



wwPDB X-ray Structure Validation Summary Report ⓘ

Apr 19, 2026 – 03:26 AM UTC

PDB ID : 5WIS / pdb_00005wis
Title : Crystal structure of the Thermus thermophilus 70S ribosome in complex with methymycin and bound to mRNA and A-, P- and E-site tRNAs at 2.7Å resolution
Authors : Almutairi, M.M.; Svetlov, M.S.; Hansen, D.A.; Khabibullina, N.F.; Klepacki, D.; Kang, H.Y.; Sherman, D.H.; Vazquez-Laslop, N.; Polikanov, Y.S.; Mankin, A.S.
Deposited on : 2017-07-20
Resolution : 2.70 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 types of validation reports are described at

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

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

MolProbity	:	4-5-2 with Phenix2.0
Mogul	:	2022.3.0, CSD as543be (2022)
Xtriage (Phenix)	:	2.0
EDS	:	3.0
Buster-report	:	wwPDB partial adaption of 1.1.7 (2018)
Percentile statistics	:	20250101.v01 (using entries in the PDB archive January 1st 2025)
CCP4	:	9.0.010 (Gargrove)
Density-Fitness	:	1.0.12
Ideal geometry (proteins)	:	Engh & Huber (2001)

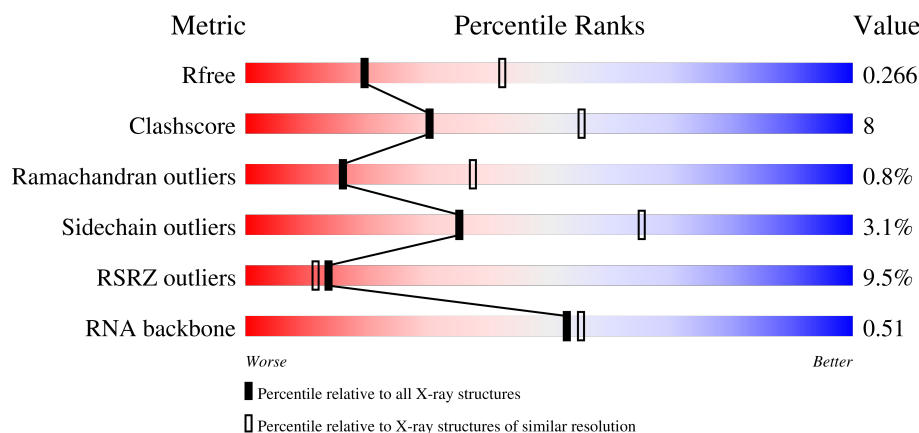
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

The reported resolution of this entry is 2.70 Å.

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



Metric	Whole archive (#Entries)	Similar resolution (#Entries, resolution range(Å))
R_{free}	180053	3538 (2.70-2.70)
Clashscore	190562	3843 (2.70-2.70)
Ramachandran outliers	187476	3778 (2.70-2.70)
Sidechain outliers	187428	3778 (2.70-2.70)
RSRZ outliers	180081	3538 (2.70-2.70)
RNA backbone	3983	1044 (2.90-2.50)

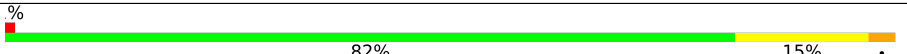
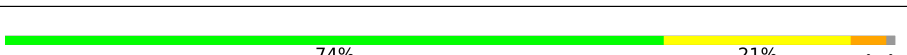
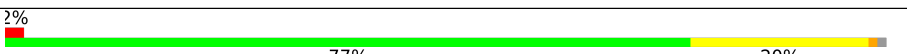
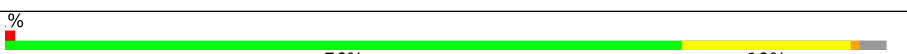
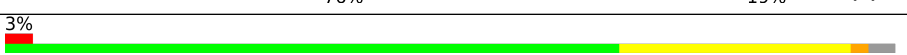
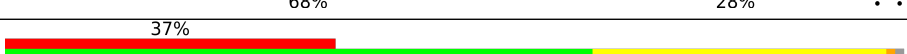
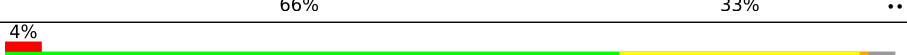
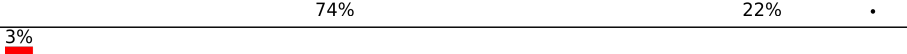
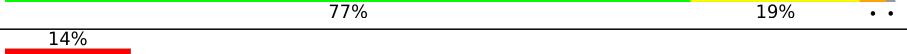
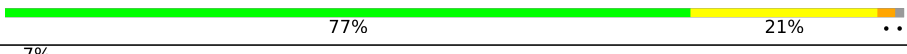
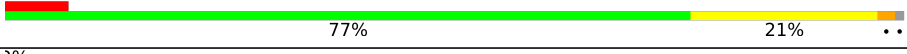
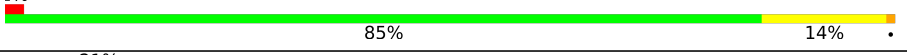
The table below summarises the geometric issues observed across the polymeric chains and their fit to the electron density. The red, orange, yellow and green segments of the lower bar indicate the fraction of residues that contain outliers for ≥ 3 , 2, 1 and 0 types of geometric quality criteria respectively.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

Mol	Chain	Length	Quality of chain
1	1A	2915	4% 64% 28% 6%
1	2A	2915	3% 58% 32% 6%

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Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.49

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Mol	Chain	Length	Quality of chain
2	1B	121	
2	2B	121	
3	1D	276	
3	2D	276	
4	1E	206	
4	2E	206	
5	1F	210	
5	2F	210	
6	1G	182	
6	2G	182	
7	1H	180	
7	2H	180	
8	1I	148	
8	2I	148	
9	1N	140	
9	2N	140	
10	1O	122	
10	2O	122	
11	1P	150	
11	2P	150	
12	1Q	141	
12	2Q	141	
13	1R	118	
13	2R	118	
14	1S	112	

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Mol	Chain	Length	Quality of chain
14	2S	112	
15	1T	146	
15	2T	146	
16	1U	118	
16	2U	118	
17	1V	101	
17	2V	101	
18	1W	113	
18	2W	113	
19	1X	96	
19	2X	96	
20	1Y	110	
20	2Y	110	
21	1Z	206	
21	2Z	206	
22	10	85	
22	20	85	
23	11	98	
23	21	98	
24	12	72	
24	22	72	
25	13	60	
25	23	60	
26	14	71	
26	24	71	

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Mol	Chain	Length	Quality of chain
27	15	60	
27	25	60	
28	16	54	
28	26	54	
29	17	49	
29	27	49	
30	18	65	
30	28	65	
31	19	37	
31	29	37	
32	1a	1521	
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	

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Mol	Chain	Length	Quality of chain
39	2h	138	
40	1i	128	
40	2i	128	
41	1j	105	
41	2j	105	
42	1k	129	
42	2k	129	
43	1l	132	
43	2l	132	
44	1m	126	
44	2m	126	
45	1n	61	
45	2n	61	
46	1o	89	
46	2o	89	
47	1p	88	
47	2p	88	
48	1q	105	
48	2q	105	
49	1r	88	
49	2r	88	
50	1s	93	
50	2s	93	
51	1t	106	
51	2t	106	

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Mol	Chain	Length	Quality of chain
52	1u	27	
52	2u	27	
53	1v	24	
53	2v	24	
54	1w	76	
54	1y	76	
54	2w	76	
54	2y	76	
55	1x	77	
55	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
56	MG	2A	3884	-	-	-	X

2 Entry composition

There are 61 unique types of molecules in this entry. The entry contains 301023 atoms, of which 0 are hydrogens and 0 are deuteriums.

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

- Molecule 1 is a RNA chain called 23S ribosomal RNA.

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
32	1a	1500	Total	C	N	O	P	0	0	0
			32246	14358	5975	10413	1500			
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	123	Total	C	N	O	S	0	0	0
			958	592	198	166	2			
44	2m	122	Total	C	N	O	S	0	0	0
			950	586	197	165	2			

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

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

- Molecule 53 is a RNA chain called mRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
53	1v	13	Total	C	N	O	P	0	0	0
			277	125	51	88	13			
53	2v	13	Total	C	N	O	P	0	0	0
			277	125	51	88	13			

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

Mol	Chain	Residues	Atoms						ZeroOcc	AltConf	Trace
54	1w	74	Total 1592	C 713	N 285	O 518	P 74	S 2	0	0	0
54	1y	74	Total 1585	C 707	N 285	O 518	P 74	S 1	0	0	0
54	2w	72	Total 1544	C 690	N 278	O 502	P 72	S 2	0	0	0
54	2y	73	Total 1565	C 698	N 283	O 510	P 73	S 1	0	0	0

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

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

- Molecule 56 is MAGNESIUM ION (CCD ID: MG) (formula: Mg).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	1A	1146	Total	Mg	0	0
			1146	1146		
56	1B	36	Total	Mg	0	0
			36	36		
56	1D	12	Total	Mg	0	0
			12	12		
56	1E	12	Total	Mg	0	0
			12	12		
56	1F	8	Total	Mg	0	0
			8	8		
56	1G	5	Total	Mg	0	0
			5	5		
56	1H	1	Total	Mg	0	0
			1	1		
56	1I	1	Total	Mg	0	0
			1	1		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	1N	6	Total 6	Mg 6	0	0
56	1O	8	Total 8	Mg 8	0	0
56	1P	4	Total 4	Mg 4	0	0
56	1Q	5	Total 5	Mg 5	0	0
56	1R	2	Total 2	Mg 2	0	0
56	1S	3	Total 3	Mg 3	0	0
56	1T	1	Total 1	Mg 1	0	0
56	1U	6	Total 6	Mg 6	0	0
56	1V	3	Total 3	Mg 3	0	0
56	1W	7	Total 7	Mg 7	0	0
56	1X	5	Total 5	Mg 5	0	0
56	1Y	3	Total 3	Mg 3	0	0
56	1Z	4	Total 4	Mg 4	0	0
56	10	9	Total 9	Mg 9	0	0
56	11	4	Total 4	Mg 4	0	0
56	12	2	Total 2	Mg 2	0	0
56	13	3	Total 3	Mg 3	0	0
56	15	2	Total 2	Mg 2	0	0
56	16	3	Total 3	Mg 3	0	0
56	17	3	Total 3	Mg 3	0	0
56	18	3	Total 3	Mg 3	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	19	1	Total 1	Mg 1	0	0
56	1a	229	Total 229	Mg 229	0	0
56	1b	2	Total 2	Mg 2	0	0
56	1d	1	Total 1	Mg 1	0	0
56	1e	2	Total 2	Mg 2	0	0
56	1f	2	Total 2	Mg 2	0	0
56	1h	1	Total 1	Mg 1	0	0
56	1l	3	Total 3	Mg 3	0	0
56	1m	1	Total 1	Mg 1	0	0
56	1n	1	Total 1	Mg 1	0	0
56	1o	1	Total 1	Mg 1	0	0
56	1q	1	Total 1	Mg 1	0	0
56	1r	1	Total 1	Mg 1	0	0
56	1s	1	Total 1	Mg 1	0	0
56	1t	1	Total 1	Mg 1	0	0
56	1v	1	Total 1	Mg 1	0	0
56	1w	11	Total 11	Mg 11	0	0
56	1x	16	Total 16	Mg 16	0	0
56	1y	4	Total 4	Mg 4	0	0
56	2A	903	Total 903	Mg 903	0	0
56	2B	21	Total 21	Mg 21	0	0

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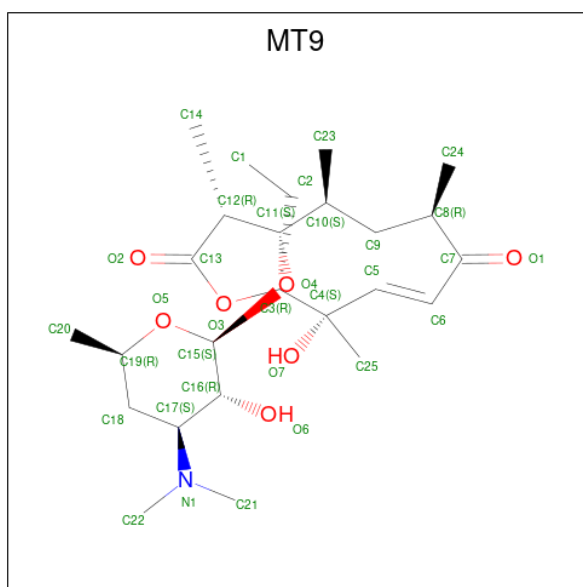
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	2D	4	Total 4	Mg 4	0	0
56	2E	8	Total 8	Mg 8	0	0
56	2F	4	Total 4	Mg 4	0	0
56	2G	1	Total 1	Mg 1	0	0
56	2N	1	Total 1	Mg 1	0	0
56	2O	1	Total 1	Mg 1	0	0
56	2Q	4	Total 4	Mg 4	0	0
56	2R	2	Total 2	Mg 2	0	0
56	2T	2	Total 2	Mg 2	0	0
56	2U	5	Total 5	Mg 5	0	0
56	2V	2	Total 2	Mg 2	0	0
56	2W	3	Total 3	Mg 3	0	0
56	2X	2	Total 2	Mg 2	0	0
56	2Y	1	Total 1	Mg 1	0	0
56	2Z	1	Total 1	Mg 1	0	0
56	20	3	Total 3	Mg 3	0	0
56	23	3	Total 3	Mg 3	0	0
56	25	4	Total 4	Mg 4	0	0
56	26	1	Total 1	Mg 1	0	0
56	27	1	Total 1	Mg 1	0	0
56	28	1	Total 1	Mg 1	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	2a	244	Total 244	Mg 244	0	0
56	2d	1	Total 1	Mg 1	0	0
56	2e	1	Total 1	Mg 1	0	0
56	2f	2	Total 2	Mg 2	0	0
56	2g	1	Total 1	Mg 1	0	0
56	2j	2	Total 2	Mg 2	0	0
56	2l	3	Total 3	Mg 3	0	0
56	2q	3	Total 3	Mg 3	0	0
56	2r	1	Total 1	Mg 1	0	0
56	2t	1	Total 1	Mg 1	0	0
56	2v	3	Total 3	Mg 3	0	0
56	2w	9	Total 9	Mg 9	0	0
56	2x	6	Total 6	Mg 6	0	0
56	2y	7	Total 7	Mg 7	0	0

- Molecule 57 is (3R,4S,5S,7R,9E,11S,12R)-12-ethyl-11-hydroxy-3,5,7,11-tetramethyl-2,8-dioxooxacyclododec-9-en-4-yl 3,4,6-trideoxy-3-(dimethylamino)-beta-D-xylo-hexopyranoside (CCD ID: MT9) (formula: C₂₅H₄₃NO₇).



Mol	Chain	Residues	Atoms				ZeroOcc	AltConf
57	1A	1	Total	C	N	O	0	0
			33	25	1	7		
57	2A	1	Total	C	N	O	0	0
			33	25	1	7		

- Molecule 58 is POTASSIUM ION (CCD ID: K) (formula: K).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
58	1A	1	Total	K	0	0
			1	1		
58	2A	1	Total	K	0	0
			1	1		

- Molecule 59 is ZINC ION (CCD ID: ZN) (formula: Zn).

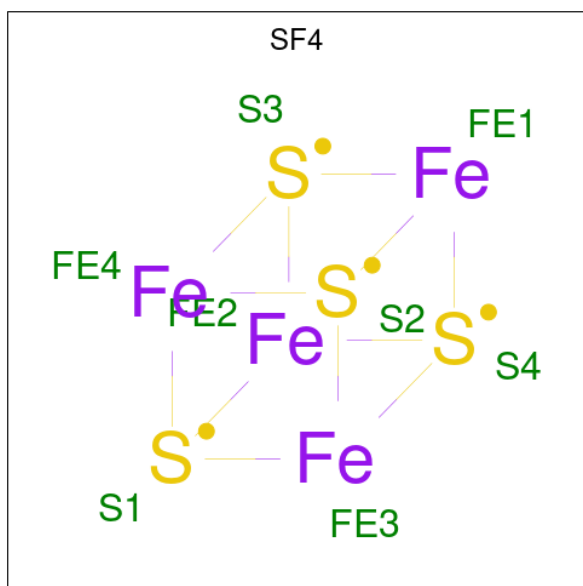
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	1Y	1	Total	Zn	0	0
			1	1		
59	14	1	Total	Zn	0	0
			1	1		
59	15	1	Total	Zn	0	0
			1	1		
59	16	1	Total	Zn	0	0
			1	1		
59	19	1	Total	Zn	0	0
			1	1		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	1n	1	Total	Zn	0	0
			1	1		
59	2Y	1	Total	Zn	0	0
			1	1		
59	24	1	Total	Zn	0	0
			1	1		
59	25	1	Total	Zn	0	0
			1	1		
59	26	1	Total	Zn	0	0
			1	1		
59	29	1	Total	Zn	0	0
			1	1		
59	2n	1	Total	Zn	0	0
			1	1		

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



Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
60	1d	1	Total	Fe	S	0	0
			8	4	4		
60	2d	1	Total	Fe	S	0	0
			8	4	4		

- Molecule 61 is water.

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
61	1A	2243	Total 2243	O 2243	0	0
61	1B	69	Total 69	O 69	0	0
61	1D	26	Total 26	O 26	0	0
61	1E	28	Total 28	O 28	0	0
61	1F	19	Total 19	O 19	0	0
61	1G	6	Total 6	O 6	0	0
61	1H	2	Total 2	O 2	0	0
61	1I	2	Total 2	O 2	0	0
61	1N	6	Total 6	O 6	0	0
61	1O	7	Total 7	O 7	0	0
61	1P	20	Total 20	O 20	0	0
61	1Q	15	Total 15	O 15	0	0
61	1R	12	Total 12	O 12	0	0
61	1S	5	Total 5	O 5	0	0
61	1T	10	Total 10	O 10	0	0
61	1U	16	Total 16	O 16	0	0
61	1V	10	Total 10	O 10	0	0
61	1W	6	Total 6	O 6	0	0
61	1X	6	Total 6	O 6	0	0
61	1Y	4	Total 4	O 4	0	0
61	1Z	1	Total 1	O 1	0	0
61	10	11	Total 11	O 11	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
61	11	12	Total 12	O 12	0	0
61	12	3	Total 3	O 3	0	0
61	13	5	Total 5	O 5	0	0
61	14	1	Total 1	O 1	0	0
61	15	4	Total 4	O 4	0	0
61	16	2	Total 2	O 2	0	0
61	17	9	Total 9	O 9	0	0
61	18	13	Total 13	O 13	0	0
61	1a	460	Total 460	O 460	0	0
61	1b	1	Total 1	O 1	0	0
61	1d	1	Total 1	O 1	0	0
61	1f	1	Total 1	O 1	0	0
61	1g	1	Total 1	O 1	0	0
61	1l	9	Total 9	O 9	0	0
61	1m	1	Total 1	O 1	0	0
61	1o	1	Total 1	O 1	0	0
61	1p	1	Total 1	O 1	0	0
61	1q	4	Total 4	O 4	0	0
61	1u	1	Total 1	O 1	0	0
61	1v	5	Total 5	O 5	0	0
61	1w	22	Total 22	O 22	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
61	1x	16	Total 16	O 16	0	0
61	1y	2	Total 2	O 2	0	0
61	2A	1395	Total 1395	O 1395	0	0
61	2B	28	Total 28	O 28	0	0
61	2D	25	Total 25	O 25	0	0
61	2E	15	Total 15	O 15	0	0
61	2F	14	Total 14	O 14	0	0
61	2I	4	Total 4	O 4	0	0
61	2N	1	Total 1	O 1	0	0
61	2O	1	Total 1	O 1	0	0
61	2P	14	Total 14	O 14	0	0
61	2Q	2	Total 2	O 2	0	0
61	2R	3	Total 3	O 3	0	0
61	2T	6	Total 6	O 6	0	0
61	2U	4	Total 4	O 4	0	0
61	2V	1	Total 1	O 1	0	0
61	2W	3	Total 3	O 3	0	0
61	2X	4	Total 4	O 4	0	0
61	2Y	1	Total 1	O 1	0	0
61	2Z	2	Total 2	O 2	0	0
61	20	4	Total 4	O 4	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
61	21	14	Total 14	O 14	0	0
61	22	2	Total 2	O 2	0	0
61	23	1	Total 1	O 1	0	0
61	25	5	Total 5	O 5	0	0
61	27	3	Total 3	O 3	0	0
61	28	4	Total 4	O 4	0	0
61	29	1	Total 1	O 1	0	0
61	2a	381	Total 381	O 381	0	0
61	2e	2	Total 2	O 2	0	0
61	2g	1	Total 1	O 1	0	0
61	2i	1	Total 1	O 1	0	0
61	2j	4	Total 4	O 4	0	0
61	2l	6	Total 6	O 6	0	0
61	2o	3	Total 3	O 3	0	0
61	2p	3	Total 3	O 3	0	0
61	2q	1	Total 1	O 1	0	0
61	2r	1	Total 1	O 1	0	0
61	2t	2	Total 2	O 2	0	0
61	2u	1	Total 1	O 1	0	0
61	2v	1	Total 1	O 1	0	0
61	2w	3	Total 3	O 3	0	0

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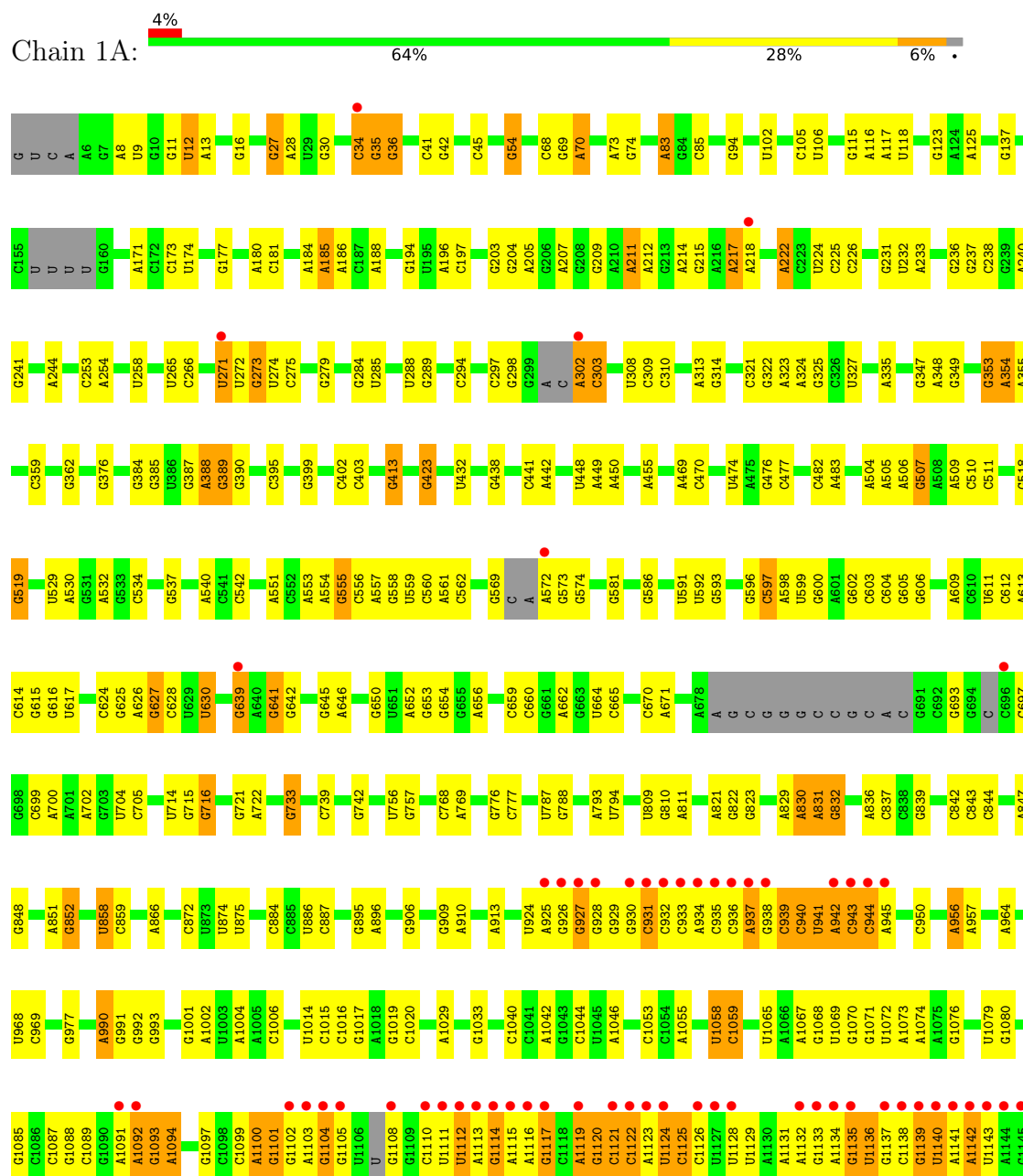
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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
61	2x	8	Total	O	0	0
			8	8		
61	2y	19	Total	O	0	0
			19	19		

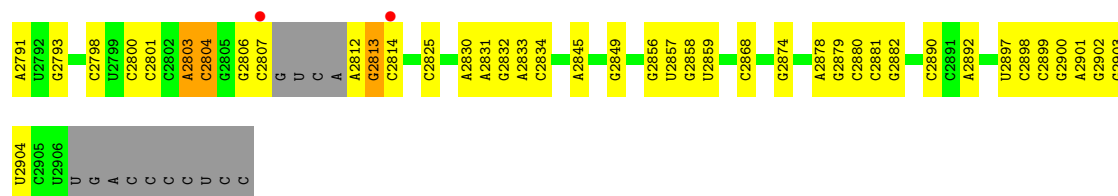
3 Residue-property plots

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.

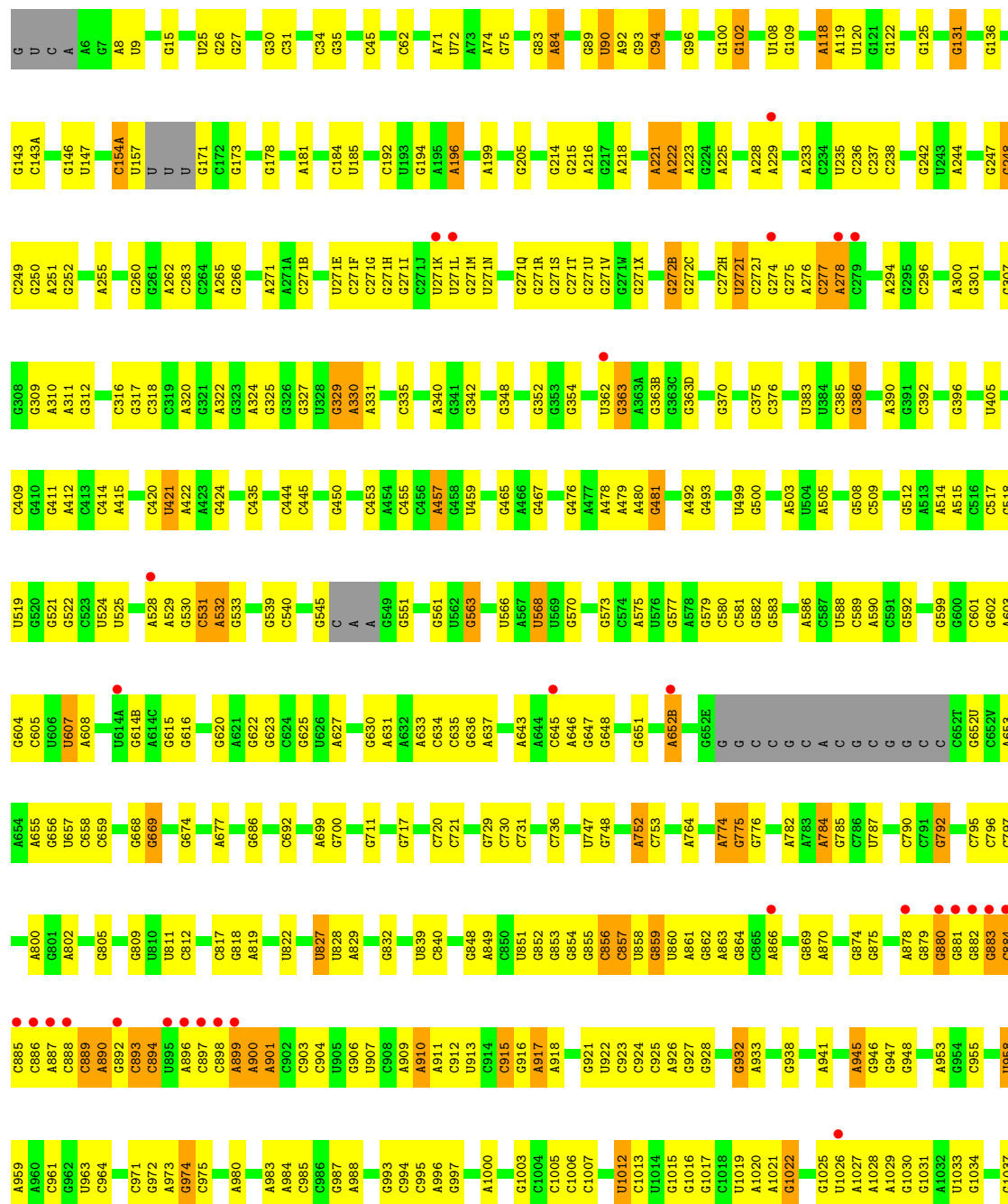
- Molecule 1: 23S ribosomal RNA



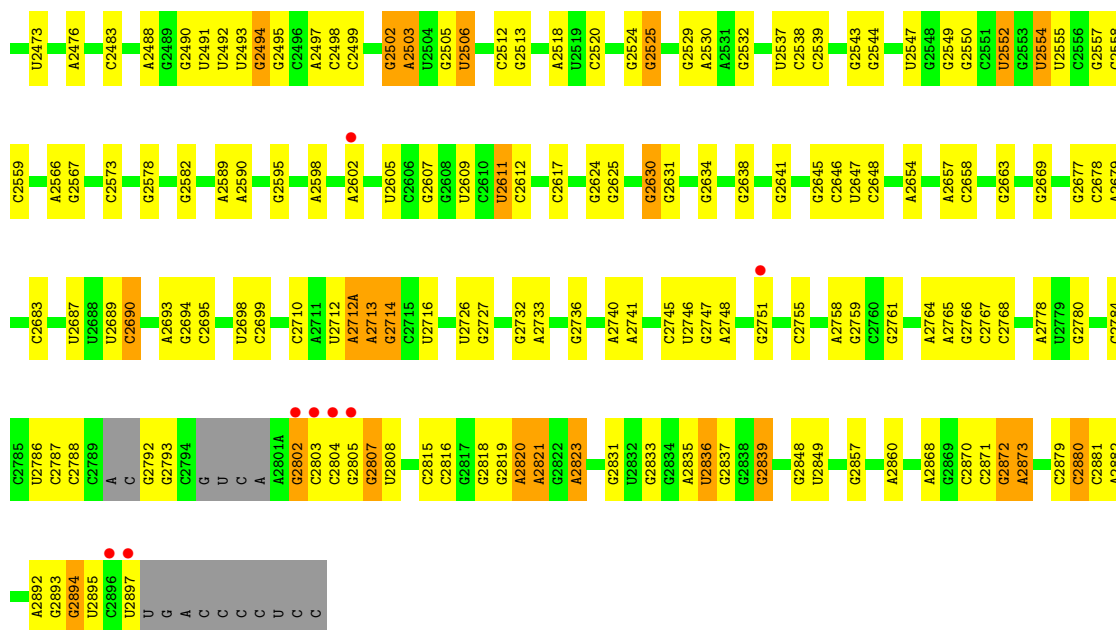
U2661	A2530	C2416	C2306	C2200	U2140	G2034	A1934	U1809	A1679	C1550	A1405	U1256	C1146
U2662	U2534	G2417	G2314	G2203	A2141	A2035	A1935	U1810	A1679	C1550	A1406	G1257	U1147
A2666	G2315	U2418	G2315	G2204	G2142	A2036	C1936	C1811	C1683	C1551	G1410	A1258	C1148
A2669	G2541	U2420	G2316	G2205	U2144	G2040	U1937	C1812	C1683	C1552	G1411	C1265	C1150
A2672	G2544	G2421	A2317	G2206	G2145	A2041	U1939	A1814	C1685	A1554	G1417	C1276	U1151
G2673	U2549	G2422	G2320	G2209	G2146	A2042	U1940	A1815	U1686	C1555	U1418	G1277	G1152
G2674	C2550	U2431	U2324	G2210	G2147	C2043	A1941	A1816	G1694	A1556	A1419	U1283	G1153
C2679	G2561	A2434	G2327	U2211	G2148	G2045	A1949	C1821	G1695	U1566	A1425	G1284	G1154
G2680	G2562	A2437	C2328	G2212	C2150	G2045	A1950	A1822	G1697	G1567	G1426	G1285	U1157
A2691	C2563	A2437	G2331	G2213	U2151	U2050	G1951	U1825	G1698	G1572	A1430	U1286	G1158
C2695	U2564	G2440	A2332	G2214	G2152	A2051	G1952	C1826	A1699	G1573	A1430	G1290	U1159
U2699	U2566	G2441	G2337	U2225	G2153	A2052	G1953	U1827	G1700	A1574	G1431	G1290	G1160
U2700	A2576	A2442	G2337	G2226	U2154	G2053	U1953	U1828	A1701	A1575	C1432	G1294	G1161
U2701	A2577	U2443	G2338	G2227	A2157	A2055	G1955	U1829	C1703	U1452	U1452	U1294	C1162
C2702	G2579	U2447	A2339	G2228	C2158	C2061	A1959	G1830	C1710	C1579	C1453	A1299	G1171
C2711	C2585	A2451	G2341	A2229	C2159	C2065	A1960	C1831	A1711	G	U	G1302	A1174
U2714	G2585	C2452	G2342	A2237	C2161	G2074	U1961	G1832	C1717	A	C	G1312	U1176
C2715	G2594	C2453	G2343	G2247	G2163	C2077	C1964	A1834	U1720	G1584	U1465	U1313	C1180
G2718	G2595	G2457	G2346	G2250	C2164	G2078	G1971	A1841	G1721	G1585	U1466	A1314	G1181
C2721	C2598	A2347	A2347	G2251	U2166	A2082	A1974	G1842	G1735	G1586	G1467	A1315	G1182
C2722	A2602	G2458	G2348	G2252	C2168	G2083	A1975	G1847	U1740	U1587	A1473	G1317	G1183
A2723	C2603	A2459	G2359	U2255	C2169	C2087	U1977	G1848	G1746	C1593	G1475	A1318	G1190
U2725	U2614	A2460	C2359	U2256	G2170	C2088	A1982	G1857	C1741	A1605	U1477	A1320	G1199
C2727	G2617	G2466	C2362	G2262	U2171	C2088	C1983	A1860	G1742	A1613	C1478	A1324	G1200
U2735	U2621	U2481	G2363	G2263	C2172	G2091	A1984	A1864	G1744	G1614	U1479	A1201	U1211
C2736	C2622	G2482	G2365	G2264	G2173	G2105	U1985	U1864	A1745	G1615	U1538	A1202	G1212
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• Molecule 1: 23S ribosomal RNA

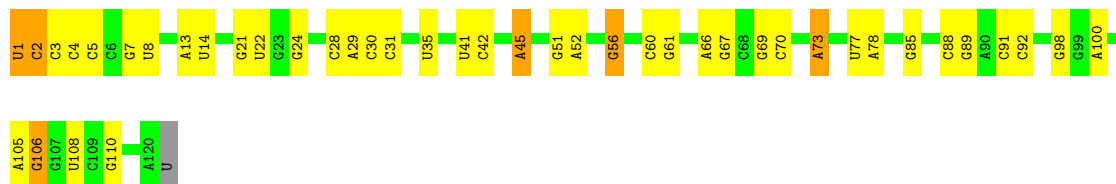


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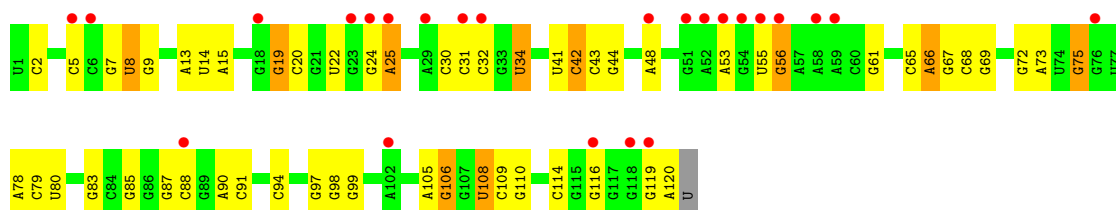
• Molecule 2: 5S Ribosomal RNA

Chain 1B: 64% 31% 5% .



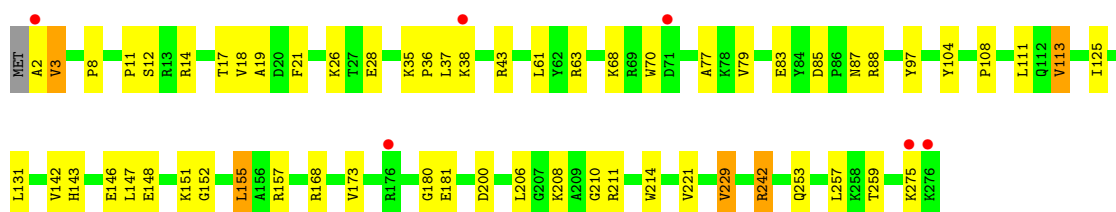
• Molecule 2: 5S Ribosomal RNA

Chain 2B: 20% 53% 38% 8% .

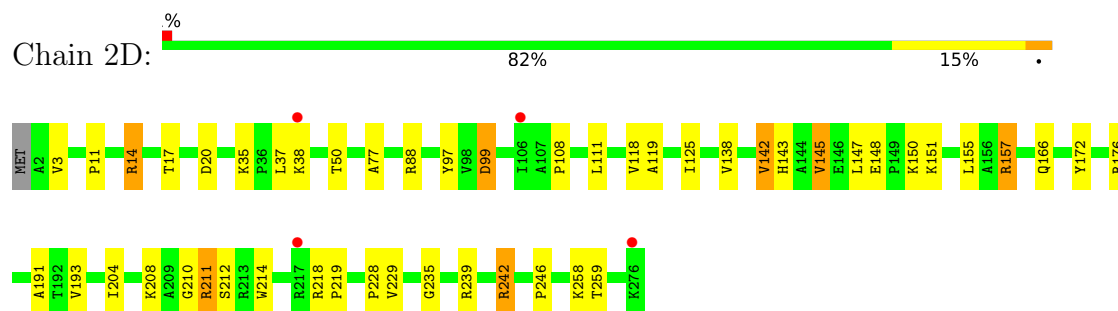


• Molecule 3: 50S ribosomal protein L2

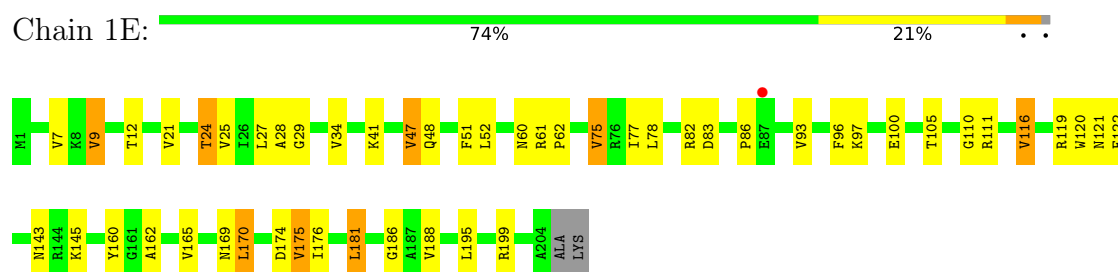
Chain 1D: 2% 78% 20% .



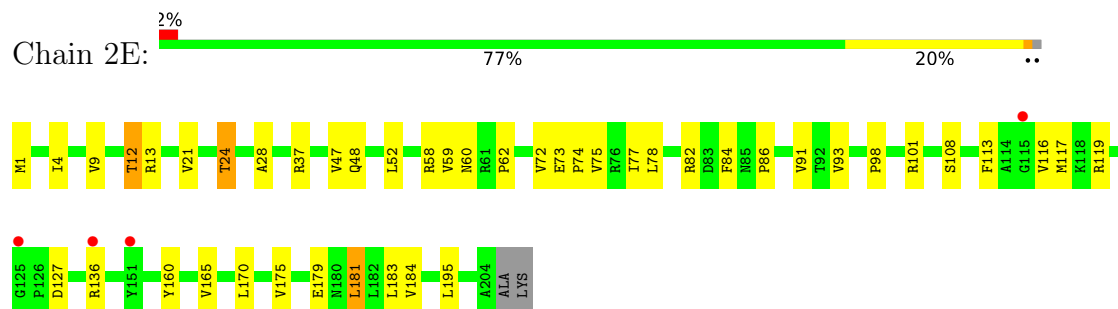
- Molecule 3: 50S ribosomal protein L2



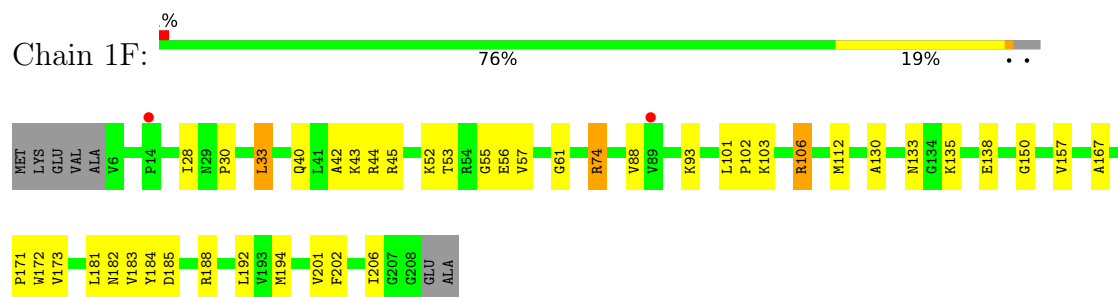
- Molecule 4: 50S ribosomal protein L3



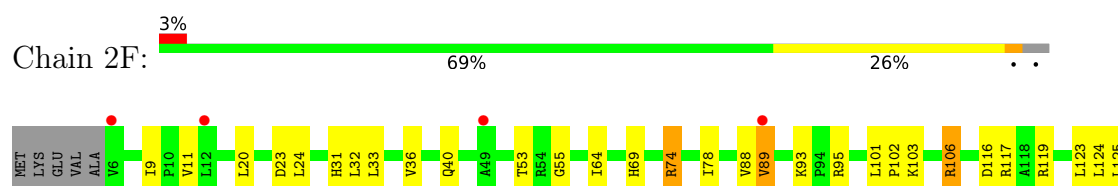
- Molecule 4: 50S ribosomal protein L3

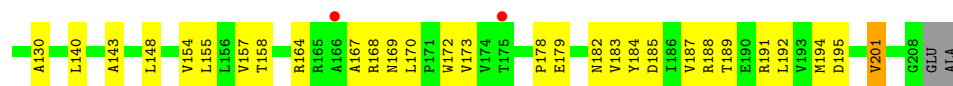


- Molecule 5: 50S ribosomal protein L4

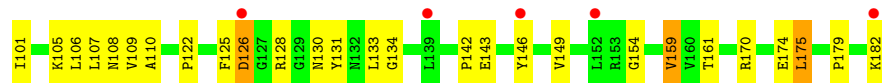
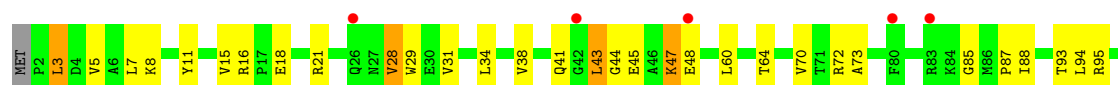


- Molecule 5: 50S ribosomal protein L4

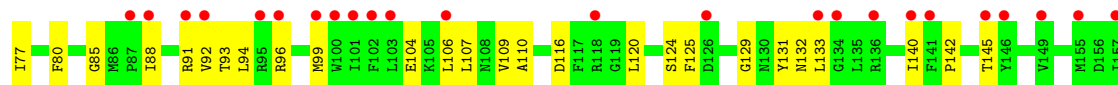
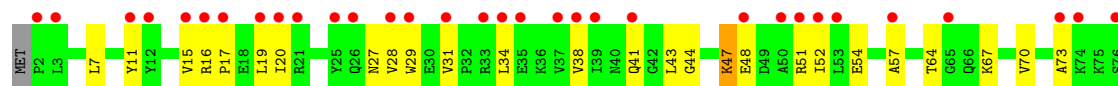




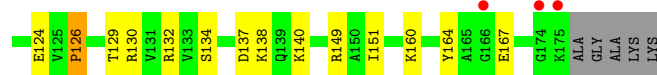
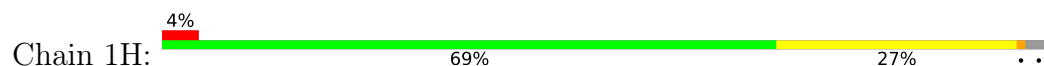
- Molecule 6: 50S ribosomal protein L5



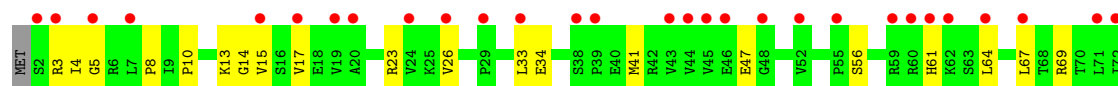
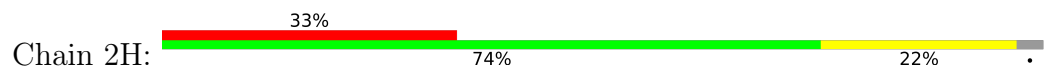
- Molecule 6: 50S ribosomal protein L5

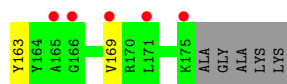


- Molecule 7: 50S ribosomal protein L6

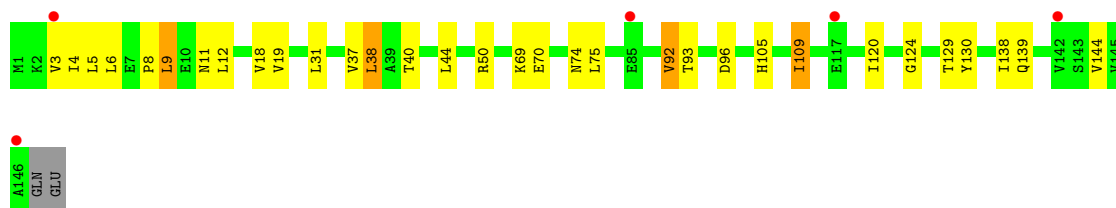
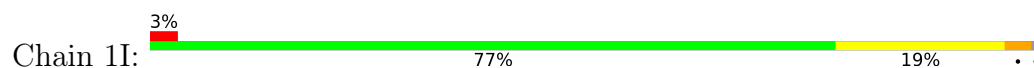


- Molecule 7: 50S ribosomal protein L6

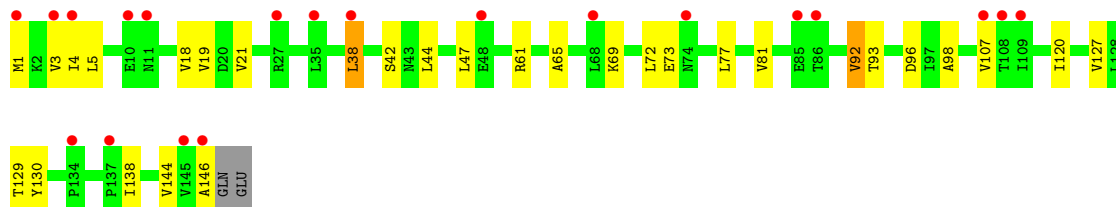
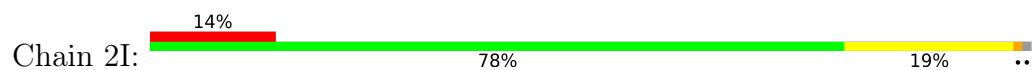




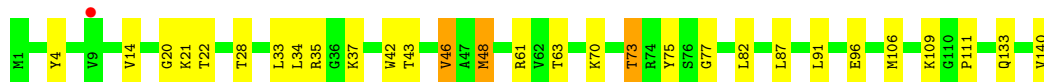
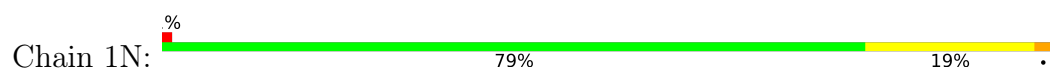
- Molecule 8: 50S ribosomal protein L9



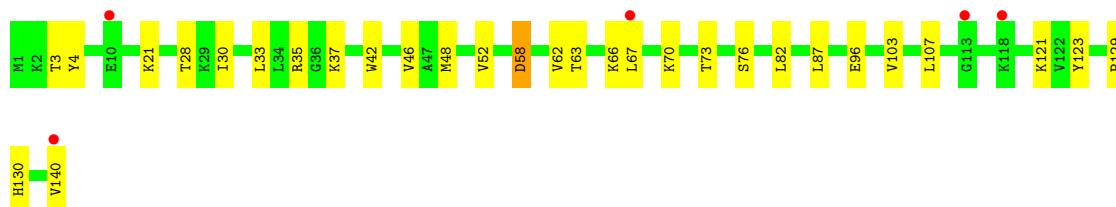
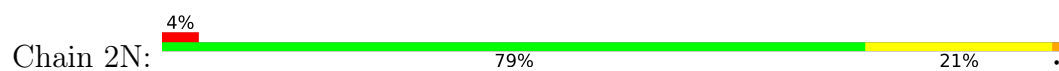
- Molecule 8: 50S ribosomal protein L9



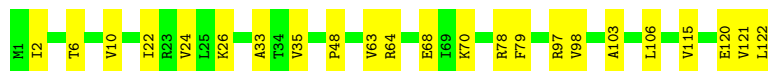
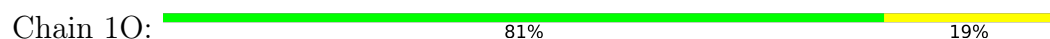
- Molecule 9: 50S ribosomal protein L13



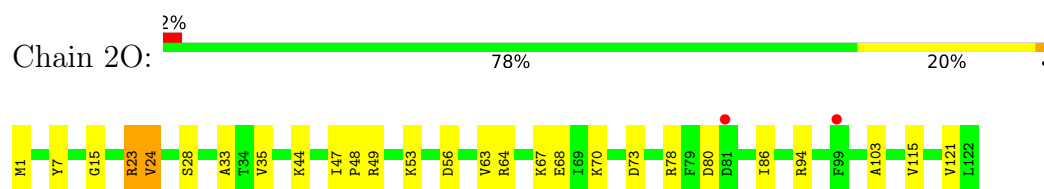
- Molecule 9: 50S ribosomal protein L13



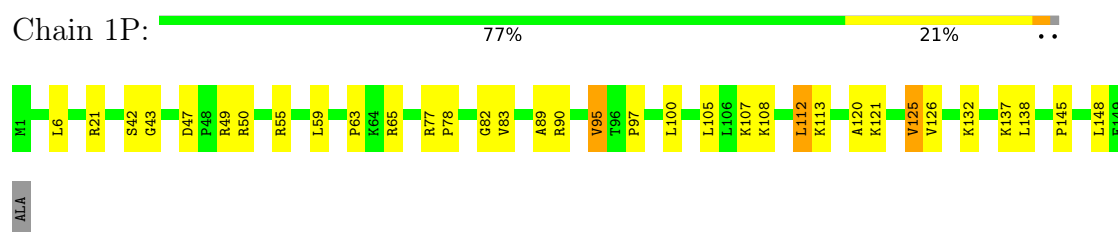
- Molecule 10: 50S ribosomal protein L14



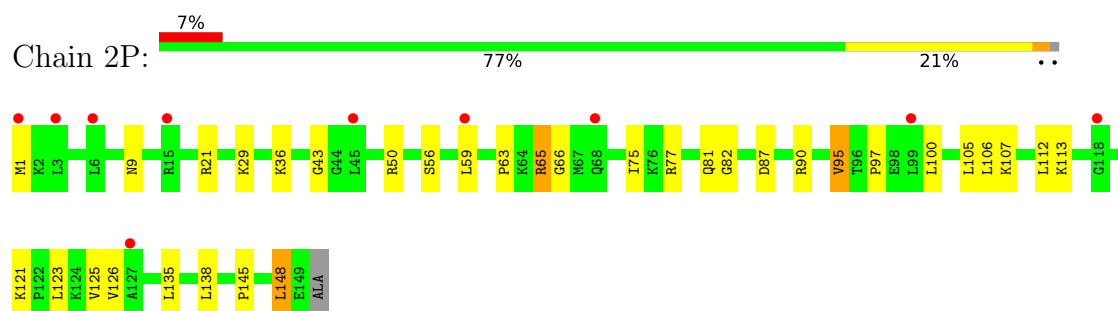
- Molecule 10: 50S ribosomal protein L14



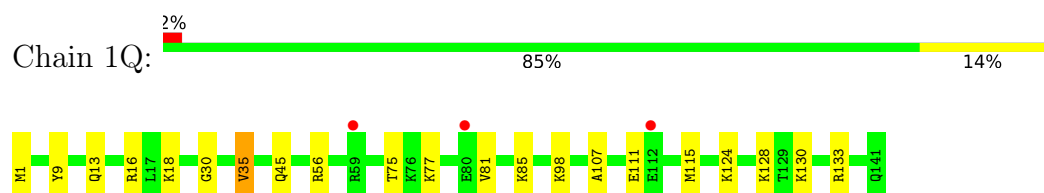
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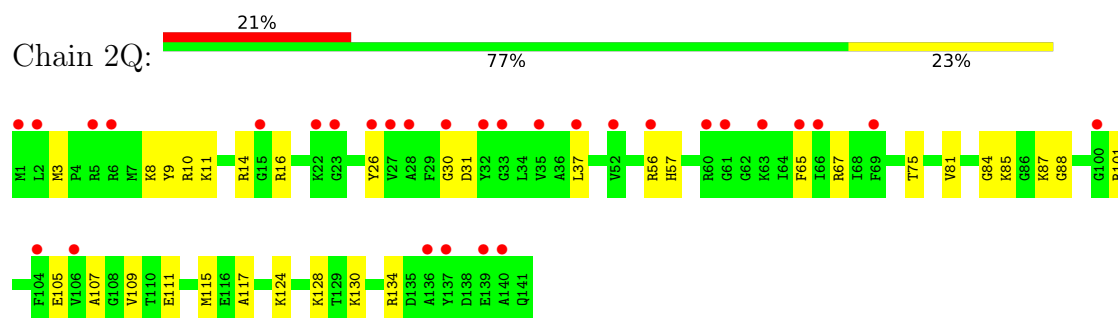
- Molecule 11: 50S ribosomal protein L15



- Molecule 12: 50S ribosomal protein L16



- Molecule 12: 50S ribosomal protein L16

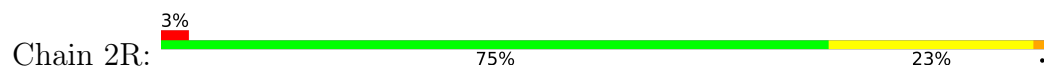


- Molecule 13: 50S ribosomal protein L17

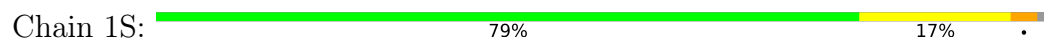




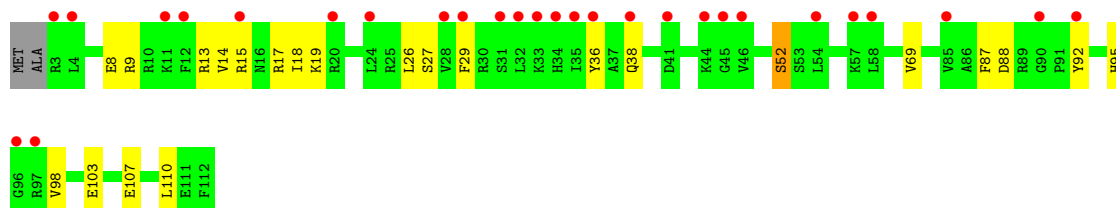
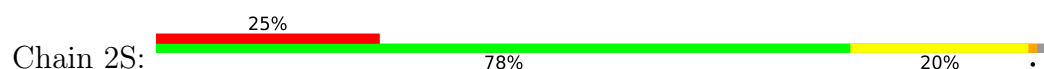
- Molecule 13: 50S ribosomal protein L17



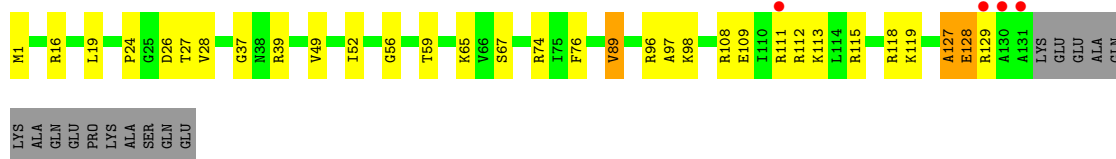
- Molecule 14: 50S ribosomal protein L18



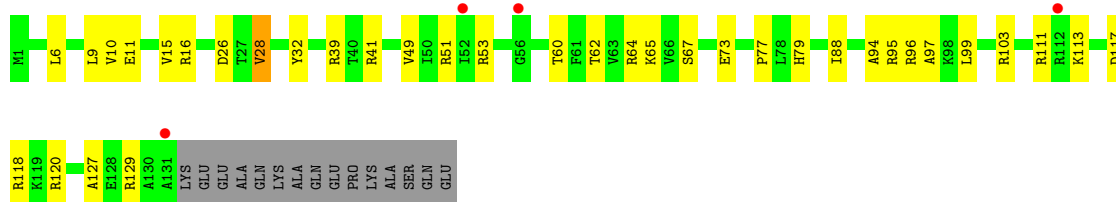
- Molecule 14: 50S ribosomal protein L18



- Molecule 15: 50S ribosomal protein L19



- Molecule 15: 50S ribosomal protein L19



- Molecule 16: 50S ribosomal protein L20

Chain 1U:  83% 14% ..



- Molecule 16: 50S ribosomal protein L20

Chain 2U:  2% 76% 21% ..



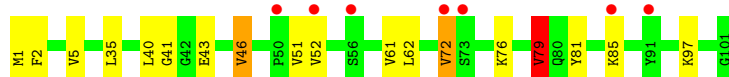
- Molecule 17: 50S ribosomal protein L21

Chain 1V:  79% 15% 5% ..



- Molecule 17: 50S ribosomal protein L21

Chain 2V:  7% 82% 15% ..



- Molecule 18: 50S ribosomal protein L22

Chain 1W:  % 77% 21% ..



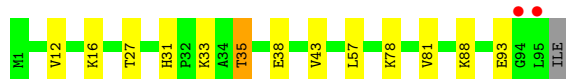
- Molecule 18: 50S ribosomal protein L22

Chain 2W:  % 79% 19% ..

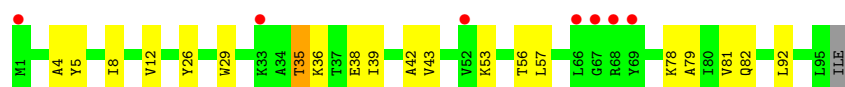
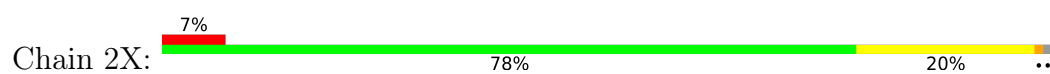


- Molecule 19: 50S ribosomal protein L23

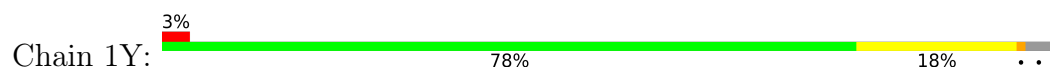
Chain 1X:  2% 85% 12% ..



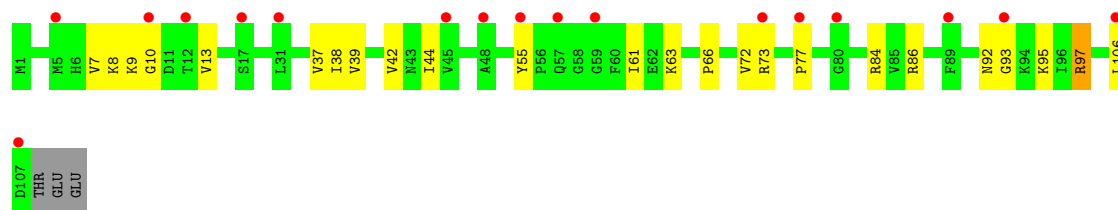
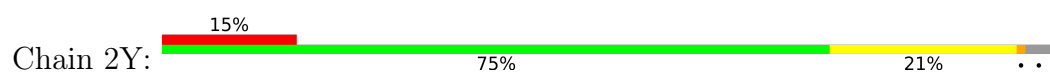
- Molecule 19: 50S ribosomal protein L23



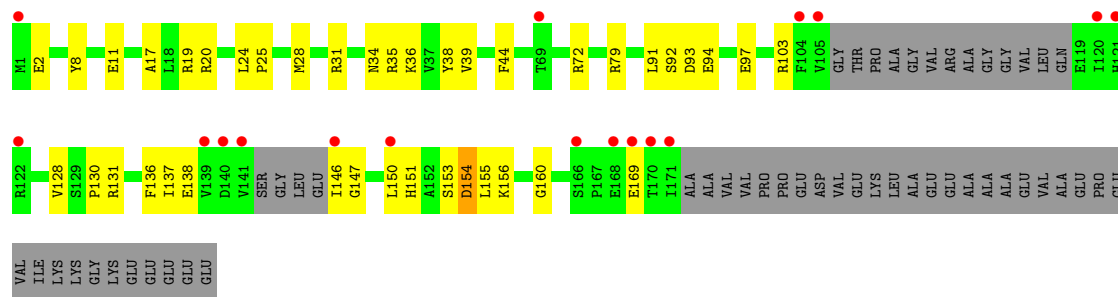
- Molecule 20: 50S ribosomal protein L24



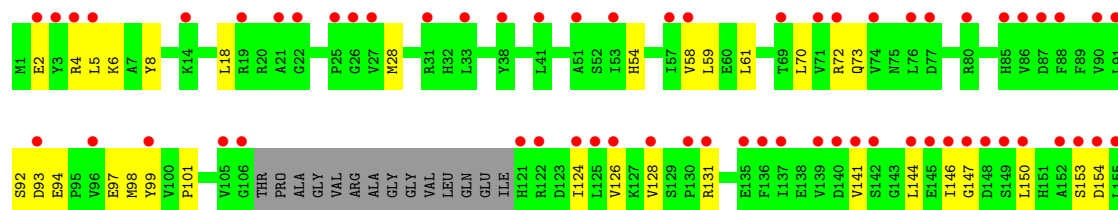
- Molecule 20: 50S ribosomal protein L24

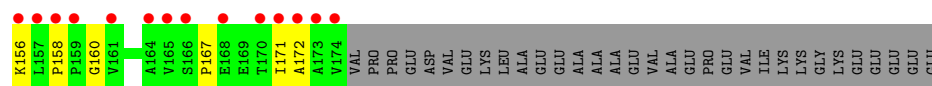


- Molecule 21: 50S ribosomal protein L25

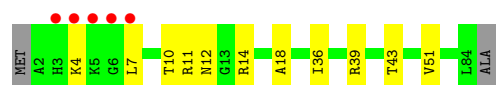
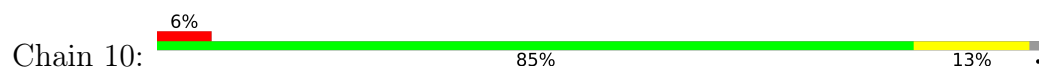


- Molecule 21: 50S ribosomal protein L25

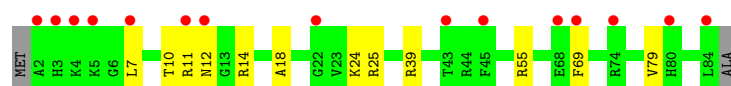
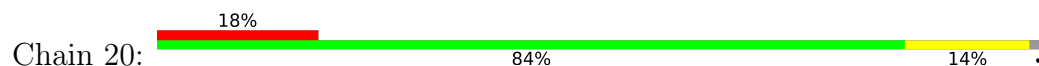




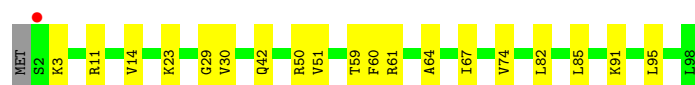
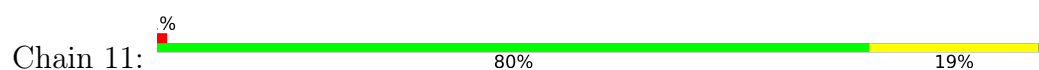
- Molecule 22: 50S ribosomal protein L27



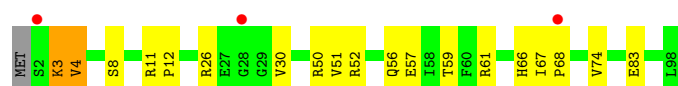
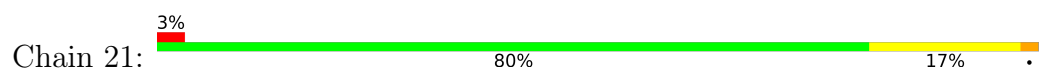
- Molecule 22: 50S ribosomal protein L27



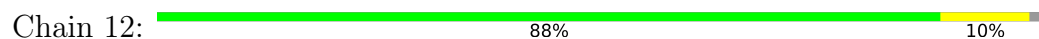
- Molecule 23: 50S ribosomal protein L28



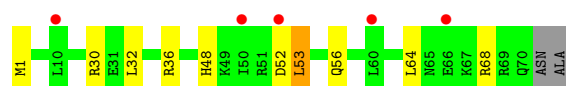
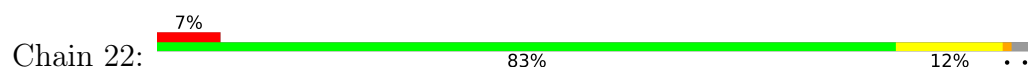
- Molecule 23: 50S ribosomal protein L28



- Molecule 24: 50S ribosomal protein L29



- Molecule 24: 50S ribosomal protein L29



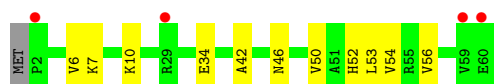
- Molecule 25: 50S ribosomal protein L30

Chain 13:  80% 15% . .



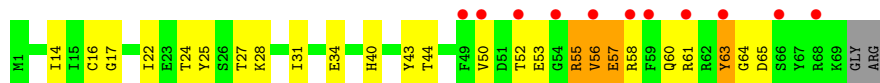
- Molecule 25: 50S ribosomal protein L30

Chain 23:  7% 80% 18% .



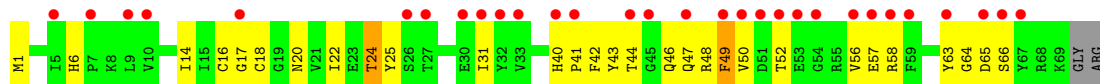
- Molecule 26: 50S ribosomal protein L31

Chain 14:  15% 62% 30% 6% .



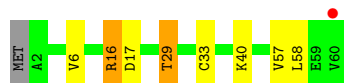
- Molecule 26: 50S ribosomal protein L31

Chain 24:  42% 56% 38% . .



- Molecule 27: 50S ribosomal protein L32

Chain 15:  2% 85% 10% . .



- Molecule 27: 50S ribosomal protein L32

Chain 25:  88% 10% .

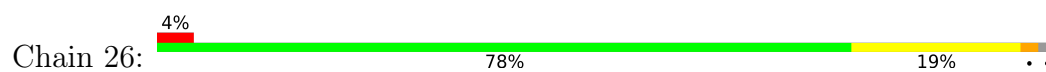


- Molecule 28: 50S ribosomal protein L33

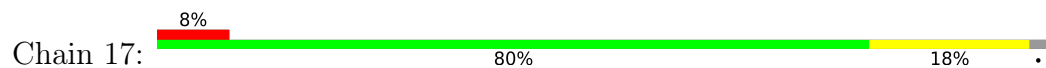
Chain 16:  72% 22% . .



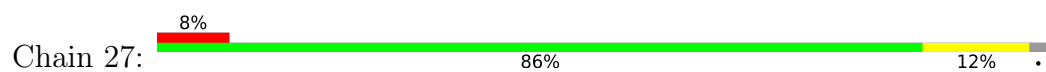
- Molecule 28: 50S ribosomal protein L33



- Molecule 29: 50S ribosomal protein L34



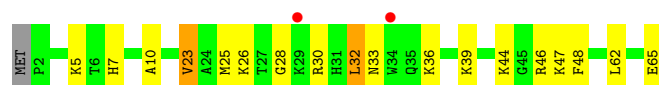
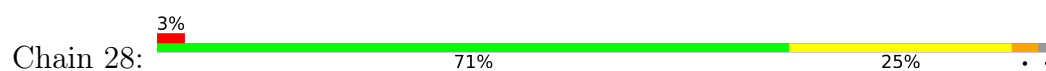
- Molecule 29: 50S ribosomal protein L34



- Molecule 30: 50S ribosomal protein L35



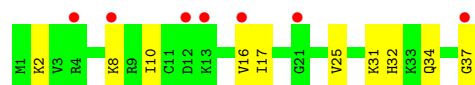
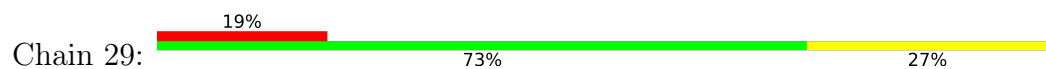
- Molecule 30: 50S ribosomal protein L35



- Molecule 31: 50S ribosomal protein L36

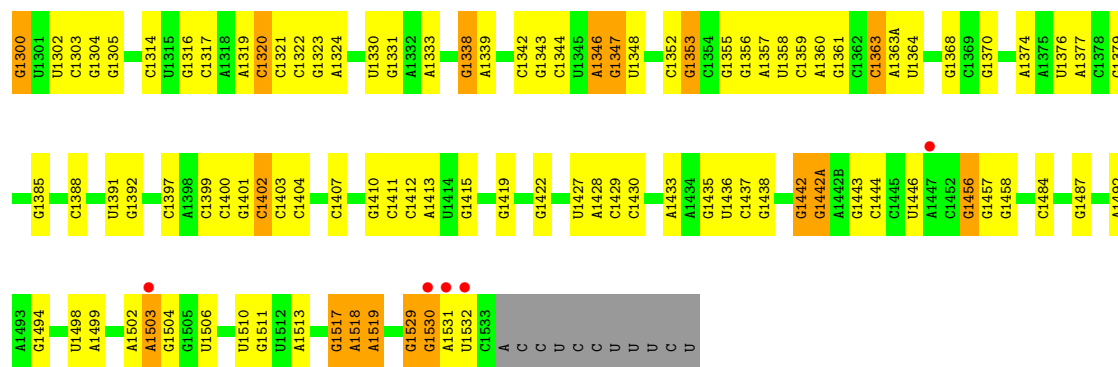


- Molecule 31: 50S ribosomal protein L36

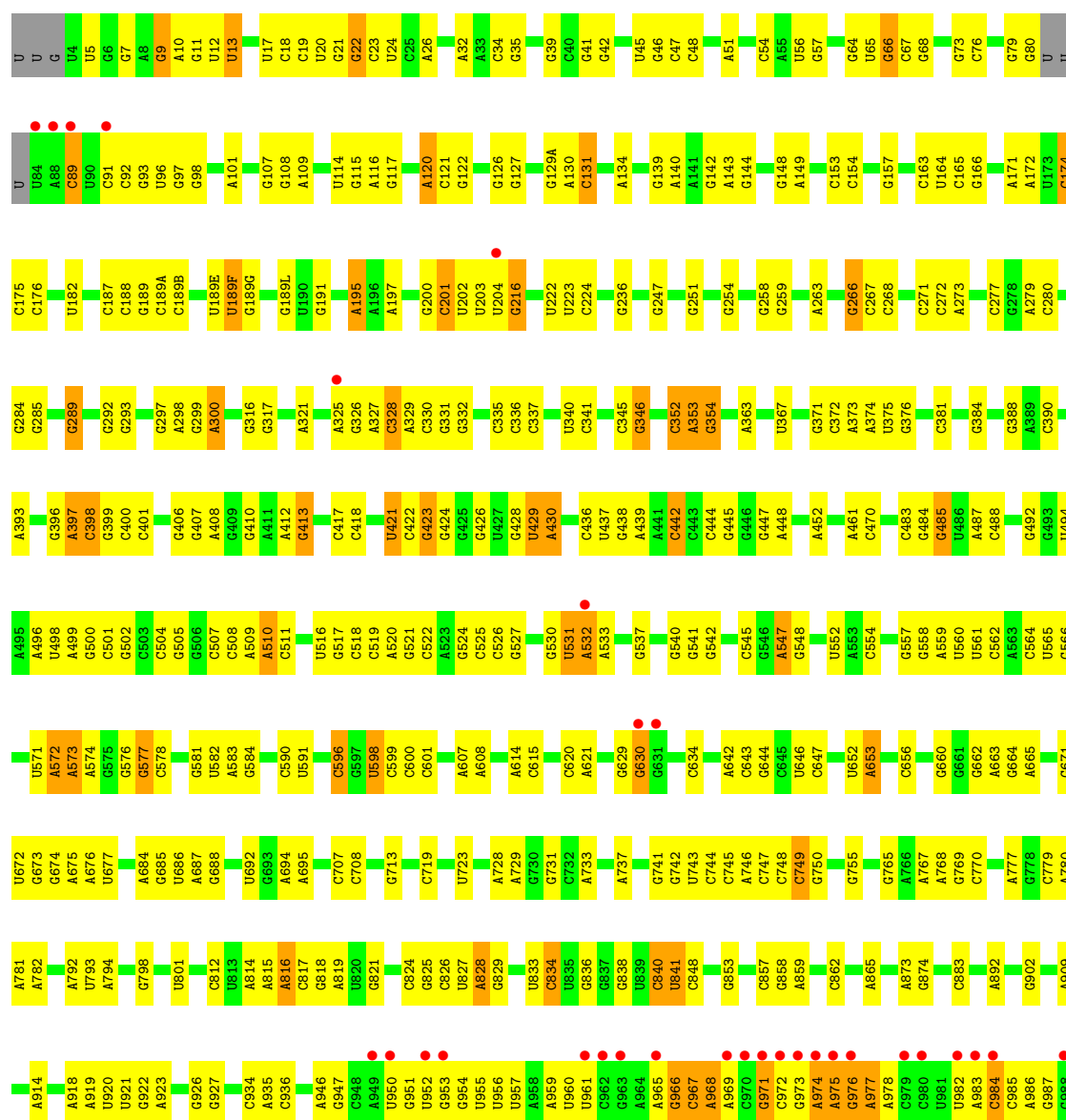


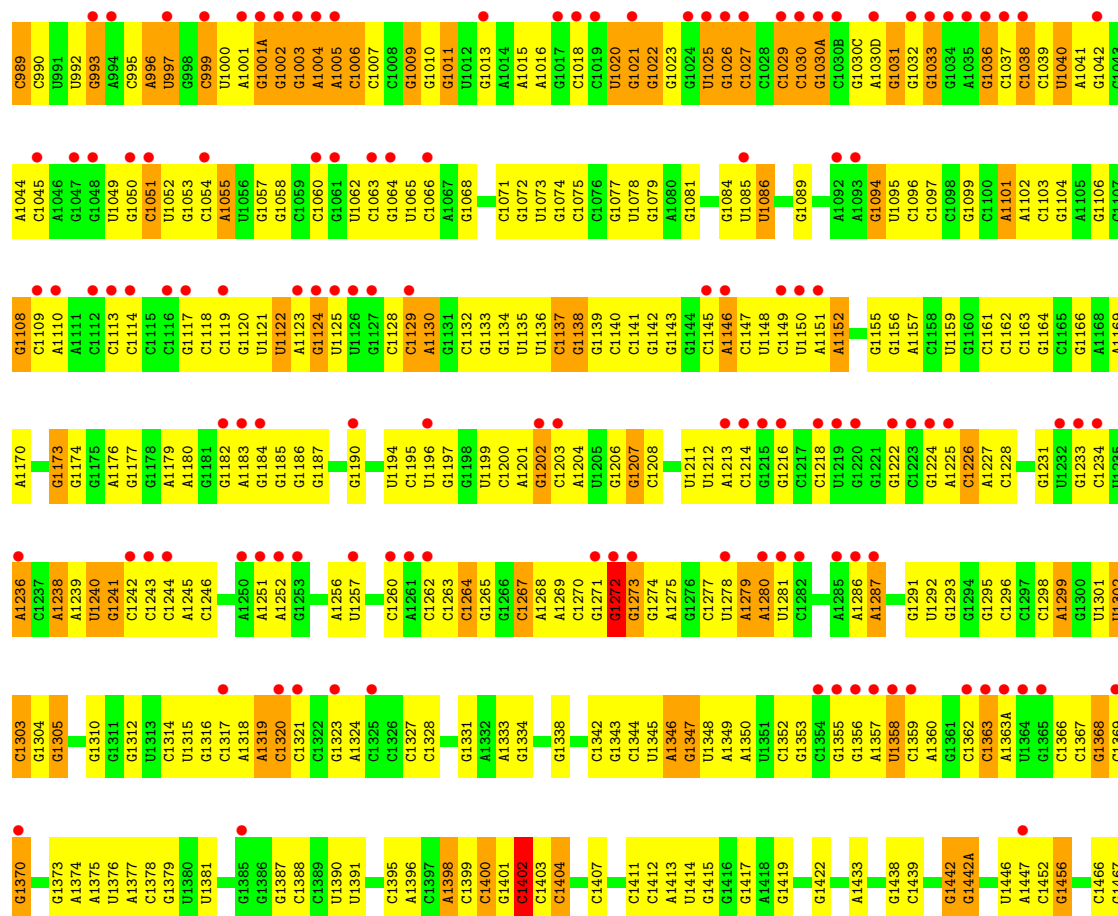
- Molecule 32: 16S Ribosomal RNA



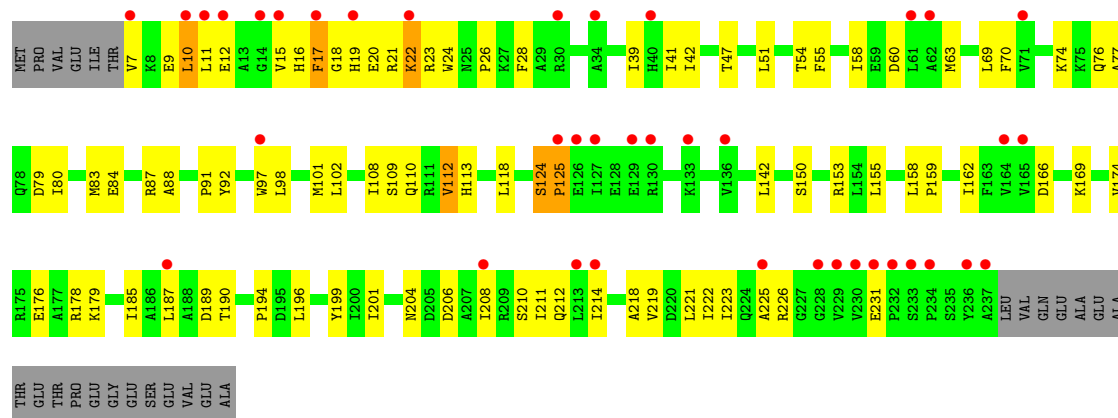


• Molecule 32: 16S Ribosomal RNA

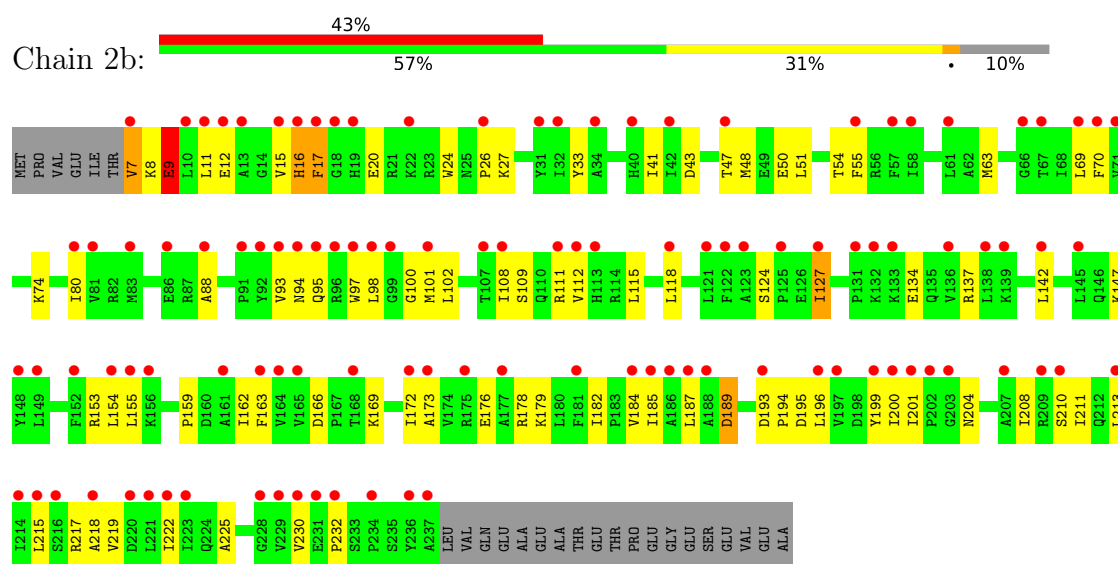




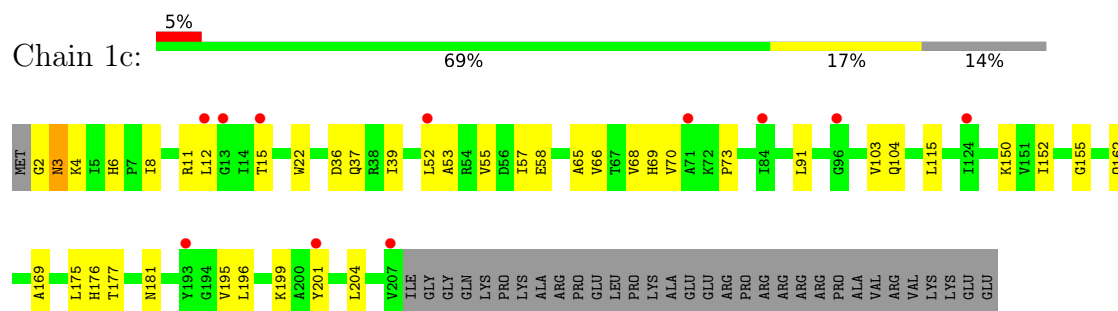
• Molecule 33: 30S ribosomal protein S2



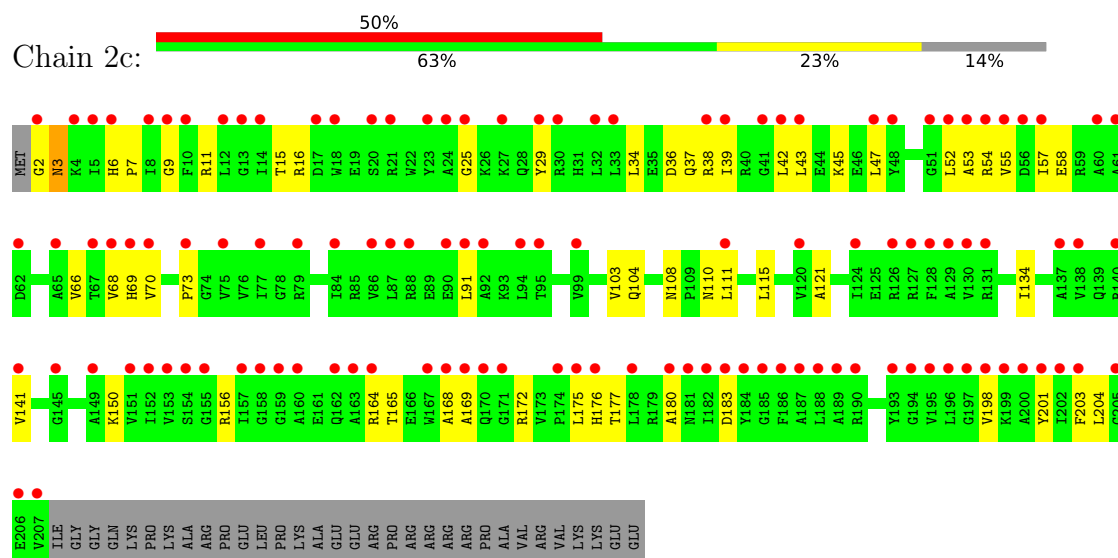
• Molecule 33: 30S ribosomal protein S2



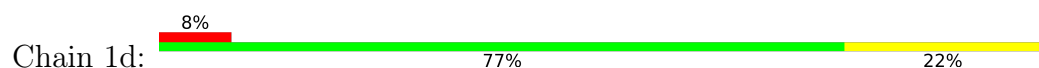
• Molecule 34: 30S ribosomal protein S3

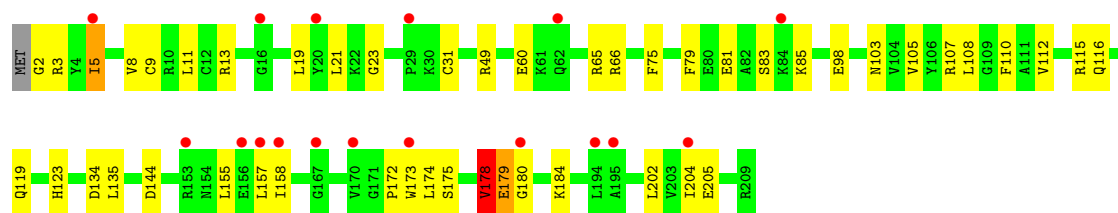


• Molecule 34: 30S ribosomal protein S3

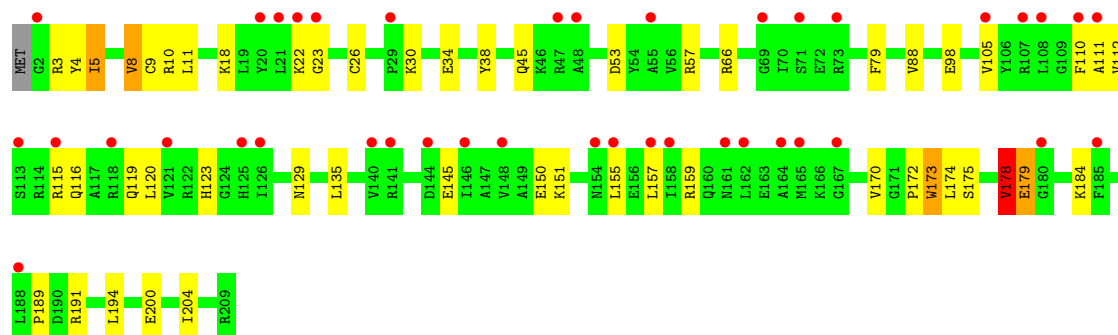
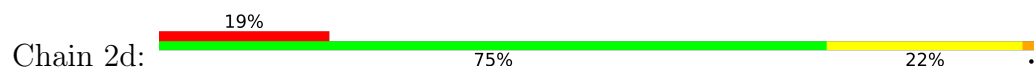


• Molecule 35: 30S ribosomal protein S4

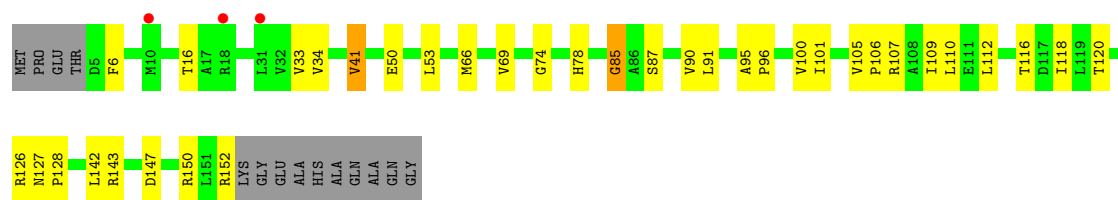




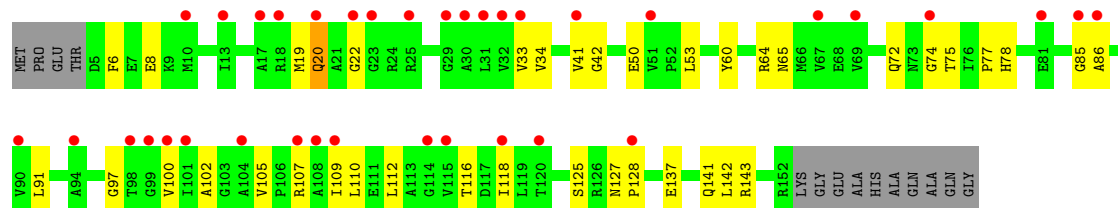
• Molecule 35: 30S ribosomal protein S4



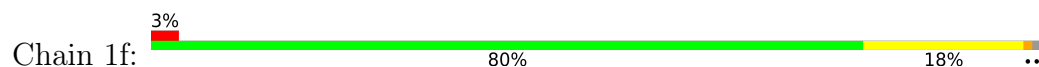
• Molecule 36: 30S ribosomal protein S5



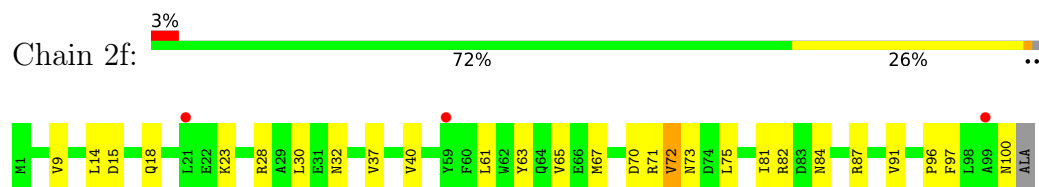
• Molecule 36: 30S ribosomal protein S5



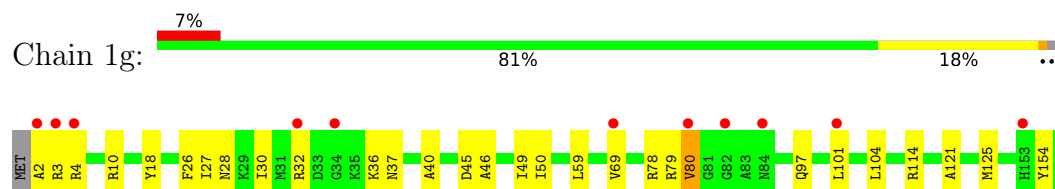
• Molecule 37: 30S ribosomal protein S6



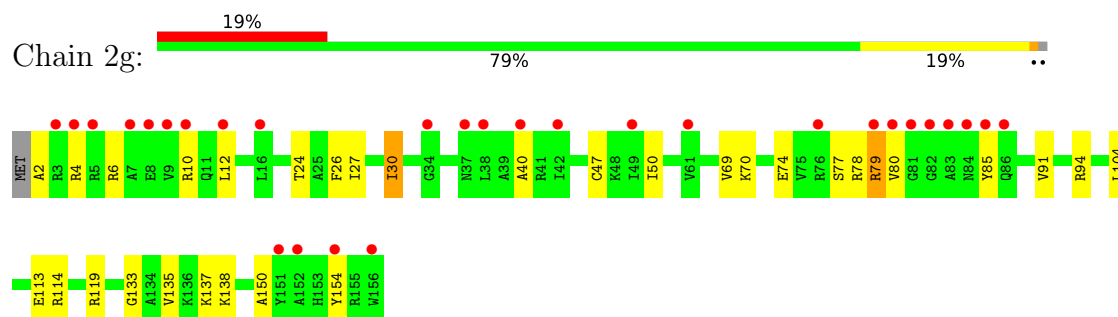
- Molecule 37: 30S ribosomal protein S6



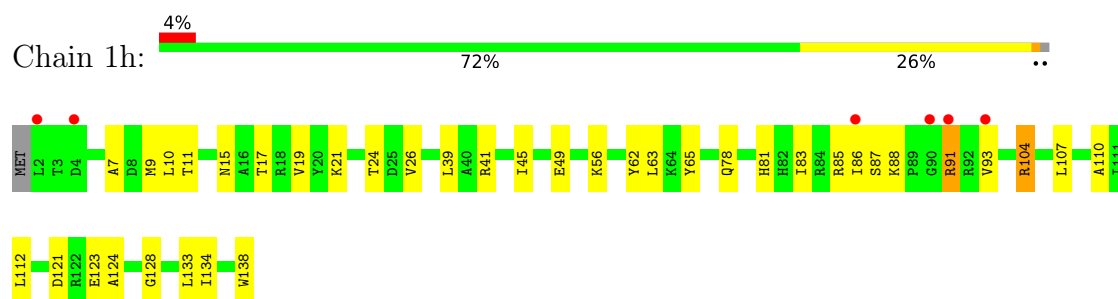
- Molecule 38: 30S ribosomal protein S7



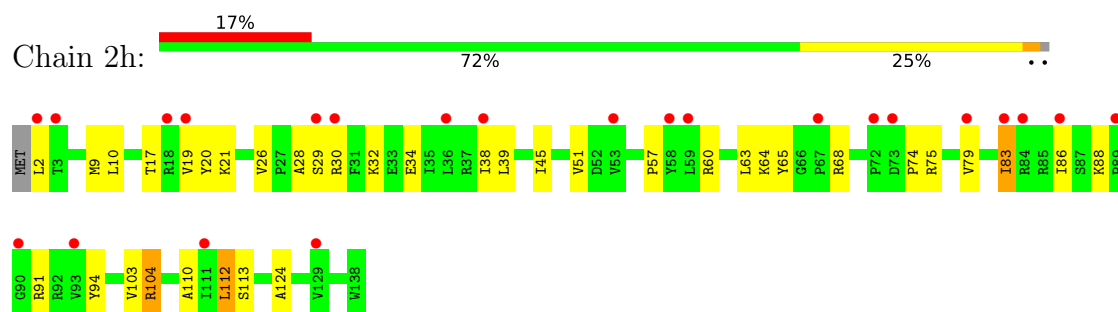
- Molecule 38: 30S ribosomal protein S7



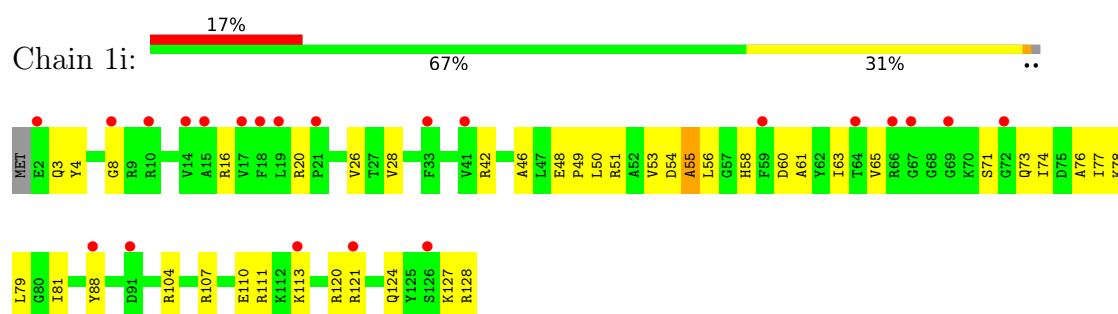
- Molecule 39: 30S ribosomal protein S8



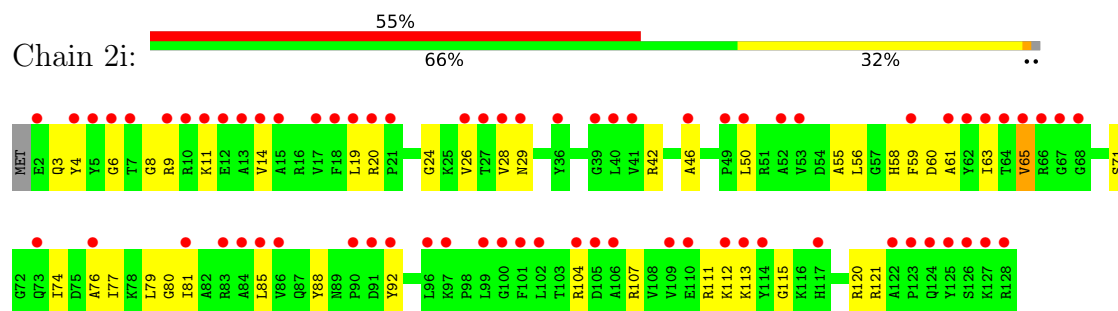
- Molecule 39: 30S ribosomal protein S8



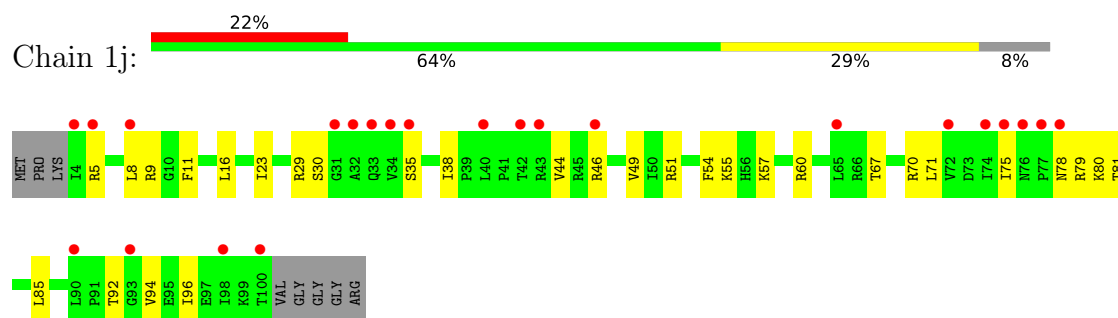
- Molecule 40: 30S ribosomal protein S9



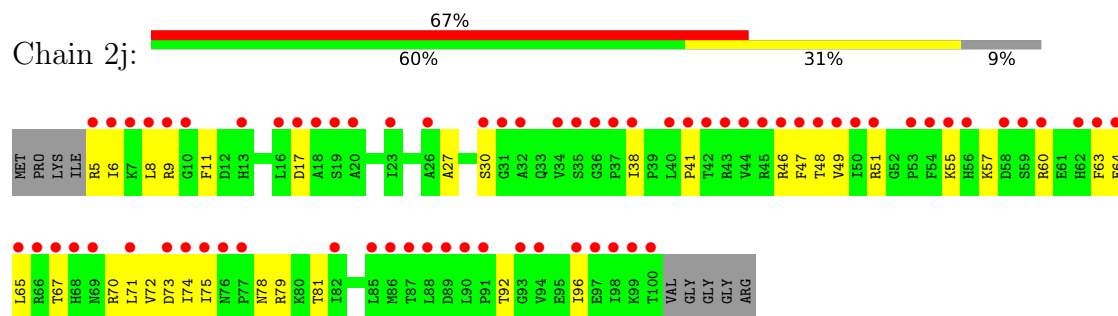
• Molecule 40: 30S ribosomal protein S9



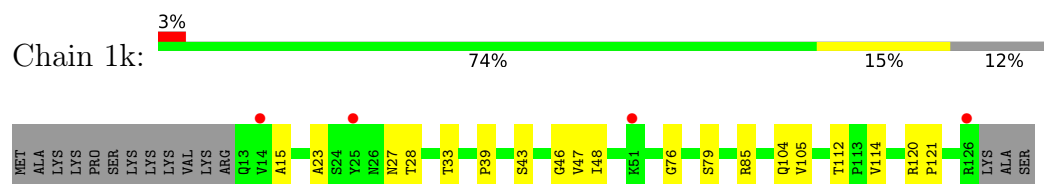
• Molecule 41: 30S ribosomal protein S10



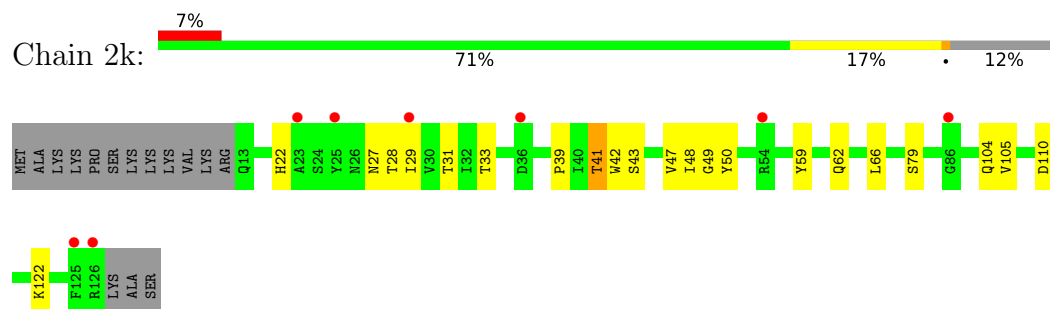
• Molecule 41: 30S ribosomal protein S10



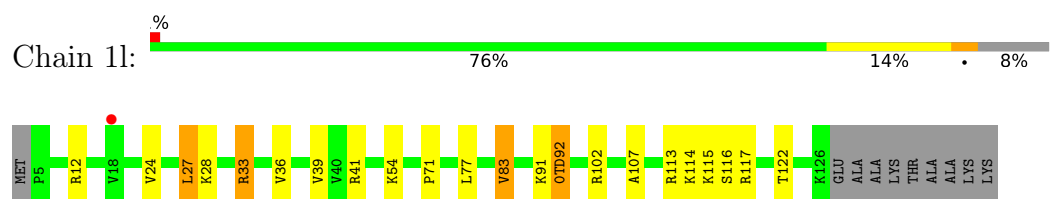
• Molecule 42: 30S ribosomal protein S11



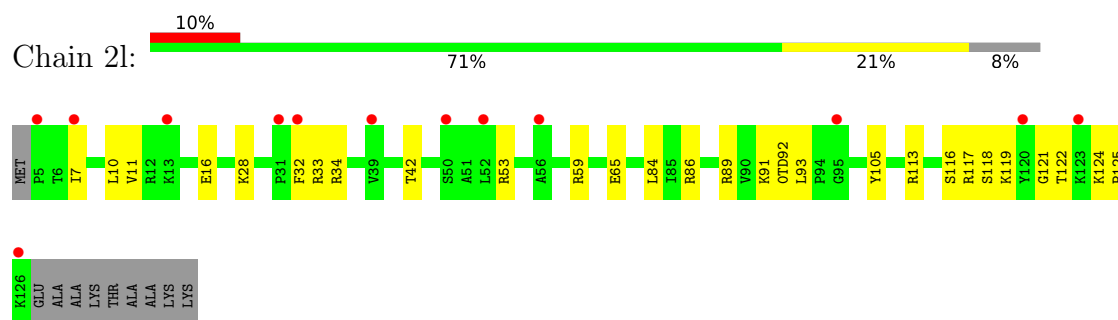
- Molecule 42: 30S ribosomal protein S11



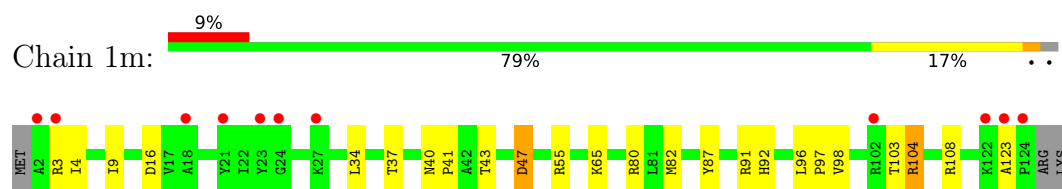
- Molecule 43: 30S ribosomal protein S12



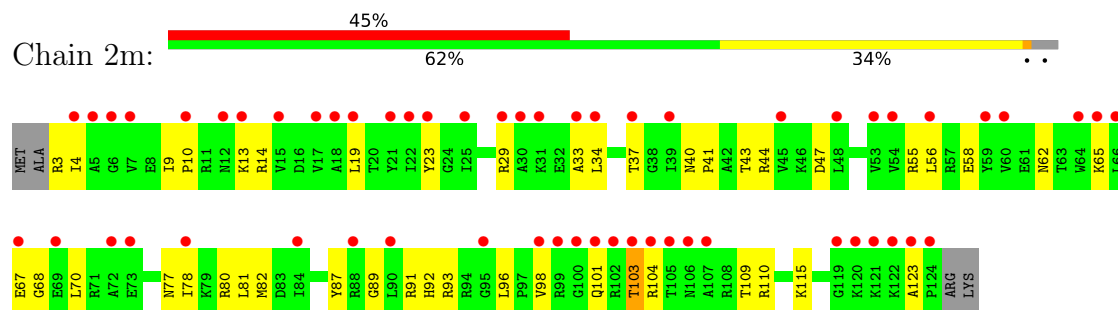
- Molecule 43: 30S ribosomal protein S12



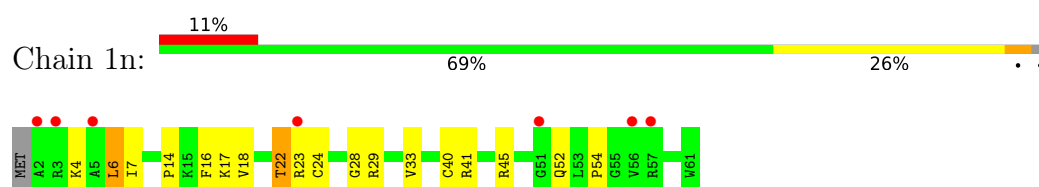
- Molecule 44: 30S ribosomal protein S13



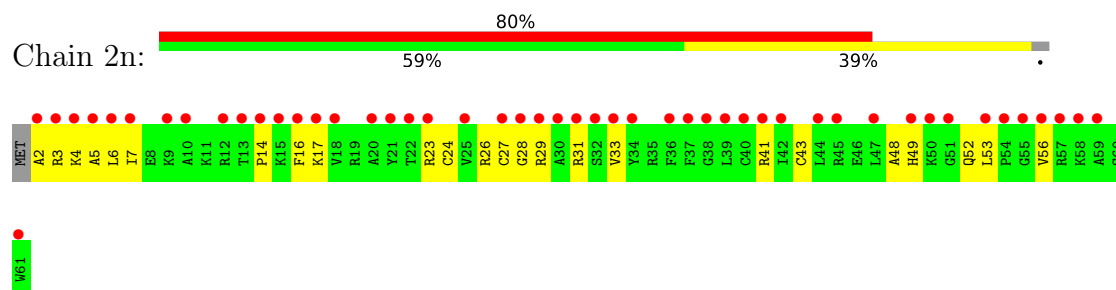
- Molecule 44: 30S ribosomal protein S13



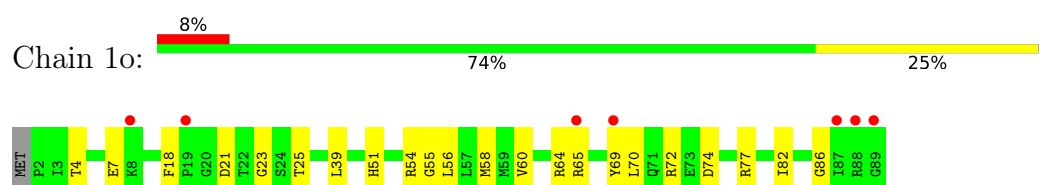
- Molecule 45: 30S ribosomal protein S14 type Z



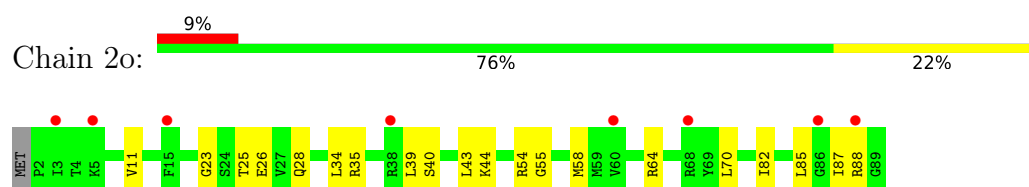
- Molecule 45: 30S ribosomal protein S14 type Z



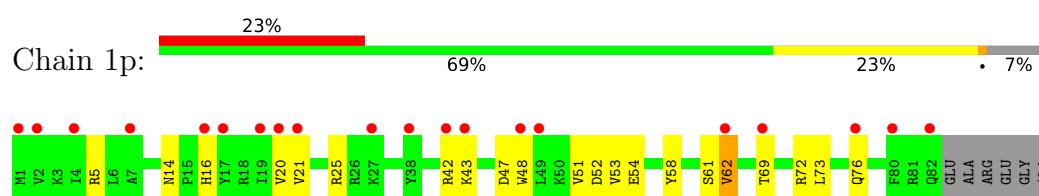
- Molecule 46: 30S ribosomal protein S15



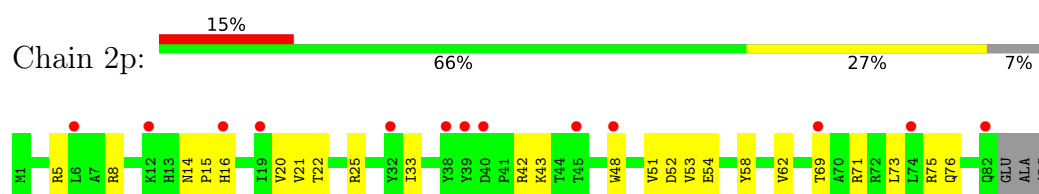
- Molecule 46: 30S ribosomal protein S15



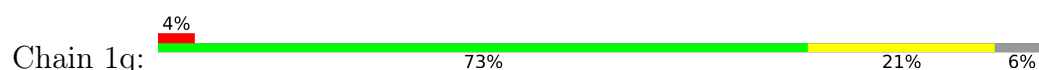
- Molecule 47: 30S ribosomal protein S16



- Molecule 47: 30S ribosomal protein S16

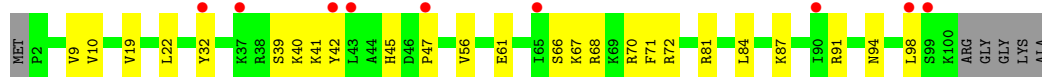
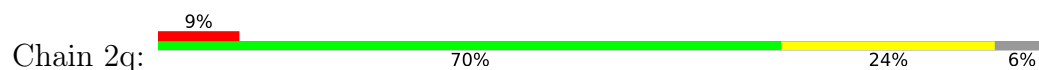


- Molecule 48: 30S ribosomal protein S17

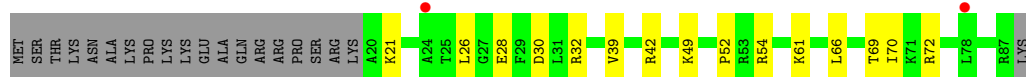




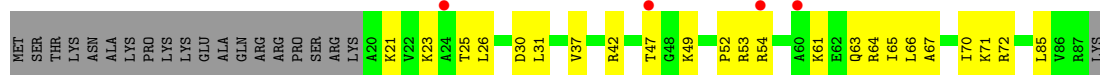
- Molecule 48: 30S ribosomal protein S17



- Molecule 49: 30S ribosomal protein S18



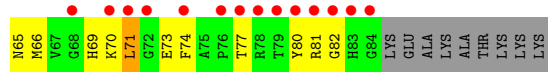
- Molecule 49: 30S ribosomal protein S18



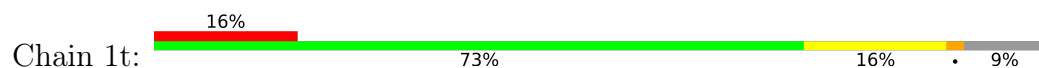
- Molecule 50: 30S ribosomal protein S19



- Molecule 50: 30S ribosomal protein S19

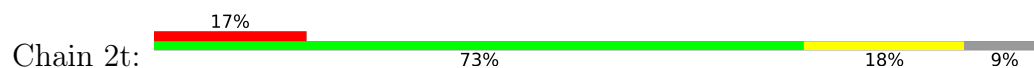


- Molecule 51: 30S ribosomal protein S20

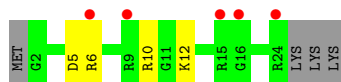




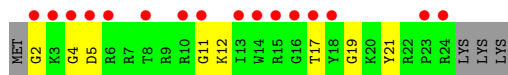
- Molecule 51: 30S ribosomal protein S20



- Molecule 52: 30S ribosomal protein Thx



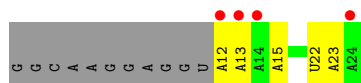
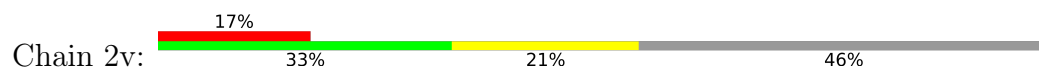
- Molecule 52: 30S ribosomal protein Thx



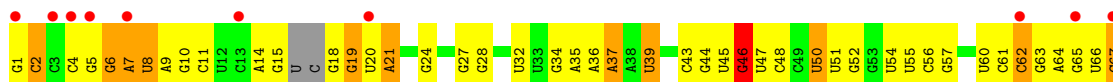
- Molecule 53: mRNA



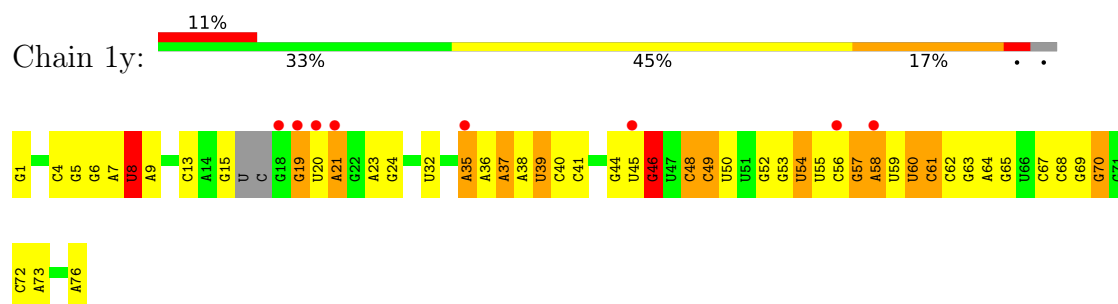
- Molecule 53: mRNA



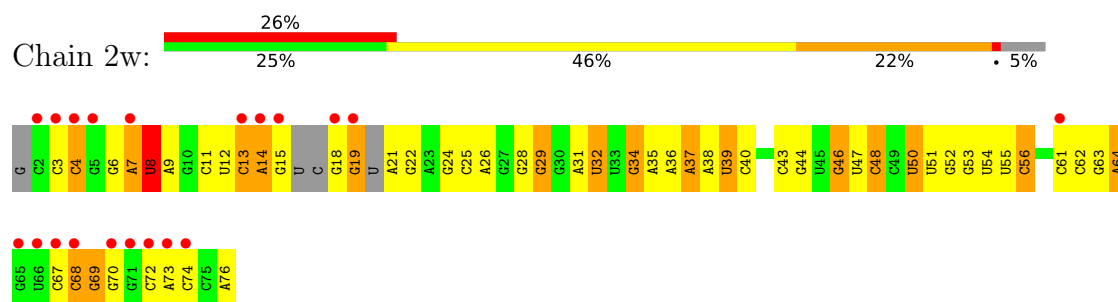
- Molecule 54: A-site and E-site tRNAs



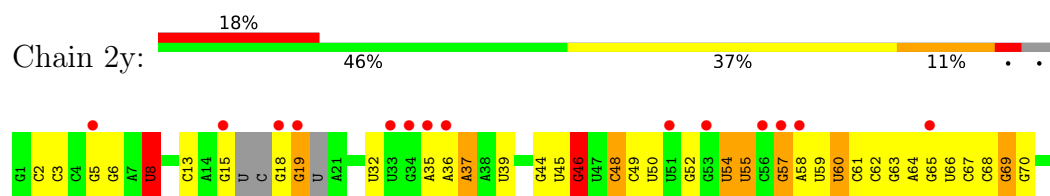
- Molecule 54: A-site and E-site tRNAs



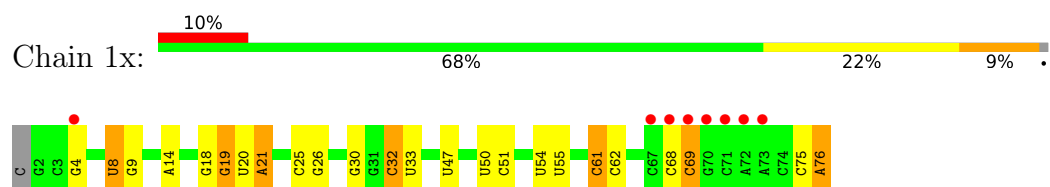
- Molecule 54: A-site and E-site tRNAs



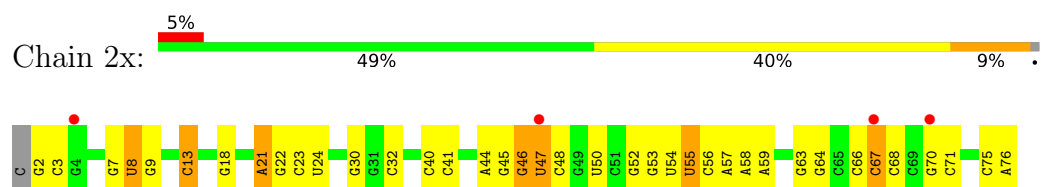
- Molecule 54: A-site and E-site tRNAs



- Molecule 55: P-site tRNA



- Molecule 55: 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	209.62Å 448.78Å 622.53Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	255.77 – 2.70 255.77 – 2.70	Depositor EDS
% Data completeness (in resolution range)	98.8 (255.77-2.70) 98.8 (255.77-2.70)	Depositor EDS
R_{merge}	0.14	Depositor
R_{sym}	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ ¹	1.21 (at 2.69Å)	Xtriage
Refinement program	PHENIX 1.8.2	Depositor
R, R_{free}	0.222 , 0.266 0.224 , 0.266	Depositor DCC
R_{free} test set	78357 reflections (4.96%)	wwPDB-VP
Wilson B-factor (Å ²)	52.2	Xtriage
Anisotropy	0.150	Xtriage
Bulk solvent k_{sol} (e/Å ³), B_{sol} (Å ²)	0.34 , 71.1	EDS
L-test for twinning ²	$\langle L \rangle = 0.41$, $\langle L^2 \rangle = 0.23$	Xtriage
Estimated twinning fraction	No twinning to report.	Xtriage
F_o, F_c correlation	0.89	EDS
Total number of atoms	301023	wwPDB-VP
Average B, all atoms (Å ²)	59.0	wwPDB-VP

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 1.50% 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: 4OC, UR3, 5MU, MIA, 5MC, MG, 2MG, MA6, 0TD, 4SU, PSU, OMG, ZN, 2MA, M2G, 7MG, SF4, K, MT9, 2MU

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	0.27	0/69009	0.45	1/107712 (0.0%)
1	2A	0.20	0/67293	0.39	1/105034 (0.0%)
2	1B	0.21	0/2882	0.39	0/4494
2	2B	0.19	0/2879	0.36	0/4487
3	1D	0.27	0/2186	0.52	0/2944
3	2D	0.22	0/2186	0.46	0/2944
4	1E	0.27	0/1592	0.55	0/2149
4	2E	0.21	0/1592	0.43	0/2149
5	1F	0.25	0/1619	0.47	0/2193
5	2F	0.18	0/1615	0.40	0/2188
6	1G	0.22	0/1448	0.44	0/1957
6	2G	0.18	0/1453	0.40	0/1963
7	1H	0.22	0/1356	0.41	0/1834
7	2H	0.18	0/1356	0.37	0/1834
8	1I	0.18	0/1112	0.40	0/1514
8	2I	0.16	0/1079	0.40	0/1475
9	1N	0.27	0/1144	0.46	0/1543
9	2N	0.20	0/1144	0.40	0/1543
10	1O	0.24	0/943	0.46	0/1269
10	2O	0.18	0/943	0.38	0/1269
11	1P	0.27	0/1152	0.47	0/1533
11	2P	0.21	0/1152	0.45	0/1533
12	1Q	0.27	0/1143	0.47	0/1527
12	2Q	0.19	0/1143	0.41	0/1527
13	1R	0.26	0/982	0.50	0/1312
13	2R	0.20	0/982	0.44	0/1312
14	1S	0.21	0/883	0.47	0/1176
14	2S	0.19	0/880	0.43	0/1172
15	1T	0.25	0/1105	0.48	0/1477
15	2T	0.19	0/1097	0.43	0/1468
16	1U	0.29	0/977	0.49	0/1301

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
16	2U	0.20	0/977	0.40	0/1301
17	1V	0.32	1/782 (0.1%)	0.46	0/1049
17	2V	0.18	0/782	0.41	0/1049
18	1W	0.27	0/897	0.47	0/1205
18	2W	0.21	0/897	0.41	0/1205
19	1X	0.26	0/764	0.52	0/1025
19	2X	0.19	0/764	0.43	0/1025
20	1Y	0.24	0/819	0.53	0/1095
20	2Y	0.20	0/819	0.41	0/1095
21	1Z	0.21	0/1267	0.46	0/1717
21	2Z	0.20	0/1299	0.44	0/1763
22	10	0.25	0/662	0.54	0/881
22	20	0.19	0/662	0.45	0/881
23	11	0.25	0/762	0.44	0/1014
23	21	0.20	0/762	0.40	0/1014
24	12	0.24	0/590	0.40	0/781
24	22	0.17	0/590	0.34	0/781
25	13	0.25	0/474	0.46	0/635
25	23	0.18	0/469	0.43	0/630
26	14	0.29	0/565	0.70	2/761 (0.3%)
26	24	0.23	0/545	0.57	0/737
27	15	0.26	0/469	0.52	0/635
27	25	0.21	0/469	0.41	0/635
28	16	0.25	0/460	0.46	0/613
28	26	0.19	0/456	0.44	0/608
29	17	0.27	0/426	0.53	0/561
29	27	0.25	0/426	0.52	0/561
30	18	0.26	0/525	0.48	0/691
30	28	0.19	0/525	0.39	0/691
31	19	0.25	0/310	0.51	0/407
31	29	0.18	0/310	0.40	0/407
32	1a	0.20	0/35795	0.38	0/55864
32	2a	0.19	0/35886	0.36	1/56005 (0.0%)
33	1b	0.20	0/1881	0.48	0/2542
33	2b	0.20	0/1860	0.44	0/2518
34	1c	0.18	0/1572	0.38	0/2126
34	2c	0.18	0/1566	0.38	0/2119
35	1d	0.19	0/1685	0.43	0/2262
35	2d	0.18	0/1704	0.43	0/2284
36	1e	0.20	0/1145	0.44	0/1543
36	2e	0.19	0/1149	0.45	0/1548
37	1f	0.19	0/823	0.37	0/1115
37	2f	0.18	0/829	0.38	0/1123

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
38	1g	0.18	0/1250	0.38	0/1679
38	2g	0.18	0/1254	0.38	0/1683
39	1h	0.19	0/1108	0.39	0/1494
39	2h	0.17	0/1108	0.39	0/1494
40	1i	0.18	0/1002	0.48	0/1346
40	2i	0.19	0/997	0.45	0/1343
41	1j	0.18	0/722	0.43	0/982
41	2j	0.19	0/727	0.43	0/988
42	1k	0.17	0/844	0.38	0/1145
42	2k	0.16	0/848	0.37	0/1149
43	1l	0.20	0/937	0.41	0/1260
43	2l	0.20	0/937	0.45	0/1260
44	1m	0.19	0/969	0.42	0/1302
44	2m	0.19	0/961	0.41	0/1291
45	1n	0.21	0/501	0.45	0/664
45	2n	0.18	0/501	0.41	0/664
46	1o	0.18	0/739	0.39	0/985
46	2o	0.16	0/739	0.37	0/985
47	1p	0.18	0/697	0.45	0/939
47	2p	0.18	0/693	0.44	0/935
48	1q	0.18	0/836	0.38	0/1117
48	2q	0.17	0/836	0.37	0/1117
49	1r	0.20	0/560	0.42	0/746
49	2r	0.16	0/560	0.38	0/746
50	1s	0.19	0/667	0.48	0/900
50	2s	0.23	0/661	0.49	0/893
51	1t	0.19	0/730	0.46	0/965
51	2t	0.18	0/729	0.46	0/965
52	1u	0.19	0/203	0.43	0/266
52	2u	0.19	0/203	0.45	0/266
53	1v	0.18	0/310	0.31	0/480
53	2v	0.21	0/310	0.32	0/480
54	1w	0.26	1/1606 (0.1%)	0.40	0/2497
54	1y	0.25	1/1606 (0.1%)	0.40	0/2497
54	2w	0.26	1/1556 (0.1%)	0.37	0/2418
54	2y	0.25	0/1583	0.39	0/2459
55	1x	0.26	0/1725	0.41	0/2689
55	2x	0.24	1/1725 (0.1%)	0.37	0/2689
All	All	0.22	5/316686 (0.0%)	0.41	5/474113 (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
6	1G	0	1
26	14	0	1
26	24	0	2
33	2b	0	1
All	All	0	5

All (5) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
17	1V	34	GLU	C-N	-5.45	1.22	1.33
54	1w	8	4SU	O3'-P	5.32	1.61	1.56
55	2x	8	4SU	O3'-P	5.29	1.61	1.56
54	1y	8	4SU	O3'-P	5.17	1.61	1.56
54	2w	8	4SU	O3'-P	5.03	1.61	1.56

All (5) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
26	14	63	TYR	N-CA-C	-7.13	102.11	111.02
26	14	64	GLY	N-CA-C	5.93	127.23	113.18
1	1A	2014	G	C2'-C3'-O3'	5.50	117.75	109.50
1	2A	1992	G	C2'-C3'-O3'	5.36	117.54	109.50
32	2a	1272	G	N1-C2-N2	-5.11	100.88	116.20

There are no chirality outliers.

All (5) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
26	14	63	TYR	Peptide
6	1G	126	ASP	Peptide
26	24	56	VAL	Peptide
26	24	63	TYR	Peptide
33	2b	9	GLU	Peptide

5.2 Too-close contacts

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	1A	61852	0	31193	596	0
1	2A	60322	0	30421	698	0
2	1B	2577	0	1305	31	0
2	2B	2575	0	1303	35	0
3	1D	2136	0	2218	51	0
3	2D	2136	0	2218	47	0
4	1E	1559	0	1618	36	0
4	2E	1559	0	1618	29	0
5	1F	1584	0	1625	34	0
5	2F	1580	0	1619	37	0
6	1G	1423	0	1436	40	0
6	2G	1428	0	1438	42	0
7	1H	1330	0	1407	31	0
7	2H	1330	0	1407	22	0
8	1I	1097	0	1140	21	0
8	2I	1064	0	1082	16	0
9	1N	1117	0	1184	22	0
9	2N	1117	0	1184	22	0
10	1O	933	0	996	11	0
10	2O	933	0	996	18	0
11	1P	1135	0	1212	28	0
11	2P	1135	0	1212	26	0
12	1Q	1122	0	1179	18	0
12	2Q	1122	0	1179	27	0
13	1R	968	0	1033	20	0
13	2R	968	0	1033	19	0
14	1S	873	0	927	14	0
14	2S	870	0	923	15	0
15	1T	1091	0	1151	24	0
15	2T	1083	0	1136	28	0
16	1U	959	0	1019	15	0
16	2U	959	0	1019	22	0
17	1V	771	0	830	13	0
17	2V	771	0	830	11	0
18	1W	886	0	940	12	0
18	2W	886	0	940	15	0
19	1X	750	0	814	9	0
19	2X	750	0	814	14	0
20	1Y	806	0	881	18	0
20	2Y	806	0	881	18	0
21	1Z	1240	0	1240	26	0
21	2Z	1271	0	1273	22	0
22	10	653	0	674	15	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
22	20	653	0	674	13	0
23	11	755	0	826	12	0
23	21	755	0	826	17	0
24	12	588	0	643	3	0
24	22	588	0	643	9	0
25	13	469	0	518	5	0
25	23	464	0	514	7	0
26	14	552	0	533	17	0
26	24	532	0	503	19	0
27	15	455	0	465	5	0
27	25	455	0	465	5	0
28	16	453	0	473	9	0
28	26	449	0	469	9	0
29	17	418	0	467	5	0
29	27	418	0	467	5	0
30	18	517	0	582	26	0
30	28	517	0	582	16	0
31	19	307	0	335	0	0
31	29	307	0	335	6	0
32	1a	32246	0	16295	439	0
32	2a	32327	0	16339	584	0
33	1b	1846	0	1867	62	0
33	2b	1825	0	1828	61	0
34	1c	1548	0	1535	30	0
34	2c	1542	0	1517	36	0
35	1d	1655	0	1672	34	0
35	2d	1674	0	1714	35	0
36	1e	1129	0	1185	30	0
36	2e	1133	0	1191	27	0
37	1f	810	0	804	12	0
37	2f	816	0	808	20	0
38	1g	1231	0	1238	19	0
38	2g	1235	0	1249	21	0
39	1h	1088	0	1126	25	0
39	2h	1088	0	1126	27	0
40	1i	983	0	986	32	0
40	2i	978	0	966	30	0
41	1j	709	0	650	22	0
41	2j	714	0	672	27	0
42	1k	829	0	825	9	0
42	2k	833	0	836	16	0
43	1l	932	0	981	19	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
43	2l	932	0	981	25	0
44	1m	958	0	1002	20	0
44	2m	950	0	988	39	0
45	1n	492	0	529	16	0
45	2n	492	0	529	23	0
46	1o	728	0	760	16	0
46	2o	728	0	760	13	0
47	1p	681	0	697	16	0
47	2p	677	0	686	16	0
48	1q	823	0	891	16	0
48	2q	823	0	891	18	0
49	1r	555	0	618	12	0
49	2r	555	0	618	22	0
50	1s	652	0	662	19	0
50	2s	646	0	644	25	0
51	1t	728	0	798	14	0
51	2t	727	0	796	11	0
52	1u	199	0	208	3	0
52	2u	199	0	208	8	0
53	1v	277	0	140	2	0
53	2v	277	0	140	2	0
54	1w	1592	0	819	39	0
54	1y	1585	0	804	28	0
54	2w	1544	0	788	32	0
54	2y	1565	0	795	25	0
55	1x	1625	0	828	12	0
55	2x	1625	0	828	21	0
56	10	9	0	0	0	0
56	11	4	0	0	0	0
56	12	2	0	0	0	0
56	13	3	0	0	0	0
56	15	2	0	0	0	0
56	16	3	0	0	0	0
56	17	3	0	0	0	0
56	18	3	0	0	0	0
56	19	1	0	0	0	0
56	1A	1146	0	0	0	0
56	1B	36	0	0	0	0
56	1D	12	0	0	0	0
56	1E	12	0	0	0	0
56	1F	8	0	0	0	0
56	1G	5	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
56	1H	1	0	0	0	0
56	1I	1	0	0	0	0
56	1N	6	0	0	0	0
56	1O	8	0	0	0	0
56	1P	4	0	0	0	0
56	1Q	5	0	0	0	0
56	1R	2	0	0	0	0
56	1S	3	0	0	0	0
56	1T	1	0	0	0	0
56	1U	6	0	0	0	0
56	1V	3	0	0	0	0
56	1W	7	0	0	0	0
56	1X	5	0	0	0	0
56	1Y	3	0	0	0	0
56	1Z	4	0	0	0	0
56	1a	229	0	0	0	0
56	1b	2	0	0	0	0
56	1d	1	0	0	0	0
56	1e	2	0	0	0	0
56	1f	2	0	0	0	0
56	1h	1	0	0	0	0
56	1l	3	0	0	0	0
56	1m	1	0	0	0	0
56	1n	1	0	0	0	0
56	1o	1	0	0	0	0
56	1q	1	0	0	0	0
56	1r	1	0	0	0	0
56	1s	1	0	0	0	0
56	1t	1	0	0	0	0
56	1v	1	0	0	0	0
56	1w	11	0	0	0	0
56	1x	16	0	0	0	0
56	1y	4	0	0	0	0
56	20	3	0	0	0	0
56	23	3	0	0	0	0
56	25	4	0	0	0	0
56	26	1	0	0	0	0
56	27	1	0	0	0	0
56	28	1	0	0	0	0
56	2A	903	0	0	0	0
56	2B	21	0	0	0	0
56	2D	4	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
56	2E	8	0	0	0	0
56	2F	4	0	0	0	0
56	2G	1	0	0	0	0
56	2N	1	0	0	0	0
56	2O	1	0	0	0	0
56	2Q	4	0	0	0	0
56	2R	2	0	0	0	0
56	2T	2	0	0	0	0
56	2U	5	0	0	0	0
56	2V	2	0	0	0	0
56	2W	3	0	0	0	0
56	2X	2	0	0	0	0
56	2Y	1	0	0	0	0
56	2Z	1	0	0	0	0
56	2a	244	0	0	0	0
56	2d	1	0	0	0	0
56	2e	1	0	0	0	0
56	2f	2	0	0	0	0
56	2g	1	0	0	0	0
56	2j	2	0	0	0	0
56	2l	3	0	0	0	0
56	2q	3	0	0	0	0
56	2r	1	0	0	0	0
56	2t	1	0	0	0	0
56	2v	3	0	0	0	0
56	2w	9	0	0	0	0
56	2x	6	0	0	0	0
56	2y	7	0	0	0	0
57	1A	33	0	43	1	0
57	2A	33	0	43	1	0
58	1A	1	0	0	0	0
58	2A	1	0	0	0	0
59	14	1	0	0	0	0
59	15	1	0	0	0	0
59	16	1	0	0	0	0
59	19	1	0	0	0	0
59	1Y	1	0	0	0	0
59	1n	1	0	0	0	0
59	24	1	0	0	0	0
59	25	1	0	0	0	0
59	26	1	0	0	0	0
59	29	1	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
59	2Y	1	0	0	0	0
59	2n	1	0	0	0	0
60	1d	8	0	0	0	0
60	2d	8	0	0	0	0
61	10	11	0	0	1	0
61	11	12	0	0	0	0
61	12	3	0	0	0	0
61	13	5	0	0	1	0
61	14	1	0	0	0	0
61	15	4	0	0	0	0
61	16	2	0	0	0	0
61	17	9	0	0	0	0
61	18	13	0	0	2	0
61	1A	2243	0	0	55	0
61	1B	69	0	0	1	0
61	1D	26	0	0	2	0
61	1E	28	0	0	6	0
61	1F	19	0	0	1	0
61	1G	6	0	0	3	0
61	1H	2	0	0	0	0
61	1I	2	0	0	0	0
61	1N	6	0	0	2	0
61	1O	7	0	0	0	0
61	1P	20	0	0	2	0
61	1Q	15	0	0	0	0
61	1R	12	0	0	0	0
61	1S	5	0	0	0	0
61	1T	10	0	0	1	0
61	1U	16	0	0	0	0
61	1V	10	0	0	1	0
61	1W	6	0	0	0	0
61	1X	6	0	0	1	0
61	1Y	4	0	0	0	0
61	1Z	1	0	0	0	0
61	1a	460	0	0	28	0
61	1b	1	0	0	0	0
61	1d	1	0	0	0	0
61	1f	1	0	0	0	0
61	1g	1	0	0	0	0
61	1l	9	0	0	0	0
61	1m	1	0	0	0	0
61	1o	1	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
6l	1p	1	0	0	0	0
6l	1q	4	0	0	0	0
6l	1u	1	0	0	1	0
6l	1v	5	0	0	0	0
6l	1w	22	0	0	0	0
6l	1x	16	0	0	0	0
6l	1y	2	0	0	0	0
6l	20	4	0	0	0	0
6l	21	14	0	0	0	0
6l	22	2	0	0	0	0
6l	23	1	0	0	0	0
6l	25	5	0	0	1	0
6l	27	3	0	0	0	0
6l	28	4	0	0	0	0
6l	29	1	0	0	0	0
6l	2A	1395	0	0	72	0
6l	2B	28	0	0	0	0
6l	2D	25	0	0	0	0
6l	2E	15	0	0	1	0
6l	2F	14	0	0	0	0
6l	2I	4	0	0	0	0
6l	2N	1	0	0	0	0
6l	2O	1	0	0	0	0
6l	2P	14	0	0	2	0
6l	2Q	2	0	0	1	0
6l	2R	3	0	0	0	0
6l	2T	6	0	0	0	0
6l	2U	4	0	0	0	0
6l	2V	1	0	0	0	0
6l	2W	3	0	0	0	0
6l	2X	4	0	0	1	0
6l	2Y	1	0	0	0	0
6l	2Z	2	0	0	0	0
6l	2a	381	0	0	27	0
6l	2e	2	0	0	0	0
6l	2g	1	0	0	0	0
6l	2i	1	0	0	0	0
6l	2j	4	0	0	1	0
6l	2l	6	0	0	0	0
6l	2o	3	0	0	0	0
6l	2p	3	0	0	0	0
6l	2q	1	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
6l	2r	1	0	0	0	0
6l	2t	2	0	0	0	0
6l	2u	1	0	0	1	0
6l	2v	1	0	0	0	0
6l	2w	3	0	0	0	0
6l	2x	8	0	0	2	0
6l	2y	19	0	0	0	0
All	All	301023	0	196772	3963	0

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

The worst 5 of 3963 close contacts within the same asymmetric unit are listed below, sorted by their clash magnitude.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:1A:1128:U:H3	1:1A:1132:A:N6	1.27	1.29
32:2a:997:U:H3	32:2a:1044:A:N6	1.39	1.21
1:2A:2138:C:N4	1:2A:2153:G:H1	1.48	1.09
32:2a:1002:G:H1	32:2a:1038:C:N4	1.52	1.05
1:1A:1128:U:O4	1:1A:1132:A:N1	1.88	1.04

There are no symmetry-related clashes.

5.3 Torsion angles [i](#)

5.3.1 Protein backbone [i](#)

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
3	1D	273/276 (99%)	259 (95%)	13 (5%)	1 (0%)	30	54
3	2D	273/276 (99%)	258 (94%)	14 (5%)	1 (0%)	30	54
4	1E	202/206 (98%)	192 (95%)	9 (4%)	1 (0%)	24	48
4	2E	202/206 (98%)	193 (96%)	8 (4%)	1 (0%)	24	48

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
5	1F	201/210 (96%)	192 (96%)	8 (4%)	1 (0%)	24	48
5	2F	201/210 (96%)	193 (96%)	6 (3%)	2 (1%)	12	32
6	1G	179/182 (98%)	170 (95%)	8 (4%)	1 (1%)	21	44
6	2G	179/182 (98%)	164 (92%)	13 (7%)	2 (1%)	11	29
7	1H	172/180 (96%)	159 (92%)	10 (6%)	3 (2%)	7	19
7	2H	172/180 (96%)	162 (94%)	7 (4%)	3 (2%)	7	19
8	1I	144/148 (97%)	131 (91%)	12 (8%)	1 (1%)	18	41
8	2I	144/148 (97%)	133 (92%)	9 (6%)	2 (1%)	9	23
9	1N	138/140 (99%)	134 (97%)	4 (3%)	0	100	100
9	2N	138/140 (99%)	133 (96%)	5 (4%)	0	100	100
10	1O	120/122 (98%)	112 (93%)	8 (7%)	0	100	100
10	2O	120/122 (98%)	109 (91%)	11 (9%)	0	100	100
11	1P	147/150 (98%)	138 (94%)	9 (6%)	0	100	100
11	2P	147/150 (98%)	140 (95%)	7 (5%)	0	100	100
12	1Q	139/141 (99%)	130 (94%)	9 (6%)	0	100	100
12	2Q	139/141 (99%)	130 (94%)	9 (6%)	0	100	100
13	1R	116/118 (98%)	113 (97%)	3 (3%)	0	100	100
13	2R	116/118 (98%)	111 (96%)	5 (4%)	0	100	100
14	1S	108/112 (96%)	104 (96%)	4 (4%)	0	100	100
14	2S	108/112 (96%)	99 (92%)	9 (8%)	0	100	100
15	1T	129/146 (88%)	124 (96%)	2 (2%)	3 (2%)	5	13
15	2T	129/146 (88%)	122 (95%)	7 (5%)	0	100	100
16	1U	114/118 (97%)	113 (99%)	1 (1%)	0	100	100
16	2U	114/118 (97%)	113 (99%)	1 (1%)	0	100	100
17	1V	99/101 (98%)	94 (95%)	4 (4%)	1 (1%)	12	32
17	2V	99/101 (98%)	96 (97%)	2 (2%)	1 (1%)	12	32
18	1W	110/113 (97%)	109 (99%)	1 (1%)	0	100	100
18	2W	110/113 (97%)	108 (98%)	2 (2%)	0	100	100
19	1X	93/96 (97%)	90 (97%)	3 (3%)	0	100	100
19	2X	93/96 (97%)	89 (96%)	4 (4%)	0	100	100
20	1Y	105/110 (96%)	98 (93%)	6 (6%)	1 (1%)	12	32

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
20	2Y	105/110 (96%)	101 (96%)	4 (4%)	0	100	100
21	1Z	148/206 (72%)	136 (92%)	10 (7%)	2 (1%)	9	23
21	2Z	156/206 (76%)	136 (87%)	17 (11%)	3 (2%)	6	17
22	10	81/85 (95%)	79 (98%)	2 (2%)	0	100	100
22	20	81/85 (95%)	79 (98%)	2 (2%)	0	100	100
23	11	95/98 (97%)	92 (97%)	3 (3%)	0	100	100
23	21	95/98 (97%)	93 (98%)	1 (1%)	1 (1%)	11	29
24	12	68/72 (94%)	68 (100%)	0	0	100	100
24	22	68/72 (94%)	68 (100%)	0	0	100	100
25	13	57/60 (95%)	56 (98%)	1 (2%)	0	100	100
25	23	57/60 (95%)	55 (96%)	2 (4%)	0	100	100
26	14	67/71 (94%)	55 (82%)	10 (15%)	2 (3%)	3	8
26	24	67/71 (94%)	53 (79%)	12 (18%)	2 (3%)	3	8
27	15	57/60 (95%)	56 (98%)	1 (2%)	0	100	100
27	25	57/60 (95%)	55 (96%)	2 (4%)	0	100	100
28	16	51/54 (94%)	50 (98%)	1 (2%)	0	100	100
28	26	51/54 (94%)	48 (94%)	3 (6%)	0	100	100
29	17	46/49 (94%)	45 (98%)	1 (2%)	0	100	100
29	27	46/49 (94%)	46 (100%)	0	0	100	100
30	18	62/65 (95%)	62 (100%)	0	0	100	100
30	28	62/65 (95%)	60 (97%)	2 (3%)	0	100	100
31	19	35/37 (95%)	34 (97%)	1 (3%)	0	100	100
31	29	35/37 (95%)	35 (100%)	0	0	100	100
33	1b	229/256 (90%)	202 (88%)	19 (8%)	8 (4%)	3	6
33	2b	229/256 (90%)	205 (90%)	19 (8%)	5 (2%)	5	14
34	1c	204/239 (85%)	190 (93%)	13 (6%)	1 (0%)	24	48
34	2c	204/239 (85%)	190 (93%)	12 (6%)	2 (1%)	12	32
35	1d	206/209 (99%)	191 (93%)	13 (6%)	2 (1%)	12	32
35	2d	206/209 (99%)	192 (93%)	11 (5%)	3 (2%)	8	22
36	1e	146/162 (90%)	136 (93%)	8 (6%)	2 (1%)	9	23
36	2e	146/162 (90%)	135 (92%)	10 (7%)	1 (1%)	18	41

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
37	1f	98/101 (97%)	95 (97%)	2 (2%)	1 (1%)	12	32
37	2f	98/101 (97%)	98 (100%)	0	0	100	100
38	1g	153/156 (98%)	143 (94%)	8 (5%)	2 (1%)	9	25
38	2g	153/156 (98%)	142 (93%)	8 (5%)	3 (2%)	6	16
39	1h	135/138 (98%)	133 (98%)	2 (2%)	0	100	100
39	2h	135/138 (98%)	130 (96%)	5 (4%)	0	100	100
40	1i	125/128 (98%)	112 (90%)	11 (9%)	2 (2%)	7	20
40	2i	125/128 (98%)	116 (93%)	8 (6%)	1 (1%)	16	37
41	1j	95/105 (90%)	85 (90%)	7 (7%)	3 (3%)	3	7
41	2j	94/105 (90%)	82 (87%)	9 (10%)	3 (3%)	3	7
42	1k	112/129 (87%)	106 (95%)	5 (4%)	1 (1%)	14	35
42	2k	112/129 (87%)	103 (92%)	7 (6%)	2 (2%)	6	18
43	1l	119/132 (90%)	112 (94%)	7 (6%)	0	100	100
43	2l	119/132 (90%)	109 (92%)	10 (8%)	0	100	100
44	1m	121/126 (96%)	113 (93%)	8 (7%)	0	100	100
44	2m	120/126 (95%)	111 (92%)	9 (8%)	0	100	100
45	1n	58/61 (95%)	52 (90%)	6 (10%)	0	100	100
45	2n	58/61 (95%)	52 (90%)	6 (10%)	0	100	100
46	1o	86/89 (97%)	84 (98%)	2 (2%)	0	100	100
46	2o	86/89 (97%)	83 (96%)	3 (4%)	0	100	100
47	1p	80/88 (91%)	75 (94%)	5 (6%)	0	100	100
47	2p	80/88 (91%)	72 (90%)	8 (10%)	0	100	100
48	1q	97/105 (92%)	93 (96%)	4 (4%)	0	100	100
48	2q	97/105 (92%)	90 (93%)	7 (7%)	0	100	100
49	1r	66/88 (75%)	64 (97%)	2 (3%)	0	100	100
49	2r	66/88 (75%)	63 (96%)	3 (4%)	0	100	100
50	1s	81/93 (87%)	73 (90%)	6 (7%)	2 (2%)	4	11
50	2s	81/93 (87%)	71 (88%)	9 (11%)	1 (1%)	10	27
51	1t	94/106 (89%)	86 (92%)	4 (4%)	4 (4%)	2	4
51	2t	94/106 (89%)	86 (92%)	4 (4%)	4 (4%)	2	4
52	1u	21/27 (78%)	21 (100%)	0	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
52	2u	21/27 (78%)	21 (100%)	0	0	100	100
All	All	11370/12128 (94%)	10694 (94%)	588 (5%)	88 (1%)	16	37

5 of 88 Ramachandran outliers are listed below:

Mol	Chain	Res	Type
6	1G	47	LYS
7	1H	126	PRO
26	14	55	ARG
33	1b	10	LEU
41	1j	75	ILE

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%)	203 (94%)	12 (6%)	19	44
3	2D	215/218 (99%)	204 (95%)	11 (5%)	21	48
4	1E	164/166 (99%)	151 (92%)	13 (8%)	11	28
4	2E	164/166 (99%)	157 (96%)	7 (4%)	26	54
5	1F	160/166 (96%)	152 (95%)	8 (5%)	22	48
5	2F	159/166 (96%)	152 (96%)	7 (4%)	25	53
6	1G	143/156 (92%)	136 (95%)	7 (5%)	22	49
6	2G	143/156 (92%)	139 (97%)	4 (3%)	38	68
7	1H	144/148 (97%)	138 (96%)	6 (4%)	26	55
7	2H	144/148 (97%)	143 (99%)	1 (1%)	76	90
8	1I	113/124 (91%)	109 (96%)	4 (4%)	32	61
8	2I	105/124 (85%)	100 (95%)	5 (5%)	23	50
9	1N	118/119 (99%)	112 (95%)	6 (5%)	21	48
9	2N	118/119 (99%)	114 (97%)	4 (3%)	32	62
10	1O	100/100 (100%)	96 (96%)	4 (4%)	28	56

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
10	2O	100/100 (100%)	97 (97%)	3 (3%)	36	66
11	1P	115/116 (99%)	111 (96%)	4 (4%)	32	61
11	2P	115/116 (99%)	111 (96%)	4 (4%)	32	61
12	1Q	111/111 (100%)	109 (98%)	2 (2%)	51	78
12	2Q	111/111 (100%)	110 (99%)	1 (1%)	70	87
13	1R	101/101 (100%)	91 (90%)	10 (10%)	7	19
13	2R	101/101 (100%)	95 (94%)	6 (6%)	18	42
14	1S	86/88 (98%)	80 (93%)	6 (7%)	14	34
14	2S	85/88 (97%)	82 (96%)	3 (4%)	32	61
15	1T	115/127 (91%)	112 (97%)	3 (3%)	40	70
15	2T	113/127 (89%)	111 (98%)	2 (2%)	51	78
16	1U	93/94 (99%)	90 (97%)	3 (3%)	34	64
16	2U	93/94 (99%)	92 (99%)	1 (1%)	65	85
17	1V	80/82 (98%)	72 (90%)	8 (10%)	7	19
17	2V	80/82 (98%)	74 (92%)	6 (8%)	12	31
18	1W	90/92 (98%)	85 (94%)	5 (6%)	19	44
18	2W	90/92 (98%)	88 (98%)	2 (2%)	45	74
19	1X	77/78 (99%)	76 (99%)	1 (1%)	61	83
19	2X	77/78 (99%)	76 (99%)	1 (1%)	61	83
20	1Y	85/91 (93%)	83 (98%)	2 (2%)	43	72
20	2Y	85/91 (93%)	84 (99%)	1 (1%)	63	84
21	1Z	135/179 (75%)	134 (99%)	1 (1%)	76	90
21	2Z	137/179 (76%)	135 (98%)	2 (2%)	57	81
22	10	65/67 (97%)	64 (98%)	1 (2%)	57	81
22	20	65/67 (97%)	65 (100%)	0	100	100
23	11	80/83 (96%)	78 (98%)	2 (2%)	42	71
23	21	80/83 (96%)	78 (98%)	2 (2%)	42	71
24	12	65/67 (97%)	64 (98%)	1 (2%)	57	81
24	22	65/67 (97%)	64 (98%)	1 (2%)	57	81
25	13	51/52 (98%)	47 (92%)	4 (8%)	11	29
25	23	50/52 (96%)	49 (98%)	1 (2%)	48	76

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
26	14	59/63 (94%)	55 (93%)	4 (7%)	14	35
26	24	53/63 (84%)	50 (94%)	3 (6%)	18	43
27	15	50/52 (96%)	45 (90%)	5 (10%)	7	19
27	25	50/52 (96%)	48 (96%)	2 (4%)	28	56
28	16	51/52 (98%)	48 (94%)	3 (6%)	18	42
28	26	50/52 (96%)	49 (98%)	1 (2%)	48	76
29	17	41/42 (98%)	40 (98%)	1 (2%)	43	72
29	27	41/42 (98%)	40 (98%)	1 (2%)	43	72
30	18	54/55 (98%)	51 (94%)	3 (6%)	19	44
30	28	54/55 (98%)	51 (94%)	3 (6%)	19	44
31	19	34/34 (100%)	32 (94%)	2 (6%)	18	42
31	29	34/34 (100%)	33 (97%)	1 (3%)	37	67
33	1b	192/220 (87%)	188 (98%)	4 (2%)	47	75
33	2b	187/220 (85%)	178 (95%)	9 (5%)	23	50
34	1c	142/188 (76%)	141 (99%)	1 (1%)	76	90
34	2c	140/188 (74%)	138 (99%)	2 (1%)	59	82
35	1d	169/181 (93%)	163 (96%)	6 (4%)	31	60
35	2d	173/181 (96%)	168 (97%)	5 (3%)	37	67
36	1e	113/123 (92%)	112 (99%)	1 (1%)	70	87
36	2e	114/123 (93%)	112 (98%)	2 (2%)	51	78
37	1f	84/90 (93%)	82 (98%)	2 (2%)	43	72
37	2f	85/90 (94%)	83 (98%)	2 (2%)	43	72
38	1g	119/127 (94%)	118 (99%)	1 (1%)	73	88
38	2g	120/127 (94%)	118 (98%)	2 (2%)	53	79
39	1h	114/119 (96%)	111 (97%)	3 (3%)	40	70
39	2h	114/119 (96%)	109 (96%)	5 (4%)	25	53
40	1i	90/99 (91%)	90 (100%)	0	100	100
40	2i	89/99 (90%)	88 (99%)	1 (1%)	65	85
41	1j	66/92 (72%)	66 (100%)	0	100	100
41	2j	69/92 (75%)	69 (100%)	0	100	100
42	1k	82/99 (83%)	80 (98%)	2 (2%)	43	72

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
42	2k	83/99 (84%)	82 (99%)	1 (1%)	63	84
43	1l	96/108 (89%)	92 (96%)	4 (4%)	26	55
43	2l	96/108 (89%)	96 (100%)	0	100	100
44	1m	93/101 (92%)	91 (98%)	2 (2%)	45	74
44	2m	92/101 (91%)	91 (99%)	1 (1%)	65	85
45	1n	49/50 (98%)	44 (90%)	5 (10%)	7	18
45	2n	49/50 (98%)	48 (98%)	1 (2%)	48	76
46	1o	78/80 (98%)	78 (100%)	0	100	100
46	2o	78/80 (98%)	77 (99%)	1 (1%)	61	83
47	1p	69/74 (93%)	65 (94%)	4 (6%)	18	42
47	2p	68/74 (92%)	66 (97%)	2 (3%)	37	67
48	1q	94/97 (97%)	93 (99%)	1 (1%)	65	85
48	2q	94/97 (97%)	94 (100%)	0	100	100
49	1r	59/77 (77%)	59 (100%)	0	100	100
49	2r	59/77 (77%)	58 (98%)	1 (2%)	53	79
50	1s	69/80 (86%)	67 (97%)	2 (3%)	37	67
50	2s	67/80 (84%)	65 (97%)	2 (3%)	36	66
51	1t	70/82 (85%)	70 (100%)	0	100	100
51	2t	70/82 (85%)	70 (100%)	0	100	100
52	1u	18/22 (82%)	18 (100%)	0	100	100
52	2u	18/22 (82%)	18 (100%)	0	100	100
All	All	9303/10064 (92%)	9011 (97%)	292 (3%)	35	65

5 of 292 residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
17	2V	62	LEU
46	2o	87	ILE
21	2Z	18	LEU
33	2b	127	ILE
18	1W	17	VAL

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. 5 of 107 such sidechains are listed below:

Mol	Chain	Res	Type
7	2H	139	GLN
26	24	46	GLN
46	2o	9	GLN
10	2O	3	GLN
21	2Z	55	HIS

5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	1A	2864/2915 (98%)	414 (14%)	34 (1%)
1	2A	2791/2915 (95%)	455 (16%)	22 (0%)
2	1B	120/121 (99%)	11 (9%)	1 (0%)
2	2B	118/121 (97%)	27 (22%)	0
32	1a	1497/1521 (98%)	232 (15%)	0
32	2a	1501/1521 (98%)	265 (17%)	0
53	1v	12/24 (50%)	2 (16%)	0
53	2v	12/24 (50%)	2 (16%)	0
54	1w	72/76 (94%)	23 (31%)	0
54	1y	72/76 (94%)	22 (30%)	0
54	2w	69/76 (90%)	24 (34%)	0
54	2y	70/76 (92%)	19 (27%)	0
55	1x	75/77 (97%)	9 (12%)	0
55	2x	75/77 (97%)	15 (20%)	0
All	All	9348/9620 (97%)	1520 (16%)	57 (0%)

5 of 1520 RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	1A	12	U
1	1A	13	A
1	1A	27	G
1	1A	34	C
1	1A	35	G

5 of 57 RNA pucker outliers are listed below:

Mol	Chain	Res	Type
1	1A	2205	C
1	2A	2126	A
1	2A	196	A
1	2A	2119	A
1	2A	1442	G

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

84 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	1w	8	54	18,21,22	1.75	4 (22%)	25,30,33	2.25	5 (20%)
32	PSU	1a	516	32	18,21,22	1.37	3 (16%)	21,30,33	1.96	4 (19%)
32	5MC	1a	1404	32	19,22,23	1.73	3 (15%)	26,32,35	1.12	3 (11%)
54	PSU	1w	39	54	18,21,22	1.40	2 (11%)	21,30,33	2.06	4 (19%)
32	MA6	1a	1519	32	23,26,27	1.53	4 (17%)	33,38,41	2.35	13 (39%)
54	MIA	2y	37	54	21,24,32	1.65	4 (19%)	30,35,47	2.02	7 (23%)
1	PSU	2A	1917	1	18,21,22	1.35	2 (11%)	21,30,33	2.05	3 (14%)
1	2MU	2A	2552	1	19,22,24	1.31	3 (15%)	25,31,36	1.88	6 (24%)
32	M2G	2a	966	32	24,27,28	1.30	3 (12%)	33,40,43	1.85	6 (18%)
1	5MC	2A	1962	56,1	19,22,23	1.45	3 (15%)	26,32,35	1.11	2 (7%)
54	7MG	2w	46	54	23,26,27	1.36	2 (8%)	27,39,42	2.55	7 (25%)
1	PSU	1A	2617	56,1	18,21,22	1.37	3 (16%)	21,30,33	2.04	3 (14%)
1	4OC	1A	1942	1	19,22,24	0.79	0	25,31,35	0.97	1 (4%)
55	5MC	2x	32	55	19,22,23	1.63	2 (10%)	26,32,35	1.20	3 (11%)
32	2MG	1a	1207	32	23,26,27	1.27	3 (13%)	33,38,41	2.15	9 (27%)
32	5MC	2a	1400	32	19,22,23	1.62	3 (15%)	26,32,35	1.21	3 (11%)
32	5MC	2a	1407	32	19,22,23	1.50	3 (15%)	26,32,35	1.17	3 (11%)
55	4SU	1x	8	55	18,21,22	2.21	4 (22%)	25,30,33	1.58	5 (20%)
54	7MG	2y	46	54	23,26,27	1.35	3 (13%)	27,39,42	2.76	7 (25%)
54	7MG	1y	46	54	23,26,27	1.36	3 (13%)	27,39,42	2.73	6 (22%)
54	PSU	2y	32	54	18,21,22	1.38	2 (11%)	21,30,33	1.89	3 (14%)
1	2MA	2A	2503	56,1	22,25,26	1.48	5 (22%)	32,37,40	2.22	9 (28%)
54	4SU	1y	8	54	18,21,22	1.78	5 (27%)	25,30,33	1.92	4 (16%)
32	4OC	1a	1402	32	20,23,24	0.75	0	25,32,35	1.03	2 (8%)
1	OMG	1A	2263	56,1,55	23,26,27	1.26	3 (13%)	32,38,41	1.97	6 (18%)
55	4SU	2x	8	55	18,21,22	2.01	6 (33%)	25,30,33	1.59	6 (24%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
32	MA6	2a	1519	32	23,26,27	1.60	4 (17%)	33,38,41	2.26	11 (33%)
54	PSU	1y	55	54	18,21,22	1.39	2 (11%)	21,30,33	2.08	3 (14%)
1	PSU	1A	1939	1	18,21,22	1.35	2 (11%)	21,30,33	1.91	4 (19%)
1	5MU	2A	1939	56,1	19,22,23	1.41	5 (26%)	27,32,35	2.18	6 (22%)
54	7MG	1w	46	54	23,26,27	1.45	4 (17%)	27,39,42	2.55	7 (25%)
54	5MU	1y	54	54	19,22,23	1.52	6 (31%)	27,32,35	1.85	6 (22%)
55	PSU	2x	55	55	18,21,22	1.36	2 (11%)	21,30,33	2.00	4 (19%)
55	5MU	2x	54	55	19,22,23	1.38	5 (26%)	27,32,35	2.34	6 (22%)
1	5MC	1A	1984	1	19,22,23	1.59	3 (15%)	26,32,35	1.11	2 (7%)
54	5MU	1w	54	54	19,22,23	1.34	5 (26%)	27,32,35	2.09	6 (22%)
1	PSU	2A	2605	1	18,21,22	1.36	3 (16%)	21,30,33	2.03	5 (23%)
54	4SU	2y	8	54	18,21,22	1.80	6 (33%)	25,30,33	1.89	4 (16%)
1	5MU	1A	1961	56,1	19,22,23	1.42	5 (26%)	27,32,35	2.25	6 (22%)
32	5MC	1a	1400	32	19,22,23	1.61	3 (15%)	26,32,35	1.14	3 (11%)
55	5MC	1x	32	55	19,22,23	1.69	3 (15%)	26,32,35	1.22	3 (11%)
54	4SU	2w	8	54	18,21,22	1.74	4 (22%)	25,30,33	2.63	6 (24%)
1	2MU	1A	2564	56,1	19,22,24	1.28	3 (15%)	25,31,36	1.94	5 (20%)
32	MA6	1a	1518	32	23,26,27	1.52	5 (21%)	33,38,41	2.27	12 (36%)
54	MIA	1y	37	54	21,24,32	1.63	3 (14%)	30,35,47	2.03	9 (30%)
32	4OC	2a	1402	32	20,23,24	0.78	0	25,32,35	1.12	2 (8%)
54	PSU	2y	39	54	18,21,22	1.31	2 (11%)	21,30,33	2.00	4 (19%)
32	M2G	1a	966	32	24,27,28	1.30	3 (12%)	33,40,43	1.86	6 (18%)
54	MIA	1w	37	54	28,31,32	2.30	6 (21%)	38,44,47	2.72	11 (28%)
54	PSU	2w	55	54	18,21,22	1.42	2 (11%)	21,30,33	2.02	3 (14%)
32	PSU	2a	516	32	18,21,22	1.36	2 (11%)	21,30,33	1.88	4 (19%)
54	MIA	2w	37	54	24,27,32	2.11	5 (20%)	32,39,47	2.35	11 (34%)
32	UR3	2a	1498	32	19,22,23	1.01	1 (5%)	26,32,35	1.76	3 (11%)
32	5MC	2a	967	32	19,22,23	1.80	3 (15%)	26,32,35	1.18	3 (11%)
32	MA6	2a	1518	32	23,26,27	1.58	3 (13%)	33,38,41	2.34	13 (39%)
54	PSU	1w	55	54	18,21,22	1.40	2 (11%)	21,30,33	1.97	4 (19%)
1	4OC	2A	1920	1	19,22,24	0.78	0	25,31,35	0.79	0
54	PSU	1y	39	54	18,21,22	1.37	2 (11%)	21,30,33	2.00	3 (14%)
1	5MC	2A	1942	1	19,22,23	1.64	3 (15%)	26,32,35	1.15	2 (7%)
1	2MA	1A	2515	56,1	22,25,26	1.39	4 (18%)	32,37,40	2.29	8 (25%)
1	OMG	2A	2251	1,55	23,26,27	1.21	3 (13%)	32,38,41	1.96	6 (18%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
54	PSU	2w	32	54	18,21,22	1.34	2 (11%)	21,30,33	2.02	4 (19%)
32	5MC	1a	967	32	19,22,23	1.71	3 (15%)	26,32,35	1.13	3 (11%)
1	PSU	1A	1933	1	18,21,22	1.34	2 (11%)	21,30,33	2.13	4 (19%)
43	0TD	1l	92	43	8,9,10	4.69	2 (25%)	6,11,13	6.06	2 (33%)
32	5MC	2a	1404	32	19,22,23	1.73	3 (15%)	26,32,35	1.23	3 (11%)
32	2MG	2a	1207	32	23,26,27	1.24	3 (13%)	33,38,41	2.14	6 (18%)
1	5MU	1A	1937	1	19,22,23	1.41	4 (21%)	27,32,35	2.10	7 (25%)
54	PSU	1y	32	54	18,21,22	1.39	2 (11%)	21,30,33	1.97	3 (14%)
54	PSU	2y	55	54	18,21,22	1.36	2 (11%)	21,30,33	2.02	4 (19%)
32	7MG	2a	527	56,32	23,26,27	1.36	3 (13%)	27,39,42	2.58	7 (25%)
54	PSU	2w	39	54	18,21,22	1.36	2 (11%)	21,30,33	1.76	3 (14%)
55	PSU	1x	55	56,55	18,21,22	1.39	2 (11%)	21,30,33	2.07	4 (19%)
1	PSU	2A	1911	1	18,21,22	1.34	2 (11%)	21,30,33	1.97	3 (14%)
54	5MU	2y	54	54	19,22,23	1.52	5 (26%)	27,32,35	2.08	8 (29%)
32	5MC	1a	1407	32	19,22,23	1.54	2 (10%)	26,32,35	1.14	2 (7%)
54	5MU	2w	54	54	19,22,23	1.40	5 (26%)	27,32,35	1.96	6 (22%)
1	5MU	2A	1915	1	19,22,23	1.45	6 (31%)	27,32,35	2.10	6 (22%)
43	0TD	2l	92	43	8,9,10	4.63	2 (25%)	6,11,13	2.92	3 (50%)
32	7MG	1a	527	32	23,26,27	1.41	4 (17%)	27,39,42	2.53	7 (25%)
54	PSU	1w	32	54	18,21,22	1.34	2 (11%)	21,30,33	1.98	3 (14%)
55	5MU	1x	54	56,55	19,22,23	1.43	6 (31%)	27,32,35	1.99	8 (29%)
1	5MC	1A	1964	1	19,22,23	1.40	3 (15%)	26,32,35	1.14	2 (7%)
32	UR3	1a	1498	32	19,22,23	1.04	1 (5%)	26,32,35	1.74	3 (11%)

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	1w	8	54	-	0/7/25/26	0/2/2/2
32	PSU	1a	516	32	-	0/7/25/26	0/2/2/2
32	5MC	1a	1404	32	-	0/7/25/26	0/2/2/2
54	PSU	1w	39	54	-	0/7/25/26	0/2/2/2
32	MA6	1a	1519	32	-	4/11/29/30	0/3/3/3
54	MIA	2y	37	54	-	2/7/25/34	0/3/3/3
1	PSU	2A	1917	1	-	2/7/25/26	0/2/2/2

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
1	2MU	2A	2552	1	-	0/9/27/28	0/2/2/2
32	M2G	2a	966	32	-	0/11/29/30	0/3/3/3
1	5MC	2A	1962	56,1	-	0/7/25/26	0/2/2/2
54	7MG	2w	46	54	-	4/7/37/38	0/3/3/3
1	PSU	1A	2617	56,1	-	0/7/25/26	0/2/2/2
1	4OC	1A	1942	1	-	1/9/27/30	0/2/2/2
55	5MC	2x	32	55	-	0/7/25/26	0/2/2/2
32	2MG	1a	1207	32	-	0/9/27/28	0/3/3/3
32	5MC	2a	1400	32	-	0/7/25/26	0/2/2/2
32	5MC	2a	1407	32	-	0/7/25/26	0/2/2/2
55	4SU	1x	8	55	-	0/7/25/26	0/2/2/2
54	7MG	2y	46	54	-	3/7/37/38	0/3/3/3
54	7MG	1y	46	54	-	3/7/37/38	0/3/3/3
54	PSU	2y	32	54	-	0/7/25/26	0/2/2/2
1	2MA	2A	2503	56,1	-	1/7/25/26	0/3/3/3
54	4SU	1y	8	54	-	2/7/25/26	0/2/2/2
32	4OC	1a	1402	32	-	3/9/29/30	0/2/2/2
1	OMG	1A	2263	56,1,55	-	0/9/27/28	0/3/3/3
55	4SU	2x	8	55	-	0/7/25/26	0/2/2/2
32	MA6	2a	1519	32	-	6/11/29/30	0/3/3/3
54	PSU	1y	55	54	-	2/7/25/26	0/2/2/2
1	PSU	1A	1939	1	-	0/7/25/26	0/2/2/2
1	5MU	2A	1939	56,1	-	0/7/25/26	0/2/2/2
54	7MG	1w	46	54	-	2/7/37/38	0/3/3/3
54	5MU	1y	54	54	-	3/7/25/26	0/2/2/2
55	PSU	2x	55	55	-	0/7/25/26	0/2/2/2
55	5MU	2x	54	55	-	0/7/25/26	0/2/2/2
1	5MC	1A	1984	1	-	2/7/25/26	0/2/2/2
54	5MU	1w	54	54	-	0/7/25/26	0/2/2/2
1	PSU	2A	2605	1	-	0/7/25/26	0/2/2/2
54	4SU	2y	8	54	-	2/7/25/26	0/2/2/2
1	5MU	1A	1961	56,1	-	0/7/25/26	0/2/2/2
32	5MC	1a	1400	32	-	2/7/25/26	0/2/2/2
55	5MC	1x	32	55	-	0/7/25/26	0/2/2/2
54	4SU	2w	8	54	-	0/7/25/26	0/2/2/2
1	2MU	1A	2564	56,1	-	0/9/27/28	0/2/2/2
32	MA6	1a	1518	32	-	1/11/29/30	0/3/3/3
54	MIA	1y	37	54	-	1/7/25/34	0/3/3/3
32	4OC	2a	1402	32	-	3/9/29/30	0/2/2/2

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
54	PSU	2y	39	54	-	0/7/25/26	0/2/2/2
32	M2G	1a	966	32	-	0/11/29/30	0/3/3/3
54	MIA	1w	37	54	-	1/15/33/34	0/3/3/3
54	PSU	2w	55	54	-	0/7/25/26	0/2/2/2
32	PSU	2a	516	32	-	0/7/25/26	0/2/2/2
54	MIA	2w	37	54	-	2/11/29/34	0/3/3/3
32	UR3	2a	1498	32	-	0/7/25/26	0/2/2/2
32	5MC	2a	967	32	-	0/7/25/26	0/2/2/2
32	MA6	2a	1518	32	-	2/11/29/30	0/3/3/3
54	PSU	1w	55	54	-	0/7/25/26	0/2/2/2
1	4OC	2A	1920	1	-	0/9/27/30	0/2/2/2
54	PSU	1y	39	54	-	0/7/25/26	0/2/2/2
1	5MC	2A	1942	1	-	0/7/25/26	0/2/2/2
1	2MA	1A	2515	56,1	-	1/7/25/26	0/3/3/3
1	OMG	2A	2251	1,55	-	0/9/27/28	0/3/3/3
54	PSU	2w	32	54	-	0/7/25/26	0/2/2/2
32	5MC	1a	967	32	-	0/7/25/26	0/2/2/2
1	PSU	1A	1933	1	-	0/7/25/26	0/2/2/2
43	0TD	1l	92	43	-	1/7/12/14	-
32	5MC	2a	1404	32	-	0/7/25/26	0/2/2/2
32	2MG	2a	1207	32	-	0/9/27/28	0/3/3/3
1	5MU	1A	1937	1	-	0/7/25/26	0/2/2/2
54	PSU	1y	32	54	-	0/7/25/26	0/2/2/2
54	PSU	2y	55	54	-	0/7/25/26	0/2/2/2
32	7MG	2a	527	56,32	-	3/7/37/38	0/3/3/3
54	PSU	2w	39	54	-	0/7/25/26	0/2/2/2
55	PSU	1x	55	56,55	-	1/7/25/26	0/2/2/2
1	PSU	2A	1911	1	-	0/7/25/26	0/2/2/2
54	5MU	2y	54	54	-	3/7/25/26	0/2/2/2
32	5MC	1a	1407	32	-	0/7/25/26	0/2/2/2
54	5MU	2w	54	54	-	0/7/25/26	0/2/2/2
1	5MU	2A	1915	1	-	0/7/25/26	0/2/2/2
43	0TD	2l	92	43	-	2/7/12/14	-
32	7MG	1a	527	32	-	2/7/37/38	0/3/3/3
54	PSU	1w	32	54	-	0/7/25/26	0/2/2/2
55	5MU	1x	54	56,55	-	0/7/25/26	0/2/2/2
1	5MC	1A	1964	1	-	0/7/25/26	0/2/2/2
32	UR3	1a	1498	32	-	0/7/25/26	0/2/2/2

The worst 5 of 260 bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
43	1l	92	0TD	CB-SB	-12.76	1.69	1.82
43	2l	92	0TD	CB-SB	-12.48	1.69	1.82
54	2w	37	MIA	C2-S10	-7.02	1.69	1.75
54	1w	37	MIA	C13-C14	6.88	1.53	1.32
32	2a	967	5MC	C5-C4	6.78	1.49	1.44

The worst 5 of 422 bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
43	1l	92	0TD	CSB-SB-CB	-14.41	76.47	102.36
54	2y	46	7MG	N9-C4-N3	9.87	139.93	125.46
54	1y	46	7MG	N9-C4-N3	9.84	139.89	125.46
54	1w	37	MIA	C12-C13-C14	-8.89	111.05	127.01
54	1w	46	7MG	N9-C4-N3	8.89	138.49	125.46

There are no chirality outliers.

5 of 67 torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
32	1a	1519	MA6	O4'-C4'-C5'-O5'
32	2a	1402	4OC	O4'-C4'-C5'-O5'
32	2a	1518	MA6	C5-C6-N6-C9
32	2a	1519	MA6	O4'-C4'-C5'-O5'
43	2l	92	0TD	O-C-CA-CB

There are no ring outliers.

36 monomers are involved in 50 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
54	1w	39	PSU	1	0
32	1a	1519	MA6	2	0
54	2y	37	MIA	1	0
1	2A	2552	2MU	2	0
32	2a	966	M2G	1	0
32	2a	1400	5MC	1	0
55	1x	8	4SU	2	0
54	2y	46	7MG	1	0
54	1y	46	7MG	1	0
1	2A	2503	2MA	2	0
54	1y	8	4SU	3	0
32	1a	1402	4OC	2	0
55	2x	8	4SU	1	0

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Mol	Chain	Res	Type	Clashes	Symm-Clashes
32	2a	1519	MA6	3	0
1	1A	1939	PSU	1	0
54	1w	46	7MG	1	0
55	2x	55	PSU	1	0
54	2y	8	4SU	1	0
55	1x	32	5MC	1	0
54	2w	8	4SU	4	0
1	1A	2564	2MU	1	0
32	1a	1518	MA6	2	0
54	1y	37	MIA	1	0
32	2a	1402	4OC	2	0
54	1w	37	MIA	1	0
54	2w	37	MIA	1	0
32	2a	967	5MC	2	0
32	2a	1518	MA6	3	0
54	1y	39	PSU	1	0
54	2w	32	PSU	1	0
43	1l	92	0TD	1	0
32	2a	1404	5MC	1	0
32	2a	1207	2MG	2	0
54	2y	55	PSU	1	0
54	2w	39	PSU	3	0
1	2A	1915	5MU	1	0

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

Of 2867 ligands modelled in this entry, 2863 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
60	SF4	1d	501	35	0,12,12	-	-	-		
57	MT9	2A	3897	-	33,34,34	2.36	4 (12%)	44,50,50	1.59	10 (22%)
57	MT9	1A	4113	-	33,34,34	2.44	4 (12%)	44,50,50	1.66	10 (22%)
60	SF4	2d	501	35	0,12,12	-	-	-		

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

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
60	SF4	1d	501	35	-	-	0/6/5/5
57	MT9	2A	3897	-	-	3/46/62/62	0/1/2/2
57	MT9	1A	4113	-	-	5/46/62/62	0/1/2/2
60	SF4	2d	501	35	-	-	0/6/5/5

The worst 5 of 8 bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
57	1A	4113	MT9	C4-C5	-10.27	1.39	1.51
57	2A	3897	MT9	C4-C5	-10.17	1.39	1.51
57	1A	4113	MT9	C8-C7	-5.87	1.42	1.51
57	1A	4113	MT9	C12-C13	-5.53	1.39	1.51
57	2A	3897	MT9	C8-C7	-5.42	1.43	1.51

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

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
57	1A	4113	MT9	C3-O3-C13	-4.22	110.83	118.20
57	1A	4113	MT9	C8-C7-C6	3.64	126.95	117.90
57	2A	3897	MT9	C8-C7-C6	3.63	126.94	117.90
57	1A	4113	MT9	O3-C3-C2	3.39	113.67	107.36
57	2A	3897	MT9	C3-O3-C13	-3.37	112.32	118.20

There are no chirality outliers.

5 of 8 torsion outliers are listed below:

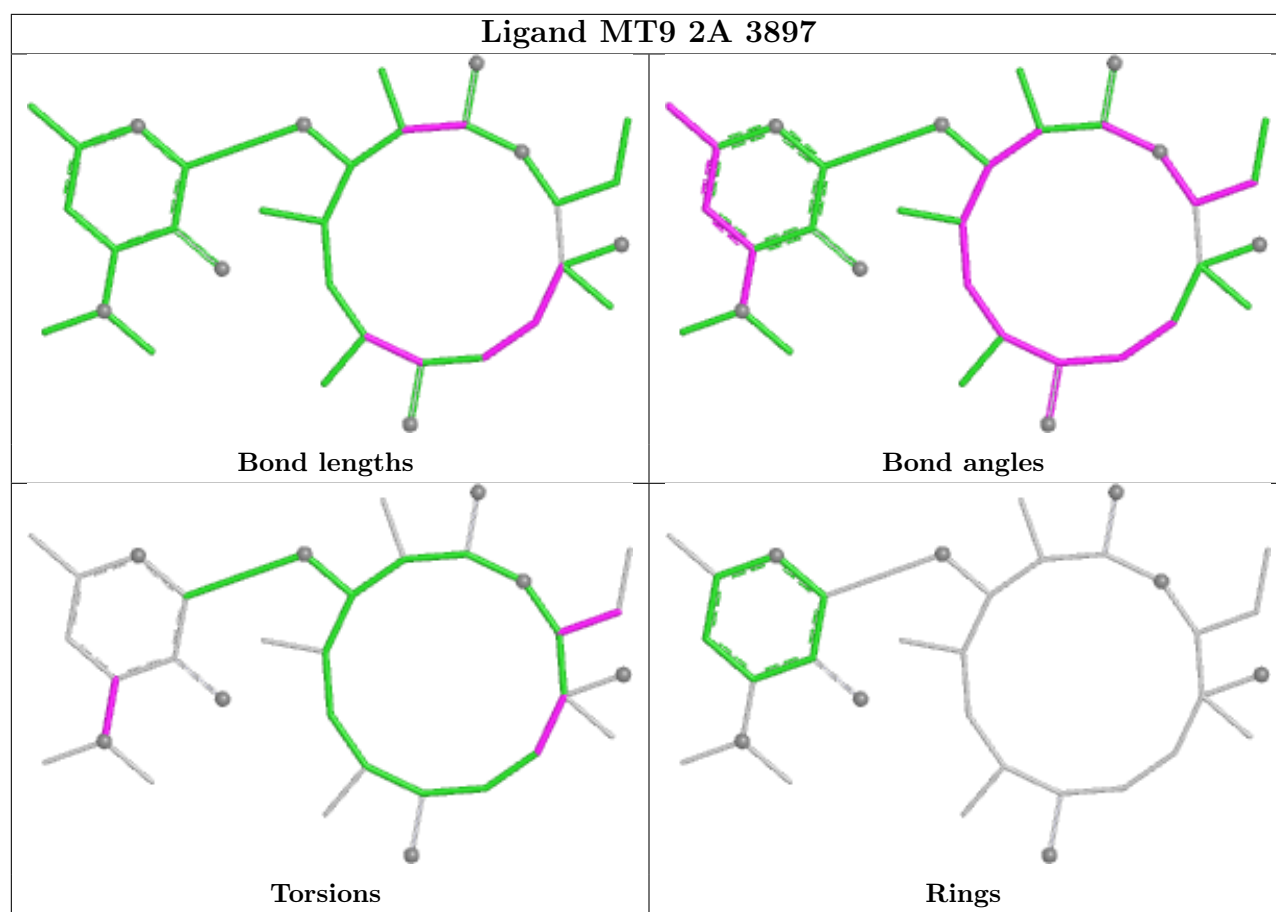
Mol	Chain	Res	Type	Atoms
57	1A	4113	MT9	C18-C17-N1-C21
57	2A	3897	MT9	C18-C17-N1-C21
57	1A	4113	MT9	C25-C4-C5-C6
57	2A	3897	MT9	C1-C2-C3-O3
57	1A	4113	MT9	O5-C15-O4-C11

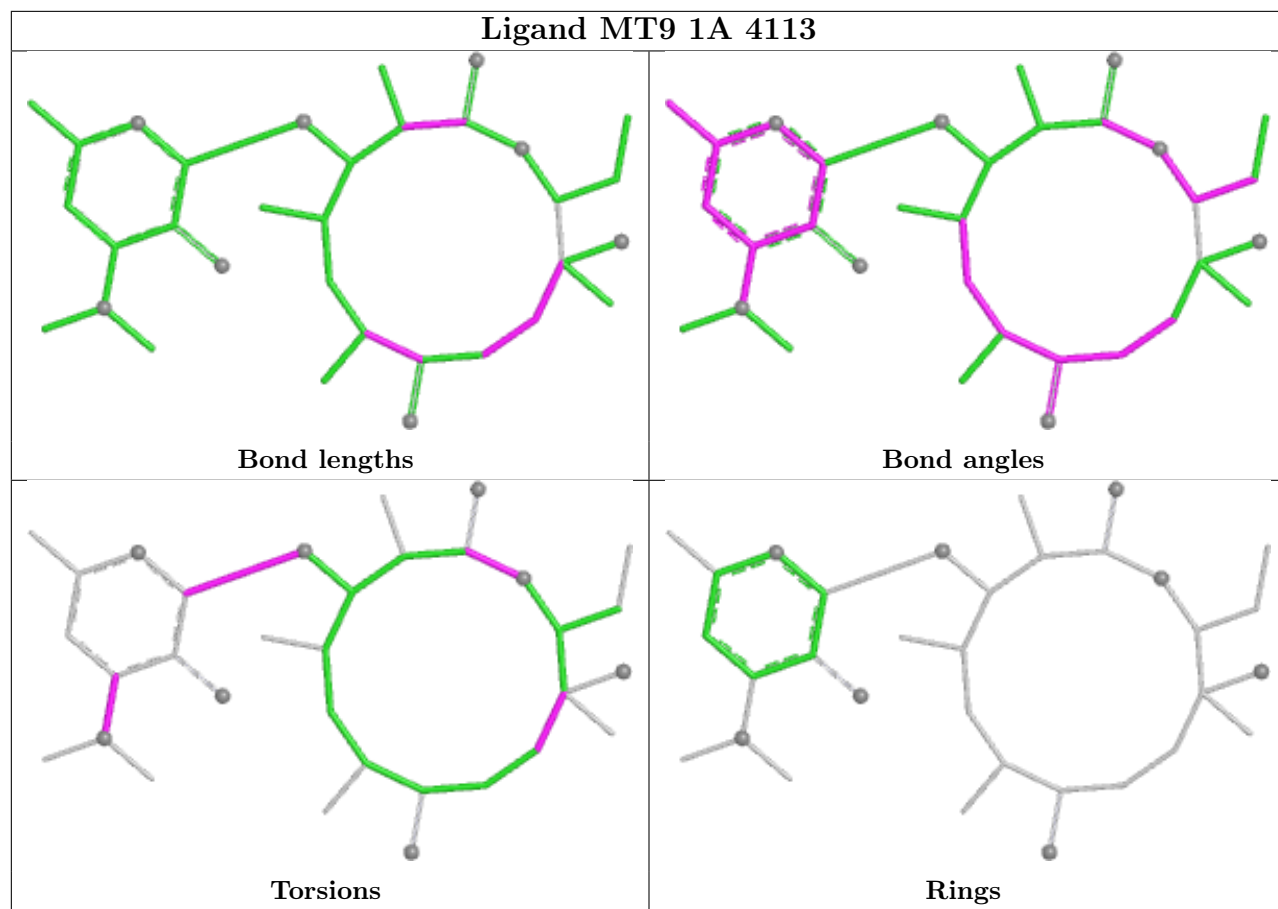
There are no ring outliers.

2 monomers are involved in 2 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
57	2A	3897	MT9	1	0
57	1A	4113	MT9	1	0

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 ⓘ

6.1 Protein, DNA and RNA chains ⓘ

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

Mol	Chain	Analysed	<RSRZ>	#RSRZ > 2			OWAB(Å ²)	Q < 0.9
1	1A	2860/2915 (98%)	-0.50	107 (3%)	45	41	16, 34, 90, 106	0
1	2A	2789/2915 (95%)	0.11	101 (3%)	46	42	32, 57, 91, 107	0
2	1B	120/121 (99%)	-0.39	0	100	100	30, 46, 58, 84	0
2	2B	120/121 (99%)	1.40	24 (20%)	3	2	58, 80, 88, 92	0
3	1D	275/276 (99%)	-0.12	6 (2%)	62	59	20, 35, 52, 85	0
3	2D	275/276 (99%)	0.32	4 (1%)	72	70	31, 50, 64, 85	0
4	1E	204/206 (99%)	-0.16	1 (0%)	87	86	18, 37, 56, 77	0
4	2E	204/206 (99%)	0.56	4 (1%)	65	62	33, 58, 71, 79	0
5	1F	203/210 (96%)	-0.01	2 (0%)	79	78	19, 39, 66, 88	0
5	2F	203/210 (96%)	0.74	6 (2%)	52	49	34, 68, 81, 88	0
6	1G	181/182 (99%)	0.61	10 (5%)	30	27	34, 56, 70, 86	0
6	2G	181/182 (99%)	1.82	67 (37%)	1	0	71, 81, 88, 96	0
7	1H	174/180 (96%)	0.26	7 (4%)	42	38	33, 51, 64, 71	0
7	2H	174/180 (96%)	1.76	60 (34%)	1	1	68, 83, 90, 96	0
8	1I	146/148 (98%)	0.84	5 (3%)	48	44	41, 72, 81, 84	0
8	2I	146/148 (98%)	1.18	20 (13%)	6	5	57, 72, 83, 91	0
9	1N	140/140 (100%)	-0.17	1 (0%)	84	83	24, 35, 58, 70	0
9	2N	140/140 (100%)	0.80	5 (3%)	46	42	44, 65, 78, 86	0
10	1O	122/122 (100%)	-0.16	0	100	100	24, 37, 56, 62	0
10	2O	122/122 (100%)	0.54	2 (1%)	70	68	45, 58, 69, 72	0
11	1P	149/150 (99%)	0.15	0	100	100	19, 42, 67, 76	0
11	2P	149/150 (99%)	0.79	10 (6%)	24	21	37, 67, 82, 89	0
12	1Q	141/141 (100%)	0.07	3 (2%)	63	61	26, 38, 54, 78	0
12	2Q	141/141 (100%)	1.30	30 (21%)	2	2	48, 66, 78, 86	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
13	1R	118/118 (100%)	-0.31	0 100 100	22, 31, 44, 59	0
13	2R	118/118 (100%)	0.46	4 (3%) 48 44	39, 52, 63, 67	0
14	1S	110/112 (98%)	0.15	0 100 100	34, 46, 60, 67	0
14	2S	110/112 (98%)	1.62	28 (25%) 1 1	63, 76, 84, 90	0
15	1T	131/146 (89%)	0.10	4 (3%) 51 48	30, 42, 63, 77	0
15	2T	131/146 (89%)	0.72	4 (3%) 51 48	51, 61, 76, 84	0
16	1U	116/118 (98%)	-0.33	0 100 100	19, 29, 45, 63	0
16	2U	116/118 (98%)	0.71	2 (1%) 69 67	43, 62, 77, 85	0
17	1V	101/101 (100%)	-0.32	0 100 100	18, 36, 53, 62	0
17	2V	101/101 (100%)	0.99	7 (6%) 23 20	43, 70, 79, 84	0
18	1W	112/113 (99%)	-0.40	1 (0%) 81 80	22, 30, 47, 82	0
18	2W	112/113 (99%)	0.19	1 (0%) 81 80	38, 49, 63, 85	0
19	1X	95/96 (98%)	-0.12	2 (2%) 63 61	26, 36, 54, 72	0
19	2X	95/96 (98%)	0.78	7 (7%) 20 18	45, 59, 73, 80	0
20	1Y	107/110 (97%)	0.20	3 (2%) 55 51	31, 45, 63, 83	0
20	2Y	107/110 (97%)	1.32	17 (15%) 5 4	60, 72, 81, 87	0
21	1Z	154/206 (74%)	0.96	17 (11%) 10 9	39, 59, 81, 89	0
21	2Z	160/206 (77%)	2.12	77 (48%) 0 0	70, 82, 92, 97	0
22	10	83/85 (97%)	0.09	5 (6%) 27 24	27, 36, 59, 68	0
22	20	83/85 (97%)	1.22	15 (18%) 3 3	44, 63, 75, 84	0
23	11	97/98 (98%)	0.13	1 (1%) 79 78	27, 41, 67, 72	0
23	21	97/98 (98%)	0.66	3 (3%) 51 48	41, 58, 76, 78	0
24	12	70/72 (97%)	0.11	0 100 100	32, 47, 56, 73	0
24	22	70/72 (97%)	0.89	5 (7%) 22 19	58, 69, 78, 81	0
25	13	59/60 (98%)	-0.22	0 100 100	21, 33, 56, 76	0
25	23	59/60 (98%)	0.82	4 (6%) 23 20	56, 65, 80, 90	0
26	14	69/71 (97%)	0.99	11 (15%) 5 4	47, 72, 86, 92	0
26	24	69/71 (97%)	2.05	30 (43%) 0 0	78, 88, 94, 95	0
27	15	59/60 (98%)	-0.28	1 (1%) 69 67	18, 29, 48, 62	0
27	25	59/60 (98%)	0.16	0 100 100	37, 51, 66, 73	0
28	16	53/54 (98%)	-0.17	0 100 100	28, 39, 53, 69	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
28	26	53/54 (98%)	0.94	2 (3%) 44 40	48, 61, 70, 75	0
29	17	48/49 (97%)	-0.24	4 (8%) 17 14	20, 26, 60, 66	0
29	27	48/49 (97%)	0.16	4 (8%) 17 14	32, 41, 65, 76	0
30	18	64/65 (98%)	-0.10	0 100 100	25, 31, 40, 53	0
30	28	64/65 (98%)	0.75	2 (3%) 51 48	47, 55, 63, 65	0
31	19	37/37 (100%)	-0.20	0 100 100	27, 37, 52, 56	0
31	29	37/37 (100%)	1.41	7 (18%) 3 3	62, 69, 75, 78	0
32	1a	1488/1521 (97%)	0.32	46 (3%) 51 48	35, 63, 89, 104	0
32	2a	1491/1521 (98%)	0.90	161 (10%) 11 9	52, 77, 95, 106	0
33	1b	231/256 (90%)	1.30	39 (16%) 4 3	66, 76, 86, 93	0
33	2b	231/256 (90%)	2.07	110 (47%) 0 0	75, 87, 94, 100	0
34	1c	206/239 (86%)	0.88	11 (5%) 32 28	58, 69, 80, 87	0
34	2c	206/239 (86%)	2.16	119 (57%) 0 0	76, 86, 91, 101	0
35	1d	208/209 (99%)	0.96	17 (8%) 17 15	52, 68, 79, 85	0
35	2d	208/209 (99%)	1.33	40 (19%) 3 3	62, 74, 82, 89	0
36	1e	148/162 (91%)	0.54	3 (2%) 65 62	52, 62, 73, 80	0
36	2e	148/162 (91%)	1.55	36 (24%) 2 1	69, 79, 86, 98	0
37	1f	100/101 (99%)	0.72	3 (3%) 52 49	50, 64, 74, 77	0
37	2f	100/101 (99%)	0.77	3 (3%) 52 49	56, 68, 79, 82	0
38	1g	155/156 (99%)	0.84	11 (7%) 22 19	56, 68, 80, 94	0
38	2g	155/156 (99%)	1.32	29 (18%) 3 3	70, 78, 88, 92	0
39	1h	137/138 (99%)	0.77	6 (4%) 39 35	55, 65, 72, 76	0
39	2h	137/138 (99%)	1.35	23 (16%) 4 3	69, 79, 85, 87	0
40	1i	127/128 (99%)	1.28	22 (17%) 4 3	51, 73, 82, 89	0
40	2i	127/128 (99%)	2.26	71 (55%) 0 0	75, 85, 91, 93	0
41	1j	97/105 (92%)	1.40	23 (23%) 2 1	57, 75, 85, 93	0
41	2j	96/105 (91%)	2.48	70 (72%) 0 0	75, 86, 94, 100	0
42	1k	114/129 (88%)	0.70	4 (3%) 47 43	46, 64, 74, 80	0
42	2k	114/129 (88%)	0.96	9 (7%) 18 16	55, 73, 81, 83	0
43	1l	121/132 (91%)	0.36	1 (0%) 82 81	42, 52, 65, 71	0
43	2l	121/132 (91%)	1.13	13 (10%) 11 9	56, 69, 76, 81	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2			OWAB(Å²)	Q<0.9
44	1m	123/126 (97%)	0.96	11 (8%)	15	13	52, 67, 77, 81	0
44	2m	122/126 (96%)	2.07	57 (46%)	0	0	71, 84, 90, 95	0
45	1n	60/61 (98%)	1.13	7 (11%)	9	7	55, 64, 71, 74	0
45	2n	60/61 (98%)	3.15	49 (81%)	0	0	77, 85, 91, 94	0
46	1o	88/89 (98%)	0.69	7 (7%)	18	15	49, 62, 73, 78	0
46	2o	88/89 (98%)	1.01	8 (9%)	15	12	60, 73, 82, 87	0
47	1p	82/88 (93%)	1.44	20 (24%)	2	1	53, 68, 76, 78	0
47	2p	82/88 (93%)	1.31	13 (15%)	5	4	62, 68, 78, 85	0
48	1q	99/105 (94%)	0.89	4 (4%)	42	38	52, 65, 74, 79	0
48	2q	99/105 (94%)	1.11	9 (9%)	15	12	63, 72, 80, 83	0
49	1r	68/88 (77%)	0.66	2 (2%)	53	50	53, 63, 76, 80	0
49	2r	68/88 (77%)	0.97	4 (5%)	28	25	63, 71, 79, 88	0
50	1s	83/93 (89%)	0.82	3 (3%)	46	42	57, 68, 78, 84	0
50	2s	83/93 (89%)	2.36	56 (67%)	0	0	78, 86, 91, 94	0
51	1t	96/106 (90%)	1.23	17 (17%)	4	3	56, 68, 75, 84	0
51	2t	96/106 (90%)	1.22	18 (18%)	3	3	59, 70, 84, 87	0
52	1u	23/27 (85%)	1.28	5 (21%)	2	2	57, 65, 70, 74	0
52	2u	23/27 (85%)	2.86	16 (69%)	0	0	78, 80, 84, 85	0
53	1v	13/24 (54%)	0.48	1 (7%)	19	17	45, 58, 78, 95	0
53	2v	13/24 (54%)	1.73	4 (30%)	1	1	67, 82, 96, 98	0
54	1w	67/76 (88%)	1.33	14 (20%)	2	2	47, 86, 97, 105	0
54	1y	67/76 (88%)	1.12	8 (11%)	9	7	37, 90, 98, 101	0
54	2w	65/76 (85%)	1.78	20 (30%)	1	1	67, 94, 102, 105	0
54	2y	66/76 (86%)	1.47	14 (21%)	2	2	56, 97, 101, 103	0
55	1x	72/77 (93%)	0.60	8 (11%)	10	8	34, 65, 81, 87	0
55	2x	72/77 (93%)	1.03	4 (5%)	30	26	55, 80, 90, 92	0
All	All	20875/21748 (95%)	0.53	1992 (9%)	14	11	16, 63, 90, 107	0

The worst 5 of 1992 RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
45	2n	2	ALA	8.0
45	1n	2	ALA	7.2

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Mol	Chain	Res	Type	RSRZ
6	2G	28	VAL	7.1
21	2Z	172	ALA	6.9
23	21	2	SER	6.5

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

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

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(Å ²)	Q<0.9
54	7MG	2y	46	24/25	0.43	0.19	92,100,108,129	0
54	PSU	2y	55	20/21	0.54	0.18	96,103,115,116	0
54	7MG	2w	46	24/25	0.59	0.15	88,94,107,122	0
54	PSU	1y	55	20/21	0.61	0.17	87,94,103,112	0
54	7MG	1y	46	24/25	0.61	0.16	86,94,105,112	0
54	5MU	1y	54	21/22	0.63	0.15	79,91,96,109	0
54	5MU	2y	54	21/22	0.63	0.16	90,96,106,120	0
54	7MG	1w	46	24/25	0.64	0.18	77,87,101,122	0
54	4SU	2y	8	20/21	0.65	0.13	93,100,109,115	0
54	PSU	2w	55	20/21	0.69	0.14	85,91,101,102	0
54	4SU	2w	8	20/21	0.70	0.13	88,96,108,111	0
54	MIA	2y	37	22/30	0.71	0.16	78,90,104,116	0
54	4SU	1y	8	20/21	0.73	0.13	88,94,109,110	0
54	PSU	2y	32	20/21	0.74	0.15	76,88,94,101	0
54	MIA	1y	37	22/30	0.76	0.13	69,81,86,91	0
54	PSU	2y	39	20/21	0.77	0.15	81,87,93,95	0
32	2MG	2a	1207	24/25	0.81	0.14	79,86,93,95	0
54	4SU	1w	8	20/21	0.81	0.14	75,83,92,101	0
54	5MU	2w	54	21/22	0.82	0.15	75,83,91,93	0
55	PSU	2x	55	20/21	0.82	0.12	76,83,88,93	0
54	PSU	1y	32	20/21	0.83	0.11	75,83,94,95	0
54	PSU	1y	39	20/21	0.84	0.12	70,77,86,87	0
32	M2G	2a	966	25/26	0.85	0.21	57,73,85,89	0
55	4SU	2x	8	20/21	0.86	0.13	77,83,87,91	0
55	5MU	2x	54	21/22	0.86	0.12	80,84,89,90	0
54	PSU	1w	55	20/21	0.86	0.11	63,77,81,85	0
43	0TD	1l	92	10/11	0.87	0.11	42,51,56,70	0
43	0TD	2l	92	10/11	0.88	0.14	62,66,69,79	0
1	PSU	2A	1917	20/21	0.88	0.12	62,70,78,83	0
55	5MU	1x	54	21/22	0.88	0.13	61,71,74,78	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
54	MIA	2w	37	25/30	0.88	0.13	74,80,87,95	0
32	PSU	2a	516	20/21	0.88	0.12	68,76,85,86	0
54	PSU	2w	32	20/21	0.89	0.12	76,85,94,94	0
1	5MU	2A	1915	21/22	0.90	0.11	70,75,80,83	0
32	5MC	2a	967	21/22	0.91	0.15	67,72,78,87	0
54	PSU	2w	39	20/21	0.91	0.10	73,81,88,88	0
1	PSU	2A	1911	20/21	0.91	0.10	61,66,73,74	0
55	5MC	2x	32	21/22	0.91	0.13	68,77,80,81	0
32	4OC	2a	1402	22/23	0.91	0.13	57,65,69,69	0
32	5MC	2a	1404	21/22	0.91	0.13	52,62,66,67	0
1	4OC	2A	1920	21/23	0.91	0.12	60,66,70,73	0
32	7MG	2a	527	24/25	0.92	0.12	63,69,75,81	0
54	PSU	1w	32	20/21	0.92	0.12	59,68,77,78	0
55	PSU	1x	55	20/21	0.92	0.10	51,65,73,78	0
32	5MC	2a	1400	21/22	0.92	0.14	70,74,79,82	0
1	5MU	1A	1937	21/22	0.93	0.10	47,53,59,65	0
32	5MC	2a	1407	21/22	0.93	0.11	51,57,62,66	0
55	4SU	1x	8	20/21	0.93	0.11	52,61,78,79	0
32	MA6	2a	1518	24/25	0.94	0.11	56,69,71,72	0
32	UR3	2a	1498	21/22	0.94	0.11	50,60,63,68	0
32	PSU	1a	516	20/21	0.95	0.09	54,58,61,63	0
54	5MU	1w	54	21/22	0.95	0.09	53,66,71,74	0
32	5MC	1a	967	21/22	0.95	0.11	41,54,60,62	0
55	5MC	1x	32	21/22	0.95	0.12	46,49,55,58	0
32	2MG	1a	1207	24/25	0.95	0.09	52,65,69,73	0
32	5MC	1a	1404	21/22	0.95	0.10	34,40,47,49	0
1	5MC	2A	1942	21/22	0.95	0.10	49,58,65,68	0
1	5MC	2A	1962	21/22	0.95	0.10	41,50,60,78	0
1	PSU	1A	1939	20/21	0.95	0.11	34,49,52,57	0
1	5MC	1A	1964	21/22	0.96	0.09	34,41,45,54	0
1	5MU	2A	1939	21/22	0.96	0.09	36,42,45,46	0
32	7MG	1a	527	24/25	0.96	0.08	42,48,56,59	0
54	PSU	1w	39	20/21	0.96	0.09	48,61,67,68	0
32	M2G	1a	966	25/26	0.96	0.10	45,51,58,60	0
32	MA6	2a	1519	24/25	0.96	0.11	52,62,69,70	0
1	OMG	2A	2251	24/25	0.96	0.09	40,45,48,53	0
32	MA6	1a	1519	24/25	0.96	0.09	36,42,47,49	0
54	MIA	1w	37	29/30	0.96	0.10	39,54,62,68	0
32	5MC	1a	1400	21/22	0.97	0.09	33,48,53,57	0
32	4OC	1a	1402	22/23	0.97	0.08	38,46,49,58	0
1	2MA	2A	2503	23/24	0.97	0.09	30,37,41,46	0
1	2MU	2A	2552	21/23	0.97	0.07	36,43,49,57	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
1	PSU	2A	2605	20/21	0.97	0.07	37,40,46,48	0
1	2MU	1A	2564	21/23	0.97	0.07	22,27,30,32	0
32	5MC	1a	1407	21/22	0.97	0.08	33,40,45,46	0
1	5MU	1A	1961	21/22	0.97	0.07	21,27,32,39	0
1	PSU	1A	1933	20/21	0.97	0.07	33,43,49,50	0
32	MA6	1a	1518	24/25	0.98	0.08	31,41,46,46	0
1	PSU	1A	2617	20/21	0.98	0.06	22,26,30,32	0
1	5MC	1A	1984	21/22	0.98	0.06	28,34,36,43	0
1	OMG	1A	2263	24/25	0.98	0.06	18,23,27,29	0
1	2MA	1A	2515	23/24	0.98	0.06	16,21,24,25	0
1	4OC	1A	1942	21/23	0.98	0.07	38,42,48,49	0
32	UR3	1a	1498	21/22	0.98	0.07	32,39,43,44	0

6.3 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

6.4 Ligands [i](#)

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

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2y	3005	1/1	-0.00	0.22	94,94,94,94	0
56	MG	2j	8001	1/1	0.32	0.20	81,81,81,81	0
56	MG	2a	1826	1/1	0.41	0.23	81,81,81,81	0
56	MG	1A	3623	1/1	0.50	0.22	72,72,72,72	0
56	MG	2A	3664	1/1	0.51	0.22	82,82,82,82	0
56	MG	2A	3285	1/1	0.52	0.15	80,80,80,80	0
56	MG	2a	1840	1/1	0.55	0.13	78,78,78,78	0
56	MG	1a	1777	1/1	0.57	0.18	65,65,65,65	0
56	MG	1A	3210	1/1	0.60	0.31	81,81,81,81	0
56	MG	2A	3725	1/1	0.60	0.20	65,65,65,65	0
56	MG	1A	3782	1/1	0.60	0.14	60,60,60,60	0
56	MG	2a	1837	1/1	0.61	0.39	89,89,89,89	0
56	MG	2w	3004	1/1	0.61	0.17	77,77,77,77	0
56	MG	2A	3023	1/1	0.61	0.21	73,73,73,73	0
56	MG	2A	3423	1/1	0.62	0.15	73,73,73,73	0
56	MG	2A	3471	1/1	0.62	0.25	61,61,61,61	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2B	3019	1/1	0.62	0.18	90,90,90,90	0
56	MG	2v	103	1/1	0.63	0.29	78,78,78,78	0
56	MG	2A	3402	1/1	0.63	0.19	80,80,80,80	0
56	MG	2a	1821	1/1	0.63	0.25	75,75,75,75	0
56	MG	2A	3100	1/1	0.64	0.17	77,77,77,77	0
56	MG	1a	1667	1/1	0.64	0.27	69,69,69,69	0
56	MG	2A	3137	1/1	0.65	0.16	61,61,61,61	0
56	MG	2A	3642	1/1	0.65	0.22	45,45,45,45	0
56	MG	2B	3016	1/1	0.65	0.25	70,70,70,70	0
56	MG	2Q	3004	1/1	0.66	0.35	58,58,58,58	0
56	MG	1y	3002	1/1	0.67	0.20	90,90,90,90	0
56	MG	2A	3382	1/1	0.67	0.25	75,75,75,75	0
56	MG	2A	3766	1/1	0.67	0.30	83,83,83,83	0
56	MG	2A	3812	1/1	0.67	0.20	93,93,93,93	0
56	MG	2A	3032	1/1	0.67	0.19	66,66,66,66	0
56	MG	2A	3705	1/1	0.68	0.14	78,78,78,78	0
56	MG	2A	3500	1/1	0.69	0.26	69,69,69,69	0
56	MG	1a	1668	1/1	0.69	0.18	68,68,68,68	0
56	MG	1A	3446	1/1	0.69	0.19	67,67,67,67	0
56	MG	2a	1655	1/1	0.69	0.17	75,75,75,75	0
56	MG	2A	3683	1/1	0.69	0.25	77,77,77,77	0
56	MG	2A	3818	1/1	0.69	0.15	70,70,70,70	0
56	MG	1a	1766	1/1	0.70	0.19	86,86,86,86	0
56	MG	2a	1714	1/1	0.70	0.29	79,79,79,79	0
56	MG	1A	3507	1/1	0.70	0.25	71,71,71,71	0
56	MG	1B	226	1/1	0.70	0.15	69,69,69,69	0
56	MG	1a	1759	1/1	0.70	0.15	81,81,81,81	0
56	MG	1A	3905	1/1	0.71	0.23	72,72,72,72	0
56	MG	2A	3830	1/1	0.71	0.15	54,54,54,54	0
56	MG	2A	3871	1/1	0.71	0.17	66,66,66,66	0
56	MG	2a	1722	1/1	0.71	0.20	66,66,66,66	0
56	MG	2A	3243	1/1	0.71	0.21	67,67,67,67	0
56	MG	2y	3001	1/1	0.71	0.19	76,76,76,76	0
56	MG	1A	4011	1/1	0.71	0.24	69,69,69,69	0
56	MG	2A	3665	1/1	0.72	0.20	69,69,69,69	0
56	MG	2a	1824	1/1	0.72	0.19	70,70,70,70	0
56	MG	2B	3021	1/1	0.72	0.29	79,79,79,79	0
56	MG	1y	3004	1/1	0.72	0.15	83,83,83,83	0
56	MG	2U	202	1/1	0.72	0.34	72,72,72,72	0
56	MG	2a	1625	1/1	0.72	0.16	80,80,80,80	0
56	MG	2A	3394	1/1	0.72	0.27	72,72,72,72	0
56	MG	2a	1670	1/1	0.72	0.18	62,62,62,62	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3907	1/1	0.72	0.21	77,77,77,77	0
56	MG	2A	3101	1/1	0.72	0.33	76,76,76,76	0
56	MG	1A	3985	1/1	0.73	0.18	78,78,78,78	0
56	MG	1A	3259	1/1	0.73	0.17	57,57,57,57	0
56	MG	1A	4066	1/1	0.73	0.13	51,51,51,51	0
56	MG	1v	3001	1/1	0.73	0.22	70,70,70,70	0
56	MG	2a	1814	1/1	0.73	0.14	62,62,62,62	0
56	MG	2A	3713	1/1	0.74	0.16	67,67,67,67	0
56	MG	2A	3416	1/1	0.74	0.17	63,63,63,63	0
56	MG	2A	3656	1/1	0.74	0.21	80,80,80,80	0
56	MG	1A	4110	1/1	0.74	0.34	49,49,49,49	0
56	MG	2A	3395	1/1	0.74	0.11	67,67,67,67	0
56	MG	1A	3405	1/1	0.74	0.21	45,45,45,45	0
56	MG	2A	3505	1/1	0.74	0.18	65,65,65,65	0
56	MG	1A	3499	1/1	0.75	0.16	55,55,55,55	0
56	MG	1A	3531	1/1	0.75	0.29	60,60,60,60	0
56	MG	1x	102	1/1	0.75	0.37	66,66,66,66	0
56	MG	1a	1680	1/1	0.75	0.22	76,76,76,76	0
56	MG	1A	3955	1/1	0.75	0.16	55,55,55,55	0
56	MG	2A	3244	1/1	0.75	0.22	74,74,74,74	0
56	MG	2A	3259	1/1	0.75	0.27	65,65,65,65	0
56	MG	1A	3801	1/1	0.75	0.11	47,47,47,47	0
56	MG	2A	3333	1/1	0.75	0.19	61,61,61,61	0
56	MG	2A	3579	1/1	0.75	0.22	65,65,65,65	0
56	MG	2A	3360	1/1	0.75	0.30	75,75,75,75	0
56	MG	2a	1715	1/1	0.76	0.19	70,70,70,70	0
56	MG	2A	3091	1/1	0.76	0.16	69,69,69,69	0
56	MG	2a	1726	1/1	0.76	0.20	71,71,71,71	0
56	MG	2a	1794	1/1	0.76	0.18	75,75,75,75	0
56	MG	2A	3879	1/1	0.76	0.16	52,52,52,52	0
56	MG	2A	3884	1/1	0.76	0.49	66,66,66,66	0
56	MG	2A	3453	1/1	0.76	0.33	64,64,64,64	0
56	MG	1A	3968	1/1	0.76	0.27	68,68,68,68	0
56	MG	1A	3645	1/1	0.76	0.14	47,47,47,47	0
56	MG	1a	1652	1/1	0.76	0.13	56,56,56,56	0
56	MG	2e	3001	1/1	0.76	0.12	76,76,76,76	0
56	MG	2A	3201	1/1	0.76	0.35	71,71,71,71	0
56	MG	1A	3406	1/1	0.76	0.32	68,68,68,68	0
56	MG	1A	3371	1/1	0.76	0.27	59,59,59,59	0
56	MG	2A	3054	1/1	0.76	0.29	73,73,73,73	0
56	MG	2A	3835	1/1	0.76	0.15	58,58,58,58	0
56	MG	2a	1756	1/1	0.77	0.10	88,88,88,88	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3719	1/1	0.77	0.22	61,61,61,61	0
56	MG	2A	3249	1/1	0.77	0.16	62,62,62,62	0
56	MG	1A	3917	1/1	0.77	0.13	50,50,50,50	0
56	MG	2A	3175	1/1	0.77	0.12	84,84,84,84	0
56	MG	1A	4112	1/1	0.77	0.17	62,62,62,62	0
56	MG	2a	1653	1/1	0.77	0.26	82,82,82,82	0
56	MG	2A	3346	1/1	0.77	0.26	69,69,69,69	0
56	MG	2A	3667	1/1	0.77	0.16	62,62,62,62	0
56	MG	2a	1700	1/1	0.77	0.13	59,59,59,59	0
56	MG	2A	3214	1/1	0.77	0.17	64,64,64,64	0
56	MG	2w	3002	1/1	0.77	0.19	77,77,77,77	0
56	MG	2A	3687	1/1	0.77	0.17	69,69,69,69	0
56	MG	2w	3005	1/1	0.77	0.13	76,76,76,76	0
56	MG	1A	4034	1/1	0.77	0.30	64,64,64,64	0
56	MG	1A	3991	1/1	0.77	0.15	32,32,32,32	0
56	MG	2a	1649	1/1	0.78	0.17	72,72,72,72	0
56	MG	1A	3525	1/1	0.78	0.21	70,70,70,70	0
56	MG	2A	3051	1/1	0.78	0.15	69,69,69,69	0
56	MG	2B	3011	1/1	0.78	0.26	72,72,72,72	0
56	MG	1a	1745	1/1	0.78	0.22	80,80,80,80	0
56	MG	2a	1703	1/1	0.78	0.32	64,64,64,64	0
56	MG	1a	1772	1/1	0.78	0.15	66,66,66,66	0
56	MG	2l	3003	1/1	0.78	0.19	74,74,74,74	0
56	MG	2t	3001	1/1	0.78	0.28	62,62,62,62	0
56	MG	2A	3520	1/1	0.78	0.24	69,69,69,69	0
56	MG	2A	3347	1/1	0.78	0.23	68,68,68,68	0
56	MG	2A	3606	1/1	0.78	0.23	65,65,65,65	0
56	MG	2Z	8001	1/1	0.78	0.17	73,73,73,73	0
56	MG	2A	3181	1/1	0.78	0.31	76,76,76,76	0
56	MG	2a	1634	1/1	0.78	0.17	85,85,85,85	0
56	MG	2A	3327	1/1	0.79	0.32	60,60,60,60	0
56	MG	2A	3171	1/1	0.79	0.17	73,73,73,73	0
56	MG	1A	4070	1/1	0.79	0.19	45,45,45,45	0
56	MG	2A	3880	1/1	0.79	0.18	65,65,65,65	0
56	MG	1A	3292	1/1	0.79	0.18	60,60,60,60	0
56	MG	2a	1768	1/1	0.79	0.15	82,82,82,82	0
56	MG	2a	1769	1/1	0.79	0.11	81,81,81,81	0
56	MG	2A	3886	1/1	0.79	0.33	54,54,54,54	0
56	MG	2A	3357	1/1	0.79	0.11	70,70,70,70	0
56	MG	1a	1773	1/1	0.79	0.14	74,74,74,74	0
56	MG	1A	3931	1/1	0.79	0.17	36,36,36,36	0
56	MG	2A	3215	1/1	0.79	0.13	71,71,71,71	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2a	1832	1/1	0.79	0.27	75,75,75,75	0
56	MG	2A	3224	1/1	0.79	0.15	69,69,69,69	0
56	MG	1a	1691	1/1	0.79	0.20	64,64,64,64	0
56	MG	1a	1716	1/1	0.79	0.21	58,58,58,58	0
56	MG	1A	4059	1/1	0.79	0.23	51,51,51,51	0
56	MG	2A	3102	1/1	0.79	0.20	72,72,72,72	0
56	MG	2A	3281	1/1	0.79	0.08	81,81,81,81	0
56	MG	2A	3477	1/1	0.79	0.30	78,78,78,78	0
56	MG	2A	3767	1/1	0.79	0.12	54,54,54,54	0
56	MG	2A	3488	1/1	0.79	0.16	71,71,71,71	0
56	MG	1A	3779	1/1	0.79	0.12	23,23,23,23	0
56	MG	2A	3286	1/1	0.79	0.16	67,67,67,67	0
56	MG	2a	1708	1/1	0.79	0.34	75,75,75,75	0
56	MG	2a	1795	1/1	0.80	0.12	77,77,77,77	0
56	MG	2A	3324	1/1	0.80	0.19	69,69,69,69	0
56	MG	1Z	303	1/1	0.80	0.08	65,65,65,65	0
56	MG	2A	3696	1/1	0.80	0.10	79,79,79,79	0
56	MG	2a	1658	1/1	0.80	0.10	73,73,73,73	0
56	MG	2a	1830	1/1	0.80	0.22	59,59,59,59	0
56	MG	1A	3535	1/1	0.80	0.17	57,57,57,57	0
56	MG	2A	3590	1/1	0.80	0.17	72,72,72,72	0
56	MG	2A	3603	1/1	0.80	0.19	67,67,67,67	0
56	MG	1a	1701	1/1	0.80	0.16	59,59,59,59	0
56	MG	2A	3625	1/1	0.80	0.19	63,63,63,63	0
56	MG	2A	3153	1/1	0.80	0.23	69,69,69,69	0
56	MG	1B	207	1/1	0.80	0.28	69,69,69,69	0
56	MG	1A	3454	1/1	0.80	0.22	61,61,61,61	0
56	MG	2a	1732	1/1	0.80	0.13	94,94,94,94	0
56	MG	2A	3292	1/1	0.80	0.13	65,65,65,65	0
56	MG	2A	3305	1/1	0.80	0.25	67,67,67,67	0
56	MG	2A	3856	1/1	0.80	0.16	58,58,58,58	0
56	MG	2a	1639	1/1	0.80	0.35	68,68,68,68	0
56	MG	1A	3058	1/1	0.81	0.18	69,69,69,69	0
56	MG	1A	3415	1/1	0.81	0.22	61,61,61,61	0
56	MG	1A	3428	1/1	0.81	0.28	68,68,68,68	0
56	MG	2A	3428	1/1	0.81	0.27	79,79,79,79	0
56	MG	2A	3345	1/1	0.81	0.13	59,59,59,59	0
56	MG	1A	3830	1/1	0.81	0.14	49,49,49,49	0
56	MG	1A	3986	1/1	0.81	0.10	59,59,59,59	0
56	MG	2A	3062	1/1	0.81	0.16	67,67,67,67	0
56	MG	2A	3681	1/1	0.81	0.15	56,56,56,56	0
56	MG	2a	1834	1/1	0.81	0.27	81,81,81,81	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3197	1/1	0.81	0.13	72,72,72,72	0
56	MG	2A	3881	1/1	0.81	0.15	69,69,69,69	0
56	MG	2A	3362	1/1	0.81	0.14	72,72,72,72	0
56	MG	2A	3693	1/1	0.81	0.15	58,58,58,58	0
56	MG	1A	3214	1/1	0.81	0.16	46,46,46,46	0
56	MG	2a	1721	1/1	0.81	0.18	76,76,76,76	0
56	MG	2A	3529	1/1	0.81	0.21	64,64,64,64	0
56	MG	2A	3573	1/1	0.81	0.12	48,48,48,48	0
56	MG	2w	3003	1/1	0.81	0.17	74,74,74,74	0
56	MG	1A	3664	1/1	0.81	0.13	66,66,66,66	0
56	MG	1A	3730	1/1	0.81	0.14	58,58,58,58	0
56	MG	2a	1761	1/1	0.81	0.12	95,95,95,95	0
56	MG	2A	3751	1/1	0.81	0.25	77,77,77,77	0
56	MG	1B	201	1/1	0.82	0.16	57,57,57,57	0
56	MG	2A	3758	1/1	0.82	0.22	64,64,64,64	0
56	MG	2A	3506	1/1	0.82	0.13	47,47,47,47	0
56	MG	1A	3723	1/1	0.82	0.20	47,47,47,47	0
56	MG	2A	3189	1/1	0.82	0.26	65,65,65,65	0
56	MG	2A	3545	1/1	0.82	0.16	58,58,58,58	0
56	MG	1A	3891	1/1	0.82	0.16	53,53,53,53	0
56	MG	1P	203	1/1	0.82	0.18	84,84,84,84	0
56	MG	2A	3204	1/1	0.82	0.29	67,67,67,67	0
56	MG	2A	3350	1/1	0.82	0.15	63,63,63,63	0
56	MG	1U	204	1/1	0.82	0.47	74,74,74,74	0
56	MG	2a	1766	1/1	0.82	0.17	87,87,87,87	0
56	MG	2A	3359	1/1	0.82	0.12	63,63,63,63	0
56	MG	2A	3628	1/1	0.82	0.11	58,58,58,58	0
56	MG	1A	3460	1/1	0.82	0.17	53,53,53,53	0
56	MG	2A	3649	1/1	0.82	0.16	84,84,84,84	0
56	MG	2A	3216	1/1	0.82	0.16	77,77,77,77	0
56	MG	1a	1651	1/1	0.82	0.36	75,75,75,75	0
56	MG	1A	3563	1/1	0.82	0.29	64,64,64,64	0
56	MG	2A	3097	1/1	0.82	0.24	74,74,74,74	0
56	MG	1a	1655	1/1	0.82	0.26	63,63,63,63	0
56	MG	2A	3257	1/1	0.82	0.32	63,63,63,63	0
56	MG	1a	1664	1/1	0.82	0.27	74,74,74,74	0
56	MG	2a	1606	1/1	0.82	0.21	58,58,58,58	0
56	MG	2a	1608	1/1	0.82	0.20	63,63,63,63	0
56	MG	2a	1843	1/1	0.82	0.23	66,66,66,66	0
56	MG	1s	3001	1/1	0.82	0.15	63,63,63,63	0
56	MG	1A	3582	1/1	0.82	0.13	49,49,49,49	0
56	MG	2j	8002	1/1	0.82	0.08	82,82,82,82	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2a	1635	1/1	0.82	0.23	71,71,71,71	0
56	MG	2A	3704	1/1	0.82	0.10	75,75,75,75	0
56	MG	1A	3666	1/1	0.82	0.11	43,43,43,43	0
56	MG	2A	3710	1/1	0.82	0.20	69,69,69,69	0
56	MG	2a	1654	1/1	0.82	0.16	85,85,85,85	0
56	MG	1x	107	1/1	0.82	0.18	66,66,66,66	0
56	MG	2A	3173	1/1	0.82	0.29	77,77,77,77	0
56	MG	2w	3008	1/1	0.82	0.10	75,75,75,75	0
56	MG	2A	3308	1/1	0.82	0.21	63,63,63,63	0
56	MG	2a	1672	1/1	0.82	0.25	68,68,68,68	0
56	MG	2A	3435	1/1	0.83	0.25	58,58,58,58	0
56	MG	2A	3445	1/1	0.83	0.28	57,57,57,57	0
56	MG	1Y	203	1/1	0.83	0.11	66,66,66,66	0
56	MG	2a	1656	1/1	0.83	0.11	59,59,59,59	0
56	MG	1a	1758	1/1	0.83	0.13	81,81,81,81	0
56	MG	2A	3265	1/1	0.83	0.14	56,56,56,56	0
56	MG	2A	3484	1/1	0.83	0.11	67,67,67,67	0
56	MG	2a	1696	1/1	0.83	0.29	76,76,76,76	0
56	MG	2A	3752	1/1	0.83	0.14	59,59,59,59	0
56	MG	2A	3753	1/1	0.83	0.16	64,64,64,64	0
56	MG	2a	1704	1/1	0.83	0.18	65,65,65,65	0
56	MG	2A	3755	1/1	0.83	0.12	71,71,71,71	0
56	MG	2A	3757	1/1	0.83	0.15	50,50,50,50	0
56	MG	1A	3017	1/1	0.83	0.14	34,34,34,34	0
56	MG	18	101	1/1	0.83	0.19	65,65,65,65	0
56	MG	1a	1604	1/1	0.83	0.14	55,55,55,55	0
56	MG	2A	3795	1/1	0.83	0.20	85,85,85,85	0
56	MG	2A	3797	1/1	0.83	0.12	73,73,73,73	0
56	MG	2a	1739	1/1	0.83	0.17	84,84,84,84	0
56	MG	2a	1742	1/1	0.83	0.14	74,74,74,74	0
56	MG	2a	1745	1/1	0.83	0.09	96,96,96,96	0
56	MG	2A	3107	1/1	0.83	0.14	62,62,62,62	0
56	MG	1a	1639	1/1	0.83	0.13	59,59,59,59	0
56	MG	1A	3753	1/1	0.83	0.12	20,20,20,20	0
56	MG	2a	1767	1/1	0.83	0.19	79,79,79,79	0
56	MG	2A	3323	1/1	0.83	0.10	68,68,68,68	0
56	MG	2A	3852	1/1	0.83	0.19	46,46,46,46	0
56	MG	2A	3556	1/1	0.83	0.15	53,53,53,53	0
56	MG	1a	1811	1/1	0.83	0.14	63,63,63,63	0
56	MG	1r	101	1/1	0.83	0.27	69,69,69,69	0
56	MG	2a	1817	1/1	0.83	0.19	59,59,59,59	0
56	MG	1A	3586	1/1	0.83	0.15	51,51,51,51	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3343	1/1	0.83	0.12	74,74,74,74	0
56	MG	1A	3355	1/1	0.83	0.13	66,66,66,66	0
56	MG	2A	3607	1/1	0.83	0.16	53,53,53,53	0
56	MG	2B	3009	1/1	0.83	0.32	66,66,66,66	0
56	MG	1w	106	1/1	0.83	0.14	66,66,66,66	0
56	MG	2A	3196	1/1	0.83	0.15	69,69,69,69	0
56	MG	1A	3631	1/1	0.83	0.15	52,52,52,52	0
56	MG	1A	3526	1/1	0.83	0.18	85,85,85,85	0
56	MG	1A	3463	1/1	0.83	0.10	62,62,62,62	0
56	MG	2A	3207	1/1	0.83	0.14	55,55,55,55	0
56	MG	2A	3212	1/1	0.83	0.12	52,52,52,52	0
56	MG	1y	3003	1/1	0.83	0.30	82,82,82,82	0
56	MG	1a	1676	1/1	0.83	0.36	72,72,72,72	0
56	MG	1A	3360	1/1	0.83	0.18	57,57,57,57	0
56	MG	1A	3503	1/1	0.83	0.09	54,54,54,54	0
56	MG	1S	3002	1/1	0.83	0.19	49,49,49,49	0
56	MG	2a	1636	1/1	0.83	0.14	81,81,81,81	0
56	MG	1A	4036	1/1	0.83	0.10	34,34,34,34	0
56	MG	2a	1640	1/1	0.83	0.11	68,68,68,68	0
56	MG	2a	1645	1/1	0.83	0.24	70,70,70,70	0
56	MG	1a	1729	1/1	0.83	0.16	79,79,79,79	0
56	MG	1a	1733	1/1	0.84	0.15	55,55,55,55	0
56	MG	1w	110	1/1	0.84	0.19	69,69,69,69	0
56	MG	1G	3004	1/1	0.84	0.12	67,67,67,67	0
56	MG	2A	3338	1/1	0.84	0.21	64,64,64,64	0
56	MG	2A	3493	1/1	0.84	0.27	74,74,74,74	0
56	MG	1a	1754	1/1	0.84	0.23	65,65,65,65	0
56	MG	2A	3217	1/1	0.84	0.25	63,63,63,63	0
56	MG	2A	3112	1/1	0.84	0.17	59,59,59,59	0
56	MG	2A	3508	1/1	0.84	0.19	61,61,61,61	0
56	MG	1x	113	1/1	0.84	0.28	69,69,69,69	0
56	MG	1A	3846	1/1	0.84	0.12	58,58,58,58	0
56	MG	1A	3858	1/1	0.84	0.22	47,47,47,47	0
56	MG	2a	1609	1/1	0.84	0.26	71,71,71,71	0
56	MG	2a	1791	1/1	0.84	0.18	71,71,71,71	0
56	MG	2a	1793	1/1	0.84	0.26	71,71,71,71	0
56	MG	1A	3051	1/1	0.84	0.34	66,66,66,66	0
56	MG	1A	4072	1/1	0.84	0.14	32,32,32,32	0
56	MG	2A	3262	1/1	0.84	0.15	69,69,69,69	0
56	MG	1A	3899	1/1	0.84	0.15	37,37,37,37	0
56	MG	2A	3185	1/1	0.84	0.18	55,55,55,55	0
56	MG	1A	3470	1/1	0.84	0.16	63,63,63,63	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3787	1/1	0.84	0.13	59,59,59,59	0
56	MG	2a	1829	1/1	0.84	0.27	77,77,77,77	0
56	MG	2A	3793	1/1	0.84	0.14	60,60,60,60	0
56	MG	1A	3191	1/1	0.84	0.12	50,50,50,50	0
56	MG	2A	3619	1/1	0.84	0.15	50,50,50,50	0
56	MG	2A	3804	1/1	0.84	0.15	61,61,61,61	0
56	MG	2A	3409	1/1	0.84	0.21	69,69,69,69	0
56	MG	1a	1637	1/1	0.84	0.15	68,68,68,68	0
56	MG	2d	502	1/1	0.84	0.28	74,74,74,74	0
56	MG	2a	1667	1/1	0.84	0.11	59,59,59,59	0
56	MG	2a	1669	1/1	0.84	0.29	59,59,59,59	0
56	MG	2A	3422	1/1	0.84	0.13	62,62,62,62	0
56	MG	2A	3303	1/1	0.84	0.17	68,68,68,68	0
56	MG	2a	1674	1/1	0.84	0.19	58,58,58,58	0
56	MG	2A	3090	1/1	0.84	0.20	76,76,76,76	0
56	MG	1A	3829	1/1	0.84	0.13	61,61,61,61	0
56	MG	2A	3862	1/1	0.84	0.19	84,84,84,84	0
56	MG	2A	3315	1/1	0.84	0.11	65,65,65,65	0
56	MG	2a	1705	1/1	0.84	0.14	86,86,86,86	0
56	MG	2A	3447	1/1	0.84	0.14	49,49,49,49	0
56	MG	2w	3009	1/1	0.84	0.17	73,73,73,73	0
56	MG	2x	104	1/1	0.84	0.15	68,68,68,68	0
56	MG	2x	105	1/1	0.84	0.23	66,66,66,66	0
56	MG	2A	3675	1/1	0.84	0.17	62,62,62,62	0
56	MG	1A	3618	1/1	0.84	0.14	50,50,50,50	0
56	MG	1a	1678	1/1	0.85	0.07	54,54,54,54	0
56	MG	1A	3638	1/1	0.85	0.16	48,48,48,48	0
56	MG	1A	3459	1/1	0.85	0.13	53,53,53,53	0
56	MG	2a	1711	1/1	0.85	0.30	64,64,64,64	0
56	MG	1A	3930	1/1	0.85	0.12	23,23,23,23	0
56	MG	2A	3671	1/1	0.85	0.10	63,63,63,63	0
56	MG	2a	1719	1/1	0.85	0.18	67,67,67,67	0
56	MG	2A	3450	1/1	0.85	0.20	53,53,53,53	0
56	MG	2A	3680	1/1	0.85	0.24	78,78,78,78	0
56	MG	1A	3262	1/1	0.85	0.25	55,55,55,55	0
56	MG	2a	1729	1/1	0.85	0.20	76,76,76,76	0
56	MG	1x	101	1/1	0.85	0.13	51,51,51,51	0
56	MG	2a	1734	1/1	0.85	0.19	62,62,62,62	0
56	MG	2A	3686	1/1	0.85	0.13	71,71,71,71	0
56	MG	1A	3440	1/1	0.85	0.18	63,63,63,63	0
56	MG	1a	1629	1/1	0.85	0.18	64,64,64,64	0
56	MG	2B	3014	1/1	0.85	0.22	78,78,78,78	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3342	1/1	0.85	0.12	65,65,65,65	0
56	MG	2A	3702	1/1	0.85	0.15	75,75,75,75	0
56	MG	2A	3227	1/1	0.85	0.20	57,57,57,57	0
56	MG	2E	304	1/1	0.85	0.30	70,70,70,70	0
56	MG	2A	3232	1/1	0.85	0.15	65,65,65,65	0
56	MG	2a	1785	1/1	0.85	0.26	70,70,70,70	0
56	MG	2A	3236	1/1	0.85	0.17	64,64,64,64	0
56	MG	1A	3966	1/1	0.85	0.15	75,75,75,75	0
56	MG	2a	1605	1/1	0.85	0.15	73,73,73,73	0
56	MG	1A	3668	1/1	0.85	0.10	56,56,56,56	0
56	MG	2a	1812	1/1	0.85	0.21	62,62,62,62	0
56	MG	2A	3355	1/1	0.85	0.14	79,79,79,79	0
56	MG	2A	3735	1/1	0.85	0.12	69,69,69,69	0
56	MG	2a	1820	1/1	0.85	0.15	58,58,58,58	0
56	MG	2a	1613	1/1	0.85	0.12	61,61,61,61	0
56	MG	2a	1823	1/1	0.85	0.09	72,72,72,72	0
56	MG	2a	1615	1/1	0.85	0.29	69,69,69,69	0
56	MG	2A	3736	1/1	0.85	0.20	69,69,69,69	0
56	MG	2a	1629	1/1	0.85	0.19	80,80,80,80	0
56	MG	2a	1630	1/1	0.85	0.17	69,69,69,69	0
56	MG	2A	3161	1/1	0.85	0.20	71,71,71,71	0
56	MG	1A	3984	1/1	0.85	0.23	81,81,81,81	0
56	MG	2A	3553	1/1	0.85	0.10	56,56,56,56	0
56	MG	1A	3690	1/1	0.85	0.12	43,43,43,43	0
56	MG	1A	3716	1/1	0.85	0.11	65,65,65,65	0
56	MG	2A	3366	1/1	0.85	0.15	61,61,61,61	0
56	MG	1A	3266	1/1	0.85	0.14	60,60,60,60	0
56	MG	2a	1650	1/1	0.85	0.15	68,68,68,68	0
56	MG	1O	207	1/1	0.85	0.15	67,67,67,67	0
56	MG	2A	3777	1/1	0.85	0.14	76,76,76,76	0
56	MG	2A	3283	1/1	0.85	0.14	69,69,69,69	0
56	MG	1A	3488	1/1	0.85	0.15	50,50,50,50	0
56	MG	1A	3257	1/1	0.85	0.31	61,61,61,61	0
56	MG	2A	3415	1/1	0.85	0.14	70,70,70,70	0
56	MG	2A	3074	1/1	0.85	0.15	66,66,66,66	0
56	MG	2A	3809	1/1	0.85	0.14	78,78,78,78	0
56	MG	2A	3088	1/1	0.85	0.20	72,72,72,72	0
56	MG	2A	3817	1/1	0.85	0.12	84,84,84,84	0
56	MG	2a	1678	1/1	0.85	0.15	72,72,72,72	0
56	MG	2A	3644	1/1	0.85	0.19	62,62,62,62	0
56	MG	2A	3824	1/1	0.85	0.11	80,80,80,80	0
56	MG	2A	3304	1/1	0.85	0.08	55,55,55,55	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3433	1/1	0.86	0.14	64,64,64,64	0
56	MG	1a	1756	1/1	0.86	0.09	87,87,87,87	0
56	MG	2A	3255	1/1	0.86	0.34	71,71,71,71	0
56	MG	10	106	1/1	0.86	0.12	58,58,58,58	0
56	MG	1A	4026	1/1	0.86	0.14	52,52,52,52	0
56	MG	1A	3872	1/1	0.86	0.13	28,28,28,28	0
56	MG	1a	1617	1/1	0.86	0.23	65,65,65,65	0
56	MG	2A	3269	1/1	0.86	0.24	63,63,63,63	0
56	MG	2a	1679	1/1	0.86	0.20	71,71,71,71	0
56	MG	2a	1680	1/1	0.86	0.19	64,64,64,64	0
56	MG	1A	3555	1/1	0.86	0.16	68,68,68,68	0
56	MG	2A	3769	1/1	0.86	0.12	56,56,56,56	0
56	MG	2A	3776	1/1	0.86	0.13	68,68,68,68	0
56	MG	1A	3699	1/1	0.86	0.13	53,53,53,53	0
56	MG	2A	3489	1/1	0.86	0.15	73,73,73,73	0
56	MG	2a	1707	1/1	0.86	0.19	76,76,76,76	0
56	MG	2A	3491	1/1	0.86	0.14	67,67,67,67	0
56	MG	1a	1797	1/1	0.86	0.08	66,66,66,66	0
56	MG	2A	3109	1/1	0.86	0.23	68,68,68,68	0
56	MG	2A	3501	1/1	0.86	0.14	62,62,62,62	0
56	MG	1a	1810	1/1	0.86	0.31	61,61,61,61	0
56	MG	2A	3302	1/1	0.86	0.16	69,69,69,69	0
56	MG	2A	3126	1/1	0.86	0.08	63,63,63,63	0
56	MG	2A	3518	1/1	0.86	0.23	67,67,67,67	0
56	MG	2A	3136	1/1	0.86	0.13	46,46,46,46	0
56	MG	1A	3392	1/1	0.86	0.12	48,48,48,48	0
56	MG	2A	3306	1/1	0.86	0.23	67,67,67,67	0
56	MG	2A	3152	1/1	0.86	0.18	55,55,55,55	0
56	MG	2A	3312	1/1	0.86	0.26	67,67,67,67	0
56	MG	2A	3568	1/1	0.86	0.10	34,34,34,34	0
56	MG	2a	1751	1/1	0.86	0.10	51,51,51,51	0
56	MG	1h	3001	1/1	0.86	0.10	64,64,64,64	0
56	MG	2A	3321	1/1	0.86	0.15	68,68,68,68	0
56	MG	2A	3154	1/1	0.86	0.20	72,72,72,72	0
56	MG	1o	3001	1/1	0.86	0.18	64,64,64,64	0
56	MG	1q	201	1/1	0.86	0.11	59,59,59,59	0
56	MG	2A	3172	1/1	0.86	0.17	66,66,66,66	0
56	MG	2a	1784	1/1	0.86	0.19	84,84,84,84	0
56	MG	2A	3905	1/1	0.86	0.12	56,56,56,56	0
56	MG	2a	1787	1/1	0.86	0.29	77,77,77,77	0
56	MG	2A	3615	1/1	0.86	0.10	49,49,49,49	0
56	MG	2A	3336	1/1	0.86	0.13	67,67,67,67	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3622	1/1	0.86	0.27	67,67,67,67	0
56	MG	1A	3449	1/1	0.86	0.07	57,57,57,57	0
56	MG	2a	1807	1/1	0.86	0.09	89,89,89,89	0
56	MG	2a	1809	1/1	0.86	0.18	72,72,72,72	0
56	MG	2a	1811	1/1	0.86	0.14	68,68,68,68	0
56	MG	1A	3298	1/1	0.86	0.18	44,44,44,44	0
56	MG	1A	3264	1/1	0.86	0.07	50,50,50,50	0
56	MG	1w	102	1/1	0.86	0.15	73,73,73,73	0
56	MG	2G	3001	1/1	0.86	0.18	62,62,62,62	0
56	MG	1a	1662	1/1	0.86	0.15	67,67,67,67	0
56	MG	2A	3194	1/1	0.86	0.12	65,65,65,65	0
56	MG	2V	202	1/1	0.86	0.14	61,61,61,61	0
56	MG	1A	3772	1/1	0.86	0.18	71,71,71,71	0
56	MG	2a	1828	1/1	0.86	0.13	68,68,68,68	0
56	MG	1A	3511	1/1	0.86	0.12	43,43,43,43	0
56	MG	1A	3356	1/1	0.86	0.23	55,55,55,55	0
56	MG	1A	3800	1/1	0.86	0.12	52,52,52,52	0
56	MG	1a	1677	1/1	0.86	0.14	68,68,68,68	0
56	MG	2a	1612	1/1	0.86	0.22	64,64,64,64	0
56	MG	1D	311	1/1	0.86	0.30	43,43,43,43	0
56	MG	2a	1841	1/1	0.86	0.29	89,89,89,89	0
56	MG	1A	3221	1/1	0.86	0.32	71,71,71,71	0
56	MG	2a	1622	1/1	0.86	0.09	65,65,65,65	0
56	MG	1A	3025	1/1	0.86	0.23	73,73,73,73	0
56	MG	2a	1627	1/1	0.86	0.22	71,71,71,71	0
56	MG	2A	3390	1/1	0.86	0.12	54,54,54,54	0
56	MG	1O	208	1/1	0.86	0.13	75,75,75,75	0
56	MG	2a	1633	1/1	0.86	0.21	57,57,57,57	0
56	MG	2A	3691	1/1	0.86	0.14	59,59,59,59	0
56	MG	2A	3025	1/1	0.86	0.17	65,65,65,65	0
56	MG	1A	3487	1/1	0.86	0.23	55,55,55,55	0
56	MG	1A	3548	1/1	0.86	0.26	56,56,56,56	0
56	MG	1A	3997	1/1	0.86	0.11	58,58,58,58	0
56	MG	1A	3553	1/1	0.86	0.10	58,58,58,58	0
56	MG	2A	3421	1/1	0.86	0.09	52,52,52,52	0
56	MG	2A	3242	1/1	0.86	0.14	67,67,67,67	0
56	MG	2A	3064	1/1	0.86	0.15	76,76,76,76	0
56	MG	2x	106	1/1	0.86	0.24	62,62,62,62	0
56	MG	1A	4023	1/1	0.86	0.12	68,68,68,68	0
56	MG	2y	3002	1/1	0.86	0.13	63,63,63,63	0
56	MG	2A	3429	1/1	0.86	0.08	62,62,62,62	0
56	MG	1A	3832	1/1	0.87	0.27	49,49,49,49	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3306	1/1	0.87	0.17	56,56,56,56	0
56	MG	2A	3096	1/1	0.87	0.11	60,60,60,60	0
56	MG	2A	3391	1/1	0.87	0.11	70,70,70,70	0
56	MG	1A	3103	1/1	0.87	0.07	23,23,23,23	0
56	MG	1A	3868	1/1	0.87	0.10	30,30,30,30	0
56	MG	1a	1614	1/1	0.87	0.11	48,48,48,48	0
56	MG	2a	1713	1/1	0.87	0.18	76,76,76,76	0
56	MG	1A	3707	1/1	0.87	0.16	78,78,78,78	0
56	MG	2A	3412	1/1	0.87	0.18	60,60,60,60	0
56	MG	2a	1716	1/1	0.87	0.21	63,63,63,63	0
56	MG	1a	1625	1/1	0.87	0.30	62,62,62,62	0
56	MG	2A	3883	1/1	0.87	0.12	42,42,42,42	0
56	MG	1A	3889	1/1	0.87	0.12	42,42,42,42	0
56	MG	1a	1819	1/1	0.87	0.15	49,49,49,49	0
56	MG	1A	3201	1/1	0.87	0.19	61,61,61,61	0
56	MG	2B	3006	1/1	0.87	0.17	72,72,72,72	0
56	MG	2A	3132	1/1	0.87	0.21	55,55,55,55	0
56	MG	2A	3678	1/1	0.87	0.11	59,59,59,59	0
56	MG	2a	1740	1/1	0.87	0.13	71,71,71,71	0
56	MG	2B	3013	1/1	0.87	0.25	68,68,68,68	0
56	MG	2A	3424	1/1	0.87	0.10	56,56,56,56	0
56	MG	1A	3357	1/1	0.87	0.09	46,46,46,46	0
56	MG	1a	1640	1/1	0.87	0.16	62,62,62,62	0
56	MG	2a	1757	1/1	0.87	0.09	66,66,66,66	0
56	MG	1A	3142	1/1	0.87	0.13	47,47,47,47	0
56	MG	2A	3288	1/1	0.87	0.39	63,63,63,63	0
56	MG	2A	3438	1/1	0.87	0.24	53,53,53,53	0
56	MG	1A	3293	1/1	0.87	0.15	54,54,54,54	0
56	MG	2A	3694	1/1	0.87	0.12	64,64,64,64	0
56	MG	2A	3695	1/1	0.87	0.16	45,45,45,45	0
56	MG	2W	201	1/1	0.87	0.11	54,54,54,54	0
56	MG	1A	4078	1/1	0.87	0.22	66,66,66,66	0
56	MG	1a	1656	1/1	0.87	0.22	64,64,64,64	0
56	MG	2a	1792	1/1	0.87	0.13	74,74,74,74	0
56	MG	2A	3165	1/1	0.87	0.16	52,52,52,52	0
56	MG	1A	3758	1/1	0.87	0.12	67,67,67,67	0
56	MG	1A	3491	1/1	0.87	0.16	60,60,60,60	0
56	MG	2a	1802	1/1	0.87	0.19	60,60,60,60	0
56	MG	1A	4125	1/1	0.87	0.14	38,38,38,38	0
56	MG	2A	3716	1/1	0.87	0.14	48,48,48,48	0
56	MG	1A	3646	1/1	0.87	0.13	46,46,46,46	0
56	MG	2a	1616	1/1	0.87	0.15	54,54,54,54	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2a	1621	1/1	0.87	0.11	67,67,67,67	0
56	MG	1x	106	1/1	0.87	0.20	68,68,68,68	0
56	MG	2a	1819	1/1	0.87	0.24	61,61,61,61	0
56	MG	2A	3734	1/1	0.87	0.13	70,70,70,70	0
56	MG	1A	3942	1/1	0.87	0.16	58,58,58,58	0
56	MG	2A	3322	1/1	0.87	0.19	60,60,60,60	0
56	MG	2A	3738	1/1	0.87	0.12	59,59,59,59	0
56	MG	2a	1632	1/1	0.87	0.22	60,60,60,60	0
56	MG	2A	3745	1/1	0.87	0.14	41,41,41,41	0
56	MG	1B	212	1/1	0.87	0.10	50,50,50,50	0
56	MG	1B	221	1/1	0.87	0.10	55,55,55,55	0
56	MG	1A	3186	1/1	0.87	0.11	45,45,45,45	0
56	MG	1A	3795	1/1	0.87	0.14	39,39,39,39	0
56	MG	2A	3006	1/1	0.87	0.30	58,58,58,58	0
56	MG	2a	1641	1/1	0.87	0.20	71,71,71,71	0
56	MG	2a	1642	1/1	0.87	0.30	63,63,63,63	0
56	MG	1A	3501	1/1	0.87	0.11	60,60,60,60	0
56	MG	1A	3983	1/1	0.87	0.14	68,68,68,68	0
56	MG	2A	3208	1/1	0.87	0.12	59,59,59,59	0
56	MG	1a	1721	1/1	0.87	0.08	68,68,68,68	0
56	MG	2A	3044	1/1	0.87	0.15	56,56,56,56	0
56	MG	1A	3399	1/1	0.87	0.14	51,51,51,51	0
56	MG	1A	3827	1/1	0.87	0.12	47,47,47,47	0
56	MG	2A	3791	1/1	0.87	0.10	64,64,64,64	0
56	MG	2a	1659	1/1	0.87	0.29	75,75,75,75	0
56	MG	2a	1660	1/1	0.87	0.24	82,82,82,82	0
56	MG	1R	202	1/1	0.87	0.16	38,38,38,38	0
56	MG	1A	3681	1/1	0.87	0.16	57,57,57,57	0
56	MG	2A	3583	1/1	0.87	0.22	68,68,68,68	0
56	MG	1A	3685	1/1	0.87	0.10	38,38,38,38	0
56	MG	2A	3230	1/1	0.87	0.23	73,73,73,73	0
56	MG	2A	3604	1/1	0.87	0.21	55,55,55,55	0
56	MG	2A	3815	1/1	0.87	0.12	67,67,67,67	0
56	MG	1A	3993	1/1	0.87	0.15	42,42,42,42	0
56	MG	2a	1684	1/1	0.87	0.17	63,63,63,63	0
56	MG	2A	3363	1/1	0.87	0.07	65,65,65,65	0
56	MG	2A	3712	1/1	0.88	0.11	60,60,60,60	0
56	MG	2a	1651	1/1	0.88	0.23	60,60,60,60	0
56	MG	1A	3695	1/1	0.88	0.15	58,58,58,58	0
56	MG	1A	3900	1/1	0.88	0.15	41,41,41,41	0
56	MG	2A	3718	1/1	0.88	0.09	57,57,57,57	0
56	MG	1a	1685	1/1	0.88	0.30	59,59,59,59	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3722	1/1	0.88	0.16	62,62,62,62	0
56	MG	1A	3347	1/1	0.88	0.13	55,55,55,55	0
56	MG	2A	3432	1/1	0.88	0.10	62,62,62,62	0
56	MG	2A	3256	1/1	0.88	0.25	63,63,63,63	0
56	MG	2A	3059	1/1	0.88	0.24	61,61,61,61	0
56	MG	1a	1693	1/1	0.88	0.23	65,65,65,65	0
56	MG	2A	3260	1/1	0.88	0.18	69,69,69,69	0
56	MG	1a	1700	1/1	0.88	0.11	56,56,56,56	0
56	MG	2A	3449	1/1	0.88	0.34	55,55,55,55	0
56	MG	1A	3482	1/1	0.88	0.11	45,45,45,45	0
56	MG	2A	3266	1/1	0.88	0.11	67,67,67,67	0
56	MG	2a	1683	1/1	0.88	0.24	66,66,66,66	0
56	MG	2A	3455	1/1	0.88	0.33	68,68,68,68	0
56	MG	2a	1690	1/1	0.88	0.19	61,61,61,61	0
56	MG	2A	3458	1/1	0.88	0.27	60,60,60,60	0
56	MG	2A	3762	1/1	0.88	0.09	58,58,58,58	0
56	MG	2A	3763	1/1	0.88	0.25	66,66,66,66	0
56	MG	2A	3461	1/1	0.88	0.14	52,52,52,52	0
56	MG	2A	3465	1/1	0.88	0.16	63,63,63,63	0
56	MG	1B	227	1/1	0.88	0.20	54,54,54,54	0
56	MG	2A	3276	1/1	0.88	0.16	62,62,62,62	0
56	MG	2a	1709	1/1	0.88	0.27	69,69,69,69	0
56	MG	2A	3481	1/1	0.88	0.14	48,48,48,48	0
56	MG	1A	3159	1/1	0.88	0.27	56,56,56,56	0
56	MG	1A	3571	1/1	0.88	0.11	35,35,35,35	0
56	MG	2A	3284	1/1	0.88	0.16	60,60,60,60	0
56	MG	1a	1730	1/1	0.88	0.13	51,51,51,51	0
56	MG	1A	3227	1/1	0.88	0.10	49,49,49,49	0
56	MG	1A	3750	1/1	0.88	0.13	41,41,41,41	0
56	MG	1A	3421	1/1	0.88	0.42	62,62,62,62	0
56	MG	2A	3810	1/1	0.88	0.12	63,63,63,63	0
56	MG	2A	3301	1/1	0.88	0.08	68,68,68,68	0
56	MG	1A	3591	1/1	0.88	0.12	29,29,29,29	0
56	MG	1A	3762	1/1	0.88	0.12	41,41,41,41	0
56	MG	1A	3495	1/1	0.88	0.11	60,60,60,60	0
56	MG	1A	3619	1/1	0.88	0.09	35,35,35,35	0
56	MG	2A	3114	1/1	0.88	0.19	49,49,49,49	0
56	MG	2A	3541	1/1	0.88	0.10	40,40,40,40	0
56	MG	2a	1748	1/1	0.88	0.10	74,74,74,74	0
56	MG	1a	1767	1/1	0.88	0.10	71,71,71,71	0
56	MG	1A	3270	1/1	0.88	0.13	53,53,53,53	0
56	MG	2A	3133	1/1	0.88	0.16	69,69,69,69	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	10	105	1/1	0.88	0.23	55,55,55,55	0
56	MG	2A	3872	1/1	0.88	0.12	47,47,47,47	0
56	MG	1A	3790	1/1	0.88	0.12	45,45,45,45	0
56	MG	2A	3574	1/1	0.88	0.13	57,57,57,57	0
56	MG	2A	3577	1/1	0.88	0.14	53,53,53,53	0
56	MG	17	101	1/1	0.88	0.09	52,52,52,52	0
56	MG	2A	3580	1/1	0.88	0.15	67,67,67,67	0
56	MG	1A	3624	1/1	0.88	0.14	63,63,63,63	0
56	MG	2A	3588	1/1	0.88	0.20	71,71,71,71	0
56	MG	1A	3798	1/1	0.88	0.15	56,56,56,56	0
56	MG	2A	3591	1/1	0.88	0.09	63,63,63,63	0
56	MG	1a	1605	1/1	0.88	0.11	57,57,57,57	0
56	MG	1A	3437	1/1	0.88	0.11	54,54,54,54	0
56	MG	2a	1801	1/1	0.88	0.21	72,72,72,72	0
56	MG	1A	3249	1/1	0.88	0.13	70,70,70,70	0
56	MG	2a	1803	1/1	0.88	0.24	65,65,65,65	0
56	MG	1A	4015	1/1	0.88	0.13	70,70,70,70	0
56	MG	2a	1808	1/1	0.88	0.33	75,75,75,75	0
56	MG	2A	3614	1/1	0.88	0.11	32,32,32,32	0
56	MG	1A	3805	1/1	0.88	0.22	34,34,34,34	0
56	MG	1a	1633	1/1	0.88	0.18	56,56,56,56	0
56	MG	1t	3001	1/1	0.88	0.26	53,53,53,53	0
56	MG	1A	3204	1/1	0.88	0.12	68,68,68,68	0
56	MG	1A	3387	1/1	0.88	0.19	37,37,37,37	0
56	MG	1w	105	1/1	0.88	0.14	67,67,67,67	0
56	MG	2A	3643	1/1	0.88	0.14	47,47,47,47	0
56	MG	1A	3648	1/1	0.88	0.14	25,25,25,25	0
56	MG	25	105	1/1	0.88	0.13	59,59,59,59	0
56	MG	2a	1603	1/1	0.88	0.12	71,71,71,71	0
56	MG	2a	1604	1/1	0.88	0.14	67,67,67,67	0
56	MG	2A	3645	1/1	0.88	0.13	56,56,56,56	0
56	MG	1a	1646	1/1	0.88	0.26	78,78,78,78	0
56	MG	1w	111	1/1	0.88	0.12	61,61,61,61	0
56	MG	2A	3663	1/1	0.88	0.11	84,84,84,84	0
56	MG	1A	3661	1/1	0.88	0.10	39,39,39,39	0
56	MG	2A	3205	1/1	0.88	0.12	53,53,53,53	0
56	MG	2a	1614	1/1	0.88	0.20	59,59,59,59	0
56	MG	1A	3842	1/1	0.88	0.12	53,53,53,53	0
56	MG	2a	1844	1/1	0.88	0.16	66,66,66,66	0
56	MG	2A	3369	1/1	0.88	0.10	68,68,68,68	0
56	MG	2A	3373	1/1	0.88	0.13	57,57,57,57	0
56	MG	2A	3377	1/1	0.88	0.10	57,57,57,57	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3380	1/1	0.88	0.11	68,68,68,68	0
56	MG	1A	4069	1/1	0.88	0.13	55,55,55,55	0
56	MG	2A	3682	1/1	0.88	0.12	71,71,71,71	0
56	MG	2v	102	1/1	0.88	0.21	67,67,67,67	0
56	MG	1A	3057	1/1	0.88	0.08	34,34,34,34	0
56	MG	1x	108	1/1	0.88	0.27	59,59,59,59	0
56	MG	1x	109	1/1	0.88	0.17	68,68,68,68	0
56	MG	1A	3394	1/1	0.88	0.10	68,68,68,68	0
56	MG	2A	3396	1/1	0.88	0.25	60,60,60,60	0
56	MG	1A	3859	1/1	0.88	0.14	29,29,29,29	0
56	MG	1A	3024	1/1	0.88	0.12	47,47,47,47	0
56	MG	1A	3461	1/1	0.88	0.14	50,50,50,50	0
56	MG	1a	1675	1/1	0.88	0.23	70,70,70,70	0
56	MG	2A	3008	1/1	0.88	0.18	56,56,56,56	0
56	MG	2a	1644	1/1	0.88	0.13	64,64,64,64	0
56	MG	1A	3547	1/1	0.88	0.10	56,56,56,56	0
56	MG	1A	3402	1/1	0.88	0.18	53,53,53,53	0
56	MG	2y	3007	1/1	0.88	0.15	75,75,75,75	0
56	MG	2a	1666	1/1	0.89	0.08	64,64,64,64	0
56	MG	2A	3110	1/1	0.89	0.26	65,65,65,65	0
56	MG	1A	3052	1/1	0.89	0.13	48,48,48,48	0
56	MG	1A	4037	1/1	0.89	0.14	48,48,48,48	0
56	MG	2A	3116	1/1	0.89	0.32	56,56,56,56	0
56	MG	2A	3772	1/1	0.89	0.11	55,55,55,55	0
56	MG	2A	3310	1/1	0.89	0.09	58,58,58,58	0
56	MG	2A	3119	1/1	0.89	0.23	62,62,62,62	0
56	MG	2A	3778	1/1	0.89	0.16	71,71,71,71	0
56	MG	2a	1682	1/1	0.89	0.43	62,62,62,62	0
56	MG	1A	4043	1/1	0.89	0.27	54,54,54,54	0
56	MG	1a	1821	1/1	0.89	0.14	51,51,51,51	0
56	MG	1A	3471	1/1	0.89	0.17	51,51,51,51	0
56	MG	1l	203	1/1	0.89	0.12	52,52,52,52	0
56	MG	2A	3546	1/1	0.89	0.10	42,42,42,42	0
56	MG	2A	3549	1/1	0.89	0.11	56,56,56,56	0
56	MG	1A	3728	1/1	0.89	0.17	62,62,62,62	0
56	MG	1A	4067	1/1	0.89	0.22	58,58,58,58	0
56	MG	1A	3873	1/1	0.89	0.08	25,25,25,25	0
56	MG	2A	3814	1/1	0.89	0.16	71,71,71,71	0
56	MG	2A	3334	1/1	0.89	0.12	61,61,61,61	0
56	MG	1A	3886	1/1	0.89	0.14	38,38,38,38	0
56	MG	1A	3473	1/1	0.89	0.18	62,62,62,62	0
56	MG	1A	3748	1/1	0.89	0.11	24,24,24,24	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3238	1/1	0.89	0.10	41,41,41,41	0
56	MG	2A	3833	1/1	0.89	0.13	47,47,47,47	0
56	MG	2A	3581	1/1	0.89	0.10	30,30,30,30	0
56	MG	1w	103	1/1	0.89	0.15	51,51,51,51	0
56	MG	1A	3630	1/1	0.89	0.10	56,56,56,56	0
56	MG	1A	3241	1/1	0.89	0.18	54,54,54,54	0
56	MG	2A	3870	1/1	0.89	0.09	53,53,53,53	0
56	MG	2a	1730	1/1	0.89	0.14	74,74,74,74	0
56	MG	1A	3538	1/1	0.89	0.26	49,49,49,49	0
56	MG	2A	3592	1/1	0.89	0.24	56,56,56,56	0
56	MG	1A	3543	1/1	0.89	0.31	62,62,62,62	0
56	MG	1A	3301	1/1	0.89	0.17	40,40,40,40	0
56	MG	2A	3358	1/1	0.89	0.17	57,57,57,57	0
56	MG	1A	3489	1/1	0.89	0.31	61,61,61,61	0
56	MG	1A	3932	1/1	0.89	0.13	60,60,60,60	0
56	MG	1A	3937	1/1	0.89	0.16	73,73,73,73	0
56	MG	2A	3892	1/1	0.89	0.11	64,64,64,64	0
56	MG	1A	3549	1/1	0.89	0.21	54,54,54,54	0
56	MG	2B	3002	1/1	0.89	0.10	63,63,63,63	0
56	MG	2A	3620	1/1	0.89	0.19	67,67,67,67	0
56	MG	2B	3007	1/1	0.89	0.12	63,63,63,63	0
56	MG	1A	3951	1/1	0.89	0.10	50,50,50,50	0
56	MG	1O	202	1/1	0.89	0.17	58,58,58,58	0
56	MG	1O	204	1/1	0.89	0.18	62,62,62,62	0
56	MG	2A	3375	1/1	0.89	0.10	61,61,61,61	0
56	MG	1A	3791	1/1	0.89	0.12	47,47,47,47	0
56	MG	1A	3452	1/1	0.89	0.13	49,49,49,49	0
56	MG	1a	1712	1/1	0.89	0.32	56,56,56,56	0
56	MG	2A	3388	1/1	0.89	0.13	58,58,58,58	0
56	MG	2E	305	1/1	0.89	0.09	53,53,53,53	0
56	MG	2A	3389	1/1	0.89	0.42	51,51,51,51	0
56	MG	1A	3366	1/1	0.89	0.31	50,50,50,50	0
56	MG	1A	3562	1/1	0.89	0.19	51,51,51,51	0
56	MG	2U	204	1/1	0.89	0.17	62,62,62,62	0
56	MG	1a	1728	1/1	0.89	0.09	53,53,53,53	0
56	MG	2A	3222	1/1	0.89	0.08	48,48,48,48	0
56	MG	2A	3030	1/1	0.89	0.12	54,54,54,54	0
56	MG	1A	3414	1/1	0.89	0.10	45,45,45,45	0
56	MG	26	502	1/1	0.89	0.37	69,69,69,69	0
56	MG	2A	3042	1/1	0.89	0.15	49,49,49,49	0
56	MG	2A	3043	1/1	0.89	0.20	53,53,53,53	0
56	MG	2A	3414	1/1	0.89	0.21	52,52,52,52	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3683	1/1	0.89	0.09	21,21,21,21	0
56	MG	2A	3046	1/1	0.89	0.20	65,65,65,65	0
56	MG	2A	3684	1/1	0.89	0.09	51,51,51,51	0
56	MG	1A	3817	1/1	0.89	0.08	23,23,23,23	0
56	MG	2a	1825	1/1	0.89	0.11	74,74,74,74	0
56	MG	1A	3820	1/1	0.89	0.15	47,47,47,47	0
56	MG	2A	3689	1/1	0.89	0.17	62,62,62,62	0
56	MG	1A	3062	1/1	0.89	0.28	55,55,55,55	0
56	MG	2A	3252	1/1	0.89	0.38	50,50,50,50	0
56	MG	2a	1620	1/1	0.89	0.19	63,63,63,63	0
56	MG	2A	3253	1/1	0.89	0.13	60,60,60,60	0
56	MG	2a	1835	1/1	0.89	0.13	72,72,72,72	0
56	MG	1A	3345	1/1	0.89	0.11	39,39,39,39	0
56	MG	2A	3430	1/1	0.89	0.17	64,64,64,64	0
56	MG	13	101	1/1	0.89	0.12	55,55,55,55	0
56	MG	2A	3065	1/1	0.89	0.25	65,65,65,65	0
56	MG	2A	3067	1/1	0.89	0.19	58,58,58,58	0
56	MG	2A	3068	1/1	0.89	0.09	59,59,59,59	0
56	MG	2A	3441	1/1	0.89	0.21	39,39,39,39	0
56	MG	1A	3692	1/1	0.89	0.13	56,56,56,56	0
56	MG	2A	3079	1/1	0.89	0.12	51,51,51,51	0
56	MG	1A	4014	1/1	0.89	0.17	54,54,54,54	0
56	MG	2A	3089	1/1	0.89	0.14	50,50,50,50	0
56	MG	2v	101	1/1	0.89	0.23	62,62,62,62	0
56	MG	2A	3720	1/1	0.89	0.08	52,52,52,52	0
56	MG	1A	3694	1/1	0.89	0.10	55,55,55,55	0
56	MG	1a	1771	1/1	0.89	0.10	73,73,73,73	0
56	MG	2A	3094	1/1	0.89	0.13	47,47,47,47	0
56	MG	1A	3049	1/1	0.89	0.14	36,36,36,36	0
56	MG	1A	3508	1/1	0.89	0.13	56,56,56,56	0
56	MG	2A	3466	1/1	0.89	0.15	54,54,54,54	0
56	MG	1A	3592	1/1	0.89	0.15	51,51,51,51	0
56	MG	1a	1785	1/1	0.89	0.12	69,69,69,69	0
56	MG	1a	1794	1/1	0.89	0.14	79,79,79,79	0
56	MG	2A	3293	1/1	0.89	0.08	69,69,69,69	0
56	MG	2A	3299	1/1	0.89	0.06	65,65,65,65	0
56	MG	2A	3105	1/1	0.89	0.15	69,69,69,69	0
56	MG	2y	3004	1/1	0.89	0.13	79,79,79,79	0
56	MG	1a	1621	1/1	0.89	0.26	56,56,56,56	0
56	MG	1a	1804	1/1	0.89	0.11	62,62,62,62	0
56	MG	1A	3510	1/1	0.90	0.08	48,48,48,48	0
56	MG	1A	3988	1/1	0.90	0.08	41,41,41,41	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3655	1/1	0.90	0.09	67,67,67,67	0
56	MG	1A	3001	1/1	0.90	0.12	38,38,38,38	0
56	MG	1A	3835	1/1	0.90	0.10	63,63,63,63	0
56	MG	1a	1809	1/1	0.90	0.10	47,47,47,47	0
56	MG	2A	3151	1/1	0.90	0.15	59,59,59,59	0
56	MG	2A	3666	1/1	0.90	0.09	34,34,34,34	0
56	MG	2A	3361	1/1	0.90	0.20	60,60,60,60	0
56	MG	1A	3837	1/1	0.90	0.08	25,25,25,25	0
56	MG	1A	3841	1/1	0.90	0.25	36,36,36,36	0
56	MG	2A	3365	1/1	0.90	0.12	71,71,71,71	0
56	MG	1a	1815	1/1	0.90	0.17	60,60,60,60	0
56	MG	1A	3520	1/1	0.90	0.12	47,47,47,47	0
56	MG	2A	3371	1/1	0.90	0.08	61,61,61,61	0
56	MG	2A	3372	1/1	0.90	0.17	50,50,50,50	0
56	MG	1a	1608	1/1	0.90	0.14	56,56,56,56	0
56	MG	2A	3168	1/1	0.90	0.26	53,53,53,53	0
56	MG	1a	1825	1/1	0.90	0.34	56,56,56,56	0
56	MG	2a	1643	1/1	0.90	0.19	63,63,63,63	0
56	MG	1b	3002	1/1	0.90	0.21	68,68,68,68	0
56	MG	1a	1611	1/1	0.90	0.14	55,55,55,55	0
56	MG	2a	1646	1/1	0.90	0.19	53,53,53,53	0
56	MG	2A	3385	1/1	0.90	0.26	59,59,59,59	0
56	MG	1A	3139	1/1	0.90	0.11	43,43,43,43	0
56	MG	1A	4018	1/1	0.90	0.09	70,70,70,70	0
56	MG	2a	1652	1/1	0.90	0.22	71,71,71,71	0
56	MG	1a	1619	1/1	0.90	0.27	51,51,51,51	0
56	MG	2A	3700	1/1	0.90	0.12	59,59,59,59	0
56	MG	2A	3187	1/1	0.90	0.16	63,63,63,63	0
56	MG	1A	4022	1/1	0.90	0.08	55,55,55,55	0
56	MG	1a	1624	1/1	0.90	0.23	59,59,59,59	0
56	MG	2A	3706	1/1	0.90	0.15	68,68,68,68	0
56	MG	1A	3851	1/1	0.90	0.12	28,28,28,28	0
56	MG	1a	1628	1/1	0.90	0.17	54,54,54,54	0
56	MG	2A	3404	1/1	0.90	0.30	52,52,52,52	0
56	MG	1w	101	1/1	0.90	0.11	59,59,59,59	0
56	MG	1A	3727	1/1	0.90	0.09	14,14,14,14	0
56	MG	1a	1631	1/1	0.90	0.24	59,59,59,59	0
56	MG	1A	3469	1/1	0.90	0.12	68,68,68,68	0
56	MG	1a	1634	1/1	0.90	0.31	72,72,72,72	0
56	MG	1w	107	1/1	0.90	0.23	70,70,70,70	0
56	MG	1w	109	1/1	0.90	0.10	76,76,76,76	0
56	MG	1A	3260	1/1	0.90	0.20	47,47,47,47	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3741	1/1	0.90	0.09	17,17,17,17	0
56	MG	1A	4042	1/1	0.90	0.09	24,24,24,24	0
56	MG	2A	3743	1/1	0.90	0.21	58,58,58,58	0
56	MG	2a	1694	1/1	0.90	0.34	67,67,67,67	0
56	MG	2A	3218	1/1	0.90	0.13	63,63,63,63	0
56	MG	1A	3240	1/1	0.90	0.18	54,54,54,54	0
56	MG	2a	1702	1/1	0.90	0.22	65,65,65,65	0
56	MG	1a	1650	1/1	0.90	0.11	46,46,46,46	0
56	MG	1A	4051	1/1	0.90	0.13	60,60,60,60	0
56	MG	2A	3754	1/1	0.90	0.14	48,48,48,48	0
56	MG	1A	4056	1/1	0.90	0.14	54,54,54,54	0
56	MG	2A	3437	1/1	0.90	0.25	51,51,51,51	0
56	MG	1A	4057	1/1	0.90	0.09	59,59,59,59	0
56	MG	1x	111	1/1	0.90	0.13	52,52,52,52	0
56	MG	1A	3064	1/1	0.90	0.12	49,49,49,49	0
56	MG	2A	3764	1/1	0.90	0.10	66,66,66,66	0
56	MG	1x	115	1/1	0.90	0.09	68,68,68,68	0
56	MG	1y	3001	1/1	0.90	0.09	62,62,62,62	0
56	MG	2A	3248	1/1	0.90	0.13	55,55,55,55	0
56	MG	1A	3310	1/1	0.90	0.31	52,52,52,52	0
56	MG	1A	3639	1/1	0.90	0.12	51,51,51,51	0
56	MG	2a	1724	1/1	0.90	0.32	53,53,53,53	0
56	MG	1A	3894	1/1	0.90	0.22	37,37,37,37	0
56	MG	2A	3459	1/1	0.90	0.20	60,60,60,60	0
56	MG	2A	3780	1/1	0.90	0.09	53,53,53,53	0
56	MG	2A	3001	1/1	0.90	0.27	56,56,56,56	0
56	MG	1A	3433	1/1	0.90	0.13	53,53,53,53	0
56	MG	1a	1672	1/1	0.90	0.23	69,69,69,69	0
56	MG	2A	3469	1/1	0.90	0.37	54,54,54,54	0
56	MG	1A	3763	1/1	0.90	0.09	48,48,48,48	0
56	MG	2a	1744	1/1	0.90	0.10	81,81,81,81	0
56	MG	2A	3798	1/1	0.90	0.09	75,75,75,75	0
56	MG	2A	3801	1/1	0.90	0.10	65,65,65,65	0
56	MG	2A	3473	1/1	0.90	0.12	59,59,59,59	0
56	MG	2a	1754	1/1	0.90	0.11	82,82,82,82	0
56	MG	1A	3901	1/1	0.90	0.20	48,48,48,48	0
56	MG	1A	3384	1/1	0.90	0.11	48,48,48,48	0
56	MG	1A	3323	1/1	0.90	0.11	43,43,43,43	0
56	MG	1A	3442	1/1	0.90	0.25	48,48,48,48	0
56	MG	2A	3268	1/1	0.90	0.13	60,60,60,60	0
56	MG	1a	1684	1/1	0.90	0.27	49,49,49,49	0
56	MG	2A	3274	1/1	0.90	0.14	60,60,60,60	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2a	1770	1/1	0.90	0.10	77,77,77,77	0
56	MG	2a	1771	1/1	0.90	0.14	68,68,68,68	0
56	MG	1A	3920	1/1	0.90	0.12	65,65,65,65	0
56	MG	1a	1690	1/1	0.90	0.17	56,56,56,56	0
56	MG	2A	3049	1/1	0.90	0.16	57,57,57,57	0
56	MG	1A	3390	1/1	0.90	0.28	43,43,43,43	0
56	MG	2A	3840	1/1	0.90	0.10	45,45,45,45	0
56	MG	2A	3850	1/1	0.90	0.10	77,77,77,77	0
56	MG	1B	211	1/1	0.90	0.10	51,51,51,51	0
56	MG	2A	3058	1/1	0.90	0.21	51,51,51,51	0
56	MG	2a	1796	1/1	0.90	0.28	69,69,69,69	0
56	MG	2a	1800	1/1	0.90	0.15	60,60,60,60	0
56	MG	1a	1698	1/1	0.90	0.17	68,68,68,68	0
56	MG	1a	1699	1/1	0.90	0.15	74,74,74,74	0
56	MG	1A	3561	1/1	0.90	0.14	41,41,41,41	0
56	MG	2A	3543	1/1	0.90	0.10	37,37,37,37	0
56	MG	1A	3328	1/1	0.90	0.14	48,48,48,48	0
56	MG	2A	3066	1/1	0.90	0.21	53,53,53,53	0
56	MG	1a	1707	1/1	0.90	0.32	62,62,62,62	0
56	MG	1A	3678	1/1	0.90	0.08	14,14,14,14	0
56	MG	2A	3071	1/1	0.90	0.14	56,56,56,56	0
56	MG	2A	3567	1/1	0.90	0.10	52,52,52,52	0
56	MG	1A	3330	1/1	0.90	0.29	67,67,67,67	0
56	MG	1A	3947	1/1	0.90	0.08	34,34,34,34	0
56	MG	2A	3307	1/1	0.90	0.41	65,65,65,65	0
56	MG	1a	1723	1/1	0.90	0.11	41,41,41,41	0
56	MG	2A	3309	1/1	0.90	0.16	72,72,72,72	0
56	MG	1F	301	1/1	0.90	0.25	62,62,62,62	0
56	MG	1F	308	1/1	0.90	0.15	54,54,54,54	0
56	MG	2a	1827	1/1	0.90	0.14	71,71,71,71	0
56	MG	1G	3003	1/1	0.90	0.14	64,64,64,64	0
56	MG	2A	3585	1/1	0.90	0.17	62,62,62,62	0
56	MG	1A	3396	1/1	0.90	0.12	36,36,36,36	0
56	MG	2B	3017	1/1	0.90	0.10	50,50,50,50	0
56	MG	1A	3954	1/1	0.90	0.19	36,36,36,36	0
56	MG	1a	1749	1/1	0.90	0.24	57,57,57,57	0
56	MG	2D	304	1/1	0.90	0.28	53,53,53,53	0
56	MG	2a	1839	1/1	0.90	0.20	69,69,69,69	0
56	MG	2E	302	1/1	0.90	0.12	55,55,55,55	0
56	MG	1A	3580	1/1	0.90	0.34	33,33,33,33	0
56	MG	2A	3595	1/1	0.90	0.09	58,58,58,58	0
56	MG	2F	303	1/1	0.90	0.12	46,46,46,46	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3601	1/1	0.90	0.09	32,32,32,32	0
56	MG	2O	8001	1/1	0.90	0.09	48,48,48,48	0
56	MG	2g	8001	1/1	0.90	0.09	69,69,69,69	0
56	MG	1A	3958	1/1	0.90	0.20	67,67,67,67	0
56	MG	2T	3002	1/1	0.90	0.12	53,53,53,53	0
56	MG	2A	3328	1/1	0.90	0.16	55,55,55,55	0
56	MG	2q	3003	1/1	0.90	0.16	68,68,68,68	0
56	MG	2A	3330	1/1	0.90	0.07	53,53,53,53	0
56	MG	2A	3331	1/1	0.90	0.15	68,68,68,68	0
56	MG	1a	1757	1/1	0.90	0.13	61,61,61,61	0
56	MG	1A	3581	1/1	0.90	0.09	42,42,42,42	0
56	MG	1A	3158	1/1	0.90	0.19	47,47,47,47	0
56	MG	1A	3971	1/1	0.90	0.12	68,68,68,68	0
56	MG	2a	1601	1/1	0.90	0.18	61,61,61,61	0
56	MG	1A	3979	1/1	0.90	0.09	21,21,21,21	0
56	MG	1A	3826	1/1	0.90	0.09	60,60,60,60	0
56	MG	2A	3344	1/1	0.90	0.34	55,55,55,55	0
56	MG	2x	103	1/1	0.90	0.18	77,77,77,77	0
56	MG	2A	3630	1/1	0.90	0.11	61,61,61,61	0
56	MG	2a	1607	1/1	0.90	0.28	73,73,73,73	0
56	MG	2A	3636	1/1	0.90	0.20	54,54,54,54	0
56	MG	1A	3584	1/1	0.90	0.08	22,22,22,22	0
56	MG	2a	1610	1/1	0.90	0.26	74,74,74,74	0
56	MG	2a	1611	1/1	0.90	0.15	75,75,75,75	0
56	MG	1Y	204	1/1	0.90	0.27	48,48,48,48	0
56	MG	1A	3254	1/1	0.90	0.14	52,52,52,52	0
56	MG	2a	1648	1/1	0.91	0.07	76,76,76,76	0
56	MG	2A	3749	1/1	0.91	0.08	52,52,52,52	0
56	MG	1m	201	1/1	0.91	0.19	40,40,40,40	0
56	MG	1a	1654	1/1	0.91	0.20	61,61,61,61	0
56	MG	2A	3314	1/1	0.91	0.11	60,60,60,60	0
56	MG	1A	3311	1/1	0.91	0.18	53,53,53,53	0
56	MG	2A	3319	1/1	0.91	0.14	52,52,52,52	0
56	MG	2A	3320	1/1	0.91	0.09	57,57,57,57	0
56	MG	2A	3504	1/1	0.91	0.11	51,51,51,51	0
56	MG	2A	3760	1/1	0.91	0.14	60,60,60,60	0
56	MG	1B	218	1/1	0.91	0.21	58,58,58,58	0
56	MG	1A	3745	1/1	0.91	0.10	34,34,34,34	0
56	MG	2a	1663	1/1	0.91	0.20	66,66,66,66	0
56	MG	1a	1663	1/1	0.91	0.16	64,64,64,64	0
56	MG	2A	3513	1/1	0.91	0.13	43,43,43,43	0
56	MG	1A	3747	1/1	0.91	0.11	38,38,38,38	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3566	1/1	0.91	0.11	38,38,38,38	0
56	MG	2A	3521	1/1	0.91	0.14	52,52,52,52	0
56	MG	1B	233	1/1	0.91	0.13	64,64,64,64	0
56	MG	2A	3532	1/1	0.91	0.21	61,61,61,61	0
56	MG	2A	3535	1/1	0.91	0.17	53,53,53,53	0
56	MG	1B	236	1/1	0.91	0.07	23,23,23,23	0
56	MG	2a	1681	1/1	0.91	0.17	57,57,57,57	0
56	MG	1a	1674	1/1	0.91	0.12	56,56,56,56	0
56	MG	2A	3332	1/1	0.91	0.13	52,52,52,52	0
56	MG	1D	305	1/1	0.91	0.18	52,52,52,52	0
56	MG	2a	1687	1/1	0.91	0.19	65,65,65,65	0
56	MG	2A	3794	1/1	0.91	0.09	61,61,61,61	0
56	MG	1A	3149	1/1	0.91	0.24	44,44,44,44	0
56	MG	2A	3169	1/1	0.91	0.14	52,52,52,52	0
56	MG	2a	1699	1/1	0.91	0.19	62,62,62,62	0
56	MG	1A	3087	1/1	0.91	0.14	43,43,43,43	0
56	MG	1A	3237	1/1	0.91	0.23	51,51,51,51	0
56	MG	1A	3429	1/1	0.91	0.08	45,45,45,45	0
56	MG	2A	3806	1/1	0.91	0.10	55,55,55,55	0
56	MG	1A	3674	1/1	0.91	0.08	22,22,22,22	0
56	MG	2A	3177	1/1	0.91	0.23	55,55,55,55	0
56	MG	1A	3875	1/1	0.91	0.09	25,25,25,25	0
56	MG	1a	1688	1/1	0.91	0.19	51,51,51,51	0
56	MG	2a	1710	1/1	0.91	0.16	70,70,70,70	0
56	MG	1A	3877	1/1	0.91	0.09	67,67,67,67	0
56	MG	1A	3878	1/1	0.91	0.10	39,39,39,39	0
56	MG	2A	3191	1/1	0.91	0.12	71,71,71,71	0
56	MG	1A	4012	1/1	0.91	0.07	71,71,71,71	0
56	MG	2A	3195	1/1	0.91	0.26	65,65,65,65	0
56	MG	1a	1694	1/1	0.91	0.24	58,58,58,58	0
56	MG	1A	3296	1/1	0.91	0.40	66,66,66,66	0
56	MG	1x	114	1/1	0.91	0.19	65,65,65,65	0
56	MG	1A	3476	1/1	0.91	0.25	56,56,56,56	0
56	MG	2a	1725	1/1	0.91	0.15	66,66,66,66	0
56	MG	2A	3599	1/1	0.91	0.12	57,57,57,57	0
56	MG	2a	1728	1/1	0.91	0.15	73,73,73,73	0
56	MG	1A	4017	1/1	0.91	0.14	56,56,56,56	0
56	MG	2A	3861	1/1	0.91	0.09	65,65,65,65	0
56	MG	1A	3203	1/1	0.91	0.07	45,45,45,45	0
56	MG	2A	3867	1/1	0.91	0.08	75,75,75,75	0
56	MG	2A	3869	1/1	0.91	0.12	46,46,46,46	0
56	MG	1W	202	1/1	0.91	0.13	50,50,50,50	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3605	1/1	0.91	0.14	56,56,56,56	0
56	MG	2A	3370	1/1	0.91	0.24	57,57,57,57	0
56	MG	1a	1710	1/1	0.91	0.19	62,62,62,62	0
56	MG	2a	1747	1/1	0.91	0.09	80,80,80,80	0
56	MG	2A	3610	1/1	0.91	0.10	59,59,59,59	0
56	MG	1Y	201	1/1	0.91	0.14	43,43,43,43	0
56	MG	2A	3004	1/1	0.91	0.14	56,56,56,56	0
56	MG	1A	3438	1/1	0.91	0.14	41,41,41,41	0
56	MG	1A	3348	1/1	0.91	0.09	41,41,41,41	0
56	MG	2A	3016	1/1	0.91	0.14	66,66,66,66	0
56	MG	2A	3894	1/1	0.91	0.10	62,62,62,62	0
56	MG	2A	3898	1/1	0.91	0.24	52,52,52,52	0
56	MG	1A	3354	1/1	0.91	0.22	50,50,50,50	0
56	MG	1A	4027	1/1	0.91	0.11	49,49,49,49	0
56	MG	2A	3027	1/1	0.91	0.52	53,53,53,53	0
56	MG	1A	3088	1/1	0.91	0.13	50,50,50,50	0
56	MG	2a	1781	1/1	0.91	0.09	63,63,63,63	0
56	MG	2B	3008	1/1	0.91	0.17	56,56,56,56	0
56	MG	10	107	1/1	0.91	0.12	53,53,53,53	0
56	MG	2A	3037	1/1	0.91	0.09	42,42,42,42	0
56	MG	2a	1790	1/1	0.91	0.10	66,66,66,66	0
56	MG	1A	3902	1/1	0.91	0.16	54,54,54,54	0
56	MG	1a	1735	1/1	0.91	0.12	60,60,60,60	0
56	MG	2A	3647	1/1	0.91	0.17	51,51,51,51	0
56	MG	1a	1738	1/1	0.91	0.10	54,54,54,54	0
56	MG	2A	3653	1/1	0.91	0.22	66,66,66,66	0
56	MG	2A	3397	1/1	0.91	0.18	53,53,53,53	0
56	MG	2a	1798	1/1	0.91	0.23	53,53,53,53	0
56	MG	2a	1799	1/1	0.91	0.15	67,67,67,67	0
56	MG	2D	301	1/1	0.91	0.18	58,58,58,58	0
56	MG	16	103	1/1	0.91	0.08	53,53,53,53	0
56	MG	2E	301	1/1	0.91	0.12	49,49,49,49	0
56	MG	2A	3658	1/1	0.91	0.12	54,54,54,54	0
56	MG	2a	1806	1/1	0.91	0.13	67,67,67,67	0
56	MG	2A	3660	1/1	0.91	0.15	57,57,57,57	0
56	MG	2A	3047	1/1	0.91	0.09	54,54,54,54	0
56	MG	2F	301	1/1	0.91	0.14	48,48,48,48	0
56	MG	1A	3303	1/1	0.91	0.26	51,51,51,51	0
56	MG	2F	304	1/1	0.91	0.37	51,51,51,51	0
56	MG	2a	1813	1/1	0.91	0.15	62,62,62,62	0
56	MG	17	103	1/1	0.91	0.08	60,60,60,60	0
56	MG	1A	3906	1/1	0.91	0.14	68,68,68,68	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3696	1/1	0.91	0.06	16,16,16,16	0
56	MG	1A	3913	1/1	0.91	0.10	25,25,25,25	0
56	MG	2A	3674	1/1	0.91	0.16	50,50,50,50	0
56	MG	2a	1822	1/1	0.91	0.18	65,65,65,65	0
56	MG	2A	3418	1/1	0.91	0.20	60,60,60,60	0
56	MG	1A	3916	1/1	0.91	0.11	39,39,39,39	0
56	MG	1a	1760	1/1	0.91	0.17	61,61,61,61	0
56	MG	1a	1610	1/1	0.91	0.08	57,57,57,57	0
56	MG	2A	3263	1/1	0.91	0.17	66,66,66,66	0
56	MG	1A	3628	1/1	0.91	0.09	32,32,32,32	0
56	MG	28	101	1/1	0.91	0.21	52,52,52,52	0
56	MG	1A	3807	1/1	0.91	0.08	28,28,28,28	0
56	MG	2a	1831	1/1	0.91	0.09	78,78,78,78	0
56	MG	1A	4064	1/1	0.91	0.16	45,45,45,45	0
56	MG	2A	3431	1/1	0.91	0.12	55,55,55,55	0
56	MG	1A	3810	1/1	0.91	0.17	55,55,55,55	0
56	MG	2A	3690	1/1	0.91	0.19	66,66,66,66	0
56	MG	1A	3702	1/1	0.91	0.12	42,42,42,42	0
56	MG	2A	3275	1/1	0.91	0.09	56,56,56,56	0
56	MG	1A	3819	1/1	0.91	0.15	52,52,52,52	0
56	MG	1A	3629	1/1	0.91	0.07	32,32,32,32	0
56	MG	1a	1796	1/1	0.91	0.09	67,67,67,67	0
56	MG	1A	3708	1/1	0.91	0.07	51,51,51,51	0
56	MG	1a	1803	1/1	0.91	0.09	63,63,63,63	0
56	MG	2f	3001	1/1	0.91	0.11	47,47,47,47	0
56	MG	1A	3164	1/1	0.91	0.18	55,55,55,55	0
56	MG	1A	4081	1/1	0.91	0.12	39,39,39,39	0
56	MG	1A	4099	1/1	0.91	0.13	51,51,51,51	0
56	MG	1A	4105	1/1	0.91	0.09	41,41,41,41	0
56	MG	2q	3001	1/1	0.91	0.12	47,47,47,47	0
56	MG	2A	3457	1/1	0.91	0.32	59,59,59,59	0
56	MG	2A	3298	1/1	0.91	0.10	53,53,53,53	0
56	MG	1A	3720	1/1	0.91	0.12	52,52,52,52	0
56	MG	1A	3098	1/1	0.91	0.25	66,66,66,66	0
56	MG	2A	3464	1/1	0.91	0.16	60,60,60,60	0
56	MG	1A	4118	1/1	0.91	0.09	27,27,27,27	0
56	MG	1A	3556	1/1	0.91	0.09	49,49,49,49	0
56	MG	1A	3956	1/1	0.91	0.20	45,45,45,45	0
56	MG	2A	3726	1/1	0.91	0.10	55,55,55,55	0
56	MG	2A	3728	1/1	0.91	0.17	69,69,69,69	0
56	MG	2A	3733	1/1	0.91	0.09	57,57,57,57	0
56	MG	2x	102	1/1	0.91	0.11	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2a	1638	1/1	0.91	0.12	78,78,78,78	0
56	MG	1e	201	1/1	0.91	0.25	51,51,51,51	0
56	MG	2A	3472	1/1	0.91	0.25	58,58,58,58	0
56	MG	1f	3001	1/1	0.91	0.21	40,40,40,40	0
56	MG	2A	3737	1/1	0.91	0.10	52,52,52,52	0
56	MG	1A	3408	1/1	0.91	0.09	48,48,48,48	0
56	MG	1A	3409	1/1	0.91	0.18	52,52,52,52	0
56	MG	2A	3118	1/1	0.91	0.21	53,53,53,53	0
56	MG	2A	3746	1/1	0.91	0.08	76,76,76,76	0
57	MT9	2A	3897	33/33	0.91	0.15	37,55,61,64	0
59	ZN	24	501	1/1	0.91	0.10	119,119,119,119	0
56	MG	1A	3970	1/1	0.92	0.08	64,64,64,64	0
56	MG	2A	3349	1/1	0.92	0.17	63,63,63,63	0
56	MG	2A	3662	1/1	0.92	0.14	54,54,54,54	0
56	MG	2a	1617	1/1	0.92	0.23	59,59,59,59	0
56	MG	1U	201	1/1	0.92	0.17	33,33,33,33	0
56	MG	1A	3032	1/1	0.92	0.08	38,38,38,38	0
56	MG	1A	3974	1/1	0.92	0.12	22,22,22,22	0
56	MG	1a	1762	1/1	0.92	0.07	52,52,52,52	0
56	MG	1a	1764	1/1	0.92	0.09	63,63,63,63	0
56	MG	2a	1628	1/1	0.92	0.28	56,56,56,56	0
56	MG	1A	3977	1/1	0.92	0.10	22,22,22,22	0
56	MG	1A	3818	1/1	0.92	0.13	51,51,51,51	0
56	MG	2a	1631	1/1	0.92	0.11	61,61,61,61	0
56	MG	2A	3122	1/1	0.92	0.10	67,67,67,67	0
56	MG	2A	3124	1/1	0.92	0.24	68,68,68,68	0
56	MG	1A	3677	1/1	0.92	0.09	31,31,31,31	0
56	MG	1A	3483	1/1	0.92	0.14	60,60,60,60	0
56	MG	1A	3486	1/1	0.92	0.09	51,51,51,51	0
56	MG	1A	3559	1/1	0.92	0.20	36,36,36,36	0
56	MG	1a	1781	1/1	0.92	0.09	53,53,53,53	0
56	MG	1A	3256	1/1	0.92	0.20	65,65,65,65	0
56	MG	10	109	1/1	0.92	0.18	58,58,58,58	0
56	MG	1A	3222	1/1	0.92	0.18	31,31,31,31	0
56	MG	1A	3226	1/1	0.92	0.14	52,52,52,52	0
56	MG	2A	3156	1/1	0.92	0.12	58,58,58,58	0
56	MG	1A	3308	1/1	0.92	0.11	47,47,47,47	0
56	MG	2A	3383	1/1	0.92	0.08	51,51,51,51	0
56	MG	2A	3162	1/1	0.92	0.12	55,55,55,55	0
56	MG	2A	3164	1/1	0.92	0.20	66,66,66,66	0
56	MG	1A	3309	1/1	0.92	0.20	53,53,53,53	0
56	MG	1a	1807	1/1	0.92	0.23	54,54,54,54	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3840	1/1	0.92	0.12	47,47,47,47	0
56	MG	1a	1601	1/1	0.92	0.20	54,54,54,54	0
56	MG	1a	1603	1/1	0.92	0.15	57,57,57,57	0
56	MG	2A	3709	1/1	0.92	0.11	26,26,26,26	0
56	MG	1a	1813	1/1	0.92	0.16	50,50,50,50	0
56	MG	1A	3573	1/1	0.92	0.17	59,59,59,59	0
56	MG	1A	3143	1/1	0.92	0.24	46,46,46,46	0
56	MG	1A	3389	1/1	0.92	0.13	43,43,43,43	0
56	MG	2A	3406	1/1	0.92	0.17	59,59,59,59	0
56	MG	2a	1665	1/1	0.92	0.20	64,64,64,64	0
56	MG	2A	3407	1/1	0.92	0.07	61,61,61,61	0
56	MG	1A	3502	1/1	0.92	0.07	40,40,40,40	0
56	MG	2A	3410	1/1	0.92	0.17	55,55,55,55	0
56	MG	2A	3724	1/1	0.92	0.08	67,67,67,67	0
56	MG	2a	1671	1/1	0.92	0.21	57,57,57,57	0
56	MG	1a	1826	1/1	0.92	0.08	61,61,61,61	0
56	MG	1a	1829	1/1	0.92	0.09	57,57,57,57	0
56	MG	1A	4020	1/1	0.92	0.09	50,50,50,50	0
56	MG	1A	3856	1/1	0.92	0.14	33,33,33,33	0
56	MG	1e	202	1/1	0.92	0.10	69,69,69,69	0
56	MG	1A	3583	1/1	0.92	0.16	47,47,47,47	0
56	MG	1A	3709	1/1	0.92	0.11	55,55,55,55	0
56	MG	2A	3199	1/1	0.92	0.15	54,54,54,54	0
56	MG	1A	3860	1/1	0.92	0.15	38,38,38,38	0
56	MG	2A	3741	1/1	0.92	0.10	71,71,71,71	0
56	MG	2A	3427	1/1	0.92	0.11	60,60,60,60	0
56	MG	1a	1623	1/1	0.92	0.15	48,48,48,48	0
56	MG	1A	3862	1/1	0.92	0.08	35,35,35,35	0
56	MG	2A	3206	1/1	0.92	0.18	59,59,59,59	0
56	MG	1A	3866	1/1	0.92	0.07	26,26,26,26	0
56	MG	1A	3023	1/1	0.92	0.08	37,37,37,37	0
56	MG	2A	3209	1/1	0.92	0.10	68,68,68,68	0
56	MG	1A	3871	1/1	0.92	0.07	37,37,37,37	0
56	MG	1A	3585	1/1	0.92	0.09	40,40,40,40	0
56	MG	1A	4047	1/1	0.92	0.08	46,46,46,46	0
56	MG	1A	3721	1/1	0.92	0.11	60,60,60,60	0
56	MG	1A	4053	1/1	0.92	0.08	43,43,43,43	0
56	MG	1a	1638	1/1	0.92	0.24	53,53,53,53	0
56	MG	2A	3219	1/1	0.92	0.26	65,65,65,65	0
56	MG	1A	3313	1/1	0.92	0.08	38,38,38,38	0
56	MG	1A	3451	1/1	0.92	0.08	49,49,49,49	0
56	MG	1A	3320	1/1	0.92	0.09	57,57,57,57	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1648	1/1	0.92	0.08	57,57,57,57	0
56	MG	1A	3879	1/1	0.92	0.24	30,30,30,30	0
56	MG	2a	1720	1/1	0.92	0.22	59,59,59,59	0
56	MG	1A	3882	1/1	0.92	0.09	57,57,57,57	0
56	MG	2A	3237	1/1	0.92	0.08	44,44,44,44	0
56	MG	2A	3463	1/1	0.92	0.17	47,47,47,47	0
56	MG	2A	3240	1/1	0.92	0.12	54,54,54,54	0
56	MG	2A	3781	1/1	0.92	0.10	74,74,74,74	0
56	MG	2a	1727	1/1	0.92	0.09	51,51,51,51	0
56	MG	2A	3782	1/1	0.92	0.17	68,68,68,68	0
56	MG	2A	3784	1/1	0.92	0.09	72,72,72,72	0
56	MG	1A	3884	1/1	0.92	0.08	45,45,45,45	0
56	MG	1a	1653	1/1	0.92	0.15	70,70,70,70	0
56	MG	1x	104	1/1	0.92	0.14	64,64,64,64	0
56	MG	2A	3245	1/1	0.92	0.10	57,57,57,57	0
56	MG	2A	3246	1/1	0.92	0.13	35,35,35,35	0
56	MG	2A	3247	1/1	0.92	0.09	47,47,47,47	0
56	MG	1A	3610	1/1	0.92	0.07	30,30,30,30	0
56	MG	1A	3733	1/1	0.92	0.09	49,49,49,49	0
56	MG	1A	4071	1/1	0.92	0.10	40,40,40,40	0
56	MG	1A	3615	1/1	0.92	0.19	58,58,58,58	0
56	MG	2A	3254	1/1	0.92	0.24	61,61,61,61	0
56	MG	2a	1752	1/1	0.92	0.15	57,57,57,57	0
56	MG	1A	4074	1/1	0.92	0.12	47,47,47,47	0
56	MG	2A	3811	1/1	0.92	0.14	59,59,59,59	0
56	MG	1A	3109	1/1	0.92	0.14	31,31,31,31	0
56	MG	2A	3813	1/1	0.92	0.09	77,77,77,77	0
56	MG	2a	1764	1/1	0.92	0.11	72,72,72,72	0
56	MG	2A	3496	1/1	0.92	0.12	53,53,53,53	0
56	MG	1a	1666	1/1	0.92	0.09	64,64,64,64	0
56	MG	1A	3895	1/1	0.92	0.22	32,32,32,32	0
56	MG	1A	4092	1/1	0.92	0.14	55,55,55,55	0
56	MG	1a	1670	1/1	0.92	0.12	43,43,43,43	0
56	MG	1a	1671	1/1	0.92	0.20	48,48,48,48	0
56	MG	2A	3507	1/1	0.92	0.20	57,57,57,57	0
56	MG	1A	3512	1/1	0.92	0.09	52,52,52,52	0
56	MG	2A	3836	1/1	0.92	0.11	44,44,44,44	0
56	MG	2A	3838	1/1	0.92	0.09	53,53,53,53	0
56	MG	2a	1789	1/1	0.92	0.21	63,63,63,63	0
56	MG	1A	4100	1/1	0.92	0.10	63,63,63,63	0
56	MG	2A	3267	1/1	0.92	0.13	62,62,62,62	0
56	MG	1A	4103	1/1	0.92	0.14	60,60,60,60	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3621	1/1	0.92	0.08	23,23,23,23	0
56	MG	2A	3528	1/1	0.92	0.12	55,55,55,55	0
56	MG	2A	3273	1/1	0.92	0.10	59,59,59,59	0
56	MG	1A	4109	1/1	0.92	0.08	30,30,30,30	0
56	MG	2A	3534	1/1	0.92	0.12	39,39,39,39	0
56	MG	2A	3013	1/1	0.92	0.10	56,56,56,56	0
56	MG	2A	3536	1/1	0.92	0.08	73,73,73,73	0
56	MG	1A	3513	1/1	0.92	0.16	56,56,56,56	0
56	MG	2A	3280	1/1	0.92	0.25	62,62,62,62	0
56	MG	2A	3544	1/1	0.92	0.12	45,45,45,45	0
56	MG	2A	3018	1/1	0.92	0.10	44,44,44,44	0
56	MG	2A	3022	1/1	0.92	0.20	45,45,45,45	0
56	MG	1a	1679	1/1	0.92	0.23	56,56,56,56	0
56	MG	1A	3239	1/1	0.92	0.20	48,48,48,48	0
56	MG	2A	3888	1/1	0.92	0.12	71,71,71,71	0
56	MG	1A	3524	1/1	0.92	0.08	59,59,59,59	0
56	MG	2A	3560	1/1	0.92	0.10	71,71,71,71	0
56	MG	2A	3895	1/1	0.92	0.33	62,62,62,62	0
56	MG	2a	1816	1/1	0.92	0.18	53,53,53,53	0
56	MG	2A	3566	1/1	0.92	0.09	34,34,34,34	0
56	MG	1A	3268	1/1	0.92	0.16	44,44,44,44	0
56	MG	1a	1687	1/1	0.92	0.23	55,55,55,55	0
56	MG	2B	3004	1/1	0.92	0.20	66,66,66,66	0
56	MG	2B	3005	1/1	0.92	0.12	57,57,57,57	0
56	MG	2A	3572	1/1	0.92	0.09	58,58,58,58	0
56	MG	2A	3033	1/1	0.92	0.07	48,48,48,48	0
56	MG	2A	3297	1/1	0.92	0.10	52,52,52,52	0
56	MG	1A	3331	1/1	0.92	0.17	50,50,50,50	0
56	MG	1A	3909	1/1	0.92	0.08	36,36,36,36	0
56	MG	1A	3770	1/1	0.92	0.08	40,40,40,40	0
56	MG	1A	3527	1/1	0.92	0.06	40,40,40,40	0
56	MG	1A	3774	1/1	0.92	0.12	49,49,49,49	0
56	MG	1a	1696	1/1	0.92	0.09	50,50,50,50	0
56	MG	1B	220	1/1	0.92	0.11	58,58,58,58	0
56	MG	1A	3637	1/1	0.92	0.08	44,44,44,44	0
56	MG	1A	3924	1/1	0.92	0.14	39,39,39,39	0
56	MG	2A	3057	1/1	0.92	0.18	66,66,66,66	0
56	MG	1A	3928	1/1	0.92	0.17	51,51,51,51	0
56	MG	1B	230	1/1	0.92	0.11	53,53,53,53	0
56	MG	2A	3061	1/1	0.92	0.22	58,58,58,58	0
56	MG	2A	3602	1/1	0.92	0.16	55,55,55,55	0
56	MG	2E	306	1/1	0.92	0.07	40,40,40,40	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1708	1/1	0.92	0.32	49,49,49,49	0
56	MG	1a	1709	1/1	0.92	0.22	48,48,48,48	0
56	MG	2A	3317	1/1	0.92	0.18	69,69,69,69	0
56	MG	1A	3111	1/1	0.92	0.07	34,34,34,34	0
56	MG	1A	3533	1/1	0.92	0.18	43,43,43,43	0
56	MG	1A	3640	1/1	0.92	0.08	28,28,28,28	0
56	MG	2R	3001	1/1	0.92	0.18	65,65,65,65	0
56	MG	2R	3002	1/1	0.92	0.10	52,52,52,52	0
56	MG	2T	3001	1/1	0.92	0.10	52,52,52,52	0
56	MG	2A	3612	1/1	0.92	0.10	58,58,58,58	0
56	MG	2U	201	1/1	0.92	0.20	47,47,47,47	0
56	MG	1A	3534	1/1	0.92	0.14	40,40,40,40	0
56	MG	1A	3796	1/1	0.92	0.14	64,64,64,64	0
56	MG	2A	3618	1/1	0.92	0.11	41,41,41,41	0
56	MG	1a	1724	1/1	0.92	0.10	43,43,43,43	0
56	MG	1a	1727	1/1	0.92	0.10	65,65,65,65	0
56	MG	2A	3621	1/1	0.92	0.12	50,50,50,50	0
56	MG	2w	3006	1/1	0.92	0.22	77,77,77,77	0
56	MG	2A	3082	1/1	0.92	0.21	54,54,54,54	0
56	MG	2A	3086	1/1	0.92	0.08	41,41,41,41	0
56	MG	1A	3272	1/1	0.92	0.19	42,42,42,42	0
56	MG	1A	3950	1/1	0.92	0.20	57,57,57,57	0
56	MG	1A	3090	1/1	0.92	0.17	47,47,47,47	0
56	MG	1I	3001	1/1	0.92	0.08	61,61,61,61	0
56	MG	1A	3413	1/1	0.92	0.09	51,51,51,51	0
56	MG	2A	3337	1/1	0.92	0.13	68,68,68,68	0
56	MG	1A	3242	1/1	0.92	0.21	51,51,51,51	0
56	MG	1a	1744	1/1	0.92	0.13	60,60,60,60	0
56	MG	1A	3806	1/1	0.92	0.11	35,35,35,35	0
56	MG	1A	3177	1/1	0.92	0.14	55,55,55,55	0
56	MG	1A	3477	1/1	0.92	0.08	44,44,44,44	0
56	MG	1A	3815	1/1	0.92	0.07	53,53,53,53	0
56	MG	1a	1752	1/1	0.93	0.09	66,66,66,66	0
56	MG	2U	205	1/1	0.93	0.10	53,53,53,53	0
56	MG	1a	1753	1/1	0.93	0.12	56,56,56,56	0
56	MG	1A	3448	1/1	0.93	0.10	50,50,50,50	0
56	MG	2W	202	1/1	0.93	0.28	40,40,40,40	0
56	MG	2A	3611	1/1	0.93	0.17	62,62,62,62	0
56	MG	25	101	1/1	0.93	0.22	56,56,56,56	0
56	MG	1A	3729	1/1	0.93	0.10	52,52,52,52	0
56	MG	2A	3613	1/1	0.93	0.09	24,24,24,24	0
56	MG	1A	3385	1/1	0.93	0.17	37,37,37,37	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3246	1/1	0.93	0.07	52,52,52,52	0
56	MG	2A	3617	1/1	0.93	0.12	44,44,44,44	0
56	MG	1A	3521	1/1	0.93	0.14	56,56,56,56	0
56	MG	1A	3522	1/1	0.93	0.10	58,58,58,58	0
56	MG	11	103	1/1	0.93	0.10	45,45,45,45	0
56	MG	1A	4019	1/1	0.93	0.11	50,50,50,50	0
56	MG	15	103	1/1	0.93	0.08	45,45,45,45	0
56	MG	1A	3388	1/1	0.93	0.17	29,29,29,29	0
56	MG	2A	3626	1/1	0.93	0.15	50,50,50,50	0
56	MG	16	104	1/1	0.93	0.17	39,39,39,39	0
56	MG	2A	3341	1/1	0.93	0.10	50,50,50,50	0
56	MG	2A	3635	1/1	0.93	0.14	39,39,39,39	0
56	MG	1A	4021	1/1	0.93	0.07	60,60,60,60	0
56	MG	17	102	1/1	0.93	0.16	38,38,38,38	0
56	MG	2A	3115	1/1	0.93	0.29	58,58,58,58	0
56	MG	1a	1776	1/1	0.93	0.10	46,46,46,46	0
56	MG	2a	1618	1/1	0.93	0.12	63,63,63,63	0
56	MG	2a	1619	1/1	0.93	0.12	48,48,48,48	0
56	MG	2A	3117	1/1	0.93	0.17	57,57,57,57	0
56	MG	1A	3319	1/1	0.93	0.18	36,36,36,36	0
56	MG	1a	1779	1/1	0.93	0.10	50,50,50,50	0
56	MG	2a	1623	1/1	0.93	0.09	69,69,69,69	0
56	MG	2a	1624	1/1	0.93	0.08	48,48,48,48	0
56	MG	1a	1780	1/1	0.93	0.10	51,51,51,51	0
56	MG	2A	3352	1/1	0.93	0.12	49,49,49,49	0
56	MG	1A	3749	1/1	0.93	0.12	19,19,19,19	0
56	MG	2A	3657	1/1	0.93	0.13	55,55,55,55	0
56	MG	1A	3248	1/1	0.93	0.09	56,56,56,56	0
56	MG	2A	3127	1/1	0.93	0.07	60,60,60,60	0
56	MG	2A	3661	1/1	0.93	0.10	63,63,63,63	0
56	MG	1A	3321	1/1	0.93	0.16	49,49,49,49	0
56	MG	1A	3273	1/1	0.93	0.13	41,41,41,41	0
56	MG	1A	3324	1/1	0.93	0.22	54,54,54,54	0
56	MG	1A	3466	1/1	0.93	0.09	48,48,48,48	0
56	MG	2A	3146	1/1	0.93	0.24	58,58,58,58	0
56	MG	2A	3147	1/1	0.93	0.14	56,56,56,56	0
56	MG	1A	4038	1/1	0.93	0.08	49,49,49,49	0
56	MG	2A	3672	1/1	0.93	0.08	42,42,42,42	0
56	MG	2A	3673	1/1	0.93	0.15	65,65,65,65	0
56	MG	1a	1806	1/1	0.93	0.12	49,49,49,49	0
56	MG	1A	3768	1/1	0.93	0.07	24,24,24,24	0
56	MG	1a	1808	1/1	0.93	0.24	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3467	1/1	0.93	0.11	41,41,41,41	0
56	MG	2A	3158	1/1	0.93	0.15	46,46,46,46	0
56	MG	2A	3374	1/1	0.93	0.12	48,48,48,48	0
56	MG	1A	3327	1/1	0.93	0.09	34,34,34,34	0
56	MG	2A	3376	1/1	0.93	0.06	62,62,62,62	0
56	MG	2A	3685	1/1	0.93	0.14	55,55,55,55	0
56	MG	1A	3283	1/1	0.93	0.10	51,51,51,51	0
56	MG	2A	3379	1/1	0.93	0.12	61,61,61,61	0
56	MG	1A	4052	1/1	0.93	0.08	18,18,18,18	0
56	MG	2A	3381	1/1	0.93	0.13	58,58,58,58	0
56	MG	2a	1657	1/1	0.93	0.12	65,65,65,65	0
56	MG	1A	3775	1/1	0.93	0.10	45,45,45,45	0
56	MG	2A	3692	1/1	0.93	0.15	65,65,65,65	0
56	MG	2A	3166	1/1	0.93	0.15	47,47,47,47	0
56	MG	2a	1661	1/1	0.93	0.12	58,58,58,58	0
56	MG	1A	3642	1/1	0.93	0.07	20,20,20,20	0
56	MG	1A	3780	1/1	0.93	0.08	36,36,36,36	0
56	MG	1a	1626	1/1	0.93	0.12	58,58,58,58	0
56	MG	1A	3546	1/1	0.93	0.20	42,42,42,42	0
56	MG	1A	3903	1/1	0.93	0.07	30,30,30,30	0
56	MG	2A	3393	1/1	0.93	0.12	55,55,55,55	0
56	MG	1A	3783	1/1	0.93	0.06	48,48,48,48	0
56	MG	2A	3176	1/1	0.93	0.09	41,41,41,41	0
56	MG	1A	3403	1/1	0.93	0.11	46,46,46,46	0
56	MG	2a	1675	1/1	0.93	0.21	63,63,63,63	0
56	MG	1A	3132	1/1	0.93	0.12	44,44,44,44	0
56	MG	2A	3398	1/1	0.93	0.17	59,59,59,59	0
56	MG	2A	3400	1/1	0.93	0.19	48,48,48,48	0
56	MG	1a	1635	1/1	0.93	0.19	49,49,49,49	0
56	MG	1a	1636	1/1	0.93	0.17	58,58,58,58	0
56	MG	1A	3091	1/1	0.93	0.13	43,43,43,43	0
56	MG	1A	3551	1/1	0.93	0.17	55,55,55,55	0
56	MG	1A	3333	1/1	0.93	0.37	69,69,69,69	0
56	MG	1A	3480	1/1	0.93	0.14	46,46,46,46	0
56	MG	1a	1645	1/1	0.93	0.20	50,50,50,50	0
56	MG	2A	3413	1/1	0.93	0.10	55,55,55,55	0
56	MG	2a	1698	1/1	0.93	0.13	51,51,51,51	0
56	MG	1A	3672	1/1	0.93	0.09	24,24,24,24	0
56	MG	2A	3198	1/1	0.93	0.27	49,49,49,49	0
56	MG	1A	3096	1/1	0.93	0.13	56,56,56,56	0
56	MG	2A	3417	1/1	0.93	0.13	64,64,64,64	0
56	MG	1A	3925	1/1	0.93	0.08	59,59,59,59	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	4094	1/1	0.93	0.11	39,39,39,39	0
56	MG	1A	4096	1/1	0.93	0.09	40,40,40,40	0
56	MG	2A	3740	1/1	0.93	0.08	45,45,45,45	0
56	MG	1A	3675	1/1	0.93	0.07	51,51,51,51	0
56	MG	1A	3002	1/1	0.93	0.09	55,55,55,55	0
56	MG	2A	3425	1/1	0.93	0.12	58,58,58,58	0
56	MG	1A	4101	1/1	0.93	0.10	43,43,43,43	0
56	MG	1A	3809	1/1	0.93	0.09	47,47,47,47	0
56	MG	1a	1660	1/1	0.93	0.08	54,54,54,54	0
56	MG	1a	1661	1/1	0.93	0.14	56,56,56,56	0
56	MG	2a	1718	1/1	0.93	0.14	69,69,69,69	0
56	MG	1A	3300	1/1	0.93	0.08	29,29,29,29	0
56	MG	1A	4106	1/1	0.93	0.16	47,47,47,47	0
56	MG	1A	3934	1/1	0.93	0.08	41,41,41,41	0
56	MG	2A	3756	1/1	0.93	0.07	55,55,55,55	0
56	MG	1x	103	1/1	0.93	0.09	56,56,56,56	0
56	MG	1A	3811	1/1	0.93	0.17	47,47,47,47	0
56	MG	2A	3220	1/1	0.93	0.10	47,47,47,47	0
56	MG	2A	3439	1/1	0.93	0.28	55,55,55,55	0
56	MG	2A	3221	1/1	0.93	0.12	42,42,42,42	0
56	MG	2A	3442	1/1	0.93	0.21	45,45,45,45	0
56	MG	2A	3443	1/1	0.93	0.21	44,44,44,44	0
56	MG	2A	3444	1/1	0.93	0.24	49,49,49,49	0
56	MG	2a	1733	1/1	0.93	0.09	61,61,61,61	0
56	MG	2A	3768	1/1	0.93	0.13	54,54,54,54	0
56	MG	2a	1737	1/1	0.93	0.12	86,86,86,86	0
56	MG	1A	4111	1/1	0.93	0.10	41,41,41,41	0
56	MG	2A	3771	1/1	0.93	0.16	65,65,65,65	0
56	MG	2a	1741	1/1	0.93	0.09	77,77,77,77	0
56	MG	2A	3446	1/1	0.93	0.17	41,41,41,41	0
56	MG	1A	3144	1/1	0.93	0.30	31,31,31,31	0
56	MG	2A	3226	1/1	0.93	0.14	54,54,54,54	0
56	MG	1a	1669	1/1	0.93	0.12	56,56,56,56	0
56	MG	2A	3452	1/1	0.93	0.23	68,68,68,68	0
56	MG	2A	3228	1/1	0.93	0.08	51,51,51,51	0
56	MG	2A	3229	1/1	0.93	0.21	58,58,58,58	0
56	MG	1A	3945	1/1	0.93	0.11	59,59,59,59	0
56	MG	1A	3417	1/1	0.93	0.13	52,52,52,52	0
56	MG	2A	3788	1/1	0.93	0.11	54,54,54,54	0
56	MG	2A	3234	1/1	0.93	0.09	62,62,62,62	0
56	MG	1x	112	1/1	0.93	0.14	64,64,64,64	0
56	MG	1A	3949	1/1	0.93	0.08	19,19,19,19	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1673	1/1	0.93	0.24	55,55,55,55	0
56	MG	2A	3241	1/1	0.93	0.12	58,58,58,58	0
56	MG	1A	3564	1/1	0.93	0.25	43,43,43,43	0
56	MG	2A	3800	1/1	0.93	0.09	70,70,70,70	0
56	MG	1x	116	1/1	0.93	0.09	62,62,62,62	0
56	MG	2a	1775	1/1	0.93	0.09	62,62,62,62	0
56	MG	2a	1779	1/1	0.93	0.11	60,60,60,60	0
56	MG	1A	3687	1/1	0.93	0.10	29,29,29,29	0
56	MG	1A	3420	1/1	0.93	0.12	50,50,50,50	0
56	MG	1A	3822	1/1	0.93	0.07	41,41,41,41	0
56	MG	2A	3476	1/1	0.93	0.11	57,57,57,57	0
56	MG	1B	219	1/1	0.93	0.15	52,52,52,52	0
56	MG	2A	3479	1/1	0.93	0.17	56,56,56,56	0
56	MG	2A	3480	1/1	0.93	0.16	65,65,65,65	0
56	MG	1A	3823	1/1	0.93	0.18	47,47,47,47	0
56	MG	2A	3482	1/1	0.93	0.18	43,43,43,43	0
56	MG	2A	3002	1/1	0.93	0.28	55,55,55,55	0
56	MG	2A	3486	1/1	0.93	0.09	66,66,66,66	0
56	MG	1A	3568	1/1	0.93	0.26	44,44,44,44	0
56	MG	2a	1797	1/1	0.93	0.10	65,65,65,65	0
56	MG	2A	3829	1/1	0.93	0.09	60,60,60,60	0
56	MG	1A	3037	1/1	0.93	0.18	54,54,54,54	0
56	MG	2A	3490	1/1	0.93	0.21	57,57,57,57	0
56	MG	1A	3427	1/1	0.93	0.21	50,50,50,50	0
56	MG	2A	3011	1/1	0.93	0.10	49,49,49,49	0
56	MG	1A	3046	1/1	0.93	0.06	15,15,15,15	0
56	MG	2A	3014	1/1	0.93	0.13	51,51,51,51	0
56	MG	1A	3307	1/1	0.93	0.07	44,44,44,44	0
56	MG	1A	3972	1/1	0.93	0.14	63,63,63,63	0
56	MG	1D	304	1/1	0.93	0.07	15,15,15,15	0
56	MG	2a	1810	1/1	0.93	0.14	78,78,78,78	0
56	MG	1a	1692	1/1	0.93	0.14	63,63,63,63	0
56	MG	1A	3700	1/1	0.93	0.14	44,44,44,44	0
56	MG	1A	3063	1/1	0.93	0.08	30,30,30,30	0
56	MG	1D	312	1/1	0.93	0.21	34,34,34,34	0
56	MG	2A	3515	1/1	0.93	0.10	64,64,64,64	0
56	MG	2A	3517	1/1	0.93	0.15	51,51,51,51	0
56	MG	2A	3031	1/1	0.93	0.15	45,45,45,45	0
56	MG	2A	3876	1/1	0.93	0.07	71,71,71,71	0
56	MG	2A	3877	1/1	0.93	0.16	47,47,47,47	0
56	MG	2A	3878	1/1	0.93	0.13	59,59,59,59	0
56	MG	1a	1697	1/1	0.93	0.17	55,55,55,55	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3271	1/1	0.93	0.20	68,68,68,68	0
56	MG	2A	3522	1/1	0.93	0.09	60,60,60,60	0
56	MG	1E	307	1/1	0.93	0.11	50,50,50,50	0
56	MG	2A	3034	1/1	0.93	0.13	46,46,46,46	0
56	MG	1E	310	1/1	0.93	0.16	35,35,35,35	0
56	MG	1A	3978	1/1	0.93	0.09	26,26,26,26	0
56	MG	2A	3891	1/1	0.93	0.08	49,49,49,49	0
56	MG	1A	3265	1/1	0.93	0.13	46,46,46,46	0
56	MG	1A	3980	1/1	0.93	0.07	83,83,83,83	0
56	MG	2A	3282	1/1	0.93	0.07	58,58,58,58	0
56	MG	1A	3504	1/1	0.93	0.08	50,50,50,50	0
56	MG	2a	1836	1/1	0.93	0.10	65,65,65,65	0
56	MG	2A	3901	1/1	0.93	0.12	67,67,67,67	0
56	MG	1A	3211	1/1	0.93	0.13	46,46,46,46	0
56	MG	1N	204	1/1	0.93	0.24	43,43,43,43	0
56	MG	1a	1711	1/1	0.93	0.17	60,60,60,60	0
56	MG	2A	3547	1/1	0.93	0.07	52,52,52,52	0
56	MG	2A	3287	1/1	0.93	0.12	55,55,55,55	0
56	MG	2A	3052	1/1	0.93	0.10	48,48,48,48	0
56	MG	2A	3554	1/1	0.93	0.10	45,45,45,45	0
56	MG	2A	3290	1/1	0.93	0.09	64,64,64,64	0
56	MG	2A	3559	1/1	0.93	0.15	47,47,47,47	0
56	MG	2B	3012	1/1	0.93	0.14	69,69,69,69	0
56	MG	2A	3053	1/1	0.93	0.18	64,64,64,64	0
56	MG	2l	3002	1/1	0.93	0.06	62,62,62,62	0
56	MG	2A	3563	1/1	0.93	0.11	65,65,65,65	0
56	MG	1A	3372	1/1	0.93	0.14	41,41,41,41	0
56	MG	2A	3295	1/1	0.93	0.18	55,55,55,55	0
56	MG	2B	3018	1/1	0.93	0.12	79,79,79,79	0
56	MG	1a	1714	1/1	0.93	0.16	48,48,48,48	0
56	MG	1O	203	1/1	0.93	0.10	53,53,53,53	0
56	MG	1A	3849	1/1	0.93	0.13	38,38,38,38	0
56	MG	1O	205	1/1	0.93	0.22	45,45,45,45	0
56	MG	1A	3590	1/1	0.93	0.09	58,58,58,58	0
56	MG	1A	3989	1/1	0.93	0.09	53,53,53,53	0
56	MG	1A	3373	1/1	0.93	0.12	40,40,40,40	0
56	MG	1A	3163	1/1	0.93	0.14	41,41,41,41	0
56	MG	1A	3996	1/1	0.93	0.14	60,60,60,60	0
56	MG	1A	3601	1/1	0.93	0.12	61,61,61,61	0
56	MG	1a	1734	1/1	0.93	0.13	56,56,56,56	0
56	MG	1A	3999	1/1	0.93	0.08	57,57,57,57	0
56	MG	1A	4002	1/1	0.93	0.06	29,29,29,29	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3311	1/1	0.93	0.15	56,56,56,56	0
56	MG	2Q	3002	1/1	0.93	0.28	54,54,54,54	0
56	MG	2A	3080	1/1	0.93	0.19	52,52,52,52	0
56	MG	2A	3081	1/1	0.93	0.09	44,44,44,44	0
56	MG	2y	3003	1/1	0.93	0.11	63,63,63,63	0
56	MG	1X	105	1/1	0.93	0.11	48,48,48,48	0
56	MG	2A	3085	1/1	0.93	0.07	41,41,41,41	0
56	MG	1A	4003	1/1	0.93	0.09	27,27,27,27	0
56	MG	2A	3087	1/1	0.93	0.09	50,50,50,50	0
56	MG	1A	4007	1/1	0.93	0.09	29,29,29,29	0
56	MG	1A	4135	1/1	0.94	0.10	36,36,36,36	0
56	MG	2A	3392	1/1	0.94	0.08	52,52,52,52	0
56	MG	1A	3290	1/1	0.94	0.13	38,38,38,38	0
56	MG	1B	206	1/1	0.94	0.06	34,34,34,34	0
56	MG	1A	3338	1/1	0.94	0.11	45,45,45,45	0
56	MG	1a	1657	1/1	0.94	0.08	54,54,54,54	0
56	MG	1a	1658	1/1	0.94	0.14	61,61,61,61	0
56	MG	2A	3200	1/1	0.94	0.15	66,66,66,66	0
56	MG	1B	208	1/1	0.94	0.17	57,57,57,57	0
56	MG	2A	3202	1/1	0.94	0.12	46,46,46,46	0
56	MG	2A	3203	1/1	0.94	0.21	47,47,47,47	0
56	MG	1A	3410	1/1	0.94	0.10	37,37,37,37	0
56	MG	1A	3825	1/1	0.94	0.07	54,54,54,54	0
56	MG	1B	216	1/1	0.94	0.09	34,34,34,34	0
56	MG	1A	3411	1/1	0.94	0.10	58,58,58,58	0
56	MG	1A	3083	1/1	0.94	0.20	34,34,34,34	0
56	MG	1A	3698	1/1	0.94	0.09	33,33,33,33	0
56	MG	2a	1626	1/1	0.94	0.13	68,68,68,68	0
56	MG	1A	3975	1/1	0.94	0.09	36,36,36,36	0
56	MG	1B	222	1/1	0.94	0.07	56,56,56,56	0
56	MG	1A	3197	1/1	0.94	0.08	31,31,31,31	0
56	MG	1A	3493	1/1	0.94	0.16	52,52,52,52	0
56	MG	1A	3244	1/1	0.94	0.07	61,61,61,61	0
56	MG	2A	3420	1/1	0.94	0.16	72,72,72,72	0
56	MG	1B	232	1/1	0.94	0.08	52,52,52,52	0
56	MG	1A	3110	1/1	0.94	0.11	41,41,41,41	0
56	MG	1A	3839	1/1	0.94	0.09	28,28,28,28	0
56	MG	1D	301	1/1	0.94	0.08	43,43,43,43	0
56	MG	2a	1637	1/1	0.94	0.38	63,63,63,63	0
56	MG	1A	3500	1/1	0.94	0.16	33,33,33,33	0
56	MG	1A	3419	1/1	0.94	0.06	39,39,39,39	0
56	MG	1A	3714	1/1	0.94	0.10	62,62,62,62	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3715	1/1	0.94	0.09	45,45,45,45	0
56	MG	1E	301	1/1	0.94	0.09	49,49,49,49	0
56	MG	1E	305	1/1	0.94	0.19	54,54,54,54	0
56	MG	1A	3848	1/1	0.94	0.07	33,33,33,33	0
56	MG	2A	3231	1/1	0.94	0.50	49,49,49,49	0
56	MG	1A	3147	1/1	0.94	0.13	33,33,33,33	0
56	MG	2a	1647	1/1	0.94	0.10	55,55,55,55	0
56	MG	2A	3436	1/1	0.94	0.21	43,43,43,43	0
56	MG	2A	3233	1/1	0.94	0.10	52,52,52,52	0
56	MG	1a	1689	1/1	0.94	0.20	65,65,65,65	0
56	MG	2A	3721	1/1	0.94	0.16	67,67,67,67	0
56	MG	2A	3235	1/1	0.94	0.14	42,42,42,42	0
56	MG	1A	3100	1/1	0.94	0.20	32,32,32,32	0
56	MG	1F	306	1/1	0.94	0.11	47,47,47,47	0
56	MG	1A	3206	1/1	0.94	0.15	46,46,46,46	0
56	MG	2A	3727	1/1	0.94	0.09	42,42,42,42	0
56	MG	1G	3002	1/1	0.94	0.08	46,46,46,46	0
56	MG	2A	3731	1/1	0.94	0.11	62,62,62,62	0
56	MG	1A	3209	1/1	0.94	0.08	51,51,51,51	0
56	MG	1A	3724	1/1	0.94	0.09	13,13,13,13	0
56	MG	1G	3005	1/1	0.94	0.12	58,58,58,58	0
56	MG	1A	4000	1/1	0.94	0.06	29,29,29,29	0
56	MG	1A	4001	1/1	0.94	0.08	26,26,26,26	0
56	MG	2A	3451	1/1	0.94	0.29	61,61,61,61	0
56	MG	1O	201	1/1	0.94	0.21	54,54,54,54	0
56	MG	1A	3362	1/1	0.94	0.22	50,50,50,50	0
56	MG	1a	1702	1/1	0.94	0.24	54,54,54,54	0
56	MG	1a	1705	1/1	0.94	0.31	50,50,50,50	0
56	MG	1A	3602	1/1	0.94	0.10	29,29,29,29	0
56	MG	2A	3747	1/1	0.94	0.09	55,55,55,55	0
56	MG	2A	3748	1/1	0.94	0.15	40,40,40,40	0
56	MG	2a	1676	1/1	0.94	0.15	63,63,63,63	0
56	MG	2a	1677	1/1	0.94	0.16	55,55,55,55	0
56	MG	1A	3605	1/1	0.94	0.14	37,37,37,37	0
56	MG	2A	3750	1/1	0.94	0.12	49,49,49,49	0
56	MG	1A	3363	1/1	0.94	0.07	32,32,32,32	0
56	MG	1A	3869	1/1	0.94	0.07	32,32,32,32	0
56	MG	1A	3436	1/1	0.94	0.12	61,61,61,61	0
56	MG	2A	3258	1/1	0.94	0.11	60,60,60,60	0
56	MG	1A	3150	1/1	0.94	0.10	45,45,45,45	0
56	MG	2a	1685	1/1	0.94	0.14	65,65,65,65	0
56	MG	2A	3467	1/1	0.94	0.08	40,40,40,40	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2a	1689	1/1	0.94	0.16	54,54,54,54	0
56	MG	1Q	204	1/1	0.94	0.07	38,38,38,38	0
56	MG	2a	1691	1/1	0.94	0.16	63,63,63,63	0
56	MG	2a	1692	1/1	0.94	0.10	52,52,52,52	0
56	MG	2a	1693	1/1	0.94	0.12	40,40,40,40	0
56	MG	2A	3261	1/1	0.94	0.13	71,71,71,71	0
56	MG	2A	3048	1/1	0.94	0.28	55,55,55,55	0
56	MG	2a	1697	1/1	0.94	0.24	70,70,70,70	0
56	MG	1A	4016	1/1	0.94	0.09	44,44,44,44	0
56	MG	1S	3001	1/1	0.94	0.09	42,42,42,42	0
56	MG	1a	1722	1/1	0.94	0.08	53,53,53,53	0
56	MG	2a	1701	1/1	0.94	0.12	55,55,55,55	0
56	MG	2A	3478	1/1	0.94	0.34	49,49,49,49	0
56	MG	1A	3742	1/1	0.94	0.12	49,49,49,49	0
56	MG	1A	3744	1/1	0.94	0.13	19,19,19,19	0
56	MG	2A	3055	1/1	0.94	0.12	56,56,56,56	0
56	MG	2a	1706	1/1	0.94	0.15	64,64,64,64	0
56	MG	2A	3056	1/1	0.94	0.09	45,45,45,45	0
56	MG	2A	3483	1/1	0.94	0.17	56,56,56,56	0
56	MG	1A	3369	1/1	0.94	0.10	56,56,56,56	0
56	MG	2A	3485	1/1	0.94	0.11	47,47,47,47	0
56	MG	1V	203	1/1	0.94	0.07	49,49,49,49	0
56	MG	2a	1712	1/1	0.94	0.09	61,61,61,61	0
56	MG	1W	201	1/1	0.94	0.21	50,50,50,50	0
56	MG	1A	3153	1/1	0.94	0.09	50,50,50,50	0
56	MG	1W	207	1/1	0.94	0.07	35,35,35,35	0
56	MG	1A	3112	1/1	0.94	0.14	42,42,42,42	0
56	MG	2A	3492	1/1	0.94	0.09	59,59,59,59	0
56	MG	1A	3445	1/1	0.94	0.11	24,24,24,24	0
56	MG	2A	3789	1/1	0.94	0.17	67,67,67,67	0
56	MG	1A	3523	1/1	0.94	0.10	53,53,53,53	0
56	MG	1a	1739	1/1	0.94	0.18	64,64,64,64	0
56	MG	2a	1723	1/1	0.94	0.16	47,47,47,47	0
56	MG	1a	1743	1/1	0.94	0.06	58,58,58,58	0
56	MG	1A	3117	1/1	0.94	0.09	35,35,35,35	0
56	MG	2A	3796	1/1	0.94	0.11	70,70,70,70	0
56	MG	2A	3072	1/1	0.94	0.10	53,53,53,53	0
56	MG	1A	3887	1/1	0.94	0.26	20,20,20,20	0
56	MG	2A	3077	1/1	0.94	0.09	53,53,53,53	0
56	MG	1A	3888	1/1	0.94	0.13	59,59,59,59	0
56	MG	2A	3803	1/1	0.94	0.07	70,70,70,70	0
56	MG	1a	1750	1/1	0.94	0.15	63,63,63,63	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1751	1/1	0.94	0.10	44,44,44,44	0
56	MG	2A	3808	1/1	0.94	0.09	42,42,42,42	0
56	MG	1A	3447	1/1	0.94	0.14	42,42,42,42	0
56	MG	2A	3083	1/1	0.94	0.13	53,53,53,53	0
56	MG	1A	3376	1/1	0.94	0.10	44,44,44,44	0
56	MG	1A	3634	1/1	0.94	0.09	34,34,34,34	0
56	MG	11	102	1/1	0.94	0.16	39,39,39,39	0
56	MG	2A	3525	1/1	0.94	0.10	41,41,41,41	0
56	MG	1A	3377	1/1	0.94	0.10	49,49,49,49	0
56	MG	11	104	1/1	0.94	0.11	69,69,69,69	0
56	MG	2a	1749	1/1	0.94	0.10	64,64,64,64	0
56	MG	1A	3528	1/1	0.94	0.08	52,52,52,52	0
56	MG	2A	3820	1/1	0.94	0.09	74,74,74,74	0
56	MG	2A	3533	1/1	0.94	0.12	48,48,48,48	0
56	MG	13	103	1/1	0.94	0.09	45,45,45,45	0
56	MG	2A	3093	1/1	0.94	0.32	53,53,53,53	0
56	MG	1a	1761	1/1	0.94	0.08	57,57,57,57	0
56	MG	1A	3380	1/1	0.94	0.12	38,38,38,38	0
56	MG	1A	4048	1/1	0.94	0.14	54,54,54,54	0
56	MG	2A	3098	1/1	0.94	0.08	51,51,51,51	0
56	MG	2A	3099	1/1	0.94	0.11	49,49,49,49	0
56	MG	2A	3313	1/1	0.94	0.20	70,70,70,70	0
56	MG	1A	4050	1/1	0.94	0.15	50,50,50,50	0
56	MG	1A	3263	1/1	0.94	0.10	34,34,34,34	0
56	MG	2A	3860	1/1	0.94	0.12	67,67,67,67	0
56	MG	2a	1777	1/1	0.94	0.22	64,64,64,64	0
56	MG	2A	3552	1/1	0.94	0.10	51,51,51,51	0
56	MG	2a	1780	1/1	0.94	0.14	57,57,57,57	0
56	MG	1A	3048	1/1	0.94	0.16	22,22,22,22	0
56	MG	2A	3103	1/1	0.94	0.17	36,36,36,36	0
56	MG	1A	3776	1/1	0.94	0.10	23,23,23,23	0
56	MG	2A	3106	1/1	0.94	0.16	49,49,49,49	0
56	MG	1A	3455	1/1	0.94	0.10	59,59,59,59	0
56	MG	18	102	1/1	0.94	0.16	40,40,40,40	0
56	MG	2A	3873	1/1	0.94	0.10	46,46,46,46	0
56	MG	1A	3315	1/1	0.94	0.16	52,52,52,52	0
56	MG	2A	3325	1/1	0.94	0.17	53,53,53,53	0
56	MG	1a	1602	1/1	0.94	0.25	64,64,64,64	0
56	MG	2A	3571	1/1	0.94	0.07	36,36,36,36	0
56	MG	1A	4058	1/1	0.94	0.09	36,36,36,36	0
56	MG	1A	3317	1/1	0.94	0.19	33,33,33,33	0
56	MG	1A	3657	1/1	0.94	0.11	29,29,29,29	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3138	1/1	0.94	0.08	37,37,37,37	0
56	MG	2A	3885	1/1	0.94	0.10	49,49,49,49	0
56	MG	2A	3578	1/1	0.94	0.07	51,51,51,51	0
56	MG	1a	1795	1/1	0.94	0.15	61,61,61,61	0
56	MG	2A	3890	1/1	0.94	0.27	52,52,52,52	0
56	MG	1A	3914	1/1	0.94	0.09	39,39,39,39	0
56	MG	1A	3174	1/1	0.94	0.16	31,31,31,31	0
56	MG	1a	1612	1/1	0.94	0.07	51,51,51,51	0
56	MG	1A	3234	1/1	0.94	0.08	34,34,34,34	0
56	MG	2A	3339	1/1	0.94	0.15	62,62,62,62	0
56	MG	2A	3899	1/1	0.94	0.11	36,36,36,36	0
56	MG	2A	3340	1/1	0.94	0.14	74,74,74,74	0
56	MG	1A	3175	1/1	0.94	0.15	41,41,41,41	0
56	MG	2A	3131	1/1	0.94	0.14	48,48,48,48	0
56	MG	2B	3003	1/1	0.94	0.16	60,60,60,60	0
56	MG	1A	3395	1/1	0.94	0.16	43,43,43,43	0
56	MG	2A	3596	1/1	0.94	0.11	57,57,57,57	0
56	MG	1a	1620	1/1	0.94	0.15	56,56,56,56	0
56	MG	2A	3600	1/1	0.94	0.21	61,61,61,61	0
56	MG	1A	3799	1/1	0.94	0.40	33,33,33,33	0
56	MG	1a	1622	1/1	0.94	0.14	41,41,41,41	0
56	MG	2B	3010	1/1	0.94	0.07	68,68,68,68	0
56	MG	2A	3140	1/1	0.94	0.19	71,71,71,71	0
56	MG	2A	3144	1/1	0.94	0.15	38,38,38,38	0
56	MG	1A	3673	1/1	0.94	0.09	16,16,16,16	0
56	MG	2A	3351	1/1	0.94	0.26	64,64,64,64	0
56	MG	1A	3552	1/1	0.94	0.09	49,49,49,49	0
56	MG	2A	3353	1/1	0.94	0.14	56,56,56,56	0
56	MG	2A	3148	1/1	0.94	0.21	58,58,58,58	0
56	MG	2A	3150	1/1	0.94	0.13	41,41,41,41	0
56	MG	2a	1833	1/1	0.94	0.10	52,52,52,52	0
56	MG	1A	4085	1/1	0.94	0.14	46,46,46,46	0
56	MG	1a	1817	1/1	0.94	0.25	59,59,59,59	0
56	MG	1A	4086	1/1	0.94	0.09	19,19,19,19	0
56	MG	1A	3802	1/1	0.94	0.07	15,15,15,15	0
56	MG	2a	1838	1/1	0.94	0.14	64,64,64,64	0
56	MG	1A	3804	1/1	0.94	0.08	28,28,28,28	0
56	MG	1A	3933	1/1	0.94	0.10	52,52,52,52	0
56	MG	1A	3107	1/1	0.94	0.04	31,31,31,31	0
56	MG	2a	1842	1/1	0.94	0.12	49,49,49,49	0
56	MG	1b	3001	1/1	0.94	0.10	82,82,82,82	0
56	MG	2A	3368	1/1	0.94	0.23	47,47,47,47	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3108	1/1	0.94	0.13	32,32,32,32	0
56	MG	1d	502	1/1	0.94	0.25	52,52,52,52	0
56	MG	1A	3939	1/1	0.94	0.13	48,48,48,48	0
56	MG	2A	3629	1/1	0.94	0.14	44,44,44,44	0
56	MG	1A	3277	1/1	0.94	0.25	31,31,31,31	0
56	MG	2A	3631	1/1	0.94	0.08	36,36,36,36	0
56	MG	1A	3679	1/1	0.94	0.09	42,42,42,42	0
56	MG	1A	3680	1/1	0.94	0.11	24,24,24,24	0
56	MG	2A	3638	1/1	0.94	0.13	34,34,34,34	0
56	MG	2q	3002	1/1	0.94	0.07	74,74,74,74	0
56	MG	2A	3639	1/1	0.94	0.12	38,38,38,38	0
56	MG	2r	101	1/1	0.94	0.31	70,70,70,70	0
56	MG	1A	4107	1/1	0.94	0.10	37,37,37,37	0
56	MG	1A	3189	1/1	0.94	0.15	34,34,34,34	0
56	MG	2U	203	1/1	0.94	0.30	63,63,63,63	0
56	MG	1n	502	1/1	0.94	0.18	55,55,55,55	0
56	MG	2A	3378	1/1	0.94	0.09	57,57,57,57	0
56	MG	2A	3646	1/1	0.94	0.07	55,55,55,55	0
56	MG	1a	1644	1/1	0.94	0.14	47,47,47,47	0
56	MG	1A	3812	1/1	0.94	0.24	54,54,54,54	0
56	MG	2X	101	1/1	0.94	0.09	46,46,46,46	0
56	MG	2w	3007	1/1	0.94	0.13	71,71,71,71	0
56	MG	2A	3650	1/1	0.94	0.14	42,42,42,42	0
56	MG	20	102	1/1	0.94	0.10	62,62,62,62	0
56	MG	2x	101	1/1	0.94	0.08	61,61,61,61	0
56	MG	2A	3652	1/1	0.94	0.11	53,53,53,53	0
56	MG	25	103	1/1	0.94	0.17	54,54,54,54	0
56	MG	25	104	1/1	0.94	0.08	52,52,52,52	0
56	MG	2A	3180	1/1	0.94	0.08	47,47,47,47	0
56	MG	2A	3654	1/1	0.94	0.12	52,52,52,52	0
56	MG	1A	3287	1/1	0.94	0.27	52,52,52,52	0
56	MG	1A	3684	1/1	0.94	0.07	14,14,14,14	0
56	MG	2A	3384	1/1	0.94	0.10	69,69,69,69	0
56	MG	1A	3332	1/1	0.94	0.06	39,39,39,39	0
56	MG	2A	3387	1/1	0.94	0.21	52,52,52,52	0
56	MG	1A	4124	1/1	0.94	0.29	39,39,39,39	0
57	MT9	1A	4113	33/33	0.94	0.10	25,34,43,56	0
56	MG	1A	3407	1/1	0.94	0.25	64,64,64,64	0
56	MG	2A	3193	1/1	0.94	0.07	62,62,62,62	0
56	MG	2A	3589	1/1	0.95	0.06	51,51,51,51	0
56	MG	1B	213	1/1	0.95	0.07	49,49,49,49	0
56	MG	2N	8001	1/1	0.95	0.07	52,52,52,52	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1681	1/1	0.95	0.07	51,51,51,51	0
56	MG	2Q	3001	1/1	0.95	0.09	53,53,53,53	0
56	MG	1a	1683	1/1	0.95	0.21	42,42,42,42	0
56	MG	1A	3595	1/1	0.95	0.12	51,51,51,51	0
56	MG	1A	3773	1/1	0.95	0.06	44,44,44,44	0
56	MG	2A	3597	1/1	0.95	0.14	54,54,54,54	0
56	MG	1A	3236	1/1	0.95	0.12	31,31,31,31	0
56	MG	1A	3481	1/1	0.95	0.11	46,46,46,46	0
56	MG	1A	3059	1/1	0.95	0.15	53,53,53,53	0
56	MG	1A	3943	1/1	0.95	0.10	53,53,53,53	0
56	MG	1B	223	1/1	0.95	0.15	62,62,62,62	0
56	MG	1A	3778	1/1	0.95	0.14	21,21,21,21	0
56	MG	2A	3316	1/1	0.95	0.07	52,52,52,52	0
56	MG	1A	3607	1/1	0.95	0.13	45,45,45,45	0
56	MG	1B	228	1/1	0.95	0.09	53,53,53,53	0
56	MG	1A	3391	1/1	0.95	0.18	32,32,32,32	0
56	MG	1A	3611	1/1	0.95	0.05	31,31,31,31	0
56	MG	2X	102	1/1	0.95	0.07	66,66,66,66	0
56	MG	2Y	502	1/1	0.95	0.13	49,49,49,49	0
56	MG	1A	3113	1/1	0.95	0.13	25,25,25,25	0
56	MG	20	101	1/1	0.95	0.09	48,48,48,48	0
56	MG	2A	3073	1/1	0.95	0.12	53,53,53,53	0
56	MG	23	102	1/1	0.95	0.10	52,52,52,52	0
56	MG	23	103	1/1	0.95	0.08	56,56,56,56	0
56	MG	1A	3786	1/1	0.95	0.09	23,23,23,23	0
56	MG	2A	3076	1/1	0.95	0.16	48,48,48,48	0
56	MG	1A	3789	1/1	0.95	0.41	36,36,36,36	0
56	MG	1D	302	1/1	0.95	0.09	51,51,51,51	0
56	MG	1A	3170	1/1	0.95	0.14	33,33,33,33	0
56	MG	27	101	1/1	0.95	0.16	48,48,48,48	0
56	MG	1a	1703	1/1	0.95	0.20	49,49,49,49	0
56	MG	1a	1704	1/1	0.95	0.12	50,50,50,50	0
56	MG	1A	3114	1/1	0.95	0.12	25,25,25,25	0
56	MG	1D	309	1/1	0.95	0.11	48,48,48,48	0
56	MG	1A	3960	1/1	0.95	0.09	67,67,67,67	0
56	MG	1A	3964	1/1	0.95	0.06	26,26,26,26	0
56	MG	1A	3792	1/1	0.95	0.07	32,32,32,32	0
56	MG	1A	3116	1/1	0.95	0.10	47,47,47,47	0
56	MG	1A	3490	1/1	0.95	0.12	44,44,44,44	0
56	MG	2A	3633	1/1	0.95	0.15	58,58,58,58	0
56	MG	1a	1713	1/1	0.95	0.16	60,60,60,60	0
56	MG	1E	309	1/1	0.95	0.11	51,51,51,51	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1715	1/1	0.95	0.10	40,40,40,40	0
56	MG	1A	3398	1/1	0.95	0.11	38,38,38,38	0
56	MG	1a	1718	1/1	0.95	0.12	48,48,48,48	0
56	MG	1E	312	1/1	0.95	0.06	34,34,34,34	0
56	MG	1A	3627	1/1	0.95	0.07	30,30,30,30	0
56	MG	1A	3036	1/1	0.95	0.14	25,25,25,25	0
56	MG	1A	3178	1/1	0.95	0.18	61,61,61,61	0
56	MG	1A	3976	1/1	0.95	0.08	32,32,32,32	0
56	MG	2A	3648	1/1	0.95	0.06	47,47,47,47	0
56	MG	1A	3496	1/1	0.95	0.09	40,40,40,40	0
56	MG	2A	3104	1/1	0.95	0.07	52,52,52,52	0
56	MG	2A	3651	1/1	0.95	0.11	41,41,41,41	0
56	MG	2A	3354	1/1	0.95	0.06	50,50,50,50	0
56	MG	1A	3119	1/1	0.95	0.16	39,39,39,39	0
56	MG	2A	3356	1/1	0.95	0.08	55,55,55,55	0
56	MG	1A	3404	1/1	0.95	0.09	49,49,49,49	0
56	MG	1A	3247	1/1	0.95	0.13	41,41,41,41	0
56	MG	1A	3127	1/1	0.95	0.07	61,61,61,61	0
56	MG	1N	205	1/1	0.95	0.17	46,46,46,46	0
56	MG	2A	3659	1/1	0.95	0.16	67,67,67,67	0
56	MG	1A	3190	1/1	0.95	0.30	30,30,30,30	0
56	MG	1A	3322	1/1	0.95	0.22	40,40,40,40	0
56	MG	1A	3505	1/1	0.95	0.07	39,39,39,39	0
56	MG	2A	3364	1/1	0.95	0.07	56,56,56,56	0
56	MG	1A	3506	1/1	0.95	0.11	43,43,43,43	0
56	MG	1A	3813	1/1	0.95	0.14	50,50,50,50	0
56	MG	1a	1747	1/1	0.95	0.06	51,51,51,51	0
56	MG	1O	206	1/1	0.95	0.08	54,54,54,54	0
56	MG	2A	3669	1/1	0.95	0.08	47,47,47,47	0
56	MG	1A	3814	1/1	0.95	0.08	48,48,48,48	0
56	MG	2A	3123	1/1	0.95	0.14	47,47,47,47	0
56	MG	1A	3251	1/1	0.95	0.10	40,40,40,40	0
56	MG	1A	3994	1/1	0.95	0.12	17,17,17,17	0
56	MG	1P	204	1/1	0.95	0.18	32,32,32,32	0
56	MG	2A	3677	1/1	0.95	0.27	48,48,48,48	0
56	MG	2A	3130	1/1	0.95	0.07	45,45,45,45	0
56	MG	2A	3679	1/1	0.95	0.06	43,43,43,43	0
56	MG	1Q	201	1/1	0.95	0.12	26,26,26,26	0
56	MG	1A	3816	1/1	0.95	0.09	36,36,36,36	0
56	MG	1A	3050	1/1	0.95	0.17	48,48,48,48	0
56	MG	2A	3135	1/1	0.95	0.11	39,39,39,39	0
56	MG	1A	3649	1/1	0.95	0.07	20,20,20,20	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3325	1/1	0.95	0.20	45,45,45,45	0
56	MG	2A	3139	1/1	0.95	0.17	46,46,46,46	0
56	MG	1S	3003	1/1	0.95	0.06	64,64,64,64	0
56	MG	2A	3688	1/1	0.95	0.08	48,48,48,48	0
56	MG	2A	3141	1/1	0.95	0.17	47,47,47,47	0
56	MG	2A	3142	1/1	0.95	0.17	59,59,59,59	0
56	MG	2A	3143	1/1	0.95	0.18	41,41,41,41	0
56	MG	2a	1662	1/1	0.95	0.07	81,81,81,81	0
56	MG	1A	3659	1/1	0.95	0.06	21,21,21,21	0
56	MG	2A	3145	1/1	0.95	0.10	52,52,52,52	0
56	MG	1A	3660	1/1	0.95	0.04	23,23,23,23	0
56	MG	1A	3255	1/1	0.95	0.21	63,63,63,63	0
56	MG	1A	3192	1/1	0.95	0.10	48,48,48,48	0
56	MG	2A	3698	1/1	0.95	0.07	69,69,69,69	0
56	MG	1A	3193	1/1	0.95	0.22	38,38,38,38	0
56	MG	2A	3701	1/1	0.95	0.08	48,48,48,48	0
56	MG	1a	1769	1/1	0.95	0.08	63,63,63,63	0
56	MG	1W	206	1/1	0.95	0.16	19,19,19,19	0
56	MG	1A	3516	1/1	0.95	0.23	34,34,34,34	0
56	MG	1X	104	1/1	0.95	0.19	39,39,39,39	0
56	MG	1a	1774	1/1	0.95	0.08	61,61,61,61	0
56	MG	1A	3828	1/1	0.95	0.07	43,43,43,43	0
56	MG	1A	3258	1/1	0.95	0.12	47,47,47,47	0
56	MG	2A	3403	1/1	0.95	0.23	55,55,55,55	0
56	MG	2A	3714	1/1	0.95	0.09	48,48,48,48	0
56	MG	1A	3418	1/1	0.95	0.15	47,47,47,47	0
56	MG	2A	3405	1/1	0.95	0.09	48,48,48,48	0
56	MG	1A	3831	1/1	0.95	0.09	54,54,54,54	0
56	MG	1A	3026	1/1	0.95	0.07	34,34,34,34	0
56	MG	2a	1688	1/1	0.95	0.11	71,71,71,71	0
56	MG	1Z	304	1/1	0.95	0.19	53,53,53,53	0
56	MG	1a	1788	1/1	0.95	0.12	65,65,65,65	0
56	MG	2A	3411	1/1	0.95	0.09	54,54,54,54	0
56	MG	10	101	1/1	0.95	0.07	44,44,44,44	0
56	MG	10	102	1/1	0.95	0.05	40,40,40,40	0
56	MG	1A	3198	1/1	0.95	0.08	39,39,39,39	0
56	MG	2a	1695	1/1	0.95	0.13	51,51,51,51	0
56	MG	1A	3334	1/1	0.95	0.16	43,43,43,43	0
56	MG	2A	3730	1/1	0.95	0.06	35,35,35,35	0
56	MG	1A	3336	1/1	0.95	0.10	39,39,39,39	0
56	MG	10	108	1/1	0.95	0.06	44,44,44,44	0
56	MG	1a	1805	1/1	0.95	0.15	59,59,59,59	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3179	1/1	0.95	0.13	35,35,35,35	0
56	MG	1A	3069	1/1	0.95	0.28	55,55,55,55	0
56	MG	1A	3340	1/1	0.95	0.15	38,38,38,38	0
56	MG	2A	3182	1/1	0.95	0.21	53,53,53,53	0
56	MG	2A	3183	1/1	0.95	0.09	53,53,53,53	0
56	MG	2A	3184	1/1	0.95	0.10	49,49,49,49	0
56	MG	2A	3742	1/1	0.95	0.13	53,53,53,53	0
56	MG	2A	3426	1/1	0.95	0.17	60,60,60,60	0
56	MG	2A	3744	1/1	0.95	0.07	65,65,65,65	0
56	MG	1A	3431	1/1	0.95	0.11	46,46,46,46	0
56	MG	1A	3843	1/1	0.95	0.11	35,35,35,35	0
56	MG	1A	4028	1/1	0.95	0.09	55,55,55,55	0
56	MG	1A	3101	1/1	0.95	0.10	45,45,45,45	0
56	MG	2A	3192	1/1	0.95	0.20	55,55,55,55	0
56	MG	1A	4035	1/1	0.95	0.08	48,48,48,48	0
56	MG	1A	3847	1/1	0.95	0.11	54,54,54,54	0
56	MG	2a	1717	1/1	0.95	0.13	58,58,58,58	0
56	MG	1a	1816	1/1	0.95	0.23	56,56,56,56	0
56	MG	1A	3102	1/1	0.95	0.09	33,33,33,33	0
56	MG	1A	3079	1/1	0.95	0.11	36,36,36,36	0
56	MG	1A	4039	1/1	0.95	0.07	48,48,48,48	0
56	MG	1A	3349	1/1	0.95	0.14	35,35,35,35	0
56	MG	2A	3440	1/1	0.95	0.20	40,40,40,40	0
56	MG	1A	3852	1/1	0.95	0.13	39,39,39,39	0
56	MG	2A	3759	1/1	0.95	0.30	39,39,39,39	0
56	MG	1a	1827	1/1	0.95	0.18	49,49,49,49	0
56	MG	1A	3853	1/1	0.95	0.05	28,28,28,28	0
56	MG	18	103	1/1	0.95	0.13	29,29,29,29	0
56	MG	1A	3854	1/1	0.95	0.11	25,25,25,25	0
56	MG	2A	3765	1/1	0.95	0.10	56,56,56,56	0
56	MG	1A	3688	1/1	0.95	0.08	44,44,44,44	0
56	MG	1A	3689	1/1	0.95	0.08	28,28,28,28	0
56	MG	1A	3146	1/1	0.95	0.13	32,32,32,32	0
56	MG	1A	3691	1/1	0.95	0.07	60,60,60,60	0
56	MG	2A	3770	1/1	0.95	0.16	55,55,55,55	0
56	MG	1a	1607	1/1	0.95	0.22	41,41,41,41	0
56	MG	2A	3210	1/1	0.95	0.07	62,62,62,62	0
56	MG	1l	201	1/1	0.95	0.15	38,38,38,38	0
56	MG	1A	3540	1/1	0.95	0.14	37,37,37,37	0
56	MG	1A	3541	1/1	0.95	0.18	47,47,47,47	0
56	MG	2A	3779	1/1	0.95	0.06	60,60,60,60	0
56	MG	1A	3104	1/1	0.95	0.11	29,29,29,29	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3443	1/1	0.95	0.18	35,35,35,35	0
56	MG	1A	4060	1/1	0.95	0.07	37,37,37,37	0
56	MG	1a	1615	1/1	0.95	0.12	60,60,60,60	0
56	MG	1a	1616	1/1	0.95	0.08	51,51,51,51	0
56	MG	1A	4062	1/1	0.95	0.07	21,21,21,21	0
56	MG	1a	1618	1/1	0.95	0.10	42,42,42,42	0
56	MG	2A	3223	1/1	0.95	0.14	54,54,54,54	0
56	MG	2A	3468	1/1	0.95	0.09	49,49,49,49	0
56	MG	2a	1765	1/1	0.95	0.05	80,80,80,80	0
56	MG	1A	3148	1/1	0.95	0.08	33,33,33,33	0
56	MG	2A	3470	1/1	0.95	0.11	51,51,51,51	0
56	MG	1A	3213	1/1	0.95	0.06	44,44,44,44	0
56	MG	1A	3040	1/1	0.95	0.27	33,33,33,33	0
56	MG	1w	104	1/1	0.95	0.17	52,52,52,52	0
56	MG	1A	3219	1/1	0.95	0.10	45,45,45,45	0
56	MG	2a	1773	1/1	0.95	0.07	76,76,76,76	0
56	MG	1A	3706	1/1	0.95	0.14	51,51,51,51	0
56	MG	2a	1776	1/1	0.95	0.08	77,77,77,77	0
56	MG	1A	3280	1/1	0.95	0.09	42,42,42,42	0
56	MG	1A	3450	1/1	0.95	0.13	46,46,46,46	0
56	MG	2A	3805	1/1	0.95	0.13	70,70,70,70	0
56	MG	1A	3365	1/1	0.95	0.19	39,39,39,39	0
56	MG	2a	1783	1/1	0.95	0.11	65,65,65,65	0
56	MG	2A	3807	1/1	0.95	0.10	65,65,65,65	0
56	MG	1a	1627	1/1	0.95	0.09	23,23,23,23	0
56	MG	1A	3282	1/1	0.95	0.19	45,45,45,45	0
56	MG	2a	1788	1/1	0.95	0.06	71,71,71,71	0
56	MG	1A	3220	1/1	0.95	0.10	45,45,45,45	0
56	MG	1A	3370	1/1	0.95	0.16	56,56,56,56	0
56	MG	1a	1632	1/1	0.95	0.17	40,40,40,40	0
56	MG	1A	3285	1/1	0.95	0.08	43,43,43,43	0
56	MG	1A	4088	1/1	0.95	0.10	36,36,36,36	0
56	MG	1A	3085	1/1	0.95	0.07	35,35,35,35	0
56	MG	1A	3086	1/1	0.95	0.22	53,53,53,53	0
56	MG	1A	3462	1/1	0.95	0.05	34,34,34,34	0
56	MG	1A	4097	1/1	0.95	0.15	39,39,39,39	0
56	MG	2A	3821	1/1	0.95	0.12	61,61,61,61	0
56	MG	2A	3822	1/1	0.95	0.09	82,82,82,82	0
56	MG	1A	3567	1/1	0.95	0.13	36,36,36,36	0
56	MG	2A	3825	1/1	0.95	0.06	55,55,55,55	0
56	MG	2A	3827	1/1	0.95	0.06	44,44,44,44	0
56	MG	2A	3495	1/1	0.95	0.20	73,73,73,73	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3375	1/1	0.95	0.17	39,39,39,39	0
56	MG	2A	3832	1/1	0.95	0.06	79,79,79,79	0
56	MG	2A	3498	1/1	0.95	0.07	62,62,62,62	0
56	MG	1a	1641	1/1	0.95	0.11	47,47,47,47	0
56	MG	2A	3250	1/1	0.95	0.07	58,58,58,58	0
56	MG	2A	3503	1/1	0.95	0.07	56,56,56,56	0
56	MG	1a	1643	1/1	0.95	0.15	47,47,47,47	0
56	MG	2A	3841	1/1	0.95	0.09	27,27,27,27	0
56	MG	2A	3843	1/1	0.95	0.07	52,52,52,52	0
56	MG	2a	1815	1/1	0.95	0.12	49,49,49,49	0
56	MG	2A	3846	1/1	0.95	0.06	61,61,61,61	0
56	MG	1A	3569	1/1	0.95	0.28	52,52,52,52	0
56	MG	2A	3851	1/1	0.95	0.11	66,66,66,66	0
56	MG	1A	3465	1/1	0.95	0.11	46,46,46,46	0
56	MG	1A	3224	1/1	0.95	0.06	39,39,39,39	0
56	MG	1a	1647	1/1	0.95	0.17	56,56,56,56	0
56	MG	2A	3510	1/1	0.95	0.11	53,53,53,53	0
56	MG	2A	3511	1/1	0.95	0.13	24,24,24,24	0
56	MG	1A	3579	1/1	0.95	0.14	43,43,43,43	0
56	MG	1A	3225	1/1	0.95	0.17	57,57,57,57	0
56	MG	2A	3003	1/1	0.95	0.18	40,40,40,40	0
56	MG	1A	3468	1/1	0.95	0.14	45,45,45,45	0
56	MG	2A	3519	1/1	0.95	0.08	46,46,46,46	0
56	MG	2A	3005	1/1	0.95	0.29	58,58,58,58	0
56	MG	1A	3379	1/1	0.95	0.16	26,26,26,26	0
56	MG	1A	3022	1/1	0.95	0.14	44,44,44,44	0
56	MG	2A	3523	1/1	0.95	0.08	37,37,37,37	0
56	MG	2A	3264	1/1	0.95	0.15	64,64,64,64	0
56	MG	2A	3010	1/1	0.95	0.22	57,57,57,57	0
56	MG	1A	3033	1/1	0.95	0.12	34,34,34,34	0
56	MG	1A	4115	1/1	0.95	0.07	33,33,33,33	0
56	MG	1A	4116	1/1	0.95	0.14	30,30,30,30	0
56	MG	1A	4117	1/1	0.95	0.20	38,38,38,38	0
56	MG	1A	3228	1/1	0.95	0.32	44,44,44,44	0
56	MG	2A	3019	1/1	0.95	0.07	37,37,37,37	0
56	MG	2A	3889	1/1	0.95	0.22	39,39,39,39	0
56	MG	2A	3539	1/1	0.95	0.11	60,60,60,60	0
56	MG	2A	3540	1/1	0.95	0.09	60,60,60,60	0
56	MG	1A	4119	1/1	0.95	0.13	34,34,34,34	0
56	MG	1A	4120	1/1	0.95	0.12	27,27,27,27	0
56	MG	2A	3024	1/1	0.95	0.12	47,47,47,47	0
56	MG	2f	3002	1/1	0.95	0.12	62,62,62,62	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3277	1/1	0.95	0.13	51,51,51,51	0
56	MG	2A	3278	1/1	0.95	0.09	42,42,42,42	0
56	MG	2A	3900	1/1	0.95	0.20	44,44,44,44	0
56	MG	2I	3001	1/1	0.95	0.13	60,60,60,60	0
56	MG	2A	3279	1/1	0.95	0.10	64,64,64,64	0
56	MG	2A	3904	1/1	0.95	0.09	40,40,40,40	0
56	MG	1A	3474	1/1	0.95	0.10	48,48,48,48	0
56	MG	2B	3001	1/1	0.95	0.14	57,57,57,57	0
56	MG	2A	3550	1/1	0.95	0.08	35,35,35,35	0
56	MG	1A	3587	1/1	0.95	0.10	27,27,27,27	0
56	MG	1A	4133	1/1	0.95	0.10	30,30,30,30	0
56	MG	1A	3755	1/1	0.95	0.09	20,20,20,20	0
56	MG	2A	3555	1/1	0.95	0.10	34,34,34,34	0
56	MG	1A	4136	1/1	0.95	0.28	39,39,39,39	0
56	MG	1A	4139	1/1	0.95	0.09	33,33,33,33	0
56	MG	1A	4140	1/1	0.95	0.19	37,37,37,37	0
56	MG	2A	3035	1/1	0.95	0.09	44,44,44,44	0
56	MG	1A	4146	1/1	0.95	0.20	38,38,38,38	0
56	MG	2A	3289	1/1	0.95	0.05	46,46,46,46	0
56	MG	2A	3039	1/1	0.95	0.13	33,33,33,33	0
56	MG	2A	3570	1/1	0.95	0.08	53,53,53,53	0
56	MG	2A	3291	1/1	0.95	0.08	58,58,58,58	0
56	MG	2A	3041	1/1	0.95	0.12	52,52,52,52	0
56	MG	1A	3589	1/1	0.95	0.06	20,20,20,20	0
56	MG	1B	203	1/1	0.95	0.18	55,55,55,55	0
56	MG	1B	205	1/1	0.95	0.26	60,60,60,60	0
56	MG	1A	3232	1/1	0.95	0.12	51,51,51,51	0
56	MG	2D	303	1/1	0.95	0.32	40,40,40,40	0
56	MG	1A	3161	1/1	0.95	0.15	33,33,33,33	0
56	MG	2A	3300	1/1	0.95	0.10	66,66,66,66	0
56	MG	1A	3765	1/1	0.95	0.07	16,16,16,16	0
56	MG	2A	3582	1/1	0.95	0.06	61,61,61,61	0
56	MG	1B	209	1/1	0.95	0.09	48,48,48,48	0
56	MG	2A	3584	1/1	0.95	0.10	62,62,62,62	0
56	MG	2E	307	1/1	0.95	0.09	36,36,36,36	0
56	MG	1A	3478	1/1	0.95	0.09	43,43,43,43	0
56	MG	1A	3593	1/1	0.95	0.07	33,33,33,33	0
56	MG	2A	3348	1/1	0.96	0.08	44,44,44,44	0
56	MG	1A	3492	1/1	0.96	0.08	45,45,45,45	0
56	MG	1F	307	1/1	0.96	0.06	44,44,44,44	0
56	MG	2A	3637	1/1	0.96	0.05	39,39,39,39	0
56	MG	1a	1720	1/1	0.96	0.06	54,54,54,54	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3400	1/1	0.96	0.06	36,36,36,36	0
56	MG	2A	3640	1/1	0.96	0.24	52,52,52,52	0
56	MG	2A	3641	1/1	0.96	0.11	60,60,60,60	0
56	MG	1A	3494	1/1	0.96	0.07	51,51,51,51	0
56	MG	1A	3622	1/1	0.96	0.10	14,14,14,14	0
56	MG	1A	3207	1/1	0.96	0.23	35,35,35,35	0
56	MG	1a	1726	1/1	0.96	0.13	46,46,46,46	0
56	MG	1A	3043	1/1	0.96	0.06	26,26,26,26	0
56	MG	23	101	1/1	0.96	0.10	62,62,62,62	0
56	MG	1H	3001	1/1	0.96	0.17	37,37,37,37	0
56	MG	1A	3797	1/1	0.96	0.05	28,28,28,28	0
56	MG	1N	203	1/1	0.96	0.07	42,42,42,42	0
56	MG	1A	3326	1/1	0.96	0.19	46,46,46,46	0
56	MG	2A	3108	1/1	0.96	0.09	42,42,42,42	0
56	MG	1A	3045	1/1	0.96	0.08	31,31,31,31	0
56	MG	1N	206	1/1	0.96	0.11	31,31,31,31	0
56	MG	1A	3016	1/1	0.96	0.12	62,62,62,62	0
56	MG	1A	3987	1/1	0.96	0.10	40,40,40,40	0
56	MG	1a	1742	1/1	0.96	0.12	53,53,53,53	0
56	MG	2a	1602	1/1	0.96	0.23	53,53,53,53	0
56	MG	1A	3329	1/1	0.96	0.09	42,42,42,42	0
56	MG	1A	3071	1/1	0.96	0.06	31,31,31,31	0
56	MG	1A	3990	1/1	0.96	0.07	39,39,39,39	0
56	MG	1a	1746	1/1	0.96	0.08	58,58,58,58	0
56	MG	1A	3633	1/1	0.96	0.08	45,45,45,45	0
56	MG	1A	3073	1/1	0.96	0.08	33,33,33,33	0
56	MG	1A	3218	1/1	0.96	0.09	47,47,47,47	0
56	MG	1P	201	1/1	0.96	0.18	26,26,26,26	0
56	MG	1P	202	1/1	0.96	0.23	28,28,28,28	0
56	MG	1A	3152	1/1	0.96	0.11	30,30,30,30	0
56	MG	1A	3412	1/1	0.96	0.09	36,36,36,36	0
56	MG	2A	3668	1/1	0.96	0.06	50,50,50,50	0
56	MG	1A	3998	1/1	0.96	0.06	68,68,68,68	0
56	MG	1Q	203	1/1	0.96	0.14	51,51,51,51	0
56	MG	1A	3074	1/1	0.96	0.12	25,25,25,25	0
56	MG	1A	3641	1/1	0.96	0.05	29,29,29,29	0
56	MG	1A	3509	1/1	0.96	0.13	38,38,38,38	0
56	MG	1A	3643	1/1	0.96	0.06	21,21,21,21	0
56	MG	2A	3676	1/1	0.96	0.07	52,52,52,52	0
56	MG	2A	3386	1/1	0.96	0.12	67,67,67,67	0
56	MG	1A	3335	1/1	0.96	0.06	43,43,43,43	0
56	MG	1a	1763	1/1	0.96	0.07	75,75,75,75	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3157	1/1	0.96	0.39	33,33,33,33	0
56	MG	1a	1765	1/1	0.96	0.06	69,69,69,69	0
56	MG	1A	3416	1/1	0.96	0.19	51,51,51,51	0
56	MG	1A	3337	1/1	0.96	0.08	38,38,38,38	0
56	MG	1a	1768	1/1	0.96	0.07	71,71,71,71	0
56	MG	1A	3651	1/1	0.96	0.07	24,24,24,24	0
56	MG	1A	3652	1/1	0.96	0.06	37,37,37,37	0
56	MG	2A	3149	1/1	0.96	0.15	34,34,34,34	0
56	MG	1W	204	1/1	0.96	0.11	30,30,30,30	0
56	MG	1A	3654	1/1	0.96	0.05	20,20,20,20	0
56	MG	1A	3515	1/1	0.96	0.11	28,28,28,28	0
56	MG	1a	1775	1/1	0.96	0.11	68,68,68,68	0
56	MG	1A	3075	1/1	0.96	0.08	47,47,47,47	0
56	MG	2A	3155	1/1	0.96	0.12	63,63,63,63	0
56	MG	1A	3518	1/1	0.96	0.17	46,46,46,46	0
56	MG	1A	3339	1/1	0.96	0.29	47,47,47,47	0
56	MG	2A	3159	1/1	0.96	0.18	37,37,37,37	0
56	MG	2A	3160	1/1	0.96	0.12	54,54,54,54	0
56	MG	1A	3663	1/1	0.96	0.07	30,30,30,30	0
56	MG	1A	3047	1/1	0.96	0.17	44,44,44,44	0
56	MG	1a	1782	1/1	0.96	0.08	66,66,66,66	0
56	MG	2A	3703	1/1	0.96	0.06	61,61,61,61	0
56	MG	1a	1783	1/1	0.96	0.05	58,58,58,58	0
56	MG	1Z	301	1/1	0.96	0.08	44,44,44,44	0
56	MG	1a	1786	1/1	0.96	0.06	67,67,67,67	0
56	MG	2A	3708	1/1	0.96	0.10	63,63,63,63	0
56	MG	1A	3341	1/1	0.96	0.10	50,50,50,50	0
56	MG	2A	3170	1/1	0.96	0.12	54,54,54,54	0
56	MG	2A	3711	1/1	0.96	0.12	52,52,52,52	0
56	MG	1A	3425	1/1	0.96	0.10	46,46,46,46	0
56	MG	2A	3419	1/1	0.96	0.23	59,59,59,59	0
56	MG	1A	3342	1/1	0.96	0.17	44,44,44,44	0
56	MG	1A	3344	1/1	0.96	0.30	36,36,36,36	0
56	MG	2A	3717	1/1	0.96	0.07	40,40,40,40	0
56	MG	10	104	1/1	0.96	0.16	41,41,41,41	0
56	MG	1a	1798	1/1	0.96	0.08	69,69,69,69	0
56	MG	1A	4030	1/1	0.96	0.06	45,45,45,45	0
56	MG	1A	4031	1/1	0.96	0.11	52,52,52,52	0
56	MG	1A	3834	1/1	0.96	0.12	40,40,40,40	0
56	MG	2a	1664	1/1	0.96	0.14	57,57,57,57	0
56	MG	1A	3279	1/1	0.96	0.23	25,25,25,25	0
56	MG	1A	3430	1/1	0.96	0.07	41,41,41,41	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3160	1/1	0.96	0.19	32,32,32,32	0
56	MG	1A	3281	1/1	0.96	0.09	46,46,46,46	0
56	MG	1A	3030	1/1	0.96	0.27	26,26,26,26	0
56	MG	12	102	1/1	0.96	0.12	37,37,37,37	0
56	MG	1a	1812	1/1	0.96	0.10	54,54,54,54	0
56	MG	2A	3434	1/1	0.96	0.13	40,40,40,40	0
56	MG	1A	4040	1/1	0.96	0.10	25,25,25,25	0
56	MG	1a	1814	1/1	0.96	0.06	55,55,55,55	0
56	MG	1A	4041	1/1	0.96	0.07	33,33,33,33	0
56	MG	1A	3350	1/1	0.96	0.06	53,53,53,53	0
56	MG	16	101	1/1	0.96	0.07	42,42,42,42	0
56	MG	2A	3739	1/1	0.96	0.07	52,52,52,52	0
56	MG	1A	3004	1/1	0.96	0.08	47,47,47,47	0
56	MG	1A	3682	1/1	0.96	0.07	25,25,25,25	0
56	MG	1a	1824	1/1	0.96	0.18	36,36,36,36	0
56	MG	1A	3005	1/1	0.96	0.13	58,58,58,58	0
56	MG	1A	3441	1/1	0.96	0.25	42,42,42,42	0
56	MG	2a	1686	1/1	0.96	0.10	62,62,62,62	0
56	MG	1A	3286	1/1	0.96	0.17	52,52,52,52	0
56	MG	1A	3542	1/1	0.96	0.15	22,22,22,22	0
56	MG	1A	3229	1/1	0.96	0.09	54,54,54,54	0
56	MG	1A	3444	1/1	0.96	0.23	36,36,36,36	0
56	MG	19	101	1/1	0.96	0.06	44,44,44,44	0
56	MG	1A	3358	1/1	0.96	0.05	42,42,42,42	0
56	MG	1A	3855	1/1	0.96	0.15	40,40,40,40	0
56	MG	1A	3359	1/1	0.96	0.12	49,49,49,49	0
56	MG	1f	3002	1/1	0.96	0.11	45,45,45,45	0
56	MG	1A	3168	1/1	0.96	0.34	32,32,32,32	0
56	MG	2A	3211	1/1	0.96	0.05	37,37,37,37	0
56	MG	1A	3693	1/1	0.96	0.06	47,47,47,47	0
56	MG	1l	202	1/1	0.96	0.11	69,69,69,69	0
56	MG	2A	3462	1/1	0.96	0.22	47,47,47,47	0
56	MG	1a	1606	1/1	0.96	0.20	62,62,62,62	0
56	MG	1A	3291	1/1	0.96	0.15	41,41,41,41	0
56	MG	2A	3761	1/1	0.96	0.10	50,50,50,50	0
56	MG	1A	3861	1/1	0.96	0.08	53,53,53,53	0
56	MG	1a	1609	1/1	0.96	0.09	27,27,27,27	0
56	MG	1A	3115	1/1	0.96	0.14	31,31,31,31	0
56	MG	1A	4068	1/1	0.96	0.07	22,22,22,22	0
56	MG	1A	3865	1/1	0.96	0.06	25,25,25,25	0
56	MG	1A	3364	1/1	0.96	0.12	42,42,42,42	0
56	MG	1A	3035	1/1	0.96	0.17	32,32,32,32	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3010	1/1	0.96	0.13	33,33,33,33	0
56	MG	2A	3225	1/1	0.96	0.23	51,51,51,51	0
56	MG	2A	3474	1/1	0.96	0.17	46,46,46,46	0
56	MG	2A	3475	1/1	0.96	0.17	46,46,46,46	0
56	MG	2A	3774	1/1	0.96	0.07	36,36,36,36	0
56	MG	1A	3870	1/1	0.96	0.09	46,46,46,46	0
56	MG	1A	4075	1/1	0.96	0.07	47,47,47,47	0
56	MG	1A	3557	1/1	0.96	0.36	33,33,33,33	0
56	MG	1A	3701	1/1	0.96	0.09	48,48,48,48	0
56	MG	1A	4082	1/1	0.96	0.11	27,27,27,27	0
56	MG	1A	4083	1/1	0.96	0.10	51,51,51,51	0
56	MG	1w	108	1/1	0.96	0.06	36,36,36,36	0
56	MG	1A	3558	1/1	0.96	0.09	23,23,23,23	0
56	MG	1A	3874	1/1	0.96	0.06	19,19,19,19	0
56	MG	1A	4087	1/1	0.96	0.16	46,46,46,46	0
56	MG	1A	3367	1/1	0.96	0.12	43,43,43,43	0
56	MG	1A	4090	1/1	0.96	0.09	48,48,48,48	0
56	MG	2A	3238	1/1	0.96	0.18	42,42,42,42	0
56	MG	2A	3239	1/1	0.96	0.20	39,39,39,39	0
56	MG	1A	3560	1/1	0.96	0.11	43,43,43,43	0
56	MG	1A	3368	1/1	0.96	0.06	36,36,36,36	0
56	MG	1x	105	1/1	0.96	0.19	53,53,53,53	0
56	MG	2A	3494	1/1	0.96	0.21	59,59,59,59	0
56	MG	2a	1735	1/1	0.96	0.11	75,75,75,75	0
56	MG	2a	1736	1/1	0.96	0.06	76,76,76,76	0
56	MG	2A	3799	1/1	0.96	0.06	54,54,54,54	0
56	MG	1a	1630	1/1	0.96	0.13	44,44,44,44	0
56	MG	1A	3118	1/1	0.96	0.14	27,27,27,27	0
56	MG	1A	3711	1/1	0.96	0.06	37,37,37,37	0
56	MG	1A	3883	1/1	0.96	0.08	47,47,47,47	0
56	MG	1x	110	1/1	0.96	0.05	53,53,53,53	0
56	MG	2A	3502	1/1	0.96	0.07	43,43,43,43	0
56	MG	2a	1746	1/1	0.96	0.11	70,70,70,70	0
56	MG	1A	3056	1/1	0.96	0.15	47,47,47,47	0
56	MG	1A	3885	1/1	0.96	0.09	40,40,40,40	0
56	MG	1A	3184	1/1	0.96	0.08	25,25,25,25	0
56	MG	1A	3120	1/1	0.96	0.05	28,28,28,28	0
56	MG	1A	3718	1/1	0.96	0.19	38,38,38,38	0
56	MG	1A	3719	1/1	0.96	0.05	42,42,42,42	0
56	MG	1A	4108	1/1	0.96	0.07	40,40,40,40	0
56	MG	1A	3304	1/1	0.96	0.05	29,29,29,29	0
56	MG	2a	1759	1/1	0.96	0.05	83,83,83,83	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3893	1/1	0.96	0.09	44,44,44,44	0
56	MG	2a	1762	1/1	0.96	0.06	51,51,51,51	0
56	MG	1A	3464	1/1	0.96	0.20	34,34,34,34	0
56	MG	2A	3516	1/1	0.96	0.15	52,52,52,52	0
56	MG	1A	3305	1/1	0.96	0.20	51,51,51,51	0
56	MG	1A	3570	1/1	0.96	0.09	27,27,27,27	0
56	MG	1A	3726	1/1	0.96	0.07	42,42,42,42	0
56	MG	2A	3823	1/1	0.96	0.08	48,48,48,48	0
56	MG	1A	3187	1/1	0.96	0.20	25,25,25,25	0
56	MG	1a	1649	1/1	0.96	0.14	62,62,62,62	0
56	MG	1A	3572	1/1	0.96	0.09	15,15,15,15	0
56	MG	2A	3828	1/1	0.96	0.09	59,59,59,59	0
56	MG	2A	3007	1/1	0.96	0.14	53,53,53,53	0
56	MG	2A	3524	1/1	0.96	0.06	46,46,46,46	0
56	MG	1A	3125	1/1	0.96	0.12	45,45,45,45	0
56	MG	1A	3575	1/1	0.96	0.10	34,34,34,34	0
56	MG	2A	3834	1/1	0.96	0.06	42,42,42,42	0
56	MG	2a	1782	1/1	0.96	0.07	54,54,54,54	0
56	MG	1A	3732	1/1	0.96	0.04	26,26,26,26	0
56	MG	2A	3531	1/1	0.96	0.07	57,57,57,57	0
56	MG	1A	3577	1/1	0.96	0.09	29,29,29,29	0
56	MG	2A	3839	1/1	0.96	0.05	71,71,71,71	0
56	MG	2A	3270	1/1	0.96	0.17	63,63,63,63	0
56	MG	1A	4129	1/1	0.96	0.10	33,33,33,33	0
56	MG	2A	3015	1/1	0.96	0.11	48,48,48,48	0
56	MG	2A	3844	1/1	0.96	0.06	39,39,39,39	0
56	MG	1A	4130	1/1	0.96	0.23	36,36,36,36	0
56	MG	2A	3849	1/1	0.96	0.06	51,51,51,51	0
56	MG	2A	3017	1/1	0.96	0.09	46,46,46,46	0
56	MG	1A	3737	1/1	0.96	0.06	34,34,34,34	0
56	MG	1A	3910	1/1	0.96	0.07	71,71,71,71	0
56	MG	2A	3020	1/1	0.96	0.09	63,63,63,63	0
56	MG	1a	1659	1/1	0.96	0.19	64,64,64,64	0
56	MG	1A	3911	1/1	0.96	0.12	72,72,72,72	0
56	MG	1A	3738	1/1	0.96	0.09	45,45,45,45	0
56	MG	2A	3863	1/1	0.96	0.09	49,49,49,49	0
56	MG	2A	3864	1/1	0.96	0.06	59,59,59,59	0
56	MG	2A	3866	1/1	0.96	0.07	34,34,34,34	0
56	MG	2a	1805	1/1	0.96	0.09	51,51,51,51	0
56	MG	1A	3739	1/1	0.96	0.08	44,44,44,44	0
56	MG	2A	3026	1/1	0.96	0.18	48,48,48,48	0
56	MG	1A	4141	1/1	0.96	0.06	30,30,30,30	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3028	1/1	0.96	0.07	39,39,39,39	0
56	MG	2A	3029	1/1	0.96	0.20	56,56,56,56	0
56	MG	1A	3740	1/1	0.96	0.13	44,44,44,44	0
56	MG	1a	1665	1/1	0.96	0.10	61,61,61,61	0
56	MG	1A	4147	1/1	0.96	0.28	40,40,40,40	0
56	MG	1A	3012	1/1	0.96	0.22	29,29,29,29	0
56	MG	1A	3128	1/1	0.96	0.13	37,37,37,37	0
56	MG	1A	3923	1/1	0.96	0.05	23,23,23,23	0
56	MG	2A	3565	1/1	0.96	0.06	45,45,45,45	0
56	MG	2a	1818	1/1	0.96	0.12	74,74,74,74	0
56	MG	2A	3882	1/1	0.96	0.09	49,49,49,49	0
56	MG	2A	3036	1/1	0.96	0.15	35,35,35,35	0
56	MG	1A	3381	1/1	0.96	0.11	48,48,48,48	0
56	MG	2A	3296	1/1	0.96	0.07	42,42,42,42	0
56	MG	2A	3569	1/1	0.96	0.13	61,61,61,61	0
56	MG	2A	3887	1/1	0.96	0.15	51,51,51,51	0
56	MG	1A	3382	1/1	0.96	0.06	35,35,35,35	0
56	MG	2A	3040	1/1	0.96	0.18	50,50,50,50	0
56	MG	1A	3472	1/1	0.96	0.08	48,48,48,48	0
56	MG	1A	3383	1/1	0.96	0.09	37,37,37,37	0
56	MG	1A	3095	1/1	0.96	0.06	42,42,42,42	0
56	MG	2A	3893	1/1	0.96	0.06	49,49,49,49	0
56	MG	2A	3575	1/1	0.96	0.07	38,38,38,38	0
56	MG	2A	3576	1/1	0.96	0.13	41,41,41,41	0
56	MG	1A	3133	1/1	0.96	0.17	26,26,26,26	0
56	MG	1A	3386	1/1	0.96	0.21	38,38,38,38	0
56	MG	1A	3312	1/1	0.96	0.13	45,45,45,45	0
56	MG	1B	217	1/1	0.96	0.16	56,56,56,56	0
56	MG	2A	3903	1/1	0.96	0.17	45,45,45,45	0
56	MG	1A	3757	1/1	0.96	0.06	30,30,30,30	0
56	MG	1A	3938	1/1	0.96	0.07	41,41,41,41	0
56	MG	1A	3479	1/1	0.96	0.16	40,40,40,40	0
56	MG	1A	3941	1/1	0.96	0.06	43,43,43,43	0
56	MG	1A	3134	1/1	0.96	0.17	35,35,35,35	0
56	MG	2A	3586	1/1	0.96	0.07	49,49,49,49	0
56	MG	1A	3252	1/1	0.96	0.09	34,34,34,34	0
56	MG	1a	1686	1/1	0.96	0.13	48,48,48,48	0
56	MG	1A	3316	1/1	0.96	0.14	34,34,34,34	0
56	MG	1A	3766	1/1	0.96	0.05	66,66,66,66	0
56	MG	1A	3594	1/1	0.96	0.10	45,45,45,45	0
56	MG	2A	3060	1/1	0.96	0.14	41,41,41,41	0
56	MG	1B	229	1/1	0.96	0.07	69,69,69,69	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3253	1/1	0.96	0.19	52,52,52,52	0
56	MG	1A	3771	1/1	0.96	0.08	18,18,18,18	0
56	MG	1A	3952	1/1	0.96	0.07	40,40,40,40	0
56	MG	2B	3015	1/1	0.96	0.20	59,59,59,59	0
56	MG	1B	234	1/1	0.96	0.10	42,42,42,42	0
56	MG	1a	1695	1/1	0.96	0.07	49,49,49,49	0
56	MG	1A	3953	1/1	0.96	0.07	46,46,46,46	0
56	MG	2A	3070	1/1	0.96	0.15	60,60,60,60	0
56	MG	2B	3020	1/1	0.96	0.26	78,78,78,78	0
56	MG	2A	3326	1/1	0.96	0.14	46,46,46,46	0
56	MG	1A	3598	1/1	0.96	0.12	12,12,12,12	0
56	MG	2D	302	1/1	0.96	0.11	33,33,33,33	0
56	MG	1A	3484	1/1	0.96	0.06	46,46,46,46	0
56	MG	2A	3329	1/1	0.96	0.09	59,59,59,59	0
56	MG	1A	3318	1/1	0.96	0.09	54,54,54,54	0
56	MG	1A	3038	1/1	0.96	0.07	25,25,25,25	0
56	MG	1A	3015	1/1	0.96	0.29	47,47,47,47	0
56	MG	1A	3608	1/1	0.96	0.04	17,17,17,17	0
56	MG	1A	3099	1/1	0.96	0.09	30,30,30,30	0
56	MG	2A	3616	1/1	0.96	0.07	34,34,34,34	0
56	MG	1A	3967	1/1	0.96	0.14	59,59,59,59	0
56	MG	1A	3060	1/1	0.96	0.10	42,42,42,42	0
56	MG	1a	1706	1/1	0.96	0.30	52,52,52,52	0
56	MG	1A	3041	1/1	0.96	0.12	35,35,35,35	0
56	MG	1A	3616	1/1	0.96	0.08	26,26,26,26	0
56	MG	1A	3784	1/1	0.96	0.07	43,43,43,43	0
56	MG	2A	3624	1/1	0.96	0.08	33,33,33,33	0
56	MG	1E	311	1/1	0.96	0.06	49,49,49,49	0
56	MG	2Q	3003	1/1	0.96	0.07	57,57,57,57	0
56	MG	1A	3973	1/1	0.96	0.11	51,51,51,51	0
56	MG	1A	3617	1/1	0.96	0.07	18,18,18,18	0
56	MG	1F	302	1/1	0.96	0.06	31,31,31,31	0
56	MG	1F	303	1/1	0.96	0.14	24,24,24,24	0
56	MG	1F	305	1/1	0.96	0.07	45,45,45,45	0
58	K	2A	3896	1/1	0.96	0.07	56,56,56,56	0
56	MG	2A	3632	1/1	0.96	0.06	47,47,47,47	0
56	MG	1Z	302	1/1	0.97	0.06	40,40,40,40	0
56	MG	1A	3904	1/1	0.97	0.05	81,81,81,81	0
56	MG	1A	4073	1/1	0.97	0.10	40,40,40,40	0
56	MG	1A	3439	1/1	0.97	0.18	29,29,29,29	0
56	MG	1A	3769	1/1	0.97	0.04	42,42,42,42	0
56	MG	2A	3084	1/1	0.97	0.08	54,54,54,54	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3135	1/1	0.97	0.05	36,36,36,36	0
56	MG	1A	4079	1/1	0.97	0.10	10,10,10,10	0
56	MG	1A	3632	1/1	0.97	0.09	33,33,33,33	0
56	MG	1A	3082	1/1	0.97	0.11	29,29,29,29	0
56	MG	1A	3188	1/1	0.97	0.21	32,32,32,32	0
56	MG	1A	3912	1/1	0.97	0.08	48,48,48,48	0
56	MG	11	101	1/1	0.97	0.10	31,31,31,31	0
56	MG	2A	3092	1/1	0.97	0.07	39,39,39,39	0
56	MG	1A	3635	1/1	0.97	0.05	50,50,50,50	0
56	MG	1A	3636	1/1	0.97	0.10	28,28,28,28	0
56	MG	2A	3095	1/1	0.97	0.07	40,40,40,40	0
56	MG	1A	3915	1/1	0.97	0.06	41,41,41,41	0
56	MG	2A	3530	1/1	0.97	0.04	33,33,33,33	0
56	MG	12	101	1/1	0.97	0.06	50,50,50,50	0
56	MG	1A	3243	1/1	0.97	0.12	35,35,35,35	0
56	MG	1A	4091	1/1	0.97	0.09	46,46,46,46	0
56	MG	13	102	1/1	0.97	0.07	38,38,38,38	0
56	MG	1A	3105	1/1	0.97	0.25	33,33,33,33	0
56	MG	1A	3009	1/1	0.97	0.09	30,30,30,30	0
56	MG	2A	3537	1/1	0.97	0.15	51,51,51,51	0
56	MG	2A	3538	1/1	0.97	0.06	47,47,47,47	0
56	MG	1A	4095	1/1	0.97	0.08	31,31,31,31	0
56	MG	1A	3084	1/1	0.97	0.06	44,44,44,44	0
56	MG	2A	3783	1/1	0.97	0.05	60,60,60,60	0
56	MG	1A	3781	1/1	0.97	0.06	36,36,36,36	0
56	MG	1A	3529	1/1	0.97	0.20	45,45,45,45	0
56	MG	2A	3318	1/1	0.97	0.10	44,44,44,44	0
56	MG	1A	3926	1/1	0.97	0.05	31,31,31,31	0
56	MG	2A	3790	1/1	0.97	0.07	86,86,86,86	0
56	MG	1A	3530	1/1	0.97	0.05	45,45,45,45	0
56	MG	2A	3792	1/1	0.97	0.07	55,55,55,55	0
56	MG	1A	3006	1/1	0.97	0.06	24,24,24,24	0
56	MG	2A	3548	1/1	0.97	0.07	32,32,32,32	0
56	MG	1A	3785	1/1	0.97	0.04	23,23,23,23	0
56	MG	2A	3111	1/1	0.97	0.17	40,40,40,40	0
56	MG	1A	3644	1/1	0.97	0.06	17,17,17,17	0
56	MG	2A	3113	1/1	0.97	0.13	65,65,65,65	0
56	MG	1A	3532	1/1	0.97	0.09	29,29,29,29	0
56	MG	1A	3374	1/1	0.97	0.05	32,32,32,32	0
56	MG	1a	1789	1/1	0.97	0.06	69,69,69,69	0
56	MG	2A	3558	1/1	0.97	0.06	30,30,30,30	0
56	MG	1a	1792	1/1	0.97	0.06	49,49,49,49	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3936	1/1	0.97	0.04	50,50,50,50	0
56	MG	2A	3561	1/1	0.97	0.06	52,52,52,52	0
56	MG	1A	3007	1/1	0.97	0.05	18,18,18,18	0
56	MG	2A	3564	1/1	0.97	0.08	36,36,36,36	0
56	MG	2A	3120	1/1	0.97	0.08	35,35,35,35	0
56	MG	1A	3194	1/1	0.97	0.07	38,38,38,38	0
56	MG	1A	3793	1/1	0.97	0.07	40,40,40,40	0
56	MG	2A	3335	1/1	0.97	0.06	55,55,55,55	0
56	MG	1A	4114	1/1	0.97	0.20	28,28,28,28	0
56	MG	1a	1802	1/1	0.97	0.07	54,54,54,54	0
56	MG	1A	3650	1/1	0.97	0.04	18,18,18,18	0
56	MG	2A	3128	1/1	0.97	0.12	47,47,47,47	0
56	MG	2A	3129	1/1	0.97	0.11	62,62,62,62	0
56	MG	2A	3819	1/1	0.97	0.05	57,57,57,57	0
56	MG	1A	3537	1/1	0.97	0.14	38,38,38,38	0
56	MG	1A	3314	1/1	0.97	0.07	39,39,39,39	0
56	MG	1A	3944	1/1	0.97	0.04	40,40,40,40	0
56	MG	1A	3378	1/1	0.97	0.16	36,36,36,36	0
56	MG	1A	3453	1/1	0.97	0.07	39,39,39,39	0
56	MG	1a	1613	1/1	0.97	0.08	47,47,47,47	0
56	MG	1A	4121	1/1	0.97	0.12	29,29,29,29	0
56	MG	2A	3138	1/1	0.97	0.18	43,43,43,43	0
56	MG	1A	3018	1/1	0.97	0.20	37,37,37,37	0
56	MG	1A	3027	1/1	0.97	0.11	32,32,32,32	0
56	MG	1A	4126	1/1	0.97	0.18	30,30,30,30	0
56	MG	1A	4127	1/1	0.97	0.11	29,29,29,29	0
56	MG	1A	3544	1/1	0.97	0.26	40,40,40,40	0
56	MG	1A	3803	1/1	0.97	0.05	25,25,25,25	0
56	MG	1A	4131	1/1	0.97	0.16	31,31,31,31	0
56	MG	2A	3837	1/1	0.97	0.05	33,33,33,33	0
56	MG	1a	1818	1/1	0.97	0.06	69,69,69,69	0
56	MG	1A	3545	1/1	0.97	0.11	22,22,22,22	0
56	MG	1A	3456	1/1	0.97	0.07	33,33,33,33	0
56	MG	1a	1822	1/1	0.97	0.06	58,58,58,58	0
56	MG	1A	3458	1/1	0.97	0.12	46,46,46,46	0
56	MG	1A	4137	1/1	0.97	0.31	39,39,39,39	0
56	MG	1A	3199	1/1	0.97	0.09	15,15,15,15	0
56	MG	1A	3808	1/1	0.97	0.06	39,39,39,39	0
56	MG	1a	1828	1/1	0.97	0.05	57,57,57,57	0
56	MG	1A	3200	1/1	0.97	0.07	27,27,27,27	0
56	MG	1A	4142	1/1	0.97	0.12	29,29,29,29	0
56	MG	2A	3853	1/1	0.97	0.08	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3157	1/1	0.97	0.23	51,51,51,51	0
56	MG	1A	3962	1/1	0.97	0.08	63,63,63,63	0
56	MG	1A	3550	1/1	0.97	0.07	54,54,54,54	0
56	MG	1A	3965	1/1	0.97	0.04	55,55,55,55	0
56	MG	1B	202	1/1	0.97	0.17	38,38,38,38	0
56	MG	1A	3029	1/1	0.97	0.15	30,30,30,30	0
56	MG	2A	3865	1/1	0.97	0.05	40,40,40,40	0
56	MG	1B	204	1/1	0.97	0.07	42,42,42,42	0
56	MG	1A	3202	1/1	0.97	0.06	28,28,28,28	0
56	MG	1A	3070	1/1	0.97	0.06	27,27,27,27	0
56	MG	2A	3167	1/1	0.97	0.33	57,57,57,57	0
56	MG	1A	3092	1/1	0.97	0.14	35,35,35,35	0
56	MG	2a	1738	1/1	0.97	0.10	69,69,69,69	0
56	MG	1A	3093	1/1	0.97	0.06	29,29,29,29	0
56	MG	1A	3154	1/1	0.97	0.17	31,31,31,31	0
56	MG	2A	3874	1/1	0.97	0.16	38,38,38,38	0
56	MG	2A	3875	1/1	0.97	0.06	60,60,60,60	0
56	MG	2a	1743	1/1	0.97	0.07	78,78,78,78	0
56	MG	1B	210	1/1	0.97	0.24	42,42,42,42	0
56	MG	1a	1642	1/1	0.97	0.12	48,48,48,48	0
56	MG	1A	3208	1/1	0.97	0.05	35,35,35,35	0
56	MG	1A	3094	1/1	0.97	0.18	21,21,21,21	0
56	MG	2A	3623	1/1	0.97	0.10	44,44,44,44	0
56	MG	1A	3019	1/1	0.97	0.14	32,32,32,32	0
56	MG	2a	1750	1/1	0.97	0.06	70,70,70,70	0
56	MG	1A	3072	1/1	0.97	0.10	40,40,40,40	0
56	MG	2A	3178	1/1	0.97	0.06	55,55,55,55	0
56	MG	2A	3627	1/1	0.97	0.05	37,37,37,37	0
56	MG	2a	1755	1/1	0.97	0.06	70,70,70,70	0
56	MG	1A	3821	1/1	0.97	0.09	44,44,44,44	0
56	MG	1A	3393	1/1	0.97	0.13	48,48,48,48	0
56	MG	2a	1758	1/1	0.97	0.05	56,56,56,56	0
56	MG	1A	3267	1/1	0.97	0.15	30,30,30,30	0
56	MG	2a	1760	1/1	0.97	0.05	72,72,72,72	0
56	MG	1A	3824	1/1	0.97	0.07	37,37,37,37	0
56	MG	1A	3981	1/1	0.97	0.06	71,71,71,71	0
56	MG	1A	3982	1/1	0.97	0.15	40,40,40,40	0
56	MG	2A	3634	1/1	0.97	0.08	33,33,33,33	0
56	MG	1A	3212	1/1	0.97	0.18	32,32,32,32	0
56	MG	1A	3565	1/1	0.97	0.24	47,47,47,47	0
56	MG	2A	3188	1/1	0.97	0.05	53,53,53,53	0
56	MG	1A	3097	1/1	0.97	0.08	33,33,33,33	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3190	1/1	0.97	0.05	49,49,49,49	0
56	MG	2A	3399	1/1	0.97	0.10	36,36,36,36	0
56	MG	1A	3475	1/1	0.97	0.12	41,41,41,41	0
56	MG	1A	3271	1/1	0.97	0.09	38,38,38,38	0
56	MG	1A	3053	1/1	0.97	0.09	33,33,33,33	0
56	MG	1A	3162	1/1	0.97	0.06	28,28,28,28	0
56	MG	2a	1778	1/1	0.97	0.09	62,62,62,62	0
56	MG	1A	3401	1/1	0.97	0.09	43,43,43,43	0
56	MG	1A	3274	1/1	0.97	0.12	20,20,20,20	0
56	MG	1B	235	1/1	0.97	0.06	31,31,31,31	0
56	MG	2A	3408	1/1	0.97	0.05	47,47,47,47	0
56	MG	1A	3992	1/1	0.97	0.07	22,22,22,22	0
56	MG	1A	3126	1/1	0.97	0.04	28,28,28,28	0
56	MG	1A	3278	1/1	0.97	0.31	33,33,33,33	0
56	MG	2a	1786	1/1	0.97	0.06	66,66,66,66	0
56	MG	1D	303	1/1	0.97	0.17	34,34,34,34	0
56	MG	1A	3838	1/1	0.97	0.12	26,26,26,26	0
56	MG	1A	3576	1/1	0.97	0.09	41,41,41,41	0
56	MG	1D	307	1/1	0.97	0.19	32,32,32,32	0
56	MG	1A	3042	1/1	0.97	0.07	29,29,29,29	0
56	MG	1A	3578	1/1	0.97	0.29	27,27,27,27	0
56	MG	1A	3165	1/1	0.97	0.06	41,41,41,41	0
56	MG	1A	3485	1/1	0.97	0.06	38,38,38,38	0
56	MG	1E	302	1/1	0.97	0.06	36,36,36,36	0
56	MG	1E	303	1/1	0.97	0.09	30,30,30,30	0
56	MG	1E	304	1/1	0.97	0.06	24,24,24,24	0
56	MG	1A	3166	1/1	0.97	0.26	40,40,40,40	0
56	MG	2A	3213	1/1	0.97	0.10	48,48,48,48	0
56	MG	1E	306	1/1	0.97	0.06	32,32,32,32	0
56	MG	1A	3223	1/1	0.97	0.19	27,27,27,27	0
56	MG	1A	3167	1/1	0.97	0.15	33,33,33,33	0
56	MG	1A	4009	1/1	0.97	0.07	35,35,35,35	0
56	MG	2a	1804	1/1	0.97	0.09	61,61,61,61	0
56	MG	1A	4010	1/1	0.97	0.06	47,47,47,47	0
56	MG	2A	3670	1/1	0.97	0.06	55,55,55,55	0
56	MG	1A	3284	1/1	0.97	0.10	33,33,33,33	0
56	MG	1A	3031	1/1	0.97	0.25	32,32,32,32	0
56	MG	1A	3346	1/1	0.97	0.09	38,38,38,38	0
56	MG	1A	3129	1/1	0.97	0.05	35,35,35,35	0
56	MG	1A	3588	1/1	0.97	0.13	48,48,48,48	0
56	MG	1A	3171	1/1	0.97	0.11	34,34,34,34	0
56	MG	2E	308	1/1	0.97	0.07	53,53,53,53	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3012	1/1	0.97	0.06	41,41,41,41	0
56	MG	2F	302	1/1	0.97	0.06	57,57,57,57	0
56	MG	1A	3289	1/1	0.97	0.04	27,27,27,27	0
56	MG	1A	3857	1/1	0.97	0.13	30,30,30,30	0
56	MG	1A	3130	1/1	0.97	0.07	37,37,37,37	0
56	MG	1A	3351	1/1	0.97	0.09	47,47,47,47	0
56	MG	1A	3497	1/1	0.97	0.13	32,32,32,32	0
56	MG	1A	3498	1/1	0.97	0.24	30,30,30,30	0
56	MG	1A	4024	1/1	0.97	0.10	46,46,46,46	0
56	MG	1A	3352	1/1	0.97	0.06	29,29,29,29	0
56	MG	1N	201	1/1	0.97	0.19	43,43,43,43	0
56	MG	1N	202	1/1	0.97	0.04	35,35,35,35	0
56	MG	1A	3864	1/1	0.97	0.10	49,49,49,49	0
56	MG	2A	3448	1/1	0.97	0.07	52,52,52,52	0
56	MG	1A	3353	1/1	0.97	0.09	65,65,65,65	0
56	MG	1A	3131	1/1	0.97	0.12	28,28,28,28	0
56	MG	1A	3731	1/1	0.97	0.06	30,30,30,30	0
56	MG	1A	4032	1/1	0.97	0.05	37,37,37,37	0
56	MG	1A	4033	1/1	0.97	0.10	56,56,56,56	0
56	MG	2A	3454	1/1	0.97	0.21	45,45,45,45	0
56	MG	2V	201	1/1	0.97	0.10	42,42,42,42	0
56	MG	1A	3230	1/1	0.97	0.14	32,32,32,32	0
56	MG	2A	3697	1/1	0.97	0.08	36,36,36,36	0
56	MG	2A	3456	1/1	0.97	0.15	47,47,47,47	0
56	MG	2W	203	1/1	0.97	0.07	54,54,54,54	0
56	MG	1A	3422	1/1	0.97	0.25	46,46,46,46	0
56	MG	1A	3606	1/1	0.97	0.10	45,45,45,45	0
56	MG	1A	3424	1/1	0.97	0.10	38,38,38,38	0
56	MG	2A	3460	1/1	0.97	0.12	39,39,39,39	0
56	MG	1A	3176	1/1	0.97	0.09	46,46,46,46	0
56	MG	1A	3426	1/1	0.97	0.14	50,50,50,50	0
56	MG	1A	3295	1/1	0.97	0.19	43,43,43,43	0
56	MG	1A	3613	1/1	0.97	0.05	29,29,29,29	0
56	MG	1A	3077	1/1	0.97	0.12	30,30,30,30	0
56	MG	2A	3251	1/1	0.97	0.16	58,58,58,58	0
56	MG	1A	3297	1/1	0.97	0.23	39,39,39,39	0
56	MG	1A	4045	1/1	0.97	0.06	32,32,32,32	0
56	MG	1a	1717	1/1	0.97	0.12	54,54,54,54	0
56	MG	1A	3880	1/1	0.97	0.06	38,38,38,38	0
56	MG	1a	1719	1/1	0.97	0.10	42,42,42,42	0
56	MG	1A	3746	1/1	0.97	0.07	22,22,22,22	0
56	MG	1A	3014	1/1	0.97	0.12	32,32,32,32	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3361	1/1	0.97	0.07	47,47,47,47	0
56	MG	1A	3179	1/1	0.97	0.11	36,36,36,36	0
56	MG	2A	3050	1/1	0.97	0.14	22,22,22,22	0
56	MG	1A	3620	1/1	0.97	0.07	38,38,38,38	0
56	MG	1T	8001	1/1	0.97	0.06	41,41,41,41	0
56	MG	1A	4055	1/1	0.97	0.04	47,47,47,47	0
56	MG	1U	202	1/1	0.97	0.21	32,32,32,32	0
56	MG	1A	3751	1/1	0.97	0.13	25,25,25,25	0
56	MG	1U	205	1/1	0.97	0.10	40,40,40,40	0
56	MG	2A	3729	1/1	0.97	0.05	36,36,36,36	0
56	MG	1U	206	1/1	0.97	0.06	28,28,28,28	0
56	MG	1A	3752	1/1	0.97	0.05	25,25,25,25	0
56	MG	1A	3434	1/1	0.97	0.26	39,39,39,39	0
56	MG	1a	1736	1/1	0.97	0.09	51,51,51,51	0
56	MG	2A	3272	1/1	0.97	0.24	47,47,47,47	0
56	MG	1a	1737	1/1	0.97	0.09	39,39,39,39	0
56	MG	1A	3514	1/1	0.97	0.17	24,24,24,24	0
56	MG	1A	3435	1/1	0.97	0.19	41,41,41,41	0
56	MG	1a	1740	1/1	0.97	0.14	33,33,33,33	0
56	MG	1a	1741	1/1	0.97	0.10	52,52,52,52	0
56	MG	1A	3183	1/1	0.97	0.13	35,35,35,35	0
56	MG	1A	3759	1/1	0.97	0.06	40,40,40,40	0
56	MG	2A	3069	1/1	0.97	0.06	29,29,29,29	0
56	MG	1X	102	1/1	0.97	0.15	29,29,29,29	0
56	MG	1A	3898	1/1	0.97	0.09	19,19,19,19	0
56	MG	1A	3625	1/1	0.97	0.14	22,22,22,22	0
56	MG	1A	3302	1/1	0.97	0.22	41,41,41,41	0
56	MG	1A	3519	1/1	0.97	0.21	29,29,29,29	0
56	MG	2A	3075	1/1	0.97	0.15	44,44,44,44	0
58	K	1A	4148	1/1	0.97	0.08	56,56,56,56	0
56	MG	1A	3080	1/1	0.97	0.20	33,33,33,33	0
56	MG	1A	3767	1/1	0.97	0.06	43,43,43,43	0
56	MG	2A	3526	1/1	0.98	0.05	45,45,45,45	0
56	MG	1A	3959	1/1	0.98	0.07	43,43,43,43	0
56	MG	1a	1778	1/1	0.98	0.06	59,59,59,59	0
56	MG	1A	3233	1/1	0.98	0.17	26,26,26,26	0
56	MG	1A	3961	1/1	0.98	0.05	60,60,60,60	0
56	MG	1A	4076	1/1	0.98	0.07	29,29,29,29	0
56	MG	2A	3367	1/1	0.98	0.15	50,50,50,50	0
56	MG	1A	4077	1/1	0.98	0.08	63,63,63,63	0
56	MG	2A	3902	1/1	0.98	0.16	51,51,51,51	0
56	MG	1A	3172	1/1	0.98	0.05	15,15,15,15	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1784	1/1	0.98	0.08	52,52,52,52	0
56	MG	1A	3963	1/1	0.98	0.12	49,49,49,49	0
56	MG	1A	4080	1/1	0.98	0.08	19,19,19,19	0
56	MG	1a	1787	1/1	0.98	0.04	33,33,33,33	0
56	MG	1A	3626	1/1	0.98	0.06	28,28,28,28	0
56	MG	1G	3001	1/1	0.98	0.11	33,33,33,33	0
56	MG	2A	3542	1/1	0.98	0.05	43,43,43,43	0
56	MG	2A	3063	1/1	0.98	0.06	68,68,68,68	0
56	MG	1a	1791	1/1	0.98	0.06	53,53,53,53	0
56	MG	1A	3151	1/1	0.98	0.18	37,37,37,37	0
56	MG	1a	1793	1/1	0.98	0.05	71,71,71,71	0
56	MG	1A	3008	1/1	0.98	0.04	15,15,15,15	0
56	MG	1A	4084	1/1	0.98	0.14	31,31,31,31	0
56	MG	1A	3275	1/1	0.98	0.22	27,27,27,27	0
56	MG	1A	3863	1/1	0.98	0.06	50,50,50,50	0
56	MG	2A	3551	1/1	0.98	0.06	33,33,33,33	0
56	MG	1A	3697	1/1	0.98	0.10	51,51,51,51	0
56	MG	1a	1799	1/1	0.98	0.06	64,64,64,64	0
56	MG	1a	1800	1/1	0.98	0.04	57,57,57,57	0
56	MG	1a	1801	1/1	0.98	0.06	63,63,63,63	0
56	MG	1A	3397	1/1	0.98	0.05	30,30,30,30	0
56	MG	2A	3557	1/1	0.98	0.04	47,47,47,47	0
56	MG	2a	1731	1/1	0.98	0.07	55,55,55,55	0
56	MG	1A	4089	1/1	0.98	0.08	30,30,30,30	0
56	MG	1A	3574	1/1	0.98	0.20	35,35,35,35	0
56	MG	1A	3276	1/1	0.98	0.20	35,35,35,35	0
56	MG	1A	3089	1/1	0.98	0.10	14,14,14,14	0
56	MG	2A	3562	1/1	0.98	0.07	41,41,41,41	0
56	MG	1A	3205	1/1	0.98	0.06	33,33,33,33	0
56	MG	1A	3704	1/1	0.98	0.06	46,46,46,46	0
56	MG	1A	3705	1/1	0.98	0.07	44,44,44,44	0
56	MG	1A	3121	1/1	0.98	0.17	28,28,28,28	0
56	MG	1A	4098	1/1	0.98	0.05	29,29,29,29	0
56	MG	1A	3155	1/1	0.98	0.10	26,26,26,26	0
56	MG	1A	3156	1/1	0.98	0.07	33,33,33,33	0
56	MG	1A	3876	1/1	0.98	0.08	22,22,22,22	0
56	MG	2A	3401	1/1	0.98	0.10	48,48,48,48	0
56	MG	1A	4102	1/1	0.98	0.08	32,32,32,32	0
56	MG	1A	3180	1/1	0.98	0.09	36,36,36,36	0
56	MG	1A	4104	1/1	0.98	0.10	37,37,37,37	0
56	MG	1A	3182	1/1	0.98	0.08	41,41,41,41	0
56	MG	1A	3794	1/1	0.98	0.08	46,46,46,46	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1820	1/1	0.98	0.16	41,41,41,41	0
56	MG	1A	3713	1/1	0.98	0.04	15,15,15,15	0
56	MG	1Q	202	1/1	0.98	0.04	39,39,39,39	0
56	MG	1a	1823	1/1	0.98	0.05	55,55,55,55	0
56	MG	1A	3881	1/1	0.98	0.09	26,26,26,26	0
56	MG	1A	3536	1/1	0.98	0.16	39,39,39,39	0
56	MG	1A	3245	1/1	0.98	0.03	37,37,37,37	0
56	MG	1A	3137	1/1	0.98	0.11	15,15,15,15	0
56	MG	1A	3717	1/1	0.98	0.05	44,44,44,44	0
56	MG	1A	3539	1/1	0.98	0.15	33,33,33,33	0
56	MG	2A	3587	1/1	0.98	0.09	44,44,44,44	0
56	MG	2a	1763	1/1	0.98	0.04	75,75,75,75	0
56	MG	1A	3122	1/1	0.98	0.38	40,40,40,40	0
56	MG	1A	3185	1/1	0.98	0.21	34,34,34,34	0
56	MG	1A	3123	1/1	0.98	0.09	39,39,39,39	0
56	MG	1U	203	1/1	0.98	0.34	40,40,40,40	0
56	MG	1A	3995	1/1	0.98	0.04	21,21,21,21	0
56	MG	2A	3594	1/1	0.98	0.04	37,37,37,37	0
56	MG	1a	1682	1/1	0.98	0.18	43,43,43,43	0
56	MG	1A	3722	1/1	0.98	0.04	53,53,53,53	0
56	MG	2a	1772	1/1	0.98	0.08	89,89,89,89	0
56	MG	2A	3773	1/1	0.98	0.05	38,38,38,38	0
56	MG	2a	1774	1/1	0.98	0.06	67,67,67,67	0
56	MG	1A	3892	1/1	0.98	0.06	38,38,38,38	0
56	MG	2A	3775	1/1	0.98	0.05	53,53,53,53	0
56	MG	2A	3598	1/1	0.98	0.06	57,57,57,57	0
56	MG	1V	201	1/1	0.98	0.16	28,28,28,28	0
56	MG	20	103	1/1	0.98	0.19	55,55,55,55	0
56	MG	1V	202	1/1	0.98	0.12	35,35,35,35	0
56	MG	1A	3215	1/1	0.98	0.06	38,38,38,38	0
56	MG	1A	4122	1/1	0.98	0.04	37,37,37,37	0
56	MG	1A	3216	1/1	0.98	0.10	40,40,40,40	0
56	MG	1W	203	1/1	0.98	0.10	30,30,30,30	0
56	MG	1A	3725	1/1	0.98	0.09	25,25,25,25	0
56	MG	1W	205	1/1	0.98	0.06	30,30,30,30	0
56	MG	2A	3785	1/1	0.98	0.05	56,56,56,56	0
56	MG	2A	3786	1/1	0.98	0.07	64,64,64,64	0
56	MG	1A	3217	1/1	0.98	0.16	33,33,33,33	0
56	MG	2A	3121	1/1	0.98	0.08	41,41,41,41	0
56	MG	1A	3457	1/1	0.98	0.08	42,42,42,42	0
56	MG	1X	101	1/1	0.98	0.07	26,26,26,26	0
56	MG	1A	4128	1/1	0.98	0.09	32,32,32,32	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3125	1/1	0.98	0.17	46,46,46,46	0
56	MG	1X	103	1/1	0.98	0.07	34,34,34,34	0
56	MG	1A	3140	1/1	0.98	0.08	34,34,34,34	0
56	MG	1A	3653	1/1	0.98	0.03	25,25,25,25	0
56	MG	1A	3141	1/1	0.98	0.19	27,27,27,27	0
56	MG	1A	3655	1/1	0.98	0.05	18,18,18,18	0
56	MG	1A	3656	1/1	0.98	0.10	19,19,19,19	0
56	MG	1A	3596	1/1	0.98	0.06	19,19,19,19	0
56	MG	1A	3734	1/1	0.98	0.05	38,38,38,38	0
56	MG	2A	3134	1/1	0.98	0.06	38,38,38,38	0
56	MG	2A	3802	1/1	0.98	0.04	60,60,60,60	0
56	MG	1A	4138	1/1	0.98	0.08	35,35,35,35	0
56	MG	1A	3735	1/1	0.98	0.05	34,34,34,34	0
56	MG	1A	3908	1/1	0.98	0.12	37,37,37,37	0
56	MG	1A	3658	1/1	0.98	0.06	27,27,27,27	0
56	MG	2A	3294	1/1	0.98	0.17	47,47,47,47	0
56	MG	10	103	1/1	0.98	0.09	35,35,35,35	0
56	MG	1A	3597	1/1	0.98	0.04	32,32,32,32	0
56	MG	1A	4143	1/1	0.98	0.29	28,28,28,28	0
56	MG	1A	4145	1/1	0.98	0.11	25,25,25,25	0
56	MG	1A	3020	1/1	0.98	0.04	40,40,40,40	0
56	MG	1A	3600	1/1	0.98	0.12	28,28,28,28	0
56	MG	1A	3081	1/1	0.98	0.13	29,29,29,29	0
56	MG	1A	3028	1/1	0.98	0.12	24,24,24,24	0
56	MG	2A	3816	1/1	0.98	0.04	59,59,59,59	0
56	MG	1A	3665	1/1	0.98	0.06	19,19,19,19	0
56	MG	1A	3603	1/1	0.98	0.06	23,23,23,23	0
56	MG	1A	3604	1/1	0.98	0.07	25,25,25,25	0
56	MG	1A	3918	1/1	0.98	0.05	44,44,44,44	0
56	MG	1A	3919	1/1	0.98	0.04	8,8,8,8	0
56	MG	1A	3670	1/1	0.98	0.07	23,23,23,23	0
56	MG	1A	3921	1/1	0.98	0.05	53,53,53,53	0
56	MG	1A	3299	1/1	0.98	0.12	23,23,23,23	0
56	MG	15	101	1/1	0.98	0.13	36,36,36,36	0
56	MG	2A	3826	1/1	0.98	0.07	50,50,50,50	0
56	MG	1A	3145	1/1	0.98	0.24	37,37,37,37	0
56	MG	1A	3554	1/1	0.98	0.05	51,51,51,51	0
56	MG	1A	3065	1/1	0.98	0.12	24,24,24,24	0
56	MG	1B	214	1/1	0.98	0.06	44,44,44,44	0
56	MG	2A	3831	1/1	0.98	0.09	52,52,52,52	0
56	MG	1a	1731	1/1	0.98	0.04	34,34,34,34	0
56	MG	1B	215	1/1	0.98	0.07	39,39,39,39	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3927	1/1	0.98	0.06	50,50,50,50	0
56	MG	2A	3163	1/1	0.98	0.12	55,55,55,55	0
56	MG	1A	3676	1/1	0.98	0.07	31,31,31,31	0
56	MG	1A	3833	1/1	0.98	0.07	14,14,14,14	0
56	MG	1A	3261	1/1	0.98	0.21	40,40,40,40	0
56	MG	1A	3754	1/1	0.98	0.06	53,53,53,53	0
56	MG	1A	3836	1/1	0.98	0.05	54,54,54,54	0
56	MG	1A	3423	1/1	0.98	0.09	39,39,39,39	0
56	MG	1A	3756	1/1	0.98	0.08	17,17,17,17	0
56	MG	1B	225	1/1	0.98	0.04	30,30,30,30	0
56	MG	1A	4044	1/1	0.98	0.12	59,59,59,59	0
56	MG	2A	3487	1/1	0.98	0.12	23,23,23,23	0
56	MG	1A	3612	1/1	0.98	0.04	18,18,18,18	0
56	MG	2A	3174	1/1	0.98	0.04	51,51,51,51	0
56	MG	1A	4046	1/1	0.98	0.04	40,40,40,40	0
56	MG	1A	3067	1/1	0.98	0.23	30,30,30,30	0
56	MG	1A	3343	1/1	0.98	0.13	37,37,37,37	0
56	MG	2A	3859	1/1	0.98	0.07	52,52,52,52	0
56	MG	1B	231	1/1	0.98	0.04	42,42,42,42	0
56	MG	1A	4049	1/1	0.98	0.05	26,26,26,26	0
56	MG	1A	3940	1/1	0.98	0.05	56,56,56,56	0
56	MG	2a	1668	1/1	0.98	0.09	46,46,46,46	0
56	MG	1A	3760	1/1	0.98	0.06	24,24,24,24	0
56	MG	2A	3497	1/1	0.98	0.04	67,67,67,67	0
56	MG	1A	3761	1/1	0.98	0.07	22,22,22,22	0
56	MG	1A	3844	1/1	0.98	0.07	38,38,38,38	0
56	MG	2a	1673	1/1	0.98	0.08	51,51,51,51	0
56	MG	1A	4054	1/1	0.98	0.07	54,54,54,54	0
56	MG	2A	3868	1/1	0.98	0.05	62,62,62,62	0
56	MG	1A	3845	1/1	0.98	0.06	32,32,32,32	0
56	MG	2A	3186	1/1	0.98	0.09	40,40,40,40	0
56	MG	1A	3195	1/1	0.98	0.11	16,16,16,16	0
56	MG	1A	3946	1/1	0.98	0.04	35,35,35,35	0
56	MG	1A	3196	1/1	0.98	0.04	36,36,36,36	0
56	MG	1A	3948	1/1	0.98	0.04	54,54,54,54	0
56	MG	1D	308	1/1	0.98	0.06	40,40,40,40	0
56	MG	2A	3509	1/1	0.98	0.07	36,36,36,36	0
56	MG	1A	3764	1/1	0.98	0.12	50,50,50,50	0
56	MG	1A	4061	1/1	0.98	0.05	33,33,33,33	0
56	MG	1A	3068	1/1	0.98	0.04	15,15,15,15	0
56	MG	1A	4063	1/1	0.98	0.05	50,50,50,50	0
56	MG	1A	3850	1/1	0.98	0.09	38,38,38,38	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3169	1/1	0.98	0.20	34,34,34,34	0
56	MG	1A	3686	1/1	0.98	0.07	27,27,27,27	0
56	MG	1A	3076	1/1	0.98	0.07	34,34,34,34	0
56	MG	1A	3231	1/1	0.98	0.10	26,26,26,26	0
56	MG	2y	3006	1/1	0.98	0.07	86,86,86,86	0
56	MG	2A	3045	1/1	0.98	0.12	42,42,42,42	0
56	MG	1A	3269	1/1	0.98	0.10	36,36,36,36	0
56	MG	1E	308	1/1	0.98	0.09	21,21,21,21	0
56	MG	1A	3957	1/1	0.98	0.04	37,37,37,37	0
56	MG	1A	3061	1/1	0.98	0.07	34,34,34,34	0
59	ZN	14	501	1/1	0.98	0.04	95,95,95,95	0
56	MG	2A	3699	1/1	0.98	0.04	59,59,59,59	0
59	ZN	29	501	1/1	0.98	0.04	74,74,74,74	0
59	ZN	2n	501	1/1	0.98	0.04	104,104,104,104	0
60	SF4	1d	501	8/8	0.98	0.05	58,66,74,76	0
60	SF4	2d	501	8/8	0.98	0.05	67,71,80,97	0
56	MG	1A	3647	1/1	0.99	0.05	29,29,29,29	0
56	MG	1A	3599	1/1	0.99	0.06	25,25,25,25	0
56	MG	1A	3003	1/1	0.99	0.04	22,22,22,22	0
56	MG	1A	3181	1/1	0.99	0.12	33,33,33,33	0
56	MG	2A	3608	1/1	0.99	0.03	63,63,63,63	0
56	MG	2A	3609	1/1	0.99	0.08	42,42,42,42	0
56	MG	2A	3842	1/1	0.99	0.05	45,45,45,45	0
56	MG	1A	3710	1/1	0.99	0.03	29,29,29,29	0
56	MG	1F	304	1/1	0.99	0.23	29,29,29,29	0
56	MG	2E	303	1/1	0.99	0.13	38,38,38,38	0
56	MG	2A	3845	1/1	0.99	0.07	40,40,40,40	0
56	MG	1A	4004	1/1	0.99	0.04	20,20,20,20	0
56	MG	2A	3847	1/1	0.99	0.04	46,46,46,46	0
56	MG	2A	3848	1/1	0.99	0.03	46,46,46,46	0
56	MG	1A	4005	1/1	0.99	0.02	28,28,28,28	0
56	MG	1A	4006	1/1	0.99	0.03	25,25,25,25	0
56	MG	1A	3922	1/1	0.99	0.04	19,19,19,19	0
56	MG	1A	4008	1/1	0.99	0.06	30,30,30,30	0
56	MG	1A	3777	1/1	0.99	0.10	20,20,20,20	0
56	MG	2A	3854	1/1	0.99	0.06	37,37,37,37	0
56	MG	2A	3855	1/1	0.99	0.03	49,49,49,49	0
56	MG	1A	3517	1/1	0.99	0.18	29,29,29,29	0
56	MG	2A	3857	1/1	0.99	0.05	32,32,32,32	0
56	MG	2A	3858	1/1	0.99	0.06	68,68,68,68	0
56	MG	1a	1748	1/1	0.99	0.07	37,37,37,37	0
56	MG	1A	3712	1/1	0.99	0.03	21,21,21,21	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3124	1/1	0.99	0.14	24,24,24,24	0
56	MG	1A	4013	1/1	0.99	0.04	54,54,54,54	0
56	MG	1A	3078	1/1	0.99	0.06	42,42,42,42	0
56	MG	1A	3969	1/1	0.99	0.03	59,59,59,59	0
56	MG	1A	3013	1/1	0.99	0.06	27,27,27,27	0
56	MG	1a	1755	1/1	0.99	0.07	51,51,51,51	0
56	MG	1A	4065	1/1	0.99	0.06	37,37,37,37	0
56	MG	1A	3929	1/1	0.99	0.03	42,42,42,42	0
56	MG	1A	3250	1/1	0.99	0.07	21,21,21,21	0
56	MG	2a	1753	1/1	0.99	0.03	48,48,48,48	0
56	MG	1B	224	1/1	0.99	0.06	34,34,34,34	0
56	MG	2A	3009	1/1	0.99	0.04	43,43,43,43	0
56	MG	2A	3707	1/1	0.99	0.04	62,62,62,62	0
56	MG	1A	3011	1/1	0.99	0.07	22,22,22,22	0
56	MG	2A	3078	1/1	0.99	0.19	31,31,31,31	0
56	MG	1A	3288	1/1	0.99	0.14	20,20,20,20	0
56	MG	1A	3609	1/1	0.99	0.05	35,35,35,35	0
56	MG	1A	3787	1/1	0.99	0.04	34,34,34,34	0
56	MG	1A	3788	1/1	0.99	0.02	30,30,30,30	0
56	MG	1A	4123	1/1	0.99	0.11	31,31,31,31	0
56	MG	1A	3896	1/1	0.99	0.03	21,21,21,21	0
56	MG	1A	3897	1/1	0.99	0.02	29,29,29,29	0
56	MG	1A	3044	1/1	0.99	0.08	31,31,31,31	0
56	MG	1A	3235	1/1	0.99	0.12	27,27,27,27	0
56	MG	1a	1770	1/1	0.99	0.04	62,62,62,62	0
56	MG	2A	3021	1/1	0.99	0.07	28,28,28,28	0
56	MG	1A	4029	1/1	0.99	0.03	20,20,20,20	0
56	MG	2A	3723	1/1	0.99	0.03	47,47,47,47	0
56	MG	2A	3499	1/1	0.99	0.04	49,49,49,49	0
56	MG	1A	3021	1/1	0.99	0.02	15,15,15,15	0
56	MG	1A	3662	1/1	0.99	0.03	38,38,38,38	0
56	MG	2w	3001	1/1	0.99	0.15	48,48,48,48	0
56	MG	1A	3039	1/1	0.99	0.06	27,27,27,27	0
56	MG	1A	4132	1/1	0.99	0.16	22,22,22,22	0
56	MG	1A	3614	1/1	0.99	0.04	22,22,22,22	0
56	MG	1Q	205	1/1	0.99	0.05	28,28,28,28	0
56	MG	1R	201	1/1	0.99	0.24	27,27,27,27	0
56	MG	2A	3732	1/1	0.99	0.04	40,40,40,40	0
56	MG	1A	4134	1/1	0.99	0.04	27,27,27,27	0
56	MG	1D	306	1/1	0.99	0.12	22,22,22,22	0
56	MG	1A	3173	1/1	0.99	0.04	24,24,24,24	0
56	MG	1A	3106	1/1	0.99	0.07	25,25,25,25	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3867	1/1	0.99	0.03	26,26,26,26	0
56	MG	2A	3512	1/1	0.99	0.03	66,66,66,66	0
56	MG	1D	310	1/1	0.99	0.09	15,15,15,15	0
56	MG	2A	3514	1/1	0.99	0.04	39,39,39,39	0
56	MG	1A	3667	1/1	0.99	0.03	21,21,21,21	0
56	MG	1A	3054	1/1	0.99	0.15	37,37,37,37	0
56	MG	2A	3038	1/1	0.99	0.04	35,35,35,35	0
56	MG	1A	3669	1/1	0.99	0.05	21,21,21,21	0
56	MG	1A	3055	1/1	0.99	0.05	23,23,23,23	0
56	MG	1a	1725	1/1	0.99	0.07	40,40,40,40	0
56	MG	2A	3593	1/1	0.99	0.06	36,36,36,36	0
56	MG	1a	1790	1/1	0.99	0.05	38,38,38,38	0
56	MG	1A	3671	1/1	0.99	0.06	30,30,30,30	0
56	MG	1A	3034	1/1	0.99	0.11	24,24,24,24	0
56	MG	1A	4144	1/1	0.99	0.10	30,30,30,30	0
59	ZN	1Y	202	1/1	0.99	0.03	66,66,66,66	0
56	MG	1A	3432	1/1	0.99	0.05	31,31,31,31	0
59	ZN	19	102	1/1	0.99	0.05	41,41,41,41	0
59	ZN	1n	501	1/1	0.99	0.05	59,59,59,59	0
59	ZN	2Y	501	1/1	0.99	0.04	87,87,87,87	0
56	MG	1A	3703	1/1	0.99	0.05	20,20,20,20	0
59	ZN	25	102	1/1	0.99	0.03	55,55,55,55	0
56	MG	2A	3527	1/1	0.99	0.06	35,35,35,35	0
56	MG	1A	4093	1/1	0.99	0.14	27,27,27,27	0
56	MG	1A	3066	1/1	0.99	0.11	16,16,16,16	0
56	MG	1A	3136	1/1	0.99	0.09	33,33,33,33	0
59	ZN	16	102	1/1	1.00	0.04	38,38,38,38	0
56	MG	1A	3743	1/1	1.00	0.07	38,38,38,38	0
56	MG	1A	3294	1/1	1.00	0.02	23,23,23,23	0
56	MG	1A	3736	1/1	1.00	0.06	20,20,20,20	0
56	MG	1A	3890	1/1	1.00	0.06	35,35,35,35	0
56	MG	1A	4025	1/1	1.00	0.04	45,45,45,45	0
59	ZN	26	501	1/1	1.00	0.04	56,56,56,56	0
56	MG	1a	1732	1/1	1.00	0.05	32,32,32,32	0
56	MG	2A	3715	1/1	1.00	0.03	26,26,26,26	0
56	MG	1A	3935	1/1	1.00	0.05	39,39,39,39	0
59	ZN	15	102	1/1	1.00	0.05	41,41,41,41	0

6.5 Other polymers ⓘ

There are no such residues in this entry.