



wwPDB X-ray Structure Validation Summary Report ⓘ

Sep 19, 2020 – 12:51 PM BST

PDB ID : 6XQE
Title : Crystal structure of the *Thermus thermophilus* 70S ribosome in complex with sarecycline, UAA-mRNA, and deacylated P-site tRNA at 3.00Å resolution
Authors : Batool, Z.; Lomakin, I.B.; Bunick, C.G.; Polikanov, Y.S.
Deposited on : 2020-07-09
Resolution : 3.00 Å(reported)

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

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

A user guide is available at

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

with specific help available everywhere you see the ⓘ symbol.

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

MolProbity	:	4.02b-467
Mogul	:	1.8.5 (274361), CSD as541be (2020)
Xtriage (Phenix)	:	1.13
EDS	:	2.14.6
buster-report	:	1.1.7 (2018)
Percentile statistics	:	20191225.v01 (using entries in the PDB archive December 25th 2019)
Refmac	:	5.8.0158
CCP4	:	7.0.044 (Gargrove)
Ideal geometry (proteins)	:	Engh & Huber (2001)
Ideal geometry (DNA, RNA)	:	Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP)	:	2.14.6

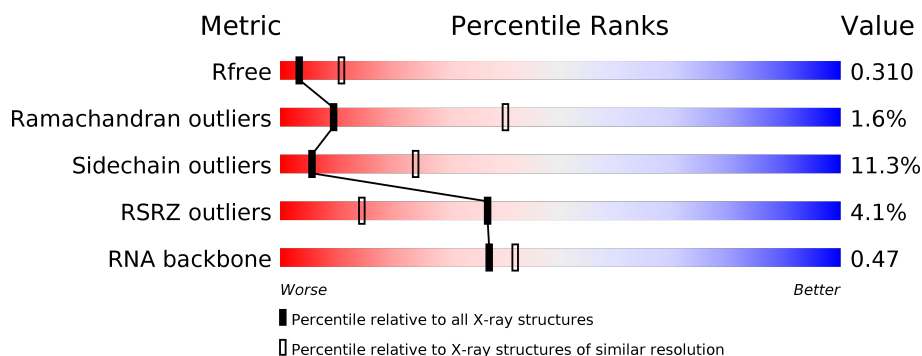
1 Overall quality at a glance ⓘ

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

The reported resolution of this entry is 3.00 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.



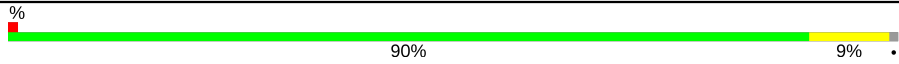

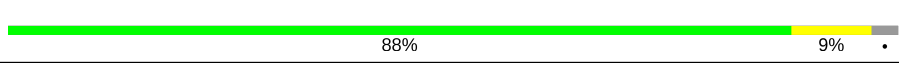

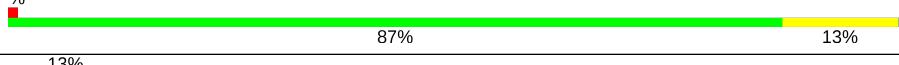
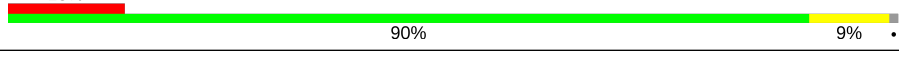
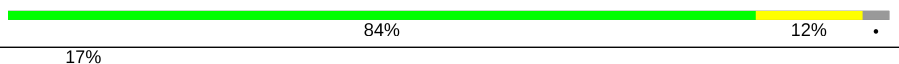

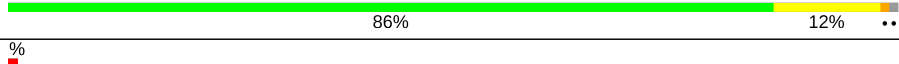

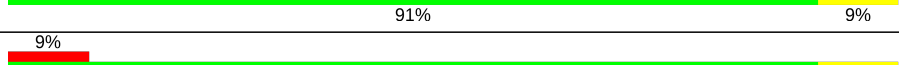
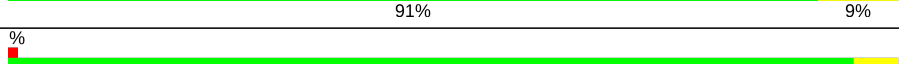
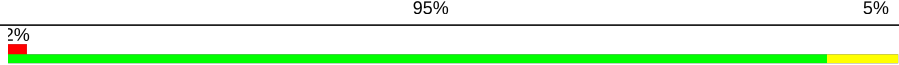
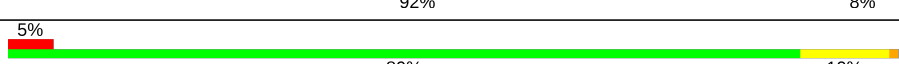
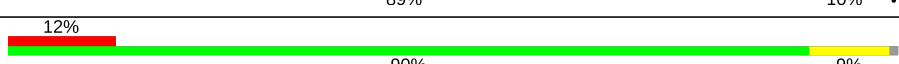
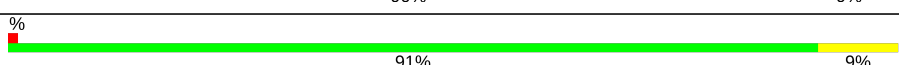
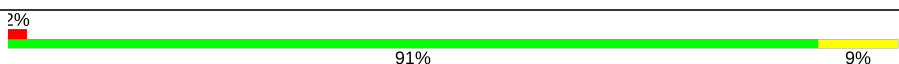
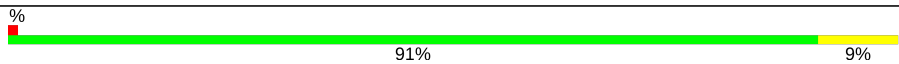
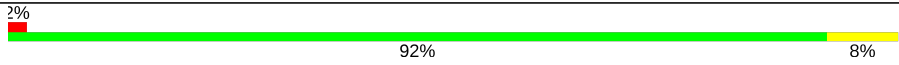

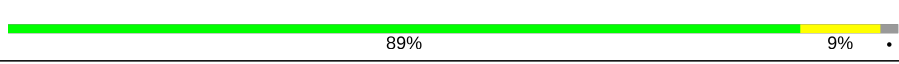
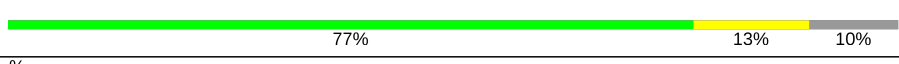
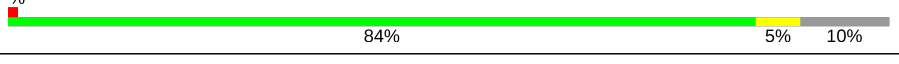
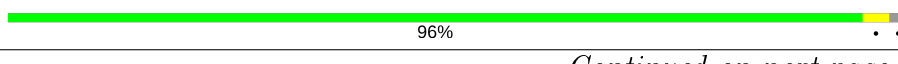

Metric	Whole archive (#Entries)	Similar resolution (#Entries, resolution range(Å))
R_{free}	130704	2092 (3.00-3.00)
Ramachandran outliers	138981	2333 (3.00-3.00)
Sidechain outliers	138945	2336 (3.00-3.00)
RSRZ outliers	127900	1990 (3.00-3.00)
RNA backbone	3102	1173 (3.30-2.70)

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 respectively. A grey segment represents the fraction of residues that are not modelled. The numeric value for each fraction is indicated below the corresponding segment, with a dot representing fractions $\leq 5\%$. The upper red bar (where present) indicates the fraction of residues that have poor fit to the electron density. The numeric value is given above the bar.

Mol	Chain	Length	Quality of chain
1	1A	2915	<div> <div>78%</div> <div>20%</div> <div>..</div> </div>
1	2A	2915	<div> <div>75%</div> <div>21%</div> <div>.</div> </div>
2	1B	121	<div> <div>91%</div> <div>7%</div> <div>..</div> </div>
2	2B	121	<div> <div>71%</div> <div>28%</div> <div>.</div> </div>
3	1D	276	<div> <div>91%</div> <div>9%</div> </div>
3	2D	276	<div> <div>%</div> <div>93%</div> <div>7%</div> </div>

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Mol	Chain	Length	Quality of chain
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	
15	2T	146	
16	1U	118	

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Mol	Chain	Length	Quality of chain
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	
28	16	54	
28	26	54	

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Mol	Chain	Length	Quality of chain
29	17	49	
29	27	49	
30	18	65	
30	28	65	
31	19	37	
31	29	37	
32	1a	1521	
32	2a	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	
41	1j	105	

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Mol	Chain	Length	Quality of chain
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	
53	2v	24	

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Mol	Chain	Length	Quality of chain
54	1x	77	
54	2x	77	

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

Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
55	MG	1A	3304	-	-	-	X
55	MG	1A	3312	-	-	-	X
55	MG	1A	3370	-	-	-	X
55	MG	1A	3842	-	-	-	X
55	MG	1O	203	-	-	-	X
55	MG	1Q	205	-	-	-	X
55	MG	1a	3061	-	-	-	X
55	MG	1a	3172	-	-	-	X
55	MG	1x	102	-	-	-	X
55	MG	2A	3195	-	-	-	X
55	MG	2A	3232	-	-	-	X
55	MG	2A	3278	-	-	-	X
55	MG	2A	3294	-	-	-	X
55	MG	2A	3539	-	-	-	X
55	MG	2a	1790	-	-	-	X
55	MG	2a	1806	-	-	-	X

2 Entry composition [i](#)

There are 60 unique types of molecules in this entry. The entry contains 292071 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	0	0	0
			1091	680	225	185			
15	2T	131	Total	C	N	O	0	0	0
			1083	675	224	183			

- 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			
32	2a	1503	Total	C	N	O	P	0	0	0
			32327	14396	5990	10438	1503			

- 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			
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			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
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	122	Total	C	N	O	S	0	0	0
			951	587	197	165	2			
44	2m	121	Total	C	N	O	S	0	0	0
			943	581	196	164	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	10	Total	C	N	O	P	0	0	0
			217	98	44	65	10			
53	2v	6	Total	C	N	O	P	0	0	0
			113	49	22	36	6			

- Molecule 54 is a RNA chain called P-site tRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
54	1x	76	Total	C	N	O	P	S	0	0
			1625	725	294	529	76	1		
54	2x	76	Total	C	N	O	P	S	0	0
			1625	725	294	529	76	1		

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

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
55	2E	5	Total	Mg	0	0
			5	5		
55	17	7	Total	Mg	0	0
			7	7		
55	2d	2	Total	Mg	0	0
			2	2		
55	1T	2	Total	Mg	0	0
			2	2		
55	1N	5	Total	Mg	0	0
			5	5		
55	2r	1	Total	Mg	0	0
			1	1		
55	18	6	Total	Mg	0	0
			6	6		
55	2W	1	Total	Mg	0	0
			1	1		
55	1Y	2	Total	Mg	0	0
			2	2		
55	13	5	Total	Mg	0	0
			5	5		
55	1f	2	Total	Mg	0	0
			2	2		
55	1P	6	Total	Mg	0	0
			6	6		
55	2B	20	Total	Mg	0	0
			20	20		
55	2a	216	Total	Mg	0	0
			216	216		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
55	1E	14	Total 14	Mg 14	0	0
55	1b	1	Total 1	Mg 1	0	0
55	2l	2	Total 2	Mg 2	0	0
55	2F	6	Total 6	Mg 6	0	0
55	16	3	Total 3	Mg 3	0	0
55	28	3	Total 3	Mg 3	0	0
55	2e	2	Total 2	Mg 2	0	0
55	1W	9	Total 9	Mg 9	0	0
55	1A	1040	Total 1040	Mg 1040	0	0
55	1t	1	Total 1	Mg 1	0	0
55	1n	3	Total 3	Mg 3	0	0
55	2P	4	Total 4	Mg 4	0	0
55	1X	4	Total 4	Mg 4	0	0
55	2q	2	Total 2	Mg 2	0	0
55	12	2	Total 2	Mg 2	0	0
55	2i	1	Total 1	Mg 1	0	0
55	1S	3	Total 3	Mg 3	0	0
55	25	4	Total 4	Mg 4	0	0
55	2T	4	Total 4	Mg 4	0	0
55	1D	10	Total 10	Mg 10	0	0
55	23	3	Total 3	Mg 3	0	0

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Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
55	1e	1	Total 1 Mg 1	0	0
55	2G	1	Total 1 Mg 1	0	0
55	1I	1	Total 1 Mg 1	0	0
55	2f	2	Total 2 Mg 2	0	0
55	1V	4	Total 4 Mg 4	0	0
55	2X	1	Total 1 Mg 1	0	0
55	1a	212	Total 212 Mg 212	0	0
55	2Q	4	Total 4 Mg 4	0	0
55	15	4	Total 4 Mg 4	0	0
55	1x	13	Total 13 Mg 13	0	0
55	2j	2	Total 2 Mg 2	0	0
55	1R	7	Total 7 Mg 7	0	0
55	26	1	Total 1 Mg 1	0	0
55	2U	2	Total 2 Mg 2	0	0
55	1G	4	Total 4 Mg 4	0	0
55	2O	1	Total 1 Mg 1	0	0
55	1I	3	Total 3 Mg 3	0	0
55	1d	1	Total 1 Mg 1	0	0
55	2n	1	Total 1 Mg 1	0	0
55	1H	2	Total 2 Mg 2	0	0
55	2I	6	Total 6 Mg 6	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
55	2g	1	Total 1	Mg 1	0	0
55	2Y	2	Total 2	Mg 2	0	0
55	1v	5	Total 5	Mg 5	0	0
55	2x	7	Total 7	Mg 7	0	0
55	2R	1	Total 1	Mg 1	0	0
55	1Z	3	Total 3	Mg 3	0	0
55	2D	8	Total 8	Mg 8	0	0
55	14	1	Total 1	Mg 1	0	0
55	2k	1	Total 1	Mg 1	0	0
55	1U	8	Total 8	Mg 8	0	0
55	1O	5	Total 5	Mg 5	0	0
55	27	2	Total 2	Mg 2	0	0
55	19	2	Total 2	Mg 2	0	0
55	1l	3	Total 3	Mg 3	0	0
55	2V	1	Total 1	Mg 1	0	0
55	1F	13	Total 13	Mg 13	0	0
55	10	7	Total 7	Mg 7	0	0
55	2t	1	Total 1	Mg 1	0	0
55	1Q	5	Total 5	Mg 5	0	0
55	2A	850	Total 850	Mg 850	0	0
55	2Z	2	Total 2	Mg 2	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
55	1B	36	Total	Mg	0	0
			36	36		

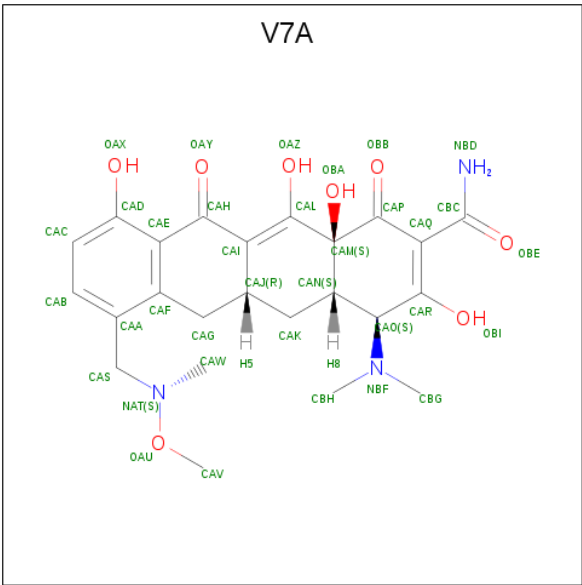
- Molecule 56 is POTASSIUM ION (three-letter code: K) (formula: K).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	1A	1	Total	K	0	0
			1	1		
56	2A	1	Total	K	0	0
			1	1		

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

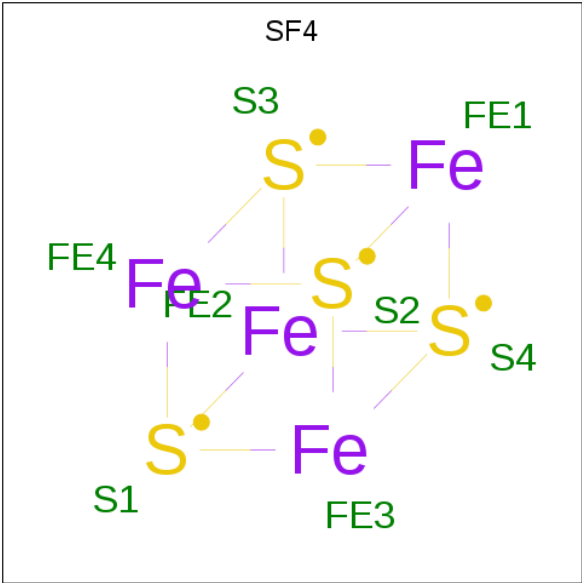
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	1Y	1	Total	Zn	0	0
			1	1		
57	14	1	Total	Zn	0	0
			1	1		
57	1n	1	Total	Zn	0	0
			1	1		
57	15	1	Total	Zn	0	0
			1	1		
57	29	1	Total	Zn	0	0
			1	1		
57	19	1	Total	Zn	0	0
			1	1		
57	26	1	Total	Zn	0	0
			1	1		
57	25	1	Total	Zn	0	0
			1	1		
57	24	1	Total	Zn	0	0
			1	1		
57	2n	1	Total	Zn	0	0
			1	1		
57	2Y	1	Total	Zn	0	0
			1	1		
57	16	1	Total	Zn	0	0
			1	1		

- Molecule 58 is Sarecycline (three-letter code: V7A) (formula: C₂₄H₂₉N₃O₈).



Mol	Chain	Residues	Atoms				ZeroOcc	AltConf
58	1a	1	Total	C	N	O	0	0
			35	24	3	8		
58	2a	1	Total	C	N	O	0	0
			35	24	3	8		

- Molecule 59 is IRON/SULFUR CLUSTER (three-letter code: SF4) (formula: Fe₄S₄).



Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
59	1d	1	Total	Fe	S	0	0
			8	4	4		
59	2d	1	Total	Fe	S	0	0
			8	4	4		

- Molecule 60 is water.

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
60	1A	1402	Total 1402	O 1402	0	0
60	1B	54	Total 54	O 54	0	0
60	1D	21	Total 21	O 21	0	0
60	1E	18	Total 18	O 18	0	0
60	1F	9	Total 9	O 9	0	0
60	1G	4	Total 4	O 4	0	0
60	1H	1	Total 1	O 1	0	0
60	1I	1	Total 1	O 1	0	0
60	1N	5	Total 5	O 5	0	0
60	1O	8	Total 8	O 8	0	0
60	1P	18	Total 18	O 18	0	0
60	1Q	7	Total 7	O 7	0	0
60	1R	9	Total 9	O 9	0	0
60	1S	4	Total 4	O 4	0	0
60	1T	4	Total 4	O 4	0	0
60	1U	14	Total 14	O 14	0	0
60	1V	10	Total 10	O 10	0	0
60	1W	8	Total 8	O 8	0	0
60	1X	4	Total 4	O 4	0	0
60	1Y	4	Total 4	O 4	0	0
60	1Z	1	Total 1	O 1	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
60	10	11	Total 11	O 11	0	0
60	11	8	Total 8	O 8	0	0
60	12	3	Total 3	O 3	0	0
60	13	3	Total 3	O 3	0	0
60	14	1	Total 1	O 1	0	0
60	15	6	Total 6	O 6	0	0
60	16	4	Total 4	O 4	0	0
60	17	10	Total 10	O 10	0	0
60	18	8	Total 8	O 8	0	0
60	1a	211	Total 211	O 211	0	0
60	1b	1	Total 1	O 1	0	0
60	1d	2	Total 2	O 2	0	0
60	1e	1	Total 1	O 1	0	0
60	1i	1	Total 1	O 1	0	0
60	1l	4	Total 4	O 4	0	0
60	1m	1	Total 1	O 1	0	0
60	1q	2	Total 2	O 2	0	0
60	1u	2	Total 2	O 2	0	0
60	1v	3	Total 3	O 3	0	0
60	1x	11	Total 11	O 11	0	0
60	2A	680	Total 680	O 680	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
60	2B	24	Total 24	O 24	0	0
60	2D	12	Total 12	O 12	0	0
60	2E	13	Total 13	O 13	0	0
60	2F	10	Total 10	O 10	0	0
60	2I	1	Total 1	O 1	0	0
60	2O	2	Total 2	O 2	0	0
60	2P	4	Total 4	O 4	0	0
60	2Q	2	Total 2	O 2	0	0
60	2R	3	Total 3	O 3	0	0
60	2T	4	Total 4	O 4	0	0
60	2U	1	Total 1	O 1	0	0
60	2W	2	Total 2	O 2	0	0
60	2X	2	Total 2	O 2	0	0
60	2Y	1	Total 1	O 1	0	0
60	2Z	3	Total 3	O 3	0	0
60	21	10	Total 10	O 10	0	0
60	23	1	Total 1	O 1	0	0
60	27	1	Total 1	O 1	0	0
60	28	2	Total 2	O 2	0	0
60	29	1	Total 1	O 1	0	0
60	2a	173	Total 173	O 173	0	0

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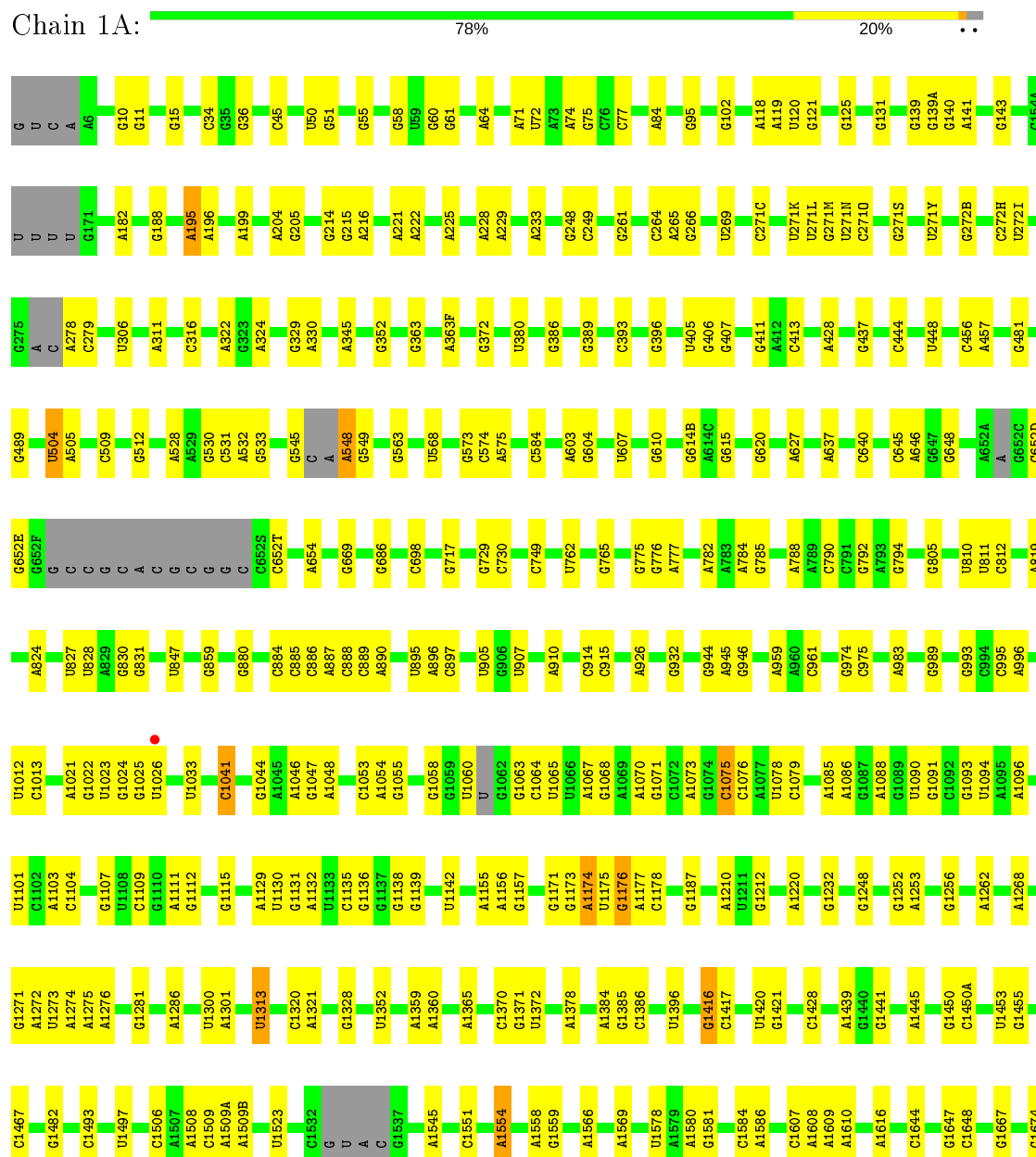
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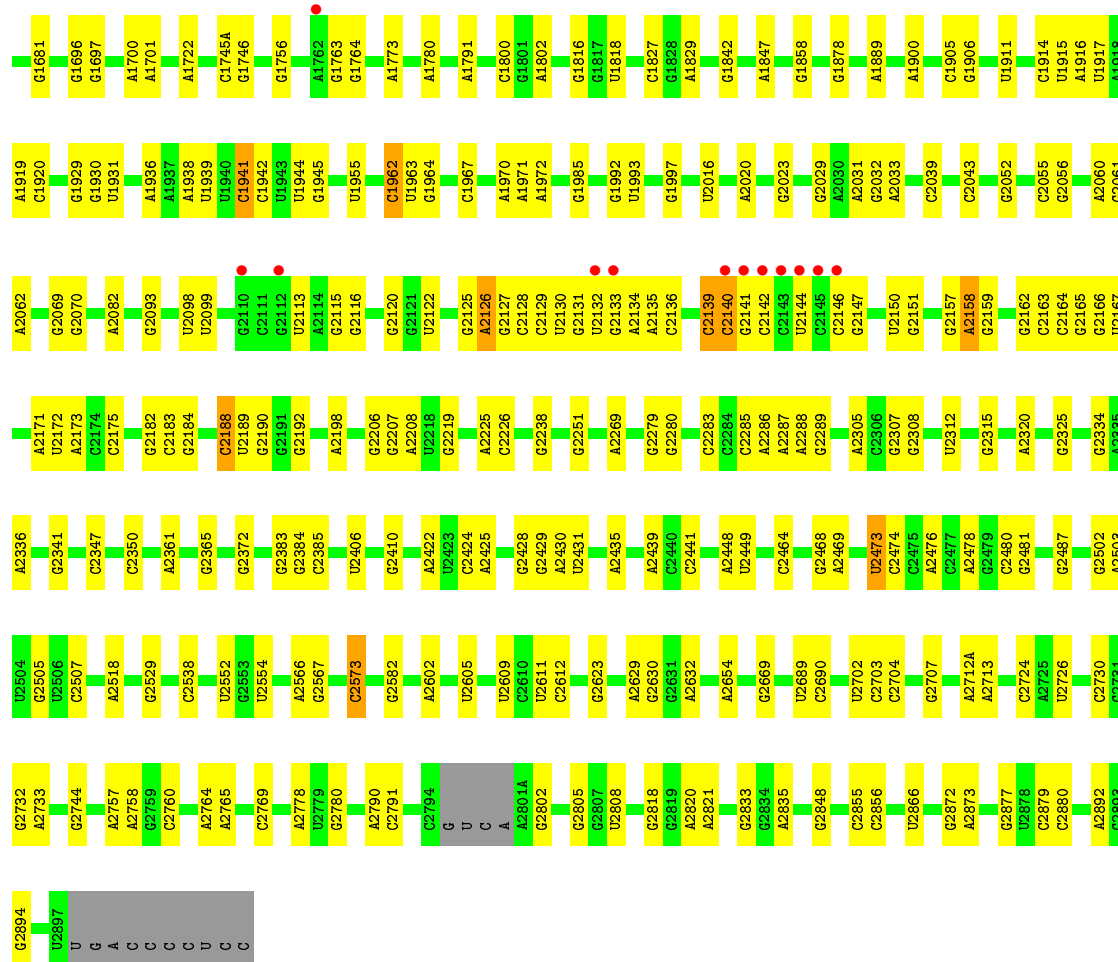
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
60	2e	2	Total	O	0	0
			2	2		
60	2g	1	Total	O	0	0
			1	1		
60	2i	1	Total	O	0	0
			1	1		
60	2j	4	Total	O	0	0
			4	4		
60	2l	3	Total	O	0	0
			3	3		
60	2n	1	Total	O	0	0
			1	1		
60	2t	3	Total	O	0	0
			3	3		
60	2v	1	Total	O	0	0
			1	1		
60	2x	5	Total	O	0	0
			5	5		

3 Residue-property plots

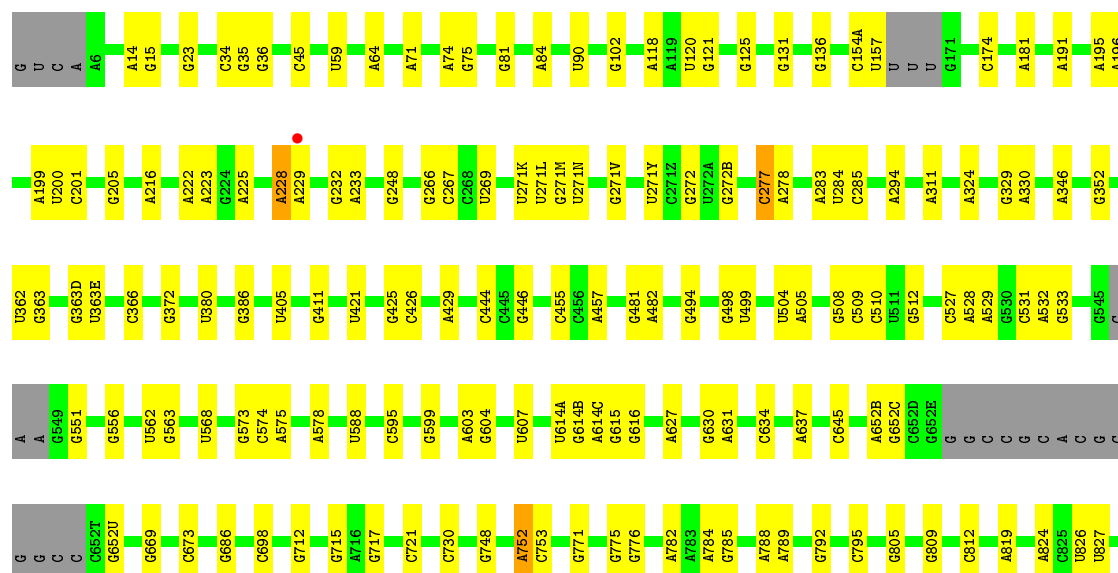
These plots are drawn for all protein, RNA, DNA and oligosaccharide chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and electron density.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

• Molecule 1: 23S Ribosomal RNA





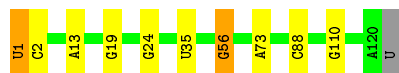
Chain 2A:



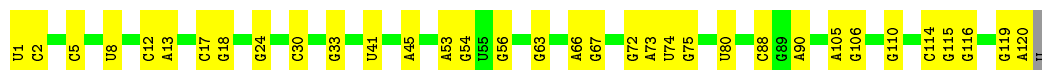
A2764	C2394	C2226	G2121	U1955	U1779	C1588	A1460	G1271	A126	U	A960	G832
A2765	C2395	G2238	U2122	C1962	A1780	G1593	G1461	A1272	A126	U	C961	G843
G2766	G2396	G2239	G2123	U1963	A1791	G1593	C1467	A1273	A1129	G	C964	C857
C2767	C2402	G2251	A2126	G1967	C1800	A1608	A1609	A1275	G1131	U	C965	C856
A2778	U2406	A2274	C2128	A1970	G1801	A1616	A1471	C1298	A1132	U	G968	U858
G2779	G2407	C2275	C2129	A1971	C1804	G1475	G1475	G1299	U1133	A	G972	G859
C2780	U2408	G2279	U2130	A1972	G1816	G1622	G1479	A1301	C1135	G	A973	C865
A	G2410	G2283	G2132	G1982	G1816	G1626	G1479	A1301	G1136	A	G974	A866
C	A2422	C2283	G2133	U1993	U1833	G1626	G1482	G1309	G1139	G	C975	G869
G2782	A2425	A2287	A2135	G1997	U1833	C1638	G1482	U1313	A1142A	A	A980	G874
G2793	A2425	A2288	C2136	U1834	G1835	U1639	A1486	C1314	A1143	G	A981	U877
U	G2429	C2297	C2137	A2001	A1847	A1496	A1496	G1319	G1144	C	A982	A878
C	A2430	G2297	C2138	A1848	A1848	U1497	A1490	G1320	G1151	A	A983	G879
A	A2435	A2305	G2140	G1861	G1861	C1493	C1493	A1321	G1151	U	A996	G880
G2801A	A2440	G2308	G2141	G2021	G1864	A1494	A1494	U1341	A1155	C	C1006	C884
G2802	A2441	G2308	C2142	U2022	U1864	A1496	A1496	G1341	G1164	U	U1012	C885
G2803	A2441	G2308	U2144	G2023	G1866	G1661	U1497	U1341	G1170	U	C1013	C886
G2807	G2446	U2311	C2145	A2031	C1866	C1662	U1503	U1352	G1171	A	U1014	A887
A2820	G2447	G2313	C2146	G2032	G1878	A1664	U1503	U1352	G	A	G1015	C888
A2821	A2448	G2318	G2147	A2033	G1878	A1664	U1503	U1352	A	A	A1020	G892
A2822	U2449	G2318	G2148	C2043	A1890	C1670	A1508	A1359	A	G	A1021	C893
A2823	G2468	G2319	C2155	G2052	G1899	G1674	C1509	A1360	U	A	G1022	C894
A2835	G2468	G2321	G2156	A1900	A1900	C1675	A1509A	A1365	A	G	U1023	A896
C2855	U2473	G2325	G2160	G1906	G1906	A1696	C1530	G1368	C1178	G	G1024	A896
A2868	A2476	A2327	C2161	G2060	G1910	A1700	C1531	G1369	G1192	C	G1025	G897
G2872	G2481	A2333	C2162	A2061	U1911	A1701	U	C1370	A1204	G	U1026	C898
C2879	C2498	A2336	C2163	A2062	A1912	G1702	G1537	A1379	U1205	U	A900	A901
C2880	C2498	G2337	C2164	C2063	A1913	G1703	G1538	G1380	G1206	A	U1033	A910
U2884	G2502	G2340	U2172	U2068	A1914	G1703	G1538	A1384	A1210	A	C1038	A910
C2885	U2503	G2345	G2181	G2069	A1917	G1721	G1541	G1385	U1211	G	G1039	A917
G2886	G2505	A2346	G2182	G2101	U1917	A1722	A1542	C1386	C1221	U	C1040	C925
A2892	U2506	C2347	C2183	C2107	A1928	G1740	A1544	U1406	G1227	C	G1042	G928
G2893	C2507	C2350	C2185	G2109	G1929	G1740	A1545	G1416	G1232	A	G	G932
U2895	A2518	A2366	C2188	U2109	G1930	G1756	A1566	G1421	A1237	G	A	A933
C2896	G2525	A2376	U2189	G2111	A1936	U1757	A1566	A1427	U1240	C	C	G934
U2897	G2529	A2379	G2192	G2112	A1937	U1757	A1569	A1428	A1241	G	A	G938
G	U2554	G2379	G2206	U2113	A1938	A1762	A1569	C1429	A1247	A	G	A941
A	U2554	G2379	G2207	G2115	U1939	G1763	U1578	G1429	G1248	C	C	A945
C	A2566	G2383	A2208	G2116	U1940	G1764	A1579	C1437	U1249	A	A	G946
C	G2567	G2384	A2208	A2117	C1941	A1773	A1580	A1445	A1253	G	G	A953
C	C2568	G2385	U2218	U2118	C1942	A1773	C1584	A1449	G1256	A	A	A959
U	G2569	G2391	A2225	A2119	U1944	G1776	A1586	A1450	G	G	G	

6

- Molecule 2: 5S Ribosomal RNA

Chain 1B:  91% 7% ..

- Molecule 2: 5S Ribosomal RNA

Chain 2B:  71% 28% .

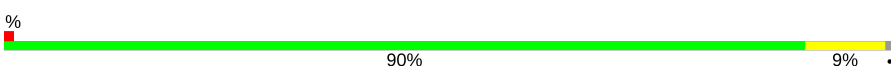
- Molecule 3: 50S ribosomal protein L2

Chain 1D:  91% 9%

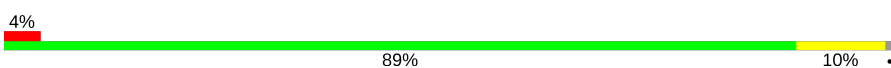
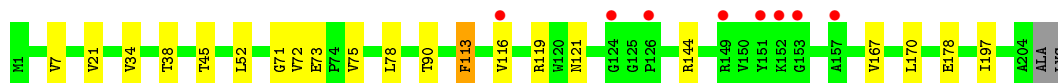
- Molecule 3: 50S ribosomal protein L2

Chain 2D:  93% 7%

- Molecule 4: 50S ribosomal protein L3

Chain 1E:  90% 9% .

- Molecule 4: 50S ribosomal protein L3

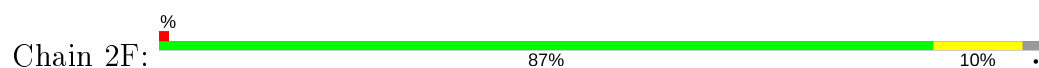
Chain 2E:  89% 10% .

- Molecule 5: 50S ribosomal protein L4

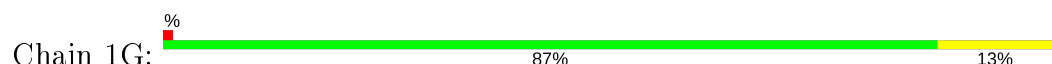
Chain 1F:  88% 9% .



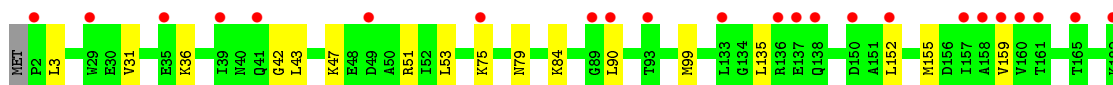
- Molecule 5: 50S ribosomal protein L4



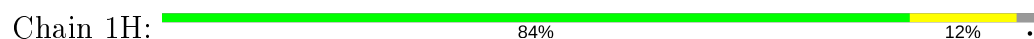
- Molecule 6: 50S ribosomal protein L5



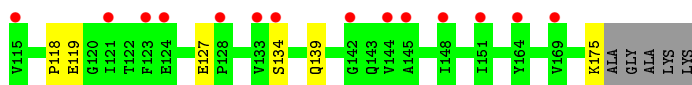
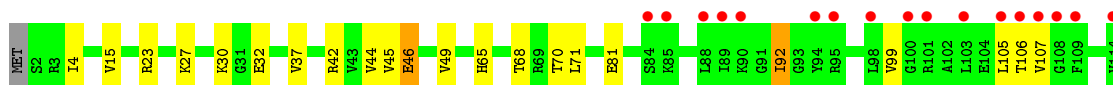
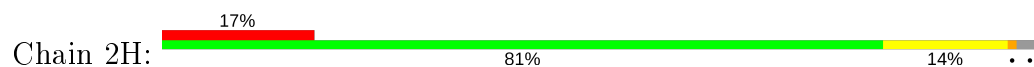
- Molecule 6: 50S ribosomal protein L5



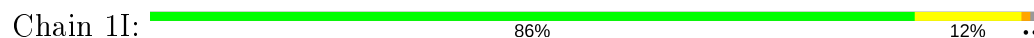
- Molecule 7: 50S ribosomal protein L6



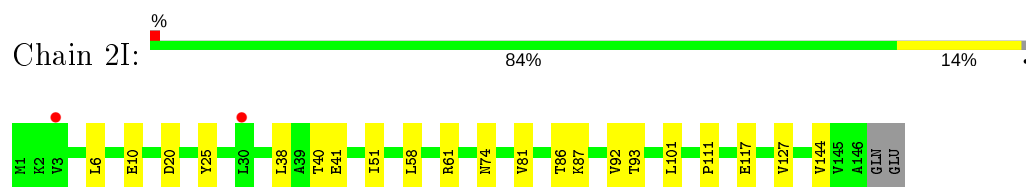
- Molecule 7: 50S ribosomal protein L6



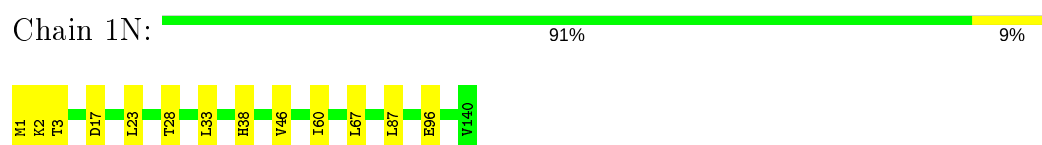
- Molecule 8: 50S ribosomal protein L9



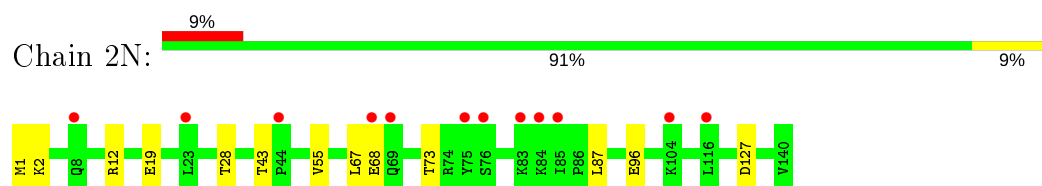
- Molecule 8: 50S ribosomal protein L9



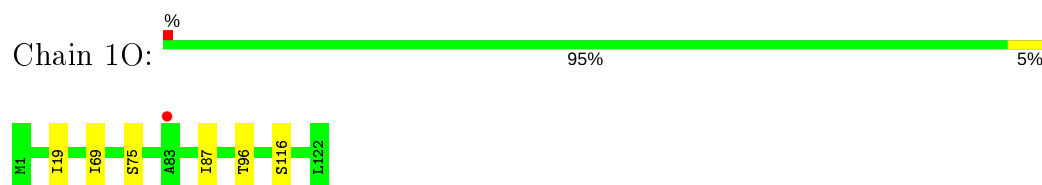
- Molecule 9: 50S ribosomal protein L13



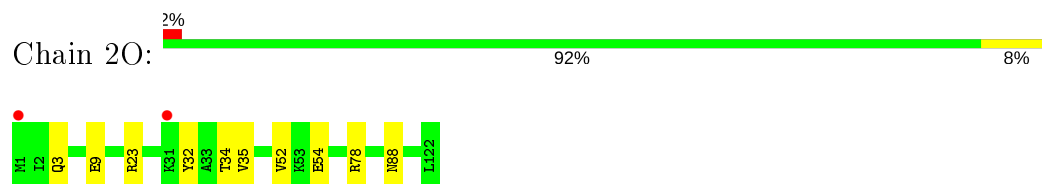
- Molecule 9: 50S ribosomal protein L13



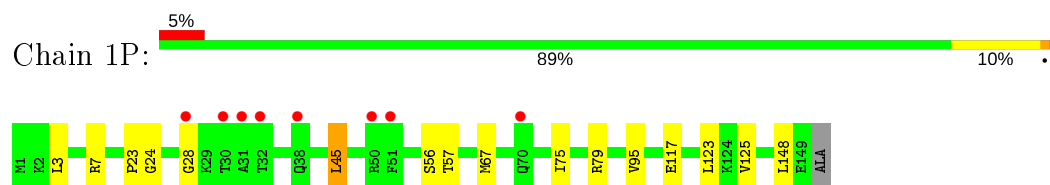
- Molecule 10: 50S ribosomal protein L14



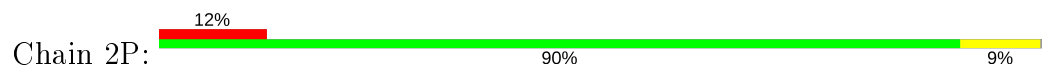
- Molecule 10: 50S ribosomal protein L14

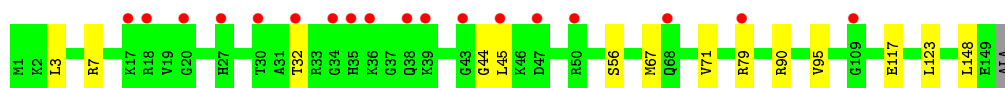


- Molecule 11: 50S ribosomal protein L15



- Molecule 11: 50S ribosomal protein L15





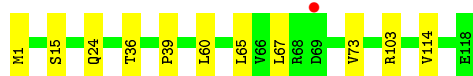
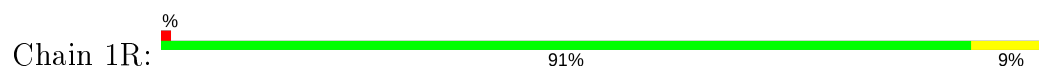
- Molecule 12: 50S ribosomal protein L16



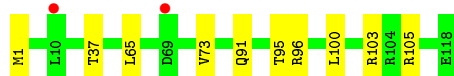
- Molecule 12: 50S ribosomal protein L16



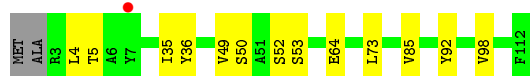
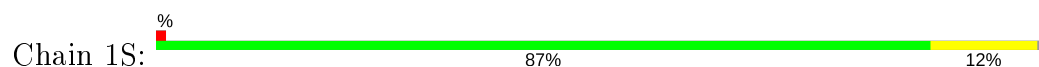
- Molecule 13: 50S ribosomal protein L17



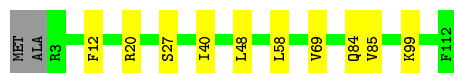
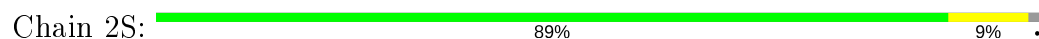
- Molecule 13: 50S ribosomal protein L17




- Molecule 14: 50S ribosomal protein L18

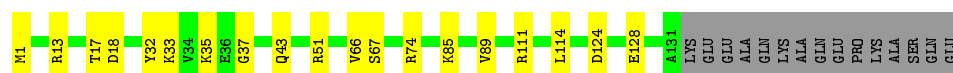


- Molecule 14: 50S ribosomal protein L18




- Molecule 15: 50S ribosomal protein L19

Chain 1T:  77% 13% 10%



- Molecule 15: 50S ribosomal protein L19

Chain 2T:  84% 5% 10%



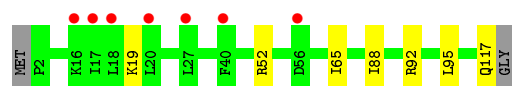
- Molecule 16: 50S ribosomal protein L20

Chain 1U:  96% . .



- Molecule 16: 50S ribosomal protein L20

Chain 2U:  92% 6% .



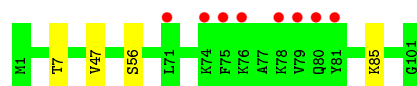
- Molecule 17: 50S ribosomal protein L21

Chain 1V:  91% 9%



- Molecule 17: 50S ribosomal protein L21

Chain 2V:  96% .

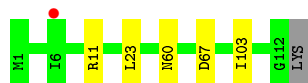


- Molecule 18: 50S ribosomal protein L22

Chain 1W:  93% 6% .



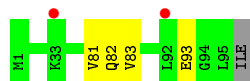
- Molecule 18: 50S ribosomal protein L22



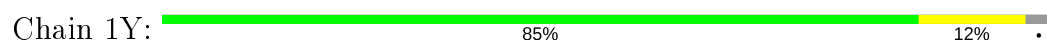
- Molecule 19: 50S ribosomal protein L23



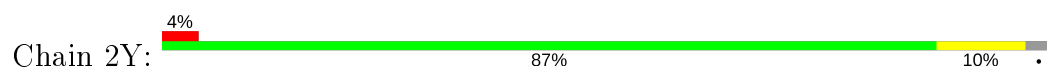
- Molecule 19: 50S ribosomal protein L23



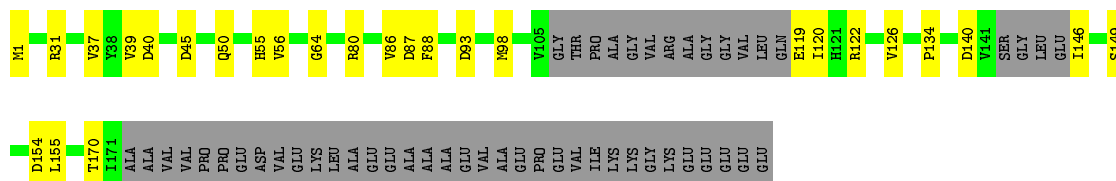
- Molecule 20: 50S ribosomal protein L24



- Molecule 20: 50S ribosomal protein L24

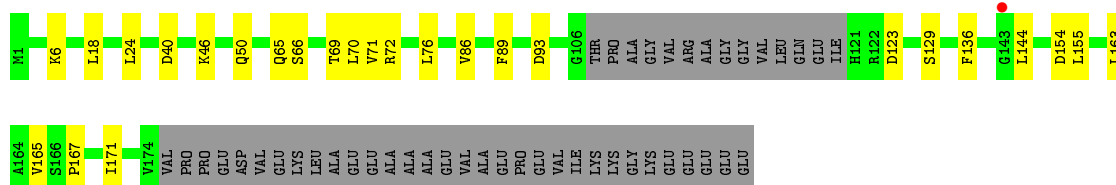


- Molecule 21: 50S ribosomal protein L25

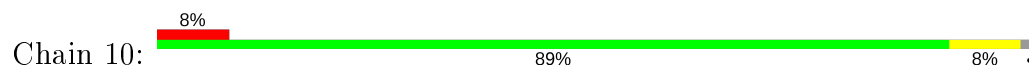


- Molecule 21: 50S ribosomal protein L25

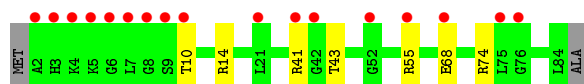
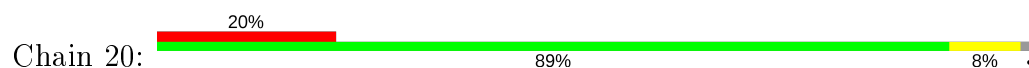




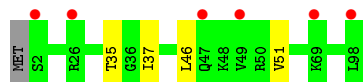
- Molecule 22: 50S ribosomal protein L27



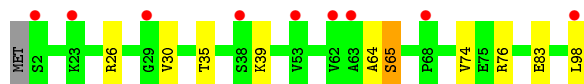
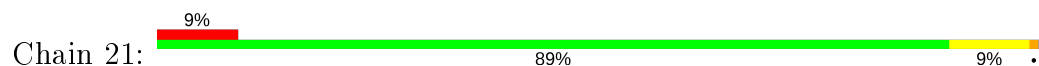
- Molecule 22: 50S ribosomal protein L27



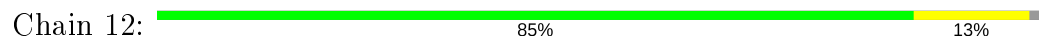
- Molecule 23: 50S ribosomal protein L28



- Molecule 23: 50S ribosomal protein L28




- Molecule 24: 50S ribosomal protein L29



- Molecule 24: 50S ribosomal protein L29




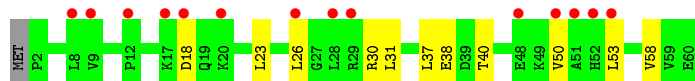
- Molecule 25: 50S ribosomal protein L30

Chain 13:  83% 15%




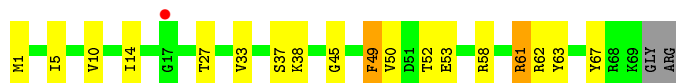
- Molecule 25: 50S ribosomal protein L30

Chain 23:  23% 80% 18%




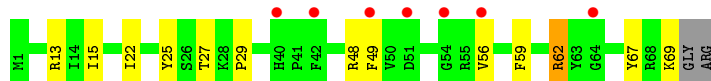
- Molecule 26: 50S ribosomal protein L31

Chain 14:  % 72% 23%




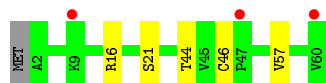
- Molecule 26: 50S ribosomal protein L31

Chain 24:  10% 79% 17%



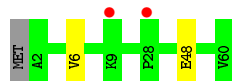
- Molecule 27: 50S ribosomal protein L32

Chain 15:  5% 90% 8%



- Molecule 27: 50S ribosomal protein L32

Chain 25:  3% 95%



- Molecule 28: 50S ribosomal protein L33

Chain 16:  89% 9%



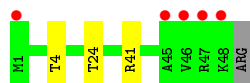
- Molecule 28: 50S ribosomal protein L33

Chain 26:  70% 28% .

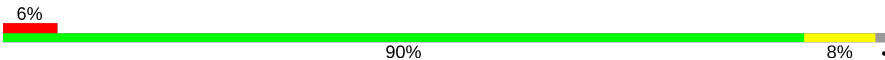


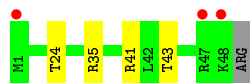
- Molecule 29: 50S ribosomal protein L34

Chain 17:  10% 92% 6% .



- Molecule 29: 50S ribosomal protein L34

Chain 27:  6% 90% 8% .




- Molecule 30: 50S ribosomal protein L35

Chain 18:  8% 94% 5% .



- Molecule 30: 50S ribosomal protein L35

Chain 28:  31% 91% 8% .

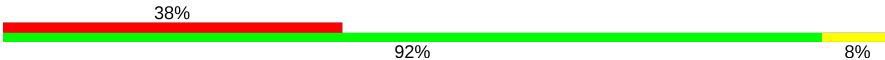


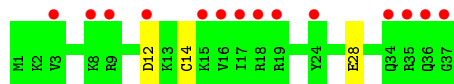
- Molecule 31: 50S ribosomal protein L36

Chain 19:  95% 5% .



- Molecule 31: 50S ribosomal protein L36

Chain 29:  38% 92% 8% .



Chain 1a:

78% 21%

U U G U4 U5 G6 G7 A8 G9 G22 A52 G39 C47 C48 U49 A50 A51 G61 G79 G U U U U A C U90 C91 C92 C93 A101 G105 A116 C121 G124 C131 G138 G139 A143 G144 G147 G156 G157 C163

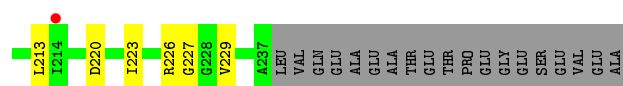
A969 C970 G971 A874 A975 A976 A977 A978 U982 U992 G993 A994 C995 A996 U997 U1000 A1001 G1001A G1002 G1003 G1004 U1005 A1006 C1007 G1008 G1013 U1020 G1021 G1022 G1023 G1024 U1025 G1026 C1027 G1028 C1029 G1030 G1030A C1030B G1030C A1030D G1031 G1032 G1033 G1034 A1035 G1036 A1041 G1042 C1043 A1044 C1045 G1053 C1054 A1055 U1065 C1066 A1067 G1068 G1081 G1084 U1085 A1092 G1093 G1094 U1095 C1096 A1101 U1121 G1124 U1125 G1134 U1135 U1136 C1137 G1138 G1139 A1146 C1149 G1152 C1158 U1159 G1184 G1185 G1186 C1195 U1196 G1197 G1198 U1199 G1202 C1203 G1204 G1207 G1208 A1213 G1224 G1226 A1227 C1236 C1237 A1238 A1239 G1256 U1257 G1258 C1270 G1274 G1275 C1276 C1277 U1278 A1279 A1280 G1286 A1287 G1294 G1299 G1300 U1301 A1302 G1305 G1312 C1320 C1321 C1322 G1323 G1338 G1347 G1363 C1354 G1355 G1356 A1357 C1363 G1365 C1366 C1367 C1368 C1369 G1370 U1381 C1388 C1395 A1396 A1397 A1398 C1399 G1400 G1401 C1402 G1403 C1404 C1407 G1415 G1419 U1436 C1440 G1441 G1442 G1443A A1442B U1446 A1447 C1452 G1455 G1487 A1492 A1493 G1494 U1498 A1503 G1504 G1505 U1506 G1517 G1518 A1519 G1529 G1530 A1531 U1532 C1533 C1534

A969 C970 G971 A874 A975 A976 A977 A978 U982 U992 G993 A994 C995 A996 U997 U1000 A1001 G1001A G1002 G1003 G1004 U1005 A1006 C1007 G1008 G1013 U1020 G1021 G1022 G1023 G1024 U1025 G1026 C1027 G1028 C1029 G1030 G1030A C1030B G1030C A1030D G1031 G1032 G1033 G1034 A1035 G1036 A1041 G1042 C1043 A1044 C1045 G1053 C1054 A1055 U1065 C1066 A1067 G1068 G1081 G1084 U1085 A1092 G1093 G1094 U1095 C1096 A1101 U1121 G1124 U1125 G1134 U1135 U1136 C1137 G1138 G1139 A1146 C1149 G1152 C1158 U1159 G1184 G1185 G1186 C1195 U1196 G1197 G1198 U1199 G1202 C1203 G1204 G1207 G1208 A1213 G1224 G1226 A1227 C1236 C1237 A1238 A1239 G1256 U1257 G1258 C1270 G1274 G1275 C1276 C1277 U1278 A1279 A1280 G1286 A1287 G1294 G1299 G1300 U1301 A1302 G1305 G1312 C1320 C1321 C1322 G1323 G1338 G1347 G1363 C1354 G1355 G1356 A1357 C1363 G1365 C1366 C1367 C1368 C1369 G1370 U1381 C1388 C1395 A1396 A1397 A1398 C1399 G1400 G1401 C1402 G1403 C1404 C1407 G1415 G1419 U1436 C1440 G1441 G1442 G1443A A1442B U1446 A1447 C1452 G1455 G1487 A1492 A1493 G1494 U1498 A1503 G1504 G1505 U1506 G1517 G1518 A1519 G1529 G1530 A1531 U1532 C1533 C1534

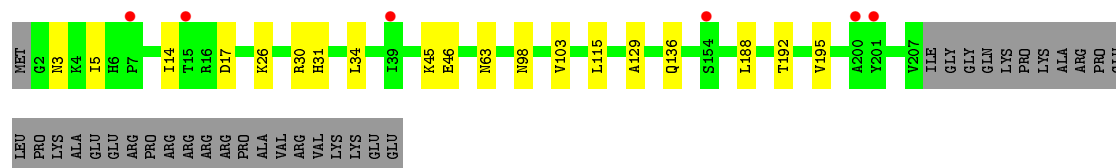
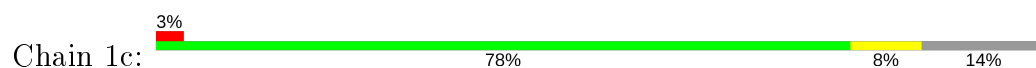
Chain 2a:

2% 75% 23%

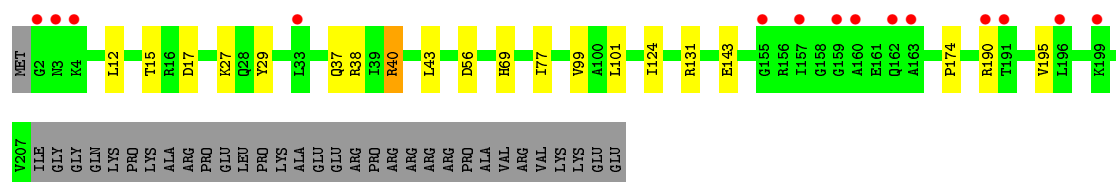
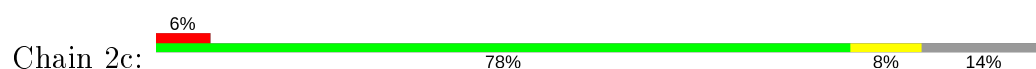
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C47	Yellow
C48	Yellow
A51	Yellow
G52	Yellow
A53	Green
C54	Yellow
A59	Yellow
G66	Yellow
G73	Yellow
G79	Yellow
G80	Yellow
U	Grey
U	Grey
U84	Green
A88	Yellow
C89	Yellow
A101	Yellow
G102	Yellow
G115	Yellow
A116	Yellow
A120	Yellow
C121	Yellow
G129A	Yellow
A130	Yellow
C131	Yellow
C150	Yellow
A151	Yellow
C163	Yellow
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U182	Yellow
C199B	Yellow
C189C	Yellow
C189D	Green
U189E	Yellow
U189F	Yellow
G189G	Yellow
G199H	Yellow
U189K	Yellow
A195	Yellow
A196	Green
A197	Yellow
U202	Yellow
U203	Green
U204	Yellow
G216	Yellow
C217	Yellow
G238	Yellow
U239	Yellow
C245	Yellow
A246	Green
G247	Yellow
A250	Yellow
G251	Yellow
G266	Orange
C267	Yellow
A274	Yellow
G281	Yellow
A288	Yellow
G289	Yellow
G306	Yellow
C307	Yellow
C308	Yellow
A321	Yellow
G324	Yellow
C328	Yellow
G332	Yellow
G348	Yellow
A349	Yellow
G350	Yellow
G351	Yellow
C352	Yellow
A353	Yellow
G354	Yellow



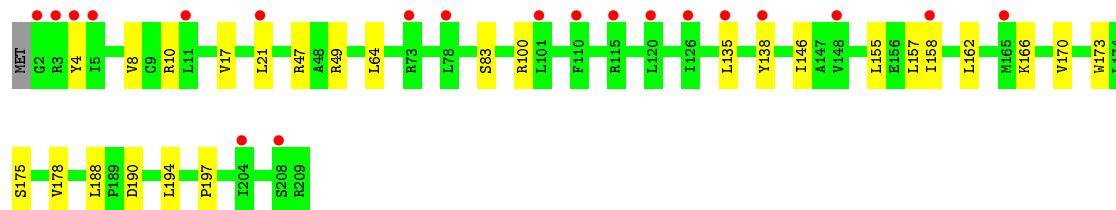
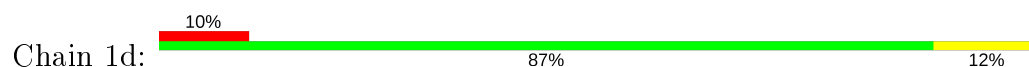
• Molecule 34: 30S ribosomal protein S3



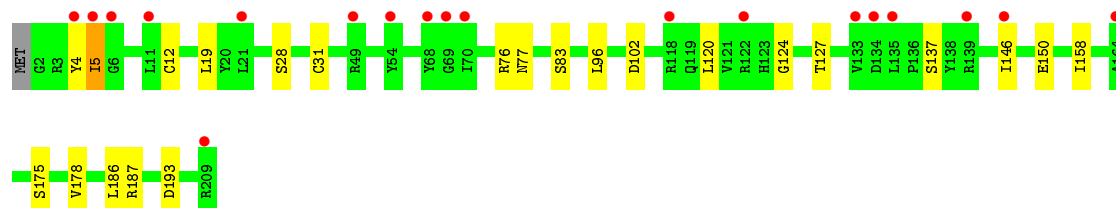
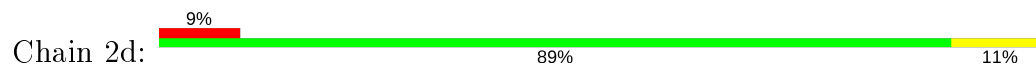
• Molecule 34: 30S ribosomal protein S3



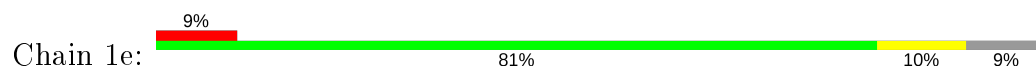
• Molecule 35: 30S ribosomal protein S4

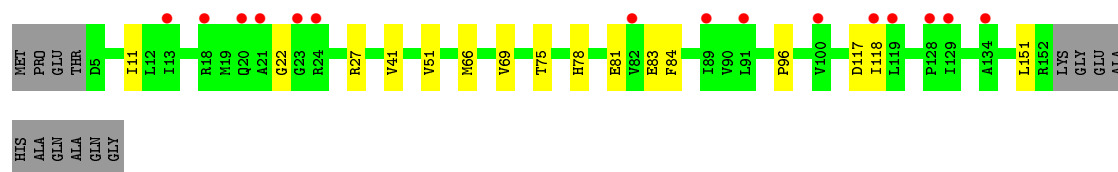


• Molecule 35: 30S ribosomal protein S4

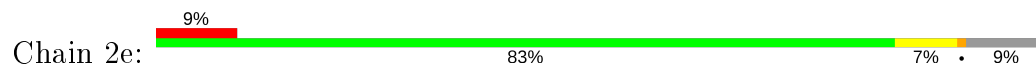


• Molecule 36: 30S ribosomal protein S5





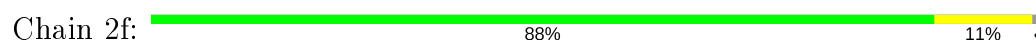
- Molecule 36: 30S ribosomal protein S5



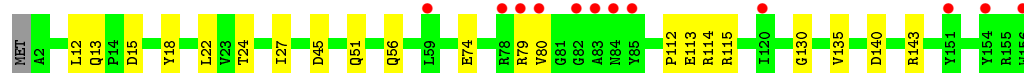
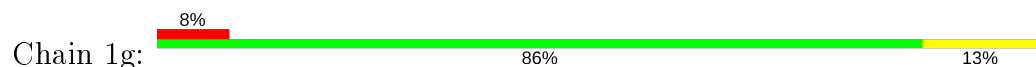
- Molecule 37: 30S ribosomal protein S6



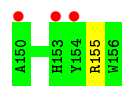
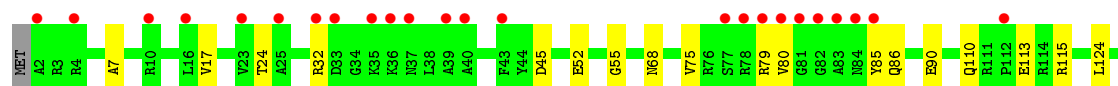
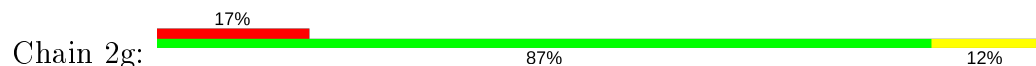
- Molecule 37: 30S ribosomal protein S6



- Molecule 38: 30S ribosomal protein S7



- Molecule 38: 30S ribosomal protein S7

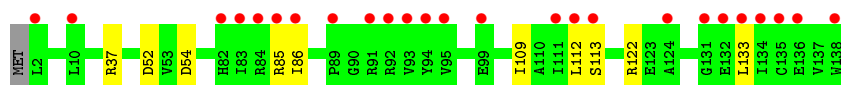


- Molecule 39: 30S ribosomal protein S8

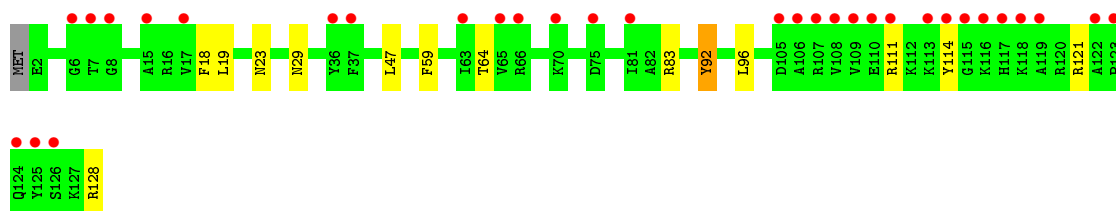
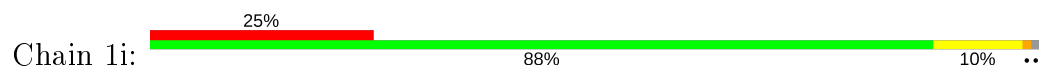




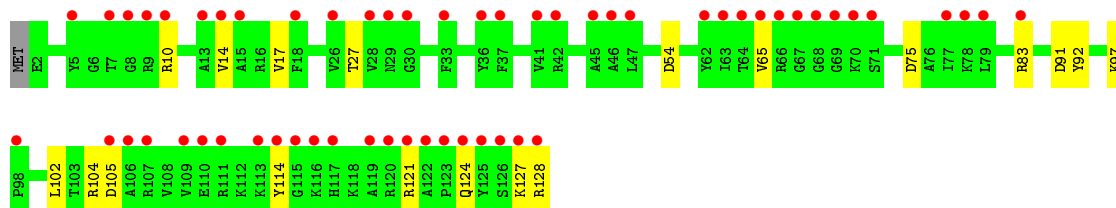
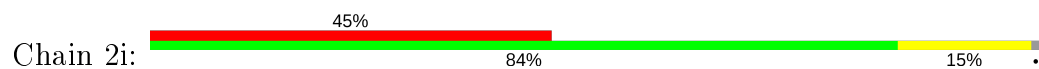
- Molecule 39: 30S ribosomal protein S8



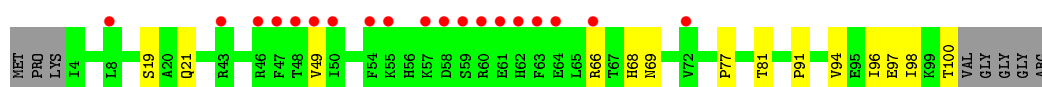
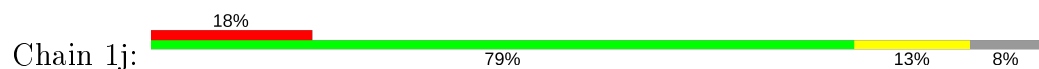
- Molecule 40: 30S ribosomal protein S9



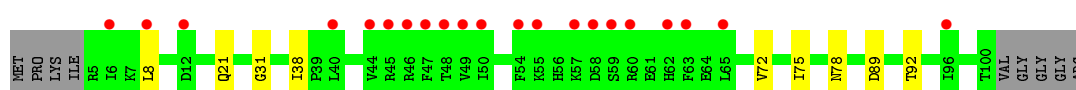
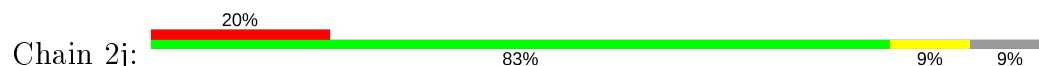
- Molecule 40: 30S ribosomal protein S9



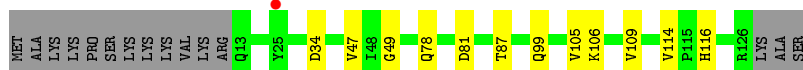
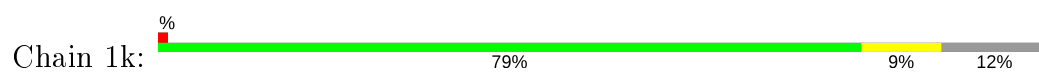
- Molecule 41: 30S ribosomal protein S10



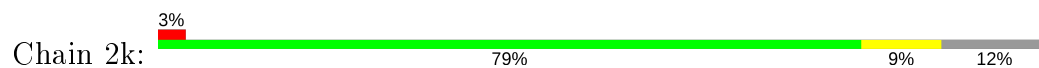
- Molecule 41: 30S ribosomal protein S10



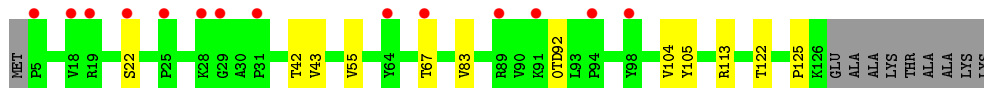
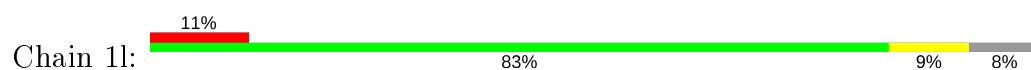
- Molecule 42: 30S ribosomal protein S11



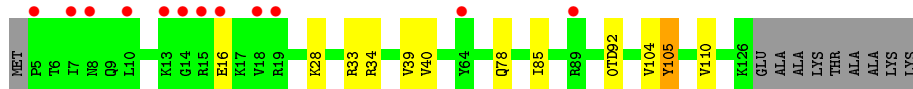
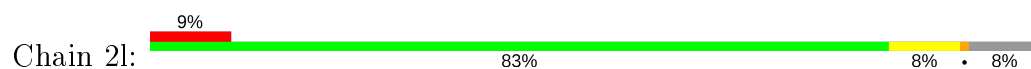
- Molecule 42: 30S ribosomal protein S11



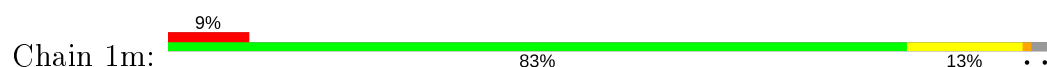
- Molecule 43: 30S ribosomal protein S12



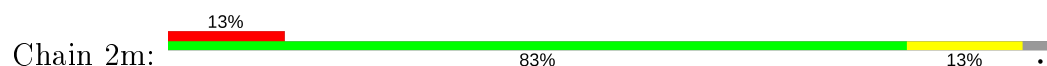
- Molecule 43: 30S ribosomal protein S12



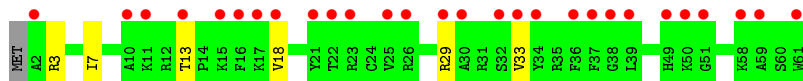
- Molecule 44: 30S ribosomal protein S13



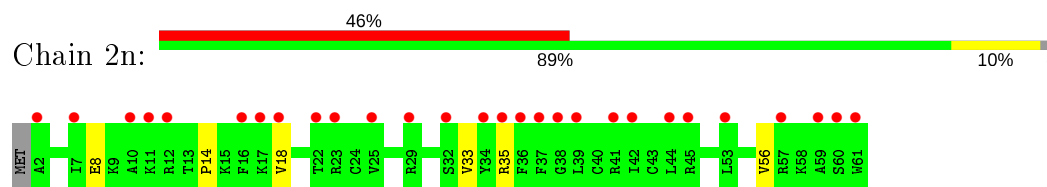
- Molecule 44: 30S ribosomal protein S13



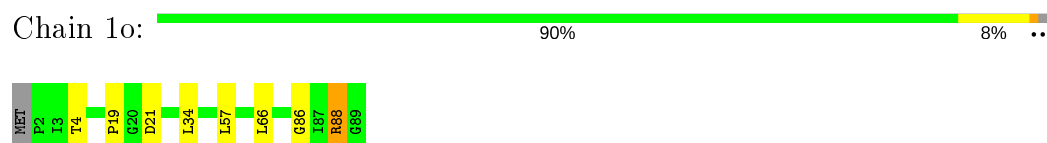
- Molecule 45: 30S ribosomal protein S14 type Z



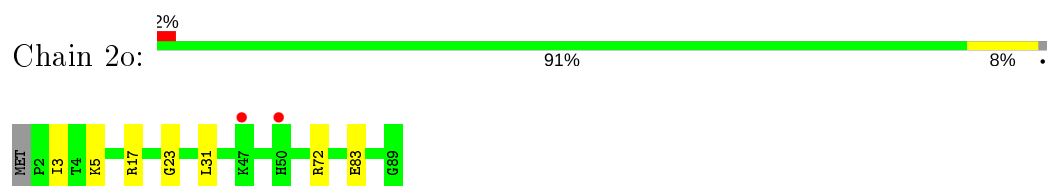
- Molecule 45: 30S ribosomal protein S14 type Z



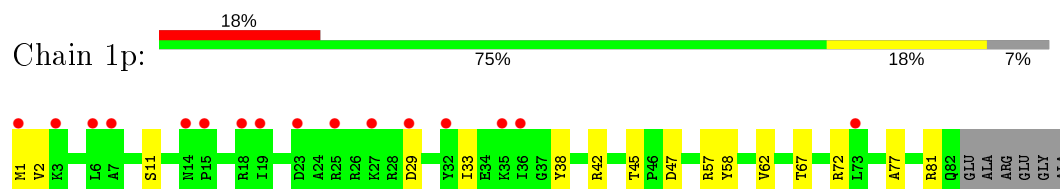
- Molecule 46: 30S ribosomal protein S15



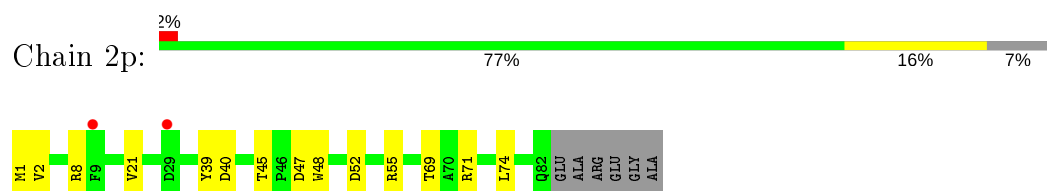
- Molecule 46: 30S ribosomal protein S15



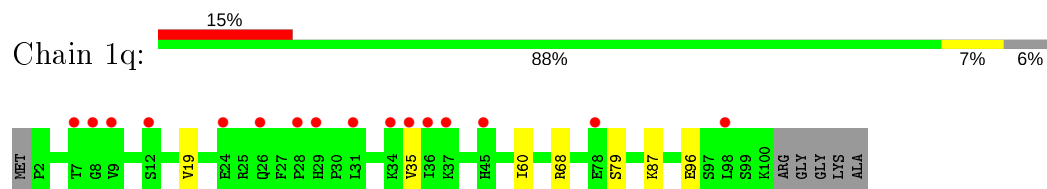
- Molecule 47: 30S ribosomal protein S16



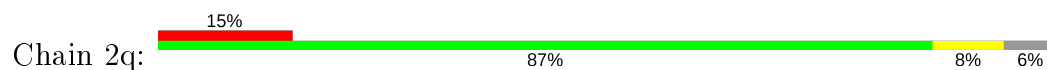
- Molecule 47: 30S ribosomal protein S16



- Molecule 48: 30S ribosomal protein S17

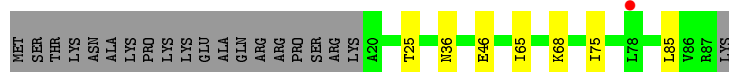


- Molecule 48: 30S ribosomal protein S17

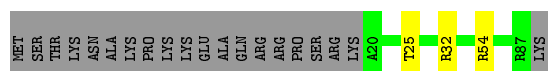




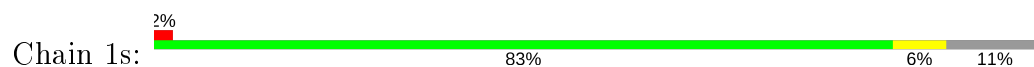
- Molecule 49: 30S ribosomal protein S18



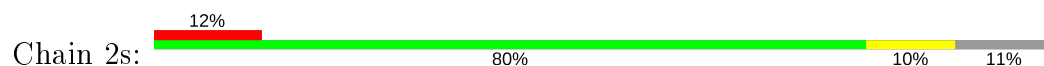
- Molecule 49: 30S ribosomal protein S18



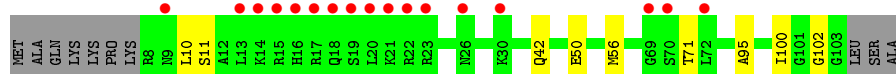
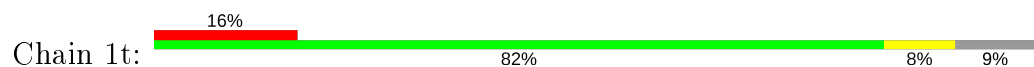
- Molecule 50: 30S ribosomal protein S19



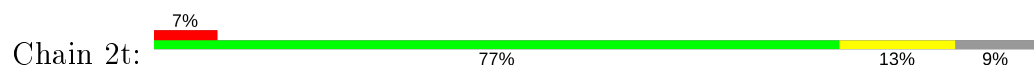
- Molecule 50: 30S ribosomal protein S19



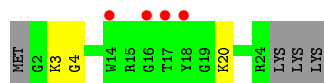
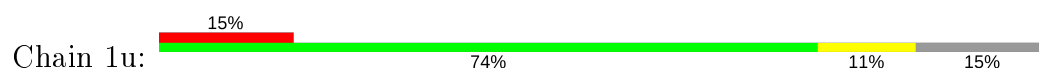
- Molecule 51: 30S ribosomal protein S20



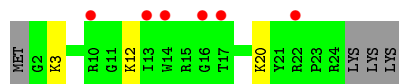
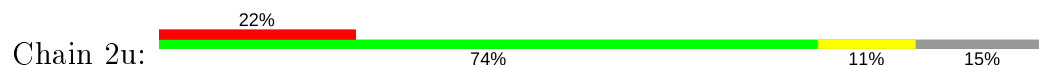
- Molecule 51: 30S ribosomal protein S20



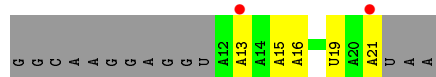
- Molecule 52: 30S ribosomal protein Thx



- Molecule 52: 30S ribosomal protein Thx



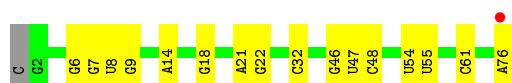
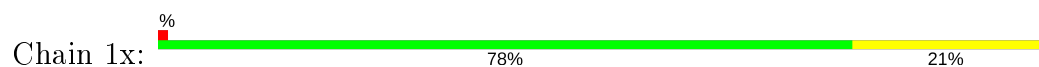
- Molecule 53: mRNA



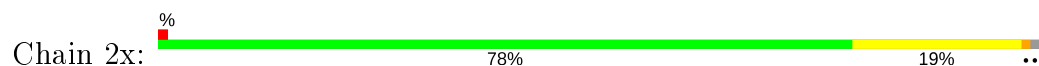
- Molecule 53: mRNA



- Molecule 54: P-site tRNA



- Molecule 54: P-site tRNA



4 Data and refinement statistics

Property	Value	Source
Space group	P 21 21 21	Depositor
Cell constants a, b, c, α , β , γ	209.13Å 448.15Å 621.94Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	86.77 – 3.00 310.97 – 3.00	Depositor EDS
% Data completeness (in resolution range)	99.0 (86.77-3.00) 99.0 (310.97-3.00)	Depositor EDS
R_{merge}	0.30	Depositor
R_{sym}	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ ¹	1.21 (at 3.01Å)	Xtriage
Refinement program	PHENIX 1.8.2	Depositor
R, R_{free}	0.246 , 0.311 0.246 , 0.310	Depositor DCC
R_{free} test set	57107 reflections (5.01%)	wwPDB-VP
Wilson B-factor (Å ²)	58.0	Xtriage
Anisotropy	0.166	Xtriage
Bulk solvent k_{sol} (e/Å ³), B_{sol} (Å ²)	0.26 , 57.7	EDS
L-test for twinning ²	$\langle L \rangle = 0.36$, $\langle L^2 \rangle = 0.19$	Xtriage
Estimated twinning fraction	No twinning to report.	Xtriage
F_o, F_c correlation	0.79	EDS
Total number of atoms	292071	wwPDB-VP
Average B, all atoms (Å ²)	54.0	wwPDB-VP

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 1.60% 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: 5MU, ZN, M2G, OMG, MA6, SF4, 0TD, MG, V7A, 2MA, 2MU, 2MG, 5MC, UR3, 4OC, 4SU, 7MG, K, 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	0.47	0/69009	0.96	54/107712 (0.1%)
1	2A	0.41	0/67293	0.92	50/105034 (0.0%)
2	1B	0.45	1/2882 (0.0%)	0.93	1/4494 (0.0%)
2	2B	0.41	1/2879 (0.0%)	0.92	1/4487 (0.0%)
3	1D	0.36	0/2186	0.54	0/2944
3	2D	0.33	0/2186	0.53	0/2944
4	1E	0.33	0/1592	0.52	0/2149
4	2E	0.31	0/1592	0.51	0/2149
5	1F	0.35	0/1619	0.52	0/2193
5	2F	0.31	0/1615	0.50	0/2188
6	1G	0.30	0/1448	0.50	0/1957
6	2G	0.28	0/1453	0.48	0/1963
7	1H	0.32	0/1356	0.50	0/1834
7	2H	0.30	0/1356	0.46	0/1834
8	1I	0.29	0/1112	0.49	0/1514
8	2I	0.29	0/1079	0.51	1/1475 (0.1%)
9	1N	0.32	0/1144	0.49	0/1543
9	2N	0.31	0/1144	0.49	0/1543
10	1O	0.34	0/943	0.51	0/1269
10	2O	0.32	0/943	0.53	0/1269
11	1P	0.32	0/1152	0.54	0/1533
11	2P	0.31	0/1152	0.54	0/1533
12	1Q	0.33	0/1143	0.53	0/1527
12	2Q	0.31	0/1143	0.51	0/1527
13	1R	0.30	0/982	0.52	0/1312
13	2R	0.29	0/982	0.48	0/1312
14	1S	0.32	0/883	0.53	0/1176
14	2S	0.31	0/880	0.48	0/1172
15	1T	0.30	0/1105	0.50	0/1477
15	2T	0.31	0/1097	0.48	0/1468
16	1U	0.34	0/977	0.45	0/1301

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
16	2U	0.28	0/977	0.42	0/1301
17	1V	0.35	0/782	0.52	0/1049
17	2V	0.28	0/782	0.51	0/1049
18	1W	0.33	0/897	0.50	0/1205
18	2W	0.31	0/897	0.48	0/1205
19	1X	0.37	0/764	0.55	0/1025
19	2X	0.31	0/764	0.50	0/1025
20	1Y	0.35	0/819	0.50	0/1095
20	2Y	0.30	0/819	0.51	0/1095
21	1Z	0.31	0/1267	0.51	0/1717
21	2Z	0.30	0/1299	0.51	0/1763
22	10	0.34	0/662	0.54	0/881
22	20	0.30	0/662	0.49	0/881
23	11	0.31	0/762	0.52	0/1014
23	21	0.31	0/762	0.52	0/1014
24	12	0.31	0/590	0.49	0/781
24	22	0.29	0/590	0.44	0/781
25	13	0.33	0/474	0.48	0/635
25	23	0.28	0/469	0.47	0/630
26	14	0.32	0/565	0.51	0/761
26	24	0.30	0/545	0.52	0/737
27	15	0.29	0/469	0.49	0/635
27	25	0.31	0/469	0.46	0/635
28	16	0.38	0/460	0.54	0/613
28	26	0.34	0/456	0.50	0/608
29	17	0.34	0/426	0.49	0/561
29	27	0.31	0/426	0.49	0/561
30	18	0.33	0/525	0.52	0/691
30	28	0.32	0/525	0.50	0/691
31	19	0.31	0/310	0.49	0/407
31	29	0.29	0/310	0.52	0/407
32	1a	0.37	0/35795	0.91	18/55864 (0.0%)
32	2a	0.37	3/35886 (0.0%)	0.93	42/56005 (0.1%)
33	1b	0.29	0/1881	0.49	0/2542
33	2b	0.30	0/1860	0.50	0/2518
34	1c	0.29	0/1572	0.48	0/2126
34	2c	0.30	0/1566	0.50	0/2119
35	1d	0.30	0/1685	0.48	0/2262
35	2d	0.28	0/1704	0.48	0/2284
36	1e	0.30	0/1145	0.51	0/1543
36	2e	0.32	0/1149	0.52	0/1548
37	1f	0.31	0/823	0.50	0/1115
37	2f	0.30	0/829	0.49	0/1123

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
38	1g	0.29	0/1250	0.46	0/1679
38	2g	0.30	0/1254	0.47	0/1683
39	1h	0.28	0/1108	0.47	0/1494
39	2h	0.27	0/1108	0.45	0/1494
40	1i	0.29	0/1002	0.51	0/1346
40	2i	0.28	0/997	0.48	0/1343
41	1j	0.29	0/722	0.48	0/982
41	2j	0.29	0/727	0.51	0/988
42	1k	0.29	0/844	0.50	0/1145
42	2k	0.29	0/848	0.46	0/1149
43	1l	0.30	0/937	0.52	0/1260
43	2l	0.30	0/937	0.53	0/1260
44	1m	0.28	0/961	0.48	0/1290
44	2m	0.30	0/953	0.48	0/1279
45	1n	0.31	0/501	0.46	0/664
45	2n	0.30	0/501	0.47	0/664
46	1o	0.27	0/739	0.46	0/985
46	2o	0.27	0/739	0.47	0/985
47	1p	0.32	0/697	0.51	0/939
47	2p	0.28	0/693	0.47	0/935
48	1q	0.29	0/836	0.48	0/1117
48	2q	0.29	0/836	0.48	0/1117
49	1r	0.27	0/560	0.45	0/746
49	2r	0.27	0/560	0.44	0/746
50	1s	0.28	0/667	0.52	1/900 (0.1%)
50	2s	0.29	0/661	0.56	0/893
51	1t	0.28	0/730	0.45	0/965
51	2t	0.27	0/729	0.45	0/965
52	1u	0.29	0/203	0.52	0/266
52	2u	0.27	0/203	0.47	0/266
53	1v	0.47	0/244	0.95	0/378
53	2v	0.57	0/126	1.13	0/195
54	1x	0.51	1/1725 (0.1%)	1.10	7/2689 (0.3%)
54	2x	0.45	0/1725	1.03	6/2689 (0.2%)
All	All	0.39	6/310069 (0.0%)	0.84	181/463831 (0.0%)

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

Mol	Chain	#Chirality outliers	#Planarity outliers
20	1Y	0	1

The worst 5 of 6 bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
2	2B	1	U	OP3-P	-10.39	1.48	1.61
2	1B	1	U	OP3-P	-10.19	1.49	1.61
32	2a	1272	G	N1-C2	-7.86	1.31	1.37
32	2a	1272	G	C6-N1	-7.51	1.34	1.39
32	2a	1263	C	N3-C4	-5.84	1.29	1.33

The worst 5 of 181 bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
32	2a	1263	C	N1-C2-O2	19.27	130.46	118.90
32	2a	1272	G	N3-C2-N2	17.88	132.41	119.90
32	2a	1272	G	C5-C6-O6	17.17	138.90	128.60
32	2a	1272	G	N1-C2-N2	-14.94	102.76	116.20
32	2a	1263	C	C2-N3-C4	11.86	125.83	119.90

There are no chirality outliers.

All (1) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
20	1Y	52	SER	Peptide

5.2 Too-close contacts [i](#)

Due to software issues we are unable to calculate clashes - this section is therefore empty.

5.3 Torsion angles [i](#)

5.3.1 Protein backbone [i](#)

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
3	1D	273/276 (99%)	249 (91%)	24 (9%)	0	100	100
3	2D	273/276 (99%)	245 (90%)	26 (10%)	2 (1%)	22	60

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
4	1E	202/206 (98%)	181 (90%)	19 (9%)	2 (1%)	15	53
4	2E	202/206 (98%)	179 (89%)	19 (9%)	4 (2%)	7	34
5	1F	201/210 (96%)	177 (88%)	21 (10%)	3 (2%)	10	42
5	2F	201/210 (96%)	175 (87%)	19 (10%)	7 (4%)	3	20
6	1G	179/182 (98%)	156 (87%)	19 (11%)	4 (2%)	6	31
6	2G	179/182 (98%)	153 (86%)	24 (13%)	2 (1%)	14	50
7	1H	172/180 (96%)	155 (90%)	12 (7%)	5 (3%)	4	24
7	2H	172/180 (96%)	150 (87%)	18 (10%)	4 (2%)	6	30
8	1I	144/148 (97%)	125 (87%)	18 (12%)	1 (1%)	22	60
8	2I	144/148 (97%)	124 (86%)	18 (12%)	2 (1%)	11	43
9	1N	138/140 (99%)	126 (91%)	11 (8%)	1 (1%)	22	60
9	2N	138/140 (99%)	126 (91%)	10 (7%)	2 (1%)	11	43
10	1O	120/122 (98%)	105 (88%)	15 (12%)	0	100	100
10	2O	120/122 (98%)	106 (88%)	13 (11%)	1 (1%)	19	57
11	1P	147/150 (98%)	122 (83%)	20 (14%)	5 (3%)	3	20
11	2P	147/150 (98%)	128 (87%)	17 (12%)	2 (1%)	11	43
12	1Q	139/141 (99%)	123 (88%)	15 (11%)	1 (1%)	22	60
12	2Q	139/141 (99%)	125 (90%)	10 (7%)	4 (3%)	4	24
13	1R	116/118 (98%)	100 (86%)	14 (12%)	2 (2%)	9	39
13	2R	116/118 (98%)	105 (90%)	11 (10%)	0	100	100
14	1S	108/112 (96%)	94 (87%)	14 (13%)	0	100	100
14	2S	108/112 (96%)	100 (93%)	7 (6%)	1 (1%)	17	55
15	1T	129/146 (88%)	109 (84%)	18 (14%)	2 (2%)	9	40
15	2T	129/146 (88%)	114 (88%)	14 (11%)	1 (1%)	19	57
16	1U	114/118 (97%)	111 (97%)	3 (3%)	0	100	100
16	2U	114/118 (97%)	107 (94%)	7 (6%)	0	100	100
17	1V	99/101 (98%)	84 (85%)	14 (14%)	1 (1%)	15	53
17	2V	99/101 (98%)	95 (96%)	4 (4%)	0	100	100
18	1W	110/113 (97%)	107 (97%)	2 (2%)	1 (1%)	17	55
18	2W	110/113 (97%)	101 (92%)	8 (7%)	1 (1%)	17	55
19	1X	93/96 (97%)	85 (91%)	8 (9%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
19	2X	93/96 (97%)	80 (86%)	12 (13%)	1 (1%)	14	50
20	1Y	105/110 (96%)	92 (88%)	12 (11%)	1 (1%)	15	53
20	2Y	105/110 (96%)	93 (89%)	8 (8%)	4 (4%)	3	18
21	1Z	148/206 (72%)	120 (81%)	23 (16%)	5 (3%)	3	20
21	2Z	156/206 (76%)	118 (76%)	35 (22%)	3 (2%)	8	36
22	10	81/85 (95%)	74 (91%)	6 (7%)	1 (1%)	13	48
22	20	81/85 (95%)	74 (91%)	7 (9%)	0	100	100
23	11	95/98 (97%)	88 (93%)	7 (7%)	0	100	100
23	21	95/98 (97%)	85 (90%)	7 (7%)	3 (3%)	4	22
24	12	68/72 (94%)	61 (90%)	7 (10%)	0	100	100
24	22	68/72 (94%)	66 (97%)	2 (3%)	0	100	100
25	13	57/60 (95%)	53 (93%)	4 (7%)	0	100	100
25	23	57/60 (95%)	52 (91%)	5 (9%)	0	100	100
26	14	67/71 (94%)	50 (75%)	12 (18%)	5 (8%)	1	5
26	24	67/71 (94%)	50 (75%)	14 (21%)	3 (4%)	2	14
27	15	57/60 (95%)	51 (90%)	5 (9%)	1 (2%)	8	37
27	25	57/60 (95%)	52 (91%)	5 (9%)	0	100	100
28	16	51/54 (94%)	46 (90%)	5 (10%)	0	100	100
28	26	51/54 (94%)	43 (84%)	8 (16%)	0	100	100
29	17	46/49 (94%)	45 (98%)	1 (2%)	0	100	100
29	27	46/49 (94%)	43 (94%)	3 (6%)	0	100	100
30	18	62/65 (95%)	59 (95%)	3 (5%)	0	100	100
30	28	62/65 (95%)	57 (92%)	5 (8%)	0	100	100
31	19	35/37 (95%)	32 (91%)	3 (9%)	0	100	100
31	29	35/37 (95%)	30 (86%)	5 (14%)	0	100	100
33	1b	229/256 (90%)	186 (81%)	36 (16%)	7 (3%)	4	23
33	2b	229/256 (90%)	193 (84%)	28 (12%)	8 (4%)	3	20
34	1c	204/239 (85%)	179 (88%)	22 (11%)	3 (2%)	10	42
34	2c	204/239 (85%)	171 (84%)	29 (14%)	4 (2%)	7	34
35	1d	206/209 (99%)	173 (84%)	31 (15%)	2 (1%)	15	53
35	2d	206/209 (99%)	185 (90%)	17 (8%)	4 (2%)	8	36

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
36	1e	146/162 (90%)	127 (87%)	15 (10%)	4 (3%)	5	26
36	2e	146/162 (90%)	129 (88%)	14 (10%)	3 (2%)	7	33
37	1f	98/101 (97%)	91 (93%)	7 (7%)	0	100	100
37	2f	98/101 (97%)	89 (91%)	9 (9%)	0	100	100
38	1g	153/156 (98%)	133 (87%)	17 (11%)	3 (2%)	7	34
38	2g	153/156 (98%)	132 (86%)	18 (12%)	3 (2%)	7	34
39	1h	135/138 (98%)	125 (93%)	10 (7%)	0	100	100
39	2h	135/138 (98%)	124 (92%)	10 (7%)	1 (1%)	22	60
40	1i	125/128 (98%)	102 (82%)	20 (16%)	3 (2%)	6	29
40	2i	125/128 (98%)	106 (85%)	15 (12%)	4 (3%)	4	22
41	1j	95/105 (90%)	78 (82%)	15 (16%)	2 (2%)	7	33
41	2j	94/105 (90%)	75 (80%)	16 (17%)	3 (3%)	4	22
42	1k	112/129 (87%)	90 (80%)	19 (17%)	3 (3%)	5	26
42	2k	112/129 (87%)	95 (85%)	13 (12%)	4 (4%)	3	19
43	1l	119/132 (90%)	103 (87%)	14 (12%)	2 (2%)	9	39
43	2l	119/132 (90%)	99 (83%)	19 (16%)	1 (1%)	19	57
44	1m	120/126 (95%)	96 (80%)	19 (16%)	5 (4%)	3	16
44	2m	119/126 (94%)	99 (83%)	17 (14%)	3 (2%)	5	28
45	1n	58/61 (95%)	52 (90%)	6 (10%)	0	100	100
45	2n	58/61 (95%)	55 (95%)	2 (3%)	1 (2%)	9	39
46	1o	86/89 (97%)	77 (90%)	6 (7%)	3 (4%)	3	20
46	2o	86/89 (97%)	77 (90%)	8 (9%)	1 (1%)	13	48
47	1p	80/88 (91%)	71 (89%)	7 (9%)	2 (2%)	5	28
47	2p	80/88 (91%)	64 (80%)	15 (19%)	1 (1%)	12	45
48	1q	97/105 (92%)	84 (87%)	12 (12%)	1 (1%)	15	53
48	2q	97/105 (92%)	88 (91%)	8 (8%)	1 (1%)	15	53
49	1r	66/88 (75%)	60 (91%)	5 (8%)	1 (2%)	10	42
49	2r	66/88 (75%)	64 (97%)	2 (3%)	0	100	100
50	1s	81/93 (87%)	65 (80%)	15 (18%)	1 (1%)	13	48
50	2s	81/93 (87%)	60 (74%)	21 (26%)	0	100	100
51	1t	94/106 (89%)	78 (83%)	13 (14%)	3 (3%)	4	22

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
51	2t	94/106 (89%)	84 (89%)	8 (8%)	2 (2%)	7	33
52	1u	21/27 (78%)	13 (62%)	6 (29%)	2 (10%)	0	3
52	2u	21/27 (78%)	16 (76%)	4 (19%)	1 (5%)	2	13
All	All	11368/12128 (94%)	9944 (88%)	1244 (11%)	180 (2%)	9	40

5 of 180 Ramachandran outliers are listed below:

Mol	Chain	Res	Type
5	1F	130	ALA
7	1H	92	ILE
11	1P	45	LEU
33	1b	17	PHE
33	1b	33	TYR

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%)	191 (89%)	24 (11%)	6	24
3	2D	215/218 (99%)	198 (92%)	17 (8%)	12	41
4	1E	164/166 (99%)	148 (90%)	16 (10%)	8	30
4	2E	164/166 (99%)	146 (89%)	18 (11%)	6	25
5	1F	160/166 (96%)	144 (90%)	16 (10%)	7	29
5	2F	159/166 (96%)	144 (91%)	15 (9%)	8	32
6	1G	143/156 (92%)	124 (87%)	19 (13%)	4	17
6	2G	143/156 (92%)	128 (90%)	15 (10%)	7	27
7	1H	144/148 (97%)	127 (88%)	17 (12%)	5	22
7	2H	144/148 (97%)	118 (82%)	26 (18%)	1	9
8	1I	113/124 (91%)	94 (83%)	19 (17%)	2	11
8	2I	105/124 (85%)	87 (83%)	18 (17%)	2	10
9	1N	118/119 (99%)	106 (90%)	12 (10%)	7	28

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
9	2N	118/119 (99%)	107 (91%)	11 (9%)	9	33
10	1O	100/100 (100%)	94 (94%)	6 (6%)	19	53
10	2O	100/100 (100%)	91 (91%)	9 (9%)	9	35
11	1P	115/116 (99%)	103 (90%)	12 (10%)	7	27
11	2P	115/116 (99%)	103 (90%)	12 (10%)	7	27
12	1Q	111/111 (100%)	100 (90%)	11 (10%)	8	30
12	2Q	111/111 (100%)	102 (92%)	9 (8%)	11	40
13	1R	101/101 (100%)	92 (91%)	9 (9%)	9	35
13	2R	101/101 (100%)	91 (90%)	10 (10%)	8	30
14	1S	86/88 (98%)	73 (85%)	13 (15%)	3	14
14	2S	85/88 (97%)	76 (89%)	9 (11%)	6	26
15	1T	115/127 (91%)	98 (85%)	17 (15%)	3	14
15	2T	113/127 (89%)	106 (94%)	7 (6%)	18	52
16	1U	93/94 (99%)	90 (97%)	3 (3%)	39	74
16	2U	93/94 (99%)	86 (92%)	7 (8%)	13	43
17	1V	80/82 (98%)	72 (90%)	8 (10%)	7	29
17	2V	80/82 (98%)	76 (95%)	4 (5%)	24	60
18	1W	90/92 (98%)	84 (93%)	6 (7%)	16	49
18	2W	90/92 (98%)	86 (96%)	4 (4%)	28	65
19	1X	77/78 (99%)	71 (92%)	6 (8%)	12	42
19	2X	77/78 (99%)	74 (96%)	3 (4%)	32	69
20	1Y	85/91 (93%)	74 (87%)	11 (13%)	4	19
20	2Y	85/91 (93%)	78 (92%)	7 (8%)	11	39
21	1Z	135/179 (75%)	113 (84%)	22 (16%)	2	11
21	2Z	137/179 (76%)	114 (83%)	23 (17%)	2	11
22	10	65/67 (97%)	59 (91%)	6 (9%)	9	34
22	20	65/67 (97%)	58 (89%)	7 (11%)	6	26
23	11	80/83 (96%)	76 (95%)	4 (5%)	24	60
23	21	80/83 (96%)	72 (90%)	8 (10%)	7	29
24	12	65/67 (97%)	56 (86%)	9 (14%)	3	17
24	22	65/67 (97%)	57 (88%)	8 (12%)	4	21

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
25	13	51/52 (98%)	42 (82%)	9 (18%)	2	10
25	23	50/52 (96%)	39 (78%)	11 (22%)	1	4
26	14	59/63 (94%)	44 (75%)	15 (25%)	0	3
26	24	53/63 (84%)	42 (79%)	11 (21%)	1	5
27	15	50/52 (96%)	46 (92%)	4 (8%)	12	40
27	25	50/52 (96%)	48 (96%)	2 (4%)	31	68
28	16	51/52 (98%)	46 (90%)	5 (10%)	8	30
28	26	50/52 (96%)	35 (70%)	15 (30%)	0	1
29	17	41/42 (98%)	38 (93%)	3 (7%)	14	44
29	27	41/42 (98%)	37 (90%)	4 (10%)	8	30
30	18	54/55 (98%)	51 (94%)	3 (6%)	21	56
30	28	54/55 (98%)	49 (91%)	5 (9%)	9	33
31	19	34/34 (100%)	32 (94%)	2 (6%)	19	54
31	29	34/34 (100%)	31 (91%)	3 (9%)	10	36
33	1b	192/220 (87%)	162 (84%)	30 (16%)	2	13
33	2b	187/220 (85%)	158 (84%)	29 (16%)	2	13
34	1c	142/188 (76%)	126 (89%)	16 (11%)	6	24
34	2c	140/188 (74%)	123 (88%)	17 (12%)	5	21
35	1d	169/181 (93%)	145 (86%)	24 (14%)	3	16
35	2d	173/181 (96%)	153 (88%)	20 (12%)	5	23
36	1e	113/123 (92%)	101 (89%)	12 (11%)	6	26
36	2e	114/123 (93%)	103 (90%)	11 (10%)	8	32
37	1f	84/90 (93%)	78 (93%)	6 (7%)	14	46
37	2f	85/90 (94%)	74 (87%)	11 (13%)	4	19
38	1g	119/127 (94%)	101 (85%)	18 (15%)	3	14
38	2g	120/127 (94%)	104 (87%)	16 (13%)	4	17
39	1h	114/119 (96%)	105 (92%)	9 (8%)	12	41
39	2h	114/119 (96%)	105 (92%)	9 (8%)	12	41
40	1i	90/99 (91%)	78 (87%)	12 (13%)	4	17
40	2i	89/99 (90%)	74 (83%)	15 (17%)	2	11
41	1j	66/92 (72%)	54 (82%)	12 (18%)	1	9

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
41	2j	69/92 (75%)	63 (91%)	6 (9%)	10	37
42	1k	82/99 (83%)	73 (89%)	9 (11%)	6	25
42	2k	83/99 (84%)	75 (90%)	8 (10%)	8	32
43	1l	96/108 (89%)	87 (91%)	9 (9%)	8	32
43	2l	96/108 (89%)	85 (88%)	11 (12%)	5	24
44	1m	92/101 (91%)	78 (85%)	14 (15%)	3	14
44	2m	91/101 (90%)	77 (85%)	14 (15%)	2	13
45	1n	49/50 (98%)	43 (88%)	6 (12%)	5	21
45	2n	49/50 (98%)	44 (90%)	5 (10%)	7	28
46	1o	78/80 (98%)	72 (92%)	6 (8%)	13	42
46	2o	78/80 (98%)	72 (92%)	6 (8%)	13	42
47	1p	69/74 (93%)	55 (80%)	14 (20%)	1	6
47	2p	68/74 (92%)	55 (81%)	13 (19%)	1	8
48	1q	94/97 (97%)	88 (94%)	6 (6%)	17	51
48	2q	94/97 (97%)	87 (93%)	7 (7%)	13	44
49	1r	59/77 (77%)	53 (90%)	6 (10%)	7	28
49	2r	59/77 (77%)	56 (95%)	3 (5%)	24	60
50	1s	69/80 (86%)	65 (94%)	4 (6%)	20	55
50	2s	67/80 (84%)	58 (87%)	9 (13%)	4	17
51	1t	70/82 (85%)	64 (91%)	6 (9%)	10	37
51	2t	70/82 (85%)	58 (83%)	12 (17%)	2	10
52	1u	18/22 (82%)	17 (94%)	1 (6%)	21	56
52	2u	18/22 (82%)	16 (89%)	2 (11%)	6	25
All	All	9301/10064 (92%)	8252 (89%)	1049 (11%)	6	24

5 of 1049 residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
44	1m	64	TRP
6	2G	99	MET
43	2l	33	ARG
46	1o	66	LEU
3	2D	88	ARG

Some sidechains can be flipped to improve hydrogen bonding and reduce clashes. 5 of 115 such sidechains are listed below:

Mol	Chain	Res	Type
41	1j	69	ASN
5	2F	69	HIS
42	2k	93	GLN
42	1k	117	ASN
50	1s	69	HIS

5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	1A	2861/2915 (98%)	554 (19%)	21 (0%)
1	2A	2788/2915 (95%)	579 (20%)	21 (0%)
2	1B	120/121 (99%)	9 (7%)	1 (0%)
2	2B	118/121 (97%)	32 (27%)	0
32	1a	1494/1521 (98%)	303 (20%)	0
32	2a	1498/1521 (98%)	336 (22%)	0
53	1v	9/24 (37%)	5 (55%)	0
53	2v	4/24 (16%)	1 (25%)	0
54	1x	75/77 (97%)	9 (12%)	0
54	2x	75/77 (97%)	9 (12%)	0
All	All	9042/9316 (97%)	1837 (20%)	43 (0%)

5 of 1837 RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	1A	10	G
1	1A	11	G
1	1A	15	G
1	1A	34	C
1	1A	36	G

5 of 43 RNA pucker outliers are listed below:

Mol	Chain	Res	Type
1	1A	2473	U
1	2A	266	G
1	2A	1913	A
1	1A	2689	U
2	1B	1	U

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

56 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
54	4SU	2x	8	54	14,21,22	1.38	2 (14%)	15,30,33	2.32	2 (13%)
32	5MC	1a	967	32	15,22,23	1.21	1 (6%)	19,32,35	1.28	2 (10%)
54	5MU	1x	54	54	15,22,23	1.04	2 (13%)	16,32,35	2.32	1 (6%)
32	PSU	1a	516	55,32	17,21,22	1.47	3 (17%)	20,30,33	3.19	6 (30%)
1	5MU	1A	1939	1	15,22,23	1.11	2 (13%)	16,32,35	1.82	2 (12%)
32	5MC	2a	967	32	15,22,23	1.33	1 (6%)	19,32,35	1.31	2 (10%)
1	2MA	2A	2503	1,55	17,25,26	1.35	2 (11%)	19,37,40	2.03	3 (15%)
1	5MU	1A	1915	1	15,22,23	1.04	1 (6%)	16,32,35	1.73	2 (12%)
32	5MC	1a	1407	32	15,22,23	1.25	1 (6%)	19,32,35	1.38	2 (10%)
32	UR3	1a	1498	32	14,22,23	0.74	1 (7%)	15,32,35	0.77	1 (6%)
32	5MC	2a	1400	32,54	15,22,23	1.46	1 (6%)	19,32,35	1.40	2 (10%)
32	7MG	1a	527	55,32	22,26,27	1.76	4 (18%)	28,39,42	2.74	8 (28%)
32	2MG	1a	1207	32	19,26,27	1.22	2 (10%)	21,38,41	2.19	8 (38%)
32	MA6	1a	1518	32	19,26,27	0.89	1 (5%)	18,38,41	1.71	4 (22%)
54	5MC	1x	32	54	15,22,23	1.28	1 (6%)	19,32,35	1.48	4 (21%)
1	PSU	1A	2605	1	17,21,22	1.57	3 (17%)	20,30,33	3.06	6 (30%)
1	2MU	2A	2552	1,55	14,22,24	0.90	0	14,31,36	0.92	0
1	5MC	2A	1942	1,55	15,22,23	1.38	1 (6%)	19,32,35	1.46	3 (15%)
32	MA6	2a	1518	32	19,26,27	0.94	1 (5%)	18,38,41	1.72	4 (22%)
32	5MC	1a	1404	32	15,22,23	1.38	1 (6%)	19,32,35	1.25	3 (15%)
32	UR3	2a	1498	32	14,22,23	0.76	1 (7%)	15,32,35	0.70	0
1	5MU	2A	1939	1	15,22,23	1.18	2 (13%)	16,32,35	1.82	2 (12%)
54	4SU	1x	8	54	14,21,22	1.47	2 (14%)	15,30,33	2.63	2 (13%)
1	PSU	2A	1911	1	17,21,22	1.56	2 (11%)	20,30,33	3.16	6 (30%)
1	4OC	2A	1920	1	15,22,24	0.67	0	17,31,35	1.41	2 (11%)
54	PSU	1x	55	54	17,21,22	1.59	2 (11%)	20,30,33	3.32	6 (30%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
32	4OC	1a	1402	32	16,23,24	0.59	0	17,32,35	1.31	1 (5%)
1	5MC	2A	1962	1,55	15,22,23	1.32	1 (6%)	19,32,35	1.37	2 (10%)
1	5MC	1A	1942	1	15,22,23	1.19	1 (6%)	19,32,35	1.39	3 (15%)
32	5MC	2a	1404	32	15,22,23	1.29	1 (6%)	19,32,35	1.40	2 (10%)
32	MA6	1a	1519	32	19,26,27	1.02	1 (5%)	18,38,41	1.60	3 (16%)
43	0TD	1l	92	43	4,9,10	3.03	1 (25%)	3,11,13	2.20	1 (33%)
32	M2G	2a	966	55,32	20,27,28	1.46	3 (15%)	22,40,43	2.19	6 (27%)
32	PSU	2a	516	32	17,21,22	1.45	3 (17%)	20,30,33	3.18	5 (25%)
1	PSU	1A	1917	1	17,21,22	1.85	4 (23%)	20,30,33	3.28	6 (30%)
32	7MG	2a	527	55,32	22,26,27	1.72	4 (18%)	28,39,42	2.67	9 (32%)
1	PSU	1A	1911	1	17,21,22	1.53	2 (11%)	20,30,33	3.22	6 (30%)
1	5MC	1A	1962	1	15,22,23	1.32	1 (6%)	19,32,35	1.30	2 (10%)
43	0TD	2l	92	43	4,9,10	3.20	1 (25%)	3,11,13	7.91	1 (33%)
1	5MU	2A	1915	1,55	15,22,23	1.16	1 (6%)	16,32,35	1.84	2 (12%)
32	M2G	1a	966	55,32	20,27,28	1.43	3 (15%)	22,40,43	2.17	7 (31%)
54	PSU	2x	55	54	17,21,22	1.66	2 (11%)	20,30,33	3.21	6 (30%)
32	2MG	2a	1207	55,32	19,26,27	1.25	2 (10%)	21,38,41	2.26	7 (33%)
1	OMG	1A	2251	1,55,54	18,26,27	1.28	2 (11%)	20,38,41	2.00	7 (35%)
1	PSU	2A	1917	1	17,21,22	1.57	3 (17%)	20,30,33	3.09	6 (30%)
32	4OC	2a	1402	32	16,23,24	0.66	0	17,32,35	1.36	3 (17%)
1	OMG	2A	2251	1,55,54	18,26,27	1.06	2 (11%)	20,38,41	2.12	5 (25%)
1	PSU	2A	2605	1	17,21,22	1.49	2 (11%)	20,30,33	3.15	6 (30%)
32	5MC	1a	1400	32	15,22,23	1.32	1 (6%)	19,32,35	1.43	2 (10%)
32	MA6	2a	1519	32	19,26,27	1.03	1 (5%)	18,38,41	1.69	5 (27%)
1	2MU	1A	2552	1,55	14,22,24	0.88	0	14,31,36	0.86	1 (7%)
54	5MU	2x	54	54	15,22,23	1.08	1 (6%)	16,32,35	1.74	2 (12%)
32	5MC	2a	1407	32	15,22,23	1.37	1 (6%)	19,32,35	1.36	2 (10%)
1	2MA	1A	2503	1,55	17,25,26	1.30	2 (11%)	19,37,40	2.30	5 (26%)
54	5MC	2x	32	54	15,22,23	1.36	1 (6%)	19,32,35	1.37	3 (15%)
1	4OC	1A	1920	1	15,22,24	0.67	0	17,31,35	1.38	1 (5%)

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
54	4SU	2x	8	54	-	0/5/25/26	0/2/2/2
32	5MC	1a	967	32	-	2/5/25/26	0/2/2/2
54	5MU	1x	54	54	-	0/5/25/26	0/2/2/2
32	PSU	1a	516	55,32	-	0/7/25/26	0/2/2/2
1	5MU	1A	1939	1	-	0/5/25/26	0/2/2/2
32	5MC	2a	967	32	-	0/5/25/26	0/2/2/2
1	2MA	2A	2503	1,55	-	1/3/25/26	0/3/3/3
1	5MU	1A	1915	1	-	0/5/25/26	0/2/2/2
32	5MC	1a	1407	32	-	0/5/25/26	0/2/2/2
32	UR3	1a	1498	32	-	0/5/25/26	0/2/2/2
32	5MC	2a	1400	32,54	-	0/5/25/26	0/2/2/2
32	7MG	1a	527	55,32	-	2/7/37/38	0/3/3/3
32	2MG	1a	1207	32	-	2/5/27/28	0/3/3/3
32	MA6	1a	1518	32	-	3/7/29/30	0/3/3/3
54	5MC	1x	32	54	-	0/5/25/26	0/2/2/2
1	PSU	1A	2605	1	-	0/7/25/26	0/2/2/2
1	2MU	2A	2552	1,55	-	2/7/27/28	0/2/2/2
1	5MC	2A	1942	1,55	-	0/5/25/26	0/2/2/2
32	MA6	2a	1518	32	-	3/7/29/30	0/3/3/3
32	5MC	1a	1404	32	-	0/5/25/26	0/2/2/2
32	UR3	2a	1498	32	-	0/5/25/26	0/2/2/2
1	5MU	2A	1939	1	-	2/5/25/26	0/2/2/2
54	4SU	1x	8	54	-	0/5/25/26	0/2/2/2
1	PSU	2A	1911	1	-	1/7/25/26	0/2/2/2
1	4OC	2A	1920	1	-	1/7/27/30	0/2/2/2
54	PSU	1x	55	54	-	0/7/25/26	0/2/2/2
32	4OC	1a	1402	32	-	4/9/29/30	0/2/2/2
1	5MC	2A	1962	1,55	-	2/5/25/26	0/2/2/2
1	5MC	1A	1942	1	-	0/5/25/26	0/2/2/2
32	5MC	2a	1404	32	-	0/5/25/26	0/2/2/2
32	MA6	1a	1519	32	-	3/7/29/30	0/3/3/3
43	0TD	1l	92	43	-	2/3/12/14	-
32	M2G	2a	966	55,32	-	0/7/29/30	0/3/3/3
32	PSU	2a	516	32	-	0/7/25/26	0/2/2/2
1	PSU	1A	1917	1	-	0/7/25/26	0/2/2/2
32	7MG	2a	527	55,32	-	2/7/37/38	0/3/3/3
1	PSU	1A	1911	1	-	0/7/25/26	0/2/2/2
1	5MC	1A	1962	1	-	2/5/25/26	0/2/2/2
43	0TD	2l	92	43	-	1/3/12/14	-
1	5MU	2A	1915	1,55	-	1/5/25/26	0/2/2/2

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
32	M2G	1a	966	55,32	-	1/7/29/30	0/3/3/3
54	PSU	2x	55	54	-	0/7/25/26	0/2/2/2
32	2MG	2a	1207	55,32	-	1/5/27/28	0/3/3/3
1	OMG	1A	2251	1,55,54	-	0/5/27/28	0/3/3/3
1	PSU	2A	1917	1	-	1/7/25/26	0/2/2/2
32	4OC	2a	1402	32	-	4/9/29/30	0/2/2/2
1	OMG	2A	2251	1,55,54	-	0/5/27/28	0/3/3/3
1	PSU	2A	2605	1	-	0/7/25/26	0/2/2/2
32	5MC	1a	1400	32	-	2/5/25/26	0/2/2/2
32	MA6	2a	1519	32	-	4/7/29/30	0/3/3/3
1	2MU	1A	2552	1,55	-	0/7/27/28	0/2/2/2
54	5MU	2x	54	54	-	0/5/25/26	0/2/2/2
32	5MC	2a	1407	32	-	0/5/25/26	0/2/2/2
1	2MA	1A	2503	1,55	-	3/3/25/26	0/3/3/3
54	5MC	2x	32	54	-	0/5/25/26	0/2/2/2
1	4OC	1A	1920	1	-	1/7/27/30	0/2/2/2

The worst 5 of 87 bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
43	2l	92	0TD	CB-SB	-6.10	1.69	1.84
1	1A	1917	PSU	C5-C1'	-5.88	1.47	1.52
43	1l	92	0TD	CB-SB	-5.72	1.70	1.84
32	2a	1400	5MC	C5-C4	5.21	1.49	1.41
1	2A	1942	5MC	C5-C4	4.95	1.49	1.41

The worst 5 of 200 bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
43	2l	92	0TD	CSB-SB-CB	-13.65	75.00	101.85
54	1x	8	4SU	C2-N3-C4	8.99	128.18	115.15
32	2a	516	PSU	N1-C2-N3	-8.95	121.32	128.43
54	1x	54	5MU	C4-N3-C2	8.93	122.68	115.14
32	1a	516	PSU	N1-C2-N3	-8.88	121.37	128.43

There are no chirality outliers.

5 of 53 torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
32	1a	1207	2MG	N1-C2-N2-CM2
32	1a	1207	2MG	N3-C2-N2-CM2

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Mol	Chain	Res	Type	Atoms
32	1a	1518	MA6	C5-C6-N6-C9
32	1a	1518	MA6	C5-C6-N6-C10
32	2a	1518	MA6	C5-C6-N6-C9

There are no ring outliers.

No monomer is involved in short contacts.

5.5 Carbohydrates [i](#)

There are no monosaccharides in this entry.

5.6 Ligand geometry [i](#)

Of 2656 ligands modelled in this entry, 2652 are monoatomic - leaving 4 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
59	SF4	1d	302	35	0,12,12	0.00	-	-		
59	SF4	2d	303	35	0,12,12	0.00	-	-		
58	V7A	1a	3213	55	36,38,38	4.06	12 (33%)	40,60,60	2.03	10 (25%)
58	V7A	2a	1817	55	36,38,38	4.08	12 (33%)	40,60,60	2.01	10 (25%)

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
59	SF4	1d	302	35	-	-	0/6/5/5
59	SF4	2d	303	35	-	-	0/6/5/5
58	V7A	1a	3213	55	-	4/13/72/72	0/4/4/4
58	V7A	2a	1817	55	-	3/13/72/72	0/4/4/4

The worst 5 of 24 bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
58	2a	1817	V7A	CAM-CAL	-13.28	1.40	1.52
58	1a	3213	V7A	CAM-CAL	-12.63	1.41	1.52
58	1a	3213	V7A	CAM-CAP	-11.44	1.39	1.55
58	2a	1817	V7A	CAM-CAP	-10.90	1.40	1.55
58	1a	3213	V7A	CAJ-CAI	-10.84	1.40	1.51

The worst 5 of 20 bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
58	1a	3213	V7A	CAV-OAU-NAT	5.69	120.33	108.11
58	1a	3213	V7A	OAZ-CAL-CAI	-5.30	116.65	123.90
58	2a	1817	V7A	CAV-OAU-NAT	5.19	119.24	108.11
58	2a	1817	V7A	CAJ-CAK-CAN	-4.67	102.28	110.49
58	2a	1817	V7A	OAZ-CAL-CAI	-4.50	117.74	123.90

There are no chirality outliers.

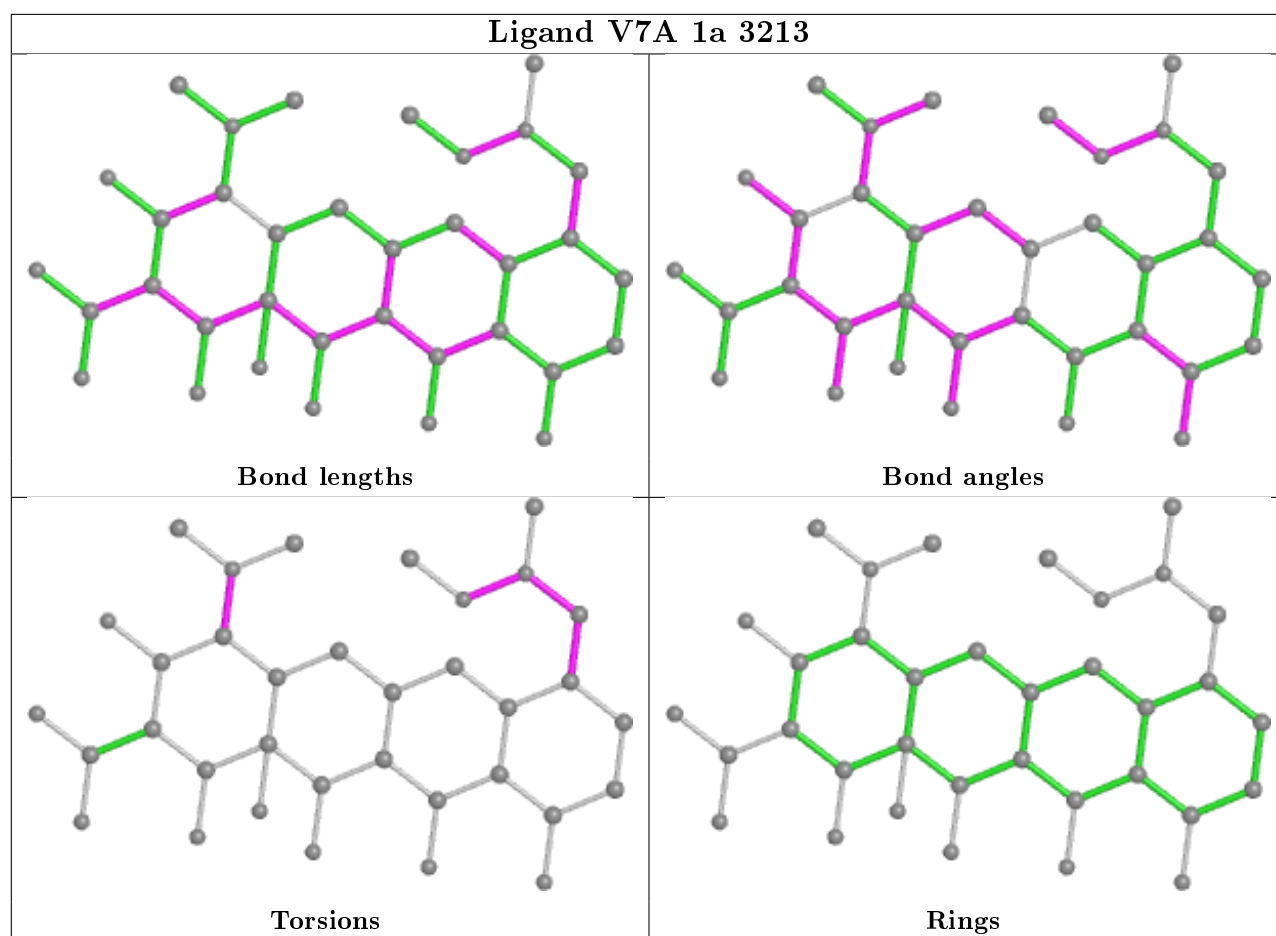
5 of 7 torsion outliers are listed below:

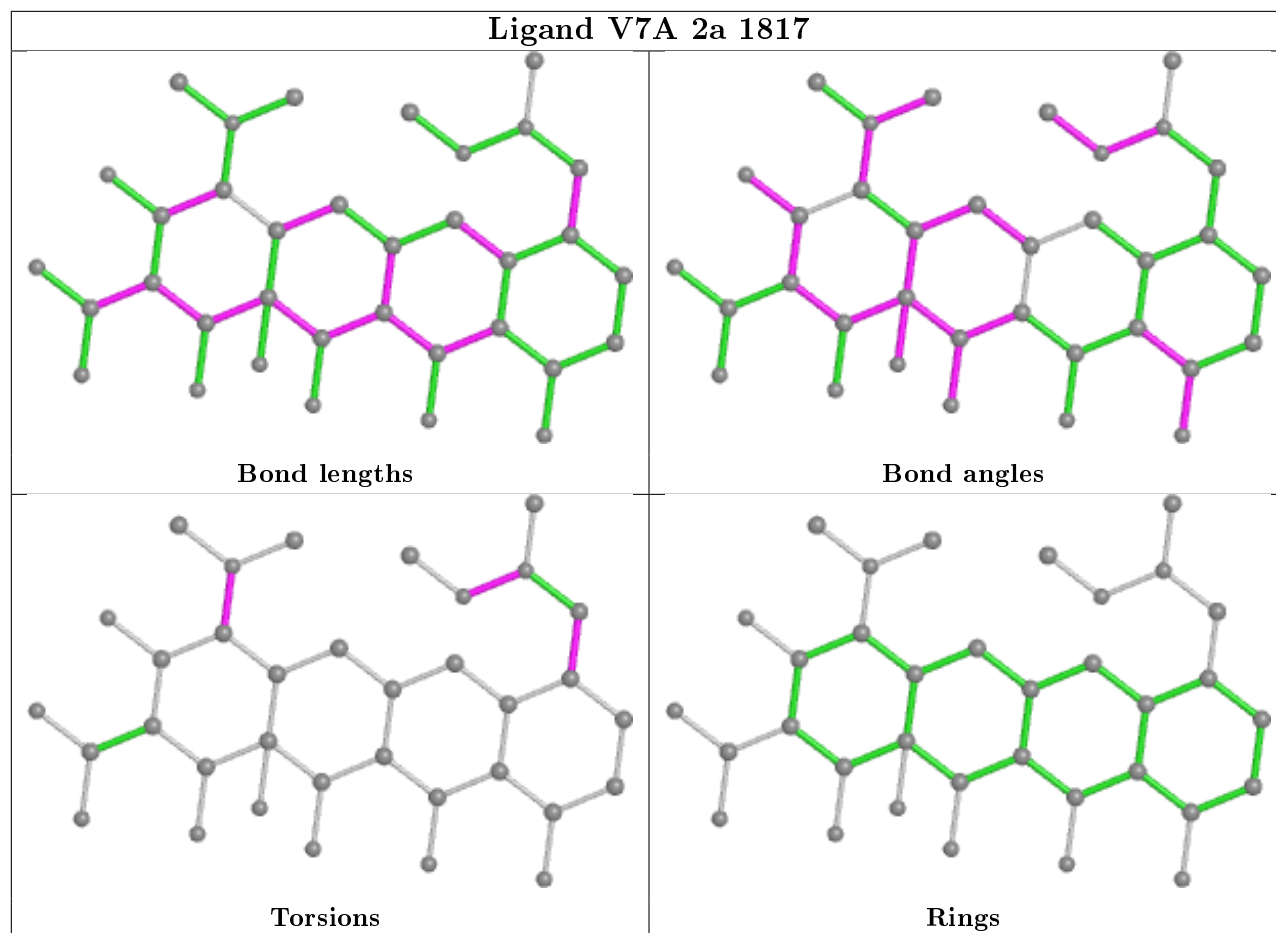
Mol	Chain	Res	Type	Atoms
58	1a	3213	V7A	CAN-CAO-NBF-CBH
58	1a	3213	V7A	CAW-NAT-OAU-CAV
58	2a	1817	V7A	CAN-CAO-NBF-CBH
58	2a	1817	V7A	CAW-NAT-OAU-CAV
58	1a	3213	V7A	CAA-CAS-NAT-CAW

There are no ring outliers.

No monomer is involved in short contacts.

The following is a two-dimensional graphical depiction of Mogul quality analysis of bond lengths, bond angles, torsion angles, and ring geometry for all instances of the Ligand of Interest. In addition, ligands with molecular weight > 250 and outliers as shown on the validation Tables will also be included. For torsion angles, if less than 5% of the Mogul distribution of torsion angles is within 10 degrees of the torsion angle in question, then that torsion angle is considered an outlier. Any bond that is central to one or more torsion angles identified as an outlier by Mogul will be highlighted in the graph. For rings, the root-mean-square deviation (RMSD) between the ring in question and similar rings identified by Mogul is calculated over all ring torsion angles. If the average RMSD is greater than 60 degrees and the minimal RMSD between the ring in question and any Mogul-identified rings is also greater than 60 degrees, then that ring is considered an outlier. The outliers are highlighted in purple. The color gray indicates Mogul did not find sufficient equivalents in the CSD to analyse the geometry.





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 [i](#)

6.1 Protein, DNA and RNA chains [i](#)

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%)	0.06	13 (0%) 91 75	11, 36, 96, 112	0
1	2A	2789/2915 (95%)	0.08	12 (0%) 92 79	21, 49, 90, 116	0
2	1B	120/121 (99%)	-0.24	0 100 100	27, 51, 64, 85	0
2	2B	120/121 (99%)	-0.13	0 100 100	49, 68, 83, 95	0
3	1D	275/276 (99%)	0.14	0 100 100	14, 31, 47, 61	0
3	2D	275/276 (99%)	0.30	4 (1%) 73 46	22, 42, 58, 84	0
4	1E	204/206 (99%)	0.03	2 (0%) 82 59	18, 43, 63, 78	0
4	2E	204/206 (99%)	0.23	8 (3%) 39 15	24, 50, 66, 77	0
5	1F	203/210 (96%)	-0.08	1 (0%) 91 75	16, 40, 64, 82	0
5	2F	203/210 (96%)	0.05	3 (1%) 73 46	27, 57, 73, 79	0
6	1G	181/182 (99%)	-0.12	1 (0%) 89 72	37, 53, 70, 83	0
6	2G	181/182 (99%)	0.64	23 (12%) 3 1	53, 68, 82, 88	0
7	1H	174/180 (96%)	-0.15	0 100 100	33, 51, 64, 71	0
7	2H	174/180 (96%)	0.78	31 (17%) 1 0	50, 75, 84, 88	0
8	1I	146/148 (98%)	-0.34	0 100 100	40, 64, 75, 81	0
8	2I	146/148 (98%)	-0.14	2 (1%) 75 49	42, 68, 80, 87	0
9	1N	140/140 (100%)	0.01	0 100 100	22, 42, 60, 68	0
9	2N	140/140 (100%)	0.51	12 (8%) 10 3	35, 58, 75, 82	0
10	1O	122/122 (100%)	0.22	1 (0%) 86 65	23, 37, 55, 65	0
10	2O	122/122 (100%)	-0.00	2 (1%) 72 44	30, 44, 60, 71	0
11	1P	149/150 (99%)	0.24	8 (5%) 25 9	17, 43, 65, 72	0
11	2P	149/150 (99%)	0.56	18 (12%) 4 1	30, 59, 74, 85	0
12	1Q	141/141 (100%)	0.17	1 (0%) 87 69	25, 41, 53, 61	0
12	2Q	141/141 (100%)	0.46	3 (2%) 63 34	38, 52, 65, 70	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
13	1R	118/118 (100%)	0.15	1 (0%) 86 65	22, 39, 53, 62	0
13	2R	118/118 (100%)	0.17	2 (1%) 70 41	31, 44, 57, 67	0
14	1S	110/112 (98%)	-0.17	1 (0%) 84 63	33, 45, 57, 65	0
14	2S	110/112 (98%)	-0.07	0 100 100	48, 60, 71, 79	0
15	1T	131/146 (89%)	-0.01	0 100 100	27, 48, 67, 79	0
15	2T	131/146 (89%)	-0.01	1 (0%) 86 65	36, 49, 67, 79	0
16	1U	116/118 (98%)	0.19	0 100 100	22, 37, 56, 63	0
16	2U	116/118 (98%)	0.44	7 (6%) 21 7	37, 54, 66, 82	0
17	1V	101/101 (100%)	-0.11	0 100 100	22, 48, 61, 69	0
17	2V	101/101 (100%)	0.07	8 (7%) 12 4	39, 63, 73, 80	0
18	1W	112/113 (99%)	0.42	3 (2%) 54 26	22, 37, 56, 88	0
18	2W	112/113 (99%)	0.35	1 (0%) 84 63	29, 44, 60, 78	0
19	1X	95/96 (98%)	0.11	2 (2%) 63 34	21, 34, 50, 73	0
19	2X	95/96 (98%)	0.33	2 (2%) 63 34	34, 50, 65, 74	0
20	1Y	107/110 (97%)	-0.16	0 100 100	31, 47, 64, 71	0
20	2Y	107/110 (97%)	0.49	4 (3%) 41 17	49, 64, 77, 84	0
21	1Z	154/206 (74%)	-0.29	0 100 100	38, 56, 70, 85	0
21	2Z	160/206 (77%)	-0.25	1 (0%) 89 72	50, 66, 85, 96	0
22	10	83/85 (97%)	0.72	7 (8%) 11 3	26, 34, 68, 92	0
22	20	83/85 (97%)	1.32	17 (20%) 1 0	37, 51, 73, 87	0
23	11	97/98 (98%)	0.68	6 (6%) 20 7	17, 39, 63, 67	0
23	21	97/98 (98%)	0.84	9 (9%) 8 3	27, 50, 67, 75	0
24	12	70/72 (97%)	-0.30	0 100 100	26, 41, 56, 62	0
24	22	70/72 (97%)	-0.13	0 100 100	43, 59, 71, 77	0
25	13	59/60 (98%)	0.13	0 100 100	27, 39, 59, 63	0
25	23	59/60 (98%)	1.26	14 (23%) 0 0	47, 56, 72, 75	0
26	14	69/71 (97%)	-0.23	1 (1%) 75 49	53, 74, 87, 101	0
26	24	69/71 (97%)	0.27	7 (10%) 7 2	61, 82, 95, 102	0
27	15	59/60 (98%)	0.44	3 (5%) 28 10	23, 44, 75, 81	0
27	25	59/60 (98%)	0.27	2 (3%) 45 19	28, 48, 77, 89	0
28	16	53/54 (98%)	-0.30	0 100 100	30, 39, 51, 56	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
28	26	53/54 (98%)	-0.16	0 100 100	40, 48, 59, 61	0
29	17	48/49 (97%)	0.74	5 (10%) 6 2	16, 24, 49, 53	0
29	27	48/49 (97%)	0.65	3 (6%) 20 6	26, 33, 51, 67	0
30	18	64/65 (98%)	0.60	5 (7%) 13 4	23, 30, 39, 58	0
30	28	64/65 (98%)	1.58	20 (31%) 0 0	35, 45, 56, 73	0
31	19	37/37 (100%)	0.21	0 100 100	31, 45, 54, 65	0
31	29	37/37 (100%)	1.80	14 (37%) 0 0	46, 61, 74, 76	0
32	1a	1488/1521 (97%)	0.08	19 (1%) 77 51	27, 65, 93, 112	0
32	2a	1491/1521 (98%)	0.13	30 (2%) 65 36	43, 69, 93, 110	0
33	1b	231/256 (90%)	-0.21	1 (0%) 92 79	55, 73, 86, 101	0
33	2b	231/256 (90%)	-0.39	1 (0%) 92 79	61, 79, 86, 93	0
34	1c	206/239 (86%)	0.22	6 (2%) 51 23	51, 66, 79, 90	0
34	2c	206/239 (86%)	0.24	14 (6%) 17 5	60, 76, 84, 91	0
35	1d	208/209 (99%)	0.66	20 (9%) 8 2	49, 67, 79, 85	0
35	2d	208/209 (99%)	0.60	19 (9%) 9 3	51, 66, 75, 82	0
36	1e	148/162 (91%)	0.64	15 (10%) 7 2	43, 57, 68, 71	0
36	2e	148/162 (91%)	0.67	15 (10%) 7 2	47, 67, 77, 81	0
37	1f	100/101 (99%)	-0.16	0 100 100	42, 61, 71, 75	0
37	2f	100/101 (99%)	-0.39	0 100 100	53, 66, 74, 80	0
38	1g	155/156 (99%)	0.29	12 (7%) 13 4	53, 66, 83, 104	0
38	2g	155/156 (99%)	0.72	27 (17%) 1 0	63, 73, 83, 91	0
39	1h	137/138 (99%)	0.13	2 (1%) 73 46	50, 61, 71, 74	0
39	2h	137/138 (99%)	0.97	25 (18%) 1 0	54, 69, 77, 84	0
40	1i	127/128 (99%)	1.00	32 (25%) 0 0	54, 69, 80, 85	0
40	2i	127/128 (99%)	2.14	57 (44%) 0 0	66, 80, 88, 94	0
41	1j	97/105 (92%)	0.62	19 (19%) 1 0	53, 68, 81, 85	0
41	2j	96/105 (91%)	0.89	21 (21%) 0 0	60, 77, 87, 97	0
42	1k	114/129 (88%)	0.16	1 (0%) 84 63	37, 60, 75, 79	0
42	2k	114/129 (88%)	0.23	4 (3%) 44 18	51, 65, 74, 81	0
43	1l	121/132 (91%)	0.39	14 (11%) 4 1	42, 52, 64, 73	0
43	2l	121/132 (91%)	0.50	12 (9%) 7 2	45, 58, 68, 78	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
44	1m	122/126 (96%)	0.16	11 (9%) 9 3	55, 70, 79, 84	0
44	2m	121/126 (96%)	0.59	17 (14%) 2 1	62, 75, 85, 92	0
45	1n	60/61 (98%)	1.85	28 (46%) 0 0	57, 66, 75, 79	0
45	2n	60/61 (98%)	2.01	28 (46%) 0 0	66, 75, 83, 88	0
46	1o	88/89 (98%)	-0.06	0 100 100	43, 58, 70, 82	0
46	2o	88/89 (98%)	-0.20	2 (2%) 60 31	55, 66, 75, 80	0
47	1p	82/88 (93%)	0.94	16 (19%) 1 0	50, 68, 80, 86	0
47	2p	82/88 (93%)	0.25	2 (2%) 59 30	51, 64, 71, 81	0
48	1q	99/105 (94%)	0.95	16 (16%) 1 0	47, 61, 71, 74	0
48	2q	99/105 (94%)	0.90	16 (16%) 1 0	50, 62, 72, 75	0
49	1r	68/88 (77%)	0.03	1 (1%) 73 46	44, 62, 72, 83	0
49	2r	68/88 (77%)	0.03	0 100 100	51, 67, 76, 80	0
50	1s	83/93 (89%)	0.06	2 (2%) 59 30	59, 70, 80, 91	0
50	2s	83/93 (89%)	0.59	11 (13%) 3 1	67, 77, 86, 90	0
51	1t	96/106 (90%)	0.82	17 (17%) 1 0	53, 67, 78, 81	0
51	2t	96/106 (90%)	0.49	7 (7%) 15 4	45, 62, 75, 83	0
52	1u	23/27 (85%)	0.97	4 (17%) 1 0	58, 65, 72, 79	0
52	2u	23/27 (85%)	1.62	6 (26%) 0 0	65, 71, 78, 79	0
53	1v	10/24 (41%)	0.98	2 (20%) 1 0	46, 82, 94, 95	0
53	2v	6/24 (25%)	0.72	2 (33%) 0 0	56, 66, 85, 85	0
54	1x	72/77 (93%)	-0.13	1 (1%) 75 49	34, 52, 70, 87	0
54	2x	72/77 (93%)	-0.16	1 (1%) 75 49	44, 68, 80, 93	0
All	All	20598/21444 (96%)	0.21	847 (4%) 37 14	11, 57, 84, 116	0

The worst 5 of 847 RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
22	10	7	LEU	11.9
22	10	6	GLY	10.1
22	20	3	HIS	8.8
44	2m	120	LYS	8.2
44	2m	122	LYS	8.2

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. The B-factors column lists the minimum, median, 95th percentile and maximum values of B factors of atoms in the group. The column labelled 'Q< 0.9' lists the number of atoms with occupancy less than 0.9.

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(Å ²)	Q<0.9
1	5MU	2A	1915	21/22	0.90	0.18	55,72,85,88	0
54	PSU	2x	55	20/21	0.90	0.15	64,72,80,82	0
32	7MG	2a	527	24/25	0.91	0.22	47,63,71,72	0
32	PSU	1a	516	20/21	0.91	0.16	53,70,76,77	0
54	PSU	1x	55	20/21	0.91	0.15	49,59,66,70	0
32	M2G	2a	966	25/26	0.92	0.36	65,73,80,82	0
32	UR3	2a	1498	21/22	0.93	0.23	31,51,56,64	0
54	4SU	2x	8	20/21	0.93	0.14	63,68,73,78	0
32	5MC	2a	967	21/22	0.93	0.28	65,71,77,81	0
32	2MG	2a	1207	24/25	0.93	0.15	69,74,78,93	0
32	4OC	2a	1402	22/23	0.93	0.23	54,61,66,69	0
54	5MU	2x	54	21/22	0.93	0.16	70,76,87,97	0
54	5MU	1x	54	21/22	0.94	0.14	45,63,65,67	0
43	0TD	2l	92	10/11	0.94	0.35	55,59,62,66	0
54	4SU	1x	8	20/21	0.94	0.17	44,55,61,70	0
1	PSU	2A	1911	20/21	0.94	0.15	51,55,79,79	0
32	5MC	2a	1400	21/22	0.94	0.28	60,73,79,95	0
32	2MG	1a	1207	24/25	0.94	0.13	63,68,73,79	0
1	PSU	1A	1917	20/21	0.94	0.18	48,59,63,65	0
32	PSU	2a	516	20/21	0.94	0.15	61,69,73,76	0
54	5MC	2x	32	21/22	0.94	0.23	57,63,70,71	0
32	MA6	1a	1518	24/25	0.95	0.23	27,35,38,40	0
32	M2G	1a	966	25/26	0.95	0.27	45,52,57,60	0
32	4OC	1a	1402	22/23	0.95	0.25	35,46,53,58	0
32	MA6	2a	1518	24/25	0.95	0.25	44,53,61,64	0
1	5MU	1A	1915	21/22	0.95	0.15	53,60,70,73	0
1	PSU	1A	1911	20/21	0.95	0.17	40,51,56,57	0
32	5MC	1a	1400	21/22	0.95	0.20	43,50,56,59	0
32	MA6	2a	1519	24/25	0.95	0.35	46,53,56,65	0
1	5MC	2A	1962	21/22	0.95	0.17	31,40,49,54	0
32	5MC	1a	967	21/22	0.95	0.34	47,57,63,65	0
1	4OC	1A	1920	21/23	0.95	0.22	27,44,55,57	0
54	5MC	1x	32	21/22	0.96	0.19	37,52,60,64	0
1	5MC	2A	1942	21/22	0.96	0.15	36,49,55,57	0
32	5MC	2a	1404	21/22	0.96	0.24	47,58,62,62	0
32	MA6	1a	1519	24/25	0.96	0.24	29,36,41,42	0
32	UR3	1a	1498	21/22	0.96	0.19	23,37,42,47	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
1	PSU	2A	2605	20/21	0.96	0.24	23,31,35,36	0
1	OMG	2A	2251	24/25	0.96	0.24	27,33,36,37	0
1	5MU	2A	1939	21/22	0.96	0.17	30,37,45,48	0
1	2MA	2A	2503	23/24	0.96	0.23	20,26,36,41	0
32	5MC	2a	1407	21/22	0.96	0.22	41,52,56,58	0
32	5MC	1a	1407	21/22	0.96	0.18	32,42,49,52	0
1	4OC	2A	1920	21/23	0.96	0.19	44,56,61,64	0
1	PSU	2A	1917	20/21	0.97	0.13	45,56,60,60	0
1	2MU	2A	2552	21/23	0.97	0.18	25,33,39,42	0
1	5MC	1A	1942	21/22	0.97	0.18	22,36,41,46	0
1	5MC	1A	1962	21/22	0.97	0.19	25,37,42,46	0
32	7MG	1a	527	24/25	0.97	0.21	34,39,51,55	0
1	5MU	1A	1939	21/22	0.97	0.19	19,25,29,32	0
1	2MU	1A	2552	21/23	0.97	0.24	26,37,41,43	0
43	0TD	1l	92	10/11	0.97	0.20	43,49,50,55	0
32	5MC	1a	1404	21/22	0.97	0.18	29,38,43,47	0
1	2MA	1A	2503	23/24	0.97	0.24	12,19,25,31	0
1	PSU	1A	2605	20/21	0.97	0.18	19,24,28,28	0
1	OMG	1A	2251	24/25	0.97	0.25	18,23,30,34	0

6.3 Carbohydrates [i](#)

There are no monosaccharides in this entry.

6.4 Ligands [i](#)

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

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	4012	1/1	0.38	0.28	47,47,47,47	0
55	MG	2A	3272	1/1	0.49	0.21	56,56,56,56	0
55	MG	1a	3102	1/1	0.52	0.18	61,61,61,61	0
55	MG	2V	201	1/1	0.57	0.21	53,53,53,53	0
55	MG	1A	3952	1/1	0.58	0.26	68,68,68,68	0
57	ZN	25	105	1/1	0.59	0.12	164,164,164,164	0
55	MG	1a	3061	1/1	0.60	0.86	61,61,61,61	0
55	MG	1A	3528	1/1	0.62	0.22	59,59,59,59	0
55	MG	1v	102	1/1	0.64	0.09	76,76,76,76	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
57	ZN	14	102	1/1	0.64	0.18	159,159,159,159	0
55	MG	2a	1687	1/1	0.65	0.15	44,44,44,44	0
55	MG	2T	203	1/1	0.65	0.23	51,51,51,51	0
55	MG	1A	3249	1/1	0.66	0.19	49,49,49,49	0
55	MG	1A	3296	1/1	0.66	0.28	48,48,48,48	0
55	MG	2A	3115	1/1	0.67	0.20	47,47,47,47	0
55	MG	1a	3066	1/1	0.68	0.18	62,62,62,62	0
55	MG	2a	1790	1/1	0.70	0.88	62,62,62,62	0
55	MG	1A	3354	1/1	0.70	0.21	37,37,37,37	0
55	MG	2A	3027	1/1	0.70	0.23	46,46,46,46	0
55	MG	1Q	205	1/1	0.71	0.53	37,37,37,37	0
55	MG	1A	3079	1/1	0.72	0.21	46,46,46,46	0
55	MG	1F	308	1/1	0.72	0.31	51,51,51,51	0
55	MG	1a	3027	1/1	0.72	0.17	45,45,45,45	0
55	MG	1P	205	1/1	0.72	0.36	56,56,56,56	0
55	MG	2A	3577	1/1	0.72	0.20	69,69,69,69	0
55	MG	1A	3842	1/1	0.74	0.86	28,28,28,28	0
55	MG	1A	3290	1/1	0.74	0.19	39,39,39,39	0
55	MG	2A	3294	1/1	0.74	0.82	33,33,33,33	0
55	MG	2A	3200	1/1	0.74	0.13	57,57,57,57	0
55	MG	2A	3787	1/1	0.74	0.29	57,57,57,57	0
55	MG	2A	3187	1/1	0.74	0.25	62,62,62,62	0
55	MG	1A	3884	1/1	0.75	0.30	68,68,68,68	0
55	MG	1A	3217	1/1	0.75	0.21	48,48,48,48	0
55	MG	2A	3751	1/1	0.75	0.21	50,50,50,50	0
55	MG	1A	3553	1/1	0.75	0.17	27,27,27,27	0
55	MG	1A	3883	1/1	0.75	0.15	55,55,55,55	0
55	MG	13	102	1/1	0.75	0.29	59,59,59,59	0
55	MG	2d	301	1/1	0.75	0.13	38,38,38,38	0
55	MG	2A	3242	1/1	0.75	0.17	62,62,62,62	0
55	MG	1A	3815	1/1	0.76	0.37	31,31,31,31	0
55	MG	2a	1650	1/1	0.76	0.21	49,49,49,49	0
55	MG	1A	3370	1/1	0.76	0.57	35,35,35,35	0
55	MG	2a	1665	1/1	0.76	0.20	57,57,57,57	0
55	MG	2B	207	1/1	0.76	0.21	60,60,60,60	0
55	MG	2a	1714	1/1	0.76	0.27	50,50,50,50	0
55	MG	1E	301	1/1	0.76	0.32	57,57,57,57	0
55	MG	2A	3558	1/1	0.76	0.12	28,28,28,28	0
55	MG	1A	3996	1/1	0.77	0.13	51,51,51,51	0
55	MG	2a	1795	1/1	0.77	0.29	65,65,65,65	0
55	MG	2A	3262	1/1	0.77	0.17	50,50,50,50	0
55	MG	2q	202	1/1	0.77	0.18	55,55,55,55	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
57	ZN	24	501	1/1	0.77	0.09	147,147,147,147	0
55	MG	2A	3225	1/1	0.77	0.35	42,42,42,42	0
55	MG	2A	3232	1/1	0.77	0.77	43,43,43,43	0
55	MG	1x	103	1/1	0.77	0.14	50,50,50,50	0
55	MG	2a	1653	1/1	0.77	0.11	42,42,42,42	0
55	MG	1a	3058	1/1	0.77	0.18	48,48,48,48	0
55	MG	1A	3048	1/1	0.77	0.24	33,33,33,33	0
55	MG	1A	3813	1/1	0.77	0.12	44,44,44,44	0
55	MG	1A	3304	1/1	0.77	0.43	35,35,35,35	0
55	MG	1a	3050	1/1	0.77	0.33	25,25,25,25	0
55	MG	1B	226	1/1	0.78	0.16	53,53,53,53	0
55	MG	1A	3887	1/1	0.78	0.23	55,55,55,55	0
55	MG	1a	3177	1/1	0.78	0.28	69,69,69,69	0
55	MG	1D	303	1/1	0.78	0.17	31,31,31,31	0
55	MG	1A	3418	1/1	0.78	0.15	36,36,36,36	0
55	MG	2a	1717	1/1	0.78	0.24	68,68,68,68	0
55	MG	2a	1746	1/1	0.78	0.20	40,40,40,40	0
55	MG	2A	3160	1/1	0.78	0.10	58,58,58,58	0
55	MG	2A	3132	1/1	0.78	0.38	38,38,38,38	0
55	MG	2a	1723	1/1	0.79	0.13	65,65,65,65	0
55	MG	2A	3539	1/1	0.79	0.57	41,41,41,41	0
55	MG	2A	3278	1/1	0.79	0.53	36,36,36,36	0
55	MG	1n	102	1/1	0.79	0.39	55,55,55,55	0
55	MG	2A	3361	1/1	0.79	0.15	50,50,50,50	0
55	MG	1a	3176	1/1	0.79	0.09	71,71,71,71	0
55	MG	1A	3582	1/1	0.79	0.11	45,45,45,45	0
55	MG	1x	102	1/1	0.79	0.56	80,80,80,80	0
55	MG	1a	3193	1/1	0.79	0.16	62,62,62,62	0
55	MG	1a	3172	1/1	0.79	0.48	67,67,67,67	0
55	MG	2a	1806	1/1	0.79	0.45	55,55,55,55	0
55	MG	1O	203	1/1	0.79	0.40	58,58,58,58	0
55	MG	1a	3032	1/1	0.79	0.21	57,57,57,57	0
55	MG	2a	1704	1/1	0.79	0.22	47,47,47,47	0
55	MG	2A	3380	1/1	0.79	0.21	49,49,49,49	0
55	MG	2A	3195	1/1	0.79	0.71	42,42,42,42	0
55	MG	2A	3676	1/1	0.79	0.12	65,65,65,65	0
55	MG	2A	3048	1/1	0.80	0.26	57,57,57,57	0
55	MG	2A	3851	1/1	0.80	0.69	47,47,47,47	0
55	MG	2a	1622	1/1	0.80	0.13	63,63,63,63	0
55	MG	1A	3881	1/1	0.80	0.20	44,44,44,44	0
55	MG	2A	3673	1/1	0.80	0.17	56,56,56,56	0
55	MG	1A	3722	1/1	0.80	0.13	44,44,44,44	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3541	1/1	0.80	0.15	39,39,39,39	0
55	MG	2a	1730	1/1	0.80	0.17	59,59,59,59	0
55	MG	2A	3246	1/1	0.80	0.46	48,48,48,48	0
55	MG	2A	3171	1/1	0.80	0.13	35,35,35,35	0
55	MG	2A	3576	1/1	0.80	0.20	48,48,48,48	0
55	MG	2A	3158	1/1	0.80	0.20	38,38,38,38	0
55	MG	1A	3312	1/1	0.80	0.48	40,40,40,40	0
55	MG	2A	3717	1/1	0.80	0.17	28,28,28,28	0
55	MG	1A	3655	1/1	0.81	0.48	41,41,41,41	0
55	MG	1A	3015	1/1	0.81	0.12	36,36,36,36	0
55	MG	1A	3695	1/1	0.81	0.12	10,10,10,10	0
55	MG	1A	3166	1/1	0.81	0.25	36,36,36,36	0
55	MG	2a	1662	1/1	0.81	0.62	52,52,52,52	0
55	MG	2A	3663	1/1	0.81	0.46	43,43,43,43	0
55	MG	1A	3935	1/1	0.81	0.13	56,56,56,56	0
55	MG	2A	3234	1/1	0.81	0.18	53,53,53,53	0
55	MG	2A	3418	1/1	0.81	0.11	46,46,46,46	0
55	MG	1a	3198	1/1	0.81	0.14	56,56,56,56	0
55	MG	2A	3455	1/1	0.81	0.14	32,32,32,32	0
55	MG	2A	3502	1/1	0.81	0.21	56,56,56,56	0
55	MG	2A	3381	1/1	0.81	0.78	68,68,68,68	0
55	MG	2A	3814	1/1	0.81	0.26	44,44,44,44	0
55	MG	1A	3962	1/1	0.81	0.45	53,53,53,53	0
55	MG	2a	1641	1/1	0.81	0.32	51,51,51,51	0
55	MG	2a	1604	1/1	0.81	0.14	47,47,47,47	0
55	MG	2a	1659	1/1	0.81	0.14	45,45,45,45	0
55	MG	1B	218	1/1	0.81	0.13	41,41,41,41	0
55	MG	2A	3722	1/1	0.82	0.17	25,25,25,25	0
55	MG	1A	3840	1/1	0.82	0.31	28,28,28,28	0
55	MG	2a	1610	1/1	0.82	0.13	55,55,55,55	0
55	MG	2A	3150	1/1	0.82	0.22	45,45,45,45	0
55	MG	1A	3162	1/1	0.82	0.14	36,36,36,36	0
55	MG	2a	1739	1/1	0.82	0.23	83,83,83,83	0
55	MG	1A	3612	1/1	0.82	0.17	14,14,14,14	0
55	MG	2a	1774	1/1	0.82	0.13	29,29,29,29	0
55	MG	1A	3173	1/1	0.82	0.33	28,28,28,28	0
55	MG	1A	3650	1/1	0.82	0.14	60,60,60,60	0
55	MG	14	101	1/1	0.82	0.18	40,40,40,40	0
55	MG	2A	3137	1/1	0.82	0.83	39,39,39,39	0
55	MG	1A	3423	1/1	0.82	0.16	34,34,34,34	0
55	MG	2A	3289	1/1	0.82	0.30	49,49,49,49	0
55	MG	2a	1740	1/1	0.82	0.26	75,75,75,75	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3793	1/1	0.82	1.61	48,48,48,48	0
55	MG	1A	3421	1/1	0.82	0.20	33,33,33,33	0
55	MG	2A	3321	1/1	0.82	0.51	37,37,37,37	0
55	MG	2A	3214	1/1	0.82	0.29	33,33,33,33	0
55	MG	1A	3740	1/1	0.82	0.11	39,39,39,39	0
55	MG	1A	3356	1/1	0.82	0.39	29,29,29,29	0
55	MG	2A	3569	1/1	0.82	0.12	35,35,35,35	0
55	MG	2A	3718	1/1	0.82	0.17	62,62,62,62	0
55	MG	2A	3169	1/1	0.82	0.20	40,40,40,40	0
55	MG	2A	3396	1/1	0.82	0.18	51,51,51,51	0
55	MG	1A	3471	1/1	0.82	0.48	47,47,47,47	0
55	MG	2a	1617	1/1	0.82	0.18	46,46,46,46	0
55	MG	2A	3662	1/1	0.82	0.26	39,39,39,39	0
55	MG	1a	3071	1/1	0.83	0.15	37,37,37,37	0
55	MG	2A	3277	1/1	0.83	0.43	42,42,42,42	0
55	MG	1B	208	1/1	0.83	0.13	43,43,43,43	0
55	MG	1A	3841	1/1	0.83	0.25	41,41,41,41	0
55	MG	2B	211	1/1	0.83	0.24	35,35,35,35	0
55	MG	1a	3096	1/1	0.83	0.19	42,42,42,42	0
55	MG	1B	224	1/1	0.83	0.16	40,40,40,40	0
55	MG	1P	203	1/1	0.83	0.41	44,44,44,44	0
55	MG	2A	3493	1/1	0.83	0.41	42,42,42,42	0
55	MG	2A	3151	1/1	0.83	0.23	56,56,56,56	0
55	MG	2A	3646	1/1	0.83	0.18	54,54,54,54	0
55	MG	2A	3350	1/1	0.83	0.34	38,38,38,38	0
55	MG	2A	3061	1/1	0.83	0.12	50,50,50,50	0
55	MG	2A	3377	1/1	0.83	0.15	49,49,49,49	0
55	MG	2A	3264	1/1	0.83	0.34	52,52,52,52	0
55	MG	1b	301	1/1	0.83	0.12	54,54,54,54	0
55	MG	25	104	1/1	0.83	0.38	34,34,34,34	0
55	MG	1A	3059	1/1	0.83	0.24	35,35,35,35	0
55	MG	1B	203	1/1	0.83	0.18	51,51,51,51	0
55	MG	2A	3707	1/1	0.83	0.13	44,44,44,44	0
55	MG	17	103	1/1	0.83	0.53	41,41,41,41	0
55	MG	1A	3914	1/1	0.83	0.46	62,62,62,62	0
55	MG	23	102	1/1	0.83	0.62	52,52,52,52	0
55	MG	2a	1608	1/1	0.83	0.67	53,53,53,53	0
55	MG	2A	3327	1/1	0.83	0.26	43,43,43,43	0
55	MG	2A	3068	1/1	0.83	0.67	32,32,32,32	0
55	MG	2A	3075	1/1	0.84	0.12	30,30,30,30	0
55	MG	1B	229	1/1	0.84	0.18	75,75,75,75	0
55	MG	1O	201	1/1	0.84	0.55	40,40,40,40	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2f	202	1/1	0.84	0.15	52,52,52,52	0
55	MG	2A	3471	1/1	0.84	0.14	43,43,43,43	0
55	MG	1A	3317	1/1	0.84	0.55	24,24,24,24	0
55	MG	1a	3171	1/1	0.84	0.31	59,59,59,59	0
55	MG	2a	1632	1/1	0.84	0.21	51,51,51,51	0
55	MG	1a	3051	1/1	0.84	0.15	78,78,78,78	0
55	MG	1A	3308	1/1	0.84	0.15	45,45,45,45	0
55	MG	1A	3310	1/1	0.84	0.19	36,36,36,36	0
55	MG	1a	3210	1/1	0.84	0.18	34,34,34,34	0
55	MG	1A	3145	1/1	0.84	0.43	45,45,45,45	0
55	MG	2A	3215	1/1	0.84	0.18	24,24,24,24	0
55	MG	2A	3126	1/1	0.84	0.24	46,46,46,46	0
55	MG	2A	3566	1/1	0.84	0.19	25,25,25,25	0
55	MG	2A	3540	1/1	0.84	0.16	32,32,32,32	0
55	MG	1A	3164	1/1	0.84	0.54	31,31,31,31	0
55	MG	25	102	1/1	0.84	0.32	28,28,28,28	0
55	MG	1A	3512	1/1	0.84	0.16	23,23,23,23	0
55	MG	2A	3255	1/1	0.84	1.19	51,51,51,51	0
57	ZN	2Y	203	1/1	0.84	0.08	98,98,98,98	0
55	MG	2A	3422	1/1	0.84	0.22	52,52,52,52	0
55	MG	2A	3619	1/1	0.84	0.17	42,42,42,42	0
55	MG	2A	3572	1/1	0.84	0.47	50,50,50,50	0
55	MG	1A	3359	1/1	0.84	0.18	33,33,33,33	0
55	MG	1a	3028	1/1	0.84	0.26	52,52,52,52	0
55	MG	2A	3549	1/1	0.84	0.47	61,61,61,61	0
55	MG	2A	3341	1/1	0.84	0.13	43,43,43,43	0
55	MG	2A	3546	1/1	0.84	0.33	54,54,54,54	0
55	MG	2A	3267	1/1	0.84	0.16	54,54,54,54	0
55	MG	1A	3910	1/1	0.84	0.13	35,35,35,35	0
55	MG	1A	3569	1/1	0.85	0.14	28,28,28,28	0
55	MG	2D	301	1/1	0.85	0.51	42,42,42,42	0
55	MG	1A	3969	1/1	0.85	0.11	45,45,45,45	0
55	MG	2A	3489	1/1	0.85	0.14	17,17,17,17	0
55	MG	1A	3332	1/1	0.85	0.32	47,47,47,47	0
55	MG	1A	3865	1/1	0.85	0.17	49,49,49,49	0
55	MG	2A	3600	1/1	0.85	0.15	37,37,37,37	0
55	MG	2A	3199	1/1	0.85	0.14	38,38,38,38	0
55	MG	1a	3060	1/1	0.85	0.10	44,44,44,44	0
55	MG	1A	3662	1/1	0.85	0.13	72,72,72,72	0
55	MG	2A	3301	1/1	0.85	0.24	44,44,44,44	0
55	MG	1A	3912	1/1	0.85	0.69	39,39,39,39	0
55	MG	1A	3530	1/1	0.85	0.15	42,42,42,42	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1a	3155	1/1	0.85	0.11	49,49,49,49	0
55	MG	10	104	1/1	0.85	0.21	46,46,46,46	0
55	MG	1B	233	1/1	0.85	0.11	42,42,42,42	0
55	MG	2A	3292	1/1	0.85	0.48	48,48,48,48	0
55	MG	2a	1605	1/1	0.85	0.47	42,42,42,42	0
55	MG	2A	3649	1/1	0.85	0.13	47,47,47,47	0
55	MG	1E	309	1/1	0.85	0.49	36,36,36,36	0
55	MG	1A	3262	1/1	0.85	0.15	27,27,27,27	0
55	MG	2A	3625	1/1	0.85	0.12	24,24,24,24	0
55	MG	1A	3182	1/1	0.85	0.27	42,42,42,42	0
55	MG	2A	3618	1/1	0.85	0.17	22,22,22,22	0
55	MG	2A	3088	1/1	0.85	0.15	45,45,45,45	0
55	MG	2a	1611	1/1	0.85	0.19	47,47,47,47	0
55	MG	1A	3362	1/1	0.85	0.56	44,44,44,44	0
55	MG	2a	1607	1/1	0.85	0.25	31,31,31,31	0
55	MG	2A	3202	1/1	0.85	0.18	45,45,45,45	0
55	MG	1A	3618	1/1	0.85	0.12	46,46,46,46	0
55	MG	2A	3312	1/1	0.85	0.58	52,52,52,52	0
55	MG	2A	3840	1/1	0.85	0.18	51,51,51,51	0
55	MG	1A	3250	1/1	0.85	0.32	46,46,46,46	0
55	MG	2A	3482	1/1	0.85	0.16	35,35,35,35	0
55	MG	2A	3838	1/1	0.85	0.13	49,49,49,49	0
55	MG	1B	219	1/1	0.85	0.22	40,40,40,40	0
55	MG	2A	3224	1/1	0.85	0.22	34,34,34,34	0
55	MG	2A	3497	1/1	0.85	0.11	51,51,51,51	0
55	MG	2a	1618	1/1	0.85	0.15	40,40,40,40	0
55	MG	27	101	1/1	0.86	0.15	31,31,31,31	0
55	MG	1A	3567	1/1	0.86	0.16	27,27,27,27	0
55	MG	2A	3008	1/1	0.86	0.43	24,24,24,24	0
55	MG	2a	1614	1/1	0.86	0.23	53,53,53,53	0
55	MG	2f	201	1/1	0.86	0.10	44,44,44,44	0
55	MG	2A	3411	1/1	0.86	0.10	43,43,43,43	0
55	MG	2A	3819	1/1	0.86	0.14	21,21,21,21	0
55	MG	2A	3367	1/1	0.86	0.31	27,27,27,27	0
55	MG	2A	3265	1/1	0.86	0.14	37,37,37,37	0
55	MG	2a	1791	1/1	0.86	0.09	66,66,66,66	0
55	MG	1A	3306	1/1	0.86	0.13	38,38,38,38	0
55	MG	2A	3769	1/1	0.86	0.18	69,69,69,69	0
55	MG	2a	1710	1/1	0.86	0.21	57,57,57,57	0
55	MG	1A	3436	1/1	0.86	0.11	53,53,53,53	0
55	MG	2A	3737	1/1	0.86	0.20	51,51,51,51	0
55	MG	2a	1670	1/1	0.86	0.16	49,49,49,49	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1W	208	1/1	0.86	0.45	27,27,27,27	0
55	MG	1A	3371	1/1	0.86	0.15	45,45,45,45	0
55	MG	1a	3065	1/1	0.86	0.18	51,51,51,51	0
55	MG	2A	3835	1/1	0.86	0.11	42,42,42,42	0
55	MG	1E	312	1/1	0.86	1.08	54,54,54,54	0
55	MG	2a	1815	1/1	0.86	0.23	53,53,53,53	0
55	MG	19	101	1/1	0.86	0.16	58,58,58,58	0
55	MG	2A	3434	1/1	0.86	0.09	15,15,15,15	0
55	MG	2A	3141	1/1	0.86	0.17	31,31,31,31	0
55	MG	1a	3197	1/1	0.86	0.46	61,61,61,61	0
55	MG	1A	3083	1/1	0.86	0.20	42,42,42,42	0
55	MG	2A	3412	1/1	0.86	0.36	45,45,45,45	0
55	MG	1A	3458	1/1	0.86	0.25	25,25,25,25	0
55	MG	1A	3443	1/1	0.86	0.14	39,39,39,39	0
55	MG	2a	1609	1/1	0.86	0.15	39,39,39,39	0
55	MG	1A	3190	1/1	0.86	0.25	41,41,41,41	0
55	MG	1A	3516	1/1	0.86	0.35	31,31,31,31	0
55	MG	1a	3184	1/1	0.86	0.26	55,55,55,55	0
55	MG	2A	3145	1/1	0.86	0.09	51,51,51,51	0
55	MG	2a	1802	1/1	0.86	0.10	58,58,58,58	0
55	MG	2A	3607	1/1	0.86	0.28	44,44,44,44	0
55	MG	2A	3286	1/1	0.86	0.45	33,33,33,33	0
55	MG	2a	1699	1/1	0.86	0.16	36,36,36,36	0
55	MG	2a	1703	1/1	0.86	0.15	38,38,38,38	0
55	MG	1A	3919	1/1	0.86	0.19	33,33,33,33	0
55	MG	2A	3347	1/1	0.86	0.17	48,48,48,48	0
55	MG	2A	3683	1/1	0.86	0.19	47,47,47,47	0
55	MG	2A	3487	1/1	0.86	0.17	44,44,44,44	0
55	MG	1A	4005	1/1	0.86	0.14	47,47,47,47	0
55	MG	2P	201	1/1	0.86	0.43	40,40,40,40	0
55	MG	2a	1698	1/1	0.86	0.40	40,40,40,40	0
55	MG	1A	3591	1/1	0.86	0.10	29,29,29,29	0
55	MG	1A	3903	1/1	0.86	0.88	37,37,37,37	0
55	MG	2A	3316	1/1	0.86	0.13	36,36,36,36	0
55	MG	2a	1705	1/1	0.87	0.29	54,54,54,54	0
55	MG	1A	3853	1/1	0.87	0.14	39,39,39,39	0
55	MG	2a	1725	1/1	0.87	0.14	59,59,59,59	0
55	MG	1A	3268	1/1	0.87	0.14	42,42,42,42	0
55	MG	2A	3738	1/1	0.87	0.17	27,27,27,27	0
55	MG	2A	3407	1/1	0.87	0.14	50,50,50,50	0
55	MG	1A	3473	1/1	0.87	0.67	27,27,27,27	0
55	MG	1A	3094	1/1	0.87	1.01	33,33,33,33	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	4031	1/1	0.87	0.21	21,21,21,21	0
55	MG	1A	3297	1/1	0.87	0.44	40,40,40,40	0
55	MG	1A	3236	1/1	0.87	0.74	28,28,28,28	0
55	MG	2a	1628	1/1	0.87	0.16	41,41,41,41	0
55	MG	2a	1761	1/1	0.87	0.16	49,49,49,49	0
55	MG	1E	313	1/1	0.87	0.17	43,43,43,43	0
55	MG	1A	3040	1/1	0.87	0.28	36,36,36,36	0
55	MG	2A	3249	1/1	0.87	0.13	25,25,25,25	0
55	MG	1A	3439	1/1	0.87	0.21	26,26,26,26	0
55	MG	1A	3121	1/1	0.87	0.32	27,27,27,27	0
55	MG	2A	3680	1/1	0.87	0.12	46,46,46,46	0
55	MG	1A	3972	1/1	0.87	0.81	37,37,37,37	0
55	MG	2a	1733	1/1	0.87	0.08	44,44,44,44	0
55	MG	1a	3100	1/1	0.87	0.18	31,31,31,31	0
55	MG	2A	3849	1/1	0.87	0.26	50,50,50,50	0
55	MG	1A	3358	1/1	0.87	0.99	35,35,35,35	0
55	MG	1A	3523	1/1	0.87	0.14	42,42,42,42	0
55	MG	2A	3660	1/1	0.87	0.34	38,38,38,38	0
55	MG	2A	3168	1/1	0.87	0.24	47,47,47,47	0
55	MG	2A	3288	1/1	0.87	0.20	30,30,30,30	0
55	MG	13	103	1/1	0.87	0.41	53,53,53,53	0
55	MG	2A	3236	1/1	0.87	0.61	38,38,38,38	0
55	MG	2A	3058	1/1	0.87	0.17	47,47,47,47	0
55	MG	1a	3035	1/1	0.87	0.19	52,52,52,52	0
55	MG	2a	1625	1/1	0.87	0.58	43,43,43,43	0
55	MG	1a	3075	1/1	0.87	0.45	43,43,43,43	0
55	MG	2A	3730	1/1	0.87	0.65	54,54,54,54	0
55	MG	2A	3332	1/1	0.87	0.10	33,33,33,33	0
55	MG	2F	301	1/1	0.87	0.15	28,28,28,28	0
55	MG	1a	3191	1/1	0.87	0.12	62,62,62,62	0
55	MG	1A	4014	1/1	0.87	0.79	51,51,51,51	0
55	MG	2A	3308	1/1	0.87	0.60	44,44,44,44	0
55	MG	2A	3078	1/1	0.87	0.84	41,41,41,41	0
55	MG	2A	3186	1/1	0.87	0.17	43,43,43,43	0
55	MG	1G	202	1/1	0.87	0.48	40,40,40,40	0
55	MG	1A	3849	1/1	0.87	0.65	22,22,22,22	0
55	MG	2A	3420	1/1	0.87	0.12	38,38,38,38	0
55	MG	1A	3623	1/1	0.87	0.13	15,15,15,15	0
55	MG	1A	3226	1/1	0.87	0.17	46,46,46,46	0
55	MG	2A	3406	1/1	0.87	0.16	30,30,30,30	0
55	MG	1A	3889	1/1	0.87	0.08	45,45,45,45	0
55	MG	1A	3746	1/1	0.87	0.33	51,51,51,51	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1a	3076	1/1	0.87	0.38	59,59,59,59	0
55	MG	1a	3029	1/1	0.87	0.11	30,30,30,30	0
55	MG	1A	3861	1/1	0.87	0.23	28,28,28,28	0
55	MG	1A	3323	1/1	0.87	0.16	30,30,30,30	0
55	MG	2A	3323	1/1	0.87	0.35	38,38,38,38	0
55	MG	1a	3041	1/1	0.88	0.14	45,45,45,45	0
55	MG	2A	3435	1/1	0.88	0.09	24,24,24,24	0
55	MG	2A	3805	1/1	0.88	0.15	46,46,46,46	0
55	MG	2A	3727	1/1	0.88	0.31	40,40,40,40	0
55	MG	1A	3376	1/1	0.88	0.17	24,24,24,24	0
55	MG	1A	3326	1/1	0.88	0.35	26,26,26,26	0
55	MG	2A	3207	1/1	0.88	0.41	48,48,48,48	0
55	MG	1A	3406	1/1	0.88	0.25	31,31,31,31	0
55	MG	1a	3093	1/1	0.88	0.20	53,53,53,53	0
55	MG	1a	3187	1/1	0.88	0.34	76,76,76,76	0
55	MG	2A	3074	1/1	0.88	0.21	44,44,44,44	0
55	MG	1A	3762	1/1	0.88	0.15	30,30,30,30	0
55	MG	1A	3133	1/1	0.88	0.10	27,27,27,27	0
55	MG	2a	1624	1/1	0.88	0.18	38,38,38,38	0
55	MG	1A	3026	1/1	0.88	0.12	52,52,52,52	0
55	MG	2A	3345	1/1	0.88	0.60	41,41,41,41	0
55	MG	1a	3047	1/1	0.88	0.11	55,55,55,55	0
55	MG	2a	1645	1/1	0.88	0.47	46,46,46,46	0
55	MG	1A	3943	1/1	0.88	0.19	41,41,41,41	0
55	MG	2A	3728	1/1	0.88	0.15	57,57,57,57	0
55	MG	1A	3587	1/1	0.88	0.11	37,37,37,37	0
55	MG	2a	1616	1/1	0.88	0.13	52,52,52,52	0
55	MG	2A	3759	1/1	0.88	0.18	43,43,43,43	0
55	MG	2A	3319	1/1	0.88	0.32	38,38,38,38	0
55	MG	2A	3349	1/1	0.88	0.18	49,49,49,49	0
55	MG	1A	3104	1/1	0.88	0.91	32,32,32,32	0
55	MG	2A	3201	1/1	0.88	0.20	51,51,51,51	0
55	MG	1x	106	1/1	0.88	0.21	39,39,39,39	0
55	MG	1a	3166	1/1	0.88	0.13	84,84,84,84	0
55	MG	2A	3587	1/1	0.88	0.25	60,60,60,60	0
55	MG	1A	3898	1/1	0.88	0.20	54,54,54,54	0
55	MG	2A	3340	1/1	0.88	0.21	43,43,43,43	0
55	MG	1Q	203	1/1	0.88	0.19	45,45,45,45	0
55	MG	2A	3172	1/1	0.88	0.28	43,43,43,43	0
55	MG	2A	3774	1/1	0.88	0.20	58,58,58,58	0
55	MG	17	102	1/1	0.88	0.33	18,18,18,18	0
55	MG	1A	4036	1/1	0.88	0.17	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3281	1/1	0.88	0.12	12,12,12,12	0
55	MG	2B	212	1/1	0.88	0.21	60,60,60,60	0
55	MG	1A	3967	1/1	0.88	0.14	38,38,38,38	0
55	MG	1A	3143	1/1	0.88	0.38	28,28,28,28	0
55	MG	2a	1776	1/1	0.88	0.11	55,55,55,55	0
55	MG	2a	1762	1/1	0.88	0.13	55,55,55,55	0
55	MG	1B	230	1/1	0.88	0.32	44,44,44,44	0
55	MG	2A	3336	1/1	0.88	0.71	59,59,59,59	0
55	MG	2A	3544	1/1	0.88	0.28	48,48,48,48	0
55	MG	2A	3304	1/1	0.88	0.52	39,39,39,39	0
55	MG	2a	1691	1/1	0.88	0.20	53,53,53,53	0
55	MG	2A	3705	1/1	0.88	0.13	55,55,55,55	0
55	MG	1a	3104	1/1	0.88	0.49	48,48,48,48	0
55	MG	2A	3850	1/1	0.88	0.34	54,54,54,54	0
55	MG	1D	309	1/1	0.88	0.45	21,21,21,21	0
55	MG	1x	104	1/1	0.88	0.12	46,46,46,46	0
55	MG	2A	3428	1/1	0.88	0.35	58,58,58,58	0
55	MG	1a	3004	1/1	0.88	0.25	57,57,57,57	0
55	MG	1A	3820	1/1	0.88	0.15	14,14,14,14	0
55	MG	1A	3518	1/1	0.88	0.72	45,45,45,45	0
55	MG	2a	1782	1/1	0.88	0.16	40,40,40,40	0
55	MG	2A	3409	1/1	0.88	0.12	28,28,28,28	0
55	MG	1B	231	1/1	0.88	0.13	39,39,39,39	0
55	MG	2A	3051	1/1	0.88	0.10	42,42,42,42	0
55	MG	1A	3506	1/1	0.88	0.16	23,23,23,23	0
55	MG	2a	1651	1/1	0.88	0.10	58,58,58,58	0
55	MG	2A	3732	1/1	0.88	0.16	28,28,28,28	0
55	MG	1A	3198	1/1	0.88	0.22	29,29,29,29	0
55	MG	2A	3251	1/1	0.88	0.41	21,21,21,21	0
55	MG	1A	3200	1/1	0.88	0.26	37,37,37,37	0
55	MG	1A	3475	1/1	0.88	0.62	27,27,27,27	0
55	MG	1a	3016	1/1	0.88	0.53	46,46,46,46	0
55	MG	2A	3243	1/1	0.88	0.32	51,51,51,51	0
55	MG	2B	216	1/1	0.88	0.15	57,57,57,57	0
55	MG	2A	3836	1/1	0.88	0.21	39,39,39,39	0
55	MG	2A	3545	1/1	0.88	0.20	57,57,57,57	0
55	MG	2a	1749	1/1	0.88	0.21	53,53,53,53	0
55	MG	1a	3094	1/1	0.88	0.66	41,41,41,41	0
55	MG	2A	3504	1/1	0.88	0.27	37,37,37,37	0
55	MG	1A	3752	1/1	0.88	0.48	24,24,24,24	0
55	MG	1A	3822	1/1	0.88	0.11	26,26,26,26	0
55	MG	2a	1603	1/1	0.88	0.10	43,43,43,43	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2t	201	1/1	0.88	0.14	36,36,36,36	0
55	MG	2F	304	1/1	0.88	0.17	54,54,54,54	0
55	MG	1a	3142	1/1	0.88	0.37	44,44,44,44	0
55	MG	2a	1649	1/1	0.88	0.19	62,62,62,62	0
55	MG	1A	3726	1/1	0.88	0.08	18,18,18,18	0
55	MG	1a	3009	1/1	0.88	0.20	28,28,28,28	0
55	MG	2A	3101	1/1	0.89	0.19	30,30,30,30	0
55	MG	1A	3602	1/1	0.89	0.12	28,28,28,28	0
55	MG	1A	3714	1/1	0.89	0.12	11,11,11,11	0
55	MG	2a	1789	1/1	0.89	0.07	44,44,44,44	0
55	MG	1A	3228	1/1	0.89	0.24	30,30,30,30	0
55	MG	1A	4003	1/1	0.89	0.20	42,42,42,42	0
55	MG	1B	235	1/1	0.89	0.10	38,38,38,38	0
55	MG	2A	3453	1/1	0.89	0.11	31,31,31,31	0
55	MG	1R	207	1/1	0.89	0.43	55,55,55,55	0
55	MG	1A	3184	1/1	0.89	0.45	43,43,43,43	0
55	MG	1a	3054	1/1	0.89	0.23	46,46,46,46	0
55	MG	1A	3030	1/1	0.89	0.68	21,21,21,21	0
55	MG	2a	1794	1/1	0.89	0.12	51,51,51,51	0
55	MG	2A	3155	1/1	0.89	0.13	54,54,54,54	0
55	MG	2A	3113	1/1	0.89	0.11	50,50,50,50	0
55	MG	2A	3764	1/1	0.89	0.17	37,37,37,37	0
55	MG	1A	3498	1/1	0.89	0.12	25,25,25,25	0
55	MG	1a	3062	1/1	0.89	0.14	51,51,51,51	0
55	MG	2A	3087	1/1	0.89	0.13	31,31,31,31	0
55	MG	1A	3189	1/1	0.89	0.43	22,22,22,22	0
55	MG	1A	3957	1/1	0.89	0.50	63,63,63,63	0
55	MG	1A	3965	1/1	0.89	0.20	27,27,27,27	0
55	MG	2A	3658	1/1	0.89	0.42	52,52,52,52	0
55	MG	2A	3584	1/1	0.89	0.11	40,40,40,40	0
55	MG	1W	204	1/1	0.89	0.43	43,43,43,43	0
55	MG	1A	3293	1/1	0.89	0.17	41,41,41,41	0
55	MG	1A	3737	1/1	0.89	0.10	33,33,33,33	0
55	MG	1A	3615	1/1	0.89	0.15	15,15,15,15	0
55	MG	1A	3760	1/1	0.89	0.13	22,22,22,22	0
55	MG	1A	4034	1/1	0.89	0.41	54,54,54,54	0
55	MG	1A	3199	1/1	0.89	0.15	43,43,43,43	0
55	MG	2A	3237	1/1	0.89	0.68	40,40,40,40	0
55	MG	2A	3672	1/1	0.89	0.20	39,39,39,39	0
55	MG	2A	3765	1/1	0.89	0.16	54,54,54,54	0
55	MG	1B	225	1/1	0.89	0.13	49,49,49,49	0
55	MG	1a	3189	1/1	0.89	0.14	44,44,44,44	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1a	3116	1/1	0.89	0.18	32,32,32,32	0
55	MG	1A	3167	1/1	0.89	0.75	30,30,30,30	0
55	MG	1A	3601	1/1	0.89	0.13	47,47,47,47	0
55	MG	1A	3524	1/1	0.89	0.12	35,35,35,35	0
55	MG	2a	1751	1/1	0.89	0.08	34,34,34,34	0
55	MG	2A	3466	1/1	0.89	0.14	45,45,45,45	0
55	MG	1A	3348	1/1	0.89	0.20	28,28,28,28	0
55	MG	1A	3355	1/1	0.89	0.45	37,37,37,37	0
55	MG	1A	3070	1/1	0.89	0.15	41,41,41,41	0
55	MG	2A	3217	1/1	0.89	0.21	50,50,50,50	0
55	MG	1A	3702	1/1	0.89	0.13	21,21,21,21	0
55	MG	2A	3456	1/1	0.89	0.16	38,38,38,38	0
55	MG	2Q	202	1/1	0.89	0.21	38,38,38,38	0
55	MG	2A	3425	1/1	0.89	0.29	42,42,42,42	0
55	MG	2E	305	1/1	0.89	0.15	27,27,27,27	0
55	MG	1A	3433	1/1	0.89	0.21	28,28,28,28	0
55	MG	2X	101	1/1	0.89	0.15	34,34,34,34	0
55	MG	1a	3088	1/1	0.89	0.19	51,51,51,51	0
55	MG	2A	3091	1/1	0.89	0.16	58,58,58,58	0
55	MG	2a	1626	1/1	0.89	0.66	46,46,46,46	0
55	MG	2A	3675	1/1	0.89	0.09	42,42,42,42	0
55	MG	2A	3403	1/1	0.89	0.17	35,35,35,35	0
55	MG	2a	1673	1/1	0.89	0.25	50,50,50,50	0
55	MG	1A	3179	1/1	0.89	0.15	44,44,44,44	0
55	MG	2a	1656	1/1	0.89	0.14	44,44,44,44	0
55	MG	1v	103	1/1	0.89	0.14	55,55,55,55	0
55	MG	1a	3089	1/1	0.89	0.15	49,49,49,49	0
55	MG	2A	3711	1/1	0.89	0.18	47,47,47,47	0
55	MG	1A	3680	1/1	0.89	0.11	37,37,37,37	0
58	V7A	2a	1817	35/35	0.89	0.51	61,80,87,88	0
55	MG	2A	3775	1/1	0.89	0.14	30,30,30,30	0
55	MG	1A	3611	1/1	0.89	0.14	59,59,59,59	0
55	MG	2a	1709	1/1	0.89	0.11	44,44,44,44	0
55	MG	16	103	1/1	0.89	0.56	42,42,42,42	0
57	ZN	15	105	1/1	0.89	0.35	155,155,155,155	0
55	MG	2A	3613	1/1	0.89	0.23	28,28,28,28	0
55	MG	2A	3364	1/1	0.89	0.21	33,33,33,33	0
55	MG	1B	211	1/1	0.89	0.35	39,39,39,39	0
55	MG	1a	3122	1/1	0.89	0.12	42,42,42,42	0
55	MG	2A	3314	1/1	0.89	0.11	39,39,39,39	0
55	MG	1A	3025	1/1	0.89	0.27	51,51,51,51	0
55	MG	2A	3376	1/1	0.89	0.18	33,33,33,33	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3356	1/1	0.90	0.20	36,36,36,36	0
55	MG	2A	3279	1/1	0.90	0.14	33,33,33,33	0
55	MG	2A	3659	1/1	0.90	0.14	41,41,41,41	0
55	MG	2a	1640	1/1	0.90	0.13	42,42,42,42	0
55	MG	2A	3015	1/1	0.90	0.14	55,55,55,55	0
55	MG	2A	3291	1/1	0.90	0.15	39,39,39,39	0
55	MG	2A	3424	1/1	0.90	0.11	40,40,40,40	0
55	MG	2A	3090	1/1	0.90	0.16	40,40,40,40	0
55	MG	2A	3162	1/1	0.90	0.23	37,37,37,37	0
55	MG	1A	3125	1/1	0.90	0.14	29,29,29,29	0
55	MG	1A	3704	1/1	0.90	0.10	41,41,41,41	0
55	MG	2A	3398	1/1	0.90	0.18	38,38,38,38	0
55	MG	1A	3314	1/1	0.90	0.19	29,29,29,29	0
55	MG	2a	1666	1/1	0.90	0.07	44,44,44,44	0
55	MG	1A	3158	1/1	0.90	0.21	58,58,58,58	0
55	MG	2a	1642	1/1	0.90	0.12	43,43,43,43	0
55	MG	1x	112	1/1	0.90	0.12	51,51,51,51	0
55	MG	1a	3149	1/1	0.90	0.18	56,56,56,56	0
55	MG	2A	3560	1/1	0.90	0.10	23,23,23,23	0
55	MG	2a	1748	1/1	0.90	0.11	61,61,61,61	0
55	MG	1A	3583	1/1	0.90	0.11	43,43,43,43	0
55	MG	2A	3039	1/1	0.90	0.11	32,32,32,32	0
55	MG	2a	1685	1/1	0.90	0.75	51,51,51,51	0
55	MG	1A	3590	1/1	0.90	0.14	21,21,21,21	0
55	MG	2A	3817	1/1	0.90	0.13	55,55,55,55	0
55	MG	1A	3287	1/1	0.90	0.20	48,48,48,48	0
55	MG	1A	3497	1/1	0.90	0.14	37,37,37,37	0
55	MG	1a	3049	1/1	0.90	0.12	63,63,63,63	0
55	MG	2A	3346	1/1	0.90	0.26	47,47,47,47	0
55	MG	1A	3447	1/1	0.90	0.24	33,33,33,33	0
55	MG	2A	3143	1/1	0.90	0.51	44,44,44,44	0
55	MG	2A	3531	1/1	0.90	0.10	43,43,43,43	0
55	MG	2A	3042	1/1	0.90	0.30	42,42,42,42	0
55	MG	1A	3270	1/1	0.90	0.14	41,41,41,41	0
55	MG	2A	3028	1/1	0.90	0.18	35,35,35,35	0
55	MG	1A	3057	1/1	0.90	0.16	18,18,18,18	0
55	MG	1A	3481	1/1	0.90	0.80	31,31,31,31	0
55	MG	1a	3012	1/1	0.90	0.17	43,43,43,43	0
55	MG	2A	3359	1/1	0.90	0.21	33,33,33,33	0
55	MG	1e	201	1/1	0.90	0.17	54,54,54,54	0
55	MG	2A	3697	1/1	0.90	0.06	30,30,30,30	0
55	MG	1A	3735	1/1	0.90	0.32	37,37,37,37	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2a	1727	1/1	0.90	0.15	44,44,44,44	0
55	MG	2A	3709	1/1	0.90	0.09	42,42,42,42	0
55	MG	1A	3004	1/1	0.90	0.59	25,25,25,25	0
55	MG	2A	3218	1/1	0.90	0.41	43,43,43,43	0
55	MG	1A	3193	1/1	0.90	0.11	15,15,15,15	0
55	MG	2a	1778	1/1	0.90	0.32	61,61,61,61	0
55	MG	2A	3282	1/1	0.90	0.40	49,49,49,49	0
55	MG	2a	1755	1/1	0.90	0.11	44,44,44,44	0
55	MG	2A	3689	1/1	0.90	0.19	51,51,51,51	0
55	MG	1A	3448	1/1	0.90	0.08	37,37,37,37	0
55	MG	2A	3244	1/1	0.90	0.09	35,35,35,35	0
55	MG	2A	3098	1/1	0.90	0.10	42,42,42,42	0
55	MG	2A	3111	1/1	0.90	0.11	56,56,56,56	0
55	MG	1a	3181	1/1	0.90	0.19	48,48,48,48	0
55	MG	1A	3006	1/1	0.90	0.52	25,25,25,25	0
55	MG	1A	3172	1/1	0.90	0.45	22,22,22,22	0
55	MG	2A	3778	1/1	0.90	0.20	50,50,50,50	0
55	MG	2A	3057	1/1	0.90	0.12	38,38,38,38	0
55	MG	2A	3460	1/1	0.90	0.09	32,32,32,32	0
55	MG	2a	1702	1/1	0.90	0.16	55,55,55,55	0
55	MG	2A	3768	1/1	0.90	0.22	35,35,35,35	0
55	MG	2A	3005	1/1	0.90	0.12	45,45,45,45	0
55	MG	1B	215	1/1	0.90	0.23	56,56,56,56	0
55	MG	2A	3494	1/1	0.90	0.47	50,50,50,50	0
55	MG	1A	3600	1/1	0.90	0.21	33,33,33,33	0
55	MG	2A	3701	1/1	0.90	0.23	48,48,48,48	0
55	MG	2A	3161	1/1	0.90	0.14	51,51,51,51	0
55	MG	2A	3621	1/1	0.90	0.15	21,21,21,21	0
55	MG	2a	1792	1/1	0.90	0.16	72,72,72,72	0
55	MG	2A	3235	1/1	0.90	0.29	34,34,34,34	0
55	MG	2A	3414	1/1	0.90	0.40	28,28,28,28	0
55	MG	2I	101	1/1	0.90	0.14	37,37,37,37	0
55	MG	2F	306	1/1	0.90	0.11	54,54,54,54	0
55	MG	2A	3326	1/1	0.90	0.11	34,34,34,34	0
55	MG	2A	3442	1/1	0.90	0.14	33,33,33,33	0
55	MG	1A	3933	1/1	0.90	0.17	51,51,51,51	0
55	MG	2a	1721	1/1	0.90	0.09	57,57,57,57	0
55	MG	1A	3252	1/1	0.90	0.59	35,35,35,35	0
55	MG	1A	3493	1/1	0.90	0.15	37,37,37,37	0
55	MG	2P	203	1/1	0.90	0.38	46,46,46,46	0
55	MG	1A	3691	1/1	0.90	0.14	26,26,26,26	0
55	MG	2A	3135	1/1	0.90	0.28	49,49,49,49	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3634	1/1	0.90	0.24	40,40,40,40	0
55	MG	2A	3156	1/1	0.90	0.74	32,32,32,32	0
55	MG	1V	203	1/1	0.90	0.17	22,22,22,22	0
55	MG	2A	3014	1/1	0.90	0.26	21,21,21,21	0
55	MG	2A	3112	1/1	0.90	0.19	42,42,42,42	0
55	MG	2a	1676	1/1	0.90	0.12	40,40,40,40	0
55	MG	2A	3092	1/1	0.91	0.17	44,44,44,44	0
55	MG	1A	3908	1/1	0.91	0.12	29,29,29,29	0
55	MG	1A	3479	1/1	0.91	0.17	48,48,48,48	0
55	MG	2a	1813	1/1	0.91	0.40	63,63,63,63	0
55	MG	2a	1621	1/1	0.91	0.23	40,40,40,40	0
55	MG	1A	3169	1/1	0.91	0.08	38,38,38,38	0
55	MG	2A	3045	1/1	0.91	0.17	45,45,45,45	0
55	MG	2g	201	1/1	0.91	0.11	65,65,65,65	0
55	MG	1A	4010	1/1	0.91	0.13	54,54,54,54	0
55	MG	1A	3013	1/1	0.91	0.28	20,20,20,20	0
55	MG	2A	3122	1/1	0.91	0.14	49,49,49,49	0
55	MG	1a	3081	1/1	0.91	0.14	41,41,41,41	0
55	MG	1a	3212	1/1	0.91	0.12	66,66,66,66	0
55	MG	2A	3222	1/1	0.91	0.24	47,47,47,47	0
55	MG	2A	3392	1/1	0.91	0.23	25,25,25,25	0
55	MG	1A	3606	1/1	0.91	0.09	28,28,28,28	0
55	MG	1a	3042	1/1	0.91	0.16	53,53,53,53	0
55	MG	2A	3612	1/1	0.91	0.15	26,26,26,26	0
55	MG	2A	3763	1/1	0.91	0.13	60,60,60,60	0
55	MG	2A	3740	1/1	0.91	0.11	42,42,42,42	0
55	MG	1A	3875	1/1	0.91	0.12	39,39,39,39	0
55	MG	2A	3595	1/1	0.91	0.13	45,45,45,45	0
55	MG	1A	3295	1/1	0.91	0.16	26,26,26,26	0
55	MG	2a	1647	1/1	0.91	0.10	60,60,60,60	0
55	MG	1a	3085	1/1	0.91	0.20	38,38,38,38	0
55	MG	2W	201	1/1	0.91	0.48	45,45,45,45	0
55	MG	2j	202	1/1	0.91	0.28	49,49,49,49	0
55	MG	2A	3223	1/1	0.91	0.67	51,51,51,51	0
55	MG	2A	3317	1/1	0.91	0.25	49,49,49,49	0
55	MG	1A	3324	1/1	0.91	0.16	31,31,31,31	0
55	MG	1a	3101	1/1	0.91	0.15	45,45,45,45	0
55	MG	2A	3461	1/1	0.91	0.10	51,51,51,51	0
55	MG	2A	3016	1/1	0.91	0.34	33,33,33,33	0
55	MG	2A	3252	1/1	0.91	0.30	36,36,36,36	0
55	MG	2A	3250	1/1	0.91	0.11	44,44,44,44	0
55	MG	2a	1667	1/1	0.91	0.12	40,40,40,40	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
58	V7A	1a	3213	35/35	0.91	0.31	47,61,72,74	0
55	MG	1A	4037	1/1	0.91	0.27	33,33,33,33	0
55	MG	2A	3429	1/1	0.91	0.25	32,32,32,32	0
55	MG	1A	3313	1/1	0.91	0.12	21,21,21,21	0
55	MG	1A	3697	1/1	0.91	0.18	32,32,32,32	0
55	MG	1a	3003	1/1	0.91	0.11	54,54,54,54	0
55	MG	2A	3064	1/1	0.91	0.13	33,33,33,33	0
55	MG	2a	1639	1/1	0.91	0.13	34,34,34,34	0
55	MG	1A	3707	1/1	0.91	0.25	24,24,24,24	0
55	MG	1A	3339	1/1	0.91	1.00	33,33,33,33	0
55	MG	2a	1606	1/1	0.91	0.15	30,30,30,30	0
55	MG	1A	3492	1/1	0.91	0.12	47,47,47,47	0
55	MG	1A	3330	1/1	0.91	0.47	33,33,33,33	0
55	MG	1A	3683	1/1	0.91	0.13	39,39,39,39	0
55	MG	1A	3720	1/1	0.91	0.09	56,56,56,56	0
55	MG	1A	3767	1/1	0.91	0.10	40,40,40,40	0
55	MG	2A	3643	1/1	0.91	0.16	43,43,43,43	0
55	MG	2A	3128	1/1	0.91	0.34	36,36,36,36	0
55	MG	2B	217	1/1	0.91	0.20	53,53,53,53	0
55	MG	1a	3059	1/1	0.91	0.25	41,41,41,41	0
55	MG	1A	3062	1/1	0.91	0.13	36,36,36,36	0
55	MG	1A	3502	1/1	0.91	0.10	24,24,24,24	0
55	MG	1A	3750	1/1	0.91	0.13	58,58,58,58	0
55	MG	1A	3161	1/1	0.91	0.52	29,29,29,29	0
55	MG	28	101	1/1	0.91	0.16	31,31,31,31	0
55	MG	2A	3256	1/1	0.91	0.29	55,55,55,55	0
55	MG	1a	3048	1/1	0.91	0.16	64,64,64,64	0
55	MG	2A	3280	1/1	0.91	0.12	39,39,39,39	0
55	MG	1a	3148	1/1	0.91	0.16	40,40,40,40	0
55	MG	1A	3932	1/1	0.91	0.19	15,15,15,15	0
55	MG	1A	3608	1/1	0.91	0.24	19,19,19,19	0
55	MG	1A	3588	1/1	0.91	0.24	35,35,35,35	0
55	MG	2A	3770	1/1	0.91	0.38	45,45,45,45	0
55	MG	2a	1756	1/1	0.91	0.16	33,33,33,33	0
55	MG	2A	3170	1/1	0.91	0.21	47,47,47,47	0
55	MG	1A	3063	1/1	0.91	0.15	21,21,21,21	0
55	MG	1B	234	1/1	0.91	0.20	48,48,48,48	0
55	MG	2A	3055	1/1	0.91	0.08	48,48,48,48	0
55	MG	1Y	201	1/1	0.91	0.25	33,33,33,33	0
55	MG	1A	3507	1/1	0.91	0.12	37,37,37,37	0
55	MG	1A	3610	1/1	0.91	0.34	24,24,24,24	0
55	MG	1X	102	1/1	0.91	0.67	34,34,34,34	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3411	1/1	0.91	0.19	29,29,29,29	0
55	MG	2A	3693	1/1	0.91	0.15	20,20,20,20	0
55	MG	1a	3030	1/1	0.91	0.21	50,50,50,50	0
55	MG	1A	3315	1/1	0.91	0.29	21,21,21,21	0
55	MG	1a	3067	1/1	0.91	0.12	34,34,34,34	0
55	MG	1A	3364	1/1	0.91	0.20	43,43,43,43	0
55	MG	1A	3271	1/1	0.91	0.23	32,32,32,32	0
55	MG	2A	3054	1/1	0.91	0.14	43,43,43,43	0
55	MG	1A	3069	1/1	0.91	0.21	29,29,29,29	0
55	MG	1P	206	1/1	0.91	0.55	34,34,34,34	0
55	MG	1A	3227	1/1	0.91	0.11	37,37,37,37	0
55	MG	2A	3480	1/1	0.91	0.13	38,38,38,38	0
55	MG	1A	3986	1/1	0.91	0.14	23,23,23,23	0
55	MG	2A	3848	1/1	0.91	0.13	31,31,31,31	0
55	MG	2A	3180	1/1	0.91	0.47	38,38,38,38	0
55	MG	1A	3496	1/1	0.91	0.25	26,26,26,26	0
55	MG	1A	3405	1/1	0.91	0.34	32,32,32,32	0
55	MG	1A	3401	1/1	0.91	0.15	37,37,37,37	0
55	MG	1A	3360	1/1	0.91	0.56	28,28,28,28	0
55	MG	2A	3630	1/1	0.91	0.20	32,32,32,32	0
55	MG	2A	3526	1/1	0.91	0.09	27,27,27,27	0
55	MG	1A	3322	1/1	0.91	0.15	20,20,20,20	0
55	MG	2A	3582	1/1	0.91	0.18	33,33,33,33	0
55	MG	1A	3113	1/1	0.91	0.13	36,36,36,36	0
55	MG	1D	305	1/1	0.91	0.42	20,20,20,20	0
55	MG	2A	3766	1/1	0.91	0.26	37,37,37,37	0
55	MG	1A	3320	1/1	0.91	0.52	28,28,28,28	0
55	MG	1a	3202	1/1	0.91	0.17	47,47,47,47	0
55	MG	2A	3593	1/1	0.91	0.16	50,50,50,50	0
55	MG	2A	3240	1/1	0.91	0.14	29,29,29,29	0
55	MG	1A	3463	1/1	0.91	0.32	45,45,45,45	0
55	MG	2A	3184	1/1	0.91	0.16	29,29,29,29	0
55	MG	2a	1724	1/1	0.91	0.11	49,49,49,49	0
55	MG	1U	202	1/1	0.91	0.17	28,28,28,28	0
55	MG	1A	3586	1/1	0.91	0.17	31,31,31,31	0
55	MG	2A	3178	1/1	0.91	0.14	42,42,42,42	0
55	MG	1A	3924	1/1	0.91	0.49	47,47,47,47	0
55	MG	1A	3080	1/1	0.91	0.53	41,41,41,41	0
55	MG	2A	3263	1/1	0.91	0.09	33,33,33,33	0
55	MG	1A	3427	1/1	0.91	0.10	22,22,22,22	0
55	MG	1A	3521	1/1	0.91	0.17	25,25,25,25	0
55	MG	1A	4039	1/1	0.91	0.53	31,31,31,31	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3379	1/1	0.91	0.17	43,43,43,43	0
55	MG	1A	3468	1/1	0.91	0.24	34,34,34,34	0
55	MG	2A	3629	1/1	0.91	0.51	39,39,39,39	0
55	MG	1A	3192	1/1	0.91	0.74	39,39,39,39	0
55	MG	2A	3528	1/1	0.91	0.15	24,24,24,24	0
55	MG	2A	3131	1/1	0.91	0.09	49,49,49,49	0
55	MG	1A	3852	1/1	0.91	0.10	62,62,62,62	0
55	MG	2a	1716	1/1	0.91	0.08	47,47,47,47	0
55	MG	1O	205	1/1	0.91	0.25	31,31,31,31	0
55	MG	2A	3149	1/1	0.91	0.20	43,43,43,43	0
55	MG	1A	3706	1/1	0.91	0.12	39,39,39,39	0
55	MG	2A	3550	1/1	0.91	0.12	10,10,10,10	0
55	MG	1A	3568	1/1	0.91	0.24	31,31,31,31	0
55	MG	2a	1601	1/1	0.91	0.29	23,23,23,23	0
55	MG	1A	3147	1/1	0.91	0.10	41,41,41,41	0
55	MG	1A	3529	1/1	0.91	0.25	38,38,38,38	0
55	MG	1A	3938	1/1	0.91	0.17	18,18,18,18	0
55	MG	1A	3066	1/1	0.91	0.23	23,23,23,23	0
55	MG	1A	4024	1/1	0.91	0.16	35,35,35,35	0
55	MG	1t	201	1/1	0.91	0.21	42,42,42,42	0
55	MG	2A	3351	1/1	0.91	0.10	38,38,38,38	0
55	MG	1A	3589	1/1	0.91	0.13	25,25,25,25	0
55	MG	2a	1805	1/1	0.91	0.30	50,50,50,50	0
55	MG	1A	3736	1/1	0.91	0.10	33,33,33,33	0
55	MG	1E	306	1/1	0.91	0.18	33,33,33,33	0
55	MG	2A	3714	1/1	0.91	0.11	56,56,56,56	0
55	MG	1a	3105	1/1	0.91	0.15	35,35,35,35	0
55	MG	2l	201	1/1	0.91	0.76	47,47,47,47	0
55	MG	1A	3994	1/1	0.91	0.13	10,10,10,10	0
55	MG	2A	3362	1/1	0.91	0.11	42,42,42,42	0
55	MG	1A	3678	1/1	0.91	0.13	16,16,16,16	0
55	MG	1a	3144	1/1	0.91	0.12	56,56,56,56	0
55	MG	1A	3027	1/1	0.92	0.40	28,28,28,28	0
55	MG	1A	3055	1/1	0.92	0.15	29,29,29,29	0
55	MG	2A	3258	1/1	0.92	0.14	23,23,23,23	0
55	MG	2a	1787	1/1	0.92	0.13	49,49,49,49	0
55	MG	2A	3205	1/1	0.92	0.14	54,54,54,54	0
55	MG	2A	3645	1/1	0.92	0.20	49,49,49,49	0
55	MG	1E	311	1/1	0.92	0.24	48,48,48,48	0
55	MG	1A	3998	1/1	0.92	0.10	44,44,44,44	0
55	MG	1a	3162	1/1	0.92	0.35	47,47,47,47	0
55	MG	1a	3200	1/1	0.92	0.13	53,53,53,53	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3733	1/1	0.92	0.20	15,15,15,15	0
55	MG	2A	3174	1/1	0.92	0.31	33,33,33,33	0
55	MG	1a	3121	1/1	0.92	0.12	54,54,54,54	0
55	MG	1A	3240	1/1	0.92	0.20	31,31,31,31	0
55	MG	1A	3787	1/1	0.92	0.10	58,58,58,58	0
55	MG	1A	3734	1/1	0.92	0.20	12,12,12,12	0
55	MG	2A	3076	1/1	0.92	0.24	30,30,30,30	0
55	MG	2A	3720	1/1	0.92	0.18	44,44,44,44	0
55	MG	2A	3134	1/1	0.92	0.15	40,40,40,40	0
55	MG	1A	3280	1/1	0.92	0.20	19,19,19,19	0
55	MG	1A	3644	1/1	0.92	0.18	28,28,28,28	0
55	MG	2A	3651	1/1	0.92	0.14	48,48,48,48	0
55	MG	1A	3009	1/1	0.92	0.10	32,32,32,32	0
55	MG	1A	3215	1/1	0.92	0.11	44,44,44,44	0
55	MG	2A	3841	1/1	0.92	0.48	53,53,53,53	0
55	MG	1A	3751	1/1	0.92	0.15	44,44,44,44	0
55	MG	1a	3170	1/1	0.92	0.11	36,36,36,36	0
55	MG	2a	1706	1/1	0.92	0.17	30,30,30,30	0
55	MG	2A	3059	1/1	0.92	0.18	32,32,32,32	0
55	MG	2A	3444	1/1	0.92	1.16	36,36,36,36	0
55	MG	1A	3284	1/1	0.92	0.13	14,14,14,14	0
55	MG	2a	1775	1/1	0.92	0.12	68,68,68,68	0
55	MG	2A	3191	1/1	0.92	0.11	50,50,50,50	0
55	MG	1a	3007	1/1	0.92	0.18	27,27,27,27	0
55	MG	1A	3715	1/1	0.92	0.09	21,21,21,21	0
55	MG	1A	3877	1/1	0.92	0.12	37,37,37,37	0
55	MG	1A	3180	1/1	0.92	0.30	56,56,56,56	0
55	MG	2A	3514	1/1	0.92	0.10	51,51,51,51	0
55	MG	2A	3716	1/1	0.92	0.09	45,45,45,45	0
55	MG	2a	1780	1/1	0.92	0.47	62,62,62,62	0
55	MG	1A	3336	1/1	0.92	0.57	32,32,32,32	0
55	MG	1A	3472	1/1	0.92	0.14	30,30,30,30	0
55	MG	1A	3915	1/1	0.92	0.22	53,53,53,53	0
55	MG	2A	3416	1/1	0.92	0.12	33,33,33,33	0
55	MG	2B	220	1/1	0.92	0.20	50,50,50,50	0
55	MG	1A	3412	1/1	0.92	0.12	27,27,27,27	0
55	MG	2A	3832	1/1	0.92	0.18	36,36,36,36	0
55	MG	2a	1712	1/1	0.92	0.15	41,41,41,41	0
55	MG	1a	3090	1/1	0.92	0.23	46,46,46,46	0
55	MG	1A	3159	1/1	0.92	0.13	28,28,28,28	0
55	MG	1G	201	1/1	0.92	0.13	30,30,30,30	0
55	MG	1B	214	1/1	0.92	0.26	30,30,30,30	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3041	1/1	0.92	0.14	29,29,29,29	0
55	MG	1a	3072	1/1	0.92	0.17	60,60,60,60	0
55	MG	2a	1798	1/1	0.92	0.15	53,53,53,53	0
55	MG	1A	3288	1/1	0.92	0.31	27,27,27,27	0
55	MG	2A	3094	1/1	0.92	0.31	49,49,49,49	0
55	MG	2A	3794	1/1	0.92	0.11	37,37,37,37	0
55	MG	1A	3333	1/1	0.92	0.41	30,30,30,30	0
55	MG	2A	3194	1/1	0.92	0.29	26,26,26,26	0
55	MG	2A	3241	1/1	0.92	0.68	32,32,32,32	0
55	MG	1A	3311	1/1	0.92	0.55	41,41,41,41	0
55	MG	1A	3400	1/1	0.92	0.15	27,27,27,27	0
55	MG	2a	1814	1/1	0.92	0.24	46,46,46,46	0
55	MG	1A	3744	1/1	0.92	0.82	21,21,21,21	0
55	MG	1A	4001	1/1	0.92	0.12	41,41,41,41	0
55	MG	1A	3868	1/1	0.92	0.11	29,29,29,29	0
55	MG	2A	3731	1/1	0.92	0.28	46,46,46,46	0
55	MG	1a	3079	1/1	0.92	0.16	41,41,41,41	0
55	MG	1A	3897	1/1	0.92	0.24	42,42,42,42	0
55	MG	2A	3762	1/1	0.92	0.46	67,67,67,67	0
55	MG	2A	3589	1/1	0.92	0.14	42,42,42,42	0
55	MG	2a	1743	1/1	0.92	0.14	38,38,38,38	0
55	MG	2A	3614	1/1	0.92	0.20	13,13,13,13	0
55	MG	2A	3495	1/1	0.92	0.11	17,17,17,17	0
55	MG	2A	3616	1/1	0.92	0.21	44,44,44,44	0
55	MG	2A	3310	1/1	0.92	0.25	28,28,28,28	0
55	MG	2A	3536	1/1	0.92	0.17	15,15,15,15	0
55	MG	2A	3108	1/1	0.92	0.31	41,41,41,41	0
55	MG	2A	3095	1/1	0.92	0.12	37,37,37,37	0
55	MG	2a	1680	1/1	0.92	0.20	40,40,40,40	0
55	MG	1A	3904	1/1	0.92	0.12	32,32,32,32	0
55	MG	1A	3307	1/1	0.92	0.17	39,39,39,39	0
55	MG	1a	3107	1/1	0.92	0.25	36,36,36,36	0
55	MG	2A	3248	1/1	0.92	0.12	33,33,33,33	0
55	MG	1A	3435	1/1	0.92	0.10	54,54,54,54	0
55	MG	2A	3824	1/1	0.92	0.48	53,53,53,53	0
55	MG	2A	3655	1/1	0.92	0.43	26,26,26,26	0
55	MG	1A	3595	1/1	0.92	0.18	24,24,24,24	0
55	MG	2A	3511	1/1	0.92	0.10	27,27,27,27	0
55	MG	1a	3017	1/1	0.92	0.13	48,48,48,48	0
55	MG	2B	203	1/1	0.92	0.14	52,52,52,52	0
55	MG	2a	1718	1/1	0.92	0.18	38,38,38,38	0
55	MG	1G	204	1/1	0.92	0.14	51,51,51,51	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3205	1/1	0.92	0.16	18,18,18,18	0
55	MG	1A	3597	1/1	0.92	0.10	34,34,34,34	0
55	MG	2a	1646	1/1	0.92	0.07	58,58,58,58	0
55	MG	2A	3181	1/1	0.92	0.09	10,10,10,10	0
55	MG	2A	3752	1/1	0.92	0.07	30,30,30,30	0
55	MG	1B	228	1/1	0.92	0.11	33,33,33,33	0
55	MG	2A	3736	1/1	0.92	0.17	54,54,54,54	0
55	MG	1a	3005	1/1	0.92	0.23	51,51,51,51	0
55	MG	1A	3893	1/1	0.92	0.13	35,35,35,35	0
55	MG	1A	3622	1/1	0.92	0.24	39,39,39,39	0
55	MG	1A	3208	1/1	0.92	0.57	28,28,28,28	0
55	MG	2a	1759	1/1	0.92	0.19	50,50,50,50	0
55	MG	1U	208	1/1	0.92	0.38	36,36,36,36	0
55	MG	2A	3440	1/1	0.92	0.35	40,40,40,40	0
55	MG	2a	1767	1/1	0.92	0.20	52,52,52,52	0
55	MG	2A	3209	1/1	0.92	0.17	24,24,24,24	0
55	MG	2A	3302	1/1	0.92	0.40	40,40,40,40	0
55	MG	2A	3183	1/1	0.92	0.30	39,39,39,39	0
55	MG	1A	3684	1/1	0.92	0.13	52,52,52,52	0
55	MG	1A	4040	1/1	0.92	0.19	33,33,33,33	0
55	MG	2A	3213	1/1	0.92	0.40	32,32,32,32	0
55	MG	1A	3807	1/1	0.92	0.34	31,31,31,31	0
55	MG	2A	3193	1/1	0.92	0.29	52,52,52,52	0
55	MG	2a	1643	1/1	0.92	0.18	49,49,49,49	0
55	MG	2a	1671	1/1	0.92	0.12	42,42,42,42	0
55	MG	2a	1688	1/1	0.92	0.39	33,33,33,33	0
55	MG	1f	202	1/1	0.92	0.19	38,38,38,38	0
55	MG	2a	1741	1/1	0.92	0.16	59,59,59,59	0
55	MG	2A	3296	1/1	0.92	0.28	35,35,35,35	0
55	MG	2A	3785	1/1	0.92	0.21	43,43,43,43	0
55	MG	2A	3006	1/1	0.92	0.17	38,38,38,38	0
55	MG	2a	1768	1/1	0.92	0.19	36,36,36,36	0
55	MG	1a	3112	1/1	0.92	0.11	54,54,54,54	0
55	MG	2A	3530	1/1	0.92	0.19	45,45,45,45	0
55	MG	2k	201	1/1	0.92	0.11	54,54,54,54	0
55	MG	1a	3174	1/1	0.92	0.10	49,49,49,49	0
55	MG	2A	3457	1/1	0.92	0.10	54,54,54,54	0
55	MG	2T	202	1/1	0.92	0.14	39,39,39,39	0
55	MG	2B	213	1/1	0.92	0.26	50,50,50,50	0
55	MG	2a	1757	1/1	0.92	0.23	44,44,44,44	0
55	MG	1A	3437	1/1	0.92	0.21	29,29,29,29	0
55	MG	2A	3086	1/1	0.92	0.17	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3142	1/1	0.92	0.24	37,37,37,37	0
55	MG	1a	3169	1/1	0.92	0.10	55,55,55,55	0
55	MG	2A	3839	1/1	0.92	0.14	30,30,30,30	0
55	MG	2a	1744	1/1	0.92	0.58	72,72,72,72	0
55	MG	1A	3366	1/1	0.92	0.17	43,43,43,43	0
55	MG	2A	3275	1/1	0.92	0.13	45,45,45,45	0
55	MG	1V	202	1/1	0.92	0.78	35,35,35,35	0
55	MG	1A	3419	1/1	0.92	0.28	49,49,49,49	0
55	MG	2a	1654	1/1	0.92	0.09	35,35,35,35	0
55	MG	1A	3204	1/1	0.92	1.09	25,25,25,25	0
55	MG	2A	3067	1/1	0.92	0.16	27,27,27,27	0
55	MG	1A	3430	1/1	0.92	0.10	32,32,32,32	0
55	MG	27	102	1/1	0.92	0.51	36,36,36,36	0
55	MG	1A	3835	1/1	0.92	0.74	24,24,24,24	0
55	MG	1A	3012	1/1	0.92	0.74	27,27,27,27	0
55	MG	1A	3218	1/1	0.93	0.24	22,22,22,22	0
55	MG	1A	3273	1/1	0.93	0.10	26,26,26,26	0
55	MG	1P	204	1/1	0.93	1.23	19,19,19,19	0
55	MG	1A	3508	1/1	0.93	0.12	33,33,33,33	0
55	MG	1A	3780	1/1	0.93	0.20	41,41,41,41	0
55	MG	1A	3274	1/1	0.93	0.13	40,40,40,40	0
55	MG	1A	3570	1/1	0.93	0.38	43,43,43,43	0
55	MG	2A	3320	1/1	0.93	0.14	30,30,30,30	0
55	MG	2a	1613	1/1	0.93	0.10	45,45,45,45	0
55	MG	1A	3146	1/1	0.93	0.31	49,49,49,49	0
55	MG	2a	1810	1/1	0.93	0.08	66,66,66,66	0
55	MG	2a	1681	1/1	0.93	0.79	54,54,54,54	0
55	MG	1A	3911	1/1	0.93	0.25	47,47,47,47	0
55	MG	2A	3436	1/1	0.93	0.64	33,33,33,33	0
55	MG	2A	3449	1/1	0.93	0.15	32,32,32,32	0
55	MG	2a	1770	1/1	0.93	0.12	40,40,40,40	0
55	MG	1A	3001	1/1	0.93	0.28	26,26,26,26	0
55	MG	2A	3331	1/1	0.93	0.21	45,45,45,45	0
55	MG	1A	3532	1/1	0.93	0.18	40,40,40,40	0
55	MG	2A	3443	1/1	0.93	0.67	46,46,46,46	0
55	MG	2A	3754	1/1	0.93	0.25	43,43,43,43	0
55	MG	1A	3719	1/1	0.93	0.15	19,19,19,19	0
55	MG	1A	3808	1/1	0.93	0.47	22,22,22,22	0
55	MG	1A	3687	1/1	0.93	0.13	59,59,59,59	0
55	MG	2A	3725	1/1	0.93	0.12	25,25,25,25	0
55	MG	2A	3268	1/1	0.93	0.11	32,32,32,32	0
55	MG	2A	3096	1/1	0.93	0.15	44,44,44,44	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3793	1/1	0.93	0.14	13,13,13,13	0
55	MG	1a	3034	1/1	0.93	0.30	49,49,49,49	0
55	MG	2A	3617	1/1	0.93	0.14	23,23,23,23	0
55	MG	2A	3103	1/1	0.93	0.91	34,34,34,34	0
55	MG	2A	3025	1/1	0.93	0.29	35,35,35,35	0
55	MG	1A	3907	1/1	0.93	0.54	61,61,61,61	0
55	MG	2A	3355	1/1	0.93	0.23	30,30,30,30	0
55	MG	1A	3073	1/1	0.93	0.27	22,22,22,22	0
55	MG	1X	101	1/1	0.93	0.25	30,30,30,30	0
55	MG	1A	3747	1/1	0.93	0.13	38,38,38,38	0
55	MG	2O	201	1/1	0.93	0.10	31,31,31,31	0
55	MG	2A	3674	1/1	0.93	0.23	36,36,36,36	0
55	MG	1A	3286	1/1	0.93	0.13	41,41,41,41	0
55	MG	1A	3191	1/1	0.93	0.29	44,44,44,44	0
55	MG	1A	3342	1/1	0.93	0.29	46,46,46,46	0
55	MG	2A	3781	1/1	0.93	0.14	43,43,43,43	0
55	MG	2A	3378	1/1	0.93	0.20	44,44,44,44	0
55	MG	2A	3695	1/1	0.93	0.12	44,44,44,44	0
55	MG	2A	3578	1/1	0.93	0.11	22,22,22,22	0
55	MG	2a	1612	1/1	0.93	0.69	39,39,39,39	0
55	MG	1A	3367	1/1	0.93	0.07	27,27,27,27	0
55	MG	2A	3580	1/1	0.93	0.08	40,40,40,40	0
55	MG	1A	3855	1/1	0.93	0.11	36,36,36,36	0
55	MG	2A	3009	1/1	0.93	0.10	27,27,27,27	0
55	MG	1a	3179	1/1	0.93	0.13	47,47,47,47	0
55	MG	28	103	1/1	0.93	0.13	53,53,53,53	0
55	MG	2D	305	1/1	0.93	0.52	41,41,41,41	0
55	MG	2A	3678	1/1	0.93	0.10	43,43,43,43	0
55	MG	2A	3085	1/1	0.93	0.35	27,27,27,27	0
55	MG	1a	3013	1/1	0.93	0.14	56,56,56,56	0
55	MG	1A	3417	1/1	0.93	0.18	30,30,30,30	0
55	MG	1A	3658	1/1	0.93	0.15	32,32,32,32	0
55	MG	1A	3988	1/1	0.93	0.08	36,36,36,36	0
55	MG	2A	3423	1/1	0.93	0.17	31,31,31,31	0
55	MG	2a	1753	1/1	0.93	0.10	33,33,33,33	0
55	MG	1a	3152	1/1	0.93	0.24	40,40,40,40	0
55	MG	2A	3023	1/1	0.93	0.83	41,41,41,41	0
55	MG	2A	3290	1/1	0.93	0.35	30,30,30,30	0
55	MG	2A	3696	1/1	0.93	0.08	50,50,50,50	0
55	MG	1a	3033	1/1	0.93	0.13	40,40,40,40	0
55	MG	2x	103	1/1	0.93	0.17	29,29,29,29	0
55	MG	2G	201	1/1	0.93	0.15	42,42,42,42	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3120	1/1	0.93	0.24	30,30,30,30	0
55	MG	1a	3069	1/1	0.93	0.12	23,23,23,23	0
55	MG	1A	3224	1/1	0.93	0.16	48,48,48,48	0
55	MG	2a	1694	1/1	0.93	0.47	46,46,46,46	0
55	MG	1A	3428	1/1	0.93	0.23	21,21,21,21	0
55	MG	1A	3377	1/1	0.93	0.10	38,38,38,38	0
55	MG	1A	3343	1/1	0.93	0.18	25,25,25,25	0
55	MG	1A	3928	1/1	0.93	0.07	33,33,33,33	0
55	MG	1a	3068	1/1	0.93	0.12	30,30,30,30	0
55	MG	2A	3353	1/1	0.93	0.21	36,36,36,36	0
55	MG	2A	3557	1/1	0.93	0.12	15,15,15,15	0
55	MG	2A	3500	1/1	0.93	0.17	33,33,33,33	0
55	MG	2A	3203	1/1	0.93	0.24	44,44,44,44	0
55	MG	1A	3033	1/1	0.93	0.14	35,35,35,35	0
55	MG	1A	3825	1/1	0.93	0.37	52,52,52,52	0
55	MG	1a	3083	1/1	0.93	0.09	40,40,40,40	0
55	MG	2a	1736	1/1	0.93	0.10	54,54,54,54	0
55	MG	1A	4035	1/1	0.93	0.26	15,15,15,15	0
55	MG	2A	3591	1/1	0.93	0.18	41,41,41,41	0
55	MG	2A	3040	1/1	0.93	0.12	28,28,28,28	0
55	MG	2A	3328	1/1	0.93	0.56	64,64,64,64	0
55	MG	1A	3413	1/1	0.93	0.39	21,21,21,21	0
55	MG	2A	3635	1/1	0.93	0.25	31,31,31,31	0
55	MG	1A	3232	1/1	0.93	0.10	26,26,26,26	0
55	MG	1S	203	1/1	0.93	0.09	44,44,44,44	0
55	MG	1A	3764	1/1	0.93	0.12	41,41,41,41	0
55	MG	1A	3098	1/1	0.93	0.56	32,32,32,32	0
55	MG	1a	3113	1/1	0.93	0.11	41,41,41,41	0
55	MG	2A	3231	1/1	0.93	0.14	55,55,55,55	0
55	MG	2A	3661	1/1	0.93	0.17	41,41,41,41	0
55	MG	1A	3585	1/1	0.93	0.11	22,22,22,22	0
55	MG	1A	3827	1/1	0.93	0.38	22,22,22,22	0
55	MG	1A	3774	1/1	0.93	0.24	37,37,37,37	0
55	MG	2a	1652	1/1	0.93	0.21	40,40,40,40	0
55	MG	1A	3718	1/1	0.93	0.24	46,46,46,46	0
55	MG	2A	3283	1/1	0.93	0.11	31,31,31,31	0
55	MG	2a	1669	1/1	0.93	0.08	40,40,40,40	0
55	MG	1A	3140	1/1	0.93	0.38	33,33,33,33	0
55	MG	1a	3168	1/1	0.93	0.16	50,50,50,50	0
55	MG	1N	203	1/1	0.93	0.31	45,45,45,45	0
55	MG	2A	3571	1/1	0.93	0.17	29,29,29,29	0
55	MG	1f	201	1/1	0.93	0.19	33,33,33,33	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3510	1/1	0.93	0.15	21,21,21,21	0
55	MG	2A	3441	1/1	0.93	0.12	31,31,31,31	0
55	MG	1A	3716	1/1	0.93	0.11	20,20,20,20	0
55	MG	2a	1615	1/1	0.93	0.17	37,37,37,37	0
55	MG	2A	3179	1/1	0.93	0.27	30,30,30,30	0
55	MG	1A	4023	1/1	0.93	0.15	40,40,40,40	0
55	MG	2A	3254	1/1	0.93	0.33	41,41,41,41	0
55	MG	1A	3331	1/1	0.93	0.13	21,21,21,21	0
55	MG	1A	3155	1/1	0.93	0.19	23,23,23,23	0
55	MG	1a	3154	1/1	0.93	0.08	41,41,41,41	0
55	MG	2a	1636	1/1	0.93	0.12	49,49,49,49	0
55	MG	2A	3053	1/1	0.93	0.17	34,34,34,34	0
55	MG	1T	202	1/1	0.93	0.22	46,46,46,46	0
55	MG	1A	3765	1/1	0.93	0.24	26,26,26,26	0
55	MG	1A	3790	1/1	0.93	0.21	28,28,28,28	0
55	MG	1A	3862	1/1	0.93	0.10	39,39,39,39	0
55	MG	1A	3803	1/1	0.93	0.44	29,29,29,29	0
55	MG	2E	304	1/1	0.93	0.20	30,30,30,30	0
55	MG	1A	3438	1/1	0.93	0.23	33,33,33,33	0
55	MG	2a	1728	1/1	0.93	0.21	41,41,41,41	0
55	MG	2A	3284	1/1	0.93	0.11	33,33,33,33	0
55	MG	2A	3391	1/1	0.93	0.11	30,30,30,30	0
55	MG	2Z	301	1/1	0.93	0.19	68,68,68,68	0
55	MG	1A	3457	1/1	0.93	0.18	38,38,38,38	0
55	MG	15	102	1/1	0.93	0.64	33,33,33,33	0
55	MG	1A	3201	1/1	0.93	0.20	36,36,36,36	0
55	MG	2a	1679	1/1	0.93	0.09	45,45,45,45	0
55	MG	2A	3681	1/1	0.93	0.21	50,50,50,50	0
55	MG	1B	213	1/1	0.93	0.11	48,48,48,48	0
55	MG	2A	3239	1/1	0.93	0.66	34,34,34,34	0
55	MG	2A	3166	1/1	0.93	0.23	38,38,38,38	0
55	MG	1A	3480	1/1	0.93	0.13	20,20,20,20	0
55	MG	1a	3108	1/1	0.93	0.13	63,63,63,63	0
55	MG	2a	1719	1/1	0.93	0.14	43,43,43,43	0
55	MG	2A	3324	1/1	0.93	0.11	26,26,26,26	0
55	MG	1l	202	1/1	0.93	0.08	18,18,18,18	0
55	MG	1A	3230	1/1	0.93	0.21	23,23,23,23	0
55	MG	1A	3564	1/1	0.93	0.18	15,15,15,15	0
55	MG	1a	3145	1/1	0.93	0.14	23,23,23,23	0
55	MG	1A	3007	1/1	0.93	0.12	27,27,27,27	0
55	MG	2Q	201	1/1	0.93	0.10	53,53,53,53	0
55	MG	1a	3117	1/1	0.93	0.16	46,46,46,46	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2I	105	1/1	0.93	0.14	49,49,49,49	0
55	MG	1A	3282	1/1	0.93	0.27	24,24,24,24	0
55	MG	1a	3209	1/1	0.93	0.09	38,38,38,38	0
55	MG	1A	3640	1/1	0.93	0.53	47,47,47,47	0
55	MG	1A	3556	1/1	0.93	0.13	26,26,26,26	0
55	MG	2a	1660	1/1	0.93	0.09	46,46,46,46	0
55	MG	1A	3316	1/1	0.93	0.37	14,14,14,14	0
55	MG	2a	1788	1/1	0.93	0.07	54,54,54,54	0
55	MG	2a	1657	1/1	0.93	0.26	45,45,45,45	0
55	MG	2a	1634	1/1	0.93	0.14	51,51,51,51	0
55	MG	1A	3831	1/1	0.93	0.12	26,26,26,26	0
55	MG	2a	1690	1/1	0.93	0.18	39,39,39,39	0
55	MG	2x	104	1/1	0.93	0.22	50,50,50,50	0
55	MG	1A	3630	1/1	0.93	0.13	18,18,18,18	0
55	MG	1A	3509	1/1	0.93	0.17	24,24,24,24	0
55	MG	1S	201	1/1	0.93	0.21	47,47,47,47	0
55	MG	1A	3918	1/1	0.93	0.13	19,19,19,19	0
55	MG	1A	3733	1/1	0.93	0.15	17,17,17,17	0
55	MG	2A	3037	1/1	0.93	0.17	17,17,17,17	0
55	MG	1A	3415	1/1	0.93	0.52	26,26,26,26	0
55	MG	2a	1797	1/1	0.93	0.15	47,47,47,47	0
55	MG	1A	3011	1/1	0.93	0.37	16,16,16,16	0
55	MG	2A	3684	1/1	0.93	0.09	36,36,36,36	0
55	MG	2A	3175	1/1	0.93	0.75	37,37,37,37	0
55	MG	1B	223	1/1	0.93	0.12	28,28,28,28	0
55	MG	2B	206	1/1	0.93	0.14	40,40,40,40	0
55	MG	1A	3981	1/1	0.93	0.15	47,47,47,47	0
55	MG	1F	305	1/1	0.93	0.10	34,34,34,34	0
55	MG	2a	1735	1/1	0.93	0.07	33,33,33,33	0
55	MG	2A	3679	1/1	0.93	0.08	40,40,40,40	0
55	MG	2A	3506	1/1	0.93	0.17	33,33,33,33	0
55	MG	1A	3885	1/1	0.93	0.23	46,46,46,46	0
55	MG	2A	3004	1/1	0.93	0.10	24,24,24,24	0
55	MG	2A	3371	1/1	0.93	0.35	27,27,27,27	0
55	MG	2B	204	1/1	0.93	0.21	47,47,47,47	0
55	MG	2A	3756	1/1	0.93	0.24	25,25,25,25	0
55	MG	2a	1742	1/1	0.93	0.19	44,44,44,44	0
55	MG	2A	3260	1/1	0.93	0.29	23,23,23,23	0
55	MG	1A	3264	1/1	0.93	0.19	42,42,42,42	0
55	MG	1A	3805	1/1	0.93	0.11	41,41,41,41	0
55	MG	1A	3216	1/1	0.93	0.11	30,30,30,30	0
55	MG	1A	3202	1/1	0.93	0.35	33,33,33,33	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3173	1/1	0.93	0.10	31,31,31,31	0
55	MG	1a	3124	1/1	0.93	0.09	34,34,34,34	0
55	MG	1A	3223	1/1	0.93	0.16	31,31,31,31	0
55	MG	1A	3247	1/1	0.94	0.20	37,37,37,37	0
55	MG	2A	3125	1/1	0.94	0.17	18,18,18,18	0
55	MG	1A	3812	1/1	0.94	0.11	39,39,39,39	0
55	MG	2A	3438	1/1	0.94	0.91	30,30,30,30	0
55	MG	2A	3596	1/1	0.94	0.17	26,26,26,26	0
55	MG	1A	3464	1/1	0.94	0.59	29,29,29,29	0
55	MG	1A	3338	1/1	0.94	0.12	55,55,55,55	0
55	MG	2A	3447	1/1	0.94	0.65	39,39,39,39	0
55	MG	1A	3886	1/1	0.94	0.22	36,36,36,36	0
55	MG	1A	3108	1/1	0.94	0.11	33,33,33,33	0
55	MG	1E	303	1/1	0.94	0.49	22,22,22,22	0
55	MG	2A	3605	1/1	0.94	0.12	33,33,33,33	0
55	MG	1A	3913	1/1	0.94	0.27	50,50,50,50	0
55	MG	2A	3012	1/1	0.94	0.25	40,40,40,40	0
55	MG	1A	3839	1/1	0.94	0.11	36,36,36,36	0
55	MG	1A	3196	1/1	0.94	0.12	24,24,24,24	0
55	MG	2B	201	1/1	0.94	0.14	40,40,40,40	0
55	MG	2A	3421	1/1	0.94	0.21	22,22,22,22	0
55	MG	2A	3542	1/1	0.94	0.09	29,29,29,29	0
55	MG	1A	3577	1/1	0.94	0.12	15,15,15,15	0
55	MG	1A	3096	1/1	0.94	0.13	23,23,23,23	0
55	MG	1A	3863	1/1	0.94	0.10	49,49,49,49	0
55	MG	2A	3773	1/1	0.94	0.35	36,36,36,36	0
55	MG	1A	3261	1/1	0.94	0.18	23,23,23,23	0
55	MG	1a	3020	1/1	0.94	0.13	34,34,34,34	0
55	MG	1A	3372	1/1	0.94	0.47	22,22,22,22	0
55	MG	1A	3351	1/1	0.94	0.26	28,28,28,28	0
55	MG	1A	3817	1/1	0.94	0.19	40,40,40,40	0
55	MG	1A	3983	1/1	0.94	0.17	25,25,25,25	0
55	MG	1A	3229	1/1	0.94	0.62	29,29,29,29	0
55	MG	2a	1777	1/1	0.94	0.17	38,38,38,38	0
55	MG	2E	301	1/1	0.94	0.12	31,31,31,31	0
55	MG	1Q	201	1/1	0.94	0.17	26,26,26,26	0
55	MG	1a	3026	1/1	0.94	0.11	27,27,27,27	0
55	MG	1A	3748	1/1	0.94	0.12	27,27,27,27	0
55	MG	1W	206	1/1	0.94	0.53	27,27,27,27	0
55	MG	2B	214	1/1	0.94	0.17	49,49,49,49	0
55	MG	2A	3809	1/1	0.94	0.08	40,40,40,40	0
55	MG	2A	3298	1/1	0.94	0.33	52,52,52,52	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3301	1/1	0.94	0.95	29,29,29,29	0
55	MG	1A	3165	1/1	0.94	0.74	40,40,40,40	0
55	MG	2A	3496	1/1	0.94	0.19	27,27,27,27	0
55	MG	2A	3036	1/1	0.94	0.18	23,23,23,23	0
55	MG	1A	3771	1/1	0.94	0.11	44,44,44,44	0
55	MG	1A	4025	1/1	0.94	0.15	45,45,45,45	0
55	MG	1a	3114	1/1	0.94	0.07	51,51,51,51	0
55	MG	2a	1720	1/1	0.94	0.11	23,23,23,23	0
55	MG	1a	3118	1/1	0.94	0.15	42,42,42,42	0
55	MG	2P	204	1/1	0.94	0.26	44,44,44,44	0
55	MG	2A	3245	1/1	0.94	0.21	24,24,24,24	0
55	MG	1A	3699	1/1	0.94	0.10	22,22,22,22	0
55	MG	1A	3160	1/1	0.94	0.32	45,45,45,45	0
55	MG	2A	3501	1/1	0.94	0.18	26,26,26,26	0
55	MG	1A	3141	1/1	0.94	0.19	37,37,37,37	0
55	MG	10	107	1/1	0.94	0.11	24,24,24,24	0
55	MG	2A	3603	1/1	0.94	0.09	45,45,45,45	0
55	MG	1N	204	1/1	0.94	0.14	33,33,33,33	0
55	MG	2A	3365	1/1	0.94	0.57	34,34,34,34	0
55	MG	2a	1771	1/1	0.94	0.14	39,39,39,39	0
55	MG	1a	3139	1/1	0.94	0.06	30,30,30,30	0
55	MG	1A	3446	1/1	0.94	0.20	42,42,42,42	0
55	MG	2A	3338	1/1	0.94	0.10	32,32,32,32	0
55	MG	1a	3024	1/1	0.94	0.11	30,30,30,30	0
55	MG	2A	3823	1/1	0.94	0.10	26,26,26,26	0
55	MG	1A	3451	1/1	0.94	0.31	47,47,47,47	0
55	MG	2A	3575	1/1	0.94	0.12	26,26,26,26	0
55	MG	2A	3519	1/1	0.94	0.14	60,60,60,60	0
55	MG	25	101	1/1	0.94	0.13	38,38,38,38	0
55	MG	2A	3034	1/1	0.94	0.44	32,32,32,32	0
55	MG	2a	1623	1/1	0.94	0.43	48,48,48,48	0
55	MG	2A	3847	1/1	0.94	0.14	47,47,47,47	0
55	MG	1A	3514	1/1	0.94	0.96	33,33,33,33	0
55	MG	2A	3776	1/1	0.94	0.32	57,57,57,57	0
55	MG	1A	3060	1/1	0.94	0.14	21,21,21,21	0
55	MG	2d	302	1/1	0.94	0.11	47,47,47,47	0
55	MG	2A	3382	1/1	0.94	0.21	22,22,22,22	0
55	MG	1A	4038	1/1	0.94	0.23	35,35,35,35	0
55	MG	1a	3206	1/1	0.94	0.11	60,60,60,60	0
55	MG	1A	3739	1/1	0.94	0.17	34,34,34,34	0
55	MG	1a	3199	1/1	0.94	0.06	55,55,55,55	0
55	MG	2A	3534	1/1	0.94	0.16	21,21,21,21	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3777	1/1	0.94	0.25	46,46,46,46	0
55	MG	1A	3398	1/1	0.94	1.12	29,29,29,29	0
55	MG	1A	3460	1/1	0.94	0.18	41,41,41,41	0
55	MG	1A	3511	1/1	0.94	0.14	21,21,21,21	0
55	MG	2A	3363	1/1	0.94	0.15	49,49,49,49	0
55	MG	2A	3521	1/1	0.94	0.08	33,33,33,33	0
55	MG	1A	3666	1/1	0.94	0.10	23,23,23,23	0
55	MG	1A	3976	1/1	0.94	0.22	26,26,26,26	0
55	MG	2A	3761	1/1	0.94	0.12	55,55,55,55	0
55	MG	2A	3499	1/1	0.94	0.24	45,45,45,45	0
55	MG	1A	3154	1/1	0.94	0.29	17,17,17,17	0
55	MG	2A	3273	1/1	0.94	0.14	24,24,24,24	0
55	MG	2A	3079	1/1	0.94	0.25	50,50,50,50	0
55	MG	1A	3634	1/1	0.94	0.20	32,32,32,32	0
55	MG	1A	3222	1/1	0.94	0.16	19,19,19,19	0
55	MG	2a	1678	1/1	0.94	0.14	40,40,40,40	0
55	MG	1A	3921	1/1	0.94	0.16	54,54,54,54	0
55	MG	1A	3305	1/1	0.94	0.15	23,23,23,23	0
55	MG	2A	3815	1/1	0.94	0.15	44,44,44,44	0
55	MG	1A	3624	1/1	0.94	0.15	43,43,43,43	0
55	MG	2A	3802	1/1	0.94	0.12	17,17,17,17	0
55	MG	2a	1772	1/1	0.94	0.27	28,28,28,28	0
55	MG	2E	303	1/1	0.94	0.29	28,28,28,28	0
55	MG	2A	3653	1/1	0.94	0.10	41,41,41,41	0
55	MG	1A	3383	1/1	0.94	0.56	27,27,27,27	0
55	MG	1A	3051	1/1	0.94	0.11	16,16,16,16	0
55	MG	1A	3652	1/1	0.94	0.26	29,29,29,29	0
55	MG	1x	107	1/1	0.94	0.13	48,48,48,48	0
55	MG	2A	3114	1/1	0.94	0.35	30,30,30,30	0
55	MG	2A	3352	1/1	0.94	0.67	31,31,31,31	0
55	MG	2a	1793	1/1	0.94	0.18	57,57,57,57	0
55	MG	2A	3639	1/1	0.94	0.05	44,44,44,44	0
55	MG	2A	3484	1/1	0.94	0.22	29,29,29,29	0
55	MG	1A	3403	1/1	0.94	0.11	30,30,30,30	0
55	MG	1A	3754	1/1	0.94	0.14	33,33,33,33	0
55	MG	2A	3820	1/1	0.94	0.11	33,33,33,33	0
55	MG	1A	3300	1/1	0.94	0.15	27,27,27,27	0
55	MG	1A	3984	1/1	0.94	0.18	43,43,43,43	0
55	MG	1A	3635	1/1	0.94	0.12	22,22,22,22	0
55	MG	1A	3694	1/1	0.94	0.10	35,35,35,35	0
55	MG	2A	3300	1/1	0.94	0.14	31,31,31,31	0
55	MG	11	102	1/1	0.94	0.12	28,28,28,28	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3402	1/1	0.94	0.16	35,35,35,35	0
55	MG	1a	3127	1/1	0.94	0.13	21,21,21,21	0
55	MG	2A	3465	1/1	0.94	0.15	50,50,50,50	0
55	MG	2A	3197	1/1	0.94	0.20	34,34,34,34	0
55	MG	2A	3388	1/1	0.94	0.17	34,34,34,34	0
55	MG	1a	3070	1/1	0.94	0.13	25,25,25,25	0
55	MG	1A	3544	1/1	0.94	0.30	40,40,40,40	0
55	MG	2Z	302	1/1	0.94	0.21	26,26,26,26	0
55	MG	2A	3758	1/1	0.94	0.15	59,59,59,59	0
55	MG	2A	3337	1/1	0.94	0.10	35,35,35,35	0
55	MG	2a	1655	1/1	0.94	0.12	57,57,57,57	0
55	MG	1A	3982	1/1	0.94	0.13	36,36,36,36	0
55	MG	1N	202	1/1	0.94	0.33	41,41,41,41	0
55	MG	1A	3778	1/1	0.94	0.08	55,55,55,55	0
55	MG	2A	3177	1/1	0.94	0.21	33,33,33,33	0
55	MG	1A	3148	1/1	0.94	0.29	26,26,26,26	0
55	MG	2A	3306	1/1	0.94	0.23	19,19,19,19	0
55	MG	2A	3622	1/1	0.94	0.16	37,37,37,37	0
55	MG	2A	3318	1/1	0.94	0.28	44,44,44,44	0
55	MG	1A	3106	1/1	0.94	0.29	33,33,33,33	0
55	MG	1A	3245	1/1	0.94	0.22	48,48,48,48	0
55	MG	1A	3035	1/1	0.94	0.18	18,18,18,18	0
55	MG	2A	3513	1/1	0.94	0.13	56,56,56,56	0
55	MG	2A	3060	1/1	0.94	0.11	41,41,41,41	0
55	MG	1A	3526	1/1	0.94	0.21	31,31,31,31	0
55	MG	1a	3023	1/1	0.94	0.18	44,44,44,44	0
55	MG	2A	3632	1/1	0.94	0.13	37,37,37,37	0
55	MG	2A	3372	1/1	0.94	0.15	42,42,42,42	0
55	MG	2a	1674	1/1	0.94	0.29	67,67,67,67	0
55	MG	2A	3790	1/1	0.94	0.25	42,42,42,42	0
55	MG	1A	3319	1/1	0.94	0.42	16,16,16,16	0
55	MG	2A	3261	1/1	0.94	0.38	43,43,43,43	0
55	MG	1a	3137	1/1	0.94	0.16	34,34,34,34	0
55	MG	1A	4018	1/1	0.94	0.11	40,40,40,40	0
55	MG	1a	3073	1/1	0.94	0.13	50,50,50,50	0
55	MG	2a	1731	1/1	0.94	0.10	33,33,33,33	0
55	MG	2A	3233	1/1	0.94	0.38	50,50,50,50	0
55	MG	1A	3837	1/1	0.94	0.14	51,51,51,51	0
55	MG	1A	3416	1/1	0.94	0.22	44,44,44,44	0
55	MG	2A	3486	1/1	0.94	0.12	29,29,29,29	0
55	MG	2A	3276	1/1	0.94	0.23	59,59,59,59	0
55	MG	1A	3948	1/1	0.94	0.13	43,43,43,43	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2a	1812	1/1	0.94	0.14	54,54,54,54	0
55	MG	1A	3758	1/1	0.94	0.87	49,49,49,49	0
55	MG	1D	301	1/1	0.94	0.55	22,22,22,22	0
55	MG	1a	3203	1/1	0.94	0.08	39,39,39,39	0
55	MG	2A	3470	1/1	0.94	0.22	21,21,21,21	0
55	MG	2A	3189	1/1	0.94	0.12	32,32,32,32	0
55	MG	1A	3789	1/1	0.94	0.15	24,24,24,24	0
55	MG	2a	1697	1/1	0.94	0.15	23,23,23,23	0
55	MG	2A	3691	1/1	0.94	0.12	24,24,24,24	0
55	MG	2A	3644	1/1	0.94	0.18	52,52,52,52	0
55	MG	2A	3164	1/1	0.94	0.14	36,36,36,36	0
55	MG	12	101	1/1	0.94	0.33	34,34,34,34	0
55	MG	2A	3146	1/1	0.94	0.14	40,40,40,40	0
55	MG	2A	3081	1/1	0.94	0.16	54,54,54,54	0
55	MG	2A	3686	1/1	0.94	0.16	31,31,31,31	0
55	MG	2i	201	1/1	0.94	0.08	70,70,70,70	0
55	MG	1A	3991	1/1	0.94	0.15	34,34,34,34	0
55	MG	2A	3724	1/1	0.94	0.19	41,41,41,41	0
55	MG	17	107	1/1	0.94	0.35	32,32,32,32	0
55	MG	1A	3392	1/1	0.94	0.72	31,31,31,31	0
55	MG	2A	3072	1/1	0.94	0.38	26,26,26,26	0
55	MG	1B	205	1/1	0.94	0.14	44,44,44,44	0
55	MG	1A	3206	1/1	0.94	0.09	28,28,28,28	0
55	MG	1A	4007	1/1	0.94	0.21	50,50,50,50	0
55	MG	2A	3666	1/1	0.94	0.10	26,26,26,26	0
55	MG	1A	3665	1/1	0.94	0.13	10,10,10,10	0
55	MG	1A	3103	1/1	0.94	0.14	32,32,32,32	0
55	MG	1A	3272	1/1	0.94	0.08	23,23,23,23	0
55	MG	1A	4002	1/1	0.94	0.24	17,17,17,17	0
55	MG	1a	3014	1/1	0.94	0.12	33,33,33,33	0
55	MG	1A	3878	1/1	0.94	0.19	20,20,20,20	0
55	MG	1A	3459	1/1	0.94	0.12	28,28,28,28	0
55	MG	2A	3669	1/1	0.94	0.38	37,37,37,37	0
55	MG	1A	4021	1/1	0.94	0.17	42,42,42,42	0
55	MG	1a	3078	1/1	0.94	0.47	19,19,19,19	0
55	MG	2A	3313	1/1	0.94	0.17	52,52,52,52	0
55	MG	2A	3405	1/1	0.94	0.45	41,41,41,41	0
55	MG	1A	3944	1/1	0.94	0.08	20,20,20,20	0
55	MG	2A	3694	1/1	0.94	0.16	21,21,21,21	0
55	MG	2A	3330	1/1	0.94	0.30	26,26,26,26	0
55	MG	21	106	1/1	0.94	0.26	35,35,35,35	0
55	MG	2B	219	1/1	0.94	0.10	63,63,63,63	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3712	1/1	0.94	0.08	36,36,36,36	0
55	MG	2A	3148	1/1	0.94	0.13	31,31,31,31	0
55	MG	2A	3779	1/1	0.94	0.14	34,34,34,34	0
55	MG	1A	3328	1/1	0.94	0.12	22,22,22,22	0
55	MG	2I	202	1/1	0.94	0.11	42,42,42,42	0
55	MG	1A	3467	1/1	0.94	0.28	28,28,28,28	0
55	MG	1A	3522	1/1	0.94	0.16	33,33,33,33	0
55	MG	1R	206	1/1	0.94	0.17	24,24,24,24	0
55	MG	1A	3127	1/1	0.94	0.63	18,18,18,18	0
55	MG	2A	3464	1/1	0.94	0.13	20,20,20,20	0
55	MG	1A	3581	1/1	0.94	0.11	15,15,15,15	0
55	MG	2A	3305	1/1	0.94	0.15	29,29,29,29	0
55	MG	1B	220	1/1	0.94	0.15	25,25,25,25	0
55	MG	1A	3761	1/1	0.94	0.12	51,51,51,51	0
55	MG	1A	3345	1/1	0.94	0.14	21,21,21,21	0
55	MG	1A	3381	1/1	0.94	0.12	22,22,22,22	0
55	MG	1A	3078	1/1	0.94	0.18	23,23,23,23	0
55	MG	1A	3255	1/1	0.94	0.26	32,32,32,32	0
55	MG	1A	3434	1/1	0.94	0.12	43,43,43,43	0
55	MG	10	106	1/1	0.94	0.08	35,35,35,35	0
55	MG	1A	3816	1/1	0.94	0.16	15,15,15,15	0
55	MG	1A	3749	1/1	0.94	0.14	53,53,53,53	0
55	MG	2A	3522	1/1	0.94	0.20	34,34,34,34	0
55	MG	1A	3361	1/1	0.94	0.31	37,37,37,37	0
55	MG	2A	3066	1/1	0.94	0.12	46,46,46,46	0
55	MG	2A	3573	1/1	0.94	0.13	23,23,23,23	0
55	MG	1A	3999	1/1	0.94	0.18	32,32,32,32	0
55	MG	2A	3671	1/1	0.94	0.09	35,35,35,35	0
55	MG	1Z	302	1/1	0.94	0.13	32,32,32,32	0
55	MG	2a	1638	1/1	0.94	0.12	42,42,42,42	0
55	MG	26	101	1/1	0.94	0.17	40,40,40,40	0
55	MG	1A	3558	1/1	0.94	0.16	21,21,21,21	0
55	MG	1a	3052	1/1	0.94	0.17	53,53,53,53	0
55	MG	1A	3334	1/1	0.94	0.80	51,51,51,51	0
55	MG	1A	3961	1/1	0.94	0.13	37,37,37,37	0
55	MG	1A	3017	1/1	0.94	0.18	10,10,10,10	0
55	MG	2A	3253	1/1	0.94	0.09	30,30,30,30	0
55	MG	1A	3555	1/1	0.94	0.53	34,34,34,34	0
55	MG	1A	3071	1/1	0.94	0.53	35,35,35,35	0
55	MG	1A	3105	1/1	0.94	0.39	19,19,19,19	0
55	MG	1a	3109	1/1	0.94	0.10	40,40,40,40	0
55	MG	1A	3490	1/1	0.94	0.17	35,35,35,35	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3334	1/1	0.94	0.07	30,30,30,30	0
55	MG	1A	3408	1/1	0.94	0.10	18,18,18,18	0
55	MG	1A	3810	1/1	0.94	0.27	23,23,23,23	0
55	MG	2A	3601	1/1	0.94	0.21	44,44,44,44	0
55	MG	1A	3426	1/1	0.94	0.29	15,15,15,15	0
55	MG	2A	3366	1/1	0.94	0.71	31,31,31,31	0
55	MG	2A	3374	1/1	0.94	0.13	35,35,35,35	0
55	MG	1x	105	1/1	0.94	0.19	54,54,54,54	0
55	MG	1a	3128	1/1	0.94	0.32	30,30,30,30	0
55	MG	2A	3402	1/1	0.94	0.10	34,34,34,34	0
55	MG	2A	3105	1/1	0.94	0.40	45,45,45,45	0
55	MG	1A	3029	1/1	0.94	0.45	23,23,23,23	0
55	MG	2a	1811	1/1	0.94	0.13	42,42,42,42	0
55	MG	2A	3089	1/1	0.94	0.12	32,32,32,32	0
55	MG	2a	1796	1/1	0.94	0.17	50,50,50,50	0
55	MG	1A	3922	1/1	0.94	0.14	68,68,68,68	0
55	MG	1a	3064	1/1	0.94	0.22	59,59,59,59	0
55	MG	1A	3763	1/1	0.94	0.09	37,37,37,37	0
55	MG	2A	3810	1/1	0.94	0.09	29,29,29,29	0
55	MG	2A	3124	1/1	0.94	0.15	29,29,29,29	0
55	MG	1A	3579	1/1	0.94	0.16	15,15,15,15	0
55	MG	2A	3035	1/1	0.94	0.11	30,30,30,30	0
55	MG	1a	3031	1/1	0.94	0.15	29,29,29,29	0
55	MG	1A	3580	1/1	0.94	0.21	42,42,42,42	0
55	MG	1A	4020	1/1	0.94	0.08	35,35,35,35	0
55	MG	2D	302	1/1	0.94	0.23	28,28,28,28	0
55	MG	2a	1773	1/1	0.95	0.23	48,48,48,48	0
55	MG	1a	3211	1/1	0.95	0.32	36,36,36,36	0
55	MG	2E	302	1/1	0.95	0.13	27,27,27,27	0
55	MG	2A	3507	1/1	0.95	0.19	10,10,10,10	0
55	MG	2A	3052	1/1	0.95	0.23	34,34,34,34	0
55	MG	1A	3872	1/1	0.95	0.26	52,52,52,52	0
55	MG	1A	3170	1/1	0.95	0.13	32,32,32,32	0
55	MG	1A	4006	1/1	0.95	0.11	42,42,42,42	0
55	MG	2a	1668	1/1	0.95	0.14	28,28,28,28	0
55	MG	1A	3671	1/1	0.95	0.13	28,28,28,28	0
55	MG	2a	1738	1/1	0.95	0.12	59,59,59,59	0
55	MG	1A	3690	1/1	0.95	0.10	27,27,27,27	0
55	MG	2a	1734	1/1	0.95	0.15	40,40,40,40	0
55	MG	1A	3137	1/1	0.95	0.27	30,30,30,30	0
55	MG	17	101	1/1	0.95	0.55	19,19,19,19	0
55	MG	2a	1783	1/1	0.95	0.08	56,56,56,56	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3775	1/1	0.95	0.10	40,40,40,40	0
55	MG	2a	1799	1/1	0.95	0.18	49,49,49,49	0
55	MG	1E	308	1/1	0.95	0.52	34,34,34,34	0
55	MG	2A	3749	1/1	0.95	0.11	35,35,35,35	0
55	MG	1A	3811	1/1	0.95	0.11	39,39,39,39	0
55	MG	2A	3159	1/1	0.95	0.10	31,31,31,31	0
55	MG	1A	3368	1/1	0.95	0.14	29,29,29,29	0
55	MG	2A	3818	1/1	0.95	0.10	30,30,30,30	0
55	MG	2A	3220	1/1	0.95	0.26	38,38,38,38	0
55	MG	2x	101	1/1	0.95	0.14	35,35,35,35	0
55	MG	2a	1804	1/1	0.95	0.12	31,31,31,31	0
55	MG	1A	3488	1/1	0.95	0.11	30,30,30,30	0
55	MG	2a	1693	1/1	0.95	0.10	57,57,57,57	0
55	MG	1A	3786	1/1	0.95	0.22	37,37,37,37	0
55	MG	1A	3797	1/1	0.95	0.13	20,20,20,20	0
55	MG	1B	221	1/1	0.95	0.09	56,56,56,56	0
55	MG	1A	3661	1/1	0.95	0.11	45,45,45,45	0
55	MG	1x	110	1/1	0.95	0.22	29,29,29,29	0
55	MG	1A	3882	1/1	0.95	0.07	24,24,24,24	0
55	MG	1A	3183	1/1	0.95	0.41	29,29,29,29	0
55	MG	2A	3547	1/1	0.95	0.18	29,29,29,29	0
55	MG	1A	3168	1/1	0.95	0.41	31,31,31,31	0
55	MG	1A	3039	1/1	0.95	0.34	38,38,38,38	0
55	MG	1A	3953	1/1	0.95	0.11	46,46,46,46	0
55	MG	2A	3608	1/1	0.95	0.14	31,31,31,31	0
55	MG	1A	3551	1/1	0.95	0.18	19,19,19,19	0
55	MG	1A	3541	1/1	0.95	0.17	32,32,32,32	0
55	MG	1A	3151	1/1	0.95	0.11	25,25,25,25	0
55	MG	2A	3631	1/1	0.95	0.14	19,19,19,19	0
55	MG	1A	3806	1/1	0.95	0.06	30,30,30,30	0
55	MG	1d	301	1/1	0.95	0.14	21,21,21,21	0
55	MG	18	103	1/1	0.95	0.81	40,40,40,40	0
55	MG	1A	3299	1/1	0.95	0.18	34,34,34,34	0
55	MG	1A	3639	1/1	0.95	0.24	15,15,15,15	0
55	MG	1A	3642	1/1	0.95	0.18	21,21,21,21	0
55	MG	15	104	1/1	0.95	0.10	17,17,17,17	0
55	MG	2A	3093	1/1	0.95	0.28	22,22,22,22	0
55	MG	1A	3476	1/1	0.95	0.40	38,38,38,38	0
55	MG	1A	3128	1/1	0.95	0.17	23,23,23,23	0
55	MG	2A	3760	1/1	0.95	0.46	50,50,50,50	0
55	MG	1a	3185	1/1	0.95	0.17	32,32,32,32	0
55	MG	1A	3397	1/1	0.95	0.21	16,16,16,16	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1a	3161	1/1	0.95	0.11	41,41,41,41	0
55	MG	1A	3292	1/1	0.95	0.14	25,25,25,25	0
55	MG	2A	3357	1/1	0.95	0.13	14,14,14,14	0
55	MG	2A	3343	1/1	0.95	0.11	25,25,25,25	0
55	MG	1A	3550	1/1	0.95	0.13	24,24,24,24	0
55	MG	1A	3870	1/1	0.95	0.08	14,14,14,14	0
55	MG	1A	3044	1/1	0.95	0.10	22,22,22,22	0
55	MG	2a	1708	1/1	0.95	0.09	69,69,69,69	0
55	MG	2F	305	1/1	0.95	0.09	27,27,27,27	0
55	MG	1D	306	1/1	0.95	0.11	35,35,35,35	0
55	MG	1A	3731	1/1	0.95	0.21	25,25,25,25	0
55	MG	1v	105	1/1	0.95	0.13	35,35,35,35	0
55	MG	1a	3196	1/1	0.95	0.08	34,34,34,34	0
55	MG	1A	3016	1/1	0.95	0.11	28,28,28,28	0
55	MG	1A	3531	1/1	0.95	0.13	30,30,30,30	0
55	MG	2A	3210	1/1	0.95	0.12	15,15,15,15	0
55	MG	1A	3394	1/1	0.95	0.36	31,31,31,31	0
55	MG	1A	4032	1/1	0.95	0.14	39,39,39,39	0
55	MG	1A	3032	1/1	0.95	0.40	16,16,16,16	0
55	MG	1W	207	1/1	0.95	0.20	23,23,23,23	0
55	MG	1H	201	1/1	0.95	0.18	32,32,32,32	0
55	MG	2A	3498	1/1	0.95	0.40	32,32,32,32	0
55	MG	2A	3729	1/1	0.95	0.08	24,24,24,24	0
55	MG	1A	3309	1/1	0.95	0.51	25,25,25,25	0
55	MG	2A	3562	1/1	0.95	0.15	29,29,29,29	0
55	MG	1A	3285	1/1	0.95	0.16	42,42,42,42	0
55	MG	1A	3638	1/1	0.95	0.12	10,10,10,10	0
55	MG	2A	3772	1/1	0.95	0.14	31,31,31,31	0
55	MG	1a	3037	1/1	0.95	0.10	39,39,39,39	0
55	MG	1W	201	1/1	0.95	0.30	35,35,35,35	0
55	MG	2a	1758	1/1	0.95	0.09	46,46,46,46	0
55	MG	1A	3395	1/1	0.95	0.22	13,13,13,13	0
55	MG	1A	4030	1/1	0.95	0.49	20,20,20,20	0
55	MG	1A	3964	1/1	0.95	0.14	40,40,40,40	0
55	MG	2a	1801	1/1	0.95	0.15	47,47,47,47	0
55	MG	1A	3942	1/1	0.95	0.24	35,35,35,35	0
55	MG	1A	3539	1/1	0.95	0.14	20,20,20,20	0
55	MG	1B	204	1/1	0.95	0.14	25,25,25,25	0
55	MG	2A	3503	1/1	0.95	0.17	26,26,26,26	0
55	MG	2A	3335	1/1	0.95	0.13	25,25,25,25	0
56	K	2A	3433	1/1	0.95	0.17	43,43,43,43	0
55	MG	1a	3021	1/1	0.95	0.10	34,34,34,34	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3860	1/1	0.95	0.24	45,45,45,45	0
55	MG	1A	3985	1/1	0.95	0.22	37,37,37,37	0
55	MG	1A	4041	1/1	0.95	0.53	29,29,29,29	0
55	MG	1B	222	1/1	0.95	0.17	22,22,22,22	0
55	MG	1A	3379	1/1	0.95	0.20	27,27,27,27	0
55	MG	2A	3138	1/1	0.95	0.34	31,31,31,31	0
55	MG	2A	3843	1/1	0.95	0.11	39,39,39,39	0
55	MG	1F	302	1/1	0.95	0.52	16,16,16,16	0
55	MG	1A	3024	1/1	0.95	0.40	38,38,38,38	0
55	MG	1A	3171	1/1	0.95	0.18	34,34,34,34	0
55	MG	1A	3380	1/1	0.95	0.50	24,24,24,24	0
55	MG	1A	3501	1/1	0.95	0.43	36,36,36,36	0
55	MG	10	102	1/1	0.95	0.17	34,34,34,34	0
55	MG	2A	3230	1/1	0.95	0.07	42,42,42,42	0
55	MG	2T	204	1/1	0.95	0.17	39,39,39,39	0
55	MG	1A	3968	1/1	0.95	0.13	42,42,42,42	0
55	MG	1A	3833	1/1	0.95	0.20	28,28,28,28	0
55	MG	1a	3015	1/1	0.95	0.14	31,31,31,31	0
55	MG	1A	3157	1/1	0.95	0.25	29,29,29,29	0
55	MG	13	104	1/1	0.95	0.11	38,38,38,38	0
55	MG	2a	1747	1/1	0.95	0.15	29,29,29,29	0
55	MG	2A	3437	1/1	0.95	0.44	37,37,37,37	0
55	MG	1A	3233	1/1	0.95	0.14	24,24,24,24	0
55	MG	2A	3703	1/1	0.95	0.12	30,30,30,30	0
55	MG	1A	3213	1/1	0.95	0.50	35,35,35,35	0
55	MG	1A	3696	1/1	0.95	0.18	25,25,25,25	0
55	MG	1A	3294	1/1	0.95	0.09	10,10,10,10	0
55	MG	2A	3140	1/1	0.95	0.16	37,37,37,37	0
55	MG	1U	203	1/1	0.95	0.36	32,32,32,32	0
55	MG	2A	3216	1/1	0.95	0.18	28,28,28,28	0
55	MG	2A	3136	1/1	0.95	0.12	39,39,39,39	0
55	MG	1A	3681	1/1	0.95	0.17	22,22,22,22	0
55	MG	1A	3347	1/1	0.95	0.35	34,34,34,34	0
55	MG	2F	303	1/1	0.95	0.09	30,30,30,30	0
55	MG	1A	3989	1/1	0.95	0.11	35,35,35,35	0
55	MG	1F	301	1/1	0.95	0.83	29,29,29,29	0
55	MG	1a	3165	1/1	0.95	0.10	48,48,48,48	0
55	MG	1A	3188	1/1	0.95	0.21	22,22,22,22	0
55	MG	1n	101	1/1	0.95	0.10	34,34,34,34	0
55	MG	1A	3836	1/1	0.95	0.08	45,45,45,45	0
55	MG	1A	3721	1/1	0.95	0.12	26,26,26,26	0
55	MG	2a	1785	1/1	0.95	0.07	37,37,37,37	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3797	1/1	0.95	0.15	21,21,21,21	0
55	MG	1A	3008	1/1	0.95	0.15	21,21,21,21	0
55	MG	2A	3119	1/1	0.95	0.26	29,29,29,29	0
55	MG	1A	3876	1/1	0.95	0.35	42,42,42,42	0
55	MG	1a	3095	1/1	0.95	0.50	28,28,28,28	0
55	MG	2A	3610	1/1	0.95	0.77	41,41,41,41	0
55	MG	2B	210	1/1	0.95	0.17	45,45,45,45	0
56	K	1A	3470	1/1	0.95	0.28	51,51,51,51	0
55	MG	1A	3344	1/1	0.95	0.40	19,19,19,19	0
55	MG	2A	3690	1/1	0.95	0.13	43,43,43,43	0
55	MG	2a	1692	1/1	0.95	0.11	46,46,46,46	0
55	MG	1B	210	1/1	0.95	0.13	30,30,30,30	0
55	MG	2a	1807	1/1	0.95	0.18	45,45,45,45	0
55	MG	2A	3564	1/1	0.95	0.10	25,25,25,25	0
55	MG	1A	3791	1/1	0.95	0.12	46,46,46,46	0
55	MG	2A	3011	1/1	0.95	0.53	41,41,41,41	0
55	MG	1A	3374	1/1	0.95	0.41	33,33,33,33	0
55	MG	2A	3808	1/1	0.95	0.08	22,22,22,22	0
55	MG	1A	3235	1/1	0.95	0.15	20,20,20,20	0
55	MG	1N	205	1/1	0.95	0.12	22,22,22,22	0
55	MG	2B	202	1/1	0.95	0.13	40,40,40,40	0
55	MG	1A	3934	1/1	0.95	0.10	49,49,49,49	0
55	MG	1A	3974	1/1	0.95	0.12	30,30,30,30	0
55	MG	1A	3766	1/1	0.95	0.10	26,26,26,26	0
55	MG	1A	3930	1/1	0.95	0.18	17,17,17,17	0
55	MG	1A	3239	1/1	0.95	0.14	43,43,43,43	0
55	MG	1A	3373	1/1	0.95	0.24	28,28,28,28	0
55	MG	2A	3080	1/1	0.95	0.30	26,26,26,26	0
55	MG	2A	3071	1/1	0.95	0.39	45,45,45,45	0
55	MG	2a	1765	1/1	0.95	0.13	42,42,42,42	0
55	MG	1A	3021	1/1	0.95	0.16	21,21,21,21	0
55	MG	2A	3393	1/1	0.95	0.15	31,31,31,31	0
55	MG	1A	3130	1/1	0.95	0.57	18,18,18,18	0
55	MG	2A	3077	1/1	0.95	0.10	23,23,23,23	0
55	MG	1A	3265	1/1	0.95	0.23	19,19,19,19	0
55	MG	1A	3329	1/1	0.95	0.53	25,25,25,25	0
55	MG	2B	209	1/1	0.95	0.08	40,40,40,40	0
55	MG	2A	3786	1/1	0.95	0.13	37,37,37,37	0
55	MG	2A	3688	1/1	0.95	0.20	53,53,53,53	0
55	MG	1F	307	1/1	0.95	0.27	30,30,30,30	0
55	MG	2j	201	1/1	0.95	0.08	64,64,64,64	0
55	MG	2A	3157	1/1	0.95	0.24	20,20,20,20	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3826	1/1	0.95	0.12	29,29,29,29	0
55	MG	1a	3080	1/1	0.95	0.09	63,63,63,63	0
55	MG	1A	3058	1/1	0.95	0.18	39,39,39,39	0
55	MG	1a	3057	1/1	0.95	0.08	31,31,31,31	0
55	MG	2A	3552	1/1	0.95	0.22	15,15,15,15	0
55	MG	1A	3517	1/1	0.95	0.15	34,34,34,34	0
55	MG	1A	3097	1/1	0.95	0.57	30,30,30,30	0
55	MG	1A	3175	1/1	0.95	0.28	26,26,26,26	0
55	MG	2A	3204	1/1	0.95	0.19	18,18,18,18	0
55	MG	2A	3842	1/1	0.95	0.15	42,42,42,42	0
55	MG	1a	3040	1/1	0.95	0.10	34,34,34,34	0
55	MG	1A	3466	1/1	0.95	1.04	38,38,38,38	0
55	MG	2A	3831	1/1	0.95	0.23	35,35,35,35	0
55	MG	1A	3990	1/1	0.95	0.10	25,25,25,25	0
55	MG	2A	3699	1/1	0.95	0.10	20,20,20,20	0
55	MG	2A	3568	1/1	0.95	0.14	50,50,50,50	0
55	MG	2A	3827	1/1	0.95	0.18	45,45,45,45	0
55	MG	2A	3139	1/1	0.95	0.20	22,22,22,22	0
55	MG	2A	3211	1/1	0.95	0.67	26,26,26,26	0
55	MG	1A	3491	1/1	0.95	0.27	37,37,37,37	0
55	MG	2A	3485	1/1	0.95	0.10	22,22,22,22	0
55	MG	10	105	1/1	0.95	0.17	29,29,29,29	0
55	MG	1A	3804	1/1	0.95	0.17	35,35,35,35	0
55	MG	1A	3538	1/1	0.95	0.27	29,29,29,29	0
55	MG	1A	4000	1/1	0.95	0.14	32,32,32,32	0
55	MG	1a	3082	1/1	0.95	0.16	28,28,28,28	0
55	MG	1A	3698	1/1	0.95	0.13	12,12,12,12	0
55	MG	1A	3949	1/1	0.95	0.08	31,31,31,31	0
55	MG	1A	3891	1/1	0.95	0.10	42,42,42,42	0
55	MG	1A	3251	1/1	0.95	0.22	26,26,26,26	0
55	MG	1A	3670	1/1	0.95	0.19	57,57,57,57	0
55	MG	2A	3130	1/1	0.95	0.23	28,28,28,28	0
55	MG	2A	3611	1/1	0.95	0.30	43,43,43,43	0
55	MG	2A	3491	1/1	0.95	0.16	27,27,27,27	0
55	MG	2A	3523	1/1	0.95	0.33	27,27,27,27	0
55	MG	1A	3087	1/1	0.95	0.09	30,30,30,30	0
55	MG	2A	3700	1/1	0.95	0.20	31,31,31,31	0
55	MG	2A	3297	1/1	0.95	0.23	17,17,17,17	0
55	MG	1A	3248	1/1	0.95	0.09	16,16,16,16	0
55	MG	2A	3828	1/1	0.95	0.13	24,24,24,24	0
55	MG	2A	3574	1/1	0.95	0.08	31,31,31,31	0
55	MG	2a	1701	1/1	0.95	0.11	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3659	1/1	0.95	0.12	31,31,31,31	0
55	MG	1A	4033	1/1	0.95	0.11	42,42,42,42	0
55	MG	1A	3866	1/1	0.95	0.15	62,62,62,62	0
55	MG	2A	3097	1/1	0.95	0.27	40,40,40,40	0
55	MG	2A	3750	1/1	0.95	0.10	41,41,41,41	0
55	MG	1a	3025	1/1	0.95	0.16	30,30,30,30	0
55	MG	2a	1763	1/1	0.95	0.08	31,31,31,31	0
55	MG	1A	3005	1/1	0.95	0.34	20,20,20,20	0
55	MG	1a	3043	1/1	0.95	0.12	35,35,35,35	0
55	MG	1A	3909	1/1	0.95	0.28	80,80,80,80	0
55	MG	2A	3144	1/1	0.95	0.17	44,44,44,44	0
55	MG	2A	3459	1/1	0.95	0.12	29,29,29,29	0
55	MG	1A	3880	1/1	0.95	0.08	55,55,55,55	0
55	MG	2A	3322	1/1	0.95	0.11	33,33,33,33	0
55	MG	2A	3354	1/1	0.95	0.07	31,31,31,31	0
55	MG	1Y	202	1/1	0.95	0.07	44,44,44,44	0
55	MG	1A	3350	1/1	0.95	0.16	28,28,28,28	0
55	MG	2A	3782	1/1	0.95	0.23	37,37,37,37	0
55	MG	1A	3955	1/1	0.95	0.11	39,39,39,39	0
55	MG	1A	3992	1/1	0.95	0.39	40,40,40,40	0
55	MG	1A	3455	1/1	0.95	0.12	35,35,35,35	0
55	MG	1A	3621	1/1	0.95	0.24	36,36,36,36	0
55	MG	2A	3757	1/1	0.95	0.09	34,34,34,34	0
55	MG	2a	1648	1/1	0.95	0.46	58,58,58,58	0
55	MG	1A	3519	1/1	0.95	0.18	26,26,26,26	0
55	MG	2B	205	1/1	0.95	0.16	34,34,34,34	0
55	MG	2A	3431	1/1	0.95	0.14	37,37,37,37	0
55	MG	2A	3657	1/1	0.95	0.36	50,50,50,50	0
55	MG	1A	3542	1/1	0.95	0.25	51,51,51,51	0
55	MG	1D	304	1/1	0.95	0.81	30,30,30,30	0
55	MG	2A	3753	1/1	0.95	0.11	31,31,31,31	0
55	MG	2A	3509	1/1	0.95	0.09	26,26,26,26	0
55	MG	1A	3851	1/1	0.95	0.11	43,43,43,43	0
55	MG	1A	3869	1/1	0.95	0.10	61,61,61,61	0
55	MG	1A	3102	1/1	0.95	0.36	17,17,17,17	0
55	MG	2B	215	1/1	0.95	0.17	60,60,60,60	0
55	MG	1A	3905	1/1	0.95	0.15	63,63,63,63	0
55	MG	1A	3619	1/1	0.95	0.10	23,23,23,23	0
55	MG	2D	303	1/1	0.95	0.29	40,40,40,40	0
55	MG	2A	3430	1/1	0.95	0.11	29,29,29,29	0
55	MG	1A	3089	1/1	0.95	0.19	39,39,39,39	0
55	MG	1A	3675	1/1	0.95	0.14	43,43,43,43	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3198	1/1	0.95	0.25	26,26,26,26	0
55	MG	1A	3561	1/1	0.95	0.24	52,52,52,52	0
55	MG	1A	3772	1/1	0.95	0.56	47,47,47,47	0
55	MG	10	101	1/1	0.95	0.38	33,33,33,33	0
55	MG	2A	3592	1/1	0.95	0.24	31,31,31,31	0
55	MG	2Y	201	1/1	0.95	0.23	27,27,27,27	0
55	MG	2A	3102	1/1	0.95	0.12	31,31,31,31	0
55	MG	1A	3959	1/1	0.95	0.13	56,56,56,56	0
55	MG	1G	203	1/1	0.95	0.12	40,40,40,40	0
55	MG	2A	3167	1/1	0.95	0.17	51,51,51,51	0
55	MG	2a	1637	1/1	0.95	0.83	39,39,39,39	0
55	MG	2A	3821	1/1	0.95	0.29	33,33,33,33	0
55	MG	1A	3346	1/1	0.95	0.29	15,15,15,15	0
55	MG	1A	3485	1/1	0.96	0.13	39,39,39,39	0
55	MG	28	102	1/1	0.96	0.12	29,29,29,29	0
55	MG	1A	3461	1/1	0.96	0.18	38,38,38,38	0
55	MG	2A	3602	1/1	0.96	0.15	36,36,36,36	0
55	MG	1A	3225	1/1	0.96	0.18	54,54,54,54	0
55	MG	2A	3410	1/1	0.96	0.10	43,43,43,43	0
55	MG	2A	3247	1/1	0.96	0.15	25,25,25,25	0
55	MG	2A	3062	1/1	0.96	0.11	17,17,17,17	0
55	MG	1U	206	1/1	0.96	0.29	19,19,19,19	0
55	MG	1A	3604	1/1	0.96	0.11	20,20,20,20	0
55	MG	1W	205	1/1	0.96	0.61	34,34,34,34	0
55	MG	1F	306	1/1	0.96	0.35	11,11,11,11	0
55	MG	2A	3620	1/1	0.96	0.26	21,21,21,21	0
55	MG	2A	3404	1/1	0.96	0.09	12,12,12,12	0
55	MG	1a	3056	1/1	0.96	0.10	37,37,37,37	0
55	MG	1A	3031	1/1	0.96	0.74	26,26,26,26	0
55	MG	2A	3563	1/1	0.96	0.18	27,27,27,27	0
55	MG	2A	3594	1/1	0.96	0.51	37,37,37,37	0
55	MG	2A	3419	1/1	0.96	0.13	31,31,31,31	0
55	MG	2B	218	1/1	0.96	0.21	37,37,37,37	0
55	MG	2A	3598	1/1	0.96	0.19	32,32,32,32	0
55	MG	2A	3833	1/1	0.96	0.09	47,47,47,47	0
55	MG	2A	3307	1/1	0.96	0.29	38,38,38,38	0
55	MG	1A	3946	1/1	0.96	0.10	30,30,30,30	0
55	MG	1A	3177	1/1	0.96	0.10	26,26,26,26	0
55	MG	2A	3609	1/1	0.96	0.11	27,27,27,27	0
55	MG	1A	3926	1/1	0.96	0.06	34,34,34,34	0
55	MG	2A	3269	1/1	0.96	0.11	30,30,30,30	0
55	MG	1A	3504	1/1	0.96	0.50	25,25,25,25	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	4022	1/1	0.96	0.13	27,27,27,27	0
55	MG	1A	4017	1/1	0.96	0.11	25,25,25,25	0
55	MG	2a	1781	1/1	0.96	0.06	53,53,53,53	0
55	MG	1A	3092	1/1	0.96	0.29	22,22,22,22	0
55	MG	1A	3682	1/1	0.96	0.11	56,56,56,56	0
55	MG	2A	3640	1/1	0.96	0.35	39,39,39,39	0
55	MG	1A	3091	1/1	0.96	0.21	32,32,32,32	0
55	MG	1a	3055	1/1	0.96	0.10	46,46,46,46	0
55	MG	18	105	1/1	0.96	0.13	35,35,35,35	0
55	MG	1A	3074	1/1	0.96	0.39	19,19,19,19	0
55	MG	15	103	1/1	0.96	0.16	23,23,23,23	0
55	MG	10	103	1/1	0.96	0.18	25,25,25,25	0
55	MG	1A	3212	1/1	0.96	0.08	10,10,10,10	0
55	MG	1A	3242	1/1	0.96	1.00	28,28,28,28	0
55	MG	1B	216	1/1	0.96	0.23	67,67,67,67	0
55	MG	1A	3449	1/1	0.96	0.21	15,15,15,15	0
55	MG	2A	3315	1/1	0.96	0.07	14,14,14,14	0
55	MG	2A	3668	1/1	0.96	0.13	29,29,29,29	0
55	MG	1A	3176	1/1	0.96	0.17	21,21,21,21	0
55	MG	2A	3147	1/1	0.96	0.26	51,51,51,51	0
55	MG	1a	3036	1/1	0.96	0.14	34,34,34,34	0
55	MG	1A	3014	1/1	0.96	0.13	30,30,30,30	0
55	MG	2A	3458	1/1	0.96	0.10	28,28,28,28	0
55	MG	2A	3348	1/1	0.96	0.62	32,32,32,32	0
55	MG	2A	3110	1/1	0.96	0.18	17,17,17,17	0
55	MG	2A	3508	1/1	0.96	0.13	15,15,15,15	0
55	MG	1A	3207	1/1	0.96	0.16	22,22,22,22	0
55	MG	1a	3183	1/1	0.96	0.12	44,44,44,44	0
55	MG	1a	3063	1/1	0.96	0.11	42,42,42,42	0
55	MG	2A	3188	1/1	0.96	0.08	30,30,30,30	0
55	MG	1A	3923	1/1	0.96	0.23	36,36,36,36	0
55	MG	2A	3044	1/1	0.96	1.07	33,33,33,33	0
55	MG	2A	3532	1/1	0.96	0.10	24,24,24,24	0
55	MG	1A	3605	1/1	0.96	0.29	53,53,53,53	0
55	MG	2a	1754	1/1	0.96	0.13	45,45,45,45	0
55	MG	1D	307	1/1	0.96	0.14	10,10,10,10	0
55	MG	1A	3424	1/1	0.96	0.10	21,21,21,21	0
55	MG	1a	3038	1/1	0.96	0.13	54,54,54,54	0
55	MG	1A	3987	1/1	0.96	0.23	34,34,34,34	0
55	MG	1A	4016	1/1	0.96	0.24	54,54,54,54	0
55	MG	23	101	1/1	0.96	0.45	21,21,21,21	0
55	MG	1A	3275	1/1	0.96	0.54	21,21,21,21	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3899	1/1	0.96	0.13	22,22,22,22	0
55	MG	23	103	1/1	0.96	0.23	17,17,17,17	0
55	MG	2a	1809	1/1	0.96	0.13	26,26,26,26	0
55	MG	2A	3046	1/1	0.96	0.13	56,56,56,56	0
55	MG	2A	3399	1/1	0.96	0.28	32,32,32,32	0
55	MG	1A	3951	1/1	0.96	0.16	33,33,33,33	0
55	MG	2A	3844	1/1	0.96	0.20	40,40,40,40	0
55	MG	2A	3638	1/1	0.96	0.09	26,26,26,26	0
55	MG	2A	3082	1/1	0.96	0.22	44,44,44,44	0
55	MG	2F	302	1/1	0.96	0.24	32,32,32,32	0
55	MG	1A	3432	1/1	0.96	0.10	20,20,20,20	0
55	MG	1A	3291	1/1	0.96	0.15	18,18,18,18	0
55	MG	1A	3152	1/1	0.96	0.12	29,29,29,29	0
55	MG	2A	3664	1/1	0.96	0.32	34,34,34,34	0
55	MG	1A	3503	1/1	0.96	0.24	40,40,40,40	0
55	MG	1A	3219	1/1	0.96	0.20	36,36,36,36	0
55	MG	2A	3656	1/1	0.96	0.09	35,35,35,35	0
55	MG	2A	3845	1/1	0.96	0.13	38,38,38,38	0
55	MG	1A	3768	1/1	0.96	0.59	24,24,24,24	0
55	MG	2A	3581	1/1	0.96	0.11	33,33,33,33	0
55	MG	2A	3654	1/1	0.96	0.31	41,41,41,41	0
55	MG	1R	204	1/1	0.96	0.26	41,41,41,41	0
55	MG	1A	3685	1/1	0.96	0.24	26,26,26,26	0
55	MG	2a	1713	1/1	0.96	0.10	31,31,31,31	0
55	MG	2a	1620	1/1	0.96	0.25	63,63,63,63	0
55	MG	1A	3651	1/1	0.96	0.14	15,15,15,15	0
55	MG	2A	3303	1/1	0.96	0.24	44,44,44,44	0
55	MG	1a	3091	1/1	0.96	0.17	53,53,53,53	0
55	MG	2a	1752	1/1	0.96	0.17	37,37,37,37	0
55	MG	1B	201	1/1	0.96	0.14	23,23,23,23	0
55	MG	1a	3123	1/1	0.96	0.13	27,27,27,27	0
55	MG	1a	3106	1/1	0.96	0.10	10,10,10,10	0
55	MG	1A	4019	1/1	0.96	0.20	41,41,41,41	0
55	MG	1A	3757	1/1	0.96	0.33	23,23,23,23	0
55	MG	2A	3073	1/1	0.96	0.12	30,30,30,30	0
55	MG	1A	3792	1/1	0.96	0.08	37,37,37,37	0
55	MG	1A	3340	1/1	0.96	0.16	39,39,39,39	0
55	MG	1A	3647	1/1	0.96	0.13	43,43,43,43	0
55	MG	2A	3599	1/1	0.96	0.14	40,40,40,40	0
55	MG	1A	3814	1/1	0.96	0.09	25,25,25,25	0
55	MG	2A	3615	1/1	0.96	0.15	25,25,25,25	0
55	MG	16	101	1/1	0.96	0.12	28,28,28,28	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1a	3111	1/1	0.96	0.12	25,25,25,25	0
55	MG	2A	3510	1/1	0.96	0.20	32,32,32,32	0
55	MG	1A	3156	1/1	0.96	0.13	24,24,24,24	0
55	MG	1A	3378	1/1	0.96	0.09	29,29,29,29	0
55	MG	1A	3375	1/1	0.96	0.79	29,29,29,29	0
55	MG	1A	3277	1/1	0.96	0.19	25,25,25,25	0
55	MG	1A	3116	1/1	0.96	0.17	29,29,29,29	0
55	MG	1A	3892	1/1	0.96	0.08	25,25,25,25	0
55	MG	1A	3474	1/1	0.96	0.38	23,23,23,23	0
55	MG	2a	1726	1/1	0.96	0.20	26,26,26,26	0
55	MG	1A	3858	1/1	0.96	0.12	41,41,41,41	0
55	MG	1A	3363	1/1	0.96	0.21	33,33,33,33	0
55	MG	19	102	1/1	0.96	0.12	31,31,31,31	0
55	MG	2A	3397	1/1	0.96	0.11	21,21,21,21	0
55	MG	1a	3150	1/1	0.96	0.09	41,41,41,41	0
55	MG	1A	3034	1/1	0.96	0.23	22,22,22,22	0
55	MG	2A	3050	1/1	0.96	0.12	31,31,31,31	0
55	MG	2r	101	1/1	0.96	0.18	40,40,40,40	0
55	MG	2A	3623	1/1	0.96	0.11	14,14,14,14	0
55	MG	1A	3084	1/1	0.96	0.09	32,32,32,32	0
55	MG	1a	3074	1/1	0.96	0.56	42,42,42,42	0
55	MG	2A	3742	1/1	0.96	0.32	50,50,50,50	0
55	MG	2A	3735	1/1	0.96	0.14	22,22,22,22	0
55	MG	1A	3019	1/1	0.96	0.44	30,30,30,30	0
55	MG	1Z	301	1/1	0.96	0.10	23,23,23,23	0
55	MG	2A	3743	1/1	0.96	0.12	35,35,35,35	0
55	MG	2A	3648	1/1	0.96	0.07	32,32,32,32	0
55	MG	2A	3454	1/1	0.96	0.15	28,28,28,28	0
55	MG	2A	3333	1/1	0.96	0.09	34,34,34,34	0
55	MG	1A	3266	1/1	0.96	0.97	52,52,52,52	0
55	MG	1A	3776	1/1	0.96	0.12	32,32,32,32	0
55	MG	1a	3156	1/1	0.96	0.17	39,39,39,39	0
55	MG	12	102	1/1	0.96	0.37	32,32,32,32	0
55	MG	1A	3136	1/1	0.96	0.54	30,30,30,30	0
55	MG	1a	3159	1/1	0.96	0.07	34,34,34,34	0
55	MG	1a	3019	1/1	0.96	0.14	50,50,50,50	0
55	MG	1A	3138	1/1	0.96	0.65	29,29,29,29	0
55	MG	1A	3107	1/1	0.96	0.08	14,14,14,14	0
55	MG	1A	3628	1/1	0.96	0.10	22,22,22,22	0
55	MG	1A	3534	1/1	0.96	0.64	37,37,37,37	0
55	MG	2a	1695	1/1	0.96	0.12	43,43,43,43	0
55	MG	1P	201	1/1	0.96	0.47	15,15,15,15	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3041	1/1	0.96	0.20	30,30,30,30	0
55	MG	1A	3134	1/1	0.96	0.99	34,34,34,34	0
55	MG	1A	3486	1/1	0.96	0.13	38,38,38,38	0
55	MG	2A	3106	1/1	0.96	0.12	23,23,23,23	0
55	MG	17	104	1/1	0.96	0.45	28,28,28,28	0
55	MG	1A	3925	1/1	0.96	0.13	20,20,20,20	0
55	MG	1B	232	1/1	0.96	0.16	59,59,59,59	0
55	MG	2A	3259	1/1	0.96	0.63	40,40,40,40	0
55	MG	2A	3476	1/1	0.96	0.23	24,24,24,24	0
55	MG	2A	3389	1/1	0.96	0.12	28,28,28,28	0
55	MG	2A	3473	1/1	0.96	0.19	19,19,19,19	0
55	MG	1a	3110	1/1	0.96	0.12	40,40,40,40	0
55	MG	1A	3552	1/1	0.96	0.07	21,21,21,21	0
55	MG	1A	3896	1/1	0.96	0.13	28,28,28,28	0
55	MG	1F	311	1/1	0.96	0.16	29,29,29,29	0
55	MG	1A	3453	1/1	0.96	0.22	24,24,24,24	0
55	MG	1A	3386	1/1	0.96	0.17	10,10,10,10	0
55	MG	1A	3799	1/1	0.96	0.66	51,51,51,51	0
55	MG	2a	1745	1/1	0.96	0.08	53,53,53,53	0
55	MG	1a	3120	1/1	0.96	0.15	42,42,42,42	0
55	MG	1A	3214	1/1	0.96	0.17	23,23,23,23	0
55	MG	2T	201	1/1	0.96	0.11	36,36,36,36	0
55	MG	2n	101	1/1	0.96	0.11	56,56,56,56	0
55	MG	2A	3813	1/1	0.96	0.13	34,34,34,34	0
55	MG	2A	3748	1/1	0.96	0.12	44,44,44,44	0
55	MG	1A	3283	1/1	0.96	0.11	12,12,12,12	0
55	MG	2A	3270	1/1	0.96	0.17	25,25,25,25	0
55	MG	1a	3143	1/1	0.96	0.10	34,34,34,34	0
55	MG	2A	3013	1/1	0.96	0.47	36,36,36,36	0
55	MG	1A	3742	1/1	0.96	0.10	22,22,22,22	0
55	MG	1A	3770	1/1	0.96	0.13	32,32,32,32	0
55	MG	2a	1737	1/1	0.96	0.17	28,28,28,28	0
55	MG	2x	106	1/1	0.96	0.14	60,60,60,60	0
55	MG	2A	3032	1/1	0.96	0.11	28,28,28,28	0
55	MG	2A	3043	1/1	0.96	0.10	20,20,20,20	0
55	MG	2A	3182	1/1	0.96	0.33	25,25,25,25	0
55	MG	1A	3390	1/1	0.96	0.12	30,30,30,30	0
55	MG	2A	3734	1/1	0.96	0.34	39,39,39,39	0
55	MG	1A	3389	1/1	0.96	0.11	21,21,21,21	0
55	MG	2A	3274	1/1	0.96	0.42	46,46,46,46	0
55	MG	1A	3220	1/1	0.96	0.22	34,34,34,34	0
55	MG	1a	3006	1/1	0.96	0.29	56,56,56,56	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3846	1/1	0.96	0.43	43,43,43,43	0
55	MG	1A	3956	1/1	0.96	0.25	39,39,39,39	0
55	MG	1F	304	1/1	0.96	0.08	30,30,30,30	0
55	MG	2a	1627	1/1	0.96	0.17	27,27,27,27	0
55	MG	2D	306	1/1	0.96	0.58	28,28,28,28	0
55	MG	1A	3777	1/1	0.96	0.04	51,51,51,51	0
55	MG	1A	3023	1/1	0.96	0.19	24,24,24,24	0
55	MG	2A	3358	1/1	0.96	0.15	17,17,17,17	0
55	MG	1A	3209	1/1	0.96	0.07	24,24,24,24	0
55	MG	1A	3149	1/1	0.96	0.28	23,23,23,23	0
55	MG	1A	3668	1/1	0.96	0.19	10,10,10,10	0
55	MG	2a	1663	1/1	0.96	0.15	31,31,31,31	0
55	MG	1A	3646	1/1	0.96	0.12	16,16,16,16	0
55	MG	1A	3673	1/1	0.96	0.41	39,39,39,39	0
55	MG	1A	3469	1/1	0.96	0.12	25,25,25,25	0
55	MG	1a	3178	1/1	0.96	0.14	36,36,36,36	0
55	MG	2A	3788	1/1	0.96	0.10	40,40,40,40	0
55	MG	2A	3807	1/1	0.96	0.17	21,21,21,21	0
55	MG	2A	3117	1/1	0.96	0.31	33,33,33,33	0
55	MG	2A	3715	1/1	0.96	0.29	46,46,46,46	0
55	MG	1a	3164	1/1	0.96	0.15	52,52,52,52	0
55	MG	1A	3657	1/1	0.96	0.10	19,19,19,19	0
55	MG	2A	3698	1/1	0.96	0.29	34,34,34,34	0
55	MG	1A	3759	1/1	0.96	0.14	20,20,20,20	0
55	MG	1A	3730	1/1	0.96	0.14	37,37,37,37	0
55	MG	1A	3090	1/1	0.96	0.10	42,42,42,42	0
55	MG	1A	4008	1/1	0.96	0.21	24,24,24,24	0
55	MG	2A	3228	1/1	0.96	0.18	45,45,45,45	0
55	MG	1A	3002	1/1	0.96	0.22	39,39,39,39	0
55	MG	1A	3357	1/1	0.96	0.10	39,39,39,39	0
55	MG	2A	3796	1/1	0.96	0.15	22,22,22,22	0
55	MG	2A	3084	1/1	0.96	0.17	28,28,28,28	0
55	MG	1A	3565	1/1	0.96	0.16	16,16,16,16	0
55	MG	2A	3652	1/1	0.96	0.30	31,31,31,31	0
55	MG	2A	3375	1/1	0.96	0.14	35,35,35,35	0
55	MG	2A	3369	1/1	0.96	0.09	19,19,19,19	0
55	MG	2A	3708	1/1	0.96	0.07	20,20,20,20	0
55	MG	1A	3028	1/1	0.96	0.24	39,39,39,39	0
55	MG	1A	3043	1/1	0.96	0.19	25,25,25,25	0
55	MG	1A	3879	1/1	0.96	0.22	41,41,41,41	0
55	MG	1A	3995	1/1	0.96	0.59	48,48,48,48	0
55	MG	1A	3632	1/1	0.96	0.10	27,27,27,27	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3597	1/1	0.96	0.14	26,26,26,26	0
55	MG	2A	3212	1/1	0.96	0.21	26,26,26,26	0
55	MG	2A	3226	1/1	0.96	0.14	32,32,32,32	0
55	MG	1A	3444	1/1	0.96	0.12	18,18,18,18	0
55	MG	1A	3410	1/1	0.96	0.24	34,34,34,34	0
55	MG	2a	1779	1/1	0.96	0.11	19,19,19,19	0
55	MG	1A	3900	1/1	0.96	0.09	22,22,22,22	0
55	MG	2A	3120	1/1	0.96	0.17	30,30,30,30	0
55	MG	1A	3185	1/1	0.96	0.10	23,23,23,23	0
55	MG	2A	3063	1/1	0.96	0.10	26,26,26,26	0
55	MG	2A	3555	1/1	0.96	0.10	32,32,32,32	0
55	MG	1A	3993	1/1	0.96	0.20	10,10,10,10	0
55	MG	1A	3713	1/1	0.96	0.15	40,40,40,40	0
55	MG	1A	4029	1/1	0.96	0.19	36,36,36,36	0
55	MG	1A	3633	1/1	0.96	0.14	26,26,26,26	0
55	MG	2A	3395	1/1	0.96	0.67	38,38,38,38	0
55	MG	1Q	204	1/1	0.96	0.14	35,35,35,35	0
55	MG	2A	3535	1/1	0.96	0.15	24,24,24,24	0
55	MG	2A	3520	1/1	0.96	0.22	48,48,48,48	0
55	MG	2A	3670	1/1	0.96	0.11	43,43,43,43	0
57	ZN	1n	104	1/1	0.96	0.11	68,68,68,68	0
55	MG	1A	3745	1/1	0.96	0.48	32,32,32,32	0
55	MG	2a	1808	1/1	0.96	0.06	56,56,56,56	0
55	MG	1A	3873	1/1	0.96	0.06	50,50,50,50	0
55	MG	1E	304	1/1	0.96	0.08	24,24,24,24	0
55	MG	1x	109	1/1	0.96	0.07	53,53,53,53	0
55	MG	1W	203	1/1	0.96	0.13	13,13,13,13	0
55	MG	2A	3019	1/1	0.96	0.36	47,47,47,47	0
55	MG	2A	3516	1/1	0.96	0.07	36,36,36,36	0
55	MG	1A	3653	1/1	0.96	0.08	21,21,21,21	0
55	MG	1A	3210	1/1	0.96	0.16	25,25,25,25	0
55	MG	1A	3420	1/1	0.96	0.18	29,29,29,29	0
55	MG	1A	3788	1/1	0.96	0.07	35,35,35,35	0
55	MG	1A	3298	1/1	0.96	0.13	38,38,38,38	0
55	MG	2a	1750	1/1	0.96	0.10	32,32,32,32	0
55	MG	2A	3789	1/1	0.96	0.17	34,34,34,34	0
55	MG	1a	3134	1/1	0.96	0.12	28,28,28,28	0
55	MG	1a	3180	1/1	0.96	0.08	55,55,55,55	0
55	MG	1a	3039	1/1	0.96	0.08	20,20,20,20	0
55	MG	1A	3494	1/1	0.96	0.15	10,10,10,10	0
55	MG	1A	3997	1/1	0.96	0.18	37,37,37,37	0
55	MG	2a	1633	1/1	0.96	0.12	51,51,51,51	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3260	1/1	0.96	0.14	35,35,35,35	0
55	MG	2A	3837	1/1	0.96	0.18	23,23,23,23	0
55	MG	1A	3489	1/1	0.96	0.11	28,28,28,28	0
55	MG	2A	3439	1/1	0.96	0.12	15,15,15,15	0
55	MG	2A	3099	1/1	0.96	0.13	26,26,26,26	0
55	MG	1R	202	1/1	0.96	0.08	32,32,32,32	0
55	MG	1A	3535	1/1	0.96	0.64	22,22,22,22	0
55	MG	1A	3901	1/1	0.96	0.25	28,28,28,28	0
55	MG	1A	3727	1/1	0.96	0.16	39,39,39,39	0
55	MG	1A	3895	1/1	0.96	0.15	48,48,48,48	0
55	MG	1A	3365	1/1	0.96	0.20	41,41,41,41	0
55	MG	2A	3451	1/1	0.96	0.12	19,19,19,19	0
55	MG	1a	3018	1/1	0.96	0.25	42,42,42,42	0
55	MG	2A	3746	1/1	0.96	0.36	31,31,31,31	0
55	MG	2A	3049	1/1	0.96	0.09	21,21,21,21	0
55	MG	2A	3537	1/1	0.96	0.11	22,22,22,22	0
55	MG	2a	1760	1/1	0.96	0.17	38,38,38,38	0
55	MG	1A	3241	1/1	0.96	0.28	34,34,34,34	0
55	MG	2A	3163	1/1	0.96	0.21	33,33,33,33	0
55	MG	1A	3462	1/1	0.96	0.77	30,30,30,30	0
55	MG	1l	103	1/1	0.96	0.16	47,47,47,47	0
55	MG	1A	3609	1/1	0.97	0.08	17,17,17,17	0
55	MG	2A	3026	1/1	0.97	0.25	27,27,27,27	0
55	MG	1l	203	1/1	0.97	0.08	35,35,35,35	0
55	MG	2A	3692	1/1	0.97	0.18	48,48,48,48	0
55	MG	2A	3479	1/1	0.97	0.15	23,23,23,23	0
55	MG	1A	3482	1/1	0.97	0.34	28,28,28,28	0
55	MG	2A	3726	1/1	0.97	0.08	52,52,52,52	0
55	MG	2A	3463	1/1	0.97	0.19	16,16,16,16	0
55	MG	1a	3201	1/1	0.97	0.09	33,33,33,33	0
55	MG	1A	3276	1/1	0.97	0.14	40,40,40,40	0
55	MG	1a	3136	1/1	0.97	0.06	32,32,32,32	0
55	MG	1a	3167	1/1	0.97	0.28	31,31,31,31	0
55	MG	2A	3448	1/1	0.97	0.10	32,32,32,32	0
55	MG	1A	3769	1/1	0.97	0.10	48,48,48,48	0
55	MG	1a	3192	1/1	0.97	0.10	33,33,33,33	0
55	MG	1A	3088	1/1	0.97	0.20	28,28,28,28	0
55	MG	1A	3845	1/1	0.97	0.12	19,19,19,19	0
55	MG	1A	3082	1/1	0.97	0.17	36,36,36,36	0
55	MG	1A	3263	1/1	0.97	0.14	15,15,15,15	0
55	MG	2A	3481	1/1	0.97	1.08	39,39,39,39	0
55	MG	2A	3745	1/1	0.97	0.18	17,17,17,17	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3677	1/1	0.97	0.27	36,36,36,36	0
55	MG	1A	3054	1/1	0.97	0.18	21,21,21,21	0
55	MG	1A	3963	1/1	0.97	0.07	42,42,42,42	0
55	MG	1I	201	1/1	0.97	0.07	52,52,52,52	0
55	MG	1A	3664	1/1	0.97	0.14	20,20,20,20	0
55	MG	1a	3086	1/1	0.97	0.48	50,50,50,50	0
55	MG	2a	1644	1/1	0.97	0.19	31,31,31,31	0
55	MG	1A	3832	1/1	0.97	0.16	17,17,17,17	0
55	MG	1A	3117	1/1	0.97	0.09	24,24,24,24	0
55	MG	2A	3432	1/1	0.97	0.19	45,45,45,45	0
55	MG	2A	3712	1/1	0.97	0.11	22,22,22,22	0
55	MG	1A	3382	1/1	0.97	0.46	26,26,26,26	0
55	MG	2A	3413	1/1	0.97	0.08	30,30,30,30	0
55	MG	1a	3010	1/1	0.97	0.16	55,55,55,55	0
55	MG	1A	3547	1/1	0.97	0.22	24,24,24,24	0
55	MG	1A	3607	1/1	0.97	0.12	15,15,15,15	0
55	MG	2A	3474	1/1	0.97	0.20	10,10,10,10	0
55	MG	1A	3110	1/1	0.97	0.81	43,43,43,43	0
55	MG	2A	3561	1/1	0.97	0.15	27,27,27,27	0
55	MG	2A	3565	1/1	0.97	0.15	22,22,22,22	0
55	MG	1A	3966	1/1	0.97	0.16	28,28,28,28	0
55	MG	1A	3708	1/1	0.97	0.07	26,26,26,26	0
55	MG	1A	3254	1/1	0.97	0.23	27,27,27,27	0
55	MG	1A	4028	1/1	0.97	0.13	17,17,17,17	0
55	MG	1S	202	1/1	0.97	0.12	33,33,33,33	0
55	MG	1A	3818	1/1	0.97	0.14	28,28,28,28	0
55	MG	1E	302	1/1	0.97	0.28	32,32,32,32	0
55	MG	1A	3431	1/1	0.97	0.26	14,14,14,14	0
55	MG	1A	3163	1/1	0.97	0.14	19,19,19,19	0
55	MG	1A	3174	1/1	0.97	0.14	23,23,23,23	0
55	MG	1A	3971	1/1	0.97	0.14	20,20,20,20	0
55	MG	2A	3007	1/1	0.97	0.08	38,38,38,38	0
55	MG	1A	3289	1/1	0.97	0.25	10,10,10,10	0
55	MG	2A	3469	1/1	0.97	0.17	21,21,21,21	0
55	MG	1a	3077	1/1	0.97	0.11	46,46,46,46	0
55	MG	2a	1658	1/1	0.97	0.08	20,20,20,20	0
55	MG	2A	3281	1/1	0.97	0.28	40,40,40,40	0
55	MG	1A	3124	1/1	0.97	0.18	27,27,27,27	0
55	MG	1A	3576	1/1	0.97	0.13	10,10,10,10	0
55	MG	1A	3064	1/1	0.97	0.46	13,13,13,13	0
55	MG	2a	1769	1/1	0.97	0.11	64,64,64,64	0
55	MG	1A	3598	1/1	0.97	0.29	35,35,35,35	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3650	1/1	0.97	0.27	41,41,41,41	0
55	MG	1a	3099	1/1	0.97	0.13	18,18,18,18	0
55	MG	1a	3045	1/1	0.97	0.07	30,30,30,30	0
55	MG	2A	3031	1/1	0.97	0.38	46,46,46,46	0
55	MG	1x	101	1/1	0.97	0.19	38,38,38,38	0
55	MG	2q	201	1/1	0.97	0.09	30,30,30,30	0
55	MG	2A	3100	1/1	0.97	0.16	22,22,22,22	0
55	MG	1A	3794	1/1	0.97	0.07	35,35,35,35	0
55	MG	1a	3140	1/1	0.97	0.23	42,42,42,42	0
55	MG	1A	3973	1/1	0.97	0.12	38,38,38,38	0
55	MG	2U	201	1/1	0.97	1.21	48,48,48,48	0
55	MG	1A	3575	1/1	0.97	0.08	27,27,27,27	0
55	MG	2A	3047	1/1	0.97	0.08	33,33,33,33	0
55	MG	2A	3834	1/1	0.97	0.11	43,43,43,43	0
55	MG	1a	3195	1/1	0.97	0.06	25,25,25,25	0
55	MG	1A	3954	1/1	0.97	0.10	38,38,38,38	0
55	MG	1A	3445	1/1	0.97	0.08	19,19,19,19	0
55	MG	1A	4026	1/1	0.97	0.12	26,26,26,26	0
55	MG	1a	3147	1/1	0.97	0.10	44,44,44,44	0
55	MG	2A	3637	1/1	0.97	0.09	31,31,31,31	0
55	MG	1A	3560	1/1	0.97	0.17	19,19,19,19	0
55	MG	1A	3256	1/1	0.97	0.27	25,25,25,25	0
55	MG	1A	3571	1/1	0.97	0.08	25,25,25,25	0
55	MG	1a	3207	1/1	0.97	0.29	50,50,50,50	0
57	ZN	29	501	1/1	0.97	0.09	60,60,60,60	0
55	MG	2A	3548	1/1	0.97	0.29	22,22,22,22	0
55	MG	1A	4009	1/1	0.97	0.08	41,41,41,41	0
55	MG	1A	3941	1/1	0.97	0.12	15,15,15,15	0
55	MG	1A	3978	1/1	0.97	0.11	21,21,21,21	0
55	MG	1A	3114	1/1	0.97	0.38	10,10,10,10	0
55	MG	1A	3487	1/1	0.97	0.14	25,25,25,25	0
57	ZN	16	104	1/1	0.97	0.19	36,36,36,36	0
55	MG	1B	209	1/1	0.97	0.12	27,27,27,27	0
55	MG	2A	3401	1/1	0.97	0.17	38,38,38,38	0
55	MG	2A	3287	1/1	0.97	0.44	46,46,46,46	0
55	MG	1A	3126	1/1	0.97	0.12	48,48,48,48	0
55	MG	1A	3513	1/1	0.97	0.18	53,53,53,53	0
55	MG	2A	3238	1/1	0.97	0.60	18,18,18,18	0
55	MG	1A	3554	1/1	0.97	0.15	18,18,18,18	0
55	MG	2U	202	1/1	0.97	0.08	23,23,23,23	0
55	MG	2A	3628	1/1	0.97	0.14	31,31,31,31	0
55	MG	2A	3665	1/1	0.97	0.50	37,37,37,37	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3689	1/1	0.97	0.17	54,54,54,54	0
55	MG	1A	3859	1/1	0.97	0.06	51,51,51,51	0
55	MG	1A	3122	1/1	0.97	0.37	22,22,22,22	0
55	MG	1A	3669	1/1	0.97	0.10	26,26,26,26	0
55	MG	2a	1619	1/1	0.97	0.47	27,27,27,27	0
55	MG	1A	3563	1/1	0.97	0.12	19,19,19,19	0
55	MG	1A	3076	1/1	0.97	0.26	42,42,42,42	0
55	MG	1A	3703	1/1	0.97	0.20	25,25,25,25	0
55	MG	13	105	1/1	0.97	0.27	26,26,26,26	0
55	MG	21	103	1/1	0.97	0.16	18,18,18,18	0
55	MG	1Z	303	1/1	0.97	0.15	28,28,28,28	0
55	MG	1A	3049	1/1	0.97	0.24	26,26,26,26	0
55	MG	2A	3739	1/1	0.97	0.11	43,43,43,43	0
55	MG	2A	3579	1/1	0.97	0.20	25,25,25,25	0
55	MG	2A	3830	1/1	0.97	0.11	31,31,31,31	0
55	MG	1A	3548	1/1	0.97	0.15	23,23,23,23	0
55	MG	1F	309	1/1	0.97	0.14	45,45,45,45	0
55	MG	1A	3700	1/1	0.97	0.08	20,20,20,20	0
55	MG	2A	3525	1/1	0.97	0.19	29,29,29,29	0
55	MG	2A	3285	1/1	0.97	0.12	21,21,21,21	0
55	MG	2A	3386	1/1	0.97	0.15	31,31,31,31	0
55	MG	2A	3682	1/1	0.97	0.11	61,61,61,61	0
55	MG	1A	3663	1/1	0.97	0.20	44,44,44,44	0
55	MG	2a	1664	1/1	0.97	0.14	37,37,37,37	0
55	MG	2R	201	1/1	0.97	0.16	17,17,17,17	0
55	MG	2A	3784	1/1	0.97	0.11	43,43,43,43	0
55	MG	2A	3792	1/1	0.97	0.07	30,30,30,30	0
55	MG	2A	3647	1/1	0.97	0.13	41,41,41,41	0
55	MG	1a	3022	1/1	0.97	0.17	16,16,16,16	0
55	MG	2a	1686	1/1	0.97	0.23	37,37,37,37	0
55	MG	1Q	202	1/1	0.97	0.32	30,30,30,30	0
55	MG	18	104	1/1	0.97	0.07	33,33,33,33	0
55	MG	2A	3627	1/1	0.97	0.20	23,23,23,23	0
55	MG	1v	104	1/1	0.97	0.10	35,35,35,35	0
55	MG	1A	3599	1/1	0.97	0.11	27,27,27,27	0
55	MG	1A	3795	1/1	0.97	0.20	27,27,27,27	0
55	MG	1A	3573	1/1	0.97	0.14	24,24,24,24	0
55	MG	1A	3045	1/1	0.97	0.10	15,15,15,15	0
55	MG	1a	3158	1/1	0.97	0.12	19,19,19,19	0
55	MG	1A	3916	1/1	0.97	0.10	21,21,21,21	0
55	MG	2A	3747	1/1	0.97	0.16	49,49,49,49	0
55	MG	2a	1635	1/1	0.97	0.12	31,31,31,31	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3221	1/1	0.97	0.12	33,33,33,33	0
55	MG	1A	3782	1/1	0.97	0.09	29,29,29,29	0
55	MG	18	101	1/1	0.97	0.58	27,27,27,27	0
55	MG	1A	3936	1/1	0.97	0.18	39,39,39,39	0
55	MG	1a	3182	1/1	0.97	0.14	39,39,39,39	0
55	MG	1A	3603	1/1	0.97	0.07	36,36,36,36	0
55	MG	1A	3203	1/1	0.97	0.64	14,14,14,14	0
55	MG	2A	3452	1/1	0.97	0.28	37,37,37,37	0
55	MG	1A	3574	1/1	0.97	0.25	46,46,46,46	0
55	MG	1A	3672	1/1	0.97	0.13	32,32,32,32	0
55	MG	1a	3053	1/1	0.97	0.19	38,38,38,38	0
55	MG	1A	3578	1/1	0.97	0.09	26,26,26,26	0
55	MG	1a	3163	1/1	0.97	0.14	42,42,42,42	0
55	MG	1a	3188	1/1	0.97	0.07	41,41,41,41	0
55	MG	2A	3153	1/1	0.97	0.18	50,50,50,50	0
55	MG	2A	3311	1/1	0.97	0.20	33,33,33,33	0
55	MG	2A	3477	1/1	0.97	0.12	16,16,16,16	0
55	MG	1A	3543	1/1	0.97	0.24	30,30,30,30	0
55	MG	2A	3517	1/1	0.97	0.37	44,44,44,44	0
55	MG	2A	3604	1/1	0.97	0.21	34,34,34,34	0
55	MG	1a	3173	1/1	0.97	0.11	51,51,51,51	0
55	MG	1A	3660	1/1	0.97	0.13	10,10,10,10	0
55	MG	2A	3677	1/1	0.97	0.09	37,37,37,37	0
55	MG	1A	3559	1/1	0.97	0.15	42,42,42,42	0
55	MG	2A	3687	1/1	0.97	0.19	49,49,49,49	0
55	MG	2x	105	1/1	0.97	0.22	31,31,31,31	0
55	MG	1A	3318	1/1	0.97	0.32	27,27,27,27	0
55	MG	2A	3806	1/1	0.97	0.16	36,36,36,36	0
55	MG	2A	3229	1/1	0.97	0.12	22,22,22,22	0
55	MG	1A	3970	1/1	0.97	0.10	22,22,22,22	0
55	MG	1X	103	1/1	0.97	0.14	34,34,34,34	0
55	MG	1A	3259	1/1	0.97	0.08	10,10,10,10	0
55	MG	1A	3425	1/1	0.97	0.10	33,33,33,33	0
55	MG	1A	3409	1/1	0.97	0.08	50,50,50,50	0
55	MG	1a	3130	1/1	0.97	0.27	38,38,38,38	0
55	MG	1A	3649	1/1	0.97	0.18	31,31,31,31	0
55	MG	1A	3725	1/1	0.97	0.11	26,26,26,26	0
55	MG	1A	3710	1/1	0.97	0.06	13,13,13,13	0
55	MG	11	101	1/1	0.97	0.52	32,32,32,32	0
55	MG	1A	3221	1/1	0.97	0.62	30,30,30,30	0
55	MG	1A	3617	1/1	0.97	0.11	27,27,27,27	0
55	MG	1A	3906	1/1	0.97	0.18	21,21,21,21	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3325	1/1	0.97	0.43	41,41,41,41	0
55	MG	1a	3011	1/1	0.97	0.09	70,70,70,70	0
55	MG	18	102	1/1	0.97	0.56	26,26,26,26	0
55	MG	1A	3515	1/1	0.97	0.19	10,10,10,10	0
55	MG	2A	3816	1/1	0.97	0.19	25,25,25,25	0
55	MG	2A	3723	1/1	0.97	0.12	36,36,36,36	0
55	MG	1A	3667	1/1	0.97	0.15	18,18,18,18	0
55	MG	1A	3038	1/1	0.97	0.28	21,21,21,21	0
55	MG	1A	3838	1/1	0.97	0.14	26,26,26,26	0
55	MG	2A	3710	1/1	0.97	0.30	27,27,27,27	0
55	MG	2A	3394	1/1	0.97	0.14	27,27,27,27	0
55	MG	1A	3829	1/1	0.97	0.17	28,28,28,28	0
55	MG	1A	3267	1/1	0.97	0.36	37,37,37,37	0
55	MG	2Q	204	1/1	0.97	0.28	33,33,33,33	0
55	MG	1v	101	1/1	0.97	0.16	36,36,36,36	0
55	MG	2A	3567	1/1	0.97	0.15	33,33,33,33	0
55	MG	1A	3429	1/1	0.97	0.09	23,23,23,23	0
55	MG	1A	3135	1/1	0.97	0.13	23,23,23,23	0
55	MG	1P	202	1/1	0.97	0.57	15,15,15,15	0
55	MG	1A	3648	1/1	0.97	0.23	34,34,34,34	0
55	MG	1A	3238	1/1	0.97	0.09	27,27,27,27	0
55	MG	2A	3475	1/1	0.97	0.14	17,17,17,17	0
55	MG	1B	212	1/1	0.97	0.15	25,25,25,25	0
55	MG	2A	3667	1/1	0.97	0.09	32,32,32,32	0
55	MG	1A	3902	1/1	0.97	0.09	25,25,25,25	0
55	MG	1A	3396	1/1	0.97	0.19	22,22,22,22	0
55	MG	1A	3592	1/1	0.97	0.09	27,27,27,27	0
55	MG	1B	207	1/1	0.97	0.13	42,42,42,42	0
55	MG	2A	3478	1/1	0.97	0.20	15,15,15,15	0
55	MG	2A	3021	1/1	0.97	0.20	55,55,55,55	0
55	MG	2A	3744	1/1	0.97	0.08	51,51,51,51	0
55	MG	1A	3100	1/1	0.97	0.16	38,38,38,38	0
55	MG	2A	3029	1/1	0.97	0.11	21,21,21,21	0
55	MG	1a	3141	1/1	0.97	0.47	45,45,45,45	0
55	MG	1U	207	1/1	0.97	0.50	31,31,31,31	0
55	MG	1A	3234	1/1	0.97	0.39	28,28,28,28	0
55	MG	1a	3001	1/1	0.97	0.07	47,47,47,47	0
55	MG	1W	209	1/1	0.97	0.21	29,29,29,29	0
55	MG	1A	3187	1/1	0.97	0.80	33,33,33,33	0
55	MG	1n	103	1/1	0.97	0.12	37,37,37,37	0
55	MG	1V	204	1/1	0.97	0.09	51,51,51,51	0
55	MG	2A	3002	1/1	0.97	0.10	26,26,26,26	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3931	1/1	0.97	0.20	20,20,20,20	0
55	MG	1A	3456	1/1	0.97	0.33	24,24,24,24	0
55	MG	2e	202	1/1	0.97	0.16	47,47,47,47	0
55	MG	1A	3525	1/1	0.97	0.12	25,25,25,25	0
55	MG	2A	3783	1/1	0.97	0.12	19,19,19,19	0
55	MG	2A	3383	1/1	0.97	0.20	37,37,37,37	0
55	MG	1A	3917	1/1	0.97	0.12	16,16,16,16	0
55	MG	1U	201	1/1	0.97	0.56	24,24,24,24	0
55	MG	1A	3414	1/1	0.97	0.35	26,26,26,26	0
55	MG	2A	3116	1/1	0.97	0.47	38,38,38,38	0
55	MG	2A	3755	1/1	0.97	0.15	42,42,42,42	0
55	MG	2A	3554	1/1	0.97	0.27	28,28,28,28	0
55	MG	2A	3118	1/1	0.97	0.07	51,51,51,51	0
55	MG	1A	3693	1/1	0.97	0.20	46,46,46,46	0
55	MG	1A	3195	1/1	0.97	0.36	31,31,31,31	0
55	MG	2a	1700	1/1	0.97	0.21	43,43,43,43	0
55	MG	2e	201	1/1	0.97	0.13	43,43,43,43	0
55	MG	1A	3533	1/1	0.97	0.12	21,21,21,21	0
55	MG	1A	3821	1/1	0.97	0.14	10,10,10,10	0
55	MG	2A	3553	1/1	0.97	0.24	45,45,45,45	0
55	MG	1A	3947	1/1	0.97	0.15	25,25,25,25	0
55	MG	1A	3819	1/1	0.97	0.13	22,22,22,22	0
55	MG	1A	3732	1/1	0.97	0.13	18,18,18,18	0
55	MG	1R	205	1/1	0.97	0.06	30,30,30,30	0
55	MG	2A	3069	1/1	0.97	0.13	28,28,28,28	0
55	MG	1A	3194	1/1	0.97	0.66	32,32,32,32	0
55	MG	1A	4011	1/1	0.97	0.26	54,54,54,54	0
55	MG	1A	3779	1/1	0.97	0.19	27,27,27,27	0
55	MG	2A	3588	1/1	0.97	0.08	42,42,42,42	0
55	MG	1x	113	1/1	0.97	0.22	50,50,50,50	0
55	MG	2A	3483	1/1	0.97	0.14	29,29,29,29	0
55	MG	1A	3785	1/1	0.97	0.14	14,14,14,14	0
55	MG	1A	3724	1/1	0.97	0.12	20,20,20,20	0
55	MG	1F	303	1/1	0.97	0.31	29,29,29,29	0
55	MG	2A	3527	1/1	0.97	0.12	40,40,40,40	0
55	MG	1A	3349	1/1	0.97	0.14	26,26,26,26	0
55	MG	2A	3342	1/1	0.97	0.08	21,21,21,21	0
55	MG	2A	3450	1/1	0.97	0.15	19,19,19,19	0
55	MG	1A	3042	1/1	0.97	0.09	18,18,18,18	0
55	MG	1a	3103	1/1	0.97	0.11	44,44,44,44	0
55	MG	2x	107	1/1	0.97	0.11	47,47,47,47	0
55	MG	1A	3037	1/1	0.97	0.28	15,15,15,15	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1x	111	1/1	0.97	0.12	32,32,32,32	0
55	MG	2A	3408	1/1	0.97	0.20	34,34,34,34	0
55	MG	2I	102	1/1	0.97	0.44	40,40,40,40	0
55	MG	1A	3231	1/1	0.97	0.12	22,22,22,22	0
55	MG	2A	3329	1/1	0.97	0.31	23,23,23,23	0
55	MG	1A	3010	1/1	0.97	0.28	33,33,33,33	0
55	MG	1a	3084	1/1	0.97	0.10	10,10,10,10	0
55	MG	1A	3656	1/1	0.97	0.09	36,36,36,36	0
55	MG	1A	3085	1/1	0.97	0.16	14,14,14,14	0
55	MG	1A	3781	1/1	0.97	0.07	41,41,41,41	0
55	MG	1a	3115	1/1	0.97	0.20	38,38,38,38	0
55	MG	1A	3809	1/1	0.97	0.68	32,32,32,32	0
55	MG	1A	3800	1/1	0.97	0.12	24,24,24,24	0
55	MG	1A	3614	1/1	0.97	0.16	14,14,14,14	0
55	MG	2A	3384	1/1	0.97	0.11	24,24,24,24	0
55	MG	2A	3719	1/1	0.97	0.20	47,47,47,47	0
55	MG	1A	3643	1/1	0.97	0.17	35,35,35,35	0
55	MG	2A	3467	1/1	0.97	0.15	34,34,34,34	0
55	MG	1A	3545	1/1	0.97	0.19	22,22,22,22	0
55	MG	2A	3780	1/1	0.97	0.23	51,51,51,51	0
55	MG	1A	3086	1/1	0.97	0.10	37,37,37,37	0
55	MG	1F	312	1/1	0.97	0.09	19,19,19,19	0
55	MG	1O	202	1/1	0.97	0.21	47,47,47,47	0
55	MG	1A	3843	1/1	0.97	0.55	35,35,35,35	0
55	MG	1A	3679	1/1	0.97	0.46	23,23,23,23	0
55	MG	1A	3369	1/1	0.97	0.13	32,32,32,32	0
55	MG	2A	3176	1/1	0.97	0.09	19,19,19,19	0
55	MG	1A	3067	1/1	0.97	0.11	21,21,21,21	0
55	MG	1A	3119	1/1	0.97	0.20	34,34,34,34	0
55	MG	1A	3594	1/1	0.97	0.24	25,25,25,25	0
55	MG	1a	3044	1/1	0.97	0.10	56,56,56,56	0
55	MG	25	103	1/1	0.97	0.16	27,27,27,27	0
55	MG	1A	3484	1/1	0.97	0.46	24,24,24,24	0
55	MG	1A	3756	1/1	0.97	0.11	23,23,23,23	0
55	MG	1A	3711	1/1	0.98	0.09	10,10,10,10	0
55	MG	1a	3138	1/1	0.98	0.12	19,19,19,19	0
55	MG	2a	1711	1/1	0.98	0.12	49,49,49,49	0
55	MG	1A	3856	1/1	0.98	0.46	39,39,39,39	0
55	MG	1a	3008	1/1	0.98	0.12	54,54,54,54	0
55	MG	2a	1683	1/1	0.98	0.11	36,36,36,36	0
55	MG	2A	3427	1/1	0.98	0.65	45,45,45,45	0
55	MG	2A	3490	1/1	0.98	0.20	23,23,23,23	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1a	3194	1/1	0.98	0.07	27,27,27,27	0
55	MG	1A	3674	1/1	0.98	0.12	32,32,32,32	0
55	MG	2A	3417	1/1	0.98	0.13	16,16,16,16	0
55	MG	1a	3092	1/1	0.98	0.15	51,51,51,51	0
55	MG	1A	3636	1/1	0.98	0.07	29,29,29,29	0
55	MG	2A	3083	1/1	0.98	0.35	27,27,27,27	0
55	MG	2A	3020	1/1	0.98	0.23	26,26,26,26	0
55	MG	2A	3633	1/1	0.98	0.23	30,30,30,30	0
55	MG	2A	3798	1/1	0.98	0.20	23,23,23,23	0
55	MG	2A	3551	1/1	0.98	0.10	32,32,32,32	0
55	MG	2A	3271	1/1	0.98	0.12	57,57,57,57	0
55	MG	2A	3685	1/1	0.98	0.11	10,10,10,10	0
55	MG	2a	1696	1/1	0.98	0.11	53,53,53,53	0
55	MG	1A	3131	1/1	0.98	0.27	21,21,21,21	0
55	MG	13	101	1/1	0.98	0.37	30,30,30,30	0
55	MG	1A	3823	1/1	0.98	0.10	16,16,16,16	0
55	MG	2A	3524	1/1	0.98	0.14	23,23,23,23	0
55	MG	1A	3890	1/1	0.98	0.28	43,43,43,43	0
55	MG	2A	3771	1/1	0.98	0.13	26,26,26,26	0
55	MG	1A	3101	1/1	0.98	0.33	34,34,34,34	0
55	MG	2A	3070	1/1	0.98	0.11	34,34,34,34	0
55	MG	1A	3129	1/1	0.98	0.64	24,24,24,24	0
55	MG	2A	3488	1/1	0.98	0.19	16,16,16,16	0
55	MG	2A	3010	1/1	0.98	0.07	42,42,42,42	0
55	MG	1A	4015	1/1	0.98	0.33	19,19,19,19	0
55	MG	1A	3093	1/1	0.98	0.12	18,18,18,18	0
55	MG	1A	3939	1/1	0.98	0.11	14,14,14,14	0
55	MG	2A	3586	1/1	0.98	0.12	31,31,31,31	0
55	MG	2A	3344	1/1	0.98	0.14	37,37,37,37	0
55	MG	2A	3721	1/1	0.98	0.13	40,40,40,40	0
55	MG	1A	3150	1/1	0.98	0.09	27,27,27,27	0
55	MG	17	106	1/1	0.98	0.22	17,17,17,17	0
55	MG	1A	3540	1/1	0.98	0.15	14,14,14,14	0
55	MG	1a	3132	1/1	0.98	0.18	31,31,31,31	0
55	MG	2A	3185	1/1	0.98	0.30	54,54,54,54	0
55	MG	1A	3979	1/1	0.98	0.21	36,36,36,36	0
55	MG	2A	3038	1/1	0.98	0.25	51,51,51,51	0
55	MG	1A	3692	1/1	0.98	0.11	44,44,44,44	0
55	MG	1A	3773	1/1	0.98	0.10	21,21,21,21	0
55	MG	1X	104	1/1	0.98	0.10	22,22,22,22	0
55	MG	1A	3980	1/1	0.98	0.08	24,24,24,24	0
55	MG	1A	3920	1/1	0.98	0.13	15,15,15,15	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2a	1722	1/1	0.98	0.10	26,26,26,26	0
55	MG	1A	3627	1/1	0.98	0.19	10,10,10,10	0
55	MG	1A	3958	1/1	0.98	0.15	27,27,27,27	0
55	MG	1a	3135	1/1	0.98	0.14	26,26,26,26	0
55	MG	1A	3068	1/1	0.98	0.27	34,34,34,34	0
55	MG	17	105	1/1	0.98	0.29	22,22,22,22	0
55	MG	1A	3854	1/1	0.98	0.08	38,38,38,38	0
55	MG	1A	3046	1/1	0.98	0.12	30,30,30,30	0
55	MG	1A	3115	1/1	0.98	0.28	20,20,20,20	0
55	MG	1a	3046	1/1	0.98	0.11	33,33,33,33	0
55	MG	1A	3686	1/1	0.98	0.14	39,39,39,39	0
55	MG	2A	3121	1/1	0.98	0.06	29,29,29,29	0
55	MG	2a	1689	1/1	0.98	0.14	42,42,42,42	0
55	MG	2B	208	1/1	0.98	0.08	22,22,22,22	0
55	MG	2A	3208	1/1	0.98	0.10	37,37,37,37	0
55	MG	1H	202	1/1	0.98	0.12	22,22,22,22	0
55	MG	2A	3129	1/1	0.98	0.22	27,27,27,27	0
55	MG	2a	1677	1/1	0.98	0.06	45,45,45,45	0
55	MG	1A	4013	1/1	0.98	0.13	28,28,28,28	0
55	MG	1A	3937	1/1	0.98	0.08	29,29,29,29	0
55	MG	1A	3654	1/1	0.98	0.11	43,43,43,43	0
55	MG	1E	314	1/1	0.98	0.11	36,36,36,36	0
55	MG	1A	3065	1/1	0.98	0.12	16,16,16,16	0
55	MG	1A	3056	1/1	0.98	0.12	42,42,42,42	0
55	MG	2A	3590	1/1	0.98	0.12	27,27,27,27	0
55	MG	2A	3512	1/1	0.98	0.14	27,27,27,27	0
55	MG	1V	201	1/1	0.98	0.40	27,27,27,27	0
55	MG	2a	1629	1/1	0.98	0.10	30,30,30,30	0
55	MG	2A	3390	1/1	0.98	0.11	37,37,37,37	0
55	MG	1A	3520	1/1	0.98	0.16	11,11,11,11	0
55	MG	2a	1816	1/1	0.98	0.05	35,35,35,35	0
55	MG	2D	308	1/1	0.98	0.24	22,22,22,22	0
55	MG	1A	3047	1/1	0.98	0.17	21,21,21,21	0
55	MG	2D	307	1/1	0.98	0.20	25,25,25,25	0
55	MG	1U	204	1/1	0.98	0.34	17,17,17,17	0
55	MG	2a	1707	1/1	0.98	0.60	43,43,43,43	0
55	MG	1A	3572	1/1	0.98	0.10	36,36,36,36	0
55	MG	1T	201	1/1	0.98	0.32	13,13,13,13	0
55	MG	1B	227	1/1	0.98	0.10	31,31,31,31	0
55	MG	2A	3825	1/1	0.98	0.10	31,31,31,31	0
55	MG	1A	3388	1/1	0.98	0.23	19,19,19,19	0
55	MG	2A	3826	1/1	0.98	0.09	35,35,35,35	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3755	1/1	0.98	0.41	26,26,26,26	0
55	MG	1A	3478	1/1	0.98	0.45	27,27,27,27	0
55	MG	1A	3052	1/1	0.98	0.08	19,19,19,19	0
55	MG	2a	1630	1/1	0.98	0.13	48,48,48,48	0
57	ZN	2n	102	1/1	0.98	0.12	52,52,52,52	0
55	MG	18	106	1/1	0.98	0.11	10,10,10,10	0
55	MG	1A	3500	1/1	0.98	0.35	34,34,34,34	0
55	MG	1a	3146	1/1	0.98	0.16	29,29,29,29	0
55	MG	1A	3798	1/1	0.98	0.11	17,17,17,17	0
55	MG	1a	3175	1/1	0.98	0.07	74,74,74,74	0
55	MG	2A	3360	1/1	0.98	0.06	38,38,38,38	0
55	MG	1A	3095	1/1	0.98	0.51	29,29,29,29	0
55	MG	1A	3278	1/1	0.98	0.28	32,32,32,32	0
55	MG	2A	3492	1/1	0.98	0.15	10,10,10,10	0
55	MG	2a	1803	1/1	0.98	0.08	29,29,29,29	0
55	MG	1A	3527	1/1	0.98	0.14	17,17,17,17	0
55	MG	2A	3800	1/1	0.98	0.12	20,20,20,20	0
55	MG	1A	3111	1/1	0.98	0.22	40,40,40,40	0
55	MG	1A	3499	1/1	0.98	0.15	39,39,39,39	0
55	MG	1E	310	1/1	0.98	0.11	20,20,20,20	0
55	MG	15	101	1/1	0.98	0.67	36,36,36,36	0
55	MG	1W	202	1/1	0.98	0.23	20,20,20,20	0
55	MG	2A	3472	1/1	0.98	0.23	15,15,15,15	0
55	MG	2A	3152	1/1	0.98	0.19	50,50,50,50	0
55	MG	2A	3266	1/1	0.98	0.27	32,32,32,32	0
55	MG	1l	201	1/1	0.98	0.19	24,24,24,24	0
55	MG	1A	3729	1/1	0.98	0.15	29,29,29,29	0
55	MG	21	104	1/1	0.98	0.11	27,27,27,27	0
55	MG	1A	3867	1/1	0.98	0.16	18,18,18,18	0
55	MG	2A	3543	1/1	0.98	0.08	23,23,23,23	0
55	MG	1A	3596	1/1	0.98	0.09	34,34,34,34	0
55	MG	1A	3864	1/1	0.98	0.10	31,31,31,31	0
55	MG	1A	3253	1/1	0.98	0.31	21,21,21,21	0
55	MG	1A	3211	1/1	0.98	0.11	37,37,37,37	0
55	MG	2A	3795	1/1	0.98	0.15	30,30,30,30	0
55	MG	1A	3404	1/1	0.98	0.11	21,21,21,21	0
55	MG	1A	3353	1/1	0.98	0.35	34,34,34,34	0
55	MG	1A	3081	1/1	0.98	0.09	15,15,15,15	0
55	MG	1A	3452	1/1	0.98	0.14	21,21,21,21	0
55	MG	1A	4027	1/1	0.98	0.09	11,11,11,11	0
55	MG	1A	3874	1/1	0.98	0.11	30,30,30,30	0
55	MG	2a	1715	1/1	0.98	0.12	36,36,36,36	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3977	1/1	0.98	0.11	23,23,23,23	0
55	MG	1A	3743	1/1	0.98	0.11	10,10,10,10	0
55	MG	2A	3468	1/1	0.98	0.30	19,19,19,19	0
55	MG	1A	3153	1/1	0.98	0.13	30,30,30,30	0
55	MG	1A	3302	1/1	0.98	0.17	26,26,26,26	0
55	MG	1A	3003	1/1	0.98	0.10	28,28,28,28	0
55	MG	1A	3407	1/1	0.98	0.16	37,37,37,37	0
55	MG	2A	3033	1/1	0.98	0.10	22,22,22,22	0
55	MG	1A	3717	1/1	0.98	0.11	25,25,25,25	0
55	MG	1A	3077	1/1	0.98	0.12	24,24,24,24	0
55	MG	1A	3940	1/1	0.98	0.20	22,22,22,22	0
55	MG	1a	3119	1/1	0.98	0.12	16,16,16,16	0
55	MG	2A	3702	1/1	0.98	0.07	31,31,31,31	0
55	MG	2A	3293	1/1	0.98	0.38	16,16,16,16	0
55	MG	1A	3178	1/1	0.98	0.09	17,17,17,17	0
55	MG	2A	3529	1/1	0.98	0.12	28,28,28,28	0
55	MG	2A	3018	1/1	0.98	0.14	10,10,10,10	0
55	MG	1A	3246	1/1	0.98	0.22	37,37,37,37	0
55	MG	2A	3804	1/1	0.98	0.09	31,31,31,31	0
55	MG	1a	3186	1/1	0.98	0.18	31,31,31,31	0
55	MG	2A	3636	1/1	0.98	0.12	28,28,28,28	0
55	MG	1A	3142	1/1	0.98	0.41	25,25,25,25	0
55	MG	2A	3803	1/1	0.98	0.18	32,32,32,32	0
55	MG	1A	3728	1/1	0.98	0.09	24,24,24,24	0
55	MG	2A	3022	1/1	0.98	0.22	27,27,27,27	0
55	MG	1x	108	1/1	0.98	0.21	28,28,28,28	0
55	MG	1A	3258	1/1	0.98	0.08	22,22,22,22	0
55	MG	1A	3705	1/1	0.98	0.12	36,36,36,36	0
55	MG	1A	3927	1/1	0.98	0.44	31,31,31,31	0
55	MG	2A	3446	1/1	0.98	0.44	59,59,59,59	0
55	MG	1A	3050	1/1	0.98	0.38	25,25,25,25	0
55	MG	2A	3400	1/1	0.98	0.22	13,13,13,13	0
55	MG	2A	3165	1/1	0.98	0.15	33,33,33,33	0
55	MG	2A	3257	1/1	0.98	0.19	35,35,35,35	0
55	MG	1A	3871	1/1	0.98	0.09	12,12,12,12	0
55	MG	1A	3237	1/1	0.98	0.09	25,25,25,25	0
55	MG	1A	3620	1/1	0.98	0.11	15,15,15,15	0
55	MG	1a	3097	1/1	0.98	0.14	23,23,23,23	0
55	MG	2A	3767	1/1	0.98	0.12	40,40,40,40	0
55	MG	1F	313	1/1	0.98	0.18	49,49,49,49	0
55	MG	2a	1672	1/1	0.98	0.10	21,21,21,21	0
55	MG	1A	3613	1/1	0.98	0.21	19,19,19,19	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3368	1/1	0.98	0.40	22,22,22,22	0
55	MG	1A	3181	1/1	0.98	0.09	27,27,27,27	0
55	MG	2A	3706	1/1	0.98	0.09	31,31,31,31	0
55	MG	2A	3295	1/1	0.98	0.42	31,31,31,31	0
55	MG	2A	3533	1/1	0.98	0.12	42,42,42,42	0
55	MG	2Q	203	1/1	0.98	0.41	49,49,49,49	0
55	MG	1A	3629	1/1	0.98	0.16	16,16,16,16	0
55	MG	2A	3626	1/1	0.98	0.07	21,21,21,21	0
55	MG	1B	217	1/1	0.98	0.23	32,32,32,32	0
55	MG	1A	3549	1/1	0.98	0.15	16,16,16,16	0
55	MG	16	102	1/1	0.98	0.12	18,18,18,18	0
55	MG	1A	3352	1/1	0.98	0.15	16,16,16,16	0
55	MG	1A	3676	1/1	0.98	0.11	25,25,25,25	0
55	MG	1a	3204	1/1	0.98	0.19	22,22,22,22	0
55	MG	2A	3192	1/1	0.98	0.11	25,25,25,25	0
55	MG	2A	3791	1/1	0.98	0.09	22,22,22,22	0
55	MG	1A	3020	1/1	0.98	0.17	24,24,24,24	0
55	MG	1A	3536	1/1	0.98	0.17	16,16,16,16	0
55	MG	1A	3645	1/1	0.98	0.16	30,30,30,30	0
55	MG	1A	3688	1/1	0.98	0.11	33,33,33,33	0
55	MG	2A	3370	1/1	0.98	0.13	25,25,25,25	0
55	MG	2A	3065	1/1	0.98	0.10	26,26,26,26	0
55	MG	1A	3321	1/1	0.98	0.22	18,18,18,18	0
55	MG	1A	3244	1/1	0.98	0.20	20,20,20,20	0
55	MG	1a	3125	1/1	0.98	0.20	21,21,21,21	0
55	MG	1A	3641	1/1	0.98	0.20	17,17,17,17	0
55	MG	2a	1764	1/1	0.98	0.36	41,41,41,41	0
55	MG	2A	3642	1/1	0.98	0.13	12,12,12,12	0
55	MG	2A	3127	1/1	0.98	0.19	32,32,32,32	0
55	MG	1R	201	1/1	0.98	0.36	40,40,40,40	0
55	MG	1A	3802	1/1	0.98	0.12	38,38,38,38	0
55	MG	2A	3190	1/1	0.98	0.20	42,42,42,42	0
55	MG	2A	3538	1/1	0.98	0.10	35,35,35,35	0
55	MG	1A	3454	1/1	0.98	0.14	21,21,21,21	0
55	MG	2A	3154	1/1	0.98	0.27	32,32,32,32	0
55	MG	1A	3950	1/1	0.98	0.15	27,27,27,27	0
55	MG	1A	3505	1/1	0.98	0.09	27,27,27,27	0
55	MG	1A	3626	1/1	0.98	0.15	14,14,14,14	0
55	MG	1A	3888	1/1	0.98	0.15	58,58,58,58	0
55	MG	1A	3844	1/1	0.98	0.18	23,23,23,23	0
55	MG	1R	203	1/1	0.98	0.34	30,30,30,30	0
55	MG	1A	3796	1/1	0.98	0.09	10,10,10,10	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2A	3024	1/1	0.98	0.12	25,25,25,25	0
55	MG	1A	3830	1/1	0.98	0.12	30,30,30,30	0
55	MG	1A	3197	1/1	0.98	0.07	16,16,16,16	0
55	MG	2x	102	1/1	0.98	0.21	41,41,41,41	0
55	MG	1E	307	1/1	0.98	0.21	23,23,23,23	0
55	MG	1A	3701	1/1	0.98	0.21	42,42,42,42	0
55	MG	1A	3537	1/1	0.98	0.07	30,30,30,30	0
55	MG	1F	310	1/1	0.98	0.56	25,25,25,25	0
55	MG	1E	305	1/1	0.98	0.20	34,34,34,34	0
55	MG	1A	3053	1/1	0.98	0.17	31,31,31,31	0
55	MG	1D	302	1/1	0.98	0.20	30,30,30,30	0
55	MG	1A	3441	1/1	0.98	0.34	38,38,38,38	0
55	MG	2A	3339	1/1	0.98	0.09	34,34,34,34	0
55	MG	1A	3631	1/1	0.98	0.08	42,42,42,42	0
55	MG	1A	4004	1/1	0.98	0.23	31,31,31,31	0
55	MG	1A	3824	1/1	0.98	0.18	14,14,14,14	0
55	MG	2A	3570	1/1	0.98	0.10	33,33,33,33	0
55	MG	2A	3559	1/1	0.98	0.12	24,24,24,24	0
55	MG	2A	3003	1/1	0.98	0.21	29,29,29,29	0
55	MG	1D	310	1/1	0.98	0.09	36,36,36,36	0
55	MG	2A	3001	1/1	0.98	0.14	29,29,29,29	0
55	MG	2A	3518	1/1	0.98	0.26	34,34,34,34	0
55	MG	1a	3208	1/1	0.98	0.10	41,41,41,41	0
55	MG	1O	204	1/1	0.98	0.38	38,38,38,38	0
55	MG	2A	3822	1/1	0.98	0.05	28,28,28,28	0
55	MG	1A	3847	1/1	0.98	0.10	21,21,21,21	0
55	MG	1A	3139	1/1	0.98	0.23	35,35,35,35	0
55	MG	2a	1631	1/1	0.98	0.12	42,42,42,42	0
55	MG	1A	3465	1/1	0.98	0.31	22,22,22,22	0
55	MG	2A	3624	1/1	0.98	0.18	27,27,27,27	0
55	MG	2a	1661	1/1	0.98	0.14	46,46,46,46	0
55	MG	1A	3566	1/1	0.98	0.07	18,18,18,18	0
55	MG	2A	3415	1/1	0.98	0.08	36,36,36,36	0
55	MG	1A	3738	1/1	0.98	0.20	42,42,42,42	0
55	MG	2A	3196	1/1	0.98	0.42	29,29,29,29	0
55	MG	1A	3741	1/1	0.98	0.13	16,16,16,16	0
55	MG	2A	3713	1/1	0.98	0.28	39,39,39,39	0
55	MG	1A	3387	1/1	0.98	0.21	24,24,24,24	0
55	MG	1a	3087	1/1	0.98	0.05	36,36,36,36	0
55	MG	2A	3133	1/1	0.98	0.23	33,33,33,33	0
55	MG	1A	3857	1/1	0.98	0.21	23,23,23,23	0
55	MG	2A	3109	1/1	0.98	0.33	41,41,41,41	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3850	1/1	0.98	0.29	17,17,17,17	0
55	MG	2A	3373	1/1	0.98	0.18	31,31,31,31	0
55	MG	2a	1684	1/1	0.98	0.09	68,68,68,68	0
55	MG	1A	3279	1/1	0.98	0.10	16,16,16,16	0
55	MG	1A	3483	1/1	0.98	0.19	40,40,40,40	0
55	MG	1a	3131	1/1	0.98	0.15	42,42,42,42	0
55	MG	2A	3829	1/1	0.98	0.19	33,33,33,33	0
55	MG	1a	3153	1/1	0.98	0.11	26,26,26,26	0
55	MG	1A	3341	1/1	0.98	0.09	43,43,43,43	0
55	MG	2a	1732	1/1	0.98	0.10	39,39,39,39	0
55	MG	1A	3257	1/1	0.98	0.16	25,25,25,25	0
55	MG	1A	3846	1/1	0.98	0.28	18,18,18,18	0
55	MG	2a	1729	1/1	0.98	0.12	48,48,48,48	0
55	MG	2A	3799	1/1	0.99	0.13	29,29,29,29	0
55	MG	2a	1800	1/1	0.99	0.13	28,28,28,28	0
55	MG	1A	3144	1/1	0.99	0.14	38,38,38,38	0
55	MG	1A	3960	1/1	0.99	0.14	14,14,14,14	0
55	MG	2A	3299	1/1	0.99	0.12	46,46,46,46	0
55	MG	1A	3975	1/1	0.99	0.10	23,23,23,23	0
55	MG	1A	3061	1/1	0.99	0.06	37,37,37,37	0
55	MG	1a	3133	1/1	0.99	0.09	47,47,47,47	0
55	MG	2a	1682	1/1	0.99	0.07	26,26,26,26	0
55	MG	2A	3556	1/1	0.99	0.11	26,26,26,26	0
55	MG	1A	3945	1/1	0.99	0.21	24,24,24,24	0
57	ZN	19	103	1/1	0.99	0.16	41,41,41,41	0
55	MG	2a	1786	1/1	0.99	0.15	39,39,39,39	0
55	MG	1A	3075	1/1	0.99	0.16	10,10,10,10	0
55	MG	1A	3625	1/1	0.99	0.12	20,20,20,20	0
55	MG	2A	3030	1/1	0.99	0.12	33,33,33,33	0
55	MG	1A	3399	1/1	0.99	0.15	17,17,17,17	0
55	MG	1a	3002	1/1	0.99	0.12	30,30,30,30	0
55	MG	1A	3929	1/1	0.99	0.12	18,18,18,18	0
55	MG	2A	3585	1/1	0.99	0.13	43,43,43,43	0
55	MG	1a	3160	1/1	0.99	0.15	39,39,39,39	0
59	SF4	2d	303	8/8	0.99	0.16	45,64,72,73	0
55	MG	1A	3894	1/1	0.99	0.15	16,16,16,16	0
55	MG	1A	3337	1/1	0.99	0.19	16,16,16,16	0
55	MG	1a	3157	1/1	0.99	0.13	34,34,34,34	0
55	MG	2A	3206	1/1	0.99	0.12	42,42,42,42	0
55	MG	1A	3584	1/1	0.99	0.14	31,31,31,31	0
55	MG	1A	3495	1/1	0.99	0.14	33,33,33,33	0
55	MG	2A	3227	1/1	0.99	0.19	29,29,29,29	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3269	1/1	0.99	0.14	25,25,25,25	0
55	MG	2A	3741	1/1	0.99	0.08	25,25,25,25	0
55	MG	1A	3072	1/1	0.99	0.11	27,27,27,27	0
55	MG	2A	3325	1/1	0.99	0.27	22,22,22,22	0
55	MG	1A	3709	1/1	0.99	0.06	28,28,28,28	0
55	MG	1A	3109	1/1	0.99	0.16	21,21,21,21	0
57	ZN	1Y	203	1/1	0.99	0.12	62,62,62,62	0
55	MG	1A	3784	1/1	0.99	0.31	24,24,24,24	0
55	MG	2A	3017	1/1	0.99	0.12	24,24,24,24	0
55	MG	1U	205	1/1	0.99	0.56	27,27,27,27	0
55	MG	1A	3036	1/1	0.99	0.27	30,30,30,30	0
55	MG	1A	3723	1/1	0.99	0.07	36,36,36,36	0
55	MG	2A	3811	1/1	0.99	0.17	10,10,10,10	0
55	MG	2a	1602	1/1	0.99	0.10	51,51,51,51	0
55	MG	1A	3385	1/1	0.99	0.16	26,26,26,26	0
55	MG	2A	3426	1/1	0.99	0.16	55,55,55,55	0
55	MG	1A	3477	1/1	0.99	0.15	25,25,25,25	0
55	MG	1A	3391	1/1	0.99	0.13	37,37,37,37	0
55	MG	1A	3303	1/1	0.99	0.19	37,37,37,37	0
55	MG	1A	3112	1/1	0.99	0.09	16,16,16,16	0
55	MG	2A	3385	1/1	0.99	0.09	21,21,21,21	0
55	MG	1a	3151	1/1	0.99	0.13	28,28,28,28	0
55	MG	1A	3099	1/1	0.99	0.12	23,23,23,23	0
55	MG	2A	3641	1/1	0.99	0.14	41,41,41,41	0
55	MG	2A	3606	1/1	0.99	0.17	34,34,34,34	0
55	MG	2P	202	1/1	0.99	0.21	29,29,29,29	0
55	MG	2A	3056	1/1	0.99	0.20	23,23,23,23	0
55	MG	1A	3834	1/1	0.99	0.26	11,11,11,11	0
55	MG	1A	3450	1/1	0.99	0.07	14,14,14,14	0
55	MG	1A	3783	1/1	0.99	0.26	21,21,21,21	0
55	MG	1A	3243	1/1	0.99	0.35	26,26,26,26	0
55	MG	1A	3422	1/1	0.99	0.11	36,36,36,36	0
55	MG	1a	3129	1/1	0.99	0.11	32,32,32,32	0
55	MG	1A	3616	1/1	0.99	0.08	12,12,12,12	0
55	MG	2a	1784	1/1	0.99	0.27	32,32,32,32	0
55	MG	2A	3801	1/1	0.99	0.05	44,44,44,44	0
55	MG	2Y	202	1/1	0.99	0.20	20,20,20,20	0
55	MG	2A	3445	1/1	0.99	0.36	32,32,32,32	0
55	MG	1A	3637	1/1	0.99	0.19	16,16,16,16	0
55	MG	2A	3123	1/1	0.99	0.14	16,16,16,16	0
55	MG	2A	3704	1/1	0.99	0.06	26,26,26,26	0
55	MG	1A	3118	1/1	0.99	0.24	24,24,24,24	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	2D	304	1/1	0.99	0.19	22,22,22,22	0
55	MG	1D	308	1/1	0.99	0.13	42,42,42,42	0
55	MG	2A	3462	1/1	0.99	0.24	15,15,15,15	0
55	MG	1A	3442	1/1	0.99	0.18	22,22,22,22	0
55	MG	2A	3104	1/1	0.99	0.33	28,28,28,28	0
55	MG	2A	3515	1/1	0.99	0.14	24,24,24,24	0
55	MG	1a	3190	1/1	0.99	0.32	44,44,44,44	0
55	MG	1A	3384	1/1	0.99	0.09	21,21,21,21	0
55	MG	2A	3583	1/1	0.99	0.09	19,19,19,19	0
55	MG	1A	3335	1/1	0.99	0.10	19,19,19,19	0
55	MG	2A	3219	1/1	0.99	0.15	24,24,24,24	0
57	ZN	26	102	1/1	0.99	0.18	40,40,40,40	0
55	MG	1B	206	1/1	0.99	0.14	45,45,45,45	0
55	MG	2a	1675	1/1	0.99	0.14	44,44,44,44	0
55	MG	1A	3848	1/1	0.99	0.18	22,22,22,22	0
55	MG	2A	3387	1/1	0.99	0.21	25,25,25,25	0
55	MG	1A	3440	1/1	0.99	0.17	19,19,19,19	0
55	MG	1A	3132	1/1	0.99	0.10	35,35,35,35	0
55	MG	1a	3126	1/1	0.99	0.09	30,30,30,30	0
55	MG	2A	3309	1/1	0.99	0.11	48,48,48,48	0
55	MG	1A	3828	1/1	0.99	0.24	28,28,28,28	0
55	MG	1N	201	1/1	0.99	0.24	47,47,47,47	0
55	MG	1A	3593	1/1	0.99	0.14	22,22,22,22	0
55	MG	2A	3505	1/1	0.99	0.13	18,18,18,18	0
55	MG	1A	3801	1/1	0.99	0.14	10,10,10,10	0
55	MG	1A	3123	1/1	0.99	0.13	10,10,10,10	0
55	MG	1A	3186	1/1	0.99	0.17	34,34,34,34	0
55	MG	1a	3098	1/1	0.99	0.10	26,26,26,26	0
59	SF4	1d	302	8/8	0.99	0.16	47,72,75,78	0
55	MG	1A	3327	1/1	0.99	0.19	26,26,26,26	0
55	MG	1B	236	1/1	0.99	0.17	21,21,21,21	0
55	MG	1A	3562	1/1	0.99	0.06	25,25,25,25	0
55	MG	2A	3107	1/1	0.99	0.09	39,39,39,39	0
55	MG	1A	3753	1/1	0.99	0.06	28,28,28,28	0
55	MG	1A	3018	1/1	0.99	0.13	16,16,16,16	0
55	MG	1a	3205	1/1	0.99	0.07	41,41,41,41	0
55	MG	1A	3546	1/1	0.99	0.13	35,35,35,35	0
55	MG	1A	3557	1/1	0.99	0.21	54,54,54,54	0
55	MG	2A	3812	1/1	0.99	0.12	28,28,28,28	0
55	MG	1B	202	1/1	0.99	0.22	48,48,48,48	0
55	MG	1A	3393	1/1	0.99	0.70	21,21,21,21	0
55	MG	2a	1766	1/1	1.00	0.06	39,39,39,39	0

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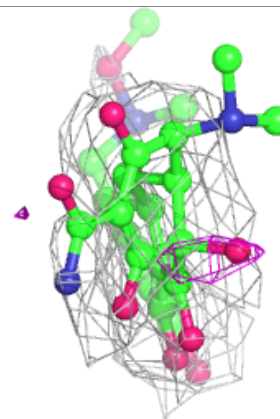
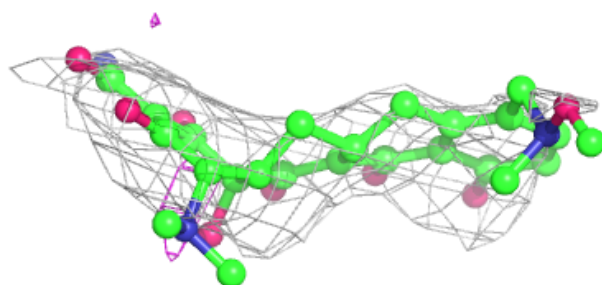
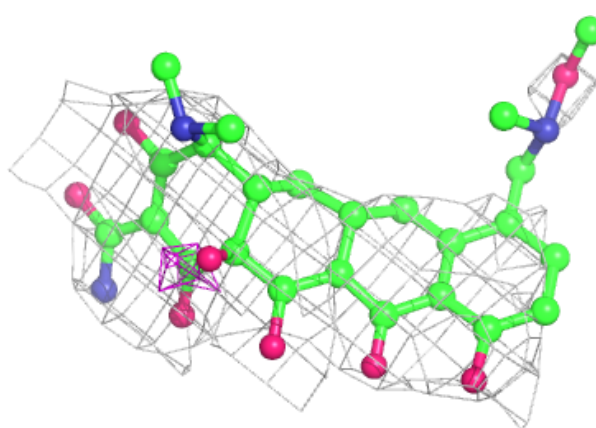
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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(\AA^2)	Q<0.9
55	MG	1A	3022	1/1	1.00	0.23	29,29,29,29	0

The following is a graphical depiction of the model fit to experimental electron density of all instances of the Ligand of Interest. In addition, ligands with molecular weight > 250 and outliers as shown on the geometry validation Tables will also be included. Each fit is shown from different orientation to approximate a three-dimensional view.

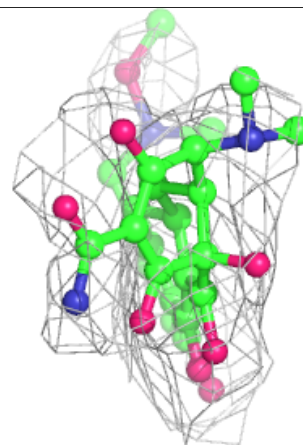
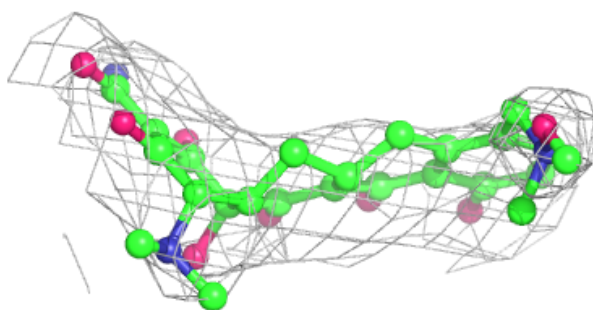
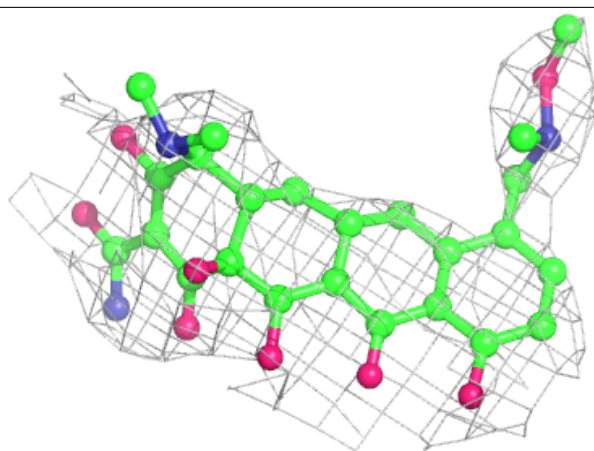
Electron density around V7A 2a 1817:

2mF_o-DF_c (at 0.7 rmsd) in gray
mF_o-DF_c (at 3 rmsd) in purple (negative)
and green (positive)



Electron density around V7A 1a 3213:

$2mF_o-DF_c$ (at 0.7 rmsd) in gray
 mF_o-DF_c (at 3 rmsd) in purple (negative)
and green (positive)



6.5 Other polymers ⓘ

There are no such residues in this entry.