



Full wwPDB X-ray Structure Validation Report ⓘ

Mar 30, 2026 – 12:09 PM UTC

PDB ID : 4Z8C / pdb_00004z8c
Title : Crystal structure of the Thermus thermophilus 70S ribosome bound to translation inhibitor oncocin
Authors : Roy, R.N.; Lomakin, I.B.; Gagnon, M.G.; Steitz, T.A.
Deposited on : 2015-04-08
Resolution : 2.90 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

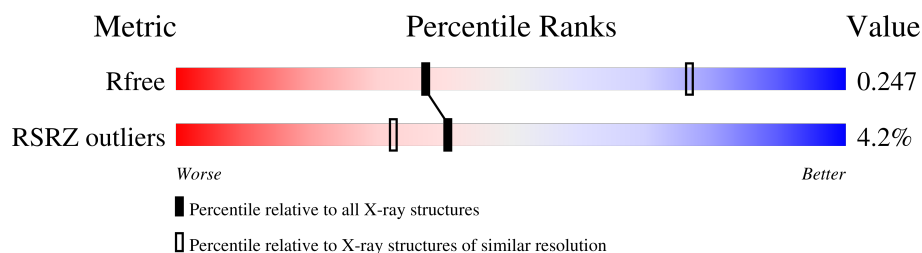
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

X-RAY DIFFRACTION

The reported resolution of this entry is 2.90 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.

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

Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
56	MG	2a	3024	-	-	-	X
56	MG	2a	3138	-	-	-	X
56	MG	2a	3139	-	-	-	X

2 Entry composition

There are 59 unique types of molecules in this entry. The entry contains 288378 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 Initiator Methionine tRNA.

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

- Molecule 55 is a protein called Oncocin.

Mol	Chain	Residues	Atoms				ZeroOcc	AltConf	Trace
55	1z	13	Total	C	N	O	0	0	0
			108	72	20	16			
55	2z	13	Total	C	N	O	0	0	0
			108	72	20	16			

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

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	1A	1021	Total	Mg	0	0
			1021	1021		
56	1B	24	Total	Mg	0	0
			24	24		
56	1D	14	Total	Mg	0	0
			14	14		
56	1E	5	Total	Mg	0	0
			5	5		
56	1F	8	Total	Mg	0	0
			8	8		
56	1G	3	Total	Mg	0	0
			3	3		
56	1H	2	Total	Mg	0	0
			2	2		
56	1N	6	Total	Mg	0	0
			6	6		
56	1O	2	Total	Mg	0	0
			2	2		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	1P	6	Total 6	Mg 6	0	0
56	1Q	4	Total 4	Mg 4	0	0
56	1R	4	Total 4	Mg 4	0	0
56	1T	5	Total 5	Mg 5	0	0
56	1U	5	Total 5	Mg 5	0	0
56	1V	1	Total 1	Mg 1	0	0
56	1W	5	Total 5	Mg 5	0	0
56	1X	2	Total 2	Mg 2	0	0
56	1Z	2	Total 2	Mg 2	0	0
56	10	7	Total 7	Mg 7	0	0
56	11	1	Total 1	Mg 1	0	0
56	12	1	Total 1	Mg 1	0	0
56	13	2	Total 2	Mg 2	0	0
56	15	4	Total 4	Mg 4	0	0
56	16	2	Total 2	Mg 2	0	0
56	17	1	Total 1	Mg 1	0	0
56	18	3	Total 3	Mg 3	0	0
56	19	3	Total 3	Mg 3	0	0
56	1a	276	Total 276	Mg 276	0	0
56	1b	2	Total 2	Mg 2	0	0
56	1d	2	Total 2	Mg 2	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	1e	3	Total 3	Mg 3	0	0
56	1f	1	Total 1	Mg 1	0	0
56	1k	1	Total 1	Mg 1	0	0
56	1l	1	Total 1	Mg 1	0	0
56	1n	1	Total 1	Mg 1	0	0
56	1o	2	Total 2	Mg 2	0	0
56	1q	4	Total 4	Mg 4	0	0
56	1r	2	Total 2	Mg 2	0	0
56	1t	1	Total 1	Mg 1	0	0
56	1x	8	Total 8	Mg 8	0	0
56	1z	1	Total 1	Mg 1	0	0
56	2A	566	Total 566	Mg 566	0	0
56	2B	11	Total 11	Mg 11	0	0
56	2D	6	Total 6	Mg 6	0	0
56	2E	4	Total 4	Mg 4	0	0
56	2F	6	Total 6	Mg 6	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

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Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
56	2U	1	Total Mg 1 1	0	0
56	2Y	1	Total Mg 1 1	0	0
56	2Z	1	Total Mg 1 1	0	0
56	20	1	Total Mg 1 1	0	0
56	25	1	Total Mg 1 1	0	0
56	26	1	Total Mg 1 1	0	0
56	28	1	Total Mg 1 1	0	0
56	2a	263	Total Mg 263 263	0	0
56	2d	1	Total Mg 1 1	0	0
56	2f	1	Total Mg 1 1	0	0
56	2h	1	Total Mg 1 1	0	0
56	2k	1	Total Mg 1 1	0	0
56	2l	4	Total Mg 4 4	0	0
56	2q	2	Total Mg 2 2	0	0
56	2t	1	Total Mg 1 1	0	0
56	2x	4	Total Mg 4 4	0	0
56	2z	1	Total Mg 1 1	0	0

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

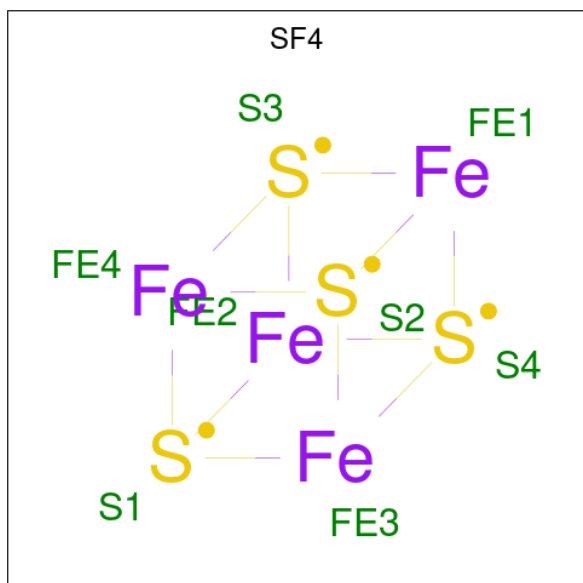
Mol	Chain	Residues	Atoms	ZeroOcc	AltConf
57	1Y	1	Total Zn 1 1	0	0
57	14	1	Total Zn 1 1	0	0

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

- Molecule 58 is IRON/SULFUR CLUSTER (CCD ID: SF4) (formula: Fe_4S_4).



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

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

- Molecule 59 is water.

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	1A	2088	Total	O	0	0
			2088	2088		
59	1B	38	Total	O	0	0
			38	38		
59	1D	22	Total	O	0	0
			22	22		
59	1E	25	Total	O	0	0
			25	25		
59	1F	16	Total	O	0	0
			16	16		
59	1G	6	Total	O	0	0
			6	6		
59	1H	5	Total	O	0	0
			5	5		
59	1I	1	Total	O	0	0
			1	1		
59	1N	5	Total	O	0	0
			5	5		
59	1O	4	Total	O	0	0
			4	4		
59	1P	21	Total	O	0	0
			21	21		
59	1Q	9	Total	O	0	0
			9	9		
59	1R	9	Total	O	0	0
			9	9		
59	1S	1	Total	O	0	0
			1	1		
59	1T	11	Total	O	0	0
			11	11		
59	1U	13	Total	O	0	0
			13	13		
59	1V	2	Total	O	0	0
			2	2		
59	1W	9	Total	O	0	0
			9	9		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	1X	5	Total 5	O 5	0	0
59	1Y	1	Total 1	O 1	0	0
59	1Z	3	Total 3	O 3	0	0
59	10	10	Total 10	O 10	0	0
59	11	4	Total 4	O 4	0	0
59	12	1	Total 1	O 1	0	0
59	13	4	Total 4	O 4	0	0
59	15	5	Total 5	O 5	0	0
59	16	8	Total 8	O 8	0	0
59	17	8	Total 8	O 8	0	0
59	18	13	Total 13	O 13	0	0
59	19	2	Total 2	O 2	0	0
59	1a	319	Total 319	O 319	0	0
59	1b	2	Total 2	O 2	0	0
59	1c	1	Total 1	O 1	0	0
59	1d	4	Total 4	O 4	0	0
59	1e	2	Total 2	O 2	0	0
59	1f	1	Total 1	O 1	0	0
59	1i	1	Total 1	O 1	0	0
59	1j	1	Total 1	O 1	0	0
59	1m	1	Total 1	O 1	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	1o	3	Total 3	O 3	0	0
59	1p	1	Total 1	O 1	0	0
59	1s	1	Total 1	O 1	0	0
59	1v	1	Total 1	O 1	0	0
59	1x	7	Total 7	O 7	0	0
59	1z	2	Total 2	O 2	0	0
59	2A	801	Total 801	O 801	0	0
59	2B	13	Total 13	O 13	0	0
59	2D	20	Total 20	O 20	0	0
59	2E	10	Total 10	O 10	0	0
59	2F	7	Total 7	O 7	0	0
59	2N	1	Total 1	O 1	0	0
59	2O	4	Total 4	O 4	0	0
59	2P	4	Total 4	O 4	0	0
59	2Q	2	Total 2	O 2	0	0
59	2R	3	Total 3	O 3	0	0
59	2T	3	Total 3	O 3	0	0
59	2U	2	Total 2	O 2	0	0
59	2V	2	Total 2	O 2	0	0
59	2W	2	Total 2	O 2	0	0
59	2X	3	Total 3	O 3	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	2Y	1	Total 1	O 1	0	0
59	2Z	2	Total 2	O 2	0	0
59	20	3	Total 3	O 3	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	2	Total 2	O 2	0	0
59	28	4	Total 4	O 4	0	0
59	2a	371	Total 371	O 371	0	0
59	2c	2	Total 2	O 2	0	0
59	2e	3	Total 3	O 3	0	0
59	2h	1	Total 1	O 1	0	0
59	2i	2	Total 2	O 2	0	0
59	2j	1	Total 1	O 1	0	0
59	2k	3	Total 3	O 3	0	0
59	2l	3	Total 3	O 3	0	0
59	2m	1	Total 1	O 1	0	0
59	2n	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	2	Total 2	O 2	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
59	2r	1	Total 1	O 1	0	0
59	2t	3	Total 3	O 3	0	0
59	2x	3	Total 3	O 3	0	0
59	2z	1	Total 1	O 1	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	210.14Å 450.73Å 623.47Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	59.98 – 2.90 59.98 – 2.90	Depositor EDS
% Data completeness (in resolution range)	98.5 (59.98-2.90) 98.5 (59.98-2.90)	Depositor EDS
R_{merge}	0.18	Depositor
R_{sym}	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ ¹	1.12 (at 2.82Å)	Xtriage
Refinement program	PHENIX	Depositor
R, R_{free}	0.203 , 0.248 0.204 , 0.247	Depositor DCC
R_{free} test set	70488 reflections (4.92%)	wwPDB-VP
Wilson B-factor (Å ²)	63.0	Xtriage
Anisotropy	0.092	Xtriage
Bulk solvent k_{sol} (e/Å ³), B_{sol} (Å ²)	0.29 , 54.0	EDS
L-test for twinning ²	$\langle L \rangle = 0.47$, $\langle L^2 \rangle = 0.30$	Xtriage
Estimated twinning fraction	No twinning to report.	Xtriage
F_o, F_c correlation	0.93	EDS
Total number of atoms	288378	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 2.07% 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	PSU	1x	55	54	18,21,22	1.39	3 (16%)	21,30,33	1.90	3 (14%)
54	4SU	2x	8	54	18,21,22	2.00	5 (27%)	25,30,33	1.36	3 (12%)
54	5MC	2x	32	54	19,22,23	1.60	3 (15%)	26,32,35	1.21	3 (11%)

Mol	Type	Chain	Res	Link	Bond lengths			Bond angles		
					Counts	RMSZ	# Z > 2	Counts	RMSZ	# Z > 2
54	5MU	1x	54	54	19,22,23	1.44	6 (31%)	27,32,35	1.88	5 (18%)
54	5MC	1x	32	54	19,22,23	1.70	3 (15%)	26,32,35	1.15	2 (7%)
54	5MU	2x	54	54	19,22,23	1.42	5 (26%)	27,32,35	2.21	6 (22%)
54	PSU	2x	55	54	18,21,22	1.38	2 (11%)	21,30,33	1.92	3 (14%)
54	4SU	1x	8	54	18,21,22	2.07	4 (22%)	25,30,33	1.36	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	PSU	1x	55	54	-	3/7/25/26	0/2/2/2
54	4SU	2x	8	54	-	0/7/25/26	0/2/2/2
54	5MC	2x	32	54	-	0/7/25/26	0/2/2/2
54	5MU	1x	54	54	-	0/7/25/26	0/2/2/2
54	5MC	1x	32	54	-	0/7/25/26	0/2/2/2
54	5MU	2x	54	54	-	0/7/25/26	0/2/2/2
54	PSU	2x	55	54	-	3/7/25/26	0/2/2/2
54	4SU	1x	8	54	-	0/7/25/26	0/2/2/2

All (31) bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
54	1x	32	5MC	C5-C4	6.15	1.48	1.44
54	2x	32	5MC	C5-C4	5.67	1.48	1.44
54	1x	8	4SU	C4-S4	-4.62	1.60	1.68
54	1x	8	4SU	C4-N3	-4.58	1.32	1.37
54	2x	8	4SU	C4-S4	-4.54	1.60	1.68
54	2x	8	4SU	C4-N3	-4.42	1.33	1.37
54	2x	55	PSU	C6-C5	3.62	1.39	1.35
54	1x	8	4SU	C2-N3	-3.59	1.31	1.38
54	1x	55	PSU	C6-C5	3.49	1.39	1.35
54	1x	8	4SU	C5-C4	-3.23	1.38	1.42
54	2x	8	4SU	C5-C4	-3.11	1.38	1.42
54	1x	32	5MC	C6-C5	2.94	1.39	1.34
54	2x	54	5MU	C6-C5	2.91	1.39	1.34
54	1x	54	5MU	C6-C5	2.90	1.39	1.34
54	2x	32	5MC	C6-C5	2.78	1.39	1.34
54	2x	8	4SU	C2-N3	-2.72	1.33	1.38
54	1x	54	5MU	C4-N3	-2.72	1.33	1.38

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Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
54	2x	54	5MU	C4-N3	-2.61	1.34	1.38
54	2x	54	5MU	C4-C5	2.57	1.49	1.44
54	1x	55	PSU	C4-N3	-2.46	1.34	1.38
54	1x	54	5MU	C4-C5	2.38	1.48	1.44
54	2x	55	PSU	C4-N3	-2.32	1.34	1.38
54	1x	32	5MC	C6-N1	-2.24	1.34	1.38
54	1x	54	5MU	C2-N3	-2.21	1.34	1.38
54	1x	54	5MU	C2-N1	2.18	1.41	1.38
54	2x	54	5MU	C6-N1	-2.18	1.34	1.38
54	2x	32	5MC	C6-N1	-2.14	1.34	1.38
54	2x	54	5MU	C2-N1	2.12	1.41	1.38
54	1x	54	5MU	C6-N1	-2.11	1.34	1.38
54	2x	8	4SU	O2-C2	2.09	1.26	1.23
54	1x	55	PSU	O4'-C1'	-2.02	1.41	1.43

All (29) bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
54	2x	55	PSU	N1-C2-N3	5.90	121.40	115.17
54	1x	55	PSU	N1-C2-N3	5.83	121.32	115.17
54	2x	54	5MU	N3-C2-N1	5.57	122.14	114.89
54	2x	54	5MU	C4-N3-C2	-5.46	120.19	127.34
54	1x	54	5MU	N3-C2-N1	4.66	120.95	114.89
54	1x	54	5MU	C4-N3-C2	-4.46	121.49	127.34
54	2x	54	5MU	C5-C4-N3	4.26	119.03	115.32
54	1x	54	5MU	C5-C4-N3	4.01	118.81	115.32
54	2x	54	5MU	O4-C4-C5	-3.89	120.46	124.92
54	2x	55	PSU	C4-N3-C2	-3.81	121.12	126.37
54	1x	55	PSU	C4-N3-C2	-3.78	121.16	126.37
54	2x	54	5MU	C5-C6-N1	-3.70	119.30	123.31
54	2x	55	PSU	O2-C2-N1	-3.60	119.08	122.79
54	1x	32	5MC	C5-C6-N1	-3.58	119.42	123.31
54	2x	8	4SU	C5-C4-N3	3.57	118.08	114.75
54	1x	8	4SU	C5-C4-N3	3.50	118.00	114.75
54	1x	55	PSU	O2-C2-N1	-3.45	119.23	122.79
54	1x	54	5MU	O4-C4-C5	-3.38	121.05	124.92
54	2x	54	5MU	O2-C2-N1	-3.27	118.53	122.80
54	1x	54	5MU	C5-C6-N1	-3.25	119.78	123.31
54	1x	8	4SU	C6-C5-C4	-3.14	117.23	119.95
54	2x	32	5MC	C5-C6-N1	-2.99	120.06	123.31
54	2x	8	4SU	C1'-N1-C2	2.97	122.92	117.59
54	1x	32	5MC	C5-C4-N3	-2.86	118.82	121.75

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Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
54	2x	32	5MC	O2-C2-N3	-2.71	118.05	122.33
54	2x	32	5MC	C5-C4-N3	-2.65	119.03	121.75
54	2x	8	4SU	C6-C5-C4	-2.64	117.66	119.95
54	1x	8	4SU	C1'-N1-C2	2.41	121.93	117.59
54	1x	8	4SU	O2-C2-N1	2.36	125.86	122.80

There are no chirality outliers.

All (6) torsion outliers are listed below:

Mol	Chain	Res	Type	Atoms
54	1x	55	PSU	C3'-C4'-C5'-O5'
54	2x	55	PSU	C3'-C4'-C5'-O5'
54	1x	55	PSU	O4'-C4'-C5'-O5'
54	2x	55	PSU	O4'-C4'-C5'-O5'
54	1x	55	PSU	C4'-C5'-O5'-P
54	2x	55	PSU	C4'-C5'-O5'-P

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 2352 ligands modelled in this entry, 2350 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	2d	501	35	0,12,12	-	-	-		
58	SF4	1d	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	2d	501	35	-	-	0/6/5/5
58	SF4	1d	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%)	-0.42	39 (1%) 73 65	24, 41, 87, 115	0
1	2A	2790/2915 (95%)	-0.34	33 (1%) 76 69	26, 45, 94, 113	0
2	1B	120/121 (99%)	0.12	0 100 100	40, 66, 79, 96	0
2	2B	120/121 (99%)	0.60	3 (2%) 58 48	45, 72, 82, 98	0
3	1D	275/276 (99%)	-0.17	2 (0%) 84 79	21, 40, 58, 81	0
3	2D	275/276 (99%)	-0.16	3 (1%) 78 71	21, 42, 60, 81	0
4	1E	204/206 (99%)	-0.44	0 100 100	17, 37, 64, 79	0
4	2E	204/206 (99%)	0.02	1 (0%) 87 83	26, 52, 72, 83	0
5	1F	203/210 (96%)	0.04	0 100 100	22, 50, 77, 92	0
5	2F	203/210 (96%)	0.13	0 100 100	25, 54, 78, 92	0
6	1G	181/182 (99%)	0.74	13 (7%) 21 17	54, 71, 84, 98	0
6	2G	181/182 (99%)	1.21	28 (15%) 5 4	59, 76, 88, 99	0
7	1H	174/180 (96%)	0.67	8 (4%) 37 29	49, 64, 79, 86	0
7	2H	174/180 (96%)	0.95	9 (5%) 33 26	52, 70, 82, 88	0
8	1I	146/148 (98%)	0.61	4 (2%) 56 47	47, 75, 87, 93	0
8	2I	146/148 (98%)	0.82	12 (8%) 17 14	48, 77, 87, 93	0
9	1N	140/140 (100%)	0.11	0 100 100	30, 46, 70, 78	0
9	2N	140/140 (100%)	0.06	2 (1%) 73 65	33, 50, 72, 80	0
10	1O	122/122 (100%)	-0.13	2 (1%) 70 62	31, 42, 64, 70	0
10	2O	122/122 (100%)	-0.11	0 100 100	34, 45, 64, 71	0
11	1P	149/150 (99%)	0.17	1 (0%) 84 79	25, 53, 75, 85	0
11	2P	149/150 (99%)	0.16	0 100 100	29, 57, 77, 87	0
12	1Q	141/141 (100%)	0.25	5 (3%) 47 38	32, 50, 65, 84	0
12	2Q	141/141 (100%)	0.30	5 (3%) 47 38	36, 53, 68, 84	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
13	1R	118/118 (100%)	-0.11	0 100 100	27, 40, 56, 67	0
13	2R	118/118 (100%)	-0.12	0 100 100	29, 43, 57, 69	0
14	1S	110/112 (98%)	0.40	4 (3%) 46 38	49, 63, 74, 82	0
14	2S	110/112 (98%)	0.85	8 (7%) 21 17	54, 67, 78, 84	0
15	1T	131/146 (89%)	0.09	4 (3%) 51 42	35, 48, 80, 88	0
15	2T	131/146 (89%)	0.05	2 (1%) 72 64	37, 50, 78, 87	0
16	1U	116/118 (98%)	0.05	2 (1%) 69 61	29, 39, 57, 78	0
16	2U	116/118 (98%)	-0.05	0 100 100	31, 44, 61, 78	0
17	1V	101/101 (100%)	0.01	0 100 100	30, 51, 69, 78	0
17	2V	101/101 (100%)	0.17	0 100 100	33, 57, 72, 79	0
18	1W	112/113 (99%)	-0.16	0 100 100	28, 37, 59, 90	0
18	2W	112/113 (99%)	-0.23	0 100 100	30, 40, 60, 90	0
19	1X	95/96 (98%)	0.05	2 (2%) 63 54	31, 47, 68, 80	0
19	2X	95/96 (98%)	0.21	2 (2%) 63 54	35, 50, 70, 81	0
20	1Y	107/110 (97%)	0.09	5 (4%) 36 28	31, 49, 74, 84	0
20	2Y	107/110 (97%)	0.85	9 (8%) 17 14	55, 72, 84, 96	0
21	1Z	186/206 (90%)	0.06	0 100 100	40, 62, 79, 96	0
21	2Z	186/206 (90%)	0.80	10 (5%) 31 24	67, 81, 94, 101	0
22	10	75/85 (88%)	-0.20	1 (1%) 75 67	23, 38, 53, 67	0
22	20	75/85 (88%)	0.51	4 (5%) 32 25	43, 63, 73, 78	0
23	11	97/98 (98%)	-0.11	1 (1%) 79 73	21, 42, 71, 80	0
23	21	97/98 (98%)	0.19	3 (3%) 51 42	33, 54, 77, 86	0
24	12	70/72 (97%)	0.52	6 (8%) 16 13	42, 59, 74, 89	0
24	22	70/72 (97%)	0.40	1 (1%) 73 65	46, 63, 75, 87	0
25	13	59/60 (98%)	0.17	3 (5%) 33 26	32, 46, 67, 80	0
25	23	59/60 (98%)	0.13	2 (3%) 48 39	36, 49, 72, 82	0
26	14	69/71 (97%)	0.90	11 (15%) 5 4	68, 84, 100, 103	0
26	24	69/71 (97%)	1.51	16 (23%) 2 2	73, 86, 100, 104	0
27	15	59/60 (98%)	-0.29	0 100 100	21, 40, 60, 78	0
27	25	59/60 (98%)	-0.17	0 100 100	24, 43, 62, 79	0
28	16	53/54 (98%)	-0.15	1 (1%) 66 58	36, 48, 63, 68	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	39, 51, 65, 70	0
29	17	48/49 (97%)	-0.12	3 (6%) 26 20	26, 31, 67, 76	0
29	27	48/49 (97%)	-0.03	2 (4%) 40 32	28, 33, 67, 78	0
30	18	64/65 (98%)	-0.03	0 100 100	32, 40, 51, 64	0
30	28	64/65 (98%)	0.12	0 100 100	34, 43, 54, 67	0
31	19	37/37 (100%)	0.36	2 (5%) 31 24	40, 49, 66, 69	0
31	29	37/37 (100%)	0.66	2 (5%) 31 24	44, 53, 67, 71	0
32	1a	1477/1521 (97%)	0.17	23 (1%) 70 62	33, 72, 99, 116	0
32	2a	1483/1521 (97%)	0.31	29 (1%) 65 56	43, 78, 102, 119	0
33	1b	231/256 (90%)	0.99	26 (11%) 10 9	60, 83, 95, 105	0
33	2b	231/256 (90%)	1.27	43 (18%) 3 3	71, 88, 97, 101	0
34	1c	206/239 (86%)	0.88	11 (5%) 32 25	70, 84, 91, 99	0
34	2c	206/239 (86%)	1.24	33 (16%) 5 4	72, 86, 93, 100	0
35	1d	208/209 (99%)	1.02	26 (12%) 8 7	55, 78, 91, 101	0
35	2d	208/209 (99%)	0.93	18 (8%) 16 13	58, 75, 87, 90	0
36	1e	148/162 (91%)	0.25	2 (1%) 73 65	47, 65, 77, 94	0
36	2e	148/162 (91%)	0.58	4 (2%) 56 47	56, 74, 86, 91	0
37	1f	100/101 (99%)	0.59	4 (4%) 42 34	57, 75, 85, 89	0
37	2f	100/101 (99%)	0.47	0 100 100	60, 76, 86, 90	0
38	1g	155/156 (99%)	0.66	13 (8%) 17 14	66, 80, 93, 101	0
38	2g	155/156 (99%)	0.89	14 (9%) 15 13	69, 81, 94, 99	0
39	1h	137/138 (99%)	0.34	1 (0%) 84 79	56, 68, 80, 85	0
39	2h	137/138 (99%)	0.73	8 (5%) 29 22	58, 75, 83, 89	0
40	1i	127/128 (99%)	1.45	31 (24%) 2 1	66, 87, 94, 99	0
40	2i	127/128 (99%)	1.77	44 (34%) 1 1	65, 88, 96, 100	0
41	1j	97/105 (92%)	1.56	26 (26%) 1 1	65, 86, 96, 104	0
41	2j	96/105 (91%)	2.08	47 (48%) 0 0	75, 92, 99, 103	0
42	1k	114/129 (88%)	0.42	3 (2%) 57 48	45, 69, 82, 94	0
42	2k	114/129 (88%)	0.63	6 (5%) 32 25	52, 76, 88, 91	0
43	1l	122/132 (92%)	0.42	7 (5%) 29 23	47, 65, 76, 88	0
43	2l	122/132 (92%)	0.42	4 (3%) 49 40	52, 66, 77, 85	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å ²)	Q<0.9
44	1m	118/126 (93%)	0.76	6 (5%) 33 26	64, 82, 89, 95	0
44	2m	116/126 (92%)	1.25	13 (11%) 10 9	66, 84, 91, 94	0
45	1n	60/61 (98%)	1.25	9 (15%) 5 4	69, 81, 88, 93	0
45	2n	60/61 (98%)	1.96	28 (46%) 0 0	71, 83, 89, 95	0
46	1o	88/89 (98%)	0.40	3 (3%) 48 39	49, 68, 84, 91	0
46	2o	88/89 (98%)	0.47	3 (3%) 48 39	56, 73, 85, 94	0
47	1p	82/88 (93%)	1.56	23 (28%) 1 1	66, 79, 92, 96	0
47	2p	82/88 (93%)	0.87	4 (4%) 35 27	59, 71, 83, 96	0
48	1q	99/105 (94%)	0.37	2 (2%) 65 56	51, 68, 81, 84	0
48	2q	99/105 (94%)	0.51	2 (2%) 65 56	57, 72, 83, 86	0
49	1r	68/88 (77%)	0.44	2 (2%) 53 45	56, 71, 84, 91	0
49	2r	68/88 (77%)	0.62	0 100 100	65, 75, 89, 94	0
50	1s	84/93 (90%)	0.80	4 (4%) 35 28	71, 82, 92, 98	0
50	2s	83/93 (89%)	1.54	25 (30%) 1 1	80, 92, 101, 108	0
51	1t	96/106 (90%)	0.83	8 (8%) 17 14	61, 74, 88, 90	0
51	2t	96/106 (90%)	0.59	3 (3%) 51 42	54, 73, 85, 90	0
52	1u	23/27 (85%)	1.41	4 (17%) 4 3	71, 80, 85, 87	0
52	2u	23/27 (85%)	2.37	16 (69%) 0 0	72, 82, 87, 89	0
53	1v	5/24 (20%)	1.01	1 (20%) 3 2	57, 67, 94, 101	0
53	2v	5/24 (20%)	0.92	1 (20%) 3 2	61, 70, 95, 102	0
54	1x	72/77 (93%)	-0.03	0 100 100	35, 68, 87, 102	0
54	2x	72/77 (93%)	0.12	0 100 100	38, 72, 89, 103	0
55	1z	13/19 (68%)	0.20	0 100 100	26, 37, 63, 72	0
55	2z	13/19 (68%)	0.12	1 (7%) 19 16	28, 39, 65, 74	0
All	All	20522/21482 (95%)	0.22	857 (4%) 40 32	17, 62, 93, 119	0

All (857) RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
23	11	2	SER	6.0
38	1g	79	ARG	5.6
1	1A	934	C	5.6
7	1H	2	SER	5.6
29	27	48	LYS	5.6

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Mol	Chain	Res	Type	RSRZ
45	2n	2	ALA	5.6
40	2i	7	THR	5.5
40	1i	106	ALA	5.3
41	2j	64	GLU	5.3
38	1g	80	VAL	5.2
41	2j	63	PHE	5.1
41	2j	76	ASN	5.0
23	2l	2	SER	5.0
40	1i	67	GLY	4.9
40	1i	13	ALA	4.9
32	2a	1510	U	4.9
32	1a	1509	A	4.8
40	1i	64	THR	4.8
45	2n	33	VAL	4.8
40	2i	102	LEU	4.7
6	1G	48	GLU	4.7
6	1G	51	ARG	4.7
50	2s	35	SER	4.7
33	2b	187	LEU	4.6
20	2Y	1	MET	4.6
44	2m	102	ARG	4.6
47	1p	19	ILE	4.5
6	1G	49	ASP	4.5
1	1A	1149	C	4.4
34	2c	186	PHE	4.4
41	2j	62	HIS	4.3
31	29	37	GLY	4.3
38	1g	83	ALA	4.2
26	24	49	PHE	4.2
41	2j	65	LEU	4.2
40	2i	2	GLU	4.2
15	1T	37	GLY	4.1
33	2b	152	PHE	4.1
33	1b	188	ALA	4.1
41	2j	78	ASN	4.1
1	1A	33	C	4.1
32	1a	981	G	4.1
38	2g	80	VAL	4.0
32	1a	980	G	4.0
20	2Y	90	LEU	4.0
26	24	50	VAL	4.0
40	2i	109	VAL	4.0

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Mol	Chain	Res	Type	RSRZ
12	2Q	59	ARG	3.9
6	2G	53	LEU	3.9
40	2i	126	SER	3.9
26	24	52	THR	3.9
1	1A	2815	G	3.9
33	2b	136	VAL	3.9
44	1m	108	ARG	3.9
52	2u	14	TRP	3.9
40	2i	17	VAL	3.9
41	1j	46	ARG	3.9
45	1n	2	ALA	3.9
6	2G	28	VAL	3.8
41	2j	38	ILE	3.8
1	1A	2905	U	3.8
39	2h	2	LEU	3.8
6	2G	76	SER	3.8
26	24	56	VAL	3.8
35	2d	146	ILE	3.8
12	1Q	60	ARG	3.7
34	2c	180	ALA	3.7
35	2d	164	ALA	3.7
38	2g	156	TRP	3.7
41	2j	47	PHE	3.7
42	2k	126	ARG	3.7
47	1p	82	GLN	3.7
40	2i	9	ARG	3.7
40	2i	10	ARG	3.7
24	12	11	GLU	3.7
43	2l	18	VAL	3.7
34	1c	28	GLN	3.6
38	2g	83	ALA	3.6
33	2b	92	TYR	3.6
41	2j	75	ILE	3.6
41	1j	62	HIS	3.6
1	1A	2904	C	3.6
41	2j	55	LYS	3.6
45	2n	34	TYR	3.6
1	1A	637	U	3.6
51	2t	74	LYS	3.6
50	2s	13	ASP	3.6
32	2a	1268	A	3.6
47	1p	42	ARG	3.5

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Mol	Chain	Res	Type	RSRZ
15	1T	130	ALA	3.5
20	2Y	57	GLN	3.5
34	1c	185	GLY	3.5
1	2A	2902	G	3.5
44	1m	2	ALA	3.5
40	2i	68	GLY	3.5
45	2n	25	VAL	3.5
24	12	12	GLU	3.5
26	24	65	ASP	3.5
40	1i	105	ASP	3.5
35	1d	168	ARG	3.5
45	1n	59	ALA	3.5
6	2G	39	ILE	3.5
21	2Z	175	VAL	3.5
38	1g	156	TRP	3.5
1	1A	1220	G	3.4
34	2c	13	GLY	3.4
35	1d	155	LEU	3.4
41	1j	66	ARG	3.4
14	2S	3	ARG	3.4
38	2g	78	ARG	3.4
41	2j	46	ARG	3.4
52	2u	23	PRO	3.4
43	1l	91	LYS	3.4
12	1Q	59	ARG	3.4
32	2a	981	G	3.4
40	2i	5	TYR	3.4
43	2l	64	TYR	3.4
35	1d	101	LEU	3.4
40	1i	2	GLU	3.4
26	14	49	PHE	3.4
33	1b	122	PHE	3.4
41	2j	58	ASP	3.4
1	2A	2815	G	3.3
38	2g	34	GLY	3.3
52	2u	6	ARG	3.3
52	2u	9	ARG	3.3
7	2H	2	SER	3.3
41	1j	42	THR	3.3
50	2s	79	THR	3.3
1	1A	2813	C	3.3
32	2a	1102	C	3.3

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Mol	Chain	Res	Type	RSRZ
40	2i	15	ALA	3.3
16	1U	117	GLN	3.3
45	2n	3	ARG	3.3
41	2j	40	LEU	3.3
8	1I	10	GLU	3.3
32	2a	1509	A	3.3
1	2A	1582	C	3.3
34	1c	207	VAL	3.3
26	24	44	THR	3.3
41	2j	100	THR	3.3
1	2A	2812	G	3.3
7	1H	124	GLU	3.2
26	24	57	GLU	3.2
34	2c	124	ILE	3.2
33	1b	16	HIS	3.2
33	2b	237	ALA	3.2
33	1b	12	GLU	3.2
25	23	2	PRO	3.2
47	1p	41	PRO	3.2
34	2c	8	ILE	3.2
44	2m	118	ALA	3.2
51	1t	18	GLN	3.2
40	1i	68	GLY	3.2
38	1g	78	ARG	3.2
37	1f	88	VAL	3.2
40	2i	11	LYS	3.2
7	2H	82	GLY	3.2
33	2b	131	PRO	3.2
38	2g	85	TYR	3.1
40	1i	10	ARG	3.1
34	2c	187	ALA	3.1
6	1G	75	LYS	3.1
33	2b	70	PHE	3.1
45	2n	18	VAL	3.1
41	2j	52	GLY	3.1
15	1T	131	ALA	3.1
50	1s	56	GLN	3.1
1	2A	2904	C	3.1
40	2i	16	ARG	3.1
50	2s	71	LEU	3.1
41	2j	24	VAL	3.1
41	2j	61	GLU	3.1

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Mol	Chain	Res	Type	RSRZ
41	2j	37	PRO	3.1
29	17	48	LYS	3.1
51	1t	74	LYS	3.1
41	2j	27	ALA	3.1
41	2j	32	ALA	3.1
45	2n	49	HIS	3.1
34	2c	10	PHE	3.1
41	1j	63	PHE	3.1
45	2n	57	ARG	3.1
45	2n	6	LEU	3.1
45	2n	58	LYS	3.1
33	2b	134	GLU	3.0
41	2j	48	THR	3.0
36	1e	22	GLY	3.0
40	2i	20	ARG	3.0
40	2i	66	ARG	3.0
33	1b	61	LEU	3.0
21	2Z	170	THR	3.0
41	1j	98	ILE	3.0
6	1G	50	ALA	3.0
46	1o	87	ILE	3.0
6	2G	172	LEU	3.0
34	2c	64	VAL	3.0
33	2b	133	LYS	2.9
40	1i	78	LYS	2.9
19	2X	68	ARG	2.9
33	2b	121	LEU	2.9
34	2c	2	GLY	2.9
43	2l	63	GLY	2.9
8	2I	11	ASN	2.9
41	1j	64	GLU	2.9
50	2s	2	PRO	2.9
1	2A	10	G	2.9
32	1a	982	G	2.9
1	1A	2806	C	2.9
40	1i	15	ALA	2.9
1	1A	8	U	2.9
7	2H	174	GLY	2.9
52	1u	4	GLY	2.9
26	24	51	ASP	2.9
41	2j	59	SER	2.9
20	2Y	45	VAL	2.9

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Mol	Chain	Res	Type	RSRZ
33	2b	123	ALA	2.9
38	1g	2	ALA	2.9
1	1A	669	C	2.9
1	2A	2901	G	2.9
45	2n	55	GLY	2.9
47	1p	17	TYR	2.9
12	1Q	33	GLY	2.9
34	2c	185	GLY	2.9
51	1t	103	GLY	2.9
35	1d	116	GLN	2.9
34	1c	184	TYR	2.9
41	2j	77	PRO	2.9
38	1g	84	ASN	2.9
26	14	51	ASP	2.9
35	1d	102	ASP	2.9
26	24	61	ARG	2.9
22	10	10	THR	2.9
41	1j	87	THR	2.9
47	1p	22	THR	2.9
34	2c	5	ILE	2.9
41	1j	31	GLY	2.8
52	2u	16	GLY	2.8
22	20	26	TYR	2.8
26	24	67	TYR	2.8
1	1A	2804	G	2.8
32	2a	1184	G	2.8
40	2i	108	VAL	2.8
6	2G	161	THR	2.8
40	2i	79	LEU	2.8
45	2n	22	THR	2.8
26	14	45	GLY	2.8
46	1o	89	GLY	2.8
52	2u	4	GLY	2.8
21	2Z	168	GLU	2.8
44	2m	64	TRP	2.8
2	2B	1	U	2.8
8	2I	81	VAL	2.8
33	1b	7	VAL	2.8
33	1b	165	VAL	2.8
33	1b	237	ALA	2.8
50	2s	40	ILE	2.8
51	1t	10	LEU	2.8

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Mol	Chain	Res	Type	RSRZ
52	2u	13	ILE	2.8
8	2l	86	THR	2.8
26	14	52	THR	2.8
50	2s	4	SER	2.8
32	2a	1356	G	2.8
1	1A	933	A	2.8
41	2j	93	GLY	2.8
47	1p	1	MET	2.8
39	2h	135	CYS	2.8
24	12	70	GLN	2.8
40	2i	21	PRO	2.8
1	1A	1071	U	2.8
29	27	46	VAL	2.8
39	2h	53	VAL	2.8
41	2j	49	VAL	2.8
34	2c	6	HIS	2.8
41	2j	8	LEU	2.8
1	1A	2901	G	2.8
40	2i	49	PRO	2.8
45	2n	40	CYS	2.8
50	2s	80	TYR	2.8
52	1u	18	TYR	2.8
19	1X	95	LEU	2.8
34	2c	62	ASP	2.8
42	2k	31	THR	2.8
47	1p	44	THR	2.8
51	1t	14	LYS	2.8
40	1i	6	GLY	2.8
50	2s	8	GLY	2.8
34	2c	7	PRO	2.8
24	12	69	ARG	2.7
39	2h	122	ARG	2.7
33	1b	15	VAL	2.7
33	2b	97	TRP	2.7
32	2a	1265	G	2.7
40	1i	81	ILE	2.7
40	2i	103	THR	2.7
50	2s	12	ASP	2.7
51	1t	69	GLY	2.7
52	1u	2	GLY	2.7
34	2c	207	VAL	2.7
47	1p	2	VAL	2.7

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Mol	Chain	Res	Type	RSRZ
6	1G	88	ILE	2.7
26	14	59	PHE	2.7
33	2b	69	LEU	2.7
41	2j	54	PHE	2.7
33	2b	140	HIS	2.7
33	2b	132	LYS	2.7
43	1l	126	LYS	2.7
45	2n	4	LYS	2.7
1	2A	2805	G	2.7
40	2i	8	GLY	2.7
26	24	66	SER	2.7
1	2A	1120	C	2.7
1	2A	1121	C	2.7
34	2c	178	LEU	2.7
26	24	63	TYR	2.7
1	1A	1934	A	2.7
40	2i	29	ASN	2.7
35	1d	23	GLY	2.7
44	1m	102	ARG	2.7
52	2u	8	THR	2.7
6	2G	2	PRO	2.7
1	1A	638	G	2.7
35	1d	152	SER	2.7
12	2Q	104	PHE	2.7
34	2c	182	ILE	2.7
33	1b	22	LYS	2.7
38	1g	153	HIS	2.7
42	1k	42	TRP	2.7
50	2s	69	HIS	2.7
4	2E	151	TYR	2.7
42	1k	25	TYR	2.7
20	2Y	50	ARG	2.7
23	21	26	ARG	2.7
43	1l	89	ARG	2.7
34	2c	44	GLU	2.7
39	2h	99	GLU	2.7
1	2A	1160	G	2.6
34	1c	23	TYR	2.6
32	1a	64	C	2.6
6	1G	53	LEU	2.6
20	1Y	57	GLN	2.6
40	1i	12	GLU	2.6

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Mol	Chain	Res	Type	RSRZ
6	2G	29	TRP	2.6
1	1A	2329	G	2.6
32	2a	982	G	2.6
34	1c	201	TYR	2.6
41	2j	70	ARG	2.6
20	2Y	58	GLY	2.6
34	1c	2	GLY	2.6
52	2u	17	THR	2.6
44	2m	40	ASN	2.6
1	1A	553	A	2.6
50	2s	14	HIS	2.6
7	1H	101	ARG	2.6
8	1I	29	TYR	2.6
6	2G	32	PRO	2.6
8	1I	35	LEU	2.6
35	1d	194	LEU	2.6
33	2b	135	GLN	2.6
47	1p	68	ASP	2.6
41	2j	5	ARG	2.6
46	1o	88	ARG	2.6
1	2A	1506	A	2.6
26	14	63	TYR	2.6
35	2d	183	GLY	2.6
21	2Z	57	ILE	2.6
41	2j	44	VAL	2.6
50	2s	9	VAL	2.6
6	2G	50	ALA	2.6
35	2d	110	PHE	2.6
44	1m	107	ALA	2.6
45	2n	5	ALA	2.6
1	2A	1089	G	2.5
1	2A	1108	G	2.5
14	1S	3	ARG	2.5
45	2n	29	ARG	2.5
1	1A	2814	C	2.5
7	2H	175	LYS	2.5
1	1A	2802	A	2.5
32	2a	1339	A	2.5
40	2i	67	GLY	2.5
45	2n	56	VAL	2.5
3	1D	28	GLU	2.5
33	2b	190	THR	2.5

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Mol	Chain	Res	Type	RSRZ
41	2j	67	THR	2.5
20	1Y	107	ASP	2.5
20	2Y	94	LYS	2.5
38	1g	77	SER	2.5
40	1i	11	LYS	2.5
41	2j	35	SER	2.5
33	2b	155	LEU	2.5
49	1r	78	LEU	2.5
49	1r	85	LEU	2.5
51	2t	103	GLY	2.5
6	2G	31	VAL	2.5
9	2N	9	VAL	2.5
26	14	50	VAL	2.5
33	2b	162	ILE	2.5
42	2k	75	TYR	2.5
40	1i	7	THR	2.5
34	1c	65	ALA	2.5
41	1j	5	ARG	2.5
43	1l	28	LYS	2.5
45	2n	32	SER	2.5
40	2i	123	PRO	2.5
41	1j	65	LEU	2.5
8	2I	79	ILE	2.5
14	2S	60	GLY	2.5
33	2b	127	ILE	2.5
35	1d	158	ILE	2.5
44	1m	4	ILE	2.5
50	2s	49	ILE	2.5
6	1G	149	VAL	2.5
7	1H	131	VAL	2.5
34	2c	66	VAL	2.5
35	1d	112	VAL	2.5
1	2A	2227	G	2.5
8	2I	48	GLU	2.5
40	2i	82	ALA	2.5
45	2n	59	ALA	2.5
40	1i	66	ARG	2.5
35	2d	154	ASN	2.5
52	2u	5	ASP	2.5
6	2G	19	LEU	2.5
22	20	84	LEU	2.5
33	1b	83	MET	2.5

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Mol	Chain	Res	Type	RSRZ
40	2i	19	LEU	2.5
44	2m	66	LEU	2.5
45	1n	44	LEU	2.5
41	1j	4	ILE	2.5
45	2n	60	SER	2.5
33	2b	165	VAL	2.5
33	2b	163	PHE	2.5
6	2G	69	ALA	2.5
15	2T	131	ALA	2.5
33	2b	137	ARG	2.5
41	2j	28	ARG	2.5
45	1n	3	ARG	2.5
1	2A	2801	C	2.5
32	1a	153	G	2.5
32	1a	979	A	2.5
41	2j	69	ASN	2.5
35	1d	135	LEU	2.4
6	2G	20	ILE	2.4
7	2H	115	VAL	2.4
33	2b	99	GLY	2.4
35	1d	167	GLY	2.4
40	1i	109	VAL	2.4
35	2d	209	ARG	2.4
3	1D	38	LYS	2.4
33	2b	236	TYR	2.4
34	2c	60	ALA	2.4
40	2i	36	TYR	2.4
40	2i	125	TYR	2.4
44	2m	75	ALA	2.4
52	2u	21	TYR	2.4
1	2A	696	C	2.4
1	2A	2813	C	2.4
32	1a	1022	C	2.4
33	2b	138	LEU	2.4
35	1d	162	LEU	2.4
35	1d	165	MET	2.4
1	2A	1155	G	2.4
1	2A	1579	G	2.4
1	2A	2816	G	2.4
33	1b	222	ILE	2.4
45	1n	7	ILE	2.4
48	2q	65	ILE	2.4

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Mol	Chain	Res	Type	RSRZ
33	1b	228	GLY	2.4
39	1h	88	LYS	2.4
40	2i	118	LYS	2.4
50	2s	74	PHE	2.4
24	22	70	GLN	2.4
26	14	67	TYR	2.4
43	1l	64	TYR	2.4
52	2u	18	TYR	2.4
22	20	10	THR	2.4
33	1b	187	LEU	2.4
38	2g	16	LEU	2.4
41	2j	71	LEU	2.4
40	1i	14	VAL	2.4
40	2i	65	VAL	2.4
50	1s	9	VAL	2.4
50	2s	68	GLY	2.4
53	2v	14	A	2.4
6	1G	47	LYS	2.4
24	12	15	LYS	2.4
26	14	55	ARG	2.4
1	2A	1092	G	2.4
33	2b	150	SER	2.4
16	1U	81	HIS	2.4
21	2Z	150	LEU	2.4
33	1b	10	LEU	2.4
45	2n	7	ILE	2.4
43	2l	5	PRO	2.4
6	2G	38	VAL	2.4
14	1S	46	VAL	2.4
26	14	56	VAL	2.4
35	2d	112	VAL	2.4
40	1i	41	VAL	2.4
47	1p	20	VAL	2.4
50	2s	11	VAL	2.4
45	2n	9	LYS	2.4
8	2I	146	ALA	2.4
1	1A	1465	U	2.4
32	1a	614	G	2.4
32	2a	1239	U	2.4
40	2i	4	TYR	2.4
51	2t	10	LEU	2.4
6	2G	17	PRO	2.4

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Mol	Chain	Res	Type	RSRZ
6	2G	33	ARG	2.3
7	1H	132	ARG	2.3
8	2I	92	VAL	2.3
35	2d	46	LYS	2.3
44	2m	80	ARG	2.3
6	1G	80	PHE	2.3
6	2G	102	PHE	2.3
38	2g	43	PHE	2.3
45	2n	27	CYS	2.3
47	2p	9	PHE	2.3
33	1b	123	ALA	2.3
40	2i	106	ALA	2.3
32	2a	1266	C	2.3
41	1j	86	MET	2.3
34	2c	48	TYR	2.3
34	2c	201	TYR	2.3
45	1n	13	THR	2.3
7	2H	79	VAL	2.3
32	1a	152	G	2.3
32	1a	176	G	2.3
32	2a	1172	G	2.3
35	1d	3	ARG	2.3
40	1i	93	ARG	2.3
41	2j	29	ARG	2.3
41	1j	69	ASN	2.3
6	2G	141	PHE	2.3
8	2I	10	GLU	2.3
8	2I	46	ALA	2.3
25	13	60	GLU	2.3
33	2b	143	GLU	2.3
35	1d	156	GLU	2.3
40	2i	91	ASP	2.3
35	1d	164	ALA	2.3
47	1p	76	GLN	2.3
1	1A	571	A	2.3
19	2X	69	TYR	2.3
32	2a	396	C	2.3
32	2a	1037	C	2.3
32	2a	1226	C	2.3
47	2p	13	HIS	2.3
40	2i	81	ILE	2.3
41	2j	81	THR	2.3

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Mol	Chain	Res	Type	RSRZ
50	2s	31	ILE	2.3
7	1H	3	ARG	2.3
26	24	29	PRO	2.3
29	17	23	ARG	2.3
33	2b	130	ARG	2.3
41	2j	66	ARG	2.3
52	2u	22	ARG	2.3
6	2G	15	VAL	2.3
21	2Z	56	VAL	2.3
40	1i	65	VAL	2.3
40	2i	6	GLY	2.3
1	2A	1216	G	2.3
6	2G	48	GLU	2.3
32	1a	65	G	2.3
32	2a	980	G	2.3
32	2a	1206	G	2.3
35	2d	179	GLU	2.3
21	2Z	1	MET	2.3
39	2h	4	ASP	2.3
33	1b	158	LEU	2.3
47	2p	82	GLN	2.3
14	2S	35	ILE	2.3
41	1j	67	THR	2.3
41	1j	100	THR	2.3
45	1n	22	THR	2.3
47	1p	45	THR	2.3
35	1d	153	ARG	2.3
38	1g	5	ARG	2.3
45	1n	29	ARG	2.3
45	2n	21	TYR	2.3
20	1Y	56	PRO	2.3
35	1d	92	VAL	2.3
40	2i	14	VAL	2.3
41	1j	34	VAL	2.3
55	2z	13	PRO	2.3
1	2A	33	C	2.3
32	1a	1120	C	2.3
32	1a	1268	A	2.3
9	2N	49	GLY	2.3
42	2k	90	GLY	2.3
47	1p	14	ASN	2.3
3	2D	153	ALA	2.3

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Mol	Chain	Res	Type	RSRZ
6	2G	86	MET	2.3
15	1T	127	ALA	2.3
33	1b	225	ALA	2.3
34	2c	137	ALA	2.3
40	1i	19	LEU	2.3
50	2s	16	LEU	2.3
34	1c	31	HIS	2.3
19	1X	68	ARG	2.3
33	1b	97	TRP	2.3
33	2b	124	SER	2.3
38	2g	115	ARG	2.3
40	1i	126	SER	2.3
8	1I	15	VAL	2.3
46	2o	2	PRO	2.3
50	1s	19	VAL	2.3
40	2i	115	GLY	2.3
32	2a	1243	A	2.2
32	2a	1344	C	2.2
40	2i	37	PHE	2.2
33	2b	129	GLU	2.2
35	1d	111	ALA	2.2
14	2S	54	LEU	2.2
37	1f	54	LYS	2.2
41	1j	33	GLN	2.2
48	1q	87	LYS	2.2
45	2n	42	ILE	2.2
38	2g	32	ARG	2.2
38	2g	76	ARG	2.2
47	1p	13	HIS	2.2
52	2u	15	ARG	2.2
33	2b	67	THR	2.2
7	1H	133	VAL	2.2
45	1n	60	SER	2.2
51	1t	11	SER	2.2
1	1A	2330	G	2.2
1	2A	2329	G	2.2
32	1a	1121	G	2.2
40	1i	18	PHE	2.2
1	1A	270	U	2.2
1	1A	1984	U	2.2
35	1d	48	ALA	2.2
1	2A	1934	A	2.2

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Mol	Chain	Res	Type	RSRZ
1	2A	2900	A	2.2
33	2b	44	LEU	2.2
38	1g	16	LEU	2.2
33	2b	222	ILE	2.2
34	2c	57	ILE	2.2
40	2i	53	VAL	2.2
41	2j	42	THR	2.2
41	1j	30	SER	2.2
23	21	28	GLY	2.2
26	14	54	GLY	2.2
34	2c	9	GLY	2.2
34	2c	158	GLY	2.2
21	2Z	136	PHE	2.2
1	2A	2330	G	2.2
6	2G	43	LEU	2.2
6	2G	175	LEU	2.2
10	1O	91	LEU	2.2
12	2Q	28	ALA	2.2
14	1S	93	LYS	2.2
33	2b	154	LEU	2.2
35	2d	156	GLU	2.2
10	1O	49	ARG	2.2
25	13	55	ARG	2.2
34	2c	179	ARG	2.2
47	1p	25	ARG	2.2
52	2u	24	ARG	2.2
1	2A	2811	A	2.2
1	1A	935	C	2.2
14	2S	34	HIS	2.2
8	2I	8	PRO	2.2
34	2c	153	VAL	2.2
36	1e	67	VAL	2.2
36	2e	82	VAL	2.2
41	1j	48	THR	2.2
47	2p	69	THR	2.2
50	2s	34	TRP	2.2
33	2b	151	GLY	2.2
6	2G	74	LYS	2.2
14	2S	29	PHE	2.2
35	1d	188	LEU	2.2
3	2D	28	GLU	2.2
25	23	60	GLU	2.2

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Mol	Chain	Res	Type	RSRZ
21	2Z	4	ARG	2.2
32	1a	1023	U	2.2
33	1b	130	ARG	2.2
39	2h	92	ARG	2.2
47	1p	5	ARG	2.2
1	1A	2816	G	2.2
1	1A	2902	G	2.2
1	2A	9	G	2.2
1	2A	2226	G	2.2
2	2B	118	G	2.2
31	29	20	HIS	2.2
32	2a	1138	G	2.2
40	2i	28	VAL	2.2
32	2a	1120	C	2.2
35	2d	124	GLY	2.2
47	1p	37	GLY	2.2
40	2i	62	TYR	2.2
47	1p	39	TYR	2.2
50	2s	52	TYR	2.2
6	1G	46	ALA	2.2
6	2G	95	ARG	2.1
41	1j	79	ARG	2.1
44	1m	110	ARG	2.1
52	1u	15	ARG	2.1
44	2m	15	VAL	2.1
1	1A	2640	A	2.1
12	2Q	30	GLY	2.1
21	2Z	143	GLY	2.1
32	2a	1267	A	2.1
38	2g	82	GLY	2.1
40	1i	8	GLY	2.1
41	1j	93	GLY	2.1
52	2u	11	GLY	2.1
1	1A	154	C	2.1
32	1a	160	C	2.1
37	1f	63	TYR	2.1
38	1g	85	TYR	2.1
3	2D	2	ALA	2.1
12	2Q	60	ARG	2.1
14	1S	13	ARG	2.1
28	16	4	GLU	2.1
34	2c	131	ARG	2.1

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Mol	Chain	Res	Type	RSRZ
45	2n	8	GLU	2.1
46	2o	68	ARG	2.1
33	1b	19	HIS	2.1
41	1j	78	ASN	2.1
1	1A	1150	U	2.1
7	2H	17	VAL	2.1
7	2H	45	VAL	2.1
35	2d	170	VAL	2.1
37	1f	6	VAL	2.1
40	1i	17	VAL	2.1
44	2m	41	PRO	2.1
46	2o	60	VAL	2.1
20	1Y	1	MET	2.1
31	19	12	ASP	2.1
47	1p	40	ASP	2.1
41	2j	31	GLY	2.1
41	1j	8	LEU	2.1
1	1A	2811	A	2.1
7	1H	130	ARG	2.1
33	2b	21	ARG	2.1
38	2g	79	ARG	2.1
40	2i	42	ARG	2.1
31	19	17	ILE	2.1
32	1a	96	G	2.1
32	1a	151	G	2.1
32	2a	951	G	2.1
33	2b	126	GLU	2.1
34	2c	14	ILE	2.1
40	1i	63	ILE	2.1
32	1a	158	C	2.1
34	2c	162	GLN	2.1
38	2g	84	ASN	2.1
41	2j	72	VAL	2.1
40	2i	98	PRO	2.1
47	1p	66	PRO	2.1
38	1g	82	GLY	2.1
44	2m	16	ASP	2.1
44	2m	100	GLY	2.1
22	20	55	ARG	2.1
45	2n	35	ARG	2.1
47	1p	48	TRP	2.1
29	17	45	ALA	2.1

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Mol	Chain	Res	Type	RSRZ
34	2c	129	ALA	2.1
39	2h	111	ILE	2.1
41	2j	6	ILE	2.1
45	2n	20	ALA	2.1
6	1G	54	GLU	2.1
8	2I	99	GLU	2.1
24	12	20	GLU	2.1
33	2b	86	GLU	2.1
44	2m	69	GLU	2.1
1	2A	5	A	2.1
1	2A	1090	A	2.1
26	24	69	LYS	2.1
32	1a	376	G	2.1
32	1a	1002	G	2.1
32	2a	1307	C	2.1
41	2j	94	VAL	2.1
42	1k	51	LYS	2.1
50	2s	47	HIS	2.1
41	2j	34	VAL	2.1
43	1l	18	VAL	2.1
48	2q	9	VAL	2.1
14	2S	61	ASN	2.1
7	2H	123	PHE	2.1
26	24	45	GLY	2.1
26	24	54	GLY	2.1
6	2G	181	ARG	2.1
12	1Q	21	THR	2.1
33	1b	107	THR	2.1
48	1q	68	ARG	2.1
50	1s	13	ASP	2.1
50	2s	78	ARG	2.1
15	2T	130	ALA	2.1
42	2k	89	ALA	2.1
41	1j	61	GLU	2.1
35	2d	20	TYR	2.1
33	1b	40	HIS	2.0
41	2j	13	HIS	2.0
14	2S	46	VAL	2.0
34	2c	106	VAL	2.0
36	2e	41	VAL	2.0
53	1v	14	A	2.0
33	1b	125	PRO	2.0

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Mol	Chain	Res	Type	RSRZ
1	1A	932	C	2.0
1	1A	1554	C	2.0
1	1A	2206	C	2.0
6	2G	3	LEU	2.0
32	2a	1345	C	2.0
40	1i	56	LEU	2.0
44	2m	70	LEU	2.0
47	1p	49	LEU	2.0
1	1A	2211	G	2.0
11	1P	116	GLY	2.0
32	2a	1469	G	2.0
33	1b	163	PHE	2.0
35	2d	180	GLY	2.0
40	1i	69	GLY	2.0
51	1t	71	THR	2.0
12	1Q	135	ASP	2.0
20	1Y	91	GLU	2.0
6	1G	84	LYS	2.0
40	2i	112	LYS	2.0
33	1b	229	VAL	2.0
33	2b	7	VAL	2.0
35	1d	133	VAL	2.0
35	1d	178	VAL	2.0
42	2k	114	VAL	2.0
25	13	2	PRO	2.0
34	1c	7	PRO	2.0
41	1j	77	PRO	2.0
41	2j	41	PRO	2.0
43	1l	94	PRO	2.0
2	2B	45	A	2.0
8	2I	38	LEU	2.0
32	2a	984	A	2.0
33	2b	196	LEU	2.0
35	2d	155	LEU	2.0
41	2j	16	LEU	2.0
35	1d	122	ARG	2.0
35	2d	122	ARG	2.0
40	1i	9	ARG	2.0
33	2b	57	PHE	2.0
35	1d	110	PHE	2.0
35	2d	87	GLY	2.0
35	2d	109	GLY	2.0

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Mol	Chain	Res	Type	RSRZ
50	2s	82	GLY	2.0
1	1A	1098	C	2.0
32	1a	220	C	2.0
36	2e	21	ALA	2.0
20	2Y	91	GLU	2.0
32	2a	1107	G	2.0
40	1i	113	LYS	2.0
20	2Y	5	MET	2.0
33	2b	83	MET	2.0
34	1c	195	VAL	2.0
36	2e	67	VAL	2.0
50	2s	67	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	0.75	0.13	69,77,85,97	0
54	4SU	2x	8	20/21	0.84	0.10	70,77,94,99	0
54	5MU	2x	54	21/22	0.87	0.11	75,81,92,101	0
54	PSU	1x	55	20/21	0.91	0.11	62,73,83,85	0
54	5MU	1x	54	21/22	0.92	0.10	57,66,80,87	0
54	5MC	2x	32	21/22	0.92	0.12	61,77,79,91	0
54	5MC	1x	32	21/22	0.93	0.10	48,59,69,81	0
54	4SU	1x	8	20/21	0.94	0.10	51,69,76,81	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	3429	1/1	0.33	0.24	70,70,70,70	0
56	MG	2a	3256	1/1	0.35	0.29	94,94,94,94	0
56	MG	2A	3455	1/1	0.48	0.30	69,69,69,69	0
56	MG	1B	213	1/1	0.53	0.15	110,110,110,110	0
56	MG	2a	3052	1/1	0.56	0.17	95,95,95,95	0
56	MG	1A	3467	1/1	0.56	0.15	71,71,71,71	0
56	MG	1A	3936	1/1	0.58	0.17	64,64,64,64	0
56	MG	2a	3207	1/1	0.58	0.22	73,73,73,73	0
56	MG	2A	3484	1/1	0.58	0.16	82,82,82,82	0
56	MG	2a	3254	1/1	0.59	0.16	71,71,71,71	0
56	MG	1A	3032	1/1	0.59	0.10	63,63,63,63	0
56	MG	1A	3378	1/1	0.61	0.21	69,69,69,69	0
56	MG	2A	3511	1/1	0.61	0.25	52,52,52,52	0
56	MG	1A	3322	1/1	0.61	0.33	50,50,50,50	0
56	MG	2A	3443	1/1	0.62	0.20	56,56,56,56	0
56	MG	2A	3036	1/1	0.62	0.24	74,74,74,74	0
56	MG	2a	3123	1/1	0.63	0.25	81,81,81,81	0
56	MG	2a	3158	1/1	0.64	0.23	74,74,74,74	0
56	MG	2A	3515	1/1	0.64	0.13	63,63,63,63	0
56	MG	2a	3235	1/1	0.64	0.28	68,68,68,68	0
56	MG	1a	1755	1/1	0.64	0.17	79,79,79,79	0
56	MG	2A	3210	1/1	0.64	0.11	63,63,63,63	0
56	MG	2A	3529	1/1	0.65	0.15	68,68,68,68	0
56	MG	2A	3540	1/1	0.65	0.28	58,58,58,58	0
56	MG	1A	3840	1/1	0.65	0.24	63,63,63,63	0
56	MG	2a	3156	1/1	0.66	0.26	72,72,72,72	0
56	MG	1A	3837	1/1	0.66	0.16	62,62,62,62	0
56	MG	2A	3497	1/1	0.66	0.17	54,54,54,54	0
56	MG	2G	3001	1/1	0.66	0.12	58,58,58,58	0
56	MG	1a	1865	1/1	0.66	0.22	69,69,69,69	0
56	MG	1A	3521	1/1	0.66	0.22	75,75,75,75	0
56	MG	1A	3433	1/1	0.67	0.12	71,71,71,71	0
56	MG	1a	1775	1/1	0.67	0.28	80,80,80,80	0
56	MG	1A	3331	1/1	0.67	0.18	57,57,57,57	0
56	MG	2a	3205	1/1	0.67	0.22	74,74,74,74	0
56	MG	1A	3838	1/1	0.67	0.22	66,66,66,66	0
56	MG	2a	3051	1/1	0.67	0.20	76,76,76,76	0
56	MG	18	101	1/1	0.67	0.23	63,63,63,63	0
56	MG	2a	3058	1/1	0.67	0.26	87,87,87,87	0
56	MG	1a	1791	1/1	0.68	0.15	99,99,99,99	0
56	MG	2a	3065	1/1	0.68	0.20	80,80,80,80	0
56	MG	1a	1854	1/1	0.68	0.19	82,82,82,82	0
56	MG	1A	3505	1/1	0.68	0.12	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	3013	1/1	0.68	0.30	70,70,70,70	0
56	MG	1A	3520	1/1	0.69	0.17	75,75,75,75	0
56	MG	1a	1750	1/1	0.69	0.15	64,64,64,64	0
56	MG	2a	3024	1/1	0.69	0.45	76,76,76,76	0
56	MG	1A	3036	1/1	0.70	0.10	55,55,55,55	0
56	MG	2A	3400	1/1	0.70	0.27	48,48,48,48	0
56	MG	1a	1754	1/1	0.70	0.23	81,81,81,81	0
56	MG	1A	3434	1/1	0.70	0.13	70,70,70,70	0
56	MG	1a	1694	1/1	0.70	0.25	78,78,78,78	0
56	MG	2A	3118	1/1	0.70	0.29	75,75,75,75	0
56	MG	2a	3151	1/1	0.71	0.17	92,92,92,92	0
56	MG	2A	3266	1/1	0.71	0.19	71,71,71,71	0
56	MG	2A	3551	1/1	0.71	0.13	57,57,57,57	0
56	MG	2A	3381	1/1	0.71	0.20	64,64,64,64	0
56	MG	1A	3403	1/1	0.72	0.14	59,59,59,59	0
56	MG	2a	3152	1/1	0.72	0.10	84,84,84,84	0
56	MG	1A	3487	1/1	0.72	0.24	63,63,63,63	0
56	MG	1A	3855	1/1	0.72	0.20	84,84,84,84	0
56	MG	2a	3191	1/1	0.72	0.20	54,54,54,54	0
56	MG	2A	3509	1/1	0.72	0.26	62,62,62,62	0
56	MG	1A	3647	1/1	0.72	0.23	74,74,74,74	0
56	MG	2A	3389	1/1	0.72	0.30	77,77,77,77	0
56	MG	1A	3327	1/1	0.72	0.18	73,73,73,73	0
56	MG	2A	3061	1/1	0.72	0.31	68,68,68,68	0
56	MG	2x	3003	1/1	0.72	0.20	73,73,73,73	0
56	MG	1a	1744	1/1	0.73	0.25	61,61,61,61	0
56	MG	1A	3769	1/1	0.73	0.23	90,90,90,90	0
56	MG	1a	1796	1/1	0.73	0.14	86,86,86,86	0
56	MG	2A	3141	1/1	0.73	0.20	74,74,74,74	0
56	MG	2a	3159	1/1	0.73	0.32	85,85,85,85	0
56	MG	2A	3187	1/1	0.73	0.29	49,49,49,49	0
56	MG	2A	3486	1/1	0.73	0.19	57,57,57,57	0
56	MG	1A	3349	1/1	0.73	0.22	73,73,73,73	0
56	MG	1A	3333	1/1	0.73	0.18	70,70,70,70	0
56	MG	2A	3380	1/1	0.73	0.23	74,74,74,74	0
56	MG	2a	3092	1/1	0.73	0.24	87,87,87,87	0
56	MG	1f	3001	1/1	0.73	0.16	68,68,68,68	0
56	MG	2A	3507	1/1	0.74	0.14	47,47,47,47	0
56	MG	2a	3171	1/1	0.74	0.21	68,68,68,68	0
56	MG	2B	3009	1/1	0.74	0.17	63,63,63,63	0
56	MG	1A	3705	1/1	0.74	0.22	34,34,34,34	0
56	MG	2a	3138	1/1	0.74	0.42	81,81,81,81	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3031	1/1	0.74	0.10	61,61,61,61	0
56	MG	2A	3218	1/1	0.74	0.24	62,62,62,62	0
56	MG	1A	3668	1/1	0.74	0.17	60,60,60,60	0
56	MG	2A	3441	1/1	0.74	0.23	58,58,58,58	0
56	MG	2A	3315	1/1	0.75	0.18	55,55,55,55	0
56	MG	1A	3392	1/1	0.75	0.22	52,52,52,52	0
56	MG	1A	3350	1/1	0.75	0.28	54,54,54,54	0
56	MG	2A	3005	1/1	0.75	0.21	54,54,54,54	0
56	MG	1N	3002	1/1	0.75	0.12	48,48,48,48	0
56	MG	1a	1760	1/1	0.75	0.18	85,85,85,85	0
56	MG	1a	1769	1/1	0.75	0.21	87,87,87,87	0
56	MG	2a	3014	1/1	0.75	0.14	65,65,65,65	0
56	MG	1a	1774	1/1	0.75	0.30	76,76,76,76	0
56	MG	2a	3031	1/1	0.75	0.27	65,65,65,65	0
56	MG	1A	3513	1/1	0.75	0.21	57,57,57,57	0
56	MG	2a	3221	1/1	0.75	0.25	66,66,66,66	0
56	MG	1A	3005	1/1	0.75	0.14	58,58,58,58	0
56	MG	2a	3239	1/1	0.75	0.32	76,76,76,76	0
56	MG	2a	3241	1/1	0.75	0.14	95,95,95,95	0
56	MG	1a	1732	1/1	0.75	0.28	74,74,74,74	0
56	MG	1A	3757	1/1	0.75	0.14	39,39,39,39	0
56	MG	2q	201	1/1	0.75	0.18	58,58,58,58	0
56	MG	2A	3273	1/1	0.75	0.16	41,41,41,41	0
56	MG	1A	3876	1/1	0.76	0.24	79,79,79,79	0
56	MG	1a	1843	1/1	0.76	0.29	74,74,74,74	0
56	MG	1A	3464	1/1	0.76	0.34	76,76,76,76	0
56	MG	2E	304	1/1	0.76	0.13	67,67,67,67	0
56	MG	1a	1864	1/1	0.76	0.26	73,73,73,73	0
56	MG	1A	3675	1/1	0.76	0.16	79,79,79,79	0
56	MG	1H	8002	1/1	0.76	0.16	79,79,79,79	0
56	MG	1A	3417	1/1	0.76	0.18	58,58,58,58	0
56	MG	2a	3045	1/1	0.76	0.16	74,74,74,74	0
56	MG	2A	3524	1/1	0.76	0.09	41,41,41,41	0
56	MG	10	101	1/1	0.76	0.12	61,61,61,61	0
56	MG	1a	1627	1/1	0.77	0.20	67,67,67,67	0
56	MG	2A	3145	1/1	0.77	0.27	77,77,77,77	0
56	MG	2A	3164	1/1	0.77	0.16	64,64,64,64	0
56	MG	1a	1689	1/1	0.77	0.38	65,65,65,65	0
56	MG	2a	3167	1/1	0.77	0.21	80,80,80,80	0
56	MG	1A	3385	1/1	0.77	0.12	59,59,59,59	0
56	MG	1A	3145	1/1	0.77	0.18	77,77,77,77	0
56	MG	1B	219	1/1	0.77	0.13	60,60,60,60	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3372	1/1	0.77	0.28	57,57,57,57	0
56	MG	2a	3055	1/1	0.77	0.09	98,98,98,98	0
56	MG	1A	3324	1/1	0.77	0.18	57,57,57,57	0
56	MG	1T	204	1/1	0.77	0.17	67,67,67,67	0
56	MG	1A	3831	1/1	0.77	0.25	70,70,70,70	0
56	MG	2a	3113	1/1	0.77	0.29	65,65,65,65	0
56	MG	1a	1761	1/1	0.77	0.23	86,86,86,86	0
56	MG	1A	3928	1/1	0.77	0.12	62,62,62,62	0
56	MG	1a	1611	1/1	0.77	0.20	78,78,78,78	0
56	MG	1A	3384	1/1	0.78	0.19	69,69,69,69	0
56	MG	2A	3258	1/1	0.78	0.18	30,30,30,30	0
56	MG	1A	3399	1/1	0.78	0.22	58,58,58,58	0
56	MG	2B	3004	1/1	0.78	0.20	75,75,75,75	0
56	MG	2a	3075	1/1	0.78	0.34	72,72,72,72	0
56	MG	2A	3119	1/1	0.78	0.18	61,61,61,61	0
56	MG	2a	3210	1/1	0.78	0.18	62,62,62,62	0
56	MG	2A	3495	1/1	0.78	0.10	71,71,71,71	0
56	MG	2a	3234	1/1	0.78	0.18	66,66,66,66	0
56	MG	1A	3779	1/1	0.78	0.19	54,54,54,54	0
56	MG	1a	1867	1/1	0.78	0.31	80,80,80,80	0
56	MG	2a	3139	1/1	0.78	0.45	70,70,70,70	0
56	MG	1a	1798	1/1	0.78	0.28	71,71,71,71	0
56	MG	1A	3787	1/1	0.78	0.12	74,74,74,74	0
56	MG	1a	1852	1/1	0.78	0.25	70,70,70,70	0
56	MG	2A	3211	1/1	0.78	0.16	63,63,63,63	0
56	MG	2a	3013	1/1	0.79	0.26	72,72,72,72	0
56	MG	1A	3132	1/1	0.79	0.20	51,51,51,51	0
56	MG	2A	3232	1/1	0.79	0.12	54,54,54,54	0
56	MG	2A	3488	1/1	0.79	0.26	63,63,63,63	0
56	MG	2A	3013	1/1	0.79	0.27	68,68,68,68	0
56	MG	1A	3352	1/1	0.79	0.16	58,58,58,58	0
56	MG	2a	3194	1/1	0.79	0.28	72,72,72,72	0
56	MG	1a	1801	1/1	0.79	0.23	59,59,59,59	0
56	MG	1A	3503	1/1	0.79	0.11	62,62,62,62	0
56	MG	2A	3326	1/1	0.79	0.17	42,42,42,42	0
56	MG	2A	3358	1/1	0.79	0.18	59,59,59,59	0
56	MG	1A	3939	1/1	0.79	0.22	69,69,69,69	0
56	MG	1A	3460	1/1	0.79	0.11	36,36,36,36	0
56	MG	1A	3371	1/1	0.79	0.23	56,56,56,56	0
56	MG	1A	3867	1/1	0.79	0.16	30,30,30,30	0
56	MG	1A	3875	1/1	0.79	0.20	83,83,83,83	0
56	MG	1N	3006	1/1	0.79	0.19	71,71,71,71	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1n	502	1/1	0.79	0.14	67,67,67,67	0
56	MG	2A	3468	1/1	0.79	0.21	56,56,56,56	0
56	MG	2a	3067	1/1	0.80	0.18	88,88,88,88	0
56	MG	2A	3499	1/1	0.80	0.21	47,47,47,47	0
56	MG	1a	1608	1/1	0.80	0.17	59,59,59,59	0
56	MG	2A	3508	1/1	0.80	0.21	56,56,56,56	0
56	MG	2A	3264	1/1	0.80	0.18	64,64,64,64	0
56	MG	1a	1765	1/1	0.80	0.23	57,57,57,57	0
56	MG	1A	3724	1/1	0.80	0.21	58,58,58,58	0
56	MG	2A	3274	1/1	0.80	0.26	64,64,64,64	0
56	MG	1a	1773	1/1	0.80	0.14	77,77,77,77	0
56	MG	2a	3153	1/1	0.80	0.14	67,67,67,67	0
56	MG	2A	3530	1/1	0.80	0.21	46,46,46,46	0
56	MG	2A	3033	1/1	0.80	0.23	61,61,61,61	0
56	MG	1A	3944	1/1	0.80	0.17	68,68,68,68	0
56	MG	2a	3160	1/1	0.80	0.22	74,74,74,74	0
56	MG	1a	1676	1/1	0.80	0.30	74,74,74,74	0
56	MG	2A	3071	1/1	0.80	0.29	68,68,68,68	0
56	MG	1A	3451	1/1	0.80	0.29	66,66,66,66	0
56	MG	1A	3455	1/1	0.80	0.29	63,63,63,63	0
56	MG	2A	3413	1/1	0.80	0.20	62,62,62,62	0
56	MG	1a	1724	1/1	0.80	0.20	76,76,76,76	0
56	MG	1A	3863	1/1	0.80	0.16	78,78,78,78	0
56	MG	2a	3214	1/1	0.80	0.28	97,97,97,97	0
56	MG	1A	3458	1/1	0.80	0.13	46,46,46,46	0
56	MG	2a	3035	1/1	0.80	0.10	69,69,69,69	0
56	MG	1a	1748	1/1	0.80	0.17	52,52,52,52	0
56	MG	1A	3105	1/1	0.80	0.19	61,61,61,61	0
56	MG	1A	3792	1/1	0.80	0.17	79,79,79,79	0
56	MG	1A	3148	1/1	0.80	0.10	62,62,62,62	0
56	MG	2A	3228	1/1	0.80	0.20	57,57,57,57	0
56	MG	2a	3257	1/1	0.80	0.23	90,90,90,90	0
56	MG	2a	3060	1/1	0.80	0.12	86,86,86,86	0
56	MG	1A	3515	1/1	0.80	0.16	48,48,48,48	0
56	MG	1T	202	1/1	0.81	0.13	55,55,55,55	0
56	MG	1A	3361	1/1	0.81	0.15	66,66,66,66	0
56	MG	1A	3509	1/1	0.81	0.26	71,71,71,71	0
56	MG	2A	3328	1/1	0.81	0.19	57,57,57,57	0
56	MG	1A	3151	1/1	0.81	0.17	47,47,47,47	0
56	MG	1A	3889	1/1	0.81	0.14	60,60,60,60	0
56	MG	1A	3165	1/1	0.81	0.20	53,53,53,53	0
56	MG	1A	3406	1/1	0.81	0.15	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	1642	1/1	0.81	0.31	69,69,69,69	0
56	MG	2A	3401	1/1	0.81	0.13	48,48,48,48	0
56	MG	1A	3806	1/1	0.81	0.26	61,61,61,61	0
56	MG	2a	3170	1/1	0.81	0.24	77,77,77,77	0
56	MG	2a	3012	1/1	0.81	0.31	66,66,66,66	0
56	MG	1a	1795	1/1	0.81	0.21	75,75,75,75	0
56	MG	1A	3377	1/1	0.81	0.18	61,61,61,61	0
56	MG	2A	3453	1/1	0.81	0.21	61,61,61,61	0
56	MG	1A	3995	1/1	0.81	0.35	42,42,42,42	0
56	MG	1a	1717	1/1	0.81	0.15	63,63,63,63	0
56	MG	1A	3998	1/1	0.81	0.09	75,75,75,75	0
56	MG	1A	3167	1/1	0.81	0.13	35,35,35,35	0
56	MG	1A	3228	1/1	0.81	0.24	59,59,59,59	0
56	MG	1A	3493	1/1	0.81	0.17	65,65,65,65	0
56	MG	1A	3499	1/1	0.81	0.15	62,62,62,62	0
56	MG	1a	1751	1/1	0.81	0.16	61,61,61,61	0
56	MG	2A	3501	1/1	0.81	0.13	85,85,85,85	0
56	MG	2A	3505	1/1	0.81	0.15	57,57,57,57	0
56	MG	1a	1869	1/1	0.81	0.31	64,64,64,64	0
56	MG	1b	3002	1/1	0.81	0.12	76,76,76,76	0
56	MG	1A	3357	1/1	0.81	0.25	68,68,68,68	0
56	MG	2a	3066	1/1	0.82	0.12	84,84,84,84	0
56	MG	1a	1857	1/1	0.82	0.22	73,73,73,73	0
56	MG	1a	1860	1/1	0.82	0.30	55,55,55,55	0
56	MG	1A	3488	1/1	0.82	0.10	65,65,65,65	0
56	MG	1a	1609	1/1	0.82	0.15	63,63,63,63	0
56	MG	2A	3240	1/1	0.82	0.12	63,63,63,63	0
56	MG	1A	3130	1/1	0.82	0.16	58,58,58,58	0
56	MG	1a	1623	1/1	0.82	0.21	72,72,72,72	0
56	MG	1a	1876	1/1	0.82	0.12	80,80,80,80	0
56	MG	1A	3459	1/1	0.82	0.17	46,46,46,46	0
56	MG	1A	3077	1/1	0.82	0.14	69,69,69,69	0
56	MG	2A	3306	1/1	0.82	0.18	48,48,48,48	0
56	MG	1a	1672	1/1	0.82	0.29	58,58,58,58	0
56	MG	1F	307	1/1	0.82	0.13	59,59,59,59	0
56	MG	2A	3533	1/1	0.82	0.16	35,35,35,35	0
56	MG	1a	1677	1/1	0.82	0.13	66,66,66,66	0
56	MG	1A	3440	1/1	0.82	0.14	66,66,66,66	0
56	MG	1a	1780	1/1	0.82	0.16	68,68,68,68	0
56	MG	2A	3039	1/1	0.82	0.14	49,49,49,49	0
56	MG	1A	3822	1/1	0.82	0.14	46,46,46,46	0
56	MG	2a	3198	1/1	0.82	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	1A	3027	1/1	0.82	0.12	54,54,54,54	0
56	MG	2A	3101	1/1	0.82	0.25	57,57,57,57	0
56	MG	1a	1723	1/1	0.82	0.16	85,85,85,85	0
56	MG	2A	3419	1/1	0.82	0.16	56,56,56,56	0
56	MG	2A	3423	1/1	0.82	0.16	66,66,66,66	0
56	MG	2a	3230	1/1	0.82	0.18	50,50,50,50	0
56	MG	2a	3231	1/1	0.82	0.14	75,75,75,75	0
56	MG	1A	3472	1/1	0.82	0.16	83,83,83,83	0
56	MG	1A	3745	1/1	0.82	0.27	71,71,71,71	0
56	MG	2A	3445	1/1	0.82	0.18	71,71,71,71	0
56	MG	1a	1802	1/1	0.82	0.20	76,76,76,76	0
56	MG	2A	3148	1/1	0.82	0.25	63,63,63,63	0
56	MG	1W	3005	1/1	0.82	0.24	49,49,49,49	0
56	MG	1A	3365	1/1	0.82	0.16	46,46,46,46	0
56	MG	2a	3259	1/1	0.82	0.24	56,56,56,56	0
56	MG	2A	3201	1/1	0.82	0.17	47,47,47,47	0
56	MG	1A	3961	1/1	0.82	0.17	60,60,60,60	0
56	MG	1A	3516	1/1	0.83	0.26	53,53,53,53	0
56	MG	2A	3298	1/1	0.83	0.14	49,49,49,49	0
56	MG	2a	3108	1/1	0.83	0.16	62,62,62,62	0
56	MG	1A	3339	1/1	0.83	0.15	56,56,56,56	0
56	MG	1a	1705	1/1	0.83	0.18	73,73,73,73	0
56	MG	1A	3491	1/1	0.83	0.23	61,61,61,61	0
56	MG	1A	3570	1/1	0.83	0.14	66,66,66,66	0
56	MG	2A	3347	1/1	0.83	0.22	60,60,60,60	0
56	MG	2A	3352	1/1	0.83	0.11	68,68,68,68	0
56	MG	2A	3525	1/1	0.83	0.22	63,63,63,63	0
56	MG	1A	3895	1/1	0.83	0.13	24,24,24,24	0
56	MG	2A	3371	1/1	0.83	0.20	56,56,56,56	0
56	MG	1a	1805	1/1	0.83	0.20	67,67,67,67	0
56	MG	1T	203	1/1	0.83	0.14	61,61,61,61	0
56	MG	2A	3124	1/1	0.83	0.27	54,54,54,54	0
56	MG	1a	1846	1/1	0.83	0.17	52,52,52,52	0
56	MG	1A	3909	1/1	0.83	0.13	62,62,62,62	0
56	MG	2A	3411	1/1	0.83	0.32	64,64,64,64	0
56	MG	2F	304	1/1	0.83	0.14	53,53,53,53	0
56	MG	1A	3023	1/1	0.83	0.22	50,50,50,50	0
56	MG	2a	3010	1/1	0.83	0.17	74,74,74,74	0
56	MG	2A	3417	1/1	0.83	0.16	70,70,70,70	0
56	MG	1A	3811	1/1	0.83	0.10	39,39,39,39	0
56	MG	1A	3164	1/1	0.83	0.39	64,64,64,64	0
56	MG	2a	3216	1/1	0.83	0.22	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	3217	1/1	0.83	0.18	83,83,83,83	0
56	MG	1A	3320	1/1	0.83	0.16	93,93,93,93	0
56	MG	1A	3955	1/1	0.83	0.15	63,63,63,63	0
56	MG	1A	3394	1/1	0.83	0.22	56,56,56,56	0
56	MG	1A	3711	1/1	0.83	0.13	48,48,48,48	0
56	MG	1A	3997	1/1	0.83	0.15	50,50,50,50	0
56	MG	1A	3375	1/1	0.83	0.26	51,51,51,51	0
56	MG	2a	3053	1/1	0.83	0.11	85,85,85,85	0
56	MG	1a	1663	1/1	0.83	0.11	76,76,76,76	0
56	MG	2A	3257	1/1	0.83	0.17	62,62,62,62	0
56	MG	1A	3445	1/1	0.83	0.16	63,63,63,63	0
56	MG	1A	3066	1/1	0.83	0.18	53,53,53,53	0
56	MG	2k	8001	1/1	0.83	0.12	83,83,83,83	0
56	MG	2A	3011	1/1	0.83	0.16	61,61,61,61	0
56	MG	1B	223	1/1	0.83	0.09	59,59,59,59	0
56	MG	1a	1618	1/1	0.84	0.28	64,64,64,64	0
56	MG	1A	3597	1/1	0.84	0.21	48,48,48,48	0
56	MG	1A	3617	1/1	0.84	0.13	27,27,27,27	0
56	MG	1a	1776	1/1	0.84	0.16	69,69,69,69	0
56	MG	2A	3080	1/1	0.84	0.22	48,48,48,48	0
56	MG	1A	3833	1/1	0.84	0.28	78,78,78,78	0
56	MG	2A	3111	1/1	0.84	0.23	68,68,68,68	0
56	MG	1A	3633	1/1	0.84	0.22	71,71,71,71	0
56	MG	1A	3243	1/1	0.84	0.28	46,46,46,46	0
56	MG	1A	3049	1/1	0.84	0.14	48,48,48,48	0
56	MG	1A	3189	1/1	0.84	0.15	39,39,39,39	0
56	MG	1a	1683	1/1	0.84	0.23	68,68,68,68	0
56	MG	2a	3080	1/1	0.84	0.13	40,40,40,40	0
56	MG	1D	301	1/1	0.84	0.12	56,56,56,56	0
56	MG	1a	1803	1/1	0.84	0.18	49,49,49,49	0
56	MG	1F	304	1/1	0.84	0.15	43,43,43,43	0
56	MG	2a	3116	1/1	0.84	0.20	62,62,62,62	0
56	MG	1a	1818	1/1	0.84	0.21	63,63,63,63	0
56	MG	1a	1834	1/1	0.84	0.24	63,63,63,63	0
56	MG	2A	3489	1/1	0.84	0.13	63,63,63,63	0
56	MG	1A	3682	1/1	0.84	0.14	51,51,51,51	0
56	MG	2A	3213	1/1	0.84	0.29	59,59,59,59	0
56	MG	1a	1844	1/1	0.84	0.13	43,43,43,43	0
56	MG	1A	3419	1/1	0.84	0.13	56,56,56,56	0
56	MG	1A	3427	1/1	0.84	0.15	53,53,53,53	0
56	MG	1A	3334	1/1	0.84	0.18	72,72,72,72	0
56	MG	1A	3338	1/1	0.84	0.19	54,54,54,54	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1737	1/1	0.84	0.22	57,57,57,57	0
56	MG	1A	3367	1/1	0.84	0.18	66,66,66,66	0
56	MG	1A	3479	1/1	0.84	0.22	38,38,38,38	0
56	MG	1T	205	1/1	0.84	0.12	65,65,65,65	0
56	MG	2a	3192	1/1	0.84	0.16	60,60,60,60	0
56	MG	1A	3924	1/1	0.84	0.14	66,66,66,66	0
56	MG	2A	3287	1/1	0.84	0.10	56,56,56,56	0
56	MG	1A	3480	1/1	0.84	0.20	56,56,56,56	0
56	MG	1A	3166	1/1	0.84	0.11	55,55,55,55	0
56	MG	1a	1757	1/1	0.84	0.15	63,63,63,63	0
56	MG	2A	3318	1/1	0.84	0.09	47,47,47,47	0
56	MG	2A	3557	1/1	0.84	0.18	64,64,64,64	0
56	MG	2B	3002	1/1	0.84	0.15	61,61,61,61	0
56	MG	2B	3003	1/1	0.84	0.20	73,73,73,73	0
56	MG	1a	1604	1/1	0.84	0.18	69,69,69,69	0
56	MG	1o	3001	1/1	0.84	0.20	74,74,74,74	0
56	MG	1z	101	1/1	0.84	0.23	76,76,76,76	0
56	MG	1A	3523	1/1	0.84	0.17	59,59,59,59	0
56	MG	2A	3357	1/1	0.84	0.24	65,65,65,65	0
56	MG	1A	3325	1/1	0.84	0.20	74,74,74,74	0
56	MG	2a	3244	1/1	0.84	0.45	67,67,67,67	0
56	MG	2a	3011	1/1	0.84	0.22	67,67,67,67	0
56	MG	2A	3365	1/1	0.84	0.14	63,63,63,63	0
56	MG	2A	3367	1/1	0.84	0.15	50,50,50,50	0
56	MG	1a	1768	1/1	0.84	0.21	72,72,72,72	0
56	MG	2a	3263	1/1	0.84	0.32	71,71,71,71	0
56	MG	2A	3016	1/1	0.84	0.23	57,57,57,57	0
56	MG	2a	3026	1/1	0.84	0.19	64,64,64,64	0
56	MG	1A	3581	1/1	0.84	0.10	37,37,37,37	0
56	MG	1A	3461	1/1	0.85	0.12	71,71,71,71	0
56	MG	2A	3113	1/1	0.85	0.25	64,64,64,64	0
56	MG	1A	3665	1/1	0.85	0.13	22,22,22,22	0
56	MG	1a	1643	1/1	0.85	0.35	76,76,76,76	0
56	MG	1A	4000	1/1	0.85	0.16	61,61,61,61	0
56	MG	2A	3427	1/1	0.85	0.19	56,56,56,56	0
56	MG	2A	3129	1/1	0.85	0.26	49,49,49,49	0
56	MG	2a	3063	1/1	0.85	0.42	77,77,77,77	0
56	MG	1a	1668	1/1	0.85	0.13	59,59,59,59	0
56	MG	1A	4011	1/1	0.85	0.26	47,47,47,47	0
56	MG	2A	3449	1/1	0.85	0.08	60,60,60,60	0
56	MG	2a	3072	1/1	0.85	0.35	77,77,77,77	0
56	MG	2a	3074	1/1	0.85	0.20	74,74,74,74	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3452	1/1	0.85	0.17	69,69,69,69	0
56	MG	1A	4017	1/1	0.85	0.31	60,60,60,60	0
56	MG	1A	3173	1/1	0.85	0.22	62,62,62,62	0
56	MG	2A	3465	1/1	0.85	0.13	31,31,31,31	0
56	MG	1A	3033	1/1	0.85	0.12	57,57,57,57	0
56	MG	2A	3474	1/1	0.85	0.18	41,41,41,41	0
56	MG	1A	3343	1/1	0.85	0.16	77,77,77,77	0
56	MG	1A	3847	1/1	0.85	0.14	46,46,46,46	0
56	MG	1D	304	1/1	0.85	0.13	58,58,58,58	0
56	MG	1A	3683	1/1	0.85	0.14	58,58,58,58	0
56	MG	2A	3217	1/1	0.85	0.24	56,56,56,56	0
56	MG	1a	1849	1/1	0.85	0.11	53,53,53,53	0
56	MG	1A	3059	1/1	0.85	0.13	46,46,46,46	0
56	MG	1A	3128	1/1	0.85	0.16	56,56,56,56	0
56	MG	1A	3721	1/1	0.85	0.13	55,55,55,55	0
56	MG	1A	3262	1/1	0.85	0.12	66,66,66,66	0
56	MG	2a	3163	1/1	0.85	0.16	70,70,70,70	0
56	MG	1a	1738	1/1	0.85	0.31	64,64,64,64	0
56	MG	2a	3168	1/1	0.85	0.15	71,71,71,71	0
56	MG	1A	3414	1/1	0.85	0.14	65,65,65,65	0
56	MG	1A	3751	1/1	0.85	0.12	39,39,39,39	0
56	MG	2a	3175	1/1	0.85	0.15	62,62,62,62	0
56	MG	1A	3558	1/1	0.85	0.09	61,61,61,61	0
56	MG	1A	3489	1/1	0.85	0.08	51,51,51,51	0
56	MG	1a	1753	1/1	0.85	0.12	70,70,70,70	0
56	MG	2A	3292	1/1	0.85	0.17	42,42,42,42	0
56	MG	2a	3204	1/1	0.85	0.26	64,64,64,64	0
56	MG	1A	3285	1/1	0.85	0.20	54,54,54,54	0
56	MG	1A	3583	1/1	0.85	0.13	55,55,55,55	0
56	MG	12	3001	1/1	0.85	0.16	61,61,61,61	0
56	MG	2a	3213	1/1	0.85	0.13	67,67,67,67	0
56	MG	1A	3418	1/1	0.85	0.12	66,66,66,66	0
56	MG	2A	3319	1/1	0.85	0.15	58,58,58,58	0
56	MG	1A	3607	1/1	0.85	0.15	40,40,40,40	0
56	MG	1A	3949	1/1	0.85	0.12	48,48,48,48	0
56	MG	2a	3227	1/1	0.85	0.17	66,66,66,66	0
56	MG	1A	3953	1/1	0.85	0.18	71,71,71,71	0
56	MG	1A	3001	1/1	0.85	0.30	42,42,42,42	0
56	MG	2a	3233	1/1	0.85	0.22	61,61,61,61	0
56	MG	1a	1772	1/1	0.85	0.08	78,78,78,78	0
56	MG	1a	1617	1/1	0.85	0.13	78,78,78,78	0
56	MG	2A	3363	1/1	0.85	0.18	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	2a	3008	1/1	0.85	0.34	67,67,67,67	0
56	MG	2A	3364	1/1	0.85	0.14	60,60,60,60	0
56	MG	1A	3815	1/1	0.85	0.23	67,67,67,67	0
56	MG	1A	3425	1/1	0.85	0.15	63,63,63,63	0
56	MG	1a	1624	1/1	0.85	0.15	62,62,62,62	0
56	MG	2A	3074	1/1	0.85	0.25	57,57,57,57	0
56	MG	1a	1626	1/1	0.85	0.19	80,80,80,80	0
56	MG	2A	3085	1/1	0.85	0.14	53,53,53,53	0
56	MG	2A	3092	1/1	0.85	0.15	54,54,54,54	0
56	MG	1a	1786	1/1	0.85	0.15	92,92,92,92	0
56	MG	2a	3050	1/1	0.86	0.14	93,93,93,93	0
56	MG	1a	1690	1/1	0.86	0.15	63,63,63,63	0
56	MG	1A	3929	1/1	0.86	0.13	58,58,58,58	0
56	MG	1a	1699	1/1	0.86	0.22	71,71,71,71	0
56	MG	2A	3424	1/1	0.86	0.31	80,80,80,80	0
56	MG	1a	1819	1/1	0.86	0.09	54,54,54,54	0
56	MG	2A	3436	1/1	0.86	0.32	58,58,58,58	0
56	MG	1A	3144	1/1	0.86	0.16	54,54,54,54	0
56	MG	1A	3396	1/1	0.86	0.22	67,67,67,67	0
56	MG	1A	3120	1/1	0.86	0.14	41,41,41,41	0
56	MG	1A	3172	1/1	0.86	0.26	44,44,44,44	0
56	MG	1A	3518	1/1	0.86	0.21	53,53,53,53	0
56	MG	2a	3073	1/1	0.86	0.19	61,61,61,61	0
56	MG	15	101	1/1	0.86	0.25	58,58,58,58	0
56	MG	1A	3024	1/1	0.86	0.10	63,63,63,63	0
56	MG	1a	1601	1/1	0.86	0.12	53,53,53,53	0
56	MG	1A	3078	1/1	0.86	0.09	62,62,62,62	0
56	MG	2a	3099	1/1	0.86	0.07	35,35,35,35	0
56	MG	1a	1606	1/1	0.86	0.21	76,76,76,76	0
56	MG	2A	3481	1/1	0.86	0.19	50,50,50,50	0
56	MG	2A	3483	1/1	0.86	0.15	54,54,54,54	0
56	MG	1A	3454	1/1	0.86	0.26	66,66,66,66	0
56	MG	2A	3233	1/1	0.86	0.27	48,48,48,48	0
56	MG	1A	3545	1/1	0.86	0.15	17,17,17,17	0
56	MG	2a	3147	1/1	0.86	0.19	73,73,73,73	0
56	MG	2A	3253	1/1	0.86	0.14	43,43,43,43	0
56	MG	2A	3490	1/1	0.86	0.11	58,58,58,58	0
56	MG	1A	3201	1/1	0.86	0.16	49,49,49,49	0
56	MG	1a	1875	1/1	0.86	0.11	82,82,82,82	0
56	MG	1A	3560	1/1	0.86	0.13	45,45,45,45	0
56	MG	1a	1756	1/1	0.86	0.19	75,75,75,75	0
56	MG	1A	3035	1/1	0.86	0.18	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	3241	1/1	0.86	0.14	42,42,42,42	0
56	MG	1A	3133	1/1	0.86	0.17	59,59,59,59	0
56	MG	2A	3289	1/1	0.86	0.15	58,58,58,58	0
56	MG	1A	3881	1/1	0.86	0.17	55,55,55,55	0
56	MG	2A	3002	1/1	0.86	0.17	47,47,47,47	0
56	MG	1a	1767	1/1	0.86	0.16	64,64,64,64	0
56	MG	2a	3180	1/1	0.86	0.18	61,61,61,61	0
56	MG	2A	3008	1/1	0.86	0.15	64,64,64,64	0
56	MG	1A	3883	1/1	0.86	0.27	58,58,58,58	0
56	MG	1a	1628	1/1	0.86	0.25	61,61,61,61	0
56	MG	1A	3363	1/1	0.86	0.18	68,68,68,68	0
56	MG	1A	3894	1/1	0.86	0.12	41,41,41,41	0
56	MG	2A	3334	1/1	0.86	0.10	58,58,58,58	0
56	MG	1a	1646	1/1	0.86	0.14	57,57,57,57	0
56	MG	1a	1657	1/1	0.86	0.13	71,71,71,71	0
56	MG	2A	3355	1/1	0.86	0.14	73,73,73,73	0
56	MG	2A	3048	1/1	0.86	0.26	56,56,56,56	0
56	MG	2A	3054	1/1	0.86	0.18	49,49,49,49	0
56	MG	2B	3011	1/1	0.86	0.24	83,83,83,83	0
56	MG	1a	1658	1/1	0.86	0.09	67,67,67,67	0
56	MG	2a	3225	1/1	0.86	0.16	59,59,59,59	0
56	MG	1A	3463	1/1	0.86	0.08	80,80,80,80	0
56	MG	1a	1781	1/1	0.86	0.14	46,46,46,46	0
56	MG	2a	3003	1/1	0.86	0.12	57,57,57,57	0
56	MG	1A	3614	1/1	0.86	0.17	37,37,37,37	0
56	MG	1A	3918	1/1	0.86	0.12	71,71,71,71	0
56	MG	1a	1792	1/1	0.86	0.12	54,54,54,54	0
56	MG	1A	3923	1/1	0.86	0.20	53,53,53,53	0
56	MG	1A	3508	1/1	0.86	0.15	57,57,57,57	0
56	MG	2A	3390	1/1	0.86	0.30	56,56,56,56	0
56	MG	2a	3248	1/1	0.86	0.20	65,65,65,65	0
56	MG	2a	3015	1/1	0.86	0.37	67,67,67,67	0
56	MG	1a	1680	1/1	0.86	0.12	53,53,53,53	0
56	MG	2A	3114	1/1	0.86	0.11	57,57,57,57	0
56	MG	2A	3409	1/1	0.86	0.09	50,50,50,50	0
56	MG	2a	3261	1/1	0.86	0.37	74,74,74,74	0
56	MG	1Q	3003	1/1	0.86	0.34	51,51,51,51	0
56	MG	2a	3041	1/1	0.86	0.21	73,73,73,73	0
56	MG	2a	3043	1/1	0.86	0.42	65,65,65,65	0
56	MG	1A	3619	1/1	0.86	0.12	48,48,48,48	0
56	MG	2A	3342	1/1	0.87	0.11	63,63,63,63	0
56	MG	1A	3358	1/1	0.87	0.23	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	3197	1/1	0.87	0.20	50,50,50,50	0
56	MG	2a	3100	1/1	0.87	0.30	66,66,66,66	0
56	MG	1Z	8002	1/1	0.87	0.12	67,67,67,67	0
56	MG	1A	3437	1/1	0.87	0.23	62,62,62,62	0
56	MG	2A	3522	1/1	0.87	0.15	70,70,70,70	0
56	MG	1A	3462	1/1	0.87	0.09	57,57,57,57	0
56	MG	2a	3125	1/1	0.87	0.32	60,60,60,60	0
56	MG	1A	3362	1/1	0.87	0.10	56,56,56,56	0
56	MG	1A	3537	1/1	0.87	0.14	32,32,32,32	0
56	MG	1a	1873	1/1	0.87	0.13	52,52,52,52	0
56	MG	1A	3141	1/1	0.87	0.30	66,66,66,66	0
56	MG	2A	3146	1/1	0.87	0.17	64,64,64,64	0
56	MG	2A	3374	1/1	0.87	0.21	37,37,37,37	0
56	MG	2a	3155	1/1	0.87	0.17	79,79,79,79	0
56	MG	2A	3376	1/1	0.87	0.17	47,47,47,47	0
56	MG	1A	3549	1/1	0.87	0.16	48,48,48,48	0
56	MG	1a	1704	1/1	0.87	0.12	54,54,54,54	0
56	MG	2A	3182	1/1	0.87	0.10	52,52,52,52	0
56	MG	1A	4013	1/1	0.87	0.14	49,49,49,49	0
56	MG	2A	3197	1/1	0.87	0.20	57,57,57,57	0
56	MG	2D	306	1/1	0.87	0.18	68,68,68,68	0
56	MG	1A	3884	1/1	0.87	0.10	53,53,53,53	0
56	MG	1a	1790	1/1	0.87	0.14	71,71,71,71	0
56	MG	2a	3172	1/1	0.87	0.26	60,60,60,60	0
56	MG	1r	3001	1/1	0.87	0.16	69,69,69,69	0
56	MG	2P	201	1/1	0.87	0.17	41,41,41,41	0
56	MG	26	101	1/1	0.87	0.15	57,57,57,57	0
56	MG	1A	3554	1/1	0.87	0.21	80,80,80,80	0
56	MG	1A	3808	1/1	0.87	0.23	51,51,51,51	0
56	MG	2a	3196	1/1	0.87	0.14	90,90,90,90	0
56	MG	1a	1731	1/1	0.87	0.28	57,57,57,57	0
56	MG	2A	3224	1/1	0.87	0.07	42,42,42,42	0
56	MG	1A	3448	1/1	0.87	0.07	49,49,49,49	0
56	MG	1A	3342	1/1	0.87	0.17	52,52,52,52	0
56	MG	1A	3816	1/1	0.87	0.13	68,68,68,68	0
56	MG	2A	3440	1/1	0.87	0.27	60,60,60,60	0
56	MG	2A	3015	1/1	0.87	0.19	52,52,52,52	0
56	MG	2A	3245	1/1	0.87	0.25	48,48,48,48	0
56	MG	1F	302	1/1	0.87	0.27	51,51,51,51	0
56	MG	1A	3818	1/1	0.87	0.16	61,61,61,61	0
56	MG	1A	3380	1/1	0.87	0.09	63,63,63,63	0
56	MG	1a	1813	1/1	0.87	0.19	59,59,59,59	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2a	3044	1/1	0.87	0.28	71,71,71,71	0
56	MG	2A	3044	1/1	0.87	0.20	48,48,48,48	0
56	MG	2A	3046	1/1	0.87	0.30	64,64,64,64	0
56	MG	1G	3002	1/1	0.87	0.18	61,61,61,61	0
56	MG	2A	3282	1/1	0.87	0.12	48,48,48,48	0
56	MG	1A	3823	1/1	0.87	0.10	43,43,43,43	0
56	MG	2a	3240	1/1	0.87	0.23	81,81,81,81	0
56	MG	1a	1823	1/1	0.87	0.10	69,69,69,69	0
56	MG	2a	3056	1/1	0.87	0.16	85,85,85,85	0
56	MG	1A	3697	1/1	0.87	0.17	58,58,58,58	0
56	MG	2a	3253	1/1	0.87	0.11	56,56,56,56	0
56	MG	2A	3295	1/1	0.87	0.13	48,48,48,48	0
56	MG	1A	3931	1/1	0.87	0.26	33,33,33,33	0
56	MG	1A	3034	1/1	0.87	0.28	73,73,73,73	0
56	MG	2A	3084	1/1	0.87	0.26	54,54,54,54	0
56	MG	1A	3707	1/1	0.87	0.18	70,70,70,70	0
56	MG	2A	3087	1/1	0.87	0.24	59,59,59,59	0
56	MG	1A	3431	1/1	0.87	0.19	64,64,64,64	0
56	MG	1A	3592	1/1	0.87	0.17	47,47,47,47	0
56	MG	2A	3103	1/1	0.87	0.16	53,53,53,53	0
56	MG	2A	3448	1/1	0.88	0.18	56,56,56,56	0
56	MG	2A	3239	1/1	0.88	0.12	60,60,60,60	0
56	MG	2A	3031	1/1	0.88	0.19	51,51,51,51	0
56	MG	1a	1702	1/1	0.88	0.35	82,82,82,82	0
56	MG	2A	3251	1/1	0.88	0.17	53,53,53,53	0
56	MG	2A	3252	1/1	0.88	0.15	60,60,60,60	0
56	MG	1a	1703	1/1	0.88	0.25	62,62,62,62	0
56	MG	2A	3038	1/1	0.88	0.19	58,58,58,58	0
56	MG	16	103	1/1	0.88	0.10	68,68,68,68	0
56	MG	2A	3040	1/1	0.88	0.11	53,53,53,53	0
56	MG	2a	3077	1/1	0.88	0.22	59,59,59,59	0
56	MG	1A	3722	1/1	0.88	0.12	22,22,22,22	0
56	MG	1A	3070	1/1	0.88	0.17	72,72,72,72	0
56	MG	1a	1721	1/1	0.88	0.27	59,59,59,59	0
56	MG	2A	3276	1/1	0.88	0.10	42,42,42,42	0
56	MG	2A	3278	1/1	0.88	0.10	31,31,31,31	0
56	MG	2a	3111	1/1	0.88	0.23	56,56,56,56	0
56	MG	2A	3281	1/1	0.88	0.17	59,59,59,59	0
56	MG	1a	1603	1/1	0.88	0.16	57,57,57,57	0
56	MG	2A	3284	1/1	0.88	0.14	62,62,62,62	0
56	MG	1A	4021	1/1	0.88	0.18	57,57,57,57	0
56	MG	1A	3634	1/1	0.88	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	2A	3073	1/1	0.88	0.25	55,55,55,55	0
56	MG	1A	3300	1/1	0.88	0.31	54,54,54,54	0
56	MG	2a	3148	1/1	0.88	0.10	72,72,72,72	0
56	MG	1B	221	1/1	0.88	0.11	68,68,68,68	0
56	MG	1B	222	1/1	0.88	0.16	70,70,70,70	0
56	MG	2A	3512	1/1	0.88	0.10	44,44,44,44	0
56	MG	2A	3311	1/1	0.88	0.19	42,42,42,42	0
56	MG	2A	3519	1/1	0.88	0.12	45,45,45,45	0
56	MG	1a	1742	1/1	0.88	0.19	51,51,51,51	0
56	MG	2A	3523	1/1	0.88	0.13	56,56,56,56	0
56	MG	1A	3575	1/1	0.88	0.17	41,41,41,41	0
56	MG	2a	3161	1/1	0.88	0.10	102,102,102,102	0
56	MG	2A	3091	1/1	0.88	0.17	49,49,49,49	0
56	MG	2A	3526	1/1	0.88	0.18	49,49,49,49	0
56	MG	1A	3497	1/1	0.88	0.15	57,57,57,57	0
56	MG	1A	3671	1/1	0.88	0.20	56,56,56,56	0
56	MG	1A	3018	1/1	0.88	0.14	59,59,59,59	0
56	MG	2A	3106	1/1	0.88	0.25	59,59,59,59	0
56	MG	2a	3173	1/1	0.88	0.12	74,74,74,74	0
56	MG	2A	3543	1/1	0.88	0.16	56,56,56,56	0
56	MG	2A	3346	1/1	0.88	0.18	62,62,62,62	0
56	MG	2a	3185	1/1	0.88	0.10	67,67,67,67	0
56	MG	1A	3933	1/1	0.88	0.09	42,42,42,42	0
56	MG	1A	3456	1/1	0.88	0.28	39,39,39,39	0
56	MG	1A	3938	1/1	0.88	0.14	44,44,44,44	0
56	MG	1H	8001	1/1	0.88	0.09	58,58,58,58	0
56	MG	1A	3067	1/1	0.88	0.12	46,46,46,46	0
56	MG	1a	1759	1/1	0.88	0.13	70,70,70,70	0
56	MG	1A	3941	1/1	0.88	0.21	65,65,65,65	0
56	MG	2A	3133	1/1	0.88	0.09	52,52,52,52	0
56	MG	2A	3136	1/1	0.88	0.19	54,54,54,54	0
56	MG	1a	1654	1/1	0.88	0.21	78,78,78,78	0
56	MG	1a	1655	1/1	0.88	0.09	56,56,56,56	0
56	MG	1A	3943	1/1	0.88	0.13	68,68,68,68	0
56	MG	1A	3871	1/1	0.88	0.26	60,60,60,60	0
56	MG	2a	3005	1/1	0.88	0.23	62,62,62,62	0
56	MG	1A	3603	1/1	0.88	0.15	60,60,60,60	0
56	MG	2A	3179	1/1	0.88	0.16	46,46,46,46	0
56	MG	1a	1665	1/1	0.88	0.14	61,61,61,61	0
56	MG	1A	3360	1/1	0.88	0.12	47,47,47,47	0
56	MG	2A	3194	1/1	0.88	0.15	44,44,44,44	0
56	MG	1A	3880	1/1	0.88	0.16	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	1t	3001	1/1	0.88	0.12	54,54,54,54	0
56	MG	2a	3237	1/1	0.88	0.19	52,52,52,52	0
56	MG	2A	3205	1/1	0.88	0.30	51,51,51,51	0
56	MG	2A	3416	1/1	0.88	0.18	48,48,48,48	0
56	MG	2a	3029	1/1	0.88	0.20	69,69,69,69	0
56	MG	1x	105	1/1	0.88	0.24	57,57,57,57	0
56	MG	1A	3442	1/1	0.88	0.20	58,58,58,58	0
56	MG	2a	3250	1/1	0.88	0.17	61,61,61,61	0
56	MG	1A	3709	1/1	0.88	0.14	45,45,45,45	0
56	MG	1Z	8001	1/1	0.88	0.09	69,69,69,69	0
56	MG	2A	3425	1/1	0.88	0.10	47,47,47,47	0
56	MG	1A	3099	1/1	0.88	0.10	39,39,39,39	0
56	MG	2a	3258	1/1	0.88	0.08	69,69,69,69	0
56	MG	1A	3819	1/1	0.88	0.10	82,82,82,82	0
56	MG	2A	3227	1/1	0.88	0.10	47,47,47,47	0
56	MG	10	106	1/1	0.88	0.15	61,61,61,61	0
56	MG	1A	3891	1/1	0.88	0.15	62,62,62,62	0
56	MG	1A	3410	1/1	0.88	0.14	49,49,49,49	0
56	MG	2q	202	1/1	0.88	0.27	69,69,69,69	0
56	MG	2A	3447	1/1	0.88	0.10	62,62,62,62	0
56	MG	1A	3268	1/1	0.89	0.08	55,55,55,55	0
56	MG	2A	3383	1/1	0.89	0.17	66,66,66,66	0
56	MG	2a	3036	1/1	0.89	0.12	72,72,72,72	0
56	MG	2A	3387	1/1	0.89	0.26	63,63,63,63	0
56	MG	2A	3122	1/1	0.89	0.21	66,66,66,66	0
56	MG	1a	1814	1/1	0.89	0.18	67,67,67,67	0
56	MG	2A	3395	1/1	0.89	0.10	33,33,33,33	0
56	MG	2A	3396	1/1	0.89	0.23	40,40,40,40	0
56	MG	1a	1685	1/1	0.89	0.14	53,53,53,53	0
56	MG	2A	3131	1/1	0.89	0.24	52,52,52,52	0
56	MG	2A	3403	1/1	0.89	0.23	62,62,62,62	0
56	MG	2A	3405	1/1	0.89	0.20	63,63,63,63	0
56	MG	1A	3407	1/1	0.89	0.09	41,41,41,41	0
56	MG	1A	3055	1/1	0.89	0.39	59,59,59,59	0
56	MG	2A	3137	1/1	0.89	0.30	51,51,51,51	0
56	MG	1a	1832	1/1	0.89	0.10	50,50,50,50	0
56	MG	2A	3144	1/1	0.89	0.30	51,51,51,51	0
56	MG	1A	3299	1/1	0.89	0.17	61,61,61,61	0
56	MG	2A	3421	1/1	0.89	0.14	59,59,59,59	0
56	MG	2a	3070	1/1	0.89	0.28	69,69,69,69	0
56	MG	1a	1696	1/1	0.89	0.22	70,70,70,70	0
56	MG	1P	201	1/1	0.89	0.13	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	2A	3158	1/1	0.89	0.17	47,47,47,47	0
56	MG	1a	1701	1/1	0.89	0.17	54,54,54,54	0
56	MG	1A	3784	1/1	0.89	0.14	33,33,33,33	0
56	MG	1A	3416	1/1	0.89	0.17	60,60,60,60	0
56	MG	1A	3081	1/1	0.89	0.12	58,58,58,58	0
56	MG	1A	3794	1/1	0.89	0.16	68,68,68,68	0
56	MG	1a	1709	1/1	0.89	0.22	43,43,43,43	0
56	MG	2a	3101	1/1	0.89	0.25	53,53,53,53	0
56	MG	1A	3083	1/1	0.89	0.09	56,56,56,56	0
56	MG	1U	201	1/1	0.89	0.12	51,51,51,51	0
56	MG	1A	3058	1/1	0.89	0.15	50,50,50,50	0
56	MG	1A	3481	1/1	0.89	0.35	40,40,40,40	0
56	MG	2a	3122	1/1	0.89	0.20	72,72,72,72	0
56	MG	1a	1725	1/1	0.89	0.22	59,59,59,59	0
56	MG	1A	3006	1/1	0.89	0.17	57,57,57,57	0
56	MG	2a	3134	1/1	0.89	0.40	59,59,59,59	0
56	MG	1A	3612	1/1	0.89	0.12	55,55,55,55	0
56	MG	1b	3001	1/1	0.89	0.13	83,83,83,83	0
56	MG	2A	3226	1/1	0.89	0.10	57,57,57,57	0
56	MG	2A	3479	1/1	0.89	0.07	50,50,50,50	0
56	MG	1A	3190	1/1	0.89	0.22	44,44,44,44	0
56	MG	1d	502	1/1	0.89	0.22	76,76,76,76	0
56	MG	1A	3115	1/1	0.89	0.10	49,49,49,49	0
56	MG	1a	1739	1/1	0.89	0.33	46,46,46,46	0
56	MG	1A	3820	1/1	0.89	0.12	57,57,57,57	0
56	MG	2a	3157	1/1	0.89	0.21	74,74,74,74	0
56	MG	1q	201	1/1	0.89	0.26	51,51,51,51	0
56	MG	1q	204	1/1	0.89	0.13	52,52,52,52	0
56	MG	2A	3491	1/1	0.89	0.13	47,47,47,47	0
56	MG	2A	3250	1/1	0.89	0.12	64,64,64,64	0
56	MG	1A	3119	1/1	0.89	0.20	65,65,65,65	0
56	MG	1A	3214	1/1	0.89	0.20	60,60,60,60	0
56	MG	1A	3826	1/1	0.89	0.13	37,37,37,37	0
56	MG	2A	3502	1/1	0.89	0.07	35,35,35,35	0
56	MG	2A	3504	1/1	0.89	0.17	56,56,56,56	0
56	MG	1a	1602	1/1	0.89	0.12	64,64,64,64	0
56	MG	2A	3001	1/1	0.89	0.09	45,45,45,45	0
56	MG	1A	3215	1/1	0.89	0.18	51,51,51,51	0
56	MG	1A	3987	1/1	0.89	0.15	38,38,38,38	0
56	MG	1A	3988	1/1	0.89	0.25	45,45,45,45	0
56	MG	1A	3637	1/1	0.89	0.07	47,47,47,47	0
56	MG	2A	3275	1/1	0.89	0.17	60,60,60,60	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3436	1/1	0.89	0.14	61,61,61,61	0
56	MG	1A	3379	1/1	0.89	0.11	61,61,61,61	0
56	MG	1A	3220	1/1	0.89	0.21	60,60,60,60	0
56	MG	2a	3199	1/1	0.89	0.23	62,62,62,62	0
56	MG	2a	3201	1/1	0.89	0.10	88,88,88,88	0
56	MG	2a	3202	1/1	0.89	0.21	72,72,72,72	0
56	MG	1A	4001	1/1	0.89	0.43	43,43,43,43	0
56	MG	1a	1762	1/1	0.89	0.17	89,89,89,89	0
56	MG	2A	3285	1/1	0.89	0.20	39,39,39,39	0
56	MG	2a	3209	1/1	0.89	0.22	70,70,70,70	0
56	MG	1A	3441	1/1	0.89	0.13	72,72,72,72	0
56	MG	1A	3074	1/1	0.89	0.32	51,51,51,51	0
56	MG	1A	3860	1/1	0.89	0.14	29,29,29,29	0
56	MG	1A	4018	1/1	0.89	0.36	50,50,50,50	0
56	MG	1a	1770	1/1	0.89	0.14	67,67,67,67	0
56	MG	2A	3045	1/1	0.89	0.11	48,48,48,48	0
56	MG	1A	3238	1/1	0.89	0.24	45,45,45,45	0
56	MG	1a	1633	1/1	0.89	0.13	38,38,38,38	0
56	MG	2a	3228	1/1	0.89	0.11	69,69,69,69	0
56	MG	1a	1638	1/1	0.89	0.19	63,63,63,63	0
56	MG	1B	209	1/1	0.89	0.26	62,62,62,62	0
56	MG	1B	211	1/1	0.89	0.15	51,51,51,51	0
56	MG	1A	3865	1/1	0.89	0.14	68,68,68,68	0
56	MG	1a	1649	1/1	0.89	0.16	63,63,63,63	0
56	MG	2A	3076	1/1	0.89	0.24	60,60,60,60	0
56	MG	2A	3079	1/1	0.89	0.35	52,52,52,52	0
56	MG	1A	3387	1/1	0.89	0.26	56,56,56,56	0
56	MG	1A	3450	1/1	0.89	0.20	45,45,45,45	0
56	MG	2Y	502	1/1	0.89	0.20	52,52,52,52	0
56	MG	2a	3245	1/1	0.89	0.28	62,62,62,62	0
56	MG	2a	3247	1/1	0.89	0.10	71,71,71,71	0
56	MG	2Z	3001	1/1	0.89	0.11	68,68,68,68	0
56	MG	1A	3390	1/1	0.89	0.32	40,40,40,40	0
56	MG	1A	3161	1/1	0.89	0.06	33,33,33,33	0
56	MG	1A	3125	1/1	0.89	0.12	60,60,60,60	0
56	MG	1A	3246	1/1	0.89	0.14	58,58,58,58	0
56	MG	1a	1797	1/1	0.89	0.11	60,60,60,60	0
56	MG	1D	313	1/1	0.89	0.15	56,56,56,56	0
56	MG	2A	3366	1/1	0.89	0.15	69,69,69,69	0
56	MG	1a	1800	1/1	0.89	0.13	57,57,57,57	0
56	MG	1A	3457	1/1	0.89	0.10	72,72,72,72	0
56	MG	2A	3372	1/1	0.89	0.23	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	1A	3256	1/1	0.89	0.22	43,43,43,43	0
56	MG	1A	3065	1/1	0.89	0.18	49,49,49,49	0
56	MG	1A	3404	1/1	0.89	0.10	48,48,48,48	0
56	MG	2a	3027	1/1	0.90	0.15	70,70,70,70	0
56	MG	1A	3824	1/1	0.90	0.09	33,33,33,33	0
56	MG	2a	3030	1/1	0.90	0.33	73,73,73,73	0
56	MG	1W	3003	1/1	0.90	0.17	35,35,35,35	0
56	MG	2a	3032	1/1	0.90	0.14	50,50,50,50	0
56	MG	2a	3034	1/1	0.90	0.14	59,59,59,59	0
56	MG	2A	3398	1/1	0.90	0.16	50,50,50,50	0
56	MG	1a	1711	1/1	0.90	0.22	49,49,49,49	0
56	MG	1A	3691	1/1	0.90	0.10	51,51,51,51	0
56	MG	2a	3042	1/1	0.90	0.17	56,56,56,56	0
56	MG	2A	3402	1/1	0.90	0.16	62,62,62,62	0
56	MG	1A	3828	1/1	0.90	0.13	67,67,67,67	0
56	MG	1a	1861	1/1	0.90	0.16	58,58,58,58	0
56	MG	1A	3346	1/1	0.90	0.13	63,63,63,63	0
56	MG	1A	3832	1/1	0.90	0.23	57,57,57,57	0
56	MG	1A	3703	1/1	0.90	0.20	49,49,49,49	0
56	MG	1a	1868	1/1	0.90	0.30	66,66,66,66	0
56	MG	1A	3043	1/1	0.90	0.12	52,52,52,52	0
56	MG	1A	3971	1/1	0.90	0.11	62,62,62,62	0
56	MG	2a	3057	1/1	0.90	0.07	96,96,96,96	0
56	MG	2A	3204	1/1	0.90	0.12	44,44,44,44	0
56	MG	15	105	1/1	0.90	0.10	49,49,49,49	0
56	MG	1A	3978	1/1	0.90	0.07	29,29,29,29	0
56	MG	17	101	1/1	0.90	0.24	57,57,57,57	0
56	MG	2A	3426	1/1	0.90	0.18	62,62,62,62	0
56	MG	1a	1741	1/1	0.90	0.21	63,63,63,63	0
56	MG	1A	3312	1/1	0.90	0.15	38,38,38,38	0
56	MG	2A	3437	1/1	0.90	0.07	57,57,57,57	0
56	MG	2A	3438	1/1	0.90	0.14	62,62,62,62	0
56	MG	1e	3001	1/1	0.90	0.14	69,69,69,69	0
56	MG	1e	3002	1/1	0.90	0.11	83,83,83,83	0
56	MG	2A	3442	1/1	0.90	0.12	50,50,50,50	0
56	MG	2A	3225	1/1	0.90	0.27	63,63,63,63	0
56	MG	2a	3086	1/1	0.90	0.29	64,64,64,64	0
56	MG	1a	1743	1/1	0.90	0.25	56,56,56,56	0
56	MG	1A	3079	1/1	0.90	0.15	58,58,58,58	0
56	MG	1a	1745	1/1	0.90	0.16	79,79,79,79	0
56	MG	1A	3044	1/1	0.90	0.23	45,45,45,45	0
56	MG	1A	3146	1/1	0.90	0.12	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	2a	3110	1/1	0.90	0.27	68,68,68,68	0
56	MG	1A	3857	1/1	0.90	0.10	32,32,32,32	0
56	MG	1A	3858	1/1	0.90	0.18	46,46,46,46	0
56	MG	2a	3114	1/1	0.90	0.31	53,53,53,53	0
56	MG	2A	3460	1/1	0.90	0.11	54,54,54,54	0
56	MG	1A	3359	1/1	0.90	0.14	53,53,53,53	0
56	MG	1x	106	1/1	0.90	0.31	63,63,63,63	0
56	MG	2A	3469	1/1	0.90	0.12	51,51,51,51	0
56	MG	1A	3179	1/1	0.90	0.10	44,44,44,44	0
56	MG	2a	3137	1/1	0.90	0.06	67,67,67,67	0
56	MG	1A	3728	1/1	0.90	0.15	52,52,52,52	0
56	MG	1a	1615	1/1	0.90	0.20	55,55,55,55	0
56	MG	2a	3143	1/1	0.90	0.09	88,88,88,88	0
56	MG	2a	3145	1/1	0.90	0.17	61,61,61,61	0
56	MG	1a	1758	1/1	0.90	0.07	82,82,82,82	0
56	MG	1A	4015	1/1	0.90	0.10	15,15,15,15	0
56	MG	2a	3149	1/1	0.90	0.20	65,65,65,65	0
56	MG	2A	3260	1/1	0.90	0.20	72,72,72,72	0
56	MG	2A	3487	1/1	0.90	0.10	32,32,32,32	0
56	MG	1A	3729	1/1	0.90	0.13	48,48,48,48	0
56	MG	1a	1620	1/1	0.90	0.08	58,58,58,58	0
56	MG	2A	3269	1/1	0.90	0.11	41,41,41,41	0
56	MG	1A	3740	1/1	0.90	0.20	36,36,36,36	0
56	MG	1A	3511	1/1	0.90	0.10	76,76,76,76	0
56	MG	1B	201	1/1	0.90	0.12	50,50,50,50	0
56	MG	1B	207	1/1	0.90	0.09	33,33,33,33	0
56	MG	1A	3393	1/1	0.90	0.14	52,52,52,52	0
56	MG	2a	3162	1/1	0.90	0.16	75,75,75,75	0
56	MG	2A	3279	1/1	0.90	0.12	62,62,62,62	0
56	MG	2a	3164	1/1	0.90	0.16	82,82,82,82	0
56	MG	2a	3166	1/1	0.90	0.08	66,66,66,66	0
56	MG	1A	3879	1/1	0.90	0.18	31,31,31,31	0
56	MG	1a	1771	1/1	0.90	0.07	63,63,63,63	0
56	MG	2a	3169	1/1	0.90	0.11	99,99,99,99	0
56	MG	2A	3283	1/1	0.90	0.08	48,48,48,48	0
56	MG	1A	3037	1/1	0.90	0.15	58,58,58,58	0
56	MG	1A	3328	1/1	0.90	0.11	65,65,65,65	0
56	MG	2A	3510	1/1	0.90	0.10	34,34,34,34	0
56	MG	1A	3398	1/1	0.90	0.24	43,43,43,43	0
56	MG	2a	3179	1/1	0.90	0.16	49,49,49,49	0
56	MG	1A	3783	1/1	0.90	0.13	37,37,37,37	0
56	MG	1A	3625	1/1	0.90	0.12	22,22,22,22	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3053	1/1	0.90	0.23	55,55,55,55	0
56	MG	2A	3520	1/1	0.90	0.11	42,42,42,42	0
56	MG	1a	1650	1/1	0.90	0.39	57,57,57,57	0
56	MG	1B	224	1/1	0.90	0.06	44,44,44,44	0
56	MG	1A	3519	1/1	0.90	0.10	62,62,62,62	0
56	MG	2A	3314	1/1	0.90	0.20	56,56,56,56	0
56	MG	1D	302	1/1	0.90	0.15	65,65,65,65	0
56	MG	1A	3060	1/1	0.90	0.26	67,67,67,67	0
56	MG	1D	311	1/1	0.90	0.12	55,55,55,55	0
56	MG	2A	3321	1/1	0.90	0.14	35,35,35,35	0
56	MG	1A	3478	1/1	0.90	0.13	65,65,65,65	0
56	MG	2A	3542	1/1	0.90	0.20	42,42,42,42	0
56	MG	1a	1667	1/1	0.90	0.12	65,65,65,65	0
56	MG	2a	3211	1/1	0.90	0.09	61,61,61,61	0
56	MG	2A	3546	1/1	0.90	0.16	44,44,44,44	0
56	MG	2A	3333	1/1	0.90	0.09	51,51,51,51	0
56	MG	2A	3556	1/1	0.90	0.13	62,62,62,62	0
56	MG	1A	3898	1/1	0.90	0.07	25,25,25,25	0
56	MG	2A	3566	1/1	0.90	0.17	62,62,62,62	0
56	MG	2A	3338	1/1	0.90	0.18	56,56,56,56	0
56	MG	2a	3226	1/1	0.90	0.10	58,58,58,58	0
56	MG	1A	3905	1/1	0.90	0.11	51,51,51,51	0
56	MG	1F	305	1/1	0.90	0.19	44,44,44,44	0
56	MG	2B	3006	1/1	0.90	0.14	58,58,58,58	0
56	MG	1A	3050	1/1	0.90	0.11	48,48,48,48	0
56	MG	2B	3010	1/1	0.90	0.30	50,50,50,50	0
56	MG	1F	308	1/1	0.90	0.21	55,55,55,55	0
56	MG	2A	3093	1/1	0.90	0.26	55,55,55,55	0
56	MG	1A	3917	1/1	0.90	0.14	31,31,31,31	0
56	MG	2F	301	1/1	0.90	0.13	57,57,57,57	0
56	MG	1A	3655	1/1	0.90	0.16	48,48,48,48	0
56	MG	2A	3361	1/1	0.90	0.12	54,54,54,54	0
56	MG	2a	3242	1/1	0.90	0.19	61,61,61,61	0
56	MG	1a	1808	1/1	0.90	0.12	52,52,52,52	0
56	MG	1a	1811	1/1	0.90	0.13	55,55,55,55	0
56	MG	1a	1688	1/1	0.90	0.11	56,56,56,56	0
56	MG	1A	3663	1/1	0.90	0.09	55,55,55,55	0
56	MG	2a	3002	1/1	0.90	0.15	65,65,65,65	0
56	MG	1A	3813	1/1	0.90	0.17	56,56,56,56	0
56	MG	1A	3527	1/1	0.90	0.10	41,41,41,41	0
56	MG	2a	3255	1/1	0.90	0.13	70,70,70,70	0
56	MG	1A	3261	1/1	0.90	0.30	62,62,62,62	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1828	1/1	0.90	0.08	61,61,61,61	0
56	MG	1A	3110	1/1	0.90	0.23	55,55,55,55	0
56	MG	1A	3206	1/1	0.90	0.18	21,21,21,21	0
56	MG	2a	3260	1/1	0.90	0.28	72,72,72,72	0
56	MG	1a	1836	1/1	0.90	0.16	41,41,41,41	0
56	MG	2a	3262	1/1	0.90	0.17	56,56,56,56	0
56	MG	1A	3209	1/1	0.90	0.14	40,40,40,40	0
56	MG	1A	3111	1/1	0.90	0.28	68,68,68,68	0
56	MG	2a	3019	1/1	0.90	0.10	73,73,73,73	0
56	MG	2A	3138	1/1	0.90	0.13	54,54,54,54	0
56	MG	1A	3685	1/1	0.90	0.20	73,73,73,73	0
56	MG	1A	3307	1/1	0.91	0.19	59,59,59,59	0
56	MG	1A	3582	1/1	0.91	0.15	45,45,45,45	0
56	MG	1A	3468	1/1	0.91	0.14	57,57,57,57	0
56	MG	1A	3586	1/1	0.91	0.14	57,57,57,57	0
56	MG	1a	1670	1/1	0.91	0.10	43,43,43,43	0
56	MG	1A	3104	1/1	0.91	0.11	55,55,55,55	0
56	MG	1a	1673	1/1	0.91	0.13	71,71,71,71	0
56	MG	1A	3477	1/1	0.91	0.08	66,66,66,66	0
56	MG	1A	4019	1/1	0.91	0.42	47,47,47,47	0
56	MG	1A	3600	1/1	0.91	0.11	54,54,54,54	0
56	MG	1A	3314	1/1	0.91	0.11	55,55,55,55	0
56	MG	1A	3606	1/1	0.91	0.12	32,32,32,32	0
56	MG	1a	1686	1/1	0.91	0.15	49,49,49,49	0
56	MG	1A	3366	1/1	0.91	0.08	46,46,46,46	0
56	MG	1A	3423	1/1	0.91	0.16	57,57,57,57	0
56	MG	1e	3003	1/1	0.91	0.13	68,68,68,68	0
56	MG	1A	3424	1/1	0.91	0.08	61,61,61,61	0
56	MG	1A	3830	1/1	0.91	0.16	63,63,63,63	0
56	MG	2a	3016	1/1	0.91	0.25	65,65,65,65	0
56	MG	2a	3017	1/1	0.91	0.09	70,70,70,70	0
56	MG	2a	3018	1/1	0.91	0.16	49,49,49,49	0
56	MG	1A	3482	1/1	0.91	0.12	48,48,48,48	0
56	MG	1A	3003	1/1	0.91	0.29	46,46,46,46	0
56	MG	1A	3624	1/1	0.91	0.08	18,18,18,18	0
56	MG	1A	3106	1/1	0.91	0.15	64,64,64,64	0
56	MG	1r	3002	1/1	0.91	0.13	55,55,55,55	0
56	MG	1A	3108	1/1	0.91	0.11	56,56,56,56	0
56	MG	1A	3839	1/1	0.91	0.13	40,40,40,40	0
56	MG	2A	3327	1/1	0.91	0.14	36,36,36,36	0
56	MG	1A	3430	1/1	0.91	0.10	62,62,62,62	0
56	MG	2A	3329	1/1	0.91	0.10	37,37,37,37	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3845	1/1	0.91	0.08	42,42,42,42	0
56	MG	1a	1710	1/1	0.91	0.23	71,71,71,71	0
56	MG	1A	3076	1/1	0.91	0.13	58,58,58,58	0
56	MG	2A	3339	1/1	0.91	0.14	31,31,31,31	0
56	MG	2A	3340	1/1	0.91	0.18	56,56,56,56	0
56	MG	1A	3850	1/1	0.91	0.14	41,41,41,41	0
56	MG	2a	3049	1/1	0.91	0.18	64,64,64,64	0
56	MG	1A	3640	1/1	0.91	0.14	24,24,24,24	0
56	MG	1A	3645	1/1	0.91	0.13	23,23,23,23	0
56	MG	1A	3496	1/1	0.91	0.08	88,88,88,88	0
56	MG	2A	3353	1/1	0.91	0.23	61,61,61,61	0
56	MG	2A	3014	1/1	0.91	0.06	66,66,66,66	0
56	MG	1A	3217	1/1	0.91	0.18	42,42,42,42	0
56	MG	1a	1726	1/1	0.91	0.26	54,54,54,54	0
56	MG	2A	3360	1/1	0.91	0.20	58,58,58,58	0
56	MG	2A	3018	1/1	0.91	0.08	50,50,50,50	0
56	MG	2A	3026	1/1	0.91	0.12	51,51,51,51	0
56	MG	2A	3030	1/1	0.91	0.15	50,50,50,50	0
56	MG	1A	3661	1/1	0.91	0.12	36,36,36,36	0
56	MG	1A	3045	1/1	0.91	0.11	70,70,70,70	0
56	MG	2a	3068	1/1	0.91	0.09	67,67,67,67	0
56	MG	1A	3500	1/1	0.91	0.09	55,55,55,55	0
56	MG	2A	3369	1/1	0.91	0.10	31,31,31,31	0
56	MG	1A	3502	1/1	0.91	0.10	63,63,63,63	0
56	MG	1N	3003	1/1	0.91	0.07	42,42,42,42	0
56	MG	1N	3005	1/1	0.91	0.06	51,51,51,51	0
56	MG	1A	3113	1/1	0.91	0.11	63,63,63,63	0
56	MG	2a	3079	1/1	0.91	0.13	69,69,69,69	0
56	MG	1A	3673	1/1	0.91	0.13	65,65,65,65	0
56	MG	2a	3082	1/1	0.91	0.13	61,61,61,61	0
56	MG	2a	3084	1/1	0.91	0.42	56,56,56,56	0
56	MG	1P	203	1/1	0.91	0.23	65,65,65,65	0
56	MG	1A	3878	1/1	0.91	0.11	43,43,43,43	0
56	MG	2a	3094	1/1	0.91	0.25	49,49,49,49	0
56	MG	2a	3096	1/1	0.91	0.32	80,80,80,80	0
56	MG	1a	1746	1/1	0.91	0.25	55,55,55,55	0
56	MG	1A	3332	1/1	0.91	0.10	55,55,55,55	0
56	MG	2A	3057	1/1	0.91	0.33	58,58,58,58	0
56	MG	2a	3103	1/1	0.91	0.39	69,69,69,69	0
56	MG	1A	3678	1/1	0.91	0.18	61,61,61,61	0
56	MG	2A	3070	1/1	0.91	0.26	32,32,32,32	0
56	MG	1A	3233	1/1	0.91	0.19	34,34,34,34	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3114	1/1	0.91	0.09	52,52,52,52	0
56	MG	1A	3684	1/1	0.91	0.10	50,50,50,50	0
56	MG	2a	3115	1/1	0.91	0.36	54,54,54,54	0
56	MG	1U	202	1/1	0.91	0.29	50,50,50,50	0
56	MG	2a	3118	1/1	0.91	0.09	38,38,38,38	0
56	MG	2A	3077	1/1	0.91	0.23	64,64,64,64	0
56	MG	1W	3002	1/1	0.91	0.30	55,55,55,55	0
56	MG	1A	3047	1/1	0.91	0.07	67,67,67,67	0
56	MG	2a	3131	1/1	0.91	0.23	44,44,44,44	0
56	MG	2A	3082	1/1	0.91	0.16	41,41,41,41	0
56	MG	2A	3083	1/1	0.91	0.17	49,49,49,49	0
56	MG	1A	3689	1/1	0.91	0.16	55,55,55,55	0
56	MG	1A	3444	1/1	0.91	0.26	36,36,36,36	0
56	MG	2a	3141	1/1	0.91	0.08	73,73,73,73	0
56	MG	2A	3086	1/1	0.91	0.30	49,49,49,49	0
56	MG	1A	3019	1/1	0.91	0.10	49,49,49,49	0
56	MG	2A	3089	1/1	0.91	0.22	40,40,40,40	0
56	MG	1A	3698	1/1	0.91	0.07	48,48,48,48	0
56	MG	10	102	1/1	0.91	0.21	60,60,60,60	0
56	MG	10	105	1/1	0.91	0.26	53,53,53,53	0
56	MG	2A	3096	1/1	0.91	0.18	40,40,40,40	0
56	MG	2A	3432	1/1	0.91	0.14	61,61,61,61	0
56	MG	1A	3446	1/1	0.91	0.18	62,62,62,62	0
56	MG	10	107	1/1	0.91	0.10	55,55,55,55	0
56	MG	2A	3105	1/1	0.91	0.23	54,54,54,54	0
56	MG	11	4000	1/1	0.91	0.10	43,43,43,43	0
56	MG	2A	3107	1/1	0.91	0.19	45,45,45,45	0
56	MG	1A	3080	1/1	0.91	0.10	50,50,50,50	0
56	MG	1A	3255	1/1	0.91	0.20	58,58,58,58	0
56	MG	15	104	1/1	0.91	0.16	62,62,62,62	0
56	MG	2A	3116	1/1	0.91	0.18	47,47,47,47	0
56	MG	1A	3040	1/1	0.91	0.12	45,45,45,45	0
56	MG	2a	3165	1/1	0.91	0.12	52,52,52,52	0
56	MG	1A	3919	1/1	0.91	0.13	49,49,49,49	0
56	MG	2A	3450	1/1	0.91	0.22	62,62,62,62	0
56	MG	1A	3025	1/1	0.91	0.24	57,57,57,57	0
56	MG	1A	3093	1/1	0.91	0.14	47,47,47,47	0
56	MG	2A	3126	1/1	0.91	0.15	50,50,50,50	0
56	MG	2A	3456	1/1	0.91	0.10	51,51,51,51	0
56	MG	1a	1779	1/1	0.91	0.24	82,82,82,82	0
56	MG	1A	3098	1/1	0.91	0.15	52,52,52,52	0
56	MG	1A	3534	1/1	0.91	0.15	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	3176	1/1	0.91	0.21	56,56,56,56	0
56	MG	1a	1784	1/1	0.91	0.10	56,56,56,56	0
56	MG	1A	3353	1/1	0.91	0.10	58,58,58,58	0
56	MG	2a	3184	1/1	0.91	0.10	80,80,80,80	0
56	MG	2A	3475	1/1	0.91	0.13	35,35,35,35	0
56	MG	1a	1788	1/1	0.91	0.10	76,76,76,76	0
56	MG	1A	3932	1/1	0.91	0.17	20,20,20,20	0
56	MG	1A	3539	1/1	0.91	0.13	44,44,44,44	0
56	MG	1a	1607	1/1	0.91	0.26	78,78,78,78	0
56	MG	2a	3197	1/1	0.91	0.24	58,58,58,58	0
56	MG	1a	1793	1/1	0.91	0.13	77,77,77,77	0
56	MG	1A	3732	1/1	0.91	0.15	45,45,45,45	0
56	MG	2A	3152	1/1	0.91	0.25	58,58,58,58	0
56	MG	2A	3157	1/1	0.91	0.12	43,43,43,43	0
56	MG	2a	3203	1/1	0.91	0.15	69,69,69,69	0
56	MG	1A	3937	1/1	0.91	0.10	58,58,58,58	0
56	MG	2A	3161	1/1	0.91	0.19	32,32,32,32	0
56	MG	2A	3162	1/1	0.91	0.35	50,50,50,50	0
56	MG	2a	3208	1/1	0.91	0.23	58,58,58,58	0
56	MG	1A	3733	1/1	0.91	0.12	35,35,35,35	0
56	MG	2A	3176	1/1	0.91	0.24	43,43,43,43	0
56	MG	1a	1613	1/1	0.91	0.11	60,60,60,60	0
56	MG	1a	1614	1/1	0.91	0.35	68,68,68,68	0
56	MG	2A	3503	1/1	0.91	0.17	59,59,59,59	0
56	MG	2A	3183	1/1	0.91	0.10	56,56,56,56	0
56	MG	1A	3543	1/1	0.91	0.15	65,65,65,65	0
56	MG	1A	3274	1/1	0.91	0.14	41,41,41,41	0
56	MG	2a	3222	1/1	0.91	0.20	46,46,46,46	0
56	MG	1A	3547	1/1	0.91	0.11	54,54,54,54	0
56	MG	1a	1804	1/1	0.91	0.10	52,52,52,52	0
56	MG	1A	3057	1/1	0.91	0.17	50,50,50,50	0
56	MG	1a	1807	1/1	0.91	0.18	60,60,60,60	0
56	MG	1A	3947	1/1	0.91	0.16	43,43,43,43	0
56	MG	1A	3140	1/1	0.91	0.16	62,62,62,62	0
56	MG	1a	1812	1/1	0.91	0.28	49,49,49,49	0
56	MG	2A	3216	1/1	0.91	0.09	45,45,45,45	0
56	MG	1A	3778	1/1	0.91	0.12	23,23,23,23	0
56	MG	1A	3408	1/1	0.91	0.12	42,42,42,42	0
56	MG	1A	3200	1/1	0.91	0.17	46,46,46,46	0
56	MG	1A	3564	1/1	0.91	0.15	35,35,35,35	0
56	MG	1a	1820	1/1	0.91	0.16	54,54,54,54	0
56	MG	1A	3785	1/1	0.91	0.09	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	1A	3982	1/1	0.91	0.10	49,49,49,49	0
56	MG	2A	3229	1/1	0.91	0.24	55,55,55,55	0
56	MG	1a	1830	1/1	0.91	0.14	61,61,61,61	0
56	MG	1A	3984	1/1	0.91	0.07	47,47,47,47	0
56	MG	2A	3235	1/1	0.91	0.22	52,52,52,52	0
56	MG	2a	3252	1/1	0.91	0.22	70,70,70,70	0
56	MG	1a	1833	1/1	0.91	0.18	59,59,59,59	0
56	MG	1a	1644	1/1	0.91	0.20	49,49,49,49	0
56	MG	1a	1645	1/1	0.91	0.15	67,67,67,67	0
56	MG	1a	1840	1/1	0.91	0.17	41,41,41,41	0
56	MG	1a	1841	1/1	0.91	0.23	64,64,64,64	0
56	MG	2B	3001	1/1	0.91	0.17	65,65,65,65	0
56	MG	1A	3412	1/1	0.91	0.22	45,45,45,45	0
56	MG	1A	3572	1/1	0.91	0.15	50,50,50,50	0
56	MG	2A	3254	1/1	0.91	0.20	40,40,40,40	0
56	MG	1A	3573	1/1	0.91	0.20	47,47,47,47	0
56	MG	1A	3805	1/1	0.91	0.17	41,41,41,41	0
56	MG	1A	3303	1/1	0.91	0.25	63,63,63,63	0
56	MG	1A	3999	1/1	0.91	0.10	21,21,21,21	0
56	MG	2D	302	1/1	0.91	0.19	54,54,54,54	0
56	MG	1A	3580	1/1	0.91	0.10	46,46,46,46	0
56	MG	2A	3156	1/1	0.92	0.10	49,49,49,49	0
56	MG	1A	3474	1/1	0.92	0.16	39,39,39,39	0
56	MG	2A	3404	1/1	0.92	0.14	43,43,43,43	0
56	MG	13	102	1/1	0.92	0.12	56,56,56,56	0
56	MG	1A	3475	1/1	0.92	0.12	53,53,53,53	0
56	MG	1A	3966	1/1	0.92	0.34	47,47,47,47	0
56	MG	1A	3821	1/1	0.92	0.12	23,23,23,23	0
56	MG	2A	3166	1/1	0.92	0.29	49,49,49,49	0
56	MG	2A	3172	1/1	0.92	0.24	49,49,49,49	0
56	MG	1A	3381	1/1	0.92	0.12	47,47,47,47	0
56	MG	1A	3980	1/1	0.92	0.14	50,50,50,50	0
56	MG	1a	1747	1/1	0.92	0.10	56,56,56,56	0
56	MG	1A	3124	1/1	0.92	0.12	56,56,56,56	0
56	MG	1a	1749	1/1	0.92	0.15	57,57,57,57	0
56	MG	2a	3062	1/1	0.92	0.12	59,59,59,59	0
56	MG	1A	3030	1/1	0.92	0.18	49,49,49,49	0
56	MG	2a	3064	1/1	0.92	0.07	77,77,77,77	0
56	MG	1k	3001	1/1	0.92	0.29	70,70,70,70	0
56	MG	2A	3430	1/1	0.92	0.21	45,45,45,45	0
56	MG	2A	3198	1/1	0.92	0.17	41,41,41,41	0
56	MG	2A	3434	1/1	0.92	0.13	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	3199	1/1	0.92	0.26	39,39,39,39	0
56	MG	1A	3677	1/1	0.92	0.16	51,51,51,51	0
56	MG	1a	1752	1/1	0.92	0.07	72,72,72,72	0
56	MG	1A	3827	1/1	0.92	0.07	31,31,31,31	0
56	MG	1q	202	1/1	0.92	0.19	62,62,62,62	0
56	MG	1q	203	1/1	0.92	0.25	51,51,51,51	0
56	MG	1A	3992	1/1	0.92	0.17	55,55,55,55	0
56	MG	1A	3082	1/1	0.92	0.08	56,56,56,56	0
56	MG	2A	3446	1/1	0.92	0.08	42,42,42,42	0
56	MG	1A	3555	1/1	0.92	0.12	46,46,46,46	0
56	MG	1A	3388	1/1	0.92	0.11	53,53,53,53	0
56	MG	2a	3089	1/1	0.92	0.13	66,66,66,66	0
56	MG	2a	3091	1/1	0.92	0.37	61,61,61,61	0
56	MG	2A	3221	1/1	0.92	0.14	47,47,47,47	0
56	MG	1x	102	1/1	0.92	0.23	67,67,67,67	0
56	MG	1x	103	1/1	0.92	0.27	70,70,70,70	0
56	MG	1A	3062	1/1	0.92	0.12	48,48,48,48	0
56	MG	1A	3345	1/1	0.92	0.06	60,60,60,60	0
56	MG	1x	108	1/1	0.92	0.14	70,70,70,70	0
56	MG	2A	3458	1/1	0.92	0.19	54,54,54,54	0
56	MG	2A	3459	1/1	0.92	0.10	49,49,49,49	0
56	MG	1A	3565	1/1	0.92	0.13	45,45,45,45	0
56	MG	2A	3461	1/1	0.92	0.07	47,47,47,47	0
56	MG	1A	4007	1/1	0.92	0.27	36,36,36,36	0
56	MG	1A	3567	1/1	0.92	0.15	39,39,39,39	0
56	MG	2A	3234	1/1	0.92	0.22	48,48,48,48	0
56	MG	2A	3003	1/1	0.92	0.10	53,53,53,53	0
56	MG	1a	1763	1/1	0.92	0.09	69,69,69,69	0
56	MG	2a	3121	1/1	0.92	0.24	63,63,63,63	0
56	MG	2A	3006	1/1	0.92	0.15	64,64,64,64	0
56	MG	1a	1764	1/1	0.92	0.17	68,68,68,68	0
56	MG	1A	3004	1/1	0.92	0.21	40,40,40,40	0
56	MG	2a	3127	1/1	0.92	0.20	61,61,61,61	0
56	MG	1A	3095	1/1	0.92	0.12	50,50,50,50	0
56	MG	2a	3132	1/1	0.92	0.20	65,65,65,65	0
56	MG	1A	3844	1/1	0.92	0.20	71,71,71,71	0
56	MG	1A	3700	1/1	0.92	0.13	63,63,63,63	0
56	MG	1A	3038	1/1	0.92	0.28	52,52,52,52	0
56	MG	2A	3255	1/1	0.92	0.17	52,52,52,52	0
56	MG	2a	3140	1/1	0.92	0.18	63,63,63,63	0
56	MG	1A	3222	1/1	0.92	0.14	50,50,50,50	0
56	MG	2A	3025	1/1	0.92	0.21	67,67,67,67	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2a	3144	1/1	0.92	0.15	56,56,56,56	0
56	MG	1A	3576	1/1	0.92	0.16	19,19,19,19	0
56	MG	1B	204	1/1	0.92	0.20	54,54,54,54	0
56	MG	1A	3022	1/1	0.92	0.32	59,59,59,59	0
56	MG	1B	208	1/1	0.92	0.22	56,56,56,56	0
56	MG	1A	3142	1/1	0.92	0.16	57,57,57,57	0
56	MG	1A	3498	1/1	0.92	0.14	67,67,67,67	0
56	MG	1A	3315	1/1	0.92	0.19	49,49,49,49	0
56	MG	1A	3318	1/1	0.92	0.23	52,52,52,52	0
56	MG	2A	3043	1/1	0.92	0.27	49,49,49,49	0
56	MG	1A	3452	1/1	0.92	0.11	56,56,56,56	0
56	MG	1A	3234	1/1	0.92	0.15	43,43,43,43	0
56	MG	1A	3504	1/1	0.92	0.22	60,60,60,60	0
56	MG	1A	3235	1/1	0.92	0.21	46,46,46,46	0
56	MG	2A	3049	1/1	0.92	0.20	52,52,52,52	0
56	MG	2A	3513	1/1	0.92	0.21	61,61,61,61	0
56	MG	2A	3514	1/1	0.92	0.08	60,60,60,60	0
56	MG	1A	3734	1/1	0.92	0.14	38,38,38,38	0
56	MG	1A	3738	1/1	0.92	0.10	37,37,37,37	0
56	MG	1A	3323	1/1	0.92	0.11	58,58,58,58	0
56	MG	1A	3178	1/1	0.92	0.27	41,41,41,41	0
56	MG	2A	3065	1/1	0.92	0.20	35,35,35,35	0
56	MG	1A	3882	1/1	0.92	0.12	49,49,49,49	0
56	MG	2A	3299	1/1	0.92	0.10	46,46,46,46	0
56	MG	2A	3302	1/1	0.92	0.13	41,41,41,41	0
56	MG	2A	3528	1/1	0.92	0.25	49,49,49,49	0
56	MG	2A	3303	1/1	0.92	0.21	55,55,55,55	0
56	MG	1F	301	1/1	0.92	0.11	52,52,52,52	0
56	MG	2A	3310	1/1	0.92	0.09	31,31,31,31	0
56	MG	2A	3539	1/1	0.92	0.30	35,35,35,35	0
56	MG	1A	3750	1/1	0.92	0.08	21,21,21,21	0
56	MG	2A	3312	1/1	0.92	0.09	28,28,28,28	0
56	MG	1A	3610	1/1	0.92	0.10	37,37,37,37	0
56	MG	2A	3544	1/1	0.92	0.21	45,45,45,45	0
56	MG	1A	3143	1/1	0.92	0.09	52,52,52,52	0
56	MG	1F	306	1/1	0.92	0.17	33,33,33,33	0
56	MG	2A	3554	1/1	0.92	0.58	51,51,51,51	0
56	MG	1A	3890	1/1	0.92	0.11	37,37,37,37	0
56	MG	1A	3761	1/1	0.92	0.13	66,66,66,66	0
56	MG	2A	3562	1/1	0.92	0.16	49,49,49,49	0
56	MG	2A	3564	1/1	0.92	0.10	59,59,59,59	0
56	MG	1A	3892	1/1	0.92	0.14	43,43,43,43	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3182	1/1	0.92	0.17	42,42,42,42	0
56	MG	1A	3771	1/1	0.92	0.13	71,71,71,71	0
56	MG	1A	3772	1/1	0.92	0.08	60,60,60,60	0
56	MG	2A	3330	1/1	0.92	0.18	47,47,47,47	0
56	MG	2A	3332	1/1	0.92	0.08	38,38,38,38	0
56	MG	1A	3100	1/1	0.92	0.14	48,48,48,48	0
56	MG	1A	3329	1/1	0.92	0.09	52,52,52,52	0
56	MG	1A	3780	1/1	0.92	0.10	43,43,43,43	0
56	MG	2a	3212	1/1	0.92	0.15	54,54,54,54	0
56	MG	1a	1817	1/1	0.92	0.13	52,52,52,52	0
56	MG	2D	304	1/1	0.92	0.13	42,42,42,42	0
56	MG	2a	3215	1/1	0.92	0.20	56,56,56,56	0
56	MG	1A	3102	1/1	0.92	0.14	60,60,60,60	0
56	MG	1A	3041	1/1	0.92	0.10	55,55,55,55	0
56	MG	2a	3220	1/1	0.92	0.19	63,63,63,63	0
56	MG	1P	204	1/1	0.92	0.10	47,47,47,47	0
56	MG	1a	1822	1/1	0.92	0.14	52,52,52,52	0
56	MG	2A	3349	1/1	0.92	0.27	59,59,59,59	0
56	MG	1A	3260	1/1	0.92	0.13	58,58,58,58	0
56	MG	1a	1827	1/1	0.92	0.08	71,71,71,71	0
56	MG	1A	3466	1/1	0.92	0.07	40,40,40,40	0
56	MG	1A	3791	1/1	0.92	0.08	20,20,20,20	0
56	MG	1A	3522	1/1	0.92	0.17	69,69,69,69	0
56	MG	1A	3147	1/1	0.92	0.10	54,54,54,54	0
56	MG	2a	3004	1/1	0.92	0.18	46,46,46,46	0
56	MG	1a	1708	1/1	0.92	0.10	61,61,61,61	0
56	MG	2a	3006	1/1	0.92	0.13	62,62,62,62	0
56	MG	2a	3238	1/1	0.92	0.21	55,55,55,55	0
56	MG	1A	3797	1/1	0.92	0.51	40,40,40,40	0
56	MG	1a	1839	1/1	0.92	0.18	56,56,56,56	0
56	MG	1A	3525	1/1	0.92	0.17	60,60,60,60	0
56	MG	1A	3335	1/1	0.92	0.08	55,55,55,55	0
56	MG	1a	1842	1/1	0.92	0.09	73,73,73,73	0
56	MG	1A	3807	1/1	0.92	0.15	57,57,57,57	0
56	MG	1a	1718	1/1	0.92	0.14	48,48,48,48	0
56	MG	1A	3648	1/1	0.92	0.08	20,20,20,20	0
56	MG	1A	3809	1/1	0.92	0.19	45,45,45,45	0
56	MG	1A	3528	1/1	0.92	0.13	54,54,54,54	0
56	MG	1A	3942	1/1	0.92	0.15	53,53,53,53	0
56	MG	2a	3023	1/1	0.92	0.10	65,65,65,65	0
56	MG	1A	3657	1/1	0.92	0.16	49,49,49,49	0
56	MG	2A	3140	1/1	0.92	0.16	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1729	1/1	0.92	0.22	35,35,35,35	0
56	MG	2A	3388	1/1	0.92	0.14	45,45,45,45	0
56	MG	1A	3471	1/1	0.92	0.09	74,74,74,74	0
56	MG	1A	3336	1/1	0.92	0.18	49,49,49,49	0
56	MG	1a	1733	1/1	0.92	0.24	65,65,65,65	0
56	MG	1A	3817	1/1	0.92	0.22	51,51,51,51	0
56	MG	2A	3151	1/1	0.92	0.19	47,47,47,47	0
56	MG	2f	3001	1/1	0.92	0.09	59,59,59,59	0
56	MG	1A	3664	1/1	0.92	0.06	15,15,15,15	0
56	MG	2l	201	1/1	0.92	0.14	63,63,63,63	0
56	MG	2a	3038	1/1	0.92	0.18	57,57,57,57	0
56	MG	2a	3039	1/1	0.92	0.26	57,57,57,57	0
56	MG	2t	3001	1/1	0.92	0.17	50,50,50,50	0
56	MG	2A	3153	1/1	0.92	0.11	33,33,33,33	0
56	MG	1A	3915	1/1	0.93	0.18	58,58,58,58	0
56	MG	2A	3451	1/1	0.93	0.16	61,61,61,61	0
56	MG	1E	302	1/1	0.93	0.06	48,48,48,48	0
56	MG	1E	303	1/1	0.93	0.14	58,58,58,58	0
56	MG	1A	3810	1/1	0.93	0.07	48,48,48,48	0
56	MG	1A	3577	1/1	0.93	0.17	60,60,60,60	0
56	MG	1A	3578	1/1	0.93	0.14	64,64,64,64	0
56	MG	2A	3256	1/1	0.93	0.11	34,34,34,34	0
56	MG	2A	3052	1/1	0.93	0.14	43,43,43,43	0
56	MG	1a	1671	1/1	0.93	0.07	38,38,38,38	0
56	MG	1A	3330	1/1	0.93	0.11	41,41,41,41	0
56	MG	2A	3467	1/1	0.93	0.14	46,46,46,46	0
56	MG	2a	3076	1/1	0.93	0.09	83,83,83,83	0
56	MG	2A	3262	1/1	0.93	0.16	54,54,54,54	0
56	MG	2A	3263	1/1	0.93	0.25	56,56,56,56	0
56	MG	2A	3473	1/1	0.93	0.14	64,64,64,64	0
56	MG	1A	3002	1/1	0.93	0.09	53,53,53,53	0
56	MG	1A	3925	1/1	0.93	0.18	42,42,42,42	0
56	MG	2A	3476	1/1	0.93	0.11	33,33,33,33	0
56	MG	2a	3087	1/1	0.93	0.19	55,55,55,55	0
56	MG	2a	3088	1/1	0.93	0.12	54,54,54,54	0
56	MG	2A	3478	1/1	0.93	0.10	44,44,44,44	0
56	MG	1A	3138	1/1	0.93	0.27	55,55,55,55	0
56	MG	1A	3236	1/1	0.93	0.15	46,46,46,46	0
56	MG	1A	3930	1/1	0.93	0.20	44,44,44,44	0
56	MG	1a	1684	1/1	0.93	0.11	48,48,48,48	0
56	MG	2A	3485	1/1	0.93	0.09	38,38,38,38	0
56	MG	1A	3368	1/1	0.93	0.13	25,25,25,25	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3590	1/1	0.93	0.10	47,47,47,47	0
56	MG	1A	3237	1/1	0.93	0.11	46,46,46,46	0
56	MG	2a	3104	1/1	0.93	0.20	62,62,62,62	0
56	MG	2A	3078	1/1	0.93	0.19	60,60,60,60	0
56	MG	1A	3935	1/1	0.93	0.09	28,28,28,28	0
56	MG	1A	3015	1/1	0.93	0.20	52,52,52,52	0
56	MG	2A	3493	1/1	0.93	0.11	58,58,58,58	0
56	MG	1A	3710	1/1	0.93	0.21	39,39,39,39	0
56	MG	2A	3496	1/1	0.93	0.11	42,42,42,42	0
56	MG	1a	1695	1/1	0.93	0.18	50,50,50,50	0
56	MG	1A	3373	1/1	0.93	0.08	48,48,48,48	0
56	MG	2a	3120	1/1	0.93	0.26	63,63,63,63	0
56	MG	2A	3500	1/1	0.93	0.07	49,49,49,49	0
56	MG	1a	1698	1/1	0.93	0.11	55,55,55,55	0
56	MG	1A	3714	1/1	0.93	0.14	46,46,46,46	0
56	MG	2a	3124	1/1	0.93	0.39	63,63,63,63	0
56	MG	1A	3715	1/1	0.93	0.08	63,63,63,63	0
56	MG	2A	3296	1/1	0.93	0.07	45,45,45,45	0
56	MG	2a	3129	1/1	0.93	0.19	65,65,65,65	0
56	MG	1R	204	1/1	0.93	0.14	36,36,36,36	0
56	MG	2A	3506	1/1	0.93	0.11	56,56,56,56	0
56	MG	1A	3420	1/1	0.93	0.07	56,56,56,56	0
56	MG	2a	3135	1/1	0.93	0.13	60,60,60,60	0
56	MG	2a	3136	1/1	0.93	0.18	51,51,51,51	0
56	MG	2A	3300	1/1	0.93	0.12	32,32,32,32	0
56	MG	2A	3301	1/1	0.93	0.09	38,38,38,38	0
56	MG	1A	3829	1/1	0.93	0.15	55,55,55,55	0
56	MG	1A	3310	1/1	0.93	0.19	44,44,44,44	0
56	MG	2A	3095	1/1	0.93	0.15	31,31,31,31	0
56	MG	1a	1706	1/1	0.93	0.18	64,64,64,64	0
56	MG	2A	3097	1/1	0.93	0.18	51,51,51,51	0
56	MG	1A	3945	1/1	0.93	0.22	60,60,60,60	0
56	MG	2A	3313	1/1	0.93	0.22	53,53,53,53	0
56	MG	1A	3311	1/1	0.93	0.10	65,65,65,65	0
56	MG	1A	3240	1/1	0.93	0.22	49,49,49,49	0
56	MG	2a	3150	1/1	0.93	0.13	51,51,51,51	0
56	MG	1U	205	1/1	0.93	0.27	52,52,52,52	0
56	MG	1A	3426	1/1	0.93	0.12	46,46,46,46	0
56	MG	2A	3110	1/1	0.93	0.12	45,45,45,45	0
56	MG	2A	3325	1/1	0.93	0.20	52,52,52,52	0
56	MG	1A	3835	1/1	0.93	0.20	66,66,66,66	0
56	MG	2A	3112	1/1	0.93	0.10	39,39,39,39	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3960	1/1	0.93	0.23	53,53,53,53	0
56	MG	2A	3531	1/1	0.93	0.06	52,52,52,52	0
56	MG	1A	3340	1/1	0.93	0.13	47,47,47,47	0
56	MG	2A	3115	1/1	0.93	0.20	35,35,35,35	0
56	MG	2A	3331	1/1	0.93	0.10	48,48,48,48	0
56	MG	1A	3008	1/1	0.93	0.10	58,58,58,58	0
56	MG	2A	3117	1/1	0.93	0.12	53,53,53,53	0
56	MG	1A	3968	1/1	0.93	0.16	50,50,50,50	0
56	MG	1A	3101	1/1	0.93	0.12	55,55,55,55	0
56	MG	2A	3120	1/1	0.93	0.29	45,45,45,45	0
56	MG	10	104	1/1	0.93	0.08	43,43,43,43	0
56	MG	1A	3973	1/1	0.93	0.08	51,51,51,51	0
56	MG	1A	3977	1/1	0.93	0.17	53,53,53,53	0
56	MG	2A	3127	1/1	0.93	0.21	57,57,57,57	0
56	MG	1A	3623	1/1	0.93	0.15	25,25,25,25	0
56	MG	2A	3565	1/1	0.93	0.08	52,52,52,52	0
56	MG	1A	3843	1/1	0.93	0.14	50,50,50,50	0
56	MG	1A	3317	1/1	0.93	0.19	34,34,34,34	0
56	MG	1a	1870	1/1	0.93	0.17	88,88,88,88	0
56	MG	13	101	1/1	0.93	0.17	49,49,49,49	0
56	MG	1A	3432	1/1	0.93	0.10	58,58,58,58	0
56	MG	1A	3749	1/1	0.93	0.08	27,27,27,27	0
56	MG	2a	3190	1/1	0.93	0.09	52,52,52,52	0
56	MG	1A	3009	1/1	0.93	0.24	56,56,56,56	0
56	MG	2A	3362	1/1	0.93	0.24	57,57,57,57	0
56	MG	1A	3319	1/1	0.93	0.17	57,57,57,57	0
56	MG	1A	3993	1/1	0.93	0.12	73,73,73,73	0
56	MG	1A	3856	1/1	0.93	0.10	69,69,69,69	0
56	MG	1A	3046	1/1	0.93	0.09	65,65,65,65	0
56	MG	18	102	1/1	0.93	0.13	57,57,57,57	0
56	MG	1A	3351	1/1	0.93	0.15	62,62,62,62	0
56	MG	2F	302	1/1	0.93	0.14	47,47,47,47	0
56	MG	1A	3546	1/1	0.93	0.13	36,36,36,36	0
56	MG	1A	3770	1/1	0.93	0.10	36,36,36,36	0
56	MG	2N	8001	1/1	0.93	0.05	53,53,53,53	0
56	MG	1A	3321	1/1	0.93	0.12	37,37,37,37	0
56	MG	2A	3375	1/1	0.93	0.14	48,48,48,48	0
56	MG	1A	4004	1/1	0.93	0.14	49,49,49,49	0
56	MG	1A	3486	1/1	0.93	0.07	67,67,67,67	0
56	MG	1A	3868	1/1	0.93	0.10	44,44,44,44	0
56	MG	1A	3776	1/1	0.93	0.09	20,20,20,20	0
56	MG	1a	1610	1/1	0.93	0.10	53,53,53,53	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3171	1/1	0.93	0.14	41,41,41,41	0
56	MG	1A	3873	1/1	0.93	0.15	48,48,48,48	0
56	MG	1A	4016	1/1	0.93	0.16	37,37,37,37	0
56	MG	2a	3009	1/1	0.93	0.20	65,65,65,65	0
56	MG	2A	3178	1/1	0.93	0.12	34,34,34,34	0
56	MG	1A	3552	1/1	0.93	0.13	50,50,50,50	0
56	MG	1A	3071	1/1	0.93	0.08	44,44,44,44	0
56	MG	1x	104	1/1	0.93	0.10	52,52,52,52	0
56	MG	1A	3354	1/1	0.93	0.12	43,43,43,43	0
56	MG	1A	3662	1/1	0.93	0.08	44,44,44,44	0
56	MG	2A	3195	1/1	0.93	0.15	48,48,48,48	0
56	MG	1A	3556	1/1	0.93	0.13	59,59,59,59	0
56	MG	1a	1621	1/1	0.93	0.23	68,68,68,68	0
56	MG	1B	203	1/1	0.93	0.15	61,61,61,61	0
56	MG	1A	3443	1/1	0.93	0.19	48,48,48,48	0
56	MG	1B	205	1/1	0.93	0.19	56,56,56,56	0
56	MG	2a	3025	1/1	0.93	0.21	60,60,60,60	0
56	MG	2A	3415	1/1	0.93	0.14	60,60,60,60	0
56	MG	1A	3786	1/1	0.93	0.12	57,57,57,57	0
56	MG	1A	3355	1/1	0.93	0.11	53,53,53,53	0
56	MG	2A	3007	1/1	0.93	0.19	46,46,46,46	0
56	MG	1a	1630	1/1	0.93	0.10	56,56,56,56	0
56	MG	2A	3214	1/1	0.93	0.16	43,43,43,43	0
56	MG	2A	3010	1/1	0.93	0.17	56,56,56,56	0
56	MG	2a	3246	1/1	0.93	0.13	62,62,62,62	0
56	MG	1A	3021	1/1	0.93	0.23	48,48,48,48	0
56	MG	1a	1637	1/1	0.93	0.09	53,53,53,53	0
56	MG	1A	3107	1/1	0.93	0.18	58,58,58,58	0
56	MG	2A	3429	1/1	0.93	0.20	58,58,58,58	0
56	MG	1A	3226	1/1	0.93	0.10	41,41,41,41	0
56	MG	1A	3569	1/1	0.93	0.08	45,45,45,45	0
56	MG	1A	3804	1/1	0.93	0.05	31,31,31,31	0
56	MG	1A	3449	1/1	0.93	0.12	66,66,66,66	0
56	MG	1A	3265	1/1	0.93	0.11	51,51,51,51	0
56	MG	2a	3047	1/1	0.93	0.29	63,63,63,63	0
56	MG	2A	3027	1/1	0.93	0.14	51,51,51,51	0
56	MG	1a	1785	1/1	0.93	0.06	63,63,63,63	0
56	MG	1A	3405	1/1	0.93	0.18	54,54,54,54	0
56	MG	1A	3903	1/1	0.93	0.10	19,19,19,19	0
56	MG	1a	1653	1/1	0.93	0.16	60,60,60,60	0
56	MG	2a	3054	1/1	0.93	0.06	89,89,89,89	0
56	MG	1A	3075	1/1	0.93	0.10	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	1A	3048	1/1	0.93	0.15	45,45,45,45	0
56	MG	2l	204	1/1	0.93	0.17	64,64,64,64	0
56	MG	2A	3241	1/1	0.93	0.15	65,65,65,65	0
56	MG	2A	3244	1/1	0.93	0.17	63,63,63,63	0
56	MG	1A	3911	1/1	0.93	0.09	39,39,39,39	0
56	MG	2x	3002	1/1	0.93	0.25	65,65,65,65	0
56	MG	2a	3061	1/1	0.93	0.20	73,73,73,73	0
56	MG	2A	3037	1/1	0.94	0.10	46,46,46,46	0
56	MG	28	8001	1/1	0.94	0.20	54,54,54,54	0
56	MG	2A	3308	1/1	0.94	0.11	40,40,40,40	0
56	MG	2A	3309	1/1	0.94	0.12	31,31,31,31	0
56	MG	1A	3383	1/1	0.94	0.13	56,56,56,56	0
56	MG	1A	3136	1/1	0.94	0.12	60,60,60,60	0
56	MG	1A	3026	1/1	0.94	0.10	53,53,53,53	0
56	MG	2A	3041	1/1	0.94	0.11	53,53,53,53	0
56	MG	2A	3042	1/1	0.94	0.19	43,43,43,43	0
56	MG	1A	3649	1/1	0.94	0.09	71,71,71,71	0
56	MG	1A	3109	1/1	0.94	0.20	55,55,55,55	0
56	MG	1A	3656	1/1	0.94	0.10	33,33,33,33	0
56	MG	1A	3812	1/1	0.94	0.07	51,51,51,51	0
56	MG	2A	3322	1/1	0.94	0.09	67,67,67,67	0
56	MG	1A	3121	1/1	0.94	0.26	38,38,38,38	0
56	MG	1A	3814	1/1	0.94	0.18	56,56,56,56	0
56	MG	1A	3168	1/1	0.94	0.17	42,42,42,42	0
56	MG	1A	3526	1/1	0.94	0.12	32,32,32,32	0
56	MG	1a	1766	1/1	0.94	0.14	60,60,60,60	0
56	MG	2a	3021	1/1	0.94	0.10	60,60,60,60	0
56	MG	1A	3958	1/1	0.94	0.11	64,64,64,64	0
56	MG	2A	3059	1/1	0.94	0.08	48,48,48,48	0
56	MG	1A	3270	1/1	0.94	0.07	34,34,34,34	0
56	MG	2A	3062	1/1	0.94	0.13	39,39,39,39	0
56	MG	1A	3170	1/1	0.94	0.08	44,44,44,44	0
56	MG	2A	3336	1/1	0.94	0.08	36,36,36,36	0
56	MG	2A	3066	1/1	0.94	0.14	55,55,55,55	0
56	MG	1A	3964	1/1	0.94	0.11	29,29,29,29	0
56	MG	1A	3530	1/1	0.94	0.14	51,51,51,51	0
56	MG	2A	3341	1/1	0.94	0.20	52,52,52,52	0
56	MG	1A	3275	1/1	0.94	0.08	25,25,25,25	0
56	MG	1A	3670	1/1	0.94	0.07	55,55,55,55	0
56	MG	2a	3037	1/1	0.94	0.22	53,53,53,53	0
56	MG	2A	3075	1/1	0.94	0.30	59,59,59,59	0
56	MG	1a	1605	1/1	0.94	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	1A	3972	1/1	0.94	0.25	42,42,42,42	0
56	MG	1A	3395	1/1	0.94	0.18	37,37,37,37	0
56	MG	1a	1778	1/1	0.94	0.10	57,57,57,57	0
56	MG	2A	3356	1/1	0.94	0.10	60,60,60,60	0
56	MG	1A	3538	1/1	0.94	0.20	57,57,57,57	0
56	MG	2a	3046	1/1	0.94	0.09	65,65,65,65	0
56	MG	2A	3081	1/1	0.94	0.14	44,44,44,44	0
56	MG	2a	3048	1/1	0.94	0.15	80,80,80,80	0
56	MG	1A	3276	1/1	0.94	0.10	38,38,38,38	0
56	MG	1A	3676	1/1	0.94	0.18	50,50,50,50	0
56	MG	1a	1782	1/1	0.94	0.09	43,43,43,43	0
56	MG	1a	1783	1/1	0.94	0.09	57,57,57,57	0
56	MG	1A	3981	1/1	0.94	0.13	50,50,50,50	0
56	MG	1A	3541	1/1	0.94	0.18	48,48,48,48	0
56	MG	2A	3088	1/1	0.94	0.20	51,51,51,51	0
56	MG	1A	3397	1/1	0.94	0.23	49,49,49,49	0
56	MG	1A	3278	1/1	0.94	0.12	56,56,56,56	0
56	MG	1A	3281	1/1	0.94	0.09	53,53,53,53	0
56	MG	1A	3990	1/1	0.94	0.14	36,36,36,36	0
56	MG	2A	3373	1/1	0.94	0.12	47,47,47,47	0
56	MG	1A	3401	1/1	0.94	0.11	46,46,46,46	0
56	MG	1A	3548	1/1	0.94	0.14	38,38,38,38	0
56	MG	1A	3282	1/1	0.94	0.30	52,52,52,52	0
56	MG	2A	3098	1/1	0.94	0.07	51,51,51,51	0
56	MG	2A	3100	1/1	0.94	0.13	50,50,50,50	0
56	MG	1A	3834	1/1	0.94	0.10	46,46,46,46	0
56	MG	2A	3385	1/1	0.94	0.10	35,35,35,35	0
56	MG	2a	3069	1/1	0.94	0.10	95,95,95,95	0
56	MG	2A	3386	1/1	0.94	0.08	30,30,30,30	0
56	MG	2A	3102	1/1	0.94	0.17	52,52,52,52	0
56	MG	1a	1625	1/1	0.94	0.14	52,52,52,52	0
56	MG	2A	3104	1/1	0.94	0.08	54,54,54,54	0
56	MG	1A	3123	1/1	0.94	0.07	44,44,44,44	0
56	MG	1A	3287	1/1	0.94	0.15	27,27,27,27	0
56	MG	1A	3291	1/1	0.94	0.19	48,48,48,48	0
56	MG	2a	3078	1/1	0.94	0.28	47,47,47,47	0
56	MG	1a	1629	1/1	0.94	0.12	49,49,49,49	0
56	MG	1A	3470	1/1	0.94	0.07	52,52,52,52	0
56	MG	2a	3081	1/1	0.94	0.15	52,52,52,52	0
56	MG	1A	3347	1/1	0.94	0.16	60,60,60,60	0
56	MG	1a	1636	1/1	0.94	0.09	44,44,44,44	0
56	MG	1a	1806	1/1	0.94	0.17	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	3842	1/1	0.94	0.07	56,56,56,56	0
56	MG	1A	4009	1/1	0.94	0.10	56,56,56,56	0
56	MG	2A	3407	1/1	0.94	0.14	40,40,40,40	0
56	MG	2A	3408	1/1	0.94	0.26	62,62,62,62	0
56	MG	1a	1810	1/1	0.94	0.14	38,38,38,38	0
56	MG	1a	1639	1/1	0.94	0.13	62,62,62,62	0
56	MG	2a	3095	1/1	0.94	0.27	45,45,45,45	0
56	MG	1A	3293	1/1	0.94	0.11	27,27,27,27	0
56	MG	2A	3414	1/1	0.94	0.10	49,49,49,49	0
56	MG	1A	3409	1/1	0.94	0.07	51,51,51,51	0
56	MG	1A	3297	1/1	0.94	0.20	70,70,70,70	0
56	MG	1A	3068	1/1	0.94	0.20	44,44,44,44	0
56	MG	2A	3125	1/1	0.94	0.37	56,56,56,56	0
56	MG	2a	3106	1/1	0.94	0.09	63,63,63,63	0
56	MG	1A	3849	1/1	0.94	0.16	73,73,73,73	0
56	MG	1a	1647	1/1	0.94	0.18	56,56,56,56	0
56	MG	1a	1648	1/1	0.94	0.12	47,47,47,47	0
56	MG	2a	3112	1/1	0.94	0.13	45,45,45,45	0
56	MG	2A	3130	1/1	0.94	0.20	39,39,39,39	0
56	MG	1A	3227	1/1	0.94	0.16	41,41,41,41	0
56	MG	1A	3301	1/1	0.94	0.26	53,53,53,53	0
56	MG	2A	3134	1/1	0.94	0.14	51,51,51,51	0
56	MG	2a	3117	1/1	0.94	0.17	58,58,58,58	0
56	MG	2A	3135	1/1	0.94	0.17	61,61,61,61	0
56	MG	1a	1825	1/1	0.94	0.11	43,43,43,43	0
56	MG	2A	3433	1/1	0.94	0.22	65,65,65,65	0
56	MG	1A	3176	1/1	0.94	0.21	40,40,40,40	0
56	MG	1A	3306	1/1	0.94	0.21	41,41,41,41	0
56	MG	1A	3574	1/1	0.94	0.10	48,48,48,48	0
56	MG	1A	3229	1/1	0.94	0.11	50,50,50,50	0
56	MG	2a	3126	1/1	0.94	0.18	41,41,41,41	0
56	MG	1A	3726	1/1	0.94	0.20	64,64,64,64	0
56	MG	1a	1660	1/1	0.94	0.24	50,50,50,50	0
56	MG	1A	3484	1/1	0.94	0.14	48,48,48,48	0
56	MG	1A	3308	1/1	0.94	0.20	50,50,50,50	0
56	MG	1A	3096	1/1	0.94	0.14	50,50,50,50	0
56	MG	1A	3869	1/1	0.94	0.08	42,42,42,42	0
56	MG	1A	3870	1/1	0.94	0.12	53,53,53,53	0
56	MG	1B	218	1/1	0.94	0.07	39,39,39,39	0
56	MG	1A	3127	1/1	0.94	0.07	53,53,53,53	0
56	MG	1B	220	1/1	0.94	0.10	70,70,70,70	0
56	MG	1A	3872	1/1	0.94	0.09	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	1a	1850	1/1	0.94	0.13	59,59,59,59	0
56	MG	1a	1851	1/1	0.94	0.20	74,74,74,74	0
56	MG	2A	3165	1/1	0.94	0.28	49,49,49,49	0
56	MG	1A	3069	1/1	0.94	0.10	39,39,39,39	0
56	MG	2a	3146	1/1	0.94	0.12	57,57,57,57	0
56	MG	2A	3170	1/1	0.94	0.06	42,42,42,42	0
56	MG	1a	1678	1/1	0.94	0.08	43,43,43,43	0
56	MG	1a	1855	1/1	0.94	0.15	55,55,55,55	0
56	MG	2A	3175	1/1	0.94	0.22	52,52,52,52	0
56	MG	1a	1679	1/1	0.94	0.24	57,57,57,57	0
56	MG	1a	1858	1/1	0.94	0.24	52,52,52,52	0
56	MG	1a	1859	1/1	0.94	0.07	59,59,59,59	0
56	MG	1A	3736	1/1	0.94	0.11	38,38,38,38	0
56	MG	1a	1681	1/1	0.94	0.20	55,55,55,55	0
56	MG	1a	1862	1/1	0.94	0.11	65,65,65,65	0
56	MG	2A	3188	1/1	0.94	0.21	35,35,35,35	0
56	MG	2A	3190	1/1	0.94	0.17	54,54,54,54	0
56	MG	1a	1863	1/1	0.94	0.14	51,51,51,51	0
56	MG	1A	3313	1/1	0.94	0.24	53,53,53,53	0
56	MG	1A	3877	1/1	0.94	0.08	42,42,42,42	0
56	MG	1A	3492	1/1	0.94	0.27	52,52,52,52	0
56	MG	1A	3741	1/1	0.94	0.12	19,19,19,19	0
56	MG	1a	1687	1/1	0.94	0.11	46,46,46,46	0
56	MG	1D	305	1/1	0.94	0.10	39,39,39,39	0
56	MG	1a	1871	1/1	0.94	0.09	46,46,46,46	0
56	MG	2A	3207	1/1	0.94	0.09	44,44,44,44	0
56	MG	1D	309	1/1	0.94	0.14	41,41,41,41	0
56	MG	1a	1874	1/1	0.94	0.06	61,61,61,61	0
56	MG	1A	3585	1/1	0.94	0.18	51,51,51,51	0
56	MG	1A	3185	1/1	0.94	0.21	41,41,41,41	0
56	MG	2A	3494	1/1	0.94	0.07	56,56,56,56	0
56	MG	1A	3187	1/1	0.94	0.16	46,46,46,46	0
56	MG	1A	3316	1/1	0.94	0.23	48,48,48,48	0
56	MG	2a	3178	1/1	0.94	0.11	59,59,59,59	0
56	MG	1A	3594	1/1	0.94	0.16	62,62,62,62	0
56	MG	1A	3886	1/1	0.94	0.20	33,33,33,33	0
56	MG	2a	3181	1/1	0.94	0.21	47,47,47,47	0
56	MG	2a	3182	1/1	0.94	0.15	51,51,51,51	0
56	MG	1A	3888	1/1	0.94	0.26	42,42,42,42	0
56	MG	1A	3759	1/1	0.94	0.05	60,60,60,60	0
56	MG	1A	3129	1/1	0.94	0.16	40,40,40,40	0
56	MG	1A	3763	1/1	0.94	0.09	43,43,43,43	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3089	1/1	0.94	0.09	55,55,55,55	0
56	MG	1G	3001	1/1	0.94	0.16	48,48,48,48	0
56	MG	1o	3002	1/1	0.94	0.10	44,44,44,44	0
56	MG	1a	1707	1/1	0.94	0.23	67,67,67,67	0
56	MG	1A	3369	1/1	0.94	0.17	40,40,40,40	0
56	MG	1G	3003	1/1	0.94	0.05	45,45,45,45	0
56	MG	2a	3200	1/1	0.94	0.13	49,49,49,49	0
56	MG	2A	3237	1/1	0.94	0.11	41,41,41,41	0
56	MG	1A	3605	1/1	0.94	0.10	33,33,33,33	0
56	MG	1A	3501	1/1	0.94	0.18	56,56,56,56	0
56	MG	1a	1715	1/1	0.94	0.15	39,39,39,39	0
56	MG	1A	3900	1/1	0.94	0.08	66,66,66,66	0
56	MG	1A	3191	1/1	0.94	0.08	37,37,37,37	0
56	MG	2A	3246	1/1	0.94	0.14	36,36,36,36	0
56	MG	2A	3248	1/1	0.94	0.21	56,56,56,56	0
56	MG	2A	3521	1/1	0.94	0.10	44,44,44,44	0
56	MG	1A	3194	1/1	0.94	0.17	32,32,32,32	0
56	MG	1A	3906	1/1	0.94	0.10	49,49,49,49	0
56	MG	1O	3002	1/1	0.94	0.14	46,46,46,46	0
56	MG	1A	3244	1/1	0.94	0.21	67,67,67,67	0
56	MG	1A	3439	1/1	0.94	0.11	51,51,51,51	0
56	MG	1a	1727	1/1	0.94	0.29	54,54,54,54	0
56	MG	1A	3782	1/1	0.94	0.07	34,34,34,34	0
56	MG	2a	3218	1/1	0.94	0.15	62,62,62,62	0
56	MG	1P	205	1/1	0.94	0.36	31,31,31,31	0
56	MG	1A	3506	1/1	0.94	0.16	58,58,58,58	0
56	MG	1R	203	1/1	0.94	0.20	43,43,43,43	0
56	MG	2a	3224	1/1	0.94	0.12	54,54,54,54	0
56	MG	2A	3538	1/1	0.94	0.13	22,22,22,22	0
56	MG	2A	3261	1/1	0.94	0.12	42,42,42,42	0
56	MG	1A	3374	1/1	0.94	0.22	64,64,64,64	0
56	MG	1A	3150	1/1	0.94	0.07	34,34,34,34	0
56	MG	1A	3920	1/1	0.94	0.15	64,64,64,64	0
56	MG	2A	3009	1/1	0.94	0.07	60,60,60,60	0
56	MG	1A	3921	1/1	0.94	0.13	55,55,55,55	0
56	MG	2A	3271	1/1	0.94	0.10	27,27,27,27	0
56	MG	2A	3272	1/1	0.94	0.20	46,46,46,46	0
56	MG	2a	3236	1/1	0.94	0.17	62,62,62,62	0
56	MG	2A	3555	1/1	0.94	0.15	36,36,36,36	0
56	MG	1A	3249	1/1	0.94	0.17	42,42,42,42	0
56	MG	2A	3012	1/1	0.94	0.08	53,53,53,53	0
56	MG	2A	3561	1/1	0.94	0.21	41,41,41,41	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3512	1/1	0.94	0.12	60,60,60,60	0
56	MG	1A	3789	1/1	0.94	0.08	26,26,26,26	0
56	MG	1A	3630	1/1	0.94	0.14	32,32,32,32	0
56	MG	1A	3253	1/1	0.94	0.17	31,31,31,31	0
56	MG	2A	3280	1/1	0.94	0.08	63,63,63,63	0
56	MG	1A	3091	1/1	0.94	0.05	51,51,51,51	0
56	MG	2A	3019	1/1	0.94	0.26	46,46,46,46	0
56	MG	2A	3020	1/1	0.94	0.11	48,48,48,48	0
56	MG	2A	3021	1/1	0.94	0.13	58,58,58,58	0
56	MG	2B	3008	1/1	0.94	0.09	47,47,47,47	0
56	MG	2A	3023	1/1	0.94	0.16	61,61,61,61	0
56	MG	1W	3004	1/1	0.94	0.18	31,31,31,31	0
56	MG	1A	3795	1/1	0.94	0.12	64,64,64,64	0
56	MG	2A	3291	1/1	0.94	0.10	36,36,36,36	0
56	MG	1A	3118	1/1	0.94	0.09	68,68,68,68	0
56	MG	2A	3294	1/1	0.94	0.10	37,37,37,37	0
56	MG	2E	303	1/1	0.94	0.18	26,26,26,26	0
56	MG	2A	3029	1/1	0.94	0.08	48,48,48,48	0
56	MG	1A	3798	1/1	0.94	0.08	61,61,61,61	0
56	MG	1A	3803	1/1	0.94	0.10	29,29,29,29	0
56	MG	2A	3032	1/1	0.94	0.07	53,53,53,53	0
56	MG	2h	3001	1/1	0.94	0.20	62,62,62,62	0
56	MG	1A	3638	1/1	0.94	0.18	41,41,41,41	0
56	MG	2A	3034	1/1	0.94	0.16	54,54,54,54	0
56	MG	2O	201	1/1	0.94	0.12	48,48,48,48	0
56	MG	2A	3035	1/1	0.94	0.26	58,58,58,58	0
56	MG	2Q	3002	1/1	0.94	0.10	39,39,39,39	0
56	MG	1A	3257	1/1	0.94	0.15	44,44,44,44	0
56	MG	2A	3305	1/1	0.94	0.07	42,42,42,42	0
56	MG	2O	8001	1/1	0.94	0.13	65,65,65,65	0
56	MG	2z	101	1/1	0.94	0.11	54,54,54,54	0
56	MG	2A	3397	1/1	0.95	0.17	66,66,66,66	0
56	MG	1A	3086	1/1	0.95	0.10	44,44,44,44	0
56	MG	2a	3040	1/1	0.95	0.16	39,39,39,39	0
56	MG	2A	3399	1/1	0.95	0.13	31,31,31,31	0
56	MG	2A	3173	1/1	0.95	0.13	52,52,52,52	0
56	MG	2A	3174	1/1	0.95	0.10	45,45,45,45	0
56	MG	1A	3959	1/1	0.95	0.09	79,79,79,79	0
56	MG	1A	3411	1/1	0.95	0.14	62,62,62,62	0
56	MG	2A	3177	1/1	0.95	0.21	48,48,48,48	0
56	MG	1A	3611	1/1	0.95	0.11	58,58,58,58	0
56	MG	1A	3962	1/1	0.95	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	3258	1/1	0.95	0.17	36,36,36,36	0
56	MG	1A	3413	1/1	0.95	0.04	41,41,41,41	0
56	MG	2A	3410	1/1	0.95	0.29	56,56,56,56	0
56	MG	2A	3184	1/1	0.95	0.20	39,39,39,39	0
56	MG	2A	3412	1/1	0.95	0.12	63,63,63,63	0
56	MG	2A	3185	1/1	0.95	0.16	52,52,52,52	0
56	MG	1A	3735	1/1	0.95	0.07	36,36,36,36	0
56	MG	1A	3014	1/1	0.95	0.28	53,53,53,53	0
56	MG	1A	3531	1/1	0.95	0.13	33,33,33,33	0
56	MG	2A	3004	1/1	0.95	0.06	52,52,52,52	0
56	MG	2a	3059	1/1	0.95	0.08	83,83,83,83	0
56	MG	2A	3418	1/1	0.95	0.11	57,57,57,57	0
56	MG	1A	3739	1/1	0.95	0.09	30,30,30,30	0
56	MG	2A	3420	1/1	0.95	0.13	53,53,53,53	0
56	MG	2A	3196	1/1	0.95	0.17	35,35,35,35	0
56	MG	1A	3974	1/1	0.95	0.09	30,30,30,30	0
56	MG	1A	3975	1/1	0.95	0.30	45,45,45,45	0
56	MG	1A	3976	1/1	0.95	0.14	69,69,69,69	0
56	MG	1A	3532	1/1	0.95	0.11	62,62,62,62	0
56	MG	2A	3203	1/1	0.95	0.17	43,43,43,43	0
56	MG	1A	3848	1/1	0.95	0.06	61,61,61,61	0
56	MG	1A	3012	1/1	0.95	0.10	47,47,47,47	0
56	MG	2a	3071	1/1	0.95	0.17	59,59,59,59	0
56	MG	1A	3742	1/1	0.95	0.12	35,35,35,35	0
56	MG	1A	3851	1/1	0.95	0.22	62,62,62,62	0
56	MG	1A	3852	1/1	0.95	0.08	16,16,16,16	0
56	MG	2A	3435	1/1	0.95	0.08	53,53,53,53	0
56	MG	1A	3854	1/1	0.95	0.11	53,53,53,53	0
56	MG	1A	3744	1/1	0.95	0.15	53,53,53,53	0
56	MG	1A	3535	1/1	0.95	0.10	41,41,41,41	0
56	MG	1A	3991	1/1	0.95	0.08	52,52,52,52	0
56	MG	1A	3628	1/1	0.95	0.11	28,28,28,28	0
56	MG	2A	3219	1/1	0.95	0.06	44,44,44,44	0
56	MG	2A	3220	1/1	0.95	0.21	45,45,45,45	0
56	MG	2a	3083	1/1	0.95	0.07	51,51,51,51	0
56	MG	1A	3536	1/1	0.95	0.11	68,68,68,68	0
56	MG	2A	3223	1/1	0.95	0.13	50,50,50,50	0
56	MG	2A	3022	1/1	0.95	0.06	51,51,51,51	0
56	MG	1A	3364	1/1	0.95	0.06	44,44,44,44	0
56	MG	1A	3861	1/1	0.95	0.17	48,48,48,48	0
56	MG	1A	3862	1/1	0.95	0.17	62,62,62,62	0
56	MG	1A	3753	1/1	0.95	0.10	68,68,68,68	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3092	1/1	0.95	0.14	49,49,49,49	0
56	MG	2A	3230	1/1	0.95	0.15	47,47,47,47	0
56	MG	1A	3758	1/1	0.95	0.09	65,65,65,65	0
56	MG	2a	3097	1/1	0.95	0.32	71,71,71,71	0
56	MG	1A	3263	1/1	0.95	0.10	37,37,37,37	0
56	MG	1A	4005	1/1	0.95	0.15	58,58,58,58	0
56	MG	1A	4006	1/1	0.95	0.10	45,45,45,45	0
56	MG	2A	3236	1/1	0.95	0.06	91,91,91,91	0
56	MG	1a	1622	1/1	0.95	0.10	90,90,90,90	0
56	MG	2a	3105	1/1	0.95	0.07	55,55,55,55	0
56	MG	2A	3464	1/1	0.95	0.06	50,50,50,50	0
56	MG	2A	3238	1/1	0.95	0.06	60,60,60,60	0
56	MG	1A	3221	1/1	0.95	0.06	37,37,37,37	0
56	MG	1A	3542	1/1	0.95	0.11	26,26,26,26	0
56	MG	1A	3766	1/1	0.95	0.10	49,49,49,49	0
56	MG	1A	3767	1/1	0.95	0.10	40,40,40,40	0
56	MG	1A	3641	1/1	0.95	0.07	39,39,39,39	0
56	MG	1A	3644	1/1	0.95	0.10	19,19,19,19	0
56	MG	2A	3247	1/1	0.95	0.08	59,59,59,59	0
56	MG	2A	3477	1/1	0.95	0.08	57,57,57,57	0
56	MG	1A	3266	1/1	0.95	0.09	44,44,44,44	0
56	MG	1A	3051	1/1	0.95	0.09	50,50,50,50	0
56	MG	1A	3773	1/1	0.95	0.22	80,80,80,80	0
56	MG	1a	1635	1/1	0.95	0.23	50,50,50,50	0
56	MG	1A	4020	1/1	0.95	0.08	32,32,32,32	0
56	MG	1A	3370	1/1	0.95	0.08	66,66,66,66	0
56	MG	1a	1794	1/1	0.95	0.07	67,67,67,67	0
56	MG	1A	3777	1/1	0.95	0.07	49,49,49,49	0
56	MG	1B	202	1/1	0.95	0.08	52,52,52,52	0
56	MG	1A	3054	1/1	0.95	0.14	47,47,47,47	0
56	MG	2a	3130	1/1	0.95	0.18	43,43,43,43	0
56	MG	1A	3653	1/1	0.95	0.08	25,25,25,25	0
56	MG	2A	3056	1/1	0.95	0.13	38,38,38,38	0
56	MG	2a	3133	1/1	0.95	0.15	60,60,60,60	0
56	MG	1A	3149	1/1	0.95	0.15	55,55,55,55	0
56	MG	2A	3058	1/1	0.95	0.13	37,37,37,37	0
56	MG	1A	3428	1/1	0.95	0.10	65,65,65,65	0
56	MG	1A	3885	1/1	0.95	0.09	53,53,53,53	0
56	MG	2A	3267	1/1	0.95	0.14	39,39,39,39	0
56	MG	1A	3135	1/1	0.95	0.13	64,64,64,64	0
56	MG	2A	3063	1/1	0.95	0.17	45,45,45,45	0
56	MG	2A	3064	1/1	0.95	0.18	56,56,56,56	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1B	210	1/1	0.95	0.05	49,49,49,49	0
56	MG	1A	3887	1/1	0.95	0.21	31,31,31,31	0
56	MG	2A	3068	1/1	0.95	0.19	39,39,39,39	0
56	MG	2A	3069	1/1	0.95	0.12	37,37,37,37	0
56	MG	2A	3277	1/1	0.95	0.18	41,41,41,41	0
56	MG	1A	3553	1/1	0.95	0.12	44,44,44,44	0
56	MG	1a	1651	1/1	0.95	0.06	54,54,54,54	0
56	MG	2A	3072	1/1	0.95	0.22	48,48,48,48	0
56	MG	1A	3063	1/1	0.95	0.07	53,53,53,53	0
56	MG	1a	1809	1/1	0.95	0.16	62,62,62,62	0
56	MG	1A	3230	1/1	0.95	0.13	46,46,46,46	0
56	MG	1A	3280	1/1	0.95	0.06	34,34,34,34	0
56	MG	1a	1656	1/1	0.95	0.19	57,57,57,57	0
56	MG	2A	3286	1/1	0.95	0.11	32,32,32,32	0
56	MG	1A	3231	1/1	0.95	0.30	49,49,49,49	0
56	MG	1A	3667	1/1	0.95	0.09	49,49,49,49	0
56	MG	1a	1659	1/1	0.95	0.12	63,63,63,63	0
56	MG	1A	3153	1/1	0.95	0.06	24,24,24,24	0
56	MG	1A	3283	1/1	0.95	0.09	44,44,44,44	0
56	MG	1A	3154	1/1	0.95	0.09	28,28,28,28	0
56	MG	1a	1821	1/1	0.95	0.06	49,49,49,49	0
56	MG	1a	1666	1/1	0.95	0.09	71,71,71,71	0
56	MG	2A	3527	1/1	0.95	0.23	51,51,51,51	0
56	MG	1A	3157	1/1	0.95	0.12	28,28,28,28	0
56	MG	1D	303	1/1	0.95	0.12	47,47,47,47	0
56	MG	1A	3904	1/1	0.95	0.09	27,27,27,27	0
56	MG	1A	3674	1/1	0.95	0.07	52,52,52,52	0
56	MG	2A	3532	1/1	0.95	0.14	51,51,51,51	0
56	MG	1a	1829	1/1	0.95	0.21	47,47,47,47	0
56	MG	2A	3534	1/1	0.95	0.09	64,64,64,64	0
56	MG	2A	3535	1/1	0.95	0.11	44,44,44,44	0
56	MG	2A	3536	1/1	0.95	0.10	38,38,38,38	0
56	MG	2A	3304	1/1	0.95	0.13	41,41,41,41	0
56	MG	1A	3801	1/1	0.95	0.08	42,42,42,42	0
56	MG	1D	310	1/1	0.95	0.07	57,57,57,57	0
56	MG	1a	1675	1/1	0.95	0.06	56,56,56,56	0
56	MG	1A	3289	1/1	0.95	0.22	43,43,43,43	0
56	MG	1a	1835	1/1	0.95	0.14	73,73,73,73	0
56	MG	1D	312	1/1	0.95	0.07	47,47,47,47	0
56	MG	2a	3186	1/1	0.95	0.14	48,48,48,48	0
56	MG	2a	3187	1/1	0.95	0.06	59,59,59,59	0
56	MG	2A	3099	1/1	0.95	0.14	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	1A	3910	1/1	0.95	0.17	65,65,65,65	0
56	MG	1D	314	1/1	0.95	0.12	63,63,63,63	0
56	MG	1A	3290	1/1	0.95	0.17	37,37,37,37	0
56	MG	1A	3912	1/1	0.95	0.09	40,40,40,40	0
56	MG	2A	3558	1/1	0.95	0.07	34,34,34,34	0
56	MG	2A	3559	1/1	0.95	0.08	54,54,54,54	0
56	MG	1a	1682	1/1	0.95	0.18	46,46,46,46	0
56	MG	1A	3337	1/1	0.95	0.13	57,57,57,57	0
56	MG	1A	3916	1/1	0.95	0.17	52,52,52,52	0
56	MG	2A	3323	1/1	0.95	0.09	48,48,48,48	0
56	MG	1a	1847	1/1	0.95	0.04	53,53,53,53	0
56	MG	1A	3158	1/1	0.95	0.12	24,24,24,24	0
56	MG	1A	3389	1/1	0.95	0.22	45,45,45,45	0
56	MG	1A	3192	1/1	0.95	0.05	35,35,35,35	0
56	MG	1A	3296	1/1	0.95	0.23	55,55,55,55	0
56	MG	1a	1853	1/1	0.95	0.22	57,57,57,57	0
56	MG	1A	3193	1/1	0.95	0.15	23,23,23,23	0
56	MG	1A	3922	1/1	0.95	0.09	19,19,19,19	0
56	MG	1A	3298	1/1	0.95	0.19	50,50,50,50	0
56	MG	1A	3159	1/1	0.95	0.06	35,35,35,35	0
56	MG	1A	3695	1/1	0.95	0.18	41,41,41,41	0
56	MG	1A	3072	1/1	0.95	0.14	56,56,56,56	0
56	MG	1A	3242	1/1	0.95	0.07	33,33,33,33	0
56	MG	2E	302	1/1	0.95	0.10	59,59,59,59	0
56	MG	2A	3123	1/1	0.95	0.14	41,41,41,41	0
56	MG	1A	3139	1/1	0.95	0.16	41,41,41,41	0
56	MG	1A	3701	1/1	0.95	0.11	68,68,68,68	0
56	MG	2A	3343	1/1	0.95	0.12	56,56,56,56	0
56	MG	2F	303	1/1	0.95	0.12	50,50,50,50	0
56	MG	1A	3304	1/1	0.95	0.08	34,34,34,34	0
56	MG	1O	3001	1/1	0.95	0.06	45,45,45,45	0
56	MG	2A	3128	1/1	0.95	0.24	60,60,60,60	0
56	MG	1a	1866	1/1	0.95	0.16	61,61,61,61	0
56	MG	1A	3704	1/1	0.95	0.08	18,18,18,18	0
56	MG	2Q	3001	1/1	0.95	0.22	48,48,48,48	0
56	MG	2A	3354	1/1	0.95	0.09	63,63,63,63	0
56	MG	1A	3305	1/1	0.95	0.09	43,43,43,43	0
56	MG	2A	3132	1/1	0.95	0.15	53,53,53,53	0
56	MG	1A	3706	1/1	0.95	0.05	22,22,22,22	0
56	MG	1A	3028	1/1	0.95	0.11	46,46,46,46	0
56	MG	1A	3591	1/1	0.95	0.09	77,77,77,77	0
56	MG	2a	3001	1/1	0.95	0.30	60,60,60,60	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1Q	3001	1/1	0.95	0.24	38,38,38,38	0
56	MG	1A	3202	1/1	0.95	0.28	43,43,43,43	0
56	MG	1R	202	1/1	0.95	0.10	52,52,52,52	0
56	MG	1A	3940	1/1	0.95	0.16	63,63,63,63	0
56	MG	1A	3205	1/1	0.95	0.13	41,41,41,41	0
56	MG	2a	3007	1/1	0.95	0.26	46,46,46,46	0
56	MG	1a	1719	1/1	0.95	0.17	53,53,53,53	0
56	MG	1A	3252	1/1	0.95	0.07	28,28,28,28	0
56	MG	2A	3368	1/1	0.95	0.15	29,29,29,29	0
56	MG	1A	3598	1/1	0.95	0.18	51,51,51,51	0
56	MG	2A	3370	1/1	0.95	0.13	44,44,44,44	0
56	MG	2A	3147	1/1	0.95	0.20	42,42,42,42	0
56	MG	1A	3717	1/1	0.95	0.10	59,59,59,59	0
56	MG	2A	3149	1/1	0.95	0.19	43,43,43,43	0
56	MG	1A	3010	1/1	0.95	0.21	42,42,42,42	0
56	MG	1A	3946	1/1	0.95	0.13	41,41,41,41	0
56	MG	1A	3126	1/1	0.95	0.20	53,53,53,53	0
56	MG	2A	3378	1/1	0.95	0.13	66,66,66,66	0
56	MG	2A	3379	1/1	0.95	0.07	42,42,42,42	0
56	MG	1U	204	1/1	0.95	0.43	42,42,42,42	0
56	MG	1A	3210	1/1	0.95	0.23	43,43,43,43	0
56	MG	1W	3001	1/1	0.95	0.20	34,34,34,34	0
56	MG	1A	3950	1/1	0.95	0.14	41,41,41,41	0
56	MG	1a	1735	1/1	0.95	0.12	66,66,66,66	0
56	MG	1A	3952	1/1	0.95	0.05	51,51,51,51	0
56	MG	1A	3524	1/1	0.95	0.16	54,54,54,54	0
56	MG	1A	3954	1/1	0.95	0.17	51,51,51,51	0
56	MG	2A	3167	1/1	0.95	0.18	25,25,25,25	0
56	MG	2A	3391	1/1	0.95	0.12	48,48,48,48	0
56	MG	2x	3001	1/1	0.95	0.13	66,66,66,66	0
56	MG	2A	3392	1/1	0.95	0.12	28,28,28,28	0
56	MG	1A	3727	1/1	0.95	0.11	26,26,26,26	0
56	MG	2x	3004	1/1	0.95	0.13	42,42,42,42	0
56	MG	1A	3956	1/1	0.95	0.07	53,53,53,53	0
57	ZN	24	501	1/1	0.95	0.06	125,125,125,125	0
56	MG	1A	3279	1/1	0.96	0.20	28,28,28,28	0
56	MG	2A	3482	1/1	0.96	0.05	60,60,60,60	0
56	MG	1d	503	1/1	0.96	0.16	52,52,52,52	0
56	MG	2A	3297	1/1	0.96	0.11	58,58,58,58	0
56	MG	18	103	1/1	0.96	0.10	52,52,52,52	0
56	MG	19	102	1/1	0.96	0.12	51,51,51,51	0
56	MG	1A	3696	1/1	0.96	0.13	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	3593	1/1	0.96	0.12	35,35,35,35	0
56	MG	1A	3198	1/1	0.96	0.18	28,28,28,28	0
56	MG	1A	3595	1/1	0.96	0.23	56,56,56,56	0
56	MG	1A	3042	1/1	0.96	0.30	44,44,44,44	0
56	MG	1A	3326	1/1	0.96	0.07	53,53,53,53	0
56	MG	1A	3465	1/1	0.96	0.08	44,44,44,44	0
56	MG	2A	3307	1/1	0.96	0.14	35,35,35,35	0
56	MG	1A	3601	1/1	0.96	0.12	44,44,44,44	0
56	MG	1A	3415	1/1	0.96	0.06	50,50,50,50	0
56	MG	2A	3498	1/1	0.96	0.06	49,49,49,49	0
56	MG	1A	3131	1/1	0.96	0.16	42,42,42,42	0
56	MG	2a	3090	1/1	0.96	0.10	42,42,42,42	0
56	MG	1A	3913	1/1	0.96	0.09	64,64,64,64	0
56	MG	1a	1612	1/1	0.96	0.04	65,65,65,65	0
56	MG	1A	3708	1/1	0.96	0.10	63,63,63,63	0
56	MG	1x	101	1/1	0.96	0.09	51,51,51,51	0
56	MG	2A	3139	1/1	0.96	0.08	45,45,45,45	0
56	MG	2A	3316	1/1	0.96	0.15	57,57,57,57	0
56	MG	2A	3317	1/1	0.96	0.07	34,34,34,34	0
56	MG	1A	3117	1/1	0.96	0.07	48,48,48,48	0
56	MG	1A	3203	1/1	0.96	0.19	32,32,32,32	0
56	MG	2A	3142	1/1	0.96	0.16	50,50,50,50	0
56	MG	2A	3143	1/1	0.96	0.23	48,48,48,48	0
56	MG	1a	1616	1/1	0.96	0.06	74,74,74,74	0
56	MG	2A	3324	1/1	0.96	0.06	34,34,34,34	0
56	MG	2a	3107	1/1	0.96	0.11	57,57,57,57	0
56	MG	1A	3609	1/1	0.96	0.07	22,22,22,22	0
56	MG	1A	3713	1/1	0.96	0.10	40,40,40,40	0
56	MG	1a	1619	1/1	0.96	0.06	54,54,54,54	0
56	MG	2A	3516	1/1	0.96	0.20	59,59,59,59	0
56	MG	2A	3517	1/1	0.96	0.17	56,56,56,56	0
56	MG	1A	3204	1/1	0.96	0.21	37,37,37,37	0
56	MG	1A	3053	1/1	0.96	0.27	53,53,53,53	0
56	MG	1B	215	1/1	0.96	0.06	31,31,31,31	0
56	MG	1A	3473	1/1	0.96	0.23	50,50,50,50	0
56	MG	1A	3422	1/1	0.96	0.06	51,51,51,51	0
56	MG	2a	3119	1/1	0.96	0.29	56,56,56,56	0
56	MG	2A	3155	1/1	0.96	0.17	33,33,33,33	0
56	MG	1A	3615	1/1	0.96	0.09	39,39,39,39	0
56	MG	1A	3616	1/1	0.96	0.07	22,22,22,22	0
56	MG	2A	3337	1/1	0.96	0.27	46,46,46,46	0
56	MG	1A	3825	1/1	0.96	0.07	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	3160	1/1	0.96	0.24	52,52,52,52	0
56	MG	1A	3134	1/1	0.96	0.15	45,45,45,45	0
56	MG	1A	3476	1/1	0.96	0.10	53,53,53,53	0
56	MG	1A	3207	1/1	0.96	0.05	30,30,30,30	0
56	MG	1a	1631	1/1	0.96	0.12	36,36,36,36	0
56	MG	1A	3376	1/1	0.96	0.10	53,53,53,53	0
56	MG	1a	1634	1/1	0.96	0.10	52,52,52,52	0
56	MG	2A	3168	1/1	0.96	0.07	47,47,47,47	0
56	MG	2A	3169	1/1	0.96	0.07	33,33,33,33	0
56	MG	1A	3245	1/1	0.96	0.14	49,49,49,49	0
56	MG	1A	3626	1/1	0.96	0.09	25,25,25,25	0
56	MG	1A	3175	1/1	0.96	0.18	35,35,35,35	0
56	MG	2A	3017	1/1	0.96	0.18	47,47,47,47	0
56	MG	1D	306	1/1	0.96	0.18	31,31,31,31	0
56	MG	1A	3247	1/1	0.96	0.19	43,43,43,43	0
56	MG	2A	3547	1/1	0.96	0.13	22,22,22,22	0
56	MG	2a	3142	1/1	0.96	0.13	51,51,51,51	0
56	MG	2A	3550	1/1	0.96	0.16	42,42,42,42	0
56	MG	1a	1641	1/1	0.96	0.09	55,55,55,55	0
56	MG	1A	3632	1/1	0.96	0.07	25,25,25,25	0
56	MG	1A	3248	1/1	0.96	0.09	26,26,26,26	0
56	MG	1A	3483	1/1	0.96	0.10	54,54,54,54	0
56	MG	2A	3181	1/1	0.96	0.12	39,39,39,39	0
56	MG	1A	3636	1/1	0.96	0.08	43,43,43,43	0
56	MG	1A	3073	1/1	0.96	0.14	51,51,51,51	0
56	MG	2A	3560	1/1	0.96	0.10	31,31,31,31	0
56	MG	1A	3251	1/1	0.96	0.13	30,30,30,30	0
56	MG	1A	3743	1/1	0.96	0.19	57,57,57,57	0
56	MG	2a	3154	1/1	0.96	0.15	87,87,87,87	0
56	MG	1E	304	1/1	0.96	0.15	57,57,57,57	0
56	MG	1E	305	1/1	0.96	0.13	31,31,31,31	0
56	MG	1A	3211	1/1	0.96	0.27	58,58,58,58	0
56	MG	2A	3192	1/1	0.96	0.13	29,29,29,29	0
56	MG	1A	3341	1/1	0.96	0.05	64,64,64,64	0
56	MG	1A	3747	1/1	0.96	0.09	29,29,29,29	0
56	MG	1A	3642	1/1	0.96	0.09	58,58,58,58	0
56	MG	2B	3005	1/1	0.96	0.28	53,53,53,53	0
56	MG	1A	3550	1/1	0.96	0.12	33,33,33,33	0
56	MG	2B	3007	1/1	0.96	0.20	56,56,56,56	0
56	MG	2A	3377	1/1	0.96	0.11	43,43,43,43	0
56	MG	1A	3302	1/1	0.96	0.21	41,41,41,41	0
56	MG	1A	3435	1/1	0.96	0.07	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	1799	1/1	0.96	0.06	61,61,61,61	0
56	MG	2D	301	1/1	0.96	0.17	57,57,57,57	0
56	MG	1A	3755	1/1	0.96	0.05	66,66,66,66	0
56	MG	2A	3382	1/1	0.96	0.07	42,42,42,42	0
56	MG	2D	305	1/1	0.96	0.10	28,28,28,28	0
56	MG	1A	3213	1/1	0.96	0.21	33,33,33,33	0
56	MG	1a	1662	1/1	0.96	0.05	61,61,61,61	0
56	MG	1A	3853	1/1	0.96	0.19	46,46,46,46	0
56	MG	1a	1664	1/1	0.96	0.14	61,61,61,61	0
56	MG	1A	3254	1/1	0.96	0.13	34,34,34,34	0
56	MG	1A	3652	1/1	0.96	0.08	36,36,36,36	0
56	MG	1A	3495	1/1	0.96	0.06	66,66,66,66	0
56	MG	1A	3654	1/1	0.96	0.11	67,67,67,67	0
56	MG	2a	3183	1/1	0.96	0.14	37,37,37,37	0
56	MG	2F	305	1/1	0.96	0.09	41,41,41,41	0
56	MG	2A	3051	1/1	0.96	0.14	40,40,40,40	0
56	MG	2A	3393	1/1	0.96	0.10	63,63,63,63	0
56	MG	1N	3004	1/1	0.96	0.06	36,36,36,36	0
56	MG	2a	3189	1/1	0.96	0.24	51,51,51,51	0
56	MG	1A	3177	1/1	0.96	0.14	39,39,39,39	0
56	MG	1A	3152	1/1	0.96	0.15	50,50,50,50	0
56	MG	1A	3965	1/1	0.96	0.13	53,53,53,53	0
56	MG	2a	3193	1/1	0.96	0.12	47,47,47,47	0
56	MG	2Q	3005	1/1	0.96	0.07	52,52,52,52	0
56	MG	2U	201	1/1	0.96	0.11	48,48,48,48	0
56	MG	1A	3348	1/1	0.96	0.09	47,47,47,47	0
56	MG	1A	3094	1/1	0.96	0.11	51,51,51,51	0
56	MG	1A	3969	1/1	0.96	0.09	18,18,18,18	0
56	MG	1A	3970	1/1	0.96	0.33	30,30,30,30	0
56	MG	1A	3084	1/1	0.96	0.11	50,50,50,50	0
56	MG	1A	3155	1/1	0.96	0.04	37,37,37,37	0
56	MG	1Q	3002	1/1	0.96	0.10	44,44,44,44	0
56	MG	1A	3866	1/1	0.96	0.04	37,37,37,37	0
56	MG	1A	3186	1/1	0.96	0.20	32,32,32,32	0
56	MG	2A	3067	1/1	0.96	0.13	52,52,52,52	0
56	MG	1A	3774	1/1	0.96	0.06	20,20,20,20	0
56	MG	1A	3223	1/1	0.96	0.21	33,33,33,33	0
56	MG	1A	3085	1/1	0.96	0.10	47,47,47,47	0
56	MG	1A	3400	1/1	0.96	0.08	45,45,45,45	0
56	MG	1A	3979	1/1	0.96	0.10	49,49,49,49	0
56	MG	1A	3097	1/1	0.96	0.06	55,55,55,55	0
56	MG	1A	3507	1/1	0.96	0.25	64,64,64,64	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1a	1692	1/1	0.96	0.21	57,57,57,57	0
56	MG	2A	3242	1/1	0.96	0.15	52,52,52,52	0
56	MG	2A	3243	1/1	0.96	0.12	63,63,63,63	0
56	MG	1A	3781	1/1	0.96	0.06	40,40,40,40	0
56	MG	1U	203	1/1	0.96	0.10	47,47,47,47	0
56	MG	1A	3983	1/1	0.96	0.15	55,55,55,55	0
56	MG	1a	1697	1/1	0.96	0.10	52,52,52,52	0
56	MG	2a	3020	1/1	0.96	0.21	49,49,49,49	0
56	MG	1A	3402	1/1	0.96	0.09	58,58,58,58	0
56	MG	2a	3022	1/1	0.96	0.08	73,73,73,73	0
56	MG	2A	3249	1/1	0.96	0.07	59,59,59,59	0
56	MG	1V	201	1/1	0.96	0.19	45,45,45,45	0
56	MG	2a	3229	1/1	0.96	0.15	51,51,51,51	0
56	MG	1A	3985	1/1	0.96	0.19	66,66,66,66	0
56	MG	1A	3017	1/1	0.96	0.08	40,40,40,40	0
56	MG	2a	3232	1/1	0.96	0.14	66,66,66,66	0
56	MG	1a	1845	1/1	0.96	0.08	46,46,46,46	0
56	MG	1A	3579	1/1	0.96	0.04	45,45,45,45	0
56	MG	1A	3087	1/1	0.96	0.13	57,57,57,57	0
56	MG	1a	1848	1/1	0.96	0.07	57,57,57,57	0
56	MG	1A	3162	1/1	0.96	0.12	38,38,38,38	0
56	MG	1A	3112	1/1	0.96	0.19	40,40,40,40	0
56	MG	2A	3090	1/1	0.96	0.14	45,45,45,45	0
56	MG	1A	3788	1/1	0.96	0.06	23,23,23,23	0
56	MG	1A	3680	1/1	0.96	0.10	42,42,42,42	0
56	MG	1A	3996	1/1	0.96	0.04	40,40,40,40	0
56	MG	2a	3243	1/1	0.96	0.07	67,67,67,67	0
56	MG	10	103	1/1	0.96	0.11	75,75,75,75	0
56	MG	1A	3514	1/1	0.96	0.09	53,53,53,53	0
56	MG	1a	1714	1/1	0.96	0.11	68,68,68,68	0
56	MG	2A	3268	1/1	0.96	0.24	55,55,55,55	0
56	MG	1A	3584	1/1	0.96	0.08	33,33,33,33	0
56	MG	2a	3249	1/1	0.96	0.12	61,61,61,61	0
56	MG	2A	3270	1/1	0.96	0.08	45,45,45,45	0
56	MG	2a	3251	1/1	0.96	0.15	57,57,57,57	0
56	MG	1A	3793	1/1	0.96	0.06	40,40,40,40	0
56	MG	1A	3232	1/1	0.96	0.23	43,43,43,43	0
56	MG	1A	3088	1/1	0.96	0.23	60,60,60,60	0
56	MG	1a	1720	1/1	0.96	0.07	50,50,50,50	0
56	MG	1A	4002	1/1	0.96	0.29	34,34,34,34	0
56	MG	1a	1722	1/1	0.96	0.06	54,54,54,54	0
56	MG	2A	3457	1/1	0.96	0.10	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	1A	4003	1/1	0.96	0.24	43,43,43,43	0
56	MG	1A	3687	1/1	0.96	0.10	48,48,48,48	0
56	MG	1A	3277	1/1	0.96	0.13	48,48,48,48	0
56	MG	2A	3108	1/1	0.96	0.17	46,46,46,46	0
56	MG	2A	3462	1/1	0.96	0.07	60,60,60,60	0
56	MG	2A	3109	1/1	0.96	0.05	64,64,64,64	0
56	MG	15	102	1/1	0.96	0.21	40,40,40,40	0
56	MG	1A	3800	1/1	0.96	0.08	50,50,50,50	0
56	MG	1a	1728	1/1	0.96	0.18	53,53,53,53	0
56	MG	2l	203	1/1	0.96	0.17	60,60,60,60	0
56	MG	1A	3039	1/1	0.96	0.10	49,49,49,49	0
56	MG	2A	3470	1/1	0.96	0.08	44,44,44,44	0
56	MG	2A	3472	1/1	0.96	0.06	39,39,39,39	0
56	MG	16	101	1/1	0.96	0.12	45,45,45,45	0
56	MG	1A	4008	1/1	0.96	0.41	48,48,48,48	0
56	MG	1A	3893	1/1	0.96	0.10	30,30,30,30	0
56	MG	1a	1734	1/1	0.96	0.19	45,45,45,45	0
56	MG	1A	3802	1/1	0.96	0.10	77,77,77,77	0
56	MG	2A	3293	1/1	0.96	0.08	62,62,62,62	0
56	MG	1a	1736	1/1	0.96	0.26	41,41,41,41	0
57	ZN	2n	501	1/1	0.96	0.06	98,98,98,98	0
56	MG	1a	1815	1/1	0.97	0.11	58,58,58,58	0
56	MG	1A	3589	1/1	0.97	0.10	29,29,29,29	0
56	MG	2A	3024	1/1	0.97	0.20	43,43,43,43	0
56	MG	1a	1712	1/1	0.97	0.12	25,25,25,25	0
56	MG	1a	1713	1/1	0.97	0.10	63,63,63,63	0
56	MG	1A	3184	1/1	0.97	0.11	52,52,52,52	0
56	MG	2A	3028	1/1	0.97	0.21	52,52,52,52	0
56	MG	1A	3103	1/1	0.97	0.21	50,50,50,50	0
56	MG	1A	3737	1/1	0.97	0.11	24,24,24,24	0
56	MG	1A	3658	1/1	0.97	0.07	18,18,18,18	0
56	MG	1A	3659	1/1	0.97	0.04	23,23,23,23	0
56	MG	2A	3422	1/1	0.97	0.07	57,57,57,57	0
56	MG	2a	3128	1/1	0.97	0.19	38,38,38,38	0
56	MG	1a	1826	1/1	0.97	0.09	36,36,36,36	0
56	MG	1A	3540	1/1	0.97	0.10	45,45,45,45	0
56	MG	1A	3016	1/1	0.97	0.06	61,61,61,61	0
56	MG	2A	3150	1/1	0.97	0.15	30,30,30,30	0
56	MG	2E	301	1/1	0.97	0.18	44,44,44,44	0
56	MG	1A	3284	1/1	0.97	0.07	41,41,41,41	0
56	MG	2A	3428	1/1	0.97	0.20	49,49,49,49	0
56	MG	1A	3344	1/1	0.97	0.05	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	1A	3596	1/1	0.97	0.08	49,49,49,49	0
56	MG	2A	3154	1/1	0.97	0.13	32,32,32,32	0
56	MG	2A	3288	1/1	0.97	0.07	57,57,57,57	0
56	MG	1A	3544	1/1	0.97	0.06	36,36,36,36	0
56	MG	2A	3290	1/1	0.97	0.07	49,49,49,49	0
56	MG	2F	306	1/1	0.97	0.08	40,40,40,40	0
56	MG	1A	3746	1/1	0.97	0.09	7,7,7,7	0
56	MG	1A	3896	1/1	0.97	0.06	19,19,19,19	0
56	MG	1A	3020	1/1	0.97	0.21	54,54,54,54	0
56	MG	2O	203	1/1	0.97	0.07	40,40,40,40	0
56	MG	2A	3439	1/1	0.97	0.11	68,68,68,68	0
56	MG	2A	3159	1/1	0.97	0.11	37,37,37,37	0
56	MG	1a	1838	1/1	0.97	0.14	66,66,66,66	0
56	MG	2Q	3003	1/1	0.97	0.12	41,41,41,41	0
56	MG	2Q	3004	1/1	0.97	0.25	42,42,42,42	0
56	MG	1A	3899	1/1	0.97	0.07	47,47,47,47	0
56	MG	1a	1730	1/1	0.97	0.07	70,70,70,70	0
56	MG	2A	3444	1/1	0.97	0.10	29,29,29,29	0
56	MG	1N	3001	1/1	0.97	0.21	42,42,42,42	0
56	MG	2A	3047	1/1	0.97	0.45	65,65,65,65	0
56	MG	25	502	1/1	0.97	0.31	47,47,47,47	0
56	MG	1A	3748	1/1	0.97	0.06	37,37,37,37	0
56	MG	1A	3986	1/1	0.97	0.25	42,42,42,42	0
56	MG	2A	3050	1/1	0.97	0.09	38,38,38,38	0
56	MG	1A	3902	1/1	0.97	0.05	45,45,45,45	0
56	MG	1a	1632	1/1	0.97	0.10	41,41,41,41	0
56	MG	1A	3669	1/1	0.97	0.11	46,46,46,46	0
56	MG	1A	3599	1/1	0.97	0.12	37,37,37,37	0
56	MG	2A	3055	1/1	0.97	0.07	40,40,40,40	0
56	MG	1A	3116	1/1	0.97	0.08	41,41,41,41	0
56	MG	1A	3029	1/1	0.97	0.24	54,54,54,54	0
56	MG	1A	3908	1/1	0.97	0.09	40,40,40,40	0
56	MG	1P	202	1/1	0.97	0.10	55,55,55,55	0
56	MG	1A	3994	1/1	0.97	0.13	47,47,47,47	0
56	MG	1a	1640	1/1	0.97	0.27	51,51,51,51	0
56	MG	2A	3180	1/1	0.97	0.15	37,37,37,37	0
56	MG	1A	3754	1/1	0.97	0.05	18,18,18,18	0
56	MG	2a	3174	1/1	0.97	0.23	69,69,69,69	0
56	MG	1A	3602	1/1	0.97	0.11	45,45,45,45	0
56	MG	2A	3466	1/1	0.97	0.07	69,69,69,69	0
56	MG	2a	3177	1/1	0.97	0.11	44,44,44,44	0
56	MG	1P	206	1/1	0.97	0.18	54,54,54,54	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3438	1/1	0.97	0.07	58,58,58,58	0
56	MG	1A	3171	1/1	0.97	0.06	38,38,38,38	0
56	MG	2A	3320	1/1	0.97	0.09	33,33,33,33	0
56	MG	2A	3186	1/1	0.97	0.11	36,36,36,36	0
56	MG	1A	3259	1/1	0.97	0.05	38,38,38,38	0
56	MG	1R	201	1/1	0.97	0.29	63,63,63,63	0
56	MG	2A	3189	1/1	0.97	0.10	34,34,34,34	0
56	MG	1A	3914	1/1	0.97	0.11	38,38,38,38	0
56	MG	2A	3191	1/1	0.97	0.28	40,40,40,40	0
56	MG	2a	3188	1/1	0.97	0.11	60,60,60,60	0
56	MG	1A	3760	1/1	0.97	0.09	36,36,36,36	0
56	MG	2A	3193	1/1	0.97	0.12	32,32,32,32	0
56	MG	1A	3064	1/1	0.97	0.11	40,40,40,40	0
56	MG	1T	201	1/1	0.97	0.07	41,41,41,41	0
56	MG	1a	1652	1/1	0.97	0.06	48,48,48,48	0
56	MG	2a	3033	1/1	0.97	0.07	53,53,53,53	0
56	MG	1A	3836	1/1	0.97	0.06	35,35,35,35	0
56	MG	1A	3762	1/1	0.97	0.04	45,45,45,45	0
56	MG	1A	3294	1/1	0.97	0.05	23,23,23,23	0
56	MG	2A	3335	1/1	0.97	0.14	40,40,40,40	0
56	MG	1A	3056	1/1	0.97	0.13	62,62,62,62	0
56	MG	2A	3202	1/1	0.97	0.12	46,46,46,46	0
56	MG	1A	3216	1/1	0.97	0.09	37,37,37,37	0
56	MG	1A	3841	1/1	0.97	0.05	36,36,36,36	0
56	MG	1A	3239	1/1	0.97	0.19	36,36,36,36	0
56	MG	1A	3613	1/1	0.97	0.08	35,35,35,35	0
56	MG	2a	3206	1/1	0.97	0.07	63,63,63,63	0
56	MG	2A	3208	1/1	0.97	0.10	40,40,40,40	0
56	MG	2A	3209	1/1	0.97	0.10	45,45,45,45	0
56	MG	1a	1661	1/1	0.97	0.05	15,15,15,15	0
56	MG	1A	4012	1/1	0.97	0.17	36,36,36,36	0
56	MG	2A	3212	1/1	0.97	0.14	23,23,23,23	0
56	MG	1A	3557	1/1	0.97	0.07	31,31,31,31	0
56	MG	1A	3264	1/1	0.97	0.07	38,38,38,38	0
56	MG	2A	3215	1/1	0.97	0.06	39,39,39,39	0
56	MG	1A	3559	1/1	0.97	0.08	51,51,51,51	0
56	MG	1A	3517	1/1	0.97	0.20	43,43,43,43	0
56	MG	1A	3561	1/1	0.97	0.09	44,44,44,44	0
56	MG	1A	3562	1/1	0.97	0.07	55,55,55,55	0
56	MG	2a	3219	1/1	0.97	0.11	52,52,52,52	0
56	MG	1X	102	1/1	0.97	0.09	53,53,53,53	0
56	MG	1A	3447	1/1	0.97	0.05	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	3222	1/1	0.97	0.17	36,36,36,36	0
56	MG	1l	201	1/1	0.97	0.11	58,58,58,58	0
56	MG	2A	3094	1/1	0.97	0.12	46,46,46,46	0
56	MG	1A	3699	1/1	0.97	0.07	47,47,47,47	0
56	MG	1A	3386	1/1	0.97	0.11	63,63,63,63	0
56	MG	1a	1777	1/1	0.97	0.16	28,28,28,28	0
56	MG	1a	1674	1/1	0.97	0.16	54,54,54,54	0
56	MG	1A	3090	1/1	0.97	0.06	53,53,53,53	0
56	MG	1A	3627	1/1	0.97	0.06	20,20,20,20	0
56	MG	2A	3518	1/1	0.97	0.06	42,42,42,42	0
56	MG	1A	3219	1/1	0.97	0.09	59,59,59,59	0
56	MG	1A	3485	1/1	0.97	0.07	55,55,55,55	0
56	MG	1B	206	1/1	0.97	0.13	46,46,46,46	0
56	MG	1A	3571	1/1	0.97	0.07	25,25,25,25	0
56	MG	1A	3859	1/1	0.97	0.05	24,24,24,24	0
56	MG	1A	3007	1/1	0.97	0.10	56,56,56,56	0
56	MG	1a	1787	1/1	0.97	0.10	58,58,58,58	0
56	MG	1A	3160	1/1	0.97	0.13	40,40,40,40	0
56	MG	1A	3421	1/1	0.97	0.04	47,47,47,47	0
56	MG	1B	212	1/1	0.97	0.16	54,54,54,54	0
56	MG	1x	107	1/1	0.97	0.11	61,61,61,61	0
56	MG	1A	3273	1/1	0.97	0.10	36,36,36,36	0
56	MG	1A	3490	1/1	0.97	0.07	66,66,66,66	0
56	MG	2A	3384	1/1	0.97	0.06	45,45,45,45	0
56	MG	1A	3122	1/1	0.97	0.10	58,58,58,58	0
56	MG	1A	3529	1/1	0.97	0.06	48,48,48,48	0
56	MG	1A	3951	1/1	0.97	0.10	70,70,70,70	0
56	MG	1A	3011	1/1	0.97	0.06	67,67,67,67	0
56	MG	2A	3537	1/1	0.97	0.09	37,37,37,37	0
56	MG	1A	3643	1/1	0.97	0.06	19,19,19,19	0
56	MG	1A	3719	1/1	0.97	0.10	38,38,38,38	0
56	MG	1A	3224	1/1	0.97	0.16	35,35,35,35	0
56	MG	2A	3121	1/1	0.97	0.14	32,32,32,32	0
56	MG	19	101	1/1	0.97	0.08	65,65,65,65	0
56	MG	2a	3093	1/1	0.97	0.08	50,50,50,50	0
56	MG	2A	3394	1/1	0.97	0.25	54,54,54,54	0
56	MG	2A	3545	1/1	0.97	0.09	39,39,39,39	0
56	MG	1A	3494	1/1	0.97	0.18	51,51,51,51	0
56	MG	1A	3957	1/1	0.97	0.12	42,42,42,42	0
56	MG	2a	3098	1/1	0.97	0.24	50,50,50,50	0
56	MG	1a	1700	1/1	0.97	0.07	36,36,36,36	0
56	MG	2d	502	1/1	0.97	0.29	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	3723	1/1	0.97	0.09	8,8,8,8	0
56	MG	1A	3533	1/1	0.97	0.10	51,51,51,51	0
56	MG	2a	3102	1/1	0.97	0.25	47,47,47,47	0
56	MG	2A	3259	1/1	0.97	0.06	72,72,72,72	0
56	MG	2l	202	1/1	0.97	0.05	40,40,40,40	0
56	MG	1A	3225	1/1	0.97	0.14	34,34,34,34	0
56	MG	1A	3180	1/1	0.97	0.08	33,33,33,33	0
56	MG	1A	3650	1/1	0.97	0.08	19,19,19,19	0
56	MG	1A	3963	1/1	0.97	0.11	14,14,14,14	0
56	MG	1A	3052	1/1	0.97	0.16	59,59,59,59	0
56	MG	2a	3109	1/1	0.97	0.17	47,47,47,47	0
56	MG	2A	3406	1/1	0.97	0.05	41,41,41,41	0
56	MG	2A	3265	1/1	0.97	0.12	37,37,37,37	0
56	MG	2A	3563	1/1	0.97	0.06	47,47,47,47	0
56	MG	1A	3730	1/1	0.97	0.07	30,30,30,30	0
57	ZN	1n	501	1/1	0.97	0.05	84,84,84,84	0
56	MG	1A	3183	1/1	0.97	0.18	44,44,44,44	0
57	ZN	29	501	1/1	0.97	0.04	62,62,62,62	0
56	MG	1A	3587	1/1	0.97	0.11	40,40,40,40	0
56	MG	1A	3629	1/1	0.98	0.08	27,27,27,27	0
56	MG	2a	3028	1/1	0.98	0.06	76,76,76,76	0
56	MG	1A	3391	1/1	0.98	0.05	35,35,35,35	0
56	MG	1A	3631	1/1	0.98	0.04	19,19,19,19	0
56	MG	1A	3288	1/1	0.98	0.14	40,40,40,40	0
56	MG	1A	3989	1/1	0.98	0.06	25,25,25,25	0
56	MG	1A	3199	1/1	0.98	0.06	37,37,37,37	0
56	MG	1A	3790	1/1	0.98	0.08	48,48,48,48	0
56	MG	1A	3666	1/1	0.98	0.11	37,37,37,37	0
56	MG	2A	3480	1/1	0.98	0.15	46,46,46,46	0
56	MG	1A	3061	1/1	0.98	0.16	44,44,44,44	0
56	MG	1D	308	1/1	0.98	0.10	35,35,35,35	0
56	MG	1A	3604	1/1	0.98	0.04	27,27,27,27	0
56	MG	1A	3250	1/1	0.98	0.17	34,34,34,34	0
56	MG	1A	3292	1/1	0.98	0.14	17,17,17,17	0
56	MG	1a	1691	1/1	0.98	0.06	27,27,27,27	0
56	MG	1A	3639	1/1	0.98	0.05	25,25,25,25	0
56	MG	1a	1693	1/1	0.98	0.11	52,52,52,52	0
56	MG	1a	1824	1/1	0.98	0.10	52,52,52,52	0
56	MG	1X	101	1/1	0.98	0.08	45,45,45,45	0
56	MG	1A	3752	1/1	0.98	0.08	50,50,50,50	0
56	MG	1A	3799	1/1	0.98	0.10	42,42,42,42	0
56	MG	1E	301	1/1	0.98	0.04	15,15,15,15	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	2A	3060	1/1	0.98	0.14	43,43,43,43	0
56	MG	1A	3672	1/1	0.98	0.04	13,13,13,13	0
56	MG	2a	3223	1/1	0.98	0.10	49,49,49,49	0
56	MG	1A	3267	1/1	0.98	0.26	27,27,27,27	0
56	MG	2D	303	1/1	0.98	0.07	49,49,49,49	0
56	MG	2A	3200	1/1	0.98	0.27	48,48,48,48	0
56	MG	1A	3948	1/1	0.98	0.04	28,28,28,28	0
56	MG	2A	3344	1/1	0.98	0.10	39,39,39,39	0
56	MG	1A	3174	1/1	0.98	0.22	37,37,37,37	0
56	MG	1A	3756	1/1	0.98	0.10	51,51,51,51	0
56	MG	1A	3382	1/1	0.98	0.03	40,40,40,40	0
56	MG	1F	303	1/1	0.98	0.04	32,32,32,32	0
56	MG	2A	3206	1/1	0.98	0.08	41,41,41,41	0
56	MG	1a	1837	1/1	0.98	0.08	37,37,37,37	0
56	MG	1A	3718	1/1	0.98	0.04	41,41,41,41	0
56	MG	1A	3563	1/1	0.98	0.08	46,46,46,46	0
56	MG	1A	3901	1/1	0.98	0.06	44,44,44,44	0
56	MG	2A	3431	1/1	0.98	0.05	68,68,68,68	0
56	MG	1A	3720	1/1	0.98	0.07	45,45,45,45	0
56	MG	2A	3359	1/1	0.98	0.07	50,50,50,50	0
56	MG	1A	3469	1/1	0.98	0.07	49,49,49,49	0
56	MG	2O	202	1/1	0.98	0.15	42,42,42,42	0
56	MG	1A	3588	1/1	0.98	0.07	33,33,33,33	0
56	MG	1A	3679	1/1	0.98	0.04	25,25,25,25	0
56	MG	1A	3764	1/1	0.98	0.10	41,41,41,41	0
56	MG	1A	3907	1/1	0.98	0.06	15,15,15,15	0
56	MG	1A	3765	1/1	0.98	0.04	44,44,44,44	0
56	MG	1A	3295	1/1	0.98	0.06	50,50,50,50	0
56	MG	1a	1716	1/1	0.98	0.07	53,53,53,53	0
56	MG	1A	3725	1/1	0.98	0.06	42,42,42,42	0
56	MG	1A	3768	1/1	0.98	0.05	10,10,10,10	0
56	MG	1A	3269	1/1	0.98	0.05	36,36,36,36	0
56	MG	1A	3568	1/1	0.98	0.17	42,42,42,42	0
56	MG	1A	3864	1/1	0.98	0.06	55,55,55,55	0
56	MG	19	104	1/1	0.98	0.04	31,31,31,31	0
56	MG	1A	3195	1/1	0.98	0.11	33,33,33,33	0
56	MG	2a	3085	1/1	0.98	0.04	40,40,40,40	0
56	MG	1A	3272	1/1	0.98	0.07	29,29,29,29	0
56	MG	1A	3621	1/1	0.98	0.06	18,18,18,18	0
56	MG	1A	3731	1/1	0.98	0.07	23,23,23,23	0
56	MG	1A	3688	1/1	0.98	0.08	41,41,41,41	0
56	MG	2A	3231	1/1	0.98	0.17	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	2A	3454	1/1	0.98	0.05	56,56,56,56	0
56	MG	1A	3510	1/1	0.98	0.09	55,55,55,55	0
56	MG	1A	3690	1/1	0.98	0.09	48,48,48,48	0
56	MG	1A	3356	1/1	0.98	0.13	42,42,42,42	0
56	MG	1A	3181	1/1	0.98	0.14	24,24,24,24	0
56	MG	1a	1669	1/1	0.98	0.16	46,46,46,46	0
56	MG	1A	3874	1/1	0.98	0.14	56,56,56,56	0
56	MG	1A	3551	1/1	0.98	0.12	33,33,33,33	0
56	MG	2A	3541	1/1	0.98	0.20	42,42,42,42	0
56	MG	1B	214	1/1	0.98	0.04	68,68,68,68	0
56	MG	2A	3463	1/1	0.98	0.09	57,57,57,57	0
56	MG	1A	3926	1/1	0.98	0.10	38,38,38,38	0
56	MG	1A	3927	1/1	0.98	0.11	50,50,50,50	0
56	MG	1a	1872	1/1	0.98	0.12	53,53,53,53	0
56	MG	1A	3218	1/1	0.98	0.14	21,21,21,21	0
56	MG	2A	3549	1/1	0.98	0.10	43,43,43,43	0
56	MG	1A	3188	1/1	0.98	0.06	43,43,43,43	0
57	ZN	1Y	201	1/1	0.98	0.04	61,61,61,61	0
56	MG	1a	1740	1/1	0.98	0.09	47,47,47,47	0
57	ZN	2Y	501	1/1	0.98	0.05	85,85,85,85	0
56	MG	2A	3552	1/1	0.98	0.10	22,22,22,22	0
56	MG	2A	3553	1/1	0.98	0.04	44,44,44,44	0
56	MG	1A	3660	1/1	0.98	0.05	11,11,11,11	0
58	SF4	1d	501	8/8	0.98	0.05	66,73,85,85	0
58	SF4	2d	501	8/8	0.98	0.05	56,76,90,91	0
56	MG	2A	3351	1/1	0.99	0.07	37,37,37,37	0
56	MG	2a	3195	1/1	0.99	0.03	43,43,43,43	0
56	MG	1A	3635	1/1	0.99	0.04	46,46,46,46	0
56	MG	1A	4014	1/1	0.99	0.07	43,43,43,43	0
56	MG	1A	3716	1/1	0.99	0.03	43,43,43,43	0
56	MG	1A	3846	1/1	0.99	0.03	78,78,78,78	0
56	MG	1A	3692	1/1	0.99	0.06	33,33,33,33	0
56	MG	1A	3693	1/1	0.99	0.07	21,21,21,21	0
56	MG	1D	307	1/1	0.99	0.06	15,15,15,15	0
56	MG	2A	3492	1/1	0.99	0.05	44,44,44,44	0
56	MG	1a	1789	1/1	0.99	0.06	66,66,66,66	0
56	MG	1A	3694	1/1	0.99	0.03	16,16,16,16	0
56	MG	1A	3208	1/1	0.99	0.21	34,34,34,34	0
56	MG	1A	3169	1/1	0.99	0.07	21,21,21,21	0
56	MG	1A	3934	1/1	0.99	0.03	33,33,33,33	0
56	MG	1a	1831	1/1	0.99	0.06	54,54,54,54	0
56	MG	1A	3622	1/1	0.99	0.07	22,22,22,22	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
56	MG	1A	3309	1/1	0.99	0.02	20,20,20,20	0
56	MG	2A	3163	1/1	0.99	0.15	33,33,33,33	0
56	MG	1A	3608	1/1	0.99	0.07	25,25,25,25	0
56	MG	2A	3548	1/1	0.99	0.13	40,40,40,40	0
56	MG	1A	3775	1/1	0.99	0.06	14,14,14,14	0
56	MG	1Q	3004	1/1	0.99	0.04	39,39,39,39	0
56	MG	1A	3967	1/1	0.99	0.04	19,19,19,19	0
56	MG	1A	3163	1/1	0.99	0.06	32,32,32,32	0
56	MG	1A	3137	1/1	0.99	0.06	51,51,51,51	0
56	MG	1A	3702	1/1	0.99	0.03	44,44,44,44	0
56	MG	1A	3212	1/1	0.99	0.04	39,39,39,39	0
56	MG	1A	3453	1/1	0.99	0.06	18,18,18,18	0
56	MG	1A	3196	1/1	0.99	0.11	25,25,25,25	0
56	MG	1A	3646	1/1	0.99	0.06	10,10,10,10	0
56	MG	1A	3286	1/1	0.99	0.08	36,36,36,36	0
56	MG	1A	3566	1/1	0.99	0.09	23,23,23,23	0
56	MG	1B	216	1/1	0.99	0.05	54,54,54,54	0
56	MG	1B	217	1/1	0.99	0.09	50,50,50,50	0
56	MG	1A	3686	1/1	0.99	0.04	46,46,46,46	0
56	MG	1A	3156	1/1	0.99	0.11	47,47,47,47	0
56	MG	1A	3271	1/1	0.99	0.06	17,17,17,17	0
56	MG	1A	3712	1/1	0.99	0.07	43,43,43,43	0
56	MG	1A	4010	1/1	0.99	0.18	41,41,41,41	0
57	ZN	14	501	1/1	0.99	0.04	106,106,106,106	0
57	ZN	15	103	1/1	0.99	0.02	43,43,43,43	0
56	MG	1a	1816	1/1	0.99	0.05	40,40,40,40	0
56	MG	2A	3345	1/1	0.99	0.06	35,35,35,35	0
56	MG	1A	3651	1/1	0.99	0.06	21,21,21,21	0
57	ZN	25	501	1/1	0.99	0.03	54,54,54,54	0
57	ZN	26	102	1/1	0.99	0.04	56,56,56,56	0
56	MG	1a	1856	1/1	0.99	0.07	43,43,43,43	0
56	MG	2A	3348	1/1	0.99	0.06	52,52,52,52	0
56	MG	1A	3618	1/1	0.99	0.09	18,18,18,18	0
56	MG	2A	3350	1/1	0.99	0.04	34,34,34,34	0
56	MG	2A	3471	1/1	1.00	0.04	34,34,34,34	0
56	MG	1A	3796	1/1	1.00	0.02	16,16,16,16	0
56	MG	1A	3620	1/1	1.00	0.04	38,38,38,38	0
57	ZN	16	102	1/1	1.00	0.02	29,29,29,29	0
57	ZN	19	103	1/1	1.00	0.04	45,45,45,45	0
56	MG	1A	3897	1/1	1.00	0.04	29,29,29,29	0
56	MG	1A	3681	1/1	1.00	0.02	14,14,14,14	0

5.5 Other polymers [i](#)

There are no such residues in this entry.