



# wwPDB X-ray Structure Validation Summary Report ⓘ

Feb 1, 2016 – 11:26 PM GMT

PDB ID : 4Z8C  
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 wwPDB X-ray Structure Validation Summary Report for a publicly released PDB entry.  
We welcome your comments at [validation@mail.wwpdb.org](mailto:validation@mail.wwpdb.org)  
A user guide is available at  
<http://wwpdb.org/validation/2016/XrayValidationReportHelp>  
with specific help available everywhere you see the ⓘ symbol.

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

MolProbity : 4.02b-467  
Mogul : 1.7 (RC4), CSD as536be (2015)  
Xtriage (Phenix) : 1.9-1692  
EDS : rb-20026688  
Percentile statistics : 20151230.v01 (using entries in the PDB archive December 30th 2015)  
Refmac : 5.8.0135  
CCP4 : 6.5.0  
Ideal geometry (proteins) : Engh & Huber (2001)  
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)  
Validation Pipeline (wwPDB-VP) : trunk26865



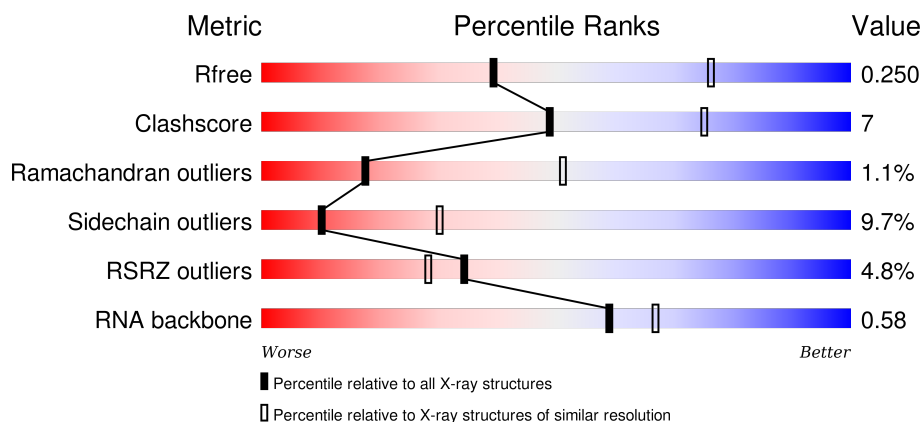
# 1 Overall quality at a glance

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))
$R_{free}$	91344	1451 (2.90-2.90)
Clashscore	102246	1668 (2.90-2.90)
Ramachandran outliers	100387	1630 (2.90-2.90)
Sidechain outliers	100360	1632 (2.90-2.90)
RSRZ outliers	91569	1456 (2.90-2.90)
RNA backbone	2183	1093 (3.30-2.50)






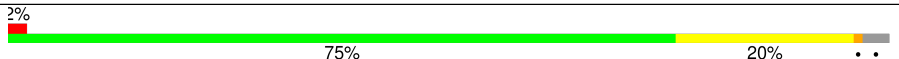

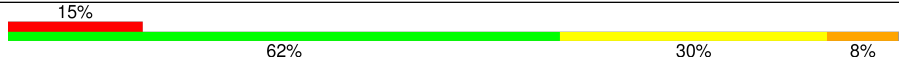

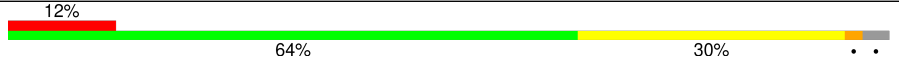







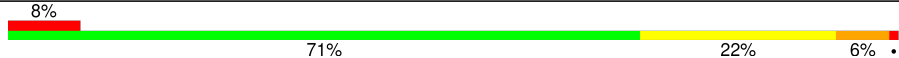

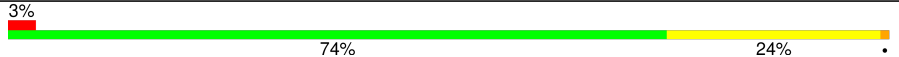





The table below summarises the geometric issues observed across the polymeric chains and their fit to the electron density. The red, orange, yellow and green segments on the lower bar indicate the fraction of residues that contain outliers for  $\geq 3$ , 2, 1 and 0 types of geometric quality criteria.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

Mol	Chain	Length	Quality of chain
1	1A	2915	
1	2A	2915	
2	1B	121	
2	2B	121	

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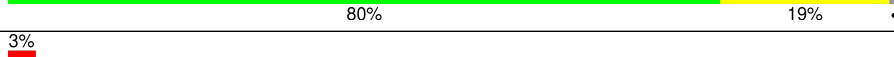




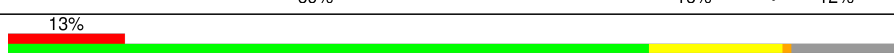


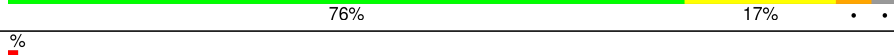


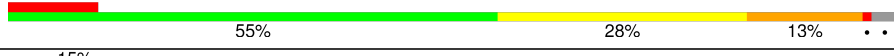

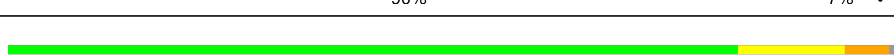




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Mol	Chain	Length	Quality of chain
3	1D	276	
3	2D	276	
4	1E	206	
4	2E	206	
5	1F	210	
5	2F	210	
6	1G	182	
6	2G	182	
7	1H	180	
7	2H	180	
8	1I	148	
8	2I	148	
9	1N	140	
9	2N	140	
10	1O	122	
10	2O	122	
11	1P	150	
11	2P	150	
12	1Q	141	
12	2Q	141	
13	1R	118	
13	2R	118	
14	1S	112	
14	2S	112	
15	1T	146	

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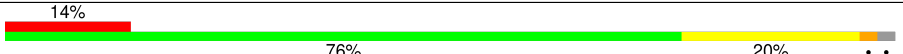
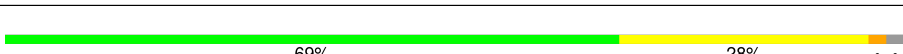
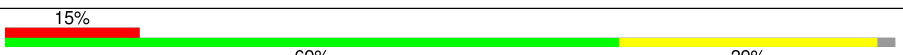
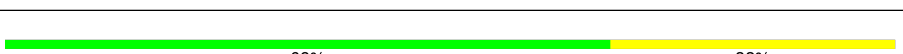
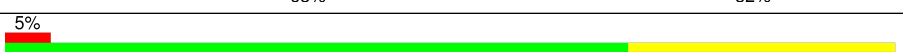

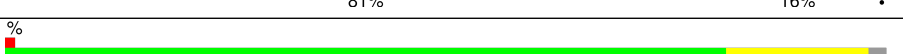
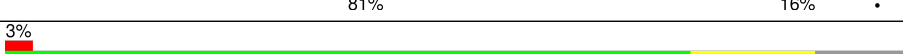
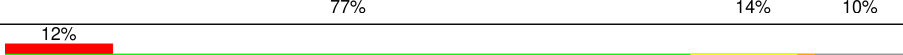
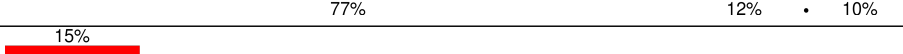
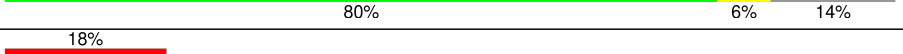





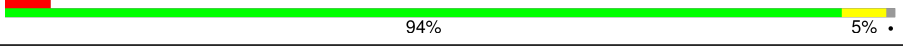

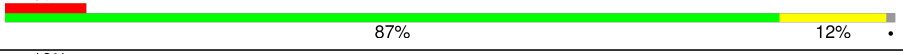
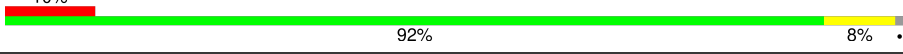


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Mol	Chain	Length	Quality of chain
15	2T	146	
16	1U	118	
16	2U	118	
17	1V	101	
17	2V	101	
18	1W	113	
18	2W	113	
19	1X	96	
19	2X	96	
20	1Y	110	
20	2Y	110	
21	1Z	206	
21	2Z	206	
22	10	85	
22	20	85	
23	11	98	
23	21	98	
24	12	72	
24	22	72	
25	13	60	
25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

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Mol	Chain	Length	Quality of chain
28	16	54	
28	26	54	
29	17	49	
29	27	49	
30	18	65	
30	28	65	
31	19	37	
31	29	37	
32	1a	1521	
32	2a	1521	
33	1b	256	
33	2b	256	
34	1c	239	
34	2c	239	
35	1d	209	
35	2d	209	
36	1e	162	
36	2e	162	
37	1f	101	
37	2f	101	
38	1g	156	
38	2g	156	
39	1h	138	
39	2h	138	
40	1i	128	

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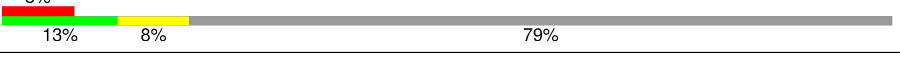

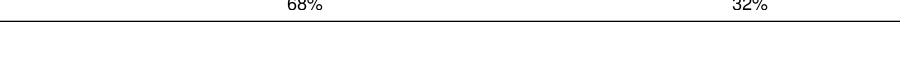
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Mol	Chain	Length	Quality of chain
40	2i	128	
41	1j	105	
41	2j	105	
42	1k	129	
42	2k	129	
43	1l	132	
43	2l	132	
44	1m	126	
44	2m	126	
45	1n	61	
45	2n	61	
46	1o	89	
46	2o	89	
47	1p	88	
47	2p	88	
48	1q	105	
48	2q	105	
49	1r	88	
49	2r	88	
50	1s	93	
50	2s	93	
51	1t	106	
51	2t	106	
52	1u	27	
52	2u	27	

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Mol	Chain	Length	Quality of chain
53	1v	24	
53	2v	24	
54	1x	77	
54	2x	77	
55	1z	19	
55	2z	19	

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	15	101	-	-	-	X
56	MG	1A	3001	-	-	-	X
56	MG	1A	3074	-	-	-	X
56	MG	1A	3092	-	-	-	X
56	MG	1A	3093	-	-	-	X
56	MG	1A	3110	-	-	-	X
56	MG	1A	3129	-	-	-	X
56	MG	1A	3131	-	-	-	X
56	MG	1A	3132	-	-	-	X
56	MG	1A	3162	-	-	-	X
56	MG	1A	3170	-	-	-	X
56	MG	1A	3172	-	-	-	X
56	MG	1A	3177	-	-	-	X
56	MG	1A	3178	-	-	-	X
56	MG	1A	3179	-	-	-	X
56	MG	1A	3182	-	-	-	X
56	MG	1A	3186	-	-	-	X
56	MG	1A	3187	-	-	-	X
56	MG	1A	3203	-	-	-	X
56	MG	1A	3208	-	-	-	X
56	MG	1A	3215	-	-	-	X
56	MG	1A	3225	-	-	-	X
56	MG	1A	3241	-	-	-	X
56	MG	1A	3253	-	-	-	X
56	MG	1A	3267	-	-	-	X
56	MG	1A	3287	-	-	-	X
56	MG	1A	3289	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
56	MG	1A	3312	-	-	-	X
56	MG	1A	3315	-	-	-	X
56	MG	1A	3322	-	-	-	X
56	MG	1A	3329	-	-	-	X
56	MG	1A	3374	-	-	-	X
56	MG	1A	3394	-	-	-	X
56	MG	1A	3411	-	-	-	X
56	MG	1A	3437	-	-	-	X
56	MG	1A	3442	-	-	-	X
56	MG	1A	3444	-	-	-	X
56	MG	1A	3449	-	-	-	X
56	MG	1A	3456	-	-	-	X
56	MG	1A	3459	-	-	-	X
56	MG	1A	3473	-	-	-	X
56	MG	1A	3481	-	-	-	X
56	MG	1A	3518	-	-	-	X
56	MG	1A	3705	-	-	-	X
56	MG	1A	3711	-	-	-	X
56	MG	1A	3797	-	-	-	X
56	MG	1A	3821	-	-	-	X
56	MG	1A	3887	-	-	-	X
56	MG	1A	3888	-	-	-	X
56	MG	1A	3911	-	-	-	X
56	MG	1A	3930	-	-	-	X
56	MG	1A	3931	-	-	-	X
56	MG	1A	3962	-	-	-	X
56	MG	1A	3964	-	-	-	X
56	MG	1A	3968	-	-	-	X
56	MG	1A	3973	-	-	-	X
56	MG	1A	3975	-	-	-	X
56	MG	1A	3980	-	-	-	X
56	MG	1A	3986	-	-	-	X
56	MG	1A	3994	-	-	-	X
56	MG	1A	3995	-	-	-	X
56	MG	1A	4002	-	-	-	X
56	MG	1A	4003	-	-	-	X
56	MG	1A	4007	-	-	-	X
56	MG	1A	4008	-	-	-	X
56	MG	1A	4010	-	-	-	X
56	MG	1A	4011	-	-	-	X
56	MG	1A	4012	-	-	-	X
56	MG	1A	4017	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
56	MG	1A	4018	-	-	-	X
56	MG	1A	4019	-	-	-	X
56	MG	1B	205	-	-	-	X
56	MG	1B	211	-	-	-	X
56	MG	1D	303	-	-	-	X
56	MG	1D	306	-	-	-	X
56	MG	1D	309	-	-	-	X
56	MG	1D	311	-	-	-	X
56	MG	1D	312	-	-	-	X
56	MG	1E	303	-	-	-	X
56	MG	1F	304	-	-	-	X
56	MG	1F	306	-	-	-	X
56	MG	1H	8002	-	-	-	X
56	MG	1P	205	-	-	-	X
56	MG	1P	206	-	-	-	X
56	MG	1R	203	-	-	-	X
56	MG	1U	202	-	-	-	X
56	MG	1U	203	-	-	-	X
56	MG	1U	204	-	-	-	X
56	MG	1U	205	-	-	-	X
56	MG	1W	3005	-	-	-	X
56	MG	1X	102	-	-	-	X
56	MG	1a	1659	-	-	-	X
56	MG	1a	1668	-	-	-	X
56	MG	1a	1739	-	-	-	X
56	MG	1a	1749	-	-	-	X
56	MG	1a	1765	-	-	-	X
56	MG	1a	1814	-	-	-	X
56	MG	1a	1858	-	-	-	X
56	MG	2A	3016	-	-	-	X
56	MG	2A	3017	-	-	-	X
56	MG	2A	3019	-	-	-	X
56	MG	2A	3024	-	-	-	X
56	MG	2A	3027	-	-	-	X
56	MG	2A	3029	-	-	-	X
56	MG	2A	3042	-	-	-	X
56	MG	2A	3063	-	-	-	X
56	MG	2A	3107	-	-	-	X
56	MG	2A	3123	-	-	-	X
56	MG	2A	3139	-	-	-	X
56	MG	2A	3155	-	-	-	X
56	MG	2A	3188	-	-	-	X

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Mol	Type	Chain	Res	Chirality	Geometry	Clashes	Electron density
56	MG	2A	3191	-	-	-	X
56	MG	2A	3199	-	-	-	X
56	MG	2A	3209	-	-	-	X
56	MG	2A	3214	-	-	-	X
56	MG	2A	3222	-	-	-	X
56	MG	2A	3232	-	-	-	X
56	MG	2A	3233	-	-	-	X
56	MG	2A	3303	-	-	-	X
56	MG	2A	3304	-	-	-	X
56	MG	2A	3318	-	-	-	X
56	MG	2A	3349	-	-	-	X
56	MG	2A	3381	-	-	-	X
56	MG	2A	3416	-	-	-	X
56	MG	2A	3430	-	-	-	X
56	MG	2A	3468	-	-	-	X
56	MG	2A	3470	-	-	-	X
56	MG	2A	3540	-	-	-	X
56	MG	2A	3550	-	-	-	X
56	MG	2A	3554	-	-	-	X
56	MG	2A	3555	-	-	-	X
56	MG	2A	3559	-	-	-	X
56	MG	2A	3561	-	-	-	X
56	MG	2A	3562	-	-	-	X
56	MG	2A	3563	-	-	-	X
56	MG	2D	303	-	-	-	X
56	MG	2F	303	-	-	-	X
56	MG	2P	201	-	-	-	X
56	MG	2U	201	-	-	-	X
56	MG	2a	3011	-	-	-	X
56	MG	2a	3022	-	-	-	X
56	MG	2a	3026	-	-	-	X
56	MG	2a	3105	-	-	-	X
56	MG	2a	3194	-	-	-	X
56	MG	2a	3228	-	-	-	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 (three-letter code: MG) (formula: Mg).

Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	2E	4	Total	Mg	0	0
			4	4		
56	17	1	Total	Mg	0	0
			1	1		
56	1z	1	Total	Mg	0	0
			1	1		
56	2d	1	Total	Mg	0	0
			1	1		
56	1T	5	Total	Mg	0	0
			5	5		
56	1N	6	Total	Mg	0	0
			6	6		
56	20	1	Total	Mg	0	0
			1	1		
56	18	3	Total	Mg	0	0
			3	3		
56	1o	2	Total	Mg	0	0
			2	2		

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	13	2	Total 2	Mg 2	0	0
56	1f	1	Total 1	Mg 1	0	0
56	2h	1	Total 1	Mg 1	0	0
56	1P	6	Total 6	Mg 6	0	0
56	2B	11	Total 11	Mg 11	0	0
56	1q	4	Total 4	Mg 4	0	0
56	2a	263	Total 263	Mg 263	0	0
56	1k	1	Total 1	Mg 1	0	0
56	1E	5	Total 5	Mg 5	0	0
56	2z	1	Total 1	Mg 1	0	0
56	1b	2	Total 2	Mg 2	0	0
56	2l	4	Total 4	Mg 4	0	0
56	2F	6	Total 6	Mg 6	0	0
56	16	2	Total 2	Mg 2	0	0
56	28	1	Total 1	Mg 1	0	0
56	1W	5	Total 5	Mg 5	0	0
56	1A	1021	Total 1021	Mg 1021	0	0
56	1t	1	Total 1	Mg 1	0	0
56	1n	1	Total 1	Mg 1	0	0
56	2P	1	Total 1	Mg 1	0	0
56	1X	2	Total 2	Mg 2	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	12	1	Total 1	Mg 1	0	0
56	25	1	Total 1	Mg 1	0	0
56	1D	14	Total 14	Mg 14	0	0
56	2N	1	Total 1	Mg 1	0	0
56	1e	3	Total 3	Mg 3	0	0
56	2G	1	Total 1	Mg 1	0	0
56	2f	1	Total 1	Mg 1	0	0
56	1V	1	Total 1	Mg 1	0	0
56	1a	276	Total 276	Mg 276	0	0
56	2Q	5	Total 5	Mg 5	0	0
56	15	4	Total 4	Mg 4	0	0
56	1x	8	Total 8	Mg 8	0	0
56	1R	4	Total 4	Mg 4	0	0
56	26	1	Total 1	Mg 1	0	0
56	2U	1	Total 1	Mg 1	0	0
56	1G	3	Total 3	Mg 3	0	0
56	2O	3	Total 3	Mg 3	0	0
56	11	1	Total 1	Mg 1	0	0
56	1d	2	Total 2	Mg 2	0	0
56	1H	2	Total 2	Mg 2	0	0
56	2Y	1	Total 1	Mg 1	0	0

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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
56	2x	4	Total 4	Mg 4	0	0
56	1Z	2	Total 2	Mg 2	0	0
56	2D	6	Total 6	Mg 6	0	0
56	2q	2	Total 2	Mg 2	0	0
56	2k	1	Total 1	Mg 1	0	0
56	1U	5	Total 5	Mg 5	0	0
56	1O	2	Total 2	Mg 2	0	0
56	1r	2	Total 2	Mg 2	0	0
56	19	3	Total 3	Mg 3	0	0
56	1l	1	Total 1	Mg 1	0	0
56	1F	8	Total 8	Mg 8	0	0
56	10	7	Total 7	Mg 7	0	0
56	2t	1	Total 1	Mg 1	0	0
56	1Q	4	Total 4	Mg 4	0	0
56	2A	566	Total 566	Mg 566	0	0
56	2Z	1	Total 1	Mg 1	0	0
56	1B	24	Total 24	Mg 24	0	0

- Molecule 57 is ZINC ION (three-letter code: ZN) (formula: Zn).

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

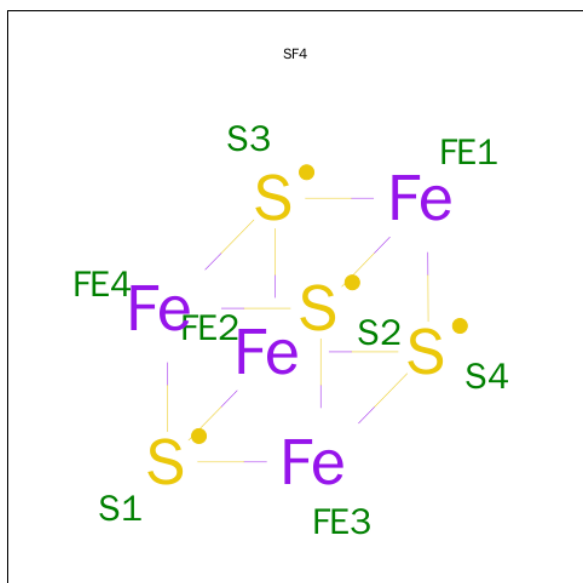
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Mol	Chain	Residues	Atoms		ZeroOcc	AltConf
57	1n	1	Total 1	Zn 1	0	0
57	15	1	Total 1	Zn 1	0	0
57	29	1	Total 1	Zn 1	0	0
57	19	1	Total 1	Zn 1	0	0
57	26	1	Total 1	Zn 1	0	0
57	25	1	Total 1	Zn 1	0	0
57	24	1	Total 1	Zn 1	0	0
57	2n	1	Total 1	Zn 1	0	0
57	2Y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

- Molecule 58 is IRON/SULFUR CLUSTER (three-letter code: SF4) (formula: Fe<sub>4</sub>S<sub>4</sub>).



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

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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
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59	2q	2	Total 2	O 2	0	0

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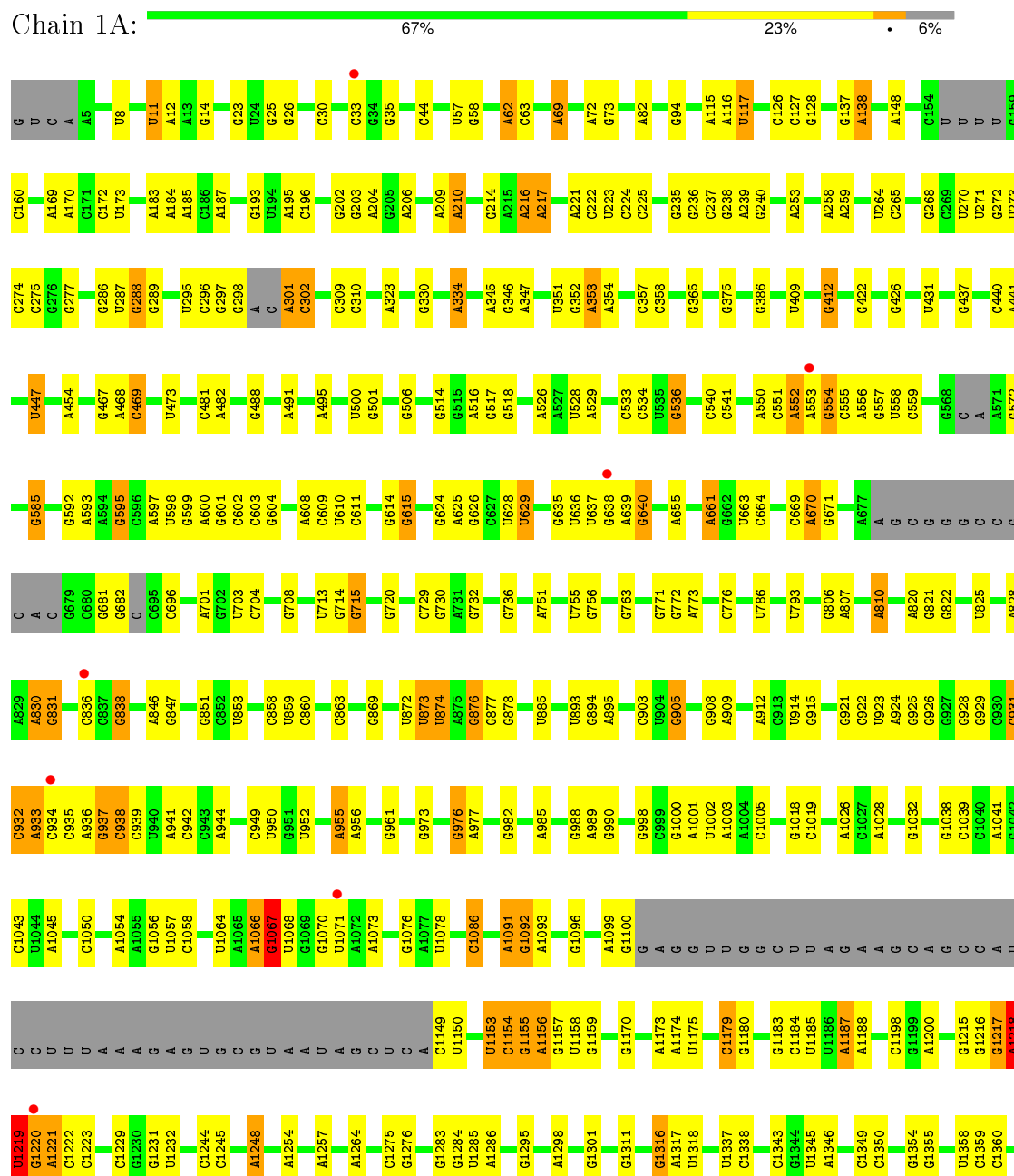
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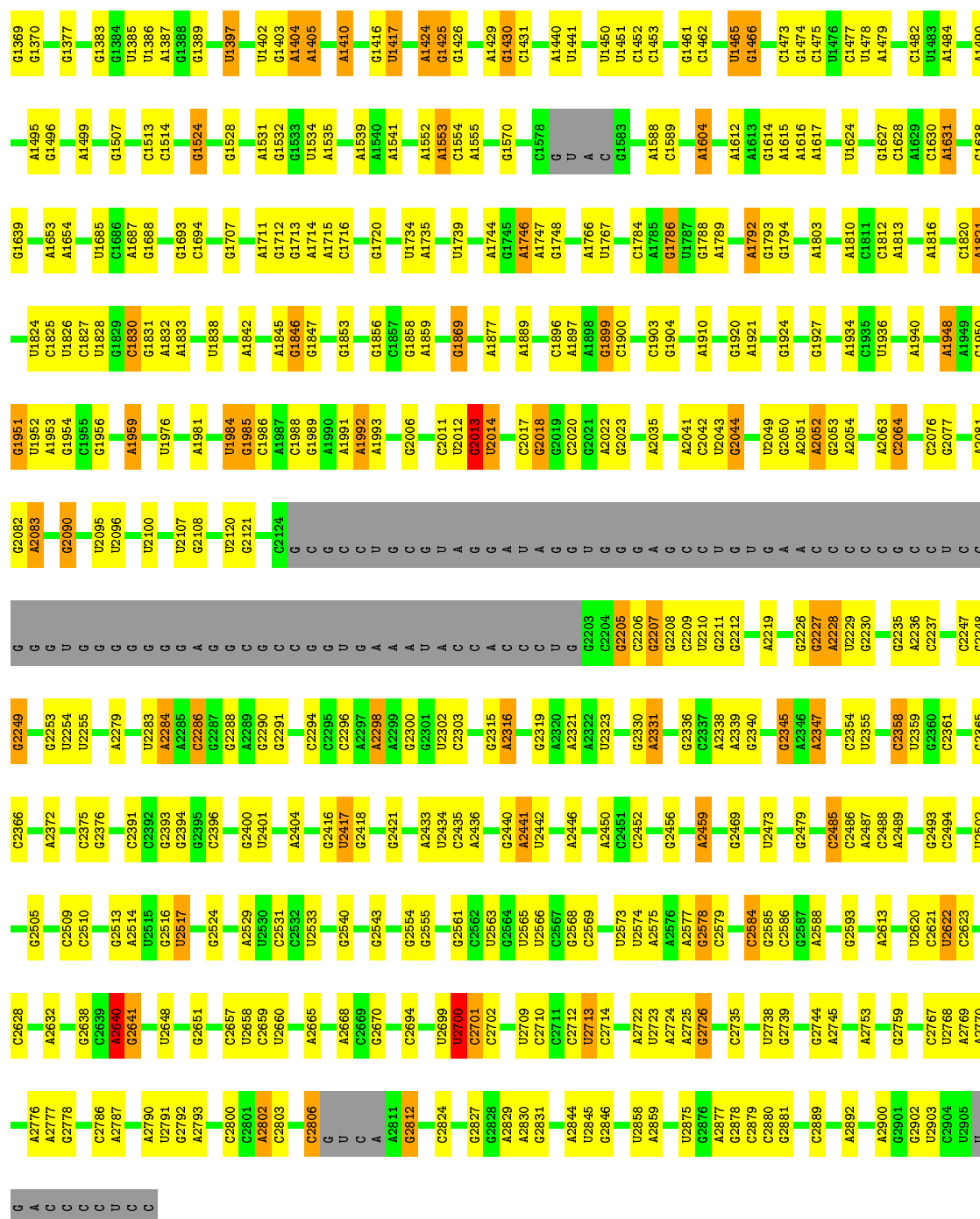
### 3 Residue-property plots

These plots are drawn for all protein, RNA and DNA chains in the entry. The first graphic for a chain summarises the proportions of errors displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and electron density. Residues are color-coded according to the number of geometric quality criteria for which they contain at least one outlier: green = 0, yellow = 1, orange = 2 and red = 3 or more. A red dot above a residue indicates a poor fit to the electron density ( $RSRZ > 2$ ). Stretches of 2 or more consecutive residues without any outlier are shown as a green connector. Residues present in the sample, but not in the model, are shown in grey.

- Molecule 1: 23S ribosomal RNA







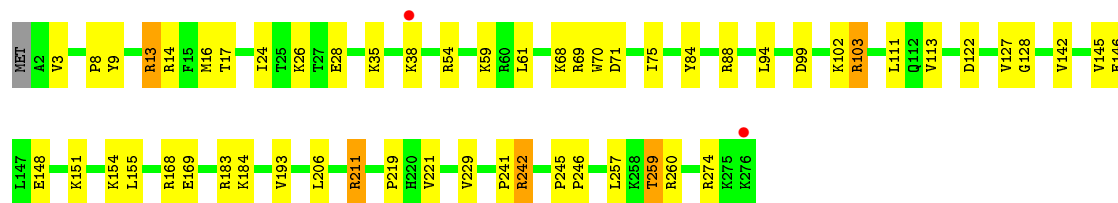


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G1712	G1846	G1461	G1585	G1461	G1355	U1068	A962	U873	C767	C858	A556	G437	A299	G193
G1713	G1847	C1462	U1586	C1462	G1356	G1069	A963	U874	U770	C859	U558	A438	A301	
C1716	G1854	U1465	C1589	U1465	U1358	U1071	U967	U879	A768	G660	C559	A448	C309	G201
G1720	C1873	G1466	A1590	G1466	C1359	A1072	C988	U885	G769	A661	A560	A454	C310	G202
U1734	A1877	U1361	C1592	U1361	U1360	A1073	U972	U886	U770	G662	C561		C311	G203
A1735	A1878	G1362	C1593	G1362	U1361	G1075	G973	C886	C776	U663		A468	C320	A204
U1739	G1880	C1473	A1604	C1473	U1361	G1076	G973	U889	C777	C664	C568	C469	C321	A206
U1744		C1474	G1605	C1474	U1362	A1077	G976	A895	C778	A668	C	C470	A322	
A1745		U1476	G1370	U1476	A1238	U1078	G976		G779	A670	A	G471	A323	A209
G1746		C1477		C1477	C1245	U1080	G984	U898	U786	G671	G573	C478	G324	A210
A1747		U1374		U1374	G1151	G1081	A985	U899	G806	G675		A479	C325	A217
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C1754		U1377		U1377	G1154	G1084	A989	U902	U808	G679	C587	A482	G332	C222
		A1381		A1381	G1155	G1085	G990	U903	G809	C680	U588		A333	U223
		G1382		G1382	U1156	G1086	G991	U904	A810	C681	U589	G493	A334	C224
		C1383		C1383	U1157	G1087	G992	U905	G811	G	U590	G494		C225
		U1397		U1397	U1158	G1088		U908	G816	C		A495	A343	
		G1403		G1403	G1161	G1089	G1000	U909		C	A593	A496	A344	G228
		A1404		A1404	G1162	A1090	A1001	A909	A820	C	A594	A497	A345	
		U1405		U1405	G1163	A1091	U1002	G910	G821	C	G595		G346	A232
		C1406		C1406	C1164	G1092	C1005	U920	G822	C	C596	G501	A347	
		C1407		C1407	G1167	A1093	G1018	G921	U825	C	U598	A504		G235
		A1413		A1413	G1170	A1094	G1019	A924	A828	C	A599	A505	G350	G236
		C1415		C1415	G1171	A1095	C1019	U925	A829	C	A600	G506	U351	C237
		A1423		A1423	G1172	C1097	G1036	C930	A834	C	G601	A507	C238	
		U1424		U1424	G1180	G1098	G1037	C931	A835	C	G602	A508	A353	C241
		G1425		G1425	G1181	A1099	G1038	C932	C836	C	G603	A516	A354	
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		U1429		U1429	G1183	G1100	C1027	U928	A838	C		G518	C360	A253
		G1430		G1430	G1184	G1101	G1028	G929	A839	C	A608	G524	G361	
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		U1441		U1441	U1186	G1103	G1037	C931	C836	C	U610	U528	G365	A259
		A1442		A1442	G1187	G1104	G1038	C932	C837	C	C611	A529	G375	A260
		C1443		C1443	G1188	U1105	G1039	C933	C838	C		G530	U885	U264
		U1444		U1444	G1189	G1106	C1040	A936	C841	C	A625	G531	U885	C265
		G1444		G1444	G1190	G1107	A1041	U936	C842	C	G626	G532	C266	C267
		U1450		U1450	G1191	G1108	G1042	C938	C843	C	U628	C533	G267	G268
		A1451		A1451	G1192	G1109	G1042	C938	C844	C	A630	G536	G398	C269
					G1193	U1110	G1047	C939	C845	C	A631	G540	G408	U270
					G1194	U1111	G1048	U940	C846	C		C541	U409	U271
					G1195	G1112	G1049	A941	C847	C		C542	U410	G272
					G1196	G1113	C1050	C946	C848	C		C543	G412	C274
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					G1199	G1116	U1057	C949	C851	C		C546	G431	U287
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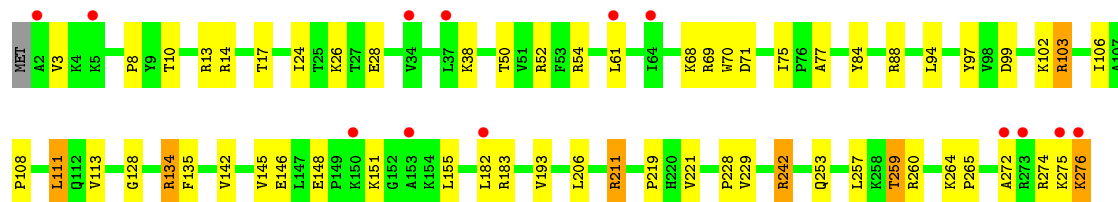
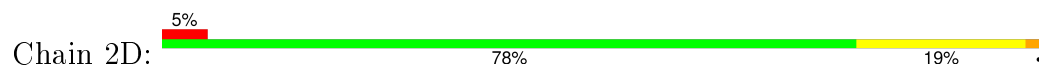




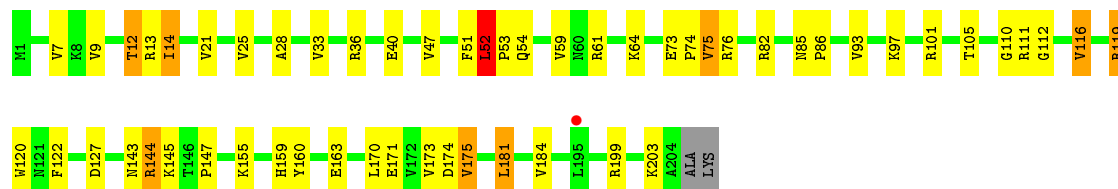




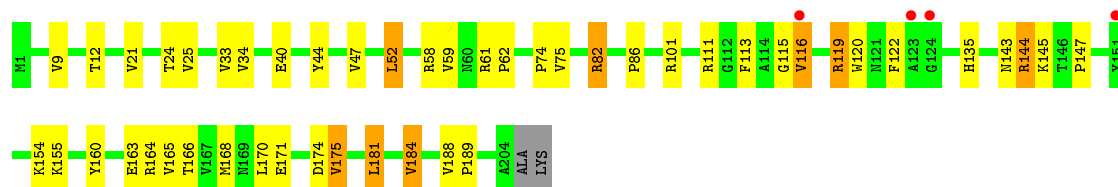
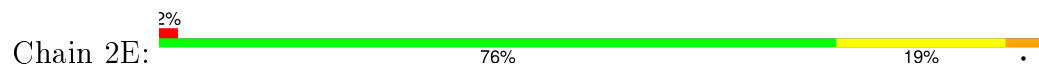
• Molecule 3: 50S ribosomal protein L2



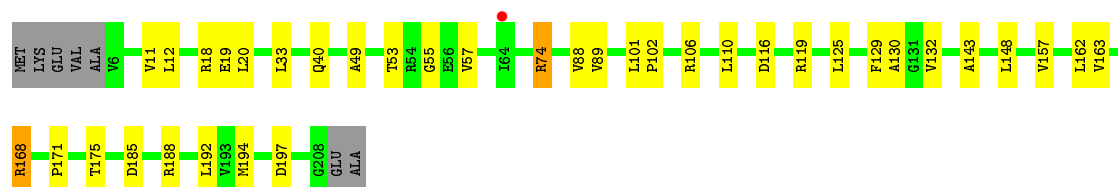
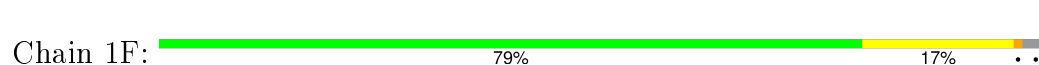
• Molecule 4: 50S ribosomal protein L3



• Molecule 4: 50S ribosomal protein L3

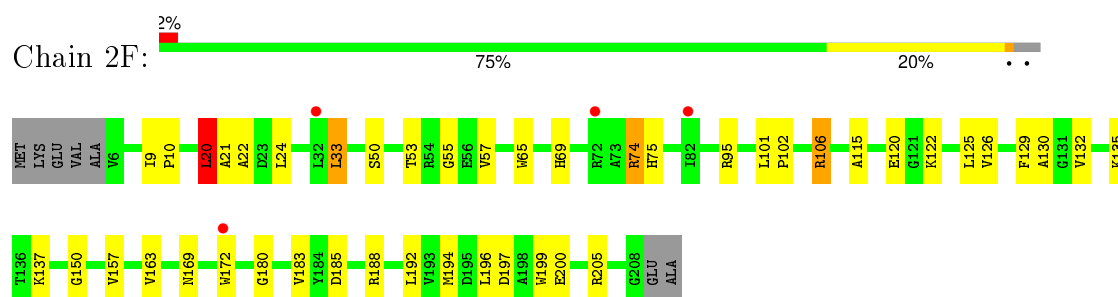


• Molecule 5: 50S ribosomal protein L4

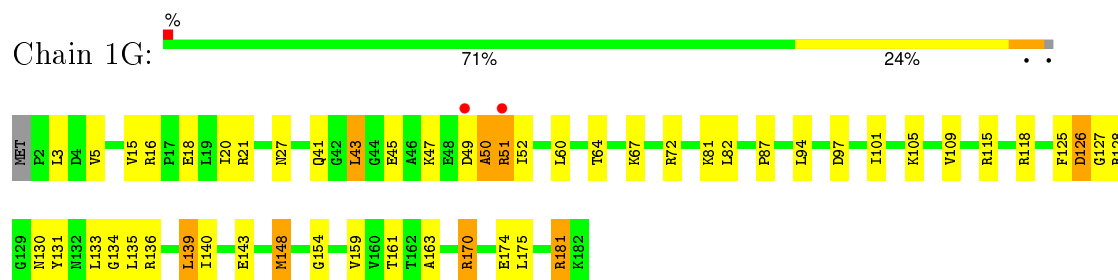


• Molecule 5: 50S ribosomal protein L4

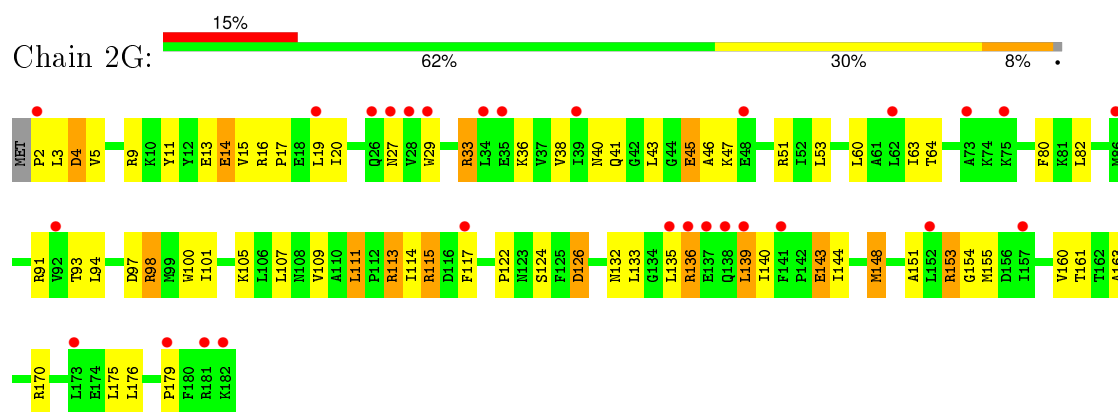




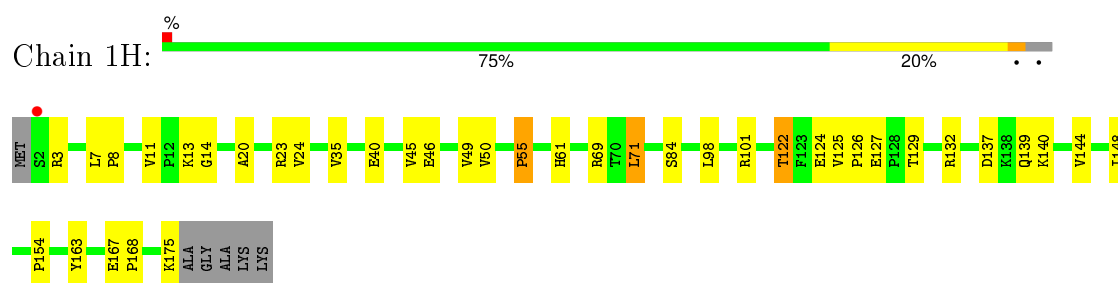
- Molecule 6: 50S ribosomal protein L5



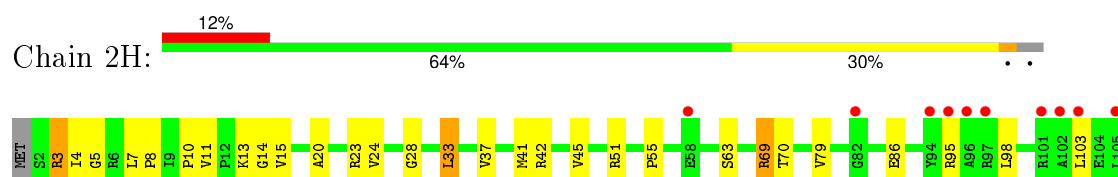
- Molecule 6: 50S ribosomal protein L5



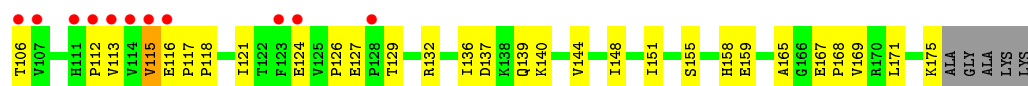
- Molecule 7: 50S ribosomal protein L6



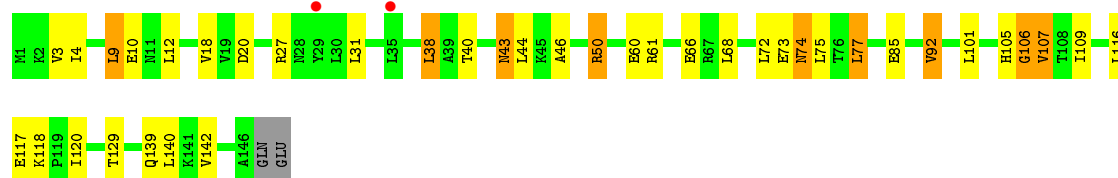
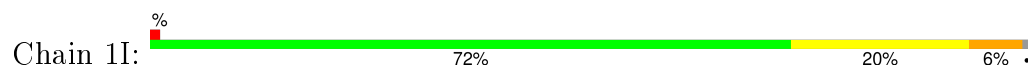
- Molecule 7: 50S ribosomal protein L6



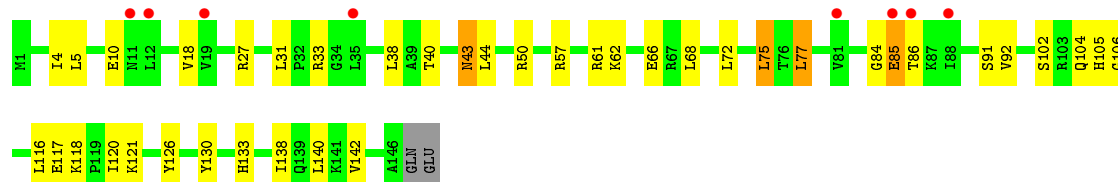




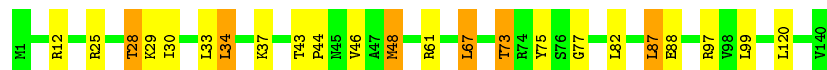
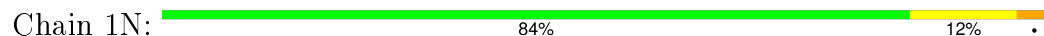
• Molecule 8: 50S ribosomal protein L9



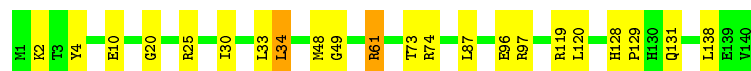
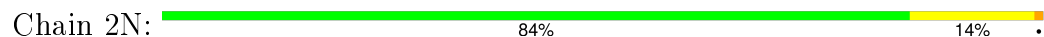
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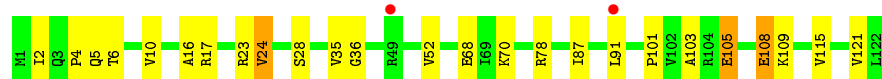
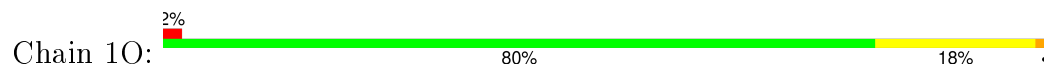
• Molecule 9: 50S ribosomal protein L13



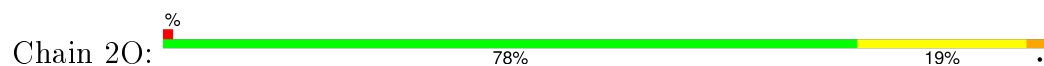
• Molecule 9: 50S ribosomal protein L13



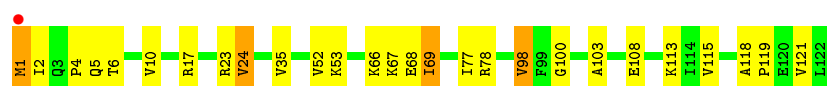
• Molecule 10: 50S ribosomal protein L14



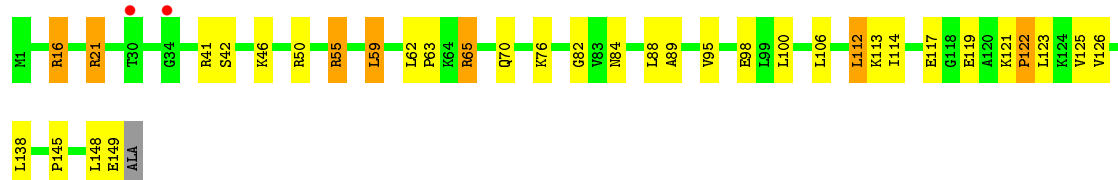
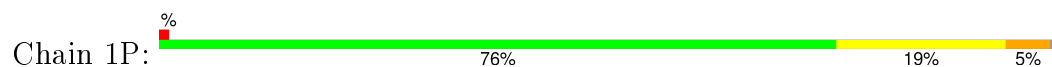
• Molecule 10: 50S ribosomal protein L14



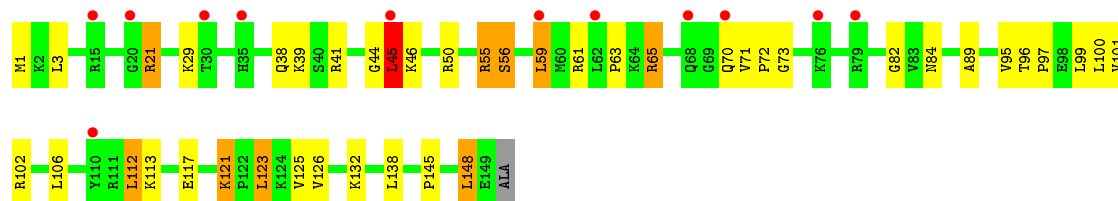




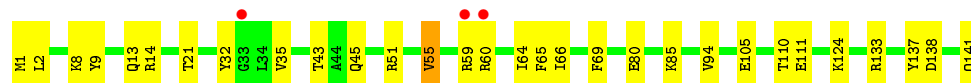
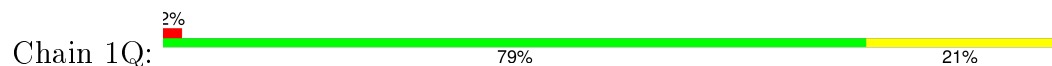
- Molecule 11: 50S ribosomal protein L15



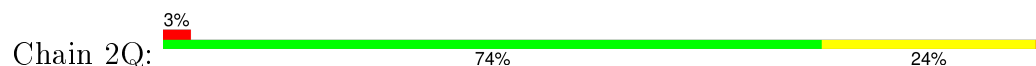
- Molecule 11: 50S ribosomal protein L15



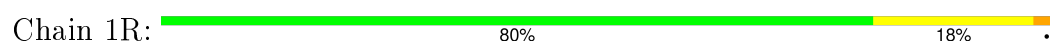
- Molecule 12: 50S ribosomal protein L16



- Molecule 12: 50S ribosomal protein L16

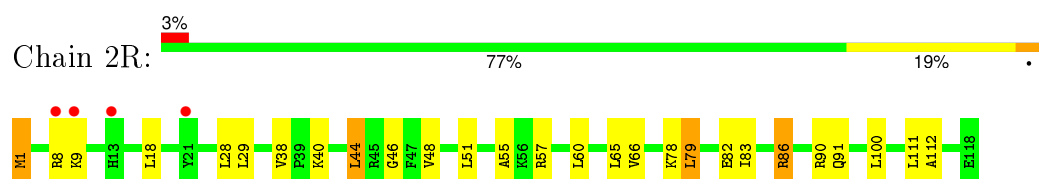


- Molecule 13: 50S ribosomal protein L17

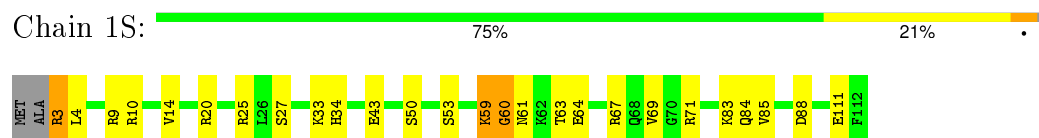


- Molecule 13: 50S ribosomal protein L17

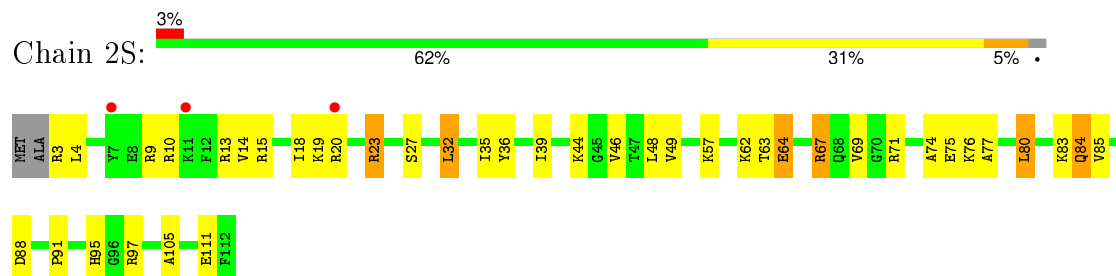




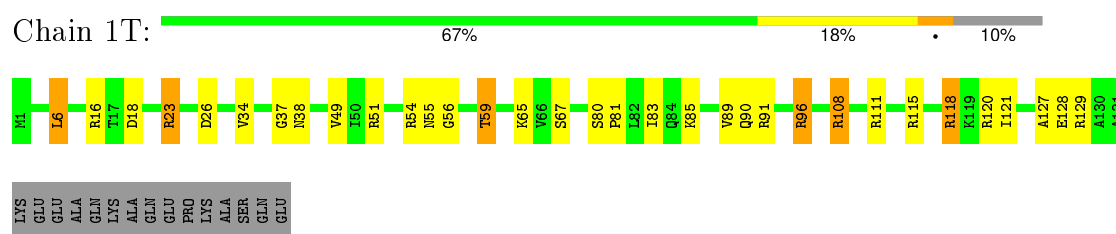
- Molecule 14: 50S ribosomal protein L18



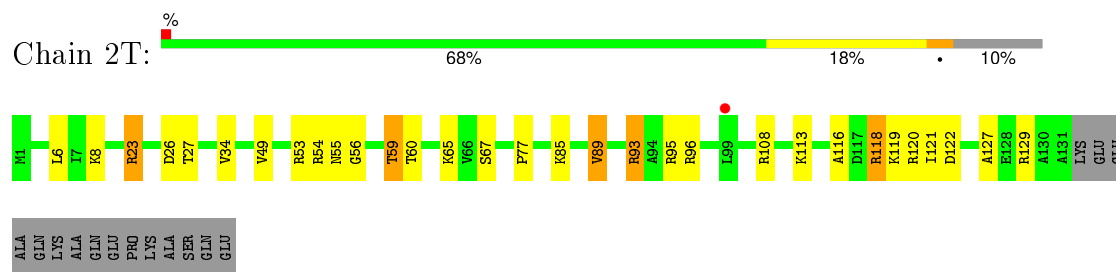
- Molecule 14: 50S ribosomal protein L18



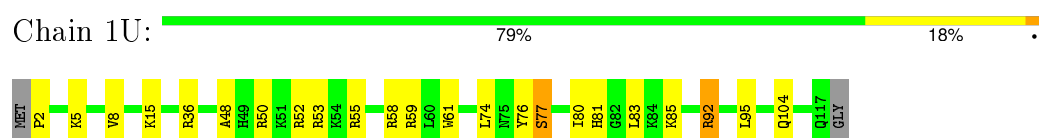
- Molecule 15: 50S ribosomal protein L19



- Molecule 15: 50S ribosomal protein L19




- Molecule 16: 50S ribosomal protein L20




- Molecule 16: 50S ribosomal protein L20



Chain 2U:  81% 16% ..



- Molecule 17: 50S ribosomal protein L21

Chain 1V:  80% 16% ..




- Molecule 17: 50S ribosomal protein L21

Chain 2V:  68% 27% ..




- Molecule 18: 50S ribosomal protein L22

Chain 1W:  83% 15% ..




- Molecule 18: 50S ribosomal protein L22

Chain 2W:  83% 14% ..




- Molecule 19: 50S ribosomal protein L23

Chain 1X:  80% 19% .



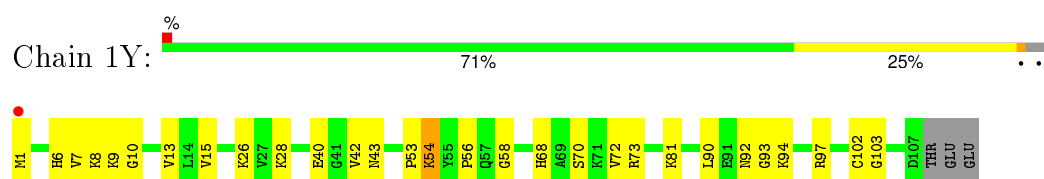
- Molecule 19: 50S ribosomal protein L23

Chain 2X:  76% 21% ..

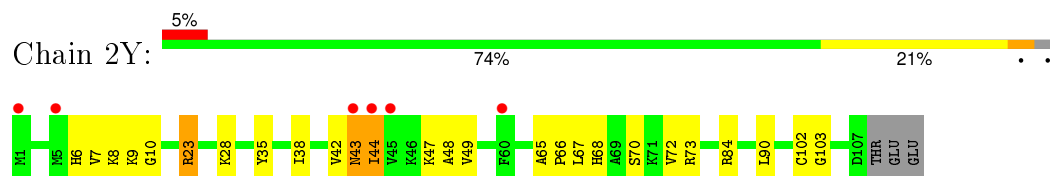


- Molecule 20: 50S ribosomal protein L24

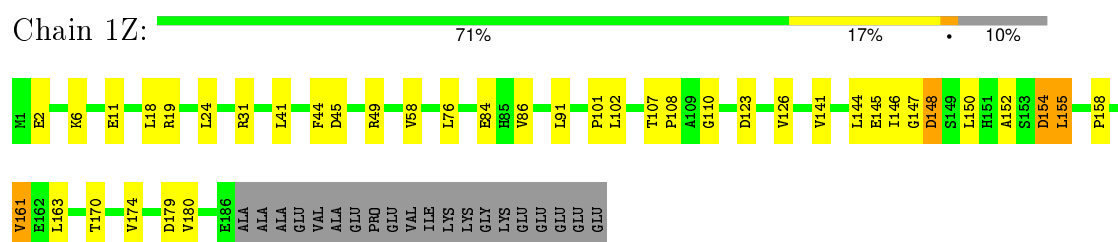




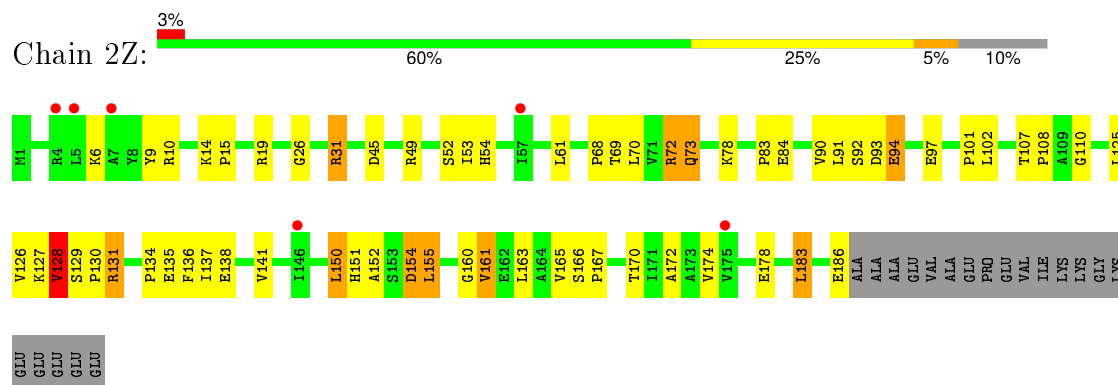
- Molecule 20: 50S ribosomal protein L24



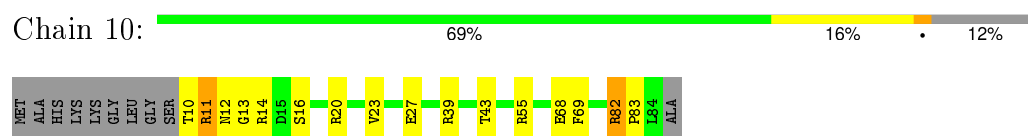
- Molecule 21: 50S ribosomal protein L25



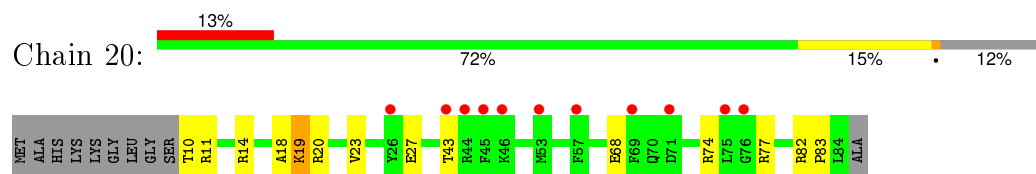
- Molecule 21: 50S ribosomal protein L25



- Molecule 22: 50S ribosomal protein L27

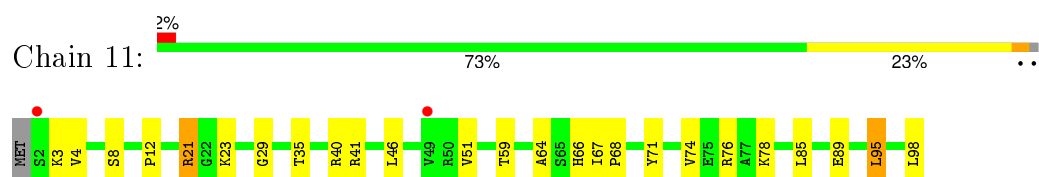


- Molecule 22: 50S ribosomal protein L27

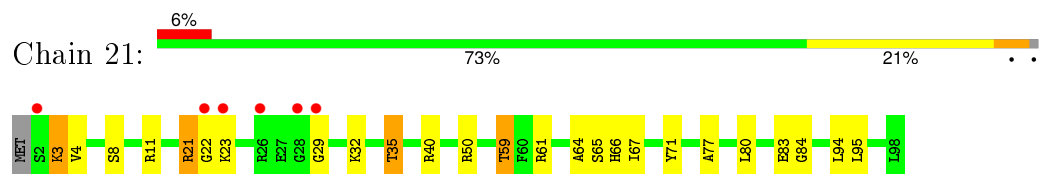


- Molecule 23: 50S ribosomal protein L28

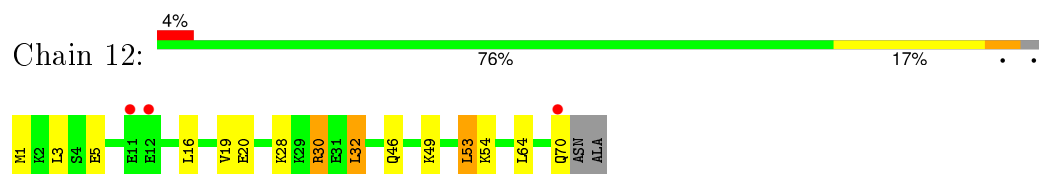




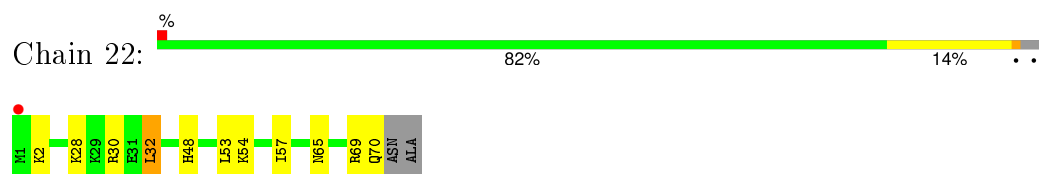
- Molecule 23: 50S ribosomal protein L28



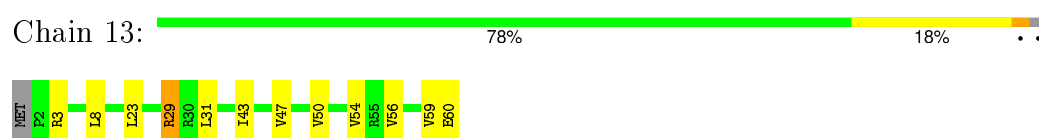
- Molecule 24: 50S ribosomal protein L29



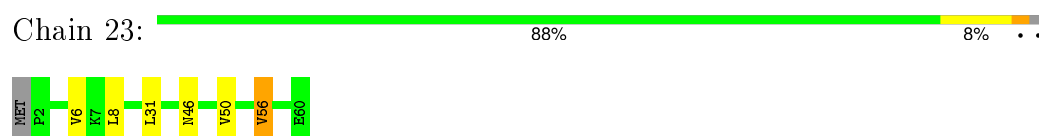
- Molecule 24: 50S ribosomal protein L29



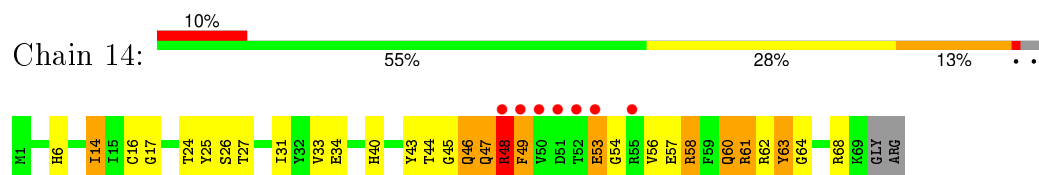
- Molecule 25: 50S ribosomal protein L30



- Molecule 25: 50S ribosomal protein L30

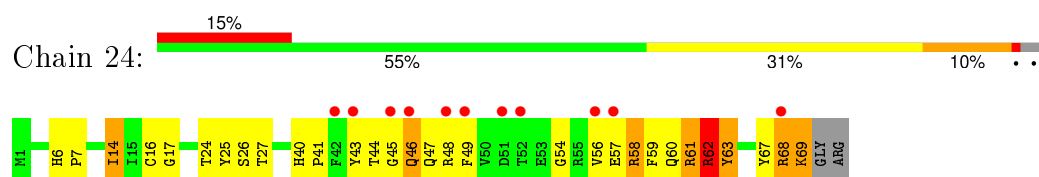


- Molecule 26: 50S ribosomal protein L31

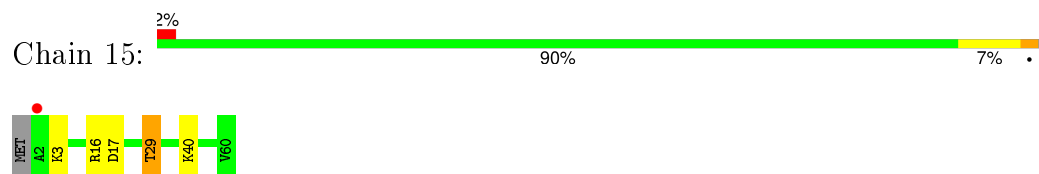


- Molecule 26: 50S ribosomal protein L31

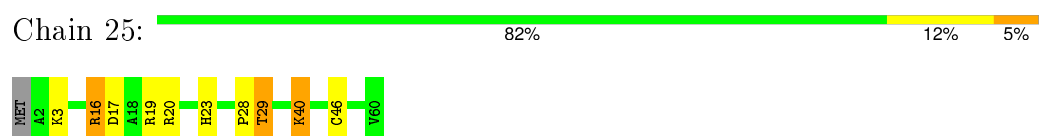




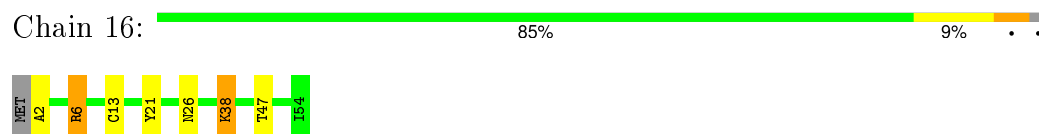
- Molecule 27: 50S ribosomal protein L32



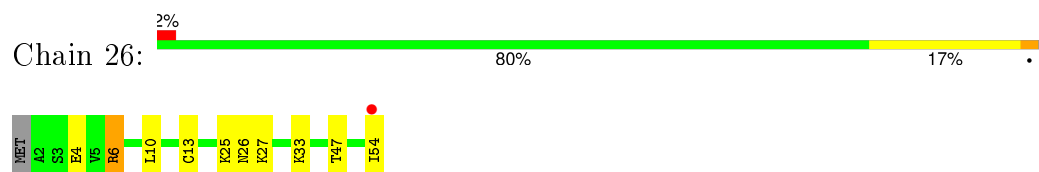
- Molecule 27: 50S ribosomal protein L32



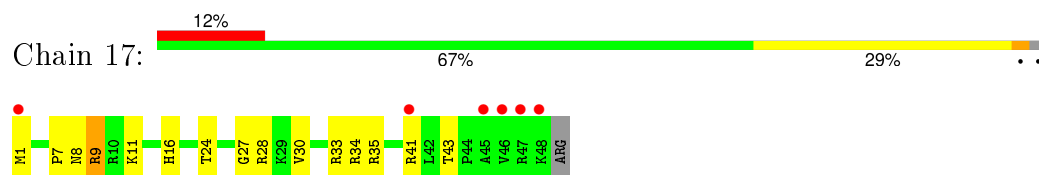
- Molecule 28: 50S ribosomal protein L33



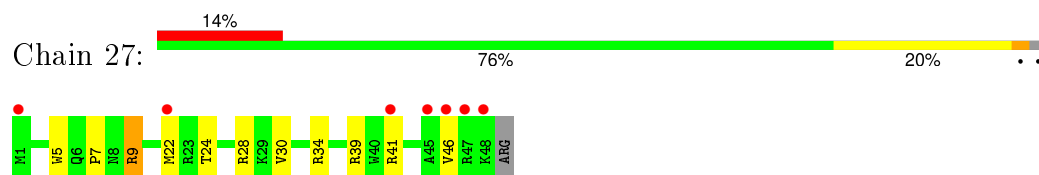
- Molecule 28: 50S ribosomal protein L33



- Molecule 29: 50S ribosomal protein L34

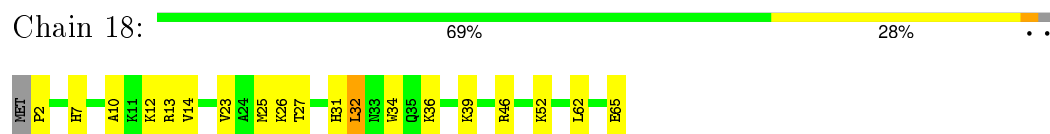


- Molecule 29: 50S ribosomal protein L34

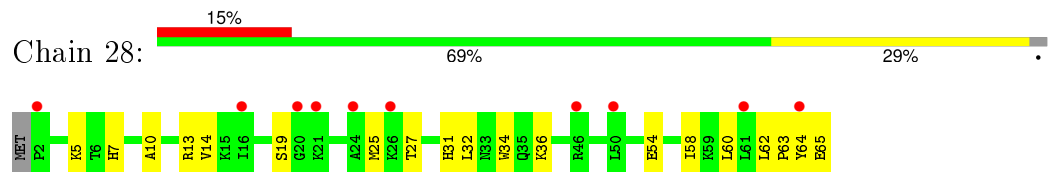


- Molecule 30: 50S ribosomal protein L35

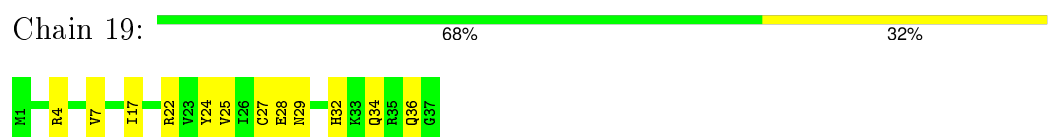




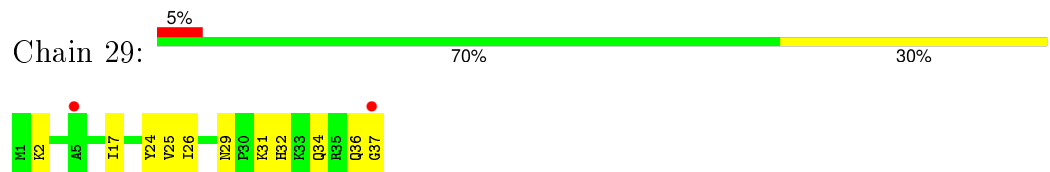
- Molecule 30: 50S ribosomal protein L35



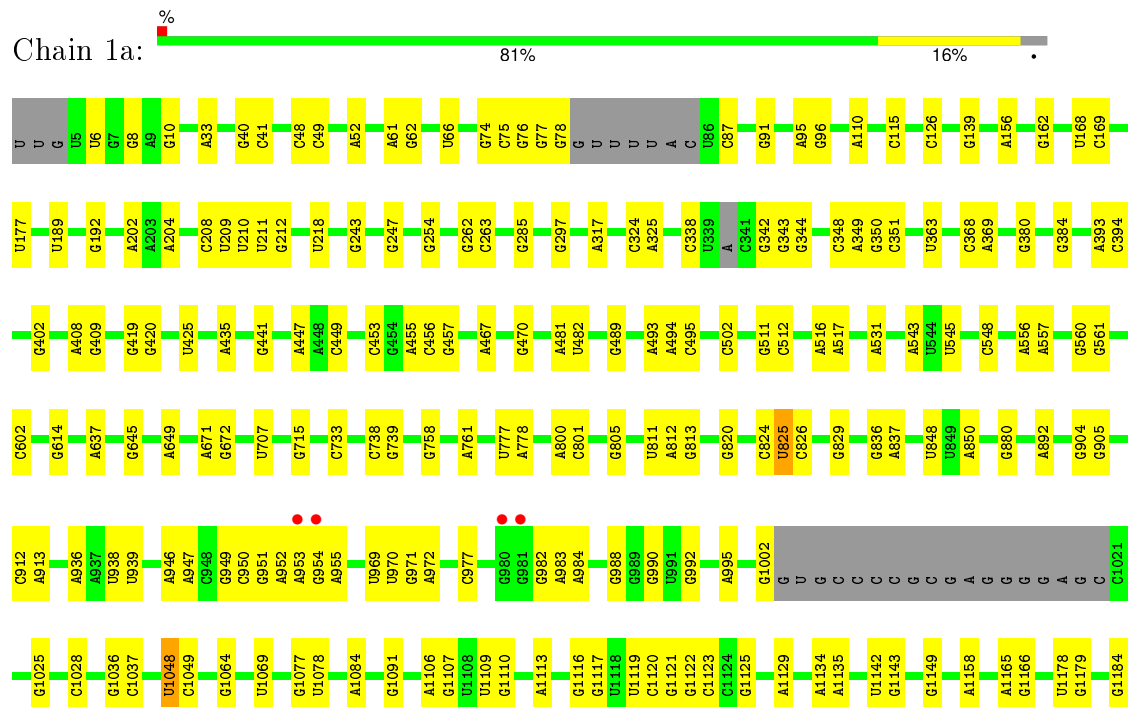
- Molecule 31: 50S ribosomal protein L36



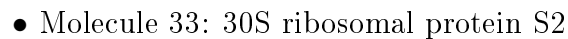
- Molecule 31: 50S ribosomal protein L36



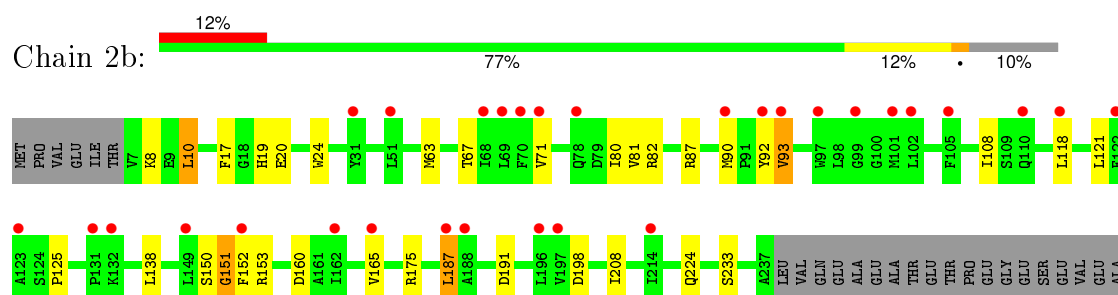
- Molecule 32: 16S ribosomal RNA



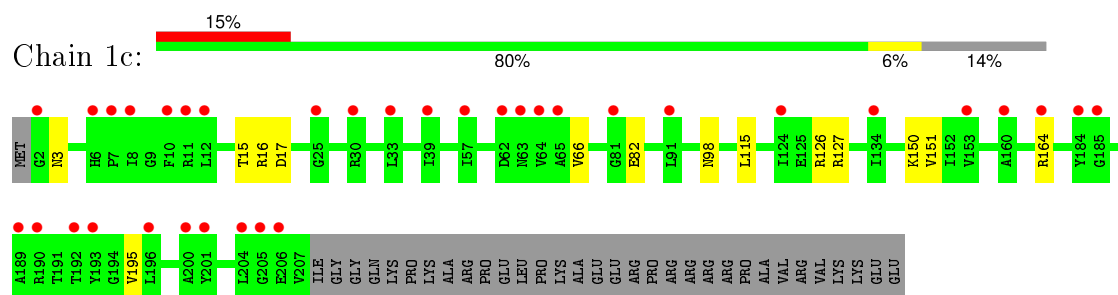




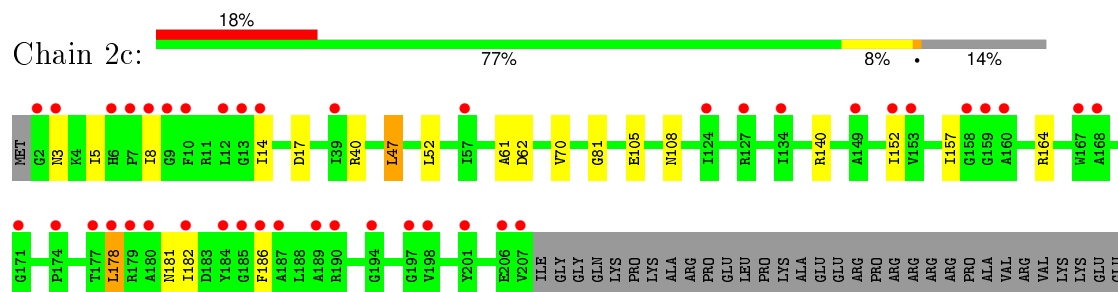




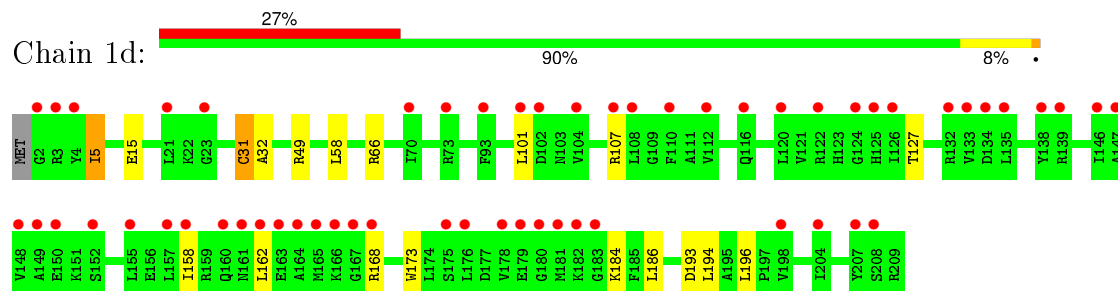
• Molecule 34: 30S ribosomal protein S3



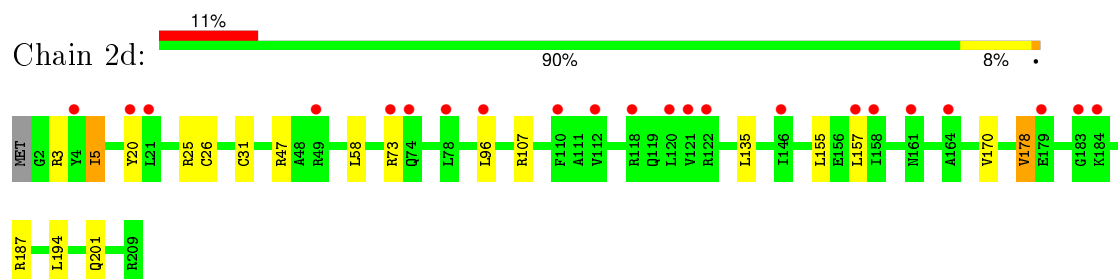
• Molecule 34: 30S ribosomal protein S3



• Molecule 35: 30S ribosomal protein S4

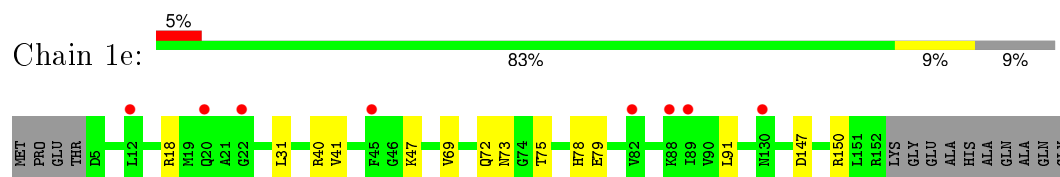


• Molecule 35: 30S ribosomal protein S4

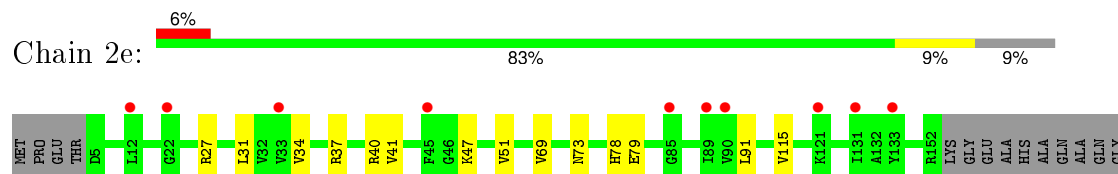




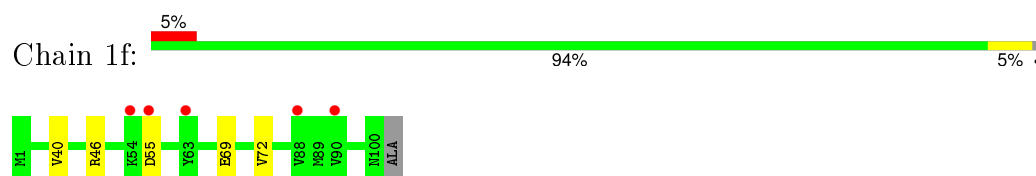
- Molecule 36: 30S ribosomal protein S5



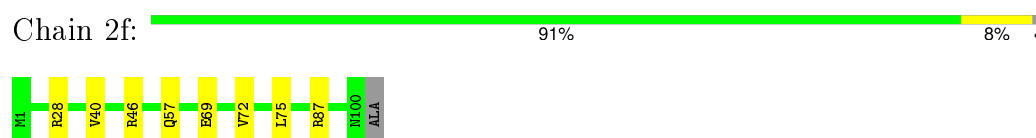
- Molecule 36: 30S ribosomal protein S5



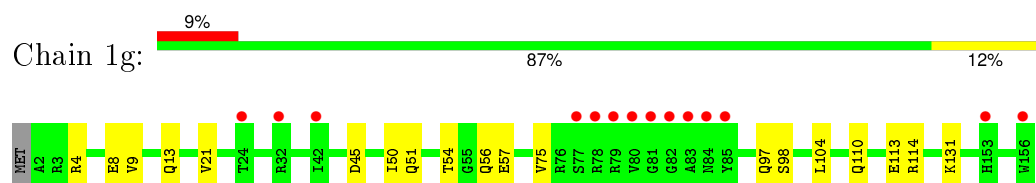
- Molecule 37: 30S ribosomal protein S6



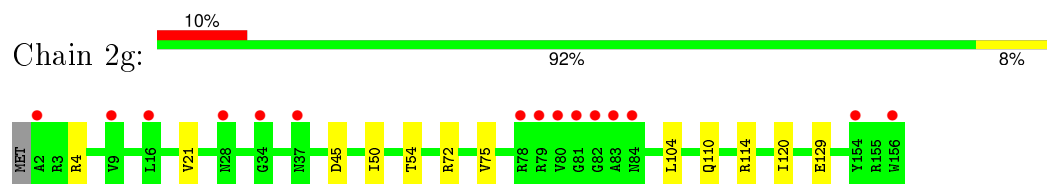
- Molecule 37: 30S ribosomal protein S6



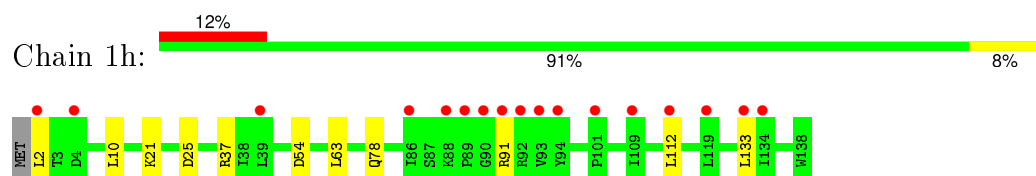
- Molecule 38: 30S ribosomal protein S7



- Molecule 38: 30S ribosomal protein S7

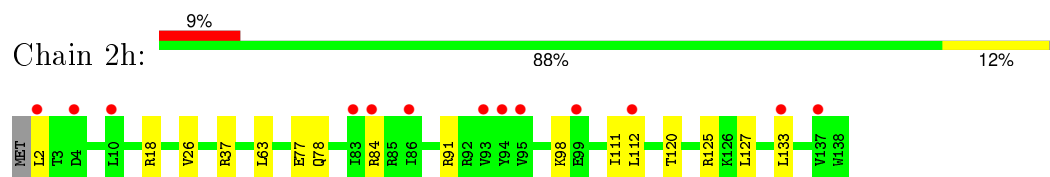


- Molecule 39: 30S ribosomal protein S8

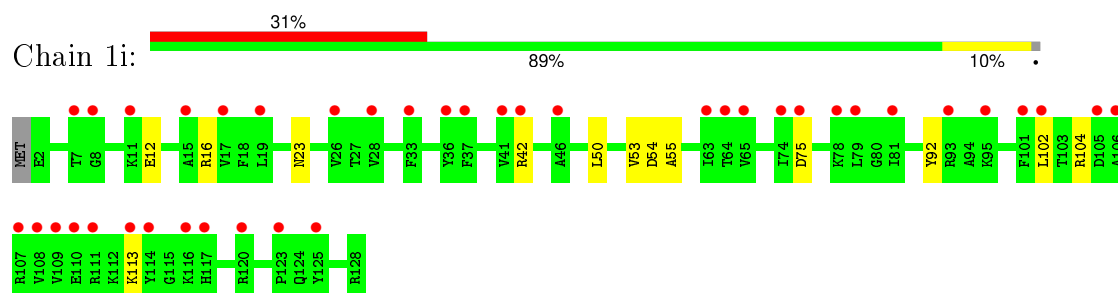




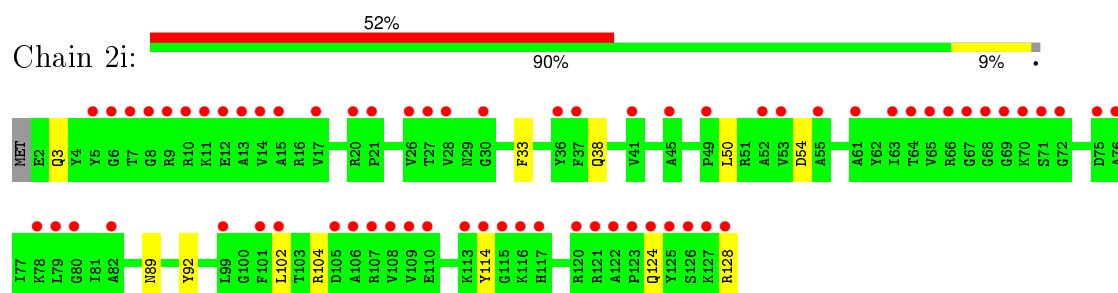
- Molecule 39: 30S ribosomal protein S8



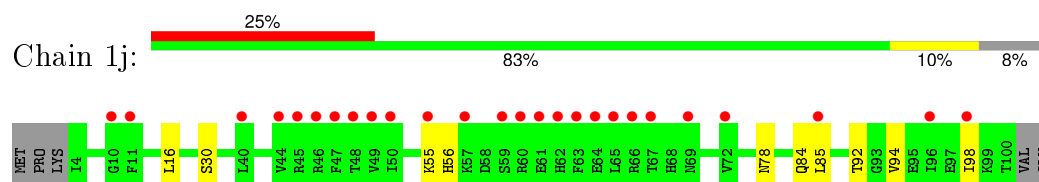
- Molecule 40: 30S ribosomal protein S9



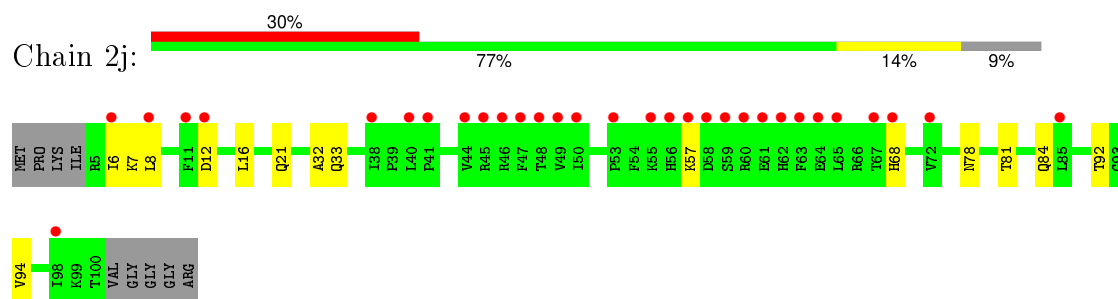
- Molecule 40: 30S ribosomal protein S9



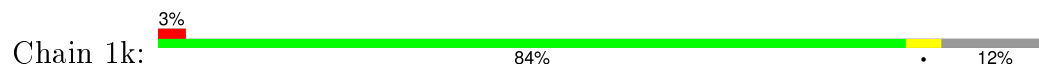
- Molecule 41: 30S ribosomal protein S10



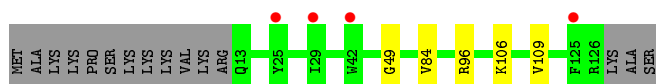
- Molecule 41: 30S ribosomal protein S10



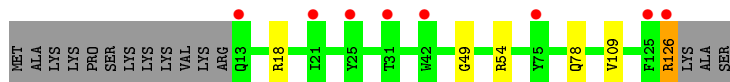
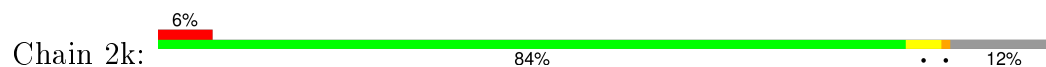
- Molecule 42: 30S ribosomal protein S11



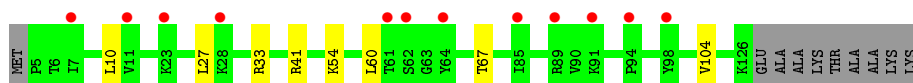
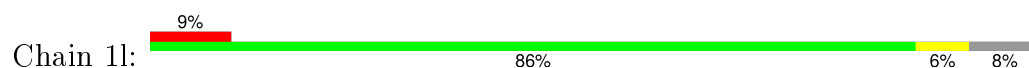




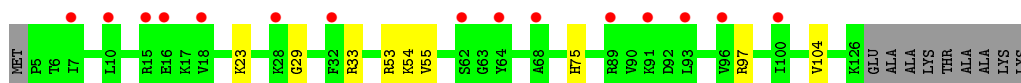
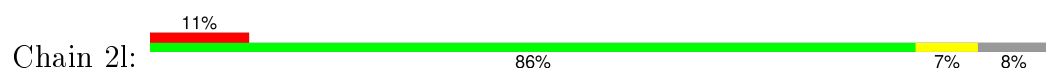
- Molecule 42: 30S ribosomal protein S11



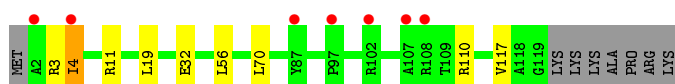
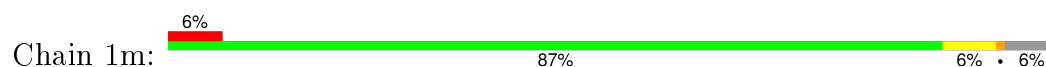
- Molecule 43: 30S ribosomal protein S12



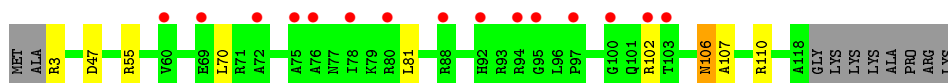
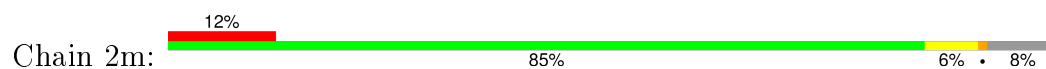
- Molecule 43: 30S ribosomal protein S12



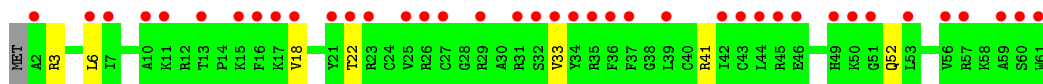
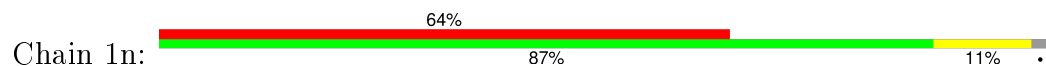
- Molecule 44: 30S ribosomal protein S13



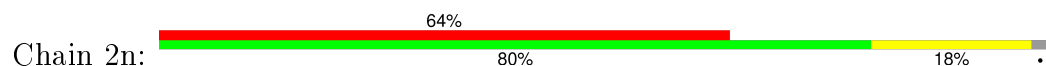
- Molecule 44: 30S ribosomal protein S13



- Molecule 45: 30S ribosomal protein S14 type Z



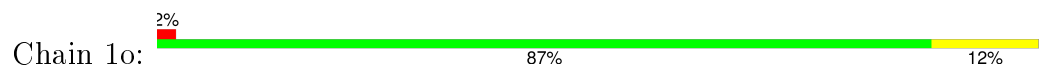
- Molecule 45: 30S ribosomal protein S14 type Z



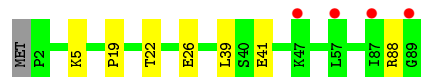
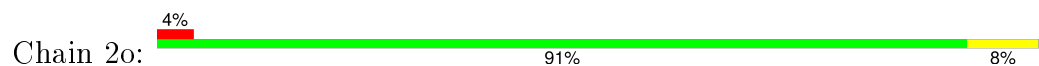




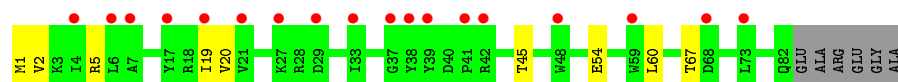
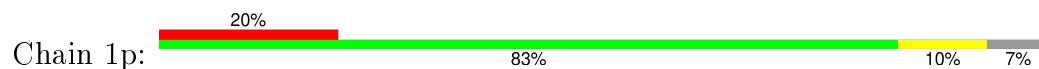
- Molecule 46: 30S ribosomal protein S15



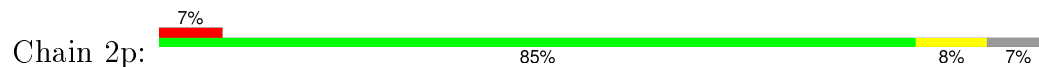
- Molecule 46: 30S ribosomal protein S15



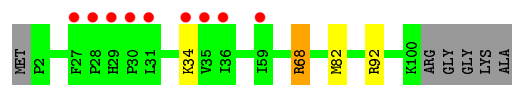
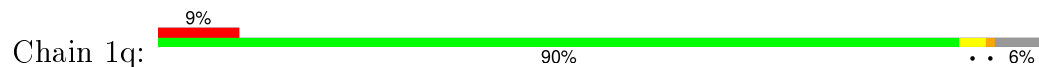
- Molecule 47: 30S ribosomal protein S16



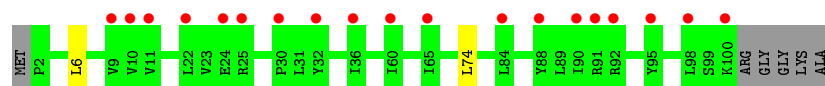
- Molecule 47: 30S ribosomal protein S16



- Molecule 48: 30S ribosomal protein S17



- Molecule 48: 30S ribosomal protein S17



- Molecule 49: 30S ribosomal protein S18



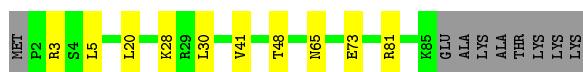
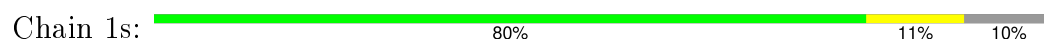




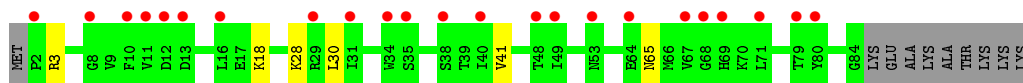
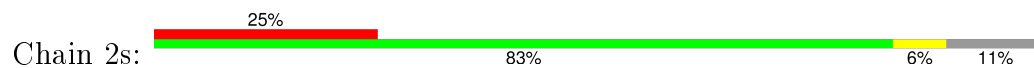
- Molecule 49: 30S ribosomal protein S18



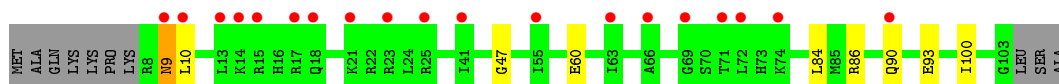
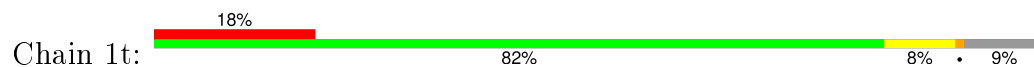
- Molecule 50: 30S ribosomal protein S19



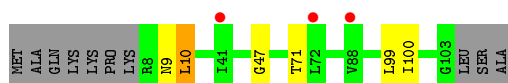
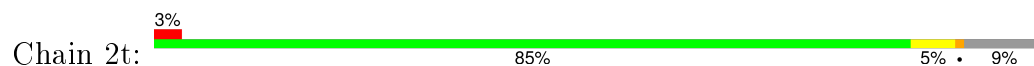
- Molecule 50: 30S ribosomal protein S19



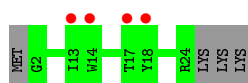
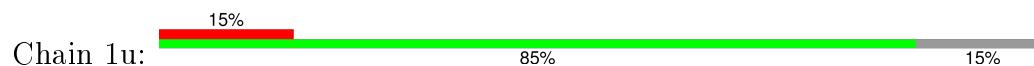
- Molecule 51: 30S ribosomal protein S20



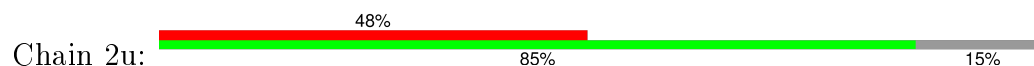
- Molecule 51: 30S ribosomal protein S20



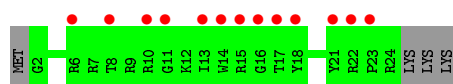
- Molecule 52: 30S ribosomal protein Thx



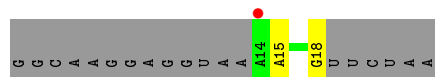
- Molecule 52: 30S ribosomal protein Thx



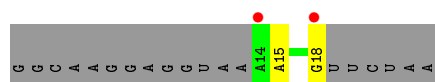




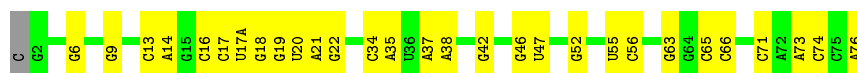
- Molecule 53: mRNA



- Molecule 53: mRNA



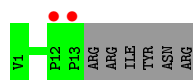
- Molecule 54: Initiator Methionine tRNA



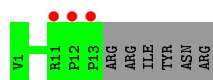
- Molecule 54: Initiator Methionine tRNA



- Molecule 55: Oncocin



- Molecule 55: Oncocin





## 4 Data and refinement statistics

Property	Value	Source
Space group	P 21 21 21	Depositor
Cell constants a, b, c, $\alpha$ , $\beta$ , $\gamma$	210.14Å 450.73Å 623.47Å 90.00° 90.00° 90.00°	Depositor
Resolution (Å)	59.98 – 2.90 311.74 – 2.80	Depositor EDS
% Data completeness (in resolution range)	98.5 (59.98-2.90) 98.0 (311.74-2.80)	Depositor EDS
$R_{merge}$	0.18	Depositor
$R_{sym}$	(Not available)	Depositor
$\langle I/\sigma(I) \rangle$ <sup>1</sup>	1.12 (at 2.82Å)	Xtriage
Refinement program	PHENIX	Depositor
R, $R_{free}$	0.203 , 0.248 0.207 , 0.250	Depositor DCC
$R_{free}$ test set	63771 reflections (5.28%)	DCC
Wilson B-factor (Å <sup>2</sup> )	63.0	Xtriage
Anisotropy	0.092	Xtriage
Bulk solvent $k_{sol}$ (e/Å <sup>3</sup> ), $B_{sol}$ (Å <sup>2</sup> )	0.28 , 56.9	EDS
Estimated twinning fraction	No twinning to report.	Xtriage
L-test for twinning <sup>2</sup>	$\langle  L  \rangle = 0.47$ , $\langle L^2 \rangle = 0.30$	Xtriage
Outliers	0 of 1404775 reflections	Xtriage
$F_o, F_c$ correlation	0.91	EDS
Total number of atoms	288378	wwPDB-VP
Average B, all atoms (Å <sup>2</sup> )	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.*

<sup>1</sup>Intensities estimated from amplitudes.

<sup>2</sup>Theoretical values of  $\langle |L| \rangle$ ,  $\langle L^2 \rangle$  for acentric reflections are 0.5, 0.375 respectively for untwinned datasets, and 0.333, 0.2 for perfectly twinned datasets.



## 5 Model quality ⓘ

### 5.1 Standard geometry ⓘ

Bond lengths and bond angles in the following residue types are not validated in this section: 5MU, ZN, SF4, MG, 5MC, 4SU, PSU

The Z score for a bond length (or angle) is the number of standard deviations the observed value is removed from the expected value. A bond length (or angle) with  $|Z| > 5$  is considered an outlier worth inspection. RMSZ is the root-mean-square of all Z scores of the bond lengths (or angles).

Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# $ Z  > 5$	RMSZ	# $ Z  > 5$
1	1A	0.34	0/66249	0.77	19/103407 (0.0%)
1	2A	0.28	0/67298	0.76	12/105044 (0.0%)
2	1B	0.28	0/2877	0.79	0/4488
2	2B	0.30	0/2878	0.79	0/4490
3	1D	0.29	0/2186	0.49	0/2944
3	2D	0.28	0/2192	0.49	0/2951
4	1E	0.36	0/1592	0.51	1/2149 (0.0%)
4	2E	0.26	0/1592	0.49	0/2149
5	1F	0.27	0/1619	0.49	0/2193
5	2F	0.25	0/1615	0.48	0/2188
6	1G	0.23	0/1450	0.46	0/1959
6	2G	0.26	0/1449	0.49	0/1958
7	1H	0.25	0/1356	0.46	0/1834
7	2H	0.24	0/1356	0.45	0/1834
8	1I	0.24	0/1100	0.49	0/1501
8	2I	0.23	0/1076	0.49	0/1471
9	1N	0.27	0/1144	0.46	0/1543
9	2N	0.25	0/1144	0.46	0/1543
10	1O	0.28	0/943	0.48	0/1269
10	2O	0.27	0/943	0.49	0/1269
11	1P	0.27	0/1156	0.50	0/1537
11	2P	0.27	0/1152	0.49	0/1533
12	1Q	0.28	0/1143	0.46	0/1527
12	2Q	0.27	0/1143	0.47	0/1527
13	1R	0.26	0/982	0.50	0/1312
13	2R	0.25	0/982	0.47	0/1312
14	1S	0.25	0/887	0.48	0/1180
14	2S	0.27	0/880	0.52	0/1172
15	1T	0.27	0/1105	0.45	0/1477
15	2T	0.26	0/1097	0.45	0/1468
16	1U	0.30	0/977	0.45	0/1301
16	2U	0.26	0/977	0.45	0/1301



Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z  >5	RMSZ	# Z  >5
17	1V	0.28	0/782	0.46	0/1049
17	2V	0.25	0/782	0.44	0/1049
18	1W	0.28	0/897	0.44	0/1205
18	2W	0.26	0/897	0.44	0/1205
19	1X	0.28	0/764	0.46	0/1025
19	2X	0.26	0/764	0.48	1/1025 (0.1%)
20	1Y	0.28	0/819	0.51	0/1095
20	2Y	0.25	0/819	0.47	0/1095
21	1Z	0.25	0/1502	0.45	0/2041
21	2Z	0.24	0/1486	0.44	0/2022
22	10	0.28	0/606	0.50	0/808
22	20	0.26	0/606	0.47	0/808
23	11	0.26	0/762	0.45	0/1014
23	21	0.25	0/762	0.43	0/1014
24	12	0.24	0/590	0.43	0/781
24	22	0.25	0/590	0.40	0/781
25	13	0.26	0/474	0.43	0/635
25	23	0.24	0/469	0.44	0/630
26	14	0.29	0/571	0.59	0/768
26	24	0.29	0/545	0.58	0/737
27	15	0.29	0/469	0.51	0/635
27	25	0.25	0/469	0.48	0/635
28	16	0.29	0/460	0.43	0/613
28	26	0.26	0/456	0.45	0/608
29	17	0.28	0/426	0.43	0/561
29	27	0.27	0/426	0.44	0/561
30	18	0.29	0/525	0.47	0/691
30	28	0.26	0/525	0.43	0/691
31	19	0.29	0/310	0.44	0/407
31	29	0.25	0/310	0.43	0/407
32	1a	0.25	0/35537	0.77	7/55456 (0.0%)
32	2a	0.25	2/35680 (0.0%)	0.78	33/55681 (0.1%)
33	1b	0.25	0/1820	0.51	0/2468
33	2b	2.71	8/1728 (0.5%)	0.68	3/2352 (0.1%)
34	1c	0.23	0/1504	0.44	0/2047
34	2c	0.25	0/1435	0.51	2/1960 (0.1%)
35	1d	0.24	0/1648	0.48	0/2222
35	2d	0.25	0/1659	0.45	0/2230
36	1e	0.25	0/1145	0.49	0/1543
36	2e	0.26	0/1111	0.50	0/1504
37	1f	0.24	0/819	0.45	0/1111
37	2f	0.24	0/830	0.44	0/1125
38	1g	0.23	0/1198	0.42	0/1613



Mol	Chain	Bond lengths		Bond angles	
		RMSZ	# Z  >5	RMSZ	# Z  >5
38	2g	0.24	0/1185	0.42	0/1602
39	1h	0.23	0/1108	0.45	0/1494
39	2h	0.23	0/1094	0.47	0/1478
40	1i	0.25	0/995	0.49	0/1339
40	2i	0.26	0/949	0.50	0/1284
41	1j	0.24	0/695	0.55	0/950
41	2j	0.25	0/690	0.50	0/943
42	1k	0.24	0/840	0.45	0/1138
42	2k	0.25	0/844	0.51	1/1145 (0.1%)
43	1l	0.24	0/936	0.46	0/1263
43	2l	0.25	0/934	0.51	1/1262 (0.1%)
44	1m	0.23	0/933	0.49	0/1254
44	2m	0.27	0/913	0.47	0/1230
45	1n	0.29	0/491	0.50	0/653
45	2n	0.27	0/467	0.41	0/624
46	1o	0.24	0/726	0.47	0/970
46	2o	0.23	0/739	0.44	0/985
47	1p	0.22	0/686	0.46	0/926
47	2p	0.23	0/693	0.47	0/935
48	1q	0.25	0/824	0.46	0/1105
48	2q	0.24	0/836	0.44	0/1117
49	1r	0.25	0/560	0.48	0/746
49	2r	0.23	0/560	0.44	0/746
50	1s	0.24	0/657	0.52	0/890
50	2s	0.26	0/661	0.53	0/893
51	1t	0.22	0/714	0.52	0/948
51	2t	0.23	0/733	0.44	0/969
52	1u	0.21	0/191	0.45	0/252
52	2u	0.24	0/203	0.42	0/266
53	1v	0.39	0/122	1.03	0/188
53	2v	0.41	0/122	1.09	0/188
54	1x	0.41	0/1725	1.03	10/2689 (0.4%)
54	2x	0.41	0/1725	1.04	11/2689 (0.4%)
55	1z	0.29	0/114	0.54	0/158
55	2z	0.24	0/114	0.48	0/158
All	All	0.35	10/306296 (0.0%)	0.71	101/458216 (0.0%)

Chiral center outliers are detected by calculating the chiral volume of a chiral center and verifying if the center is modelled as a planar moiety or with the opposite hand. A planarity outlier is detected by checking planarity of atoms in a peptide group, atoms in a mainchain group or atoms of a sidechain that are expected to be planar.



Mol	Chain	#Chirality outliers	#Planarity outliers
5	2F	0	1
41	2j	0	1
All	All	0	2

The worst 5 of 10 bond length outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
33	2b	92	TYR	CD1-CE1	58.43	2.27	1.39
33	2b	92	TYR	CD2-CE2	54.24	2.20	1.39
33	2b	92	TYR	CE2-CZ	42.19	1.93	1.38
33	2b	92	TYR	CE1-CZ	41.20	1.92	1.38
33	2b	92	TYR	CG-CD1	33.29	1.82	1.39

The worst 5 of 101 bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
33	2b	150	SER	C-N-CA	16.60	157.16	122.30
32	2a	1102	C	N1-C2-O2	14.44	127.57	118.90
32	2a	1137	G	N3-C2-N2	13.03	129.02	119.90
32	2a	1137	G	N1-C2-N2	-12.03	105.38	116.20
32	2a	1137	G	C4-N9-C1'	11.15	141.00	126.50

There are no chirality outliers.

All (2) planarity outliers are listed below:

Mol	Chain	Res	Type	Group
5	2F	20	LEU	Peptide
41	2j	32	ALA	Peptide

## 5.2 Too-close contacts [\(i\)](#)

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

Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
1	1A	59154	0	29830	434	0
1	2A	60091	0	30300	592	0
2	1B	2572	0	1306	18	0
2	2B	2573	0	1306	34	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
3	1D	2136	0	2218	38	0
3	2D	2142	0	2229	39	0
4	1E	1559	0	1618	31	0
4	2E	1559	0	1618	27	0
5	1F	1584	0	1625	21	0
5	2F	1580	0	1619	25	0
6	1G	1425	0	1443	28	0
6	2G	1424	0	1434	48	0
7	1H	1330	0	1407	19	0
7	2H	1330	0	1407	31	0
8	1I	1085	0	1114	17	0
8	2I	1061	0	1080	24	0
9	1N	1117	0	1184	13	0
9	2N	1117	0	1184	11	0
10	1O	933	0	996	16	0
10	2O	933	0	996	17	0
11	1P	1139	0	1223	27	0
11	2P	1135	0	1212	38	0
12	1Q	1122	0	1178	19	0
12	2Q	1122	0	1179	22	0
13	1R	968	0	1033	8	0
13	2R	968	0	1033	12	0
14	1S	877	0	938	18	0
14	2S	870	0	923	23	0
15	1T	1091	0	1151	21	0
15	2T	1083	0	1136	19	0
16	1U	959	0	1019	10	0
16	2U	959	0	1019	15	0
17	1V	771	0	830	9	0
17	2V	771	0	830	18	0
18	1W	886	0	940	10	0
18	2W	886	0	940	8	0
19	1X	750	0	814	13	0
19	2X	750	0	814	14	0
20	1Y	806	0	881	16	0
20	2Y	806	0	881	18	0
21	1Z	1470	0	1478	20	0
21	2Z	1454	0	1452	41	0
22	10	598	0	614	11	0
22	20	598	0	614	11	0
23	11	755	0	826	16	0
23	21	755	0	826	15	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
24	12	588	0	643	6	0
24	22	588	0	643	4	0
25	13	469	0	518	8	0
25	23	464	0	514	2	0
26	14	558	0	544	20	0
26	24	532	0	503	23	0
27	15	455	0	465	7	0
27	25	455	0	465	9	0
28	16	453	0	473	5	0
28	26	449	0	469	6	0
29	17	418	0	467	10	0
29	27	418	0	467	8	0
30	18	517	0	582	16	0
30	28	517	0	582	14	0
31	19	307	0	335	8	0
31	29	307	0	335	8	0
32	1a	31750	0	16028	0	0
32	2a	31877	0	16086	0	0
33	1b	1786	0	1744	0	0
33	2b	1697	0	1574	0	0
34	1c	1480	0	1400	0	0
34	2c	1412	0	1246	0	0
35	1d	1618	0	1579	0	0
35	2d	1630	0	1633	0	0
36	1e	1129	0	1184	0	0
36	2e	1095	0	1124	0	0
37	1f	806	0	793	0	0
37	2f	817	0	808	0	0
38	1g	1183	0	1165	0	0
38	2g	1167	0	1119	0	0
39	1h	1088	0	1126	0	0
39	2h	1074	0	1100	0	0
40	1i	976	0	973	0	0
40	2i	932	0	891	0	0
41	1j	682	0	598	0	0
41	2j	678	0	612	0	0
42	1k	826	0	829	0	0
42	2k	829	0	825	0	0
43	1l	920	0	958	0	0
43	2l	918	0	947	0	0
44	1m	923	0	962	0	0
44	2m	903	0	923	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
45	1n	482	0	507	0	0
45	2n	459	0	467	0	0
46	1o	715	0	729	0	0
46	2o	728	0	760	0	0
47	1p	671	0	679	0	0
47	2p	677	0	686	0	0
48	1q	811	0	858	0	0
48	2q	823	0	891	0	0
49	1r	555	0	618	0	0
49	2r	555	0	618	0	0
50	1s	642	0	629	0	0
50	2s	646	0	644	0	0
51	1t	712	0	759	0	0
51	2t	731	0	807	0	0
52	1u	187	0	186	0	0
52	2u	199	0	208	0	0
53	1v	109	0	55	0	0
53	2v	109	0	55	0	0
54	1x	1625	0	829	0	0
54	2x	1625	0	829	0	0
55	1z	108	0	116	0	0
55	2z	108	0	116	0	0
56	10	7	0	0	0	0
56	11	1	0	0	0	0
56	12	1	0	0	0	0
56	13	2	0	0	0	0
56	15	4	0	0	0	0
56	16	2	0	0	0	0
56	17	1	0	0	0	0
56	18	3	0	0	0	0
56	19	3	0	0	0	0
56	1A	1021	0	0	0	0
56	1B	24	0	0	0	0
56	1D	14	0	0	0	0
56	1E	5	0	0	0	0
56	1F	8	0	0	0	0
56	1G	3	0	0	0	0
56	1H	2	0	0	0	0
56	1N	6	0	0	0	0
56	1O	2	0	0	0	0
56	1P	6	0	0	0	0
56	1Q	4	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
56	1R	4	0	0	0	0
56	1T	5	0	0	0	0
56	1U	5	0	0	0	0
56	1V	1	0	0	0	0
56	1W	5	0	0	0	0
56	1X	2	0	0	0	0
56	1Z	2	0	0	0	0
56	1a	276	0	0	0	0
56	1b	2	0	0	0	0
56	1d	2	0	0	0	0
56	1e	3	0	0	0	0
56	1f	1	0	0	0	0
56	1k	1	0	0	0	0
56	1l	1	0	0	0	0
56	1n	1	0	0	0	0
56	1o	2	0	0	0	0
56	1q	4	0	0	0	0
56	1r	2	0	0	0	0
56	1t	1	0	0	0	0
56	1x	8	0	0	0	0
56	1z	1	0	0	0	0
56	20	1	0	0	0	0
56	25	1	0	0	0	0
56	26	1	0	0	0	0
56	28	1	0	0	0	0
56	2A	566	0	0	0	0
56	2B	11	0	0	0	0
56	2D	6	0	0	0	0
56	2E	4	0	0	0	0
56	2F	6	0	0	0	0
56	2G	1	0	0	0	0
56	2N	1	0	0	0	0
56	2O	3	0	0	0	0
56	2P	1	0	0	0	0
56	2Q	5	0	0	0	0
56	2U	1	0	0	0	0
56	2Y	1	0	0	0	0
56	2Z	1	0	0	0	0
56	2a	263	0	0	0	0
56	2d	1	0	0	0	0
56	2f	1	0	0	0	0
56	2h	1	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
56	2k	1	0	0	0	0
56	2l	4	0	0	0	0
56	2q	2	0	0	0	0
56	2t	1	0	0	0	0
56	2x	4	0	0	0	0
56	2z	1	0	0	0	0
57	14	1	0	0	0	0
57	15	1	0	0	0	0
57	16	1	0	0	0	0
57	19	1	0	0	0	0
57	1Y	1	0	0	0	0
57	1n	1	0	0	0	0
57	24	1	0	0	0	0
57	25	1	0	0	0	0
57	26	1	0	0	0	0
57	29	1	0	0	0	0
57	2Y	1	0	0	0	0
57	2n	1	0	0	0	0
58	1d	8	0	0	0	0
58	2d	8	0	0	0	0
59	10	10	0	0	1	0
59	11	4	0	0	0	0
59	12	1	0	0	0	0
59	13	4	0	0	1	0
59	15	5	0	0	0	0
59	16	8	0	0	1	0
59	17	8	0	0	2	0
59	18	13	0	0	0	0
59	19	2	0	0	0	0
59	1A	2088	0	0	61	0
59	1B	38	0	0	4	0
59	1D	22	0	0	0	0
59	1E	25	0	0	4	0
59	1F	16	0	0	0	0
59	1G	6	0	0	0	0
59	1H	5	0	0	1	0
59	1I	1	0	0	0	0
59	1N	5	0	0	0	0
59	1O	4	0	0	0	0
59	1P	21	0	0	1	0
59	1Q	9	0	0	0	0
59	1R	9	0	0	1	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
59	1S	1	0	0	0	0
59	1T	11	0	0	2	0
59	1U	13	0	0	0	0
59	1V	2	0	0	0	0
59	1W	9	0	0	0	0
59	1X	5	0	0	0	0
59	1Y	1	0	0	0	0
59	1Z	3	0	0	0	0
59	1a	319	0	0	0	0
59	1b	2	0	0	0	0
59	1c	1	0	0	0	0
59	1d	4	0	0	0	0
59	1e	2	0	0	0	0
59	1f	1	0	0	0	0
59	1i	1	0	0	0	0
59	1j	1	0	0	0	0
59	1m	1	0	0	0	0
59	1o	3	0	0	0	0
59	1p	1	0	0	0	0
59	1s	1	0	0	0	0
59	1v	1	0	0	0	0
59	1x	7	0	0	0	0
59	1z	2	0	0	0	0
59	20	3	0	0	0	0
59	21	1	0	0	0	0
59	23	2	0	0	0	0
59	25	1	0	0	0	0
59	27	2	0	0	0	0
59	28	4	0	0	0	0
59	2A	801	0	0	45	0
59	2B	13	0	0	0	0
59	2D	20	0	0	1	0
59	2E	10	0	0	1	0
59	2F	7	0	0	0	0
59	2N	1	0	0	0	0
59	2O	4	0	0	0	0
59	2P	4	0	0	0	0
59	2Q	2	0	0	0	0
59	2R	3	0	0	0	0
59	2T	3	0	0	0	0
59	2U	2	0	0	1	0
59	2V	2	0	0	0	0

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Mol	Chain	Non-H	H(model)	H(added)	Clashes	Symm-Clashes
59	2W	2	0	0	0	0
59	2X	3	0	0	0	0
59	2Y	1	0	0	0	0
59	2Z	2	0	0	0	0
59	2a	371	0	0	0	0
59	2c	2	0	0	0	0
59	2e	3	0	0	0	0
59	2h	1	0	0	0	0
59	2i	2	0	0	0	0
59	2j	1	0	0	0	0
59	2k	3	0	0	0	0
59	2l	3	0	0	0	0
59	2m	1	0	0	0	0
59	2n	1	0	0	0	0
59	2o	1	0	0	0	0
59	2p	2	0	0	0	0
59	2q	2	0	0	0	0
59	2r	1	0	0	0	0
59	2t	3	0	0	0	0
59	2x	3	0	0	0	0
59	2z	1	0	0	0	0
All	All	288378	0	189976	1793	0

The all-atom clashscore is defined as the number of clashes found per 1000 atoms (including hydrogen atoms). The all-atom clashscore for this structure is 7.

The worst 5 of 1793 close contacts within the same asymmetric unit are listed below, sorted by their clash magnitude.

Atom-1	Atom-2	Interatomic distance (Å)	Clash overlap (Å)
1:2A:1098:C:N3	1:2A:1099:A:C8	2.28	1.01
1:2A:1096:G:C5	1:2A:1097:C:C5	2.56	0.94
1:2A:1151:G:C2	1:2A:1152:G:N7	2.38	0.92
1:2A:1096:G:C4	1:2A:1097:C:C5	2.61	0.88
1:2A:1151:G:H2'	1:2A:1152:G:H5'	1.55	0.88

There are no symmetry-related clashes.



## 5.3 Torsion angles ⓘ

### 5.3.1 Protein backbone ⓘ

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

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

Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
3	1D	273/276 (99%)	262 (96%)	11 (4%)	0	100	100
3	2D	273/276 (99%)	262 (96%)	10 (4%)	1 (0%)	39	74
4	1E	202/206 (98%)	194 (96%)	7 (4%)	1 (0%)	34	71
4	2E	202/206 (98%)	196 (97%)	5 (2%)	1 (0%)	34	71
5	1F	201/210 (96%)	193 (96%)	7 (4%)	1 (0%)	34	71
5	2F	201/210 (96%)	194 (96%)	6 (3%)	1 (0%)	34	71
6	1G	179/182 (98%)	161 (90%)	14 (8%)	4 (2%)	8	31
6	2G	179/182 (98%)	159 (89%)	16 (9%)	4 (2%)	8	31
7	1H	172/180 (96%)	156 (91%)	14 (8%)	2 (1%)	16	48
7	2H	172/180 (96%)	157 (91%)	13 (8%)	2 (1%)	16	48
8	1I	144/148 (97%)	124 (86%)	15 (10%)	5 (4%)	4	18
8	2I	144/148 (97%)	123 (85%)	19 (13%)	2 (1%)	14	44
9	1N	138/140 (99%)	135 (98%)	3 (2%)	0	100	100
9	2N	138/140 (99%)	133 (96%)	4 (3%)	1 (1%)	26	63
10	1O	120/122 (98%)	116 (97%)	4 (3%)	0	100	100
10	2O	120/122 (98%)	116 (97%)	4 (3%)	0	100	100
11	1P	147/150 (98%)	137 (93%)	9 (6%)	1 (1%)	26	63
11	2P	147/150 (98%)	136 (92%)	10 (7%)	1 (1%)	26	63
12	1Q	139/141 (99%)	131 (94%)	8 (6%)	0	100	100
12	2Q	139/141 (99%)	130 (94%)	8 (6%)	1 (1%)	26	63
13	1R	116/118 (98%)	111 (96%)	5 (4%)	0	100	100
13	2R	116/118 (98%)	111 (96%)	5 (4%)	0	100	100
14	1S	108/112 (96%)	99 (92%)	8 (7%)	1 (1%)	21	57
14	2S	108/112 (96%)	97 (90%)	6 (6%)	5 (5%)	3	11
15	1T	129/146 (88%)	125 (97%)	4 (3%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
15	2T	129/146 (88%)	124 (96%)	5 (4%)	0	100	100
16	1U	114/118 (97%)	113 (99%)	1 (1%)	0	100	100
16	2U	114/118 (97%)	112 (98%)	2 (2%)	0	100	100
17	1V	99/101 (98%)	95 (96%)	3 (3%)	1 (1%)	19	54
17	2V	99/101 (98%)	96 (97%)	2 (2%)	1 (1%)	19	54
18	1W	110/113 (97%)	110 (100%)	0	0	100	100
18	2W	110/113 (97%)	110 (100%)	0	0	100	100
19	1X	93/96 (97%)	92 (99%)	0	1 (1%)	17	51
19	2X	93/96 (97%)	91 (98%)	0	2 (2%)	8	31
20	1Y	105/110 (96%)	98 (93%)	5 (5%)	2 (2%)	10	35
20	2Y	105/110 (96%)	100 (95%)	5 (5%)	0	100	100
21	1Z	184/206 (89%)	172 (94%)	12 (6%)	0	100	100
21	2Z	184/206 (89%)	172 (94%)	11 (6%)	1 (0%)	34	71
22	10	73/85 (86%)	69 (94%)	3 (4%)	1 (1%)	14	44
22	20	73/85 (86%)	69 (94%)	4 (6%)	0	100	100
23	11	95/98 (97%)	93 (98%)	2 (2%)	0	100	100
23	21	95/98 (97%)	92 (97%)	2 (2%)	1 (1%)	17	51
24	12	68/72 (94%)	67 (98%)	1 (2%)	0	100	100
24	22	68/72 (94%)	67 (98%)	1 (2%)	0	100	100
25	13	57/60 (95%)	55 (96%)	2 (4%)	0	100	100
25	23	57/60 (95%)	55 (96%)	2 (4%)	0	100	100
26	14	67/71 (94%)	51 (76%)	8 (12%)	8 (12%)	0	1
26	24	67/71 (94%)	53 (79%)	7 (10%)	7 (10%)	1	1
27	15	57/60 (95%)	55 (96%)	2 (4%)	0	100	100
27	25	57/60 (95%)	55 (96%)	2 (4%)	0	100	100
28	16	51/54 (94%)	50 (98%)	1 (2%)	0	100	100
28	26	51/54 (94%)	50 (98%)	1 (2%)	0	100	100
29	17	46/49 (94%)	46 (100%)	0	0	100	100
29	27	46/49 (94%)	44 (96%)	1 (2%)	1 (2%)	8	31
30	18	62/65 (95%)	61 (98%)	1 (2%)	0	100	100
30	28	62/65 (95%)	61 (98%)	1 (2%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
31	19	35/37 (95%)	35 (100%)	0	0	100	100
31	29	35/37 (95%)	35 (100%)	0	0	100	100
33	1b	229/256 (90%)	195 (85%)	28 (12%)	6 (3%)	7	26
33	2b	229/256 (90%)	190 (83%)	33 (14%)	6 (3%)	7	26
34	1c	204/239 (85%)	171 (84%)	30 (15%)	3 (2%)	13	42
34	2c	204/239 (85%)	169 (83%)	29 (14%)	6 (3%)	6	23
35	1d	206/209 (99%)	187 (91%)	16 (8%)	3 (2%)	13	42
35	2d	206/209 (99%)	189 (92%)	14 (7%)	3 (2%)	13	42
36	1e	146/162 (90%)	128 (88%)	16 (11%)	2 (1%)	14	44
36	2e	146/162 (90%)	130 (89%)	13 (9%)	3 (2%)	9	32
37	1f	98/101 (97%)	93 (95%)	5 (5%)	0	100	100
37	2f	98/101 (97%)	94 (96%)	4 (4%)	0	100	100
38	1g	153/156 (98%)	139 (91%)	12 (8%)	2 (1%)	15	46
38	2g	153/156 (98%)	138 (90%)	13 (8%)	2 (1%)	15	46
39	1h	135/138 (98%)	131 (97%)	4 (3%)	0	100	100
39	2h	135/138 (98%)	129 (96%)	6 (4%)	0	100	100
40	1i	125/128 (98%)	107 (86%)	16 (13%)	2 (2%)	12	40
40	2i	125/128 (98%)	108 (86%)	16 (13%)	1 (1%)	24	60
41	1j	95/105 (90%)	83 (87%)	9 (10%)	3 (3%)	5	20
41	2j	94/105 (90%)	85 (90%)	8 (8%)	1 (1%)	17	51
42	1k	112/129 (87%)	101 (90%)	10 (9%)	1 (1%)	21	57
42	2k	112/129 (87%)	102 (91%)	9 (8%)	1 (1%)	21	57
43	1l	120/132 (91%)	113 (94%)	7 (6%)	0	100	100
43	2l	120/132 (91%)	113 (94%)	7 (6%)	0	100	100
44	1m	116/126 (92%)	103 (89%)	12 (10%)	1 (1%)	21	57
44	2m	114/126 (90%)	100 (88%)	12 (10%)	2 (2%)	11	37
45	1n	58/61 (95%)	54 (93%)	2 (3%)	2 (3%)	5	19
45	2n	58/61 (95%)	54 (93%)	3 (5%)	1 (2%)	11	38
46	1o	86/89 (97%)	83 (96%)	2 (2%)	1 (1%)	16	48
46	2o	86/89 (97%)	81 (94%)	3 (4%)	2 (2%)	8	30
47	1p	80/88 (91%)	66 (82%)	14 (18%)	0	100	100

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Mol	Chain	Analysed	Favoured	Allowed	Outliers	Percentiles	
47	2p	80/88 (91%)	69 (86%)	11 (14%)	0	100	100
48	1q	97/105 (92%)	92 (95%)	4 (4%)	1 (1%)	19	54
48	2q	97/105 (92%)	93 (96%)	4 (4%)	0	100	100
49	1r	66/88 (75%)	61 (92%)	5 (8%)	0	100	100
49	2r	66/88 (75%)	61 (92%)	5 (8%)	0	100	100
50	1s	82/93 (88%)	70 (85%)	12 (15%)	0	100	100
50	2s	81/93 (87%)	66 (82%)	15 (18%)	0	100	100
51	1t	94/106 (89%)	84 (89%)	7 (7%)	3 (3%)	5	20
51	2t	94/106 (89%)	83 (88%)	7 (7%)	4 (4%)	3	13
52	1u	21/27 (78%)	18 (86%)	3 (14%)	0	100	100
52	2u	21/27 (78%)	18 (86%)	3 (14%)	0	100	100
55	1z	11/19 (58%)	11 (100%)	0	0	100	100
55	2z	11/19 (58%)	11 (100%)	0	0	100	100
All	All	11432/12166 (94%)	10579 (92%)	731 (6%)	122 (1%)	17	51

5 of 122 Ramachandran outliers are listed below:

Mol	Chain	Res	Type
6	1G	126	ASP
7	1H	126	PRO
14	1S	60	GLY
22	10	13	GLY
26	14	45	GLY

### 5.3.2 Protein sidechains ⓘ

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

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

Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
3	1D	215/218 (99%)	197 (92%)	18 (8%)	14	37
3	2D	216/218 (99%)	198 (92%)	18 (8%)	14	38
4	1E	164/166 (99%)	147 (90%)	17 (10%)	9	26

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
4	2E	164/166 (99%)	146 (89%)	18 (11%)	8	23
5	1F	160/166 (96%)	148 (92%)	12 (8%)	17	44
5	2F	159/166 (96%)	146 (92%)	13 (8%)	14	39
6	1G	143/156 (92%)	129 (90%)	14 (10%)	10	30
6	2G	142/156 (91%)	126 (89%)	16 (11%)	7	22
7	1H	144/148 (97%)	135 (94%)	9 (6%)	22	54
7	2H	144/148 (97%)	135 (94%)	9 (6%)	22	54
8	1I	110/124 (89%)	92 (84%)	18 (16%)	3	8
8	2I	104/124 (84%)	97 (93%)	7 (7%)	20	50
9	1N	118/119 (99%)	107 (91%)	11 (9%)	11	32
9	2N	118/119 (99%)	108 (92%)	10 (8%)	13	37
10	1O	100/100 (100%)	95 (95%)	5 (5%)	30	65
10	2O	100/100 (100%)	92 (92%)	8 (8%)	15	40
11	1P	116/116 (100%)	103 (89%)	13 (11%)	7	22
11	2P	115/116 (99%)	100 (87%)	15 (13%)	5	15
12	1Q	111/111 (100%)	103 (93%)	8 (7%)	18	46
12	2Q	111/111 (100%)	102 (92%)	9 (8%)	15	39
13	1R	101/101 (100%)	88 (87%)	13 (13%)	5	16
13	2R	101/101 (100%)	87 (86%)	14 (14%)	4	12
14	1S	87/88 (99%)	80 (92%)	7 (8%)	15	40
14	2S	85/88 (97%)	73 (86%)	12 (14%)	4	12
15	1T	115/127 (91%)	104 (90%)	11 (10%)	10	31
15	2T	113/127 (89%)	103 (91%)	10 (9%)	12	35
16	1U	93/94 (99%)	82 (88%)	11 (12%)	6	19
16	2U	93/94 (99%)	86 (92%)	7 (8%)	17	44
17	1V	80/82 (98%)	66 (82%)	14 (18%)	2	7
17	2V	80/82 (98%)	69 (86%)	11 (14%)	4	13
18	1W	90/92 (98%)	84 (93%)	6 (7%)	20	50
18	2W	90/92 (98%)	83 (92%)	7 (8%)	16	41
19	1X	77/78 (99%)	75 (97%)	2 (3%)	54	85
19	2X	77/78 (99%)	74 (96%)	3 (4%)	39	75

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
20	1Y	85/91 (93%)	81 (95%)	4 (5%)	32	68
20	2Y	85/91 (93%)	79 (93%)	6 (7%)	18	47
21	1Z	159/179 (89%)	144 (91%)	15 (9%)	11	32
21	2Z	156/179 (87%)	138 (88%)	18 (12%)	7	21
22	10	60/67 (90%)	55 (92%)	5 (8%)	14	38
22	20	60/67 (90%)	58 (97%)	2 (3%)	45	80
23	11	80/83 (96%)	75 (94%)	5 (6%)	22	54
23	21	80/83 (96%)	74 (92%)	6 (8%)	17	44
24	12	65/67 (97%)	57 (88%)	8 (12%)	6	17
24	22	65/67 (97%)	60 (92%)	5 (8%)	16	42
25	13	51/52 (98%)	47 (92%)	4 (8%)	16	41
25	23	50/52 (96%)	47 (94%)	3 (6%)	24	57
26	14	60/63 (95%)	50 (83%)	10 (17%)	3	8
26	24	53/63 (84%)	45 (85%)	8 (15%)	3	11
27	15	50/52 (96%)	48 (96%)	2 (4%)	38	74
27	25	50/52 (96%)	47 (94%)	3 (6%)	24	57
28	16	51/52 (98%)	49 (96%)	2 (4%)	39	75
28	26	50/52 (96%)	48 (96%)	2 (4%)	38	74
29	17	41/42 (98%)	38 (93%)	3 (7%)	17	45
29	27	41/42 (98%)	40 (98%)	1 (2%)	57	86
30	18	54/55 (98%)	48 (89%)	6 (11%)	8	22
30	28	54/55 (98%)	51 (94%)	3 (6%)	26	60
31	19	34/34 (100%)	34 (100%)	0	100	100
31	29	34/34 (100%)	33 (97%)	1 (3%)	50	83
33	1b	177/220 (80%)	148 (84%)	29 (16%)	3	8
33	2b	158/220 (72%)	130 (82%)	28 (18%)	2	7
34	1c	127/188 (68%)	116 (91%)	11 (9%)	13	36
34	2c	108/188 (57%)	92 (85%)	16 (15%)	4	11
35	1d	161/181 (89%)	143 (89%)	18 (11%)	7	22
35	2d	164/181 (91%)	146 (89%)	18 (11%)	8	23
36	1e	113/123 (92%)	101 (89%)	12 (11%)	8	25

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
36	2e	106/123 (86%)	95 (90%)	11 (10%)	9	26
37	1f	83/90 (92%)	78 (94%)	5 (6%)	24	57
37	2f	86/90 (96%)	78 (91%)	8 (9%)	11	32
38	1g	111/127 (87%)	94 (85%)	17 (15%)	3	10
38	2g	107/127 (84%)	97 (91%)	10 (9%)	11	32
39	1h	114/119 (96%)	103 (90%)	11 (10%)	10	31
39	2h	111/119 (93%)	95 (86%)	16 (14%)	4	12
40	1i	89/99 (90%)	78 (88%)	11 (12%)	6	17
40	2i	80/99 (81%)	69 (86%)	11 (14%)	4	13
41	1j	60/92 (65%)	53 (88%)	7 (12%)	7	19
41	2j	62/92 (67%)	49 (79%)	13 (21%)	1	4
42	1k	82/99 (83%)	78 (95%)	4 (5%)	31	67
42	2k	82/99 (83%)	77 (94%)	5 (6%)	23	56
43	1l	95/109 (87%)	87 (92%)	8 (8%)	14	37
43	2l	94/109 (86%)	86 (92%)	8 (8%)	13	37
44	1m	90/101 (89%)	81 (90%)	9 (10%)	9	28
44	2m	87/101 (86%)	79 (91%)	8 (9%)	11	33
45	1n	47/50 (94%)	42 (89%)	5 (11%)	8	25
45	2n	43/50 (86%)	33 (77%)	10 (23%)	1	3
46	1o	75/80 (94%)	65 (87%)	10 (13%)	5	14
46	2o	78/80 (98%)	73 (94%)	5 (6%)	22	53
47	1p	67/74 (90%)	58 (87%)	9 (13%)	5	13
47	2p	68/74 (92%)	61 (90%)	7 (10%)	9	26
48	1q	91/97 (94%)	87 (96%)	4 (4%)	35	70
48	2q	94/97 (97%)	92 (98%)	2 (2%)	61	88
49	1r	59/77 (77%)	51 (86%)	8 (14%)	5	13
49	2r	59/77 (77%)	52 (88%)	7 (12%)	6	19
50	1s	65/80 (81%)	55 (85%)	10 (15%)	3	10
50	2s	67/80 (84%)	61 (91%)	6 (9%)	12	34
51	1t	66/82 (80%)	59 (89%)	7 (11%)	8	25
51	2t	71/82 (87%)	68 (96%)	3 (4%)	36	73

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Mol	Chain	Analysed	Rotameric	Outliers	Percentiles	
52	1u	16/22 (73%)	16 (100%)	0	100	100
52	2u	18/22 (82%)	18 (100%)	0	100	100
55	1z	13/19 (68%)	13 (100%)	0	100	100
55	2z	13/19 (68%)	13 (100%)	0	100	100
All	All	9161/10104 (91%)	8276 (90%)	885 (10%)	10	30

5 of 885 residues with a non-rotameric sidechain are listed below:

Mol	Chain	Res	Type
46	1o	3	ILE
5	2F	197	ASP
41	2j	33	GLN
47	1p	5	ARG
3	2D	61	LEU

Some sidechains can be flipped to improve hydrogen bonding and reduce clashes. 5 of 128 such sidechains are listed below:

Mol	Chain	Res	Type
3	2D	253	GLN
15	2T	58	ASN
44	2m	77	ASN
5	2F	69	HIS
8	2I	43	ASN

### 5.3.3 RNA ⓘ

Mol	Chain	Analysed	Backbone Outliers	Pucker Outliers
1	1A	2737/2915 (93%)	357 (13%)	17 (0%)
1	2A	2781/2915 (95%)	405 (14%)	26 (0%)
2	1B	119/121 (98%)	13 (10%)	2 (1%)
2	2B	119/121 (98%)	23 (19%)	2 (1%)
32	1a	1472/1521 (96%)	238 (16%)	0
32	2a	1479/1521 (97%)	240 (16%)	0
53	1v	4/24 (16%)	2 (50%)	0
53	2v	4/24 (16%)	2 (50%)	0
54	1x	75/77 (97%)	26 (34%)	0
54	2x	75/77 (97%)	26 (34%)	0
All	All	8865/9316 (95%)	1332 (15%)	47 (0%)



5 of 1332 RNA backbone outliers are listed below:

Mol	Chain	Res	Type
1	1A	11	U
1	1A	12	A
1	1A	14	G
1	1A	33	C
1	1A	44	C

5 of 47 RNA pucker outliers are listed below:

Mol	Chain	Res	Type
1	2A	424	G
1	2A	1047	G
1	2A	2700	U
1	2A	669	C
1	2A	1087	G

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

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	5MC	1x	32	54	13,22,23	1.38	1 (7%)	15,32,35	0.99	1 (6%)
54	5MU	1x	54	54	12,22,23	0.27	0	14,32,35	2.50	2 (14%)
54	PSU	1x	55	54	13,21,22	1.33	2 (15%)	18,30,33	3.44	5 (27%)
54	4SU	1x	8	54	11,21,22	1.14	1 (9%)	13,30,33	1.68	1 (7%)
54	5MC	2x	32	54	13,22,23	1.31	1 (7%)	15,32,35	0.98	1 (6%)
54	5MU	2x	54	54	12,22,23	0.32	0	14,32,35	2.17	2 (14%)
54	PSU	2x	55	54	13,21,22	1.17	2 (15%)	18,30,33	3.39	5 (27%)
54	4SU	2x	8	54	11,21,22	1.13	1 (9%)	13,30,33	1.68	1 (7%)

In the following table, the Chirals column lists the number of chiral outliers, the number of chiral centers analysed, the number of these observed in the model and the number defined in the chemical



component dictionary. Similar counts are reported in the Torsion and Rings columns. '-' means no outliers of that kind were identified.

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
54	5MC	1x	32	54	-	0/3/25/26	0/2/2/2
54	5MU	1x	54	54	-	0/3/25/26	0/2/2/2
54	PSU	1x	55	54	-	0/7/25/26	0/2/2/2
54	4SU	1x	8	54	-	0/3/25/26	0/2/2/2
54	5MC	2x	32	54	-	0/3/25/26	0/2/2/2
54	5MU	2x	54	54	-	0/3/25/26	0/2/2/2
54	PSU	2x	55	54	-	0/7/25/26	0/2/2/2
54	4SU	2x	8	54	-	0/3/25/26	0/2/2/2

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

Mol	Chain	Res	Type	Atoms	Z	Observed(Å)	Ideal(Å)
54	1x	55	PSU	C5-C1'	-4.12	1.48	1.52
54	1x	8	4SU	C4-S4	-3.58	1.60	1.67
54	2x	8	4SU	C4-S4	-3.50	1.60	1.67
54	2x	55	PSU	C5-C1'	-3.43	1.49	1.52
54	1x	55	PSU	O4'-C1'	-2.13	1.41	1.44

The worst 5 of 18 bond angle outliers are listed below:

Mol	Chain	Res	Type	Atoms	Z	Observed(°)	Ideal(°)
54	1x	55	PSU	N1-C2-N3	-10.99	121.32	128.33
54	2x	55	PSU	N1-C2-N3	-10.87	121.40	128.33
54	1x	8	4SU	C5-C4-N3	-5.75	118.00	123.63
54	1x	54	5MU	C5-C4-N3	-5.68	118.81	125.14
54	2x	8	4SU	C5-C4-N3	-5.67	118.08	123.63

There are no chirality outliers.

There are no torsion outliers.

There are no ring outliers.

No monomer is involved in short contacts.

## 5.5 Carbohydrates ⓘ

There are no carbohydrates in this entry.



## 5.6 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	1d	501	35	0,12,12	0.00	-	0,24,24	0.00	-
58	SF4	2d	501	35	0,12,12	0.00	-	0,24,24	0.00	-

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

Mol	Type	Chain	Res	Link	Chirals	Torsions	Rings
58	SF4	1d	501	35	-	0/0/48/48	0/6/5/5
58	SF4	2d	501	35	-	0/0/48/48	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.

## 5.7 Other polymers [i](#)

There are no such residues in this entry.

## 5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.



## 6 Fit of model and data ⓘ

### 6.1 Protein, DNA and RNA chains ⓘ

In the following table, the column labelled ‘#RSRZ> 2’ contains the number (and percentage) of RSRZ outliers, followed by percent RSRZ outliers for the chain as percentile scores relative to all X-ray entries and entries of similar resolution. The OWAB column contains the minimum, median, 95<sup>th</sup> 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 <sup>2</sup> )	Q<0.9
1	1A	2746/2915 (94%)	0.66	7 (0%) 94 94	24, 41, 87, 115	0
1	2A	2790/2915 (95%)	0.25	12 (0%) 93 92	26, 45, 94, 113	0
2	1B	120/121 (99%)	0.49	0 100 100	40, 66, 79, 96	0
2	2B	120/121 (99%)	-0.10	0 100 100	45, 72, 82, 98	0
3	1D	275/276 (99%)	0.56	2 (0%) 89 88	21, 40, 58, 81	0
3	2D	275/276 (99%)	0.54	13 (4%) 35 29	21, 42, 60, 81	0
4	1E	204/206 (99%)	0.36	1 (0%) 91 90	17, 37, 64, 79	0
4	2E	204/206 (99%)	0.41	4 (1%) 68 64	26, 52, 72, 83	0
5	1F	203/210 (96%)	0.39	1 (0%) 91 90	22, 50, 77, 92	0
5	2F	203/210 (96%)	0.13	4 (1%) 68 64	25, 54, 78, 92	0
6	1G	181/182 (99%)	0.21	2 (1%) 82 80	54, 71, 84, 98	0
6	2G	181/182 (99%)	0.85	28 (15%) 3 1	59, 76, 88, 99	0
7	1H	174/180 (96%)	0.37	1 (0%) 90 89	49, 64, 79, 86	0
7	2H	174/180 (96%)	0.51	21 (12%) 6 3	52, 70, 82, 88	0
8	1I	146/148 (98%)	0.13	2 (1%) 78 76	47, 75, 87, 93	0
8	2I	146/148 (98%)	0.30	8 (5%) 29 22	48, 77, 87, 93	0
9	1N	140/140 (100%)	0.60	0 100 100	30, 46, 70, 78	0
9	2N	140/140 (100%)	0.28	0 100 100	33, 50, 72, 80	0
10	1O	122/122 (100%)	0.53	2 (1%) 74 72	31, 42, 64, 70	0
10	2O	122/122 (100%)	0.24	1 (0%) 87 86	34, 45, 64, 71	0
11	1P	149/150 (99%)	0.53	2 (1%) 79 78	25, 53, 75, 85	0
11	2P	149/150 (99%)	0.73	12 (8%) 15 9	29, 57, 77, 87	0
12	1Q	141/141 (100%)	0.57	3 (2%) 67 62	32, 50, 65, 84	0
12	2Q	141/141 (100%)	0.32	4 (2%) 56 50	36, 53, 68, 84	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å <sup>2</sup> )	Q<0.9
13	1R	118/118 (100%)	0.51	0 100 100	27, 40, 56, 67	0
13	2R	118/118 (100%)	0.47	4 (3%) 49 41	29, 43, 57, 69	0
14	1S	110/112 (98%)	0.29	0 100 100	49, 63, 74, 82	0
14	2S	110/112 (98%)	0.09	3 (2%) 58 52	54, 67, 78, 84	0
15	1T	131/146 (89%)	0.38	0 100 100	35, 48, 80, 88	0
15	2T	131/146 (89%)	0.23	1 (0%) 87 86	37, 50, 78, 87	0
16	1U	116/118 (98%)	0.60	0 100 100	29, 39, 57, 78	0
16	2U	116/118 (98%)	0.23	0 100 100	31, 44, 61, 78	0
17	1V	101/101 (100%)	0.43	0 100 100	30, 51, 69, 78	0
17	2V	101/101 (100%)	-0.07	3 (2%) 54 47	33, 57, 72, 79	0
18	1W	112/113 (99%)	0.52	2 (1%) 71 68	28, 37, 59, 90	0
18	2W	112/113 (99%)	0.25	2 (1%) 71 68	30, 40, 60, 90	0
19	1X	95/96 (98%)	0.55	1 (1%) 82 80	31, 47, 68, 80	0
19	2X	95/96 (98%)	0.22	3 (3%) 51 43	35, 50, 71, 81	0
20	1Y	107/110 (97%)	0.18	1 (0%) 85 84	31, 49, 74, 84	0
20	2Y	107/110 (97%)	0.73	6 (5%) 28 21	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%)	0.27	6 (3%) 51 43	67, 81, 94, 101	0
22	10	75/85 (88%)	0.33	0 100 100	23, 38, 53, 67	0
22	20	75/85 (88%)	1.18	11 (14%) 3 2	43, 63, 73, 78	0
23	11	97/98 (98%)	0.59	2 (2%) 67 62	21, 42, 71, 80	0
23	21	97/98 (98%)	0.52	6 (6%) 24 17	33, 54, 77, 86	0
24	12	70/72 (97%)	0.53	3 (4%) 39 32	42, 59, 74, 89	0
24	22	70/72 (97%)	-0.02	1 (1%) 78 76	46, 63, 75, 87	0
25	13	59/60 (98%)	0.45	0 100 100	32, 46, 67, 80	0
25	23	59/60 (98%)	0.39	0 100 100	36, 49, 72, 82	0
26	14	69/71 (97%)	0.32	7 (10%) 9 5	68, 84, 100, 103	0
26	24	69/71 (97%)	0.78	11 (15%) 3 1	73, 86, 100, 104	0
27	15	59/60 (98%)	0.54	1 (1%) 73 70	21, 40, 60, 78	0
27	25	59/60 (98%)	0.19	0 100 100	24, 43, 62, 79	0
28	16	53/54 (98%)	0.24	0 100 100	36, 48, 63, 68	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å <sup>2</sup> )	Q<0.9
28	26	53/54 (98%)	0.06	1 (1%) 70 66	39, 51, 65, 70	0
29	17	48/49 (97%)	1.22	6 (12%) 5 3	26, 31, 67, 76	0
29	27	48/49 (97%)	1.11	7 (14%) 3 2	28, 33, 67, 78	0
30	18	64/65 (98%)	0.72	0 100 100	32, 40, 51, 64	0
30	28	64/65 (98%)	1.18	10 (15%) 3 1	34, 43, 54, 67	0
31	19	37/37 (100%)	0.64	0 100 100	40, 49, 66, 69	0
31	29	37/37 (100%)	0.79	2 (5%) 29 23	44, 53, 67, 71	0
32	1a	1477/1521 (97%)	0.32	16 (1%) 82 80	33, 72, 99, 116	0
32	2a	1483/1521 (97%)	0.35	20 (1%) 79 78	43, 78, 102, 119	0
33	1b	231/256 (90%)	0.12	7 (3%) 54 47	60, 83, 95, 105	0
33	2b	231/256 (90%)	0.79	30 (12%) 5 3	71, 88, 97, 101	0
34	1c	206/239 (86%)	0.84	35 (16%) 2 1	70, 84, 91, 99	0
34	2c	206/239 (86%)	0.82	42 (20%) 1 1	72, 86, 93, 100	0
35	1d	208/209 (99%)	1.28	57 (27%) 1 0	55, 78, 91, 101	0
35	2d	208/209 (99%)	0.66	22 (10%) 8 5	58, 75, 87, 90	0
36	1e	148/162 (91%)	0.60	8 (5%) 29 23	47, 65, 77, 94	0
36	2e	148/162 (91%)	0.59	10 (6%) 20 14	56, 74, 86, 91	0
37	1f	100/101 (99%)	0.37	5 (5%) 32 26	57, 75, 85, 89	0
37	2f	100/101 (99%)	-0.20	0 100 100	60, 76, 86, 90	0
38	1g	155/156 (99%)	0.55	14 (9%) 12 7	66, 80, 93, 101	0
38	2g	155/156 (99%)	0.42	15 (9%) 10 6	69, 81, 94, 99	0
39	1h	137/138 (99%)	0.72	17 (12%) 5 3	56, 68, 80, 85	0
39	2h	137/138 (99%)	0.62	13 (9%) 10 6	58, 75, 83, 89	0
40	1i	127/128 (99%)	1.45	40 (31%) 1 0	66, 87, 94, 99	0
40	2i	127/128 (99%)	2.37	66 (51%) 0 0	65, 88, 96, 100	0
41	1j	97/105 (92%)	1.17	26 (26%) 1 0	65, 86, 96, 104	0
41	2j	96/105 (91%)	1.55	31 (32%) 1 0	75, 92, 99, 103	0
42	1k	114/129 (88%)	0.41	4 (3%) 48 40	45, 69, 82, 94	0
42	2k	114/129 (88%)	0.38	8 (7%) 19 13	52, 76, 88, 91	0
43	1l	122/132 (92%)	0.54	12 (9%) 10 6	47, 65, 76, 88	0
43	2l	122/132 (92%)	0.88	15 (12%) 5 3	52, 67, 77, 85	0

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Mol	Chain	Analysed	<RSRZ>	#RSRZ>2	OWAB(Å <sup>2</sup> )	Q<0.9
44	1m	118/126 (93%)	0.45	7 (5%) 26 19	64, 82, 89, 95	0
44	2m	116/126 (92%)	0.53	15 (12%) 5 3	66, 84, 91, 94	0
45	1n	60/61 (98%)	2.37	39 (65%) 0 0	69, 81, 88, 93	0
45	2n	60/61 (98%)	2.77	39 (65%) 0 0	71, 83, 89, 95	0
46	1o	88/89 (98%)	0.28	2 (2%) 64 59	49, 68, 84, 91	0
46	2o	88/89 (98%)	0.29	4 (4%) 37 31	56, 73, 85, 94	0
47	1p	82/88 (93%)	1.18	18 (21%) 1 0	66, 79, 92, 96	0
47	2p	82/88 (93%)	0.76	6 (7%) 18 12	59, 71, 83, 96	0
48	1q	99/105 (94%)	0.60	9 (9%) 11 7	51, 68, 81, 84	0
48	2q	99/105 (94%)	1.10	19 (19%) 2 1	57, 72, 83, 86	0
49	1r	68/88 (77%)	0.40	2 (2%) 55 49	56, 71, 84, 91	0
49	2r	68/88 (77%)	0.70	6 (8%) 12 8	65, 75, 89, 94	0
50	1s	84/93 (90%)	0.01	0 100 100	71, 83, 92, 98	0
50	2s	83/93 (89%)	1.41	23 (27%) 1 0	80, 92, 101, 108	0
51	1t	96/106 (90%)	0.92	19 (19%) 1 1	61, 74, 88, 90	0
51	2t	96/106 (90%)	0.41	3 (3%) 52 45	54, 73, 85, 90	0
52	1u	23/27 (85%)	1.16	4 (17%) 2 1	71, 80, 85, 87	0
52	2u	23/27 (85%)	2.49	13 (56%) 0 0	72, 82, 87, 89	0
53	1v	5/24 (20%)	1.13	1 (20%) 1 1	57, 67, 94, 101	0
53	2v	5/24 (20%)	2.17	2 (40%) 0 0	61, 70, 95, 102	0
54	1x	72/77 (93%)	0.27	0 100 100	35, 68, 87, 102	0
54	2x	72/77 (93%)	0.24	0 100 100	38, 72, 89, 103	0
55	1z	13/19 (68%)	0.96	2 (15%) 3 1	26, 37, 63, 72	0
55	2z	13/19 (68%)	1.78	3 (23%) 1 0	28, 39, 65, 74	0
All	All	20522/21482 (95%)	0.50	979 (4%) 34 28	17, 62, 93, 119	0

The worst 5 of 979 RSRZ outliers are listed below:

Mol	Chain	Res	Type	RSRZ
29	17	48	LYS	9.8
40	2i	7	THR	8.1
52	2u	14	TRP	7.9
26	24	49	PHE	7.6
38	1g	79	ARG	7.6



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

In the following table, the Atoms column lists the number of modelled atoms in the group and the number defined in the chemical component dictionary. LLDF column lists the quality of electron density of the group with respect to its neighbouring residues in protein, DNA or RNA chains. The B-factors column lists the minimum, median, 95<sup>th</sup> 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	LLDF	B-factors(Å <sup>2</sup> )	Q<0.9
54	PSU	2x	55	20/21	0.86	0.12	-	69,77,85,97	0
54	5MU	2x	54	21/22	0.94	0.16	-	75,81,92,101	0
54	5MC	2x	32	21/22	0.96	0.22	-	61,77,79,91	0
54	4SU	1x	8	20/21	0.95	0.20	-	51,69,76,81	0
54	4SU	2x	8	20/21	0.92	0.15	-	70,77,94,99	0
54	5MC	1x	32	21/22	0.97	0.19	-	48,59,69,81	0
54	5MU	1x	54	21/22	0.94	0.17	-	57,66,80,87	0
54	PSU	1x	55	20/21	0.92	0.17	-	62,73,83,85	0

## 6.3 Carbohydrates [i](#)

There are no carbohydrates in this entry.

## 6.4 Ligands [i](#)

In the following table, the Atoms column lists the number of modelled atoms in the group and the number defined in the chemical component dictionary. LLDF column lists the quality of electron density of the group with respect to its neighbouring residues in protein, DNA or RNA chains. The B-factors column lists the minimum, median, 95<sup>th</sup> 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	LLDF	B-factors(Å <sup>2</sup> )	Q<0.9
56	MG	1A	3481	1/1	0.93	0.89	57.89	40,40,40,40	0
56	MG	1A	3930	1/1	0.95	0.72	49.16	44,44,44,44	0
56	MG	1A	3374	1/1	0.89	0.79	38.56	64,64,64,64	0
56	MG	1A	3267	1/1	0.98	0.72	38.07	27,27,27,27	0
56	MG	1A	3289	1/1	0.94	0.85	34.56	43,43,43,43	0
56	MG	1A	3074	1/1	0.97	0.71	33.30	51,51,51,51	0
56	MG	1A	3092	1/1	0.89	0.60	31.90	49,49,49,49	0
56	MG	1A	4008	1/1	0.89	1.45	31.40	48,48,48,48	0
56	MG	2A	3554	1/1	0.77	2.20	29.91	51,51,51,51	0
56	MG	1U	205	1/1	0.89	1.04	28.87	52,52,52,52	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	2A	3042	1/1	0.96	0.53	28.41	43,43,43,43	0
56	MG	1A	3962	1/1	0.91	1.01	27.70	52,52,52,52	0
56	MG	1E	303	1/1	0.94	0.99	27.39	58,58,58,58	0
56	MG	2A	3107	1/1	0.92	0.75	25.71	45,45,45,45	0
56	MG	1A	3001	1/1	0.97	0.85	25.68	42,42,42,42	0
56	MG	2a	3022	1/1	0.85	0.70	25.01	73,73,73,73	0
56	MG	1A	3473	1/1	0.94	0.87	24.99	50,50,50,50	0
56	MG	1A	3986	1/1	0.94	0.85	24.95	42,42,42,42	0
56	MG	1A	3178	1/1	0.89	0.71	24.78	41,41,41,41	0
56	MG	1A	3975	1/1	0.96	0.91	24.72	45,45,45,45	0
56	MG	1U	202	1/1	0.90	0.85	24.56	50,50,50,50	0
56	MG	1A	4007	1/1	0.90	0.90	23.77	36,36,36,36	0
56	MG	1F	306	1/1	0.91	0.77	22.72	33,33,33,33	0
56	MG	1A	3187	1/1	0.91	0.64	22.66	46,46,46,46	0
56	MG	1A	3132	1/1	0.68	0.72	22.56	51,51,51,51	0
56	MG	1A	3456	1/1	0.97	0.53	21.83	39,39,39,39	0
56	MG	1D	309	1/1	0.97	0.82	21.81	41,41,41,41	0
56	MG	1A	3093	1/1	0.95	0.52	20.89	47,47,47,47	0
56	MG	1A	4003	1/1	0.94	0.89	20.72	43,43,43,43	0
56	MG	1A	4018	1/1	0.85	0.93	19.89	50,50,50,50	0
56	MG	1A	4017	1/1	0.81	0.93	19.88	60,60,60,60	0
56	MG	2A	3214	1/1	0.90	0.78	18.94	43,43,43,43	0
56	MG	1W	3005	1/1	0.91	0.53	18.81	49,49,49,49	0
56	MG	1A	3110	1/1	0.93	0.70	17.31	55,55,55,55	0
56	MG	1U	204	1/1	0.89	1.10	17.28	42,42,42,42	0
56	MG	1A	4011	1/1	0.95	0.95	17.13	47,47,47,47	0
56	MG	1A	4002	1/1	0.94	0.89	17.09	34,34,34,34	0
56	MG	1A	3322	1/1	0.81	0.62	15.93	50,50,50,50	0
56	MG	2A	3063	1/1	0.95	0.46	15.72	45,45,45,45	0
56	MG	1A	3394	1/1	0.67	0.46	15.30	56,56,56,56	0
56	MG	1A	4010	1/1	0.87	0.57	15.25	41,41,41,41	0
56	MG	1A	3995	1/1	0.82	0.89	14.63	42,42,42,42	0
56	MG	1A	3182	1/1	0.97	0.47	14.25	42,42,42,42	0
56	MG	1a	1765	1/1	0.78	0.51	14.02	57,57,57,57	0
56	MG	1A	3179	1/1	0.94	0.40	13.95	44,44,44,44	0
56	MG	1A	3931	1/1	0.91	0.45	13.72	33,33,33,33	0
56	MG	1A	3241	1/1	0.89	0.39	13.46	42,42,42,42	0
56	MG	2U	201	1/1	0.96	0.69	13.20	48,48,48,48	0
56	MG	1A	3129	1/1	0.89	0.67	12.98	40,40,40,40	0
56	MG	1A	3994	1/1	0.97	0.62	12.00	47,47,47,47	0
56	MG	1P	206	1/1	0.97	0.76	11.97	54,54,54,54	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3131	1/1	0.86	0.57	11.92	42,42,42,42	0
56	MG	1A	3315	1/1	0.90	0.46	11.67	49,49,49,49	0
56	MG	2A	3027	1/1	0.74	0.89	11.60	51,51,51,51	0
56	MG	2A	3563	1/1	0.94	0.70	11.17	47,47,47,47	0
56	MG	2A	3562	1/1	0.77	0.65	11.15	49,49,49,49	0
56	MG	1F	304	1/1	0.88	0.55	11.12	43,43,43,43	0
56	MG	2A	3139	1/1	0.94	0.31	11.06	45,45,45,45	0
56	MG	1A	3442	1/1	0.73	0.38	10.87	58,58,58,58	0
56	MG	1A	3980	1/1	0.86	0.44	10.80	50,50,50,50	0
56	MG	1A	3186	1/1	0.94	0.43	10.46	32,32,32,32	0
56	MG	2A	3016	1/1	0.93	0.26	10.14	57,57,57,57	0
56	MG	1R	203	1/1	0.91	0.63	9.96	43,43,43,43	0
56	MG	1A	3177	1/1	0.93	0.62	9.90	39,39,39,39	0
56	MG	1A	3444	1/1	0.91	0.59	9.74	36,36,36,36	0
56	MG	1A	3797	1/1	0.96	0.80	9.72	40,40,40,40	0
56	MG	1D	311	1/1	0.87	0.47	9.67	55,55,55,55	0
56	MG	1a	1749	1/1	0.96	0.88	9.66	57,57,57,57	0
56	MG	1A	3215	1/1	0.95	0.23	9.63	51,51,51,51	0
56	MG	1A	4019	1/1	0.94	0.78	9.28	47,47,47,47	0
56	MG	1A	3968	1/1	0.91	0.62	9.12	50,50,50,50	0
56	MG	1A	3203	1/1	0.98	0.28	9.05	32,32,32,32	0
56	MG	1A	3449	1/1	0.93	0.31	8.92	66,66,66,66	0
56	MG	1A	3973	1/1	0.90	0.45	8.74	51,51,51,51	0
56	MG	2A	3222	1/1	0.94	0.61	8.74	36,36,36,36	0
56	MG	2A	3561	1/1	0.96	0.93	8.65	41,41,41,41	0
56	MG	2A	3349	1/1	0.95	0.24	8.65	59,59,59,59	0
56	MG	1D	306	1/1	0.97	0.49	8.52	31,31,31,31	0
56	MG	2a	3228	1/1	0.84	0.40	8.45	69,69,69,69	0
56	MG	2A	3233	1/1	0.91	0.54	8.29	48,48,48,48	0
56	MG	15	101	1/1	0.66	0.49	8.29	58,58,58,58	0
56	MG	1X	102	1/1	0.91	0.47	7.70	53,53,53,53	0
56	MG	1U	203	1/1	0.93	0.51	7.52	47,47,47,47	0
56	MG	1A	3887	1/1	0.92	0.34	7.19	31,31,31,31	0
56	MG	1A	3225	1/1	0.99	0.55	7.11	34,34,34,34	0
56	MG	1a	1668	1/1	0.89	0.30	6.95	59,59,59,59	0
56	MG	1a	1659	1/1	0.92	0.44	6.82	63,63,63,63	0
56	MG	2A	3416	1/1	0.90	0.19	6.73	48,48,48,48	0
56	MG	2A	3430	1/1	0.92	0.30	6.49	45,45,45,45	0
56	MG	1D	303	1/1	0.89	0.43	6.44	47,47,47,47	0
56	MG	1A	3518	1/1	0.90	0.36	6.29	53,53,53,53	0
56	MG	1A	3411	1/1	0.94	0.30	6.25	62,62,62,62	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1D	312	1/1	0.98	0.44	6.23	47,47,47,47	0
56	MG	2F	303	1/1	0.89	0.62	6.11	50,50,50,50	0
56	MG	2a	3011	1/1	0.80	0.22	6.07	67,67,67,67	0
56	MG	1A	3287	1/1	0.93	0.35	6.01	27,27,27,27	0
56	MG	2A	3540	1/1	0.77	0.33	5.89	58,58,58,58	0
56	MG	1A	3705	1/1	0.84	0.40	5.82	34,34,34,34	0
56	MG	1a	1858	1/1	0.91	0.26	5.67	52,52,52,52	0
56	MG	1A	3208	1/1	0.97	0.29	5.34	34,34,34,34	0
56	MG	2A	3024	1/1	0.90	0.26	5.33	43,43,43,43	0
56	MG	1A	3162	1/1	0.91	0.41	5.24	38,38,38,38	0
56	MG	1A	3459	1/1	0.83	0.43	5.15	46,46,46,46	0
56	MG	1P	205	1/1	0.96	0.53	5.14	31,31,31,31	0
56	MG	2P	201	1/1	0.96	0.57	5.02	41,41,41,41	0
56	MG	1A	4012	1/1	0.97	0.45	4.88	36,36,36,36	0
56	MG	1H	8002	1/1	0.80	0.42	4.71	79,79,79,79	0
56	MG	1A	3711	1/1	0.94	0.34	4.67	48,48,48,48	0
56	MG	1A	3172	1/1	0.85	0.63	4.66	44,44,44,44	0
56	MG	2A	3029	1/1	0.92	0.24	4.36	48,48,48,48	0
56	MG	1a	1814	1/1	0.95	0.30	4.34	67,67,67,67	0
56	MG	2A	3381	1/1	0.63	0.20	4.06	64,64,64,64	0
56	MG	2A	3550	1/1	0.92	0.50	4.01	42,42,42,42	0
56	MG	2A	3019	1/1	0.98	0.30	4.01	46,46,46,46	0
56	MG	2a	3105	1/1	0.94	0.33	3.99	55,55,55,55	0
56	MG	1A	3253	1/1	0.98	0.26	3.81	31,31,31,31	0
56	MG	1A	3888	1/1	0.86	0.48	3.72	42,42,42,42	0
56	MG	1A	3329	1/1	0.96	0.26	3.68	52,52,52,52	0
56	MG	2A	3304	1/1	0.97	0.28	3.52	41,41,41,41	0
56	MG	2A	3555	1/1	0.81	0.52	3.34	36,36,36,36	0
56	MG	2A	3199	1/1	0.96	0.23	3.32	39,39,39,39	0
56	MG	1A	3821	1/1	0.92	0.29	3.18	23,23,23,23	0
56	MG	1A	3437	1/1	0.92	0.25	3.16	62,62,62,62	0
56	MG	1A	3312	1/1	0.95	0.26	3.10	38,38,38,38	0
56	MG	2A	3470	1/1	0.98	0.27	3.09	44,44,44,44	0
56	MG	2A	3191	1/1	0.90	0.29	3.00	40,40,40,40	0
56	MG	2A	3123	1/1	0.99	0.19	2.92	41,41,41,41	0
56	MG	1B	211	1/1	0.89	0.23	2.87	51,51,51,51	0
56	MG	2a	3194	1/1	0.97	0.21	2.82	72,72,72,72	0
56	MG	2A	3155	1/1	0.96	0.22	2.76	33,33,33,33	0
56	MG	2D	303	1/1	0.92	0.31	2.74	49,49,49,49	0
56	MG	2A	3232	1/1	0.83	0.24	2.72	54,54,54,54	0
56	MG	1a	1739	1/1	0.94	0.29	2.69	46,46,46,46	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3964	1/1	0.96	0.30	2.65	29,29,29,29	0
56	MG	2A	3559	1/1	0.86	0.31	2.60	54,54,54,54	0
56	MG	2A	3303	1/1	0.98	0.23	2.50	55,55,55,55	0
56	MG	1A	3911	1/1	0.94	0.27	2.47	39,39,39,39	0
56	MG	2a	3026	1/1	0.83	0.26	2.37	64,64,64,64	0
56	MG	2A	3017	1/1	0.89	0.36	2.23	47,47,47,47	0
56	MG	2A	3209	1/1	0.79	0.19	2.22	45,45,45,45	0
56	MG	2A	3468	1/1	0.91	0.20	2.19	56,56,56,56	0
56	MG	1A	3170	1/1	0.83	0.23	2.11	44,44,44,44	0
56	MG	2A	3318	1/1	0.93	0.25	2.11	47,47,47,47	0
56	MG	1B	205	1/1	0.86	0.22	2.07	56,56,56,56	0
56	MG	2A	3188	1/1	0.96	0.23	2.04	35,35,35,35	0
56	MG	1a	1730	1/1	0.97	0.24	1.88	70,70,70,70	0
56	MG	1W	3003	1/1	0.95	0.28	1.86	35,35,35,35	0
56	MG	1A	3860	1/1	0.97	0.23	1.85	29,29,29,29	0
56	MG	2A	3556	1/1	0.84	0.19	1.79	62,62,62,62	0
56	MG	2a	3065	1/1	0.86	0.34	1.77	80,80,80,80	0
56	MG	15	104	1/1	0.87	0.27	1.74	62,62,62,62	0
56	MG	2A	3396	1/1	0.92	0.23	1.74	40,40,40,40	0
56	MG	2A	3558	1/1	0.98	0.32	1.70	34,34,34,34	0
57	ZN	25	501	1/1	0.98	0.21	1.69	54,54,54,54	0
56	MG	1A	3408	1/1	0.81	0.24	1.63	42,42,42,42	0
56	MG	1A	3167	1/1	0.82	0.27	1.56	35,35,35,35	0
56	MG	1A	3292	1/1	0.95	0.27	1.48	17,17,17,17	0
57	ZN	26	102	1/1	0.95	0.20	1.44	56,56,56,56	0
56	MG	1A	3222	1/1	0.95	0.23	1.43	50,50,50,50	0
56	MG	2a	3030	1/1	0.42	0.26	1.41	73,73,73,73	0
56	MG	2A	3427	1/1	0.92	0.24	1.40	56,56,56,56	0
56	MG	2a	3111	1/1	0.98	0.25	1.36	56,56,56,56	0
56	MG	2a	3068	1/1	0.94	0.33	1.33	67,67,67,67	0
56	MG	19	101	1/1	0.98	0.29	1.33	65,65,65,65	0
56	MG	1A	4020	1/1	0.99	0.30	1.31	32,32,32,32	0
56	MG	2a	3027	1/1	0.84	0.28	1.27	70,70,70,70	0
56	MG	1A	3912	1/1	0.94	0.33	1.19	40,40,40,40	0
56	MG	2A	3147	1/1	0.89	0.16	1.18	42,42,42,42	0
56	MG	1x	104	1/1	0.96	0.15	1.14	52,52,52,52	0
56	MG	1A	3809	1/1	0.98	0.21	1.14	45,45,45,45	0
56	MG	1R	202	1/1	0.96	0.29	1.13	52,52,52,52	0
56	MG	1a	1710	1/1	0.88	0.18	1.07	71,71,71,71	0
56	MG	1a	1795	1/1	0.95	0.21	1.04	75,75,75,75	0
56	MG	1q	203	1/1	0.90	0.24	0.99	51,51,51,51	0
56	MG	1A	3258	1/1	0.95	0.24	0.98	36,36,36,36	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1q	202	1/1	0.91	0.26	0.95	62,62,62,62	0
56	MG	2A	3078	1/1	0.94	0.19	0.91	60,60,60,60	0
56	MG	2a	3257	1/1	0.91	0.16	0.90	90,90,90,90	0
56	MG	1A	3740	1/1	0.94	0.25	0.84	36,36,36,36	0
57	ZN	2Y	501	1/1	0.92	0.19	0.75	85,85,85,85	0
56	MG	2a	3187	1/1	0.88	0.24	0.63	59,59,59,59	0
56	MG	1a	1722	1/1	0.89	0.22	0.57	54,54,54,54	0
56	MG	1A	3813	1/1	0.94	0.21	0.54	56,56,56,56	0
56	MG	2A	3376	1/1	0.93	0.18	0.51	47,47,47,47	0
56	MG	2A	3149	1/1	0.96	0.22	0.48	43,43,43,43	0
56	MG	1A	3250	1/1	0.98	0.20	0.43	34,34,34,34	0
56	MG	2a	3028	1/1	0.99	0.23	0.36	76,76,76,76	0
56	MG	2a	3259	1/1	0.72	0.23	0.34	56,56,56,56	0
56	MG	2A	3486	1/1	0.82	0.18	0.26	57,57,57,57	0
56	MG	1a	1870	1/1	0.95	0.30	0.24	88,88,88,88	0
56	MG	1D	310	1/1	0.85	0.22	0.21	57,57,57,57	0
56	MG	1a	1810	1/1	0.95	0.20	0.19	38,38,38,38	0
56	MG	1A	3683	1/1	0.88	0.22	0.16	58,58,58,58	0
56	MG	2a	3007	1/1	0.98	0.20	0.12	46,46,46,46	0
56	MG	1A	3882	1/1	0.97	0.22	0.11	49,49,49,49	0
56	MG	2A	3326	1/1	0.72	0.21	0.11	42,42,42,42	0
56	MG	2A	3100	1/1	0.95	0.21	0.10	50,50,50,50	0
56	MG	1A	3168	1/1	0.92	0.25	0.07	42,42,42,42	0
56	MG	2A	3292	1/1	0.91	0.19	0.05	42,42,42,42	0
56	MG	2A	3539	1/1	0.95	0.22	0.01	35,35,35,35	0
56	MG	2A	3059	1/1	0.91	0.18	-0.06	48,48,48,48	0
56	MG	1a	1846	1/1	0.92	0.18	-0.06	52,52,52,52	0
56	MG	1A	3010	1/1	0.89	0.22	-0.09	42,42,42,42	0
57	ZN	1Y	201	1/1	0.97	0.19	-0.11	61,61,61,61	0
56	MG	2A	3048	1/1	0.93	0.18	-0.12	56,56,56,56	0
56	MG	2A	3507	1/1	0.94	0.17	-0.15	47,47,47,47	0
56	MG	1A	3321	1/1	0.95	0.24	-0.16	37,37,37,37	0
56	MG	2A	3186	1/1	0.92	0.15	-0.16	36,36,36,36	0
56	MG	1e	3002	1/1	0.95	0.18	-0.18	83,83,83,83	0
56	MG	1a	1673	1/1	0.90	0.16	-0.18	71,71,71,71	0
56	MG	2A	3565	1/1	0.93	0.15	-0.25	52,52,52,52	0
56	MG	1A	3548	1/1	0.97	0.20	-0.26	38,38,38,38	0
56	MG	2a	3255	1/1	0.72	0.18	-0.27	70,70,70,70	0
56	MG	2A	3405	1/1	0.84	0.18	-0.28	63,63,63,63	0
56	MG	1A	3198	1/1	0.95	0.21	-0.30	28,28,28,28	0
57	ZN	16	102	1/1	0.99	0.19	-0.30	29,29,29,29	0
56	MG	1D	308	1/1	0.89	0.20	-0.30	35,35,35,35	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	2A	3373	1/1	0.92	0.17	-0.40	47,47,47,47	0
56	MG	2a	3005	1/1	0.93	0.20	-0.40	62,62,62,62	0
56	MG	1a	1859	1/1	0.95	0.18	-0.40	59,59,59,59	0
56	MG	2a	3184	1/1	0.91	0.14	-0.41	80,80,80,80	0
56	MG	2a	3261	1/1	0.76	0.23	-0.46	74,74,74,74	0
57	ZN	19	103	1/1	0.99	0.22	-0.47	45,45,45,45	0
56	MG	2E	304	1/1	0.94	0.18	-0.48	67,67,67,67	0
56	MG	2A	3262	1/1	0.96	0.18	-0.50	54,54,54,54	0
56	MG	1A	3202	1/1	0.93	0.22	-0.55	43,43,43,43	0
56	MG	2D	302	1/1	0.87	0.19	-0.56	54,54,54,54	0
56	MG	2A	3339	1/1	0.90	0.22	-0.58	31,31,31,31	0
56	MG	1A	3613	1/1	0.84	0.19	-0.62	35,35,35,35	0
56	MG	2A	3171	1/1	0.94	0.15	-0.63	41,41,41,41	0
56	MG	2A	3368	1/1	0.83	0.21	-0.63	29,29,29,29	0
57	ZN	29	501	1/1	0.93	0.16	-0.63	62,62,62,62	0
57	ZN	14	501	1/1	0.97	0.14	-0.68	106,106,106,106	0
56	MG	1A	3645	1/1	0.82	0.21	-0.68	23,23,23,23	0
56	MG	2q	202	1/1	0.79	0.26	-0.70	69,69,69,69	0
56	MG	1A	3022	1/1	0.90	0.23	-0.70	59,59,59,59	0
56	MG	2a	3013	1/1	0.87	0.15	-0.71	72,72,72,72	0
56	MG	2A	3312	1/1	0.97	0.20	-0.72	28,28,28,28	0
56	MG	2A	3293	1/1	0.92	0.16	-0.73	62,62,62,62	0
56	MG	1a	1799	1/1	0.93	0.19	-0.73	61,61,61,61	0
56	MG	1A	3993	1/1	0.97	0.20	-0.75	73,73,73,73	0
56	MG	1A	3567	1/1	0.88	0.20	-0.79	39,39,39,39	0
56	MG	1A	3990	1/1	0.96	0.23	-0.80	36,36,36,36	0
58	SF4	1d	501	8/8	0.98	0.18	-0.81	66,73,85,85	0
56	MG	1A	3843	1/1	0.95	0.18	-0.82	50,50,50,50	0
56	MG	2A	3508	1/1	0.85	0.21	-0.83	56,56,56,56	0
56	MG	2a	3163	1/1	0.90	0.12	-0.84	70,70,70,70	0
56	MG	2A	3106	1/1	0.73	0.14	-0.84	59,59,59,59	0
56	MG	1A	3617	1/1	0.91	0.23	-0.88	27,27,27,27	0
56	MG	2l	202	1/1	0.95	0.24	-0.89	40,40,40,40	0
56	MG	2O	202	1/1	0.99	0.16	-0.90	42,42,42,42	0
56	MG	2a	3232	1/1	0.98	0.13	-0.92	66,66,66,66	0
56	MG	1A	3588	1/1	0.97	0.22	-0.92	33,33,33,33	0
56	MG	1d	502	1/1	0.85	0.13	-0.98	76,76,76,76	0
58	SF4	2d	501	8/8	0.99	0.17	-0.99	56,76,90,91	0
56	MG	1a	1790	1/1	0.94	0.17	-1.01	71,71,71,71	0
56	MG	1A	3538	1/1	0.95	0.21	-1.01	57,57,57,57	0
56	MG	1a	1711	1/1	0.96	0.19	-1.03	49,49,49,49	0
56	MG	2D	301	1/1	0.93	0.16	-1.05	57,57,57,57	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3575	1/1	0.79	0.21	-1.07	41,41,41,41	0
56	MG	1a	1837	1/1	0.99	0.16	-1.08	37,37,37,37	0
56	MG	2A	3235	1/1	0.78	0.16	-1.10	52,52,52,52	0
56	MG	1B	207	1/1	0.92	0.19	-1.11	33,33,33,33	0
56	MG	2A	3564	1/1	0.81	0.18	-1.11	59,59,59,59	0
56	MG	1a	1629	1/1	0.91	0.13	-1.14	49,49,49,49	0
56	MG	2G	3001	1/1	0.58	0.18	-1.14	58,58,58,58	0
56	MG	2E	301	1/1	0.90	0.18	-1.17	44,44,44,44	0
56	MG	1A	3188	1/1	0.93	0.21	-1.17	43,43,43,43	0
56	MG	2a	3129	1/1	0.95	0.14	-1.18	65,65,65,65	0
56	MG	2A	3533	1/1	0.89	0.22	-1.19	35,35,35,35	0
56	MG	2a	3113	1/1	0.91	0.10	-1.19	65,65,65,65	0
56	MG	2A	3300	1/1	0.84	0.17	-1.20	32,32,32,32	0
57	ZN	15	103	1/1	0.99	0.18	-1.20	43,43,43,43	0
56	MG	1A	3895	1/1	0.82	0.22	-1.23	24,24,24,24	0
57	ZN	2n	501	1/1	0.92	0.12	-1.23	98,98,98,98	0
56	MG	1A	3528	1/1	0.96	0.20	-1.23	54,54,54,54	0
56	MG	1A	3632	1/1	0.82	0.20	-1.26	25,25,25,25	0
56	MG	1A	3607	1/1	0.85	0.20	-1.27	40,40,40,40	0
56	MG	1A	3731	1/1	0.93	0.22	-1.28	23,23,23,23	0
56	MG	1A	4004	1/1	0.95	0.22	-1.30	49,49,49,49	0
56	MG	1a	1665	1/1	0.93	0.14	-1.30	61,61,61,61	0
56	MG	2A	3546	1/1	0.92	0.16	-1.31	44,44,44,44	0
57	ZN	24	501	1/1	0.77	0.10	-1.33	125,125,125,125	0
56	MG	2B	3004	1/1	0.82	0.14	-1.34	75,75,75,75	0
56	MG	1A	3637	1/1	0.94	0.18	-1.36	47,47,47,47	0
56	MG	2A	3398	1/1	0.82	0.16	-1.38	50,50,50,50	0
56	MG	2A	3220	1/1	0.94	0.13	-1.38	45,45,45,45	0
56	MG	2A	3167	1/1	0.94	0.20	-1.38	25,25,25,25	0
56	MG	1A	3578	1/1	0.97	0.17	-1.40	64,64,64,64	0
56	MG	2t	3001	1/1	0.94	0.12	-1.40	50,50,50,50	0
56	MG	2A	3261	1/1	0.94	0.18	-1.41	42,42,42,42	0
56	MG	2a	3108	1/1	0.79	0.09	-1.44	62,62,62,62	0
56	MG	1Q	3004	1/1	0.98	0.18	-1.45	39,39,39,39	0
56	MG	2a	3191	1/1	0.85	0.12	-1.46	54,54,54,54	0
56	MG	1A	3589	1/1	0.96	0.20	-1.46	29,29,29,29	0
56	MG	1A	3767	1/1	0.93	0.20	-1.46	40,40,40,40	0
56	MG	2A	3299	1/1	0.87	0.17	-1.47	46,46,46,46	0
56	MG	2a	3153	1/1	0.81	0.11	-1.51	67,67,67,67	0
56	MG	1A	3196	1/1	0.97	0.20	-1.51	25,25,25,25	0
56	MG	2A	3306	1/1	0.87	0.19	-1.52	48,48,48,48	0
56	MG	1n	502	1/1	0.83	0.24	-1.53	67,67,67,67	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	2a	3048	1/1	0.94	0.17	-1.53	80,80,80,80	0
56	MG	2A	3067	1/1	0.89	0.18	-1.57	52,52,52,52	0
56	MG	2A	3338	1/1	0.93	0.14	-1.65	56,56,56,56	0
57	ZN	1n	501	1/1	0.92	0.11	-1.66	84,84,84,84	0
56	MG	2A	3168	1/1	0.94	0.20	-1.67	47,47,47,47	0
56	MG	2A	3296	1/1	0.96	0.18	-1.69	45,45,45,45	0
56	MG	2A	3301	1/1	0.82	0.15	-1.69	38,38,38,38	0
56	MG	1a	1738	1/1	0.95	0.12	-1.69	64,64,64,64	0
56	MG	2a	3104	1/1	0.86	0.14	-1.70	62,62,62,62	0
56	MG	2A	3184	1/1	0.89	0.19	-1.70	39,39,39,39	0
56	MG	2A	3370	1/1	0.97	0.15	-1.74	44,44,44,44	0
56	MG	1a	1873	1/1	0.72	0.09	-1.76	52,52,52,52	0
56	MG	2a	3034	1/1	0.91	0.17	-1.77	59,59,59,59	0
56	MG	1a	1604	1/1	0.93	0.12	-1.77	69,69,69,69	0
56	MG	1b	3001	1/1	0.85	0.11	-1.79	83,83,83,83	0
56	MG	1A	3233	1/1	0.87	0.19	-1.80	34,34,34,34	0
56	MG	2A	3337	1/1	0.92	0.18	-1.81	46,46,46,46	0
56	MG	2A	3481	1/1	0.97	0.15	-1.82	50,50,50,50	0
56	MG	1A	3935	1/1	0.96	0.19	-1.83	28,28,28,28	0
56	MG	2A	3054	1/1	0.95	0.15	-1.83	49,49,49,49	0
56	MG	2A	3070	1/1	0.94	0.17	-1.88	32,32,32,32	0
56	MG	2Q	3003	1/1	0.95	0.14	-1.89	41,41,41,41	0
56	MG	2a	3076	1/1	0.97	0.16	-1.90	83,83,83,83	0
56	MG	1A	4015	1/1	0.90	0.20	-1.92	15,15,15,15	0
56	MG	1A	3159	1/1	0.97	0.16	-1.93	35,35,35,35	0
56	MG	1a	1664	1/1	0.96	0.12	-1.93	61,61,61,61	0
56	MG	2A	3164	1/1	0.90	0.15	-1.94	64,64,64,64	0
56	MG	2a	3099	1/1	0.92	0.16	-1.96	35,35,35,35	0
56	MG	2A	3208	1/1	0.96	0.13	-1.99	40,40,40,40	0
56	MG	1G	3001	1/1	0.87	0.17	-2.01	48,48,48,48	0
56	MG	1a	1709	1/1	0.92	0.16	-2.03	43,43,43,43	0
56	MG	1a	1815	1/1	0.99	0.10	-2.03	58,58,58,58	0
56	MG	2a	3082	1/1	0.85	0.13	-2.06	61,61,61,61	0
56	MG	2A	3401	1/1	0.69	0.13	-2.07	48,48,48,48	0
56	MG	2F	306	1/1	0.94	0.11	-2.08	40,40,40,40	0
56	MG	2A	3500	1/1	0.98	0.12	-2.14	49,49,49,49	0
56	MG	1t	3001	1/1	0.81	0.13	-2.16	54,54,54,54	0
56	MG	1A	3033	1/1	0.83	0.13	-2.17	57,57,57,57	0
56	MG	1A	3539	1/1	0.86	0.22	-2.17	44,44,44,44	0
56	MG	1r	3001	1/1	0.80	0.15	-2.18	69,69,69,69	0
56	MG	1A	3282	1/1	0.95	0.17	-2.22	52,52,52,52	0
56	MG	1A	3501	1/1	0.95	0.18	-2.23	56,56,56,56	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	10	101	1/1	0.80	0.14	-2.23	61,61,61,61	0
56	MG	2A	3133	1/1	0.94	0.12	-2.23	52,52,52,52	0
56	MG	1A	3054	1/1	0.85	0.18	-2.24	47,47,47,47	0
56	MG	2a	3131	1/1	0.96	0.17	-2.28	44,44,44,44	0
56	MG	1a	1789	1/1	0.97	0.08	-2.30	66,66,66,66	0
56	MG	1A	3819	1/1	0.93	0.10	-2.32	82,82,82,82	0
56	MG	2A	3395	1/1	0.87	0.18	-2.36	33,33,33,33	0
56	MG	1A	3150	1/1	0.86	0.15	-2.37	34,34,34,34	0
56	MG	2A	3547	1/1	0.90	0.15	-2.37	22,22,22,22	0
56	MG	1a	1872	1/1	0.92	0.10	-2.37	53,53,53,53	0
56	MG	1A	3775	1/1	0.99	0.17	-2.41	14,14,14,14	0
56	MG	2l	201	1/1	0.76	0.10	-2.42	63,63,63,63	0
56	MG	2D	305	1/1	0.98	0.14	-2.42	28,28,28,28	0
56	MG	1a	1671	1/1	0.91	0.18	-2.42	38,38,38,38	0
56	MG	1a	1822	1/1	0.80	0.15	-2.42	52,52,52,52	0
56	MG	1a	1874	1/1	0.94	0.07	-2.43	61,61,61,61	0
56	MG	2A	3173	1/1	0.88	0.12	-2.45	52,52,52,52	0
56	MG	2a	3066	1/1	0.90	0.09	-2.46	84,84,84,84	0
56	MG	1A	3734	1/1	0.87	0.21	-2.46	38,38,38,38	0
56	MG	2A	3154	1/1	0.96	0.16	-2.47	32,32,32,32	0
56	MG	2A	3553	1/1	0.91	0.12	-2.47	44,44,44,44	0
56	MG	1a	1728	1/1	0.97	0.13	-2.48	53,53,53,53	0
56	MG	2A	3196	1/1	0.98	0.17	-2.49	35,35,35,35	0
56	MG	1A	3704	1/1	0.79	0.18	-2.59	18,18,18,18	0
56	MG	1A	3612	1/1	0.89	0.16	-2.61	55,55,55,55	0
56	MG	2a	3116	1/1	0.74	0.12	-2.62	62,62,62,62	0
56	MG	2A	3297	1/1	0.92	0.11	-2.62	58,58,58,58	0
56	MG	1a	1687	1/1	0.91	0.16	-2.63	46,46,46,46	0
56	MG	1A	3385	1/1	0.77	0.20	-2.63	59,59,59,59	0
56	MG	2A	3548	1/1	0.91	0.11	-2.63	40,40,40,40	0
56	MG	28	8001	1/1	0.95	0.16	-2.64	54,54,54,54	0
56	MG	1A	4014	1/1	0.99	0.16	-2.64	43,43,43,43	0
56	MG	1a	1847	1/1	0.99	0.10	-2.66	53,53,53,53	0
56	MG	1a	1824	1/1	0.98	0.16	-2.68	52,52,52,52	0
56	MG	1A	3885	1/1	0.96	0.17	-2.69	53,53,53,53	0
56	MG	1A	3221	1/1	0.84	0.16	-2.69	37,37,37,37	0
56	MG	2a	3203	1/1	0.94	0.16	-2.70	69,69,69,69	0
56	MG	2A	3465	1/1	0.78	0.16	-2.70	31,31,31,31	0
56	MG	1A	3756	1/1	0.98	0.15	-2.70	51,51,51,51	0
56	MG	1A	3273	1/1	0.90	0.18	-2.71	36,36,36,36	0
56	MG	2a	3087	1/1	0.88	0.12	-2.71	55,55,55,55	0
56	MG	1B	221	1/1	0.79	0.17	-2.71	68,68,68,68	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	2A	3386	1/1	0.91	0.16	-2.72	30,30,30,30	0
56	MG	1A	3268	1/1	0.86	0.14	-2.72	55,55,55,55	0
56	MG	2A	3392	1/1	0.96	0.13	-2.74	28,28,28,28	0
56	MG	1A	3979	1/1	0.94	0.19	-2.74	49,49,49,49	0
56	MG	2A	3560	1/1	0.90	0.17	-2.75	31,31,31,31	0
56	MG	1A	3195	1/1	0.99	0.17	-2.76	33,33,33,33	0
56	MG	2A	3194	1/1	0.74	0.14	-2.78	44,44,44,44	0
56	MG	2A	3321	1/1	0.85	0.15	-2.78	35,35,35,35	0
56	MG	1A	3192	1/1	0.96	0.19	-2.82	35,35,35,35	0
56	MG	2A	3215	1/1	0.97	0.18	-2.83	39,39,39,39	0
56	MG	2x	3004	1/1	0.95	0.13	-2.85	42,42,42,42	0
56	MG	2A	3536	1/1	0.93	0.12	-2.89	38,38,38,38	0
56	MG	1A	3566	1/1	0.96	0.17	-2.90	23,23,23,23	0
56	MG	1A	3658	1/1	0.88	0.19	-2.91	18,18,18,18	0
56	MG	1A	3954	1/1	0.95	0.11	-2.91	51,51,51,51	0
56	MG	2a	3212	1/1	0.95	0.10	-2.92	54,54,54,54	0
56	MG	1A	3907	1/1	0.97	0.15	-2.93	15,15,15,15	0
56	MG	2A	3329	1/1	0.91	0.14	-2.93	37,37,37,37	0
56	MG	2A	3090	1/1	0.97	0.15	-2.94	45,45,45,45	0
56	MG	1a	1718	1/1	0.97	0.15	-2.95	48,48,48,48	0
56	MG	1A	3753	1/1	0.96	0.19	-2.95	68,68,68,68	0
56	MG	1A	3651	1/1	0.98	0.15	-2.95	21,21,21,21	0
56	MG	1A	3618	1/1	0.91	0.19	-2.98	18,18,18,18	0
56	MG	2F	305	1/1	0.84	0.12	-2.99	41,41,41,41	0
56	MG	1A	3551	1/1	0.95	0.20	-2.99	33,33,33,33	0
56	MG	1A	3608	1/1	0.94	0.19	-3.01	25,25,25,25	0
56	MG	2A	3112	1/1	0.85	0.13	-3.02	39,39,39,39	0
56	MG	2a	3208	1/1	0.95	0.13	-3.03	58,58,58,58	0
56	MG	1a	1677	1/1	0.74	0.12	-3.03	66,66,66,66	0
56	MG	2a	3244	1/1	0.94	0.17	-3.03	67,67,67,67	0
56	MG	1A	3750	1/1	0.91	0.15	-3.05	21,21,21,21	0
56	MG	1A	3615	1/1	0.88	0.18	-3.06	39,39,39,39	0
56	MG	2a	3077	1/1	0.78	0.11	-3.07	59,59,59,59	0
56	MG	1a	1755	1/1	0.92	0.09	-3.10	79,79,79,79	0
56	MG	1a	1635	1/1	0.94	0.16	-3.11	50,50,50,50	0
56	MG	2A	3212	1/1	0.94	0.15	-3.11	23,23,23,23	0
56	MG	2A	3428	1/1	0.93	0.14	-3.14	49,49,49,49	0
56	MG	2A	3552	1/1	0.95	0.11	-3.15	22,22,22,22	0
56	MG	2A	3320	1/1	0.93	0.14	-3.17	33,33,33,33	0
56	MG	1A	3989	1/1	0.95	0.13	-3.18	25,25,25,25	0
56	MG	2A	3335	1/1	0.97	0.17	-3.23	40,40,40,40	0
56	MG	2A	3317	1/1	0.92	0.15	-3.25	34,34,34,34	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3790	1/1	0.97	0.18	-3.27	48,48,48,48	0
56	MG	2A	3206	1/1	0.93	0.12	-3.29	41,41,41,41	0
56	MG	2A	3189	1/1	0.93	0.13	-3.32	34,34,34,34	0
56	MG	2A	3534	1/1	0.98	0.09	-3.33	64,64,64,64	0
56	MG	1A	3722	1/1	0.89	0.19	-3.34	22,22,22,22	0
56	MG	2a	3183	1/1	0.98	0.14	-3.34	37,37,37,37	0
56	MG	2A	3058	1/1	0.94	0.16	-3.37	37,37,37,37	0
56	MG	1A	3606	1/1	0.79	0.19	-3.37	32,32,32,32	0
56	MG	1a	1684	1/1	0.94	0.10	-3.38	48,48,48,48	0
56	MG	2A	3359	1/1	0.94	0.12	-3.38	50,50,50,50	0
56	MG	2a	3193	1/1	0.97	0.12	-3.43	47,47,47,47	0
56	MG	1a	1724	1/1	0.84	0.09	-3.45	76,76,76,76	0
56	MG	1F	305	1/1	0.90	0.15	-3.49	44,44,44,44	0
56	MG	2A	3446	1/1	0.93	0.10	-3.50	42,42,42,42	0
56	MG	1A	3621	1/1	0.90	0.14	-3.53	18,18,18,18	0
56	MG	1X	101	1/1	0.96	0.18	-3.55	45,45,45,45	0
56	MG	1A	3893	1/1	0.95	0.18	-3.55	30,30,30,30	0
56	MG	1A	3727	1/1	0.95	0.19	-3.57	26,26,26,26	0
56	MG	1A	3280	1/1	0.90	0.14	-3.60	34,34,34,34	0
56	MG	1a	1715	1/1	0.95	0.07	-3.60	39,39,39,39	0
56	MG	1A	3780	1/1	0.63	0.14	-3.61	43,43,43,43	0
56	MG	2A	3052	1/1	0.99	0.14	-3.64	43,43,43,43	0
56	MG	1A	3565	1/1	0.94	0.18	-3.64	45,45,45,45	0
56	MG	2B	3006	1/1	0.93	0.09	-3.68	58,58,58,58	0
56	MG	1a	1666	1/1	0.90	0.11	-3.69	71,71,71,71	0
56	MG	1A	3733	1/1	0.93	0.13	-3.70	35,35,35,35	0
56	MG	2A	3369	1/1	0.83	0.15	-3.71	31,31,31,31	0
56	MG	1A	3193	1/1	0.94	0.16	-3.72	23,23,23,23	0
56	MG	1a	1716	1/1	0.99	0.08	-3.73	53,53,53,53	0
56	MG	1Q	3002	1/1	0.97	0.11	-3.74	44,44,44,44	0
56	MG	2a	3260	1/1	0.93	0.08	-3.74	72,72,72,72	0
56	MG	2A	3542	1/1	0.88	0.15	-3.75	42,42,42,42	0
56	MG	2A	3499	1/1	0.89	0.14	-3.77	47,47,47,47	0
56	MG	1A	3963	1/1	0.97	0.16	-3.78	14,14,14,14	0
56	MG	1a	1813	1/1	0.93	0.16	-3.81	59,59,59,59	0
56	MG	1A	3792	1/1	0.91	0.16	-3.82	79,79,79,79	0
56	MG	1A	3665	1/1	0.81	0.20	-3.89	22,22,22,22	0
56	MG	2a	3118	1/1	0.90	0.17	-3.91	38,38,38,38	0
56	MG	1A	3547	1/1	0.83	0.16	-3.93	54,54,54,54	0
56	MG	2f	3001	1/1	0.96	0.09	-3.97	59,59,59,59	0
56	MG	1A	3317	1/1	0.95	0.18	-3.97	34,34,34,34	0
56	MG	1A	3910	1/1	0.94	0.14	-3.99	65,65,65,65	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	2A	3377	1/1	0.95	0.11	-4.01	43,43,43,43	0
56	MG	1A	3181	1/1	0.97	0.16	-4.02	24,24,24,24	0
56	MG	1A	3672	1/1	0.98	0.12	-4.05	13,13,13,13	0
56	MG	1A	3791	1/1	0.97	0.17	-4.09	20,20,20,20	0
56	MG	1A	3897	1/1	0.97	0.18	-4.09	29,29,29,29	0
56	MG	1A	3401	1/1	0.96	0.14	-4.14	46,46,46,46	0
56	MG	2a	3230	1/1	0.92	0.14	-4.16	50,50,50,50	0
56	MG	1A	3537	1/1	0.95	0.15	-4.16	32,32,32,32	0
56	MG	1A	3774	1/1	0.97	0.12	-4.22	20,20,20,20	0
56	MG	2A	3205	1/1	0.96	0.12	-4.22	51,51,51,51	0
56	MG	1A	3730	1/1	0.91	0.15	-4.23	30,30,30,30	0
56	MG	1a	1720	1/1	0.94	0.10	-4.23	50,50,50,50	0
56	MG	2A	3385	1/1	0.95	0.14	-4.26	35,35,35,35	0
56	MG	2a	3213	1/1	0.96	0.10	-4.26	67,67,67,67	0
56	MG	2A	3310	1/1	0.88	0.14	-4.27	31,31,31,31	0
56	MG	2A	3295	1/1	0.83	0.14	-4.27	48,48,48,48	0
56	MG	1A	4009	1/1	0.97	0.17	-4.28	56,56,56,56	0
56	MG	1A	3770	1/1	0.97	0.17	-4.30	36,36,36,36	0
56	MG	1A	3205	1/1	0.94	0.17	-4.31	41,41,41,41	0
56	MG	1A	3768	1/1	0.97	0.15	-4.32	10,10,10,10	0
56	MG	2A	3420	1/1	0.98	0.12	-4.33	53,53,53,53	0
56	MG	2A	3268	1/1	0.88	0.16	-4.35	55,55,55,55	0
56	MG	1a	1731	1/1	0.95	0.11	-4.41	57,57,57,57	0
56	MG	1E	305	1/1	0.93	0.14	-4.42	31,31,31,31	0
56	MG	1A	3309	1/1	0.97	0.10	-4.43	20,20,20,20	0
56	MG	2A	3531	1/1	0.95	0.09	-4.46	52,52,52,52	0
56	MG	1A	3999	1/1	0.89	0.14	-4.46	21,21,21,21	0
56	MG	1A	3327	1/1	0.80	0.13	-4.51	73,73,73,73	0
56	MG	2A	3502	1/1	0.89	0.13	-4.54	35,35,35,35	0
56	MG	2A	3429	1/1	0.90	0.12	-4.54	58,58,58,58	0
56	MG	1A	3822	1/1	0.78	0.15	-4.55	46,46,46,46	0
56	MG	1A	3761	1/1	0.93	0.10	-4.59	66,66,66,66	0
56	MG	1A	3597	1/1	0.93	0.12	-4.60	48,48,48,48	0
56	MG	2A	3126	1/1	0.80	0.11	-4.60	50,50,50,50	0
56	MG	2a	3142	1/1	0.96	0.14	-4.61	51,51,51,51	0
56	MG	1A	3604	1/1	0.97	0.14	-4.72	27,27,27,27	0
56	MG	2A	3324	1/1	0.95	0.12	-4.74	34,34,34,34	0
56	MG	1a	1697	1/1	0.94	0.10	-4.79	52,52,52,52	0
56	MG	2A	3115	1/1	0.93	0.13	-4.79	35,35,35,35	0
56	MG	1A	3933	1/1	0.93	0.13	-4.80	42,42,42,42	0
56	MG	1F	302	1/1	0.96	0.15	-4.85	51,51,51,51	0
56	MG	1D	307	1/1	0.95	0.13	-4.88	15,15,15,15	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1a	1663	1/1	0.98	0.08	-4.88	76,76,76,76	0
56	MG	1a	1641	1/1	0.97	0.08	-4.90	55,55,55,55	0
56	MG	1a	1675	1/1	0.92	0.15	-4.92	56,56,56,56	0
56	MG	1A	3227	1/1	0.94	0.15	-4.93	41,41,41,41	0
56	MG	2A	3270	1/1	0.90	0.11	-4.94	45,45,45,45	0
56	MG	2a	3134	1/1	0.86	0.11	-4.94	59,59,59,59	0
56	MG	1A	3169	1/1	0.98	0.17	-4.94	21,21,21,21	0
56	MG	1A	3283	1/1	0.90	0.13	-4.96	44,44,44,44	0
56	MG	1A	3624	1/1	0.94	0.17	-4.99	18,18,18,18	0
56	MG	2A	3110	1/1	0.80	0.17	-4.99	45,45,45,45	0
56	MG	1A	3550	1/1	0.98	0.16	-5.03	33,33,33,33	0
56	MG	2A	3150	1/1	0.98	0.15	-5.05	30,30,30,30	0
56	MG	1A	3248	1/1	0.94	0.12	-5.07	26,26,26,26	0
56	MG	1A	3194	1/1	0.98	0.17	-5.07	32,32,32,32	0
56	MG	2a	3093	1/1	0.89	0.10	-5.11	50,50,50,50	0
56	MG	1A	3723	1/1	0.92	0.18	-5.16	8,8,8,8	0
56	MG	2A	3068	1/1	0.92	0.12	-5.26	39,39,39,39	0
56	MG	2A	3302	1/1	0.90	0.16	-5.26	41,41,41,41	0
56	MG	1A	3739	1/1	0.96	0.16	-5.27	30,30,30,30	0
56	MG	2a	3132	1/1	0.96	0.10	-5.28	65,65,65,65	0
56	MG	1a	1856	1/1	0.96	0.16	-5.29	43,43,43,43	0
56	MG	1A	3826	1/1	0.93	0.16	-5.36	37,37,37,37	0
56	MG	2A	3130	1/1	0.90	0.12	-5.39	39,39,39,39	0
56	MG	1A	3908	1/1	0.99	0.16	-5.42	40,40,40,40	0
56	MG	1a	1876	1/1	0.83	0.12	-5.43	80,80,80,80	0
56	MG	1A	3969	1/1	0.96	0.11	-5.43	18,18,18,18	0
56	MG	2A	3221	1/1	0.92	0.09	-5.43	47,47,47,47	0
56	MG	2A	3278	1/1	0.93	0.12	-5.44	31,31,31,31	0
56	MG	1A	3859	1/1	0.96	0.12	-5.50	24,24,24,24	0
56	MG	1A	3629	1/1	0.98	0.14	-5.51	27,27,27,27	0
56	MG	1A	3996	1/1	0.91	0.09	-5.54	40,40,40,40	0
56	MG	2A	3471	1/1	0.95	0.14	-5.57	34,34,34,34	0
56	MG	1A	3157	1/1	0.98	0.15	-5.67	28,28,28,28	0
56	MG	1a	1669	1/1	0.93	0.12	-5.70	46,46,46,46	0
56	MG	1A	3158	1/1	0.97	0.13	-5.73	24,24,24,24	0
56	MG	2a	3139	1/1	0.87	0.12	-5.74	70,70,70,70	0
56	MG	2a	3179	1/1	0.95	0.12	-5.75	49,49,49,49	0
56	MG	1A	3288	1/1	0.96	0.15	-5.76	40,40,40,40	0
56	MG	1A	3460	1/1	0.90	0.15	-5.78	36,36,36,36	0
56	MG	1A	3659	1/1	0.96	0.09	-5.80	23,23,23,23	0
56	MG	2A	3291	1/1	0.94	0.15	-5.81	36,36,36,36	0
56	MG	1A	3670	1/1	0.93	0.13	-5.89	55,55,55,55	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3596	1/1	0.94	0.11	-5.91	49,49,49,49	0
56	MG	1A	3839	1/1	0.95	0.14	-5.92	40,40,40,40	0
56	MG	2A	3518	1/1	0.96	0.11	-5.96	42,42,42,42	0
56	MG	2A	3519	1/1	0.97	0.06	-5.97	45,45,45,45	0
56	MG	1A	3622	1/1	0.97	0.13	-6.00	22,22,22,22	0
56	MG	1A	3535	1/1	0.95	0.11	-6.01	41,41,41,41	0
56	MG	2A	3263	1/1	0.95	0.09	-6.08	56,56,56,56	0
56	MG	2a	3049	1/1	0.90	0.10	-6.09	64,64,64,64	0
56	MG	1A	3559	1/1	0.97	0.14	-6.21	51,51,51,51	0
56	MG	2A	3442	1/1	0.92	0.11	-6.25	50,50,50,50	0
56	MG	1A	3616	1/1	0.93	0.14	-6.25	22,22,22,22	0
56	MG	2A	3283	1/1	0.94	0.11	-6.28	48,48,48,48	0
56	MG	1A	3611	1/1	0.91	0.11	-6.28	58,58,58,58	0
56	MG	2A	3051	1/1	0.95	0.10	-6.29	40,40,40,40	0
56	MG	1A	3626	1/1	0.95	0.16	-6.33	25,25,25,25	0
56	MG	1A	3906	1/1	0.93	0.16	-6.44	49,49,49,49	0
56	MG	2A	3273	1/1	0.88	0.06	-6.44	41,41,41,41	0
56	MG	1A	3546	1/1	0.97	0.19	-6.45	36,36,36,36	0
56	MG	2A	3344	1/1	0.97	0.14	-6.54	39,39,39,39	0
56	MG	1A	3811	1/1	0.93	0.09	-6.55	39,39,39,39	0
56	MG	1A	3694	1/1	0.99	0.13	-6.58	16,16,16,16	0
56	MG	1a	1607	1/1	0.85	0.10	-6.72	78,78,78,78	0
56	MG	2A	3351	1/1	0.99	0.11	-6.75	37,37,37,37	0
56	MG	1a	1691	1/1	0.88	0.12	-6.83	27,27,27,27	0
56	MG	1A	3610	1/1	0.79	0.10	-6.83	37,37,37,37	0
56	MG	2A	3062	1/1	0.95	0.10	-6.86	39,39,39,39	0
56	MG	1A	3922	1/1	0.97	0.13	-6.86	19,19,19,19	0
56	MG	1A	3581	1/1	0.96	0.10	-6.88	37,37,37,37	0
56	MG	1A	3263	1/1	0.89	0.14	-6.90	37,37,37,37	0
56	MG	1A	3293	1/1	0.95	0.13	-6.98	27,27,27,27	0
56	MG	1a	1752	1/1	0.97	0.06	-6.98	72,72,72,72	0
56	MG	1B	210	1/1	0.95	0.14	-7.03	49,49,49,49	0
56	MG	1a	1808	1/1	0.95	0.11	-7.08	52,52,52,52	0
56	MG	2A	3345	1/1	0.97	0.12	-7.12	35,35,35,35	0
56	MG	2A	3421	1/1	0.98	0.09	-7.13	59,59,59,59	0
56	MG	1A	3681	1/1	0.98	0.13	-7.16	14,14,14,14	0
56	MG	1a	1871	1/1	0.91	0.11	-7.17	46,46,46,46	0
56	MG	2A	3460	1/1	0.70	0.10	-7.17	54,54,54,54	0
56	MG	1a	1632	1/1	0.97	0.13	-7.17	41,41,41,41	0
56	MG	2A	3099	1/1	0.97	0.10	-7.19	30,30,30,30	0
56	MG	1A	3679	1/1	0.95	0.13	-7.26	25,25,25,25	0
56	MG	1A	3874	1/1	0.88	0.16	-7.29	56,56,56,56	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3825	1/1	0.96	0.12	-7.32	30,30,30,30	0
56	MG	1A	3793	1/1	0.93	0.10	-7.43	40,40,40,40	0
56	MG	2a	3090	1/1	0.93	0.10	-7.45	42,42,42,42	0
56	MG	2A	3487	1/1	0.90	0.08	-7.51	32,32,32,32	0
56	MG	1A	3161	1/1	0.96	0.09	-7.60	33,33,33,33	0
56	MG	1A	3336	1/1	0.93	0.14	-7.83	49,49,49,49	0
56	MG	1A	3639	1/1	0.93	0.12	-7.98	25,25,25,25	0
56	MG	1a	1603	1/1	0.95	0.09	-8.01	57,57,57,57	0
56	MG	1A	3151	1/1	0.94	0.08	-8.23	47,47,47,47	0
56	MG	2a	3251	1/1	0.90	0.08	-8.24	57,57,57,57	0
56	MG	1A	3676	1/1	0.88	0.16	-8.25	50,50,50,50	0
56	MG	1B	218	1/1	0.95	0.12	-8.30	39,39,39,39	0
56	MG	1A	3660	1/1	0.97	0.10	-8.35	11,11,11,11	0
56	MG	20	8001	1/1	0.92	0.08	-8.52	65,65,65,65	0
56	MG	1A	3749	1/1	0.92	0.12	-8.52	27,27,27,27	0
56	MG	1A	3217	1/1	0.94	0.12	-8.55	42,42,42,42	0
56	MG	1A	3576	1/1	0.98	0.12	-8.58	19,19,19,19	0
56	MG	1A	3553	1/1	0.99	0.14	-8.65	44,44,44,44	0
56	MG	1A	3190	1/1	0.95	0.15	-8.79	44,44,44,44	0
56	MG	1A	3803	1/1	0.96	0.08	-9.12	29,29,29,29	0
56	MG	1B	220	1/1	0.92	0.14	-9.38	70,70,70,70	0
56	MG	1A	3900	1/1	0.93	0.11	-9.46	66,66,66,66	0
56	MG	2A	3485	1/1	0.90	0.13	-9.70	38,38,38,38	0
56	MG	1A	3262	1/1	0.85	0.06	-9.86	66,66,66,66	0
56	MG	1A	3648	1/1	0.83	0.13	-9.99	20,20,20,20	0
56	MG	1A	3569	1/1	0.93	0.07	-10.32	45,45,45,45	0
56	MG	2a	3103	1/1	0.93	0.12	-10.37	69,69,69,69	0
56	MG	1A	3631	1/1	0.94	0.12	-10.54	19,19,19,19	0
56	MG	1A	3978	1/1	0.95	0.12	-10.93	29,29,29,29	0
56	MG	1a	1700	1/1	0.94	0.10	-10.94	36,36,36,36	0
56	MG	1A	3275	1/1	0.94	0.09	-10.97	25,25,25,25	0
56	MG	1A	3872	1/1	0.92	0.12	-11.13	28,28,28,28	0
56	MG	1A	3757	1/1	0.85	0.11	-11.42	39,39,39,39	0
56	MG	1A	3690	1/1	0.98	0.15	-11.74	48,48,48,48	0
56	MG	1A	3752	1/1	0.95	0.10	-11.96	50,50,50,50	0
56	MG	1A	3643	1/1	0.95	0.13	-12.02	19,19,19,19	0
56	MG	1A	3056	1/1	0.93	0.12	-12.05	62,62,62,62	0
56	MG	1A	3716	1/1	0.96	0.12	-12.19	43,43,43,43	0
56	MG	1A	3789	1/1	0.96	0.08	-12.81	26,26,26,26	0
56	MG	1A	3294	1/1	0.92	0.10	-13.29	23,23,23,23	0
56	MG	1A	3561	1/1	0.97	0.10	-13.90	44,44,44,44	0
56	MG	2A	3406	1/1	0.96	0.07	-15.50	41,41,41,41	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1a	1661	1/1	0.94	0.09	-16.18	15,15,15,15	0
56	MG	1A	3271	1/1	0.97	0.08	-16.19	17,17,17,17	0
56	MG	1A	3602	1/1	0.93	0.10	-16.32	45,45,45,45	0
56	MG	2A	3163	1/1	0.98	0.08	-21.60	33,33,33,33	0
56	MG	1A	3541	1/1	0.91	0.07	-26.45	48,48,48,48	0
56	MG	2A	3011	1/1	0.75	0.35	-	61,61,61,61	0
56	MG	1A	3156	1/1	0.97	0.16	-	47,47,47,47	0
56	MG	1a	1775	1/1	0.81	0.35	-	80,80,80,80	0
56	MG	1A	3745	1/1	0.85	0.68	-	71,71,71,71	0
56	MG	1a	1865	1/1	0.91	0.17	-	69,69,69,69	0
56	MG	1A	3045	1/1	0.89	0.10	-	70,70,70,70	0
56	MG	2A	3475	1/1	0.94	0.18	-	35,35,35,35	0
56	MG	2A	3342	1/1	0.79	0.26	-	63,63,63,63	0
56	MG	1A	3136	1/1	0.92	0.16	-	60,60,60,60	0
56	MG	2A	3148	1/1	0.83	0.26	-	63,63,63,63	0
56	MG	1A	3871	1/1	0.91	0.26	-	60,60,60,60	0
56	MG	1A	3710	1/1	0.97	0.24	-	39,39,39,39	0
56	MG	1A	3113	1/1	0.92	0.14	-	63,63,63,63	0
56	MG	2A	3236	1/1	0.83	0.06	-	91,91,91,91	0
56	MG	1A	3592	1/1	0.96	0.26	-	47,47,47,47	0
56	MG	1a	1746	1/1	0.95	0.23	-	55,55,55,55	0
56	MG	2F	301	1/1	0.93	0.11	-	57,57,57,57	0
56	MG	2A	3135	1/1	0.91	0.13	-	61,61,61,61	0
56	MG	2A	3219	1/1	0.90	0.30	-	44,44,44,44	0
56	MG	1A	3965	1/1	0.87	0.43	-	53,53,53,53	0
56	MG	1A	3357	1/1	0.70	0.21	-	68,68,68,68	0
56	MG	2A	3549	1/1	0.97	0.27	-	43,43,43,43	0
56	MG	1a	1809	1/1	0.94	0.18	-	62,62,62,62	0
56	MG	1A	3542	1/1	0.87	0.15	-	26,26,26,26	0
56	MG	1A	3837	1/1	0.93	0.08	-	62,62,62,62	0
56	MG	2a	3015	1/1	0.80	0.15	-	67,67,67,67	0
56	MG	1a	1605	1/1	0.96	0.13	-	55,55,55,55	0
56	MG	1A	3960	1/1	0.94	0.67	-	53,53,53,53	0
56	MG	1A	3974	1/1	0.91	0.12	-	30,30,30,30	0
56	MG	1A	3720	1/1	0.97	0.10	-	45,45,45,45	0
56	MG	2a	3130	1/1	0.98	0.08	-	43,43,43,43	0
56	MG	1A	3423	1/1	0.90	0.22	-	57,57,57,57	0
56	MG	2a	3165	1/1	0.93	0.15	-	52,52,52,52	0
56	MG	1A	3122	1/1	0.93	0.23	-	58,58,58,58	0
56	MG	2a	3055	1/1	0.90	0.05	-	98,98,98,98	0
56	MG	2A	3180	1/1	0.92	0.09	-	37,37,37,37	0
56	MG	2A	3095	1/1	0.95	0.19	-	31,31,31,31	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3759	1/1	0.92	0.15	-	60,60,60,60	0
56	MG	1A	3834	1/1	0.95	0.14	-	46,46,46,46	0
56	MG	2A	3443	1/1	0.48	0.23	-	56,56,56,56	0
56	MG	2a	3125	1/1	0.81	0.11	-	60,60,60,60	0
56	MG	1A	3005	1/1	0.80	0.24	-	58,58,58,58	0
56	MG	2A	3527	1/1	0.91	0.26	-	51,51,51,51	0
56	MG	1A	3098	1/1	0.87	0.18	-	52,52,52,52	0
56	MG	1a	1863	1/1	0.95	0.11	-	51,51,51,51	0
56	MG	1A	3375	1/1	0.91	0.62	-	51,51,51,51	0
56	MG	1A	3032	1/1	0.61	0.16	-	63,63,63,63	0
56	MG	1A	3290	1/1	0.80	0.51	-	37,37,37,37	0
56	MG	2A	3473	1/1	0.81	0.14	-	64,64,64,64	0
56	MG	2A	3239	1/1	0.84	0.11	-	60,60,60,60	0
56	MG	1a	1828	1/1	0.83	0.11	-	61,61,61,61	0
56	MG	1A	3035	1/1	0.67	0.67	-	53,53,53,53	0
56	MG	1A	3012	1/1	0.94	0.21	-	47,47,47,47	0
56	MG	2a	3035	1/1	0.85	0.10	-	69,69,69,69	0
56	MG	2A	3031	1/1	0.82	0.15	-	51,51,51,51	0
56	MG	2a	3078	1/1	0.96	0.14	-	47,47,47,47	0
56	MG	2A	3325	1/1	0.93	0.18	-	52,52,52,52	0
56	MG	2a	3033	1/1	0.98	0.41	-	53,53,53,53	0
56	MG	2A	3388	1/1	0.79	0.21	-	45,45,45,45	0
56	MG	1A	3719	1/1	0.97	0.13	-	38,38,38,38	0
56	MG	2A	3279	1/1	0.93	0.12	-	62,62,62,62	0
56	MG	2a	3147	1/1	0.95	0.08	-	73,73,73,73	0
56	MG	1A	3695	1/1	0.95	0.23	-	41,41,41,41	0
56	MG	1x	108	1/1	0.88	0.14	-	70,70,70,70	0
56	MG	2a	3140	1/1	0.93	0.08	-	63,63,63,63	0
56	MG	2a	3189	1/1	0.90	0.21	-	51,51,51,51	0
56	MG	1a	1682	1/1	0.92	0.12	-	46,46,46,46	0
56	MG	1a	1771	1/1	0.82	0.16	-	63,63,63,63	0
56	MG	1A	3350	1/1	0.89	0.14	-	54,54,54,54	0
56	MG	2A	3049	1/1	0.82	0.14	-	52,52,52,52	0
56	MG	1a	1791	1/1	0.90	0.07	-	99,99,99,99	0
56	MG	1D	305	1/1	0.92	0.28	-	39,39,39,39	0
56	MG	1A	3089	1/1	0.91	0.45	-	55,55,55,55	0
56	MG	2A	3414	1/1	0.94	0.07	-	49,49,49,49	0
56	MG	1A	3024	1/1	0.84	0.20	-	63,63,63,63	0
56	MG	2A	3002	1/1	0.94	0.15	-	47,47,47,47	0
56	MG	1B	209	1/1	0.84	0.32	-	62,62,62,62	0
56	MG	1a	1616	1/1	0.94	0.27	-	74,74,74,74	0
56	MG	2A	3444	1/1	0.99	0.21	-	29,29,29,29	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1B	214	1/1	0.92	0.08	-	68,68,68,68	0
56	MG	1B	208	1/1	0.87	0.25	-	56,56,56,56	0
56	MG	1a	1844	1/1	0.97	0.10	-	43,43,43,43	0
56	MG	1A	3609	1/1	0.92	0.17	-	22,22,22,22	0
56	MG	1A	3832	1/1	0.91	0.11	-	57,57,57,57	0
56	MG	19	102	1/1	0.94	0.13	-	51,51,51,51	0
56	MG	2A	3435	1/1	0.94	0.07	-	53,53,53,53	0
56	MG	2A	3526	1/1	0.97	0.09	-	49,49,49,49	0
56	MG	1A	3020	1/1	0.89	0.13	-	54,54,54,54	0
56	MG	1A	3865	1/1	0.85	0.25	-	68,68,68,68	0
56	MG	2A	3020	1/1	0.96	0.15	-	48,48,48,48	0
56	MG	2a	3196	1/1	0.92	0.17	-	90,90,90,90	0
56	MG	1A	3414	1/1	0.93	0.11	-	65,65,65,65	0
56	MG	1A	3988	1/1	0.91	0.74	-	45,45,45,45	0
56	MG	1A	3476	1/1	0.95	0.23	-	53,53,53,53	0
56	MG	2a	3058	1/1	0.80	0.17	-	87,87,87,87	0
56	MG	1A	3453	1/1	0.96	0.17	-	18,18,18,18	0
56	MG	1A	3077	1/1	0.86	0.24	-	69,69,69,69	0
56	MG	2A	3394	1/1	0.97	0.22	-	54,54,54,54	0
56	MG	2A	3026	1/1	0.92	0.17	-	51,51,51,51	0
56	MG	1A	3507	1/1	0.89	0.89	-	64,64,64,64	0
56	MG	1A	3139	1/1	0.93	0.41	-	41,41,41,41	0
56	MG	2A	3340	1/1	0.94	0.16	-	56,56,56,56	0
56	MG	1A	3101	1/1	0.86	0.29	-	55,55,55,55	0
56	MG	1a	1727	1/1	0.88	0.13	-	54,54,54,54	0
56	MG	1a	1774	1/1	0.94	0.09	-	76,76,76,76	0
56	MG	2a	3158	1/1	0.84	0.11	-	74,74,74,74	0
56	MG	1A	3855	1/1	0.83	0.19	-	84,84,84,84	0
56	MG	1A	3061	1/1	0.94	0.17	-	44,44,44,44	0
56	MG	1A	3244	1/1	0.96	0.12	-	67,67,67,67	0
56	MG	1A	3878	1/1	0.87	0.19	-	43,43,43,43	0
56	MG	1A	3876	1/1	0.89	0.22	-	79,79,79,79	0
56	MG	1A	3482	1/1	0.82	0.18	-	48,48,48,48	0
56	MG	2a	3052	1/1	0.81	0.08	-	95,95,95,95	0
56	MG	2A	3264	1/1	0.87	0.10	-	64,64,64,64	0
56	MG	1A	3049	1/1	0.93	0.28	-	48,48,48,48	0
56	MG	1A	3047	1/1	0.92	0.17	-	67,67,67,67	0
56	MG	2a	3240	1/1	0.87	0.23	-	81,81,81,81	0
56	MG	2a	3135	1/1	0.89	0.11	-	60,60,60,60	0
56	MG	2a	3071	1/1	0.93	0.30	-	59,59,59,59	0
56	MG	1A	3118	1/1	0.84	0.15	-	68,68,68,68	0
56	MG	1G	3002	1/1	0.84	0.15	-	61,61,61,61	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	2h	3001	1/1	0.90	0.15	-	62,62,62,62	0
56	MG	2A	3372	1/1	0.97	0.18	-	45,45,45,45	0
56	MG	1A	3896	1/1	0.91	0.18	-	19,19,19,19	0
56	MG	1A	3633	1/1	0.89	0.22	-	71,71,71,71	0
56	MG	2a	3253	1/1	0.95	0.31	-	56,56,56,56	0
56	MG	2a	3210	1/1	0.81	0.11	-	62,62,62,62	0
56	MG	1A	3216	1/1	0.97	0.14	-	37,37,37,37	0
56	MG	2a	3143	1/1	0.95	0.20	-	88,88,88,88	0
56	MG	1A	3094	1/1	0.93	0.33	-	51,51,51,51	0
56	MG	1A	3088	1/1	0.98	0.44	-	60,60,60,60	0
56	MG	1A	3276	1/1	0.91	0.28	-	38,38,38,38	0
56	MG	2a	3010	1/1	0.91	0.12	-	74,74,74,74	0
56	MG	1A	3255	1/1	0.90	0.12	-	58,58,58,58	0
56	MG	1A	3781	1/1	0.97	0.12	-	40,40,40,40	0
56	MG	2A	3566	1/1	0.97	0.11	-	62,62,62,62	0
56	MG	1a	1816	1/1	0.97	0.12	-	40,40,40,40	0
56	MG	1a	1723	1/1	0.77	0.16	-	85,85,85,85	0
56	MG	1A	3123	1/1	0.88	0.19	-	44,44,44,44	0
56	MG	2A	3089	1/1	0.94	0.18	-	40,40,40,40	0
56	MG	1A	3701	1/1	0.96	0.09	-	68,68,68,68	0
56	MG	2k	8001	1/1	0.97	0.05	-	83,83,83,83	0
56	MG	1a	1734	1/1	0.95	0.10	-	45,45,45,45	0
56	MG	2A	3014	1/1	0.93	0.06	-	66,66,66,66	0
56	MG	1A	3947	1/1	0.93	0.36	-	43,43,43,43	0
56	MG	1a	1681	1/1	0.94	0.06	-	55,55,55,55	0
56	MG	1A	3055	1/1	0.95	0.65	-	59,59,59,59	0
56	MG	1A	3738	1/1	0.91	0.17	-	37,37,37,37	0
56	MG	2z	101	1/1	0.96	0.68	-	54,54,54,54	0
56	MG	1H	8001	1/1	0.85	0.09	-	58,58,58,58	0
56	MG	2A	3367	1/1	0.93	0.17	-	50,50,50,50	0
56	MG	2A	3424	1/1	0.96	0.14	-	80,80,80,80	0
56	MG	1a	1802	1/1	0.82	0.13	-	76,76,76,76	0
56	MG	1A	3247	1/1	0.97	0.61	-	43,43,43,43	0
56	MG	2A	3494	1/1	0.95	0.10	-	56,56,56,56	0
56	MG	1A	3155	1/1	0.97	0.19	-	37,37,37,37	0
56	MG	1A	3345	1/1	0.92	0.09	-	60,60,60,60	0
56	MG	2a	3155	1/1	0.90	0.10	-	79,79,79,79	0
56	MG	2A	3521	1/1	0.94	0.16	-	44,44,44,44	0
56	MG	2a	3164	1/1	0.91	0.10	-	82,82,82,82	0
56	MG	2A	3450	1/1	0.91	0.08	-	62,62,62,62	0
56	MG	1A	3175	1/1	0.95	0.39	-	35,35,35,35	0
56	MG	1A	3620	1/1	0.98	0.11	-	38,38,38,38	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3765	1/1	0.91	0.10	-	44,44,44,44	0
56	MG	2A	3314	1/1	0.93	0.19	-	56,56,56,56	0
56	MG	2A	3012	1/1	0.94	0.24	-	53,53,53,53	0
56	MG	1A	3270	1/1	0.82	0.18	-	34,34,34,34	0
56	MG	2A	3528	1/1	0.94	0.25	-	49,49,49,49	0
56	MG	2a	3144	1/1	0.96	0.09	-	56,56,56,56	0
56	MG	1a	1679	1/1	0.89	0.20	-	57,57,57,57	0
56	MG	1A	3487	1/1	0.96	0.31	-	63,63,63,63	0
56	MG	1A	3976	1/1	0.93	0.28	-	69,69,69,69	0
56	MG	1A	3957	1/1	0.94	0.13	-	42,42,42,42	0
56	MG	1A	3590	1/1	0.96	0.06	-	47,47,47,47	0
56	MG	1A	3462	1/1	0.84	0.07	-	57,57,57,57	0
56	MG	2a	3019	1/1	0.93	0.11	-	73,73,73,73	0
56	MG	1A	3848	1/1	0.95	0.24	-	61,61,61,61	0
56	MG	1a	1741	1/1	0.84	0.14	-	63,63,63,63	0
56	MG	1Q	3001	1/1	0.90	0.29	-	38,38,38,38	0
56	MG	1A	3226	1/1	0.83	0.10	-	41,41,41,41	0
56	MG	1A	3396	1/1	0.81	0.24	-	67,67,67,67	0
56	MG	2a	3258	1/1	0.89	0.21	-	69,69,69,69	0
56	MG	2a	3032	1/1	0.90	0.22	-	50,50,50,50	0
56	MG	1A	3398	1/1	0.90	0.56	-	43,43,43,43	0
56	MG	1A	3016	1/1	0.96	0.23	-	61,61,61,61	0
56	MG	1A	3923	1/1	0.73	0.24	-	53,53,53,53	0
56	MG	1A	3545	1/1	0.90	0.17	-	17,17,17,17	0
56	MG	2A	3319	1/1	0.89	0.17	-	58,58,58,58	0
56	MG	1A	3773	1/1	0.94	0.27	-	80,80,80,80	0
56	MG	1a	1623	1/1	0.74	0.19	-	72,72,72,72	0
56	MG	2A	3379	1/1	0.97	0.09	-	42,42,42,42	0
56	MG	1a	1707	1/1	0.85	0.11	-	67,67,67,67	0
56	MG	1A	3771	1/1	0.92	0.49	-	71,71,71,71	0
56	MG	1a	1751	1/1	0.75	0.18	-	61,61,61,61	0
56	MG	1A	3387	1/1	0.88	1.26	-	56,56,56,56	0
56	MG	1a	1758	1/1	0.91	0.07	-	82,82,82,82	0
56	MG	2A	3039	1/1	0.80	0.20	-	49,49,49,49	0
56	MG	1A	3135	1/1	0.87	0.34	-	64,64,64,64	0
56	MG	2a	3243	1/1	0.95	0.10	-	67,67,67,67	0
56	MG	1B	203	1/1	0.90	0.21	-	61,61,61,61	0
56	MG	1a	1729	1/1	0.99	0.22	-	35,35,35,35	0
56	MG	2A	3466	1/1	0.99	0.15	-	69,69,69,69	0
56	MG	1A	3028	1/1	0.78	0.43	-	46,46,46,46	0
56	MG	2A	3226	1/1	0.94	0.21	-	57,57,57,57	0
56	MG	1A	3619	1/1	0.79	0.14	-	48,48,48,48	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1a	1606	1/1	0.89	0.12	-	76,76,76,76	0
56	MG	1a	1811	1/1	0.94	0.14	-	55,55,55,55	0
56	MG	2A	3267	1/1	0.92	0.15	-	39,39,39,39	0
56	MG	2A	3478	1/1	0.96	0.16	-	44,44,44,44	0
56	MG	2a	3220	1/1	0.93	0.10	-	63,63,63,63	0
56	MG	1A	3812	1/1	0.96	0.07	-	51,51,51,51	0
56	MG	1A	3446	1/1	0.82	0.32	-	62,62,62,62	0
56	MG	2A	3162	1/1	0.85	0.25	-	50,50,50,50	0
56	MG	1A	3458	1/1	0.89	0.23	-	46,46,46,46	0
56	MG	2A	3336	1/1	0.89	0.12	-	36,36,36,36	0
56	MG	2A	3079	1/1	0.95	0.33	-	52,52,52,52	0
56	MG	1A	3393	1/1	0.94	0.50	-	52,52,52,52	0
56	MG	1A	3531	1/1	0.93	0.22	-	33,33,33,33	0
56	MG	2a	3122	1/1	0.97	0.18	-	72,72,72,72	0
56	MG	1A	3953	1/1	0.80	0.30	-	71,71,71,71	0
56	MG	1A	3977	1/1	0.72	0.57	-	53,53,53,53	0
56	MG	1A	3472	1/1	0.85	0.67	-	83,83,83,83	0
56	MG	2Q	3005	1/1	0.94	0.52	-	52,52,52,52	0
56	MG	1A	3391	1/1	0.92	0.17	-	35,35,35,35	0
56	MG	1A	3506	1/1	0.95	0.14	-	58,58,58,58	0
56	MG	1O	3002	1/1	0.95	0.21	-	46,46,46,46	0
56	MG	1A	3220	1/1	0.89	0.15	-	60,60,60,60	0
56	MG	2A	3094	1/1	0.89	0.25	-	46,46,46,46	0
56	MG	1l	201	1/1	0.90	0.14	-	58,58,58,58	0
56	MG	1A	3373	1/1	0.97	0.06	-	48,48,48,48	0
56	MG	2A	3159	1/1	0.93	0.18	-	37,37,37,37	0
56	MG	1a	1851	1/1	0.97	0.11	-	74,74,74,74	0
56	MG	1a	1797	1/1	0.94	0.11	-	60,60,60,60	0
56	MG	1a	1609	1/1	0.81	0.20	-	63,63,63,63	0
56	MG	1a	1647	1/1	0.94	0.19	-	56,56,56,56	0
56	MG	2A	3145	1/1	0.64	0.22	-	77,77,77,77	0
56	MG	1a	1820	1/1	0.90	0.14	-	54,54,54,54	0
56	MG	1a	1776	1/1	0.77	0.18	-	69,69,69,69	0
56	MG	1A	3699	1/1	0.88	0.12	-	47,47,47,47	0
56	MG	2A	3334	1/1	0.75	0.18	-	58,58,58,58	0
56	MG	2a	3177	1/1	0.96	0.06	-	44,44,44,44	0
56	MG	1A	3041	1/1	0.92	0.22	-	55,55,55,55	0
56	MG	2A	3037	1/1	0.94	0.07	-	46,46,46,46	0
56	MG	1a	1806	1/1	0.97	0.14	-	47,47,47,47	0
56	MG	1A	3836	1/1	0.93	0.19	-	35,35,35,35	0
56	MG	1A	3316	1/1	0.95	0.26	-	48,48,48,48	0
56	MG	1A	3879	1/1	0.97	0.21	-	31,31,31,31	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	4001	1/1	0.88	0.81	-	43,43,43,43	0
56	MG	1a	1832	1/1	0.95	0.13	-	50,50,50,50	0
56	MG	2A	3417	1/1	0.96	0.13	-	70,70,70,70	0
56	MG	1A	3277	1/1	0.93	0.87	-	48,48,48,48	0
56	MG	1A	3432	1/1	0.92	0.15	-	58,58,58,58	0
56	MG	2A	3438	1/1	0.97	0.18	-	62,62,62,62	0
56	MG	1A	3380	1/1	0.85	0.39	-	63,63,63,63	0
56	MG	1A	3689	1/1	0.94	0.12	-	55,55,55,55	0
56	MG	2A	3142	1/1	0.84	0.19	-	50,50,50,50	0
56	MG	1A	3485	1/1	0.93	0.13	-	55,55,55,55	0
56	MG	1A	3905	1/1	0.94	0.12	-	51,51,51,51	0
56	MG	2A	3457	1/1	0.98	0.12	-	45,45,45,45	0
56	MG	1A	3291	1/1	0.91	0.52	-	48,48,48,48	0
56	MG	2A	3201	1/1	0.90	0.12	-	47,47,47,47	0
56	MG	2A	3085	1/1	0.92	0.27	-	53,53,53,53	0
56	MG	2a	3127	1/1	0.95	0.18	-	61,61,61,61	0
56	MG	2a	3085	1/1	0.98	0.07	-	40,40,40,40	0
56	MG	1A	3209	1/1	0.94	0.60	-	40,40,40,40	0
56	MG	2A	3129	1/1	0.95	0.16	-	49,49,49,49	0
56	MG	1a	1852	1/1	0.92	0.13	-	70,70,70,70	0
56	MG	1A	3246	1/1	0.87	0.20	-	58,58,58,58	0
56	MG	2A	3004	1/1	0.95	0.14	-	52,52,52,52	0
56	MG	2A	3210	1/1	0.30	0.17	-	63,63,63,63	0
56	MG	1A	3591	1/1	0.91	0.12	-	77,77,77,77	0
56	MG	1A	3742	1/1	0.95	0.20	-	35,35,35,35	0
56	MG	1A	3758	1/1	0.89	0.12	-	65,65,65,65	0
56	MG	1a	1794	1/1	0.90	0.16	-	67,67,67,67	0
56	MG	1A	3025	1/1	0.97	0.15	-	57,57,57,57	0
56	MG	1A	3071	1/1	0.97	0.18	-	44,44,44,44	0
56	MG	1A	3925	1/1	0.95	0.16	-	42,42,42,42	0
56	MG	1A	3909	1/1	0.94	0.12	-	62,62,62,62	0
56	MG	1A	3845	1/1	0.89	0.12	-	42,42,42,42	0
56	MG	1A	3383	1/1	0.95	0.20	-	56,56,56,56	0
56	MG	2a	3133	1/1	0.97	0.06	-	60,60,60,60	0
56	MG	1A	3053	1/1	0.92	0.22	-	53,53,53,53	0
56	MG	1A	3741	1/1	0.90	0.20	-	19,19,19,19	0
56	MG	1a	1753	1/1	0.94	0.08	-	70,70,70,70	0
56	MG	1A	3861	1/1	0.95	0.18	-	48,48,48,48	0
56	MG	1a	1689	1/1	0.83	0.11	-	65,65,65,65	0
56	MG	2A	3350	1/1	0.96	0.14	-	34,34,34,34	0
56	MG	1A	3478	1/1	0.91	0.39	-	65,65,65,65	0
56	MG	1a	1760	1/1	0.84	0.12	-	85,85,85,85	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3286	1/1	0.98	0.38	-	36,36,36,36	0
56	MG	1A	3264	1/1	0.93	0.39	-	38,38,38,38	0
56	MG	2A	3274	1/1	0.92	0.12	-	64,64,64,64	0
56	MG	2a	3185	1/1	0.56	0.16	-	67,67,67,67	0
56	MG	2A	3557	1/1	0.88	0.09	-	64,64,64,64	0
56	MG	1A	3429	1/1	0.48	0.32	-	70,70,70,70	0
56	MG	2A	3093	1/1	0.90	0.10	-	55,55,55,55	0
56	MG	1a	1818	1/1	0.95	0.16	-	63,63,63,63	0
56	MG	1A	3206	1/1	0.97	0.16	-	21,21,21,21	0
56	MG	1A	3579	1/1	0.93	0.14	-	45,45,45,45	0
56	MG	1A	3185	1/1	0.86	0.10	-	41,41,41,41	0
56	MG	1A	3483	1/1	0.89	0.49	-	54,54,54,54	0
56	MG	1a	1849	1/1	0.86	0.10	-	53,53,53,53	0
56	MG	1A	3653	1/1	0.92	0.15	-	25,25,25,25	0
56	MG	2a	3238	1/1	0.87	0.15	-	55,55,55,55	0
56	MG	1a	1645	1/1	0.96	0.11	-	67,67,67,67	0
56	MG	1A	3829	1/1	0.95	0.26	-	55,55,55,55	0
56	MG	1a	1785	1/1	0.94	0.26	-	63,63,63,63	0
56	MG	2B	3007	1/1	0.83	0.24	-	56,56,56,56	0
56	MG	1A	3493	1/1	0.93	0.15	-	65,65,65,65	0
56	MG	2A	3124	1/1	0.92	0.25	-	54,54,54,54	0
56	MG	1a	1639	1/1	0.94	0.15	-	62,62,62,62	0
56	MG	1A	3296	1/1	0.70	0.18	-	55,55,55,55	0
56	MG	1A	3087	1/1	0.92	0.57	-	57,57,57,57	0
56	MG	1a	1798	1/1	0.93	0.10	-	71,71,71,71	0
56	MG	1A	3838	1/1	0.76	0.21	-	66,66,66,66	0
56	MG	2A	3216	1/1	0.86	0.44	-	45,45,45,45	0
56	MG	1A	3021	1/1	0.93	1.05	-	48,48,48,48	0
56	MG	1A	3367	1/1	0.85	0.11	-	66,66,66,66	0
56	MG	1A	3967	1/1	0.93	0.11	-	19,19,19,19	0
56	MG	1A	3928	1/1	0.91	0.09	-	62,62,62,62	0
56	MG	1A	3103	1/1	0.89	0.21	-	50,50,50,50	0
56	MG	1A	3564	1/1	0.82	0.19	-	35,35,35,35	0
56	MG	2A	3363	1/1	0.94	0.17	-	61,61,61,61	0
56	MG	1A	3555	1/1	0.89	0.13	-	46,46,46,46	0
56	MG	1A	3755	1/1	0.82	0.10	-	66,66,66,66	0
56	MG	1A	3496	1/1	0.96	0.13	-	88,88,88,88	0
56	MG	2A	3140	1/1	0.93	0.10	-	58,58,58,58	0
56	MG	1A	3684	1/1	0.93	0.11	-	50,50,50,50	0
56	MG	2A	3127	1/1	0.88	0.16	-	57,57,57,57	0
56	MG	1A	3530	1/1	0.94	0.26	-	51,51,51,51	0
56	MG	1A	3532	1/1	0.97	0.10	-	62,62,62,62	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3574	1/1	0.97	0.21	-	48,48,48,48	0
56	MG	1A	3249	1/1	0.90	0.25	-	42,42,42,42	0
56	MG	1A	3338	1/1	0.70	0.34	-	54,54,54,54	0
56	MG	1A	3764	1/1	0.97	0.16	-	41,41,41,41	0
56	MG	1A	3802	1/1	0.96	0.11	-	77,77,77,77	0
56	MG	2A	3493	1/1	0.82	0.23	-	58,58,58,58	0
56	MG	1A	3744	1/1	0.92	0.11	-	53,53,53,53	0
56	MG	1A	3696	1/1	0.95	0.12	-	65,65,65,65	0
56	MG	1A	3265	1/1	0.96	0.14	-	51,51,51,51	0
56	MG	1A	3433	1/1	0.73	0.14	-	71,71,71,71	0
56	MG	2A	3244	1/1	0.94	0.10	-	63,63,63,63	0
56	MG	1A	3894	1/1	0.90	0.10	-	41,41,41,41	0
56	MG	2a	3018	1/1	0.81	0.19	-	49,49,49,49	0
56	MG	2A	3153	1/1	0.94	0.17	-	33,33,33,33	0
56	MG	1A	3076	1/1	0.89	0.48	-	58,58,58,58	0
56	MG	1A	3779	1/1	0.85	0.27	-	54,54,54,54	0
56	MG	2a	3242	1/1	0.93	0.12	-	61,61,61,61	0
56	MG	2a	3205	1/1	0.86	0.22	-	74,74,74,74	0
56	MG	2A	3246	1/1	0.86	0.17	-	36,36,36,36	0
56	MG	1A	3332	1/1	0.86	0.11	-	55,55,55,55	0
56	MG	1A	3036	1/1	0.90	0.42	-	55,55,55,55	0
56	MG	2A	3413	1/1	0.97	0.23	-	62,62,62,62	0
56	MG	2a	3069	1/1	0.89	0.34	-	95,95,95,95	0
56	MG	2A	3409	1/1	0.96	0.16	-	50,50,50,50	0
56	MG	1A	3392	1/1	0.62	0.77	-	52,52,52,52	0
56	MG	2a	3115	1/1	0.92	0.14	-	54,54,54,54	0
56	MG	1A	3067	1/1	0.84	0.19	-	46,46,46,46	0
56	MG	1A	3434	1/1	0.82	0.20	-	70,70,70,70	0
56	MG	1A	3095	1/1	0.81	0.32	-	50,50,50,50	0
56	MG	1A	3662	1/1	0.95	0.28	-	44,44,44,44	0
56	MG	1A	3390	1/1	0.86	0.72	-	40,40,40,40	0
56	MG	1a	1646	1/1	0.90	0.15	-	57,57,57,57	0
56	MG	1A	3199	1/1	0.96	0.20	-	37,37,37,37	0
56	MG	1A	3365	1/1	0.87	0.24	-	46,46,46,46	0
56	MG	1A	3027	1/1	0.79	0.34	-	54,54,54,54	0
56	MG	2a	3150	1/1	0.97	0.18	-	51,51,51,51	0
56	MG	1A	3614	1/1	0.86	0.23	-	37,37,37,37	0
56	MG	1a	1721	1/1	0.84	0.13	-	59,59,59,59	0
56	MG	10	103	1/1	0.92	0.10	-	75,75,75,75	0
56	MG	1W	3002	1/1	0.80	0.20	-	55,55,55,55	0
56	MG	2A	3461	1/1	0.91	0.14	-	47,47,47,47	0
56	MG	1A	3200	1/1	0.88	0.12	-	46,46,46,46	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3426	1/1	0.93	0.28	-	46,46,46,46	0
56	MG	2A	3121	1/1	0.96	0.18	-	32,32,32,32	0
56	MG	10	107	1/1	0.94	0.16	-	55,55,55,55	0
56	MG	2A	3015	1/1	0.91	0.22	-	52,52,52,52	0
56	MG	1A	3646	1/1	0.98	0.14	-	10,10,10,10	0
56	MG	2A	3358	1/1	0.82	0.27	-	59,59,59,59	0
56	MG	1a	1784	1/1	0.92	0.14	-	56,56,56,56	0
56	MG	1A	3445	1/1	0.69	0.29	-	63,63,63,63	0
56	MG	1A	3842	1/1	0.89	0.22	-	56,56,56,56	0
56	MG	1A	3295	1/1	0.92	0.18	-	50,50,50,50	0
56	MG	1B	206	1/1	0.98	0.26	-	46,46,46,46	0
56	MG	2A	3227	1/1	0.87	0.17	-	47,47,47,47	0
56	MG	2A	3515	1/1	0.70	0.09	-	63,63,63,63	0
56	MG	2a	3110	1/1	0.92	0.13	-	68,68,68,68	0
56	MG	2A	3080	1/1	0.90	0.16	-	48,48,48,48	0
56	MG	1A	3798	1/1	0.91	0.07	-	61,61,61,61	0
56	MG	2A	3479	1/1	0.94	0.38	-	50,50,50,50	0
56	MG	1A	3521	1/1	0.53	0.33	-	75,75,75,75	0
56	MG	1A	3174	1/1	0.98	0.66	-	37,37,37,37	0
56	MG	1A	3048	1/1	0.96	0.24	-	45,45,45,45	0
56	MG	1A	3344	1/1	0.89	0.10	-	72,72,72,72	0
56	MG	1A	3807	1/1	0.91	0.12	-	57,57,57,57	0
56	MG	2a	3256	1/1	0.77	0.13	-	94,94,94,94	0
56	MG	2a	3079	1/1	0.85	0.06	-	69,69,69,69	0
56	MG	1a	1750	1/1	0.84	0.38	-	64,64,64,64	0
56	MG	1A	3961	1/1	0.72	0.21	-	60,60,60,60	0
56	MG	1a	1869	1/1	0.89	0.16	-	64,64,64,64	0
56	MG	1A	3404	1/1	0.87	0.34	-	48,48,48,48	0
56	MG	2A	3366	1/1	0.98	0.20	-	69,69,69,69	0
56	MG	1A	3691	1/1	0.95	0.09	-	51,51,51,51	0
56	MG	2a	3186	1/1	0.91	0.14	-	48,48,48,48	0
56	MG	2A	3354	1/1	0.92	0.11	-	63,63,63,63	0
56	MG	1A	3418	1/1	0.94	0.10	-	66,66,66,66	0
56	MG	1a	1829	1/1	0.96	0.24	-	47,47,47,47	0
56	MG	2A	3211	1/1	0.85	0.29	-	63,63,63,63	0
56	MG	1a	1867	1/1	0.91	0.11	-	80,80,80,80	0
56	MG	1A	3788	1/1	0.94	0.15	-	23,23,23,23	0
56	MG	2Z	3001	1/1	0.98	0.11	-	68,68,68,68	0
56	MG	2A	3256	1/1	0.86	0.15	-	34,34,34,34	0
56	MG	2A	3175	1/1	0.93	0.24	-	52,52,52,52	0
56	MG	1A	3623	1/1	0.97	0.25	-	25,25,25,25	0
56	MG	1A	3543	1/1	0.93	0.18	-	65,65,65,65	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	4016	1/1	0.93	0.47	-	37,37,37,37	0
56	MG	1F	303	1/1	0.97	0.14	-	32,32,32,32	0
56	MG	1a	1617	1/1	0.82	0.30	-	78,78,78,78	0
56	MG	1a	1701	1/1	0.89	0.17	-	54,54,54,54	0
56	MG	2A	3285	1/1	0.96	0.19	-	39,39,39,39	0
56	MG	2A	3480	1/1	0.97	0.14	-	46,46,46,46	0
56	MG	2I	204	1/1	0.89	0.12	-	64,64,64,64	0
56	MG	1A	3514	1/1	0.90	0.18	-	53,53,53,53	0
56	MG	2A	3071	1/1	0.80	0.17	-	68,68,68,68	0
56	MG	1A	3866	1/1	0.96	0.15	-	37,37,37,37	0
56	MG	2A	3357	1/1	0.90	0.20	-	65,65,65,65	0
56	MG	1a	1836	1/1	0.92	0.16	-	41,41,41,41	0
56	MG	2a	3062	1/1	0.95	0.11	-	59,59,59,59	0
56	MG	1a	1841	1/1	0.91	0.18	-	64,64,64,64	0
56	MG	2A	3489	1/1	0.90	0.07	-	63,63,63,63	0
56	MG	1A	3144	1/1	0.89	0.36	-	54,54,54,54	0
56	MG	1A	3810	1/1	0.90	0.06	-	48,48,48,48	0
56	MG	2A	3535	1/1	0.97	0.12	-	44,44,44,44	0
56	MG	1a	1712	1/1	0.97	0.16	-	25,25,25,25	0
56	MG	1A	3259	1/1	0.98	0.16	-	38,38,38,38	0
56	MG	1a	1840	1/1	0.95	0.20	-	41,41,41,41	0
56	MG	1a	1713	1/1	0.98	0.13	-	63,63,63,63	0
56	MG	2A	3469	1/1	0.93	0.11	-	51,51,51,51	0
56	MG	1A	3278	1/1	0.82	0.13	-	56,56,56,56	0
56	MG	1A	3732	1/1	0.93	0.22	-	45,45,45,45	0
56	MG	1A	3402	1/1	0.88	0.15	-	58,58,58,58	0
56	MG	2A	3213	1/1	0.68	0.15	-	59,59,59,59	0
56	MG	2A	3152	1/1	0.97	0.10	-	58,58,58,58	0
56	MG	1a	1800	1/1	0.83	0.18	-	57,57,57,57	0
56	MG	1a	1733	1/1	0.92	0.12	-	65,65,65,65	0
56	MG	2a	3235	1/1	0.61	0.15	-	68,68,68,68	0
56	MG	2A	3459	1/1	0.97	0.11	-	49,49,49,49	0
56	MG	1A	3070	1/1	0.65	0.17	-	72,72,72,72	0
56	MG	2a	3128	1/1	0.93	0.11	-	38,38,38,38	0
56	MG	2B	3005	1/1	0.92	0.16	-	53,53,53,53	0
56	MG	1a	1672	1/1	0.82	0.15	-	58,58,58,58	0
56	MG	2A	3426	1/1	0.94	0.12	-	62,62,62,62	0
56	MG	2B	3010	1/1	0.93	0.32	-	50,50,50,50	0
56	MG	2a	3247	1/1	0.92	0.11	-	71,71,71,71	0
56	MG	1a	1633	1/1	0.90	0.14	-	38,38,38,38	0
56	MG	2A	3390	1/1	0.93	0.18	-	56,56,56,56	0
56	MG	1A	3636	1/1	0.96	0.16	-	43,43,43,43	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1a	1868	1/1	0.91	0.12	-	66,66,66,66	0
56	MG	1A	3323	1/1	0.90	0.36	-	58,58,58,58	0
56	MG	1A	3166	1/1	0.69	0.15	-	55,55,55,55	0
56	MG	1F	308	1/1	0.80	0.62	-	55,55,55,55	0
56	MG	1A	3841	1/1	0.95	0.10	-	36,36,36,36	0
56	MG	1A	3938	1/1	0.96	0.18	-	44,44,44,44	0
56	MG	2F	304	1/1	0.73	0.24	-	53,53,53,53	0
56	MG	1A	3347	1/1	0.95	0.16	-	60,60,60,60	0
56	MG	1A	3835	1/1	0.98	0.10	-	66,66,66,66	0
56	MG	1A	3795	1/1	0.94	0.15	-	64,64,64,64	0
56	MG	1a	1630	1/1	0.87	0.28	-	56,56,56,56	0
56	MG	1A	3729	1/1	0.87	0.17	-	48,48,48,48	0
56	MG	1A	3498	1/1	0.71	0.37	-	67,67,67,67	0
56	MG	18	101	1/1	0.87	0.11	-	63,63,63,63	0
56	MG	1A	3668	1/1	0.85	0.12	-	60,60,60,60	0
56	MG	2a	3136	1/1	0.98	0.09	-	51,51,51,51	0
56	MG	1A	3058	1/1	0.97	0.12	-	50,50,50,50	0
56	MG	1A	3862	1/1	0.85	0.21	-	62,62,62,62	0
56	MG	1a	1803	1/1	0.95	0.16	-	49,49,49,49	0
56	MG	2a	3029	1/1	0.87	0.28	-	69,69,69,69	0
56	MG	1a	1614	1/1	0.77	0.33	-	68,68,68,68	0
56	MG	2A	3257	1/1	0.75	0.19	-	62,62,62,62	0
56	MG	2A	3097	1/1	0.86	0.13	-	51,51,51,51	0
56	MG	2A	3445	1/1	0.64	0.16	-	71,71,71,71	0
56	MG	1A	3504	1/1	0.96	0.11	-	60,60,60,60	0
56	MG	2A	3030	1/1	0.88	0.09	-	50,50,50,50	0
56	MG	1A	3164	1/1	0.59	0.15	-	64,64,64,64	0
56	MG	2A	3289	1/1	0.91	0.20	-	58,58,58,58	0
56	MG	2a	3206	1/1	0.85	0.09	-	63,63,63,63	0
56	MG	2a	3017	1/1	0.96	0.23	-	70,70,70,70	0
56	MG	2A	3204	1/1	0.93	0.11	-	44,44,44,44	0
56	MG	1a	1862	1/1	0.83	0.13	-	65,65,65,65	0
56	MG	1D	313	1/1	0.90	0.58	-	56,56,56,56	0
56	MG	1A	3475	1/1	0.94	0.55	-	53,53,53,53	0
56	MG	1A	3698	1/1	0.88	0.07	-	48,48,48,48	0
56	MG	1A	3853	1/1	0.95	0.14	-	46,46,46,46	0
56	MG	1A	3138	1/1	0.94	0.59	-	55,55,55,55	0
56	MG	1A	3284	1/1	0.88	0.10	-	41,41,41,41	0
56	MG	2A	3309	1/1	0.83	0.18	-	31,31,31,31	0
56	MG	1A	3509	1/1	0.83	0.23	-	71,71,71,71	0
56	MG	2A	3437	1/1	0.92	0.12	-	57,57,57,57	0
56	MG	1A	3817	1/1	0.95	0.23	-	51,51,51,51	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3664	1/1	0.94	0.14	-	15,15,15,15	0
56	MG	2q	201	1/1	0.90	0.17	-	58,58,58,58	0
56	MG	2A	3050	1/1	0.96	0.16	-	38,38,38,38	0
56	MG	2A	3234	1/1	0.87	0.39	-	48,48,48,48	0
56	MG	2A	3404	1/1	0.94	0.10	-	43,43,43,43	0
56	MG	1B	222	1/1	0.90	0.09	-	70,70,70,70	0
56	MG	2A	3286	1/1	0.95	0.13	-	32,32,32,32	0
56	MG	1A	3465	1/1	0.90	0.19	-	44,44,44,44	0
56	MG	2a	3250	1/1	0.90	0.15	-	61,61,61,61	0
56	MG	2A	3383	1/1	0.93	0.22	-	66,66,66,66	0
56	MG	1A	3864	1/1	0.96	0.07	-	55,55,55,55	0
56	MG	1A	3508	1/1	0.71	0.44	-	57,57,57,57	0
56	MG	1E	302	1/1	0.93	0.10	-	48,48,48,48	0
56	MG	1a	1678	1/1	0.92	0.07	-	43,43,43,43	0
56	MG	2A	3200	1/1	0.96	0.17	-	48,48,48,48	0
56	MG	1A	3356	1/1	0.97	0.16	-	42,42,42,42	0
56	MG	2A	3423	1/1	0.95	0.06	-	66,66,66,66	0
56	MG	1a	1875	1/1	0.86	0.26	-	82,82,82,82	0
56	MG	2O	203	1/1	0.94	0.13	-	40,40,40,40	0
56	MG	2A	3119	1/1	0.88	0.13	-	61,61,61,61	0
56	MG	1A	3502	1/1	0.93	0.35	-	63,63,63,63	0
56	MG	2A	3096	1/1	0.79	0.16	-	40,40,40,40	0
56	MG	1A	3776	1/1	0.94	0.13	-	20,20,20,20	0
56	MG	26	101	1/1	0.90	0.10	-	57,57,57,57	0
56	MG	1A	4006	1/1	0.95	0.16	-	45,45,45,45	0
56	MG	2a	3149	1/1	0.93	0.12	-	65,65,65,65	0
56	MG	2A	3240	1/1	0.81	0.15	-	63,63,63,63	0
56	MG	2A	3125	1/1	0.98	0.20	-	56,56,56,56	0
56	MG	2a	3061	1/1	0.96	0.10	-	73,73,73,73	0
56	MG	2A	3008	1/1	0.73	0.12	-	64,64,64,64	0
56	MG	2A	3364	1/1	0.80	0.15	-	60,60,60,60	0
56	MG	2A	3076	1/1	0.69	0.20	-	60,60,60,60	0
56	MG	2A	3488	1/1	0.94	0.20	-	63,63,63,63	0
56	MG	2a	3039	1/1	0.94	0.17	-	57,57,57,57	0
56	MG	1A	3573	1/1	0.94	0.24	-	47,47,47,47	0
56	MG	1a	1756	1/1	0.90	0.11	-	75,75,75,75	0
56	MG	2B	3011	1/1	0.88	0.18	-	83,83,83,83	0
56	MG	2a	3109	1/1	0.95	0.12	-	47,47,47,47	0
56	MG	2A	3245	1/1	0.97	0.17	-	48,48,48,48	0
56	MG	1A	3490	1/1	0.89	0.12	-	66,66,66,66	0
56	MG	1A	3652	1/1	0.98	0.10	-	36,36,36,36	0
56	MG	2a	3045	1/1	0.91	0.22	-	74,74,74,74	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	2A	3229	1/1	0.76	0.21	-	55,55,55,55	0
56	MG	2a	3084	1/1	0.97	0.22	-	56,56,56,56	0
56	MG	13	101	1/1	0.95	0.13	-	49,49,49,49	0
56	MG	2A	3280	1/1	0.94	0.11	-	63,63,63,63	0
56	MG	1A	3685	1/1	0.87	0.26	-	73,73,73,73	0
56	MG	2a	3234	1/1	0.91	0.13	-	66,66,66,66	0
56	MG	1A	3650	1/1	0.95	0.18	-	19,19,19,19	0
56	MG	1a	1719	1/1	0.99	0.14	-	53,53,53,53	0
56	MG	2a	3214	1/1	0.68	0.20	-	97,97,97,97	0
56	MG	1A	3858	1/1	0.99	0.23	-	46,46,46,46	0
56	MG	1A	3427	1/1	0.83	0.15	-	53,53,53,53	0
56	MG	1A	3450	1/1	0.81	0.19	-	45,45,45,45	0
56	MG	2a	3046	1/1	0.85	0.09	-	65,65,65,65	0
56	MG	1A	3050	1/1	0.78	0.28	-	48,48,48,48	0
56	MG	1A	3527	1/1	0.63	0.19	-	41,41,41,41	0
56	MG	1A	3510	1/1	0.96	0.24	-	55,55,55,55	0
56	MG	1a	1656	1/1	0.96	0.06	-	57,57,57,57	0
56	MG	1A	3355	1/1	0.89	0.13	-	53,53,53,53	0
56	MG	1A	3471	1/1	0.84	0.09	-	74,74,74,74	0
56	MG	15	102	1/1	0.94	0.68	-	40,40,40,40	0
56	MG	2A	3266	1/1	0.81	0.13	-	71,71,71,71	0
56	MG	1A	3352	1/1	0.83	0.12	-	58,58,58,58	0
56	MG	1A	3751	1/1	0.84	0.13	-	39,39,39,39	0
56	MG	2A	3512	1/1	0.94	0.11	-	44,44,44,44	0
56	MG	1A	3148	1/1	0.94	0.15	-	62,62,62,62	0
56	MG	1A	3197	1/1	0.81	0.13	-	50,50,50,50	0
56	MG	2a	3094	1/1	0.91	0.17	-	49,49,49,49	0
56	MG	1A	3700	1/1	0.93	0.07	-	63,63,63,63	0
56	MG	2A	3484	1/1	0.86	0.14	-	82,82,82,82	0
56	MG	1P	202	1/1	0.91	0.20	-	55,55,55,55	0
56	MG	1A	3627	1/1	0.93	0.14	-	20,20,20,20	0
56	MG	2A	3380	1/1	0.85	0.27	-	74,74,74,74	0
56	MG	1A	3085	1/1	0.98	0.60	-	47,47,47,47	0
56	MG	1A	3484	1/1	0.97	0.17	-	48,48,48,48	0
56	MG	2A	3007	1/1	0.95	0.60	-	46,46,46,46	0
56	MG	2a	3053	1/1	0.94	0.14	-	85,85,85,85	0
56	MG	2a	3146	1/1	0.83	0.15	-	57,57,57,57	0
56	MG	2a	3126	1/1	0.96	0.21	-	41,41,41,41	0
56	MG	1A	3536	1/1	0.92	0.20	-	68,68,68,68	0
56	MG	1B	223	1/1	0.83	0.12	-	59,59,59,59	0
56	MG	1a	1667	1/1	0.88	0.15	-	65,65,65,65	0
56	MG	1A	3937	1/1	0.91	0.16	-	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3505	1/1	0.91	0.10	-	65,65,65,65	0
56	MG	1a	1834	1/1	0.95	0.20	-	63,63,63,63	0
56	MG	1q	204	1/1	0.94	0.27	-	52,52,52,52	0
56	MG	2A	3503	1/1	0.92	0.08	-	59,59,59,59	0
56	MG	2A	3003	1/1	0.88	0.11	-	53,53,53,53	0
56	MG	1a	1644	1/1	0.91	0.14	-	49,49,49,49	0
56	MG	2A	3514	1/1	0.89	0.13	-	60,60,60,60	0
56	MG	1A	3034	1/1	0.60	0.23	-	73,73,73,73	0
56	MG	1A	3470	1/1	0.83	0.13	-	52,52,52,52	0
56	MG	1a	1745	1/1	0.81	0.19	-	79,79,79,79	0
56	MG	1a	1650	1/1	0.96	0.26	-	57,57,57,57	0
56	MG	2A	3218	1/1	0.82	0.56	-	62,62,62,62	0
56	MG	1A	3421	1/1	0.81	0.14	-	47,47,47,47	0
56	MG	1A	3229	1/1	0.98	0.12	-	50,50,50,50	0
56	MG	1A	3230	1/1	0.83	0.56	-	46,46,46,46	0
56	MG	1a	1655	1/1	0.95	0.11	-	56,56,56,56	0
56	MG	2A	3472	1/1	0.92	0.09	-	39,39,39,39	0
56	MG	1A	3457	1/1	0.82	0.15	-	72,72,72,72	0
56	MG	1A	3068	1/1	0.91	0.70	-	44,44,44,44	0
56	MG	1A	3556	1/1	0.92	0.12	-	59,59,59,59	0
56	MG	2A	3131	1/1	0.80	0.17	-	52,52,52,52	0
56	MG	1a	1736	1/1	0.94	0.22	-	41,41,41,41	0
56	MG	1A	3360	1/1	0.86	0.23	-	47,47,47,47	0
56	MG	2x	3001	1/1	0.89	0.14	-	66,66,66,66	0
56	MG	2A	3044	1/1	0.93	0.75	-	48,48,48,48	0
56	MG	2A	3407	1/1	0.87	0.18	-	40,40,40,40	0
56	MG	1A	3339	1/1	0.90	0.18	-	56,56,56,56	0
56	MG	2A	3276	1/1	0.98	0.15	-	42,42,42,42	0
56	MG	1A	3308	1/1	0.94	0.34	-	50,50,50,50	0
56	MG	1a	1768	1/1	0.96	0.18	-	72,72,72,72	0
56	MG	2a	3041	1/1	0.92	0.28	-	73,73,73,73	0
56	MG	1a	1839	1/1	0.97	0.12	-	56,56,56,56	0
56	MG	2A	3252	1/1	0.89	0.16	-	60,60,60,60	0
56	MG	2A	3491	1/1	0.95	0.04	-	47,47,47,47	0
56	MG	1A	3212	1/1	0.96	0.07	-	39,39,39,39	0
56	MG	2a	3003	1/1	0.93	0.19	-	57,57,57,57	0
56	MG	1A	3783	1/1	0.90	0.18	-	37,37,37,37	0
56	MG	1a	1638	1/1	0.91	0.11	-	63,63,63,63	0
56	MG	2A	3408	1/1	0.92	0.16	-	62,62,62,62	0
56	MG	1A	3039	1/1	0.84	0.30	-	49,49,49,49	0
56	MG	2a	3217	1/1	0.87	0.12	-	83,83,83,83	0
56	MG	1A	3818	1/1	0.87	0.16	-	61,61,61,61	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3702	1/1	0.98	0.14	-	44,44,44,44	0
56	MG	1A	3480	1/1	0.95	0.09	-	56,56,56,56	0
56	MG	1a	1651	1/1	0.84	0.28	-	54,54,54,54	0
56	MG	2A	3282	1/1	0.91	0.11	-	48,48,48,48	0
56	MG	2a	3254	1/1	0.56	0.23	-	71,71,71,71	0
56	MG	2A	3073	1/1	0.90	0.18	-	55,55,55,55	0
56	MG	1a	1622	1/1	0.98	0.17	-	90,90,90,90	0
56	MG	2x	3003	1/1	0.83	0.12	-	73,73,73,73	0
56	MG	1E	304	1/1	0.91	0.17	-	57,57,57,57	0
56	MG	2A	3197	1/1	0.92	0.24	-	57,57,57,57	0
56	MG	1A	3210	1/1	0.94	0.21	-	43,43,43,43	0
56	MG	1A	3673	1/1	0.95	0.26	-	65,65,65,65	0
56	MG	1A	3331	1/1	0.88	0.20	-	57,57,57,57	0
56	MG	1A	3125	1/1	0.89	0.16	-	60,60,60,60	0
56	MG	1A	3715	1/1	0.87	0.15	-	63,63,63,63	0
56	MG	1A	3325	1/1	0.91	0.10	-	74,74,74,74	0
56	MG	1A	3425	1/1	0.90	0.14	-	63,63,63,63	0
56	MG	1A	3171	1/1	0.98	0.25	-	38,38,38,38	0
56	MG	2a	3162	1/1	0.80	0.20	-	75,75,75,75	0
56	MG	1E	301	1/1	0.92	0.14	-	15,15,15,15	0
56	MG	2A	3375	1/1	0.93	0.17	-	48,48,48,48	0
56	MG	1A	3945	1/1	0.96	0.21	-	60,60,60,60	0
56	MG	1A	3666	1/1	0.98	0.12	-	37,37,37,37	0
56	MG	1A	3915	1/1	0.90	0.22	-	58,58,58,58	0
56	MG	1Z	8002	1/1	0.91	0.12	-	67,67,67,67	0
56	MG	1a	1821	1/1	0.85	0.07	-	49,49,49,49	0
56	MG	1A	3314	1/1	0.89	0.30	-	55,55,55,55	0
56	MG	1A	3191	1/1	0.95	0.16	-	37,37,37,37	0
56	MG	1a	1864	1/1	0.90	0.15	-	73,73,73,73	0
56	MG	1A	3515	1/1	0.89	0.37	-	48,48,48,48	0
56	MG	2a	3223	1/1	0.98	0.11	-	49,49,49,49	0
56	MG	1A	3359	1/1	0.86	0.32	-	53,53,53,53	0
56	MG	2A	3108	1/1	0.96	0.12	-	46,46,46,46	0
56	MG	1A	3464	1/1	0.75	1.11	-	76,76,76,76	0
56	MG	2A	3203	1/1	0.89	0.09	-	43,43,43,43	0
56	MG	1A	3982	1/1	0.92	0.32	-	49,49,49,49	0
56	MG	1a	1683	1/1	0.98	0.12	-	68,68,68,68	0
56	MG	1A	3066	1/1	0.82	0.61	-	53,53,53,53	0
56	MG	1A	3815	1/1	0.70	0.13	-	67,67,67,67	0
56	MG	1A	3997	1/1	0.85	0.22	-	50,50,50,50	0
56	MG	2A	3482	1/1	0.96	0.26	-	60,60,60,60	0
56	MG	2a	3262	1/1	0.94	0.09	-	56,56,56,56	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3563	1/1	0.88	0.14	-	46,46,46,46	0
56	MG	1B	217	1/1	0.97	0.18	-	50,50,50,50	0
56	MG	2A	3464	1/1	0.95	0.10	-	50,50,50,50	0
56	MG	1W	3001	1/1	0.95	0.32	-	34,34,34,34	0
56	MG	2a	3152	1/1	0.81	0.15	-	84,84,84,84	0
56	MG	2A	3315	1/1	0.83	0.22	-	55,55,55,55	0
56	MG	2a	3120	1/1	0.94	0.12	-	63,63,63,63	0
56	MG	1A	3409	1/1	0.82	0.12	-	51,51,51,51	0
56	MG	1A	3522	1/1	0.85	0.12	-	69,69,69,69	0
56	MG	1A	3747	1/1	0.91	0.17	-	29,29,29,29	0
56	MG	2A	3066	1/1	0.89	0.18	-	55,55,55,55	0
56	MG	2A	3225	1/1	0.86	0.14	-	63,63,63,63	0
56	MG	2A	3504	1/1	0.93	0.08	-	56,56,56,56	0
56	MG	2A	3313	1/1	0.94	0.09	-	53,53,53,53	0
56	MG	1A	3763	1/1	0.89	0.09	-	43,43,43,43	0
56	MG	1A	3903	1/1	0.94	0.17	-	19,19,19,19	0
56	MG	1A	3800	1/1	0.97	0.10	-	50,50,50,50	0
56	MG	2a	3222	1/1	0.96	0.16	-	46,46,46,46	0
56	MG	1A	3305	1/1	0.90	0.20	-	43,43,43,43	0
56	MG	1A	3096	1/1	0.91	0.36	-	50,50,50,50	0
56	MG	1D	314	1/1	0.96	0.35	-	63,63,63,63	0
56	MG	1A	3641	1/1	0.93	0.11	-	39,39,39,39	0
56	MG	2A	3074	1/1	0.80	0.26	-	57,57,57,57	0
56	MG	1A	3932	1/1	0.91	0.35	-	20,20,20,20	0
56	MG	1a	1843	1/1	0.90	0.08	-	74,74,74,74	0
56	MG	2A	3378	1/1	0.95	0.17	-	66,66,66,66	0
56	MG	2a	3036	1/1	0.82	0.27	-	72,72,72,72	0
56	MG	2A	3105	1/1	0.93	0.27	-	54,54,54,54	0
56	MG	1A	3726	1/1	0.88	0.24	-	64,64,64,64	0
56	MG	2A	3255	1/1	0.88	0.15	-	52,52,52,52	0
56	MG	2a	3123	1/1	0.79	0.15	-	81,81,81,81	0
56	MG	2A	3516	1/1	0.91	0.15	-	59,59,59,59	0
56	MG	2A	3458	1/1	0.98	0.08	-	54,54,54,54	0
56	MG	1A	3431	1/1	0.89	0.29	-	64,64,64,64	0
56	MG	1A	3785	1/1	0.84	0.14	-	72,72,72,72	0
56	MG	2a	3073	1/1	0.65	0.14	-	61,61,61,61	0
56	MG	10	102	1/1	0.94	0.14	-	60,60,60,60	0
56	MG	2A	3374	1/1	0.96	0.25	-	37,37,37,37	0
56	MG	2a	3168	1/1	0.91	0.08	-	71,71,71,71	0
56	MG	1A	3709	1/1	0.78	0.20	-	45,45,45,45	0
56	MG	1A	3311	1/1	0.89	0.17	-	65,65,65,65	0
56	MG	2a	3002	1/1	0.94	0.24	-	65,65,65,65	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	2A	3043	1/1	0.99	0.19	-	49,49,49,49	0
56	MG	1A	3371	1/1	0.90	0.15	-	56,56,56,56	0
56	MG	2a	3025	1/1	0.90	0.10	-	60,60,60,60	0
56	MG	1a	1757	1/1	0.77	0.14	-	63,63,63,63	0
56	MG	2a	3202	1/1	0.92	0.18	-	72,72,72,72	0
56	MG	2A	3195	1/1	0.71	0.15	-	48,48,48,48	0
56	MG	2A	3330	1/1	0.97	0.23	-	47,47,47,47	0
56	MG	2a	3070	1/1	0.71	0.14	-	69,69,69,69	0
56	MG	1A	3407	1/1	0.93	0.10	-	41,41,41,41	0
56	MG	1A	3091	1/1	0.96	0.11	-	51,51,51,51	0
56	MG	1A	3142	1/1	0.87	0.61	-	57,57,57,57	0
56	MG	1A	3760	1/1	0.97	0.17	-	36,36,36,36	0
56	MG	1a	1624	1/1	0.89	0.60	-	62,62,62,62	0
56	MG	1R	201	1/1	0.95	0.27	-	63,63,63,63	0
56	MG	1A	3201	1/1	0.96	0.15	-	49,49,49,49	0
56	MG	2A	3436	1/1	0.91	0.16	-	58,58,58,58	0
56	MG	2A	3092	1/1	0.80	0.12	-	54,54,54,54	0
56	MG	1A	3165	1/1	0.73	0.25	-	53,53,53,53	0
56	MG	1A	3492	1/1	0.96	0.57	-	52,52,52,52	0
56	MG	1a	1825	1/1	0.97	0.07	-	43,43,43,43	0
56	MG	1A	3707	1/1	0.96	0.20	-	70,70,70,70	0
56	MG	1a	1769	1/1	0.89	0.09	-	87,87,87,87	0
56	MG	1A	3235	1/1	0.87	0.16	-	46,46,46,46	0
56	MG	1A	3306	1/1	0.86	0.26	-	41,41,41,41	0
56	MG	2a	3207	1/1	0.85	0.15	-	73,73,73,73	0
56	MG	1T	204	1/1	0.89	0.19	-	67,67,67,67	0
56	MG	1A	3272	1/1	0.98	0.22	-	29,29,29,29	0
56	MG	2A	3389	1/1	0.81	0.24	-	77,77,77,77	0
56	MG	25	502	1/1	0.95	0.29	-	47,47,47,47	0
56	MG	1A	3218	1/1	0.97	0.16	-	21,21,21,21	0
56	MG	1A	3104	1/1	0.90	0.09	-	55,55,55,55	0
56	MG	2a	3248	1/1	0.91	0.15	-	65,65,65,65	0
56	MG	2a	3229	1/1	0.89	0.18	-	51,51,51,51	0
56	MG	1A	3468	1/1	0.86	0.24	-	57,57,57,57	0
56	MG	1A	3919	1/1	0.95	0.30	-	49,49,49,49	0
56	MG	2A	3141	1/1	0.79	0.15	-	74,74,74,74	0
56	MG	2A	3075	1/1	0.93	0.14	-	59,59,59,59	0
56	MG	1a	1804	1/1	0.94	0.11	-	52,52,52,52	0
56	MG	1A	3674	1/1	0.96	0.15	-	52,52,52,52	0
56	MG	1a	1627	1/1	0.72	0.14	-	67,67,67,67	0
56	MG	1A	3828	1/1	0.98	0.08	-	67,67,67,67	0
56	MG	2A	3172	1/1	0.94	0.27	-	49,49,49,49	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3160	1/1	0.93	0.12	-	40,40,40,40	0
56	MG	1A	3970	1/1	0.97	1.04	-	30,30,30,30	0
56	MG	1A	3724	1/1	0.94	0.20	-	58,58,58,58	0
56	MG	1A	3605	1/1	0.86	0.20	-	33,33,33,33	0
56	MG	1A	3337	1/1	0.91	0.32	-	57,57,57,57	0
56	MG	1A	3869	1/1	0.92	0.09	-	42,42,42,42	0
56	MG	1A	3419	1/1	0.93	0.20	-	56,56,56,56	0
56	MG	2A	3399	1/1	0.96	0.12	-	31,31,31,31	0
56	MG	2A	3284	1/1	0.91	0.10	-	62,62,62,62	0
56	MG	2a	3180	1/1	0.81	0.19	-	61,61,61,61	0
56	MG	1A	3951	1/1	0.93	0.31	-	70,70,70,70	0
56	MG	1A	3784	1/1	0.93	0.21	-	33,33,33,33	0
56	MG	1A	3013	1/1	0.80	0.26	-	70,70,70,70	0
56	MG	2E	303	1/1	0.96	0.19	-	26,26,26,26	0
56	MG	2A	3331	1/1	0.97	0.16	-	48,48,48,48	0
56	MG	2A	3509	1/1	0.78	0.19	-	62,62,62,62	0
56	MG	1A	3890	1/1	0.87	0.21	-	37,37,37,37	0
56	MG	1A	3863	1/1	0.90	0.21	-	78,78,78,78	0
56	MG	2A	3046	1/1	0.94	0.26	-	64,64,64,64	0
56	MG	1A	3877	1/1	0.93	0.13	-	42,42,42,42	0
56	MG	1A	3857	1/1	0.80	0.21	-	32,32,32,32	0
56	MG	2A	3009	1/1	0.85	0.41	-	60,60,60,60	0
56	MG	2A	3028	1/1	0.89	0.14	-	52,52,52,52	0
56	MG	1a	1652	1/1	0.98	0.20	-	48,48,48,48	0
56	MG	2a	3145	1/1	0.91	0.10	-	61,61,61,61	0
56	MG	16	101	1/1	0.95	0.17	-	45,45,45,45	0
56	MG	1A	3941	1/1	0.89	0.12	-	65,65,65,65	0
56	MG	2A	3122	1/1	0.71	0.17	-	66,66,66,66	0
56	MG	1A	3769	1/1	0.83	0.20	-	90,90,90,90	0
56	MG	1A	3934	1/1	0.97	0.12	-	33,33,33,33	0
56	MG	1A	3920	1/1	0.94	0.15	-	64,64,64,64	0
56	MG	1A	3899	1/1	0.96	0.24	-	47,47,47,47	0
56	MG	2A	3497	1/1	0.85	0.18	-	54,54,54,54	0
56	MG	1A	3956	1/1	0.93	0.18	-	53,53,53,53	0
56	MG	2A	3525	1/1	0.91	0.15	-	63,63,63,63	0
56	MG	1A	3983	1/1	0.91	0.18	-	55,55,55,55	0
56	MG	2a	3138	1/1	0.79	0.13	-	81,81,81,81	0
56	MG	1a	1779	1/1	0.95	0.11	-	82,82,82,82	0
56	MG	2a	3173	1/1	0.78	0.56	-	74,74,74,74	0
56	MG	1A	3823	1/1	0.94	0.29	-	43,43,43,43	0
56	MG	1A	3998	1/1	0.89	0.27	-	75,75,75,75	0
56	MG	1x	103	1/1	0.96	0.13	-	70,70,70,70	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1q	201	1/1	0.90	0.18	-	51,51,51,51	0
56	MG	2a	3042	1/1	0.95	0.14	-	56,56,56,56	0
56	MG	2a	3233	1/1	0.97	0.10	-	61,61,61,61	0
56	MG	2A	3247	1/1	0.91	0.54	-	59,59,59,59	0
56	MG	1a	1801	1/1	0.92	0.19	-	59,59,59,59	0
56	MG	1a	1688	1/1	0.92	0.11	-	56,56,56,56	0
56	MG	2A	3332	1/1	0.78	0.13	-	38,38,38,38	0
56	MG	1A	3376	1/1	0.96	0.19	-	53,53,53,53	0
56	MG	2a	3161	1/1	0.68	0.12	-	102,102,102,102	0
56	MG	1A	3153	1/1	0.96	0.11	-	24,24,24,24	0
56	MG	2a	3012	1/1	0.91	0.13	-	66,66,66,66	0
56	MG	1A	3257	1/1	0.92	0.17	-	44,44,44,44	0
56	MG	1A	3266	1/1	0.93	0.10	-	44,44,44,44	0
56	MG	1A	3918	1/1	0.85	0.40	-	71,71,71,71	0
56	MG	2a	3031	1/1	0.53	0.17	-	65,65,65,65	0
56	MG	1A	3361	1/1	0.84	0.19	-	66,66,66,66	0
56	MG	1A	3717	1/1	0.97	0.17	-	59,59,59,59	0
56	MG	1A	3313	1/1	0.96	0.37	-	53,53,53,53	0
56	MG	1A	3413	1/1	0.94	0.13	-	41,41,41,41	0
56	MG	2A	3134	1/1	0.93	0.07	-	51,51,51,51	0
56	MG	1A	3219	1/1	0.86	0.27	-	59,59,59,59	0
56	MG	2A	3537	1/1	0.97	0.16	-	37,37,37,37	0
56	MG	2a	3209	1/1	0.95	0.14	-	70,70,70,70	0
56	MG	1A	3677	1/1	0.98	0.15	-	51,51,51,51	0
56	MG	1x	105	1/1	0.93	0.21	-	57,57,57,57	0
56	MG	1A	3240	1/1	0.94	0.16	-	49,49,49,49	0
56	MG	1A	3599	1/1	0.94	0.23	-	37,37,37,37	0
56	MG	2a	3014	1/1	0.52	0.12	-	65,65,65,65	0
56	MG	1A	3728	1/1	0.97	0.20	-	52,52,52,52	0
56	MG	1A	3735	1/1	0.93	0.16	-	36,36,36,36	0
56	MG	1a	1780	1/1	0.94	0.29	-	68,68,68,68	0
56	MG	1A	3080	1/1	0.95	0.31	-	50,50,50,50	0
56	MG	2A	3418	1/1	0.90	0.13	-	57,57,57,57	0
56	MG	1A	3236	1/1	0.89	0.45	-	46,46,46,46	0
56	MG	1A	3319	1/1	0.96	0.10	-	57,57,57,57	0
56	MG	2A	3352	1/1	0.70	0.17	-	68,68,68,68	0
56	MG	1a	1845	1/1	0.97	0.11	-	46,46,46,46	0
56	MG	1A	3520	1/1	0.62	0.14	-	75,75,75,75	0
56	MG	1A	3152	1/1	0.93	0.14	-	50,50,50,50	0
56	MG	1N	3006	1/1	0.82	0.64	-	71,71,71,71	0
56	MG	2A	3207	1/1	0.91	0.12	-	44,44,44,44	0
56	MG	10	105	1/1	0.75	0.17	-	53,53,53,53	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	2A	3538	1/1	0.95	0.19	-	22,22,22,22	0
56	MG	12	3001	1/1	0.70	0.42	-	61,61,61,61	0
56	MG	2A	3265	1/1	0.93	0.19	-	37,37,37,37	0
56	MG	1A	3552	1/1	0.69	0.22	-	50,50,50,50	0
56	MG	1A	3452	1/1	0.93	0.13	-	56,56,56,56	0
56	MG	1A	3721	1/1	0.90	0.18	-	55,55,55,55	0
56	MG	1A	3598	1/1	0.89	0.16	-	51,51,51,51	0
56	MG	1A	3102	1/1	0.88	0.13	-	60,60,60,60	0
56	MG	2a	3075	1/1	0.66	0.15	-	72,72,72,72	0
56	MG	1A	3713	1/1	0.95	0.14	-	40,40,40,40	0
56	MG	1a	1796	1/1	0.90	0.38	-	86,86,86,86	0
56	MG	1A	3638	1/1	0.95	0.29	-	41,41,41,41	0
56	MG	2A	3513	1/1	0.82	0.14	-	61,61,61,61	0
56	MG	2A	3520	1/1	0.97	0.08	-	42,42,42,42	0
56	MG	2A	3517	1/1	0.97	0.15	-	56,56,56,56	0
56	MG	1A	3173	1/1	0.95	0.19	-	62,62,62,62	0
56	MG	2Y	502	1/1	0.94	0.21	-	52,52,52,52	0
56	MG	16	103	1/1	0.96	0.10	-	68,68,68,68	0
56	MG	1A	3400	1/1	0.88	0.15	-	45,45,45,45	0
56	MG	1a	1732	1/1	0.67	0.24	-	74,74,74,74	0
56	MG	1a	1680	1/1	0.97	0.14	-	53,53,53,53	0
56	MG	1a	1835	1/1	0.98	0.07	-	73,73,73,73	0
56	MG	2a	3192	1/1	0.90	0.10	-	60,60,60,60	0
56	MG	2a	3114	1/1	0.93	0.23	-	53,53,53,53	0
56	MG	1A	3105	1/1	0.83	0.22	-	61,61,61,61	0
56	MG	1P	203	1/1	0.96	0.11	-	65,65,65,65	0
56	MG	1T	203	1/1	0.70	0.18	-	61,61,61,61	0
56	MG	1A	3466	1/1	0.92	0.09	-	40,40,40,40	0
56	MG	1A	3870	1/1	0.96	0.12	-	53,53,53,53	0
56	MG	1A	3850	1/1	0.94	0.15	-	41,41,41,41	0
56	MG	1A	3844	1/1	0.94	0.13	-	71,71,71,71	0
56	MG	1A	3007	1/1	0.87	0.28	-	56,56,56,56	0
56	MG	1A	3568	1/1	0.93	0.21	-	42,42,42,42	0
56	MG	1A	3586	1/1	0.94	0.14	-	57,57,57,57	0
56	MG	1A	3580	1/1	0.97	0.22	-	46,46,46,46	0
56	MG	1a	1610	1/1	0.89	0.13	-	53,53,53,53	0
56	MG	1A	3856	1/1	0.95	0.08	-	69,69,69,69	0
56	MG	2A	3136	1/1	0.96	0.09	-	54,54,54,54	0
56	MG	2A	3322	1/1	0.92	0.14	-	67,67,67,67	0
56	MG	1A	3649	1/1	0.92	0.14	-	71,71,71,71	0
56	MG	1a	1782	1/1	0.93	0.18	-	43,43,43,43	0
56	MG	2A	3241	1/1	0.84	0.17	-	65,65,65,65	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3519	1/1	0.86	0.31	-	62,62,62,62	0
56	MG	2a	3117	1/1	0.90	0.10	-	58,58,58,58	0
56	MG	2x	3002	1/1	0.84	0.13	-	65,65,65,65	0
56	MG	1a	1817	1/1	0.96	0.30	-	52,52,52,52	0
56	MG	2A	3384	1/1	0.95	0.06	-	45,45,45,45	0
56	MG	2a	3218	1/1	0.97	0.14	-	62,62,62,62	0
56	MG	1A	3256	1/1	0.98	0.17	-	43,43,43,43	0
56	MG	2a	3188	1/1	0.94	0.11	-	60,60,60,60	0
56	MG	1Q	3003	1/1	0.77	0.77	-	51,51,51,51	0
56	MG	2a	3021	1/1	0.94	0.48	-	60,60,60,60	0
56	MG	2A	3346	1/1	0.89	0.18	-	62,62,62,62	0
56	MG	1A	3342	1/1	0.95	0.36	-	52,52,52,52	0
56	MG	1A	3682	1/1	0.87	0.14	-	51,51,51,51	0
56	MG	1A	3326	1/1	0.96	0.09	-	53,53,53,53	0
56	MG	1A	3540	1/1	0.98	0.32	-	45,45,45,45	0
56	MG	1a	1693	1/1	0.96	0.14	-	52,52,52,52	0
56	MG	1r	3002	1/1	0.83	0.22	-	55,55,55,55	0
56	MG	1a	1714	1/1	0.98	0.09	-	68,68,68,68	0
56	MG	2A	3151	1/1	0.96	0.27	-	47,47,47,47	0
56	MG	1A	3251	1/1	0.90	0.20	-	30,30,30,30	0
56	MG	2A	3387	1/1	0.96	0.21	-	63,63,63,63	0
56	MG	1A	3926	1/1	0.99	0.31	-	38,38,38,38	0
56	MG	1A	3078	1/1	0.94	0.07	-	62,62,62,62	0
56	MG	1a	1773	1/1	0.81	0.21	-	77,77,77,77	0
56	MG	1A	3243	1/1	0.85	0.86	-	46,46,46,46	0
56	MG	1A	3992	1/1	0.87	0.16	-	55,55,55,55	0
56	MG	1A	3655	1/1	0.91	0.29	-	48,48,48,48	0
56	MG	1A	3073	1/1	0.96	0.24	-	51,51,51,51	0
56	MG	1A	3778	1/1	0.80	0.20	-	23,23,23,23	0
56	MG	1A	3354	1/1	0.93	0.09	-	43,43,43,43	0
56	MG	1A	3119	1/1	0.80	0.37	-	65,65,65,65	0
56	MG	2A	3137	1/1	0.96	0.12	-	51,51,51,51	0
56	MG	1A	3991	1/1	0.90	0.15	-	52,52,52,52	0
56	MG	2A	3316	1/1	0.98	0.21	-	57,57,57,57	0
56	MG	2A	3541	1/1	0.96	0.33	-	42,42,42,42	0
56	MG	1a	1717	1/1	0.81	0.10	-	63,63,63,63	0
56	MG	1a	1670	1/1	0.94	0.18	-	43,43,43,43	0
56	MG	2A	3157	1/1	0.90	0.17	-	43,43,43,43	0
56	MG	1A	3601	1/1	0.97	0.22	-	44,44,44,44	0
56	MG	1A	3517	1/1	0.96	0.65	-	43,43,43,43	0
56	MG	1a	1631	1/1	0.97	0.12	-	36,36,36,36	0
56	MG	2a	3169	1/1	0.96	0.69	-	99,99,99,99	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	2A	3455	1/1	0.31	0.27	-	69,69,69,69	0
56	MG	1A	3489	1/1	0.84	0.11	-	51,51,51,51	0
56	MG	1a	1737	1/1	0.92	0.07	-	57,57,57,57	0
56	MG	2a	3252	1/1	0.96	0.12	-	70,70,70,70	0
56	MG	1A	3766	1/1	0.91	0.19	-	49,49,49,49	0
56	MG	2A	3371	1/1	0.84	0.13	-	56,56,56,56	0
56	MG	18	102	1/1	0.84	0.29	-	57,57,57,57	0
56	MG	1Z	8001	1/1	0.90	0.22	-	69,69,69,69	0
56	MG	2A	3453	1/1	0.90	0.15	-	61,61,61,61	0
56	MG	1A	3420	1/1	0.83	0.11	-	56,56,56,56	0
56	MG	1A	3009	1/1	0.92	0.28	-	56,56,56,56	0
56	MG	2a	3040	1/1	0.94	0.15	-	39,39,39,39	0
56	MG	10	104	1/1	0.84	0.22	-	43,43,43,43	0
56	MG	1A	3901	1/1	0.91	0.12	-	44,44,44,44	0
56	MG	1a	1692	1/1	0.95	0.09	-	57,57,57,57	0
56	MG	2A	3410	1/1	0.89	0.21	-	56,56,56,56	0
56	MG	1A	3851	1/1	0.95	0.17	-	62,62,62,62	0
56	MG	1a	1763	1/1	0.92	0.36	-	69,69,69,69	0
56	MG	1A	3403	1/1	0.85	0.08	-	59,59,59,59	0
56	MG	1A	3558	1/1	0.96	0.10	-	61,61,61,61	0
56	MG	1A	3772	1/1	0.91	0.12	-	60,60,60,60	0
56	MG	1a	1812	1/1	0.94	0.22	-	49,49,49,49	0
56	MG	2A	3179	1/1	0.88	0.13	-	46,46,46,46	0
56	MG	2A	3447	1/1	0.91	0.12	-	62,62,62,62	0
56	MG	2A	3391	1/1	0.97	0.18	-	48,48,48,48	0
56	MG	1A	3958	1/1	0.87	0.16	-	64,64,64,64	0
56	MG	2a	3083	1/1	0.92	0.26	-	51,51,51,51	0
56	MG	2a	3063	1/1	0.89	0.11	-	77,77,77,77	0
56	MG	1a	1634	1/1	0.92	0.10	-	52,52,52,52	0
56	MG	1A	3107	1/1	0.93	0.27	-	58,58,58,58	0
56	MG	1A	3948	1/1	0.97	0.15	-	28,28,28,28	0
56	MG	2A	3348	1/1	0.96	0.17	-	52,52,52,52	0
56	MG	1A	3461	1/1	0.88	0.25	-	71,71,71,71	0
56	MG	1A	3463	1/1	0.97	0.08	-	80,80,80,80	0
56	MG	2A	3328	1/1	0.84	0.19	-	57,57,57,57	0
56	MG	2A	3025	1/1	0.91	0.45	-	67,67,67,67	0
56	MG	1A	3224	1/1	0.93	0.18	-	35,35,35,35	0
56	MG	2A	3452	1/1	0.85	0.16	-	69,69,69,69	0
56	MG	2A	3111	1/1	0.94	0.28	-	68,68,68,68	0
56	MG	1T	201	1/1	0.95	0.19	-	41,41,41,41	0
56	MG	2a	3124	1/1	0.94	0.15	-	63,63,63,63	0
56	MG	2A	3492	1/1	0.92	0.08	-	44,44,44,44	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	19	104	1/1	0.98	0.13	-	31,31,31,31	0
56	MG	2A	3298	1/1	0.87	0.13	-	49,49,49,49	0
56	MG	2a	3198	1/1	0.90	0.06	-	63,63,63,63	0
56	MG	1D	302	1/1	0.91	0.70	-	65,65,65,65	0
56	MG	1A	3952	1/1	0.94	0.15	-	51,51,51,51	0
56	MG	1a	1805	1/1	0.94	0.19	-	67,67,67,67	0
56	MG	1a	1640	1/1	0.86	0.20	-	51,51,51,51	0
56	MG	1A	3384	1/1	0.90	0.13	-	69,69,69,69	0
56	MG	1A	3069	1/1	0.96	0.28	-	39,39,39,39	0
56	MG	1a	1649	1/1	0.84	0.37	-	63,63,63,63	0
56	MG	1A	3827	1/1	0.94	0.13	-	31,31,31,31	0
56	MG	1A	3630	1/1	0.97	0.14	-	32,32,32,32	0
56	MG	1A	3635	1/1	0.98	0.11	-	46,46,46,46	0
56	MG	2a	3107	1/1	0.97	0.07	-	57,57,57,57	0
56	MG	2A	3072	1/1	0.91	0.17	-	48,48,48,48	0
56	MG	1A	3847	1/1	0.89	0.17	-	46,46,46,46	0
56	MG	2A	3132	1/1	0.93	0.26	-	53,53,53,53	0
56	MG	1A	3940	1/1	0.92	0.17	-	63,63,63,63	0
56	MG	1A	3816	1/1	0.95	0.18	-	68,68,68,68	0
56	MG	2A	3522	1/1	0.87	0.11	-	70,70,70,70	0
56	MG	2A	3128	1/1	0.90	0.17	-	60,60,60,60	0
56	MG	1a	1744	1/1	0.89	0.17	-	61,61,61,61	0
56	MG	2a	3081	1/1	0.94	0.16	-	52,52,52,52	0
56	MG	2a	3241	1/1	0.77	0.42	-	95,95,95,95	0
56	MG	1A	4005	1/1	0.93	0.12	-	58,58,58,58	0
56	MG	1A	3204	1/1	0.96	0.26	-	37,37,37,37	0
56	MG	1A	3324	1/1	0.90	0.24	-	57,57,57,57	0
56	MG	1a	1842	1/1	0.94	0.21	-	73,73,73,73	0
56	MG	1A	3086	1/1	0.96	0.53	-	44,44,44,44	0
56	MG	1a	1625	1/1	0.97	0.07	-	52,52,52,52	0
56	MG	2A	3308	1/1	0.92	0.16	-	40,40,40,40	0
56	MG	2A	3551	1/1	0.72	0.23	-	57,57,57,57	0
56	MG	1A	3052	1/1	0.86	0.13	-	59,59,59,59	0
56	MG	2a	3102	1/1	0.93	0.24	-	47,47,47,47	0
56	MG	1a	1735	1/1	0.96	0.13	-	66,66,66,66	0
56	MG	2A	3529	1/1	0.95	0.10	-	68,68,68,68	0
56	MG	2A	3146	1/1	0.95	0.14	-	64,64,64,64	0
56	MG	1A	3985	1/1	0.88	0.17	-	66,66,66,66	0
56	MG	1A	3443	1/1	0.93	0.47	-	48,48,48,48	0
56	MG	1A	3663	1/1	0.96	0.12	-	55,55,55,55	0
56	MG	1A	3351	1/1	0.93	0.18	-	62,62,62,62	0
56	MG	1A	3927	1/1	0.90	0.18	-	50,50,50,50	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	2A	3402	1/1	0.89	0.40	-	62,62,62,62	0
56	MG	1A	3099	1/1	0.89	0.12	-	39,39,39,39	0
56	MG	1a	1833	1/1	0.97	0.13	-	59,59,59,59	0
56	MG	2a	3092	1/1	0.71	0.18	-	87,87,87,87	0
56	MG	2A	3193	1/1	0.95	0.11	-	32,32,32,32	0
56	MG	2a	3121	1/1	0.89	0.19	-	63,63,63,63	0
56	MG	1A	3364	1/1	0.94	0.15	-	44,44,44,44	0
56	MG	1B	201	1/1	0.87	0.10	-	50,50,50,50	0
56	MG	2a	3237	1/1	0.96	0.07	-	52,52,52,52	0
56	MG	1a	1708	1/1	0.91	0.12	-	61,61,61,61	0
56	MG	1A	3043	1/1	0.74	0.37	-	52,52,52,52	0
56	MG	11	4000	1/1	0.91	0.09	-	43,43,43,43	0
56	MG	1A	3786	1/1	0.92	0.15	-	57,57,57,57	0
56	MG	2a	3176	1/1	0.88	0.17	-	56,56,56,56	0
56	MG	1A	3004	1/1	0.92	0.16	-	40,40,40,40	0
56	MG	2A	3432	1/1	0.97	0.16	-	61,61,61,61	0
56	MG	2A	3523	1/1	0.86	0.39	-	56,56,56,56	0
56	MG	1A	3824	1/1	0.95	0.18	-	33,33,33,33	0
56	MG	1A	3852	1/1	0.94	0.17	-	16,16,16,16	0
56	MG	2A	3182	1/1	0.91	0.28	-	52,52,52,52	0
56	MG	1A	3884	1/1	0.75	0.14	-	53,53,53,53	0
56	MG	2A	3327	1/1	0.96	0.19	-	36,36,36,36	0
56	MG	1a	1850	1/1	0.95	0.10	-	59,59,59,59	0
56	MG	1A	3310	1/1	0.98	0.30	-	44,44,44,44	0
56	MG	2A	3198	1/1	0.87	0.18	-	41,41,41,41	0
56	MG	1A	3603	1/1	0.90	0.10	-	60,60,60,60	0
56	MG	2A	3393	1/1	0.94	0.13	-	63,63,63,63	0
56	MG	1A	3111	1/1	0.75	0.22	-	68,68,68,68	0
56	MG	1a	1783	1/1	0.81	0.16	-	57,57,57,57	0
56	MG	1a	1766	1/1	0.91	0.28	-	60,60,60,60	0
56	MG	2A	3088	1/1	0.88	0.17	-	51,51,51,51	0
56	MG	1A	3405	1/1	0.83	0.14	-	54,54,54,54	0
56	MG	1R	204	1/1	0.97	0.20	-	36,36,36,36	0
56	MG	1A	3304	1/1	0.97	0.17	-	34,34,34,34	0
56	MG	2a	3016	1/1	0.87	0.16	-	65,65,65,65	0
56	MG	1a	1690	1/1	0.83	0.16	-	63,63,63,63	0
56	MG	1A	3040	1/1	0.91	0.39	-	45,45,45,45	0
56	MG	1A	3708	1/1	0.95	0.18	-	63,63,63,63	0
56	MG	2A	3156	1/1	0.83	0.17	-	49,49,49,49	0
56	MG	1A	3984	1/1	0.87	0.12	-	47,47,47,47	0
56	MG	1a	1855	1/1	0.93	0.10	-	55,55,55,55	0
56	MG	1A	3334	1/1	0.92	0.17	-	72,72,72,72	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1a	1620	1/1	0.91	0.09	-	58,58,58,58	0
56	MG	1A	3006	1/1	0.90	0.24	-	57,57,57,57	0
56	MG	1A	3748	1/1	0.94	0.13	-	37,37,37,37	0
56	MG	1A	3386	1/1	0.93	0.28	-	63,63,63,63	0
56	MG	2A	3056	1/1	0.88	0.13	-	38,38,38,38	0
56	MG	2A	3356	1/1	0.88	0.13	-	60,60,60,60	0
56	MG	1A	3972	1/1	0.97	0.66	-	42,42,42,42	0
56	MG	1A	3570	1/1	0.78	0.18	-	66,66,66,66	0
56	MG	1a	1706	1/1	0.82	0.22	-	64,64,64,64	0
56	MG	1A	3395	1/1	0.97	0.50	-	37,37,37,37	0
56	MG	2A	3250	1/1	0.74	0.16	-	64,64,64,64	0
56	MG	2A	3006	1/1	0.82	0.10	-	64,64,64,64	0
56	MG	1a	1861	1/1	0.93	0.17	-	58,58,58,58	0
56	MG	1A	3146	1/1	0.86	0.18	-	65,65,65,65	0
56	MG	2A	3415	1/1	0.93	0.16	-	60,60,60,60	0
56	MG	2A	3454	1/1	0.96	0.12	-	56,56,56,56	0
56	MG	1a	1637	1/1	0.91	0.31	-	53,53,53,53	0
56	MG	1A	3656	1/1	0.98	0.20	-	33,33,33,33	0
56	MG	1b	3002	1/1	0.82	0.12	-	76,76,76,76	0
56	MG	17	101	1/1	0.87	0.28	-	57,57,57,57	0
56	MG	1A	3495	1/1	0.93	0.11	-	66,66,66,66	0
56	MG	1A	3854	1/1	0.89	0.26	-	53,53,53,53	0
56	MG	2A	3081	1/1	0.94	0.08	-	44,44,44,44	0
56	MG	1A	3686	1/1	0.94	0.13	-	46,46,46,46	0
56	MG	1A	3808	1/1	0.94	0.21	-	51,51,51,51	0
56	MG	1x	107	1/1	0.86	0.38	-	61,61,61,61	0
56	MG	1A	3714	1/1	0.98	0.21	-	46,46,46,46	0
56	MG	2l	203	1/1	0.83	0.22	-	60,60,60,60	0
56	MG	1A	3447	1/1	0.93	0.15	-	63,63,63,63	0
56	MG	1A	3137	1/1	0.91	0.34	-	51,51,51,51	0
56	MG	2a	3088	1/1	0.96	0.22	-	54,54,54,54	0
56	MG	2A	3040	1/1	0.89	0.15	-	53,53,53,53	0
56	MG	2A	3495	1/1	0.89	0.12	-	71,71,71,71	0
56	MG	1A	3242	1/1	0.85	0.12	-	33,33,33,33	0
56	MG	1A	3261	1/1	0.79	0.34	-	62,62,62,62	0
56	MG	1A	3042	1/1	0.93	0.83	-	44,44,44,44	0
56	MG	1A	3801	1/1	0.94	0.15	-	42,42,42,42	0
56	MG	2a	3221	1/1	0.83	0.12	-	66,66,66,66	0
56	MG	1A	3939	1/1	0.82	0.40	-	69,69,69,69	0
56	MG	1A	3706	1/1	0.94	0.13	-	22,22,22,22	0
56	MG	1A	3697	1/1	0.88	0.14	-	58,58,58,58	0
56	MG	2a	3227	1/1	0.94	0.19	-	66,66,66,66	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3737	1/1	0.92	0.19	-	24,24,24,24	0
56	MG	1A	3516	1/1	0.83	0.27	-	53,53,53,53	0
56	MG	1A	3297	1/1	0.87	0.31	-	70,70,70,70	0
56	MG	2A	3449	1/1	0.86	0.12	-	60,60,60,60	0
56	MG	2a	3236	1/1	0.93	0.14	-	62,62,62,62	0
56	MG	2A	3023	1/1	0.89	0.18	-	61,61,61,61	0
56	MG	1A	3916	1/1	0.98	0.17	-	52,52,52,52	0
56	MG	2a	3190	1/1	0.94	0.20	-	52,52,52,52	0
56	MG	1a	1657	1/1	0.89	0.10	-	71,71,71,71	0
56	MG	1A	3143	1/1	0.95	0.26	-	52,52,52,52	0
56	MG	1A	3417	1/1	0.63	0.48	-	58,58,58,58	0
56	MG	1A	3366	1/1	0.96	0.09	-	46,46,46,46	0
56	MG	1a	1611	1/1	0.88	0.10	-	78,78,78,78	0
56	MG	1A	4000	1/1	0.90	0.14	-	61,61,61,61	0
56	MG	2A	3425	1/1	0.94	0.19	-	47,47,47,47	0
56	MG	2a	3038	1/1	0.82	0.17	-	57,57,57,57	0
56	MG	1B	224	1/1	0.96	0.12	-	44,44,44,44	0
56	MG	1a	1767	1/1	0.95	0.14	-	64,64,64,64	0
56	MG	1a	1704	1/1	0.82	0.18	-	54,54,54,54	0
56	MG	1A	3512	1/1	0.91	0.28	-	60,60,60,60	0
56	MG	2A	3498	1/1	0.93	0.13	-	49,49,49,49	0
56	MG	1a	1838	1/1	0.92	0.33	-	66,66,66,66	0
56	MG	1A	3880	1/1	0.97	0.14	-	55,55,55,55	0
56	MG	1A	3112	1/1	0.95	0.27	-	40,40,40,40	0
56	MG	1a	1636	1/1	0.97	0.20	-	44,44,44,44	0
56	MG	1e	3003	1/1	0.92	0.21	-	68,68,68,68	0
56	MG	1A	3127	1/1	0.93	0.50	-	53,53,53,53	0
56	MG	1A	3917	1/1	0.94	0.15	-	31,31,31,31	0
56	MG	2A	3038	1/1	0.83	0.11	-	58,58,58,58	0
56	MG	1a	1725	1/1	0.87	0.17	-	59,59,59,59	0
56	MG	1A	3260	1/1	0.95	0.22	-	58,58,58,58	0
56	MG	1x	101	1/1	0.97	0.30	-	51,51,51,51	0
56	MG	1A	3422	1/1	0.97	0.09	-	51,51,51,51	0
56	MG	1A	3526	1/1	0.85	0.21	-	32,32,32,32	0
56	MG	2A	3143	1/1	0.88	0.11	-	48,48,48,48	0
56	MG	2A	3183	1/1	0.92	0.14	-	56,56,56,56	0
56	MG	1A	3867	1/1	0.88	0.21	-	30,30,30,30	0
56	MG	1A	3415	1/1	0.88	0.11	-	50,50,50,50	0
56	MG	1a	1754	1/1	0.88	0.10	-	81,81,81,81	0
56	MG	1A	3084	1/1	0.93	0.12	-	50,50,50,50	0
56	MG	1A	3438	1/1	0.97	0.29	-	58,58,58,58	0
56	MG	2a	3201	1/1	0.83	0.39	-	88,88,88,88	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1a	1827	1/1	0.91	0.10	-	71,71,71,71	0
56	MG	1A	3886	1/1	0.93	0.35	-	33,33,33,33	0
56	MG	1a	1748	1/1	0.87	0.20	-	52,52,52,52	0
56	MG	1A	3725	1/1	0.87	0.18	-	42,42,42,42	0
56	MG	1O	3001	1/1	0.88	0.26	-	45,45,45,45	0
56	MG	2a	3054	1/1	0.92	0.07	-	89,89,89,89	0
56	MG	1a	1793	1/1	0.92	0.09	-	77,77,77,77	0
56	MG	1A	3881	1/1	0.85	0.10	-	55,55,55,55	0
56	MG	2A	3231	1/1	0.93	0.20	-	45,45,45,45	0
56	MG	1A	3130	1/1	0.85	0.36	-	58,58,58,58	0
56	MG	2A	3451	1/1	0.97	0.17	-	61,61,61,61	0
56	MG	2A	3277	1/1	0.94	0.24	-	41,41,41,41	0
56	MG	1B	202	1/1	0.93	0.19	-	52,52,52,52	0
56	MG	2a	3200	1/1	0.96	0.11	-	49,49,49,49	0
56	MG	1A	3180	1/1	0.94	0.47	-	33,33,33,33	0
56	MG	2a	3020	1/1	0.85	0.16	-	49,49,49,49	0
56	MG	2A	3259	1/1	0.96	0.15	-	72,72,72,72	0
56	MG	2A	3064	1/1	0.94	0.23	-	56,56,56,56	0
56	MG	1a	1764	1/1	0.91	0.44	-	68,68,68,68	0
56	MG	1A	3163	1/1	0.83	0.14	-	32,32,32,32	0
56	MG	1a	1612	1/1	0.92	0.09	-	65,65,65,65	0
56	MG	1a	1788	1/1	0.91	0.64	-	76,76,76,76	0
56	MG	1A	3593	1/1	0.96	0.22	-	35,35,35,35	0
56	MG	2A	3441	1/1	0.93	0.08	-	58,58,58,58	0
56	MG	1A	4013	1/1	0.89	0.23	-	49,49,49,49	0
56	MG	2A	3018	1/1	0.87	0.19	-	50,50,50,50	0
56	MG	1A	3736	1/1	0.96	0.17	-	38,38,38,38	0
56	MG	1A	3072	1/1	0.96	0.16	-	56,56,56,56	0
56	MG	2a	3249	1/1	0.95	0.12	-	61,61,61,61	0
56	MG	1A	3299	1/1	0.92	0.13	-	61,61,61,61	0
56	MG	1a	1628	1/1	0.79	0.15	-	61,61,61,61	0
56	MG	1A	3097	1/1	0.89	0.29	-	55,55,55,55	0
56	MG	1A	3499	1/1	0.81	0.16	-	62,62,62,62	0
56	MG	1B	215	1/1	0.95	0.16	-	31,31,31,31	0
56	MG	1f	3001	1/1	0.65	0.13	-	68,68,68,68	0
56	MG	2A	3102	1/1	0.85	0.15	-	52,52,52,52	0
56	MG	2a	3225	1/1	0.94	0.14	-	59,59,59,59	0
56	MG	2a	3009	1/1	0.97	0.23	-	65,65,65,65	0
56	MG	2a	3050	1/1	0.90	0.41	-	93,93,93,93	0
56	MG	2A	3287	1/1	0.75	0.14	-	56,56,56,56	0
56	MG	2a	3167	1/1	0.89	0.58	-	80,80,80,80	0
56	MG	2A	3353	1/1	0.70	0.14	-	61,61,61,61	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3023	1/1	0.58	0.19	-	50,50,50,50	0
56	MG	2A	3434	1/1	0.96	0.23	-	63,63,63,63	0
56	MG	1A	3430	1/1	0.90	0.12	-	62,62,62,62	0
56	MG	1A	3410	1/1	0.89	0.14	-	49,49,49,49	0
56	MG	1A	3079	1/1	0.79	0.14	-	58,58,58,58	0
56	MG	2a	3072	1/1	0.69	0.12	-	77,77,77,77	0
56	MG	1A	3549	1/1	0.85	0.19	-	48,48,48,48	0
56	MG	2A	3238	1/1	0.84	0.14	-	60,60,60,60	0
56	MG	1B	212	1/1	0.96	0.20	-	54,54,54,54	0
56	MG	2A	3001	1/1	0.76	0.21	-	45,45,45,45	0
56	MG	2A	3158	1/1	0.88	0.17	-	47,47,47,47	0
56	MG	2A	3035	1/1	0.87	0.12	-	58,58,58,58	0
56	MG	2a	3089	1/1	0.93	0.09	-	66,66,66,66	0
56	MG	1A	3030	1/1	0.95	0.24	-	49,49,49,49	0
56	MG	2A	3333	1/1	0.85	0.16	-	51,51,51,51	0
56	MG	1x	106	1/1	0.87	0.21	-	63,63,63,63	0
56	MG	1a	1743	1/1	0.97	0.11	-	56,56,56,56	0
56	MG	1B	216	1/1	0.99	0.22	-	54,54,54,54	0
56	MG	2A	3113	1/1	0.67	0.17	-	64,64,64,64	0
56	MG	2a	3098	1/1	0.88	0.30	-	50,50,50,50	0
56	MG	2a	3211	1/1	0.95	0.12	-	61,61,61,61	0
56	MG	2a	3224	1/1	0.94	0.17	-	54,54,54,54	0
56	MG	1a	1759	1/1	0.91	0.18	-	70,70,70,70	0
56	MG	1A	3534	1/1	0.86	0.23	-	35,35,35,35	0
56	MG	1A	3943	1/1	0.94	0.17	-	68,68,68,68	0
56	MG	2A	3248	1/1	0.95	0.22	-	56,56,56,56	0
56	MG	1a	1602	1/1	0.86	0.35	-	64,64,64,64	0
56	MG	1A	3820	1/1	0.90	0.11	-	57,57,57,57	0
56	MG	2B	3008	1/1	0.96	0.19	-	47,47,47,47	0
56	MG	1a	1726	1/1	0.88	0.20	-	54,54,54,54	0
56	MG	1A	3642	1/1	0.92	0.19	-	58,58,58,58	0
56	MG	2d	502	1/1	0.96	0.23	-	40,40,40,40	0
56	MG	1A	3494	1/1	0.88	0.34	-	51,51,51,51	0
56	MG	1A	3298	1/1	0.95	0.28	-	50,50,50,50	0
56	MG	2a	3215	1/1	0.92	0.24	-	56,56,56,56	0
56	MG	1A	3274	1/1	0.85	0.21	-	41,41,41,41	0
56	MG	1a	1787	1/1	0.94	0.16	-	58,58,58,58	0
56	MG	1A	3075	1/1	0.92	0.18	-	71,71,71,71	0
56	MG	1A	3062	1/1	0.91	0.13	-	48,48,48,48	0
56	MG	1A	3833	1/1	0.76	0.15	-	78,78,78,78	0
56	MG	1a	1613	1/1	0.97	0.06	-	60,60,60,60	0
56	MG	1A	3026	1/1	0.85	0.14	-	53,53,53,53	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	2A	3061	1/1	0.42	0.27	-	68,68,68,68	0
56	MG	10	106	1/1	0.90	0.37	-	61,61,61,61	0
56	MG	2Q	3001	1/1	0.93	0.33	-	48,48,48,48	0
56	MG	1A	3044	1/1	0.79	1.11	-	45,45,45,45	0
56	MG	1A	3562	1/1	0.93	0.12	-	55,55,55,55	0
56	MG	2A	3166	1/1	0.96	0.26	-	49,49,49,49	0
56	MG	1A	3348	1/1	0.97	0.13	-	47,47,47,47	0
56	MG	1A	3372	1/1	0.81	0.53	-	57,57,57,57	0
56	MG	2A	3403	1/1	0.92	0.18	-	62,62,62,62	0
56	MG	2A	3033	1/1	0.92	0.32	-	61,61,61,61	0
56	MG	1A	3279	1/1	0.93	0.19	-	28,28,28,28	0
56	MG	1A	3211	1/1	0.93	0.14	-	58,58,58,58	0
56	MG	1A	3399	1/1	0.70	0.31	-	58,58,58,58	0
56	MG	1A	3149	1/1	0.85	0.16	-	55,55,55,55	0
56	MG	1A	3743	1/1	0.96	0.38	-	57,57,57,57	0
56	MG	2a	3171	1/1	0.94	0.11	-	68,68,68,68	0
56	MG	1a	1826	1/1	0.97	0.15	-	36,36,36,36	0
56	MG	2A	3253	1/1	0.96	0.24	-	43,43,43,43	0
56	MG	2A	3467	1/1	0.97	0.11	-	46,46,46,46	0
56	MG	2A	3021	1/1	0.88	0.28	-	58,58,58,58	0
56	MG	2A	3077	1/1	0.94	0.20	-	64,64,64,64	0
56	MG	1a	1747	1/1	0.97	0.09	-	56,56,56,56	0
56	MG	2A	3165	1/1	0.93	0.22	-	49,49,49,49	0
56	MG	1o	3002	1/1	0.95	0.11	-	44,44,44,44	0
56	MG	1A	3239	1/1	0.91	0.56	-	36,36,36,36	0
56	MG	2A	3419	1/1	0.93	0.07	-	56,56,56,56	0
56	MG	1A	3585	1/1	0.97	0.13	-	51,51,51,51	0
56	MG	1A	3346	1/1	0.92	0.10	-	63,63,63,63	0
56	MG	2B	3009	1/1	0.92	0.23	-	63,63,63,63	0
56	MG	1a	1848	1/1	0.94	0.23	-	57,57,57,57	0
56	MG	1A	3448	1/1	0.87	0.17	-	49,49,49,49	0
56	MG	2A	3439	1/1	0.95	0.08	-	68,68,68,68	0
56	MG	2A	3169	1/1	0.84	0.25	-	33,33,33,33	0
56	MG	2A	3055	1/1	0.89	0.30	-	40,40,40,40	0
56	MG	1A	3525	1/1	0.89	0.15	-	60,60,60,60	0
56	MG	2A	3290	1/1	0.91	0.14	-	49,49,49,49	0
56	MG	2A	3036	1/1	0.70	0.10	-	74,74,74,74	0
56	MG	1A	3479	1/1	0.82	0.41	-	38,38,38,38	0
56	MG	2a	3001	1/1	0.80	0.19	-	60,60,60,60	0
56	MG	1A	3349	1/1	0.70	0.17	-	73,73,73,73	0
56	MG	2A	3382	1/1	0.97	0.11	-	42,42,42,42	0
56	MG	2A	3490	1/1	0.90	0.30	-	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3184	1/1	0.96	0.20	-	52,52,52,52	0
56	MG	1A	3949	1/1	0.95	0.15	-	48,48,48,48	0
56	MG	1A	3377	1/1	0.77	0.24	-	61,61,61,61	0
56	MG	1A	3051	1/1	0.96	0.61	-	50,50,50,50	0
56	MG	1A	3389	1/1	0.96	0.61	-	45,45,45,45	0
56	MG	2a	3080	1/1	0.98	0.07	-	40,40,40,40	0
56	MG	2a	3056	1/1	0.76	0.12	-	85,85,85,85	0
56	MG	1A	3379	1/1	0.85	0.28	-	61,61,61,61	0
56	MG	2a	3239	1/1	0.97	0.05	-	76,76,76,76	0
56	MG	2A	3362	1/1	0.97	0.17	-	57,57,57,57	0
56	MG	2A	3524	1/1	0.91	0.11	-	41,41,41,41	0
56	MG	2A	3254	1/1	0.79	0.28	-	40,40,40,40	0
56	MG	1A	3687	1/1	0.99	0.13	-	48,48,48,48	0
56	MG	1A	3213	1/1	0.89	0.13	-	33,33,33,33	0
56	MG	1A	3654	1/1	0.86	0.18	-	67,67,67,67	0
56	MG	1A	3688	1/1	0.98	0.16	-	41,41,41,41	0
56	MG	1A	3412	1/1	0.93	0.56	-	45,45,45,45	0
56	MG	1a	1770	1/1	0.94	0.07	-	67,67,67,67	0
56	MG	1A	3513	1/1	0.75	0.34	-	57,57,57,57	0
56	MG	2A	3223	1/1	0.96	0.42	-	50,50,50,50	0
56	MG	2a	3067	1/1	0.91	0.10	-	88,88,88,88	0
56	MG	2A	3360	1/1	0.92	0.14	-	58,58,58,58	0
56	MG	1A	3320	1/1	0.87	0.06	-	93,93,93,93	0
56	MG	2A	3177	1/1	0.90	0.22	-	48,48,48,48	0
56	MG	2A	3116	1/1	0.93	0.12	-	47,47,47,47	0
56	MG	2a	3178	1/1	0.94	0.15	-	59,59,59,59	0
56	MG	1A	3269	1/1	0.98	0.11	-	36,36,36,36	0
56	MG	1A	3814	1/1	0.96	0.24	-	56,56,56,56	0
56	MG	2A	3456	1/1	0.92	0.13	-	51,51,51,51	0
56	MG	2B	3003	1/1	0.78	0.21	-	73,73,73,73	0
56	MG	2A	3034	1/1	0.82	0.25	-	54,54,54,54	0
56	MG	2a	3182	1/1	0.97	0.15	-	51,51,51,51	0
56	MG	2A	3190	1/1	0.92	0.20	-	54,54,54,54	0
56	MG	1A	3189	1/1	0.92	0.18	-	39,39,39,39	0
56	MG	1A	3577	1/1	0.97	0.23	-	60,60,60,60	0
56	MG	2A	3082	1/1	0.88	0.19	-	41,41,41,41	0
56	MG	1A	3363	1/1	0.78	0.18	-	68,68,68,68	0
56	MG	1A	3929	1/1	0.89	0.20	-	58,58,58,58	0
56	MG	2A	3098	1/1	0.90	0.12	-	51,51,51,51	0
56	MG	1N	3005	1/1	0.94	0.10	-	51,51,51,51	0
56	MG	1a	1807	1/1	0.83	0.17	-	60,60,60,60	0
56	MG	2A	3202	1/1	0.96	0.16	-	46,46,46,46	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3234	1/1	0.92	0.10	-	43,43,43,43	0
56	MG	2a	3096	1/1	0.68	0.22	-	80,80,80,80	0
56	MG	2a	3119	1/1	0.96	0.28	-	56,56,56,56	0
56	MG	1T	202	1/1	0.84	0.13	-	55,55,55,55	0
56	MG	1T	205	1/1	0.89	0.27	-	65,65,65,65	0
56	MG	1a	1703	1/1	0.80	0.14	-	62,62,62,62	0
56	MG	1A	3428	1/1	0.88	0.11	-	65,65,65,65	0
56	MG	1a	1685	1/1	0.94	0.16	-	53,53,53,53	0
56	MG	1A	3667	1/1	0.96	0.13	-	49,49,49,49	0
56	MG	2A	3187	1/1	0.82	1.30	-	49,49,49,49	0
56	MG	2A	3104	1/1	0.82	0.56	-	54,54,54,54	0
56	MG	2A	3275	1/1	0.95	0.12	-	60,60,60,60	0
56	MG	1D	304	1/1	0.71	0.52	-	58,58,58,58	0
56	MG	1a	1695	1/1	0.92	0.23	-	50,50,50,50	0
56	MG	2A	3013	1/1	0.84	0.17	-	68,68,68,68	0
56	MG	1a	1615	1/1	0.90	0.18	-	55,55,55,55	0
56	MG	1A	3120	1/1	0.81	0.50	-	41,41,41,41	0
56	MG	2A	3411	1/1	0.93	0.14	-	64,64,64,64	0
56	MG	2A	3109	1/1	0.95	0.15	-	64,64,64,64	0
56	MG	1A	3330	1/1	0.84	0.13	-	41,41,41,41	0
56	MG	2A	3144	1/1	0.91	0.31	-	51,51,51,51	0
56	MG	1A	3544	1/1	0.89	0.13	-	36,36,36,36	0
56	MG	1A	3223	1/1	0.97	0.24	-	33,33,33,33	0
56	MG	1A	3680	1/1	0.98	0.10	-	42,42,42,42	0
56	MG	1a	1642	1/1	0.84	0.12	-	69,69,69,69	0
56	MG	1a	1658	1/1	0.91	0.35	-	67,67,67,67	0
56	MG	1A	3115	1/1	0.91	0.13	-	49,49,49,49	0
56	MG	2A	3010	1/1	0.89	0.10	-	56,56,56,56	0
56	MG	1A	3114	1/1	0.93	0.14	-	52,52,52,52	0
56	MG	2a	3006	1/1	0.84	0.10	-	62,62,62,62	0
56	MG	1A	3944	1/1	0.90	0.08	-	68,68,68,68	0
56	MG	2A	3288	1/1	0.89	0.14	-	57,57,57,57	0
56	MG	1A	3406	1/1	0.88	0.10	-	50,50,50,50	0
56	MG	1a	1860	1/1	0.94	0.21	-	55,55,55,55	0
56	MG	2A	3343	1/1	0.95	0.14	-	56,56,56,56	0
56	MG	2A	3181	1/1	0.96	0.10	-	39,39,39,39	0
56	MG	1d	503	1/1	0.99	0.11	-	52,52,52,52	0
56	MG	1A	3474	1/1	0.91	0.37	-	39,39,39,39	0
56	MG	1A	3693	1/1	0.98	0.17	-	21,21,21,21	0
56	MG	1A	3595	1/1	0.95	0.31	-	56,56,56,56	0
56	MG	2A	3463	1/1	0.95	0.04	-	57,57,57,57	0
56	MG	1a	1830	1/1	0.96	0.25	-	61,61,61,61	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1a	1866	1/1	0.95	0.17	-	61,61,61,61	0
56	MG	1A	3488	1/1	0.89	0.13	-	65,65,65,65	0
56	MG	1A	3846	1/1	0.85	0.08	-	78,78,78,78	0
56	MG	2A	3281	1/1	0.94	0.22	-	59,59,59,59	0
56	MG	1A	3038	1/1	0.92	0.10	-	52,52,52,52	0
56	MG	1A	3703	1/1	0.86	0.15	-	49,49,49,49	0
56	MG	2A	3086	1/1	0.83	0.30	-	49,49,49,49	0
56	MG	1A	3368	1/1	0.90	0.24	-	25,25,25,25	0
56	MG	2a	3097	1/1	0.90	0.21	-	71,71,71,71	0
56	MG	2A	3474	1/1	0.89	0.21	-	41,41,41,41	0
56	MG	1a	1823	1/1	0.94	0.18	-	69,69,69,69	0
56	MG	1A	3806	1/1	0.91	0.28	-	61,61,61,61	0
56	MG	1A	3644	1/1	0.96	0.20	-	19,19,19,19	0
56	MG	2a	3023	1/1	0.82	0.17	-	65,65,65,65	0
56	MG	2A	3400	1/1	0.90	0.49	-	48,48,48,48	0
56	MG	1A	3583	1/1	0.83	0.10	-	55,55,55,55	0
56	MG	2A	3501	1/1	0.68	0.16	-	85,85,85,85	0
56	MG	2A	3272	1/1	0.90	0.23	-	46,46,46,46	0
56	MG	1z	101	1/1	0.77	0.40	-	76,76,76,76	0
56	MG	2A	3341	1/1	0.86	0.17	-	52,52,52,52	0
56	MG	1a	1621	1/1	0.86	0.23	-	68,68,68,68	0
56	MG	2N	8001	1/1	0.96	0.15	-	53,53,53,53	0
56	MG	2a	3197	1/1	0.95	0.12	-	58,58,58,58	0
56	MG	2a	3195	1/1	0.97	0.06	-	43,43,43,43	0
56	MG	1A	3109	1/1	0.94	0.31	-	55,55,55,55	0
56	MG	1A	3307	1/1	0.95	0.30	-	59,59,59,59	0
56	MG	2A	3271	1/1	0.83	0.15	-	27,27,27,27	0
56	MG	1A	3830	1/1	0.93	0.26	-	63,63,63,63	0
56	MG	1A	3134	1/1	0.93	0.51	-	45,45,45,45	0
56	MG	1A	3397	1/1	0.91	0.71	-	49,49,49,49	0
56	MG	1A	3237	1/1	0.94	0.13	-	46,46,46,46	0
56	MG	1a	1698	1/1	0.92	0.14	-	55,55,55,55	0
56	MG	2A	3230	1/1	0.93	0.09	-	47,47,47,47	0
56	MG	2a	3246	1/1	0.94	0.07	-	62,62,62,62	0
56	MG	1A	3090	1/1	0.94	0.43	-	53,53,53,53	0
56	MG	1A	3214	1/1	0.96	0.25	-	60,60,60,60	0
56	MG	1A	3777	1/1	0.91	0.13	-	49,49,49,49	0
56	MG	1D	301	1/1	0.89	0.18	-	56,56,56,56	0
56	MG	1A	3381	1/1	0.89	0.17	-	47,47,47,47	0
56	MG	2a	3156	1/1	0.76	0.10	-	72,72,72,72	0
56	MG	2a	3141	1/1	0.87	0.18	-	73,73,73,73	0
56	MG	1A	3640	1/1	0.86	0.20	-	24,24,24,24	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3942	1/1	0.89	0.12	-	53,53,53,53	0
56	MG	2A	3532	1/1	0.96	0.09	-	51,51,51,51	0
56	MG	2A	3114	1/1	0.84	0.13	-	57,57,57,57	0
56	MG	2A	3005	1/1	0.87	0.13	-	54,54,54,54	0
56	MG	2A	3060	1/1	0.95	0.14	-	43,43,43,43	0
56	MG	2A	3530	1/1	0.84	0.12	-	46,46,46,46	0
56	MG	2a	3175	1/1	0.91	0.08	-	62,62,62,62	0
56	MG	1A	3511	1/1	0.81	0.13	-	76,76,76,76	0
56	MG	1A	3587	1/1	0.97	0.11	-	40,40,40,40	0
56	MG	1a	1778	1/1	0.93	0.26	-	57,57,57,57	0
56	MG	2A	3170	1/1	0.94	0.12	-	42,42,42,42	0
56	MG	1A	3018	1/1	0.77	0.12	-	59,59,59,59	0
56	MG	1a	1854	1/1	0.83	0.09	-	82,82,82,82	0
56	MG	1a	1686	1/1	0.90	0.17	-	49,49,49,49	0
56	MG	1A	3469	1/1	0.96	0.13	-	49,49,49,49	0
56	MG	1A	3378	1/1	0.86	0.49	-	69,69,69,69	0
56	MG	1A	3647	1/1	0.83	0.21	-	74,74,74,74	0
56	MG	2a	3064	1/1	0.94	0.08	-	77,77,77,77	0
56	MG	1A	3231	1/1	0.94	1.03	-	49,49,49,49	0
56	MG	2a	3091	1/1	0.81	0.25	-	61,61,61,61	0
56	MG	2A	3433	1/1	0.90	0.14	-	65,65,65,65	0
56	MG	1a	1694	1/1	0.79	0.16	-	78,78,78,78	0
56	MG	1A	3003	1/1	0.91	0.18	-	46,46,46,46	0
56	MG	1A	3424	1/1	0.95	0.15	-	61,61,61,61	0
56	MG	1A	3082	1/1	0.67	0.17	-	56,56,56,56	0
56	MG	2a	3044	1/1	0.84	0.21	-	71,71,71,71	0
56	MG	1a	1761	1/1	0.92	0.21	-	86,86,86,86	0
56	MG	1A	3600	1/1	0.86	0.20	-	54,54,54,54	0
56	MG	2A	3087	1/1	0.82	0.21	-	59,59,59,59	0
56	MG	1a	1601	1/1	0.93	0.10	-	53,53,53,53	0
56	MG	2A	3477	1/1	0.91	0.19	-	57,57,57,57	0
56	MG	1A	3121	1/1	0.88	0.85	-	38,38,38,38	0
56	MG	1a	1853	1/1	0.96	0.14	-	57,57,57,57	0
56	MG	1A	3455	1/1	0.90	0.16	-	63,63,63,63	0
56	MG	1A	3328	1/1	0.82	0.11	-	65,65,65,65	0
56	MG	2A	3323	1/1	0.89	0.10	-	48,48,48,48	0
56	MG	1A	3891	1/1	0.94	0.08	-	62,62,62,62	0
56	MG	1A	3254	1/1	0.92	0.33	-	34,34,34,34	0
56	MG	1A	3981	1/1	0.91	0.24	-	50,50,50,50	0
56	MG	1A	3029	1/1	0.94	0.14	-	54,54,54,54	0
56	MG	2A	3032	1/1	0.93	0.14	-	53,53,53,53	0
56	MG	2A	3057	1/1	0.95	0.23	-	58,58,58,58	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1a	1742	1/1	0.90	0.18	-	51,51,51,51	0
56	MG	13	102	1/1	0.87	0.14	-	56,56,56,56	0
56	MG	1A	3063	1/1	0.98	0.05	-	53,53,53,53	0
56	MG	2A	3422	1/1	0.98	0.30	-	57,57,57,57	0
56	MG	2A	3176	1/1	0.91	0.14	-	43,43,43,43	0
56	MG	1A	3491	1/1	0.94	0.42	-	61,61,61,61	0
56	MG	1a	1786	1/1	0.92	0.52	-	92,92,92,92	0
56	MG	1A	3503	1/1	0.87	0.11	-	62,62,62,62	0
56	MG	1a	1740	1/1	0.98	0.07	-	47,47,47,47	0
56	MG	2a	3263	1/1	0.96	0.23	-	71,71,71,71	0
56	MG	1a	1626	1/1	0.76	0.12	-	80,80,80,80	0
56	MG	1A	3116	1/1	0.95	0.29	-	41,41,41,41	0
56	MG	1A	3117	1/1	0.90	0.19	-	48,48,48,48	0
56	MG	1A	3133	1/1	0.88	0.33	-	59,59,59,59	0
56	MG	1A	3799	1/1	0.98	0.33	-	42,42,42,42	0
56	MG	2a	3157	1/1	0.81	0.14	-	74,74,74,74	0
56	MG	1A	3524	1/1	0.92	0.16	-	54,54,54,54	0
56	MG	15	105	1/1	0.81	0.10	-	49,49,49,49	0
56	MG	1A	3081	1/1	0.93	0.30	-	58,58,58,58	0
56	MG	1A	3746	1/1	0.97	0.18	-	7,7,7,7	0
56	MG	1a	1648	1/1	0.95	0.11	-	47,47,47,47	0
56	MG	1A	3883	1/1	0.84	0.59	-	58,58,58,58	0
56	MG	1A	3059	1/1	0.90	0.13	-	46,46,46,46	0
56	MG	1A	3358	1/1	0.90	0.70	-	44,44,44,44	0
56	MG	1A	3014	1/1	0.88	0.33	-	53,53,53,53	0
56	MG	2a	3204	1/1	0.92	0.14	-	64,64,64,64	0
56	MG	1a	1762	1/1	0.93	0.07	-	89,89,89,89	0
56	MG	2A	3448	1/1	0.87	0.14	-	56,56,56,56	0
56	MG	1A	3529	1/1	0.98	0.24	-	48,48,48,48	0
56	MG	1A	3571	1/1	0.99	0.22	-	25,25,25,25	0
56	MG	1A	3628	1/1	0.94	0.18	-	28,28,28,28	0
56	MG	2A	3069	1/1	0.93	0.11	-	37,37,37,37	0
56	MG	2a	3172	1/1	0.96	0.14	-	60,60,60,60	0
56	MG	2a	3199	1/1	0.99	0.16	-	62,62,62,62	0
56	MG	1P	201	1/1	0.69	0.40	-	47,47,47,47	0
56	MG	2A	3347	1/1	0.92	0.10	-	60,60,60,60	0
56	MG	1A	3950	1/1	0.96	0.16	-	41,41,41,41	0
56	MG	2A	3258	1/1	0.74	0.13	-	30,30,30,30	0
56	MG	2a	3037	1/1	0.87	0.15	-	53,53,53,53	0
56	MG	1N	3004	1/1	0.92	0.23	-	36,36,36,36	0
56	MG	2A	3084	1/1	0.93	0.20	-	54,54,54,54	0
56	MG	2A	3496	1/1	0.91	0.08	-	42,42,42,42	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3300	1/1	0.91	0.14	-	54,54,54,54	0
56	MG	1A	3669	1/1	0.95	0.11	-	46,46,46,46	0
56	MG	2a	3137	1/1	0.92	0.09	-	67,67,67,67	0
56	MG	1A	3064	1/1	0.93	0.41	-	40,40,40,40	0
56	MG	2A	3242	1/1	0.97	0.15	-	52,52,52,52	0
56	MG	1A	3971	1/1	0.91	0.10	-	62,62,62,62	0
56	MG	1A	3782	1/1	0.96	0.17	-	34,34,34,34	0
56	MG	1A	3302	1/1	0.89	0.48	-	41,41,41,41	0
56	MG	1A	3017	1/1	0.93	0.15	-	40,40,40,40	0
56	MG	1A	3477	1/1	0.94	0.29	-	66,66,66,66	0
56	MG	2D	304	1/1	0.88	0.30	-	42,42,42,42	0
56	MG	1B	204	1/1	0.79	0.28	-	54,54,54,54	0
56	MG	1A	3889	1/1	0.86	0.37	-	60,60,60,60	0
56	MG	1a	1608	1/1	0.85	0.28	-	59,59,59,59	0
56	MG	1A	3183	1/1	0.85	0.73	-	44,44,44,44	0
56	MG	2a	3154	1/1	0.95	0.45	-	87,87,87,87	0
56	MG	1A	3382	1/1	0.98	0.08	-	40,40,40,40	0
56	MG	1A	3353	1/1	0.93	0.24	-	58,58,58,58	0
56	MG	2A	3545	1/1	0.98	0.19	-	39,39,39,39	0
56	MG	1A	3904	1/1	0.96	0.12	-	27,27,27,27	0
56	MG	1A	3946	1/1	0.97	0.23	-	41,41,41,41	0
56	MG	2A	3397	1/1	0.91	0.17	-	66,66,66,66	0
56	MG	1A	3718	1/1	0.91	0.13	-	41,41,41,41	0
56	MG	2a	3024	1/1	0.87	0.17	-	76,76,76,76	0
56	MG	1A	3333	1/1	0.91	0.26	-	70,70,70,70	0
56	MG	1A	3369	1/1	0.97	0.51	-	40,40,40,40	0
56	MG	2A	3260	1/1	0.94	0.14	-	72,72,72,72	0
56	MG	1A	3388	1/1	0.87	0.10	-	53,53,53,53	0
56	MG	2A	3365	1/1	0.62	0.13	-	63,63,63,63	0
56	MG	1a	1831	1/1	0.96	0.12	-	54,54,54,54	0
56	MG	2a	3100	1/1	0.77	0.15	-	66,66,66,66	0
56	MG	1A	3671	1/1	0.94	0.32	-	56,56,56,56	0
56	MG	1A	3057	1/1	0.66	0.70	-	50,50,50,50	0
56	MG	1A	3678	1/1	0.95	0.16	-	61,61,61,61	0
56	MG	1V	201	1/1	0.89	0.30	-	45,45,45,45	0
56	MG	1A	3523	1/1	0.82	0.17	-	59,59,59,59	0
56	MG	2A	3120	1/1	0.99	0.25	-	45,45,45,45	0
56	MG	2a	3043	1/1	0.65	0.20	-	65,65,65,65	0
56	MG	1A	3560	1/1	0.70	0.13	-	45,45,45,45	0
56	MG	1A	3584	1/1	0.95	0.17	-	33,33,33,33	0
56	MG	2A	3311	1/1	0.93	0.21	-	42,42,42,42	0
56	MG	1A	4021	1/1	0.95	0.12	-	57,57,57,57	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1e	3001	1/1	0.92	0.13	-	69,69,69,69	0
56	MG	2a	3219	1/1	0.94	0.14	-	52,52,52,52	0
56	MG	2A	3294	1/1	0.89	0.17	-	37,37,37,37	0
56	MG	1A	3245	1/1	0.94	0.14	-	49,49,49,49	0
56	MG	1A	3440	1/1	0.80	0.10	-	66,66,66,66	0
56	MG	1A	3451	1/1	0.80	0.88	-	66,66,66,66	0
56	MG	2a	3004	1/1	0.92	0.16	-	46,46,46,46	0
56	MG	1A	3675	1/1	0.82	0.11	-	79,79,79,79	0
56	MG	2A	3174	1/1	0.85	0.12	-	45,45,45,45	0
56	MG	1a	1705	1/1	0.74	0.09	-	73,73,73,73	0
56	MG	2A	3065	1/1	0.91	0.12	-	35,35,35,35	0
56	MG	1A	3902	1/1	0.94	0.10	-	45,45,45,45	0
56	MG	1A	3500	1/1	0.92	0.23	-	55,55,55,55	0
56	MG	2A	3045	1/1	0.92	0.46	-	48,48,48,48	0
56	MG	1B	219	1/1	0.87	0.07	-	60,60,60,60	0
56	MG	1a	1781	1/1	0.91	0.28	-	46,46,46,46	0
56	MG	2a	3060	1/1	0.93	0.20	-	86,86,86,86	0
56	MG	1A	3657	1/1	0.88	0.17	-	49,49,49,49	0
56	MG	2a	3095	1/1	0.95	0.23	-	45,45,45,45	0
56	MG	1A	3362	1/1	0.89	0.18	-	56,56,56,56	0
56	MG	2A	3185	1/1	0.90	0.19	-	52,52,52,52	0
56	MG	2A	3192	1/1	0.99	0.14	-	29,29,29,29	0
56	MG	1A	3019	1/1	0.94	0.13	-	49,49,49,49	0
56	MG	1A	3435	1/1	0.97	0.12	-	56,56,56,56	0
56	MG	2A	3103	1/1	0.79	0.37	-	53,53,53,53	0
56	MG	1A	3924	1/1	0.73	0.14	-	66,66,66,66	0
56	MG	1A	3065	1/1	0.89	0.34	-	49,49,49,49	0
56	MG	2A	3307	1/1	0.94	0.20	-	35,35,35,35	0
56	MG	1A	3692	1/1	0.94	0.14	-	33,33,33,33	0
56	MG	1A	3762	1/1	0.99	0.16	-	45,45,45,45	0
56	MG	1A	3875	1/1	0.85	0.30	-	83,83,83,83	0
56	MG	1A	3126	1/1	0.64	0.16	-	53,53,53,53	0
56	MG	1A	3625	1/1	0.91	0.15	-	22,22,22,22	0
56	MG	1A	3303	1/1	0.89	0.30	-	63,63,63,63	0
56	MG	2a	3047	1/1	0.83	0.21	-	63,63,63,63	0
56	MG	2a	3008	1/1	0.73	0.19	-	67,67,67,67	0
56	MG	1A	3557	1/1	0.93	0.19	-	31,31,31,31	0
56	MG	1A	3285	1/1	0.95	0.20	-	54,54,54,54	0
56	MG	2A	3053	1/1	0.91	0.14	-	55,55,55,55	0
56	MG	2A	3117	1/1	0.75	0.14	-	53,53,53,53	0
56	MG	1A	3661	1/1	0.93	0.19	-	36,36,36,36	0
56	MG	1W	3004	1/1	0.95	0.46	-	31,31,31,31	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1A	3572	1/1	0.97	0.17	-	50,50,50,50	0
56	MG	2A	3505	1/1	0.89	0.15	-	57,57,57,57	0
56	MG	2D	306	1/1	0.94	0.08	-	68,68,68,68	0
56	MG	2A	3161	1/1	0.97	0.24	-	32,32,32,32	0
56	MG	1A	3318	1/1	0.90	0.29	-	52,52,52,52	0
56	MG	2a	3181	1/1	0.89	0.18	-	47,47,47,47	0
56	MG	1A	3754	1/1	0.97	0.18	-	18,18,18,18	0
56	MG	2A	3228	1/1	0.81	0.19	-	57,57,57,57	0
56	MG	1A	3141	1/1	0.90	0.19	-	66,66,66,66	0
56	MG	1A	3281	1/1	0.84	0.42	-	53,53,53,53	0
56	MG	1A	3340	1/1	0.94	0.24	-	47,47,47,47	0
56	MG	2A	3462	1/1	0.98	0.22	-	60,60,60,60	0
56	MG	2a	3086	1/1	0.82	0.21	-	64,64,64,64	0
56	MG	1A	3486	1/1	0.82	0.14	-	67,67,67,67	0
56	MG	1A	3232	1/1	0.98	0.61	-	43,43,43,43	0
56	MG	2A	3269	1/1	0.86	0.14	-	41,41,41,41	0
56	MG	1A	3913	1/1	0.86	0.16	-	64,64,64,64	0
56	MG	1A	3840	1/1	0.87	0.17	-	63,63,63,63	0
56	MG	2a	3170	1/1	0.81	0.14	-	77,77,77,77	0
56	MG	1a	1699	1/1	0.94	0.07	-	71,71,71,71	0
56	MG	1a	1676	1/1	0.65	0.11	-	74,74,74,74	0
56	MG	1A	3533	1/1	0.91	0.22	-	51,51,51,51	0
56	MG	2a	3216	1/1	0.86	0.25	-	56,56,56,56	0
56	MG	2A	3361	1/1	0.88	0.20	-	54,54,54,54	0
56	MG	2a	3166	1/1	0.90	0.13	-	66,66,66,66	0
56	MG	1A	3936	1/1	0.78	0.19	-	64,64,64,64	0
56	MG	1A	3106	1/1	0.80	0.17	-	64,64,64,64	0
56	MG	1a	1772	1/1	0.94	0.23	-	78,78,78,78	0
56	MG	1A	3343	1/1	0.92	0.18	-	77,77,77,77	0
56	MG	1A	3416	1/1	0.91	0.12	-	60,60,60,60	0
56	MG	1A	3228	1/1	0.89	0.15	-	59,59,59,59	0
56	MG	2a	3226	1/1	0.97	0.07	-	58,58,58,58	0
56	MG	1A	3002	1/1	0.77	0.25	-	53,53,53,53	0
56	MG	2A	3224	1/1	0.89	0.13	-	42,42,42,42	0
56	MG	2A	3041	1/1	0.84	0.41	-	53,53,53,53	0
56	MG	1A	3921	1/1	0.91	0.26	-	55,55,55,55	0
56	MG	1A	3787	1/1	0.93	0.14	-	74,74,74,74	0
56	MG	2A	3243	1/1	0.94	0.17	-	63,63,63,63	0
56	MG	2A	3091	1/1	0.76	0.23	-	49,49,49,49	0
56	MG	2A	3355	1/1	0.94	0.20	-	73,73,73,73	0
56	MG	1A	3335	1/1	0.96	0.15	-	55,55,55,55	0
56	MG	2A	3440	1/1	0.94	0.29	-	60,60,60,60	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	2A	3178	1/1	0.90	0.09	-	34,34,34,34	0
56	MG	1A	3436	1/1	0.94	0.17	-	61,61,61,61	0
56	MG	1a	1654	1/1	0.89	0.13	-	78,78,78,78	0
56	MG	1A	3796	1/1	0.99	0.12	-	16,16,16,16	0
56	MG	1A	3008	1/1	0.92	0.46	-	58,58,58,58	0
56	MG	1A	3898	1/1	0.94	0.15	-	25,25,25,25	0
56	MG	2A	3249	1/1	0.89	0.14	-	59,59,59,59	0
56	MG	2A	3543	1/1	0.94	0.21	-	56,56,56,56	0
56	MG	1A	3207	1/1	0.90	0.10	-	30,30,30,30	0
56	MG	1A	3497	1/1	0.93	0.20	-	57,57,57,57	0
56	MG	1A	3959	1/1	0.77	0.08	-	79,79,79,79	0
56	MG	1A	3011	1/1	0.95	0.10	-	67,67,67,67	0
56	MG	1N	3003	1/1	0.88	0.20	-	42,42,42,42	0
56	MG	2a	3101	1/1	0.95	0.16	-	53,53,53,53	0
56	MG	1A	3015	1/1	0.97	0.10	-	52,52,52,52	0
56	MG	2A	3101	1/1	0.92	0.23	-	57,57,57,57	0
56	MG	1k	3001	1/1	0.64	0.17	-	70,70,70,70	0
56	MG	1A	3252	1/1	0.95	0.15	-	28,28,28,28	0
56	MG	1A	3892	1/1	0.97	0.15	-	43,43,43,43	0
56	MG	1A	3454	1/1	0.93	0.48	-	66,66,66,66	0
56	MG	2A	3506	1/1	0.94	0.33	-	56,56,56,56	0
56	MG	1A	3794	1/1	0.94	0.14	-	68,68,68,68	0
56	MG	1A	3031	1/1	0.80	0.36	-	61,61,61,61	0
56	MG	1A	3140	1/1	0.92	0.41	-	62,62,62,62	0
56	MG	2a	3159	1/1	0.72	0.20	-	85,85,85,85	0
56	MG	2A	3237	1/1	0.95	0.15	-	41,41,41,41	0
56	MG	2a	3160	1/1	0.92	0.25	-	74,74,74,74	0
56	MG	2O	201	1/1	0.88	0.10	-	48,48,48,48	0
56	MG	2A	3412	1/1	0.94	0.15	-	63,63,63,63	0
56	MG	1a	1618	1/1	0.86	0.12	-	64,64,64,64	0
56	MG	1a	1674	1/1	0.94	0.13	-	54,54,54,54	0
56	MG	2A	3510	1/1	0.81	0.15	-	34,34,34,34	0
56	MG	1A	3046	1/1	0.92	0.32	-	65,65,65,65	0
56	MG	1A	3554	1/1	0.95	0.14	-	80,80,80,80	0
56	MG	2A	3160	1/1	0.91	0.21	-	52,52,52,52	0
56	MG	2a	3074	1/1	0.99	0.18	-	74,74,74,74	0
56	MG	1a	1702	1/1	0.79	0.34	-	82,82,82,82	0
56	MG	2a	3151	1/1	0.93	0.16	-	92,92,92,92	0
56	MG	2A	3138	1/1	0.90	0.14	-	54,54,54,54	0
56	MG	1N	3001	1/1	0.91	0.27	-	42,42,42,42	0
56	MG	1A	3124	1/1	0.83	0.58	-	56,56,56,56	0
56	MG	1x	102	1/1	0.90	0.30	-	67,67,67,67	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1G	3003	1/1	0.91	0.16	-	45,45,45,45	0
56	MG	1A	3712	1/1	0.94	0.14	-	43,43,43,43	0
56	MG	2A	3305	1/1	0.90	0.12	-	42,42,42,42	0
56	MG	2A	3431	1/1	0.98	0.19	-	68,68,68,68	0
56	MG	2a	3112	1/1	0.95	0.13	-	45,45,45,45	0
56	MG	1P	204	1/1	0.94	0.12	-	47,47,47,47	0
56	MG	2A	3483	1/1	0.77	0.18	-	54,54,54,54	0
56	MG	1a	1777	1/1	0.91	0.24	-	28,28,28,28	0
56	MG	1A	3831	1/1	0.94	0.15	-	70,70,70,70	0
56	MG	1B	213	1/1	0.81	0.18	-	110,110,110,110	0
56	MG	2F	302	1/1	0.93	0.21	-	47,47,47,47	0
56	MG	2A	3544	1/1	0.93	0.18	-	45,45,45,45	0
56	MG	1a	1653	1/1	0.91	0.13	-	60,60,60,60	0
56	MG	2a	3051	1/1	0.89	0.10	-	76,76,76,76	0
56	MG	1A	3370	1/1	0.94	0.13	-	66,66,66,66	0
56	MG	1N	3002	1/1	0.72	0.51	-	48,48,48,48	0
56	MG	1A	3873	1/1	0.94	0.19	-	48,48,48,48	0
56	MG	1a	1819	1/1	0.97	0.14	-	54,54,54,54	0
56	MG	1A	3914	1/1	0.95	0.15	-	38,38,38,38	0
56	MG	1U	201	1/1	0.89	0.19	-	51,51,51,51	0
56	MG	1F	307	1/1	0.68	0.26	-	59,59,59,59	0
56	MG	1A	3439	1/1	0.74	0.12	-	51,51,51,51	0
56	MG	1o	3001	1/1	0.76	0.15	-	74,74,74,74	0
56	MG	2B	3002	1/1	0.86	0.14	-	61,61,61,61	0
56	MG	2A	3118	1/1	0.84	0.21	-	75,75,75,75	0
56	MG	1A	3805	1/1	0.95	0.28	-	41,41,41,41	0
56	MG	2B	3001	1/1	0.60	0.20	-	65,65,65,65	0
56	MG	2a	3106	1/1	0.97	0.22	-	63,63,63,63	0
56	MG	2a	3245	1/1	0.95	0.22	-	62,62,62,62	0
56	MG	1A	3301	1/1	0.86	0.16	-	53,53,53,53	0
56	MG	1A	3083	1/1	0.81	0.33	-	56,56,56,56	0
56	MG	1A	3987	1/1	0.95	0.27	-	38,38,38,38	0
56	MG	2Q	3004	1/1	0.95	0.26	-	42,42,42,42	0
56	MG	1A	3966	1/1	0.82	0.98	-	47,47,47,47	0
56	MG	1A	3594	1/1	0.96	0.11	-	62,62,62,62	0
56	MG	1A	3849	1/1	0.92	0.17	-	73,73,73,73	0
56	MG	1A	3128	1/1	0.85	0.15	-	56,56,56,56	0
56	MG	1a	1696	1/1	0.95	0.16	-	70,70,70,70	0
56	MG	2A	3476	1/1	0.95	0.19	-	33,33,33,33	0
56	MG	1A	3154	1/1	0.91	0.13	-	28,28,28,28	0
56	MG	1A	3060	1/1	0.81	0.55	-	67,67,67,67	0
56	MG	2A	3251	1/1	0.95	0.09	-	53,53,53,53	0

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	LLDF	B-factors( $\text{\AA}^2$ )	Q<0.9
56	MG	1a	1857	1/1	0.92	0.25	-	73,73,73,73	0
56	MG	1A	3634	1/1	0.94	0.18	-	56,56,56,56	0
56	MG	1A	3341	1/1	0.93	0.10	-	64,64,64,64	0
56	MG	1a	1643	1/1	0.83	0.14	-	76,76,76,76	0
56	MG	1A	3804	1/1	0.93	0.12	-	31,31,31,31	0
56	MG	1A	3100	1/1	0.85	0.12	-	48,48,48,48	0
56	MG	2a	3059	1/1	0.98	0.11	-	83,83,83,83	0
56	MG	18	103	1/1	0.93	0.30	-	52,52,52,52	0
56	MG	2A	3022	1/1	0.89	0.15	-	51,51,51,51	0
56	MG	2a	3148	1/1	0.96	0.07	-	72,72,72,72	0
56	MG	1A	3955	1/1	0.85	0.15	-	63,63,63,63	0
56	MG	2A	3217	1/1	0.85	0.12	-	56,56,56,56	0
56	MG	1a	1619	1/1	0.97	0.23	-	54,54,54,54	0
56	MG	1A	3868	1/1	0.93	0.18	-	44,44,44,44	0
56	MG	1a	1792	1/1	0.83	0.10	-	54,54,54,54	0
56	MG	1A	3238	1/1	0.86	0.98	-	45,45,45,45	0
56	MG	2a	3057	1/1	0.95	0.12	-	96,96,96,96	0
56	MG	1A	3147	1/1	0.92	0.23	-	54,54,54,54	0
56	MG	2A	3083	1/1	0.84	0.14	-	49,49,49,49	0
56	MG	1A	3467	1/1	0.81	0.23	-	71,71,71,71	0
56	MG	1A	3037	1/1	0.89	0.13	-	58,58,58,58	0
56	MG	1a	1660	1/1	0.89	0.14	-	50,50,50,50	0
56	MG	2a	3231	1/1	0.87	0.17	-	75,75,75,75	0
56	MG	1A	3441	1/1	0.92	0.25	-	72,72,72,72	0
56	MG	1A	3582	1/1	0.96	0.16	-	45,45,45,45	0
56	MG	1F	301	1/1	0.89	0.12	-	52,52,52,52	0
56	MG	2A	3047	1/1	0.79	0.21	-	65,65,65,65	0
56	MG	1A	3108	1/1	0.95	0.30	-	56,56,56,56	0
56	MG	1A	3145	1/1	0.49	0.18	-	77,77,77,77	0
56	MG	1A	3176	1/1	0.96	0.86	-	40,40,40,40	0
56	MG	2Q	3002	1/1	0.97	0.13	-	39,39,39,39	0
56	MG	2A	3511	1/1	0.78	0.14	-	52,52,52,52	0
56	MG	2E	302	1/1	0.93	0.15	-	59,59,59,59	0
56	MG	2a	3174	1/1	0.93	0.19	-	69,69,69,69	0
56	MG	1a	1662	1/1	0.87	0.15	-	61,61,61,61	0

## 6.5 Other polymers

There are no such residues in this entry.