
34	RA	1566	A
34	RA	1567	A
34	RA	1569	A
34	RA	1578	U
34	RA	1579	A
34	RA	1582	C
34	RA	1586	A
34	RA	1598	C
34	RA	1607	C
34	RA	1608	A
34	RA	1609	A
34	RA	1610	A
34	RA	1611	C
34	RA	1613	G
34	RA	1616	A
34	RA	1617	C
34	RA	1618	A

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Mol	Chain	Res	Type
34	RA	1640	C
34	RA	1644	C
34	RA	1646	C
34	RA	1648	C
34	RA	1649	G
34	RA	1654	A
34	RA	1664	A
34	RA	1665	A
34	RA	1667	G
34	RA	1668	A
34	RA	1674	G
34	RA	1675	C
34	RA	1688	U
34	RA	1694	C
34	RA	1695	G
34	RA	1698	A
34	RA	1718	G
34	RA	1725	G
34	RA	1729	A
34	RA	1731	G
34	RA	1732	A
34	RA	1733	G
34	RA	1742	C
34	RA	1743	G
34	RA	1756	G
34	RA	1762	A
34	RA	1763	G
34	RA	1764	G
34	RA	1773	A
34	RA	1774	C
34	RA	1780	A
34	RA	1784	A
34	RA	1787	A
34	RA	1791	A
34	RA	1799	G
34	RA	1800	C
34	RA	1816	G
34	RA	1820	U
34	RA	1829	A
34	RA	1835	G
34	RA	1846	G
34	RA	1847	A

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Mol	Chain	Res	Type
34	RA	1848	A
34	RA	1858	G
34	RA	1869	G
34	RA	1872	A
34	RA	1878	G
34	RA	1881	C
34	RA	1882	C
34	RA	1884	A
34	RA	1888	G
34	RA	1889	A
34	RA	1896	G
34	RA	1900	A
34	RA	1903	G
34	RA	1905	C
34	RA	1906	G
34	RA	1913	A
34	RA	1929	G
34	RA	1930	G
34	RA	1931	U
34	RA	1932	A
34	RA	1936	A
34	RA	1937	A
34	RA	1938	A
34	RA	1939	U
34	RA	1955	U
34	RA	1963	U
34	RA	1965	C
34	RA	1966	A
34	RA	1967	C
34	RA	1969	A
34	RA	1970	A
34	RA	1971	A
34	RA	1972	A
34	RA	1981	A
34	RA	1982	C
34	RA	1992	G
34	RA	1993	U
34	RA	2020	A
34	RA	2021	C
34	RA	2023	G
34	RA	2031	A
34	RA	2032	G

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Mol	Chain	Res	Type
34	RA	2033	A
34	RA	2043	C
34	RA	2049	G
34	RA	2052	G
34	RA	2055	C
34	RA	2056	G
34	RA	2059	A
34	RA	2060	A
34	RA	2061	G
34	RA	2062	A
34	RA	2069	G
34	RA	2077	A
34	RA	2080	G
34	RA	2089	U
34	RA	2092	U
34	RA	2093	G
34	RA	2095	C
34	RA	2096	U
34	RA	2097	C
34	RA	2099	U
34	RA	2110	G
34	RA	2111	C
34	RA	2112	G
34	RA	2113	U
34	RA	2114	A
34	RA	2115	G
34	RA	2116	G
34	RA	2117	A
34	RA	2118	U
34	RA	2119	A
34	RA	2120	G
34	RA	2124	G
34	RA	2126	A
34	RA	2127	G
34	RA	2128	C
34	RA	2131	G
34	RA	2132	U
34	RA	2133	G
34	RA	2134	A
34	RA	2136	C
34	RA	2137	C
34	RA	2142	C

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Mol	Chain	Res	Type
34	RA	2145	C
34	RA	2146	C
34	RA	2147	G
34	RA	2148	G
34	RA	2158	A
34	RA	2164	C
34	RA	2166	G
34	RA	2169	A
34	RA	2173	A
34	RA	2176	A
34	RA	2178	C
34	RA	2189	U
34	RA	2190	G
34	RA	2191	G
34	RA	2192	G
34	RA	2194	G
34	RA	2197	U
34	RA	2198	A
34	RA	2209	C
34	RA	2210	G
34	RA	2211	G
34	RA	2212	A
34	RA	2215	G
34	RA	2225	A
34	RA	2238	G
34	RA	2239	G
34	RA	2243	U
34	RA	2244	U
34	RA	2266	A
34	RA	2275	C
34	RA	2283	C
34	RA	2287	A
34	RA	2288	A
34	RA	2302	G
34	RA	2304	G
34	RA	2305	A
34	RA	2307	G
34	RA	2308	G
34	RA	2310	A
34	RA	2311	A
34	RA	2319	G
34	RA	2320	A

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Mol	Chain	Res	Type
34	RA	2321	G
34	RA	2325	G
34	RA	2334	G
34	RA	2335	A
34	RA	2342	C
34	RA	2345	G
34	RA	2346	A
34	RA	2347	C
34	RA	2350	C
34	RA	2354	G
34	RA	2372	G
34	RA	2383	G
34	RA	2385	C
34	RA	2392	A
34	RA	2396	G
34	RA	2402	C
34	RA	2403	C
34	RA	2406	U
34	RA	2414	G
34	RA	2423	U
34	RA	2424	C
34	RA	2425	A
34	RA	2427	C
34	RA	2428	G
34	RA	2429	G
34	RA	2430	A
34	RA	2432	A
34	RA	2435	A
34	RA	2439	A
34	RA	2440	C
34	RA	2441	C
34	RA	2445	G
34	RA	2447	G
34	RA	2448	A
34	RA	2469	A
34	RA	2470	G
34	RA	2476	A
34	RA	2478	A
34	RA	2482	G
34	RA	2484	G
34	RA	2487	G
34	RA	2491	U

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Mol	Chain	Res	Type
34	RA	2494	G
34	RA	2499	C
34	RA	2502	G
34	RA	2503	A
34	RA	2504	U
34	RA	2505	G
34	RA	2506	U
34	RA	2518	A
34	RA	2520	C
34	RA	2529	G
34	RA	2535	G
34	RA	2542	A
34	RA	2543	G
34	RA	2554	U
34	RA	2562	U
34	RA	2566	A
34	RA	2567	G
34	RA	2569	G
34	RA	2572	A
34	RA	2573	C
34	RA	2577	A
34	RA	2584	U
34	RA	2585	U
34	RA	2602	A
34	RA	2609	U
34	RA	2610	C
34	RA	2611	U
34	RA	2612	C
34	RA	2614	A
34	RA	2615	U
34	RA	2621	A
34	RA	2629	A
34	RA	2630	G
34	RA	2655	G
34	RA	2665	A
34	RA	2673	G
34	RA	2675	A
34	RA	2682	U
34	RA	2686	G
34	RA	2689	U
34	RA	2690	C
34	RA	2691	C

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Mol	Chain	Res	Type
34	RA	2702	U
34	RA	2712(A)	U
34	RA	2712(B)	A
34	RA	2713	A
34	RA	2714	G
34	RA	2718	G
34	RA	2726	U
34	RA	2732	G
34	RA	2733	A
34	RA	2734	A
34	RA	2744	G
34	RA	2748	A
34	RA	2751	G
34	RA	2755	C
34	RA	2757	A
34	RA	2758	A
34	RA	2761	G
34	RA	2764	A
34	RA	2765	A
34	RA	2766	G
34	RA	2777	G
34	RA	2778	A
34	RA	2779	U
34	RA	2780	G
34	RA	2790	A
34	RA	2791	C
34	RA	2792	G
34	RA	2794	C
34	RA	2797	U
34	RA	2798	C
34	RA	2802	G
34	RA	2807	G
34	RA	2808	U
34	RA	2813	A
34	RA	2818	G
34	RA	2820	A
34	RA	2821	A
34	RA	2830	G
34	RA	2833	G
34	RA	2834	G
34	RA	2835	A
34	RA	2847	U

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Mol	Chain	Res	Type
34	RA	2849	U
34	RA	2850	A
34	RA	2867	G
34	RA	2872	G
34	RA	2873	A
34	RA	2880	C
34	RA	2883	A
34	RA	2886	G
34	RA	2892	A
34	RA	2894	G
34	RA	2895	U
35	RB	8	U
35	RB	9	G
35	RB	12	C
35	RB	13	A
35	RB	15	A
35	RB	22	U
35	RB	25	A
35	RB	32	C
35	RB	41	U
35	RB	42	C
35	RB	44	G
35	RB	45	A
35	RB	53	A
35	RB	56	G
35	RB	67	G
35	RB	73	A
35	RB	77	U
35	RB	81	G
35	RB	96	G
35	RB	109	G
1	XA	6	G
1	XA	9	G
1	XA	16	A
1	XA	18	C
1	XA	19	C
1	XA	22	G
1	XA	31	G
1	XA	32	A
1	XA	39	G
1	XA	41	G
1	XA	44	G

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Mol	Chain	Res	Type
1	XA	47	C
1	XA	48	C
1	XA	51	A
1	XA	59	A
1	XA	65	U
1	XA	66	G
1	XA	68(E)	C
1	XA	68(F)	G
1	XA	68(I)	G
1	XA	68(K)	G
1	XA	68(M)	U
1	XA	68(N)	U
1	XA	68(P)	A
1	XA	68(Q)	C
1	XA	68(T)	C
1	XA	68(V)	U
1	XA	68(W)	G
1	XA	101	A
1	XA	108	G
1	XA	109	A
1	XA	116	A
1	XA	121	C
1	XA	129(B)	G
1	XA	131	C
1	XA	142	G
1	XA	144	G
1	XA	147	G
1	XA	151	A
1	XA	160	A
1	XA	161	A
1	XA	163	C
1	XA	169	C
1	XA	173	U
1	XA	177	C
1	XA	182	U
1	XA	183	G
1	XA	186(A)	C
1	XA	186(H)	C
1	XA	186(K)	G
1	XA	186(L)	G
1	XA	186(N)	G
1	XA	195	A

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Mol	Chain	Res	Type
1	XA	197	A
1	XA	201(B)	U
1	XA	201(C)	U
1	XA	201(D)	U
1	XA	216	G
1	XA	231	G
1	XA	247	G
1	XA	251	G
1	XA	259	G
1	XA	261	U
1	XA	263	A
1	XA	266	G
1	XA	267	C
1	XA	278	G
1	XA	281	G
1	XA	289	G
1	XA	309	G
1	XA	310	G
1	XA	315	A
1	XA	328	C
1	XA	329	A
1	XA	330	C
1	XA	332	G
1	XA	345	C
1	XA	346	G
1	XA	347	G
1	XA	348	G
1	XA	351	G
1	XA	352	C
1	XA	353	A
1	XA	354	G
1	XA	356	A
1	XA	367	U
1	XA	368	U
1	XA	369	C
1	XA	372	C
1	XA	373	A
1	XA	378	G
1	XA	379	C
1	XA	386	C
1	XA	387	U
1	XA	390	C

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Mol	Chain	Res	Type
1	XA	391	G
1	XA	392	G
1	XA	393	A
1	XA	397	A
1	XA	398	C
1	XA	412	A
1	XA	413	G
1	XA	414	A
1	XA	422	C
1	XA	423	G
1	XA	424	G
1	XA	428	G
1	XA	429	U
1	XA	440	A
1	XA	443	C
1	XA	444	C
1	XA	451	A
1	XA	452	A
1	XA	453	A
1	XA	458(C)	A
1	XA	458(D)	C
1	XA	481	G
1	XA	484	G
1	XA	485	G
1	XA	495	A
1	XA	497	A
1	XA	498	U
1	XA	509	A
1	XA	511	C
1	XA	517	G
1	XA	518	C
1	XA	521	G
1	XA	524	G
1	XA	527	G
1	XA	530	G
1	XA	531	U
1	XA	532	A
1	XA	533	A
1	XA	536	C
1	XA	537	G
1	XA	547	A
1	XA	549	C

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Mol	Chain	Res	Type
1	XA	559	A
1	XA	560	U
1	XA	562	C
1	XA	568	G
1	XA	571	U
1	XA	572	A
1	XA	573	A
1	XA	576	G
1	XA	577	G
1	XA	581	G
1	XA	587	G
1	XA	588	G
1	XA	590	C
1	XA	596	C
1	XA	608	A
1	XA	610	G
1	XA	613	C
1	XA	617	G
1	XA	618	C
1	XA	620	C
1	XA	625	G
1	XA	629	G
1	XA	634	C
1	XA	637	G
1	XA	642	A
1	XA	653	A
1	XA	654	G
1	XA	661	G
1	XA	665	A
1	XA	671	G
1	XA	674	G
1	XA	686	U
1	XA	688	G
1	XA	695	A
1	XA	702	A
1	XA	703	G
1	XA	718	G
1	XA	721	G
1	XA	723	U
1	XA	724	G
1	XA	739	C
1	XA	741	G

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Mol	Chain	Res	Type
1	XA	749	C
1	XA	751	U
1	XA	753	A
1	XA	755	G
1	XA	758	G
1	XA	760	G
1	XA	777	A
1	XA	781	A
1	XA	789	U
1	XA	793	U
1	XA	794	A
1	XA	799	G
1	XA	809	G
1	XA	811	C
1	XA	812	C
1	XA	815	A
1	XA	816	A
1	XA	817	C
1	XA	818	G
1	XA	821	G
1	XA	828	A
1	XA	829	G
1	XA	838(B)	U
1	XA	838(C)	C
1	XA	838(D)	U
1	XA	848	C
1	XA	853	G
1	XA	855	G
1	XA	859	A
1	XA	867	G
1	XA	871	U
1	XA	872	A
1	XA	876	G
1	XA	877	C
1	XA	884	U
1	XA	885	G
1	XA	889	A
1	XA	898	G
1	XA	899	C
1	XA	902	G
1	XA	914	A
1	XA	916	G

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Mol	Chain	Res	Type
1	XA	926	G
1	XA	927	G
1	XA	934	C
1	XA	935	A
1	XA	943	U
1	XA	946	A
1	XA	948	C
1	XA	957	U
1	XA	958	A
1	XA	960	U
1	XA	961	U
1	XA	966	G
1	XA	967	C
1	XA	968	A
1	XA	969	A
1	XA	971	G
1	XA	972	C
1	XA	973	G
1	XA	974	A
1	XA	975	A
1	XA	976	G
1	XA	977	A
1	XA	979	C
1	XA	980	C
1	XA	981	U
1	XA	991	U
1	XA	992	U
1	XA	993	G
1	XA	1004	A
1	XA	1006	C
1	XA	1015	A
1	XA	1023	G
1	XA	1025	U
1	XA	1026	G
1	XA	1028(C)	C
1	XA	1028(D)	G
1	XA	1036	G
1	XA	1037	C
1	XA	1043	C
1	XA	1046	A
1	XA	1048	G
1	XA	1050	G

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Mol	Chain	Res	Type
1	XA	1053	G
1	XA	1054	C
1	XA	1055	A
1	XA	1058	G
1	XA	1060	C
1	XA	1064	G
1	XA	1065	U
1	XA	1066	C
1	XA	1076	C
1	XA	1080	A
1	XA	1081	G
1	XA	1083	U
1	XA	1089	G
1	XA	1094	G
1	XA	1095	U
1	XA	1099	G
1	XA	1101	A
1	XA	1106	G
1	XA	1107	C
1	XA	1114	C
1	XA	1116	C
1	XA	1118	C
1	XA	1123	A
1	XA	1124	G
1	XA	1125	U
1	XA	1126	U
1	XA	1127	G
1	XA	1128	C
1	XA	1129	C
1	XA	1130	A
1	XA	1131	G
1	XA	1132	C
1	XA	1134	G
1	XA	1136	U
1	XA	1137	C
1	XA	1138	G
1	XA	1139	G
1	XA	1145	C
1	XA	1146	A
1	XA	1147	C
1	XA	1150	U
1	XA	1151	A

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Mol	Chain	Res	Type
1	XA	1154	G
1	XA	1157	A
1	XA	1158	C
1	XA	1159	U
1	XA	1161	C
1	XA	1163	C
1	XA	1177	G
1	XA	1178	G
1	XA	1179	A
1	XA	1181	G
1	XA	1182	G
1	XA	1183	A
1	XA	1184	G
1	XA	1187	G
1	XA	1190	G
1	XA	1193	G
1	XA	1196	U
1	XA	1197	G
1	XA	1200	C
1	XA	1201	A
1	XA	1202	G
1	XA	1203	C
1	XA	1211	U
1	XA	1212	U
1	XA	1213	A
1	XA	1216	G
1	XA	1221	G
1	XA	1222	G
1	XA	1225	A
1	XA	1226	C
1	XA	1227	A
1	XA	1228	C
1	XA	1232	U
1	XA	1236	A
1	XA	1237	C
1	XA	1238	A
1	XA	1240	U
1	XA	1241	G
1	XA	1243	C
1	XA	1246	C
1	XA	1252	A
1	XA	1253	G

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Mol	Chain	Res	Type
1	XA	1254	C
1	XA	1256	A
1	XA	1257	U
1	XA	1258	G
1	XA	1260	C
1	XA	1263	C
1	XA	1265	G
1	XA	1269	A
1	XA	1270	C
1	XA	1272	G
1	XA	1273	G
1	XA	1274	G
1	XA	1275	A
1	XA	1277	C
1	XA	1278	U
1	XA	1280	A
1	XA	1281	U
1	XA	1282	C
1	XA	1285	A
1	XA	1286	A
1	XA	1287	A
1	XA	1290	G
1	XA	1293	G
1	XA	1296	C
1	XA	1297	C
1	XA	1298	C
1	XA	1299	A
1	XA	1300	G
1	XA	1302	U
1	XA	1303	C
1	XA	1304	G
1	XA	1305	G
1	XA	1306	A
1	XA	1307	U
1	XA	1312	G
1	XA	1319	A
1	XA	1320	C
1	XA	1321	C
1	XA	1322	C
1	XA	1323	G
1	XA	1325	C
1	XA	1327	C

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Mol	Chain	Res	Type
1	XA	1328	C
1	XA	1329	A
1	XA	1330	U
1	XA	1331	G
1	XA	1335	C
1	XA	1336	C
1	XA	1337	G
1	XA	1340	A
1	XA	1346	A
1	XA	1347	G
1	XA	1348	U
1	XA	1349	A
1	XA	1353	G
1	XA	1355	G
1	XA	1356	G
1	XA	1359	C
1	XA	1360	A
1	XA	1361	G
1	XA	1363	A
1	XA	1364	U
1	XA	1365	G
1	XA	1368	G
1	XA	1370	G
1	XA	1373	G
1	XA	1375	A
1	XA	1378	C
1	XA	1379	G
1	XA	1381	U
1	XA	1382	C
1	XA	1384	C
1	XA	1386	G
1	XA	1394	A
1	XA	1397	C
1	XA	1398	A
1	XA	1413	A
1	XA	1419	G
1	XA	1426	C
1	XA	1432	G
1	XA	1433	A
1	XA	1435	G
1	XA	1438	G
1	XA	1440(B)	G

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Mol	Chain	Res	Type
1	XA	1440(C)	G
1	XA	1440(D)	G
1	XA	1440(E)	A
1	XA	1440(I)	U
1	XA	1440(J)	A
1	XA	1440(K)	C
1	XA	1440(L)	G
1	XA	1475	G
1	XA	1477	C
1	XA	1487	G
1	XA	1492	A
1	XA	1493	A
1	XA	1494	G
1	XA	1495	U
1	XA	1496	C
1	XA	1497	G
1	XA	1498	U
1	XA	1499	A
1	XA	1500	A
1	XA	1502	A
1	XA	1503	A
1	XA	1504	G
1	XA	1505	G
1	XA	1506	U
1	XA	1507	A
1	XA	1517	G
1	XA	1519	A
1	XA	1520	G
1	XA	1528	U
1	XA	1529	G
1	XA	1530	G
1	XA	1531	A
1	XA	1532	U
1	XA	1533	C
1	XA	1534	A
1	XA	1538	C
1	XA	1539	C
1	XA	1541	U
22	XV	8	U
22	XV	11	C
22	XV	16	C
22	XV	17	C

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Mol	Chain	Res	Type
22	XV	18	U
22	XV	19	G
22	XV	20	G
22	XV	21(B)	A
22	XV	42	A
22	XV	43	G
22	XV	46	G
22	XV	47	U
22	XV	48	C
22	XV	58	A
22	XV	59	A
22	XV	61	C
23	XX	9	G
23	XX	10	G
23	XX	12	A
23	XX	13	A
23	XX	15	A
34	YA	9	U
34	YA	15	G
34	YA	23	G
34	YA	28	A
34	YA	34	C
34	YA	35	G
34	YA	36	G
34	YA	46	C
34	YA	51	G
34	YA	54	G
34	YA	55	G
34	YA	64	A
34	YA	72	U
34	YA	74	A
34	YA	75	G
34	YA	78	A
34	YA	90	U
34	YA	96	G
34	YA	101	G
34	YA	102	G
34	YA	103	A
34	YA	118	A
34	YA	120	U
34	YA	125	G
34	YA	131	G

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Mol	Chain	Res	Type
34	YA	138	G
34	YA	142	G
34	YA	155	C
34	YA	161	U
34	YA	162	U
34	YA	178	G
34	YA	188	G
34	YA	196	A
34	YA	199	A
34	YA	204	A
34	YA	205	G
34	YA	215	G
34	YA	216	A
34	YA	221	A
34	YA	222	A
34	YA	223	A
34	YA	226	G
34	YA	228	A
34	YA	229	A
34	YA	230	U
34	YA	232	G
34	YA	233	A
34	YA	242	G
34	YA	243	U
34	YA	245	G
34	YA	248	G
34	YA	252	G
34	YA	261	G
34	YA	265	A
34	YA	266	G
34	YA	267	C
34	YA	269	U
34	YA	270(L)	C
34	YA	270(M)	U
34	YA	270(N)	U
34	YA	270(Q)	C
34	YA	271(D)	U
34	YA	271(E)	G
34	YA	274	G
34	YA	275	G
34	YA	277	C
34	YA	278	A

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Mol	Chain	Res	Type
34	YA	279	C
34	YA	283	A
34	YA	299	A
34	YA	300	A
34	YA	311	A
34	YA	316	C
34	YA	317	G
34	YA	323	G
34	YA	329	G
34	YA	330	A
34	YA	332	A
34	YA	338	G
34	YA	342	G
34	YA	343	C
34	YA	352	G
34	YA	363(A)	G
34	YA	363(F)	U
34	YA	364	C
34	YA	371	A
34	YA	372	G
34	YA	374	A
34	YA	380	U
34	YA	386	G
34	YA	387	U
34	YA	399	G
34	YA	405	U
34	YA	407	G
34	YA	411	G
34	YA	412	A
34	YA	420	C
34	YA	428	A
34	YA	429	A
34	YA	444	C
34	YA	448	U
34	YA	454	A
34	YA	455	C
34	YA	457	A
34	YA	470	A
34	YA	473	G
34	YA	479	A
34	YA	480	A
34	YA	481	G

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Mol	Chain	Res	Type
34	YA	494	G
34	YA	496	G
34	YA	504	U
34	YA	505	A
34	YA	508	G
34	YA	509	C
34	YA	512	G
34	YA	518	G
34	YA	527	C
34	YA	528	A
34	YA	529	A
34	YA	530	G
34	YA	531	C
34	YA	532	A
34	YA	533	G
34	YA	537	C
34	YA	539	G
34	YA	540	G
34	YA	546	C
34	YA	547	A
34	YA	554	U
34	YA	563	G
34	YA	568	U
34	YA	573	G
34	YA	574	C
34	YA	575	A
34	YA	599	G
34	YA	603	A
34	YA	604	G
34	YA	607	U
34	YA	614	U
34	YA	615	G
34	YA	617	G
34	YA	621	A
34	YA	622	G
34	YA	627	A
34	YA	629	G
34	YA	634	C
34	YA	637	A
34	YA	638	G
34	YA	645	C
34	YA	646	A

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Mol	Chain	Res	Type
34	YA	647	G
34	YA	650	C
34	YA	651	G
34	YA	654(A)	A
34	YA	654(B)	G
34	YA	658	C
34	YA	686	G
34	YA	695	G
34	YA	700	G
34	YA	701	G
34	YA	704	G
34	YA	714	U
34	YA	721	C
34	YA	722	A
34	YA	726	G
34	YA	730	C
34	YA	731	C
34	YA	738	G
34	YA	747	U
34	YA	748	G
34	YA	753	C
34	YA	764	A
34	YA	771	G
34	YA	775	G
34	YA	777	A
34	YA	782	A
34	YA	784	A
34	YA	785	G
34	YA	789	A
34	YA	790	C
34	YA	792	G
34	YA	805	G
34	YA	812	C
34	YA	819	A
34	YA	827	U
34	YA	828	U
34	YA	847	U
34	YA	854	G
34	YA	856	C
34	YA	857	C
34	YA	859	G
34	YA	860	U

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Mol	Chain	Res	Type
34	YA	865	C
34	YA	866	A
34	YA	869	G
34	YA	872	A
34	YA	877	U
34	YA	878	A
34	YA	879	G
34	YA	889	C
34	YA	890	A
34	YA	896	A
34	YA	897	C
34	YA	899	A
34	YA	901	A
34	YA	904	C
34	YA	905	U
34	YA	906	G
34	YA	907	U
34	YA	910	A
34	YA	915	C
34	YA	917	A
34	YA	926	A
34	YA	932	G
34	YA	933	A
34	YA	941	A
34	YA	945	A
34	YA	946	G
34	YA	953	A
34	YA	957	A
34	YA	959	A
34	YA	961	C
34	YA	973	A
34	YA	974(A)	G
34	YA	974(B)	C
34	YA	975	G
34	YA	983	A
34	YA	989	G
34	YA	1005	C
34	YA	1008	C
34	YA	1011	G
34	YA	1012	U
34	YA	1013	C
34	YA	1015	G

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Mol	Chain	Res	Type
34	YA	1017	G
34	YA	1020	A
34	YA	1022	G
34	YA	1023	U
34	YA	1024	G
34	YA	1026	U
34	YA	1027	A
34	YA	1033	U
34	YA	1037	G
34	YA	1044	G
34	YA	1045	A
34	YA	1046	A
34	YA	1048	A
34	YA	1050	A
34	YA	1051	G
34	YA	1054	A
34	YA	1058	G
34	YA	1059	G
34	YA	1061	U
34	YA	1062	G
34	YA	1065	U
34	YA	1067	A
34	YA	1068	G
34	YA	1071	G
34	YA	1073	A
34	YA	1074	G
34	YA	1076	C
34	YA	1077	A
34	YA	1078	U
34	YA	1081	U
34	YA	1082	U
34	YA	1083	U
34	YA	1084	A
34	YA	1085	A
34	YA	1086	A
34	YA	1087	G
34	YA	1088	A
34	YA	1090	U
34	YA	1093	G
34	YA	1095	A
34	YA	1096	A
34	YA	1097	U

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Mol	Chain	Res	Type
34	YA	1099	G
34	YA	1103	A
34	YA	1104	C
34	YA	1110	G
34	YA	1111	A
34	YA	1112	G
34	YA	1115	G
34	YA	1122	G
34	YA	1126	A
34	YA	1130	U
34	YA	1131	G
34	YA	1132	A
34	YA	1135	C
34	YA	1136	G
34	YA	1139	G
34	YA	1142(A)	U
34	YA	1142(B)	A
34	YA	1156	A
34	YA	1168	G
34	YA	1170	G
34	YA	1173	G
34	YA	1174	A
34	YA	1175	U
34	YA	1176	G
34	YA	1179	C
34	YA	1180	C
34	YA	1183	G
34	YA	1186	G
34	YA	1190	G
34	YA	1195	G
34	YA	1204	A
34	YA	1205	U
34	YA	1206	G
34	YA	1210	A
34	YA	1212	G
34	YA	1218	C
34	YA	1220	A
34	YA	1221	C
34	YA	1236	G
34	YA	1238	G
34	YA	1240	U
34	YA	1241	A

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Mol	Chain	Res	Type
34	YA	1243	G
34	YA	1244	G
34	YA	1247	A
34	YA	1248	G
34	YA	1253	A
34	YA	1256	G
34	YA	1265	A
34	YA	1267	U
34	YA	1271	G
34	YA	1272	A
34	YA	1273	U
34	YA	1275	A
34	YA	1281	G
34	YA	1288	U
34	YA	1300	U
34	YA	1301	A
34	YA	1313	U
34	YA	1319	G
34	YA	1329	U
34	YA	1341	U
34	YA	1349	A
34	YA	1352	U
34	YA	1365	A
34	YA	1368	G
34	YA	1370	C
34	YA	1378	A
34	YA	1379	A
34	YA	1384	A
34	YA	1385	G
34	YA	1386	C
34	YA	1388	G
34	YA	1395	A
34	YA	1403	C
34	YA	1404	C
34	YA	1407	C
34	YA	1408	C
34	YA	1411	C
34	YA	1412	A
34	YA	1416	G
34	YA	1419	A
34	YA	1420	U
34	YA	1421	G

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Mol	Chain	Res	Type
34	YA	1427	A
34	YA	1428	C
34	YA	1437	C
34	YA	1444(B)	A
34	YA	1445	C
34	YA	1448	G
34	YA	1449(A)	A
34	YA	1449(B)	G
34	YA	1455	G
34	YA	1461	G
34	YA	1464	C
34	YA	1467	C
34	YA	1471	A
34	YA	1472	A
34	YA	1475	G
34	YA	1478	G
34	YA	1480	G
34	YA	1482	U
34	YA	1483	G
34	YA	1485	G
34	YA	1487	G
34	YA	1490	A
34	YA	1493	C
34	YA	1496	A
34	YA	1497	U
34	YA	1506	C
34	YA	1507	A
34	YA	1508	A
34	YA	1509	C
34	YA	1510	A
34	YA	1511	A
34	YA	1514	U
34	YA	1521	G
34	YA	1523	U
34	YA	1534	G
34	YA	1535	U
34	YA	1536	A
34	YA	1537	C
34	YA	1538	G
34	YA	1543	A
34	YA	1544	C
34	YA	1545(A)	A

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Mol	Chain	Res	Type
34	YA	1554	A
34	YA	1555	G
34	YA	1558	A
34	YA	1559	G
34	YA	1560	G
34	YA	1566	A
34	YA	1567	A
34	YA	1569	A
34	YA	1578	U
34	YA	1580	A
34	YA	1585	C
34	YA	1586	A
34	YA	1607	C
34	YA	1608	A
34	YA	1609	A
34	YA	1610	A
34	YA	1613	G
34	YA	1616	A
34	YA	1617	C
34	YA	1618	A
34	YA	1634	A
34	YA	1638	C
34	YA	1640	C
34	YA	1644	C
34	YA	1648	C
34	YA	1651	G
34	YA	1654	A
34	YA	1665	A
34	YA	1670	C
34	YA	1674	G
34	YA	1693	U
34	YA	1694	C
34	YA	1695	G
34	YA	1718	G
34	YA	1725	G
34	YA	1728	G
34	YA	1729	A
34	YA	1730	U
34	YA	1731	G
34	YA	1732	A
34	YA	1741	C
34	YA	1742	C

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Mol	Chain	Res	Type
34	YA	1743	G
34	YA	1750	G
34	YA	1753	G
34	YA	1756	G
34	YA	1762	A
34	YA	1763	G
34	YA	1764	G
34	YA	1773	A
34	YA	1774	C
34	YA	1776	G
34	YA	1779	U
34	YA	1780	A
34	YA	1781	C
34	YA	1782	C
34	YA	1784	A
34	YA	1787	A
34	YA	1791	A
34	YA	1799	G
34	YA	1800	C
34	YA	1802	A
34	YA	1816	G
34	YA	1820	U
34	YA	1847	A
34	YA	1853	A
34	YA	1858	G
34	YA	1864	U
34	YA	1869	G
34	YA	1872	A
34	YA	1878	G
34	YA	1881	C
34	YA	1882	C
34	YA	1885	A
34	YA	1888	G
34	YA	1889	A
34	YA	1900	A
34	YA	1903	G
34	YA	1906	G
34	YA	1913	A
34	YA	1914	C
34	YA	1929	G
34	YA	1930	G
34	YA	1931	U

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Mol	Chain	Res	Type
34	YA	1932	A
34	YA	1936	A
34	YA	1937	A
34	YA	1938	A
34	YA	1939	U
34	YA	1944	U
34	YA	1955	U
34	YA	1963	U
34	YA	1965	C
34	YA	1967	C
34	YA	1969	A
34	YA	1970	A
34	YA	1971	A
34	YA	1972	A
34	YA	1981	A
34	YA	1982	C
34	YA	1989	G
34	YA	1991	U
34	YA	1992	G
34	YA	1993	U
34	YA	2004	G
34	YA	2013	A
34	YA	2020	A
34	YA	2021	C
34	YA	2022	U
34	YA	2023	G
34	YA	2031	A
34	YA	2032	G
34	YA	2033	A
34	YA	2039	C
34	YA	2043	C
34	YA	2052	G
34	YA	2055	C
34	YA	2056	G
34	YA	2059	A
34	YA	2060	A
34	YA	2061	G
34	YA	2062	A
34	YA	2069	G
34	YA	2075	U
34	YA	2080	G
34	YA	2092	U

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Mol	Chain	Res	Type
34	YA	2093	G
34	YA	2094	G
34	YA	2099	U
34	YA	2107	C
34	YA	2108	C
34	YA	2111	C
34	YA	2113	U
34	YA	2114	A
34	YA	2115	G
34	YA	2116	G
34	YA	2118	U
34	YA	2119	A
34	YA	2120	G
34	YA	2126	A
34	YA	2127	G
34	YA	2128	C
34	YA	2131	G
34	YA	2132	U
34	YA	2133	G
34	YA	2135	A
34	YA	2137	C
34	YA	2138	C
34	YA	2145	C
34	YA	2146	C
34	YA	2148	G
34	YA	2158	A
34	YA	2159	G
34	YA	2164	C
34	YA	2165	G
34	YA	2166	G
34	YA	2172	U
34	YA	2173	A
34	YA	2176	A
34	YA	2190	G
34	YA	2191	G
34	YA	2194	G
34	YA	2197	U
34	YA	2198	A
34	YA	2199	A
34	YA	2209	C
34	YA	2210	G
34	YA	2211	G

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Mol	Chain	Res	Type
34	YA	2212	A
34	YA	2215	G
34	YA	2225	A
34	YA	2234	G
34	YA	2238	G
34	YA	2266	A
34	YA	2275	C
34	YA	2278	A
34	YA	2279	G
34	YA	2280	G
34	YA	2283	C
34	YA	2287	A
34	YA	2288	A
34	YA	2289	G
34	YA	2299	G
34	YA	2302	G
34	YA	2307	G
34	YA	2308	G
34	YA	2309	A
34	YA	2310	A
34	YA	2311	A
34	YA	2312	U
34	YA	2314	C
34	YA	2318	G
34	YA	2319	G
34	YA	2320	A
34	YA	2321	G
34	YA	2322	A
34	YA	2325	G
34	YA	2327	A
34	YA	2329	G
34	YA	2330	G
34	YA	2334	G
34	YA	2335	A
34	YA	2336	A
34	YA	2337	G
34	YA	2340	G
34	YA	2342	C
34	YA	2343	C
34	YA	2346	A
34	YA	2347	C
34	YA	2350	C

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Mol	Chain	Res	Type
34	YA	2354	G
34	YA	2372	G
34	YA	2377	A
34	YA	2379	G
34	YA	2383	G
34	YA	2385	C
34	YA	2391	G
34	YA	2392	A
34	YA	2396	G
34	YA	2400	G
34	YA	2402	C
34	YA	2403	C
34	YA	2406	U
34	YA	2410	G
34	YA	2413	G
34	YA	2414	G
34	YA	2423	U
34	YA	2424	C
34	YA	2425	A
34	YA	2427	C
34	YA	2428	G
34	YA	2429	G
34	YA	2430	A
34	YA	2435	A
34	YA	2439	A
34	YA	2440	C
34	YA	2441	C
34	YA	2445	G
34	YA	2448	A
34	YA	2469	A
34	YA	2470	G
34	YA	2471	C
34	YA	2475	C
34	YA	2476	A
34	YA	2480	C
34	YA	2484	G
34	YA	2487	G
34	YA	2491	U
34	YA	2498	C
34	YA	2500	U
34	YA	2502	G
34	YA	2504	U

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Mol	Chain	Res	Type
34	YA	2505	G
34	YA	2506	U
34	YA	2513	G
34	YA	2518	A
34	YA	2524	G
34	YA	2525	G
34	YA	2529	G
34	YA	2530	A
34	YA	2535	G
34	YA	2542	A
34	YA	2550	G
34	YA	2551	C
34	YA	2554	U
34	YA	2559	C
34	YA	2562	U
34	YA	2567	G
34	YA	2569	G
34	YA	2572	A
34	YA	2573	C
34	YA	2574	G
34	YA	2577	A
34	YA	2578	G
34	YA	2584	U
34	YA	2585	U
34	YA	2586	C
34	YA	2602	A
34	YA	2609	U
34	YA	2611	U
34	YA	2612	C
34	YA	2615	U
34	YA	2621	A
34	YA	2623	G
34	YA	2629	A
34	YA	2630	G
34	YA	2632	A
34	YA	2638	G
34	YA	2655	G
34	YA	2665	A
34	YA	2673	G
34	YA	2675	A
34	YA	2682	U
34	YA	2689	U

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Mol	Chain	Res	Type
34	YA	2690	C
34	YA	2691	C
34	YA	2701	C
34	YA	2702	U
34	YA	2703	C
34	YA	2712(A)	U
34	YA	2712(B)	A
34	YA	2713	A
34	YA	2714	G
34	YA	2718	G
34	YA	2724	C
34	YA	2726	U
34	YA	2733	A
34	YA	2734	A
34	YA	2739	U
34	YA	2744	G
34	YA	2758	A
34	YA	2761	G
34	YA	2762	G
34	YA	2765	A
34	YA	2766	G
34	YA	2777	G
34	YA	2778	A
34	YA	2779	U
34	YA	2780	G
34	YA	2787	C
34	YA	2789	C
34	YA	2790	A
34	YA	2791	C
34	YA	2792	G
34	YA	2794	C
34	YA	2797	U
34	YA	2798	C
34	YA	2807	G
34	YA	2808	U
34	YA	2818	G
34	YA	2820	A
34	YA	2821	A
34	YA	2830	G
34	YA	2832	U
34	YA	2833	G
34	YA	2834	G

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Mol	Chain	Res	Type
34	YA	2835	A
34	YA	2847	U
34	YA	2848	G
34	YA	2851	A
34	YA	2867	G
34	YA	2872	G
34	YA	2873	A
34	YA	2879	C
34	YA	2880	C
34	YA	2883	A
34	YA	2886	G
34	YA	2891	G
34	YA	2892	A
34	YA	2893	G
34	YA	2895	U
35	YB	7	G
35	YB	8	U
35	YB	9	G
35	YB	13	A
35	YB	15	A
35	YB	16	G
35	YB	19	G
35	YB	21	G
35	YB	22	U
35	YB	25	A
35	YB	35	U
35	YB	41	U
35	YB	42	C
35	YB	44	G
35	YB	45	A
35	YB	47	C
35	YB	52	A
35	YB	53	A
35	YB	56	G
35	YB	67	G
35	YB	73	A
35	YB	81	G
35	YB	109	G

All (125) RNA pucker outliers are listed below:

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Mol	Chain	Res	Type
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Mol	Chain	Res	Type
1	QA	8	A
1	QA	65	U
1	QA	251	G
1	QA	328	C
1	QA	428	G
1	QA	687	A
1	QA	748	C
1	QA	992	U
1	QA	1054	C
1	QA	1225	A
1	QA	1278	U
1	QA	1285	A
1	QA	1504	G
1	QA	1538	C
22	QV	10	G
34	RA	99	U
34	RA	102	G
34	RA	222	A
34	RA	229	A
34	RA	242	G
34	RA	271(C)	G
34	RA	277	C
34	RA	404	C
34	RA	503	A
34	RA	512	G
34	RA	637	A
34	RA	752	A
34	RA	846	C
34	RA	856	C
34	RA	1022	G
34	RA	1026	U
34	RA	1045	A
34	RA	1069	A
34	RA	1130	U
34	RA	1178	C
34	RA	1312	U
34	RA	1427	A
34	RA	1558	A
34	RA	1653	G
34	RA	1694	C
34	RA	1799	G

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Mol	Chain	Res	Type
34	RA	1819	A
34	RA	1930	G
34	RA	1992	G
34	RA	2060	A
34	RA	2126	A
34	RA	2320	A
34	RA	2439	A
34	RA	2468	G
34	RA	2566	A
34	RA	2610	C
34	RA	2689	U
34	RA	2776	A
34	RA	2791	C
34	RA	2832	U
35	RB	66	A
1	XA	115	G
1	XA	266	G
1	XA	309	G
1	XA	328	C
1	XA	428	G
1	XA	608	A
1	XA	617	G
1	XA	635	G
1	XA	748	C
1	XA	980	C
1	XA	992	U
1	XA	1054	C
1	XA	1225	A
1	XA	1253	G
1	XA	1285	A
1	XA	1304	G
1	XA	1320	C
1	XA	1346	A
1	XA	1348	U
1	XA	1359	C
1	XA	1377	A
1	XA	1379	G
1	XA	1381	U
1	XA	1440(B)	G
1	XA	1504	G
1	XA	1505	G
1	XA	1537	U

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Mol	Chain	Res	Type
22	XV	10	G
34	YA	102	G
34	YA	177	G
34	YA	221	A
34	YA	229	A
34	YA	241	A
34	YA	242	G
34	YA	271(C)	G
34	YA	278	A
34	YA	404	C
34	YA	503	A
34	YA	637	A
34	YA	653	A
34	YA	752	A
34	YA	846	C
34	YA	856	C
34	YA	859	G
34	YA	1022	G
34	YA	1026	U
34	YA	1045	A
34	YA	1085	A
34	YA	1178	C
34	YA	1427	A
34	YA	1508	A
34	YA	1535	U
34	YA	1558	A
34	YA	1608	A
34	YA	1653	G
34	YA	1694	C
34	YA	1799	G
34	YA	1819	A
34	YA	1930	G
34	YA	1992	G
34	YA	2126	A
34	YA	2439	A
34	YA	2566	A
34	YA	2681	C
34	YA	2689	U
34	YA	2712(A)	U
34	YA	2712(B)	A
34	YA	2776	A
35	YB	66	A

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

There are no non-standard protein/DNA/RNA residues in this entry.

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

Of 1038 ligands modelled in this entry, 1036 are monoatomic - leaving 2 for Mogul analysis.

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

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
56	SF4	XD	301	4	0,12,12	-	-	-		
56	SF4	QD	301	4	0,12,12	-	-	-		

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

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
56	SF4	XD	301	4	-	-	0/6/5/5
56	SF4	QD	301	4	-	-	0/6/5/5

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

There are no torsion outliers.

There are no ring outliers.

1 monomer is involved in 2 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
56	QD	301	SF4	2	0

5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

6 Fit of model and data ⓘ

6.1 Protein, DNA and RNA chains ⓘ

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

Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
1	QA	1511/1521 (99%)	-0.06	20 (1%) 74 60	186, 276, 733, 884	0
1	XA	1515/1521 (99%)	-0.17	16 (1%) 77 64	158, 261, 672, 775	0
2	QB	235/256 (91%)	0.03	10 (4%) 40 32	230, 330, 393, 429	0
2	XB	236/256 (92%)	0.28	18 (7%) 21 19	237, 291, 340, 364	0
3	QC	205/239 (85%)	0.26	13 (6%) 27 24	271, 379, 421, 449	0
3	XC	205/239 (85%)	0.35	12 (5%) 29 25	275, 359, 467, 497	0
4	QD	208/209 (99%)	0.70	23 (11%) 12 12	231, 275, 325, 369	0
4	XD	208/209 (99%)	0.60	18 (8%) 17 16	216, 258, 306, 370	0
5	QE	151/162 (93%)	0.43	10 (6%) 26 23	208, 268, 318, 333	0
5	XE	151/162 (93%)	0.12	9 (5%) 29 24	174, 222, 277, 313	0
6	QF	101/101 (100%)	0.32	6 (5%) 29 25	224, 285, 333, 355	0
6	XF	101/101 (100%)	0.46	7 (6%) 24 21	278, 343, 376, 406	0
7	QG	155/156 (99%)	-0.05	4 (2%) 57 43	332, 425, 492, 522	0
7	XG	155/156 (99%)	0.08	5 (3%) 50 38	327, 458, 559, 580	0
8	QH	137/138 (99%)	1.15	27 (19%) 3 6	230, 290, 341, 368	0
8	XH	137/138 (99%)	0.61	12 (8%) 17 16	197, 259, 299, 342	0
9	QI	105/128 (82%)	0.07	1 (0%) 79 65	289, 373, 413, 449	0
9	XI	107/128 (83%)	0.21	3 (2%) 55 41	364, 448, 487, 522	0
10	QJ	99/105 (94%)	0.41	3 (3%) 52 40	354, 404, 458, 476	0
10	XJ	96/105 (91%)	0.73	9 (9%) 15 15	340, 397, 495, 509	0
11	QK	119/129 (92%)	0.45	8 (6%) 25 22	172, 269, 315, 333	0
11	XK	116/129 (89%)	0.55	9 (7%) 20 19	213, 279, 334, 364	0
12	QL	125/132 (94%)	0.79	14 (11%) 11 12	183, 237, 288, 334	0
12	XL	122/132 (92%)	0.87	21 (17%) 5 7	168, 221, 289, 338	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
13	QM	115/126 (91%)	0.02	2 (1%) 69 53	298, 386, 443, 469	0
13	XM	114/126 (90%)	0.34	7 (6%) 28 24	317, 414, 450, 463	0
14	QN	60/61 (98%)	0.74	7 (11%) 10 11	295, 353, 479, 493	0
14	XN	60/61 (98%)	1.00	12 (20%) 3 6	312, 556, 587, 612	0
15	QO	88/89 (98%)	0.73	13 (14%) 7 9	196, 272, 314, 335	0
15	XO	87/89 (97%)	0.57	9 (10%) 13 13	182, 221, 267, 284	0
16	QP	84/88 (95%)	0.97	14 (16%) 5 7	189, 247, 303, 348	0
16	XP	84/88 (95%)	1.56	30 (35%) 1 2	264, 358, 461, 493	0
17	QQ	100/105 (95%)	1.19	19 (19%) 4 6	199, 251, 286, 319	0
17	XQ	100/105 (95%)	0.61	16 (16%) 6 8	174, 231, 273, 307	0
18	QR	70/88 (79%)	0.20	3 (4%) 40 32	240, 297, 336, 375	0
18	XR	70/88 (79%)	0.21	3 (4%) 40 32	168, 247, 321, 337	0
19	QS	83/93 (89%)	0.81	4 (4%) 36 30	308, 459, 605, 672	0
19	XS	84/93 (90%)	0.77	6 (7%) 23 21	341, 414, 456, 474	0
20	QT	99/106 (93%)	1.29	31 (31%) 1 3	205, 271, 358, 377	0
20	XT	99/106 (93%)	1.14	19 (19%) 4 6	198, 242, 278, 302	0
21	QU	25/27 (92%)	0.81	2 (8%) 20 18	300, 347, 463, 563	0
21	XU	25/27 (92%)	1.18	5 (20%) 3 6	336, 374, 530, 549	0
22	QV	68/77 (88%)	-0.19	1 (1%) 71 57	294, 387, 435, 446	0
22	XV	68/77 (88%)	-0.34	0 100 100	281, 411, 474, 504	0
23	QX	19/19 (100%)	0.21	0 100 100	368, 432, 461, 464	0
23	XX	19/19 (100%)	0.29	0 100 100	356, 422, 471, 476	0
24	R0	81/85 (95%)	0.79	15 (18%) 4 6	166, 206, 296, 364	0
24	Y0	82/85 (96%)	0.57	6 (7%) 22 20	120, 168, 274, 329	0
25	R1	95/98 (96%)	1.23	20 (21%) 3 5	162, 201, 262, 291	0
25	Y1	93/98 (94%)	1.06	15 (16%) 5 8	117, 176, 219, 232	0
26	R2	69/72 (95%)	0.38	4 (5%) 30 25	124, 192, 238, 283	0
26	Y2	68/72 (94%)	0.88	11 (16%) 5 8	124, 171, 214, 231	0
27	R3	59/60 (98%)	0.51	7 (11%) 10 11	175, 228, 281, 338	0
27	Y3	59/60 (98%)	0.48	4 (6%) 25 22	115, 168, 241, 269	0
28	R4	45/71 (63%)	-0.29	0 100 100	237, 382, 418, 429	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2		OWAB(Å ²)	Q<0.9
28	Y4	46/71 (64%)	-0.31	0	100 100	206, 273, 315, 322	0
29	R5	59/60 (98%)	0.19	1 (1%)	69 53	142, 193, 225, 291	0
29	Y5	59/60 (98%)	0.15	1 (1%)	69 53	122, 172, 225, 300	0
30	R6	53/54 (98%)	-0.27	1 (1%)	66 50	175, 230, 267, 357	0
30	Y6	53/54 (98%)	0.22	2 (3%)	44 35	144, 194, 248, 260	0
31	R7	47/49 (95%)	0.96	7 (14%)	7 9	138, 167, 186, 207	0
31	Y7	48/49 (97%)	0.75	9 (18%)	4 6	111, 146, 192, 224	0
32	R8	64/65 (98%)	0.75	7 (10%)	12 13	154, 201, 248, 284	0
32	Y8	64/65 (98%)	1.21	15 (23%)	2 4	120, 172, 209, 259	0
33	R9	37/37 (100%)	1.48	10 (27%)	2 4	260, 289, 340, 357	0
33	Y9	37/37 (100%)	1.25	6 (16%)	5 8	174, 205, 233, 250	0
34	RA	2882/2905 (99%)	-0.31	3 (0%)	92 89	123, 197, 357, 505	0
34	YA	2883/2905 (99%)	-0.32	13 (0%)	87 76	96, 149, 276, 493	0
35	RB	120/122 (98%)	-0.35	0	100 100	205, 256, 303, 350	0
35	YB	120/122 (98%)	-0.41	1 (0%)	82 69	135, 175, 210, 248	0
36	RD	272/276 (98%)	0.71	38 (13%)	7 10	127, 186, 231, 284	0
36	YD	272/276 (98%)	0.62	25 (9%)	16 15	107, 159, 207, 243	0
37	RE	205/206 (99%)	0.78	30 (14%)	7 9	128, 211, 265, 310	0
37	YE	205/206 (99%)	0.68	24 (11%)	10 11	102, 157, 205, 254	0
38	RF	202/210 (96%)	0.25	9 (4%)	39 31	143, 199, 272, 328	0
38	YF	202/210 (96%)	0.47	16 (7%)	20 18	101, 160, 215, 256	0
39	RG	181/182 (99%)	-0.04	5 (2%)	55 41	231, 307, 360, 388	0
39	YG	181/182 (99%)	0.06	9 (4%)	35 29	178, 226, 291, 327	0
40	RH	174/180 (96%)	0.23	12 (6%)	24 21	189, 269, 313, 358	0
40	YH	174/180 (96%)	0.36	11 (6%)	27 24	123, 175, 217, 258	0
41	RI	146/148 (98%)	0.29	11 (7%)	22 20	188, 254, 300, 327	0
41	YI	146/148 (98%)	0.06	7 (4%)	36 30	158, 226, 271, 293	0
42	RN	138/140 (98%)	0.49	14 (10%)	14 13	139, 219, 260, 268	0
42	YN	138/140 (98%)	0.71	18 (13%)	9 10	94, 160, 205, 231	0
43	RO	122/122 (100%)	0.34	8 (6%)	26 23	158, 202, 247, 286	0
43	YO	122/122 (100%)	0.11	1 (0%)	82 69	113, 159, 199, 213	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
44	RP	150/150 (100%)	0.72	27 (18%) 4 7	160, 228, 287, 331	0
44	YP	147/150 (98%)	0.66	16 (10%) 12 13	128, 166, 217, 255	0
45	RQ	141/141 (100%)	0.36	13 (9%) 16 15	194, 231, 267, 309	0
45	YQ	141/141 (100%)	0.37	5 (3%) 47 36	132, 174, 238, 314	0
46	RR	117/118 (99%)	0.94	19 (16%) 5 8	121, 180, 224, 257	0
46	YR	117/118 (99%)	0.98	27 (23%) 2 4	104, 155, 212, 237	0
47	RS	111/112 (99%)	0.40	10 (9%) 17 15	191, 262, 311, 357	0
47	YS	111/112 (99%)	0.70	22 (19%) 3 6	146, 196, 282, 328	0
48	RT	137/146 (93%)	0.33	11 (8%) 20 18	158, 225, 307, 320	0
48	YT	137/146 (93%)	0.46	14 (10%) 13 13	127, 185, 257, 282	0
49	RU	117/118 (99%)	0.70	13 (11%) 12 12	149, 243, 286, 304	0
49	YU	117/118 (99%)	0.56	7 (5%) 29 24	101, 147, 194, 223	0
50	RV	101/101 (100%)	0.19	10 (9%) 14 14	165, 222, 278, 288	0
50	YV	101/101 (100%)	0.35	12 (11%) 10 11	118, 170, 240, 280	0
51	RW	113/113 (100%)	0.59	10 (8%) 17 16	117, 160, 226, 339	0
51	YW	113/113 (100%)	0.62	12 (10%) 13 13	92, 137, 192, 239	0
52	RX	92/96 (95%)	0.58	6 (6%) 26 23	127, 177, 212, 235	0
52	YX	92/96 (95%)	0.76	11 (11%) 10 11	110, 146, 190, 259	0
53	RY	107/110 (97%)	0.50	6 (5%) 31 26	170, 219, 264, 315	0
53	YY	107/110 (97%)	0.29	3 (2%) 55 41	107, 170, 205, 244	0
54	RZ	183/206 (88%)	-0.07	4 (2%) 62 47	211, 258, 298, 354	0
54	YZ	193/206 (93%)	0.14	9 (4%) 37 30	124, 196, 287, 350	0
All	All	20769/21416 (96%)	0.16	1144 (5%) 32 26	92, 221, 460, 884	0

All (1144) RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
4	XD	4	TYR	11.2
17	QQ	24	GLU	9.4
2	XB	101	MET	9.1
8	QH	111	ILE	9.0
12	XL	28	LYS	8.6
17	QQ	7	THR	8.6
10	XJ	63	PHE	8.5

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Mol	Chain	Res	Type	RSRZ
45	YQ	38	GLU	8.5
13	QM	93	ARG	8.4
17	QQ	35	VAL	8.3
11	QK	129	SER	8.0
17	QQ	36	ILE	7.9
11	XK	126	ARG	7.8
36	RD	185	VAL	7.8
54	RZ	79	ARG	7.8
20	QT	70	SER	7.7
41	RI	37	VAL	7.7
47	YS	9	ARG	7.6
47	YS	4	LEU	7.4
4	QD	128	VAL	7.4
17	QQ	33	GLY	7.2
3	QC	129	ALA	7.1
26	Y2	15	LYS	7.1
11	XK	122	LYS	7.1
24	R0	3	HIS	7.1
36	RD	147	LEU	7.1
16	XP	29	ASP	7.0
20	XT	9	ASN	7.0
26	R2	8	LYS	6.9
50	RV	72	VAL	6.8
48	RT	1	MET	6.8
53	RY	1	MET	6.8
36	YD	184	LYS	6.7
47	RS	2	ALA	6.6
4	XD	2	GLY	6.6
48	RT	99	LEU	6.5
24	R0	76	GLY	6.5
19	XS	56	GLN	6.4
19	QS	33	THR	6.4
11	QK	128	ALA	6.3
38	RF	81	PRO	6.3
46	YR	48	VAL	6.3
8	QH	101	PRO	6.3
17	QQ	32	TYR	6.3
12	XL	20	LYS	6.1
3	QC	124	ILE	6.1
21	XU	4	GLY	6.1
17	QQ	34	LYS	6.0
16	QP	29	ASP	5.9

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Mol	Chain	Res	Type	RSRZ
4	XD	11	LEU	5.9
20	XT	11	SER	5.9
11	XK	125	PHE	5.9
47	YS	8	GLU	5.9
47	RS	3	ARG	5.8
14	XN	53	LEU	5.8
52	RX	34	ALA	5.8
20	QT	71	THR	5.7
6	QF	72	VAL	5.7
37	YE	113	PHE	5.7
3	XC	158	GLY	5.6
16	XP	1	MET	5.6
34	YA	2602	A	5.6
47	RS	8	GLU	5.5
8	QH	135	CYS	5.5
47	YS	3	ARG	5.5
46	RR	48	VAL	5.5
10	XJ	47	PHE	5.5
45	RQ	39	PRO	5.4
5	QE	26	PHE	5.4
14	XN	48	ALA	5.4
47	YS	7	TYR	5.4
46	RR	69	ASP	5.4
3	XC	178	LEU	5.4
12	XL	29	GLY	5.3
40	RH	165	ALA	5.3
38	RF	41	LEU	5.3
46	RR	66	VAL	5.3
44	YP	35	HIS	5.3
36	RD	6	PHE	5.3
3	QC	158	GLY	5.3
4	QD	146	ILE	5.3
16	QP	24	ALA	5.3
20	QT	56	MET	5.2
26	Y2	12	GLU	5.2
41	RI	38	LEU	5.2
10	QJ	58	ASP	5.2
3	QC	189	ALA	5.2
24	Y0	2	ALA	5.2
36	RD	5	LYS	5.2
3	XC	154	SER	5.2
37	RE	188	VAL	5.2

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Mol	Chain	Res	Type	RSRZ
16	XP	30	GLY	5.2
1	XA	1375	A	5.2
2	XB	71	VAL	5.1
37	RE	79	ARG	5.1
40	YH	64	LEU	5.1
8	QH	112	LEU	5.1
14	QN	11	LYS	5.1
17	QQ	37	LYS	5.1
37	RE	182	LEU	5.1
40	RH	150	ALA	5.1
5	XE	13	ILE	5.1
10	XJ	46	ARG	5.0
48	YT	106	SER	5.0
42	YN	75	TYR	5.0
25	R1	12	PRO	5.0
38	YF	181	LEU	5.0
48	YT	104	ASN	5.0
18	XR	31	LEU	5.0
8	QH	99	GLU	5.0
36	YD	169	GLU	5.0
16	XP	25	ARG	4.9
3	XC	165	THR	4.9
8	QH	134	ILE	4.9
16	XP	28	ARG	4.9
16	XP	60	LEU	4.9
11	QK	127	LYS	4.9
49	RU	37	GLU	4.9
2	XB	98	LEU	4.9
44	RP	110	TYR	4.9
52	RX	33	LYS	4.9
50	RV	73	SER	4.9
41	RI	36	ALA	4.9
37	RE	152	LYS	4.9
36	RD	184	LYS	4.8
46	YR	6	SER	4.8
37	RE	26	ILE	4.8
44	RP	71	VAL	4.8
48	YT	1	MET	4.8
3	XC	175	LEU	4.8
10	XJ	65	LEU	4.8
16	QP	25	ARG	4.8
20	QT	72	LEU	4.8

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Mol	Chain	Res	Type	RSRZ
20	XT	10	LEU	4.8
15	XO	60	VAL	4.8
16	XP	27	LYS	4.7
15	QO	66	LEU	4.7
20	XT	72	LEU	4.7
16	XP	31	LYS	4.7
46	RR	70	LEU	4.7
47	YS	11	LYS	4.7
50	YV	81	TYR	4.7
4	QD	94	LEU	4.7
44	RP	50	ARG	4.7
1	QA	1065	U	4.7
41	YI	21	VAL	4.6
12	QL	32	PHE	4.6
8	XH	112	LEU	4.6
46	YR	69	ASP	4.6
16	QP	70	ALA	4.6
34	RA	2319	G	4.6
36	RD	211	ARG	4.6
10	XJ	58	ASP	4.6
3	XC	155	GLY	4.6
46	YR	29	LEU	4.6
15	XO	63	ARG	4.6
47	YS	17	ARG	4.6
16	XP	65	GLN	4.6
33	R9	28	GLU	4.6
52	YX	62	LYS	4.5
40	RH	105	LEU	4.5
4	QD	68	TYR	4.5
46	RR	65	LEU	4.5
5	XE	81	GLU	4.5
20	QT	9	ASN	4.5
2	XB	97	TRP	4.5
3	QC	130	VAL	4.5
4	XD	5	ILE	4.5
51	YW	98	LYS	4.5
16	XP	4	ILE	4.4
4	XD	64	LEU	4.4
5	QE	43	LEU	4.4
4	QD	139	ARG	4.4
46	RR	68	ARG	4.4
15	QO	54	ARG	4.4

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Mol	Chain	Res	Type	RSRZ
16	QP	28	ARG	4.4
47	YS	92	TYR	4.4
31	R7	10	ARG	4.4
48	RT	102	ILE	4.4
19	XS	55	LYS	4.4
13	XM	82	MET	4.4
4	QD	67	ILE	4.4
20	XT	68	LYS	4.4
5	XE	25	ARG	4.4
20	QT	11	SER	4.4
31	R7	46	VAL	4.3
1	XA	134	A	4.3
40	RH	151	ILE	4.3
12	XL	89	ARG	4.3
50	YV	74	LYS	4.3
36	RD	226	MET	4.3
36	RD	36	PRO	4.3
17	QQ	26	GLN	4.3
3	QC	128	PHE	4.3
8	XH	91	ARG	4.3
36	RD	181	GLU	4.3
48	YT	109	GLU	4.3
42	RN	99	LEU	4.2
37	YE	156	MET	4.2
8	QH	120	THR	4.2
36	YD	170	GLY	4.2
8	QH	19	VAL	4.2
11	XK	124	LYS	4.2
31	R7	14	LYS	4.2
36	YD	38	LYS	4.2
37	RE	111	ARG	4.2
47	YS	13	ARG	4.2
48	RT	50	ILE	4.2
42	YN	82	LEU	4.2
14	QN	53	LEU	4.2
37	RE	121	ASN	4.2
50	RV	81	TYR	4.2
20	XT	60	GLU	4.2
37	RE	123	ALA	4.2
11	QK	84	VAL	4.2
20	QT	8	ARG	4.2
12	XL	120	TYR	4.2

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Mol	Chain	Res	Type	RSRZ
20	XT	12	ALA	4.2
6	XF	79	LEU	4.2
12	QL	93	LEU	4.2
2	QB	101	MET	4.2
42	RN	84	LYS	4.2
48	RT	94	ALA	4.2
5	XE	24	ARG	4.1
54	RZ	78	LYS	4.1
1	QA	60	A	4.1
20	XT	30	LYS	4.1
6	XF	72	VAL	4.1
31	Y7	15	THR	4.1
10	XJ	62	HIS	4.1
33	R9	25	VAL	4.1
18	QR	66	LEU	4.1
16	XP	3	LYS	4.1
53	YY	63	LYS	4.1
8	QH	95	VAL	4.1
12	XL	10	LEU	4.1
49	RU	35	ALA	4.1
8	XH	111	ILE	4.0
37	RE	156	MET	4.0
49	RU	91	ASP	4.0
4	QD	64	LEU	4.0
37	YE	195	LEU	4.0
10	XJ	64	GLU	4.0
25	R1	26	ARG	4.0
47	RS	4	LEU	4.0
2	QB	163	PHE	4.0
42	YN	81	GLY	4.0
31	Y7	48	LYS	4.0
32	Y8	11	LYS	4.0
36	RD	155	LEU	4.0
37	RE	195	LEU	4.0
24	R0	74	ARG	4.0
41	RI	3	VAL	4.0
2	XB	163	PHE	4.0
8	XH	80	ILE	4.0
50	RV	74	LYS	3.9
20	QT	67	ALA	3.9
52	YX	28	PHE	3.9
7	QG	8	GLU	3.9

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Mol	Chain	Res	Type	RSRZ
8	XH	133	LEU	3.9
51	YW	96	ILE	3.9
12	QL	86	ARG	3.9
20	QT	22	ARG	3.9
36	RD	259	THR	3.9
30	R6	4	GLU	3.9
53	YY	64	GLU	3.9
1	XA	1376	U	3.9
20	QT	13	LEU	3.9
24	R0	71	ASP	3.9
31	R7	22	MET	3.9
34	YA	2181	G	3.9
31	R7	45	ALA	3.9
42	RN	115	ARG	3.9
25	R1	3	LYS	3.9
45	YQ	39	PRO	3.9
12	QL	10	LEU	3.8
24	R0	75	LEU	3.8
41	RI	92	VAL	3.8
27	R3	14	GLY	3.8
25	R1	21	ARG	3.8
5	XE	129	ILE	3.8
7	XG	8	GLU	3.8
25	R1	27	GLU	3.8
11	XK	121	PRO	3.8
36	RD	173	VAL	3.8
14	QN	13	THR	3.8
52	RX	39	ILE	3.8
50	RV	77	ALA	3.8
12	XL	7	ILE	3.8
16	XP	2	VAL	3.8
47	YS	28	VAL	3.8
42	YN	76	SER	3.8
2	XB	108	ILE	3.7
17	XQ	83	ASP	3.7
24	R0	4	LYS	3.7
37	RE	128	SER	3.7
20	QT	12	ALA	3.7
36	RD	258	LYS	3.7
42	YN	46	VAL	3.7
44	RP	3	LEU	3.7
46	RR	71	GLN	3.7

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Mol	Chain	Res	Type	RSRZ
3	XC	153	VAL	3.7
47	YS	12	PHE	3.7
37	YE	76	ARG	3.7
16	QP	7	ALA	3.7
38	YF	37	VAL	3.7
24	R0	77	ARG	3.7
37	RE	151	TYR	3.7
40	YH	67	LEU	3.7
41	RI	21	VAL	3.7
41	YI	30	LEU	3.7
3	QC	120	VAL	3.7
17	XQ	24	GLU	3.7
25	R1	11	ARG	3.7
54	YZ	79	ARG	3.7
36	RD	16	MET	3.6
6	XF	95	GLU	3.6
3	QC	152	ILE	3.6
8	QH	100	ILE	3.6
41	RI	1	MET	3.6
2	QB	173	ALA	3.6
31	R7	18	PHE	3.6
17	QQ	30	PRO	3.6
12	QL	18	VAL	3.6
20	XT	64	ASP	3.6
11	XK	120	ARG	3.6
15	XO	68	ARG	3.6
45	RQ	100	GLY	3.6
52	RX	68	ARG	3.6
46	RR	47	PHE	3.6
46	RR	67	LEU	3.6
24	R0	45	PHE	3.6
20	QT	74	LYS	3.6
16	XP	74	LEU	3.6
3	XC	162	GLN	3.6
51	RW	99	ARG	3.6
44	RP	51	PHE	3.6
16	QP	68	ASP	3.6
3	QC	178	LEU	3.6
44	YP	3	LEU	3.6
33	R9	24	TYR	3.5
12	XL	27	LEU	3.5
26	Y2	20	GLU	3.5

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Mol	Chain	Res	Type	RSRZ
4	QD	77	ASN	3.5
7	XG	28	ASN	3.5
25	Y1	4	VAL	3.5
48	RT	22	PHE	3.5
36	YD	174	ILE	3.5
15	XO	65	ARG	3.5
49	RU	36	ARG	3.5
36	YD	5	LYS	3.5
15	XO	32	LEU	3.5
48	RT	114	LEU	3.5
47	YS	16	ASN	3.5
13	XM	90	LEU	3.5
17	QQ	39	SER	3.5
20	QT	73	HIS	3.5
7	QG	33	ASP	3.5
10	XJ	60	ARG	3.5
37	YE	7	VAL	3.5
43	RO	38	VAL	3.5
51	YW	86	LEU	3.5
15	QO	62	GLN	3.5
36	RD	28	GLU	3.4
16	XP	9	PHE	3.4
38	YF	41	LEU	3.4
1	XA	1367	C	3.4
16	QP	5	ARG	3.4
53	RY	33	LYS	3.4
52	YX	54	VAL	3.4
16	XP	22	THR	3.4
25	Y1	11	ARG	3.4
40	RH	166	GLY	3.4
14	QN	14	PRO	3.4
26	Y2	21	LEU	3.4
38	YF	50	SER	3.4
51	RW	98	LYS	3.4
2	QB	214	ILE	3.4
44	RP	26	GLY	3.4
8	QH	133	LEU	3.4
11	QK	125	PHE	3.4
5	XE	14	ARG	3.4
12	XL	19	ARG	3.4
37	YE	149	ARG	3.4
54	YZ	166	SER	3.4

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Mol	Chain	Res	Type	RSRZ
17	XQ	36	ILE	3.4
44	RP	35	HIS	3.4
47	RS	5	THR	3.4
20	XT	56	MET	3.4
1	QA	1112	C	3.4
51	RW	81	ALA	3.4
25	Y1	38	SER	3.4
46	YR	44	LEU	3.4
11	QK	11	LYS	3.4
36	RD	40	THR	3.4
4	QD	93	PHE	3.4
7	QG	37	ASN	3.3
20	QT	26	ASN	3.3
20	XT	63	ILE	3.3
25	R1	10	LYS	3.3
25	Y1	92	LYS	3.3
42	YN	83	LYS	3.3
15	XO	36	ILE	3.3
15	XO	66	LEU	3.3
3	XC	167	TRP	3.3
36	RD	52	ARG	3.3
36	RD	169	GLU	3.3
17	XQ	37	LYS	3.3
33	Y9	15	LYS	3.3
36	YD	168	ARG	3.3
12	QL	7	ILE	3.3
21	XU	3	LYS	3.3
36	YD	175	LEU	3.3
44	RP	53	GLY	3.3
33	Y9	26	ILE	3.3
20	QT	27	LYS	3.3
36	RD	42	GLY	3.3
47	YS	2	ALA	3.3
36	YD	44	ASN	3.3
20	QT	30	LYS	3.3
6	QF	53	ALA	3.3
40	YH	57	ASP	3.3
2	QB	97	TRP	3.3
7	XG	11	GLN	3.3
14	XN	61	TRP	3.3
33	R9	23	VAL	3.3
37	RE	163	GLU	3.3

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Mol	Chain	Res	Type	RSRZ
15	QO	58	MET	3.3
21	QU	16	GLY	3.2
32	Y8	20	GLY	3.2
4	XD	135	LEU	3.2
46	RR	44	LEU	3.2
12	QL	19	ARG	3.2
26	Y2	13	ALA	3.2
54	YZ	193	GLU	3.2
47	YS	20	ARG	3.2
51	RW	42	ARG	3.2
49	YU	7	GLY	3.2
26	Y2	66	GLU	3.2
48	RT	11	GLU	3.2
50	RV	76	LYS	3.2
1	QA	1111	A	3.2
20	XT	18	GLN	3.2
36	YD	41	GLY	3.2
50	RV	78	LYS	3.2
32	Y8	54	GLU	3.2
36	RD	175	LEU	3.2
5	XE	18	ARG	3.2
17	XQ	84	LEU	3.2
45	YQ	91	GLU	3.2
38	RF	184	TYR	3.2
47	RS	20	ARG	3.2
53	RY	20	TYR	3.2
32	Y8	63	PRO	3.2
25	R1	95	LEU	3.2
46	YR	47	PHE	3.2
33	R9	29	ASN	3.2
49	YU	91	ASP	3.1
25	Y1	37	ILE	3.1
44	RP	40	SER	3.1
8	QH	93	VAL	3.1
16	XP	68	ASP	3.1
33	Y9	25	VAL	3.1
8	XH	18	ARG	3.1
25	Y1	70	VAL	3.1
37	YE	188	VAL	3.1
54	YZ	168	GLU	3.1
17	QQ	22	LEU	3.1
12	XL	126	LYS	3.1

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Mol	Chain	Res	Type	RSRZ
2	XB	68	ILE	3.1
25	R1	7	ILE	3.1
36	RD	41	GLY	3.1
1	XA	135	C	3.1
6	QF	90	VAL	3.1
34	YA	2139	C	3.1
16	XP	34	GLU	3.1
16	QP	27	LYS	3.1
1	XA	1374	A	3.1
49	RU	79	PHE	3.1
15	QO	56	LEU	3.1
34	YA	2334	G	3.1
8	QH	67	PRO	3.1
37	RE	126	PRO	3.1
39	YG	11	TYR	3.1
31	Y7	47	ARG	3.1
33	Y9	34	GLN	3.1
36	YD	271	ILE	3.1
24	R0	46	LYS	3.1
51	YW	46	PHE	3.0
39	YG	34	LEU	3.0
42	YN	45	ASN	3.0
42	YN	109	LYS	3.0
44	RP	43	GLY	3.0
27	Y3	57	GLU	3.0
46	YR	24	GLN	3.0
7	XG	5	ARG	3.0
25	R1	41	ARG	3.0
49	RU	28	ARG	3.0
1	QA	378	G	3.0
8	XH	101	PRO	3.0
50	YV	76	LYS	3.0
12	XL	14	GLY	3.0
46	YR	7	GLY	3.0
20	QT	37	SER	3.0
38	RF	98	SER	3.0
12	XL	16	GLU	3.0
34	RA	2602	A	3.0
4	XD	66	ARG	3.0
16	XP	71	ARG	3.0
36	YD	134	ARG	3.0
48	YT	105	LEU	3.0

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Mol	Chain	Res	Type	RSRZ
45	RQ	40	ALA	3.0
24	R0	53	MET	3.0
16	XP	32	TYR	3.0
40	RH	86	GLU	3.0
8	QH	123	GLU	3.0
16	QP	4	ILE	3.0
20	QT	18	GLN	3.0
38	YF	55	GLY	3.0
40	YH	71	LEU	3.0
36	RD	35	LYS	3.0
44	YP	65	ARG	3.0
5	QE	81	GLU	3.0
20	QT	64	ASP	3.0
7	QG	9	VAL	3.0
16	XP	24	ALA	3.0
46	YR	25	ALA	3.0
2	XB	51	LEU	3.0
20	XT	71	THR	2.9
4	QD	80	GLU	2.9
6	XF	53	ALA	2.9
38	RF	37	VAL	2.9
46	YR	65	LEU	2.9
6	QF	96	PRO	2.9
4	XD	3	ARG	2.9
48	RT	103	ARG	2.9
25	Y1	12	PRO	2.9
37	RE	153	GLY	2.9
49	RU	94	ASN	2.9
49	YU	44	ASN	2.9
54	YZ	74	VAL	2.9
45	RQ	21	THR	2.9
49	RU	19	LYS	2.9
42	RN	73	THR	2.9
32	R8	11	LYS	2.9
32	R8	54	GLU	2.9
16	XP	66	PRO	2.9
33	R9	30	PRO	2.9
48	YT	102	ILE	2.9
50	RV	71	LEU	2.9
36	RD	38	LYS	2.9
42	YN	68	GLU	2.9
49	YU	37	GLU	2.9

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Mol	Chain	Res	Type	RSRZ
16	XP	26	ARG	2.9
31	R7	23	ARG	2.9
25	R1	44	PRO	2.9
46	RR	51	LEU	2.9
1	QA	1246	C	2.9
20	QT	23	ARG	2.9
14	XN	54	PRO	2.9
4	XD	70	ILE	2.9
17	XQ	4	LYS	2.9
8	XH	93	VAL	2.9
14	XN	56	VAL	2.9
44	YP	75	ILE	2.9
20	XT	13	LEU	2.9
49	RU	34	LYS	2.8
37	YE	157	ALA	2.8
3	QC	131	ARG	2.8
9	XI	71	SER	2.8
4	XD	93	PHE	2.8
2	QB	102	LEU	2.8
40	YH	107	VAL	2.8
4	QD	4	TYR	2.8
11	XK	11	LYS	2.8
24	Y0	45	PHE	2.8
16	QP	66	PRO	2.8
8	QH	36	LEU	2.8
11	XK	123	LYS	2.8
15	QO	67	LEU	2.8
32	Y8	65	GLU	2.8
44	RP	70	GLN	2.8
33	R9	9	ARG	2.8
37	YE	79	ARG	2.8
46	YR	2	ARG	2.8
32	Y8	19	SER	2.8
49	RU	30	LYS	2.8
12	QL	30	ALA	2.8
25	Y1	41	ARG	2.8
33	R9	20	HIS	2.8
33	Y9	12	ASP	2.8
2	XB	200	ILE	2.8
4	QD	126	ILE	2.8
25	Y1	91	LYS	2.8
46	RR	42	LYS	2.8

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Mol	Chain	Res	Type	RSRZ
51	YW	21	VAL	2.8
4	XD	15	GLU	2.8
44	RP	41	ARG	2.8
38	YF	39	TRP	2.8
1	XA	1368	G	2.8
36	YD	15	PHE	2.8
41	RI	30	LEU	2.8
44	RP	46	LYS	2.8
14	XN	51	GLY	2.8
36	RD	183	ARG	2.8
46	RR	10	LEU	2.8
2	QB	71	VAL	2.8
16	QP	2	VAL	2.8
50	YV	70	ILE	2.8
20	QT	15	ARG	2.8
21	XU	6	ARG	2.8
39	YG	35	GLU	2.8
16	XP	70	ALA	2.8
20	QT	66	ALA	2.8
20	XT	58	LYS	2.8
8	QH	119	LEU	2.8
45	RQ	17	LEU	2.8
40	YH	151	ILE	2.8
42	YN	115	ARG	2.8
20	QT	60	GLU	2.8
26	R2	60	LEU	2.7
2	XB	58	ILE	2.7
4	XD	133	VAL	2.7
14	QN	25	VAL	2.7
44	YP	43	GLY	2.7
31	Y7	1	MET	2.7
50	YV	78	LYS	2.7
8	QH	127	LEU	2.7
37	YE	182	LEU	2.7
36	YD	173	VAL	2.7
37	RE	3	GLY	2.7
37	RE	76	ARG	2.7
38	YF	44	ARG	2.7
12	QL	98	TYR	2.7
14	XN	44	LEU	2.7
37	RE	25	VAL	2.7
1	QA	966	G	2.7

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Mol	Chain	Res	Type	RSRZ
27	R3	12	PRO	2.7
3	QC	199	LYS	2.7
32	Y8	40	GLU	2.7
2	XB	165	VAL	2.7
39	YG	161	THR	2.7
19	XS	20	LEU	2.7
12	XL	11	VAL	2.7
39	RG	157	ILE	2.7
37	RE	96	PHE	2.7
42	RN	68	GLU	2.7
45	RQ	47	ILE	2.7
44	YP	36	LYS	2.7
49	RU	22	LYS	2.7
15	XO	59	MET	2.7
43	RO	2	ILE	2.7
47	RS	7	TYR	2.7
48	YT	22	PHE	2.7
26	R2	11	GLU	2.7
37	YE	152	LYS	2.7
1	QA	134	A	2.7
5	QE	106	PRO	2.7
25	Y1	73	LEU	2.7
44	YP	56	SER	2.7
20	XT	55	ILE	2.7
7	XG	10	ARG	2.7
14	QN	12	ARG	2.7
38	YF	40	GLN	2.7
48	YT	46	GLU	2.7
24	Y0	3	HIS	2.7
51	RW	82	LEU	2.7
34	YA	949	C	2.7
37	YE	68	ALA	2.7
5	QE	15	ARG	2.7
17	QQ	25	ARG	2.7
51	YW	92	ARG	2.7
47	YS	88	ASP	2.6
37	YE	121	ASN	2.6
30	Y6	11	LEU	2.6
36	RD	133	LEU	2.6
45	RQ	37	LEU	2.6
48	RT	96	ARG	2.6
40	RH	153	LYS	2.6

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Mol	Chain	Res	Type	RSRZ
33	R9	12	ASP	2.6
37	YE	27	LEU	2.6
42	YN	26	LEU	2.6
50	YV	80	GLN	2.6
52	YX	94	GLY	2.6
12	XL	90	VAL	2.6
26	Y2	68	ARG	2.6
9	XI	70	LYS	2.6
20	QT	29	LYS	2.6
38	YF	110	LEU	2.6
40	RH	120	GLY	2.6
42	YN	77	GLY	2.6
43	RO	74	GLY	2.6
38	YF	56	GLU	2.6
51	YW	40	ASN	2.6
5	QE	128	PRO	2.6
5	XE	21	ALA	2.6
8	QH	13	ILE	2.6
25	Y1	90	ILE	2.6
51	YW	17	VAL	2.6
48	YT	76	PHE	2.6
38	YF	184	TYR	2.6
37	YE	153	GLY	2.6
46	YR	72	ASP	2.6
4	XD	24	GLU	2.6
12	XL	17	LYS	2.6
31	Y7	11	LYS	2.6
6	XF	75	LEU	2.6
37	YE	71	GLY	2.6
37	RE	155	LYS	2.6
39	RG	75	LYS	2.6
45	YQ	139	GLU	2.6
2	XB	26	PRO	2.6
4	XD	37	PRO	2.6
42	RN	79	PRO	2.6
36	YD	133	LEU	2.6
46	YR	14	SER	2.6
52	RX	65	ARG	2.6
19	QS	50	ALA	2.6
42	RN	103	VAL	2.6
43	RO	1	MET	2.6
27	R3	26	LEU	2.6

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Mol	Chain	Res	Type	RSRZ
21	QU	14	TRP	2.6
38	YF	34	TRP	2.6
13	XM	85	GLY	2.6
3	XC	4	LYS	2.6
44	RP	33	ARG	2.6
10	QJ	59	SER	2.6
51	YW	97	LYS	2.6
44	RP	57	THR	2.6
44	RP	52	GLU	2.6
46	YR	41	ALA	2.6
24	Y0	46	LYS	2.5
51	YW	74	ALA	2.5
1	QA	108	G	2.5
36	RD	154	LYS	2.5
36	YD	183	ARG	2.5
44	YP	24	GLY	2.5
46	YR	8	ARG	2.5
38	YF	82	ILE	2.5
36	RD	215	LEU	2.5
45	RQ	80	GLU	2.5
46	YR	102	GLU	2.5
17	XQ	34	LYS	2.5
26	Y2	69	ARG	2.5
1	QA	1183	A	2.5
35	YB	88	C	2.5
47	YS	18	ILE	2.5
44	YP	6	LEU	2.5
13	QM	113	PRO	2.5
54	RZ	166	SER	2.5
16	XP	23	ASP	2.5
12	QL	99	HIS	2.5
37	RE	2	LYS	2.5
31	Y7	23	ARG	2.5
44	RP	79	ARG	2.5
1	QA	325	A	2.5
44	RP	54	GLY	2.5
1	XA	769	G	2.5
1	QA	133	U	2.5
12	QL	128	ALA	2.5
53	RY	43	ASN	2.5
36	YD	36	PRO	2.5
36	YD	14	ARG	2.5

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Mol	Chain	Res	Type	RSRZ
52	YX	56	THR	2.5
1	XA	1054	C	2.5
1	QA	1248	A	2.5
4	QD	5	ILE	2.5
52	YX	71	GLY	2.5
27	R3	15	TYR	2.5
46	YR	51	LEU	2.5
16	XP	7	ALA	2.5
21	XU	14	TRP	2.5
25	Y1	3	LYS	2.5
50	YV	84	LYS	2.5
24	R0	72	ARG	2.5
25	Y1	27	GLU	2.5
1	XA	1248	A	2.5
4	QD	133	VAL	2.5
25	R1	22	GLY	2.5
8	XH	119	LEU	2.5
20	QT	53	LEU	2.5
32	Y8	60	LEU	2.5
37	RE	160	TYR	2.5
2	XB	152	PHE	2.5
16	QP	26	ARG	2.5
26	Y2	70	GLN	2.5
8	XH	120	THR	2.5
36	YD	185	VAL	2.5
37	YE	134	ILE	2.5
45	RQ	90	VAL	2.5
47	YS	14	VAL	2.5
19	XS	16	LEU	2.5
37	RE	1	MET	2.5
15	QO	35	ARG	2.5
17	QQ	27	PHE	2.5
3	QC	198	VAL	2.5
47	YS	98	VAL	2.5
9	QI	70	LYS	2.4
32	Y8	51	ALA	2.4
37	YE	28	ALA	2.4
37	YE	205	ALA	2.4
50	YV	83	ARG	2.4
17	XQ	30	PRO	2.4
22	QV	75	C	2.4
34	YA	2180	U	2.4

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Mol	Chain	Res	Type	RSRZ
46	YR	79	LEU	2.4
42	RN	80	GLY	2.4
46	YR	99	LYS	2.4
31	Y7	45	ALA	2.4
44	YP	110	TYR	2.4
45	RQ	25	ASP	2.4
1	QA	1186	G	2.4
2	XB	70	PHE	2.4
15	QO	68	ARG	2.4
45	RQ	104	PHE	2.4
51	RW	46	PHE	2.4
1	XA	1249	C	2.4
2	XB	196	LEU	2.4
19	XS	9	VAL	2.4
41	YI	44	LEU	2.4
41	YI	114	LEU	2.4
32	Y8	5	LYS	2.4
24	R0	78	TYR	2.4
25	R1	71	TYR	2.4
39	YG	178	PHE	2.4
51	RW	63	ASP	2.4
40	RH	172	LYS	2.4
41	YI	99	GLU	2.4
17	QQ	29	HIS	2.4
48	RT	100	TYR	2.4
43	RO	81	ASP	2.4
4	QD	137	SER	2.4
15	QO	70	LEU	2.4
25	R1	70	VAL	2.4
40	RH	103	LEU	2.4
42	YN	138	LEU	2.4
43	YO	86	ILE	2.4
44	RP	39	LYS	2.4
5	QE	22	GLY	2.4
4	XD	10	ARG	2.4
45	RQ	38	GLU	2.4
4	QD	104	VAL	2.4
27	R3	6	VAL	2.4
44	RP	36	LYS	2.4
46	YR	5	LYS	2.4
3	QC	10	PHE	2.4
37	RE	122	PHE	2.4

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Mol	Chain	Res	Type	RSRZ
54	YZ	88	PHE	2.4
8	XH	135	CYS	2.4
17	QQ	56	VAL	2.4
1	XA	1538	C	2.4
27	R3	20	LYS	2.4
46	RR	73	VAL	2.4
41	RI	20	ASP	2.4
48	YT	86	ILE	2.4
52	YX	53	LYS	2.4
4	QD	71	SER	2.4
38	YF	61	GLY	2.4
46	YR	71	GLN	2.4
54	YZ	192	ALA	2.4
43	RO	7	TYR	2.4
12	XL	13	LYS	2.4
37	RE	184	VAL	2.4
37	YE	133	LYS	2.4
42	RN	54	VAL	2.4
49	YU	16	LYS	2.4
2	XB	194	PRO	2.4
4	QD	51	PRO	2.4
24	R0	43	THR	2.4
9	XI	106	ALA	2.3
52	RX	70	LEU	2.3
17	XQ	56	VAL	2.3
20	QT	63	ILE	2.3
3	XC	164	ARG	2.3
24	R0	6	GLY	2.3
20	QT	59	ALA	2.3
38	YF	192	LEU	2.3
40	YH	105	LEU	2.3
42	YN	101	HIS	2.3
1	QA	107	G	2.3
20	QT	69	GLY	2.3
4	QD	103	ASN	2.3
5	QE	31	LEU	2.3
14	XN	60	SER	2.3
11	QK	32	ILE	2.3
49	YU	62	ILE	2.3
8	QH	14	ARG	2.3
46	YR	17	ARG	2.3
20	XT	69	GLY	2.3

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Mol	Chain	Res	Type	RSRZ
32	R8	51	ALA	2.3
41	RI	35	LEU	2.3
14	XN	37	PHE	2.3
44	RP	38	GLN	2.3
46	RR	24	GLN	2.3
39	RG	39	ILE	2.3
50	RV	79	VAL	2.3
50	YV	79	VAL	2.3
2	XB	137	ARG	2.3
4	QD	3	ARG	2.3
25	Y1	6	GLU	2.3
38	RF	44	ARG	2.3
14	XN	15	LYS	2.3
16	XP	6	LEU	2.3
39	YG	173	LEU	2.3
42	RN	2	LYS	2.3
38	YF	73	ALA	2.3
47	YS	6	ALA	2.3
26	Y2	63	VAL	2.3
44	YP	23	PRO	2.3
54	RZ	169	GLU	2.3
5	QE	110	LEU	2.3
19	QS	30	LEU	2.3
32	Y8	59	LYS	2.3
42	YN	87	LEU	2.3
25	R1	4	VAL	2.3
36	RD	174	ILE	2.3
39	RG	41	GLN	2.3
51	YW	11	ARG	2.3
8	QH	132	GLU	2.3
39	YG	168	GLU	2.3
4	XD	78	LEU	2.3
24	Y0	49	LYS	2.3
37	RE	118	LYS	2.3
49	RU	27	LEU	2.3
51	RW	86	LEU	2.3
3	XC	163	ALA	2.3
20	QT	41	ILE	2.3
42	RN	122	VAL	2.3
45	YQ	47	ILE	2.3
8	QH	84	ARG	2.3
32	Y8	57	ARG	2.3

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Mol	Chain	Res	Type	RSRZ
37	RE	74	PRO	2.3
50	YV	85	LYS	2.3
26	Y2	16	LEU	2.3
31	Y7	31	LEU	2.3
32	R8	50	LEU	2.3
38	RF	71	GLY	2.3
15	QO	60	VAL	2.2
40	YH	115	VAL	2.2
41	RI	4	ILE	2.2
6	QF	75	LEU	2.2
39	YG	32	PRO	2.2
48	YT	11	GLU	2.2
36	RD	267	SER	2.2
14	XN	59	ALA	2.2
32	R8	48	PHE	2.2
32	Y8	10	ALA	2.2
40	YH	113	VAL	2.2
43	RO	111	PHE	2.2
52	YX	59	VAL	2.2
13	XM	93	ARG	2.2
34	YA	2142	C	2.2
34	YA	2178	C	2.2
32	R8	59	LYS	2.2
32	Y8	3	LYS	2.2
18	QR	76	LEU	2.2
36	RD	230	ASP	2.2
4	QD	98	GLU	2.2
8	QH	136	GLU	2.2
10	QJ	62	HIS	2.2
27	R3	60	GLU	2.2
44	YP	54	GLY	2.2
17	QQ	11	VAL	2.2
25	R1	20	ARG	2.2
40	RH	170	ARG	2.2
44	YP	61	ARG	2.2
47	YS	30	ARG	2.2
1	QA	135	C	2.2
15	QO	57	LEU	2.2
31	Y7	42	LEU	2.2
8	QH	9	MET	2.2
11	XK	113	PRO	2.2
38	RF	70	THR	2.2

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Mol	Chain	Res	Type	RSRZ
46	RR	72	ASP	2.2
19	XS	17	GLU	2.2
20	XT	25	ARG	2.2
36	YD	224	ALA	2.2
41	YI	3	VAL	2.2
48	YT	61	PHE	2.2
52	YX	52	VAL	2.2
34	YA	2319	G	2.2
42	RN	83	LYS	2.2
44	RP	45	LEU	2.2
42	YN	72	TYR	2.2
6	QF	83	ASP	2.2
8	QH	137	VAL	2.2
12	XL	95	GLY	2.2
16	XP	5	ARG	2.2
17	XQ	80	GLY	2.2
20	QT	16	HIS	2.2
20	XT	66	ALA	2.2
29	Y5	2	ALA	2.2
20	QT	24	LEU	2.2
36	RD	177	LEU	2.2
46	YR	70	LEU	2.2
4	QD	207	TYR	2.2
24	Y0	53	MET	2.2
12	QL	101	VAL	2.2
16	QP	80	PHE	2.2
29	R5	24	ALA	2.2
37	YE	204	ALA	2.2
17	QQ	86	GLU	2.2
18	XR	38	GLU	2.2
24	R0	19	LYS	2.2
34	YA	985	C	2.2
48	YT	114	LEU	2.2
17	XQ	32	TYR	2.2
42	RN	46	VAL	2.2
1	XA	1364	U	2.2
37	YE	127	ASP	2.2
38	RF	100	THR	2.2
44	YP	76	LYS	2.2
4	XD	144	ASP	2.2
15	XO	57	LEU	2.2
36	RD	186	HIS	2.2

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Mol	Chain	Res	Type	RSRZ
13	XM	113	PRO	2.2
18	XR	80	PRO	2.2
39	RG	21	ARG	2.2
14	XN	11	LYS	2.2
44	RP	29	LYS	2.2
21	XU	2	GLY	2.2
10	XJ	67	THR	2.2
18	QR	58	LEU	2.2
37	RE	24	THR	2.2
12	XL	23	LYS	2.1
16	XP	33	ILE	2.1
37	RE	198	VAL	2.1
39	YG	20	ILE	2.1
15	QO	55	GLY	2.1
37	YE	6	GLY	2.1
34	YA	2179	C	2.1
44	RP	6	LEU	2.1
47	RS	32	LEU	2.1
1	XA	1187	G	2.1
2	QB	175	ARG	2.1
17	QQ	38	ARG	2.1
17	XQ	101	ARG	2.1
5	XE	88	LYS	2.1
46	RR	52	ILE	2.1
36	YD	22	SER	2.1
47	RS	12	PHE	2.1
36	RD	237	GLU	2.1
42	YN	73	THR	2.1
36	RD	273	ARG	2.1
2	QB	93	VAL	2.1
41	YI	1	MET	2.1
17	XQ	6	LEU	2.1
37	YE	123	ALA	2.1
46	YR	21	TYR	2.1
34	YA	2107	C	2.1
8	QH	15	ASN	2.1
12	XL	99	HIS	2.1
11	QK	110	ASP	2.1
51	RW	22	ASP	2.1
27	Y3	50	VAL	2.1
37	RE	116	VAL	2.1
44	YP	114	ILE	2.1

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Mol	Chain	Res	Type	RSRZ
43	RO	84	ALA	2.1
25	Y1	20	ARG	2.1
27	Y3	20	LYS	2.1
33	Y9	22	ARG	2.1
44	RP	16	ARG	2.1
2	XB	174	VAL	2.1
12	XL	18	VAL	2.1
13	XM	78	ILE	2.1
26	R2	52	ASP	2.1
49	RU	110	VAL	2.1
53	YY	24	VAL	2.1
1	QA	1187	G	2.1
34	YA	2182	G	2.1
5	QE	132	ALA	2.1
16	XP	64	ALA	2.1
36	YD	240	ALA	2.1
44	YP	68	GLN	2.1
30	Y6	53	LYS	2.1
49	YU	28	ARG	2.1
40	RH	147	ASN	2.1
8	QH	83	ILE	2.1
15	QO	4	THR	2.1
25	R1	46	LEU	2.1
1	QA	1367	C	2.1
4	XD	130	GLY	2.1
32	R8	23	VAL	2.1
50	YV	72	VAL	2.1
53	RY	6	HIS	2.1
32	Y8	48	PHE	2.1
36	YD	124	PRO	2.1
44	RP	31	ALA	2.1
1	QA	551	U	2.1
46	YR	40	LYS	2.1
47	RS	11	LYS	2.1
50	YV	82	ARG	2.1
8	QH	6	ILE	2.0
19	QS	49	ILE	2.0
33	R9	11	CYS	2.0
42	YN	99	LEU	2.0
46	RR	28	LEU	2.0
36	RD	32	SER	2.0
1	QA	1366	C	2.0

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Mol	Chain	Res	Type	RSRZ
44	RP	28	GLY	2.0
47	YS	5	THR	2.0
47	YS	91	PRO	2.0
51	RW	83	LYS	2.0
36	RD	13	ARG	2.0
46	RR	8	ARG	2.0
40	YH	125	VAL	2.0
40	YH	41	MET	2.0
54	YZ	150	LEU	2.0
17	XQ	86	GLU	2.0
25	R1	6	GLU	2.0
1	XA	1369	C	2.0
6	XF	51	PRO	2.0
6	XF	96	PRO	2.0
25	R1	77	ALA	2.0
36	RD	232	PRO	2.0
53	RY	65	ALA	2.0
16	XP	17	TYR	2.0
2	QB	174	VAL	2.0
27	Y3	23	LEU	2.0
46	YR	52	ILE	2.0
52	YX	70	LEU	2.0
42	RN	1	MET	2.0
52	YX	89	ILE	2.0
13	XM	99	ARG	2.0
17	XQ	68	ARG	2.0
36	YD	153	ALA	2.0
45	RQ	15	GLY	2.0
54	YZ	80	ARG	2.0
1	XA	1290	G	2.0
12	XL	8	ASN	2.0
8	XH	95	VAL	2.0
14	QN	47	LEU	2.0
17	XQ	31	LEU	2.0
25	R1	47	GLN	2.0
36	RD	46	GLN	2.0
36	YD	155	LEU	2.0
48	YT	30	VAL	2.0
50	RV	40	LEU	2.0
51	YW	85	VAL	2.0
46	YR	9	LYS	2.0
12	QL	129	ALA	2.0

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Mol	Chain	Res	Type	RSRZ
34	RA	2798	C	2.0
4	QD	7	PRO	2.0
12	QL	31	PRO	2.0

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

There are no non-standard protein/DNA/RNA residues in this entry.

6.3 Carbohydrates [i](#)

There are no monosaccharides in this entry.

6.4 Ligands [i](#)

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

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(Å ²)	Q<0.9
55	MG	YA	3277	1/1	-0.39	0.38	213,213,213,213	0
55	MG	QA	1634	1/1	-0.27	0.29	175,175,175,175	0
55	MG	YA	3192	1/1	-0.25	0.25	219,219,219,219	0
55	MG	RA	3139	1/1	-0.24	0.42	290,290,290,290	0
55	MG	RA	3164	1/1	-0.24	0.31	199,199,199,199	0
55	MG	YA	3145	1/1	-0.17	0.29	84,84,84,84	0
55	MG	RA	3186	1/1	-0.16	0.20	205,205,205,205	0
55	MG	YA	3142	1/1	-0.14	0.38	158,158,158,158	0
55	MG	RA	3011	1/1	-0.14	0.22	129,129,129,129	0
55	MG	QE	201	1/1	-0.12	0.43	198,198,198,198	0
55	MG	QH	201	1/1	-0.11	0.17	188,188,188,188	0
55	MG	RA	3163	1/1	-0.07	0.22	165,165,165,165	0
55	MG	RA	3169	1/1	-0.06	0.38	131,131,131,131	0
55	MG	QA	1654	1/1	-0.05	0.31	189,189,189,189	0
55	MG	RA	3224	1/1	-0.04	0.24	181,181,181,181	0
55	MG	RA	3075	1/1	-0.01	0.49	147,147,147,147	0
55	MG	RA	3272	1/1	0.02	0.26	188,188,188,188	0
55	MG	YA	3091	1/1	0.02	0.29	145,145,145,145	0
55	MG	RA	2907	1/1	0.05	0.23	158,158,158,158	0
55	MG	YA	3202	1/1	0.07	0.39	164,164,164,164	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	RA	3211	1/1	0.07	0.27	145,145,145,145	0
55	MG	YA	3254	1/1	0.08	0.22	181,181,181,181	0
55	MG	RA	3217	1/1	0.09	0.13	205,205,205,205	0
55	MG	RA	2921	1/1	0.11	0.42	136,136,136,136	0
55	MG	XA	1675	1/1	0.12	0.16	137,137,137,137	0
55	MG	RA	3078	1/1	0.13	0.43	208,208,208,208	0
55	MG	QA	1615	1/1	0.13	0.28	139,139,139,139	0
55	MG	RA	3326	1/1	0.13	0.21	165,165,165,165	0
55	MG	XA	1651	1/1	0.13	0.25	162,162,162,162	0
55	MG	YA	3250	1/1	0.14	0.34	148,148,148,148	0
55	MG	RA	3013	1/1	0.14	0.42	143,143,143,143	0
55	MG	RA	3223	1/1	0.14	0.41	122,122,122,122	0
55	MG	YA	3126	1/1	0.15	0.52	84,84,84,84	0
55	MG	XA	1647	1/1	0.15	0.14	199,199,199,199	0
55	MG	RA	3008	1/1	0.16	0.29	150,150,150,150	0
55	MG	RA	3172	1/1	0.17	0.24	107,107,107,107	0
55	MG	RA	3275	1/1	0.17	0.26	156,156,156,156	0
55	MG	RA	3138	1/1	0.17	0.19	185,185,185,185	0
55	MG	RA	3206	1/1	0.18	0.28	101,101,101,101	0
55	MG	RA	3030	1/1	0.18	0.22	157,157,157,157	0
55	MG	YA	2962	1/1	0.20	0.41	124,124,124,124	0
55	MG	QA	1627	1/1	0.20	0.20	173,173,173,173	0
55	MG	YA	3234	1/1	0.21	0.31	123,123,123,123	0
55	MG	YA	3176	1/1	0.21	0.21	111,111,111,111	0
55	MG	XA	1620	1/1	0.22	0.23	138,138,138,138	0
55	MG	RA	3035	1/1	0.22	0.24	113,113,113,113	0
55	MG	RA	3179	1/1	0.23	0.31	121,121,121,121	0
55	MG	QA	1605	1/1	0.23	0.31	188,188,188,188	0
55	MG	QA	1647	1/1	0.24	0.23	225,225,225,225	0
55	MG	RA	2909	1/1	0.24	0.30	186,186,186,186	0
55	MG	RA	3023	1/1	0.24	0.36	98,98,98,98	0
55	MG	YA	2965	1/1	0.25	0.35	94,94,94,94	0
55	MG	RA	2937	1/1	0.26	0.29	119,119,119,119	0
55	MG	YA	3093	1/1	0.26	0.26	127,127,127,127	0
55	MG	XA	1680	1/1	0.27	0.36	144,144,144,144	0
55	MG	QA	1624	1/1	0.27	0.23	92,92,92,92	0
55	MG	RA	3005	1/1	0.27	0.31	174,174,174,174	0
55	MG	RA	3122	1/1	0.28	0.20	97,97,97,97	0
55	MG	RA	3286	1/1	0.28	0.51	126,126,126,126	0
55	MG	YA	3262	1/1	0.28	0.44	150,150,150,150	0
55	MG	YA	3015	1/1	0.28	0.25	123,123,123,123	0
55	MG	RA	2989	1/1	0.29	0.27	136,136,136,136	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	RA	3193	1/1	0.29	0.21	141,141,141,141	0
55	MG	RA	3018	1/1	0.29	0.12	136,136,136,136	0
55	MG	YA	3175	1/1	0.29	0.49	117,117,117,117	0
55	MG	XA	1679	1/1	0.29	0.24	124,124,124,124	0
55	MG	QA	1664	1/1	0.29	0.14	156,156,156,156	0
55	MG	YA	3213	1/1	0.30	0.20	169,169,169,169	0
55	MG	YA	3117	1/1	0.30	0.83	120,120,120,120	0
55	MG	XA	1676	1/1	0.31	0.34	143,143,143,143	0
55	MG	RA	2935	1/1	0.31	0.24	132,132,132,132	0
55	MG	YA	3281	1/1	0.31	0.45	151,151,151,151	0
55	MG	QA	1602	1/1	0.32	0.20	148,148,148,148	0
55	MG	RA	3294	1/1	0.32	0.14	143,143,143,143	0
55	MG	XA	1657	1/1	0.32	0.47	236,236,236,236	0
55	MG	YA	3245	1/1	0.33	0.52	84,84,84,84	0
55	MG	YA	2980	1/1	0.33	0.23	84,84,84,84	0
55	MG	YA	2956	1/1	0.33	0.32	114,114,114,114	0
55	MG	YA	3290	1/1	0.33	0.41	211,211,211,211	0
55	MG	XA	1650	1/1	0.34	0.19	189,189,189,189	0
55	MG	XA	1660	1/1	0.34	0.32	114,114,114,114	0
55	MG	RA	2963	1/1	0.34	0.20	256,256,256,256	0
55	MG	YA	2904	1/1	0.34	0.50	122,122,122,122	0
55	MG	YA	3164	1/1	0.35	0.36	84,84,84,84	0
55	MG	YA	3137	1/1	0.35	0.31	110,110,110,110	0
55	MG	QA	1662	1/1	0.35	0.30	151,151,151,151	0
55	MG	YA	3178	1/1	0.35	0.40	84,84,84,84	0
55	MG	RA	2938	1/1	0.35	0.37	159,159,159,159	0
55	MG	YA	2954	1/1	0.36	0.40	135,135,135,135	0
55	MG	RA	3083	1/1	0.36	0.69	150,150,150,150	0
55	MG	RA	2906	1/1	0.36	0.39	150,150,150,150	0
55	MG	YA	2905	1/1	0.36	0.57	175,175,175,175	0
55	MG	RA	3173	1/1	0.37	0.26	130,130,130,130	0
55	MG	RA	3140	1/1	0.37	0.38	88,88,88,88	0
55	MG	XA	1672	1/1	0.37	0.24	160,160,160,160	0
55	MG	YA	2948	1/1	0.37	0.33	84,84,84,84	0
55	MG	YA	3162	1/1	0.38	0.13	112,112,112,112	0
55	MG	RA	3127	1/1	0.38	0.15	132,132,132,132	0
55	MG	QA	1613	1/1	0.39	0.35	127,127,127,127	0
55	MG	YA	2916	1/1	0.39	0.37	121,121,121,121	0
55	MG	RA	3161	1/1	0.39	0.29	116,116,116,116	0
55	MG	YA	3120	1/1	0.39	0.40	84,84,84,84	0
55	MG	YA	3127	1/1	0.40	0.24	84,84,84,84	0
55	MG	YA	3233	1/1	0.40	0.17	153,153,153,153	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(Å ²)	Q<0.9
55	MG	RA	3124	1/1	0.40	0.14	115,115,115,115	0
55	MG	YA	3141	1/1	0.40	0.32	102,102,102,102	0
55	MG	RA	3332	1/1	0.40	0.40	120,120,120,120	0
55	MG	XA	1613	1/1	0.41	0.49	171,171,171,171	0
55	MG	YA	2972	1/1	0.41	0.31	100,100,100,100	0
55	MG	YA	3279	1/1	0.41	0.21	122,122,122,122	0
55	MG	YA	3039	1/1	0.41	0.11	123,123,123,123	0
55	MG	YA	3179	1/1	0.41	0.32	84,84,84,84	0
55	MG	YB	201	1/1	0.41	0.35	104,104,104,104	0
55	MG	RA	3029	1/1	0.42	0.34	100,100,100,100	0
55	MG	YA	3194	1/1	0.42	0.52	163,163,163,163	0
55	MG	YA	3148	1/1	0.42	0.18	102,102,102,102	0
55	MG	RA	3238	1/1	0.42	0.29	157,157,157,157	0
55	MG	YA	3060	1/1	0.42	0.19	177,177,177,177	0
55	MG	RA	3117	1/1	0.43	0.12	243,243,243,243	0
55	MG	YA	3068	1/1	0.43	0.20	158,158,158,158	0
55	MG	RA	2923	1/1	0.43	0.27	174,174,174,174	0
55	MG	RA	3302	1/1	0.43	0.20	116,116,116,116	0
55	MG	RA	2962	1/1	0.43	0.27	124,124,124,124	0
55	MG	RA	3028	1/1	0.43	0.23	96,96,96,96	0
55	MG	XA	1601	1/1	0.43	0.25	174,174,174,174	0
55	MG	YA	3171	1/1	0.43	0.28	84,84,84,84	0
55	MG	YA	3059	1/1	0.43	0.15	189,189,189,189	0
55	MG	RA	3170	1/1	0.44	0.17	132,132,132,132	0
55	MG	XA	1628	1/1	0.44	0.20	109,109,109,109	0
55	MG	QA	1628	1/1	0.45	0.12	238,238,238,238	0
55	MG	RA	3314	1/1	0.45	0.39	137,137,137,137	0
55	MG	RA	2943	1/1	0.46	0.37	132,132,132,132	0
55	MG	YA	3003	1/1	0.46	0.28	97,97,97,97	0
55	MG	XA	1649	1/1	0.46	0.16	123,123,123,123	0
55	MG	QA	1636	1/1	0.46	0.26	134,134,134,134	0
55	MG	XA	1634	1/1	0.46	0.27	107,107,107,107	0
55	MG	YA	2951	1/1	0.47	0.35	101,101,101,101	0
55	MG	YA	3109	1/1	0.47	0.27	95,95,95,95	0
55	MG	YA	3186	1/1	0.47	0.23	187,187,187,187	0
55	MG	QA	1655	1/1	0.47	0.16	121,121,121,121	0
55	MG	XA	1612	1/1	0.47	0.51	191,191,191,191	0
55	MG	RA	3106	1/1	0.47	0.23	90,90,90,90	0
55	MG	RA	3074	1/1	0.47	0.16	149,149,149,149	0
55	MG	RA	3213	1/1	0.47	0.13	157,157,157,157	0
55	MG	XA	1678	1/1	0.47	0.15	173,173,173,173	0
55	MG	RA	3137	1/1	0.48	0.10	177,177,177,177	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	RA	2930	1/1	0.48	0.33	136,136,136,136	0
55	MG	RA	2947	1/1	0.48	0.24	118,118,118,118	0
55	MG	RA	2929	1/1	0.48	0.20	211,211,211,211	0
55	MG	YA	3199	1/1	0.48	0.38	186,186,186,186	0
55	MG	YA	3140	1/1	0.48	0.16	84,84,84,84	0
55	MG	XA	1630	1/1	0.49	0.24	180,180,180,180	0
55	MG	RA	3132	1/1	0.49	0.35	238,238,238,238	0
55	MG	YA	3205	1/1	0.49	0.41	152,152,152,152	0
55	MG	RA	3092	1/1	0.50	0.28	98,98,98,98	0
55	MG	RA	3104	1/1	0.50	0.37	84,84,84,84	0
55	MG	RA	3293	1/1	0.50	0.13	121,121,121,121	0
55	MG	YA	3131	1/1	0.50	0.20	141,141,141,141	0
55	MG	QA	1643	1/1	0.50	0.24	180,180,180,180	0
55	MG	RA	3052	1/1	0.50	0.23	84,84,84,84	0
55	MG	RA	3034	1/1	0.50	0.19	111,111,111,111	0
55	MG	YU	201	1/1	0.50	0.14	97,97,97,97	0
55	MG	QA	1614	1/1	0.51	0.34	120,120,120,120	0
55	MG	YA	3267	1/1	0.51	0.15	84,84,84,84	0
55	MG	YA	3097	1/1	0.51	0.68	135,135,135,135	0
55	MG	XA	1606	1/1	0.51	0.14	165,165,165,165	0
55	MG	YA	3090	1/1	0.52	0.22	84,84,84,84	0
55	MG	YA	3115	1/1	0.52	0.30	148,148,148,148	0
55	MG	RA	3026	1/1	0.52	0.36	84,84,84,84	0
55	MG	RA	3059	1/1	0.52	0.37	153,153,153,153	0
55	MG	QA	1604	1/1	0.52	0.29	153,153,153,153	0
55	MG	YA	3258	1/1	0.53	0.35	88,88,88,88	0
55	MG	RA	3324	1/1	0.53	0.32	120,120,120,120	0
55	MG	YA	2940	1/1	0.53	0.22	84,84,84,84	0
55	MG	XA	1665	1/1	0.53	0.44	123,123,123,123	0
55	MG	QA	1637	1/1	0.53	0.26	126,126,126,126	0
55	MG	YA	2989	1/1	0.53	0.15	120,120,120,120	0
55	MG	XA	1618	1/1	0.53	0.28	103,103,103,103	0
55	MG	YA	3011	1/1	0.53	0.30	84,84,84,84	0
55	MG	RA	3328	1/1	0.53	0.25	145,145,145,145	0
55	MG	YA	3259	1/1	0.54	0.17	84,84,84,84	0
55	MG	XA	1652	1/1	0.54	0.09	143,143,143,143	0
55	MG	RA	3129	1/1	0.54	0.28	126,126,126,126	0
55	MG	RA	3086	1/1	0.54	0.36	84,84,84,84	0
55	MG	RA	3320	1/1	0.54	0.26	84,84,84,84	0
55	MG	QA	1625	1/1	0.54	0.22	104,104,104,104	0
55	MG	YA	2955	1/1	0.54	0.50	110,110,110,110	0
55	MG	YA	3256	1/1	0.54	0.23	94,94,94,94	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	RA	3101	1/1	0.54	0.71	120,120,120,120	0
55	MG	QA	1670	1/1	0.55	0.19	190,190,190,190	0
55	MG	RA	2953	1/1	0.55	0.32	101,101,101,101	0
55	MG	RA	3226	1/1	0.55	0.30	84,84,84,84	0
55	MG	XA	1688	1/1	0.55	0.15	122,122,122,122	0
55	MG	XA	1632	1/1	0.55	0.19	104,104,104,104	0
55	MG	RA	3039	1/1	0.55	0.16	256,256,256,256	0
55	MG	RA	3118	1/1	0.55	0.11	186,186,186,186	0
55	MG	RA	3162	1/1	0.55	0.57	139,139,139,139	0
55	MG	RA	2916	1/1	0.55	0.27	97,97,97,97	0
55	MG	YA	3048	1/1	0.56	0.38	84,84,84,84	0
55	MG	QA	1663	1/1	0.56	0.29	128,128,128,128	0
55	MG	RA	3095	1/1	0.56	0.13	258,258,258,258	0
55	MG	RA	3190	1/1	0.56	0.18	143,143,143,143	0
55	MG	RA	3057	1/1	0.56	0.72	180,180,180,180	0
55	MG	RA	3242	1/1	0.56	0.24	101,101,101,101	0
55	MG	YA	3211	1/1	0.56	0.44	101,101,101,101	0
55	MG	RA	3221	1/1	0.56	0.52	182,182,182,182	0
57	ZN	QN	101	1/1	0.56	0.17	439,439,439,439	0
55	MG	RA	3281	1/1	0.57	0.28	105,105,105,105	0
55	MG	YA	3050	1/1	0.57	0.29	84,84,84,84	0
55	MG	YA	3004	1/1	0.57	0.48	84,84,84,84	0
55	MG	RA	2913	1/1	0.57	0.15	107,107,107,107	0
55	MG	XA	1621	1/1	0.57	0.52	128,128,128,128	0
55	MG	RA	2969	1/1	0.57	0.25	104,104,104,104	0
55	MG	RA	3174	1/1	0.58	0.22	142,142,142,142	0
55	MG	RA	3208	1/1	0.58	0.29	84,84,84,84	0
55	MG	RA	2992	1/1	0.58	0.15	123,123,123,123	0
55	MG	RA	3180	1/1	0.58	0.20	168,168,168,168	0
55	MG	RA	3183	1/1	0.58	0.33	201,201,201,201	0
55	MG	RA	3220	1/1	0.58	0.78	166,166,166,166	0
55	MG	RA	3002	1/1	0.58	0.27	132,132,132,132	0
55	MG	YA	3242	1/1	0.58	0.14	84,84,84,84	0
55	MG	RA	3073	1/1	0.58	0.22	89,89,89,89	0
55	MG	YA	3049	1/1	0.58	0.28	84,84,84,84	0
55	MG	YA	3253	1/1	0.58	0.36	84,84,84,84	0
55	MG	RA	3144	1/1	0.58	0.48	126,126,126,126	0
55	MG	RA	3308	1/1	0.59	0.24	161,161,161,161	0
55	MG	RA	3289	1/1	0.59	0.30	125,125,125,125	0
55	MG	YA	2982	1/1	0.59	0.33	84,84,84,84	0
55	MG	RA	3093	1/1	0.59	0.17	131,131,131,131	0
55	MG	YA	3183	1/1	0.59	0.14	89,89,89,89	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	XA	1642	1/1	0.59	0.37	128,128,128,128	0
55	MG	RA	2912	1/1	0.59	0.50	132,132,132,132	0
55	MG	RA	3136	1/1	0.59	0.37	84,84,84,84	0
55	MG	YA	3057	1/1	0.60	0.29	105,105,105,105	0
55	MG	RA	3088	1/1	0.60	0.29	101,101,101,101	0
55	MG	RE	304	1/1	0.60	0.53	96,96,96,96	0
55	MG	R0	101	1/1	0.60	0.26	147,147,147,147	0
55	MG	YA	3077	1/1	0.60	0.22	87,87,87,87	0
55	MG	RA	2941	1/1	0.60	0.39	167,167,167,167	0
55	MG	YA	3044	1/1	0.60	0.31	84,84,84,84	0
55	MG	RA	3198	1/1	0.60	0.17	144,144,144,144	0
55	MG	YA	3239	1/1	0.60	0.41	84,84,84,84	0
55	MG	YA	3288	1/1	0.60	0.46	84,84,84,84	0
55	MG	XA	1674	1/1	0.60	0.16	159,159,159,159	0
55	MG	RA	2924	1/1	0.60	0.32	98,98,98,98	0
55	MG	YA	3187	1/1	0.60	0.26	117,117,117,117	0
55	MG	YA	3114	1/1	0.60	0.29	116,116,116,116	0
55	MG	RA	3076	1/1	0.61	0.26	114,114,114,114	0
55	MG	YA	2974	1/1	0.61	0.42	84,84,84,84	0
55	MG	YA	3032	1/1	0.61	0.21	84,84,84,84	0
55	MG	RA	3065	1/1	0.61	0.24	105,105,105,105	0
55	MG	RA	3252	1/1	0.61	0.30	121,121,121,121	0
55	MG	RA	3054	1/1	0.61	0.37	84,84,84,84	0
55	MG	XA	1669	1/1	0.61	0.08	187,187,187,187	0
55	MG	RA	3134	1/1	0.61	0.26	117,117,117,117	0
55	MG	YA	3128	1/1	0.61	0.38	134,134,134,134	0
55	MG	YA	3276	1/1	0.62	0.51	115,115,115,115	0
55	MG	RA	2903	1/1	0.62	0.14	100,100,100,100	0
55	MG	YA	3069	1/1	0.62	0.26	176,176,176,176	0
55	MG	RA	3167	1/1	0.62	0.28	165,165,165,165	0
55	MG	RA	2931	1/1	0.62	0.18	125,125,125,125	0
55	MG	YA	3174	1/1	0.62	0.25	94,94,94,94	0
55	MG	XA	1607	1/1	0.62	0.24	108,108,108,108	0
55	MG	YA	2961	1/1	0.62	0.27	117,117,117,117	0
55	MG	YA	3061	1/1	0.62	0.14	146,146,146,146	0
55	MG	QA	1618	1/1	0.63	0.38	176,176,176,176	0
55	MG	XA	1682	1/1	0.63	0.19	95,95,95,95	0
55	MG	RA	3304	1/1	0.63	0.28	155,155,155,155	0
55	MG	YA	3135	1/1	0.63	0.31	84,84,84,84	0
55	MG	YA	3232	1/1	0.63	0.25	91,91,91,91	0
55	MG	YA	2984	1/1	0.63	0.52	84,84,84,84	0
55	MG	RA	2965	1/1	0.63	0.20	211,211,211,211	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	YA	3237	1/1	0.63	0.42	115,115,115,115	0
55	MG	RA	2942	1/1	0.63	0.24	109,109,109,109	0
55	MG	YA	3287	1/1	0.63	0.36	84,84,84,84	0
55	MG	RA	3064	1/1	0.63	0.47	110,110,110,110	0
55	MG	YA	3010	1/1	0.63	0.41	84,84,84,84	0
55	MG	YA	2964	1/1	0.63	0.36	115,115,115,115	0
55	MG	XA	1646	1/1	0.63	0.19	148,148,148,148	0
55	MG	QA	1629	1/1	0.63	0.24	142,142,142,142	0
55	MG	YA	2981	1/1	0.64	0.31	103,103,103,103	0
55	MG	YA	3155	1/1	0.64	0.36	99,99,99,99	0
55	MG	XA	1670	1/1	0.64	0.35	136,136,136,136	0
55	MG	YA	2952	1/1	0.64	0.41	101,101,101,101	0
55	MG	YA	3122	1/1	0.64	0.42	84,84,84,84	0
55	MG	XA	1619	1/1	0.64	0.33	111,111,111,111	0
55	MG	YA	3017	1/1	0.64	0.45	107,107,107,107	0
55	MG	YA	3144	1/1	0.64	0.32	93,93,93,93	0
55	MG	QA	1619	1/1	0.64	0.28	123,123,123,123	0
55	MG	Y8	101	1/1	0.65	0.59	197,197,197,197	0
55	MG	RA	2910	1/1	0.65	0.18	149,149,149,149	0
55	MG	RA	3123	1/1	0.65	0.28	115,115,115,115	0
55	MG	RA	2952	1/1	0.65	0.16	143,143,143,143	0
55	MG	YA	3024	1/1	0.65	0.24	84,84,84,84	0
55	MG	XA	1616	1/1	0.65	0.34	291,291,291,291	0
55	MG	RA	3103	1/1	0.65	0.41	84,84,84,84	0
55	MG	QA	1669	1/1	0.65	0.26	158,158,158,158	0
55	MG	RA	3147	1/1	0.65	0.51	104,104,104,104	0
55	MG	YA	3107	1/1	0.65	0.32	204,204,204,204	0
55	MG	QA	1645	1/1	0.65	0.18	101,101,101,101	0
55	MG	RA	3295	1/1	0.65	0.28	129,129,129,129	0
55	MG	RA	2905	1/1	0.65	0.29	184,184,184,184	0
55	MG	RA	3020	1/1	0.65	0.33	126,126,126,126	0
55	MG	YA	3196	1/1	0.65	0.20	84,84,84,84	0
55	MG	YA	3151	1/1	0.66	0.30	94,94,94,94	0
55	MG	RA	3143	1/1	0.66	0.19	109,109,109,109	0
55	MG	YA	2935	1/1	0.66	0.39	84,84,84,84	0
55	MG	RA	3260	1/1	0.66	0.41	104,104,104,104	0
55	MG	QA	1632	1/1	0.66	0.11	84,84,84,84	0
55	MG	R0	102	1/1	0.66	0.11	120,120,120,120	0
55	MG	YA	3273	1/1	0.66	0.24	102,102,102,102	0
55	MG	YA	3218	1/1	0.66	0.32	84,84,84,84	0
55	MG	RA	3313	1/1	0.66	0.24	203,203,203,203	0
55	MG	YA	3035	1/1	0.66	0.28	122,122,122,122	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	RA	3051	1/1	0.66	0.46	170,170,170,170	0
55	MG	YA	3282	1/1	0.66	0.15	116,116,116,116	0
55	MG	QA	1607	1/1	0.66	0.36	84,84,84,84	0
55	MG	QA	1617	1/1	0.66	0.36	159,159,159,159	0
55	MG	YA	2995	1/1	0.66	0.28	116,116,116,116	0
55	MG	QA	1609	1/1	0.66	0.10	120,120,120,120	0
55	MG	RA	3111	1/1	0.66	0.34	102,102,102,102	0
55	MG	YA	3005	1/1	0.66	0.32	84,84,84,84	0
55	MG	RA	2939	1/1	0.67	0.14	84,84,84,84	0
55	MG	RA	3071	1/1	0.67	0.64	288,288,288,288	0
55	MG	YA	3180	1/1	0.67	0.29	90,90,90,90	0
55	MG	YA	3182	1/1	0.67	0.15	84,84,84,84	0
55	MG	XA	1633	1/1	0.67	0.26	99,99,99,99	0
55	MG	YA	3055	1/1	0.67	0.12	92,92,92,92	0
55	MG	RA	3048	1/1	0.67	0.25	168,168,168,168	0
55	MG	YA	3018	1/1	0.67	0.91	120,120,120,120	0
55	MG	YA	3224	1/1	0.67	0.34	84,84,84,84	0
55	MG	YA	2933	1/1	0.68	0.45	97,97,97,97	0
55	MG	RA	3210	1/1	0.68	0.18	121,121,121,121	0
55	MG	RA	3094	1/1	0.68	0.10	106,106,106,106	0
55	MG	YA	3085	1/1	0.68	0.30	84,84,84,84	0
55	MG	YA	3086	1/1	0.68	0.31	84,84,84,84	0
55	MG	RA	3171	1/1	0.68	0.40	115,115,115,115	0
55	MG	RA	2986	1/1	0.68	0.33	84,84,84,84	0
55	MG	RA	2920	1/1	0.68	0.17	113,113,113,113	0
55	MG	YA	2953	1/1	0.68	0.44	113,113,113,113	0
55	MG	RA	2957	1/1	0.68	0.33	95,95,95,95	0
55	MG	YA	3209	1/1	0.68	0.18	111,111,111,111	0
55	MG	QA	1601	1/1	0.68	0.26	123,123,123,123	0
55	MG	RA	3079	1/1	0.68	0.32	135,135,135,135	0
55	MG	YA	2959	1/1	0.68	0.32	84,84,84,84	0
55	MG	RA	3053	1/1	0.69	0.34	122,122,122,122	0
55	MG	YA	3095	1/1	0.69	0.15	84,84,84,84	0
55	MG	YA	3268	1/1	0.69	0.27	89,89,89,89	0
55	MG	QA	1626	1/1	0.69	0.18	104,104,104,104	0
55	MG	YA	2957	1/1	0.69	0.19	84,84,84,84	0
55	MG	RA	3222	1/1	0.69	0.39	166,166,166,166	0
55	MG	RA	2918	1/1	0.69	0.27	176,176,176,176	0
55	MG	XA	1658	1/1	0.69	0.35	179,179,179,179	0
55	MG	YA	3247	1/1	0.69	0.20	87,87,87,87	0
55	MG	RA	3182	1/1	0.69	0.12	150,150,150,150	0
55	MG	RA	3214	1/1	0.69	0.30	157,157,157,157	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	QA	1638	1/1	0.69	0.27	84,84,84,84	0
55	MG	RA	3240	1/1	0.69	0.21	89,89,89,89	0
55	MG	XA	1623	1/1	0.69	0.19	127,127,127,127	0
55	MG	YA	3228	1/1	0.69	0.28	115,115,115,115	0
55	MG	RA	3102	1/1	0.70	0.29	102,102,102,102	0
55	MG	XA	1671	1/1	0.70	0.08	149,149,149,149	0
55	MG	YA	2944	1/1	0.70	0.19	106,106,106,106	0
55	MG	QL	201	1/1	0.70	0.19	258,258,258,258	0
55	MG	RA	3157	1/1	0.70	0.20	156,156,156,156	0
55	MG	RA	3327	1/1	0.70	0.32	84,84,84,84	0
55	MG	QA	1621	1/1	0.70	0.32	159,159,159,159	0
55	MG	RA	3031	1/1	0.70	0.15	94,94,94,94	0
55	MG	YR	202	1/1	0.70	0.39	119,119,119,119	0
55	MG	YA	3189	1/1	0.70	0.20	93,93,93,93	0
55	MG	RA	3237	1/1	0.70	0.32	84,84,84,84	0
55	MG	YA	2931	1/1	0.71	0.42	84,84,84,84	0
55	MG	QA	1648	1/1	0.71	0.27	139,139,139,139	0
55	MG	R8	102	1/1	0.71	0.14	109,109,109,109	0
55	MG	YA	2936	1/1	0.71	0.42	84,84,84,84	0
55	MG	QA	1623	1/1	0.71	0.24	84,84,84,84	0
55	MG	YA	3001	1/1	0.71	0.11	84,84,84,84	0
55	MG	XA	1656	1/1	0.71	0.13	98,98,98,98	0
55	MG	YA	2963	1/1	0.71	0.35	84,84,84,84	0
55	MG	QA	1606	1/1	0.71	0.23	116,116,116,116	0
55	MG	YA	3094	1/1	0.71	0.17	104,104,104,104	0
55	MG	RA	3305	1/1	0.71	0.22	134,134,134,134	0
55	MG	QA	1642	1/1	0.71	0.07	157,157,157,157	0
55	MG	YA	3104	1/1	0.71	0.27	105,105,105,105	0
55	MG	YA	2908	1/1	0.71	0.30	84,84,84,84	0
55	MG	RA	3133	1/1	0.71	0.87	169,169,169,169	0
55	MG	YA	3156	1/1	0.72	0.25	84,84,84,84	0
55	MG	YA	3132	1/1	0.72	0.10	124,124,124,124	0
55	MG	YA	3271	1/1	0.72	0.30	92,92,92,92	0
55	MG	YA	3133	1/1	0.72	0.16	84,84,84,84	0
55	MG	YA	3134	1/1	0.72	0.16	120,120,120,120	0
55	MG	YA	3111	1/1	0.72	0.16	119,119,119,119	0
55	MG	RA	2954	1/1	0.72	0.61	306,306,306,306	0
55	MG	QA	1622	1/1	0.72	0.49	169,169,169,169	0
55	MG	RA	2904	1/1	0.72	0.29	219,219,219,219	0
55	MG	RA	3001	1/1	0.72	0.16	95,95,95,95	0
55	MG	YA	2969	1/1	0.72	0.36	84,84,84,84	0
55	MG	RA	3045	1/1	0.72	0.32	169,169,169,169	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	YA	3291	1/1	0.72	0.34	84,84,84,84	0
55	MG	YA	3082	1/1	0.72	0.38	84,84,84,84	0
55	MG	XA	1685	1/1	0.72	0.17	113,113,113,113	0
55	MG	XA	1668	1/1	0.72	0.13	131,131,131,131	0
55	MG	YA	3265	1/1	0.72	0.34	107,107,107,107	0
55	MG	Y5	102	1/1	0.73	0.24	84,84,84,84	0
55	MG	RA	3125	1/1	0.73	0.37	174,174,174,174	0
55	MG	RA	2936	1/1	0.73	0.11	87,87,87,87	0
55	MG	RA	2968	1/1	0.73	0.13	176,176,176,176	0
55	MG	YA	3087	1/1	0.73	0.43	84,84,84,84	0
55	MG	RA	3322	1/1	0.73	0.23	104,104,104,104	0
55	MG	RA	2917	1/1	0.73	0.15	100,100,100,100	0
55	MG	YA	2926	1/1	0.73	0.22	84,84,84,84	0
55	MG	YA	3143	1/1	0.73	0.39	84,84,84,84	0
55	MG	YA	3123	1/1	0.73	0.23	84,84,84,84	0
55	MG	XA	1602	1/1	0.73	0.20	96,96,96,96	0
55	MG	YA	3067	1/1	0.73	0.10	102,102,102,102	0
55	MG	XA	1603	1/1	0.73	0.15	189,189,189,189	0
55	MG	YA	3153	1/1	0.73	0.41	84,84,84,84	0
55	MG	RA	2998	1/1	0.73	0.18	84,84,84,84	0
55	MG	YA	3083	1/1	0.74	0.33	84,84,84,84	0
55	MG	RA	3046	1/1	0.74	0.22	150,150,150,150	0
55	MG	QA	1660	1/1	0.74	0.16	127,127,127,127	0
55	MG	YA	3173	1/1	0.74	0.28	84,84,84,84	0
55	MG	RA	2972	1/1	0.74	0.33	84,84,84,84	0
55	MG	YA	3054	1/1	0.74	0.14	93,93,93,93	0
55	MG	YA	2976	1/1	0.74	0.28	84,84,84,84	0
55	MG	RA	3311	1/1	0.74	0.26	84,84,84,84	0
55	MG	RA	3204	1/1	0.74	0.34	167,167,167,167	0
55	MG	YA	3161	1/1	0.74	0.22	132,132,132,132	0
57	ZN	XN	101	1/1	0.74	0.14	489,489,489,489	0
55	MG	YA	2915	1/1	0.75	0.28	107,107,107,107	0
55	MG	RA	2908	1/1	0.75	0.12	102,102,102,102	0
55	MG	RA	2966	1/1	0.75	0.14	149,149,149,149	0
55	MG	YA	3065	1/1	0.75	0.14	107,107,107,107	0
55	MG	RA	3115	1/1	0.75	0.10	130,130,130,130	0
55	MG	YA	3110	1/1	0.75	0.20	177,177,177,177	0
55	MG	YA	2932	1/1	0.75	0.72	84,84,84,84	0
55	MG	QA	1666	1/1	0.75	0.13	178,178,178,178	0
55	MG	YA	2983	1/1	0.75	0.55	84,84,84,84	0
55	MG	YA	2934	1/1	0.75	0.55	84,84,84,84	0
55	MG	QA	1631	1/1	0.75	0.11	110,110,110,110	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	RA	2964	1/1	0.75	0.17	84,84,84,84	0
55	MG	XA	1664	1/1	0.75	0.31	84,84,84,84	0
55	MG	YA	3191	1/1	0.75	0.27	84,84,84,84	0
55	MG	RA	3330	1/1	0.75	0.22	84,84,84,84	0
55	MG	YA	3051	1/1	0.75	0.32	84,84,84,84	0
55	MG	YA	2907	1/1	0.75	0.47	151,151,151,151	0
55	MG	YA	2968	1/1	0.75	0.32	84,84,84,84	0
55	MG	RA	3200	1/1	0.75	0.24	141,141,141,141	0
55	MG	YA	3257	1/1	0.75	0.19	84,84,84,84	0
55	MG	YA	3203	1/1	0.75	0.28	84,84,84,84	0
55	MG	RA	3152	1/1	0.76	0.46	122,122,122,122	0
55	MG	YA	3210	1/1	0.76	0.41	121,121,121,121	0
55	MG	RA	3280	1/1	0.76	0.13	84,84,84,84	0
55	MG	RA	3156	1/1	0.76	0.20	106,106,106,106	0
55	MG	RA	3285	1/1	0.76	0.17	161,161,161,161	0
55	MG	RA	2975	1/1	0.76	0.50	104,104,104,104	0
55	MG	QA	1658	1/1	0.76	0.10	84,84,84,84	0
55	MG	RA	2926	1/1	0.76	0.31	124,124,124,124	0
55	MG	YA	2912	1/1	0.76	0.30	84,84,84,84	0
55	MG	YA	3027	1/1	0.76	0.12	102,102,102,102	0
55	MG	YA	3075	1/1	0.76	0.39	84,84,84,84	0
55	MG	RA	3112	1/1	0.76	0.23	84,84,84,84	0
55	MG	YA	3286	1/1	0.76	0.32	144,144,144,144	0
55	MG	RA	3066	1/1	0.76	0.22	117,117,117,117	0
55	MG	RA	2928	1/1	0.76	0.18	138,138,138,138	0
55	MG	YA	3154	1/1	0.76	0.14	138,138,138,138	0
55	MG	YA	2930	1/1	0.76	0.41	84,84,84,84	0
55	MG	RA	3056	1/1	0.76	0.30	139,139,139,139	0
55	MG	RA	3146	1/1	0.76	0.09	84,84,84,84	0
55	MG	RA	2950	1/1	0.76	0.09	108,108,108,108	0
55	MG	YA	3002	1/1	0.76	0.28	132,132,132,132	0
55	MG	RA	3310	1/1	0.76	0.55	116,116,116,116	0
55	MG	YA	3195	1/1	0.77	0.16	84,84,84,84	0
55	MG	RA	3307	1/1	0.77	0.41	132,132,132,132	0
55	MG	YA	3158	1/1	0.77	0.31	84,84,84,84	0
55	MG	YA	3159	1/1	0.77	0.29	84,84,84,84	0
55	MG	YA	2945	1/1	0.77	0.18	86,86,86,86	0
55	MG	XA	1641	1/1	0.77	0.17	112,112,112,112	0
55	MG	YA	2922	1/1	0.77	0.22	84,84,84,84	0
55	MG	XE	201	1/1	0.77	0.21	157,157,157,157	0
55	MG	YA	2929	1/1	0.77	0.40	84,84,84,84	0
55	MG	YA	2979	1/1	0.77	0.38	95,95,95,95	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	RA	3273	1/1	0.77	0.30	113,113,113,113	0
55	MG	QF	301	1/1	0.77	0.12	145,145,145,145	0
55	MG	RA	3199	1/1	0.77	0.26	84,84,84,84	0
55	MG	RA	2976	1/1	0.77	0.46	139,139,139,139	0
55	MG	YA	2906	1/1	0.77	0.29	84,84,84,84	0
55	MG	YA	3038	1/1	0.77	0.26	84,84,84,84	0
55	MG	RA	3282	1/1	0.77	0.28	149,149,149,149	0
55	MG	YA	3042	1/1	0.77	0.34	84,84,84,84	0
55	MG	YA	3240	1/1	0.77	0.25	84,84,84,84	0
55	MG	YA	3118	1/1	0.77	0.33	84,84,84,84	0
55	MG	YA	3243	1/1	0.77	0.28	108,108,108,108	0
55	MG	YA	3119	1/1	0.77	0.39	84,84,84,84	0
55	MG	RA	3271	1/1	0.77	0.22	102,102,102,102	0
55	MG	YA	3248	1/1	0.77	0.26	134,134,134,134	0
55	MG	YA	2938	1/1	0.77	0.21	84,84,84,84	0
55	MG	RA	2999	1/1	0.77	0.39	121,121,121,121	0
55	MG	YA	3130	1/1	0.78	0.22	84,84,84,84	0
55	MG	XA	1625	1/1	0.78	0.14	190,190,190,190	0
55	MG	YA	2921	1/1	0.78	0.59	84,84,84,84	0
55	MG	QA	1665	1/1	0.78	0.17	84,84,84,84	0
55	MG	RA	3318	1/1	0.78	0.40	84,84,84,84	0
55	MG	XA	1605	1/1	0.78	0.43	161,161,161,161	0
55	MG	QA	1616	1/1	0.78	0.10	128,128,128,128	0
55	MG	YA	3272	1/1	0.78	0.26	98,98,98,98	0
55	MG	YA	3025	1/1	0.78	0.11	98,98,98,98	0
55	MG	YA	3275	1/1	0.78	0.23	112,112,112,112	0
55	MG	YA	3066	1/1	0.78	0.23	113,113,113,113	0
55	MG	RA	3299	1/1	0.78	0.10	102,102,102,102	0
55	MG	XA	1637	1/1	0.78	0.17	84,84,84,84	0
55	MG	RA	3278	1/1	0.78	0.12	113,113,113,113	0
55	MG	YA	3073	1/1	0.78	0.15	84,84,84,84	0
55	MG	YA	3284	1/1	0.78	0.11	128,128,128,128	0
55	MG	YA	2988	1/1	0.78	0.32	84,84,84,84	0
55	MG	RA	3241	1/1	0.78	0.28	84,84,84,84	0
55	MG	RA	3177	1/1	0.78	0.15	156,156,156,156	0
55	MG	RA	2970	1/1	0.78	0.52	102,102,102,102	0
55	MG	RA	3256	1/1	0.78	0.10	84,84,84,84	0
55	MG	RA	3205	1/1	0.78	0.27	100,100,100,100	0
55	MG	YE	302	1/1	0.78	0.26	131,131,131,131	0
55	MG	RA	3038	1/1	0.78	0.20	119,119,119,119	0
55	MG	YA	3088	1/1	0.78	0.25	84,84,84,84	0
55	MG	RA	3218	1/1	0.78	0.17	196,196,196,196	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	YA	3129	1/1	0.78	0.29	84,84,84,84	0
55	MG	YA	3043	1/1	0.79	0.29	84,84,84,84	0
55	MG	RA	3036	1/1	0.79	0.07	136,136,136,136	0
55	MG	QA	1651	1/1	0.79	0.26	116,116,116,116	0
55	MG	YA	3212	1/1	0.79	0.29	84,84,84,84	0
55	MG	RA	3251	1/1	0.79	0.16	177,177,177,177	0
55	MG	RA	3022	1/1	0.79	0.30	84,84,84,84	0
55	MG	YA	3188	1/1	0.79	0.14	124,124,124,124	0
55	MG	QA	1641	1/1	0.79	0.18	155,155,155,155	0
55	MG	YA	3096	1/1	0.79	0.26	92,92,92,92	0
55	MG	YA	3168	1/1	0.79	0.50	84,84,84,84	0
55	MG	YA	3053	1/1	0.79	0.22	93,93,93,93	0
55	MG	YA	3292	1/1	0.79	0.12	109,109,109,109	0
55	MG	RA	3225	1/1	0.79	0.32	93,93,93,93	0
55	MG	YD	302	1/1	0.79	0.36	99,99,99,99	0
55	MG	RA	3202	1/1	0.79	0.18	134,134,134,134	0
55	MG	RA	2933	1/1	0.79	0.16	194,194,194,194	0
55	MG	YA	3006	1/1	0.79	0.34	84,84,84,84	0
55	MG	QA	1649	1/1	0.79	0.25	109,109,109,109	0
55	MG	RA	3176	1/1	0.79	0.15	152,152,152,152	0
55	MG	YA	3150	1/1	0.80	0.68	84,84,84,84	0
55	MG	YA	2911	1/1	0.80	0.24	84,84,84,84	0
55	MG	YA	2975	1/1	0.80	0.13	97,97,97,97	0
55	MG	RA	3166	1/1	0.80	0.23	124,124,124,124	0
55	MG	YA	3026	1/1	0.80	0.28	84,84,84,84	0
55	MG	YA	3263	1/1	0.80	0.56	88,88,88,88	0
55	MG	YA	2950	1/1	0.80	0.17	111,111,111,111	0
55	MG	RA	3243	1/1	0.80	0.17	84,84,84,84	0
55	MG	RA	3017	1/1	0.80	0.18	84,84,84,84	0
55	MG	RA	2959	1/1	0.80	0.30	84,84,84,84	0
55	MG	RA	3037	1/1	0.80	0.09	145,145,145,145	0
55	MG	RA	2987	1/1	0.80	0.23	84,84,84,84	0
55	MG	YA	3167	1/1	0.80	0.19	84,84,84,84	0
55	MG	QA	1603	1/1	0.80	0.10	121,121,121,121	0
55	MG	RA	3116	1/1	0.80	0.18	89,89,89,89	0
55	MG	YA	2958	1/1	0.80	0.40	84,84,84,84	0
55	MG	YA	3231	1/1	0.80	0.19	84,84,84,84	0
55	MG	RA	3297	1/1	0.80	0.21	167,167,167,167	0
55	MG	XA	1610	1/1	0.80	0.57	237,237,237,237	0
55	MG	RA	3195	1/1	0.80	0.53	170,170,170,170	0
55	MG	YA	2903	1/1	0.80	0.18	90,90,90,90	0
55	MG	RA	3141	1/1	0.80	0.26	84,84,84,84	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	YA	3136	1/1	0.80	0.24	84,84,84,84	0
55	MG	XA	1614	1/1	0.80	0.07	106,106,106,106	0
55	MG	YA	2967	1/1	0.80	0.08	84,84,84,84	0
55	MG	YA	3101	1/1	0.80	0.69	139,139,139,139	0
55	MG	XA	1644	1/1	0.80	0.32	235,235,235,235	0
55	MG	YA	3013	1/1	0.80	0.24	84,84,84,84	0
55	MG	RA	2982	1/1	0.80	0.15	87,87,87,87	0
55	MG	YA	3062	1/1	0.80	0.20	119,119,119,119	0
55	MG	RA	3025	1/1	0.80	0.12	84,84,84,84	0
55	MG	YA	3255	1/1	0.80	0.24	84,84,84,84	0
55	MG	QA	1610	1/1	0.81	1.26	211,211,211,211	0
55	MG	YA	3270	1/1	0.81	0.26	84,84,84,84	0
55	MG	YA	3030	1/1	0.81	0.26	110,110,110,110	0
55	MG	YA	3031	1/1	0.81	0.49	84,84,84,84	0
55	MG	YA	3236	1/1	0.81	0.27	84,84,84,84	0
55	MG	YA	2999	1/1	0.81	0.19	84,84,84,84	0
55	MG	RA	3175	1/1	0.81	0.13	84,84,84,84	0
55	MG	XA	1686	1/1	0.81	0.22	108,108,108,108	0
55	MG	RA	2911	1/1	0.81	0.17	111,111,111,111	0
55	MG	RA	3207	1/1	0.81	0.13	151,151,151,151	0
55	MG	YA	3072	1/1	0.81	0.32	106,106,106,106	0
55	MG	YA	3197	1/1	0.81	0.13	84,84,84,84	0
55	MG	YA	3165	1/1	0.81	0.23	84,84,84,84	0
55	MG	Y2	101	1/1	0.81	0.20	133,133,133,133	0
55	MG	RA	3194	1/1	0.81	0.13	111,111,111,111	0
55	MG	RA	3107	1/1	0.81	0.25	94,94,94,94	0
55	MG	RA	3265	1/1	0.81	0.34	84,84,84,84	0
55	MG	RA	2995	1/1	0.81	0.29	260,260,260,260	0
55	MG	RA	2974	1/1	0.81	0.38	84,84,84,84	0
55	MG	RA	3010	1/1	0.81	0.13	138,138,138,138	0
55	MG	XA	1654	1/1	0.81	0.13	118,118,118,118	0
55	MG	YP	201	1/1	0.81	0.16	113,113,113,113	0
55	MG	QA	1620	1/1	0.81	0.21	159,159,159,159	0
55	MG	YA	3149	1/1	0.81	0.16	93,93,93,93	0
55	MG	RA	3276	1/1	0.81	0.14	138,138,138,138	0
55	MG	RA	3203	1/1	0.81	0.14	193,193,193,193	0
55	MG	YA	2985	1/1	0.82	0.48	84,84,84,84	0
55	MG	XA	1673	1/1	0.82	0.13	114,114,114,114	0
55	MG	YA	2943	1/1	0.82	0.18	93,93,93,93	0
55	MG	YA	3274	1/1	0.82	0.28	84,84,84,84	0
55	MG	YA	2919	1/1	0.82	0.36	84,84,84,84	0
55	MG	YA	2997	1/1	0.82	0.38	84,84,84,84	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	XA	1640	1/1	0.82	0.08	101,101,101,101	0
55	MG	YA	2946	1/1	0.82	0.15	87,87,87,87	0
55	MG	RA	2983	1/1	0.82	0.30	84,84,84,84	0
55	MG	YA	3198	1/1	0.82	0.44	124,124,124,124	0
55	MG	RA	3004	1/1	0.82	0.06	205,205,205,205	0
55	MG	RA	3021	1/1	0.82	0.10	96,96,96,96	0
55	MG	QA	1657	1/1	0.82	0.38	157,157,157,157	0
55	MG	XA	1629	1/1	0.82	0.10	135,135,135,135	0
55	MG	YA	3074	1/1	0.82	0.16	84,84,84,84	0
55	MG	YA	3112	1/1	0.82	0.34	96,96,96,96	0
55	MG	RA	3323	1/1	0.82	0.29	205,205,205,205	0
55	MG	RA	3047	1/1	0.82	0.07	139,139,139,139	0
55	MG	RA	3325	1/1	0.82	0.18	84,84,84,84	0
55	MG	RA	2901	1/1	0.82	0.14	98,98,98,98	0
55	MG	YA	2913	1/1	0.82	0.29	84,84,84,84	0
55	MG	YQ	201	1/1	0.82	0.08	111,111,111,111	0
55	MG	YA	2937	1/1	0.82	0.22	84,84,84,84	0
55	MG	RA	3120	1/1	0.82	0.27	129,129,129,129	0
55	MG	YA	3056	1/1	0.82	0.26	84,84,84,84	0
55	MG	YA	3124	1/1	0.82	0.37	84,84,84,84	0
55	MG	RA	3098	1/1	0.83	0.41	212,212,212,212	0
55	MG	RA	3003	1/1	0.83	0.17	108,108,108,108	0
55	MG	RA	3306	1/1	0.83	0.12	111,111,111,111	0
55	MG	RA	3080	1/1	0.83	0.15	108,108,108,108	0
55	MG	RA	3150	1/1	0.83	0.17	84,84,84,84	0
55	MG	RA	2914	1/1	0.83	0.15	129,129,129,129	0
55	MG	YA	3034	1/1	0.83	0.22	84,84,84,84	0
55	MG	YA	2971	1/1	0.83	0.25	84,84,84,84	0
55	MG	RE	303	1/1	0.83	0.18	109,109,109,109	0
55	MG	RA	2927	1/1	0.83	0.10	135,135,135,135	0
55	MG	YA	3252	1/1	0.83	0.31	118,118,118,118	0
55	MG	YA	3041	1/1	0.83	0.25	84,84,84,84	0
55	MG	RF	301	1/1	0.83	0.25	118,118,118,118	0
55	MG	RA	3192	1/1	0.83	0.12	105,105,105,105	0
55	MG	YA	3007	1/1	0.83	0.57	84,84,84,84	0
55	MG	RA	3058	1/1	0.83	0.31	147,147,147,147	0
55	MG	YA	3177	1/1	0.83	0.11	90,90,90,90	0
55	MG	YA	3078	1/1	0.83	0.27	140,140,140,140	0
55	MG	RA	3027	1/1	0.83	0.23	92,92,92,92	0
55	MG	YE	303	1/1	0.83	0.40	124,124,124,124	0
55	MG	RA	3228	1/1	0.83	0.21	105,105,105,105	0
55	MG	RA	3042	1/1	0.83	0.08	111,111,111,111	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	RA	3077	1/1	0.83	0.54	185,185,185,185	0
55	MG	R8	101	1/1	0.83	0.18	111,111,111,111	0
55	MG	YA	3021	1/1	0.83	0.58	84,84,84,84	0
55	MG	YA	2942	1/1	0.83	0.26	120,120,120,120	0
55	MG	RA	3168	1/1	0.84	0.06	206,206,206,206	0
55	MG	RA	3329	1/1	0.84	0.20	93,93,93,93	0
55	MG	RA	2994	1/1	0.84	0.12	90,90,90,90	0
55	MG	YA	3103	1/1	0.84	0.49	84,84,84,84	0
55	MG	YA	3070	1/1	0.84	0.23	84,84,84,84	0
55	MG	QA	1635	1/1	0.84	0.11	112,112,112,112	0
55	MG	YA	2990	1/1	0.84	0.09	84,84,84,84	0
55	MG	RA	3160	1/1	0.84	0.12	109,109,109,109	0
55	MG	YA	3022	1/1	0.84	0.27	84,84,84,84	0
55	MG	YA	3169	1/1	0.84	0.15	119,119,119,119	0
55	MG	YA	2996	1/1	0.84	0.13	84,84,84,84	0
55	MG	YA	3052	1/1	0.84	0.10	84,84,84,84	0
55	MG	XA	1687	1/1	0.84	0.55	172,172,172,172	0
55	MG	RA	2985	1/1	0.84	0.32	139,139,139,139	0
55	MG	XA	1615	1/1	0.84	0.09	162,162,162,162	0
55	MG	RA	3135	1/1	0.84	0.15	100,100,100,100	0
55	MG	RA	2961	1/1	0.84	0.15	101,101,101,101	0
55	MG	YA	2941	1/1	0.84	0.11	106,106,106,106	0
55	MG	RA	2945	1/1	0.84	0.25	84,84,84,84	0
55	MG	YA	3223	1/1	0.84	0.14	84,84,84,84	0
55	MG	YA	3181	1/1	0.84	0.14	115,115,115,115	0
55	MG	YA	3227	1/1	0.84	0.29	90,90,90,90	0
55	MG	YR	201	1/1	0.84	0.14	95,95,95,95	0
55	MG	Y8	102	1/1	0.84	0.32	123,123,123,123	0
55	MG	RA	2934	1/1	0.84	0.23	94,94,94,94	0
55	MG	QA	1608	1/1	0.84	0.18	97,97,97,97	0
55	MG	YA	3040	1/1	0.84	0.18	84,84,84,84	0
55	MG	QA	1656	1/1	0.85	0.17	105,105,105,105	0
55	MG	RA	3110	1/1	0.85	0.14	108,108,108,108	0
55	MG	RA	3151	1/1	0.85	0.12	84,84,84,84	0
55	MG	RA	3298	1/1	0.85	0.17	95,95,95,95	0
55	MG	YA	2910	1/1	0.85	0.20	84,84,84,84	0
55	MG	YA	3009	1/1	0.85	0.24	84,84,84,84	0
55	MG	RA	3012	1/1	0.85	0.13	106,106,106,106	0
55	MG	QH	202	1/1	0.85	0.32	211,211,211,211	0
55	MG	XA	1635	1/1	0.85	0.11	159,159,159,159	0
55	MG	YA	3200	1/1	0.85	0.39	142,142,142,142	0
55	MG	RA	3315	1/1	0.85	0.20	100,100,100,100	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	RA	3119	1/1	0.85	0.11	137,137,137,137	0
55	MG	YA	3157	1/1	0.85	0.25	84,84,84,84	0
55	MG	YE	304	1/1	0.85	0.32	84,84,84,84	0
55	MG	YA	2973	1/1	0.85	0.14	90,90,90,90	0
55	MG	YA	3020	1/1	0.85	0.18	110,110,110,110	0
55	MG	RA	3201	1/1	0.85	0.34	125,125,125,125	0
55	MG	RA	3230	1/1	0.85	0.18	84,84,84,84	0
55	MG	XA	1643	1/1	0.85	0.19	105,105,105,105	0
55	MG	XA	1661	1/1	0.85	0.20	105,105,105,105	0
55	MG	YA	3222	1/1	0.85	0.15	84,84,84,84	0
55	MG	YA	3206	1/1	0.86	0.11	84,84,84,84	0
55	MG	RA	2981	1/1	0.86	0.21	109,109,109,109	0
55	MG	YA	3098	1/1	0.86	0.26	174,174,174,174	0
55	MG	RA	2956	1/1	0.86	0.27	198,198,198,198	0
55	MG	RA	3283	1/1	0.86	0.42	117,117,117,117	0
55	MG	YA	2914	1/1	0.86	0.47	84,84,84,84	0
55	MG	YA	3185	1/1	0.86	0.12	87,87,87,87	0
55	MG	XA	1655	1/1	0.86	0.26	84,84,84,84	0
55	MG	YA	3108	1/1	0.86	0.12	117,117,117,117	0
55	MG	RA	3043	1/1	0.86	0.09	133,133,133,133	0
55	MG	YA	2918	1/1	0.86	0.21	84,84,84,84	0
55	MG	XA	1608	1/1	0.86	0.17	108,108,108,108	0
55	MG	RA	2967	1/1	0.86	0.10	230,230,230,230	0
55	MG	RA	3097	1/1	0.86	0.04	98,98,98,98	0
55	MG	QL	202	1/1	0.86	0.09	95,95,95,95	0
55	MG	RA	3087	1/1	0.86	0.29	164,164,164,164	0
55	MG	YA	2986	1/1	0.86	0.20	84,84,84,84	0
55	MG	YA	3063	1/1	0.86	0.19	104,104,104,104	0
55	MG	XA	1681	1/1	0.86	0.10	171,171,171,171	0
55	MG	XA	1631	1/1	0.86	0.32	167,167,167,167	0
55	MG	RA	2977	1/1	0.86	0.11	84,84,84,84	0
55	MG	YA	2970	1/1	0.86	0.45	84,84,84,84	0
55	MG	RA	3089	1/1	0.86	0.17	198,198,198,198	0
55	MG	RA	2951	1/1	0.87	0.24	255,255,255,255	0
55	MG	RA	3055	1/1	0.87	0.24	123,123,123,123	0
55	MG	RA	2919	1/1	0.87	0.10	130,130,130,130	0
55	MG	YA	2901	1/1	0.87	0.32	242,242,242,242	0
55	MG	RA	3236	1/1	0.87	0.23	84,84,84,84	0
55	MG	RA	2979	1/1	0.87	0.11	135,135,135,135	0
55	MG	RA	3303	1/1	0.87	0.20	256,256,256,256	0
55	MG	QA	1661	1/1	0.87	0.12	108,108,108,108	0
55	MG	XA	1645	1/1	0.87	0.21	106,106,106,106	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	YA	3283	1/1	0.87	0.29	131,131,131,131	0
55	MG	YA	2939	1/1	0.87	0.11	84,84,84,84	0
55	MG	YA	3285	1/1	0.87	0.13	94,94,94,94	0
55	MG	RA	3121	1/1	0.87	0.30	177,177,177,177	0
55	MG	RA	3181	1/1	0.87	0.14	180,180,180,180	0
55	MG	QA	1646	1/1	0.87	0.07	84,84,84,84	0
55	MG	RA	3062	1/1	0.87	0.26	84,84,84,84	0
55	MG	RA	3246	1/1	0.87	0.20	84,84,84,84	0
55	MG	XA	1626	1/1	0.87	0.05	112,112,112,112	0
55	MG	YA	3071	1/1	0.87	0.23	84,84,84,84	0
55	MG	RA	3250	1/1	0.87	0.21	84,84,84,84	0
55	MG	RA	3006	1/1	0.87	0.17	84,84,84,84	0
55	MG	RA	3288	1/1	0.87	0.14	99,99,99,99	0
55	MG	RA	3041	1/1	0.87	0.07	131,131,131,131	0
55	MG	RA	3292	1/1	0.87	0.14	105,105,105,105	0
55	MG	XA	1659	1/1	0.87	0.15	84,84,84,84	0
55	MG	YA	3080	1/1	0.87	0.12	94,94,94,94	0
55	MG	YA	3226	1/1	0.87	0.06	100,100,100,100	0
55	MG	YA	3081	1/1	0.87	0.18	84,84,84,84	0
55	MG	RA	3191	1/1	0.87	0.39	148,148,148,148	0
55	MG	RA	3033	1/1	0.87	0.16	95,95,95,95	0
55	MG	RA	3231	1/1	0.88	0.10	141,141,141,141	0
55	MG	QA	1639	1/1	0.88	0.08	135,135,135,135	0
55	MG	YA	3147	1/1	0.88	0.19	84,84,84,84	0
55	MG	RA	3215	1/1	0.88	0.10	102,102,102,102	0
55	MG	RA	2960	1/1	0.88	0.22	108,108,108,108	0
55	MG	XA	1627	1/1	0.88	0.41	165,165,165,165	0
55	MG	RA	3185	1/1	0.88	0.24	84,84,84,84	0
55	MG	QA	1612	1/1	0.88	0.09	141,141,141,141	0
55	MG	RA	3277	1/1	0.88	0.13	182,182,182,182	0
55	MG	RA	3301	1/1	0.88	0.17	142,142,142,142	0
55	MG	YA	3217	1/1	0.88	0.17	104,104,104,104	0
55	MG	XA	1609	1/1	0.88	0.20	96,96,96,96	0
55	MG	YA	3220	1/1	0.88	0.17	84,84,84,84	0
55	MG	QA	1659	1/1	0.88	0.04	125,125,125,125	0
55	MG	RA	3016	1/1	0.88	0.18	84,84,84,84	0
55	MG	RA	3100	1/1	0.88	0.60	165,165,165,165	0
55	MG	RA	3247	1/1	0.88	0.19	84,84,84,84	0
55	MG	YE	301	1/1	0.88	0.24	113,113,113,113	0
55	MG	YA	2966	1/1	0.88	0.28	84,84,84,84	0
55	MG	RA	3148	1/1	0.88	0.21	89,89,89,89	0
55	MG	YA	3266	1/1	0.88	0.22	102,102,102,102	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(Å ²)	Q<0.9
55	MG	RA	2988	1/1	0.88	0.16	101,101,101,101	0
55	MG	QA	1652	1/1	0.88	0.09	162,162,162,162	0
55	MG	XA	1667	1/1	0.88	0.13	173,173,173,173	0
55	MG	YA	3029	1/1	0.88	0.18	84,84,84,84	0
55	MG	YA	3235	1/1	0.88	0.25	84,84,84,84	0
55	MG	RA	3044	1/1	0.88	0.38	156,156,156,156	0
57	ZN	R9	101	1/1	0.88	0.10	314,314,314,314	0
55	MG	RA	3155	1/1	0.88	0.13	95,95,95,95	0
55	MG	YA	2920	1/1	0.89	0.12	84,84,84,84	0
55	MG	RA	3068	1/1	0.89	0.16	113,113,113,113	0
55	MG	YA	2993	1/1	0.89	0.19	84,84,84,84	0
55	MG	RA	3113	1/1	0.89	0.12	84,84,84,84	0
55	MG	RA	3149	1/1	0.89	0.42	85,85,85,85	0
55	MG	YA	2909	1/1	0.89	0.25	84,84,84,84	0
55	MG	YA	2998	1/1	0.89	0.33	84,84,84,84	0
55	MG	RE	302	1/1	0.89	0.25	84,84,84,84	0
55	MG	YA	2978	1/1	0.89	0.10	90,90,90,90	0
55	MG	YA	3269	1/1	0.89	0.25	84,84,84,84	0
55	MG	RA	2948	1/1	0.89	0.12	84,84,84,84	0
55	MG	RA	3209	1/1	0.89	0.14	165,165,165,165	0
55	MG	XA	1611	1/1	0.89	0.41	278,278,278,278	0
55	MG	YA	2949	1/1	0.89	0.17	84,84,84,84	0
55	MG	XA	1666	1/1	0.89	0.15	113,113,113,113	0
55	MG	RA	3165	1/1	0.89	0.13	197,197,197,197	0
55	MG	RA	2946	1/1	0.89	0.17	96,96,96,96	0
55	MG	YA	3036	1/1	0.89	0.23	84,84,84,84	0
55	MG	RA	3197	1/1	0.89	0.34	84,84,84,84	0
55	MG	YA	2987	1/1	0.89	0.23	84,84,84,84	0
55	MG	YA	3064	1/1	0.89	0.17	84,84,84,84	0
57	ZN	RY	201	1/1	0.89	0.07	344,344,344,344	0
55	MG	RA	3317	1/1	0.89	0.11	84,84,84,84	0
55	MG	YA	3201	1/1	0.90	0.36	139,139,139,139	0
55	MG	RA	3114	1/1	0.90	0.10	84,84,84,84	0
55	MG	RA	3153	1/1	0.90	0.08	117,117,117,117	0
55	MG	RA	3184	1/1	0.90	0.10	103,103,103,103	0
55	MG	YA	3246	1/1	0.90	0.10	129,129,129,129	0
55	MG	RA	3096	1/1	0.90	0.11	122,122,122,122	0
55	MG	RA	3126	1/1	0.90	0.08	137,137,137,137	0
55	MG	RA	3105	1/1	0.90	0.18	107,107,107,107	0
55	MG	RA	3032	1/1	0.90	0.20	84,84,84,84	0
55	MG	RA	3131	1/1	0.90	0.12	117,117,117,117	0
55	MG	RA	3254	1/1	0.90	0.11	130,130,130,130	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	YA	3289	1/1	0.90	0.11	84,84,84,84	0
55	MG	RA	2993	1/1	0.90	0.07	141,141,141,141	0
55	MG	RA	3109	1/1	0.90	0.12	84,84,84,84	0
55	MG	YA	2923	1/1	0.90	0.16	84,84,84,84	0
55	MG	YA	3125	1/1	0.90	0.26	89,89,89,89	0
55	MG	YA	3037	1/1	0.90	0.26	84,84,84,84	0
55	MG	RA	3264	1/1	0.90	0.14	148,148,148,148	0
55	MG	YA	2947	1/1	0.90	0.10	84,84,84,84	0
55	MG	QA	1644	1/1	0.90	0.09	84,84,84,84	0
55	MG	RA	3178	1/1	0.90	0.41	232,232,232,232	0
55	MG	YA	3230	1/1	0.90	0.20	84,84,84,84	0
55	MG	YA	3016	1/1	0.90	0.05	84,84,84,84	0
55	MG	RA	3050	1/1	0.90	0.22	110,110,110,110	0
55	MG	YA	2992	1/1	0.90	0.12	84,84,84,84	0
55	MG	YA	3045	1/1	0.90	0.23	84,84,84,84	0
55	MG	YA	3166	1/1	0.90	0.21	144,144,144,144	0
55	MG	YA	3019	1/1	0.90	0.07	84,84,84,84	0
55	MG	RA	3009	1/1	0.90	0.11	101,101,101,101	0
55	MG	RA	3040	1/1	0.90	0.12	89,89,89,89	0
55	MG	RA	3287	1/1	0.91	0.11	123,123,123,123	0
55	MG	YA	3238	1/1	0.91	0.10	129,129,129,129	0
55	MG	YA	3105	1/1	0.91	0.17	135,135,135,135	0
55	MG	RA	3099	1/1	0.91	0.29	84,84,84,84	0
55	MG	YA	3280	1/1	0.91	0.06	120,120,120,120	0
55	MG	RA	3232	1/1	0.91	0.06	99,99,99,99	0
55	MG	RA	3291	1/1	0.91	0.15	84,84,84,84	0
55	MG	YA	3138	1/1	0.91	0.17	98,98,98,98	0
55	MG	RA	3234	1/1	0.91	0.07	101,101,101,101	0
55	MG	QA	1650	1/1	0.91	0.24	108,108,108,108	0
55	MG	QA	1653	1/1	0.91	0.31	130,130,130,130	0
55	MG	XA	1662	1/1	0.91	0.10	141,141,141,141	0
55	MG	XA	1636	1/1	0.91	0.12	95,95,95,95	0
55	MG	RA	2980	1/1	0.91	0.18	84,84,84,84	0
55	MG	RA	3239	1/1	0.91	0.10	84,84,84,84	0
55	MG	YA	3216	1/1	0.91	0.16	84,84,84,84	0
55	MG	RA	3128	1/1	0.91	0.21	187,187,187,187	0
55	MG	XA	1617	1/1	0.91	0.16	84,84,84,84	0
55	MG	YA	3121	1/1	0.91	0.09	84,84,84,84	0
55	MG	RA	3262	1/1	0.91	0.26	194,194,194,194	0
55	MG	RA	3300	1/1	0.91	0.13	117,117,117,117	0
55	MG	RF	302	1/1	0.91	0.08	158,158,158,158	0
55	MG	YA	3008	1/1	0.91	0.26	84,84,84,84	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	RA	3319	1/1	0.91	0.11	114,114,114,114	0
55	MG	XA	1622	1/1	0.91	0.11	132,132,132,132	0
55	MG	RA	2996	1/1	0.91	0.18	177,177,177,177	0
55	MG	RA	3321	1/1	0.91	0.12	88,88,88,88	0
55	MG	YA	3193	1/1	0.91	0.33	176,176,176,176	0
55	MG	YX	101	1/1	0.91	0.08	84,84,84,84	0
55	MG	YA	3160	1/1	0.91	0.09	112,112,112,112	0
55	MG	XA	1604	1/1	0.91	0.12	176,176,176,176	0
55	MG	RA	3219	1/1	0.91	0.38	143,143,143,143	0
55	MG	RA	2915	1/1	0.91	0.10	84,84,84,84	0
55	MG	XA	1663	1/1	0.92	0.11	115,115,115,115	0
55	MG	XA	1684	1/1	0.92	0.13	210,210,210,210	0
55	MG	RA	3019	1/1	0.92	0.13	84,84,84,84	0
55	MG	RA	3267	1/1	0.92	0.16	143,143,143,143	0
55	MG	RA	3014	1/1	0.92	0.13	84,84,84,84	0
55	MG	XA	1648	1/1	0.92	0.05	137,137,137,137	0
55	MG	RA	2997	1/1	0.92	0.27	161,161,161,161	0
55	MG	YA	3264	1/1	0.92	0.20	84,84,84,84	0
55	MG	YA	3046	1/1	0.92	0.19	124,124,124,124	0
55	MG	YA	3047	1/1	0.92	0.15	84,84,84,84	0
55	MG	RN	201	1/1	0.92	0.13	98,98,98,98	0
55	MG	YA	3100	1/1	0.92	0.18	84,84,84,84	0
55	MG	QA	1640	1/1	0.92	0.30	101,101,101,101	0
55	MG	RA	3188	1/1	0.92	0.22	140,140,140,140	0
55	MG	YA	3028	1/1	0.92	0.20	84,84,84,84	0
55	MG	RA	3069	1/1	0.92	0.57	189,189,189,189	0
55	MG	RA	3091	1/1	0.92	0.28	132,132,132,132	0
55	MG	RA	2958	1/1	0.92	0.19	91,91,91,91	0
55	MG	RA	3257	1/1	0.92	0.19	86,86,86,86	0
55	MG	RA	3258	1/1	0.92	0.25	165,165,165,165	0
55	MG	RA	3063	1/1	0.92	0.17	84,84,84,84	0
55	MG	YA	3278	1/1	0.92	0.06	85,85,85,85	0
55	MG	YA	3190	1/1	0.92	0.28	99,99,99,99	0
55	MG	XA	1624	1/1	0.92	0.06	115,115,115,115	0
55	MG	RA	3227	1/1	0.92	0.26	123,123,123,123	0
55	MG	RA	3081	1/1	0.92	0.20	84,84,84,84	0
55	MG	RA	3284	1/1	0.93	0.10	108,108,108,108	0
55	MG	RA	2949	1/1	0.93	0.13	96,96,96,96	0
55	MG	RA	3248	1/1	0.93	0.09	105,105,105,105	0
55	MG	RA	3249	1/1	0.93	0.10	84,84,84,84	0
55	MG	YA	3139	1/1	0.93	0.07	84,84,84,84	0
55	MG	RO	201	1/1	0.93	0.35	186,186,186,186	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	QA	1667	1/1	0.93	0.13	154,154,154,154	0
55	MG	YA	3000	1/1	0.93	0.10	93,93,93,93	0
55	MG	YA	2917	1/1	0.93	0.23	98,98,98,98	0
55	MG	RA	2978	1/1	0.93	0.22	84,84,84,84	0
55	MG	YA	3092	1/1	0.93	0.09	149,149,149,149	0
55	MG	YD	301	1/1	0.93	0.14	86,86,86,86	0
55	MG	RA	3212	1/1	0.93	0.10	170,170,170,170	0
55	MG	RA	3253	1/1	0.93	0.23	84,84,84,84	0
55	MG	RA	3007	1/1	0.93	0.13	87,87,87,87	0
55	MG	RA	2944	1/1	0.93	0.11	84,84,84,84	0
55	MG	RA	3189	1/1	0.93	0.31	174,174,174,174	0
55	MG	RA	3216	1/1	0.93	0.12	122,122,122,122	0
55	MG	YA	2928	1/1	0.93	0.18	103,103,103,103	0
55	MG	RA	3259	1/1	0.93	0.19	97,97,97,97	0
55	MG	YA	3076	1/1	0.93	0.24	158,158,158,158	0
55	MG	RD	301	1/1	0.93	0.19	128,128,128,128	0
55	MG	RA	3244	1/1	0.93	0.08	144,144,144,144	0
55	MG	YA	3106	1/1	0.93	0.12	84,84,84,84	0
55	MG	YA	3079	1/1	0.93	0.12	84,84,84,84	0
55	MG	YA	3221	1/1	0.93	0.10	84,84,84,84	0
55	MG	RA	3235	1/1	0.93	0.14	84,84,84,84	0
55	MG	RA	3279	1/1	0.94	0.04	169,169,169,169	0
55	MG	XA	1677	1/1	0.94	0.05	147,147,147,147	0
55	MG	YA	3204	1/1	0.94	0.13	135,135,135,135	0
55	MG	XA	1653	1/1	0.94	0.11	84,84,84,84	0
55	MG	RA	2990	1/1	0.94	0.14	103,103,103,103	0
55	MG	RA	3268	1/1	0.94	0.13	84,84,84,84	0
55	MG	RA	3024	1/1	0.94	0.08	159,159,159,159	0
55	MG	YA	3099	1/1	0.94	0.18	84,84,84,84	0
55	MG	QA	1611	1/1	0.94	0.15	124,124,124,124	0
55	MG	RA	2922	1/1	0.94	0.15	105,105,105,105	0
55	MG	YA	3241	1/1	0.94	0.18	84,84,84,84	0
55	MG	YA	3215	1/1	0.94	0.18	84,84,84,84	0
55	MG	RA	3060	1/1	0.94	0.12	106,106,106,106	0
55	MG	RA	3061	1/1	0.94	0.18	91,91,91,91	0
55	MG	RA	3158	1/1	0.94	0.07	117,117,117,117	0
55	MG	YF	301	1/1	0.94	0.09	90,90,90,90	0
55	MG	YA	3170	1/1	0.94	0.09	84,84,84,84	0
55	MG	RE	301	1/1	0.94	0.10	84,84,84,84	0
55	MG	YA	3249	1/1	0.94	0.07	131,131,131,131	0
55	MG	YA	2991	1/1	0.94	0.10	95,95,95,95	0
55	MG	RA	3266	1/1	0.94	0.14	305,305,305,305	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	XO	101	1/1	0.94	0.05	151,151,151,151	0
55	MG	YA	2960	1/1	0.94	0.25	84,84,84,84	0
55	MG	YA	3014	1/1	0.94	0.04	84,84,84,84	0
55	MG	Y1	101	1/1	0.94	0.07	117,117,117,117	0
55	MG	YA	3113	1/1	0.94	0.05	84,84,84,84	0
55	MG	RA	3312	1/1	0.95	0.16	115,115,115,115	0
55	MG	RA	3269	1/1	0.95	0.06	200,200,200,200	0
55	MG	YA	3163	1/1	0.95	0.10	104,104,104,104	0
55	MG	YA	2924	1/1	0.95	0.12	84,84,84,84	0
55	MG	YA	3116	1/1	0.95	0.29	133,133,133,133	0
55	MG	RA	3130	1/1	0.95	0.10	119,119,119,119	0
55	MG	RA	3072	1/1	0.95	0.05	142,142,142,142	0
55	MG	RA	2925	1/1	0.95	0.18	84,84,84,84	0
55	MG	YA	3260	1/1	0.95	0.04	84,84,84,84	0
55	MG	XA	1639	1/1	0.95	0.12	84,84,84,84	0
55	MG	RA	3274	1/1	0.95	0.09	149,149,149,149	0
55	MG	YA	3293	1/1	0.95	0.09	102,102,102,102	0
55	MG	RA	3015	1/1	0.95	0.16	141,141,141,141	0
55	MG	YA	3172	1/1	0.95	0.18	84,84,84,84	0
55	MG	YA	3146	1/1	0.95	0.08	84,84,84,84	0
55	MG	RA	3142	1/1	0.95	0.19	152,152,152,152	0
55	MG	RA	3263	1/1	0.95	0.06	136,136,136,136	0
55	MG	RA	3090	1/1	0.95	0.18	138,138,138,138	0
55	MG	YA	3033	1/1	0.95	0.09	84,84,84,84	0
55	MG	Y7	101	1/1	0.95	0.26	208,208,208,208	0
55	MG	YA	3152	1/1	0.95	0.11	84,84,84,84	0
55	MG	YA	3012	1/1	0.95	0.12	84,84,84,84	0
55	MG	RA	2973	1/1	0.95	0.17	84,84,84,84	0
55	MG	RA	3000	1/1	0.95	0.04	109,109,109,109	0
55	MG	YA	2994	1/1	0.95	0.52	84,84,84,84	0
55	MG	RA	3309	1/1	0.95	0.07	120,120,120,120	0
55	MG	RA	3233	1/1	0.95	0.11	84,84,84,84	0
55	MG	YA	2977	1/1	0.95	0.09	84,84,84,84	0
55	MG	RA	2991	1/1	0.95	0.02	110,110,110,110	0
55	MG	YA	3219	1/1	0.95	0.07	91,91,91,91	0
55	MG	RA	3145	1/1	0.96	0.14	112,112,112,112	0
55	MG	RA	3261	1/1	0.96	0.07	191,191,191,191	0
55	MG	RA	2955	1/1	0.96	0.09	192,192,192,192	0
55	MG	YA	3023	1/1	0.96	0.07	84,84,84,84	0
55	MG	QA	1630	1/1	0.96	0.10	137,137,137,137	0
55	MG	YA	3225	1/1	0.96	0.14	100,100,100,100	0
55	MG	YA	3184	1/1	0.96	0.08	95,95,95,95	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	RA	3290	1/1	0.96	0.06	95,95,95,95	0
55	MG	YA	3251	1/1	0.96	0.06	99,99,99,99	0
55	MG	YA	3058	1/1	0.96	0.05	84,84,84,84	0
55	MG	R3	101	1/1	0.96	0.10	94,94,94,94	0
55	MG	YA	3207	1/1	0.96	0.07	84,84,84,84	0
55	MG	XA	1638	1/1	0.96	0.18	84,84,84,84	0
55	MG	RA	2932	1/1	0.96	0.05	131,131,131,131	0
55	MG	RA	3196	1/1	0.96	0.24	168,168,168,168	0
55	MG	RA	3159	1/1	0.96	0.12	104,104,104,104	0
55	MG	RA	3187	1/1	0.96	0.14	130,130,130,130	0
55	MG	RA	3067	1/1	0.96	0.05	101,101,101,101	0
55	MG	YA	3261	1/1	0.96	0.07	190,190,190,190	0
56	SF4	QD	301	8/8	0.96	0.06	181,237,336,420	0
55	MG	RA	3270	1/1	0.96	0.11	84,84,84,84	0
55	MG	YA	3084	1/1	0.96	0.17	84,84,84,84	0
55	MG	RA	3049	1/1	0.96	0.10	84,84,84,84	0
55	MG	RA	2984	1/1	0.96	0.07	84,84,84,84	0
57	ZN	Y9	101	1/1	0.96	0.05	162,162,162,162	0
55	MG	RA	3331	1/1	0.97	0.09	84,84,84,84	0
55	MG	QA	1633	1/1	0.97	0.05	190,190,190,190	0
55	MG	XA	1683	1/1	0.97	0.06	120,120,120,120	0
55	MG	YA	3102	1/1	0.97	0.16	178,178,178,178	0
55	MG	YA	3214	1/1	0.97	0.04	95,95,95,95	0
55	MG	RA	3082	1/1	0.97	0.10	84,84,84,84	0
55	MG	RA	3070	1/1	0.97	0.13	181,181,181,181	0
55	MG	YA	3294	1/1	0.97	0.04	103,103,103,103	0
55	MG	RA	3084	1/1	0.97	0.11	150,150,150,150	0
56	SF4	XD	301	8/8	0.97	0.05	163,216,235,295	0
55	MG	RA	3255	1/1	0.97	0.05	99,99,99,99	0
57	ZN	R5	101	1/1	0.97	0.03	268,268,268,268	0
57	ZN	R6	101	1/1	0.97	0.05	149,149,149,149	0
55	MG	YA	3244	1/1	0.97	0.12	84,84,84,84	0
55	MG	YA	2925	1/1	0.97	0.11	84,84,84,84	0
55	MG	RA	3245	1/1	0.97	0.19	93,93,93,93	0
57	ZN	Y5	101	1/1	0.97	0.07	308,308,308,308	0
55	MG	YA	2927	1/1	0.97	0.10	103,103,103,103	0
55	MG	RA	2940	1/1	0.98	0.04	84,84,84,84	0
55	MG	YA	2902	1/1	0.98	0.09	86,86,86,86	0
55	MG	RA	2902	1/1	0.98	0.04	84,84,84,84	0
55	MG	RA	3108	1/1	0.98	0.06	117,117,117,117	0
55	MG	RA	3296	1/1	0.98	0.07	121,121,121,121	0
55	MG	RA	3154	1/1	0.98	0.06	127,127,127,127	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	YA	3229	1/1	0.98	0.07	84,84,84,84	0
55	MG	QA	1668	1/1	0.98	0.06	114,114,114,114	0
55	MG	YA	3208	1/1	0.98	0.09	100,100,100,100	0
55	MG	RA	3085	1/1	0.98	0.09	140,140,140,140	0
55	MG	YA	3089	1/1	0.98	0.07	84,84,84,84	0
55	MG	RA	3229	1/1	0.98	0.09	84,84,84,84	0
55	MG	RA	3316	1/1	0.99	0.03	84,84,84,84	0
57	ZN	Y6	101	1/1	0.99	0.02	144,144,144,144	0
55	MG	RA	2971	1/1	0.99	0.04	84,84,84,84	0
57	ZN	YY	201	1/1	0.99	0.02	193,193,193,193	0

6.5 Other polymers [i](#)

There are no such residues in this entry.