



Full wwPDB X-ray Structure Validation Report ⓘ

Mar 31, 2026 – 02:34 AM UTC

PDB ID : 5HCP / pdb_00005hcp
Title : Crystal structure of antimicrobial peptide Metalnikowin bound to the *Thermus thermophilus* 70S ribosome
Authors : Gagnon, M.G.; Roy, R.N.; Lomakin, I.B.; Florin, T.; Mankin, A.S.; Steitz, T.A.
Deposited on : 2016-01-04
Resolution : 2.89 Å(reported)

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

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

A user guide is available at

<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

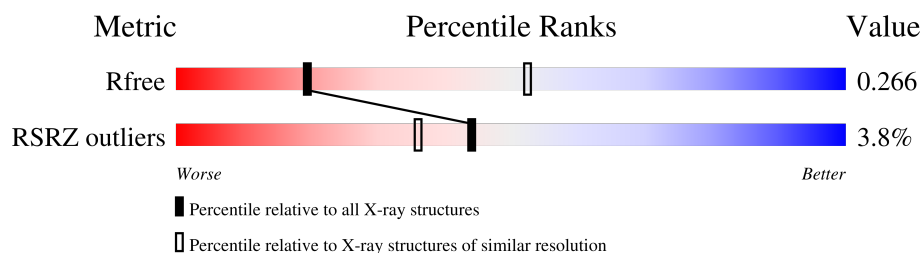
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

The reported resolution of this entry is 2.89 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	2481 (2.90-2.90)
RSRZ outliers	180081	2481 (2.90-2.90)

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

2 Entry composition [i](#)

There are 59 unique types of molecules in this entry. The entry contains 288976 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	2746	Total	C	N	O	P	0	0	0
			59154	26327	11077	19005	2745			
1	2A	2790	Total	C	N	O	P	0	0	0
			60091	26746	11243	19313	2789			

- 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	1146	476	831	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
			2136	1349	423	361	3			
3	2D	275	Total	C	N	O	S	0	0	0
			2142	1352	426	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
			1425	914	256	251	4			
6	2G	181	Total	C	N	O	S	0	0	0
			1424	911	258	251	4			

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

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

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
8	1I	146	Total	C	N	O	S	0	0	0
			1085	693	189	202	1			
8	2I	146	Total	C	N	O	S	0	0	0
			1061	680	186	194	1			

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

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

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

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

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

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
11	1P	149	Total	C	N	O	S	0	0	0
			1139	709	231	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
			771	495	140	135	1			
17	2V	101	Total	C	N	O	S	0	0	0
			771	495	140	135	1			

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

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

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

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

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

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

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
21	1Z	186	Total	C	N	O	S	0	0	0
			1470	937	262	269	2			
21	2Z	186	Total	C	N	O	S	0	0	0
			1454	929	256	267	2			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
22	10	75	Total	C	N	O	S	0	0	0
			598	370	127	100	1			
22	20	75	Total	C	N	O	S	0	0	0
			598	370	127	100	1			

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

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

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

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

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

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

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
26	14	69	Total	C	N	O	S	0	0	0
			558	352	102	99	5			

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

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

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

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

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

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

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

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

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

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

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

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
32	1a	1477	Total	C	N	O	P	0	0	0
			31750	14131	5883	10259	1477			
32	2a	1483	Total	C	N	O	P	0	0	0
			31877	14188	5905	10301	1483			

- 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
			1786	1136	321	325	4			
33	2b	231	Total	C	N	O	S	0	0	0
			1697	1079	292	321	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
			1480	932	281	266	1			
34	2c	206	Total	C	N	O	S	0	0	0
			1412	883	269	259	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
			1618	1013	312	286	7			
35	2d	208	Total	C	N	O	S	0	0	0
			1630	1022	321	280	7			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
36	1e	148	Total	C	N	O	S	0	0	0
			1129	714	213	198	4			
36	2e	148	Total	C	N	O	S	0	0	0
			1095	695	203	193	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
			806	511	143	149	3			
37	2f	100	Total	C	N	O	S	0	0	0
			817	516	146	152	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
			1183	732	232	213	6			
38	2g	155	Total	C	N	O	S	0	0	0
			1167	728	220	213	6			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
39	1h	137	Total	C	N	O	S	0	0	0
			1088	689	206	191	2			
39	2h	137	Total	C	N	O	S	0	0	0
			1074	681	202	189	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
			976	620	189	167			
40	2i	127	Total	C	N	O	0	0	0
			932	589	177	166			

- 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
			682	424	130	128			
41	2j	96	Total	C	N	O	0	0	0
			678	424	126	128			

- 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
			826	513	156	154	3			

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Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
42	2k	114	Total	C	N	O	S	0	0	0
			829	516	155	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
			920	579	181	159	1			
43	2l	122	Total	C	N	O	S	0	0	0
			918	576	182	159	1			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
44	1m	118	Total	C	N	O	S	0	0	0
			923	569	191	161	2			
44	2m	116	Total	C	N	O	S	0	0	0
			903	555	187	159	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
			482	306	100	72	4			
45	2n	60	Total	C	N	O	S	0	0	0
			459	291	93	71	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
			715	447	140	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
			671	424	133	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
			811	519	148	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	84	Total	C	N	O	S	0	0	0
			642	409	119	112	2			
50	2s	83	Total	C	N	O	S	0	0	0
			646	412	119	113	2			

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

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
51	1t	96	Total	C	N	O	S	0	0	0
			712	435	152	123	2			
51	2t	96	Total	C	N	O	S	0	0	0
			731	449	156	124	2			

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

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
52	1u	23	Total	C	N	O	0	0	0
			187	116	42	29			
52	2u	23	Total	C	N	O	0	0	0
			199	122	48	29			

- Molecule 53 is a RNA chain called mRNA.

Mol	Chain	Residues	Atoms					ZeroOcc	AltConf	Trace
53	1v	5	Total	C	N	O	P	0	0	0
			109	49	22	33	5			
53	2v	5	Total	C	N	O	P	0	0	0
			109	49	22	33	5			

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

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

- Molecule 55 is a protein called Metalnikowin-1.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
55	1z	12	Total	C	N	O	0	0	0
			105	66	22	17			
55	2z	13	Total	C	N	O	0	0	0
			112	71	23	18			

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

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	1A	1090	Total	Mg	0	0
			1090	1090		
56	1B	30	Total	Mg	0	0
			30	30		
56	1D	13	Total	Mg	0	0
			13	13		
56	1E	12	Total	Mg	0	0
			12	12		
56	1F	8	Total	Mg	0	0
			8	8		
56	1G	3	Total	Mg	0	0
			3	3		
56	1H	4	Total	Mg	0	0
			4	4		
56	1N	7	Total	Mg	0	0
			7	7		
56	1O	3	Total	Mg	0	0
			3	3		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	1P	6	Total 6	Mg 6	0	0
56	1Q	9	Total 9	Mg 9	0	0
56	1R	6	Total 6	Mg 6	0	0
56	1T	6	Total 6	Mg 6	0	0
56	1U	6	Total 6	Mg 6	0	0
56	1V	2	Total 2	Mg 2	0	0
56	1W	6	Total 6	Mg 6	0	0
56	1X	1	Total 1	Mg 1	0	0
56	1Y	1	Total 1	Mg 1	0	0
56	1Z	2	Total 2	Mg 2	0	0
56	10	6	Total 6	Mg 6	0	0
56	11	3	Total 3	Mg 3	0	0
56	12	1	Total 1	Mg 1	0	0
56	13	2	Total 2	Mg 2	0	0
56	14	1	Total 1	Mg 1	0	0
56	15	3	Total 3	Mg 3	0	0
56	16	4	Total 4	Mg 4	0	0
56	17	2	Total 2	Mg 2	0	0
56	18	3	Total 3	Mg 3	0	0
56	19	5	Total 5	Mg 5	0	0
56	1a	309	Total 309	Mg 309	0	0

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Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
56	1b	1	Total Mg 1 1	0	0
56	1d	2	Total Mg 2 2	0	0
56	1e	3	Total Mg 3 3	0	0
56	1f	1	Total Mg 1 1	0	0
56	1h	2	Total Mg 2 2	0	0
56	1k	1	Total Mg 1 1	0	0
56	1l	2	Total Mg 2 2	0	0
56	1m	1	Total Mg 1 1	0	0
56	1n	1	Total Mg 1 1	0	0
56	1o	3	Total Mg 3 3	0	0
56	1p	2	Total Mg 2 2	0	0
56	1q	3	Total Mg 3 3	0	0
56	1r	3	Total Mg 3 3	0	0
56	1t	1	Total Mg 1 1	0	0
56	1u	1	Total Mg 1 1	0	0
56	1v	1	Total Mg 1 1	0	0
56	1x	10	Total Mg 10 10	0	0
56	1z	1	Total Mg 1 1	0	0
56	2A	561	Total Mg 561 561	0	0
56	2B	10	Total Mg 10 10	0	0
56	2D	5	Total Mg 5 5	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	2E	4	Total 4	Mg 4	0	0
56	2F	4	Total 4	Mg 4	0	0
56	2G	1	Total 1	Mg 1	0	0
56	2N	1	Total 1	Mg 1	0	0
56	2O	3	Total 3	Mg 3	0	0
56	2P	1	Total 1	Mg 1	0	0
56	2Q	5	Total 5	Mg 5	0	0
56	2R	2	Total 2	Mg 2	0	0
56	2T	1	Total 1	Mg 1	0	0
56	2U	1	Total 1	Mg 1	0	0
56	2X	1	Total 1	Mg 1	0	0
56	2Y	1	Total 1	Mg 1	0	0
56	2Z	1	Total 1	Mg 1	0	0
56	20	1	Total 1	Mg 1	0	0
56	23	1	Total 1	Mg 1	0	0
56	25	1	Total 1	Mg 1	0	0
56	28	3	Total 3	Mg 3	0	0
56	2a	255	Total 255	Mg 255	0	0
56	2d	2	Total 2	Mg 2	0	0
56	2e	2	Total 2	Mg 2	0	0
56	2f	1	Total 1	Mg 1	0	0

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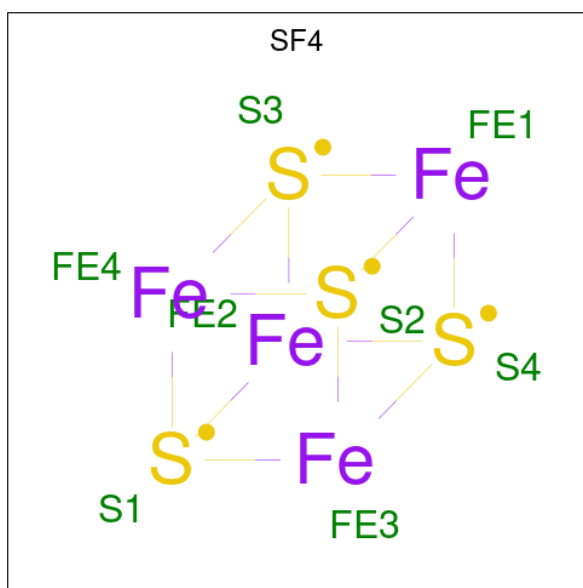
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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	2l	5	Total 5	Mg 5	0	0
56	2q	3	Total 3	Mg 3	0	0
56	2r	1	Total 1	Mg 1	0	0
56	2t	1	Total 1	Mg 1	0	0
56	2v	1	Total 1	Mg 1	0	0
56	2x	7	Total 7	Mg 7	0	0
56	2z	1	Total 1	Mg 1	0	0

- Molecule 57 is ZINC ION (CCD ID: ZN) (formula: Zn).

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

- Molecule 58 is IRON/SULFUR CLUSTER (CCD ID: SF4) (formula: Fe₄S₄).



Mol	Chain	Residues	Atoms			ZeroOcc	AltConf
58	1d	1	Total	Fe	S	0	0
			8	4	4		
58	2d	1	Total	Fe	S	0	0
			8	4	4		

- Molecule 59 is water.

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	1A	2412	Total	O	0	0
			2412	2412		
59	1B	46	Total	O	0	0
			46	46		
59	1D	25	Total	O	0	0
			25	25		
59	1E	30	Total	O	0	0
			30	30		
59	1F	21	Total	O	0	0
			21	21		
59	1G	8	Total	O	0	0
			8	8		
59	1H	4	Total	O	0	0
			4	4		
59	1I	3	Total	O	0	0
			3	3		
59	1N	8	Total	O	0	0
			8	8		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	1O	5	Total 5	O 5	0	0
59	1P	25	Total 25	O 25	0	0
59	1Q	13	Total 13	O 13	0	0
59	1R	18	Total 18	O 18	0	0
59	1S	2	Total 2	O 2	0	0
59	1T	13	Total 13	O 13	0	0
59	1U	18	Total 18	O 18	0	0
59	1V	5	Total 5	O 5	0	0
59	1W	14	Total 14	O 14	0	0
59	1X	5	Total 5	O 5	0	0
59	1Y	1	Total 1	O 1	0	0
59	1Z	6	Total 6	O 6	0	0
59	10	8	Total 8	O 8	0	0
59	11	7	Total 7	O 7	0	0
59	12	2	Total 2	O 2	0	0
59	13	3	Total 3	O 3	0	0
59	15	7	Total 7	O 7	0	0
59	16	11	Total 11	O 11	0	0
59	17	9	Total 9	O 9	0	0
59	18	16	Total 16	O 16	0	0
59	19	3	Total 3	O 3	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	1a	394	Total 394	O 394	0	0
59	1b	1	Total 1	O 1	0	0
59	1c	1	Total 1	O 1	0	0
59	1d	3	Total 3	O 3	0	0
59	1e	4	Total 4	O 4	0	0
59	1f	1	Total 1	O 1	0	0
59	1h	2	Total 2	O 2	0	0
59	1j	1	Total 1	O 1	0	0
59	1k	2	Total 2	O 2	0	0
59	1l	3	Total 3	O 3	0	0
59	1m	1	Total 1	O 1	0	0
59	1p	1	Total 1	O 1	0	0
59	1q	1	Total 1	O 1	0	0
59	1s	1	Total 1	O 1	0	0
59	1u	1	Total 1	O 1	0	0
59	1v	3	Total 3	O 3	0	0
59	1x	10	Total 10	O 10	0	0
59	1z	5	Total 5	O 5	0	0
59	2A	824	Total 824	O 824	0	0
59	2B	9	Total 9	O 9	0	0
59	2D	18	Total 18	O 18	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	2E	11	Total 11	O 11	0	0
59	2F	6	Total 6	O 6	0	0
59	2N	2	Total 2	O 2	0	0
59	2O	1	Total 1	O 1	0	0
59	2P	6	Total 6	O 6	0	0
59	2R	1	Total 1	O 1	0	0
59	2T	1	Total 1	O 1	0	0
59	2U	3	Total 3	O 3	0	0
59	2W	1	Total 1	O 1	0	0
59	2X	3	Total 3	O 3	0	0
59	2Y	2	Total 2	O 2	0	0
59	2Z	5	Total 5	O 5	0	0
59	20	2	Total 2	O 2	0	0
59	21	1	Total 1	O 1	0	0
59	23	2	Total 2	O 2	0	0
59	25	1	Total 1	O 1	0	0
59	27	1	Total 1	O 1	0	0
59	28	4	Total 4	O 4	0	0
59	2a	329	Total 329	O 329	0	0
59	2c	1	Total 1	O 1	0	0
59	2d	1	Total 1	O 1	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	2e	4	Total 4	O 4	0	0
59	2g	1	Total 1	O 1	0	0
59	2i	2	Total 2	O 2	0	0
59	2l	2	Total 2	O 2	0	0
59	2m	1	Total 1	O 1	0	0
59	2o	1	Total 1	O 1	0	0
59	2p	2	Total 2	O 2	0	0
59	2q	1	Total 1	O 1	0	0
59	2t	4	Total 4	O 4	0	0
59	2u	1	Total 1	O 1	0	0
59	2x	6	Total 6	O 6	0	0
59	2z	2	Total 2	O 2	0	0

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	209.87Å 450.49Å 623.11Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	49.85 – 2.89 49.85 – 2.89	Depositor EDS
% Data completeness (in resolution range)	98.6 (49.85-2.89) 98.6 (49.85-2.89)	Depositor EDS
R_{merge}	0.22	Depositor
R_{sym}	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ ¹	1.25 (at 2.91Å)	Xtriage
Refinement program	PHENIX	Depositor
R, R_{free}	0.212 , 0.268 0.212 , 0.266	Depositor DCC
R_{free} test set	64043 reflections (4.94%)	wwPDB-VP
Wilson B-factor (Å ²)	56.9	Xtriage
Anisotropy	0.146	Xtriage
Bulk solvent k_{sol} (e/Å ³), B_{sol} (Å ²)	0.29 , 61.9	EDS
L-test for twinning ²	$\langle L \rangle = 0.42$, $\langle L^2 \rangle = 0.24$	Xtriage
Estimated twinning fraction	No twinning to report.	Xtriage
F_o, F_c correlation	0.91	EDS
Total number of atoms	288976	wwPDB-VP
Average B, all atoms (Å ²)	59.0	wwPDB-VP

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 1.69% 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](#)

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

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

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
54	5MU	2x	54	54	19,22,23	1.43	5 (26%)	27,32,35	1.98	6 (22%)
54	5MC	2x	32	54	19,22,23	1.70	2 (10%)	26,32,35	1.27	3 (11%)
54	5MU	1x	54	54	19,22,23	1.54	5 (26%)	27,32,35	1.90	5 (18%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
54	PSU	2x	55	54	18,21,22	1.36	2 (11%)	21,30,33	1.95	4 (19%)
54	5MC	1x	32	54	19,22,23	1.62	3 (15%)	26,32,35	1.16	2 (7%)
54	4SU	2x	8	54	18,21,22	2.00	5 (27%)	25,30,33	1.25	4 (16%)
54	PSU	1x	55	54	18,21,22	1.33	2 (11%)	21,30,33	1.87	3 (14%)
54	4SU	1x	8	54	18,21,22	2.13	4 (22%)	25,30,33	1.34	4 (16%)

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

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
54	5MU	2x	54	54	-	1/7/25/26	0/2/2/2
54	5MC	2x	32	54	-	1/7/25/26	0/2/2/2
54	5MU	1x	54	54	-	0/7/25/26	0/2/2/2
54	PSU	2x	55	54	-	2/7/25/26	0/2/2/2
54	5MC	1x	32	54	-	0/7/25/26	0/2/2/2
54	4SU	2x	8	54	-	0/7/25/26	0/2/2/2
54	PSU	1x	55	54	-	2/7/25/26	0/2/2/2
54	4SU	1x	8	54	-	0/7/25/26	0/2/2/2

All (28) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
54	2x	32	5MC	C5-C4	5.99	1.48	1.44
54	1x	32	5MC	C5-C4	5.62	1.48	1.44
54	1x	8	4SU	C4-N3	-4.92	1.32	1.37
54	2x	8	4SU	C4-N3	-4.73	1.32	1.37
54	1x	8	4SU	C4-S4	-4.44	1.60	1.68
54	2x	8	4SU	C4-S4	-4.28	1.61	1.68
54	1x	8	4SU	C2-N3	-4.17	1.30	1.38
54	2x	55	PSU	C6-C5	3.65	1.39	1.35
54	1x	54	5MU	C6-C5	3.37	1.40	1.34
54	1x	55	PSU	C6-C5	3.13	1.38	1.35
54	2x	32	5MC	C6-C5	3.11	1.39	1.34
54	1x	32	5MC	C6-C5	3.09	1.39	1.34
54	2x	8	4SU	C5-C4	-3.01	1.38	1.42
54	2x	8	4SU	C2-N3	-3.01	1.32	1.38
54	1x	8	4SU	C5-C4	-2.97	1.39	1.42
54	1x	54	5MU	C4-N3	-2.93	1.33	1.38
54	2x	54	5MU	C6-C5	2.91	1.39	1.34

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
54	1x	54	5MU	C2-N1	2.65	1.42	1.38
54	1x	54	5MU	C4-C5	2.59	1.49	1.44
54	2x	54	5MU	C4-C5	2.57	1.49	1.44
54	1x	55	PSU	C4-N3	-2.44	1.34	1.38
54	2x	54	5MU	C4-N3	-2.42	1.34	1.38
54	2x	55	PSU	C4-N3	-2.40	1.34	1.38
54	2x	54	5MU	C2-N1	2.38	1.42	1.38
54	1x	32	5MC	C6-N1	-2.23	1.34	1.38
54	2x	8	4SU	C6-C5	2.08	1.39	1.35
54	2x	54	5MU	C6-N1	-2.06	1.34	1.38
54	1x	54	5MU	C2-N3	-2.05	1.34	1.38

All (31) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
54	2x	55	PSU	N1-C2-N3	6.07	121.57	115.17
54	1x	55	PSU	N1-C2-N3	5.52	120.99	115.17
54	1x	54	5MU	N3-C2-N1	5.25	121.72	114.89
54	2x	54	5MU	N3-C2-N1	4.82	121.16	114.89
54	2x	54	5MU	C4-N3-C2	-4.78	121.07	127.34
54	1x	54	5MU	C4-N3-C2	-4.45	121.51	127.34
54	2x	54	5MU	C5-C4-N3	3.96	118.77	115.32
54	2x	55	PSU	C4-N3-C2	-3.92	120.97	126.37
54	1x	54	5MU	C5-C4-N3	3.71	118.55	115.32
54	1x	55	PSU	C4-N3-C2	-3.69	121.29	126.37
54	1x	55	PSU	O2-C2-N1	-3.59	119.08	122.79
54	2x	54	5MU	O4-C4-C5	-3.53	120.88	124.92
54	2x	32	5MC	O2-C2-N3	-3.35	117.04	122.33
54	2x	54	5MU	C5-C6-N1	-3.31	119.72	123.31
54	1x	32	5MC	C5-C6-N1	-3.21	119.82	123.31
54	1x	54	5MU	C5-C6-N1	-3.18	119.86	123.31
54	1x	8	4SU	C6-C5-C4	-3.10	117.27	119.95
54	2x	55	PSU	O2-C2-N1	-3.09	119.61	122.79
54	1x	8	4SU	O2-C2-N1	3.06	126.78	122.80
54	2x	32	5MC	C5-C4-N3	-3.05	118.63	121.75
54	1x	32	5MC	C5-C4-N3	-3.00	118.68	121.75
54	2x	32	5MC	C5-C6-N1	-2.94	120.12	123.31
54	1x	8	4SU	C5-C4-N3	2.91	117.46	114.75
54	2x	8	4SU	C6-C5-C4	-2.82	117.51	119.95
54	2x	8	4SU	C1'-N1-C2	2.70	122.43	117.59
54	2x	8	4SU	C5-C4-N3	2.56	117.13	114.75
54	1x	54	5MU	O4-C4-C5	-2.55	122.00	124.92

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
54	2x	54	5MU	O2-C2-N1	-2.44	119.62	122.80
54	1x	8	4SU	O2-C2-N3	-2.22	117.39	121.49
54	2x	55	PSU	C5-C6-N1	-2.16	119.14	122.14
54	2x	8	4SU	C6-N1-C2	-2.05	118.50	121.00

There are no chirality outliers.

All (6) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
54	1x	55	PSU	O4'-C4'-C5'-O5'
54	2x	55	PSU	O4'-C4'-C5'-O5'
54	2x	54	5MU	C3'-C4'-C5'-O5'
54	1x	55	PSU	C3'-C4'-C5'-O5'
54	2x	32	5MC	C2'-C1'-N1-C2
54	2x	55	PSU	C3'-C4'-C5'-O5'

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 2494 ligands modelled in this entry, 2492 are monoatomic - leaving 2 for Mogul analysis.

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

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	$\# Z > 2$	Counts	RMSZ	$\# Z > 2$
58	SF4	1d	501	35	0,12,12	-	-	-		
58	SF4	2d	501	35	0,12,12	-	-	-		

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

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

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

There are no torsion outliers.

There are no ring outliers.

No monomer is involved in short contacts.

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	2746/2915 (94%)	-0.38	28 (1%) 79 73	22, 41, 87, 114	0
1	2A	2790/2915 (95%)	-0.28	22 (0%) 82 77	25, 46, 93, 111	0
2	1B	120/121 (99%)	0.15	1 (0%) 82 77	36, 65, 80, 91	0
2	2B	120/121 (99%)	0.65	2 (1%) 69 61	41, 70, 83, 92	0
3	1D	275/276 (99%)	-0.21	0 100 100	23, 40, 57, 77	0
3	2D	275/276 (99%)	-0.17	2 (0%) 84 79	24, 43, 59, 78	0
4	1E	204/206 (99%)	-0.12	0 100 100	21, 43, 66, 81	0
4	2E	204/206 (99%)	0.01	0 100 100	23, 45, 67, 82	0
5	1F	203/210 (96%)	0.09	1 (0%) 87 83	22, 50, 75, 90	0
5	2F	203/210 (96%)	0.14	1 (0%) 87 83	24, 54, 76, 93	0
6	1G	181/182 (99%)	0.56	6 (3%) 49 40	50, 70, 83, 98	0
6	2G	181/182 (99%)	1.09	14 (7%) 19 16	54, 73, 85, 100	0
7	1H	174/180 (96%)	0.64	4 (2%) 61 52	48, 64, 77, 82	0
7	2H	174/180 (96%)	0.88	8 (4%) 37 29	56, 69, 81, 83	0
8	1I	146/148 (98%)	0.55	3 (2%) 63 54	45, 75, 84, 89	0
8	2I	146/148 (98%)	0.69	6 (4%) 41 33	47, 76, 85, 92	0
9	1N	140/140 (100%)	0.18	0 100 100	27, 46, 66, 79	0
9	2N	140/140 (100%)	0.16	0 100 100	30, 50, 70, 81	0
10	1O	122/122 (100%)	-0.00	0 100 100	33, 45, 63, 70	0
10	2O	122/122 (100%)	-0.06	0 100 100	34, 47, 63, 73	0
11	1P	149/150 (99%)	0.23	0 100 100	22, 52, 75, 86	0
11	2P	149/150 (99%)	0.30	2 (1%) 75 67	25, 56, 78, 89	0
12	1Q	141/141 (100%)	0.33	4 (2%) 55 46	31, 49, 63, 87	0
12	2Q	141/141 (100%)	0.36	2 (1%) 73 65	35, 52, 66, 85	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
13	1R	118/118 (100%)	-0.39	0 100 100	19, 30, 47, 57	0
13	2R	118/118 (100%)	0.14	1 (0%) 82 77	35, 51, 65, 76	0
14	1S	110/112 (98%)	-0.06	1 (0%) 81 75	33, 49, 65, 68	0
14	2S	110/112 (98%)	1.33	23 (20%) 2 2	65, 79, 88, 93	0
15	1T	131/146 (89%)	-0.10	3 (2%) 61 52	29, 41, 70, 90	0
15	2T	131/146 (89%)	0.22	0 100 100	47, 58, 78, 90	0
16	1U	116/118 (98%)	-0.58	1 (0%) 81 75	19, 27, 44, 64	0
16	2U	116/118 (98%)	0.34	1 (0%) 81 75	37, 59, 77, 88	0
17	1V	101/101 (100%)	0.08	1 (0%) 79 73	27, 49, 65, 74	0
17	2V	101/101 (100%)	0.20	2 (1%) 65 56	30, 54, 71, 78	0
18	1W	112/113 (99%)	-0.25	0 100 100	23, 35, 55, 84	0
18	2W	112/113 (99%)	-0.20	1 (0%) 81 75	26, 37, 56, 87	0
19	1X	95/96 (98%)	0.29	3 (3%) 50 41	31, 46, 66, 78	0
19	2X	95/96 (98%)	0.44	4 (4%) 40 32	34, 49, 69, 78	0
20	1Y	107/110 (97%)	-0.07	1 (0%) 81 75	29, 47, 68, 85	0
20	2Y	107/110 (97%)	0.88	7 (6%) 25 20	55, 72, 83, 98	0
21	1Z	186/206 (90%)	0.53	1 (0%) 87 83	51, 68, 82, 92	0
21	2Z	186/206 (90%)	0.91	3 (1%) 70 62	55, 71, 84, 93	0
22	10	75/85 (88%)	0.25	1 (1%) 75 67	33, 45, 59, 66	0
22	20	75/85 (88%)	0.51	4 (5%) 32 25	37, 50, 64, 68	0
23	11	97/98 (98%)	0.13	1 (1%) 79 73	28, 47, 71, 78	0
23	21	97/98 (98%)	0.25	3 (3%) 51 42	30, 49, 73, 78	0
24	12	70/72 (97%)	0.47	2 (2%) 53 45	42, 58, 72, 80	0
24	22	70/72 (97%)	0.48	0 100 100	45, 61, 74, 79	0
25	13	59/60 (98%)	0.21	1 (1%) 69 61	32, 45, 65, 78	0
25	23	59/60 (98%)	0.13	1 (1%) 69 61	35, 50, 69, 80	0
26	14	69/71 (97%)	0.66	7 (10%) 12 10	52, 75, 99, 102	0
26	24	69/71 (97%)	1.33	11 (15%) 5 4	78, 93, 98, 103	0
27	15	59/60 (98%)	-0.58	0 100 100	15, 30, 45, 72	0
27	25	59/60 (98%)	-0.09	0 100 100	33, 50, 67, 72	0
28	16	53/54 (98%)	-0.08	0 100 100	38, 48, 59, 70	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
28	26	53/54 (98%)	0.01	0 100 100	42, 51, 62, 72	0
29	17	48/49 (97%)	-0.07	2 (4%) 40 32	24, 31, 64, 81	0
29	27	48/49 (97%)	0.07	2 (4%) 40 32	26, 34, 64, 82	0
30	18	64/65 (98%)	0.07	0 100 100	30, 41, 49, 63	0
30	28	64/65 (98%)	0.17	0 100 100	33, 43, 52, 64	0
31	19	37/37 (100%)	0.45	3 (8%) 18 15	38, 48, 70, 74	0
31	29	37/37 (100%)	0.77	2 (5%) 31 24	43, 53, 71, 77	0
32	1a	1477/1521 (97%)	0.19	23 (1%) 70 62	34, 72, 98, 114	0
32	2a	1483/1521 (97%)	0.38	33 (2%) 62 53	43, 78, 101, 115	0
33	1b	231/256 (90%)	1.04	34 (14%) 6 4	67, 86, 96, 115	0
33	2b	231/256 (90%)	1.16	32 (13%) 6 5	71, 88, 97, 106	0
34	1c	206/239 (86%)	1.07	25 (12%) 8 7	70, 84, 93, 99	0
34	2c	206/239 (86%)	1.34	27 (13%) 7 6	73, 85, 94, 96	0
35	1d	208/209 (99%)	0.89	10 (4%) 35 28	56, 77, 88, 100	0
35	2d	208/209 (99%)	0.82	11 (5%) 32 25	56, 73, 84, 92	0
36	1e	148/162 (91%)	0.30	3 (2%) 65 56	47, 67, 79, 87	0
36	2e	148/162 (91%)	0.67	3 (2%) 65 56	56, 74, 84, 90	0
37	1f	100/101 (99%)	0.44	0 100 100	55, 73, 84, 90	0
37	2f	100/101 (99%)	0.52	3 (3%) 52 43	56, 74, 83, 87	0
38	1g	155/156 (99%)	0.82	9 (5%) 29 22	66, 79, 91, 98	0
38	2g	155/156 (99%)	0.97	20 (12%) 7 6	67, 81, 91, 97	0
39	1h	137/138 (99%)	0.40	1 (0%) 84 79	53, 69, 77, 82	0
39	2h	137/138 (99%)	0.79	6 (4%) 39 30	62, 75, 84, 91	0
40	1i	127/128 (99%)	1.40	27 (21%) 2 2	61, 87, 95, 97	0
40	2i	127/128 (99%)	2.00	52 (40%) 0 0	67, 88, 95, 102	0
41	1j	97/105 (92%)	1.45	21 (21%) 2 2	57, 87, 95, 98	0
41	2j	96/105 (91%)	1.83	41 (42%) 0 0	76, 91, 101, 102	0
42	1k	114/129 (88%)	0.34	2 (1%) 67 59	38, 67, 80, 94	0
42	2k	114/129 (88%)	0.58	6 (5%) 32 25	52, 76, 86, 92	0
43	1l	122/132 (92%)	0.54	5 (4%) 41 33	49, 63, 76, 89	0
43	2l	122/132 (92%)	0.57	3 (2%) 58 48	56, 67, 76, 82	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
44	1m	118/126 (93%)	0.80	5 (4%) 40 32	58, 80, 87, 92	0
44	2m	116/126 (92%)	1.42	27 (23%) 2 2	70, 83, 89, 96	0
45	1n	60/61 (98%)	1.24	5 (8%) 17 14	60, 77, 86, 92	0
45	2n	60/61 (98%)	2.38	40 (66%) 0 0	77, 88, 94, 96	0
46	1o	88/89 (98%)	0.52	1 (1%) 78 71	50, 65, 82, 86	0
46	2o	88/89 (98%)	0.52	0 100 100	60, 74, 87, 92	0
47	1p	82/88 (93%)	1.22	11 (13%) 7 5	62, 77, 88, 93	0
47	2p	82/88 (93%)	0.89	3 (3%) 45 37	59, 70, 79, 91	0
48	1q	99/105 (94%)	0.66	3 (3%) 52 43	47, 68, 80, 83	0
48	2q	99/105 (94%)	0.57	3 (3%) 52 43	58, 72, 81, 85	0
49	1r	68/88 (77%)	0.50	0 100 100	58, 68, 83, 87	0
49	2r	68/88 (77%)	0.54	2 (2%) 53 45	63, 74, 84, 88	0
50	1s	84/93 (90%)	1.18	10 (11%) 9 7	67, 81, 92, 98	0
50	2s	83/93 (89%)	1.95	32 (38%) 1 0	76, 91, 102, 105	0
51	1t	96/106 (90%)	1.08	12 (12%) 8 7	57, 75, 84, 90	0
51	2t	96/106 (90%)	0.82	10 (10%) 11 10	56, 73, 84, 87	0
52	1u	23/27 (85%)	1.57	5 (21%) 2 2	68, 77, 82, 85	0
52	2u	23/27 (85%)	1.91	9 (39%) 1 0	68, 80, 85, 88	0
53	1v	5/24 (20%)	0.55	1 (20%) 3 2	50, 55, 86, 88	0
53	2v	5/24 (20%)	1.03	1 (20%) 3 2	73, 74, 89, 103	0
54	1x	72/77 (93%)	0.03	0 100 100	34, 68, 85, 99	0
54	2x	72/77 (93%)	0.15	0 100 100	36, 71, 87, 101	0
55	1z	12/15 (80%)	0.42	2 (16%) 4 3	29, 36, 72, 81	0
55	2z	13/15 (86%)	1.47	4 (30%) 1 1	31, 37, 84, 91	0
All	All	20521/21474 (95%)	0.26	782 (3%) 44 36	15, 62, 92, 115	0

All (782) RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
55	2z	13	PRO	7.2
7	1H	2	SER	5.9
55	2z	12	PRO	5.7
38	1g	79	ARG	5.5
29	17	48	LYS	5.2

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Mol	Chain	Res	Type	RSRZ
32	2a	981	G	5.1
32	1a	980	G	5.1
40	2i	16	ARG	5.0
32	1a	981	G	5.0
50	2s	38	SER	4.9
41	1j	62	HIS	4.6
40	2i	15	ALA	4.6
1	1A	2815	G	4.6
45	2n	18	VAL	4.6
50	2s	8	GLY	4.5
45	2n	25	VAL	4.5
50	2s	71	LEU	4.4
50	2s	11	VAL	4.3
45	2n	17	LYS	4.3
40	1i	13	ALA	4.2
41	1j	98	ILE	4.2
12	1Q	59	ARG	4.2
34	2c	180	ALA	4.2
1	1A	8	U	4.2
44	2m	102	ARG	4.1
20	2Y	5	MET	4.1
52	2u	14	TRP	4.1
34	2c	2	GLY	4.1
52	2u	11	GLY	4.1
40	2i	66	ARG	4.1
41	2j	5	ARG	4.1
6	2G	28	VAL	4.1
52	2u	6	ARG	4.1
32	2a	980	G	4.1
41	2j	40	LEU	4.0
40	2i	14	VAL	4.0
34	2c	155	GLY	4.0
40	2i	17	VAL	4.0
40	2i	102	LEU	3.9
34	1c	28	GLN	3.9
20	2Y	1	MET	3.9
38	2g	78	ARG	3.9
40	2i	42	ARG	3.9
1	1A	1149	C	3.9
50	2s	52	TYR	3.9
50	2s	33	THR	3.9
38	2g	5	ARG	3.9

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Mol	Chain	Res	Type	RSRZ
38	2g	83	ALA	3.9
40	1i	2	GLU	3.9
1	1A	2813	C	3.9
29	27	48	LYS	3.8
40	2i	126	SER	3.8
50	2s	9	VAL	3.8
47	1p	41	PRO	3.8
6	1G	51	ARG	3.8
38	1g	78	ARG	3.7
45	1n	59	ALA	3.7
40	2i	115	GLY	3.7
32	2a	1184	G	3.7
38	1g	80	VAL	3.7
40	2i	29	ASN	3.6
33	2b	187	LEU	3.6
50	2s	35	SER	3.6
26	24	49	PHE	3.6
41	2j	63	PHE	3.6
45	2n	16	PHE	3.6
50	2s	76	PRO	3.6
45	2n	37	PHE	3.6
26	14	53	GLU	3.6
38	1g	156	TRP	3.6
34	2c	129	ALA	3.5
40	2i	61	ALA	3.5
45	2n	35	ARG	3.5
40	2i	68	GLY	3.5
40	1i	106	ALA	3.5
40	2i	65	VAL	3.5
50	2s	51	VAL	3.5
51	1t	10	LEU	3.5
45	2n	3	ARG	3.5
1	1A	2814	C	3.5
34	2c	186	PHE	3.5
40	2i	28	VAL	3.5
40	2i	67	GLY	3.5
33	2b	97	TRP	3.5
15	1T	130	ALA	3.4
35	2d	2	GLY	3.4
43	1l	18	VAL	3.4
34	2c	179	ARG	3.4
50	2s	12	ASP	3.4

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Mol	Chain	Res	Type	RSRZ
40	1i	102	LEU	3.4
15	1T	37	GLY	3.4
40	2i	119	ALA	3.4
45	2n	32	SER	3.4
32	2a	1268	A	3.4
50	2s	13	ASP	3.4
32	2a	1206	G	3.3
55	2z	10	PRO	3.3
43	1l	126	LYS	3.3
38	2g	16	LEU	3.3
41	2j	44	VAL	3.3
44	2m	90	LEU	3.3
41	2j	49	VAL	3.3
40	2i	36	TYR	3.3
44	2m	87	TYR	3.3
45	2n	34	TYR	3.3
1	2A	1582	C	3.3
45	2n	2	ALA	3.3
50	2s	34	TRP	3.3
33	1b	227	GLY	3.3
50	1s	85	LYS	3.3
50	2s	14	HIS	3.3
51	1t	67	ALA	3.3
52	2u	17	THR	3.3
32	1a	982	G	3.3
7	2H	2	SER	3.3
34	2c	8	ILE	3.2
52	2u	13	ILE	3.2
15	1T	131	ALA	3.2
45	2n	33	VAL	3.2
35	2d	49	ARG	3.2
26	24	56	VAL	3.2
1	1A	1098	C	3.2
14	2S	35	ILE	3.2
41	2j	37	PRO	3.2
35	2d	164	ALA	3.2
45	2n	49	HIS	3.2
47	1p	19	ILE	3.2
40	2i	20	ARG	3.1
40	1i	105	ASP	3.1
45	1n	2	ALA	3.1
1	1A	33	C	3.1

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Mol	Chain	Res	Type	RSRZ
40	1i	109	VAL	3.1
20	1Y	1	MET	3.1
34	2c	172	ARG	3.1
7	2H	111	HIS	3.1
24	12	15	LYS	3.1
32	2a	1207	A	3.1
21	2Z	1	MET	3.1
33	2b	101	MET	3.1
51	1t	103	GLY	3.1
40	2i	63	ILE	3.1
45	2n	9	LYS	3.1
8	2I	46	ALA	3.1
33	2b	165	VAL	3.1
45	1n	16	PHE	3.1
34	1c	2	GLY	3.1
34	1c	9	GLY	3.1
40	2i	30	GLY	3.1
33	1b	131	PRO	3.1
38	2g	38	LEU	3.1
40	2i	10	ARG	3.1
51	1t	14	LYS	3.1
38	2g	156	TRP	3.1
44	2m	60	VAL	3.1
1	1A	9	G	3.1
34	2c	190	ARG	3.1
41	2j	55	LYS	3.1
1	1A	934	C	3.1
45	2n	21	TYR	3.0
8	2I	11	ASN	3.0
45	2n	60	SER	3.0
44	2m	6	GLY	3.0
41	1j	70	ARG	3.0
40	2i	110	GLU	3.0
32	2a	1509	A	3.0
33	1b	237	ALA	3.0
1	1A	669	C	3.0
2	2B	88	C	3.0
42	2k	13	GLN	3.0
55	1z	11	ARG	3.0
55	2z	11	ARG	3.0
38	2g	12	LEU	3.0
40	1i	64	THR	3.0

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Mol	Chain	Res	Type	RSRZ
1	2A	33	C	3.0
12	1Q	24	GLY	3.0
45	1n	7	ILE	3.0
50	2s	15	LEU	3.0
41	2j	72	VAL	3.0
32	1a	979	A	3.0
14	2S	32	LEU	3.0
1	1A	2904	C	3.0
1	2A	1149	C	3.0
41	1j	63	PHE	3.0
40	2i	9	ARG	2.9
43	1l	89	ARG	2.9
40	2i	5	TYR	2.9
40	1i	69	GLY	2.9
50	2s	16	LEU	2.9
38	2g	84	ASN	2.9
1	2A	2815	G	2.9
45	2n	15	LYS	2.9
44	2m	64	TRP	2.9
33	1b	167	PRO	2.9
40	2i	2	GLU	2.9
37	2f	89	MET	2.9
33	2b	152	PHE	2.9
34	1c	207	VAL	2.9
33	1b	130	ARG	2.9
14	2S	58	LEU	2.9
41	2j	65	LEU	2.9
45	2n	6	LEU	2.9
39	2h	131	GLY	2.9
45	2n	38	GLY	2.9
14	2S	93	LYS	2.9
40	2i	11	LYS	2.9
40	2i	103	THR	2.9
45	2n	42	ILE	2.9
51	1t	55	ILE	2.9
6	2G	53	LEU	2.9
35	1d	104	VAL	2.9
20	2Y	90	LEU	2.9
40	2i	99	LEU	2.9
33	2b	99	GLY	2.8
41	2j	93	GLY	2.8
41	2j	64	GLU	2.8

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Mol	Chain	Res	Type	RSRZ
1	2A	1216	G	2.8
8	2I	146	ALA	2.8
8	2I	96	ASP	2.8
38	1g	77	SER	2.8
45	2n	7	ILE	2.8
1	1A	1934	A	2.8
20	2Y	57	GLN	2.8
50	1s	56	GLN	2.8
33	2b	197	VAL	2.8
40	1i	10	ARG	2.8
45	2n	29	ARG	2.8
51	1t	12	ALA	2.8
52	1u	10	ARG	2.8
33	1b	195	ASP	2.8
34	2c	5	ILE	2.8
44	2m	9	ILE	2.8
34	2c	41	GLY	2.8
45	1n	38	GLY	2.8
50	2s	68	GLY	2.8
40	1i	12	GLU	2.8
41	1j	64	GLU	2.8
33	2b	92	TYR	2.8
14	2S	29	PHE	2.8
40	2i	106	ALA	2.8
34	1c	178	LEU	2.8
6	1G	48	GLU	2.8
33	1b	30	ARG	2.8
26	14	56	VAL	2.8
29	27	46	VAL	2.8
1	1A	270	U	2.8
51	2t	10	LEU	2.8
41	1j	76	ASN	2.8
40	1i	11	LYS	2.8
35	1d	80	GLU	2.7
52	1u	15	ARG	2.7
8	2I	81	VAL	2.7
33	2b	136	VAL	2.7
40	2i	62	TYR	2.7
20	2Y	34	LYS	2.7
43	1l	91	LYS	2.7
45	2n	22	THR	2.7
40	1i	68	GLY	2.7

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Mol	Chain	Res	Type	RSRZ
6	2G	17	PRO	2.7
8	1I	10	GLU	2.7
32	2a	958	C	2.7
33	2b	188	ALA	2.7
34	2c	189	ALA	2.7
40	2i	76	ALA	2.7
47	2p	7	ALA	2.7
38	2g	42	ILE	2.7
1	2A	2901	G	2.7
32	2a	1202	G	2.7
6	2G	76	SER	2.7
14	2S	3	ARG	2.7
23	1l	2	SER	2.7
42	2k	126	ARG	2.7
41	1j	21	GLN	2.7
33	1b	207	ALA	2.7
40	2i	37	PHE	2.7
50	2s	49	ILE	2.7
7	1H	85	LYS	2.7
34	2c	167	TRP	2.7
33	2b	236	TYR	2.7
40	2i	125	TYR	2.7
36	1e	22	GLY	2.7
40	2i	64	THR	2.7
41	2j	10	GLY	2.7
47	1p	68	ASP	2.7
26	14	57	GLU	2.7
39	2h	53	VAL	2.7
39	2h	2	LEU	2.7
40	2i	96	LEU	2.7
40	1i	15	ALA	2.7
50	2s	40	ILE	2.7
40	2i	117	HIS	2.7
34	1c	201	TYR	2.7
31	29	37	GLY	2.7
44	2m	103	THR	2.7
1	1A	2905	U	2.7
36	2e	20	GLN	2.7
26	24	9	LEU	2.7
33	2b	150	SER	2.7
49	2r	66	LEU	2.7
21	1Z	164	ALA	2.6

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Mol	Chain	Res	Type	RSRZ
14	2S	34	HIS	2.6
14	2S	7	TYR	2.6
44	2m	3	ARG	2.6
32	2a	957	C	2.6
43	2l	18	VAL	2.6
44	2m	101	GLN	2.6
45	2n	8	GLU	2.6
51	2t	24	LEU	2.6
14	2S	52	SER	2.6
14	2S	55	ALA	2.6
33	2b	237	ALA	2.6
45	2n	61	TRP	2.6
1	1A	1220	G	2.6
1	1A	2901	G	2.6
14	2S	92	TYR	2.6
24	12	69	ARG	2.6
41	2j	9	ARG	2.6
5	2F	208	GLY	2.6
35	1d	109	GLY	2.6
44	2m	100	GLY	2.6
34	2c	182	ILE	2.6
32	2a	1201	U	2.6
34	2c	11	ARG	2.6
40	1i	42	ARG	2.6
51	2t	25	ARG	2.6
11	2P	116	GLY	2.6
41	1j	77	PRO	2.6
32	1a	1121	G	2.6
45	2n	44	LEU	2.6
25	23	60	GLU	2.6
33	2b	134	GLU	2.6
40	1i	81	ILE	2.6
38	1g	83	ALA	2.6
45	2n	20	ALA	2.6
32	1a	209	U	2.6
34	1c	6	HIS	2.6
40	1i	117	HIS	2.6
1	1A	2811	A	2.6
32	2a	1165	A	2.6
41	1j	65	LEU	2.6
33	1b	136	VAL	2.6
40	2i	7	THR	2.6

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Mol	Chain	Res	Type	RSRZ
47	1p	22	THR	2.6
26	24	57	GLU	2.6
26	14	59	PHE	2.6
41	2j	78	ASN	2.6
1	2A	10	G	2.5
50	2s	29	ARG	2.5
32	1a	516	A	2.5
36	2e	22	GLY	2.5
50	2s	2	PRO	2.5
19	1X	95	LEU	2.5
35	1d	176	LEU	2.5
51	1t	69	GLY	2.5
52	2u	16	GLY	2.5
50	1s	9	VAL	2.5
34	2c	62	ASP	2.5
42	1k	36	ASP	2.5
50	1s	38	SER	2.5
32	1a	158	C	2.5
34	2c	185	GLY	2.5
26	14	63	TYR	2.5
34	2c	207	VAL	2.5
44	2m	7	VAL	2.5
41	1j	33	GLN	2.5
41	2j	48	THR	2.5
38	2g	2	ALA	2.5
41	1j	32	ALA	2.5
45	2n	30	ALA	2.5
39	2h	135	CYS	2.5
41	2j	59	SER	2.5
14	2S	4	LEU	2.5
33	2b	196	LEU	2.5
40	2i	21	PRO	2.5
41	2j	41	PRO	2.5
50	1s	59	PRO	2.5
7	1H	133	VAL	2.5
1	1A	1554	C	2.5
34	2c	10	PHE	2.5
47	1p	80	PHE	2.5
41	2j	27	ALA	2.5
35	2d	118	ARG	2.5
17	1V	63	GLY	2.5
32	1a	978	U	2.5

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Mol	Chain	Res	Type	RSRZ
41	2j	34	VAL	2.5
34	1c	184	TYR	2.5
40	2i	114	TYR	2.5
45	2n	13	THR	2.5
45	2n	58	LYS	2.5
51	2t	74	LYS	2.5
1	2A	1160	G	2.5
2	2B	118	G	2.5
32	2a	1172	G	2.5
31	19	12	ASP	2.4
50	1s	13	ASP	2.4
26	14	54	GLY	2.4
33	2b	70	PHE	2.4
44	1m	94	ARG	2.4
44	1m	102	ARG	2.4
12	2Q	22	LYS	2.4
1	2A	2801	C	2.4
2	1B	88	C	2.4
32	2a	1205	C	2.4
35	2d	154	ASN	2.4
35	1d	155	LEU	2.4
45	2n	27	CYS	2.4
1	1A	2804	G	2.4
33	1b	234	PRO	2.4
45	2n	14	PRO	2.4
8	2I	92	VAL	2.4
18	2W	112	GLY	2.4
26	24	50	VAL	2.4
37	2f	6	VAL	2.4
32	1a	1023	U	2.4
34	1c	206	GLU	2.4
33	1b	29	ALA	2.4
45	2n	59	ALA	2.4
50	1s	33	THR	2.4
12	1Q	1	MET	2.4
31	19	17	ILE	2.4
41	2j	38	ILE	2.4
23	21	29	GLY	2.4
36	1e	85	GLY	2.4
42	1k	14	VAL	2.4
32	1a	1268	A	2.4
53	2v	14	A	2.4

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Mol	Chain	Res	Type	RSRZ
14	2S	20	ARG	2.4
45	2n	4	LYS	2.4
1	2A	1579	G	2.4
32	1a	988	G	2.4
33	1b	12	GLU	2.4
33	1b	70	PHE	2.4
41	2j	32	ALA	2.4
16	1U	117	GLN	2.4
26	24	67	TYR	2.4
41	2j	87	THR	2.4
50	2s	79	THR	2.4
41	2j	13	HIS	2.4
50	2s	66	MET	2.4
33	1b	41	ILE	2.4
41	2j	50	ILE	2.4
41	1j	93	GLY	2.4
19	1X	68	ARG	2.4
20	2Y	2	ARG	2.4
41	2j	89	ASP	2.4
48	2q	91	ARG	2.4
32	2a	979	A	2.4
32	1a	1239	U	2.4
32	2a	982	G	2.4
47	1p	39	TYR	2.4
34	2c	42	LEU	2.4
34	1c	39	ILE	2.4
35	2d	146	ILE	2.4
44	1m	4	ILE	2.4
34	1c	7	PRO	2.4
6	2G	42	GLY	2.3
14	2S	45	GLY	2.3
35	2d	122	ARG	2.3
41	1j	46	ARG	2.3
45	2n	11	LYS	2.3
51	1t	9	ASN	2.3
51	1t	79	ARG	2.3
26	24	65	ASP	2.3
33	1b	188	ALA	2.3
33	2b	129	GLU	2.3
41	2j	83	GLU	2.3
48	1q	66	SER	2.3
53	1v	14	A	2.3

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Mol	Chain	Res	Type	RSRZ
32	1a	339	U	2.3
40	2i	81	ILE	2.3
41	1j	96	ILE	2.3
44	2m	78	ILE	2.3
6	2G	74	LYS	2.3
55	1z	12	PRO	2.3
1	1A	2812	G	2.3
14	2S	65	VAL	2.3
32	1a	376	G	2.3
32	1a	614	G	2.3
41	2j	66	ARG	2.3
44	2m	94	ARG	2.3
47	1p	79	VAL	2.3
39	1h	15	ASN	2.3
40	1i	67	GLY	2.3
52	1u	2	GLY	2.3
34	2c	60	ALA	2.3
47	1p	82	GLN	2.3
19	2X	92	LEU	2.3
33	1b	187	LEU	2.3
35	1d	162	LEU	2.3
33	1b	97	TRP	2.3
1	1A	1097	C	2.3
1	2A	1120	C	2.3
47	2p	16	HIS	2.3
41	2j	99	LYS	2.3
40	1i	41	VAL	2.3
45	2n	36	PHE	2.3
32	1a	1002	G	2.3
38	2g	117	ALA	2.3
42	2k	89	ALA	2.3
44	2m	5	ALA	2.3
50	2s	50	ALA	2.3
33	2b	118	LEU	2.3
41	1j	71	LEU	2.3
50	1s	22	LEU	2.3
31	29	17	ILE	2.3
35	2d	71	SER	2.3
19	2X	68	ARG	2.3
1	2A	678	A	2.3
7	2H	44	VAL	2.3
21	2Z	175	VAL	2.3

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Mol	Chain	Res	Type	RSRZ
26	24	29	PRO	2.3
52	1u	23	PRO	2.3
32	2a	1099	C	2.3
7	2H	14	GLY	2.3
19	2X	67	GLY	2.3
40	2i	101	PHE	2.3
33	1b	225	ALA	2.3
38	2g	127	ALA	2.3
41	2j	61	GLU	2.3
34	2c	178	LEU	2.3
35	1d	157	LEU	2.3
36	1e	20	GLN	2.3
40	2i	79	LEU	2.3
41	2j	8	LEU	2.3
51	1t	62	LEU	2.3
33	1b	27	LYS	2.3
43	1l	28	LYS	2.3
1	1A	10	G	2.3
14	2S	39	ILE	2.3
32	2a	1204	G	2.3
34	2c	201	TYR	2.3
40	2i	109	VAL	2.3
44	2m	97	PRO	2.3
8	1I	90	GLY	2.3
20	2Y	58	GLY	2.3
38	1g	81	GLY	2.3
32	2a	1239	U	2.3
1	1A	2802	A	2.2
1	2A	1088	C	2.2
6	2G	35	GLU	2.2
42	2k	117	ASN	2.2
44	2m	106	ASN	2.2
50	2s	24	ALA	2.2
39	2h	86	ILE	2.2
44	2m	4	ILE	2.2
6	2G	15	VAL	2.2
47	2p	39	TYR	2.2
50	2s	80	TYR	2.2
34	2c	171	GLY	2.2
40	2i	8	GLY	2.2
41	2j	47	PHE	2.2
34	1c	189	ALA	2.2

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Mol	Chain	Res	Type	RSRZ
44	2m	51	ALA	2.2
51	2t	21	LYS	2.2
52	2u	3	LYS	2.2
16	2U	88	ILE	2.2
51	2t	100	ILE	2.2
12	1Q	5	ARG	2.2
14	2S	5	THR	2.2
33	2b	67	THR	2.2
40	1i	65	VAL	2.2
41	1j	44	VAL	2.2
50	2s	67	VAL	2.2
33	2b	148	TYR	2.2
33	1b	22	LYS	2.2
1	2A	9	G	2.2
1	2A	2812	G	2.2
3	2D	2	ALA	2.2
22	20	75	LEU	2.2
37	2f	61	LEU	2.2
38	2g	36	LYS	2.2
40	1i	19	LEU	2.2
45	2n	10	ALA	2.2
51	1t	74	LYS	2.2
31	19	28	GLU	2.2
32	2a	996	G	2.2
32	2a	1025	G	2.2
32	2a	1265	G	2.2
33	2b	170	GLU	2.2
41	2j	74	ILE	2.2
47	1p	42	ARG	2.2
50	2s	31	ILE	2.2
50	2s	36	ARG	2.2
32	1a	160	C	2.2
33	2b	189	ASP	2.2
6	2G	109	VAL	2.2
6	2G	160	VAL	2.2
7	1H	45	VAL	2.2
14	2S	28	VAL	2.2
14	2S	91	PRO	2.2
41	2j	24	VAL	2.2
26	24	52	THR	2.2
33	1b	228	GLY	2.2
38	2g	154	TYR	2.2

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Mol	Chain	Res	Type	RSRZ
43	2l	69	TYR	2.2
52	1u	18	TYR	2.2
40	2i	112	LYS	2.2
41	2j	71	LEU	2.2
44	2m	81	LEU	2.2
33	2b	123	ALA	2.2
33	1b	214	ILE	2.2
33	2b	42	ILE	2.2
33	2b	144	ARG	2.2
33	2b	201	ILE	2.2
34	1c	8	ILE	2.2
38	2g	4	ARG	2.2
51	2t	9	ASN	2.2
1	1A	2205	G	2.2
1	2A	159	G	2.2
32	2a	1356	G	2.2
1	1A	571	A	2.2
32	1a	1120	C	2.2
33	1b	101	MET	2.2
40	2i	27	THR	2.2
22	20	76	GLY	2.2
34	1c	171	GLY	2.2
12	2Q	104	PHE	2.2
34	1c	23	TYR	2.2
22	20	84	LEU	2.2
33	1b	9	GLU	2.2
44	2m	42	ALA	2.2
44	2m	75	ALA	2.2
38	1g	5	ARG	2.1
41	1j	5	ARG	2.1
1	2A	1071	U	2.1
33	1b	19	HIS	2.1
7	2H	8	PRO	2.1
14	2S	98	VAL	2.1
34	1c	195	VAL	2.1
34	2c	55	VAL	2.1
41	2j	77	PRO	2.1
1	1A	2902	G	2.1
1	2A	1108	G	2.1
1	2A	1134	G	2.1
32	1a	176	G	2.1
32	1a	1509	A	2.1

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Mol	Chain	Res	Type	RSRZ
32	2a	1469	G	2.1
35	2d	124	GLY	2.1
45	2n	28	GLY	2.1
32	2a	1303	C	2.1
34	1c	33	LEU	2.1
45	2n	39	LEU	2.1
48	2q	98	LEU	2.1
40	2i	92	TYR	2.1
26	24	66	SER	2.1
40	1i	126	SER	2.1
40	2i	13	ALA	2.1
34	1c	125	GLU	2.1
45	2n	31	ARG	2.1
48	2q	24	GLU	2.1
51	2t	23	ARG	2.1
52	2u	15	ARG	2.1
1	1A	1465	U	2.1
7	2H	131	VAL	2.1
51	2t	98	PRO	2.1
3	2D	275	LYS	2.1
14	2S	59	LYS	2.1
40	2i	105	ASP	2.1
35	1d	69	GLY	2.1
35	2d	167	GLY	2.1
40	1i	8	GLY	2.1
50	2s	63	THR	2.1
35	1d	101	LEU	2.1
32	2a	956	A	2.1
33	2b	186	ALA	2.1
44	2m	118	ALA	2.1
6	1G	35	GLU	2.1
32	2a	1203	G	2.1
33	1b	201	ILE	2.1
1	2A	1121	C	2.1
23	21	2	SER	2.1
11	2P	68	GLN	2.1
17	2V	72	VAL	2.1
5	1F	14	PRO	2.1
6	2G	142	PRO	2.1
33	1b	133	LYS	2.1
43	2l	49	ASN	2.1
51	2t	29	LYS	2.1

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Mol	Chain	Res	Type	RSRZ
1	1A	1150	U	2.1
6	2G	175	LEU	2.1
33	1b	11	LEU	2.1
34	2c	159	GLY	2.1
34	2c	197	GLY	2.1
40	1i	39	GLY	2.1
41	2j	36	GLY	2.1
41	2j	58	ASP	2.1
46	1o	89	GLY	2.1
14	1S	3	ARG	2.1
44	2m	104	ARG	2.1
52	2u	10	ARG	2.1
6	1G	46	ALA	2.1
6	2G	151	ALA	2.1
38	2g	7	ALA	2.1
34	1c	90	GLU	2.1
38	1g	151	TYR	2.1
32	2a	952	A	2.1
19	2X	31	HIS	2.1
1	2A	2330	G	2.1
32	2a	1235	G	2.1
8	1I	8	PRO	2.1
41	2j	91	PRO	2.1
40	2i	40	LEU	2.1
14	2S	12	PHE	2.1
32	1a	211	U	2.1
34	1c	81	GLY	2.1
48	1q	33	GLY	2.1
22	20	43	THR	2.1
26	14	55	ARG	2.1
34	1c	30	ARG	2.1
39	2h	122	ARG	2.1
41	2j	42	THR	2.1
45	2n	12	ARG	2.1
50	2s	3	ARG	2.1
6	2G	61	ALA	2.1
36	2e	13	ILE	2.1
49	2r	60	ALA	2.1
40	2i	12	GLU	2.1
33	2b	31	TYR	2.1
34	1c	193	TYR	2.1
33	2b	133	LYS	2.0

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Mol	Chain	Res	Type	RSRZ
40	1i	113	LYS	2.0
33	1b	16	HIS	2.0
33	2b	140	HIS	2.0
42	2k	104	GLN	2.0
47	1p	16	HIS	2.0
7	2H	45	VAL	2.0
7	2H	169	VAL	2.0
29	17	46	VAL	2.0
32	1a	1432	A	2.0
34	1c	64	VAL	2.0
1	2A	1154	C	2.0
32	2a	1304	C	2.0
35	2d	155	LEU	2.0
40	2i	85	LEU	2.0
41	1j	8	LEU	2.0
44	2m	48	LEU	2.0
6	1G	80	PHE	2.0
14	2S	60	GLY	2.0
35	1d	139	ARG	2.0
38	2g	82	GLY	2.0
41	1j	47	PHE	2.0
42	2k	90	GLY	2.0
44	2m	95	GLY	2.0
48	1q	72	ARG	2.0
33	1b	222	ILE	2.0
6	1G	50	ALA	2.0
38	2g	39	ALA	2.0
44	1m	2	ALA	2.0
47	1p	70	ALA	2.0
33	2b	12	GLU	2.0
40	1i	4	TYR	2.0
44	1m	59	TYR	2.0
50	1s	6	LYS	2.0
33	2b	48	MET	2.0
17	2V	73	SER	2.0
33	1b	7	VAL	2.0
41	1j	72	VAL	2.0
51	1t	16	HIS	2.0
25	13	2	PRO	2.0
32	2a	1339	A	2.0
33	1b	10	LEU	2.0
41	1j	90	LEU	2.0

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Mol	Chain	Res	Type	RSRZ
19	1X	65	ARG	2.0
38	2g	79	ARG	2.0
41	2j	76	ASN	2.0
50	1s	29	ARG	2.0
23	21	28	GLY	2.0
26	24	45	GLY	2.0
33	1b	122	PHE	2.0
22	10	10	THR	2.0
34	1c	169	ALA	2.0
40	1i	122	ALA	2.0
41	2j	67	THR	2.0
13	2R	69	ASP	2.0
32	2a	1510	U	2.0
32	2a	1000	G	2.0
33	1b	134	GLU	2.0
34	1c	62	ASP	2.0
40	1i	62	TYR	2.0
21	2Z	96	VAL	2.0
38	2g	141	VAL	2.0
44	2m	15	VAL	2.0
50	2s	41	VAL	2.0

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

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

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors(Å ²)	Q<0.9
54	PSU	2x	55	20/21	0.83	0.11	62,68,78,88	0
54	4SU	2x	8	20/21	0.88	0.10	57,78,89,92	0
54	5MU	2x	54	21/22	0.89	0.10	72,76,86,101	0
54	5MC	2x	32	21/22	0.89	0.14	68,80,88,91	0
54	5MU	1x	54	21/22	0.92	0.10	50,67,76,78	0
54	PSU	1x	55	20/21	0.93	0.09	57,65,79,80	0
54	5MC	1x	32	21/22	0.93	0.12	51,62,76,86	0
54	4SU	1x	8	20/21	0.95	0.09	47,65,79,80	0

5.3 Carbohydrates ⓘ

There are no oligosaccharides in this entry.

5.4 Ligands ⓘ

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

Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3298	1/1	0.50	0.25	81,81,81,81	0
56	MG	1A	3901	1/1	0.53	0.27	86,86,86,86	0
56	MG	2a	3145	1/1	0.53	0.18	95,95,95,95	0
56	MG	1A	3517	1/1	0.54	0.18	69,69,69,69	0
56	MG	1A	4076	1/1	0.57	0.25	72,72,72,72	0
56	MG	1a	1784	1/1	0.58	0.26	80,80,80,80	0
56	MG	1A	3348	1/1	0.58	0.15	52,52,52,52	0
56	MG	1A	4037	1/1	0.59	0.31	75,75,75,75	0
56	MG	1A	3525	1/1	0.60	0.19	74,74,74,74	0
56	MG	2A	3224	1/1	0.62	0.19	73,73,73,73	0
56	MG	1a	1800	1/1	0.62	0.22	79,79,79,79	0
56	MG	2a	3167	1/1	0.63	0.20	59,59,59,59	0
56	MG	2q	201	1/1	0.63	0.14	58,58,58,58	0
56	MG	1a	1773	1/1	0.64	0.36	77,77,77,77	0
56	MG	1A	3511	1/1	0.64	0.18	55,55,55,55	0
56	MG	2a	3121	1/1	0.64	0.21	88,88,88,88	0
56	MG	2a	3150	1/1	0.65	0.20	76,76,76,76	0
56	MG	1A	4016	1/1	0.65	0.17	77,77,77,77	0
56	MG	2A	3476	1/1	0.65	0.26	66,66,66,66	0
56	MG	1A	3369	1/1	0.68	0.21	67,67,67,67	0
56	MG	1a	1615	1/1	0.68	0.16	66,66,66,66	0
56	MG	1a	1752	1/1	0.68	0.15	61,61,61,61	0
56	MG	1A	4058	1/1	0.68	0.17	48,48,48,48	0
56	MG	2a	3053	1/1	0.68	0.13	99,99,99,99	0
56	MG	1A	3411	1/1	0.69	0.25	51,51,51,51	0
56	MG	1A	3433	1/1	0.69	0.16	51,51,51,51	0
56	MG	1A	4041	1/1	0.69	0.13	57,57,57,57	0
56	MG	2a	3234	1/1	0.69	0.31	82,82,82,82	0
56	MG	1a	1738	1/1	0.69	0.22	73,73,73,73	0
56	MG	2B	202	1/1	0.70	0.17	55,55,55,55	0
56	MG	1A	3847	1/1	0.70	0.18	55,55,55,55	0
56	MG	1a	1783	1/1	0.70	0.15	73,73,73,73	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3242	1/1	0.70	0.19	67,67,67,67	0
56	MG	2x	107	1/1	0.70	0.17	78,78,78,78	0
56	MG	1m	201	1/1	0.71	0.13	95,95,95,95	0
56	MG	2a	3230	1/1	0.71	0.15	67,67,67,67	0
56	MG	2A	3498	1/1	0.71	0.16	69,69,69,69	0
56	MG	1A	3333	1/1	0.71	0.24	57,57,57,57	0
56	MG	2x	106	1/1	0.71	0.14	82,82,82,82	0
56	MG	2a	3006	1/1	0.71	0.22	77,77,77,77	0
56	MG	2a	3223	1/1	0.72	0.28	77,77,77,77	0
56	MG	1a	1848	1/1	0.72	0.19	77,77,77,77	0
56	MG	1a	1645	1/1	0.72	0.15	78,78,78,78	0
56	MG	2a	3051	1/1	0.73	0.13	91,91,91,91	0
56	MG	1A	3318	1/1	0.73	0.17	65,65,65,65	0
56	MG	2A	3432	1/1	0.73	0.15	55,55,55,55	0
56	MG	2a	3114	1/1	0.74	0.17	77,77,77,77	0
56	MG	2a	3157	1/1	0.74	0.32	108,108,108,108	0
56	MG	1A	4034	1/1	0.74	0.18	72,72,72,72	0
56	MG	2q	202	1/1	0.74	0.18	59,59,59,59	0
56	MG	2a	3168	1/1	0.74	0.23	67,67,67,67	0
56	MG	2A	3456	1/1	0.74	0.31	64,64,64,64	0
56	MG	2a	3162	1/1	0.75	0.18	79,79,79,79	0
56	MG	1A	3510	1/1	0.75	0.12	46,46,46,46	0
56	MG	2a	3146	1/1	0.75	0.16	71,71,71,71	0
56	MG	2a	3049	1/1	0.75	0.15	79,79,79,79	0
56	MG	2A	3445	1/1	0.75	0.28	74,74,74,74	0
56	MG	1A	3974	1/1	0.76	0.25	59,59,59,59	0
56	MG	1A	4000	1/1	0.76	0.20	69,69,69,69	0
56	MG	2a	3019	1/1	0.76	0.11	72,72,72,72	0
56	MG	2a	3047	1/1	0.76	0.17	79,79,79,79	0
56	MG	2A	3339	1/1	0.76	0.27	63,63,63,63	0
56	MG	2A	3380	1/1	0.76	0.15	59,59,59,59	0
56	MG	1A	3557	1/1	0.76	0.14	71,71,71,71	0
56	MG	2a	3054	1/1	0.76	0.11	96,96,96,96	0
56	MG	1B	214	1/1	0.76	0.16	56,56,56,56	0
56	MG	1Q	206	1/1	0.76	0.17	53,53,53,53	0
56	MG	1A	3408	1/1	0.76	0.32	58,58,58,58	0
56	MG	1A	3493	1/1	0.76	0.10	64,64,64,64	0
56	MG	2A	3404	1/1	0.77	0.21	64,64,64,64	0
56	MG	1A	4027	1/1	0.77	0.17	62,62,62,62	0
56	MG	2a	3039	1/1	0.77	0.22	67,67,67,67	0
56	MG	2A	3443	1/1	0.77	0.26	63,63,63,63	0
56	MG	2A	3197	1/1	0.77	0.12	63,63,63,63	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1F	304	1/1	0.77	0.21	73,73,73,73	0
56	MG	1A	3644	1/1	0.77	0.12	45,45,45,45	0
56	MG	2A	3490	1/1	0.77	0.32	66,66,66,66	0
56	MG	2a	3101	1/1	0.77	0.21	77,77,77,77	0
56	MG	1a	1852	1/1	0.77	0.19	82,82,82,82	0
56	MG	2A	3391	1/1	0.77	0.17	52,52,52,52	0
56	MG	2a	3142	1/1	0.77	0.26	80,80,80,80	0
56	MG	2a	3004	1/1	0.77	0.28	61,61,61,61	0
56	MG	1A	3491	1/1	0.78	0.09	54,54,54,54	0
56	MG	2A	3370	1/1	0.78	0.25	65,65,65,65	0
56	MG	1A	3989	1/1	0.78	0.39	31,31,31,31	0
56	MG	1A	3513	1/1	0.78	0.30	59,59,59,59	0
56	MG	2a	3151	1/1	0.78	0.11	97,97,97,97	0
56	MG	2a	3031	1/1	0.78	0.14	88,88,88,88	0
56	MG	2a	3161	1/1	0.78	0.14	63,63,63,63	0
56	MG	1Z	301	1/1	0.78	0.15	94,94,94,94	0
56	MG	2a	3165	1/1	0.78	0.20	74,74,74,74	0
56	MG	10	106	1/1	0.78	0.19	54,54,54,54	0
56	MG	18	3002	1/1	0.78	0.11	58,58,58,58	0
56	MG	1A	4054	1/1	0.78	0.25	69,69,69,69	0
56	MG	1A	3218	1/1	0.78	0.14	58,58,58,58	0
56	MG	2A	3171	1/1	0.78	0.11	49,49,49,49	0
56	MG	2a	3070	1/1	0.78	0.28	66,66,66,66	0
56	MG	2A	3483	1/1	0.78	0.12	72,72,72,72	0
56	MG	1A	3465	1/1	0.78	0.11	59,59,59,59	0
56	MG	1a	1747	1/1	0.78	0.17	64,64,64,64	0
56	MG	1a	1782	1/1	0.79	0.13	73,73,73,73	0
56	MG	2A	3194	1/1	0.79	0.17	56,56,56,56	0
56	MG	1H	3002	1/1	0.79	0.11	62,62,62,62	0
56	MG	11	102	1/1	0.79	0.16	65,65,65,65	0
56	MG	2A	3308	1/1	0.79	0.21	53,53,53,53	0
56	MG	2a	3055	1/1	0.79	0.13	81,81,81,81	0
56	MG	12	3001	1/1	0.79	0.19	57,57,57,57	0
56	MG	2a	3072	1/1	0.79	0.18	65,65,65,65	0
56	MG	2a	3206	1/1	0.79	0.17	56,56,56,56	0
56	MG	2a	3208	1/1	0.79	0.17	65,65,65,65	0
56	MG	2a	3076	1/1	0.79	0.19	65,65,65,65	0
56	MG	1a	1811	1/1	0.79	0.12	66,66,66,66	0
56	MG	1A	4066	1/1	0.79	0.25	35,35,35,35	0
56	MG	1a	1764	1/1	0.79	0.14	59,59,59,59	0
56	MG	1a	1871	1/1	0.79	0.34	69,69,69,69	0
56	MG	1A	3345	1/1	0.79	0.17	63,63,63,63	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3114	1/1	0.79	0.20	50,50,50,50	0
56	MG	1A	3101	1/1	0.80	0.13	50,50,50,50	0
56	MG	1A	3935	1/1	0.80	0.13	52,52,52,52	0
56	MG	1a	1718	1/1	0.80	0.14	75,75,75,75	0
56	MG	1P	204	1/1	0.80	0.15	61,61,61,61	0
56	MG	1A	3138	1/1	0.80	0.17	48,48,48,48	0
56	MG	1T	206	1/1	0.80	0.17	76,76,76,76	0
56	MG	2A	3104	1/1	0.80	0.11	59,59,59,59	0
56	MG	1A	3041	1/1	0.80	0.21	51,51,51,51	0
56	MG	2a	3187	1/1	0.80	0.28	77,77,77,77	0
56	MG	2a	3189	1/1	0.80	0.27	76,76,76,76	0
56	MG	1A	3728	1/1	0.80	0.19	62,62,62,62	0
56	MG	2A	3172	1/1	0.80	0.12	63,63,63,63	0
56	MG	2a	3210	1/1	0.80	0.16	78,78,78,78	0
56	MG	1A	3417	1/1	0.80	0.08	76,76,76,76	0
56	MG	1A	3891	1/1	0.80	0.28	68,68,68,68	0
56	MG	2A	3561	1/1	0.80	0.12	55,55,55,55	0
56	MG	2a	3254	1/1	0.80	0.29	76,76,76,76	0
56	MG	1A	4033	1/1	0.80	0.18	62,62,62,62	0
56	MG	2A	3261	1/1	0.80	0.11	43,43,43,43	0
56	MG	1a	1789	1/1	0.80	0.18	60,60,60,60	0
56	MG	1a	1792	1/1	0.80	0.19	92,92,92,92	0
56	MG	2A	3128	1/1	0.81	0.14	56,56,56,56	0
56	MG	1a	1820	1/1	0.81	0.14	79,79,79,79	0
56	MG	1a	1786	1/1	0.81	0.20	49,49,49,49	0
56	MG	2A	3433	1/1	0.81	0.14	43,43,43,43	0
56	MG	2a	3193	1/1	0.81	0.12	65,65,65,65	0
56	MG	2a	3138	1/1	0.81	0.20	81,81,81,81	0
56	MG	2a	3141	1/1	0.81	0.14	70,70,70,70	0
56	MG	1A	3019	1/1	0.81	0.22	46,46,46,46	0
56	MG	1a	1791	1/1	0.81	0.28	82,82,82,82	0
56	MG	2a	3226	1/1	0.81	0.16	77,77,77,77	0
56	MG	2a	3229	1/1	0.81	0.17	57,57,57,57	0
56	MG	1a	1872	1/1	0.81	0.12	57,57,57,57	0
56	MG	1a	1880	1/1	0.81	0.20	69,69,69,69	0
56	MG	2A	3278	1/1	0.81	0.14	43,43,43,43	0
56	MG	2l	3002	1/1	0.81	0.25	68,68,68,68	0
56	MG	1A	3368	1/1	0.81	0.17	65,65,65,65	0
56	MG	1o	101	1/1	0.81	0.15	81,81,81,81	0
56	MG	1B	223	1/1	0.81	0.14	65,65,65,65	0
56	MG	14	502	1/1	0.81	0.20	64,64,64,64	0
56	MG	1a	1842	1/1	0.82	0.23	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2a	3153	1/1	0.82	0.12	92,92,92,92	0
56	MG	1a	1761	1/1	0.82	0.15	65,65,65,65	0
56	MG	1A	4025	1/1	0.82	0.19	80,80,80,80	0
56	MG	1B	210	1/1	0.82	0.12	56,56,56,56	0
56	MG	2a	3164	1/1	0.82	0.19	80,80,80,80	0
56	MG	1A	3515	1/1	0.82	0.38	41,41,41,41	0
56	MG	1A	3351	1/1	0.82	0.11	46,46,46,46	0
56	MG	1a	1905	1/1	0.82	0.13	67,67,67,67	0
56	MG	1e	201	1/1	0.82	0.14	66,66,66,66	0
56	MG	1E	307	1/1	0.82	0.09	57,57,57,57	0
56	MG	1A	3916	1/1	0.82	0.23	45,45,45,45	0
56	MG	2a	3058	1/1	0.82	0.17	66,66,66,66	0
56	MG	2A	3054	1/1	0.82	0.21	58,58,58,58	0
56	MG	1A	3416	1/1	0.82	0.18	66,66,66,66	0
56	MG	1a	1656	1/1	0.82	0.10	84,84,84,84	0
56	MG	1A	3494	1/1	0.82	0.34	68,68,68,68	0
56	MG	1a	1798	1/1	0.82	0.15	71,71,71,71	0
56	MG	2a	3117	1/1	0.82	0.36	66,66,66,66	0
56	MG	1A	3383	1/1	0.82	0.13	67,67,67,67	0
56	MG	1A	3405	1/1	0.82	0.29	41,41,41,41	0
56	MG	2A	3505	1/1	0.82	0.18	50,50,50,50	0
56	MG	1A	3121	1/1	0.82	0.07	72,72,72,72	0
56	MG	1a	1823	1/1	0.82	0.12	71,71,71,71	0
56	MG	2a	3001	1/1	0.82	0.27	62,62,62,62	0
56	MG	2A	3230	1/1	0.82	0.18	47,47,47,47	0
56	MG	2A	3087	1/1	0.83	0.11	53,53,53,53	0
56	MG	2A	3093	1/1	0.83	0.18	57,57,57,57	0
56	MG	2A	3486	1/1	0.83	0.11	42,42,42,42	0
56	MG	1A	3550	1/1	0.83	0.18	68,68,68,68	0
56	MG	2a	3149	1/1	0.83	0.13	63,63,63,63	0
56	MG	1H	3004	1/1	0.83	0.10	66,66,66,66	0
56	MG	1a	1668	1/1	0.83	0.15	63,63,63,63	0
56	MG	2a	3152	1/1	0.83	0.16	67,67,67,67	0
56	MG	1A	3456	1/1	0.83	0.10	71,71,71,71	0
56	MG	1a	1732	1/1	0.83	0.27	63,63,63,63	0
56	MG	2a	3158	1/1	0.83	0.24	65,65,65,65	0
56	MG	1A	3366	1/1	0.83	0.11	42,42,42,42	0
56	MG	1a	1739	1/1	0.83	0.14	70,70,70,70	0
56	MG	1a	1827	1/1	0.83	0.19	68,68,68,68	0
56	MG	1A	3056	1/1	0.83	0.10	59,59,59,59	0
56	MG	1A	3747	1/1	0.83	0.21	50,50,50,50	0
56	MG	2a	3032	1/1	0.83	0.15	59,59,59,59	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2a	3169	1/1	0.83	0.25	67,67,67,67	0
56	MG	2a	3170	1/1	0.83	0.21	72,72,72,72	0
56	MG	2A	3272	1/1	0.83	0.19	67,67,67,67	0
56	MG	1A	3444	1/1	0.83	0.17	57,57,57,57	0
56	MG	2a	3191	1/1	0.83	0.10	88,88,88,88	0
56	MG	1A	3881	1/1	0.83	0.14	64,64,64,64	0
56	MG	2a	3197	1/1	0.83	0.13	78,78,78,78	0
56	MG	2a	3198	1/1	0.83	0.16	65,65,65,65	0
56	MG	2A	3336	1/1	0.83	0.15	74,74,74,74	0
56	MG	1A	3451	1/1	0.83	0.12	66,66,66,66	0
56	MG	2A	3364	1/1	0.83	0.11	44,44,44,44	0
56	MG	1a	1781	1/1	0.83	0.11	75,75,75,75	0
56	MG	1B	229	1/1	0.83	0.10	36,36,36,36	0
56	MG	1b	3001	1/1	0.83	0.19	70,70,70,70	0
56	MG	1A	3509	1/1	0.83	0.19	68,68,68,68	0
56	MG	1h	8002	1/1	0.83	0.28	80,80,80,80	0
56	MG	2a	3240	1/1	0.83	0.27	66,66,66,66	0
56	MG	2a	3243	1/1	0.83	0.24	74,74,74,74	0
56	MG	1A	3529	1/1	0.83	0.16	61,61,61,61	0
56	MG	1a	1627	1/1	0.83	0.18	70,70,70,70	0
56	MG	2A	3035	1/1	0.83	0.20	54,54,54,54	0
56	MG	1a	1639	1/1	0.83	0.31	77,77,77,77	0
56	MG	2a	3126	1/1	0.83	0.25	60,60,60,60	0
56	MG	2A	3467	1/1	0.83	0.10	47,47,47,47	0
56	MG	1A	3749	1/1	0.84	0.20	43,43,43,43	0
56	MG	2A	3374	1/1	0.84	0.12	58,58,58,58	0
56	MG	1A	3449	1/1	0.84	0.14	67,67,67,67	0
56	MG	1H	3003	1/1	0.84	0.12	65,65,65,65	0
56	MG	2A	3401	1/1	0.84	0.29	70,70,70,70	0
56	MG	1A	3115	1/1	0.84	0.15	56,56,56,56	0
56	MG	2A	3162	1/1	0.84	0.07	40,40,40,40	0
56	MG	1A	3146	1/1	0.84	0.24	62,62,62,62	0
56	MG	2A	3439	1/1	0.84	0.10	60,60,60,60	0
56	MG	2a	3061	1/1	0.84	0.13	71,71,71,71	0
56	MG	1a	1756	1/1	0.84	0.21	58,58,58,58	0
56	MG	2A	3177	1/1	0.84	0.16	54,54,54,54	0
56	MG	2A	3189	1/1	0.84	0.21	52,52,52,52	0
56	MG	1a	1909	1/1	0.84	0.09	71,71,71,71	0
56	MG	1A	3619	1/1	0.84	0.14	56,56,56,56	0
56	MG	1d	503	1/1	0.84	0.17	77,77,77,77	0
56	MG	1a	1806	1/1	0.84	0.13	51,51,51,51	0
56	MG	2a	3122	1/1	0.84	0.32	71,71,71,71	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2a	3217	1/1	0.84	0.21	68,68,68,68	0
56	MG	2a	3218	1/1	0.84	0.19	67,67,67,67	0
56	MG	2A	3239	1/1	0.84	0.23	59,59,59,59	0
56	MG	2a	3130	1/1	0.84	0.12	50,50,50,50	0
56	MG	2A	3256	1/1	0.84	0.15	43,43,43,43	0
56	MG	1A	3506	1/1	0.84	0.25	75,75,75,75	0
56	MG	1a	1818	1/1	0.84	0.13	81,81,81,81	0
56	MG	2A	3275	1/1	0.84	0.15	51,51,51,51	0
56	MG	2G	3001	1/1	0.84	0.09	56,56,56,56	0
56	MG	2Z	301	1/1	0.84	0.15	70,70,70,70	0
56	MG	2e	3001	1/1	0.84	0.20	74,74,74,74	0
56	MG	1A	3508	1/1	0.84	0.13	68,68,68,68	0
56	MG	1A	3948	1/1	0.84	0.27	41,41,41,41	0
56	MG	1A	3004	1/1	0.84	0.11	54,54,54,54	0
56	MG	2A	3074	1/1	0.84	0.09	54,54,54,54	0
56	MG	1a	1719	1/1	0.84	0.28	68,68,68,68	0
56	MG	2a	3118	1/1	0.85	0.18	60,60,60,60	0
56	MG	1a	1751	1/1	0.85	0.17	64,64,64,64	0
56	MG	1N	3002	1/1	0.85	0.11	48,48,48,48	0
56	MG	1A	3514	1/1	0.85	0.32	59,59,59,59	0
56	MG	1a	1757	1/1	0.85	0.15	69,69,69,69	0
56	MG	1A	3485	1/1	0.85	0.21	57,57,57,57	0
56	MG	1A	3402	1/1	0.85	0.27	60,60,60,60	0
56	MG	1X	3001	1/1	0.85	0.23	56,56,56,56	0
56	MG	1r	3002	1/1	0.85	0.17	70,70,70,70	0
56	MG	1a	1776	1/1	0.85	0.07	67,67,67,67	0
56	MG	1A	3016	1/1	0.85	0.10	56,56,56,56	0
56	MG	1A	3198	1/1	0.85	0.12	50,50,50,50	0
56	MG	2A	3482	1/1	0.85	0.14	52,52,52,52	0
56	MG	1A	3922	1/1	0.85	0.13	25,25,25,25	0
56	MG	1A	4061	1/1	0.85	0.16	61,61,61,61	0
56	MG	1A	3927	1/1	0.85	0.29	66,66,66,66	0
56	MG	2A	3494	1/1	0.85	0.17	38,38,38,38	0
56	MG	1A	3502	1/1	0.85	0.10	70,70,70,70	0
56	MG	1A	4083	1/1	0.85	0.18	65,65,65,65	0
56	MG	2A	3559	1/1	0.85	0.09	56,56,56,56	0
56	MG	1a	1623	1/1	0.85	0.09	67,67,67,67	0
56	MG	1a	1795	1/1	0.85	0.14	64,64,64,64	0
56	MG	1A	3941	1/1	0.85	0.11	60,60,60,60	0
56	MG	1A	3301	1/1	0.85	0.15	64,64,64,64	0
56	MG	1a	1802	1/1	0.85	0.25	68,68,68,68	0
56	MG	2a	3175	1/1	0.85	0.16	66,66,66,66	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1644	1/1	0.85	0.22	75,75,75,75	0
56	MG	1A	3970	1/1	0.85	0.23	64,64,64,64	0
56	MG	2a	3014	1/1	0.85	0.25	64,64,64,64	0
56	MG	1a	1815	1/1	0.85	0.14	73,73,73,73	0
56	MG	1B	227	1/1	0.85	0.08	69,69,69,69	0
56	MG	1a	1819	1/1	0.85	0.20	73,73,73,73	0
56	MG	2A	3246	1/1	0.85	0.16	55,55,55,55	0
56	MG	2a	3043	1/1	0.85	0.22	78,78,78,78	0
56	MG	2a	3045	1/1	0.85	0.23	68,68,68,68	0
56	MG	1A	3593	1/1	0.85	0.16	62,62,62,62	0
56	MG	2A	3257	1/1	0.85	0.18	67,67,67,67	0
56	MG	1a	1713	1/1	0.85	0.12	66,66,66,66	0
56	MG	1B	230	1/1	0.85	0.08	67,67,67,67	0
56	MG	1a	1831	1/1	0.85	0.15	84,84,84,84	0
56	MG	1A	3453	1/1	0.85	0.11	72,72,72,72	0
56	MG	2a	3057	1/1	0.85	0.21	70,70,70,70	0
56	MG	2A	3298	1/1	0.85	0.12	52,52,52,52	0
56	MG	1A	3087	1/1	0.85	0.15	45,45,45,45	0
56	MG	1A	3384	1/1	0.85	0.32	59,59,59,59	0
56	MG	1A	3473	1/1	0.85	0.17	55,55,55,55	0
56	MG	1a	1744	1/1	0.85	0.32	58,58,58,58	0
56	MG	1a	1746	1/1	0.85	0.16	59,59,59,59	0
56	MG	2a	3111	1/1	0.85	0.22	60,60,60,60	0
56	MG	1a	1899	1/1	0.85	0.16	64,64,64,64	0
56	MG	1A	3479	1/1	0.85	0.11	59,59,59,59	0
56	MG	1T	202	1/1	0.86	0.15	51,51,51,51	0
56	MG	2A	3366	1/1	0.86	0.15	50,50,50,50	0
56	MG	1A	3914	1/1	0.86	0.17	41,41,41,41	0
56	MG	1A	3092	1/1	0.86	0.23	44,44,44,44	0
56	MG	1A	3450	1/1	0.86	0.25	52,52,52,52	0
56	MG	1Z	302	1/1	0.86	0.11	65,65,65,65	0
56	MG	1A	4063	1/1	0.86	0.14	47,47,47,47	0
56	MG	1A	3281	1/1	0.86	0.24	62,62,62,62	0
56	MG	2A	3408	1/1	0.86	0.11	62,62,62,62	0
56	MG	1A	3012	1/1	0.86	0.24	66,66,66,66	0
56	MG	2a	3133	1/1	0.86	0.08	63,63,63,63	0
56	MG	1A	3174	1/1	0.86	0.15	47,47,47,47	0
56	MG	2A	3030	1/1	0.86	0.26	64,64,64,64	0
56	MG	1A	3646	1/1	0.86	0.12	65,65,65,65	0
56	MG	1a	1602	1/1	0.86	0.15	61,61,61,61	0
56	MG	1a	1603	1/1	0.86	0.22	69,69,69,69	0
56	MG	2a	3148	1/1	0.86	0.20	56,56,56,56	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1605	1/1	0.86	0.24	68,68,68,68	0
56	MG	1A	3689	1/1	0.86	0.15	65,65,65,65	0
56	MG	2A	3479	1/1	0.86	0.16	47,47,47,47	0
56	MG	2A	3099	1/1	0.86	0.07	61,61,61,61	0
56	MG	2A	3101	1/1	0.86	0.12	56,56,56,56	0
56	MG	1B	216	1/1	0.86	0.11	70,70,70,70	0
56	MG	1A	3727	1/1	0.86	0.12	49,49,49,49	0
56	MG	2a	3159	1/1	0.86	0.12	58,58,58,58	0
56	MG	1a	1797	1/1	0.86	0.08	73,73,73,73	0
56	MG	2A	3496	1/1	0.86	0.10	35,35,35,35	0
56	MG	2A	3131	1/1	0.86	0.16	56,56,56,56	0
56	MG	1A	3023	1/1	0.86	0.09	55,55,55,55	0
56	MG	2A	3510	1/1	0.86	0.20	61,61,61,61	0
56	MG	2A	3515	1/1	0.86	0.14	55,55,55,55	0
56	MG	1A	3326	1/1	0.86	0.10	62,62,62,62	0
56	MG	1a	1801	1/1	0.86	0.12	65,65,65,65	0
56	MG	1A	3421	1/1	0.86	0.20	38,38,38,38	0
56	MG	2A	3186	1/1	0.86	0.23	59,59,59,59	0
56	MG	1E	301	1/1	0.86	0.31	40,40,40,40	0
56	MG	1E	304	1/1	0.86	0.15	65,65,65,65	0
56	MG	1a	1694	1/1	0.86	0.20	51,51,51,51	0
56	MG	2A	3208	1/1	0.86	0.11	48,48,48,48	0
56	MG	1A	3038	1/1	0.86	0.16	55,55,55,55	0
56	MG	2a	3199	1/1	0.86	0.25	64,64,64,64	0
56	MG	1A	3876	1/1	0.86	0.10	33,33,33,33	0
56	MG	1F	307	1/1	0.86	0.29	59,59,59,59	0
56	MG	1A	3435	1/1	0.86	0.15	66,66,66,66	0
56	MG	2a	3038	1/1	0.86	0.24	58,58,58,58	0
56	MG	1a	1733	1/1	0.86	0.22	52,52,52,52	0
56	MG	1a	1736	1/1	0.86	0.16	51,51,51,51	0
56	MG	2A	3258	1/1	0.86	0.20	62,62,62,62	0
56	MG	2a	3228	1/1	0.86	0.21	58,58,58,58	0
56	MG	1A	3337	1/1	0.86	0.14	44,44,44,44	0
56	MG	1A	3897	1/1	0.86	0.19	60,60,60,60	0
56	MG	2A	3274	1/1	0.86	0.10	37,37,37,37	0
56	MG	1A	4038	1/1	0.86	0.13	54,54,54,54	0
56	MG	1A	3446	1/1	0.86	0.13	58,58,58,58	0
56	MG	2a	3253	1/1	0.86	0.17	54,54,54,54	0
56	MG	1A	4044	1/1	0.86	0.29	44,44,44,44	0
56	MG	2a	3056	1/1	0.86	0.16	58,58,58,58	0
56	MG	1a	1873	1/1	0.86	0.14	60,60,60,60	0
56	MG	1a	1750	1/1	0.86	0.22	69,69,69,69	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1R	204	1/1	0.86	0.19	70,70,70,70	0
56	MG	2A	3342	1/1	0.86	0.17	61,61,61,61	0
56	MG	2A	3359	1/1	0.86	0.20	57,57,57,57	0
56	MG	1A	3512	1/1	0.87	0.15	50,50,50,50	0
56	MG	1a	1775	1/1	0.87	0.09	65,65,65,65	0
56	MG	1a	1864	1/1	0.87	0.30	69,69,69,69	0
56	MG	1a	1686	1/1	0.87	0.18	62,62,62,62	0
56	MG	1a	1692	1/1	0.87	0.24	64,64,64,64	0
56	MG	2a	3135	1/1	0.87	0.10	73,73,73,73	0
56	MG	1A	3322	1/1	0.87	0.18	53,53,53,53	0
56	MG	2A	3495	1/1	0.87	0.18	60,60,60,60	0
56	MG	2A	3217	1/1	0.87	0.08	41,41,41,41	0
56	MG	1A	3083	1/1	0.87	0.10	45,45,45,45	0
56	MG	1a	1716	1/1	0.87	0.22	54,54,54,54	0
56	MG	1A	3278	1/1	0.87	0.21	50,50,50,50	0
56	MG	1A	3084	1/1	0.87	0.17	53,53,53,53	0
56	MG	2A	3526	1/1	0.87	0.10	50,50,50,50	0
56	MG	2A	3547	1/1	0.87	0.15	52,52,52,52	0
56	MG	1a	1729	1/1	0.87	0.22	75,75,75,75	0
56	MG	19	102	1/1	0.87	0.13	50,50,50,50	0
56	MG	1A	3432	1/1	0.87	0.11	53,53,53,53	0
56	MG	2B	208	1/1	0.87	0.14	58,58,58,58	0
56	MG	2B	210	1/1	0.87	0.15	65,65,65,65	0
56	MG	1e	203	1/1	0.87	0.13	84,84,84,84	0
56	MG	2A	3265	1/1	0.87	0.12	47,47,47,47	0
56	MG	1a	1796	1/1	0.87	0.10	85,85,85,85	0
56	MG	1A	3050	1/1	0.87	0.10	48,48,48,48	0
56	MG	1A	3024	1/1	0.87	0.15	56,56,56,56	0
56	MG	2a	3009	1/1	0.87	0.14	84,84,84,84	0
56	MG	1a	1608	1/1	0.87	0.10	70,70,70,70	0
56	MG	2A	3009	1/1	0.87	0.12	50,50,50,50	0
56	MG	2a	3020	1/1	0.87	0.19	44,44,44,44	0
56	MG	2a	3022	1/1	0.87	0.16	77,77,77,77	0
56	MG	2A	3011	1/1	0.87	0.08	48,48,48,48	0
56	MG	1a	1610	1/1	0.87	0.16	69,69,69,69	0
56	MG	1A	3758	1/1	0.87	0.19	49,49,49,49	0
56	MG	1A	3237	1/1	0.87	0.12	58,58,58,58	0
56	MG	2a	3042	1/1	0.87	0.13	72,72,72,72	0
56	MG	2A	3073	1/1	0.87	0.08	38,38,38,38	0
56	MG	2a	3203	1/1	0.87	0.15	58,58,58,58	0
56	MG	2a	3044	1/1	0.87	0.12	71,71,71,71	0
56	MG	1A	3865	1/1	0.87	0.14	72,72,72,72	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3077	1/1	0.87	0.12	43,43,43,43	0
56	MG	2a	3212	1/1	0.87	0.14	46,46,46,46	0
56	MG	1a	1813	1/1	0.87	0.18	75,75,75,75	0
56	MG	1a	1814	1/1	0.87	0.14	67,67,67,67	0
56	MG	1A	4048	1/1	0.87	0.13	48,48,48,48	0
56	MG	1a	1640	1/1	0.87	0.27	78,78,78,78	0
56	MG	2A	3102	1/1	0.87	0.11	51,51,51,51	0
56	MG	1a	1643	1/1	0.87	0.15	68,68,68,68	0
56	MG	1D	306	1/1	0.87	0.17	49,49,49,49	0
56	MG	2A	3410	1/1	0.87	0.12	52,52,52,52	0
56	MG	1a	1760	1/1	0.87	0.21	79,79,79,79	0
56	MG	2A	3130	1/1	0.87	0.15	49,49,49,49	0
56	MG	2A	3435	1/1	0.87	0.14	46,46,46,46	0
56	MG	1a	1825	1/1	0.87	0.10	73,73,73,73	0
56	MG	2a	3100	1/1	0.87	0.36	80,80,80,80	0
56	MG	2A	3147	1/1	0.87	0.18	62,62,62,62	0
56	MG	2A	3151	1/1	0.87	0.27	54,54,54,54	0
56	MG	1A	3356	1/1	0.87	0.10	63,63,63,63	0
56	MG	1A	3879	1/1	0.87	0.14	49,49,49,49	0
56	MG	1a	1767	1/1	0.87	0.10	71,71,71,71	0
56	MG	2A	3057	1/1	0.88	0.09	52,52,52,52	0
56	MG	2A	3372	1/1	0.88	0.10	33,33,33,33	0
56	MG	1R	202	1/1	0.88	0.42	55,55,55,55	0
56	MG	1A	3464	1/1	0.88	0.11	57,57,57,57	0
56	MG	1A	4047	1/1	0.88	0.24	47,47,47,47	0
56	MG	2A	3392	1/1	0.88	0.10	55,55,55,55	0
56	MG	2A	3399	1/1	0.88	0.26	64,64,64,64	0
56	MG	1T	203	1/1	0.88	0.12	75,75,75,75	0
56	MG	2a	3092	1/1	0.88	0.21	62,62,62,62	0
56	MG	2a	3093	1/1	0.88	0.27	66,66,66,66	0
56	MG	1A	3556	1/1	0.88	0.22	58,58,58,58	0
56	MG	1A	3104	1/1	0.88	0.16	52,52,52,52	0
56	MG	2A	3409	1/1	0.88	0.14	64,64,64,64	0
56	MG	1A	3587	1/1	0.88	0.20	42,42,42,42	0
56	MG	2A	3415	1/1	0.88	0.23	57,57,57,57	0
56	MG	2A	3425	1/1	0.88	0.10	48,48,48,48	0
56	MG	1A	3899	1/1	0.88	0.16	77,77,77,77	0
56	MG	1A	3472	1/1	0.88	0.11	53,53,53,53	0
56	MG	1A	3614	1/1	0.88	0.20	57,57,57,57	0
56	MG	2A	3124	1/1	0.88	0.29	65,65,65,65	0
56	MG	1A	3239	1/1	0.88	0.10	50,50,50,50	0
56	MG	1A	3643	1/1	0.88	0.13	49,49,49,49	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1B	208	1/1	0.88	0.25	63,63,63,63	0
56	MG	2A	3463	1/1	0.88	0.18	44,44,44,44	0
56	MG	2A	3135	1/1	0.88	0.11	43,43,43,43	0
56	MG	2A	3141	1/1	0.88	0.21	49,49,49,49	0
56	MG	1B	209	1/1	0.88	0.19	56,56,56,56	0
56	MG	2a	3147	1/1	0.88	0.14	68,68,68,68	0
56	MG	1a	1755	1/1	0.88	0.13	57,57,57,57	0
56	MG	2A	3153	1/1	0.88	0.10	52,52,52,52	0
56	MG	1A	3074	1/1	0.88	0.18	63,63,63,63	0
56	MG	1A	3370	1/1	0.88	0.09	49,49,49,49	0
56	MG	1A	3263	1/1	0.88	0.12	42,42,42,42	0
56	MG	1A	3699	1/1	0.88	0.20	58,58,58,58	0
56	MG	2A	3181	1/1	0.88	0.10	48,48,48,48	0
56	MG	1A	3969	1/1	0.88	0.16	77,77,77,77	0
56	MG	1A	3355	1/1	0.88	0.09	64,64,64,64	0
56	MG	1a	1874	1/1	0.88	0.11	48,48,48,48	0
56	MG	2A	3514	1/1	0.88	0.10	71,71,71,71	0
56	MG	1a	1772	1/1	0.88	0.20	62,62,62,62	0
56	MG	2A	3516	1/1	0.88	0.09	53,53,53,53	0
56	MG	1a	1892	1/1	0.88	0.12	58,58,58,58	0
56	MG	2A	3214	1/1	0.88	0.35	45,45,45,45	0
56	MG	1A	3387	1/1	0.88	0.13	54,54,54,54	0
56	MG	1A	3496	1/1	0.88	0.16	42,42,42,42	0
56	MG	1A	3998	1/1	0.88	0.16	59,59,59,59	0
56	MG	2B	205	1/1	0.88	0.32	59,59,59,59	0
56	MG	1E	302	1/1	0.88	0.17	41,41,41,41	0
56	MG	2A	3244	1/1	0.88	0.21	85,85,85,85	0
56	MG	2a	3192	1/1	0.88	0.18	61,61,61,61	0
56	MG	2D	301	1/1	0.88	0.12	49,49,49,49	0
56	MG	2D	302	1/1	0.88	0.09	57,57,57,57	0
56	MG	1A	3521	1/1	0.88	0.13	46,46,46,46	0
56	MG	1A	3754	1/1	0.88	0.24	39,39,39,39	0
56	MG	1A	3500	1/1	0.88	0.11	53,53,53,53	0
56	MG	1A	3838	1/1	0.88	0.14	64,64,64,64	0
56	MG	2a	3005	1/1	0.88	0.07	72,72,72,72	0
56	MG	2a	3209	1/1	0.88	0.12	53,53,53,53	0
56	MG	1A	3336	1/1	0.88	0.17	58,58,58,58	0
56	MG	1a	1671	1/1	0.88	0.10	77,77,77,77	0
56	MG	2a	3214	1/1	0.88	0.12	73,73,73,73	0
56	MG	2A	3270	1/1	0.88	0.13	59,59,59,59	0
56	MG	1q	202	1/1	0.88	0.17	53,53,53,53	0
56	MG	1a	1678	1/1	0.88	0.18	51,51,51,51	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2a	3225	1/1	0.88	0.15	59,59,59,59	0
56	MG	1x	108	1/1	0.88	0.23	49,49,49,49	0
56	MG	2a	3030	1/1	0.88	0.12	51,51,51,51	0
56	MG	1x	110	1/1	0.88	0.08	73,73,73,73	0
56	MG	2A	3279	1/1	0.88	0.15	51,51,51,51	0
56	MG	2A	3294	1/1	0.88	0.22	50,50,50,50	0
56	MG	2A	3006	1/1	0.88	0.15	59,59,59,59	0
56	MG	1A	3860	1/1	0.88	0.10	38,38,38,38	0
56	MG	2a	3252	1/1	0.88	0.15	67,67,67,67	0
56	MG	2A	3010	1/1	0.88	0.23	68,68,68,68	0
56	MG	1A	3532	1/1	0.88	0.08	62,62,62,62	0
56	MG	2A	3015	1/1	0.88	0.15	57,57,57,57	0
56	MG	2A	3345	1/1	0.88	0.14	60,60,60,60	0
56	MG	1A	3873	1/1	0.88	0.13	41,41,41,41	0
56	MG	1A	4039	1/1	0.88	0.12	56,56,56,56	0
56	MG	1A	3544	1/1	0.88	0.12	61,61,61,61	0
56	MG	2A	3368	1/1	0.88	0.23	48,48,48,48	0
56	MG	1A	3443	1/1	0.89	0.14	72,72,72,72	0
56	MG	1A	3590	1/1	0.89	0.12	51,51,51,51	0
56	MG	2A	3344	1/1	0.89	0.22	59,59,59,59	0
56	MG	2A	3012	1/1	0.89	0.21	71,71,71,71	0
56	MG	2A	3353	1/1	0.89	0.14	41,41,41,41	0
56	MG	1A	3020	1/1	0.89	0.19	40,40,40,40	0
56	MG	2A	3021	1/1	0.89	0.18	58,58,58,58	0
56	MG	1A	3123	1/1	0.89	0.16	49,49,49,49	0
56	MG	1a	1688	1/1	0.89	0.16	61,61,61,61	0
56	MG	2A	3040	1/1	0.89	0.17	57,57,57,57	0
56	MG	1A	3448	1/1	0.89	0.09	56,56,56,56	0
56	MG	1A	3339	1/1	0.89	0.16	56,56,56,56	0
56	MG	2a	3064	1/1	0.89	0.13	79,79,79,79	0
56	MG	2a	3069	1/1	0.89	0.14	65,65,65,65	0
56	MG	2A	3065	1/1	0.89	0.17	55,55,55,55	0
56	MG	2A	3388	1/1	0.89	0.13	55,55,55,55	0
56	MG	1A	3396	1/1	0.89	0.16	55,55,55,55	0
56	MG	1A	3908	1/1	0.89	0.12	51,51,51,51	0
56	MG	2A	3075	1/1	0.89	0.18	48,48,48,48	0
56	MG	1a	1803	1/1	0.89	0.11	66,66,66,66	0
56	MG	2A	3082	1/1	0.89	0.14	57,57,57,57	0
56	MG	2a	3106	1/1	0.89	0.23	52,52,52,52	0
56	MG	1A	3235	1/1	0.89	0.19	53,53,53,53	0
56	MG	1a	1810	1/1	0.89	0.08	59,59,59,59	0
56	MG	1A	3403	1/1	0.89	0.23	50,50,50,50	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1726	1/1	0.89	0.14	68,68,68,68	0
56	MG	1A	3404	1/1	0.89	0.20	54,54,54,54	0
56	MG	1A	4074	1/1	0.89	0.14	47,47,47,47	0
56	MG	1A	3107	1/1	0.89	0.18	53,53,53,53	0
56	MG	2a	3128	1/1	0.89	0.20	62,62,62,62	0
56	MG	10	101	1/1	0.89	0.17	49,49,49,49	0
56	MG	1A	3316	1/1	0.89	0.19	51,51,51,51	0
56	MG	2A	3129	1/1	0.89	0.08	33,33,33,33	0
56	MG	1B	202	1/1	0.89	0.20	63,63,63,63	0
56	MG	2A	3452	1/1	0.89	0.12	60,60,60,60	0
56	MG	1a	1740	1/1	0.89	0.23	63,63,63,63	0
56	MG	2A	3134	1/1	0.89	0.11	44,44,44,44	0
56	MG	1a	1741	1/1	0.89	0.08	48,48,48,48	0
56	MG	2A	3469	1/1	0.89	0.09	49,49,49,49	0
56	MG	2A	3471	1/1	0.89	0.20	63,63,63,63	0
56	MG	1A	3059	1/1	0.89	0.21	32,32,32,32	0
56	MG	1a	1839	1/1	0.89	0.20	53,53,53,53	0
56	MG	2A	3149	1/1	0.89	0.14	63,63,63,63	0
56	MG	1A	3319	1/1	0.89	0.17	75,75,75,75	0
56	MG	2A	3485	1/1	0.89	0.11	56,56,56,56	0
56	MG	1A	3954	1/1	0.89	0.10	16,16,16,16	0
56	MG	1a	1749	1/1	0.89	0.12	64,64,64,64	0
56	MG	1A	3320	1/1	0.89	0.19	70,70,70,70	0
56	MG	1A	3481	1/1	0.89	0.23	46,46,46,46	0
56	MG	1A	3764	1/1	0.89	0.12	68,68,68,68	0
56	MG	1A	3769	1/1	0.89	0.19	55,55,55,55	0
56	MG	1a	1606	1/1	0.89	0.14	71,71,71,71	0
56	MG	1a	1878	1/1	0.89	0.15	67,67,67,67	0
56	MG	1a	1879	1/1	0.89	0.10	52,52,52,52	0
56	MG	1A	3991	1/1	0.89	0.30	38,38,38,38	0
56	MG	1a	1881	1/1	0.89	0.19	52,52,52,52	0
56	MG	2A	3519	1/1	0.89	0.13	52,52,52,52	0
56	MG	2a	3186	1/1	0.89	0.13	50,50,50,50	0
56	MG	1a	1886	1/1	0.89	0.13	56,56,56,56	0
56	MG	2A	3535	1/1	0.89	0.14	52,52,52,52	0
56	MG	2A	3537	1/1	0.89	0.14	47,47,47,47	0
56	MG	1a	1887	1/1	0.89	0.14	47,47,47,47	0
56	MG	2A	3557	1/1	0.89	0.10	54,54,54,54	0
56	MG	2A	3223	1/1	0.89	0.12	70,70,70,70	0
56	MG	1A	3119	1/1	0.89	0.18	59,59,59,59	0
56	MG	1a	1898	1/1	0.89	0.10	59,59,59,59	0
56	MG	2a	3202	1/1	0.89	0.13	52,52,52,52	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3234	1/1	0.89	0.08	65,65,65,65	0
56	MG	1A	3427	1/1	0.89	0.11	51,51,51,51	0
56	MG	1a	1762	1/1	0.89	0.28	71,71,71,71	0
56	MG	1D	311	1/1	0.89	0.10	51,51,51,51	0
56	MG	2A	3247	1/1	0.89	0.14	59,59,59,59	0
56	MG	1a	1626	1/1	0.89	0.17	68,68,68,68	0
56	MG	2a	3213	1/1	0.89	0.14	71,71,71,71	0
56	MG	1A	3855	1/1	0.89	0.13	48,48,48,48	0
56	MG	1A	3857	1/1	0.89	0.14	51,51,51,51	0
56	MG	1a	1774	1/1	0.89	0.13	62,62,62,62	0
56	MG	1A	3259	1/1	0.89	0.13	53,53,53,53	0
56	MG	1k	201	1/1	0.89	0.10	49,49,49,49	0
56	MG	1A	3187	1/1	0.89	0.23	52,52,52,52	0
56	MG	1a	1777	1/1	0.89	0.12	54,54,54,54	0
56	MG	2a	3017	1/1	0.89	0.19	61,61,61,61	0
56	MG	1A	3868	1/1	0.89	0.10	84,84,84,84	0
56	MG	1A	4036	1/1	0.89	0.10	54,54,54,54	0
56	MG	1a	1650	1/1	0.89	0.26	49,49,49,49	0
56	MG	2a	3023	1/1	0.89	0.17	66,66,66,66	0
56	MG	2a	3025	1/1	0.89	0.18	60,60,60,60	0
56	MG	2A	3281	1/1	0.89	0.09	45,45,45,45	0
56	MG	2A	3288	1/1	0.89	0.09	41,41,41,41	0
56	MG	2d	503	1/1	0.89	0.08	62,62,62,62	0
56	MG	1a	1655	1/1	0.89	0.24	60,60,60,60	0
56	MG	2a	3036	1/1	0.89	0.09	59,59,59,59	0
56	MG	2A	3003	1/1	0.89	0.10	43,43,43,43	0
56	MG	1A	3373	1/1	0.89	0.14	44,44,44,44	0
56	MG	2t	201	1/1	0.89	0.16	58,58,58,58	0
56	MG	2A	3318	1/1	0.89	0.09	43,43,43,43	0
56	MG	1a	1661	1/1	0.89	0.09	59,59,59,59	0
56	MG	2A	3343	1/1	0.90	0.11	55,55,55,55	0
56	MG	1a	1680	1/1	0.90	0.13	55,55,55,55	0
56	MG	2A	3052	1/1	0.90	0.14	37,37,37,37	0
56	MG	1a	1685	1/1	0.90	0.17	55,55,55,55	0
56	MG	2A	3357	1/1	0.90	0.16	52,52,52,52	0
56	MG	2A	3055	1/1	0.90	0.15	35,35,35,35	0
56	MG	2a	3052	1/1	0.90	0.09	81,81,81,81	0
56	MG	1A	3558	1/1	0.90	0.22	43,43,43,43	0
56	MG	1A	4023	1/1	0.90	0.17	52,52,52,52	0
56	MG	1A	3856	1/1	0.90	0.11	55,55,55,55	0
56	MG	1a	1693	1/1	0.90	0.11	53,53,53,53	0
56	MG	1A	3567	1/1	0.90	0.10	40,40,40,40	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1696	1/1	0.90	0.13	48,48,48,48	0
56	MG	2a	3060	1/1	0.90	0.10	81,81,81,81	0
56	MG	1a	1700	1/1	0.90	0.13	60,60,60,60	0
56	MG	1A	4029	1/1	0.90	0.14	42,42,42,42	0
56	MG	2a	3065	1/1	0.90	0.11	93,93,93,93	0
56	MG	2a	3068	1/1	0.90	0.09	56,56,56,56	0
56	MG	2A	3089	1/1	0.90	0.17	36,36,36,36	0
56	MG	1N	3007	1/1	0.90	0.14	54,54,54,54	0
56	MG	1A	3418	1/1	0.90	0.10	48,48,48,48	0
56	MG	1Q	202	1/1	0.90	0.16	32,32,32,32	0
56	MG	1A	3272	1/1	0.90	0.20	58,58,58,58	0
56	MG	1A	3462	1/1	0.90	0.10	47,47,47,47	0
56	MG	1a	1834	1/1	0.90	0.14	57,57,57,57	0
56	MG	1a	1837	1/1	0.90	0.09	48,48,48,48	0
56	MG	2A	3412	1/1	0.90	0.20	54,54,54,54	0
56	MG	1A	3870	1/1	0.90	0.09	44,44,44,44	0
56	MG	1A	3599	1/1	0.90	0.09	38,38,38,38	0
56	MG	1A	3423	1/1	0.90	0.10	42,42,42,42	0
56	MG	1A	3878	1/1	0.90	0.08	37,37,37,37	0
56	MG	1a	1858	1/1	0.90	0.09	56,56,56,56	0
56	MG	1a	1859	1/1	0.90	0.17	59,59,59,59	0
56	MG	1a	1860	1/1	0.90	0.14	61,61,61,61	0
56	MG	1U	205	1/1	0.90	0.35	41,41,41,41	0
56	MG	1W	3006	1/1	0.90	0.19	40,40,40,40	0
56	MG	1A	3273	1/1	0.90	0.11	37,37,37,37	0
56	MG	1A	3350	1/1	0.90	0.19	60,60,60,60	0
56	MG	2a	3136	1/1	0.90	0.21	56,56,56,56	0
56	MG	2A	3160	1/1	0.90	0.16	49,49,49,49	0
56	MG	1A	3884	1/1	0.90	0.23	58,58,58,58	0
56	MG	2A	3167	1/1	0.90	0.14	45,45,45,45	0
56	MG	2A	3168	1/1	0.90	0.10	33,33,33,33	0
56	MG	1A	3889	1/1	0.90	0.15	55,55,55,55	0
56	MG	1A	3184	1/1	0.90	0.27	52,52,52,52	0
56	MG	1A	3478	1/1	0.90	0.14	49,49,49,49	0
56	MG	2A	3178	1/1	0.90	0.14	54,54,54,54	0
56	MG	1A	3030	1/1	0.90	0.09	46,46,46,46	0
56	MG	2A	3487	1/1	0.90	0.06	40,40,40,40	0
56	MG	13	101	1/1	0.90	0.15	47,47,47,47	0
56	MG	1A	3436	1/1	0.90	0.19	51,51,51,51	0
56	MG	2A	3192	1/1	0.90	0.19	43,43,43,43	0
56	MG	15	104	1/1	0.90	0.14	53,53,53,53	0
56	MG	1a	1894	1/1	0.90	0.28	63,63,63,63	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3504	1/1	0.90	0.14	42,42,42,42	0
56	MG	2A	3198	1/1	0.90	0.08	56,56,56,56	0
56	MG	2a	3163	1/1	0.90	0.12	94,94,94,94	0
56	MG	1a	1895	1/1	0.90	0.26	59,59,59,59	0
56	MG	1A	3713	1/1	0.90	0.08	61,61,61,61	0
56	MG	1A	3289	1/1	0.90	0.10	61,61,61,61	0
56	MG	2A	3219	1/1	0.90	0.27	56,56,56,56	0
56	MG	2A	3220	1/1	0.90	0.12	72,72,72,72	0
56	MG	2A	3525	1/1	0.90	0.22	36,36,36,36	0
56	MG	2A	3221	1/1	0.90	0.09	39,39,39,39	0
56	MG	1a	1601	1/1	0.90	0.14	63,63,63,63	0
56	MG	1A	4078	1/1	0.90	0.20	65,65,65,65	0
56	MG	1A	3488	1/1	0.90	0.10	43,43,43,43	0
56	MG	2A	3549	1/1	0.90	0.13	49,49,49,49	0
56	MG	1A	3733	1/1	0.90	0.18	49,49,49,49	0
56	MG	2A	3558	1/1	0.90	0.32	43,43,43,43	0
56	MG	1B	206	1/1	0.90	0.15	51,51,51,51	0
56	MG	2A	3241	1/1	0.90	0.14	56,56,56,56	0
56	MG	1A	3926	1/1	0.90	0.14	51,51,51,51	0
56	MG	2a	3200	1/1	0.90	0.12	71,71,71,71	0
56	MG	1A	3094	1/1	0.90	0.10	54,54,54,54	0
56	MG	1A	3335	1/1	0.90	0.34	58,58,58,58	0
56	MG	2a	3204	1/1	0.90	0.08	57,57,57,57	0
56	MG	1A	3936	1/1	0.90	0.18	33,33,33,33	0
56	MG	1A	3533	1/1	0.90	0.14	56,56,56,56	0
56	MG	1p	3002	1/1	0.90	0.14	77,77,77,77	0
56	MG	1B	221	1/1	0.90	0.10	30,30,30,30	0
56	MG	2O	201	1/1	0.90	0.13	54,54,54,54	0
56	MG	2Q	3001	1/1	0.90	0.12	45,45,45,45	0
56	MG	2X	8001	1/1	0.90	0.11	45,45,45,45	0
56	MG	2a	3216	1/1	0.90	0.14	55,55,55,55	0
56	MG	1a	1628	1/1	0.90	0.13	57,57,57,57	0
56	MG	1x	105	1/1	0.90	0.18	64,64,64,64	0
56	MG	1x	106	1/1	0.90	0.09	42,42,42,42	0
56	MG	1A	3540	1/1	0.90	0.31	48,48,48,48	0
56	MG	1A	3063	1/1	0.90	0.13	57,57,57,57	0
56	MG	1A	3313	1/1	0.90	0.08	42,42,42,42	0
56	MG	2a	3012	1/1	0.90	0.16	58,58,58,58	0
56	MG	1A	3771	1/1	0.90	0.14	55,55,55,55	0
56	MG	2a	3231	1/1	0.90	0.19	59,59,59,59	0
56	MG	2A	3007	1/1	0.90	0.14	77,77,77,77	0
56	MG	2a	3235	1/1	0.90	0.15	73,73,73,73	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2a	3236	1/1	0.90	0.23	83,83,83,83	0
56	MG	1A	3793	1/1	0.90	0.11	55,55,55,55	0
56	MG	1A	3983	1/1	0.90	0.13	31,31,31,31	0
56	MG	2a	3250	1/1	0.90	0.13	63,63,63,63	0
56	MG	1A	3794	1/1	0.90	0.09	19,19,19,19	0
56	MG	2A	3304	1/1	0.90	0.09	66,66,66,66	0
56	MG	1A	3090	1/1	0.90	0.14	47,47,47,47	0
56	MG	2a	3255	1/1	0.90	0.09	57,57,57,57	0
56	MG	2d	502	1/1	0.90	0.31	61,61,61,61	0
56	MG	2a	3027	1/1	0.90	0.16	47,47,47,47	0
56	MG	2A	3314	1/1	0.90	0.11	43,43,43,43	0
56	MG	2A	3316	1/1	0.90	0.11	55,55,55,55	0
56	MG	1A	3845	1/1	0.90	0.42	35,35,35,35	0
56	MG	2A	3329	1/1	0.90	0.15	55,55,55,55	0
56	MG	1A	3999	1/1	0.90	0.17	51,51,51,51	0
56	MG	2v	101	1/1	0.90	0.12	65,65,65,65	0
56	MG	2x	101	1/1	0.90	0.12	72,72,72,72	0
56	MG	2x	104	1/1	0.90	0.22	84,84,84,84	0
56	MG	1F	302	1/1	0.90	0.17	39,39,39,39	0
56	MG	1A	3376	1/1	0.90	0.10	49,49,49,49	0
56	MG	1A	3516	1/1	0.91	0.23	52,52,52,52	0
56	MG	2a	3026	1/1	0.91	0.20	54,54,54,54	0
56	MG	2A	3259	1/1	0.91	0.14	39,39,39,39	0
56	MG	1A	3997	1/1	0.91	0.08	32,32,32,32	0
56	MG	1A	3162	1/1	0.91	0.21	38,38,38,38	0
56	MG	1A	3397	1/1	0.91	0.09	36,36,36,36	0
56	MG	1R	203	1/1	0.91	0.08	35,35,35,35	0
56	MG	1A	3400	1/1	0.91	0.11	37,37,37,37	0
56	MG	1A	4002	1/1	0.91	0.14	65,65,65,65	0
56	MG	2a	3040	1/1	0.91	0.16	58,58,58,58	0
56	MG	1a	1748	1/1	0.91	0.15	55,55,55,55	0
56	MG	1A	4010	1/1	0.91	0.13	37,37,37,37	0
56	MG	1A	3792	1/1	0.91	0.10	62,62,62,62	0
56	MG	1U	202	1/1	0.91	0.27	51,51,51,51	0
56	MG	1A	3076	1/1	0.91	0.10	51,51,51,51	0
56	MG	1A	3276	1/1	0.91	0.17	51,51,51,51	0
56	MG	2A	3301	1/1	0.91	0.06	42,42,42,42	0
56	MG	1A	4026	1/1	0.91	0.23	63,63,63,63	0
56	MG	1A	3827	1/1	0.91	0.12	30,30,30,30	0
56	MG	2A	3005	1/1	0.91	0.10	59,59,59,59	0
56	MG	1A	3835	1/1	0.91	0.10	47,47,47,47	0
56	MG	1A	3343	1/1	0.91	0.10	55,55,55,55	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3328	1/1	0.91	0.13	70,70,70,70	0
56	MG	10	103	1/1	0.91	0.24	62,62,62,62	0
56	MG	1A	3535	1/1	0.91	0.14	59,59,59,59	0
56	MG	1a	1765	1/1	0.91	0.11	58,58,58,58	0
56	MG	2a	3063	1/1	0.91	0.07	85,85,85,85	0
56	MG	1A	4035	1/1	0.91	0.11	46,46,46,46	0
56	MG	1a	1769	1/1	0.91	0.13	63,63,63,63	0
56	MG	2A	3017	1/1	0.91	0.23	46,46,46,46	0
56	MG	1A	3467	1/1	0.91	0.15	54,54,54,54	0
56	MG	2A	3348	1/1	0.91	0.14	60,60,60,60	0
56	MG	2A	3026	1/1	0.91	0.17	49,49,49,49	0
56	MG	2a	3075	1/1	0.91	0.06	59,59,59,59	0
56	MG	2A	3029	1/1	0.91	0.10	57,57,57,57	0
56	MG	2a	3088	1/1	0.91	0.12	76,76,76,76	0
56	MG	1A	3849	1/1	0.91	0.09	52,52,52,52	0
56	MG	2A	3031	1/1	0.91	0.08	47,47,47,47	0
56	MG	2a	3096	1/1	0.91	0.15	55,55,55,55	0
56	MG	1A	3103	1/1	0.91	0.24	32,32,32,32	0
56	MG	1A	3055	1/1	0.91	0.23	45,45,45,45	0
56	MG	2A	3041	1/1	0.91	0.08	41,41,41,41	0
56	MG	1A	3285	1/1	0.91	0.16	32,32,32,32	0
56	MG	1A	3858	1/1	0.91	0.23	53,53,53,53	0
56	MG	2a	3115	1/1	0.91	0.21	50,50,50,50	0
56	MG	1A	3035	1/1	0.91	0.09	44,44,44,44	0
56	MG	1A	3206	1/1	0.91	0.23	45,45,45,45	0
56	MG	2a	3119	1/1	0.91	0.10	69,69,69,69	0
56	MG	1A	3866	1/1	0.91	0.17	42,42,42,42	0
56	MG	1A	3114	1/1	0.91	0.13	48,48,48,48	0
56	MG	2a	3124	1/1	0.91	0.15	62,62,62,62	0
56	MG	2A	3396	1/1	0.91	0.14	49,49,49,49	0
56	MG	1A	4060	1/1	0.91	0.08	31,31,31,31	0
56	MG	1A	3579	1/1	0.91	0.13	41,41,41,41	0
56	MG	1A	3486	1/1	0.91	0.07	57,57,57,57	0
56	MG	2A	3081	1/1	0.91	0.18	54,54,54,54	0
56	MG	1a	1611	1/1	0.91	0.07	46,46,46,46	0
56	MG	1A	4064	1/1	0.91	0.15	57,57,57,57	0
56	MG	1a	1619	1/1	0.91	0.07	75,75,75,75	0
56	MG	2A	3091	1/1	0.91	0.12	49,49,49,49	0
56	MG	2A	3092	1/1	0.91	0.19	48,48,48,48	0
56	MG	1a	1620	1/1	0.91	0.17	54,54,54,54	0
56	MG	2A	3098	1/1	0.91	0.13	41,41,41,41	0
56	MG	1a	1622	1/1	0.91	0.08	49,49,49,49	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3588	1/1	0.91	0.13	60,60,60,60	0
56	MG	1A	3877	1/1	0.91	0.13	39,39,39,39	0
56	MG	1A	3419	1/1	0.91	0.21	52,52,52,52	0
56	MG	2A	3106	1/1	0.91	0.11	47,47,47,47	0
56	MG	2A	3107	1/1	0.91	0.14	49,49,49,49	0
56	MG	2a	3155	1/1	0.91	0.14	87,87,87,87	0
56	MG	2A	3457	1/1	0.91	0.15	46,46,46,46	0
56	MG	2A	3109	1/1	0.91	0.18	54,54,54,54	0
56	MG	2A	3466	1/1	0.91	0.09	45,45,45,45	0
56	MG	1A	3229	1/1	0.91	0.13	50,50,50,50	0
56	MG	1a	1629	1/1	0.91	0.17	63,63,63,63	0
56	MG	1a	1633	1/1	0.91	0.07	77,77,77,77	0
56	MG	1A	3422	1/1	0.91	0.08	40,40,40,40	0
56	MG	2A	3478	1/1	0.91	0.10	50,50,50,50	0
56	MG	1A	3367	1/1	0.91	0.10	48,48,48,48	0
56	MG	2A	3481	1/1	0.91	0.10	46,46,46,46	0
56	MG	1a	1641	1/1	0.91	0.21	47,47,47,47	0
56	MG	1B	205	1/1	0.91	0.19	56,56,56,56	0
56	MG	2a	3173	1/1	0.91	0.09	43,43,43,43	0
56	MG	1A	3616	1/1	0.91	0.09	29,29,29,29	0
56	MG	2a	3177	1/1	0.91	0.20	69,69,69,69	0
56	MG	2a	3183	1/1	0.91	0.14	60,60,60,60	0
56	MG	2A	3136	1/1	0.91	0.15	61,61,61,61	0
56	MG	1A	3010	1/1	0.91	0.11	66,66,66,66	0
56	MG	1a	1646	1/1	0.91	0.14	80,80,80,80	0
56	MG	1A	3631	1/1	0.91	0.11	43,43,43,43	0
56	MG	1a	1824	1/1	0.91	0.12	54,54,54,54	0
56	MG	1a	1653	1/1	0.91	0.23	56,56,56,56	0
56	MG	2A	3159	1/1	0.91	0.16	49,49,49,49	0
56	MG	1A	3632	1/1	0.91	0.13	67,67,67,67	0
56	MG	1A	3497	1/1	0.91	0.15	55,55,55,55	0
56	MG	2A	3163	1/1	0.91	0.16	56,56,56,56	0
56	MG	2A	3512	1/1	0.91	0.14	38,38,38,38	0
56	MG	1A	3009	1/1	0.91	0.32	42,42,42,42	0
56	MG	1a	1662	1/1	0.91	0.14	60,60,60,60	0
56	MG	1a	1667	1/1	0.91	0.10	64,64,64,64	0
56	MG	2A	3518	1/1	0.91	0.10	32,32,32,32	0
56	MG	1A	3238	1/1	0.91	0.17	52,52,52,52	0
56	MG	2A	3174	1/1	0.91	0.13	56,56,56,56	0
56	MG	1a	1669	1/1	0.91	0.14	37,37,37,37	0
56	MG	1A	3666	1/1	0.91	0.07	25,25,25,25	0
56	MG	1B	225	1/1	0.91	0.14	63,63,63,63	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3184	1/1	0.91	0.08	45,45,45,45	0
56	MG	1A	3918	1/1	0.91	0.13	61,61,61,61	0
56	MG	1A	3091	1/1	0.91	0.27	54,54,54,54	0
56	MG	1A	3066	1/1	0.91	0.11	56,56,56,56	0
56	MG	2A	3193	1/1	0.91	0.10	37,37,37,37	0
56	MG	1a	1870	1/1	0.91	0.14	75,75,75,75	0
56	MG	2A	3196	1/1	0.91	0.10	47,47,47,47	0
56	MG	1A	3711	1/1	0.91	0.12	61,61,61,61	0
56	MG	1A	3439	1/1	0.91	0.07	50,50,50,50	0
56	MG	1A	3014	1/1	0.91	0.12	55,55,55,55	0
56	MG	2a	3232	1/1	0.91	0.23	54,54,54,54	0
56	MG	2A	3211	1/1	0.91	0.21	33,33,33,33	0
56	MG	1A	3940	1/1	0.91	0.20	65,65,65,65	0
56	MG	2A	3215	1/1	0.91	0.11	31,31,31,31	0
56	MG	1a	1876	1/1	0.91	0.19	62,62,62,62	0
56	MG	2a	3241	1/1	0.91	0.15	73,73,73,73	0
56	MG	2a	3242	1/1	0.91	0.11	73,73,73,73	0
56	MG	2O	202	1/1	0.91	0.09	37,37,37,37	0
56	MG	2a	3248	1/1	0.91	0.16	56,56,56,56	0
56	MG	1A	3095	1/1	0.91	0.12	46,46,46,46	0
56	MG	2Q	3005	1/1	0.91	0.09	54,54,54,54	0
56	MG	1a	1697	1/1	0.91	0.14	56,56,56,56	0
56	MG	1A	3271	1/1	0.91	0.20	54,54,54,54	0
56	MG	25	502	1/1	0.91	0.31	54,54,54,54	0
56	MG	1a	1704	1/1	0.91	0.23	48,48,48,48	0
56	MG	1a	1706	1/1	0.91	0.18	52,52,52,52	0
56	MG	2A	3226	1/1	0.91	0.11	42,42,42,42	0
56	MG	1A	3953	1/1	0.91	0.12	38,38,38,38	0
56	MG	1A	3390	1/1	0.91	0.09	52,52,52,52	0
56	MG	1F	306	1/1	0.91	0.20	59,59,59,59	0
56	MG	1A	3963	1/1	0.91	0.09	27,27,27,27	0
56	MG	1A	3391	1/1	0.91	0.10	51,51,51,51	0
56	MG	1A	3750	1/1	0.91	0.08	28,28,28,28	0
56	MG	1A	3751	1/1	0.91	0.14	74,74,74,74	0
56	MG	1A	3753	1/1	0.91	0.09	34,34,34,34	0
56	MG	1A	3395	1/1	0.91	0.12	46,46,46,46	0
56	MG	1A	3027	1/1	0.92	0.13	47,47,47,47	0
56	MG	2A	3255	1/1	0.92	0.13	43,43,43,43	0
56	MG	1A	3156	1/1	0.92	0.17	50,50,50,50	0
56	MG	11	101	1/1	0.92	0.11	50,50,50,50	0
56	MG	1A	3623	1/1	0.92	0.13	45,45,45,45	0
56	MG	1A	3329	1/1	0.92	0.20	53,53,53,53	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1r	3001	1/1	0.92	0.12	61,61,61,61	0
56	MG	1A	3120	1/1	0.92	0.10	59,59,59,59	0
56	MG	1r	3003	1/1	0.92	0.06	49,49,49,49	0
56	MG	1t	3001	1/1	0.92	0.17	53,53,53,53	0
56	MG	1x	103	1/1	0.92	0.15	48,48,48,48	0
56	MG	13	102	1/1	0.92	0.07	46,46,46,46	0
56	MG	1a	1754	1/1	0.92	0.20	51,51,51,51	0
56	MG	1A	3077	1/1	0.92	0.13	48,48,48,48	0
56	MG	1A	4050	1/1	0.92	0.26	84,84,84,84	0
56	MG	2A	3002	1/1	0.92	0.09	40,40,40,40	0
56	MG	1A	4053	1/1	0.92	0.29	52,52,52,52	0
56	MG	1a	1759	1/1	0.92	0.36	66,66,66,66	0
56	MG	1A	3372	1/1	0.92	0.10	50,50,50,50	0
56	MG	1A	3175	1/1	0.92	0.24	43,43,43,43	0
56	MG	1A	3375	1/1	0.92	0.08	44,44,44,44	0
56	MG	1A	3007	1/1	0.92	0.15	73,73,73,73	0
56	MG	1A	3378	1/1	0.92	0.14	35,35,35,35	0
56	MG	1A	3701	1/1	0.92	0.05	29,29,29,29	0
56	MG	2A	3320	1/1	0.92	0.17	53,53,53,53	0
56	MG	1A	3890	1/1	0.92	0.13	31,31,31,31	0
56	MG	1A	3704	1/1	0.92	0.08	45,45,45,45	0
56	MG	1A	3706	1/1	0.92	0.07	15,15,15,15	0
56	MG	1A	3476	1/1	0.92	0.20	56,56,56,56	0
56	MG	2a	3059	1/1	0.92	0.07	69,69,69,69	0
56	MG	1a	1616	1/1	0.92	0.21	55,55,55,55	0
56	MG	1a	1618	1/1	0.92	0.10	77,77,77,77	0
56	MG	1A	3527	1/1	0.92	0.20	69,69,69,69	0
56	MG	1A	4088	1/1	0.92	0.15	45,45,45,45	0
56	MG	1B	201	1/1	0.92	0.09	40,40,40,40	0
56	MG	2a	3066	1/1	0.92	0.19	64,64,64,64	0
56	MG	1A	3716	1/1	0.92	0.10	38,38,38,38	0
56	MG	1a	1625	1/1	0.92	0.06	34,34,34,34	0
56	MG	2A	3358	1/1	0.92	0.08	60,60,60,60	0
56	MG	1A	3719	1/1	0.92	0.09	56,56,56,56	0
56	MG	2a	3073	1/1	0.92	0.16	63,63,63,63	0
56	MG	1a	1787	1/1	0.92	0.13	52,52,52,52	0
56	MG	1A	3722	1/1	0.92	0.13	49,49,49,49	0
56	MG	2a	3087	1/1	0.92	0.18	57,57,57,57	0
56	MG	2A	3064	1/1	0.92	0.27	56,56,56,56	0
56	MG	2A	3369	1/1	0.92	0.15	56,56,56,56	0
56	MG	1A	3477	1/1	0.92	0.20	40,40,40,40	0
56	MG	2A	3371	1/1	0.92	0.12	52,52,52,52	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3070	1/1	0.92	0.17	53,53,53,53	0
56	MG	1A	3920	1/1	0.92	0.15	65,65,65,65	0
56	MG	1a	1632	1/1	0.92	0.11	32,32,32,32	0
56	MG	2a	3109	1/1	0.92	0.23	42,42,42,42	0
56	MG	1A	3186	1/1	0.92	0.16	24,24,24,24	0
56	MG	1A	3425	1/1	0.92	0.11	50,50,50,50	0
56	MG	1A	3740	1/1	0.92	0.15	68,68,68,68	0
56	MG	2A	3393	1/1	0.92	0.12	56,56,56,56	0
56	MG	2A	3395	1/1	0.92	0.14	47,47,47,47	0
56	MG	1a	1799	1/1	0.92	0.14	57,57,57,57	0
56	MG	2a	3120	1/1	0.92	0.18	62,62,62,62	0
56	MG	2A	3084	1/1	0.92	0.08	26,26,26,26	0
56	MG	1A	3741	1/1	0.92	0.09	63,63,63,63	0
56	MG	2a	3123	1/1	0.92	0.11	39,39,39,39	0
56	MG	1A	3341	1/1	0.92	0.07	53,53,53,53	0
56	MG	1A	3484	1/1	0.92	0.08	45,45,45,45	0
56	MG	1A	3541	1/1	0.92	0.09	56,56,56,56	0
56	MG	1A	3942	1/1	0.92	0.10	91,91,91,91	0
56	MG	1a	1809	1/1	0.92	0.10	54,54,54,54	0
56	MG	2a	3134	1/1	0.92	0.27	55,55,55,55	0
56	MG	1a	1647	1/1	0.92	0.17	53,53,53,53	0
56	MG	2A	3419	1/1	0.92	0.26	61,61,61,61	0
56	MG	2a	3137	1/1	0.92	0.14	62,62,62,62	0
56	MG	2A	3421	1/1	0.92	0.23	41,41,41,41	0
56	MG	1a	1649	1/1	0.92	0.10	59,59,59,59	0
56	MG	2A	3431	1/1	0.92	0.12	50,50,50,50	0
56	MG	2a	3144	1/1	0.92	0.12	66,66,66,66	0
56	MG	1a	1812	1/1	0.92	0.09	53,53,53,53	0
56	MG	1A	3429	1/1	0.92	0.10	52,52,52,52	0
56	MG	1A	3547	1/1	0.92	0.14	61,61,61,61	0
56	MG	2A	3436	1/1	0.92	0.10	44,44,44,44	0
56	MG	2A	3437	1/1	0.92	0.10	69,69,69,69	0
56	MG	1D	310	1/1	0.92	0.06	42,42,42,42	0
56	MG	1A	3430	1/1	0.92	0.24	41,41,41,41	0
56	MG	1a	1660	1/1	0.92	0.10	23,23,23,23	0
56	MG	2A	3448	1/1	0.92	0.09	62,62,62,62	0
56	MG	2A	3116	1/1	0.92	0.10	54,54,54,54	0
56	MG	2A	3117	1/1	0.92	0.20	46,46,46,46	0
56	MG	2A	3118	1/1	0.92	0.20	52,52,52,52	0
56	MG	2A	3461	1/1	0.92	0.15	53,53,53,53	0
56	MG	2A	3462	1/1	0.92	0.10	46,46,46,46	0
56	MG	1A	3554	1/1	0.92	0.16	65,65,65,65	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1822	1/1	0.92	0.18	56,56,56,56	0
56	MG	1A	3964	1/1	0.92	0.10	50,50,50,50	0
56	MG	1a	1666	1/1	0.92	0.09	72,72,72,72	0
56	MG	1A	3555	1/1	0.92	0.12	61,61,61,61	0
56	MG	2A	3133	1/1	0.92	0.08	35,35,35,35	0
56	MG	1A	3765	1/1	0.92	0.10	45,45,45,45	0
56	MG	1A	3300	1/1	0.92	0.24	41,41,41,41	0
56	MG	1A	3125	1/1	0.92	0.13	52,52,52,52	0
56	MG	1a	1673	1/1	0.92	0.10	46,46,46,46	0
56	MG	2a	3176	1/1	0.92	0.22	50,50,50,50	0
56	MG	1a	1674	1/1	0.92	0.15	62,62,62,62	0
56	MG	2a	3178	1/1	0.92	0.14	55,55,55,55	0
56	MG	2a	3182	1/1	0.92	0.11	78,78,78,78	0
56	MG	1A	3984	1/1	0.92	0.22	32,32,32,32	0
56	MG	1a	1679	1/1	0.92	0.11	45,45,45,45	0
56	MG	1a	1851	1/1	0.92	0.09	56,56,56,56	0
56	MG	2A	3489	1/1	0.92	0.11	52,52,52,52	0
56	MG	2A	3154	1/1	0.92	0.19	42,42,42,42	0
56	MG	1A	3252	1/1	0.92	0.10	56,56,56,56	0
56	MG	1a	1853	1/1	0.92	0.10	70,70,70,70	0
56	MG	1a	1857	1/1	0.92	0.12	71,71,71,71	0
56	MG	1G	3001	1/1	0.92	0.09	50,50,50,50	0
56	MG	2A	3164	1/1	0.92	0.29	51,51,51,51	0
56	MG	1A	3563	1/1	0.92	0.12	54,54,54,54	0
56	MG	1a	1687	1/1	0.92	0.10	42,42,42,42	0
56	MG	1A	3995	1/1	0.92	0.12	54,54,54,54	0
56	MG	2A	3513	1/1	0.92	0.13	26,26,26,26	0
56	MG	1a	1866	1/1	0.92	0.17	73,73,73,73	0
56	MG	2a	3207	1/1	0.92	0.11	59,59,59,59	0
56	MG	1A	3394	1/1	0.92	0.11	42,42,42,42	0
56	MG	1A	3806	1/1	0.92	0.12	38,38,38,38	0
56	MG	1N	3003	1/1	0.92	0.10	38,38,38,38	0
56	MG	1A	3813	1/1	0.92	0.10	48,48,48,48	0
56	MG	2A	3522	1/1	0.92	0.18	43,43,43,43	0
56	MG	1O	3003	1/1	0.92	0.14	60,60,60,60	0
56	MG	1P	201	1/1	0.92	0.14	49,49,49,49	0
56	MG	2A	3528	1/1	0.92	0.14	46,46,46,46	0
56	MG	1a	1702	1/1	0.92	0.13	60,60,60,60	0
56	MG	2a	3222	1/1	0.92	0.13	62,62,62,62	0
56	MG	2A	3190	1/1	0.92	0.15	48,48,48,48	0
56	MG	2A	3539	1/1	0.92	0.18	57,57,57,57	0
56	MG	1A	3826	1/1	0.92	0.09	17,17,17,17	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3575	1/1	0.92	0.18	57,57,57,57	0
56	MG	1A	4007	1/1	0.92	0.12	35,35,35,35	0
56	MG	1a	1882	1/1	0.92	0.27	60,60,60,60	0
56	MG	1A	3578	1/1	0.92	0.10	48,48,48,48	0
56	MG	1A	3837	1/1	0.92	0.09	26,26,26,26	0
56	MG	2a	3233	1/1	0.92	0.13	58,58,58,58	0
56	MG	1a	1888	1/1	0.92	0.18	60,60,60,60	0
56	MG	1a	1889	1/1	0.92	0.06	42,42,42,42	0
56	MG	1A	4017	1/1	0.92	0.12	61,61,61,61	0
56	MG	1A	3188	1/1	0.92	0.12	53,53,53,53	0
56	MG	1A	3585	1/1	0.92	0.10	14,14,14,14	0
56	MG	1a	1896	1/1	0.92	0.21	46,46,46,46	0
56	MG	2F	302	1/1	0.92	0.17	48,48,48,48	0
56	MG	1A	3846	1/1	0.92	0.11	66,66,66,66	0
56	MG	2N	8001	1/1	0.92	0.09	63,63,63,63	0
56	MG	1A	3194	1/1	0.92	0.07	39,39,39,39	0
56	MG	1a	1900	1/1	0.92	0.16	47,47,47,47	0
56	MG	1a	1903	1/1	0.92	0.14	56,56,56,56	0
56	MG	1a	1734	1/1	0.92	0.22	53,53,53,53	0
56	MG	2A	3228	1/1	0.92	0.11	44,44,44,44	0
56	MG	1A	3195	1/1	0.92	0.12	44,44,44,44	0
56	MG	23	101	1/1	0.92	0.16	50,50,50,50	0
56	MG	2A	3233	1/1	0.92	0.07	52,52,52,52	0
56	MG	1A	3005	1/1	0.92	0.13	57,57,57,57	0
56	MG	2A	3236	1/1	0.92	0.07	60,60,60,60	0
56	MG	1A	3504	1/1	0.92	0.13	62,62,62,62	0
56	MG	2A	3240	1/1	0.92	0.11	34,34,34,34	0
56	MG	1A	3358	1/1	0.92	0.07	61,61,61,61	0
56	MG	1A	3610	1/1	0.92	0.08	48,48,48,48	0
56	MG	1A	3362	1/1	0.92	0.11	49,49,49,49	0
56	MG	2a	3015	1/1	0.92	0.14	62,62,62,62	0
56	MG	2z	8001	1/1	0.92	0.17	69,69,69,69	0
56	MG	2A	3235	1/1	0.93	0.10	57,57,57,57	0
56	MG	1A	3992	1/1	0.93	0.09	36,36,36,36	0
56	MG	28	103	1/1	0.93	0.14	56,56,56,56	0
56	MG	1a	1717	1/1	0.93	0.07	47,47,47,47	0
56	MG	1P	202	1/1	0.93	0.17	59,59,59,59	0
56	MG	1A	3360	1/1	0.93	0.26	57,57,57,57	0
56	MG	2A	3243	1/1	0.93	0.08	69,69,69,69	0
56	MG	2a	3008	1/1	0.93	0.09	53,53,53,53	0
56	MG	1a	1720	1/1	0.93	0.16	45,45,45,45	0
56	MG	2a	3010	1/1	0.93	0.17	44,44,44,44	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2a	3011	1/1	0.93	0.21	58,58,58,58	0
56	MG	1A	3996	1/1	0.93	0.10	37,37,37,37	0
56	MG	1a	1727	1/1	0.93	0.09	56,56,56,56	0
56	MG	2A	3252	1/1	0.93	0.09	43,43,43,43	0
56	MG	1Q	204	1/1	0.93	0.10	45,45,45,45	0
56	MG	2a	3018	1/1	0.93	0.07	62,62,62,62	0
56	MG	1A	3307	1/1	0.93	0.10	63,63,63,63	0
56	MG	1Q	209	1/1	0.93	0.13	59,59,59,59	0
56	MG	1A	3228	1/1	0.93	0.12	48,48,48,48	0
56	MG	1A	3026	1/1	0.93	0.10	47,47,47,47	0
56	MG	1A	3420	1/1	0.93	0.09	54,54,54,54	0
56	MG	2A	3262	1/1	0.93	0.06	53,53,53,53	0
56	MG	1A	3772	1/1	0.93	0.17	43,43,43,43	0
56	MG	1A	3777	1/1	0.93	0.10	30,30,30,30	0
56	MG	1A	3562	1/1	0.93	0.07	54,54,54,54	0
56	MG	1a	1742	1/1	0.93	0.21	68,68,68,68	0
56	MG	2a	3033	1/1	0.93	0.29	64,64,64,64	0
56	MG	1A	4015	1/1	0.93	0.15	46,46,46,46	0
56	MG	1A	3157	1/1	0.93	0.16	43,43,43,43	0
56	MG	1V	202	1/1	0.93	0.07	28,28,28,28	0
56	MG	1A	3236	1/1	0.93	0.14	58,58,58,58	0
56	MG	2A	3284	1/1	0.93	0.09	34,34,34,34	0
56	MG	1A	4020	1/1	0.93	0.06	26,26,26,26	0
56	MG	1A	4022	1/1	0.93	0.08	43,43,43,43	0
56	MG	1x	109	1/1	0.93	0.16	66,66,66,66	0
56	MG	1A	3800	1/1	0.93	0.06	32,32,32,32	0
56	MG	1A	3804	1/1	0.93	0.06	66,66,66,66	0
56	MG	1a	1753	1/1	0.93	0.10	48,48,48,48	0
56	MG	2A	3310	1/1	0.93	0.20	55,55,55,55	0
56	MG	2A	3311	1/1	0.93	0.12	41,41,41,41	0
56	MG	1A	3490	1/1	0.93	0.08	57,57,57,57	0
56	MG	1A	3576	1/1	0.93	0.10	65,65,65,65	0
56	MG	2A	3317	1/1	0.93	0.17	36,36,36,36	0
56	MG	1A	3825	1/1	0.93	0.21	45,45,45,45	0
56	MG	1A	4030	1/1	0.93	0.12	43,43,43,43	0
56	MG	1A	3008	1/1	0.93	0.18	51,51,51,51	0
56	MG	1A	3492	1/1	0.93	0.23	47,47,47,47	0
56	MG	1A	3832	1/1	0.93	0.08	34,34,34,34	0
56	MG	2A	3013	1/1	0.93	0.05	60,60,60,60	0
56	MG	1A	3583	1/1	0.93	0.15	49,49,49,49	0
56	MG	15	102	1/1	0.93	0.15	52,52,52,52	0
56	MG	2A	3019	1/1	0.93	0.07	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3020	1/1	0.93	0.15	39,39,39,39	0
56	MG	1A	3371	1/1	0.93	0.06	46,46,46,46	0
56	MG	1a	1766	1/1	0.93	0.10	50,50,50,50	0
56	MG	2a	3071	1/1	0.93	0.17	72,72,72,72	0
56	MG	1A	3165	1/1	0.93	0.18	23,23,23,23	0
56	MG	1a	1768	1/1	0.93	0.11	56,56,56,56	0
56	MG	1A	3323	1/1	0.93	0.08	41,41,41,41	0
56	MG	2A	3360	1/1	0.93	0.17	43,43,43,43	0
56	MG	2a	3081	1/1	0.93	0.13	65,65,65,65	0
56	MG	2A	3363	1/1	0.93	0.08	49,49,49,49	0
56	MG	2A	3032	1/1	0.93	0.18	62,62,62,62	0
56	MG	1A	3324	1/1	0.93	0.14	57,57,57,57	0
56	MG	1A	3499	1/1	0.93	0.08	64,64,64,64	0
56	MG	1A	3594	1/1	0.93	0.12	40,40,40,40	0
56	MG	1A	3595	1/1	0.93	0.11	66,66,66,66	0
56	MG	1A	4049	1/1	0.93	0.12	49,49,49,49	0
56	MG	2a	3104	1/1	0.93	0.19	54,54,54,54	0
56	MG	1a	1607	1/1	0.93	0.12	60,60,60,60	0
56	MG	1A	3086	1/1	0.93	0.14	52,52,52,52	0
56	MG	1A	3605	1/1	0.93	0.10	31,31,31,31	0
56	MG	2a	3112	1/1	0.93	0.19	53,53,53,53	0
56	MG	1A	3501	1/1	0.93	0.11	51,51,51,51	0
56	MG	2A	3390	1/1	0.93	0.11	35,35,35,35	0
56	MG	2A	3066	1/1	0.93	0.16	48,48,48,48	0
56	MG	1A	3029	1/1	0.93	0.05	71,71,71,71	0
56	MG	2A	3071	1/1	0.93	0.09	46,46,46,46	0
56	MG	1A	3861	1/1	0.93	0.05	42,42,42,42	0
56	MG	1A	3331	1/1	0.93	0.15	62,62,62,62	0
56	MG	1a	1788	1/1	0.93	0.10	51,51,51,51	0
56	MG	1A	3618	1/1	0.93	0.15	68,68,68,68	0
56	MG	2A	3079	1/1	0.93	0.09	50,50,50,50	0
56	MG	2A	3405	1/1	0.93	0.11	46,46,46,46	0
56	MG	2A	3406	1/1	0.93	0.14	55,55,55,55	0
56	MG	2a	3129	1/1	0.93	0.13	60,60,60,60	0
56	MG	2A	3080	1/1	0.93	0.15	59,59,59,59	0
56	MG	1A	3505	1/1	0.93	0.11	49,49,49,49	0
56	MG	1a	1621	1/1	0.93	0.07	57,57,57,57	0
56	MG	1A	3621	1/1	0.93	0.11	40,40,40,40	0
56	MG	1A	4072	1/1	0.93	0.12	48,48,48,48	0
56	MG	1a	1624	1/1	0.93	0.07	80,80,80,80	0
56	MG	1A	3178	1/1	0.93	0.10	37,37,37,37	0
56	MG	2A	3423	1/1	0.93	0.07	53,53,53,53	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	4075	1/1	0.93	0.16	61,61,61,61	0
56	MG	2a	3143	1/1	0.93	0.10	42,42,42,42	0
56	MG	1A	3624	1/1	0.93	0.10	40,40,40,40	0
56	MG	1A	4077	1/1	0.93	0.06	47,47,47,47	0
56	MG	1A	3437	1/1	0.93	0.07	46,46,46,46	0
56	MG	2A	3434	1/1	0.93	0.10	42,42,42,42	0
56	MG	1a	1630	1/1	0.93	0.07	42,42,42,42	0
56	MG	1A	3386	1/1	0.93	0.10	68,68,68,68	0
56	MG	1A	4086	1/1	0.93	0.12	63,63,63,63	0
56	MG	2A	3438	1/1	0.93	0.14	53,53,53,53	0
56	MG	1A	3633	1/1	0.93	0.10	50,50,50,50	0
56	MG	2A	3440	1/1	0.93	0.10	50,50,50,50	0
56	MG	1A	4090	1/1	0.93	0.17	26,26,26,26	0
56	MG	1A	3635	1/1	0.93	0.14	50,50,50,50	0
56	MG	2A	3112	1/1	0.93	0.15	59,59,59,59	0
56	MG	1A	3638	1/1	0.93	0.10	38,38,38,38	0
56	MG	2A	3455	1/1	0.93	0.15	62,62,62,62	0
56	MG	1A	3117	1/1	0.93	0.13	62,62,62,62	0
56	MG	1A	3389	1/1	0.93	0.23	56,56,56,56	0
56	MG	1a	1816	1/1	0.93	0.12	76,76,76,76	0
56	MG	1A	3057	1/1	0.93	0.09	47,47,47,47	0
56	MG	2a	3166	1/1	0.93	0.13	60,60,60,60	0
56	MG	1A	3022	1/1	0.93	0.14	49,49,49,49	0
56	MG	2A	3465	1/1	0.93	0.12	45,45,45,45	0
56	MG	1A	3670	1/1	0.93	0.10	37,37,37,37	0
56	MG	1A	3393	1/1	0.93	0.14	42,42,42,42	0
56	MG	1a	1651	1/1	0.93	0.10	62,62,62,62	0
56	MG	1A	3691	1/1	0.93	0.08	54,54,54,54	0
56	MG	2A	3474	1/1	0.93	0.10	51,51,51,51	0
56	MG	2A	3475	1/1	0.93	0.08	39,39,39,39	0
56	MG	1A	3338	1/1	0.93	0.09	52,52,52,52	0
56	MG	1A	3017	1/1	0.93	0.10	53,53,53,53	0
56	MG	1a	1829	1/1	0.93	0.07	45,45,45,45	0
56	MG	2A	3138	1/1	0.93	0.11	47,47,47,47	0
56	MG	1a	1830	1/1	0.93	0.18	68,68,68,68	0
56	MG	2A	3142	1/1	0.93	0.22	56,56,56,56	0
56	MG	2A	3484	1/1	0.93	0.11	47,47,47,47	0
56	MG	1A	3452	1/1	0.93	0.13	50,50,50,50	0
56	MG	1B	226	1/1	0.93	0.15	71,71,71,71	0
56	MG	2a	3196	1/1	0.93	0.12	70,70,70,70	0
56	MG	1A	3519	1/1	0.93	0.09	48,48,48,48	0
56	MG	1A	3707	1/1	0.93	0.11	32,32,32,32	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1841	1/1	0.93	0.17	52,52,52,52	0
56	MG	2A	3491	1/1	0.93	0.13	46,46,46,46	0
56	MG	1A	3710	1/1	0.93	0.08	48,48,48,48	0
56	MG	1a	1845	1/1	0.93	0.11	62,62,62,62	0
56	MG	1A	3193	1/1	0.93	0.10	53,53,53,53	0
56	MG	1A	3522	1/1	0.93	0.09	50,50,50,50	0
56	MG	1a	1670	1/1	0.93	0.08	50,50,50,50	0
56	MG	2A	3166	1/1	0.93	0.09	32,32,32,32	0
56	MG	2A	3508	1/1	0.93	0.09	42,42,42,42	0
56	MG	1A	3015	1/1	0.93	0.17	51,51,51,51	0
56	MG	1a	1854	1/1	0.93	0.11	60,60,60,60	0
56	MG	1D	312	1/1	0.93	0.11	45,45,45,45	0
56	MG	1A	3460	1/1	0.93	0.23	52,52,52,52	0
56	MG	1a	1675	1/1	0.93	0.10	42,42,42,42	0
56	MG	1A	3039	1/1	0.93	0.10	71,71,71,71	0
56	MG	1A	3025	1/1	0.93	0.14	55,55,55,55	0
56	MG	2a	3221	1/1	0.93	0.24	57,57,57,57	0
56	MG	1a	1865	1/1	0.93	0.15	51,51,51,51	0
56	MG	1A	3204	1/1	0.93	0.23	33,33,33,33	0
56	MG	1a	1867	1/1	0.93	0.16	64,64,64,64	0
56	MG	2A	3187	1/1	0.93	0.17	40,40,40,40	0
56	MG	1a	1682	1/1	0.93	0.07	47,47,47,47	0
56	MG	2A	3529	1/1	0.93	0.24	53,53,53,53	0
56	MG	1a	1684	1/1	0.93	0.12	57,57,57,57	0
56	MG	1E	310	1/1	0.93	0.13	49,49,49,49	0
56	MG	1A	3729	1/1	0.93	0.19	60,60,60,60	0
56	MG	1A	3466	1/1	0.93	0.11	55,55,55,55	0
56	MG	2A	3195	1/1	0.93	0.15	47,47,47,47	0
56	MG	1A	3047	1/1	0.93	0.15	54,54,54,54	0
56	MG	1a	1690	1/1	0.93	0.06	57,57,57,57	0
56	MG	1A	3214	1/1	0.93	0.10	34,34,34,34	0
56	MG	2A	3207	1/1	0.93	0.17	49,49,49,49	0
56	MG	2B	201	1/1	0.93	0.14	57,57,57,57	0
56	MG	1A	3744	1/1	0.93	0.09	43,43,43,43	0
56	MG	1H	3001	1/1	0.93	0.14	35,35,35,35	0
56	MG	1A	3542	1/1	0.93	0.24	63,63,63,63	0
56	MG	2a	3251	1/1	0.93	0.15	43,43,43,43	0
56	MG	2B	209	1/1	0.93	0.31	57,57,57,57	0
56	MG	1a	1885	1/1	0.93	0.12	54,54,54,54	0
56	MG	1A	3215	1/1	0.93	0.11	49,49,49,49	0
56	MG	1A	3410	1/1	0.93	0.07	42,42,42,42	0
56	MG	2E	301	1/1	0.93	0.22	36,36,36,36	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2E	303	1/1	0.93	0.05	60,60,60,60	0
56	MG	2E	304	1/1	0.93	0.04	56,56,56,56	0
56	MG	2I	3001	1/1	0.93	0.10	65,65,65,65	0
56	MG	1A	3549	1/1	0.93	0.14	73,73,73,73	0
56	MG	2I	3004	1/1	0.93	0.10	60,60,60,60	0
56	MG	1A	3987	1/1	0.93	0.11	58,58,58,58	0
56	MG	1a	1891	1/1	0.93	0.14	51,51,51,51	0
56	MG	1A	3153	1/1	0.93	0.14	20,20,20,20	0
56	MG	1a	1707	1/1	0.93	0.09	31,31,31,31	0
56	MG	1a	1710	1/1	0.93	0.13	45,45,45,45	0
56	MG	2x	103	1/1	0.93	0.13	65,65,65,65	0
56	MG	1A	3552	1/1	0.93	0.10	51,51,51,51	0
56	MG	2x	105	1/1	0.93	0.13	52,52,52,52	0
56	MG	1a	1897	1/1	0.93	0.20	84,84,84,84	0
56	MG	2Y	502	1/1	0.93	0.14	51,51,51,51	0
56	MG	1a	1715	1/1	0.93	0.17	50,50,50,50	0
57	ZN	24	501	1/1	0.93	0.08	133,133,133,133	0
56	MG	1A	3169	1/1	0.94	0.07	55,55,55,55	0
56	MG	1A	3447	1/1	0.94	0.08	54,54,54,54	0
56	MG	2R	201	1/1	0.94	0.17	49,49,49,49	0
56	MG	1A	3093	1/1	0.94	0.07	48,48,48,48	0
56	MG	2A	3249	1/1	0.94	0.12	56,56,56,56	0
56	MG	1A	3002	1/1	0.94	0.18	38,38,38,38	0
56	MG	1A	3060	1/1	0.94	0.08	46,46,46,46	0
56	MG	1u	8001	1/1	0.94	0.08	53,53,53,53	0
56	MG	28	101	1/1	0.94	0.14	45,45,45,45	0
56	MG	1x	101	1/1	0.94	0.06	64,64,64,64	0
56	MG	1x	102	1/1	0.94	0.16	67,67,67,67	0
56	MG	2a	3003	1/1	0.94	0.18	55,55,55,55	0
56	MG	1A	3634	1/1	0.94	0.10	38,38,38,38	0
56	MG	1x	104	1/1	0.94	0.12	56,56,56,56	0
56	MG	1A	3098	1/1	0.94	0.13	40,40,40,40	0
56	MG	16	101	1/1	0.94	0.16	55,55,55,55	0
56	MG	2A	3267	1/1	0.94	0.18	38,38,38,38	0
56	MG	2A	3268	1/1	0.94	0.13	44,44,44,44	0
56	MG	17	102	1/1	0.94	0.13	60,60,60,60	0
56	MG	2A	3271	1/1	0.94	0.08	50,50,50,50	0
56	MG	2a	3013	1/1	0.94	0.16	50,50,50,50	0
56	MG	18	3001	1/1	0.94	0.17	50,50,50,50	0
56	MG	1A	3636	1/1	0.94	0.21	49,49,49,49	0
56	MG	1A	3256	1/1	0.94	0.42	44,44,44,44	0
56	MG	2A	3277	1/1	0.94	0.10	44,44,44,44	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	19	104	1/1	0.94	0.08	42,42,42,42	0
56	MG	1A	3862	1/1	0.94	0.15	45,45,45,45	0
56	MG	1A	3639	1/1	0.94	0.13	47,47,47,47	0
56	MG	1A	3641	1/1	0.94	0.09	52,52,52,52	0
56	MG	1A	3867	1/1	0.94	0.13	60,60,60,60	0
56	MG	2A	3289	1/1	0.94	0.10	37,37,37,37	0
56	MG	1A	3392	1/1	0.94	0.08	33,33,33,33	0
56	MG	2a	3028	1/1	0.94	0.07	38,38,38,38	0
56	MG	1A	3869	1/1	0.94	0.06	40,40,40,40	0
56	MG	1A	3524	1/1	0.94	0.07	68,68,68,68	0
56	MG	1A	3454	1/1	0.94	0.07	64,64,64,64	0
56	MG	2A	3305	1/1	0.94	0.10	50,50,50,50	0
56	MG	2A	3306	1/1	0.94	0.06	32,32,32,32	0
56	MG	2a	3037	1/1	0.94	0.15	50,50,50,50	0
56	MG	2A	3014	1/1	0.94	0.13	52,52,52,52	0
56	MG	1A	3649	1/1	0.94	0.11	26,26,26,26	0
56	MG	1a	1612	1/1	0.94	0.09	57,57,57,57	0
56	MG	2a	3041	1/1	0.94	0.08	50,50,50,50	0
56	MG	2A	3312	1/1	0.94	0.09	38,38,38,38	0
56	MG	1a	1614	1/1	0.94	0.06	66,66,66,66	0
56	MG	1A	3659	1/1	0.94	0.08	17,17,17,17	0
56	MG	1A	3526	1/1	0.94	0.12	47,47,47,47	0
56	MG	2A	3025	1/1	0.94	0.26	55,55,55,55	0
56	MG	1A	3100	1/1	0.94	0.10	51,51,51,51	0
56	MG	2A	3322	1/1	0.94	0.08	49,49,49,49	0
56	MG	1A	3680	1/1	0.94	0.16	44,44,44,44	0
56	MG	1A	3457	1/1	0.94	0.09	61,61,61,61	0
56	MG	2A	3331	1/1	0.94	0.17	45,45,45,45	0
56	MG	1A	3885	1/1	0.94	0.12	54,54,54,54	0
56	MG	1A	3887	1/1	0.94	0.15	45,45,45,45	0
56	MG	2A	3340	1/1	0.94	0.14	52,52,52,52	0
56	MG	2A	3034	1/1	0.94	0.18	44,44,44,44	0
56	MG	1A	3690	1/1	0.94	0.07	21,21,21,21	0
56	MG	1A	3261	1/1	0.94	0.38	68,68,68,68	0
56	MG	1A	3697	1/1	0.94	0.12	41,41,41,41	0
56	MG	2a	3062	1/1	0.94	0.07	81,81,81,81	0
56	MG	2A	3051	1/1	0.94	0.12	35,35,35,35	0
56	MG	2A	3352	1/1	0.94	0.09	38,38,38,38	0
56	MG	1A	3895	1/1	0.94	0.05	40,40,40,40	0
56	MG	1A	3124	1/1	0.94	0.20	36,36,36,36	0
56	MG	1A	3534	1/1	0.94	0.11	55,55,55,55	0
56	MG	1a	1794	1/1	0.94	0.06	56,56,56,56	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3061	1/1	0.94	0.13	29,29,29,29	0
56	MG	2A	3361	1/1	0.94	0.10	34,34,34,34	0
56	MG	2A	3063	1/1	0.94	0.21	48,48,48,48	0
56	MG	1A	3267	1/1	0.94	0.12	22,22,22,22	0
56	MG	2a	3074	1/1	0.94	0.20	47,47,47,47	0
56	MG	2A	3365	1/1	0.94	0.06	41,41,41,41	0
56	MG	1B	207	1/1	0.94	0.15	52,52,52,52	0
56	MG	2a	3077	1/1	0.94	0.10	51,51,51,51	0
56	MG	2a	3080	1/1	0.94	0.31	55,55,55,55	0
56	MG	1A	3907	1/1	0.94	0.10	50,50,50,50	0
56	MG	2a	3083	1/1	0.94	0.10	49,49,49,49	0
56	MG	2A	3067	1/1	0.94	0.12	49,49,49,49	0
56	MG	2A	3068	1/1	0.94	0.10	52,52,52,52	0
56	MG	2a	3090	1/1	0.94	0.19	47,47,47,47	0
56	MG	1A	3537	1/1	0.94	0.10	53,53,53,53	0
56	MG	1a	1636	1/1	0.94	0.25	53,53,53,53	0
56	MG	2a	3094	1/1	0.94	0.24	55,55,55,55	0
56	MG	2a	3095	1/1	0.94	0.08	41,41,41,41	0
56	MG	2A	3072	1/1	0.94	0.08	29,29,29,29	0
56	MG	2a	3097	1/1	0.94	0.14	55,55,55,55	0
56	MG	2A	3378	1/1	0.94	0.09	41,41,41,41	0
56	MG	1A	3270	1/1	0.94	0.09	41,41,41,41	0
56	MG	1B	212	1/1	0.94	0.25	59,59,59,59	0
56	MG	1B	213	1/1	0.94	0.07	50,50,50,50	0
56	MG	2A	3076	1/1	0.94	0.17	56,56,56,56	0
56	MG	1A	3709	1/1	0.94	0.15	38,38,38,38	0
56	MG	1A	3049	1/1	0.94	0.07	37,37,37,37	0
56	MG	2a	3113	1/1	0.94	0.30	50,50,50,50	0
56	MG	1a	1807	1/1	0.94	0.09	55,55,55,55	0
56	MG	1A	3127	1/1	0.94	0.14	40,40,40,40	0
56	MG	2a	3116	1/1	0.94	0.12	45,45,45,45	0
56	MG	1A	3543	1/1	0.94	0.12	46,46,46,46	0
56	MG	2A	3400	1/1	0.94	0.11	48,48,48,48	0
56	MG	1A	3471	1/1	0.94	0.12	55,55,55,55	0
56	MG	2A	3403	1/1	0.94	0.09	57,57,57,57	0
56	MG	2A	3085	1/1	0.94	0.12	38,38,38,38	0
56	MG	2A	3086	1/1	0.94	0.13	47,47,47,47	0
56	MG	1A	3718	1/1	0.94	0.11	36,36,36,36	0
56	MG	1A	3018	1/1	0.94	0.11	53,53,53,53	0
56	MG	1A	3548	1/1	0.94	0.09	72,72,72,72	0
56	MG	2a	3127	1/1	0.94	0.12	50,50,50,50	0
56	MG	1A	3139	1/1	0.94	0.12	52,52,52,52	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1654	1/1	0.94	0.07	45,45,45,45	0
56	MG	1D	301	1/1	0.94	0.15	32,32,32,32	0
56	MG	2a	3132	1/1	0.94	0.10	57,57,57,57	0
56	MG	2A	3417	1/1	0.94	0.08	44,44,44,44	0
56	MG	2A	3418	1/1	0.94	0.13	52,52,52,52	0
56	MG	1D	305	1/1	0.94	0.15	13,13,13,13	0
56	MG	2A	3420	1/1	0.94	0.35	72,72,72,72	0
56	MG	1a	1657	1/1	0.94	0.10	62,62,62,62	0
56	MG	1A	3145	1/1	0.94	0.17	38,38,38,38	0
56	MG	2a	3140	1/1	0.94	0.18	49,49,49,49	0
56	MG	1A	3407	1/1	0.94	0.17	40,40,40,40	0
56	MG	1A	3947	1/1	0.94	0.17	42,42,42,42	0
56	MG	1a	1663	1/1	0.94	0.11	52,52,52,52	0
56	MG	1a	1664	1/1	0.94	0.15	62,62,62,62	0
56	MG	1A	3199	1/1	0.94	0.14	44,44,44,44	0
56	MG	1A	3089	1/1	0.94	0.05	52,52,52,52	0
56	MG	1A	3147	1/1	0.94	0.12	28,28,28,28	0
56	MG	1a	1833	1/1	0.94	0.10	62,62,62,62	0
56	MG	1A	3483	1/1	0.94	0.12	41,41,41,41	0
56	MG	2A	3122	1/1	0.94	0.20	51,51,51,51	0
56	MG	1A	3414	1/1	0.94	0.15	50,50,50,50	0
56	MG	2A	3442	1/1	0.94	0.06	66,66,66,66	0
56	MG	2A	3125	1/1	0.94	0.15	52,52,52,52	0
56	MG	2A	3127	1/1	0.94	0.14	43,43,43,43	0
56	MG	1A	3559	1/1	0.94	0.07	55,55,55,55	0
56	MG	2A	3449	1/1	0.94	0.08	57,57,57,57	0
56	MG	1a	1672	1/1	0.94	0.17	47,47,47,47	0
56	MG	2A	3454	1/1	0.94	0.09	57,57,57,57	0
56	MG	1A	3290	1/1	0.94	0.12	30,30,30,30	0
56	MG	1F	303	1/1	0.94	0.35	32,32,32,32	0
56	MG	1A	3972	1/1	0.94	0.08	41,41,41,41	0
56	MG	1a	1850	1/1	0.94	0.13	44,44,44,44	0
56	MG	1A	3297	1/1	0.94	0.17	49,49,49,49	0
56	MG	1A	3975	1/1	0.94	0.12	44,44,44,44	0
56	MG	1A	3977	1/1	0.94	0.06	31,31,31,31	0
56	MG	1A	3208	1/1	0.94	0.05	55,55,55,55	0
56	MG	1a	1855	1/1	0.94	0.18	48,48,48,48	0
56	MG	2a	3171	1/1	0.94	0.12	46,46,46,46	0
56	MG	2A	3143	1/1	0.94	0.07	41,41,41,41	0
56	MG	1A	3105	1/1	0.94	0.16	47,47,47,47	0
56	MG	2A	3473	1/1	0.94	0.10	41,41,41,41	0
56	MG	1A	3757	1/1	0.94	0.08	49,49,49,49	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3006	1/1	0.94	0.11	54,54,54,54	0
56	MG	1A	3990	1/1	0.94	0.16	17,17,17,17	0
56	MG	2A	3477	1/1	0.94	0.07	47,47,47,47	0
56	MG	1A	3302	1/1	0.94	0.14	59,59,59,59	0
56	MG	2A	3158	1/1	0.94	0.06	39,39,39,39	0
56	MG	2a	3188	1/1	0.94	0.16	53,53,53,53	0
56	MG	1N	3005	1/1	0.94	0.11	45,45,45,45	0
56	MG	1N	3006	1/1	0.94	0.08	55,55,55,55	0
56	MG	1A	3306	1/1	0.94	0.31	69,69,69,69	0
56	MG	1a	1869	1/1	0.94	0.18	68,68,68,68	0
56	MG	2a	3195	1/1	0.94	0.09	48,48,48,48	0
56	MG	1A	3111	1/1	0.94	0.20	35,35,35,35	0
56	MG	1A	3495	1/1	0.94	0.11	41,41,41,41	0
56	MG	1A	3424	1/1	0.94	0.10	37,37,37,37	0
56	MG	2A	3488	1/1	0.94	0.09	69,69,69,69	0
56	MG	1a	1698	1/1	0.94	0.18	63,63,63,63	0
56	MG	1a	1699	1/1	0.94	0.18	58,58,58,58	0
56	MG	1A	3773	1/1	0.94	0.16	48,48,48,48	0
56	MG	1a	1701	1/1	0.94	0.10	63,63,63,63	0
56	MG	2a	3205	1/1	0.94	0.22	57,57,57,57	0
56	MG	1Q	201	1/1	0.94	0.08	33,33,33,33	0
56	MG	1a	1703	1/1	0.94	0.16	52,52,52,52	0
56	MG	1A	3311	1/1	0.94	0.12	44,44,44,44	0
56	MG	2A	3499	1/1	0.94	0.08	49,49,49,49	0
56	MG	2A	3501	1/1	0.94	0.08	37,37,37,37	0
56	MG	2a	3211	1/1	0.94	0.16	55,55,55,55	0
56	MG	2A	3182	1/1	0.94	0.05	36,36,36,36	0
56	MG	1A	3786	1/1	0.94	0.13	23,23,23,23	0
56	MG	1A	3589	1/1	0.94	0.12	36,36,36,36	0
56	MG	2A	3509	1/1	0.94	0.07	46,46,46,46	0
56	MG	1A	3426	1/1	0.94	0.09	54,54,54,54	0
56	MG	2A	3188	1/1	0.94	0.12	41,41,41,41	0
56	MG	2a	3219	1/1	0.94	0.13	62,62,62,62	0
56	MG	1A	3161	1/1	0.94	0.22	47,47,47,47	0
56	MG	1A	3315	1/1	0.94	0.10	65,65,65,65	0
56	MG	2A	3191	1/1	0.94	0.16	37,37,37,37	0
56	MG	1A	3802	1/1	0.94	0.14	65,65,65,65	0
56	MG	1R	205	1/1	0.94	0.11	28,28,28,28	0
56	MG	1A	3033	1/1	0.94	0.10	50,50,50,50	0
56	MG	2A	3521	1/1	0.94	0.08	41,41,41,41	0
56	MG	1A	4019	1/1	0.94	0.12	39,39,39,39	0
56	MG	1T	204	1/1	0.94	0.12	50,50,50,50	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3374	1/1	0.94	0.14	26,26,26,26	0
56	MG	1A	3808	1/1	0.94	0.08	73,73,73,73	0
56	MG	2A	3200	1/1	0.94	0.11	50,50,50,50	0
56	MG	2A	3531	1/1	0.94	0.15	52,52,52,52	0
56	MG	1a	1728	1/1	0.94	0.19	46,46,46,46	0
56	MG	2a	3237	1/1	0.94	0.08	59,59,59,59	0
56	MG	2a	3239	1/1	0.94	0.24	55,55,55,55	0
56	MG	1A	3317	1/1	0.94	0.24	63,63,63,63	0
56	MG	1A	3232	1/1	0.94	0.07	44,44,44,44	0
56	MG	2A	3541	1/1	0.94	0.15	38,38,38,38	0
56	MG	2A	3213	1/1	0.94	0.16	42,42,42,42	0
56	MG	1W	3004	1/1	0.94	0.09	40,40,40,40	0
56	MG	2A	3555	1/1	0.94	0.17	49,49,49,49	0
56	MG	1a	1904	1/1	0.94	0.23	76,76,76,76	0
56	MG	1A	3613	1/1	0.94	0.11	41,41,41,41	0
56	MG	1a	1908	1/1	0.94	0.08	53,53,53,53	0
56	MG	1A	3164	1/1	0.94	0.08	40,40,40,40	0
56	MG	1a	1737	1/1	0.94	0.22	48,48,48,48	0
56	MG	1d	502	1/1	0.94	0.12	82,82,82,82	0
56	MG	2B	204	1/1	0.94	0.29	65,65,65,65	0
56	MG	1A	3379	1/1	0.94	0.27	43,43,43,43	0
56	MG	2f	201	1/1	0.94	0.11	48,48,48,48	0
56	MG	2B	206	1/1	0.94	0.13	37,37,37,37	0
56	MG	1A	3380	1/1	0.94	0.09	49,49,49,49	0
56	MG	1e	202	1/1	0.94	0.12	65,65,65,65	0
56	MG	2A	3229	1/1	0.94	0.17	59,59,59,59	0
56	MG	1A	4031	1/1	0.94	0.20	66,66,66,66	0
56	MG	2q	203	1/1	0.94	0.20	53,53,53,53	0
56	MG	2r	3001	1/1	0.94	0.10	55,55,55,55	0
56	MG	1A	3381	1/1	0.94	0.15	54,54,54,54	0
56	MG	10	105	1/1	0.94	0.10	52,52,52,52	0
56	MG	1l	202	1/1	0.94	0.05	58,58,58,58	0
56	MG	1A	3044	1/1	0.94	0.21	60,60,60,60	0
56	MG	2A	3238	1/1	0.94	0.12	30,30,30,30	0
56	MG	2F	304	1/1	0.94	0.06	41,41,41,41	0
56	MG	1n	102	1/1	0.94	0.08	53,53,53,53	0
56	MG	1a	1745	1/1	0.94	0.06	55,55,55,55	0
56	MG	1o	103	1/1	0.94	0.16	47,47,47,47	0
56	MG	1A	3622	1/1	0.94	0.06	36,36,36,36	0
56	MG	1A	3080	1/1	0.95	0.12	53,53,53,53	0
56	MG	1A	3937	1/1	0.95	0.13	52,52,52,52	0
56	MG	1a	1906	1/1	0.95	0.18	45,45,45,45	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2F	301	1/1	0.95	0.10	49,49,49,49	0
56	MG	1A	3938	1/1	0.95	0.07	45,45,45,45	0
56	MG	1F	305	1/1	0.95	0.07	40,40,40,40	0
56	MG	1A	3939	1/1	0.95	0.11	19,19,19,19	0
56	MG	1A	3568	1/1	0.95	0.11	45,45,45,45	0
56	MG	1A	3569	1/1	0.95	0.06	40,40,40,40	0
56	MG	1G	3003	1/1	0.95	0.06	43,43,43,43	0
56	MG	2P	201	1/1	0.95	0.10	33,33,33,33	0
56	MG	1A	3409	1/1	0.95	0.10	42,42,42,42	0
56	MG	2Q	3002	1/1	0.95	0.15	55,55,55,55	0
56	MG	2Q	3004	1/1	0.95	0.15	41,41,41,41	0
56	MG	1A	3489	1/1	0.95	0.08	41,41,41,41	0
56	MG	1h	8001	1/1	0.95	0.07	60,60,60,60	0
56	MG	1a	1714	1/1	0.95	0.14	57,57,57,57	0
56	MG	1A	3257	1/1	0.95	0.15	39,39,39,39	0
56	MG	2A	3245	1/1	0.95	0.14	52,52,52,52	0
56	MG	1A	3952	1/1	0.95	0.08	40,40,40,40	0
56	MG	1N	3001	1/1	0.95	0.13	44,44,44,44	0
56	MG	1A	3752	1/1	0.95	0.06	63,63,63,63	0
56	MG	2A	3251	1/1	0.95	0.10	49,49,49,49	0
56	MG	1A	3046	1/1	0.95	0.14	45,45,45,45	0
56	MG	2a	3002	1/1	0.95	0.15	58,58,58,58	0
56	MG	1o	102	1/1	0.95	0.05	72,72,72,72	0
56	MG	1A	3959	1/1	0.95	0.08	53,53,53,53	0
56	MG	1a	1721	1/1	0.95	0.15	45,45,45,45	0
56	MG	1A	3580	1/1	0.95	0.10	36,36,36,36	0
56	MG	2a	3007	1/1	0.95	0.23	40,40,40,40	0
56	MG	1A	3128	1/1	0.95	0.09	50,50,50,50	0
56	MG	1O	3001	1/1	0.95	0.10	39,39,39,39	0
56	MG	1O	3002	1/1	0.95	0.10	47,47,47,47	0
56	MG	2A	3263	1/1	0.95	0.06	42,42,42,42	0
56	MG	1a	1730	1/1	0.95	0.15	44,44,44,44	0
56	MG	2A	3266	1/1	0.95	0.06	41,41,41,41	0
56	MG	1A	3968	1/1	0.95	0.10	26,26,26,26	0
56	MG	1v	3001	1/1	0.95	0.12	41,41,41,41	0
56	MG	1A	3415	1/1	0.95	0.07	39,39,39,39	0
56	MG	1A	3759	1/1	0.95	0.10	68,68,68,68	0
56	MG	1P	203	1/1	0.95	0.33	28,28,28,28	0
56	MG	1A	3971	1/1	0.95	0.05	27,27,27,27	0
56	MG	1A	3761	1/1	0.95	0.07	67,67,67,67	0
56	MG	1A	3262	1/1	0.95	0.15	28,28,28,28	0
56	MG	2a	3024	1/1	0.95	0.10	78,78,78,78	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1x	107	1/1	0.95	0.23	48,48,48,48	0
56	MG	1A	3129	1/1	0.95	0.07	42,42,42,42	0
56	MG	2A	3280	1/1	0.95	0.09	58,58,58,58	0
56	MG	1A	3976	1/1	0.95	0.14	42,42,42,42	0
56	MG	1A	3766	1/1	0.95	0.12	37,37,37,37	0
56	MG	2A	3001	1/1	0.95	0.07	35,35,35,35	0
56	MG	1A	3979	1/1	0.95	0.11	55,55,55,55	0
56	MG	2A	3290	1/1	0.95	0.09	36,36,36,36	0
56	MG	2a	3034	1/1	0.95	0.09	43,43,43,43	0
56	MG	2A	3293	1/1	0.95	0.09	30,30,30,30	0
56	MG	1A	3982	1/1	0.95	0.09	21,21,21,21	0
56	MG	2A	3296	1/1	0.95	0.17	57,57,57,57	0
56	MG	1A	3768	1/1	0.95	0.07	8,8,8,8	0
56	MG	1A	3265	1/1	0.95	0.14	27,27,27,27	0
56	MG	1A	3136	1/1	0.95	0.10	35,35,35,35	0
56	MG	1A	3013	1/1	0.95	0.23	43,43,43,43	0
56	MG	1A	3085	1/1	0.95	0.12	48,48,48,48	0
56	MG	1A	3774	1/1	0.95	0.09	43,43,43,43	0
56	MG	2A	3309	1/1	0.95	0.10	37,37,37,37	0
56	MG	1A	3061	1/1	0.95	0.07	51,51,51,51	0
56	MG	2a	3048	1/1	0.95	0.14	66,66,66,66	0
56	MG	1U	204	1/1	0.95	0.15	40,40,40,40	0
56	MG	2a	3050	1/1	0.95	0.06	86,86,86,86	0
56	MG	1A	3779	1/1	0.95	0.09	35,35,35,35	0
56	MG	2A	3313	1/1	0.95	0.14	46,46,46,46	0
56	MG	1A	3780	1/1	0.95	0.09	26,26,26,26	0
56	MG	2A	3016	1/1	0.95	0.09	51,51,51,51	0
56	MG	1W	3001	1/1	0.95	0.14	53,53,53,53	0
56	MG	2A	3018	1/1	0.95	0.09	47,47,47,47	0
56	MG	1A	3106	1/1	0.95	0.06	48,48,48,48	0
56	MG	2A	3321	1/1	0.95	0.14	26,26,26,26	0
56	MG	1W	3005	1/1	0.95	0.20	42,42,42,42	0
56	MG	1A	3787	1/1	0.95	0.07	26,26,26,26	0
56	MG	1A	3601	1/1	0.95	0.09	36,36,36,36	0
56	MG	1A	3001	1/1	0.95	0.17	31,31,31,31	0
56	MG	2A	3334	1/1	0.95	0.10	53,53,53,53	0
56	MG	2A	3027	1/1	0.95	0.10	56,56,56,56	0
56	MG	2A	3028	1/1	0.95	0.12	53,53,53,53	0
56	MG	1a	1763	1/1	0.95	0.17	55,55,55,55	0
56	MG	2a	3067	1/1	0.95	0.30	57,57,57,57	0
56	MG	1A	3200	1/1	0.95	0.12	39,39,39,39	0
56	MG	1A	4003	1/1	0.95	0.09	44,44,44,44	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	4005	1/1	0.95	0.10	42,42,42,42	0
56	MG	10	104	1/1	0.95	0.12	47,47,47,47	0
56	MG	1A	4006	1/1	0.95	0.08	35,35,35,35	0
56	MG	2A	3350	1/1	0.95	0.09	62,62,62,62	0
56	MG	2A	3351	1/1	0.95	0.12	49,49,49,49	0
56	MG	1A	3280	1/1	0.95	0.11	43,43,43,43	0
56	MG	1A	3201	1/1	0.95	0.07	40,40,40,40	0
56	MG	2A	3043	1/1	0.95	0.13	42,42,42,42	0
56	MG	2a	3078	1/1	0.95	0.16	61,61,61,61	0
56	MG	2A	3045	1/1	0.95	0.15	48,48,48,48	0
56	MG	2A	3046	1/1	0.95	0.30	60,60,60,60	0
56	MG	2a	3082	1/1	0.95	0.21	54,54,54,54	0
56	MG	2A	3048	1/1	0.95	0.07	45,45,45,45	0
56	MG	1A	4014	1/1	0.95	0.10	57,57,57,57	0
56	MG	2A	3362	1/1	0.95	0.11	42,42,42,42	0
56	MG	1A	3361	1/1	0.95	0.12	45,45,45,45	0
56	MG	2a	3091	1/1	0.95	0.14	46,46,46,46	0
56	MG	2A	3053	1/1	0.95	0.10	43,43,43,43	0
56	MG	1A	3283	1/1	0.95	0.25	43,43,43,43	0
56	MG	1A	3364	1/1	0.95	0.10	64,64,64,64	0
56	MG	2A	3367	1/1	0.95	0.14	54,54,54,54	0
56	MG	2A	3056	1/1	0.95	0.13	52,52,52,52	0
56	MG	1A	4018	1/1	0.95	0.07	45,45,45,45	0
56	MG	2a	3098	1/1	0.95	0.26	55,55,55,55	0
56	MG	1a	1779	1/1	0.95	0.07	66,66,66,66	0
56	MG	2A	3062	1/1	0.95	0.10	46,46,46,46	0
56	MG	1A	3620	1/1	0.95	0.06	55,55,55,55	0
56	MG	2A	3373	1/1	0.95	0.11	48,48,48,48	0
56	MG	1A	3818	1/1	0.95	0.10	59,59,59,59	0
56	MG	2A	3376	1/1	0.95	0.07	19,19,19,19	0
56	MG	1A	3820	1/1	0.95	0.12	71,71,71,71	0
56	MG	1A	3824	1/1	0.95	0.09	49,49,49,49	0
56	MG	2A	3381	1/1	0.95	0.17	41,41,41,41	0
56	MG	1a	1785	1/1	0.95	0.08	60,60,60,60	0
56	MG	2A	3389	1/1	0.95	0.09	39,39,39,39	0
56	MG	1A	3202	1/1	0.95	0.15	50,50,50,50	0
56	MG	2A	3069	1/1	0.95	0.21	42,42,42,42	0
56	MG	1A	3286	1/1	0.95	0.10	40,40,40,40	0
56	MG	1A	3150	1/1	0.95	0.24	44,44,44,44	0
56	MG	1A	3828	1/1	0.95	0.07	49,49,49,49	0
56	MG	1A	3830	1/1	0.95	0.05	34,34,34,34	0
56	MG	1A	3831	1/1	0.95	0.07	18,18,18,18	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	4032	1/1	0.95	0.08	38,38,38,38	0
56	MG	1A	3205	1/1	0.95	0.12	33,33,33,33	0
56	MG	2A	3402	1/1	0.95	0.15	44,44,44,44	0
56	MG	1A	3834	1/1	0.95	0.09	63,63,63,63	0
56	MG	2A	3078	1/1	0.95	0.09	48,48,48,48	0
56	MG	1A	3438	1/1	0.95	0.08	49,49,49,49	0
56	MG	1A	3836	1/1	0.95	0.05	25,25,25,25	0
56	MG	1a	1609	1/1	0.95	0.12	58,58,58,58	0
56	MG	1A	3152	1/1	0.95	0.08	30,30,30,30	0
56	MG	2A	3083	1/1	0.95	0.09	39,39,39,39	0
56	MG	1A	3442	1/1	0.95	0.07	39,39,39,39	0
56	MG	2A	3413	1/1	0.95	0.07	55,55,55,55	0
56	MG	1A	3520	1/1	0.95	0.08	61,61,61,61	0
56	MG	2a	3139	1/1	0.95	0.13	53,53,53,53	0
56	MG	2A	3416	1/1	0.95	0.07	36,36,36,36	0
56	MG	1a	1613	1/1	0.95	0.09	77,77,77,77	0
56	MG	1a	1805	1/1	0.95	0.11	61,61,61,61	0
56	MG	1A	3207	1/1	0.95	0.16	42,42,42,42	0
56	MG	1A	3021	1/1	0.95	0.06	56,56,56,56	0
56	MG	1A	4046	1/1	0.95	0.13	45,45,45,45	0
56	MG	1A	3209	1/1	0.95	0.17	39,39,39,39	0
56	MG	2A	3096	1/1	0.95	0.11	47,47,47,47	0
56	MG	2A	3428	1/1	0.95	0.09	59,59,59,59	0
56	MG	2A	3429	1/1	0.95	0.10	53,53,53,53	0
56	MG	2A	3097	1/1	0.95	0.17	51,51,51,51	0
56	MG	1A	3850	1/1	0.95	0.10	47,47,47,47	0
56	MG	1A	3853	1/1	0.95	0.07	35,35,35,35	0
56	MG	1A	3210	1/1	0.95	0.07	40,40,40,40	0
56	MG	1A	4052	1/1	0.95	0.31	33,33,33,33	0
56	MG	2a	3156	1/1	0.95	0.12	50,50,50,50	0
56	MG	2A	3103	1/1	0.95	0.08	46,46,46,46	0
56	MG	1A	3640	1/1	0.95	0.12	38,38,38,38	0
56	MG	1A	3305	1/1	0.95	0.15	50,50,50,50	0
56	MG	2a	3160	1/1	0.95	0.09	61,61,61,61	0
56	MG	1A	4057	1/1	0.95	0.15	50,50,50,50	0
56	MG	1A	3112	1/1	0.95	0.20	51,51,51,51	0
56	MG	2A	3110	1/1	0.95	0.22	53,53,53,53	0
56	MG	1A	4059	1/1	0.95	0.15	28,28,28,28	0
56	MG	1a	1821	1/1	0.95	0.14	52,52,52,52	0
56	MG	2A	3446	1/1	0.95	0.09	55,55,55,55	0
56	MG	1A	3859	1/1	0.95	0.08	52,52,52,52	0
56	MG	1A	3067	1/1	0.95	0.19	56,56,56,56	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	4062	1/1	0.95	0.10	49,49,49,49	0
56	MG	2A	3120	1/1	0.95	0.13	58,58,58,58	0
56	MG	2A	3121	1/1	0.95	0.14	38,38,38,38	0
56	MG	1A	3530	1/1	0.95	0.06	47,47,47,47	0
56	MG	2a	3174	1/1	0.95	0.11	51,51,51,51	0
56	MG	1a	1826	1/1	0.95	0.09	59,59,59,59	0
56	MG	1A	3648	1/1	0.95	0.07	26,26,26,26	0
56	MG	1a	1828	1/1	0.95	0.08	70,70,70,70	0
56	MG	1a	1634	1/1	0.95	0.24	55,55,55,55	0
56	MG	1A	4065	1/1	0.95	0.07	33,33,33,33	0
56	MG	1a	1638	1/1	0.95	0.07	52,52,52,52	0
56	MG	1a	1832	1/1	0.95	0.10	52,52,52,52	0
56	MG	1A	3863	1/1	0.95	0.14	38,38,38,38	0
56	MG	1A	4070	1/1	0.95	0.14	47,47,47,47	0
56	MG	1A	3864	1/1	0.95	0.06	38,38,38,38	0
56	MG	1A	3308	1/1	0.95	0.18	49,49,49,49	0
56	MG	1A	3309	1/1	0.95	0.16	59,59,59,59	0
56	MG	1A	3310	1/1	0.95	0.10	52,52,52,52	0
56	MG	1a	1844	1/1	0.95	0.11	56,56,56,56	0
56	MG	1A	3382	1/1	0.95	0.07	53,53,53,53	0
56	MG	2A	3144	1/1	0.95	0.17	41,41,41,41	0
56	MG	1A	3216	1/1	0.95	0.10	39,39,39,39	0
56	MG	1A	3687	1/1	0.95	0.11	17,17,17,17	0
56	MG	1A	4085	1/1	0.95	0.30	34,34,34,34	0
56	MG	1A	3871	1/1	0.95	0.07	31,31,31,31	0
56	MG	1a	1652	1/1	0.95	0.25	60,60,60,60	0
56	MG	1A	3539	1/1	0.95	0.07	45,45,45,45	0
56	MG	1A	3312	1/1	0.95	0.12	41,41,41,41	0
56	MG	1A	3217	1/1	0.95	0.10	56,56,56,56	0
56	MG	2A	3161	1/1	0.95	0.08	38,38,38,38	0
56	MG	1A	3314	1/1	0.95	0.08	51,51,51,51	0
56	MG	1A	3160	1/1	0.95	0.08	46,46,46,46	0
56	MG	2A	3493	1/1	0.95	0.06	39,39,39,39	0
56	MG	1a	1659	1/1	0.95	0.08	52,52,52,52	0
56	MG	2A	3165	1/1	0.95	0.16	42,42,42,42	0
56	MG	1a	1862	1/1	0.95	0.10	35,35,35,35	0
56	MG	2A	3497	1/1	0.95	0.12	46,46,46,46	0
56	MG	1A	3880	1/1	0.95	0.16	38,38,38,38	0
56	MG	1A	3070	1/1	0.95	0.13	48,48,48,48	0
56	MG	1A	3031	1/1	0.95	0.14	60,60,60,60	0
56	MG	1A	3705	1/1	0.95	0.10	59,59,59,59	0
56	MG	1a	1868	1/1	0.95	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
56	MG	2A	3176	1/1	0.95	0.12	38,38,38,38	0
56	MG	1A	3075	1/1	0.95	0.07	46,46,46,46	0
56	MG	1A	3888	1/1	0.95	0.08	43,43,43,43	0
56	MG	2A	3511	1/1	0.95	0.11	40,40,40,40	0
56	MG	2A	3179	1/1	0.95	0.23	29,29,29,29	0
56	MG	1A	3003	1/1	0.95	0.07	44,44,44,44	0
56	MG	1A	3708	1/1	0.95	0.11	59,59,59,59	0
56	MG	2A	3183	1/1	0.95	0.14	46,46,46,46	0
56	MG	1A	3167	1/1	0.95	0.12	25,25,25,25	0
56	MG	1A	3321	1/1	0.95	0.15	33,33,33,33	0
56	MG	1A	3553	1/1	0.95	0.12	63,63,63,63	0
56	MG	1a	1877	1/1	0.95	0.10	52,52,52,52	0
56	MG	1A	3712	1/1	0.95	0.13	56,56,56,56	0
56	MG	1A	3045	1/1	0.95	0.06	39,39,39,39	0
56	MG	1A	3905	1/1	0.95	0.18	45,45,45,45	0
56	MG	1A	3171	1/1	0.95	0.27	51,51,51,51	0
56	MG	1A	3172	1/1	0.95	0.07	31,31,31,31	0
56	MG	1a	1883	1/1	0.95	0.15	53,53,53,53	0
56	MG	2A	3532	1/1	0.95	0.12	46,46,46,46	0
56	MG	2a	3245	1/1	0.95	0.12	64,64,64,64	0
56	MG	2a	3246	1/1	0.95	0.13	60,60,60,60	0
56	MG	2a	3247	1/1	0.95	0.14	50,50,50,50	0
56	MG	1A	3909	1/1	0.95	0.08	46,46,46,46	0
56	MG	1A	3912	1/1	0.95	0.11	56,56,56,56	0
56	MG	1A	3241	1/1	0.95	0.12	23,23,23,23	0
56	MG	1D	307	1/1	0.95	0.16	37,37,37,37	0
56	MG	2A	3542	1/1	0.95	0.12	56,56,56,56	0
56	MG	2A	3543	1/1	0.95	0.14	49,49,49,49	0
56	MG	2A	3545	1/1	0.95	0.10	53,53,53,53	0
56	MG	2A	3546	1/1	0.95	0.11	49,49,49,49	0
56	MG	1D	308	1/1	0.95	0.10	54,54,54,54	0
56	MG	2A	3204	1/1	0.95	0.19	53,53,53,53	0
56	MG	2e	3002	1/1	0.95	0.06	64,64,64,64	0
56	MG	2A	3553	1/1	0.95	0.11	41,41,41,41	0
56	MG	2A	3554	1/1	0.95	0.13	59,59,59,59	0
56	MG	2A	3205	1/1	0.95	0.10	43,43,43,43	0
56	MG	1a	1890	1/1	0.95	0.16	40,40,40,40	0
56	MG	1A	3327	1/1	0.95	0.07	44,44,44,44	0
56	MG	1A	3078	1/1	0.95	0.10	61,61,61,61	0
56	MG	2A	3212	1/1	0.95	0.19	34,34,34,34	0
56	MG	1A	3560	1/1	0.95	0.06	63,63,63,63	0
56	MG	1D	313	1/1	0.95	0.09	39,39,39,39	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2B	203	1/1	0.95	0.13	51,51,51,51	0
56	MG	1A	3921	1/1	0.95	0.06	45,45,45,45	0
56	MG	1A	3561	1/1	0.95	0.09	72,72,72,72	0
56	MG	2A	3218	1/1	0.95	0.21	54,54,54,54	0
56	MG	1A	3099	1/1	0.95	0.15	55,55,55,55	0
56	MG	1E	306	1/1	0.95	0.09	57,57,57,57	0
56	MG	1A	3735	1/1	0.95	0.09	63,63,63,63	0
56	MG	1a	1902	1/1	0.95	0.09	62,62,62,62	0
56	MG	1A	3332	1/1	0.95	0.10	35,35,35,35	0
57	ZN	2n	501	1/1	0.95	0.07	110,110,110,110	0
56	MG	1A	3848	1/1	0.96	0.07	45,45,45,45	0
56	MG	2A	3283	1/1	0.96	0.09	35,35,35,35	0
56	MG	1A	3656	1/1	0.96	0.09	20,20,20,20	0
56	MG	2A	3287	1/1	0.96	0.12	33,33,33,33	0
56	MG	2A	3033	1/1	0.96	0.22	47,47,47,47	0
56	MG	20	8001	1/1	0.96	0.07	62,62,62,62	0
56	MG	16	102	1/1	0.96	0.10	40,40,40,40	0
56	MG	1A	3538	1/1	0.96	0.08	39,39,39,39	0
56	MG	2A	3037	1/1	0.96	0.17	42,42,42,42	0
56	MG	2A	3038	1/1	0.96	0.06	45,45,45,45	0
56	MG	2A	3295	1/1	0.96	0.09	38,38,38,38	0
56	MG	2A	3039	1/1	0.96	0.17	38,38,38,38	0
56	MG	2A	3297	1/1	0.96	0.07	40,40,40,40	0
56	MG	1A	3661	1/1	0.96	0.07	30,30,30,30	0
56	MG	2A	3299	1/1	0.96	0.12	64,64,64,64	0
56	MG	1A	3131	1/1	0.96	0.12	31,31,31,31	0
56	MG	2A	3302	1/1	0.96	0.11	41,41,41,41	0
56	MG	2A	3042	1/1	0.96	0.16	46,46,46,46	0
56	MG	1A	3133	1/1	0.96	0.24	48,48,48,48	0
56	MG	2A	3044	1/1	0.96	0.12	51,51,51,51	0
56	MG	1A	3672	1/1	0.96	0.12	29,29,29,29	0
56	MG	1A	3676	1/1	0.96	0.11	40,40,40,40	0
56	MG	2A	3047	1/1	0.96	0.12	37,37,37,37	0
56	MG	1A	3679	1/1	0.96	0.06	38,38,38,38	0
56	MG	2A	3049	1/1	0.96	0.12	56,56,56,56	0
56	MG	2a	3016	1/1	0.96	0.27	65,65,65,65	0
56	MG	2A	3050	1/1	0.96	0.29	55,55,55,55	0
56	MG	1a	1790	1/1	0.96	0.09	62,62,62,62	0
56	MG	1A	3463	1/1	0.96	0.07	39,39,39,39	0
56	MG	1A	3682	1/1	0.96	0.12	29,29,29,29	0
56	MG	2a	3021	1/1	0.96	0.08	58,58,58,58	0
56	MG	1A	3684	1/1	0.96	0.08	50,50,50,50	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3319	1/1	0.96	0.20	51,51,51,51	0
56	MG	1A	3686	1/1	0.96	0.11	28,28,28,28	0
56	MG	1A	3266	1/1	0.96	0.10	18,18,18,18	0
56	MG	1A	3116	1/1	0.96	0.05	52,52,52,52	0
56	MG	2A	3324	1/1	0.96	0.04	43,43,43,43	0
56	MG	2A	3058	1/1	0.96	0.10	51,51,51,51	0
56	MG	2a	3029	1/1	0.96	0.09	66,66,66,66	0
56	MG	2A	3060	1/1	0.96	0.12	40,40,40,40	0
56	MG	1A	3268	1/1	0.96	0.15	23,23,23,23	0
56	MG	1A	3269	1/1	0.96	0.12	53,53,53,53	0
56	MG	1A	3470	1/1	0.96	0.07	51,51,51,51	0
56	MG	2A	3338	1/1	0.96	0.12	46,46,46,46	0
56	MG	2a	3035	1/1	0.96	0.11	59,59,59,59	0
56	MG	1A	3166	1/1	0.96	0.13	35,35,35,35	0
56	MG	1A	3137	1/1	0.96	0.10	36,36,36,36	0
56	MG	1A	3051	1/1	0.96	0.28	47,47,47,47	0
56	MG	1A	3872	1/1	0.96	0.08	39,39,39,39	0
56	MG	1a	1617	1/1	0.96	0.18	55,55,55,55	0
56	MG	1A	3474	1/1	0.96	0.09	58,58,58,58	0
56	MG	1A	3875	1/1	0.96	0.07	38,38,38,38	0
56	MG	1A	3475	1/1	0.96	0.11	66,66,66,66	0
56	MG	1A	3052	1/1	0.96	0.11	51,51,51,51	0
56	MG	1A	3334	1/1	0.96	0.09	73,73,73,73	0
56	MG	2a	3046	1/1	0.96	0.05	70,70,70,70	0
56	MG	1A	3140	1/1	0.96	0.08	36,36,36,36	0
56	MG	1A	3141	1/1	0.96	0.09	39,39,39,39	0
56	MG	1A	3279	1/1	0.96	0.20	41,41,41,41	0
56	MG	1A	3406	1/1	0.96	0.13	49,49,49,49	0
56	MG	1A	4068	1/1	0.96	0.10	39,39,39,39	0
56	MG	1A	3143	1/1	0.96	0.06	34,34,34,34	0
56	MG	1A	4071	1/1	0.96	0.15	44,44,44,44	0
56	MG	1A	3886	1/1	0.96	0.05	42,42,42,42	0
56	MG	1A	3714	1/1	0.96	0.15	49,49,49,49	0
56	MG	1A	3042	1/1	0.96	0.10	42,42,42,42	0
56	MG	1A	3717	1/1	0.96	0.15	55,55,55,55	0
56	MG	1A	3340	1/1	0.96	0.10	59,59,59,59	0
56	MG	1a	1637	1/1	0.96	0.10	51,51,51,51	0
56	MG	1A	3564	1/1	0.96	0.06	36,36,36,36	0
56	MG	2A	3088	1/1	0.96	0.06	44,44,44,44	0
56	MG	1A	3720	1/1	0.96	0.12	45,45,45,45	0
56	MG	2A	3090	1/1	0.96	0.20	58,58,58,58	0
56	MG	1A	4084	1/1	0.96	0.24	32,32,32,32	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3721	1/1	0.96	0.06	45,45,45,45	0
56	MG	1a	1642	1/1	0.96	0.04	58,58,58,58	0
56	MG	2A	3377	1/1	0.96	0.07	33,33,33,33	0
56	MG	2A	3094	1/1	0.96	0.11	43,43,43,43	0
56	MG	1A	3565	1/1	0.96	0.08	41,41,41,41	0
56	MG	1A	4087	1/1	0.96	0.17	42,42,42,42	0
56	MG	2A	3385	1/1	0.96	0.07	61,61,61,61	0
56	MG	2A	3387	1/1	0.96	0.20	38,38,38,38	0
56	MG	1A	3726	1/1	0.96	0.07	46,46,46,46	0
56	MG	1a	1836	1/1	0.96	0.07	51,51,51,51	0
56	MG	1A	3903	1/1	0.96	0.18	71,71,71,71	0
56	MG	1A	3487	1/1	0.96	0.10	40,40,40,40	0
56	MG	1A	3282	1/1	0.96	0.16	54,54,54,54	0
56	MG	1B	203	1/1	0.96	0.11	39,39,39,39	0
56	MG	2a	3079	1/1	0.96	0.07	52,52,52,52	0
56	MG	2A	3394	1/1	0.96	0.16	54,54,54,54	0
56	MG	1a	1843	1/1	0.96	0.09	55,55,55,55	0
56	MG	1A	3220	1/1	0.96	0.30	41,41,41,41	0
56	MG	1A	3574	1/1	0.96	0.11	37,37,37,37	0
56	MG	2a	3085	1/1	0.96	0.15	48,48,48,48	0
56	MG	2a	3086	1/1	0.96	0.08	50,50,50,50	0
56	MG	1a	1847	1/1	0.96	0.07	53,53,53,53	0
56	MG	2A	3111	1/1	0.96	0.15	36,36,36,36	0
56	MG	1A	3910	1/1	0.96	0.10	61,61,61,61	0
56	MG	1a	1849	1/1	0.96	0.06	38,38,38,38	0
56	MG	2A	3115	1/1	0.96	0.15	41,41,41,41	0
56	MG	1A	3911	1/1	0.96	0.09	30,30,30,30	0
56	MG	1A	3222	1/1	0.96	0.10	19,19,19,19	0
56	MG	2A	3407	1/1	0.96	0.14	46,46,46,46	0
56	MG	1A	3913	1/1	0.96	0.06	28,28,28,28	0
56	MG	1A	3739	1/1	0.96	0.13	34,34,34,34	0
56	MG	1a	1658	1/1	0.96	0.10	58,58,58,58	0
56	MG	2A	3411	1/1	0.96	0.04	47,47,47,47	0
56	MG	1A	3915	1/1	0.96	0.14	52,52,52,52	0
56	MG	2a	3102	1/1	0.96	0.12	44,44,44,44	0
56	MG	2A	3123	1/1	0.96	0.19	57,57,57,57	0
56	MG	1A	3223	1/1	0.96	0.14	29,29,29,29	0
56	MG	1A	3917	1/1	0.96	0.04	35,35,35,35	0
56	MG	2A	3126	1/1	0.96	0.26	43,43,43,43	0
56	MG	1A	3577	1/1	0.96	0.11	30,30,30,30	0
56	MG	1A	3349	1/1	0.96	0.07	35,35,35,35	0
56	MG	1A	3745	1/1	0.96	0.06	57,57,57,57	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3287	1/1	0.96	0.16	38,38,38,38	0
56	MG	2A	3422	1/1	0.96	0.09	73,73,73,73	0
56	MG	1A	3923	1/1	0.96	0.08	40,40,40,40	0
56	MG	1A	3924	1/1	0.96	0.12	52,52,52,52	0
56	MG	2A	3427	1/1	0.96	0.22	54,54,54,54	0
56	MG	1A	3288	1/1	0.96	0.22	42,42,42,42	0
56	MG	1A	3352	1/1	0.96	0.20	57,57,57,57	0
56	MG	2A	3430	1/1	0.96	0.21	50,50,50,50	0
56	MG	1D	302	1/1	0.96	0.13	52,52,52,52	0
56	MG	2A	3137	1/1	0.96	0.08	33,33,33,33	0
56	MG	1D	304	1/1	0.96	0.12	34,34,34,34	0
56	MG	1A	3928	1/1	0.96	0.11	29,29,29,29	0
56	MG	1A	3929	1/1	0.96	0.07	29,29,29,29	0
56	MG	1A	3930	1/1	0.96	0.09	55,55,55,55	0
56	MG	1a	1676	1/1	0.96	0.07	55,55,55,55	0
56	MG	1a	1875	1/1	0.96	0.08	58,58,58,58	0
56	MG	2A	3148	1/1	0.96	0.11	38,38,38,38	0
56	MG	1A	3931	1/1	0.96	0.06	32,32,32,32	0
56	MG	2A	3441	1/1	0.96	0.12	55,55,55,55	0
56	MG	2A	3150	1/1	0.96	0.11	30,30,30,30	0
56	MG	1D	309	1/1	0.96	0.14	56,56,56,56	0
56	MG	1A	3932	1/1	0.96	0.10	59,59,59,59	0
56	MG	1a	1681	1/1	0.96	0.12	47,47,47,47	0
56	MG	2A	3155	1/1	0.96	0.17	37,37,37,37	0
56	MG	2A	3157	1/1	0.96	0.06	31,31,31,31	0
56	MG	2A	3450	1/1	0.96	0.06	56,56,56,56	0
56	MG	1A	3933	1/1	0.96	0.08	46,46,46,46	0
56	MG	2A	3453	1/1	0.96	0.08	31,31,31,31	0
56	MG	1a	1683	1/1	0.96	0.07	58,58,58,58	0
56	MG	1A	3584	1/1	0.96	0.07	39,39,39,39	0
56	MG	1A	3354	1/1	0.96	0.10	84,84,84,84	0
56	MG	1a	1884	1/1	0.96	0.10	46,46,46,46	0
56	MG	1A	3179	1/1	0.96	0.16	22,22,22,22	0
56	MG	1A	3498	1/1	0.96	0.09	53,53,53,53	0
56	MG	1A	3048	1/1	0.96	0.17	51,51,51,51	0
56	MG	1A	3294	1/1	0.96	0.17	55,55,55,55	0
56	MG	1A	3591	1/1	0.96	0.12	32,32,32,32	0
56	MG	1E	308	1/1	0.96	0.17	35,35,35,35	0
56	MG	2A	3468	1/1	0.96	0.10	47,47,47,47	0
56	MG	2A	3169	1/1	0.96	0.11	37,37,37,37	0
56	MG	1A	3296	1/1	0.96	0.09	49,49,49,49	0
56	MG	2A	3472	1/1	0.96	0.06	50,50,50,50	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1695	1/1	0.96	0.17	53,53,53,53	0
56	MG	1a	1893	1/1	0.96	0.18	69,69,69,69	0
56	MG	2A	3175	1/1	0.96	0.14	34,34,34,34	0
56	MG	1E	312	1/1	0.96	0.05	44,44,44,44	0
56	MG	1F	301	1/1	0.96	0.19	30,30,30,30	0
56	MG	1A	3944	1/1	0.96	0.20	39,39,39,39	0
56	MG	1A	3945	1/1	0.96	0.17	30,30,30,30	0
56	MG	2A	3180	1/1	0.96	0.09	29,29,29,29	0
56	MG	1A	3946	1/1	0.96	0.13	24,24,24,24	0
56	MG	1A	3231	1/1	0.96	0.06	42,42,42,42	0
56	MG	1A	3122	1/1	0.96	0.07	56,56,56,56	0
56	MG	1a	1901	1/1	0.96	0.12	47,47,47,47	0
56	MG	1A	3949	1/1	0.96	0.07	35,35,35,35	0
56	MG	1A	3596	1/1	0.96	0.16	44,44,44,44	0
56	MG	1G	3002	1/1	0.96	0.14	52,52,52,52	0
56	MG	1A	3767	1/1	0.96	0.10	33,33,33,33	0
56	MG	1A	3597	1/1	0.96	0.14	71,71,71,71	0
56	MG	1a	1907	1/1	0.96	0.13	36,36,36,36	0
56	MG	2a	3179	1/1	0.96	0.05	58,58,58,58	0
56	MG	1a	1711	1/1	0.96	0.11	62,62,62,62	0
56	MG	1A	3957	1/1	0.96	0.11	34,34,34,34	0
56	MG	2a	3185	1/1	0.96	0.12	57,57,57,57	0
56	MG	1A	3363	1/1	0.96	0.06	49,49,49,49	0
56	MG	1A	3428	1/1	0.96	0.06	43,43,43,43	0
56	MG	1A	3602	1/1	0.96	0.07	41,41,41,41	0
56	MG	1A	3603	1/1	0.96	0.14	62,62,62,62	0
56	MG	1A	3507	1/1	0.96	0.06	45,45,45,45	0
56	MG	2A	3500	1/1	0.96	0.06	45,45,45,45	0
56	MG	2A	3199	1/1	0.96	0.12	21,21,21,21	0
56	MG	2a	3194	1/1	0.96	0.16	44,44,44,44	0
56	MG	2A	3503	1/1	0.96	0.06	37,37,37,37	0
56	MG	1A	3234	1/1	0.96	0.09	46,46,46,46	0
56	MG	1f	3001	1/1	0.96	0.16	57,57,57,57	0
56	MG	2A	3506	1/1	0.96	0.09	51,51,51,51	0
56	MG	1A	3148	1/1	0.96	0.16	51,51,51,51	0
56	MG	1A	3149	1/1	0.96	0.20	32,32,32,32	0
56	MG	1a	1724	1/1	0.96	0.07	62,62,62,62	0
56	MG	1A	3973	1/1	0.96	0.10	50,50,50,50	0
56	MG	1A	3303	1/1	0.96	0.12	50,50,50,50	0
56	MG	1A	3304	1/1	0.96	0.10	45,45,45,45	0
56	MG	1A	3791	1/1	0.96	0.19	51,51,51,51	0
56	MG	1A	3097	1/1	0.96	0.15	49,49,49,49	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1731	1/1	0.96	0.10	37,37,37,37	0
56	MG	2A	3517	1/1	0.96	0.06	47,47,47,47	0
56	MG	1p	3001	1/1	0.96	0.21	52,52,52,52	0
56	MG	1A	3151	1/1	0.96	0.08	45,45,45,45	0
56	MG	1q	201	1/1	0.96	0.17	58,58,58,58	0
56	MG	1A	3980	1/1	0.96	0.18	59,59,59,59	0
56	MG	2A	3524	1/1	0.96	0.08	50,50,50,50	0
56	MG	1A	3110	1/1	0.96	0.10	65,65,65,65	0
56	MG	1A	3799	1/1	0.96	0.06	20,20,20,20	0
56	MG	2A	3225	1/1	0.96	0.16	58,58,58,58	0
56	MG	1A	3197	1/1	0.96	0.06	23,23,23,23	0
56	MG	2A	3227	1/1	0.96	0.11	38,38,38,38	0
56	MG	1A	3441	1/1	0.96	0.08	49,49,49,49	0
56	MG	2A	3534	1/1	0.96	0.12	39,39,39,39	0
56	MG	2a	3224	1/1	0.96	0.10	60,60,60,60	0
56	MG	1A	3988	1/1	0.96	0.25	39,39,39,39	0
56	MG	1A	3518	1/1	0.96	0.07	51,51,51,51	0
56	MG	2A	3231	1/1	0.96	0.12	57,57,57,57	0
56	MG	2A	3540	1/1	0.96	0.10	46,46,46,46	0
56	MG	1A	3625	1/1	0.96	0.05	31,31,31,31	0
56	MG	1A	3627	1/1	0.96	0.10	41,41,41,41	0
56	MG	1a	1743	1/1	0.96	0.12	44,44,44,44	0
56	MG	2A	3544	1/1	0.96	0.06	48,48,48,48	0
56	MG	1A	3629	1/1	0.96	0.10	33,33,33,33	0
56	MG	1T	201	1/1	0.96	0.09	41,41,41,41	0
56	MG	1A	3814	1/1	0.96	0.08	49,49,49,49	0
56	MG	1A	3072	1/1	0.96	0.10	46,46,46,46	0
56	MG	2A	3550	1/1	0.96	0.06	39,39,39,39	0
56	MG	2A	3551	1/1	0.96	0.08	32,32,32,32	0
56	MG	2A	3552	1/1	0.96	0.13	44,44,44,44	0
56	MG	1A	3819	1/1	0.96	0.05	43,43,43,43	0
56	MG	1T	205	1/1	0.96	0.06	56,56,56,56	0
56	MG	2a	3244	1/1	0.96	0.09	64,64,64,64	0
56	MG	1A	3247	1/1	0.96	0.17	35,35,35,35	0
56	MG	2A	3556	1/1	0.96	0.06	55,55,55,55	0
56	MG	1z	8001	1/1	0.96	0.11	56,56,56,56	0
56	MG	1A	3823	1/1	0.96	0.10	17,17,17,17	0
56	MG	1A	3248	1/1	0.96	0.09	47,47,47,47	0
56	MG	2A	3560	1/1	0.96	0.08	41,41,41,41	0
56	MG	1A	4001	1/1	0.96	0.11	35,35,35,35	0
56	MG	1A	3377	1/1	0.96	0.14	24,24,24,24	0
56	MG	1A	3154	1/1	0.96	0.21	30,30,30,30	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3253	1/1	0.96	0.06	59,59,59,59	0
56	MG	1A	3255	1/1	0.96	0.06	45,45,45,45	0
56	MG	1A	3126	1/1	0.96	0.10	27,27,27,27	0
56	MG	1A	3829	1/1	0.96	0.06	18,18,18,18	0
56	MG	1A	3079	1/1	0.96	0.10	53,53,53,53	0
56	MG	1A	4011	1/1	0.96	0.08	68,68,68,68	0
56	MG	1A	4013	1/1	0.96	0.09	42,42,42,42	0
56	MG	1A	3258	1/1	0.96	0.16	63,63,63,63	0
56	MG	2I	3003	1/1	0.96	0.14	57,57,57,57	0
56	MG	1A	3073	1/1	0.96	0.08	46,46,46,46	0
56	MG	1A	3260	1/1	0.96	0.12	31,31,31,31	0
56	MG	1A	3385	1/1	0.96	0.14	45,45,45,45	0
56	MG	1A	3645	1/1	0.96	0.07	36,36,36,36	0
56	MG	1A	3203	1/1	0.96	0.28	47,47,47,47	0
56	MG	2A	3269	1/1	0.96	0.13	38,38,38,38	0
56	MG	1A	3081	1/1	0.96	0.15	40,40,40,40	0
56	MG	1a	1771	1/1	0.96	0.06	48,48,48,48	0
56	MG	2x	102	1/1	0.96	0.27	68,68,68,68	0
56	MG	2A	3022	1/1	0.96	0.05	46,46,46,46	0
56	MG	1A	3840	1/1	0.96	0.15	59,59,59,59	0
56	MG	1A	3841	1/1	0.96	0.06	44,44,44,44	0
56	MG	1A	3458	1/1	0.96	0.10	48,48,48,48	0
56	MG	1A	3650	1/1	0.96	0.09	28,28,28,28	0
56	MG	1A	3654	1/1	0.96	0.08	44,44,44,44	0
56	MG	2Q	3003	1/1	0.96	0.15	44,44,44,44	0
56	MG	15	103	1/1	0.96	0.19	28,28,28,28	0
56	MG	2A	3206	1/1	0.97	0.15	50,50,50,50	0
56	MG	1A	3192	1/1	0.97	0.18	46,46,46,46	0
56	MG	1A	3839	1/1	0.97	0.07	18,18,18,18	0
56	MG	2A	3210	1/1	0.97	0.21	31,31,31,31	0
56	MG	2A	3023	1/1	0.97	0.14	45,45,45,45	0
56	MG	2A	3024	1/1	0.97	0.11	55,55,55,55	0
56	MG	1A	3011	1/1	0.97	0.06	47,47,47,47	0
56	MG	1B	228	1/1	0.97	0.07	36,36,36,36	0
56	MG	1a	1635	1/1	0.97	0.08	56,56,56,56	0
56	MG	2A	3216	1/1	0.97	0.08	44,44,44,44	0
56	MG	1A	3455	1/1	0.97	0.06	58,58,58,58	0
56	MG	2A	3444	1/1	0.97	0.08	63,63,63,63	0
56	MG	1A	3274	1/1	0.97	0.15	47,47,47,47	0
56	MG	1A	3275	1/1	0.97	0.09	41,41,41,41	0
56	MG	1A	3357	1/1	0.97	0.05	54,54,54,54	0
56	MG	1A	3459	1/1	0.97	0.13	49,49,49,49	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3211	1/1	0.97	0.17	37,37,37,37	0
56	MG	1A	3978	1/1	0.97	0.07	36,36,36,36	0
56	MG	1A	3461	1/1	0.97	0.17	48,48,48,48	0
56	MG	1A	3851	1/1	0.97	0.06	65,65,65,65	0
56	MG	1A	3852	1/1	0.97	0.08	30,30,30,30	0
56	MG	1A	3277	1/1	0.97	0.11	34,34,34,34	0
56	MG	1A	3213	1/1	0.97	0.23	44,44,44,44	0
56	MG	2A	3460	1/1	0.97	0.05	45,45,45,45	0
56	MG	1a	1648	1/1	0.97	0.10	48,48,48,48	0
56	MG	1A	3985	1/1	0.97	0.06	34,34,34,34	0
56	MG	1A	3986	1/1	0.97	0.13	49,49,49,49	0
56	MG	1A	3523	1/1	0.97	0.14	50,50,50,50	0
56	MG	1A	3053	1/1	0.97	0.10	36,36,36,36	0
56	MG	1E	303	1/1	0.97	0.06	31,31,31,31	0
56	MG	2A	3237	1/1	0.97	0.10	37,37,37,37	0
56	MG	1A	3243	1/1	0.97	0.10	38,38,38,38	0
56	MG	2A	3470	1/1	0.97	0.05	64,64,64,64	0
56	MG	1A	3412	1/1	0.97	0.07	42,42,42,42	0
56	MG	1A	3413	1/1	0.97	0.10	60,60,60,60	0
56	MG	1A	3528	1/1	0.97	0.13	42,42,42,42	0
56	MG	2a	3089	1/1	0.97	0.05	46,46,46,46	0
56	MG	2A	3242	1/1	0.97	0.14	22,22,22,22	0
56	MG	1E	309	1/1	0.97	0.05	54,54,54,54	0
56	MG	1A	3724	1/1	0.97	0.09	38,38,38,38	0
56	MG	1E	311	1/1	0.97	0.09	40,40,40,40	0
56	MG	1A	3468	1/1	0.97	0.07	39,39,39,39	0
56	MG	1A	3611	1/1	0.97	0.07	35,35,35,35	0
56	MG	1A	3469	1/1	0.97	0.16	33,33,33,33	0
56	MG	1A	3244	1/1	0.97	0.05	35,35,35,35	0
56	MG	1a	1665	1/1	0.97	0.07	50,50,50,50	0
56	MG	2A	3059	1/1	0.97	0.09	37,37,37,37	0
56	MG	1A	3731	1/1	0.97	0.06	45,45,45,45	0
56	MG	1A	3245	1/1	0.97	0.11	18,18,18,18	0
56	MG	2a	3103	1/1	0.97	0.12	48,48,48,48	0
56	MG	1a	1838	1/1	0.97	0.13	39,39,39,39	0
56	MG	1A	3734	1/1	0.97	0.06	46,46,46,46	0
56	MG	1A	3108	1/1	0.97	0.06	52,52,52,52	0
56	MG	2A	3260	1/1	0.97	0.13	39,39,39,39	0
56	MG	1F	308	1/1	0.97	0.05	36,36,36,36	0
56	MG	1A	4004	1/1	0.97	0.10	49,49,49,49	0
56	MG	1A	3284	1/1	0.97	0.11	52,52,52,52	0
56	MG	1A	3536	1/1	0.97	0.18	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3196	1/1	0.97	0.16	37,37,37,37	0
56	MG	1A	3874	1/1	0.97	0.04	28,28,28,28	0
56	MG	1A	3742	1/1	0.97	0.07	52,52,52,52	0
56	MG	1A	4012	1/1	0.97	0.05	47,47,47,47	0
56	MG	1A	3109	1/1	0.97	0.04	45,45,45,45	0
56	MG	1A	3253	1/1	0.97	0.16	32,32,32,32	0
56	MG	1A	3325	1/1	0.97	0.05	41,41,41,41	0
56	MG	1N	3004	1/1	0.97	0.03	26,26,26,26	0
56	MG	1A	3068	1/1	0.97	0.17	45,45,45,45	0
56	MG	2a	3125	1/1	0.97	0.15	33,33,33,33	0
56	MG	2A	3276	1/1	0.97	0.04	60,60,60,60	0
56	MG	1a	1856	1/1	0.97	0.09	41,41,41,41	0
56	MG	1A	3219	1/1	0.97	0.10	56,56,56,56	0
56	MG	1A	3480	1/1	0.97	0.08	54,54,54,54	0
56	MG	1A	3882	1/1	0.97	0.12	36,36,36,36	0
56	MG	2a	3131	1/1	0.97	0.29	48,48,48,48	0
56	MG	1A	3883	1/1	0.97	0.09	36,36,36,36	0
56	MG	2A	3282	1/1	0.97	0.06	35,35,35,35	0
56	MG	1A	4021	1/1	0.97	0.09	59,59,59,59	0
56	MG	1A	3630	1/1	0.97	0.08	37,37,37,37	0
56	MG	1a	1691	1/1	0.97	0.05	66,66,66,66	0
56	MG	1A	3054	1/1	0.97	0.15	53,53,53,53	0
56	MG	1A	4024	1/1	0.97	0.10	39,39,39,39	0
56	MG	1A	3545	1/1	0.97	0.11	50,50,50,50	0
56	MG	2A	3292	1/1	0.97	0.06	21,21,21,21	0
56	MG	1P	205	1/1	0.97	0.07	53,53,53,53	0
56	MG	1P	206	1/1	0.97	0.13	42,42,42,42	0
56	MG	1A	3546	1/1	0.97	0.09	66,66,66,66	0
56	MG	1A	3330	1/1	0.97	0.13	39,39,39,39	0
56	MG	2A	3527	1/1	0.97	0.07	39,39,39,39	0
56	MG	1A	4028	1/1	0.97	0.14	63,63,63,63	0
56	MG	1A	3291	1/1	0.97	0.17	41,41,41,41	0
56	MG	2A	3530	1/1	0.97	0.19	43,43,43,43	0
56	MG	2A	3095	1/1	0.97	0.07	46,46,46,46	0
56	MG	1Q	207	1/1	0.97	0.17	41,41,41,41	0
56	MG	1A	3293	1/1	0.97	0.10	11,11,11,11	0
56	MG	2A	3303	1/1	0.97	0.11	30,30,30,30	0
56	MG	1A	3637	1/1	0.97	0.06	43,43,43,43	0
56	MG	2a	3154	1/1	0.97	0.10	71,71,71,71	0
56	MG	2A	3538	1/1	0.97	0.12	39,39,39,39	0
56	MG	1A	3892	1/1	0.97	0.06	46,46,46,46	0
56	MG	2A	3100	1/1	0.97	0.05	35,35,35,35	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3307	1/1	0.97	0.15	60,60,60,60	0
56	MG	1A	3102	1/1	0.97	0.22	49,49,49,49	0
56	MG	1A	3551	1/1	0.97	0.10	55,55,55,55	0
56	MG	1a	1708	1/1	0.97	0.10	67,67,67,67	0
56	MG	1R	206	1/1	0.97	0.11	32,32,32,32	0
56	MG	2A	3105	1/1	0.97	0.15	28,28,28,28	0
56	MG	1A	3295	1/1	0.97	0.10	42,42,42,42	0
56	MG	2A	3548	1/1	0.97	0.06	38,38,38,38	0
56	MG	1A	3900	1/1	0.97	0.12	39,39,39,39	0
56	MG	2A	3108	1/1	0.97	0.15	52,52,52,52	0
56	MG	1A	3182	1/1	0.97	0.17	32,32,32,32	0
56	MG	1A	3902	1/1	0.97	0.13	42,42,42,42	0
56	MG	1A	3642	1/1	0.97	0.13	38,38,38,38	0
56	MG	1A	4040	1/1	0.97	0.13	32,32,32,32	0
56	MG	1A	3431	1/1	0.97	0.06	43,43,43,43	0
56	MG	1A	3224	1/1	0.97	0.05	33,33,33,33	0
56	MG	2A	3323	1/1	0.97	0.13	43,43,43,43	0
56	MG	1A	4045	1/1	0.97	0.06	23,23,23,23	0
56	MG	2A	3325	1/1	0.97	0.11	48,48,48,48	0
56	MG	1U	206	1/1	0.97	0.29	38,38,38,38	0
56	MG	1a	1722	1/1	0.97	0.13	47,47,47,47	0
56	MG	2a	3180	1/1	0.97	0.13	46,46,46,46	0
56	MG	2a	3181	1/1	0.97	0.08	58,58,58,58	0
56	MG	2A	3119	1/1	0.97	0.08	35,35,35,35	0
56	MG	1A	3227	1/1	0.97	0.11	26,26,26,26	0
56	MG	2A	3335	1/1	0.97	0.09	35,35,35,35	0
56	MG	1a	1725	1/1	0.97	0.20	51,51,51,51	0
56	MG	1A	3434	1/1	0.97	0.09	50,50,50,50	0
56	MG	1W	3002	1/1	0.97	0.14	35,35,35,35	0
56	MG	1A	3775	1/1	0.97	0.06	31,31,31,31	0
56	MG	1A	3776	1/1	0.97	0.07	31,31,31,31	0
56	MG	1A	3647	1/1	0.97	0.08	19,19,19,19	0
56	MG	1A	3778	1/1	0.97	0.07	19,19,19,19	0
56	MG	1A	3299	1/1	0.97	0.14	59,59,59,59	0
56	MG	2D	304	1/1	0.97	0.06	36,36,36,36	0
56	MG	1A	3183	1/1	0.97	0.13	28,28,28,28	0
56	MG	2A	3349	1/1	0.97	0.06	67,67,67,67	0
56	MG	1A	4055	1/1	0.97	0.06	43,43,43,43	0
56	MG	1A	4056	1/1	0.97	0.04	29,29,29,29	0
56	MG	2A	3132	1/1	0.97	0.08	45,45,45,45	0
56	MG	1A	3783	1/1	0.97	0.08	28,28,28,28	0
56	MG	2A	3356	1/1	0.97	0.06	38,38,38,38	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3071	1/1	0.97	0.23	37,37,37,37	0
56	MG	1A	3651	1/1	0.97	0.06	14,14,14,14	0
56	MG	1A	3788	1/1	0.97	0.13	30,30,30,30	0
56	MG	2O	203	1/1	0.97	0.06	41,41,41,41	0
56	MG	1A	3789	1/1	0.97	0.07	14,14,14,14	0
56	MG	1I	103	1/1	0.97	0.06	42,42,42,42	0
56	MG	2A	3140	1/1	0.97	0.14	33,33,33,33	0
56	MG	1A	3790	1/1	0.97	0.09	19,19,19,19	0
56	MG	1A	3230	1/1	0.97	0.14	35,35,35,35	0
56	MG	1A	3388	1/1	0.97	0.07	55,55,55,55	0
56	MG	1A	3925	1/1	0.97	0.13	56,56,56,56	0
56	MG	2T	201	1/1	0.97	0.06	41,41,41,41	0
56	MG	2A	3146	1/1	0.97	0.07	35,35,35,35	0
56	MG	1A	3658	1/1	0.97	0.04	13,13,13,13	0
56	MG	1A	3440	1/1	0.97	0.06	39,39,39,39	0
56	MG	1A	3342	1/1	0.97	0.13	53,53,53,53	0
56	MG	1A	3665	1/1	0.97	0.13	24,24,24,24	0
56	MG	1I	201	1/1	0.97	0.07	35,35,35,35	0
56	MG	1A	3036	1/1	0.97	0.17	40,40,40,40	0
56	MG	28	102	1/1	0.97	0.15	50,50,50,50	0
56	MG	16	104	1/1	0.97	0.06	53,53,53,53	0
56	MG	2a	3227	1/1	0.97	0.09	62,62,62,62	0
56	MG	16	105	1/1	0.97	0.10	44,44,44,44	0
56	MG	2A	3156	1/1	0.97	0.09	23,23,23,23	0
56	MG	17	101	1/1	0.97	0.05	26,26,26,26	0
56	MG	2A	3379	1/1	0.97	0.05	35,35,35,35	0
56	MG	1A	4073	1/1	0.97	0.25	48,48,48,48	0
56	MG	1A	3667	1/1	0.97	0.13	14,14,14,14	0
56	MG	2A	3382	1/1	0.97	0.08	46,46,46,46	0
56	MG	1A	3805	1/1	0.97	0.08	49,49,49,49	0
56	MG	1a	1758	1/1	0.97	0.14	67,67,67,67	0
56	MG	18	3003	1/1	0.97	0.08	35,35,35,35	0
56	MG	19	101	1/1	0.97	0.14	51,51,51,51	0
56	MG	1q	203	1/1	0.97	0.19	53,53,53,53	0
56	MG	1A	3669	1/1	0.97	0.06	17,17,17,17	0
56	MG	1A	3566	1/1	0.97	0.09	34,34,34,34	0
56	MG	19	106	1/1	0.97	0.07	45,45,45,45	0
56	MG	1A	3812	1/1	0.97	0.04	48,48,48,48	0
56	MG	1A	4080	1/1	0.97	0.23	36,36,36,36	0
56	MG	2A	3170	1/1	0.97	0.06	33,33,33,33	0
56	MG	2A	3398	1/1	0.97	0.07	32,32,32,32	0
56	MG	1A	3344	1/1	0.97	0.07	42,42,42,42	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1604	1/1	0.97	0.08	60,60,60,60	0
56	MG	1A	3675	1/1	0.97	0.10	42,42,42,42	0
56	MG	1A	3037	1/1	0.97	0.17	40,40,40,40	0
56	MG	1a	1770	1/1	0.97	0.06	70,70,70,70	0
56	MG	1A	3678	1/1	0.97	0.06	39,39,39,39	0
56	MG	1A	3503	1/1	0.97	0.07	30,30,30,30	0
56	MG	1A	3570	1/1	0.97	0.13	46,46,46,46	0
56	MG	1A	3571	1/1	0.97	0.12	32,32,32,32	0
56	MG	1A	3445	1/1	0.97	0.07	46,46,46,46	0
56	MG	1A	3346	1/1	0.97	0.06	55,55,55,55	0
56	MG	1A	3347	1/1	0.97	0.05	56,56,56,56	0
56	MG	1a	1778	1/1	0.97	0.04	48,48,48,48	0
56	MG	2A	3185	1/1	0.97	0.12	32,32,32,32	0
56	MG	1A	3233	1/1	0.97	0.11	36,36,36,36	0
56	MG	1a	1780	1/1	0.97	0.05	56,56,56,56	0
56	MG	1A	3168	1/1	0.97	0.12	35,35,35,35	0
56	MG	1A	3950	1/1	0.97	0.14	64,64,64,64	0
56	MG	1A	3189	1/1	0.97	0.14	34,34,34,34	0
56	MG	1A	3399	1/1	0.97	0.20	45,45,45,45	0
56	MG	1A	3698	1/1	0.97	0.06	26,26,26,26	0
56	MG	1B	211	1/1	0.97	0.05	39,39,39,39	0
56	MG	1A	3833	1/1	0.97	0.04	50,50,50,50	0
56	MG	1A	3581	1/1	0.97	0.15	54,54,54,54	0
56	MG	1A	3962	1/1	0.97	0.08	14,14,14,14	0
56	MG	2A	3426	1/1	0.97	0.05	46,46,46,46	0
56	MG	1B	215	1/1	0.97	0.08	74,74,74,74	0
56	MG	1A	3582	1/1	0.97	0.09	25,25,25,25	0
56	MG	1B	218	1/1	0.97	0.07	34,34,34,34	0
56	MG	1A	3702	1/1	0.97	0.05	20,20,20,20	0
56	MG	1B	222	1/1	0.97	0.09	59,59,59,59	0
57	ZN	29	501	1/1	0.97	0.05	66,66,66,66	0
56	MG	1A	3190	1/1	0.97	0.10	42,42,42,42	0
56	MG	1A	3809	1/1	0.98	0.09	33,33,33,33	0
56	MG	2a	3105	1/1	0.98	0.10	62,62,62,62	0
56	MG	1A	4089	1/1	0.98	0.10	29,29,29,29	0
56	MG	2a	3108	1/1	0.98	0.10	68,68,68,68	0
56	MG	2A	3383	1/1	0.98	0.08	29,29,29,29	0
56	MG	2a	3110	1/1	0.98	0.07	55,55,55,55	0
56	MG	2A	3384	1/1	0.98	0.05	39,39,39,39	0
56	MG	1A	3810	1/1	0.98	0.04	30,30,30,30	0
56	MG	2A	3386	1/1	0.98	0.09	37,37,37,37	0
56	MG	1A	3811	1/1	0.98	0.11	28,28,28,28	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1V	201	1/1	0.98	0.18	39,39,39,39	0
56	MG	1A	3893	1/1	0.98	0.10	51,51,51,51	0
56	MG	2A	3232	1/1	0.98	0.06	47,47,47,47	0
56	MG	1a	1793	1/1	0.98	0.07	83,83,83,83	0
56	MG	1A	3894	1/1	0.98	0.10	50,50,50,50	0
56	MG	1A	3163	1/1	0.98	0.04	32,32,32,32	0
56	MG	1W	3003	1/1	0.98	0.15	33,33,33,33	0
56	MG	1a	1677	1/1	0.98	0.10	48,48,48,48	0
56	MG	1A	3660	1/1	0.98	0.04	24,24,24,24	0
56	MG	2A	3397	1/1	0.98	0.04	58,58,58,58	0
56	MG	1A	3993	1/1	0.98	0.04	34,34,34,34	0
56	MG	1A	3994	1/1	0.98	0.07	23,23,23,23	0
56	MG	1A	3898	1/1	0.98	0.06	52,52,52,52	0
56	MG	1Y	502	1/1	0.98	0.08	40,40,40,40	0
56	MG	1A	3398	1/1	0.98	0.05	46,46,46,46	0
56	MG	1a	1804	1/1	0.98	0.15	37,37,37,37	0
56	MG	1A	3817	1/1	0.98	0.07	40,40,40,40	0
56	MG	2B	207	1/1	0.98	0.07	67,67,67,67	0
56	MG	1A	3730	1/1	0.98	0.04	34,34,34,34	0
56	MG	10	102	1/1	0.98	0.04	48,48,48,48	0
56	MG	1a	1808	1/1	0.98	0.08	33,33,33,33	0
56	MG	2A	3250	1/1	0.98	0.16	30,30,30,30	0
56	MG	1A	3118	1/1	0.98	0.05	47,47,47,47	0
56	MG	2D	303	1/1	0.98	0.24	45,45,45,45	0
56	MG	1A	3212	1/1	0.98	0.18	36,36,36,36	0
56	MG	2D	305	1/1	0.98	0.09	29,29,29,29	0
56	MG	1a	1689	1/1	0.98	0.08	48,48,48,48	0
56	MG	2A	3254	1/1	0.98	0.06	30,30,30,30	0
56	MG	1A	3904	1/1	0.98	0.07	17,17,17,17	0
56	MG	2A	3414	1/1	0.98	0.20	50,50,50,50	0
56	MG	1A	3821	1/1	0.98	0.06	21,21,21,21	0
56	MG	2F	303	1/1	0.98	0.09	52,52,52,52	0
56	MG	1A	3906	1/1	0.98	0.07	53,53,53,53	0
56	MG	1B	219	1/1	0.98	0.08	62,62,62,62	0
56	MG	1B	220	1/1	0.98	0.04	42,42,42,42	0
56	MG	1A	3617	1/1	0.98	0.12	13,13,13,13	0
56	MG	1A	3130	1/1	0.98	0.11	21,21,21,21	0
56	MG	1A	3240	1/1	0.98	0.07	32,32,32,32	0
56	MG	1B	224	1/1	0.98	0.06	58,58,58,58	0
56	MG	1A	3572	1/1	0.98	0.06	52,52,52,52	0
56	MG	2A	3424	1/1	0.98	0.07	63,63,63,63	0
56	MG	1A	4009	1/1	0.98	0.07	47,47,47,47	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3673	1/1	0.98	0.04	25,25,25,25	0
56	MG	1A	3674	1/1	0.98	0.05	24,24,24,24	0
56	MG	1A	3032	1/1	0.98	0.16	44,44,44,44	0
56	MG	2R	202	1/1	0.98	0.05	35,35,35,35	0
56	MG	1A	3132	1/1	0.98	0.08	33,33,33,33	0
56	MG	2U	201	1/1	0.98	0.11	49,49,49,49	0
56	MG	1A	3746	1/1	0.98	0.05	46,46,46,46	0
56	MG	1A	3677	1/1	0.98	0.05	39,39,39,39	0
56	MG	2A	3008	1/1	0.98	0.05	51,51,51,51	0
56	MG	1D	303	1/1	0.98	0.11	38,38,38,38	0
56	MG	1a	1709	1/1	0.98	0.14	61,61,61,61	0
56	MG	1A	3748	1/1	0.98	0.05	13,13,13,13	0
56	MG	1A	3058	1/1	0.98	0.10	45,45,45,45	0
56	MG	2A	3139	1/1	0.98	0.14	35,35,35,35	0
56	MG	1a	1712	1/1	0.98	0.07	40,40,40,40	0
56	MG	1a	1835	1/1	0.98	0.06	51,51,51,51	0
56	MG	1A	3919	1/1	0.98	0.06	51,51,51,51	0
56	MG	1A	3043	1/1	0.98	0.07	55,55,55,55	0
56	MG	1A	3170	1/1	0.98	0.07	25,25,25,25	0
56	MG	2A	3285	1/1	0.98	0.07	33,33,33,33	0
56	MG	2A	3286	1/1	0.98	0.13	47,47,47,47	0
56	MG	2A	3145	1/1	0.98	0.09	25,25,25,25	0
56	MG	1A	3626	1/1	0.98	0.10	45,45,45,45	0
56	MG	2A	3447	1/1	0.98	0.11	43,43,43,43	0
56	MG	1a	1840	1/1	0.98	0.10	27,27,27,27	0
56	MG	1A	3683	1/1	0.98	0.04	42,42,42,42	0
56	MG	2a	3184	1/1	0.98	0.17	53,53,53,53	0
56	MG	2A	3291	1/1	0.98	0.06	37,37,37,37	0
56	MG	2A	3451	1/1	0.98	0.07	59,59,59,59	0
56	MG	1A	3069	1/1	0.98	0.07	39,39,39,39	0
56	MG	1A	3755	1/1	0.98	0.03	26,26,26,26	0
56	MG	1A	3628	1/1	0.98	0.12	43,43,43,43	0
56	MG	2a	3190	1/1	0.98	0.04	47,47,47,47	0
56	MG	1A	3842	1/1	0.98	0.06	54,54,54,54	0
56	MG	1A	3843	1/1	0.98	0.12	53,53,53,53	0
56	MG	1A	3028	1/1	0.98	0.07	50,50,50,50	0
56	MG	2A	3458	1/1	0.98	0.05	54,54,54,54	0
56	MG	1A	3249	1/1	0.98	0.04	20,20,20,20	0
56	MG	1E	305	1/1	0.98	0.08	14,14,14,14	0
56	MG	1A	3250	1/1	0.98	0.08	22,22,22,22	0
56	MG	1A	3763	1/1	0.98	0.06	42,42,42,42	0
56	MG	2A	3464	1/1	0.98	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
56	MG	1A	3251	1/1	0.98	0.09	31,31,31,31	0
56	MG	2a	3201	1/1	0.98	0.04	66,66,66,66	0
56	MG	1A	3934	1/1	0.98	0.06	35,35,35,35	0
56	MG	1A	3692	1/1	0.98	0.05	18,18,18,18	0
56	MG	1A	3693	1/1	0.98	0.05	25,25,25,25	0
56	MG	1A	3694	1/1	0.98	0.05	28,28,28,28	0
56	MG	2A	3036	1/1	0.98	0.12	30,30,30,30	0
56	MG	1A	3695	1/1	0.98	0.06	13,13,13,13	0
56	MG	1a	1735	1/1	0.98	0.07	45,45,45,45	0
56	MG	1A	3854	1/1	0.98	0.10	42,42,42,42	0
56	MG	1A	3696	1/1	0.98	0.12	32,32,32,32	0
56	MG	1a	1863	1/1	0.98	0.12	42,42,42,42	0
56	MG	1A	3770	1/1	0.98	0.05	50,50,50,50	0
56	MG	2A	3315	1/1	0.98	0.07	39,39,39,39	0
56	MG	1A	3221	1/1	0.98	0.18	28,28,28,28	0
56	MG	2A	3173	1/1	0.98	0.11	27,27,27,27	0
56	MG	1A	4042	1/1	0.98	0.13	44,44,44,44	0
56	MG	1A	4043	1/1	0.98	0.09	26,26,26,26	0
56	MG	1A	3943	1/1	0.98	0.11	44,44,44,44	0
56	MG	2a	3220	1/1	0.98	0.11	85,85,85,85	0
56	MG	1A	3040	1/1	0.98	0.08	36,36,36,36	0
56	MG	1A	3062	1/1	0.98	0.05	56,56,56,56	0
56	MG	1A	3700	1/1	0.98	0.06	31,31,31,31	0
56	MG	1A	3176	1/1	0.98	0.07	30,30,30,30	0
56	MG	1A	3226	1/1	0.98	0.09	23,23,23,23	0
56	MG	2A	3326	1/1	0.98	0.09	43,43,43,43	0
56	MG	1A	3703	1/1	0.98	0.07	18,18,18,18	0
56	MG	1A	3177	1/1	0.98	0.04	33,33,33,33	0
56	MG	2A	3492	1/1	0.98	0.12	67,67,67,67	0
56	MG	1a	1631	1/1	0.98	0.11	44,44,44,44	0
56	MG	1A	3951	1/1	0.98	0.08	36,36,36,36	0
56	MG	1A	3082	1/1	0.98	0.04	33,33,33,33	0
56	MG	1A	3158	1/1	0.98	0.15	32,32,32,32	0
56	MG	2A	3337	1/1	0.98	0.06	59,59,59,59	0
56	MG	1A	3782	1/1	0.98	0.05	19,19,19,19	0
56	MG	1A	3955	1/1	0.98	0.05	21,21,21,21	0
56	MG	1A	3180	1/1	0.98	0.06	34,34,34,34	0
56	MG	2a	3238	1/1	0.98	0.05	57,57,57,57	0
56	MG	1A	3958	1/1	0.98	0.09	40,40,40,40	0
56	MG	1A	3784	1/1	0.98	0.10	20,20,20,20	0
56	MG	1A	3785	1/1	0.98	0.06	26,26,26,26	0
56	MG	1A	3292	1/1	0.98	0.10	62,62,62,62	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3346	1/1	0.98	0.04	69,69,69,69	0
56	MG	1A	3181	1/1	0.98	0.16	38,38,38,38	0
56	MG	1A	3967	1/1	0.98	0.11	57,57,57,57	0
56	MG	1A	3159	1/1	0.98	0.13	36,36,36,36	0
56	MG	1A	3598	1/1	0.98	0.08	30,30,30,30	0
56	MG	1A	3359	1/1	0.98	0.07	37,37,37,37	0
56	MG	2a	3249	1/1	0.98	0.07	36,36,36,36	0
56	MG	1A	3600	1/1	0.98	0.06	56,56,56,56	0
56	MG	2A	3354	1/1	0.98	0.04	48,48,48,48	0
56	MG	2A	3355	1/1	0.98	0.10	45,45,45,45	0
56	MG	2A	3201	1/1	0.98	0.13	47,47,47,47	0
56	MG	2A	3202	1/1	0.98	0.07	38,38,38,38	0
56	MG	1A	3264	1/1	0.98	0.18	27,27,27,27	0
56	MG	1A	3034	1/1	0.98	0.12	49,49,49,49	0
56	MG	2A	3520	1/1	0.98	0.07	37,37,37,37	0
56	MG	1Q	203	1/1	0.98	0.10	41,41,41,41	0
56	MG	1A	3144	1/1	0.98	0.09	48,48,48,48	0
56	MG	2A	3523	1/1	0.98	0.13	34,34,34,34	0
56	MG	1A	3064	1/1	0.98	0.07	48,48,48,48	0
56	MG	2A	3209	1/1	0.98	0.28	54,54,54,54	0
56	MG	2a	3084	1/1	0.98	0.10	50,50,50,50	0
56	MG	1A	3652	1/1	0.98	0.07	24,24,24,24	0
56	MG	2l	3005	1/1	0.98	0.09	44,44,44,44	0
56	MG	1A	3801	1/1	0.98	0.06	40,40,40,40	0
56	MG	1R	201	1/1	0.98	0.03	32,32,32,32	0
56	MG	1A	3653	1/1	0.98	0.12	48,48,48,48	0
56	MG	1A	3803	1/1	0.98	0.06	22,22,22,22	0
56	MG	1A	4079	1/1	0.98	0.11	33,33,33,33	0
56	MG	1A	3608	1/1	0.98	0.10	26,26,26,26	0
56	MG	2A	3533	1/1	0.98	0.10	47,47,47,47	0
56	MG	1A	4081	1/1	0.98	0.12	42,42,42,42	0
56	MG	1A	4082	1/1	0.98	0.06	66,66,66,66	0
56	MG	2A	3536	1/1	0.98	0.10	42,42,42,42	0
56	MG	1A	3981	1/1	0.98	0.11	42,42,42,42	0
56	MG	1A	3609	1/1	0.98	0.09	38,38,38,38	0
56	MG	1A	3657	1/1	0.98	0.04	34,34,34,34	0
56	MG	2A	3222	1/1	0.98	0.08	62,62,62,62	0
57	ZN	14	501	1/1	0.98	0.04	103,103,103,103	0
57	ZN	1n	101	1/1	0.98	0.04	83,83,83,83	0
57	ZN	2Y	501	1/1	0.98	0.04	99,99,99,99	0
56	MG	1A	3807	1/1	0.98	0.05	46,46,46,46	0
57	ZN	26	501	1/1	0.98	0.03	61,61,61,61	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3531	1/1	0.98	0.06	45,45,45,45	0
56	MG	1U	201	1/1	0.98	0.04	26,26,26,26	0
58	SF4	1d	501	8/8	0.98	0.06	64,72,77,86	0
56	MG	1A	3715	1/1	0.99	0.04	9,9,9,9	0
56	MG	1A	4067	1/1	0.99	0.04	21,21,21,21	0
56	MG	2a	3107	1/1	0.99	0.03	57,57,57,57	0
56	MG	1A	3185	1/1	0.99	0.08	38,38,38,38	0
56	MG	1A	4069	1/1	0.99	0.04	47,47,47,47	0
56	MG	1A	3482	1/1	0.99	0.05	12,12,12,12	0
56	MG	1A	3096	1/1	0.99	0.16	40,40,40,40	0
56	MG	1A	3762	1/1	0.99	0.03	39,39,39,39	0
56	MG	1A	3960	1/1	0.99	0.03	28,28,28,28	0
56	MG	1A	3961	1/1	0.99	0.03	40,40,40,40	0
56	MG	2A	3375	1/1	0.99	0.09	46,46,46,46	0
56	MG	2A	3459	1/1	0.99	0.04	29,29,29,29	0
56	MG	1A	3685	1/1	0.99	0.05	16,16,16,16	0
56	MG	1A	3328	1/1	0.99	0.04	27,27,27,27	0
56	MG	2A	3004	1/1	0.99	0.05	59,59,59,59	0
56	MG	1A	3088	1/1	0.99	0.05	53,53,53,53	0
56	MG	1U	203	1/1	0.99	0.08	29,29,29,29	0
56	MG	1A	3965	1/1	0.99	0.06	36,36,36,36	0
56	MG	1a	1817	1/1	0.99	0.04	54,54,54,54	0
56	MG	2a	3215	1/1	0.99	0.08	53,53,53,53	0
56	MG	1A	3966	1/1	0.99	0.05	19,19,19,19	0
56	MG	2A	3300	1/1	0.99	0.03	38,38,38,38	0
56	MG	1A	3688	1/1	0.99	0.06	8,8,8,8	0
56	MG	2A	3152	1/1	0.99	0.14	30,30,30,30	0
56	MG	1A	3723	1/1	0.99	0.03	29,29,29,29	0
56	MG	1A	3655	1/1	0.99	0.05	24,24,24,24	0
56	MG	1A	3815	1/1	0.99	0.04	20,20,20,20	0
56	MG	1A	3816	1/1	0.99	0.04	51,51,51,51	0
56	MG	1A	3725	1/1	0.99	0.03	18,18,18,18	0
56	MG	1A	3604	1/1	0.99	0.06	36,36,36,36	0
56	MG	1A	3113	1/1	0.99	0.05	35,35,35,35	0
56	MG	1A	3606	1/1	0.99	0.05	28,28,28,28	0
56	MG	1A	3246	1/1	0.99	0.04	23,23,23,23	0
56	MG	2A	3480	1/1	0.99	0.05	21,21,21,21	0
56	MG	1A	3822	1/1	0.99	0.06	21,21,21,21	0
56	MG	1A	3134	1/1	0.99	0.07	23,23,23,23	0
56	MG	1A	3155	1/1	0.99	0.15	39,39,39,39	0
56	MG	1A	3732	1/1	0.99	0.05	26,26,26,26	0
56	MG	1B	204	1/1	0.99	0.04	49,49,49,49	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3662	1/1	0.99	0.06	30,30,30,30	0
56	MG	1A	3663	1/1	0.99	0.02	19,19,19,19	0
56	MG	1A	3664	1/1	0.99	0.07	19,19,19,19	0
56	MG	1A	3737	1/1	0.99	0.03	18,18,18,18	0
56	MG	1a	1705	1/1	0.99	0.09	58,58,58,58	0
56	MG	2E	302	1/1	0.99	0.17	36,36,36,36	0
56	MG	1A	3781	1/1	0.99	0.04	24,24,24,24	0
56	MG	1A	3365	1/1	0.99	0.04	30,30,30,30	0
56	MG	1A	3612	1/1	0.99	0.07	23,23,23,23	0
56	MG	2A	3248	1/1	0.99	0.06	35,35,35,35	0
56	MG	1A	3191	1/1	0.99	0.12	28,28,28,28	0
56	MG	2A	3327	1/1	0.99	0.06	44,44,44,44	0
56	MG	1A	3401	1/1	0.99	0.09	30,30,30,30	0
56	MG	1A	3743	1/1	0.99	0.04	30,30,30,30	0
56	MG	2A	3330	1/1	0.99	0.06	40,40,40,40	0
56	MG	1A	3615	1/1	0.99	0.04	39,39,39,39	0
56	MG	2A	3333	1/1	0.99	0.04	35,35,35,35	0
56	MG	2A	3502	1/1	0.99	0.03	37,37,37,37	0
56	MG	1a	1846	1/1	0.99	0.03	25,25,25,25	0
56	MG	1A	3671	1/1	0.99	0.08	27,27,27,27	0
56	MG	1B	217	1/1	0.99	0.03	38,38,38,38	0
56	MG	1A	3225	1/1	0.99	0.03	21,21,21,21	0
56	MG	2A	3507	1/1	0.99	0.03	38,38,38,38	0
56	MG	1A	3135	1/1	0.99	0.16	30,30,30,30	0
56	MG	1A	3592	1/1	0.99	0.07	25,25,25,25	0
56	MG	2A	3113	1/1	0.99	0.12	29,29,29,29	0
56	MG	2A	3341	1/1	0.99	0.06	51,51,51,51	0
56	MG	1A	4051	1/1	0.99	0.13	15,15,15,15	0
56	MG	1A	3173	1/1	0.99	0.04	40,40,40,40	0
56	MG	2a	3172	1/1	0.99	0.05	42,42,42,42	0
56	MG	1A	3353	1/1	0.99	0.06	40,40,40,40	0
56	MG	1A	3573	1/1	0.99	0.10	41,41,41,41	0
56	MG	2A	3264	1/1	0.99	0.06	29,29,29,29	0
56	MG	2A	3347	1/1	0.99	0.05	56,56,56,56	0
56	MG	1A	3795	1/1	0.99	0.08	13,13,13,13	0
56	MG	1a	1723	1/1	0.99	0.07	62,62,62,62	0
56	MG	1A	3796	1/1	0.99	0.09	24,24,24,24	0
56	MG	1A	3896	1/1	0.99	0.04	72,72,72,72	0
56	MG	1A	3797	1/1	0.99	0.03	28,28,28,28	0
56	MG	1a	1861	1/1	0.99	0.07	51,51,51,51	0
56	MG	1A	3798	1/1	0.99	0.04	19,19,19,19	0
56	MG	1A	3142	1/1	0.99	0.03	20,20,20,20	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3273	1/1	0.99	0.06	53,53,53,53	0
56	MG	19	105	1/1	0.99	0.13	34,34,34,34	0
56	MG	1Q	205	1/1	0.99	0.03	40,40,40,40	0
57	ZN	1Y	501	1/1	0.99	0.02	62,62,62,62	0
56	MG	1A	3254	1/1	0.99	0.08	41,41,41,41	0
57	ZN	15	101	1/1	0.99	0.07	54,54,54,54	0
57	ZN	16	103	1/1	0.99	0.07	45,45,45,45	0
56	MG	1A	3065	1/1	0.99	0.03	39,39,39,39	0
56	MG	2a	3099	1/1	0.99	0.18	41,41,41,41	0
56	MG	1Q	208	1/1	0.99	0.02	37,37,37,37	0
56	MG	2A	3203	1/1	0.99	0.08	43,43,43,43	0
56	MG	1A	3681	1/1	0.99	0.04	19,19,19,19	0
56	MG	1A	4008	1/1	0.99	0.03	20,20,20,20	0
56	MG	1A	3756	1/1	0.99	0.06	30,30,30,30	0
58	SF4	2d	501	8/8	0.99	0.04	58,72,80,87	0
56	MG	1A	3844	1/1	1.00	0.03	18,18,18,18	0
57	ZN	19	103	1/1	1.00	0.05	47,47,47,47	0
56	MG	1A	3760	1/1	1.00	0.02	34,34,34,34	0
56	MG	1A	3586	1/1	1.00	0.08	33,33,33,33	0
56	MG	1A	3736	1/1	1.00	0.06	15,15,15,15	0
57	ZN	25	501	1/1	1.00	0.03	50,50,50,50	0
56	MG	1A	3607	1/1	1.00	0.07	26,26,26,26	0
56	MG	1A	3738	1/1	1.00	0.05	17,17,17,17	0
56	MG	1A	3668	1/1	1.00	0.04	17,17,17,17	0
56	MG	1A	3956	1/1	1.00	0.06	12,12,12,12	0
56	MG	2A	3332	1/1	1.00	0.02	42,42,42,42	0

5.5 Other polymers ⓘ

There are no such residues in this entry.