



wwPDB X-ray Structure Validation Summary Report ⓘ

Mar 18, 2026 – 12:36 PM UTC

PDB ID : 7RQC / pdb_00007rqc
Title : Crystal structure of the Thermus thermophilus 70S ribosome in complex with protein Y, A-site aminoacyl-tRNA analog ACC-PMN, and P-site MFI-tripeptidyl-tRNA analog ACCA-IFM at 2.50Å resolution
Authors : Syroegin, E.A.; Flemmich, L.; Klepacki, D.; Vazquez-Laslop, N.; Micura, R.; Polikanov, Y.S.
Deposited on : 2021-08-06
Resolution : 2.50 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	:	FAILED
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	:	9.0.010 (Gargrove)
Density-Fitness	:	1.0.12
Ideal geometry (proteins)	:	Engh & Huber (2001)
Ideal geometry (DNA, RNA)	:	Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP)	:	2.49

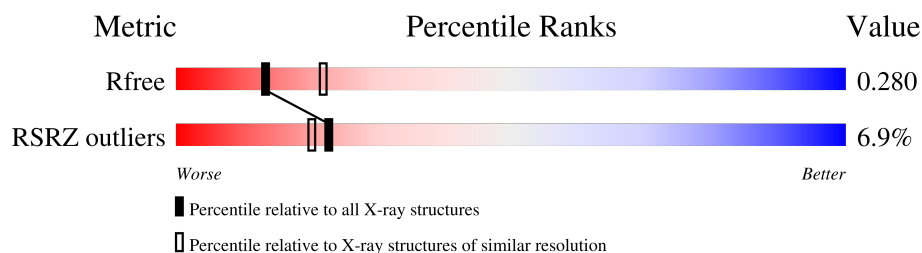
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

The reported resolution of this entry is 2.50 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	5829 (2.50-2.50)
RSRZ outliers	180081	5833 (2.50-2.50)

MolProbity failed to run properly - the sequence quality summary graphics cannot be shown.

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	3638	-	-	-	X
57	MG	1A	4019	-	-	-	X

2 Entry composition

There are 62 unique types of molecules in this entry. The entry contains 297682 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
			27	20	3	3	1			
56	2v	3	Total	C	N	O	S	0	0	0
			27	20	3	3	1			

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

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	1A	1022	Total	Mg	0	0
			1022	1022		
57	1B	30	Total	Mg	0	0
			30	30		
57	1D	15	Total	Mg	0	0
			15	15		
57	1E	12	Total	Mg	0	0
			12	12		
57	1F	16	Total	Mg	0	0
			16	16		

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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	5	Total 5	Mg 5	0	0
57	1Q	7	Total 7	Mg 7	0	0
57	1R	4	Total 4	Mg 4	0	0
57	1S	1	Total 1	Mg 1	0	0
57	1T	5	Total 5	Mg 5	0	0
57	1U	4	Total 4	Mg 4	0	0
57	1V	7	Total 7	Mg 7	0	0
57	1W	3	Total 3	Mg 3	0	0
57	1X	1	Total 1	Mg 1	0	0
57	1Y	1	Total 1	Mg 1	0	0
57	1Z	1	Total 1	Mg 1	0	0
57	10	9	Total 9	Mg 9	0	0
57	11	5	Total 5	Mg 5	0	0
57	13	3	Total 3	Mg 3	0	0
57	15	9	Total 9	Mg 9	0	0
57	17	6	Total 6	Mg 6	0	0
57	18	3	Total 3	Mg 3	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	19	2	Total 2	Mg 2	0	0
57	1a	274	Total 274	Mg 274	0	0
57	1b	1	Total 1	Mg 1	0	0
57	1d	6	Total 6	Mg 6	0	0
57	1e	3	Total 3	Mg 3	0	0
57	1f	2	Total 2	Mg 2	0	0
57	1g	1	Total 1	Mg 1	0	0
57	1h	2	Total 2	Mg 2	0	0
57	1i	1	Total 1	Mg 1	0	0
57	1l	3	Total 3	Mg 3	0	0
57	1m	1	Total 1	Mg 1	0	0
57	1n	3	Total 3	Mg 3	0	0
57	1o	2	Total 2	Mg 2	0	0
57	1t	1	Total 1	Mg 1	0	0
57	1u	1	Total 1	Mg 1	0	0
57	1y	4	Total 4	Mg 4	0	0
57	1w	1	Total 1	Mg 1	0	0
57	1x	2	Total 2	Mg 2	0	0
57	2A	733	Total 733	Mg 733	0	0
57	2B	19	Total 19	Mg 19	0	0
57	2D	10	Total 10	Mg 10	0	0

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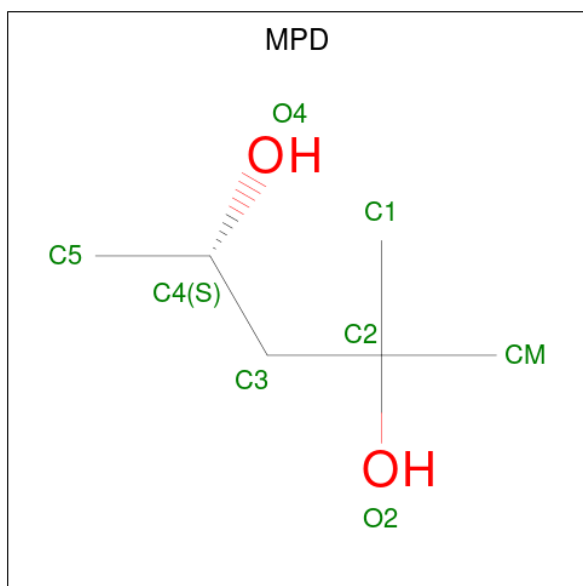
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	2E	7	Total 7	Mg 7	0	0
57	2F	4	Total 4	Mg 4	0	0
57	2G	3	Total 3	Mg 3	0	0
57	2I	1	Total 1	Mg 1	0	0
57	2O	2	Total 2	Mg 2	0	0
57	2P	2	Total 2	Mg 2	0	0
57	2Q	4	Total 4	Mg 4	0	0
57	2R	3	Total 3	Mg 3	0	0
57	2T	3	Total 3	Mg 3	0	0
57	2U	1	Total 1	Mg 1	0	0
57	2V	3	Total 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	21	1	Total 1	Mg 1	0	0
57	25	3	Total 3	Mg 3	0	0
57	27	2	Total 2	Mg 2	0	0
57	28	2	Total 2	Mg 2	0	0
57	2a	190	Total 190	Mg 190	0	0
57	2e	1	Total 1	Mg 1	0	0

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

- Molecule 58 is (4S)-2-METHYL-2,4-PENTANEDIOL (CCD ID: MPD) (formula: C₆H₁₄O₂).



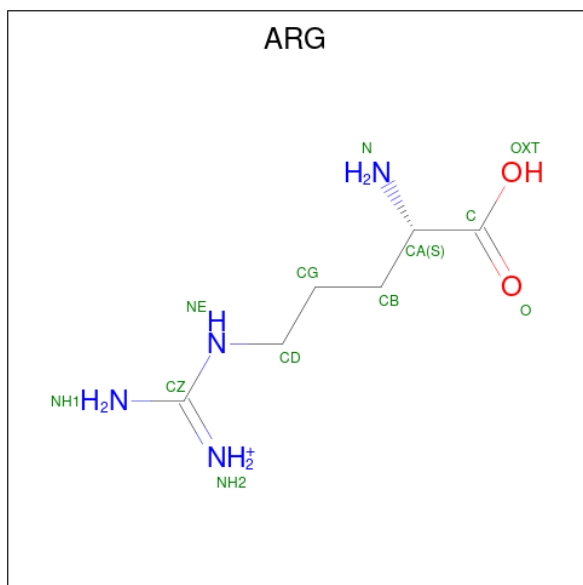
Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
58	1A	1	Total C O 8 6 2	0	0
58	18	1	Total C O 8 6 2	0	0
58	1a	1	Total C O 8 6 2	0	0
58	1a	1	Total C O 8 6 2	0	0

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Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
58	2A	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).

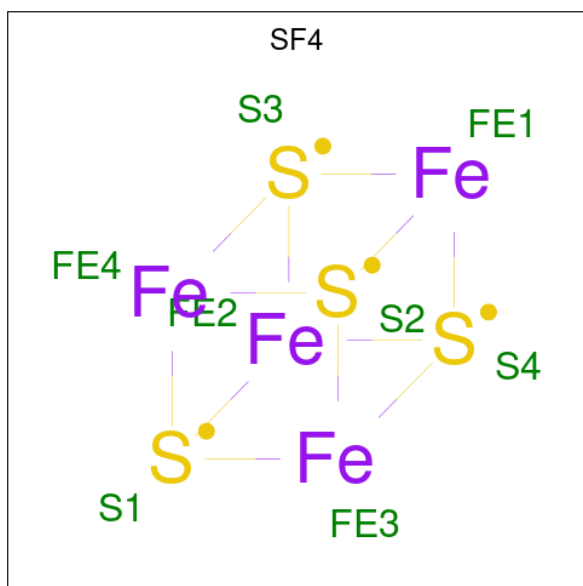
Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
60	1Y	1	Total	Zn	0	0
			1	1		
60	14	1	Total	Zn	0	0
			1	1		
60	15	1	Total	Zn	0	0
			1	1		
60	16	1	Total	Zn	0	0
			1	1		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
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		
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	3972	Total 3972	O 3972	0	0
62	1B	98	Total 98	O 98	0	0
62	1D	107	Total 107	O 107	0	0
62	1E	75	Total 75	O 75	0	0
62	1F	60	Total 60	O 60	0	0
62	1G	20	Total 20	O 20	0	0
62	1H	15	Total 15	O 15	0	0
62	1I	4	Total 4	O 4	0	0
62	1N	47	Total 47	O 47	0	0
62	1O	26	Total 26	O 26	0	0
62	1P	63	Total 63	O 63	0	0
62	1Q	37	Total 37	O 37	0	0
62	1R	34	Total 34	O 34	0	0
62	1S	15	Total 15	O 15	0	0
62	1T	32	Total 32	O 32	0	0
62	1U	45	Total 45	O 45	0	0
62	1V	36	Total 36	O 36	0	0
62	1W	25	Total 25	O 25	0	0
62	1X	26	Total 26	O 26	0	0
62	1Y	16	Total 16	O 16	0	0
62	1Z	8	Total 8	O 8	0	0
62	10	21	Total 21	O 21	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
62	11	26	Total 26	O 26	0	0
62	12	12	Total 12	O 12	0	0
62	13	21	Total 21	O 21	0	0
62	14	2	Total 2	O 2	0	0
62	15	25	Total 25	O 25	0	0
62	16	18	Total 18	O 18	0	0
62	17	13	Total 13	O 13	0	0
62	18	26	Total 26	O 26	0	0
62	19	5	Total 5	O 5	0	0
62	1a	509	Total 509	O 509	0	0
62	1b	1	Total 1	O 1	0	0
62	1d	10	Total 10	O 10	0	0
62	1e	3	Total 3	O 3	0	0
62	1f	3	Total 3	O 3	0	0
62	1h	1	Total 1	O 1	0	0
62	1j	2	Total 2	O 2	0	0
62	1l	4	Total 4	O 4	0	0
62	1m	1	Total 1	O 1	0	0
62	1o	4	Total 4	O 4	0	0
62	1p	3	Total 3	O 3	0	0
62	1t	1	Total 1	O 1	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
62	1u	1	Total 1	O 1	0	0
62	1y	8	Total 8	O 8	0	0
62	1w	3	Total 3	O 3	0	0
62	1x	3	Total 3	O 3	0	0
62	2A	2105	Total 2105	O 2105	0	0
62	2B	46	Total 46	O 46	0	0
62	2D	48	Total 48	O 48	0	0
62	2E	26	Total 26	O 26	0	0
62	2F	24	Total 24	O 24	0	0
62	2G	4	Total 4	O 4	0	0
62	2H	2	Total 2	O 2	0	0
62	2I	4	Total 4	O 4	0	0
62	2N	5	Total 5	O 5	0	0
62	2O	18	Total 18	O 18	0	0
62	2P	19	Total 19	O 19	0	0
62	2Q	18	Total 18	O 18	0	0
62	2R	22	Total 22	O 22	0	0
62	2S	3	Total 3	O 3	0	0
62	2T	7	Total 7	O 7	0	0
62	2U	10	Total 10	O 10	0	0
62	2V	7	Total 7	O 7	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
62	2W	17	Total 17	O 17	0	0
62	2X	8	Total 8	O 8	0	0
62	2Y	5	Total 5	O 5	0	0
62	2Z	11	Total 11	O 11	0	0
62	20	12	Total 12	O 12	0	0
62	21	20	Total 20	O 20	0	0
62	22	1	Total 1	O 1	0	0
62	23	4	Total 4	O 4	0	0
62	25	10	Total 10	O 10	0	0
62	26	5	Total 5	O 5	0	0
62	27	8	Total 8	O 8	0	0
62	28	12	Total 12	O 12	0	0
62	29	2	Total 2	O 2	0	0
62	2a	291	Total 291	O 291	0	0
62	2d	3	Total 3	O 3	0	0
62	2f	3	Total 3	O 3	0	0
62	2j	2	Total 2	O 2	0	0
62	2l	2	Total 2	O 2	0	0
62	2m	2	Total 2	O 2	0	0
62	2o	4	Total 4	O 4	0	0
62	2p	2	Total 2	O 2	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
62	2q	1	Total	O	0	0
			1	1		
62	2r	3	Total	O	0	0
			3	3		
62	2t	2	Total	O	0	0
			2	2		
62	2y	3	Total	O	0	0
			3	3		
62	2w	3	Total	O	0	0
			3	3		
62	2x	1	Total	O	0	0
			1	1		

MolProbity failed to run properly - this section is therefore empty.

3 Data and refinement statistics

Property	Value	Source
Space group	P 21 21 21	Depositor
Cell constants a, b, c, α , β , γ	209.32Å 449.63Å 620.21Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	155.05 – 2.50 155.05 – 2.50	Depositor EDS
% Data completeness (in resolution range)	99.9 (155.05-2.50) 99.9 (155.05-2.50)	Depositor EDS
R_{merge}	0.25	Depositor
R_{sym}	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ ¹	1.29 (at 2.52Å)	Xtriage
Refinement program	PHENIX 1.8.2	Depositor
R, R_{free}	0.230 , 0.278 0.233 , 0.280	Depositor DCC
R_{free} test set	99670 reflections (5.01%)	wwPDB-VP
Wilson B-factor (Å ²)	48.3	Xtriage
Anisotropy	0.142	Xtriage
Bulk solvent k_{sol} (e/Å ³), B_{sol} (Å ²)	0.33 , 62.4	EDS
L-test for twinning ²	$\langle L \rangle = 0.35$, $\langle L^2 \rangle = 0.18$	Xtriage
Estimated twinning fraction	No twinning to report.	Xtriage
F_o, F_c correlation	0.90	EDS
Total number of atoms	297682	wwPDB-VP
Average B, all atoms (Å ²)	56.0	wwPDB-VP

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 1.62% 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.

4 Model quality [i](#)

4.1 Standard geometry [i](#)

MolProbity failed to run properly - this section is therefore empty.

4.2 Too-close contacts [i](#)

MolProbity failed to run properly - this section is therefore empty.

4.3 Torsion angles [i](#)

4.3.1 Protein backbone [i](#)

MolProbity failed to run properly - this section is therefore empty.

4.3.2 Protein sidechains [i](#)

MolProbity failed to run properly - this section is therefore empty.

4.3.3 RNA [i](#)

MolProbity failed to run properly - this section is therefore empty.

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

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	$\# Z > 2$
32	UR3	1a	1498	32	19,22,23	0.97	1 (5%)	26,32,35	1.85	3 (11%)
32	4OC	2a	1402	32	20,23,24	0.77	0	25,32,35	0.94	1 (4%)
32	PSU	2a	516	57,32	18,21,22	1.37	2 (11%)	21,30,33	2.06	5 (23%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
32	MA6	2a	1518	32	23,26,27	0.39	0	33,38,41	2.05	9 (27%)
1	PSU	1A	2605	1	18,21,22	1.43	4 (22%)	21,30,33	2.15	5 (23%)
43	0TD	1l	92	43	8,9,10	4.46	1 (12%)	6,11,13	3.17	2 (33%)
1	PSU	1A	1911	1	18,21,22	1.36	2 (11%)	21,30,33	2.05	4 (19%)
32	5MC	2a	1407	32	19,22,23	1.75	2 (10%)	26,32,35	1.13	3 (11%)
54	PPU	2w	76	1,54	37,40,41	1.33	4 (10%)	47,57,60	1.95	12 (25%)
1	5MU	1A	1915	1	19,22,23	1.43	4 (21%)	27,32,35	2.29	9 (33%)
1	5MU	1A	1939	57,1	19,22,23	1.41	5 (26%)	27,32,35	2.27	6 (22%)
1	PSU	2A	1917	1	18,21,22	1.34	3 (16%)	21,30,33	2.00	4 (19%)
32	5MC	2a	967	32	19,22,23	1.83	2 (10%)	26,32,35	1.02	2 (7%)
32	G7M	1a	527	57,32	23,26,27	1.53	3 (13%)	34,39,42	1.63	4 (11%)
32	5MC	1a	1404	32	19,22,23	1.71	3 (15%)	26,32,35	1.14	3 (11%)
32	5MC	1a	1407	32	19,22,23	1.69	2 (10%)	26,32,35	1.16	3 (11%)
1	OMG	2A	2251	57,1,55	23,26,27	1.17	3 (13%)	32,38,41	2.01	6 (18%)
32	5MC	2a	1400	32	19,22,23	1.67	3 (15%)	26,32,35	1.24	3 (11%)
1	2MA	1A	2503	57,1	22,25,26	1.48	6 (27%)	32,37,40	2.43	9 (28%)
32	M2G	2a	966	57,32	24,27,28	1.29	3 (12%)	33,40,43	1.80	5 (15%)
32	MA6	1a	1519	32	23,26,27	0.41	0	33,38,41	2.09	10 (30%)
1	5MC	2A	1962	57,1	19,22,23	1.46	3 (15%)	26,32,35	1.20	4 (15%)
32	MA6	1a	1518	32	23,26,27	0.41	0	33,38,41	1.93	9 (27%)
32	2MG	2a	1207	32	23,26,27	1.22	3 (13%)	33,38,41	2.13	8 (24%)
1	PSU	2A	1911	1	18,21,22	1.39	2 (11%)	21,30,33	2.03	3 (14%)
43	0TD	2l	92	43	8,9,10	4.45	1 (12%)	6,11,13	6.67	3 (50%)
1	5MC	2A	1942	1	19,22,23	1.80	3 (15%)	26,32,35	1.30	2 (7%)
1	2MA	2A	2503	57,1	22,25,26	1.45	3 (13%)	32,37,40	2.36	9 (28%)
32	PSU	1a	516	57,32	18,21,22	1.36	2 (11%)	21,30,33	1.94	5 (23%)
32	4OC	1a	1402	32	20,23,24	0.76	0	25,32,35	0.90	1 (4%)
1	PSU	2A	2605	1	18,21,22	1.38	3 (16%)	21,30,33	1.95	4 (19%)
1	OMU	2A	2552	57,1	19,22,23	1.22	2 (10%)	25,31,34	1.83	6 (24%)
54	PPU	1w	76	1,54	37,40,41	1.23	3 (8%)	47,57,60	2.11	14 (29%)
32	5MC	1a	967	32	19,22,23	1.59	3 (15%)	26,32,35	1.14	3 (11%)
1	OMU	1A	2552	57,1	19,22,23	1.32	3 (15%)	25,31,34	1.79	6 (24%)
1	PSU	1A	1917	1	18,21,22	1.34	2 (11%)	21,30,33	1.95	3 (14%)
1	5MU	2A	1939	57,1	19,22,23	1.53	5 (26%)	27,32,35	2.24	8 (29%)
55	8AN	1x	76	57,55,56	21,24,25	1.42	3 (14%)	26,35,38	2.97	10 (38%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
1	5MC	1A	1962	57,1	19,22,23	1.59	3 (15%)	26,32,35	1.12	3 (11%)
1	OMG	1A	2251	57,1,55	23,26,27	1.22	3 (13%)	32,38,41	2.02	6 (18%)
32	G7M	2a	527	57,32	23,26,27	1.49	3 (13%)	34,39,42	1.62	4 (11%)
32	5MC	1a	1400	32	19,22,23	1.75	3 (15%)	26,32,35	1.23	2 (7%)
32	MA6	2a	1519	32	23,26,27	0.42	0	33,38,41	2.11	9 (27%)
32	M2G	1a	966	32	24,27,28	1.34	3 (12%)	33,40,43	1.87	5 (15%)
32	UR3	2a	1498	32	19,22,23	1.02	2 (10%)	26,32,35	1.64	3 (11%)
55	8AN	2x	76	57,55,56	21,24,25	1.38	3 (14%)	26,35,38	2.45	11 (42%)
32	5MC	2a	1404	32	19,22,23	1.79	3 (15%)	26,32,35	1.15	2 (7%)
32	2MG	1a	1207	32	23,26,27	1.27	3 (13%)	33,38,41	2.21	7 (21%)
1	OMC	2A	1920	1	19,22,23	0.81	0	25,31,34	1.03	1 (4%)
1	5MU	2A	1915	1	19,22,23	1.35	4 (21%)	27,32,35	2.13	7 (25%)
1	OMC	1A	1920	1	19,22,23	0.85	0	25,31,34	1.15	2 (8%)
1	5MC	1A	1942	57,1	19,22,23	1.57	3 (15%)	26,32,35	1.13	2 (7%)

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
32	UR3	1a	1498	32	-	0/7/25/26	0/2/2/2
32	4OC	2a	1402	32	-	0/9/29/30	0/2/2/2
32	PSU	2a	516	57,32	-	0/7/25/26	0/2/2/2
32	MA6	2a	1518	32	-	0/11/29/30	0/3/3/3
1	PSU	1A	2605	1	-	0/7/25/26	0/2/2/2
43	0TD	1l	92	43	-	1/7/12/14	-
1	PSU	1A	1911	1	-	2/7/25/26	0/2/2/2
32	5MC	2a	1407	32	-	0/7/25/26	0/2/2/2
54	PPU	2w	76	1,54	-	2/25/43/44	0/4/4/4
1	5MU	1A	1915	1	-	2/7/25/26	0/2/2/2
1	5MU	1A	1939	57,1	-	0/7/25/26	0/2/2/2
1	PSU	2A	1917	1	-	0/7/25/26	0/2/2/2
32	5MC	2a	967	32	-	0/7/25/26	0/2/2/2
32	G7M	1a	527	57,32	-	1/7/25/26	0/3/3/3
32	5MC	1a	1404	32	-	0/7/25/26	0/2/2/2
32	5MC	1a	1407	32	-	0/7/25/26	0/2/2/2
1	OMG	2A	2251	57,1,55	-	1/9/27/28	0/3/3/3
32	5MC	2a	1400	32	-	6/7/25/26	0/2/2/2

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Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
1	2MA	1A	2503	57,1	-	1/7/25/26	0/3/3/3
32	M2G	2a	966	57,32	-	0/11/29/30	0/3/3/3
32	MA6	1a	1519	32	-	2/11/29/30	0/3/3/3
1	5MC	2A	1962	57,1	-	0/7/25/26	0/2/2/2
32	MA6	1a	1518	32	-	0/11/29/30	0/3/3/3
32	2MG	2a	1207	32	-	0/9/27/28	0/3/3/3
1	PSU	2A	1911	1	-	0/7/25/26	0/2/2/2
43	0TD	2l	92	43	-	1/7/12/14	-
1	5MC	2A	1942	1	-	0/7/25/26	0/2/2/2
1	2MA	2A	2503	57,1	-	0/7/25/26	0/3/3/3
32	PSU	1a	516	57,32	-	1/7/25/26	0/2/2/2
32	4OC	1a	1402	32	-	3/9/29/30	0/2/2/2
1	PSU	2A	2605	1	-	0/7/25/26	0/2/2/2
1	OMU	2A	2552	57,1	-	0/9/27/28	0/2/2/2
54	PPU	1w	76	1,54	-	2/25/43/44	0/4/4/4
32	5MC	1a	967	32	-	0/7/25/26	0/2/2/2
1	OMU	1A	2552	57,1	-	0/9/27/28	0/2/2/2
1	PSU	1A	1917	1	-	0/7/25/26	0/2/2/2
1	5MU	2A	1939	57,1	-	0/7/25/26	0/2/2/2
55	8AN	1x	76	57,55,56	-	3/7/25/26	0/3/3/3
1	5MC	1A	1962	57,1	-	0/7/25/26	0/2/2/2
1	OMG	1A	2251	57,1,55	-	0/9/27/28	0/3/3/3
32	G7M	2a	527	57,32	-	2/7/25/26	0/3/3/3
32	5MC	1a	1400	32	-	0/7/25/26	0/2/2/2
32	MA6	2a	1519	32	-	3/11/29/30	0/3/3/3
32	M2G	1a	966	32	-	0/11/29/30	0/3/3/3
32	UR3	2a	1498	32	-	0/7/25/26	0/2/2/2
55	8AN	2x	76	57,55,56	-	3/7/25/26	0/3/3/3
32	5MC	2a	1404	32	-	0/7/25/26	0/2/2/2
32	2MG	1a	1207	32	-	1/9/27/28	0/3/3/3
1	OMC	2A	1920	1	-	1/9/27/28	0/2/2/2
1	5MU	2A	1915	1	-	2/7/25/26	0/2/2/2
1	OMC	1A	1920	1	-	2/9/27/28	0/2/2/2
1	5MC	1A	1942	57,1	-	0/7/25/26	0/2/2/2

The worst 5 of 127 bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
43	2l	92	0TD	CB-SB	-12.23	1.69	1.82
43	1l	92	0TD	CB-SB	-12.15	1.70	1.82

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
32	2a	967	5MC	C5-C4	6.83	1.49	1.44
1	2A	1942	5MC	C5-C4	6.67	1.49	1.44
32	2a	1404	5MC	C5-C4	6.66	1.49	1.44

The worst 5 of 270 bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
43	2l	92	0TD	CSB-SB-CB	-15.74	74.08	102.36
55	1x	76	8AN	O4'-C1'-N9	10.22	127.73	108.09
1	2A	2503	2MA	C5-C4-N3	-8.00	118.75	127.18
1	1A	2503	2MA	C5-C4-N3	-7.97	118.78	127.18
32	1a	1498	UR3	C4-N3-C2	-7.43	118.60	124.58

There are no chirality outliers.

5 of 42 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
1	2A	1915	5MU	O4'-C1'-N1-C2
1	2A	1915	5MU	O4'-C1'-N1-C6
32	2a	1400	5MC	C2'-C1'-N1-C6

There are no ring outliers.

No monomer is involved in short contacts.

4.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

4.6 Ligand geometry [i](#)

Of 2521 ligands modelled in this entry, 2510 are monoatomic - leaving 11 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	2A	3735	-	7,7,7	0.31	0	9,10,10	0.24	0
59	ARG	1B	231	57	10,11,11	0.82	1 (10%)	9,13,13	1.05	1 (11%)
61	SF4	2d	501	35	0,12,12	-	-	-	-	-
58	MPD	1A	4023	-	7,7,7	0.29	0	9,10,10	0.38	0
61	SF4	1d	307	35	0,12,12	-	-	-	-	-
58	MPD	18	104	-	7,7,7	0.33	0	9,10,10	0.55	0
59	ARG	1F	317	57	10,11,11	0.75	1 (10%)	9,13,13	1.06	1 (11%)
58	MPD	1a	3276	-	7,7,7	0.39	0	9,10,10	0.43	0
58	MPD	2B	220	-	7,7,7	0.35	0	9,10,10	0.32	0
58	MPD	2A	3734	-	7,7,7	0.33	0	9,10,10	0.36	0
58	MPD	1a	3275	-	7,7,7	0.30	0	9,10,10	0.40	0

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	2A	3735	-	-	3/5/5/5	-
59	ARG	1B	231	57	-	2/11/11/11	-
61	SF4	2d	501	35	-	-	0/6/5/5
58	MPD	1A	4023	-	-	1/5/5/5	-
61	SF4	1d	307	35	-	-	0/6/5/5
58	MPD	18	104	-	-	1/5/5/5	-
59	ARG	1F	317	57	-	2/11/11/11	-
58	MPD	1a	3276	-	-	5/5/5/5	-
58	MPD	2B	220	-	-	0/5/5/5	-
58	MPD	2A	3734	-	-	0/5/5/5	-
58	MPD	1a	3275	-	-	0/5/5/5	-

All (2) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
59	1B	231	ARG	OXT-C	-2.44	1.22	1.30
59	1F	317	ARG	OXT-C	-2.11	1.23	1.30

All (2) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
59	1F	317	ARG	OXT-C-O	-3.10	117.04	124.08
59	1B	231	ARG	OXT-C-O	-2.84	117.63	124.08

There are no chirality outliers.

5 of 14 torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
58	18	104	MPD	C2-C3-C4-C5
58	1a	3276	MPD	C2-C3-C4-O4
58	1a	3276	MPD	C2-C3-C4-C5
59	1F	317	ARG	O-C-CA-N
59	1F	317	ARG	OXT-C-CA-N

There are no ring outliers.

No monomer is involved in short contacts.

4.7 Other polymers [i](#)

There are no such residues in this entry.

4.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

5 Fit of model and data ⓘ

5.1 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	110 (3%) 44 39	19, 36, 84, 98	0
1	2A	2856/2915 (97%)	0.03	111 (3%) 43 38	33, 55, 87, 99	0
2	1B	120/121 (99%)	-0.27	1 (0%) 82 80	29, 47, 61, 78	0
2	2B	120/121 (99%)	0.65	2 (1%) 69 65	57, 71, 79, 84	0
3	1D	275/276 (99%)	-0.04	2 (0%) 84 81	22, 36, 51, 61	0
3	2D	275/276 (99%)	0.44	4 (1%) 72 68	29, 50, 61, 74	0
4	1E	204/206 (99%)	0.07	1 (0%) 87 85	20, 40, 58, 70	0
4	2E	204/206 (99%)	0.39	1 (0%) 87 85	31, 53, 68, 76	0
5	1F	203/210 (96%)	0.10	3 (1%) 72 68	19, 41, 64, 77	0
5	2F	203/210 (96%)	0.72	8 (3%) 43 38	33, 62, 74, 79	0
6	1G	181/182 (99%)	0.89	14 (7%) 19 17	43, 59, 71, 78	0
6	2G	181/182 (99%)	1.55	46 (25%) 1 1	64, 74, 79, 83	0
7	1H	174/180 (96%)	0.45	6 (3%) 48 43	38, 51, 62, 66	0
7	2H	173/180 (96%)	1.37	27 (15%) 5 4	65, 72, 77, 84	0
8	1I	147/148 (99%)	0.96	10 (6%) 23 20	43, 67, 74, 77	0
8	2I	146/148 (98%)	1.49	34 (23%) 2 1	57, 70, 78, 80	0
9	1N	140/140 (100%)	0.02	0 100 100	23, 38, 56, 67	0
9	2N	140/140 (100%)	0.74	3 (2%) 63 59	43, 60, 71, 75	0
10	1O	122/122 (100%)	0.00	0 100 100	29, 40, 58, 68	0
10	2O	122/122 (100%)	0.42	1 (0%) 82 80	41, 53, 65, 68	0
11	1P	149/150 (99%)	0.23	0 100 100	19, 45, 64, 71	0
11	2P	149/150 (99%)	0.86	9 (6%) 27 24	37, 61, 74, 82	0
12	1Q	141/141 (100%)	0.04	1 (0%) 84 81	25, 39, 52, 60	0
12	2Q	141/141 (100%)	0.92	7 (4%) 34 30	45, 60, 68, 72	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
13	1R	118/118 (100%)	-0.03	0 100 100	25, 35, 47, 60	0
13	2R	118/118 (100%)	0.51	3 (2%) 58 54	38, 50, 60, 67	0
14	1S	110/112 (98%)	0.54	1 (0%) 81 78	35, 48, 58, 63	0
14	2S	110/112 (98%)	1.37	16 (14%) 6 5	60, 67, 73, 74	0
15	1T	131/146 (89%)	0.17	3 (2%) 61 57	31, 45, 62, 73	0
15	2T	131/146 (89%)	0.48	1 (0%) 82 80	45, 55, 69, 75	0
16	1U	116/118 (98%)	-0.18	1 (0%) 81 78	21, 29, 46, 59	0
16	2U	116/118 (98%)	0.74	3 (2%) 57 52	38, 56, 68, 71	0
17	1V	101/101 (100%)	0.06	0 100 100	20, 42, 56, 64	0
17	2V	101/101 (100%)	0.84	1 (0%) 79 76	40, 63, 70, 74	0
18	1W	112/113 (99%)	-0.14	0 100 100	23, 33, 50, 74	0
18	2W	112/113 (99%)	0.39	4 (3%) 46 41	37, 48, 62, 80	0
19	1X	95/96 (98%)	0.10	3 (3%) 50 46	25, 36, 57, 71	0
19	2X	95/96 (98%)	0.87	6 (6%) 26 23	48, 57, 68, 75	0
20	1Y	107/110 (97%)	0.44	2 (1%) 66 62	36, 47, 61, 71	0
20	2Y	107/110 (97%)	1.12	9 (8%) 17 15	56, 65, 72, 73	0
21	1Z	203/206 (98%)	0.59	5 (2%) 58 54	37, 55, 67, 76	0
21	2Z	201/206 (97%)	1.33	30 (14%) 5 4	60, 70, 76, 81	0
22	10	83/85 (97%)	0.06	0 100 100	26, 35, 50, 56	0
22	20	83/85 (97%)	1.09	7 (8%) 17 15	41, 57, 66, 70	0
23	11	97/98 (98%)	0.35	1 (1%) 79 76	28, 43, 61, 68	0
23	21	97/98 (98%)	0.68	2 (2%) 63 59	38, 54, 69, 73	0
24	12	70/72 (97%)	0.44	1 (1%) 73 70	36, 47, 57, 73	0
24	22	70/72 (97%)	0.88	4 (5%) 29 26	56, 63, 68, 70	0
25	13	59/60 (98%)	0.06	0 100 100	26, 36, 59, 65	0
25	23	59/60 (98%)	0.84	3 (5%) 33 29	48, 58, 69, 75	0
26	14	69/71 (97%)	1.21	14 (20%) 3 2	50, 70, 81, 84	0
26	24	69/71 (97%)	1.76	19 (27%) 1 1	73, 79, 86, 89	0
27	15	59/60 (98%)	-0.12	1 (1%) 69 65	20, 34, 49, 59	0
27	25	59/60 (98%)	0.23	1 (1%) 69 65	34, 49, 63, 69	0
28	16	53/54 (98%)	0.10	1 (1%) 66 62	31, 41, 52, 61	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
28	26	53/54 (98%)	0.62	1 (1%) 66 62	48, 56, 65, 66	0
29	17	48/49 (97%)	-0.14	1 (2%) 63 59	20, 28, 47, 52	0
29	27	48/49 (97%)	0.18	3 (6%) 26 23	32, 41, 57, 65	0
30	18	64/65 (98%)	-0.13	0 100 100	25, 34, 40, 51	0
30	28	64/65 (98%)	0.72	1 (1%) 70 67	45, 51, 58, 64	0
31	19	37/37 (100%)	-0.01	0 100 100	28, 39, 54, 59	0
31	29	37/37 (100%)	1.06	2 (5%) 31 27	51, 61, 68, 71	0
32	1a	1488/1521 (97%)	0.25	28 (1%) 66 62	35, 64, 84, 98	0
32	2a	1492/1521 (98%)	0.53	39 (2%) 57 52	48, 71, 86, 97	0
33	1b	231/256 (90%)	1.16	36 (15%) 5 4	60, 71, 79, 84	0
33	2b	231/256 (90%)	1.61	65 (28%) 1 1	64, 76, 81, 84	0
34	1c	206/239 (86%)	1.01	14 (6%) 23 20	56, 67, 74, 82	0
34	2c	206/239 (86%)	1.57	62 (30%) 1 1	66, 76, 81, 84	0
35	1d	208/209 (99%)	1.05	25 (12%) 9 7	52, 65, 74, 82	0
35	2d	208/209 (99%)	1.29	34 (16%) 4 3	56, 67, 73, 78	0
36	1e	148/162 (91%)	0.79	5 (3%) 48 43	48, 62, 69, 76	0
36	2e	148/162 (91%)	1.09	15 (10%) 12 10	55, 68, 74, 82	0
37	1f	100/101 (99%)	0.68	2 (2%) 65 61	50, 60, 70, 72	0
37	2f	100/101 (99%)	0.76	2 (2%) 65 61	56, 66, 72, 78	0
38	1g	155/156 (99%)	0.78	6 (3%) 43 38	55, 66, 71, 76	0
38	2g	155/156 (99%)	1.55	41 (26%) 1 1	66, 74, 79, 81	0
39	1h	137/138 (99%)	0.84	9 (6%) 24 21	50, 63, 69, 71	0
39	2h	137/138 (99%)	1.14	7 (5%) 33 29	62, 68, 74, 77	0
40	1i	127/128 (99%)	1.44	20 (15%) 5 4	55, 70, 76, 79	0
40	2i	126/128 (98%)	2.21	69 (54%) 0 0	69, 77, 82, 87	0
41	1j	97/105 (92%)	1.37	20 (20%) 2 2	56, 69, 79, 81	0
41	2j	96/105 (91%)	2.20	50 (52%) 0 0	68, 77, 81, 85	0
42	1k	114/129 (88%)	0.62	4 (3%) 47 42	44, 60, 68, 73	0
42	2k	114/129 (88%)	1.08	14 (12%) 8 7	57, 68, 75, 77	0
43	1l	121/132 (91%)	0.85	7 (5%) 29 25	48, 57, 68, 72	0
43	2l	121/132 (91%)	0.99	9 (7%) 20 18	54, 62, 68, 73	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
44	1m	116/126 (92%)	1.07	11 (9%) 14 12	56, 67, 74, 79	0
44	2m	114/126 (90%)	1.76	41 (35%) 1 1	69, 76, 80, 81	0
45	1n	60/61 (98%)	1.28	9 (15%) 5 4	56, 65, 71, 72	0
45	2n	60/61 (98%)	2.19	38 (63%) 0 0	69, 75, 80, 85	0
46	1o	88/89 (98%)	0.84	3 (3%) 48 43	45, 60, 70, 75	0
46	2o	88/89 (98%)	1.12	7 (7%) 18 16	56, 66, 74, 77	0
47	1p	82/88 (93%)	1.31	12 (14%) 6 4	60, 66, 75, 83	0
47	2p	82/88 (93%)	1.19	6 (7%) 21 19	57, 66, 72, 79	0
48	1q	99/105 (94%)	1.02	8 (8%) 18 15	51, 62, 69, 74	0
48	2q	99/105 (94%)	0.87	2 (2%) 65 61	57, 65, 73, 75	0
49	1r	68/88 (77%)	0.61	1 (1%) 72 68	52, 59, 72, 76	0
49	2r	68/88 (77%)	1.10	6 (8%) 15 14	62, 70, 76, 80	0
50	1s	83/93 (89%)	1.03	6 (7%) 21 19	57, 68, 75, 79	0
50	2s	83/93 (89%)	1.86	32 (38%) 1 0	70, 77, 82, 84	0
51	1t	96/106 (90%)	1.11	14 (14%) 6 4	57, 65, 73, 74	0
51	2t	98/106 (92%)	0.92	3 (3%) 51 47	54, 65, 74, 78	0
52	1u	23/27 (85%)	1.52	5 (21%) 2 2	61, 67, 71, 73	0
52	2u	23/27 (85%)	2.63	15 (65%) 0 0	72, 75, 79, 80	0
53	1y	97/113 (85%)	1.01	9 (9%) 14 12	46, 59, 68, 79	0
53	2y	96/113 (84%)	2.44	62 (64%) 0 0	69, 78, 83, 91	0
54	1w	3/4 (75%)	-0.10	0 100 100	36, 36, 42, 43	0
54	2w	3/4 (75%)	-0.20	0 100 100	49, 49, 52, 64	0
55	1x	3/4 (75%)	-0.09	0 100 100	36, 36, 37, 48	0
55	2x	3/4 (75%)	1.05	1 (33%) 1 1	49, 49, 54, 65	0
56	1v	3/3 (100%)	0.20	0 100 100	32, 32, 32, 38	0
56	2v	3/3 (100%)	1.47	1 (33%) 1 1	48, 48, 54, 60	0
All	All	20796/21490 (96%)	0.47	1426 (6%) 23 20	19, 60, 80, 99	0

The worst 5 of 1426 RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
53	1y	98	ALA	9.2
1	1A	1087	G	6.4

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Mol	Chain	Res	Type	RSRZ
44	1m	2	ALA	6.1
45	2n	2	ALA	6.1
32	2a	1030(A)	G	6.0

5.2 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	PSU	2A	1917	20/21	0.82	0.11	64,74,85,92	0
1	5MU	2A	1915	21/22	0.83	0.12	73,80,85,96	0
32	2MG	2a	1207	24/25	0.84	0.13	72,77,87,90	0
1	5MU	1A	1915	21/22	0.86	0.11	62,71,74,87	0
32	PSU	2a	516	20/21	0.88	0.11	68,70,79,81	0
32	M2G	2a	966	25/26	0.88	0.16	58,71,83,94	0
1	PSU	2A	1911	20/21	0.88	0.11	63,68,81,82	0
1	PSU	1A	1917	20/21	0.89	0.10	62,67,70,75	0
32	5MC	2a	967	21/22	0.90	0.14	65,69,74,76	0
32	2MG	1a	1207	24/25	0.90	0.13	59,68,73,75	0
1	OMC	2A	1920	21/22	0.91	0.12	57,65,69,72	0
1	PSU	1A	1911	20/21	0.92	0.10	59,61,66,67	0
43	0TD	1l	92	10/11	0.92	0.11	54,59,62,71	0
43	0TD	2l	92	10/11	0.92	0.13	63,65,69,72	0
32	G7M	2a	527	24/25	0.93	0.12	60,63,67,71	0
32	MA6	1a	1519	24/25	0.94	0.11	41,49,53,55	0
32	5MC	1a	967	21/22	0.94	0.11	56,64,70,75	0
32	5MC	2a	1400	21/22	0.94	0.14	61,70,75,78	0
32	4OC	2a	1402	22/23	0.94	0.11	60,66,69,73	0
32	5MC	2a	1404	21/22	0.94	0.11	51,61,66,72	0
32	5MC	2a	1407	21/22	0.94	0.11	54,65,68,69	0
32	UR3	2a	1498	21/22	0.94	0.11	57,61,66,69	0
1	5MC	2A	1942	21/22	0.94	0.10	43,52,58,58	0
32	G7M	1a	527	24/25	0.95	0.10	43,55,60,61	0
32	UR3	1a	1498	21/22	0.95	0.11	43,49,52,55	0
32	M2G	1a	966	25/26	0.95	0.10	53,57,62,65	0
32	MA6	2a	1518	24/25	0.95	0.10	58,63,66,69	0
32	MA6	2a	1519	24/25	0.95	0.11	57,65,69,74	0
32	PSU	1a	516	20/21	0.95	0.09	53,62,66,70	0
55	8AN	2x	76	22/23	0.95	0.09	41,45,53,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
32	5MC	1a	1404	21/22	0.96	0.10	44,50,54,56	0
1	5MU	2A	1939	21/22	0.96	0.09	33,39,43,43	0
32	5MC	1a	1407	21/22	0.96	0.10	38,51,57,60	0
1	5MC	2A	1962	21/22	0.96	0.09	41,47,54,57	0
1	OMC	1A	1920	21/22	0.96	0.10	38,54,60,63	0
54	PPU	2w	76	37/38	0.96	0.09	30,41,45,51	0
32	4OC	1a	1402	22/23	0.96	0.09	43,50,52,68	0
1	PSU	2A	2605	20/21	0.97	0.08	29,38,42,44	0
32	5MC	1a	1400	21/22	0.97	0.10	48,52,57,60	0
1	PSU	1A	2605	20/21	0.97	0.08	24,28,32,35	0
32	MA6	1a	1518	24/25	0.97	0.09	42,47,50,50	0
1	5MC	1A	1942	21/22	0.97	0.07	30,36,42,46	0
1	OMG	2A	2251	24/25	0.97	0.08	30,38,42,45	0
1	2MA	2A	2503	23/24	0.97	0.08	30,35,39,41	0
55	8AN	1x	76	22/23	0.97	0.08	28,35,41,48	0
1	OMU	2A	2552	21/22	0.97	0.08	33,40,44,45	0
1	5MC	1A	1962	21/22	0.98	0.06	33,38,43,44	0
1	OMG	1A	2251	24/25	0.98	0.07	20,25,29,30	0
54	PPU	1w	76	37/38	0.98	0.07	23,28,32,33	0
1	2MA	1A	2503	23/24	0.98	0.06	17,21,24,28	0
1	OMU	1A	2552	21/22	0.98	0.07	21,27,33,33	0
1	5MU	1A	1939	21/22	0.98	0.07	23,28,31,34	0

5.3 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.4 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	4019	1/1	0.48	0.42	80,80,80,80	0
57	MG	2A	3677	1/1	0.59	0.21	63,63,63,63	0
57	MG	2A	3530	1/1	0.63	0.30	75,75,75,75	0
57	MG	1a	3135	1/1	0.67	0.22	79,79,79,79	0
57	MG	1a	3153	1/1	0.67	0.19	71,71,71,71	0
57	MG	1B	227	1/1	0.67	0.21	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	1P	203	1/1	0.67	0.22	51,51,51,51	0
57	MG	2a	3132	1/1	0.67	0.25	81,81,81,81	0
58	MPD	2B	220	8/8	0.67	0.29	53,65,71,77	0
57	MG	1A	3658	1/1	0.68	0.15	44,44,44,44	0
57	MG	1A	3885	1/1	0.69	0.27	53,53,53,53	0
57	MG	2A	3619	1/1	0.70	0.24	60,60,60,60	0
57	MG	2A	3728	1/1	0.71	0.15	76,76,76,76	0
57	MG	2A	3666	1/1	0.71	0.16	75,75,75,75	0
57	MG	1a	3273	1/1	0.71	0.19	63,63,63,63	0
57	MG	2A	3091	1/1	0.72	0.20	84,84,84,84	0
57	MG	2A	3457	1/1	0.72	0.20	71,71,71,71	0
57	MG	2a	3184	1/1	0.72	0.21	67,67,67,67	0
57	MG	1A	3507	1/1	0.72	0.19	76,76,76,76	0
57	MG	2a	3164	1/1	0.73	0.15	71,71,71,71	0
57	MG	2a	3010	1/1	0.73	0.26	81,81,81,81	0
57	MG	2B	206	1/1	0.73	0.21	74,74,74,74	0
57	MG	1a	3199	1/1	0.74	0.18	59,59,59,59	0
57	MG	1a	3145	1/1	0.74	0.19	67,67,67,67	0
57	MG	2j	201	1/1	0.74	0.12	71,71,71,71	0
57	MG	1A	3866	1/1	0.74	0.22	70,70,70,70	0
57	MG	2A	3731	1/1	0.75	0.20	64,64,64,64	0
57	MG	1a	3228	1/1	0.75	0.18	72,72,72,72	0
57	MG	1A	3746	1/1	0.75	0.23	63,63,63,63	0
57	MG	1A	3694	1/1	0.75	0.20	63,63,63,63	0
57	MG	1a	3133	1/1	0.76	0.16	63,63,63,63	0
57	MG	1A	3621	1/1	0.76	0.19	37,37,37,37	0
57	MG	1A	3868	1/1	0.76	0.12	40,40,40,40	0
57	MG	2A	3174	1/1	0.76	0.16	63,63,63,63	0
57	MG	2B	207	1/1	0.76	0.20	70,70,70,70	0
57	MG	2A	3301	1/1	0.76	0.38	64,64,64,64	0
57	MG	2a	3064	1/1	0.76	0.29	75,75,75,75	0
57	MG	1A	3812	1/1	0.76	0.18	55,55,55,55	0
57	MG	1a	3156	1/1	0.76	0.13	64,64,64,64	0
57	MG	2A	3534	1/1	0.76	0.20	70,70,70,70	0
57	MG	1a	3189	1/1	0.76	0.12	77,77,77,77	0
57	MG	1a	3052	1/1	0.76	0.16	76,76,76,76	0
57	MG	1B	222	1/1	0.77	0.22	74,74,74,74	0
57	MG	1A	3638	1/1	0.77	0.45	38,38,38,38	0
57	MG	2B	203	1/1	0.77	0.22	71,71,71,71	0
57	MG	1a	3176	1/1	0.78	0.16	74,74,74,74	0
57	MG	2a	3068	1/1	0.78	0.19	75,75,75,75	0
57	MG	1A	3968	1/1	0.78	0.23	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	3994	1/1	0.78	0.16	72,72,72,72	0
57	MG	2A	3216	1/1	0.78	0.14	56,56,56,56	0
57	MG	2A	3287	1/1	0.78	0.22	67,67,67,67	0
57	MG	1A	3865	1/1	0.78	0.24	46,46,46,46	0
57	MG	1A	3850	1/1	0.79	0.13	63,63,63,63	0
57	MG	1A	3671	1/1	0.79	0.18	52,52,52,52	0
57	MG	2A	3313	1/1	0.79	0.24	68,68,68,68	0
57	MG	2U	201	1/1	0.79	0.12	64,64,64,64	0
57	MG	1a	3209	1/1	0.79	0.17	68,68,68,68	0
57	MG	2a	3062	1/1	0.79	0.35	69,69,69,69	0
57	MG	1A	3779	1/1	0.79	0.12	34,34,34,34	0
57	MG	1A	4004	1/1	0.79	0.17	51,51,51,51	0
57	MG	1a	3164	1/1	0.79	0.29	79,79,79,79	0
57	MG	2A	3114	1/1	0.79	0.19	70,70,70,70	0
57	MG	2A	3157	1/1	0.79	0.20	63,63,63,63	0
57	MG	1a	3175	1/1	0.79	0.13	63,63,63,63	0
57	MG	1A	3483	1/1	0.79	0.14	31,31,31,31	0
57	MG	1A	3768	1/1	0.80	0.16	55,55,55,55	0
57	MG	1A	3461	1/1	0.80	0.09	16,16,16,16	0
57	MG	1a	3081	1/1	0.80	0.27	72,72,72,72	0
57	MG	2A	3533	1/1	0.80	0.19	71,71,71,71	0
57	MG	1a	3100	1/1	0.80	0.23	78,78,78,78	0
57	MG	2A	3541	1/1	0.80	0.17	60,60,60,60	0
57	MG	2A	3581	1/1	0.80	0.17	62,62,62,62	0
57	MG	2A	3613	1/1	0.80	0.11	43,43,43,43	0
57	MG	1A	3933	1/1	0.80	0.11	51,51,51,51	0
57	MG	1A	3687	1/1	0.80	0.14	56,56,56,56	0
57	MG	1a	3141	1/1	0.80	0.13	60,60,60,60	0
57	MG	1a	3205	1/1	0.80	0.18	77,77,77,77	0
57	MG	1A	3979	1/1	0.80	0.12	29,29,29,29	0
57	MG	2A	3326	1/1	0.81	0.16	63,63,63,63	0
57	MG	1a	3040	1/1	0.81	0.20	66,66,66,66	0
57	MG	1a	3263	1/1	0.81	0.16	55,55,55,55	0
57	MG	1A	3608	1/1	0.81	0.22	70,70,70,70	0
57	MG	1A	3951	1/1	0.81	0.10	60,60,60,60	0
57	MG	1a	3099	1/1	0.81	0.19	59,59,59,59	0
57	MG	1A	4020	1/1	0.81	0.22	59,59,59,59	0
57	MG	2A	3599	1/1	0.81	0.15	49,49,49,49	0
57	MG	1a	3108	1/1	0.81	0.17	56,56,56,56	0
57	MG	1a	3119	1/1	0.81	0.17	69,69,69,69	0
57	MG	1A	3502	1/1	0.81	0.13	50,50,50,50	0
57	MG	1A	3433	1/1	0.81	0.13	32,32,32,32	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3718	1/1	0.81	0.16	71,71,71,71	0
57	MG	1A	3605	1/1	0.81	0.11	52,52,52,52	0
57	MG	1A	3378	1/1	0.82	0.15	60,60,60,60	0
57	MG	1a	3010	1/1	0.82	0.26	58,58,58,58	0
57	MG	2A	3386	1/1	0.82	0.15	62,62,62,62	0
57	MG	1a	3218	1/1	0.82	0.27	71,71,71,71	0
57	MG	2A	3476	1/1	0.82	0.20	63,63,63,63	0
57	MG	2A	3508	1/1	0.82	0.16	72,72,72,72	0
57	MG	1a	3178	1/1	0.82	0.19	70,70,70,70	0
57	MG	1a	3104	1/1	0.82	0.19	61,61,61,61	0
57	MG	1a	3011	1/1	0.82	0.14	69,69,69,69	0
57	MG	2A	3043	1/1	0.82	0.23	71,71,71,71	0
57	MG	2A	3575	1/1	0.82	0.12	62,62,62,62	0
57	MG	2A	3327	1/1	0.83	0.19	65,65,65,65	0
57	MG	2A	3368	1/1	0.83	0.15	60,60,60,60	0
57	MG	1B	228	1/1	0.83	0.12	52,52,52,52	0
57	MG	1G	203	1/1	0.83	0.13	67,67,67,67	0
57	MG	2A	3001	1/1	0.83	0.23	55,55,55,55	0
57	MG	2A	3503	1/1	0.83	0.14	63,63,63,63	0
57	MG	1A	3196	1/1	0.83	0.14	66,66,66,66	0
57	MG	2G	203	1/1	0.83	0.25	72,72,72,72	0
57	MG	2A	3090	1/1	0.83	0.20	70,70,70,70	0
57	MG	1T	203	1/1	0.83	0.14	65,65,65,65	0
57	MG	2a	3015	1/1	0.83	0.20	61,61,61,61	0
57	MG	1A	3513	1/1	0.83	0.14	31,31,31,31	0
57	MG	1a	3123	1/1	0.83	0.22	78,78,78,78	0
57	MG	1A	3572	1/1	0.83	0.13	42,42,42,42	0
57	MG	1a	3203	1/1	0.83	0.27	75,75,75,75	0
57	MG	2a	3134	1/1	0.83	0.12	63,63,63,63	0
57	MG	1A	3468	1/1	0.83	0.14	50,50,50,50	0
57	MG	1A	3389	1/1	0.83	0.08	31,31,31,31	0
57	MG	1a	3067	1/1	0.83	0.12	54,54,54,54	0
57	MG	1A	3221	1/1	0.83	0.12	65,65,65,65	0
57	MG	1A	3218	1/1	0.84	0.20	64,64,64,64	0
57	MG	1A	3518	1/1	0.84	0.12	50,50,50,50	0
57	MG	2A	3611	1/1	0.84	0.16	43,43,43,43	0
57	MG	2A	3246	1/1	0.84	0.34	62,62,62,62	0
57	MG	1G	204	1/1	0.84	0.13	50,50,50,50	0
57	MG	1A	3969	1/1	0.84	0.21	59,59,59,59	0
57	MG	1A	3976	1/1	0.84	0.17	55,55,55,55	0
57	MG	2A	3314	1/1	0.84	0.14	37,37,37,37	0
57	MG	1W	201	1/1	0.84	0.20	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	3627	1/1	0.84	0.11	22,22,22,22	0
57	MG	1a	3140	1/1	0.84	0.26	58,58,58,58	0
57	MG	2A	3376	1/1	0.84	0.15	69,69,69,69	0
57	MG	2A	3377	1/1	0.84	0.10	39,39,39,39	0
57	MG	2B	211	1/1	0.84	0.10	59,59,59,59	0
57	MG	1a	3256	1/1	0.84	0.20	61,61,61,61	0
57	MG	2A	3398	1/1	0.84	0.17	59,59,59,59	0
57	MG	2a	3009	1/1	0.84	0.33	70,70,70,70	0
57	MG	1A	3530	1/1	0.84	0.11	56,56,56,56	0
57	MG	2A	3462	1/1	0.84	0.18	53,53,53,53	0
57	MG	2a	3023	1/1	0.84	0.19	69,69,69,69	0
57	MG	2a	3059	1/1	0.84	0.25	71,71,71,71	0
57	MG	1A	3547	1/1	0.84	0.11	44,44,44,44	0
57	MG	1A	3571	1/1	0.84	0.11	65,65,65,65	0
57	MG	1A	3419	1/1	0.84	0.12	25,25,25,25	0
57	MG	2a	3120	1/1	0.84	0.15	71,71,71,71	0
57	MG	2A	3522	1/1	0.84	0.14	35,35,35,35	0
57	MG	1a	3159	1/1	0.84	0.11	56,56,56,56	0
57	MG	2a	3148	1/1	0.84	0.12	67,67,67,67	0
57	MG	1a	3070	1/1	0.84	0.22	73,73,73,73	0
57	MG	1A	3578	1/1	0.84	0.12	40,40,40,40	0
57	MG	2A	3137	1/1	0.84	0.27	67,67,67,67	0
57	MG	1A	3425	1/1	0.84	0.11	35,35,35,35	0
57	MG	1A	3306	1/1	0.85	0.24	70,70,70,70	0
57	MG	2A	3322	1/1	0.85	0.12	52,52,52,52	0
57	MG	1l	201	1/1	0.85	0.19	68,68,68,68	0
57	MG	1a	3191	1/1	0.85	0.12	69,69,69,69	0
57	MG	1a	3195	1/1	0.85	0.14	63,63,63,63	0
57	MG	2A	3699	1/1	0.85	0.11	71,71,71,71	0
57	MG	2A	3717	1/1	0.85	0.20	63,63,63,63	0
57	MG	2A	3062	1/1	0.85	0.21	70,70,70,70	0
57	MG	2A	3724	1/1	0.85	0.20	66,66,66,66	0
57	MG	1A	3990	1/1	0.85	0.11	47,47,47,47	0
57	MG	2A	3730	1/1	0.85	0.12	52,52,52,52	0
57	MG	1A	3792	1/1	0.85	0.11	49,49,49,49	0
57	MG	2A	3104	1/1	0.85	0.23	63,63,63,63	0
57	MG	2A	3442	1/1	0.85	0.13	56,56,56,56	0
57	MG	2A	3447	1/1	0.85	0.18	64,64,64,64	0
57	MG	1A	3959	1/1	0.85	0.12	65,65,65,65	0
57	MG	2B	212	1/1	0.85	0.16	67,67,67,67	0
57	MG	1A	3330	1/1	0.85	0.16	61,61,61,61	0
57	MG	2A	3468	1/1	0.85	0.19	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	2A	3469	1/1	0.85	0.09	67,67,67,67	0
57	MG	1A	3222	1/1	0.85	0.10	59,59,59,59	0
57	MG	2A	3160	1/1	0.85	0.23	63,63,63,63	0
57	MG	2A	3167	1/1	0.85	0.21	56,56,56,56	0
57	MG	1a	3220	1/1	0.85	0.15	76,76,76,76	0
57	MG	2A	3523	1/1	0.85	0.10	48,48,48,48	0
57	MG	2A	3528	1/1	0.85	0.15	65,65,65,65	0
57	MG	2A	3175	1/1	0.85	0.20	70,70,70,70	0
57	MG	2A	3181	1/1	0.85	0.31	74,74,74,74	0
57	MG	1A	3893	1/1	0.85	0.13	54,54,54,54	0
57	MG	1a	3229	1/1	0.85	0.14	66,66,66,66	0
57	MG	2a	3143	1/1	0.85	0.21	64,64,64,64	0
57	MG	2A	3561	1/1	0.85	0.13	58,58,58,58	0
57	MG	2A	3568	1/1	0.85	0.14	43,43,43,43	0
57	MG	1a	3242	1/1	0.85	0.19	64,64,64,64	0
57	MG	1a	3094	1/1	0.85	0.24	69,69,69,69	0
57	MG	1a	3096	1/1	0.85	0.27	69,69,69,69	0
59	ARG	1F	317	12/12	0.85	0.15	48,60,72,73	0
57	MG	2A	3588	1/1	0.86	0.11	54,54,54,54	0
57	MG	2A	3363	1/1	0.86	0.09	48,48,48,48	0
57	MG	1A	3808	1/1	0.86	0.14	70,70,70,70	0
57	MG	1A	3643	1/1	0.86	0.23	58,58,58,58	0
57	MG	1a	3109	1/1	0.86	0.26	72,72,72,72	0
57	MG	1A	3843	1/1	0.86	0.12	56,56,56,56	0
57	MG	2A	3668	1/1	0.86	0.09	51,51,51,51	0
57	MG	2A	3397	1/1	0.86	0.10	37,37,37,37	0
57	MG	1A	3292	1/1	0.86	0.15	67,67,67,67	0
57	MG	2A	3700	1/1	0.86	0.12	80,80,80,80	0
57	MG	2A	3408	1/1	0.86	0.15	52,52,52,52	0
57	MG	2A	3437	1/1	0.86	0.36	49,49,49,49	0
57	MG	2A	3723	1/1	0.86	0.15	65,65,65,65	0
57	MG	1A	3582	1/1	0.86	0.37	40,40,40,40	0
57	MG	2A	3726	1/1	0.86	0.12	69,69,69,69	0
57	MG	2A	3443	1/1	0.86	0.15	51,51,51,51	0
57	MG	2A	3131	1/1	0.86	0.27	63,63,63,63	0
57	MG	2A	3451	1/1	0.86	0.14	53,53,53,53	0
57	MG	2A	3732	1/1	0.86	0.27	66,66,66,66	0
57	MG	1A	3675	1/1	0.86	0.10	71,71,71,71	0
57	MG	1A	3475	1/1	0.86	0.10	20,20,20,20	0
57	MG	2A	3465	1/1	0.86	0.13	63,63,63,63	0
57	MG	1A	3477	1/1	0.86	0.08	18,18,18,18	0
57	MG	1a	3069	1/1	0.86	0.10	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	3707	1/1	0.86	0.14	65,65,65,65	0
57	MG	2O	202	1/1	0.86	0.11	69,69,69,69	0
57	MG	1A	3393	1/1	0.86	0.09	29,29,29,29	0
57	MG	1a	3232	1/1	0.86	0.16	47,47,47,47	0
57	MG	2A	3510	1/1	0.86	0.13	58,58,58,58	0
57	MG	2a	3011	1/1	0.86	0.22	58,58,58,58	0
57	MG	2A	3511	1/1	0.86	0.17	73,73,73,73	0
57	MG	2A	3198	1/1	0.86	0.31	61,61,61,61	0
57	MG	2a	3047	1/1	0.86	0.24	69,69,69,69	0
57	MG	2a	3056	1/1	0.86	0.16	74,74,74,74	0
57	MG	1A	3624	1/1	0.86	0.12	29,29,29,29	0
57	MG	2a	3060	1/1	0.86	0.25	76,76,76,76	0
57	MG	2A	3526	1/1	0.86	0.35	52,52,52,52	0
57	MG	1a	3253	1/1	0.86	0.10	51,51,51,51	0
57	MG	1G	202	1/1	0.86	0.09	42,42,42,42	0
57	MG	2A	3291	1/1	0.86	0.19	62,62,62,62	0
57	MG	1a	3167	1/1	0.86	0.09	39,39,39,39	0
57	MG	2A	3535	1/1	0.86	0.17	60,60,60,60	0
57	MG	2A	3539	1/1	0.86	0.15	67,67,67,67	0
57	MG	1a	3168	1/1	0.86	0.20	60,60,60,60	0
57	MG	1a	3274	1/1	0.86	0.19	68,68,68,68	0
57	MG	2A	3564	1/1	0.86	0.12	66,66,66,66	0
57	MG	1b	301	1/1	0.86	0.09	75,75,75,75	0
57	MG	1A	3444	1/1	0.86	0.11	55,55,55,55	0
57	MG	1A	3275	1/1	0.86	0.28	62,62,62,62	0
57	MG	1A	3697	1/1	0.87	0.18	39,39,39,39	0
57	MG	1A	3705	1/1	0.87	0.18	57,57,57,57	0
57	MG	1a	3219	1/1	0.87	0.15	63,63,63,63	0
57	MG	2A	3189	1/1	0.87	0.17	59,59,59,59	0
57	MG	2A	3196	1/1	0.87	0.17	55,55,55,55	0
57	MG	1a	3032	1/1	0.87	0.16	60,60,60,60	0
57	MG	1A	3595	1/1	0.87	0.17	69,69,69,69	0
57	MG	1a	3144	1/1	0.87	0.11	61,61,61,61	0
57	MG	2A	3733	1/1	0.87	0.11	59,59,59,59	0
57	MG	2A	3283	1/1	0.87	0.10	75,75,75,75	0
57	MG	1A	3597	1/1	0.87	0.13	41,41,41,41	0
57	MG	1a	3150	1/1	0.87	0.14	56,56,56,56	0
57	MG	1a	3251	1/1	0.87	0.13	61,61,61,61	0
57	MG	1a	3059	1/1	0.87	0.30	60,60,60,60	0
57	MG	1A	4013	1/1	0.87	0.10	47,47,47,47	0
57	MG	1A	3747	1/1	0.87	0.08	61,61,61,61	0
57	MG	2R	203	1/1	0.87	0.13	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	3883	1/1	0.87	0.07	46,46,46,46	0
57	MG	2V	201	1/1	0.87	0.10	62,62,62,62	0
57	MG	1a	3165	1/1	0.87	0.21	70,70,70,70	0
57	MG	2A	3335	1/1	0.87	0.13	63,63,63,63	0
57	MG	2A	3553	1/1	0.87	0.13	53,53,53,53	0
57	MG	1A	3766	1/1	0.87	0.13	41,41,41,41	0
57	MG	1A	3123	1/1	0.87	0.22	49,49,49,49	0
57	MG	2a	3031	1/1	0.87	0.22	65,65,65,65	0
57	MG	2a	3040	1/1	0.87	0.25	66,66,66,66	0
57	MG	2A	3373	1/1	0.87	0.13	74,74,74,74	0
57	MG	1A	3776	1/1	0.87	0.08	22,22,22,22	0
57	MG	1D	309	1/1	0.87	0.45	48,48,48,48	0
57	MG	1A	3942	1/1	0.87	0.11	67,67,67,67	0
57	MG	1a	3188	1/1	0.87	0.10	80,80,80,80	0
57	MG	1A	3462	1/1	0.87	0.08	35,35,35,35	0
57	MG	2a	3066	1/1	0.87	0.15	68,68,68,68	0
57	MG	1A	3480	1/1	0.87	0.11	27,27,27,27	0
57	MG	2a	3069	1/1	0.87	0.17	57,57,57,57	0
57	MG	2a	3073	1/1	0.87	0.26	69,69,69,69	0
57	MG	2A	3614	1/1	0.87	0.14	62,62,62,62	0
57	MG	2a	3131	1/1	0.87	0.15	72,72,72,72	0
57	MG	1a	3193	1/1	0.87	0.13	59,59,59,59	0
57	MG	2A	3640	1/1	0.87	0.10	59,59,59,59	0
57	MG	2A	3646	1/1	0.87	0.12	55,55,55,55	0
57	MG	1A	3197	1/1	0.87	0.26	64,64,64,64	0
57	MG	2a	3153	1/1	0.87	0.13	63,63,63,63	0
57	MG	1A	3494	1/1	0.87	0.11	32,32,32,32	0
57	MG	1a	3201	1/1	0.87	0.20	72,72,72,72	0
57	MG	2a	3190	1/1	0.87	0.33	64,64,64,64	0
57	MG	1A	3814	1/1	0.87	0.12	43,43,43,43	0
58	MPD	1a	3275	8/8	0.87	0.20	57,66,67,71	0
57	MG	2A	3165	1/1	0.87	0.23	59,59,59,59	0
57	MG	1a	3128	1/1	0.87	0.17	64,64,64,64	0
57	MG	2A	3411	1/1	0.88	0.19	41,41,41,41	0
57	MG	2A	3430	1/1	0.88	0.17	64,64,64,64	0
57	MG	2A	3722	1/1	0.88	0.10	54,54,54,54	0
57	MG	2A	3149	1/1	0.88	0.23	56,56,56,56	0
57	MG	1B	217	1/1	0.88	0.10	54,54,54,54	0
57	MG	2A	3158	1/1	0.88	0.10	63,63,63,63	0
57	MG	2A	3727	1/1	0.88	0.14	56,56,56,56	0
57	MG	1a	3147	1/1	0.88	0.18	64,64,64,64	0
57	MG	1B	219	1/1	0.88	0.09	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	1a	3225	1/1	0.88	0.26	71,71,71,71	0
57	MG	1A	3769	1/1	0.88	0.13	51,51,51,51	0
57	MG	2A	3463	1/1	0.88	0.13	61,61,61,61	0
57	MG	2B	201	1/1	0.88	0.08	64,64,64,64	0
57	MG	1A	3271	1/1	0.88	0.15	49,49,49,49	0
57	MG	2A	3178	1/1	0.88	0.16	46,46,46,46	0
57	MG	1A	3588	1/1	0.88	0.11	40,40,40,40	0
57	MG	2A	3183	1/1	0.88	0.15	61,61,61,61	0
57	MG	2A	3478	1/1	0.88	0.16	64,64,64,64	0
57	MG	2B	218	1/1	0.88	0.09	70,70,70,70	0
57	MG	2E	304	1/1	0.88	0.11	38,38,38,38	0
57	MG	2E	307	1/1	0.88	0.15	63,63,63,63	0
57	MG	2A	3481	1/1	0.88	0.16	67,67,67,67	0
57	MG	1a	3087	1/1	0.88	0.21	57,57,57,57	0
57	MG	2A	3191	1/1	0.88	0.09	72,72,72,72	0
57	MG	1A	3203	1/1	0.88	0.20	44,44,44,44	0
57	MG	1E	304	1/1	0.88	0.09	26,26,26,26	0
57	MG	1A	3463	1/1	0.88	0.21	59,59,59,59	0
57	MG	2A	3219	1/1	0.88	0.20	63,63,63,63	0
57	MG	2A	3231	1/1	0.88	0.18	59,59,59,59	0
57	MG	2A	3233	1/1	0.88	0.28	63,63,63,63	0
57	MG	1A	3600	1/1	0.88	0.08	33,33,33,33	0
57	MG	2a	3028	1/1	0.88	0.13	56,56,56,56	0
57	MG	2A	3248	1/1	0.88	0.18	67,67,67,67	0
57	MG	2a	3033	1/1	0.88	0.13	57,57,57,57	0
57	MG	2a	3037	1/1	0.88	0.14	66,66,66,66	0
57	MG	2a	3038	1/1	0.88	0.16	61,61,61,61	0
57	MG	2A	3277	1/1	0.88	0.29	73,73,73,73	0
57	MG	2a	3042	1/1	0.88	0.23	63,63,63,63	0
57	MG	1A	3516	1/1	0.88	0.11	52,52,52,52	0
57	MG	2a	3049	1/1	0.88	0.26	57,57,57,57	0
57	MG	1N	203	1/1	0.88	0.12	58,58,58,58	0
57	MG	2a	3057	1/1	0.88	0.23	68,68,68,68	0
57	MG	1A	3193	1/1	0.88	0.22	61,61,61,61	0
57	MG	1A	3298	1/1	0.88	0.13	63,63,63,63	0
57	MG	2A	3304	1/1	0.88	0.22	54,54,54,54	0
57	MG	2A	3307	1/1	0.88	0.43	70,70,70,70	0
57	MG	1w	101	1/1	0.88	0.12	52,52,52,52	0
57	MG	1a	3121	1/1	0.88	0.11	52,52,52,52	0
57	MG	2A	3003	1/1	0.88	0.15	62,62,62,62	0
57	MG	2A	3037	1/1	0.88	0.20	62,62,62,62	0
57	MG	1A	3736	1/1	0.88	0.11	47,47,47,47	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3086	1/1	0.88	0.11	44,44,44,44	0
57	MG	2A	3075	1/1	0.88	0.18	64,64,64,64	0
57	MG	2A	3084	1/1	0.88	0.18	63,63,63,63	0
57	MG	1A	3027	1/1	0.88	0.07	65,65,65,65	0
57	MG	2a	3147	1/1	0.88	0.13	53,53,53,53	0
57	MG	2A	3620	1/1	0.88	0.12	54,54,54,54	0
57	MG	1A	3353	1/1	0.88	0.10	31,31,31,31	0
57	MG	1A	3449	1/1	0.88	0.14	58,58,58,58	0
57	MG	2a	3170	1/1	0.88	0.16	82,82,82,82	0
57	MG	1a	3204	1/1	0.88	0.09	63,63,63,63	0
57	MG	2A	3395	1/1	0.88	0.14	60,60,60,60	0
57	MG	1a	3048	1/1	0.88	0.23	61,61,61,61	0
57	MG	2A	3689	1/1	0.88	0.12	55,55,55,55	0
58	MPD	2A	3734	8/8	0.88	0.22	38,49,52,57	0
57	MG	1A	3886	1/1	0.88	0.13	48,48,48,48	0
57	MG	2A	3143	1/1	0.88	0.12	53,53,53,53	0
57	MG	2A	3225	1/1	0.89	0.10	55,55,55,55	0
57	MG	1A	3974	1/1	0.89	0.08	44,44,44,44	0
57	MG	1A	3439	1/1	0.89	0.15	57,57,57,57	0
57	MG	2A	3237	1/1	0.89	0.17	57,57,57,57	0
57	MG	1a	3068	1/1	0.89	0.24	67,67,67,67	0
57	MG	1a	3211	1/1	0.89	0.24	68,68,68,68	0
57	MG	2A	3268	1/1	0.89	0.17	55,55,55,55	0
57	MG	1A	3506	1/1	0.89	0.07	29,29,29,29	0
57	MG	1A	3988	1/1	0.89	0.28	48,48,48,48	0
57	MG	1a	3077	1/1	0.89	0.13	60,60,60,60	0
57	MG	1A	3372	1/1	0.89	0.16	65,65,65,65	0
57	MG	2A	3300	1/1	0.89	0.34	64,64,64,64	0
57	MG	1A	3010	1/1	0.89	0.12	50,50,50,50	0
57	MG	1A	3995	1/1	0.89	0.11	57,57,57,57	0
57	MG	1a	3095	1/1	0.89	0.12	63,63,63,63	0
57	MG	2A	3312	1/1	0.89	0.16	55,55,55,55	0
57	MG	1A	4002	1/1	0.89	0.12	54,54,54,54	0
57	MG	1a	3097	1/1	0.89	0.34	60,60,60,60	0
57	MG	1A	3625	1/1	0.89	0.15	52,52,52,52	0
57	MG	1A	3457	1/1	0.89	0.08	44,44,44,44	0
57	MG	1A	3381	1/1	0.89	0.10	44,44,44,44	0
57	MG	1a	3266	1/1	0.89	0.10	61,61,61,61	0
57	MG	2A	3340	1/1	0.89	0.15	68,68,68,68	0
57	MG	2A	3361	1/1	0.89	0.12	55,55,55,55	0
57	MG	1A	3816	1/1	0.89	0.11	56,56,56,56	0
57	MG	1B	203	1/1	0.89	0.18	64,64,64,64	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1B	207	1/1	0.89	0.16	53,53,53,53	0
57	MG	1d	303	1/1	0.89	0.17	64,64,64,64	0
57	MG	1d	306	1/1	0.89	0.10	78,78,78,78	0
57	MG	2B	210	1/1	0.89	0.30	66,66,66,66	0
57	MG	1h	202	1/1	0.89	0.19	66,66,66,66	0
57	MG	1A	3842	1/1	0.89	0.12	45,45,45,45	0
57	MG	1l	202	1/1	0.89	0.11	56,56,56,56	0
57	MG	1y	203	1/1	0.89	0.14	73,73,73,73	0
57	MG	2A	3404	1/1	0.89	0.09	32,32,32,32	0
57	MG	2G	202	1/1	0.89	0.14	66,66,66,66	0
57	MG	2A	3407	1/1	0.89	0.13	45,45,45,45	0
57	MG	1B	218	1/1	0.89	0.11	46,46,46,46	0
57	MG	2R	201	1/1	0.89	0.15	59,59,59,59	0
57	MG	1a	3126	1/1	0.89	0.14	61,61,61,61	0
57	MG	1A	3210	1/1	0.89	0.11	52,52,52,52	0
57	MG	1A	3531	1/1	0.89	0.09	52,52,52,52	0
57	MG	2W	202	1/1	0.89	0.18	52,52,52,52	0
57	MG	28	102	1/1	0.89	0.13	52,52,52,52	0
57	MG	1A	3857	1/1	0.89	0.08	51,51,51,51	0
57	MG	2A	3048	1/1	0.89	0.19	53,53,53,53	0
57	MG	2A	3049	1/1	0.89	0.17	57,57,57,57	0
57	MG	1A	3860	1/1	0.89	0.10	50,50,50,50	0
57	MG	2a	3021	1/1	0.89	0.16	62,62,62,62	0
57	MG	1A	3533	1/1	0.89	0.10	44,44,44,44	0
57	MG	2a	3024	1/1	0.89	0.25	70,70,70,70	0
57	MG	2a	3026	1/1	0.89	0.30	69,69,69,69	0
57	MG	1A	3537	1/1	0.89	0.22	61,61,61,61	0
57	MG	1A	3680	1/1	0.89	0.14	58,58,58,58	0
57	MG	1A	3313	1/1	0.89	0.18	39,39,39,39	0
57	MG	1A	3466	1/1	0.89	0.10	62,62,62,62	0
57	MG	2A	3112	1/1	0.89	0.11	57,57,57,57	0
57	MG	1a	3151	1/1	0.89	0.17	64,64,64,64	0
57	MG	2A	3120	1/1	0.89	0.27	55,55,55,55	0
57	MG	2A	3126	1/1	0.89	0.26	65,65,65,65	0
57	MG	2A	3484	1/1	0.89	0.10	28,28,28,28	0
57	MG	2A	3490	1/1	0.89	0.14	56,56,56,56	0
57	MG	2A	3499	1/1	0.89	0.11	44,44,44,44	0
57	MG	1A	3401	1/1	0.89	0.08	19,19,19,19	0
57	MG	1A	3418	1/1	0.89	0.11	26,26,26,26	0
57	MG	1A	3917	1/1	0.89	0.09	21,21,21,21	0
57	MG	1A	3041	1/1	0.89	0.23	58,58,58,58	0
57	MG	10	104	1/1	0.89	0.12	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	10	107	1/1	0.89	0.10	55,55,55,55	0
57	MG	10	109	1/1	0.89	0.16	58,58,58,58	0
57	MG	1a	3002	1/1	0.89	0.12	60,60,60,60	0
57	MG	2a	3078	1/1	0.89	0.27	69,69,69,69	0
57	MG	2a	3086	1/1	0.89	0.19	54,54,54,54	0
57	MG	2a	3090	1/1	0.89	0.12	66,66,66,66	0
57	MG	2a	3101	1/1	0.89	0.39	64,64,64,64	0
57	MG	2a	3112	1/1	0.89	0.33	65,65,65,65	0
57	MG	2a	3115	1/1	0.89	0.23	65,65,65,65	0
57	MG	2a	3119	1/1	0.89	0.12	50,50,50,50	0
57	MG	1a	3006	1/1	0.89	0.10	63,63,63,63	0
57	MG	1A	3585	1/1	0.89	0.09	36,36,36,36	0
57	MG	1a	3181	1/1	0.89	0.16	63,63,63,63	0
57	MG	1A	3189	1/1	0.89	0.16	66,66,66,66	0
57	MG	1a	3020	1/1	0.89	0.14	57,57,57,57	0
57	MG	1a	3022	1/1	0.89	0.11	73,73,73,73	0
57	MG	2A	3552	1/1	0.89	0.08	50,50,50,50	0
57	MG	1a	3023	1/1	0.89	0.23	61,61,61,61	0
57	MG	1A	3952	1/1	0.89	0.12	58,58,58,58	0
57	MG	2a	3168	1/1	0.89	0.10	66,66,66,66	0
57	MG	2A	3193	1/1	0.89	0.12	58,58,58,58	0
57	MG	1A	3358	1/1	0.89	0.06	22,22,22,22	0
57	MG	1a	3200	1/1	0.89	0.13	66,66,66,66	0
57	MG	2A	3576	1/1	0.89	0.11	50,50,50,50	0
57	MG	1A	3486	1/1	0.89	0.09	18,18,18,18	0
57	MG	2A	3218	1/1	0.89	0.12	53,53,53,53	0
58	MPD	2A	3735	8/8	0.89	0.19	53,57,64,68	0
57	MG	2A	3596	1/1	0.89	0.09	39,39,39,39	0
57	MG	1A	3438	1/1	0.89	0.10	51,51,51,51	0
57	MG	2A	3130	1/1	0.90	0.20	47,47,47,47	0
57	MG	1A	3441	1/1	0.90	0.14	48,48,48,48	0
57	MG	2A	3132	1/1	0.90	0.17	60,60,60,60	0
57	MG	2A	3412	1/1	0.90	0.09	59,59,59,59	0
57	MG	1A	3601	1/1	0.90	0.11	50,50,50,50	0
57	MG	1A	3604	1/1	0.90	0.18	54,54,54,54	0
57	MG	2A	3438	1/1	0.90	0.19	60,60,60,60	0
57	MG	1a	3089	1/1	0.90	0.14	55,55,55,55	0
57	MG	1A	3398	1/1	0.90	0.08	44,44,44,44	0
57	MG	2B	208	1/1	0.90	0.25	55,55,55,55	0
57	MG	1A	3322	1/1	0.90	0.31	59,59,59,59	0
57	MG	1A	3753	1/1	0.90	0.10	52,52,52,52	0
57	MG	1A	3924	1/1	0.90	0.12	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	3932	1/1	0.90	0.12	51,51,51,51	0
57	MG	2D	302	1/1	0.90	0.13	49,49,49,49	0
57	MG	2E	301	1/1	0.90	0.16	47,47,47,47	0
57	MG	1A	3374	1/1	0.90	0.13	46,46,46,46	0
57	MG	1E	311	1/1	0.90	0.10	38,38,38,38	0
57	MG	1F	315	1/1	0.90	0.11	53,53,53,53	0
57	MG	1A	3940	1/1	0.90	0.09	39,39,39,39	0
57	MG	2I	201	1/1	0.90	0.15	64,64,64,64	0
57	MG	1A	3460	1/1	0.90	0.16	43,43,43,43	0
57	MG	2Q	202	1/1	0.90	0.19	53,53,53,53	0
57	MG	2A	3187	1/1	0.90	0.16	63,63,63,63	0
57	MG	2A	3188	1/1	0.90	0.15	60,60,60,60	0
57	MG	2T	201	1/1	0.90	0.15	66,66,66,66	0
57	MG	1a	3231	1/1	0.90	0.10	53,53,53,53	0
57	MG	1A	3949	1/1	0.90	0.12	38,38,38,38	0
57	MG	1a	3236	1/1	0.90	0.09	65,65,65,65	0
57	MG	1A	3558	1/1	0.90	0.18	51,51,51,51	0
57	MG	2a	3005	1/1	0.90	0.25	66,66,66,66	0
57	MG	2a	3008	1/1	0.90	0.11	64,64,64,64	0
57	MG	1A	3129	1/1	0.90	0.15	47,47,47,47	0
57	MG	2A	3214	1/1	0.90	0.15	60,60,60,60	0
57	MG	1A	3955	1/1	0.90	0.09	40,40,40,40	0
57	MG	1A	3635	1/1	0.90	0.15	44,44,44,44	0
57	MG	2a	3019	1/1	0.90	0.06	86,86,86,86	0
57	MG	1X	101	1/1	0.90	0.22	48,48,48,48	0
57	MG	1a	3139	1/1	0.90	0.18	66,66,66,66	0
57	MG	1Z	301	1/1	0.90	0.13	61,61,61,61	0
57	MG	2a	3025	1/1	0.90	0.18	52,52,52,52	0
57	MG	1A	3207	1/1	0.90	0.15	58,58,58,58	0
57	MG	1A	3642	1/1	0.90	0.19	63,63,63,63	0
57	MG	1A	3973	1/1	0.90	0.12	55,55,55,55	0
57	MG	15	3108	1/1	0.90	0.08	48,48,48,48	0
57	MG	2a	3035	1/1	0.90	0.21	59,59,59,59	0
57	MG	2A	3253	1/1	0.90	0.23	55,55,55,55	0
57	MG	2A	3254	1/1	0.90	0.24	57,57,57,57	0
57	MG	2a	3039	1/1	0.90	0.30	82,82,82,82	0
57	MG	2A	3551	1/1	0.90	0.09	60,60,60,60	0
57	MG	1f	202	1/1	0.90	0.06	40,40,40,40	0
57	MG	1A	3575	1/1	0.90	0.12	61,61,61,61	0
57	MG	2A	3281	1/1	0.90	0.19	66,66,66,66	0
57	MG	1a	3004	1/1	0.90	0.12	55,55,55,55	0
57	MG	2A	3566	1/1	0.90	0.12	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	3058	1/1	0.90	0.25	64,64,64,64	0
57	MG	1A	3975	1/1	0.90	0.13	52,52,52,52	0
57	MG	1n	102	1/1	0.90	0.08	53,53,53,53	0
57	MG	1A	3650	1/1	0.90	0.09	49,49,49,49	0
57	MG	1a	3158	1/1	0.90	0.11	55,55,55,55	0
57	MG	2A	3584	1/1	0.90	0.07	55,55,55,55	0
57	MG	1x	102	1/1	0.90	0.09	41,41,41,41	0
57	MG	1A	3977	1/1	0.90	0.09	64,64,64,64	0
57	MG	2A	3597	1/1	0.90	0.11	64,64,64,64	0
57	MG	1a	3161	1/1	0.90	0.19	65,65,65,65	0
57	MG	1A	3270	1/1	0.90	0.26	65,65,65,65	0
57	MG	1A	3824	1/1	0.90	0.23	26,26,26,26	0
57	MG	1A	3989	1/1	0.90	0.14	32,32,32,32	0
57	MG	2A	3618	1/1	0.90	0.11	44,44,44,44	0
57	MG	1A	3436	1/1	0.90	0.09	51,51,51,51	0
57	MG	2A	3055	1/1	0.90	0.13	56,56,56,56	0
57	MG	2A	3057	1/1	0.90	0.11	51,51,51,51	0
57	MG	2a	3122	1/1	0.90	0.14	62,62,62,62	0
57	MG	2a	3124	1/1	0.90	0.09	56,56,56,56	0
57	MG	2a	3129	1/1	0.90	0.10	62,62,62,62	0
57	MG	2A	3336	1/1	0.90	0.17	54,54,54,54	0
57	MG	2A	3657	1/1	0.90	0.09	58,58,58,58	0
57	MG	1a	3174	1/1	0.90	0.09	76,76,76,76	0
57	MG	2A	3347	1/1	0.90	0.13	47,47,47,47	0
57	MG	2a	3145	1/1	0.90	0.22	66,66,66,66	0
57	MG	1A	3583	1/1	0.90	0.19	48,48,48,48	0
57	MG	1A	3847	1/1	0.90	0.11	36,36,36,36	0
57	MG	1A	3679	1/1	0.90	0.16	66,66,66,66	0
57	MG	2a	3154	1/1	0.90	0.08	69,69,69,69	0
57	MG	1a	3053	1/1	0.90	0.22	71,71,71,71	0
57	MG	2A	3704	1/1	0.90	0.09	40,40,40,40	0
57	MG	2A	3716	1/1	0.90	0.15	55,55,55,55	0
57	MG	2a	3173	1/1	0.90	0.09	71,71,71,71	0
57	MG	2A	3095	1/1	0.90	0.11	73,73,73,73	0
57	MG	2A	3103	1/1	0.90	0.16	56,56,56,56	0
57	MG	2A	3721	1/1	0.90	0.14	49,49,49,49	0
57	MG	1A	3853	1/1	0.90	0.10	27,27,27,27	0
57	MG	1A	3365	1/1	0.90	0.09	22,22,22,22	0
57	MG	1A	3397	1/1	0.90	0.08	35,35,35,35	0
57	MG	1A	3519	1/1	0.90	0.10	25,25,25,25	0
57	MG	1A	3523	1/1	0.90	0.11	54,54,54,54	0
57	MG	1A	3340	1/1	0.91	0.10	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	3848	1/1	0.91	0.14	43,43,43,43	0
57	MG	1a	3041	1/1	0.91	0.09	64,64,64,64	0
57	MG	1a	3044	1/1	0.91	0.16	57,57,57,57	0
57	MG	2A	3094	1/1	0.91	0.18	52,52,52,52	0
57	MG	1A	3849	1/1	0.91	0.11	40,40,40,40	0
57	MG	2A	3096	1/1	0.91	0.17	54,54,54,54	0
57	MG	1A	3274	1/1	0.91	0.15	49,49,49,49	0
57	MG	2A	3381	1/1	0.91	0.13	48,48,48,48	0
57	MG	1A	3696	1/1	0.91	0.20	39,39,39,39	0
57	MG	2A	3105	1/1	0.91	0.18	56,56,56,56	0
57	MG	1A	3854	1/1	0.91	0.07	50,50,50,50	0
57	MG	1A	3527	1/1	0.91	0.14	59,59,59,59	0
57	MG	2A	3403	1/1	0.91	0.10	34,34,34,34	0
57	MG	1A	3145	1/1	0.91	0.15	50,50,50,50	0
57	MG	2A	3121	1/1	0.91	0.17	60,60,60,60	0
57	MG	1A	3362	1/1	0.91	0.09	37,37,37,37	0
57	MG	1A	3728	1/1	0.91	0.11	50,50,50,50	0
57	MG	1A	3421	1/1	0.91	0.07	33,33,33,33	0
57	MG	2A	3413	1/1	0.91	0.13	55,55,55,55	0
57	MG	1a	3078	1/1	0.91	0.15	65,65,65,65	0
57	MG	2A	3136	1/1	0.91	0.18	59,59,59,59	0
57	MG	1A	3869	1/1	0.91	0.11	52,52,52,52	0
57	MG	2B	213	1/1	0.91	0.08	69,69,69,69	0
57	MG	2A	3142	1/1	0.91	0.13	46,46,46,46	0
57	MG	1a	3082	1/1	0.91	0.14	80,80,80,80	0
57	MG	2D	306	1/1	0.91	0.15	44,44,44,44	0
57	MG	2D	308	1/1	0.91	0.13	59,59,59,59	0
57	MG	1a	3083	1/1	0.91	0.14	76,76,76,76	0
57	MG	1a	3210	1/1	0.91	0.21	65,65,65,65	0
57	MG	1A	3875	1/1	0.91	0.10	39,39,39,39	0
57	MG	1a	3215	1/1	0.91	0.12	62,62,62,62	0
57	MG	1A	3741	1/1	0.91	0.08	48,48,48,48	0
57	MG	1A	3744	1/1	0.91	0.18	51,51,51,51	0
57	MG	1A	3164	1/1	0.91	0.14	51,51,51,51	0
57	MG	1A	3889	1/1	0.91	0.11	56,56,56,56	0
57	MG	2A	3470	1/1	0.91	0.14	61,61,61,61	0
57	MG	2A	3475	1/1	0.91	0.10	58,58,58,58	0
57	MG	1A	3368	1/1	0.91	0.08	39,39,39,39	0
57	MG	2T	203	1/1	0.91	0.12	60,60,60,60	0
57	MG	2A	3477	1/1	0.91	0.09	39,39,39,39	0
57	MG	1A	3898	1/1	0.91	0.09	56,56,56,56	0
57	MG	1A	3901	1/1	0.91	0.09	26,26,26,26	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	20	102	1/1	0.91	0.09	56,56,56,56	0
57	MG	1E	310	1/1	0.91	0.10	58,58,58,58	0
57	MG	2a	3002	1/1	0.91	0.17	57,57,57,57	0
57	MG	2a	3004	1/1	0.91	0.17	59,59,59,59	0
57	MG	1a	3106	1/1	0.91	0.16	59,59,59,59	0
57	MG	2A	3491	1/1	0.91	0.15	56,56,56,56	0
57	MG	1a	3238	1/1	0.91	0.12	56,56,56,56	0
57	MG	1A	3557	1/1	0.91	0.09	46,46,46,46	0
57	MG	2A	3192	1/1	0.91	0.31	55,55,55,55	0
57	MG	1A	3758	1/1	0.91	0.15	70,70,70,70	0
57	MG	2a	3016	1/1	0.91	0.18	62,62,62,62	0
57	MG	2A	3195	1/1	0.91	0.17	58,58,58,58	0
57	MG	2A	3512	1/1	0.91	0.14	67,67,67,67	0
57	MG	2A	3518	1/1	0.91	0.16	59,59,59,59	0
57	MG	1a	3116	1/1	0.91	0.10	61,61,61,61	0
57	MG	1A	3761	1/1	0.91	0.13	49,49,49,49	0
57	MG	2A	3524	1/1	0.91	0.07	35,35,35,35	0
57	MG	2a	3027	1/1	0.91	0.08	68,68,68,68	0
57	MG	2A	3209	1/1	0.91	0.13	53,53,53,53	0
57	MG	2A	3527	1/1	0.91	0.12	59,59,59,59	0
57	MG	1a	3120	1/1	0.91	0.11	63,63,63,63	0
57	MG	1A	3171	1/1	0.91	0.14	45,45,45,45	0
57	MG	1A	3243	1/1	0.91	0.16	57,57,57,57	0
57	MG	1A	3264	1/1	0.91	0.12	34,34,34,34	0
57	MG	1N	204	1/1	0.91	0.13	51,51,51,51	0
57	MG	1A	3321	1/1	0.91	0.29	66,66,66,66	0
57	MG	2a	3041	1/1	0.91	0.24	56,56,56,56	0
57	MG	1A	3577	1/1	0.91	0.08	27,27,27,27	0
57	MG	2A	3543	1/1	0.91	0.10	67,67,67,67	0
57	MG	1T	204	1/1	0.91	0.12	64,64,64,64	0
57	MG	2a	3054	1/1	0.91	0.10	60,60,60,60	0
57	MG	2A	3238	1/1	0.91	0.09	58,58,58,58	0
57	MG	1g	201	1/1	0.91	0.23	62,62,62,62	0
57	MG	2A	3555	1/1	0.91	0.20	48,48,48,48	0
57	MG	1A	3646	1/1	0.91	0.23	52,52,52,52	0
57	MG	2A	3251	1/1	0.91	0.11	53,53,53,53	0
57	MG	1A	3802	1/1	0.91	0.07	26,26,26,26	0
57	MG	1A	3059	1/1	0.91	0.17	47,47,47,47	0
57	MG	2A	3569	1/1	0.91	0.10	45,45,45,45	0
57	MG	2A	3570	1/1	0.91	0.11	63,63,63,63	0
57	MG	2A	3571	1/1	0.91	0.10	44,44,44,44	0
57	MG	2a	3071	1/1	0.91	0.16	64,64,64,64	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2a	3072	1/1	0.91	0.15	59,59,59,59	0
57	MG	2A	3255	1/1	0.91	0.09	49,49,49,49	0
57	MG	1l	203	1/1	0.91	0.10	63,63,63,63	0
57	MG	2a	3083	1/1	0.91	0.21	64,64,64,64	0
57	MG	1A	3965	1/1	0.91	0.07	35,35,35,35	0
57	MG	2A	3582	1/1	0.91	0.10	50,50,50,50	0
57	MG	2a	3094	1/1	0.91	0.21	55,55,55,55	0
57	MG	1A	3447	1/1	0.91	0.13	55,55,55,55	0
57	MG	1A	3666	1/1	0.91	0.13	21,21,21,21	0
57	MG	2a	3113	1/1	0.91	0.29	53,53,53,53	0
57	MG	2a	3114	1/1	0.91	0.33	62,62,62,62	0
57	MG	1A	3972	1/1	0.91	0.10	56,56,56,56	0
57	MG	17	105	1/1	0.91	0.23	62,62,62,62	0
57	MG	1A	3448	1/1	0.91	0.12	61,61,61,61	0
57	MG	2A	3600	1/1	0.91	0.07	47,47,47,47	0
57	MG	2A	3033	1/1	0.91	0.23	52,52,52,52	0
57	MG	2a	3127	1/1	0.91	0.10	71,71,71,71	0
57	MG	1A	3817	1/1	0.91	0.08	21,21,21,21	0
57	MG	2A	3306	1/1	0.91	0.12	43,43,43,43	0
57	MG	1A	3392	1/1	0.91	0.09	35,35,35,35	0
57	MG	2A	3310	1/1	0.91	0.13	58,58,58,58	0
57	MG	2a	3137	1/1	0.91	0.14	55,55,55,55	0
57	MG	2A	3044	1/1	0.91	0.14	53,53,53,53	0
57	MG	1A	3835	1/1	0.91	0.08	44,44,44,44	0
57	MG	1A	3328	1/1	0.91	0.13	51,51,51,51	0
57	MG	2A	3317	1/1	0.91	0.07	50,50,50,50	0
57	MG	1A	3034	1/1	0.91	0.15	51,51,51,51	0
57	MG	1A	3983	1/1	0.91	0.08	55,55,55,55	0
57	MG	2a	3160	1/1	0.91	0.15	74,74,74,74	0
57	MG	2A	3669	1/1	0.91	0.21	75,75,75,75	0
57	MG	2a	3166	1/1	0.91	0.12	58,58,58,58	0
57	MG	2A	3671	1/1	0.91	0.09	53,53,53,53	0
57	MG	2A	3675	1/1	0.91	0.11	45,45,45,45	0
57	MG	1A	3987	1/1	0.91	0.12	32,32,32,32	0
57	MG	2a	3179	1/1	0.91	0.13	60,60,60,60	0
57	MG	2A	3678	1/1	0.91	0.09	54,54,54,54	0
57	MG	2a	3189	1/1	0.91	0.25	55,55,55,55	0
57	MG	2A	3685	1/1	0.91	0.14	55,55,55,55	0
57	MG	2f	201	1/1	0.91	0.13	60,60,60,60	0
57	MG	2A	3688	1/1	0.91	0.09	59,59,59,59	0
57	MG	2A	3334	1/1	0.91	0.12	52,52,52,52	0
57	MG	2A	3693	1/1	0.91	0.10	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	3064	1/1	0.91	0.14	49,49,49,49	0
57	MG	2A	3067	1/1	0.91	0.09	47,47,47,47	0
57	MG	2A	3702	1/1	0.91	0.08	62,62,62,62	0
57	MG	2A	3698	1/1	0.92	0.08	72,72,72,72	0
57	MG	1D	311	1/1	0.92	0.13	52,52,52,52	0
57	MG	1D	314	1/1	0.92	0.07	67,67,67,67	0
57	MG	1a	3107	1/1	0.92	0.21	52,52,52,52	0
57	MG	1D	315	1/1	0.92	0.11	48,48,48,48	0
57	MG	1n	103	1/1	0.92	0.17	60,60,60,60	0
57	MG	1y	201	1/1	0.92	0.15	66,66,66,66	0
57	MG	1y	202	1/1	0.92	0.14	56,56,56,56	0
57	MG	1A	3751	1/1	0.92	0.08	25,25,25,25	0
57	MG	1a	3110	1/1	0.92	0.17	53,53,53,53	0
57	MG	1E	308	1/1	0.92	0.14	44,44,44,44	0
57	MG	1A	3545	1/1	0.92	0.12	41,41,41,41	0
57	MG	1A	3130	1/1	0.92	0.29	37,37,37,37	0
57	MG	1F	314	1/1	0.92	0.10	41,41,41,41	0
57	MG	1A	3908	1/1	0.92	0.24	55,55,55,55	0
57	MG	2A	3729	1/1	0.92	0.19	53,53,53,53	0
57	MG	2A	3039	1/1	0.92	0.27	58,58,58,58	0
57	MG	1A	3911	1/1	0.92	0.06	20,20,20,20	0
57	MG	1A	3555	1/1	0.92	0.11	45,45,45,45	0
57	MG	2A	3375	1/1	0.92	0.10	40,40,40,40	0
57	MG	1a	3129	1/1	0.92	0.12	64,64,64,64	0
57	MG	1A	3326	1/1	0.92	0.11	41,41,41,41	0
57	MG	2A	3050	1/1	0.92	0.17	66,66,66,66	0
57	MG	2A	3053	1/1	0.92	0.08	48,48,48,48	0
57	MG	1A	3384	1/1	0.92	0.13	56,56,56,56	0
57	MG	1A	3562	1/1	0.92	0.10	57,57,57,57	0
57	MG	2A	3060	1/1	0.92	0.09	43,43,43,43	0
57	MG	2A	3400	1/1	0.92	0.16	55,55,55,55	0
57	MG	1A	3935	1/1	0.92	0.14	58,58,58,58	0
57	MG	2B	215	1/1	0.92	0.07	72,72,72,72	0
57	MG	1Q	206	1/1	0.92	0.10	41,41,41,41	0
57	MG	1a	3142	1/1	0.92	0.09	58,58,58,58	0
57	MG	1R	204	1/1	0.92	0.16	50,50,50,50	0
57	MG	2A	3409	1/1	0.92	0.10	40,40,40,40	0
57	MG	2A	3083	1/1	0.92	0.13	55,55,55,55	0
57	MG	1A	3487	1/1	0.92	0.08	53,53,53,53	0
57	MG	2A	3085	1/1	0.92	0.11	41,41,41,41	0
57	MG	2G	201	1/1	0.92	0.10	70,70,70,70	0
57	MG	1A	3388	1/1	0.92	0.07	27,27,27,27	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3435	1/1	0.92	0.18	56,56,56,56	0
57	MG	1V	206	1/1	0.92	0.20	48,48,48,48	0
57	MG	1A	3948	1/1	0.92	0.12	39,39,39,39	0
57	MG	1a	3152	1/1	0.92	0.08	65,65,65,65	0
57	MG	2Q	203	1/1	0.92	0.10	62,62,62,62	0
57	MG	1A	3785	1/1	0.92	0.12	48,48,48,48	0
57	MG	1A	3787	1/1	0.92	0.07	28,28,28,28	0
57	MG	1A	3573	1/1	0.92	0.08	43,43,43,43	0
57	MG	2A	3455	1/1	0.92	0.11	57,57,57,57	0
57	MG	1A	3794	1/1	0.92	0.10	59,59,59,59	0
57	MG	2A	3111	1/1	0.92	0.11	55,55,55,55	0
57	MG	1A	3956	1/1	0.92	0.10	51,51,51,51	0
57	MG	1A	3796	1/1	0.92	0.07	70,70,70,70	0
57	MG	25	102	1/1	0.92	0.21	45,45,45,45	0
57	MG	2A	3116	1/1	0.92	0.17	51,51,51,51	0
57	MG	2A	3119	1/1	0.92	0.09	54,54,54,54	0
57	MG	1A	3138	1/1	0.92	0.07	43,43,43,43	0
57	MG	1A	3659	1/1	0.92	0.05	19,19,19,19	0
57	MG	1a	3003	1/1	0.92	0.13	53,53,53,53	0
57	MG	1A	3064	1/1	0.92	0.11	59,59,59,59	0
57	MG	1A	3069	1/1	0.92	0.29	29,29,29,29	0
57	MG	1A	3512	1/1	0.92	0.09	17,17,17,17	0
57	MG	2a	3013	1/1	0.92	0.18	66,66,66,66	0
57	MG	1a	3177	1/1	0.92	0.07	59,59,59,59	0
57	MG	2A	3489	1/1	0.92	0.18	67,67,67,67	0
57	MG	1A	3678	1/1	0.92	0.08	55,55,55,55	0
57	MG	2A	3138	1/1	0.92	0.11	44,44,44,44	0
57	MG	2A	3494	1/1	0.92	0.11	42,42,42,42	0
57	MG	1a	3019	1/1	0.92	0.15	58,58,58,58	0
57	MG	2A	3502	1/1	0.92	0.10	60,60,60,60	0
57	MG	1a	3184	1/1	0.92	0.09	69,69,69,69	0
57	MG	2A	3148	1/1	0.92	0.09	55,55,55,55	0
57	MG	1a	3185	1/1	0.92	0.19	72,72,72,72	0
57	MG	2A	3153	1/1	0.92	0.27	58,58,58,58	0
57	MG	2a	3032	1/1	0.92	0.25	57,57,57,57	0
57	MG	1A	3822	1/1	0.92	0.15	55,55,55,55	0
57	MG	1A	3347	1/1	0.92	0.09	46,46,46,46	0
57	MG	1a	3190	1/1	0.92	0.10	55,55,55,55	0
57	MG	1A	3832	1/1	0.92	0.08	36,36,36,36	0
57	MG	1a	3025	1/1	0.92	0.14	60,60,60,60	0
57	MG	2A	3525	1/1	0.92	0.14	50,50,50,50	0
57	MG	2A	3169	1/1	0.92	0.12	62,62,62,62	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3046	1/1	0.92	0.22	54,54,54,54	0
57	MG	2a	3043	1/1	0.92	0.18	70,70,70,70	0
57	MG	2a	3045	1/1	0.92	0.09	54,54,54,54	0
57	MG	1a	3035	1/1	0.92	0.18	52,52,52,52	0
57	MG	1a	3038	1/1	0.92	0.18	64,64,64,64	0
57	MG	2A	3531	1/1	0.92	0.12	63,63,63,63	0
57	MG	1A	3682	1/1	0.92	0.08	68,68,68,68	0
57	MG	1A	3683	1/1	0.92	0.09	66,66,66,66	0
57	MG	1a	3042	1/1	0.92	0.20	71,71,71,71	0
57	MG	1A	3088	1/1	0.92	0.09	46,46,46,46	0
57	MG	1A	3590	1/1	0.92	0.12	54,54,54,54	0
57	MG	1A	3695	1/1	0.92	0.10	49,49,49,49	0
57	MG	2A	3547	1/1	0.92	0.10	32,32,32,32	0
57	MG	2A	3550	1/1	0.92	0.07	62,62,62,62	0
57	MG	1A	3991	1/1	0.92	0.09	50,50,50,50	0
57	MG	1a	3214	1/1	0.92	0.16	71,71,71,71	0
57	MG	1A	3011	1/1	0.92	0.09	50,50,50,50	0
57	MG	1a	3060	1/1	0.92	0.21	65,65,65,65	0
57	MG	2A	3557	1/1	0.92	0.09	60,60,60,60	0
57	MG	2a	3077	1/1	0.92	0.32	60,60,60,60	0
57	MG	1a	3062	1/1	0.92	0.11	57,57,57,57	0
57	MG	2a	3082	1/1	0.92	0.15	58,58,58,58	0
57	MG	1a	3064	1/1	0.92	0.14	58,58,58,58	0
57	MG	2a	3084	1/1	0.92	0.17	79,79,79,79	0
57	MG	2a	3085	1/1	0.92	0.14	60,60,60,60	0
57	MG	2A	3213	1/1	0.92	0.14	56,56,56,56	0
57	MG	1A	3851	1/1	0.92	0.11	61,61,61,61	0
57	MG	1A	3060	1/1	0.92	0.12	36,36,36,36	0
57	MG	1A	3309	1/1	0.92	0.20	25,25,25,25	0
57	MG	2a	3102	1/1	0.92	0.18	51,51,51,51	0
57	MG	1A	4009	1/1	0.92	0.12	45,45,45,45	0
57	MG	2A	3221	1/1	0.92	0.07	53,53,53,53	0
57	MG	1a	3074	1/1	0.92	0.18	60,60,60,60	0
57	MG	2A	3226	1/1	0.92	0.18	59,59,59,59	0
57	MG	1a	3234	1/1	0.92	0.11	63,63,63,63	0
57	MG	1A	3249	1/1	0.92	0.15	42,42,42,42	0
57	MG	1A	3711	1/1	0.92	0.09	33,33,33,33	0
57	MG	1a	3241	1/1	0.92	0.28	62,62,62,62	0
57	MG	1A	3714	1/1	0.92	0.21	30,30,30,30	0
57	MG	2A	3247	1/1	0.92	0.08	54,54,54,54	0
57	MG	2a	3130	1/1	0.92	0.11	65,65,65,65	0
57	MG	1A	3720	1/1	0.92	0.09	35,35,35,35	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3602	1/1	0.92	0.09	54,54,54,54	0
57	MG	1A	3603	1/1	0.92	0.09	30,30,30,30	0
57	MG	2A	3252	1/1	0.92	0.24	66,66,66,66	0
57	MG	2a	3138	1/1	0.92	0.07	67,67,67,67	0
57	MG	1a	3086	1/1	0.92	0.22	61,61,61,61	0
57	MG	1A	3474	1/1	0.92	0.10	59,59,59,59	0
57	MG	1a	3265	1/1	0.92	0.20	64,64,64,64	0
57	MG	2A	3266	1/1	0.92	0.13	63,63,63,63	0
57	MG	1a	3088	1/1	0.92	0.07	58,58,58,58	0
57	MG	2A	3644	1/1	0.92	0.08	48,48,48,48	0
57	MG	2a	3155	1/1	0.92	0.15	61,61,61,61	0
57	MG	2a	3156	1/1	0.92	0.10	72,72,72,72	0
57	MG	2a	3157	1/1	0.92	0.09	68,68,68,68	0
57	MG	1a	3267	1/1	0.92	0.13	55,55,55,55	0
57	MG	2A	3654	1/1	0.92	0.07	23,23,23,23	0
57	MG	2A	3278	1/1	0.92	0.20	45,45,45,45	0
57	MG	2a	3167	1/1	0.92	0.11	67,67,67,67	0
57	MG	1A	3263	1/1	0.92	0.11	59,59,59,59	0
57	MG	2A	3667	1/1	0.92	0.09	50,50,50,50	0
57	MG	1A	3534	1/1	0.92	0.08	38,38,38,38	0
57	MG	2a	3177	1/1	0.92	0.12	59,59,59,59	0
57	MG	1A	3884	1/1	0.92	0.09	42,42,42,42	0
57	MG	2a	3183	1/1	0.92	0.21	66,66,66,66	0
57	MG	1A	3614	1/1	0.92	0.11	43,43,43,43	0
57	MG	2A	3299	1/1	0.92	0.20	53,53,53,53	0
57	MG	1d	304	1/1	0.92	0.10	63,63,63,63	0
57	MG	2e	201	1/1	0.92	0.25	61,61,61,61	0
57	MG	1A	3377	1/1	0.92	0.08	66,66,66,66	0
57	MG	1B	230	1/1	0.92	0.08	61,61,61,61	0
57	MG	2n	101	1/1	0.92	0.22	58,58,58,58	0
57	MG	2A	3686	1/1	0.92	0.10	47,47,47,47	0
57	MG	1A	3748	1/1	0.92	0.09	49,49,49,49	0
57	MG	1a	3103	1/1	0.92	0.13	67,67,67,67	0
57	MG	2A	3309	1/1	0.92	0.10	55,55,55,55	0
59	ARG	1B	231	12/12	0.92	0.10	29,44,51,55	0
57	MG	2A	3695	1/1	0.92	0.12	61,61,61,61	0
57	MG	1B	210	1/1	0.93	0.09	49,49,49,49	0
57	MG	2A	3651	1/1	0.93	0.08	37,37,37,37	0
57	MG	2A	3263	1/1	0.93	0.16	51,51,51,51	0
57	MG	2A	3265	1/1	0.93	0.27	50,50,50,50	0
57	MG	2A	3663	1/1	0.93	0.09	70,70,70,70	0
57	MG	1B	211	1/1	0.93	0.19	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	3718	1/1	0.93	0.35	38,38,38,38	0
57	MG	1A	3619	1/1	0.93	0.12	57,57,57,57	0
57	MG	1A	3726	1/1	0.93	0.10	42,42,42,42	0
57	MG	2A	3279	1/1	0.93	0.26	44,44,44,44	0
57	MG	1e	201	1/1	0.93	0.18	63,63,63,63	0
57	MG	2A	3282	1/1	0.93	0.14	55,55,55,55	0
57	MG	1B	220	1/1	0.93	0.07	28,28,28,28	0
57	MG	2A	3286	1/1	0.93	0.28	44,44,44,44	0
57	MG	1A	3867	1/1	0.93	0.08	38,38,38,38	0
57	MG	1A	3727	1/1	0.93	0.09	42,42,42,42	0
57	MG	2A	3292	1/1	0.93	0.31	64,64,64,64	0
57	MG	2A	3298	1/1	0.93	0.34	56,56,56,56	0
57	MG	1A	3556	1/1	0.93	0.09	39,39,39,39	0
57	MG	2A	3696	1/1	0.93	0.11	62,62,62,62	0
57	MG	1B	229	1/1	0.93	0.13	56,56,56,56	0
57	MG	1A	3730	1/1	0.93	0.10	52,52,52,52	0
57	MG	1n	101	1/1	0.93	0.25	66,66,66,66	0
57	MG	1A	3879	1/1	0.93	0.07	28,28,28,28	0
57	MG	1A	3880	1/1	0.93	0.12	54,54,54,54	0
57	MG	2A	3707	1/1	0.93	0.09	46,46,46,46	0
57	MG	2A	3709	1/1	0.93	0.07	44,44,44,44	0
57	MG	1o	101	1/1	0.93	0.20	58,58,58,58	0
57	MG	1o	102	1/1	0.93	0.10	56,56,56,56	0
57	MG	1A	3881	1/1	0.93	0.07	29,29,29,29	0
57	MG	2A	3719	1/1	0.93	0.09	47,47,47,47	0
57	MG	1A	3357	1/1	0.93	0.07	25,25,25,25	0
57	MG	1a	3111	1/1	0.93	0.10	56,56,56,56	0
57	MG	1a	3113	1/1	0.93	0.10	41,41,41,41	0
57	MG	2A	3320	1/1	0.93	0.12	45,45,45,45	0
57	MG	1A	3740	1/1	0.93	0.06	24,24,24,24	0
57	MG	2A	3323	1/1	0.93	0.10	57,57,57,57	0
57	MG	1E	305	1/1	0.93	0.18	47,47,47,47	0
57	MG	1E	306	1/1	0.93	0.09	42,42,42,42	0
57	MG	2A	3332	1/1	0.93	0.06	35,35,35,35	0
57	MG	2A	3023	1/1	0.93	0.16	54,54,54,54	0
57	MG	2A	3028	1/1	0.93	0.09	59,59,59,59	0
57	MG	2A	3032	1/1	0.93	0.09	50,50,50,50	0
57	MG	1A	3496	1/1	0.93	0.10	53,53,53,53	0
57	MG	2B	202	1/1	0.93	0.20	62,62,62,62	0
57	MG	2A	3342	1/1	0.93	0.12	45,45,45,45	0
57	MG	1E	309	1/1	0.93	0.08	25,25,25,25	0
57	MG	2A	3348	1/1	0.93	0.14	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	3358	1/1	0.93	0.07	31,31,31,31	0
57	MG	1A	3626	1/1	0.93	0.13	49,49,49,49	0
57	MG	2A	3041	1/1	0.93	0.14	44,44,44,44	0
57	MG	2A	3366	1/1	0.93	0.09	40,40,40,40	0
57	MG	2A	3367	1/1	0.93	0.12	63,63,63,63	0
57	MG	1A	3559	1/1	0.93	0.17	61,61,61,61	0
57	MG	1F	312	1/1	0.93	0.15	52,52,52,52	0
57	MG	1A	3391	1/1	0.93	0.08	43,43,43,43	0
57	MG	1A	3636	1/1	0.93	0.10	43,43,43,43	0
57	MG	1G	201	1/1	0.93	0.10	62,62,62,62	0
57	MG	1A	3563	1/1	0.93	0.07	51,51,51,51	0
57	MG	2A	3054	1/1	0.93	0.10	60,60,60,60	0
57	MG	2A	3388	1/1	0.93	0.16	63,63,63,63	0
57	MG	2F	303	1/1	0.93	0.16	57,57,57,57	0
57	MG	2A	3394	1/1	0.93	0.07	58,58,58,58	0
57	MG	1A	3902	1/1	0.93	0.10	20,20,20,20	0
57	MG	1A	3639	1/1	0.93	0.24	30,30,30,30	0
57	MG	1A	3503	1/1	0.93	0.08	37,37,37,37	0
57	MG	1A	3505	1/1	0.93	0.06	42,42,42,42	0
57	MG	1A	3921	1/1	0.93	0.10	50,50,50,50	0
57	MG	1A	3644	1/1	0.93	0.15	60,60,60,60	0
57	MG	2A	3406	1/1	0.93	0.10	63,63,63,63	0
57	MG	1A	3930	1/1	0.93	0.09	55,55,55,55	0
57	MG	2A	3078	1/1	0.93	0.14	53,53,53,53	0
57	MG	1T	201	1/1	0.93	0.08	42,42,42,42	0
57	MG	1A	3767	1/1	0.93	0.07	31,31,31,31	0
57	MG	1A	3014	1/1	0.93	0.09	39,39,39,39	0
57	MG	1a	3157	1/1	0.93	0.12	58,58,58,58	0
57	MG	2A	3429	1/1	0.93	0.08	44,44,44,44	0
57	MG	1A	3317	1/1	0.93	0.15	37,37,37,37	0
57	MG	27	101	1/1	0.93	0.20	49,49,49,49	0
57	MG	2A	3431	1/1	0.93	0.08	48,48,48,48	0
57	MG	2A	3432	1/1	0.93	0.14	56,56,56,56	0
57	MG	1A	3938	1/1	0.93	0.09	52,52,52,52	0
57	MG	1A	3652	1/1	0.93	0.08	35,35,35,35	0
57	MG	2a	3007	1/1	0.93	0.21	60,60,60,60	0
57	MG	1A	3395	1/1	0.93	0.07	18,18,18,18	0
57	MG	2A	3439	1/1	0.93	0.14	52,52,52,52	0
57	MG	2A	3098	1/1	0.93	0.12	54,54,54,54	0
57	MG	1A	3782	1/1	0.93	0.11	44,44,44,44	0
57	MG	1A	3039	1/1	0.93	0.10	38,38,38,38	0
57	MG	2A	3448	1/1	0.93	0.16	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	3090	1/1	0.93	0.13	43,43,43,43	0
57	MG	2a	3018	1/1	0.93	0.21	54,54,54,54	0
57	MG	1a	3173	1/1	0.93	0.13	68,68,68,68	0
57	MG	11	102	1/1	0.93	0.18	48,48,48,48	0
57	MG	2A	3459	1/1	0.93	0.07	47,47,47,47	0
57	MG	2A	3461	1/1	0.93	0.10	61,61,61,61	0
57	MG	1A	3667	1/1	0.93	0.07	46,46,46,46	0
57	MG	1A	3370	1/1	0.93	0.07	23,23,23,23	0
57	MG	2A	3464	1/1	0.93	0.12	58,58,58,58	0
57	MG	2A	3117	1/1	0.93	0.17	58,58,58,58	0
57	MG	18	103	1/1	0.93	0.08	50,50,50,50	0
57	MG	1A	3674	1/1	0.93	0.14	31,31,31,31	0
57	MG	1A	3957	1/1	0.93	0.11	49,49,49,49	0
57	MG	2A	3473	1/1	0.93	0.06	52,52,52,52	0
57	MG	2a	3036	1/1	0.93	0.07	66,66,66,66	0
57	MG	1A	3800	1/1	0.93	0.12	48,48,48,48	0
57	MG	2A	3129	1/1	0.93	0.11	46,46,46,46	0
57	MG	1A	3402	1/1	0.93	0.10	14,14,14,14	0
57	MG	1A	3805	1/1	0.93	0.08	34,34,34,34	0
57	MG	2A	3479	1/1	0.93	0.09	45,45,45,45	0
57	MG	1A	3406	1/1	0.93	0.17	57,57,57,57	0
57	MG	1A	3809	1/1	0.93	0.12	24,24,24,24	0
57	MG	1A	3589	1/1	0.93	0.09	43,43,43,43	0
57	MG	1a	3192	1/1	0.93	0.09	63,63,63,63	0
57	MG	2a	3048	1/1	0.93	0.27	63,63,63,63	0
57	MG	1A	3228	1/1	0.93	0.09	38,38,38,38	0
57	MG	2a	3053	1/1	0.93	0.10	56,56,56,56	0
57	MG	1a	3194	1/1	0.93	0.10	51,51,51,51	0
57	MG	2A	3495	1/1	0.93	0.16	57,57,57,57	0
57	MG	1A	3591	1/1	0.93	0.11	56,56,56,56	0
57	MG	2A	3500	1/1	0.93	0.09	42,42,42,42	0
57	MG	1a	3197	1/1	0.93	0.10	64,64,64,64	0
57	MG	1a	3198	1/1	0.93	0.07	46,46,46,46	0
57	MG	2A	3154	1/1	0.93	0.07	47,47,47,47	0
57	MG	2A	3509	1/1	0.93	0.12	64,64,64,64	0
57	MG	2a	3065	1/1	0.93	0.12	76,76,76,76	0
57	MG	2A	3156	1/1	0.93	0.10	55,55,55,55	0
57	MG	2a	3067	1/1	0.93	0.21	69,69,69,69	0
57	MG	1A	3184	1/1	0.93	0.09	43,43,43,43	0
57	MG	1A	3685	1/1	0.93	0.09	28,28,28,28	0
57	MG	2a	3070	1/1	0.93	0.07	51,51,51,51	0
57	MG	1A	3296	1/1	0.93	0.18	46,46,46,46	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3981	1/1	0.93	0.07	66,66,66,66	0
57	MG	1a	3039	1/1	0.93	0.34	64,64,64,64	0
57	MG	1A	3827	1/1	0.93	0.08	45,45,45,45	0
57	MG	1a	3206	1/1	0.93	0.12	56,56,56,56	0
57	MG	2a	3079	1/1	0.93	0.11	59,59,59,59	0
57	MG	1A	3986	1/1	0.93	0.26	46,46,46,46	0
57	MG	2A	3176	1/1	0.93	0.09	57,57,57,57	0
57	MG	1A	3830	1/1	0.93	0.07	45,45,45,45	0
57	MG	2A	3529	1/1	0.93	0.20	62,62,62,62	0
57	MG	2A	3179	1/1	0.93	0.12	52,52,52,52	0
57	MG	2a	3088	1/1	0.93	0.12	67,67,67,67	0
57	MG	1A	3691	1/1	0.93	0.13	53,53,53,53	0
57	MG	2a	3093	1/1	0.93	0.22	65,65,65,65	0
57	MG	1A	3834	1/1	0.93	0.10	26,26,26,26	0
57	MG	2a	3095	1/1	0.93	0.13	53,53,53,53	0
57	MG	1a	3051	1/1	0.93	0.19	56,56,56,56	0
57	MG	1A	3599	1/1	0.93	0.12	48,48,48,48	0
57	MG	2A	3536	1/1	0.93	0.15	69,69,69,69	0
57	MG	1A	3331	1/1	0.93	0.07	21,21,21,21	0
57	MG	2A	3190	1/1	0.93	0.15	58,58,58,58	0
57	MG	1a	3056	1/1	0.93	0.12	52,52,52,52	0
57	MG	2A	3544	1/1	0.93	0.08	29,29,29,29	0
57	MG	1a	3221	1/1	0.93	0.20	62,62,62,62	0
57	MG	1a	3223	1/1	0.93	0.13	61,61,61,61	0
57	MG	2a	3123	1/1	0.93	0.08	55,55,55,55	0
57	MG	1A	3992	1/1	0.93	0.23	37,37,37,37	0
57	MG	1a	3226	1/1	0.93	0.08	60,60,60,60	0
57	MG	1A	3188	1/1	0.93	0.18	42,42,42,42	0
57	MG	2A	3199	1/1	0.93	0.12	51,51,51,51	0
57	MG	2A	3202	1/1	0.93	0.11	46,46,46,46	0
57	MG	2A	3204	1/1	0.93	0.18	64,64,64,64	0
57	MG	1A	3536	1/1	0.93	0.12	50,50,50,50	0
57	MG	2A	3210	1/1	0.93	0.21	59,59,59,59	0
57	MG	1A	3996	1/1	0.93	0.12	56,56,56,56	0
57	MG	1A	3997	1/1	0.93	0.19	55,55,55,55	0
57	MG	2a	3144	1/1	0.93	0.24	68,68,68,68	0
57	MG	1A	3703	1/1	0.93	0.17	40,40,40,40	0
57	MG	1A	3382	1/1	0.93	0.08	43,43,43,43	0
57	MG	2A	3572	1/1	0.93	0.08	47,47,47,47	0
57	MG	1a	3237	1/1	0.93	0.17	48,48,48,48	0
57	MG	1A	3437	1/1	0.93	0.11	44,44,44,44	0
57	MG	2A	3579	1/1	0.93	0.12	55,55,55,55	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3580	1/1	0.93	0.06	43,43,43,43	0
57	MG	2A	3223	1/1	0.93	0.18	57,57,57,57	0
57	MG	1a	3071	1/1	0.93	0.10	62,62,62,62	0
57	MG	2a	3161	1/1	0.93	0.07	65,65,65,65	0
57	MG	2a	3163	1/1	0.93	0.11	64,64,64,64	0
57	MG	2A	3583	1/1	0.93	0.07	52,52,52,52	0
57	MG	1a	3073	1/1	0.93	0.19	59,59,59,59	0
57	MG	1A	3709	1/1	0.93	0.08	53,53,53,53	0
57	MG	2A	3591	1/1	0.93	0.09	50,50,50,50	0
57	MG	2a	3169	1/1	0.93	0.07	56,56,56,56	0
57	MG	1A	3095	1/1	0.93	0.09	45,45,45,45	0
57	MG	2A	3234	1/1	0.93	0.15	46,46,46,46	0
57	MG	2A	3236	1/1	0.93	0.09	55,55,55,55	0
57	MG	1A	3713	1/1	0.93	0.09	36,36,36,36	0
57	MG	2A	3601	1/1	0.93	0.10	62,62,62,62	0
57	MG	1B	201	1/1	0.93	0.08	43,43,43,43	0
57	MG	2A	3606	1/1	0.93	0.08	56,56,56,56	0
57	MG	2A	3245	1/1	0.93	0.10	55,55,55,55	0
57	MG	1a	3264	1/1	0.93	0.11	77,77,77,77	0
57	MG	1A	3216	1/1	0.93	0.16	48,48,48,48	0
57	MG	1B	205	1/1	0.93	0.31	39,39,39,39	0
57	MG	1B	206	1/1	0.93	0.08	47,47,47,47	0
57	MG	2p	101	1/1	0.93	0.25	60,60,60,60	0
58	MPD	1A	4023	8/8	0.93	0.16	41,51,57,57	0
57	MG	1a	3269	1/1	0.93	0.15	75,75,75,75	0
58	MPD	1a	3276	8/8	0.93	0.11	54,57,61,63	0
57	MG	2A	3623	1/1	0.93	0.07	71,71,71,71	0
57	MG	2A	3626	1/1	0.93	0.14	61,61,61,61	0
57	MG	2A	3630	1/1	0.93	0.08	55,55,55,55	0
57	MG	1a	3271	1/1	0.93	0.11	52,52,52,52	0
57	MG	1A	3858	1/1	0.93	0.09	40,40,40,40	0
57	MG	2A	3270	1/1	0.94	0.21	56,56,56,56	0
57	MG	2A	3271	1/1	0.94	0.10	50,50,50,50	0
57	MG	1A	3361	1/1	0.94	0.18	60,60,60,60	0
57	MG	1B	223	1/1	0.94	0.07	44,44,44,44	0
57	MG	1B	226	1/1	0.94	0.06	32,32,32,32	0
57	MG	2A	3280	1/1	0.94	0.23	56,56,56,56	0
57	MG	1A	3283	1/1	0.94	0.16	51,51,51,51	0
57	MG	2A	3645	1/1	0.94	0.08	70,70,70,70	0
57	MG	1a	3098	1/1	0.94	0.17	57,57,57,57	0
57	MG	2A	3647	1/1	0.94	0.07	46,46,46,46	0
57	MG	2A	3650	1/1	0.94	0.08	30,30,30,30	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3743	1/1	0.94	0.07	41,41,41,41	0
57	MG	1A	3290	1/1	0.94	0.15	67,67,67,67	0
57	MG	1a	3101	1/1	0.94	0.24	57,57,57,57	0
57	MG	2A	3662	1/1	0.94	0.07	61,61,61,61	0
57	MG	2A	3288	1/1	0.94	0.15	57,57,57,57	0
57	MG	2A	3665	1/1	0.94	0.07	37,37,37,37	0
57	MG	2A	3290	1/1	0.94	0.16	49,49,49,49	0
57	MG	1A	3022	1/1	0.94	0.08	47,47,47,47	0
57	MG	1A	3293	1/1	0.94	0.12	62,62,62,62	0
57	MG	2A	3293	1/1	0.94	0.20	57,57,57,57	0
57	MG	2A	3296	1/1	0.94	0.07	40,40,40,40	0
57	MG	2A	3672	1/1	0.94	0.07	33,33,33,33	0
57	MG	2A	3297	1/1	0.94	0.11	52,52,52,52	0
57	MG	1a	3105	1/1	0.94	0.11	46,46,46,46	0
57	MG	1t	201	1/1	0.94	0.15	65,65,65,65	0
57	MG	1A	3629	1/1	0.94	0.14	42,42,42,42	0
57	MG	1A	3633	1/1	0.94	0.14	57,57,57,57	0
57	MG	2A	3687	1/1	0.94	0.08	43,43,43,43	0
57	MG	1A	3194	1/1	0.94	0.08	45,45,45,45	0
57	MG	1y	204	1/1	0.94	0.08	63,63,63,63	0
57	MG	2A	3691	1/1	0.94	0.14	51,51,51,51	0
57	MG	1A	3755	1/1	0.94	0.13	48,48,48,48	0
57	MG	2A	3308	1/1	0.94	0.21	51,51,51,51	0
57	MG	1A	3195	1/1	0.94	0.15	51,51,51,51	0
57	MG	1A	3759	1/1	0.94	0.07	35,35,35,35	0
57	MG	2A	3311	1/1	0.94	0.07	43,43,43,43	0
57	MG	1a	3112	1/1	0.94	0.11	72,72,72,72	0
57	MG	1A	3224	1/1	0.94	0.07	41,41,41,41	0
57	MG	2A	3026	1/1	0.94	0.11	55,55,55,55	0
57	MG	2A	3316	1/1	0.94	0.08	40,40,40,40	0
57	MG	2A	3708	1/1	0.94	0.12	59,59,59,59	0
57	MG	1A	3023	1/1	0.94	0.09	28,28,28,28	0
57	MG	2A	3711	1/1	0.94	0.09	49,49,49,49	0
57	MG	2A	3713	1/1	0.94	0.09	24,24,24,24	0
57	MG	2A	3714	1/1	0.94	0.15	49,49,49,49	0
57	MG	2A	3030	1/1	0.94	0.08	26,26,26,26	0
57	MG	1A	3918	1/1	0.94	0.10	34,34,34,34	0
57	MG	1A	3553	1/1	0.94	0.16	59,59,59,59	0
57	MG	2A	3036	1/1	0.94	0.10	46,46,46,46	0
57	MG	1E	312	1/1	0.94	0.10	59,59,59,59	0
57	MG	2A	3330	1/1	0.94	0.08	38,38,38,38	0
57	MG	1A	3312	1/1	0.94	0.18	26,26,26,26	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1a	3124	1/1	0.94	0.10	73,73,73,73	0
57	MG	1A	3926	1/1	0.94	0.06	51,51,51,51	0
57	MG	1A	3231	1/1	0.94	0.13	32,32,32,32	0
57	MG	1A	3770	1/1	0.94	0.06	26,26,26,26	0
57	MG	1a	3132	1/1	0.94	0.07	57,57,57,57	0
57	MG	2A	3343	1/1	0.94	0.10	56,56,56,56	0
57	MG	2A	3345	1/1	0.94	0.10	63,63,63,63	0
57	MG	1A	3772	1/1	0.94	0.05	37,37,37,37	0
57	MG	1A	3773	1/1	0.94	0.08	26,26,26,26	0
57	MG	2A	3351	1/1	0.94	0.10	38,38,38,38	0
57	MG	2A	3354	1/1	0.94	0.08	38,38,38,38	0
57	MG	2A	3355	1/1	0.94	0.13	65,65,65,65	0
57	MG	2B	204	1/1	0.94	0.20	56,56,56,56	0
57	MG	1a	3137	1/1	0.94	0.09	51,51,51,51	0
57	MG	1A	3775	1/1	0.94	0.05	32,32,32,32	0
57	MG	1A	3234	1/1	0.94	0.12	34,34,34,34	0
57	MG	2B	209	1/1	0.94	0.14	61,61,61,61	0
57	MG	2A	3059	1/1	0.94	0.28	46,46,46,46	0
57	MG	1A	3941	1/1	0.94	0.07	47,47,47,47	0
57	MG	1O	201	1/1	0.94	0.17	48,48,48,48	0
57	MG	2A	3369	1/1	0.94	0.06	36,36,36,36	0
57	MG	2B	214	1/1	0.94	0.09	52,52,52,52	0
57	MG	2A	3372	1/1	0.94	0.08	35,35,35,35	0
57	MG	2B	217	1/1	0.94	0.10	62,62,62,62	0
57	MG	1a	3143	1/1	0.94	0.18	65,65,65,65	0
57	MG	1A	3647	1/1	0.94	0.13	36,36,36,36	0
57	MG	2D	305	1/1	0.94	0.12	44,44,44,44	0
57	MG	2A	3072	1/1	0.94	0.15	49,49,49,49	0
57	MG	1A	3943	1/1	0.94	0.09	23,23,23,23	0
57	MG	2D	309	1/1	0.94	0.10	46,46,46,46	0
57	MG	2D	310	1/1	0.94	0.12	47,47,47,47	0
57	MG	1R	202	1/1	0.94	0.12	35,35,35,35	0
57	MG	2E	303	1/1	0.94	0.07	49,49,49,49	0
57	MG	2A	3382	1/1	0.94	0.09	58,58,58,58	0
57	MG	2A	3079	1/1	0.94	0.07	43,43,43,43	0
57	MG	2F	302	1/1	0.94	0.12	51,51,51,51	0
57	MG	2A	3080	1/1	0.94	0.17	56,56,56,56	0
57	MG	2A	3082	1/1	0.94	0.09	53,53,53,53	0
57	MG	1A	3387	1/1	0.94	0.09	24,24,24,24	0
57	MG	1A	3108	1/1	0.94	0.11	51,51,51,51	0
57	MG	1A	3786	1/1	0.94	0.11	45,45,45,45	0
57	MG	2A	3399	1/1	0.94	0.18	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	2P	201	1/1	0.94	0.10	60,60,60,60	0
57	MG	2A	3086	1/1	0.94	0.35	42,42,42,42	0
57	MG	2A	3087	1/1	0.94	0.13	45,45,45,45	0
57	MG	2Q	204	1/1	0.94	0.08	67,67,67,67	0
57	MG	1A	3200	1/1	0.94	0.09	38,38,38,38	0
57	MG	1U	201	1/1	0.94	0.22	33,33,33,33	0
57	MG	1V	204	1/1	0.94	0.08	43,43,43,43	0
57	MG	1A	3473	1/1	0.94	0.12	47,47,47,47	0
57	MG	1A	3663	1/1	0.94	0.08	24,24,24,24	0
57	MG	1W	203	1/1	0.94	0.10	58,58,58,58	0
57	MG	1a	3162	1/1	0.94	0.25	57,57,57,57	0
57	MG	1a	3163	1/1	0.94	0.10	57,57,57,57	0
57	MG	2A	3420	1/1	0.94	0.08	30,30,30,30	0
57	MG	2A	3422	1/1	0.94	0.08	31,31,31,31	0
57	MG	2A	3426	1/1	0.94	0.09	65,65,65,65	0
57	MG	2A	3428	1/1	0.94	0.07	42,42,42,42	0
57	MG	1A	3795	1/1	0.94	0.10	47,47,47,47	0
57	MG	2A	3108	1/1	0.94	0.09	51,51,51,51	0
57	MG	2A	3110	1/1	0.94	0.12	56,56,56,56	0
57	MG	1A	3567	1/1	0.94	0.12	55,55,55,55	0
57	MG	2A	3433	1/1	0.94	0.13	49,49,49,49	0
57	MG	10	101	1/1	0.94	0.20	45,45,45,45	0
57	MG	1A	3962	1/1	0.94	0.10	46,46,46,46	0
57	MG	1A	3260	1/1	0.94	0.13	39,39,39,39	0
57	MG	10	108	1/1	0.94	0.10	47,47,47,47	0
57	MG	2A	3440	1/1	0.94	0.10	75,75,75,75	0
57	MG	1A	3966	1/1	0.94	0.09	52,52,52,52	0
57	MG	1A	3202	1/1	0.94	0.05	40,40,40,40	0
57	MG	2A	3444	1/1	0.94	0.09	41,41,41,41	0
57	MG	11	103	1/1	0.94	0.08	36,36,36,36	0
57	MG	2A	3122	1/1	0.94	0.07	48,48,48,48	0
57	MG	1A	3329	1/1	0.94	0.12	29,29,29,29	0
57	MG	1A	3082	1/1	0.94	0.13	47,47,47,47	0
57	MG	2A	3456	1/1	0.94	0.08	39,39,39,39	0
57	MG	1a	3183	1/1	0.94	0.08	66,66,66,66	0
57	MG	2a	3030	1/1	0.94	0.10	57,57,57,57	0
57	MG	2A	3458	1/1	0.94	0.08	49,49,49,49	0
57	MG	1A	3676	1/1	0.94	0.10	48,48,48,48	0
57	MG	2A	3460	1/1	0.94	0.11	63,63,63,63	0
57	MG	19	102	1/1	0.94	0.09	60,60,60,60	0
57	MG	2A	3133	1/1	0.94	0.15	64,64,64,64	0
57	MG	2A	3134	1/1	0.94	0.21	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	2A	3135	1/1	0.94	0.08	27,27,27,27	0
57	MG	1a	3187	1/1	0.94	0.11	60,60,60,60	0
57	MG	1A	3204	1/1	0.94	0.09	50,50,50,50	0
57	MG	1A	3333	1/1	0.94	0.23	48,48,48,48	0
57	MG	1A	3339	1/1	0.94	0.08	36,36,36,36	0
57	MG	1A	3681	1/1	0.94	0.07	58,58,58,58	0
57	MG	1A	3818	1/1	0.94	0.12	37,37,37,37	0
57	MG	1A	3491	1/1	0.94	0.07	59,59,59,59	0
57	MG	1a	3015	1/1	0.94	0.17	64,64,64,64	0
57	MG	1A	3126	1/1	0.94	0.11	62,62,62,62	0
57	MG	2a	3050	1/1	0.94	0.17	54,54,54,54	0
57	MG	2a	3052	1/1	0.94	0.11	59,59,59,59	0
57	MG	1A	3587	1/1	0.94	0.07	48,48,48,48	0
57	MG	1A	3405	1/1	0.94	0.07	34,34,34,34	0
57	MG	2A	3483	1/1	0.94	0.07	64,64,64,64	0
57	MG	1A	3342	1/1	0.94	0.07	48,48,48,48	0
57	MG	2A	3485	1/1	0.94	0.07	61,61,61,61	0
57	MG	2A	3487	1/1	0.94	0.11	62,62,62,62	0
57	MG	2A	3488	1/1	0.94	0.07	64,64,64,64	0
57	MG	1A	3415	1/1	0.94	0.06	35,35,35,35	0
57	MG	2a	3063	1/1	0.94	0.25	66,66,66,66	0
57	MG	2A	3164	1/1	0.94	0.09	48,48,48,48	0
57	MG	1a	3027	1/1	0.94	0.16	56,56,56,56	0
57	MG	1a	3029	1/1	0.94	0.09	49,49,49,49	0
57	MG	2A	3168	1/1	0.94	0.08	48,48,48,48	0
57	MG	2A	3498	1/1	0.94	0.08	64,64,64,64	0
57	MG	1A	3346	1/1	0.94	0.07	41,41,41,41	0
57	MG	1A	3021	1/1	0.94	0.15	39,39,39,39	0
57	MG	1a	3036	1/1	0.94	0.29	52,52,52,52	0
57	MG	1a	3207	1/1	0.94	0.08	69,69,69,69	0
57	MG	2A	3507	1/1	0.94	0.09	56,56,56,56	0
57	MG	1a	3208	1/1	0.94	0.14	58,58,58,58	0
57	MG	1A	3420	1/1	0.94	0.10	53,53,53,53	0
57	MG	2A	3180	1/1	0.94	0.16	56,56,56,56	0
57	MG	1A	3845	1/1	0.94	0.07	42,42,42,42	0
57	MG	1A	3846	1/1	0.94	0.13	31,31,31,31	0
57	MG	2A	3513	1/1	0.94	0.07	42,42,42,42	0
57	MG	2A	3514	1/1	0.94	0.07	62,62,62,62	0
57	MG	2A	3517	1/1	0.94	0.09	49,49,49,49	0
57	MG	2a	3087	1/1	0.94	0.13	54,54,54,54	0
57	MG	2A	3185	1/1	0.94	0.10	55,55,55,55	0
57	MG	2A	3519	1/1	0.94	0.09	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	1A	3699	1/1	0.94	0.17	45,45,45,45	0
57	MG	1A	3348	1/1	0.94	0.16	52,52,52,52	0
57	MG	1a	3217	1/1	0.94	0.09	66,66,66,66	0
57	MG	2a	3098	1/1	0.94	0.31	66,66,66,66	0
57	MG	2a	3099	1/1	0.94	0.27	45,45,45,45	0
57	MG	1A	3999	1/1	0.94	0.13	30,30,30,30	0
57	MG	1a	3047	1/1	0.94	0.15	56,56,56,56	0
57	MG	2a	3103	1/1	0.94	0.22	49,49,49,49	0
57	MG	2a	3108	1/1	0.94	0.20	51,51,51,51	0
57	MG	2a	3109	1/1	0.94	0.15	62,62,62,62	0
57	MG	2a	3111	1/1	0.94	0.28	64,64,64,64	0
57	MG	1A	4000	1/1	0.94	0.11	15,15,15,15	0
57	MG	1A	4001	1/1	0.94	0.11	47,47,47,47	0
57	MG	1A	3063	1/1	0.94	0.07	35,35,35,35	0
57	MG	1A	3706	1/1	0.94	0.10	57,57,57,57	0
57	MG	2a	3116	1/1	0.94	0.23	49,49,49,49	0
57	MG	1A	3515	1/1	0.94	0.09	21,21,21,21	0
57	MG	2A	3532	1/1	0.94	0.08	62,62,62,62	0
57	MG	1a	3057	1/1	0.94	0.08	58,58,58,58	0
57	MG	2A	3200	1/1	0.94	0.07	41,41,41,41	0
57	MG	1A	4010	1/1	0.94	0.09	36,36,36,36	0
57	MG	2a	3126	1/1	0.94	0.17	61,61,61,61	0
57	MG	1A	4012	1/1	0.94	0.17	33,33,33,33	0
57	MG	2a	3128	1/1	0.94	0.15	71,71,71,71	0
57	MG	2A	3205	1/1	0.94	0.07	50,50,50,50	0
57	MG	2A	3207	1/1	0.94	0.15	52,52,52,52	0
57	MG	2A	3208	1/1	0.94	0.19	54,54,54,54	0
57	MG	1A	3426	1/1	0.94	0.09	37,37,37,37	0
57	MG	2a	3133	1/1	0.94	0.12	66,66,66,66	0
57	MG	1A	3276	1/1	0.94	0.10	43,43,43,43	0
57	MG	2a	3136	1/1	0.94	0.09	71,71,71,71	0
57	MG	1a	3065	1/1	0.94	0.08	58,58,58,58	0
57	MG	1a	3066	1/1	0.94	0.10	63,63,63,63	0
57	MG	2a	3140	1/1	0.94	0.10	69,69,69,69	0
57	MG	1A	3435	1/1	0.94	0.09	44,44,44,44	0
57	MG	2A	3217	1/1	0.94	0.31	65,65,65,65	0
57	MG	1A	3606	1/1	0.94	0.08	27,27,27,27	0
57	MG	1B	202	1/1	0.94	0.13	52,52,52,52	0
57	MG	2A	3558	1/1	0.94	0.08	40,40,40,40	0
57	MG	1a	3245	1/1	0.94	0.08	46,46,46,46	0
57	MG	2A	3222	1/1	0.94	0.07	43,43,43,43	0
57	MG	1A	3520	1/1	0.94	0.27	47,47,47,47	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1a	3252	1/1	0.94	0.17	54,54,54,54	0
57	MG	1A	3861	1/1	0.94	0.10	49,49,49,49	0
57	MG	1A	3862	1/1	0.94	0.06	54,54,54,54	0
57	MG	1a	3260	1/1	0.94	0.10	55,55,55,55	0
57	MG	1A	3610	1/1	0.94	0.16	32,32,32,32	0
57	MG	1a	3076	1/1	0.94	0.15	47,47,47,47	0
57	MG	1B	208	1/1	0.94	0.18	50,50,50,50	0
57	MG	1B	209	1/1	0.94	0.07	38,38,38,38	0
57	MG	2A	3239	1/1	0.94	0.17	52,52,52,52	0
57	MG	2A	3240	1/1	0.94	0.34	59,59,59,59	0
57	MG	2A	3241	1/1	0.94	0.07	54,54,54,54	0
57	MG	2a	3172	1/1	0.94	0.12	55,55,55,55	0
57	MG	1A	3521	1/1	0.94	0.09	46,46,46,46	0
57	MG	1A	3616	1/1	0.94	0.24	38,38,38,38	0
57	MG	2A	3585	1/1	0.94	0.18	55,55,55,55	0
57	MG	1B	216	1/1	0.94	0.16	43,43,43,43	0
57	MG	1a	3084	1/1	0.94	0.07	50,50,50,50	0
57	MG	2a	3185	1/1	0.94	0.10	77,77,77,77	0
57	MG	2a	3186	1/1	0.94	0.10	67,67,67,67	0
57	MG	2a	3188	1/1	0.94	0.09	58,58,58,58	0
57	MG	2A	3249	1/1	0.94	0.23	49,49,49,49	0
57	MG	1a	3085	1/1	0.94	0.27	64,64,64,64	0
57	MG	1A	3617	1/1	0.94	0.16	34,34,34,34	0
57	MG	1d	302	1/1	0.94	0.19	63,63,63,63	0
57	MG	1A	3618	1/1	0.94	0.07	30,30,30,30	0
57	MG	1A	3734	1/1	0.94	0.08	41,41,41,41	0
57	MG	2A	3603	1/1	0.94	0.06	51,51,51,51	0
57	MG	2A	3604	1/1	0.94	0.08	41,41,41,41	0
57	MG	2A	3258	1/1	0.94	0.10	48,48,48,48	0
57	MG	1d	305	1/1	0.94	0.12	55,55,55,55	0
57	MG	1A	3278	1/1	0.94	0.07	47,47,47,47	0
57	MG	1a	3092	1/1	0.94	0.09	52,52,52,52	0
57	MG	2A	3616	1/1	0.94	0.10	49,49,49,49	0
57	MG	2A	3617	1/1	0.94	0.07	75,75,75,75	0
57	MG	1e	203	1/1	0.94	0.24	59,59,59,59	0
60	ZN	24	501	1/1	0.94	0.14	119,119,119,119	0
57	MG	1A	3971	1/1	0.95	0.04	27,27,27,27	0
57	MG	2A	3344	1/1	0.95	0.12	53,53,53,53	0
57	MG	11	104	1/1	0.95	0.17	48,48,48,48	0
57	MG	13	102	1/1	0.95	0.08	51,51,51,51	0
57	MG	15	3107	1/1	0.95	0.17	25,25,25,25	0
57	MG	2A	3350	1/1	0.95	0.07	46,46,46,46	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3478	1/1	0.95	0.08	29,29,29,29	0
57	MG	2A	3352	1/1	0.95	0.09	44,44,44,44	0
57	MG	1A	3806	1/1	0.95	0.06	48,48,48,48	0
57	MG	1a	3169	1/1	0.95	0.08	66,66,66,66	0
57	MG	2A	3356	1/1	0.95	0.06	40,40,40,40	0
57	MG	2A	3697	1/1	0.95	0.33	39,39,39,39	0
57	MG	1a	3171	1/1	0.95	0.14	61,61,61,61	0
57	MG	18	101	1/1	0.95	0.07	43,43,43,43	0
57	MG	1A	3161	1/1	0.95	0.15	35,35,35,35	0
57	MG	2A	3364	1/1	0.95	0.08	44,44,44,44	0
57	MG	1A	3481	1/1	0.95	0.07	25,25,25,25	0
57	MG	2A	3705	1/1	0.95	0.10	56,56,56,56	0
57	MG	2A	3706	1/1	0.95	0.08	39,39,39,39	0
57	MG	2A	3099	1/1	0.95	0.05	48,48,48,48	0
57	MG	2A	3101	1/1	0.95	0.09	43,43,43,43	0
57	MG	1a	3001	1/1	0.95	0.14	68,68,68,68	0
57	MG	1A	3581	1/1	0.95	0.11	47,47,47,47	0
57	MG	2A	3712	1/1	0.95	0.06	47,47,47,47	0
57	MG	1A	3813	1/1	0.95	0.06	37,37,37,37	0
57	MG	2A	3106	1/1	0.95	0.07	39,39,39,39	0
57	MG	1a	3180	1/1	0.95	0.07	49,49,49,49	0
57	MG	1A	3482	1/1	0.95	0.07	45,45,45,45	0
57	MG	2A	3378	1/1	0.95	0.10	66,66,66,66	0
57	MG	1a	3182	1/1	0.95	0.12	59,59,59,59	0
57	MG	1A	3265	1/1	0.95	0.09	46,46,46,46	0
57	MG	2A	3383	1/1	0.95	0.11	53,53,53,53	0
57	MG	2A	3384	1/1	0.95	0.14	57,57,57,57	0
57	MG	1A	3584	1/1	0.95	0.10	39,39,39,39	0
57	MG	2A	3115	1/1	0.95	0.24	54,54,54,54	0
57	MG	2A	3393	1/1	0.95	0.08	59,59,59,59	0
57	MG	1A	3485	1/1	0.95	0.08	29,29,29,29	0
57	MG	1A	3267	1/1	0.95	0.09	28,28,28,28	0
57	MG	2A	3396	1/1	0.95	0.08	59,59,59,59	0
57	MG	1a	3016	1/1	0.95	0.10	60,60,60,60	0
57	MG	1a	3017	1/1	0.95	0.14	57,57,57,57	0
57	MG	1A	3688	1/1	0.95	0.09	53,53,53,53	0
57	MG	1A	3689	1/1	0.95	0.11	56,56,56,56	0
57	MG	2A	3123	1/1	0.95	0.14	38,38,38,38	0
57	MG	2A	3125	1/1	0.95	0.13	48,48,48,48	0
57	MG	2A	3405	1/1	0.95	0.07	49,49,49,49	0
57	MG	1A	3828	1/1	0.95	0.09	34,34,34,34	0
57	MG	1A	3829	1/1	0.95	0.08	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	1a	3024	1/1	0.95	0.07	50,50,50,50	0
57	MG	1A	3690	1/1	0.95	0.07	37,37,37,37	0
57	MG	1A	3993	1/1	0.95	0.11	24,24,24,24	0
57	MG	1a	3028	1/1	0.95	0.14	59,59,59,59	0
57	MG	1A	3163	1/1	0.95	0.13	50,50,50,50	0
57	MG	2A	3414	1/1	0.95	0.09	47,47,47,47	0
57	MG	2A	3416	1/1	0.95	0.19	50,50,50,50	0
57	MG	1a	3030	1/1	0.95	0.08	46,46,46,46	0
57	MG	2A	3421	1/1	0.95	0.12	49,49,49,49	0
57	MG	1A	3332	1/1	0.95	0.12	20,20,20,20	0
57	MG	2D	301	1/1	0.95	0.47	46,46,46,46	0
57	MG	1a	3034	1/1	0.95	0.12	48,48,48,48	0
57	MG	2A	3427	1/1	0.95	0.07	40,40,40,40	0
57	MG	1A	3206	1/1	0.95	0.22	41,41,41,41	0
57	MG	2A	3140	1/1	0.95	0.14	53,53,53,53	0
57	MG	2A	3141	1/1	0.95	0.09	60,60,60,60	0
57	MG	1A	3028	1/1	0.95	0.25	26,26,26,26	0
57	MG	1A	3594	1/1	0.95	0.09	50,50,50,50	0
57	MG	2A	3145	1/1	0.95	0.20	46,46,46,46	0
57	MG	2A	3434	1/1	0.95	0.09	50,50,50,50	0
57	MG	1A	3698	1/1	0.95	0.12	48,48,48,48	0
57	MG	1A	3500	1/1	0.95	0.10	47,47,47,47	0
57	MG	2A	3150	1/1	0.95	0.05	41,41,41,41	0
57	MG	2A	3152	1/1	0.95	0.17	46,46,46,46	0
57	MG	1A	3700	1/1	0.95	0.22	53,53,53,53	0
57	MG	2A	3441	1/1	0.95	0.07	43,43,43,43	0
57	MG	1A	3168	1/1	0.95	0.07	45,45,45,45	0
57	MG	1A	4005	1/1	0.95	0.09	47,47,47,47	0
57	MG	1A	3031	1/1	0.95	0.04	14,14,14,14	0
57	MG	2P	202	1/1	0.95	0.15	64,64,64,64	0
57	MG	2Q	201	1/1	0.95	0.08	56,56,56,56	0
57	MG	1A	3408	1/1	0.95	0.08	28,28,28,28	0
57	MG	1a	3216	1/1	0.95	0.08	74,74,74,74	0
57	MG	2A	3450	1/1	0.95	0.07	51,51,51,51	0
57	MG	2A	3162	1/1	0.95	0.20	41,41,41,41	0
57	MG	2A	3453	1/1	0.95	0.07	58,58,58,58	0
57	MG	1A	3345	1/1	0.95	0.10	48,48,48,48	0
57	MG	1A	3033	1/1	0.95	0.08	36,36,36,36	0
57	MG	1A	4016	1/1	0.95	0.08	47,47,47,47	0
57	MG	1a	3055	1/1	0.95	0.13	44,44,44,44	0
57	MG	2V	202	1/1	0.95	0.14	54,54,54,54	0
57	MG	2V	203	1/1	0.95	0.09	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	4017	1/1	0.95	0.10	58,58,58,58	0
57	MG	2W	203	1/1	0.95	0.11	61,61,61,61	0
57	MG	2Y	201	1/1	0.95	0.13	53,53,53,53	0
57	MG	20	101	1/1	0.95	0.11	54,54,54,54	0
57	MG	2A	3170	1/1	0.95	0.19	44,44,44,44	0
57	MG	2A	3171	1/1	0.95	0.08	42,42,42,42	0
57	MG	2A	3172	1/1	0.95	0.06	39,39,39,39	0
57	MG	1a	3222	1/1	0.95	0.09	49,49,49,49	0
57	MG	1A	3186	1/1	0.95	0.23	38,38,38,38	0
57	MG	1A	3112	1/1	0.95	0.14	32,32,32,32	0
57	MG	2A	3467	1/1	0.95	0.08	53,53,53,53	0
57	MG	1A	3071	1/1	0.95	0.11	40,40,40,40	0
57	MG	1a	3061	1/1	0.95	0.11	60,60,60,60	0
57	MG	1A	3607	1/1	0.95	0.23	30,30,30,30	0
57	MG	1a	3063	1/1	0.95	0.18	68,68,68,68	0
57	MG	2A	3182	1/1	0.95	0.09	64,64,64,64	0
57	MG	1A	3423	1/1	0.95	0.11	20,20,20,20	0
57	MG	1B	204	1/1	0.95	0.07	45,45,45,45	0
57	MG	2A	3186	1/1	0.95	0.09	49,49,49,49	0
57	MG	1A	3722	1/1	0.95	0.10	43,43,43,43	0
57	MG	2A	3480	1/1	0.95	0.09	48,48,48,48	0
57	MG	1A	3723	1/1	0.95	0.07	39,39,39,39	0
57	MG	2A	3482	1/1	0.95	0.09	45,45,45,45	0
57	MG	1A	3424	1/1	0.95	0.09	32,32,32,32	0
57	MG	1A	3354	1/1	0.95	0.12	55,55,55,55	0
57	MG	1A	3355	1/1	0.95	0.08	23,23,23,23	0
57	MG	2A	3486	1/1	0.95	0.06	64,64,64,64	0
57	MG	1a	3243	1/1	0.95	0.07	47,47,47,47	0
57	MG	1A	3075	1/1	0.95	0.20	40,40,40,40	0
57	MG	1a	3248	1/1	0.95	0.09	54,54,54,54	0
57	MG	1A	3874	1/1	0.95	0.09	46,46,46,46	0
57	MG	1B	212	1/1	0.95	0.06	48,48,48,48	0
57	MG	1a	3075	1/1	0.95	0.17	66,66,66,66	0
57	MG	1a	3254	1/1	0.95	0.07	49,49,49,49	0
57	MG	1B	215	1/1	0.95	0.10	56,56,56,56	0
57	MG	2A	3203	1/1	0.95	0.08	52,52,52,52	0
57	MG	1a	3258	1/1	0.95	0.10	68,68,68,68	0
57	MG	1A	3020	1/1	0.95	0.33	28,28,28,28	0
57	MG	1a	3262	1/1	0.95	0.08	57,57,57,57	0
57	MG	1A	3876	1/1	0.95	0.05	27,27,27,27	0
57	MG	1a	3080	1/1	0.95	0.14	65,65,65,65	0
57	MG	1A	3877	1/1	0.95	0.09	62,62,62,62	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2a	3046	1/1	0.95	0.34	62,62,62,62	0
57	MG	2A	3211	1/1	0.95	0.18	52,52,52,52	0
57	MG	1A	3359	1/1	0.95	0.09	32,32,32,32	0
57	MG	1A	3739	1/1	0.95	0.09	42,42,42,42	0
57	MG	1a	3268	1/1	0.95	0.09	62,62,62,62	0
57	MG	2a	3051	1/1	0.95	0.21	54,54,54,54	0
57	MG	1B	221	1/1	0.95	0.10	34,34,34,34	0
57	MG	2A	3515	1/1	0.95	0.07	59,59,59,59	0
57	MG	1A	3083	1/1	0.95	0.22	28,28,28,28	0
57	MG	1A	3882	1/1	0.95	0.06	27,27,27,27	0
57	MG	1B	225	1/1	0.95	0.12	54,54,54,54	0
57	MG	2A	3520	1/1	0.95	0.14	57,57,57,57	0
57	MG	1A	3300	1/1	0.95	0.08	31,31,31,31	0
57	MG	1d	301	1/1	0.95	0.24	67,67,67,67	0
57	MG	2a	3061	1/1	0.95	0.30	67,67,67,67	0
57	MG	2A	3224	1/1	0.95	0.10	42,42,42,42	0
57	MG	1A	3742	1/1	0.95	0.10	41,41,41,41	0
57	MG	1a	3090	1/1	0.95	0.13	47,47,47,47	0
57	MG	1A	3532	1/1	0.95	0.07	25,25,25,25	0
57	MG	1A	3304	1/1	0.95	0.10	33,33,33,33	0
57	MG	1A	3887	1/1	0.95	0.13	27,27,27,27	0
57	MG	2A	3235	1/1	0.95	0.10	58,58,58,58	0
57	MG	1D	308	1/1	0.95	0.16	35,35,35,35	0
57	MG	1A	3745	1/1	0.95	0.06	58,58,58,58	0
57	MG	1A	3240	1/1	0.95	0.16	32,32,32,32	0
57	MG	1A	3896	1/1	0.95	0.07	45,45,45,45	0
57	MG	1h	201	1/1	0.95	0.20	49,49,49,49	0
57	MG	2a	3074	1/1	0.95	0.24	50,50,50,50	0
57	MG	1A	3628	1/1	0.95	0.08	46,46,46,46	0
57	MG	2A	3243	1/1	0.95	0.34	47,47,47,47	0
57	MG	2A	3540	1/1	0.95	0.08	53,53,53,53	0
57	MG	1A	3900	1/1	0.95	0.08	48,48,48,48	0
57	MG	2A	3542	1/1	0.95	0.09	64,64,64,64	0
57	MG	1A	3535	1/1	0.95	0.10	49,49,49,49	0
57	MG	1A	3631	1/1	0.95	0.07	30,30,30,30	0
57	MG	1m	201	1/1	0.95	0.06	62,62,62,62	0
57	MG	1A	3905	1/1	0.95	0.06	27,27,27,27	0
57	MG	1A	3443	1/1	0.95	0.08	42,42,42,42	0
57	MG	2a	3089	1/1	0.95	0.16	60,60,60,60	0
57	MG	1A	3909	1/1	0.95	0.09	28,28,28,28	0
57	MG	1A	3754	1/1	0.95	0.14	56,56,56,56	0
57	MG	1A	3307	1/1	0.95	0.06	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	3556	1/1	0.95	0.06	53,53,53,53	0
57	MG	2a	3096	1/1	0.95	0.20	51,51,51,51	0
57	MG	1F	310	1/1	0.95	0.06	41,41,41,41	0
57	MG	1u	101	1/1	0.95	0.10	64,64,64,64	0
57	MG	1A	3038	1/1	0.95	0.06	26,26,26,26	0
57	MG	1A	3373	1/1	0.95	0.09	44,44,44,44	0
57	MG	1A	3548	1/1	0.95	0.11	41,41,41,41	0
57	MG	2a	3107	1/1	0.95	0.23	50,50,50,50	0
57	MG	1A	3310	1/1	0.95	0.07	32,32,32,32	0
57	MG	1A	3554	1/1	0.95	0.08	37,37,37,37	0
57	MG	1A	3375	1/1	0.95	0.09	36,36,36,36	0
57	MG	2A	3272	1/1	0.95	0.10	38,38,38,38	0
57	MG	2A	3273	1/1	0.95	0.08	42,42,42,42	0
57	MG	2A	3275	1/1	0.95	0.26	38,38,38,38	0
57	MG	1A	3244	1/1	0.95	0.23	51,51,51,51	0
57	MG	1a	3122	1/1	0.95	0.11	58,58,58,58	0
57	MG	2A	3007	1/1	0.95	0.09	42,42,42,42	0
57	MG	2A	3015	1/1	0.95	0.21	50,50,50,50	0
57	MG	2A	3018	1/1	0.95	0.23	37,37,37,37	0
57	MG	2A	3019	1/1	0.95	0.25	53,53,53,53	0
57	MG	1H	201	1/1	0.95	0.11	40,40,40,40	0
57	MG	2a	3125	1/1	0.95	0.13	56,56,56,56	0
57	MG	2A	3284	1/1	0.95	0.08	51,51,51,51	0
57	MG	2A	3587	1/1	0.95	0.11	54,54,54,54	0
57	MG	2A	3024	1/1	0.95	0.14	50,50,50,50	0
57	MG	1A	3245	1/1	0.95	0.16	49,49,49,49	0
57	MG	2A	3594	1/1	0.95	0.06	67,67,67,67	0
57	MG	1A	3936	1/1	0.95	0.05	20,20,20,20	0
57	MG	1A	3062	1/1	0.95	0.08	46,46,46,46	0
57	MG	2A	3031	1/1	0.95	0.13	59,59,59,59	0
57	MG	1A	3651	1/1	0.95	0.07	34,34,34,34	0
57	MG	1a	3130	1/1	0.95	0.08	55,55,55,55	0
57	MG	2A	3035	1/1	0.95	0.08	45,45,45,45	0
57	MG	1Q	204	1/1	0.95	0.06	30,30,30,30	0
57	MG	1A	3259	1/1	0.95	0.19	57,57,57,57	0
57	MG	2a	3141	1/1	0.95	0.08	68,68,68,68	0
57	MG	2A	3605	1/1	0.95	0.10	34,34,34,34	0
57	MG	1A	3653	1/1	0.95	0.36	44,44,44,44	0
57	MG	2A	3607	1/1	0.95	0.08	43,43,43,43	0
57	MG	2a	3146	1/1	0.95	0.10	49,49,49,49	0
57	MG	2A	3609	1/1	0.95	0.09	48,48,48,48	0
57	MG	1A	3654	1/1	0.95	0.05	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	3151	1/1	0.95	0.10	60,60,60,60	0
57	MG	2a	3152	1/1	0.95	0.08	65,65,65,65	0
57	MG	2A	3612	1/1	0.95	0.07	59,59,59,59	0
57	MG	1A	3944	1/1	0.95	0.05	35,35,35,35	0
57	MG	2A	3303	1/1	0.95	0.18	45,45,45,45	0
57	MG	2A	3615	1/1	0.95	0.06	71,71,71,71	0
57	MG	1A	3946	1/1	0.95	0.07	23,23,23,23	0
57	MG	2a	3159	1/1	0.95	0.10	62,62,62,62	0
57	MG	1A	3781	1/1	0.95	0.31	49,49,49,49	0
57	MG	1A	3656	1/1	0.95	0.28	48,48,48,48	0
57	MG	1A	3152	1/1	0.95	0.21	31,31,31,31	0
57	MG	1A	3385	1/1	0.95	0.11	32,32,32,32	0
57	MG	2a	3165	1/1	0.95	0.10	49,49,49,49	0
57	MG	2A	3621	1/1	0.95	0.09	62,62,62,62	0
57	MG	1A	3953	1/1	0.95	0.12	50,50,50,50	0
57	MG	2A	3625	1/1	0.95	0.07	44,44,44,44	0
57	MG	1A	3661	1/1	0.95	0.07	21,21,21,21	0
57	MG	2A	3627	1/1	0.95	0.08	38,38,38,38	0
57	MG	2a	3171	1/1	0.95	0.07	66,66,66,66	0
57	MG	1a	3149	1/1	0.95	0.11	48,48,48,48	0
57	MG	2A	3633	1/1	0.95	0.12	61,61,61,61	0
57	MG	2a	3174	1/1	0.95	0.06	54,54,54,54	0
57	MG	2A	3636	1/1	0.95	0.07	53,53,53,53	0
57	MG	1A	3789	1/1	0.95	0.08	20,20,20,20	0
57	MG	2A	3642	1/1	0.95	0.09	45,45,45,45	0
57	MG	1Y	201	1/1	0.95	0.10	53,53,53,53	0
57	MG	2A	3315	1/1	0.95	0.14	53,53,53,53	0
57	MG	2A	3061	1/1	0.95	0.08	43,43,43,43	0
57	MG	1A	3564	1/1	0.95	0.08	45,45,45,45	0
57	MG	2A	3648	1/1	0.95	0.12	53,53,53,53	0
57	MG	2A	3063	1/1	0.95	0.09	46,46,46,46	0
57	MG	1A	3565	1/1	0.95	0.10	49,49,49,49	0
57	MG	2A	3065	1/1	0.95	0.10	54,54,54,54	0
57	MG	2A	3325	1/1	0.95	0.14	50,50,50,50	0
57	MG	2A	3661	1/1	0.95	0.07	57,57,57,57	0
57	MG	1a	3154	1/1	0.95	0.13	60,60,60,60	0
57	MG	2r	101	1/1	0.95	0.12	63,63,63,63	0
57	MG	2x	101	1/1	0.95	0.10	50,50,50,50	0
57	MG	2A	3069	1/1	0.95	0.20	50,50,50,50	0
58	MPD	18	104	8/8	0.95	0.14	31,40,43,44	0
57	MG	1A	3323	1/1	0.95	0.08	27,27,27,27	0
57	MG	1A	3324	1/1	0.95	0.06	35,35,35,35	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3333	1/1	0.95	0.06	31,31,31,31	0
57	MG	1A	3158	1/1	0.95	0.14	27,27,27,27	0
57	MG	1A	3327	1/1	0.95	0.09	41,41,41,41	0
57	MG	1a	3160	1/1	0.95	0.08	60,60,60,60	0
57	MG	2A	3081	1/1	0.95	0.10	73,73,73,73	0
57	MG	1A	3803	1/1	0.95	0.06	52,52,52,52	0
57	MG	1a	3014	1/1	0.96	0.05	47,47,47,47	0
57	MG	1A	3156	1/1	0.96	0.15	39,39,39,39	0
57	MG	1A	3793	1/1	0.96	0.07	45,45,45,45	0
57	MG	1A	3549	1/1	0.96	0.05	43,43,43,43	0
57	MG	1A	3551	1/1	0.96	0.17	29,29,29,29	0
57	MG	1A	3552	1/1	0.96	0.06	40,40,40,40	0
57	MG	1A	3978	1/1	0.96	0.07	46,46,46,46	0
57	MG	2A	3144	1/1	0.96	0.14	28,28,28,28	0
57	MG	1a	3213	1/1	0.96	0.06	54,54,54,54	0
57	MG	1A	3797	1/1	0.96	0.08	39,39,39,39	0
57	MG	1A	3798	1/1	0.96	0.06	31,31,31,31	0
57	MG	1A	3081	1/1	0.96	0.11	31,31,31,31	0
57	MG	2A	3419	1/1	0.96	0.08	51,51,51,51	0
57	MG	2A	3151	1/1	0.96	0.05	56,56,56,56	0
57	MG	1A	3801	1/1	0.96	0.07	35,35,35,35	0
57	MG	1A	3349	1/1	0.96	0.09	31,31,31,31	0
57	MG	2A	3424	1/1	0.96	0.11	54,54,54,54	0
57	MG	1A	3660	1/1	0.96	0.07	52,52,52,52	0
57	MG	1A	3350	1/1	0.96	0.05	36,36,36,36	0
57	MG	1A	3662	1/1	0.96	0.08	49,49,49,49	0
57	MG	1A	3807	1/1	0.96	0.18	40,40,40,40	0
57	MG	2A	3159	1/1	0.96	0.07	38,38,38,38	0
57	MG	1A	3352	1/1	0.96	0.07	26,26,26,26	0
57	MG	1A	3002	1/1	0.96	0.06	44,44,44,44	0
57	MG	2A	3163	1/1	0.96	0.09	47,47,47,47	0
57	MG	1A	3277	1/1	0.96	0.13	34,34,34,34	0
57	MG	1a	3227	1/1	0.96	0.11	45,45,45,45	0
57	MG	2A	3166	1/1	0.96	0.12	70,70,70,70	0
57	MG	1A	3016	1/1	0.96	0.17	28,28,28,28	0
57	MG	1A	3672	1/1	0.96	0.08	44,44,44,44	0
57	MG	1A	3673	1/1	0.96	0.12	56,56,56,56	0
57	MG	1A	3998	1/1	0.96	0.13	47,47,47,47	0
57	MG	1A	3560	1/1	0.96	0.06	38,38,38,38	0
57	MG	1a	3235	1/1	0.96	0.12	61,61,61,61	0
57	MG	2B	216	1/1	0.96	0.07	63,63,63,63	0
57	MG	1a	3045	1/1	0.96	0.19	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	3445	1/1	0.96	0.24	71,71,71,71	0
57	MG	1A	3280	1/1	0.96	0.06	33,33,33,33	0
57	MG	1A	3819	1/1	0.96	0.10	33,33,33,33	0
57	MG	2A	3449	1/1	0.96	0.08	43,43,43,43	0
57	MG	1A	3125	1/1	0.96	0.12	45,45,45,45	0
57	MG	2D	307	1/1	0.96	0.07	40,40,40,40	0
57	MG	1A	3823	1/1	0.96	0.06	12,12,12,12	0
57	MG	1A	3677	1/1	0.96	0.11	55,55,55,55	0
57	MG	1a	3244	1/1	0.96	0.08	50,50,50,50	0
57	MG	1A	4006	1/1	0.96	0.14	42,42,42,42	0
57	MG	1A	3456	1/1	0.96	0.09	41,41,41,41	0
57	MG	1a	3249	1/1	0.96	0.13	68,68,68,68	0
57	MG	1A	3286	1/1	0.96	0.17	32,32,32,32	0
57	MG	1A	3566	1/1	0.96	0.17	49,49,49,49	0
57	MG	1A	3458	1/1	0.96	0.07	46,46,46,46	0
57	MG	1A	4014	1/1	0.96	0.13	37,37,37,37	0
57	MG	1a	3255	1/1	0.96	0.06	48,48,48,48	0
57	MG	1A	4015	1/1	0.96	0.09	36,36,36,36	0
57	MG	1A	3288	1/1	0.96	0.09	50,50,50,50	0
57	MG	2O	201	1/1	0.96	0.05	52,52,52,52	0
57	MG	2A	3466	1/1	0.96	0.12	42,42,42,42	0
57	MG	1A	3004	1/1	0.96	0.08	37,37,37,37	0
57	MG	2A	3194	1/1	0.96	0.15	40,40,40,40	0
57	MG	1A	3291	1/1	0.96	0.08	31,31,31,31	0
57	MG	1A	3837	1/1	0.96	0.08	38,38,38,38	0
57	MG	2A	3471	1/1	0.96	0.07	36,36,36,36	0
57	MG	2A	3197	1/1	0.96	0.30	59,59,59,59	0
57	MG	1A	4021	1/1	0.96	0.07	57,57,57,57	0
57	MG	2R	202	1/1	0.96	0.17	51,51,51,51	0
57	MG	1A	3839	1/1	0.96	0.24	42,42,42,42	0
57	MG	1A	3128	1/1	0.96	0.06	54,54,54,54	0
57	MG	2A	3201	1/1	0.96	0.08	61,61,61,61	0
57	MG	1A	3576	1/1	0.96	0.05	57,57,57,57	0
57	MG	1A	3844	1/1	0.96	0.06	38,38,38,38	0
57	MG	1A	3464	1/1	0.96	0.09	52,52,52,52	0
57	MG	1A	3173	1/1	0.96	0.09	22,22,22,22	0
57	MG	1A	3579	1/1	0.96	0.06	21,21,21,21	0
57	MG	1A	3692	1/1	0.96	0.07	47,47,47,47	0
57	MG	1A	3177	1/1	0.96	0.18	47,47,47,47	0
57	MG	1A	3472	1/1	0.96	0.13	49,49,49,49	0
57	MG	1A	3225	1/1	0.96	0.14	28,28,28,28	0
57	MG	2I	101	1/1	0.96	0.08	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	3212	1/1	0.96	0.05	47,47,47,47	0
57	MG	25	103	1/1	0.96	0.12	57,57,57,57	0
57	MG	1A	3299	1/1	0.96	0.08	44,44,44,44	0
57	MG	27	102	1/1	0.96	0.21	32,32,32,32	0
57	MG	1A	3183	1/1	0.96	0.33	32,32,32,32	0
57	MG	2a	3001	1/1	0.96	0.07	50,50,50,50	0
57	MG	2A	3215	1/1	0.96	0.21	53,53,53,53	0
57	MG	1A	3855	1/1	0.96	0.08	35,35,35,35	0
57	MG	1A	3229	1/1	0.96	0.32	39,39,39,39	0
57	MG	2a	3006	1/1	0.96	0.09	50,50,50,50	0
57	MG	2A	3497	1/1	0.96	0.06	63,63,63,63	0
57	MG	1A	3051	1/1	0.96	0.19	42,42,42,42	0
57	MG	1A	3379	1/1	0.96	0.06	33,33,33,33	0
57	MG	2A	3220	1/1	0.96	0.06	44,44,44,44	0
57	MG	2A	3501	1/1	0.96	0.12	50,50,50,50	0
57	MG	1A	3380	1/1	0.96	0.07	50,50,50,50	0
57	MG	1A	3233	1/1	0.96	0.09	44,44,44,44	0
57	MG	2A	3505	1/1	0.96	0.09	59,59,59,59	0
57	MG	1A	3308	1/1	0.96	0.09	49,49,49,49	0
57	MG	1A	3035	1/1	0.96	0.15	42,42,42,42	0
57	MG	1A	3235	1/1	0.96	0.25	34,34,34,34	0
57	MG	1A	3598	1/1	0.96	0.09	50,50,50,50	0
57	MG	1A	3236	1/1	0.96	0.08	35,35,35,35	0
57	MG	1A	3870	1/1	0.96	0.09	25,25,25,25	0
57	MG	1A	3871	1/1	0.96	0.06	50,50,50,50	0
57	MG	1A	3716	1/1	0.96	0.08	33,33,33,33	0
57	MG	1D	301	1/1	0.96	0.24	33,33,33,33	0
57	MG	2a	3029	1/1	0.96	0.14	53,53,53,53	0
57	MG	1D	303	1/1	0.96	0.08	36,36,36,36	0
57	MG	1A	3490	1/1	0.96	0.06	40,40,40,40	0
57	MG	1a	3102	1/1	0.96	0.30	57,57,57,57	0
57	MG	1A	3719	1/1	0.96	0.17	33,33,33,33	0
57	MG	1A	3238	1/1	0.96	0.14	33,33,33,33	0
57	MG	1D	312	1/1	0.96	0.21	40,40,40,40	0
57	MG	2A	3244	1/1	0.96	0.11	53,53,53,53	0
57	MG	1D	313	1/1	0.96	0.07	50,50,50,50	0
57	MG	1A	3314	1/1	0.96	0.11	26,26,26,26	0
57	MG	1A	3187	1/1	0.96	0.06	42,42,42,42	0
57	MG	1A	3319	1/1	0.96	0.07	35,35,35,35	0
57	MG	1A	3501	1/1	0.96	0.09	45,45,45,45	0
57	MG	1A	3320	1/1	0.96	0.42	59,59,59,59	0
57	MG	2a	3044	1/1	0.96	0.14	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	2A	3006	1/1	0.96	0.07	38,38,38,38	0
57	MG	1A	3132	1/1	0.96	0.17	33,33,33,33	0
57	MG	1A	3731	1/1	0.96	0.09	26,26,26,26	0
57	MG	2A	3016	1/1	0.96	0.10	38,38,38,38	0
57	MG	2A	3017	1/1	0.96	0.11	47,47,47,47	0
57	MG	2A	3260	1/1	0.96	0.24	40,40,40,40	0
57	MG	2A	3537	1/1	0.96	0.14	47,47,47,47	0
57	MG	2A	3538	1/1	0.96	0.12	59,59,59,59	0
57	MG	1a	3114	1/1	0.96	0.06	30,30,30,30	0
57	MG	2A	3264	1/1	0.96	0.20	50,50,50,50	0
57	MG	1a	3115	1/1	0.96	0.10	47,47,47,47	0
57	MG	2A	3020	1/1	0.96	0.22	42,42,42,42	0
57	MG	2A	3022	1/1	0.96	0.14	51,51,51,51	0
57	MG	1A	3732	1/1	0.96	0.10	25,25,25,25	0
57	MG	2A	3546	1/1	0.96	0.10	36,36,36,36	0
57	MG	1a	3117	1/1	0.96	0.08	55,55,55,55	0
57	MG	1A	3733	1/1	0.96	0.07	46,46,46,46	0
57	MG	1A	3609	1/1	0.96	0.08	45,45,45,45	0
57	MG	2A	3274	1/1	0.96	0.26	55,55,55,55	0
57	MG	1F	305	1/1	0.96	0.12	23,23,23,23	0
57	MG	1F	307	1/1	0.96	0.07	28,28,28,28	0
57	MG	1F	308	1/1	0.96	0.11	24,24,24,24	0
57	MG	1A	3137	1/1	0.96	0.09	45,45,45,45	0
57	MG	2A	3034	1/1	0.96	0.12	63,63,63,63	0
57	MG	2A	3559	1/1	0.96	0.09	57,57,57,57	0
57	MG	2A	3560	1/1	0.96	0.06	30,30,30,30	0
57	MG	1A	3737	1/1	0.96	0.18	44,44,44,44	0
57	MG	1A	3738	1/1	0.96	0.05	32,32,32,32	0
57	MG	1A	3613	1/1	0.96	0.14	29,29,29,29	0
57	MG	2A	3567	1/1	0.96	0.07	50,50,50,50	0
57	MG	2A	3038	1/1	0.96	0.08	43,43,43,43	0
57	MG	1F	316	1/1	0.96	0.10	50,50,50,50	0
57	MG	2a	3080	1/1	0.96	0.12	55,55,55,55	0
57	MG	1a	3131	1/1	0.96	0.14	57,57,57,57	0
57	MG	2A	3042	1/1	0.96	0.06	27,27,27,27	0
57	MG	1A	3190	1/1	0.96	0.05	28,28,28,28	0
57	MG	1A	3615	1/1	0.96	0.16	35,35,35,35	0
57	MG	2A	3045	1/1	0.96	0.10	52,52,52,52	0
57	MG	2A	3046	1/1	0.96	0.22	63,63,63,63	0
57	MG	2A	3294	1/1	0.96	0.13	50,50,50,50	0
57	MG	2A	3295	1/1	0.96	0.07	42,42,42,42	0
57	MG	2A	3047	1/1	0.96	0.12	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	2a	3091	1/1	0.96	0.15	64,64,64,64	0
57	MG	1A	3092	1/1	0.96	0.16	26,26,26,26	0
57	MG	1A	3508	1/1	0.96	0.07	46,46,46,46	0
57	MG	1A	3510	1/1	0.96	0.04	16,16,16,16	0
57	MG	2A	3586	1/1	0.96	0.06	54,54,54,54	0
57	MG	2A	3051	1/1	0.96	0.12	53,53,53,53	0
57	MG	1H	202	1/1	0.96	0.07	49,49,49,49	0
57	MG	2A	3302	1/1	0.96	0.14	44,44,44,44	0
57	MG	2A	3593	1/1	0.96	0.09	54,54,54,54	0
57	MG	1A	3910	1/1	0.96	0.06	37,37,37,37	0
57	MG	2a	3105	1/1	0.96	0.09	48,48,48,48	0
57	MG	2a	3106	1/1	0.96	0.26	54,54,54,54	0
57	MG	1A	3325	1/1	0.96	0.08	29,29,29,29	0
57	MG	2A	3056	1/1	0.96	0.19	60,60,60,60	0
57	MG	1A	3914	1/1	0.96	0.21	39,39,39,39	0
57	MG	2a	3110	1/1	0.96	0.33	51,51,51,51	0
57	MG	1A	3916	1/1	0.96	0.05	47,47,47,47	0
57	MG	1P	204	1/1	0.96	0.06	69,69,69,69	0
57	MG	1Q	201	1/1	0.96	0.23	35,35,35,35	0
57	MG	1A	3250	1/1	0.96	0.26	30,30,30,30	0
57	MG	1A	3622	1/1	0.96	0.07	23,23,23,23	0
57	MG	1A	3255	1/1	0.96	0.14	36,36,36,36	0
57	MG	1R	203	1/1	0.96	0.07	45,45,45,45	0
57	MG	2A	3066	1/1	0.96	0.08	44,44,44,44	0
57	MG	2a	3121	1/1	0.96	0.09	58,58,58,58	0
57	MG	1A	3922	1/1	0.96	0.11	52,52,52,52	0
57	MG	2A	3068	1/1	0.96	0.09	48,48,48,48	0
57	MG	1A	3407	1/1	0.96	0.13	45,45,45,45	0
57	MG	2A	3070	1/1	0.96	0.11	47,47,47,47	0
57	MG	1A	3257	1/1	0.96	0.08	39,39,39,39	0
57	MG	2A	3074	1/1	0.96	0.07	48,48,48,48	0
57	MG	1A	3412	1/1	0.96	0.13	48,48,48,48	0
57	MG	2A	3077	1/1	0.96	0.08	57,57,57,57	0
57	MG	2A	3329	1/1	0.96	0.05	38,38,38,38	0
57	MG	1A	3012	1/1	0.96	0.10	30,30,30,30	0
57	MG	2A	3331	1/1	0.96	0.05	41,41,41,41	0
57	MG	1U	202	1/1	0.96	0.17	43,43,43,43	0
57	MG	2A	3622	1/1	0.96	0.07	53,53,53,53	0
57	MG	1A	3146	1/1	0.96	0.06	31,31,31,31	0
57	MG	1A	3934	1/1	0.96	0.06	38,38,38,38	0
57	MG	1V	207	1/1	0.96	0.06	45,45,45,45	0
57	MG	2a	3139	1/1	0.96	0.19	76,76,76,76	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3148	1/1	0.96	0.08	40,40,40,40	0
57	MG	1A	3760	1/1	0.96	0.09	35,35,35,35	0
57	MG	2a	3142	1/1	0.96	0.09	60,60,60,60	0
57	MG	1A	3150	1/1	0.96	0.07	61,61,61,61	0
57	MG	1a	3166	1/1	0.96	0.10	51,51,51,51	0
57	MG	2A	3639	1/1	0.96	0.10	55,55,55,55	0
57	MG	1A	3762	1/1	0.96	0.10	48,48,48,48	0
57	MG	2A	3641	1/1	0.96	0.06	43,43,43,43	0
57	MG	2A	3089	1/1	0.96	0.10	45,45,45,45	0
57	MG	1A	3099	1/1	0.96	0.22	38,38,38,38	0
57	MG	1A	3266	1/1	0.96	0.12	37,37,37,37	0
57	MG	1A	3637	1/1	0.96	0.14	41,41,41,41	0
57	MG	1a	3172	1/1	0.96	0.15	58,58,58,58	0
57	MG	10	106	1/1	0.96	0.08	47,47,47,47	0
57	MG	2A	3649	1/1	0.96	0.08	46,46,46,46	0
57	MG	1A	3201	1/1	0.96	0.09	41,41,41,41	0
57	MG	2a	3158	1/1	0.96	0.06	67,67,67,67	0
57	MG	1A	3269	1/1	0.96	0.09	48,48,48,48	0
57	MG	1A	3771	1/1	0.96	0.04	36,36,36,36	0
57	MG	2A	3357	1/1	0.96	0.06	47,47,47,47	0
57	MG	2a	3162	1/1	0.96	0.07	57,57,57,57	0
57	MG	2A	3658	1/1	0.96	0.07	66,66,66,66	0
57	MG	2A	3659	1/1	0.96	0.05	50,50,50,50	0
57	MG	1A	3640	1/1	0.96	0.14	34,34,34,34	0
57	MG	1A	3641	1/1	0.96	0.11	32,32,32,32	0
57	MG	1a	3179	1/1	0.96	0.05	65,65,65,65	0
57	MG	1A	3344	1/1	0.96	0.06	25,25,25,25	0
57	MG	2A	3365	1/1	0.96	0.06	34,34,34,34	0
57	MG	11	105	1/1	0.96	0.06	26,26,26,26	0
57	MG	1A	3428	1/1	0.96	0.06	20,20,20,20	0
57	MG	15	3102	1/1	0.96	0.12	34,34,34,34	0
57	MG	1A	3430	1/1	0.96	0.05	15,15,15,15	0
57	MG	2A	3113	1/1	0.96	0.14	52,52,52,52	0
57	MG	1A	3432	1/1	0.96	0.14	46,46,46,46	0
57	MG	2A	3374	1/1	0.96	0.09	46,46,46,46	0
57	MG	2a	3182	1/1	0.96	0.05	71,71,71,71	0
57	MG	1a	3186	1/1	0.96	0.08	50,50,50,50	0
57	MG	2A	3679	1/1	0.96	0.09	48,48,48,48	0
57	MG	2A	3683	1/1	0.96	0.06	38,38,38,38	0
57	MG	1A	3544	1/1	0.96	0.05	36,36,36,36	0
57	MG	1A	3783	1/1	0.96	0.14	35,35,35,35	0
57	MG	18	102	1/1	0.96	0.17	36,36,36,36	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3379	1/1	0.96	0.09	42,42,42,42	0
57	MG	1A	3960	1/1	0.96	0.08	38,38,38,38	0
57	MG	19	101	1/1	0.96	0.14	46,46,46,46	0
57	MG	1A	3961	1/1	0.96	0.10	47,47,47,47	0
57	MG	1A	3784	1/1	0.96	0.09	29,29,29,29	0
57	MG	2A	3124	1/1	0.96	0.06	38,38,38,38	0
57	MG	1A	3648	1/1	0.96	0.08	32,32,32,32	0
57	MG	2A	3391	1/1	0.96	0.07	53,53,53,53	0
57	MG	2A	3392	1/1	0.96	0.08	40,40,40,40	0
57	MG	1A	3649	1/1	0.96	0.10	49,49,49,49	0
57	MG	1A	3153	1/1	0.96	0.04	31,31,31,31	0
57	MG	2A	3703	1/1	0.96	0.11	55,55,55,55	0
57	MG	1A	3788	1/1	0.96	0.15	39,39,39,39	0
57	MG	1a	3007	1/1	0.96	0.08	47,47,47,47	0
57	MG	1a	3008	1/1	0.96	0.07	55,55,55,55	0
57	MG	1A	3154	1/1	0.96	0.06	48,48,48,48	0
57	MG	1A	3790	1/1	0.96	0.10	47,47,47,47	0
57	MG	1a	3013	1/1	0.96	0.08	54,54,54,54	0
57	MG	1A	3939	1/1	0.97	0.08	44,44,44,44	0
57	MG	10	103	1/1	0.97	0.06	38,38,38,38	0
57	MG	2A	3701	1/1	0.97	0.06	50,50,50,50	0
57	MG	2A	3385	1/1	0.97	0.06	37,37,37,37	0
57	MG	2A	3118	1/1	0.97	0.20	57,57,57,57	0
57	MG	2A	3387	1/1	0.97	0.10	41,41,41,41	0
57	MG	1A	3756	1/1	0.97	0.05	40,40,40,40	0
57	MG	10	105	1/1	0.97	0.07	43,43,43,43	0
57	MG	1A	3757	1/1	0.97	0.05	31,31,31,31	0
57	MG	1A	3611	1/1	0.97	0.17	31,31,31,31	0
57	MG	1A	3208	1/1	0.97	0.16	39,39,39,39	0
57	MG	2A	3710	1/1	0.97	0.07	53,53,53,53	0
57	MG	1A	3289	1/1	0.97	0.05	45,45,45,45	0
57	MG	11	101	1/1	0.97	0.10	39,39,39,39	0
57	MG	1A	3484	1/1	0.97	0.07	29,29,29,29	0
57	MG	2A	3128	1/1	0.97	0.08	33,33,33,33	0
57	MG	1A	3209	1/1	0.97	0.17	42,42,42,42	0
57	MG	1A	3764	1/1	0.97	0.05	30,30,30,30	0
57	MG	2A	3401	1/1	0.97	0.05	56,56,56,56	0
57	MG	2A	3402	1/1	0.97	0.05	45,45,45,45	0
57	MG	1A	3950	1/1	0.97	0.05	43,43,43,43	0
57	MG	13	101	1/1	0.97	0.11	43,43,43,43	0
57	MG	1A	3765	1/1	0.97	0.05	38,38,38,38	0
57	MG	15	3101	1/1	0.97	0.10	26,26,26,26	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3151	1/1	0.97	0.08	43,43,43,43	0
57	MG	1A	3213	1/1	0.97	0.21	37,37,37,37	0
57	MG	1A	3954	1/1	0.97	0.06	55,55,55,55	0
57	MG	15	3109	1/1	0.97	0.08	51,51,51,51	0
57	MG	17	104	1/1	0.97	0.14	30,30,30,30	0
57	MG	1A	3488	1/1	0.97	0.08	27,27,27,27	0
57	MG	17	106	1/1	0.97	0.07	40,40,40,40	0
57	MG	1A	3489	1/1	0.97	0.06	23,23,23,23	0
57	MG	1A	3215	1/1	0.97	0.45	35,35,35,35	0
57	MG	1A	3295	1/1	0.97	0.06	15,15,15,15	0
57	MG	1A	3065	1/1	0.97	0.04	38,38,38,38	0
57	MG	1a	3212	1/1	0.97	0.05	69,69,69,69	0
57	MG	2B	205	1/1	0.97	0.18	54,54,54,54	0
57	MG	1A	3495	1/1	0.97	0.06	48,48,48,48	0
57	MG	1A	3217	1/1	0.97	0.18	31,31,31,31	0
57	MG	1A	3497	1/1	0.97	0.12	22,22,22,22	0
57	MG	1A	3777	1/1	0.97	0.06	41,41,41,41	0
57	MG	1A	3498	1/1	0.97	0.05	52,52,52,52	0
57	MG	2A	3155	1/1	0.97	0.12	52,52,52,52	0
57	MG	1a	3005	1/1	0.97	0.23	54,54,54,54	0
57	MG	1A	3780	1/1	0.97	0.13	43,43,43,43	0
57	MG	1A	3499	1/1	0.97	0.05	32,32,32,32	0
57	MG	1A	3632	1/1	0.97	0.08	24,24,24,24	0
57	MG	1A	3066	1/1	0.97	0.08	42,42,42,42	0
57	MG	2A	3436	1/1	0.97	0.06	48,48,48,48	0
57	MG	1A	3634	1/1	0.97	0.07	35,35,35,35	0
57	MG	2B	219	1/1	0.97	0.08	66,66,66,66	0
57	MG	1a	3224	1/1	0.97	0.04	46,46,46,46	0
57	MG	1A	3096	1/1	0.97	0.08	50,50,50,50	0
57	MG	2D	303	1/1	0.97	0.34	39,39,39,39	0
57	MG	2D	304	1/1	0.97	0.09	34,34,34,34	0
57	MG	1A	3386	1/1	0.97	0.06	31,31,31,31	0
57	MG	1A	3302	1/1	0.97	0.05	46,46,46,46	0
57	MG	1A	3504	1/1	0.97	0.07	42,42,42,42	0
57	MG	1A	3098	1/1	0.97	0.10	49,49,49,49	0
57	MG	1a	3230	1/1	0.97	0.08	64,64,64,64	0
57	MG	1A	3980	1/1	0.97	0.06	48,48,48,48	0
57	MG	2A	3446	1/1	0.97	0.11	57,57,57,57	0
57	MG	1A	3305	1/1	0.97	0.12	45,45,45,45	0
57	MG	1a	3233	1/1	0.97	0.14	52,52,52,52	0
57	MG	2A	3173	1/1	0.97	0.09	40,40,40,40	0
57	MG	1A	3390	1/1	0.97	0.04	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	1A	3984	1/1	0.97	0.05	47,47,47,47	0
57	MG	2A	3452	1/1	0.97	0.04	36,36,36,36	0
57	MG	1A	3157	1/1	0.97	0.09	44,44,44,44	0
57	MG	2A	3454	1/1	0.97	0.04	45,45,45,45	0
57	MG	1A	3025	1/1	0.97	0.12	33,33,33,33	0
57	MG	1a	3026	1/1	0.97	0.05	49,49,49,49	0
57	MG	1a	3239	1/1	0.97	0.08	58,58,58,58	0
57	MG	1A	3159	1/1	0.97	0.12	22,22,22,22	0
57	MG	1A	3645	1/1	0.97	0.11	37,37,37,37	0
57	MG	1A	3394	1/1	0.97	0.08	22,22,22,22	0
57	MG	1A	3105	1/1	0.97	0.06	31,31,31,31	0
57	MG	1A	3396	1/1	0.97	0.10	27,27,27,27	0
57	MG	1a	3246	1/1	0.97	0.04	50,50,50,50	0
57	MG	1A	3517	1/1	0.97	0.05	36,36,36,36	0
57	MG	1A	3230	1/1	0.97	0.12	33,33,33,33	0
57	MG	1a	3250	1/1	0.97	0.05	57,57,57,57	0
57	MG	1A	3311	1/1	0.97	0.06	19,19,19,19	0
57	MG	2T	202	1/1	0.97	0.07	51,51,51,51	0
57	MG	1A	3106	1/1	0.97	0.06	43,43,43,43	0
57	MG	1A	3232	1/1	0.97	0.09	31,31,31,31	0
57	MG	1A	3522	1/1	0.97	0.08	24,24,24,24	0
57	MG	1A	3404	1/1	0.97	0.04	22,22,22,22	0
57	MG	2A	3472	1/1	0.97	0.05	66,66,66,66	0
57	MG	1A	3657	1/1	0.97	0.10	38,38,38,38	0
57	MG	1a	3043	1/1	0.97	0.05	54,54,54,54	0
57	MG	2X	101	1/1	0.97	0.08	51,51,51,51	0
57	MG	1A	3811	1/1	0.97	0.16	42,42,42,42	0
57	MG	1A	3525	1/1	0.97	0.04	25,25,25,25	0
57	MG	1a	3046	1/1	0.97	0.23	49,49,49,49	0
57	MG	1A	4003	1/1	0.97	0.34	44,44,44,44	0
57	MG	1A	3048	1/1	0.97	0.10	37,37,37,37	0
57	MG	1a	3049	1/1	0.97	0.16	41,41,41,41	0
57	MG	1A	3528	1/1	0.97	0.05	39,39,39,39	0
57	MG	1A	3529	1/1	0.97	0.04	23,23,23,23	0
57	MG	28	101	1/1	0.97	0.06	51,51,51,51	0
57	MG	2A	3206	1/1	0.97	0.18	41,41,41,41	0
57	MG	1A	4007	1/1	0.97	0.05	54,54,54,54	0
57	MG	1A	4008	1/1	0.97	0.13	43,43,43,43	0
57	MG	1a	3272	1/1	0.97	0.13	58,58,58,58	0
57	MG	1A	3165	1/1	0.97	0.09	24,24,24,24	0
57	MG	1A	3318	1/1	0.97	0.06	21,21,21,21	0
57	MG	1A	4011	1/1	0.97	0.07	53,53,53,53	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3664	1/1	0.97	0.04	46,46,46,46	0
57	MG	1A	3665	1/1	0.97	0.11	45,45,45,45	0
57	MG	1A	3166	1/1	0.97	0.07	42,42,42,42	0
57	MG	1A	3409	1/1	0.97	0.06	54,54,54,54	0
57	MG	1A	3825	1/1	0.97	0.04	26,26,26,26	0
57	MG	2a	3014	1/1	0.97	0.17	55,55,55,55	0
57	MG	1A	3668	1/1	0.97	0.06	49,49,49,49	0
57	MG	1A	4018	1/1	0.97	0.06	39,39,39,39	0
57	MG	1e	202	1/1	0.97	0.06	51,51,51,51	0
57	MG	1A	3669	1/1	0.97	0.06	46,46,46,46	0
57	MG	2a	3020	1/1	0.97	0.08	59,59,59,59	0
57	MG	1A	3410	1/1	0.97	0.06	18,18,18,18	0
57	MG	2A	3504	1/1	0.97	0.05	35,35,35,35	0
57	MG	1A	3167	1/1	0.97	0.04	27,27,27,27	0
57	MG	1A	4022	1/1	0.97	0.05	36,36,36,36	0
57	MG	1A	3414	1/1	0.97	0.09	46,46,46,46	0
57	MG	1i	201	1/1	0.97	0.04	51,51,51,51	0
57	MG	2A	3227	1/1	0.97	0.11	44,44,44,44	0
57	MG	2A	3228	1/1	0.97	0.10	33,33,33,33	0
57	MG	1a	3072	1/1	0.97	0.14	44,44,44,44	0
57	MG	2A	3232	1/1	0.97	0.10	52,52,52,52	0
57	MG	1A	3237	1/1	0.97	0.14	26,26,26,26	0
57	MG	1A	3538	1/1	0.97	0.21	31,31,31,31	0
57	MG	2a	3034	1/1	0.97	0.06	45,45,45,45	0
57	MG	2A	3516	1/1	0.97	0.08	74,74,74,74	0
57	MG	1A	3836	1/1	0.97	0.05	36,36,36,36	0
57	MG	1A	3539	1/1	0.97	0.07	41,41,41,41	0
57	MG	1A	3838	1/1	0.97	0.05	42,42,42,42	0
57	MG	1A	3540	1/1	0.97	0.05	22,22,22,22	0
57	MG	2A	3521	1/1	0.97	0.06	53,53,53,53	0
57	MG	1A	3840	1/1	0.97	0.06	44,44,44,44	0
57	MG	1A	3541	1/1	0.97	0.04	23,23,23,23	0
57	MG	1A	3416	1/1	0.97	0.05	38,38,38,38	0
57	MG	2A	3242	1/1	0.97	0.20	37,37,37,37	0
57	MG	1A	3417	1/1	0.97	0.04	21,21,21,21	0
57	MG	1A	3072	1/1	0.97	0.11	31,31,31,31	0
57	MG	1A	3169	1/1	0.97	0.33	35,35,35,35	0
57	MG	1A	3114	1/1	0.97	0.10	39,39,39,39	0
57	MG	1A	3172	1/1	0.97	0.12	46,46,46,46	0
57	MG	1A	3422	1/1	0.97	0.05	44,44,44,44	0
57	MG	1A	3120	1/1	0.97	0.06	31,31,31,31	0
57	MG	2A	3250	1/1	0.97	0.08	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	3248	1/1	0.97	0.20	39,39,39,39	0
57	MG	2A	3002	1/1	0.97	0.07	41,41,41,41	0
57	MG	2a	3055	1/1	0.97	0.09	51,51,51,51	0
57	MG	1a	3091	1/1	0.97	0.10	51,51,51,51	0
57	MG	1A	3852	1/1	0.97	0.09	36,36,36,36	0
57	MG	1a	3093	1/1	0.97	0.09	47,47,47,47	0
57	MG	2A	3257	1/1	0.97	0.07	36,36,36,36	0
57	MG	2A	3009	1/1	0.97	0.07	45,45,45,45	0
57	MG	2A	3259	1/1	0.97	0.10	24,24,24,24	0
57	MG	2A	3011	1/1	0.97	0.05	34,34,34,34	0
57	MG	2A	3261	1/1	0.97	0.11	55,55,55,55	0
57	MG	2A	3262	1/1	0.97	0.14	33,33,33,33	0
57	MG	2A	3013	1/1	0.97	0.06	31,31,31,31	0
57	MG	2A	3014	1/1	0.97	0.06	49,49,49,49	0
57	MG	2A	3549	1/1	0.97	0.05	45,45,45,45	0
57	MG	1A	3121	1/1	0.97	0.10	31,31,31,31	0
57	MG	1A	3179	1/1	0.97	0.12	42,42,42,42	0
57	MG	2A	3267	1/1	0.97	0.04	25,25,25,25	0
57	MG	1B	224	1/1	0.97	0.04	38,38,38,38	0
57	MG	2A	3269	1/1	0.97	0.06	45,45,45,45	0
57	MG	1A	3427	1/1	0.97	0.09	16,16,16,16	0
57	MG	1A	3856	1/1	0.97	0.06	40,40,40,40	0
57	MG	2a	3075	1/1	0.97	0.13	53,53,53,53	0
57	MG	2a	3076	1/1	0.97	0.06	61,61,61,61	0
57	MG	1A	3693	1/1	0.97	0.16	40,40,40,40	0
57	MG	1A	3251	1/1	0.97	0.05	37,37,37,37	0
57	MG	1A	3429	1/1	0.97	0.11	27,27,27,27	0
57	MG	1A	3253	1/1	0.97	0.14	37,37,37,37	0
57	MG	2a	3081	1/1	0.97	0.30	51,51,51,51	0
57	MG	2A	3025	1/1	0.97	0.07	34,34,34,34	0
57	MG	1A	3431	1/1	0.97	0.07	32,32,32,32	0
57	MG	2A	3027	1/1	0.97	0.08	38,38,38,38	0
57	MG	1A	3863	1/1	0.97	0.04	28,28,28,28	0
57	MG	1D	304	1/1	0.97	0.06	34,34,34,34	0
57	MG	1D	307	1/1	0.97	0.08	47,47,47,47	0
57	MG	1A	3182	1/1	0.97	0.12	30,30,30,30	0
57	MG	1A	3074	1/1	0.97	0.09	29,29,29,29	0
57	MG	2A	3573	1/1	0.97	0.05	49,49,49,49	0
57	MG	1A	3334	1/1	0.97	0.06	25,25,25,25	0
57	MG	2a	3092	1/1	0.97	0.29	52,52,52,52	0
57	MG	1A	3702	1/1	0.97	0.09	54,54,54,54	0
57	MG	2A	3577	1/1	0.97	0.07	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	3335	1/1	0.97	0.07	20,20,20,20	0
57	MG	2A	3289	1/1	0.97	0.10	42,42,42,42	0
57	MG	2a	3097	1/1	0.97	0.29	55,55,55,55	0
57	MG	1A	3337	1/1	0.97	0.10	14,14,14,14	0
57	MG	1A	3258	1/1	0.97	0.09	47,47,47,47	0
57	MG	2a	3100	1/1	0.97	0.18	41,41,41,41	0
57	MG	1E	302	1/1	0.97	0.18	35,35,35,35	0
57	MG	2A	3040	1/1	0.97	0.06	44,44,44,44	0
57	MG	1A	3872	1/1	0.97	0.11	27,27,27,27	0
57	MG	1A	3037	1/1	0.97	0.07	44,44,44,44	0
57	MG	1A	3440	1/1	0.97	0.03	14,14,14,14	0
57	MG	1a	3118	1/1	0.97	0.06	42,42,42,42	0
57	MG	2A	3589	1/1	0.97	0.06	42,42,42,42	0
57	MG	1E	307	1/1	0.97	0.06	45,45,45,45	0
57	MG	1A	3710	1/1	0.97	0.05	37,37,37,37	0
57	MG	1A	3574	1/1	0.97	0.10	34,34,34,34	0
57	MG	1A	3878	1/1	0.97	0.07	39,39,39,39	0
57	MG	1A	3712	1/1	0.97	0.03	32,32,32,32	0
57	MG	1A	3341	1/1	0.97	0.06	34,34,34,34	0
57	MG	1a	3125	1/1	0.97	0.05	50,50,50,50	0
57	MG	2A	3305	1/1	0.97	0.18	46,46,46,46	0
57	MG	2a	3118	1/1	0.97	0.07	67,67,67,67	0
57	MG	2A	3052	1/1	0.97	0.04	32,32,32,32	0
57	MG	1F	303	1/1	0.97	0.15	28,28,28,28	0
57	MG	1A	3077	1/1	0.97	0.07	27,27,27,27	0
57	MG	1A	3715	1/1	0.97	0.07	35,35,35,35	0
57	MG	1A	3262	1/1	0.97	0.06	44,44,44,44	0
57	MG	1A	3078	1/1	0.97	0.06	44,44,44,44	0
57	MG	2A	3058	1/1	0.97	0.05	32,32,32,32	0
57	MG	1F	311	1/1	0.97	0.05	33,33,33,33	0
57	MG	1A	3056	1/1	0.97	0.08	38,38,38,38	0
57	MG	1A	3032	1/1	0.97	0.08	40,40,40,40	0
57	MG	1A	3450	1/1	0.97	0.08	40,40,40,40	0
57	MG	1a	3138	1/1	0.97	0.08	48,48,48,48	0
57	MG	1A	3454	1/1	0.97	0.09	56,56,56,56	0
57	MG	1A	3009	1/1	0.97	0.04	26,26,26,26	0
57	MG	1A	3133	1/1	0.97	0.08	32,32,32,32	0
57	MG	2A	3324	1/1	0.97	0.10	26,26,26,26	0
57	MG	1A	3268	1/1	0.97	0.12	31,31,31,31	0
57	MG	1A	3459	1/1	0.97	0.05	53,53,53,53	0
57	MG	1A	3134	1/1	0.97	0.07	37,37,37,37	0
57	MG	1A	3084	1/1	0.97	0.15	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	3903	1/1	0.97	0.04	30,30,30,30	0
57	MG	2A	3073	1/1	0.97	0.05	24,24,24,24	0
57	MG	1A	3040	1/1	0.97	0.05	37,37,37,37	0
57	MG	2A	3628	1/1	0.97	0.05	49,49,49,49	0
57	MG	2A	3629	1/1	0.97	0.05	69,69,69,69	0
57	MG	1A	3592	1/1	0.97	0.15	52,52,52,52	0
57	MG	2A	3632	1/1	0.97	0.07	70,70,70,70	0
57	MG	1P	201	1/1	0.97	0.22	31,31,31,31	0
57	MG	2A	3634	1/1	0.97	0.04	40,40,40,40	0
57	MG	2a	3149	1/1	0.97	0.11	54,54,54,54	0
57	MG	2a	3150	1/1	0.97	0.07	49,49,49,49	0
57	MG	2A	3635	1/1	0.97	0.05	68,68,68,68	0
57	MG	1A	3593	1/1	0.97	0.03	18,18,18,18	0
57	MG	1A	3139	1/1	0.97	0.12	38,38,38,38	0
57	MG	2A	3337	1/1	0.97	0.07	42,42,42,42	0
57	MG	2A	3339	1/1	0.97	0.10	53,53,53,53	0
57	MG	1A	3140	1/1	0.97	0.08	32,32,32,32	0
57	MG	2A	3341	1/1	0.97	0.10	38,38,38,38	0
57	MG	1Q	203	1/1	0.97	0.16	29,29,29,29	0
57	MG	1A	3913	1/1	0.97	0.12	28,28,28,28	0
57	MG	1Q	205	1/1	0.97	0.06	47,47,47,47	0
57	MG	1A	3465	1/1	0.97	0.07	55,55,55,55	0
57	MG	2A	3346	1/1	0.97	0.09	49,49,49,49	0
57	MG	1A	3087	1/1	0.97	0.10	41,41,41,41	0
57	MG	1A	3008	1/1	0.97	0.06	27,27,27,27	0
57	MG	2A	3349	1/1	0.97	0.07	52,52,52,52	0
57	MG	1A	3471	1/1	0.97	0.04	26,26,26,26	0
57	MG	2A	3088	1/1	0.97	0.14	60,60,60,60	0
57	MG	1S	201	1/1	0.97	0.10	48,48,48,48	0
57	MG	1A	3920	1/1	0.97	0.08	49,49,49,49	0
57	MG	1T	202	1/1	0.97	0.09	41,41,41,41	0
57	MG	2A	3092	1/1	0.97	0.08	42,42,42,42	0
57	MG	2A	3664	1/1	0.97	0.04	47,47,47,47	0
57	MG	2A	3093	1/1	0.97	0.06	45,45,45,45	0
57	MG	1A	3147	1/1	0.97	0.13	29,29,29,29	0
57	MG	2a	3175	1/1	0.97	0.09	55,55,55,55	0
57	MG	2a	3176	1/1	0.97	0.12	63,63,63,63	0
57	MG	2A	3359	1/1	0.97	0.11	35,35,35,35	0
57	MG	2a	3178	1/1	0.97	0.09	74,74,74,74	0
57	MG	2A	3360	1/1	0.97	0.12	33,33,33,33	0
57	MG	2a	3180	1/1	0.97	0.05	61,61,61,61	0
57	MG	2a	3181	1/1	0.97	0.04	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	3602	1/1	0.97	0.04	33,33,33,33	0
57	MG	2A	3670	1/1	0.97	0.09	50,50,50,50	0
57	MG	1A	3042	1/1	0.97	0.07	28,28,28,28	0
57	MG	2A	3097	1/1	0.97	0.08	48,48,48,48	0
57	MG	2A	3673	1/1	0.97	0.06	49,49,49,49	0
57	MG	2a	3187	1/1	0.97	0.28	57,57,57,57	0
57	MG	1A	3364	1/1	0.97	0.05	42,42,42,42	0
57	MG	2A	3676	1/1	0.97	0.05	40,40,40,40	0
57	MG	1a	3170	1/1	0.97	0.07	57,57,57,57	0
57	MG	1U	203	1/1	0.97	0.11	36,36,36,36	0
57	MG	1V	202	1/1	0.97	0.16	23,23,23,23	0
57	MG	2A	3680	1/1	0.97	0.07	63,63,63,63	0
57	MG	2A	3682	1/1	0.97	0.07	53,53,53,53	0
57	MG	1A	3928	1/1	0.97	0.08	32,32,32,32	0
57	MG	2A	3684	1/1	0.97	0.07	45,45,45,45	0
57	MG	2t	201	1/1	0.97	0.07	35,35,35,35	0
57	MG	1V	205	1/1	0.97	0.04	44,44,44,44	0
57	MG	1A	3282	1/1	0.97	0.08	33,33,33,33	0
57	MG	1A	3149	1/1	0.97	0.06	32,32,32,32	0
57	MG	2A	3109	1/1	0.97	0.07	50,50,50,50	0
57	MG	1A	3369	1/1	0.97	0.04	25,25,25,25	0
57	MG	1W	202	1/1	0.97	0.05	38,38,38,38	0
57	MG	1A	3752	1/1	0.97	0.12	34,34,34,34	0
57	MG	1A	3479	1/1	0.97	0.04	28,28,28,28	0
57	MG	2A	3380	1/1	0.97	0.09	42,42,42,42	0
57	MG	1A	3091	1/1	0.97	0.19	31,31,31,31	0
57	MG	1A	3287	1/1	0.97	0.26	44,44,44,44	0
60	ZN	29	501	1/1	0.97	0.05	78,78,78,78	0
57	MG	2A	3076	1/1	0.98	0.09	35,35,35,35	0
57	MG	2E	305	1/1	0.98	0.04	38,38,38,38	0
57	MG	2E	306	1/1	0.98	0.07	30,30,30,30	0
57	MG	1A	3729	1/1	0.98	0.09	23,23,23,23	0
57	MG	2F	301	1/1	0.98	0.23	40,40,40,40	0
57	MG	1A	3135	1/1	0.98	0.22	24,24,24,24	0
57	MG	1A	3036	1/1	0.98	0.05	27,27,27,27	0
57	MG	2F	304	1/1	0.98	0.19	41,41,41,41	0
57	MG	1A	3067	1/1	0.98	0.05	27,27,27,27	0
57	MG	1A	3338	1/1	0.98	0.08	44,44,44,44	0
57	MG	1A	3859	1/1	0.98	0.05	38,38,38,38	0
57	MG	1A	3620	1/1	0.98	0.05	37,37,37,37	0
57	MG	1A	3735	1/1	0.98	0.07	27,27,27,27	0
57	MG	1a	3009	1/1	0.98	0.27	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	1A	3191	1/1	0.98	0.18	34,34,34,34	0
57	MG	1A	3192	1/1	0.98	0.15	45,45,45,45	0
57	MG	1A	3864	1/1	0.98	0.04	31,31,31,31	0
57	MG	1A	3094	1/1	0.98	0.04	14,14,14,14	0
57	MG	1A	3049	1/1	0.98	0.06	35,35,35,35	0
57	MG	1A	3343	1/1	0.98	0.05	31,31,31,31	0
57	MG	1a	3202	1/1	0.98	0.04	70,70,70,70	0
57	MG	1A	3524	1/1	0.98	0.05	36,36,36,36	0
57	MG	1a	3018	1/1	0.98	0.06	46,46,46,46	0
57	MG	1A	3142	1/1	0.98	0.03	31,31,31,31	0
57	MG	1A	3526	1/1	0.98	0.24	26,26,26,26	0
57	MG	1a	3021	1/1	0.98	0.04	48,48,48,48	0
57	MG	1A	3630	1/1	0.98	0.04	22,22,22,22	0
57	MG	1A	3143	1/1	0.98	0.05	34,34,34,34	0
57	MG	2A	3100	1/1	0.98	0.10	45,45,45,45	0
57	MG	1A	3144	1/1	0.98	0.05	34,34,34,34	0
57	MG	2W	201	1/1	0.98	0.14	42,42,42,42	0
57	MG	1A	3199	1/1	0.98	0.10	27,27,27,27	0
57	MG	1A	3070	1/1	0.98	0.16	28,28,28,28	0
57	MG	1A	3097	1/1	0.98	0.05	34,34,34,34	0
57	MG	1A	3434	1/1	0.98	0.03	19,19,19,19	0
57	MG	1A	3273	1/1	0.98	0.22	37,37,37,37	0
57	MG	2A	3545	1/1	0.98	0.07	41,41,41,41	0
57	MG	1A	3351	1/1	0.98	0.06	51,51,51,51	0
57	MG	25	101	1/1	0.98	0.20	39,39,39,39	0
57	MG	1a	3031	1/1	0.98	0.15	52,52,52,52	0
57	MG	2A	3548	1/1	0.98	0.05	51,51,51,51	0
57	MG	2A	3321	1/1	0.98	0.05	37,37,37,37	0
57	MG	1A	3050	1/1	0.98	0.19	39,39,39,39	0
57	MG	1a	3033	1/1	0.98	0.03	28,28,28,28	0
57	MG	1A	3029	1/1	0.98	0.07	32,32,32,32	0
57	MG	1B	214	1/1	0.98	0.05	37,37,37,37	0
57	MG	2A	3554	1/1	0.98	0.04	69,69,69,69	0
57	MG	2a	3003	1/1	0.98	0.03	46,46,46,46	0
57	MG	1A	3100	1/1	0.98	0.15	27,27,27,27	0
57	MG	1a	3037	1/1	0.98	0.08	36,36,36,36	0
57	MG	2A	3328	1/1	0.98	0.07	50,50,50,50	0
57	MG	1A	3205	1/1	0.98	0.10	40,40,40,40	0
57	MG	1A	3356	1/1	0.98	0.07	18,18,18,18	0
57	MG	1A	3102	1/1	0.98	0.05	30,30,30,30	0
57	MG	1A	3279	1/1	0.98	0.06	41,41,41,41	0
57	MG	2A	3562	1/1	0.98	0.04	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	3012	1/1	0.98	0.05	48,48,48,48	0
57	MG	1A	3888	1/1	0.98	0.08	27,27,27,27	0
57	MG	2A	3565	1/1	0.98	0.05	51,51,51,51	0
57	MG	1A	3542	1/1	0.98	0.05	29,29,29,29	0
57	MG	1A	3891	1/1	0.98	0.04	46,46,46,46	0
57	MG	2a	3017	1/1	0.98	0.04	55,55,55,55	0
57	MG	1A	3892	1/1	0.98	0.05	32,32,32,32	0
57	MG	1A	3446	1/1	0.98	0.17	31,31,31,31	0
57	MG	2A	3338	1/1	0.98	0.05	33,33,33,33	0
57	MG	1A	3895	1/1	0.98	0.04	28,28,28,28	0
57	MG	2a	3022	1/1	0.98	0.06	50,50,50,50	0
57	MG	2A	3127	1/1	0.98	0.15	34,34,34,34	0
57	MG	1A	3103	1/1	0.98	0.25	27,27,27,27	0
57	MG	1A	3546	1/1	0.98	0.09	27,27,27,27	0
57	MG	1A	3052	1/1	0.98	0.08	27,27,27,27	0
57	MG	1A	3054	1/1	0.98	0.05	23,23,23,23	0
57	MG	2A	3578	1/1	0.98	0.03	50,50,50,50	0
57	MG	1A	3363	1/1	0.98	0.05	10,10,10,10	0
57	MG	1a	3054	1/1	0.98	0.05	40,40,40,40	0
57	MG	1a	3240	1/1	0.98	0.08	51,51,51,51	0
57	MG	1A	3550	1/1	0.98	0.18	46,46,46,46	0
57	MG	1D	302	1/1	0.98	0.05	42,42,42,42	0
57	MG	1A	3451	1/1	0.98	0.06	35,35,35,35	0
57	MG	1a	3058	1/1	0.98	0.21	47,47,47,47	0
57	MG	2A	3139	1/1	0.98	0.17	36,36,36,36	0
57	MG	2A	3353	1/1	0.98	0.05	41,41,41,41	0
57	MG	1A	3906	1/1	0.98	0.06	30,30,30,30	0
57	MG	1D	305	1/1	0.98	0.19	42,42,42,42	0
57	MG	1a	3247	1/1	0.98	0.04	51,51,51,51	0
57	MG	1D	306	1/1	0.98	0.07	20,20,20,20	0
57	MG	1A	3907	1/1	0.98	0.04	14,14,14,14	0
57	MG	2A	3595	1/1	0.98	0.05	40,40,40,40	0
57	MG	1A	3655	1/1	0.98	0.11	44,44,44,44	0
57	MG	2A	3146	1/1	0.98	0.09	61,61,61,61	0
57	MG	2A	3598	1/1	0.98	0.08	39,39,39,39	0
57	MG	2A	3147	1/1	0.98	0.09	54,54,54,54	0
57	MG	1A	3453	1/1	0.98	0.06	21,21,21,21	0
57	MG	1D	310	1/1	0.98	0.11	26,26,26,26	0
57	MG	1A	3774	1/1	0.98	0.04	34,34,34,34	0
57	MG	1A	3076	1/1	0.98	0.23	30,30,30,30	0
57	MG	1A	3912	1/1	0.98	0.06	42,42,42,42	0
57	MG	1A	3455	1/1	0.98	0.06	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	1a	3257	1/1	0.98	0.06	51,51,51,51	0
57	MG	2A	3371	1/1	0.98	0.04	43,43,43,43	0
57	MG	1A	3211	1/1	0.98	0.12	40,40,40,40	0
57	MG	1E	301	1/1	0.98	0.09	28,28,28,28	0
57	MG	1a	3261	1/1	0.98	0.04	51,51,51,51	0
57	MG	1A	3915	1/1	0.98	0.04	47,47,47,47	0
57	MG	1E	303	1/1	0.98	0.05	28,28,28,28	0
57	MG	1A	3778	1/1	0.98	0.04	22,22,22,22	0
57	MG	2A	3161	1/1	0.98	0.17	43,43,43,43	0
57	MG	1A	3366	1/1	0.98	0.06	19,19,19,19	0
57	MG	1A	3367	1/1	0.98	0.13	45,45,45,45	0
57	MG	1A	3919	1/1	0.98	0.05	47,47,47,47	0
57	MG	1A	3155	1/1	0.98	0.24	40,40,40,40	0
57	MG	1A	3214	1/1	0.98	0.05	29,29,29,29	0
57	MG	1A	3110	1/1	0.98	0.10	26,26,26,26	0
57	MG	1A	3561	1/1	0.98	0.10	38,38,38,38	0
57	MG	2A	3624	1/1	0.98	0.04	41,41,41,41	0
57	MG	1A	3925	1/1	0.98	0.05	27,27,27,27	0
57	MG	1F	301	1/1	0.98	0.12	28,28,28,28	0
57	MG	1F	302	1/1	0.98	0.07	25,25,25,25	0
57	MG	2A	3389	1/1	0.98	0.04	37,37,37,37	0
57	MG	1A	3371	1/1	0.98	0.11	58,58,58,58	0
57	MG	1A	3927	1/1	0.98	0.06	47,47,47,47	0
57	MG	2A	3631	1/1	0.98	0.08	58,58,58,58	0
57	MG	1A	3111	1/1	0.98	0.11	37,37,37,37	0
57	MG	1A	3929	1/1	0.98	0.04	42,42,42,42	0
57	MG	1A	3055	1/1	0.98	0.10	38,38,38,38	0
57	MG	2A	3177	1/1	0.98	0.10	32,32,32,32	0
57	MG	1A	3931	1/1	0.98	0.09	60,60,60,60	0
57	MG	1A	3113	1/1	0.98	0.08	21,21,21,21	0
57	MG	1F	313	1/1	0.98	0.07	31,31,31,31	0
57	MG	1A	3670	1/1	0.98	0.07	38,38,38,38	0
57	MG	1f	201	1/1	0.98	0.28	62,62,62,62	0
57	MG	2A	3643	1/1	0.98	0.07	55,55,55,55	0
57	MG	1A	3219	1/1	0.98	0.11	26,26,26,26	0
57	MG	1A	3791	1/1	0.98	0.06	27,27,27,27	0
57	MG	1A	3220	1/1	0.98	0.09	17,17,17,17	0
57	MG	1A	3568	1/1	0.98	0.11	34,34,34,34	0
57	MG	1A	3569	1/1	0.98	0.03	19,19,19,19	0
57	MG	1A	3570	1/1	0.98	0.05	34,34,34,34	0
57	MG	1A	3470	1/1	0.98	0.05	31,31,31,31	0
57	MG	1A	3297	1/1	0.98	0.13	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	2A	3652	1/1	0.98	0.04	36,36,36,36	0
57	MG	2A	3410	1/1	0.98	0.04	38,38,38,38	0
57	MG	2A	3656	1/1	0.98	0.04	38,38,38,38	0
57	MG	1N	201	1/1	0.98	0.05	38,38,38,38	0
57	MG	1N	202	1/1	0.98	0.10	38,38,38,38	0
57	MG	1A	3160	1/1	0.98	0.19	29,29,29,29	0
57	MG	2A	3660	1/1	0.98	0.06	36,36,36,36	0
57	MG	1A	3799	1/1	0.98	0.10	45,45,45,45	0
57	MG	2A	3415	1/1	0.98	0.08	49,49,49,49	0
57	MG	1A	3030	1/1	0.98	0.04	26,26,26,26	0
57	MG	2A	3417	1/1	0.98	0.07	41,41,41,41	0
57	MG	2A	3418	1/1	0.98	0.08	35,35,35,35	0
57	MG	1A	3223	1/1	0.98	0.15	34,34,34,34	0
57	MG	1A	3301	1/1	0.98	0.10	32,32,32,32	0
57	MG	1A	3162	1/1	0.98	0.08	33,33,33,33	0
57	MG	1P	205	1/1	0.98	0.07	27,27,27,27	0
57	MG	1A	3804	1/1	0.98	0.05	39,39,39,39	0
57	MG	2A	3425	1/1	0.98	0.07	35,35,35,35	0
57	MG	1Q	202	1/1	0.98	0.03	29,29,29,29	0
57	MG	1A	3117	1/1	0.98	0.19	27,27,27,27	0
57	MG	2a	3117	1/1	0.98	0.06	68,68,68,68	0
57	MG	2A	3674	1/1	0.98	0.07	61,61,61,61	0
57	MG	1A	3226	1/1	0.98	0.06	37,37,37,37	0
57	MG	1x	101	1/1	0.98	0.04	46,46,46,46	0
57	MG	1A	3686	1/1	0.98	0.04	39,39,39,39	0
57	MG	1A	3580	1/1	0.98	0.08	34,34,34,34	0
57	MG	1R	201	1/1	0.98	0.07	30,30,30,30	0
57	MG	1A	3118	1/1	0.98	0.07	21,21,21,21	0
57	MG	2A	3681	1/1	0.98	0.06	36,36,36,36	0
57	MG	2A	3004	1/1	0.98	0.04	35,35,35,35	0
57	MG	1A	3810	1/1	0.98	0.08	38,38,38,38	0
57	MG	1A	3080	1/1	0.98	0.06	33,33,33,33	0
57	MG	2A	3008	1/1	0.98	0.06	29,29,29,29	0
57	MG	1A	3057	1/1	0.98	0.04	26,26,26,26	0
57	MG	2A	3010	1/1	0.98	0.07	61,61,61,61	0
57	MG	1A	3122	1/1	0.98	0.14	37,37,37,37	0
57	MG	2A	3012	1/1	0.98	0.25	57,57,57,57	0
57	MG	2A	3690	1/1	0.98	0.06	56,56,56,56	0
57	MG	2a	3135	1/1	0.98	0.05	46,46,46,46	0
57	MG	1A	3018	1/1	0.98	0.14	23,23,23,23	0
57	MG	2A	3692	1/1	0.98	0.06	50,50,50,50	0
57	MG	1A	3963	1/1	0.98	0.06	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	3964	1/1	0.98	0.06	28,28,28,28	0
57	MG	1T	205	1/1	0.98	0.05	40,40,40,40	0
57	MG	1A	3815	1/1	0.98	0.04	38,38,38,38	0
57	MG	1A	3586	1/1	0.98	0.07	38,38,38,38	0
57	MG	1A	3967	1/1	0.98	0.08	39,39,39,39	0
57	MG	1A	3124	1/1	0.98	0.08	40,40,40,40	0
57	MG	2A	3021	1/1	0.98	0.04	34,34,34,34	0
57	MG	1V	203	1/1	0.98	0.09	34,34,34,34	0
57	MG	1a	3134	1/1	0.98	0.04	43,43,43,43	0
57	MG	2A	3229	1/1	0.98	0.06	50,50,50,50	0
57	MG	2A	3230	1/1	0.98	0.13	45,45,45,45	0
57	MG	1A	3170	1/1	0.98	0.09	41,41,41,41	0
57	MG	1a	3136	1/1	0.98	0.08	34,34,34,34	0
57	MG	1A	3970	1/1	0.98	0.06	22,22,22,22	0
57	MG	1A	3024	1/1	0.98	0.04	15,15,15,15	0
57	MG	1A	3820	1/1	0.98	0.16	28,28,28,28	0
57	MG	2A	3029	1/1	0.98	0.04	39,39,39,39	0
57	MG	1A	3821	1/1	0.98	0.23	26,26,26,26	0
57	MG	1A	3061	1/1	0.98	0.04	30,30,30,30	0
57	MG	1A	3316	1/1	0.98	0.04	21,21,21,21	0
57	MG	2A	3715	1/1	0.98	0.07	40,40,40,40	0
57	MG	1A	3127	1/1	0.98	0.03	24,24,24,24	0
57	MG	1A	3174	1/1	0.98	0.18	30,30,30,30	0
57	MG	1A	3826	1/1	0.98	0.05	24,24,24,24	0
57	MG	1A	3701	1/1	0.98	0.12	34,34,34,34	0
57	MG	2A	3720	1/1	0.98	0.08	46,46,46,46	0
57	MG	1a	3148	1/1	0.98	0.05	56,56,56,56	0
57	MG	10	102	1/1	0.98	0.04	27,27,27,27	0
57	MG	1A	3400	1/1	0.98	0.07	25,25,25,25	0
57	MG	1A	3239	1/1	0.98	0.09	33,33,33,33	0
57	MG	2A	3725	1/1	0.98	0.11	45,45,45,45	0
57	MG	1A	3982	1/1	0.98	0.08	39,39,39,39	0
57	MG	1A	3175	1/1	0.98	0.14	32,32,32,32	0
57	MG	2A	3474	1/1	0.98	0.06	40,40,40,40	0
57	MG	1A	3403	1/1	0.98	0.08	21,21,21,21	0
57	MG	1a	3155	1/1	0.98	0.06	46,46,46,46	0
57	MG	1A	3985	1/1	0.98	0.05	20,20,20,20	0
57	MG	1A	3833	1/1	0.98	0.04	41,41,41,41	0
57	MG	1A	3241	1/1	0.98	0.16	31,31,31,31	0
57	MG	1A	3176	1/1	0.98	0.12	24,24,24,24	0
57	MG	2A	3256	1/1	0.98	0.19	44,44,44,44	0
57	MG	1A	3085	1/1	0.98	0.33	38,38,38,38	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3013	1/1	0.98	0.05	22,22,22,22	0
57	MG	1A	3246	1/1	0.98	0.20	32,32,32,32	0
57	MG	1A	3180	1/1	0.98	0.15	29,29,29,29	0
57	MG	1A	3006	1/1	0.98	0.07	25,25,25,25	0
57	MG	13	103	1/1	0.98	0.10	34,34,34,34	0
57	MG	1A	3841	1/1	0.98	0.03	22,22,22,22	0
57	MG	1A	3411	1/1	0.98	0.03	33,33,33,33	0
57	MG	15	3103	1/1	0.98	0.14	22,22,22,22	0
57	MG	15	3104	1/1	0.98	0.16	30,30,30,30	0
57	MG	2A	3493	1/1	0.98	0.06	49,49,49,49	0
57	MG	15	3105	1/1	0.98	0.11	33,33,33,33	0
57	MG	15	3106	1/1	0.98	0.04	40,40,40,40	0
57	MG	1A	3131	1/1	0.98	0.15	30,30,30,30	0
57	MG	1A	3717	1/1	0.98	0.08	58,58,58,58	0
57	MG	1A	3044	1/1	0.98	0.09	25,25,25,25	0
57	MG	17	101	1/1	0.98	0.04	32,32,32,32	0
57	MG	17	102	1/1	0.98	0.10	28,28,28,28	0
57	MG	17	103	1/1	0.98	0.14	32,32,32,32	0
57	MG	1A	3252	1/1	0.98	0.41	38,38,38,38	0
57	MG	1A	3509	1/1	0.98	0.04	44,44,44,44	0
57	MG	1A	3185	1/1	0.98	0.12	38,38,38,38	0
57	MG	1A	3612	1/1	0.98	0.21	35,35,35,35	0
57	MG	2A	3071	1/1	0.98	0.11	39,39,39,39	0
57	MG	1A	3725	1/1	0.98	0.07	36,36,36,36	0
57	MG	1A	3089	1/1	0.98	0.09	36,36,36,36	0
57	MG	1A	3001	1/1	0.98	0.04	26,26,26,26	0
57	MG	1A	3514	1/1	0.98	0.04	36,36,36,36	0
60	ZN	14	501	1/1	0.98	0.08	105,105,105,105	0
60	ZN	2Y	202	1/1	0.98	0.04	82,82,82,82	0
57	MG	2E	302	1/1	0.98	0.07	39,39,39,39	0
57	MG	2A	3285	1/1	0.98	0.11	37,37,37,37	0
60	ZN	2n	102	1/1	0.98	0.06	76,76,76,76	0
61	SF4	2d	501	8/8	0.98	0.04	64,74,77,79	0
57	MG	2A	3563	1/1	0.99	0.06	53,53,53,53	0
57	MG	1A	3256	1/1	0.99	0.07	38,38,38,38	0
57	MG	1A	3079	1/1	0.99	0.14	37,37,37,37	0
57	MG	1A	3053	1/1	0.99	0.17	23,23,23,23	0
57	MG	1A	3119	1/1	0.99	0.07	32,32,32,32	0
57	MG	1A	3043	1/1	0.99	0.09	21,21,21,21	0
57	MG	2a	3104	1/1	0.99	0.34	50,50,50,50	0
57	MG	1A	3945	1/1	0.99	0.13	23,23,23,23	0
57	MG	1a	3012	1/1	0.99	0.12	27,27,27,27	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3362	1/1	0.99	0.07	27,27,27,27	0
57	MG	1P	202	1/1	0.99	0.17	23,23,23,23	0
57	MG	1A	3492	1/1	0.99	0.04	25,25,25,25	0
57	MG	2A	3574	1/1	0.99	0.07	31,31,31,31	0
57	MG	1A	3947	1/1	0.99	0.07	45,45,45,45	0
57	MG	1A	3493	1/1	0.99	0.06	24,24,24,24	0
57	MG	1A	3261	1/1	0.99	0.12	26,26,26,26	0
57	MG	1A	3227	1/1	0.99	0.09	30,30,30,30	0
57	MG	2A	3694	1/1	0.99	0.05	49,49,49,49	0
57	MG	1A	3873	1/1	0.99	0.05	30,30,30,30	0
57	MG	2A	3370	1/1	0.99	0.03	35,35,35,35	0
57	MG	1A	3303	1/1	0.99	0.12	30,30,30,30	0
57	MG	1A	3198	1/1	0.99	0.16	36,36,36,36	0
57	MG	1A	3442	1/1	0.99	0.08	27,27,27,27	0
57	MG	1Q	207	1/1	0.99	0.11	45,45,45,45	0
57	MG	1A	3003	1/1	0.99	0.03	28,28,28,28	0
57	MG	1B	213	1/1	0.99	0.06	34,34,34,34	0
57	MG	2A	3276	1/1	0.99	0.18	29,29,29,29	0
57	MG	1A	3068	1/1	0.99	0.18	29,29,29,29	0
57	MG	1A	3445	1/1	0.99	0.03	22,22,22,22	0
57	MG	1A	3958	1/1	0.99	0.03	37,37,37,37	0
57	MG	2A	3592	1/1	0.99	0.07	57,57,57,57	0
57	MG	1A	3045	1/1	0.99	0.10	8,8,8,8	0
57	MG	2A	3184	1/1	0.99	0.02	41,41,41,41	0
57	MG	1A	3101	1/1	0.99	0.03	45,45,45,45	0
57	MG	1A	3017	1/1	0.99	0.19	27,27,27,27	0
57	MG	1A	3399	1/1	0.99	0.08	15,15,15,15	0
57	MG	1A	3623	1/1	0.99	0.06	30,30,30,30	0
57	MG	1A	3749	1/1	0.99	0.06	26,26,26,26	0
57	MG	1A	3750	1/1	0.99	0.09	29,29,29,29	0
57	MG	2A	3492	1/1	0.99	0.05	35,35,35,35	0
57	MG	1A	3684	1/1	0.99	0.06	34,34,34,34	0
57	MG	1U	204	1/1	0.99	0.17	32,32,32,32	0
57	MG	1a	3127	1/1	0.99	0.04	32,32,32,32	0
57	MG	2A	3496	1/1	0.99	0.06	47,47,47,47	0
57	MG	2A	3005	1/1	0.99	0.05	62,62,62,62	0
57	MG	1V	201	1/1	0.99	0.14	27,27,27,27	0
57	MG	2A	3608	1/1	0.99	0.03	40,40,40,40	0
57	MG	1A	3058	1/1	0.99	0.17	42,42,42,42	0
57	MG	2A	3610	1/1	0.99	0.04	25,25,25,25	0
57	MG	1A	3104	1/1	0.99	0.08	35,35,35,35	0
57	MG	2A	3102	1/1	0.99	0.11	55,55,55,55	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	1A	3890	1/1	0.99	0.03	21,21,21,21	0
57	MG	1A	3452	1/1	0.99	0.07	22,22,22,22	0
57	MG	1A	3178	1/1	0.99	0.23	28,28,28,28	0
57	MG	1A	3272	1/1	0.99	0.17	31,31,31,31	0
57	MG	2A	3506	1/1	0.99	0.04	44,44,44,44	0
57	MG	2A	3107	1/1	0.99	0.04	30,30,30,30	0
57	MG	1A	3894	1/1	0.99	0.03	36,36,36,36	0
57	MG	1A	3511	1/1	0.99	0.07	18,18,18,18	0
57	MG	1A	3047	1/1	0.99	0.04	24,24,24,24	0
57	MG	1A	3897	1/1	0.99	0.03	48,48,48,48	0
57	MG	1A	3315	1/1	0.99	0.09	29,29,29,29	0
57	MG	1a	3050	1/1	0.99	0.07	46,46,46,46	0
57	MG	1A	3899	1/1	0.99	0.07	49,49,49,49	0
57	MG	1A	3073	1/1	0.99	0.04	33,33,33,33	0
57	MG	1A	3360	1/1	0.99	0.03	19,19,19,19	0
57	MG	1A	3181	1/1	0.99	0.11	26,26,26,26	0
57	MG	1A	3763	1/1	0.99	0.04	31,31,31,31	0
57	MG	1a	3146	1/1	0.99	0.05	52,52,52,52	0
57	MG	1A	3904	1/1	0.99	0.03	49,49,49,49	0
57	MG	1A	3831	1/1	0.99	0.05	46,46,46,46	0
57	MG	1A	3107	1/1	0.99	0.08	35,35,35,35	0
57	MG	1A	3007	1/1	0.99	0.06	33,33,33,33	0
57	MG	1A	3242	1/1	0.99	0.17	21,21,21,21	0
57	MG	2A	3318	1/1	0.99	0.08	17,17,17,17	0
57	MG	2A	3637	1/1	0.99	0.08	33,33,33,33	0
57	MG	2A	3638	1/1	0.99	0.07	30,30,30,30	0
57	MG	2A	3319	1/1	0.99	0.05	57,57,57,57	0
57	MG	1A	3212	1/1	0.99	0.05	39,39,39,39	0
57	MG	2A	3423	1/1	0.99	0.05	37,37,37,37	0
57	MG	1A	3413	1/1	0.99	0.02	26,26,26,26	0
57	MG	1A	3109	1/1	0.99	0.18	27,27,27,27	0
57	MG	1A	3281	1/1	0.99	0.06	32,32,32,32	0
57	MG	1A	3467	1/1	0.99	0.04	32,32,32,32	0
57	MG	1A	3019	1/1	0.99	0.15	23,23,23,23	0
57	MG	1A	3469	1/1	0.99	0.05	40,40,40,40	0
57	MG	1A	3026	1/1	0.99	0.11	29,29,29,29	0
57	MG	1A	3708	1/1	0.99	0.05	21,21,21,21	0
57	MG	1A	3284	1/1	0.99	0.04	23,23,23,23	0
57	MG	1A	3285	1/1	0.99	0.09	31,31,31,31	0
57	MG	1A	3247	1/1	0.99	0.14	28,28,28,28	0
57	MG	2A	3653	1/1	0.99	0.04	34,34,34,34	0
57	MG	1A	3005	1/1	0.99	0.09	22,22,22,22	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
57	MG	2A	3655	1/1	0.99	0.04	69,69,69,69	0
57	MG	1A	3136	1/1	0.99	0.12	24,24,24,24	0
57	MG	1A	3923	1/1	0.99	0.03	46,46,46,46	0
57	MG	1a	3259	1/1	0.99	0.06	33,33,33,33	0
57	MG	1F	304	1/1	0.99	0.10	28,28,28,28	0
57	MG	1A	3476	1/1	0.99	0.08	42,42,42,42	0
57	MG	1F	306	1/1	0.99	0.06	35,35,35,35	0
57	MG	1a	3079	1/1	0.99	0.16	43,43,43,43	0
57	MG	1A	3093	1/1	0.99	0.03	31,31,31,31	0
57	MG	1A	3376	1/1	0.99	0.04	30,30,30,30	0
57	MG	1A	3015	1/1	0.99	0.04	26,26,26,26	0
57	MG	1A	3115	1/1	0.99	0.04	28,28,28,28	0
57	MG	1A	3596	1/1	0.99	0.06	41,41,41,41	0
57	MG	1A	3116	1/1	0.99	0.14	31,31,31,31	0
57	MG	1A	3721	1/1	0.99	0.05	22,22,22,22	0
57	MG	1A	3254	1/1	0.99	0.11	34,34,34,34	0
57	MG	1A	3336	1/1	0.99	0.04	30,30,30,30	0
60	ZN	1Y	202	1/1	0.99	0.03	53,53,53,53	0
57	MG	1A	3724	1/1	0.99	0.03	35,35,35,35	0
60	ZN	15	3110	1/1	0.99	0.02	43,43,43,43	0
60	ZN	16	501	1/1	0.99	0.05	40,40,40,40	0
60	ZN	19	103	1/1	0.99	0.04	41,41,41,41	0
60	ZN	1n	104	1/1	0.99	0.03	65,65,65,65	0
57	MG	1A	3294	1/1	0.99	0.04	20,20,20,20	0
57	MG	1A	3383	1/1	0.99	0.09	42,42,42,42	0
60	ZN	25	104	1/1	0.99	0.02	49,49,49,49	0
57	MG	1A	3937	1/1	0.99	0.06	35,35,35,35	0
57	MG	1A	3543	1/1	0.99	0.03	30,30,30,30	0
61	SF4	1d	307	8/8	0.99	0.04	54,58,64,66	0
57	MG	1A	3141	1/1	0.99	0.08	21,21,21,21	0
57	MG	1A	3704	1/1	1.00	0.03	31,31,31,31	0
57	MG	1a	3196	1/1	1.00	0.06	52,52,52,52	0
60	ZN	26	501	1/1	1.00	0.04	59,59,59,59	0
57	MG	1F	309	1/1	1.00	0.04	33,33,33,33	0
57	MG	2A	3590	1/1	1.00	0.03	32,32,32,32	0
57	MG	2A	3390	1/1	1.00	0.04	44,44,44,44	0
57	MG	1a	3270	1/1	1.00	0.03	56,56,56,56	0

5.5 Other polymers ⓘ

There are no such residues in this entry.