



wwPDB X-ray Structure Validation Summary Report ⓘ

Aug 4, 2026 – 06:40 PM EDT

PDB ID : 7RQB / pdb_00007rqb
Title : Crystal structure of the Thermus thermophilus 70S ribosome in complex with protein Y, A-site aminoacyl-tRNA analog ACC-PMN, and P-site MAI-tripeptidyl-tRNA analog ACCA-IAM at 2.45Å resolution
Authors : Syroegin, E.A.; Flemmich, L.; Klepacki, D.; Vazquez-Laslop, N.; Micura, R.; Polikanov, Y.S.
Deposited on : 2021-08-06
Resolution : 2.45 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

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

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

MolProbity	:	4-5-2 with Phenix2.0
Mogul	:	2022.3.0, CSD as543be (2022)
Xtriage (Phenix)	:	2.0
EDS	:	3.0
Percentile statistics	:	20250101.v01 (using entries in the PDB archive January 1st 2025)
CCP4	:	9.0.015 (Gargrove)
Density-Fitness	:	1.0.12
Ideal geometry (proteins)	:	Engh & Huber (2001)
Ideal geometry (DNA, RNA)	:	Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP)	:	2.50

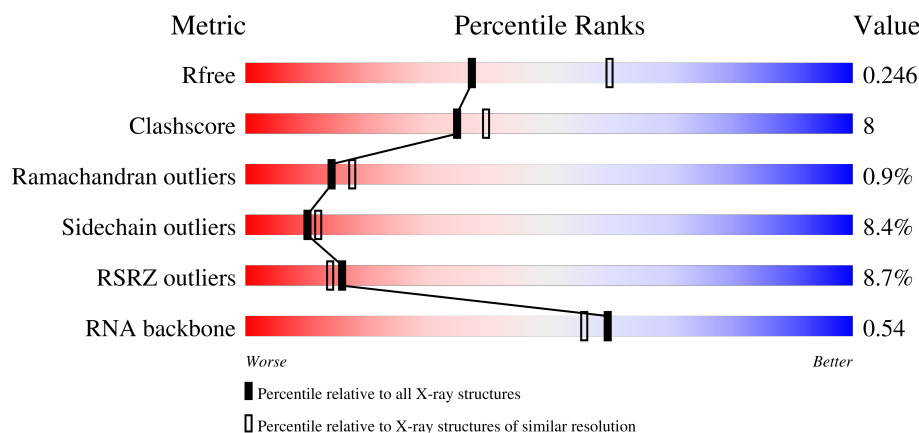
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

The reported resolution of this entry is 2.45 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	1190 (2.46-2.46)
Clashscore	190562	1229 (2.46-2.46)
Ramachandran outliers	187476	1218 (2.46-2.46)
Sidechain outliers	187428	1218 (2.46-2.46)
RSRZ outliers	180081	1190 (2.46-2.46)
RNA backbone	3983	1023 (2.72-2.20)

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	6% 69% 24% 5% •
1	2A	2915	6% 63% 29% 7% •
2	1B	121	% 77% 19% ••

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Mol	Chain	Length	Quality of chain
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	
14	2S	112	

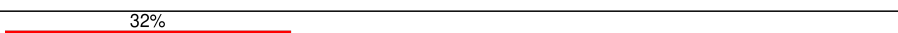
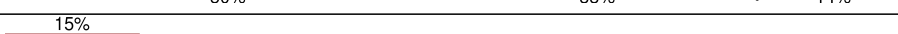
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Mol	Chain	Length	Quality of chain
15	1T	146	
15	2T	146	
16	1U	118	
16	2U	118	
17	1V	101	
17	2V	101	
18	1W	113	
18	2W	113	
19	1X	96	
19	2X	96	
20	1Y	110	
20	2Y	110	
21	1Z	206	
21	2Z	206	
22	10	85	
22	20	85	
23	11	98	
23	21	98	
24	12	72	
24	22	72	
25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

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Mol	Chain	Length	Quality of chain
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	
39	2h	138	

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Mol	Chain	Length	Quality of chain
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	
52	1u	27	

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Mol	Chain	Length	Quality of chain
52	2u	27	
53	1y	113	
53	2y	113	
54	1w	4	
54	2w	4	
55	1x	4	
55	2x	4	
56	1v	3	
56	2v	3	

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

Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
57	MG	1A	3576	-	-	-	X
57	MG	1A	3698	-	-	-	X
57	MG	1a	1702	-	-	-	X
57	MG	1a	1711	-	-	-	X
57	MG	1a	1822	-	-	-	X
57	MG	2A	3047	-	-	-	X
57	MG	2A	3159	-	-	-	X
57	MG	2a	3009	-	-	-	X
57	MG	2a	3045	-	-	-	X

2 Entry composition

There are 62 unique types of molecules in this entry. The entry contains 297350 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	2872	Total	C	N	O	P	0	0	0
			61869	27540	11574	19884	2871			
1	2A	2867	Total	C	N	O	P	0	0	0
			61758	27491	11552	19850	2865			

- 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
			2572	1145	476	832	119			
2	2B	120	Total	C	N	O	P	0	0	0
			2573	1146	476	832	119			

- 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
			2131	1346	422	360	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
			1426	916	253	253	4			
6	2G	181	Total	C	N	O	S	0	0	0
			1424	912	259	249	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	173	Total	C	N	O	S	0	0	0
			1324	842	247	234	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
8	1I	147	Total	C	N	O	S	0	0	0
			1094	699	191	203	1			
8	2I	146	Total	C	N	O	S	0	0	0
			1076	687	186	202	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
			1121	722	208	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
			877	553	175	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
			775	498	141	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
			810	520	153	131	6			
20	2Y	107	Total	C	N	O	S	0	0	0
			810	519	153	132	6			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
21	1Z	203	Total	C	N	O	S	0	0	0
			1587	1011	282	292	2			
21	2Z	201	Total	C	N	O	S	0	0	0
			1557	995	274	286	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
			650	401	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
			754	475	148	130	1			
23	21	97	Total	C	N	O	S	0	0	0
			759	478	149	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
			592	368	119	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
			546	346	96	99	5			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
26	24	69	Total	C	N	O	S	0	0	0
			536	342	98	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
			459	288	90	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	1504	Total	C	N	O	P	0	0	0
			32331	14396	5990	10441	1504			

- 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
			1842	1175	330	332	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
			1558	979	305	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
			1665	1043	329	286	7			
35	2d	208	Total	C	N	O	S	0	0	0
			1668	1047	330	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
			1133	716	214	199	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
			814	516	144	151	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
			1235	769	244	216	6			
38	2g	155	Total	C	N	O	S	0	0	0
			1229	766	241	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
			1098	694	210	192	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
			986	625	193	168			
40	2i	126	Total	C	N	O	0	0	0
			966	613	186	167			

- 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
			719	446	142	131			
41	2j	96	Total	C	N	O	0	0	0
			710	442	137	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
			834	520	156	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	116	Total	C	N	O	S	0	0	0
			914	564	189	159	2			
44	2m	114	Total	C	N	O	S	0	0	0
			895	550	186	157	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
			648	415	120	111	2			
50	2s	83	Total	C	N	O	S	0	0	0
			645	410	118	115	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
			732	449	157	124	2			
51	2t	98	Total	C	N	O	S	0	0	0
			733	451	154	126	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 protein called Ribosome-associated inhibitor A.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
53	1y	97	Total	C	N	O	S	0	0	0
			764	478	144	139	3			
53	2y	96	Total	C	N	O	S	0	0	0
			749	468	141	137	3			

- Molecule 54 is a RNA chain called A-site Aminoacyl-tRNA Analog.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
54	1w	4	Total	C	N	O	P	0	0	1
			78	40	13	22	3			
54	2w	4	Total	C	N	O	P	0	0	1
			78	40	13	22	3			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
55	1x	4	Total	C	N	O	P	0	0	1
			63	28	12	20	3			
55	2x	4	Total	C	N	O	P	0	0	1
			63	28	12	20	3			

- Molecule 56 is a protein called P-site Peptidyl-tRNA Analog Peptide.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
56	1v	3	Total	C	N	O	S	0	0	0
			21	14	3	3	1			
56	2v	3	Total	C	N	O	S	0	0	0
			21	14	3	3	1			

- Molecule 57 is MAGNESIUM ION (CCD ID: MG) (formula: Mg).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	1A	1013	Total	Mg	0	0
			1013	1013		
57	1B	30	Total	Mg	0	0
			30	30		
57	1D	17	Total	Mg	0	0
			17	17		
57	1E	8	Total	Mg	0	0
			8	8		
57	1F	17	Total	Mg	0	0
			17	17		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	1G	4	Total 4	Mg 4	0	0
57	1H	2	Total 2	Mg 2	0	0
57	1N	4	Total 4	Mg 4	0	0
57	1O	1	Total 1	Mg 1	0	0
57	1P	3	Total 3	Mg 3	0	0
57	1Q	5	Total 5	Mg 5	0	0
57	1R	7	Total 7	Mg 7	0	0
57	1T	7	Total 7	Mg 7	0	0
57	1U	5	Total 5	Mg 5	0	0
57	1V	7	Total 7	Mg 7	0	0
57	1W	3	Total 3	Mg 3	0	0
57	1Y	1	Total 1	Mg 1	0	0
57	1Z	1	Total 1	Mg 1	0	0
57	10	8	Total 8	Mg 8	0	0
57	11	5	Total 5	Mg 5	0	0
57	13	3	Total 3	Mg 3	0	0
57	15	7	Total 7	Mg 7	0	0
57	17	6	Total 6	Mg 6	0	0
57	18	3	Total 3	Mg 3	0	0
57	19	1	Total 1	Mg 1	0	0
57	1a	267	Total 267	Mg 267	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	1b	1	Total 1	Mg 1	0	0
57	1d	5	Total 5	Mg 5	0	0
57	1e	5	Total 5	Mg 5	0	0
57	1f	2	Total 2	Mg 2	0	0
57	1g	2	Total 2	Mg 2	0	0
57	1h	2	Total 2	Mg 2	0	0
57	1i	1	Total 1	Mg 1	0	0
57	1l	2	Total 2	Mg 2	0	0
57	1m	1	Total 1	Mg 1	0	0
57	1n	3	Total 3	Mg 3	0	0
57	1o	3	Total 3	Mg 3	0	0
57	1t	2	Total 2	Mg 2	0	0
57	1y	2	Total 2	Mg 2	0	0
57	1x	2	Total 2	Mg 2	0	0
57	2A	731	Total 731	Mg 731	0	0
57	2B	19	Total 19	Mg 19	0	0
57	2D	10	Total 10	Mg 10	0	0
57	2E	6	Total 6	Mg 6	0	0
57	2F	5	Total 5	Mg 5	0	0
57	2G	2	Total 2	Mg 2	0	0
57	2I	1	Total 1	Mg 1	0	0

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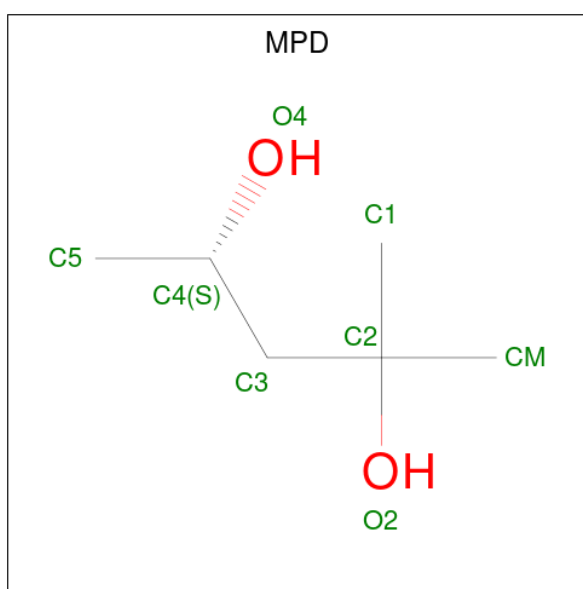
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	2O	2	Total 2	Mg 2	0	0
57	2P	1	Total 1	Mg 1	0	0
57	2Q	3	Total 3	Mg 3	0	0
57	2R	3	Total 3	Mg 3	0	0
57	2T	4	Total 4	Mg 4	0	0
57	2U	2	Total 2	Mg 2	0	0
57	2V	3	Total 3	Mg 3	0	0
57	2W	3	Total 3	Mg 3	0	0
57	2X	1	Total 1	Mg 1	0	0
57	2Y	1	Total 1	Mg 1	0	0
57	20	2	Total 2	Mg 2	0	0
57	23	1	Total 1	Mg 1	0	0
57	25	2	Total 2	Mg 2	0	0
57	27	2	Total 2	Mg 2	0	0
57	28	2	Total 2	Mg 2	0	0
57	29	1	Total 1	Mg 1	0	0
57	2a	183	Total 183	Mg 183	0	0
57	2e	1	Total 1	Mg 1	0	0
57	2f	2	Total 2	Mg 2	0	0
57	2j	1	Total 1	Mg 1	0	0
57	2k	1	Total 1	Mg 1	0	0

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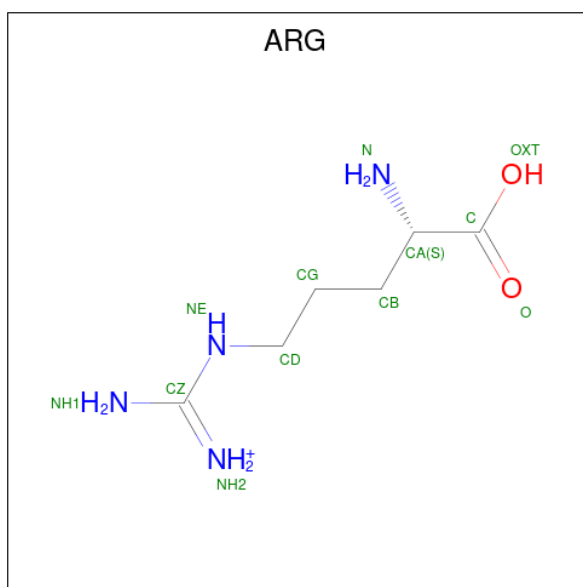
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	2p	1	Total	Mg	0	0
			1	1		
57	2r	1	Total	Mg	0	0
			1	1		
57	2t	1	Total	Mg	0	0
			1	1		
57	2x	2	Total	Mg	0	0
			2	2		

- Molecule 58 is (4S)-2-METHYL-2,4-PENTANEDIOL (CCD ID: MPD) (formula: $C_6H_{14}O_2$).



Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
58	1A	1	Total	C	O	0	0
			8	6	2		
58	1T	1	Total	C	O	0	0
			8	6	2		
58	18	1	Total	C	O	0	0
			8	6	2		
58	1a	1	Total	C	O	0	0
			8	6	2		
58	2A	1	Total	C	O	0	0
			8	6	2		
58	2B	1	Total	C	O	0	0
			8	6	2		

- Molecule 59 is ARGinine (CCD ID: ARG) (formula: $C_6H_{15}N_4O_2$).



Mol	Chain	Residues	Atoms				ZeroOcc	AltConf
59	1B	1	Total	C	N	O	0	0
			12	6	4	2		
59	1F	1	Total	C	N	O	0	0
			12	6	4	2		

- Molecule 60 is ZINC ION (CCD ID: ZN) (formula: Zn).

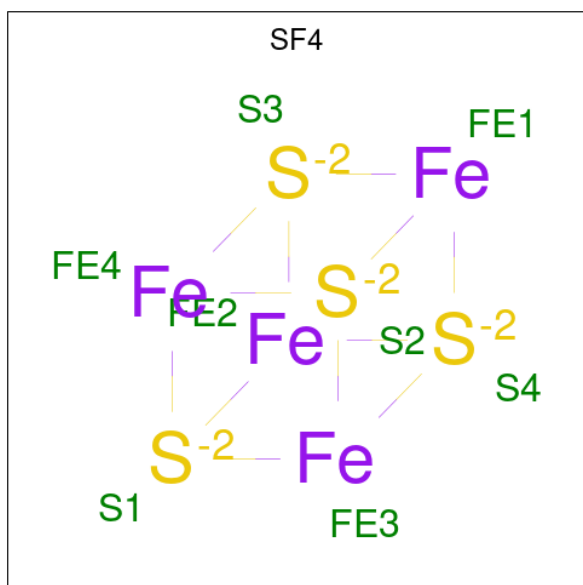
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
60	1Y	1	Total	Zn	0	0
			1	1		
60	14	1	Total	Zn	0	0
			1	1		
60	15	1	Total	Zn	0	0
			1	1		
60	16	1	Total	Zn	0	0
			1	1		
60	19	1	Total	Zn	0	0
			1	1		
60	1n	1	Total	Zn	0	0
			1	1		
60	2Y	1	Total	Zn	0	0
			1	1		
60	24	1	Total	Zn	0	0
			1	1		
60	25	1	Total	Zn	0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

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
60	29	1	Total	Zn	0	0
			1	1		
60	2n	1	Total	Zn	0	0
			1	1		

- Molecule 61 is IRON/SULFUR CLUSTER (CCD ID: SF4) (formula: Fe_4S_4).



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

- Molecule 62 is water.

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
62	1A	3935	Total	O	0	0
			3935	3935		
62	1B	80	Total	O	0	0
			80	80		
62	1D	108	Total	O	0	0
			108	108		
62	1E	74	Total	O	0	0
			74	74		
62	1F	60	Total	O	0	0
			60	60		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
62	1G	16	Total 16	O 16	0	0
62	1H	15	Total 15	O 15	0	0
62	1I	5	Total 5	O 5	0	0
62	1N	54	Total 54	O 54	0	0
62	1O	21	Total 21	O 21	0	0
62	1P	63	Total 63	O 63	0	0
62	1Q	39	Total 39	O 39	0	0
62	1R	28	Total 28	O 28	0	0
62	1S	12	Total 12	O 12	0	0
62	1T	32	Total 32	O 32	0	0
62	1U	53	Total 53	O 53	0	0
62	1V	33	Total 33	O 33	0	0
62	1W	24	Total 24	O 24	0	0
62	1X	24	Total 24	O 24	0	0
62	1Y	15	Total 15	O 15	0	0
62	1Z	7	Total 7	O 7	0	0
62	10	23	Total 23	O 23	0	0
62	11	26	Total 26	O 26	0	0
62	12	10	Total 10	O 10	0	0
62	13	25	Total 25	O 25	0	0
62	14	2	Total 2	O 2	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
62	15	18	Total 18	O 18	0	0
62	16	18	Total 18	O 18	0	0
62	17	17	Total 17	O 17	0	0
62	18	25	Total 25	O 25	0	0
62	19	5	Total 5	O 5	0	0
62	1a	424	Total 424	O 424	0	0
62	1b	1	Total 1	O 1	0	0
62	1d	8	Total 8	O 8	0	0
62	1e	2	Total 2	O 2	0	0
62	1f	2	Total 2	O 2	0	0
62	1g	1	Total 1	O 1	0	0
62	1h	1	Total 1	O 1	0	0
62	1l	2	Total 2	O 2	0	0
62	1o	4	Total 4	O 4	0	0
62	1p	1	Total 1	O 1	0	0
62	1t	1	Total 1	O 1	0	0
62	1y	2	Total 2	O 2	0	0
62	1w	6	Total 6	O 6	0	0
62	1x	2	Total 2	O 2	0	0
62	2A	2058	Total 2058	O 2058	0	0
62	2B	39	Total 39	O 39	0	0

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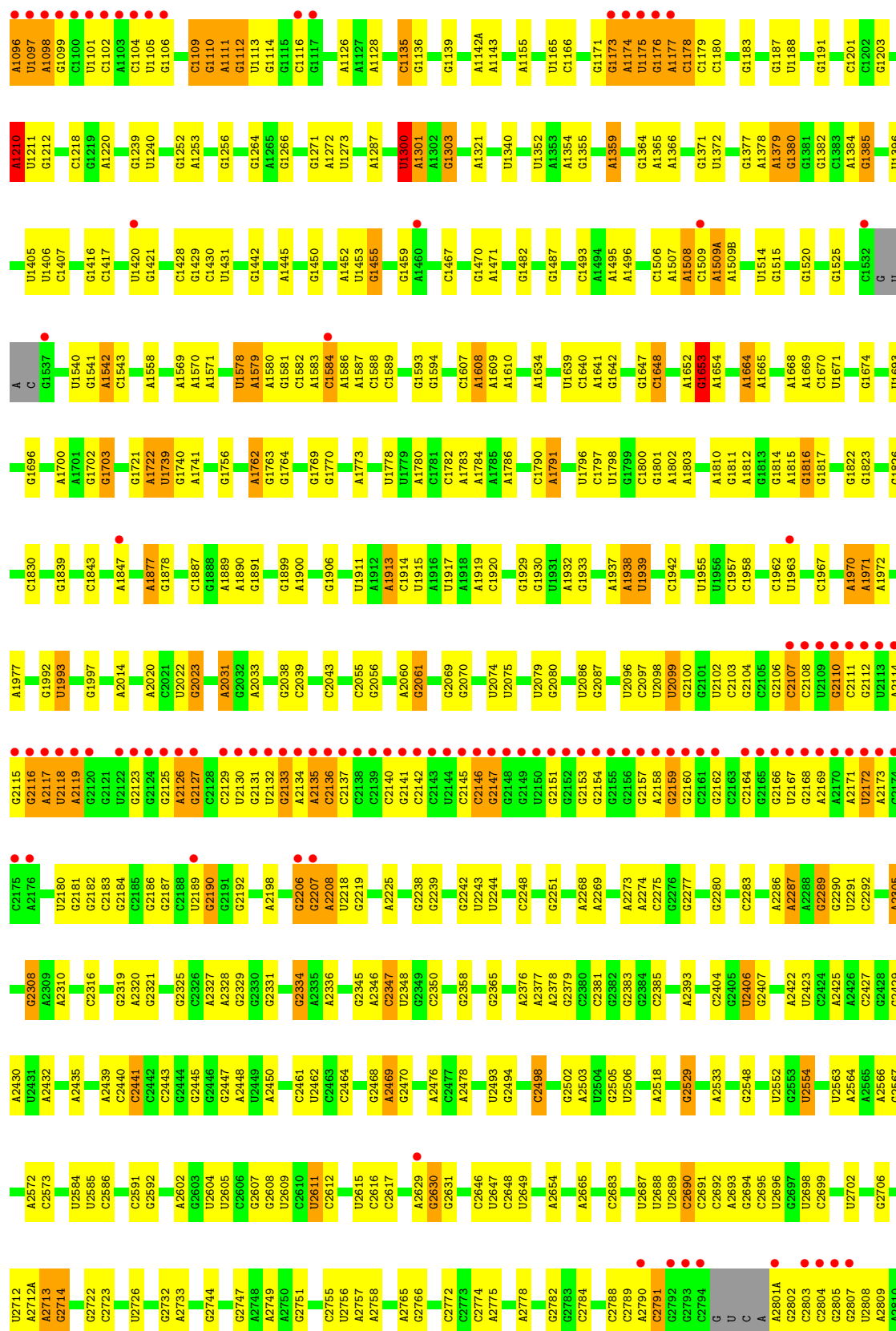
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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
62	2D	51	Total 51	O 51	0	0
62	2E	32	Total 32	O 32	0	0
62	2F	26	Total 26	O 26	0	0
62	2G	7	Total 7	O 7	0	0
62	2H	1	Total 1	O 1	0	0
62	2I	2	Total 2	O 2	0	0
62	2N	5	Total 5	O 5	0	0
62	2O	16	Total 16	O 16	0	0
62	2P	26	Total 26	O 26	0	0
62	2Q	14	Total 14	O 14	0	0
62	2R	17	Total 17	O 17	0	0
62	2S	3	Total 3	O 3	0	0
62	2T	7	Total 7	O 7	0	0
62	2U	15	Total 15	O 15	0	0
62	2V	3	Total 3	O 3	0	0
62	2W	17	Total 17	O 17	0	0
62	2X	8	Total 8	O 8	0	0
62	2Y	2	Total 2	O 2	0	0
62	2Z	6	Total 6	O 6	0	0
62	20	8	Total 8	O 8	0	0
62	21	15	Total 15	O 15	0	0

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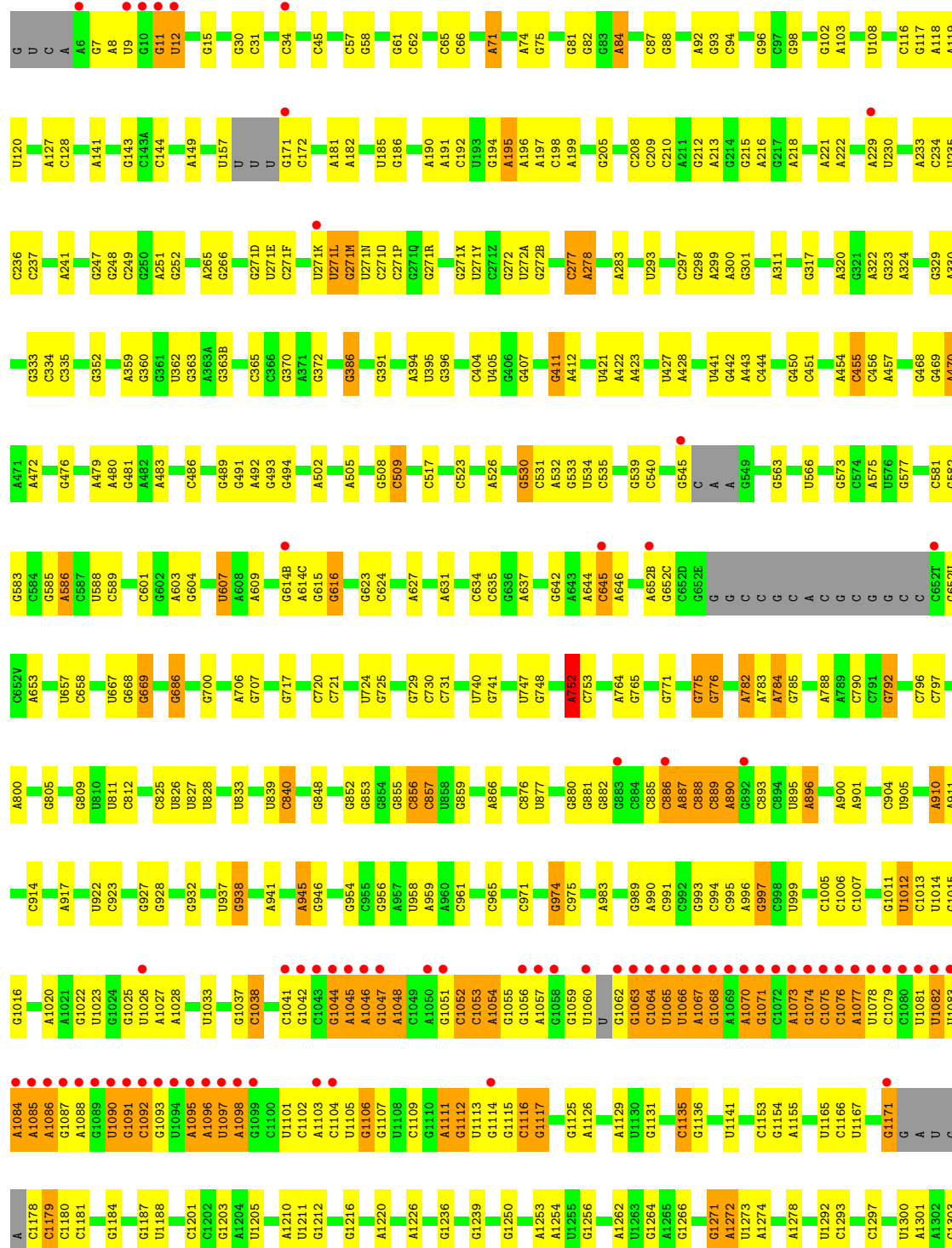
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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
62	23	2	Total 2	O 2	0	0
62	25	11	Total 11	O 11	0	0
62	26	7	Total 7	O 7	0	0
62	27	5	Total 5	O 5	0	0
62	28	16	Total 16	O 16	0	0
62	2a	243	Total 243	O 243	0	0
62	2e	2	Total 2	O 2	0	0
62	2l	3	Total 3	O 3	0	0
62	2m	1	Total 1	O 1	0	0
62	2n	2	Total 2	O 2	0	0
62	2o	2	Total 2	O 2	0	0
62	2p	1	Total 1	O 1	0	0
62	2r	4	Total 4	O 4	0	0
62	2t	2	Total 2	O 2	0	0
62	2w	4	Total 4	O 4	0	0
62	2x	3	Total 3	O 3	0	0

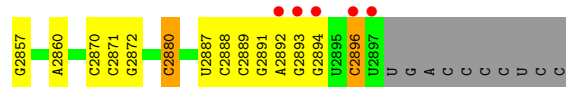




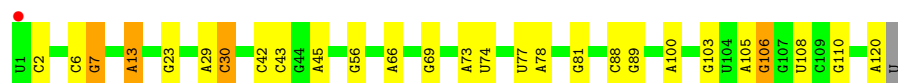
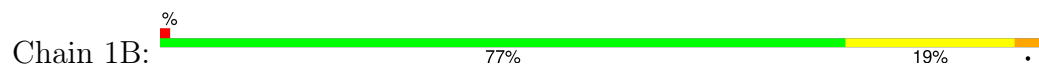
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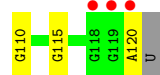
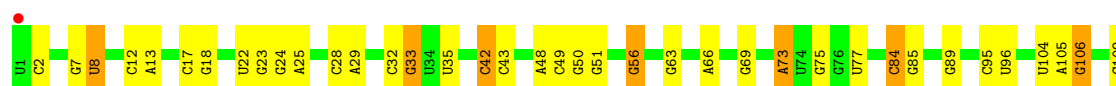
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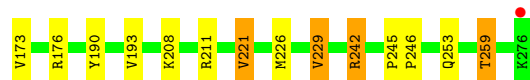
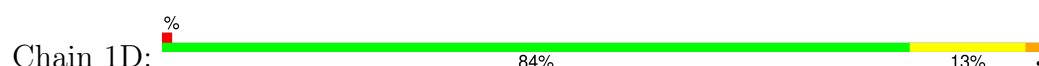
- Molecule 2: 5S Ribosomal RNA



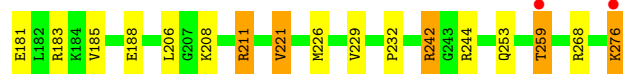
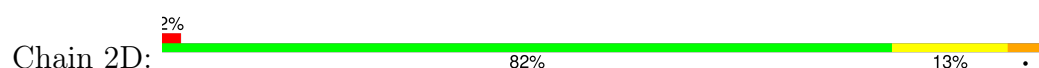
- Molecule 2: 5S Ribosomal RNA



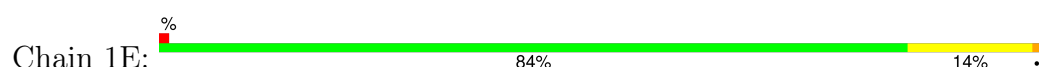
- Molecule 3: 50S ribosomal protein L2



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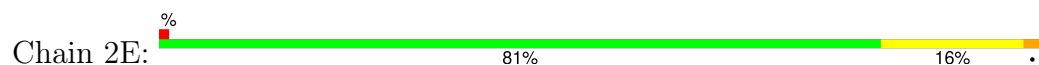


- Molecule 4: 50S ribosomal protein L3

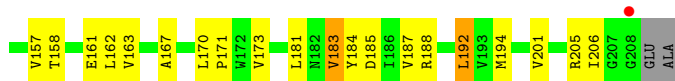
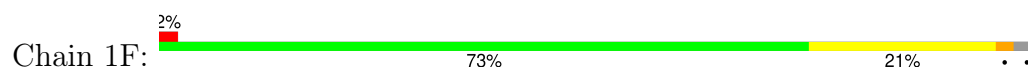




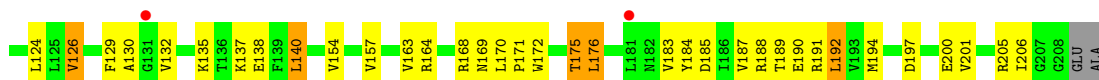
- Molecule 4: 50S ribosomal protein L3



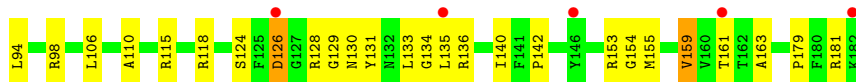
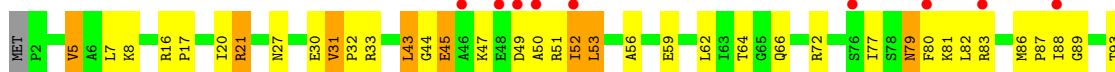
- Molecule 5: 50S ribosomal protein L4



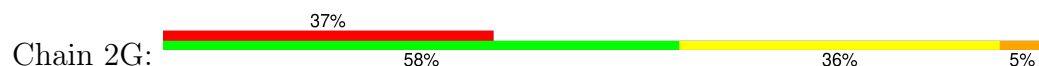
- Molecule 5: 50S ribosomal protein L4

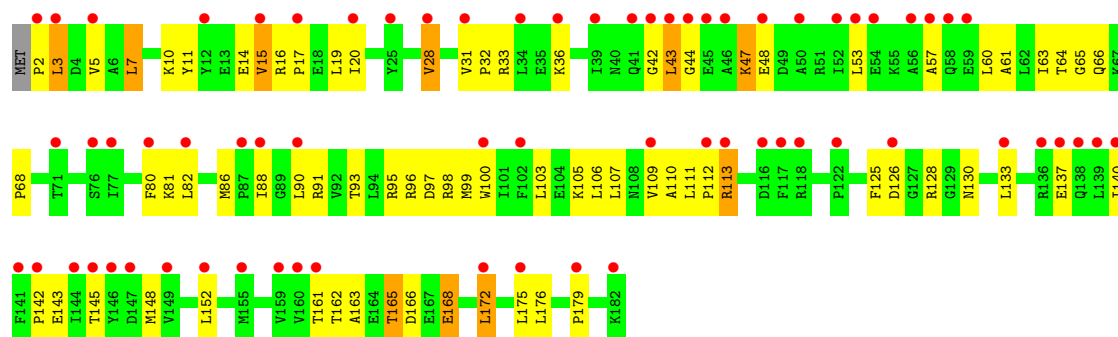


- Molecule 6: 50S ribosomal protein L5

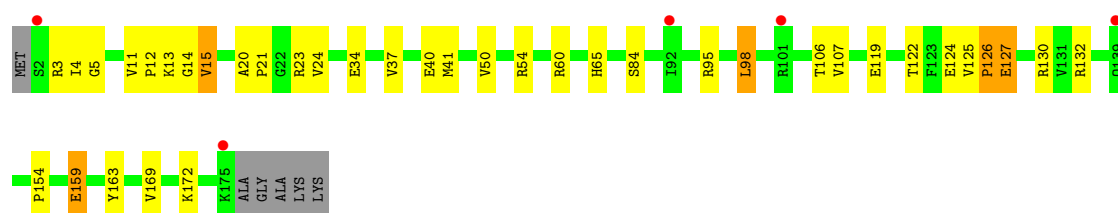
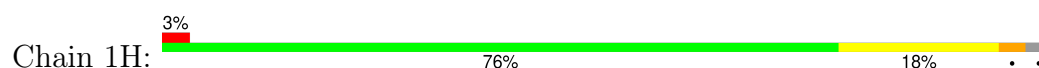


- Molecule 6: 50S ribosomal protein L5

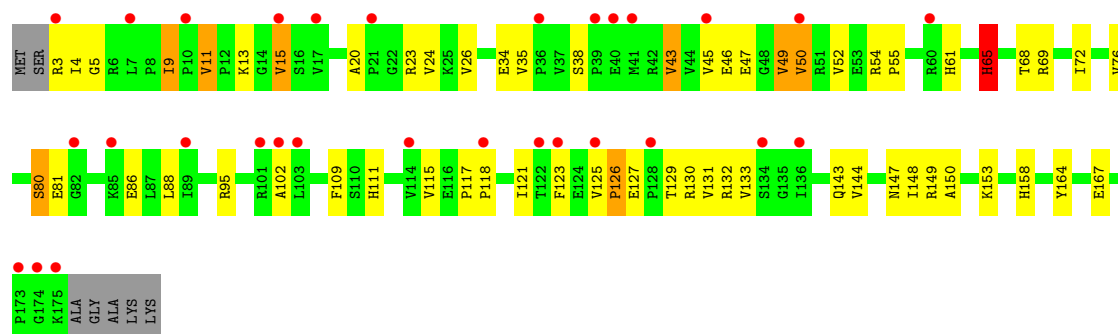




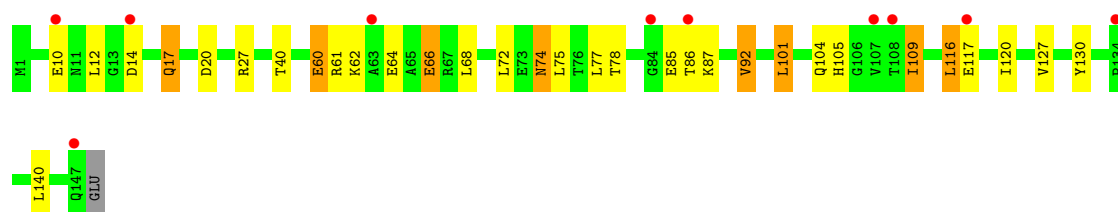
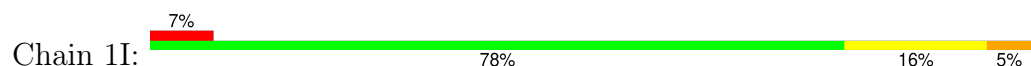
• Molecule 7: 50S ribosomal protein L6



• Molecule 7: 50S ribosomal protein L6

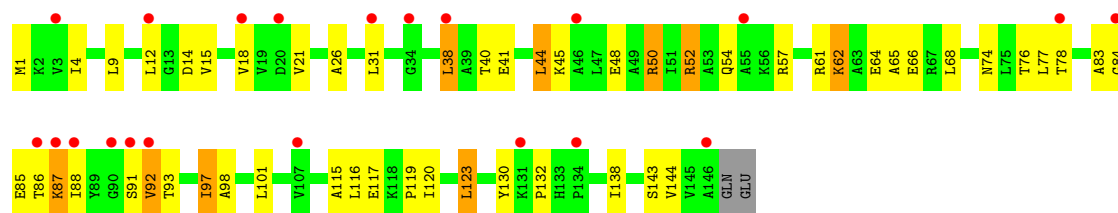


• Molecule 8: 50S ribosomal protein L9

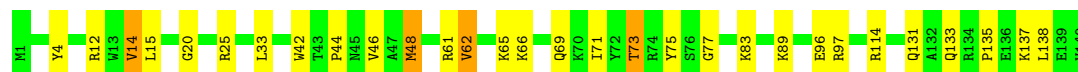
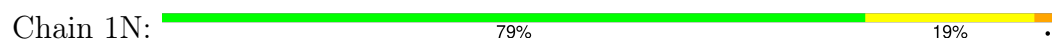


• Molecule 8: 50S ribosomal protein L9

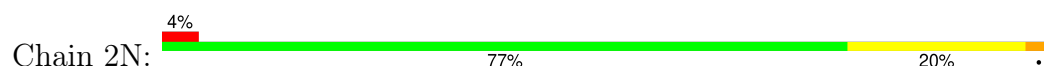




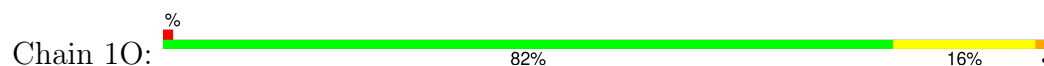
• Molecule 9: 50S ribosomal protein L13



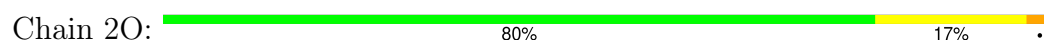
• Molecule 9: 50S ribosomal protein L13



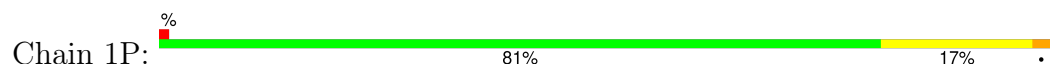
• Molecule 10: 50S ribosomal protein L14



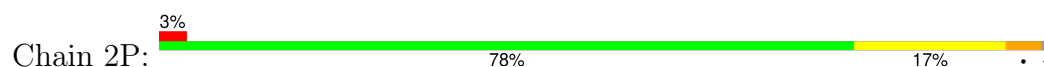
• Molecule 10: 50S ribosomal protein L14



• Molecule 11: 50S ribosomal protein L15

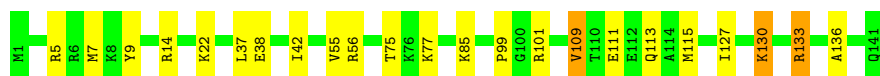


• Molecule 11: 50S ribosomal protein L15



- Molecule 12: 50S ribosomal protein L16

Chain 1Q:  84% 14% .



- Molecule 12: 50S ribosomal protein L16

Chain 2Q:  6% 81% 18% .



- Molecule 13: 50S ribosomal protein L17

Chain 1R:  84% 15% .



- Molecule 13: 50S ribosomal protein L17

Chain 2R:  82% 16% .



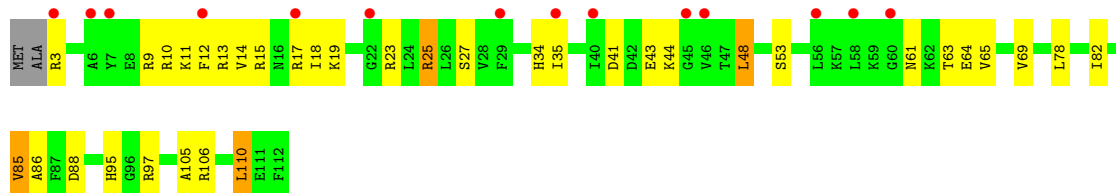
- Molecule 14: 50S ribosomal protein L18

Chain 1S:  79% 18% . .

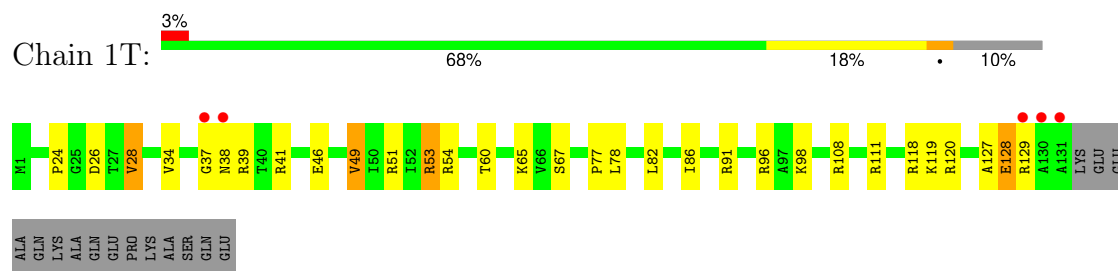


- Molecule 14: 50S ribosomal protein L18

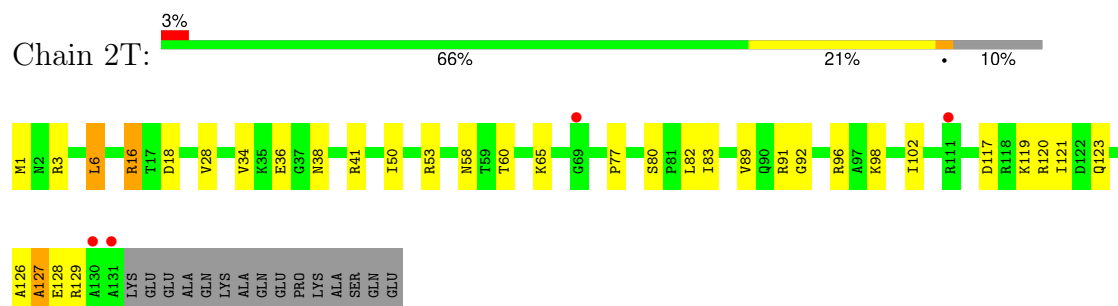
Chain 2S:  12% 66% 29% . .



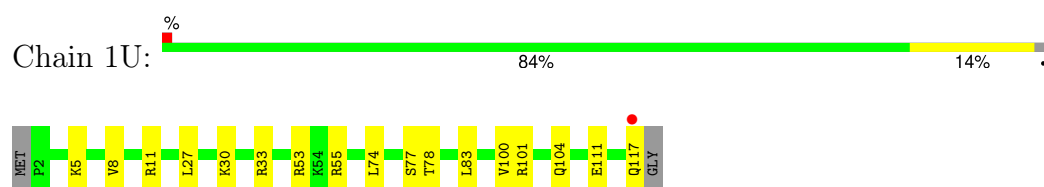
● Molecule 15: 50S ribosomal protein L19



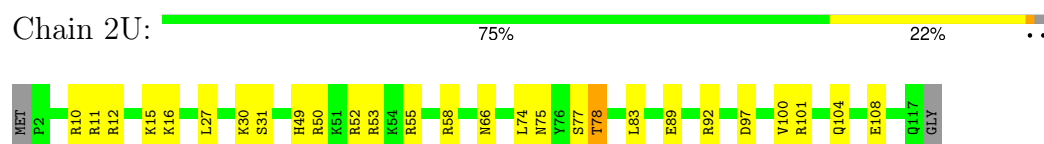
● Molecule 15: 50S ribosomal protein L19



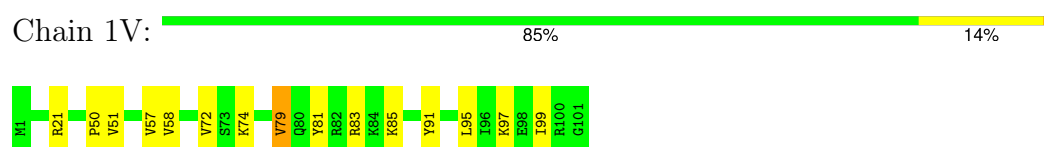
● Molecule 16: 50S ribosomal protein L20



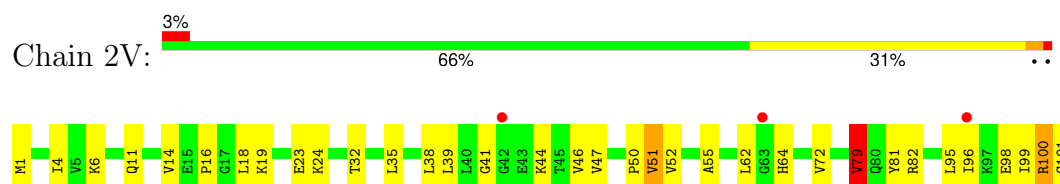
● Molecule 16: 50S ribosomal protein L20



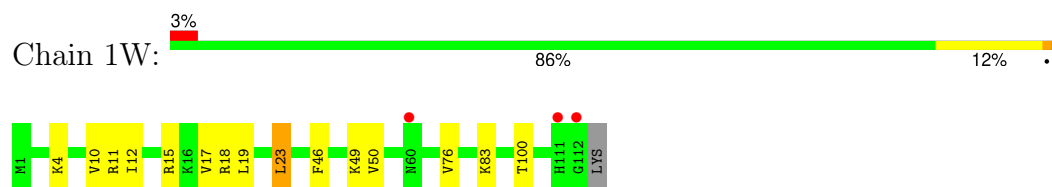
● Molecule 17: 50S ribosomal protein L21



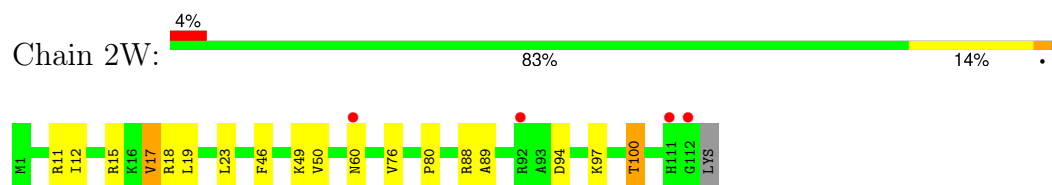
● Molecule 17: 50S ribosomal protein L21



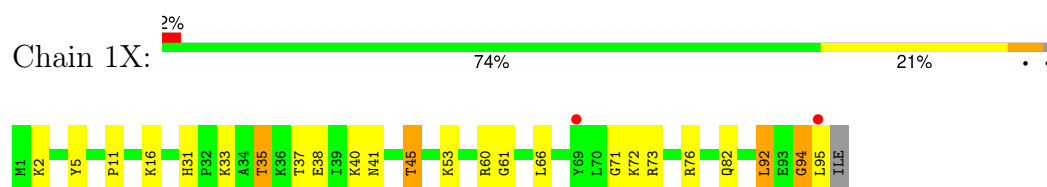
- Molecule 18: 50S ribosomal protein L22



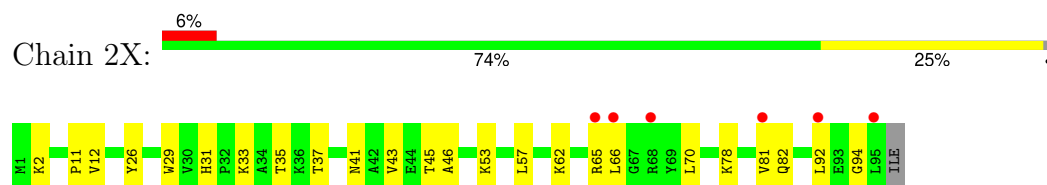
- Molecule 18: 50S ribosomal protein L22



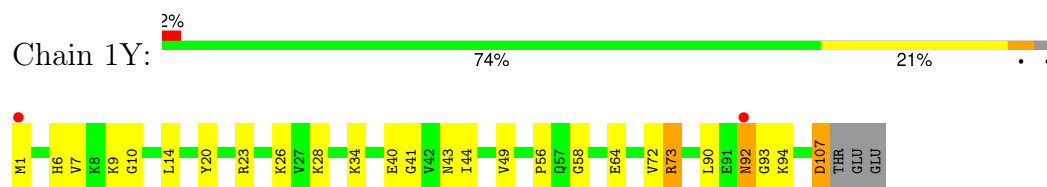
- Molecule 19: 50S ribosomal protein L23



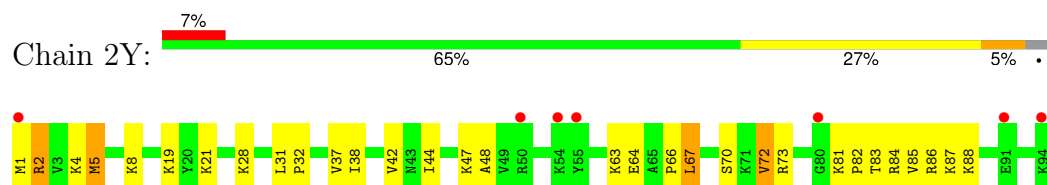
- Molecule 19: 50S ribosomal protein L23



- Molecule 20: 50S ribosomal protein L24

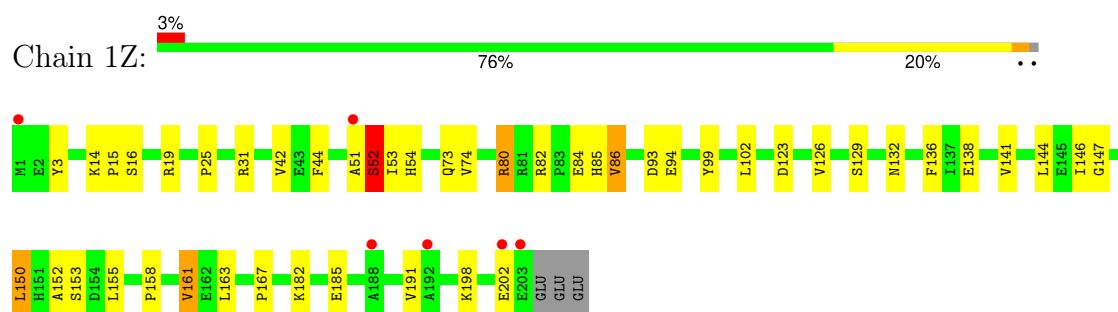


- Molecule 20: 50S ribosomal protein L24

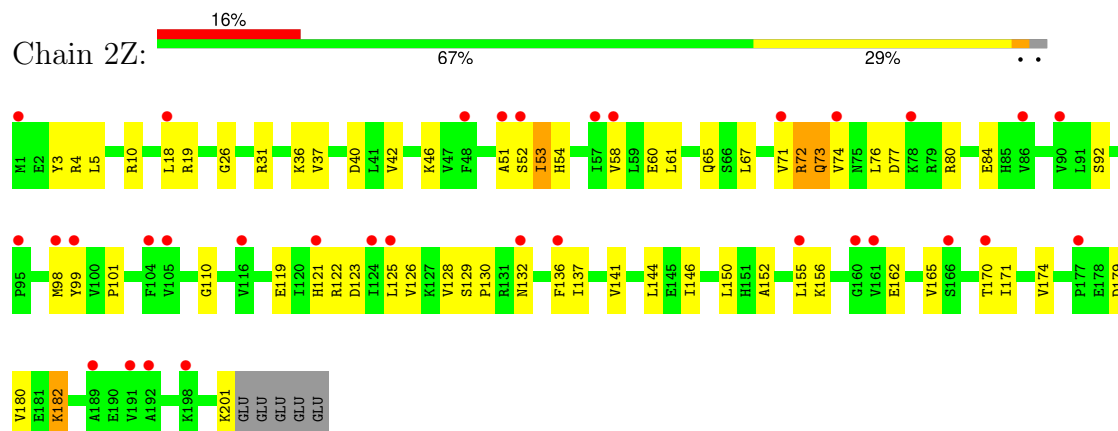


- Molecule 21: 50S ribosomal protein L25

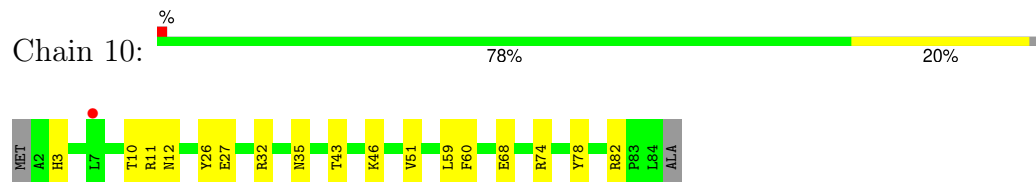




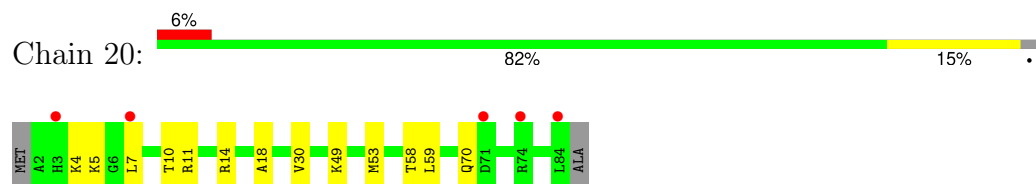
- Molecule 21: 50S ribosomal protein L25



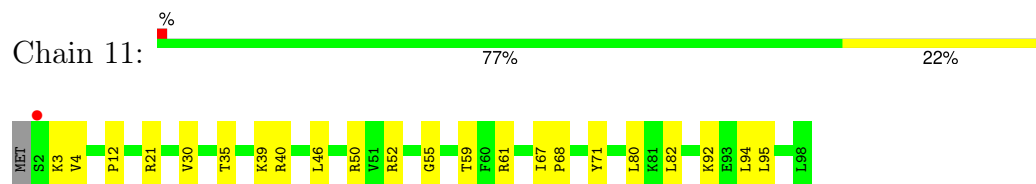
- Molecule 22: 50S ribosomal protein L27



- Molecule 22: 50S ribosomal protein L27



- Molecule 23: 50S ribosomal protein L28

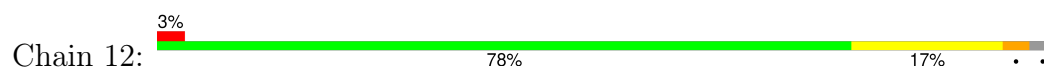


- Molecule 23: 50S ribosomal protein L28





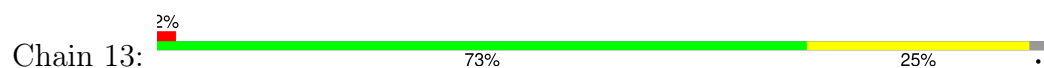
- Molecule 24: 50S ribosomal protein L29



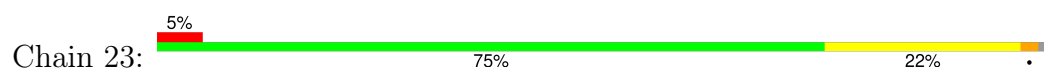
- Molecule 24: 50S ribosomal protein L29



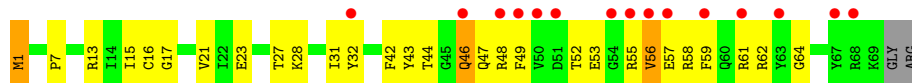
- Molecule 25: 50S ribosomal protein L30



- Molecule 25: 50S ribosomal protein L30

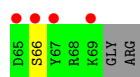


- Molecule 26: 50S ribosomal protein L31

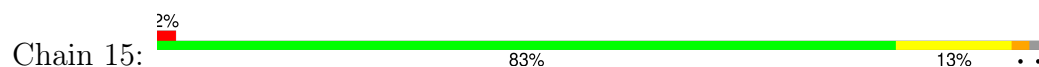


- Molecule 26: 50S ribosomal protein L31

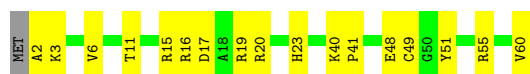




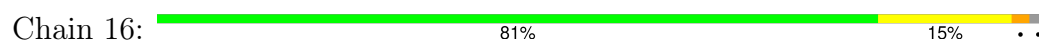
- Molecule 27: 50S ribosomal protein L32



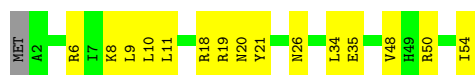
- Molecule 27: 50S ribosomal protein L32



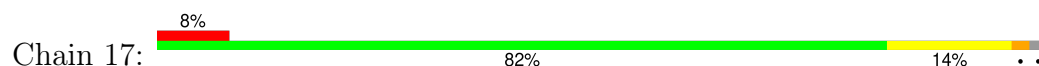
- Molecule 28: 50S ribosomal protein L33



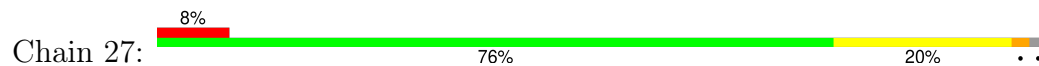
- Molecule 28: 50S ribosomal protein L33



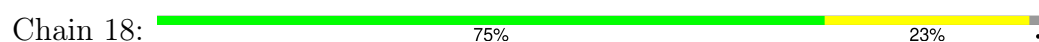
- Molecule 29: 50S ribosomal protein L34

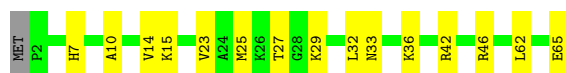


- Molecule 29: 50S ribosomal protein L34

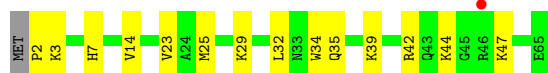
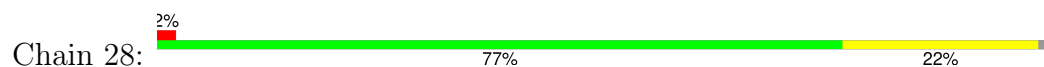


- Molecule 30: 50S ribosomal protein L35





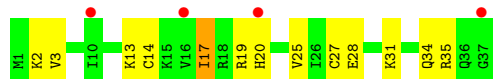
- Molecule 30: 50S ribosomal protein L35



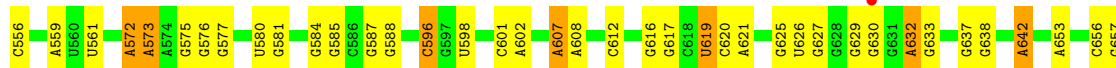
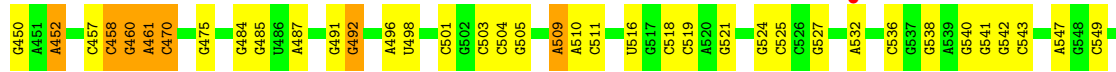
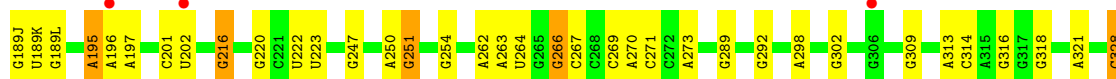
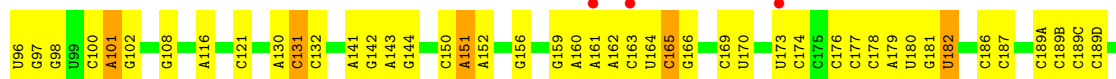
- Molecule 31: 50S ribosomal protein L36

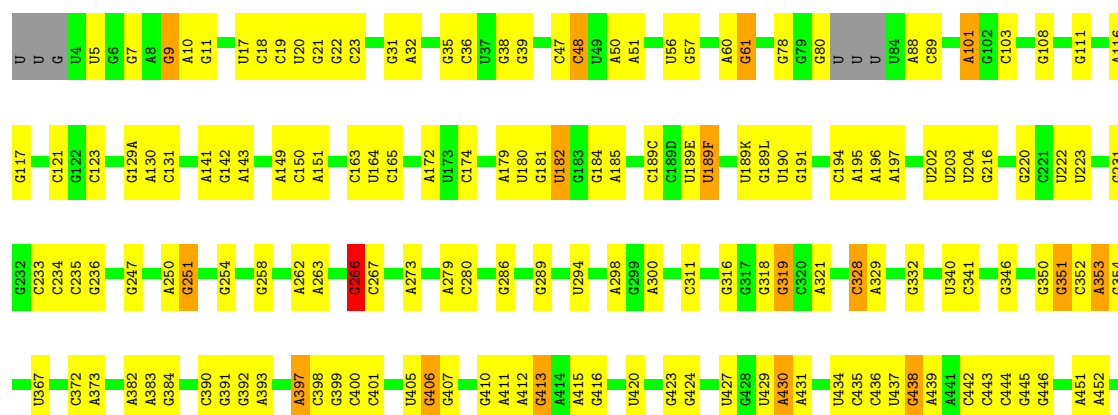
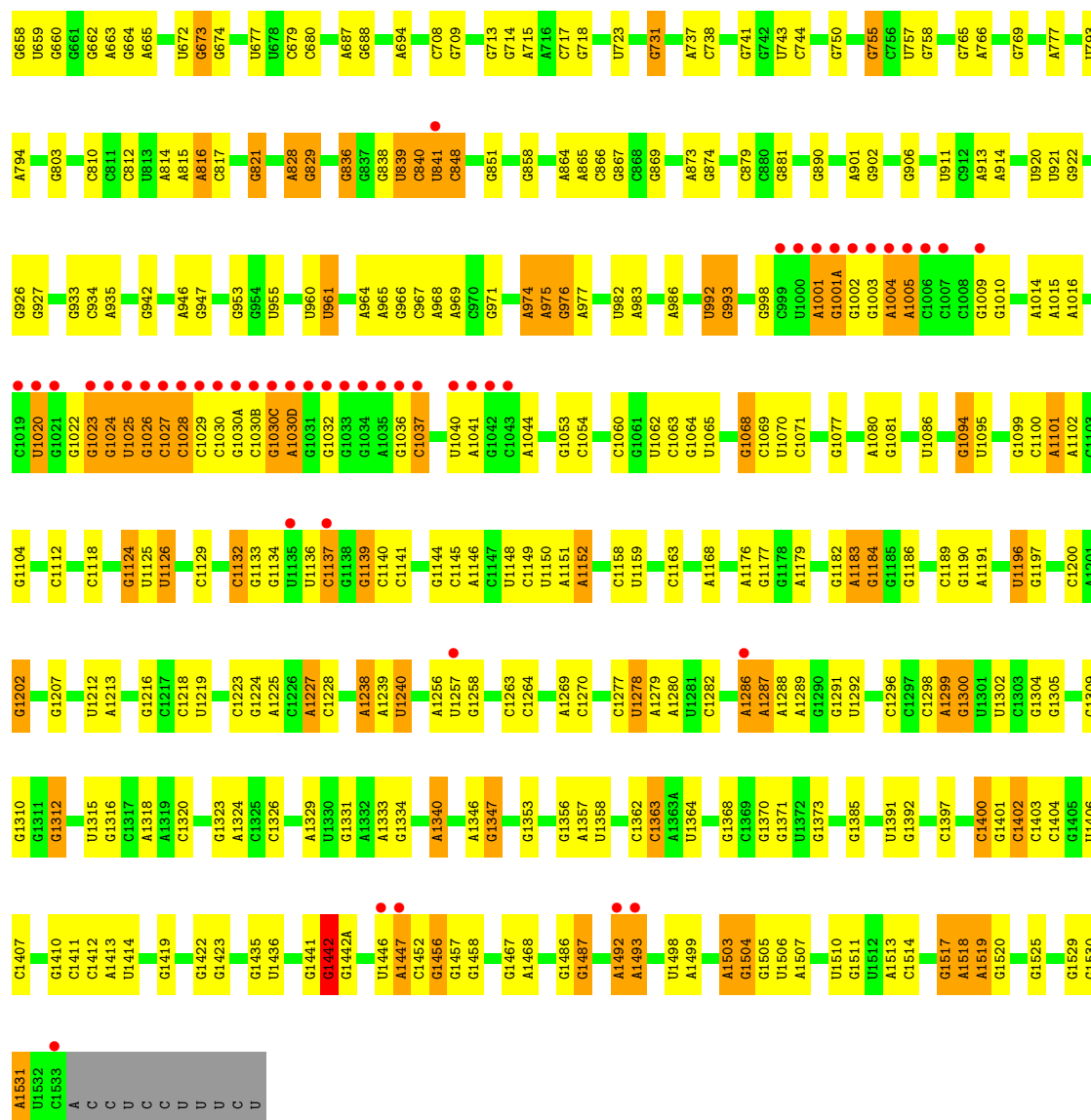


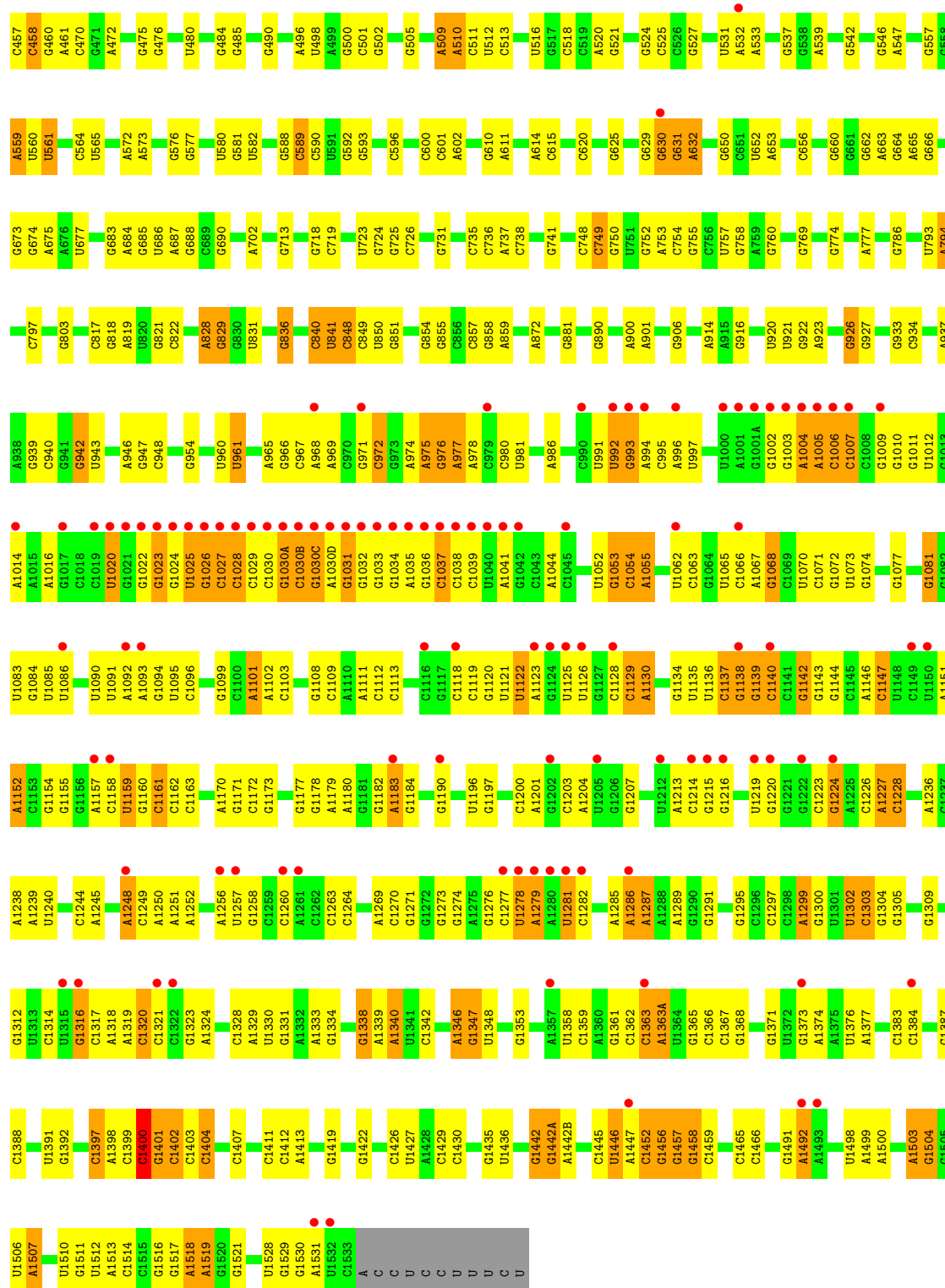
- Molecule 31: 50S ribosomal protein L36



- Molecule 32: 16S Ribosomal RNA

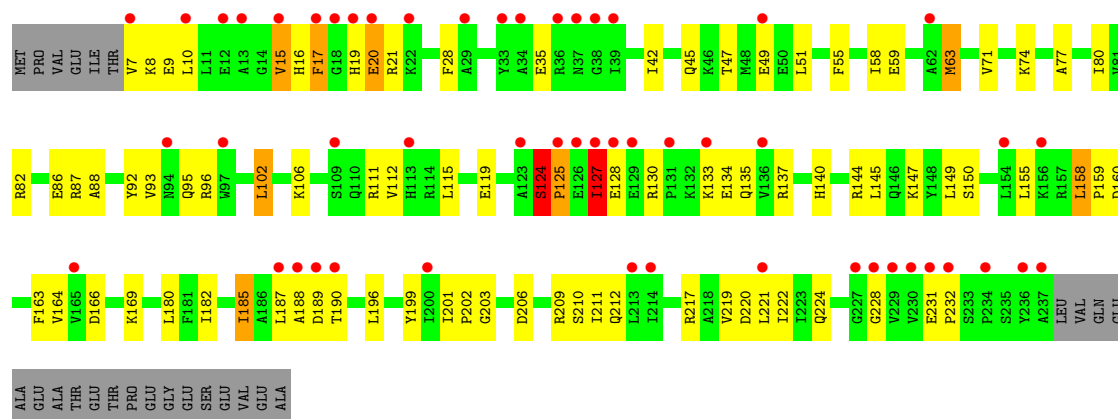




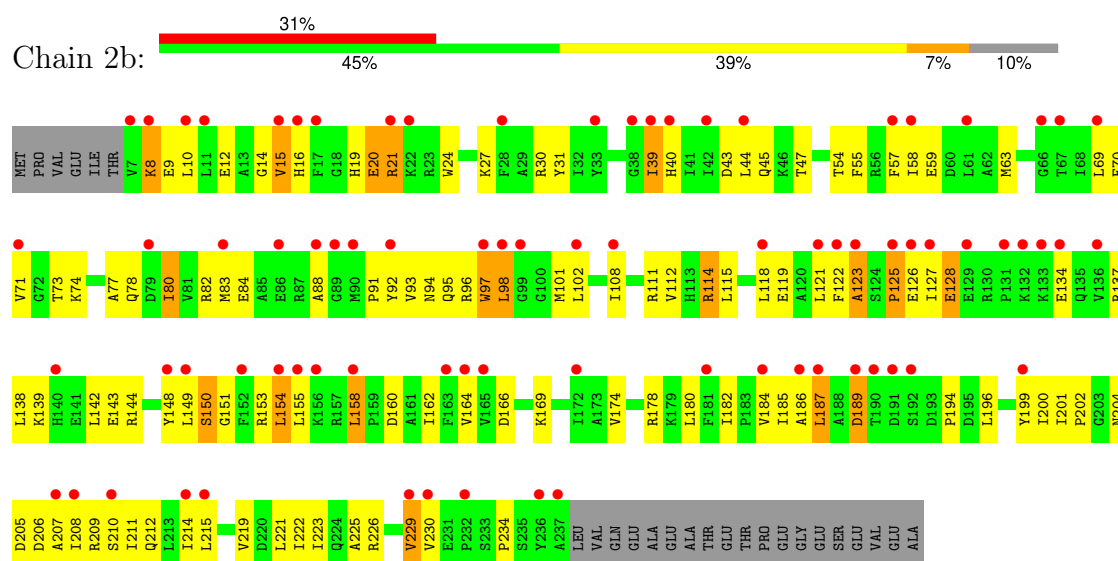


• Molecule 33: 30S ribosomal protein S2

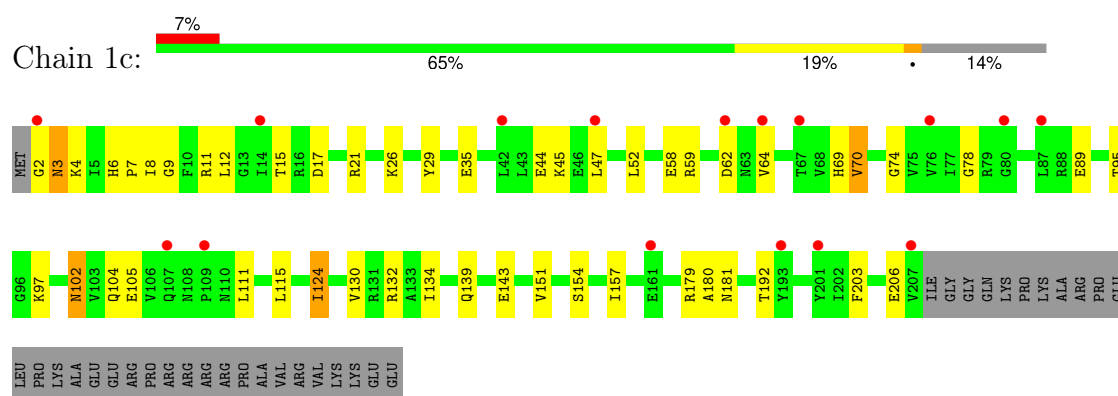




• Molecule 33: 30S ribosomal protein S2

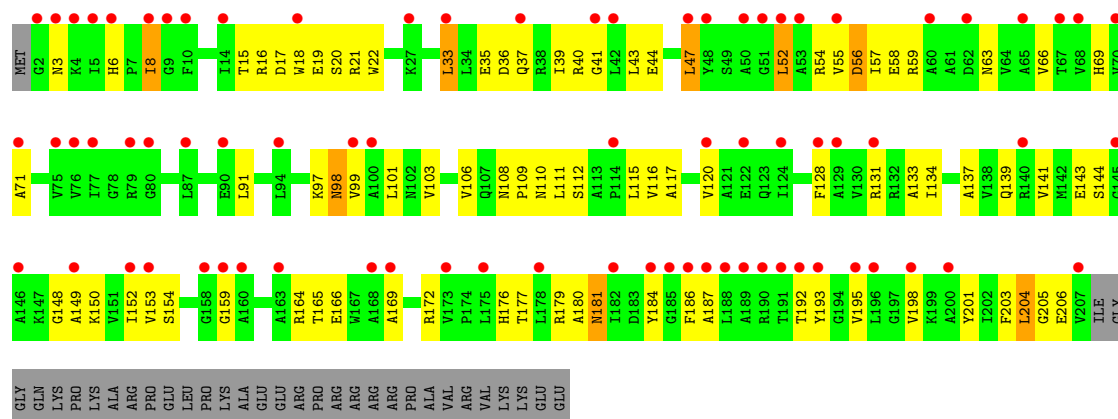


• Molecule 34: 30S ribosomal protein S3

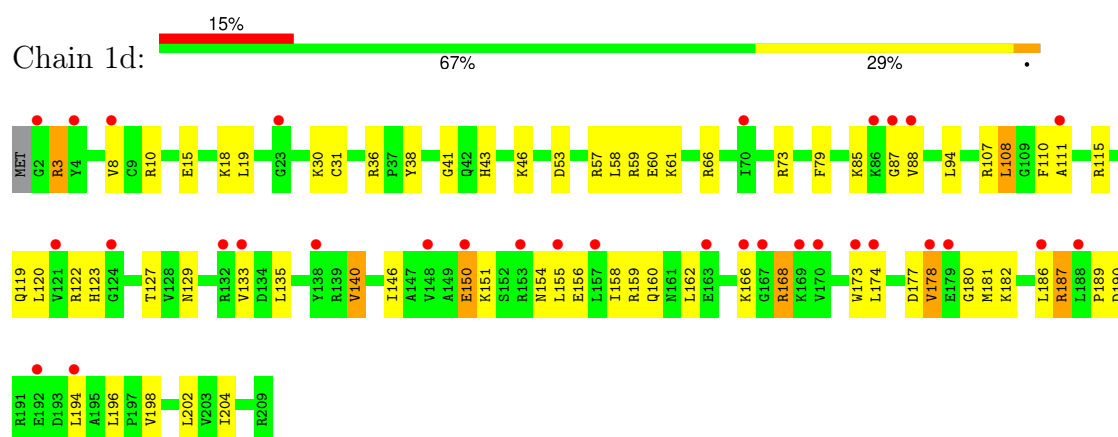


• Molecule 34: 30S ribosomal protein S3

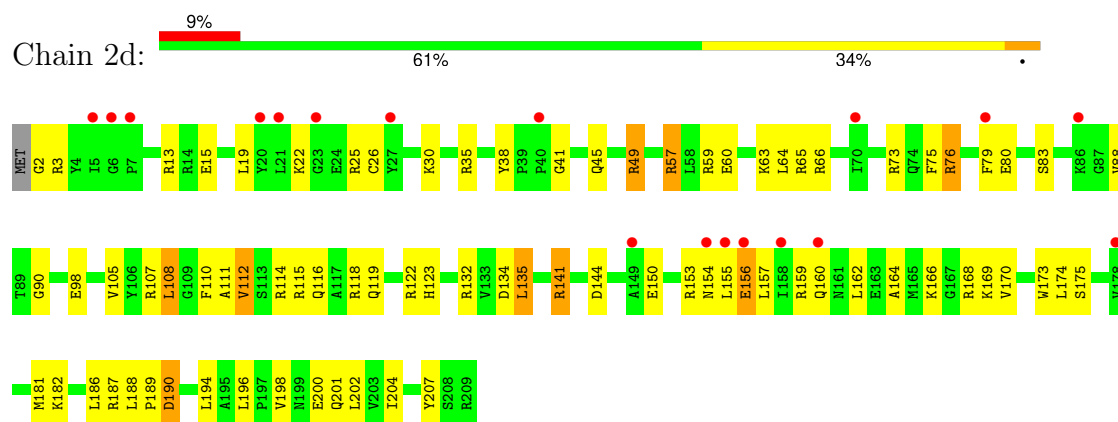




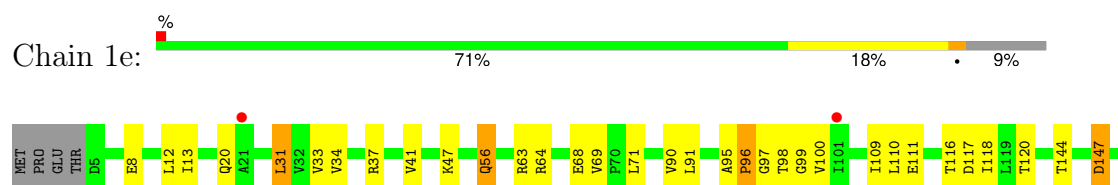
• Molecule 35: 30S ribosomal protein S4

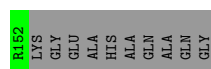


• Molecule 35: 30S ribosomal protein S4

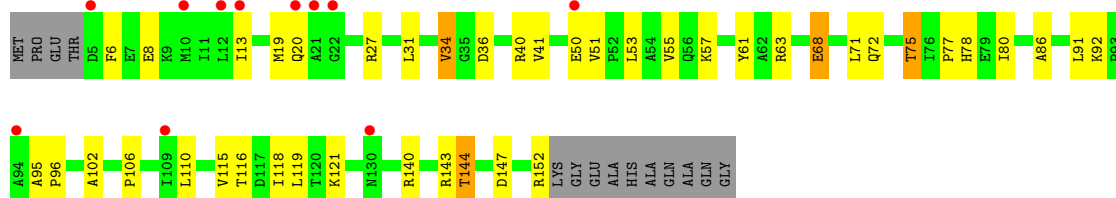


• Molecule 36: 30S ribosomal protein S5

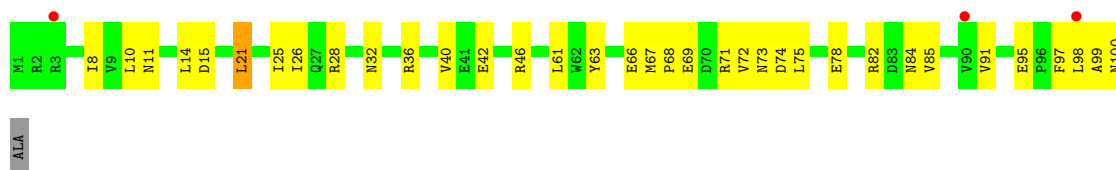




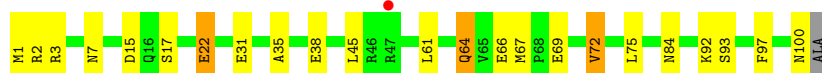
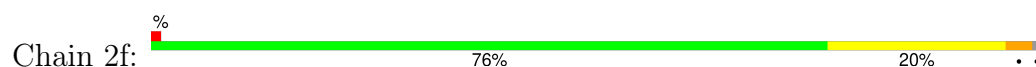
- Molecule 36: 30S ribosomal protein S5



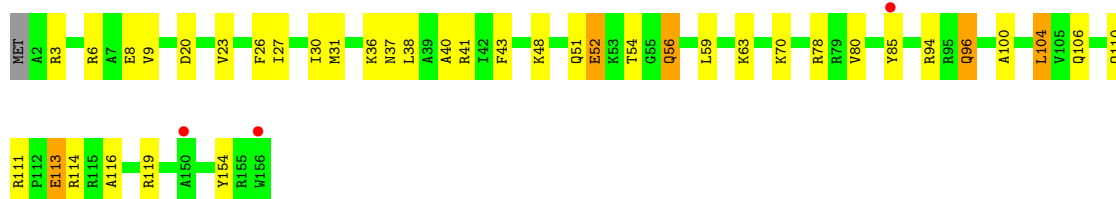
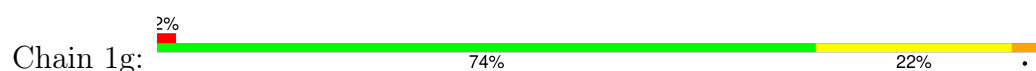
- Molecule 37: 30S ribosomal protein S6



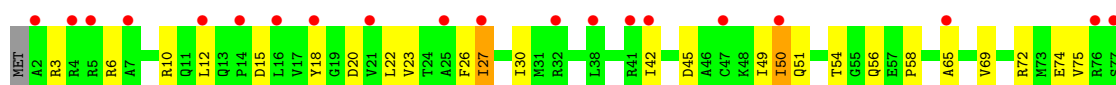
- Molecule 37: 30S ribosomal protein S6



- Molecule 38: 30S ribosomal protein S7

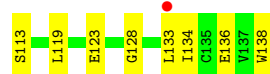
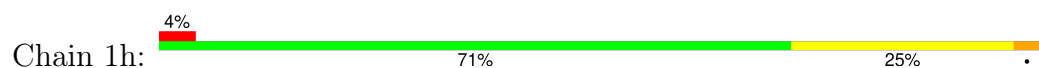


- Molecule 38: 30S ribosomal protein S7

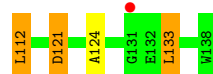
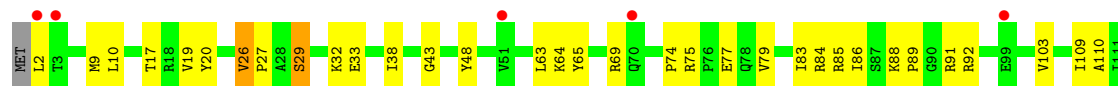
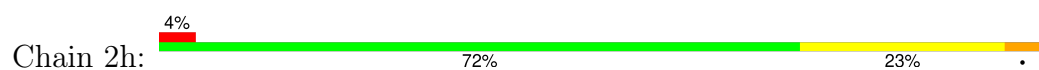




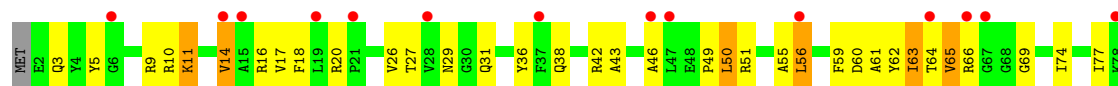
• Molecule 39: 30S ribosomal protein S8



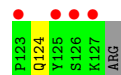
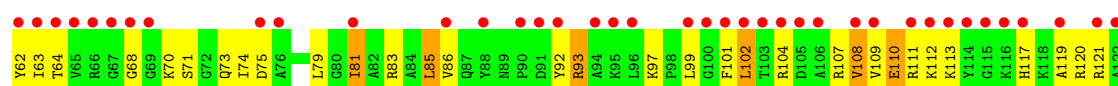
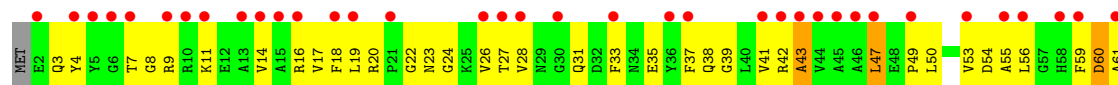
• Molecule 39: 30S ribosomal protein S8



• Molecule 40: 30S ribosomal protein S9



• Molecule 40: 30S ribosomal protein S9



Chain 1j:

Sequence logo for Chain 1j. The y-axis represents information content in bits. The x-axis shows positions 1 to 200. A color scale at the top indicates conservation levels: red (0-27%), green (27-50%), yellow (50-35%), orange (35-7%), and grey (7-8%).

Position	Amino Acid	Information Content (bits)
1	VAL	0.15
2	GLY	0.15
3	GLY	0.15
4	ARG	0.15
5	GLY	0.15
6	GLY	0.15
7	GLY	0.15
8	GLY	0.15
9	GLY	0.15
10	GLY	0.15
11	GLY	0.15
12	GLY	0.15
13	GLY	0.15
14	GLY	0.15
15	GLY	0.15
16	GLY	0.15
17	GLY	0.15
18	GLY	0.15
19	GLY	0.15
20	GLY	0.15
21	GLY	0.15
22	GLY	0.15
23	GLY	0.15
24	GLY	0.15
25	GLY	0.15
26	GLY	0.15
27	GLY	0.15
28	GLY	0.15
29	GLY	0.15
30	GLY	0.15
31	GLY	0.15
32	GLY	0.15
33	GLY	0.15
34	GLY	0.15
35	GLY	0.15
36	GLY	0.15
37	GLY	0.15
38	GLY	0.15
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133	GLY	0.15
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135	GLY	0.15
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137	GLY	0.15
138	GLY	0.15
139	GLY	0.15
140	GLY	0.15
141	GLY	0.15
142	GLY	0.15
143	GLY	0.15</

Chain 2j:

Amino Acid	Segment 1 (65%)	Segment 2 (54%)	Segment 3 (32%)	Segment 4 (5%)	Segment 5 (9%)
MET					
PRO					
LYS					
ILE					
RS					
IG					
K7					
L8					
R9					
G10					
H13					
K14					
T15					
L16					
D17					
A18					
S19					
V24					
E25					
A26					
A27					
R28					
R29					
S30					
G31					
A32					
D33					
V34					
S35					
G36					
P37					
I38					
P39					
L40					
P41					
T42					
R43					
V44					
R45					
R46					
F47					
T48					
V49					
I50					
R51					
G52					
P53					
F54					
K55					
H56					
K57					
D58					
S59					
R60					
E61					
H62					
F63					
F64					

Chain 1k:

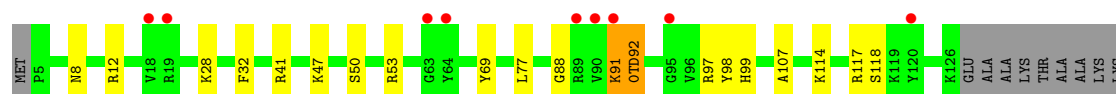
3% 68% 19% 12%

MET ALA LYS LYS PRO SER LYS LYS LYS VAL LYS ARG Q13 Q14 A15 S16 N27 I28 D34 G37 R38 Y50 S53 R54 Q62 L66 A69 K70 M73 A74 Y75 S79 V84 R91 E92 R96 Q99 L103 A104 V105 V109 V114 M17 R120 R126 LYS ALA SER

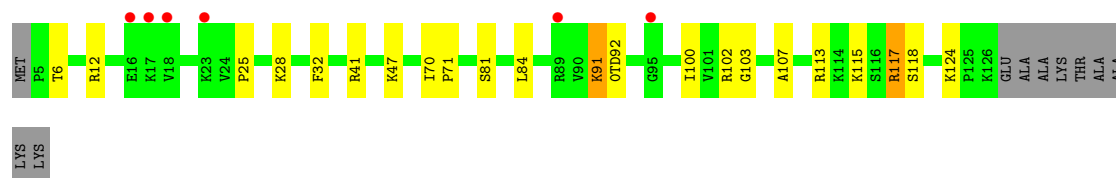
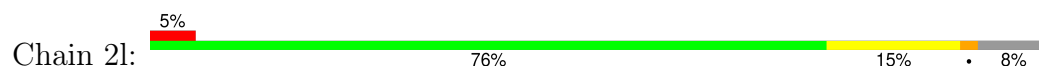
Chain 2k:

Amino Acid	Percentage
LYS	69%
ALA	18%
VAL	5%
MET	12%

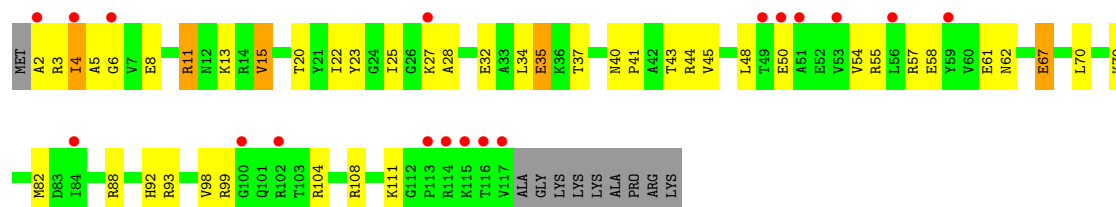
Chain 11: 7% 77% 14% 8%



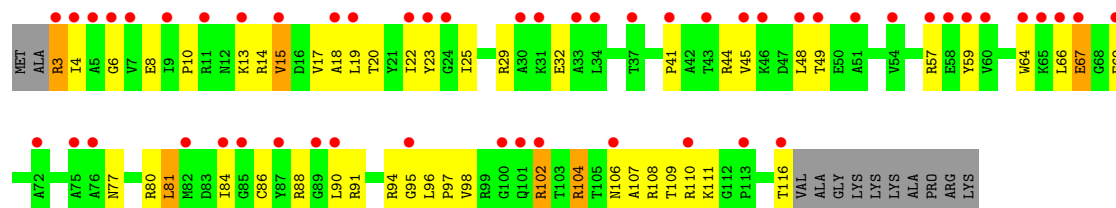
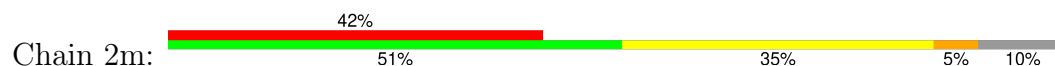
- Molecule 43: 30S ribosomal protein S12



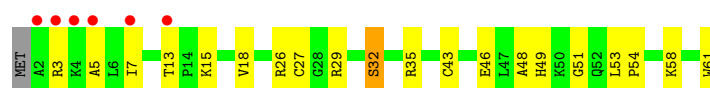
- Molecule 44: 30S ribosomal protein S13



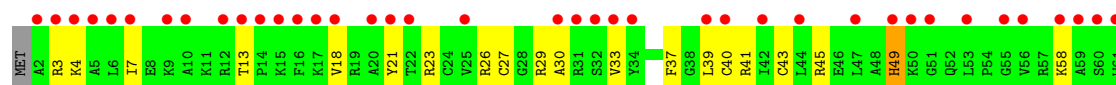
- Molecule 44: 30S ribosomal protein S13



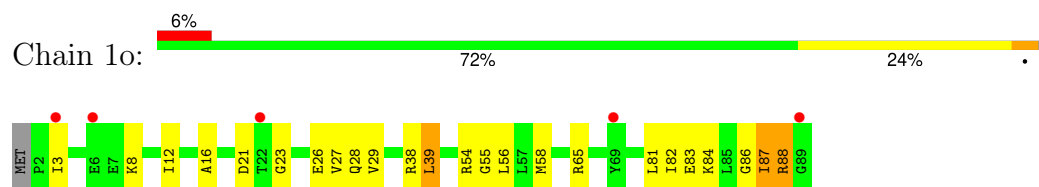
- Molecule 45: 30S ribosomal protein S14 type Z



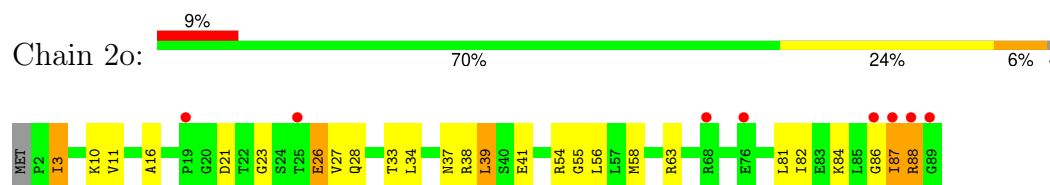
- Molecule 45: 30S ribosomal protein S14 type Z



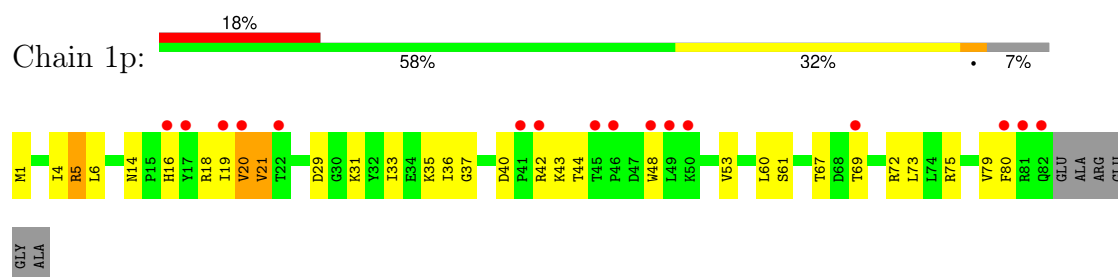
- Molecule 46: 30S ribosomal protein S15



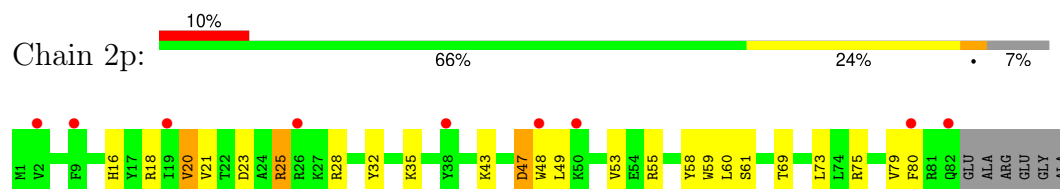
- Molecule 46: 30S ribosomal protein S15



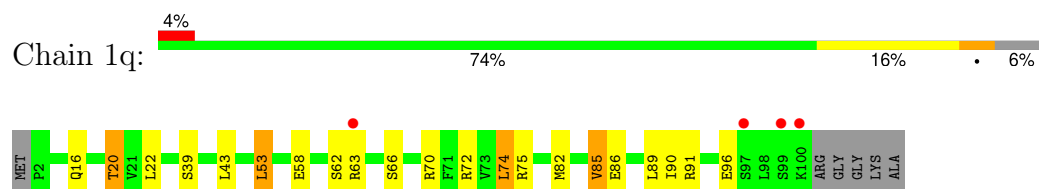
- Molecule 47: 30S ribosomal protein S16



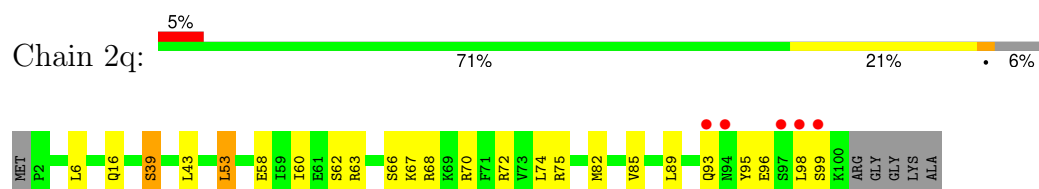
- Molecule 47: 30S ribosomal protein S16



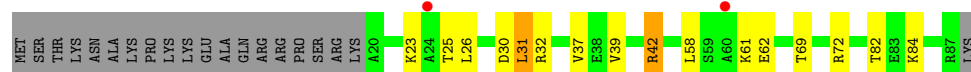
- Molecule 48: 30S ribosomal protein S17



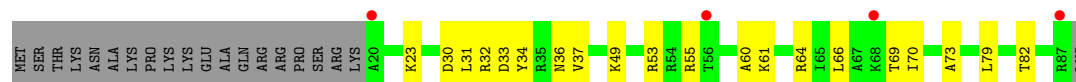
- Molecule 48: 30S ribosomal protein S17



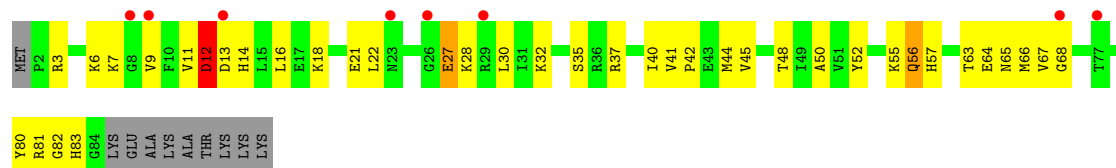
- Molecule 49: 30S ribosomal protein S18



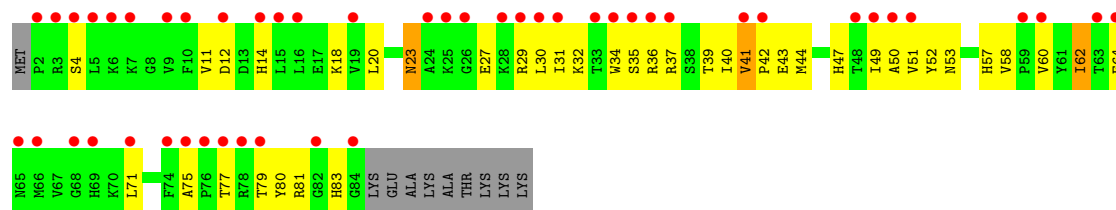
• Molecule 49: 30S ribosomal protein S18



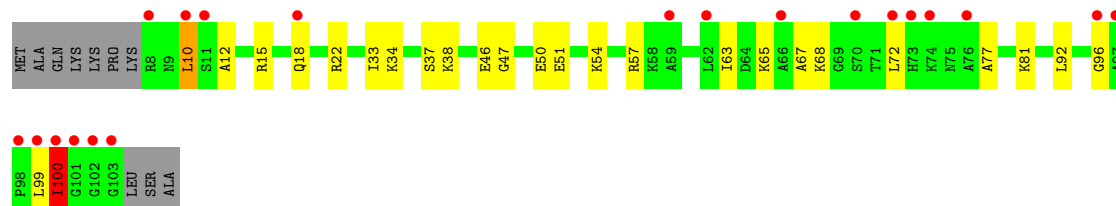
• Molecule 50: 30S ribosomal protein S19



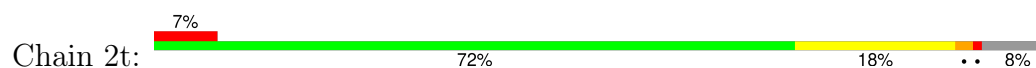
• Molecule 50: 30S ribosomal protein S19



• Molecule 51: 30S ribosomal protein S20

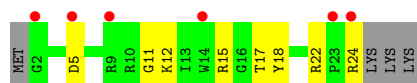


• Molecule 51: 30S ribosomal protein S20





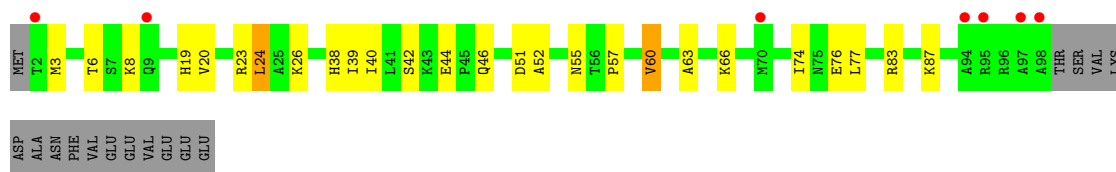
- Molecule 52: 30S ribosomal protein Thx



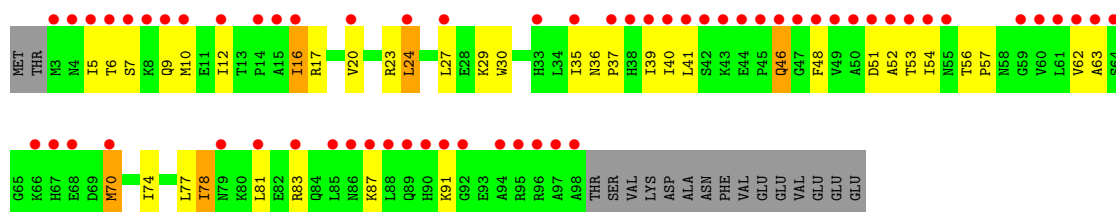
- Molecule 52: 30S ribosomal protein Thx



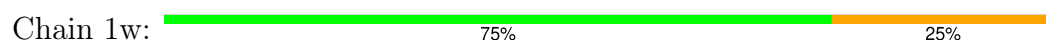
- Molecule 53: Ribosome-associated inhibitor A



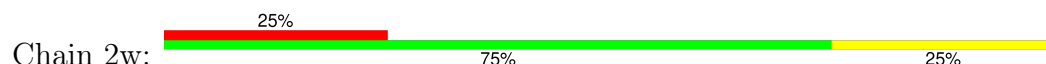
- Molecule 53: Ribosome-associated inhibitor A



- Molecule 54: A-site Aminoacyl-tRNA Analog



- Molecule 54: A-site Aminoacyl-tRNA Analog





- Molecule 55: P-site Peptidyl-tRNA Analog RNA

Chain 1x:
75% 25%

A horizontal bar chart for Chain 1x. The bar is 75% green and 25% yellow.



- Molecule 55: P-site Peptidyl-tRNA Analog RNA

Chain 2x:
25% 75% 25%

A horizontal bar chart for Chain 2x. The bar is 25% red, 75% green, and 25% orange.



- Molecule 56: P-site Peptidyl-tRNA Analog Peptide

Chain 1v:
100%

A horizontal bar chart for Chain 1v. The bar is 100% green.

There are no outlier residues recorded for this chain.

- Molecule 56: P-site Peptidyl-tRNA Analog Peptide

Chain 2v:
100%

A horizontal bar chart for Chain 2v. The bar is 100% green.

There are no outlier residues recorded for this chain.

4 Data and refinement statistics

Property	Value	Source
Space group	P 21 21 21	Depositor
Cell constants a, b, c, α , β , γ	209.61Å 448.45Å 618.45Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	187.31 – 2.45 187.31 – 2.45	Depositor EDS
% Data completeness (in resolution range)	99.9 (187.31-2.45) 99.9 (187.31-2.45)	Depositor EDS
R_{merge}	0.21	Depositor
R_{sym}	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ ¹	1.27 (at 2.45Å)	Xtriage
Refinement program	PHENIX 1.8.2	Depositor
R, R_{free}	0.205 , 0.244 0.207 , 0.246	Depositor DCC
R_{free} test set	105390 reflections (5.01%)	wwPDB-VP
Wilson B-factor (Å ²)	51.3	Xtriage
Anisotropy	0.140	Xtriage
Bulk solvent k_{sol} (e/Å ³), B_{sol} (Å ²)	0.35 , 59.3	EDS
L-test for twinning ²	$\langle L \rangle = 0.45$, $\langle L^2 \rangle = 0.28$	Xtriage
Estimated twinning fraction	No twinning to report.	Xtriage
F_o, F_c correlation	0.93	EDS
Total number of atoms	297350	wwPDB-VP
Average B, all atoms (Å ²)	58.0	wwPDB-VP

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

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, G7M, OMG, ZN, OMC, OMU, 8AN, MG, 0TD, MA6, PSU, PPU, 2MA, M2G, 5MU, 2MG, SF4, UR3, 5MC, MPD

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	0.29	0/69031	0.48	5/107754 (0.0%)
1	2A	0.23	0/68903	0.43	2/107552 (0.0%)
2	1B	0.23	0/2876	0.43	0/4486
2	2B	0.19	0/2878	0.36	0/4490
3	1D	0.29	0/2181	0.54	0/2940
3	2D	0.22	0/2186	0.47	0/2944
4	1E	0.25	0/1592	0.50	0/2149
4	2E	0.21	0/1592	0.45	0/2149
5	1F	0.26	0/1619	0.47	0/2193
5	2F	0.22	0/1615	0.46	0/2188
6	1G	0.20	0/1451	0.46	1/1961 (0.1%)
6	2G	0.22	0/1449	0.42	0/1957
7	1H	0.22	0/1356	0.41	0/1834
7	2H	0.20	0/1350	0.41	0/1826
8	1I	0.20	0/1109	0.44	0/1512
8	2I	0.18	0/1091	0.40	0/1490
9	1N	0.25	0/1148	0.47	0/1547
9	2N	0.19	0/1144	0.40	0/1543
10	1O	0.27	0/943	0.48	0/1269
10	2O	0.22	0/943	0.44	0/1269
11	1P	0.26	0/1152	0.54	0/1533
11	2P	0.21	0/1152	0.51	0/1533
12	1Q	0.29	0/1143	0.53	0/1527
12	2Q	0.21	0/1143	0.42	0/1527
13	1R	0.28	0/982	0.48	0/1312
13	2R	0.22	0/982	0.45	0/1312
14	1S	0.23	0/887	0.54	1/1180 (0.1%)
14	2S	0.20	0/880	0.44	0/1172
15	1T	0.25	0/1105	0.48	0/1477
15	2T	0.22	0/1097	0.45	0/1468
16	1U	0.29	0/977	0.50	0/1301

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
16	2U	0.20	0/977	0.41	0/1301
17	1V	0.25	0/786	0.47	0/1053
17	2V	0.19	0/782	0.43	0/1049
18	1W	0.28	0/897	0.49	0/1205
18	2W	0.22	0/897	0.43	0/1205
19	1X	0.27	0/764	0.49	0/1025
19	2X	0.21	0/764	0.42	0/1025
20	1Y	0.24	0/823	0.53	0/1099
20	2Y	0.19	0/823	0.47	0/1100
21	1Z	0.21	0/1620	0.44	0/2200
21	2Z	0.19	0/1590	0.40	0/2162
22	10	0.27	0/662	0.50	0/881
22	20	0.22	0/659	0.48	0/877
23	11	0.25	0/761	0.44	0/1013
23	21	0.24	0/766	0.43	0/1018
24	12	0.23	0/590	0.47	0/781
24	22	0.21	0/594	0.45	0/785
25	13	0.24	0/474	0.47	0/635
25	23	0.19	0/469	0.40	0/630
26	14	0.25	0/559	0.59	0/754
26	24	0.27	0/549	0.65	2/741 (0.3%)
27	15	0.26	0/473	0.52	0/639
27	25	0.21	0/469	0.45	0/635
28	16	0.25	0/460	0.49	0/613
28	26	0.21	0/456	0.42	0/608
29	17	0.29	0/426	0.55	0/561
29	27	0.24	0/426	0.51	0/561
30	18	0.26	0/525	0.50	0/691
30	28	0.21	0/525	0.43	0/691
31	19	0.29	0/310	0.48	0/407
31	29	0.21	0/310	0.41	0/407
32	1a	0.20	0/35795	0.39	1/55864 (0.0%)
32	2a	0.20	1/35890 (0.0%)	0.38	2/56012 (0.0%)
33	1b	0.22	0/1876	0.47	0/2533
33	2b	0.23	0/1860	0.48	0/2518
34	1c	0.19	0/1582	0.39	0/2137
34	2c	0.21	0/1566	0.39	0/2119
35	1d	0.20	0/1695	0.44	0/2274
35	2d	0.19	0/1698	0.42	0/2277
36	1e	0.21	0/1149	0.43	0/1548
36	2e	0.21	0/1149	0.42	0/1548
37	1f	0.20	0/827	0.38	0/1120
37	2f	0.19	0/829	0.40	0/1123

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z >5	RMSZ	# Z >5
38	1g	0.18	0/1254	0.38	0/1683
38	2g	0.19	0/1248	0.36	0/1676
39	1h	0.19	0/1118	0.41	0/1506
39	2h	0.17	0/1108	0.40	0/1494
40	1i	0.20	0/1005	0.45	0/1351
40	2i	0.21	0/985	0.45	0/1329
41	1j	0.20	0/732	0.38	0/993
41	2j	0.22	0/723	0.49	0/984
42	1k	0.19	0/849	0.43	0/1150
42	2k	0.21	0/848	0.49	2/1149 (0.2%)
43	1l	0.20	0/937	0.44	0/1260
43	2l	0.20	0/937	0.43	0/1260
44	1m	0.20	0/924	0.45	0/1242
44	2m	0.23	0/905	0.48	0/1217
45	1n	0.20	0/501	0.41	0/664
45	2n	0.21	0/501	0.40	0/664
46	1o	0.21	0/739	0.39	0/985
46	2o	0.18	0/739	0.38	0/985
47	1p	0.21	0/697	0.50	0/939
47	2p	0.20	0/693	0.44	0/935
48	1q	0.19	0/836	0.40	0/1117
48	2q	0.18	0/836	0.37	0/1117
49	1r	0.19	0/560	0.43	0/746
49	2r	0.19	0/560	0.40	0/746
50	1s	0.18	0/663	0.42	0/895
50	2s	0.20	0/660	0.45	0/893
51	1t	0.20	0/734	0.45	0/969
51	2t	0.19	0/736	0.42	0/976
52	1u	0.20	0/203	0.43	0/266
52	2u	0.19	0/203	0.44	0/266
53	1y	0.18	0/776	0.40	0/1048
53	2y	0.19	0/761	0.36	0/1030
54	1w	0.30	0/44	0.39	0/67
54	2w	0.25	0/44	0.35	0/67
55	1x	0.38	0/44	0.57	0/67
55	2x	0.29	0/44	0.53	0/67
56	1v	0.18	0/20	0.32	0/25
56	2v	0.22	0/20	0.33	0/25
All	All	0.23	1/310246 (0.0%)	0.44	16/463673 (0.0%)

All (1) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
32	2a	527	G7M	O3'-P	5.04	1.61	1.56

The worst 5 of 16 bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
14	1S	60	GLY	N-CA-C	7.93	117.42	110.21
1	2A	1992	G	C2'-C3'-O3'	6.61	119.41	109.50
6	1G	89	GLY	N-CA-C	6.59	118.84	111.85
42	2k	116	HIS	CA-C-N	6.38	133.19	121.70
42	2k	116	HIS	C-N-CA	6.38	133.19	121.70

There are no chirality outliers.

There are no planarity outliers.

5.2 Too-close contacts [i](#)

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	1A	61869	0	31206	510	0
1	2A	61758	0	31154	637	0
2	1B	2572	0	1305	16	0
2	2B	2573	0	1306	31	0
3	1D	2131	0	2207	32	0
3	2D	2136	0	2218	31	0
4	1E	1559	0	1618	16	0
4	2E	1559	0	1618	20	0
5	1F	1584	0	1625	31	0
5	2F	1580	0	1619	49	0
6	1G	1426	0	1445	47	0
6	2G	1424	0	1441	47	0
7	1H	1330	0	1407	24	0
7	2H	1324	0	1402	41	0
8	1I	1094	0	1127	15	0
8	2I	1076	0	1094	27	0
9	1N	1121	0	1195	15	0
9	2N	1117	0	1184	21	0
10	1O	933	0	996	19	0
10	2O	933	0	996	15	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
11	1P	1135	0	1212	24	0
11	2P	1135	0	1212	27	0
12	1Q	1122	0	1177	15	0
12	2Q	1122	0	1177	17	0
13	1R	968	0	1033	13	0
13	2R	968	0	1033	16	0
14	1S	877	0	938	13	0
14	2S	870	0	923	25	0
15	1T	1091	0	1151	19	0
15	2T	1083	0	1136	22	0
16	1U	959	0	1019	12	0
16	2U	959	0	1019	17	0
17	1V	775	0	841	8	0
17	2V	771	0	830	19	0
18	1W	886	0	940	8	0
18	2W	886	0	940	10	0
19	1X	750	0	814	20	0
19	2X	750	0	814	18	0
20	1Y	810	0	892	15	0
20	2Y	810	0	887	22	0
21	1Z	1587	0	1598	26	0
21	2Z	1557	0	1564	40	0
22	10	653	0	674	12	0
22	20	650	0	665	7	0
23	11	754	0	823	15	0
23	21	759	0	837	20	0
24	12	588	0	643	8	0
24	22	592	0	654	12	0
25	13	469	0	518	10	0
25	23	464	0	514	9	0
26	14	546	0	522	26	0
26	24	536	0	514	30	0
27	15	459	0	476	5	0
27	25	455	0	465	12	0
28	16	453	0	473	3	0
28	26	449	0	469	10	0
29	17	418	0	467	7	0
29	27	418	0	467	7	0
30	18	517	0	582	10	0
30	28	517	0	582	9	0
31	19	307	0	335	1	0
31	29	307	0	335	9	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
32	1a	32246	0	16293	336	0
32	2a	32331	0	16338	412	0
33	1b	1842	0	1862	52	0
33	2b	1825	0	1828	80	0
34	1c	1558	0	1557	29	0
34	2c	1542	0	1517	64	0
35	1d	1665	0	1687	46	0
35	2d	1668	0	1703	57	0
36	1e	1133	0	1191	18	0
36	2e	1133	0	1191	29	0
37	1f	814	0	808	22	0
37	2f	816	0	808	14	0
38	1g	1235	0	1249	27	0
38	2g	1229	0	1238	29	0
39	1h	1098	0	1143	25	0
39	2h	1088	0	1126	20	0
40	1i	986	0	990	40	0
40	2i	966	0	953	51	0
41	1j	719	0	672	28	0
41	2j	710	0	661	26	0
42	1k	834	0	838	14	0
42	2k	833	0	836	12	0
43	1l	932	0	981	16	0
43	2l	932	0	981	15	0
44	1m	914	0	954	27	0
44	2m	895	0	920	33	0
45	1n	492	0	528	17	0
45	2n	492	0	529	14	0
46	1o	728	0	760	19	0
46	2o	728	0	760	13	0
47	1p	681	0	697	16	0
47	2p	677	0	686	17	0
48	1q	823	0	891	16	0
48	2q	823	0	891	17	0
49	1r	555	0	618	15	0
49	2r	555	0	618	11	0
50	1s	648	0	658	23	0
50	2s	645	0	635	33	0
51	1t	732	0	809	17	0
51	2t	733	0	795	14	0
52	1u	199	0	208	4	0
52	2u	199	0	208	2	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
53	1y	764	0	786	19	0
53	2y	749	0	757	27	0
54	1w	78	0	50	2	0
54	2w	78	0	50	0	0
55	1x	63	0	33	0	0
55	2x	63	0	33	1	0
56	1v	21	0	27	0	0
56	2v	21	0	27	0	0
57	10	8	0	0	0	0
57	11	5	0	0	0	0
57	13	3	0	0	0	0
57	15	7	0	0	0	0
57	17	6	0	0	0	0
57	18	3	0	0	0	0
57	19	1	0	0	0	0
57	1A	1013	0	0	0	0
57	1B	30	0	0	0	0
57	1D	17	0	0	0	0
57	1E	8	0	0	0	0
57	1F	17	0	0	0	0
57	1G	4	0	0	0	0
57	1H	2	0	0	0	0
57	1N	4	0	0	0	0
57	1O	1	0	0	0	0
57	1P	3	0	0	0	0
57	1Q	5	0	0	0	0
57	1R	7	0	0	0	0
57	1T	7	0	0	0	0
57	1U	5	0	0	0	0
57	1V	7	0	0	0	0
57	1W	3	0	0	0	0
57	1Y	1	0	0	0	0
57	1Z	1	0	0	0	0
57	1a	267	0	0	0	0
57	1b	1	0	0	0	0
57	1d	5	0	0	0	0
57	1e	5	0	0	0	0
57	1f	2	0	0	0	0
57	1g	2	0	0	0	0
57	1h	2	0	0	0	0
57	1i	1	0	0	0	0
57	1l	2	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
57	1m	1	0	0	0	0
57	1n	3	0	0	0	0
57	1o	3	0	0	0	0
57	1t	2	0	0	0	0
57	1x	2	0	0	0	0
57	1y	2	0	0	0	0
57	20	2	0	0	0	0
57	23	1	0	0	0	0
57	25	2	0	0	0	0
57	27	2	0	0	0	0
57	28	2	0	0	0	0
57	29	1	0	0	0	0
57	2A	731	0	0	0	0
57	2B	19	0	0	0	0
57	2D	10	0	0	0	0
57	2E	6	0	0	0	0
57	2F	5	0	0	0	0
57	2G	2	0	0	0	0
57	2I	1	0	0	0	0
57	2O	2	0	0	0	0
57	2P	1	0	0	0	0
57	2Q	3	0	0	0	0
57	2R	3	0	0	0	0
57	2T	4	0	0	0	0
57	2U	2	0	0	0	0
57	2V	3	0	0	0	0
57	2W	3	0	0	0	0
57	2X	1	0	0	0	0
57	2Y	1	0	0	0	0
57	2a	183	0	0	0	0
57	2e	1	0	0	0	0
57	2f	2	0	0	0	0
57	2j	1	0	0	0	0
57	2k	1	0	0	0	0
57	2p	1	0	0	0	0
57	2r	1	0	0	0	0
57	2t	1	0	0	0	0
57	2x	2	0	0	0	0
58	18	8	0	14	1	0
58	1A	8	0	14	0	0
58	1T	8	0	14	1	0
58	1a	8	0	12	4	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
58	2A	8	0	14	0	0
58	2B	8	0	14	0	0
59	1B	12	0	12	1	0
59	1F	12	0	12	0	0
60	14	1	0	0	0	0
60	15	1	0	0	0	0
60	16	1	0	0	0	0
60	19	1	0	0	0	0
60	1Y	1	0	0	0	0
60	1n	1	0	0	0	0
60	24	1	0	0	0	0
60	25	1	0	0	0	0
60	26	1	0	0	0	0
60	29	1	0	0	0	0
60	2Y	1	0	0	0	0
60	2n	1	0	0	0	0
61	1d	8	0	0	0	0
61	2d	8	0	0	0	0
62	10	23	0	0	1	0
62	11	26	0	0	0	0
62	12	10	0	0	2	0
62	13	25	0	0	0	0
62	14	2	0	0	0	0
62	15	18	0	0	0	0
62	16	18	0	0	1	0
62	17	17	0	0	1	0
62	18	25	0	0	1	0
62	19	5	0	0	0	0
62	1A	3935	0	0	72	0
62	1B	80	0	0	2	0
62	1D	108	0	0	1	0
62	1E	74	0	0	1	0
62	1F	60	0	0	1	0
62	1G	16	0	0	0	0
62	1H	15	0	0	0	0
62	1I	5	0	0	0	0
62	1N	54	0	0	1	0
62	1O	21	0	0	1	0
62	1P	63	0	0	2	0
62	1Q	39	0	0	2	0
62	1R	28	0	0	1	0
62	1S	12	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
62	1T	32	0	0	2	0
62	1U	53	0	0	4	0
62	1V	33	0	0	0	0
62	1W	24	0	0	1	0
62	1X	24	0	0	1	0
62	1Y	15	0	0	2	0
62	1Z	7	0	0	0	0
62	1a	424	0	0	17	0
62	1b	1	0	0	0	0
62	1d	8	0	0	1	0
62	1e	2	0	0	0	0
62	1f	2	0	0	0	0
62	1g	1	0	0	0	0
62	1h	1	0	0	0	0
62	1l	2	0	0	1	0
62	1o	4	0	0	0	0
62	1p	1	0	0	0	0
62	1t	1	0	0	0	0
62	1w	6	0	0	0	0
62	1x	2	0	0	0	0
62	1y	2	0	0	0	0
62	20	8	0	0	0	0
62	21	15	0	0	2	0
62	23	2	0	0	1	0
62	25	11	0	0	1	0
62	26	7	0	0	1	0
62	27	5	0	0	0	0
62	28	16	0	0	0	0
62	2A	2058	0	0	72	0
62	2B	39	0	0	2	0
62	2D	51	0	0	1	0
62	2E	32	0	0	3	0
62	2F	26	0	0	2	0
62	2G	7	0	0	0	0
62	2H	1	0	0	0	0
62	2I	2	0	0	1	0
62	2N	5	0	0	0	0
62	2O	16	0	0	0	0
62	2P	26	0	0	2	0
62	2Q	14	0	0	0	0
62	2R	17	0	0	3	0
62	2S	3	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
62	2T	7	0	0	0	0
62	2U	15	0	0	0	0
62	2V	3	0	0	1	0
62	2W	17	0	0	0	0
62	2X	8	0	0	0	0
62	2Y	2	0	0	0	0
62	2Z	6	0	0	0	0
62	2a	243	0	0	12	0
62	2e	2	0	0	0	0
62	2l	3	0	0	0	0
62	2m	1	0	0	0	0
62	2n	2	0	0	0	0
62	2o	2	0	0	0	0
62	2p	1	0	0	0	0
62	2r	4	0	0	0	0
62	2t	2	0	0	0	0
62	2w	4	0	0	0	0
62	2x	3	0	0	1	0
All	All	297350	0	194813	3545	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 3545 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:1082:U:H3	1:1A:1086:A:N6	1.37	1.20
1:2A:2122:U:H3	1:2A:2176:A:N6	1.58	1.00
1:1A:1082:U:O4	1:1A:1086:A:N1	1.98	0.95
22:10:10:THR:HG22	22:10:12:ASN:H	1.27	0.95
32:1a:1003:G:H2'	32:1a:1004:A:H4'	1.46	0.95

There are no symmetry-related clashes.

5.3 Torsion angles

5.3.1 Protein backbone

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
3	1D	273/276 (99%)	260 (95%)	13 (5%)	0	100	100
3	2D	273/276 (99%)	259 (95%)	14 (5%)	0	100	100
4	1E	202/206 (98%)	193 (96%)	8 (4%)	1 (0%)	24	32
4	2E	202/206 (98%)	192 (95%)	9 (4%)	1 (0%)	24	32
5	1F	201/210 (96%)	193 (96%)	7 (4%)	1 (0%)	24	32
5	2F	201/210 (96%)	192 (96%)	8 (4%)	1 (0%)	24	32
6	1G	179/182 (98%)	159 (89%)	18 (10%)	2 (1%)	11	12
6	2G	179/182 (98%)	157 (88%)	19 (11%)	3 (2%)	7	6
7	1H	172/180 (96%)	162 (94%)	8 (5%)	2 (1%)	10	11
7	2H	171/180 (95%)	148 (86%)	21 (12%)	2 (1%)	10	11
8	1I	145/148 (98%)	130 (90%)	14 (10%)	1 (1%)	18	24
8	2I	144/148 (97%)	128 (89%)	13 (9%)	3 (2%)	5	4
9	1N	138/140 (99%)	133 (96%)	5 (4%)	0	100	100
9	2N	138/140 (99%)	133 (96%)	5 (4%)	0	100	100
10	1O	120/122 (98%)	113 (94%)	6 (5%)	1 (1%)	16	20
10	2O	120/122 (98%)	114 (95%)	4 (3%)	2 (2%)	7	6
11	1P	147/150 (98%)	137 (93%)	9 (6%)	1 (1%)	18	24
11	2P	147/150 (98%)	139 (95%)	6 (4%)	2 (1%)	9	8
12	1Q	139/141 (99%)	133 (96%)	6 (4%)	0	100	100
12	2Q	139/141 (99%)	131 (94%)	8 (6%)	0	100	100
13	1R	116/118 (98%)	114 (98%)	2 (2%)	0	100	100
13	2R	116/118 (98%)	113 (97%)	3 (3%)	0	100	100
14	1S	108/112 (96%)	102 (94%)	6 (6%)	0	100	100
14	2S	108/112 (96%)	102 (94%)	5 (5%)	1 (1%)	14	17
15	1T	129/146 (88%)	121 (94%)	8 (6%)	0	100	100
15	2T	129/146 (88%)	121 (94%)	6 (5%)	2 (2%)	7	7
16	1U	114/118 (97%)	114 (100%)	0	0	100	100
16	2U	114/118 (97%)	112 (98%)	2 (2%)	0	100	100
17	1V	99/101 (98%)	96 (97%)	3 (3%)	0	100	100
17	2V	99/101 (98%)	91 (92%)	6 (6%)	2 (2%)	6	4

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
18	1W	110/113 (97%)	110 (100%)	0	0	100	100
18	2W	110/113 (97%)	109 (99%)	1 (1%)	0	100	100
19	1X	93/96 (97%)	90 (97%)	2 (2%)	1 (1%)	11	12
19	2X	93/96 (97%)	90 (97%)	2 (2%)	1 (1%)	11	12
20	1Y	105/110 (96%)	97 (92%)	8 (8%)	0	100	100
20	2Y	105/110 (96%)	95 (90%)	10 (10%)	0	100	100
21	1Z	201/206 (98%)	187 (93%)	12 (6%)	2 (1%)	12	14
21	2Z	199/206 (97%)	186 (94%)	11 (6%)	2 (1%)	12	14
22	10	81/85 (95%)	79 (98%)	2 (2%)	0	100	100
22	20	81/85 (95%)	78 (96%)	2 (2%)	1 (1%)	10	11
23	11	95/98 (97%)	92 (97%)	3 (3%)	0	100	100
23	21	95/98 (97%)	92 (97%)	2 (2%)	1 (1%)	11	12
24	12	68/72 (94%)	67 (98%)	1 (2%)	0	100	100
24	22	68/72 (94%)	66 (97%)	2 (3%)	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%)	47 (70%)	16 (24%)	4 (6%)	1	0
26	24	67/71 (94%)	47 (70%)	18 (27%)	2 (3%)	3	2
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%)	49 (96%)	2 (4%)	0	100	100
28	26	51/54 (94%)	48 (94%)	3 (6%)	0	100	100
29	17	46/49 (94%)	45 (98%)	0	1 (2%)	5	3
29	27	46/49 (94%)	45 (98%)	1 (2%)	0	100	100
30	18	62/65 (95%)	61 (98%)	1 (2%)	0	100	100
30	28	62/65 (95%)	62 (100%)	0	0	100	100
31	19	35/37 (95%)	35 (100%)	0	0	100	100
31	29	35/37 (95%)	34 (97%)	1 (3%)	0	100	100
33	1b	229/256 (90%)	191 (83%)	28 (12%)	10 (4%)	2	0
33	2b	229/256 (90%)	199 (87%)	21 (9%)	9 (4%)	2	1
34	1c	204/239 (85%)	191 (94%)	12 (6%)	1 (0%)	24	32

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
34	2c	204/239 (85%)	179 (88%)	24 (12%)	1 (0%)	24	32
35	1d	206/209 (99%)	196 (95%)	10 (5%)	0	100	100
35	2d	206/209 (99%)	200 (97%)	6 (3%)	0	100	100
36	1e	146/162 (90%)	141 (97%)	4 (3%)	1 (1%)	18	24
36	2e	146/162 (90%)	141 (97%)	5 (3%)	0	100	100
37	1f	98/101 (97%)	94 (96%)	4 (4%)	0	100	100
37	2f	98/101 (97%)	96 (98%)	2 (2%)	0	100	100
38	1g	153/156 (98%)	146 (95%)	7 (5%)	0	100	100
38	2g	153/156 (98%)	144 (94%)	9 (6%)	0	100	100
39	1h	135/138 (98%)	127 (94%)	7 (5%)	1 (1%)	18	24
39	2h	135/138 (98%)	126 (93%)	9 (7%)	0	100	100
40	1i	125/128 (98%)	114 (91%)	10 (8%)	1 (1%)	16	20
40	2i	124/128 (97%)	106 (86%)	15 (12%)	3 (2%)	4	3
41	1j	95/105 (90%)	82 (86%)	9 (10%)	4 (4%)	2	0
41	2j	94/105 (90%)	81 (86%)	9 (10%)	4 (4%)	2	0
42	1k	112/129 (87%)	104 (93%)	8 (7%)	0	100	100
42	2k	112/129 (87%)	106 (95%)	3 (3%)	3 (3%)	4	2
43	1l	119/132 (90%)	113 (95%)	5 (4%)	1 (1%)	16	20
43	2l	119/132 (90%)	111 (93%)	7 (6%)	1 (1%)	16	20
44	1m	114/126 (90%)	104 (91%)	9 (8%)	1 (1%)	14	17
44	2m	112/126 (89%)	96 (86%)	13 (12%)	3 (3%)	4	2
45	1n	58/61 (95%)	57 (98%)	1 (2%)	0	100	100
45	2n	58/61 (95%)	54 (93%)	4 (7%)	0	100	100
46	1o	86/89 (97%)	82 (95%)	2 (2%)	2 (2%)	5	3
46	2o	86/89 (97%)	81 (94%)	3 (4%)	2 (2%)	5	3
47	1p	80/88 (91%)	73 (91%)	7 (9%)	0	100	100
47	2p	80/88 (91%)	75 (94%)	5 (6%)	0	100	100
48	1q	97/105 (92%)	90 (93%)	7 (7%)	0	100	100
48	2q	97/105 (92%)	91 (94%)	6 (6%)	0	100	100
49	1r	66/88 (75%)	64 (97%)	2 (3%)	0	100	100
49	2r	66/88 (75%)	62 (94%)	4 (6%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
50	1s	81/93 (87%)	76 (94%)	3 (4%)	2 (2%)	4	3
50	2s	81/93 (87%)	70 (86%)	10 (12%)	1 (1%)	10	11
51	1t	94/106 (89%)	88 (94%)	3 (3%)	3 (3%)	3	1
51	2t	96/106 (91%)	89 (93%)	4 (4%)	3 (3%)	3	1
52	1u	21/27 (78%)	20 (95%)	1 (5%)	0	100	100
52	2u	21/27 (78%)	19 (90%)	2 (10%)	0	100	100
53	1y	95/113 (84%)	92 (97%)	3 (3%)	0	100	100
53	2y	94/113 (83%)	88 (94%)	6 (6%)	0	100	100
56	1v	1/3 (33%)	1 (100%)	0	0	100	100
56	2v	1/3 (33%)	1 (100%)	0	0	100	100
All	All	11643/12360 (94%)	10886 (94%)	657 (6%)	100 (1%)	14	17

5 of 100 Ramachandran outliers are listed below:

Mol	Chain	Res	Type
5	1F	130	ALA
6	1G	51	ARG
8	1I	86	THR
21	1Z	53	ILE
33	1b	21	ARG

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	214/218 (98%)	198 (92%)	16 (8%)	12	16
3	2D	215/218 (99%)	197 (92%)	18 (8%)	10	12
4	1E	164/166 (99%)	151 (92%)	13 (8%)	11	14
4	2E	164/166 (99%)	151 (92%)	13 (8%)	11	14
5	1F	160/166 (96%)	143 (89%)	17 (11%)	6	6
5	2F	159/166 (96%)	143 (90%)	16 (10%)	7	7

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
6	1G	144/156 (92%)	130 (90%)	14 (10%)	8	8
6	2G	142/156 (91%)	128 (90%)	14 (10%)	7	7
7	1H	144/148 (97%)	136 (94%)	8 (6%)	19	27
7	2H	143/148 (97%)	126 (88%)	17 (12%)	5	4
8	1I	111/124 (90%)	94 (85%)	17 (15%)	3	2
8	2I	108/124 (87%)	90 (83%)	18 (17%)	2	1
9	1N	119/119 (100%)	108 (91%)	11 (9%)	8	9
9	2N	118/119 (99%)	109 (92%)	9 (8%)	12	15
10	1O	100/100 (100%)	95 (95%)	5 (5%)	22	32
10	2O	100/100 (100%)	95 (95%)	5 (5%)	22	32
11	1P	115/116 (99%)	110 (96%)	5 (4%)	26	38
11	2P	115/116 (99%)	107 (93%)	8 (7%)	14	18
12	1Q	111/111 (100%)	103 (93%)	8 (7%)	13	17
12	2Q	111/111 (100%)	104 (94%)	7 (6%)	16	23
13	1R	101/101 (100%)	95 (94%)	6 (6%)	18	26
13	2R	101/101 (100%)	96 (95%)	5 (5%)	22	32
14	1S	87/88 (99%)	75 (86%)	12 (14%)	3	3
14	2S	85/88 (97%)	75 (88%)	10 (12%)	5	4
15	1T	115/127 (91%)	107 (93%)	8 (7%)	14	18
15	2T	113/127 (89%)	105 (93%)	8 (7%)	13	18
16	1U	93/94 (99%)	86 (92%)	7 (8%)	12	16
16	2U	93/94 (99%)	86 (92%)	7 (8%)	12	16
17	1V	81/82 (99%)	79 (98%)	2 (2%)	42	56
17	2V	80/82 (98%)	72 (90%)	8 (10%)	7	7
18	1W	90/92 (98%)	85 (94%)	5 (6%)	19	27
18	2W	90/92 (98%)	85 (94%)	5 (6%)	19	27
19	1X	77/78 (99%)	72 (94%)	5 (6%)	15	21
19	2X	77/78 (99%)	76 (99%)	1 (1%)	61	73
20	1Y	86/91 (94%)	75 (87%)	11 (13%)	4	3
20	2Y	86/91 (94%)	74 (86%)	12 (14%)	3	3
21	1Z	169/179 (94%)	154 (91%)	15 (9%)	9	10

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
21	2Z	165/179 (92%)	151 (92%)	14 (8%)	10	11
22	10	65/67 (97%)	61 (94%)	4 (6%)	16	23
22	20	64/67 (96%)	61 (95%)	3 (5%)	23	35
23	11	79/83 (95%)	73 (92%)	6 (8%)	12	15
23	21	81/83 (98%)	72 (89%)	9 (11%)	6	5
24	12	65/67 (97%)	61 (94%)	4 (6%)	16	23
24	22	66/67 (98%)	60 (91%)	6 (9%)	9	10
25	13	51/52 (98%)	48 (94%)	3 (6%)	18	26
25	23	50/52 (96%)	46 (92%)	4 (8%)	11	13
26	14	58/63 (92%)	54 (93%)	4 (7%)	14	19
26	24	54/63 (86%)	43 (80%)	11 (20%)	1	0
27	15	51/52 (98%)	48 (94%)	3 (6%)	18	26
27	25	50/52 (96%)	48 (96%)	2 (4%)	28	41
28	16	51/52 (98%)	46 (90%)	5 (10%)	7	7
28	26	50/52 (96%)	48 (96%)	2 (4%)	28	41
29	17	41/42 (98%)	38 (93%)	3 (7%)	13	17
29	27	41/42 (98%)	37 (90%)	4 (10%)	7	7
30	18	54/55 (98%)	49 (91%)	5 (9%)	8	9
30	28	54/55 (98%)	51 (94%)	3 (6%)	19	27
31	19	34/34 (100%)	33 (97%)	1 (3%)	37	52
31	29	34/34 (100%)	33 (97%)	1 (3%)	37	52
33	1b	191/220 (87%)	171 (90%)	20 (10%)	6	6
33	2b	187/220 (85%)	166 (89%)	21 (11%)	6	5
34	1c	144/188 (77%)	131 (91%)	13 (9%)	9	10
34	2c	140/188 (74%)	126 (90%)	14 (10%)	7	7
35	1d	171/181 (94%)	155 (91%)	16 (9%)	8	8
35	2d	172/181 (95%)	151 (88%)	21 (12%)	5	4
36	1e	114/123 (93%)	104 (91%)	10 (9%)	9	10
36	2e	114/123 (93%)	107 (94%)	7 (6%)	17	24
37	1f	85/90 (94%)	76 (89%)	9 (11%)	6	6
37	2f	85/90 (94%)	77 (91%)	8 (9%)	8	8

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
38	1g	120/127 (94%)	112 (93%)	8 (7%)	15	20
38	2g	119/127 (94%)	113 (95%)	6 (5%)	22	32
39	1h	116/119 (98%)	111 (96%)	5 (4%)	26	38
39	2h	114/119 (96%)	103 (90%)	11 (10%)	8	8
40	1i	91/99 (92%)	81 (89%)	10 (11%)	6	5
40	2i	88/99 (89%)	78 (89%)	10 (11%)	5	5
41	1j	68/92 (74%)	57 (84%)	11 (16%)	2	1
41	2j	68/92 (74%)	61 (90%)	7 (10%)	7	6
42	1k	83/99 (84%)	78 (94%)	5 (6%)	17	26
42	2k	83/99 (84%)	77 (93%)	6 (7%)	13	17
43	1l	96/108 (89%)	95 (99%)	1 (1%)	68	77
43	2l	96/108 (89%)	92 (96%)	4 (4%)	26	39
44	1m	90/101 (89%)	83 (92%)	7 (8%)	11	14
44	2m	87/101 (86%)	76 (87%)	11 (13%)	4	4
45	1n	49/50 (98%)	46 (94%)	3 (6%)	17	24
45	2n	49/50 (98%)	45 (92%)	4 (8%)	10	13
46	1o	78/80 (98%)	75 (96%)	3 (4%)	29	43
46	2o	78/80 (98%)	71 (91%)	7 (9%)	9	10
47	1p	69/74 (93%)	60 (87%)	9 (13%)	4	3
47	2p	68/74 (92%)	63 (93%)	5 (7%)	13	16
48	1q	94/97 (97%)	89 (95%)	5 (5%)	20	30
48	2q	94/97 (97%)	89 (95%)	5 (5%)	20	30
49	1r	59/77 (77%)	54 (92%)	5 (8%)	10	11
49	2r	59/77 (77%)	55 (93%)	4 (7%)	14	20
50	1s	68/80 (85%)	56 (82%)	12 (18%)	2	1
50	2s	67/80 (84%)	58 (87%)	9 (13%)	4	3
51	1t	71/82 (87%)	68 (96%)	3 (4%)	26	39
51	2t	70/82 (85%)	63 (90%)	7 (10%)	7	7
52	1u	18/22 (82%)	17 (94%)	1 (6%)	19	27
52	2u	18/22 (82%)	18 (100%)	0	100	100
53	1y	82/98 (84%)	78 (95%)	4 (5%)	22	33

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
53	2y	79/98 (81%)	70 (89%)	9 (11%)	5	5
56	1v	2/2 (100%)	2 (100%)	0	100	100
56	2v	2/2 (100%)	2 (100%)	0	100	100
All	All	9535/10264 (93%)	8736 (92%)	799 (8%)	10	12

5 of 799 residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
8	2I	41	GLU
21	2Z	141	VAL
8	2I	116	LEU
8	2I	40	THR
14	2S	78	LEU

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. 5 of 171 such sidechains are listed below:

Mol	Chain	Res	Type
21	2Z	65	GLN
37	2f	13	ASN
26	24	40	HIS
34	2c	136	GLN
40	2i	3	GLN

5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	1A	2865/2915 (98%)	389 (13%)	36 (1%)
1	2A	2858/2915 (98%)	457 (15%)	33 (1%)
2	1B	119/121 (98%)	12 (10%)	0
2	2B	119/121 (98%)	15 (12%)	0
32	1a	1497/1521 (98%)	232 (15%)	0
32	2a	1501/1521 (98%)	263 (17%)	0
54	1w	1/4 (25%)	0	0
54	2w	1/4 (25%)	0	0
55	1x	1/4 (25%)	0	0
55	2x	1/4 (25%)	0	0
All	All	8963/9130 (98%)	1368 (15%)	69 (0%)

5 of 1368 RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	1A	34	C
1	1A	35	G
1	1A	45	C
1	1A	71	A
1	1A	74	A

5 of 69 RNA pucker outliers are listed below:

Mol	Chain	Res	Type
1	2A	1420	U
1	2A	1491	G
1	2A	2406	U
1	1A	1442	G
1	1A	1379	A

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

52 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	$\# Z > 2$
43	0TD	1l	92	43	8,9,10	4.78	2 (25%)	6,11,13	2.20	3 (50%)
1	OMG	1A	2251	12,55,1	23,26,27	1.26	3 (13%)	32,38,41	2.01	5 (15%)
43	0TD	2l	92	43	8,9,10	4.37	2 (25%)	6,11,13	4.35	3 (50%)
1	PSU	1A	1911	1	18,21,22	1.39	2 (11%)	21,30,33	1.96	4 (19%)
32	UR3	1a	1498	32	19,22,23	1.04	1 (5%)	26,32,35	1.89	4 (15%)
1	5MU	1A	1939	1,57	19,22,23	1.38	5 (26%)	27,32,35	2.28	6 (22%)
1	OMU	1A	2552	1,57	19,22,23	1.26	3 (15%)	25,31,34	1.92	6 (24%)
32	5MC	1a	1407	32	19,22,23	1.69	2 (10%)	26,32,35	1.14	3 (11%)
32	MA6	1a	1519	32	23,26,27	0.44	0	33,38,41	1.99	9 (27%)
54	PPU	2w	76	1,54	37,40,41	1.23	3 (8%)	47,57,60	2.04	12 (25%)
32	MA6	2a	1518	32	23,26,27	0.38	0	33,38,41	2.03	9 (27%)
32	4OC	2a	1402	32	20,23,24	0.76	0	25,32,35	0.93	1 (4%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
32	MA6	1a	1518	32	23,26,27	0.37	0	33,38,41	1.96	9 (27%)
1	OMC	2A	1920	1	19,22,23	0.79	0	25,31,34	0.95	1 (4%)
1	2MA	2A	2503	1,57	22,25,26	1.45	3 (13%)	32,37,40	2.44	9 (28%)
1	5MU	2A	1939	1,57	19,22,23	1.44	6 (31%)	27,32,35	2.21	6 (22%)
1	PSU	2A	1917	1	18,21,22	1.34	2 (11%)	21,30,33	1.97	3 (14%)
1	5MC	2A	1962	1,57	19,22,23	1.67	3 (15%)	26,32,35	1.11	2 (7%)
1	PSU	2A	1911	1	18,21,22	1.43	2 (11%)	21,30,33	2.01	3 (14%)
54	PPU	1w	76	1,54	37,40,41	1.21	5 (13%)	47,57,60	1.93	11 (23%)
55	8AN	2x	76	55,57,56	21,24,25	1.42	3 (14%)	26,35,38	2.22	9 (34%)
1	5MU	2A	1915	1	19,22,23	1.46	4 (21%)	27,32,35	2.18	9 (33%)
32	5MC	2a	1400	32	19,22,23	1.56	3 (15%)	26,32,35	1.28	4 (15%)
1	2MA	1A	2503	1,57	22,25,26	1.47	6 (27%)	32,37,40	2.29	7 (21%)
1	5MC	1A	1962	1	19,22,23	1.71	3 (15%)	26,32,35	1.14	4 (15%)
32	2MG	1a	1207	32	23,26,27	1.23	3 (13%)	33,38,41	2.30	8 (24%)
32	5MC	2a	1407	32	19,22,23	1.57	3 (15%)	26,32,35	1.13	3 (11%)
32	5MC	1a	967	32	19,22,23	1.67	3 (15%)	26,32,35	1.10	2 (7%)
32	UR3	2a	1498	32	19,22,23	0.98	1 (5%)	26,32,35	1.75	3 (11%)
32	PSU	1a	516	32,57	18,21,22	1.40	2 (11%)	21,30,33	1.93	3 (14%)
32	PSU	2a	516	32,57	18,21,22	1.32	2 (11%)	21,30,33	2.02	5 (23%)
1	5MC	2A	1942	1	19,22,23	1.83	3 (15%)	26,32,35	1.19	2 (7%)
1	OMC	1A	1920	1	19,22,23	0.79	0	25,31,34	0.96	1 (4%)
1	OMG	2A	2251	12,55,1,57	23,26,27	1.18	3 (13%)	32,38,41	1.93	6 (18%)
32	M2G	2a	966	32	24,27,28	1.28	3 (12%)	33,40,43	1.82	5 (15%)
32	M2G	1a	966	32	24,27,28	1.31	3 (12%)	33,40,43	1.84	5 (15%)
32	4OC	1a	1402	32	20,23,24	0.72	0	25,32,35	1.01	1 (4%)
32	G7M	1a	527	32,57	23,26,27	1.49	3 (13%)	34,39,42	1.67	5 (14%)
1	PSU	1A	1917	1	18,21,22	1.33	2 (11%)	21,30,33	2.00	4 (19%)
1	PSU	2A	2605	1	18,21,22	1.28	2 (11%)	21,30,33	2.07	4 (19%)
32	MA6	2a	1519	32	23,26,27	0.42	0	33,38,41	2.09	9 (27%)
1	OMU	2A	2552	1,57	19,22,23	1.23	4 (21%)	25,31,34	1.97	6 (24%)
1	5MU	1A	1915	1	19,22,23	1.47	4 (21%)	27,32,35	2.29	9 (33%)
32	5MC	1a	1404	32	19,22,23	1.60	3 (15%)	26,32,35	1.08	2 (7%)
32	5MC	1a	1400	32	19,22,23	1.76	3 (15%)	26,32,35	1.17	2 (7%)
32	5MC	2a	1404	32	19,22,23	1.78	3 (15%)	26,32,35	1.24	4 (15%)
32	G7M	2a	527	32,57	23,26,27	1.50	3 (13%)	34,39,42	1.68	5 (14%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
1	5MC	1A	1942	1,57	19,22,23	1.42	3 (15%)	26,32,35	1.09	2 (7%)
1	PSU	1A	2605	1	18,21,22	1.36	3 (16%)	21,30,33	2.19	4 (19%)
32	5MC	2a	967	32	19,22,23	1.83	2 (10%)	26,32,35	1.09	2 (7%)
32	2MG	2a	1207	32	23,26,27	1.21	2 (8%)	33,38,41	2.20	7 (21%)
55	8AN	1x	76	55,57,56	21,24,25	1.35	3 (14%)	26,35,38	2.32	10 (38%)

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
43	0TD	1l	92	43	-	2/7/12/14	-
1	OMG	1A	2251	12,55,1	-	0/9/27/28	0/3/3/3
43	0TD	2l	92	43	-	2/7/12/14	-
1	PSU	1A	1911	1	-	0/7/25/26	0/2/2/2
32	UR3	1a	1498	32	-	0/7/25/26	0/2/2/2
1	5MU	1A	1939	1,57	-	0/7/25/26	0/2/2/2
1	OMU	1A	2552	1,57	-	0/9/27/28	0/2/2/2
32	5MC	1a	1407	32	-	0/7/25/26	0/2/2/2
32	MA6	1a	1519	32	-	2/11/29/30	0/3/3/3
54	PPU	2w	76	1,54	-	2/25/43/44	0/4/4/4
32	MA6	2a	1518	32	-	0/11/29/30	0/3/3/3
32	4OC	2a	1402	32	-	0/9/29/30	0/2/2/2
32	MA6	1a	1518	32	-	0/11/29/30	0/3/3/3
1	OMC	2A	1920	1	-	0/9/27/28	0/2/2/2
1	2MA	2A	2503	1,57	-	2/7/25/26	0/3/3/3
1	5MU	2A	1939	1,57	-	0/7/25/26	0/2/2/2
1	PSU	2A	1917	1	-	0/7/25/26	0/2/2/2
1	5MC	2A	1962	1,57	-	0/7/25/26	0/2/2/2
1	PSU	2A	1911	1	-	0/7/25/26	0/2/2/2
54	PPU	1w	76	1,54	-	0/25/43/44	0/4/4/4
55	8AN	2x	76	55,57,56	-	1/7/25/26	0/3/3/3
1	5MU	2A	1915	1	-	2/7/25/26	0/2/2/2
32	5MC	2a	1400	32	-	2/7/25/26	0/2/2/2
1	2MA	1A	2503	1,57	-	1/7/25/26	0/3/3/3
1	5MC	1A	1962	1	-	0/7/25/26	0/2/2/2
32	2MG	1a	1207	32	-	0/9/27/28	0/3/3/3
32	5MC	2a	1407	32	-	0/7/25/26	0/2/2/2
32	5MC	1a	967	32	-	0/7/25/26	0/2/2/2

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
32	UR3	2a	1498	32	-	0/7/25/26	0/2/2/2
32	PSU	1a	516	32,57	-	0/7/25/26	0/2/2/2
32	PSU	2a	516	32,57	-	0/7/25/26	0/2/2/2
1	5MC	2A	1942	1	-	0/7/25/26	0/2/2/2
1	OMC	1A	1920	1	-	1/9/27/28	0/2/2/2
1	OMG	2A	2251	12,55,1,57	-	1/9/27/28	0/3/3/3
32	M2G	2a	966	32	-	0/11/29/30	0/3/3/3
32	M2G	1a	966	32	-	0/11/29/30	0/3/3/3
32	4OC	1a	1402	32	-	0/9/29/30	0/2/2/2
32	G7M	1a	527	32,57	-	2/7/25/26	0/3/3/3
1	PSU	1A	1917	1	-	0/7/25/26	0/2/2/2
1	PSU	2A	2605	1	-	0/7/25/26	0/2/2/2
32	MA6	2a	1519	32	-	2/11/29/30	0/3/3/3
1	OMU	2A	2552	1,57	-	0/9/27/28	0/2/2/2
1	5MU	1A	1915	1	-	2/7/25/26	0/2/2/2
32	5MC	1a	1404	32	-	0/7/25/26	0/2/2/2
32	5MC	1a	1400	32	-	0/7/25/26	0/2/2/2
32	5MC	2a	1404	32	-	0/7/25/26	0/2/2/2
32	G7M	2a	527	32,57	-	1/7/25/26	0/3/3/3
1	5MC	1A	1942	1,57	-	0/7/25/26	0/2/2/2
1	PSU	1A	2605	1	-	0/7/25/26	0/2/2/2
32	5MC	2a	967	32	-	0/7/25/26	0/2/2/2
32	2MG	2a	1207	32	-	0/9/27/28	0/3/3/3
55	8AN	1x	76	55,57,56	-	1/7/25/26	0/3/3/3

The worst 5 of 129 bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
43	1l	92	0TD	CB-SB	-12.67	1.69	1.82
43	2l	92	0TD	CB-SB	-11.69	1.70	1.82
32	2a	967	5MC	C5-C4	6.91	1.49	1.44
1	2A	1942	5MC	C5-C4	6.85	1.49	1.44
32	2a	1404	5MC	C5-C4	6.60	1.49	1.44

The worst 5 of 261 bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
43	2l	92	0TD	CSB-SB-CB	-9.74	84.85	102.36
1	2A	2503	2MA	C5-C4-N3	-8.03	118.73	127.18
1	1A	2503	2MA	C5-C4-N3	-7.90	118.86	127.18
32	1a	1498	UR3	C4-N3-C2	-7.60	118.46	124.58
32	2a	1498	UR3	C4-N3-C2	-7.15	118.82	124.58

There are no chirality outliers.

5 of 26 torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
1	1A	1915	5MU	O4'-C1'-N1-C2
1	1A	1915	5MU	O4'-C1'-N1-C6
43	1l	92	0TD	CA-CB-SB-CSB
43	1l	92	0TD	CG-CB-SB-CSB
1	2A	1915	5MU	O4'-C1'-N1-C2

There are no ring outliers.

17 monomers are involved in 18 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
43	1l	92	0TD	1	0
1	1A	1939	5MU	1	0
32	1a	1519	MA6	1	0
32	2a	1518	MA6	1	0
32	2a	1402	4OC	2	0
32	1a	1518	MA6	1	0
1	2A	1920	OMC	1	0
1	2A	1939	5MU	1	0
1	2A	1917	PSU	1	0
54	1w	76	PPU	2	0
55	2x	76	8AN	1	0
1	2A	1915	5MU	1	0
32	2a	1400	5MC	1	0
32	1a	1402	4OC	1	0
32	2a	1519	MA6	2	0
32	1a	1400	5MC	1	0
32	2a	1404	5MC	1	0

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

Of 2490 ligands modelled in this entry, 2480 are monoatomic - leaving 10 for Mogul analysis.

In the following table, the Counts columns list the number of bonds (or angles) for which Mogul statistics could be retrieved, the number of bonds (or angles) that are observed in the model and the number of bonds (or angles) that are defined in the Chemical Component Dictionary. The

Link column lists molecule types, if any, to which the group is linked. The Z score for a bond length (or angle) is the number of standard deviations the observed value is removed from the expected value. A bond length (or angle) with $|Z| > 2$ is considered an outlier worth inspection. RMSZ is the root-mean-square of all Z scores of the bond lengths (or angles).

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
58	MPD	1A	4014	-	7,7,7	0.36	0	9,10,10	0.44	0
58	MPD	18	104	-	7,7,7	0.30	0	9,10,10	0.30	0
58	MPD	1a	1868	32	7,7,7	0.41	0	9,10,10	0.56	0
58	MPD	2A	3732	-	7,7,7	0.38	0	9,10,10	0.22	0
58	MPD	2B	220	-	7,7,7	0.34	0	9,10,10	0.28	0
61	SF4	2d	501	35	0,12,12	-	-	-	-	-
61	SF4	1d	306	35	0,12,12	-	-	-	-	-
59	ARG	1F	318	-	10,11,11	0.73	1 (10%)	9,13,13	0.94	1 (11%)
58	MPD	1T	208	-	7,7,7	0.29	0	9,10,10	0.32	0
59	ARG	1B	231	-	10,11,11	0.71	1 (10%)	9,13,13	0.93	1 (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
58	MPD	1A	4014	-	-	1/5/5/5	-
58	MPD	18	104	-	-	1/5/5/5	-
58	MPD	1a	1868	32	-	4/5/5/5	-
58	MPD	2A	3732	-	-	2/5/5/5	-
58	MPD	2B	220	-	-	4/5/5/5	-
61	SF4	2d	501	35	-	-	0/6/5/5
61	SF4	1d	306	35	-	-	0/6/5/5
59	ARG	1F	318	-	-	2/11/11/11	-
58	MPD	1T	208	-	-	5/5/5/5	-
59	ARG	1B	231	-	-	3/11/11/11	-

All (2) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
59	1F	318	ARG	OXT-C	-2.13	1.23	1.30
59	1B	231	ARG	OXT-C	-2.03	1.24	1.30

All (2) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
59	1F	318	ARG	OXT-C-O	-2.71	117.93	124.08
59	1B	231	ARG	OXT-C-O	-2.56	118.26	124.08

There are no chirality outliers.

5 of 22 torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
58	1T	208	MPD	C1-C2-C3-C4
58	1a	1868	MPD	C2-C3-C4-O4
58	1a	1868	MPD	C2-C3-C4-C5
59	1B	231	ARG	C-CA-CB-CG
59	1F	318	ARG	NE-CD-CG-CB

There are no ring outliers.

4 monomers are involved in 7 short contacts:

Mol	Chain	Res	Type	Clashes	Symm-Clashes
58	18	104	MPD	1	0
58	1a	1868	MPD	4	0
58	1T	208	MPD	1	0
59	1B	231	ARG	1	0

5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

6 Fit of model and data ⓘ

6.1 Protein, DNA and RNA chains ⓘ

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

Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
1	1A	2861/2915 (98%)	-0.48	172 (6%) 27 24	22, 38, 90, 100	0
1	2A	2856/2915 (97%)	-0.00	185 (6%) 25 22	32, 55, 93, 102	0
2	1B	120/121 (99%)	-0.26	1 (0%) 82 83	34, 51, 67, 82	0
2	2B	120/121 (99%)	0.77	4 (3%) 49 50	59, 75, 82, 89	0
3	1D	275/276 (99%)	-0.20	4 (1%) 72 74	24, 37, 52, 71	0
3	2D	275/276 (99%)	0.14	5 (1%) 67 69	30, 49, 61, 81	0
4	1E	204/206 (99%)	-0.11	2 (0%) 79 80	22, 41, 61, 71	0
4	2E	204/206 (99%)	0.25	2 (0%) 79 80	33, 54, 68, 77	0
5	1F	203/210 (96%)	0.04	4 (1%) 65 66	23, 43, 68, 84	0
5	2F	203/210 (96%)	0.60	6 (2%) 52 53	33, 64, 76, 85	0
6	1G	181/182 (99%)	0.73	14 (7%) 19 17	49, 64, 75, 84	0
6	2G	181/182 (99%)	1.81	68 (37%) 1 0	70, 79, 85, 89	0
7	1H	174/180 (96%)	0.32	5 (2%) 53 55	40, 54, 66, 72	0
7	2H	173/180 (96%)	1.33	30 (17%) 4 3	65, 77, 83, 86	0
8	1I	147/148 (99%)	0.92	10 (6%) 23 21	42, 68, 78, 82	0
8	2I	146/148 (98%)	1.26	21 (14%) 6 4	56, 73, 81, 83	0
9	1N	140/140 (100%)	-0.06	0 100 100	29, 40, 59, 74	0
9	2N	140/140 (100%)	0.63	5 (3%) 46 46	45, 62, 73, 77	0
10	1O	122/122 (100%)	-0.13	1 (0%) 82 83	31, 41, 59, 64	0
10	2O	122/122 (100%)	0.25	0 100 100	43, 52, 64, 72	0
11	1P	149/150 (99%)	0.10	1 (0%) 84 86	23, 47, 67, 80	0
11	2P	149/150 (99%)	0.57	4 (2%) 56 57	39, 65, 78, 83	0
12	1Q	141/141 (100%)	-0.11	0 100 100	27, 40, 52, 68	0
12	2Q	141/141 (100%)	0.85	8 (5%) 29 26	41, 63, 72, 78	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
13	1R	118/118 (100%)	-0.23	0 100 100	27, 37, 54, 63	0
13	2R	118/118 (100%)	0.24	0 100 100	40, 50, 59, 71	0
14	1S	110/112 (98%)	0.28	1 (0%) 81 82	40, 51, 63, 69	0
14	2S	110/112 (98%)	1.33	14 (12%) 8 6	64, 71, 76, 79	0
15	1T	131/146 (89%)	0.09	5 (3%) 44 44	34, 45, 67, 79	0
15	2T	131/146 (89%)	0.38	4 (3%) 51 52	44, 56, 72, 81	0
16	1U	116/118 (98%)	-0.30	1 (0%) 81 82	25, 33, 48, 67	0
16	2U	116/118 (98%)	0.47	0 100 100	40, 58, 70, 75	0
17	1V	101/101 (100%)	-0.10	0 100 100	26, 43, 59, 68	0
17	2V	101/101 (100%)	0.70	3 (2%) 52 53	43, 68, 73, 80	0
18	1W	112/113 (99%)	-0.13	3 (2%) 56 57	28, 34, 53, 76	0
18	2W	112/113 (99%)	0.29	4 (3%) 46 46	38, 49, 65, 85	0
19	1X	95/96 (98%)	0.02	2 (2%) 63 65	30, 39, 61, 74	0
19	2X	95/96 (98%)	0.74	6 (6%) 26 23	45, 59, 72, 78	0
20	1Y	107/110 (97%)	0.33	2 (1%) 66 67	38, 50, 65, 75	0
20	2Y	107/110 (97%)	1.01	8 (7%) 20 17	58, 68, 76, 83	0
21	1Z	203/206 (98%)	0.56	6 (2%) 52 53	42, 59, 73, 82	0
21	2Z	201/206 (97%)	1.40	33 (16%) 4 3	63, 74, 81, 84	0
22	10	83/85 (97%)	-0.06	1 (1%) 76 78	30, 38, 50, 62	0
22	20	83/85 (97%)	0.97	5 (6%) 27 24	45, 60, 69, 74	0
23	11	97/98 (98%)	0.16	1 (1%) 79 80	29, 45, 67, 72	0
23	21	97/98 (98%)	0.45	2 (2%) 63 65	39, 55, 73, 77	0
24	12	70/72 (97%)	0.27	2 (2%) 53 55	39, 50, 59, 81	0
24	22	70/72 (97%)	0.89	4 (5%) 29 26	61, 67, 73, 77	0
25	13	59/60 (98%)	-0.06	1 (1%) 69 71	27, 38, 63, 74	0
25	23	59/60 (98%)	0.74	3 (5%) 33 31	51, 60, 77, 79	0
26	14	69/71 (97%)	1.22	15 (21%) 2 2	56, 76, 88, 95	0
26	24	69/71 (97%)	2.20	35 (50%) 0 0	78, 84, 90, 93	0
27	15	59/60 (98%)	-0.21	1 (1%) 69 71	25, 36, 54, 63	0
27	25	59/60 (98%)	0.11	0 100 100	35, 49, 65, 73	0
28	16	53/54 (98%)	-0.04	0 100 100	33, 43, 58, 60	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
28	26	53/54 (98%)	0.42	0 100 100	50, 59, 65, 71	0
29	17	48/49 (97%)	-0.14	4 (8%) 17 15	24, 31, 56, 61	0
29	27	48/49 (97%)	0.20	4 (8%) 17 15	35, 40, 64, 70	0
30	18	64/65 (98%)	-0.26	0 100 100	30, 36, 42, 53	0
30	28	64/65 (98%)	0.39	1 (1%) 70 73	45, 54, 61, 65	0
31	19	37/37 (100%)	0.07	0 100 100	33, 42, 54, 58	0
31	29	37/37 (100%)	0.96	4 (10%) 11 9	55, 64, 72, 74	0
32	1a	1488/1521 (97%)	0.28	62 (4%) 40 40	36, 67, 89, 100	0
32	2a	1492/1521 (98%)	0.62	106 (7%) 22 19	44, 73, 91, 102	0
33	1b	231/256 (90%)	1.41	52 (22%) 2 1	63, 75, 83, 88	0
33	2b	231/256 (90%)	1.77	79 (34%) 1 1	70, 80, 86, 91	0
34	1c	206/239 (86%)	0.95	16 (7%) 19 16	62, 70, 80, 84	0
34	2c	206/239 (86%)	1.85	77 (37%) 1 0	72, 81, 85, 92	0
35	1d	208/209 (99%)	1.10	32 (15%) 5 3	56, 69, 78, 84	0
35	2d	208/209 (99%)	1.13	18 (8%) 16 14	59, 69, 77, 82	0
36	1e	148/162 (91%)	0.65	2 (1%) 73 75	52, 63, 72, 80	0
36	2e	148/162 (91%)	1.08	11 (7%) 20 18	57, 70, 79, 85	0
37	1f	100/101 (99%)	0.67	3 (3%) 52 53	49, 64, 71, 79	0
37	2f	100/101 (99%)	0.71	1 (1%) 79 80	58, 67, 75, 82	0
38	1g	155/156 (99%)	0.68	3 (1%) 66 67	60, 69, 76, 81	0
38	2g	155/156 (99%)	1.54	39 (25%) 1 1	69, 79, 83, 87	0
39	1h	137/138 (99%)	0.71	5 (3%) 46 46	56, 66, 71, 78	0
39	2h	137/138 (99%)	0.95	6 (4%) 39 38	62, 71, 76, 79	0
40	1i	127/128 (99%)	1.39	23 (18%) 3 3	58, 75, 80, 82	0
40	2i	126/128 (98%)	2.40	79 (62%) 0 0	74, 82, 87, 89	0
41	1j	97/105 (92%)	1.58	28 (28%) 1 1	60, 75, 83, 89	0
41	2j	96/105 (91%)	2.66	68 (70%) 0 0	72, 82, 86, 91	0
42	1k	114/129 (88%)	0.60	4 (3%) 47 47	44, 62, 72, 77	0
42	2k	114/129 (88%)	0.86	6 (5%) 32 30	58, 70, 78, 81	0
43	1l	121/132 (91%)	0.68	9 (7%) 20 18	49, 59, 68, 76	0
43	2l	121/132 (91%)	0.74	6 (4%) 34 32	54, 63, 70, 75	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
44	1m	116/126 (92%)	1.11	18 (15%) 5 3	60, 72, 78, 80	0
44	2m	114/126 (90%)	1.99	53 (46%) 0 0	72, 81, 85, 87	0
45	1n	60/61 (98%)	1.12	6 (10%) 12 10	63, 68, 76, 79	0
45	2n	60/61 (98%)	2.48	39 (65%) 0 0	75, 80, 85, 89	0
46	1o	88/89 (98%)	0.86	5 (5%) 29 26	46, 63, 74, 76	0
46	2o	88/89 (98%)	1.03	8 (9%) 15 12	59, 69, 78, 81	0
47	1p	82/88 (93%)	1.41	16 (19%) 3 2	59, 69, 76, 83	0
47	2p	82/88 (93%)	1.26	9 (10%) 10 8	61, 68, 76, 79	0
48	1q	99/105 (94%)	1.02	4 (4%) 42 41	55, 67, 75, 80	0
48	2q	99/105 (94%)	0.73	5 (5%) 33 31	57, 67, 75, 79	0
49	1r	68/88 (77%)	0.62	2 (2%) 53 55	54, 61, 75, 82	0
49	2r	68/88 (77%)	0.91	4 (5%) 28 25	61, 69, 78, 85	0
50	1s	83/93 (89%)	1.17	8 (9%) 13 11	63, 74, 79, 82	0
50	2s	83/93 (89%)	2.26	48 (57%) 0 0	68, 82, 86, 89	0
51	1t	96/106 (90%)	1.34	20 (20%) 2 2	60, 70, 78, 82	0
51	2t	98/106 (92%)	0.97	7 (7%) 22 19	56, 67, 78, 80	0
52	1u	23/27 (85%)	1.51	6 (26%) 1 1	65, 69, 72, 74	0
52	2u	23/27 (85%)	2.48	15 (65%) 0 0	74, 79, 82, 82	0
53	1y	97/113 (85%)	0.68	7 (7%) 21 19	53, 62, 72, 76	0
53	2y	96/113 (84%)	2.39	62 (64%) 0 0	71, 80, 87, 89	0
54	1w	3/4 (75%)	0.19	0 100 100	30, 30, 40, 55	0
54	2w	3/4 (75%)	0.74	1 (33%) 1 1	41, 41, 48, 62	0
55	1x	3/4 (75%)	-0.45	0 100 100	28, 28, 30, 41	0
55	2x	3/4 (75%)	0.77	1 (33%) 1 1	37, 37, 40, 55	0
56	1v	3/3 (100%)	-0.46	0 100 100	25, 25, 27, 33	0
56	2v	3/3 (100%)	0.27	0 100 100	40, 40, 44, 51	0
All	All	20796/21490 (96%)	0.44	1806 (8%) 16 14	22, 62, 85, 102	0

The worst 5 of 1806 RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
45	2n	2	ALA	9.3
1	1A	1087	G	7.6

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Mol	Chain	Res	Type	RSRZ
1	2A	2147	G	6.5
41	2j	37	PRO	6.1
45	1n	2	ALA	6.1

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
1	5MU	2A	1915	21/22	0.82	0.12	79,87,90,101	0
32	M2G	2a	966	25/26	0.86	0.16	63,71,84,97	0
1	5MU	1A	1915	21/22	0.88	0.12	74,79,83,87	0
32	2MG	2a	1207	24/25	0.88	0.12	77,81,86,93	0
1	PSU	2A	1911	20/21	0.89	0.10	67,72,77,80	0
1	PSU	2A	1917	20/21	0.89	0.10	73,77,92,95	0
43	0TD	2l	92	10/11	0.90	0.12	59,66,69,74	0
1	PSU	1A	1917	20/21	0.91	0.11	65,69,75,77	0
32	5MC	2a	967	21/22	0.91	0.15	66,72,78,80	0
43	0TD	1l	92	10/11	0.92	0.13	49,59,64,67	0
32	PSU	2a	516	20/21	0.92	0.10	66,72,76,79	0
1	OMC	2A	1920	21/22	0.93	0.10	65,68,71,76	0
1	PSU	1A	1911	20/21	0.93	0.09	57,64,69,70	0
32	5MC	2a	1404	21/22	0.93	0.10	54,61,64,66	0
32	5MC	1a	967	21/22	0.93	0.11	57,66,73,82	0
32	5MC	2a	1400	21/22	0.94	0.13	62,75,78,80	0
32	4OC	2a	1402	22/23	0.94	0.11	59,64,71,73	0
32	G7M	2a	527	24/25	0.94	0.10	60,67,70,72	0
32	UR3	2a	1498	21/22	0.94	0.10	52,62,66,69	0
32	M2G	1a	966	25/26	0.94	0.10	55,61,68,71	0
32	5MC	2a	1407	21/22	0.95	0.10	56,64,67,69	0
32	2MG	1a	1207	24/25	0.95	0.09	62,70,74,78	0
32	MA6	2a	1518	24/25	0.95	0.10	58,64,68,69	0
32	PSU	1a	516	20/21	0.95	0.08	60,64,67,71	0
32	G7M	1a	527	24/25	0.96	0.10	48,55,59,67	0
1	OMC	1A	1920	21/22	0.96	0.09	50,57,61,63	0
32	5MC	1a	1400	21/22	0.96	0.10	49,57,60,65	0
32	4OC	1a	1402	22/23	0.96	0.09	46,53,56,64	0
32	5MC	1a	1407	21/22	0.96	0.09	44,54,58,63	0
1	5MC	2A	1942	21/22	0.96	0.09	44,49,54,60	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
32	UR3	1a	1498	21/22	0.96	0.08	42,50,53,59	0
32	MA6	1a	1518	24/25	0.96	0.10	43,47,52,53	0
32	MA6	2a	1519	24/25	0.96	0.12	49,63,67,69	0
32	MA6	1a	1519	24/25	0.96	0.10	42,48,52,54	0
32	5MC	1a	1404	21/22	0.97	0.08	45,48,51,56	0
1	5MU	2A	1939	21/22	0.97	0.08	35,39,42,46	0
1	2MA	1A	2503	23/24	0.97	0.07	20,24,27,29	0
1	OMG	2A	2251	24/25	0.97	0.08	38,40,43,45	0
1	2MA	2A	2503	23/24	0.97	0.08	29,34,37,39	0
1	5MC	1A	1942	21/22	0.97	0.08	33,39,44,47	0
1	5MC	1A	1962	21/22	0.98	0.07	30,36,40,48	0
1	OMU	2A	2552	21/22	0.98	0.07	33,40,43,46	0
1	PSU	2A	2605	20/21	0.98	0.08	32,38,43,44	0
1	OMG	1A	2251	24/25	0.98	0.07	22,28,32,43	0
1	5MU	1A	1939	21/22	0.98	0.07	27,31,36,39	0
1	OMU	1A	2552	21/22	0.98	0.06	26,31,33,35	0
1	5MC	2A	1962	21/22	0.98	0.07	33,46,53,56	0
1	PSU	1A	2605	20/21	0.98	0.07	27,30,33,36	0
54	PPU	1w	76	37/38	0.98	0.07	21,26,30,30	0
54	PPU	2w	76	37/38	0.98	0.07	30,36,41,44	0
55	8AN	1x	76	22/23	0.98	0.06	19,26,28,32	0
55	8AN	2x	76	22/23	0.98	0.07	33,36,39,44	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
57	MG	1A	3741	1/1	0.43	0.21	71,71,71,71	0
57	MG	2A	3542	1/1	0.49	0.32	70,70,70,70	0
57	MG	1a	1822	1/1	0.52	0.40	89,89,89,89	0
57	MG	1A	3968	1/1	0.53	0.21	40,40,40,40	0
57	MG	1A	3683	1/1	0.55	0.17	79,79,79,79	0
57	MG	2a	3176	1/1	0.55	0.36	74,74,74,74	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3319	1/1	0.56	0.33	83,83,83,83	0
57	MG	1a	1802	1/1	0.58	0.34	76,76,76,76	0
57	MG	2A	3707	1/1	0.60	0.23	86,86,86,86	0
57	MG	1A	3506	1/1	0.60	0.29	76,76,76,76	0
57	MG	2a	3128	1/1	0.61	0.23	74,74,74,74	0
57	MG	2A	3399	1/1	0.61	0.20	79,79,79,79	0
57	MG	1a	1835	1/1	0.63	0.31	67,67,67,67	0
57	MG	1A	3637	1/1	0.64	0.29	79,79,79,79	0
57	MG	2A	3121	1/1	0.64	0.34	79,79,79,79	0
57	MG	1a	1702	1/1	0.64	0.44	79,79,79,79	0
57	MG	2a	3131	1/1	0.64	0.23	76,76,76,76	0
57	MG	2A	3511	1/1	0.64	0.14	75,75,75,75	0
57	MG	1A	3874	1/1	0.65	0.25	45,45,45,45	0
57	MG	2A	3414	1/1	0.65	0.24	70,70,70,70	0
57	MG	2A	3474	1/1	0.65	0.22	64,64,64,64	0
57	MG	2A	3326	1/1	0.66	0.27	72,72,72,72	0
57	MG	2A	3540	1/1	0.66	0.17	70,70,70,70	0
57	MG	2G	202	1/1	0.66	0.26	80,80,80,80	0
57	MG	1A	3857	1/1	0.67	0.19	71,71,71,71	0
57	MG	1a	1867	1/1	0.68	0.23	76,76,76,76	0
57	MG	2A	3433	1/1	0.68	0.28	69,69,69,69	0
57	MG	2a	3089	1/1	0.68	0.37	83,83,83,83	0
57	MG	2A	3304	1/1	0.69	0.32	73,73,73,73	0
57	MG	1a	1798	1/1	0.69	0.16	77,77,77,77	0
57	MG	2a	3099	1/1	0.69	0.33	75,75,75,75	0
57	MG	1A	3981	1/1	0.69	0.27	73,73,73,73	0
57	MG	1a	1784	1/1	0.69	0.22	79,79,79,79	0
57	MG	2A	3172	1/1	0.69	0.27	80,80,80,80	0
57	MG	2A	3164	1/1	0.70	0.28	86,86,86,86	0
57	MG	1A	3501	1/1	0.70	0.23	69,69,69,69	0
57	MG	1a	1841	1/1	0.70	0.25	74,74,74,74	0
57	MG	1A	3518	1/1	0.70	0.15	58,58,58,58	0
57	MG	2A	3116	1/1	0.70	0.25	91,91,91,91	0
57	MG	1A	3554	1/1	0.70	0.28	65,65,65,65	0
57	MG	1A	3596	1/1	0.71	0.16	47,47,47,47	0
57	MG	1A	3602	1/1	0.71	0.22	71,71,71,71	0
57	MG	1a	1740	1/1	0.71	0.24	91,91,91,91	0
57	MG	1A	3913	1/1	0.71	0.24	64,64,64,64	0
57	MG	1A	3613	1/1	0.71	0.23	72,72,72,72	0
57	MG	2A	3467	1/1	0.72	0.21	66,66,66,66	0
57	MG	2a	3025	1/1	0.72	0.19	69,69,69,69	0
57	MG	1t	202	1/1	0.72	0.20	74,74,74,74	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1a	1612	1/1	0.72	0.34	90,90,90,90	0
57	MG	1a	1795	1/1	0.72	0.15	69,69,69,69	0
57	MG	1A	3589	1/1	0.72	0.23	66,66,66,66	0
57	MG	1A	3690	1/1	0.72	0.18	70,70,70,70	0
57	MG	2B	214	1/1	0.73	0.19	65,65,65,65	0
57	MG	1a	1718	1/1	0.73	0.34	77,77,77,77	0
57	MG	2a	3019	1/1	0.73	0.25	85,85,85,85	0
57	MG	1d	303	1/1	0.73	0.31	77,77,77,77	0
57	MG	2A	3502	1/1	0.73	0.23	58,58,58,58	0
57	MG	1O	201	1/1	0.73	0.16	61,61,61,61	0
57	MG	2a	3120	1/1	0.73	0.30	80,80,80,80	0
57	MG	1a	1773	1/1	0.73	0.20	75,75,75,75	0
57	MG	1A	3433	1/1	0.73	0.20	51,51,51,51	0
57	MG	1A	3528	1/1	0.73	0.18	57,57,57,57	0
57	MG	2B	218	1/1	0.74	0.20	85,85,85,85	0
57	MG	1A	3572	1/1	0.74	0.23	51,51,51,51	0
57	MG	2a	3009	1/1	0.74	0.51	81,81,81,81	0
57	MG	2A	3311	1/1	0.74	0.20	79,79,79,79	0
57	MG	1A	3327	1/1	0.74	0.18	61,61,61,61	0
57	MG	1A	3695	1/1	0.74	0.26	65,65,65,65	0
57	MG	1A	3698	1/1	0.74	0.45	45,45,45,45	0
57	MG	2A	3675	1/1	0.74	0.14	64,64,64,64	0
57	MG	2A	3698	1/1	0.74	0.25	65,65,65,65	0
57	MG	1A	3413	1/1	0.74	0.17	70,70,70,70	0
57	MG	1A	3674	1/1	0.74	0.21	69,69,69,69	0
57	MG	2A	3132	1/1	0.75	0.24	65,65,65,65	0
57	MG	2A	3159	1/1	0.75	0.44	59,59,59,59	0
57	MG	1a	1792	1/1	0.75	0.30	83,83,83,83	0
57	MG	2A	3047	1/1	0.75	0.44	79,79,79,79	0
57	MG	2A	3663	1/1	0.75	0.13	71,71,71,71	0
57	MG	2A	3445	1/1	0.75	0.36	76,76,76,76	0
57	MG	2A	3691	1/1	0.75	0.19	50,50,50,50	0
57	MG	1a	1678	1/1	0.75	0.33	78,78,78,78	0
57	MG	2A	3703	1/1	0.75	0.14	67,67,67,67	0
57	MG	1A	3380	1/1	0.75	0.17	57,57,57,57	0
57	MG	2a	3164	1/1	0.75	0.19	82,82,82,82	0
57	MG	2A	3491	1/1	0.75	0.28	72,72,72,72	0
57	MG	1A	3924	1/1	0.76	0.15	61,61,61,61	0
57	MG	2A	3726	1/1	0.76	0.24	64,64,64,64	0
57	MG	1A	3846	1/1	0.76	0.15	62,62,62,62	0
57	MG	1A	3708	1/1	0.76	0.18	41,41,41,41	0
57	MG	1A	3531	1/1	0.76	0.14	50,50,50,50	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3032	1/1	0.76	0.33	80,80,80,80	0
57	MG	1T	202	1/1	0.76	0.25	69,69,69,69	0
57	MG	2A	3075	1/1	0.76	0.32	74,74,74,74	0
57	MG	2a	3037	1/1	0.76	0.17	77,77,77,77	0
57	MG	2a	3043	1/1	0.76	0.24	74,74,74,74	0
57	MG	2a	3047	1/1	0.76	0.30	82,82,82,82	0
57	MG	2A	3655	1/1	0.76	0.12	77,77,77,77	0
57	MG	2A	3397	1/1	0.76	0.17	66,66,66,66	0
57	MG	2a	3109	1/1	0.76	0.25	85,85,85,85	0
57	MG	2A	3671	1/1	0.76	0.16	58,58,58,58	0
57	MG	2A	3082	1/1	0.76	0.21	78,78,78,78	0
57	MG	2A	3085	1/1	0.76	0.38	77,77,77,77	0
57	MG	1a	1827	1/1	0.76	0.20	77,77,77,77	0
57	MG	1A	3762	1/1	0.76	0.18	62,62,62,62	0
57	MG	2f	202	1/1	0.76	0.19	75,75,75,75	0
57	MG	1A	3618	1/1	0.77	0.15	43,43,43,43	0
57	MG	1A	3419	1/1	0.77	0.17	76,76,76,76	0
57	MG	2A	3495	1/1	0.77	0.20	71,71,71,71	0
57	MG	1a	1746	1/1	0.77	0.20	67,67,67,67	0
57	MG	2A	3403	1/1	0.77	0.19	78,78,78,78	0
57	MG	2A	3704	1/1	0.77	0.19	64,64,64,64	0
57	MG	2A	3525	1/1	0.77	0.22	75,75,75,75	0
57	MG	1a	1855	1/1	0.77	0.21	79,79,79,79	0
57	MG	1A	3783	1/1	0.77	0.23	67,67,67,67	0
57	MG	2A	3572	1/1	0.77	0.21	50,50,50,50	0
57	MG	2A	3624	1/1	0.77	0.19	71,71,71,71	0
57	MG	1A	3785	1/1	0.77	0.16	66,66,66,66	0
57	MG	2a	3178	1/1	0.77	0.34	69,69,69,69	0
57	MG	1n	101	1/1	0.77	0.36	78,78,78,78	0
57	MG	2j	201	1/1	0.77	0.23	79,79,79,79	0
57	MG	2A	3473	1/1	0.78	0.21	71,71,71,71	0
57	MG	1A	3799	1/1	0.78	0.21	30,30,30,30	0
57	MG	2A	3481	1/1	0.78	0.34	51,51,51,51	0
57	MG	1y	201	1/1	0.78	0.14	70,70,70,70	0
57	MG	2A	3345	1/1	0.78	0.21	67,67,67,67	0
57	MG	2A	3127	1/1	0.78	0.37	78,78,78,78	0
57	MG	2A	3699	1/1	0.78	0.22	65,65,65,65	0
57	MG	2a	3093	1/1	0.78	0.39	69,69,69,69	0
57	MG	2A	3700	1/1	0.78	0.19	70,70,70,70	0
57	MG	1a	1611	1/1	0.78	0.35	69,69,69,69	0
57	MG	2A	3146	1/1	0.78	0.25	70,70,70,70	0
57	MG	2a	3121	1/1	0.78	0.17	77,77,77,77	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1a	1787	1/1	0.78	0.30	75,75,75,75	0
57	MG	2A	3722	1/1	0.78	0.31	75,75,75,75	0
57	MG	2A	3541	1/1	0.78	0.23	79,79,79,79	0
57	MG	2a	3174	1/1	0.78	0.28	66,66,66,66	0
57	MG	1A	3556	1/1	0.78	0.14	56,56,56,56	0
57	MG	1d	305	1/1	0.78	0.12	85,85,85,85	0
57	MG	1A	3661	1/1	0.78	0.16	64,64,64,64	0
57	MG	2O	202	1/1	0.78	0.28	76,76,76,76	0
58	MPD	2B	220	8/8	0.78	0.28	63,66,74,77	0
59	ARG	1F	318	12/12	0.78	0.22	58,66,77,78	0
57	MG	2A	3551	1/1	0.79	0.15	61,61,61,61	0
57	MG	2Q	203	1/1	0.79	0.34	62,62,62,62	0
57	MG	1A	3121	1/1	0.79	0.31	62,62,62,62	0
57	MG	2A	3620	1/1	0.79	0.15	79,79,79,79	0
57	MG	2A	3415	1/1	0.79	0.16	53,53,53,53	0
57	MG	1A	3179	1/1	0.79	0.23	77,77,77,77	0
57	MG	1A	3424	1/1	0.79	0.19	53,53,53,53	0
57	MG	1A	3821	1/1	0.79	0.19	62,62,62,62	0
57	MG	2A	3471	1/1	0.79	0.16	69,69,69,69	0
57	MG	1A	3280	1/1	0.79	0.11	80,80,80,80	0
57	MG	1A	3986	1/1	0.79	0.19	52,52,52,52	0
57	MG	1a	1743	1/1	0.79	0.22	74,74,74,74	0
57	MG	1a	1831	1/1	0.79	0.26	80,80,80,80	0
57	MG	1B	217	1/1	0.79	0.14	63,63,63,63	0
57	MG	1A	3853	1/1	0.79	0.11	69,69,69,69	0
57	MG	2A	3377	1/1	0.79	0.17	72,72,72,72	0
57	MG	2A	3711	1/1	0.79	0.22	78,78,78,78	0
57	MG	2A	3517	1/1	0.79	0.24	77,77,77,77	0
57	MG	2A	3089	1/1	0.79	0.26	76,76,76,76	0
57	MG	2B	212	1/1	0.79	0.26	68,68,68,68	0
57	MG	1a	1775	1/1	0.79	0.27	72,72,72,72	0
57	MG	1A	3576	1/1	0.79	0.46	46,46,46,46	0
57	MG	2D	304	1/1	0.79	0.24	58,58,58,58	0
57	MG	2A	3413	1/1	0.79	0.21	77,77,77,77	0
57	MG	1A	3952	1/1	0.80	0.15	49,49,49,49	0
57	MG	1A	3160	1/1	0.80	0.19	52,52,52,52	0
57	MG	1A	3430	1/1	0.80	0.22	62,62,62,62	0
57	MG	1a	1642	1/1	0.80	0.21	67,67,67,67	0
57	MG	2A	3670	1/1	0.80	0.17	69,69,69,69	0
57	MG	1A	3038	1/1	0.80	0.17	53,53,53,53	0
57	MG	2a	3045	1/1	0.80	0.43	71,71,71,71	0
57	MG	1A	4009	1/1	0.80	0.22	68,68,68,68	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2a	3060	1/1	0.80	0.40	75,75,75,75	0
57	MG	2a	3080	1/1	0.80	0.19	62,62,62,62	0
57	MG	2A	3163	1/1	0.80	0.18	74,74,74,74	0
57	MG	1a	1711	1/1	0.80	0.41	76,76,76,76	0
57	MG	1A	3611	1/1	0.80	0.28	61,61,61,61	0
57	MG	2A	3193	1/1	0.80	0.41	73,73,73,73	0
57	MG	2A	3209	1/1	0.80	0.28	73,73,73,73	0
57	MG	2A	3219	1/1	0.80	0.25	69,69,69,69	0
57	MG	2A	3250	1/1	0.80	0.17	68,68,68,68	0
57	MG	1A	3781	1/1	0.80	0.21	59,59,59,59	0
57	MG	2a	3162	1/1	0.80	0.24	78,78,78,78	0
57	MG	2A	3045	1/1	0.80	0.13	67,67,67,67	0
57	MG	1a	1824	1/1	0.80	0.31	76,76,76,76	0
57	MG	1R	207	1/1	0.80	0.25	53,53,53,53	0
57	MG	1T	201	1/1	0.80	0.23	68,68,68,68	0
57	MG	1a	1758	1/1	0.80	0.31	71,71,71,71	0
57	MG	1a	1838	1/1	0.80	0.22	71,71,71,71	0
57	MG	2A	3111	1/1	0.80	0.20	58,58,58,58	0
57	MG	2A	3598	1/1	0.80	0.18	74,74,74,74	0
57	MG	1a	1675	1/1	0.81	0.20	60,60,60,60	0
57	MG	1A	3673	1/1	0.81	0.22	69,69,69,69	0
57	MG	2A	3723	1/1	0.81	0.23	57,57,57,57	0
57	MG	1A	3514	1/1	0.81	0.15	65,65,65,65	0
57	MG	2B	201	1/1	0.81	0.31	79,79,79,79	0
57	MG	1a	1703	1/1	0.81	0.20	72,72,72,72	0
57	MG	1A	3997	1/1	0.81	0.34	42,42,42,42	0
57	MG	1A	3646	1/1	0.81	0.17	54,54,54,54	0
57	MG	1a	1722	1/1	0.81	0.21	76,76,76,76	0
57	MG	1A	3873	1/1	0.81	0.11	57,57,57,57	0
57	MG	1a	1863	1/1	0.81	0.18	57,57,57,57	0
57	MG	2A	3516	1/1	0.81	0.10	79,79,79,79	0
57	MG	2A	3179	1/1	0.81	0.17	73,73,73,73	0
57	MG	2A	3182	1/1	0.81	0.29	73,73,73,73	0
57	MG	2a	3021	1/1	0.81	0.29	74,74,74,74	0
57	MG	2a	3023	1/1	0.81	0.40	79,79,79,79	0
57	MG	2A	3539	1/1	0.81	0.19	73,73,73,73	0
57	MG	1A	3211	1/1	0.81	0.14	64,64,64,64	0
57	MG	1P	202	1/1	0.81	0.22	43,43,43,43	0
57	MG	2A	3212	1/1	0.81	0.28	66,66,66,66	0
57	MG	1a	1748	1/1	0.81	0.30	72,72,72,72	0
57	MG	2A	3571	1/1	0.81	0.17	44,44,44,44	0
57	MG	1e	205	1/1	0.81	0.15	70,70,70,70	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3577	1/1	0.81	0.31	60,60,60,60	0
57	MG	1A	3816	1/1	0.81	0.26	37,37,37,37	0
57	MG	1a	1772	1/1	0.81	0.12	69,69,69,69	0
57	MG	1A	3691	1/1	0.81	0.20	73,73,73,73	0
57	MG	1A	3943	1/1	0.81	0.19	74,74,74,74	0
57	MG	2A	3037	1/1	0.81	0.29	69,69,69,69	0
57	MG	1A	3835	1/1	0.81	0.14	61,61,61,61	0
57	MG	1A	3953	1/1	0.81	0.15	71,71,71,71	0
57	MG	2a	3154	1/1	0.81	0.30	80,80,80,80	0
57	MG	2A	3051	1/1	0.81	0.22	68,68,68,68	0
57	MG	1A	3842	1/1	0.81	0.15	71,71,71,71	0
57	MG	1a	1647	1/1	0.81	0.27	70,70,70,70	0
57	MG	1a	1651	1/1	0.81	0.29	72,72,72,72	0
57	MG	2A	3419	1/1	0.81	0.15	62,62,62,62	0
57	MG	2A	3701	1/1	0.81	0.17	55,55,55,55	0
57	MG	2A	3426	1/1	0.81	0.19	68,68,68,68	0
57	MG	1a	1662	1/1	0.81	0.22	75,75,75,75	0
57	MG	1a	1819	1/1	0.81	0.27	80,80,80,80	0
57	MG	2A	3683	1/1	0.82	0.27	57,57,57,57	0
57	MG	2A	3381	1/1	0.82	0.19	55,55,55,55	0
57	MG	2A	3697	1/1	0.82	0.19	84,84,84,84	0
57	MG	2A	3026	1/1	0.82	0.21	70,70,70,70	0
57	MG	1A	3868	1/1	0.82	0.17	67,67,67,67	0
57	MG	1A	3632	1/1	0.82	0.28	45,45,45,45	0
57	MG	2A	3405	1/1	0.82	0.17	72,72,72,72	0
57	MG	1a	1603	1/1	0.82	0.22	72,72,72,72	0
57	MG	1A	3634	1/1	0.82	0.20	48,48,48,48	0
57	MG	1A	3752	1/1	0.82	0.21	69,69,69,69	0
57	MG	1A	3636	1/1	0.82	0.13	75,75,75,75	0
57	MG	1A	3567	1/1	0.82	0.20	50,50,50,50	0
57	MG	1A	3946	1/1	0.82	0.18	65,65,65,65	0
57	MG	2A	3441	1/1	0.82	0.18	70,70,70,70	0
57	MG	2A	3442	1/1	0.82	0.31	75,75,75,75	0
57	MG	1A	3366	1/1	0.82	0.26	77,77,77,77	0
57	MG	2A	3447	1/1	0.82	0.23	62,62,62,62	0
57	MG	2A	3449	1/1	0.82	0.18	92,92,92,92	0
57	MG	1A	3178	1/1	0.82	0.18	55,55,55,55	0
57	MG	1A	3663	1/1	0.82	0.12	52,52,52,52	0
57	MG	1A	3977	1/1	0.82	0.26	55,55,55,55	0
57	MG	1a	1830	1/1	0.82	0.23	77,77,77,77	0
57	MG	2U	201	1/1	0.82	0.16	62,62,62,62	0
57	MG	1A	3582	1/1	0.82	0.16	37,37,37,37	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3134	1/1	0.82	0.20	68,68,68,68	0
57	MG	1A	3985	1/1	0.82	0.18	72,72,72,72	0
57	MG	1a	1713	1/1	0.82	0.36	60,60,60,60	0
57	MG	1A	3323	1/1	0.82	0.39	74,74,74,74	0
57	MG	2a	3026	1/1	0.82	0.36	75,75,75,75	0
57	MG	2A	3514	1/1	0.82	0.18	68,68,68,68	0
57	MG	1a	1842	1/1	0.82	0.18	88,88,88,88	0
57	MG	1a	1719	1/1	0.82	0.18	58,58,58,58	0
57	MG	1a	1857	1/1	0.82	0.23	59,59,59,59	0
57	MG	2a	3053	1/1	0.82	0.20	67,67,67,67	0
57	MG	1A	3444	1/1	0.82	0.15	42,42,42,42	0
57	MG	1A	3458	1/1	0.82	0.21	42,42,42,42	0
57	MG	2A	3204	1/1	0.82	0.33	67,67,67,67	0
57	MG	1A	3544	1/1	0.82	0.12	48,48,48,48	0
57	MG	2A	3544	1/1	0.82	0.12	64,64,64,64	0
57	MG	2A	3545	1/1	0.82	0.21	78,78,78,78	0
57	MG	2A	3549	1/1	0.82	0.28	65,65,65,65	0
57	MG	1B	220	1/1	0.82	0.17	69,69,69,69	0
57	MG	1e	201	1/1	0.82	0.22	76,76,76,76	0
57	MG	1e	202	1/1	0.82	0.36	73,73,73,73	0
57	MG	2A	3573	1/1	0.82	0.14	47,47,47,47	0
57	MG	2A	3289	1/1	0.82	0.35	65,65,65,65	0
57	MG	2A	3292	1/1	0.82	0.33	76,76,76,76	0
57	MG	2a	3171	1/1	0.82	0.24	75,75,75,75	0
57	MG	2A	3298	1/1	0.82	0.47	78,78,78,78	0
57	MG	1e	204	1/1	0.82	0.22	70,70,70,70	0
57	MG	2a	3177	1/1	0.82	0.34	61,61,61,61	0
57	MG	1A	3467	1/1	0.82	0.14	52,52,52,52	0
57	MG	1a	1750	1/1	0.82	0.15	82,82,82,82	0
57	MG	2A	3334	1/1	0.82	0.19	64,64,64,64	0
58	MPD	1T	208	8/8	0.82	0.22	68,70,74,81	0
58	MPD	1a	1868	8/8	0.82	0.17	53,64,70,74	0
57	MG	1A	3027	1/1	0.82	0.13	70,70,70,70	0
57	MG	1A	3865	1/1	0.82	0.12	51,51,51,51	0
57	MG	2B	207	1/1	0.83	0.14	67,67,67,67	0
57	MG	1A	3775	1/1	0.83	0.18	45,45,45,45	0
57	MG	2A	3069	1/1	0.83	0.37	65,65,65,65	0
57	MG	2B	216	1/1	0.83	0.15	83,83,83,83	0
57	MG	2A	3307	1/1	0.83	0.18	65,65,65,65	0
57	MG	2A	3531	1/1	0.83	0.19	77,77,77,77	0
57	MG	1R	203	1/1	0.83	0.29	47,47,47,47	0
57	MG	2A	3312	1/1	0.83	0.15	39,39,39,39	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1a	1714	1/1	0.83	0.20	72,72,72,72	0
57	MG	1a	1717	1/1	0.83	0.28	81,81,81,81	0
57	MG	1A	3776	1/1	0.83	0.19	45,45,45,45	0
57	MG	1A	3508	1/1	0.83	0.20	62,62,62,62	0
57	MG	2A	3379	1/1	0.83	0.18	70,70,70,70	0
57	MG	2A	3550	1/1	0.83	0.13	73,73,73,73	0
57	MG	1A	3511	1/1	0.83	0.14	35,35,35,35	0
57	MG	1a	1732	1/1	0.83	0.15	85,85,85,85	0
57	MG	18	102	1/1	0.83	0.24	40,40,40,40	0
57	MG	1A	3641	1/1	0.83	0.14	45,45,45,45	0
57	MG	1A	3448	1/1	0.83	0.11	49,49,49,49	0
57	MG	1A	3992	1/1	0.83	0.16	60,60,60,60	0
57	MG	1a	1632	1/1	0.83	0.27	63,63,63,63	0
57	MG	1a	1638	1/1	0.83	0.26	67,67,67,67	0
57	MG	2a	3063	1/1	0.83	0.25	78,78,78,78	0
57	MG	1a	1761	1/1	0.83	0.20	70,70,70,70	0
57	MG	2A	3420	1/1	0.83	0.20	55,55,55,55	0
57	MG	2a	3091	1/1	0.83	0.33	69,69,69,69	0
57	MG	1A	3656	1/1	0.83	0.21	75,75,75,75	0
57	MG	2A	3178	1/1	0.83	0.21	65,65,65,65	0
57	MG	1A	4002	1/1	0.83	0.16	71,71,71,71	0
57	MG	2A	3677	1/1	0.83	0.32	56,56,56,56	0
57	MG	1A	3252	1/1	0.83	0.14	47,47,47,47	0
57	MG	1a	1658	1/1	0.83	0.14	69,69,69,69	0
57	MG	1A	3376	1/1	0.83	0.11	65,65,65,65	0
57	MG	1A	3434	1/1	0.83	0.26	63,63,63,63	0
57	MG	2a	3157	1/1	0.83	0.19	73,73,73,73	0
57	MG	2A	3456	1/1	0.83	0.17	78,78,78,78	0
57	MG	1E	304	1/1	0.83	0.14	28,28,28,28	0
57	MG	2A	3217	1/1	0.83	0.12	69,69,69,69	0
57	MG	1a	1701	1/1	0.83	0.25	72,72,72,72	0
57	MG	1G	202	1/1	0.83	0.13	60,60,60,60	0
57	MG	2A	3253	1/1	0.83	0.27	75,75,75,75	0
57	MG	2A	3259	1/1	0.83	0.29	57,57,57,57	0
57	MG	1A	3263	1/1	0.83	0.48	77,77,77,77	0
57	MG	2A	3290	1/1	0.83	0.47	68,68,68,68	0
57	MG	1a	1709	1/1	0.83	0.21	60,60,60,60	0
57	MG	2A	3729	1/1	0.83	0.18	69,69,69,69	0
57	MG	2A	3295	1/1	0.83	0.38	65,65,65,65	0
57	MG	2B	202	1/1	0.83	0.16	75,75,75,75	0
57	MG	1e	203	1/1	0.84	0.14	61,61,61,61	0
57	MG	2G	201	1/1	0.84	0.21	77,77,77,77	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3660	1/1	0.84	0.29	36,36,36,36	0
57	MG	2A	3546	1/1	0.84	0.23	66,66,66,66	0
57	MG	1A	3465	1/1	0.84	0.12	62,62,62,62	0
57	MG	1A	3875	1/1	0.84	0.23	62,62,62,62	0
57	MG	2a	3004	1/1	0.84	0.19	76,76,76,76	0
57	MG	15	105	1/1	0.84	0.23	35,35,35,35	0
57	MG	2a	3010	1/1	0.84	0.38	81,81,81,81	0
57	MG	1A	3525	1/1	0.84	0.21	64,64,64,64	0
57	MG	1a	1810	1/1	0.84	0.20	72,72,72,72	0
57	MG	1a	1816	1/1	0.84	0.12	73,73,73,73	0
57	MG	2A	3034	1/1	0.84	0.16	58,58,58,58	0
57	MG	1A	3321	1/1	0.84	0.38	70,70,70,70	0
57	MG	2A	3612	1/1	0.84	0.18	44,44,44,44	0
57	MG	2A	3614	1/1	0.84	0.16	38,38,38,38	0
57	MG	2A	3232	1/1	0.84	0.27	69,69,69,69	0
57	MG	2A	3249	1/1	0.84	0.33	69,69,69,69	0
57	MG	1A	3926	1/1	0.84	0.13	69,69,69,69	0
57	MG	1B	206	1/1	0.84	0.12	53,53,53,53	0
57	MG	1B	207	1/1	0.84	0.24	69,69,69,69	0
57	MG	2a	3073	1/1	0.84	0.40	71,71,71,71	0
57	MG	2A	3451	1/1	0.84	0.20	64,64,64,64	0
57	MG	2A	3452	1/1	0.84	0.17	58,58,58,58	0
57	MG	2A	3454	1/1	0.84	0.32	82,82,82,82	0
57	MG	2A	3286	1/1	0.84	0.31	67,67,67,67	0
57	MG	2A	3461	1/1	0.84	0.18	56,56,56,56	0
57	MG	2a	3107	1/1	0.84	0.31	80,80,80,80	0
57	MG	1A	3063	1/1	0.84	0.13	52,52,52,52	0
57	MG	2a	3117	1/1	0.84	0.15	69,69,69,69	0
57	MG	1A	3677	1/1	0.84	0.14	83,83,83,83	0
57	MG	1D	313	1/1	0.84	0.25	33,33,33,33	0
57	MG	1A	3272	1/1	0.84	0.13	67,67,67,67	0
57	MG	1a	1657	1/1	0.84	0.12	72,72,72,72	0
57	MG	2A	3483	1/1	0.84	0.18	64,64,64,64	0
57	MG	2A	3301	1/1	0.84	0.21	70,70,70,70	0
57	MG	2A	3492	1/1	0.84	0.19	78,78,78,78	0
57	MG	2A	3105	1/1	0.84	0.22	65,65,65,65	0
57	MG	1A	3856	1/1	0.84	0.13	45,45,45,45	0
57	MG	1N	203	1/1	0.84	0.13	61,61,61,61	0
57	MG	1A	3454	1/1	0.84	0.15	55,55,55,55	0
57	MG	1A	3330	1/1	0.84	0.13	58,58,58,58	0
57	MG	1a	1866	1/1	0.84	0.29	72,72,72,72	0
57	MG	2a	3181	1/1	0.84	0.26	70,70,70,70	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1a	1769	1/1	0.84	0.13	69,69,69,69	0
57	MG	1a	1691	1/1	0.84	0.30	69,69,69,69	0
57	MG	1a	1692	1/1	0.84	0.40	64,64,64,64	0
57	MG	1a	1697	1/1	0.84	0.27	69,69,69,69	0
57	MG	2A	3394	1/1	0.84	0.16	75,75,75,75	0
57	MG	1A	3463	1/1	0.84	0.13	46,46,46,46	0
57	MG	1A	3476	1/1	0.85	0.10	24,24,24,24	0
57	MG	2A	3206	1/1	0.85	0.17	67,67,67,67	0
57	MG	1B	210	1/1	0.85	0.16	60,60,60,60	0
57	MG	2A	3728	1/1	0.85	0.15	56,56,56,56	0
57	MG	2A	3468	1/1	0.85	0.14	65,65,65,65	0
57	MG	1B	213	1/1	0.85	0.25	69,69,69,69	0
57	MG	1a	1768	1/1	0.85	0.17	62,62,62,62	0
57	MG	2B	204	1/1	0.85	0.21	75,75,75,75	0
57	MG	2B	205	1/1	0.85	0.32	72,72,72,72	0
57	MG	1a	1652	1/1	0.85	0.32	74,74,74,74	0
57	MG	2B	210	1/1	0.85	0.13	66,66,66,66	0
57	MG	2A	3228	1/1	0.85	0.46	73,73,73,73	0
57	MG	1A	3823	1/1	0.85	0.14	50,50,50,50	0
57	MG	1A	3442	1/1	0.85	0.13	55,55,55,55	0
57	MG	1D	309	1/1	0.85	0.33	47,47,47,47	0
57	MG	1a	1674	1/1	0.85	0.18	69,69,69,69	0
57	MG	2A	3499	1/1	0.85	0.20	78,78,78,78	0
57	MG	1A	3093	1/1	0.85	0.17	57,57,57,57	0
57	MG	2A	3505	1/1	0.85	0.11	71,71,71,71	0
57	MG	2A	3264	1/1	0.85	0.35	60,60,60,60	0
57	MG	1A	3206	1/1	0.85	0.26	68,68,68,68	0
57	MG	1E	307	1/1	0.85	0.14	62,62,62,62	0
57	MG	1A	3852	1/1	0.85	0.14	41,41,41,41	0
57	MG	2A	3521	1/1	0.85	0.24	66,66,66,66	0
57	MG	1A	3010	1/1	0.85	0.16	52,52,52,52	0
57	MG	1A	3568	1/1	0.85	0.17	52,52,52,52	0
57	MG	2A	3532	1/1	0.85	0.16	67,67,67,67	0
57	MG	1A	3151	1/1	0.85	0.25	62,62,62,62	0
57	MG	2A	3300	1/1	0.85	0.27	66,66,66,66	0
57	MG	2a	3034	1/1	0.85	0.39	68,68,68,68	0
57	MG	2A	3058	1/1	0.85	0.15	67,67,67,67	0
57	MG	2a	3039	1/1	0.85	0.18	76,76,76,76	0
57	MG	2A	3062	1/1	0.85	0.15	56,56,56,56	0
57	MG	1A	3863	1/1	0.85	0.17	35,35,35,35	0
57	MG	1a	1821	1/1	0.85	0.14	68,68,68,68	0
57	MG	1a	1708	1/1	0.85	0.37	70,70,70,70	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2a	3058	1/1	0.85	0.26	79,79,79,79	0
57	MG	2A	3084	1/1	0.85	0.32	74,74,74,74	0
57	MG	1A	3982	1/1	0.85	0.21	50,50,50,50	0
57	MG	2a	3064	1/1	0.85	0.26	71,71,71,71	0
57	MG	1A	3155	1/1	0.85	0.29	71,71,71,71	0
57	MG	2A	3373	1/1	0.85	0.15	44,44,44,44	0
57	MG	1A	3040	1/1	0.85	0.18	64,64,64,64	0
57	MG	1l	104	1/1	0.85	0.15	64,64,64,64	0
57	MG	2a	3092	1/1	0.85	0.20	60,60,60,60	0
57	MG	1A	3786	1/1	0.85	0.23	65,65,65,65	0
57	MG	2A	3385	1/1	0.85	0.15	78,78,78,78	0
57	MG	2a	3101	1/1	0.85	0.25	61,61,61,61	0
57	MG	2a	3102	1/1	0.85	0.30	66,66,66,66	0
57	MG	2A	3389	1/1	0.85	0.18	76,76,76,76	0
57	MG	2A	3120	1/1	0.85	0.16	65,65,65,65	0
57	MG	2A	3619	1/1	0.85	0.17	59,59,59,59	0
57	MG	1A	3035	1/1	0.85	0.15	54,54,54,54	0
57	MG	1A	4001	1/1	0.85	0.17	34,34,34,34	0
57	MG	2A	3651	1/1	0.85	0.12	76,76,76,76	0
57	MG	1A	3529	1/1	0.85	0.11	56,56,56,56	0
57	MG	2a	3148	1/1	0.85	0.14	69,69,69,69	0
57	MG	1a	1844	1/1	0.85	0.12	69,69,69,69	0
57	MG	2A	3139	1/1	0.85	0.21	55,55,55,55	0
57	MG	1a	1846	1/1	0.85	0.21	64,64,64,64	0
57	MG	2A	3150	1/1	0.85	0.34	63,63,63,63	0
57	MG	2a	3167	1/1	0.85	0.21	72,72,72,72	0
57	MG	2A	3155	1/1	0.85	0.10	66,66,66,66	0
57	MG	1a	1724	1/1	0.85	0.20	63,63,63,63	0
57	MG	1A	3876	1/1	0.85	0.16	55,55,55,55	0
57	MG	1a	1861	1/1	0.85	0.21	74,74,74,74	0
57	MG	1a	1621	1/1	0.85	0.25	61,61,61,61	0
57	MG	2A	3175	1/1	0.85	0.17	56,56,56,56	0
57	MG	1A	4011	1/1	0.85	0.23	59,59,59,59	0
57	MG	1a	1635	1/1	0.85	0.29	65,65,65,65	0
57	MG	2r	101	1/1	0.85	0.18	67,67,67,67	0
57	MG	1b	301	1/1	0.85	0.27	78,78,78,78	0
57	MG	2A	3191	1/1	0.85	0.37	71,71,71,71	0
58	MPD	2A	3732	8/8	0.85	0.22	51,60,66,69	0
57	MG	1A	3890	1/1	0.85	0.27	37,37,37,37	0
57	MG	2A	3199	1/1	0.85	0.13	59,59,59,59	0
57	MG	1A	3659	1/1	0.86	0.12	51,51,51,51	0
57	MG	1a	1694	1/1	0.86	0.40	67,67,67,67	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3500	1/1	0.86	0.19	63,63,63,63	0
57	MG	1F	314	1/1	0.86	0.10	42,42,42,42	0
57	MG	1A	3372	1/1	0.86	0.17	74,74,74,74	0
57	MG	1A	3585	1/1	0.86	0.18	72,72,72,72	0
57	MG	1A	3373	1/1	0.86	0.21	53,53,53,53	0
57	MG	1a	1706	1/1	0.86	0.31	62,62,62,62	0
57	MG	1A	3672	1/1	0.86	0.16	69,69,69,69	0
57	MG	1A	3485	1/1	0.86	0.10	24,24,24,24	0
57	MG	2A	3108	1/1	0.86	0.21	72,72,72,72	0
57	MG	2A	3308	1/1	0.86	0.21	64,64,64,64	0
57	MG	1A	3535	1/1	0.86	0.24	78,78,78,78	0
57	MG	1a	1840	1/1	0.86	0.13	76,76,76,76	0
57	MG	2A	3325	1/1	0.86	0.19	65,65,65,65	0
57	MG	1A	3223	1/1	0.86	0.15	54,54,54,54	0
57	MG	1A	3549	1/1	0.86	0.11	39,39,39,39	0
57	MG	2A	3335	1/1	0.86	0.14	65,65,65,65	0
57	MG	2a	3032	1/1	0.86	0.25	73,73,73,73	0
57	MG	2A	3340	1/1	0.86	0.17	75,75,75,75	0
57	MG	1W	201	1/1	0.86	0.28	55,55,55,55	0
57	MG	1A	3824	1/1	0.86	0.11	54,54,54,54	0
57	MG	2a	3040	1/1	0.86	0.33	72,72,72,72	0
57	MG	2A	3133	1/1	0.86	0.34	65,65,65,65	0
57	MG	13	102	1/1	0.86	0.12	73,73,73,73	0
57	MG	2A	3553	1/1	0.86	0.13	39,39,39,39	0
57	MG	2A	3380	1/1	0.86	0.14	47,47,47,47	0
57	MG	1A	3120	1/1	0.86	0.24	55,55,55,55	0
57	MG	1a	1858	1/1	0.86	0.26	77,77,77,77	0
57	MG	1A	3326	1/1	0.86	0.17	45,45,45,45	0
57	MG	1A	3558	1/1	0.86	0.14	55,55,55,55	0
57	MG	2A	3600	1/1	0.86	0.14	53,53,53,53	0
57	MG	2a	3077	1/1	0.86	0.36	69,69,69,69	0
57	MG	1A	3849	1/1	0.86	0.10	57,57,57,57	0
57	MG	2a	3082	1/1	0.86	0.38	66,66,66,66	0
57	MG	2A	3398	1/1	0.86	0.17	66,66,66,66	0
57	MG	1A	3417	1/1	0.86	0.11	22,22,22,22	0
57	MG	1a	1618	1/1	0.86	0.14	62,62,62,62	0
57	MG	2A	3165	1/1	0.86	0.18	60,60,60,60	0
57	MG	2A	3640	1/1	0.86	0.13	41,41,41,41	0
57	MG	1A	3281	1/1	0.86	0.19	46,46,46,46	0
57	MG	2A	3173	1/1	0.86	0.12	52,52,52,52	0
57	MG	2a	3105	1/1	0.86	0.30	65,65,65,65	0
57	MG	1A	3713	1/1	0.86	0.11	44,44,44,44	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1a	1754	1/1	0.86	0.28	67,67,67,67	0
57	MG	1A	3715	1/1	0.86	0.17	59,59,59,59	0
57	MG	1A	3730	1/1	0.86	0.12	54,54,54,54	0
57	MG	2A	3183	1/1	0.86	0.34	78,78,78,78	0
57	MG	2a	3127	1/1	0.86	0.15	68,68,68,68	0
57	MG	1A	3317	1/1	0.86	0.20	52,52,52,52	0
57	MG	2a	3130	1/1	0.86	0.29	73,73,73,73	0
57	MG	1A	3746	1/1	0.86	0.23	50,50,50,50	0
57	MG	2a	3134	1/1	0.86	0.22	77,77,77,77	0
57	MG	1A	3575	1/1	0.86	0.14	54,54,54,54	0
57	MG	2A	3446	1/1	0.86	0.26	47,47,47,47	0
57	MG	2a	3156	1/1	0.86	0.14	84,84,84,84	0
57	MG	1n	102	1/1	0.86	0.15	53,53,53,53	0
57	MG	1n	103	1/1	0.86	0.29	67,67,67,67	0
57	MG	1A	3756	1/1	0.86	0.24	68,68,68,68	0
57	MG	2A	3210	1/1	0.86	0.31	65,65,65,65	0
57	MG	1A	3759	1/1	0.86	0.15	59,59,59,59	0
57	MG	2A	3001	1/1	0.86	0.34	65,65,65,65	0
57	MG	2A	3016	1/1	0.86	0.29	76,76,76,76	0
57	MG	2A	3463	1/1	0.86	0.16	41,41,41,41	0
57	MG	1B	218	1/1	0.86	0.17	53,53,53,53	0
57	MG	2A	3230	1/1	0.86	0.23	67,67,67,67	0
57	MG	1A	3652	1/1	0.86	0.10	43,43,43,43	0
57	MG	1a	1669	1/1	0.86	0.17	66,66,66,66	0
57	MG	1D	301	1/1	0.86	0.17	65,65,65,65	0
57	MG	2A	3480	1/1	0.86	0.16	56,56,56,56	0
57	MG	1A	3193	1/1	0.86	0.42	64,64,64,64	0
57	MG	1A	3892	1/1	0.86	0.13	28,28,28,28	0
57	MG	1A	3907	1/1	0.86	0.14	29,29,29,29	0
57	MG	2A	3272	1/1	0.86	0.25	78,78,78,78	0
57	MG	1a	1860	1/1	0.87	0.21	68,68,68,68	0
57	MG	1F	316	1/1	0.87	0.20	53,53,53,53	0
57	MG	2F	301	1/1	0.87	0.15	66,66,66,66	0
57	MG	1A	3124	1/1	0.87	0.21	63,63,63,63	0
57	MG	1A	3446	1/1	0.87	0.13	83,83,83,83	0
57	MG	2I	201	1/1	0.87	0.14	74,74,74,74	0
57	MG	1a	1663	1/1	0.87	0.13	61,61,61,61	0
57	MG	2A	3529	1/1	0.87	0.14	75,75,75,75	0
57	MG	2R	201	1/1	0.87	0.26	51,51,51,51	0
57	MG	1a	1759	1/1	0.87	0.21	67,67,67,67	0
57	MG	29	101	1/1	0.87	0.14	66,66,66,66	0
57	MG	2A	3336	1/1	0.87	0.21	60,60,60,60	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3533	1/1	0.87	0.21	68,68,68,68	0
57	MG	1A	3171	1/1	0.87	0.19	68,68,68,68	0
57	MG	2a	3016	1/1	0.87	0.27	80,80,80,80	0
57	MG	1a	1763	1/1	0.87	0.19	65,65,65,65	0
57	MG	2A	3366	1/1	0.87	0.18	64,64,64,64	0
57	MG	2A	3161	1/1	0.87	0.27	74,74,74,74	0
57	MG	1A	3083	1/1	0.87	0.13	55,55,55,55	0
57	MG	1A	3062	1/1	0.87	0.20	68,68,68,68	0
57	MG	1a	1770	1/1	0.87	0.26	67,67,67,67	0
57	MG	2A	3547	1/1	0.87	0.17	80,80,80,80	0
57	MG	2A	3170	1/1	0.87	0.14	52,52,52,52	0
57	MG	1A	3246	1/1	0.87	0.12	68,68,68,68	0
57	MG	1a	1681	1/1	0.87	0.32	72,72,72,72	0
57	MG	2A	3391	1/1	0.87	0.16	65,65,65,65	0
57	MG	2A	3561	1/1	0.87	0.09	63,63,63,63	0
57	MG	1l	3101	1/1	0.87	0.17	73,73,73,73	0
57	MG	1a	1686	1/1	0.87	0.24	72,72,72,72	0
57	MG	2a	3055	1/1	0.87	0.14	68,68,68,68	0
57	MG	2a	3056	1/1	0.87	0.28	77,77,77,77	0
57	MG	2a	3057	1/1	0.87	0.30	68,68,68,68	0
57	MG	1a	1688	1/1	0.87	0.18	63,63,63,63	0
57	MG	1A	3431	1/1	0.87	0.12	32,32,32,32	0
57	MG	1o	101	1/1	0.87	0.27	73,73,73,73	0
57	MG	2A	3186	1/1	0.87	0.20	74,74,74,74	0
57	MG	2a	3065	1/1	0.87	0.12	65,65,65,65	0
57	MG	2a	3066	1/1	0.87	0.30	65,65,65,65	0
57	MG	2A	3406	1/1	0.87	0.12	74,74,74,74	0
57	MG	2a	3074	1/1	0.87	0.12	75,75,75,75	0
57	MG	1A	3700	1/1	0.87	0.24	60,60,60,60	0
57	MG	1A	3891	1/1	0.87	0.09	20,20,20,20	0
57	MG	2a	3081	1/1	0.87	0.37	75,75,75,75	0
57	MG	2A	3196	1/1	0.87	0.21	64,64,64,64	0
57	MG	2A	3418	1/1	0.87	0.21	56,56,56,56	0
57	MG	1a	1796	1/1	0.87	0.09	80,80,80,80	0
57	MG	2A	3012	1/1	0.87	0.18	68,68,68,68	0
57	MG	10	107	1/1	0.87	0.17	65,65,65,65	0
57	MG	2a	3098	1/1	0.87	0.18	63,63,63,63	0
57	MG	2A	3024	1/1	0.87	0.11	51,51,51,51	0
57	MG	2A	3668	1/1	0.87	0.20	57,57,57,57	0
57	MG	1A	3378	1/1	0.87	0.14	44,44,44,44	0
57	MG	1a	1808	1/1	0.87	0.23	72,72,72,72	0
57	MG	2A	3444	1/1	0.87	0.24	70,70,70,70	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3296	1/1	0.87	0.18	66,66,66,66	0
57	MG	2a	3110	1/1	0.87	0.12	69,69,69,69	0
57	MG	2a	3114	1/1	0.87	0.15	59,59,59,59	0
57	MG	1A	3480	1/1	0.87	0.15	39,39,39,39	0
57	MG	2A	3222	1/1	0.87	0.12	65,65,65,65	0
57	MG	1a	1704	1/1	0.87	0.30	66,66,66,66	0
57	MG	17	105	1/1	0.87	0.19	59,59,59,59	0
57	MG	1A	3408	1/1	0.87	0.14	68,68,68,68	0
57	MG	1a	1601	1/1	0.87	0.13	80,80,80,80	0
57	MG	1A	3740	1/1	0.87	0.17	65,65,65,65	0
57	MG	1a	1829	1/1	0.87	0.12	46,46,46,46	0
57	MG	1A	3490	1/1	0.87	0.09	71,71,71,71	0
57	MG	1A	3546	1/1	0.87	0.17	58,58,58,58	0
57	MG	2a	3155	1/1	0.87	0.24	66,66,66,66	0
57	MG	2A	3270	1/1	0.87	0.16	64,64,64,64	0
57	MG	2A	3713	1/1	0.87	0.25	68,68,68,68	0
57	MG	2a	3161	1/1	0.87	0.17	70,70,70,70	0
57	MG	2A	3719	1/1	0.87	0.21	58,58,58,58	0
57	MG	1B	225	1/1	0.87	0.09	53,53,53,53	0
57	MG	2A	3278	1/1	0.87	0.14	55,55,55,55	0
57	MG	2a	3169	1/1	0.87	0.21	72,72,72,72	0
57	MG	1A	3667	1/1	0.87	0.18	72,72,72,72	0
57	MG	2A	3288	1/1	0.87	0.39	62,62,62,62	0
57	MG	1A	3548	1/1	0.87	0.27	40,40,40,40	0
57	MG	2A	3090	1/1	0.87	0.30	66,66,66,66	0
57	MG	2A	3093	1/1	0.87	0.10	57,57,57,57	0
57	MG	2a	3179	1/1	0.87	0.29	77,77,77,77	0
57	MG	1A	3757	1/1	0.87	0.17	50,50,50,50	0
57	MG	1a	1723	1/1	0.87	0.30	68,68,68,68	0
57	MG	1D	315	1/1	0.87	0.18	74,74,74,74	0
57	MG	2p	101	1/1	0.87	0.26	61,61,61,61	0
57	MG	1A	3500	1/1	0.87	0.12	58,58,58,58	0
57	MG	2x	101	1/1	0.87	0.15	44,44,44,44	0
57	MG	1A	3979	1/1	0.87	0.13	85,85,85,85	0
57	MG	2B	213	1/1	0.87	0.29	74,74,74,74	0
57	MG	1F	312	1/1	0.87	0.13	37,37,37,37	0
57	MG	2B	215	1/1	0.87	0.14	68,68,68,68	0
57	MG	1A	3760	1/1	0.87	0.17	50,50,50,50	0
57	MG	1A	3958	1/1	0.88	0.12	50,50,50,50	0
57	MG	2Q	202	1/1	0.88	0.38	65,65,65,65	0
57	MG	2A	3060	1/1	0.88	0.30	54,54,54,54	0
57	MG	1A	3963	1/1	0.88	0.13	60,60,60,60	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3130	1/1	0.88	0.13	60,60,60,60	0
57	MG	1a	1805	1/1	0.88	0.28	71,71,71,71	0
57	MG	2A	3078	1/1	0.88	0.35	63,63,63,63	0
57	MG	2a	3005	1/1	0.88	0.31	73,73,73,73	0
57	MG	1A	3361	1/1	0.88	0.11	39,39,39,39	0
57	MG	1H	201	1/1	0.88	0.10	72,72,72,72	0
57	MG	1A	3850	1/1	0.88	0.12	49,49,49,49	0
57	MG	2A	3088	1/1	0.88	0.16	65,65,65,65	0
57	MG	2A	3535	1/1	0.88	0.18	37,37,37,37	0
57	MG	1A	3980	1/1	0.88	0.18	58,58,58,58	0
57	MG	2A	3310	1/1	0.88	0.15	64,64,64,64	0
57	MG	1A	3287	1/1	0.88	0.18	54,54,54,54	0
57	MG	2a	3031	1/1	0.88	0.21	69,69,69,69	0
57	MG	1a	1695	1/1	0.88	0.30	66,66,66,66	0
57	MG	2a	3033	1/1	0.88	0.32	64,64,64,64	0
57	MG	2A	3321	1/1	0.88	0.16	61,61,61,61	0
57	MG	2a	3036	1/1	0.88	0.19	67,67,67,67	0
57	MG	1A	3474	1/1	0.88	0.09	28,28,28,28	0
57	MG	2a	3038	1/1	0.88	0.23	65,65,65,65	0
57	MG	1a	1825	1/1	0.88	0.11	67,67,67,67	0
57	MG	1A	3184	1/1	0.88	0.15	55,55,55,55	0
57	MG	1A	3479	1/1	0.88	0.12	30,30,30,30	0
57	MG	1A	3987	1/1	0.88	0.22	77,77,77,77	0
57	MG	2a	3046	1/1	0.88	0.22	73,73,73,73	0
57	MG	2A	3339	1/1	0.88	0.19	73,73,73,73	0
57	MG	2A	3552	1/1	0.88	0.10	59,59,59,59	0
57	MG	1V	204	1/1	0.88	0.20	64,64,64,64	0
57	MG	1A	3603	1/1	0.88	0.14	59,59,59,59	0
57	MG	2A	3348	1/1	0.88	0.18	61,61,61,61	0
57	MG	2A	3130	1/1	0.88	0.14	54,54,54,54	0
57	MG	2a	3059	1/1	0.88	0.37	70,70,70,70	0
57	MG	1A	3993	1/1	0.88	0.26	70,70,70,70	0
57	MG	1A	3312	1/1	0.88	0.19	64,64,64,64	0
57	MG	1A	3866	1/1	0.88	0.12	40,40,40,40	0
57	MG	1A	3676	1/1	0.88	0.11	74,74,74,74	0
57	MG	2A	3608	1/1	0.88	0.13	36,36,36,36	0
57	MG	2a	3070	1/1	0.88	0.37	62,62,62,62	0
57	MG	1A	3059	1/1	0.88	0.21	44,44,44,44	0
57	MG	1a	1716	1/1	0.88	0.13	70,70,70,70	0
57	MG	2A	3617	1/1	0.88	0.12	82,82,82,82	0
57	MG	2A	3618	1/1	0.88	0.14	31,31,31,31	0
57	MG	1A	3197	1/1	0.88	0.34	68,68,68,68	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1B	203	1/1	0.88	0.17	68,68,68,68	0
57	MG	2A	3623	1/1	0.88	0.15	66,66,66,66	0
57	MG	1a	1602	1/1	0.88	0.18	74,74,74,74	0
57	MG	1a	1859	1/1	0.88	0.33	71,71,71,71	0
57	MG	1A	3685	1/1	0.88	0.17	72,72,72,72	0
57	MG	1A	3626	1/1	0.88	0.12	23,23,23,23	0
57	MG	1A	3200	1/1	0.88	0.27	63,63,63,63	0
57	MG	2A	3171	1/1	0.88	0.15	53,53,53,53	0
57	MG	1a	1615	1/1	0.88	0.13	63,63,63,63	0
57	MG	1a	1735	1/1	0.88	0.23	70,70,70,70	0
57	MG	1a	1736	1/1	0.88	0.20	72,72,72,72	0
57	MG	1B	212	1/1	0.88	0.28	72,72,72,72	0
57	MG	1A	3555	1/1	0.88	0.20	60,60,60,60	0
57	MG	1A	3788	1/1	0.88	0.20	64,64,64,64	0
57	MG	2A	3694	1/1	0.88	0.26	65,65,65,65	0
57	MG	1A	3898	1/1	0.88	0.17	57,57,57,57	0
57	MG	1A	3383	1/1	0.88	0.17	62,62,62,62	0
57	MG	2A	3429	1/1	0.88	0.23	69,69,69,69	0
57	MG	1a	1753	1/1	0.88	0.16	67,67,67,67	0
57	MG	1a	1639	1/1	0.88	0.41	74,74,74,74	0
57	MG	2A	3194	1/1	0.88	0.16	62,62,62,62	0
57	MG	1h	202	1/1	0.88	0.15	67,67,67,67	0
57	MG	1a	1755	1/1	0.88	0.14	47,47,47,47	0
57	MG	1A	3808	1/1	0.88	0.18	73,73,73,73	0
57	MG	1B	230	1/1	0.88	0.10	72,72,72,72	0
57	MG	1A	3014	1/1	0.88	0.27	62,62,62,62	0
57	MG	1A	3279	1/1	0.88	0.09	75,75,75,75	0
57	MG	2A	3211	1/1	0.88	0.32	70,70,70,70	0
57	MG	1o	102	1/1	0.88	0.24	68,68,68,68	0
57	MG	2a	3163	1/1	0.88	0.19	81,81,81,81	0
57	MG	1a	1653	1/1	0.88	0.29	70,70,70,70	0
57	MG	2a	3165	1/1	0.88	0.24	67,67,67,67	0
57	MG	1A	3711	1/1	0.88	0.35	40,40,40,40	0
57	MG	1A	3208	1/1	0.88	0.11	58,58,58,58	0
57	MG	2a	3170	1/1	0.88	0.27	59,59,59,59	0
57	MG	1A	3459	1/1	0.88	0.08	23,23,23,23	0
57	MG	2A	3229	1/1	0.88	0.17	65,65,65,65	0
57	MG	1A	3461	1/1	0.88	0.20	63,63,63,63	0
57	MG	1a	1664	1/1	0.88	0.26	68,68,68,68	0
57	MG	2B	208	1/1	0.88	0.18	69,69,69,69	0
57	MG	2A	3235	1/1	0.88	0.19	72,72,72,72	0
57	MG	1a	1777	1/1	0.88	0.18	75,75,75,75	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2a	3183	1/1	0.88	0.21	70,70,70,70	0
57	MG	1a	1667	1/1	0.88	0.39	65,65,65,65	0
57	MG	1a	1668	1/1	0.88	0.16	58,58,58,58	0
57	MG	2A	3035	1/1	0.88	0.28	70,70,70,70	0
57	MG	1a	1791	1/1	0.88	0.17	66,66,66,66	0
57	MG	1E	308	1/1	0.88	0.09	36,36,36,36	0
57	MG	1a	1793	1/1	0.88	0.18	66,66,66,66	0
57	MG	2A	3276	1/1	0.88	0.30	69,69,69,69	0
57	MG	2A	3050	1/1	0.88	0.15	61,61,61,61	0
57	MG	1F	311	1/1	0.88	0.12	55,55,55,55	0
59	ARG	1B	231	12/12	0.88	0.14	40,47,58,61	0
57	MG	2A	3508	1/1	0.88	0.17	71,71,71,71	0
57	MG	1a	1727	1/1	0.89	0.13	54,54,54,54	0
57	MG	2A	3189	1/1	0.89	0.14	48,48,48,48	0
57	MG	1a	1728	1/1	0.89	0.16	77,77,77,77	0
57	MG	1A	3533	1/1	0.89	0.14	69,69,69,69	0
57	MG	2A	3458	1/1	0.89	0.32	64,64,64,64	0
57	MG	1A	3623	1/1	0.89	0.12	53,53,53,53	0
57	MG	2E	304	1/1	0.89	0.13	43,43,43,43	0
57	MG	1A	3728	1/1	0.89	0.11	64,64,64,64	0
57	MG	2F	303	1/1	0.89	0.10	49,49,49,49	0
57	MG	1A	3534	1/1	0.89	0.33	57,57,57,57	0
57	MG	1a	1741	1/1	0.89	0.16	67,67,67,67	0
57	MG	1B	204	1/1	0.89	0.21	60,60,60,60	0
57	MG	2O	201	1/1	0.89	0.14	66,66,66,66	0
57	MG	1a	1624	1/1	0.89	0.20	60,60,60,60	0
57	MG	1i	201	1/1	0.89	0.18	68,68,68,68	0
57	MG	2A	3475	1/1	0.89	0.30	60,60,60,60	0
57	MG	1a	1630	1/1	0.89	0.37	67,67,67,67	0
57	MG	2T	202	1/1	0.89	0.22	68,68,68,68	0
57	MG	1A	3328	1/1	0.89	0.16	62,62,62,62	0
57	MG	2A	3216	1/1	0.89	0.14	62,62,62,62	0
57	MG	2A	3485	1/1	0.89	0.12	59,59,59,59	0
57	MG	1A	3542	1/1	0.89	0.09	36,36,36,36	0
57	MG	1A	3870	1/1	0.89	0.09	27,27,27,27	0
57	MG	1A	3871	1/1	0.89	0.08	37,37,37,37	0
57	MG	2a	3011	1/1	0.89	0.18	65,65,65,65	0
57	MG	2a	3015	1/1	0.89	0.23	61,61,61,61	0
57	MG	2A	3223	1/1	0.89	0.12	62,62,62,62	0
57	MG	1a	1640	1/1	0.89	0.21	69,69,69,69	0
57	MG	1A	3267	1/1	0.89	0.14	76,76,76,76	0
57	MG	1a	1645	1/1	0.89	0.30	63,63,63,63	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1a	1646	1/1	0.89	0.27	60,60,60,60	0
57	MG	1a	1765	1/1	0.89	0.14	61,61,61,61	0
57	MG	2A	3239	1/1	0.89	0.23	58,58,58,58	0
57	MG	1a	1767	1/1	0.89	0.18	53,53,53,53	0
57	MG	2A	3017	1/1	0.89	0.15	56,56,56,56	0
57	MG	2A	3252	1/1	0.89	0.17	65,65,65,65	0
57	MG	1A	3298	1/1	0.89	0.30	82,82,82,82	0
57	MG	1a	1648	1/1	0.89	0.26	68,68,68,68	0
57	MG	2A	3530	1/1	0.89	0.13	72,72,72,72	0
57	MG	2A	3261	1/1	0.89	0.13	53,53,53,53	0
57	MG	1A	3639	1/1	0.89	0.16	39,39,39,39	0
57	MG	2A	3266	1/1	0.89	0.46	63,63,63,63	0
57	MG	2a	3044	1/1	0.89	0.33	71,71,71,71	0
57	MG	2A	3033	1/1	0.89	0.30	68,68,68,68	0
57	MG	1A	3421	1/1	0.89	0.10	67,67,67,67	0
57	MG	1A	3645	1/1	0.89	0.10	29,29,29,29	0
57	MG	1A	3232	1/1	0.89	0.21	76,76,76,76	0
57	MG	2A	3281	1/1	0.89	0.29	72,72,72,72	0
57	MG	1A	3369	1/1	0.89	0.10	32,32,32,32	0
57	MG	2A	3046	1/1	0.89	0.20	63,63,63,63	0
57	MG	1A	3002	1/1	0.89	0.11	52,52,52,52	0
57	MG	1D	311	1/1	0.89	0.21	43,43,43,43	0
57	MG	1a	1790	1/1	0.89	0.16	74,74,74,74	0
57	MG	2a	3062	1/1	0.89	0.12	83,83,83,83	0
57	MG	2A	3053	1/1	0.89	0.26	73,73,73,73	0
57	MG	1A	3482	1/1	0.89	0.11	32,32,32,32	0
57	MG	2A	3299	1/1	0.89	0.34	75,75,75,75	0
57	MG	1a	1666	1/1	0.89	0.27	74,74,74,74	0
57	MG	2a	3068	1/1	0.89	0.17	71,71,71,71	0
57	MG	2A	3061	1/1	0.89	0.10	46,46,46,46	0
57	MG	2a	3071	1/1	0.89	0.22	69,69,69,69	0
57	MG	1A	3778	1/1	0.89	0.17	63,63,63,63	0
57	MG	1A	3318	1/1	0.89	0.31	65,65,65,65	0
57	MG	2A	3074	1/1	0.89	0.19	55,55,55,55	0
57	MG	1A	3560	1/1	0.89	0.13	57,57,57,57	0
57	MG	2A	3582	1/1	0.89	0.18	63,63,63,63	0
57	MG	1A	3929	1/1	0.89	0.12	44,44,44,44	0
57	MG	1a	1801	1/1	0.89	0.24	58,58,58,58	0
57	MG	1A	3247	1/1	0.89	0.18	64,64,64,64	0
57	MG	1A	3944	1/1	0.89	0.11	57,57,57,57	0
57	MG	1A	3377	1/1	0.89	0.14	63,63,63,63	0
57	MG	2a	3097	1/1	0.89	0.33	60,60,60,60	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3187	1/1	0.89	0.31	68,68,68,68	0
57	MG	1a	1811	1/1	0.89	0.25	70,70,70,70	0
57	MG	1A	3504	1/1	0.89	0.09	53,53,53,53	0
57	MG	2A	3099	1/1	0.89	0.37	66,66,66,66	0
57	MG	1A	3956	1/1	0.89	0.12	48,48,48,48	0
57	MG	1a	1820	1/1	0.89	0.11	60,60,60,60	0
57	MG	1A	3282	1/1	0.89	0.19	67,67,67,67	0
57	MG	2A	3352	1/1	0.89	0.12	51,51,51,51	0
57	MG	2a	3111	1/1	0.89	0.16	71,71,71,71	0
57	MG	2A	3112	1/1	0.89	0.22	63,63,63,63	0
57	MG	2A	3662	1/1	0.89	0.20	58,58,58,58	0
57	MG	2a	3118	1/1	0.89	0.12	62,62,62,62	0
57	MG	2A	3113	1/1	0.89	0.27	72,72,72,72	0
57	MG	1A	3962	1/1	0.89	0.11	54,54,54,54	0
57	MG	2a	3122	1/1	0.89	0.15	63,63,63,63	0
57	MG	1A	3283	1/1	0.89	0.11	62,62,62,62	0
57	MG	1a	1696	1/1	0.89	0.37	71,71,71,71	0
57	MG	1A	3449	1/1	0.89	0.15	51,51,51,51	0
57	MG	2A	3676	1/1	0.89	0.14	65,65,65,65	0
57	MG	2a	3132	1/1	0.89	0.22	62,62,62,62	0
57	MG	2A	3384	1/1	0.89	0.15	49,49,49,49	0
57	MG	2A	3678	1/1	0.89	0.07	77,77,77,77	0
57	MG	2A	3682	1/1	0.89	0.14	59,59,59,59	0
57	MG	1a	1698	1/1	0.89	0.39	68,68,68,68	0
57	MG	1a	1699	1/1	0.89	0.20	70,70,70,70	0
57	MG	1A	3452	1/1	0.89	0.14	71,71,71,71	0
57	MG	1A	3591	1/1	0.89	0.12	39,39,39,39	0
57	MG	2A	3136	1/1	0.89	0.20	66,66,66,66	0
57	MG	1a	1837	1/1	0.89	0.29	72,72,72,72	0
57	MG	1A	3832	1/1	0.89	0.12	53,53,53,53	0
57	MG	1T	206	1/1	0.89	0.09	71,71,71,71	0
57	MG	2A	3153	1/1	0.89	0.41	62,62,62,62	0
57	MG	2A	3154	1/1	0.89	0.18	52,52,52,52	0
57	MG	1A	3386	1/1	0.89	0.11	29,29,29,29	0
57	MG	1A	3837	1/1	0.89	0.17	56,56,56,56	0
57	MG	1A	3838	1/1	0.89	0.11	37,37,37,37	0
57	MG	2a	3175	1/1	0.89	0.12	65,65,65,65	0
57	MG	10	108	1/1	0.89	0.13	64,64,64,64	0
57	MG	1a	1848	1/1	0.89	0.17	65,65,65,65	0
57	MG	1a	1850	1/1	0.89	0.21	70,70,70,70	0
57	MG	2A	3422	1/1	0.89	0.20	78,78,78,78	0
57	MG	2A	3727	1/1	0.89	0.19	69,69,69,69	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3600	1/1	0.89	0.15	31,31,31,31	0
57	MG	1A	3456	1/1	0.89	0.12	53,53,53,53	0
57	MG	2A	3730	1/1	0.89	0.29	68,68,68,68	0
57	MG	1A	3387	1/1	0.89	0.08	24,24,24,24	0
57	MG	1A	3610	1/1	0.89	0.11	56,56,56,56	0
57	MG	1A	3996	1/1	0.89	0.12	58,58,58,58	0
57	MG	1A	3405	1/1	0.89	0.23	69,69,69,69	0
57	MG	1A	3999	1/1	0.89	0.09	47,47,47,47	0
57	MG	1A	3174	1/1	0.89	0.42	62,62,62,62	0
57	MG	2B	209	1/1	0.89	0.30	71,71,71,71	0
57	MG	1a	1604	1/1	0.89	0.29	72,72,72,72	0
57	MG	2A	3185	1/1	0.89	0.25	66,66,66,66	0
57	MG	2A	3493	1/1	0.90	0.18	60,60,60,60	0
57	MG	1A	3435	1/1	0.90	0.13	54,54,54,54	0
57	MG	1a	1605	1/1	0.90	0.16	64,64,64,64	0
57	MG	1A	3436	1/1	0.90	0.08	61,61,61,61	0
57	MG	1a	1712	1/1	0.90	0.25	63,63,63,63	0
57	MG	1A	3764	1/1	0.90	0.10	27,27,27,27	0
57	MG	1A	3886	1/1	0.90	0.10	63,63,63,63	0
57	MG	2A	3510	1/1	0.90	0.13	55,55,55,55	0
57	MG	20	101	1/1	0.90	0.16	63,63,63,63	0
57	MG	25	101	1/1	0.90	0.18	62,62,62,62	0
57	MG	25	102	1/1	0.90	0.18	70,70,70,70	0
57	MG	1A	3584	1/1	0.90	0.11	47,47,47,47	0
57	MG	1A	3437	1/1	0.90	0.24	68,68,68,68	0
57	MG	1A	3588	1/1	0.90	0.12	49,49,49,49	0
57	MG	2a	3006	1/1	0.90	0.12	55,55,55,55	0
57	MG	2A	3092	1/1	0.90	0.21	68,68,68,68	0
57	MG	1A	3530	1/1	0.90	0.12	30,30,30,30	0
57	MG	2A	3097	1/1	0.90	0.28	58,58,58,58	0
57	MG	2A	3098	1/1	0.90	0.36	72,72,72,72	0
57	MG	1A	3290	1/1	0.90	0.16	61,61,61,61	0
57	MG	2a	3018	1/1	0.90	0.29	68,68,68,68	0
57	MG	2A	3291	1/1	0.90	0.39	65,65,65,65	0
57	MG	1A	3592	1/1	0.90	0.12	69,69,69,69	0
57	MG	1A	3915	1/1	0.90	0.14	55,55,55,55	0
57	MG	2A	3110	1/1	0.90	0.23	63,63,63,63	0
57	MG	2A	3536	1/1	0.90	0.10	67,67,67,67	0
57	MG	1A	3918	1/1	0.90	0.12	63,63,63,63	0
57	MG	1A	3595	1/1	0.90	0.10	47,47,47,47	0
57	MG	1A	3411	1/1	0.90	0.22	56,56,56,56	0
57	MG	1a	1643	1/1	0.90	0.19	59,59,59,59	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3793	1/1	0.90	0.11	43,43,43,43	0
57	MG	1a	1738	1/1	0.90	0.15	67,67,67,67	0
57	MG	2A	3125	1/1	0.90	0.18	85,85,85,85	0
57	MG	1A	3680	1/1	0.90	0.13	37,37,37,37	0
57	MG	2A	3548	1/1	0.90	0.25	73,73,73,73	0
57	MG	2a	3041	1/1	0.90	0.31	61,61,61,61	0
57	MG	2a	3042	1/1	0.90	0.22	78,78,78,78	0
57	MG	1A	3322	1/1	0.90	0.26	62,62,62,62	0
57	MG	2A	3313	1/1	0.90	0.30	66,66,66,66	0
57	MG	1A	3811	1/1	0.90	0.20	48,48,48,48	0
57	MG	2A	3322	1/1	0.90	0.18	75,75,75,75	0
57	MG	1a	1745	1/1	0.90	0.17	66,66,66,66	0
57	MG	2a	3049	1/1	0.90	0.37	66,66,66,66	0
57	MG	2a	3052	1/1	0.90	0.16	74,74,74,74	0
57	MG	1A	3947	1/1	0.90	0.15	63,63,63,63	0
57	MG	1A	3684	1/1	0.90	0.13	68,68,68,68	0
57	MG	2A	3137	1/1	0.90	0.09	45,45,45,45	0
57	MG	2A	3138	1/1	0.90	0.12	63,63,63,63	0
57	MG	1A	3819	1/1	0.90	0.10	57,57,57,57	0
57	MG	2A	3143	1/1	0.90	0.25	57,57,57,57	0
57	MG	2A	3588	1/1	0.90	0.15	50,50,50,50	0
57	MG	1A	3447	1/1	0.90	0.18	58,58,58,58	0
57	MG	2A	3148	1/1	0.90	0.12	69,69,69,69	0
57	MG	2A	3602	1/1	0.90	0.10	68,68,68,68	0
57	MG	1d	301	1/1	0.90	0.23	73,73,73,73	0
57	MG	2A	3355	1/1	0.90	0.14	65,65,65,65	0
57	MG	1F	313	1/1	0.90	0.30	45,45,45,45	0
57	MG	2A	3367	1/1	0.90	0.19	70,70,70,70	0
57	MG	1a	1661	1/1	0.90	0.17	58,58,58,58	0
57	MG	2A	3376	1/1	0.90	0.09	40,40,40,40	0
57	MG	1A	3144	1/1	0.90	0.09	69,69,69,69	0
57	MG	2A	3378	1/1	0.90	0.09	40,40,40,40	0
57	MG	2A	3157	1/1	0.90	0.13	64,64,64,64	0
57	MG	2A	3629	1/1	0.90	0.18	71,71,71,71	0
57	MG	1A	3543	1/1	0.90	0.11	52,52,52,52	0
57	MG	2a	3086	1/1	0.90	0.27	78,78,78,78	0
57	MG	2A	3644	1/1	0.90	0.12	68,68,68,68	0
57	MG	2a	3090	1/1	0.90	0.45	67,67,67,67	0
57	MG	1A	3486	1/1	0.90	0.08	57,57,57,57	0
57	MG	2A	3162	1/1	0.90	0.24	65,65,65,65	0
57	MG	1a	1762	1/1	0.90	0.09	60,60,60,60	0
57	MG	1a	1665	1/1	0.90	0.25	65,65,65,65	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1G	204	1/1	0.90	0.23	65,65,65,65	0
57	MG	2A	3169	1/1	0.90	0.21	56,56,56,56	0
57	MG	1a	1766	1/1	0.90	0.20	71,71,71,71	0
57	MG	1A	3696	1/1	0.90	0.19	44,44,44,44	0
57	MG	2a	3103	1/1	0.90	0.16	61,61,61,61	0
57	MG	1l	3102	1/1	0.90	0.18	75,75,75,75	0
57	MG	1A	3418	1/1	0.90	0.13	28,28,28,28	0
57	MG	2a	3108	1/1	0.90	0.12	52,52,52,52	0
57	MG	1A	3493	1/1	0.90	0.12	34,34,34,34	0
57	MG	2A	3679	1/1	0.90	0.12	77,77,77,77	0
57	MG	1a	1671	1/1	0.90	0.21	62,62,62,62	0
57	MG	2A	3409	1/1	0.90	0.10	56,56,56,56	0
57	MG	1A	3705	1/1	0.90	0.10	60,60,60,60	0
57	MG	1A	3499	1/1	0.90	0.11	63,63,63,63	0
57	MG	1R	205	1/1	0.90	0.17	56,56,56,56	0
57	MG	1a	1679	1/1	0.90	0.17	77,77,77,77	0
57	MG	1a	1680	1/1	0.90	0.11	63,63,63,63	0
57	MG	2a	3124	1/1	0.90	0.14	77,77,77,77	0
57	MG	2A	3187	1/1	0.90	0.26	53,53,53,53	0
57	MG	2A	3188	1/1	0.90	0.17	63,63,63,63	0
57	MG	2A	3423	1/1	0.90	0.23	77,77,77,77	0
57	MG	1A	3550	1/1	0.90	0.11	58,58,58,58	0
57	MG	1a	1788	1/1	0.90	0.13	66,66,66,66	0
57	MG	2a	3133	1/1	0.90	0.12	66,66,66,66	0
57	MG	2A	3710	1/1	0.90	0.19	62,62,62,62	0
57	MG	2a	3142	1/1	0.90	0.09	51,51,51,51	0
57	MG	2A	3430	1/1	0.90	0.14	59,59,59,59	0
57	MG	2a	3150	1/1	0.90	0.20	71,71,71,71	0
57	MG	2a	3153	1/1	0.90	0.15	60,60,60,60	0
57	MG	2A	3431	1/1	0.90	0.14	41,41,41,41	0
57	MG	2A	3718	1/1	0.90	0.36	63,63,63,63	0
57	MG	1a	1682	1/1	0.90	0.23	68,68,68,68	0
57	MG	1A	3094	1/1	0.90	0.10	57,57,57,57	0
57	MG	1A	3191	1/1	0.90	0.14	61,61,61,61	0
57	MG	2A	3724	1/1	0.90	0.19	69,69,69,69	0
57	MG	2A	3031	1/1	0.90	0.26	56,56,56,56	0
57	MG	2A	3201	1/1	0.90	0.32	74,74,74,74	0
57	MG	1a	1689	1/1	0.90	0.27	65,65,65,65	0
57	MG	1A	3315	1/1	0.90	0.17	58,58,58,58	0
57	MG	2a	3168	1/1	0.90	0.28	64,64,64,64	0
57	MG	2A	3207	1/1	0.90	0.36	53,53,53,53	0
57	MG	1A	3234	1/1	0.90	0.29	57,57,57,57	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3268	1/1	0.90	0.10	62,62,62,62	0
57	MG	2B	203	1/1	0.90	0.21	77,77,77,77	0
57	MG	1A	3153	1/1	0.90	0.28	52,52,52,52	0
57	MG	2A	3039	1/1	0.90	0.17	49,49,49,49	0
57	MG	2A	3041	1/1	0.90	0.31	72,72,72,72	0
57	MG	1A	3643	1/1	0.90	0.13	56,56,56,56	0
57	MG	1A	3512	1/1	0.90	0.10	24,24,24,24	0
57	MG	1A	3753	1/1	0.90	0.12	44,44,44,44	0
57	MG	1A	3571	1/1	0.90	0.11	22,22,22,22	0
57	MG	2e	201	1/1	0.90	0.25	73,73,73,73	0
57	MG	1A	4003	1/1	0.90	0.10	29,29,29,29	0
57	MG	1a	1813	1/1	0.90	0.21	70,70,70,70	0
57	MG	2A	3057	1/1	0.90	0.12	61,61,61,61	0
57	MG	1A	3320	1/1	0.90	0.25	64,64,64,64	0
57	MG	1a	1818	1/1	0.90	0.14	68,68,68,68	0
58	MPD	1A	4014	8/8	0.90	0.27	55,58,65,69	0
57	MG	2A	3237	1/1	0.90	0.16	46,46,46,46	0
57	MG	1A	3464	1/1	0.90	0.16	57,57,57,57	0
57	MG	1A	4012	1/1	0.90	0.17	64,64,64,64	0
57	MG	2A	3489	1/1	0.90	0.10	73,73,73,73	0
57	MG	2A	3065	1/1	0.90	0.42	48,48,48,48	0
57	MG	1B	202	1/1	0.90	0.22	60,60,60,60	0
57	MG	23	101	1/1	0.91	0.14	59,59,59,59	0
57	MG	1A	3266	1/1	0.91	0.12	56,56,56,56	0
57	MG	1B	221	1/1	0.91	0.09	50,50,50,50	0
57	MG	2A	3303	1/1	0.91	0.25	67,67,67,67	0
57	MG	1A	3803	1/1	0.91	0.09	40,40,40,40	0
57	MG	1B	228	1/1	0.91	0.08	58,58,58,58	0
57	MG	2A	3128	1/1	0.91	0.21	60,60,60,60	0
57	MG	1a	1739	1/1	0.91	0.10	60,60,60,60	0
57	MG	1a	1862	1/1	0.91	0.27	67,67,67,67	0
57	MG	1A	3382	1/1	0.91	0.24	61,61,61,61	0
57	MG	1A	3709	1/1	0.91	0.12	44,44,44,44	0
57	MG	1a	1644	1/1	0.91	0.22	61,61,61,61	0
57	MG	1A	3815	1/1	0.91	0.08	16,16,16,16	0
57	MG	1A	3932	1/1	0.91	0.08	44,44,44,44	0
57	MG	1A	3352	1/1	0.91	0.07	36,36,36,36	0
57	MG	1A	3353	1/1	0.91	0.18	62,62,62,62	0
57	MG	1A	3133	1/1	0.91	0.11	51,51,51,51	0
57	MG	1A	3722	1/1	0.91	0.14	60,60,60,60	0
57	MG	1A	3657	1/1	0.91	0.11	26,26,26,26	0
57	MG	1a	1655	1/1	0.91	0.21	60,60,60,60	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3826	1/1	0.91	0.08	47,47,47,47	0
57	MG	1f	201	1/1	0.91	0.37	72,72,72,72	0
57	MG	2A	3562	1/1	0.91	0.20	57,57,57,57	0
57	MG	1g	202	1/1	0.91	0.21	68,68,68,68	0
57	MG	1h	201	1/1	0.91	0.28	59,59,59,59	0
57	MG	2A	3362	1/1	0.91	0.11	48,48,48,48	0
57	MG	1A	3401	1/1	0.91	0.15	29,29,29,29	0
57	MG	1A	3731	1/1	0.91	0.15	55,55,55,55	0
57	MG	1A	3959	1/1	0.91	0.14	55,55,55,55	0
57	MG	2A	3590	1/1	0.91	0.08	46,46,46,46	0
57	MG	1A	3738	1/1	0.91	0.20	52,52,52,52	0
57	MG	2A	3599	1/1	0.91	0.10	41,41,41,41	0
57	MG	1A	3363	1/1	0.91	0.08	52,52,52,52	0
57	MG	2A	3166	1/1	0.91	0.17	68,68,68,68	0
57	MG	2A	3603	1/1	0.91	0.13	59,59,59,59	0
57	MG	2a	3051	1/1	0.91	0.27	69,69,69,69	0
57	MG	1A	3964	1/1	0.91	0.12	58,58,58,58	0
57	MG	1A	3841	1/1	0.91	0.20	59,59,59,59	0
57	MG	1A	3973	1/1	0.91	0.20	52,52,52,52	0
57	MG	2A	3383	1/1	0.91	0.11	34,34,34,34	0
57	MG	1A	3974	1/1	0.91	0.15	44,44,44,44	0
57	MG	1A	3141	1/1	0.91	0.18	64,64,64,64	0
57	MG	1P	203	1/1	0.91	0.16	72,72,72,72	0
57	MG	1A	3742	1/1	0.91	0.21	59,59,59,59	0
57	MG	2A	3003	1/1	0.91	0.11	64,64,64,64	0
57	MG	2A	3396	1/1	0.91	0.28	70,70,70,70	0
57	MG	2A	3638	1/1	0.91	0.14	58,58,58,58	0
57	MG	1A	3604	1/1	0.91	0.17	35,35,35,35	0
57	MG	1a	1778	1/1	0.91	0.15	72,72,72,72	0
57	MG	2A	3184	1/1	0.91	0.10	60,60,60,60	0
57	MG	1a	1782	1/1	0.91	0.16	71,71,71,71	0
57	MG	2A	3659	1/1	0.91	0.25	79,79,79,79	0
57	MG	2A	3660	1/1	0.91	0.11	42,42,42,42	0
57	MG	1A	3751	1/1	0.91	0.08	50,50,50,50	0
57	MG	1A	3609	1/1	0.91	0.15	32,32,32,32	0
57	MG	2A	3667	1/1	0.91	0.18	54,54,54,54	0
57	MG	1A	3669	1/1	0.91	0.08	61,61,61,61	0
57	MG	2A	3669	1/1	0.91	0.12	53,53,53,53	0
57	MG	2A	3411	1/1	0.91	0.10	35,35,35,35	0
57	MG	2a	3088	1/1	0.91	0.44	74,74,74,74	0
57	MG	1A	3755	1/1	0.91	0.11	59,59,59,59	0
57	MG	2A	3190	1/1	0.91	0.16	74,74,74,74	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3367	1/1	0.91	0.09	42,42,42,42	0
57	MG	2A	3192	1/1	0.91	0.16	66,66,66,66	0
57	MG	1A	3858	1/1	0.91	0.09	45,45,45,45	0
57	MG	1W	203	1/1	0.91	0.14	48,48,48,48	0
57	MG	1Y	201	1/1	0.91	0.18	57,57,57,57	0
57	MG	2A	3198	1/1	0.91	0.35	75,75,75,75	0
57	MG	2A	3688	1/1	0.91	0.19	55,55,55,55	0
57	MG	10	101	1/1	0.91	0.10	43,43,43,43	0
57	MG	1a	1797	1/1	0.91	0.20	70,70,70,70	0
57	MG	2A	3042	1/1	0.91	0.20	48,48,48,48	0
57	MG	2A	3044	1/1	0.91	0.16	67,67,67,67	0
57	MG	10	102	1/1	0.91	0.29	47,47,47,47	0
57	MG	2A	3435	1/1	0.91	0.21	76,76,76,76	0
57	MG	1a	1693	1/1	0.91	0.26	70,70,70,70	0
57	MG	10	104	1/1	0.91	0.11	59,59,59,59	0
57	MG	1A	3860	1/1	0.91	0.10	58,58,58,58	0
57	MG	1A	3240	1/1	0.91	0.13	50,50,50,50	0
57	MG	2A	3052	1/1	0.91	0.23	66,66,66,66	0
57	MG	1a	1809	1/1	0.91	0.12	60,60,60,60	0
57	MG	2A	3712	1/1	0.91	0.29	68,68,68,68	0
57	MG	2A	3448	1/1	0.91	0.35	61,61,61,61	0
57	MG	2A	3054	1/1	0.91	0.25	65,65,65,65	0
57	MG	11	101	1/1	0.91	0.17	44,44,44,44	0
57	MG	11	102	1/1	0.91	0.18	59,59,59,59	0
57	MG	1A	3371	1/1	0.91	0.15	64,64,64,64	0
57	MG	1a	1700	1/1	0.91	0.41	60,60,60,60	0
57	MG	1A	3532	1/1	0.91	0.14	53,53,53,53	0
57	MG	2A	3064	1/1	0.91	0.17	57,57,57,57	0
57	MG	1A	3761	1/1	0.91	0.07	29,29,29,29	0
57	MG	17	104	1/1	0.91	0.14	53,53,53,53	0
57	MG	2a	3145	1/1	0.91	0.10	52,52,52,52	0
57	MG	1A	3619	1/1	0.91	0.10	57,57,57,57	0
57	MG	2A	3731	1/1	0.91	0.14	63,63,63,63	0
57	MG	2A	3469	1/1	0.91	0.08	62,62,62,62	0
57	MG	1A	3621	1/1	0.91	0.08	24,24,24,24	0
57	MG	18	103	1/1	0.91	0.08	54,54,54,54	0
57	MG	1A	3773	1/1	0.91	0.17	45,45,45,45	0
57	MG	1A	3297	1/1	0.91	0.28	65,65,65,65	0
57	MG	2a	3159	1/1	0.91	0.09	70,70,70,70	0
57	MG	2B	206	1/1	0.91	0.22	65,65,65,65	0
57	MG	2A	3476	1/1	0.91	0.12	63,63,63,63	0
57	MG	1A	3123	1/1	0.91	0.19	54,54,54,54	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3086	1/1	0.91	0.13	51,51,51,51	0
57	MG	2A	3263	1/1	0.91	0.16	64,64,64,64	0
57	MG	1A	3076	1/1	0.91	0.16	55,55,55,55	0
57	MG	1A	3577	1/1	0.91	0.26	65,65,65,65	0
57	MG	2A	3267	1/1	0.91	0.20	54,54,54,54	0
57	MG	1a	1607	1/1	0.91	0.09	63,63,63,63	0
57	MG	2A	3271	1/1	0.91	0.29	61,61,61,61	0
57	MG	2a	3172	1/1	0.91	0.25	68,68,68,68	0
57	MG	2a	3173	1/1	0.91	0.26	65,65,65,65	0
57	MG	1a	1836	1/1	0.91	0.10	58,58,58,58	0
57	MG	1A	3889	1/1	0.91	0.12	57,57,57,57	0
57	MG	1A	3635	1/1	0.91	0.19	39,39,39,39	0
57	MG	1A	3537	1/1	0.91	0.11	56,56,56,56	0
57	MG	2A	3504	1/1	0.91	0.10	55,55,55,55	0
57	MG	1A	3001	1/1	0.91	0.09	36,36,36,36	0
57	MG	1A	3194	1/1	0.91	0.29	65,65,65,65	0
57	MG	2a	3182	1/1	0.91	0.17	59,59,59,59	0
57	MG	1A	3900	1/1	0.91	0.14	63,63,63,63	0
57	MG	2A	3109	1/1	0.91	0.27	74,74,74,74	0
57	MG	1a	1726	1/1	0.91	0.17	57,57,57,57	0
57	MG	1A	3379	1/1	0.91	0.18	53,53,53,53	0
57	MG	1A	3797	1/1	0.91	0.09	60,60,60,60	0
57	MG	2A	3297	1/1	0.91	0.42	60,60,60,60	0
57	MG	2R	203	1/1	0.91	0.14	62,62,62,62	0
57	MG	2A	3522	1/1	0.91	0.26	71,71,71,71	0
57	MG	1a	1852	1/1	0.91	0.07	61,61,61,61	0
57	MG	2V	201	1/1	0.91	0.18	69,69,69,69	0
57	MG	2V	203	1/1	0.91	0.27	72,72,72,72	0
57	MG	2X	101	1/1	0.91	0.12	52,52,52,52	0
57	MG	2Y	201	1/1	0.91	0.23	64,64,64,64	0
57	MG	1a	1729	1/1	0.91	0.15	53,53,53,53	0
60	ZN	24	501	1/1	0.91	0.17	129,129,129,129	0
57	MG	1A	3594	1/1	0.92	0.09	42,42,42,42	0
57	MG	2A	3342	1/1	0.92	0.18	42,42,42,42	0
57	MG	1A	3923	1/1	0.92	0.14	47,47,47,47	0
57	MG	2A	3346	1/1	0.92	0.16	58,58,58,58	0
57	MG	1a	1800	1/1	0.92	0.09	68,68,68,68	0
57	MG	2a	3008	1/1	0.92	0.11	68,68,68,68	0
57	MG	1a	1705	1/1	0.92	0.27	63,63,63,63	0
57	MG	1A	3654	1/1	0.92	0.10	59,59,59,59	0
57	MG	2A	3357	1/1	0.92	0.11	51,51,51,51	0
57	MG	1A	3087	1/1	0.92	0.09	56,56,56,56	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1a	1806	1/1	0.92	0.18	82,82,82,82	0
57	MG	1B	227	1/1	0.92	0.20	66,66,66,66	0
57	MG	1A	3269	1/1	0.92	0.08	45,45,45,45	0
57	MG	2A	3374	1/1	0.92	0.19	70,70,70,70	0
57	MG	2A	3375	1/1	0.92	0.13	43,43,43,43	0
57	MG	1A	3597	1/1	0.92	0.20	39,39,39,39	0
57	MG	1a	1626	1/1	0.92	0.19	57,57,57,57	0
57	MG	2a	3027	1/1	0.92	0.15	73,73,73,73	0
57	MG	2a	3029	1/1	0.92	0.23	70,70,70,70	0
57	MG	1a	1812	1/1	0.92	0.21	70,70,70,70	0
57	MG	1A	3937	1/1	0.92	0.07	26,26,26,26	0
57	MG	2A	3580	1/1	0.92	0.12	52,52,52,52	0
57	MG	2A	3043	1/1	0.92	0.14	61,61,61,61	0
57	MG	2A	3585	1/1	0.92	0.14	58,58,58,58	0
57	MG	1A	3553	1/1	0.92	0.09	41,41,41,41	0
57	MG	1A	3364	1/1	0.92	0.14	27,27,27,27	0
57	MG	2A	3595	1/1	0.92	0.09	70,70,70,70	0
57	MG	1A	3825	1/1	0.92	0.11	31,31,31,31	0
57	MG	1A	3241	1/1	0.92	0.17	38,38,38,38	0
57	MG	2A	3387	1/1	0.92	0.23	63,63,63,63	0
57	MG	1A	3948	1/1	0.92	0.12	48,48,48,48	0
57	MG	1A	3388	1/1	0.92	0.06	29,29,29,29	0
57	MG	2A	3606	1/1	0.92	0.17	38,38,38,38	0
57	MG	1A	3392	1/1	0.92	0.09	38,38,38,38	0
57	MG	2A	3610	1/1	0.92	0.11	50,50,50,50	0
57	MG	2a	3048	1/1	0.92	0.35	69,69,69,69	0
57	MG	2A	3395	1/1	0.92	0.14	61,61,61,61	0
57	MG	2A	3613	1/1	0.92	0.10	60,60,60,60	0
57	MG	1A	3058	1/1	0.92	0.15	49,49,49,49	0
57	MG	2A	3203	1/1	0.92	0.11	63,63,63,63	0
57	MG	1A	3192	1/1	0.92	0.12	48,48,48,48	0
57	MG	1A	3840	1/1	0.92	0.12	51,51,51,51	0
57	MG	2A	3401	1/1	0.92	0.07	36,36,36,36	0
57	MG	2A	3402	1/1	0.92	0.13	64,64,64,64	0
57	MG	1A	3022	1/1	0.92	0.08	49,49,49,49	0
57	MG	1A	3614	1/1	0.92	0.07	44,44,44,44	0
57	MG	1a	1833	1/1	0.92	0.09	68,68,68,68	0
57	MG	1A	3845	1/1	0.92	0.13	30,30,30,30	0
57	MG	2A	3410	1/1	0.92	0.07	32,32,32,32	0
57	MG	1A	3966	1/1	0.92	0.12	62,62,62,62	0
57	MG	2A	3412	1/1	0.92	0.12	48,48,48,48	0
57	MG	2A	3657	1/1	0.92	0.12	45,45,45,45	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3214	1/1	0.92	0.31	72,72,72,72	0
57	MG	1a	1737	1/1	0.92	0.10	67,67,67,67	0
57	MG	1A	3316	1/1	0.92	0.12	60,60,60,60	0
57	MG	1A	3847	1/1	0.92	0.15	39,39,39,39	0
57	MG	2A	3220	1/1	0.92	0.21	67,67,67,67	0
57	MG	1A	3758	1/1	0.92	0.08	33,33,33,33	0
57	MG	1A	3975	1/1	0.92	0.19	48,48,48,48	0
57	MG	1a	1843	1/1	0.92	0.08	56,56,56,56	0
57	MG	1A	3483	1/1	0.92	0.09	29,29,29,29	0
57	MG	1A	3573	1/1	0.92	0.06	23,23,23,23	0
57	MG	2A	3231	1/1	0.92	0.45	74,74,74,74	0
57	MG	1A	3080	1/1	0.92	0.19	55,55,55,55	0
57	MG	2A	3234	1/1	0.92	0.17	66,66,66,66	0
57	MG	1a	1747	1/1	0.92	0.11	66,66,66,66	0
57	MG	2A	3436	1/1	0.92	0.08	53,53,53,53	0
57	MG	2a	3096	1/1	0.92	0.20	63,63,63,63	0
57	MG	2A	3440	1/1	0.92	0.12	53,53,53,53	0
57	MG	1R	206	1/1	0.92	0.20	53,53,53,53	0
57	MG	1a	1854	1/1	0.92	0.17	67,67,67,67	0
57	MG	2A	3243	1/1	0.92	0.15	47,47,47,47	0
57	MG	2A	3696	1/1	0.92	0.13	70,70,70,70	0
57	MG	2A	3245	1/1	0.92	0.19	57,57,57,57	0
57	MG	2a	3104	1/1	0.92	0.33	61,61,61,61	0
57	MG	2A	3248	1/1	0.92	0.12	55,55,55,55	0
57	MG	1A	3375	1/1	0.92	0.08	38,38,38,38	0
57	MG	1A	3763	1/1	0.92	0.12	60,60,60,60	0
57	MG	1A	3984	1/1	0.92	0.10	16,16,16,16	0
57	MG	2A	3450	1/1	0.92	0.10	45,45,45,45	0
57	MG	1A	3630	1/1	0.92	0.17	57,57,57,57	0
57	MG	2A	3705	1/1	0.92	0.18	76,76,76,76	0
57	MG	2a	3116	1/1	0.92	0.20	71,71,71,71	0
57	MG	2A	3256	1/1	0.92	0.29	62,62,62,62	0
57	MG	2A	3708	1/1	0.92	0.18	62,62,62,62	0
57	MG	1a	1756	1/1	0.92	0.09	58,58,58,58	0
57	MG	1A	3631	1/1	0.92	0.15	49,49,49,49	0
57	MG	2A	3457	1/1	0.92	0.11	61,61,61,61	0
57	MG	1A	3345	1/1	0.92	0.14	42,42,42,42	0
57	MG	2a	3126	1/1	0.92	0.09	76,76,76,76	0
57	MG	2A	3459	1/1	0.92	0.10	43,43,43,43	0
57	MG	1a	1673	1/1	0.92	0.27	63,63,63,63	0
57	MG	1A	3349	1/1	0.92	0.17	60,60,60,60	0
57	MG	2A	3466	1/1	0.92	0.16	71,71,71,71	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3583	1/1	0.92	0.14	60,60,60,60	0
57	MG	2A	3725	1/1	0.92	0.17	56,56,56,56	0
57	MG	1a	1677	1/1	0.92	0.14	66,66,66,66	0
57	MG	2a	3137	1/1	0.92	0.08	63,63,63,63	0
57	MG	1A	3455	1/1	0.92	0.12	52,52,52,52	0
57	MG	2a	3144	1/1	0.92	0.13	67,67,67,67	0
57	MG	1A	3782	1/1	0.92	0.08	26,26,26,26	0
57	MG	2A	3275	1/1	0.92	0.37	58,58,58,58	0
57	MG	1A	3701	1/1	0.92	0.14	69,69,69,69	0
57	MG	10	105	1/1	0.92	0.14	56,56,56,56	0
57	MG	1A	3196	1/1	0.92	0.19	56,56,56,56	0
57	MG	2A	3282	1/1	0.92	0.34	58,58,58,58	0
57	MG	2A	3283	1/1	0.92	0.29	56,56,56,56	0
57	MG	1A	3044	1/1	0.92	0.34	68,68,68,68	0
57	MG	1A	3545	1/1	0.92	0.26	56,56,56,56	0
57	MG	2A	3487	1/1	0.92	0.19	52,52,52,52	0
57	MG	1a	1774	1/1	0.92	0.15	69,69,69,69	0
57	MG	1A	3789	1/1	0.92	0.08	43,43,43,43	0
57	MG	1f	202	1/1	0.92	0.14	51,51,51,51	0
57	MG	1g	201	1/1	0.92	0.31	67,67,67,67	0
57	MG	2A	3494	1/1	0.92	0.13	57,57,57,57	0
57	MG	1A	3882	1/1	0.92	0.13	59,59,59,59	0
57	MG	2A	3296	1/1	0.92	0.45	69,69,69,69	0
57	MG	1A	3791	1/1	0.92	0.22	52,52,52,52	0
57	MG	1a	1779	1/1	0.92	0.10	50,50,50,50	0
57	MG	1A	4013	1/1	0.92	0.11	61,61,61,61	0
57	MG	1A	3792	1/1	0.92	0.10	51,51,51,51	0
57	MG	2D	308	1/1	0.92	0.12	69,69,69,69	0
57	MG	2A	3507	1/1	0.92	0.12	44,44,44,44	0
57	MG	1a	1785	1/1	0.92	0.14	75,75,75,75	0
57	MG	1a	1786	1/1	0.92	0.14	68,68,68,68	0
57	MG	1A	3710	1/1	0.92	0.07	62,62,62,62	0
57	MG	1A	3796	1/1	0.92	0.11	56,56,56,56	0
57	MG	2a	3180	1/1	0.92	0.29	72,72,72,72	0
57	MG	1a	1789	1/1	0.92	0.14	78,78,78,78	0
57	MG	1A	3356	1/1	0.92	0.10	33,33,33,33	0
57	MG	1t	201	1/1	0.92	0.12	57,57,57,57	0
57	MG	1A	3460	1/1	0.92	0.09	30,30,30,30	0
57	MG	1A	3801	1/1	0.92	0.10	30,30,30,30	0
57	MG	2A	3526	1/1	0.92	0.13	45,45,45,45	0
57	MG	2A	3527	1/1	0.92	0.08	72,72,72,72	0
57	MG	2A	3320	1/1	0.92	0.10	51,51,51,51	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2t	201	1/1	0.92	0.16	46,46,46,46	0
57	MG	1y	202	1/1	0.92	0.10	84,84,84,84	0
57	MG	2U	202	1/1	0.92	0.11	84,84,84,84	0
57	MG	1x	101	1/1	0.92	0.13	47,47,47,47	0
57	MG	1x	102	1/1	0.92	0.12	32,32,32,32	0
57	MG	1A	3593	1/1	0.92	0.12	67,67,67,67	0
57	MG	1A	3719	1/1	0.92	0.09	33,33,33,33	0
57	MG	2A	3005	1/1	0.92	0.09	63,63,63,63	0
57	MG	2A	3010	1/1	0.92	0.15	65,65,65,65	0
57	MG	1A	3810	1/1	0.92	0.18	36,36,36,36	0
57	MG	1a	1710	1/1	0.93	0.34	64,64,64,64	0
57	MG	1A	3127	1/1	0.93	0.14	40,40,40,40	0
57	MG	1A	3976	1/1	0.93	0.13	59,59,59,59	0
57	MG	2R	202	1/1	0.93	0.20	62,62,62,62	0
57	MG	1A	3526	1/1	0.93	0.13	54,54,54,54	0
57	MG	2T	201	1/1	0.93	0.21	73,73,73,73	0
57	MG	2A	3515	1/1	0.93	0.14	56,56,56,56	0
57	MG	1A	3829	1/1	0.93	0.08	45,45,45,45	0
57	MG	2A	3124	1/1	0.93	0.41	47,47,47,47	0
57	MG	1a	1715	1/1	0.93	0.42	63,63,63,63	0
57	MG	2A	3126	1/1	0.93	0.19	49,49,49,49	0
57	MG	2W	202	1/1	0.93	0.20	56,56,56,56	0
57	MG	2A	3524	1/1	0.93	0.10	73,73,73,73	0
57	MG	1A	3440	1/1	0.93	0.10	46,46,46,46	0
57	MG	1A	3606	1/1	0.93	0.32	42,42,42,42	0
57	MG	15	102	1/1	0.93	0.20	51,51,51,51	0
57	MG	1A	3607	1/1	0.93	0.19	35,35,35,35	0
57	MG	1a	1721	1/1	0.93	0.33	69,69,69,69	0
57	MG	28	102	1/1	0.93	0.24	62,62,62,62	0
57	MG	2A	3314	1/1	0.93	0.07	37,37,37,37	0
57	MG	2A	3315	1/1	0.93	0.07	50,50,50,50	0
57	MG	1A	3245	1/1	0.93	0.13	54,54,54,54	0
57	MG	1A	3011	1/1	0.93	0.11	60,60,60,60	0
57	MG	17	106	1/1	0.93	0.10	58,58,58,58	0
57	MG	1a	1725	1/1	0.93	0.24	66,66,66,66	0
57	MG	1A	3185	1/1	0.93	0.30	60,60,60,60	0
57	MG	1a	1865	1/1	0.93	0.21	59,59,59,59	0
57	MG	1A	3251	1/1	0.93	0.09	69,69,69,69	0
57	MG	2A	3543	1/1	0.93	0.13	59,59,59,59	0
57	MG	1A	3988	1/1	0.93	0.11	53,53,53,53	0
57	MG	1A	3186	1/1	0.93	0.21	78,78,78,78	0
57	MG	2A	3151	1/1	0.93	0.07	43,43,43,43	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3341	1/1	0.93	0.17	59,59,59,59	0
57	MG	1A	3615	1/1	0.93	0.12	24,24,24,24	0
57	MG	1A	3253	1/1	0.93	0.12	55,55,55,55	0
57	MG	1A	3030	1/1	0.93	0.08	31,31,31,31	0
57	MG	1A	3733	1/1	0.93	0.06	45,45,45,45	0
57	MG	2a	3030	1/1	0.93	0.10	58,58,58,58	0
57	MG	2A	3351	1/1	0.93	0.10	37,37,37,37	0
57	MG	1a	1608	1/1	0.93	0.11	61,61,61,61	0
57	MG	2A	3556	1/1	0.93	0.15	34,34,34,34	0
57	MG	2A	3354	1/1	0.93	0.08	39,39,39,39	0
57	MG	1A	4000	1/1	0.93	0.24	42,42,42,42	0
57	MG	2A	3564	1/1	0.93	0.09	32,32,32,32	0
57	MG	2A	3568	1/1	0.93	0.10	75,75,75,75	0
57	MG	1A	3453	1/1	0.93	0.10	61,61,61,61	0
57	MG	2A	3358	1/1	0.93	0.12	35,35,35,35	0
57	MG	1A	3032	1/1	0.93	0.34	62,62,62,62	0
57	MG	1a	1617	1/1	0.93	0.12	67,67,67,67	0
57	MG	1A	3855	1/1	0.93	0.08	36,36,36,36	0
57	MG	2A	3368	1/1	0.93	0.09	32,32,32,32	0
57	MG	1a	1620	1/1	0.93	0.08	55,55,55,55	0
57	MG	2A	3167	1/1	0.93	0.15	42,42,42,42	0
57	MG	1A	3034	1/1	0.93	0.08	50,50,50,50	0
57	MG	1A	3628	1/1	0.93	0.08	43,43,43,43	0
57	MG	1A	3148	1/1	0.93	0.11	56,56,56,56	0
57	MG	1a	1629	1/1	0.93	0.22	72,72,72,72	0
57	MG	1A	3747	1/1	0.93	0.08	52,52,52,52	0
57	MG	1A	3750	1/1	0.93	0.07	50,50,50,50	0
57	MG	2A	3177	1/1	0.93	0.12	65,65,65,65	0
57	MG	1A	3095	1/1	0.93	0.09	41,41,41,41	0
57	MG	1A	3105	1/1	0.93	0.10	68,68,68,68	0
57	MG	1A	3547	1/1	0.93	0.27	55,55,55,55	0
57	MG	2A	3386	1/1	0.93	0.20	62,62,62,62	0
57	MG	1A	3389	1/1	0.93	0.08	40,40,40,40	0
57	MG	1A	3118	1/1	0.93	0.11	38,38,38,38	0
57	MG	1A	3395	1/1	0.93	0.08	31,31,31,31	0
57	MG	1a	1764	1/1	0.93	0.18	70,70,70,70	0
57	MG	1A	3551	1/1	0.93	0.13	51,51,51,51	0
57	MG	1A	3396	1/1	0.93	0.07	41,41,41,41	0
57	MG	2a	3067	1/1	0.93	0.24	71,71,71,71	0
57	MG	1A	3400	1/1	0.93	0.08	26,26,26,26	0
57	MG	2a	3069	1/1	0.93	0.35	70,70,70,70	0
57	MG	1B	219	1/1	0.93	0.11	66,66,66,66	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3626	1/1	0.93	0.06	54,54,54,54	0
57	MG	1A	3331	1/1	0.93	0.12	30,30,30,30	0
57	MG	2A	3631	1/1	0.93	0.09	44,44,44,44	0
57	MG	2A	3636	1/1	0.93	0.07	60,60,60,60	0
57	MG	2a	3079	1/1	0.93	0.10	59,59,59,59	0
57	MG	2A	3002	1/1	0.93	0.19	53,53,53,53	0
57	MG	1a	1649	1/1	0.93	0.14	44,44,44,44	0
57	MG	1A	3473	1/1	0.93	0.15	63,63,63,63	0
57	MG	2A	3647	1/1	0.93	0.07	66,66,66,66	0
57	MG	2A	3650	1/1	0.93	0.08	45,45,45,45	0
57	MG	1B	224	1/1	0.93	0.19	65,65,65,65	0
57	MG	1A	3403	1/1	0.93	0.09	29,29,29,29	0
57	MG	1A	3332	1/1	0.93	0.10	26,26,26,26	0
57	MG	2A	3658	1/1	0.93	0.08	70,70,70,70	0
57	MG	1a	1776	1/1	0.93	0.18	59,59,59,59	0
57	MG	2a	3094	1/1	0.93	0.33	67,67,67,67	0
57	MG	2a	3095	1/1	0.93	0.48	71,71,71,71	0
57	MG	2A	3019	1/1	0.93	0.25	48,48,48,48	0
57	MG	2A	3022	1/1	0.93	0.24	44,44,44,44	0
57	MG	1A	3769	1/1	0.93	0.07	44,44,44,44	0
57	MG	1A	3655	1/1	0.93	0.06	16,16,16,16	0
57	MG	1a	1660	1/1	0.93	0.17	64,64,64,64	0
57	MG	2A	3416	1/1	0.93	0.08	43,43,43,43	0
57	MG	2A	3417	1/1	0.93	0.08	50,50,50,50	0
57	MG	1A	3341	1/1	0.93	0.10	55,55,55,55	0
57	MG	1A	3899	1/1	0.93	0.12	30,30,30,30	0
57	MG	1A	3343	1/1	0.93	0.09	30,30,30,30	0
57	MG	1A	3570	1/1	0.93	0.08	63,63,63,63	0
57	MG	1D	314	1/1	0.93	0.09	69,69,69,69	0
57	MG	1A	3909	1/1	0.93	0.08	63,63,63,63	0
57	MG	2A	3680	1/1	0.93	0.10	76,76,76,76	0
57	MG	1A	3779	1/1	0.93	0.13	55,55,55,55	0
57	MG	2a	3115	1/1	0.93	0.18	66,66,66,66	0
57	MG	1E	306	1/1	0.93	0.09	25,25,25,25	0
57	MG	1A	3156	1/1	0.93	0.15	56,56,56,56	0
57	MG	1A	3416	1/1	0.93	0.07	27,27,27,27	0
57	MG	2a	3119	1/1	0.93	0.14	65,65,65,65	0
57	MG	1A	3205	1/1	0.93	0.29	38,38,38,38	0
57	MG	1A	3157	1/1	0.93	0.09	36,36,36,36	0
57	MG	2A	3437	1/1	0.93	0.08	48,48,48,48	0
57	MG	1A	3668	1/1	0.93	0.25	41,41,41,41	0
57	MG	1a	1676	1/1	0.93	0.18	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3787	1/1	0.93	0.17	69,69,69,69	0
57	MG	1F	315	1/1	0.93	0.12	60,60,60,60	0
57	MG	1A	3070	1/1	0.93	0.09	38,38,38,38	0
57	MG	1A	3934	1/1	0.93	0.09	72,72,72,72	0
57	MG	2A	3055	1/1	0.93	0.14	47,47,47,47	0
57	MG	2A	3706	1/1	0.93	0.14	66,66,66,66	0
57	MG	2A	3241	1/1	0.93	0.10	54,54,54,54	0
57	MG	2a	3135	1/1	0.93	0.14	62,62,62,62	0
57	MG	1A	3671	1/1	0.93	0.09	55,55,55,55	0
57	MG	2A	3244	1/1	0.93	0.10	60,60,60,60	0
57	MG	1A	3940	1/1	0.93	0.13	57,57,57,57	0
57	MG	1a	1683	1/1	0.93	0.15	59,59,59,59	0
57	MG	1a	1684	1/1	0.93	0.12	56,56,56,56	0
57	MG	1A	3420	1/1	0.93	0.07	32,32,32,32	0
57	MG	1a	1687	1/1	0.93	0.10	45,45,45,45	0
57	MG	2A	3720	1/1	0.93	0.15	77,77,77,77	0
57	MG	2A	3721	1/1	0.93	0.13	73,73,73,73	0
57	MG	1A	3497	1/1	0.93	0.07	62,62,62,62	0
57	MG	2A	3068	1/1	0.93	0.13	54,54,54,54	0
57	MG	1A	3286	1/1	0.93	0.17	50,50,50,50	0
57	MG	1A	3163	1/1	0.93	0.27	58,58,58,58	0
57	MG	1a	1817	1/1	0.93	0.07	83,83,83,83	0
57	MG	1A	3425	1/1	0.93	0.13	29,29,29,29	0
57	MG	1R	204	1/1	0.93	0.13	39,39,39,39	0
57	MG	1A	3426	1/1	0.93	0.09	25,25,25,25	0
57	MG	2a	3166	1/1	0.93	0.14	70,70,70,70	0
57	MG	1A	3505	1/1	0.93	0.08	25,25,25,25	0
57	MG	2A	3472	1/1	0.93	0.21	55,55,55,55	0
57	MG	1A	3167	1/1	0.93	0.15	33,33,33,33	0
57	MG	1A	3071	1/1	0.93	0.30	54,54,54,54	0
57	MG	2A	3273	1/1	0.93	0.27	55,55,55,55	0
57	MG	1A	3686	1/1	0.93	0.14	46,46,46,46	0
57	MG	2A	3477	1/1	0.93	0.20	64,64,64,64	0
57	MG	2A	3478	1/1	0.93	0.19	58,58,58,58	0
57	MG	1A	3054	1/1	0.93	0.13	44,44,44,44	0
57	MG	1A	3055	1/1	0.93	0.14	58,58,58,58	0
57	MG	1A	3693	1/1	0.93	0.18	56,56,56,56	0
57	MG	1W	202	1/1	0.93	0.17	52,52,52,52	0
57	MG	2A	3486	1/1	0.93	0.06	59,59,59,59	0
57	MG	1A	3303	1/1	0.93	0.17	49,49,49,49	0
57	MG	2A	3488	1/1	0.93	0.12	74,74,74,74	0
57	MG	2A	3284	1/1	0.93	0.29	67,67,67,67	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3285	1/1	0.93	0.25	53,53,53,53	0
57	MG	1A	3516	1/1	0.93	0.08	53,53,53,53	0
57	MG	2f	201	1/1	0.93	0.16	58,58,58,58	0
57	MG	2D	303	1/1	0.93	0.29	45,45,45,45	0
57	MG	2A	3287	1/1	0.93	0.49	70,70,70,70	0
57	MG	2D	305	1/1	0.93	0.14	55,55,55,55	0
57	MG	1A	3969	1/1	0.93	0.11	49,49,49,49	0
57	MG	2D	309	1/1	0.93	0.08	35,35,35,35	0
57	MG	2A	3106	1/1	0.93	0.18	62,62,62,62	0
57	MG	2x	102	1/1	0.93	0.17	52,52,52,52	0
57	MG	2A	3496	1/1	0.93	0.09	56,56,56,56	0
57	MG	2F	302	1/1	0.93	0.30	47,47,47,47	0
58	MPD	18	104	8/8	0.93	0.14	32,40,42,51	0
57	MG	2A	3107	1/1	0.93	0.14	62,62,62,62	0
57	MG	1A	3971	1/1	0.93	0.06	49,49,49,49	0
57	MG	1a	1707	1/1	0.93	0.43	64,64,64,64	0
57	MG	2A	3293	1/1	0.93	0.10	55,55,55,55	0
57	MG	1A	3307	1/1	0.93	0.23	45,45,45,45	0
57	MG	1A	3519	1/1	0.93	0.08	24,24,24,24	0
57	MG	1Q	204	1/1	0.94	0.10	39,39,39,39	0
57	MG	2A	3356	1/1	0.94	0.08	49,49,49,49	0
57	MG	1Q	205	1/1	0.94	0.14	55,55,55,55	0
57	MG	1A	3665	1/1	0.94	0.13	49,49,49,49	0
57	MG	1A	3970	1/1	0.94	0.13	50,50,50,50	0
57	MG	20	102	1/1	0.94	0.11	64,64,64,64	0
57	MG	2A	3363	1/1	0.94	0.10	43,43,43,43	0
57	MG	2A	3364	1/1	0.94	0.08	35,35,35,35	0
57	MG	1A	3666	1/1	0.94	0.07	41,41,41,41	0
57	MG	27	102	1/1	0.94	0.15	42,42,42,42	0
57	MG	1A	3844	1/1	0.94	0.08	37,37,37,37	0
57	MG	1A	3165	1/1	0.94	0.15	29,29,29,29	0
57	MG	1A	3262	1/1	0.94	0.09	55,55,55,55	0
57	MG	1A	3212	1/1	0.94	0.11	56,56,56,56	0
57	MG	2A	3554	1/1	0.94	0.09	47,47,47,47	0
57	MG	1A	3502	1/1	0.94	0.11	52,52,52,52	0
57	MG	1T	207	1/1	0.94	0.14	58,58,58,58	0
57	MG	1a	1794	1/1	0.94	0.12	59,59,59,59	0
57	MG	1U	205	1/1	0.94	0.25	38,38,38,38	0
57	MG	2a	3013	1/1	0.94	0.26	71,71,71,71	0
57	MG	1V	203	1/1	0.94	0.10	42,42,42,42	0
57	MG	2A	3569	1/1	0.94	0.10	71,71,71,71	0
57	MG	2A	3038	1/1	0.94	0.16	49,49,49,49	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1a	1685	1/1	0.94	0.17	56,56,56,56	0
57	MG	1A	3978	1/1	0.94	0.19	40,40,40,40	0
57	MG	2A	3576	1/1	0.94	0.11	58,58,58,58	0
57	MG	2a	3024	1/1	0.94	0.17	65,65,65,65	0
57	MG	1a	1799	1/1	0.94	0.07	66,66,66,66	0
57	MG	1V	207	1/1	0.94	0.08	59,59,59,59	0
57	MG	2A	3205	1/1	0.94	0.21	61,61,61,61	0
57	MG	2a	3028	1/1	0.94	0.12	57,57,57,57	0
57	MG	2A	3583	1/1	0.94	0.07	48,48,48,48	0
57	MG	1A	3552	1/1	0.94	0.08	52,52,52,52	0
57	MG	1A	3503	1/1	0.94	0.07	49,49,49,49	0
57	MG	2A	3390	1/1	0.94	0.08	53,53,53,53	0
57	MG	2A	3593	1/1	0.94	0.07	61,61,61,61	0
57	MG	1A	3300	1/1	0.94	0.14	54,54,54,54	0
57	MG	1A	3333	1/1	0.94	0.15	51,51,51,51	0
57	MG	1Z	301	1/1	0.94	0.13	62,62,62,62	0
57	MG	1A	3983	1/1	0.94	0.16	57,57,57,57	0
57	MG	1A	3339	1/1	0.94	0.09	35,35,35,35	0
57	MG	1A	3767	1/1	0.94	0.07	32,32,32,32	0
57	MG	1A	3507	1/1	0.94	0.10	51,51,51,51	0
57	MG	1A	3681	1/1	0.94	0.06	49,49,49,49	0
57	MG	1A	3616	1/1	0.94	0.10	27,27,27,27	0
57	MG	2A	3221	1/1	0.94	0.10	52,52,52,52	0
57	MG	2A	3404	1/1	0.94	0.09	50,50,50,50	0
57	MG	1A	3301	1/1	0.94	0.24	35,35,35,35	0
57	MG	1A	3046	1/1	0.94	0.26	51,51,51,51	0
57	MG	2A	3224	1/1	0.94	0.17	60,60,60,60	0
57	MG	2A	3227	1/1	0.94	0.22	51,51,51,51	0
57	MG	2a	3050	1/1	0.94	0.22	57,57,57,57	0
57	MG	1A	3422	1/1	0.94	0.14	31,31,31,31	0
57	MG	2A	3622	1/1	0.94	0.07	57,57,57,57	0
57	MG	1A	3622	1/1	0.94	0.09	52,52,52,52	0
57	MG	13	103	1/1	0.94	0.09	45,45,45,45	0
57	MG	1A	3423	1/1	0.94	0.11	38,38,38,38	0
57	MG	1A	3625	1/1	0.94	0.09	37,37,37,37	0
57	MG	15	107	1/1	0.94	0.12	71,71,71,71	0
57	MG	2A	3073	1/1	0.94	0.08	51,51,51,51	0
57	MG	2A	3637	1/1	0.94	0.10	53,53,53,53	0
57	MG	2A	3236	1/1	0.94	0.22	59,59,59,59	0
57	MG	1a	1826	1/1	0.94	0.08	65,65,65,65	0
57	MG	2A	3238	1/1	0.94	0.08	60,60,60,60	0
57	MG	2A	3646	1/1	0.94	0.08	61,61,61,61	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3227	1/1	0.94	0.18	36,36,36,36	0
57	MG	1A	3627	1/1	0.94	0.17	71,71,71,71	0
57	MG	2A	3080	1/1	0.94	0.41	50,50,50,50	0
57	MG	1A	3169	1/1	0.94	0.18	64,64,64,64	0
57	MG	2A	3656	1/1	0.94	0.07	52,52,52,52	0
57	MG	1A	3629	1/1	0.94	0.15	56,56,56,56	0
57	MG	1A	4010	1/1	0.94	0.20	51,51,51,51	0
57	MG	2A	3432	1/1	0.94	0.08	51,51,51,51	0
57	MG	2a	3075	1/1	0.94	0.13	50,50,50,50	0
57	MG	2a	3076	1/1	0.94	0.23	69,69,69,69	0
57	MG	1a	1834	1/1	0.94	0.12	60,60,60,60	0
57	MG	1A	3233	1/1	0.94	0.29	66,66,66,66	0
57	MG	1A	3790	1/1	0.94	0.10	33,33,33,33	0
57	MG	1A	3522	1/1	0.94	0.10	62,62,62,62	0
57	MG	2A	3091	1/1	0.94	0.17	53,53,53,53	0
57	MG	2a	3083	1/1	0.94	0.16	67,67,67,67	0
57	MG	1A	3049	1/1	0.94	0.19	44,44,44,44	0
57	MG	2a	3087	1/1	0.94	0.17	55,55,55,55	0
57	MG	2A	3260	1/1	0.94	0.11	32,32,32,32	0
57	MG	2A	3443	1/1	0.94	0.06	61,61,61,61	0
57	MG	1A	3470	1/1	0.94	0.07	33,33,33,33	0
57	MG	2A	3262	1/1	0.94	0.27	50,50,50,50	0
57	MG	1A	3795	1/1	0.94	0.12	59,59,59,59	0
57	MG	1B	205	1/1	0.94	0.10	50,50,50,50	0
57	MG	1a	1609	1/1	0.94	0.09	60,60,60,60	0
57	MG	2A	3103	1/1	0.94	0.18	67,67,67,67	0
57	MG	2A	3269	1/1	0.94	0.12	45,45,45,45	0
57	MG	1a	1610	1/1	0.94	0.29	60,60,60,60	0
57	MG	2A	3685	1/1	0.94	0.11	59,59,59,59	0
57	MG	2A	3686	1/1	0.94	0.06	43,43,43,43	0
57	MG	2a	3100	1/1	0.94	0.37	63,63,63,63	0
57	MG	1A	3471	1/1	0.94	0.12	55,55,55,55	0
57	MG	2A	3690	1/1	0.94	0.10	46,46,46,46	0
57	MG	2A	3453	1/1	0.94	0.07	37,37,37,37	0
57	MG	2A	3693	1/1	0.94	0.17	57,57,57,57	0
57	MG	1A	3274	1/1	0.94	0.19	39,39,39,39	0
57	MG	2A	3695	1/1	0.94	0.13	62,62,62,62	0
57	MG	2A	3455	1/1	0.94	0.11	65,65,65,65	0
57	MG	1A	3390	1/1	0.94	0.09	40,40,40,40	0
57	MG	2A	3274	1/1	0.94	0.22	51,51,51,51	0
57	MG	1A	3102	1/1	0.94	0.12	38,38,38,38	0
57	MG	1A	3911	1/1	0.94	0.10	66,66,66,66	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3277	1/1	0.94	0.10	50,50,50,50	0
57	MG	2A	3702	1/1	0.94	0.09	47,47,47,47	0
57	MG	1A	3640	1/1	0.94	0.08	65,65,65,65	0
57	MG	2A	3279	1/1	0.94	0.11	53,53,53,53	0
57	MG	2A	3280	1/1	0.94	0.11	61,61,61,61	0
57	MG	1A	3807	1/1	0.94	0.09	48,48,48,48	0
57	MG	1a	1731	1/1	0.94	0.15	48,48,48,48	0
57	MG	1a	1623	1/1	0.94	0.15	54,54,54,54	0
57	MG	2A	3709	1/1	0.94	0.09	65,65,65,65	0
57	MG	2A	3118	1/1	0.94	0.17	37,37,37,37	0
57	MG	1a	1733	1/1	0.94	0.07	56,56,56,56	0
57	MG	1A	3721	1/1	0.94	0.12	46,46,46,46	0
57	MG	2a	3129	1/1	0.94	0.22	69,69,69,69	0
57	MG	1a	1625	1/1	0.94	0.09	58,58,58,58	0
57	MG	2A	3714	1/1	0.94	0.19	67,67,67,67	0
57	MG	1A	3919	1/1	0.94	0.16	45,45,45,45	0
57	MG	1a	1628	1/1	0.94	0.41	65,65,65,65	0
57	MG	1A	3809	1/1	0.94	0.12	27,27,27,27	0
57	MG	2A	3479	1/1	0.94	0.10	46,46,46,46	0
57	MG	2a	3136	1/1	0.94	0.13	67,67,67,67	0
57	MG	1A	3073	1/1	0.94	0.18	51,51,51,51	0
57	MG	2A	3129	1/1	0.94	0.19	66,66,66,66	0
57	MG	2A	3482	1/1	0.94	0.08	71,71,71,71	0
57	MG	1A	3158	1/1	0.94	0.12	45,45,45,45	0
57	MG	2A	3484	1/1	0.94	0.09	35,35,35,35	0
57	MG	1A	3813	1/1	0.94	0.20	35,35,35,35	0
57	MG	2a	3151	1/1	0.94	0.16	54,54,54,54	0
57	MG	1a	1636	1/1	0.94	0.44	65,65,65,65	0
57	MG	1A	3930	1/1	0.94	0.09	54,54,54,54	0
57	MG	1B	229	1/1	0.94	0.19	77,77,77,77	0
57	MG	1A	3814	1/1	0.94	0.15	75,75,75,75	0
57	MG	1a	1641	1/1	0.94	0.15	69,69,69,69	0
57	MG	1a	1752	1/1	0.94	0.07	55,55,55,55	0
57	MG	2A	3140	1/1	0.94	0.12	50,50,50,50	0
57	MG	1A	3933	1/1	0.94	0.09	62,62,62,62	0
57	MG	1A	3644	1/1	0.94	0.10	59,59,59,59	0
57	MG	1A	3590	1/1	0.94	0.07	48,48,48,48	0
57	MG	1A	3938	1/1	0.94	0.11	49,49,49,49	0
57	MG	1A	3183	1/1	0.94	0.16	72,72,72,72	0
57	MG	2A	3501	1/1	0.94	0.16	60,60,60,60	0
57	MG	1A	3734	1/1	0.94	0.06	22,22,22,22	0
57	MG	2B	211	1/1	0.94	0.09	69,69,69,69	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3822	1/1	0.94	0.10	53,53,53,53	0
57	MG	1A	3945	1/1	0.94	0.08	62,62,62,62	0
57	MG	2A	3156	1/1	0.94	0.13	66,66,66,66	0
57	MG	2A	3318	1/1	0.94	0.13	38,38,38,38	0
57	MG	1A	3159	1/1	0.94	0.07	35,35,35,35	0
57	MG	1A	3248	1/1	0.94	0.07	43,43,43,43	0
57	MG	2A	3512	1/1	0.94	0.12	40,40,40,40	0
57	MG	1A	3126	1/1	0.94	0.17	43,43,43,43	0
57	MG	1A	3950	1/1	0.94	0.13	82,82,82,82	0
57	MG	2D	307	1/1	0.94	0.22	49,49,49,49	0
57	MG	1A	3488	1/1	0.94	0.07	30,30,30,30	0
57	MG	2A	3330	1/1	0.94	0.09	42,42,42,42	0
57	MG	2E	303	1/1	0.94	0.50	49,49,49,49	0
57	MG	1A	3827	1/1	0.94	0.21	58,58,58,58	0
57	MG	1A	3406	1/1	0.94	0.13	47,47,47,47	0
57	MG	2A	3523	1/1	0.94	0.19	64,64,64,64	0
57	MG	1A	3830	1/1	0.94	0.10	52,52,52,52	0
57	MG	2A	3337	1/1	0.94	0.13	46,46,46,46	0
57	MG	1G	201	1/1	0.94	0.11	65,65,65,65	0
57	MG	1A	3407	1/1	0.94	0.07	27,27,27,27	0
57	MG	2A	3528	1/1	0.94	0.10	63,63,63,63	0
57	MG	1A	3834	1/1	0.94	0.10	49,49,49,49	0
57	MG	1A	3748	1/1	0.94	0.09	54,54,54,54	0
57	MG	2A	3343	1/1	0.94	0.12	58,58,58,58	0
57	MG	1A	3598	1/1	0.94	0.12	56,56,56,56	0
57	MG	1A	3494	1/1	0.94	0.07	48,48,48,48	0
57	MG	1A	3967	1/1	0.94	0.06	62,62,62,62	0
57	MG	1A	3066	1/1	0.94	0.21	41,41,41,41	0
57	MG	2A	3537	1/1	0.94	0.11	37,37,37,37	0
57	MG	1a	1781	1/1	0.94	0.08	72,72,72,72	0
57	MG	2A	3353	1/1	0.94	0.12	47,47,47,47	0
57	MG	2A	3006	1/1	0.94	0.15	51,51,51,51	0
57	MG	1A	3539	1/1	0.95	0.07	37,37,37,37	0
57	MG	1A	3112	1/1	0.95	0.26	41,41,41,41	0
57	MG	2A	3197	1/1	0.95	0.10	53,53,53,53	0
57	MG	1Q	203	1/1	0.95	0.10	53,53,53,53	0
57	MG	2A	3559	1/1	0.95	0.09	69,69,69,69	0
57	MG	1a	1670	1/1	0.95	0.32	61,61,61,61	0
57	MG	1A	3703	1/1	0.95	0.07	29,29,29,29	0
57	MG	2A	3563	1/1	0.95	0.07	59,59,59,59	0
57	MG	1A	3275	1/1	0.95	0.25	36,36,36,36	0
57	MG	2A	3382	1/1	0.95	0.05	40,40,40,40	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	28	101	1/1	0.95	0.09	65,65,65,65	0
57	MG	1R	201	1/1	0.95	0.17	41,41,41,41	0
57	MG	1A	3276	1/1	0.95	0.25	43,43,43,43	0
57	MG	2a	3001	1/1	0.95	0.07	46,46,46,46	0
57	MG	2a	3003	1/1	0.95	0.08	56,56,56,56	0
57	MG	1A	3278	1/1	0.95	0.13	47,47,47,47	0
57	MG	1A	3620	1/1	0.95	0.15	54,54,54,54	0
57	MG	2A	3574	1/1	0.95	0.08	54,54,54,54	0
57	MG	1A	3117	1/1	0.95	0.12	37,37,37,37	0
57	MG	1A	3712	1/1	0.95	0.19	38,38,38,38	0
57	MG	1A	3346	1/1	0.95	0.11	50,50,50,50	0
57	MG	2A	3581	1/1	0.95	0.09	48,48,48,48	0
57	MG	1A	3478	1/1	0.95	0.06	28,28,28,28	0
57	MG	2a	3014	1/1	0.95	0.23	57,57,57,57	0
57	MG	1T	205	1/1	0.95	0.08	57,57,57,57	0
57	MG	1A	3716	1/1	0.95	0.06	33,33,33,33	0
57	MG	1A	3624	1/1	0.95	0.09	30,30,30,30	0
57	MG	1A	3347	1/1	0.95	0.10	52,52,52,52	0
57	MG	2A	3591	1/1	0.95	0.08	50,50,50,50	0
57	MG	1A	3061	1/1	0.95	0.07	35,35,35,35	0
57	MG	1A	3724	1/1	0.95	0.10	57,57,57,57	0
57	MG	2A	3597	1/1	0.95	0.07	37,37,37,37	0
57	MG	1V	205	1/1	0.95	0.07	55,55,55,55	0
57	MG	2A	3056	1/1	0.95	0.15	55,55,55,55	0
57	MG	1A	3078	1/1	0.95	0.23	44,44,44,44	0
57	MG	2A	3226	1/1	0.95	0.14	54,54,54,54	0
57	MG	1a	1690	1/1	0.95	0.17	57,57,57,57	0
57	MG	2A	3605	1/1	0.95	0.07	44,44,44,44	0
57	MG	1A	3972	1/1	0.95	0.06	20,20,20,20	0
57	MG	1A	3214	1/1	0.95	0.13	49,49,49,49	0
57	MG	1A	3354	1/1	0.95	0.05	34,34,34,34	0
57	MG	2a	3035	1/1	0.95	0.12	41,41,41,41	0
57	MG	2A	3611	1/1	0.95	0.06	26,26,26,26	0
57	MG	1A	3732	1/1	0.95	0.10	30,30,30,30	0
57	MG	1A	3839	1/1	0.95	0.18	40,40,40,40	0
57	MG	1A	3218	1/1	0.95	0.06	39,39,39,39	0
57	MG	2A	3615	1/1	0.95	0.12	59,59,59,59	0
57	MG	2A	3616	1/1	0.95	0.19	64,64,64,64	0
57	MG	1A	3487	1/1	0.95	0.15	29,29,29,29	0
57	MG	2A	3072	1/1	0.95	0.21	51,51,51,51	0
57	MG	1A	3012	1/1	0.95	0.10	41,41,41,41	0
57	MG	1A	3633	1/1	0.95	0.17	33,33,33,33	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3226	1/1	0.95	0.24	38,38,38,38	0
57	MG	2A	3240	1/1	0.95	0.30	44,44,44,44	0
57	MG	2A	3076	1/1	0.95	0.15	58,58,58,58	0
57	MG	2A	3242	1/1	0.95	0.09	60,60,60,60	0
57	MG	1A	3559	1/1	0.95	0.07	63,63,63,63	0
57	MG	2A	3630	1/1	0.95	0.08	57,57,57,57	0
57	MG	1A	3745	1/1	0.95	0.08	26,26,26,26	0
57	MG	2A	3427	1/1	0.95	0.09	48,48,48,48	0
57	MG	1A	3288	1/1	0.95	0.09	33,33,33,33	0
57	MG	1A	3563	1/1	0.95	0.08	52,52,52,52	0
57	MG	2A	3639	1/1	0.95	0.14	33,33,33,33	0
57	MG	11	105	1/1	0.95	0.09	43,43,43,43	0
57	MG	1A	3851	1/1	0.95	0.07	63,63,63,63	0
57	MG	2A	3251	1/1	0.95	0.31	65,65,65,65	0
57	MG	2a	3061	1/1	0.95	0.32	74,74,74,74	0
57	MG	2A	3434	1/1	0.95	0.10	38,38,38,38	0
57	MG	2A	3649	1/1	0.95	0.07	42,42,42,42	0
57	MG	1A	3638	1/1	0.95	0.12	62,62,62,62	0
57	MG	1A	3564	1/1	0.95	0.18	45,45,45,45	0
57	MG	2A	3654	1/1	0.95	0.12	69,69,69,69	0
57	MG	15	104	1/1	0.95	0.21	38,38,38,38	0
57	MG	2A	3439	1/1	0.95	0.14	56,56,56,56	0
57	MG	1A	3122	1/1	0.95	0.09	46,46,46,46	0
57	MG	1A	3495	1/1	0.95	0.17	60,60,60,60	0
57	MG	1A	3042	1/1	0.95	0.09	18,18,18,18	0
57	MG	1A	3368	1/1	0.95	0.09	33,33,33,33	0
57	MG	1A	3084	1/1	0.95	0.09	51,51,51,51	0
57	MG	1A	3861	1/1	0.95	0.06	25,25,25,25	0
57	MG	1A	3085	1/1	0.95	0.10	37,37,37,37	0
57	MG	2A	3104	1/1	0.95	0.14	47,47,47,47	0
57	MG	2a	3078	1/1	0.95	0.41	65,65,65,65	0
57	MG	2A	3268	1/1	0.95	0.31	61,61,61,61	0
57	MG	1A	3238	1/1	0.95	0.23	44,44,44,44	0
57	MG	1A	3653	1/1	0.95	0.06	23,23,23,23	0
57	MG	2A	3673	1/1	0.95	0.16	54,54,54,54	0
57	MG	1a	1845	1/1	0.95	0.16	77,77,77,77	0
57	MG	1A	3064	1/1	0.95	0.27	54,54,54,54	0
57	MG	1A	3374	1/1	0.95	0.12	38,38,38,38	0
57	MG	1a	1849	1/1	0.95	0.12	58,58,58,58	0
57	MG	1A	3581	1/1	0.95	0.11	47,47,47,47	0
57	MG	1a	1851	1/1	0.95	0.09	44,44,44,44	0
57	MG	2A	3681	1/1	0.95	0.08	70,70,70,70	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1a	1606	1/1	0.95	0.24	57,57,57,57	0
57	MG	2A	3115	1/1	0.95	0.23	50,50,50,50	0
57	MG	2A	3684	1/1	0.95	0.07	40,40,40,40	0
57	MG	1A	3872	1/1	0.95	0.09	49,49,49,49	0
57	MG	2A	3460	1/1	0.95	0.12	62,62,62,62	0
57	MG	1A	3008	1/1	0.95	0.06	38,38,38,38	0
57	MG	1A	3658	1/1	0.95	0.06	47,47,47,47	0
57	MG	1A	3439	1/1	0.95	0.18	55,55,55,55	0
57	MG	2A	3122	1/1	0.95	0.25	44,44,44,44	0
57	MG	1A	3768	1/1	0.95	0.08	50,50,50,50	0
57	MG	1A	3242	1/1	0.95	0.22	47,47,47,47	0
57	MG	2A	3470	1/1	0.95	0.06	56,56,56,56	0
57	MG	1A	3772	1/1	0.95	0.05	30,30,30,30	0
57	MG	1A	3131	1/1	0.95	0.29	32,32,32,32	0
57	MG	1B	208	1/1	0.95	0.35	65,65,65,65	0
57	MG	1a	1864	1/1	0.95	0.29	57,57,57,57	0
57	MG	1a	1619	1/1	0.95	0.13	55,55,55,55	0
57	MG	1A	3587	1/1	0.95	0.06	23,23,23,23	0
57	MG	1A	3313	1/1	0.95	0.15	60,60,60,60	0
57	MG	1A	3777	1/1	0.95	0.08	39,39,39,39	0
57	MG	2A	3135	1/1	0.95	0.09	37,37,37,37	0
57	MG	1A	3893	1/1	0.95	0.07	35,35,35,35	0
57	MG	1A	3896	1/1	0.95	0.06	32,32,32,32	0
57	MG	1A	3445	1/1	0.95	0.09	51,51,51,51	0
57	MG	1A	3513	1/1	0.95	0.06	37,37,37,37	0
57	MG	1A	3056	1/1	0.95	0.14	28,28,28,28	0
57	MG	1B	222	1/1	0.95	0.08	35,35,35,35	0
57	MG	2A	3144	1/1	0.95	0.20	45,45,45,45	0
57	MG	2A	3145	1/1	0.95	0.13	47,47,47,47	0
57	MG	2A	3305	1/1	0.95	0.11	52,52,52,52	0
57	MG	2A	3716	1/1	0.95	0.11	66,66,66,66	0
57	MG	1a	1631	1/1	0.95	0.12	43,43,43,43	0
57	MG	2A	3490	1/1	0.95	0.11	69,69,69,69	0
57	MG	1A	3902	1/1	0.95	0.05	46,46,46,46	0
57	MG	1A	3903	1/1	0.95	0.21	41,41,41,41	0
57	MG	1a	1751	1/1	0.95	0.15	42,42,42,42	0
57	MG	2A	3152	1/1	0.95	0.18	61,61,61,61	0
57	MG	1A	3138	1/1	0.95	0.12	43,43,43,43	0
57	MG	1a	1637	1/1	0.95	0.09	44,44,44,44	0
57	MG	2A	3497	1/1	0.95	0.08	35,35,35,35	0
57	MG	2A	3498	1/1	0.95	0.07	67,67,67,67	0
57	MG	2a	3139	1/1	0.95	0.07	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2a	3140	1/1	0.95	0.07	77,77,77,77	0
57	MG	2a	3141	1/1	0.95	0.09	72,72,72,72	0
57	MG	1A	3009	1/1	0.95	0.07	32,32,32,32	0
57	MG	1A	3142	1/1	0.95	0.10	53,53,53,53	0
57	MG	1A	3097	1/1	0.95	0.20	42,42,42,42	0
57	MG	1A	3145	1/1	0.95	0.12	51,51,51,51	0
57	MG	1A	3254	1/1	0.95	0.20	36,36,36,36	0
57	MG	2A	3324	1/1	0.95	0.19	55,55,55,55	0
57	MG	2A	3506	1/1	0.95	0.15	28,28,28,28	0
57	MG	1A	3147	1/1	0.95	0.09	40,40,40,40	0
57	MG	1A	3099	1/1	0.95	0.12	47,47,47,47	0
57	MG	2A	3327	1/1	0.95	0.10	61,61,61,61	0
57	MG	1A	3150	1/1	0.95	0.08	41,41,41,41	0
57	MG	2A	3332	1/1	0.95	0.06	36,36,36,36	0
57	MG	2A	3333	1/1	0.95	0.14	58,58,58,58	0
57	MG	1A	3925	1/1	0.95	0.10	54,54,54,54	0
57	MG	1A	3072	1/1	0.95	0.06	39,39,39,39	0
57	MG	1A	3927	1/1	0.95	0.06	58,58,58,58	0
57	MG	2A	3519	1/1	0.95	0.10	62,62,62,62	0
57	MG	2A	3168	1/1	0.95	0.15	51,51,51,51	0
57	MG	1A	3103	1/1	0.95	0.12	45,45,45,45	0
57	MG	1A	3794	1/1	0.95	0.17	40,40,40,40	0
57	MG	1A	3398	1/1	0.95	0.10	27,27,27,27	0
57	MG	2B	219	1/1	0.95	0.08	69,69,69,69	0
57	MG	2D	301	1/1	0.95	0.22	39,39,39,39	0
57	MG	1A	3048	1/1	0.95	0.23	51,51,51,51	0
57	MG	1a	1771	1/1	0.95	0.14	60,60,60,60	0
57	MG	2A	3174	1/1	0.95	0.11	38,38,38,38	0
57	MG	1A	3687	1/1	0.95	0.08	59,59,59,59	0
57	MG	2A	3347	1/1	0.95	0.16	59,59,59,59	0
57	MG	1a	1656	1/1	0.95	0.28	58,58,58,58	0
57	MG	2A	3350	1/1	0.95	0.08	52,52,52,52	0
57	MG	1A	3936	1/1	0.95	0.08	34,34,34,34	0
57	MG	1A	3689	1/1	0.95	0.20	50,50,50,50	0
57	MG	2A	3180	1/1	0.95	0.08	55,55,55,55	0
57	MG	2A	3181	1/1	0.95	0.07	46,46,46,46	0
57	MG	1A	3608	1/1	0.95	0.10	46,46,46,46	0
57	MG	2A	3538	1/1	0.95	0.12	57,57,57,57	0
57	MG	1A	3270	1/1	0.95	0.22	43,43,43,43	0
57	MG	1A	3942	1/1	0.95	0.07	60,60,60,60	0
57	MG	2A	3014	1/1	0.95	0.10	54,54,54,54	0
57	MG	2P	201	1/1	0.95	0.09	61,61,61,61	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3361	1/1	0.95	0.10	37,37,37,37	0
57	MG	1A	3536	1/1	0.95	0.26	31,31,31,31	0
57	MG	1a	1780	1/1	0.95	0.17	64,64,64,64	0
57	MG	1A	3109	1/1	0.95	0.18	44,44,44,44	0
57	MG	2A	3365	1/1	0.95	0.08	37,37,37,37	0
57	MG	1A	3612	1/1	0.95	0.11	33,33,33,33	0
57	MG	2A	3023	1/1	0.95	0.20	61,61,61,61	0
57	MG	2T	203	1/1	0.95	0.13	68,68,68,68	0
57	MG	2T	204	1/1	0.95	0.09	59,59,59,59	0
57	MG	1a	1783	1/1	0.95	0.10	79,79,79,79	0
57	MG	1A	3538	1/1	0.95	0.05	26,26,26,26	0
57	MG	2A	3027	1/1	0.95	0.05	36,36,36,36	0
57	MG	2V	202	1/1	0.95	0.25	49,49,49,49	0
57	MG	1A	3182	1/1	0.96	0.09	51,51,51,51	0
57	MG	1A	3143	1/1	0.96	0.09	41,41,41,41	0
57	MG	2A	3344	1/1	0.96	0.07	56,56,56,56	0
57	MG	1A	3243	1/1	0.96	0.17	44,44,44,44	0
57	MG	1A	3784	1/1	0.96	0.09	35,35,35,35	0
57	MG	1A	3662	1/1	0.96	0.07	54,54,54,54	0
57	MG	1A	3569	1/1	0.96	0.13	52,52,52,52	0
57	MG	2A	3349	1/1	0.96	0.06	51,51,51,51	0
57	MG	1A	3928	1/1	0.96	0.06	22,22,22,22	0
57	MG	1A	3477	1/1	0.96	0.06	28,28,28,28	0
57	MG	2A	3160	1/1	0.96	0.23	45,45,45,45	0
57	MG	1A	3108	1/1	0.96	0.15	44,44,44,44	0
57	MG	2W	203	1/1	0.96	0.07	54,54,54,54	0
57	MG	1A	3931	1/1	0.96	0.08	48,48,48,48	0
57	MG	1A	3081	1/1	0.96	0.20	46,46,46,46	0
57	MG	1A	3391	1/1	0.96	0.09	37,37,37,37	0
57	MG	2A	3557	1/1	0.96	0.06	54,54,54,54	0
57	MG	1A	3481	1/1	0.96	0.11	47,47,47,47	0
57	MG	1A	3935	1/1	0.96	0.07	29,29,29,29	0
57	MG	1T	203	1/1	0.96	0.11	58,58,58,58	0
57	MG	1T	204	1/1	0.96	0.07	63,63,63,63	0
57	MG	1d	302	1/1	0.96	0.12	71,71,71,71	0
57	MG	1A	3110	1/1	0.96	0.11	56,56,56,56	0
57	MG	1d	304	1/1	0.96	0.07	70,70,70,70	0
57	MG	1A	3082	1/1	0.96	0.33	48,48,48,48	0
57	MG	2a	3002	1/1	0.96	0.26	60,60,60,60	0
57	MG	1A	3578	1/1	0.96	0.07	52,52,52,52	0
57	MG	1U	203	1/1	0.96	0.18	37,37,37,37	0
57	MG	1U	204	1/1	0.96	0.23	29,29,29,29	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3176	1/1	0.96	0.11	65,65,65,65	0
57	MG	2a	3007	1/1	0.96	0.24	56,56,56,56	0
57	MG	1A	3484	1/1	0.96	0.10	34,34,34,34	0
57	MG	1V	201	1/1	0.96	0.23	36,36,36,36	0
57	MG	1A	3941	1/1	0.96	0.07	58,58,58,58	0
57	MG	1A	3250	1/1	0.96	0.10	48,48,48,48	0
57	MG	2a	3012	1/1	0.96	0.08	50,50,50,50	0
57	MG	1A	3188	1/1	0.96	0.15	37,37,37,37	0
57	MG	1V	206	1/1	0.96	0.11	59,59,59,59	0
57	MG	1A	3798	1/1	0.96	0.22	46,46,46,46	0
57	MG	2A	3589	1/1	0.96	0.13	49,49,49,49	0
57	MG	2a	3017	1/1	0.96	0.13	49,49,49,49	0
57	MG	1A	3189	1/1	0.96	0.15	42,42,42,42	0
57	MG	1a	1720	1/1	0.96	0.13	50,50,50,50	0
57	MG	1A	3800	1/1	0.96	0.08	48,48,48,48	0
57	MG	1A	3325	1/1	0.96	0.06	37,37,37,37	0
57	MG	1A	3802	1/1	0.96	0.09	59,59,59,59	0
57	MG	1A	3586	1/1	0.96	0.20	56,56,56,56	0
57	MG	1A	3489	1/1	0.96	0.07	38,38,38,38	0
57	MG	1A	3402	1/1	0.96	0.07	27,27,27,27	0
57	MG	10	103	1/1	0.96	0.07	47,47,47,47	0
57	MG	1A	3955	1/1	0.96	0.05	28,28,28,28	0
57	MG	2A	3604	1/1	0.96	0.08	46,46,46,46	0
57	MG	1A	3190	1/1	0.96	0.19	61,61,61,61	0
57	MG	10	106	1/1	0.96	0.07	44,44,44,44	0
57	MG	2A	3607	1/1	0.96	0.06	65,65,65,65	0
57	MG	1A	3047	1/1	0.96	0.21	41,41,41,41	0
57	MG	1A	3257	1/1	0.96	0.08	54,54,54,54	0
57	MG	1A	3496	1/1	0.96	0.09	27,27,27,27	0
57	MG	2A	3400	1/1	0.96	0.13	65,65,65,65	0
57	MG	1A	3329	1/1	0.96	0.13	30,30,30,30	0
57	MG	2A	3202	1/1	0.96	0.29	65,65,65,65	0
57	MG	1A	3069	1/1	0.96	0.06	35,35,35,35	0
57	MG	1A	3965	1/1	0.96	0.15	56,56,56,56	0
57	MG	1A	3409	1/1	0.96	0.05	22,22,22,22	0
57	MG	1A	3818	1/1	0.96	0.05	23,23,23,23	0
57	MG	2A	3407	1/1	0.96	0.16	61,61,61,61	0
57	MG	15	101	1/1	0.96	0.09	41,41,41,41	0
57	MG	2A	3621	1/1	0.96	0.14	60,60,60,60	0
57	MG	2A	3208	1/1	0.96	0.17	59,59,59,59	0
57	MG	1A	3119	1/1	0.96	0.09	41,41,41,41	0
57	MG	1A	3820	1/1	0.96	0.11	53,53,53,53	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3697	1/1	0.96	0.14	57,57,57,57	0
57	MG	1A	3020	1/1	0.96	0.23	38,38,38,38	0
57	MG	2A	3213	1/1	0.96	0.09	53,53,53,53	0
57	MG	2A	3018	1/1	0.96	0.39	44,44,44,44	0
57	MG	2A	3634	1/1	0.96	0.07	49,49,49,49	0
57	MG	2A	3635	1/1	0.96	0.10	60,60,60,60	0
57	MG	2A	3215	1/1	0.96	0.11	52,52,52,52	0
57	MG	17	103	1/1	0.96	0.16	38,38,38,38	0
57	MG	1a	1749	1/1	0.96	0.06	54,54,54,54	0
57	MG	1A	3414	1/1	0.96	0.06	37,37,37,37	0
57	MG	2A	3421	1/1	0.96	0.10	41,41,41,41	0
57	MG	2A	3641	1/1	0.96	0.07	41,41,41,41	0
57	MG	1A	3599	1/1	0.96	0.09	62,62,62,62	0
57	MG	1A	3004	1/1	0.96	0.09	41,41,41,41	0
57	MG	2A	3424	1/1	0.96	0.09	37,37,37,37	0
57	MG	2A	3425	1/1	0.96	0.10	37,37,37,37	0
57	MG	18	101	1/1	0.96	0.07	42,42,42,42	0
57	MG	2A	3030	1/1	0.96	0.16	60,60,60,60	0
57	MG	2A	3652	1/1	0.96	0.10	62,62,62,62	0
57	MG	2A	3428	1/1	0.96	0.05	33,33,33,33	0
57	MG	1A	3704	1/1	0.96	0.10	67,67,67,67	0
57	MG	2a	3072	1/1	0.96	0.09	64,64,64,64	0
57	MG	1A	3601	1/1	0.96	0.21	35,35,35,35	0
57	MG	1A	3334	1/1	0.96	0.08	30,30,30,30	0
57	MG	1a	1757	1/1	0.96	0.21	55,55,55,55	0
57	MG	1A	3335	1/1	0.96	0.07	26,26,26,26	0
57	MG	1A	3831	1/1	0.96	0.18	54,54,54,54	0
57	MG	2A	3661	1/1	0.96	0.05	44,44,44,44	0
57	MG	1A	3336	1/1	0.96	0.04	31,31,31,31	0
57	MG	1A	3088	1/1	0.96	0.15	35,35,35,35	0
57	MG	2A	3664	1/1	0.96	0.09	56,56,56,56	0
57	MG	2A	3666	1/1	0.96	0.09	32,32,32,32	0
57	MG	2A	3233	1/1	0.96	0.14	65,65,65,65	0
57	MG	2a	3085	1/1	0.96	0.24	51,51,51,51	0
57	MG	1A	3199	1/1	0.96	0.11	41,41,41,41	0
57	MG	1A	3342	1/1	0.96	0.09	35,35,35,35	0
57	MG	1A	3051	1/1	0.96	0.15	32,32,32,32	0
57	MG	1A	3344	1/1	0.96	0.13	47,47,47,47	0
57	MG	2A	3672	1/1	0.96	0.06	62,62,62,62	0
57	MG	1A	3204	1/1	0.96	0.14	39,39,39,39	0
57	MG	1A	3720	1/1	0.96	0.12	60,60,60,60	0
57	MG	1A	3517	1/1	0.96	0.05	56,56,56,56	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3990	1/1	0.96	0.04	38,38,38,38	0
57	MG	1a	1616	1/1	0.96	0.14	65,65,65,65	0
57	MG	1A	3843	1/1	0.96	0.06	67,67,67,67	0
57	MG	1A	3053	1/1	0.96	0.07	41,41,41,41	0
57	MG	1A	3427	1/1	0.96	0.09	34,34,34,34	0
57	MG	2A	3246	1/1	0.96	0.25	46,46,46,46	0
57	MG	2A	3247	1/1	0.96	0.06	59,59,59,59	0
57	MG	1A	3074	1/1	0.96	0.22	37,37,37,37	0
57	MG	1A	3162	1/1	0.96	0.12	45,45,45,45	0
57	MG	1A	3351	1/1	0.96	0.07	31,31,31,31	0
57	MG	1A	3096	1/1	0.96	0.12	47,47,47,47	0
57	MG	2A	3689	1/1	0.96	0.17	63,63,63,63	0
57	MG	2A	3059	1/1	0.96	0.14	44,44,44,44	0
57	MG	1A	3075	1/1	0.96	0.13	39,39,39,39	0
57	MG	2A	3692	1/1	0.96	0.09	39,39,39,39	0
57	MG	1A	3166	1/1	0.96	0.33	42,42,42,42	0
57	MG	1A	4004	1/1	0.96	0.10	56,56,56,56	0
57	MG	2a	3112	1/1	0.96	0.07	50,50,50,50	0
57	MG	1A	4005	1/1	0.96	0.09	53,53,53,53	0
57	MG	2A	3462	1/1	0.96	0.12	39,39,39,39	0
57	MG	1A	4008	1/1	0.96	0.15	68,68,68,68	0
57	MG	1A	3735	1/1	0.96	0.06	46,46,46,46	0
57	MG	1A	3854	1/1	0.96	0.10	31,31,31,31	0
57	MG	2A	3070	1/1	0.96	0.26	48,48,48,48	0
57	MG	1a	1633	1/1	0.96	0.10	40,40,40,40	0
57	MG	1a	1634	1/1	0.96	0.10	49,49,49,49	0
57	MG	1A	3023	1/1	0.96	0.19	34,34,34,34	0
57	MG	1A	3357	1/1	0.96	0.11	26,26,26,26	0
57	MG	2a	3125	1/1	0.96	0.12	58,58,58,58	0
57	MG	1A	3219	1/1	0.96	0.14	38,38,38,38	0
57	MG	1B	201	1/1	0.96	0.16	61,61,61,61	0
57	MG	1A	3441	1/1	0.96	0.08	53,53,53,53	0
57	MG	2A	3081	1/1	0.96	0.09	43,43,43,43	0
57	MG	1A	3221	1/1	0.96	0.10	38,38,38,38	0
57	MG	1A	3101	1/1	0.96	0.12	39,39,39,39	0
57	MG	1A	3862	1/1	0.96	0.05	56,56,56,56	0
57	MG	1A	3224	1/1	0.96	0.16	40,40,40,40	0
57	MG	2A	3087	1/1	0.96	0.12	52,52,52,52	0
57	MG	1A	3135	1/1	0.96	0.23	47,47,47,47	0
57	MG	2A	3715	1/1	0.96	0.19	63,63,63,63	0
57	MG	1A	3289	1/1	0.96	0.07	42,42,42,42	0
57	MG	2a	3138	1/1	0.96	0.08	75,75,75,75	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3867	1/1	0.96	0.08	29,29,29,29	0
57	MG	1A	3541	1/1	0.96	0.08	36,36,36,36	0
57	MG	1A	3039	1/1	0.96	0.17	41,41,41,41	0
57	MG	1B	214	1/1	0.96	0.07	57,57,57,57	0
57	MG	1A	3370	1/1	0.96	0.07	33,33,33,33	0
57	MG	1A	3294	1/1	0.96	0.18	41,41,41,41	0
57	MG	2a	3147	1/1	0.96	0.11	77,77,77,77	0
57	MG	1A	3295	1/1	0.96	0.23	58,58,58,58	0
57	MG	2a	3149	1/1	0.96	0.08	75,75,75,75	0
57	MG	1A	3230	1/1	0.96	0.15	37,37,37,37	0
57	MG	1A	3175	1/1	0.96	0.21	41,41,41,41	0
57	MG	1A	3177	1/1	0.96	0.07	52,52,52,52	0
57	MG	1A	3299	1/1	0.96	0.34	44,44,44,44	0
57	MG	1a	1659	1/1	0.96	0.23	50,50,50,50	0
57	MG	1a	1815	1/1	0.96	0.07	66,66,66,66	0
57	MG	1A	3079	1/1	0.96	0.13	37,37,37,37	0
57	MG	1A	3887	1/1	0.96	0.07	63,63,63,63	0
57	MG	2a	3160	1/1	0.96	0.06	71,71,71,71	0
57	MG	1A	3888	1/1	0.96	0.05	57,57,57,57	0
57	MG	1A	3235	1/1	0.96	0.24	40,40,40,40	0
57	MG	1A	3065	1/1	0.96	0.13	43,43,43,43	0
57	MG	2A	3114	1/1	0.96	0.12	49,49,49,49	0
57	MG	1A	3462	1/1	0.96	0.12	57,57,57,57	0
57	MG	1D	302	1/1	0.96	0.25	34,34,34,34	0
57	MG	2A	3117	1/1	0.96	0.10	54,54,54,54	0
57	MG	1D	305	1/1	0.96	0.24	44,44,44,44	0
57	MG	1D	307	1/1	0.96	0.05	39,39,39,39	0
57	MG	1A	3766	1/1	0.96	0.05	38,38,38,38	0
57	MG	1A	3239	1/1	0.96	0.19	33,33,33,33	0
57	MG	1a	1828	1/1	0.96	0.08	51,51,51,51	0
57	MG	2A	3513	1/1	0.96	0.07	35,35,35,35	0
57	MG	1A	3895	1/1	0.96	0.10	31,31,31,31	0
57	MG	1a	1672	1/1	0.96	0.18	69,69,69,69	0
57	MG	1A	3381	1/1	0.96	0.10	48,48,48,48	0
57	MG	1A	3897	1/1	0.96	0.09	26,26,26,26	0
57	MG	1A	3649	1/1	0.96	0.10	53,53,53,53	0
57	MG	1A	3771	1/1	0.96	0.07	42,42,42,42	0
57	MG	1A	3308	1/1	0.96	0.21	33,33,33,33	0
57	MG	1A	3180	1/1	0.96	0.15	33,33,33,33	0
57	MG	1A	3774	1/1	0.96	0.11	42,42,42,42	0
57	MG	1a	1839	1/1	0.96	0.06	44,44,44,44	0
57	MG	1A	3904	1/1	0.96	0.20	43,43,43,43	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2D	310	1/1	0.96	0.14	66,66,66,66	0
57	MG	2E	302	1/1	0.96	0.08	60,60,60,60	0
57	MG	1A	3905	1/1	0.96	0.05	55,55,55,55	0
57	MG	2A	3329	1/1	0.96	0.07	33,33,33,33	0
57	MG	1A	3384	1/1	0.96	0.08	36,36,36,36	0
57	MG	2A	3331	1/1	0.96	0.10	32,32,32,32	0
57	MG	1A	3908	1/1	0.96	0.11	46,46,46,46	0
57	MG	2F	304	1/1	0.96	0.15	53,53,53,53	0
57	MG	1A	3385	1/1	0.96	0.08	32,32,32,32	0
57	MG	1A	3561	1/1	0.96	0.06	53,53,53,53	0
57	MG	2A	3534	1/1	0.96	0.07	57,57,57,57	0
57	MG	1A	3912	1/1	0.96	0.07	56,56,56,56	0
57	MG	1A	3562	1/1	0.96	0.21	62,62,62,62	0
57	MG	1A	3472	1/1	0.96	0.10	42,42,42,42	0
57	MG	1H	202	1/1	0.96	0.06	53,53,53,53	0
57	MG	2A	3149	1/1	0.96	0.18	51,51,51,51	0
57	MG	1A	3780	1/1	0.96	0.06	30,30,30,30	0
57	MG	1A	3989	1/1	0.97	0.11	54,54,54,54	0
57	MG	2A	3020	1/1	0.97	0.11	40,40,40,40	0
57	MG	1A	3220	1/1	0.97	0.16	36,36,36,36	0
57	MG	1A	3521	1/1	0.97	0.11	23,23,23,23	0
57	MG	1A	3304	1/1	0.97	0.22	56,56,56,56	0
57	MG	2A	3025	1/1	0.97	0.08	46,46,46,46	0
57	MG	1A	3994	1/1	0.97	0.11	37,37,37,37	0
57	MG	2A	3323	1/1	0.97	0.11	43,43,43,43	0
57	MG	1A	3995	1/1	0.97	0.05	61,61,61,61	0
57	MG	2A	3029	1/1	0.97	0.15	31,31,31,31	0
57	MG	1A	3258	1/1	0.97	0.07	53,53,53,53	0
57	MG	1A	3725	1/1	0.97	0.08	30,30,30,30	0
57	MG	1A	3726	1/1	0.97	0.05	35,35,35,35	0
57	MG	1A	3259	1/1	0.97	0.10	38,38,38,38	0
57	MG	1A	3647	1/1	0.97	0.24	49,49,49,49	0
57	MG	1A	3311	1/1	0.97	0.12	35,35,35,35	0
57	MG	1A	3650	1/1	0.97	0.28	48,48,48,48	0
57	MG	2A	3674	1/1	0.97	0.06	51,51,51,51	0
57	MG	1A	3906	1/1	0.97	0.04	54,54,54,54	0
57	MG	1A	3410	1/1	0.97	0.04	38,38,38,38	0
57	MG	2A	3040	1/1	0.97	0.06	23,23,23,23	0
57	MG	1A	3360	1/1	0.97	0.06	45,45,45,45	0
57	MG	2A	3338	1/1	0.97	0.06	30,30,30,30	0
57	MG	1A	3146	1/1	0.97	0.23	37,37,37,37	0
57	MG	1A	3910	1/1	0.97	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
57	MG	1A	3736	1/1	0.97	0.10	50,50,50,50	0
57	MG	1A	3737	1/1	0.97	0.15	44,44,44,44	0
57	MG	1A	3222	1/1	0.97	0.11	37,37,37,37	0
57	MG	1A	3739	1/1	0.97	0.07	61,61,61,61	0
57	MG	2A	3048	1/1	0.97	0.15	48,48,48,48	0
57	MG	2A	3687	1/1	0.97	0.05	67,67,67,67	0
57	MG	1A	3916	1/1	0.97	0.05	31,31,31,31	0
57	MG	2A	3195	1/1	0.97	0.09	51,51,51,51	0
57	MG	1A	3917	1/1	0.97	0.04	50,50,50,50	0
57	MG	1A	3415	1/1	0.97	0.06	48,48,48,48	0
57	MG	1A	3314	1/1	0.97	0.05	30,30,30,30	0
57	MG	13	101	1/1	0.97	0.07	35,35,35,35	0
57	MG	1A	3921	1/1	0.97	0.09	62,62,62,62	0
57	MG	1A	3264	1/1	0.97	0.19	45,45,45,45	0
57	MG	1A	3744	1/1	0.97	0.13	42,42,42,42	0
57	MG	1A	3013	1/1	0.97	0.06	19,19,19,19	0
57	MG	15	103	1/1	0.97	0.21	33,33,33,33	0
57	MG	1B	211	1/1	0.97	0.09	39,39,39,39	0
57	MG	1A	3125	1/1	0.97	0.05	31,31,31,31	0
57	MG	2A	3520	1/1	0.97	0.06	44,44,44,44	0
57	MG	2A	3359	1/1	0.97	0.12	34,34,34,34	0
57	MG	15	106	1/1	0.97	0.04	52,52,52,52	0
57	MG	1A	3225	1/1	0.97	0.22	36,36,36,36	0
57	MG	17	102	1/1	0.97	0.11	35,35,35,35	0
57	MG	2A	3067	1/1	0.97	0.06	22,22,22,22	0
57	MG	1A	3041	1/1	0.97	0.09	29,29,29,29	0
57	MG	1A	3173	1/1	0.97	0.12	37,37,37,37	0
57	MG	1a	1832	1/1	0.97	0.15	70,70,70,70	0
57	MG	1A	3229	1/1	0.97	0.12	40,40,40,40	0
57	MG	2A	3370	1/1	0.97	0.06	46,46,46,46	0
57	MG	2A	3371	1/1	0.97	0.20	64,64,64,64	0
57	MG	1A	3031	1/1	0.97	0.07	42,42,42,42	0
57	MG	1A	3231	1/1	0.97	0.12	30,30,30,30	0
57	MG	2A	3218	1/1	0.97	0.10	53,53,53,53	0
57	MG	2a	3084	1/1	0.97	0.33	59,59,59,59	0
57	MG	1A	3754	1/1	0.97	0.07	36,36,36,36	0
57	MG	2A	3717	1/1	0.97	0.16	48,48,48,48	0
57	MG	1A	3324	1/1	0.97	0.09	30,30,30,30	0
57	MG	2A	3077	1/1	0.97	0.09	52,52,52,52	0
57	MG	19	101	1/1	0.97	0.07	62,62,62,62	0
57	MG	1A	3021	1/1	0.97	0.08	36,36,36,36	0
57	MG	1A	3670	1/1	0.97	0.11	59,59,59,59	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3605	1/1	0.97	0.26	35,35,35,35	0
57	MG	2A	3083	1/1	0.97	0.05	43,43,43,43	0
57	MG	1A	3277	1/1	0.97	0.18	49,49,49,49	0
57	MG	1A	3939	1/1	0.97	0.06	46,46,46,46	0
57	MG	1A	3111	1/1	0.97	0.14	36,36,36,36	0
57	MG	1A	3057	1/1	0.97	0.09	47,47,47,47	0
57	MG	1A	3033	1/1	0.97	0.10	46,46,46,46	0
57	MG	1a	1847	1/1	0.97	0.08	55,55,55,55	0
57	MG	1D	303	1/1	0.97	0.08	41,41,41,41	0
57	MG	1A	3491	1/1	0.97	0.05	32,32,32,32	0
57	MG	1A	3203	1/1	0.97	0.14	44,44,44,44	0
57	MG	1A	3848	1/1	0.97	0.04	41,41,41,41	0
57	MG	2A	3094	1/1	0.97	0.09	53,53,53,53	0
57	MG	2A	3095	1/1	0.97	0.12	46,46,46,46	0
57	MG	2A	3096	1/1	0.97	0.15	52,52,52,52	0
57	MG	1a	1613	1/1	0.97	0.09	29,29,29,29	0
57	MG	2A	3558	1/1	0.97	0.06	73,73,73,73	0
57	MG	1a	1853	1/1	0.97	0.08	52,52,52,52	0
57	MG	2A	3560	1/1	0.97	0.07	65,65,65,65	0
57	MG	1A	3007	1/1	0.97	0.07	33,33,33,33	0
57	MG	2a	3113	1/1	0.97	0.07	62,62,62,62	0
57	MG	2A	3102	1/1	0.97	0.05	33,33,33,33	0
57	MG	1a	1730	1/1	0.97	0.05	32,32,32,32	0
57	MG	1A	3181	1/1	0.97	0.17	37,37,37,37	0
57	MG	2A	3566	1/1	0.97	0.05	23,23,23,23	0
57	MG	1A	3438	1/1	0.97	0.07	15,15,15,15	0
57	MG	1A	3100	1/1	0.97	0.20	36,36,36,36	0
57	MG	1D	316	1/1	0.97	0.07	57,57,57,57	0
57	MG	1D	317	1/1	0.97	0.05	45,45,45,45	0
57	MG	2D	302	1/1	0.97	0.14	50,50,50,50	0
57	MG	2a	3123	1/1	0.97	0.12	57,57,57,57	0
57	MG	1E	301	1/1	0.97	0.07	36,36,36,36	0
57	MG	1E	302	1/1	0.97	0.15	37,37,37,37	0
57	MG	2A	3575	1/1	0.97	0.07	31,31,31,31	0
57	MG	1A	3770	1/1	0.97	0.11	28,28,28,28	0
57	MG	2A	3254	1/1	0.97	0.12	42,42,42,42	0
57	MG	2A	3255	1/1	0.97	0.10	40,40,40,40	0
57	MG	1E	305	1/1	0.97	0.12	50,50,50,50	0
57	MG	2E	301	1/1	0.97	0.18	42,42,42,42	0
57	MG	2A	3258	1/1	0.97	0.14	42,42,42,42	0
57	MG	1A	3015	1/1	0.97	0.10	32,32,32,32	0
57	MG	1A	3954	1/1	0.97	0.09	45,45,45,45	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2E	305	1/1	0.97	0.05	54,54,54,54	0
57	MG	2A	3586	1/1	0.97	0.05	57,57,57,57	0
57	MG	1A	3210	1/1	0.97	0.12	33,33,33,33	0
57	MG	1F	301	1/1	0.97	0.10	34,34,34,34	0
57	MG	1F	305	1/1	0.97	0.16	36,36,36,36	0
57	MG	2F	305	1/1	0.97	0.28	50,50,50,50	0
57	MG	1A	3244	1/1	0.97	0.10	44,44,44,44	0
57	MG	2A	3265	1/1	0.97	0.03	23,23,23,23	0
57	MG	2A	3594	1/1	0.97	0.06	59,59,59,59	0
57	MG	2A	3119	1/1	0.97	0.09	36,36,36,36	0
57	MG	2a	3146	1/1	0.97	0.14	52,52,52,52	0
57	MG	1A	3337	1/1	0.97	0.10	15,15,15,15	0
57	MG	1A	3161	1/1	0.97	0.17	29,29,29,29	0
57	MG	2Q	201	1/1	0.97	0.08	52,52,52,52	0
57	MG	1A	3960	1/1	0.97	0.07	30,30,30,30	0
57	MG	2A	3123	1/1	0.97	0.18	43,43,43,43	0
57	MG	1A	3859	1/1	0.97	0.08	40,40,40,40	0
57	MG	1A	3692	1/1	0.97	0.23	41,41,41,41	0
57	MG	1F	317	1/1	0.97	0.06	51,51,51,51	0
57	MG	1A	3293	1/1	0.97	0.17	39,39,39,39	0
57	MG	1A	3016	1/1	0.97	0.23	34,34,34,34	0
57	MG	1G	203	1/1	0.97	0.12	44,44,44,44	0
57	MG	1A	3565	1/1	0.97	0.08	22,22,22,22	0
57	MG	1A	3566	1/1	0.97	0.07	48,48,48,48	0
57	MG	2A	3438	1/1	0.97	0.10	46,46,46,46	0
57	MG	1A	3213	1/1	0.97	0.15	34,34,34,34	0
57	MG	1N	202	1/1	0.97	0.17	45,45,45,45	0
57	MG	1A	3394	1/1	0.97	0.10	27,27,27,27	0
57	MG	1A	3029	1/1	0.97	0.07	41,41,41,41	0
57	MG	1A	3702	1/1	0.97	0.06	48,48,48,48	0
57	MG	1m	201	1/1	0.97	0.07	67,67,67,67	0
57	MG	1A	3509	1/1	0.97	0.07	21,21,21,21	0
57	MG	1a	1650	1/1	0.97	0.14	49,49,49,49	0
57	MG	1Q	202	1/1	0.97	0.07	34,34,34,34	0
57	MG	1A	3215	1/1	0.97	0.16	38,38,38,38	0
57	MG	1A	3397	1/1	0.97	0.06	42,42,42,42	0
57	MG	1a	1654	1/1	0.97	0.07	41,41,41,41	0
57	MG	27	101	1/1	0.97	0.18	46,46,46,46	0
57	MG	1A	3707	1/1	0.97	0.04	38,38,38,38	0
57	MG	1A	3216	1/1	0.97	0.07	52,52,52,52	0
57	MG	2A	3627	1/1	0.97	0.08	37,37,37,37	0
57	MG	1R	202	1/1	0.97	0.11	39,39,39,39	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3294	1/1	0.97	0.07	40,40,40,40	0
57	MG	1A	3574	1/1	0.97	0.11	36,36,36,36	0
57	MG	2A	3633	1/1	0.97	0.09	56,56,56,56	0
57	MG	1A	3877	1/1	0.97	0.12	30,30,30,30	0
57	MG	1A	3878	1/1	0.97	0.07	50,50,50,50	0
57	MG	1A	3217	1/1	0.97	0.13	36,36,36,36	0
57	MG	1A	3515	1/1	0.97	0.10	22,22,22,22	0
57	MG	1A	3348	1/1	0.97	0.09	26,26,26,26	0
57	MG	2k	201	1/1	0.97	0.10	57,57,57,57	0
57	MG	1A	3164	1/1	0.97	0.08	47,47,47,47	0
57	MG	2A	3302	1/1	0.97	0.25	46,46,46,46	0
57	MG	2A	3008	1/1	0.97	0.12	41,41,41,41	0
57	MG	2A	3009	1/1	0.97	0.06	47,47,47,47	0
57	MG	1A	3714	1/1	0.97	0.08	29,29,29,29	0
57	MG	2A	3306	1/1	0.97	0.10	65,65,65,65	0
57	MG	1A	3579	1/1	0.97	0.08	40,40,40,40	0
57	MG	1A	3580	1/1	0.97	0.10	60,60,60,60	0
57	MG	2A	3309	1/1	0.97	0.06	39,39,39,39	0
57	MG	1A	3718	1/1	0.97	0.06	72,72,72,72	0
57	MG	2A	3653	1/1	0.97	0.07	43,43,43,43	0
57	MG	2a	3020	1/1	0.97	0.07	49,49,49,49	0
57	MG	1A	3104	1/1	0.97	0.18	41,41,41,41	0
60	ZN	2Y	202	1/1	0.97	0.06	93,93,93,93	0
57	MG	1U	201	1/1	0.97	0.15	38,38,38,38	0
60	ZN	2n	501	1/1	0.97	0.05	95,95,95,95	0
57	MG	2A	3079	1/1	0.98	0.11	44,44,44,44	0
57	MG	1A	3176	1/1	0.98	0.22	38,38,38,38	0
57	MG	1A	3019	1/1	0.98	0.19	29,29,29,29	0
57	MG	1A	3836	1/1	0.98	0.05	46,46,46,46	0
57	MG	1A	3260	1/1	0.98	0.26	34,34,34,34	0
57	MG	1A	3261	1/1	0.98	0.31	35,35,35,35	0
57	MG	2A	3225	1/1	0.98	0.13	41,41,41,41	0
57	MG	1A	3090	1/1	0.98	0.05	45,45,45,45	0
57	MG	1A	3113	1/1	0.98	0.08	43,43,43,43	0
57	MG	1A	3749	1/1	0.98	0.08	44,44,44,44	0
57	MG	1A	3114	1/1	0.98	0.13	38,38,38,38	0
57	MG	1D	304	1/1	0.98	0.05	40,40,40,40	0
57	MG	1A	3664	1/1	0.98	0.08	46,46,46,46	0
57	MG	1D	306	1/1	0.98	0.07	22,22,22,22	0
57	MG	1A	3265	1/1	0.98	0.07	42,42,42,42	0
57	MG	1D	308	1/1	0.98	0.19	38,38,38,38	0
57	MG	1A	3116	1/1	0.98	0.09	23,23,23,23	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1D	310	1/1	0.98	0.16	35,35,35,35	0
57	MG	1A	3025	1/1	0.98	0.16	40,40,40,40	0
57	MG	1D	312	1/1	0.98	0.10	54,54,54,54	0
57	MG	1A	3060	1/1	0.98	0.04	34,34,34,34	0
57	MG	1A	3949	1/1	0.98	0.09	45,45,45,45	0
57	MG	2A	3100	1/1	0.98	0.11	71,71,71,71	0
57	MG	1A	3524	1/1	0.98	0.06	31,31,31,31	0
57	MG	1a	1734	1/1	0.98	0.08	45,45,45,45	0
57	MG	1A	3951	1/1	0.98	0.17	43,43,43,43	0
57	MG	1a	1614	1/1	0.98	0.07	54,54,54,54	0
57	MG	1A	3149	1/1	0.98	0.21	44,44,44,44	0
57	MG	1A	3451	1/1	0.98	0.05	26,26,26,26	0
57	MG	1A	3527	1/1	0.98	0.05	30,30,30,30	0
57	MG	1E	303	1/1	0.98	0.11	35,35,35,35	0
57	MG	1A	3003	1/1	0.98	0.04	23,23,23,23	0
57	MG	1A	3271	1/1	0.98	0.06	39,39,39,39	0
57	MG	1A	3957	1/1	0.98	0.09	54,54,54,54	0
57	MG	1A	3050	1/1	0.98	0.14	31,31,31,31	0
57	MG	1A	3273	1/1	0.98	0.06	32,32,32,32	0
57	MG	2A	3555	1/1	0.98	0.09	31,31,31,31	0
57	MG	1A	3678	1/1	0.98	0.11	29,29,29,29	0
57	MG	1F	302	1/1	0.98	0.11	37,37,37,37	0
57	MG	2A	3408	1/1	0.98	0.04	68,68,68,68	0
57	MG	1a	1627	1/1	0.98	0.06	49,49,49,49	0
57	MG	1A	3152	1/1	0.98	0.17	34,34,34,34	0
57	MG	1F	306	1/1	0.98	0.17	39,39,39,39	0
57	MG	1F	307	1/1	0.98	0.13	37,37,37,37	0
57	MG	1F	308	1/1	0.98	0.16	35,35,35,35	0
57	MG	1F	309	1/1	0.98	0.11	34,34,34,34	0
57	MG	2A	3565	1/1	0.98	0.05	63,63,63,63	0
57	MG	1F	310	1/1	0.98	0.08	30,30,30,30	0
57	MG	1A	3077	1/1	0.98	0.22	40,40,40,40	0
57	MG	1A	3682	1/1	0.98	0.04	43,43,43,43	0
57	MG	2A	3570	1/1	0.98	0.07	56,56,56,56	0
57	MG	1A	3228	1/1	0.98	0.05	42,42,42,42	0
57	MG	1A	3098	1/1	0.98	0.25	35,35,35,35	0
57	MG	1A	3393	1/1	0.98	0.09	26,26,26,26	0
57	MG	1A	3005	1/1	0.98	0.08	23,23,23,23	0
57	MG	1A	3052	1/1	0.98	0.26	42,42,42,42	0
57	MG	2A	3131	1/1	0.98	0.10	30,30,30,30	0
57	MG	1A	3688	1/1	0.98	0.04	53,53,53,53	0
57	MG	1A	3017	1/1	0.98	0.21	34,34,34,34	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1o	103	1/1	0.98	0.04	61,61,61,61	0
57	MG	1A	3540	1/1	0.98	0.05	33,33,33,33	0
57	MG	1A	3869	1/1	0.98	0.07	41,41,41,41	0
57	MG	2B	217	1/1	0.98	0.07	68,68,68,68	0
57	MG	2A	3584	1/1	0.98	0.04	58,58,58,58	0
57	MG	1A	3043	1/1	0.98	0.17	17,17,17,17	0
57	MG	1A	3466	1/1	0.98	0.05	47,47,47,47	0
57	MG	1A	3067	1/1	0.98	0.29	33,33,33,33	0
57	MG	1A	3694	1/1	0.98	0.14	50,50,50,50	0
57	MG	2A	3141	1/1	0.98	0.07	37,37,37,37	0
57	MG	2A	3142	1/1	0.98	0.18	49,49,49,49	0
57	MG	2D	306	1/1	0.98	0.04	37,37,37,37	0
57	MG	2A	3592	1/1	0.98	0.07	34,34,34,34	0
57	MG	1A	3399	1/1	0.98	0.10	26,26,26,26	0
57	MG	1P	201	1/1	0.98	0.28	37,37,37,37	0
57	MG	1A	3340	1/1	0.98	0.08	26,26,26,26	0
57	MG	1A	3195	1/1	0.98	0.18	39,39,39,39	0
57	MG	2A	3147	1/1	0.98	0.06	40,40,40,40	0
57	MG	1Q	201	1/1	0.98	0.07	36,36,36,36	0
57	MG	1A	3236	1/1	0.98	0.18	34,34,34,34	0
57	MG	2A	3601	1/1	0.98	0.05	42,42,42,42	0
57	MG	1A	3237	1/1	0.98	0.06	35,35,35,35	0
57	MG	1A	3404	1/1	0.98	0.04	24,24,24,24	0
57	MG	1A	3128	1/1	0.98	0.24	34,34,34,34	0
57	MG	1A	3129	1/1	0.98	0.26	39,39,39,39	0
57	MG	2A	3015	1/1	0.98	0.22	40,40,40,40	0
57	MG	1A	3068	1/1	0.98	0.15	33,33,33,33	0
57	MG	1A	3291	1/1	0.98	0.15	44,44,44,44	0
57	MG	1A	3706	1/1	0.98	0.10	39,39,39,39	0
57	MG	2A	3158	1/1	0.98	0.15	46,46,46,46	0
57	MG	1A	3292	1/1	0.98	0.05	49,49,49,49	0
57	MG	1A	3036	1/1	0.98	0.04	32,32,32,32	0
57	MG	2A	3021	1/1	0.98	0.04	35,35,35,35	0
57	MG	1A	3991	1/1	0.98	0.04	57,57,57,57	0
57	MG	1A	3201	1/1	0.98	0.19	41,41,41,41	0
57	MG	1A	3557	1/1	0.98	0.10	42,42,42,42	0
57	MG	1A	3412	1/1	0.98	0.04	27,27,27,27	0
57	MG	1A	3202	1/1	0.98	0.14	41,41,41,41	0
57	MG	1A	3106	1/1	0.98	0.20	33,33,33,33	0
57	MG	2A	3028	1/1	0.98	0.05	44,44,44,44	0
57	MG	1A	3134	1/1	0.98	0.06	49,49,49,49	0
57	MG	1A	3998	1/1	0.98	0.05	54,54,54,54	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3355	1/1	0.98	0.06	25,25,25,25	0
57	MG	2A	3625	1/1	0.98	0.03	44,44,44,44	0
57	MG	2A	3464	1/1	0.98	0.04	53,53,53,53	0
57	MG	2A	3465	1/1	0.98	0.04	54,54,54,54	0
57	MG	1U	202	1/1	0.98	0.24	40,40,40,40	0
57	MG	2W	201	1/1	0.98	0.31	48,48,48,48	0
57	MG	1A	3045	1/1	0.98	0.04	23,23,23,23	0
57	MG	1A	3717	1/1	0.98	0.05	52,52,52,52	0
57	MG	2a	3152	1/1	0.98	0.06	59,59,59,59	0
57	MG	2A	3316	1/1	0.98	0.12	23,23,23,23	0
57	MG	1A	3168	1/1	0.98	0.14	28,28,28,28	0
57	MG	2A	3319	1/1	0.98	0.05	39,39,39,39	0
57	MG	2A	3036	1/1	0.98	0.14	50,50,50,50	0
57	MG	1A	3804	1/1	0.98	0.04	38,38,38,38	0
57	MG	1V	202	1/1	0.98	0.30	35,35,35,35	0
57	MG	1a	1804	1/1	0.98	0.07	60,60,60,60	0
57	MG	1A	3805	1/1	0.98	0.05	47,47,47,47	0
57	MG	1A	3806	1/1	0.98	0.10	50,50,50,50	0
57	MG	2A	3642	1/1	0.98	0.04	22,22,22,22	0
57	MG	1a	1807	1/1	0.98	0.08	62,62,62,62	0
57	MG	1A	4006	1/1	0.98	0.07	35,35,35,35	0
57	MG	1A	4007	1/1	0.98	0.04	48,48,48,48	0
57	MG	1A	3358	1/1	0.98	0.10	32,32,32,32	0
57	MG	1A	3359	1/1	0.98	0.04	22,22,22,22	0
57	MG	1A	3136	1/1	0.98	0.05	37,37,37,37	0
57	MG	1A	3209	1/1	0.98	0.04	22,22,22,22	0
57	MG	2A	3049	1/1	0.98	0.04	40,40,40,40	0
57	MG	1A	3723	1/1	0.98	0.07	26,26,26,26	0
57	MG	1A	3812	1/1	0.98	0.20	39,39,39,39	0
57	MG	1A	3362	1/1	0.98	0.05	14,14,14,14	0
57	MG	1A	3170	1/1	0.98	0.14	32,32,32,32	0
57	MG	1A	3498	1/1	0.98	0.06	46,46,46,46	0
57	MG	1A	3727	1/1	0.98	0.07	55,55,55,55	0
57	MG	1A	3086	1/1	0.98	0.10	41,41,41,41	0
57	MG	1A	3920	1/1	0.98	0.11	47,47,47,47	0
57	MG	1A	3729	1/1	0.98	0.13	36,36,36,36	0
57	MG	1A	3365	1/1	0.98	0.06	17,17,17,17	0
57	MG	2A	3200	1/1	0.98	0.11	48,48,48,48	0
57	MG	2A	3665	1/1	0.98	0.04	51,51,51,51	0
57	MG	1B	209	1/1	0.98	0.06	41,41,41,41	0
57	MG	1A	3306	1/1	0.98	0.13	38,38,38,38	0
57	MG	1I	103	1/1	0.98	0.06	52,52,52,52	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2a	3022	1/1	0.98	0.10	58,58,58,58	0
57	MG	2A	3063	1/1	0.98	0.07	48,48,48,48	0
57	MG	1A	3429	1/1	0.98	0.07	44,44,44,44	0
57	MG	1A	3648	1/1	0.98	0.06	54,54,54,54	0
57	MG	1A	3139	1/1	0.98	0.06	33,33,33,33	0
57	MG	1A	3140	1/1	0.98	0.05	34,34,34,34	0
57	MG	1B	215	1/1	0.98	0.12	49,49,49,49	0
57	MG	1A	3651	1/1	0.98	0.07	30,30,30,30	0
57	MG	2A	3071	1/1	0.98	0.06	50,50,50,50	0
57	MG	2A	3509	1/1	0.98	0.07	54,54,54,54	0
57	MG	1A	3432	1/1	0.98	0.04	30,30,30,30	0
57	MG	1A	3310	1/1	0.98	0.07	14,14,14,14	0
57	MG	1A	3255	1/1	0.98	0.25	37,37,37,37	0
57	MG	2A	3360	1/1	0.98	0.10	31,31,31,31	0
57	MG	1A	3256	1/1	0.98	0.06	27,27,27,27	0
60	ZN	14	501	1/1	0.98	0.07	105,105,105,105	0
60	ZN	15	108	1/1	0.98	0.03	49,49,49,49	0
57	MG	1A	3037	1/1	0.98	0.07	48,48,48,48	0
57	MG	1B	223	1/1	0.98	0.12	45,45,45,45	0
60	ZN	26	501	1/1	0.98	0.04	62,62,62,62	0
60	ZN	29	102	1/1	0.98	0.04	71,71,71,71	0
57	MG	17	101	1/1	0.98	0.05	45,45,45,45	0
61	SF4	1d	306	8/8	0.98	0.06	63,67,73,74	0
61	SF4	2d	501	8/8	0.98	0.05	68,78,79,82	0
57	MG	1A	3922	1/1	0.99	0.04	62,62,62,62	0
57	MG	1A	3137	1/1	0.99	0.03	23,23,23,23	0
57	MG	1N	204	1/1	0.99	0.03	51,51,51,51	0
57	MG	2A	3011	1/1	0.99	0.06	33,33,33,33	0
57	MG	1A	3024	1/1	0.99	0.03	17,17,17,17	0
57	MG	2A	3388	1/1	0.99	0.08	41,41,41,41	0
57	MG	2A	3643	1/1	0.99	0.04	49,49,49,49	0
57	MG	2A	3013	1/1	0.99	0.05	22,22,22,22	0
57	MG	2A	3645	1/1	0.99	0.05	34,34,34,34	0
57	MG	1A	3743	1/1	0.99	0.03	25,25,25,25	0
57	MG	1A	3492	1/1	0.99	0.06	21,21,21,21	0
57	MG	2A	3648	1/1	0.99	0.04	60,60,60,60	0
57	MG	2A	3392	1/1	0.99	0.04	31,31,31,31	0
57	MG	2A	3393	1/1	0.99	0.06	38,38,38,38	0
57	MG	1A	3828	1/1	0.99	0.04	41,41,41,41	0
57	MG	1A	3207	1/1	0.99	0.38	32,32,32,32	0
57	MG	1A	3107	1/1	0.99	0.18	30,30,30,30	0
57	MG	1A	3338	1/1	0.99	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
57	MG	1A	3091	1/1	0.99	0.12	39,39,39,39	0
57	MG	1A	3092	1/1	0.99	0.05	14,14,14,14	0
57	MG	1A	3879	1/1	0.99	0.04	34,34,34,34	0
57	MG	1A	3880	1/1	0.99	0.04	45,45,45,45	0
57	MG	1A	3881	1/1	0.99	0.08	45,45,45,45	0
57	MG	2A	3567	1/1	0.99	0.06	56,56,56,56	0
57	MG	1A	3154	1/1	0.99	0.16	37,37,37,37	0
57	MG	1A	3883	1/1	0.99	0.04	51,51,51,51	0
57	MG	1A	3884	1/1	0.99	0.06	31,31,31,31	0
57	MG	2A	3101	1/1	0.99	0.04	51,51,51,51	0
57	MG	1a	1856	1/1	0.99	0.05	61,61,61,61	0
57	MG	1A	3885	1/1	0.99	0.05	42,42,42,42	0
57	MG	2a	3143	1/1	0.99	0.09	53,53,53,53	0
57	MG	2A	3328	1/1	0.99	0.04	37,37,37,37	0
57	MG	1A	3018	1/1	0.99	0.14	29,29,29,29	0
57	MG	1A	3469	1/1	0.99	0.07	26,26,26,26	0
57	MG	1A	3026	1/1	0.99	0.17	34,34,34,34	0
57	MG	2A	3578	1/1	0.99	0.03	52,52,52,52	0
57	MG	2A	3579	1/1	0.99	0.02	46,46,46,46	0
57	MG	2E	306	1/1	0.99	0.10	33,33,33,33	0
57	MG	2a	3054	1/1	0.99	0.08	49,49,49,49	0
57	MG	1A	3675	1/1	0.99	0.04	62,62,62,62	0
57	MG	1A	3198	1/1	0.99	0.18	41,41,41,41	0
57	MG	2A	3257	1/1	0.99	0.11	23,23,23,23	0
57	MG	1A	3132	1/1	0.99	0.21	32,32,32,32	0
57	MG	1A	3443	1/1	0.99	0.03	24,24,24,24	0
57	MG	1A	3679	1/1	0.99	0.04	38,38,38,38	0
57	MG	2a	3158	1/1	0.99	0.03	55,55,55,55	0
57	MG	1A	3894	1/1	0.99	0.03	65,65,65,65	0
57	MG	2A	3587	1/1	0.99	0.03	49,49,49,49	0
57	MG	1A	3642	1/1	0.99	0.07	37,37,37,37	0
57	MG	1A	3006	1/1	0.99	0.03	23,23,23,23	0
57	MG	1A	3475	1/1	0.99	0.08	44,44,44,44	0
57	MG	1A	3302	1/1	0.99	0.15	30,30,30,30	0
57	MG	1a	1803	1/1	0.99	0.03	69,69,69,69	0
57	MG	1A	3249	1/1	0.99	0.12	29,29,29,29	0
57	MG	1A	3284	1/1	0.99	0.11	23,23,23,23	0
57	MG	1A	3901	1/1	0.99	0.04	15,15,15,15	0
57	MG	2A	3596	1/1	0.99	0.03	51,51,51,51	0
57	MG	1a	1742	1/1	0.99	0.04	37,37,37,37	0
57	MG	1A	3765	1/1	0.99	0.05	43,43,43,43	0
57	MG	1a	1744	1/1	0.99	0.05	42,42,42,42	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3510	1/1	0.99	0.07	21,21,21,21	0
57	MG	1a	1622	1/1	0.99	0.05	49,49,49,49	0
57	MG	1A	3350	1/1	0.99	0.04	52,52,52,52	0
57	MG	1F	303	1/1	0.99	0.07	32,32,32,32	0
57	MG	1a	1814	1/1	0.99	0.07	73,73,73,73	0
57	MG	2A	3518	1/1	0.99	0.06	30,30,30,30	0
57	MG	1F	304	1/1	0.99	0.16	33,33,33,33	0
57	MG	1A	3305	1/1	0.99	0.08	22,22,22,22	0
57	MG	1A	3450	1/1	0.99	0.07	26,26,26,26	0
57	MG	2A	3609	1/1	0.99	0.04	44,44,44,44	0
57	MG	1A	3961	1/1	0.99	0.04	37,37,37,37	0
57	MG	1A	3285	1/1	0.99	0.07	17,17,17,17	0
57	MG	1A	3617	1/1	0.99	0.09	32,32,32,32	0
57	MG	1A	3172	1/1	0.99	0.15	33,33,33,33	0
57	MG	1A	3028	1/1	0.99	0.22	30,30,30,30	0
57	MG	1a	1823	1/1	0.99	0.04	74,74,74,74	0
57	MG	1A	3309	1/1	0.99	0.11	34,34,34,34	0
57	MG	1A	3428	1/1	0.99	0.03	14,14,14,14	0
57	MG	2A	3066	1/1	0.99	0.04	44,44,44,44	0
57	MG	1A	3817	1/1	0.99	0.07	29,29,29,29	0
57	MG	1a	1760	1/1	0.99	0.03	55,55,55,55	0
57	MG	2A	3369	1/1	0.99	0.04	33,33,33,33	0
57	MG	1A	3914	1/1	0.99	0.03	49,49,49,49	0
57	MG	1A	3089	1/1	0.99	0.21	33,33,33,33	0
57	MG	2A	3372	1/1	0.99	0.07	42,42,42,42	0
57	MG	1A	3520	1/1	0.99	0.04	33,33,33,33	0
57	MG	1A	3864	1/1	0.99	0.04	34,34,34,34	0
57	MG	1A	3457	1/1	0.99	0.08	52,52,52,52	0
57	MG	2A	3628	1/1	0.99	0.03	38,38,38,38	0
60	ZN	1Y	202	1/1	0.99	0.04	60,60,60,60	0
57	MG	1B	216	1/1	0.99	0.06	44,44,44,44	0
57	MG	1A	3699	1/1	0.99	0.02	40,40,40,40	0
60	ZN	19	102	1/1	0.99	0.05	44,44,44,44	0
60	ZN	1n	104	1/1	0.99	0.03	68,68,68,68	0
57	MG	2a	3106	1/1	0.99	0.06	67,67,67,67	0
57	MG	1A	3115	1/1	0.99	0.21	36,36,36,36	0
60	ZN	25	103	1/1	0.99	0.04	57,57,57,57	0
57	MG	2A	3632	1/1	0.99	0.06	48,48,48,48	0
57	MG	2A	3004	1/1	0.99	0.02	33,33,33,33	0
57	MG	1A	3523	1/1	0.99	0.05	44,44,44,44	0
57	MG	1N	201	1/1	0.99	0.05	41,41,41,41	0
57	MG	2A	3007	1/1	0.99	0.04	42,42,42,42	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3317	1/1	1.00	0.05	59,59,59,59	0
60	ZN	16	501	1/1	1.00	0.05	44,44,44,44	0
57	MG	1A	3468	1/1	1.00	0.05	52,52,52,52	0
57	MG	2A	3503	1/1	1.00	0.04	37,37,37,37	0
57	MG	1B	226	1/1	1.00	0.01	47,47,47,47	0
57	MG	1A	3833	1/1	1.00	0.04	22,22,22,22	0

6.5 Other polymers [i](#)

There are no such residues in this entry.