

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

There are no non-standard protein/DNA/RNA residues in this entry.

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

Of 9 ligands modelled in this entry, 9 are monoatomic - leaving 0 for Mogul analysis.

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.

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Mol	Type	Chain	Res	Atoms	RSCC	RSR	B-factors($\text{\AA}^2$)	Q<0.9
2	MG	A	305	1/1	0.96	0.15	35,35,35,35	0
2	MG	A	302	1/1	0.96	0.10	31,31,31,31	0
2	MG	A	304	1/1	0.96	0.09	43,43,43,43	0
2	MG	B	301	1/1	0.98	0.08	26,26,26,26	0
2	MG	A	301	1/1	0.98	0.13	27,27,27,27	0
2	MG	B	302	1/1	0.99	0.14	31,31,31,31	0

6.5 Other polymers [i](#)

There are no such residues in this entry.