



Full wwPDB X-ray Structure Validation Report ⓘ

Sep 21, 2017 – 08:26 PM EDT

PDB ID : 5W4K
Title : Crystal structure of the *Thermus thermophilus* 70S ribosome in complex with Klebsazolicin and bound to mRNA and A-, P- and E-site tRNAs at 2.7Å resolution
Authors : Metelev, M.; Osterman, I.A.; Ghilarov, D.; Khabibullina, N.F.; Yakimov, A.; Shabalin, K.; Utkina, I.; Travin, D.Y.; Komarova, E.S.; Serebryakova, M.; Artamonova, T.; Khodorkovskii, M.; Konevega, A.L.; Sergiev, P.V.; Severinov, K.; Polikanov, Y.S.
Deposited on : unknown
Resolution : 2.70 Å(reported)

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

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

A user guide is available at

<http://wwpdb.org/validation/2016/XrayValidationReportHelp>
with specific help available everywhere you see the ⓘ symbol.

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

MolProbity : 4.02b-467
Mogul : 1.7.2 (RC1), CSD as538be (2017)
Xtriage (Phenix) : 1.9-1692
EDS : rb-20029824
Percentile statistics : 20161228.v01 (using entries in the PDB archive December 28th 2016)
Refmac : 5.8.0135
CCP4 : 6.5.0
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : rb-20029824

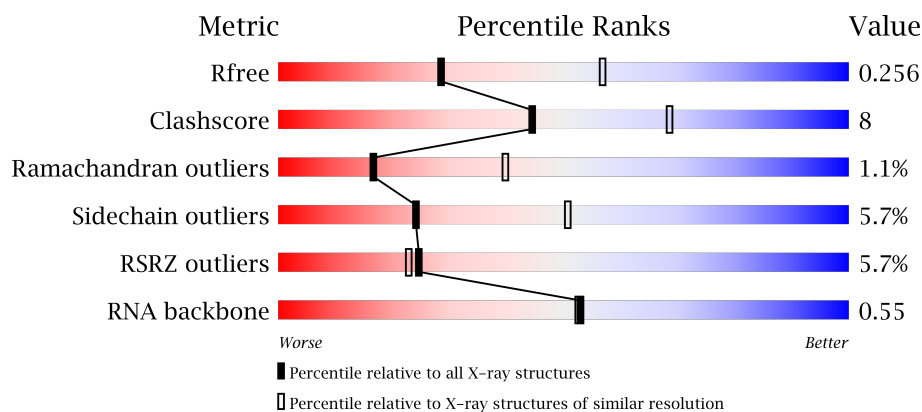
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

The reported resolution of this entry is 2.70 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}	100719	2259 (2.70-2.70)
Clashscore	112137	2590 (2.70-2.70)
Ramachandran outliers	110173	2550 (2.70-2.70)
Sidechain outliers	110143	2550 (2.70-2.70)
RSRZ outliers	101464	2275 (2.70-2.70)
RNA backbone	2435	1011 (3.06-2.34)

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

Mol	Chain	Length	Quality of chain
1	1A	2915	<div> <div>2%</div> <div>67% 25% 6% ..</div> </div>
1	2A	2915	<div> <div>3%</div> <div>59% 31% 6% .</div> </div>
2	1B	121	<div> <div></div> <div>79% 17% ...</div> </div>
2	2B	121	<div> <div></div> <div>43% 43% 13% .</div> </div>









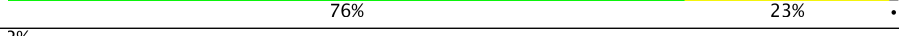

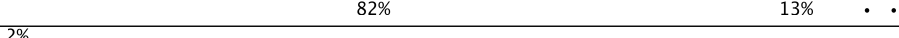
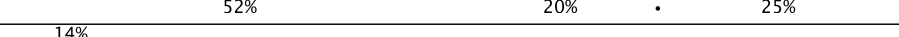

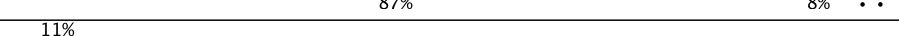


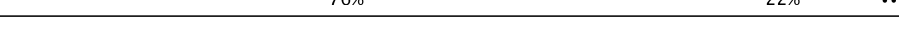

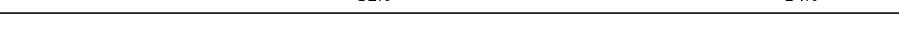






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Mol	Chain	Length	Quality of chain
3	1D	276	
3	2D	276	
4	1E	206	
4	2E	206	
5	1F	210	
5	2F	210	
6	1G	182	
6	2G	182	
7	1H	180	
7	2H	180	
8	1I	148	
8	2I	148	
9	1N	140	
9	2N	140	
10	1O	122	
10	2O	122	
11	1P	150	
11	2P	150	
12	1Q	141	
12	2Q	141	
13	1R	118	
13	2R	118	
14	1S	112	
14	2S	112	
15	1T	146	

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Mol	Chain	Length	Quality of chain
15	2T	146	
16	1U	118	
16	2U	118	
17	1V	101	
17	2V	101	
18	1W	113	
18	2W	113	
19	1X	96	
19	2X	96	
20	1Y	110	
20	2Y	110	
21	1Z	206	
21	2Z	206	
22	10	85	
22	20	85	
23	11	98	
23	21	98	
24	12	72	
24	22	72	
25	13	60	
25	23	60	
26	14	71	
26	24	71	
27	15	60	
27	25	60	

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Mol	Chain	Length	Quality of chain
28	16	54	
28	26	54	
29	17	49	
29	27	49	
30	18	65	
30	28	65	
31	19	37	
31	29	37	
32	1a	1521	
33	1b	256	
33	2b	256	
34	1c	239	
34	2c	239	
35	1d	209	
35	2d	209	
36	1e	162	
36	2e	162	
37	1f	101	
37	2f	101	
38	1g	156	
38	2g	156	
39	1h	138	
39	2h	138	
40	1i	128	
40	2i	128	

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Mol	Chain	Length	Quality of chain
41	1j	105	
41	2j	105	
42	1k	129	
42	2k	129	
43	1l	132	
43	2l	132	
44	1m	126	
44	2m	126	
45	1n	61	
45	2n	61	
46	1o	89	
46	2o	89	
47	1p	88	
47	2p	88	
48	1q	105	
48	2q	105	
49	1r	88	
49	2r	88	
50	1s	93	
50	2s	93	
51	1t	106	
51	2t	106	
52	1u	27	
52	2u	27	
53	1v	24	

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Mol	Chain	Length	Quality of chain
53	2v	24	
54	1w	76	
54	2w	76	
55	1x	77	
55	2x	77	
56	1y	76	
56	2y	76	
57	2a	1521	
58	A	18	
58	B	18	

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

Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
59	MG	10	105	-	-	-	X
59	MG	12	3002	-	-	-	X
59	MG	18	102	-	-	-	X
59	MG	1A	3011	-	-	-	X
59	MG	1A	3026	-	-	-	X
59	MG	1A	3027	-	-	-	X
59	MG	1A	3029	-	-	-	X
59	MG	1A	3033	-	-	-	X
59	MG	1A	3035	-	-	-	X
59	MG	1A	3036	-	-	-	X
59	MG	1A	3037	-	-	-	X
59	MG	1A	3065	-	-	-	X
59	MG	1A	3082	-	-	-	X
59	MG	1A	3090	-	-	-	X
59	MG	1A	3103	-	-	-	X
59	MG	1A	3105	-	-	-	X
59	MG	1A	3106	-	-	-	X
59	MG	1A	3108	-	-	-	X
59	MG	1A	3128	-	-	-	X
59	MG	1A	3129	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
59	MG	1A	3134	-	-	-	X
59	MG	1A	3139	-	-	-	X
59	MG	1A	3140	-	-	-	X
59	MG	1A	3142	-	-	-	X
59	MG	1A	3144	-	-	-	X
59	MG	1A	3147	-	-	-	X
59	MG	1A	3148	-	-	-	X
59	MG	1A	3159	-	-	-	X
59	MG	1A	3160	-	-	-	X
59	MG	1A	3161	-	-	-	X
59	MG	1A	3164	-	-	-	X
59	MG	1A	3165	-	-	-	X
59	MG	1A	3170	-	-	-	X
59	MG	1A	3177	-	-	-	X
59	MG	1A	3186	-	-	-	X
59	MG	1A	3197	-	-	-	X
59	MG	1A	3198	-	-	-	X
59	MG	1A	3204	-	-	-	X
59	MG	1A	3207	-	-	-	X
59	MG	1A	3209	-	-	-	X
59	MG	1A	3218	-	-	-	X
59	MG	1A	3234	-	-	-	X
59	MG	1A	3244	-	-	-	X
59	MG	1A	3277	-	-	-	X
59	MG	1A	3288	-	-	-	X
59	MG	1A	3289	-	-	-	X
59	MG	1A	3290	-	-	-	X
59	MG	1A	3294	-	-	-	X
59	MG	1A	3317	-	-	-	X
59	MG	1A	3327	-	-	-	X
59	MG	1A	3354	-	-	-	X
59	MG	1A	3370	-	-	-	X
59	MG	1A	3371	-	-	-	X
59	MG	1A	3373	-	-	-	X
59	MG	1A	3389	-	-	-	X
59	MG	1A	3403	-	-	-	X
59	MG	1A	3413	-	-	-	X
59	MG	1A	3417	-	-	-	X
59	MG	1A	3427	-	-	-	X
59	MG	1A	3434	-	-	-	X
59	MG	1A	3435	-	-	-	X
59	MG	1A	3450	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
59	MG	1A	3464	-	-	-	X
59	MG	1A	3466	-	-	-	X
59	MG	1A	3467	-	-	-	X
59	MG	1A	3514	-	-	-	X
59	MG	1A	3523	-	-	-	X
59	MG	1A	3529	-	-	-	X
59	MG	1A	3531	-	-	-	X
59	MG	1A	3532	-	-	-	X
59	MG	1A	3535	-	-	-	X
59	MG	1A	3675	-	-	-	X
59	MG	1A	3702	-	-	-	X
59	MG	1A	3739	-	-	-	X
59	MG	1A	3748	-	-	-	X
59	MG	1A	3754	-	-	-	X
59	MG	1A	3788	-	-	-	X
59	MG	1A	3798	-	-	-	X
59	MG	1A	3800	-	-	-	X
59	MG	1A	3802	-	-	-	X
59	MG	1A	3804	-	-	-	X
59	MG	1A	3806	-	-	-	X
59	MG	1A	3808	-	-	-	X
59	MG	1A	3825	-	-	-	X
59	MG	1A	3831	-	-	-	X
59	MG	1A	3844	-	-	-	X
59	MG	1A	3995	-	-	-	X
59	MG	1A	4019	-	-	-	X
59	MG	1A	4020	-	-	-	X
59	MG	1A	4087	-	-	-	X
59	MG	1A	4135	-	-	-	X
59	MG	1A	4169	-	-	-	X
59	MG	1A	4170	-	-	-	X
59	MG	1A	4171	-	-	-	X
59	MG	1A	4172	-	-	-	X
59	MG	1A	4174	-	-	-	X
59	MG	1A	4177	-	-	-	X
59	MG	1A	4180	-	-	-	X
59	MG	1A	4182	-	-	-	X
59	MG	1A	4190	-	-	-	X
59	MG	1A	4195	-	-	-	X
59	MG	1A	4196	-	-	-	X
59	MG	1A	4197	-	-	-	X
59	MG	1A	4198	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
59	MG	1A	4199	-	-	-	X
59	MG	1A	4200	-	-	-	X
59	MG	1A	4202	-	-	-	X
59	MG	1A	4203	-	-	-	X
59	MG	1A	4206	-	-	-	X
59	MG	1A	4207	-	-	-	X
59	MG	1A	4208	-	-	-	X
59	MG	1A	4209	-	-	-	X
59	MG	1A	4210	-	-	-	X
59	MG	1A	4211	-	-	-	X
59	MG	1A	4212	-	-	-	X
59	MG	1A	4214	-	-	-	X
59	MG	1A	4218	-	-	-	X
59	MG	1A	4219	-	-	-	X
59	MG	1A	4220	-	-	-	X
59	MG	1A	4225	-	-	-	X
59	MG	1A	4227	-	-	-	X
59	MG	1A	4228	-	-	-	X
59	MG	1B	211	-	-	-	X
59	MG	1B	223	-	-	-	X
59	MG	1D	301	-	-	-	X
59	MG	1D	304	-	-	-	X
59	MG	1D	305	-	-	-	X
59	MG	1D	307	-	-	-	X
59	MG	1D	308	-	-	-	X
59	MG	1D	310	-	-	-	X
59	MG	1D	311	-	-	-	X
59	MG	1D	313	-	-	-	X
59	MG	1E	313	-	-	-	X
59	MG	1F	304	-	-	-	X
59	MG	1F	305	-	-	-	X
59	MG	1F	310	-	-	-	X
59	MG	1N	3003	-	-	-	X
59	MG	1N	3006	-	-	-	X
59	MG	1N	3007	-	-	-	X
59	MG	1N	3009	-	-	-	X
59	MG	1O	3001	-	-	-	X
59	MG	1O	3002	-	-	-	X
59	MG	1P	201	-	-	-	X
59	MG	1P	202	-	-	-	X
59	MG	1R	202	-	-	-	X
59	MG	1S	3001	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
59	MG	1S	3002	-	-	-	X
59	MG	1T	8002	-	-	-	X
59	MG	1U	201	-	-	-	X
59	MG	1U	202	-	-	-	X
59	MG	1U	203	-	-	-	X
59	MG	1U	205	-	-	-	X
59	MG	1W	203	-	-	-	X
59	MG	1W	204	-	-	-	X
59	MG	1X	3001	-	-	-	X
59	MG	1X	3003	-	-	-	X
59	MG	1Y	504	-	-	-	X
59	MG	1Z	301	-	-	-	X
59	MG	1a	3313	-	-	-	X
59	MG	1a	3333	-	-	-	X
59	MG	1a	3335	-	-	-	X
59	MG	1a	3340	-	-	-	X
59	MG	1a	3360	-	-	-	X
59	MG	1a	3383	-	-	-	X
59	MG	1a	3411	-	-	-	X
59	MG	1a	3426	-	-	-	X
59	MG	1a	3576	-	-	-	X
59	MG	1f	3001	-	-	-	X
59	MG	1f	3003	-	-	-	X
59	MG	1x	103	-	-	-	X
59	MG	1x	117	-	-	-	X
59	MG	2A	3002	-	-	-	X
59	MG	2A	3014	-	-	-	X
59	MG	2A	3030	-	-	-	X
59	MG	2A	3040	-	-	-	X
59	MG	2A	3056	-	-	-	X
59	MG	2A	3066	-	-	-	X
59	MG	2A	3069	-	-	-	X
59	MG	2A	3076	-	-	-	X
59	MG	2A	3091	-	-	-	X
59	MG	2A	3102	-	-	-	X
59	MG	2A	3108	-	-	-	X
59	MG	2A	3136	-	-	-	X
59	MG	2A	3155	-	-	-	X
59	MG	2A	3187	-	-	-	X
59	MG	2A	3200	-	-	-	X
59	MG	2A	3286	-	-	-	X
59	MG	2A	3301	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
59	MG	2A	3331	-	-	-	X
59	MG	2A	3345	-	-	-	X
59	MG	2A	3355	-	-	-	X
59	MG	2A	3366	-	-	-	X
59	MG	2A	3495	-	-	-	X
59	MG	2A	3514	-	-	-	X
59	MG	2A	3532	-	-	-	X
59	MG	2A	3618	-	-	-	X
59	MG	2A	3667	-	-	-	X
59	MG	2A	3772	-	-	-	X
59	MG	2A	3774	-	-	-	X
59	MG	2A	3823	-	-	-	X
59	MG	2A	3824	-	-	-	X
59	MG	2A	3830	-	-	-	X
59	MG	2A	3831	-	-	-	X
59	MG	2A	3842	-	-	-	X
59	MG	2A	3844	-	-	-	X
59	MG	2A	3845	-	-	-	X
59	MG	2A	3847	-	-	-	X
59	MG	2B	3001	-	-	-	X
59	MG	2D	304	-	-	-	X
59	MG	2D	305	-	-	-	X
59	MG	2F	303	-	-	-	X
59	MG	2T	202	-	-	-	X
59	MG	2U	203	-	-	-	X
59	MG	2U	204	-	-	-	X
59	MG	2X	3001	-	-	-	X
59	MG	2a	3026	-	-	-	X
59	MG	2a	3085	-	-	-	X
59	MG	2a	3112	-	-	-	X
59	MG	2a	3117	-	-	-	X
59	MG	2a	3188	-	-	-	X
60	ZN	16	102	-	-	-	X

2 Entry composition

There are 63 unique types of molecules in this entry. The entry contains 302030 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 23S Ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
1	1A	2871	Total	C	N	O	P	0	0	0
			61852	27531	11572	19878	2871			
1	2A	2800	Total	C	N	O	P	0	0	0
			60322	26848	11284	19390	2800			

- Molecule 2 is a RNA chain called 5S Ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
2	1B	120	Total	C	N	O	P	0	0	0
			2577	1146	476	835	120			
2	2B	120	Total	C	N	O	P	0	0	0
			2575	1146	476	833	120			

- Molecule 3 is a protein called 50S ribosomal protein L2.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
3	1D	275	Total	C	N	O	S	0	0	0
			2136	1349	423	361	3			
3	2D	275	Total	C	N	O	S	0	0	0
			2136	1349	423	361	3			

- Molecule 4 is a protein called 50S ribosomal protein L3.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
4	1E	204	Total	C	N	O	S	0	0	0
			1559	985	298	270	6			
4	2E	204	Total	C	N	O	S	0	0	0
			1559	985	298	270	6			

- Molecule 5 is a protein called 50S ribosomal protein L4.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
5	1F	203	Total	C	N	O	S	0	0	1
			1584	1009	298	275	2			
5	2F	203	Total	C	N	O	S	0	0	1
			1580	1007	297	274	2			

- Molecule 6 is a protein called 50S ribosomal protein L5.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
6	1G	181	Total	C	N	O	S	0	0	0
			1423	913	253	253	4			
6	2G	181	Total	C	N	O	S	0	0	0
			1428	913	258	253	4			

- Molecule 7 is a protein called 50S ribosomal protein L6.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
7	1H	174	Total	C	N	O	S	0	0	0
			1330	845	248	236	1			
7	2H	174	Total	C	N	O	S	0	0	0
			1330	845	248	236	1			

- Molecule 8 is a protein called 50S ribosomal protein L9.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
8	1I	146	Total	C	N	O	S	0	0	0
			1097	701	191	204	1			
8	2I	146	Total	C	N	O	S	0	0	0
			1064	681	186	196	1			

- Molecule 9 is a protein called 50S ribosomal protein L13.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
9	1N	140	Total	C	N	O	S	0	0	0
			1117	719	207	187	4			
9	2N	140	Total	C	N	O	S	0	0	0
			1117	719	207	187	4			

- Molecule 10 is a protein called 50S ribosomal protein L14.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
10	1O	122	Total	C	N	O	S	0	0	0
			933	588	171	170	4			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
10	2O	122	Total	C	N	O	S	0	0	0
			933	588	171	170	4			

- Molecule 11 is a protein called 50S ribosomal protein L15.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
11	1P	149	Total	C	N	O	S	0	0	0
			1135	706	230	196	3			
11	2P	149	Total	C	N	O	S	0	0	0
			1135	706	230	196	3			

- Molecule 12 is a protein called 50S ribosomal protein L16.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
12	1Q	141	Total	C	N	O	S	0	0	0
			1122	715	212	188	7			
12	2Q	141	Total	C	N	O	S	0	0	0
			1122	715	212	188	7			

- Molecule 13 is a protein called 50S ribosomal protein L17.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
13	1R	118	Total	C	N	O	S	0	0	0
			968	604	203	160	1			
13	2R	118	Total	C	N	O	S	0	0	0
			968	604	203	160	1			

- Molecule 14 is a protein called 50S ribosomal protein L18.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
14	1S	110	Total	C	N	O	0	0	0
			873	550	174	149			
14	2S	110	Total	C	N	O	0	0	0
			870	549	173	148			

- Molecule 15 is a protein called 50S ribosomal protein L19.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
15	1T	131	Total	C	N	O	S	0	0	0
			1091	680	225	185	1			
15	2T	131	Total	C	N	O	S	0	0	0
			1083	675	224	183	1			

- Molecule 16 is a protein called 50S ribosomal protein L20.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
16	1U	116	Total	C	N	O	S	0	0	0
			959	608	201	149	1			
16	2U	116	Total	C	N	O	S	0	0	0
			959	608	201	149	1			

- Molecule 17 is a protein called 50S ribosomal protein L21.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
17	1V	101	Total	C	N	O	S	0	0	0
			771	495	140	135	1			
17	2V	101	Total	C	N	O	S	0	0	0
			771	495	140	135	1			

- Molecule 18 is a protein called 50S ribosomal protein L22.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
18	1W	112	Total	C	N	O	S	0	0	0
			886	557	174	153	2			
18	2W	112	Total	C	N	O	S	0	0	0
			886	557	174	153	2			

- Molecule 19 is a protein called 50S ribosomal protein L23.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
19	1X	95	Total	C	N	O	S	0	0	0
			750	488	135	126	1			
19	2X	95	Total	C	N	O	S	0	0	0
			750	488	135	126	1			

- Molecule 20 is a protein called 50S ribosomal protein L24.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
20	1Y	107	Total	C	N	O	S	0	0	0
			806	517	152	131	6			
20	2Y	107	Total	C	N	O	S	0	0	0
			806	517	152	131	6			

- Molecule 21 is a protein called 50S ribosomal protein L25.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
21	1Z	154	Total	C	N	O	S	0	0	0
			1240	795	222	220	3			
21	2Z	160	Total	C	N	O	S	0	0	0
			1271	814	228	227	2			

- Molecule 22 is a protein called 50S ribosomal protein L27.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
22	10	83	Total	C	N	O	S	0	0	0
			653	404	139	109	1			
22	20	83	Total	C	N	O	S	0	0	0
			653	404	139	109	1			

- Molecule 23 is a protein called 50S ribosomal protein L28.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
23	11	97	Total	C	N	O	S	0	0	0
			755	475	148	131	1			
23	21	97	Total	C	N	O	S	0	0	0
			755	475	148	131	1			

- Molecule 24 is a protein called 50S ribosomal protein L29.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
24	12	70	Total	C	N	O	S	0	0	0
			588	365	118	103	2			
24	22	70	Total	C	N	O	S	0	0	0
			588	365	118	103	2			

- Molecule 25 is a protein called 50S ribosomal protein L30.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
25	13	59	Total	C	N	O	0	0	0
			469	298	90	81			
25	23	59	Total	C	N	O	0	0	0
			464	296	90	78			

- Molecule 26 is a protein called 50S ribosomal protein L31.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
26	14	69	Total	C	N	O	S	0	0	0
			552	349	99	99	5			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
26	24	69	Total	C	N	O	S	0	0	0
			532	339	97	91	5			

- Molecule 27 is a protein called 50S ribosomal protein L32.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
27	15	59	Total	C	N	O	S	0	0	0
			455	285	89	76	5			
27	25	59	Total	C	N	O	S	0	0	0
			455	285	89	76	5			

- Molecule 28 is a protein called 50S ribosomal protein L33.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
28	16	53	Total	C	N	O	S	0	0	0
			453	281	91	77	4			
28	26	53	Total	C	N	O	S	0	0	0
			449	279	91	75	4			

- Molecule 29 is a protein called 50S ribosomal protein L34.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
29	17	48	Total	C	N	O	S	0	0	0
			418	257	104	55	2			
29	27	48	Total	C	N	O	S	0	0	0
			418	257	104	55	2			

- Molecule 30 is a protein called 50S ribosomal protein L35.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
30	18	64	Total	C	N	O	S	0	0	0
			517	331	102	82	2			
30	28	64	Total	C	N	O	S	0	0	0
			517	331	102	82	2			

- Molecule 31 is a protein called 50S ribosomal protein L36.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
31	19	37	Total	C	N	O	S	0	0	0
			307	188	68	47	4			
31	29	37	Total	C	N	O	S	0	0	0
			307	188	68	47	4			

- Molecule 32 is a RNA chain called 16S Ribosomal RNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
32	1a	1500	Total	C	N	O	P	0	0	0
			32246	14358	5975	10413	1500			

- Molecule 33 is a protein called 30S ribosomal protein S2.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
33	1b	231	Total	C	N	O	S	0	0	0
			1846	1179	331	331	5			
33	2b	231	Total	C	N	O	S	0	0	0
			1825	1167	326	327	5			

- Molecule 34 is a protein called 30S ribosomal protein S3.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
34	1c	206	Total	C	N	O	S	0	0	0
			1548	973	301	273	1			
34	2c	206	Total	C	N	O	S	0	0	0
			1542	968	300	273	1			

- Molecule 35 is a protein called 30S ribosomal protein S4.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
35	1d	208	Total	C	N	O	S	0	0	0
			1655	1038	326	284	7			
35	2d	208	Total	C	N	O	S	0	0	0
			1674	1050	333	284	7			

- Molecule 36 is a protein called 30S ribosomal protein S5.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
36	1e	148	Total	C	N	O	S	0	0	0
			1129	714	213	198	4			
36	2e	148	Total	C	N	O	S	0	0	0
			1133	716	214	199	4			

- Molecule 37 is a protein called 30S ribosomal protein S6.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
37	1f	100	Total	C	N	O	S	0	0	0
			810	514	144	149	3			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
37	2f	100	Total	C	N	O	S	0	0	0
			816	516	146	151	3			

- Molecule 38 is a protein called 30S ribosomal protein S7.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
38	1g	155	Total	C	N	O	S	0	0	0
			1231	766	243	216	6			
38	2g	155	Total	C	N	O	S	0	0	0
			1235	769	244	216	6			

- Molecule 39 is a protein called 30S ribosomal protein S8.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
39	1h	137	Total	C	N	O	S	0	0	0
			1088	689	206	191	2			
39	2h	137	Total	C	N	O	S	0	0	0
			1088	689	206	191	2			

- Molecule 40 is a protein called 30S ribosomal protein S9.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
40	1i	127	Total	C	N	O	0	0	0
			983	623	193	167			
40	2i	127	Total	C	N	O	0	0	0
			978	619	190	169			

- Molecule 41 is a protein called 30S ribosomal protein S10.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
41	1j	97	Total	C	N	O	0	0	0
			709	440	138	131			
41	2j	96	Total	C	N	O	0	0	0
			714	445	138	131			

- Molecule 42 is a protein called 30S ribosomal protein S11.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
42	1k	114	Total	C	N	O	S	0	0	0
			829	516	155	155	3			
42	2k	114	Total	C	N	O	S	0	0	0
			833	519	156	155	3			

- Molecule 43 is a protein called 30S ribosomal protein S12.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
43	1l	122	Total	C	N	O	S	0	0	0
			932	586	185	159	2			
43	2l	122	Total	C	N	O	S	0	0	0
			932	586	185	159	2			

- Molecule 44 is a protein called 30S ribosomal protein S13.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
44	1m	123	Total	C	N	O	S	0	0	0
			958	592	198	166	2			
44	2m	122	Total	C	N	O	S	0	0	0
			950	586	197	165	2			

- Molecule 45 is a protein called 30S ribosomal protein S14 type Z.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
45	1n	60	Total	C	N	O	S	0	0	0
			492	312	104	72	4			
45	2n	60	Total	C	N	O	S	0	0	0
			492	312	104	72	4			

- Molecule 46 is a protein called 30S ribosomal protein S15.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
46	1o	88	Total	C	N	O	S	0	0	0
			728	456	144	126	2			
46	2o	88	Total	C	N	O	S	0	0	0
			728	456	144	126	2			

- Molecule 47 is a protein called 30S ribosomal protein S16.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
47	1p	82	Total	C	N	O	S	0	0	0
			681	433	134	113	1			
47	2p	82	Total	C	N	O	S	0	0	0
			677	430	133	113	1			

- Molecule 48 is a protein called 30S ribosomal protein S17.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
48	1q	99	Total	C	N	O	S	0	0	0
			823	528	151	142	2			
48	2q	99	Total	C	N	O	S	0	0	0
			823	528	151	142	2			

- Molecule 49 is a protein called 30S ribosomal protein S18.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
49	1r	68	Total	C	N	O		0	0	0
			555	355	108	92				
49	2r	68	Total	C	N	O		0	0	0
			555	355	108	92				

- Molecule 50 is a protein called 30S ribosomal protein S19.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
50	1s	83	Total	C	N	O	S	0	0	0
			652	417	120	113	2			
50	2s	83	Total	C	N	O	S	0	0	0
			646	412	119	113	2			

- Molecule 51 is a protein called 30S ribosomal protein S20.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
51	1t	96	Total	C	N	O	S	0	0	0
			728	446	156	124	2			
51	2t	96	Total	C	N	O	S	0	0	0
			727	446	155	124	2			

- Molecule 52 is a protein called 30S ribosomal protein Thx.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
52	1u	23	Total	C	N	O		0	0	0
			199	122	48	29				
52	2u	23	Total	C	N	O		0	0	0
			199	122	48	29				

- Molecule 53 is a RNA chain called mRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
53	1v	13	Total	C	N	O	P	0	0	0
			277	125	51	88	13			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
53	2v	13	Total	C	N	O	P	0	0	0
			277	125	51	88	13			

- Molecule 54 is a RNA chain called tRNA, A-site and E-site tRNAs.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
54	1w	74	Total	C	N	O	P	S	0	0
			1592	713	285	518	74	2		
54	2w	72	Total	C	N	O	P	S	0	0
			1544	690	278	502	72	2		

- Molecule 55 is a RNA chain called P-site tRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
55	1x	76	Total	C	N	O	P	S	0	0
			1625	725	294	529	76	1		
55	2x	76	Total	C	N	O	P	S	0	0
			1625	725	294	529	76	1		

- Molecule 56 is a RNA chain called tRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
56	1y	74	Total	C	N	O	P	S	0	0
			1585	707	285	518	74	1		
56	2y	73	Total	C	N	O	P	S	0	0
			1565	698	283	510	73	1		

- Molecule 57 is a RNA chain called tRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
57	2a	1503	Total	C	N	O	P	0	0	0
			32327	14396	5990	10438	1503			

- Molecule 58 is a protein called Klebsazolicin.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
58	A	18	Total	C	N	O	S	0	0	0
			115	64	23	25	3			
58	B	18	Total	C	N	O	S	0	0	0
			115	64	23	25	3			

- Molecule 59 is MAGNESIUM ION (three-letter code: MG) (formula: Mg).

Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
59	2E	7	Total Mg 7 7	0	0
59	17	2	Total Mg 2 2	0	0
59	2d	1	Total Mg 1 1	0	0
59	1T	2	Total Mg 2 2	0	0
59	1N	9	Total Mg 9 9	0	0
59	20	3	Total Mg 3 3	0	0
59	18	3	Total Mg 3 3	0	0
59	2l	2	Total Mg 2 2	0	0
59	1Y	3	Total Mg 3 3	0	0
59	13	3	Total Mg 3 3	0	0
59	1f	3	Total Mg 3 3	0	0
59	A	1	Total Mg 1 1	0	0
59	1P	4	Total Mg 4 4	0	0
59	2B	22	Total Mg 22 22	0	0
59	2a	239	Total Mg 239 239	0	0
59	1k	1	Total Mg 1 1	0	0
59	1E	14	Total Mg 14 14	0	0
59	1b	2	Total Mg 2 2	0	0
59	25	3	Total Mg 3 3	0	0
59	2F	4	Total Mg 4 4	0	0
59	16	3	Total Mg 3 3	0	0
59	28	2	Total Mg 2 2	0	0

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Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
59	2e	1	Total Mg 1 1	0	0
59	1W	5	Total Mg 5 5	0	0
59	1A	1229	Total Mg 1229 1229	0	0
59	1t	1	Total Mg 1 1	0	0
59	2p	1	Total Mg 1 1	0	0
59	1n	2	Total Mg 2 2	0	0
59	2P	1	Total Mg 1 1	0	0
59	1X	5	Total Mg 5 5	0	0
59	12	2	Total Mg 2 2	0	0
59	1y	5	Total Mg 5 5	0	0
59	1S	3	Total Mg 3 3	0	0
59	1p	1	Total Mg 1 1	0	0
59	2T	3	Total Mg 3 3	0	0
59	1D	13	Total Mg 13 13	0	0
59	2N	1	Total Mg 1 1	0	0
59	1e	1	Total Mg 1 1	0	0
59	2G	1	Total Mg 1 1	0	0
59	1I	1	Total Mg 1 1	0	0
59	2f	2	Total Mg 2 2	0	0
59	1V	2	Total Mg 2 2	0	0
59	2X	2	Total Mg 2 2	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	1w	12	Total 12	Mg 12	0	0
59	1a	278	Total 278	Mg 278	0	0
59	2Q	4	Total 4	Mg 4	0	0
59	15	2	Total 2	Mg 2	0	0
59	1x	17	Total 17	Mg 17	0	0
59	2j	2	Total 2	Mg 2	0	0
59	1R	3	Total 3	Mg 3	0	0
59	1s	1	Total 1	Mg 1	0	0
59	B	1	Total 1	Mg 1	0	0
59	2v	4	Total 4	Mg 4	0	0
59	2U	4	Total 4	Mg 4	0	0
59	1G	5	Total 5	Mg 5	0	0
59	2O	2	Total 2	Mg 2	0	0
59	1l	3	Total 3	Mg 3	0	0
59	1d	1	Total 1	Mg 1	0	0
59	2r	1	Total 1	Mg 1	0	0
59	2g	1	Total 1	Mg 1	0	0
59	1v	1	Total 1	Mg 1	0	0
59	2x	4	Total 4	Mg 4	0	0
59	2R	3	Total 3	Mg 3	0	0
59	1Z	4	Total 4	Mg 4	0	0

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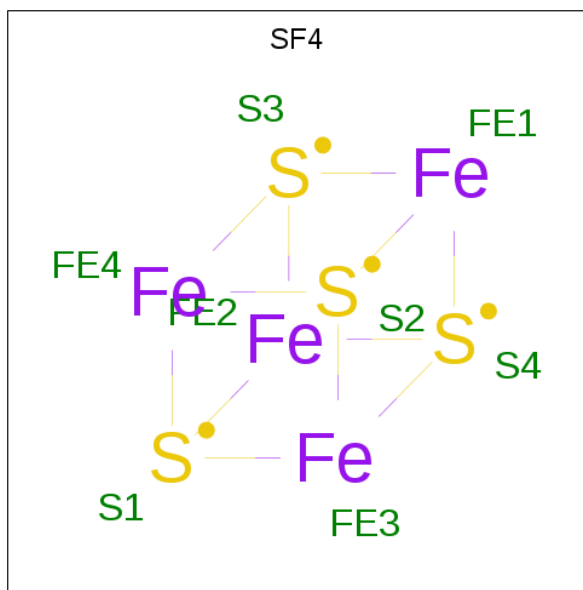
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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	2D	5	Total 5	Mg 5	0	0
59	2q	5	Total 5	Mg 5	0	0
59	2k	1	Total 1	Mg 1	0	0
59	1U	5	Total 5	Mg 5	0	0
59	1O	6	Total 6	Mg 6	0	0
59	27	1	Total 1	Mg 1	0	0
59	19	1	Total 1	Mg 1	0	0
59	1l	3	Total 3	Mg 3	0	0
59	2V	1	Total 1	Mg 1	0	0
59	1F	10	Total 10	Mg 10	0	0
59	10	8	Total 8	Mg 8	0	0
59	2t	1	Total 1	Mg 1	0	0
59	1Q	6	Total 6	Mg 6	0	0
59	2A	848	Total 848	Mg 848	0	0
59	23	2	Total 2	Mg 2	0	0
59	2Z	1	Total 1	Mg 1	0	0
59	1B	38	Total 38	Mg 38	0	0
59	2y	7	Total 7	Mg 7	0	0
59	2w	7	Total 7	Mg 7	0	0
59	2S	1	Total 1	Mg 1	0	0

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

Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
60	1Y	1	Total Zn 1 1	0	0
60	14	1	Total Zn 1 1	0	0
60	1n	1	Total Zn 1 1	0	0
60	15	1	Total Zn 1 1	0	0
60	29	1	Total Zn 1 1	0	0
60	19	1	Total Zn 1 1	0	0
60	26	1	Total Zn 1 1	0	0
60	25	1	Total Zn 1 1	0	0
60	24	1	Total Zn 1 1	0	0
60	2n	1	Total Zn 1 1	0	0
60	2Y	1	Total Zn 1 1	0	0
60	16	1	Total Zn 1 1	0	0

- Molecule 61 is IRON/SULFUR CLUSTER (three-letter code: SF4) (formula: Fe₄S₄).



Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
61	1d	1	Total	Fe	S	0	0
			8	4	4		
61	2d	1	Total	Fe	S	0	0
			8	4	4		

- Molecule 62 is POTASSIUM ION (three-letter code: K) (formula: K).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
62	2A	1	Total	K	0	0
			1	1		

- Molecule 63 is water.

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
63	1A	2636	Total	O	0	0
			2636	2636		
63	1B	71	Total	O	0	0
			71	71		
63	1D	31	Total	O	0	0
			31	31		
63	1E	35	Total	O	0	0
			35	35		
63	1F	18	Total	O	0	0
			18	18		
63	1G	6	Total	O	0	0
			6	6		
63	1H	3	Total	O	0	0
			3	3		
63	1I	2	Total	O	0	0
			2	2		
63	1N	10	Total	O	0	0
			10	10		
63	1O	8	Total	O	0	0
			8	8		
63	1P	23	Total	O	0	0
			23	23		
63	1Q	16	Total	O	0	0
			16	16		
63	1R	14	Total	O	0	0
			14	14		
63	1S	4	Total	O	0	0
			4	4		
63	1T	9	Total	O	0	0
			9	9		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
63	1U	13	Total 13	O 13	0	0
63	1V	9	Total 9	O 9	0	0
63	1W	11	Total 11	O 11	0	0
63	1X	8	Total 8	O 8	0	0
63	1Y	7	Total 7	O 7	0	0
63	1Z	2	Total 2	O 2	0	0
63	10	12	Total 12	O 12	0	0
63	11	13	Total 13	O 13	0	0
63	12	4	Total 4	O 4	0	0
63	13	5	Total 5	O 5	0	0
63	14	1	Total 1	O 1	0	0
63	15	5	Total 5	O 5	0	0
63	16	5	Total 5	O 5	0	0
63	17	9	Total 9	O 9	0	0
63	18	13	Total 13	O 13	0	0
63	19	1	Total 1	O 1	0	0
63	1a	523	Total 523	O 523	0	0
63	1b	1	Total 1	O 1	0	0
63	1d	3	Total 3	O 3	0	0
63	1e	3	Total 3	O 3	0	0
63	1g	1	Total 1	O 1	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
63	1k	3	Total 3	O 3	0	0
63	1l	10	Total 10	O 10	0	0
63	1m	1	Total 1	O 1	0	0
63	1o	2	Total 2	O 2	0	0
63	1p	1	Total 1	O 1	0	0
63	1q	4	Total 4	O 4	0	0
63	1r	1	Total 1	O 1	0	0
63	1t	2	Total 2	O 2	0	0
63	1v	6	Total 6	O 6	0	0
63	1w	26	Total 26	O 26	0	0
63	1x	18	Total 18	O 18	0	0
63	1y	2	Total 2	O 2	0	0
63	2A	1602	Total 1602	O 1602	0	0
63	2B	28	Total 28	O 28	0	0
63	2D	29	Total 29	O 29	0	0
63	2E	15	Total 15	O 15	0	0
63	2F	20	Total 20	O 20	0	0
63	2I	4	Total 4	O 4	0	0
63	2N	2	Total 2	O 2	0	0
63	2P	15	Total 15	O 15	0	0
63	2Q	2	Total 2	O 2	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
63	2R	5	Total 5	O 5	0	0
63	2T	6	Total 6	O 6	0	0
63	2U	6	Total 6	O 6	0	0
63	2V	4	Total 4	O 4	0	0
63	2W	4	Total 4	O 4	0	0
63	2X	4	Total 4	O 4	0	0
63	2Y	1	Total 1	O 1	0	0
63	2Z	2	Total 2	O 2	0	0
63	20	5	Total 5	O 5	0	0
63	21	14	Total 14	O 14	0	0
63	22	1	Total 1	O 1	0	0
63	23	2	Total 2	O 2	0	0
63	25	4	Total 4	O 4	0	0
63	27	7	Total 7	O 7	0	0
63	28	7	Total 7	O 7	0	0
63	29	1	Total 1	O 1	0	0
63	2a	389	Total 389	O 389	0	0
63	2d	3	Total 3	O 3	0	0
63	2e	3	Total 3	O 3	0	0
63	2f	1	Total 1	O 1	0	0
63	2g	1	Total 1	O 1	0	0

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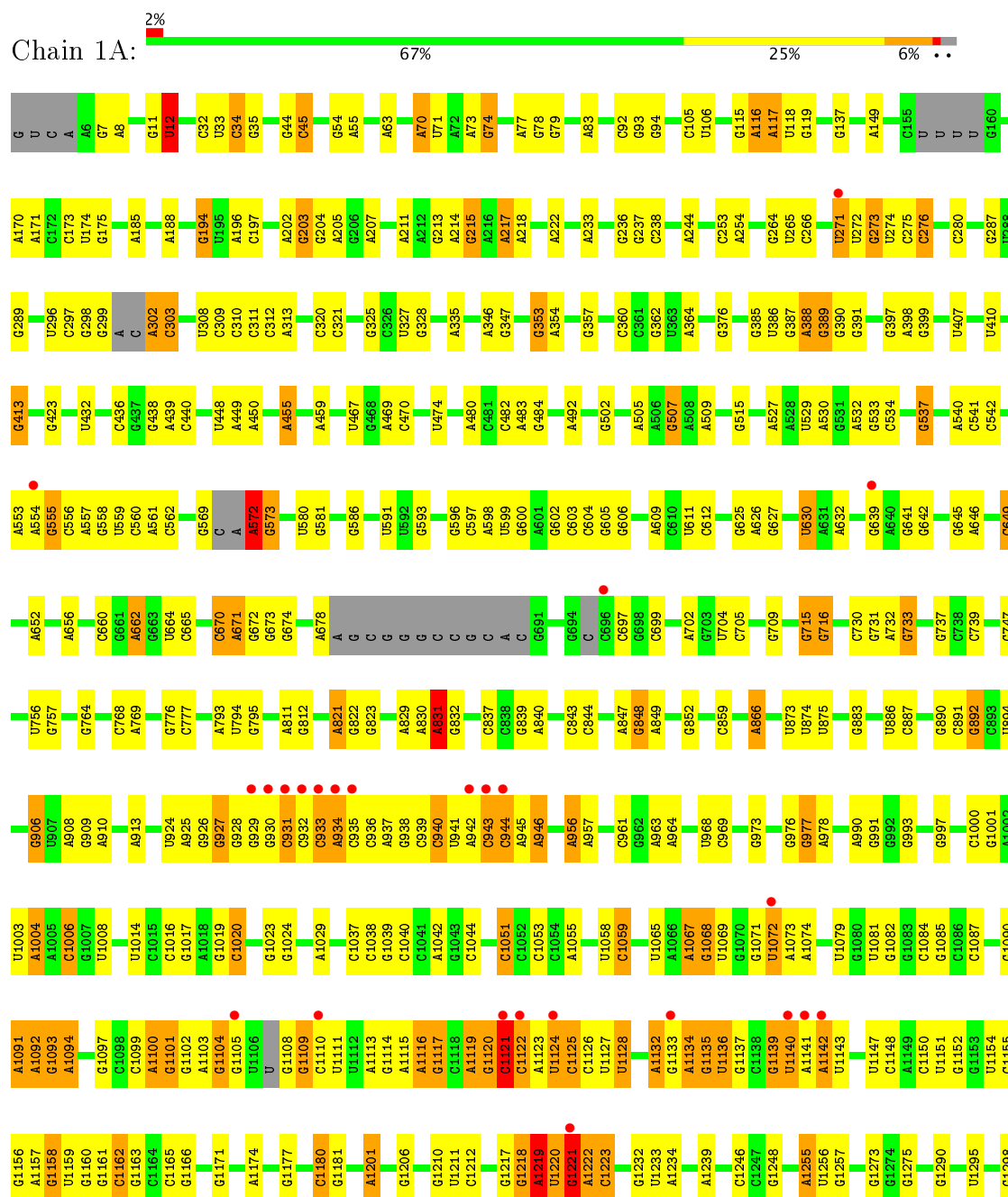
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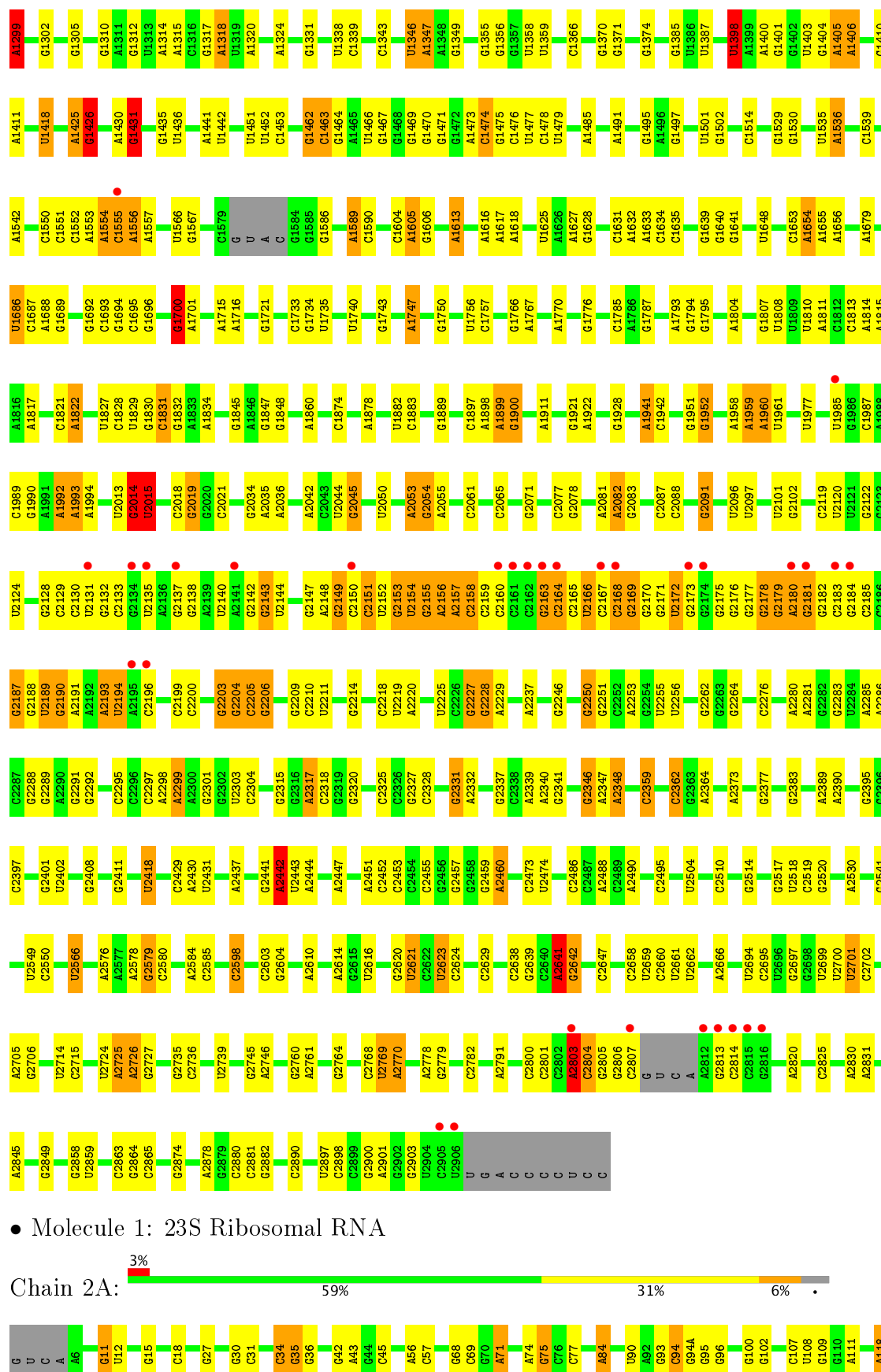
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
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			2	2		
63	2i	1	Total	O	0	0
			1	1		
63	2j	4	Total	O	0	0
			4	4		
63	2l	5	Total	O	0	0
			5	5		
63	2o	1	Total	O	0	0
			1	1		
63	2p	3	Total	O	0	0
			3	3		
63	2q	2	Total	O	0	0
			2	2		
63	2r	1	Total	O	0	0
			1	1		
63	2t	3	Total	O	0	0
			3	3		
63	2u	1	Total	O	0	0
			1	1		
63	2v	1	Total	O	0	0
			1	1		
63	2w	3	Total	O	0	0
			3	3		
63	2x	10	Total	O	0	0
			10	10		
63	2y	20	Total	O	0	0
			20	20		
63	A	4	Total	O	0	0
			4	4		
63	B	2	Total	O	0	0
			2	2		

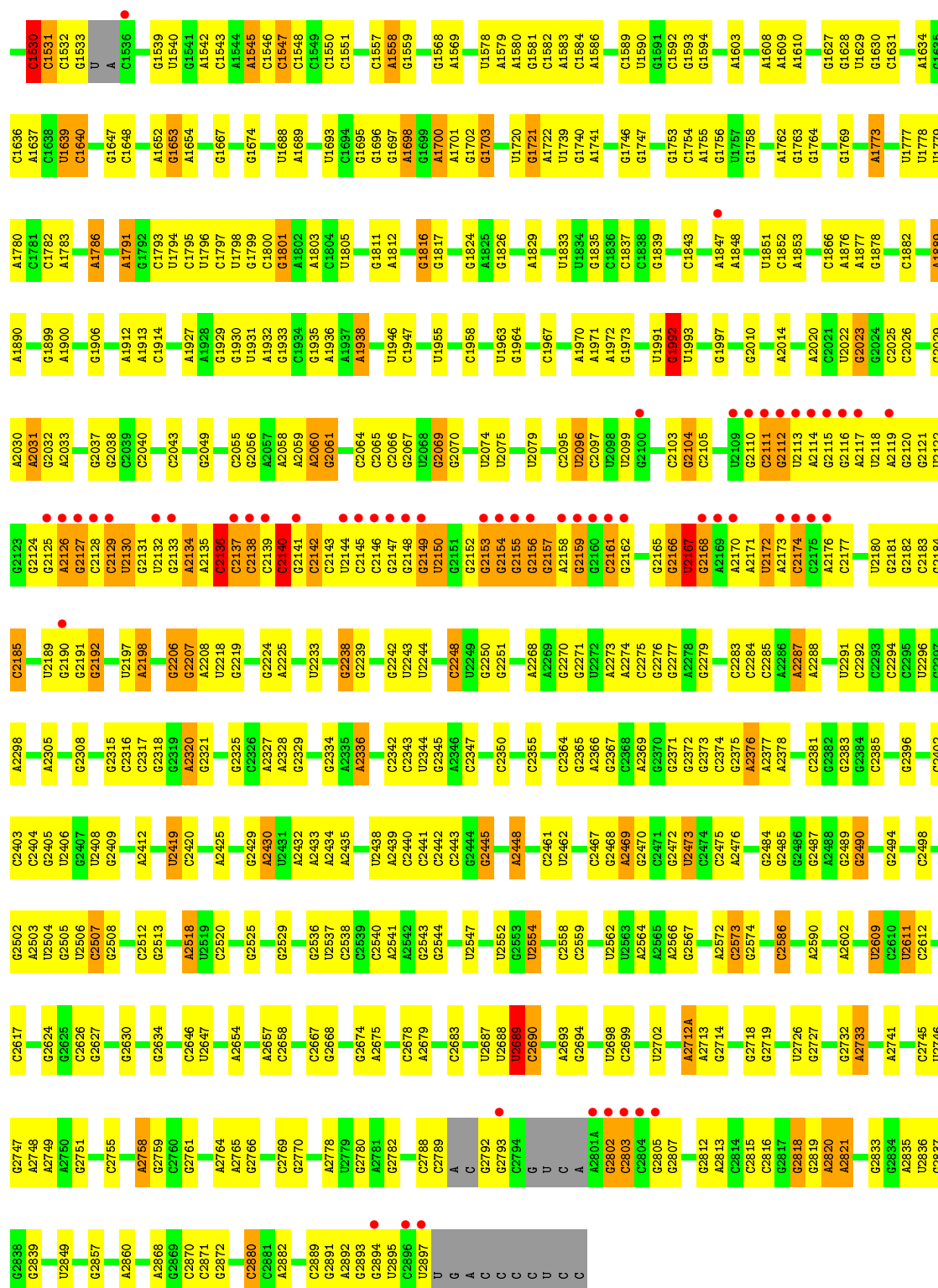
3 Residue-property plots

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

- Molecule 1: 23S Ribosomal RNA

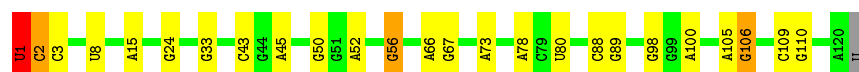




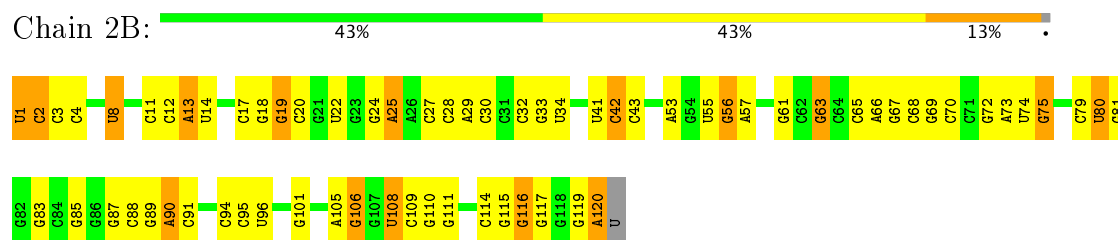


• Molecule 2: 5S Ribosomal RNA

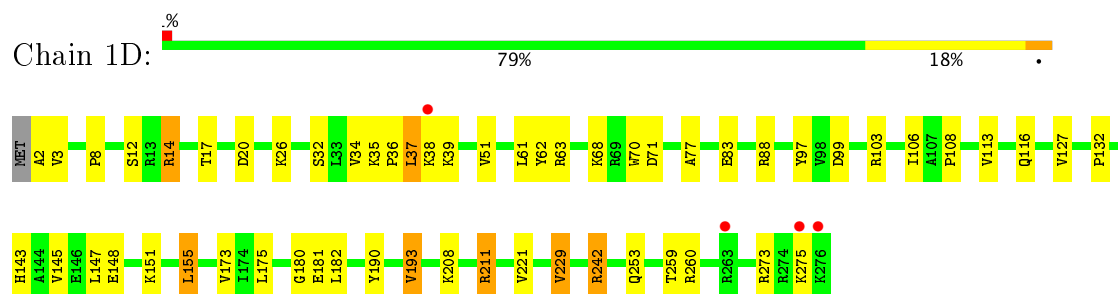
Chain 1B: 79% 17% ...



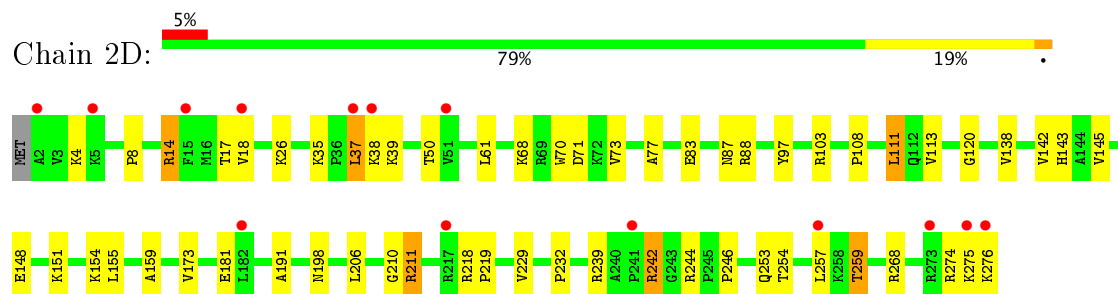
- Molecule 2: 5S Ribosomal RNA



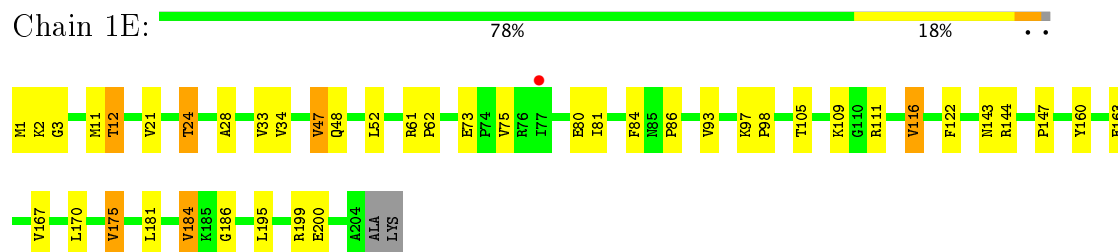
- Molecule 3: 50S ribosomal protein L2



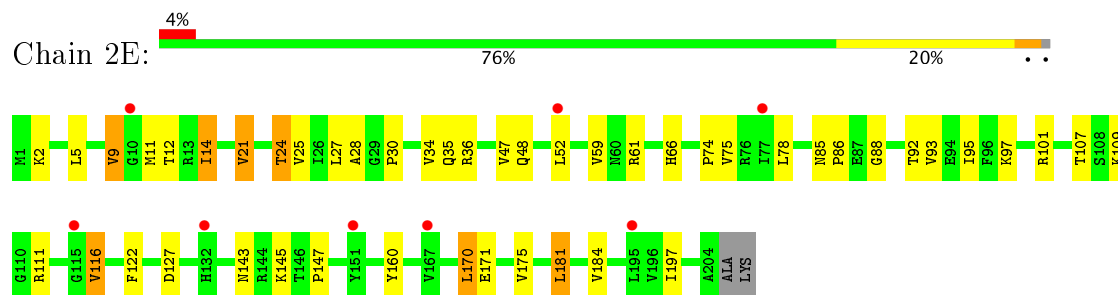
- Molecule 3: 50S ribosomal protein L2



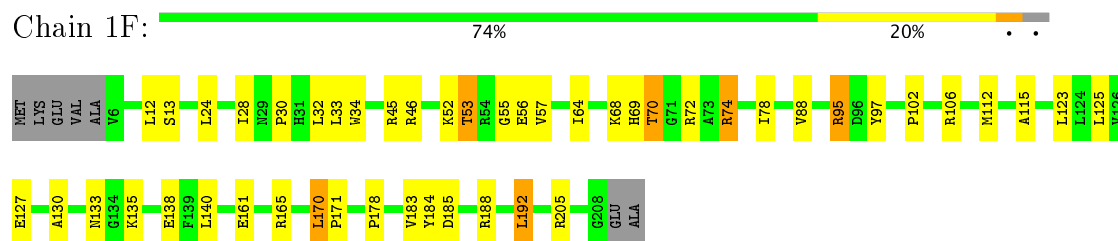
- Molecule 4: 50S ribosomal protein L3



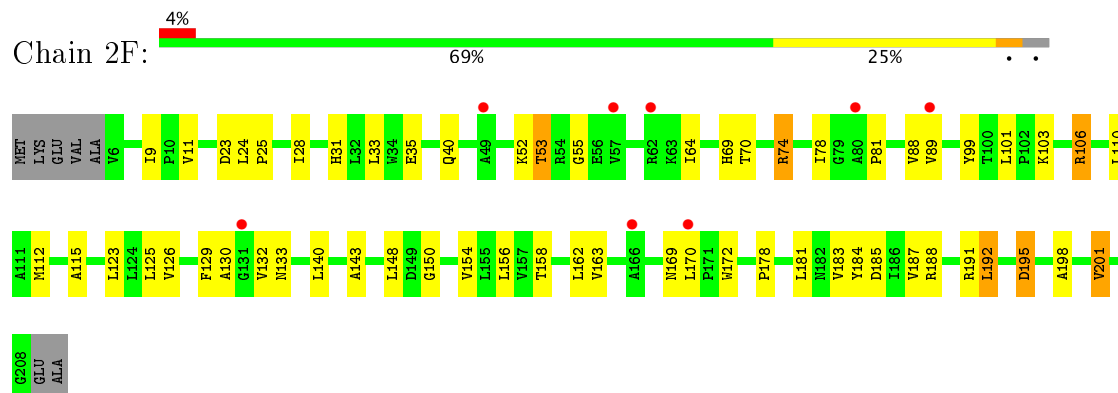
- Molecule 4: 50S ribosomal protein L3



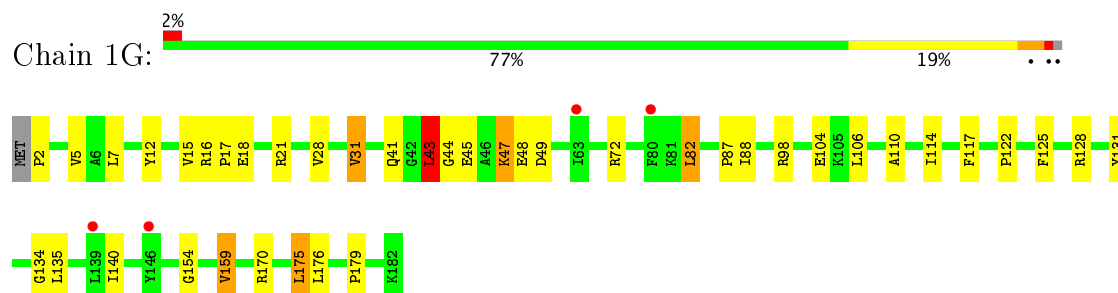
- Molecule 5: 50S ribosomal protein L4



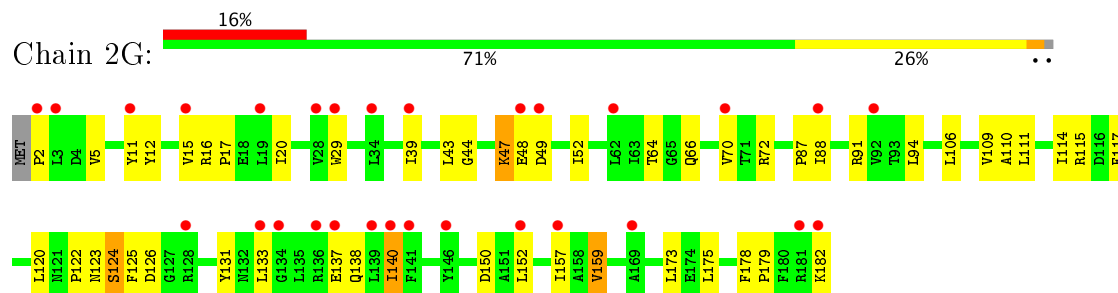
- Molecule 5: 50S ribosomal protein L4



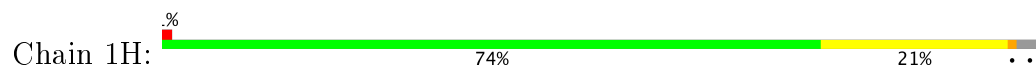
- Molecule 6: 50S ribosomal protein L5

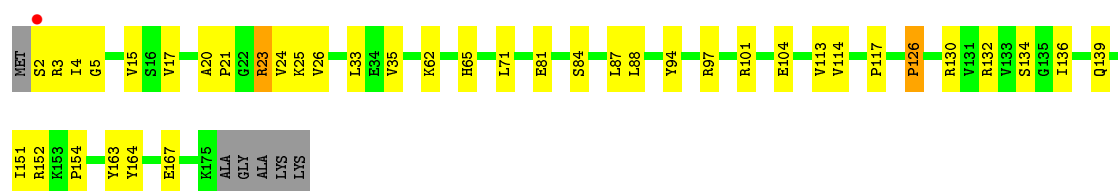


- Molecule 6: 50S ribosomal protein L5

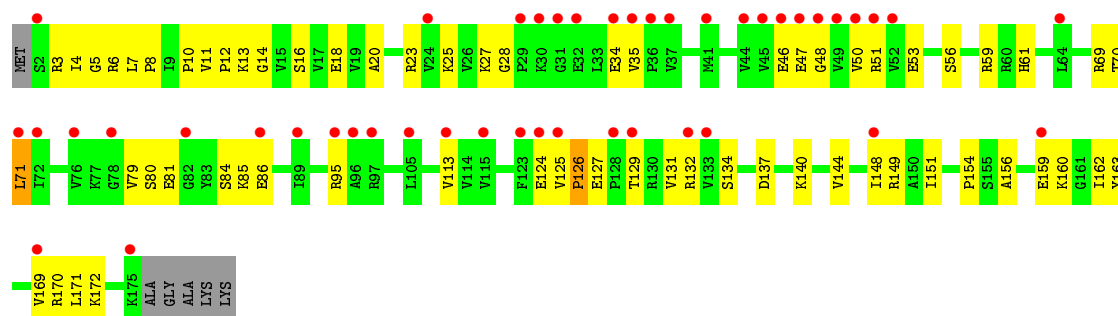


- Molecule 7: 50S ribosomal protein L6

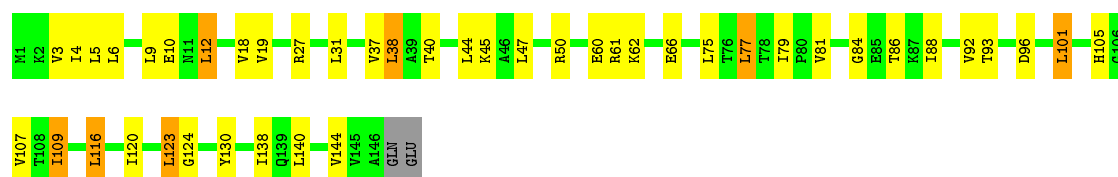




- Molecule 7: 50S ribosomal protein L6



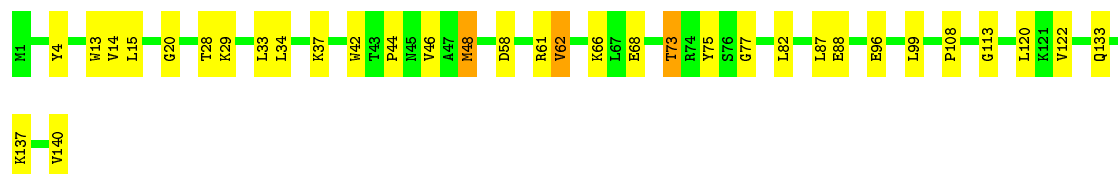
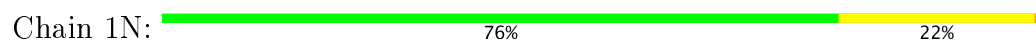
- Molecule 8: 50S ribosomal protein L9



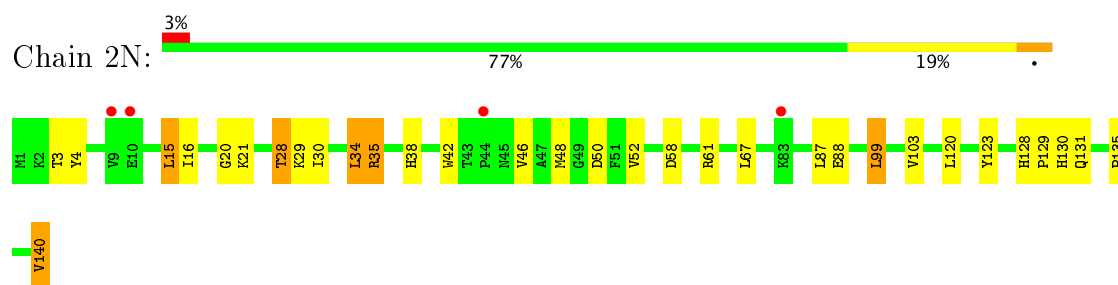
- Molecule 8: 50S ribosomal protein L9



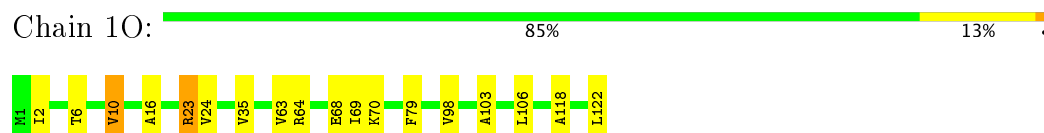
- Molecule 9: 50S ribosomal protein L13



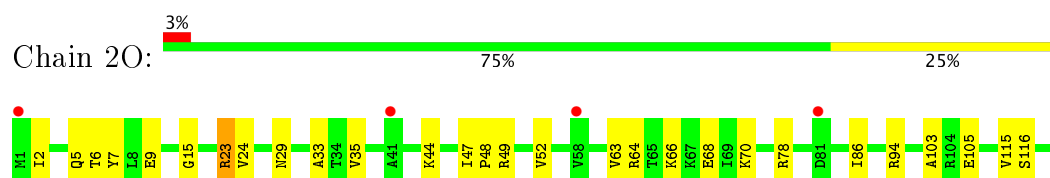
- Molecule 9: 50S ribosomal protein L13



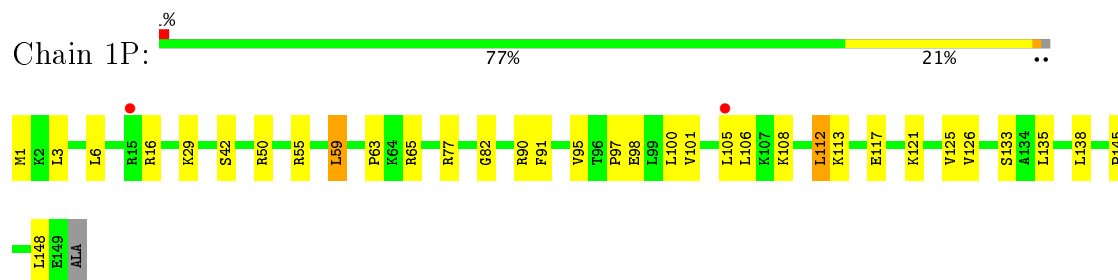
- Molecule 10: 50S ribosomal protein L14



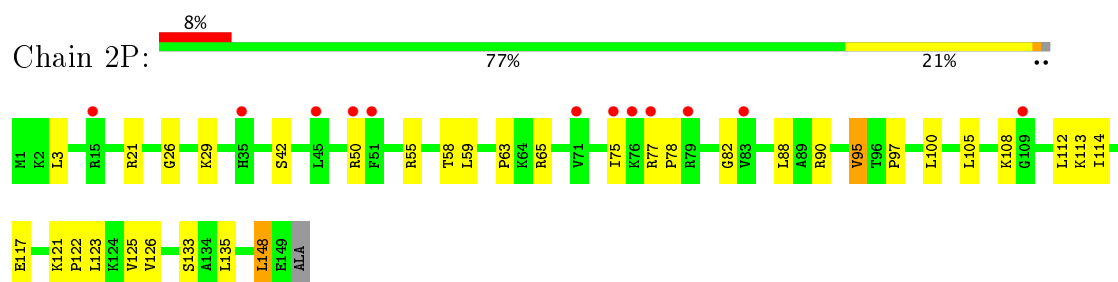
- Molecule 10: 50S ribosomal protein L14



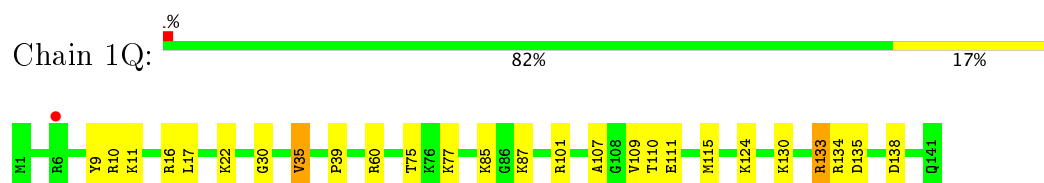
- Molecule 11: 50S ribosomal protein L15



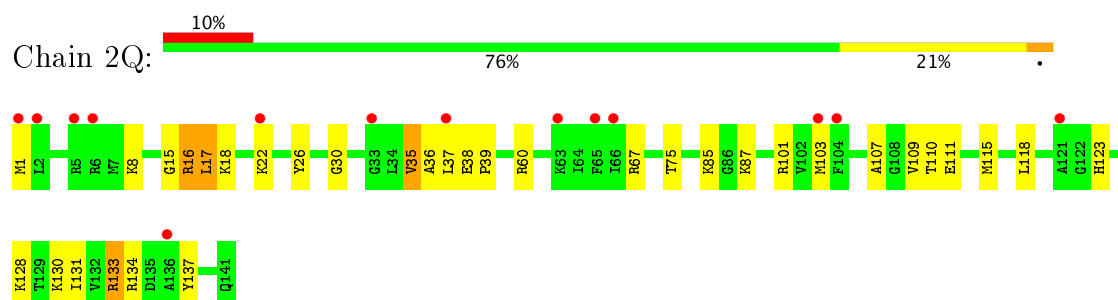
- Molecule 11: 50S ribosomal protein L15



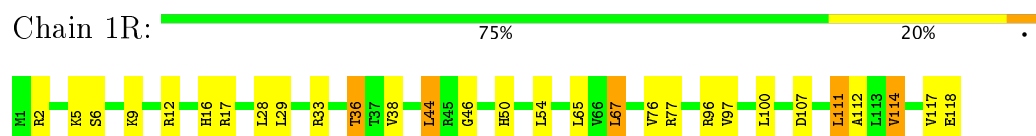
- Molecule 12: 50S ribosomal protein L16



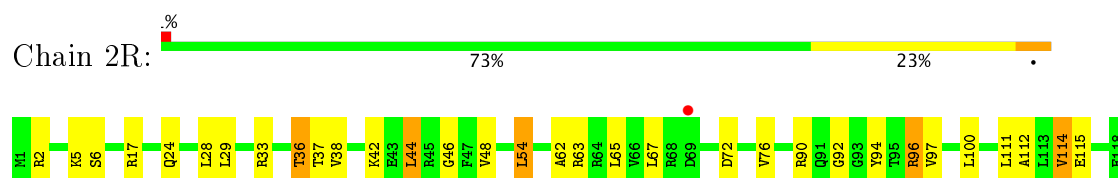
- Molecule 12: 50S ribosomal protein L16



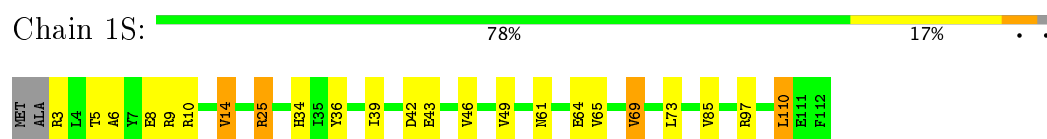
- Molecule 13: 50S ribosomal protein L17



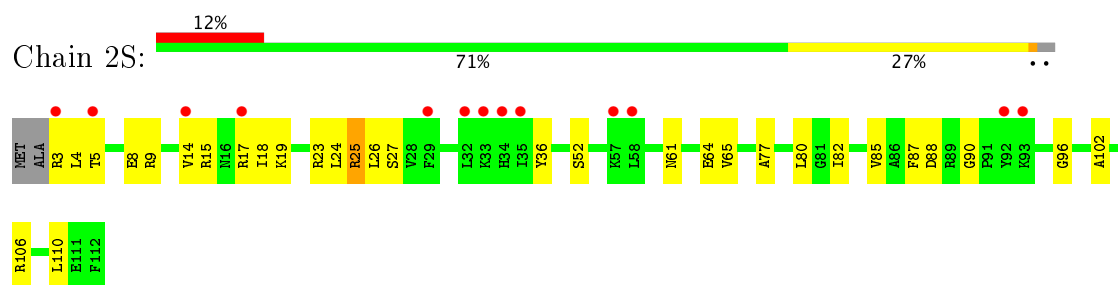
- Molecule 13: 50S ribosomal protein L17



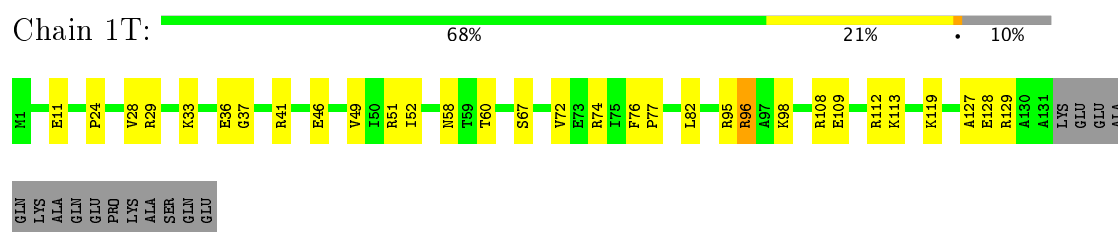
- Molecule 14: 50S ribosomal protein L18



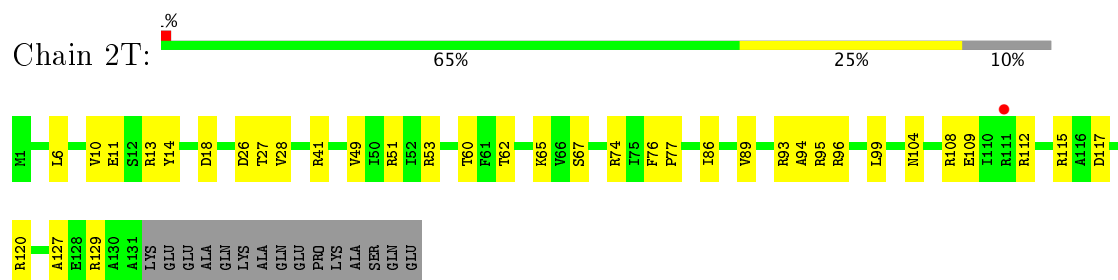
- Molecule 14: 50S ribosomal protein L18



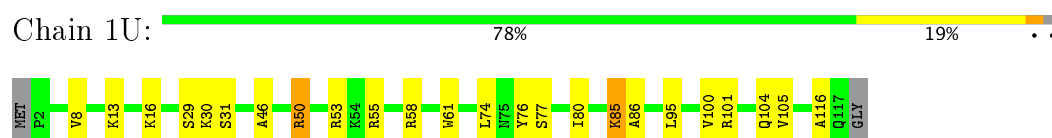
- Molecule 15: 50S ribosomal protein L19



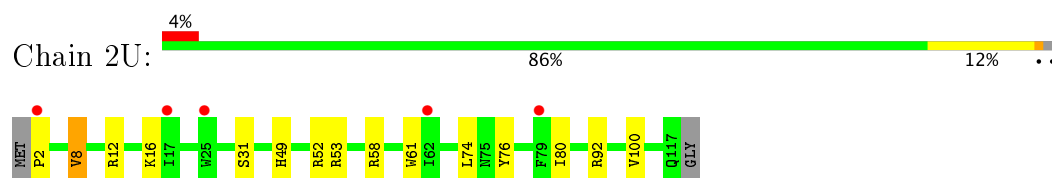
- Molecule 15: 50S ribosomal protein L19



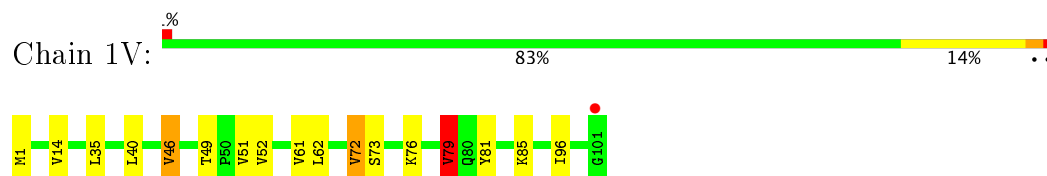
- Molecule 16: 50S ribosomal protein L20



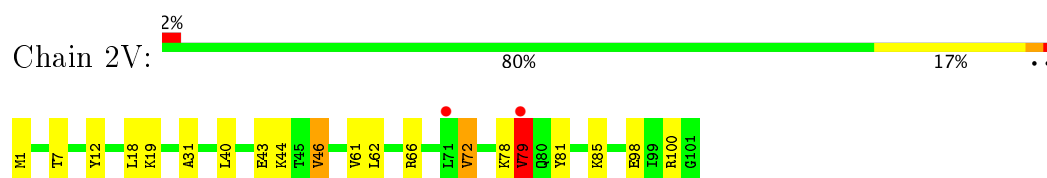
- Molecule 16: 50S ribosomal protein L20



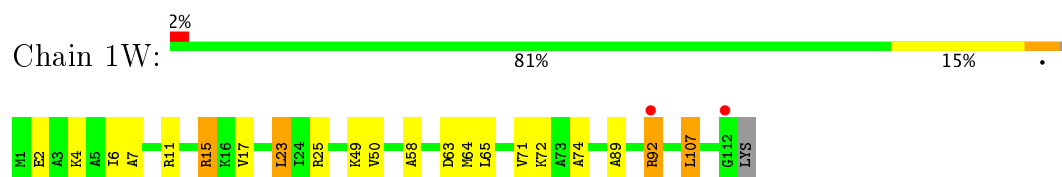
- Molecule 17: 50S ribosomal protein L21



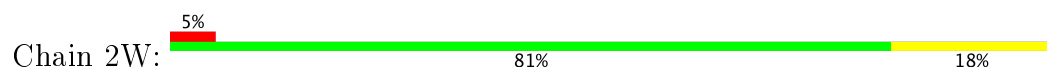
- Molecule 17: 50S ribosomal protein L21

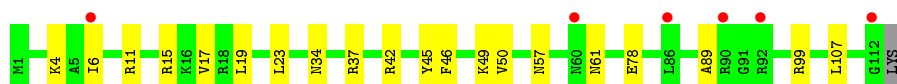


- Molecule 18: 50S ribosomal protein L22

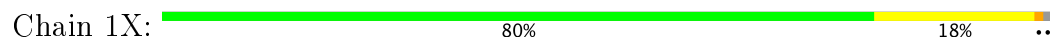


- Molecule 18: 50S ribosomal protein L22

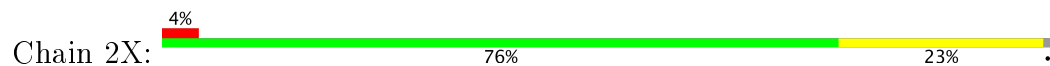




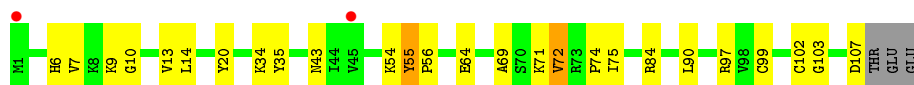
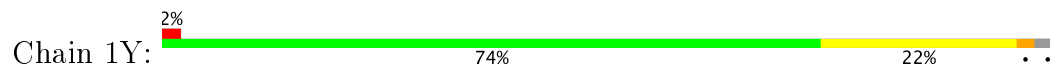
- Molecule 19: 50S ribosomal protein L23



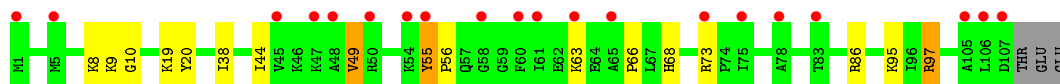
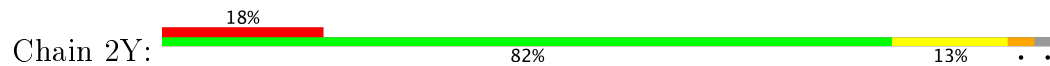
- Molecule 19: 50S ribosomal protein L23



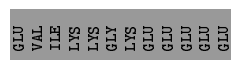
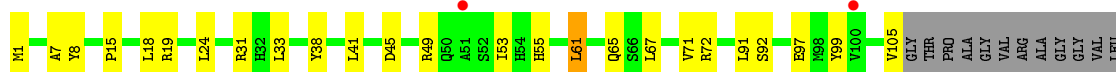
- Molecule 20: 50S ribosomal protein L24



- Molecule 20: 50S ribosomal protein L24

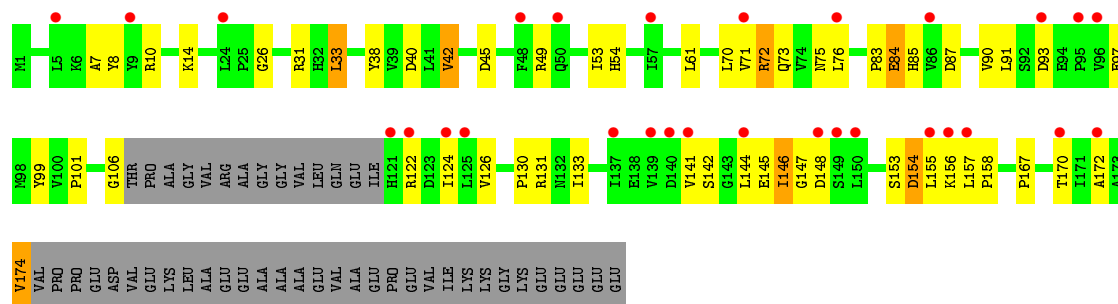


- Molecule 21: 50S ribosomal protein L25

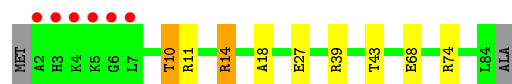
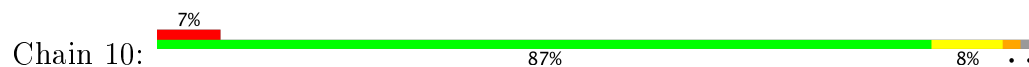


- Molecule 21: 50S ribosomal protein L25

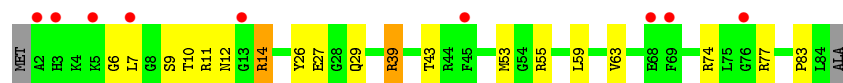
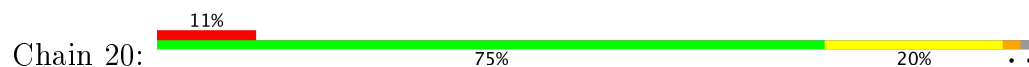




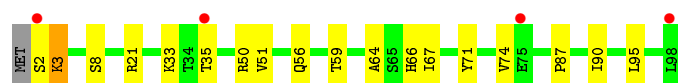
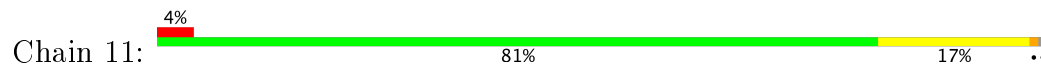
- Molecule 22: 50S ribosomal protein L27



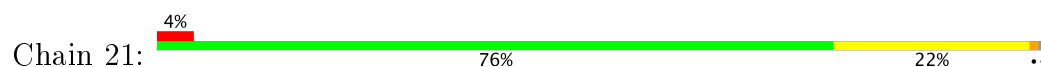
- Molecule 22: 50S ribosomal protein L27



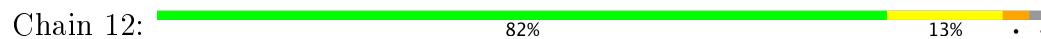
- Molecule 23: 50S ribosomal protein L28



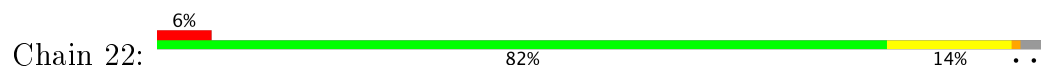
- Molecule 23: 50S ribosomal protein L28



- Molecule 24: 50S ribosomal protein L29

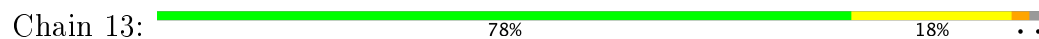


- Molecule 24: 50S ribosomal protein L29

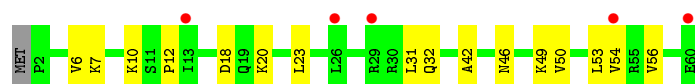




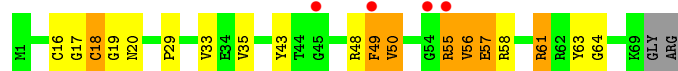
- Molecule 25: 50S ribosomal protein L30



- Molecule 25: 50S ribosomal protein L30



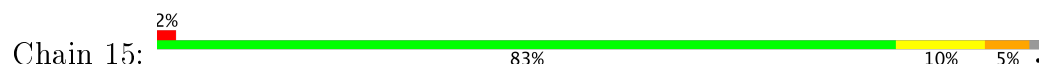
- Molecule 26: 50S ribosomal protein L31



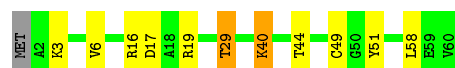
- Molecule 26: 50S ribosomal protein L31



- Molecule 27: 50S ribosomal protein L32



- Molecule 27: 50S ribosomal protein L32

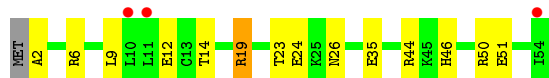
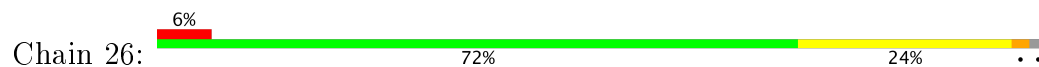


- Molecule 28: 50S ribosomal protein L33

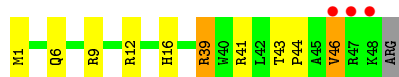
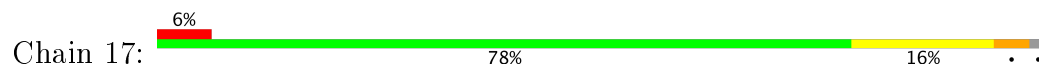




- Molecule 28: 50S ribosomal protein L33



- Molecule 29: 50S ribosomal protein L34



- Molecule 29: 50S ribosomal protein L34



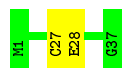
- Molecule 30: 50S ribosomal protein L35



- Molecule 30: 50S ribosomal protein L35

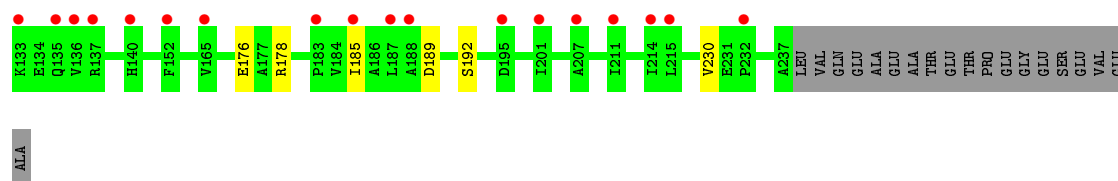


- Molecule 31: 50S ribosomal protein L36

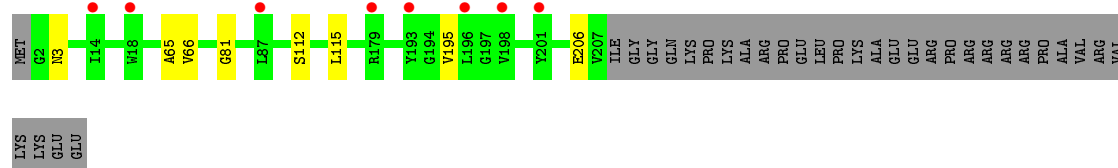
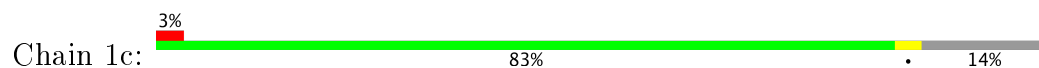


- Molecule 31: 50S ribosomal protein L36

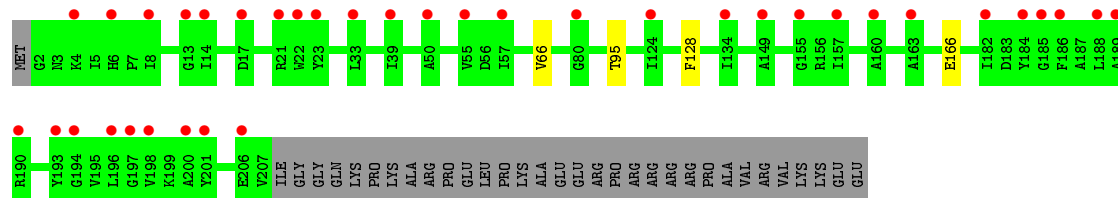
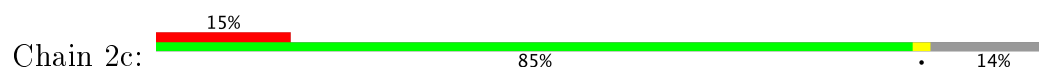




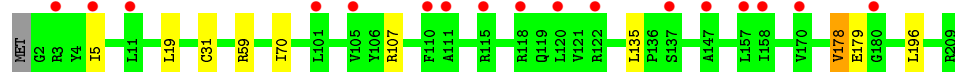
- Molecule 34: 30S ribosomal protein S3



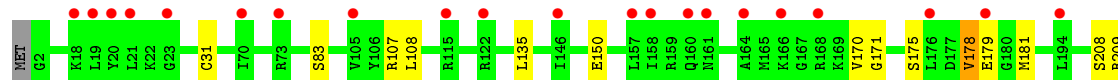
- Molecule 34: 30S ribosomal protein S3



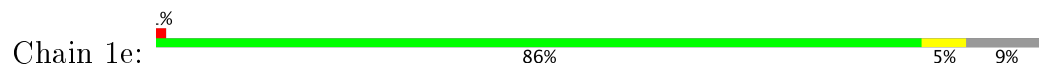
- Molecule 35: 30S ribosomal protein S4



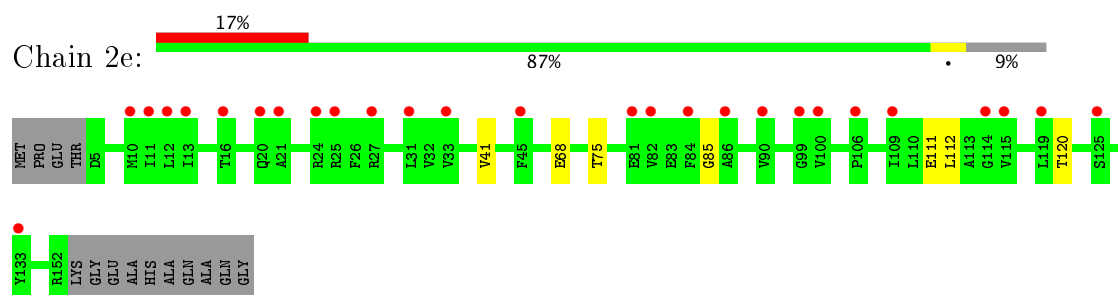
- Molecule 35: 30S ribosomal protein S4



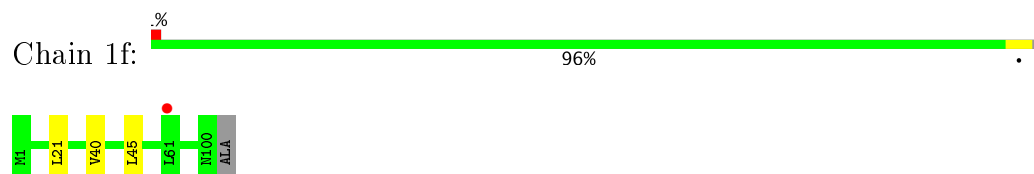
- Molecule 36: 30S ribosomal protein S5



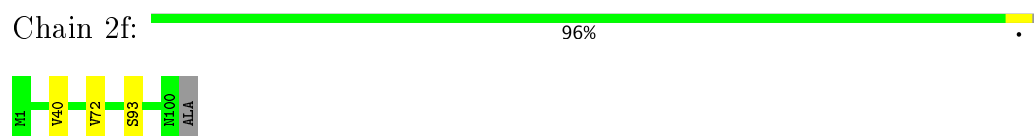
- Molecule 36: 30S ribosomal protein S5



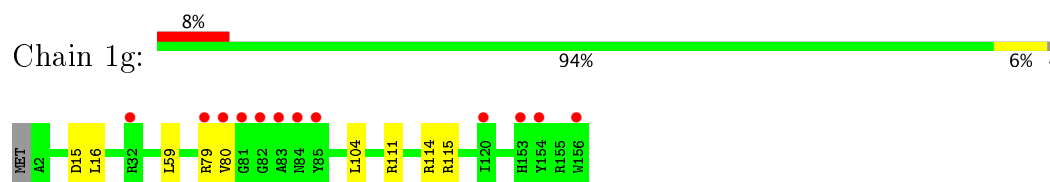
- Molecule 37: 30S ribosomal protein S6



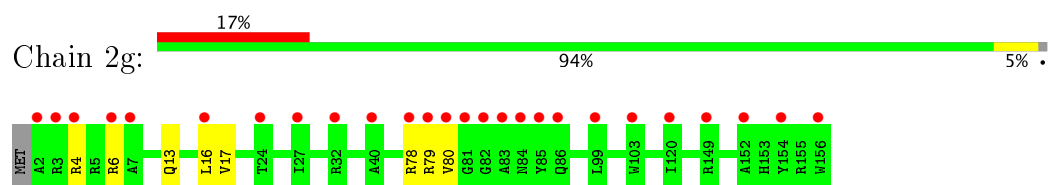
- Molecule 37: 30S ribosomal protein S6



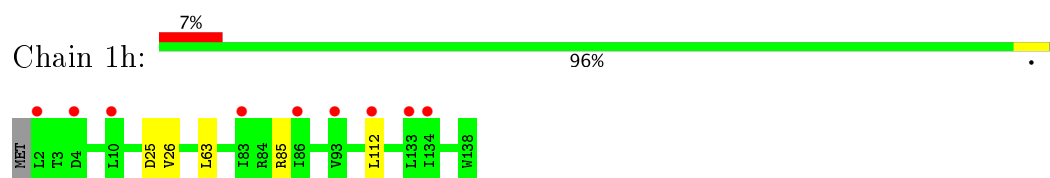
- Molecule 38: 30S ribosomal protein S7



- Molecule 38: 30S ribosomal protein S7

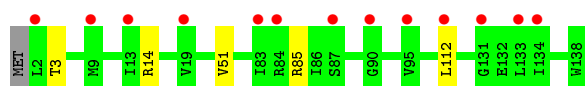


- Molecule 39: 30S ribosomal protein S8

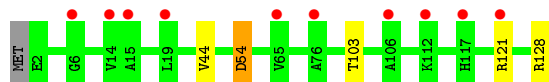


- Molecule 39: 30S ribosomal protein S8

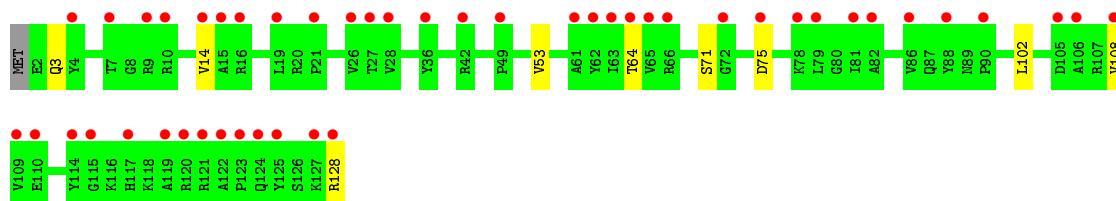
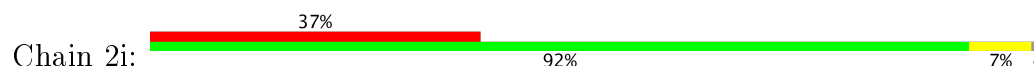




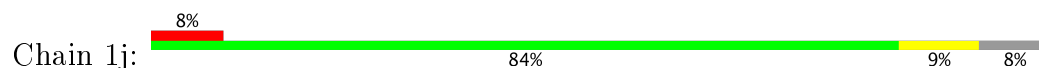
- Molecule 40: 30S ribosomal protein S9



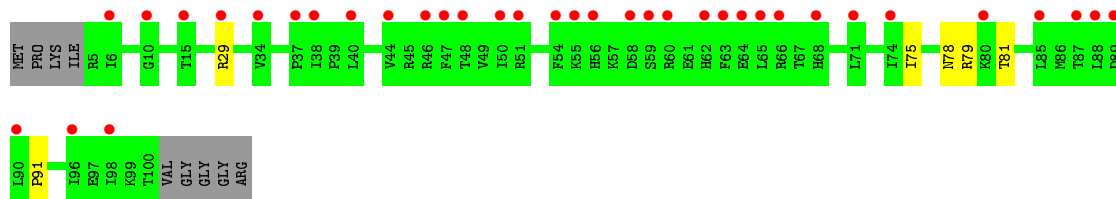
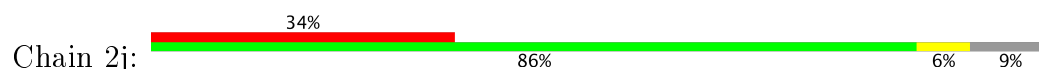
- Molecule 40: 30S ribosomal protein S9



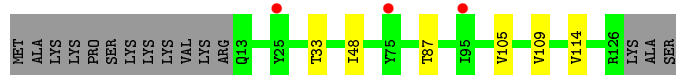
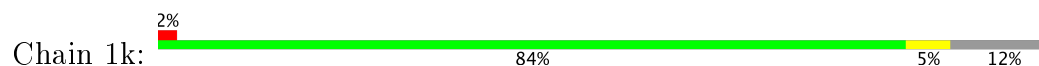
- Molecule 41: 30S ribosomal protein S10



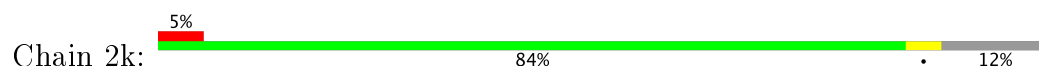
- Molecule 41: 30S ribosomal protein S10

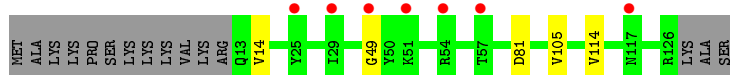


- Molecule 42: 30S ribosomal protein S11

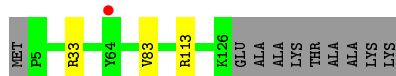
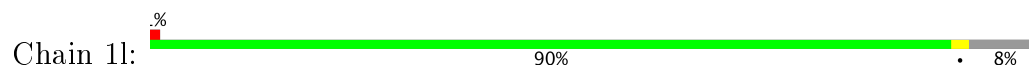


- Molecule 42: 30S ribosomal protein S11

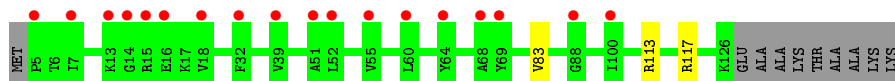
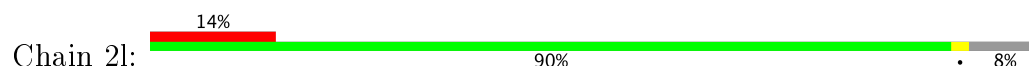




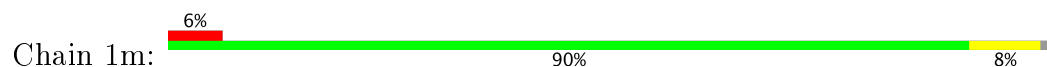
- Molecule 43: 30S ribosomal protein S12



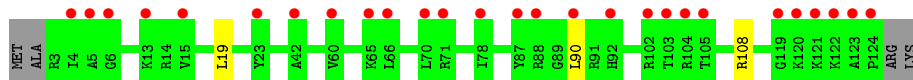
- Molecule 43: 30S ribosomal protein S12



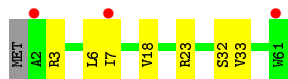
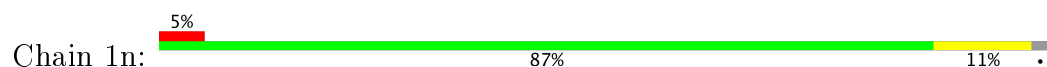
- Molecule 44: 30S ribosomal protein S13



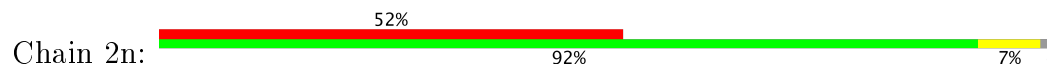
- Molecule 44: 30S ribosomal protein S13



- Molecule 45: 30S ribosomal protein S14 type Z

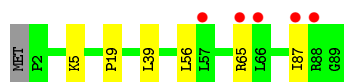


- Molecule 45: 30S ribosomal protein S14 type Z

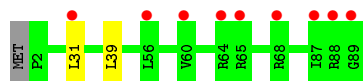


- Molecule 46: 30S ribosomal protein S15

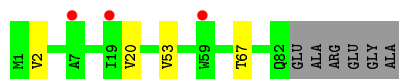
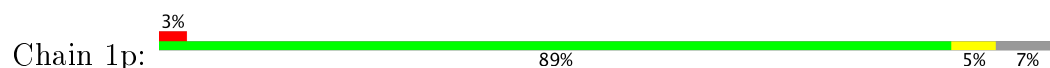




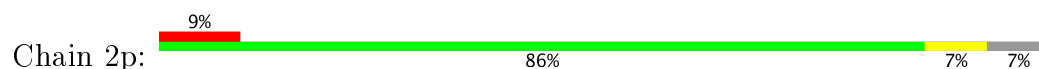
- Molecule 46: 30S ribosomal protein S15



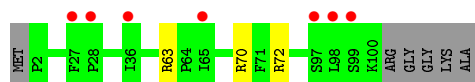
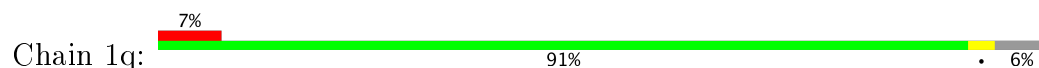
- Molecule 47: 30S ribosomal protein S16



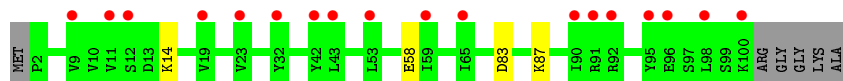
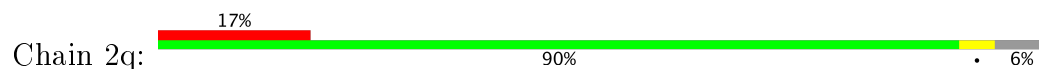
- Molecule 47: 30S ribosomal protein S16



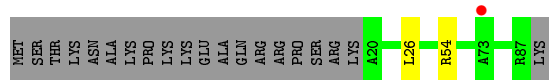
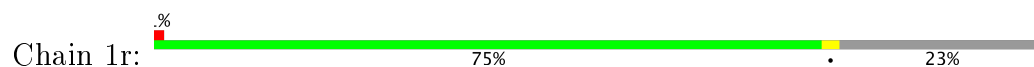
- Molecule 48: 30S ribosomal protein S17



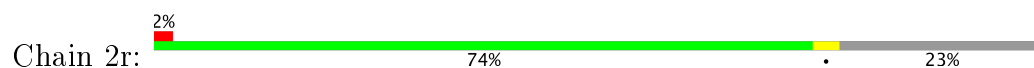
- Molecule 48: 30S ribosomal protein S17

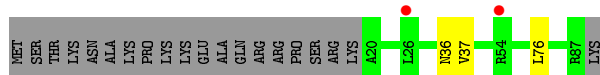


- Molecule 49: 30S ribosomal protein S18

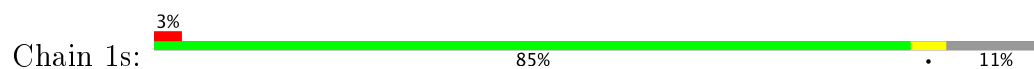


- Molecule 49: 30S ribosomal protein S18

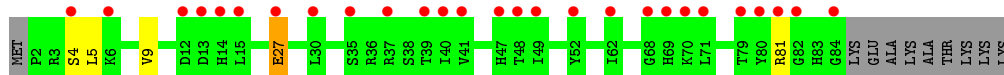
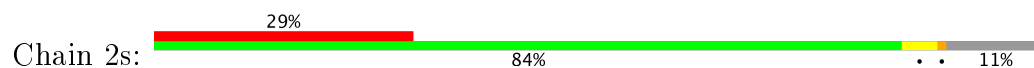




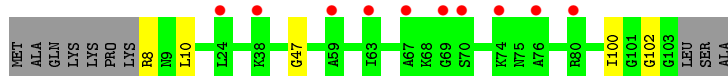
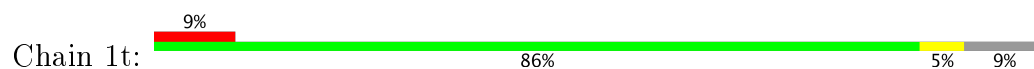
- Molecule 50: 30S ribosomal protein S19



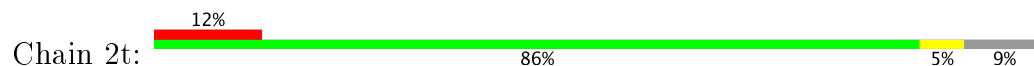
- Molecule 50: 30S ribosomal protein S19



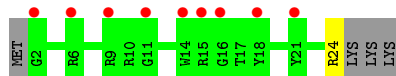
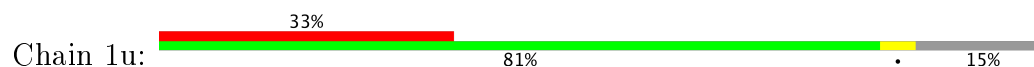
- Molecule 51: 30S ribosomal protein S20



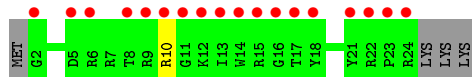
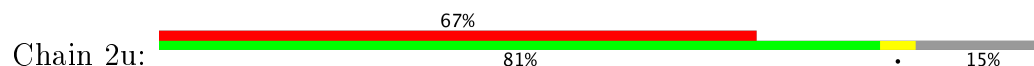
- Molecule 51: 30S ribosomal protein S20



- Molecule 52: 30S ribosomal protein Thx



- Molecule 52: 30S ribosomal protein Thx



- Molecule 53: mRNA

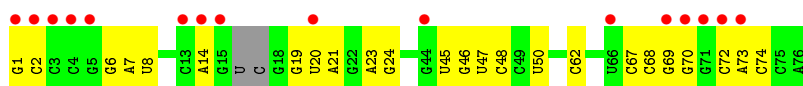




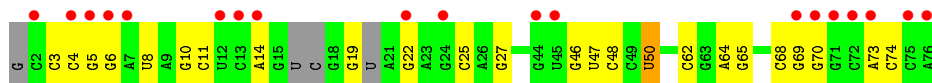
• Molecule 53: mRNA



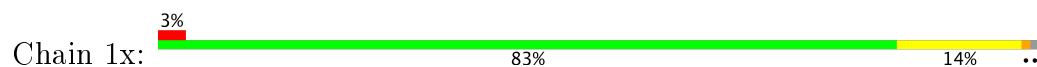
• Molecule 54: tRNA, A-site and E-site tRNAs



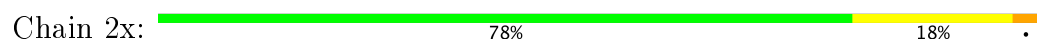
• Molecule 54: tRNA, A-site and E-site tRNAs



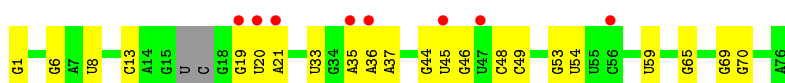
• Molecule 55: P-site tRNA



• Molecule 55: P-site tRNA



• Molecule 56: tRNA



• Molecule 56: tRNA



4 Data and refinement statistics

Property	Value	Source
Space group	P 21 21 21	Depositor
Cell constants a, b, c, α , β , γ	209.64Å 449.06Å 622.01Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	155.50 – 2.70 364.09 – 2.70	Depositor EDS
% Data completeness (in resolution range)	99.2 (155.50-2.70) 99.3 (364.09-2.70)	Depositor EDS
R_{merge}	0.16	Depositor
R_{sym}	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ ¹	1.25 (at 2.69Å)	Xtriage
Refinement program	PHENIX 1.8.2	Depositor
R, R_{free}	0.207 , 0.252 0.213 , 0.256	Depositor DCC
R_{free} test set	78960 reflections (5.28%)	DCC
Wilson B-factor (Å ²)	51.0	Xtriage
Anisotropy	0.239	Xtriage
Bulk solvent k_{sol} (e/Å ³), B_{sol} (Å ²)	0.30 , 54.2	EDS
L-test for twinning ²	$\langle L \rangle = 0.46$, $\langle L^2 \rangle = 0.29$	Xtriage
Estimated twinning fraction	No twinning to report.	Xtriage
F_o, F_c correlation	0.90	EDS
Total number of atoms	302030	wwPDB-VP
Average B, all atoms (Å ²)	53.0	wwPDB-VP

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 1.51% 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: 2MU, MIA, SF4, 0TD, 2MA, 2MG, G7M, MA6, 9VD, 9VA, BB9, UR3, M2G, ZN, K, 5MU, MG, OMG, 5MC, 4OC, 4SU, PSU

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

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# $ Z > 5$	RMSZ	# $ Z > 5$
1	1A	0.54	0/69009	0.98	82/107712 (0.1%)
1	2A	0.41	0/67293	0.88	41/105034 (0.0%)
2	1B	0.46	1/2882 (0.0%)	0.88	0/4494
2	2B	0.41	1/2879 (0.0%)	0.87	1/4487 (0.0%)
3	1D	0.36	0/2186	0.57	0/2944
3	2D	0.33	0/2186	0.54	0/2944
4	1E	0.36	0/1592	0.54	0/2149
4	2E	0.31	0/1592	0.50	0/2149
5	1F	0.34	0/1619	0.57	0/2193
5	2F	0.30	0/1615	0.49	0/2188
6	1G	0.29	0/1448	0.49	0/1957
6	2G	0.30	0/1453	0.52	0/1963
7	1H	0.33	0/1356	0.50	0/1834
7	2H	0.29	0/1356	0.45	0/1834
8	1I	0.29	0/1112	0.51	0/1514
8	2I	0.27	0/1079	0.50	0/1475
9	1N	0.34	0/1144	0.53	0/1543
9	2N	0.29	0/1144	0.46	0/1543
10	1O	0.38	0/943	0.56	0/1269
10	2O	0.34	0/943	0.52	0/1269
11	1P	0.34	0/1152	0.56	0/1533
11	2P	0.31	0/1152	0.52	0/1533
12	1Q	0.35	0/1143	0.54	0/1527
12	2Q	0.30	0/1143	0.48	0/1527
13	1R	0.33	0/982	0.54	0/1312
13	2R	0.27	0/982	0.53	0/1312
14	1S	0.32	0/883	0.56	0/1176
14	2S	0.32	0/880	0.47	0/1172
15	1T	0.35	0/1105	0.54	0/1477
15	2T	0.29	0/1097	0.46	0/1468
16	1U	0.39	0/977	0.55	0/1301

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
16	2U	0.31	0/977	0.46	0/1301
17	1V	0.38	0/782	0.57	0/1049
17	2V	0.30	0/782	0.51	0/1049
18	1W	0.35	0/897	0.54	0/1205
18	2W	0.32	0/897	0.49	0/1205
19	1X	0.38	0/764	0.57	0/1025
19	2X	0.31	0/764	0.48	0/1025
20	1Y	0.36	0/819	0.57	1/1095 (0.1%)
20	2Y	0.32	0/819	0.53	0/1095
21	1Z	0.32	0/1267	0.53	0/1717
21	2Z	0.30	0/1299	0.50	0/1763
22	10	0.37	0/662	0.58	0/881
22	20	0.30	0/662	0.50	0/881
23	11	0.35	0/762	0.53	0/1014
23	21	0.32	0/762	0.53	0/1014
24	12	0.32	0/590	0.49	0/781
24	22	0.28	0/590	0.42	0/781
25	13	0.32	0/474	0.53	0/635
25	23	0.28	0/469	0.46	0/630
26	14	0.35	0/565	0.53	0/761
26	24	0.31	0/545	0.50	0/737
27	15	0.35	0/469	0.59	1/635 (0.2%)
27	25	0.32	0/469	0.50	0/635
28	16	0.35	0/460	0.55	0/613
28	26	0.29	0/456	0.49	0/608
29	17	0.34	0/426	0.54	0/561
29	27	0.31	0/426	0.52	0/561
30	18	0.32	0/525	0.55	0/691
30	28	0.30	0/525	0.49	0/691
31	19	0.34	0/310	0.53	0/407
31	29	0.30	0/310	0.52	0/407
32	1a	0.39	0/35795	0.89	39/55864 (0.1%)
33	1b	0.29	0/1881	0.48	0/2542
33	2b	0.31	0/1860	0.47	0/2518
34	1c	0.29	0/1572	0.47	0/2126
34	2c	0.29	0/1566	0.46	0/2119
35	1d	0.30	0/1685	0.49	0/2262
35	2d	0.29	0/1704	0.47	0/2284
36	1e	0.31	0/1145	0.51	0/1543
36	2e	0.31	0/1149	0.51	0/1548
37	1f	0.30	0/823	0.47	0/1115
37	2f	0.31	0/829	0.48	0/1123
38	1g	0.28	0/1250	0.44	0/1679

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
38	2g	0.28	0/1254	0.44	0/1683
39	1h	0.28	0/1108	0.48	0/1494
39	2h	0.28	0/1108	0.46	0/1494
40	1i	0.29	0/1002	0.50	0/1346
40	2i	0.30	0/997	0.50	0/1343
41	1j	0.28	0/722	0.48	0/982
41	2j	0.30	0/727	0.52	0/988
42	1k	0.29	0/844	0.48	0/1145
42	2k	0.29	0/848	0.48	0/1149
43	1l	0.31	0/937	0.53	0/1260
43	2l	0.29	0/937	0.49	0/1260
44	1m	0.29	0/969	0.49	0/1302
44	2m	0.28	0/961	0.51	0/1291
45	1n	0.30	0/501	0.48	0/664
45	2n	0.32	0/501	0.50	0/664
46	1o	0.28	0/739	0.43	0/985
46	2o	0.26	0/739	0.42	0/985
47	1p	0.28	0/697	0.51	0/939
47	2p	0.29	0/693	0.50	0/935
48	1q	0.30	0/836	0.48	0/1117
48	2q	0.30	0/836	0.47	0/1117
49	1r	0.28	0/560	0.48	0/746
49	2r	0.28	0/560	0.44	0/746
50	1s	0.28	0/667	0.54	0/900
50	2s	0.30	0/661	0.54	0/893
51	1t	0.28	0/730	0.47	0/965
51	2t	0.28	0/729	0.43	0/965
52	1u	0.27	0/203	0.44	0/266
52	2u	0.36	0/203	0.48	0/266
53	1v	0.42	0/310	0.86	0/480
53	2v	0.47	0/310	0.86	0/480
54	1w	0.54	1/1606 (0.1%)	1.09	0/2497
54	2w	0.51	0/1556	1.13	4/2418 (0.2%)
55	1x	0.55	1/1725 (0.1%)	1.13	13/2689 (0.5%)
55	2x	0.46	0/1725	1.06	7/2689 (0.3%)
56	1y	0.72	6/1632 (0.4%)	1.22	13/2540 (0.5%)
56	2y	0.75	6/1609 (0.4%)	1.24	12/2502 (0.5%)
57	2a	0.38	3/35886 (0.0%)	0.90	50/56005 (0.1%)
58	A	2.13	4/74 (5.4%)	2.00	5/97 (5.2%)
58	B	2.13	3/74 (4.1%)	2.04	4/97 (4.1%)
All	All	0.42	26/316886 (0.0%)	0.84	273/474393 (0.1%)

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
26	24	0	1
33	1b	0	1
50	2s	0	1
58	A	1	0
58	B	1	0
All	All	2	3

All (26) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
56	2y	37	A	C5-C4	13.66	1.48	1.38
56	1y	37	A	C5-C4	13.39	1.48	1.38
58	B	12	SER	CA-CB	-13.07	1.33	1.52
58	A	12	SER	CA-CB	-12.57	1.34	1.52
56	2y	1	G	OP3-P	-10.39	1.48	1.61
56	1y	1	G	OP3-P	-10.30	1.48	1.61
2	2B	1	U	OP3-P	-10.05	1.49	1.61
54	1w	1	G	OP3-P	-10.03	1.49	1.61
2	1B	1	U	OP3-P	-9.80	1.49	1.61
57	2a	1272	G	N1-C2	-9.26	1.30	1.37
56	2y	37	A	C5-C6	9.05	1.49	1.41
56	1y	37	A	C5-C6	8.92	1.49	1.41
57	2a	1272	G	C6-N1	-8.85	1.33	1.39
58	B	12	SER	CA-C	-7.53	1.33	1.52
58	A	12	SER	CA-C	-7.29	1.34	1.52
56	1y	37	A	C8-N7	6.86	1.36	1.31
56	2y	37	A	C8-N7	6.80	1.36	1.31
56	2y	37	A	N7-C5	-6.56	1.35	1.39
58	A	12	SER	N-CA	-6.48	1.33	1.46
58	B	12	SER	N-CA	-6.44	1.33	1.46
56	1y	37	A	N7-C5	-6.31	1.35	1.39
58	A	12	SER	CB-OG	-6.19	1.34	1.42
56	1y	37	A	N9-C4	-5.51	1.34	1.37
56	2y	37	A	N9-C4	-5.27	1.34	1.37
57	2a	1263	C	N3-C4	-5.20	1.30	1.33
55	1x	22	G	N7-C5	5.10	1.42	1.39

All (273) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
57	2a	1263	C	N1-C2-O2	22.88	132.63	118.90
57	2a	1272	G	N3-C2-N2	20.83	134.48	119.90
56	2y	37	A	C2-N3-C4	20.81	121.00	110.60
56	1y	37	A	C2-N3-C4	20.56	120.88	110.60
57	2a	1272	G	C5-C6-O6	20.05	140.63	128.60
57	2a	1272	G	N1-C2-N2	-18.59	99.47	116.20
57	2a	1263	C	C2-N3-C4	14.28	127.04	119.90
57	2a	1263	C	N3-C2-O2	-13.11	112.72	121.90
57	2a	1272	G	N1-C6-O6	-12.89	112.16	119.90
56	2y	37	A	N1-C2-N3	-12.78	122.91	129.30
1	1A	1132	A	N1-C6-N6	-12.72	110.97	118.60
56	1y	37	A	N1-C2-N3	-12.44	123.08	129.30
56	2y	37	A	N3-C4-C5	-11.59	118.69	126.80
56	1y	37	A	N3-C4-C5	-11.35	118.85	126.80
1	1A	1686	U	O5'-P-OP2	-10.96	95.84	105.70
32	1a	1027	C	C5-C4-N4	10.85	127.79	120.20
32	1a	1027	C	N3-C2-O2	-10.81	114.33	121.90
1	1A	1121	C	N1-C2-O2	10.74	125.34	118.90
1	1A	1121	C	C2-N3-C4	10.60	125.20	119.90
57	2a	1272	G	C6-N1-C2	10.44	131.36	125.10
1	2A	2136	C	N1-C2-O2	9.82	124.79	118.90
56	2y	37	A	N3-C4-N9	9.80	135.24	127.40
57	2a	1263	C	C5-C6-N1	9.43	125.72	121.00
56	1y	37	A	N3-C4-N9	9.18	134.74	127.40
1	1A	1109	G	C5-C6-O6	8.94	133.96	128.60
57	2a	1263	C	C5-C4-N4	8.91	126.44	120.20
1	1A	537	G	O4'-C1'-N9	8.91	115.33	108.20
57	2a	1001(A)	G	N3-C4-N9	8.86	131.31	126.00
55	1x	46	G	C6-N1-C2	-8.76	119.84	125.10
56	1y	37	A	C4-C5-N7	-8.76	106.32	110.70
57	2a	1272	G	C5-C6-N1	-8.68	107.16	111.50
55	1x	14	A	C4-C5-C6	8.67	121.33	117.00
1	1A	1109	G	C6-N1-C2	8.49	130.19	125.10
56	1y	33	U	C2-N1-C1'	8.34	127.71	117.70
1	1A	2189	U	C2-N1-C1'	8.27	127.62	117.70
32	1a	1027	C	N3-C4-C5	-8.23	118.61	121.90
55	1x	14	A	C5-N7-C8	8.17	107.98	103.90
56	2y	37	A	C4-C5-N7	-8.09	106.65	110.70
32	1a	1025	U	N1-C2-O2	8.03	128.42	122.80
1	2A	2473	U	C2-N1-C1'	7.99	127.29	117.70
32	1a	1027	C	C6-N1-C2	-7.96	117.12	120.30
1	1A	2694	U	O5'-P-OP2	-7.94	98.56	105.70
57	2a	1263	C	N3-C4-N4	-7.85	112.50	118.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
2	2B	80	U	O4'-C1'-N1	7.84	114.48	108.20
56	2y	33	U	C2-N1-C1'	7.83	127.09	117.70
32	1a	1034	G	N3-C2-N2	7.81	125.37	119.90
1	1A	215	G	O4'-C1'-N9	7.81	114.44	108.20
56	1y	37	A	C5-N7-C8	7.71	107.76	103.90
55	1x	22	G	C5-N7-C8	-7.71	100.45	104.30
32	1a	1034	G	N9-C4-C5	-7.64	102.34	105.40
57	2a	1272	G	C2-N3-C4	-7.63	108.08	111.90
1	1A	591	U	C5-C4-O4	-7.62	121.33	125.90
1	1A	1807	G	O5'-P-OP2	-7.56	98.89	105.70
32	1a	1030(B)	C	C2-N1-C1'	7.53	127.08	118.80
57	2a	1263	C	C2-N1-C1'	7.53	127.08	118.80
1	1A	1109	G	N3-C2-N2	7.51	125.16	119.90
1	1A	2504	U	O5'-P-OP1	-7.43	99.02	105.70
1	1A	1132	A	C5-C6-N6	7.37	129.60	123.70
55	2x	14	A	C5-N7-C8	7.34	107.57	103.90
1	1A	2189	U	N1-C2-O2	7.30	127.91	122.80
58	B	12	SER	CB-CA-C	7.27	123.92	110.10
32	1a	1027	C	N1-C2-O2	7.26	123.25	118.90
57	2a	1263	C	C6-N1-C2	-7.15	117.44	120.30
1	1A	2189	U	N3-C2-O2	-7.14	117.20	122.20
58	A	12	SER	CB-CA-C	7.13	123.65	110.10
56	2y	37	A	C5-N7-C8	7.10	107.45	103.90
57	2a	1039	C	C5-C4-N4	-7.07	115.25	120.20
1	1A	2807	C	N1-C2-O2	7.04	123.12	118.90
55	1x	14	A	C5-C6-N1	-6.98	114.21	117.70
1	2A	2136	C	N3-C2-O2	-6.94	117.04	121.90
55	2x	14	A	C4-C5-C6	6.94	120.47	117.00
1	2A	2473	U	N1-C2-O2	6.87	127.61	122.80
1	1A	1222	A	O5'-P-OP1	-6.84	99.54	105.70
32	1a	1034	G	N3-C4-N9	6.84	130.10	126.00
1	2A	2248	C	O5'-P-OP2	-6.81	99.57	105.70
32	1a	1034	G	C4-C5-N7	6.80	113.52	110.80
57	2a	1001(A)	G	C4-N9-C1'	6.75	135.28	126.50
1	1A	2566	U	O5'-P-OP1	-6.74	99.64	105.70
57	2a	1263	C	C4-C5-C6	-6.73	114.03	117.40
57	2a	79	G	C5-C6-O6	6.69	132.62	128.60
1	2A	2473	U	N3-C2-O2	-6.68	117.52	122.20
58	A	11	ASN	C-N-CA	-6.68	105.00	121.70
54	2w	50	U	C5-C4-O4	-6.67	121.90	125.90
1	1A	2050	U	N3-C4-O4	-6.61	114.77	119.40
56	1y	33	U	N1-C2-O2	6.59	127.41	122.80

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
56	2y	37	A	C8-N9-C4	6.58	108.43	105.80
57	2a	1263	C	N1-C2-N3	-6.55	114.61	119.20
1	1A	1815	A	O5'-P-OP2	-6.54	99.81	105.70
32	1a	558	G	O5'-P-OP1	-6.54	99.82	105.70
1	1A	892	G	O4'-C1'-N9	6.52	113.42	108.20
57	2a	1039	C	C2-N1-C1'	6.51	125.96	118.80
1	2A	2140	C	N1-C2-O2	6.49	122.79	118.90
56	1y	37	A	C8-N9-C4	6.49	108.39	105.80
32	1a	1030(B)	C	C6-N1-C2	-6.47	117.71	120.30
58	B	11	ASN	C-N-CA	-6.46	105.55	121.70
55	1x	22	G	C4-C5-C6	-6.41	114.95	118.80
1	2A	2139	C	C2-N1-C1'	6.37	125.81	118.80
1	1A	2359	C	N1-C2-O2	6.35	122.71	118.90
1	2A	2140	C	C2-N1-C1'	6.32	125.75	118.80
1	1A	848	G	O5'-P-OP2	-6.31	100.02	105.70
32	1a	1027	C	N3-C4-N4	-6.30	113.59	118.00
1	2A	512	G	O4'-C1'-N9	6.28	113.23	108.20
57	2a	1001(A)	G	N3-C4-C5	-6.28	125.46	128.60
32	1a	267	C	O5'-P-OP1	-6.27	100.06	105.70
57	2a	1001(A)	G	C8-N9-C1'	-6.26	118.86	127.00
56	1y	33	U	C6-N1-C1'	-6.26	112.43	121.20
57	2a	1272	G	C4-N9-C1'	6.26	134.64	126.50
1	1A	2701	U	P-O3'-C3'	6.26	127.21	119.70
57	2a	1272	G	C8-N9-C1'	-6.24	118.89	127.00
1	1A	2858	G	O4'-C1'-N9	6.24	113.19	108.20
32	1a	1030(B)	C	N1-C2-O2	6.22	122.64	118.90
55	2x	46	G	C6-N1-C2	-6.20	121.38	125.10
57	2a	754	C	C2-N1-C1'	6.19	125.61	118.80
55	1x	22	G	N3-C4-N9	-6.19	122.28	126.00
32	1a	299	G	C5-C6-O6	-6.19	124.89	128.60
1	1A	840	A	O5'-P-OP2	-6.15	100.17	105.70
1	2A	1313	U	C2-N1-C1'	6.12	125.05	117.70
1	2A	1352	U	O5'-P-OP1	-6.11	100.20	105.70
1	1A	2383	G	C5-C6-N1	6.09	114.55	111.50
1	2A	945	A	N1-C6-N6	6.07	122.24	118.60
56	2y	37	A	C6-N1-C2	6.07	122.24	118.60
57	2a	1001(A)	G	C6-C5-N7	-6.07	126.76	130.40
32	1a	1030	C	N1-C2-O2	6.06	122.53	118.90
1	2A	2167	U	C2-N1-C1'	6.03	124.94	117.70
55	1x	46	G	C5-C6-N1	6.01	114.50	111.50
1	2A	2149	G	N3-C4-N9	5.99	129.59	126.00
57	2a	299	G	C5-C6-O6	-5.99	125.01	128.60

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	1A	894	U	C2-N1-C1'	5.96	124.86	117.70
32	1a	1002	G	C4-N9-C1'	5.96	134.25	126.50
1	2A	1992	G	P-O3'-C3'	5.96	126.86	119.70
1	2A	2167	U	N1-C2-O2	5.96	126.97	122.80
1	2A	1698	A	O4'-C1'-N9	5.95	112.96	108.20
32	1a	266	G	P-O3'-C3'	5.92	126.80	119.70
1	1A	1128	U	N3-C4-O4	-5.89	115.27	119.40
1	1A	2641	A	P-O3'-C3'	5.87	126.74	119.70
57	2a	754	C	N1-C2-O2	5.85	122.41	118.90
57	2a	1039	C	N3-C4-N4	5.85	122.10	118.00
1	1A	2014	G	C8-N9-C4	-5.85	104.06	106.40
32	1a	1030	C	C2-N3-C4	5.84	122.82	119.90
1	2A	141	A	N7-C8-N9	5.83	116.71	113.80
1	1A	1121	C	C5-C4-N4	5.83	124.28	120.20
55	1x	22	G	N7-C8-N9	5.83	116.01	113.10
1	1A	1128	U	N3-C4-C5	5.82	118.09	114.60
55	1x	22	G	C8-N9-C1'	5.82	134.57	127.00
32	1a	1025	U	N3-C2-O2	-5.82	118.12	122.20
54	2w	25	C	C5-C4-N4	5.82	124.27	120.20
56	1y	37	A	C6-N1-C2	5.79	122.07	118.60
1	1A	1958	A	O4'-C1'-N9	5.75	112.80	108.20
1	1A	593	G	C5-C6-O6	-5.74	125.15	128.60
56	2y	47	U	C2-N1-C1'	5.74	124.59	117.70
1	1A	194	G	O5'-P-OP2	-5.73	100.54	105.70
1	1A	399	G	O4'-C1'-N9	5.73	112.78	108.20
56	1y	37	A	C6-C5-N7	5.72	136.31	132.30
57	2a	266	G	P-O3'-C3'	5.72	126.56	119.70
1	1A	1462	G	O4'-C1'-N9	5.71	112.77	108.20
57	2a	1043	C	N1-C2-O2	5.70	122.32	118.90
1	2A	141	A	C8-N9-C4	-5.69	103.53	105.80
1	1A	1221	G	OP1-P-O3'	5.66	117.66	105.20
1	1A	1295	U	O5'-P-OP1	-5.66	100.61	105.70
32	1a	1067	A	P-O3'-C3'	5.65	126.47	119.70
32	1a	1002	G	C8-N9-C1'	-5.64	119.67	127.00
1	1A	2014	G	P-O3'-C3'	5.63	126.46	119.70
1	1A	1398	U	O5'-P-OP1	-5.63	100.63	105.70
1	2A	228	A	P-O3'-C3'	5.63	126.45	119.70
1	1A	1431	G	O4'-C1'-N9	5.61	112.68	108.20
32	1a	841	U	C5-C6-N1	5.60	125.50	122.70
1	2A	1639	U	O5'-P-OP2	-5.59	100.67	105.70
1	1A	2189	U	C5-C6-N1	5.59	125.49	122.70
1	1A	2015	U	O5'-P-OP1	-5.58	100.67	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
55	1x	14	A	C4-N9-C1'	5.57	136.32	126.30
1	2A	2139	C	N1-C2-O2	5.56	122.24	118.90
55	2x	46	G	N3-C2-N2	-5.55	116.01	119.90
57	2a	1130	A	O5'-P-OP1	-5.55	100.71	105.70
57	2a	687	A	P-O3'-C3'	5.54	126.34	119.70
1	2A	1530	C	P-O3'-C3'	5.51	126.32	119.70
1	1A	1219	A	P-O3'-C3'	5.51	126.31	119.70
20	1Y	54	LYS	C-N-CA	5.50	135.46	121.70
57	2a	65	U	P-O3'-C3'	5.50	126.30	119.70
57	2a	1028	C	C2-N3-C4	5.48	122.64	119.90
27	15	58	LEU	CA-CB-CG	5.48	127.90	115.30
58	A	4	GLY	N-CA-C	-5.47	99.42	113.10
32	1a	1027	C	C6-N1-C1'	5.47	127.37	120.80
1	1A	572	A	P-O3'-C3'	5.47	126.26	119.70
57	2a	299	G	N1-C6-O6	5.47	123.18	119.90
32	1a	1034	G	C8-N9-C1'	-5.46	119.91	127.00
1	1A	2082	A	C8-N9-C4	5.46	107.98	105.80
57	2a	1001(A)	G	N9-C4-C5	-5.44	103.22	105.40
58	B	3	PRO	N-CA-C	5.43	126.22	112.10
1	2A	2139	C	C6-N1-C1'	-5.43	114.29	120.80
1	1A	2803	A	C2-N3-C4	5.42	113.31	110.60
1	1A	961	C	N1-C2-O2	5.42	122.15	118.90
55	1x	14	A	C4-C5-N7	-5.42	107.99	110.70
1	1A	1359	U	N3-C2-O2	-5.42	118.41	122.20
1	1A	410	U	C2-N1-C1'	-5.41	111.21	117.70
58	B	12	SER	N-CA-C	5.40	125.57	111.00
1	2A	2174	C	N1-C2-O2	5.38	122.13	118.90
55	1x	14	A	C8-N9-C1'	-5.37	118.03	127.70
1	2A	2128	C	C2-N3-C4	5.36	122.58	119.90
56	1y	37	A	N7-C8-N9	-5.36	111.12	113.80
55	2x	14	A	C5-C6-N1	-5.36	115.02	117.70
1	2A	2167	U	N3-C2-O2	-5.35	118.45	122.20
1	2A	383	U	O4'-C1'-N1	5.35	112.48	108.20
57	2a	254	G	O5'-P-OP1	-5.35	100.89	105.70
32	1a	266	G	O4'-C1'-N9	-5.34	103.92	108.20
56	2y	33	U	C6-N1-C1'	-5.34	113.72	121.20
1	1A	1020	C	N1-C2-O2	-5.32	115.71	118.90
1	1A	2442	A	C2-N3-C4	5.31	113.26	110.60
1	1A	2610	A	O5'-P-OP1	-5.31	100.92	105.70
57	2a	1505	G	N3-C4-N9	-5.31	122.81	126.00
1	1A	1648	U	N3-C4-O4	-5.30	115.69	119.40
1	1A	649	C	O5'-P-OP1	-5.28	100.95	105.70

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
55	2x	22	G	C5-N7-C8	-5.27	101.66	104.30
1	2A	528	A	P-O3'-C3'	5.27	126.02	119.70
57	2a	1039	C	C6-N1-C1'	-5.27	114.48	120.80
1	1A	2331	G	C5-N7-C8	-5.26	101.67	104.30
58	A	3	PRO	C-N-CA	5.25	133.32	122.30
1	1A	1694	G	O4'-C1'-N9	-5.24	104.00	108.20
1	2A	801	G	O5'-P-OP2	-5.24	100.99	105.70
1	1A	1441	A	C8-N9-C4	5.23	107.89	105.80
1	2A	2149	G	C4-N9-C1'	5.23	133.30	126.50
1	1A	2359	C	N3-C2-O2	-5.23	118.24	121.90
1	2A	845	G	O4'-C1'-N9	5.22	112.38	108.20
1	1A	12	U	C2-N1-C1'	5.22	123.97	117.70
57	2a	913	A	P-O3'-C3'	5.22	125.96	119.70
54	2w	50	U	N3-C4-O4	5.21	123.05	119.40
1	2A	845	G	C4-N9-C1'	5.20	133.27	126.50
1	1A	1700	G	P-O3'-C3'	5.20	125.94	119.70
1	1A	2331	G	O4'-C1'-N9	5.19	112.35	108.20
57	2a	1125	U	C2-N1-C1'	5.18	123.92	117.70
1	2A	2689	U	P-O3'-C3'	5.18	125.91	119.70
1	2A	1298	C	O5'-P-OP2	-5.17	101.04	105.70
1	2A	2149	G	C8-N9-C1'	-5.17	120.27	127.00
1	1A	831	A	O4'-C1'-N9	5.17	112.34	108.20
32	1a	560	U	C3'-C2'-C1'	5.17	105.64	101.50
57	2a	1264	C	N1-C2-O2	5.17	122.00	118.90
58	A	12	SER	N-CA-C	5.17	124.95	111.00
55	2x	34	C	C2-N1-C1'	5.16	124.48	118.80
57	2a	563	A	O4'-C1'-N9	5.16	112.33	108.20
1	1A	1219	A	OP1-P-O3'	5.16	116.54	105.20
56	2y	33	U	N1-C2-O2	5.15	126.40	122.80
1	1A	591	U	N3-C4-C5	5.14	117.68	114.60
57	2a	299	G	C4-C5-N7	5.13	112.85	110.80
32	1a	1225	A	C5-C6-N6	5.13	127.80	123.70
57	2a	1065	U	P-O3'-C3'	5.13	125.86	119.70
1	1A	1128	U	C2-N3-C4	-5.13	123.92	127.00
1	1A	2054	G	C5-N7-C8	5.13	106.86	104.30
32	1a	1030(B)	C	C5-C6-N1	5.13	123.56	121.00
1	1A	1109	G	C5-C6-N1	-5.12	108.94	111.50
57	2a	1001(A)	G	C4-C5-N7	5.12	112.85	110.80
1	1A	849	A	O5'-P-OP1	-5.12	101.09	105.70
32	1a	1030(B)	C	N3-C2-O2	-5.12	118.32	121.90
1	1A	1700	G	C8-N9-C4	-5.11	104.36	106.40
1	2A	2473	U	C6-N1-C1'	-5.11	114.05	121.20

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	1A	2253	A	OP1-P-O3'	5.11	116.43	105.20
1	1A	1298	G	N3-C4-N9	-5.10	122.94	126.00
1	2A	645	C	C2-N1-C1'	5.09	124.40	118.80
1	1A	715	G	OP2-P-O3'	5.08	116.38	105.20
32	1a	560	U	C2-N1-C1'	5.08	123.80	117.70
1	1A	1426	G	O5'-P-OP2	-5.07	101.14	105.70
1	1A	1121	C	C5-C6-N1	5.06	123.53	121.00
32	1a	1225	A	C6-N1-C2	5.06	121.64	118.60
32	1a	1034	G	C4-N9-C1'	5.06	133.07	126.50
1	2A	645	C	N1-C2-O2	5.05	121.93	118.90
57	2a	1039	C	C5-C6-N1	5.04	123.52	121.00
1	1A	2189	U	C6-N1-C2	-5.04	117.98	121.00
32	1a	1065	U	P-O3'-C3'	5.03	125.74	119.70
1	1A	1808	U	OP1-P-O3'	5.03	116.26	105.20
32	1a	1030	C	C5-C4-N4	5.02	123.72	120.20
32	1a	1020	U	C2-N1-C1'	5.01	123.72	117.70
1	2A	228	A	OP1-P-O3'	5.01	116.22	105.20
1	1A	1299	A	N7-C8-N9	-5.01	111.30	113.80
32	1a	1442	G	C2-N3-C4	5.00	114.40	111.90
54	2w	10	G	C5-C6-N1	-5.00	109.00	111.50

All (2) chirality outliers are listed below:

Mol	Chain	Res	Type	Atom
58	A	12	SER	CA
58	B	12	SER	CA

All (3) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
33	1b	8	LYS	Peptide
26	24	56	VAL	Peptide
50	2s	27	GLU	Peptide

5.2 Too-close contacts [i](#)

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	1A	61852	0	31194	537	0
1	2A	60322	0	30423	646	0
2	1B	2577	0	1305	14	0
2	2B	2575	0	1303	43	0
3	1D	2136	0	2218	41	0
3	2D	2136	0	2218	45	0
4	1E	1559	0	1618	25	0
4	2E	1559	0	1618	33	0
5	1F	1584	0	1625	30	0
5	2F	1580	0	1619	38	0
6	1G	1423	0	1436	24	0
6	2G	1428	0	1438	38	0
7	1H	1330	0	1407	24	0
7	2H	1330	0	1407	41	0
8	1I	1097	0	1140	28	0
8	2I	1064	0	1082	25	0
9	1N	1117	0	1184	20	0
9	2N	1117	0	1184	24	0
10	1O	933	0	996	11	0
10	2O	933	0	996	18	0
11	1P	1135	0	1212	27	0
11	2P	1135	0	1212	29	0
12	1Q	1122	0	1179	17	0
12	2Q	1122	0	1179	20	0
13	1R	968	0	1033	17	0
13	2R	968	0	1032	19	0
14	1S	873	0	927	17	0
14	2S	870	0	923	19	0
15	1T	1091	0	1151	19	0
15	2T	1083	0	1136	21	0
16	1U	959	0	1019	15	0
16	2U	959	0	1019	10	0
17	1V	771	0	830	8	0
17	2V	771	0	830	14	0
18	1W	886	0	940	12	0
18	2W	886	0	940	9	0
19	1X	750	0	814	12	0
19	2X	750	0	814	17	0
20	1Y	806	0	881	14	0
20	2Y	806	0	881	15	0
21	1Z	1240	0	1240	23	0
21	2Z	1271	0	1273	36	0
22	10	653	0	674	7	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
22	20	653	0	674	13	0
23	11	755	0	826	14	0
23	21	755	0	826	17	0
24	12	588	0	643	10	0
24	22	588	0	643	8	0
25	13	469	0	518	9	0
25	23	464	0	514	9	0
26	14	552	0	533	12	0
26	24	532	0	503	13	0
27	15	455	0	465	6	0
27	25	455	0	465	7	0
28	16	453	0	473	7	0
28	26	449	0	469	8	0
29	17	418	0	467	5	0
29	27	418	0	467	11	0
30	18	517	0	582	15	0
30	28	517	0	582	15	0
31	19	307	0	335	1	0
31	29	307	0	335	11	0
32	1a	32246	0	16295	0	0
33	1b	1846	0	1867	0	0
33	2b	1825	0	1828	0	0
34	1c	1548	0	1535	0	0
34	2c	1542	0	1517	0	0
35	1d	1655	0	1672	0	0
35	2d	1674	0	1714	0	0
36	1e	1129	0	1185	0	0
36	2e	1133	0	1191	0	0
37	1f	810	0	804	0	0
37	2f	816	0	808	0	0
38	1g	1231	0	1238	0	0
38	2g	1235	0	1249	0	0
39	1h	1088	0	1126	0	0
39	2h	1088	0	1126	0	0
40	1i	983	0	986	0	0
40	2i	978	0	966	0	0
41	1j	709	0	650	0	0
41	2j	714	0	672	0	0
42	1k	829	0	825	0	0
42	2k	833	0	836	0	0
43	1l	932	0	981	0	0
43	2l	932	0	981	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
44	1m	958	0	1002	0	0
44	2m	950	0	988	0	0
45	1n	492	0	529	0	0
45	2n	492	0	529	0	0
46	1o	728	0	760	0	0
46	2o	728	0	760	0	0
47	1p	681	0	697	0	0
47	2p	677	0	686	0	0
48	1q	823	0	891	0	0
48	2q	823	0	891	0	0
49	1r	555	0	618	0	0
49	2r	555	0	618	0	0
50	1s	652	0	662	0	0
50	2s	646	0	644	0	0
51	1t	728	0	798	0	0
51	2t	727	0	796	0	0
52	1u	199	0	208	0	0
52	2u	199	0	208	0	0
53	1v	277	0	140	0	0
53	2v	277	0	140	0	0
54	1w	1592	0	818	0	0
54	2w	1544	0	787	0	0
55	1x	1625	0	829	0	0
55	2x	1625	0	828	0	0
56	1y	1585	0	806	0	0
56	2y	1565	0	797	0	0
57	2a	32327	0	16338	0	0
58	A	115	0	72	4	0
58	B	115	0	72	6	0
59	10	8	0	0	0	0
59	11	3	0	0	0	0
59	12	2	0	0	0	0
59	13	3	0	0	0	0
59	15	2	0	0	0	0
59	16	3	0	0	0	0
59	17	2	0	0	0	0
59	18	3	0	0	0	0
59	19	1	0	0	0	0
59	1A	1229	0	0	0	0
59	1B	38	0	0	0	0
59	1D	13	0	0	0	0
59	1E	14	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
59	1F	10	0	0	0	0
59	1G	5	0	0	0	0
59	1I	1	0	0	0	0
59	1N	9	0	0	0	0
59	1O	6	0	0	0	0
59	1P	4	0	0	0	0
59	1Q	6	0	0	0	0
59	1R	3	0	0	0	0
59	1S	3	0	0	0	0
59	1T	2	0	0	0	0
59	1U	5	0	0	0	0
59	1V	2	0	0	0	0
59	1W	5	0	0	0	0
59	1X	5	0	0	0	0
59	1Y	3	0	0	0	0
59	1Z	4	0	0	0	0
59	1a	278	0	0	0	0
59	1b	2	0	0	0	0
59	1d	1	0	0	0	0
59	1e	1	0	0	0	0
59	1f	3	0	0	0	0
59	1k	1	0	0	0	0
59	1l	3	0	0	0	0
59	1n	2	0	0	0	0
59	1p	1	0	0	0	0
59	1s	1	0	0	0	0
59	1t	1	0	0	0	0
59	1v	1	0	0	0	0
59	1w	12	0	0	0	0
59	1x	17	0	0	0	0
59	1y	5	0	0	0	0
59	20	3	0	0	0	0
59	23	2	0	0	0	0
59	25	3	0	0	0	0
59	27	1	0	0	0	0
59	28	2	0	0	0	0
59	2A	848	0	0	0	0
59	2B	22	0	0	0	0
59	2D	5	0	0	0	0
59	2E	7	0	0	0	0
59	2F	4	0	0	0	0
59	2G	1	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
59	2N	1	0	0	0	0
59	2O	2	0	0	0	0
59	2P	1	0	0	0	0
59	2Q	4	0	0	0	0
59	2R	3	0	0	0	0
59	2S	1	0	0	0	0
59	2T	3	0	0	0	0
59	2U	4	0	0	0	0
59	2V	1	0	0	0	0
59	2X	2	0	0	0	0
59	2Z	1	0	0	0	0
59	2a	239	0	0	0	0
59	2d	1	0	0	0	0
59	2e	1	0	0	0	0
59	2f	2	0	0	0	0
59	2g	1	0	0	0	0
59	2j	2	0	0	0	0
59	2k	1	0	0	0	0
59	2l	2	0	0	0	0
59	2p	1	0	0	0	0
59	2q	5	0	0	0	0
59	2r	1	0	0	0	0
59	2t	1	0	0	0	0
59	2v	4	0	0	0	0
59	2w	7	0	0	0	0
59	2x	4	0	0	0	0
59	2y	7	0	0	0	0
59	A	1	0	0	0	0
59	B	1	0	0	0	0
60	14	1	0	0	0	0
60	15	1	0	0	0	0
60	16	1	0	0	0	0
60	19	1	0	0	0	0
60	1Y	1	0	0	0	0
60	1n	1	0	0	0	0
60	24	1	0	0	0	0
60	25	1	0	0	0	0
60	26	1	0	0	0	0
60	29	1	0	0	0	0
60	2Y	1	0	0	0	0
60	2n	1	0	0	0	0
61	1d	8	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
61	2d	8	0	0	0	0
62	2A	1	0	0	0	0
63	10	12	0	0	0	0
63	11	13	0	0	0	0
63	12	4	0	0	1	0
63	13	5	0	0	1	0
63	14	1	0	0	0	0
63	15	5	0	0	0	0
63	16	5	0	0	1	0
63	17	9	0	0	0	0
63	18	13	0	0	1	0
63	19	1	0	0	0	0
63	1A	2636	0	0	45	0
63	1B	71	0	0	0	0
63	1D	31	0	0	0	0
63	1E	35	0	0	1	0
63	1F	18	0	0	0	0
63	1G	6	0	0	3	0
63	1H	3	0	0	0	0
63	1I	2	0	0	0	0
63	1N	10	0	0	0	0
63	1O	8	0	0	0	0
63	1P	23	0	0	1	0
63	1Q	16	0	0	0	0
63	1R	14	0	0	1	0
63	1S	4	0	0	0	0
63	1T	9	0	0	0	0
63	1U	13	0	0	0	0
63	1V	9	0	0	0	0
63	1W	11	0	0	1	0
63	1X	8	0	0	0	0
63	1Y	7	0	0	1	0
63	1Z	2	0	0	0	0
63	1a	523	0	0	0	0
63	1b	1	0	0	0	0
63	1d	3	0	0	0	0
63	1e	3	0	0	0	0
63	1g	1	0	0	0	0
63	1k	3	0	0	0	0
63	1l	10	0	0	0	0
63	1m	1	0	0	0	0
63	1o	2	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
63	1p	1	0	0	0	0
63	1q	4	0	0	0	0
63	1r	1	0	0	0	0
63	1t	2	0	0	0	0
63	1v	6	0	0	0	0
63	1w	26	0	0	0	0
63	1x	18	0	0	0	0
63	1y	2	0	0	0	0
63	20	5	0	0	0	0
63	21	14	0	0	0	0
63	22	1	0	0	0	0
63	23	2	0	0	0	0
63	25	4	0	0	0	0
63	27	7	0	0	0	0
63	28	7	0	0	0	0
63	29	1	0	0	0	0
63	2A	1602	0	0	42	0
63	2B	28	0	0	1	0
63	2D	29	0	0	0	0
63	2E	15	0	0	1	0
63	2F	20	0	0	0	0
63	2I	4	0	0	1	0
63	2N	2	0	0	0	0
63	2P	15	0	0	2	0
63	2Q	2	0	0	0	0
63	2R	5	0	0	0	0
63	2T	6	0	0	0	0
63	2U	6	0	0	0	0
63	2V	4	0	0	0	0
63	2W	4	0	0	0	0
63	2X	4	0	0	0	0
63	2Y	1	0	0	0	0
63	2Z	2	0	0	0	0
63	2a	389	0	0	0	0
63	2d	3	0	0	0	0
63	2e	3	0	0	0	0
63	2f	1	0	0	0	0
63	2g	1	0	0	0	0
63	2h	2	0	0	0	0
63	2i	1	0	0	0	0
63	2j	4	0	0	0	0
63	2l	5	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
63	2o	1	0	0	0	0
63	2p	3	0	0	0	0
63	2q	2	0	0	0	0
63	2r	1	0	0	0	0
63	2t	3	0	0	0	0
63	2u	1	0	0	0	0
63	2v	1	0	0	0	0
63	2w	3	0	0	0	0
63	2x	10	0	0	0	0
63	2y	20	0	0	0	0
63	A	4	0	0	0	0
63	B	2	0	0	0	0
All	All	302030	0	196834	1998	0

The all-atom clashscore is defined as the number of clashes found per 1000 atoms (including hydrogen atoms). The all-atom clashscore for this structure is 8.

All (1998) 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)
1:1A:1128:U:H3	1:1A:1132:A:N6	1.32	1.23
1:2A:2138:C:N4	1:2A:2153:G:H1	1.52	1.06
1:1A:1128:U:O4	1:1A:1132:A:N1	1.87	1.06
1:2A:2129:C:N4	1:2A:2159:G:H1	1.55	1.04
1:2A:2136:C:N4	1:2A:2155:G:H1	1.57	1.02
9:2N:123:TYR:HH	9:2N:130:HIS:HE2	1.02	0.99
1:1A:1104:G:H1	1:1A:1126:C:H42	0.98	0.98
1:1A:1104:G:H1	1:1A:1126:C:N4	1.67	0.92
1:1A:2122:G:H1	1:1A:2211:U:H3	1.18	0.90
1:1A:2158:C:N3	1:1A:2177:G:N2	2.23	0.87
1:1A:929:G:H1	1:1A:940:C:H42	1.21	0.87
1:2A:994:C:OP1	16:2U:53:ARG:NH2	2.07	0.87
10:1O:35:VAL:HG11	10:1O:103:ALA:HB3	1.56	0.86
1:1A:1004:A:N6	1:1A:1037:C:N3	55.32	0.86
22:20:10:THR:HG22	22:20:12:ASN:H	1.42	0.85
10:2O:35:VAL:HG11	10:2O:103:ALA:HB3	1.59	0.85
4:1E:28:ALA:HB3	4:1E:93:VAL:HG12	1.58	0.85
1:2A:1204:A:H2	1:2A:1241:A:H62	1.25	0.84
1:1A:1006:C:H42	1:1A:1023:G:H1	21.33	0.82
1:1A:2158:C:N4	1:1A:2177:G:N1	2.28	0.82
1:2A:2136:C:H42	1:2A:2155:G:H1	1.27	0.82

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:11:50:ARG:HG2	23:11:59:THR:HG22	1.63	0.81
1:2A:2807:G:N1	1:2A:2893:G:O6	2.13	0.81
1:2A:1422:G:H5''	10:2O:48:PRO:HB3	99.93	0.81
1:2A:2127:G:N1	1:2A:2161:C:C4	2.49	0.80
1:1A:715:G:N7	63:1A:4347:HOH:O	2.14	0.80
1:1A:2801:C:OP1	4:1E:61:ARG:NH2	2.15	0.80
1:1A:1111:U:O2	1:1A:1119:A:N6	2.15	0.79
1:2A:1002:G:H1	1:2A:1038:C:H42	43.35	0.79
1:2A:2138:C:N3	1:2A:2153:G:N2	2.28	0.79
1:2A:2127:G:N2	1:2A:2161:C:C2	2.51	0.79
1:2A:2079:U:OP1	23:21:21:ARG:NH2	2.15	0.79
1:2A:2127:G:C6	1:2A:2161:C:N4	2.51	0.79
2:2B:20:C:N4	2:2B:63:G:O6	2.17	0.78
1:1A:1104:G:N2	1:1A:1126:C:N3	2.30	0.78
1:2A:740:U:OP2	63:2A:3934:HOH:O	2.00	0.78
4:2E:47:VAL:HG11	4:2E:86:PRO:HD2	1.66	0.77
1:2A:631:A:OP1	11:2P:65:ARG:NH1	2.16	0.77
1:1A:325:G:OP2	20:1Y:84:ARG:NH2	2.17	0.77
11:2P:59:LEU:HD11	30:28:10:ALA:HB2	1.66	0.77
1:2A:2136:C:N4	1:2A:2155:G:N1	2.32	0.77
1:1A:973:G:N7	63:1A:4354:HOH:O	2.18	0.77
1:1A:2158:C:N4	1:1A:2177:G:H1	1.80	0.77
1:2A:2127:G:C2	1:2A:2161:C:N3	2.52	0.77
1:1A:1700:G:H3'	13:1R:2:ARG:HD3	1.66	0.77
1:1A:2460:A:OP1	63:1A:4304:HOH:O	2.01	0.76
1:1A:656:A:OP1	11:1P:65:ARG:NH1	2.18	0.76
1:2A:2129:C:H42	1:2A:2159:G:H1	0.80	0.76
1:1A:11:G:H2'	1:1A:12:U:H5''	1.68	0.76
1:2A:2129:C:N3	1:2A:2159:G:N2	2.31	0.76
1:2A:2287:A:H62	1:2A:2344:U:H3	1.32	0.76
1:2A:1530:C:O2'	1:2A:1531:C:O5'	2.03	0.76
21:2Z:145:GLU:H	21:2Z:148:ASP:HB2	1.50	0.76
8:1I:92:VAL:HG13	8:1I:120:ILE:HB	1.67	0.75
1:2A:2586:C:N4	58:B:5:ASN:HB3	2.01	0.75
1:1A:2598:C:N4	58:A:5:ASN:HB3	2.01	0.75
1:1A:927:G:H2'	1:1A:928:G:H8	1.51	0.75
5:1F:53:THR:HG23	5:1F:55:GLY:H	1.51	0.75
1:2A:882:G:H1	1:2A:894:C:H42	1.34	0.75
1:1A:1166:G:O6	63:1A:4336:HOH:O	2.04	0.74
27:25:16:ARG:NH1	27:25:17:ASP:OD1	2.19	0.74
22:10:11:ARG:O	22:10:14:ARG:NH2	2.20	0.74

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:2206:G:H3'	1:2A:2207:G:C8	2.23	0.74
1:1A:542:C:OP1	27:15:16:ARG:NH2	2.20	0.74
1:2A:1783:A:OP1	63:2A:3934:HOH:O	2.04	0.74
1:2A:2126:A:N6	1:2A:2162:G:O2'	2.17	0.74
1:1A:2149:G:H1	1:1A:2183:C:H42	1.33	0.74
3:1D:71:ASP:HB3	3:1D:103:ARG:HH12	1.53	0.74
1:1A:943:C:N3	1:1A:944:C:N4	2.36	0.74
8:1I:109:ILE:HG13	8:1I:130:TYR:CZ	2.23	0.73
28:16:13:CYS:SG	28:16:47:THR:HG21	2.28	0.73
1:1A:2641:A:O2'	1:1A:2642:G:OP2	2.07	0.73
10:1O:63:VAL:HG12	10:1O:106:LEU:HD11	1.70	0.73
14:1S:25:ARG:NH1	14:1S:42:ASP:OD1	2.21	0.73
1:2A:852:G:H2'	1:2A:853:G:H8	1.54	0.73
1:1A:2149:G:H1	1:1A:2183:C:N4	1.86	0.73
1:2A:1693:U:O2'	3:2D:14:ARG:NH2	2.22	0.73
1:2A:2104:G:H1	1:2A:2185:C:H42	1.37	0.73
5:2F:185:ASP:HA	5:2F:188:ARG:HD3	1.71	0.73
1:1A:1053:C:OP2	63:1A:4337:HOH:O	2.07	0.73
1:1A:1165:C:N4	63:1A:4356:HOH:O	2.20	0.73
31:29:25:VAL:HB	31:29:34:GLN:HB2	1.70	0.72
2:2B:22:U:H3	2:2B:61:G:H1	1.36	0.72
1:2A:1801:G:OP2	3:2D:154:LYS:NZ	2.22	0.72
11:1P:126:VAL:HG12	11:1P:148:LEU:HD23	1.70	0.72
1:2A:1689:A:H62	1:2A:1698:A:H2	1.35	0.72
9:2N:128:HIS:O	9:2N:131:GLN:NE2	2.22	0.72
1:1A:1613:A:OP1	3:1D:211:ARG:NH1	2.23	0.72
17:1V:40:LEU:HB2	17:1V:46:VAL:HG13	1.72	0.72
11:1P:91:PHE:O	11:1P:121:LYS:NZ	2.23	0.71
1:2A:1153:C:OP1	16:2U:92:ARG:NH2	2.24	0.71
5:2F:53:THR:HG23	5:2F:55:GLY:H	1.54	0.71
11:1P:50:ARG:HD3	30:18:7:HIS:CD2	2.25	0.71
1:2A:531:C:OP1	1:2A:561:G:N1	2.23	0.71
1:1A:1006:C:N3	1:1A:1023:G:N2	22.81	0.71
19:1X:35:THR:HG22	19:1X:38:GLU:HB2	1.71	0.71
1:2A:1798:U:H5'	3:2D:259:THR:HG22	1.72	0.71
26:14:55:ARG:H	26:14:56:VAL:HA	1.56	0.70
23:21:51:VAL:HG11	23:21:74:VAL:HG21	1.72	0.70
21:2Z:106:GLY:HA3	21:2Z:141:VAL:HG22	1.74	0.70
1:2A:2782:G:OP2	63:2A:3936:HOH:O	2.09	0.70
2:1B:33:G:H5'	6:1G:2:PRO:HD3	1.74	0.70
6:1G:45:GLU:OE2	63:1G:5001:HOH:O	2.10	0.70

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:2136:C:N3	1:2A:2155:G:N2	2.34	0.70
24:12:55:ARG:NH2	63:12:3101:HOH:O	2.25	0.69
13:1R:67:LEU:HD13	13:1R:76:VAL:HG21	1.73	0.69
1:2A:2127:G:C2	1:2A:2161:C:C4	2.80	0.69
1:2A:863:A:H2'	1:2A:864:G:H8	1.56	0.69
1:1A:1044:C:OP1	63:1A:4338:HOH:O	2.09	0.69
6:2G:15:VAL:HG22	6:2G:175:LEU:HB3	1.74	0.69
11:1P:100:LEU:HD12	11:1P:112:LEU:HD11	1.75	0.69
1:2A:400:G:N7	63:2A:3974:HOH:O	2.25	0.69
11:1P:59:LEU:HD11	30:18:10:ALA:HB2	1.75	0.69
1:1A:1485:A:OP1	63:1A:4339:HOH:O	2.10	0.69
1:1A:1059:C:OP2	63:1A:4301:HOH:O	2.11	0.69
1:2A:2127:G:N1	1:2A:2161:C:N4	2.41	0.68
1:2A:2805:G:H2'	1:2A:2807:G:C8	2.29	0.68
1:2A:376:C:OP1	63:2A:3937:HOH:O	2.12	0.68
1:1A:2101:U:OP1	23:11:21:ARG:NH2	2.23	0.68
1:2A:2712(A):A:OP2	63:2A:3938:HOH:O	2.12	0.68
1:2A:1434:A:H61	1:2A:1558:A:H62	1.41	0.68
1:1A:1151:U:H2'	1:1A:1152:G:H8	1.57	0.68
1:2A:2448:A:N6	63:2A:3982:HOH:O	2.26	0.68
23:21:50:ARG:HG2	23:21:59:THR:HG22	1.74	0.68
4:2E:48:GLN:HE21	4:2E:78:LEU:HD23	1.57	0.68
1:1A:2124:U:H3	1:1A:2209:G:H1	1.42	0.68
4:1E:163:GLU:OE2	63:1E:401:HOH:O	2.11	0.68
1:2A:1796:U:H2'	1:2A:1797:C:C6	2.29	0.68
1:2A:2138:C:H42	1:2A:2153:G:H1	0.76	0.68
1:1A:2227:G:H3'	1:1A:2228:G:C8	2.29	0.67
1:1A:737:G:O6	63:1A:4340:HOH:O	2.11	0.67
8:1I:93:THR:H	8:1I:96:ASP:HB2	1.60	0.67
19:1X:57:LEU:HD11	19:1X:78:LYS:HE2	1.76	0.67
26:24:46:GLN:HE21	26:24:48:ARG:HH21	1.40	0.67
19:2X:35:THR:HG22	19:2X:38:GLU:H	1.59	0.67
6:2G:179:PRO:HB2	26:24:42:PHE:HE2	1.58	0.67
1:1A:2408:G:N7	63:1A:4382:HOH:O	2.26	0.67
1:1A:931:C:H42	1:1A:938:G:H1	1.43	0.67
4:1E:105:THR:OG1	4:1E:199:ARG:NH2	2.28	0.67
1:1A:1069:U:OP2	63:1A:4341:HOH:O	2.11	0.67
21:2Z:72:ARG:NH2	21:2Z:97:GLU:O	2.27	0.67
1:2A:2504:U:OP2	63:2A:3941:HOH:O	2.13	0.67
1:2A:2624:G:N7	63:2A:3984:HOH:O	2.26	0.67
2:2B:41:U:H5	6:2G:70:VAL:H	1.43	0.67

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:2B:106:G:H5'	21:2Z:31:ARG:HG2	1.75	0.67
26:14:16:CYS:SG	26:14:17:GLY:N	2.67	0.67
1:1A:1310:G:OP1	27:15:19:ARG:NH2	2.19	0.67
1:2A:1253:A:OP1	63:2A:3940:HOH:O	2.12	0.67
1:1A:1464:G:OP2	63:1A:4343:HOH:O	2.13	0.66
2:2B:42:C:H2'	6:2G:66:GLN:HE21	1.59	0.66
28:16:37:ARG:NH1	63:16:5001:HOH:O	2.29	0.66
1:2A:981:A:OP1	63:2A:3939:HOH:O	2.12	0.66
1:1A:2362:C:OP2	63:1A:4342:HOH:O	2.12	0.66
1:1A:1037:C:H2'	1:1A:1038:C:H6	2.14	0.66
1:1A:1055:A:OP2	9:1N:37:LYS:NZ	2.26	0.66
1:2A:1816:G:O6	3:2D:35:LYS:NZ	2.26	0.66
1:2A:863:A:H2'	1:2A:864:G:C8	2.31	0.66
12:1Q:111:GLU:OE2	12:1Q:133:ARG:NH2	2.29	0.66
23:21:3:LYS:HB2	23:21:61:ARG:HH22	1.60	0.66
1:2A:2755:C:H2'	31:29:19:ARG:HD3	1.78	0.66
9:2N:34:LEU:HD21	9:2N:120:LEU:HB2	1.78	0.66
20:1Y:102:CYS:SG	20:1Y:103:GLY:N	2.67	0.66
1:2A:2148:G:H2'	1:2A:2149:G:C8	2.31	0.66
1:2A:2839:G:H5'	13:2R:46:GLY:HA2	1.78	0.66
1:1A:436:C:OP1	63:1A:4302:HOH:O	2.13	0.66
1:2A:276:A:H5''	1:2A:277:C:H5'	1.76	0.66
1:1A:1016:C:OP2	63:1A:4345:HOH:O	2.13	0.66
1:1A:2459:G:OP2	63:1A:4344:HOH:O	2.13	0.66
1:2A:1627:G:OP1	63:2A:3943:HOH:O	2.14	0.66
1:2A:266:G:H5''	1:2A:268:C:H41	11.54	0.66
1:2A:1125:G:H5'	31:29:37:GLY:HA2	1.77	0.65
1:2A:637:A:OP1	11:2P:133:SER:OG	2.14	0.65
17:2V:43:GLU:OE1	17:2V:43:GLU:N	2.28	0.65
1:1A:2133:C:N3	1:1A:2167:C:O2'	2.25	0.65
1:1A:2185:C:OP1	1:1A:2187:G:N2	2.29	0.65
1:1A:455:A:OP1	63:1A:4346:HOH:O	2.14	0.65
1:2A:2049:G:N7	63:2A:3993:HOH:O	2.29	0.65
1:2A:2206:G:H3'	1:2A:2207:G:H8	1.58	0.65
1:2A:330:A:H2	1:2A:1210:A:HO2'	1.43	0.65
1:2A:2445:G:OP1	5:2F:74:ARG:NH2	2.29	0.65
18:2W:34:ASN:OD1	18:2W:37:ARG:NH2	2.29	0.65
1:1A:1101:G:H1	1:1A:1150:C:H42	1.45	0.65
1:2A:1448:G:N7	63:2A:3998:HOH:O	2.29	0.65
13:2R:67:LEU:HD13	13:2R:76:VAL:HG21	1.78	0.65
1:2A:1022:G:H22	1:2A:1142(A):A:H2	1.42	0.65

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:2125:G:H1'	1:2A:2173:A:H61	1.61	0.65
1:2A:143:G:H4'	19:2X:35:THR:HG21	1.77	0.65
1:2A:2242:G:OP1	63:2A:3945:HOH:O	2.14	0.65
1:2A:825:C:O2	11:2P:55:ARG:NH1	2.29	0.65
1:2A:1568:G:N7	63:2A:4005:HOH:O	2.30	0.64
1:2A:2364:C:OP1	22:20:55:ARG:NH1	2.29	0.64
1:1A:1435:G:H2'	1:1A:1436:U:C6	2.93	0.64
1:1A:2071:G:N7	63:1A:4394:HOH:O	2.29	0.64
1:1A:287:G:O2'	63:1A:4349:HOH:O	2.15	0.64
1:2A:2206:G:H5''	1:2A:2207:G:N7	2.12	0.64
1:2A:2371:G:O2'	28:26:46:HIS:ND1	2.29	0.64
1:2A:851:U:O2'	25:23:42:ALA:O	2.12	0.64
21:2Z:154:ASP:N	21:2Z:154:ASP:OD1	2.30	0.64
1:2A:2060:A:N3	63:2A:4001:HOH:O	2.29	0.64
1:1A:1431:G:O2'	1:1A:1442:U:O2	2.14	0.64
1:1A:2297:C:OP2	28:16:6:ARG:NH1	2.30	0.64
1:1A:1072:U:OP1	63:1A:4341:HOH:O	2.14	0.64
1:1A:2150:C:H2'	1:1A:2151:C:O4'	1.97	0.64
1:1A:739:C:O2'	3:1D:38:LYS:NZ	2.30	0.64
5:1F:185:ASP:HA	5:1F:188:ARG:HD3	1.80	0.64
1:2A:1958:C:OP2	63:2A:3944:HOH:O	2.14	0.64
4:2E:48:GLN:NE2	4:2E:78:LEU:HD23	2.13	0.64
1:1A:2255:U:OP1	63:1A:4348:HOH:O	2.14	0.64
17:2V:98:GLU:OE2	17:2V:100:ARG:NH1	2.30	0.64
1:1A:2289:G:OP2	22:10:10:THR:HG21	1.97	0.64
1:1A:2143:G:H1	1:1A:2199:C:H42	1.45	0.64
4:1E:111:ARG:HG3	4:1E:160:TYR:CD2	2.33	0.64
1:2A:2127:G:N2	1:2A:2161:C:N3	2.46	0.64
1:2A:2448:A:OP1	63:2A:3942:HOH:O	2.13	0.64
7:2H:20:ALA:HB3	7:2H:23:ARG:HG3	1.78	0.64
1:1A:2695:C:O2	10:1O:70:LYS:NZ	2.21	0.63
1:2A:955:C:OP1	12:2Q:87:LYS:HE3	1.98	0.63
1:1A:1085:G:H1	1:1A:1162:C:H42	1.46	0.63
5:2F:143:ALA:HB1	5:2F:148:LEU:HB2	1.78	0.63
1:2A:34:C:H2'	1:2A:35:G:H8	4.17	0.63
1:2A:907:U:H4'	12:2Q:101:ARG:HH22	1.61	0.63
15:2T:117:ASP:OD2	15:2T:120:ARG:NE	2.24	0.63
1:2A:994:C:O2'	1:2A:996:A:OP1	2.11	0.63
2:2B:87:G:N2	2:2B:90:A:OP2	2.30	0.63
1:1A:2803:A:N3	1:1A:2803:A:H3'	2.14	0.63
8:1I:130:TYR:HB3	8:1I:138:ILE:HB	1.79	0.63

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:1420:U:O2'	1:2A:1421:G:OP1	2.15	0.63
1:2A:692:C:O2'	3:2D:38:LYS:NZ	2.30	0.63
4:2E:59:VAL:HG21	4:2E:74:PRO:HB3	1.81	0.63
1:2A:1002:G:H1	1:2A:1038:C:N4	43.06	0.63
1:2A:1793:C:OP1	63:2A:3946:HOH:O	2.16	0.63
7:1H:17:VAL:HG22	7:1H:26:VAL:HG22	1.81	0.63
1:1A:1040:C:OP1	16:1U:53:ARG:NH2	2.32	0.63
1:2A:1007:C:OP1	9:2N:35:ARG:NH1	2.32	0.63
5:2F:28:ILE:HG12	5:2F:112:MET:HG2	1.78	0.62
1:1A:2147:G:N1	1:1A:2194:U:OP1	2.21	0.62
1:1A:1115:A:H4'	1:1A:1116:A:H8	1.64	0.62
1:1A:1123:A:H2'	1:1A:1124:U:H4'	1.81	0.62
19:1X:31:HIS:CD2	19:1X:33:LYS:H	2.18	0.62
1:2A:2136:C:O2'	1:2A:2137:C:O5'	2.16	0.62
1:2A:2296:U:OP2	14:2S:9:ARG:NH2	2.33	0.62
1:1A:2658:C:OP2	1:1A:2745:G:O2'	2.17	0.62
1:2A:2143:C:H2'	1:2A:2144:U:O4'	1.99	0.62
1:2A:2683:C:OP1	15:2T:53:ARG:NH2	2.31	0.62
11:2P:50:ARG:HD3	30:28:7:HIS:CD2	2.35	0.62
1:1A:1115:A:H4'	1:1A:1116:A:C8	2.35	0.62
1:1A:1346:U:H4'	1:1A:1347:A:H5''	1.82	0.62
2:2B:24:G:N7	2:2B:56:G:H2'	2.15	0.62
1:2A:1311:G:H2'	29:27:47:ARG:HH22	1.65	0.62
1:1A:1151:U:H2'	1:1A:1152:G:C8	2.33	0.62
1:2A:2727:G:O2'	10:2O:70:LYS:NZ	2.31	0.62
4:2E:28:ALA:HB3	4:2E:93:VAL:HG12	1.82	0.62
21:1Z:72:ARG:NH2	21:1Z:97:GLU:O	2.33	0.61
1:2A:2690:C:OP1	13:2R:17:ARG:NH2	2.32	0.61
10:1O:122:LEU:HD13	15:1T:72:VAL:HG11	1.81	0.61
2:1B:106:G:H5'	21:1Z:31:ARG:HG2	1.82	0.61
1:2A:2144:U:H1'	1:2A:2148:G:N2	2.15	0.61
7:1H:3:ARG:HH22	7:1H:65:HIS:HB3	1.64	0.61
1:2A:1631:C:H2'	63:2A:4620:HOH:O	1.99	0.61
26:24:16:CYS:SG	26:24:17:GLY:N	2.73	0.61
1:2A:900:A:O2'	1:2A:901:A:OP1	2.17	0.61
7:2H:125:VAL:HG22	7:2H:131:VAL:HG22	1.82	0.61
7:2H:113:VAL:HG11	7:2H:151:ILE:HG21	1.83	0.61
1:1A:1232:G:H5''	17:1V:81:TYR:CE1	2.35	0.61
1:1A:1405:A:H2	1:1A:1418:U:O4	1.84	0.61
1:2A:2498:C:H3'	63:2A:3926:HOH:O	1.97	0.61
21:2Z:146:ILE:HG12	21:2Z:174:VAL:HG13	1.82	0.61

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:2457:G:OP1	5:1F:74:ARG:NH2	2.32	0.61
31:29:2:LYS:NZ	31:29:31:LYS:O	2.33	0.61
1:1A:1006:C:N4	1:1A:1023:G:H1	21.92	0.61
1:1A:1740:U:O2'	3:1D:14:ARG:NH2	2.33	0.61
1:1A:928:G:N2	1:1A:943:C:O2	2.34	0.61
3:1D:147:LEU:HD13	3:1D:155:LEU:HD21	1.82	0.61
7:2H:23:ARG:NH1	7:2H:34:GLU:OE1	2.33	0.61
24:12:32:LEU:HD22	24:12:36:ARG:HH11	1.66	0.60
1:1A:1829:U:H5'	3:1D:259:THR:HG22	1.81	0.60
1:2A:467:G:OP2	29:27:34:ARG:HD3	2.01	0.60
1:2A:2268:A:OP1	63:2A:3947:HOH:O	2.16	0.60
6:2G:64:THR:HB	6:2G:94:LEU:HD21	1.83	0.60
1:2A:884:C:N3	1:2A:893:C:O2'	2.34	0.60
1:2A:1769:G:O2'	1:2A:1958:C:OP1	2.17	0.60
18:2W:57:ASN:HA	18:2W:61:ASN:HD22	1.65	0.60
21:1Z:152:ALA:HB1	21:1Z:163:LEU:HD11	1.83	0.60
3:2D:17:THR:O	3:2D:211:ARG:NH2	2.34	0.60
5:2F:101:LEU:O	5:2F:106:ARG:NH1	2.34	0.60
1:2A:1286:A:C8	1:2A:1287:A:H4'	8.14	0.60
19:2X:57:LEU:HD11	19:2X:78:LYS:HE2	1.84	0.60
1:2A:2741:A:OP1	31:29:22:ARG:NH2	2.32	0.60
1:2A:1031:G:H5''	31:29:8:LYS:HE3	1.83	0.60
11:2P:26:GLY:O	63:2P:301:HOH:O	2.16	0.60
1:1A:2182:G:O6	1:1A:2183:C:N4	2.35	0.60
1:1A:1219:A:H4'	1:1A:1220:U:OP1	2.01	0.60
7:1H:94:TYR:OH	7:1H:152:ARG:NH1	2.34	0.60
1:2A:854:G:H2'	1:2A:855:G:H8	1.67	0.60
1:1A:1006:C:N4	1:1A:1023:G:N1	22.35	0.59
1:1A:1385:G:N3	63:1A:4402:HOH:O	2.32	0.59
1:1A:2015:U:OP2	63:1A:4350:HOH:O	2.16	0.59
1:1A:625:G:O2'	1:1A:702:A:N6	2.35	0.59
2:1B:105:A:OP1	21:1Z:72:ARG:NH1	2.34	0.59
23:21:83:GLU:N	23:21:83:GLU:OE1	2.34	0.59
1:2A:307:G:H21	1:2A:330:A:H62	1.49	0.59
1:2A:784:A:C6	3:2D:229:VAL:HG11	2.36	0.59
15:2T:65:LYS:HE3	15:2T:67:SER:HB2	1.84	0.59
1:1A:1104:G:N2	1:1A:1127:U:O2	2.35	0.59
1:2A:947:G:H2'	1:2A:948:G:C8	2.37	0.59
5:2F:184:TYR:CE2	5:2F:188:ARG:HD2	2.36	0.59
1:2A:1452:A:OP2	63:2A:3948:HOH:O	2.16	0.59
8:2I:3:VAL:HG12	8:2I:38:LEU:HA	1.83	0.59

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:2110:G:H3'	1:2A:2111:C:H5'	1.84	0.59
1:2A:2171:A:N3	1:2A:2172:U:N3	2.50	0.59
11:2P:100:LEU:HD12	11:2P:112:LEU:HD11	1.84	0.59
19:1X:2:LYS:NZ	19:1X:38:GLU:OE2	2.29	0.59
5:2F:25:PRO:HD2	5:2F:115:ALA:HB2	1.84	0.59
26:24:24:THR:OG1	26:24:25:TYR:N	2.30	0.59
12:2Q:16:ARG:O	12:2Q:18:LYS:N	2.31	0.59
1:1A:843:C:H2'	1:1A:844:C:C6	2.38	0.59
8:1I:101:LEU:HD22	8:1I:107:VAL:HB	1.84	0.59
17:1V:76:LYS:HB2	17:1V:81:TYR:HB3	1.83	0.59
20:1Y:55:TYR:H	20:1Y:56:PRO:HD3	1.68	0.59
7:2H:159:GLU:HG3	7:2H:169:VAL:HG11	1.84	0.59
1:1A:1140:U:H1'	1:1A:1143:U:H5	1.68	0.59
12:2Q:111:GLU:O	12:2Q:115:MET:HG2	2.02	0.59
13:2R:97:VAL:HG22	13:2R:114:VAL:HG13	1.84	0.59
1:1A:1108:G:H1	1:1A:1123:A:H61	1.49	0.59
1:2A:2022:U:O2'	1:2A:2617:C:H5'	2.03	0.59
1:2A:2152:G:C2	1:2A:2153:G:H1'	2.38	0.59
1:2A:2857:G:N2	1:2A:2860:A:OP2	2.34	0.59
19:2X:60:ARG:HH22	29:27:47:ARG:HH12	1.51	0.59
1:1A:2849:G:H5'	13:1R:46:GLY:HA2	1.85	0.58
1:1A:1000:C:OP1	12:1Q:87:LYS:HE3	2.03	0.58
1:2A:2355:C:H1'	22:20:39:ARG:HH21	1.68	0.58
1:2A:300:A:OP2	20:2Y:86:ARG:NH2	2.36	0.58
1:1A:1117:G:H1'	1:1A:1135:G:H2'	1.83	0.58
6:1G:16:ARG:HB2	6:1G:17:PRO:HD3	1.85	0.58
25:23:6:VAL:HG22	25:23:56:VAL:HG13	1.85	0.58
1:2A:1266:G:O5'	18:2W:15:ARG:NH2	2.36	0.58
30:18:33:ASN:HA	30:18:36:LYS:HD2	1.85	0.58
1:1A:1071:G:C4	1:1A:1180:C:H1'	2.38	0.58
1:2A:1634:A:OP1	63:2A:3949:HOH:O	2.16	0.58
3:2D:71:ASP:HB2	3:2D:103:ARG:HH12	1.68	0.58
1:2A:2749:A:OP1	7:2H:3:ARG:NH1	2.36	0.58
1:1A:303:C:H42	1:1A:385:G:H1	1.50	0.58
1:2A:2104:G:H1	1:2A:2185:C:N4	2.00	0.58
1:2A:34:C:H2'	1:2A:35:G:C8	4.97	0.58
22:10:27:GLU:HG3	22:10:68:GLU:HA	1.85	0.58
1:1A:1115:A:H1'	1:1A:1142:A:H4'	1.86	0.58
1:1A:1398:U:OP2	63:1A:4351:HOH:O	2.17	0.58
1:1A:2804:C:H2'	1:1A:2805:G:C8	2.39	0.58
15:1T:29:ARG:NH1	15:1T:46:GLU:OE1	2.35	0.58

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:2H:3:ARG:NH1	7:2H:4:ILE:H	2.01	0.58
1:1A:2348:A:H61	22:10:43:THR:HG22	1.69	0.58
1:1A:2825:C:H5'	27:15:29:THR:HG21	1.85	0.58
1:2A:1442:G:N3	1:2A:1442:G:H2'	2.86	0.58
1:1A:276:C:OP1	8:1I:45:LYS:NZ	2.22	0.58
8:1I:75:LEU:HD22	8:1I:105:HIS:CD2	2.39	0.58
21:1Z:24:LEU:HD22	21:1Z:41:LEU:HD12	1.86	0.58
1:2A:2693:A:H2'	1:2A:2694:G:C8	2.39	0.58
11:1P:98:GLU:O	11:1P:101:VAL:HG12	2.04	0.58
30:28:23:VAL:HG11	30:28:47:LYS:HD3	1.86	0.58
1:2A:861:A:N3	2:2B:79:C:O2'	2.35	0.58
1:2A:2529:G:O6	31:29:31:LYS:NZ	2.37	0.57
1:2A:2176:A:H2'	1:2A:2177:C:C6	2.39	0.57
1:2A:2058:A:N7	63:2A:4016:HOH:O	2.32	0.57
1:1A:1108:G:P	1:1A:1116:A:H1'	2.44	0.57
15:1T:109:GLU:O	15:1T:113:LYS:HG2	2.05	0.57
1:2A:2317:C:N4	1:2A:2318:G:O6	2.36	0.57
1:2A:271(H):G:H2'	1:2A:271(I):G:C8	2.39	0.57
1:1A:2486:C:OP1	63:1A:4353:HOH:O	2.17	0.57
7:1H:164:TYR:HB2	7:1H:167:GLU:HB2	1.86	0.57
22:20:63:VAL:HG21	22:20:83:PRO:HG3	1.85	0.57
1:2A:2345:G:N3	1:2A:2381:C:H2'	2.19	0.57
1:2A:2693:A:H2'	1:2A:2694:G:H8	1.67	0.57
1:1A:1140:U:H1'	1:1A:1143:U:C5	2.39	0.57
1:1A:1211:U:H2'	1:1A:1212:C:C6	2.40	0.57
15:1T:24:PRO:HA	15:1T:49:VAL:HG23	1.85	0.57
22:20:26:TYR:N	22:20:29:GLN:OE1	2.31	0.57
1:2A:2343:C:HO2'	1:2A:2373:G:HO2'	1.47	0.57
1:2A:1019:U:H3	1:2A:1142(A):A:H62	1.52	0.57
4:2E:181:LEU:HD11	15:2T:6:LEU:HD23	1.87	0.57
11:2P:95:VAL:HG13	11:2P:125:VAL:HA	1.86	0.57
1:1A:1004:A:H5''	1:1A:1024:G:H1	28.05	0.57
6:1G:18:GLU:OE2	6:1G:21:ARG:NH1	2.38	0.57
1:2A:2507:C:H2'	1:2A:2508:G:O4'	2.03	0.57
1:2A:2788:C:OP1	4:2E:61:ARG:NH2	2.37	0.57
1:2A:75:G:H4'	24:22:55:ARG:NH1	2.20	0.57
4:1E:12:THR:HG21	15:1T:11:GLU:OE2	2.04	0.57
1:1A:957:A:H2'	12:1Q:9:TYR:OH	2.05	0.57
20:1Y:34:LYS:NZ	63:1Y:601:HOH:O	2.36	0.57
1:2A:2590:A:O3'	3:2D:239:ARG:NH2	2.38	0.57
8:2I:93:THR:H	8:2I:96:ASP:HB2	1.69	0.57

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:2Z:10:ARG:NH2	21:2Z:26:GLY:O	2.37	0.57
11:1P:82:GLY:HA2	11:1P:113:LYS:O	2.05	0.57
14:1S:61:ASN:HB3	14:1S:64:GLU:HB2	1.87	0.57
1:2A:1340:U:OP1	19:2X:16:LYS:NZ	2.38	0.57
3:2D:148:GLU:HB2	3:2D:151:LYS:HD2	1.87	0.57
5:2F:195:ASP:HB3	5:2F:198:ALA:H	1.70	0.57
1:1A:1159:U:H2'	1:1A:1160:G:H8	1.69	0.56
1:1A:1827:U:H2'	1:1A:1828:C:C6	2.39	0.56
1:1A:469:A:H1'	1:1A:1246:C:O4'	2.05	0.56
11:1P:42:SER:O	63:1P:301:HOH:O	2.18	0.56
1:2A:1015:G:H2'	1:2A:1016:G:H8	1.70	0.56
1:2A:1495:A:H2'	1:2A:1496:A:C8	2.41	0.56
1:1A:1133:G:H2'	1:1A:1135:G:C8	2.39	0.56
1:1A:302:A:H2'	1:1A:303:C:C6	2.40	0.56
21:2Z:153:SER:HB3	21:2Z:167:PRO:HB3	1.87	0.56
1:1A:1831:C:OP1	3:1D:260:ARG:NH2	2.38	0.56
9:1N:15:LEU:HD22	9:1N:137:LYS:HD2	1.85	0.56
18:1W:58:ALA:HB1	18:1W:64:MET:HB2	1.88	0.56
1:1A:2262:G:OP1	12:1Q:85:LYS:NZ	2.38	0.56
1:2A:1030:G:OP2	12:2Q:128:LYS:NZ	2.39	0.56
1:2A:796:C:H2'	1:2A:797:C:C6	2.40	0.56
1:1A:297:C:H2'	1:1A:298:G:H8	1.70	0.56
3:1D:12:SER:HB3	3:1D:208:LYS:HB3	1.87	0.56
5:1F:178:PRO:O	5:1F:205:ARG:NH2	2.38	0.56
26:24:41:PRO:HG3	26:24:49:PHE:CZ	2.40	0.56
1:2A:1889:A:H2'	1:2A:1890:A:C8	2.39	0.56
1:2A:1278:A:OP1	13:2R:36:THR:HG23	2.05	0.56
1:2A:1188:U:H4'	17:2V:79:VAL:HG22	1.87	0.56
1:1A:1094:A:N1	1:1A:1158:G:O2'	2.34	0.56
1:1A:1139:G:H3'	1:1A:1140:U:H5''	1.87	0.56
1:1A:776:G:C6	3:1D:208:LYS:HB2	2.40	0.56
1:2A:2404:C:O3'	11:2P:77:ARG:NH2	2.39	0.56
1:2A:483:A:O2'	20:2Y:49:VAL:O	2.15	0.56
21:2Z:45:ASP:OD1	21:2Z:49:ARG:NE	2.35	0.56
1:1A:1102:G:H4'	1:1A:1132:A:H8	1.71	0.56
1:1A:939:C:H2'	1:1A:940:C:C6	2.40	0.56
5:1F:140:LEU:HD11	5:1F:170:LEU:HD11	1.88	0.56
1:2A:668:G:H5'	1:2A:669:G:OP2	2.04	0.56
14:2S:14:VAL:O	14:2S:18:ILE:HG12	2.05	0.56
19:2X:12:VAL:HG22	19:2X:29:TRP:CE2	2.40	0.56
1:1A:976:G:H5'	1:1A:1358:U:O2'	103.20	0.56

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:1874:C:H5'	3:1D:253:GLN:OE1	2.05	0.56
1:1A:611:U:H2'	1:1A:612:C:C6	2.41	0.56
1:2A:2469:A:H5'	1:2A:2470:G:OP2	2.06	0.56
1:2A:928:G:H8	1:2A:928:G:O5'	1.89	0.56
28:16:6:ARG:NH1	28:16:26:ASN:HB2	2.21	0.56
10:2O:2:ILE:HD12	10:2O:6:THR:HG21	1.87	0.56
1:1A:1159:U:H2'	1:1A:1160:G:C8	2.41	0.55
30:28:33:ASN:HA	30:28:36:LYS:HD2	1.88	0.55
1:2A:1119:C:H2'	1:2A:1120:G:C8	3.25	0.55
1:2A:84:A:H5''	20:2Y:8:LYS:HE3	1.88	0.55
1:1A:1633:A:H2'	1:1A:1634:C:C6	2.40	0.55
1:2A:566:U:H5''	11:2P:29:LYS:HE3	1.88	0.55
1:2A:882:G:H2'	1:2A:883:G:H8	1.71	0.55
15:2T:127:ALA:C	15:2T:129:ARG:H	2.10	0.55
20:2Y:44:ILE:HA	20:2Y:63:LYS:O	2.06	0.55
1:2A:1113:U:H2'	1:2A:1114:G:C8	2.41	0.55
1:2A:1899:G:N3	1:2A:1899:G:H2'	2.22	0.55
3:2D:8:PRO:HB3	3:2D:14:ARG:HG2	1.89	0.55
1:1A:1475:G:H2'	1:1A:1476:C:C6	2.40	0.55
1:1A:1634:C:H2'	1:1A:1635:C:H6	1.70	0.55
21:1Z:92:SER:O	21:1Z:130:PRO:HG2	2.06	0.55
1:2A:1507:A:O2'	1:2A:1508:A:O5'	2.25	0.55
1:2A:271(K):U:O2	8:2I:50:ARG:NE	2.39	0.55
3:1D:37:LEU:HD12	3:1D:62:TYR:HB2	1.87	0.55
17:1V:72:VAL:HG13	17:1V:85:LYS:HG2	1.88	0.55
28:26:6:ARG:NH1	28:26:26:ASN:HB2	2.21	0.55
1:2A:2507:C:H5''	1:2A:2573:C:N4	2.22	0.55
2:2B:14:U:OP2	2:2B:70:C:O2'	2.21	0.55
1:1A:1071:G:O2'	63:1A:4341:HOH:O	2.17	0.55
1:1A:2442:A:H2'	1:1A:2442:A:N3	2.21	0.55
4:1E:47:VAL:HG11	4:1E:86:PRO:HD2	1.89	0.55
18:1W:71:VAL:HA	18:1W:107:LEU:HD12	1.89	0.55
26:24:40:HIS:HB3	26:24:43:TYR:HB2	1.89	0.55
1:2A:143:G:H2'	1:2A:143(A):C:C6	2.42	0.55
7:2H:144:VAL:O	7:2H:148:ILE:HG13	2.07	0.55
1:1A:1039:G:OP1	16:1U:50:ARG:NH2	2.39	0.55
1:2A:2183:C:H2'	1:2A:2184:G:H8	1.72	0.55
8:2I:129:THR:HA	8:2I:138:ILE:O	2.07	0.55
9:2N:123:TYR:HH	9:2N:130:HIS:CD2	2.22	0.55
1:1A:1452:U:H2'	1:1A:1453:C:C6	2.42	0.55
1:1A:215:G:H21	1:1A:217:A:H62	1.54	0.55

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:1786:A:H1'	1:2A:1938:A:N6	2.21	0.55
9:2N:30:ILE:HG22	9:2N:34:LEU:HD22	1.88	0.55
11:1P:59:LEU:HD21	30:18:10:ALA:HA	1.88	0.55
1:1A:929:G:H3'	1:1A:930:G:H8	1.72	0.55
3:1D:145:VAL:HG11	3:1D:175:LEU:HD11	1.88	0.55
8:1I:3:VAL:HG12	8:1I:38:LEU:HA	1.89	0.55
1:2A:1028:A:N6	1:2A:1125:G:H2'	2.22	0.55
1:2A:1149:G:H2'	1:2A:1150:C:C6	2.42	0.55
18:1W:2:GLU:OE2	18:1W:72:LYS:NZ	2.25	0.54
1:2A:223:A:O2'	1:2A:420:C:O2	2.26	0.54
1:2A:322:A:OP2	5:2F:169:ASN:HB2	2.07	0.54
30:18:42:ARG:HD2	63:18:202:HOH:O	2.07	0.54
1:1A:1068:G:OP2	1:1A:1068:G:H8	6.81	0.54
1:1A:173:C:H2'	1:1A:174:U:C6	2.41	0.54
1:2A:1119:C:H2'	1:2A:1120:G:H8	2.53	0.54
1:2A:307:G:N1	1:2A:310:A:OP2	2.37	0.54
1:2A:652(T):C:H2'	1:2A:652(U):G:H8	1.73	0.54
1:2A:857:C:OP2	22:20:77:ARG:NH2	2.40	0.54
1:2A:859:G:N2	1:2A:917:A:OP2	2.39	0.54
21:1Z:149:SER:OG	21:1Z:171:ILE:O	2.24	0.54
5:2F:140:LEU:HD11	5:2F:170:LEU:HD11	1.88	0.54
6:2G:72:ARG:NH1	6:2G:87:PRO:HG3	2.22	0.54
19:2X:60:ARG:HH22	29:27:47:ARG:NH1	2.06	0.54
7:1H:33:LEU:HD21	7:1H:136:ILE:HG13	1.88	0.54
1:2A:2148:G:H2'	1:2A:2149:G:H8	1.72	0.54
1:2A:2698:U:H2'	1:2A:2699:C:C6	2.43	0.54
6:2G:114:ILE:HA	6:2G:140:ILE:HD11	1.90	0.54
7:2H:137:ASP:HB3	7:2H:140:LYS:HB3	1.89	0.54
7:1H:126:PRO:HG2	7:1H:130:ARG:HH21	1.71	0.54
7:1H:88:LEU:HD23	7:1H:130:ARG:HG3	1.90	0.54
16:1U:76:TYR:CZ	16:1U:80:ILE:HG13	2.43	0.54
1:2A:2133:G:HO2'	1:2A:2157:G:N2	2.05	0.54
7:2H:113:VAL:HG11	7:2H:151:ILE:HD13	1.87	0.54
11:2P:42:SER:O	63:2P:302:HOH:O	2.19	0.54
15:1T:108:ARG:HH22	15:1T:112:ARG:HD3	1.71	0.54
1:2A:96:G:H4'	24:22:48:HIS:CD2	2.43	0.54
25:23:10:LYS:HB3	25:23:53:LEU:HA	1.89	0.54
1:2A:1379:A:H4'	1:2A:1380:G:OP2	2.06	0.54
1:2A:2114:A:N6	1:2A:2115:G:H21	2.05	0.54
1:2A:1507:A:O2'	1:2A:1508:A:O4'	2.25	0.54
6:2G:137:GLU:HG2	6:2G:152:LEU:HD22	1.88	0.54

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
9:2N:67:LEU:HB3	9:2N:88:GLU:HG3	1.90	0.54
1:1A:715:G:H5'	1:1A:716:G:OP2	2.08	0.54
1:1A:1829:U:H5'	3:1D:259:THR:CG2	2.37	0.54
1:2A:995:C:O2	9:2N:3:THR:OG1	2.22	0.54
5:2F:123:LEU:HD13	5:2F:192:LEU:HD13	1.89	0.54
6:2G:16:ARG:O	6:2G:20:ILE:HG13	2.08	0.54
12:2Q:36:ALA:HB2	12:2Q:103:MET:SD	2.48	0.54
1:2A:674:G:O2'	5:2F:74:ARG:HD3	2.07	0.54
7:2H:124:GLU:OE2	7:2H:132:ARG:HD2	2.07	0.54
1:2A:1422:G:O3'	10:2O:49:ARG:NH1	99.99	0.54
1:2A:236:C:H2'	1:2A:237:C:H6	1.72	0.54
2:2B:101:G:OP2	63:2B:3101:HOH:O	2.17	0.54
1:1A:1814:A:H5'	1:1A:2620:G:H4'	1.89	0.53
1:2A:888:C:H2'	1:2A:889:C:C4	2.43	0.53
1:2A:451:C:H4'	5:2F:52:LYS:HE3	1.89	0.53
15:2T:6:LEU:O	15:2T:10:VAL:HG23	2.08	0.53
2:1B:2:C:H2'	2:1B:3:C:C6	2.42	0.53
8:1I:92:VAL:HG11	8:1I:144:VAL:HG11	1.90	0.53
15:1T:24:PRO:HD3	15:1T:52:ILE:HD12	1.90	0.53
1:2A:1116:C:H2'	1:2A:1117:G:C8	2.43	0.53
1:2A:586:A:N1	1:2A:809:G:O2'	2.35	0.53
1:2A:568:U:H5'	1:2A:945:A:N1	2.22	0.53
29:17:9:ARG:HG2	29:17:46:VAL:HG23	1.90	0.53
1:1A:238:C:O2	30:18:12:LYS:NZ	2.39	0.53
9:1N:62:VAL:HG22	9:1N:66:LYS:HD2	1.89	0.53
1:2A:577:G:O2'	1:2A:1254:A:OP1	2.25	0.53
1:2A:2328:A:H2'	1:2A:2329:G:C8	2.43	0.53
1:2A:615:G:OP1	5:2F:40:GLN:NE2	2.40	0.53
2:2B:83:G:N1	2:2B:94:C:N3	2.41	0.53
13:2R:44:LEU:HD22	13:2R:48:VAL:HG23	1.90	0.53
28:26:12:GLU:OE1	28:26:19:ARG:NH1	2.42	0.53
11:2P:63:PRO:HD3	30:28:27:THR:HG22	1.90	0.53
1:2A:880:G:N1	1:2A:898:C:O2	2.39	0.53
10:2O:7:TYR:CZ	10:2O:44:LYS:HG3	2.44	0.53
1:1A:1425:A:H4'	1:1A:1426:G:OP2	2.09	0.53
1:1A:2340:A:H2'	1:1A:2341:G:C8	2.43	0.53
1:1A:92:C:H2'	1:1A:93:G:C8	3.45	0.53
4:1E:24:THR:HG23	4:1E:184:VAL:HG12	1.91	0.53
1:2A:1446:C:H42	1:2A:1465:G:H1	1.55	0.53
1:1A:2401:G:H5''	1:1A:2402:U:O4'	2.09	0.53
1:1A:964:A:N3	2:1B:80:U:O2'	2.38	0.53

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:2315:G:H2'	1:2A:2316:C:C6	2.44	0.53
1:2A:774:A:N3	1:2A:774:A:H2'	2.24	0.53
23:11:51:VAL:HG11	23:11:74:VAL:HG21	1.90	0.53
1:1A:1140:U:O2'	1:1A:1141:A:N7	2.39	0.53
1:1A:2389:A:H2'	1:1A:2390:A:C8	2.44	0.53
5:1F:34:TRP:HA	11:1P:6:LEU:HD13	1.91	0.53
12:1Q:10:ARG:HH11	12:1Q:11:LYS:HE3	1.74	0.53
21:1Z:155:LEU:HD12	21:1Z:156:LYS:H	1.74	0.53
1:2A:1758:G:O2'	63:2A:3952:HOH:O	2.19	0.53
1:1A:1105:G:H1	1:1A:1125:C:H42	1.57	0.53
1:1A:1239:A:H62	1:1A:1299:A:N6	21.10	0.53
1:1A:2303:U:H2'	1:1A:2304:C:C6	2.44	0.53
1:1A:2760:G:O6	1:1A:2768:C:H5''	2.07	0.53
1:1A:1001:G:H5''	12:1Q:77:LYS:HD2	1.91	0.53
1:2A:1218:C:H42	1:2A:1231:G:H1	1.56	0.53
1:2A:2141:G:H2'	1:2A:2142:C:O4'	2.08	0.53
3:2D:242:ARG:HG2	3:2D:246:PRO:HG3	1.91	0.53
14:2S:61:ASN:O	14:2S:65:VAL:HG23	2.09	0.53
1:1A:2348:A:H61	22:10:43:THR:CG2	2.21	0.53
1:1A:1305:G:N2	1:1A:1331:G:H1'	40.06	0.53
1:1A:1501:U:OP1	13:1R:77:ARG:NH1	2.39	0.53
1:1A:1848:G:OP1	3:1D:88:ARG:NH2	2.41	0.53
1:1A:2013:U:H2'	1:1A:2014:G:H5''	1.91	0.53
1:1A:2227:G:H5''	1:1A:2228:G:C5	2.43	0.53
6:1G:72:ARG:NH1	6:1G:87:PRO:HG3	2.24	0.53
1:2A:1817:G:OP1	3:2D:88:ARG:NH2	2.41	0.53
1:1A:2804:C:H2'	1:1A:2805:G:H8	1.73	0.53
14:1S:39:ILE:HB	14:1S:49:VAL:HG12	1.91	0.53
1:2A:2285:C:OP2	28:26:6:ARG:NH1	2.42	0.53
1:2A:1721:G:H8	1:2A:1741:A:H62	1.56	0.53
1:2A:900:A:H2'	1:2A:901:A:C8	2.44	0.53
5:2F:103:LYS:HA	5:2F:106:ARG:HG3	1.90	0.53
5:1F:32:LEU:HD13	5:1F:112:MET:HE1	1.90	0.52
1:2A:987:G:O2'	1:2A:1000:A:N3	2.41	0.52
1:2A:2114:A:O2'	1:2A:2167:U:H1'	2.08	0.52
1:2A:2287:A:N6	1:2A:2344:U:H3	2.04	0.52
12:1Q:35:VAL:HG13	12:1Q:130:LYS:HB3	1.91	0.52
13:1R:2:ARG:HA	13:1R:5:LYS:HD2	1.91	0.52
1:2A:2292:C:OP1	14:2S:17:ARG:NH2	2.42	0.52
19:2X:4:ALA:HB1	19:2X:42:ALA:HA	1.91	0.52
1:1A:2180:A:O2'	1:1A:2181:G:OP2	2.23	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:2880:C:H2'	1:1A:2881:C:O4'	2.10	0.52
9:1N:20:GLY:HA2	9:1N:61:ARG:HG2	1.91	0.52
21:1Z:1:MET:HE2	21:1Z:55:HIS:HA	1.90	0.52
1:2A:893:C:H2'	1:2A:894:C:C5	2.45	0.52
6:2G:11:TYR:HA	6:2G:15:VAL:HB	1.92	0.52
17:2V:40:LEU:HB2	17:2V:46:VAL:HG13	1.92	0.52
21:2Z:156:LYS:HE3	21:2Z:158:PRO:HD3	1.91	0.52
29:27:9:ARG:HD3	29:27:47:ARG:HB2	1.91	0.52
1:2A:271(H):G:H2'	1:2A:271(I):G:H8	1.73	0.52
2:2B:75:G:H1	21:2Z:73:GLN:NE2	2.08	0.52
6:2G:44:GLY:N	6:2G:88:ILE:O	2.42	0.52
58:A:3:PRO:HG2	58:A:12:SER:HB2	1.91	0.52
7:1H:3:ARG:NH2	7:1H:65:HIS:HB3	2.23	0.52
8:1I:116:LEU:HD11	8:1I:120:ILE:HG13	1.90	0.52
1:2A:1973:G:OP1	63:2A:3955:HOH:O	2.19	0.52
1:2A:2141:G:O6	1:2A:2150:U:O2	2.27	0.52
1:2A:854:G:O6	63:2A:3953:HOH:O	2.19	0.52
4:2E:36:ARG:NH1	4:2E:85:ASN:OD1	2.42	0.52
6:2G:11:TYR:CZ	6:2G:16:ARG:HD3	2.44	0.52
1:1A:1099:C:H42	1:1A:1152:G:H1	1.56	0.52
1:1A:2584:A:C8	4:1E:144:ARG:HD3	2.45	0.52
21:1Z:45:ASP:CG	21:1Z:49:ARG:HE	2.13	0.52
1:2A:2689:U:OP2	1:2A:2719:G:N2	2.40	0.52
3:2D:108:PRO:HB3	3:2D:143:HIS:CE1	2.45	0.52
11:2P:63:PRO:HG2	30:28:25:MET:HB2	1.91	0.52
1:1A:2163:G:C6	1:1A:2164:C:C2	2.98	0.52
1:1A:673:G:H2'	1:1A:674:G:C8	3.06	0.52
1:2A:1170:G:O6	1:2A:1180:C:N4	2.43	0.52
2:2B:24:G:H4'	2:2B:25:A:C8	2.45	0.52
5:2F:53:THR:HG23	5:2F:55:GLY:N	2.24	0.52
1:1A:704:U:H2'	1:1A:705:C:C6	2.45	0.52
1:1A:2317:A:H5''	6:1G:134:GLY:HA3	1.90	0.52
13:1R:97:VAL:HG22	13:1R:114:VAL:HG13	1.92	0.52
21:1Z:53:ILE:HG22	21:1Z:71:VAL:HG12	1.92	0.52
1:1A:174:U:H2'	1:1A:175:G:H8	1.75	0.52
10:1O:2:ILE:HD12	10:1O:6:THR:HG21	1.91	0.52
15:1T:108:ARG:NH2	15:1T:112:ARG:HD3	2.25	0.52
15:1T:74:ARG:HG2	15:1T:76:PHE:CZ	2.45	0.52
1:1A:63:A:O3'	19:1X:71:GLY:HA3	2.10	0.52
1:2A:1794:U:H2'	1:2A:1795:C:C6	2.45	0.52
1:2A:2342:C:O2'	1:2A:2374:C:OP1	2.27	0.52

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:403:U:H4'	1:2A:404:C:H5'	1.90	0.52
19:2X:8:ILE:O	24:22:36:ARG:NH2	2.43	0.52
1:1A:2175:G:H2'	1:1A:2176:G:C8	2.45	0.52
3:1D:180:GLY:HA3	3:1D:275:LYS:HD3	1.91	0.52
1:2A:2064:C:H2'	1:2A:2065:C:C6	2.45	0.52
6:2G:47:LYS:HG3	6:2G:48:GLU:H	1.74	0.52
17:2V:72:VAL:HG13	17:2V:85:LYS:HB3	1.91	0.52
21:2Z:7:ALA:HB3	21:2Z:61:LEU:HD12	1.92	0.52
26:14:55:ARG:N	26:14:56:VAL:HA	2.23	0.51
6:1G:131:TYR:HB3	6:1G:159:VAL:HG13	1.91	0.51
1:2A:2121:G:H1	1:2A:2177:C:H42	1.57	0.51
1:1A:1556:A:H3'	1:1A:1557:A:H8	1.75	0.51
1:1A:2859:U:H4'	1:1A:2878:A:C2	2.46	0.51
3:1D:148:GLU:HB2	3:1D:151:LYS:HD2	1.93	0.51
8:1I:12:LEU:HD22	8:1I:19:VAL:HG21	1.92	0.51
14:1S:43:GLU:OE1	14:1S:43:GLU:N	4.97	0.51
20:1Y:55:TYR:H	20:1Y:56:PRO:CD	2.22	0.51
1:2A:1005:C:H2'	1:2A:1006:C:C6	2.46	0.51
1:2A:2099:U:H3	1:2A:2190:G:H1	1.58	0.51
1:2A:2133:G:O2'	1:2A:2157:G:N2	2.44	0.51
3:2D:145:VAL:HG13	3:2D:191:ALA:HB2	1.90	0.51
4:2E:36:ARG:NH2	4:2E:88:GLY:O	2.43	0.51
9:2N:16:ILE:HD13	9:2N:140:VAL:HG21	1.93	0.51
21:2Z:40:ASP:OD2	21:2Z:42:VAL:HG13	2.10	0.51
1:1A:2339:A:H2'	1:1A:2340:A:C8	2.45	0.51
1:2A:2117:A:O2'	1:2A:2118:U:H5''	2.11	0.51
1:2A:287:C:H2'	1:2A:288:C:H6	1.75	0.51
8:1I:40:THR:O	8:1I:44:LEU:HB2	2.10	0.51
22:20:11:ARG:O	22:20:14:ARG:NH2	2.43	0.51
1:2A:11:G:O5'	1:2A:11:G:H8	1.92	0.51
1:2A:1518:U:H2'	1:2A:1519:G:O4'	2.11	0.51
1:2A:2537:U:H2'	1:2A:2538:C:C6	2.45	0.51
6:2G:109:VAL:HG21	26:24:14:ILE:HD13	1.93	0.51
1:2A:296:C:O3'	20:2Y:95:LYS:NZ	2.44	0.51
1:1A:927:G:N2	1:1A:944:C:N3	2.58	0.51
9:1N:13:TRP:CE2	9:1N:133:GLN:HG2	2.46	0.51
20:1Y:6:HIS:HE1	20:1Y:72:VAL:O	1.93	0.51
4:2E:116:VAL:HG13	4:2E:122:PHE:HB2	1.92	0.51
8:2I:124:GLY:H	8:2I:144:VAL:HG23	1.74	0.51
8:2I:29:TYR:HD2	8:2I:30:LEU:HD23	1.76	0.51
26:14:57:GLU:HB3	26:14:58:ARG:HD2	1.92	0.51

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:1085:G:H1	1:1A:1162:C:N4	2.09	0.51
1:1A:2328:C:O2'	6:1G:128:ARG:NH2	2.26	0.51
4:1E:109:LYS:O	4:1E:111:ARG:NH1	2.43	0.51
5:1F:70:THR:HG23	5:1F:72:ARG:H	1.75	0.51
1:2A:854:G:H2'	1:2A:855:G:C8	2.45	0.51
1:1A:1688:A:H2'	1:1A:1689:G:O4'	2.10	0.51
1:1A:264:G:H1	1:1A:280:C:H42	1.59	0.51
1:2A:1017:G:N7	63:2A:4025:HOH:O	2.34	0.51
1:2A:2137:C:O2'	1:2A:2138:C:OP2	2.28	0.51
1:2A:2646:C:H2'	1:2A:2647:U:O4'	2.11	0.51
1:2A:30:G:H2'	1:2A:31:C:C6	2.45	0.51
1:2A:882:G:H2'	1:2A:883:G:C8	2.45	0.51
1:1A:1037:C:H2'	1:1A:1038:C:C6	2.88	0.51
15:1T:127:ALA:C	15:1T:129:ARG:H	2.13	0.51
1:2A:1796:U:H2'	1:2A:1797:C:H6	1.75	0.51
1:2A:2870:C:H2'	1:2A:2871:C:O4'	2.10	0.51
7:2H:51:ARG:NH1	7:2H:53:GLU:OE2	2.43	0.51
21:2Z:93:ASP:HB2	21:2Z:131:ARG:HH12	1.74	0.51
1:1A:92:C:H2'	1:1A:93:G:H8	3.01	0.51
1:1A:997:G:OP1	12:1Q:16:ARG:NH2	2.44	0.51
3:1D:17:THR:O	3:1D:211:ARG:NH2	2.44	0.51
7:1H:20:ALA:HB1	7:1H:21:PRO:HD2	1.93	0.51
19:1X:53:LYS:HB3	19:1X:82:GLN:HB3	1.92	0.51
6:2G:124:SER:HB2	6:2G:131:TYR:CE1	2.46	0.51
6:2G:39:ILE:HG12	6:2G:157:ILE:HG12	1.93	0.51
7:2H:80:SER:OG	7:2H:81:GLU:OE1	2.28	0.51
1:1A:2168:C:H4'	1:1A:2169:G:C4	2.46	0.51
1:1A:2264:G:OP2	63:1A:4352:HOH:O	2.17	0.51
9:2N:67:LEU:HA	9:2N:87:LEU:HD22	1.93	0.51
1:1A:1552:C:H2'	1:1A:1553:A:H8	1.77	0.50
1:1A:362:G:OP2	63:1A:4355:HOH:O	2.19	0.50
1:1A:1830:G:O2'	3:1D:181:GLU:OE2	2.23	0.50
5:1F:13:SER:HB2	5:1F:127:GLU:OE1	2.11	0.50
19:1X:31:HIS:HD2	19:1X:33:LYS:H	1.59	0.50
1:2A:1364:G:P	23:21:3:LYS:HG3	2.51	0.50
1:2A:2815:C:H5'	27:25:29:THR:HG21	1.92	0.50
30:18:63:PRO:HG2	30:18:64:TYR:CE2	2.46	0.50
1:1A:1218:G:N1	1:1A:1221:G:OP2	2.43	0.50
1:1A:1550:C:H2'	1:1A:1551:C:C6	2.46	0.50
1:1A:265:U:H2'	1:1A:266:C:C6	2.46	0.50
9:1N:4:TYR:HB2	16:1U:101:ARG:NH1	2.27	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:1011:G:H1	1:2A:1018:C:H42	17.91	0.50
1:2A:1181:C:H2'	1:2A:1182:A:C8	2.46	0.50
1:2A:1826:G:H4'	3:2D:242:ARG:CZ	2.41	0.50
1:1A:1110:C:H1'	1:1A:1122:C:H5	1.77	0.50
1:1A:1921:G:H2'	1:1A:1921:G:N3	2.26	0.50
1:1A:2204:G:H2'	1:1A:2205:C:C6	2.46	0.50
1:1A:532:A:N6	1:1A:1206:G:O2'	79.95	0.50
26:24:46:GLN:HE21	26:24:48:ARG:NH2	2.08	0.50
26:24:57:GLU:CB	26:24:58:ARG:HA	2.42	0.50
1:2A:645:C:H5''	1:2A:646:A:OP2	2.11	0.50
1:2A:868:U:C4	1:2A:869:G:N7	2.79	0.50
1:2A:927:G:H2'	1:2A:928:G:O4'	2.11	0.50
3:2D:108:PRO:HB3	3:2D:143:HIS:HE1	1.77	0.50
5:2F:150:GLY:HA2	5:2F:172:TRP:CD2	2.46	0.50
1:2A:2748:A:H5'	7:2H:4:ILE:HD12	1.93	0.50
1:1A:1210:G:H2'	1:1A:1211:U:C6	2.46	0.50
1:1A:1275:G:N7	63:1A:4416:HOH:O	2.34	0.50
1:1A:1374:G:N7	63:1A:4421:HOH:O	2.35	0.50
6:1G:106:LEU:HD12	6:1G:110:ALA:HB3	1.93	0.50
1:2A:2168:G:H8	1:2A:2170:A:N7	2.10	0.50
1:2A:903:C:H2'	1:2A:904:C:C6	2.46	0.50
12:2Q:35:VAL:HG13	12:2Q:130:LYS:HB3	1.93	0.50
11:1P:63:PRO:HG2	30:18:25:MET:HB2	1.93	0.50
1:1A:2495:C:N3	12:1Q:124:LYS:NZ	2.56	0.50
1:1A:1233:U:H4'	17:1V:79:VAL:HG22	1.93	0.50
1:2A:1036:G:OP1	7:2H:59:ARG:HG3	2.12	0.50
1:2A:848:G:H2'	1:2A:849:A:C8	2.46	0.50
1:1A:2576:A:C2	1:1A:2659:U:H4'	2.46	0.50
1:1A:2820:A:N6	1:1A:2900:G:O2'	2.34	0.50
21:1Z:7:ALA:HB3	21:1Z:61:LEU:HD12	1.92	0.50
1:2A:2166:G:H3'	1:2A:2167:U:H5''	1.94	0.50
10:2O:115:VAL:HG13	10:2O:121:VAL:HG21	1.94	0.50
21:2Z:33:LEU:HD21	21:2Z:90:VAL:HG21	1.93	0.50
1:1A:2044:U:O2'	1:1A:2629:C:H5'	2.11	0.50
6:1G:12:TYR:HA	6:1G:16:ARG:HG3	1.92	0.50
9:1N:29:LYS:HD2	9:1N:140:VAL:HB	1.94	0.50
21:1Z:105:VAL:N	21:1Z:139:VAL:O	2.39	0.50
1:2A:1021:A:H62	1:2A:1141:U:H3	1.59	0.50
1:2A:1364:G:OP2	23:21:3:LYS:HG3	2.12	0.50
5:2F:110:LEU:HD11	5:2F:181:LEU:HG	1.93	0.50
1:1A:1817:A:H1'	1:1A:1960:A:N6	2.26	0.50

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:215:G:N2	1:1A:217:A:H62	2.09	0.50
1:1A:909:G:H2'	1:1A:910:A:O4'	2.12	0.50
1:1A:968:U:H2'	1:1A:969:C:C6	2.46	0.50
22:20:10:THR:HG22	22:20:12:ASN:N	2.20	0.50
1:2A:247:G:H4'	1:2A:386:G:C5	2.47	0.50
17:2V:18:LEU:HD12	17:2V:19:LYS:H	1.76	0.50
5:1F:24:LEU:HD23	5:1F:115:ALA:HA	1.94	0.50
1:1A:1001:G:OP2	12:1Q:87:LYS:HE2	2.12	0.50
14:1S:6:ALA:O	14:1S:10:ARG:HG3	2.11	0.50
1:2A:118:A:N3	1:2A:178:G:H1'	2.27	0.50
2:2B:14:U:O3'	2:2B:108:U:O2'	2.27	0.50
2:2B:75:G:N2	21:2Z:87:ASP:OD1	2.44	0.50
1:2A:2818:G:OP2	13:2R:42:LYS:NZ	2.43	0.50
1:1A:1121:C:C2'	1:1A:1122:C:H5'	2.42	0.49
1:1A:1634:C:H2'	1:1A:1635:C:C6	2.46	0.49
1:2A:2125:G:H1'	1:2A:2173:A:N6	2.27	0.49
1:2A:2689:U:P	1:2A:2719:G:H22	2.35	0.49
1:2A:2889:C:H2'	1:2A:2891:G:O4'	2.12	0.49
1:2A:458:G:O2'	1:2A:469:G:O6	2.25	0.49
1:2A:479:A:N3	1:2A:481:G:H5''	2.26	0.49
1:2A:682:G:H1	1:2A:708:C:H42	69.28	0.49
1:2A:731:C:H5''	63:2A:4039:HOH:O	2.12	0.49
9:2N:15:LEU:HB2	9:2N:135:PRO:HB2	1.93	0.49
20:2Y:55:TYR:H	20:2Y:56:PRO:HD3	1.77	0.49
4:1E:24:THR:HG22	4:1E:186:GLY:O	2.11	0.49
14:1S:10:ARG:O	14:1S:14:VAL:HG13	2.12	0.49
1:2A:1579:A:H2'	1:2A:1580:A:C8	2.47	0.49
1:2A:2074:U:H2'	1:2A:2075:U:C6	2.47	0.49
1:2A:2154:G:H2'	1:2A:2155:G:H5'	1.94	0.49
8:2I:5:LEU:HD11	8:2I:19:VAL:HG22	1.93	0.49
1:2A:2882:A:OP1	13:2R:96:ARG:HD3	2.12	0.49
19:2X:53:LYS:HB3	19:2X:82:GLN:HB3	1.94	0.49
30:18:23:VAL:HG11	30:18:47:LYS:HD3	1.94	0.49
1:1A:2346:G:H5'	14:1S:9:ARG:HG2	1.94	0.49
1:1A:469:A:C5	5:1F:45:ARG:HD2	2.47	0.49
9:1N:42:TRP:CH2	9:1N:44:PRO:HB3	2.47	0.49
1:2A:1882:C:H5''	23:21:26:ARG:HH21	1.76	0.49
1:2A:2543:G:H2'	1:2A:2544:G:C8	2.47	0.49
9:2N:30:ILE:HG21	9:2N:120:LEU:HD13	1.95	0.49
1:1A:1735:U:O2	1:1A:1747:A:H5'	2.13	0.49
1:1A:630:U:OP1	5:1F:102:PRO:HA	2.12	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:2156:G:H2'	1:2A:2157:G:C5	2.47	0.49
3:2D:159:ALA:HB1	3:2D:198:ASN:O	2.12	0.49
1:2A:1824:G:N3	3:2D:254:THR:OG1	2.45	0.49
1:1A:174:U:H2'	1:1A:175:G:C8	2.48	0.49
17:1V:14:VAL:HB	17:1V:96:ILE:HG13	1.93	0.49
26:24:33:VAL:HG12	26:24:35:VAL:H	1.77	0.49
29:27:12:ARG:NH2	29:27:44:PRO:HB3	2.26	0.49
1:2A:832:G:OP1	63:2A:3951:HOH:O	2.18	0.49
7:2H:86:GLU:OE2	7:2H:132:ARG:NH2	2.45	0.49
21:2Z:155:LEU:HB3	21:2Z:157:LEU:HG	1.94	0.49
24:12:65:ASN:OD1	24:12:69:ARG:NH1	2.45	0.49
1:1A:55:A:N7	1:1A:357:G:N2	59.51	0.49
1:1A:599:U:H2'	1:1A:600:G:C8	2.47	0.49
1:2A:2156:G:H2'	1:2A:2157:G:C4	2.48	0.49
1:2A:2819:G:N7	63:2A:4032:HOH:O	2.35	0.49
1:1A:2155:G:H21	1:1A:2156:A:H62	1.61	0.49
1:1A:2699:U:H2'	1:1A:2700:U:O4'	2.13	0.49
1:1A:709:G:H5''	11:1P:16:ARG:HG2	1.93	0.49
1:1A:2800:C:H1'	4:1E:62:PRO:HG3	1.93	0.49
5:1F:70:THR:CG2	5:1F:72:ARG:H	2.24	0.49
21:1Z:136:PHE:O	21:1Z:137:ILE:HG13	2.13	0.49
1:2A:857:C:H1'	22:20:26:TYR:CE1	2.48	0.49
1:2A:2165:G:H2'	1:2A:2166:G:O4'	2.13	0.49
1:2A:2815:C:H2'	1:2A:2816:C:H6	1.77	0.49
1:2A:2572:A:N7	4:2E:145:LYS:HB2	2.28	0.49
1:1A:561:A:H2'	1:1A:562:C:C6	2.47	0.49
7:1H:24:VAL:HG22	7:1H:35:VAL:HB	1.95	0.49
8:1I:31:LEU:HD21	8:1I:38:LEU:HG	1.94	0.49
11:1P:106:LEU:HD22	11:1P:112:LEU:HG	1.94	0.49
15:1T:51:ARG:HG3	15:1T:98:LYS:HD2	1.95	0.49
1:2A:1589:C:H2'	1:2A:1590:U:C6	2.48	0.49
1:2A:1794:U:H2'	1:2A:1795:C:H6	1.78	0.49
1:2A:2025:C:H2'	1:2A:2026:C:C6	2.48	0.49
1:2A:2298:A:N6	1:2A:2318:G:C8	2.80	0.49
1:2A:2849:U:H4'	1:2A:2868:A:C2	2.48	0.49
13:2R:2:ARG:NH1	13:2R:5:LYS:O	2.45	0.49
14:2S:15:ARG:HB3	14:2S:19:LYS:NZ	2.28	0.49
1:1A:1370:G:N7	63:1A:4417:HOH:O	2.34	0.49
1:1A:2143:G:H1	1:1A:2199:C:N4	2.10	0.49
4:1E:116:VAL:HG13	4:1E:122:PHE:HB2	1.93	0.49
1:2A:1360:A:OP1	1:2A:1360:A:H8	4.98	0.49

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:1150:C:H2'	1:1A:1151:U:C6	2.48	0.49
1:1A:1834:A:O2'	3:1D:259:THR:HG21	2.13	0.49
1:2A:27:G:N2	1:2A:512:G:H1'	2.28	0.49
1:2A:500:G:N1	1:2A:503:A:OP2	2.46	0.49
4:2E:14:ILE:HG13	4:2E:21:VAL:HG13	1.94	0.49
21:2Z:130:PRO:HA	21:2Z:133:ILE:HG13	1.95	0.49
1:1A:1177:G:H21	9:1N:73:THR:CG2	2.26	0.48
1:1A:2579:G:H2'	1:1A:2580:C:C6	2.48	0.48
6:1G:122:PRO:HB3	6:1G:170:ARG:HH12	1.77	0.48
8:1I:77:LEU:HD21	8:1I:79:ILE:HD11	1.95	0.48
1:2A:1359:A:H2'	1:2A:1360:A:H5'	1.95	0.48
1:2A:1405:U:H2'	1:2A:1406:U:C6	2.48	0.48
1:2A:2250:G:OP1	12:2Q:85:LYS:NZ	2.41	0.48
1:2A:2273:A:H2'	1:2A:2274:A:C8	2.48	0.48
1:2A:646:A:H2'	1:2A:647:G:O4'	2.13	0.48
6:2G:123:ASN:C	6:2G:125:PHE:H	2.17	0.48
10:2O:86:ILE:HG22	10:2O:94:ARG:HD3	1.94	0.48
1:1A:2193:A:O2'	1:1A:2194:U:H5''	2.13	0.48
1:1A:2549:U:H2'	1:1A:2550:C:C6	2.48	0.48
1:1A:308:U:H2'	1:1A:309:C:C6	2.48	0.48
12:1Q:135:ASP:N	12:1Q:138:ASP:OD2	2.43	0.48
15:1T:36:GLU:OE1	15:1T:41:ARG:HD3	2.13	0.48
1:2A:111:A:O2'	24:22:65:ASN:ND2	2.43	0.48
1:2A:912:C:OP1	12:2Q:8:LYS:NZ	2.30	0.48
1:2A:1803:A:O2'	3:2D:259:THR:HG21	2.13	0.48
18:2W:4:LYS:HE3	18:2W:6:ILE:HD11	1.94	0.48
26:14:58:ARG:O	26:14:61:ARG:HB3	2.12	0.48
30:18:23:VAL:CG1	30:18:47:LYS:HD3	2.43	0.48
16:1U:85:LYS:HD3	16:1U:116:ALA:O	2.13	0.48
25:23:23:LEU:HD13	25:23:50:VAL:HG11	1.96	0.48
1:2A:1213:A:N3	1:2A:1238:G:O2'	2.42	0.48
1:2A:300:A:H2'	1:2A:334:C:H1'	1.96	0.48
1:2A:597:U:H2'	1:2A:598:G:C8	2.49	0.48
1:2A:811:U:H2'	11:2P:21:ARG:HA	1.95	0.48
1:1A:2444:A:C4	23:11:33:LYS:HG2	2.48	0.48
26:14:63:TYR:CD1	26:14:63:TYR:N	2.81	0.48
1:1A:2705:A:H2'	1:1A:2706:G:H8	1.78	0.48
8:1I:27:ARG:HG2	23:11:71:TYR:CZ	2.47	0.48
25:23:46:ASN:O	25:23:50:VAL:HG22	2.13	0.48
26:24:57:GLU:CB	26:24:58:ARG:HD2	2.43	0.48
1:2A:1791:A:H5'	3:2D:206:LEU:HD12	1.94	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:196:A:O2'	1:2A:805:G:O6	2.29	0.48
21:2Z:93:ASP:CB	21:2Z:131:ARG:HH22	2.27	0.48
6:1G:179:PRO:HG3	26:14:43:TYR:OH	2.13	0.48
1:1A:605:G:H2'	1:1A:606:G:C8	2.48	0.48
1:1A:883:G:N7	63:1A:4426:HOH:O	2.35	0.48
2:2B:4:C:H42	2:2B:117:G:H1	1.60	0.48
8:2I:4:ILE:HG12	8:2I:18:VAL:HG22	1.95	0.48
9:2N:42:TRP:HA	9:2N:48:MET:SD	2.54	0.48
12:2Q:30:GLY:HA2	12:2Q:107:ALA:HB2	1.95	0.48
1:1A:731:G:OP1	29:17:16:HIS:ND1	2.41	0.48
1:1A:1993:A:OP2	3:1D:242:ARG:NH2	2.47	0.48
8:1I:81:VAL:HG21	8:1I:88:ILE:HD13	1.94	0.48
1:2A:851:U:H5'	25:23:49:LYS:HD2	1.95	0.48
1:2A:1469:A:H2'	1:2A:1470:G:O4'	2.14	0.48
1:2A:910:A:N1	1:2A:2277:G:H1'	2.27	0.48
19:2X:31:HIS:CD2	19:2X:33:LYS:H	2.32	0.48
1:1A:1121:C:H2'	1:1A:1122:C:H5'	1.95	0.48
1:1A:2227:G:OP2	1:1A:2227:G:H4'	2.14	0.48
1:1A:2603:C:H2'	1:1A:2604:G:C8	2.48	0.48
1:1A:1201:A:OP1	16:1U:55:ARG:HD3	2.14	0.48
1:1A:2288:G:N7	63:1A:4423:HOH:O	2.35	0.48
1:1A:821:A:H2'	1:1A:821:A:N3	2.28	0.48
1:1A:2225:U:O4'	3:1D:151:LYS:HE2	2.13	0.48
3:1D:77:ALA:HB2	3:1D:97:TYR:CD1	2.48	0.48
1:2A:1354:A:H5''	3:2D:38:LYS:HD3	1.96	0.48
1:2A:2065:C:H2'	1:2A:2066:C:C6	2.48	0.48
1:2A:2366:A:H2'	1:2A:2367:G:O4'	2.13	0.48
1:2A:2674:G:H2'	1:2A:2675:A:C8	2.49	0.48
1:2A:277:C:H4'	1:2A:278:A:C8	2.49	0.48
1:2A:71:A:N7	19:2X:31:HIS:HE1	2.11	0.48
2:2B:66:A:H61	2:2B:109:C:H5''	1.78	0.48
1:2A:1843:C:H5'	3:2D:253:GLN:OE1	2.14	0.48
4:2E:111:ARG:HD3	4:2E:160:TYR:CD2	2.48	0.48
4:2E:9:VAL:HG13	4:2E:25:VAL:O	2.14	0.48
5:2F:31:HIS:NE2	5:2F:35:GLU:OE2	2.47	0.48
8:2I:140:LEU:HD22	8:2I:142:VAL:HG22	1.96	0.48
20:2Y:9:LYS:HA	20:2Y:10:GLY:HA2	1.57	0.48
20:2Y:38:ILE:HD11	20:2Y:66:PRO:HG3	1.96	0.48
1:1A:1121:C:H42	1:1A:1123:A:H62	1.62	0.48
1:2A:1753:G:H5''	15:2T:95:ARG:HD2	1.96	0.48
1:2A:1853:A:N3	1:2A:2233:U:O2'	2.45	0.48

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:2206:G:H5''	1:2A:2207:G:C8	2.48	0.48
1:2A:2224:G:OP1	3:2D:268:ARG:NE	2.47	0.48
1:2A:588:U:H2'	1:2A:589:C:C6	2.49	0.48
11:2P:82:GLY:HA2	11:2P:113:LYS:O	2.14	0.48
17:2V:1:MET:CE	17:2V:44:LYS:H	2.27	0.48
1:1A:1627:A:H8	1:1A:1627:A:OP2	1.97	0.48
1:1A:925:A:N6	1:1A:945:A:O2'	2.47	0.48
15:1T:33:LYS:O	15:1T:82:LEU:HD23	2.14	0.48
1:2A:1138:G:C6	1:2A:1140:C:H1'	6.31	0.48
1:2A:1286:A:H8	1:2A:1287:A:H4'	8.04	0.48
1:2A:1429:G:H2'	1:2A:1430:C:C6	2.48	0.48
1:2A:2144:U:H1'	1:2A:2148:G:H22	1.79	0.48
1:2A:2315:G:H2'	1:2A:2316:C:H6	1.79	0.48
1:2A:2609:U:H2'	58:B:5:ASN:HD21	1.79	0.48
1:2A:800:A:OP1	1:2A:800:A:H8	1.97	0.48
2:2B:3:C:H2'	2:2B:4:C:C6	2.48	0.48
4:2E:143:ASN:HD22	4:2E:147:PRO:HD3	1.79	0.48
12:2Q:134:ARG:CZ	21:2Z:122:ARG:HH21	2.27	0.48
26:14:49:PHE:HB3	26:14:50:VAL:H	1.52	0.47
1:1A:2178:G:H2'	1:1A:2179:G:C2	2.48	0.47
1:1A:2705:A:H2'	1:1A:2706:G:C8	2.49	0.47
1:1A:312:C:H2'	1:1A:313:A:H8	1.77	0.47
4:1E:11:MET:HG2	4:1E:24:THR:HB	1.96	0.47
1:2A:131:G:OP1	63:2A:3954:HOH:O	2.19	0.47
5:2F:187:VAL:HG12	11:2P:3:LEU:HD12	1.96	0.47
1:1A:2133:C:N4	1:1A:2166:U:O2'	2.46	0.47
1:1A:2175:G:H2'	1:1A:2176:G:H8	1.78	0.47
1:1A:2190:G:N1	1:1A:2193:A:C8	2.81	0.47
1:1A:236:G:H4'	1:1A:413:G:C5	2.49	0.47
28:26:23:THR:OG1	28:26:24:GLU:N	2.46	0.47
1:2A:1147:C:H2'	1:2A:1148:A:H8	1.79	0.47
1:2A:2609:U:H2'	58:B:5:ASN:ND2	2.29	0.47
1:2A:2657:A:O3'	7:2H:160:LYS:NZ	2.38	0.47
1:2A:287:C:H2'	1:2A:288:C:C6	2.49	0.47
1:2A:947:G:H2'	1:2A:948:G:H8	1.77	0.47
13:2R:92:GLY:HA2	13:2R:94:TYR:CZ	2.49	0.47
1:2A:1754:C:OP1	15:2T:96:ARG:NH1	2.47	0.47
28:16:14:THR:HG21	28:16:48:VAL:HG13	1.96	0.47
23:21:52:ARG:HH21	23:21:57:GLU:HB2	1.78	0.47
25:23:18:ASP:OD1	25:23:18:ASP:N	2.40	0.47
1:2A:1019:U:OP1	1:2A:1035:U:O2'	2.21	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:1477:A:H2'	1:2A:1478:G:O4'	2.14	0.47
1:2A:829:A:N7	1:2A:2248:C:H5'	2.29	0.47
1:1A:1312:G:O5'	18:1W:15:ARG:NH2	2.48	0.47
1:1A:928:G:C2	1:1A:929:G:H1'	2.50	0.47
4:1E:1:MET:HE3	4:1E:199:ARG:HD2	1.96	0.47
23:21:4:VAL:HG11	23:21:11:ARG:NH2	2.28	0.47
1:2A:821:A:N1	63:2A:4036:HOH:O	2.35	0.47
8:2I:123:LEU:HD21	8:2I:145:VAL:HA	1.96	0.47
1:1A:606:G:N2	1:1A:632:A:N7	49.66	0.47
13:1R:28:LEU:HD22	13:1R:44:LEU:HD13	1.94	0.47
1:2A:212:G:H2'	1:2A:213:A:O4'	2.14	0.47
1:2A:2180:U:H2'	1:2A:2181:G:O4'	2.13	0.47
1:2A:2732:G:H3'	1:2A:2733:A:O4'	2.15	0.47
1:2A:1695:G:N7	3:2D:14:ARG:NH2	2.62	0.47
6:2G:120:LEU:HD12	6:2G:178:PHE:HB3	1.95	0.47
1:1A:1470:G:H2'	1:1A:1471:G:O4'	2.14	0.47
1:1A:347:G:C8	5:1F:171:PRO:HG3	2.49	0.47
1:1A:660:C:O2'	1:1A:664:U:OP1	2.25	0.47
1:1A:768:C:H2'	1:1A:769:A:C8	2.50	0.47
11:1P:138:LEU:HD23	11:1P:145:PRO:HB3	1.96	0.47
13:1R:38:VAL:HG22	13:1R:112:ALA:HB2	1.96	0.47
17:1V:49:THR:HG22	17:1V:49:THR:O	2.15	0.47
1:2A:1006:C:H2'	1:2A:1007:C:C6	3.15	0.47
1:2A:2438:U:O2'	1:2A:2440:C:OP1	2.26	0.47
1:2A:288:C:H2'	1:2A:289:A:H8	1.80	0.47
5:2F:11:VAL:HG22	5:2F:125:LEU:HB2	1.97	0.47
21:2Z:145:GLU:C	21:2Z:147:GLY:H	2.16	0.47
1:1A:1004:A:C5'	1:1A:1024:G:H1	27.68	0.47
1:1A:1074:A:N6	1:1A:1171:G:H2'	2.30	0.47
5:1F:28:ILE:O	5:1F:30:PRO:HD3	2.15	0.47
7:1H:154:PRO:HB3	7:1H:163:TYR:CZ	2.48	0.47
11:1P:101:VAL:HG21	11:1P:108:LYS:HG2	1.96	0.47
27:25:40:LYS:NZ	27:25:44:THR:O	2.44	0.47
1:2A:1406:U:H2'	1:2A:1407:C:C6	2.49	0.47
1:2A:2473:U:H2'	1:2A:2473:U:O2	2.14	0.47
5:2F:9:ILE:HG21	5:2F:125:LEU:HD13	1.97	0.47
7:2H:3:ARG:HH11	7:2H:4:ILE:H	1.61	0.47
12:2Q:26:TYR:O	12:2Q:67:ARG:NH1	2.40	0.47
1:2A:1754:C:H5	15:2T:96:ARG:NH2	2.12	0.47
18:2W:46:PHE:O	18:2W:50:VAL:HG23	2.15	0.47
28:16:38:LYS:HE3	28:16:38:LYS:HB3	1.51	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
7:1H:97:ARG:NE	7:1H:104:GLU:OE1	2.47	0.47
1:2A:1833:U:OP1	63:2A:3956:HOH:O	2.20	0.47
1:2A:265:A:C8	1:2A:266:G:H1'	2.49	0.47
1:2A:323:G:O2'	1:2A:1205:U:N3	2.33	0.47
2:2B:105:A:H5'	2:2B:106:G:OP2	2.15	0.47
4:2E:111:ARG:HD3	4:2E:160:TYR:CE2	2.49	0.47
4:2E:9:VAL:HG22	4:2E:25:VAL:HB	1.95	0.47
21:2Z:73:GLN:HB3	21:2Z:87:ASP:HB2	1.97	0.47
1:1A:1136:U:C2	1:1A:1148:C:H1'	2.50	0.47
7:1H:84:SER:OG	7:1H:132:ARG:NH1	2.48	0.47
31:29:7:VAL:HG12	31:29:34:GLN:HB3	1.97	0.47
1:2A:336:C:H2'	1:2A:337:C:C6	2.92	0.47
1:2A:643:A:N1	1:2A:2369:A:O2'	2.38	0.47
24:12:32:LEU:HD22	24:12:36:ARG:NH1	2.30	0.47
25:13:18:ASP:OD1	25:13:18:ASP:N	2.48	0.47
26:14:48:ARG:HD3	26:14:48:ARG:HA	1.66	0.47
1:1A:2157:A:H5'	1:1A:2182:G:H4'	1.97	0.47
1:1A:956:A:N1	1:1A:2289:G:H1'	2.29	0.47
1:1A:2318:C:O2	63:1G:5001:HOH:O	2.15	0.47
1:1A:886:U:H2'	1:1A:887:C:C6	2.50	0.47
6:1G:15:VAL:HG22	6:1G:175:LEU:HB3	1.96	0.47
11:1P:90:ARG:NH1	11:1P:105:LEU:HD11	2.30	0.47
1:2A:328:U:H4'	20:2Y:68:HIS:CD2	2.50	0.47
4:2E:5:LEU:HD22	4:2E:197:ILE:HG12	1.96	0.47
6:2G:123:ASN:O	6:2G:125:PHE:N	2.48	0.47
8:2I:130:TYR:HB3	8:2I:138:ILE:HB	1.97	0.47
10:2O:120:GLU:HG2	10:2O:122:LEU:HG	1.97	0.47
9:2N:4:TYR:CD2	16:2U:100:VAL:HG11	2.50	0.47
21:2Z:73:GLN:O	21:2Z:87:ASP:N	2.41	0.47
1:1A:1091:A:OP1	1:1A:1092:A:H3'	2.15	0.47
1:1A:2053:A:C6	1:1A:2510:C:H1'	2.50	0.47
5:1F:133:ASN:N	5:1F:138:GLU:OE1	2.31	0.47
12:1Q:60:ARG:HH11	12:1Q:60:ARG:HB2	1.78	0.47
1:2A:1005:C:O2'	9:2N:28:THR:HG21	2.15	0.47
1:2A:1025:G:C4	1:2A:1135:C:H1'	2.49	0.47
1:2A:1932:A:H2'	1:2A:1933:G:O4'	2.14	0.47
1:2A:2112:G:H2'	1:2A:2113:U:O4'	2.15	0.47
1:2A:2184:G:H2'	1:2A:2185:C:C6	2.50	0.47
1:2A:652(B):A:N6	1:2A:655:A:H1'	2.30	0.47
3:2D:73:VAL:HG13	3:2D:120:GLY:HA3	1.97	0.47
7:2H:18:GLU:HG3	7:2H:25:LYS:HB3	1.96	0.47

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:1410:G:P	23:11:3:LYS:HG3	2.55	0.46
1:1A:1451:U:H2'	1:1A:1452:U:C6	2.51	0.46
1:1A:1639:G:H2'	1:1A:1640:G:C8	2.50	0.46
1:1A:2193:A:HO2'	1:1A:2194:U:H6	1.61	0.46
1:1A:646:A:OP2	11:1P:108:LYS:NZ	2.48	0.46
1:2A:1639:U:H2'	1:2A:1640:C:H5''	1.96	0.46
1:2A:2134:A:H2'	1:2A:2134:A:N3	2.29	0.46
1:2A:434:U:H2'	1:2A:435:C:C6	6.27	0.46
1:1A:1132:A:H5'	1:1A:1133:G:OP1	2.15	0.46
18:1W:4:LYS:HD2	18:1W:6:ILE:HD11	1.97	0.46
1:2A:1169:G:N2	1:2A:1181:C:N3	2.62	0.46
8:2I:122:GLU:O	8:2I:126:TYR:OH	2.31	0.46
1:2A:445:C:OP1	16:2U:2:PRO:HA	2.14	0.46
21:2Z:97:GLU:HA	21:2Z:126:VAL:O	2.15	0.46
30:18:62:LEU:HB3	30:18:65:GLU:HG2	1.95	0.46
1:1A:1314:A:C2	1:1A:2035:A:C4	3.04	0.46
1:1A:1473:A:H4'	1:1A:1474:C:O4'	2.16	0.46
1:1A:1495:G:H4'	1:1A:1589:A:OP1	2.16	0.46
1:1A:2091:G:OP2	63:1A:4358:HOH:O	2.21	0.46
9:1N:108:PRO:O	9:1N:113:GLY:HA3	2.15	0.46
1:1A:360:C:OP1	20:1Y:84:ARG:HG2	2.15	0.46
1:2A:1593:G:H2'	1:2A:1594:G:C8	2.51	0.46
14:2S:14:VAL:HG21	14:2S:90:GLY:O	2.15	0.46
25:13:7:LYS:HE3	25:13:32:GLN:NE2	2.31	0.46
1:1A:1101:G:H1	1:1A:1150:C:N4	2.10	0.46
1:1A:1405:A:C2	1:1A:1418:U:O4	2.65	0.46
1:1A:555:G:N1	1:1A:2045:G:OP1	2.33	0.46
1:1A:2130:C:H2'	1:1A:2131:U:H6	1.81	0.46
1:1A:2298:A:H4'	1:1A:2299:A:O4'	2.15	0.46
1:1A:605:G:OP2	63:1A:4357:HOH:O	2.20	0.46
1:2A:582:G:H2'	1:2A:583:G:C8	2.51	0.46
5:2F:178:PRO:HB2	5:2F:201:VAL:HG21	1.96	0.46
14:2S:3:ARG:HD3	14:2S:4:LEU:H	1.80	0.46
1:1A:2623:U:H5'	1:1A:2623:U:H6	1.80	0.46
1:2A:242:G:C8	30:28:5:LYS:HG2	2.50	0.46
1:2A:2558:C:H2'	1:2A:2559:C:O4'	2.16	0.46
1:2A:274:G:H2'	1:2A:275:G:C8	2.50	0.46
1:2A:302:C:OP2	20:2Y:73:ARG:NH1	2.47	0.46
30:18:8:LYS:O	30:18:12:LYS:HG3	2.16	0.46
1:1A:2143:G:H2'	1:1A:2144:U:C6	2.51	0.46
3:1D:127:VAL:HA	3:1D:193:VAL:HG22	1.97	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:1F:74:ARG:H	5:1F:74:ARG:HG3	1.35	0.46
9:1N:75:TYR:CE2	9:1N:77:GLY:HA2	2.51	0.46
1:2A:1667:G:O2'	1:2A:1991:U:O4	2.27	0.46
1:2A:2010:G:H5''	18:2W:42:ARG:HB2	1.98	0.46
1:2A:2115:G:H4'	1:2A:2167:U:C4	2.51	0.46
1:2A:2320:A:H2'	1:2A:2320:A:N3	2.31	0.46
1:2A:2376:A:N3	14:2S:106:ARG:NH2	2.58	0.46
1:2A:274:G:H2'	1:2A:275:G:H8	1.80	0.46
2:2B:1:U:H2'	2:2B:2:C:C6	2.50	0.46
6:2G:131:TYR:HB3	6:2G:159:VAL:HG13	1.96	0.46
15:2T:109:GLU:HG2	15:2T:112:ARG:NH2	2.31	0.46
1:1A:2142:G:C2'	1:1A:2143:G:H5'	2.46	0.46
1:1A:2178:G:H8	1:1A:2178:G:OP2	1.98	0.46
1:1A:927:G:H1	1:1A:944:C:N4	2.12	0.46
1:1A:2764:G:C4	7:1H:2:SER:HA	2.51	0.46
12:1Q:134:ARG:NH2	21:1Z:122:ARG:HE	2.14	0.46
1:2A:1239:G:H2'	1:2A:1240:U:O4'	2.15	0.46
1:2A:581:C:H2'	1:2A:582:G:C8	2.51	0.46
4:2E:101:ARG:CZ	4:2E:171:GLU:HB2	2.46	0.46
1:1A:1221:G:N2	1:1A:1223:C:OP2	2.48	0.46
1:1A:1554:A:O2'	1:1A:1555:C:H5''	2.15	0.46
4:1E:12:THR:HG22	15:1T:58:ASN:OD1	2.15	0.46
6:1G:114:ILE:HB	6:1G:117:PHE:HB2	1.98	0.46
8:1I:124:GLY:H	8:1I:144:VAL:HG23	1.81	0.46
16:1U:46:ALA:O	16:1U:50:ARG:HG2	2.16	0.46
20:1Y:9:LYS:HA	20:1Y:10:GLY:HA2	1.66	0.46
1:2A:2129:C:H5'	1:2A:2130:U:OP2	2.16	0.46
1:2A:35:G:H2'	1:2A:36:G:O4'	2.16	0.46
1:2A:492:A:H2'	1:2A:493:G:O4'	2.16	0.46
14:2S:27:SER:HA	14:2S:88:ASP:HB3	1.97	0.46
21:2Z:93:ASP:HB3	21:2Z:131:ARG:HH22	1.81	0.46
1:1A:1324:A:OP1	13:1R:36:THR:HG23	2.16	0.46
1:1A:1476:C:H2'	1:1A:1477:U:C6	2.50	0.46
1:1A:2658:C:H2'	1:1A:2659:U:O4'	2.15	0.46
1:1A:2863:C:H2'	1:1A:2864:G:C8	2.51	0.46
1:1A:78:G:O2'	1:1A:79:G:H5'	2.63	0.46
5:1F:64:ILE:HG21	5:1F:78:ILE:HG23	1.98	0.46
11:1P:97:PRO:HD3	11:1P:126:VAL:O	2.15	0.46
16:1U:104:GLN:NE2	16:1U:105:VAL:HG23	2.31	0.46
25:23:7:LYS:HE3	25:23:32:GLN:NE2	2.31	0.46
1:2A:2059:A:O2'	5:2F:69:HIS:HD2	1.98	0.46

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:2079:U:O3'	23:21:35:THR:OG1	2.33	0.46
1:2A:639:U:H2'	1:2A:640:C:C6	2.51	0.46
2:2B:55:U:H1'	6:2G:29:TRP:CD1	2.51	0.46
7:2H:4:ILE:O	7:2H:69:ARG:HD2	2.15	0.46
9:2N:99:LEU:HD22	9:2N:103:VAL:HG23	1.98	0.46
9:2N:30:ILE:HG23	9:2N:52:VAL:HG11	1.96	0.46
15:2T:18:ASP:OD1	15:2T:18:ASP:N	2.36	0.46
21:2Z:75:ASN:O	21:2Z:84:GLU:HG3	2.16	0.46
25:13:8:LEU:HG	25:13:31:LEU:HD23	1.98	0.46
26:14:33:VAL:HG12	26:14:35:VAL:H	1.81	0.46
27:15:48:GLU:O	27:15:60:VAL:HG11	2.15	0.46
1:1A:1051:C:H5''	63:1A:4773:HOH:O	2.16	0.46
1:1A:1469:G:H2'	1:1A:1470:G:H8	2.08	0.46
1:1A:2418:U:H2'	1:1A:2418:U:OP2	2.17	0.46
1:1A:602:G:H2'	1:1A:603:C:C6	2.51	0.46
1:2A:1274:A:N3	1:2A:1297:C:H1'	2.31	0.46
1:2A:2472:G:H2'	1:2A:2475:C:H42	1.81	0.46
1:2A:2746:U:OP1	7:2H:85:LYS:NZ	2.45	0.46
1:2A:839:U:H2'	1:2A:840:C:C6	2.51	0.46
4:2E:2:LYS:HB2	4:2E:95:ILE:HD12	1.98	0.46
1:1A:1091:A:O2'	1:1A:1093:G:C4	2.68	0.45
1:1A:1103:A:N6	1:1A:1133:G:OP2	2.49	0.45
1:1A:1845:G:H4'	3:1D:51:VAL:HG21	1.98	0.45
1:1A:2285:A:H2'	1:1A:2286:A:C8	2.51	0.45
1:1A:2291:G:O6	22:10:14:ARG:HG3	2.15	0.45
1:1A:2638:C:H2'	1:1A:2639:G:O4'	2.15	0.45
1:1A:44:G:H5''	1:1A:45:C:OP1	2.15	0.45
1:2A:1688:U:O2	1:2A:1700:A:H5'	2.15	0.45
1:2A:271(X):G:C2	1:2A:271(Y):U:O4	2.68	0.45
1:2A:2836:U:H2'	1:2A:2837:G:C8	2.51	0.45
8:2I:93:THR:N	8:2I:96:ASP:HB2	2.31	0.45
10:2O:15:GLY:O	10:2O:47:ILE:HG12	2.17	0.45
13:2R:63:ARG:O	13:2R:67:LEU:HB2	2.16	0.45
13:2R:72:ASP:O	13:2R:76:VAL:HG23	2.16	0.45
1:1A:2660:C:H2'	1:1A:2661:U:C6	2.52	0.45
1:1A:2801:C:P	4:1E:61:ARG:HH21	2.38	0.45
9:1N:4:TYR:CD2	16:1U:100:VAL:HG11	2.51	0.45
1:2A:1410:G:H2'	1:2A:1411:C:C6	2.51	0.45
1:2A:686:G:N2	1:2A:788:A:H61	2.13	0.45
23:11:56:GLN:HE21	23:11:87:PRO:HG3	1.80	0.45
25:13:29:ARG:HD2	63:13:3104:HOH:O	2.17	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:1081:U:H2'	1:1A:1082:G:C8	2.51	0.45
1:1A:1766:G:H1'	1:1A:1770:A:H61	1.81	0.45
1:1A:310:C:H2'	1:1A:311:C:H6	1.81	0.45
1:1A:927:G:H2'	1:1A:928:G:C8	2.41	0.45
3:1D:242:ARG:HD3	3:1D:242:ARG:N	2.31	0.45
4:1E:3:GLY:HA3	4:1E:81:ILE:HD12	1.98	0.45
1:2A:1042:G:C5	1:2A:1043:C:H5	2.35	0.45
1:2A:1636:C:H2'	1:2A:1637:A:C8	2.51	0.45
1:2A:1837:C:O2'	1:2A:1927:A:N3	2.42	0.45
1:2A:2162:G:H4'	1:2A:2172:U:H2'	1.99	0.45
1:2A:2815:C:H2'	1:2A:2816:C:C6	2.51	0.45
1:2A:902:C:H2'	1:2A:903:C:C6	2.51	0.45
8:2I:50:ARG:NH2	63:2I:5001:HOH:O	2.45	0.45
1:2A:2467:C:H4'	12:2Q:123:HIS:CD2	2.51	0.45
24:12:1:MET:HE2	24:12:1:MET:HB3	1.82	0.45
1:1A:2255:U:H2'	1:1A:2256:U:C6	2.51	0.45
5:1F:95:ARG:HD3	5:1F:97:TYR:CZ	2.52	0.45
23:21:81:LYS:HE2	23:21:81:LYS:HB3	1.78	0.45
1:2A:2067:G:O2'	1:2A:2069:G:H5'	2.15	0.45
1:2A:2184:G:O2'	1:2A:2185:C:H5'	2.16	0.45
1:2A:2758:A:C2	1:2A:2759:G:H1'	2.52	0.45
1:2A:1805:U:O2	3:2D:50:THR:HB	2.17	0.45
2:2B:33:G:H5'	6:2G:2:PRO:HD3	1.97	0.45
14:2S:77:ALA:HB1	14:2S:82:ILE:HB	1.98	0.45
16:2U:49:HIS:HA	16:2U:52:ARG:HB2	1.97	0.45
28:16:18:ARG:HD2	28:16:42:TRP:CD1	2.51	0.45
1:1A:1140:U:H2'	1:1A:1141:A:C8	2.51	0.45
1:1A:1821:C:H2'	1:1A:1822:A:C5	2.51	0.45
1:1A:2218:C:O2'	1:1A:2219:U:H5'	2.15	0.45
20:1Y:20:TYR:CE2	20:1Y:43:ASN:HA	2.50	0.45
19:2X:5:TYR:CE2	24:22:30:ARG:HB2	2.51	0.45
1:2A:621:A:OP2	11:2P:108:LYS:NZ	2.45	0.45
2:2B:94:C:H2'	2:2B:95:C:C6	2.52	0.45
3:2D:274:ARG:O	3:2D:275:LYS:HD2	2.16	0.45
6:2G:150:ASP:OD1	6:2G:150:ASP:N	2.48	0.45
8:2I:116:LEU:HD11	8:2I:120:ILE:HG13	1.99	0.45
20:2Y:19:LYS:HE2	20:2Y:20:TYR:CE2	2.52	0.45
1:1A:1403:U:H2'	1:1A:1404:G:O4'	2.16	0.45
1:1A:1686:U:O2'	1:1A:1687:C:H5'	2.17	0.45
1:1A:704:U:H2'	1:1A:705:C:H6	1.82	0.45
6:1G:82:LEU:HD21	6:1G:88:ILE:HG21	1.98	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
21:1Z:8:TYR:HB2	21:1Z:38:TYR:CE2	2.51	0.45
1:2A:1240:U:OP2	6:2G:115:ARG:HA	150.25	0.45
1:2A:2611:U:C4	27:25:3:LYS:HG2	2.52	0.45
1:2A:2747:G:O6	1:2A:2755:C:H5''	2.17	0.45
1:2A:920:G:H2'	1:2A:921:G:H8	1.82	0.45
2:2B:13:A:O2'	2:2B:14:U:H3'	2.17	0.45
3:2D:242:ARG:N	3:2D:242:ARG:HD3	2.32	0.45
7:2H:149:ARG:HG3	7:2H:162:ILE:O	2.16	0.45
9:2N:58:ASP:N	9:2N:58:ASP:OD1	2.47	0.45
16:2U:16:LYS:HB3	16:2U:16:LYS:HE2	1.64	0.45
23:11:64:ALA:HA	23:11:67:ILE:HG13	1.98	0.45
1:1A:1067:A:H3'	1:1A:1067:A:N3	2.32	0.45
1:1A:1604:C:H5''	1:1A:1605:A:OP2	2.16	0.45
1:1A:1733:C:H2'	1:1A:1734:G:O4'	2.16	0.45
1:1A:1952:G:O2'	1:1A:1990:G:O6	2.23	0.45
6:1G:47:LYS:HG3	6:1G:48:GLU:H	1.81	0.45
1:1A:2331:G:N2	14:1S:3:ARG:HA	2.32	0.45
1:2A:1374:G:H2'	1:2A:1375:C:C6	2.52	0.45
1:2A:2408:U:H2'	1:2A:2409:G:C8	2.51	0.45
1:2A:644:A:H4'	1:2A:645:C:C4	2.52	0.45
13:2R:33:ARG:NH1	13:2R:115:GLU:OE1	2.43	0.45
23:11:56:GLN:HB2	23:11:90:ILE:HD12	1.99	0.45
1:1A:1140:U:N3	1:1A:1142:A:H5'	2.31	0.45
1:1A:273:G:O2'	1:1A:274:U:H5''	2.17	0.45
1:1A:945:A:N3	1:1A:945:A:H2'	2.32	0.45
3:1D:8:PRO:HB3	3:1D:14:ARG:HG2	1.98	0.45
19:1X:12:VAL:HG21	19:1X:27:THR:HG22	1.99	0.45
1:2A:1300:U:H4'	1:2A:1301:A:H5''	1.98	0.45
1:2A:191:A:H2'	1:2A:192:C:C6	2.51	0.45
1:2A:2029:G:H2'	1:2A:2031:A:OP1	2.15	0.45
1:2A:2137:C:N3	1:2A:2155:G:N1	2.65	0.45
1:2A:2512:C:H2'	1:2A:2513:G:O4'	2.16	0.45
11:2P:121:LYS:HD3	11:2P:123:LEU:HD11	1.99	0.45
12:2Q:111:GLU:OE1	12:2Q:133:ARG:NH2	2.49	0.45
16:2U:8:VAL:O	16:2U:12:ARG:HG3	2.17	0.45
1:1A:1115:A:H2	1:1A:1141:A:HO2'	1.65	0.45
1:1A:2148:A:N3	1:1A:2149:G:H1'	2.32	0.45
1:1A:2159:C:H2'	1:1A:2160:C:C6	2.52	0.45
1:1A:2661:U:H2'	1:1A:2662:U:C6	2.51	0.45
21:1Z:121:HIS:HB3	21:1Z:123:ASP:O	2.17	0.45
1:2A:2405:G:H5'	11:2P:75:ILE:HD13	1.99	0.45

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:2769:C:H2'	1:2A:2770:G:O4'	2.17	0.45
1:2A:320:A:H4'	1:2A:322:A:N7	2.32	0.45
1:2A:521:G:H2'	1:2A:522:G:H8	1.82	0.45
1:2A:709:U:H2'	1:2A:710:G:C8	2.52	0.45
1:2A:882:G:H1	1:2A:894:C:N4	2.08	0.45
2:2B:57:A:N3	6:2G:29:TRP:HB3	2.32	0.45
7:2H:46:GLU:O	7:2H:48:GLY:N	2.50	0.45
8:2I:93:THR:HG22	8:2I:119:PRO:HB3	1.98	0.45
9:2N:123:TYR:CZ	9:2N:129:PRO:HD2	2.52	0.45
10:2O:63:VAL:HG23	10:2O:64:ARG:HG3	1.98	0.45
1:1A:1104:G:N1	1:1A:1126:C:N4	2.36	0.45
1:1A:1128:U:C4	1:1A:1132:A:N1	2.76	0.45
1:1A:1882:U:H2'	1:1A:1883:C:O4'	2.17	0.45
1:1A:756:U:H2'	1:1A:757:G:C8	2.52	0.45
8:1I:5:LEU:HD11	8:1I:19:VAL:HG22	1.98	0.45
10:1O:10:VAL:HG21	10:1O:16:ALA:C	2.36	0.45
14:1S:34:HIS:O	14:1S:97:ARG:NH2	2.50	0.45
16:1U:58:ARG:HA	16:1U:61:TRP:CE3	2.52	0.45
1:1A:353:G:OP2	20:1Y:71:LYS:HD2	2.17	0.45
1:2A:108:U:H2'	1:2A:109:G:C8	2.52	0.45
1:2A:1991:U:H2'	1:2A:1992:G:H5''	1.99	0.45
1:2A:2291:U:H2'	1:2A:2292:C:C6	2.51	0.45
1:2A:455:C:N3	1:2A:472:A:H2'	2.32	0.45
3:2D:108:PRO:HG2	3:2D:111:LEU:HB2	1.99	0.45
4:2E:30:PRO:HB3	4:2E:92:THR:HG22	1.99	0.45
13:2R:54:LEU:HB3	13:2R:62:ALA:HB1	1.98	0.45
58:B:4:GLY:HA2	58:B:13:ALA:HB2	1.98	0.45
1:1A:1715:A:H4'	1:1A:1716:A:O5'	2.17	0.44
1:1A:2087:C:H2'	1:1A:2088:C:C6	2.52	0.44
1:1A:2102:G:OP1	23:1I:35:THR:HG21	2.17	0.44
1:1A:2315:G:O6	63:1A:4359:HOH:O	2.21	0.44
1:1A:645:G:H5'	1:1A:645:G:N3	2.31	0.44
3:1D:108:PRO:HB3	3:1D:143:HIS:CE1	2.51	0.44
5:1F:184:TYR:CE2	5:1F:188:ARG:HD2	2.52	0.44
8:1I:60:GLU:HG3	8:1I:61:ARG:NH1	2.32	0.44
13:1R:117:VAL:HG12	13:1R:118:GLU:H	1.82	0.44
14:1S:110:LEU:HA	14:1S:110:LEU:HD12	1.77	0.44
15:1T:60:THR:HG22	15:1T:77:PRO:HA	1.98	0.44
1:2A:857:C:H1'	22:20:26:TYR:HE1	1.81	0.44
1:2A:1851:U:H2'	1:2A:1852:C:O4'	2.17	0.44
1:2A:2238:G:N3	1:2A:2238:G:H2'	2.32	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:236:C:H2'	1:2A:237:C:C6	2.51	0.44
3:2D:77:ALA:HB2	3:2D:97:TYR:CD1	2.51	0.44
4:2E:36:ARG:HG2	4:2E:47:VAL:HG12	1.99	0.44
5:2F:106:ARG:HG2	5:2F:106:ARG:H	1.50	0.44
24:12:32:LEU:HD12	24:12:57:ILE:HD12	1.98	0.44
1:1A:1552:C:H2'	1:1A:1553:A:C8	2.52	0.44
1:1A:672:G:H2'	1:1A:673:G:O4'	2.17	0.44
3:1D:34:VAL:HG12	3:1D:63:ARG:HG3	1.99	0.44
22:20:53:MET:HG3	22:20:59:LEU:HD23	1.98	0.44
1:2A:1002:G:N3	1:2A:1003:G:H1'	2.95	0.44
1:2A:1448:G:H4'	1:2A:1542:A:OP1	2.18	0.44
1:2A:2168:G:C8	1:2A:2170:A:N7	2.85	0.44
1:2A:77:C:OP1	24:22:59:ARG:NE	2.46	0.44
2:2B:27:C:N4	2:2B:56:G:O6	2.51	0.44
3:2D:37:LEU:HD13	3:2D:87:ASN:ND2	2.32	0.44
8:2I:110:ASP:N	8:2I:130:TYR:OH	2.36	0.44
8:2I:38:LEU:H	8:2I:38:LEU:HG	1.50	0.44
13:2R:38:VAL:HG22	13:2R:112:ALA:HB2	1.99	0.44
1:1A:1338:U:H2'	1:1A:1339:C:C6	2.53	0.44
1:1A:1469:G:H2'	1:1A:1470:G:C8	2.82	0.44
1:1A:2473:C:H2'	1:1A:2474:U:C6	2.52	0.44
2:1B:78:A:C2	2:1B:100:A:C4	3.04	0.44
2:1B:24:G:N7	2:1B:56:G:H2'	2.32	0.44
5:1F:53:THR:HG22	5:1F:56:GLU:HG3	1.99	0.44
16:1U:29:SER:O	16:1U:30:LYS:HD3	2.17	0.44
16:1U:86:ALA:O	17:1V:49:THR:HG23	2.17	0.44
29:27:47:ARG:HA	29:27:47:ARG:HD3	1.60	0.44
1:2A:2507:C:H5''	1:2A:2573:C:C4	2.52	0.44
1:2A:375:C:H2'	1:2A:376:C:C6	2.53	0.44
8:2I:9:LEU:HD21	8:2I:35:LEU:HD13	2.00	0.44
21:2Z:76:LEU:HA	21:2Z:83:PRO:HA	1.99	0.44
26:14:17:GLY:C	26:14:19:GLY:H	2.20	0.44
1:1A:1405:A:H2'	1:1A:1406:A:H5'	1.99	0.44
1:1A:2346:G:H4'	1:1A:2347:A:OP2	2.17	0.44
1:1A:34:C:H5''	1:1A:35:G:OP2	2.17	0.44
1:1A:664:U:H2'	1:1A:665:C:C6	2.53	0.44
1:2A:2484:G:C2	1:2A:2485:G:C8	3.05	0.44
1:2A:258:G:H2'	1:2A:259:G:H8	2.33	0.44
1:2A:660:G:H5'	5:2F:99:TYR:CE1	2.53	0.44
1:2A:754:C:H2'	1:2A:755:C:C6	2.51	0.44
1:2A:878:A:H61	1:2A:899:A:H1'	1.82	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
14:2S:64:GLU:CD	14:2S:64:GLU:H	4.02	0.44
16:2U:76:TYR:CE2	16:2U:80:ILE:HG13	2.53	0.44
17:2V:40:LEU:HB2	17:2V:46:VAL:CG1	2.47	0.44
20:2Y:8:LYS:HD3	20:2Y:97:ARG:NH1	2.33	0.44
27:15:16:ARG:HD2	27:15:20:ARG:NH1	2.33	0.44
1:1A:1810:U:H2'	63:1A:4705:HOH:O	2.17	0.44
1:1A:2150:C:H42	1:1A:2182:G:H1	1.66	0.44
8:1I:4:ILE:HG12	8:1I:18:VAL:HG22	2.00	0.44
12:1Q:60:ARG:HB2	12:1Q:60:ARG:NH1	2.31	0.44
1:2A:141:A:H8	1:2A:1408:C:HO2'	1.56	0.44
1:2A:1839:G:C8	1:2A:1927:A:H1'	2.52	0.44
1:2A:983:A:H3'	1:2A:983:A:N3	5.13	0.44
13:2R:28:LEU:HD23	13:2R:48:VAL:HG21	1.99	0.44
21:2Z:33:LEU:HD21	21:2Z:90:VAL:HG11	1.98	0.44
1:1A:1827:U:H2'	1:1A:1828:C:H6	1.83	0.44
1:1A:2735:G:H2'	1:1A:2736:C:C6	2.52	0.44
1:1A:541:C:OP1	27:15:13:LYS:NZ	2.38	0.44
2:1B:66:A:H61	2:1B:109:C:H5'	1.83	0.44
3:1D:77:ALA:HA	3:1D:97:TYR:HA	1.98	0.44
5:1F:52:LYS:HD3	5:1F:56:GLU:O	2.17	0.44
6:1G:7:LEU:HD21	6:1G:176:LEU:HD22	2.00	0.44
21:1Z:15:PRO:O	21:1Z:19:ARG:HG3	2.17	0.44
23:21:67:ILE:N	23:21:68:PRO:HD2	2.32	0.44
1:2A:1530:C:H1'	1:2A:1531:C:OP1	2.18	0.44
1:2A:2327:A:H2'	1:2A:2328:A:O4'	2.18	0.44
1:2A:251:A:C5	1:2A:252:G:H1'	2.53	0.44
1:2A:271(T):C:H2'	1:2A:271(U):G:H8	1.82	0.44
1:2A:340:A:H2'	1:2A:341:G:O4'	2.17	0.44
2:2B:75:G:H22	21:2Z:73:GLN:NE2	2.14	0.44
4:2E:170:LEU:HB3	4:2E:184:VAL:HG22	2.00	0.44
11:2P:90:ARG:NH1	11:2P:105:LEU:HD11	2.33	0.44
1:1A:2018:C:H4'	1:1A:2019:G:OP1	2.18	0.44
1:1A:2140:U:OP1	1:1A:2170:G:H4'	2.17	0.44
1:1A:928:G:H3'	1:1A:929:G:H8	1.81	0.44
11:1P:101:VAL:HG23	11:1P:106:LEU:O	2.18	0.44
28:26:35:GLU:OE2	28:26:50:ARG:NH1	2.45	0.44
1:2A:1263:U:C4	1:2A:1264:G:C6	3.06	0.44
1:2A:2540:C:H2'	1:2A:2541:A:O4'	2.18	0.44
1:2A:277:C:H4'	1:2A:278:A:H8	1.82	0.44
2:2B:94:C:H2'	2:2B:95:C:H6	1.82	0.44
6:2G:106:LEU:O	6:2G:111:LEU:HG	2.18	0.44

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:2G:122:PRO:HG3	6:2G:182:LYS:H	1.83	0.44
6:2G:12:TYR:HA	6:2G:16:ARG:HG3	2.00	0.44
7:2H:127:GLU:HB3	7:2H:129:THR:HG22	2.00	0.44
8:2I:92:VAL:HG13	8:2I:120:ILE:HB	1.98	0.44
1:1A:1108:G:N2	1:1A:1134:A:C6	2.85	0.44
1:1A:747:G:O2'	1:1A:1679:A:N3	2.38	0.44
1:1A:2724:U:O2'	1:1A:2726:A:H5'	2.18	0.44
1:1A:2761:A:H5'	7:1H:4:ILE:HD12	1.99	0.44
11:2P:59:LEU:HD12	30:28:58:ILE:HG12	2.00	0.44
1:2A:1777:U:H2'	1:2A:1778:U:C6	2.52	0.44
1:2A:184:C:H2'	1:2A:185:U:C6	2.53	0.44
1:2A:673:C:H5''	5:2F:81:PRO:HD2	1.99	0.44
11:2P:126:VAL:HG12	11:2P:148:LEU:HD22	2.00	0.44
12:2Q:118:LEU:HD12	12:2Q:131:ILE:HG23	1.99	0.44
1:2A:952:G:OP1	12:2Q:16:ARG:NH2	2.51	0.44
15:2T:93:ARG:HH22	15:2T:95:ARG:HH21	1.65	0.44
18:2W:45:TYR:CZ	18:2W:49:LYS:HE2	2.53	0.44
2:2B:96:U:OP1	21:2Z:14:LYS:NZ	2.50	0.44
1:1A:1653:C:H4'	1:1A:1654:A:O5'	2.18	0.44
1:1A:459:A:N6	63:1A:4551:HOH:O	2.47	0.44
2:1B:2:C:H2'	2:1B:3:C:H6	1.83	0.44
4:1E:97:LYS:HE2	4:1E:97:LYS:HB3	1.68	0.44
10:1O:64:ARG:HD2	10:1O:79:PHE:CD1	2.53	0.44
14:1S:5:THR:OG1	14:1S:8:GLU:HG2	2.18	0.44
21:1Z:65:GLN:OE1	21:1Z:67:LEU:HD21	2.18	0.44
1:2A:579:G:H2'	1:2A:580:C:C6	2.53	0.44
1:2A:740:U:H2'	1:2A:741:G:C8	2.53	0.44
2:2B:19:G:H2'	2:2B:20:C:O4'	2.18	0.44
3:2D:71:ASP:CB	3:2D:103:ARG:HH12	2.30	0.44
4:2E:181:LEU:HA	4:2E:181:LEU:HD12	1.68	0.44
19:2X:94:GLY:H	19:2X:95:LEU:HA	1.83	0.44
1:1A:1090:G:H5'	1:1A:1091:A:OP2	2.17	0.43
1:1A:1248:G:H5'	11:1P:3:LEU:HD23	1.99	0.43
1:1A:1535:U:O3'	1:1A:1536:A:H8	2.01	0.43
1:1A:2193:A:O2'	1:1A:2194:U:H6	2.01	0.43
1:1A:2210:C:H2'	1:1A:2211:U:O4'	2.18	0.43
1:1A:2697:G:H5'	10:1O:68:GLU:OE2	2.18	0.43
8:1I:62:LYS:O	8:1I:66:GLU:HG2	2.18	0.43
9:1N:46:VAL:HG23	9:1N:48:MET:HE3	2.00	0.43
13:1R:44:LEU:HA	13:1R:44:LEU:HD23	1.80	0.43
30:28:62:LEU:HB3	30:28:65:GLU:HG2	2.00	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:108:U:H2'	1:2A:109:G:H8	1.82	0.43
1:2A:2031:A:C6	1:2A:2498:C:H1'	2.53	0.43
1:2A:42:G:H2'	1:2A:43:A:O4'	2.18	0.43
2:2B:83:G:N2	2:2B:94:C:O2	2.33	0.43
19:2X:94:GLY:N	19:2X:95:LEU:HA	2.33	0.43
58:B:2:SER:HA	58:B:3:PRO:HD3	1.74	0.43
1:1A:2327:G:H2'	1:1A:2328:C:C6	2.52	0.43
1:1A:933:C:N3	1:1A:934:A:H8	2.15	0.43
1:1A:925:A:N6	1:1A:946:A:C8	2.86	0.43
1:1A:977:G:OP2	25:13:29:ARG:NH2	2.51	0.43
4:1E:47:VAL:O	4:1E:80:GLU:HA	2.19	0.43
1:2A:1935:G:H1'	1:2A:1964:G:N2	2.33	0.43
1:2A:2432:A:C6	1:2A:2433:A:C6	3.06	0.43
1:2A:271(E):U:H2'	1:2A:271(F):C:C6	2.53	0.43
1:2A:932:G:H4'	1:2A:933:A:O5'	2.18	0.43
7:2H:35:VAL:HG13	7:2H:71:LEU:HD23	2.00	0.43
11:2P:97:PRO:HD3	11:2P:126:VAL:O	2.18	0.43
13:2R:36:THR:HG22	13:2R:37:THR:H	1.83	0.43
29:17:12:ARG:NH2	29:17:44:PRO:HB3	2.33	0.43
1:1A:1093:G:O2'	1:1A:1094:A:O5'	2.31	0.43
1:1A:320:C:H2'	1:1A:321:C:C6	2.54	0.43
1:1A:455:A:H8	1:1A:455:A:OP2	2.01	0.43
1:1A:831:A:C5	3:1D:229:VAL:HG21	2.54	0.43
21:1Z:105:VAL:O	21:1Z:141:VAL:HG22	2.18	0.43
1:2A:1779:U:H2'	63:2A:4223:HOH:O	2.18	0.43
1:2A:2161:C:H2'	1:2A:2162:G:O4'	2.18	0.43
1:2A:2687:U:H2'	1:2A:2688:U:O4'	2.19	0.43
1:2A:2812:G:H2'	1:2A:2813:A:C8	2.53	0.43
1:2A:2819:G:H2'	1:2A:2821:A:N7	2.33	0.43
1:2A:18:C:O2'	1:2A:554:U:OP1	2.32	0.43
6:2G:111:LEU:HB3	6:2G:117:PHE:CE1	2.53	0.43
7:2H:25:LYS:HB2	7:2H:25:LYS:HE3	1.69	0.43
1:1A:1091:A:OP1	1:1A:1091:A:H4'	2.17	0.43
1:1A:1370:G:C2	1:1A:1371:G:C8	13.53	0.43
1:1A:196:A:H2'	1:1A:197:C:O4'	2.17	0.43
1:1A:386:U:H2'	1:1A:386:U:H6	1.63	0.43
1:1A:390:G:H2'	1:1A:391:G:C8	2.52	0.43
1:1A:2455:C:OP1	5:1F:68:LYS:HD3	2.19	0.43
1:1A:1696:G:O2'	13:1R:107:ASP:OD2	2.21	0.43
1:2A:686:G:H8	29:27:6:GLN:O	2.02	0.43
1:2A:1359:A:C2	1:2A:1372:U:O4	2.71	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
10:2O:105:GLU:N	10:2O:105:GLU:OE1	2.50	0.43
1:2A:1187:G:H5'	17:2V:81:TYR:CE1	2.53	0.43
1:1A:1530:G:OP1	1:1A:1530:G:H4'	4.96	0.43
1:1A:2177:G:H3'	1:1A:2178:G:C8	2.53	0.43
1:1A:2331:G:H22	14:1S:3:ARG:NE	2.17	0.43
1:2A:2336:A:H61	22:20:43:THR:HG22	1.83	0.43
9:2N:21:LYS:NZ	9:2N:140:VAL:OXT	2.45	0.43
15:2T:51:ARG:HG2	15:2T:62:THR:HB	2.01	0.43
21:2Z:54:HIS:HB3	21:2Z:101:PRO:HD3	2.00	0.43
1:1A:1318:A:H5''	14:1S:3:ARG:NH1	127.34	0.43
1:1A:1566:U:H2'	1:1A:1567:G:O4'	2.18	0.43
1:1A:1961:5MU:OP1	1:1A:2616:U:O2'	2.29	0.43
1:1A:540:A:H1'	1:1A:604:C:H1'	2.00	0.43
1:1A:649:C:O2'	1:1A:704:U:OP1	2.35	0.43
7:1H:62:LYS:HB3	7:1H:62:LYS:HE2	1.84	0.43
1:2A:783:A:OP2	63:2A:3959:HOH:O	2.21	0.43
1:2A:686:G:H21	1:2A:788:A:H61	1.65	0.43
7:2H:11:VAL:HG21	7:2H:50:VAL:HG23	2.01	0.43
7:2H:28:GLY:HA3	7:2H:79:VAL:HB	2.00	0.43
11:1P:63:PRO:HD3	30:18:27:THR:HG22	2.00	0.43
1:1A:1273:G:OP1	16:1U:13:LYS:HG2	2.18	0.43
1:1A:2519:C:H2'	1:1A:2520:G:O4'	2.19	0.43
1:1A:2647:C:H4'	4:1E:48:GLN:HE21	1.83	0.43
30:28:6:THR:HG22	30:28:63:PRO:HD2	2.01	0.43
1:2A:2191:G:H2'	1:2A:2192:G:O4'	2.18	0.43
1:2A:2547:U:O2	10:2O:23:ARG:NH2	2.46	0.43
1:2A:493:G:H2'	1:2A:494:G:O4'	2.18	0.43
1:2A:570:G:H2'	1:2A:2030:A:C5	2.53	0.43
1:2A:826:U:H4'	11:2P:55:ARG:HB3	2.00	0.43
2:2B:11:C:H3'	2:2B:12:C:C6	2.53	0.43
9:2N:28:THR:HG22	9:2N:29:LYS:N	2.33	0.43
9:2N:20:GLY:HA2	9:2N:61:ARG:HG3	2.00	0.43
1:2A:571:A:O2'	17:2V:78:LYS:HE2	2.19	0.43
20:2Y:97:ARG:HG2	20:2Y:97:ARG:H	1.62	0.43
1:1A:1410:G:OP2	23:11:3:LYS:HG3	2.19	0.43
1:1A:202:A:H2'	1:1A:203:G:O4'	2.18	0.43
1:1A:2158:C:N3	1:1A:2177:G:C2	2.84	0.43
1:1A:2155:G:HO2'	1:1A:2178:G:N2	2.17	0.43
1:1A:271:U:H1'	8:1I:50:ARG:NH1	2.33	0.43
1:1A:515:G:N7	18:1W:49:LYS:NZ	2.67	0.43
1:1A:956:A:N3	1:1A:2276:C:O2'	2.45	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
5:1F:32:LEU:HD22	5:1F:112:MET:HE3	2.01	0.43
19:1X:44:GLU:HG2	19:1X:49:VAL:O	2.18	0.43
1:2A:2037:G:H2'	1:2A:2038:G:C8	2.54	0.43
1:2A:2103:C:H2'	1:2A:2104:G:C8	2.54	0.43
6:2G:106:LEU:HA	6:2G:110:ALA:HB3	2.01	0.43
6:2G:111:LEU:HA	6:2G:114:ILE:HD12	2.01	0.43
6:2G:17:PRO:HA	6:2G:20:ILE:HD12	2.00	0.43
10:2O:68:GLU:HB3	10:2O:78:ARG:HB2	2.01	0.43
15:2T:26:ASP:O	15:2T:49:VAL:HG22	2.19	0.43
1:2A:748:G:C8	18:2W:89:ALA:HB1	2.54	0.43
19:2X:41:ASN:O	19:2X:45:THR:HG23	2.19	0.43
21:2Z:99:TYR:HA	21:2Z:124:ILE:O	2.18	0.43
3:1D:182:LEU:HD23	3:1D:182:LEU:HA	1.86	0.43
7:1H:25:LYS:HB2	7:1H:25:LYS:HE3	1.86	0.43
7:1H:3:ARG:NH1	7:1H:5:GLY:H	2.17	0.43
8:1I:61:ARG:HA	8:1I:61:ARG:HD3	1.78	0.43
20:1Y:14:LEU:HB2	20:1Y:75:ILE:HD11	2.00	0.43
21:1Z:138:GLU:H	21:1Z:156:LYS:HE2	1.84	0.43
22:20:6:GLY:O	22:20:7:LEU:HD23	2.18	0.43
1:2A:107:C:H2'	1:2A:108:U:H6	1.84	0.43
1:2A:1231:G:H2'	1:2A:1232:G:C8	2.54	0.43
1:2A:2023:G:H5'	1:2A:2617:C:H4'	2.01	0.43
1:2A:2065:C:H4'	1:2A:2251:OMG:HM22	2.01	0.43
2:2B:68:C:H2'	2:2B:69:G:H8	1.83	0.43
6:2G:114:ILE:HG13	6:2G:140:ILE:HG12	2.01	0.43
7:2H:16:SER:OG	7:2H:27:LYS:HB2	2.19	0.43
1:1A:1735:U:O2'	1:1A:1747:A:N7	2.43	0.43
1:1A:1785:C:OP1	15:1T:96:ARG:NH1	2.48	0.43
1:1A:1817:A:H8	63:1A:4619:HOH:O	2.02	0.43
1:1A:559:U:H2'	1:1A:560:C:C6	2.54	0.43
1:1A:891:C:C2'	1:1A:892:G:H5'	2.49	0.43
8:1I:62:LYS:HB2	8:1I:62:LYS:HE3	1.67	0.43
1:2A:143:G:H2'	1:2A:143(A):C:H6	1.82	0.43
1:2A:1493:C:N4	1:2A:2206:G:O2'	2.49	0.43
1:2A:197:A:N6	1:2A:2430:A:H2'	2.34	0.43
1:2A:2683:C:O2	10:2O:70:LYS:NZ	2.48	0.43
1:2A:311:A:C6	1:2A:328:U:C4	3.06	0.43
1:2A:415:A:H2'	1:2A:416:C:O4'	2.19	0.43
1:2A:612:C:C2	1:2A:616:G:N2	2.87	0.43
1:2A:862:G:H2'	1:2A:863:A:O4'	2.19	0.43
2:2B:28:C:H2'	2:2B:29:A:O4'	2.19	0.43

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:2318:G:H21	14:2S:3:ARG:HE	1.67	0.43
17:2V:1:MET:HE3	17:2V:44:LYS:H	1.83	0.43
21:2Z:70:LEU:HA	21:2Z:70:LEU:HD23	1.75	0.43
1:1A:2331:G:C2	14:1S:3:ARG:HA	2.53	0.42
1:1A:2430:A:H2'	1:1A:2431:U:C6	2.54	0.42
1:1A:388:A:H2'	1:1A:389:G:C8	2.53	0.42
3:1D:70:TRP:HB3	3:1D:190:TYR:CE2	2.53	0.42
7:1H:104:GLU:HG3	7:1H:114:VAL:HG22	2.01	0.42
7:1H:113:VAL:HG11	7:1H:151:ILE:HG21	2.00	0.42
13:1R:50:HIS:ND1	63:1R:301:HOH:O	2.22	0.42
16:1U:76:TYR:CE2	16:1U:80:ILE:HG13	2.54	0.42
1:2A:1412:A:H2'	1:2A:1413:G:C8	2.53	0.42
1:2A:2145:C:O2'	1:2A:2147:G:N7	2.50	0.42
1:2A:2327:A:H2'	1:2A:2328:A:C8	2.54	0.42
1:2A:848:G:C4	1:2A:933:A:H8	2.37	0.42
1:2A:993:G:H2'	1:2A:993:G:N3	2.85	0.42
1:2A:1799:G:O2'	3:2D:181:GLU:OE2	2.28	0.42
3:2D:77:ALA:HA	3:2D:97:TYR:HA	2.01	0.42
1:2A:1188:U:C4'	17:2V:79:VAL:HG22	2.48	0.42
1:1A:484:G:O2'	29:17:39:ARG:HD2	2.18	0.42
1:1A:1016:C:H2'	1:1A:1017:G:O4'	2.18	0.42
1:1A:2164:C:N3	1:1A:2171:G:O6	2.52	0.42
1:1A:449:A:H2'	1:1A:450:A:C8	2.54	0.42
1:1A:572:A:H1'	1:1A:573:G:OP1	2.20	0.42
1:1A:662:A:OP1	11:1P:133:SER:OG	2.34	0.42
1:1A:964:A:H5''	2:1B:98:G:O2'	2.18	0.42
3:1D:2:ALA:O	3:1D:20:ASP:HB3	2.19	0.42
8:1I:86:THR:O	8:1I:123:LEU:HB2	2.19	0.42
9:1N:96:GLU:CD	9:1N:96:GLU:H	2.23	0.42
31:29:33:LYS:HE3	31:29:33:LYS:HB2	1.86	0.42
1:2A:1628:G:H2'	1:2A:1629:U:C6	2.54	0.42
1:2A:524:U:H2'	1:2A:525:U:C6	2.54	0.42
1:2A:652(T):C:H2'	1:2A:652(U):G:C8	2.54	0.42
2:2B:119:G:H5'	2:2B:120:A:OP2	2.19	0.42
3:2D:26:LYS:HB3	3:2D:83:GLU:HG2	2.01	0.42
5:2F:64:ILE:HG21	5:2F:78:ILE:HG23	2.00	0.42
1:2A:586:A:H5'	5:2F:89:VAL:HG21	2.01	0.42
30:18:6:THR:HG22	30:18:64:TYR:HD2	1.84	0.42
1:1A:119:G:H4'	1:1A:149:A:H5'	2.00	0.42
1:1A:2149:G:N2	1:1A:2183:C:N3	2.56	0.42
2:1B:8:U:O3'	14:1S:25:ARG:NH2	2.52	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:1834:A:H4'	3:1D:259:THR:HG23	2.01	0.42
30:28:63:PRO:HG2	30:28:64:TYR:CE2	2.54	0.42
1:2A:2140:C:H2'	1:2A:2141:G:H5'	2.01	0.42
1:2A:271(S):G:C6	1:2A:271(T):C:C4	3.08	0.42
1:2A:412:A:N6	1:2A:2412:A:O4'	2.52	0.42
1:2A:478:A:N1	1:2A:500:G:H4'	2.35	0.42
9:2N:103:VAL:HG11	9:2N:120:LEU:HD22	2.01	0.42
19:2X:44:GLU:HG2	19:2X:49:VAL:O	2.20	0.42
58:A:2:SER:HA	58:A:3:PRO:HD3	1.63	0.42
1:1A:1314:A:H2'	1:1A:1315:A:O4'	2.19	0.42
1:1A:174:U:H4'	1:1A:207:A:H4'	2.02	0.42
1:1A:2429:C:OP1	11:1P:65:ARG:NH2	2.51	0.42
6:1G:43:LEU:HB3	6:1G:44:GLY:H	1.61	0.42
8:1I:84:GLY:O	8:1I:86:THR:N	2.51	0.42
1:1A:70:A:N7	19:1X:31:HIS:HE1	2.17	0.42
1:2A:2061:G:H5''	1:2A:2503:2MA:C2	2.49	0.42
1:2A:244:A:C2	1:2A:255:A:C4	3.08	0.42
9:2N:38:HIS:NE2	9:2N:50:ASP:OD2	2.53	0.42
11:2P:135:LEU:HA	11:2P:135:LEU:HD23	1.80	0.42
14:2S:87:PHE:CE1	14:2S:102:ALA:HB2	2.54	0.42
58:B:3:PRO:HG2	58:B:12:SER:HB2	2.01	0.42
25:13:31:LEU:HD23	25:13:31:LEU:HA	1.90	0.42
1:1A:1110:C:N3	1:1A:1120:G:O6	2.53	0.42
1:1A:116:A:C8	1:1A:117:A:C8	3.07	0.42
1:1A:1941:A:OP2	1:1A:1942:4OC:H5	2.19	0.42
1:1A:467:U:O2	5:1F:46:ARG:NH2	2.40	0.42
6:1G:7:LEU:HD12	6:1G:104:GLU:HA	2.01	0.42
9:1N:73:THR:HG23	9:1N:82:LEU:HD11	2.02	0.42
9:1N:68:GLU:HG3	9:1N:88:GLU:OE2	2.19	0.42
13:1R:111:LEU:HD12	13:1R:111:LEU:HA	1.86	0.42
1:2A:1006:C:C2	1:2A:1138:G:N2	2.87	0.42
1:2A:1507:A:O2'	1:2A:1508:A:C8	2.72	0.42
1:2A:1581:G:H2'	1:2A:1582:C:O4'	2.19	0.42
1:2A:2243:U:H2'	1:2A:2244:U:C6	2.55	0.42
1:2A:2070:G:C2	1:2A:2442:C:C2	3.08	0.42
1:2A:288:C:H2'	1:2A:289:A:C8	2.54	0.42
1:2A:956:G:H2'	1:2A:957:A:H2'	2.01	0.42
6:2G:173:LEU:O	6:2G:178:PHE:N	2.43	0.42
7:2H:56:SER:OG	7:2H:61:HIS:ND1	2.48	0.42
14:2S:5:THR:OG1	14:2S:8:GLU:HG2	2.20	0.42
25:13:7:LYS:HB2	25:13:34:GLU:HG2	2.02	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:1387:U:O4'	19:1X:57:LEU:HD23	2.19	0.42
1:1A:1400:A:H4'	3:1D:38:LYS:HZ3	1.85	0.42
1:1A:1452:U:H2'	1:1A:1453:C:H6	1.82	0.42
1:1A:1692:G:H5''	1:1A:1693:C:H5'	2.01	0.42
1:1A:1766:G:H1'	1:1A:1770:A:N6	2.35	0.42
1:1A:1959:A:H1'	1:1A:1961:5MU:H73	2.01	0.42
1:1A:794:U:O2	1:1A:2036:A:H1'	2.20	0.42
5:1F:161:GLU:O	5:1F:165:ARG:HB2	2.19	0.42
9:1N:61:ARG:HD3	9:1N:61:ARG:HA	1.82	0.42
1:1A:795:G:C8	18:1W:89:ALA:HB1	2.55	0.42
1:2A:300:A:P	20:2Y:86:ARG:NH2	2.92	0.42
6:2G:11:TYR:OH	6:2G:16:ARG:HD3	2.18	0.42
7:2H:154:PRO:HB3	7:2H:163:TYR:CE1	2.55	0.42
16:2U:58:ARG:HA	16:2U:61:TRP:CE3	2.55	0.42
1:1A:2897:U:H2'	1:1A:2898:C:C6	2.55	0.42
1:1A:320:C:H2'	1:1A:321:C:H6	1.85	0.42
1:1A:32:C:O2'	1:1A:33:U:H5'	2.20	0.42
1:1A:848:G:O6	5:1F:53:THR:OG1	2.36	0.42
1:1A:993:G:H2'	1:1A:993:G:N3	2.91	0.42
10:1O:23:ARG:HA	10:1O:23:ARG:HD2	1.85	0.42
30:28:55:ALA:O	30:28:59:LYS:HG3	2.19	0.42
1:2A:1011:G:H1	1:2A:1018:C:N4	18.19	0.42
1:2A:1423:G:OP1	10:2O:49:ARG:NH2	98.10	0.42
1:2A:208:C:H2'	1:2A:209:C:C6	2.55	0.42
1:2A:2136:C:O2'	1:2A:2137:C:H6	2.03	0.42
1:2A:717:G:H2'	1:2A:718:A:O4'	2.20	0.42
1:2A:765:G:N1	1:2A:812:C:O2'	83.84	0.42
1:2A:921:G:C6	1:2A:922:U:C4	3.08	0.42
4:2E:35:GLN:OE1	4:2E:66:HIS:HE1	2.03	0.42
7:2H:156:ALA:O	7:2H:172:LYS:HG2	2.19	0.42
8:2I:124:GLY:N	8:2I:144:VAL:HG23	2.35	0.42
14:2S:24:LEU:HB2	14:2S:85:VAL:HG23	2.02	0.42
24:12:52:ASP:O	24:12:56:GLN:HG3	2.20	0.42
1:1A:105:C:H2'	1:1A:106:U:H6	1.83	0.42
1:1A:866:A:C4	1:1A:1234:A:C2	3.08	0.42
1:1A:1463:C:H2'	1:1A:1464:G:O4'	2.19	0.42
1:1A:1897:C:H2'	1:1A:1898:A:O4'	2.19	0.42
1:1A:78:G:O6	1:1A:92:C:N4	28.83	0.42
1:1A:977:G:H4'	1:1A:978:A:O5'	2.20	0.42
9:1N:58:ASP:OD1	9:1N:58:ASP:N	2.52	0.42
31:29:10:ILE:HD12	31:29:32:HIS:HA	2.00	0.42

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:1791:A:OP2	1:2A:1791:A:H8	2.03	0.42
1:2A:884:C:H3'	1:2A:885:C:C6	2.55	0.42
2:2B:17:C:H2'	2:2B:18:G:O4'	2.20	0.42
5:2F:178:PRO:HB3	5:2F:198:ALA:HA	2.02	0.42
7:2H:3:ARG:CZ	7:2H:5:GLY:H	2.32	0.42
21:2Z:8:TYR:HB2	21:2Z:38:TYR:CE2	2.55	0.42
1:1A:1140:U:H3	1:1A:1142:A:H5'	1.85	0.42
1:1A:2044:U:OP1	63:1A:4361:HOH:O	2.22	0.42
1:1A:2148:A:N1	1:1A:2184:G:O2'	2.35	0.42
1:1A:2190:G:C6	1:1A:2193:A:C8	3.08	0.42
1:1A:733:G:H8	29:17:6:GLN:O	2.03	0.42
3:1D:68:LYS:HB2	3:1D:70:TRP:CE2	2.55	0.42
6:1G:125:PHE:HB2	63:1G:5005:HOH:O	2.19	0.42
23:21:23:LYS:HB3	23:21:29:GLY:HA3	2.02	0.42
31:29:17:ILE:HA	31:29:17:ILE:HD12	1.78	0.42
1:2A:2375:G:O2'	1:2A:2377:A:N7	2.36	0.42
1:2A:2419:U:H2'	1:2A:2420:C:C6	2.55	0.42
1:2A:271(K):U:H4'	1:2A:271(L):U:OP2	2.20	0.42
1:2A:606:U:H4'	1:2A:658:C:H4'	2.00	0.42
1:2A:747:U:O2	1:2A:2014:A:H1'	2.19	0.42
7:2H:154:PRO:HB3	7:2H:163:TYR:CZ	2.55	0.42
8:2I:53:ALA:O	8:2I:57:ARG:HG2	2.20	0.42
12:2Q:37:LEU:HD21	12:2Q:130:LYS:HE2	2.02	0.42
1:1A:74:G:H4'	24:12:55:ARG:NH1	2.34	0.42
30:18:26:LYS:HG2	30:18:46:ARG:O	2.20	0.42
1:1A:1810:U:OP2	63:1A:4360:HOH:O	2.22	0.42
1:1A:2874:G:OP1	15:1T:119:LYS:HD2	2.20	0.42
3:1D:38:LYS:HA	3:1D:38:LYS:HD2	1.82	0.42
6:1G:28:VAL:O	6:1G:31:VAL:HG13	2.20	0.42
15:1T:108:ARG:HG3	15:1T:109:GLU:N	2.34	0.42
1:2A:1169:G:C2	1:2A:1170:G:N7	2.88	0.42
1:2A:1495:A:H2'	1:2A:1496:A:H8	1.83	0.42
1:2A:1592:C:H2'	1:2A:1593:G:H8	1.85	0.42
1:2A:234:C:H2'	1:2A:235:U:C6	2.55	0.42
1:2A:2377:A:H2'	1:2A:2378:A:C8	2.55	0.42
1:2A:910:A:H2'	1:2A:911:A:C8	2.55	0.42
7:2H:13:LYS:HA	7:2H:14:GLY:HA2	1.79	0.42
11:2P:59:LEU:HD11	30:28:10:ALA:CB	2.45	0.42
1:1A:1120:G:N1	1:1A:1121:C:H1'	2.35	0.41
1:1A:1132:A:N3	1:1A:1132:A:H5''	2.35	0.41
1:1A:1821:C:H5''	1:1A:1822:A:OP1	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:253:C:O2'	1:1A:254:A:H2'	2.20	0.41
1:1A:7:G:H2'	1:1A:8:A:O4'	2.21	0.41
3:1D:132:PRO:HD3	3:1D:190:TYR:CZ	2.55	0.41
10:1O:106:LEU:HA	10:1O:106:LEU:HD23	1.90	0.41
1:2A:1545:A:H2'	1:2A:1546:C:O4'	2.20	0.41
1:2A:383:U:H2'	1:2A:385:C:H5	1.85	0.41
1:2A:68:G:H2'	1:2A:69:C:O4'	2.19	0.41
1:2A:956:G:N2	1:2A:959:A:H3'	2.35	0.41
1:2A:988:A:N7	63:2A:4041:HOH:O	2.37	0.41
4:2E:34:VAL:HG23	4:2E:66:HIS:HE2	1.85	0.41
10:2O:9:GLU:H	10:2O:9:GLU:HG2	1.67	0.41
15:2T:28:VAL:HG13	15:2T:86:ILE:HG23	2.01	0.41
1:1A:1320:A:N3	1:1A:1343:C:H1'	2.35	0.41
1:1A:1617:A:H2'	1:1A:1618:A:C8	2.56	0.41
1:1A:385:G:N1	1:1A:386:U:O4	2.53	0.41
7:1H:20:ALA:HB3	7:1H:23:ARG:HG3	2.02	0.41
23:21:50:ARG:HD2	23:21:57:GLU:OE2	2.18	0.41
27:25:49:CYS:SG	27:25:51:TYR:HB2	2.61	0.41
1:2A:1264:G:OP1	27:25:19:ARG:NH2	2.31	0.41
1:2A:1430:C:H2'	1:2A:1431:U:C6	2.55	0.41
1:2A:1866:C:H2'	1:2A:1876:A:O4'	2.20	0.41
1:2A:2137:C:N3	1:2A:2155:G:C6	2.88	0.41
1:2A:676:A:H1'	1:2A:2443:C:H1'	2.02	0.41
1:2A:2518:A:OP2	63:2A:3961:HOH:O	2.22	0.41
1:2A:315:G:H2'	1:2A:316:C:C6	2.55	0.41
1:2A:598:G:C6	1:2A:599:G:C5	3.08	0.41
1:2A:652(A):A:H2'	1:2A:652(A):A:N3	2.34	0.41
1:2A:684:G:OP1	29:27:16:HIS:ND1	2.52	0.41
1:2A:852:G:H2'	1:2A:853:G:C8	2.44	0.41
1:2A:2315:G:H1'	6:2G:126:ASP:OD2	2.19	0.41
8:2I:87:LYS:HE3	8:2I:87:LYS:HB2	1.73	0.41
1:2A:637:A:H2'	11:2P:117:GLU:OE1	2.19	0.41
4:2E:12:THR:HG21	15:2T:11:GLU:OE2	2.19	0.41
15:2T:27:THR:HB	15:2T:89:VAL:HG23	2.02	0.41
1:1A:1093:G:H2'	1:1A:1156:G:N2	2.35	0.41
1:1A:2081:A:O3'	5:1F:69:HIS:HA	2.21	0.41
1:1A:2155:G:N2	1:1A:2156:A:H62	2.19	0.41
1:1A:2205:C:H2'	1:1A:2206:G:H8	1.85	0.41
1:1A:2246:G:N7	63:1A:4443:HOH:O	2.37	0.41
1:1A:502:G:H4'	1:1A:527:A:N1	2.35	0.41
12:1Q:111:GLU:O	12:1Q:115:MET:HG2	2.20	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
15:1T:127:ALA:O	15:1T:128:GLU:HG2	2.20	0.41
18:1W:92:ARG:NH1	63:1W:303:HOH:O	2.52	0.41
21:1Z:99:TYR:HB3	21:1Z:123:ASP:OD2	2.19	0.41
29:27:12:ARG:HB3	29:27:46:VAL:HG21	2.03	0.41
1:2A:1480:G:C6	1:2A:1481:U:C4	3.08	0.41
1:2A:1550:C:H2'	1:2A:1551:C:C6	2.55	0.41
1:2A:2165:G:H8	1:2A:2165:G:O5'	2.03	0.41
1:2A:2321:G:N3	1:2A:2321:G:H2'	2.35	0.41
1:2A:2646:C:OP2	1:2A:2732:G:O2'	2.35	0.41
1:2A:2745:C:H2'	1:2A:2746:U:O4'	2.21	0.41
2:2B:43:C:O4'	6:2G:66:GLN:NE2	2.54	0.41
24:12:53:LEU:HA	24:12:53:LEU:HD23	1.80	0.41
1:1A:1400:A:H2'	1:1A:1401:G:O4'	2.20	0.41
1:1A:2250:G:H2'	1:1A:2250:G:N3	2.34	0.41
1:1A:908:A:C2	1:1A:963:A:C4	3.09	0.41
4:1E:143:ASN:HD22	4:1E:147:PRO:HD3	1.85	0.41
18:1W:7:ALA:HB2	18:1W:50:VAL:HG22	2.01	0.41
1:2A:1589:C:H2'	1:2A:1590:U:H6	1.85	0.41
1:2A:1630:G:H2'	1:2A:1631:C:C6	2.56	0.41
1:2A:1652:A:C2'	1:2A:1653:G:H5'	2.50	0.41
1:2A:1697:G:OP2	1:2A:1698:A:O2'	2.27	0.41
1:2A:1131:G:O6	1:2A:2040:C:H1'	2.20	0.41
1:2A:2103:C:O2'	1:2A:2104:G:H5'	2.20	0.41
1:2A:2802:G:H2'	1:2A:2803:C:O4'	2.20	0.41
1:2A:839:U:H2'	1:2A:840:C:H6	1.85	0.41
2:2B:11:C:OP2	2:2B:12:C:N4	2.41	0.41
2:2B:80:U:H2'	2:2B:81:G:C8	2.55	0.41
2:2B:8:U:O3'	14:2S:25:ARG:NH2	2.53	0.41
4:2E:109:LYS:O	4:2E:111:ARG:NH1	2.53	0.41
5:2F:132:VAL:HG21	5:2F:163:VAL:HG22	2.03	0.41
5:2F:133:ASN:O	5:2F:162:LEU:HD23	2.19	0.41
8:2I:59:ALA:HA	8:2I:62:LYS:HB3	2.00	0.41
11:2P:77:ARG:HG3	11:2P:78:PRO:HD2	2.02	0.41
1:1A:1108:G:C5	1:1A:1134:A:H2'	2.56	0.41
1:1A:943:C:C2	1:1A:944:C:C4	3.08	0.41
4:1E:47:VAL:HG22	4:1E:84:PHE:O	2.21	0.41
10:1O:98:VAL:HG22	10:1O:118:ALA:HA	2.03	0.41
11:1P:135:LEU:HA	11:1P:135:LEU:HD23	1.77	0.41
13:1R:12:ARG:HG2	13:1R:16:HIS:CE1	2.55	0.41
14:1S:65:VAL:O	14:1S:69:VAL:HG12	2.20	0.41
18:1W:23:LEU:HD12	18:1W:23:LEU:HA	1.93	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
23:21:8:SER:HB3	23:21:66:HIS:CD2	2.56	0.41
1:2A:2284:C:OP2	28:26:2:ALA:N	2.54	0.41
1:2A:1372:U:H2'	1:2A:1373:A:O4'	2.21	0.41
1:2A:1473:G:C6	1:2A:1474:C:C4	3.09	0.41
1:2A:407:G:H2'	1:2A:408:G:H8	1.85	0.41
1:2A:536:A:H2'	1:2A:537:C:C6	2.55	0.41
1:2A:569:U:C4	1:2A:570:G:C6	3.09	0.41
1:2A:93:G:H2'	1:2A:94:C:C6	2.54	0.41
2:2B:115:G:H2'	2:2B:116:G:O4'	2.20	0.41
2:2B:90:A:N7	2:2B:91:C:H1'	2.36	0.41
5:2F:154:VAL:HG22	5:2F:191:ARG:HB2	2.02	0.41
7:2H:3:ARG:HG2	7:2H:6:ARG:HE	1.84	0.41
7:2H:7:LEU:HD23	7:2H:69:ARG:NH2	2.35	0.41
12:2Q:17:LEU:HB3	12:2Q:39:PRO:HB2	2.02	0.41
17:2V:31:ALA:O	17:2V:61:VAL:HG12	2.19	0.41
24:12:32:LEU:HD11	24:12:54:LYS:HG2	2.02	0.41
1:1A:1162:C:H5'	1:1A:1163:G:OP2	2.20	0.41
1:1A:173:C:H2'	1:1A:174:U:H6	1.84	0.41
1:1A:2177:G:H3'	1:1A:2178:G:H8	1.86	0.41
1:1A:233:A:C2	1:1A:244:A:C4	3.09	0.41
1:1A:645:G:H2'	1:1A:645:G:N3	2.35	0.41
1:1A:670:C:O5'	1:1A:670:C:H6	2.03	0.41
2:1B:88:C:H2'	2:1B:89:G:O4'	2.20	0.41
6:1G:72:ARG:HH12	6:1G:87:PRO:HG3	1.85	0.41
1:1A:873:U:H4'	11:1P:55:ARG:HB3	2.03	0.41
20:1Y:35:TYR:CE2	20:1Y:69:ALA:HB3	2.56	0.41
1:2A:1773:A:H5''	63:2A:4614:HOH:O	2.19	0.41
1:2A:2095:C:H2'	1:2A:2096:U:O4'	2.20	0.41
1:2A:2792:G:N3	1:2A:2792:G:H2'	2.36	0.41
1:2A:460:A:C2	1:2A:470:A:C4	3.09	0.41
1:2A:855:G:C6	1:2A:856:C:C4	3.09	0.41
1:2A:764:A:H5'	3:2D:210:GLY:HA2	2.03	0.41
31:19:27:CYS:SG	31:19:28:GLU:N	2.94	0.41
1:1A:1053:C:OP1	9:1N:37:LYS:NZ	2.54	0.41
1:1A:1255:A:H5''	1:1A:1257:G:O4'	2.20	0.41
1:1A:2096:U:H2'	1:1A:2097:U:C6	2.55	0.41
1:1A:2621:U:H5'	58:A:5:ASN:HD21	1.85	0.41
1:1A:327:U:H2'	1:1A:328:G:H8	1.86	0.41
1:1A:397:G:H4'	1:1A:398:A:OP2	2.20	0.41
1:1A:505:A:N3	1:1A:507:G:H5''	2.35	0.41
1:1A:929:G:H1	1:1A:940:C:N4	2.03	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
2:1B:1:U:HO2'	2:1B:2:C:P	2.43	0.41
20:1Y:6:HIS:CD2	20:1Y:6:HIS:H	2.37	0.41
1:2A:11:G:C2'	1:2A:12:U:H5'	2.51	0.41
1:2A:1702:G:H2'	1:2A:1703:G:O4'	2.21	0.41
1:2A:2130:U:O2'	1:2A:2133:G:H4'	2.21	0.41
1:2A:2197:U:H1'	1:2A:2198:A:C8	2.56	0.41
1:2A:2364:C:H2'	1:2A:2365:G:O4'	2.21	0.41
1:2A:531:C:H4'	1:2A:532:A:H5''	2.03	0.41
1:2A:848:G:N9	1:2A:933:A:H8	2.19	0.41
3:2D:218:ARG:HB3	3:2D:219:PRO:HD2	2.03	0.41
1:2A:784:A:N6	3:2D:229:VAL:HG11	2.35	0.41
26:14:18:CYS:SG	26:14:20:ASN:HB2	2.61	0.41
1:1A:1014:U:O5'	1:1A:1014:U:H6	2.04	0.41
1:1A:2128:G:H2'	1:1A:2129:C:C6	2.55	0.41
1:1A:580:U:H2'	1:1A:581:G:O4'	2.64	0.41
1:1A:670:C:H5''	1:1A:671:A:OP2	2.20	0.41
1:1A:847:A:OP1	1:1A:847:A:H8	2.03	0.41
18:1W:25:ARG:NH2	18:1W:74:ALA:O	2.48	0.41
30:28:23:VAL:HG13	30:28:47:LYS:HB3	2.02	0.41
1:2A:127:A:H5''	1:2A:128:C:C6	2.55	0.41
1:2A:1539:G:H2'	1:2A:1540:U:O4'	2.20	0.41
1:2A:1557:C:H5''	1:2A:1558:A:OP2	2.21	0.41
1:2A:2678:C:H2'	1:2A:2679:A:O4'	2.21	0.41
5:2F:129:PHE:CD2	5:2F:163:VAL:HG21	2.56	0.41
10:2O:24:VAL:HG12	10:2O:33:ALA:HB2	2.03	0.41
1:1A:2153:G:H5''	1:1A:2154:U:H3'	2.02	0.41
1:1A:2724:U:H1'	1:1A:2725:A:C8	2.56	0.41
1:1A:492:A:N3	1:1A:730:C:H1'	2.35	0.41
1:1A:63:A:C5	19:1X:66:LEU:HD13	2.56	0.41
2:1B:43:C:H4'	6:1G:98:ARG:HH21	1.86	0.41
7:1H:154:PRO:HB3	7:1H:163:TYR:CE1	2.56	0.41
9:1N:120:LEU:HG	9:1N:122:VAL:HG23	2.01	0.41
12:1Q:17:LEU:HB3	12:1Q:39:PRO:HB2	2.02	0.41
28:26:9:LEU:HD13	28:26:51:GLU:HB2	2.02	0.41
1:2A:1653:G:H3'	13:2R:2:ARG:HD3	2.02	0.41
1:2A:218:A:C2	1:2A:235:U:H4'	2.56	0.41
1:2A:2552:2MU:H2'	1:2A:2554:U:OP2	2.21	0.41
1:2A:2689:U:H4'	1:2A:2690:C:H5'	2.03	0.41
1:2A:2880:C:O3'	13:2R:90:ARG:NH1	2.53	0.41
3:2D:4:LYS:HB3	3:2D:18:VAL:HG23	2.02	0.41
4:2E:97:LYS:HB3	4:2E:97:LYS:HE2	1.80	0.41

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
6:2G:138:GLN:OE1	6:2G:138:GLN:N	2.49	0.41
1:1A:1899:A:H5''	1:1A:1900:G:OP2	2.21	0.41
1:1A:2171:G:N1	1:1A:2172:U:O2	2.54	0.41
1:1A:2364:A:N6	1:1A:2377:G:O2'	2.53	0.41
3:1D:26:LYS:HD3	3:1D:83:GLU:OE2	2.21	0.41
3:1D:35:LYS:HB2	3:1D:36:PRO:HD2	2.02	0.41
5:1F:135:LYS:HB2	5:1F:138:GLU:HG3	2.03	0.41
26:24:47:GLN:C	26:24:49:PHE:H	2.24	0.41
1:2A:2489:G:C2'	1:2A:2490:G:H5'	2.51	0.41
1:2A:2626:C:H2'	1:2A:2627:G:O4'	2.21	0.41
1:2A:322:A:H5'	1:2A:340:A:H1'	2.03	0.41
1:2A:27:G:C2	1:2A:512:G:N3	2.89	0.41
3:2D:232:PRO:HB3	3:2D:244:ARG:CZ	2.51	0.41
7:2H:170:ARG:O	7:2H:171:LEU:HD23	2.20	0.41
7:2H:8:PRO:O	7:2H:10:PRO:HD3	2.20	0.41
8:2I:134:PRO:C	8:2I:136:VAL:H	2.24	0.41
14:2S:26:LEU:HD22	14:2S:87:PHE:HD1	1.86	0.41
15:2T:60:THR:HG22	15:2T:77:PRO:HA	2.02	0.41
18:2W:78:GLU:OE2	18:2W:99:ARG:HD3	2.21	0.41
1:1A:1640:G:H2'	1:1A:1641:G:O4'	2.21	0.41
1:1A:662:A:H2'	11:1P:117:GLU:OE1	2.21	0.41
1:1A:890:G:O2'	1:1A:906:G:O6	46.32	0.41
4:1E:2:LYS:HG3	4:1E:200:GLU:HB2	2.02	0.41
6:1G:41:GLN:HG2	6:1G:154:GLY:O	2.21	0.41
24:22:1:MET:SD	24:22:56:GLN:NE2	2.94	0.41
1:2A:2274:A:C5	1:2A:2276:G:C8	3.09	0.41
1:2A:2667:C:H2'	1:2A:2668:G:O4'	2.21	0.41
1:2A:866:A:C6	1:2A:914:C:C5	3.09	0.41
4:2E:11:MET:HG2	4:2E:24:THR:HB	2.03	0.41
5:2F:156:LEU:HD21	5:2F:163:VAL:HG12	2.02	0.41
5:2F:23:ASP:O	5:2F:24:LEU:HD12	2.21	0.41
12:2Q:137:TYR:CE1	21:2Z:83:PRO:HG3	2.56	0.41
23:11:8:SER:HB3	23:11:66:HIS:CD2	2.57	0.40
1:1A:1102:G:H5''	1:1A:1103:A:O4'	2.21	0.40
1:1A:1355:G:H2'	1:1A:1356:G:C8	3.71	0.40
1:1A:1478:C:H2'	1:1A:1479:U:O4'	2.22	0.40
1:1A:2177:G:N3	1:1A:2177:G:H2'	2.37	0.40
1:1A:346:A:H5'	1:1A:364:A:H1'	2.01	0.40
4:1E:98:PRO:HD3	4:1E:175:VAL:HG13	2.03	0.40
7:1H:101:ARG:NH1	7:1H:117:PRO:HG2	2.36	0.40
7:1H:87:LEU:HD23	7:1H:164:TYR:HA	2.02	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
12:1Q:30:GLY:HA2	12:1Q:107:ALA:HB2	2.03	0.40
13:1R:12:ARG:HG2	13:1R:16:HIS:ND1	2.36	0.40
16:1U:16:LYS:HE2	16:1U:16:LYS:HB3	1.77	0.40
20:1Y:13:VAL:HG12	20:1Y:74:PRO:HA	2.03	0.40
1:2A:1777:U:H2'	1:2A:1778:U:H6	1.84	0.40
1:2A:1946:U:H2'	1:2A:1947:C:C6	2.56	0.40
1:2A:2126:A:N3	1:2A:2127:G:H1'	2.36	0.40
1:2A:2396:G:OP1	23:21:25:LYS:NZ	2.39	0.40
1:2A:2461:C:H2'	1:2A:2462:U:C6	2.56	0.40
1:2A:656:G:H2'	1:2A:657:U:O4'	2.21	0.40
1:2A:856:C:HO2'	1:2A:857:C:P	2.44	0.40
3:2D:39:LYS:HB3	3:2D:39:LYS:HE2	1.86	0.40
3:2D:68:LYS:HB2	3:2D:70:TRP:CE2	2.56	0.40
4:2E:170:LEU:HB3	4:2E:184:VAL:CG2	2.51	0.40
11:2P:88:LEU:HD11	11:2P:114:ILE:HD12	2.03	0.40
15:2T:94:ALA:HB1	15:2T:99:LEU:HD21	2.03	0.40
21:2Z:75:ASN:HB2	21:2Z:85:HIS:HB3	2.03	0.40
25:13:23:LEU:HD13	25:13:50:VAL:HG11	2.03	0.40
1:1A:1992:A:H4'	1:1A:1993:A:OP1	2.21	0.40
1:1A:1312:G:O2'	1:1A:2034:G:O6	2.21	0.40
1:1A:2119:C:H2'	1:1A:2120:U:O4'	2.21	0.40
1:1A:2283:G:OP1	22:10:18:ALA:HB1	2.21	0.40
1:1A:2864:G:H2'	1:1A:2865:C:C6	2.56	0.40
1:1A:327:U:H2'	1:1A:328:G:C8	2.56	0.40
18:1W:65:LEU:HA	18:1W:65:LEU:HD23	1.85	0.40
25:23:12:PRO:HB2	25:23:20:LYS:HG2	2.02	0.40
1:2A:1198:U:H2'	1:2A:1199:U:C6	2.57	0.40
1:2A:1434:A:H2'	1:2A:1435:G:O4'	2.31	0.40
1:2A:1754:C:H2'	1:2A:1755:A:O4'	2.21	0.40
1:2A:196:A:N3	1:2A:196:A:H2'	2.36	0.40
1:2A:265:A:H1'	1:2A:266:G:O4'	2.22	0.40
1:2A:573:G:O2'	1:2A:574:C:H3'	2.21	0.40
4:2E:127:ASP:OD2	63:2E:401:HOH:O	2.22	0.40
1:2A:2658:C:P	7:2H:160:LYS:HZ1	2.43	0.40
20:2Y:8:LYS:HD3	20:2Y:97:ARG:HH11	1.86	0.40
8:1I:27:ARG:HD3	23:11:71:TYR:CE2	2.56	0.40
25:13:59:VAL:O	25:13:60:GLU:HG3	2.21	0.40
1:1A:2203:G:O2'	1:1A:2204:G:OP1	2.33	0.40
1:1A:2331:G:N1	14:1S:3:ARG:HA	2.36	0.40
1:1A:297:C:H2'	1:1A:298:G:C8	2.53	0.40
1:1A:312:C:H2'	1:1A:313:A:C8	2.56	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:440:C:OP2	63:1A:4362:HOH:O	2.22	0.40
24:22:53:LEU:HA	24:22:53:LEU:HD23	1.95	0.40
26:24:68:ARG:HD3	26:24:69:LYS:H	1.86	0.40
30:28:34:TRP:CG	30:28:35:GLN:N	2.90	0.40
1:2A:2820:A:O2'	1:2A:2821:A:OP1	2.34	0.40
1:2A:517:C:OP1	27:25:16:ARG:NH2	2.54	0.40
1:2A:56:A:H2'	1:2A:57:C:O4'	2.21	0.40
1:2A:94(A):G:H2'	1:2A:95:G:O4'	2.22	0.40
1:1A:1154:U:H2'	1:1A:1155:C:O4'	2.22	0.40
1:1A:1410:G:OP1	23:11:2:SER:HA	2.21	0.40
1:1A:2769:U:H1'	1:1A:2770:A:H5''	2.04	0.40
1:1A:831:A:C6	3:1D:229:VAL:HG11	2.56	0.40
5:1F:123:LEU:HD13	5:1F:192:LEU:HD13	2.04	0.40
13:1R:9:LYS:O	13:1R:17:ARG:HD3	2.21	0.40
21:1Z:72:ARG:HD3	21:1Z:72:ARG:HA	1.95	0.40
29:27:26:GLY:O	29:27:30:VAL:HG23	2.21	0.40
1:2A:1221(A):C:C2	1:2A:1229:G:C2	3.10	0.40
1:2A:1507:A:O2'	1:2A:1508:A:H8	2.03	0.40
1:2A:1547:C:H2'	1:2A:1548:C:H6	1.85	0.40
1:2A:262:A:H2'	1:2A:263:C:O4'	2.21	0.40
1:2A:271(D):G:H2'	1:2A:271(E):U:C6	2.57	0.40
1:2A:463:G:N2	1:2A:466:A:OP2	2.50	0.40
1:2A:647:G:H8	1:2A:647:G:O5'	2.03	0.40
1:2A:900:A:H2'	1:2A:901:A:H8	1.84	0.40
3:2D:155:LEU:HA	3:2D:155:LEU:HD23	4.52	0.40
3:2D:275:LYS:HA	3:2D:276:LYS:C	2.42	0.40
7:2H:126:PRO:HB2	7:2H:127:GLU:H	1.65	0.40
1:2A:864:G:OP2	12:2Q:22:LYS:HE3	2.21	0.40
14:2S:8:GLU:HG2	14:2S:8:GLU:H	1.65	0.40
16:2U:76:TYR:CZ	16:2U:80:ILE:HG13	2.56	0.40
1:1A:1100:A:N6	1:1A:1101:G:C6	2.89	0.40
1:1A:1756:U:H2'	1:1A:1757:C:C6	2.56	0.40
1:1A:213:G:H2'	1:1A:214:A:O4'	2.21	0.40
1:1A:2158:C:N4	1:1A:2177:G:C6	2.84	0.40
1:1A:2021:C:H4'	1:1A:2736:C:O2	2.22	0.40
8:1I:6:LEU:HD11	8:1I:37:VAL:HG23	2.04	0.40
1:2A:1232:G:H2'	1:2A:1233:C:C6	2.56	0.40
1:2A:1720:U:H2'	1:2A:1721:G:O4'	2.21	0.40
1:2A:2270:G:H2'	1:2A:2271:G:O4'	2.20	0.40
1:2A:2375:G:N2	1:2A:2378:A:OP2	2.49	0.40
1:2A:253:C:O2'	63:2A:3957:HOH:O	2.21	0.40

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Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:336:C:H2'	1:2A:337:C:H6	2.26	0.40
1:2A:473:G:C2	1:2A:474:G:C5	3.81	0.40
2:2B:88:C:H2'	2:2B:89:G:O4'	2.21	0.40
1:2A:2788:C:H5''	4:2E:61:ARG:HH21	1.87	0.40
5:2F:126:VAL:HG21	5:2F:129:PHE:CZ	2.57	0.40
14:2S:3:ARG:HD3	14:2S:4:LEU:N	2.36	0.40
15:2T:13:ARG:HB2	15:2T:14:TYR:CD2	2.57	0.40
15:2T:74:ARG:HG2	15:2T:76:PHE:CZ	2.57	0.40
17:2V:18:LEU:HD12	17:2V:19:LYS:N	2.37	0.40

There are no symmetry-related clashes.

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	
3	1D	273/276 (99%)	261 (96%)	12 (4%)	0	100	100
3	2D	273/276 (99%)	259 (95%)	14 (5%)	0	100	100
4	1E	202/206 (98%)	194 (96%)	7 (4%)	1 (0%)	32	60
4	2E	202/206 (98%)	193 (96%)	8 (4%)	1 (0%)	32	60
5	1F	201/210 (96%)	195 (97%)	5 (2%)	1 (0%)	32	60
5	2F	201/210 (96%)	193 (96%)	7 (4%)	1 (0%)	32	60
6	1G	179/182 (98%)	168 (94%)	8 (4%)	3 (2%)	11	27
6	2G	179/182 (98%)	162 (90%)	13 (7%)	4 (2%)	8	20
7	1H	172/180 (96%)	161 (94%)	10 (6%)	1 (1%)	28	56
7	2H	172/180 (96%)	156 (91%)	13 (8%)	3 (2%)	11	27
8	1I	144/148 (97%)	130 (90%)	13 (9%)	1 (1%)	25	53
8	2I	144/148 (97%)	129 (90%)	13 (9%)	2 (1%)	13	33
9	1N	138/140 (99%)	133 (96%)	5 (4%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
9	2N	138/140 (99%)	132 (96%)	6 (4%)	0	100	100
10	1O	120/122 (98%)	115 (96%)	5 (4%)	0	100	100
10	2O	120/122 (98%)	112 (93%)	6 (5%)	2 (2%)	11	27
11	1P	147/150 (98%)	134 (91%)	12 (8%)	1 (1%)	25	53
11	2P	147/150 (98%)	137 (93%)	9 (6%)	1 (1%)	25	53
12	1Q	139/141 (99%)	130 (94%)	9 (6%)	0	100	100
12	2Q	139/141 (99%)	130 (94%)	6 (4%)	3 (2%)	8	20
13	1R	116/118 (98%)	109 (94%)	7 (6%)	0	100	100
13	2R	116/118 (98%)	108 (93%)	8 (7%)	0	100	100
14	1S	108/112 (96%)	102 (94%)	6 (6%)	0	100	100
14	2S	108/112 (96%)	102 (94%)	5 (5%)	1 (1%)	20	46
15	1T	129/146 (88%)	123 (95%)	5 (4%)	1 (1%)	22	49
15	2T	129/146 (88%)	124 (96%)	5 (4%)	0	100	100
16	1U	114/118 (97%)	114 (100%)	0	0	100	100
16	2U	114/118 (97%)	111 (97%)	3 (3%)	0	100	100
17	1V	99/101 (98%)	96 (97%)	2 (2%)	1 (1%)	18	43
17	2V	99/101 (98%)	92 (93%)	6 (6%)	1 (1%)	18	43
18	1W	110/113 (97%)	109 (99%)	1 (1%)	0	100	100
18	2W	110/113 (97%)	110 (100%)	0	0	100	100
19	1X	93/96 (97%)	91 (98%)	1 (1%)	1 (1%)	17	40
19	2X	93/96 (97%)	89 (96%)	4 (4%)	0	100	100
20	1Y	105/110 (96%)	97 (92%)	7 (7%)	1 (1%)	18	43
20	2Y	105/110 (96%)	100 (95%)	4 (4%)	1 (1%)	18	43
21	1Z	148/206 (72%)	134 (90%)	12 (8%)	2 (1%)	13	33
21	2Z	156/206 (76%)	138 (88%)	15 (10%)	3 (2%)	9	23
22	10	81/85 (95%)	79 (98%)	2 (2%)	0	100	100
22	20	81/85 (95%)	76 (94%)	5 (6%)	0	100	100
23	11	95/98 (97%)	92 (97%)	2 (2%)	1 (1%)	17	40
23	21	95/98 (97%)	94 (99%)	1 (1%)	0	100	100
24	12	68/72 (94%)	67 (98%)	1 (2%)	0	100	100
24	22	68/72 (94%)	66 (97%)	2 (3%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
25	13	57/60 (95%)	57 (100%)	0	0	100	100
25	23	57/60 (95%)	54 (95%)	3 (5%)	0	100	100
26	14	67/71 (94%)	53 (79%)	9 (13%)	5 (8%)	1	1
26	24	67/71 (94%)	53 (79%)	13 (19%)	1 (2%)	12	30
27	15	57/60 (95%)	55 (96%)	2 (4%)	0	100	100
27	25	57/60 (95%)	56 (98%)	1 (2%)	0	100	100
28	16	51/54 (94%)	49 (96%)	2 (4%)	0	100	100
28	26	51/54 (94%)	50 (98%)	1 (2%)	0	100	100
29	17	46/49 (94%)	46 (100%)	0	0	100	100
29	27	46/49 (94%)	45 (98%)	1 (2%)	0	100	100
30	18	62/65 (95%)	62 (100%)	0	0	100	100
30	28	62/65 (95%)	59 (95%)	3 (5%)	0	100	100
31	19	35/37 (95%)	35 (100%)	0	0	100	100
31	29	35/37 (95%)	34 (97%)	1 (3%)	0	100	100
33	1b	229/256 (90%)	194 (85%)	26 (11%)	9 (4%)	3	8
33	2b	229/256 (90%)	201 (88%)	21 (9%)	7 (3%)	5	11
34	1c	204/239 (85%)	188 (92%)	13 (6%)	3 (2%)	12	30
34	2c	204/239 (85%)	178 (87%)	24 (12%)	2 (1%)	18	43
35	1d	206/209 (99%)	193 (94%)	11 (5%)	2 (1%)	18	43
35	2d	206/209 (99%)	189 (92%)	14 (7%)	3 (2%)	12	30
36	1e	146/162 (90%)	135 (92%)	8 (6%)	3 (2%)	8	21
36	2e	146/162 (90%)	134 (92%)	10 (7%)	2 (1%)	13	33
37	1f	98/101 (97%)	94 (96%)	3 (3%)	1 (1%)	18	43
37	2f	98/101 (97%)	94 (96%)	3 (3%)	1 (1%)	18	43
38	1g	153/156 (98%)	142 (93%)	10 (6%)	1 (1%)	25	53
38	2g	153/156 (98%)	140 (92%)	9 (6%)	4 (3%)	6	15
39	1h	135/138 (98%)	130 (96%)	5 (4%)	0	100	100
39	2h	135/138 (98%)	118 (87%)	17 (13%)	0	100	100
40	1i	125/128 (98%)	109 (87%)	14 (11%)	2 (2%)	11	28
40	2i	125/128 (98%)	110 (88%)	15 (12%)	0	100	100
41	1j	95/105 (90%)	84 (88%)	6 (6%)	5 (5%)	2	4

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
41	2j	94/105 (90%)	79 (84%)	10 (11%)	5 (5%)	2	4
42	1k	112/129 (87%)	105 (94%)	6 (5%)	1 (1%)	20	46
42	2k	112/129 (87%)	105 (94%)	5 (4%)	2 (2%)	10	25
43	1l	119/132 (90%)	114 (96%)	5 (4%)	0	100	100
43	2l	119/132 (90%)	110 (92%)	9 (8%)	0	100	100
44	1m	121/126 (96%)	111 (92%)	9 (7%)	1 (1%)	22	49
44	2m	120/126 (95%)	108 (90%)	12 (10%)	0	100	100
45	1n	58/61 (95%)	56 (97%)	2 (3%)	0	100	100
45	2n	58/61 (95%)	51 (88%)	5 (9%)	2 (3%)	4	10
46	1o	86/89 (97%)	83 (96%)	2 (2%)	1 (1%)	15	37
46	2o	86/89 (97%)	83 (96%)	3 (4%)	0	100	100
47	1p	80/88 (91%)	72 (90%)	7 (9%)	1 (1%)	14	35
47	2p	80/88 (91%)	70 (88%)	9 (11%)	1 (1%)	14	35
48	1q	97/105 (92%)	91 (94%)	6 (6%)	0	100	100
48	2q	97/105 (92%)	90 (93%)	7 (7%)	0	100	100
49	1r	66/88 (75%)	63 (96%)	3 (4%)	0	100	100
49	2r	66/88 (75%)	63 (96%)	2 (3%)	1 (2%)	12	30
50	1s	81/93 (87%)	72 (89%)	8 (10%)	1 (1%)	15	37
50	2s	81/93 (87%)	68 (84%)	10 (12%)	3 (4%)	4	8
51	1t	94/106 (89%)	85 (90%)	5 (5%)	4 (4%)	3	6
51	2t	94/106 (89%)	85 (90%)	4 (4%)	5 (5%)	2	4
52	1u	21/27 (78%)	19 (90%)	2 (10%)	0	100	100
52	2u	21/27 (78%)	15 (71%)	6 (29%)	0	100	100
58	A	12/18 (67%)	5 (42%)	5 (42%)	2 (17%)	0	0
58	B	12/18 (67%)	6 (50%)	4 (33%)	2 (17%)	0	0
All	All	11394/12164 (94%)	10597 (93%)	676 (6%)	121 (1%)	17	40

All (121) Ramachandran outliers are listed below:

Mol	Chain	Res	Type
6	1G	47	LYS
7	1H	126	PRO
8	1I	10	GLU

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Mol	Chain	Res	Type
20	1Y	55	TYR
23	1l	3	LYS
33	1b	10	LEU
33	1b	17	PHE
40	1i	54	ASP
41	1j	79	ARG
51	1t	100	ILE
5	2F	130	ALA
6	2G	47	LYS
7	2H	47	GLU
7	2H	126	PRO
20	2Y	55	TYR
38	2g	6	ARG
50	2s	81	ARG
51	2t	10	LEU
51	2t	47	GLY
58	A	12	SER
5	1F	130	ALA
15	1T	37	GLY
19	1X	93	GLU
21	1Z	156	LYS
33	1b	22	LYS
33	1b	126	GLU
36	1e	85	GLY
41	1j	55	LYS
44	1m	67	GLU
51	1t	47	GLY
6	2G	49	ASP
6	2G	124	SER
8	2I	10	GLU
10	2O	5	GLN
12	2Q	17	LEU
17	2V	79	VAL
21	2Z	172	ALA
33	2b	17	PHE
34	2c	95	THR
41	2j	29	ARG
41	2j	75	ILE
42	2k	49	GLY
49	2r	36	ASN
51	2t	100	ILE
58	A	16	ASN

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Mol	Chain	Res	Type
58	B	12	SER
58	B	16	ASN
26	14	18	CYS
33	1b	8	LYS
37	1f	40	VAL
41	1j	77	PRO
46	1o	19	PRO
51	1t	10	LEU
7	2H	12	PRO
8	2I	85	GLU
14	2S	96	GLY
21	2Z	142	SER
33	2b	8	LYS
33	2b	20	GLU
35	2d	178	VAL
35	2d	179	GLU
38	2g	4	ARG
41	2j	78	ASN
50	2s	27	GLU
51	2t	99	LEU
4	1E	52	LEU
6	1G	49	ASP
26	14	55	ARG
26	14	57	GLU
26	14	64	GLY
34	1c	65	ALA
36	1e	86	ALA
38	1g	80	VAL
41	1j	29	ARG
41	1j	78	ASN
50	1s	27	GLU
4	2E	52	LEU
33	2b	16	HIS
33	2b	74	LYS
33	2b	125	PRO
36	2e	85	GLY
38	2g	80	VAL
45	2n	19	ARG
50	2s	9	VAL
6	1G	43	LEU
11	1P	29	LYS
17	1V	79	VAL

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Mol	Chain	Res	Type
21	1Z	147	GLY
33	1b	9	GLU
33	1b	16	HIS
33	1b	231	GLU
35	1d	178	VAL
35	1d	179	GLU
10	2O	29	ASN
12	2Q	16	ARG
26	24	64	GLY
36	2e	112	LEU
41	2j	79	ARG
6	2G	52	ILE
11	2P	122	PRO
21	2Z	146	ILE
34	2c	66	VAL
45	2n	60	SER
34	1c	66	VAL
42	1k	105	VAL
47	1p	53	VAL
12	2Q	15	GLY
33	2b	127	ILE
42	2k	105	VAL
35	2d	171	GLY
51	2t	102	GLY
51	1t	102	GLY
38	2g	17	VAL
41	2j	91	PRO
47	2p	53	VAL
26	14	29	PRO
33	1b	234	PRO
36	1e	69	VAL
40	1i	44	VAL
37	2f	40	VAL
34	1c	81	GLY

5.3.2 Protein sidechains ⓘ

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
3	1D	215/218 (99%)	197 (92%)	18 (8%)	13	29
3	2D	215/218 (99%)	203 (94%)	12 (6%)	25	51
4	1E	164/166 (99%)	149 (91%)	15 (9%)	11	25
4	2E	164/166 (99%)	153 (93%)	11 (7%)	19	42
5	1F	160/166 (96%)	147 (92%)	13 (8%)	14	31
5	2F	159/166 (96%)	148 (93%)	11 (7%)	18	41
6	1G	143/156 (92%)	135 (94%)	8 (6%)	25	51
6	2G	143/156 (92%)	137 (96%)	6 (4%)	34	65
7	1H	144/148 (97%)	138 (96%)	6 (4%)	34	65
7	2H	144/148 (97%)	139 (96%)	5 (4%)	41	72
8	1I	113/124 (91%)	103 (91%)	10 (9%)	12	27
8	2I	105/124 (85%)	95 (90%)	10 (10%)	10	23
9	1N	118/119 (99%)	109 (92%)	9 (8%)	15	35
9	2N	118/119 (99%)	111 (94%)	7 (6%)	23	49
10	1O	100/100 (100%)	96 (96%)	4 (4%)	36	67
10	2O	100/100 (100%)	96 (96%)	4 (4%)	36	67
11	1P	115/116 (99%)	109 (95%)	6 (5%)	27	55
11	2P	115/116 (99%)	112 (97%)	3 (3%)	51	81
12	1Q	111/111 (100%)	104 (94%)	7 (6%)	21	46
12	2Q	111/111 (100%)	103 (93%)	8 (7%)	17	39
13	1R	101/101 (100%)	89 (88%)	12 (12%)	6	14
13	2R	101/101 (100%)	90 (89%)	11 (11%)	7	17
14	1S	86/88 (98%)	78 (91%)	8 (9%)	10	24
14	2S	85/88 (97%)	79 (93%)	6 (7%)	17	39
15	1T	115/127 (91%)	111 (96%)	4 (4%)	41	72
15	2T	113/127 (89%)	109 (96%)	4 (4%)	41	72
16	1U	93/94 (99%)	86 (92%)	7 (8%)	16	36
16	2U	93/94 (99%)	90 (97%)	3 (3%)	44	75
17	1V	80/82 (98%)	70 (88%)	10 (12%)	5	12
17	2V	80/82 (98%)	73 (91%)	7 (9%)	12	27
18	1W	90/92 (98%)	83 (92%)	7 (8%)	15	33
18	2W	90/92 (98%)	85 (94%)	5 (6%)	25	51

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
19	1X	77/78 (99%)	74 (96%)	3 (4%)	37	68
19	2X	77/78 (99%)	77 (100%)	0	100	100
20	1Y	85/91 (93%)	78 (92%)	7 (8%)	13	30
20	2Y	85/91 (93%)	83 (98%)	2 (2%)	54	83
21	1Z	135/179 (75%)	124 (92%)	11 (8%)	14	31
21	2Z	137/179 (76%)	126 (92%)	11 (8%)	14	32
22	10	65/67 (97%)	61 (94%)	4 (6%)	21	46
22	20	65/67 (97%)	60 (92%)	5 (8%)	15	34
23	11	80/83 (96%)	79 (99%)	1 (1%)	73	91
23	21	80/83 (96%)	78 (98%)	2 (2%)	53	82
24	12	65/67 (97%)	63 (97%)	2 (3%)	45	75
24	22	65/67 (97%)	62 (95%)	3 (5%)	31	61
25	13	51/52 (98%)	49 (96%)	2 (4%)	37	68
25	23	50/52 (96%)	48 (96%)	2 (4%)	36	67
26	14	59/63 (94%)	55 (93%)	4 (7%)	18	41
26	24	53/63 (84%)	51 (96%)	2 (4%)	38	68
27	15	50/52 (96%)	46 (92%)	4 (8%)	14	32
27	25	50/52 (96%)	46 (92%)	4 (8%)	14	32
28	16	51/52 (98%)	45 (88%)	6 (12%)	6	14
28	26	50/52 (96%)	47 (94%)	3 (6%)	22	48
29	17	41/42 (98%)	36 (88%)	5 (12%)	6	13
29	27	41/42 (98%)	37 (90%)	4 (10%)	9	21
30	18	54/55 (98%)	47 (87%)	7 (13%)	5	11
30	28	54/55 (98%)	50 (93%)	4 (7%)	16	37
31	19	34/34 (100%)	34 (100%)	0	100	100
31	29	34/34 (100%)	33 (97%)	1 (3%)	48	77
33	1b	192/220 (87%)	184 (96%)	8 (4%)	34	65
33	2b	187/220 (85%)	175 (94%)	12 (6%)	20	45
34	1c	142/188 (76%)	137 (96%)	5 (4%)	41	72
34	2c	140/188 (74%)	138 (99%)	2 (1%)	71	90
35	1d	169/181 (93%)	160 (95%)	9 (5%)	26	54

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
35	2d	173/181 (96%)	161 (93%)	12 (7%)	18	41
36	1e	113/123 (92%)	108 (96%)	5 (4%)	33	63
36	2e	114/123 (93%)	109 (96%)	5 (4%)	33	63
37	1f	84/90 (93%)	82 (98%)	2 (2%)	54	83
37	2f	85/90 (94%)	83 (98%)	2 (2%)	54	83
38	1g	119/127 (94%)	111 (93%)	8 (7%)	19	42
38	2g	120/127 (94%)	116 (97%)	4 (3%)	43	73
39	1h	114/119 (96%)	109 (96%)	5 (4%)	33	63
39	2h	114/119 (96%)	109 (96%)	5 (4%)	33	63
40	1i	90/99 (91%)	86 (96%)	4 (4%)	33	63
40	2i	89/99 (90%)	80 (90%)	9 (10%)	9	20
41	1j	66/92 (72%)	62 (94%)	4 (6%)	22	47
41	2j	69/92 (75%)	68 (99%)	1 (1%)	71	90
42	1k	82/99 (83%)	77 (94%)	5 (6%)	22	47
42	2k	83/99 (84%)	80 (96%)	3 (4%)	40	70
43	1l	96/108 (89%)	93 (97%)	3 (3%)	45	75
43	2l	96/108 (89%)	93 (97%)	3 (3%)	45	75
44	1m	93/101 (92%)	84 (90%)	9 (10%)	9	22
44	2m	92/101 (91%)	89 (97%)	3 (3%)	43	73
45	1n	49/50 (98%)	42 (86%)	7 (14%)	4	9
45	2n	49/50 (98%)	47 (96%)	2 (4%)	35	66
46	1o	78/80 (98%)	73 (94%)	5 (6%)	20	45
46	2o	78/80 (98%)	76 (97%)	2 (3%)	51	81
47	1p	69/74 (93%)	66 (96%)	3 (4%)	33	64
47	2p	68/74 (92%)	63 (93%)	5 (7%)	16	37
48	1q	94/97 (97%)	91 (97%)	3 (3%)	44	75
48	2q	94/97 (97%)	90 (96%)	4 (4%)	33	64
49	1r	59/77 (77%)	57 (97%)	2 (3%)	42	73
49	2r	59/77 (77%)	57 (97%)	2 (3%)	42	73
50	1s	69/80 (86%)	66 (96%)	3 (4%)	33	64
50	2s	67/80 (84%)	65 (97%)	2 (3%)	46	76

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
51	1t	70/82 (85%)	69 (99%)	1 (1%)	71	90
51	2t	70/82 (85%)	70 (100%)	0	100	100
52	1u	18/22 (82%)	17 (94%)	1 (6%)	25	51
52	2u	18/22 (82%)	17 (94%)	1 (6%)	25	51
58	A	9/9 (100%)	7 (78%)	2 (22%)	1	3
58	B	9/9 (100%)	7 (78%)	2 (22%)	1	3
All	All	9321/10082 (92%)	8790 (94%)	531 (6%)	24	51

All (531) residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
3	1D	3	VAL
3	1D	14	ARG
3	1D	32	SER
3	1D	37	LEU
3	1D	39	LYS
3	1D	61	LEU
3	1D	99	ASP
3	1D	106	ILE
3	1D	113	VAL
3	1D	116	GLN
3	1D	155	LEU
3	1D	173	VAL
3	1D	193	VAL
3	1D	211	ARG
3	1D	221	VAL
3	1D	229	VAL
3	1D	242	ARG
3	1D	273	ARG
4	1E	12	THR
4	1E	21	VAL
4	1E	24	THR
4	1E	33	VAL
4	1E	34	VAL
4	1E	47	VAL
4	1E	73	GLU
4	1E	75	VAL
4	1E	116	VAL
4	1E	167	VAL
4	1E	170	LEU

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Mol	Chain	Res	Type
4	1E	175	VAL
4	1E	181	LEU
4	1E	184	VAL
4	1E	195	LEU
5	1F	12	LEU
5	1F	33	LEU
5	1F	53	THR
5	1F	57	VAL
5	1F	70	THR
5	1F	74	ARG
5	1F	88	VAL
5	1F	95	ARG
5	1F	106	ARG
5	1F	125	LEU
5	1F	170	LEU
5	1F	183	VAL
5	1F	192	LEU
6	1G	5	VAL
6	1G	31	VAL
6	1G	43	LEU
6	1G	82	LEU
6	1G	135	LEU
6	1G	140	ILE
6	1G	159	VAL
6	1G	175	LEU
7	1H	15	VAL
7	1H	23	ARG
7	1H	71	LEU
7	1H	81	GLU
7	1H	134	SER
7	1H	139	GLN
8	1I	9	LEU
8	1I	12	LEU
8	1I	38	LEU
8	1I	47	LEU
8	1I	77	LEU
8	1I	101	LEU
8	1I	109	ILE
8	1I	116	LEU
8	1I	123	LEU
8	1I	140	LEU
9	1N	14	VAL

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Mol	Chain	Res	Type
9	1N	28	THR
9	1N	33	LEU
9	1N	34	LEU
9	1N	48	MET
9	1N	62	VAL
9	1N	73	THR
9	1N	87	LEU
9	1N	99	LEU
10	1O	10	VAL
10	1O	23	ARG
10	1O	24	VAL
10	1O	69	ILE
11	1P	1	MET
11	1P	59	LEU
11	1P	77	ARG
11	1P	95	VAL
11	1P	112	LEU
11	1P	125	VAL
12	1Q	22	LYS
12	1Q	35	VAL
12	1Q	75	THR
12	1Q	101	ARG
12	1Q	109	VAL
12	1Q	110	THR
12	1Q	133	ARG
13	1R	6	SER
13	1R	29	LEU
13	1R	33	ARG
13	1R	36	THR
13	1R	44	LEU
13	1R	54	LEU
13	1R	65	LEU
13	1R	67	LEU
13	1R	96	ARG
13	1R	100	LEU
13	1R	111	LEU
13	1R	114	VAL
14	1S	14	VAL
14	1S	25	ARG
14	1S	36	TYR
14	1S	46	VAL
14	1S	69	VAL

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Mol	Chain	Res	Type
14	1S	73	LEU
14	1S	85	VAL
14	1S	110	LEU
15	1T	28	VAL
15	1T	67	SER
15	1T	95	ARG
15	1T	96	ARG
16	1U	8	VAL
16	1U	31	SER
16	1U	50	ARG
16	1U	74	LEU
16	1U	77	SER
16	1U	85	LYS
16	1U	95	LEU
17	1V	1	MET
17	1V	35	LEU
17	1V	46	VAL
17	1V	51	VAL
17	1V	52	VAL
17	1V	61	VAL
17	1V	62	LEU
17	1V	72	VAL
17	1V	73	SER
17	1V	79	VAL
18	1W	11	ARG
18	1W	15	ARG
18	1W	17	VAL
18	1W	23	LEU
18	1W	63	ASP
18	1W	92	ARG
18	1W	107	LEU
19	1X	35	THR
19	1X	52	VAL
19	1X	81	VAL
20	1Y	7	VAL
20	1Y	64	GLU
20	1Y	72	VAL
20	1Y	90	LEU
20	1Y	97	ARG
20	1Y	99	CYS
20	1Y	107	ASP
21	1Z	18	LEU

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Mol	Chain	Res	Type
21	1Z	33	LEU
21	1Z	61	LEU
21	1Z	91	LEU
21	1Z	126	VAL
21	1Z	129	SER
21	1Z	146	ILE
21	1Z	154	ASP
21	1Z	155	LEU
21	1Z	170	THR
21	1Z	171	ILE
22	10	10	THR
22	10	14	ARG
22	10	39	ARG
22	10	74	ARG
23	11	95	LEU
24	12	53	LEU
24	12	55	ARG
25	13	23	LEU
25	13	54	VAL
26	14	49	PHE
26	14	50	VAL
26	14	56	VAL
26	14	61	ARG
27	15	6	VAL
27	15	16	ARG
27	15	29	THR
27	15	58	LEU
28	16	6	ARG
28	16	9	LEU
28	16	19	ARG
28	16	38	LYS
28	16	44	ARG
28	16	48	VAL
29	17	1	MET
29	17	39	ARG
29	17	41	ARG
29	17	43	THR
29	17	46	VAL
30	18	6	THR
30	18	14	VAL
30	18	23	VAL
30	18	29	LYS

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Mol	Chain	Res	Type
30	18	31	HIS
30	18	32	LEU
30	18	34	TRP
33	1b	54	THR
33	1b	56	ARG
33	1b	78	GLN
33	1b	83	MET
33	1b	94	ASN
33	1b	111	ARG
33	1b	178	ARG
33	1b	185	ILE
34	1c	3	ASN
34	1c	112	SER
34	1c	115	LEU
34	1c	195	VAL
34	1c	206	GLU
35	1d	5	ILE
35	1d	19	LEU
35	1d	31	CYS
35	1d	59	ARG
35	1d	70	ILE
35	1d	107	ARG
35	1d	135	LEU
35	1d	178	VAL
35	1d	196	LEU
36	1e	16	THR
36	1e	20	GLN
36	1e	41	VAL
36	1e	67	VAL
36	1e	91	LEU
37	1f	21	LEU
37	1f	45	LEU
38	1g	15	ASP
38	1g	16	LEU
38	1g	59	LEU
38	1g	79	ARG
38	1g	104	LEU
38	1g	111	ARG
38	1g	114	ARG
38	1g	115	ARG
39	1h	25	ASP
39	1h	26	VAL

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Mol	Chain	Res	Type
39	1h	63	LEU
39	1h	85	ARG
39	1h	112	LEU
40	1i	54	ASP
40	1i	103	THR
40	1i	121	ARG
40	1i	128	ARG
41	1j	38	ILE
41	1j	43	ARG
41	1j	81	THR
41	1j	92	THR
42	1k	33	THR
42	1k	48	ILE
42	1k	87	THR
42	1k	109	VAL
42	1k	114	VAL
43	1l	33	ARG
43	1l	83	VAL
43	1l	113	ARG
44	1m	14	ARG
44	1m	19	LEU
44	1m	43	THR
44	1m	49	THR
44	1m	70	LEU
44	1m	102	ARG
44	1m	104	ARG
44	1m	105	THR
44	1m	117	VAL
45	1n	3	ARG
45	1n	6	LEU
45	1n	7	ILE
45	1n	18	VAL
45	1n	23	ARG
45	1n	32	SER
45	1n	33	VAL
46	1o	5	LYS
46	1o	39	LEU
46	1o	56	LEU
46	1o	65	ARG
46	1o	87	ILE
47	1p	2	VAL
47	1p	20	VAL

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Mol	Chain	Res	Type
47	1p	67	THR
48	1q	63	ARG
48	1q	70	ARG
48	1q	72	ARG
49	1r	26	LEU
49	1r	54	ARG
50	1s	70	LYS
50	1s	77	THR
50	1s	79	THR
51	1t	8	ARG
52	1u	24	ARG
3	2D	14	ARG
3	2D	37	LEU
3	2D	61	LEU
3	2D	111	LEU
3	2D	113	VAL
3	2D	138	VAL
3	2D	142	VAL
3	2D	173	VAL
3	2D	211	ARG
3	2D	242	ARG
3	2D	257	LEU
3	2D	259	THR
4	2E	9	VAL
4	2E	14	ILE
4	2E	21	VAL
4	2E	24	THR
4	2E	27	LEU
4	2E	75	VAL
4	2E	107	THR
4	2E	116	VAL
4	2E	170	LEU
4	2E	175	VAL
4	2E	181	LEU
5	2F	33	LEU
5	2F	53	THR
5	2F	70	THR
5	2F	74	ARG
5	2F	88	VAL
5	2F	106	ARG
5	2F	158	THR
5	2F	183	VAL

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Mol	Chain	Res	Type
5	2F	192	LEU
5	2F	195	ASP
5	2F	201	VAL
6	2G	5	VAL
6	2G	43	LEU
6	2G	91	ARG
6	2G	133	LEU
6	2G	140	ILE
6	2G	159	VAL
7	2H	70	THR
7	2H	71	LEU
7	2H	84	SER
7	2H	95	ARG
7	2H	134	SER
8	2I	38	LEU
8	2I	44	LEU
8	2I	47	LEU
8	2I	92	VAL
8	2I	101	LEU
8	2I	102	SER
8	2I	116	LEU
8	2I	123	LEU
8	2I	127	VAL
8	2I	144	VAL
9	2N	15	LEU
9	2N	28	THR
9	2N	34	LEU
9	2N	35	ARG
9	2N	46	VAL
9	2N	99	LEU
9	2N	140	VAL
10	2O	23	ARG
10	2O	52	VAL
10	2O	66	LYS
10	2O	116	SER
11	2P	58	THR
11	2P	95	VAL
11	2P	148	LEU
12	2Q	1	MET
12	2Q	35	VAL
12	2Q	38	GLU
12	2Q	60	ARG

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Mol	Chain	Res	Type
12	2Q	75	THR
12	2Q	109	VAL
12	2Q	110	THR
12	2Q	133	ARG
13	2R	6	SER
13	2R	24	GLN
13	2R	29	LEU
13	2R	36	THR
13	2R	44	LEU
13	2R	54	LEU
13	2R	65	LEU
13	2R	96	ARG
13	2R	100	LEU
13	2R	111	LEU
13	2R	114	VAL
14	2S	23	ARG
14	2S	25	ARG
14	2S	36	TYR
14	2S	52	SER
14	2S	80	LEU
14	2S	110	LEU
15	2T	41	ARG
15	2T	104	ASN
15	2T	108	ARG
15	2T	115	ARG
16	2U	8	VAL
16	2U	31	SER
16	2U	74	LEU
17	2V	7	THR
17	2V	12	TYR
17	2V	46	VAL
17	2V	62	LEU
17	2V	66	ARG
17	2V	72	VAL
17	2V	79	VAL
18	2W	11	ARG
18	2W	17	VAL
18	2W	19	LEU
18	2W	23	LEU
18	2W	107	LEU
20	2Y	49	VAL
20	2Y	97	ARG

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Mol	Chain	Res	Type
21	2Z	33	LEU
21	2Z	42	VAL
21	2Z	53	ILE
21	2Z	71	VAL
21	2Z	72	ARG
21	2Z	84	GLU
21	2Z	91	LEU
21	2Z	144	LEU
21	2Z	154	ASP
21	2Z	170	THR
21	2Z	174	VAL
22	20	9	SER
22	20	14	ARG
22	20	27	GLU
22	20	39	ARG
22	20	74	ARG
23	21	35	THR
23	21	95	LEU
24	22	28	LYS
24	22	52	ASP
24	22	59	ARG
25	23	31	LEU
25	23	54	VAL
26	24	49	PHE
26	24	53	GLU
27	25	6	VAL
27	25	29	THR
27	25	40	LYS
27	25	58	LEU
28	26	14	THR
28	26	19	ARG
28	26	44	ARG
29	27	1	MET
29	27	39	ARG
29	27	41	ARG
29	27	43	THR
30	28	14	VAL
30	28	31	HIS
30	28	32	LEU
30	28	37	SER
31	29	7	VAL
33	2b	23	ARG

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Mol	Chain	Res	Type
33	2b	93	VAL
33	2b	94	ASN
33	2b	111	ARG
33	2b	124	SER
33	2b	127	ILE
33	2b	176	GLU
33	2b	178	ARG
33	2b	185	ILE
33	2b	189	ASP
33	2b	192	SER
33	2b	230	VAL
34	2c	128	PHE
34	2c	166	GLU
35	2d	31	CYS
35	2d	83	SER
35	2d	107	ARG
35	2d	108	LEU
35	2d	135	LEU
35	2d	150	GLU
35	2d	170	VAL
35	2d	175	SER
35	2d	178	VAL
35	2d	181	MET
35	2d	208	SER
35	2d	209	ARG
36	2e	41	VAL
36	2e	68	GLU
36	2e	75	THR
36	2e	111	GLU
36	2e	120	THR
37	2f	72	VAL
37	2f	93	SER
38	2g	13	GLN
38	2g	16	LEU
38	2g	78	ARG
38	2g	79	ARG
39	2h	3	THR
39	2h	14	ARG
39	2h	51	VAL
39	2h	85	ARG
39	2h	112	LEU
40	2i	3	GLN

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Mol	Chain	Res	Type
40	2i	14	VAL
40	2i	53	VAL
40	2i	64	THR
40	2i	71	SER
40	2i	75	ASP
40	2i	102	LEU
40	2i	108	VAL
40	2i	128	ARG
41	2j	81	THR
42	2k	14	VAL
42	2k	81	ASP
42	2k	114	VAL
43	2l	83	VAL
43	2l	113	ARG
43	2l	117	ARG
44	2m	19	LEU
44	2m	90	LEU
44	2m	108	ARG
45	2n	18	VAL
45	2n	33	VAL
46	2o	31	LEU
46	2o	39	LEU
47	2p	1	MET
47	2p	2	VAL
47	2p	21	VAL
47	2p	67	THR
47	2p	69	THR
48	2q	14	LYS
48	2q	58	GLU
48	2q	83	ASP
48	2q	87	LYS
49	2r	37	VAL
49	2r	76	LEU
50	2s	4	SER
50	2s	5	LEU
52	2u	10	ARG
58	A	12	SER
58	A	14	SER
58	B	12	SER
58	B	14	SER

Some sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (78) such sidechains are listed below:

Mol	Chain	Res	Type
3	1D	126	GLN
5	1F	203	GLN
6	1G	26	GLN
12	1Q	57	HIS
14	1S	95	HIS
19	1X	31	HIS
19	1X	82	GLN
20	1Y	6	HIS
20	1Y	43	ASN
21	1Z	73	GLN
21	1Z	151	HIS
22	10	50	ASN
23	11	56	GLN
25	13	32	GLN
34	1c	6	HIS
34	1c	37	GLN
34	1c	162	GLN
35	1d	42	GLN
35	1d	77	ASN
35	1d	119	GLN
35	1d	123	HIS
35	1d	125	HIS
36	1e	78	HIS
37	1f	100	ASN
38	1g	28	ASN
40	1i	3	GLN
40	1i	31	GLN
40	1i	58	HIS
40	1i	87	GLN
40	1i	124	GLN
41	1j	56	HIS
43	1l	99	HIS
45	1n	49	HIS
46	1o	28	GLN
47	1p	13	HIS
50	1s	23	ASN
50	1s	69	HIS
50	1s	83	HIS
51	1t	16	HIS
3	2D	87	ASN
4	2E	48	GLN
5	2F	69	HIS
6	2G	66	GLN

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Mol	Chain	Res	Type
12	2Q	123	HIS
13	2R	31	HIS
13	2R	71	GLN
17	2V	80	GLN
18	2W	60	ASN
19	2X	31	HIS
21	2Z	73	GLN
22	20	50	ASN
23	21	56	GLN
25	23	32	GLN
26	24	46	GLN
33	2b	40	HIS
33	2b	94	ASN
34	2c	6	HIS
34	2c	102	ASN
35	2d	125	HIS
36	2e	78	HIS
37	2f	73	ASN
37	2f	100	ASN
38	2g	28	ASN
38	2g	86	GLN
38	2g	109	ASN
40	2i	3	GLN
40	2i	58	HIS
40	2i	87	GLN
41	2j	56	HIS
44	2m	77	ASN
46	2o	62	GLN
49	2r	63	GLN
50	2s	23	ASN
50	2s	47	HIS
50	2s	69	HIS
50	2s	83	HIS
58	A	5	ASN
58	B	5	ASN

5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	1A	2861/2915 (98%)	407 (14%)	38 (1%)
1	2A	2788/2915 (95%)	458 (16%)	25 (0%)

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Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
2	1B	120/121 (99%)	10 (8%)	1 (0%)
2	2B	118/121 (97%)	27 (22%)	0
32	1a	1494/1521 (98%)	227 (15%)	0
53	1v	12/24 (50%)	2 (16%)	0
53	2v	12/24 (50%)	1 (8%)	0
54	1w	71/76 (93%)	23 (32%)	0
54	2w	68/76 (89%)	22 (32%)	0
55	1x	75/77 (97%)	10 (13%)	0
55	2x	75/77 (97%)	14 (18%)	0
56	1y	72/76 (94%)	19 (26%)	0
56	2y	70/76 (92%)	18 (25%)	0
57	2a	1498/1521 (98%)	260 (17%)	0
All	All	9334/9620 (97%)	1498 (16%)	64 (0%)

All (1498) RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	1A	12	U
1	1A	34	C
1	1A	45	C
1	1A	54	G
1	1A	70	A
1	1A	71	U
1	1A	73	A
1	1A	74	G
1	1A	77	A
1	1A	83	A
1	1A	94	G
1	1A	116	A
1	1A	117	A
1	1A	118	U
1	1A	137	G
1	1A	170	A
1	1A	171	A
1	1A	185	A
1	1A	188	A
1	1A	194	G
1	1A	203	G
1	1A	204	G
1	1A	205	A
1	1A	211	A
1	1A	217	A

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Mol	Chain	Res	Type
1	1A	218	A
1	1A	222	A
1	1A	237	G
1	1A	271	U
1	1A	272	U
1	1A	273	G
1	1A	275	C
1	1A	276	C
1	1A	289	G
1	1A	296	U
1	1A	299	G
1	1A	303	C
1	1A	335	A
1	1A	353	G
1	1A	354	A
1	1A	376	G
1	1A	387	G
1	1A	388	A
1	1A	389	G
1	1A	407	U
1	1A	413	G
1	1A	423	G
1	1A	432	U
1	1A	438	G
1	1A	439	A
1	1A	448	U
1	1A	455	A
1	1A	470	C
1	1A	474	U
1	1A	480	A
1	1A	482	C
1	1A	483	A
1	1A	507	G
1	1A	529	U
1	1A	530	A
1	1A	533	G
1	1A	534	C
1	1A	537	G
1	1A	553	A
1	1A	554	A
1	1A	555	G
1	1A	556	C

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Mol	Chain	Res	Type
1	1A	557	A
1	1A	558	G
1	1A	569	G
1	1A	573	G
1	1A	586	G
1	1A	596	G
1	1A	597	C
1	1A	598	A
1	1A	609	A
1	1A	626	A
1	1A	627	G
1	1A	630	U
1	1A	639	G
1	1A	641	G
1	1A	642	G
1	1A	652	A
1	1A	662	A
1	1A	670	C
1	1A	671	A
1	1A	678	A
1	1A	697	C
1	1A	699	C
1	1A	716	G
1	1A	733	G
1	1A	764	G
1	1A	777	C
1	1A	811	A
1	1A	812	G
1	1A	822	G
1	1A	823	G
1	1A	829	A
1	1A	830	A
1	1A	831	A
1	1A	832	G
1	1A	837	C
1	1A	839	G
1	1A	852	G
1	1A	859	C
1	1A	866	A
1	1A	874	U
1	1A	875	U
1	1A	906	G

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Mol	Chain	Res	Type
1	1A	913	A
1	1A	924	U
1	1A	926	G
1	1A	927	G
1	1A	931	C
1	1A	932	C
1	1A	933	C
1	1A	934	A
1	1A	935	C
1	1A	936	C
1	1A	937	A
1	1A	940	C
1	1A	942	A
1	1A	943	C
1	1A	944	C
1	1A	946	A
1	1A	956	A
1	1A	977	G
1	1A	990	A
1	1A	991	G
1	1A	1003	U
1	1A	1004	A
1	1A	1006	C
1	1A	1008	U
1	1A	1019	G
1	1A	1020	C
1	1A	1029	A
1	1A	1042	A
1	1A	1051	C
1	1A	1058	U
1	1A	1059	C
1	1A	1068	G
1	1A	1072	U
1	1A	1073	A
1	1A	1079	U
1	1A	1084	C
1	1A	1087	C
1	1A	1091	A
1	1A	1092	A
1	1A	1093	G
1	1A	1094	A
1	1A	1097	G

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Mol	Chain	Res	Type
1	1A	1100	A
1	1A	1101	G
1	1A	1104	G
1	1A	1109	G
1	1A	1113	A
1	1A	1114	G
1	1A	1116	A
1	1A	1117	G
1	1A	1119	A
1	1A	1120	G
1	1A	1121	C
1	1A	1122	C
1	1A	1124	U
1	1A	1125	C
1	1A	1134	A
1	1A	1135	G
1	1A	1136	U
1	1A	1137	G
1	1A	1139	G
1	1A	1140	U
1	1A	1142	A
1	1A	1147	U
1	1A	1157	A
1	1A	1158	G
1	1A	1161	G
1	1A	1162	C
1	1A	1174	A
1	1A	1180	C
1	1A	1181	G
1	1A	1217	G
1	1A	1218	G
1	1A	1219	A
1	1A	1220	U
1	1A	1221	G
1	1A	1222	A
1	1A	1223	C
1	1A	1256	U
1	1A	1290	G
1	1A	1299	A
1	1A	1302	G
1	1A	1317	G
1	1A	1318	A

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Mol	Chain	Res	Type
1	1A	1346	U
1	1A	1347	A
1	1A	1349	G
1	1A	1366	C
1	1A	1398	U
1	1A	1405	A
1	1A	1406	A
1	1A	1411	A
1	1A	1418	U
1	1A	1426	G
1	1A	1430	A
1	1A	1431	G
1	1A	1462	G
1	1A	1463	C
1	1A	1466	U
1	1A	1467	G
1	1A	1474	C
1	1A	1491	A
1	1A	1497	G
1	1A	1502	G
1	1A	1514	C
1	1A	1529	G
1	1A	1536	A
1	1A	1539	C
1	1A	1542	A
1	1A	1554	A
1	1A	1555	C
1	1A	1556	A
1	1A	1586	G
1	1A	1589	A
1	1A	1590	C
1	1A	1605	A
1	1A	1606	G
1	1A	1613	A
1	1A	1616	A
1	1A	1625	U
1	1A	1628	G
1	1A	1631	C
1	1A	1632	A
1	1A	1654	A
1	1A	1655	A
1	1A	1656	A

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Mol	Chain	Res	Type
1	1A	1695	C
1	1A	1701	A
1	1A	1721	G
1	1A	1743	G
1	1A	1747	A
1	1A	1750	G
1	1A	1767	A
1	1A	1776	G
1	1A	1787	G
1	1A	1794	G
1	1A	1795	G
1	1A	1804	A
1	1A	1811	A
1	1A	1813	C
1	1A	1822	A
1	1A	1831	C
1	1A	1832	G
1	1A	1847	G
1	1A	1860	A
1	1A	1878	A
1	1A	1889	G
1	1A	1899	A
1	1A	1900	G
1	1A	1911	A
1	1A	1922	A
1	1A	1928	G
1	1A	1941	A
1	1A	1951	G
1	1A	1952	G
1	1A	1959	A
1	1A	1960	A
1	1A	1977	U
1	1A	1985	U
1	1A	1987	C
1	1A	1989	C
1	1A	1992	A
1	1A	1993	A
1	1A	1994	A
1	1A	2014	G
1	1A	2015	U
1	1A	2019	G
1	1A	2042	A

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Mol	Chain	Res	Type
1	1A	2045	G
1	1A	2053	A
1	1A	2054	G
1	1A	2055	A
1	1A	2061	C
1	1A	2065	C
1	1A	2077	C
1	1A	2078	G
1	1A	2082	A
1	1A	2083	G
1	1A	2091	G
1	1A	2132	G
1	1A	2135	U
1	1A	2137	G
1	1A	2138	G
1	1A	2143	G
1	1A	2149	G
1	1A	2151	C
1	1A	2152	U
1	1A	2153	G
1	1A	2154	U
1	1A	2155	G
1	1A	2156	A
1	1A	2157	A
1	1A	2158	C
1	1A	2163	G
1	1A	2164	C
1	1A	2165	C
1	1A	2166	U
1	1A	2168	C
1	1A	2169	G
1	1A	2172	U
1	1A	2173	G
1	1A	2178	G
1	1A	2179	G
1	1A	2180	A
1	1A	2181	G
1	1A	2187	G
1	1A	2188	G
1	1A	2189	U
1	1A	2190	G
1	1A	2191	A

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Mol	Chain	Res	Type
1	1A	2193	A
1	1A	2194	U
1	1A	2196	C
1	1A	2200	C
1	1A	2204	G
1	1A	2206	G
1	1A	2214	G
1	1A	2220	A
1	1A	2227	G
1	1A	2228	G
1	1A	2229	A
1	1A	2237	A
1	1A	2250	G
1	1A	2251	G
1	1A	2280	A
1	1A	2281	A
1	1A	2292	G
1	1A	2295	C
1	1A	2299	A
1	1A	2301	G
1	1A	2317	A
1	1A	2320	G
1	1A	2325	C
1	1A	2332	A
1	1A	2337	G
1	1A	2346	G
1	1A	2348	A
1	1A	2359	C
1	1A	2362	C
1	1A	2373	A
1	1A	2395	G
1	1A	2397	C
1	1A	2411	G
1	1A	2418	U
1	1A	2437	A
1	1A	2441	G
1	1A	2442	A
1	1A	2443	U
1	1A	2447	A
1	1A	2451	A
1	1A	2452	C
1	1A	2453	C

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Mol	Chain	Res	Type
1	1A	2460	A
1	1A	2488	A
1	1A	2490	A
1	1A	2514	G
1	1A	2517	G
1	1A	2518	U
1	1A	2530	A
1	1A	2541	G
1	1A	2566	U
1	1A	2578	A
1	1A	2579	G
1	1A	2585	C
1	1A	2598	C
1	1A	2614	A
1	1A	2621	U
1	1A	2623	U
1	1A	2624	C
1	1A	2641	A
1	1A	2642	G
1	1A	2666	A
1	1A	2701	U
1	1A	2702	C
1	1A	2714	U
1	1A	2715	C
1	1A	2725	A
1	1A	2726	A
1	1A	2727	G
1	1A	2739	U
1	1A	2746	A
1	1A	2770	A
1	1A	2778	A
1	1A	2779	G
1	1A	2782	C
1	1A	2791	A
1	1A	2803	A
1	1A	2804	C
1	1A	2806	G
1	1A	2813	G
1	1A	2814	C
1	1A	2830	A
1	1A	2831	A
1	1A	2845	A

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Mol	Chain	Res	Type
1	1A	2882	G
1	1A	2890	C
1	1A	2901	A
1	1A	2903	G
2	1B	2	C
2	1B	15	A
2	1B	45	A
2	1B	50	G
2	1B	52	A
2	1B	56	G
2	1B	67	G
2	1B	73	A
2	1B	106	G
2	1B	110	G
32	1a	7	G
32	1a	9	G
32	1a	32	A
32	1a	39	G
32	1a	48	C
32	1a	51	A
32	1a	54	C
32	1a	61	G
32	1a	69	G
32	1a	79	G
32	1a	91	C
32	1a	96	U
32	1a	98	G
32	1a	101	A
32	1a	105	G
32	1a	116	A
32	1a	121	C
32	1a	131	C
32	1a	144	G
32	1a	145	G
32	1a	162	A
32	1a	163	C
32	1a	174	C
32	1a	182	U
32	1a	189(J)	G
32	1a	195	A
32	1a	197	A
32	1a	201	C

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Mol	Chain	Res	Type
32	1a	202	U
32	1a	203	U
32	1a	204	U
32	1a	216	G
32	1a	218	C
32	1a	247	G
32	1a	251	G
32	1a	258	G
32	1a	266	G
32	1a	267	C
32	1a	289	G
32	1a	301	G
32	1a	321	A
32	1a	328	C
32	1a	332	G
32	1a	342	C
32	1a	348	G
32	1a	351	G
32	1a	352	C
32	1a	353	A
32	1a	354	G
32	1a	367	U
32	1a	372	C
32	1a	373	A
32	1a	384	G
32	1a	392	G
32	1a	397	A
32	1a	398	C
32	1a	406	G
32	1a	412	A
32	1a	421	U
32	1a	424	G
32	1a	429	U
32	1a	439	A
32	1a	442	C
32	1a	452	A
32	1a	458	C
32	1a	461	A
32	1a	470	C
32	1a	485	G
32	1a	496	A
32	1a	498	U

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Mol	Chain	Res	Type
32	1a	505	G
32	1a	509	A
32	1a	510	A
32	1a	511	C
32	1a	518	C
32	1a	524	G
32	1a	527	G7M
32	1a	531	U
32	1a	532	A
32	1a	533	A
32	1a	547	A
32	1a	559	A
32	1a	560	U
32	1a	561	U
32	1a	562	C
32	1a	568	G
32	1a	572	A
32	1a	573	A
32	1a	576	G
32	1a	577	G
32	1a	596	C
32	1a	630	G
32	1a	653	A
32	1a	659	U
32	1a	665	A
32	1a	673	G
32	1a	687	A
32	1a	688	G
32	1a	695	A
32	1a	723	U
32	1a	731	G
32	1a	749	C
32	1a	752	G
32	1a	755	G
32	1a	774	G
32	1a	777	A
32	1a	792	A
32	1a	793	U
32	1a	794	A
32	1a	815	A
32	1a	816	A
32	1a	817	C

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Mol	Chain	Res	Type
32	1a	821	G
32	1a	828	A
32	1a	840	C
32	1a	841	U
32	1a	851	G
32	1a	870	U
32	1a	902	G
32	1a	914	A
32	1a	926	G
32	1a	927	G
32	1a	934	C
32	1a	935	A
32	1a	960	U
32	1a	961	U
32	1a	968	A
32	1a	969	A
32	1a	971	G
32	1a	972	C
32	1a	974	A
32	1a	975	A
32	1a	976	G
32	1a	977	A
32	1a	992	U
32	1a	993	G
32	1a	1000	U
32	1a	1001(A)	G
32	1a	1003	G
32	1a	1005	A
32	1a	1006	C
32	1a	1009	G
32	1a	1022	G
32	1a	1023	G
32	1a	1024	G
32	1a	1025	U
32	1a	1026	G
32	1a	1027	C
32	1a	1029	C
32	1a	1030	C
32	1a	1030(A)	G
32	1a	1030(C)	G
32	1a	1030(D)	A
32	1a	1031	G

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Mol	Chain	Res	Type
32	1a	1033	G
32	1a	1039	C
32	1a	1044	A
32	1a	1054	C
32	1a	1065	U
32	1a	1066	C
32	1a	1068	G
32	1a	1081	G
32	1a	1094	G
32	1a	1095	U
32	1a	1101	A
32	1a	1125	U
32	1a	1134	G
32	1a	1137	C
32	1a	1138	G
32	1a	1139	G
32	1a	1140	C
32	1a	1141	C
32	1a	1152	A
32	1a	1157	A
32	1a	1159	U
32	1a	1160	G
32	1a	1168	A
32	1a	1172	C
32	1a	1184	G
32	1a	1196	U
32	1a	1197	G
32	1a	1201	A
32	1a	1202	G
32	1a	1212	U
32	1a	1213	A
32	1a	1227	A
32	1a	1236	A
32	1a	1238	A
32	1a	1256	A
32	1a	1257	U
32	1a	1258	G
32	1a	1260	C
32	1a	1270	C
32	1a	1273	G
32	1a	1275	A
32	1a	1278	U

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Mol	Chain	Res	Type
32	1a	1279	A
32	1a	1280	A
32	1a	1286	A
32	1a	1287	A
32	1a	1299	A
32	1a	1300	G
32	1a	1302	U
32	1a	1320	C
32	1a	1338	G
32	1a	1347	G
32	1a	1353	G
32	1a	1363	C
32	1a	1364	U
32	1a	1370	G
32	1a	1381	U
32	1a	1397	C
32	1a	1419	G
32	1a	1442	G
32	1a	1442(A)	G
32	1a	1446	U
32	1a	1456	G
32	1a	1487	G
32	1a	1492	A
32	1a	1494	G
32	1a	1503	A
32	1a	1504	G
32	1a	1506	U
32	1a	1507	A
32	1a	1517	G
32	1a	1529	G
32	1a	1530	G
53	1v	13	A
53	1v	24	A
54	1w	2	C
54	1w	6	G
54	1w	7	A
54	1w	8	4SU
54	1w	14	A
54	1w	19	G
54	1w	20	U
54	1w	21	A
54	1w	23	A

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Mol	Chain	Res	Type
54	1w	24	G
54	1w	45	U
54	1w	46	G7M
54	1w	47	U
54	1w	48	C
54	1w	50	U
54	1w	62	C
54	1w	67	C
54	1w	68	C
54	1w	69	G
54	1w	70	G
54	1w	72	C
54	1w	73	A
54	1w	74	C
55	1x	9	G
55	1x	14	A
55	1x	18	G
55	1x	19	G
55	1x	21	A
55	1x	47	U
55	1x	61	C
55	1x	63	G
55	1x	69	C
55	1x	76	A
56	1y	6	G
56	1y	8	4SU
56	1y	13	C
56	1y	19	G
56	1y	20	U
56	1y	21	A
56	1y	35	A
56	1y	36	A
56	1y	44	G
56	1y	45	U
56	1y	46	G7M
56	1y	48	C
56	1y	49	C
56	1y	53	G
56	1y	54	5MU
56	1y	59	U
56	1y	65	G
56	1y	69	G

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Mol	Chain	Res	Type
56	1y	70	G
1	2A	11	G
1	2A	15	G
1	2A	34	C
1	2A	35	G
1	2A	45	C
1	2A	71	A
1	2A	74	A
1	2A	75	G
1	2A	84	A
1	2A	90	U
1	2A	94	C
1	2A	100	G
1	2A	102	G
1	2A	118	A
1	2A	119	A
1	2A	120	U
1	2A	141	A
1	2A	154(A)	C
1	2A	157	U
1	2A	173	G
1	2A	181	A
1	2A	196	A
1	2A	199	A
1	2A	205	G
1	2A	214	G
1	2A	215	G
1	2A	216	A
1	2A	222	A
1	2A	225	A
1	2A	228	A
1	2A	229	A
1	2A	233	A
1	2A	248	G
1	2A	249	C
1	2A	250	G
1	2A	266	G
1	2A	267	C
1	2A	271(K)	U
1	2A	271(L)	U
1	2A	271(M)	G
1	2A	271(N)	U

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Mol	Chain	Res	Type
1	2A	272(B)	G
1	2A	277	C
1	2A	278	A
1	2A	279	C
1	2A	283	A
1	2A	294	A
1	2A	311	A
1	2A	327	G
1	2A	329	G
1	2A	330	A
1	2A	342	G
1	2A	352	G
1	2A	354	G
1	2A	362	U
1	2A	363	G
1	2A	363(B)	G
1	2A	363(D)	G
1	2A	386	G
1	2A	396	G
1	2A	405	U
1	2A	406	G
1	2A	411	G
1	2A	412	A
1	2A	421	U
1	2A	422	A
1	2A	435	C
1	2A	444	C
1	2A	455	C
1	2A	457	A
1	2A	481	G
1	2A	494	G
1	2A	505	A
1	2A	508	G
1	2A	509	C
1	2A	528	A
1	2A	529	A
1	2A	531	C
1	2A	532	A
1	2A	533	G
1	2A	563	G
1	2A	568	U
1	2A	573	G

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Mol	Chain	Res	Type
1	2A	575	A
1	2A	588	U
1	2A	592	G
1	2A	599	G
1	2A	603	A
1	2A	604	G
1	2A	607	U
1	2A	614(B)	G
1	2A	615	G
1	2A	616	G
1	2A	627	A
1	2A	634	C
1	2A	637	A
1	2A	645	C
1	2A	651	G
1	2A	652(B)	A
1	2A	652(C)	G
1	2A	669	G
1	2A	686	G
1	2A	717	G
1	2A	726	G
1	2A	730	C
1	2A	752	A
1	2A	753	C
1	2A	771	G
1	2A	775	G
1	2A	776	G
1	2A	782	A
1	2A	784	A
1	2A	785	G
1	2A	792	G
1	2A	805	G
1	2A	812	C
1	2A	819	A
1	2A	827	U
1	2A	828	U
1	2A	852	G
1	2A	857	C
1	2A	859	G
1	2A	866	A
1	2A	869	G
1	2A	870	A

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Mol	Chain	Res	Type
1	2A	874	G
1	2A	875	G
1	2A	877	U
1	2A	878	A
1	2A	879	G
1	2A	880	G
1	2A	883	G
1	2A	884	C
1	2A	886	C
1	2A	887	A
1	2A	888	C
1	2A	889	C
1	2A	893	C
1	2A	894	C
1	2A	895	U
1	2A	896	A
1	2A	900	A
1	2A	901	A
1	2A	910	A
1	2A	914	C
1	2A	917	A
1	2A	923	C
1	2A	932	G
1	2A	933	A
1	2A	941	A
1	2A	945	A
1	2A	946	G
1	2A	953	A
1	2A	958	U
1	2A	959	A
1	2A	961	C
1	2A	974	G
1	2A	975	C
1	2A	983	A
1	2A	996	A
1	2A	997	G
1	2A	1012	U
1	2A	1013	C
1	2A	1017	G
1	2A	1020	A
1	2A	1022	G
1	2A	1025	G

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Mol	Chain	Res	Type
1	2A	1026	U
1	2A	1033	U
1	2A	1037	G
1	2A	1038	C
1	2A	1039	G
1	2A	1041	C
1	2A	1043	C
1	2A	1116	C
1	2A	1135	C
1	2A	1136	G
1	2A	1137	G
1	2A	1139	G
1	2A	1142(A)	A
1	2A	1144	G
1	2A	1155	A
1	2A	1171	G
1	2A	1188	U
1	2A	1210	A
1	2A	1211	U
1	2A	1220	A
1	2A	1248	G
1	2A	1253	A
1	2A	1256	G
1	2A	1271	G
1	2A	1272	A
1	2A	1273	U
1	2A	1284	A
1	2A	1287	A
1	2A	1300	U
1	2A	1301	A
1	2A	1308	A
1	2A	1314	C
1	2A	1345	C
1	2A	1352	U
1	2A	1359	A
1	2A	1360	A
1	2A	1365	A
1	2A	1368	G
1	2A	1370	C
1	2A	1373	A
1	2A	1379	A
1	2A	1384	A

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Mol	Chain	Res	Type
1	2A	1385	G
1	2A	1416	G
1	2A	1417	C
1	2A	1420	U
1	2A	1421	G
1	2A	1427	A
1	2A	1428	C
1	2A	1437	C
1	2A	1445	A
1	2A	1449	A
1	2A	1450	G
1	2A	1460	A
1	2A	1461	G
1	2A	1467	C
1	2A	1471	A
1	2A	1482	G
1	2A	1490	A
1	2A	1493	C
1	2A	1496	A
1	2A	1497	U
1	2A	1508	A
1	2A	1509	C
1	2A	1509(A)	A
1	2A	1531	C
1	2A	1532	C
1	2A	1533	G
1	2A	1543	C
1	2A	1545	A
1	2A	1547	C
1	2A	1558	A
1	2A	1559	G
1	2A	1569	A
1	2A	1578	U
1	2A	1583	A
1	2A	1584	C
1	2A	1586	A
1	2A	1603	A
1	2A	1608	A
1	2A	1609	A
1	2A	1610	A
1	2A	1640	C
1	2A	1647	G

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Mol	Chain	Res	Type
1	2A	1648	C
1	2A	1653	G
1	2A	1654	A
1	2A	1674	G
1	2A	1696	G
1	2A	1700	A
1	2A	1701	A
1	2A	1703	G
1	2A	1721	G
1	2A	1722	A
1	2A	1739	U
1	2A	1740	G
1	2A	1746	G
1	2A	1747	G
1	2A	1756	G
1	2A	1762	A
1	2A	1763	G
1	2A	1764	G
1	2A	1773	A
1	2A	1780	A
1	2A	1782	C
1	2A	1786	A
1	2A	1791	A
1	2A	1800	C
1	2A	1801	G
1	2A	1811	G
1	2A	1812	A
1	2A	1816	G
1	2A	1829	A
1	2A	1835	G
1	2A	1847	A
1	2A	1848	A
1	2A	1877	A
1	2A	1878	G
1	2A	1889	A
1	2A	1900	A
1	2A	1906	G
1	2A	1912	A
1	2A	1914	C
1	2A	1929	G
1	2A	1930	G
1	2A	1931	U

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Mol	Chain	Res	Type
1	2A	1936	A
1	2A	1938	A
1	2A	1955	U
1	2A	1963	U
1	2A	1967	C
1	2A	1970	A
1	2A	1971	A
1	2A	1972	A
1	2A	1993	U
1	2A	1997	G
1	2A	2020	A
1	2A	2023	G
1	2A	2031	A
1	2A	2032	G
1	2A	2033	A
1	2A	2043	C
1	2A	2055	C
1	2A	2056	G
1	2A	2060	A
1	2A	2061	G
1	2A	2069	G
1	2A	2096	U
1	2A	2097	C
1	2A	2104	G
1	2A	2105	C
1	2A	2111	C
1	2A	2112	G
1	2A	2116	G
1	2A	2119	A
1	2A	2120	G
1	2A	2122	U
1	2A	2124	G
1	2A	2126	A
1	2A	2127	G
1	2A	2129	C
1	2A	2130	U
1	2A	2131	G
1	2A	2132	U
1	2A	2134	A
1	2A	2135	A
1	2A	2136	C
1	2A	2137	C

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Mol	Chain	Res	Type
1	2A	2138	C
1	2A	2140	C
1	2A	2142	C
1	2A	2146	C
1	2A	2150	U
1	2A	2153	G
1	2A	2154	G
1	2A	2155	G
1	2A	2156	G
1	2A	2157	G
1	2A	2158	A
1	2A	2159	G
1	2A	2161	C
1	2A	2166	G
1	2A	2167	U
1	2A	2168	G
1	2A	2172	U
1	2A	2174	C
1	2A	2182	G
1	2A	2185	C
1	2A	2189	U
1	2A	2192	G
1	2A	2198	A
1	2A	2206	G
1	2A	2207	G
1	2A	2208	A
1	2A	2218	U
1	2A	2219	G
1	2A	2225	A
1	2A	2238	G
1	2A	2239	G
1	2A	2275	C
1	2A	2279	G
1	2A	2283	C
1	2A	2287	A
1	2A	2288	A
1	2A	2294	C
1	2A	2305	A
1	2A	2308	G
1	2A	2320	A
1	2A	2325	G
1	2A	2334	G

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Mol	Chain	Res	Type
1	2A	2336	A
1	2A	2347	C
1	2A	2350	C
1	2A	2372	G
1	2A	2376	A
1	2A	2383	G
1	2A	2385	C
1	2A	2402	C
1	2A	2403	C
1	2A	2406	U
1	2A	2419	U
1	2A	2425	A
1	2A	2429	G
1	2A	2430	A
1	2A	2434	A
1	2A	2435	A
1	2A	2439	A
1	2A	2441	C
1	2A	2445	G
1	2A	2448	A
1	2A	2468	G
1	2A	2469	A
1	2A	2476	A
1	2A	2487	G
1	2A	2490	G
1	2A	2494	G
1	2A	2502	G
1	2A	2505	G
1	2A	2506	U
1	2A	2507	C
1	2A	2518	A
1	2A	2520	C
1	2A	2525	G
1	2A	2536	G
1	2A	2554	U
1	2A	2562	U
1	2A	2564	A
1	2A	2566	A
1	2A	2567	G
1	2A	2573	C
1	2A	2574	G
1	2A	2586	C

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Mol	Chain	Res	Type
1	2A	2602	A
1	2A	2609	U
1	2A	2611	U
1	2A	2612	C
1	2A	2630	G
1	2A	2634	G
1	2A	2654	A
1	2A	2689	U
1	2A	2690	C
1	2A	2702	U
1	2A	2712(A)	A
1	2A	2713	A
1	2A	2714	G
1	2A	2718	G
1	2A	2726	U
1	2A	2733	A
1	2A	2751	G
1	2A	2758	A
1	2A	2761	G
1	2A	2764	A
1	2A	2765	A
1	2A	2766	G
1	2A	2778	A
1	2A	2780	G
1	2A	2789	C
1	2A	2793	G
1	2A	2802	G
1	2A	2803	C
1	2A	2818	G
1	2A	2820	A
1	2A	2821	A
1	2A	2833	G
1	2A	2835	A
1	2A	2872	G
1	2A	2880	C
1	2A	2892	A
1	2A	2894	G
1	2A	2895	U
1	2A	2897	U
2	2B	2	C
2	2B	8	U
2	2B	13	A

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Mol	Chain	Res	Type
2	2B	19	G
2	2B	25	A
2	2B	30	C
2	2B	32	C
2	2B	34	U
2	2B	42	C
2	2B	53	A
2	2B	56	G
2	2B	63	G
2	2B	65	C
2	2B	67	G
2	2B	72	G
2	2B	73	A
2	2B	74	U
2	2B	75	G
2	2B	85	G
2	2B	90	A
2	2B	106	G
2	2B	108	U
2	2B	110	G
2	2B	111	G
2	2B	114	C
2	2B	116	G
2	2B	120	A
57	2a	9	G
57	2a	22	G
57	2a	31	G
57	2a	32	A
57	2a	39	G
57	2a	47	C
57	2a	48	C
57	2a	50	A
57	2a	51	A
57	2a	66	G
57	2a	73	G
57	2a	79	G
57	2a	89	C
57	2a	98	G
57	2a	101	A
57	2a	116	A
57	2a	121	C
57	2a	131	C

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Mol	Chain	Res	Type
57	2a	144	G
57	2a	163	C
57	2a	174	C
57	2a	182	U
57	2a	189(F)	U
57	2a	195	A
57	2a	197	A
57	2a	201	C
57	2a	202	U
57	2a	203	U
57	2a	204	U
57	2a	216	G
57	2a	247	G
57	2a	251	G
57	2a	258	G
57	2a	266	G
57	2a	267	C
57	2a	281	G
57	2a	289	G
57	2a	300	A
57	2a	321	A
57	2a	328	C
57	2a	332	G
57	2a	342	C
57	2a	351	G
57	2a	352	C
57	2a	353	A
57	2a	354	G
57	2a	367	U
57	2a	372	C
57	2a	384	G
57	2a	397	A
57	2a	398	C
57	2a	406	G
57	2a	412	A
57	2a	423	G
57	2a	429	U
57	2a	430	A
57	2a	439	A
57	2a	442	C
57	2a	452	A
57	2a	461	A

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Mol	Chain	Res	Type
57	2a	470	C
57	2a	477	A
57	2a	482	A
57	2a	484	G
57	2a	485	G
57	2a	496	A
57	2a	498	U
57	2a	499	A
57	2a	505	G
57	2a	510	A
57	2a	511	C
57	2a	518	C
57	2a	521	G
57	2a	531	U
57	2a	532	A
57	2a	533	A
57	2a	547	A
57	2a	559	A
57	2a	564	C
57	2a	568	G
57	2a	572	A
57	2a	573	A
57	2a	576	G
57	2a	577	G
57	2a	596	C
57	2a	630	G
57	2a	653	A
57	2a	665	A
57	2a	666	G
57	2a	687	A
57	2a	688	G
57	2a	695	A
57	2a	721	G
57	2a	723	U
57	2a	724	G
57	2a	731	G
57	2a	749	C
57	2a	755	G
57	2a	777	A
57	2a	792	A
57	2a	793	U
57	2a	794	A

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Mol	Chain	Res	Type
57	2a	816	A
57	2a	817	C
57	2a	821	G
57	2a	828	A
57	2a	840	C
57	2a	841	U
57	2a	853	G
57	2a	859	A
57	2a	874	G
57	2a	885	G
57	2a	902	G
57	2a	914	A
57	2a	926	G
57	2a	927	G
57	2a	934	C
57	2a	935	A
57	2a	942	G
57	2a	960	U
57	2a	961	U
57	2a	968	A
57	2a	969	A
57	2a	971	G
57	2a	972	C
57	2a	974	A
57	2a	975	A
57	2a	976	G
57	2a	977	A
57	2a	982	U
57	2a	984	C
57	2a	989	C
57	2a	992	U
57	2a	993	G
57	2a	997	U
57	2a	998	G
57	2a	999	C
57	2a	1001	A
57	2a	1001(A)	G
57	2a	1002	G
57	2a	1003	G
57	2a	1004	A
57	2a	1005	A
57	2a	1006	C

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Mol	Chain	Res	Type
57	2a	1009	G
57	2a	1011	G
57	2a	1016	A
57	2a	1020	U
57	2a	1021	G
57	2a	1022	G
57	2a	1025	U
57	2a	1026	G
57	2a	1027	C
57	2a	1029	C
57	2a	1030	C
57	2a	1030(A)	G
57	2a	1031	G
57	2a	1032	G
57	2a	1035	A
57	2a	1038	C
57	2a	1039	C
57	2a	1040	U
57	2a	1045	C
57	2a	1046	A
57	2a	1051	C
57	2a	1053	G
57	2a	1054	C
57	2a	1055	A
57	2a	1065	U
57	2a	1066	C
57	2a	1068	G
57	2a	1077	G
57	2a	1078	U
57	2a	1079	G
57	2a	1081	G
57	2a	1086	U
57	2a	1094	G
57	2a	1095	U
57	2a	1101	A
57	2a	1108	G
57	2a	1113	C
57	2a	1117	G
57	2a	1122	U
57	2a	1125	U
57	2a	1129	C
57	2a	1130	A

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Mol	Chain	Res	Type
57	2a	1132	C
57	2a	1136	U
57	2a	1137	C
57	2a	1138	G
57	2a	1139	G
57	2a	1146	A
57	2a	1147	C
57	2a	1152	A
57	2a	1157	A
57	2a	1158	C
57	2a	1159	U
57	2a	1160	G
57	2a	1182	G
57	2a	1183	A
57	2a	1184	G
57	2a	1193	G
57	2a	1196	U
57	2a	1197	G
57	2a	1202	G
57	2a	1211	U
57	2a	1213	A
57	2a	1227	A
57	2a	1236	A
57	2a	1238	A
57	2a	1240	U
57	2a	1241	G
57	2a	1256	A
57	2a	1257	U
57	2a	1258	G
57	2a	1260	C
57	2a	1261	A
57	2a	1264	C
57	2a	1267	C
57	2a	1268	A
57	2a	1270	C
57	2a	1272	G
57	2a	1273	G
57	2a	1276	G
57	2a	1277	C
57	2a	1278	U
57	2a	1279	A
57	2a	1280	A

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Mol	Chain	Res	Type
57	2a	1286	A
57	2a	1287	A
57	2a	1302	U
57	2a	1303	C
57	2a	1305	G
57	2a	1323	G
57	2a	1338	G
57	2a	1340	A
57	2a	1346	A
57	2a	1347	G
57	2a	1358	U
57	2a	1359	C
57	2a	1363	C
57	2a	1368	G
57	2a	1370	G
57	2a	1378	C
57	2a	1419	G
57	2a	1442	G
57	2a	1442(A)	G
57	2a	1447	A
57	2a	1452	C
57	2a	1456	G
57	2a	1492	A
57	2a	1497	G
57	2a	1504	G
57	2a	1506	U
57	2a	1517	G
57	2a	1520	G
57	2a	1529	G
57	2a	1530	G
57	2a	1531	A
57	2a	1532	U
53	2v	13	A
54	2w	3	C
54	2w	4	C
54	2w	5	G
54	2w	6	G
54	2w	8	4SU
54	2w	11	C
54	2w	14	A
54	2w	19	G
54	2w	22	G

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Mol	Chain	Res	Type
54	2w	27	G
54	2w	46	G7M
54	2w	47	U
54	2w	48	C
54	2w	50	U
54	2w	62	C
54	2w	64	A
54	2w	65	G
54	2w	68	C
54	2w	69	G
54	2w	70	G
54	2w	73	A
54	2w	74	C
55	2x	9	G
55	2x	13	C
55	2x	18	G
55	2x	21	A
55	2x	22	G
55	2x	46	G
55	2x	47	U
55	2x	48	C
55	2x	51	C
55	2x	52	G
55	2x	53	G
55	2x	67	C
55	2x	68	C
55	2x	76	A
56	2y	5	G
56	2y	8	4SU
56	2y	13	C
56	2y	19	G
56	2y	35	A
56	2y	37	A
56	2y	44	G
56	2y	45	U
56	2y	46	G7M
56	2y	48	C
56	2y	49	C
56	2y	53	G
56	2y	54	5MU
56	2y	59	U
56	2y	65	G

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Mol	Chain	Res	Type
56	2y	66	U
56	2y	69	G
56	2y	70	G

All (64) RNA pucker outliers are listed below:

Mol	Chain	Res	Type
1	1A	115	G
1	1A	185	A
1	1A	271	U
1	1A	302	A
1	1A	509	A
1	1A	572	A
1	1A	596	G
1	1A	716	G
1	1A	732	A
1	1A	793	A
1	1A	811	A
1	1A	821	A
1	1A	913	A
1	1A	941	U
1	1A	1003	U
1	1A	1019	G
1	1A	1065	U
1	1A	1067	A
1	1A	1093	G
1	1A	1201	A
1	1A	1219	A
1	1A	1221	G
1	1A	1255	A
1	1A	1425	A
1	1A	1554	A
1	1A	1654	A
1	1A	1700	G
1	1A	1793	A
1	1A	2014	G
1	1A	2156	A
1	1A	2203	G
1	1A	2205	C
1	1A	2418	U
1	1A	2442	A
1	1A	2451	A

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Mol	Chain	Res	Type
1	1A	2641	A
1	1A	2701	U
1	1A	2769	U
2	1B	1	U
1	2A	34	C
1	2A	196	A
1	2A	228	A
1	2A	266	G
1	2A	271(K)	U
1	2A	271(M)	G
1	2A	277	C
1	2A	528	A
1	2A	752	A
1	2A	774	A
1	2A	827	U
1	2A	856	C
1	2A	900	A
1	2A	958	U
1	2A	1210	A
1	2A	1420	U
1	2A	1442	G
1	2A	1530	C
1	2A	1608	A
1	2A	1653	G
1	2A	1913	A
1	2A	1992	G
1	2A	2119	A
1	2A	2126	A
1	2A	2689	U

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

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

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

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
1	PSU	1A	1933	1	16,21,22	1.32	2 (12%)	20,30,33	3.45	6 (30%)
1	5MU	1A	1937	1	14,22,23	0.71	0	16,32,35	2.02	3 (18%)
1	PSU	1A	1939	1	16,21,22	1.46	2 (12%)	20,30,33	3.62	6 (30%)
1	4OC	1A	1942	1	15,22,24	0.79	0	19,31,35	0.84	0
1	5MU	1A	1961	1	14,22,23	0.86	1 (7%)	16,32,35	2.06	3 (18%)
1	5MC	1A	1964	1,59	15,22,23	1.33	1 (6%)	17,32,35	1.16	2 (11%)
1	5MC	1A	1984	1,59	15,22,23	1.38	1 (6%)	17,32,35	1.02	2 (11%)
1	OMG	1A	2263	1,55,59	18,26,27	1.31	3 (16%)	22,38,41	2.13	6 (27%)
1	2MA	1A	2515	1,59	18,25,26	1.59	4 (22%)	17,37,40	1.96	2 (11%)
1	2MU	1A	2564	1,59	14,22,24	0.91	1 (7%)	18,31,36	1.98	2 (11%)
1	PSU	1A	2617	1,59	16,21,22	1.70	3 (18%)	20,30,33	3.61	6 (30%)
32	2MG	1a	1207	32	19,26,27	1.28	2 (10%)	20,38,41	2.48	8 (40%)
32	5MC	1a	1400	32	15,22,23	1.38	1 (6%)	17,32,35	1.07	2 (11%)
32	4OC	1a	1402	32	16,23,24	0.77	0	19,32,35	1.25	1 (5%)
32	5MC	1a	1404	32	15,22,23	1.47	1 (6%)	17,32,35	0.89	0
32	5MC	1a	1407	32	15,22,23	1.36	1 (6%)	17,32,35	1.08	2 (11%)
32	UR3	1a	1498	32	14,22,23	0.93	1 (7%)	16,32,35	0.82	1 (6%)
32	MA6	1a	1518	32	16,26,27	1.00	1 (6%)	18,38,41	2.29	5 (27%)
32	MA6	1a	1519	32	16,26,27	1.03	1 (6%)	18,38,41	2.20	5 (27%)
32	PSU	1a	516	32,59	16,21,22	1.38	2 (12%)	20,30,33	3.48	6 (30%)
32	G7M	1a	527	32,59	19,26,27	2.67	3 (15%)	19,39,42	2.08	5 (26%)
32	M2G	1a	966	32	20,27,28	1.45	3 (15%)	21,40,43	2.22	6 (28%)
32	5MC	1a	967	32	15,22,23	1.35	1 (6%)	17,32,35	1.15	2 (11%)
43	0TD	1l	92	43	5,9,10	2.86	2 (40%)	3,11,13	2.24	1 (33%)
54	PSU	1w	32	54,59	16,21,22	1.28	1 (6%)	20,30,33	3.62	6 (30%)
54	MIA	1w	37	54	23,31,32	1.74	2 (8%)	25,44,47	1.50	5 (20%)
54	PSU	1w	39	54	16,21,22	1.45	1 (6%)	20,30,33	3.62	7 (35%)
54	G7M	1w	46	54	19,26,27	2.57	3 (15%)	19,39,42	2.49	5 (26%)
54	5MU	1w	54	54	14,22,23	0.77	0	16,32,35	2.24	3 (18%)
54	PSU	1w	55	54	16,21,22	1.19	1 (6%)	20,30,33	3.79	6 (30%)
54	4SU	1w	8	54	14,21,22	1.24	1 (7%)	15,30,33	1.62	2 (13%)
55	5MC	1x	32	55	15,22,23	1.41	1 (6%)	17,32,35	1.24	2 (11%)
55	5MU	1x	54	55,59	14,22,23	0.79	0	16,32,35	2.35	3 (18%)
55	PSU	1x	55	55,59	16,21,22	1.57	2 (12%)	20,30,33	3.56	6 (30%)
55	4SU	1x	8	55	14,21,22	1.39	2 (14%)	15,30,33	2.66	2 (13%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
56	PSU	1y	32	56	16,21,22	1.21	1 (6%)	20,30,33	3.55	5 (25%)
56	PSU	1y	39	56	16,21,22	1.29	1 (6%)	20,30,33	3.49	6 (30%)
56	G7M	1y	46	56	19,26,27	2.59	3 (15%)	19,39,42	2.38	6 (31%)
56	5MU	1y	54	56	14,22,23	0.76	1 (7%)	16,32,35	2.57	3 (18%)
56	PSU	1y	55	56	16,21,22	1.34	1 (6%)	20,30,33	3.60	7 (35%)
56	4SU	1y	8	56	14,21,22	1.20	1 (7%)	15,30,33	1.63	2 (13%)
1	PSU	2A	1911	1	16,21,22	1.32	1 (6%)	20,30,33	3.54	7 (35%)
1	5MU	2A	1915	1	14,22,23	0.75	0	16,32,35	2.13	3 (18%)
1	PSU	2A	1917	1	16,21,22	1.48	1 (6%)	20,30,33	3.68	7 (35%)
1	4OC	2A	1920	1	15,22,24	0.72	0	19,31,35	0.82	0
1	5MU	2A	1939	1,59	14,22,23	0.84	1 (7%)	16,32,35	2.08	3 (18%)
1	5MC	2A	1942	1	15,22,23	1.39	1 (6%)	17,32,35	1.07	1 (5%)
1	5MC	2A	1962	1,59	15,22,23	1.46	1 (6%)	17,32,35	1.14	2 (11%)
1	OMG	2A	2251	1,55,59	18,26,27	1.26	2 (11%)	22,38,41	1.96	6 (27%)
1	2MA	2A	2503	1,59	18,25,26	1.48	4 (22%)	17,37,40	1.74	2 (11%)
1	2MU	2A	2552	1,59	14,22,24	0.98	1 (7%)	18,31,36	1.78	1 (5%)
1	PSU	2A	2605	1	16,21,22	1.51	2 (12%)	20,30,33	3.47	6 (30%)
57	2MG	2a	1207	57	19,26,27	1.29	2 (10%)	20,38,41	2.33	7 (35%)
57	5MC	2a	1400	57	15,22,23	1.48	1 (6%)	17,32,35	1.14	2 (11%)
57	4OC	2a	1402	57	16,23,24	0.71	0	19,32,35	1.33	1 (5%)
57	5MC	2a	1404	57	15,22,23	1.45	1 (6%)	17,32,35	1.06	2 (11%)
57	5MC	2a	1407	57	15,22,23	1.36	1 (6%)	17,32,35	1.37	3 (17%)
57	UR3	2a	1498	57	14,22,23	0.86	1 (7%)	16,32,35	0.72	1 (6%)
57	MA6	2a	1518	57	16,26,27	1.04	1 (6%)	18,38,41	2.37	5 (27%)
57	MA6	2a	1519	57	16,26,27	1.06	1 (6%)	18,38,41	2.18	4 (22%)
57	PSU	2a	516	57	16,21,22	1.42	1 (6%)	20,30,33	3.49	7 (35%)
57	G7M	2a	527	57,59	19,26,27	2.61	3 (15%)	19,39,42	2.13	5 (26%)
57	M2G	2a	966	57	20,27,28	1.42	3 (15%)	21,40,43	2.12	6 (28%)
57	5MC	2a	967	57	15,22,23	1.47	1 (6%)	17,32,35	1.00	2 (11%)
43	0TD	2l	92	43	5,9,10	3.15	2 (40%)	3,11,13	3.92	1 (33%)
54	PSU	2w	32	54	16,21,22	1.40	1 (6%)	20,30,33	3.61	6 (30%)
54	MIA	2w	37	54	20,27,32	1.93	2 (10%)	21,39,47	1.70	5 (23%)
54	PSU	2w	39	54	16,21,22	1.33	1 (6%)	20,30,33	3.78	6 (30%)
54	G7M	2w	46	54	19,26,27	2.66	3 (15%)	19,39,42	2.21	6 (31%)
54	5MU	2w	54	54	14,22,23	0.70	0	16,32,35	2.68	2 (12%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
54	PSU	2w	55	54	16,21,22	1.15	1 (6%)	20,30,33	3.63	6 (30%)
54	4SU	2w	8	54	14,21,22	1.24	1 (7%)	15,30,33	1.23	2 (13%)
55	5MC	2x	32	55	15,22,23	1.40	1 (6%)	17,32,35	1.17	2 (11%)
55	5MU	2x	54	55	14,22,23	0.76	0	16,32,35	2.22	3 (18%)
55	PSU	2x	55	55	16,21,22	1.44	1 (6%)	20,30,33	3.54	7 (35%)
55	4SU	2x	8	55,59	14,21,22	1.35	2 (14%)	15,30,33	2.42	2 (13%)
56	PSU	2y	32	56	16,21,22	1.13	1 (6%)	20,30,33	3.60	5 (25%)
56	PSU	2y	39	56	16,21,22	1.31	1 (6%)	20,30,33	3.62	6 (30%)
56	G7M	2y	46	56	19,26,27	2.61	3 (15%)	19,39,42	2.59	6 (31%)
56	5MU	2y	54	56	14,22,23	0.70	0	16,32,35	2.43	2 (12%)
56	PSU	2y	55	56	16,21,22	1.26	1 (6%)	20,30,33	3.52	7 (35%)
56	4SU	2y	8	56,59	14,21,22	1.26	1 (7%)	15,30,33	1.50	2 (13%)
58	BB9	A	17	58	3,5,6	3.74	1 (33%)	1,5,7	4.27	1 (100%)
58	BB9	A	6	58	3,5,6	3.46	1 (33%)	1,5,7	2.57	1 (100%)
58	BB9	A	9	58	3,5,6	3.76	1 (33%)	1,5,7	2.58	1 (100%)
58	BB9	B	17	58	3,5,6	3.50	1 (33%)	1,5,7	4.12	1 (100%)
58	BB9	B	6	58	3,5,6	3.33	1 (33%)	1,5,7	2.76	1 (100%)
58	BB9	B	9	58	3,5,6	4.03	1 (33%)	1,5,7	2.90	1 (100%)

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
1	PSU	1A	1933	1	-	0/7/25/26	0/2/2/2
1	5MU	1A	1937	1	-	0/3/25/26	0/2/2/2
1	PSU	1A	1939	1	-	0/7/25/26	0/2/2/2
1	4OC	1A	1942	1	-	0/5/27/30	0/2/2/2
1	5MU	1A	1961	1	-	0/3/25/26	0/2/2/2
1	5MC	1A	1964	1,59	-	0/3/25/26	0/2/2/2
1	5MC	1A	1984	1,59	-	0/3/25/26	0/2/2/2
1	OMG	1A	2263	1,55,59	-	0/5/27/28	0/3/3/3
1	2MA	1A	2515	1,59	-	0/3/25/26	0/3/3/3
1	2MU	1A	2564	1,59	-	0/5/27/28	0/2/2/2
1	PSU	1A	2617	1,59	-	0/7/25/26	0/2/2/2
32	2MG	1a	1207	32	-	0/5/27/28	0/3/3/3
32	5MC	1a	1400	32	-	0/3/25/26	0/2/2/2

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
32	4OC	1a	1402	32	-	0/7/29/30	0/2/2/2
32	5MC	1a	1404	32	-	0/3/25/26	0/2/2/2
32	5MC	1a	1407	32	-	0/3/25/26	0/2/2/2
32	UR3	1a	1498	32	-	0/3/25/26	0/2/2/2
32	MA6	1a	1518	32	-	0/7/29/30	0/3/3/3
32	MA6	1a	1519	32	-	0/7/29/30	0/3/3/3
32	PSU	1a	516	32,59	-	0/7/25/26	0/2/2/2
32	G7M	1a	527	32,59	-	0/3/25/26	0/3/3/3
32	M2G	1a	966	32	-	0/7/29/30	0/3/3/3
32	5MC	1a	967	32	-	0/3/25/26	0/2/2/2
43	0TD	1l	92	43	-	0/2/12/14	0/0/0/0
54	PSU	1w	32	54,59	-	0/7/25/26	0/2/2/2
54	MIA	1w	37	54	-	0/11/33/34	0/3/3/3
54	PSU	1w	39	54	-	0/7/25/26	0/2/2/2
54	G7M	1w	46	54	-	0/3/25/26	0/3/3/3
54	5MU	1w	54	54	-	0/3/25/26	0/2/2/2
54	PSU	1w	55	54	-	0/7/25/26	0/2/2/2
54	4SU	1w	8	54	-	0/3/25/26	0/2/2/2
55	5MC	1x	32	55	-	0/3/25/26	0/2/2/2
55	5MU	1x	54	55,59	-	0/3/25/26	0/2/2/2
55	PSU	1x	55	55,59	-	0/7/25/26	0/2/2/2
55	4SU	1x	8	55	-	0/3/25/26	0/2/2/2
56	PSU	1y	32	56	-	0/7/25/26	0/2/2/2
56	PSU	1y	39	56	-	0/7/25/26	0/2/2/2
56	G7M	1y	46	56	-	0/3/25/26	0/3/3/3
56	5MU	1y	54	56	-	0/3/25/26	0/2/2/2
56	PSU	1y	55	56	-	0/7/25/26	0/2/2/2
56	4SU	1y	8	56	-	0/3/25/26	0/2/2/2
1	PSU	2A	1911	1	-	0/7/25/26	0/2/2/2
1	5MU	2A	1915	1	-	0/3/25/26	0/2/2/2
1	PSU	2A	1917	1	-	0/7/25/26	0/2/2/2
1	4OC	2A	1920	1	-	0/5/27/30	0/2/2/2
1	5MU	2A	1939	1,59	-	0/3/25/26	0/2/2/2
1	5MC	2A	1942	1	-	0/3/25/26	0/2/2/2
1	5MC	2A	1962	1,59	-	0/3/25/26	0/2/2/2
1	OMG	2A	2251	1,55,59	-	0/5/27/28	0/3/3/3
1	2MA	2A	2503	1,59	-	0/3/25/26	0/3/3/3
1	2MU	2A	2552	1,59	-	0/5/27/28	0/2/2/2
1	PSU	2A	2605	1	-	0/7/25/26	0/2/2/2
57	2MG	2a	1207	57	-	0/5/27/28	0/3/3/3
57	5MC	2a	1400	57	-	0/3/25/26	0/2/2/2
57	4OC	2a	1402	57	-	0/7/29/30	0/2/2/2

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
57	5MC	2a	1404	57	-	0/3/25/26	0/2/2/2
57	5MC	2a	1407	57	-	0/3/25/26	0/2/2/2
57	UR3	2a	1498	57	-	0/3/25/26	0/2/2/2
57	MA6	2a	1518	57	-	0/7/29/30	0/3/3/3
57	MA6	2a	1519	57	-	0/7/29/30	0/3/3/3
57	PSU	2a	516	57	-	0/7/25/26	0/2/2/2
57	G7M	2a	527	57,59	-	0/3/25/26	0/3/3/3
57	M2G	2a	966	57	-	0/7/29/30	0/3/3/3
57	5MC	2a	967	57	-	0/3/25/26	0/2/2/2
43	0TD	2l	92	43	-	0/2/12/14	0/0/0/0
54	PSU	2w	32	54	-	0/7/25/26	0/2/2/2
54	MIA	2w	37	54	-	0/7/29/34	0/3/3/3
54	PSU	2w	39	54	-	0/7/25/26	0/2/2/2
54	G7M	2w	46	54	-	0/3/25/26	0/3/3/3
54	5MU	2w	54	54	-	0/3/25/26	0/2/2/2
54	PSU	2w	55	54	-	0/7/25/26	0/2/2/2
54	4SU	2w	8	54	-	0/3/25/26	0/2/2/2
55	5MC	2x	32	55	-	0/3/25/26	0/2/2/2
55	5MU	2x	54	55	-	0/3/25/26	0/2/2/2
55	PSU	2x	55	55	-	0/7/25/26	0/2/2/2
55	4SU	2x	8	55,59	-	0/3/25/26	0/2/2/2
56	PSU	2y	32	56	-	0/7/25/26	0/2/2/2
56	PSU	2y	39	56	-	0/7/25/26	0/2/2/2
56	G7M	2y	46	56	-	0/3/25/26	0/3/3/3
56	5MU	2y	54	56	-	0/3/25/26	0/2/2/2
56	PSU	2y	55	56	-	0/7/25/26	0/2/2/2
56	4SU	2y	8	56,59	-	0/3/25/26	0/2/2/2
58	BB9	A	17	58	-	0/0/4/6	0/0/0/0
58	BB9	A	6	58	-	0/0/4/6	0/0/0/0
58	BB9	A	9	58	-	0/0/4/6	0/0/0/0
58	BB9	B	17	58	-	0/0/4/6	0/0/0/0
58	BB9	B	6	58	-	0/0/4/6	0/0/0/0
58	BB9	B	9	58	-	0/0/4/6	0/0/0/0

All (117) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
54	2w	37	MIA	C2-S10	-7.10	1.69	1.75
58	B	9	BB9	C-CA	-6.95	1.33	1.45
54	1w	37	MIA	C2-S10	-6.81	1.70	1.75
58	A	9	BB9	C-CA	-6.48	1.34	1.45
58	A	17	BB9	C-CA	-6.41	1.34	1.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
58	B	17	BB9	C-CA	-6.00	1.35	1.45
58	A	6	BB9	C-CA	-5.95	1.35	1.45
43	2l	92	0TD	CB-SB	-5.89	1.69	1.84
43	1l	92	0TD	CB-SB	-5.74	1.69	1.84
58	B	6	BB9	C-CA	-5.73	1.35	1.45
1	1A	2617	PSU	C5-C1'	-5.20	1.47	1.52
55	1x	55	PSU	C5-C1'	-4.88	1.48	1.52
1	2A	1917	PSU	C5-C1'	-4.52	1.48	1.52
55	2x	55	PSU	C5-C1'	-4.51	1.48	1.52
1	2A	2605	PSU	C5-C1'	-4.50	1.48	1.52
57	2a	516	PSU	C5-C1'	-4.32	1.48	1.52
54	1w	39	PSU	C5-C1'	-4.30	1.48	1.52
54	2w	32	PSU	C5-C1'	-4.28	1.48	1.52
1	1A	1939	PSU	C5-C1'	-4.23	1.48	1.52
56	1y	55	PSU	C5-C1'	-4.00	1.48	1.52
56	2y	39	PSU	C5-C1'	-3.98	1.48	1.52
32	1a	516	PSU	C5-C1'	-3.94	1.48	1.52
56	2y	55	PSU	C5-C1'	-3.86	1.48	1.52
54	2w	8	4SU	C4-S4	-3.80	1.60	1.67
56	1y	39	PSU	C5-C1'	-3.79	1.49	1.52
56	2y	8	4SU	C4-S4	-3.76	1.60	1.67
54	2w	39	PSU	C5-C1'	-3.76	1.49	1.52
1	2A	1911	PSU	C5-C1'	-3.74	1.49	1.52
54	1w	32	PSU	C5-C1'	-3.68	1.49	1.52
54	1w	8	4SU	C4-S4	-3.68	1.60	1.67
55	1x	8	4SU	C4-S4	-3.49	1.60	1.67
55	2x	8	4SU	C4-S4	-3.46	1.60	1.67
56	1y	8	4SU	C4-S4	-3.46	1.60	1.67
55	1x	8	4SU	C2-N3	-3.45	1.31	1.38
1	1A	1933	PSU	C5-C1'	-3.40	1.49	1.52
56	1y	32	PSU	C5-C1'	-3.37	1.49	1.52
55	2x	8	4SU	C2-N3	-3.13	1.32	1.38
54	2w	55	PSU	C5-C1'	-3.00	1.49	1.52
54	1w	55	PSU	C5-C1'	-2.98	1.49	1.52
56	2y	32	PSU	C5-C1'	-2.85	1.49	1.52
1	1A	1961	5MU	C2-N3	-2.47	1.33	1.38
1	1A	2617	PSU	C2-N1	-2.43	1.33	1.38
1	2A	1939	5MU	C2-N3	-2.31	1.33	1.38
1	1A	2564	2MU	C2-N3	-2.20	1.33	1.38
1	1A	2515	2MA	O5'-C5'	-2.15	1.41	1.44
1	2A	2503	2MA	O5'-C5'	-2.12	1.41	1.44
56	1y	54	5MU	C2-N3	-2.11	1.34	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
1	2A	2605	PSU	O5'-C5'	-2.10	1.41	1.44
1	1A	1933	PSU	O4'-C1'	-2.10	1.41	1.44
1	1A	2263	OMG	O5'-C5'	-2.10	1.41	1.44
32	1a	516	PSU	C2-N3	-2.07	1.34	1.38
55	1x	55	PSU	O5'-C5'	-2.02	1.41	1.44
1	1A	1939	PSU	C2-N1	-2.02	1.34	1.38
1	2A	2552	2MU	O5'-C5'	-2.01	1.42	1.44
1	1A	2617	PSU	C2-N3	-2.01	1.34	1.38
57	2a	1498	UR3	C4-N3	2.00	1.41	1.38
43	1l	92	0TD	CA-C	2.26	1.53	1.50
1	1A	2515	2MA	C5-C4	2.62	1.46	1.40
32	1a	1498	UR3	C4-N3	2.64	1.42	1.38
1	2A	2503	2MA	C5-C4	2.74	1.46	1.40
32	1a	1519	MA6	C5-C4	2.89	1.47	1.40
32	1a	966	M2G	C5-C4	2.90	1.47	1.40
32	1a	1207	2MG	C5-C4	2.93	1.47	1.40
54	1w	46	G7M	C6-C5	3.04	1.47	1.41
1	2A	2251	OMG	C5-C4	3.09	1.47	1.40
1	2A	2503	2MA	C6-N6	3.09	1.34	1.27
32	1a	1518	MA6	C5-C4	3.10	1.47	1.40
57	2a	1207	2MG	C5-C4	3.12	1.47	1.40
1	1A	2263	OMG	C5-C4	3.15	1.47	1.40
54	1w	37	MIA	C5-C4	3.20	1.47	1.40
57	2a	1519	MA6	C5-C4	3.22	1.47	1.40
57	2a	966	M2G	C5-C4	3.22	1.47	1.40
54	2w	37	MIA	C5-C4	3.31	1.48	1.40
57	2a	966	M2G	C2-N2	3.33	1.40	1.34
57	2a	1518	MA6	C5-C4	3.35	1.48	1.40
56	2y	46	G7M	C6-C5	3.53	1.48	1.41
43	2l	92	0TD	CA-C	3.53	1.54	1.50
1	1A	2263	OMG	C6-C5	3.55	1.48	1.41
1	1A	2515	2MA	C6-N6	3.61	1.35	1.27
54	2w	46	G7M	C6-C5	3.64	1.48	1.41
1	2A	2251	OMG	C6-C5	3.65	1.48	1.41
32	1a	966	M2G	C6-C5	3.66	1.48	1.41
32	1a	527	G7M	C6-C5	3.66	1.48	1.41
57	2a	966	M2G	C6-C5	3.68	1.48	1.41
32	1a	966	M2G	C2-N2	3.70	1.40	1.34
57	2a	527	G7M	C6-C5	3.73	1.48	1.41
56	1y	46	G7M	C6-C5	3.74	1.48	1.41
1	2A	2503	2MA	C6-C5	3.78	1.47	1.41
32	1a	1207	2MG	C6-C5	3.89	1.48	1.41

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
57	2a	1207	2MG	C6-C5	4.07	1.49	1.41
1	1A	2515	2MA	C6-C5	4.20	1.48	1.41
1	1A	1964	5MC	C5-C4	4.45	1.47	1.41
57	2a	1407	5MC	C5-C4	4.51	1.48	1.41
32	1a	1407	5MC	C5-C4	4.61	1.48	1.41
32	1a	967	5MC	C5-C4	4.77	1.48	1.41
32	1a	1400	5MC	C5-C4	4.85	1.48	1.41
55	2x	32	5MC	C5-C4	4.86	1.48	1.41
1	1A	1984	5MC	C5-C4	4.87	1.48	1.41
55	1x	32	5MC	C5-C4	4.87	1.48	1.41
1	2A	1942	5MC	C5-C4	4.90	1.48	1.41
32	1a	1404	5MC	C5-C4	5.03	1.48	1.41
57	2a	1404	5MC	C5-C4	5.08	1.48	1.41
1	2A	1962	5MC	C5-C4	5.11	1.48	1.41
57	2a	1400	5MC	C5-C4	5.12	1.48	1.41
57	2a	967	5MC	C5-C4	5.18	1.49	1.41
56	2y	46	G7M	C8-N7	7.06	1.46	1.33
56	1y	46	G7M	C8-N7	7.08	1.46	1.33
54	1w	46	G7M	C8-N7	7.16	1.46	1.33
57	2a	527	G7M	C8-N7	7.20	1.46	1.33
32	1a	527	G7M	C8-N7	7.32	1.46	1.33
54	2w	46	G7M	C8-N7	7.50	1.46	1.33
54	2w	46	G7M	C8-N9	7.58	1.47	1.33
57	2a	527	G7M	C8-N9	7.65	1.47	1.33
56	1y	46	G7M	C8-N9	7.65	1.47	1.33
54	1w	46	G7M	C8-N9	7.74	1.47	1.33
56	2y	46	G7M	C8-N9	7.74	1.47	1.33
32	1a	527	G7M	C8-N9	7.91	1.47	1.33

All (323) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
54	2w	39	PSU	N1-C2-N3	-10.38	120.93	128.40
54	1w	39	PSU	N1-C2-N3	-9.85	121.31	128.40
56	2y	32	PSU	N1-C2-N3	-9.80	121.35	128.40
32	1a	516	PSU	N1-C2-N3	-9.75	121.39	128.40
54	1w	32	PSU	N1-C2-N3	-9.75	121.39	128.40
54	1w	55	PSU	N1-C2-N3	-9.73	121.40	128.40
56	1y	55	PSU	C5-C4-N3	-9.68	117.49	125.43
56	2y	39	PSU	N1-C2-N3	-9.56	121.52	128.40
54	2w	32	PSU	N1-C2-N3	-9.54	121.54	128.40
1	1A	1939	PSU	N1-C2-N3	-9.49	121.57	128.40

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
55	1x	55	PSU	N1-C2-N3	-9.43	121.62	128.40
1	1A	2617	PSU	N1-C2-N3	-9.33	121.69	128.40
1	2A	1917	PSU	C5-C4-N3	-9.32	117.78	125.43
1	2A	1911	PSU	N1-C2-N3	-9.32	121.70	128.40
1	2A	2605	PSU	N1-C2-N3	-9.32	121.70	128.40
56	1y	39	PSU	N1-C2-N3	-9.29	121.72	128.40
56	1y	32	PSU	N1-C2-N3	-9.27	121.73	128.40
57	2a	516	PSU	N1-C2-N3	-9.26	121.74	128.40
54	2w	55	PSU	N1-C2-N3	-9.19	121.79	128.40
56	2y	55	PSU	N1-C2-N3	-9.13	121.83	128.40
55	2x	55	PSU	N1-C2-N3	-9.10	121.85	128.40
1	1A	1933	PSU	N1-C2-N3	-9.09	121.86	128.40
1	2A	1917	PSU	N1-C2-N3	-8.97	121.95	128.40
54	1w	55	PSU	C5-C4-N3	-8.78	118.23	125.43
56	1y	32	PSU	C5-C4-N3	-8.77	118.24	125.43
56	1y	55	PSU	N1-C2-N3	-8.71	122.13	128.40
55	2x	55	PSU	C5-C4-N3	-8.70	118.29	125.43
54	2w	39	PSU	C5-C4-N3	-8.63	118.35	125.43
56	2y	32	PSU	C5-C4-N3	-8.61	118.37	125.43
56	2y	55	PSU	C5-C4-N3	-8.61	118.37	125.43
1	1A	2617	PSU	C5-C4-N3	-8.53	118.43	125.43
54	1w	32	PSU	C5-C4-N3	-8.50	118.45	125.43
54	1w	39	PSU	C5-C4-N3	-8.46	118.49	125.43
54	2w	32	PSU	C5-C4-N3	-8.43	118.52	125.43
1	1A	1939	PSU	C5-C4-N3	-8.42	118.52	125.43
57	2a	516	PSU	C5-C4-N3	-8.33	118.59	125.43
55	1x	55	PSU	C5-C4-N3	-8.33	118.60	125.43
56	2y	39	PSU	C5-C4-N3	-8.28	118.64	125.43
1	1A	1933	PSU	C5-C4-N3	-8.25	118.66	125.43
1	2A	1911	PSU	C5-C4-N3	-8.20	118.70	125.43
56	1y	39	PSU	C5-C4-N3	-8.13	118.76	125.43
54	2w	55	PSU	C5-C4-N3	-8.06	118.82	125.43
32	1a	516	PSU	C5-C4-N3	-7.72	119.10	125.43
1	2A	2605	PSU	C5-C4-N3	-7.57	119.22	125.43
54	1w	46	G7M	C5-C6-N1	-6.54	114.17	123.48
43	2l	92	0TD	CSB-SB-CB	-6.43	89.59	101.60
54	2w	54	5MU	C5-C4-N3	-6.30	118.30	125.24
57	2a	1518	MA6	N3-C2-N1	-6.28	123.39	128.86
56	1y	54	5MU	C5-C4-N3	-6.14	118.47	125.24
56	2y	46	G7M	C5-C6-N1	-6.01	114.92	123.48
32	1a	1518	MA6	N3-C2-N1	-5.95	123.67	128.86
56	1y	46	G7M	C5-C6-N1	-5.82	115.20	123.48

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
32	1a	527	G7M	C5-C6-N1	-5.74	115.31	123.48
55	1x	54	5MU	C5-C4-N3	-5.70	118.96	125.24
55	2x	54	5MU	C5-C4-N3	-5.64	119.02	125.24
32	1a	1519	MA6	N3-C2-N1	-5.63	123.95	128.86
57	2a	1519	MA6	N3-C2-N1	-5.62	123.96	128.86
54	2w	46	G7M	C5-C6-N1	-5.57	115.56	123.48
54	1w	54	5MU	C5-C4-N3	-5.53	119.14	125.24
56	2y	54	5MU	C5-C4-N3	-5.48	119.20	125.24
57	2a	527	G7M	C5-C6-N1	-5.34	115.88	123.48
54	2w	55	PSU	C5-C1'-C2'	-5.29	106.42	115.55
1	2A	1915	5MU	C5-C4-N3	-5.19	119.52	125.24
1	1A	1937	5MU	C5-C4-N3	-4.85	119.90	125.24
54	2w	37	MIA	C12-N6-C6	-4.78	118.76	122.85
1	2A	1939	5MU	C5-C4-N3	-4.78	119.97	125.24
54	1w	55	PSU	C5-C1'-C2'	-4.65	107.53	115.55
55	1x	8	4SU	C5-C4-N3	-4.61	117.91	123.73
55	2x	8	4SU	C5-C4-N3	-4.57	117.96	123.73
1	1A	1961	5MU	C5-C4-N3	-4.56	120.21	125.24
32	1a	1402	4OC	CM4-N4-C4	-4.54	119.02	122.94
57	2a	1402	4OC	CM4-N4-C4	-4.52	119.04	122.94
1	2A	2605	PSU	C5-C6-N1	-4.46	118.61	124.39
1	1A	1939	PSU	C5-C1'-C2'	-4.44	107.89	115.55
56	2y	39	PSU	C5-C1'-C2'	-4.34	108.06	115.55
58	A	17	BB9	O-C-CA	-4.27	120.01	125.47
32	1a	1207	2MG	C5-C6-N1	-4.25	117.43	123.48
1	2A	1917	PSU	C5-C1'-C2'	-4.24	108.24	115.55
55	2x	55	PSU	C5-C6-N1	-4.23	118.91	124.39
57	2a	1207	2MG	C5-C6-N1	-4.15	117.57	123.48
54	2w	32	PSU	C5-C6-N1	-4.13	119.04	124.39
58	B	17	BB9	O-C-CA	-4.12	120.20	125.47
32	1a	966	M2G	C5-C6-N1	-4.09	117.67	123.48
32	1a	516	PSU	C5-C6-N1	-4.06	119.12	124.39
32	1a	1207	2MG	C4-C5-N7	-4.06	105.49	109.41
56	1y	39	PSU	C5-C6-N1	-4.03	119.16	124.39
1	1A	2263	OMG	C5-C6-N1	-4.01	117.77	123.48
55	1x	55	PSU	C5-C6-N1	-4.01	119.20	124.39
54	1w	37	MIA	C12-N6-C6	-4.00	118.11	123.26
1	2A	1939	5MU	C5-C6-N1	-3.96	117.87	122.15
1	1A	2617	PSU	C5-C6-N1	-3.92	119.31	124.39
1	1A	2617	PSU	C5-C1'-C2'	-3.89	108.84	115.55
57	2a	1518	MA6	C4-C5-N7	-3.87	105.67	109.41
57	2a	516	PSU	C5-C6-N1	-3.86	119.39	124.39

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
57	2a	966	M2G	C5-C6-N1	-3.85	118.00	123.48
57	2a	1207	2MG	C6-C5-C4	-3.83	117.03	120.84
57	2a	1519	MA6	C4-C5-N7	-3.82	105.72	109.41
54	1w	39	PSU	C5-C6-N1	-3.81	119.44	124.39
56	2y	55	PSU	C5-C6-N1	-3.81	119.45	124.39
54	2w	55	PSU	C5-C6-N1	-3.81	119.45	124.39
1	2A	1917	PSU	C5-C6-N1	-3.80	119.46	124.39
1	2A	1911	PSU	C5-C6-N1	-3.78	119.49	124.39
32	1a	966	M2G	C6-C5-C4	-3.78	117.09	120.84
1	2A	2251	OMG	C5-C6-N1	-3.75	118.14	123.48
56	1y	8	4SU	C5-C4-N3	-3.68	119.08	123.73
54	2w	39	PSU	C5-C6-N1	-3.68	119.62	124.39
56	2y	39	PSU	C5-C6-N1	-3.68	119.62	124.39
32	1a	1207	2MG	C6-C5-C4	-3.66	117.21	120.84
1	1A	1939	PSU	C5-C6-N1	-3.65	119.65	124.39
1	1A	1933	PSU	C5-C6-N1	-3.64	119.67	124.39
57	2a	1518	MA6	C9-N6-C6	-3.62	108.55	119.51
1	1A	1961	5MU	C5-C6-N1	-3.61	118.24	122.15
32	1a	1518	MA6	C4-C5-N7	-3.61	105.92	109.41
32	1a	1207	2MG	CM2-N2-C2	-3.58	119.28	123.63
1	2A	1911	PSU	C5-C1'-C2'	-3.57	109.39	115.55
55	1x	55	PSU	C5-C1'-C2'	-3.54	109.44	115.55
57	2a	1207	2MG	C4-C5-N7	-3.52	106.01	109.41
43	1l	92	0TD	CSB-SB-CB	-3.51	95.05	101.60
54	2w	32	PSU	C5-C1'-C2'	-3.50	109.51	115.55
57	2a	1519	MA6	C9-N6-C6	-3.50	108.92	119.51
56	1y	32	PSU	C5-C6-N1	-3.49	119.87	124.39
56	2y	46	G7M	N3-C2-N1	-3.48	122.38	127.46
54	1w	8	4SU	C5-C4-N3	-3.45	119.38	123.73
1	1A	2263	OMG	N3-C2-N1	-3.44	122.43	127.46
54	1w	32	PSU	C5-C6-N1	-3.44	119.93	124.39
1	1A	2263	OMG	C6-C5-C4	-3.44	117.43	120.84
32	1a	1519	MA6	C9-N6-C6	-3.38	109.29	119.51
32	1a	1519	MA6	C4-C5-N7	-3.37	106.16	109.41
56	1y	55	PSU	C5-C6-N1	-3.36	120.03	124.39
1	2A	2605	PSU	C5-C1'-C2'	-3.27	109.91	115.55
57	2a	1207	2MG	CM2-N2-C2	-3.18	119.76	123.63
54	1w	32	PSU	C5-C1'-C2'	-3.16	110.09	115.55
54	1w	46	G7M	N3-C2-N1	-3.13	122.88	127.46
1	2A	2251	OMG	C6-C5-C4	-3.12	117.74	120.84
56	2y	32	PSU	C5-C6-N1	-3.08	120.40	124.39
57	2a	966	M2G	C6-C5-C4	-3.08	117.78	120.84

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
32	1a	1518	MA6	C9-N6-C6	-3.07	110.23	119.51
56	1y	46	G7M	N3-C2-N1	-3.04	123.02	127.46
54	2w	37	MIA	C4-C5-N7	-2.99	106.52	109.41
56	2y	8	4SU	C5-C4-N3	-2.98	119.97	123.73
1	1A	2515	2MA	C4-C5-N7	-2.95	106.56	109.41
54	1w	37	MIA	C4-C5-N7	-2.94	106.57	109.41
57	2a	516	PSU	C5-C1'-C2'	-2.93	110.50	115.55
54	1w	37	MIA	C5-C6-N1	-2.93	117.72	120.64
58	B	9	BB9	O-C-CA	-2.90	121.76	125.47
1	2A	2251	OMG	N3-C2-N1	-2.88	123.25	127.46
1	1A	2263	OMG	C4-C5-N7	-2.87	106.64	109.41
54	1w	55	PSU	C5-C6-N1	-2.84	120.71	124.39
54	2w	37	MIA	C5-C6-N1	-2.84	117.81	120.64
32	1a	966	M2G	C4-C5-N7	-2.83	106.67	109.41
57	2a	1400	5MC	C5-C6-N1	-2.82	119.09	122.15
1	2A	2503	2MA	C4-C5-N7	-2.78	106.73	109.41
57	2a	527	G7M	CN7-N7-C8	-2.77	112.27	125.45
58	B	6	BB9	O-C-CA	-2.76	121.95	125.47
54	2w	46	G7M	CN7-N7-C8	-2.72	112.50	125.45
56	2y	55	PSU	C5-C1'-C2'	-2.71	110.87	115.55
1	2A	2251	OMG	C4-C5-N7	-2.69	106.81	109.41
55	1x	32	5MC	C5-C6-N1	-2.68	119.25	122.15
56	1y	46	G7M	CN7-N7-C8	-2.68	112.73	125.45
57	2a	1407	5MC	CM5-C5-C4	-2.67	118.90	121.65
56	1y	39	PSU	C5-C1'-C2'	-2.66	110.95	115.55
57	2a	966	M2G	C4-C5-N7	-2.66	106.84	109.41
56	2y	46	G7M	CN7-N7-C8	-2.64	112.91	125.45
32	1a	527	G7M	CN7-N7-C8	-2.63	112.95	125.45
54	2w	8	4SU	C5-C4-N3	-2.63	120.41	123.73
1	2A	1962	5MC	C5-C6-N1	-2.61	119.33	122.15
58	A	9	BB9	O-C-CA	-2.58	122.18	125.47
58	A	6	BB9	O-C-CA	-2.57	122.19	125.47
56	1y	55	PSU	C5-C1'-C2'	-2.52	111.20	115.55
54	1w	37	MIA	N3-C2-N1	-2.52	122.33	126.85
54	2w	46	G7M	N3-C2-N1	-2.50	123.81	127.46
55	2x	54	5MU	C5-C6-N1	-2.47	119.47	122.15
55	1x	54	5MU	C5-C6-N1	-2.46	119.48	122.15
1	1A	1933	PSU	C5-C1'-C2'	-2.46	111.30	115.55
55	2x	32	5MC	C5-C6-N1	-2.44	119.51	122.15
57	2a	1518	MA6	C10-N6-C9	-2.44	108.14	116.03
57	2a	527	G7M	N3-C2-N1	-2.44	123.90	127.46
54	1w	46	G7M	CN7-N7-C8	-2.42	113.93	125.45

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	2A	1915	5MU	C5-C6-N1	-2.38	119.57	122.15
1	1A	1964	5MC	C5-C6-N1	-2.38	119.58	122.15
1	1A	1984	5MC	C5-C6-N1	-2.35	119.60	122.15
54	1w	54	5MU	C5-C6-N1	-2.33	119.62	122.15
57	2a	967	5MC	C5-C6-N1	-2.33	119.63	122.15
56	1y	54	5MU	C5-C6-N1	-2.31	119.65	122.15
57	2a	1407	5MC	C5-C6-N1	-2.31	119.65	122.15
57	2a	1404	5MC	C5-C6-N1	-2.29	119.67	122.15
54	2w	39	PSU	C5-C1'-C2'	-2.29	111.60	115.55
32	1a	1407	5MC	C5-C6-N1	-2.20	119.77	122.15
55	2x	55	PSU	C5-C1'-C2'	-2.20	111.76	115.55
54	2w	37	MIA	N3-C2-N1	-2.14	123.01	126.85
57	2a	966	M2G	CM1-N2-C2	-2.12	119.32	121.34
32	1a	967	5MC	C5-C6-N1	-2.11	119.86	122.15
54	1w	39	PSU	C5-C1'-C2'	-2.09	111.95	115.55
1	1A	1937	5MU	C5-C6-N1	-2.07	119.91	122.15
54	2w	46	G7M	C6-C5-C4	-2.07	118.79	120.84
32	1a	527	G7M	N3-C2-N1	-2.06	124.44	127.46
32	1a	1207	2MG	N3-C2-N1	-2.06	123.12	126.23
32	1a	1400	5MC	C5-C6-N1	-2.04	119.94	122.15
32	1a	1518	MA6	C10-N6-C6	-2.03	113.37	119.51
1	1A	2564	2MU	C5-C4-N3	-2.00	118.33	123.12
54	1w	39	PSU	O4'-C1'-C2'	2.02	107.69	104.45
55	2x	55	PSU	O4'-C1'-C2'	2.04	107.72	104.45
56	2y	55	PSU	O4'-C1'-C2'	2.05	107.75	104.45
56	1y	55	PSU	O4'-C1'-C2'	2.07	107.77	104.45
1	2A	1917	PSU	O4'-C1'-C2'	2.12	107.86	104.45
57	2a	1498	UR3	C3U-N3-C4	2.16	121.01	118.15
57	2a	967	5MC	N4-C4-N3	2.20	120.26	117.00
57	2a	1404	5MC	N4-C4-N3	2.23	120.30	117.00
32	1a	516	PSU	O4'-C1'-C2'	2.24	108.05	104.45
1	2A	1911	PSU	O4'-C1'-C2'	2.25	108.06	104.45
57	2a	516	PSU	O4'-C1'-C2'	2.25	108.07	104.45
32	1a	966	M2G	N1-C2-N2	2.26	119.48	117.16
57	2a	1207	2MG	N2-C2-N1	2.32	119.21	116.95
56	1y	46	G7M	C1'-N9-C4	2.35	130.69	126.64
1	1A	1984	5MC	N4-C4-N3	2.37	120.50	117.00
57	2a	1400	5MC	N4-C4-N3	2.39	120.53	117.00
55	2x	32	5MC	N4-C4-N3	2.47	120.65	117.00
32	1a	1407	5MC	N4-C4-N3	2.52	120.73	117.00
32	1a	1400	5MC	N4-C4-N3	2.58	120.81	117.00
55	1x	32	5MC	N4-C4-N3	2.59	120.83	117.00

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
32	1a	1498	UR3	C3U-N3-C4	2.62	121.62	118.15
56	2y	46	G7M	C1'-N9-C4	2.80	131.47	126.64
57	2a	1407	5MC	N4-C4-N3	2.82	121.17	117.00
1	1A	1964	5MC	N4-C4-N3	2.82	121.17	117.00
1	2A	1942	5MC	N4-C4-N3	2.84	121.19	117.00
54	2w	37	MIA	C2-N1-C6	2.89	121.98	113.47
32	1a	967	5MC	N4-C4-N3	2.90	121.29	117.00
1	2A	1962	5MC	N4-C4-N3	2.92	121.32	117.00
54	1w	37	MIA	C2-N1-C6	3.00	122.31	113.47
32	1a	1519	MA6	N1-C6-N6	3.03	120.22	117.00
54	1w	46	G7M	C2-N3-C4	3.10	118.77	115.16
54	2w	8	4SU	C2-N3-C4	3.25	119.90	115.11
32	1a	1207	2MG	N2-C2-N1	3.37	120.23	116.95
32	1a	527	G7M	C2-N3-C4	3.48	119.23	115.16
54	2w	46	G7M	C2-N3-C4	3.54	119.29	115.16
56	1y	55	PSU	C6-N1-C2	3.66	121.22	115.36
32	1a	1519	MA6	C2-N1-C6	3.81	121.16	111.82
54	1w	55	PSU	C6-N1-C2	3.92	121.64	115.36
56	2y	32	PSU	C6-N1-C2	3.97	121.71	115.36
57	2a	1207	2MG	C6-N1-C2	3.97	122.29	115.18
56	1y	32	PSU	C6-N1-C2	3.99	121.75	115.36
1	2A	1917	PSU	C6-N1-C2	3.99	121.75	115.36
32	1a	1207	2MG	C6-N1-C2	4.01	122.35	115.18
57	2a	527	G7M	C2-N3-C4	4.02	119.85	115.16
56	2y	55	PSU	C6-N1-C2	4.05	121.83	115.36
57	2a	1519	MA6	C2-N1-C6	4.07	121.81	111.82
56	1y	46	G7M	C2-N3-C4	4.07	119.91	115.16
54	1w	32	PSU	C6-N1-C2	4.08	121.89	115.36
1	1A	1933	PSU	C6-N1-C2	4.12	121.96	115.36
1	2A	1911	PSU	C6-N1-C2	4.13	121.97	115.36
1	2A	2251	OMG	C6-N1-C2	4.13	122.00	116.06
54	1w	39	PSU	C6-N1-C2	4.19	122.06	115.36
56	2y	39	PSU	C6-N1-C2	4.25	122.16	115.36
55	1x	55	PSU	C6-N1-C2	4.28	122.21	115.36
1	1A	1939	PSU	C6-N1-C2	4.28	122.21	115.36
55	2x	55	PSU	C6-N1-C2	4.29	122.22	115.36
57	2a	516	PSU	C6-N1-C2	4.29	122.23	115.36
54	2w	39	PSU	C6-N1-C2	4.29	122.23	115.36
56	1y	39	PSU	C6-N1-C2	4.32	122.27	115.36
1	1A	2617	PSU	C6-N1-C2	4.32	122.27	115.36
56	2y	8	4SU	C2-N3-C4	4.32	121.48	115.11
54	2w	55	PSU	C6-N1-C2	4.33	122.29	115.36

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
32	1a	1518	MA6	C2-N1-C6	4.40	122.61	111.82
54	2w	32	PSU	C6-N1-C2	4.43	122.45	115.36
56	2y	46	G7M	C2-N3-C4	4.44	120.34	115.16
56	1y	8	4SU	C2-N3-C4	4.47	121.70	115.11
32	1a	516	PSU	C6-N1-C2	4.49	122.55	115.36
57	2a	1518	MA6	C2-N1-C6	4.50	122.86	111.82
1	2A	2605	PSU	C6-N1-C2	4.54	122.62	115.36
1	1A	2263	OMG	C6-N1-C2	4.67	122.78	116.06
32	1a	527	G7M	C6-N1-C2	4.72	122.85	116.06
57	2a	527	G7M	C6-N1-C2	4.72	122.85	116.06
1	2A	2251	OMG	C2-N3-C4	4.72	120.68	115.16
57	2a	1207	2MG	C2-N3-C4	4.75	120.53	115.11
54	1w	8	4SU	C2-N3-C4	4.75	122.11	115.11
54	2w	46	G7M	C6-N1-C2	4.78	122.93	116.06
57	2a	966	M2G	C6-N1-C2	4.82	121.92	116.18
32	1a	966	M2G	C2-N3-C4	4.93	120.73	115.11
57	2a	966	M2G	C2-N3-C4	4.98	120.80	115.11
32	1a	1207	2MG	C2-N3-C4	4.99	120.80	115.11
1	2A	1939	5MU	C4-N3-C2	5.06	119.58	115.16
1	1A	2263	OMG	C2-N3-C4	5.09	121.10	115.16
32	1a	966	M2G	C6-N1-C2	5.16	122.33	116.18
1	1A	1961	5MU	C4-N3-C2	5.30	119.79	115.16
56	1y	46	G7M	C6-N1-C2	5.55	124.04	116.06
1	1A	1937	5MU	C4-N3-C2	5.64	120.09	115.16
1	2A	2605	PSU	C4-N3-C2	5.71	120.15	115.16
1	2A	1915	5MU	C4-N3-C2	5.87	120.29	115.16
55	2x	54	5MU	C4-N3-C2	5.94	120.36	115.16
1	2A	2503	2MA	C2-N3-C4	5.95	120.55	115.41
57	2a	516	PSU	C4-N3-C2	6.16	120.55	115.16
56	2y	46	G7M	C6-N1-C2	6.17	124.93	116.06
55	1x	55	PSU	C4-N3-C2	6.18	120.56	115.16
56	1y	39	PSU	C4-N3-C2	6.27	120.64	115.16
54	1w	54	5MU	C4-N3-C2	6.28	120.65	115.16
32	1a	516	PSU	C4-N3-C2	6.30	120.67	115.16
55	2x	55	PSU	C4-N3-C2	6.30	120.67	115.16
54	2w	55	PSU	C4-N3-C2	6.31	120.68	115.16
1	1A	2617	PSU	C4-N3-C2	6.31	120.68	115.16
56	2y	39	PSU	C4-N3-C2	6.34	120.70	115.16
1	1A	1939	PSU	C4-N3-C2	6.42	120.77	115.16
54	2w	32	PSU	C4-N3-C2	6.43	120.79	115.16
54	1w	46	G7M	C6-N1-C2	6.44	125.33	116.06
56	2y	55	PSU	C4-N3-C2	6.47	120.82	115.16

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
1	1A	1933	PSU	C4-N3-C2	6.53	120.87	115.16
1	2A	1911	PSU	C4-N3-C2	6.59	120.92	115.16
1	1A	2515	2MA	C2-N3-C4	6.61	121.13	115.41
55	1x	54	5MU	C4-N3-C2	6.73	121.05	115.16
1	2A	1917	PSU	C4-N3-C2	6.80	121.11	115.16
54	1w	32	PSU	C4-N3-C2	6.87	121.17	115.16
56	1y	32	PSU	C4-N3-C2	6.88	121.18	115.16
1	2A	2552	2MU	C4-N3-C2	6.90	120.06	114.13
54	1w	39	PSU	C4-N3-C2	6.92	121.21	115.16
56	1y	55	PSU	C4-N3-C2	6.98	121.27	115.16
56	2y	32	PSU	C4-N3-C2	7.17	121.43	115.16
54	1w	55	PSU	C4-N3-C2	7.37	121.60	115.16
56	2y	54	5MU	C4-N3-C2	7.46	121.69	115.16
54	2w	39	PSU	C4-N3-C2	7.49	121.71	115.16
56	1y	54	5MU	C4-N3-C2	7.58	121.79	115.16
1	1A	2564	2MU	C4-N3-C2	7.65	120.70	114.13
55	2x	8	4SU	C2-N3-C4	7.89	126.75	115.11
54	2w	54	5MU	C4-N3-C2	8.21	122.34	115.16
55	1x	8	4SU	C2-N3-C4	8.92	128.27	115.11

There are no chirality outliers.

There are no torsion outliers.

There are no ring outliers.

5 monomers are involved in 6 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
1	1A	1942	4OC	1	0
1	1A	1961	5MU	2	0
1	2A	2251	OMG	1	0
1	2A	2503	2MA	1	0
1	2A	2552	2MU	1	0

5.5 Carbohydrates [i](#)

There are no carbohydrates in this entry.

5.6 Ligand geometry [i](#)

Of 2940 ligands modelled in this entry, 2938 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	$\# Z > 2$
61	SF4	1d	501	35	0,12,12	0.00	-	0,24,24	0.00	-
61	SF4	2d	501	35	0,12,12	0.00	-	0,24,24	0.00	-

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

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
61	SF4	1d	501	35	-	0/0/48/48	0/6/5/5
61	SF4	2d	501	35	-	0/0/48/48	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.

No monomer is involved in short contacts.

5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

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	1A	2860/2915 (98%)	0.57	57 (1%) 65 66	13, 29, 85, 99	0
1	2A	2789/2915 (95%)	0.47	79 (2%) 53 54	25, 51, 87, 102	0
2	1B	120/121 (99%)	0.11	0 100 100	23, 41, 55, 80	0
2	2B	120/121 (99%)	-0.01	0 100 100	55, 72, 80, 89	0
3	1D	275/276 (99%)	0.62	4 (1%) 74 75	14, 29, 45, 72	0
3	2D	275/276 (99%)	0.83	14 (5%) 29 27	25, 43, 57, 71	0
4	1E	204/206 (99%)	0.53	1 (0%) 90 92	14, 32, 53, 71	0
4	2E	204/206 (99%)	0.57	8 (3%) 40 39	28, 53, 65, 73	0
5	1F	203/210 (96%)	0.47	0 100 100	15, 35, 60, 77	0
5	2F	203/210 (96%)	0.60	8 (3%) 40 39	30, 61, 75, 83	0
6	1G	181/182 (99%)	0.29	4 (2%) 62 63	33, 50, 66, 81	0
6	2G	181/182 (99%)	0.94	29 (16%) 2 1	62, 73, 81, 89	0
7	1H	174/180 (96%)	0.32	1 (0%) 89 90	31, 45, 59, 63	0
7	2H	174/180 (96%)	1.29	45 (25%) 1 0	65, 78, 84, 88	0
8	1I	146/148 (98%)	0.29	0 100 100	37, 66, 76, 80	0
8	2I	146/148 (98%)	0.16	0 100 100	48, 65, 75, 81	0
9	1N	140/140 (100%)	0.46	0 100 100	18, 31, 53, 63	0
9	2N	140/140 (100%)	0.58	4 (2%) 52 52	40, 57, 71, 79	0
10	1O	122/122 (100%)	0.53	0 100 100	21, 32, 48, 54	0
10	2O	122/122 (100%)	0.73	4 (3%) 47 46	41, 53, 63, 68	0
11	1P	149/150 (99%)	0.52	2 (1%) 77 78	14, 37, 63, 72	0
11	2P	149/150 (99%)	0.88	12 (8%) 13 10	29, 61, 75, 82	0
12	1Q	141/141 (100%)	0.49	1 (0%) 87 88	21, 32, 50, 67	0
12	2Q	141/141 (100%)	0.77	14 (9%) 8 6	40, 61, 72, 76	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
13	1R	118/118 (100%)	0.58	0 100 100	18, 27, 41, 48	0
13	2R	118/118 (100%)	0.56	1 (0%) 86 86	34, 45, 55, 65	0
14	1S	110/112 (98%)	0.25	0 100 100	30, 41, 54, 59	0
14	2S	110/112 (98%)	0.64	13 (11%) 5 4	56, 67, 75, 80	0
15	1T	131/146 (89%)	0.48	0 100 100	24, 35, 61, 70	0
15	2T	131/146 (89%)	0.51	1 (0%) 86 86	44, 55, 73, 77	0
16	1U	116/118 (98%)	0.57	0 100 100	15, 23, 37, 54	0
16	2U	116/118 (98%)	0.67	5 (4%) 36 34	38, 55, 69, 76	0
17	1V	101/101 (100%)	0.31	1 (0%) 82 82	15, 32, 47, 57	0
17	2V	101/101 (100%)	0.40	2 (1%) 65 66	35, 64, 70, 75	0
18	1W	112/113 (99%)	0.58	2 (1%) 69 70	17, 25, 44, 78	0
18	2W	112/113 (99%)	0.72	6 (5%) 26 25	31, 42, 59, 87	0
19	1X	95/96 (98%)	0.48	0 100 100	17, 31, 52, 73	0
19	2X	95/96 (98%)	0.65	4 (4%) 37 35	38, 53, 65, 76	0
20	1Y	107/110 (97%)	0.34	2 (1%) 67 68	28, 42, 62, 77	0
20	2Y	107/110 (97%)	1.28	20 (18%) 1 1	50, 65, 73, 80	0
21	1Z	154/206 (74%)	0.36	5 (3%) 48 48	33, 56, 77, 85	0
21	2Z	160/206 (77%)	1.01	29 (18%) 1 1	63, 76, 86, 89	0
22	10	83/85 (97%)	0.74	6 (7%) 16 14	22, 29, 51, 60	0
22	20	83/85 (97%)	0.99	9 (10%) 6 5	39, 58, 67, 74	0
23	11	97/98 (98%)	0.69	4 (4%) 38 36	20, 36, 62, 70	0
23	21	97/98 (98%)	0.83	4 (4%) 38 36	33, 49, 69, 73	0
24	12	70/72 (97%)	0.46	0 100 100	29, 41, 53, 67	0
24	22	70/72 (97%)	0.56	4 (5%) 24 23	52, 63, 71, 76	0
25	13	59/60 (98%)	0.39	0 100 100	17, 28, 50, 75	0
25	23	59/60 (98%)	0.77	5 (8%) 11 9	46, 58, 75, 84	0
26	14	69/71 (97%)	0.38	4 (5%) 24 22	41, 65, 81, 87	0
26	24	69/71 (97%)	0.95	15 (21%) 1 1	71, 79, 87, 89	0
27	15	59/60 (98%)	0.58	1 (1%) 70 72	15, 24, 44, 53	0
27	25	59/60 (98%)	0.56	0 100 100	31, 46, 56, 72	0
28	16	53/54 (98%)	0.39	1 (1%) 67 68	25, 34, 49, 56	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
28	26	53/54 (98%)	0.81	3 (5%) 24 23	42, 54, 63, 68	0
29	17	48/49 (97%)	0.85	3 (6%) 21 19	16, 22, 47, 54	0
29	27	48/49 (97%)	1.20	8 (16%) 2 1	27, 34, 57, 66	0
30	18	64/65 (98%)	0.47	0 100 100	21, 26, 36, 48	0
30	28	64/65 (98%)	1.01	7 (10%) 6 5	40, 49, 57, 60	0
31	19	37/37 (100%)	0.59	0 100 100	22, 31, 48, 53	0
31	29	37/37 (100%)	1.66	13 (35%) 0 0	55, 63, 72, 73	0
32	1a	1488/1521 (97%)	0.37	33 (2%) 62 63	28, 57, 86, 101	0
33	1b	231/256 (90%)	0.54	13 (5%) 25 23	57, 72, 82, 90	0
33	2b	231/256 (90%)	0.97	40 (17%) 2 1	67, 80, 86, 91	0
34	1c	206/239 (86%)	0.50	8 (3%) 40 39	51, 64, 75, 82	0
34	2c	206/239 (86%)	1.01	37 (17%) 2 1	67, 78, 83, 88	0
35	1d	208/209 (99%)	0.66	17 (8%) 12 10	49, 61, 73, 81	0
35	2d	208/209 (99%)	0.84	21 (10%) 8 6	56, 66, 74, 81	0
36	1e	148/162 (91%)	0.48	1 (0%) 87 88	44, 55, 68, 72	0
36	2e	148/162 (91%)	0.98	27 (18%) 1 1	60, 72, 79, 84	0
37	1f	100/101 (99%)	0.24	1 (1%) 82 82	44, 60, 69, 77	0
37	2f	100/101 (99%)	0.17	0 100 100	50, 61, 69, 78	0
38	1g	155/156 (99%)	0.64	12 (7%) 14 12	46, 61, 74, 86	0
38	2g	155/156 (99%)	0.98	26 (16%) 2 1	62, 71, 80, 85	0
39	1h	137/138 (99%)	0.54	9 (6%) 19 17	48, 58, 65, 72	0
39	2h	137/138 (99%)	0.76	13 (9%) 9 7	60, 71, 76, 81	0
40	1i	127/128 (99%)	0.73	10 (7%) 13 11	47, 67, 76, 80	0
40	2i	127/128 (99%)	1.71	47 (37%) 0 0	66, 76, 82, 84	0
41	1j	97/105 (92%)	0.64	8 (8%) 12 10	48, 70, 80, 85	0
41	2j	96/105 (91%)	1.74	36 (37%) 0 0	70, 79, 87, 91	0
42	1k	114/129 (88%)	0.60	3 (2%) 56 56	36, 57, 68, 76	0
42	2k	114/129 (88%)	0.73	7 (6%) 22 20	48, 65, 74, 78	0
43	1l	121/132 (91%)	0.58	1 (0%) 86 86	36, 45, 57, 62	0
43	2l	121/132 (91%)	0.95	18 (14%) 3 2	48, 61, 71, 76	0
44	1m	123/126 (97%)	0.38	7 (5%) 24 23	46, 60, 69, 73	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
44	2m	122/126 (96%)	1.16	27 (22%) 1 1	65, 75, 80, 82	0
45	1n	60/61 (98%)	0.88	3 (5%) 30 28	50, 57, 66, 69	0
45	2n	60/61 (98%)	2.46	32 (53%) 0 0	70, 77, 81, 83	0
46	1o	88/89 (98%)	0.56	5 (5%) 24 23	42, 56, 67, 74	0
46	2o	88/89 (98%)	0.76	9 (10%) 7 5	53, 67, 75, 77	0
47	1p	82/88 (93%)	0.80	3 (3%) 42 41	48, 59, 69, 71	0
47	2p	82/88 (93%)	0.82	8 (9%) 8 6	55, 63, 70, 73	0
48	1q	99/105 (94%)	0.77	7 (7%) 17 14	47, 58, 70, 72	0
48	2q	99/105 (94%)	1.14	18 (18%) 1 1	56, 65, 74, 79	0
49	1r	68/88 (77%)	0.44	1 (1%) 74 75	49, 58, 70, 72	0
49	2r	68/88 (77%)	0.50	2 (2%) 52 52	53, 64, 74, 79	0
50	1s	83/93 (89%)	0.32	3 (3%) 43 42	52, 62, 70, 77	0
50	2s	83/93 (89%)	1.53	27 (32%) 0 0	71, 79, 85, 86	0
51	1t	96/106 (90%)	0.74	10 (10%) 7 5	49, 60, 71, 77	0
51	2t	96/106 (90%)	0.94	13 (13%) 3 2	54, 64, 75, 77	0
52	1u	23/27 (85%)	1.70	9 (39%) 0 0	53, 57, 63, 66	0
52	2u	23/27 (85%)	3.26	18 (78%) 0 0	68, 73, 77, 79	0
53	1v	13/24 (54%)	1.47	3 (23%) 1 1	41, 47, 71, 95	0
53	2v	13/24 (54%)	2.31	6 (46%) 0 0	59, 72, 89, 91	0
54	1w	67/76 (88%)	1.38	16 (23%) 1 1	34, 80, 94, 98	0
54	2w	65/76 (85%)	1.85	19 (29%) 1 0	52, 89, 95, 101	0
55	1x	72/77 (93%)	0.47	2 (2%) 53 54	28, 59, 75, 84	0
55	2x	72/77 (93%)	0.37	0 100 100	43, 72, 83, 85	0
56	1y	68/76 (89%)	0.71	8 (11%) 5 4	28, 84, 91, 95	0
56	2y	67/76 (88%)	1.68	23 (34%) 0 0	46, 90, 94, 97	0
57	2a	1491/1521 (98%)	0.47	55 (3%) 42 41	42, 71, 89, 101	0
58	A	13/18 (72%)	1.81	3 (23%) 1 1	17, 21, 65, 77	0
58	B	13/18 (72%)	1.95	4 (30%) 0 0	32, 35, 69, 80	0
All	All	20903/21784 (95%)	0.61	1194 (5%) 24 23	13, 56, 83, 102	0

All (1194) RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
54	2w	71	G	12.1
38	1g	80	VAL	11.4
54	1w	70	G	9.9
23	21	2	SER	9.1
45	2n	34	TYR	8.8
57	2a	1030(B)	C	7.9
54	2w	70	G	7.6
58	A	16	ASN	7.6
1	2A	883	G	7.6
33	2b	165	VAL	7.5
54	1w	71	G	7.4
21	2Z	144	LEU	7.4
38	2g	82	GLY	7.3
45	2n	39	LEU	7.2
58	B	16	ASN	6.9
45	1n	2	ALA	6.7
44	2m	102	ARG	6.7
41	2j	47	PHE	6.7
41	2j	85	LEU	6.6
52	2u	6	ARG	6.5
38	2g	81	GLY	6.4
54	2w	72	C	6.4
1	2A	2802	G	6.4
57	2a	1034	G	6.4
3	2D	38	LYS	6.2
1	2A	896	A	6.2
1	2A	2146	C	6.2
44	2m	123	ALA	6.2
56	2y	36	A	6.1
45	2n	2	ALA	6.1
31	29	37	GLY	6.0
35	2d	164	ALA	6.0
56	1y	20	U	5.9
57	2a	1030(A)	G	5.8
1	2A	2155	G	5.8
41	2j	55	LYS	5.8
45	2n	25	VAL	5.7
45	2n	51	GLY	5.6
57	2a	1033	G	5.6
26	24	49	PHE	5.6
43	2l	18	VAL	5.5
53	2v	14	A	5.5
39	2h	2	LEU	5.5

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Mol	Chain	Res	Type	RSRZ
22	10	6	GLY	5.4
3	2D	2	ALA	5.4
41	2j	59	SER	5.4
1	1A	931	C	5.4
38	1g	85	TYR	5.4
36	2e	12	LEU	5.3
40	2i	19	LEU	5.3
38	2g	80	VAL	5.3
53	1v	12	A	5.3
40	2i	14	VAL	5.3
54	2w	4	C	5.3
56	2y	34	G	5.2
52	2u	14	TRP	5.2
40	2i	124	GLN	5.1
54	1w	72	C	5.1
57	2a	1036	G	5.1
45	2n	6	LEU	5.1
57	2a	1257	U	5.1
1	2A	2145	C	5.1
34	2c	182	ILE	5.0
38	1g	82	GLY	5.0
52	2u	11	GLY	5.0
40	2i	125	TYR	5.0
44	2m	122	LYS	4.9
1	2A	2154	G	4.9
48	1q	98	LEU	4.9
44	2m	124	PRO	4.9
53	2v	24	A	4.9
52	2u	16	GLY	4.9
34	2c	198	VAL	4.9
38	1g	79	ARG	4.9
54	1w	69	G	4.9
29	17	48	LYS	4.9
21	2Z	170	THR	4.8
52	2u	24	ARG	4.8
44	2m	104	ARG	4.8
1	1A	1221	G	4.8
32	1a	1001(A)	G	4.8
32	1a	1036	G	4.8
32	1a	1030	C	4.7
1	2A	882	G	4.7
1	1A	942	A	4.7

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Mol	Chain	Res	Type	RSRZ
26	24	56	VAL	4.7
45	2n	12	ARG	4.7
50	2s	13	ASP	4.7
42	2k	25	TYR	4.7
6	2G	28	VAL	4.7
54	2w	73	A	4.7
1	2A	2160	G	4.6
32	1a	1030(A)	G	4.6
38	2g	4	ARG	4.6
12	2Q	104	PHE	4.6
54	2w	6	G	4.5
50	2s	12	ASP	4.5
58	A	15	ALA	4.5
52	2u	10	ARG	4.5
21	2Z	149	SER	4.5
32	1a	1030(C)	G	4.5
33	2b	187	LEU	4.5
48	2q	100	LYS	4.5
1	2A	2127	G	4.5
57	2a	1030	C	4.4
50	2s	80	TYR	4.4
26	24	63	TYR	4.4
38	1g	84	ASN	4.4
33	2b	123	ALA	4.4
32	1a	1003	G	4.4
25	23	60	GLU	4.4
27	15	60	VAL	4.3
7	2H	48	GLY	4.3
19	2X	92	LEU	4.3
40	1i	19	LEU	4.3
44	2m	66	LEU	4.3
38	2g	83	ALA	4.3
1	2A	888	C	4.3
40	2i	66	ARG	4.3
47	2p	9	PHE	4.3
1	2A	2803	C	4.3
40	2i	28	VAL	4.3
33	2b	101	MET	4.3
23	11	2	SER	4.3
54	1w	3	C	4.2
38	2g	79	ARG	4.2
1	1A	2167	C	4.2

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Mol	Chain	Res	Type	RSRZ
54	1w	20	U	4.2
57	2a	1035	A	4.2
54	2w	45	U	4.2
29	27	47	ARG	4.2
1	1A	1555	C	4.2
54	2w	13	C	4.2
56	2y	53	G	4.2
44	2m	6	GLY	4.1
45	2n	29	ARG	4.1
22	20	7	LEU	4.1
53	2v	12	A	4.1
32	1a	1035	A	4.1
29	27	1	MET	4.1
6	2G	48	GLU	4.1
52	2u	5	ASP	4.1
1	2A	2896	C	4.1
50	2s	48	THR	4.1
44	2m	120	LYS	4.1
52	2u	15	ARG	4.1
33	2b	92	TYR	4.1
35	2d	23	GLY	4.1
45	2n	38	GLY	4.1
20	2Y	1	MET	4.1
1	1A	932	C	4.0
41	2j	98	ILE	4.0
50	2s	79	THR	4.0
54	1w	44	G	4.0
47	2p	19	ILE	4.0
1	1A	943	C	4.0
7	2H	35	VAL	4.0
33	2b	118	LEU	4.0
26	24	50	VAL	4.0
54	2w	5	G	4.0
57	2a	1021	G	4.0
38	1g	83	ALA	4.0
22	10	7	LEU	4.0
40	2i	81	ILE	4.0
36	2e	10	MET	4.0
1	2A	2897	U	4.0
32	1a	1030(B)	C	4.0
20	2Y	106	LEU	4.0
1	2A	2144	U	4.0

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Mol	Chain	Res	Type	RSRZ
1	2A	2159	G	4.0
20	2Y	45	VAL	4.0
1	2A	2112	G	3.9
3	2D	276	LYS	3.9
32	1a	1032	G	3.9
41	2j	88	LEU	3.9
11	2P	79	ARG	3.9
31	29	15	LYS	3.9
7	2H	159	GLU	3.9
20	1Y	1	MET	3.9
40	2i	115	GLY	3.9
34	2c	8	ILE	3.9
1	2A	2115	G	3.9
41	2j	66	ARG	3.9
34	2c	194	GLY	3.9
32	1a	1531	A	3.9
40	2i	9	ARG	3.8
41	2j	65	LEU	3.8
33	1b	133	LYS	3.8
1	2A	885	C	3.8
32	1a	1001	A	3.8
34	2c	193	TYR	3.8
3	2D	275	LYS	3.8
40	2i	90	PRO	3.8
38	2g	7	ALA	3.8
41	2j	40	LEU	3.8
43	2l	32	PHE	3.8
21	2Z	172	ALA	3.8
40	1i	106	ALA	3.8
23	21	28	GLY	3.8
44	2m	87	TYR	3.8
32	1a	204	U	3.8
7	2H	89	ILE	3.8
50	2s	30	LEU	3.8
53	2v	13	A	3.7
33	2b	152	PHE	3.7
44	2m	4	ILE	3.7
40	2i	109	VAL	3.7
1	2A	2113	U	3.7
45	2n	42	ILE	3.7
50	2s	71	LEU	3.7
26	24	58	ARG	3.7

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Mol	Chain	Res	Type	RSRZ
1	2A	884	C	3.7
50	1s	71	LEU	3.7
1	2A	2147	G	3.7
3	1D	276	LYS	3.7
35	2d	161	ASN	3.7
40	2i	36	TYR	3.7
34	2c	157	ILE	3.7
5	2F	62	ARG	3.6
50	2s	68	GLY	3.6
57	2a	1001(A)	G	3.6
12	2Q	22	LYS	3.6
40	2i	82	ALA	3.6
41	2j	44	VAL	3.6
21	2Z	122	ARG	3.6
32	1a	1029	C	3.6
7	2H	2	SER	3.6
11	2P	77	ARG	3.6
26	24	51	ASP	3.6
32	1a	162	A	3.6
41	2j	62	HIS	3.6
38	2g	152	ALA	3.6
36	2e	115	VAL	3.6
41	2j	38	ILE	3.6
1	2A	2111	C	3.6
40	2i	7	THR	3.6
14	2S	32	LEU	3.6
34	2c	197	GLY	3.6
38	1g	153	HIS	3.6
22	20	2	ALA	3.6
20	2Y	75	ILE	3.6
33	2b	37	ASN	3.6
1	1A	271	U	3.5
44	2m	60	VAL	3.5
32	1a	1257	U	3.5
38	2g	156	TRP	3.5
1	2A	2173	A	3.5
44	2m	92	HIS	3.5
38	2g	6	ARG	3.5
40	2i	49	PRO	3.5
45	2n	55	GLY	3.5
45	2n	50	LYS	3.5
52	2u	9	ARG	3.5

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Mol	Chain	Res	Type	RSRZ
54	2w	76	A	3.5
18	2W	60	ASN	3.5
36	2e	31	LEU	3.5
14	2S	35	ILE	3.5
43	1l	64	TYR	3.5
45	2n	35	ARG	3.5
1	2A	897	C	3.5
22	10	3	HIS	3.5
54	2w	44	G	3.5
56	1y	35	A	3.5
34	2c	4	LYS	3.5
40	2i	110	GLU	3.5
40	2i	105	ASP	3.5
52	2u	17	THR	3.5
29	17	47	ARG	3.5
1	2A	2139	C	3.4
20	2Y	55	TYR	3.4
41	2j	58	ASP	3.4
1	2A	2128	C	3.4
40	2i	123	PRO	3.4
55	1x	67	C	3.4
45	2n	31	ARG	3.4
7	2H	124	GLU	3.4
1	2A	1509	C	3.4
50	2s	41	VAL	3.4
54	2w	69	G	3.4
21	1Z	149	SER	3.4
5	2F	166	ALA	3.4
34	2c	189	ALA	3.4
31	29	25	VAL	3.4
32	1a	1034	G	3.4
7	2H	128	PRO	3.4
23	1l	98	LEU	3.4
35	1d	170	VAL	3.4
56	2y	45	U	3.4
52	1u	14	TRP	3.4
6	2G	169	ALA	3.4
6	2G	2	PRO	3.4
32	1a	1031	G	3.4
16	2U	2	PRO	3.4
15	2T	111	ARG	3.4
32	1a	161	A	3.4

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Mol	Chain	Res	Type	RSRZ
20	2Y	65	ALA	3.3
7	2H	49	VAL	3.3
44	2m	78	ILE	3.3
57	2a	1531	A	3.3
32	1a	1532	U	3.3
45	2n	61	TRP	3.3
45	2n	57	ARG	3.3
1	1A	944	C	3.3
33	2b	70	PHE	3.3
34	2c	124	ILE	3.3
32	1a	160	A	3.3
1	2A	2149	G	3.3
29	27	46	VAL	3.3
34	2c	188	LEU	3.3
7	2H	52	VAL	3.3
41	2j	63	PHE	3.3
45	2n	37	PHE	3.3
54	1w	73	A	3.3
6	2G	29	TRP	3.3
41	2j	51	ARG	3.3
51	2t	86	ARG	3.3
1	2A	652(B)	A	3.3
7	2H	72	ILE	3.3
34	2c	196	LEU	3.3
7	2H	76	VAL	3.2
33	2b	19	HIS	3.2
41	2j	48	THR	3.2
48	2q	90	ILE	3.2
14	2S	3	ARG	3.2
56	2y	52	G	3.2
57	2a	1030(C)	G	3.2
12	2Q	66	ILE	3.2
12	2Q	103	MET	3.2
32	1a	1503	A	3.2
7	2H	45	VAL	3.2
33	2b	40	HIS	3.2
50	2s	52	TYR	3.2
46	2o	68	ARG	3.2
7	2H	24	VAL	3.2
34	2c	6	HIS	3.2
40	2i	114	TYR	3.2
56	2y	56	C	3.2

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Mol	Chain	Res	Type	RSRZ
52	2u	8	THR	3.2
18	2W	6	ILE	3.2
1	1A	934	A	3.2
53	1v	13	A	3.2
52	1u	2	GLY	3.2
41	2j	96	ILE	3.2
33	1b	188	ALA	3.2
56	2y	35	A	3.2
9	2N	83	LYS	3.2
51	1t	38	LYS	3.2
20	2Y	5	MET	3.2
56	2y	57	G	3.2
1	1A	1141	A	3.1
54	2w	7	A	3.1
7	2H	44	VAL	3.1
1	2A	2156	G	3.1
20	2Y	50	ARG	3.1
21	1Z	141	VAL	3.1
6	2G	157	ILE	3.1
20	2Y	78	ALA	3.1
38	2g	85	TYR	3.1
40	2i	4	TYR	3.1
51	2t	13	LEU	3.1
44	1m	124	PRO	3.1
3	1D	38	LYS	3.1
46	2o	60	VAL	3.1
34	2c	14	ILE	3.1
40	2i	127	LYS	3.1
21	2Z	139	VAL	3.1
1	2A	2116	G	3.1
18	2W	112	GLY	3.1
41	2j	56	HIS	3.1
31	29	16	VAL	3.1
40	2i	61	ALA	3.1
12	2Q	121	ALA	3.1
36	2e	86	ALA	3.1
36	2e	119	LEU	3.1
44	1m	2	ALA	3.1
41	2j	6	ILE	3.1
31	29	19	ARG	3.1
43	2l	64	TYR	3.1
57	2a	1150	U	3.1

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Mol	Chain	Res	Type	RSRZ
35	2d	166	LYS	3.1
56	2y	65	G	3.1
50	2s	84	GLY	3.1
36	2e	16	THR	3.1
1	2A	886	C	3.1
32	1a	1030(D)	A	3.1
45	2n	10	ALA	3.1
22	10	5	LYS	3.1
26	14	54	GLY	3.0
51	1t	70	SER	3.0
1	1A	935	C	3.0
1	1A	2160	C	3.0
52	2u	13	ILE	3.0
57	2a	1357	A	3.0
40	2i	27	THR	3.0
44	2m	121	LYS	3.0
1	2A	614(B)	G	3.0
1	2A	2133	G	3.0
36	2e	109	ILE	3.0
44	2m	119	GLY	3.0
34	2c	186	PHE	3.0
53	1v	24	A	3.0
6	2G	49	ASP	3.0
38	2g	86	GLN	3.0
44	2m	70	LEU	3.0
48	2q	98	LEU	3.0
34	2c	39	ILE	3.0
20	2Y	54	LYS	3.0
40	2i	62	TYR	3.0
40	2i	128	ARG	3.0
54	1w	4	C	3.0
7	2H	86	GLU	3.0
6	2G	136	ARG	3.0
38	1g	154	TYR	3.0
21	2Z	5	LEU	3.0
35	1d	157	LEU	3.0
3	1D	275	LYS	3.0
32	1a	1447	A	3.0
54	2w	14	A	3.0
35	2d	160	GLN	3.0
22	20	45	PHE	3.0
48	1q	27	PHE	3.0

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Mol	Chain	Res	Type	RSRZ
32	1a	1002	G	3.0
54	1w	2	C	3.0
35	2d	18	LYS	3.0
52	2u	18	TYR	3.0
35	1d	120	LEU	3.0
36	2e	90	VAL	3.0
41	2j	71	LEU	3.0
43	2l	39	VAL	3.0
52	2u	2	GLY	3.0
33	2b	232	PRO	3.0
1	1A	2814	C	2.9
51	1t	63	ILE	2.9
57	2a	1026	G	2.9
36	2e	84	PHE	2.9
26	24	53	GLU	2.9
57	2a	1287	A	2.9
5	2F	131	GLY	2.9
38	1g	81	GLY	2.9
33	2b	97	TRP	2.9
6	2G	88	ILE	2.9
42	2k	29	ILE	2.9
1	2A	1536	C	2.9
24	22	1	MET	2.9
36	2e	21	ALA	2.9
50	2s	14	HIS	2.9
36	2e	125	SER	2.9
1	1A	2906	U	2.9
22	10	4	LYS	2.9
34	2c	190	ARG	2.9
57	2a	1027	C	2.9
33	2b	31	TYR	2.9
21	2Z	141	VAL	2.9
33	2b	48	MET	2.9
44	2m	23	TYR	2.9
34	2c	160	ALA	2.9
51	1t	74	LYS	2.9
1	2A	1026	U	2.9
6	2G	146	TYR	2.9
41	2j	34	VAL	2.9
1	2A	881	G	2.9
41	2j	74	ILE	2.9
56	2y	15	G	2.9

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Mol	Chain	Res	Type	RSRZ
57	2a	1202	G	2.9
1	2A	229	A	2.9
32	1a	1025	U	2.9
34	1c	198	VAL	2.9
41	2j	64	GLU	2.9
57	2a	1532	U	2.9
1	2A	2138	C	2.9
38	2g	32	ARG	2.9
34	2c	57	ILE	2.9
56	2y	19	G	2.9
1	1A	1142	A	2.9
56	1y	21	A	2.9
21	2Z	140	ASP	2.9
38	2g	84	ASN	2.9
33	2b	185	ILE	2.9
57	2a	485	G	2.9
1	2A	614(A)	U	2.8
50	2s	49	ILE	2.8
1	2A	2175	C	2.8
34	1c	87	LEU	2.8
40	2i	88	TYR	2.8
44	2m	13	LYS	2.8
57	2a	1032	G	2.8
1	1A	1072	U	2.8
48	2q	65	ILE	2.8
40	2i	121	ARG	2.8
41	2j	10	GLY	2.8
1	1A	2135	U	2.8
32	1a	1033	G	2.8
57	2a	1220	G	2.8
1	2A	2174	C	2.8
20	2Y	105	ALA	2.8
40	1i	65	VAL	2.8
57	2a	1149	C	2.8
12	2Q	6	ARG	2.8
12	2Q	37	LEU	2.8
28	26	11	LEU	2.8
1	1A	1124	U	2.8
1	2A	2158	A	2.8
29	27	45	ALA	2.8
44	2m	5	ALA	2.8
57	2a	1001	A	2.8

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Mol	Chain	Res	Type	RSRZ
46	2o	65	ARG	2.8
48	2q	43	LEU	2.8
34	2c	50	ALA	2.8
40	2i	15	ALA	2.8
43	2l	15	ARG	2.8
41	2j	68	HIS	2.8
56	1y	36	A	2.8
57	2a	1031	G	2.8
1	1A	1110	C	2.8
45	2n	44	LEU	2.8
1	2A	2132	U	2.8
20	2Y	48	ALA	2.8
1	1A	1133	G	2.8
1	1A	2134	G	2.8
21	2Z	57	ILE	2.8
24	22	60	LEU	2.8
36	2e	13	ILE	2.8
38	2g	16	LEU	2.8
32	1a	163	C	2.8
41	1j	10	GLY	2.8
51	2t	9	ASN	2.8
7	2H	71	LEU	2.7
12	2Q	2	LEU	2.7
34	2c	185	GLY	2.7
1	2A	2801(A)	A	2.7
1	1A	696	C	2.7
21	2Z	50	GLN	2.7
54	1w	15	G	2.7
57	2a	80	G	2.7
51	1t	67	ALA	2.7
51	2t	59	ALA	2.7
29	27	48	LYS	2.7
35	2d	194	LEU	2.7
39	2h	131	GLY	2.7
42	1k	75	TYR	2.7
40	2i	63	ILE	2.7
7	2H	96	ALA	2.7
1	1A	2196	C	2.7
36	2e	81	GLU	2.7
46	1o	88	ARG	2.7
50	2s	82	GLY	2.7
6	2G	11	TYR	2.7

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Mol	Chain	Res	Type	RSRZ
21	2Z	155	LEU	2.7
33	1b	61	LEU	2.7
21	2Z	96	VAL	2.7
52	2u	22	ARG	2.7
1	1A	930	G	2.7
6	2G	19	LEU	2.7
10	2O	41	ALA	2.7
41	2j	37	PRO	2.7
45	2n	30	ALA	2.7
35	1d	3	ARG	2.7
38	2g	3	ARG	2.7
39	1h	112	LEU	2.7
41	2j	54	PHE	2.7
1	2A	898	C	2.7
56	2y	48	C	2.7
1	2A	2169	A	2.7
28	26	54	ILE	2.7
39	1h	86	ILE	2.7
40	2i	119	ALA	2.7
44	2m	71	ARG	2.7
52	2u	23	PRO	2.7
6	2G	34	LEU	2.7
40	2i	79	LEU	2.7
4	2E	77	ILE	2.7
1	1A	2168	C	2.7
42	2k	54	ARG	2.7
56	2y	21	A	2.7
14	2S	33	LYS	2.7
57	2a	1224	G	2.7
12	2Q	1	MET	2.7
7	2H	64	LEU	2.7
12	2Q	5	ARG	2.7
34	1c	14	ILE	2.7
35	1d	158	ILE	2.7
45	2n	58	LYS	2.7
1	1A	2815	C	2.6
32	1a	1027	C	2.6
44	1m	123	ALA	2.6
33	2b	71	VAL	2.6
1	2A	1847	A	2.6
51	2t	20	LEU	2.6
33	2b	55	PHE	2.6

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Mol	Chain	Res	Type	RSRZ
12	2Q	63	LYS	2.6
20	2Y	73	ARG	2.6
35	1d	122	ARG	2.6
31	29	17	ILE	2.6
33	1b	227	GLY	2.6
36	2e	114	GLY	2.6
44	2m	105	THR	2.6
7	2H	113	VAL	2.6
33	2b	81	VAL	2.6
48	2q	9	VAL	2.6
48	2q	23	VAL	2.6
57	2a	202	U	2.6
7	2H	34	GLU	2.6
10	2O	1	MET	2.6
34	2c	33	LEU	2.6
51	1t	24	LEU	2.6
45	2n	32	SER	2.6
50	1s	40	ILE	2.6
45	2n	13	THR	2.6
50	2s	69	HIS	2.6
7	2H	30	LYS	2.6
31	29	2	LYS	2.6
12	1Q	6	ARG	2.6
1	1A	2162	C	2.6
30	28	16	ILE	2.6
36	2e	11	ILE	2.6
39	2h	83	ILE	2.6
20	2Y	63	LYS	2.6
31	29	13	LYS	2.6
42	2k	57	THR	2.6
40	2i	10	ARG	2.6
33	2b	122	PHE	2.6
56	2y	51	U	2.6
34	2c	155	GLY	2.6
43	2l	13	LYS	2.6
6	1G	146	TYR	2.6
43	2l	69	TYR	2.6
5	2F	80	ALA	2.6
19	2X	68	ARG	2.6
57	2a	1286	A	2.6
6	2G	15	VAL	2.6
41	2j	89	ASP	2.6

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Mol	Chain	Res	Type	RSRZ
16	2U	17	ILE	2.6
35	2d	168	ARG	2.6
41	2j	29	ARG	2.6
34	2c	200	ALA	2.6
51	2t	73	HIS	2.6
36	2e	100	VAL	2.6
49	2r	26	LEU	2.6
23	1l	75	GLU	2.6
33	2b	17	PHE	2.6
7	1H	2	SER	2.6
39	2h	87	SER	2.6
18	2W	92	ARG	2.6
56	2y	18	G	2.6
21	1Z	51	ALA	2.6
51	1t	59	ALA	2.6
7	2H	37	VAL	2.6
29	17	46	VAL	2.6
57	2a	89	C	2.6
11	2P	109	GLY	2.6
17	1V	101	GLY	2.6
21	2Z	48	PHE	2.6
42	2k	49	GLY	2.6
57	2a	1503	A	2.6
46	2o	87	ILE	2.5
1	1A	2184	G	2.5
38	2g	154	TYR	2.5
44	1m	87	TYR	2.5
26	14	45	GLY	2.5
35	1d	180	GLY	2.5
56	1y	56	C	2.5
12	2Q	136	ALA	2.5
43	2l	16	GLU	2.5
20	2Y	107	ASP	2.5
57	2a	1020	U	2.5
57	2a	1219	U	2.5
3	2D	51	VAL	2.5
5	2F	170	LEU	2.5
43	2l	51	ALA	2.5
6	2G	3	LEU	2.5
6	2G	62	LEU	2.5
7	2H	115	VAL	2.5
18	1W	92	ARG	2.5

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Mol	Chain	Res	Type	RSRZ
35	2d	157	LEU	2.5
39	1h	93	VAL	2.5
40	2i	26	VAL	2.5
50	2s	37	ARG	2.5
6	2G	141	PHE	2.5
45	2n	36	PHE	2.5
1	2A	2125	G	2.5
57	2a	1131	G	2.5
33	2b	188	ALA	2.5
40	2i	120	ARG	2.5
4	2E	167	VAL	2.5
5	2F	89	VAL	2.5
34	1c	193	TYR	2.5
9	2N	10	GLU	2.5
26	24	57	GLU	2.5
30	28	46	ARG	2.5
40	2i	64	THR	2.5
41	1j	46	ARG	2.5
1	1A	2905	C	2.5
40	2i	72	GLY	2.5
51	2t	24	LEU	2.5
7	2H	47	GLU	2.5
34	2c	206	GLU	2.5
35	2d	179	GLU	2.5
39	2h	19	VAL	2.5
35	1d	110	PHE	2.5
44	1m	121	LYS	2.5
1	2A	887	A	2.5
33	2b	130	ARG	2.5
39	1h	4	ASP	2.5
56	2y	58	A	2.5
20	2Y	58	GLY	2.5
26	24	64	GLY	2.5
33	2b	183	PRO	2.5
58	A	3	PRO	2.5
40	2i	106	ALA	2.5
45	2n	22	THR	2.5
6	1G	63	ILE	2.5
33	2b	214	ILE	2.5
48	2q	59	ILE	2.5
3	2D	257	LEU	2.5
14	2S	58	LEU	2.5

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Mol	Chain	Res	Type	RSRZ
45	2n	46	GLU	2.5
45	2n	53	LEU	2.5
54	2w	2	C	2.5
57	2a	1066	C	2.5
32	1a	1026	G	2.5
33	1b	233	SER	2.5
56	2y	22	G	2.5
32	1a	841	U	2.5
33	2b	135	GLN	2.5
11	2P	75	ILE	2.5
34	2c	149	ALA	2.5
35	2d	158	ILE	2.5
38	2g	27	ILE	2.5
30	28	29	LYS	2.5
5	2F	57	VAL	2.5
45	2n	56	VAL	2.5
11	2P	15	ARG	2.5
1	1A	929	G	2.5
1	2A	2148	G	2.5
57	2a	1356	G	2.5
33	2b	95	GLN	2.4
46	2o	89	GLY	2.4
39	2h	13	ILE	2.4
1	2A	2119	A	2.4
11	2P	83	VAL	2.4
35	2d	105	VAL	2.4
11	2P	51	PHE	2.4
36	2e	45	PHE	2.4
38	2g	149	ARG	2.4
40	2i	75	ASP	2.4
36	2e	20	GLN	2.4
40	1i	6	GLY	2.4
1	1A	2163	G	2.4
40	1i	15	ALA	2.4
54	2w	24	G	2.4
6	2G	140	ILE	2.4
23	21	67	ILE	2.4
33	2b	215	LEU	2.4
34	2c	201	TYR	2.4
32	1a	1286	A	2.4
57	2a	1092	A	2.4
31	29	26	ILE	2.4

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Mol	Chain	Res	Type	RSRZ
45	1n	7	ILE	2.4
1	1A	1105	G	2.4
40	2i	86	VAL	2.4
57	2a	570	G	2.4
43	2l	14	GLY	2.4
1	1A	2180	A	2.4
53	2v	15	A	2.4
1	1A	1121	C	2.4
11	2P	76	LYS	2.4
16	2U	62	ILE	2.4
54	1w	66	U	2.4
56	1y	45	U	2.4
57	2a	1196	U	2.4
38	1g	156	TRP	2.4
43	2l	55	VAL	2.4
1	2A	2110	G	2.4
1	2A	2153	G	2.4
1	2A	2793	G	2.4
48	2q	32	TYR	2.4
52	1u	21	TYR	2.4
52	2u	21	TYR	2.4
6	2G	181	ARG	2.4
30	28	30	ARG	2.4
48	2q	91	ARG	2.4
14	2S	34	HIS	2.4
34	2c	163	ALA	2.4
41	2j	50	ILE	2.4
46	1o	87	ILE	2.4
54	1w	14	A	2.4
57	2a	781	A	2.4
34	2c	13	GLY	2.4
34	2c	55	VAL	2.4
35	1d	105	VAL	2.4
21	2Z	9	TYR	2.4
21	2Z	156	LYS	2.4
33	2b	133	LYS	2.4
41	1j	55	LYS	2.4
7	2H	29	PRO	2.4
1	2A	2168	G	2.4
40	2i	117	HIS	2.4
22	10	2	ALA	2.4
44	2m	90	LEU	2.4

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Mol	Chain	Res	Type	RSRZ
39	2h	134	ILE	2.4
50	2s	40	ILE	2.4
3	2D	15	PHE	2.4
1	1A	1140	U	2.4
22	20	5	LYS	2.4
42	2k	51	LYS	2.4
56	2y	37	A	2.4
32	1a	1028	C	2.4
56	2y	59	U	2.4
34	2c	184	TYR	2.4
6	1G	139	LEU	2.4
20	2Y	83	THR	2.4
21	2Z	150	LEU	2.4
4	1E	77	ILE	2.4
7	2H	31	GLY	2.4
33	2b	201	ILE	2.4
52	1u	11	GLY	2.4
52	2u	12	LYS	2.4
54	1w	1	G	2.4
45	2n	16	PHE	2.4
3	2D	241	PRO	2.3
33	1b	19	HIS	2.3
10	2O	81	ASP	2.3
21	2Z	148	ASP	2.3
40	2i	78	LYS	2.3
41	2j	87	THR	2.3
48	2q	53	LEU	2.3
51	1t	76	ALA	2.3
7	2H	132	ARG	2.3
35	2d	122	ARG	2.3
40	2i	16	ARG	2.3
1	1A	639	G	2.3
1	1A	2174	G	2.3
54	1w	5	G	2.3
57	2a	79	G	2.3
7	2H	175	LYS	2.3
40	1i	117	HIS	2.3
1	1A	933	C	2.3
35	1d	115	ARG	2.3
48	2q	92	ARG	2.3
47	1p	59	TRP	2.3
21	2Z	95	PRO	2.3

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Mol	Chain	Res	Type	RSRZ
26	24	32	TYR	2.3
31	29	12	ASP	2.3
40	2i	21	PRO	2.3
6	2G	128	ARG	2.3
6	2G	134	GLY	2.3
7	2H	82	GLY	2.3
11	2P	45	LEU	2.3
48	1q	97	SER	2.3
48	2q	95	TYR	2.3
56	2y	6	G	2.3
44	1m	102	ARG	2.3
44	1m	103	THR	2.3
20	2Y	60	PHE	2.3
57	2a	978	A	2.3
21	1Z	100	VAL	2.3
36	2e	24	ARG	2.3
48	1q	28	PRO	2.3
51	1t	69	GLY	2.3
42	2k	117	ASN	2.3
56	2y	44	G	2.3
22	20	69	PHE	2.3
1	2A	2804	C	2.3
57	2a	91	C	2.3
1	2A	2170	A	2.3
52	1u	6	ARG	2.3
26	24	54	GLY	2.3
3	2D	37	LEU	2.3
11	1P	105	LEU	2.3
28	26	10	LEU	2.3
4	2E	151	TYR	2.3
35	2d	20	TYR	2.3
50	2s	70	LYS	2.3
43	2l	100	ILE	2.3
47	1p	19	ILE	2.3
21	2Z	86	VAL	2.3
33	2b	136	VAL	2.3
48	2q	96	GLU	2.3
33	2b	21	ARG	2.3
50	2s	81	ARG	2.3
1	2A	2129	C	2.3
57	2a	1039	C	2.3
58	B	2	SER	2.3

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Mol	Chain	Res	Type	RSRZ
33	2b	207	ALA	2.3
45	2n	24	CYS	2.3
22	20	68	GLU	2.3
50	2s	62	ILE	2.3
36	2e	27	ARG	2.3
41	1j	44	VAL	2.3
13	2R	69	ASP	2.3
7	2H	41	MET	2.3
50	2s	15	LEU	2.3
35	1d	111	ALA	2.3
1	1A	2141	A	2.3
1	1A	2812	A	2.3
57	2a	1004	A	2.3
57	2a	1447	A	2.3
7	2H	123	PHE	2.3
34	2c	17	ASP	2.3
48	1q	99	SER	2.2
21	2Z	24	LEU	2.2
51	2t	72	LEU	2.2
44	2m	42	ALA	2.2
58	B	13	ALA	2.2
1	1A	1122	C	2.2
1	2A	2162	G	2.2
3	2D	217	ARG	2.2
6	2G	182	LYS	2.2
14	2S	93	LYS	2.2
54	2w	22	G	2.2
55	1x	68	C	2.2
7	2H	148	ILE	2.2
14	2S	29	PHE	2.2
21	2Z	93	ASP	2.2
12	2Q	33	GLY	2.2
39	2h	95	VAL	2.2
40	2i	65	VAL	2.2
11	2P	35	HIS	2.2
21	2Z	125	LEU	2.2
24	22	24	LEU	2.2
40	1i	121	ARG	2.2
49	1r	73	ALA	2.2
14	2S	5	THR	2.2
1	2A	2137	C	2.2
1	2A	2161	C	2.2

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Mol	Chain	Res	Type	RSRZ
33	1b	28	PHE	2.2
10	2O	58	VAL	2.2
21	2Z	124	ILE	2.2
33	2b	211	ILE	2.2
34	2c	134	ILE	2.2
48	2q	19	VAL	2.2
51	2t	63	ILE	2.2
25	23	26	LEU	2.2
18	2W	90	ARG	2.2
29	27	5	TRP	2.2
29	27	7	PRO	2.2
38	2g	99	LEU	2.2
40	1i	112	LYS	2.2
58	B	3	PRO	2.2
34	2c	21	ARG	2.2
40	1i	76	ALA	2.2
7	2H	129	THR	2.2
50	2s	39	THR	2.2
48	2q	42	TYR	2.2
52	1u	16	GLY	2.2
52	1u	18	TYR	2.2
6	2G	39	ILE	2.2
33	1b	126	GLU	2.2
17	2V	79	VAL	2.2
21	2Z	137	ILE	2.2
3	2D	5	LYS	2.2
47	2p	62	VAL	2.2
50	2s	35	SER	2.2
1	2A	2805	G	2.2
41	2j	90	LEU	2.2
1	1A	2195	A	2.2
38	2g	2	ALA	2.2
53	2v	23	A	2.2
57	2a	1093	A	2.2
4	2E	115	GLY	2.2
9	2N	9	VAL	2.2
25	23	54	VAL	2.2
36	1e	89	ILE	2.2
36	2e	82	VAL	2.2
40	1i	14	VAL	2.2
47	2p	20	VAL	2.2
11	1P	15	ARG	2.2

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Mol	Chain	Res	Type	RSRZ
26	14	55	ARG	2.2
38	2g	78	ARG	2.2
3	2D	182	LEU	2.2
26	24	40	HIS	2.2
50	2s	47	HIS	2.2
34	1c	18	TRP	2.2
1	1A	2173	G	2.2
1	2A	2141	G	2.2
51	2t	30	LYS	2.2
57	2a	793	U	2.2
57	2a	1002	G	2.2
34	2c	23	TYR	2.2
3	2D	18	VAL	2.2
45	2n	41	ARG	2.2
22	20	3	HIS	2.2
39	1h	133	LEU	2.2
1	1A	2161	C	2.2
33	2b	34	ALA	2.2
45	1n	61	TRP	2.2
51	2t	78	ALA	2.2
23	11	35	THR	2.2
1	1A	2131	U	2.2
41	2j	60	ARG	2.2
46	1o	65	ARG	2.2
1	1A	2813	G	2.2
20	2Y	61	ILE	2.2
39	1h	134	ILE	2.2
48	2q	11	VAL	2.2
1	2A	2176	A	2.2
33	1b	128	GLU	2.2
47	1p	7	ALA	2.2
3	1D	263	ARG	2.2
14	2S	17	ARG	2.2
40	2i	42	ARG	2.2
52	1u	15	ARG	2.2
1	1A	2183	C	2.2
16	2U	79	PHE	2.2
35	1d	137	SER	2.1
44	2m	65	LYS	2.1
48	1q	36	ILE	2.1
50	2s	4	SER	2.1
56	1y	47	U	2.1

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Mol	Chain	Res	Type	RSRZ
57	2a	204	U	2.1
38	2g	120	ILE	2.1
7	2H	105	LEU	2.1
35	1d	11	LEU	2.1
35	2d	176	LEU	2.1
43	2l	60	LEU	2.1
1	1A	2181	G	2.1
1	2A	2190	G	2.1
1	2A	2894	G	2.1
33	2b	195	ASP	2.1
1	2A	2114	A	2.1
1	2A	2117	A	2.1
51	2t	101	GLY	2.1
35	2d	73	ARG	2.1
40	2i	122	ALA	2.1
14	2S	57	LYS	2.1
50	2s	6	LYS	2.1
41	1j	47	PHE	2.1
1	1A	2150	C	2.1
11	2P	71	VAL	2.1
14	2S	92	TYR	2.1
33	1b	222	ILE	2.1
42	1k	25	TYR	2.1
6	2G	139	LEU	2.1
23	2l	98	LEU	2.1
46	1o	66	LEU	2.1
54	2w	12	U	2.1
4	2E	10	GLY	2.1
7	2H	97	ARG	2.1
36	2e	25	ARG	2.1
51	2t	83	ARG	2.1
1	1A	554	A	2.1
57	2a	969	A	2.1
6	2G	70	VAL	2.1
7	2H	133	VAL	2.1
7	2H	169	VAL	2.1
34	1c	201	TYR	2.1
1	2A	652(T)	C	2.1
6	2G	133	LEU	2.1
6	2G	152	LEU	2.1
35	2d	146	ILE	2.1
33	1b	228	GLY	2.1

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Mol	Chain	Res	Type	RSRZ
33	2b	137	ARG	2.1
49	2r	54	ARG	2.1
57	2a	1192	C	2.1
28	16	2	ALA	2.1
35	1d	147	ALA	2.1
43	2l	68	ALA	2.1
1	1A	2137	G	2.1
1	1A	2816	G	2.1
1	2A	652(C)	G	2.1
1	2A	2126	A	2.1
7	2H	51	ARG	2.1
33	1b	200	ILE	2.1
35	2d	21	LEU	2.1
36	2e	33	VAL	2.1
39	1h	83	ILE	2.1
39	2h	84	ARG	2.1
41	1j	40	LEU	2.1
43	2l	52	LEU	2.1
43	2l	88	GLY	2.1
51	1t	80	ARG	2.1
32	1a	91	C	2.1
54	2w	75	C	2.1
21	1Z	170	THR	2.1
6	1G	80	PHE	2.1
26	24	59	PHE	2.1
20	2Y	47	LYS	2.1
48	2q	12	SER	2.1
4	2E	52	LEU	2.1
35	2d	70	ILE	2.1
43	2l	7	ILE	2.1
46	1o	57	LEU	2.1
47	2p	38	TYR	2.1
50	2s	27	GLU	2.1
1	1A	1985	U	2.1
38	2g	24	THR	2.1
39	2h	9	MET	2.1
12	2Q	65	PHE	2.1
25	23	29	ARG	2.1
35	1d	118	ARG	2.1
22	20	13	GLY	2.1
4	2E	195	LEU	2.1
17	2V	71	LEU	2.1

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Mol	Chain	Res	Type	RSRZ
18	2W	86	LEU	2.1
30	28	41	ILE	2.1
33	1b	165	VAL	2.1
33	2b	140	HIS	2.1
35	2d	19	LEU	2.1
37	1f	61	LEU	2.1
36	2e	133	TYR	2.1
1	2A	901	A	2.1
56	1y	19	G	2.1
29	27	23	ARG	2.1
38	1g	32	ARG	2.1
46	2o	88	ARG	2.1
56	2y	33	U	2.1
1	1A	2164	C	2.1
26	24	45	GLY	2.1
6	2G	92	VAL	2.1
7	2H	50	VAL	2.1
20	1Y	45	VAL	2.1
21	2Z	157	LEU	2.1
35	1d	101	LEU	2.1
39	1h	2	LEU	2.1
46	2o	31	LEU	2.1
35	1d	5	ILE	2.1
38	1g	120	ILE	2.1
43	2l	5	PRO	2.1
19	2X	69	TYR	2.1
38	2g	40	ALA	2.1
33	2b	94	ASN	2.1
46	2o	64	ARG	2.1
52	1u	9	ARG	2.1
41	2j	15	THR	2.1
1	2A	2100	G	2.0
22	20	76	GLY	2.1
38	2g	103	TRP	2.1
39	2h	90	GLY	2.1
56	2y	63	G	2.0
1	1A	2807	C	2.0
1	2A	280	C	2.0
34	1c	196	LEU	2.0
39	1h	10	LEU	2.0
39	2h	112	LEU	2.0
4	2E	132	HIS	2.0

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Mol	Chain	Res	Type	RSRZ
21	2Z	71	VAL	2.0
30	28	7	HIS	2.0
54	1w	13	C	2.0
57	2a	1114	C	2.0
9	2N	44	PRO	2.0
42	1k	95	ILE	2.0
47	2p	39	TYR	2.0
3	2D	273	ARG	2.0
35	2d	115	ARG	2.0
41	1j	66	ARG	2.0
7	2H	32	GLU	2.0
7	2H	46	GLU	2.0
33	2b	35	GLU	2.0
7	2H	78	GLY	2.0
44	2m	103	THR	2.0
26	24	42	PHE	2.0
41	2j	80	LYS	2.0
1	2A	2109	U	2.0
1	1A	2803	A	2.0
19	2X	43	VAL	2.0
40	2i	108	VAL	2.0
41	1j	72	VAL	2.0
48	1q	65	ILE	2.0
57	2a	1061	G	2.0
44	2m	88	ARG	2.0
18	1W	112	GLY	2.0
24	22	62	THR	2.0
34	2c	80	GLY	2.0
36	2e	99	GLY	2.0
26	14	49	PHE	2.0
34	2c	22	TRP	2.0
39	2h	133	LEU	2.0
6	2G	137	GLU	2.0
7	2H	95	ARG	2.0
11	2P	50	ARG	2.0
21	2Z	121	HIS	2.0
31	29	9	ARG	2.0
34	1c	179	ARG	2.0
36	2e	106	PRO	2.0
41	2j	46	ARG	2.0
25	23	13	ILE	2.0
1	2A	899	A	2.0

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Mol	Chain	Res	Type	RSRZ
1	2A	652(U)	G	2.0
5	2F	49	ALA	2.0
31	29	33	LYS	2.0
57	2a	1018	C	2.0
16	2U	25	TRP	2.0
30	28	34	TRP	2.0
21	2Z	76	LEU	2.0
46	2o	56	LEU	2.0
47	2p	73	LEU	2.0
47	2p	74	LEU	2.0
50	1s	15	LEU	2.0
7	2H	36	PRO	2.0
7	2H	125	VAL	2.0
14	2S	14	VAL	2.0
31	29	23	VAL	2.0
44	2m	15	VAL	2.0
45	2n	11	LYS	2.0

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

In the following table, the Atoms column lists the number of modelled atoms in the group and the number defined in the chemical component dictionary. LLDF column lists the quality of electron density of the group with respect to its neighbouring residues in protein, DNA or RNA chains. The B-factors column lists the minimum, median, 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	LLDF	B-factors(Å ²)	Q<0.9
55	5MC	2x	32	21/22	0.95	0.21	-	62,66,70,75	0
56	PSU	2y	32	20/21	0.80	0.25	-	74,82,94,96	0
55	5MU	2x	54	21/22	0.94	0.25	-	70,77,80,92	0
32	5MC	1a	967	21/22	0.97	0.22	-	42,45,51,54	0
1	5MU	2A	1939	21/22	0.98	0.25	-	30,33,39,42	0
58	BB9	B	6	6/7	0.98	0.15	-	23,30,33,35	0
1	4OC	2A	1920	21/23	0.96	0.22	-	50,55,58,59	0
43	0TD	1l	92	10/11	0.92	0.23	-	41,42,49,76	0
55	4SU	2x	8	20/21	0.92	0.18	-	70,75,79,81	0
32	2MG	1a	1207	24/25	0.96	0.17	-	50,60,65,66	0
54	4SU	1w	8	20/21	0.83	0.20	-	72,81,89,91	0
32	G7M	1a	527	24/25	0.97	0.20	-	31,38,45,46	0
58	BB9	A	17	6/7	0.77	0.52	-	54,69,80,92	0
1	OMG	2A	2251	24/25	0.98	0.23	-	33,37,41,47	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
57	PSU	2a	516	20/21	0.92	0.19	-	58,66,72,72	0
57	5MC	2a	967	21/22	0.93	0.23	-	60,65,76,77	0
1	2MU	1A	2564	21/23	0.98	0.24	-	16,22,26,31	0
32	MA6	1a	1519	24/25	0.98	0.27	-	30,35,38,42	0
32	MA6	1a	1518	24/25	0.98	0.25	-	25,33,36,39	0
55	PSU	2x	55	20/21	0.90	0.18	-	69,73,79,80	0
58	BB9	B	9	6/7	0.98	0.23	-	32,35,36,37	0
56	PSU	1y	32	20/21	0.90	0.25	-	70,78,86,87	0
57	MA6	2a	1518	24/25	0.95	0.28	-	42,58,62,63	0
54	G7M	1w	46	24/25	0.82	0.20	-	70,80,100,123	0
32	5MC	1a	1407	21/22	0.98	0.23	-	27,33,37,40	0
54	PSU	2w	32	20/21	0.93	0.21	-	70,77,85,91	0
57	UR3	2a	1498	21/22	0.96	0.23	-	44,50,56,60	0
56	4SU	2y	8	20/21	0.85	0.23	-	85,90,98,107	0
54	MIA	2w	37	25/30	0.95	0.24	-	63,70,76,77	0
32	5MC	1a	1400	21/22	0.98	0.21	-	30,41,43,48	0
1	5MC	2A	1942	21/22	0.97	0.21	-	39,50,56,63	0
1	2MA	2A	2503	23/24	0.97	0.21	-	26,30,33,36	0
32	UR3	1a	1498	21/22	0.99	0.24	-	27,33,37,39	0
54	5MU	1w	54	21/22	0.96	0.17	-	45,59,64,65	0
56	4SU	1y	8	20/21	0.81	0.26	-	83,88,102,104	0
32	PSU	1a	516	20/21	0.96	0.21	-	45,49,50,51	0
1	PSU	2A	2605	20/21	0.97	0.22	-	24,33,36,37	0
56	5MU	2y	54	21/22	0.71	0.36	-	86,91,105,122	0
1	4OC	1A	1942	21/23	0.98	0.23	-	29,34,37,40	0
55	5MU	1x	54	21/22	0.96	0.17	-	54,61,65,73	0
1	5MU	1A	1937	21/22	0.94	0.21	-	39,44,52,54	0
32	5MC	1a	1404	21/22	0.98	0.21	-	29,33,37,40	0
56	PSU	2y	39	20/21	0.87	0.32	-	72,80,86,98	0
57	M2G	2a	966	25/26	0.91	0.26	-	56,62,75,78	0
57	5MC	2a	1407	21/22	0.95	0.23	-	38,49,54,63	0
54	G7M	2w	46	24/25	0.73	0.26	-	80,88,101,118	0
56	PSU	1y	39	20/21	0.92	0.20	-	69,73,78,82	0
1	PSU	2A	1911	20/21	0.94	0.21	-	50,55,60,77	0
58	BB9	B	17	6/7	0.65	0.76	-	68,79,86,100	0
54	MIA	1w	37	29/30	0.97	0.24	-	36,44,57,62	0
1	PSU	2A	1917	20/21	0.94	0.18	-	51,60,68,71	0
1	OMG	1A	2263	24/25	0.99	0.21	-	15,19,22,25	0
1	5MU	2A	1915	21/22	0.95	0.17	-	58,65,68,73	0
1	PSU	1A	2617	20/21	0.98	0.25	-	15,18,23,24	0
56	PSU	2y	55	20/21	0.56	0.42	-	89,99,107,116	0
57	G7M	2a	527	24/25	0.95	0.20	-	46,55,64,74	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
56	G7M	2y	46	24/25	0.74	0.25	-	82,90,99,118	0
57	MA6	2a	1519	24/25	0.95	0.28	-	47,54,62,63	0
57	5MC	2a	1404	21/22	0.95	0.20	-	46,51,59,64	0
32	M2G	1a	966	25/26	0.97	0.23	-	39,44,51,60	0
57	5MC	2a	1400	21/22	0.94	0.27	-	58,64,70,74	0
55	PSU	1x	55	20/21	0.95	0.19	-	52,56,67,69	0
55	5MC	1x	32	21/22	0.97	0.22	-	39,46,52,57	0
57	4OC	2a	1402	22/23	0.93	0.24	-	48,57,64,65	0
1	5MC	1A	1984	21/22	0.98	0.18	-	18,26,30,37	0
57	2MG	2a	1207	24/25	0.89	0.19	-	72,80,87,88	0
1	PSU	1A	1939	20/21	0.97	0.22	-	36,41,45,46	0
56	5MU	1y	54	21/22	0.78	0.23	-	76,85,92,104	0
56	PSU	1y	55	20/21	0.75	0.27	-	84,90,100,105	0
58	BB9	A	9	6/7	0.98	0.18	-	14,16,16,19	0
55	4SU	1x	8	20/21	0.96	0.18	-	48,55,66,78	0
54	PSU	1w	55	20/21	0.93	0.20	-	53,69,78,80	0
1	2MU	2A	2552	21/23	0.98	0.23	-	32,37,41,46	0
58	BB9	A	6	6/7	0.99	0.18	-	20,21,22,23	0
1	2MA	1A	2515	23/24	0.99	0.24	-	10,14,17,22	0
56	G7M	1y	46	24/25	0.83	0.26	-	80,88,98,108	0
1	PSU	1A	1933	20/21	0.97	0.21	-	31,38,41,42	0
54	PSU	1w	39	20/21	0.97	0.21	-	44,50,59,59	0
32	4OC	1a	1402	22/23	0.96	0.21	-	33,38,43,44	0
1	5MU	1A	1961	21/22	0.99	0.24	-	15,20,24,26	0
54	PSU	2w	55	20/21	0.85	0.22	-	73,83,90,98	0
54	4SU	2w	8	20/21	0.83	0.24	-	82,88,103,107	0
54	5MU	2w	54	21/22	0.92	0.15	-	67,75,81,86	0
1	5MC	1A	1964	21/22	0.98	0.24	-	24,31,36,43	0
43	0TD	2l	92	10/11	0.90	0.28	-	58,62,66,84	0
54	PSU	2w	39	20/21	0.95	0.24	-	65,72,79,80	0
54	PSU	1w	32	20/21	0.95	0.20	-	53,59,72,72	0
1	5MC	2A	1962	21/22	0.97	0.21	-	31,43,52,61	0

6.3 Carbohydrates [i](#)

There are no carbohydrates in this entry.

6.4 Ligands [i](#)

In the following table, the Atoms column lists the number of modelled atoms in the group and the number defined in the chemical component dictionary. LLDF column lists the quality of electron

density of the group with respect to its neighbouring residues in protein, DNA or RNA chains. The B-factors column lists the minimum, median, 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	LLDF	B-factors(Å ²)	Q<0.9
59	MG	1A	3165	1/1	0.96	0.72	63.45	22,22,22,22	0
59	MG	1A	3413	1/1	0.89	0.61	61.54	49,49,49,49	0
59	MG	1A	4135	1/1	0.86	0.62	52.16	40,40,40,40	0
59	MG	1A	3160	1/1	0.95	0.78	50.46	36,36,36,36	0
59	MG	1A	3026	1/1	0.97	0.65	48.01	28,28,28,28	0
59	MG	1A	3529	1/1	0.96	0.85	46.07	33,33,33,33	0
59	MG	1A	3198	1/1	0.94	0.59	42.19	27,27,27,27	0
59	MG	1A	4214	1/1	0.97	0.72	36.66	36,36,36,36	0
59	MG	1A	3389	1/1	0.91	0.60	35.48	36,36,36,36	0
59	MG	1N	3007	1/1	0.97	0.53	35.40	49,49,49,49	0
59	MG	1A	3808	1/1	0.97	0.40	33.18	33,33,33,33	0
59	MG	1A	3327	1/1	0.81	0.98	33.14	42,42,42,42	0
59	MG	2A	3056	1/1	0.95	0.48	32.69	56,56,56,56	0
59	MG	1A	3186	1/1	0.87	0.71	31.87	35,35,35,35	0
59	MG	1A	3370	1/1	0.76	0.73	31.79	51,51,51,51	0
59	MG	1A	3289	1/1	0.89	0.72	31.03	29,29,29,29	0
59	MG	1A	3027	1/1	0.99	0.45	28.22	30,30,30,30	0
59	MG	2D	305	1/1	0.87	0.71	27.84	44,44,44,44	0
59	MG	1A	3035	1/1	0.98	0.72	26.14	24,24,24,24	0
59	MG	2A	3200	1/1	0.90	0.58	25.82	37,37,37,37	0
59	MG	1A	3317	1/1	0.95	0.69	23.49	32,32,32,32	0
59	MG	1R	202	1/1	0.96	0.84	23.39	33,33,33,33	0
59	MG	1S	3001	1/1	0.88	0.74	22.49	39,39,39,39	0
59	MG	1a	3411	1/1	0.93	0.70	21.92	47,47,47,47	0
59	MG	1A	3466	1/1	0.97	0.52	21.68	24,24,24,24	0
59	MG	2U	204	1/1	0.97	0.75	21.35	49,49,49,49	0
59	MG	1A	3197	1/1	0.96	0.58	21.32	32,32,32,32	0
59	MG	1A	3134	1/1	0.96	0.43	21.30	24,24,24,24	0
59	MG	2A	3091	1/1	0.90	0.39	21.16	44,44,44,44	0
59	MG	1A	3371	1/1	0.96	0.55	20.69	39,39,39,39	0
59	MG	2D	304	1/1	0.84	1.22	20.35	44,44,44,44	0
59	MG	2F	303	1/1	0.80	1.18	20.29	54,54,54,54	0
59	MG	1A	3106	1/1	0.91	0.61	19.91	33,33,33,33	0
59	MG	2A	3845	1/1	0.81	1.19	19.84	49,49,49,49	0
59	MG	2A	3076	1/1	0.92	0.55	19.47	37,37,37,37	0
59	MG	1U	205	1/1	0.97	0.55	19.34	29,29,29,29	0
59	MG	1A	3144	1/1	0.96	0.73	18.91	35,35,35,35	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3748	1/1	0.94	0.54	18.27	24,24,24,24	0
59	MG	1A	3108	1/1	0.96	0.41	18.09	26,26,26,26	0
59	MG	2A	3366	1/1	0.93	1.25	17.97	70,70,70,70	0
59	MG	1A	3798	1/1	0.76	0.60	17.77	44,44,44,44	0
59	MG	1A	3403	1/1	0.91	0.47	17.77	33,33,33,33	0
59	MG	1A	4208	1/1	0.92	0.72	17.73	29,29,29,29	0
59	MG	1A	3218	1/1	0.94	0.41	17.72	38,38,38,38	0
59	MG	1P	202	1/1	0.94	0.65	17.36	25,25,25,25	0
59	MG	2A	3847	1/1	0.96	0.74	17.35	37,37,37,37	0
59	MG	1A	3535	1/1	0.97	0.47	17.23	37,37,37,37	0
59	MG	1A	4019	1/1	0.86	0.64	16.66	33,33,33,33	0
59	MG	2A	3187	1/1	0.98	1.04	16.61	37,37,37,37	0
59	MG	1A	4172	1/1	0.95	0.54	16.53	44,44,44,44	0
59	MG	1A	3531	1/1	0.99	0.52	16.39	28,28,28,28	0
59	MG	2A	3345	1/1	0.86	0.47	16.38	59,59,59,59	0
59	MG	1A	3128	1/1	0.94	0.36	16.23	30,30,30,30	0
59	MG	1A	3147	1/1	0.97	0.61	15.84	26,26,26,26	0
59	MG	1A	3164	1/1	0.93	0.58	15.77	27,27,27,27	0
59	MG	1A	4199	1/1	0.99	0.69	15.62	32,32,32,32	0
59	MG	1A	3159	1/1	0.89	0.47	15.24	32,32,32,32	0
59	MG	1A	4228	1/1	0.94	0.62	15.02	34,34,34,34	0
59	MG	1D	310	1/1	0.88	0.50	14.86	43,43,43,43	0
59	MG	1A	3036	1/1	0.97	0.44	14.49	23,23,23,23	0
59	MG	1A	3129	1/1	0.88	0.36	14.32	36,36,36,36	0
59	MG	1A	4190	1/1	0.92	0.65	14.00	42,42,42,42	0
59	MG	1A	4174	1/1	0.62	0.58	13.75	52,52,52,52	0
59	MG	1a	3576	1/1	0.94	0.56	13.61	53,53,53,53	0
59	MG	1B	211	1/1	0.98	0.33	13.47	52,52,52,52	0
59	MG	2U	203	1/1	0.94	0.83	13.46	56,56,56,56	0
59	MG	1F	310	1/1	0.89	0.50	13.26	47,47,47,47	0
59	MG	1N	3003	1/1	0.98	0.55	13.22	37,37,37,37	0
59	MG	1D	313	1/1	0.91	0.61	13.11	34,34,34,34	0
59	MG	1A	4177	1/1	0.97	0.50	12.84	29,29,29,29	0
59	MG	1A	4195	1/1	0.91	0.44	12.80	27,27,27,27	0
59	MG	1A	3806	1/1	0.80	0.51	12.61	47,47,47,47	0
59	MG	1A	3065	1/1	0.94	0.59	12.59	37,37,37,37	0
59	MG	1A	3427	1/1	0.79	0.41	12.48	43,43,43,43	0
59	MG	1A	3739	1/1	0.96	0.50	12.43	26,26,26,26	0
59	MG	1A	3804	1/1	0.97	0.43	12.27	27,27,27,27	0
59	MG	1A	3354	1/1	0.91	0.36	12.04	37,37,37,37	0
59	MG	2A	3842	1/1	0.94	0.78	11.90	46,46,46,46	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3148	1/1	0.97	0.46	11.89	26,26,26,26	0
59	MG	1A	3532	1/1	0.95	0.42	11.85	25,25,25,25	0
59	MG	2A	3286	1/1	0.96	0.31	11.80	58,58,58,58	0
59	MG	1A	4219	1/1	0.96	0.41	11.15	32,32,32,32	0
59	MG	1A	3788	1/1	0.87	0.36	10.74	27,27,27,27	0
59	MG	1U	202	1/1	0.89	0.42	10.70	27,27,27,27	0
59	MG	1a	3340	1/1	0.91	0.36	10.69	53,53,53,53	0
59	MG	18	102	1/1	0.95	0.50	10.58	37,37,37,37	0
59	MG	1A	3831	1/1	0.97	0.38	10.56	17,17,17,17	0
59	MG	1A	3209	1/1	0.90	0.39	10.47	41,41,41,41	0
59	MG	1A	3373	1/1	0.96	0.33	10.43	29,29,29,29	0
59	MG	1N	3006	1/1	0.94	0.63	10.42	41,41,41,41	0
59	MG	1A	4211	1/1	0.98	0.33	10.41	23,23,23,23	0
59	MG	1A	3140	1/1	0.89	0.45	10.39	28,28,28,28	0
59	MG	1X	3003	1/1	0.99	0.55	10.39	32,32,32,32	0
59	MG	2A	3844	1/1	0.93	0.66	10.03	48,48,48,48	0
59	MG	1A	4200	1/1	0.96	0.55	9.98	27,27,27,27	0
59	MG	1S	3002	1/1	0.94	0.47	9.82	48,48,48,48	0
59	MG	1A	4171	1/1	0.94	0.45	9.76	26,26,26,26	0
59	MG	1A	4197	1/1	0.92	0.58	9.57	25,25,25,25	0
59	MG	1a	3360	1/1	0.91	0.45	9.43	45,45,45,45	0
59	MG	1O	3002	1/1	0.96	0.50	9.30	41,41,41,41	0
59	MG	2A	3030	1/1	0.92	0.41	9.19	50,50,50,50	0
59	MG	2A	3824	1/1	0.95	0.38	9.15	43,43,43,43	0
59	MG	1A	4206	1/1	0.99	0.44	9.14	21,21,21,21	0
59	MG	1A	3514	1/1	0.94	0.29	9.04	53,53,53,53	0
59	MG	1Y	504	1/1	0.97	0.59	8.98	45,45,45,45	0
59	MG	1A	3802	1/1	0.94	0.44	8.78	31,31,31,31	0
59	MG	1U	201	1/1	0.96	0.39	8.73	27,27,27,27	0
59	MG	1A	3142	1/1	0.97	0.42	8.70	36,36,36,36	0
59	MG	1P	201	1/1	0.97	0.43	8.50	23,23,23,23	0
59	MG	1A	3434	1/1	0.98	0.35	8.23	21,21,21,21	0
59	MG	1A	4180	1/1	0.97	0.40	8.19	28,28,28,28	0
59	MG	2A	3830	1/1	0.98	0.58	8.10	32,32,32,32	0
59	MG	1A	4196	1/1	0.95	0.48	8.05	29,29,29,29	0
59	MG	1A	3290	1/1	0.95	0.39	8.01	37,37,37,37	0
59	MG	1A	4182	1/1	0.98	0.40	7.89	23,23,23,23	0
59	MG	1U	203	1/1	0.96	0.36	7.89	22,22,22,22	0
59	MG	1A	4212	1/1	0.98	0.28	7.82	32,32,32,32	0
59	MG	1A	3244	1/1	0.99	0.41	7.70	24,24,24,24	0
59	MG	2A	3831	1/1	0.90	0.84	7.63	39,39,39,39	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	4209	1/1	0.98	0.31	7.53	21,21,21,21	0
59	MG	1x	117	1/1	0.83	0.33	7.44	63,63,63,63	0
59	MG	1A	4169	1/1	0.97	0.33	7.40	22,22,22,22	0
59	MG	1a	3383	1/1	0.99	0.29	7.38	43,43,43,43	0
59	MG	1A	3207	1/1	0.95	0.44	7.36	24,24,24,24	0
59	MG	1A	3037	1/1	0.96	0.41	7.27	37,37,37,37	0
59	MG	2A	3069	1/1	0.85	0.55	7.22	45,45,45,45	0
59	MG	1A	3435	1/1	0.94	0.39	7.05	28,28,28,28	0
59	MG	1A	3995	1/1	0.87	0.31	6.85	49,49,49,49	0
59	MG	2X	3001	1/1	0.96	0.43	6.83	59,59,59,59	0
59	MG	1A	4203	1/1	0.96	0.53	6.72	33,33,33,33	0
59	MG	1A	4225	1/1	0.94	0.51	6.72	34,34,34,34	0
59	MG	1D	311	1/1	0.97	0.43	6.55	19,19,19,19	0
59	MG	1E	313	1/1	0.94	0.39	6.43	34,34,34,34	0
59	MG	2A	3155	1/1	0.96	0.27	6.42	44,44,44,44	0
59	MG	10	105	1/1	0.94	0.35	6.24	40,40,40,40	0
59	MG	1A	3011	1/1	0.94	0.35	6.21	16,16,16,16	0
59	MG	2A	3066	1/1	0.94	0.30	6.16	54,54,54,54	0
59	MG	2A	3108	1/1	0.97	0.26	5.86	43,43,43,43	0
59	MG	1A	3161	1/1	0.96	0.41	5.81	25,25,25,25	0
59	MG	1D	301	1/1	0.93	0.41	5.77	24,24,24,24	0
59	MG	1A	4202	1/1	0.95	0.40	5.69	27,27,27,27	0
59	MG	1W	203	1/1	0.97	0.35	5.56	29,29,29,29	0
59	MG	2A	3102	1/1	0.95	0.42	5.55	41,41,41,41	0
59	MG	1A	4227	1/1	0.96	0.37	5.52	32,32,32,32	0
59	MG	2a	3117	1/1	0.97	0.24	5.45	53,53,53,53	0
59	MG	2A	3495	1/1	0.95	0.29	5.43	37,37,37,37	0
59	MG	2A	3514	1/1	0.97	0.44	5.36	35,35,35,35	0
59	MG	1A	4218	1/1	0.97	0.36	5.34	24,24,24,24	0
59	MG	1A	4220	1/1	0.97	0.31	5.32	24,24,24,24	0
59	MG	1A	3288	1/1	0.89	0.27	5.25	50,50,50,50	0
59	MG	1D	307	1/1	0.98	0.39	5.21	24,24,24,24	0
59	MG	1A	3082	1/1	0.89	0.22	5.14	48,48,48,48	0
59	MG	1D	308	1/1	0.93	0.38	5.13	27,27,27,27	0
59	MG	1Z	301	1/1	0.96	0.33	5.13	34,34,34,34	0
59	MG	1D	304	1/1	0.96	0.49	5.12	38,38,38,38	0
59	MG	2A	3136	1/1	0.94	0.49	5.11	33,33,33,33	0
59	MG	2A	3532	1/1	0.92	0.27	5.00	53,53,53,53	0
59	MG	1A	3204	1/1	0.95	0.36	4.96	23,23,23,23	0
59	MG	1A	3177	1/1	0.78	0.32	4.92	47,47,47,47	0
59	MG	1A	3825	1/1	0.95	0.36	4.79	27,27,27,27	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3234	1/1	0.91	0.30	4.76	33,33,33,33	0
59	MG	1A	4170	1/1	0.98	0.32	4.75	23,23,23,23	0
59	MG	1T	8002	1/1	0.90	0.33	4.74	51,51,51,51	0
59	MG	2A	3823	1/1	0.95	0.40	4.63	47,47,47,47	0
59	MG	1A	3139	1/1	0.87	0.31	4.62	29,29,29,29	0
59	MG	1A	3702	1/1	0.80	0.29	4.59	20,20,20,20	0
59	MG	1A	3417	1/1	0.95	0.34	4.55	36,36,36,36	0
59	MG	1A	3277	1/1	0.89	0.28	4.40	48,48,48,48	0
59	MG	1a	3426	1/1	0.91	0.27	4.40	55,55,55,55	0
59	MG	2A	3331	1/1	0.94	0.29	4.38	58,58,58,58	0
59	MG	1A	3105	1/1	0.93	0.27	3.98	30,30,30,30	0
59	MG	1A	3029	1/1	0.95	0.28	3.94	29,29,29,29	0
59	MG	1A	3170	1/1	0.98	0.23	3.73	10,10,10,10	0
59	MG	1a	3313	1/1	0.97	0.27	3.67	42,42,42,42	0
59	MG	1A	4210	1/1	0.79	0.30	3.64	30,30,30,30	0
59	MG	1A	3675	1/1	0.96	0.29	3.63	21,21,21,21	0
59	MG	12	3002	1/1	0.95	0.34	3.60	36,36,36,36	0
59	MG	1f	3003	1/1	0.83	0.33	3.52	57,57,57,57	0
59	MG	1F	305	1/1	0.95	0.28	3.44	27,27,27,27	0
59	MG	2B	3001	1/1	0.87	0.23	3.42	53,53,53,53	0
59	MG	1x	103	1/1	0.94	0.33	3.40	60,60,60,60	0
59	MG	2A	3040	1/1	0.90	0.38	3.40	40,40,40,40	0
59	MG	1A	3090	1/1	0.99	0.30	3.31	29,29,29,29	0
59	MG	2T	202	1/1	0.90	0.36	3.30	61,61,61,61	0
59	MG	2A	3618	1/1	0.95	0.55	3.27	36,36,36,36	0
59	MG	2A	3774	1/1	0.76	0.28	3.26	64,64,64,64	0
59	MG	1A	4198	1/1	0.97	0.28	3.25	22,22,22,22	0
59	MG	2A	3002	1/1	0.88	0.23	3.07	46,46,46,46	0
59	MG	2A	3014	1/1	0.94	0.25	3.02	29,29,29,29	0
59	MG	1A	3103	1/1	0.89	0.25	3.02	34,34,34,34	0
59	MG	1F	304	1/1	0.93	0.28	2.95	21,21,21,21	0
59	MG	1A	4087	1/1	0.89	0.23	2.91	34,34,34,34	0
59	MG	1B	223	1/1	0.89	0.21	2.89	45,45,45,45	0
59	MG	1A	3464	1/1	0.77	0.33	2.89	51,51,51,51	0
59	MG	1A	4020	1/1	0.94	0.30	2.86	34,34,34,34	0
59	MG	1A	3467	1/1	0.96	0.29	2.85	28,28,28,28	0
59	MG	1A	3844	1/1	0.80	0.26	2.83	35,35,35,35	0
59	MG	1A	3450	1/1	0.97	0.24	2.78	31,31,31,31	0
59	MG	1a	3333	1/1	0.87	0.27	2.78	54,54,54,54	0
59	MG	1A	3294	1/1	0.89	0.27	2.78	30,30,30,30	0
59	MG	2A	3355	1/1	0.94	0.32	2.65	44,44,44,44	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3301	1/1	0.89	0.28	2.62	41,41,41,41	0
59	MG	1f	3001	1/1	0.98	0.27	2.54	29,29,29,29	0
59	MG	1A	3033	1/1	0.86	0.21	2.50	57,57,57,57	0
59	MG	2A	3667	1/1	0.96	0.30	2.48	47,47,47,47	0
59	MG	1W	204	1/1	0.95	0.28	2.45	32,32,32,32	0
59	MG	1A	3754	1/1	0.92	0.24	2.42	32,32,32,32	0
59	MG	2a	3112	1/1	0.73	0.20	2.42	51,51,51,51	0
59	MG	1O	3001	1/1	0.76	0.36	2.32	54,54,54,54	0
59	MG	1N	3009	1/1	0.95	0.27	2.24	28,28,28,28	0
59	MG	1A	3800	1/1	0.78	0.26	2.21	30,30,30,30	0
60	ZN	16	102	1/1	0.99	0.23	2.20	36,36,36,36	0
59	MG	1A	4207	1/1	0.91	0.28	2.18	26,26,26,26	0
59	MG	2a	3188	1/1	0.96	0.26	2.12	64,64,64,64	0
59	MG	1A	3523	1/1	0.89	0.26	2.10	37,37,37,37	0
59	MG	1X	3001	1/1	0.91	0.28	2.08	30,30,30,30	0
59	MG	1a	3335	1/1	0.78	0.20	2.08	57,57,57,57	0
59	MG	2a	3085	1/1	0.85	0.26	2.07	69,69,69,69	0
59	MG	2a	3026	1/1	0.96	0.26	2.07	47,47,47,47	0
59	MG	1D	305	1/1	0.97	0.27	2.05	29,29,29,29	0
59	MG	2A	3772	1/1	0.96	0.25	2.00	35,35,35,35	0
59	MG	2A	3138	1/1	0.93	0.20	1.85	45,45,45,45	0
59	MG	1A	3973	1/1	0.91	0.24	1.73	28,28,28,28	0
59	MG	2A	3486	1/1	0.93	0.16	1.70	61,61,61,61	0
59	MG	2A	3541	1/1	0.94	0.25	1.67	30,30,30,30	0
59	MG	2A	3159	1/1	0.89	0.21	1.66	62,62,62,62	0
60	ZN	19	102	1/1	0.99	0.24	1.56	40,40,40,40	0
59	MG	2A	3287	1/1	0.72	0.18	1.55	65,65,65,65	0
59	MG	1b	3001	1/1	0.94	0.29	1.54	74,74,74,74	0
59	MG	1D	309	1/1	0.98	0.28	1.54	37,37,37,37	0
59	MG	2A	3837	1/1	0.83	0.32	1.46	61,61,61,61	0
59	MG	1A	4107	1/1	0.99	0.21	1.46	17,17,17,17	0
59	MG	2S	8001	1/1	0.61	0.38	1.45	73,73,73,73	0
59	MG	1D	306	1/1	0.88	0.24	1.40	40,40,40,40	0
59	MG	1a	3570	1/1	0.94	0.23	1.32	47,47,47,47	0
59	MG	2A	3619	1/1	0.54	0.25	1.24	60,60,60,60	0
59	MG	2A	3126	1/1	0.92	0.20	1.24	59,59,59,59	0
59	MG	1a	3346	1/1	0.94	0.22	1.21	57,57,57,57	0
59	MG	1Q	202	1/1	0.98	0.28	1.16	37,37,37,37	0
59	MG	2F	302	1/1	0.94	0.30	1.10	41,41,41,41	0
59	MG	2A	3279	1/1	0.87	0.22	0.99	43,43,43,43	0
59	MG	1A	3534	1/1	0.93	0.22	0.98	38,38,38,38	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1a	3362	1/1	0.92	0.19	0.98	51,51,51,51	0
59	MG	1A	3363	1/1	0.93	0.33	0.97	30,30,30,30	0
59	MG	1N	3004	1/1	0.92	0.27	0.97	41,41,41,41	0
59	MG	2A	3843	1/1	0.95	0.23	0.96	63,63,63,63	0
59	MG	1A	4143	1/1	0.98	0.21	0.90	16,16,16,16	0
59	MG	2R	201	1/1	0.76	0.21	0.86	63,63,63,63	0
59	MG	1a	3519	1/1	0.91	0.25	0.84	65,65,65,65	0
60	ZN	25	101	1/1	0.99	0.21	0.77	49,49,49,49	0
59	MG	1A	3584	1/1	0.90	0.22	0.69	33,33,33,33	0
59	MG	1X	3004	1/1	0.94	0.23	0.66	49,49,49,49	0
59	MG	2A	3134	1/1	0.93	0.22	0.64	58,58,58,58	0
59	MG	1A	3775	1/1	0.85	0.23	0.62	46,46,46,46	0
59	MG	1a	3463	1/1	0.96	0.21	0.58	44,44,44,44	0
59	MG	2A	3383	1/1	0.96	0.20	0.55	31,31,31,31	0
59	MG	1A	4223	1/1	0.94	0.22	0.53	39,39,39,39	0
60	ZN	15	101	1/1	0.99	0.21	0.52	34,34,34,34	0
59	MG	1a	3573	1/1	0.96	0.21	0.51	37,37,37,37	0
59	MG	2A	3011	1/1	0.97	0.25	0.50	38,38,38,38	0
59	MG	13	3002	1/1	0.93	0.25	0.50	37,37,37,37	0
59	MG	1A	4204	1/1	0.97	0.25	0.49	29,29,29,29	0
59	MG	1A	3696	1/1	0.95	0.23	0.49	30,30,30,30	0
59	MG	1A	3145	1/1	0.94	0.24	0.49	44,44,44,44	0
59	MG	2Q	3004	1/1	0.83	0.26	0.49	45,45,45,45	0
59	MG	23	102	1/1	0.88	0.28	0.47	45,45,45,45	0
59	MG	1V	201	1/1	0.96	0.21	0.34	33,33,33,33	0
59	MG	1A	3038	1/1	0.93	0.25	0.32	30,30,30,30	0
59	MG	1A	3149	1/1	0.95	0.23	0.25	20,20,20,20	0
59	MG	1A	4091	1/1	0.81	0.23	0.23	42,42,42,42	0
59	MG	1A	3054	1/1	0.96	0.18	0.21	24,24,24,24	0
59	MG	2A	3323	1/1	0.90	0.20	0.16	47,47,47,47	0
59	MG	1a	3328	1/1	0.96	0.23	0.14	17,17,17,17	0
59	MG	2A	3625	1/1	0.92	0.22	0.12	67,67,67,67	0
59	MG	1B	220	1/1	0.64	0.18	0.09	66,66,66,66	0
59	MG	2a	3157	1/1	0.64	0.18	0.07	77,77,77,77	0
60	ZN	1Y	501	1/1	0.96	0.17	0.06	58,58,58,58	0
59	MG	1A	3095	1/1	0.93	0.24	0.05	23,23,23,23	0
59	MG	2X	3002	1/1	0.87	0.18	-0.01	47,47,47,47	0
59	MG	1A	3412	1/1	0.95	0.21	-0.07	37,37,37,37	0
59	MG	2A	3672	1/1	0.89	0.20	-0.08	58,58,58,58	0
59	MG	2A	3547	1/1	0.93	0.18	-0.11	67,67,67,67	0
59	MG	2q	205	1/1	0.71	0.21	-0.13	69,69,69,69	0
59	MG	1D	303	1/1	0.96	0.23	-0.20	35,35,35,35	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3708	1/1	0.93	0.24	-0.21	29,29,29,29	0
59	MG	2A	3556	1/1	0.81	0.20	-0.23	60,60,60,60	0
59	MG	2A	3171	1/1	0.68	0.22	-0.26	44,44,44,44	0
60	ZN	26	501	1/1	0.99	0.18	-0.27	57,57,57,57	0
59	MG	1A	3429	1/1	0.72	0.24	-0.29	50,50,50,50	0
59	MG	1A	3698	1/1	0.93	0.21	-0.30	16,16,16,16	0
59	MG	2A	3447	1/1	0.76	0.17	-0.31	57,57,57,57	0
59	MG	2A	3094	1/1	0.95	0.16	-0.32	43,43,43,43	0
59	MG	1A	4185	1/1	0.72	0.25	-0.32	46,46,46,46	0
59	MG	2a	3007	1/1	0.84	0.14	-0.34	66,66,66,66	0
59	MG	2A	3015	1/1	0.89	0.19	-0.37	57,57,57,57	0
59	MG	1a	3302	1/1	0.94	0.19	-0.38	53,53,53,53	0
59	MG	1a	3494	1/1	0.94	0.24	-0.38	48,48,48,48	0
59	MG	2a	3119	1/1	0.96	0.19	-0.40	68,68,68,68	0
59	MG	1A	3875	1/1	0.88	0.21	-0.43	54,54,54,54	0
59	MG	1A	3270	1/1	0.97	0.21	-0.45	45,45,45,45	0
59	MG	2a	3220	1/1	0.93	0.18	-0.45	65,65,65,65	0
59	MG	1A	3936	1/1	0.87	0.20	-0.46	19,19,19,19	0
59	MG	1A	3785	1/1	0.97	0.19	-0.47	18,18,18,18	0
59	MG	2j	8002	1/1	0.27	0.31	-0.48	85,85,85,85	0
59	MG	2B	3011	1/1	0.88	0.15	-0.48	62,62,62,62	0
59	MG	1A	3556	1/1	0.89	0.18	-0.49	30,30,30,30	0
59	MG	2A	3817	1/1	0.95	0.19	-0.53	32,32,32,32	0
59	MG	1a	3449	1/1	0.96	0.18	-0.55	53,53,53,53	0
59	MG	1a	3467	1/1	0.91	0.19	-0.61	53,53,53,53	0
59	MG	2a	3223	1/1	0.90	0.22	-0.62	69,69,69,69	0
59	MG	1E	314	1/1	0.91	0.18	-0.63	47,47,47,47	0
59	MG	1A	3089	1/1	0.92	0.21	-0.64	28,28,28,28	0
59	MG	2a	3237	1/1	0.86	0.16	-0.67	56,56,56,56	0
59	MG	2w	3006	1/1	0.88	0.17	-0.68	72,72,72,72	0
59	MG	2A	3557	1/1	0.77	0.18	-0.70	53,53,53,53	0
60	ZN	2Y	501	1/1	0.89	0.13	-0.70	86,86,86,86	0
59	MG	2A	3529	1/1	0.94	0.21	-0.72	47,47,47,47	0
59	MG	2a	3144	1/1	0.93	0.20	-0.74	50,50,50,50	0
59	MG	2a	3216	1/1	0.96	0.18	-0.75	63,63,63,63	0
59	MG	1D	302	1/1	0.89	0.21	-0.75	17,17,17,17	0
59	MG	1A	3852	1/1	0.94	0.22	-0.75	36,36,36,36	0
59	MG	2G	3001	1/1	0.76	0.20	-0.76	64,64,64,64	0
59	MG	28	101	1/1	0.95	0.21	-0.78	54,54,54,54	0
59	MG	1A	3926	1/1	0.84	0.20	-0.81	22,22,22,22	0
59	MG	2e	3001	1/1	0.96	0.13	-0.81	62,62,62,62	0
59	MG	1a	3412	1/1	0.92	0.19	-0.83	51,51,51,51	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3256	1/1	0.92	0.21	-0.85	27,27,27,27	0
61	SF4	1d	501	8/8	0.98	0.18	-0.91	52,55,64,66	0
59	MG	1A	3689	1/1	0.93	0.18	-0.92	41,41,41,41	0
59	MG	1A	3285	1/1	0.92	0.18	-0.96	42,42,42,42	0
59	MG	2a	3066	1/1	0.89	0.16	-0.96	54,54,54,54	0
59	MG	1A	3053	1/1	0.91	0.19	-0.96	44,44,44,44	0
59	MG	2a	3073	1/1	0.90	0.17	-0.97	71,71,71,71	0
59	MG	2q	201	1/1	0.96	0.14	-0.99	50,50,50,50	0
59	MG	2a	3113	1/1	0.52	0.15	-1.06	69,69,69,69	0
59	MG	2A	3690	1/1	0.80	0.20	-1.06	37,37,37,37	0
59	MG	2A	3104	1/1	0.85	0.21	-1.08	52,52,52,52	0
59	MG	2B	3008	1/1	0.79	0.17	-1.11	66,66,66,66	0
59	MG	2a	3049	1/1	0.91	0.15	-1.13	55,55,55,55	0
60	ZN	1n	501	1/1	0.99	0.17	-1.14	54,54,54,54	0
59	MG	1a	3307	1/1	0.94	0.18	-1.15	51,51,51,51	0
59	MG	1A	3183	1/1	0.89	0.19	-1.16	39,39,39,39	0
59	MG	2a	3052	1/1	0.89	0.19	-1.16	52,52,52,52	0
59	MG	2T	201	1/1	0.85	0.19	-1.16	54,54,54,54	0
59	MG	1A	4176	1/1	0.98	0.21	-1.17	15,15,15,15	0
60	ZN	29	501	1/1	0.93	0.13	-1.17	69,69,69,69	0
59	MG	1p	101	1/1	0.96	0.19	-1.17	48,48,48,48	0
61	SF4	2d	501	8/8	0.99	0.16	-1.18	56,66,80,82	0
59	MG	2l	201	1/1	0.98	0.21	-1.23	65,65,65,65	0
59	MG	1A	3727	1/1	0.94	0.20	-1.23	15,15,15,15	0
59	MG	2A	3036	1/1	0.95	0.16	-1.23	49,49,49,49	0
59	MG	1A	4146	1/1	0.98	0.22	-1.25	26,26,26,26	0
60	ZN	2n	501	1/1	0.93	0.10	-1.31	90,90,90,90	0
59	MG	1G	3001	1/1	0.92	0.17	-1.32	32,32,32,32	0
59	MG	2A	3297	1/1	0.89	0.17	-1.32	52,52,52,52	0
59	MG	1A	3028	1/1	0.95	0.19	-1.33	34,34,34,34	0
59	MG	1w	101	1/1	0.92	0.18	-1.34	56,56,56,56	0
59	MG	2B	3010	1/1	0.86	0.13	-1.35	60,60,60,60	0
59	MG	2Z	8001	1/1	0.95	0.19	-1.35	68,68,68,68	0
59	MG	1A	4156	1/1	0.95	0.20	-1.36	37,37,37,37	0
59	MG	1A	3115	1/1	0.98	0.18	-1.37	24,24,24,24	0
59	MG	2a	3141	1/1	0.93	0.10	-1.39	56,56,56,56	0
59	MG	1A	3067	1/1	0.98	0.17	-1.41	19,19,19,19	0
59	MG	1A	3260	1/1	0.98	0.17	-1.41	32,32,32,32	0
59	MG	2a	3062	1/1	0.86	0.19	-1.44	54,54,54,54	0
59	MG	2A	3278	1/1	0.94	0.20	-1.47	40,40,40,40	0
59	MG	1A	4094	1/1	0.90	0.18	-1.47	35,35,35,35	0
59	MG	1A	4075	1/1	0.82	0.21	-1.47	21,21,21,21	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2a	3142	1/1	0.91	0.10	-1.48	67,67,67,67	0
59	MG	2A	3582	1/1	0.79	0.18	-1.50	29,29,29,29	0
59	MG	2A	3311	1/1	0.86	0.19	-1.51	44,44,44,44	0
59	MG	1A	4226	1/1	0.98	0.18	-1.52	24,24,24,24	0
59	MG	2a	3146	1/1	0.94	0.12	-1.52	79,79,79,79	0
59	MG	B	3001	1/1	0.85	0.22	-1.56	46,46,46,46	0
59	MG	1A	3007	1/1	0.98	0.18	-1.59	17,17,17,17	0
59	MG	1A	3009	1/1	0.92	0.19	-1.59	25,25,25,25	0
59	MG	1A	3030	1/1	0.97	0.20	-1.60	14,14,14,14	0
59	MG	2B	3016	1/1	0.86	0.14	-1.61	77,77,77,77	0
59	MG	1A	3554	1/1	0.96	0.19	-1.61	26,26,26,26	0
59	MG	2k	3001	1/1	0.96	0.15	-1.63	50,50,50,50	0
59	MG	2a	3094	1/1	0.98	0.15	-1.63	48,48,48,48	0
59	MG	1A	4186	1/1	0.96	0.18	-1.64	27,27,27,27	0
59	MG	1A	3690	1/1	0.83	0.21	-1.64	14,14,14,14	0
59	MG	1A	3041	1/1	0.95	0.19	-1.64	20,20,20,20	0
59	MG	1a	3403	1/1	0.96	0.14	-1.74	28,28,28,28	0
59	MG	1G	3003	1/1	0.70	0.14	-1.74	56,56,56,56	0
59	MG	1A	3156	1/1	0.93	0.12	-1.74	58,58,58,58	0
59	MG	16	101	1/1	0.97	0.16	-1.74	32,32,32,32	0
59	MG	2a	3196	1/1	0.86	0.15	-1.75	63,63,63,63	0
59	MG	2A	3421	1/1	0.97	0.19	-1.77	42,42,42,42	0
60	ZN	24	501	1/1	0.75	0.06	-1.79	111,111,111,111	0
59	MG	2A	3767	1/1	0.87	0.15	-1.79	59,59,59,59	0
59	MG	2A	3008	1/1	0.83	0.14	-1.82	44,44,44,44	0
59	MG	2A	3742	1/1	0.92	0.19	-1.84	69,69,69,69	0
59	MG	2A	3524	1/1	0.72	0.17	-1.84	51,51,51,51	0
59	MG	1A	3616	1/1	0.95	0.20	-1.87	16,16,16,16	0
59	MG	1A	3697	1/1	0.95	0.14	-1.90	18,18,18,18	0
59	MG	1A	3913	1/1	0.76	0.19	-1.91	21,21,21,21	0
59	MG	1A	3172	1/1	0.98	0.20	-1.92	25,25,25,25	0
59	MG	2f	3001	1/1	0.96	0.15	-1.97	38,38,38,38	0
59	MG	1O	3006	1/1	0.92	0.16	-1.98	64,64,64,64	0
59	MG	2a	3121	1/1	0.91	0.15	-1.99	61,61,61,61	0
59	MG	1B	228	1/1	0.95	0.16	-1.99	30,30,30,30	0
59	MG	2A	3589	1/1	0.94	0.12	-2.00	46,46,46,46	0
59	MG	2A	3210	1/1	0.95	0.17	-2.00	62,62,62,62	0
59	MG	1a	3343	1/1	0.93	0.16	-2.02	40,40,40,40	0
59	MG	2a	3135	1/1	0.79	0.16	-2.03	60,60,60,60	0
59	MG	1A	3499	1/1	0.80	0.16	-2.04	46,46,46,46	0
59	MG	2a	3228	1/1	0.96	0.10	-2.04	56,56,56,56	0
59	MG	2a	3070	1/1	0.96	0.12	-2.07	60,60,60,60	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
60	ZN	14	501	1/1	0.91	0.11	-2.08	87,87,87,87	0
59	MG	2a	3056	1/1	0.96	0.19	-2.09	47,47,47,47	0
59	MG	1a	3348	1/1	0.91	0.17	-2.11	41,41,41,41	0
59	MG	1a	3352	1/1	0.94	0.18	-2.11	42,42,42,42	0
59	MG	2A	3500	1/1	0.96	0.18	-2.13	27,27,27,27	0
59	MG	1t	3001	1/1	0.87	0.17	-2.13	43,43,43,43	0
59	MG	1B	224	1/1	0.93	0.14	-2.14	52,52,52,52	0
59	MG	2A	3510	1/1	0.89	0.18	-2.16	58,58,58,58	0
59	MG	2a	3065	1/1	0.95	0.15	-2.17	58,58,58,58	0
59	MG	2A	3257	1/1	0.93	0.18	-2.17	49,49,49,49	0
59	MG	2A	3432	1/1	0.97	0.16	-2.18	26,26,26,26	0
59	MG	1a	3351	1/1	0.93	0.18	-2.22	42,42,42,42	0
59	MG	1A	4216	1/1	0.98	0.18	-2.23	24,24,24,24	0
59	MG	2A	3414	1/1	0.90	0.19	-2.25	43,43,43,43	0
59	MG	2A	3820	1/1	0.94	0.17	-2.26	35,35,35,35	0
59	MG	2A	3834	1/1	0.94	0.13	-2.26	42,42,42,42	0
59	MG	1A	3276	1/1	0.85	0.12	-2.27	52,52,52,52	0
59	MG	2a	3150	1/1	0.92	0.15	-2.28	57,57,57,57	0
59	MG	1A	3597	1/1	0.88	0.17	-2.29	34,34,34,34	0
59	MG	2A	3149	1/1	0.95	0.17	-2.30	29,29,29,29	0
59	MG	1A	4175	1/1	0.91	0.15	-2.31	36,36,36,36	0
59	MG	2A	3027	1/1	0.96	0.19	-2.33	39,39,39,39	0
59	MG	2A	3417	1/1	0.93	0.14	-2.35	31,31,31,31	0
59	MG	1A	3453	1/1	0.90	0.20	-2.35	18,18,18,18	0
59	MG	2A	3099	1/1	0.83	0.16	-2.35	51,51,51,51	0
59	MG	2a	3182	1/1	0.85	0.15	-2.36	67,67,67,67	0
59	MG	2a	3103	1/1	0.84	0.18	-2.37	62,62,62,62	0
59	MG	2A	3749	1/1	0.87	0.12	-2.38	55,55,55,55	0
59	MG	2t	3001	1/1	0.92	0.15	-2.39	53,53,53,53	0
59	MG	2a	3075	1/1	0.98	0.18	-2.39	33,33,33,33	0
59	MG	2A	3537	1/1	0.95	0.14	-2.39	59,59,59,59	0
59	MG	2U	201	1/1	0.97	0.18	-2.40	47,47,47,47	0
59	MG	1A	3216	1/1	0.94	0.15	-2.43	36,36,36,36	0
59	MG	2A	3004	1/1	0.99	0.15	-2.44	46,46,46,46	0
59	MG	2A	3120	1/1	0.97	0.17	-2.51	34,34,34,34	0
59	MG	1A	4166	1/1	0.96	0.17	-2.51	11,11,11,11	0
59	MG	1A	3530	1/1	0.95	0.17	-2.51	32,32,32,32	0
59	MG	1x	107	1/1	0.94	0.11	-2.52	50,50,50,50	0
59	MG	1A	3614	1/1	0.94	0.14	-2.53	24,24,24,24	0
59	MG	1a	3468	1/1	0.95	0.16	-2.53	51,51,51,51	0
59	MG	1A	3981	1/1	0.92	0.19	-2.55	13,13,13,13	0
59	MG	2A	3053	1/1	0.89	0.14	-2.55	61,61,61,61	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2a	3061	1/1	0.93	0.13	-2.56	60,60,60,60	0
59	MG	1A	3969	1/1	0.95	0.19	-2.59	36,36,36,36	0
59	MG	2a	3203	1/1	0.80	0.13	-2.62	57,57,57,57	0
59	MG	2A	3835	1/1	0.94	0.14	-2.62	48,48,48,48	0
59	MG	2A	3480	1/1	0.75	0.15	-2.63	24,24,24,24	0
59	MG	1k	201	1/1	0.98	0.12	-2.63	39,39,39,39	0
59	MG	1A	3934	1/1	0.89	0.18	-2.63	16,16,16,16	0
59	MG	2A	3802	1/1	0.86	0.12	-2.64	35,35,35,35	0
59	MG	1a	3508	1/1	0.85	0.17	-2.65	51,51,51,51	0
59	MG	1A	3622	1/1	0.83	0.19	-2.68	17,17,17,17	0
59	MG	1A	3042	1/1	0.97	0.19	-2.74	17,17,17,17	0
59	MG	2A	3118	1/1	0.99	0.17	-2.77	37,37,37,37	0
59	MG	2A	3410	1/1	0.82	0.16	-2.80	25,25,25,25	0
59	MG	2A	3501	1/1	0.95	0.18	-2.80	33,33,33,33	0
59	MG	2A	3292	1/1	0.73	0.16	-2.81	66,66,66,66	0
59	MG	2A	3445	1/1	0.95	0.17	-2.82	42,42,42,42	0
59	MG	2A	3841	1/1	0.93	0.09	-2.83	59,59,59,59	0
59	MG	1A	3150	1/1	0.98	0.17	-2.83	30,30,30,30	0
59	MG	2A	3566	1/1	0.95	0.07	-2.88	71,71,71,71	0
59	MG	1A	4039	1/1	0.98	0.17	-2.89	50,50,50,50	0
59	MG	1G	3004	1/1	0.94	0.12	-2.89	40,40,40,40	0
59	MG	2A	3608	1/1	0.95	0.13	-2.91	60,60,60,60	0
59	MG	2A	3407	1/1	0.94	0.14	-2.94	34,34,34,34	0
59	MG	2a	3239	1/1	0.91	0.11	-2.95	42,42,42,42	0
59	MG	1A	3503	1/1	0.73	0.14	-2.96	53,53,53,53	0
59	MG	1A	3633	1/1	0.91	0.20	-2.97	22,22,22,22	0
59	MG	1A	3570	1/1	0.89	0.18	-3.00	24,24,24,24	0
59	MG	2a	3189	1/1	0.89	0.14	-3.01	57,57,57,57	0
59	MG	1A	3676	1/1	0.93	0.13	-3.03	29,29,29,29	0
59	MG	1w	107	1/1	0.93	0.11	-3.07	62,62,62,62	0
59	MG	1A	3974	1/1	0.89	0.15	-3.08	17,17,17,17	0
59	MG	2A	3840	1/1	0.83	0.15	-3.08	30,30,30,30	0
59	MG	2A	3371	1/1	0.94	0.16	-3.09	48,48,48,48	0
59	MG	1A	3736	1/1	0.98	0.19	-3.12	15,15,15,15	0
59	MG	1A	3211	1/1	0.90	0.16	-3.12	47,47,47,47	0
59	MG	1A	3695	1/1	0.88	0.17	-3.18	20,20,20,20	0
59	MG	1Q	201	1/1	0.92	0.13	-3.19	25,25,25,25	0
59	MG	1A	4229	1/1	0.90	0.16	-3.21	24,24,24,24	0
59	MG	2A	3561	1/1	0.97	0.15	-3.21	31,31,31,31	0
59	MG	1A	3903	1/1	0.98	0.16	-3.22	29,29,29,29	0
59	MG	1a	3577	1/1	0.86	0.10	-3.23	50,50,50,50	0
59	MG	2A	3105	1/1	0.84	0.10	-3.23	54,54,54,54	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3521	1/1	0.83	0.18	-3.24	42,42,42,42	0
59	MG	1A	3258	1/1	0.92	0.17	-3.25	26,26,26,26	0
59	MG	2A	3081	1/1	0.89	0.12	-3.25	45,45,45,45	0
59	MG	2A	3709	1/1	0.91	0.14	-3.27	46,46,46,46	0
59	MG	1A	3646	1/1	0.98	0.14	-3.29	8,8,8,8	0
59	MG	1a	3453	1/1	0.90	0.09	-3.35	64,64,64,64	0
59	MG	1a	3513	1/1	0.97	0.06	-3.35	61,61,61,61	0
59	MG	2A	3430	1/1	0.90	0.13	-3.36	29,29,29,29	0
59	MG	1A	4149	1/1	0.91	0.19	-3.36	47,47,47,47	0
59	MG	2A	3481	1/1	0.85	0.13	-3.37	37,37,37,37	0
59	MG	2A	3356	1/1	0.98	0.17	-3.38	18,18,18,18	0
59	MG	1a	3461	1/1	0.92	0.14	-3.39	41,41,41,41	0
59	MG	2A	3519	1/1	0.96	0.10	-3.43	62,62,62,62	0
59	MG	2A	3433	1/1	0.90	0.18	-3.43	38,38,38,38	0
59	MG	1A	4201	1/1	0.95	0.12	-3.43	32,32,32,32	0
59	MG	2A	3761	1/1	0.92	0.07	-3.44	61,61,61,61	0
59	MG	2a	3170	1/1	0.84	0.16	-3.47	49,49,49,49	0
59	MG	1A	3640	1/1	0.97	0.15	-3.51	25,25,25,25	0
59	MG	2A	3825	1/1	0.95	0.12	-3.57	42,42,42,42	0
59	MG	1a	3567	1/1	0.71	0.08	-3.58	83,83,83,83	0
59	MG	1A	3179	1/1	0.96	0.13	-3.65	25,25,25,25	0
59	MG	1a	3564	1/1	0.88	0.12	-3.68	54,54,54,54	0
59	MG	2A	3816	1/1	0.92	0.16	-3.69	43,43,43,43	0
59	MG	10	102	1/1	0.96	0.09	-3.71	43,43,43,43	0
59	MG	2a	3082	1/1	0.91	0.09	-3.73	65,65,65,65	0
59	MG	2A	3503	1/1	0.92	0.10	-3.73	32,32,32,32	0
59	MG	2a	3200	1/1	0.94	0.14	-3.74	65,65,65,65	0
59	MG	1A	3606	1/1	0.95	0.17	-3.76	17,17,17,17	0
59	MG	1A	4016	1/1	0.88	0.14	-3.76	24,24,24,24	0
59	MG	1a	3442	1/1	0.94	0.10	-3.82	59,59,59,59	0
59	MG	1a	3316	1/1	0.94	0.12	-3.83	40,40,40,40	0
59	MG	2A	3558	1/1	0.81	0.14	-3.84	56,56,56,56	0
59	MG	2a	3207	1/1	0.94	0.16	-3.85	55,55,55,55	0
59	MG	1e	3001	1/1	0.90	0.12	-3.85	67,67,67,67	0
59	MG	1a	3551	1/1	0.97	0.11	-3.91	54,54,54,54	0
59	MG	1A	3939	1/1	0.88	0.13	-3.91	36,36,36,36	0
59	MG	1a	3325	1/1	0.90	0.14	-3.92	50,50,50,50	0
59	MG	1E	301	1/1	0.89	0.17	-3.94	33,33,33,33	0
59	MG	1A	3691	1/1	0.88	0.14	-3.94	39,39,39,39	0
59	MG	2A	3160	1/1	0.91	0.16	-3.94	48,48,48,48	0
59	MG	1A	3820	1/1	0.95	0.16	-3.96	13,13,13,13	0
59	MG	2A	3752	1/1	0.67	0.08	-3.97	61,61,61,61	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3821	1/1	0.97	0.09	-3.98	35,35,35,35	0
59	MG	1A	3724	1/1	0.92	0.14	-4.00	46,46,46,46	0
59	MG	2A	3209	1/1	0.93	0.11	-4.04	60,60,60,60	0
59	MG	1a	3359	1/1	0.92	0.10	-4.07	66,66,66,66	0
59	MG	2a	3235	1/1	0.89	0.07	-4.13	80,80,80,80	0
59	MG	2a	3159	1/1	0.87	0.14	-4.14	62,62,62,62	0
59	MG	1A	3864	1/1	0.96	0.13	-4.16	43,43,43,43	0
59	MG	2a	3131	1/1	0.87	0.09	-4.17	77,77,77,77	0
59	MG	1a	3560	1/1	0.85	0.11	-4.18	53,53,53,53	0
59	MG	2A	3025	1/1	0.91	0.18	-4.18	42,42,42,42	0
59	MG	1A	3620	1/1	0.95	0.18	-4.18	15,15,15,15	0
59	MG	1A	3674	1/1	0.91	0.18	-4.19	14,14,14,14	0
59	MG	1A	3589	1/1	0.87	0.16	-4.23	40,40,40,40	0
59	MG	2A	3567	1/1	0.94	0.12	-4.24	63,63,63,63	0
59	MG	1A	3842	1/1	0.93	0.18	-4.26	15,15,15,15	0
59	MG	2A	3010	1/1	0.99	0.13	-4.28	36,36,36,36	0
59	MG	2a	3172	1/1	0.91	0.15	-4.30	42,42,42,42	0
59	MG	1A	3016	1/1	0.98	0.18	-4.31	20,20,20,20	0
59	MG	1A	4152	1/1	0.94	0.16	-4.32	21,21,21,21	0
59	MG	1A	3857	1/1	0.74	0.15	-4.33	22,22,22,22	0
59	MG	2A	3685	1/1	0.84	0.12	-4.35	62,62,62,62	0
59	MG	1a	3312	1/1	0.97	0.09	-4.36	62,62,62,62	0
59	MG	2A	3404	1/1	0.94	0.09	-4.39	48,48,48,48	0
59	MG	2A	3391	1/1	0.91	0.17	-4.39	41,41,41,41	0
59	MG	1n	503	1/1	0.93	0.14	-4.39	49,49,49,49	0
59	MG	1A	3710	1/1	0.97	0.17	-4.42	18,18,18,18	0
59	MG	2A	3745	1/1	0.88	0.15	-4.42	48,48,48,48	0
59	MG	1A	3730	1/1	0.83	0.14	-4.45	33,33,33,33	0
59	MG	2a	3137	1/1	0.97	0.09	-4.46	68,68,68,68	0
59	MG	1a	3310	1/1	0.96	0.17	-4.46	19,19,19,19	0
59	MG	1A	3843	1/1	0.87	0.15	-4.46	19,19,19,19	0
59	MG	1A	3166	1/1	0.89	0.14	-4.48	54,54,54,54	0
59	MG	2A	3384	1/1	0.97	0.15	-4.48	28,28,28,28	0
59	MG	1a	3490	1/1	0.88	0.11	-4.50	57,57,57,57	0
59	MG	2A	3387	1/1	0.93	0.11	-4.50	47,47,47,47	0
59	MG	2A	3607	1/1	0.93	0.14	-4.51	30,30,30,30	0
59	MG	2a	3166	1/1	0.91	0.10	-4.52	65,65,65,65	0
59	MG	2A	3499	1/1	0.94	0.13	-4.54	53,53,53,53	0
59	MG	1A	3548	1/1	0.93	0.14	-4.54	38,38,38,38	0
59	MG	2A	3490	1/1	0.89	0.14	-4.57	30,30,30,30	0
59	MG	1A	3489	1/1	0.79	0.14	-4.57	33,33,33,33	0
59	MG	16	103	1/1	0.90	0.12	-4.58	49,49,49,49	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2d	502	1/1	0.95	0.13	-4.58	59,59,59,59	0
59	MG	1A	3110	1/1	0.91	0.15	-4.59	29,29,29,29	0
59	MG	1a	3323	1/1	0.94	0.13	-4.65	43,43,43,43	0
59	MG	1A	4069	1/1	0.95	0.12	-4.66	48,48,48,48	0
59	MG	1A	3502	1/1	0.94	0.13	-4.68	35,35,35,35	0
59	MG	2A	3156	1/1	0.94	0.12	-4.69	39,39,39,39	0
59	MG	2a	3067	1/1	0.93	0.13	-4.71	56,56,56,56	0
59	MG	1a	3504	1/1	0.84	0.13	-4.74	57,57,57,57	0
59	MG	2A	3413	1/1	0.80	0.11	-4.74	43,43,43,43	0
59	MG	1a	3314	1/1	0.90	0.11	-4.78	53,53,53,53	0
59	MG	2A	3744	1/1	0.72	0.10	-4.80	47,47,47,47	0
59	MG	1A	3663	1/1	0.99	0.16	-4.82	14,14,14,14	0
59	MG	1A	3047	1/1	0.97	0.15	-4.82	19,19,19,19	0
59	MG	2A	3613	1/1	0.83	0.16	-4.82	36,36,36,36	0
59	MG	1A	3046	1/1	0.95	0.14	-4.84	28,28,28,28	0
59	MG	1A	3545	1/1	0.94	0.18	-4.84	21,21,21,21	0
59	MG	1a	3474	1/1	0.93	0.13	-4.85	42,42,42,42	0
59	MG	1A	3006	1/1	0.94	0.19	-4.86	23,23,23,23	0
59	MG	2A	3400	1/1	0.97	0.16	-4.86	21,21,21,21	0
59	MG	1A	3010	1/1	0.99	0.13	-4.88	16,16,16,16	0
59	MG	1A	3558	1/1	0.97	0.16	-4.90	15,15,15,15	0
59	MG	1a	3407	1/1	0.86	0.14	-4.91	36,36,36,36	0
59	MG	1a	3317	1/1	0.93	0.09	-4.92	41,41,41,41	0
59	MG	2A	3689	1/1	0.84	0.10	-4.92	65,65,65,65	0
59	MG	2A	3026	1/1	0.89	0.15	-4.93	32,32,32,32	0
59	MG	1A	3677	1/1	0.90	0.14	-4.94	10,10,10,10	0
59	MG	1A	3559	1/1	0.97	0.14	-4.96	28,28,28,28	0
59	MG	1A	3173	1/1	0.93	0.12	-4.97	36,36,36,36	0
59	MG	2A	3509	1/1	0.96	0.12	-4.97	40,40,40,40	0
59	MG	2A	3462	1/1	0.87	0.11	-5.01	62,62,62,62	0
59	MG	2A	3479	1/1	0.89	0.14	-5.03	26,26,26,26	0
59	MG	2A	3023	1/1	0.94	0.12	-5.05	35,35,35,35	0
59	MG	2r	3001	1/1	0.86	0.11	-5.05	77,77,77,77	0
59	MG	2A	3461	1/1	0.98	0.13	-5.05	27,27,27,27	0
59	MG	1A	3169	1/1	0.93	0.14	-5.06	34,34,34,34	0
59	MG	2D	303	1/1	0.94	0.10	-5.08	20,20,20,20	0
59	MG	1A	3827	1/1	0.98	0.15	-5.08	23,23,23,23	0
59	MG	1F	303	1/1	0.97	0.15	-5.13	24,24,24,24	0
59	MG	2A	3086	1/1	0.93	0.12	-5.13	30,30,30,30	0
59	MG	1E	306	1/1	0.96	0.13	-5.13	31,31,31,31	0
59	MG	1A	3954	1/1	0.95	0.07	-5.14	49,49,49,49	0
59	MG	1A	3182	1/1	0.91	0.14	-5.14	38,38,38,38	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1a	3518	1/1	0.95	0.13	-5.14	32,32,32,32	0
59	MG	1A	3577	1/1	0.76	0.12	-5.14	57,57,57,57	0
59	MG	1a	3416	1/1	0.70	0.17	-5.15	62,62,62,62	0
59	MG	1A	4062	1/1	0.95	0.13	-5.16	34,34,34,34	0
59	MG	1A	3257	1/1	0.98	0.17	-5.16	24,24,24,24	0
59	MG	2A	3101	1/1	0.96	0.13	-5.18	27,27,27,27	0
59	MG	2A	3130	1/1	0.97	0.13	-5.18	46,46,46,46	0
59	MG	1a	3349	1/1	0.93	0.11	-5.19	45,45,45,45	0
59	MG	1A	3993	1/1	0.86	0.13	-5.20	20,20,20,20	0
59	MG	2A	3497	1/1	0.94	0.15	-5.20	30,30,30,30	0
59	MG	1A	3765	1/1	0.88	0.16	-5.21	33,33,33,33	0
59	MG	2A	3756	1/1	0.94	0.12	-5.21	51,51,51,51	0
59	MG	2A	3687	1/1	0.78	0.12	-5.24	39,39,39,39	0
59	MG	2A	3150	1/1	0.74	0.12	-5.26	42,42,42,42	0
59	MG	2A	3590	1/1	0.89	0.13	-5.28	35,35,35,35	0
59	MG	2Q	3001	1/1	0.91	0.09	-5.28	52,52,52,52	0
59	MG	2A	3419	1/1	0.93	0.12	-5.31	28,28,28,28	0
59	MG	2A	3693	1/1	0.96	0.13	-5.31	32,32,32,32	0
59	MG	1A	3568	1/1	0.93	0.16	-5.35	15,15,15,15	0
59	MG	1P	203	1/1	0.88	0.14	-5.36	49,49,49,49	0
59	MG	2a	3068	1/1	0.93	0.15	-5.38	52,52,52,52	0
59	MG	2A	3449	1/1	0.96	0.15	-5.40	17,17,17,17	0
59	MG	1A	3043	1/1	0.92	0.16	-5.41	21,21,21,21	0
59	MG	2A	3411	1/1	0.96	0.12	-5.42	40,40,40,40	0
59	MG	1a	3532	1/1	0.70	0.12	-5.42	57,57,57,57	0
59	MG	1A	3571	1/1	0.99	0.17	-5.42	13,13,13,13	0
59	MG	1A	3726	1/1	0.95	0.15	-5.45	14,14,14,14	0
59	MG	2E	302	1/1	0.95	0.10	-5.47	38,38,38,38	0
59	MG	2A	3020	1/1	0.97	0.10	-5.50	24,24,24,24	0
59	MG	1a	3530	1/1	0.80	0.11	-5.52	62,62,62,62	0
59	MG	1A	3653	1/1	0.99	0.15	-5.57	12,12,12,12	0
59	MG	1A	3546	1/1	0.91	0.15	-5.65	37,37,37,37	0
59	MG	1A	3937	1/1	0.97	0.15	-5.65	22,22,22,22	0
59	MG	1A	3773	1/1	0.92	0.07	-5.66	57,57,57,57	0
59	MG	2a	3054	1/1	0.96	0.14	-5.67	41,41,41,41	0
59	MG	1A	3592	1/1	0.98	0.16	-5.68	17,17,17,17	0
59	MG	1A	3598	1/1	0.98	0.15	-5.69	24,24,24,24	0
59	MG	1A	4064	1/1	0.92	0.14	-5.69	30,30,30,30	0
59	MG	1a	3347	1/1	0.98	0.16	-5.69	40,40,40,40	0
59	MG	1A	3008	1/1	0.95	0.13	-5.69	16,16,16,16	0
59	MG	1a	3458	1/1	0.95	0.09	-5.70	51,51,51,51	0
59	MG	1A	3883	1/1	0.48	0.10	-5.74	52,52,52,52	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1a	3308	1/1	0.93	0.15	-5.75	44,44,44,44	0
59	MG	2A	3386	1/1	0.96	0.11	-5.79	45,45,45,45	0
59	MG	1A	4165	1/1	0.99	0.18	-5.83	9,9,9,9	0
59	MG	1A	3157	1/1	0.86	0.12	-5.84	37,37,37,37	0
59	MG	1A	3701	1/1	0.96	0.14	-5.88	17,17,17,17	0
59	MG	1A	3868	1/1	0.92	0.08	-5.88	30,30,30,30	0
59	MG	2A	3078	1/1	0.88	0.17	-5.88	33,33,33,33	0
59	MG	1a	3332	1/1	0.91	0.11	-5.89	38,38,38,38	0
59	MG	2A	3814	1/1	0.87	0.10	-5.96	44,44,44,44	0
59	MG	1A	3721	1/1	0.98	0.18	-5.96	14,14,14,14	0
59	MG	1A	4224	1/1	0.99	0.10	-5.97	23,23,23,23	0
59	MG	1A	3602	1/1	0.93	0.15	-5.98	19,19,19,19	0
59	MG	2A	3572	1/1	0.94	0.15	-6.00	24,24,24,24	0
59	MG	1A	3960	1/1	0.95	0.17	-6.01	16,16,16,16	0
59	MG	2a	3195	1/1	0.89	0.11	-6.04	54,54,54,54	0
59	MG	2a	3181	1/1	0.92	0.11	-6.05	55,55,55,55	0
59	MG	1A	3072	1/1	0.92	0.11	-6.08	34,34,34,34	0
59	MG	1A	3591	1/1	0.92	0.12	-6.12	26,26,26,26	0
59	MG	2A	3700	1/1	0.91	0.16	-6.13	35,35,35,35	0
59	MG	1A	3686	1/1	0.84	0.09	-6.15	29,29,29,29	0
59	MG	1A	3935	1/1	0.93	0.12	-6.25	24,24,24,24	0
59	MG	1B	219	1/1	0.92	0.16	-6.25	28,28,28,28	0
59	MG	2A	3424	1/1	0.93	0.12	-6.29	31,31,31,31	0
59	MG	2A	3484	1/1	0.92	0.15	-6.31	33,33,33,33	0
59	MG	1A	3618	1/1	0.96	0.15	-6.31	16,16,16,16	0
59	MG	1A	3609	1/1	0.97	0.13	-6.34	17,17,17,17	0
59	MG	1A	3601	1/1	0.98	0.14	-6.35	12,12,12,12	0
59	MG	1A	3915	1/1	0.77	0.15	-6.42	24,24,24,24	0
59	MG	1A	4058	1/1	0.91	0.13	-6.44	47,47,47,47	0
59	MG	2A	3416	1/1	0.88	0.14	-6.45	34,34,34,34	0
59	MG	1A	3693	1/1	0.91	0.18	-6.46	13,13,13,13	0
59	MG	1a	3487	1/1	0.94	0.11	-6.46	62,62,62,62	0
59	MG	1A	3449	1/1	0.77	0.12	-6.53	54,54,54,54	0
59	MG	1a	3531	1/1	0.93	0.10	-6.55	49,49,49,49	0
59	MG	2A	3372	1/1	0.94	0.10	-6.55	50,50,50,50	0
59	MG	2A	3308	1/1	0.87	0.12	-6.55	50,50,50,50	0
59	MG	2A	3468	1/1	0.82	0.13	-6.58	32,32,32,32	0
59	MG	2A	3578	1/1	0.93	0.14	-6.58	25,25,25,25	0
59	MG	1A	3575	1/1	0.94	0.14	-6.67	13,13,13,13	0
59	MG	1A	3720	1/1	0.95	0.11	-6.68	34,34,34,34	0
59	MG	2A	3512	1/1	0.92	0.06	-6.68	45,45,45,45	0
59	MG	1a	3470	1/1	0.89	0.09	-6.74	49,49,49,49	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3795	1/1	0.95	0.14	-6.74	23,23,23,23	0
59	MG	2A	3436	1/1	0.75	0.13	-6.81	28,28,28,28	0
59	MG	1A	3709	1/1	0.97	0.14	-6.83	20,20,20,20	0
59	MG	2A	3622	1/1	0.84	0.12	-6.86	58,58,58,58	0
59	MG	1A	3607	1/1	0.93	0.13	-6.86	17,17,17,17	0
59	MG	2A	3096	1/1	0.91	0.10	-6.89	55,55,55,55	0
59	MG	2A	3633	1/1	0.96	0.11	-6.89	37,37,37,37	0
59	MG	1A	3861	1/1	0.86	0.12	-6.96	39,39,39,39	0
59	MG	2A	3482	1/1	0.94	0.13	-7.01	40,40,40,40	0
59	MG	2A	3485	1/1	0.69	0.14	-7.16	36,36,36,36	0
59	MG	1A	4168	1/1	0.79	0.10	-7.21	36,36,36,36	0
59	MG	2A	3394	1/1	0.97	0.15	-7.23	40,40,40,40	0
59	MG	1A	3984	1/1	0.95	0.15	-7.27	10,10,10,10	0
59	MG	1A	3605	1/1	0.95	0.16	-7.31	14,14,14,14	0
59	MG	2A	3385	1/1	0.86	0.12	-7.32	63,63,63,63	0
59	MG	2A	3492	1/1	0.90	0.15	-7.33	44,44,44,44	0
59	MG	2A	3694	1/1	0.93	0.13	-7.37	28,28,28,28	0
59	MG	2A	3125	1/1	0.94	0.13	-7.37	30,30,30,30	0
59	MG	1A	3611	1/1	0.97	0.16	-7.44	17,17,17,17	0
59	MG	2E	305	1/1	0.89	0.14	-7.48	43,43,43,43	0
59	MG	2A	3530	1/1	0.90	0.13	-7.50	35,35,35,35	0
59	MG	1A	3018	1/1	0.96	0.13	-7.53	8,8,8,8	0
59	MG	1A	3821	1/1	0.94	0.16	-7.60	21,21,21,21	0
59	MG	1a	3445	1/1	0.97	0.13	-7.67	32,32,32,32	0
59	MG	1A	3555	1/1	0.96	0.12	-7.69	48,48,48,48	0
59	MG	2A	3489	1/1	0.97	0.13	-7.90	32,32,32,32	0
59	MG	1A	3712	1/1	0.94	0.12	-7.91	43,43,43,43	0
59	MG	1a	3421	1/1	0.81	0.13	-7.97	64,64,64,64	0
59	MG	1A	3638	1/1	0.82	0.15	-7.98	28,28,28,28	0
59	MG	2A	3439	1/1	0.96	0.11	-8.03	32,32,32,32	0
59	MG	1a	3495	1/1	0.93	0.08	-8.17	63,63,63,63	0
59	MG	1A	3206	1/1	0.88	0.14	-8.27	35,35,35,35	0
59	MG	2a	3088	1/1	0.94	0.11	-8.34	61,61,61,61	0
59	MG	2A	3707	1/1	0.95	0.11	-8.35	64,64,64,64	0
59	MG	1a	3457	1/1	0.99	0.11	-8.38	28,28,28,28	0
59	MG	1A	3001	1/1	0.90	0.11	-8.49	35,35,35,35	0
59	MG	2A	3009	1/1	0.96	0.11	-8.50	31,31,31,31	0
59	MG	1A	3557	1/1	0.97	0.10	-8.51	23,23,23,23	0
59	MG	2A	3538	1/1	0.92	0.07	-8.55	47,47,47,47	0
59	MG	1A	3717	1/1	0.90	0.11	-8.66	21,21,21,21	0
59	MG	1A	3017	1/1	0.95	0.10	-8.73	28,28,28,28	0
59	MG	1A	4193	1/1	0.86	0.13	-8.74	14,14,14,14	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	4092	1/1	0.91	0.15	-8.74	45,45,45,45	0
59	MG	1a	3574	1/1	0.96	0.10	-8.76	43,43,43,43	0
59	MG	1A	3057	1/1	0.90	0.15	-8.84	26,26,26,26	0
59	MG	2A	3550	1/1	0.88	0.12	-8.88	56,56,56,56	0
59	MG	1A	3600	1/1	0.96	0.14	-8.95	21,21,21,21	0
59	MG	2A	3598	1/1	0.82	0.06	-8.97	63,63,63,63	0
59	MG	1A	3576	1/1	0.99	0.14	-8.99	9,9,9,9	0
59	MG	2A	3275	1/1	0.94	0.07	-9.01	66,66,66,66	0
59	MG	1A	3635	1/1	0.96	0.13	-9.07	8,8,8,8	0
59	MG	1A	3735	1/1	0.95	0.13	-9.10	20,20,20,20	0
59	MG	2a	3177	1/1	0.81	0.11	-9.14	58,58,58,58	0
59	MG	2A	3746	1/1	0.79	0.13	-9.14	48,48,48,48	0
59	MG	1A	3755	1/1	0.93	0.11	-9.21	24,24,24,24	0
59	MG	2A	3643	1/1	0.92	0.09	-9.34	48,48,48,48	0
59	MG	1A	3199	1/1	0.91	0.15	-9.35	32,32,32,32	0
59	MG	2A	3698	1/1	0.92	0.07	-9.43	49,49,49,49	0
59	MG	1A	3777	1/1	0.98	0.09	-9.47	45,45,45,45	0
59	MG	2A	3716	1/1	0.92	0.12	-9.63	56,56,56,56	0
59	MG	1A	3650	1/1	0.89	0.14	-9.82	36,36,36,36	0
59	MG	2A	3392	1/1	0.95	0.09	-9.82	58,58,58,58	0
59	MG	2A	3661	1/1	0.96	0.08	-9.85	54,54,54,54	0
59	MG	1B	207	1/1	0.96	0.10	-9.89	32,32,32,32	0
59	MG	1A	3923	1/1	0.85	0.07	-9.92	46,46,46,46	0
59	MG	1A	3757	1/1	0.97	0.09	-9.94	39,39,39,39	0
59	MG	1A	3814	1/1	0.97	0.13	-9.95	22,22,22,22	0
59	MG	2A	3396	1/1	0.91	0.09	-10.02	41,41,41,41	0
59	MG	1A	3873	1/1	0.86	0.09	-10.03	30,30,30,30	0
59	MG	1A	4090	1/1	0.98	0.16	-10.11	34,34,34,34	0
59	MG	1A	3786	1/1	0.92	0.11	-10.13	26,26,26,26	0
59	MG	1A	3681	1/1	0.97	0.08	-10.14	23,23,23,23	0
59	MG	1A	3594	1/1	0.95	0.14	-10.18	15,15,15,15	0
59	MG	1A	3840	1/1	0.97	0.14	-10.18	16,16,16,16	0
59	MG	2A	3399	1/1	0.92	0.10	-10.38	55,55,55,55	0
59	MG	2A	3600	1/1	0.97	0.09	-10.54	55,55,55,55	0
59	MG	1a	3318	1/1	0.94	0.14	-10.58	43,43,43,43	0
59	MG	1a	3492	1/1	0.85	0.08	-10.76	75,75,75,75	0
59	MG	1A	4037	1/1	0.94	0.08	-11.02	40,40,40,40	0
59	MG	1A	3547	1/1	0.90	0.12	-11.20	29,29,29,29	0
59	MG	1A	3884	1/1	0.92	0.15	-11.54	41,41,41,41	0
59	MG	2A	3168	1/1	0.92	0.09	-11.59	59,59,59,59	0
59	MG	1A	4059	1/1	0.91	0.15	-11.62	57,57,57,57	0
59	MG	1A	4167	1/1	0.86	0.11	-11.65	19,19,19,19	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3375	1/1	0.94	0.06	-12.15	45,45,45,45	0
59	MG	1A	3756	1/1	0.94	0.11	-12.25	26,26,26,26	0
59	MG	2A	3049	1/1	0.95	0.08	-12.94	44,44,44,44	0
59	MG	1A	3977	1/1	0.93	0.12	-13.05	37,37,37,37	0
59	MG	2A	3750	1/1	0.91	0.08	-13.24	51,51,51,51	0
59	MG	2A	3420	1/1	0.94	0.15	-13.50	28,28,28,28	0
59	MG	1A	3751	1/1	0.96	0.11	-13.95	21,21,21,21	0
59	MG	1A	3987	1/1	0.80	0.07	-14.51	47,47,47,47	0
59	MG	2A	3077	1/1	0.95	0.14	-14.63	46,46,46,46	0
59	MG	1A	3752	1/1	0.93	0.08	-14.65	27,27,27,27	0
59	MG	2A	3429	1/1	0.96	0.09	-14.73	56,56,56,56	0
59	MG	2A	3701	1/1	0.96	0.12	-14.89	26,26,26,26	0
59	MG	1A	3847	1/1	0.94	0.11	-15.31	25,25,25,25	0
59	MG	1A	3683	1/1	0.95	0.10	-16.98	29,29,29,29	0
59	MG	1A	3737	1/1	0.93	0.07	-17.00	34,34,34,34	0
59	MG	1A	3940	1/1	0.94	0.11	-19.14	27,27,27,27	0
59	MG	1A	3084	1/1	0.90	0.11	-20.37	31,31,31,31	0
59	MG	1A	4111	1/1	0.82	0.10	-	63,63,63,63	0
59	MG	1A	3081	1/1	0.86	0.20	-	51,51,51,51	0
59	MG	2A	3511	1/1	0.91	0.40	-	50,50,50,50	0
59	MG	2A	3038	1/1	0.93	0.16	-	48,48,48,48	0
59	MG	2A	3109	1/1	0.90	0.41	-	43,43,43,43	0
59	MG	2A	3819	1/1	0.92	0.11	-	47,47,47,47	0
59	MG	2A	3320	1/1	0.75	0.17	-	59,59,59,59	0
59	MG	1a	3446	1/1	0.94	0.11	-	48,48,48,48	0
59	MG	2a	3087	1/1	0.74	0.17	-	65,65,65,65	0
59	MG	1A	4123	1/1	0.94	0.12	-	46,46,46,46	0
59	MG	2A	3195	1/1	0.96	0.50	-	36,36,36,36	0
59	MG	2A	3702	1/1	0.54	0.10	-	64,64,64,64	0
59	MG	1A	4003	1/1	0.69	0.18	-	54,54,54,54	0
59	MG	2A	3454	1/1	0.84	0.14	-	36,36,36,36	0
59	MG	2a	3083	1/1	0.93	0.14	-	59,59,59,59	0
59	MG	1a	3469	1/1	0.96	0.10	-	48,48,48,48	0
59	MG	1A	3550	1/1	0.96	0.15	-	14,14,14,14	0
59	MG	1A	3604	1/1	0.95	0.12	-	27,27,27,27	0
59	MG	2A	3605	1/1	0.89	0.15	-	55,55,55,55	0
59	MG	2A	3017	1/1	0.90	0.56	-	39,39,39,39	0
59	MG	1A	3205	1/1	0.93	0.31	-	24,24,24,24	0
59	MG	1A	3411	1/1	0.89	0.14	-	48,48,48,48	0
59	MG	2A	3457	1/1	0.71	0.16	-	65,65,65,65	0
59	MG	1A	3369	1/1	0.97	0.64	-	31,31,31,31	0
59	MG	2a	3173	1/1	0.94	0.10	-	61,61,61,61	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3898	1/1	0.89	0.12	-	49,49,49,49	0
59	MG	1A	4097	1/1	0.90	0.10	-	45,45,45,45	0
59	MG	1A	3668	1/1	0.98	0.10	-	38,38,38,38	0
59	MG	2A	3072	1/1	0.82	0.29	-	42,42,42,42	0
59	MG	2a	3180	1/1	0.52	0.16	-	74,74,74,74	0
59	MG	2A	3727	1/1	0.97	0.09	-	31,31,31,31	0
59	MG	2a	3202	1/1	0.92	0.31	-	67,67,67,67	0
59	MG	1A	3938	1/1	0.98	0.12	-	22,22,22,22	0
59	MG	1A	3421	1/1	0.92	0.34	-	33,33,33,33	0
59	MG	2U	202	1/1	0.75	0.36	-	69,69,69,69	0
59	MG	2A	3317	1/1	0.81	0.14	-	51,51,51,51	0
59	MG	1a	3350	1/1	0.65	0.15	-	61,61,61,61	0
59	MG	1A	4117	1/1	0.77	0.14	-	63,63,63,63	0
59	MG	2q	203	1/1	0.91	0.14	-	53,53,53,53	0
59	MG	1D	312	1/1	0.96	0.32	-	49,49,49,49	0
59	MG	2A	3747	1/1	0.85	0.22	-	61,61,61,61	0
59	MG	1A	4036	1/1	0.84	0.10	-	46,46,46,46	0
59	MG	2a	3108	1/1	0.77	0.12	-	62,62,62,62	0
59	MG	1a	3460	1/1	0.88	0.12	-	58,58,58,58	0
59	MG	2B	3013	1/1	0.96	0.23	-	63,63,63,63	0
59	MG	1A	4108	1/1	0.97	0.11	-	32,32,32,32	0
59	MG	1A	3813	1/1	0.92	0.14	-	15,15,15,15	0
59	MG	2a	3037	1/1	0.88	0.11	-	74,74,74,74	0
59	MG	1A	3885	1/1	0.88	0.12	-	47,47,47,47	0
59	MG	2A	3448	1/1	0.68	0.15	-	70,70,70,70	0
59	MG	1A	4145	1/1	0.81	0.15	-	45,45,45,45	0
59	MG	2a	3072	1/1	0.97	0.27	-	50,50,50,50	0
59	MG	1a	3400	1/1	0.94	0.16	-	54,54,54,54	0
59	MG	1A	3880	1/1	0.76	0.13	-	64,64,64,64	0
59	MG	1A	3599	1/1	0.94	0.17	-	21,21,21,21	0
59	MG	1A	3333	1/1	0.75	0.47	-	35,35,35,35	0
59	MG	2A	3393	1/1	0.90	0.12	-	27,27,27,27	0
59	MG	1A	3283	1/1	0.93	0.15	-	45,45,45,45	0
59	MG	2A	3309	1/1	0.70	0.09	-	57,57,57,57	0
59	MG	1A	4184	1/1	0.89	0.38	-	31,31,31,31	0
59	MG	1A	4014	1/1	0.56	0.13	-	65,65,65,65	0
59	MG	1A	3790	1/1	0.93	0.09	-	35,35,35,35	0
59	MG	2a	3198	1/1	0.84	0.20	-	56,56,56,56	0
59	MG	2A	3527	1/1	0.98	0.10	-	65,65,65,65	0
59	MG	2A	3813	1/1	0.79	0.13	-	62,62,62,62	0
59	MG	1A	3324	1/1	0.94	0.50	-	39,39,39,39	0
59	MG	1A	3745	1/1	0.85	0.18	-	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3228	1/1	0.95	0.17	-	36,36,36,36	0
59	MG	2A	3057	1/1	0.72	0.13	-	51,51,51,51	0
59	MG	2A	3302	1/1	0.91	0.32	-	57,57,57,57	0
59	MG	1A	3933	1/1	0.96	0.16	-	25,25,25,25	0
59	MG	2A	3080	1/1	0.91	0.09	-	74,74,74,74	0
59	MG	2A	3261	1/1	0.91	0.13	-	59,59,59,59	0
59	MG	2a	3133	1/1	0.60	0.24	-	67,67,67,67	0
59	MG	1A	3262	1/1	0.91	0.16	-	41,41,41,41	0
59	MG	2A	3166	1/1	0.85	0.18	-	56,56,56,56	0
59	MG	2A	3300	1/1	0.99	0.24	-	46,46,46,46	0
59	MG	2A	3762	1/1	0.97	0.13	-	47,47,47,47	0
59	MG	2a	3058	1/1	0.86	0.18	-	70,70,70,70	0
59	MG	2A	3540	1/1	0.88	0.26	-	48,48,48,48	0
59	MG	2A	3359	1/1	0.94	0.09	-	48,48,48,48	0
59	MG	2A	3585	1/1	0.86	0.12	-	59,59,59,59	0
59	MG	2a	3018	1/1	0.52	0.26	-	67,67,67,67	0
59	MG	1A	3185	1/1	0.93	0.53	-	36,36,36,36	0
59	MG	1A	4045	1/1	0.89	0.07	-	53,53,53,53	0
59	MG	1a	3562	1/1	0.91	0.10	-	63,63,63,63	0
59	MG	2Q	3003	1/1	0.73	0.23	-	54,54,54,54	0
59	MG	2A	3062	1/1	0.80	0.68	-	43,43,43,43	0
59	MG	2O	8001	1/1	0.90	0.12	-	53,53,53,53	0
59	MG	12	3001	1/1	0.88	0.22	-	40,40,40,40	0
59	MG	1A	3522	1/1	0.96	0.47	-	28,28,28,28	0
59	MG	1A	3527	1/1	0.95	0.44	-	22,22,22,22	0
59	MG	2A	3270	1/1	0.88	0.13	-	62,62,62,62	0
59	MG	1w	108	1/1	0.80	0.18	-	59,59,59,59	0
59	MG	2a	3143	1/1	0.83	0.11	-	43,43,43,43	0
59	MG	2a	3011	1/1	0.89	0.19	-	53,53,53,53	0
59	MG	1A	3023	1/1	0.94	0.47	-	24,24,24,24	0
59	MG	1A	3747	1/1	0.92	0.27	-	26,26,26,26	0
59	MG	1A	3610	1/1	0.89	0.14	-	19,19,19,19	0
59	MG	2A	3158	1/1	0.91	0.18	-	50,50,50,50	0
59	MG	1A	3667	1/1	0.94	0.12	-	36,36,36,36	0
59	MG	2A	3488	1/1	0.91	0.27	-	54,54,54,54	0
59	MG	1A	3978	1/1	0.98	0.11	-	38,38,38,38	0
59	MG	1a	3438	1/1	0.81	0.11	-	55,55,55,55	0
59	MG	2A	3612	1/1	0.94	0.13	-	44,44,44,44	0
59	MG	2A	3240	1/1	0.63	0.27	-	62,62,62,62	0
59	MG	1A	4005	1/1	0.85	0.17	-	52,52,52,52	0
59	MG	1A	3195	1/1	0.70	0.23	-	48,48,48,48	0
59	MG	1A	3501	1/1	0.97	0.15	-	41,41,41,41	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1a	3534	1/1	0.89	0.10	-	45,45,45,45	0
59	MG	2a	3136	1/1	0.82	0.13	-	64,64,64,64	0
59	MG	1A	3853	1/1	0.87	0.08	-	33,33,33,33	0
59	MG	1W	205	1/1	0.97	0.33	-	23,23,23,23	0
59	MG	2A	3634	1/1	0.90	0.31	-	41,41,41,41	0
59	MG	2A	3390	1/1	0.96	0.08	-	60,60,60,60	0
59	MG	2A	3764	1/1	0.95	0.14	-	38,38,38,38	0
59	MG	1A	4073	1/1	0.91	0.09	-	40,40,40,40	0
59	MG	2A	3405	1/1	0.93	0.13	-	65,65,65,65	0
59	MG	1E	311	1/1	0.92	0.21	-	15,15,15,15	0
59	MG	1a	3454	1/1	0.92	0.10	-	46,46,46,46	0
59	MG	1A	3217	1/1	0.83	0.15	-	47,47,47,47	0
59	MG	1A	4013	1/1	0.83	0.14	-	50,50,50,50	0
59	MG	1A	3586	1/1	0.84	0.23	-	49,49,49,49	0
59	MG	1A	3789	1/1	0.88	0.16	-	56,56,56,56	0
59	MG	1A	4101	1/1	0.90	0.28	-	41,41,41,41	0
59	MG	2A	3438	1/1	0.92	0.11	-	77,77,77,77	0
59	MG	1A	3920	1/1	0.93	0.08	-	45,45,45,45	0
59	MG	1A	3163	1/1	0.88	0.22	-	48,48,48,48	0
59	MG	2A	3206	1/1	0.93	0.17	-	50,50,50,50	0
59	MG	1A	3803	1/1	0.97	0.32	-	33,33,33,33	0
59	MG	1A	3863	1/1	0.92	0.14	-	9,9,9,9	0
59	MG	1A	3846	1/1	0.84	0.18	-	57,57,57,57	0
59	MG	2a	3205	1/1	0.71	0.13	-	60,60,60,60	0
59	MG	2A	3799	1/1	0.88	0.13	-	60,60,60,60	0
59	MG	2A	3245	1/1	0.91	0.30	-	48,48,48,48	0
59	MG	2A	3695	1/1	0.87	0.09	-	53,53,53,53	0
59	MG	1A	3295	1/1	0.90	0.29	-	47,47,47,47	0
59	MG	2a	3006	1/1	0.80	0.32	-	76,76,76,76	0
59	MG	2A	3793	1/1	0.89	0.08	-	62,62,62,62	0
59	MG	1A	3448	1/1	0.94	0.63	-	36,36,36,36	0
59	MG	1A	4183	1/1	0.91	0.17	-	34,34,34,34	0
59	MG	1A	3628	1/1	0.93	0.12	-	21,21,21,21	0
59	MG	1a	3338	1/1	0.93	0.14	-	51,51,51,51	0
59	MG	2a	3033	1/1	0.94	0.18	-	56,56,56,56	0
59	MG	1A	3085	1/1	0.98	0.21	-	15,15,15,15	0
59	MG	1A	3325	1/1	0.91	0.46	-	31,31,31,31	0
59	MG	1A	3758	1/1	0.96	0.11	-	37,37,37,37	0
59	MG	2A	3084	1/1	0.74	0.14	-	76,76,76,76	0
59	MG	2A	3236	1/1	0.88	0.30	-	57,57,57,57	0
59	MG	1A	3654	1/1	0.82	0.14	-	54,54,54,54	0
59	MG	2l	202	1/1	0.84	0.17	-	71,71,71,71	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1a	3331	1/1	0.76	0.41	-	60,60,60,60	0
59	MG	1A	3381	1/1	0.97	0.39	-	41,41,41,41	0
59	MG	1A	3669	1/1	0.97	0.07	-	38,38,38,38	0
59	MG	1a	3441	1/1	0.78	0.12	-	53,53,53,53	0
59	MG	2A	3678	1/1	0.55	0.17	-	64,64,64,64	0
59	MG	1A	4047	1/1	0.96	0.26	-	31,31,31,31	0
59	MG	1A	4098	1/1	0.93	0.28	-	37,37,37,37	0
59	MG	2A	3494	1/1	0.71	0.13	-	58,58,58,58	0
59	MG	1A	3014	1/1	0.89	0.20	-	38,38,38,38	0
59	MG	1A	3091	1/1	0.94	0.14	-	17,17,17,17	0
59	MG	1A	3374	1/1	0.97	0.50	-	27,27,27,27	0
59	MG	1A	3109	1/1	0.88	0.18	-	46,46,46,46	0
59	MG	2A	3555	1/1	0.62	0.16	-	61,61,61,61	0
59	MG	2A	3310	1/1	0.83	0.16	-	53,53,53,53	0
59	MG	2A	3733	1/1	0.79	0.30	-	71,71,71,71	0
59	MG	1A	3326	1/1	0.79	0.22	-	49,49,49,49	0
59	MG	2a	3163	1/1	0.54	0.20	-	91,91,91,91	0
59	MG	2a	3020	1/1	0.93	0.28	-	61,61,61,61	0
59	MG	2A	3588	1/1	0.90	0.09	-	60,60,60,60	0
59	MG	1x	101	1/1	0.94	0.14	-	38,38,38,38	0
59	MG	1A	3463	1/1	0.90	0.34	-	24,24,24,24	0
59	MG	1A	3970	1/1	0.73	0.09	-	41,41,41,41	0
59	MG	2A	3312	1/1	0.83	0.13	-	55,55,55,55	0
59	MG	1A	3680	1/1	0.87	0.13	-	53,53,53,53	0
59	MG	1A	3906	1/1	0.95	0.09	-	36,36,36,36	0
59	MG	1a	3537	1/1	0.84	0.10	-	46,46,46,46	0
59	MG	1A	3284	1/1	0.89	0.15	-	37,37,37,37	0
59	MG	1A	3431	1/1	0.96	0.21	-	45,45,45,45	0
59	MG	2A	3258	1/1	0.93	0.14	-	68,68,68,68	0
59	MG	1a	3547	1/1	0.85	0.10	-	48,48,48,48	0
59	MG	2A	3225	1/1	0.87	0.11	-	53,53,53,53	0
59	MG	1A	3287	1/1	0.95	0.51	-	38,38,38,38	0
59	MG	1A	3332	1/1	0.91	0.51	-	35,35,35,35	0
59	MG	2A	3647	1/1	0.91	0.05	-	39,39,39,39	0
59	MG	2A	3197	1/1	0.86	0.20	-	50,50,50,50	0
59	MG	2A	3483	1/1	0.92	0.16	-	36,36,36,36	0
59	MG	1A	3719	1/1	0.91	0.09	-	44,44,44,44	0
59	MG	1A	4181	1/1	0.93	0.56	-	33,33,33,33	0
59	MG	1l	201	1/1	0.96	0.19	-	30,30,30,30	0
59	MG	1a	3462	1/1	0.85	0.11	-	72,72,72,72	0
59	MG	2A	3357	1/1	0.85	0.33	-	55,55,55,55	0
59	MG	1a	3388	1/1	0.76	0.13	-	61,61,61,61	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2a	3048	1/1	0.93	0.16	-	42,42,42,42	0
59	MG	1a	3501	1/1	0.80	0.07	-	57,57,57,57	0
59	MG	1A	3841	1/1	0.94	0.14	-	24,24,24,24	0
59	MG	2A	3715	1/1	0.82	0.09	-	61,61,61,61	0
59	MG	1A	3642	1/1	0.83	0.17	-	56,56,56,56	0
59	MG	1A	4032	1/1	0.96	0.11	-	48,48,48,48	0
59	MG	1a	3552	1/1	0.95	0.18	-	55,55,55,55	0
59	MG	1A	3896	1/1	0.82	0.26	-	54,54,54,54	0
59	MG	2A	3459	1/1	0.98	0.17	-	55,55,55,55	0
59	MG	1A	3998	1/1	0.95	0.15	-	13,13,13,13	0
59	MG	1A	3671	1/1	0.92	0.09	-	57,57,57,57	0
59	MG	1A	3948	1/1	0.61	0.11	-	69,69,69,69	0
59	MG	1A	3479	1/1	0.73	0.30	-	69,69,69,69	0
59	MG	1A	3838	1/1	0.96	0.35	-	26,26,26,26	0
59	MG	1A	4021	1/1	0.95	0.14	-	27,27,27,27	0
59	MG	2a	3078	1/1	0.89	0.12	-	70,70,70,70	0
59	MG	2A	3506	1/1	0.99	0.17	-	42,42,42,42	0
59	MG	1B	218	1/1	0.94	0.19	-	36,36,36,36	0
59	MG	1A	3990	1/1	0.63	0.43	-	46,46,46,46	0
59	MG	2a	3081	1/1	0.86	0.14	-	67,67,67,67	0
59	MG	1A	3782	1/1	0.80	0.08	-	60,60,60,60	0
59	MG	2A	3656	1/1	0.88	0.08	-	40,40,40,40	0
59	MG	2w	3007	1/1	0.72	0.15	-	72,72,72,72	0
59	MG	1A	3399	1/1	0.92	0.12	-	67,67,67,67	0
59	MG	1A	3442	1/1	0.87	0.20	-	63,63,63,63	0
59	MG	2A	3346	1/1	0.79	0.18	-	62,62,62,62	0
59	MG	2A	3013	1/1	0.94	0.29	-	47,47,47,47	0
59	MG	1a	3344	1/1	0.83	0.11	-	56,56,56,56	0
59	MG	1A	3341	1/1	0.91	0.13	-	45,45,45,45	0
59	MG	1a	3439	1/1	0.90	0.08	-	63,63,63,63	0
59	MG	1A	4011	1/1	0.88	0.17	-	59,59,59,59	0
59	MG	2A	3839	1/1	0.88	0.49	-	39,39,39,39	0
59	MG	1B	214	1/1	0.84	0.09	-	55,55,55,55	0
59	MG	2A	3688	1/1	0.87	0.07	-	38,38,38,38	0
59	MG	2a	3199	1/1	0.88	0.13	-	58,58,58,58	0
59	MG	2A	3833	1/1	0.92	0.10	-	47,47,47,47	0
59	MG	2A	3239	1/1	0.77	0.18	-	61,61,61,61	0
59	MG	2A	3255	1/1	0.86	0.13	-	65,65,65,65	0
59	MG	1Q	206	1/1	0.79	0.10	-	39,39,39,39	0
59	MG	1A	4006	1/1	0.91	0.09	-	31,31,31,31	0
59	MG	1a	3366	1/1	0.82	0.19	-	57,57,57,57	0
59	MG	2N	8001	1/1	0.94	0.13	-	45,45,45,45	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1a	3538	1/1	0.98	0.06	-	44,44,44,44	0
59	MG	1A	3353	1/1	0.81	0.84	-	51,51,51,51	0
59	MG	1A	3025	1/1	0.94	0.28	-	21,21,21,21	0
59	MG	1A	3151	1/1	0.96	0.16	-	15,15,15,15	0
59	MG	1A	3595	1/1	0.83	0.20	-	18,18,18,18	0
59	MG	2A	3044	1/1	0.72	0.32	-	62,62,62,62	0
59	MG	2A	3470	1/1	0.92	0.17	-	53,53,53,53	0
59	MG	1A	4126	1/1	0.75	0.12	-	53,53,53,53	0
59	MG	1A	3181	1/1	0.96	0.45	-	25,25,25,25	0
59	MG	1A	4147	1/1	0.91	0.31	-	50,50,50,50	0
59	MG	1A	4132	1/1	0.93	0.31	-	37,37,37,37	0
59	MG	1F	308	1/1	0.85	0.12	-	45,45,45,45	0
59	MG	1A	4139	1/1	0.91	0.10	-	28,28,28,28	0
59	MG	2x	102	1/1	0.84	0.12	-	64,64,64,64	0
59	MG	1A	3372	1/1	0.99	0.53	-	37,37,37,37	0
59	MG	2a	3238	1/1	0.72	0.27	-	62,62,62,62	0
59	MG	2A	3288	1/1	0.86	0.47	-	67,67,67,67	0
59	MG	1A	3658	1/1	0.94	0.08	-	39,39,39,39	0
59	MG	2a	3023	1/1	0.93	0.24	-	54,54,54,54	0
59	MG	2E	303	1/1	0.81	0.14	-	53,53,53,53	0
59	MG	2A	3827	1/1	0.97	0.43	-	62,62,62,62	0
59	MG	1A	4055	1/1	0.93	0.13	-	32,32,32,32	0
59	MG	2a	3120	1/1	0.92	0.11	-	51,51,51,51	0
59	MG	1A	3872	1/1	0.93	0.14	-	48,48,48,48	0
59	MG	1a	3301	1/1	0.96	0.22	-	49,49,49,49	0
59	MG	1A	3811	1/1	0.98	0.10	-	30,30,30,30	0
59	MG	2A	3313	1/1	0.89	0.14	-	54,54,54,54	0
59	MG	1I	3001	1/1	0.84	0.17	-	62,62,62,62	0
59	MG	17	102	1/1	0.94	0.12	-	46,46,46,46	0
59	MG	1A	3243	1/1	0.95	0.66	-	28,28,28,28	0
59	MG	2A	3402	1/1	0.86	0.18	-	57,57,57,57	0
59	MG	1A	4215	1/1	0.96	0.75	-	31,31,31,31	0
59	MG	1A	3113	1/1	0.86	0.40	-	37,37,37,37	0
59	MG	2A	3721	1/1	0.95	0.09	-	17,17,17,17	0
59	MG	2A	3754	1/1	0.89	0.11	-	64,64,64,64	0
59	MG	1A	3497	1/1	0.94	0.24	-	38,38,38,38	0
59	MG	2A	3340	1/1	0.84	0.17	-	59,59,59,59	0
59	MG	2a	3174	1/1	0.91	0.08	-	74,74,74,74	0
59	MG	1a	3408	1/1	0.79	0.25	-	60,60,60,60	0
59	MG	2A	3336	1/1	0.91	0.18	-	57,57,57,57	0
59	MG	2A	3583	1/1	0.82	0.43	-	48,48,48,48	0
59	MG	2A	3458	1/1	0.88	0.07	-	53,53,53,53	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3176	1/1	0.95	0.30	-	35,35,35,35	0
59	MG	2A	3663	1/1	0.43	0.14	-	60,60,60,60	0
59	MG	1A	3031	1/1	0.97	0.63	-	24,24,24,24	0
59	MG	1A	3801	1/1	0.78	0.14	-	34,34,34,34	0
59	MG	1A	3320	1/1	0.95	0.24	-	41,41,41,41	0
59	MG	2A	3815	1/1	0.93	0.05	-	56,56,56,56	0
59	MG	2a	3013	1/1	0.62	0.23	-	68,68,68,68	0
59	MG	1A	3131	1/1	0.92	0.68	-	33,33,33,33	0
59	MG	1a	3367	1/1	0.89	0.38	-	53,53,53,53	0
59	MG	1A	3282	1/1	0.96	0.23	-	36,36,36,36	0
59	MG	2a	3222	1/1	0.95	0.11	-	57,57,57,57	0
59	MG	1A	3515	1/1	0.74	0.31	-	71,71,71,71	0
59	MG	1A	3612	1/1	0.78	0.13	-	45,45,45,45	0
59	MG	1A	3228	1/1	0.86	0.20	-	56,56,56,56	0
59	MG	1A	3634	1/1	0.95	0.14	-	12,12,12,12	0
59	MG	2A	3167	1/1	0.91	0.17	-	57,57,57,57	0
59	MG	2B	3002	1/1	0.94	0.15	-	56,56,56,56	0
59	MG	1A	4057	1/1	0.72	0.17	-	50,50,50,50	0
59	MG	1A	3379	1/1	0.91	0.23	-	37,37,37,37	0
59	MG	2A	3770	1/1	0.96	0.16	-	53,53,53,53	0
59	MG	1A	3227	1/1	0.79	0.18	-	50,50,50,50	0
59	MG	1a	3337	1/1	0.94	0.25	-	55,55,55,55	0
59	MG	2a	3003	1/1	0.82	0.19	-	54,54,54,54	0
59	MG	1A	3505	1/1	0.83	0.17	-	51,51,51,51	0
59	MG	1A	4217	1/1	0.94	0.49	-	35,35,35,35	0
59	MG	1A	3740	1/1	0.76	0.12	-	49,49,49,49	0
59	MG	1A	3432	1/1	0.91	0.45	-	47,47,47,47	0
59	MG	1A	3783	1/1	0.92	0.11	-	64,64,64,64	0
59	MG	1a	3386	1/1	0.93	0.27	-	50,50,50,50	0
59	MG	2a	3093	1/1	0.83	0.25	-	71,71,71,71	0
59	MG	1A	3716	1/1	0.98	0.15	-	34,34,34,34	0
59	MG	1a	3391	1/1	0.95	0.21	-	55,55,55,55	0
59	MG	2A	3250	1/1	0.81	0.23	-	56,56,56,56	0
59	MG	2a	3210	1/1	0.92	0.09	-	65,65,65,65	0
59	MG	1A	3828	1/1	0.84	0.14	-	46,46,46,46	0
59	MG	2A	3112	1/1	0.86	0.11	-	33,33,33,33	0
59	MG	2a	3060	1/1	0.95	0.14	-	49,49,49,49	0
59	MG	2a	3118	1/1	0.74	0.29	-	71,71,71,71	0
59	MG	1a	3409	1/1	0.82	0.12	-	74,74,74,74	0
59	MG	2A	3325	1/1	0.69	0.14	-	68,68,68,68	0
59	MG	1a	3306	1/1	0.97	0.34	-	48,48,48,48	0
59	MG	1A	3816	1/1	0.98	0.20	-	35,35,35,35	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3251	1/1	0.90	0.47	-	47,47,47,47	0
59	MG	2A	3726	1/1	0.76	0.35	-	56,56,56,56	0
59	MG	1A	3750	1/1	0.94	0.18	-	42,42,42,42	0
59	MG	1a	3304	1/1	0.81	0.10	-	58,58,58,58	0
59	MG	1A	3823	1/1	0.96	0.14	-	59,59,59,59	0
59	MG	1A	4159	1/1	0.96	0.19	-	34,34,34,34	0
59	MG	1a	3536	1/1	0.93	0.08	-	58,58,58,58	0
59	MG	2A	3803	1/1	0.79	0.11	-	50,50,50,50	0
59	MG	2A	3606	1/1	0.43	0.12	-	73,73,73,73	0
59	MG	2a	3227	1/1	0.95	0.15	-	45,45,45,45	0
59	MG	1A	3259	1/1	0.96	0.17	-	41,41,41,41	0
59	MG	1A	3384	1/1	0.88	0.16	-	40,40,40,40	0
59	MG	1A	3112	1/1	0.93	0.37	-	33,33,33,33	0
59	MG	1a	3575	1/1	0.85	0.22	-	55,55,55,55	0
59	MG	1A	3626	1/1	0.93	0.16	-	24,24,24,24	0
59	MG	18	103	1/1	0.94	0.21	-	38,38,38,38	0
59	MG	2w	3004	1/1	0.74	0.21	-	75,75,75,75	0
59	MG	1A	3945	1/1	0.82	0.09	-	59,59,59,59	0
59	MG	1A	3573	1/1	0.84	0.14	-	32,32,32,32	0
59	MG	2A	3710	1/1	0.96	0.15	-	34,34,34,34	0
59	MG	1a	3374	1/1	0.92	0.15	-	53,53,53,53	0
59	MG	2A	3093	1/1	0.87	0.16	-	56,56,56,56	0
59	MG	1A	3357	1/1	0.97	0.80	-	45,45,45,45	0
59	MG	2A	3735	1/1	0.85	0.14	-	62,62,62,62	0
59	MG	1a	3431	1/1	0.96	0.13	-	38,38,38,38	0
59	MG	1A	3688	1/1	0.91	0.21	-	43,43,43,43	0
59	MG	1A	3659	1/1	0.98	0.15	-	50,50,50,50	0
59	MG	1U	204	1/1	0.93	0.10	-	36,36,36,36	0
59	MG	1A	3796	1/1	0.90	0.10	-	45,45,45,45	0
59	MG	1A	3526	1/1	0.80	0.16	-	59,59,59,59	0
59	MG	2A	3581	1/1	0.94	0.16	-	51,51,51,51	0
59	MG	1a	3578	1/1	0.90	0.50	-	59,59,59,59	0
59	MG	2A	3304	1/1	0.78	0.24	-	56,56,56,56	0
59	MG	1a	3520	1/1	0.78	0.07	-	60,60,60,60	0
59	MG	1A	4189	1/1	0.94	0.12	-	44,44,44,44	0
59	MG	1A	3361	1/1	0.95	0.33	-	35,35,35,35	0
59	MG	1A	3746	1/1	0.81	0.23	-	48,48,48,48	0
59	MG	1a	3514	1/1	0.71	0.13	-	50,50,50,50	0
59	MG	1A	4105	1/1	0.96	0.12	-	53,53,53,53	0
59	MG	2A	3185	1/1	0.72	0.49	-	53,53,53,53	0
59	MG	2A	3246	1/1	0.89	0.16	-	67,67,67,67	0
59	MG	1N	3005	1/1	0.94	0.24	-	51,51,51,51	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3408	1/1	0.85	0.27	-	32,32,32,32	0
59	MG	1A	3871	1/1	0.72	0.09	-	54,54,54,54	0
59	MG	1A	3892	1/1	0.91	0.29	-	62,62,62,62	0
59	MG	2A	3154	1/1	0.84	0.15	-	58,58,58,58	0
59	MG	1A	4188	1/1	0.83	0.21	-	55,55,55,55	0
59	MG	1A	4103	1/1	0.90	0.13	-	42,42,42,42	0
59	MG	1a	3565	1/1	0.94	0.14	-	35,35,35,35	0
59	MG	1A	4134	1/1	0.87	0.15	-	52,52,52,52	0
59	MG	1A	3377	1/1	0.93	0.55	-	62,62,62,62	0
59	MG	1A	3890	1/1	0.84	0.11	-	50,50,50,50	0
59	MG	1A	4142	1/1	0.93	0.11	-	47,47,47,47	0
59	MG	1A	3446	1/1	0.93	0.38	-	35,35,35,35	0
59	MG	1a	3368	1/1	0.88	0.17	-	63,63,63,63	0
59	MG	1A	3733	1/1	0.79	0.07	-	47,47,47,47	0
59	MG	1A	3848	1/1	0.87	0.10	-	72,72,72,72	0
59	MG	1A	3829	1/1	0.96	0.17	-	38,38,38,38	0
59	MG	1A	3439	1/1	0.93	0.50	-	39,39,39,39	0
59	MG	1A	3392	1/1	0.69	0.45	-	52,52,52,52	0
59	MG	2A	3141	1/1	0.85	0.13	-	38,38,38,38	0
59	MG	1A	3805	1/1	0.94	0.39	-	30,30,30,30	0
59	MG	2A	3720	1/1	0.80	0.09	-	42,42,42,42	0
59	MG	1a	3436	1/1	0.75	0.17	-	56,56,56,56	0
59	MG	2A	3636	1/1	0.86	0.17	-	64,64,64,64	0
59	MG	1a	3361	1/1	0.75	0.19	-	68,68,68,68	0
59	MG	1G	3002	1/1	0.91	0.19	-	53,53,53,53	0
59	MG	1A	3544	1/1	0.96	0.14	-	44,44,44,44	0
59	MG	2A	3626	1/1	0.93	0.15	-	68,68,68,68	0
59	MG	13	3003	1/1	0.93	0.24	-	42,42,42,42	0
59	MG	2A	3426	1/1	0.93	0.14	-	23,23,23,23	0
59	MG	2A	3805	1/1	0.94	0.12	-	53,53,53,53	0
59	MG	2a	3151	1/1	0.78	0.06	-	73,73,73,73	0
59	MG	1F	301	1/1	0.87	0.09	-	50,50,50,50	0
59	MG	2A	3176	1/1	0.93	0.15	-	44,44,44,44	0
59	MG	1A	4056	1/1	0.92	0.14	-	43,43,43,43	0
59	MG	2A	3129	1/1	0.95	0.12	-	48,48,48,48	0
59	MG	1A	3481	1/1	0.77	0.23	-	38,38,38,38	0
59	MG	2A	3513	1/1	0.96	0.14	-	67,67,67,67	0
59	MG	1a	3572	1/1	0.77	0.36	-	63,63,63,63	0
59	MG	2A	3113	1/1	0.95	0.36	-	38,38,38,38	0
59	MG	2A	3068	1/1	0.91	0.10	-	52,52,52,52	0
59	MG	1B	227	1/1	0.94	0.17	-	31,31,31,31	0
59	MG	1a	3539	1/1	0.95	0.13	-	50,50,50,50	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	4099	1/1	0.96	0.09	-	50,50,50,50	0
59	MG	2A	3650	1/1	0.93	0.43	-	60,60,60,60	0
59	MG	1A	3636	1/1	0.93	0.12	-	46,46,46,46	0
59	MG	2A	3645	1/1	0.86	0.10	-	51,51,51,51	0
59	MG	2A	3144	1/1	0.97	0.09	-	65,65,65,65	0
59	MG	2V	201	1/1	0.81	0.25	-	60,60,60,60	0
59	MG	1V	202	1/1	0.90	0.10	-	54,54,54,54	0
59	MG	1a	3515	1/1	0.92	0.09	-	39,39,39,39	0
59	MG	1A	4089	1/1	0.87	0.16	-	42,42,42,42	0
59	MG	1A	3312	1/1	0.91	0.65	-	33,33,33,33	0
59	MG	2A	3406	1/1	0.98	0.12	-	39,39,39,39	0
59	MG	2a	3116	1/1	0.95	0.09	-	65,65,65,65	0
59	MG	1A	3648	1/1	0.95	0.20	-	31,31,31,31	0
59	MG	2A	3039	1/1	0.97	0.15	-	23,23,23,23	0
59	MG	1A	3664	1/1	0.93	0.09	-	46,46,46,46	0
59	MG	2A	3332	1/1	0.79	0.29	-	64,64,64,64	0
59	MG	1A	3248	1/1	0.96	0.30	-	36,36,36,36	0
59	MG	1A	3299	1/1	0.90	0.19	-	49,49,49,49	0
59	MG	1A	3356	1/1	0.90	0.21	-	38,38,38,38	0
59	MG	1A	3012	1/1	0.98	0.16	-	18,18,18,18	0
59	MG	2A	3705	1/1	0.90	0.07	-	49,49,49,49	0
59	MG	2A	3654	1/1	0.85	0.26	-	62,62,62,62	0
59	MG	2A	3751	1/1	0.95	0.14	-	50,50,50,50	0
59	MG	1A	3419	1/1	0.94	0.65	-	30,30,30,30	0
59	MG	1A	4066	1/1	0.94	0.12	-	47,47,47,47	0
59	MG	2A	3114	1/1	0.95	0.08	-	39,39,39,39	0
59	MG	1A	3456	1/1	0.89	0.41	-	50,50,50,50	0
59	MG	1A	3922	1/1	0.97	0.11	-	41,41,41,41	0
59	MG	2A	3071	1/1	0.85	0.35	-	50,50,50,50	0
59	MG	1A	3621	1/1	0.96	0.17	-	25,25,25,25	0
59	MG	1A	3563	1/1	0.99	0.08	-	37,37,37,37	0
59	MG	1a	3558	1/1	0.56	0.14	-	63,63,63,63	0
59	MG	1x	112	1/1	0.92	0.20	-	47,47,47,47	0
59	MG	1A	3725	1/1	0.97	0.16	-	41,41,41,41	0
59	MG	1A	4041	1/1	0.95	0.10	-	42,42,42,42	0
59	MG	2A	3059	1/1	0.89	0.15	-	37,37,37,37	0
59	MG	2A	3153	1/1	0.89	0.27	-	57,57,57,57	0
59	MG	1A	3637	1/1	0.97	0.15	-	18,18,18,18	0
59	MG	2A	3028	1/1	0.95	0.17	-	25,25,25,25	0
59	MG	2A	3254	1/1	0.84	0.28	-	47,47,47,47	0
59	MG	2a	3134	1/1	0.73	0.11	-	80,80,80,80	0
59	MG	10	104	1/1	0.97	0.23	-	39,39,39,39	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	4025	1/1	0.92	0.12	-	37,37,37,37	0
59	MG	1A	3375	1/1	0.94	0.22	-	29,29,29,29	0
59	MG	1A	3770	1/1	0.96	0.15	-	40,40,40,40	0
59	MG	1A	3454	1/1	0.84	0.17	-	55,55,55,55	0
59	MG	1A	3574	1/1	0.90	0.19	-	52,52,52,52	0
59	MG	2A	3722	1/1	0.81	0.23	-	72,72,72,72	0
59	MG	1A	3044	1/1	0.92	0.35	-	45,45,45,45	0
59	MG	1a	3523	1/1	0.97	0.06	-	54,54,54,54	0
59	MG	1A	3178	1/1	0.79	0.23	-	61,61,61,61	0
59	MG	2A	3146	1/1	0.52	0.16	-	78,78,78,78	0
59	MG	1A	3079	1/1	0.94	0.14	-	46,46,46,46	0
59	MG	1A	4084	1/1	0.88	0.21	-	45,45,45,45	0
59	MG	2A	3201	1/1	0.88	0.57	-	43,43,43,43	0
59	MG	1A	3866	1/1	0.97	0.13	-	15,15,15,15	0
59	MG	1A	4033	1/1	0.92	0.09	-	45,45,45,45	0
59	MG	1A	4127	1/1	0.93	0.58	-	44,44,44,44	0
59	MG	1A	4028	1/1	0.93	0.18	-	50,50,50,50	0
59	MG	1A	3428	1/1	0.92	0.13	-	56,56,56,56	0
59	MG	1A	3430	1/1	0.91	0.57	-	32,32,32,32	0
59	MG	1A	3578	1/1	0.86	0.10	-	73,73,73,73	0
59	MG	2A	3224	1/1	0.75	0.23	-	49,49,49,49	0
59	MG	1A	3093	1/1	0.98	0.11	-	35,35,35,35	0
59	MG	2a	3079	1/1	0.74	0.14	-	67,67,67,67	0
59	MG	2a	3095	1/1	0.92	0.31	-	52,52,52,52	0
59	MG	1a	3406	1/1	0.84	0.21	-	50,50,50,50	0
59	MG	2A	3368	1/1	0.89	0.39	-	63,63,63,63	0
59	MG	2q	202	1/1	0.95	0.15	-	57,57,57,57	0
59	MG	2a	3014	1/1	0.74	0.20	-	57,57,57,57	0
59	MG	1A	4002	1/1	0.88	0.11	-	44,44,44,44	0
59	MG	2a	3139	1/1	0.83	0.17	-	65,65,65,65	0
59	MG	2A	3208	1/1	0.94	0.11	-	71,71,71,71	0
59	MG	1P	204	1/1	0.82	0.43	-	41,41,41,41	0
59	MG	1A	3013	1/1	0.92	0.22	-	44,44,44,44	0
59	MG	2A	3674	1/1	0.93	0.15	-	61,61,61,61	0
59	MG	1A	3069	1/1	0.97	0.34	-	28,28,28,28	0
59	MG	2A	3460	1/1	0.99	0.13	-	25,25,25,25	0
59	MG	10	101	1/1	0.91	0.46	-	41,41,41,41	0
59	MG	1A	3685	1/1	0.99	0.13	-	14,14,14,14	0
59	MG	2A	3648	1/1	0.82	0.24	-	60,60,60,60	0
59	MG	1A	3930	1/1	0.87	0.06	-	71,71,71,71	0
59	MG	1a	3540	1/1	0.84	0.09	-	36,36,36,36	0
59	MG	1A	3905	1/1	0.87	0.22	-	71,71,71,71	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	15	103	1/1	0.97	0.10	-	54,54,54,54	0
59	MG	1A	4024	1/1	0.93	0.14	-	46,46,46,46	0
59	MG	2A	3173	1/1	0.82	0.20	-	62,62,62,62	0
59	MG	2A	3067	1/1	0.68	0.30	-	55,55,55,55	0
59	MG	2A	3703	1/1	0.95	0.09	-	30,30,30,30	0
59	MG	1A	3080	1/1	0.95	0.16	-	16,16,16,16	0
59	MG	1A	3732	1/1	0.81	0.12	-	60,60,60,60	0
59	MG	1A	3056	1/1	0.98	0.19	-	24,24,24,24	0
59	MG	1A	4130	1/1	0.85	0.09	-	65,65,65,65	0
59	MG	1G	3005	1/1	0.97	0.12	-	51,51,51,51	0
59	MG	1A	3794	1/1	0.95	0.14	-	57,57,57,57	0
59	MG	2A	3330	1/1	0.86	0.12	-	53,53,53,53	0
59	MG	1A	3152	1/1	0.99	0.15	-	25,25,25,25	0
59	MG	2A	3219	1/1	0.85	0.20	-	54,54,54,54	0
59	MG	1A	3314	1/1	0.85	0.10	-	47,47,47,47	0
59	MG	1A	3521	1/1	0.75	0.12	-	54,54,54,54	0
59	MG	1B	234	1/1	0.88	0.18	-	54,54,54,54	0
59	MG	2A	3604	1/1	0.90	0.12	-	46,46,46,46	0
59	MG	2A	3465	1/1	0.96	0.08	-	54,54,54,54	0
59	MG	1A	4110	1/1	0.66	0.11	-	62,62,62,62	0
59	MG	2a	3053	1/1	0.91	0.16	-	58,58,58,58	0
59	MG	1A	3480	1/1	0.81	0.14	-	40,40,40,40	0
59	MG	1A	3098	1/1	0.95	0.61	-	30,30,30,30	0
59	MG	1A	3302	1/1	0.83	0.43	-	54,54,54,54	0
59	MG	1A	4077	1/1	0.81	0.16	-	59,59,59,59	0
59	MG	2a	3175	1/1	0.95	0.20	-	58,58,58,58	0
59	MG	2A	3358	1/1	0.76	0.17	-	56,56,56,56	0
59	MG	2A	3294	1/1	0.86	0.18	-	47,47,47,47	0
59	MG	1A	3692	1/1	0.97	0.15	-	37,37,37,37	0
59	MG	2A	3428	1/1	0.93	0.17	-	63,63,63,63	0
59	MG	2a	3029	1/1	0.90	0.15	-	58,58,58,58	0
59	MG	1a	3557	1/1	0.60	0.18	-	63,63,63,63	0
59	MG	2a	3140	1/1	0.95	0.09	-	68,68,68,68	0
59	MG	2A	3281	1/1	0.82	0.12	-	67,67,67,67	0
59	MG	2a	3028	1/1	0.92	0.14	-	64,64,64,64	0
59	MG	2a	3186	1/1	0.91	0.20	-	69,69,69,69	0
59	MG	1A	3321	1/1	0.67	0.20	-	49,49,49,49	0
59	MG	1A	3291	1/1	0.85	0.14	-	43,43,43,43	0
59	MG	1A	3780	1/1	0.88	0.16	-	10,10,10,10	0
59	MG	1A	4162	1/1	0.83	0.15	-	54,54,54,54	0
59	MG	1A	3830	1/1	0.91	0.20	-	40,40,40,40	0
59	MG	2w	3005	1/1	0.77	0.12	-	73,73,73,73	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3054	1/1	0.96	0.13	-	43,43,43,43	0
59	MG	1a	3471	1/1	0.90	0.13	-	51,51,51,51	0
59	MG	2a	3036	1/1	0.59	0.21	-	71,71,71,71	0
59	MG	2a	3160	1/1	0.88	0.11	-	73,73,73,73	0
59	MG	2a	3225	1/1	0.94	0.08	-	59,59,59,59	0
59	MG	1a	3324	1/1	0.93	0.10	-	57,57,57,57	0
59	MG	1A	4076	1/1	0.76	0.12	-	43,43,43,43	0
59	MG	2A	3493	1/1	0.87	0.14	-	29,29,29,29	0
59	MG	1A	4102	1/1	0.91	0.13	-	50,50,50,50	0
59	MG	1a	3377	1/1	0.96	0.18	-	61,61,61,61	0
59	MG	2A	3403	1/1	0.98	0.14	-	38,38,38,38	0
59	MG	1A	3239	1/1	0.95	0.20	-	49,49,49,49	0
59	MG	1A	3566	1/1	0.97	0.14	-	13,13,13,13	0
59	MG	2A	3796	1/1	0.92	0.10	-	70,70,70,70	0
59	MG	2A	3714	1/1	0.88	0.07	-	66,66,66,66	0
59	MG	1A	4070	1/1	0.92	0.14	-	33,33,33,33	0
59	MG	1a	3517	1/1	0.81	0.10	-	60,60,60,60	0
59	MG	2A	3089	1/1	0.91	0.16	-	78,78,78,78	0
59	MG	20	3002	1/1	0.98	0.31	-	51,51,51,51	0
59	MG	2A	3006	1/1	0.99	0.17	-	30,30,30,30	0
59	MG	1A	3201	1/1	0.80	0.19	-	55,55,55,55	0
59	MG	1A	3153	1/1	0.95	0.48	-	25,25,25,25	0
59	MG	1A	4157	1/1	0.94	0.12	-	24,24,24,24	0
59	MG	1a	3381	1/1	0.75	0.18	-	62,62,62,62	0
59	MG	1A	3124	1/1	0.94	0.16	-	29,29,29,29	0
59	MG	2B	3014	1/1	0.90	0.17	-	67,67,67,67	0
59	MG	2A	3327	1/1	0.76	0.14	-	62,62,62,62	0
59	MG	2g	8001	1/1	0.89	0.09	-	74,74,74,74	0
59	MG	2A	3165	1/1	0.94	0.66	-	61,61,61,61	0
59	MG	1A	3835	1/1	0.67	0.16	-	51,51,51,51	0
59	MG	2A	3379	1/1	0.93	0.14	-	24,24,24,24	0
59	MG	1A	3319	1/1	0.77	0.17	-	38,38,38,38	0
59	MG	1a	3358	1/1	0.74	0.11	-	73,73,73,73	0
59	MG	25	104	1/1	0.84	0.40	-	61,61,61,61	0
59	MG	2A	3615	1/1	0.83	0.19	-	46,46,46,46	0
59	MG	1A	3615	1/1	0.95	0.15	-	55,55,55,55	0
59	MG	2a	3132	1/1	0.93	0.31	-	68,68,68,68	0
59	MG	2a	3080	1/1	0.91	0.15	-	58,58,58,58	0
59	MG	2a	3069	1/1	0.88	0.25	-	68,68,68,68	0
59	MG	1a	3373	1/1	0.82	0.14	-	59,59,59,59	0
59	MG	1A	3596	1/1	0.86	0.15	-	54,54,54,54	0
59	MG	1A	4222	1/1	0.98	0.52	-	31,31,31,31	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3104	1/1	0.95	0.53	-	32,32,32,32	0
59	MG	1A	3703	1/1	0.97	0.09	-	55,55,55,55	0
59	MG	1A	3032	1/1	0.95	0.48	-	26,26,26,26	0
59	MG	20	3003	1/1	0.97	0.12	-	50,50,50,50	0
59	MG	2A	3552	1/1	0.94	0.08	-	54,54,54,54	0
59	MG	1A	4067	1/1	0.95	0.33	-	36,36,36,36	0
59	MG	2A	3045	1/1	0.83	0.12	-	49,49,49,49	0
59	MG	1A	3662	1/1	0.97	0.15	-	15,15,15,15	0
59	MG	2A	3227	1/1	0.97	0.42	-	47,47,47,47	0
59	MG	1A	3641	1/1	0.88	0.13	-	40,40,40,40	0
59	MG	1A	4007	1/1	0.96	0.06	-	54,54,54,54	0
59	MG	2A	3251	1/1	0.94	0.22	-	40,40,40,40	0
59	MG	2A	3284	1/1	0.95	0.18	-	59,59,59,59	0
59	MG	1A	3992	1/1	0.93	0.08	-	30,30,30,30	0
59	MG	1a	3555	1/1	0.92	0.18	-	47,47,47,47	0
59	MG	1A	3229	1/1	0.96	0.15	-	45,45,45,45	0
59	MG	1A	3699	1/1	0.90	0.16	-	37,37,37,37	0
59	MG	1A	3850	1/1	0.83	0.12	-	46,46,46,46	0
59	MG	1A	3158	1/1	0.95	0.42	-	44,44,44,44	0
59	MG	2a	3179	1/1	0.86	0.18	-	64,64,64,64	0
59	MG	2A	3188	1/1	0.86	0.54	-	53,53,53,53	0
59	MG	2A	3204	1/1	0.93	0.08	-	58,58,58,58	0
59	MG	1a	3342	1/1	0.96	0.12	-	42,42,42,42	0
59	MG	1a	3443	1/1	0.94	0.23	-	49,49,49,49	0
59	MG	1X	3005	1/1	0.98	0.14	-	26,26,26,26	0
59	MG	1A	3943	1/1	0.95	0.15	-	33,33,33,33	0
59	MG	2A	3127	1/1	0.96	0.25	-	57,57,57,57	0
59	MG	1A	3476	1/1	0.94	0.33	-	44,44,44,44	0
59	MG	1A	3040	1/1	0.96	0.14	-	30,30,30,30	0
59	MG	1A	3525	1/1	0.95	0.17	-	48,48,48,48	0
59	MG	2a	3211	1/1	0.79	0.45	-	65,65,65,65	0
59	MG	1A	4040	1/1	0.98	0.09	-	36,36,36,36	0
59	MG	2w	3003	1/1	0.80	0.45	-	74,74,74,74	0
59	MG	1a	3459	1/1	0.85	0.17	-	66,66,66,66	0
59	MG	1A	3305	1/1	0.91	0.32	-	50,50,50,50	0
59	MG	2A	3427	1/1	0.88	0.16	-	49,49,49,49	0
59	MG	2A	3370	1/1	0.88	0.11	-	54,54,54,54	0
59	MG	2a	3050	1/1	0.96	0.14	-	54,54,54,54	0
59	MG	1a	3357	1/1	0.91	0.09	-	58,58,58,58	0
59	MG	1B	205	1/1	0.76	0.20	-	40,40,40,40	0
59	MG	2A	3548	1/1	0.94	0.13	-	36,36,36,36	0
59	MG	2A	3382	1/1	0.94	0.07	-	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3322	1/1	0.89	0.69	-	42,42,42,42	0
59	MG	2A	3328	1/1	0.86	0.20	-	51,51,51,51	0
59	MG	2a	3230	1/1	0.85	0.10	-	68,68,68,68	0
59	MG	1A	3127	1/1	0.92	0.51	-	37,37,37,37	0
59	MG	2a	3043	1/1	0.90	0.17	-	66,66,66,66	0
59	MG	1A	3625	1/1	0.96	0.09	-	63,63,63,63	0
59	MG	1a	3480	1/1	0.96	0.10	-	37,37,37,37	0
59	MG	2A	3133	1/1	0.94	0.26	-	48,48,48,48	0
59	MG	1A	3286	1/1	0.89	0.30	-	64,64,64,64	0
59	MG	1A	3997	1/1	0.29	0.17	-	41,41,41,41	0
59	MG	1A	4112	1/1	0.84	0.15	-	56,56,56,56	0
59	MG	1a	3566	1/1	0.91	0.08	-	57,57,57,57	0
59	MG	2a	3024	1/1	0.92	0.21	-	59,59,59,59	0
59	MG	1A	4008	1/1	0.95	0.08	-	54,54,54,54	0
59	MG	2A	3319	1/1	0.94	0.61	-	36,36,36,36	0
59	MG	2A	3719	1/1	0.81	0.09	-	64,64,64,64	0
59	MG	2B	3021	1/1	0.94	0.20	-	68,68,68,68	0
59	MG	1A	4164	1/1	0.91	0.28	-	39,39,39,39	0
59	MG	1A	3951	1/1	0.92	0.09	-	52,52,52,52	0
59	MG	1B	212	1/1	0.92	0.40	-	42,42,42,42	0
59	MG	1A	3240	1/1	0.87	0.33	-	31,31,31,31	0
59	MG	1A	3189	1/1	0.95	0.65	-	43,43,43,43	0
59	MG	2A	3046	1/1	0.92	0.14	-	58,58,58,58	0
59	MG	1a	3327	1/1	0.88	0.19	-	60,60,60,60	0
59	MG	2A	3070	1/1	0.97	0.23	-	42,42,42,42	0
59	MG	1A	3528	1/1	0.76	0.23	-	35,35,35,35	0
59	MG	1a	3330	1/1	0.94	0.16	-	58,58,58,58	0
59	MG	2A	3364	1/1	0.56	0.30	-	63,63,63,63	0
59	MG	1A	3914	1/1	0.92	0.13	-	21,21,21,21	0
59	MG	2A	3012	1/1	0.96	0.38	-	35,35,35,35	0
59	MG	1A	3339	1/1	0.88	0.37	-	45,45,45,45	0
59	MG	1A	3856	1/1	0.98	0.10	-	39,39,39,39	0
59	MG	2A	3365	1/1	0.93	0.57	-	56,56,56,56	0
59	MG	1y	103	1/1	0.90	0.20	-	88,88,88,88	0
59	MG	1a	3429	1/1	0.82	0.17	-	61,61,61,61	0
59	MG	1A	3331	1/1	0.83	0.29	-	41,41,41,41	0
59	MG	2A	3131	1/1	0.72	0.19	-	51,51,51,51	0
59	MG	1a	3430	1/1	0.96	0.11	-	69,69,69,69	0
59	MG	1A	4018	1/1	0.95	0.13	-	47,47,47,47	0
59	MG	2a	3031	1/1	0.95	0.08	-	57,57,57,57	0
59	MG	2A	3207	1/1	0.96	0.05	-	56,56,56,56	0
59	MG	2A	3640	1/1	0.86	0.11	-	49,49,49,49	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2a	3167	1/1	0.82	0.17	-	74,74,74,74	0
59	MG	1A	3068	1/1	0.90	0.10	-	53,53,53,53	0
59	MG	1A	3395	1/1	0.50	0.27	-	54,54,54,54	0
59	MG	2A	3442	1/1	0.97	0.18	-	24,24,24,24	0
59	MG	1A	3213	1/1	0.96	0.50	-	27,27,27,27	0
59	MG	2a	3229	1/1	0.78	0.16	-	60,60,60,60	0
59	MG	2A	3122	1/1	0.75	0.20	-	67,67,67,67	0
59	MG	2A	3617	1/1	0.91	0.21	-	40,40,40,40	0
59	MG	1a	3345	1/1	0.85	0.18	-	62,62,62,62	0
59	MG	1A	4042	1/1	0.97	0.12	-	42,42,42,42	0
59	MG	1A	3482	1/1	0.78	0.15	-	42,42,42,42	0
59	MG	2A	3422	1/1	0.92	0.09	-	67,67,67,67	0
59	MG	2A	3107	1/1	0.89	0.26	-	44,44,44,44	0
59	MG	1A	3055	1/1	0.82	0.10	-	50,50,50,50	0
59	MG	2A	3369	1/1	0.92	0.07	-	67,67,67,67	0
59	MG	2A	3526	1/1	0.95	0.13	-	54,54,54,54	0
59	MG	1a	3396	1/1	0.74	0.67	-	62,62,62,62	0
59	MG	1a	3447	1/1	0.91	0.16	-	51,51,51,51	0
59	MG	2A	3339	1/1	0.91	0.17	-	50,50,50,50	0
59	MG	1A	3264	1/1	0.81	0.24	-	56,56,56,56	0
59	MG	2A	3642	1/1	0.92	0.26	-	63,63,63,63	0
59	MG	1A	3200	1/1	0.96	0.33	-	37,37,37,37	0
59	MG	2a	3022	1/1	0.89	0.24	-	62,62,62,62	0
59	MG	1A	3897	1/1	0.98	0.14	-	38,38,38,38	0
59	MG	2A	3797	1/1	0.93	0.06	-	49,49,49,49	0
59	MG	1A	4187	1/1	0.91	0.19	-	47,47,47,47	0
59	MG	2A	3757	1/1	0.95	0.12	-	54,54,54,54	0
59	MG	2a	3213	1/1	0.96	0.10	-	66,66,66,66	0
59	MG	1A	3123	1/1	0.97	0.47	-	28,28,28,28	0
59	MG	1A	3631	1/1	0.97	0.25	-	16,16,16,16	0
59	MG	1a	3414	1/1	0.74	0.32	-	57,57,57,57	0
59	MG	1A	3944	1/1	0.94	0.13	-	39,39,39,39	0
59	MG	1a	3424	1/1	0.80	0.10	-	66,66,66,66	0
59	MG	2A	3248	1/1	0.85	0.19	-	52,52,52,52	0
59	MG	1A	3989	1/1	0.83	0.14	-	49,49,49,49	0
59	MG	27	3001	1/1	0.91	0.18	-	56,56,56,56	0
59	MG	1a	3425	1/1	0.89	0.30	-	58,58,58,58	0
59	MG	1A	3860	1/1	0.98	0.12	-	37,37,37,37	0
59	MG	1A	3318	1/1	0.82	0.27	-	49,49,49,49	0
59	MG	2A	3082	1/1	0.90	0.13	-	44,44,44,44	0
59	MG	2a	3010	1/1	0.93	0.19	-	62,62,62,62	0
59	MG	1A	3021	1/1	0.94	0.13	-	67,67,67,67	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3559	1/1	0.90	0.09	-	38,38,38,38	0
59	MG	1a	3529	1/1	0.93	0.07	-	53,53,53,53	0
59	MG	1a	3550	1/1	0.83	0.14	-	78,78,78,78	0
59	MG	2a	3193	1/1	0.89	0.15	-	70,70,70,70	0
59	MG	1A	4154	1/1	0.96	0.12	-	38,38,38,38	0
59	MG	2A	3124	1/1	0.93	0.22	-	32,32,32,32	0
59	MG	1A	3424	1/1	0.94	0.14	-	53,53,53,53	0
59	MG	2A	3016	1/1	0.97	0.10	-	21,21,21,21	0
59	MG	1A	3070	1/1	0.95	0.32	-	25,25,25,25	0
59	MG	1A	4078	1/1	0.91	0.09	-	58,58,58,58	0
59	MG	2A	3736	1/1	0.88	0.56	-	58,58,58,58	0
59	MG	2A	3262	1/1	0.94	0.13	-	56,56,56,56	0
59	MG	1A	3608	1/1	0.94	0.18	-	16,16,16,16	0
59	MG	1A	3458	1/1	0.85	0.18	-	38,38,38,38	0
59	MG	1A	3799	1/1	0.96	0.21	-	22,22,22,22	0
59	MG	2A	3542	1/1	0.66	0.10	-	52,52,52,52	0
59	MG	1A	3297	1/1	0.97	0.41	-	43,43,43,43	0
59	MG	1A	4138	1/1	0.89	0.12	-	76,76,76,76	0
59	MG	2a	3089	1/1	0.99	0.15	-	67,67,67,67	0
59	MG	1A	3491	1/1	0.73	0.29	-	47,47,47,47	0
59	MG	2a	3100	1/1	0.82	0.20	-	51,51,51,51	0
59	MG	2A	3031	1/1	0.89	0.17	-	44,44,44,44	0
59	MG	1A	3924	1/1	0.94	0.20	-	22,22,22,22	0
59	MG	1A	3836	1/1	0.94	0.20	-	36,36,36,36	0
59	MG	2A	3085	1/1	0.85	0.12	-	62,62,62,62	0
59	MG	1B	238	1/1	0.83	0.15	-	30,30,30,30	0
59	MG	1a	3379	1/1	0.95	0.06	-	49,49,49,49	0
59	MG	1a	3329	1/1	0.90	0.12	-	64,64,64,64	0
59	MG	1a	3305	1/1	0.87	0.14	-	54,54,54,54	0
59	MG	2A	3079	1/1	0.93	0.13	-	60,60,60,60	0
59	MG	2A	3412	1/1	0.70	0.16	-	40,40,40,40	0
59	MG	1A	3477	1/1	0.94	0.15	-	31,31,31,31	0
59	MG	1A	3404	1/1	0.91	0.34	-	34,34,34,34	0
59	MG	1a	3482	1/1	0.45	0.27	-	52,52,52,52	0
59	MG	2A	3487	1/1	0.97	0.19	-	48,48,48,48	0
59	MG	1a	3451	1/1	0.96	0.16	-	43,43,43,43	0
59	MG	1A	3851	1/1	0.93	0.17	-	43,43,43,43	0
59	MG	2A	3415	1/1	0.87	0.14	-	36,36,36,36	0
59	MG	2a	3057	1/1	0.89	0.09	-	60,60,60,60	0
59	MG	1A	3119	1/1	0.93	0.22	-	33,33,33,33	0
59	MG	23	101	1/1	0.91	0.20	-	51,51,51,51	0
59	MG	1a	3427	1/1	0.94	0.14	-	64,64,64,64	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3593	1/1	0.89	0.15	-	18,18,18,18	0
59	MG	1a	3528	1/1	0.91	0.17	-	59,59,59,59	0
59	MG	1A	3958	1/1	0.74	0.23	-	45,45,45,45	0
59	MG	2a	3164	1/1	0.86	0.16	-	75,75,75,75	0
59	MG	2A	3798	1/1	0.86	0.13	-	57,57,57,57	0
59	MG	1A	4030	1/1	0.85	0.11	-	58,58,58,58	0
59	MG	10	103	1/1	0.97	0.24	-	35,35,35,35	0
59	MG	1A	3519	1/1	0.78	0.34	-	46,46,46,46	0
59	MG	1A	3266	1/1	0.95	0.22	-	45,45,45,45	0
59	MG	2A	3349	1/1	0.74	0.21	-	50,50,50,50	0
59	MG	1A	3793	1/1	0.96	0.08	-	46,46,46,46	0
59	MG	2A	3573	1/1	0.82	0.13	-	64,64,64,64	0
59	MG	2A	3787	1/1	0.89	0.13	-	58,58,58,58	0
59	MG	1A	3462	1/1	0.79	0.14	-	50,50,50,50	0
59	MG	1A	4151	1/1	0.96	0.09	-	45,45,45,45	0
59	MG	1A	3100	1/1	0.95	0.49	-	23,23,23,23	0
59	MG	1A	3019	1/1	0.94	0.39	-	33,33,33,33	0
59	MG	1A	3271	1/1	0.78	0.76	-	35,35,35,35	0
59	MG	1a	3522	1/1	0.93	0.12	-	66,66,66,66	0
59	MG	1A	3196	1/1	0.97	0.49	-	26,26,26,26	0
59	MG	2A	3562	1/1	0.87	0.11	-	57,57,57,57	0
59	MG	1A	3402	1/1	0.83	0.31	-	42,42,42,42	0
59	MG	2a	3041	1/1	0.90	0.20	-	63,63,63,63	0
59	MG	2A	3651	1/1	0.59	0.25	-	72,72,72,72	0
59	MG	2A	3260	1/1	0.88	0.21	-	76,76,76,76	0
59	MG	1a	3554	1/1	0.97	0.07	-	58,58,58,58	0
59	MG	1N	3008	1/1	0.95	0.18	-	36,36,36,36	0
59	MG	1a	3448	1/1	0.86	0.18	-	60,60,60,60	0
59	MG	1A	4161	1/1	0.92	0.14	-	35,35,35,35	0
59	MG	25	103	1/1	0.84	0.38	-	42,42,42,42	0
59	MG	2a	3214	1/1	0.86	0.23	-	68,68,68,68	0
59	MG	2A	3425	1/1	0.95	0.12	-	47,47,47,47	0
59	MG	1A	3623	1/1	0.85	0.16	-	15,15,15,15	0
59	MG	1Z	304	1/1	0.95	0.23	-	50,50,50,50	0
59	MG	2A	3234	1/1	0.93	0.17	-	64,64,64,64	0
59	MG	1A	3891	1/1	0.90	0.09	-	15,15,15,15	0
59	MG	1A	3474	1/1	0.83	0.16	-	55,55,55,55	0
59	MG	2A	3334	1/1	0.89	0.13	-	61,61,61,61	0
59	MG	1A	3350	1/1	0.90	0.21	-	57,57,57,57	0
59	MG	2A	3794	1/1	0.95	0.08	-	61,61,61,61	0
59	MG	2w	3002	1/1	0.87	0.10	-	73,73,73,73	0
59	MG	2A	3601	1/1	0.77	0.15	-	60,60,60,60	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3832	1/1	0.98	0.22	-	53,53,53,53	0
59	MG	1a	3571	1/1	0.81	0.11	-	57,57,57,57	0
59	MG	1a	3497	1/1	0.96	0.08	-	65,65,65,65	0
59	MG	1A	3763	1/1	0.97	0.17	-	46,46,46,46	0
59	MG	2A	3659	1/1	0.79	0.09	-	66,66,66,66	0
59	MG	1A	4191	1/1	0.93	0.11	-	27,27,27,27	0
59	MG	2a	3030	1/1	0.95	0.12	-	60,60,60,60	0
59	MG	1A	3461	1/1	0.91	0.11	-	58,58,58,58	0
59	MG	2A	3324	1/1	0.70	0.21	-	53,53,53,53	0
59	MG	1A	3358	1/1	0.84	0.91	-	49,49,49,49	0
59	MG	1A	4046	1/1	0.91	0.19	-	36,36,36,36	0
59	MG	2a	3115	1/1	0.89	0.20	-	59,59,59,59	0
59	MG	1A	3385	1/1	0.90	0.48	-	39,39,39,39	0
59	MG	1A	3972	1/1	0.95	0.16	-	24,24,24,24	0
59	MG	2A	3271	1/1	0.66	0.20	-	71,71,71,71	0
59	MG	2A	3546	1/1	0.89	0.14	-	60,60,60,60	0
59	MG	1A	3300	1/1	0.86	0.11	-	42,42,42,42	0
59	MG	1A	3250	1/1	0.95	0.13	-	49,49,49,49	0
59	MG	2A	3682	1/1	0.87	0.13	-	56,56,56,56	0
59	MG	2R	203	1/1	0.87	0.19	-	52,52,52,52	0
59	MG	1A	3996	1/1	0.81	0.11	-	35,35,35,35	0
59	MG	2a	3064	1/1	0.74	0.12	-	70,70,70,70	0
59	MG	2a	3149	1/1	0.86	0.12	-	58,58,58,58	0
59	MG	1l	202	1/1	0.61	0.26	-	82,82,82,82	0
59	MG	2A	3775	1/1	0.68	0.17	-	63,63,63,63	0
59	MG	1A	3682	1/1	0.95	0.17	-	48,48,48,48	0
59	MG	1A	3336	1/1	0.78	0.24	-	36,36,36,36	0
59	MG	1a	3452	1/1	0.96	0.11	-	40,40,40,40	0
59	MG	2v	102	1/1	0.85	0.17	-	69,69,69,69	0
59	MG	2A	3491	1/1	0.86	0.13	-	53,53,53,53	0
59	MG	10	107	1/1	0.94	0.14	-	38,38,38,38	0
59	MG	2A	3611	1/1	0.69	0.14	-	66,66,66,66	0
59	MG	1A	3004	1/1	0.93	0.36	-	47,47,47,47	0
59	MG	2a	3232	1/1	0.95	0.14	-	60,60,60,60	0
59	MG	1A	3465	1/1	0.98	0.36	-	27,27,27,27	0
59	MG	1A	3194	1/1	0.84	0.39	-	45,45,45,45	0
59	MG	2a	3171	1/1	0.82	0.12	-	55,55,55,55	0
59	MG	1A	3349	1/1	0.89	0.26	-	49,49,49,49	0
59	MG	2A	3621	1/1	0.92	0.07	-	57,57,57,57	0
59	MG	1A	3587	1/1	0.97	0.11	-	51,51,51,51	0
59	MG	2A	3630	1/1	0.94	0.10	-	66,66,66,66	0
59	MG	1A	3063	1/1	0.97	0.13	-	15,15,15,15	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1a	3303	1/1	0.89	0.08	-	55,55,55,55	0
59	MG	1A	3865	1/1	0.94	0.11	-	42,42,42,42	0
59	MG	1a	3399	1/1	0.94	0.15	-	42,42,42,42	0
59	MG	1A	4096	1/1	0.94	0.56	-	26,26,26,26	0
59	MG	1A	3203	1/1	0.90	0.19	-	53,53,53,53	0
59	MG	2A	3811	1/1	0.98	0.33	-	62,62,62,62	0
59	MG	2a	3045	1/1	0.80	0.23	-	58,58,58,58	0
59	MG	1A	3666	1/1	0.93	0.28	-	49,49,49,49	0
59	MG	2B	3012	1/1	0.98	0.14	-	63,63,63,63	0
59	MG	1a	3392	1/1	0.98	0.27	-	45,45,45,45	0
59	MG	2A	3145	1/1	0.80	0.36	-	62,62,62,62	0
59	MG	1x	111	1/1	0.94	0.07	-	65,65,65,65	0
59	MG	2y	3001	1/1	0.91	0.14	-	65,65,65,65	0
59	MG	2a	3055	1/1	0.81	0.19	-	58,58,58,58	0
59	MG	2A	3570	1/1	0.86	0.10	-	64,64,64,64	0
59	MG	1A	3787	1/1	0.88	0.14	-	49,49,49,49	0
59	MG	2A	3560	1/1	0.86	0.11	-	61,61,61,61	0
59	MG	1a	3382	1/1	0.85	0.19	-	63,63,63,63	0
59	MG	1A	3508	1/1	0.91	0.12	-	53,53,53,53	0
59	MG	1A	3348	1/1	0.95	0.24	-	56,56,56,56	0
59	MG	2y	3002	1/1	0.88	0.17	-	80,80,80,80	0
59	MG	2A	3739	1/1	0.90	0.11	-	49,49,49,49	0
59	MG	1A	3652	1/1	0.91	0.19	-	35,35,35,35	0
59	MG	2A	3088	1/1	0.83	0.20	-	65,65,65,65	0
59	MG	1a	3339	1/1	0.93	0.13	-	54,54,54,54	0
59	MG	2a	3086	1/1	0.94	0.19	-	60,60,60,60	0
59	MG	1A	3355	1/1	0.96	0.42	-	30,30,30,30	0
59	MG	1A	3961	1/1	0.82	0.11	-	37,37,37,37	0
59	MG	2A	3623	1/1	0.95	0.07	-	51,51,51,51	0
59	MG	2A	3758	1/1	0.95	0.20	-	63,63,63,63	0
59	MG	1A	3673	1/1	0.86	0.23	-	33,33,33,33	0
59	MG	1A	3470	1/1	0.86	0.28	-	47,47,47,47	0
59	MG	1A	3094	1/1	0.95	0.54	-	28,28,28,28	0
59	MG	2A	3291	1/1	0.81	0.14	-	74,74,74,74	0
59	MG	1a	3499	1/1	0.97	0.15	-	61,61,61,61	0
59	MG	2A	3518	1/1	0.96	0.14	-	51,51,51,51	0
59	MG	1B	232	1/1	0.70	0.13	-	67,67,67,67	0
59	MG	1A	3141	1/1	0.98	0.19	-	26,26,26,26	0
59	MG	2A	3267	1/1	0.95	0.22	-	55,55,55,55	0
59	MG	1a	3401	1/1	0.93	0.13	-	48,48,48,48	0
59	MG	1A	3249	1/1	0.82	0.21	-	43,43,43,43	0
59	MG	1A	3310	1/1	0.78	0.75	-	40,40,40,40	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3021	1/1	0.79	0.14	-	54,54,54,54	0
59	MG	2A	3111	1/1	0.78	0.34	-	60,60,60,60	0
59	MG	1A	3394	1/1	0.75	0.64	-	60,60,60,60	0
59	MG	1F	306	1/1	0.93	0.19	-	40,40,40,40	0
59	MG	2A	3029	1/1	0.88	0.17	-	66,66,66,66	0
59	MG	2a	3125	1/1	0.88	0.32	-	79,79,79,79	0
59	MG	1A	3722	1/1	0.78	0.15	-	72,72,72,72	0
59	MG	1A	4061	1/1	0.93	0.13	-	40,40,40,40	0
59	MG	2A	3205	1/1	0.86	0.16	-	73,73,73,73	0
59	MG	2A	3213	1/1	0.90	0.17	-	52,52,52,52	0
59	MG	2A	3474	1/1	0.94	0.15	-	55,55,55,55	0
59	MG	1a	3465	1/1	0.94	0.10	-	26,26,26,26	0
59	MG	1A	4053	1/1	0.95	0.15	-	69,69,69,69	0
59	MG	1A	3812	1/1	0.82	0.13	-	57,57,57,57	0
59	MG	1A	3882	1/1	0.87	0.05	-	45,45,45,45	0
59	MG	1l	103	1/1	0.84	0.14	-	70,70,70,70	0
59	MG	1A	3879	1/1	0.85	0.15	-	51,51,51,51	0
59	MG	2A	3769	1/1	0.88	0.16	-	56,56,56,56	0
59	MG	2a	3138	1/1	0.85	0.10	-	88,88,88,88	0
59	MG	1A	3077	1/1	0.87	0.43	-	42,42,42,42	0
59	MG	2A	3692	1/1	0.84	0.12	-	40,40,40,40	0
59	MG	2A	3018	1/1	0.54	0.18	-	71,71,71,71	0
59	MG	2A	3063	1/1	0.95	0.18	-	40,40,40,40	0
59	MG	2A	3095	1/1	0.79	0.16	-	60,60,60,60	0
59	MG	1A	3513	1/1	0.89	0.18	-	44,44,44,44	0
59	MG	1w	109	1/1	0.74	0.10	-	68,68,68,68	0
59	MG	2A	3242	1/1	0.85	0.17	-	56,56,56,56	0
59	MG	2A	3671	1/1	0.93	0.17	-	66,66,66,66	0
59	MG	1A	3121	1/1	0.93	0.13	-	37,37,37,37	0
59	MG	2A	3782	1/1	0.83	0.09	-	60,60,60,60	0
59	MG	2v	101	1/1	0.93	0.19	-	49,49,49,49	0
59	MG	2a	3109	1/1	0.88	0.17	-	55,55,55,55	0
59	MG	1F	302	1/1	0.79	0.23	-	50,50,50,50	0
59	MG	1A	3553	1/1	0.94	0.17	-	20,20,20,20	0
59	MG	2A	3203	1/1	0.96	0.78	-	61,61,61,61	0
59	MG	1A	3759	1/1	0.93	0.12	-	41,41,41,41	0
59	MG	1A	3219	1/1	0.93	0.30	-	51,51,51,51	0
59	MG	1A	4131	1/1	0.84	0.23	-	40,40,40,40	0
59	MG	2A	3142	1/1	0.94	0.11	-	58,58,58,58	0
59	MG	1A	3490	1/1	0.98	0.26	-	58,58,58,58	0
59	MG	1a	3355	1/1	0.83	0.17	-	51,51,51,51	0
59	MG	1A	4072	1/1	0.79	0.13	-	55,55,55,55	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3502	1/1	0.98	0.16	-	42,42,42,42	0
59	MG	1A	3359	1/1	0.85	0.33	-	56,56,56,56	0
59	MG	1A	3764	1/1	0.91	0.12	-	54,54,54,54	0
59	MG	2A	3553	1/1	0.96	0.10	-	63,63,63,63	0
59	MG	1A	3679	1/1	0.90	0.11	-	50,50,50,50	0
59	MG	1A	3792	1/1	0.94	0.20	-	42,42,42,42	0
59	MG	1A	3766	1/1	0.95	0.12	-	21,21,21,21	0
59	MG	2A	3191	1/1	0.68	0.63	-	59,59,59,59	0
59	MG	2A	3293	1/1	0.74	0.62	-	55,55,55,55	0
59	MG	2A	3592	1/1	0.97	0.12	-	23,23,23,23	0
59	MG	1a	3541	1/1	0.94	0.06	-	61,61,61,61	0
59	MG	2A	3539	1/1	0.86	0.27	-	57,57,57,57	0
59	MG	1B	231	1/1	0.88	0.15	-	63,63,63,63	0
59	MG	2a	3002	1/1	0.83	0.13	-	73,73,73,73	0
59	MG	2A	3119	1/1	0.91	0.13	-	61,61,61,61	0
59	MG	1A	3492	1/1	0.87	0.22	-	55,55,55,55	0
59	MG	1A	3191	1/1	0.89	0.38	-	45,45,45,45	0
59	MG	2B	3019	1/1	0.90	0.10	-	61,61,61,61	0
59	MG	2A	3007	1/1	0.97	0.15	-	40,40,40,40	0
59	MG	1A	3405	1/1	0.63	0.17	-	55,55,55,55	0
59	MG	1w	112	1/1	0.77	0.23	-	65,65,65,65	0
59	MG	1a	3320	1/1	0.94	0.12	-	51,51,51,51	0
59	MG	1A	4060	1/1	0.90	0.25	-	27,27,27,27	0
59	MG	2A	3233	1/1	0.84	0.15	-	66,66,66,66	0
59	MG	1A	3062	1/1	0.95	0.17	-	47,47,47,47	0
59	MG	1A	4052	1/1	0.89	0.11	-	60,60,60,60	0
59	MG	2A	3498	1/1	0.91	0.13	-	45,45,45,45	0
59	MG	1A	3524	1/1	0.93	0.18	-	31,31,31,31	0
59	MG	1A	3613	1/1	0.96	0.14	-	36,36,36,36	0
59	MG	1A	3670	1/1	0.94	0.20	-	43,43,43,43	0
59	MG	2A	3398	1/1	0.93	0.20	-	50,50,50,50	0
59	MG	2A	3373	1/1	0.95	0.12	-	31,31,31,31	0
59	MG	2A	3723	1/1	0.96	0.07	-	47,47,47,47	0
59	MG	1A	3919	1/1	0.93	0.05	-	67,67,67,67	0
59	MG	2a	3111	1/1	0.89	0.13	-	54,54,54,54	0
59	MG	1O	3003	1/1	0.83	0.11	-	54,54,54,54	0
59	MG	2A	3244	1/1	0.91	0.74	-	47,47,47,47	0
59	MG	2a	3126	1/1	0.86	0.12	-	59,59,59,59	0
59	MG	1A	3422	1/1	0.75	0.17	-	56,56,56,56	0
59	MG	1a	3484	1/1	0.90	0.16	-	41,41,41,41	0
59	MG	2A	3337	1/1	0.91	0.12	-	44,44,44,44	0
59	MG	1A	3303	1/1	0.82	0.38	-	50,50,50,50	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	4136	1/1	0.96	0.09	-	50,50,50,50	0
59	MG	2a	3148	1/1	0.95	0.07	-	73,73,73,73	0
59	MG	2A	3576	1/1	0.90	0.19	-	64,64,64,64	0
59	MG	2A	3170	1/1	0.84	0.23	-	60,60,60,60	0
59	MG	2A	3098	1/1	0.88	0.20	-	47,47,47,47	0
59	MG	1A	3274	1/1	0.87	0.15	-	51,51,51,51	0
59	MG	1a	3365	1/1	0.81	0.18	-	62,62,62,62	0
59	MG	1A	3867	1/1	0.84	0.15	-	36,36,36,36	0
59	MG	1A	3457	1/1	0.87	0.25	-	55,55,55,55	0
59	MG	2A	3123	1/1	0.88	0.18	-	46,46,46,46	0
59	MG	2A	3172	1/1	0.88	0.27	-	54,54,54,54	0
59	MG	1A	3049	1/1	0.84	0.18	-	54,54,54,54	0
59	MG	1a	3464	1/1	0.97	0.10	-	31,31,31,31	0
59	MG	2A	3679	1/1	0.82	0.18	-	79,79,79,79	0
59	MG	2B	3022	1/1	0.73	0.14	-	72,72,72,72	0
59	MG	1A	3215	1/1	0.95	0.69	-	33,33,33,33	0
59	MG	2A	3335	1/1	0.89	0.35	-	56,56,56,56	0
59	MG	1A	4093	1/1	0.96	0.13	-	40,40,40,40	0
59	MG	2A	3389	1/1	0.95	0.12	-	33,33,33,33	0
59	MG	2A	3660	1/1	0.94	0.17	-	76,76,76,76	0
59	MG	1A	3908	1/1	0.90	0.13	-	55,55,55,55	0
59	MG	2A	3666	1/1	0.94	0.21	-	62,62,62,62	0
59	MG	2A	3243	1/1	0.91	0.13	-	66,66,66,66	0
59	MG	1A	4051	1/1	0.93	0.07	-	38,38,38,38	0
59	MG	1A	3778	1/1	0.86	0.15	-	38,38,38,38	0
59	MG	1A	3707	1/1	0.88	0.09	-	57,57,57,57	0
59	MG	1B	233	1/1	0.79	0.34	-	64,64,64,64	0
59	MG	1A	3365	1/1	0.96	0.22	-	40,40,40,40	0
59	MG	2A	3307	1/1	0.87	0.14	-	60,60,60,60	0
59	MG	2A	3139	1/1	0.93	0.15	-	33,33,33,33	0
59	MG	1a	3516	1/1	0.94	0.20	-	64,64,64,64	0
59	MG	1A	3388	1/1	0.91	0.09	-	47,47,47,47	0
59	MG	1A	3925	1/1	0.96	0.12	-	10,10,10,10	0
59	MG	2a	3027	1/1	0.85	0.32	-	63,63,63,63	0
59	MG	1E	303	1/1	0.92	0.29	-	33,33,33,33	0
59	MG	1A	3988	1/1	0.70	0.28	-	62,62,62,62	0
59	MG	2A	3628	1/1	0.79	0.11	-	43,43,43,43	0
59	MG	1A	3252	1/1	0.80	0.14	-	28,28,28,28	0
59	MG	2A	3256	1/1	0.90	0.74	-	53,53,53,53	0
59	MG	2a	3234	1/1	0.90	0.10	-	62,62,62,62	0
59	MG	1A	3705	1/1	0.96	0.16	-	17,17,17,17	0
59	MG	1A	4140	1/1	0.93	0.11	-	31,31,31,31	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3766	1/1	0.89	0.17	-	53,53,53,53	0
59	MG	2A	3418	1/1	0.88	0.10	-	46,46,46,46	0
59	MG	2a	3105	1/1	0.96	0.12	-	64,64,64,64	0
59	MG	1A	3917	1/1	0.91	0.20	-	15,15,15,15	0
59	MG	1A	3738	1/1	0.97	0.11	-	27,27,27,27	0
59	MG	2A	3110	1/1	0.78	0.15	-	60,60,60,60	0
59	MG	1A	3982	1/1	0.92	0.18	-	57,57,57,57	0
59	MG	1A	3471	1/1	0.92	0.23	-	30,30,30,30	0
59	MG	2A	3551	1/1	0.86	0.27	-	60,60,60,60	0
59	MG	1A	3839	1/1	0.96	0.46	-	25,25,25,25	0
59	MG	1A	3767	1/1	0.87	0.18	-	53,53,53,53	0
59	MG	1A	3075	1/1	0.89	0.69	-	29,29,29,29	0
59	MG	2A	3186	1/1	0.86	0.09	-	64,64,64,64	0
59	MG	2A	3189	1/1	0.88	0.20	-	54,54,54,54	0
59	MG	1Y	503	1/1	0.86	0.22	-	69,69,69,69	0
59	MG	2a	3090	1/1	0.95	0.08	-	65,65,65,65	0
59	MG	1A	3132	1/1	0.89	0.26	-	30,30,30,30	0
59	MG	1A	3415	1/1	0.79	0.57	-	45,45,45,45	0
59	MG	1a	3423	1/1	0.91	0.18	-	53,53,53,53	0
59	MG	1A	3912	1/1	0.94	0.08	-	47,47,47,47	0
59	MG	1A	3655	1/1	0.83	0.13	-	42,42,42,42	0
59	MG	2a	3236	1/1	0.93	0.11	-	75,75,75,75	0
59	MG	1A	3344	1/1	0.85	0.17	-	57,57,57,57	0
59	MG	1A	3543	1/1	0.93	0.17	-	17,17,17,17	0
59	MG	2a	3001	1/1	0.89	0.12	-	52,52,52,52	0
59	MG	2A	3781	1/1	0.46	0.29	-	63,63,63,63	0
59	MG	2A	3528	1/1	0.69	0.25	-	81,81,81,81	0
59	MG	2A	3362	1/1	0.83	0.20	-	57,57,57,57	0
59	MG	1A	3932	1/1	0.89	0.07	-	49,49,49,49	0
59	MG	1a	3479	1/1	0.92	0.12	-	48,48,48,48	0
59	MG	2A	3609	1/1	0.96	0.07	-	44,44,44,44	0
59	MG	1W	202	1/1	0.91	0.15	-	39,39,39,39	0
59	MG	2D	302	1/1	0.93	0.12	-	39,39,39,39	0
59	MG	2A	3296	1/1	0.95	0.22	-	33,33,33,33	0
59	MG	2a	3004	1/1	0.93	0.18	-	56,56,56,56	0
59	MG	2A	3543	1/1	0.88	0.24	-	46,46,46,46	0
59	MG	1a	3354	1/1	0.92	0.10	-	66,66,66,66	0
59	MG	2A	3657	1/1	0.82	0.10	-	60,60,60,60	0
59	MG	1A	3582	1/1	0.96	0.13	-	24,24,24,24	0
59	MG	2B	3006	1/1	0.96	0.17	-	66,66,66,66	0
59	MG	1A	3347	1/1	0.90	0.51	-	40,40,40,40	0
59	MG	1A	3874	1/1	0.91	0.13	-	13,13,13,13	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2a	3184	1/1	0.94	0.18	-	61,61,61,61	0
59	MG	2A	3090	1/1	0.97	0.08	-	53,53,53,53	0
59	MG	1A	3436	1/1	0.86	0.82	-	34,34,34,34	0
59	MG	1A	3818	1/1	0.94	0.15	-	14,14,14,14	0
59	MG	1A	4081	1/1	0.96	0.07	-	14,14,14,14	0
59	MG	1A	3146	1/1	0.96	0.35	-	34,34,34,34	0
59	MG	1A	3994	1/1	0.94	0.08	-	31,31,31,31	0
59	MG	1A	3452	1/1	0.86	0.22	-	32,32,32,32	0
59	MG	2A	3179	1/1	0.87	0.15	-	46,46,46,46	0
59	MG	1A	3533	1/1	0.91	0.08	-	51,51,51,51	0
59	MG	1B	206	1/1	0.89	0.19	-	50,50,50,50	0
59	MG	1A	3281	1/1	0.96	0.27	-	43,43,43,43	0
59	MG	1A	3572	1/1	0.95	0.15	-	38,38,38,38	0
59	MG	1A	3226	1/1	0.83	0.20	-	56,56,56,56	0
59	MG	1B	230	1/1	0.83	0.16	-	59,59,59,59	0
59	MG	2A	3351	1/1	0.84	0.25	-	56,56,56,56	0
59	MG	2A	3515	1/1	0.86	0.14	-	39,39,39,39	0
59	MG	1E	308	1/1	0.81	0.20	-	59,59,59,59	0
59	MG	1a	3544	1/1	0.83	0.17	-	66,66,66,66	0
59	MG	1a	3322	1/1	0.94	0.27	-	51,51,51,51	0
59	MG	1A	4104	1/1	0.90	0.58	-	53,53,53,53	0
59	MG	1A	3039	1/1	0.91	0.08	-	22,22,22,22	0
59	MG	1A	3700	1/1	0.91	0.14	-	20,20,20,20	0
59	MG	2A	3508	1/1	0.91	0.15	-	43,43,43,43	0
59	MG	2a	3046	1/1	0.78	0.11	-	70,70,70,70	0
59	MG	1A	3188	1/1	0.89	0.42	-	37,37,37,37	0
59	MG	1A	4125	1/1	0.89	0.31	-	40,40,40,40	0
59	MG	2a	3183	1/1	0.85	0.12	-	65,65,65,65	0
59	MG	1A	3807	1/1	0.96	0.25	-	21,21,21,21	0
59	MG	1b	3002	1/1	0.94	0.16	-	69,69,69,69	0
59	MG	1R	201	1/1	0.96	0.16	-	28,28,28,28	0
59	MG	2A	3673	1/1	0.89	0.23	-	55,55,55,55	0
59	MG	1a	3341	1/1	0.90	0.11	-	50,50,50,50	0
59	MG	1A	3617	1/1	0.87	0.27	-	47,47,47,47	0
59	MG	1A	3126	1/1	0.87	0.57	-	29,29,29,29	0
59	MG	2A	3157	1/1	0.97	0.10	-	47,47,47,47	0
59	MG	2a	3104	1/1	0.74	0.19	-	54,54,54,54	0
59	MG	1Y	502	1/1	0.95	0.15	-	53,53,53,53	0
59	MG	2A	3305	1/1	0.86	0.20	-	52,52,52,52	0
59	MG	1A	3210	1/1	0.94	0.29	-	35,35,35,35	0
59	MG	1A	3517	1/1	0.86	0.23	-	41,41,41,41	0
59	MG	2A	3741	1/1	0.94	0.09	-	60,60,60,60	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3366	1/1	0.86	0.34	-	58,58,58,58	0
59	MG	2A	3037	1/1	0.80	0.15	-	59,59,59,59	0
59	MG	2A	3507	1/1	0.86	0.16	-	33,33,33,33	0
59	MG	2A	3222	1/1	0.95	0.13	-	45,45,45,45	0
59	MG	1A	4192	1/1	0.97	0.49	-	29,29,29,29	0
59	MG	1A	3083	1/1	0.94	0.19	-	36,36,36,36	0
59	MG	2A	3575	1/1	0.95	0.10	-	60,60,60,60	0
59	MG	1A	3298	1/1	0.79	0.20	-	57,57,57,57	0
59	MG	1A	4027	1/1	0.93	0.05	-	45,45,45,45	0
59	MG	1A	3193	1/1	0.72	0.51	-	56,56,56,56	0
59	MG	1A	3092	1/1	0.95	0.47	-	32,32,32,32	0
59	MG	2a	3107	1/1	0.94	0.19	-	68,68,68,68	0
59	MG	2F	301	1/1	0.88	0.21	-	70,70,70,70	0
59	MG	1A	3858	1/1	0.99	0.15	-	22,22,22,22	0
59	MG	1A	3247	1/1	0.94	0.45	-	47,47,47,47	0
59	MG	1E	305	1/1	0.92	0.32	-	42,42,42,42	0
59	MG	1Q	204	1/1	0.77	0.19	-	46,46,46,46	0
59	MG	2A	3342	1/1	0.97	0.14	-	41,41,41,41	0
59	MG	1w	106	1/1	0.98	0.13	-	61,61,61,61	0
59	MG	1A	4100	1/1	0.94	0.32	-	50,50,50,50	0
59	MG	1A	4082	1/1	0.86	0.12	-	50,50,50,50	0
59	MG	2a	3021	1/1	0.82	0.21	-	62,62,62,62	0
59	MG	2A	3810	1/1	0.90	0.44	-	59,59,59,59	0
59	MG	2A	3253	1/1	0.94	0.56	-	61,61,61,61	0
59	MG	2A	3314	1/1	0.67	0.17	-	60,60,60,60	0
59	MG	1A	3888	1/1	0.93	0.11	-	31,31,31,31	0
59	MG	1a	3533	1/1	0.98	0.05	-	59,59,59,59	0
59	MG	1A	3460	1/1	0.92	0.13	-	50,50,50,50	0
59	MG	1A	4001	1/1	0.98	0.23	-	37,37,37,37	0
59	MG	1a	3404	1/1	0.88	0.15	-	65,65,65,65	0
59	MG	1A	4035	1/1	0.97	0.17	-	35,35,35,35	0
59	MG	2a	3221	1/1	0.81	0.26	-	76,76,76,76	0
59	MG	1A	3819	1/1	0.80	0.22	-	23,23,23,23	0
59	MG	2A	3812	1/1	0.92	0.08	-	72,72,72,72	0
59	MG	1A	3154	1/1	0.97	0.15	-	42,42,42,42	0
59	MG	1A	3393	1/1	0.93	0.60	-	47,47,47,47	0
59	MG	2A	3388	1/1	0.95	0.17	-	60,60,60,60	0
59	MG	1E	309	1/1	0.93	0.17	-	47,47,47,47	0
59	MG	1a	3455	1/1	0.91	0.11	-	27,27,27,27	0
59	MG	2a	3084	1/1	0.78	0.17	-	71,71,71,71	0
59	MG	1a	3563	1/1	0.97	0.18	-	50,50,50,50	0
59	MG	1Z	303	1/1	0.72	0.35	-	69,69,69,69	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3003	1/1	0.99	0.18	-	16,16,16,16	0
59	MG	1A	3438	1/1	0.74	0.22	-	57,57,57,57	0
59	MG	1A	4004	1/1	0.97	0.10	-	43,43,43,43	0
59	MG	1A	3107	1/1	0.95	0.77	-	32,32,32,32	0
59	MG	1A	4116	1/1	0.90	0.15	-	50,50,50,50	0
59	MG	2A	3734	1/1	0.85	0.16	-	54,54,54,54	0
59	MG	1A	3441	1/1	0.76	0.33	-	62,62,62,62	0
59	MG	1A	3052	1/1	0.97	0.15	-	45,45,45,45	0
59	MG	1A	3330	1/1	0.84	0.79	-	36,36,36,36	0
59	MG	2A	3596	1/1	0.79	0.34	-	60,60,60,60	0
59	MG	2a	3169	1/1	0.95	0.13	-	51,51,51,51	0
59	MG	1A	4163	1/1	0.89	0.22	-	53,53,53,53	0
59	MG	2A	3680	1/1	0.90	0.19	-	55,55,55,55	0
59	MG	2A	3738	1/1	0.94	0.11	-	60,60,60,60	0
59	MG	2A	3152	1/1	0.80	0.23	-	48,48,48,48	0
59	MG	1A	3224	1/1	0.97	0.13	-	49,49,49,49	0
59	MG	1A	3015	1/1	0.89	0.24	-	43,43,43,43	0
59	MG	2a	3047	1/1	0.87	0.17	-	60,60,60,60	0
59	MG	2A	3755	1/1	0.95	0.12	-	64,64,64,64	0
59	MG	2A	3505	1/1	0.93	0.20	-	35,35,35,35	0
59	MG	2A	3175	1/1	0.74	0.28	-	63,63,63,63	0
59	MG	2A	3198	1/1	0.91	0.20	-	71,71,71,71	0
59	MG	1a	3491	1/1	0.92	0.17	-	75,75,75,75	0
59	MG	2A	3229	1/1	0.83	0.16	-	54,54,54,54	0
59	MG	2A	3476	1/1	0.83	0.12	-	43,43,43,43	0
59	MG	1A	3956	1/1	0.97	0.09	-	26,26,26,26	0
59	MG	2A	3574	1/1	0.93	0.12	-	56,56,56,56	0
59	MG	1a	3419	1/1	0.92	0.20	-	49,49,49,49	0
59	MG	1A	3437	1/1	0.94	0.08	-	48,48,48,48	0
59	MG	1A	4148	1/1	0.87	0.15	-	60,60,60,60	0
59	MG	2a	3101	1/1	0.92	0.12	-	59,59,59,59	0
59	MG	2a	3156	1/1	0.80	0.09	-	77,77,77,77	0
59	MG	1A	3275	1/1	0.89	0.14	-	38,38,38,38	0
59	MG	1A	3269	1/1	0.93	0.55	-	27,27,27,27	0
59	MG	1a	3394	1/1	0.77	0.18	-	61,61,61,61	0
59	MG	2A	3024	1/1	0.94	0.12	-	43,43,43,43	0
59	MG	2a	3009	1/1	0.87	0.13	-	67,67,67,67	0
59	MG	1A	3644	1/1	0.89	0.08	-	56,56,56,56	0
59	MG	1A	3632	1/1	0.86	0.17	-	53,53,53,53	0
59	MG	2A	3587	1/1	0.80	0.09	-	62,62,62,62	0
59	MG	1A	3397	1/1	0.89	0.37	-	38,38,38,38	0
59	MG	2A	3790	1/1	0.95	0.14	-	37,37,37,37	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3259	1/1	0.51	0.20	-	82,82,82,82	0
59	MG	2A	3147	1/1	0.85	0.10	-	53,53,53,53	0
59	MG	2A	3151	1/1	0.76	0.15	-	74,74,74,74	0
59	MG	1a	3489	1/1	0.87	0.16	-	85,85,85,85	0
59	MG	2A	3276	1/1	0.93	0.17	-	32,32,32,32	0
59	MG	1A	3360	1/1	0.62	0.24	-	42,42,42,42	0
59	MG	1a	3435	1/1	0.86	0.20	-	76,76,76,76	0
59	MG	1A	3122	1/1	0.96	0.46	-	28,28,28,28	0
59	MG	1A	3060	1/1	0.96	0.55	-	26,26,26,26	0
59	MG	1B	216	1/1	0.92	0.20	-	38,38,38,38	0
59	MG	1A	3401	1/1	0.96	0.20	-	28,28,28,28	0
59	MG	2A	3467	1/1	0.94	0.19	-	54,54,54,54	0
59	MG	1A	3352	1/1	0.92	0.10	-	43,43,43,43	0
59	MG	2a	3032	1/1	0.96	0.44	-	54,54,54,54	0
59	MG	1A	3538	1/1	0.95	0.18	-	20,20,20,20	0
59	MG	2A	3344	1/1	0.88	0.16	-	56,56,56,56	0
59	MG	1A	3991	1/1	0.93	0.30	-	40,40,40,40	0
59	MG	2A	3652	1/1	0.85	0.11	-	60,60,60,60	0
59	MG	1A	3088	1/1	0.97	0.53	-	44,44,44,44	0
59	MG	1A	3797	1/1	0.97	0.22	-	35,35,35,35	0
59	MG	1A	3391	1/1	0.94	0.34	-	41,41,41,41	0
59	MG	1A	3212	1/1	0.81	0.40	-	49,49,49,49	0
59	MG	1A	3950	1/1	0.77	0.21	-	47,47,47,47	0
59	MG	1A	3639	1/1	0.90	0.24	-	51,51,51,51	0
59	MG	1A	3661	1/1	0.95	0.08	-	36,36,36,36	0
59	MG	2A	3043	1/1	0.88	0.12	-	57,57,57,57	0
59	MG	1A	3952	1/1	0.68	0.17	-	48,48,48,48	0
59	MG	2a	3231	1/1	0.97	0.10	-	57,57,57,57	0
59	MG	1A	3168	1/1	0.97	0.72	-	38,38,38,38	0
59	MG	2A	3828	1/1	0.88	0.20	-	60,60,60,60	0
59	MG	2D	301	1/1	0.93	0.17	-	47,47,47,47	0
59	MG	1A	3050	1/1	0.93	0.37	-	31,31,31,31	0
59	MG	1A	3715	1/1	0.97	0.07	-	62,62,62,62	0
59	MG	2A	3743	1/1	0.96	0.10	-	40,40,40,40	0
59	MG	2A	3247	1/1	0.80	1.14	-	52,52,52,52	0
59	MG	1A	3630	1/1	0.97	0.18	-	16,16,16,16	0
59	MG	1A	3647	1/1	0.84	0.15	-	56,56,56,56	0
59	MG	1A	3723	1/1	0.92	0.12	-	53,53,53,53	0
59	MG	2a	3162	1/1	0.93	0.10	-	55,55,55,55	0
59	MG	2A	3367	1/1	0.81	0.09	-	52,52,52,52	0
59	MG	2A	3042	1/1	0.80	0.13	-	43,43,43,43	0
59	MG	1A	4122	1/1	0.72	0.18	-	44,44,44,44	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3022	1/1	0.68	0.14	-	46,46,46,46	0
59	MG	2A	3629	1/1	0.93	0.14	-	67,67,67,67	0
59	MG	1A	3059	1/1	0.98	0.21	-	10,10,10,10	0
59	MG	1A	4120	1/1	0.93	0.14	-	33,33,33,33	0
59	MG	1a	3415	1/1	0.83	0.26	-	56,56,56,56	0
59	MG	1a	3432	1/1	0.84	0.16	-	44,44,44,44	0
59	MG	2B	3004	1/1	0.85	0.21	-	53,53,53,53	0
59	MG	2a	3110	1/1	0.82	0.18	-	71,71,71,71	0
59	MG	1A	3478	1/1	0.93	0.30	-	42,42,42,42	0
59	MG	2x	103	1/1	0.85	0.13	-	74,74,74,74	0
59	MG	2B	3018	1/1	0.92	0.17	-	62,62,62,62	0
59	MG	1A	3959	1/1	0.94	0.19	-	55,55,55,55	0
59	MG	1A	3810	1/1	0.84	0.08	-	38,38,38,38	0
59	MG	2A	3789	1/1	0.79	0.12	-	66,66,66,66	0
59	MG	1A	3235	1/1	0.94	0.44	-	34,34,34,34	0
59	MG	2A	3306	1/1	0.97	0.12	-	60,60,60,60	0
59	MG	1A	3382	1/1	0.84	0.17	-	57,57,57,57	0
59	MG	1A	3927	1/1	0.95	0.16	-	12,12,12,12	0
59	MG	1F	307	1/1	0.88	0.15	-	44,44,44,44	0
59	MG	1A	3101	1/1	0.94	0.46	-	30,30,30,30	0
59	MG	1A	3238	1/1	0.85	0.49	-	41,41,41,41	0
59	MG	2A	3354	1/1	0.93	0.12	-	61,61,61,61	0
59	MG	2F	304	1/1	0.91	0.27	-	74,74,74,74	0
59	MG	2a	3097	1/1	0.84	0.16	-	48,48,48,48	0
59	MG	2a	3122	1/1	0.72	0.29	-	67,67,67,67	0
59	MG	1n	502	1/1	0.78	0.25	-	58,58,58,58	0
59	MG	1A	3472	1/1	0.96	0.19	-	26,26,26,26	0
59	MG	2A	3163	1/1	0.91	0.31	-	57,57,57,57	0
59	MG	1T	8001	1/1	0.94	0.14	-	43,43,43,43	0
59	MG	2A	3593	1/1	0.92	0.16	-	37,37,37,37	0
59	MG	2E	304	1/1	0.96	0.11	-	59,59,59,59	0
59	MG	2A	3132	1/1	0.97	0.24	-	21,21,21,21	0
59	MG	1x	104	1/1	0.93	0.15	-	61,61,61,61	0
59	MG	2A	3594	1/1	0.87	0.18	-	64,64,64,64	0
59	MG	2A	3193	1/1	0.81	0.43	-	40,40,40,40	0
59	MG	1A	3910	1/1	0.50	0.23	-	64,64,64,64	0
59	MG	2a	3178	1/1	0.91	0.11	-	52,52,52,52	0
59	MG	2a	3019	1/1	0.85	0.19	-	63,63,63,63	0
59	MG	1A	3117	1/1	0.96	0.43	-	25,25,25,25	0
59	MG	1A	3180	1/1	0.95	0.20	-	39,39,39,39	0
59	MG	2A	3478	1/1	0.98	0.19	-	56,56,56,56	0
59	MG	2A	3644	1/1	0.98	0.07	-	69,69,69,69	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3005	1/1	0.94	0.11	-	47,47,47,47	0
59	MG	1x	106	1/1	0.92	0.11	-	57,57,57,57	0
59	MG	2A	3252	1/1	0.82	0.15	-	58,58,58,58	0
59	MG	2a	3218	1/1	0.89	0.08	-	63,63,63,63	0
59	MG	1A	3845	1/1	0.97	0.42	-	33,33,33,33	0
59	MG	2A	3341	1/1	0.95	0.32	-	53,53,53,53	0
59	MG	2A	3466	1/1	0.95	0.20	-	55,55,55,55	0
59	MG	1A	3665	1/1	0.95	0.14	-	51,51,51,51	0
59	MG	2A	3103	1/1	0.95	0.16	-	62,62,62,62	0
59	MG	1a	3440	1/1	0.84	0.19	-	50,50,50,50	0
59	MG	2A	3440	1/1	0.97	0.15	-	58,58,58,58	0
59	MG	1B	208	1/1	0.88	0.17	-	63,63,63,63	0
59	MG	1A	3064	1/1	0.96	0.17	-	19,19,19,19	0
59	MG	1a	3378	1/1	0.79	0.38	-	63,63,63,63	0
59	MG	1A	3624	1/1	0.94	0.12	-	14,14,14,14	0
59	MG	1A	3267	1/1	0.89	0.13	-	42,42,42,42	0
59	MG	1A	3791	1/1	0.97	0.13	-	34,34,34,34	0
59	MG	1a	3496	1/1	0.80	0.13	-	66,66,66,66	0
59	MG	2a	3071	1/1	0.95	0.11	-	51,51,51,51	0
59	MG	2A	3565	1/1	0.86	0.21	-	49,49,49,49	0
59	MG	2A	3035	1/1	0.90	0.23	-	67,67,67,67	0
59	MG	1a	3527	1/1	0.95	0.18	-	62,62,62,62	0
59	MG	1A	4015	1/1	0.97	0.13	-	52,52,52,52	0
59	MG	2A	3624	1/1	0.96	0.09	-	44,44,44,44	0
59	MG	1A	3487	1/1	0.88	0.40	-	55,55,55,55	0
59	MG	2a	3209	1/1	0.86	0.12	-	57,57,57,57	0
59	MG	2A	3097	1/1	0.69	0.20	-	74,74,74,74	0
59	MG	1B	217	1/1	0.89	0.12	-	41,41,41,41	0
59	MG	2A	3571	1/1	0.96	0.14	-	68,68,68,68	0
59	MG	1a	3542	1/1	0.91	0.09	-	53,53,53,53	0
59	MG	19	101	1/1	0.89	0.16	-	46,46,46,46	0
59	MG	2A	3231	1/1	0.92	0.36	-	75,75,75,75	0
59	MG	1A	3907	1/1	0.92	0.38	-	74,74,74,74	0
59	MG	1A	3309	1/1	0.66	0.39	-	56,56,56,56	0
59	MG	1A	3506	1/1	0.93	0.24	-	41,41,41,41	0
59	MG	2A	3032	1/1	0.93	0.13	-	54,54,54,54	0
59	MG	2A	3216	1/1	0.77	0.33	-	47,47,47,47	0
59	MG	1A	3237	1/1	0.80	0.19	-	52,52,52,52	0
59	MG	1a	3375	1/1	0.94	0.16	-	37,37,37,37	0
59	MG	2a	3099	1/1	0.95	0.16	-	48,48,48,48	0
59	MG	2a	3129	1/1	0.74	0.14	-	75,75,75,75	0
59	MG	1A	3410	1/1	0.89	0.12	-	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3212	1/1	0.92	0.24	-	49,49,49,49	0
59	MG	2A	3192	1/1	0.86	0.35	-	46,46,46,46	0
59	MG	2A	3431	1/1	0.94	0.13	-	54,54,54,54	0
59	MG	2a	3155	1/1	0.94	0.10	-	72,72,72,72	0
59	MG	1a	3476	1/1	0.91	0.07	-	55,55,55,55	0
59	MG	1A	3078	1/1	0.81	0.12	-	48,48,48,48	0
59	MG	1a	3410	1/1	0.92	0.10	-	59,59,59,59	0
59	MG	1A	3426	1/1	0.73	0.53	-	46,46,46,46	0
59	MG	2E	307	1/1	0.84	0.14	-	62,62,62,62	0
59	MG	2a	3153	1/1	0.70	0.09	-	80,80,80,80	0
59	MG	2A	3704	1/1	0.88	0.09	-	53,53,53,53	0
59	MG	2A	3299	1/1	0.98	0.28	-	41,41,41,41	0
59	MG	2A	3003	1/1	0.81	0.23	-	61,61,61,61	0
59	MG	1A	3337	1/1	0.93	0.23	-	48,48,48,48	0
59	MG	1A	3400	1/1	0.95	0.28	-	51,51,51,51	0
59	MG	1A	3184	1/1	0.89	0.14	-	42,42,42,42	0
59	MG	2A	3580	1/1	0.92	0.13	-	29,29,29,29	0
59	MG	1A	3855	1/1	0.93	0.08	-	51,51,51,51	0
59	MG	2A	3760	1/1	0.92	0.13	-	61,61,61,61	0
59	MG	2A	3655	1/1	0.96	0.08	-	62,62,62,62	0
59	MG	1d	502	1/1	0.73	0.21	-	52,52,52,52	0
59	MG	2y	3006	1/1	0.33	0.09	-	89,89,89,89	0
59	MG	1A	3099	1/1	0.87	0.44	-	31,31,31,31	0
59	MG	1A	3230	1/1	0.93	0.64	-	40,40,40,40	0
59	MG	1A	3292	1/1	0.53	0.23	-	71,71,71,71	0
59	MG	1A	4054	1/1	0.89	0.20	-	57,57,57,57	0
59	MG	1a	3486	1/1	0.89	0.11	-	59,59,59,59	0
59	MG	2A	3453	1/1	0.85	0.26	-	60,60,60,60	0
59	MG	2A	3338	1/1	0.91	0.38	-	66,66,66,66	0
59	MG	1a	3456	1/1	0.90	0.18	-	32,32,32,32	0
59	MG	1a	3509	1/1	0.90	0.06	-	61,61,61,61	0
59	MG	1A	4121	1/1	0.97	0.15	-	30,30,30,30	0
59	MG	2A	3784	1/1	0.88	0.14	-	56,56,56,56	0
59	MG	1B	203	1/1	0.97	0.22	-	49,49,49,49	0
59	MG	2A	3196	1/1	0.92	0.34	-	51,51,51,51	0
59	MG	2Q	3002	1/1	0.81	0.25	-	48,48,48,48	0
59	MG	2a	3192	1/1	0.93	0.17	-	57,57,57,57	0
59	MG	2a	3042	1/1	0.92	0.15	-	70,70,70,70	0
59	MG	2A	3353	1/1	0.89	0.14	-	57,57,57,57	0
59	MG	1a	3526	1/1	0.95	0.17	-	50,50,50,50	0
59	MG	1A	3254	1/1	0.71	0.20	-	64,64,64,64	0
59	MG	1A	3949	1/1	0.78	0.18	-	52,52,52,52	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3265	1/1	0.94	0.32	-	73,73,73,73	0
59	MG	2A	3135	1/1	0.95	0.60	-	58,58,58,58	0
59	MG	1O	3005	1/1	0.96	0.17	-	40,40,40,40	0
59	MG	1A	3886	1/1	0.92	0.06	-	38,38,38,38	0
59	MG	1B	202	1/1	0.93	0.28	-	37,37,37,37	0
59	MG	2a	3034	1/1	0.92	0.17	-	53,53,53,53	0
59	MG	1v	3001	1/1	0.77	0.55	-	60,60,60,60	0
59	MG	1A	3364	1/1	0.96	0.50	-	34,34,34,34	0
59	MG	1a	3309	1/1	0.89	0.09	-	61,61,61,61	0
59	MG	1A	3985	1/1	0.95	0.17	-	53,53,53,53	0
59	MG	2A	3083	1/1	0.83	0.17	-	61,61,61,61	0
59	MG	1B	225	1/1	0.67	0.15	-	59,59,59,59	0
59	MG	1y	104	1/1	0.82	0.29	-	80,80,80,80	0
59	MG	2a	3154	1/1	0.79	0.16	-	42,42,42,42	0
59	MG	2B	3020	1/1	0.76	0.30	-	80,80,80,80	0
59	MG	2a	3017	1/1	0.80	0.18	-	76,76,76,76	0
59	MG	2A	3333	1/1	0.82	0.24	-	60,60,60,60	0
59	MG	2A	3326	1/1	0.89	0.14	-	46,46,46,46	0
59	MG	2A	3818	1/1	0.91	0.11	-	63,63,63,63	0
59	MG	2A	3664	1/1	0.90	0.10	-	68,68,68,68	0
59	MG	1A	3102	1/1	0.92	0.47	-	46,46,46,46	0
59	MG	1A	3187	1/1	0.80	0.20	-	39,39,39,39	0
59	MG	1a	3511	1/1	0.85	0.16	-	46,46,46,46	0
59	MG	1A	3711	1/1	0.83	0.13	-	34,34,34,34	0
59	MG	1a	3371	1/1	0.85	0.48	-	55,55,55,55	0
59	MG	2f	3002	1/1	0.91	0.11	-	69,69,69,69	0
59	MG	1A	3718	1/1	0.93	0.08	-	39,39,39,39	0
59	MG	2a	3012	1/1	0.83	0.11	-	63,63,63,63	0
59	MG	1A	3500	1/1	0.94	0.29	-	54,54,54,54	0
59	MG	1B	210	1/1	0.84	0.38	-	51,51,51,51	0
59	MG	1A	3459	1/1	0.91	0.34	-	58,58,58,58	0
59	MG	2A	3051	1/1	0.85	0.15	-	67,67,67,67	0
59	MG	2A	3681	1/1	0.90	0.08	-	62,62,62,62	0
59	MG	1Z	302	1/1	0.90	0.20	-	48,48,48,48	0
59	MG	1A	3876	1/1	0.71	0.20	-	60,60,60,60	0
59	MG	1A	3822	1/1	0.98	0.14	-	11,11,11,11	0
59	MG	1A	4029	1/1	0.90	0.07	-	46,46,46,46	0
59	MG	1A	3368	1/1	0.94	0.34	-	39,39,39,39	0
59	MG	1x	116	1/1	0.85	0.19	-	70,70,70,70	0
59	MG	1B	226	1/1	0.87	0.16	-	56,56,56,56	0
59	MG	15	102	1/1	0.93	0.22	-	25,25,25,25	0
59	MG	2A	3523	1/1	0.91	0.10	-	63,63,63,63	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3533	1/1	0.93	0.16	-	55,55,55,55	0
59	MG	1A	4022	1/1	0.95	0.25	-	39,39,39,39	0
59	MG	1a	3498	1/1	0.92	0.13	-	63,63,63,63	0
59	MG	1A	4221	1/1	0.94	0.39	-	31,31,31,31	0
59	MG	2A	3807	1/1	0.96	0.14	-	38,38,38,38	0
59	MG	1A	3473	1/1	0.92	0.22	-	54,54,54,54	0
59	MG	2A	3740	1/1	0.92	0.13	-	48,48,48,48	0
59	MG	2A	3443	1/1	0.94	0.12	-	30,30,30,30	0
59	MG	1A	3383	1/1	0.92	0.15	-	38,38,38,38	0
59	MG	1A	3957	1/1	0.96	0.09	-	33,33,33,33	0
59	MG	1a	3372	1/1	0.84	0.14	-	64,64,64,64	0
59	MG	1A	3894	1/1	0.97	0.13	-	43,43,43,43	0
59	MG	1A	4071	1/1	0.87	0.09	-	50,50,50,50	0
59	MG	1A	3231	1/1	0.94	0.45	-	39,39,39,39	0
59	MG	2A	3452	1/1	0.87	0.13	-	56,56,56,56	0
59	MG	2A	3800	1/1	0.89	0.08	-	62,62,62,62	0
59	MG	2A	3214	1/1	0.78	0.23	-	58,58,58,58	0
59	MG	1A	3588	1/1	0.96	0.16	-	30,30,30,30	0
59	MG	2A	3584	1/1	0.86	0.25	-	52,52,52,52	0
59	MG	1A	3469	1/1	0.92	0.21	-	33,33,33,33	0
59	MG	1A	3483	1/1	0.91	0.13	-	56,56,56,56	0
59	MG	2A	3778	1/1	0.89	0.10	-	63,63,63,63	0
59	MG	1A	3246	1/1	0.90	0.33	-	47,47,47,47	0
59	MG	2A	3536	1/1	0.96	0.19	-	41,41,41,41	0
59	MG	1A	3496	1/1	0.93	0.16	-	50,50,50,50	0
59	MG	1a	3370	1/1	0.76	0.19	-	67,67,67,67	0
59	MG	1a	3477	1/1	0.94	0.18	-	67,67,67,67	0
59	MG	1a	3510	1/1	0.65	0.15	-	53,53,53,53	0
59	MG	1y	102	1/1	0.74	0.50	-	77,77,77,77	0
59	MG	1A	3899	1/1	0.95	0.09	-	50,50,50,50	0
59	MG	2A	3763	1/1	0.94	0.06	-	66,66,66,66	0
59	MG	2A	3295	1/1	0.87	0.13	-	58,58,58,58	0
59	MG	1A	3488	1/1	0.83	0.18	-	45,45,45,45	0
59	MG	1A	3520	1/1	0.99	0.24	-	16,16,16,16	0
59	MG	1A	3909	1/1	0.92	0.09	-	58,58,58,58	0
59	MG	1w	111	1/1	0.74	0.15	-	80,80,80,80	0
59	MG	2A	3106	1/1	0.91	0.16	-	57,57,57,57	0
59	MG	1A	3130	1/1	0.98	0.44	-	32,32,32,32	0
59	MG	2A	3161	1/1	0.88	0.33	-	67,67,67,67	0
59	MG	1x	110	1/1	0.81	0.10	-	59,59,59,59	0
59	MG	2a	3215	1/1	0.76	0.18	-	75,75,75,75	0
59	MG	1A	3955	1/1	0.52	0.09	-	68,68,68,68	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3591	1/1	0.88	0.15	-	63,63,63,63	0
59	MG	1A	4080	1/1	0.98	0.21	-	28,28,28,28	0
59	MG	1A	4065	1/1	0.89	0.19	-	45,45,45,45	0
59	MG	1a	3466	1/1	0.92	0.21	-	72,72,72,72	0
59	MG	2A	3217	1/1	0.82	0.40	-	48,48,48,48	0
59	MG	1A	3171	1/1	0.96	0.21	-	25,25,25,25	0
59	MG	1A	3585	1/1	0.98	0.16	-	33,33,33,33	0
59	MG	1A	4009	1/1	0.93	0.13	-	71,71,71,71	0
59	MG	2B	3015	1/1	0.69	0.21	-	64,64,64,64	0
59	MG	1A	4050	1/1	0.92	0.07	-	45,45,45,45	0
59	MG	2A	3549	1/1	0.93	0.14	-	55,55,55,55	0
59	MG	2A	3610	1/1	0.92	0.48	-	40,40,40,40	0
59	MG	1A	3002	1/1	0.76	0.22	-	53,53,53,53	0
59	MG	1A	3268	1/1	0.98	0.57	-	32,32,32,32	0
59	MG	1A	3443	1/1	0.78	0.61	-	49,49,49,49	0
59	MG	1a	3402	1/1	0.92	0.12	-	60,60,60,60	0
59	MG	1A	3552	1/1	0.97	0.14	-	7,7,7,7	0
59	MG	1A	3744	1/1	0.88	0.11	-	44,44,44,44	0
59	MG	1A	3895	1/1	0.85	0.15	-	39,39,39,39	0
59	MG	2A	3464	1/1	0.94	0.13	-	46,46,46,46	0
59	MG	1A	3518	1/1	0.86	0.36	-	63,63,63,63	0
59	MG	2a	3145	1/1	0.98	0.08	-	45,45,45,45	0
59	MG	1Q	205	1/1	0.96	0.09	-	37,37,37,37	0
59	MG	2E	306	1/1	0.93	0.16	-	29,29,29,29	0
59	MG	20	3001	1/1	0.90	0.19	-	66,66,66,66	0
59	MG	1A	3549	1/1	0.94	0.20	-	40,40,40,40	0
59	MG	1x	109	1/1	0.59	0.18	-	68,68,68,68	0
59	MG	2A	3321	1/1	0.86	0.24	-	58,58,58,58	0
59	MG	1A	4144	1/1	0.92	0.22	-	42,42,42,42	0
59	MG	2A	3050	1/1	0.91	0.44	-	48,48,48,48	0
59	MG	2A	3377	1/1	0.84	0.19	-	36,36,36,36	0
59	MG	1a	3507	1/1	0.95	0.31	-	64,64,64,64	0
59	MG	2A	3838	1/1	0.90	0.11	-	51,51,51,51	0
59	MG	2A	3516	1/1	0.67	0.63	-	64,64,64,64	0
59	MG	18	101	1/1	0.80	0.38	-	62,62,62,62	0
59	MG	2A	3202	1/1	0.75	0.17	-	51,51,51,51	0
59	MG	1A	3753	1/1	0.93	0.15	-	25,25,25,25	0
59	MG	1A	3809	1/1	0.84	0.17	-	48,48,48,48	0
59	MG	2A	3724	1/1	0.92	0.19	-	34,34,34,34	0
59	MG	1A	4141	1/1	0.88	0.19	-	36,36,36,36	0
59	MG	11	101	1/1	0.83	0.19	-	45,45,45,45	0
59	MG	1B	215	1/1	0.94	0.09	-	44,44,44,44	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	4023	1/1	0.78	0.10	-	49,49,49,49	0
59	MG	2A	3801	1/1	0.86	0.24	-	55,55,55,55	0
59	MG	25	102	1/1	0.94	0.80	-	53,53,53,53	0
59	MG	1A	3713	1/1	0.89	0.24	-	49,49,49,49	0
59	MG	2A	3544	1/1	0.90	0.16	-	64,64,64,64	0
59	MG	2A	3568	1/1	0.74	0.15	-	74,74,74,74	0
59	MG	2A	3822	1/1	0.79	0.17	-	33,33,33,33	0
59	MG	2A	3788	1/1	0.94	0.31	-	57,57,57,57	0
59	MG	1A	3516	1/1	0.88	0.14	-	49,49,49,49	0
59	MG	1A	3340	1/1	0.85	0.83	-	46,46,46,46	0
59	MG	1A	4043	1/1	0.92	0.14	-	39,39,39,39	0
59	MG	1a	3548	1/1	0.86	0.20	-	74,74,74,74	0
59	MG	1A	3953	1/1	0.87	0.07	-	64,64,64,64	0
59	MG	1A	3942	1/1	0.93	0.25	-	45,45,45,45	0
59	MG	1A	3296	1/1	0.91	0.34	-	55,55,55,55	0
59	MG	2w	3001	1/1	0.82	0.21	-	56,56,56,56	0
59	MG	2A	3052	1/1	0.76	0.26	-	56,56,56,56	0
59	MG	2A	3520	1/1	0.87	0.15	-	79,79,79,79	0
59	MG	1A	4150	1/1	0.94	0.10	-	35,35,35,35	0
59	MG	1A	3649	1/1	0.86	0.09	-	55,55,55,55	0
59	MG	1A	3980	1/1	0.84	0.15	-	39,39,39,39	0
59	MG	1a	3356	1/1	0.84	0.14	-	41,41,41,41	0
59	MG	2A	3266	1/1	0.89	0.15	-	42,42,42,42	0
59	MG	1A	3346	1/1	0.85	0.14	-	52,52,52,52	0
59	MG	1A	3066	1/1	0.97	0.25	-	27,27,27,27	0
59	MG	2A	3697	1/1	0.92	0.11	-	25,25,25,25	0
59	MG	1A	3445	1/1	0.61	0.27	-	55,55,55,55	0
59	MG	1A	3125	1/1	0.93	0.62	-	32,32,32,32	0
59	MG	1a	3393	1/1	0.85	0.13	-	70,70,70,70	0
59	MG	2a	3219	1/1	0.83	0.11	-	73,73,73,73	0
59	MG	1y	105	1/1	0.83	0.17	-	80,80,80,80	0
59	MG	2A	3409	1/1	0.93	0.16	-	50,50,50,50	0
59	MG	1a	3521	1/1	0.95	0.08	-	46,46,46,46	0
59	MG	2A	3792	1/1	0.95	0.18	-	63,63,63,63	0
59	MG	1A	3396	1/1	0.88	0.24	-	46,46,46,46	0
59	MG	2a	3185	1/1	0.93	0.28	-	64,64,64,64	0
59	MG	1a	3336	1/1	0.84	0.15	-	60,60,60,60	0
59	MG	1A	3877	1/1	0.93	0.10	-	44,44,44,44	0
59	MG	2y	3004	1/1	0.80	0.27	-	66,66,66,66	0
59	MG	2A	3525	1/1	0.92	0.09	-	55,55,55,55	0
59	MG	2A	3140	1/1	0.91	0.25	-	72,72,72,72	0
59	MG	1A	3136	1/1	0.95	0.47	-	26,26,26,26	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3329	1/1	0.83	0.35	-	59,59,59,59	0
59	MG	2A	3737	1/1	0.87	0.09	-	47,47,47,47	0
59	MG	1A	3407	1/1	0.94	0.14	-	46,46,46,46	0
59	MG	1a	3376	1/1	0.71	0.18	-	62,62,62,62	0
59	MG	13	3001	1/1	0.88	0.19	-	39,39,39,39	0
59	MG	1A	3774	1/1	0.60	0.17	-	52,52,52,52	0
59	MG	2A	3522	1/1	0.94	0.10	-	48,48,48,48	0
59	MG	1A	3684	1/1	0.97	0.12	-	28,28,28,28	0
59	MG	1A	3902	1/1	0.80	0.21	-	45,45,45,45	0
59	MG	2A	3280	1/1	0.98	0.08	-	49,49,49,49	0
59	MG	2a	3212	1/1	0.72	0.21	-	73,73,73,73	0
59	MG	1a	3506	1/1	0.97	0.10	-	59,59,59,59	0
59	MG	1A	3539	1/1	0.89	0.17	-	26,26,26,26	0
59	MG	1A	3301	1/1	0.86	0.59	-	37,37,37,37	0
59	MG	2a	3005	1/1	0.80	0.13	-	74,74,74,74	0
59	MG	1A	3116	1/1	0.95	0.07	-	43,43,43,43	0
59	MG	2A	3554	1/1	0.98	0.10	-	62,62,62,62	0
59	MG	2A	3033	1/1	0.91	0.08	-	58,58,58,58	0
59	MG	2A	3779	1/1	0.91	0.07	-	67,67,67,67	0
59	MG	2A	3318	1/1	0.93	0.19	-	53,53,53,53	0
59	MG	2A	3725	1/1	0.67	0.37	-	64,64,64,64	0
59	MG	2A	3264	1/1	0.83	0.12	-	60,60,60,60	0
59	MG	2v	103	1/1	0.88	0.11	-	70,70,70,70	0
59	MG	1A	3743	1/1	0.96	0.19	-	38,38,38,38	0
59	MG	2A	3182	1/1	0.83	0.15	-	61,61,61,61	0
59	MG	2A	3316	1/1	0.94	0.13	-	49,49,49,49	0
59	MG	1A	3645	1/1	0.84	0.27	-	63,63,63,63	0
59	MG	2A	3272	1/1	0.84	0.56	-	44,44,44,44	0
59	MG	2A	3668	1/1	0.98	0.25	-	43,43,43,43	0
59	MG	1A	3345	1/1	0.92	0.35	-	37,37,37,37	0
59	MG	2A	3058	1/1	0.86	0.21	-	45,45,45,45	0
59	MG	2A	3100	1/1	0.94	0.10	-	65,65,65,65	0
59	MG	1A	4095	1/1	0.84	0.25	-	61,61,61,61	0
59	MG	1E	307	1/1	0.73	0.28	-	40,40,40,40	0
59	MG	1A	4026	1/1	0.94	0.12	-	45,45,45,45	0
59	MG	1a	3475	1/1	0.96	0.11	-	48,48,48,48	0
59	MG	1A	3143	1/1	0.94	0.20	-	56,56,56,56	0
59	MG	2A	3263	1/1	0.94	0.19	-	47,47,47,47	0
59	MG	1A	3887	1/1	0.97	0.13	-	50,50,50,50	0
59	MG	2A	3282	1/1	0.91	0.20	-	54,54,54,54	0
59	MG	2a	3106	1/1	0.75	0.11	-	67,67,67,67	0
59	MG	1A	3565	1/1	0.95	0.11	-	28,28,28,28	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3783	1/1	0.95	0.13	-	73,73,73,73	0
59	MG	1A	3351	1/1	0.95	0.29	-	31,31,31,31	0
59	MG	1B	237	1/1	0.97	0.07	-	39,39,39,39	0
59	MG	1A	3975	1/1	0.96	0.30	-	23,23,23,23	0
59	MG	1a	3353	1/1	0.98	0.23	-	58,58,58,58	0
59	MG	2A	3397	1/1	0.92	0.12	-	23,23,23,23	0
59	MG	1A	3504	1/1	0.88	0.11	-	53,53,53,53	0
59	MG	1E	310	1/1	0.96	0.14	-	34,34,34,34	0
59	MG	2A	3670	1/1	0.95	0.12	-	64,64,64,64	0
59	MG	2A	3597	1/1	0.61	0.16	-	65,65,65,65	0
59	MG	1A	3826	1/1	0.96	0.12	-	42,42,42,42	0
59	MG	1x	115	1/1	0.95	0.14	-	56,56,56,56	0
59	MG	1A	3086	1/1	0.96	0.27	-	31,31,31,31	0
59	MG	1B	213	1/1	0.92	0.17	-	43,43,43,43	0
59	MG	2x	104	1/1	0.87	0.12	-	71,71,71,71	0
59	MG	1w	103	1/1	0.83	0.10	-	56,56,56,56	0
59	MG	2A	3226	1/1	0.83	0.36	-	53,53,53,53	0
59	MG	2a	3063	1/1	0.96	0.09	-	55,55,55,55	0
59	MG	1A	4160	1/1	0.83	0.14	-	53,53,53,53	0
59	MG	2A	3711	1/1	0.93	0.08	-	50,50,50,50	0
59	MG	1a	3500	1/1	0.94	0.12	-	64,64,64,64	0
59	MG	1A	4017	1/1	0.91	0.10	-	44,44,44,44	0
59	MG	2a	3176	1/1	0.88	0.14	-	75,75,75,75	0
59	MG	2A	3599	1/1	0.91	0.18	-	65,65,65,65	0
59	MG	1A	3223	1/1	0.76	0.39	-	40,40,40,40	0
59	MG	2A	3434	1/1	0.95	0.13	-	24,24,24,24	0
59	MG	2a	3194	1/1	0.89	0.11	-	68,68,68,68	0
59	MG	1A	3976	1/1	0.88	0.11	-	56,56,56,56	0
59	MG	2a	3197	1/1	0.96	0.12	-	54,54,54,54	0
59	MG	1A	3414	1/1	0.92	0.24	-	42,42,42,42	0
59	MG	1a	3485	1/1	0.96	0.09	-	49,49,49,49	0
59	MG	1A	3862	1/1	0.89	0.08	-	38,38,38,38	0
59	MG	2A	3473	1/1	0.83	0.36	-	63,63,63,63	0
59	MG	2A	3249	1/1	0.77	0.33	-	62,62,62,62	0
59	MG	1R	203	1/1	0.77	0.19	-	33,33,33,33	0
59	MG	2a	3102	1/1	0.88	0.12	-	66,66,66,66	0
59	MG	2A	3729	1/1	0.72	0.15	-	57,57,57,57	0
59	MG	1A	3447	1/1	0.88	0.71	-	38,38,38,38	0
59	MG	1A	4079	1/1	0.92	0.19	-	68,68,68,68	0
59	MG	2A	3586	1/1	0.89	0.08	-	46,46,46,46	0
59	MG	1A	3878	1/1	0.90	0.10	-	38,38,38,38	0
59	MG	1A	3328	1/1	0.88	0.50	-	27,27,27,27	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2a	3187	1/1	0.90	0.13	-	71,71,71,71	0
59	MG	1A	3311	1/1	0.96	0.36	-	39,39,39,39	0
59	MG	1a	3413	1/1	0.94	0.13	-	54,54,54,54	0
59	MG	1a	3420	1/1	0.76	0.15	-	65,65,65,65	0
59	MG	1A	3629	1/1	0.90	0.15	-	17,17,17,17	0
59	MG	10	108	1/1	0.98	0.08	-	39,39,39,39	0
59	MG	1A	3849	1/1	0.91	0.10	-	54,54,54,54	0
59	MG	2A	3496	1/1	0.97	0.13	-	49,49,49,49	0
59	MG	2A	3034	1/1	0.97	0.32	-	50,50,50,50	0
59	MG	1A	3137	1/1	0.92	0.14	-	43,43,43,43	0
59	MG	1a	3398	1/1	0.85	0.20	-	58,58,58,58	0
59	MG	1A	3313	1/1	0.93	0.19	-	38,38,38,38	0
59	MG	1A	4129	1/1	0.92	0.16	-	55,55,55,55	0
59	MG	1A	3406	1/1	0.80	0.17	-	50,50,50,50	0
59	MG	1w	105	1/1	0.95	0.10	-	39,39,39,39	0
59	MG	1A	3928	1/1	0.98	0.12	-	16,16,16,16	0
59	MG	2A	3569	1/1	0.80	0.15	-	59,59,59,59	0
59	MG	2A	3616	1/1	0.98	0.21	-	37,37,37,37	0
59	MG	2A	3684	1/1	0.75	0.07	-	33,33,33,33	0
59	MG	2B	3005	1/1	0.96	0.24	-	61,61,61,61	0
59	MG	1A	3380	1/1	0.85	0.14	-	33,33,33,33	0
59	MG	2a	3025	1/1	0.81	0.13	-	52,52,52,52	0
59	MG	2A	3471	1/1	0.93	0.22	-	58,58,58,58	0
59	MG	1A	4194	1/1	0.91	0.52	-	33,33,33,33	0
59	MG	1a	3422	1/1	0.91	0.20	-	50,50,50,50	0
59	MG	1a	3472	1/1	0.99	0.12	-	64,64,64,64	0
59	MG	1A	3493	1/1	0.90	0.20	-	55,55,55,55	0
59	MG	2A	3285	1/1	0.79	0.45	-	52,52,52,52	0
59	MG	2A	3535	1/1	0.66	0.12	-	63,63,63,63	0
59	MG	2A	3691	1/1	0.96	0.10	-	29,29,29,29	0
59	MG	1A	3263	1/1	0.88	0.20	-	45,45,45,45	0
59	MG	1A	3420	1/1	0.73	0.50	-	60,60,60,60	0
59	MG	2A	3728	1/1	0.86	0.09	-	24,24,24,24	0
59	MG	2a	3233	1/1	0.91	0.08	-	50,50,50,50	0
59	MG	1A	3694	1/1	0.96	0.14	-	29,29,29,29	0
59	MG	1A	3278	1/1	0.74	0.19	-	62,62,62,62	0
59	MG	2A	3846	1/1	0.94	0.37	-	38,38,38,38	0
59	MG	1a	3326	1/1	0.80	0.21	-	56,56,56,56	0
59	MG	1A	3342	1/1	0.85	0.22	-	42,42,42,42	0
59	MG	1A	3918	1/1	0.87	0.17	-	27,27,27,27	0
59	MG	1A	3784	1/1	0.94	0.14	-	14,14,14,14	0
59	MG	2A	3064	1/1	0.98	0.17	-	16,16,16,16	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3155	1/1	0.73	0.14	-	61,61,61,61	0
59	MG	1A	3731	1/1	0.96	0.12	-	33,33,33,33	0
59	MG	1A	3222	1/1	0.94	0.21	-	38,38,38,38	0
59	MG	1A	3096	1/1	0.87	0.34	-	32,32,32,32	0
59	MG	1A	3893	1/1	0.71	0.37	-	39,39,39,39	0
59	MG	1A	4115	1/1	0.65	0.28	-	60,60,60,60	0
59	MG	1A	4068	1/1	0.96	0.12	-	38,38,38,38	0
59	MG	2A	3378	1/1	0.91	0.16	-	55,55,55,55	0
59	MG	2A	3732	1/1	0.92	0.18	-	47,47,47,47	0
59	MG	2A	3184	1/1	0.96	0.22	-	45,45,45,45	0
59	MG	1A	3964	1/1	0.95	0.12	-	36,36,36,36	0
59	MG	2A	3444	1/1	0.96	0.15	-	31,31,31,31	0
59	MG	1A	3742	1/1	0.95	0.12	-	29,29,29,29	0
59	MG	1A	3242	1/1	0.93	0.43	-	24,24,24,24	0
59	MG	1A	3510	1/1	0.88	0.23	-	41,41,41,41	0
59	MG	1A	3273	1/1	0.87	0.18	-	49,49,49,49	0
59	MG	1A	3022	1/1	0.71	0.32	-	39,39,39,39	0
59	MG	1a	3503	1/1	0.85	0.15	-	71,71,71,71	0
59	MG	2a	3008	1/1	0.86	0.14	-	63,63,63,63	0
59	MG	1A	3190	1/1	0.92	0.18	-	39,39,39,39	0
59	MG	2A	3631	1/1	0.95	0.13	-	54,54,54,54	0
59	MG	2A	3658	1/1	0.88	0.21	-	68,68,68,68	0
59	MG	1A	4034	1/1	0.77	0.09	-	61,61,61,61	0
59	MG	1A	3087	1/1	0.81	0.14	-	56,56,56,56	0
59	MG	2A	3343	1/1	0.85	0.15	-	60,60,60,60	0
59	MG	2A	3517	1/1	0.84	0.11	-	58,58,58,58	0
59	MG	2A	3577	1/1	0.93	0.08	-	36,36,36,36	0
59	MG	2A	3116	1/1	0.92	0.46	-	35,35,35,35	0
59	MG	2A	3463	1/1	0.97	0.13	-	43,43,43,43	0
59	MG	1A	3120	1/1	0.97	0.19	-	10,10,10,10	0
59	MG	2a	3059	1/1	0.94	0.08	-	53,53,53,53	0
59	MG	1a	3473	1/1	0.96	0.14	-	37,37,37,37	0
59	MG	1A	3511	1/1	0.86	0.17	-	51,51,51,51	0
59	MG	2A	3712	1/1	0.80	0.13	-	40,40,40,40	0
59	MG	2A	3019	1/1	0.92	0.27	-	54,54,54,54	0
59	MG	2A	3641	1/1	0.92	0.09	-	75,75,75,75	0
59	MG	2A	3669	1/1	0.85	0.28	-	69,69,69,69	0
59	MG	2A	3435	1/1	0.89	0.11	-	47,47,47,47	0
59	MG	1A	3138	1/1	0.89	0.14	-	59,59,59,59	0
59	MG	2v	104	1/1	0.98	0.15	-	52,52,52,52	0
59	MG	1A	3307	1/1	0.86	0.15	-	50,50,50,50	0
59	MG	1a	3524	1/1	0.91	0.10	-	71,71,71,71	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1f	3002	1/1	0.95	0.18	-	33,33,33,33	0
59	MG	1A	3779	1/1	0.95	0.33	-	36,36,36,36	0
59	MG	1A	3983	1/1	0.93	0.23	-	63,63,63,63	0
59	MG	2a	3114	1/1	0.77	0.23	-	59,59,59,59	0
59	MG	2A	3047	1/1	0.97	0.14	-	47,47,47,47	0
59	MG	10	106	1/1	0.83	0.17	-	63,63,63,63	0
59	MG	2A	3602	1/1	0.84	0.19	-	51,51,51,51	0
59	MG	2A	3073	1/1	0.78	0.36	-	63,63,63,63	0
59	MG	2A	3808	1/1	0.91	0.10	-	54,54,54,54	0
59	MG	1A	3656	1/1	0.92	0.12	-	57,57,57,57	0
59	MG	2a	3040	1/1	0.95	0.16	-	47,47,47,47	0
59	MG	1A	3509	1/1	0.95	0.16	-	40,40,40,40	0
59	MG	2A	3446	1/1	0.93	0.10	-	44,44,44,44	0
59	MG	2A	3649	1/1	0.97	0.09	-	47,47,47,47	0
59	MG	1A	3024	1/1	0.96	0.43	-	19,19,19,19	0
59	MG	1A	3425	1/1	0.92	0.10	-	49,49,49,49	0
59	MG	1A	3931	1/1	0.86	0.10	-	64,64,64,64	0
59	MG	2A	3676	1/1	0.82	0.09	-	60,60,60,60	0
59	MG	1A	3304	1/1	0.85	0.35	-	36,36,36,36	0
59	MG	1S	3003	1/1	0.95	0.14	-	61,61,61,61	0
59	MG	2a	3038	1/1	0.94	0.10	-	78,78,78,78	0
59	MG	1A	3245	1/1	0.99	0.43	-	21,21,21,21	0
59	MG	2A	3469	1/1	0.96	0.15	-	60,60,60,60	0
59	MG	1A	3859	1/1	0.88	0.14	-	37,37,37,37	0
59	MG	1A	3232	1/1	0.83	0.15	-	51,51,51,51	0
59	MG	1A	3772	1/1	0.91	0.22	-	46,46,46,46	0
59	MG	2R	202	1/1	0.83	0.27	-	61,61,61,61	0
59	MG	1A	3484	1/1	0.97	0.35	-	27,27,27,27	0
59	MG	1A	3815	1/1	0.97	0.18	-	25,25,25,25	0
59	MG	1E	304	1/1	0.98	0.42	-	27,27,27,27	0
59	MG	1a	3417	1/1	0.93	0.38	-	50,50,50,50	0
59	MG	1A	3911	1/1	0.75	0.21	-	62,62,62,62	0
59	MG	1A	4178	1/1	0.67	0.20	-	60,60,60,60	0
59	MG	2A	3730	1/1	0.94	0.26	-	53,53,53,53	0
59	MG	1A	3162	1/1	0.96	0.63	-	30,30,30,30	0
59	MG	2A	3143	1/1	0.84	0.11	-	77,77,77,77	0
59	MG	1a	3334	1/1	0.97	0.13	-	49,49,49,49	0
59	MG	1A	4158	1/1	0.98	0.16	-	20,20,20,20	0
59	MG	1A	3051	1/1	0.98	0.18	-	49,49,49,49	0
59	MG	1A	4063	1/1	0.78	0.18	-	45,45,45,45	0
59	MG	2A	3791	1/1	0.76	0.21	-	62,62,62,62	0
59	MG	1A	4133	1/1	0.94	0.12	-	35,35,35,35	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3097	1/1	0.96	0.15	-	35,35,35,35	0
59	MG	1A	3315	1/1	0.80	0.47	-	34,34,34,34	0
59	MG	1A	3334	1/1	0.92	0.41	-	39,39,39,39	0
59	MG	2A	3777	1/1	0.91	0.11	-	55,55,55,55	0
59	MG	1F	309	1/1	0.92	0.15	-	36,36,36,36	0
59	MG	2A	3696	1/1	0.73	0.07	-	73,73,73,73	0
59	MG	1A	3494	1/1	0.88	0.19	-	49,49,49,49	0
59	MG	2A	3780	1/1	0.91	0.09	-	62,62,62,62	0
59	MG	1a	3545	1/1	0.92	0.10	-	46,46,46,46	0
59	MG	1A	3058	1/1	0.85	0.17	-	50,50,50,50	0
59	MG	2a	3208	1/1	0.75	0.25	-	63,63,63,63	0
59	MG	1A	3323	1/1	0.86	0.73	-	28,28,28,28	0
59	MG	2a	3096	1/1	0.92	0.34	-	56,56,56,56	0
59	MG	11	102	1/1	0.96	0.13	-	25,25,25,25	0
59	MG	2A	3376	1/1	0.90	0.16	-	54,54,54,54	0
59	MG	1A	3367	1/1	0.71	0.16	-	59,59,59,59	0
59	MG	1A	3135	1/1	0.97	0.38	-	22,22,22,22	0
59	MG	2q	204	1/1	0.87	0.17	-	70,70,70,70	0
59	MG	2A	3579	1/1	0.88	0.23	-	32,32,32,32	0
59	MG	2B	3009	1/1	0.92	0.11	-	54,54,54,54	0
59	MG	1A	3776	1/1	0.94	0.04	-	55,55,55,55	0
59	MG	2A	3677	1/1	0.62	0.11	-	63,63,63,63	0
59	MG	1A	3255	1/1	0.83	0.12	-	54,54,54,54	0
59	MG	2A	3087	1/1	0.75	0.13	-	53,53,53,53	0
59	MG	1A	3444	1/1	0.96	0.25	-	45,45,45,45	0
59	MG	2A	3075	1/1	0.65	0.24	-	73,73,73,73	0
59	MG	2A	3061	1/1	0.95	0.39	-	42,42,42,42	0
59	MG	2A	3632	1/1	0.94	0.13	-	29,29,29,29	0
59	MG	2A	3686	1/1	0.80	0.11	-	48,48,48,48	0
59	MG	2A	3190	1/1	0.90	0.35	-	37,37,37,37	0
59	MG	2A	3753	1/1	0.91	0.18	-	52,52,52,52	0
59	MG	1A	3280	1/1	0.83	0.18	-	39,39,39,39	0
59	MG	1a	3568	1/1	0.89	0.07	-	53,53,53,53	0
59	MG	2A	3450	1/1	0.96	0.10	-	53,53,53,53	0
59	MG	2A	3401	1/1	0.90	0.16	-	59,59,59,59	0
59	MG	1A	3687	1/1	0.97	0.19	-	54,54,54,54	0
59	MG	2A	3298	1/1	0.96	0.23	-	41,41,41,41	0
59	MG	1A	3881	1/1	0.87	0.19	-	47,47,47,47	0
59	MG	1a	3418	1/1	0.82	0.30	-	64,64,64,64	0
59	MG	1B	236	1/1	0.94	0.26	-	57,57,57,57	0
59	MG	2a	3127	1/1	0.69	0.11	-	74,74,74,74	0
59	MG	1A	3771	1/1	0.87	0.12	-	33,33,33,33	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3475	1/1	0.93	0.19	-	53,53,53,53	0
59	MG	2A	3786	1/1	0.96	0.15	-	55,55,55,55	0
59	MG	1A	3579	1/1	0.95	0.15	-	30,30,30,30	0
59	MG	1A	3387	1/1	0.78	0.18	-	45,45,45,45	0
59	MG	2a	3226	1/1	0.91	0.19	-	59,59,59,59	0
59	MG	1A	3338	1/1	0.85	0.40	-	42,42,42,42	0
59	MG	2A	3363	1/1	0.90	0.46	-	47,47,47,47	0
59	MG	1A	4155	1/1	0.90	0.11	-	54,54,54,54	0
59	MG	2A	3620	1/1	0.97	0.16	-	50,50,50,50	0
59	MG	2a	3016	1/1	0.92	0.13	-	46,46,46,46	0
59	MG	1A	3495	1/1	0.85	0.20	-	46,46,46,46	0
59	MG	1A	3734	1/1	0.94	0.10	-	45,45,45,45	0
59	MG	2A	3614	1/1	0.88	0.17	-	63,63,63,63	0
59	MG	2j	8001	1/1	0.93	0.20	-	63,63,63,63	0
59	MG	1A	3541	1/1	0.93	0.17	-	20,20,20,20	0
59	MG	1A	3118	1/1	0.99	0.55	-	33,33,33,33	0
59	MG	1A	3076	1/1	0.93	0.20	-	30,30,30,30	0
59	MG	1A	3968	1/1	0.92	0.15	-	27,27,27,27	0
59	MG	2A	3177	1/1	0.91	0.14	-	63,63,63,63	0
59	MG	2A	3005	1/1	0.75	0.33	-	62,62,62,62	0
59	MG	1a	3364	1/1	0.88	0.30	-	53,53,53,53	0
59	MG	1A	3418	1/1	0.95	0.33	-	22,22,22,22	0
59	MG	1A	3962	1/1	0.91	0.38	-	55,55,55,55	0
59	MG	2a	3168	1/1	0.93	0.17	-	54,54,54,54	0
59	MG	1l	203	1/1	0.90	0.23	-	64,64,64,64	0
59	MG	2a	3206	1/1	0.74	0.39	-	74,74,74,74	0
59	MG	2A	3148	1/1	0.85	0.49	-	65,65,65,65	0
59	MG	2a	3074	1/1	0.92	0.14	-	64,64,64,64	0
59	MG	1A	3947	1/1	0.93	0.10	-	58,58,58,58	0
59	MG	2a	3123	1/1	0.96	0.18	-	55,55,55,55	0
59	MG	2a	3165	1/1	0.63	0.10	-	68,68,68,68	0
59	MG	1A	3904	1/1	0.85	0.13	-	58,58,58,58	0
59	MG	2a	3051	1/1	0.96	0.19	-	55,55,55,55	0
59	MG	1A	4106	1/1	0.88	0.12	-	58,58,58,58	0
59	MG	1A	3486	1/1	0.97	0.46	-	31,31,31,31	0
59	MG	1A	3728	1/1	0.94	0.17	-	16,16,16,16	0
59	MG	1A	3965	1/1	0.83	0.09	-	46,46,46,46	0
59	MG	1A	3979	1/1	0.89	0.10	-	35,35,35,35	0
59	MG	2A	3408	1/1	0.93	0.12	-	56,56,56,56	0
59	MG	1W	201	1/1	0.96	0.30	-	39,39,39,39	0
59	MG	1A	3929	1/1	0.95	0.18	-	52,52,52,52	0
59	MG	1A	3619	1/1	0.95	0.15	-	19,19,19,19	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3455	1/1	0.82	0.16	-	65,65,65,65	0
59	MG	1A	4124	1/1	0.83	0.13	-	56,56,56,56	0
59	MG	2A	3451	1/1	0.94	0.23	-	46,46,46,46	0
59	MG	1A	4010	1/1	0.95	0.08	-	50,50,50,50	0
59	MG	2A	3809	1/1	0.93	0.18	-	55,55,55,55	0
59	MG	1A	3045	1/1	0.89	0.41	-	53,53,53,53	0
59	MG	2A	3232	1/1	0.92	0.15	-	63,63,63,63	0
59	MG	2a	3039	1/1	0.87	0.34	-	55,55,55,55	0
59	MG	2A	3836	1/1	0.85	0.17	-	65,65,65,65	0
59	MG	1A	3900	1/1	0.84	0.14	-	53,53,53,53	0
59	MG	2A	3437	1/1	0.94	0.12	-	48,48,48,48	0
59	MG	1A	4173	1/1	0.93	0.66	-	36,36,36,36	0
59	MG	2A	3456	1/1	0.92	0.09	-	50,50,50,50	0
59	MG	1A	4088	1/1	0.95	0.17	-	42,42,42,42	0
59	MG	1A	3761	1/1	0.90	0.13	-	49,49,49,49	0
59	MG	28	102	1/1	0.82	0.22	-	58,58,58,58	0
59	MG	1A	3542	1/1	0.78	0.10	-	41,41,41,41	0
59	MG	2A	3804	1/1	0.96	0.34	-	59,59,59,59	0
59	MG	2A	3361	1/1	0.88	0.30	-	66,66,66,66	0
59	MG	2A	3215	1/1	0.89	0.14	-	49,49,49,49	0
59	MG	1A	3202	1/1	0.93	0.24	-	43,43,43,43	0
59	MG	2a	3098	1/1	0.58	0.14	-	69,69,69,69	0
59	MG	2A	3164	1/1	0.90	0.14	-	56,56,56,56	0
59	MG	2A	3283	1/1	0.87	0.39	-	60,60,60,60	0
59	MG	1a	3369	1/1	0.86	0.14	-	64,64,64,64	0
59	MG	2B	3017	1/1	0.92	0.14	-	57,57,57,57	0
59	MG	1A	3562	1/1	0.97	0.17	-	18,18,18,18	0
59	MG	1a	3433	1/1	0.95	0.29	-	61,61,61,61	0
59	MG	2A	3199	1/1	0.86	0.34	-	54,54,54,54	0
59	MG	2A	3181	1/1	0.91	0.17	-	57,57,57,57	0
59	MG	2a	3128	1/1	0.93	0.13	-	59,59,59,59	0
59	MG	2A	3504	1/1	0.95	0.12	-	27,27,27,27	0
59	MG	2A	3603	1/1	0.95	0.12	-	49,49,49,49	0
59	MG	1A	4048	1/1	0.91	0.12	-	33,33,33,33	0
59	MG	1w	102	1/1	0.60	0.25	-	63,63,63,63	0
59	MG	2A	3776	1/1	0.78	0.21	-	56,56,56,56	0
59	MG	1A	4085	1/1	0.86	0.14	-	64,64,64,64	0
59	MG	1A	3293	1/1	0.97	0.23	-	49,49,49,49	0
59	MG	1B	222	1/1	0.95	0.23	-	50,50,50,50	0
59	MG	1a	3493	1/1	0.87	0.14	-	51,51,51,51	0
59	MG	1x	105	1/1	0.93	0.20	-	47,47,47,47	0
59	MG	1A	4083	1/1	0.44	0.15	-	68,68,68,68	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	4000	1/1	0.88	0.17	-	47,47,47,47	0
59	MG	2P	201	1/1	0.93	0.21	-	51,51,51,51	0
59	MG	2A	3637	1/1	0.95	0.14	-	61,61,61,61	0
59	MG	1a	3405	1/1	0.82	0.15	-	47,47,47,47	0
59	MG	2A	3237	1/1	0.93	0.30	-	59,59,59,59	0
59	MG	2A	3115	1/1	0.93	0.65	-	36,36,36,36	0
59	MG	2A	3268	1/1	0.91	0.21	-	61,61,61,61	0
59	MG	1A	3941	1/1	0.96	0.16	-	30,30,30,30	0
59	MG	1A	4118	1/1	0.89	0.15	-	69,69,69,69	0
59	MG	1a	3481	1/1	0.98	0.12	-	44,44,44,44	0
59	MG	2A	3235	1/1	0.69	0.20	-	62,62,62,62	0
59	MG	1a	3384	1/1	0.94	0.24	-	46,46,46,46	0
59	MG	1A	3946	1/1	0.98	0.08	-	46,46,46,46	0
59	MG	1A	3660	1/1	0.95	0.11	-	40,40,40,40	0
59	MG	1A	3073	1/1	0.96	0.28	-	26,26,26,26	0
59	MG	1A	3760	1/1	0.95	0.23	-	32,32,32,32	0
59	MG	1A	3440	1/1	0.92	0.12	-	70,70,70,70	0
59	MG	1B	235	1/1	0.93	0.26	-	63,63,63,63	0
59	MG	1A	3672	1/1	0.88	0.11	-	42,42,42,42	0
59	MG	2A	3595	1/1	0.97	0.10	-	34,34,34,34	0
59	MG	2B	3003	1/1	0.70	0.21	-	71,71,71,71	0
59	MG	2A	3706	1/1	0.82	0.11	-	70,70,70,70	0
59	MG	1A	3335	1/1	0.85	0.17	-	46,46,46,46	0
59	MG	A	8001	1/1	0.93	0.25	-	57,57,57,57	0
59	MG	2a	3015	1/1	0.98	0.18	-	66,66,66,66	0
59	MG	1a	3389	1/1	0.91	0.12	-	48,48,48,48	0
59	MG	2A	3074	1/1	0.82	0.15	-	49,49,49,49	0
59	MG	1B	209	1/1	0.88	0.09	-	58,58,58,58	0
59	MG	1A	3208	1/1	0.91	0.53	-	33,33,33,33	0
59	MG	1a	3428	1/1	0.90	0.11	-	56,56,56,56	0
59	MG	1A	3261	1/1	0.73	0.19	-	48,48,48,48	0
59	MG	1A	3627	1/1	0.96	0.14	-	28,28,28,28	0
59	MG	1w	110	1/1	0.47	0.14	-	81,81,81,81	0
59	MG	1A	4205	1/1	0.81	0.42	-	45,45,45,45	0
59	MG	1A	3236	1/1	0.95	0.41	-	34,34,34,34	0
59	MG	16	104	1/1	0.92	0.15	-	43,43,43,43	0
59	MG	1A	4049	1/1	0.93	0.12	-	36,36,36,36	0
59	MG	1A	3921	1/1	0.87	0.17	-	37,37,37,37	0
59	MG	1A	4012	1/1	0.96	0.15	-	53,53,53,53	0
59	MG	2A	3731	1/1	0.89	0.18	-	72,72,72,72	0
59	MG	1A	3768	1/1	0.88	0.13	-	56,56,56,56	0
59	MG	2A	3350	1/1	0.92	0.10	-	57,57,57,57	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3507	1/1	0.83	0.12	-	45,45,45,45	0
59	MG	1B	204	1/1	0.83	0.27	-	48,48,48,48	0
59	MG	1a	3569	1/1	0.85	0.15	-	66,66,66,66	0
59	MG	2A	3545	1/1	0.91	0.16	-	61,61,61,61	0
59	MG	1a	3525	1/1	0.36	0.17	-	76,76,76,76	0
59	MG	1a	3502	1/1	0.97	0.07	-	67,67,67,67	0
59	MG	1a	3395	1/1	0.85	0.20	-	40,40,40,40	0
59	MG	1A	3537	1/1	0.92	0.20	-	49,49,49,49	0
59	MG	1a	3543	1/1	0.86	0.09	-	68,68,68,68	0
59	MG	2A	3220	1/1	0.79	0.19	-	68,68,68,68	0
59	MG	2A	3759	1/1	0.97	0.09	-	52,52,52,52	0
59	MG	1a	3535	1/1	0.89	0.12	-	59,59,59,59	0
59	MG	1a	3397	1/1	0.69	0.20	-	60,60,60,60	0
59	MG	1a	3385	1/1	0.84	0.29	-	57,57,57,57	0
59	MG	1N	3001	1/1	0.94	0.32	-	52,52,52,52	0
59	MG	2A	3683	1/1	0.87	0.14	-	53,53,53,53	0
59	MG	2a	3191	1/1	0.93	0.08	-	55,55,55,55	0
59	MG	2A	3423	1/1	0.86	0.10	-	33,33,33,33	0
59	MG	2A	3174	1/1	0.85	0.21	-	60,60,60,60	0
59	MG	1A	3854	1/1	0.90	0.10	-	52,52,52,52	0
59	MG	2A	3230	1/1	0.94	0.11	-	42,42,42,42	0
59	MG	1A	3512	1/1	0.90	0.22	-	40,40,40,40	0
59	MG	1A	3837	1/1	0.93	0.09	-	48,48,48,48	0
59	MG	2A	3137	1/1	0.86	0.15	-	60,60,60,60	0
59	MG	2a	3091	1/1	0.81	0.13	-	77,77,77,77	0
59	MG	1a	3387	1/1	0.93	0.27	-	40,40,40,40	0
59	MG	2E	301	1/1	0.93	0.14	-	66,66,66,66	0
59	MG	2a	3161	1/1	0.92	0.09	-	72,72,72,72	0
59	MG	2y	3007	1/1	0.86	0.17	-	78,78,78,78	0
59	MG	1A	3567	1/1	0.91	0.17	-	27,27,27,27	0
62	K	2A	3849	1/1	0.98	0.09	-	47,47,47,47	0
59	MG	1x	114	1/1	0.86	0.21	-	58,58,58,58	0
59	MG	1A	3561	1/1	0.94	0.06	-	48,48,48,48	0
59	MG	1A	3967	1/1	0.85	0.22	-	42,42,42,42	0
59	MG	1a	3478	1/1	0.68	0.16	-	68,68,68,68	0
59	MG	1A	3061	1/1	0.98	0.21	-	8,8,8,8	0
59	MG	1a	3380	1/1	0.85	0.10	-	48,48,48,48	0
59	MG	1A	3175	1/1	0.88	0.11	-	44,44,44,44	0
59	MG	2a	3130	1/1	0.82	0.06	-	79,79,79,79	0
59	MG	1A	3560	1/1	0.95	0.09	-	45,45,45,45	0
59	MG	2A	3771	1/1	0.90	0.10	-	67,67,67,67	0
59	MG	1A	3451	1/1	0.87	0.18	-	38,38,38,38	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3303	1/1	0.94	0.40	-	50,50,50,50	0
59	MG	2A	3178	1/1	0.70	0.16	-	50,50,50,50	0
59	MG	1A	4074	1/1	0.85	0.12	-	63,63,63,63	0
59	MG	1A	3423	1/1	0.87	0.15	-	41,41,41,41	0
59	MG	1A	4113	1/1	0.92	0.10	-	35,35,35,35	0
59	MG	1A	3020	1/1	0.92	0.22	-	45,45,45,45	0
59	MG	1a	3319	1/1	0.96	0.10	-	34,34,34,34	0
59	MG	1A	3651	1/1	0.84	0.13	-	37,37,37,37	0
59	MG	1A	3536	1/1	0.86	0.21	-	48,48,48,48	0
59	MG	1A	3114	1/1	0.95	0.34	-	36,36,36,36	0
59	MG	1A	3111	1/1	0.96	0.12	-	56,56,56,56	0
59	MG	2A	3675	1/1	0.72	0.12	-	51,51,51,51	0
59	MG	2A	3352	1/1	0.93	0.09	-	64,64,64,64	0
59	MG	1A	3714	1/1	0.89	0.14	-	10,10,10,10	0
59	MG	2a	3044	1/1	0.92	0.11	-	66,66,66,66	0
59	MG	1A	3322	1/1	0.93	0.73	-	47,47,47,47	0
59	MG	1A	3889	1/1	0.91	0.08	-	22,22,22,22	0
59	MG	1A	3729	1/1	0.92	0.18	-	11,11,11,11	0
59	MG	2A	3381	1/1	0.94	0.17	-	17,17,17,17	0
59	MG	2A	3055	1/1	0.95	0.11	-	24,24,24,24	0
59	MG	2B	3007	1/1	0.93	0.10	-	69,69,69,69	0
59	MG	2A	3653	1/1	0.63	0.23	-	60,60,60,60	0
59	MG	2A	3238	1/1	0.89	0.29	-	53,53,53,53	0
59	MG	1a	3450	1/1	0.94	0.10	-	51,51,51,51	0
59	MG	1a	3363	1/1	0.88	0.23	-	50,50,50,50	0
59	MG	1a	3483	1/1	0.90	0.12	-	58,58,58,58	0
59	MG	1s	3001	1/1	0.85	0.22	-	62,62,62,62	0
59	MG	1A	3603	1/1	0.85	0.13	-	39,39,39,39	0
59	MG	2A	3211	1/1	0.84	0.19	-	61,61,61,61	0
59	MG	2A	3273	1/1	0.62	0.10	-	66,66,66,66	0
59	MG	1B	221	1/1	0.92	0.14	-	52,52,52,52	0
59	MG	1A	3074	1/1	0.85	0.76	-	28,28,28,28	0
59	MG	1A	3071	1/1	0.81	0.71	-	34,34,34,34	0
59	MG	1x	108	1/1	0.83	0.17	-	53,53,53,53	0
59	MG	2A	3169	1/1	0.90	0.15	-	53,53,53,53	0
59	MG	2A	3162	1/1	0.87	0.28	-	56,56,56,56	0
59	MG	1A	3316	1/1	0.91	0.18	-	15,15,15,15	0
59	MG	1A	3781	1/1	0.95	0.17	-	42,42,42,42	0
59	MG	2A	3785	1/1	0.91	0.08	-	53,53,53,53	0
59	MG	1A	4137	1/1	0.90	0.22	-	64,64,64,64	0
59	MG	1A	3580	1/1	0.97	0.11	-	23,23,23,23	0
59	MG	1A	3220	1/1	0.92	0.46	-	25,25,25,25	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2A	3048	1/1	0.88	0.13	-	53,53,53,53	0
59	MG	1x	113	1/1	0.97	0.13	-	61,61,61,61	0
59	MG	1A	3308	1/1	0.95	0.14	-	53,53,53,53	0
59	MG	2A	3627	1/1	0.84	0.14	-	53,53,53,53	0
59	MG	1A	3306	1/1	0.89	0.44	-	47,47,47,47	0
59	MG	1O	3004	1/1	0.94	0.14	-	48,48,48,48	0
59	MG	1A	4153	1/1	0.93	0.08	-	49,49,49,49	0
59	MG	1E	302	1/1	0.91	0.22	-	47,47,47,47	0
59	MG	1A	3455	1/1	0.78	0.17	-	51,51,51,51	0
59	MG	1A	3398	1/1	0.76	0.44	-	49,49,49,49	0
59	MG	1A	3870	1/1	0.95	0.12	-	29,29,29,29	0
59	MG	2A	3795	1/1	0.95	0.12	-	66,66,66,66	0
59	MG	1A	3265	1/1	0.85	0.15	-	51,51,51,51	0
59	MG	2A	3194	1/1	0.91	0.93	-	38,38,38,38	0
59	MG	2A	3223	1/1	0.77	0.23	-	59,59,59,59	0
59	MG	2A	3348	1/1	0.91	0.11	-	66,66,66,66	0
59	MG	2A	3646	1/1	0.89	0.07	-	53,53,53,53	0
59	MG	2A	3290	1/1	0.65	0.23	-	67,67,67,67	0
59	MG	2A	3472	1/1	0.96	0.27	-	50,50,50,50	0
59	MG	1A	3214	1/1	0.87	0.12	-	46,46,46,46	0
59	MG	1A	3971	1/1	0.81	0.10	-	45,45,45,45	0
59	MG	1A	3966	1/1	0.95	0.28	-	44,44,44,44	0
59	MG	2A	3699	1/1	0.91	0.13	-	38,38,38,38	0
59	MG	1a	3553	1/1	0.78	0.14	-	60,60,60,60	0
59	MG	2a	3077	1/1	0.88	0.21	-	57,57,57,57	0
59	MG	1A	4119	1/1	0.99	0.12	-	22,22,22,22	0
59	MG	2A	3848	1/1	0.96	0.12	-	47,47,47,47	0
59	MG	1A	3378	1/1	0.62	0.17	-	62,62,62,62	0
59	MG	1A	3916	1/1	0.88	0.15	-	12,12,12,12	0
59	MG	1a	3561	1/1	0.90	0.17	-	62,62,62,62	0
59	MG	1A	3192	1/1	0.96	0.38	-	27,27,27,27	0
59	MG	2A	3380	1/1	0.83	0.14	-	41,41,41,41	0
59	MG	1a	3434	1/1	0.93	0.14	-	48,48,48,48	0
59	MG	1a	3505	1/1	0.88	0.25	-	56,56,56,56	0
59	MG	1A	3167	1/1	0.90	0.18	-	38,38,38,38	0
59	MG	1a	3311	1/1	0.94	0.16	-	45,45,45,45	0
59	MG	1a	3546	1/1	0.86	0.11	-	46,46,46,46	0
59	MG	2A	3531	1/1	0.65	0.22	-	57,57,57,57	0
59	MG	1A	3708	1/1	0.89	0.19	-	37,37,37,37	0
59	MG	1A	3498	1/1	0.89	0.14	-	51,51,51,51	0
59	MG	1A	3769	1/1	0.83	0.15	-	48,48,48,48	0
59	MG	2O	8002	1/1	0.89	0.17	-	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3253	1/1	0.92	0.17	-	37,37,37,37	0
59	MG	1A	3762	1/1	0.94	0.12	-	48,48,48,48	0
59	MG	2A	3241	1/1	0.69	0.41	-	66,66,66,66	0
59	MG	1A	3901	1/1	0.92	0.07	-	59,59,59,59	0
59	MG	2a	3147	1/1	0.94	0.09	-	59,59,59,59	0
59	MG	2A	3638	1/1	0.92	0.10	-	55,55,55,55	0
59	MG	1A	3643	1/1	0.89	0.12	-	42,42,42,42	0
59	MG	2A	3713	1/1	0.98	0.10	-	29,29,29,29	0
59	MG	1x	102	1/1	0.90	0.23	-	59,59,59,59	0
59	MG	2A	3639	1/1	0.85	0.10	-	47,47,47,47	0
59	MG	2A	3274	1/1	0.96	0.20	-	59,59,59,59	0
59	MG	1A	3048	1/1	0.87	0.20	-	39,39,39,39	0
59	MG	1A	3551	1/1	0.94	0.16	-	29,29,29,29	0
59	MG	1w	104	1/1	0.88	0.11	-	66,66,66,66	0
59	MG	2A	3826	1/1	0.94	0.13	-	54,54,54,54	0
59	MG	1a	3488	1/1	0.95	0.12	-	39,39,39,39	0
59	MG	1A	4038	1/1	0.98	0.21	-	25,25,25,25	0
59	MG	2A	3128	1/1	0.92	0.34	-	50,50,50,50	0
59	MG	2A	3060	1/1	0.71	0.38	-	63,63,63,63	0
59	MG	1a	3556	1/1	0.93	0.11	-	46,46,46,46	0
59	MG	1A	4044	1/1	0.97	0.09	-	54,54,54,54	0
59	MG	2y	3005	1/1	0.38	0.35	-	98,98,98,98	0
59	MG	2A	3360	1/1	0.70	0.18	-	50,50,50,50	0
59	MG	1A	3832	1/1	0.95	0.13	-	47,47,47,47	0
59	MG	2A	3718	1/1	0.79	0.13	-	52,52,52,52	0
59	MG	2A	3289	1/1	0.73	0.17	-	70,70,70,70	0
59	MG	2A	3395	1/1	0.94	0.12	-	28,28,28,28	0
59	MG	1a	3390	1/1	0.94	0.24	-	53,53,53,53	0
59	MG	1A	3225	1/1	0.88	0.16	-	64,64,64,64	0
59	MG	1A	3233	1/1	0.79	0.21	-	51,51,51,51	0
59	MG	2x	101	1/1	0.92	0.20	-	72,72,72,72	0
59	MG	1A	4179	1/1	0.85	0.28	-	60,60,60,60	0
59	MG	1A	3343	1/1	0.97	0.20	-	43,43,43,43	0
59	MG	1A	3583	1/1	0.96	0.10	-	50,50,50,50	0
59	MG	2A	3534	1/1	0.94	0.13	-	58,58,58,58	0
59	MG	2A	3117	1/1	0.83	0.10	-	60,60,60,60	0
59	MG	2a	3158	1/1	0.62	0.31	-	71,71,71,71	0
59	MG	2A	3065	1/1	0.79	0.23	-	47,47,47,47	0
59	MG	2A	3347	1/1	0.79	0.14	-	57,57,57,57	0
59	MG	1y	101	1/1	0.98	0.40	-	32,32,32,32	0
59	MG	1A	3133	1/1	0.92	0.18	-	41,41,41,41	0
59	MG	2A	3564	1/1	0.78	0.28	-	54,54,54,54	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1a	3315	1/1	0.94	0.17	-	48,48,48,48	0
59	MG	1a	3444	1/1	0.95	0.16	-	33,33,33,33	0
59	MG	2a	3190	1/1	0.96	0.17	-	56,56,56,56	0
59	MG	2A	3441	1/1	0.74	0.14	-	29,29,29,29	0
59	MG	1A	4086	1/1	0.86	0.12	-	33,33,33,33	0
59	MG	1A	3433	1/1	0.79	0.21	-	56,56,56,56	0
59	MG	1A	3706	1/1	0.94	0.11	-	28,28,28,28	0
59	MG	2y	3003	1/1	0.90	0.54	-	83,83,83,83	0
59	MG	1A	3279	1/1	0.78	0.17	-	45,45,45,45	0
59	MG	1A	3390	1/1	0.98	0.37	-	43,43,43,43	0
59	MG	2A	3374	1/1	0.95	0.11	-	49,49,49,49	0
59	MG	1A	3386	1/1	0.71	0.23	-	47,47,47,47	0
59	MG	1E	312	1/1	0.86	0.11	-	56,56,56,56	0
59	MG	1A	3485	1/1	0.83	0.56	-	41,41,41,41	0
59	MG	1A	3590	1/1	0.95	0.14	-	40,40,40,40	0
59	MG	2a	3092	1/1	0.90	0.20	-	61,61,61,61	0
59	MG	1A	3272	1/1	0.90	0.14	-	36,36,36,36	0
59	MG	2T	203	1/1	0.89	0.21	-	50,50,50,50	0
59	MG	2A	3092	1/1	0.96	0.14	-	32,32,32,32	0
59	MG	2a	3204	1/1	0.92	0.19	-	62,62,62,62	0
59	MG	2A	3806	1/1	0.85	0.10	-	54,54,54,54	0
59	MG	1A	4128	1/1	0.95	0.39	-	53,53,53,53	0
59	MG	1a	3549	1/1	0.93	0.18	-	67,67,67,67	0
59	MG	1A	3678	1/1	0.93	0.19	-	57,57,57,57	0
59	MG	1A	3749	1/1	0.98	0.09	-	26,26,26,26	0
59	MG	1A	3221	1/1	0.92	0.19	-	40,40,40,40	0
59	MG	1a	3512	1/1	0.95	0.10	-	45,45,45,45	0
59	MG	2A	3218	1/1	0.82	0.12	-	63,63,63,63	0
59	MG	1a	3321	1/1	0.88	0.11	-	50,50,50,50	0
59	MG	2a	3076	1/1	0.92	0.20	-	56,56,56,56	0
59	MG	1A	3416	1/1	0.86	0.19	-	52,52,52,52	0
59	MG	2A	3765	1/1	0.77	0.28	-	70,70,70,70	0
59	MG	2A	3269	1/1	0.83	0.30	-	50,50,50,50	0
59	MG	1a	3559	1/1	0.90	0.19	-	69,69,69,69	0
59	MG	1A	3540	1/1	0.97	0.13	-	47,47,47,47	0
59	MG	1A	3999	1/1	0.95	0.10	-	20,20,20,20	0
59	MG	2a	3201	1/1	0.94	0.19	-	46,46,46,46	0
59	MG	1N	3002	1/1	0.94	0.22	-	44,44,44,44	0
59	MG	1A	3869	1/1	0.86	0.17	-	54,54,54,54	0
59	MG	2A	3180	1/1	0.89	0.13	-	53,53,53,53	0
59	MG	2A	3768	1/1	0.90	0.24	-	58,58,58,58	0
59	MG	1A	3817	1/1	0.93	0.11	-	31,31,31,31	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	1A	3741	1/1	0.79	0.23	-	46,46,46,46	0
59	MG	2a	3035	1/1	0.90	0.41	-	57,57,57,57	0
59	MG	1A	3241	1/1	0.97	0.29	-	24,24,24,24	0
59	MG	17	101	1/1	0.87	0.33	-	49,49,49,49	0
59	MG	1A	3329	1/1	0.90	0.16	-	51,51,51,51	0
59	MG	1A	3581	1/1	0.97	0.16	-	22,22,22,22	0
59	MG	1X	3002	1/1	0.97	0.24	-	28,28,28,28	0
59	MG	1A	4114	1/1	0.88	0.11	-	57,57,57,57	0
59	MG	2A	3183	1/1	0.89	0.14	-	52,52,52,52	0
59	MG	2a	3217	1/1	0.79	0.12	-	58,58,58,58	0
59	MG	2A	3717	1/1	0.93	0.15	-	63,63,63,63	0
59	MG	1A	3986	1/1	0.65	0.09	-	61,61,61,61	0
59	MG	1A	3468	1/1	0.85	0.61	-	41,41,41,41	0
59	MG	2A	3221	1/1	0.82	0.16	-	59,59,59,59	0
59	MG	1A	4031	1/1	0.98	0.08	-	47,47,47,47	0
59	MG	1B	229	1/1	0.81	0.11	-	69,69,69,69	0
59	MG	2a	3124	1/1	0.96	0.10	-	62,62,62,62	0
59	MG	1A	4109	1/1	0.89	0.19	-	42,42,42,42	0
59	MG	2A	3041	1/1	0.88	0.17	-	60,60,60,60	0
59	MG	1A	3657	1/1	0.94	0.10	-	67,67,67,67	0
59	MG	2A	3121	1/1	0.95	0.22	-	49,49,49,49	0
59	MG	1A	3376	1/1	0.93	0.13	-	48,48,48,48	0
59	MG	2A	3477	1/1	0.89	0.20	-	57,57,57,57	0
59	MG	1A	3824	1/1	0.93	0.16	-	35,35,35,35	0
59	MG	2p	3001	1/1	0.74	0.19	-	63,63,63,63	0
59	MG	2A	3773	1/1	0.73	0.15	-	40,40,40,40	0
59	MG	2A	3748	1/1	0.81	0.15	-	55,55,55,55	0
59	MG	2A	3563	1/1	0.97	0.09	-	52,52,52,52	0
59	MG	2A	3635	1/1	0.90	0.10	-	62,62,62,62	0
59	MG	1A	3834	1/1	0.92	0.15	-	34,34,34,34	0
59	MG	1B	201	1/1	0.83	0.18	-	44,44,44,44	0
59	MG	1A	3833	1/1	0.95	0.09	-	36,36,36,36	0
59	MG	2a	3152	1/1	0.86	0.10	-	65,65,65,65	0
59	MG	2A	3001	1/1	0.93	0.31	-	49,49,49,49	0
59	MG	1A	3704	1/1	0.97	0.16	-	13,13,13,13	0
59	MG	1A	3034	1/1	0.93	0.17	-	27,27,27,27	0
59	MG	2A	3277	1/1	0.94	0.26	-	51,51,51,51	0
59	MG	1A	3963	1/1	0.93	0.08	-	37,37,37,37	0
59	MG	1A	3569	1/1	0.93	0.14	-	42,42,42,42	0
59	MG	2A	3829	1/1	0.79	0.67	-	60,60,60,60	0
59	MG	1A	3409	1/1	0.80	0.42	-	51,51,51,51	0
59	MG	1a	3437	1/1	0.91	0.31	-	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors(\AA^2)	Q<0.9
59	MG	2a	3224	1/1	0.93	0.12	-	58,58,58,58	0
59	MG	1A	4213	1/1	0.96	0.34	-	24,24,24,24	0
59	MG	2A	3662	1/1	0.77	0.15	-	69,69,69,69	0
59	MG	1A	3564	1/1	0.91	0.09	-	26,26,26,26	0
59	MG	1A	3174	1/1	0.88	0.16	-	23,23,23,23	0
59	MG	2A	3475	1/1	0.96	0.11	-	48,48,48,48	0
59	MG	2A	3665	1/1	0.90	0.19	-	38,38,38,38	0
59	MG	1A	3362	1/1	0.93	0.30	-	53,53,53,53	0
59	MG	2A	3315	1/1	0.92	0.15	-	63,63,63,63	0
59	MG	1Q	203	1/1	0.94	0.18	-	51,51,51,51	0

6.5 Other polymers [i](#)

There are no such residues in this entry.