



wwPDB X-ray Structure Validation Summary Report ⓘ

Jun 17, 2025 – 07:59 PM EDT

PDB ID : 9O3L / pdb_00009o3l
Title : Crystal structure of the wild-type *Thermus thermophilus* 70S ribosome in complex with macrolide erythromycin, mRNA, deacylated A-site tRNA^{phe}, P-site fMRC-peptidyl-tRNA^{met}, and deacylated E-site tRNA^{phe} at 2.75Å resolution
Authors : Syroegin, E.A.; Aleksandrova, E.V.; Kruglov, A.A.; Paranjpe, M.N.; Svetlov, M.S.; Polikanov, Y.S.
Deposited on : 2025-04-07
Resolution : 2.75 Å(reported)

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

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

A user guide is available at

<https://www.wwpdb.org/validation/2017/XrayValidationReportHelp>

with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

<http://www.wwpdb.org/validation/2017/FAQs#types>.

The following versions of software and data (see [references](#) ①) were used in the production of this report:

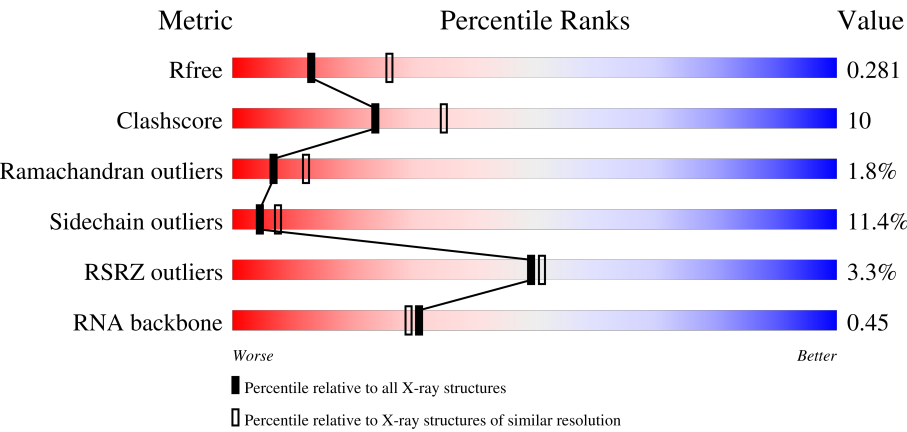
| | | |
|--------------------------------|---|--------------------------------------------------------------------|
| MolProbity | : | 4-5-2 with Phenix2.0rc1 |
| Mogul | : | 2022.3.0, CSD as543be (2022) |
| Xtriage (Phenix) | : | 2.0rc1 |
| EDS | : | 3.0 |
| buster-report | : | 1.1.7 (2018) |
| Percentile statistics | : | 20231227.v01 (using entries in the PDB archive December 27th 2023) |
| CCP4 | : | 9.0.006 (Gargrove) |
| Density-Fitness | : | 1.0.12 |
| Ideal geometry (proteins) | : | Engh & Huber (2001) |
| Ideal geometry (DNA, RNA) | : | Parkinson et al. (1996) |
| Validation Pipeline (wwPDB-VP) | : | 2.44 |

1 Overall quality at a glance i

The following experimental techniques were used to determine the structure:
X-RAY DIFFRACTION

The reported resolution of this entry is 2.75 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.













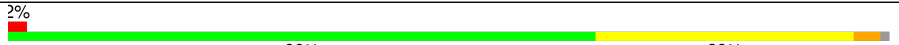
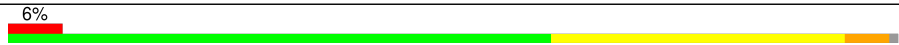

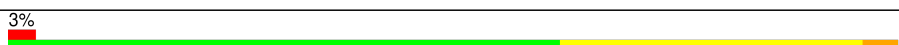



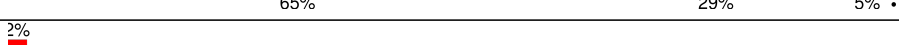
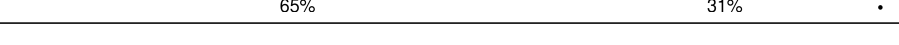
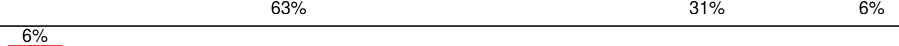





| Metric | Whole archive (#Entries) | Similar resolution (#Entries, resolution range(Å)) |
|-----------------------|-----------------------------|-------------------------------------------------------|
| R _{free} | 164625 | 1606 (2.78-2.74) |
| Clashscore | 180529 | 1689 (2.78-2.74) |
| Ramachandran outliers | 177936 | 1665 (2.78-2.74) |
| Sidechain outliers | 177891 | 1665 (2.78-2.74) |
| RSRZ outliers | 164620 | 1606 (2.78-2.74) |
| RNA backbone | 3690 | 1094 (3.00-2.52) |

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 of the lower bar indicate the fraction of residues that contain outliers for ≥ 3 , 2, 1 and 0 types of geometric quality criteria respectively. 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 | <div><div>2%</div><div><div></div><div>59%</div><div>33%</div><div>7%</div><div>.</div></div></div> |
| 1 | 2A | 2915 | <div><div>2%</div><div><div></div><div>53%</div><div>35%</div><div>8%</div><div>.</div></div></div> |
| 2 | 1B | 121 | <div><div></div><div><div>60%</div><div>38%</div><div>..</div></div></div> |
| 2 | 2B | 121 | <div><div>2%</div><div><div></div><div>43%</div><div>47%</div><div>9%</div><div>.</div></div></div> |






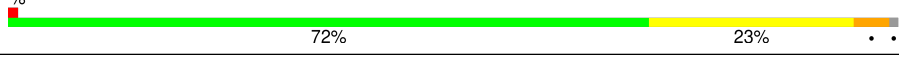

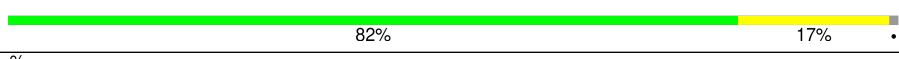
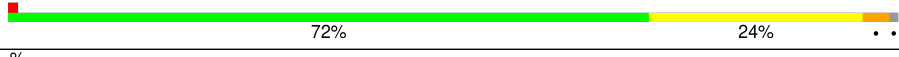


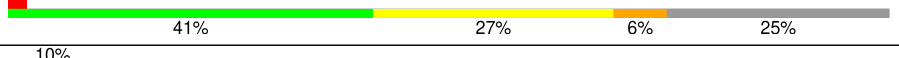
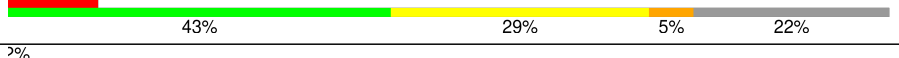






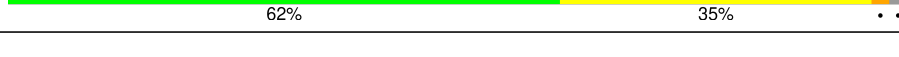
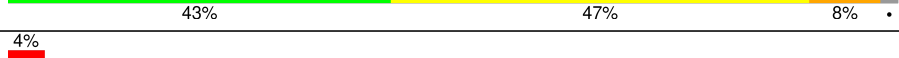


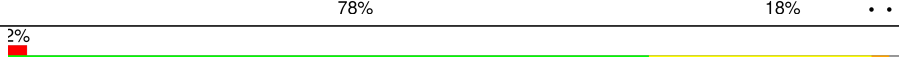

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| Mol | Chain | Length | Quality of chain |
|-----|-------|--------|--------------------------------------------------------------------------------------|
| 3 | 1D | 276 |  |
| 3 | 2D | 276 |  |
| 4 | 1E | 206 |  |
| 4 | 2E | 206 |  |
| 5 | 1F | 210 |  |
| 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 |  |












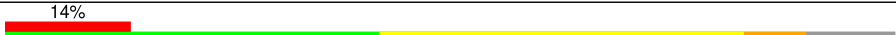

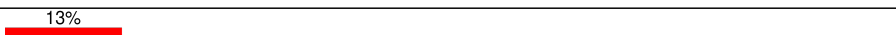
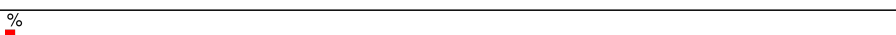
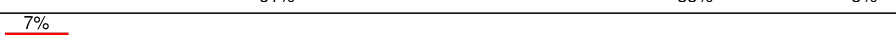

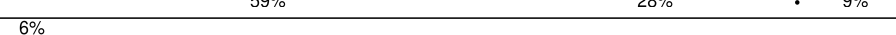







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| Mol | Chain | Length | Quality of chain |
|-----|-------|--------|--------------------------------------------------------------------------------------|
| 15 | 2T | 146 |  |
| 16 | 1U | 118 |  |
| 16 | 2U | 118 |  |
| 17 | 1V | 101 |  |
| 17 | 2V | 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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| 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 | 1w | 76 | |
| 54 | 1y | 76 | |
| 54 | 2w | 76 | |
| 54 | 2y | 76 | |
| 55 | 1x | 77 | |
| 55 | 2x | 77 | |
| 56 | 1z | 3 | |
| 56 | 2z | 3 | |

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

| Mol | Type | Chain | Res | Chirality | Geometry | Clashes | Electron density |
|-----|------|-------|------|-----------|----------|---------|------------------|
| 57 | MG | 2a | 1695 | - | - | - | X |
| 61 | SF4 | 2d | 303 | - | - | X | - |

2 Entry composition

There are 62 unique types of molecules in this entry. The entry contains 299368 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 | 2871 | Total | C | N | O | P | 0 | 0 | 0 |
| | | | 61852 | 27531 | 11572 | 19878 | 2871 | | | |
| 1 | 2A | 2800 | Total | C | N | O | P | 0 | 0 | 0 |
| | | | 60322 | 26848 | 11284 | 19390 | 2800 | | | |

- 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 |
| | | | 2577 | 1146 | 476 | 835 | 120 | | | |
| 2 | 2B | 120 | Total | C | N | O | P | 0 | 0 | 0 |
| | | | 2575 | 1146 | 476 | 833 | 120 | | | |

- 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 |
| | | | 2136 | 1349 | 423 | 361 | 3 | | | |

- Molecule 4 is a protein called 50S ribosomal protein L3.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 4 | 1E | 204 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1559 | 985 | 298 | 270 | 6 | | | |
| 4 | 2E | 204 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1559 | 985 | 298 | 270 | 6 | | | |

- Molecule 5 is a protein called 50S ribosomal protein L4.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|---------|-------|
| 5 | 1F | 202 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1583 | 1009 | 297 | 275 | 2 | | | |
| 5 | 2F | 202 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1579 | 1007 | 296 | 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 |
| | | | 1423 | 913 | 253 | 253 | 4 | | | |
| 6 | 2G | 181 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1428 | 913 | 258 | 253 | 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 |
| | | | 1097 | 701 | 191 | 204 | 1 | | | |
| 8 | 2I | 146 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1064 | 681 | 186 | 196 | 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 |
| | | | 1135 | 706 | 230 | 196 | 3 | | | |
| 11 | 2P | 149 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1135 | 706 | 230 | 196 | 3 | | | |

- Molecule 12 is a protein called 50S ribosomal protein L16.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 12 | 1Q | 141 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1122 | 715 | 212 | 188 | 7 | | | |
| 12 | 2Q | 141 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1122 | 715 | 212 | 188 | 7 | | | |

- Molecule 13 is a protein called 50S ribosomal protein L17.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 13 | 1R | 118 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 968 | 604 | 203 | 160 | 1 | | | |
| 13 | 2R | 118 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 968 | 604 | 203 | 160 | 1 | | | |

- Molecule 14 is a protein called 50S ribosomal protein L18.

| Mol | Chain | Residues | Atoms | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---------|---------|-------|
| 14 | 1S | 110 | Total | C | N | O | 0 | 0 | 0 |
| | | | 873 | 550 | 174 | 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 | 154 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1240 | 795 | 222 | 220 | 3 | | | |
| 21 | 2Z | 160 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1271 | 814 | 228 | 227 | 2 | | | |

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

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 22 | 10 | 83 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 653 | 404 | 139 | 109 | 1 | | | |
| 22 | 20 | 83 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 653 | 404 | 139 | 109 | 1 | | | |

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

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 23 | 11 | 97 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 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 |
| | | | 552 | 349 | 99 | 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 | 1500 | Total | C | N | O | P | 0 | 0 | 0 |
| | | | 32246 | 14358 | 5975 | 10413 | 1500 | | | |
| 32 | 2a | 1503 | Total | C | N | O | P | 0 | 0 | 0 |
| | | | 32327 | 14396 | 5990 | 10438 | 1503 | | | |

- 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 |
| | | | 1846 | 1179 | 331 | 331 | 5 | | | |
| 33 | 2b | 231 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1825 | 1167 | 326 | 327 | 5 | | | |

- Molecule 34 is a protein called 30S ribosomal protein S3.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 34 | 1c | 206 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1548 | 973 | 301 | 273 | 1 | | | |
| 34 | 2c | 206 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1542 | 968 | 300 | 273 | 1 | | | |

- Molecule 35 is a protein called 30S ribosomal protein S4.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|---------|-------|
| 35 | 1d | 208 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1655 | 1038 | 326 | 284 | 7 | | | |
| 35 | 2d | 208 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1674 | 1050 | 333 | 284 | 7 | | | |

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

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 36 | 1e | 148 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1129 | 714 | 213 | 198 | 4 | | | |
| 36 | 2e | 148 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1133 | 716 | 214 | 199 | 4 | | | |

- Molecule 37 is a protein called 30S ribosomal protein S6.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 37 | 1f | 100 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 810 | 514 | 144 | 149 | 3 | | | |
| 37 | 2f | 100 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 816 | 516 | 146 | 151 | 3 | | | |

- Molecule 38 is a protein called 30S ribosomal protein S7.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 38 | 1g | 155 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1231 | 766 | 243 | 216 | 6 | | | |
| 38 | 2g | 155 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1235 | 769 | 244 | 216 | 6 | | | |

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

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 39 | 1h | 137 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1088 | 689 | 206 | 191 | 2 | | | |
| 39 | 2h | 137 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 1088 | 689 | 206 | 191 | 2 | | | |

- Molecule 40 is a protein called 30S ribosomal protein S9.

| Mol | Chain | Residues | Atoms | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---------|---------|-------|
| 40 | 1i | 127 | Total | C | N | O | 0 | 0 | 0 |
| | | | 983 | 623 | 193 | 167 | | | |
| 40 | 2i | 127 | Total | C | N | O | 0 | 0 | 0 |
| | | | 978 | 619 | 190 | 169 | | | |

- 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 |
| | | | 709 | 440 | 138 | 131 | | | |
| 41 | 2j | 96 | Total | C | N | O | 0 | 0 | 0 |
| | | | 714 | 445 | 138 | 131 | | | |

- Molecule 42 is a protein called 30S ribosomal protein S11.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 42 | 1k | 114 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 829 | 516 | 155 | 155 | 3 | | | |

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| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 42 | 2k | 114 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 833 | 519 | 156 | 155 | 3 | | | |

- Molecule 43 is a protein called 30S ribosomal protein S12.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 43 | 1l | 122 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 932 | 586 | 185 | 159 | 2 | | | |
| 43 | 2l | 122 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 932 | 586 | 185 | 159 | 2 | | | |

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

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 44 | 1m | 123 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 958 | 592 | 198 | 166 | 2 | | | |
| 44 | 2m | 122 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 950 | 586 | 197 | 165 | 2 | | | |

- Molecule 45 is a protein called 30S ribosomal protein S14 type Z.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|----|---|---------|---------|-------|
| 45 | 1n | 60 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 492 | 312 | 104 | 72 | 4 | | | |
| 45 | 2n | 60 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 492 | 312 | 104 | 72 | 4 | | | |

- Molecule 46 is a protein called 30S ribosomal protein S15.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 46 | 1o | 88 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 728 | 456 | 144 | 126 | 2 | | | |
| 46 | 2o | 88 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 728 | 456 | 144 | 126 | 2 | | | |

- Molecule 47 is a protein called 30S ribosomal protein S16.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 47 | 1p | 82 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 681 | 433 | 134 | 113 | 1 | | | |
| 47 | 2p | 82 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 677 | 430 | 133 | 113 | 1 | | | |

- Molecule 48 is a protein called 30S ribosomal protein S17.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 48 | 1q | 99 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 823 | 528 | 151 | 142 | 2 | | | |
| 48 | 2q | 99 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 823 | 528 | 151 | 142 | 2 | | | |

- Molecule 49 is a protein called 30S ribosomal protein S18.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|----|--|---------|---------|-------|
| 49 | 1r | 68 | Total | C | N | O | | 0 | 0 | 0 |
| | | | 555 | 355 | 108 | 92 | | | | |
| 49 | 2r | 68 | Total | C | N | O | | 0 | 0 | 0 |
| | | | 555 | 355 | 108 | 92 | | | | |

- Molecule 50 is a protein called 30S ribosomal protein S19.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|---------|-------|
| 50 | 1s | 83 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 652 | 417 | 120 | 113 | 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 |
| | | | 728 | 446 | 156 | 124 | 2 | | | |
| 51 | 2t | 96 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 727 | 446 | 155 | 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 |
| | | | 199 | 122 | 48 | 29 | | | |
| 52 | 2u | 23 | Total | C | N | O | 0 | 0 | 0 |
| | | | 199 | 122 | 48 | 29 | | | |

- Molecule 53 is a RNA chain called MET-PHE-mRNA.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|----|---------|---------|-------|
| 53 | 1v | 13 | Total | C | N | O | P | 0 | 0 | 0 |
| | | | 277 | 125 | 51 | 88 | 13 | | | |
| 53 | 2v | 13 | Total | C | N | O | P | 0 | 0 | 0 |
| | | | 277 | 125 | 51 | 88 | 13 | | | |

- Molecule 54 is a RNA chain called A-site and E-site Deacylated tRNAphe.

| Mol | Chain | Residues | Atoms | | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|----|---|---------|---------|-------|
| 54 | 1w | 74 | Total | C | N | O | P | S | 0 | 0 | 0 |
| | | | 1592 | 713 | 285 | 518 | 74 | 2 | | | |
| 54 | 1y | 74 | Total | C | N | O | P | S | 0 | 0 | 0 |
| | | | 1585 | 707 | 285 | 518 | 74 | 1 | | | |
| 54 | 2w | 72 | Total | C | N | O | P | S | 0 | 0 | 0 |
| | | | 1544 | 690 | 278 | 502 | 72 | 2 | | | |
| 54 | 2y | 73 | Total | C | N | O | P | S | 0 | 0 | 0 |
| | | | 1565 | 698 | 283 | 510 | 73 | 1 | | | |

- Molecule 55 is a RNA chain called P-site Peptidyl-tRNA fMRC-tRNAcys RNA-part.

| Mol | Chain | Residues | Atoms | | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|----|---|---------|---------|-------|
| 55 | 1x | 77 | Total | C | N | O | P | S | 0 | 0 | 0 |
| | | | 1646 | 734 | 298 | 536 | 77 | 1 | | | |
| 55 | 2x | 77 | Total | C | N | O | P | S | 0 | 0 | 0 |
| | | | 1646 | 734 | 298 | 536 | 77 | 1 | | | |

- Molecule 56 is a protein called P-site Peptidyl-tRNA fMRC-tRNAcys Peptide-part.

| Mol | Chain | Residues | Atoms | | | | | ZeroOcc | AltConf | Trace |
|-----|-------|----------|-------|----|---|---|---|---------|---------|-------|
| 56 | 1z | 3 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 27 | 15 | 6 | 4 | 2 | | | |
| 56 | 2z | 3 | Total | C | N | O | S | 0 | 0 | 0 |
| | | | 27 | 15 | 6 | 4 | 2 | | | |

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

| Mol | Chain | Residues | Atoms | | ZeroOcc | AltConf |
|-----|-------|----------|-------|------|---------|---------|
| 57 | 1A | 1085 | Total | Mg | 0 | 0 |
| | | | 1085 | 1085 | | |
| 57 | 1B | 38 | Total | Mg | 0 | 0 |
| | | | 38 | 38 | | |
| 57 | 1D | 12 | Total | Mg | 0 | 0 |
| | | | 12 | 12 | | |

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| Mol | Chain | Residues | Atoms | | ZeroOcc | AltConf |
|-----|-------|----------|-------------|----------|---------|---------|
| 57 | 1E | 13 | Total 13 | Mg 13 | 0 | 0 |
| 57 | 1F | 13 | Total 13 | Mg 13 | 0 | 0 |
| 57 | 1G | 4 | Total 4 | Mg 4 | 0 | 0 |
| 57 | 1H | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 1I | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 1N | 4 | Total 4 | Mg 4 | 0 | 0 |
| 57 | 1O | 4 | Total 4 | Mg 4 | 0 | 0 |
| 57 | 1P | 6 | Total 6 | Mg 6 | 0 | 0 |
| 57 | 1Q | 7 | Total 7 | Mg 7 | 0 | 0 |
| 57 | 1R | 6 | Total 6 | Mg 6 | 0 | 0 |
| 57 | 1S | 2 | Total 2 | Mg 2 | 0 | 0 |
| 57 | 1T | 3 | Total 3 | Mg 3 | 0 | 0 |
| 57 | 1U | 11 | Total 11 | Mg 11 | 0 | 0 |
| 57 | 1V | 7 | Total 7 | Mg 7 | 0 | 0 |
| 57 | 1W | 7 | Total 7 | Mg 7 | 0 | 0 |
| 57 | 1X | 7 | Total 7 | Mg 7 | 0 | 0 |
| 57 | 1Y | 2 | Total 2 | Mg 2 | 0 | 0 |
| 57 | 1Z | 2 | Total 2 | Mg 2 | 0 | 0 |
| 57 | 10 | 10 | Total 10 | Mg 10 | 0 | 0 |
| 57 | 11 | 3 | Total 3 | Mg 3 | 0 | 0 |
| 57 | 12 | 2 | Total 2 | Mg 2 | 0 | 0 |

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| Mol | Chain | Residues | Atoms | | ZeroOcc | AltConf |
|-----|-------|----------|--------------|-----------|---------|---------|
| 57 | 13 | 5 | Total 5 | Mg 5 | 0 | 0 |
| 57 | 14 | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 15 | 9 | Total 9 | Mg 9 | 0 | 0 |
| 57 | 16 | 2 | Total 2 | Mg 2 | 0 | 0 |
| 57 | 17 | 6 | Total 6 | Mg 6 | 0 | 0 |
| 57 | 18 | 7 | Total 7 | Mg 7 | 0 | 0 |
| 57 | 1a | 211 | Total 211 | Mg 211 | 0 | 0 |
| 57 | 1b | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 1d | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 1e | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 1f | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 1j | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 1l | 2 | Total 2 | Mg 2 | 0 | 0 |
| 57 | 1m | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 1n | 3 | Total 3 | Mg 3 | 0 | 0 |
| 57 | 1p | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 1r | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 1t | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 1w | 6 | Total 6 | Mg 6 | 0 | 0 |
| 57 | 1x | 12 | Total 12 | Mg 12 | 0 | 0 |
| 57 | 2A | 855 | Total 855 | Mg 855 | 0 | 0 |

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| Mol | Chain | Residues | Atoms | | ZeroOcc | AltConf |
|-----|-------|----------|-------------|----------|---------|---------|
| 57 | 2B | 20 | Total 20 | Mg 20 | 0 | 0 |
| 57 | 2D | 6 | Total 6 | Mg 6 | 0 | 0 |
| 57 | 2E | 9 | Total 9 | Mg 9 | 0 | 0 |
| 57 | 2F | 6 | Total 6 | Mg 6 | 0 | 0 |
| 57 | 2N | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 2O | 2 | Total 2 | Mg 2 | 0 | 0 |
| 57 | 2P | 2 | Total 2 | Mg 2 | 0 | 0 |
| 57 | 2Q | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 2R | 4 | Total 4 | Mg 4 | 0 | 0 |
| 57 | 2T | 3 | Total 3 | Mg 3 | 0 | 0 |
| 57 | 2U | 2 | Total 2 | Mg 2 | 0 | 0 |
| 57 | 2V | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 2W | 2 | Total 2 | Mg 2 | 0 | 0 |
| 57 | 2X | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 2Y | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 2Z | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 20 | 3 | Total 3 | Mg 3 | 0 | 0 |
| 57 | 21 | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 23 | 2 | Total 2 | Mg 2 | 0 | 0 |
| 57 | 25 | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 26 | 1 | Total 1 | Mg 1 | 0 | 0 |

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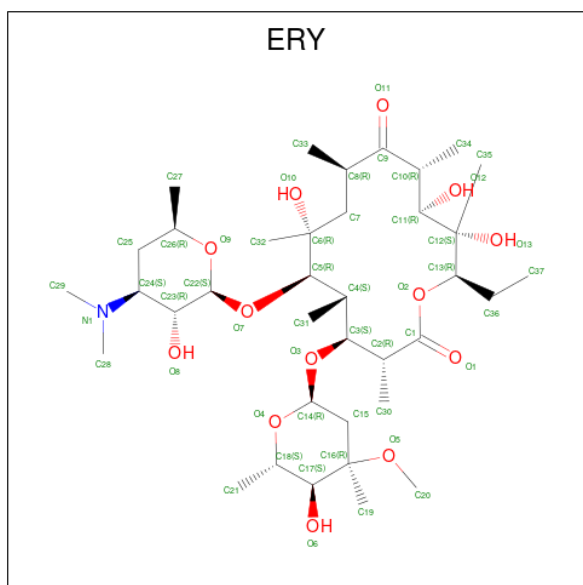
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| Mol | Chain | Residues | Atoms | | ZeroOcc | AltConf |
|-----|-------|----------|--------------|-----------|---------|---------|
| 57 | 27 | 4 | Total 4 | Mg 4 | 0 | 0 |
| 57 | 28 | 3 | Total 3 | Mg 3 | 0 | 0 |
| 57 | 2a | 225 | Total 225 | Mg 225 | 0 | 0 |
| 57 | 2d | 2 | Total 2 | Mg 2 | 0 | 0 |
| 57 | 2e | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 2f | 2 | Total 2 | Mg 2 | 0 | 0 |
| 57 | 2g | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 2j | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 2k | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 2l | 4 | Total 4 | Mg 4 | 0 | 0 |
| 57 | 2o | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 2q | 3 | Total 3 | Mg 3 | 0 | 0 |
| 57 | 2r | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 2t | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 2v | 2 | Total 2 | Mg 2 | 0 | 0 |
| 57 | 2w | 5 | Total 5 | Mg 5 | 0 | 0 |
| 57 | 2x | 6 | Total 6 | Mg 6 | 0 | 0 |
| 57 | 2z | 1 | Total 1 | Mg 1 | 0 | 0 |
| 57 | 2y | 1 | Total 1 | Mg 1 | 0 | 0 |

- Molecule 58 is POTASSIUM ION (CCD ID: K) (formula: K).

| Mol | Chain | Residues | Atoms | ZeroOcc | AltConf |
|-----|-------|----------|----------------|---------|---------|
| 58 | 1A | 1 | Total K 1 1 | 0 | 0 |
| 58 | 2A | 1 | Total K 1 1 | 0 | 0 |

- Molecule 59 is ERYTHROMYCIN A (CCD ID: ERY) (formula: $C_{37}H_{67}NO_{13}$) (labeled as "Ligand of Interest" by depositor).



| Mol | Chain | Residues | Atoms | ZeroOcc | AltConf |
|-----|-------|----------|---------------------------|---------|---------|
| 59 | 1A | 1 | Total C N O 51 37 1 13 | 0 | 0 |
| 59 | 2A | 1 | Total C N O 51 37 1 13 | 0 | 0 |

- Molecule 60 is ZINC ION (CCD ID: ZN) (formula: Zn).

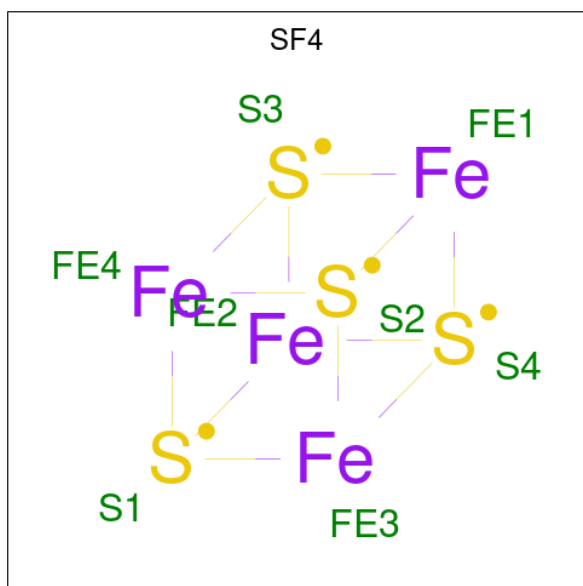
| Mol | Chain | Residues | Atoms | ZeroOcc | AltConf |
|-----|-------|----------|-----------------|---------|---------|
| 60 | 1Y | 1 | Total Zn 1 1 | 0 | 0 |
| 60 | 14 | 1 | Total Zn 1 1 | 0 | 0 |
| 60 | 15 | 1 | Total Zn 1 1 | 0 | 0 |
| 60 | 16 | 1 | Total Zn 1 1 | 0 | 0 |
| 60 | 19 | 1 | Total Zn 1 1 | 0 | 0 |

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| Mol | Chain | Residues | Atoms | | ZeroOcc | AltConf |
|-----|-------|----------|-------|----|---------|---------|
| 60 | 1n | 1 | Total | Zn | 0 | 0 |
| | | | 1 | 1 | | |
| 60 | 2Y | 1 | Total | Zn | 0 | 0 |
| | | | 1 | 1 | | |
| 60 | 24 | 1 | Total | Zn | 0 | 0 |
| | | | 1 | 1 | | |
| 60 | 25 | 1 | Total | Zn | 0 | 0 |
| | | | 1 | 1 | | |
| 60 | 26 | 1 | Total | Zn | 0 | 0 |
| | | | 1 | 1 | | |
| 60 | 29 | 1 | Total | Zn | 0 | 0 |
| | | | 1 | 1 | | |
| 60 | 2n | 1 | Total | Zn | 0 | 0 |
| | | | 1 | 1 | | |

- Molecule 61 is IRON/SULFUR CLUSTER (CCD ID: SF4) (formula: Fe_4S_4).



| Mol | Chain | Residues | Atoms | | | ZeroOcc | AltConf |
|-----|-------|----------|-------|----|---|---------|---------|
| 61 | 1d | 1 | Total | Fe | S | 0 | 0 |
| | | | 8 | 4 | 4 | | |
| 61 | 2d | 1 | Total | Fe | S | 0 | 0 |
| | | | 8 | 4 | 4 | | |

- Molecule 62 is water.

| Mol | Chain | Residues | Atoms | | ZeroOcc | AltConf |
|-----|-------|----------|---------------|-----------|---------|---------|
| 62 | 1A | 1680 | Total 1680 | O 1680 | 0 | 0 |
| 62 | 1B | 54 | Total 54 | O 54 | 0 | 0 |
| 62 | 1D | 32 | Total 32 | O 32 | 0 | 0 |
| 62 | 1E | 23 | Total 23 | O 23 | 0 | 0 |
| 62 | 1F | 17 | Total 17 | O 17 | 0 | 0 |
| 62 | 1G | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 1H | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 1N | 4 | Total 4 | O 4 | 0 | 0 |
| 62 | 1O | 3 | Total 3 | O 3 | 0 | 0 |
| 62 | 1P | 15 | Total 15 | O 15 | 0 | 0 |
| 62 | 1Q | 8 | Total 8 | O 8 | 0 | 0 |
| 62 | 1R | 12 | Total 12 | O 12 | 0 | 0 |
| 62 | 1S | 3 | Total 3 | O 3 | 0 | 0 |
| 62 | 1T | 6 | Total 6 | O 6 | 0 | 0 |
| 62 | 1U | 8 | Total 8 | O 8 | 0 | 0 |
| 62 | 1V | 9 | Total 9 | O 9 | 0 | 0 |
| 62 | 1W | 9 | Total 9 | O 9 | 0 | 0 |
| 62 | 1X | 6 | Total 6 | O 6 | 0 | 0 |
| 62 | 1Y | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 1Z | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 10 | 8 | Total 8 | O 8 | 0 | 0 |
| 62 | 11 | 7 | Total 7 | O 7 | 0 | 0 |

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| Mol | Chain | Residues | Atoms | | ZeroOcc | AltConf |
|-----|-------|----------|--------------|----------|---------|---------|
| 62 | 12 | 3 | Total 3 | O 3 | 0 | 0 |
| 62 | 13 | 4 | Total 4 | O 4 | 0 | 0 |
| 62 | 14 | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 15 | 6 | Total 6 | O 6 | 0 | 0 |
| 62 | 16 | 3 | Total 3 | O 3 | 0 | 0 |
| 62 | 17 | 10 | Total 10 | O 10 | 0 | 0 |
| 62 | 18 | 8 | Total 8 | O 8 | 0 | 0 |
| 62 | 1a | 252 | Total 252 | O 252 | 0 | 0 |
| 62 | 1b | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 1d | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 1f | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 1l | 3 | Total 3 | O 3 | 0 | 0 |
| 62 | 1m | 2 | Total 2 | O 2 | 0 | 0 |
| 62 | 1o | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 1p | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 1q | 2 | Total 2 | O 2 | 0 | 0 |
| 62 | 1u | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 1v | 3 | Total 3 | O 3 | 0 | 0 |
| 62 | 1w | 8 | Total 8 | O 8 | 0 | 0 |
| 62 | 1x | 7 | Total 7 | O 7 | 0 | 0 |
| 62 | 1z | 1 | Total 1 | O 1 | 0 | 0 |

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| Mol | Chain | Residues | Atoms | | ZeroOcc | AltConf |
|-----|-------|----------|--------------|----------|---------|---------|
| 62 | 1y | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 2A | 952 | Total 952 | O 952 | 0 | 0 |
| 62 | 2B | 21 | Total 21 | O 21 | 0 | 0 |
| 62 | 2D | 15 | Total 15 | O 15 | 0 | 0 |
| 62 | 2E | 5 | Total 5 | O 5 | 0 | 0 |
| 62 | 2F | 13 | Total 13 | O 13 | 0 | 0 |
| 62 | 2O | 3 | Total 3 | O 3 | 0 | 0 |
| 62 | 2P | 6 | Total 6 | O 6 | 0 | 0 |
| 62 | 2Q | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 2R | 4 | Total 4 | O 4 | 0 | 0 |
| 62 | 2T | 2 | Total 2 | O 2 | 0 | 0 |
| 62 | 2U | 2 | Total 2 | O 2 | 0 | 0 |
| 62 | 2V | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 2W | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 2X | 5 | Total 5 | O 5 | 0 | 0 |
| 62 | 20 | 2 | Total 2 | O 2 | 0 | 0 |
| 62 | 21 | 6 | Total 6 | O 6 | 0 | 0 |
| 62 | 23 | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 25 | 2 | Total 2 | O 2 | 0 | 0 |
| 62 | 27 | 2 | Total 2 | O 2 | 0 | 0 |
| 62 | 28 | 3 | Total 3 | O 3 | 0 | 0 |

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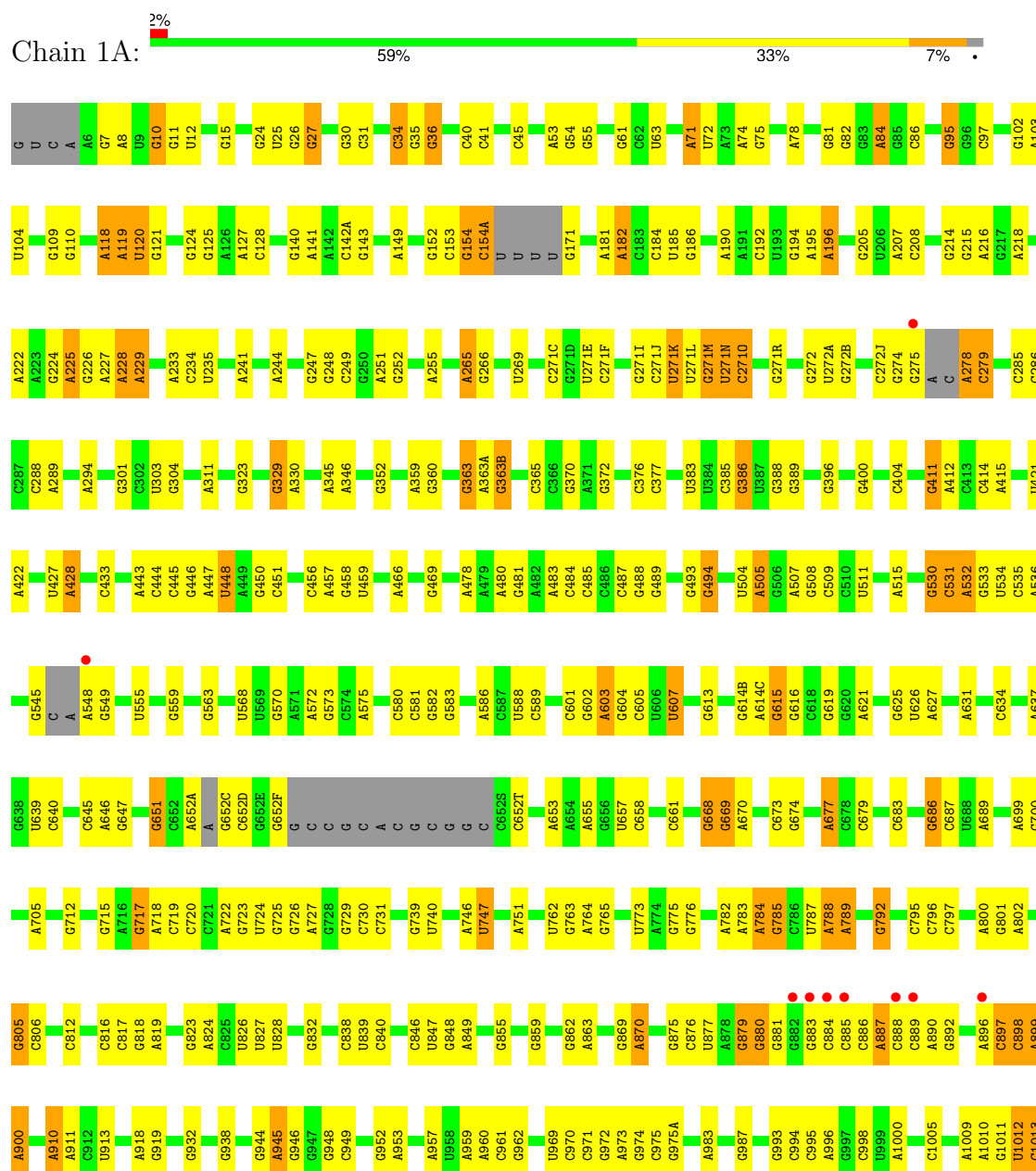
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| Mol | Chain | Residues | Atoms | | ZeroOcc | AltConf |
|-----|-------|----------|--------------|----------|---------|---------|
| 62 | 29 | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 2a | 150 | Total 150 | O 150 | 0 | 0 |
| 62 | 2e | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 2j | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 2l | 4 | Total 4 | O 4 | 0 | 0 |
| 62 | 2q | 1 | Total 1 | O 1 | 0 | 0 |
| 62 | 2v | 3 | Total 3 | O 3 | 0 | 0 |
| 62 | 2w | 3 | Total 3 | O 3 | 0 | 0 |
| 62 | 2x | 4 | Total 4 | O 4 | 0 | 0 |
| 62 | 2z | 1 | Total 1 | O 1 | 0 | 0 |

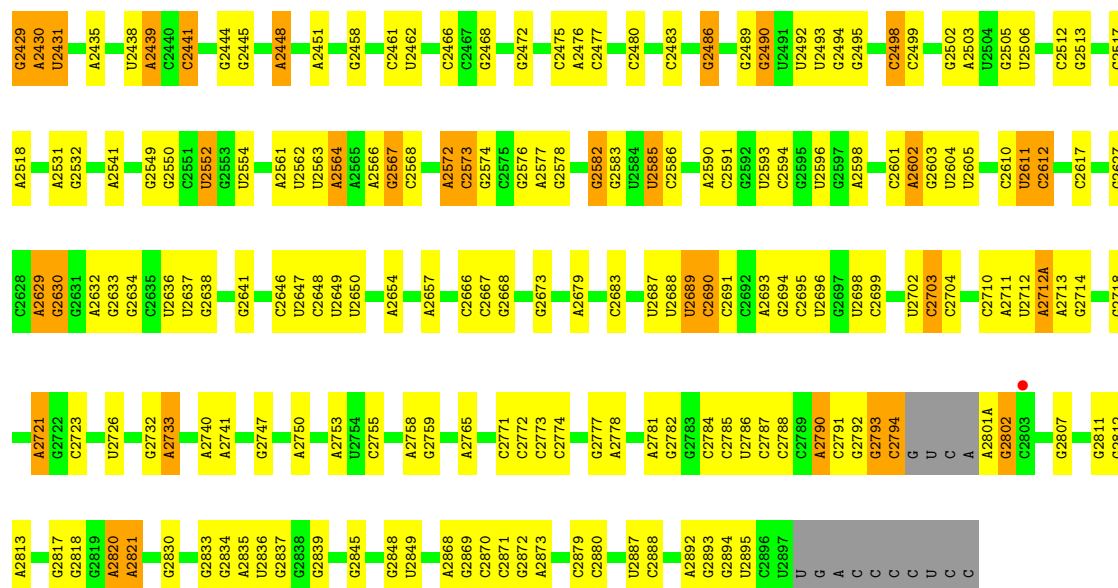
3 Residue-property plots

These plots are drawn for all protein, RNA, DNA and oligosaccharide chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes 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.

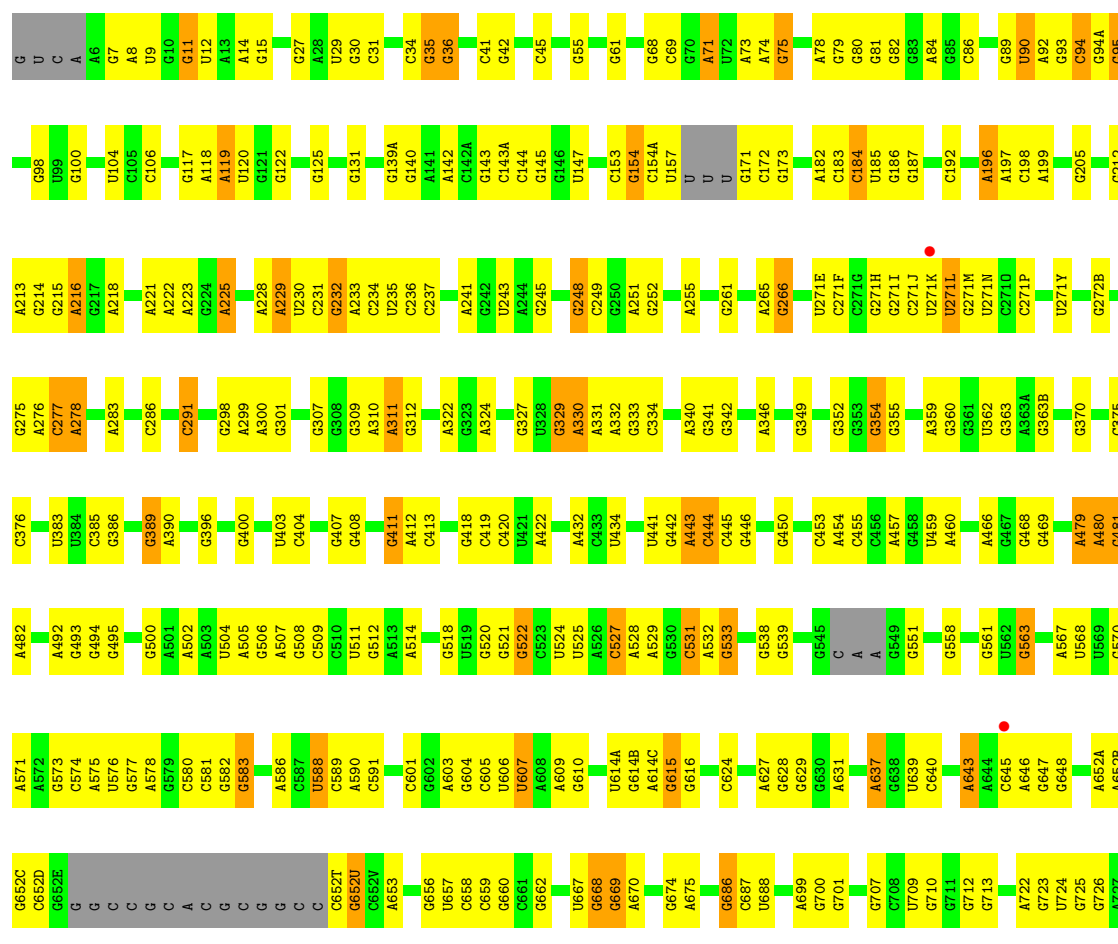
• Molecule 1: 23S Ribosomal RNA



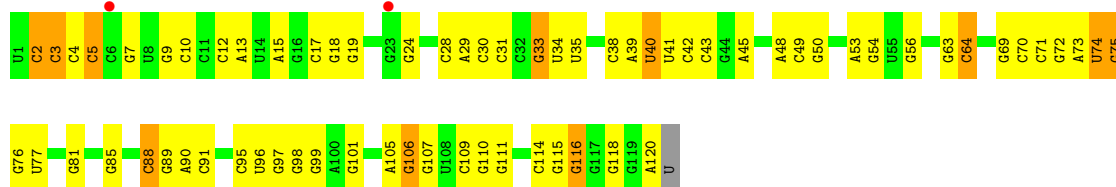
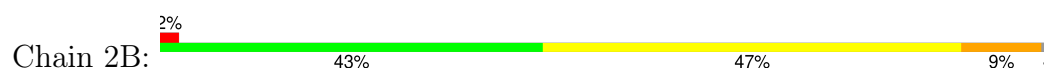
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|--------|--------|-------|-------|-------|-------|-------|-------|--------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|
| G2331 | G2334 | G2335 | G2336 | G2337 | G2341 | G2342 | G2343 | G2344 | G2351 | G2357 | G2361 | G2362 | G2363 | G2364 | G2365 | G2371 | G2372 | G2375 | G2376 | G2377 | G2378 | G2379 | G2383 | G2384 | G2385 | G2390 | G2393 | G2396 | G2404 | G2405 | G2406 | G2407 | G2417 | G2418 | G2419 | G2422 | G2423 | G2424 | G2425 | G2426 | G2427 | G2428 | | | | | | | | | | | | |
| C2231 | G2238 | A2235 | G2239 | G2237 | U2243 | U2244 | U2245 | G2246 | G2251 | G2257 | G2262 | G2263 | G2264 | U2265 | G2266 | A2267 | G2268 | G2269 | G2270 | G2271 | G2272 | G2273 | G2274 | G2275 | G2276 | G2277 | G2278 | G2279 | G2280 | G2281 | G2282 | G2283 | A2286 | A2287 | U2291 | G2292 | U2296 | A2305 | G2306 | G2307 | G2308 | A2309 | G2313 | G2314 | G2316 | G2319 | G2320 | G2325 | G2326 | A2327 | A2328 | G2329 | G2330 | |
| U2150 | G2151 | G2152 | G2153 | G2154 | G2155 | G2156 | G2157 | A2158 | G2159 | G2160 | G2161 | G2162 | G2163 | G2164 | G2165 | G2166 | G2167 | G2168 | A2169 | A2170 | A2171 | U2172 | A2173 | A2174 | G2175 | G2176 | G2177 | G2178 | G2179 | G2180 | G2181 | G2182 | G2183 | G2184 | G2187 | G2188 | U2189 | G2190 | G2191 | G2192 | G2193 | A2198 | G2206 | G2207 | A2208 | U2218 | G2219 | G2222 | G2223 | G2224 | A2225 | A2226 | A2227 | G2228 |
| A1970 | A1971 | A1972 | G1975 | G1976 | A1986 | G1992 | U1993 | G1996 | G1997 | G1998 | G1999 | G2000 | A2001 | G2002 | G2010 | G2017 | G2108 | G2109 | G2110 | G2111 | G2112 | A2113 | G2114 | G2115 | G2116 | A2117 | G2118 | A2119 | G2120 | G2121 | U2122 | A2126 | G2127 | G2128 | G2129 | U2130 | G2131 | U2132 | G2133 | A2134 | A2135 | G2207 | G2208 | G2218 | G2219 | G2222 | G2223 | G2224 | A2225 | A2226 | A2227 | G2228 | | |
| G1861 | G1862 | G1865 | G1866 | A1876 | A1877 | G1878 | G1881 | A1889 | A1890 | G1900 | G1906 | U1911 | G1914 | U1915 | A1916 | U1917 | A1918 | A1919 | G1920 | G1929 | U1930 | U1931 | A1932 | G1933 | A1937 | A1938 | U1939 | G1942 | U1943 | G1949 | G1950 | A2042 | C2043 | G1952 | A1953 | G1954 | U1955 | U1956 | C1957 | G1958 | G1962 | U1963 | G1964 | A1967 | G1968 | A1969 | | | | | | | | |
| G1772 | A1773 | U1778 | U1779 | A1780 | G1781 | G1782 | A1783 | A1784 | A1785 | A1786 | C1790 | A1791 | U1794 | G1795 | U1796 | G1797 | U1798 | G1799 | C1800 | G1801 | A1802 | A1803 | G1804 | U1805 | G1810 | G1811 | G1814 | A1815 | G1816 | G1817 | U1818 | A1819 | U1820 | G1823 | G1824 | A1825 | G1826 | C1827 | U1833 | U1837 | C1843 | G1846 | A1847 | A1848 | A1853 | G1857 | G1858 | | | | | | | |
| G1647 | C1648 | A1655 | C1656 | C1662 | C1663 | A1664 | A1665 | A1669 | C1670 | U1671 | C1672 | G1674 | U1680 | C1683 | C1684 | C1685 | C1686 | G1687 | U1688 | C1691 | U1692 | U1693 | C1694 | G1695 | G1696 | A1700 | A1701 | G1702 | G1703 | U1713 | G1714 | U1720 | G1721 | A1722 | U1739 | G1740 | A1741 | G1746 | G1756 | A1762 | G1763 | G1764 | G1769 | C1770 | G1771 | | | | | | | | | |
| U | A | C | G1537 | U1540 | G1541 | A1542 | C1543 | A1554 | C1557 | A1558 | G1559 | A1566 | A1569 | U1578 | A1579 | A1580 | G1581 | C1584 | A1586 | A1587 | C1588 | C1589 | G1594 | G1595 | U1602 | A1603 | C1604 | C1607 | A1608 | A1609 | A1610 | A1614 | C1617 | A1618 | G1619 | G1627 | G1633 | A1634 | G1635 | C1636 | A1637 | G1643 | C1644 | | | | | | | | | | | |
| G1424 | G1425 | G1426 | A1427 | C1428 | A1439 | G1442 | A1445 | G1448 | A1449 | G1450 | A1452 | U1453 | G1455 | A1467 | C1468 | G1482 | G1484 | A1490 | C1493 | A1494 | A1495 | A1496 | U1497 | G1500 | C1501 | C1504 | C1505 | A1508 | C1509 | A1509A | A1509B | G1510 | U1512 | C1513 | U1518 | G1519 | G1520 | A1528A | G1529 | C1530 | C1531 | C1532 | G1533 | | | | | | | | | | | |
| G1319 | C1320 | A1321 | A1322 | U1326 | C1327 | G1332 | C1333 | A1342 | G1343 | G1344 | C1345 | U1352 | G1355 | G1356 | U1357 | G1358 | A1359 | A1360 | A1365 | C1370 | G1371 | U1372 | A1373 | G1374 | C1375 | C1376 | A1379 | G1380 | A1384 | G1385 | C1386 | U1391 | A1395 | U1396 | U1405 | U1406 | C1407 | C1411 | A1412 | G1416 | G1418 | A1419 | U1420 | C1421 | | | | | | | | | | |
| G1173 | A1174 | U1175 | G1176 | A1177 | C1178 | C1179 | G1186 | G1187 | C1201 | G1209 | A1213 | C1218 | G1223 | A1226 | G1235 | G1236 | A1237 | G1238 | G1239 | C1244 | A1253 | G1256 | U1263 | G1266 | U1267 | A1268 | A1269 | C1270 | G1271 | A1272 | U1273 | G1278 | G1283 | A1284 | G1285 | U1300 | A1301 | A1302 | G1303 | C1315 | G1318 | G1319 | C1316 | | | | | | | | | | | |
| A1088 | G1089 | U1090 | G1091 | C1092 | G1093 | U1094 | A1095 | A1096 | U1097 | A1098 | G1099 | C1100 | U1101 | C1102 | A1103 | C1104 | U1105 | G1106 | G1107 | U1108 | C1109 | G1110 | A1111 | G1112 | U1113 | G1114 | C1115 | C1116 | G1125 | U1130 | U1133 | C1135 | G1136 | G1137 | G1138 | G1139 | C1140 | U1141 | U1142 | G1144 | C1145 | G1149 | C1153 | A1156 | G1164 | U1165 | C1166 | U1167 | G1170 | G1171 | | | | |



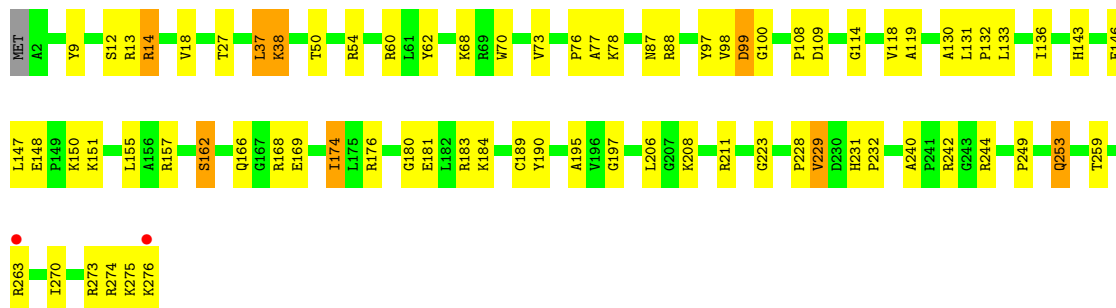
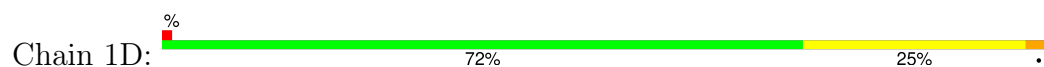
• Molecule 1: 23S Ribosomal RNA



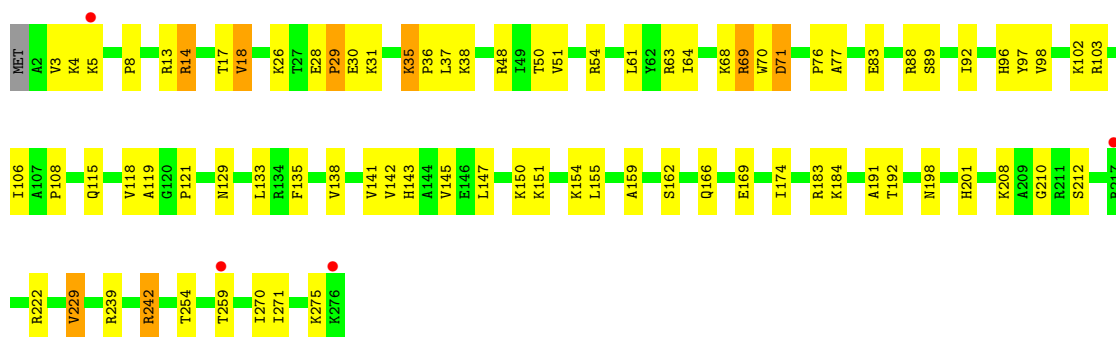
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|-------|-------|--------|-------|--------|--------|-------|-------|--------|---|-------|------|------|------|
| C1942 | G1826 | G1721 | G1628 | G1527 | G1427 | A1317 | G1263 | U1130 | A | A996 | U913 | U828 | G728 |
| A1952 | C1827 | A1722 | U1629 | G1528 | C1428 | C1320 | A1204 | G1131 | G | U999 | C914 | A829 | G729 |
| U1955 | G1828 | U1739 | G1633 | A1528A | U1431 | G1324 | U1205 | A1132 | A | G | A917 | U839 | C730 |
| G1959 | C1837 | G1740 | A1634 | C1529 | C1432 | G1325 | G1206 | G1135 | G | G1003 | A918 | C840 | C731 |
| C1962 | U1847 | A1741 | G1635 | C1530 | U1433 | G1326 | A1210 | G1136 | C | G1004 | C924 | U847 | G733 |
| U1963 | A1848 | G1747 | C1636 | C1531 | A1434 | U1326 | U1211 | G1139 | A | C1005 | C925 | U747 | G748 |
| C1967 | A1849 | G1747A | A1637 | C1532 | G1435 | C1327 | U1212 | G1140 | G | C1006 | C926 | C850 | |
| | A1853 | C1754 | G1638 | G1533 | G1436 | U1340 | C1218 | U1141 | C | U1012 | A926 | U851 | A751 |
| | A1854 | U1755 | U1639 | A | C1437 | U1340 | G1219 | U1142 | A | C1013 | G927 | C852 | A752 |
| | G1855 | C1756 | C1640 | C1536 | G1442 | U1352 | A1220 | A1142A | U | G1017 | G928 | C853 | C753 |
| A1970 | A1856 | U1757 | G1645 | G1539 | G1443 | U1353 | G1223 | G1149 | C | G1018 | C856 | C857 | A764 |
| A1972 | G1857 | G1758 | C1646 | U1540 | G1444 | A1354 | G1224 | C1150 | C | U1019 | C857 | C858 | |
| | G1858 | A1762 | G1647 | A1541 | A1445 | G1356 | G1225 | G1151 | U | G938 | U858 | U859 | U773 |
| | C1866 | G1763 | G1649 | A1542 | G1448 | A1359 | G1226 | C1152 | U | G939 | C859 | U860 | A774 |
| G1979 | A1876 | A1764 | A1652 | C1546 | A1449 | A1360 | G1228 | C1153 | A | G1022 | G940 | U861 | A775 |
| G1980 | A1877 | | G1653 | C1547 | A1450 | G1367 | G1229 | G1156 | A | U1023 | A941 | C862 | G776 |
| A1986 | G1878 | A1773 | A1654 | C1547 | G1450 | A1368 | G1230 | A1157 | A | G1024 | G942 | A863 | |
| U1991 | C1882 | G1776 | C1660 | C1550 | A1452 | G1364 | G1231 | C1158 | G | G1025 | U943 | U868 | U779 |
| G1992 | G1883 | U1777 | G1661 | C1557 | U1453 | A1366 | G1231 | U1159 | A | U1026 | G944 | C869 | G780 |
| U1993 | C1886 | U1778 | G1666 | A1558 | G1455 | A1367 | G1235 | G1160 | G | A1027 | G945 | U871 | A781 |
| | | A1780 | G1667 | A1562 | A1460 | G1369 | G1236 | G1164 | U | A1028 | G946 | C874 | A782 |
| | A1889 | C1781 | G1667 | A1562 | G1461 | G1370 | A1237 | U1165 | C | U1029 | G947 | A783 | A784 |
| A1890 | A1890 | G1782 | C1670 | A1566 | G1466 | G1371 | G1238 | G1166 | G | U1033 | G948 | A785 | |
| G1999 | C1999 | A1783 | U1671 | A1566 | C1467 | U1372 | G1239 | U1167 | U | G1034 | A953 | C879 | G786 |
| G2000 | G1899 | G1784 | U1671 | A1569 | G1467 | A1373 | U1241 | G1168 | A | U1035 | G954 | C880 | C786 |
| A2001 | A1900 | A1785 | G1674 | A1569 | C1468 | A1373 | A1241 | G1169 | U | G1036 | C955 | C881 | C787 |
| G2002 | | A1786 | G1674 | A1569 | G1468 | A1373 | A1241 | G1170 | A | G1037 | G956 | C882 | A788 |
| | | A1787 | G1674 | A1578 | A1469 | A1379 | G1243 | G1171 | A | C1038 | A957 | C883 | A789 |
| C2006 | G1906 | C1788 | U1680 | A1579 | G1470 | G1380 | G1248 | G | C | G1039 | U958 | C884 | |
| G2010 | C1908 | A1789 | G1688 | A1580 | A1472 | A1384 | A1253 | A | C | C1040 | C960 | C885 | G792 |
| | | C1790 | A1689 | C1581 | G1473 | G1385 | A1253 | U | C | G1041 | C961 | C886 | A793 |
| A2014 | U1911 | A1791 | A1689 | C1582 | G1473 | G1386 | G1256 | G | A | G1042 | G962 | C887 | |
| | A1912 | U1794 | U1693 | A1583 | A1477 | C1386 | G1256 | A | C | C1043 | U963 | C888 | C797 |
| U2017 | C1913 | C1795 | C1694 | C1584 | G1478 | A1392 | G1264 | C1178 | U | G | C970 | C889 | G798 |
| G2018 | U1914 | U1796 | G1695 | A1586 | G1478 | A1393 | G1264 | C1179 | G | A | C971 | A890 | G799 |
| A2019 | U1915 | C1797 | G1696 | A1587 | G1482 | U1394 | A1268 | C1180 | G | A | G972 | C891 | A800 |
| | A1916 | U1798 | G1697 | C1588 | A1490 | A1395 | A1268 | C1181 | U | A | C973 | C892 | A802 |
| U2022 | U1917 | G1799 | A1698 | G1593 | A1490 | U1396 | G1271 | A1182 | C | A | C974 | C893 | |
| G2023 | A1918 | C1800 | G1699 | G1594 | C1493 | G1401 | A1272 | G1183 | G | A | C975 | U895 | G805 |
| G2024 | A1919 | A1803 | A1700 | G1595 | U1497 | C1402 | U1273 | G1184 | A | G | C976 | A896 | C806 |
| C2025 | C1920 | C1804 | G1702 | A1596 | U1497 | U1406 | A1278 | C1185 | C | C | G977 | C897 | U807 |
| C2026 | G1924 | U1805 | G1703 | C1600 | C1505 | C1407 | G1286 | G1186 | C | C | G978 | C898 | G808 |
| G2027 | | | | C1607 | G1506 | C1407 | G1286 | G1187 | C | C | G979 | A899 | U809 |
| A2030 | G1929 | A1812 | C1708 | A1607 | A1507 | G1416 | U1300 | U1188 | U | A | A980 | A900 | U810 |
| A2031 | G1930 | G1813 | U1709 | A1608 | A1508 | C1417 | A1301 | G1191 | A | G | A981 | A901 | U811 |
| G2032 | | A1814 | C1710 | A1609 | C1509 | G1418 | A1301 | G1192 | G | A | C982 | C902 | C812 |
| A2033 | G1935 | A1815 | C1711 | A1610 | A1509A | G1419 | A1306 | G1193 | G | G | A983 | C903 | C815 |
| G2034 | A1936 | G1816 | G1712 | A1610 | A1509A | U1420 | G1306 | A1194 | U | U | C985 | C904 | C816 |
| G2035 | A1937 | G1817 | U1713 | G1613 | U1514 | G1421 | U1313 | G1195 | G | A990 | C986 | U907 | C817 |
| C2036 | A1938 | U1818 | G1714 | G1515 | G1515 | G1422 | C1314 | G1199 | G | C | C987 | C908 | G818 |
| G2037 | U1939 | G1824 | G1719 | G1518 | U1518 | G1423 | C1315 | C1200 | C | G | A989 | A909 | A819 |
| G2038 | A1940 | A1825 | U1720 | G1622 | U1518 | G1424 | U1316 | G1125 | U | C | C993 | A910 | U826 |
| C2039 | | | | G1627 | G1519 | | | C1202 | U | U | C995 | C912 | U827 |



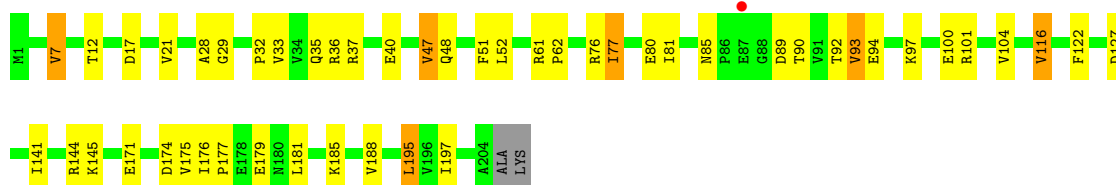
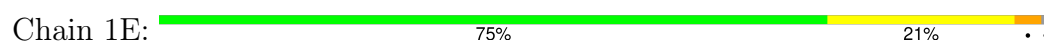
• Molecule 3: 50S ribosomal protein L2



• Molecule 3: 50S ribosomal protein L2

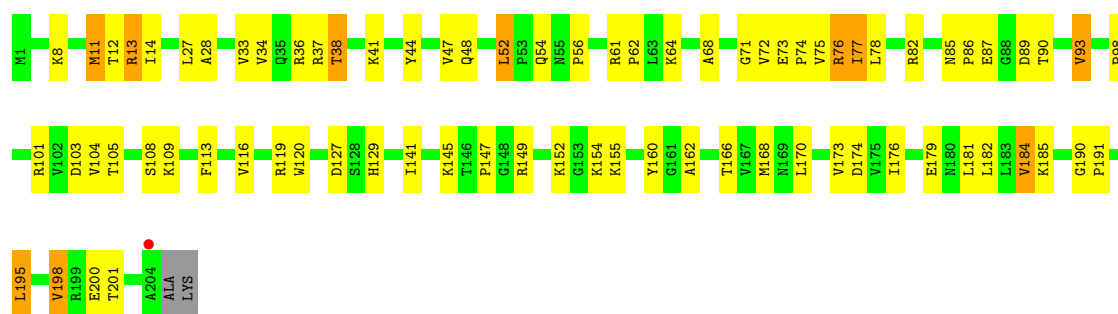


• Molecule 4: 50S ribosomal protein L3



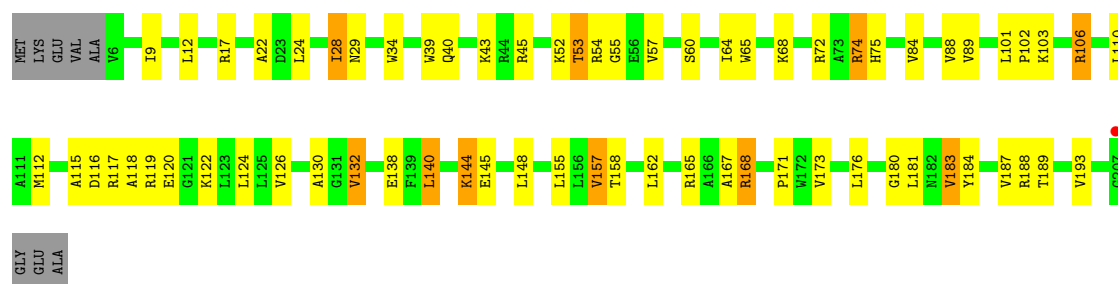
• Molecule 4: 50S ribosomal protein L3

Chain 2E: 



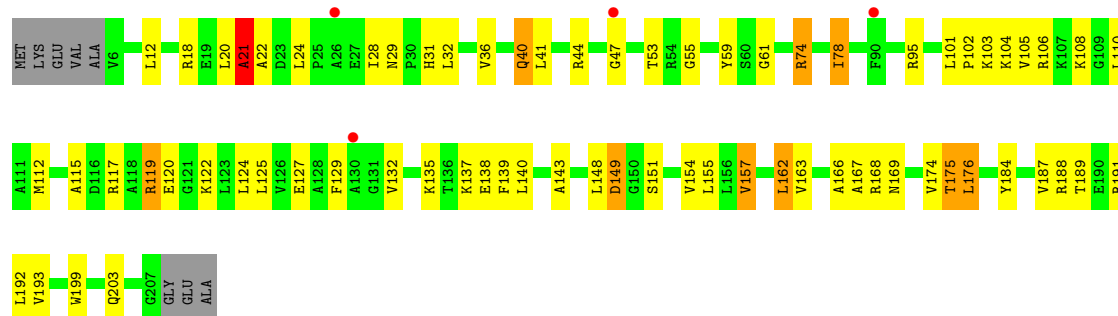
• Molecule 5: 50S ribosomal protein L4

Chain 1F: 



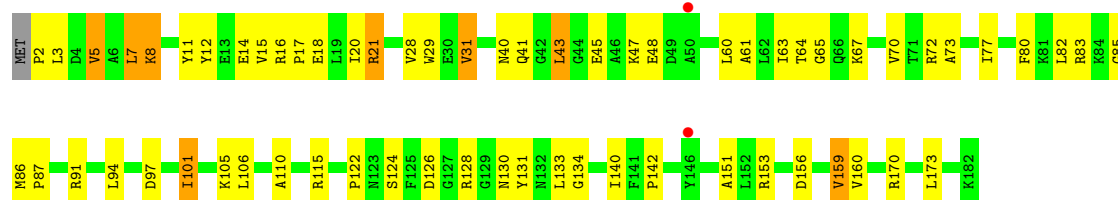
• Molecule 5: 50S ribosomal protein L4

Chain 2F: 

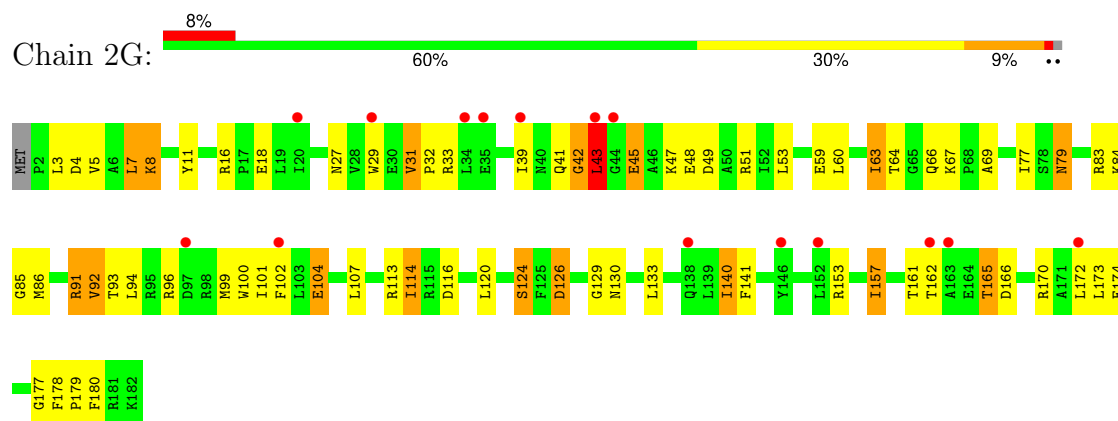


• Molecule 6: 50S ribosomal protein L5

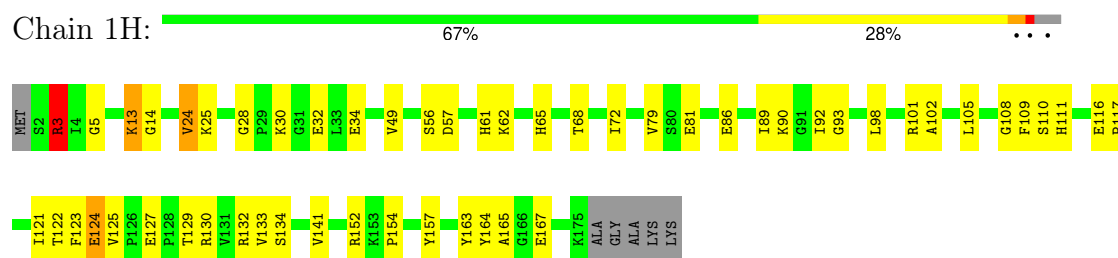
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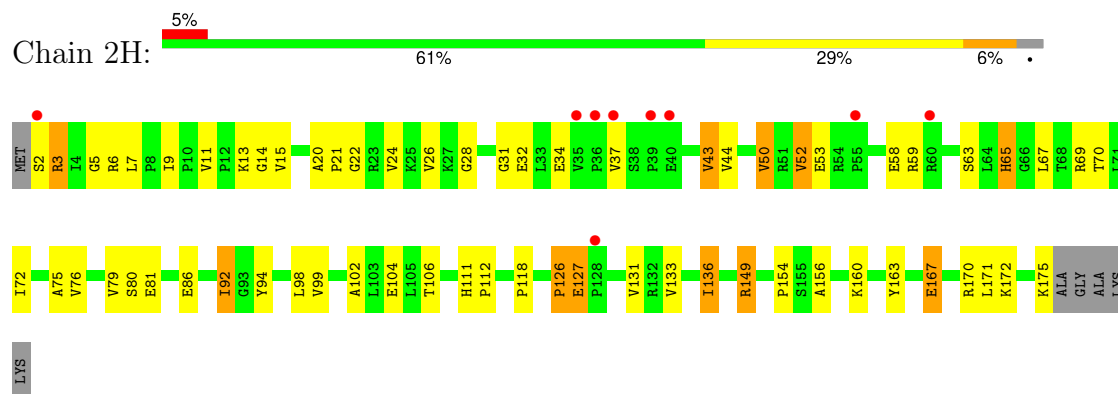
- Molecule 6: 50S ribosomal protein L5



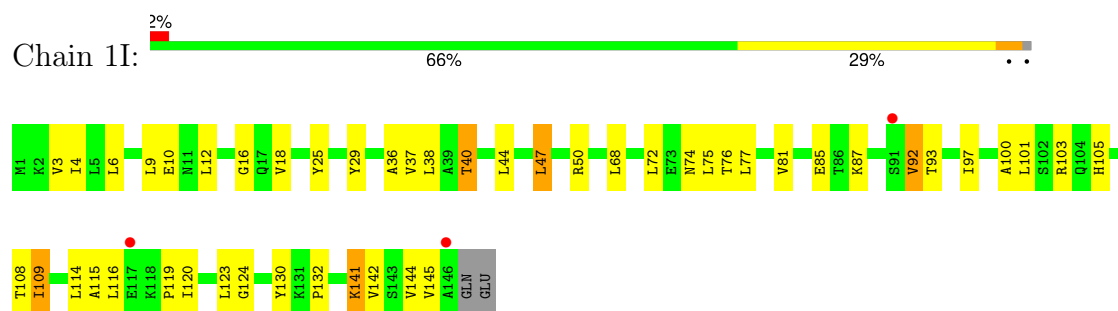
- Molecule 7: 50S ribosomal protein L6



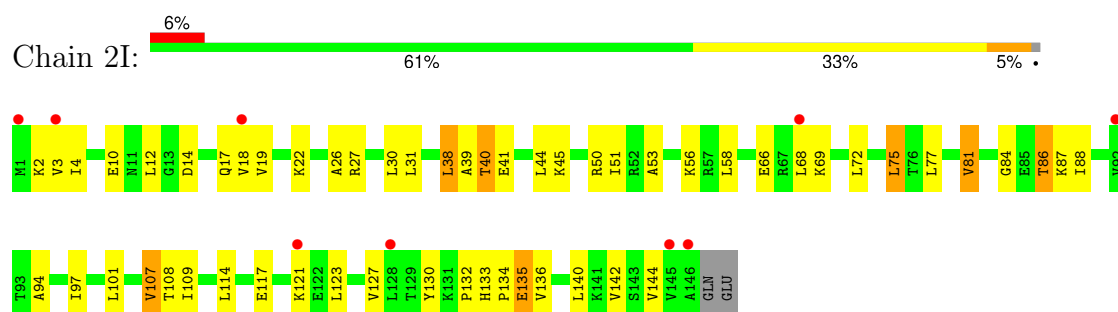
- Molecule 7: 50S ribosomal protein L6



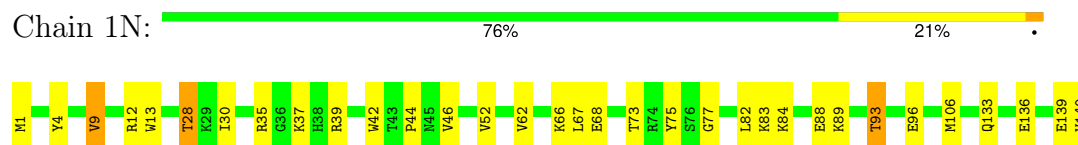
- Molecule 8: 50S ribosomal protein L9



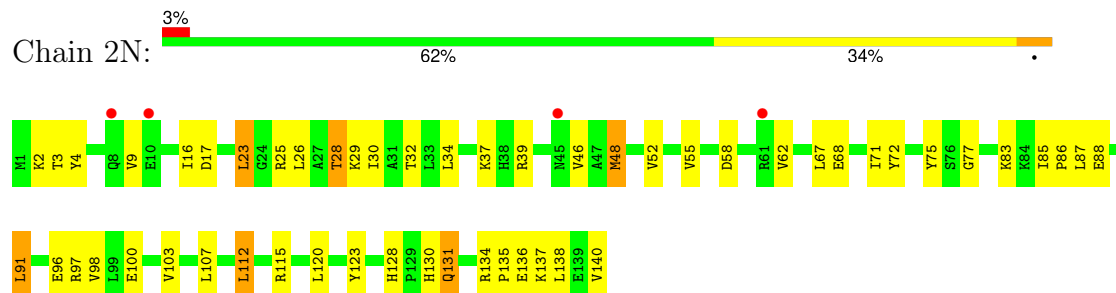
- Molecule 8: 50S ribosomal protein L9



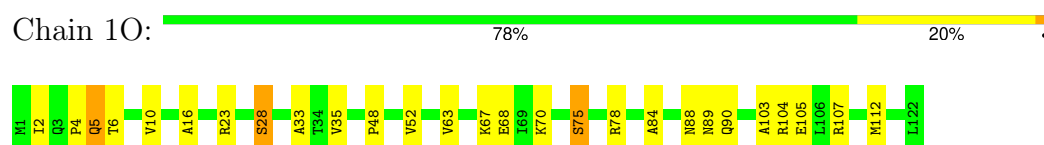
- Molecule 9: 50S ribosomal protein L13



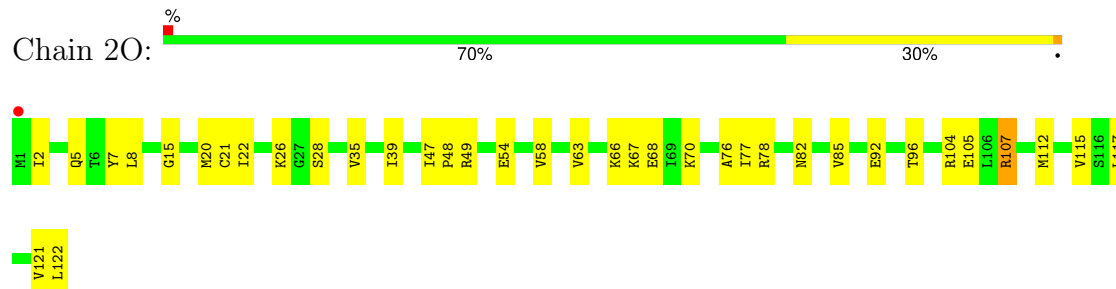
- Molecule 9: 50S ribosomal protein L13



- Molecule 10: 50S ribosomal protein L14

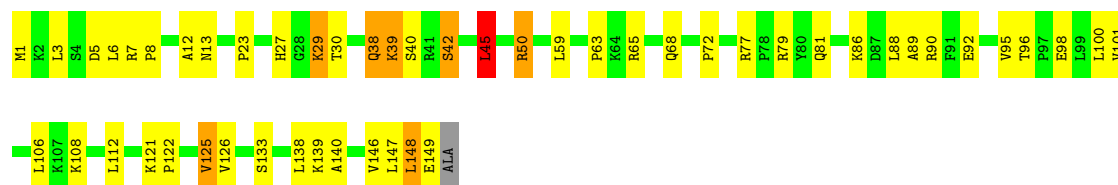


- Molecule 10: 50S ribosomal protein L14

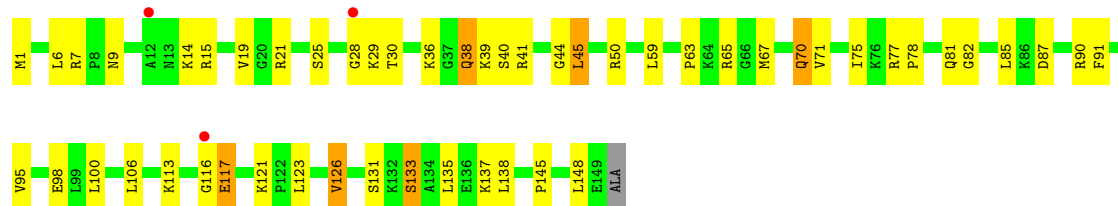


- Molecule 11: 50S ribosomal protein L15

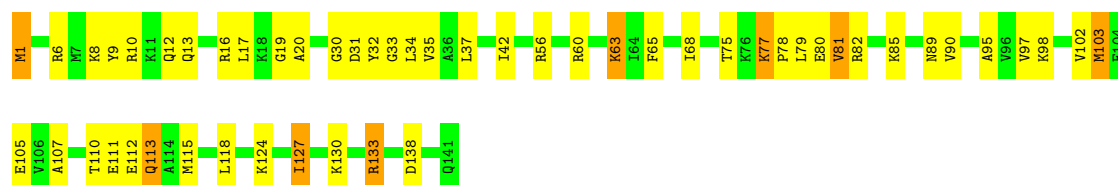




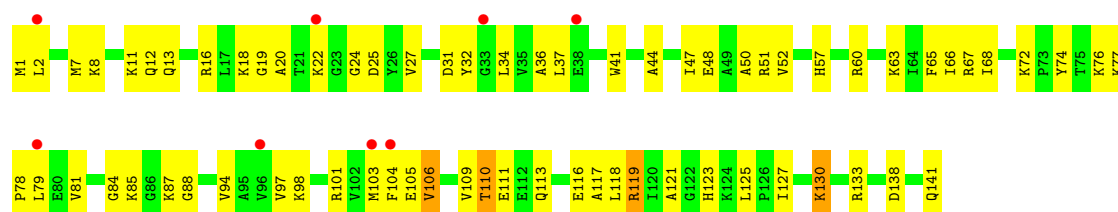
- Molecule 11: 50S ribosomal protein L15



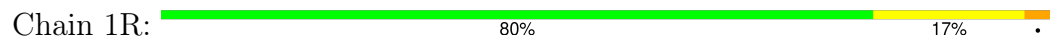
- Molecule 12: 50S ribosomal protein L16



- Molecule 12: 50S ribosomal protein L16

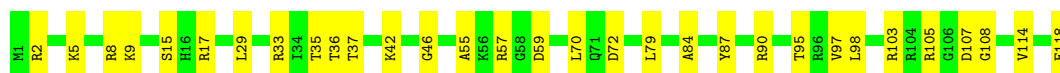


- Molecule 13: 50S ribosomal protein L17

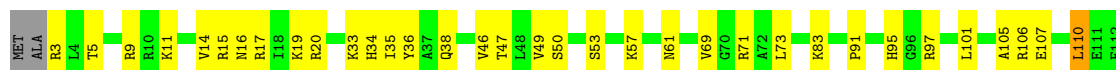


- Molecule 13: 50S ribosomal protein L17

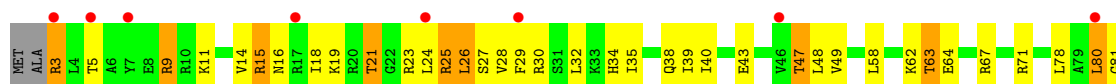




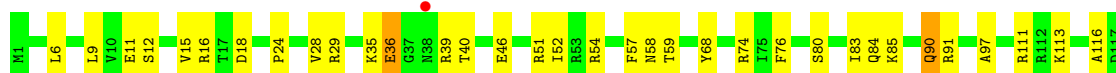
• Molecule 14: 50S ribosomal protein L18



• Molecule 14: 50S ribosomal protein L18



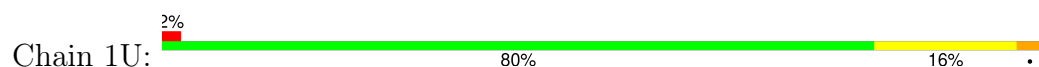
• Molecule 15: 50S ribosomal protein L19



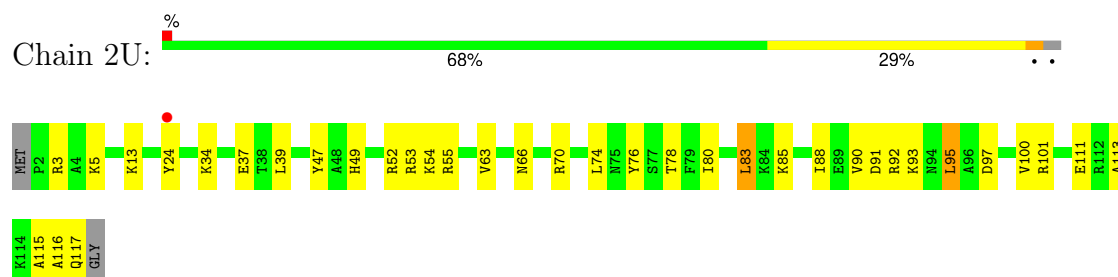
• Molecule 15: 50S ribosomal protein L19



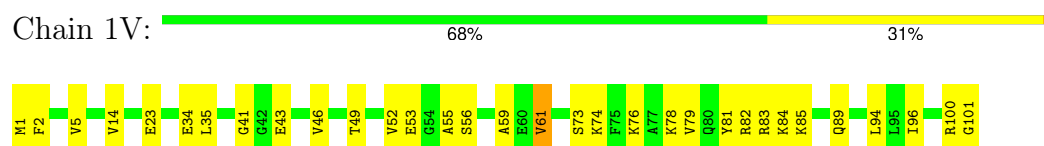
• Molecule 16: 50S ribosomal protein L20



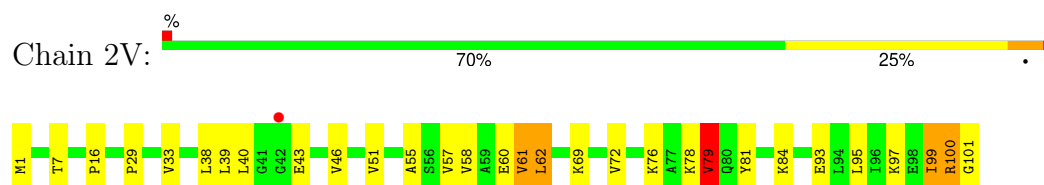
- Molecule 16: 50S ribosomal protein L20



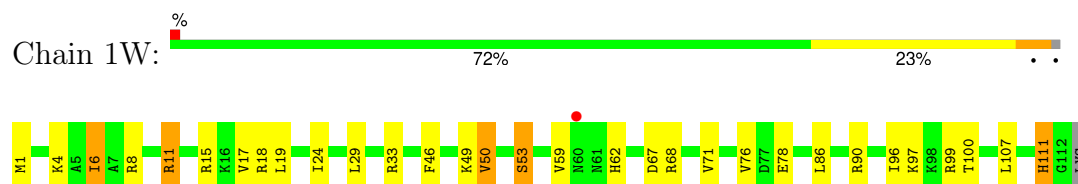
- Molecule 17: 50S ribosomal protein L21



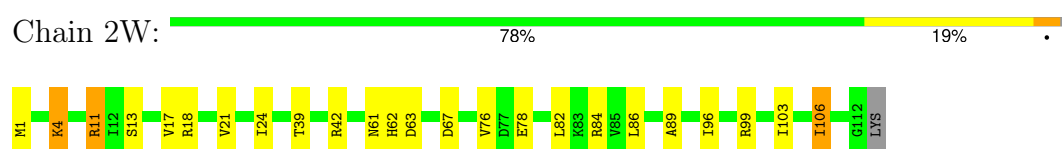
- Molecule 17: 50S ribosomal protein L21



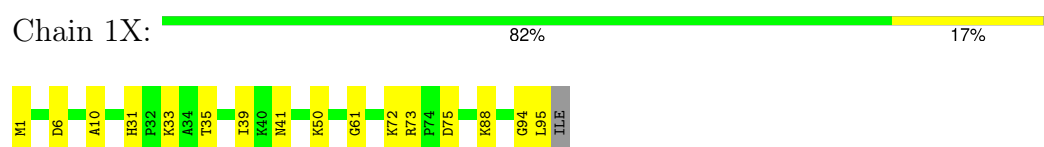
- Molecule 18: 50S ribosomal protein L22



- Molecule 18: 50S ribosomal protein L22

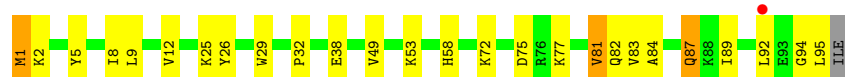


- Molecule 19: 50S ribosomal protein L23

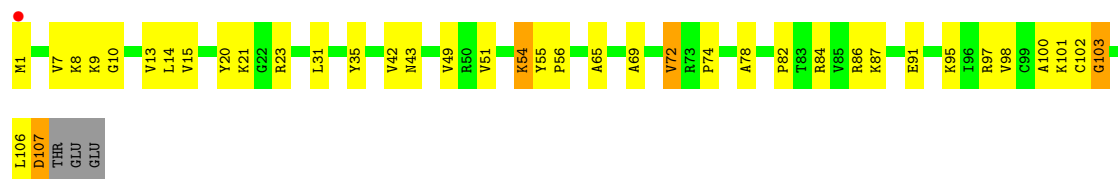


- Molecule 19: 50S ribosomal protein L23

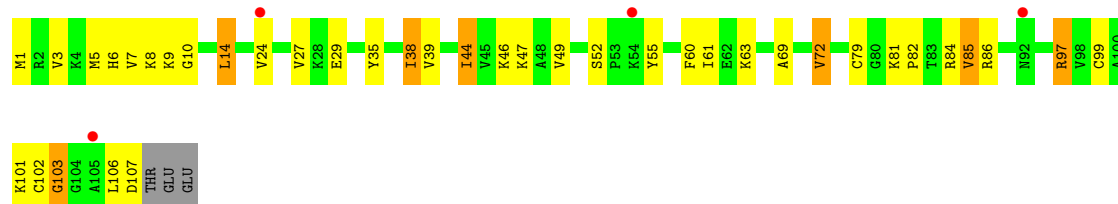




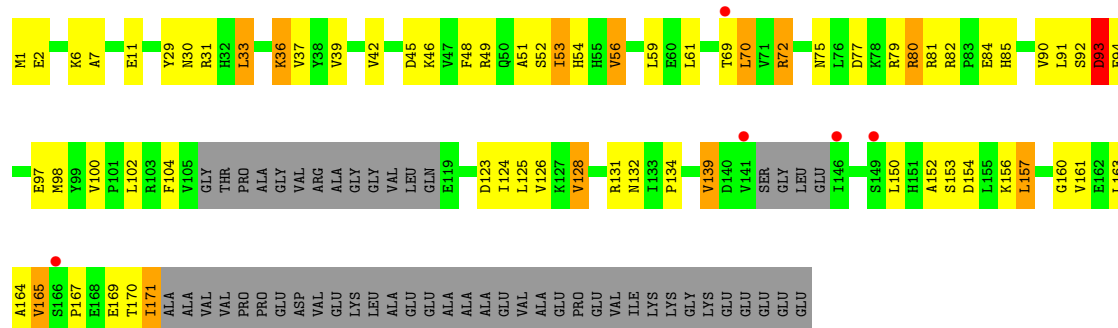
• Molecule 20: 50S ribosomal protein L24



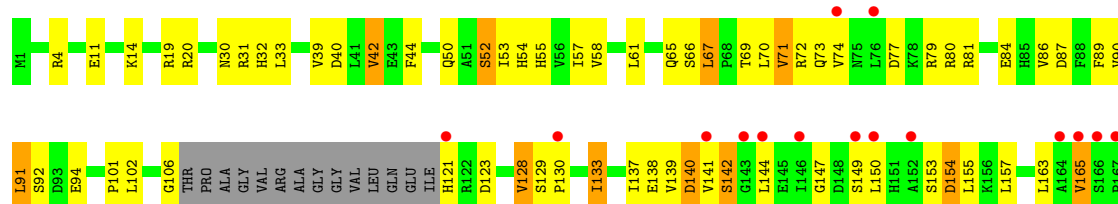
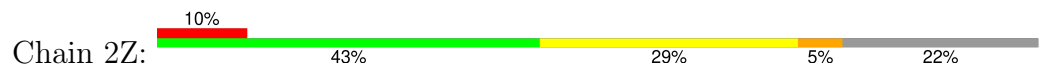
• Molecule 20: 50S ribosomal protein L24

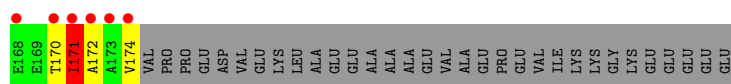


• Molecule 21: 50S ribosomal protein L25

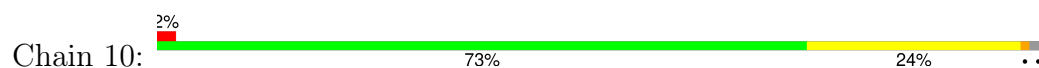


• Molecule 21: 50S ribosomal protein L25

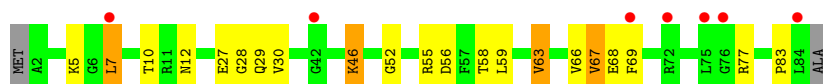
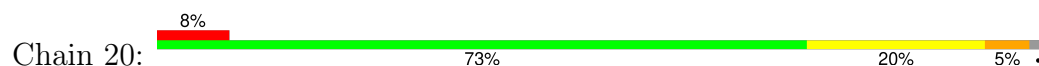




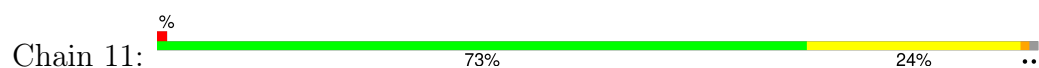
- Molecule 22: 50S ribosomal protein L27



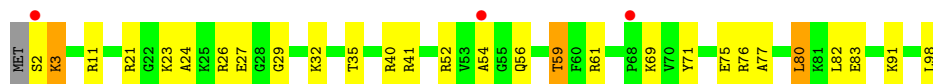
- Molecule 22: 50S ribosomal protein L27



- Molecule 23: 50S ribosomal protein L28



- Molecule 23: 50S ribosomal protein L28



- Molecule 24: 50S ribosomal protein L29

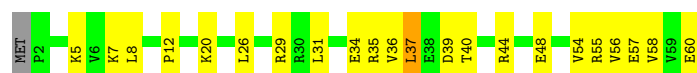


- Molecule 24: 50S ribosomal protein L29



- Molecule 25: 50S ribosomal protein L30

Chain 13:  62% 35% ..



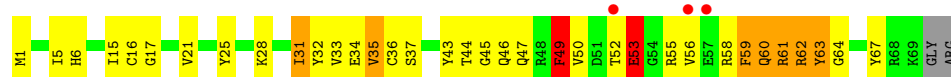
- Molecule 25: 50S ribosomal protein L30

Chain 23:  43% 47% 8% .



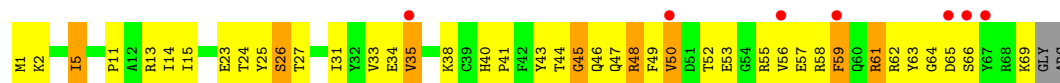
- Molecule 26: 50S ribosomal protein L31

Chain 14:  4% 48% 37% 10% ..




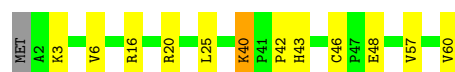
- Molecule 26: 50S ribosomal protein L31

Chain 24:  10% 39% 46% 11% .



- Molecule 27: 50S ribosomal protein L32

Chain 15:  78% 18% ..



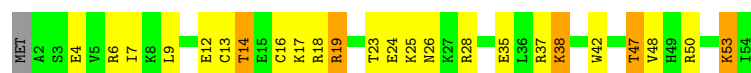
- Molecule 27: 50S ribosomal protein L32

Chain 25:  2% 72% 25% ..



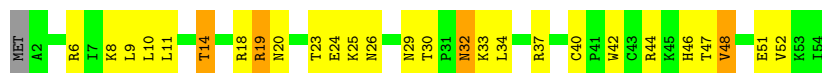
- Molecule 28: 50S ribosomal protein L33

Chain 16:  54% 35% 9% .



- Molecule 28: 50S ribosomal protein L33

Chain 26:  48% 43% 7% .



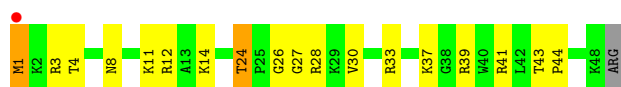
- Molecule 29: 50S ribosomal protein L34

Chain 17:  71% 24% ..




- Molecule 29: 50S ribosomal protein L34

Chain 27:  2% 61% 33% . .



- Molecule 30: 50S ribosomal protein L35

Chain 18:  2% 75% 22% . .




- Molecule 30: 50S ribosomal protein L35

Chain 28:  2% 60% 35% . .



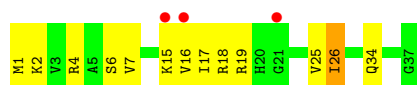
- Molecule 31: 50S ribosomal protein L36

Chain 19:  81% 19%



- Molecule 31: 50S ribosomal protein L36

Chain 29:  8% 65% 32% .



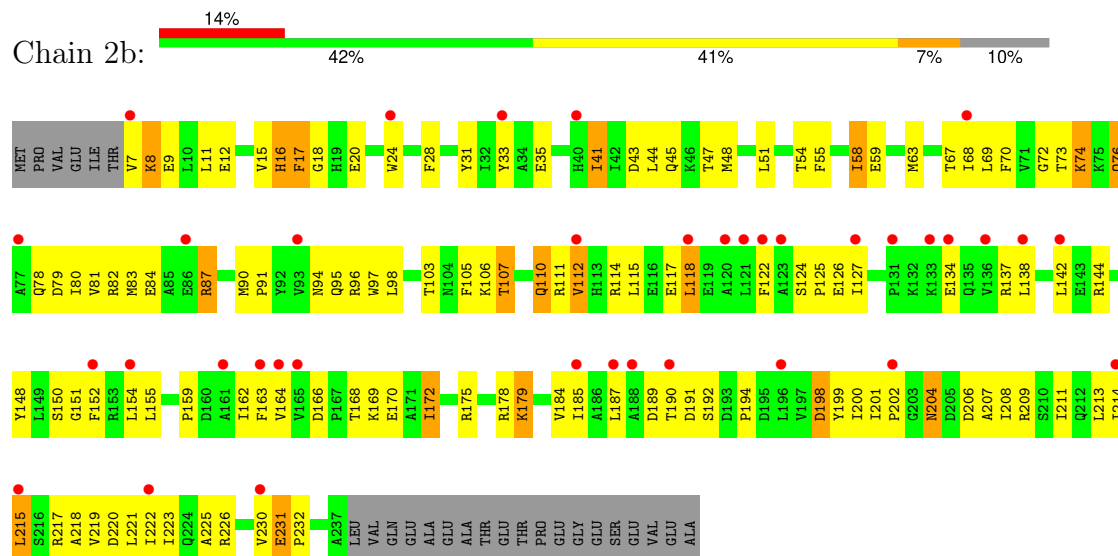
- Molecule 32: 16S Ribosomal RNA



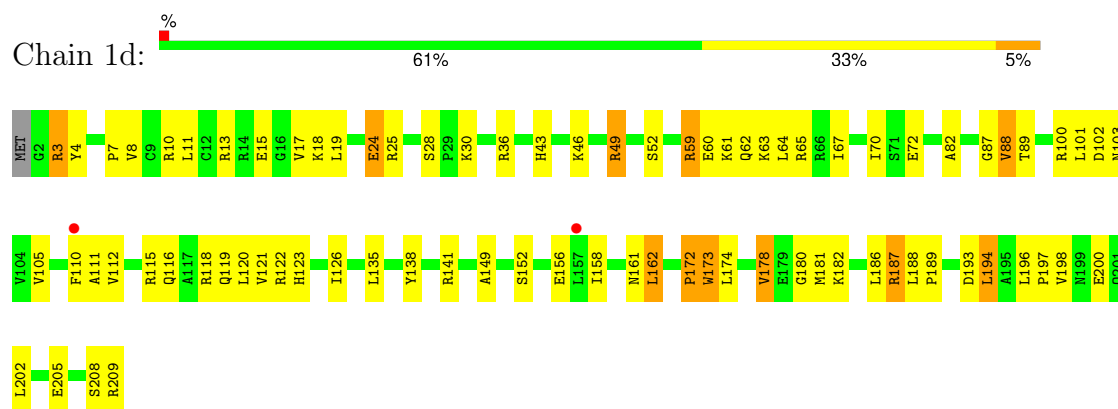


GLU
VAL
GLU
ALA

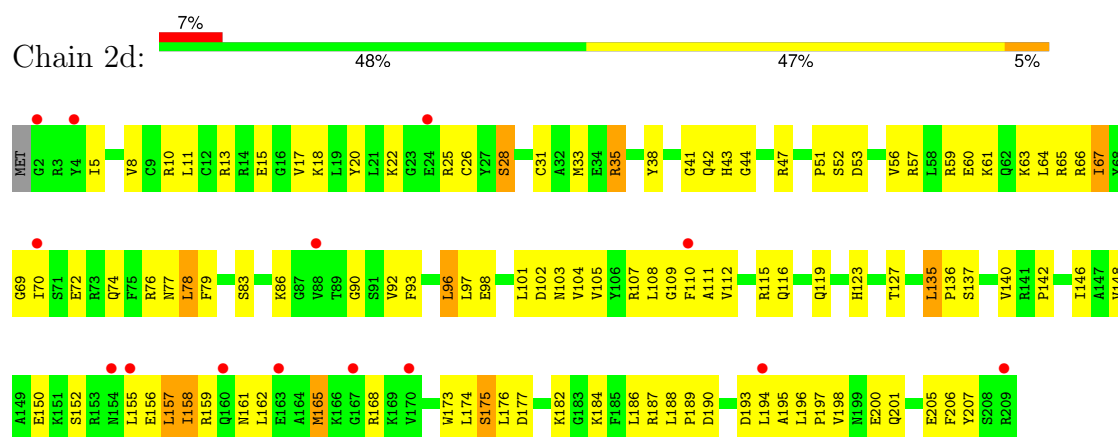
• Molecule 33: 30S ribosomal protein S2



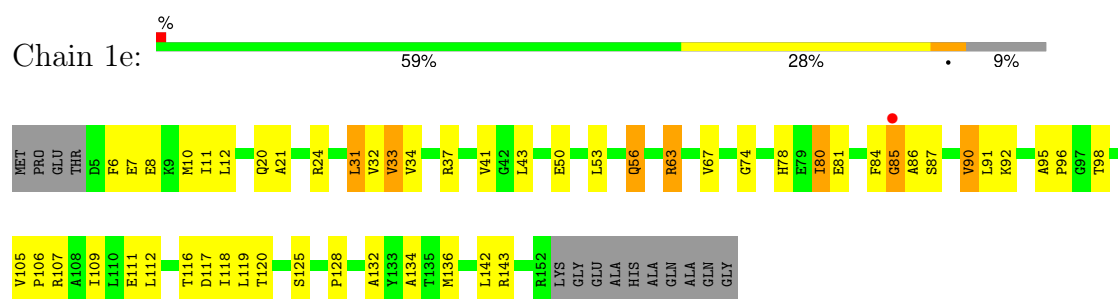
- Molecule 35: 30S ribosomal protein S4



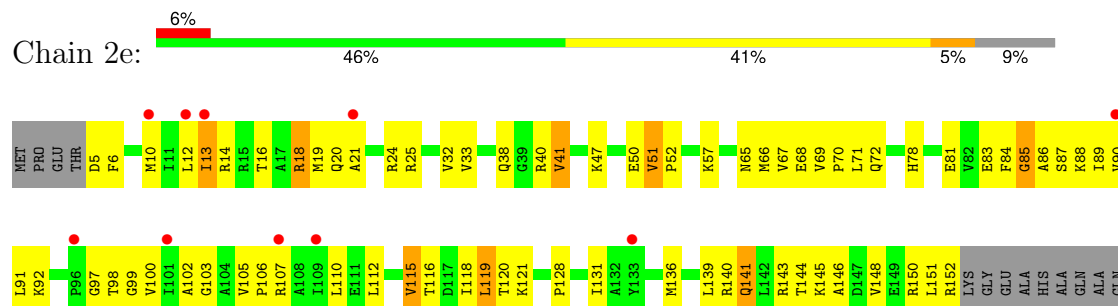
- Molecule 35: 30S ribosomal protein S4



- Molecule 36: 30S ribosomal protein S5

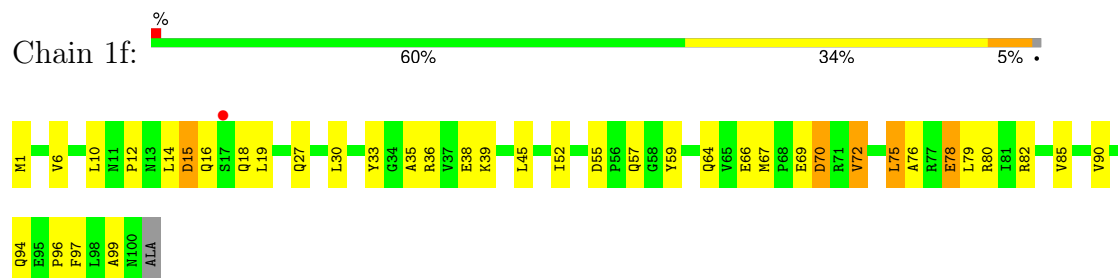


- Molecule 36: 30S ribosomal protein S5

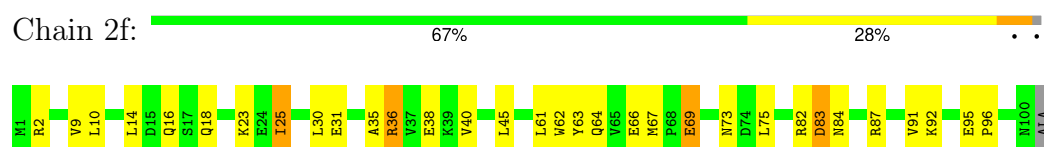


GLY

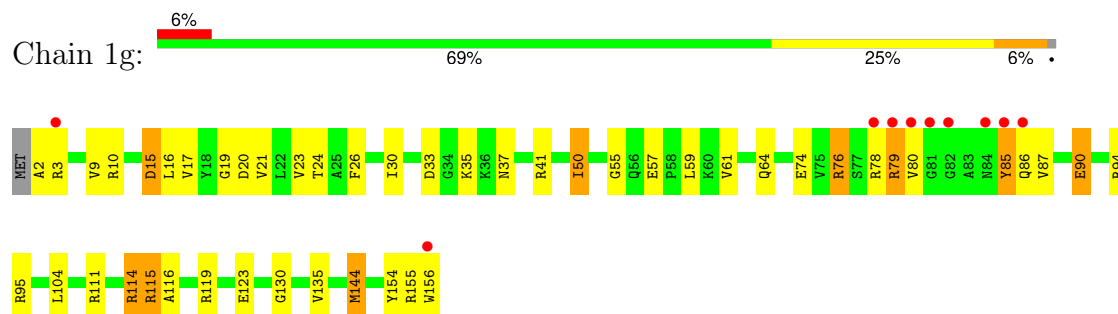
- Molecule 37: 30S ribosomal protein S6



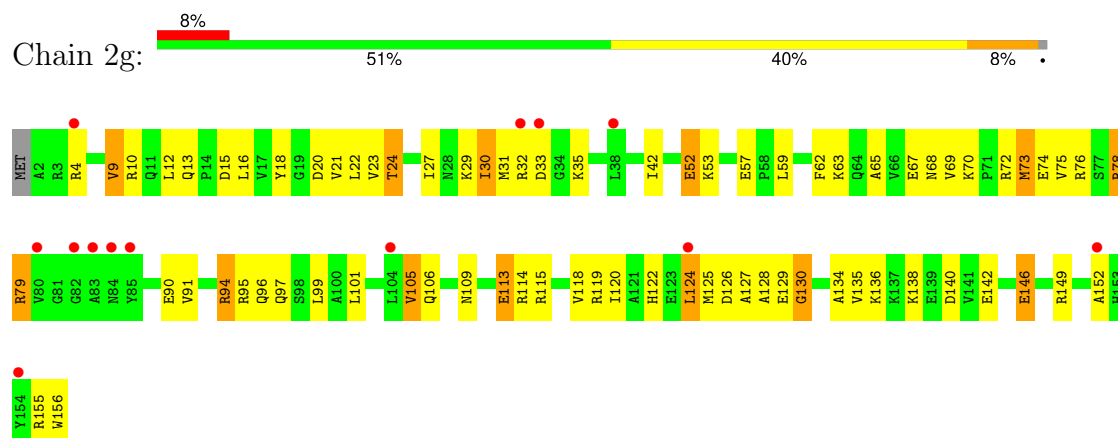
- Molecule 37: 30S ribosomal protein S6



- Molecule 38: 30S ribosomal protein S7

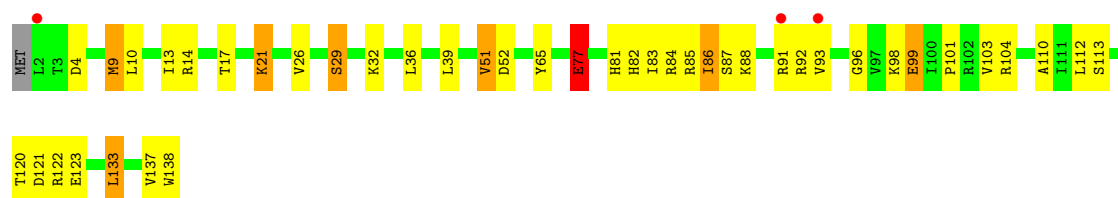


- Molecule 38: 30S ribosomal protein S7

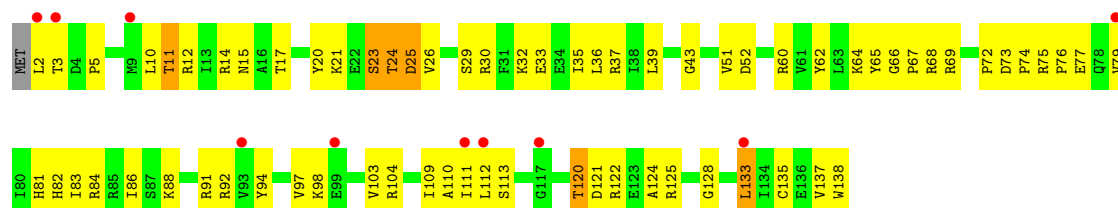


- Molecule 39: 30S ribosomal protein S8

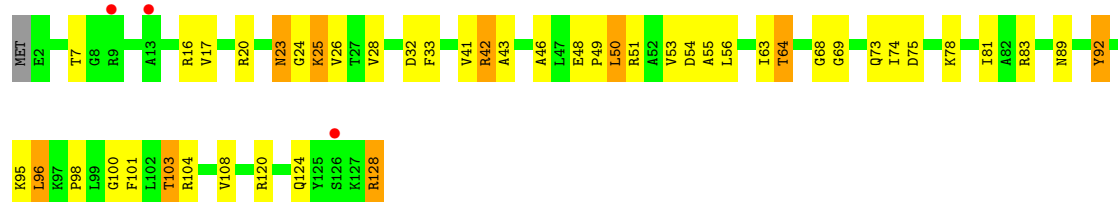




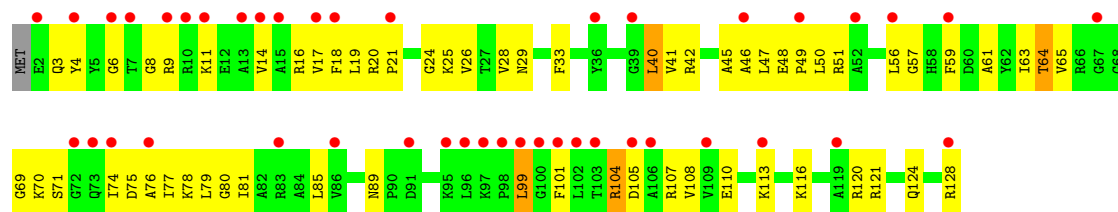
• Molecule 39: 30S ribosomal protein S8



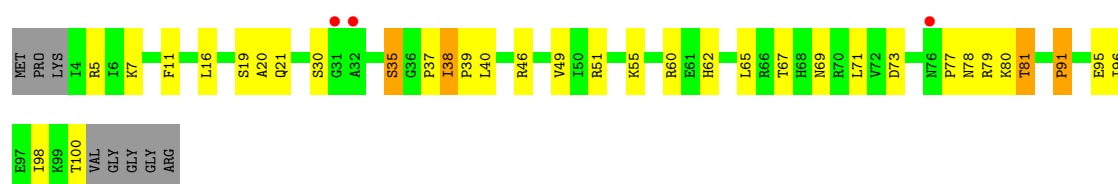
• Molecule 40: 30S ribosomal protein S9



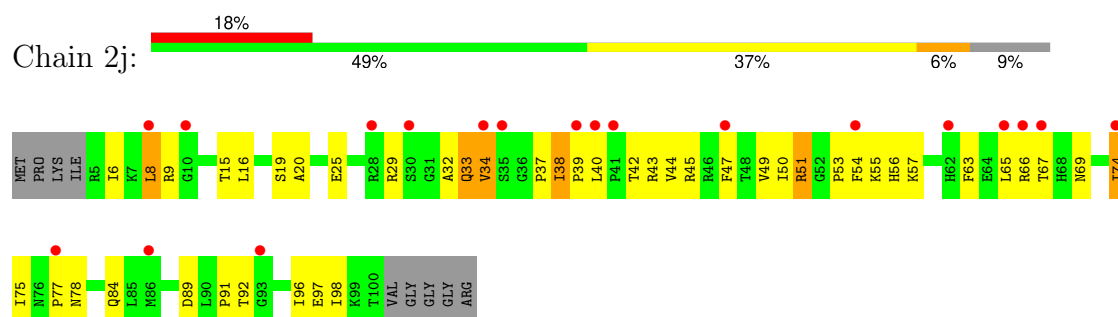
• Molecule 40: 30S ribosomal protein S9



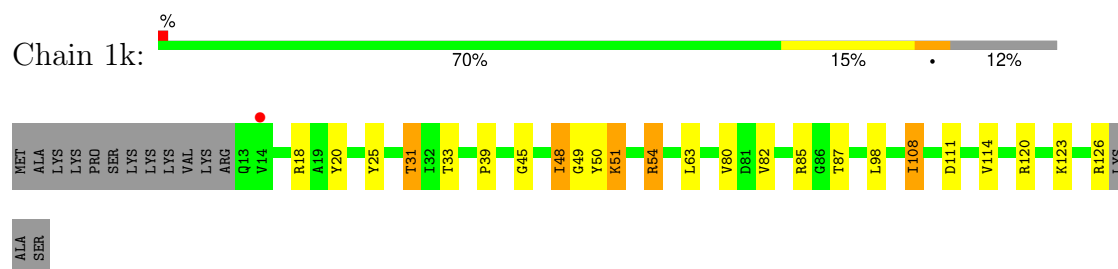
• Molecule 41: 30S ribosomal protein S10



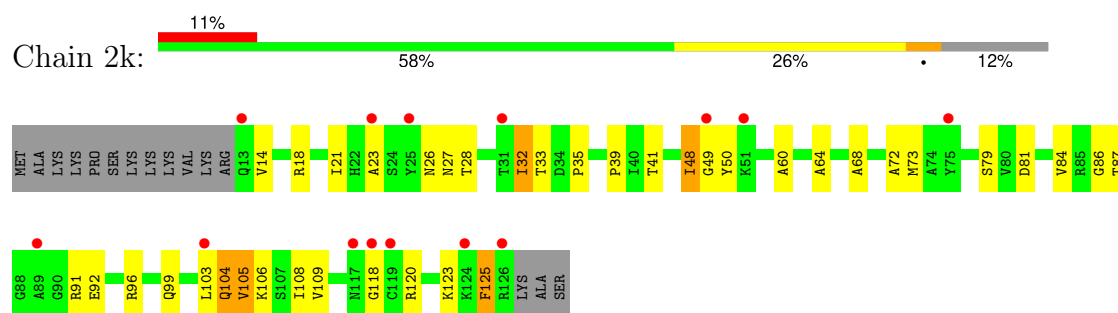
- Molecule 41: 30S ribosomal protein S10



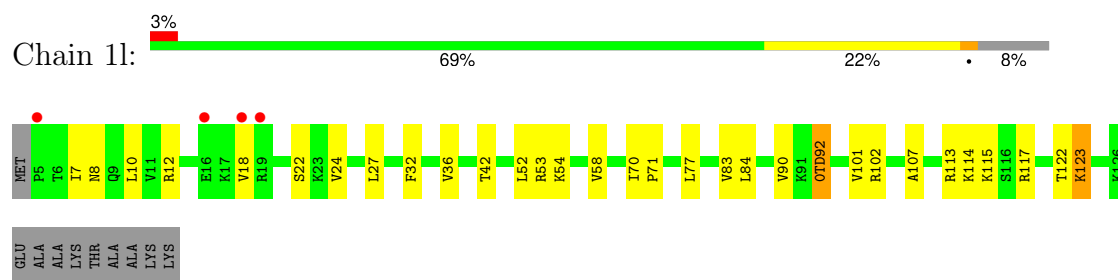
- Molecule 42: 30S ribosomal protein S11



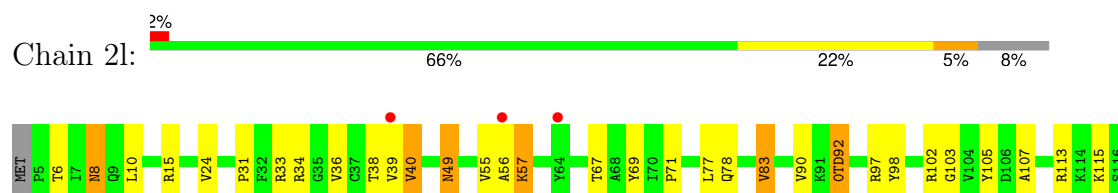
- Molecule 42: 30S ribosomal protein S11

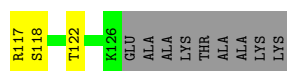


- Molecule 43: 30S ribosomal protein S12



- Molecule 43: 30S ribosomal protein S12





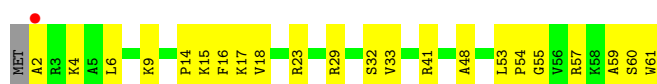
- Molecule 44: 30S ribosomal protein S13



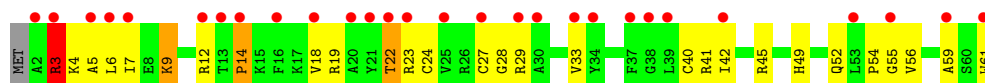
- Molecule 44: 30S ribosomal protein S13



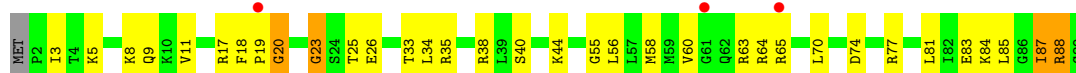
- Molecule 45: 30S ribosomal protein S14 type Z



- Molecule 45: 30S ribosomal protein S14 type Z



- Molecule 46: 30S ribosomal protein S15



- Molecule 46: 30S ribosomal protein S15



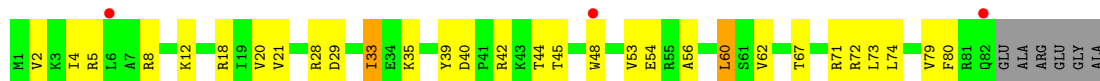


- Molecule 47: 30S ribosomal protein S16



ARG
GLU
GLY
ALA

- Molecule 47: 30S ribosomal protein S16

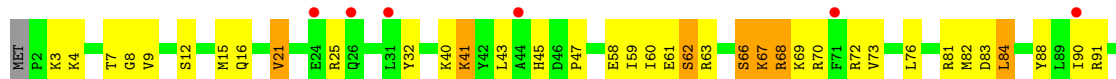


- Molecule 48: 30S ribosomal protein S17



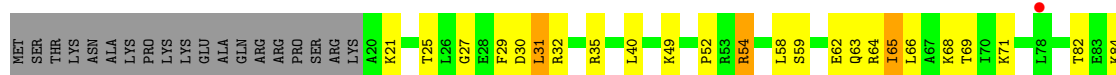
I90
R91
R92
Q93
N94
Y95
E96
S97
S99
K100
ARG
GLY
GLY
LYS
ALA

- Molecule 48: 30S ribosomal protein S17



S97
L98
S99
K100
ARG
GLY
GLY
LYS
ALA

- Molecule 49: 30S ribosomal protein S18





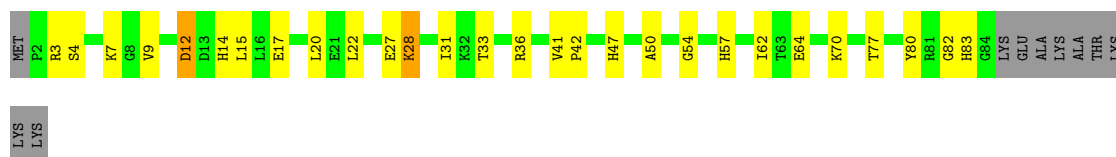
- Molecule 49: 30S ribosomal protein S18

Chain 2r: 48% 23% 7% 23%



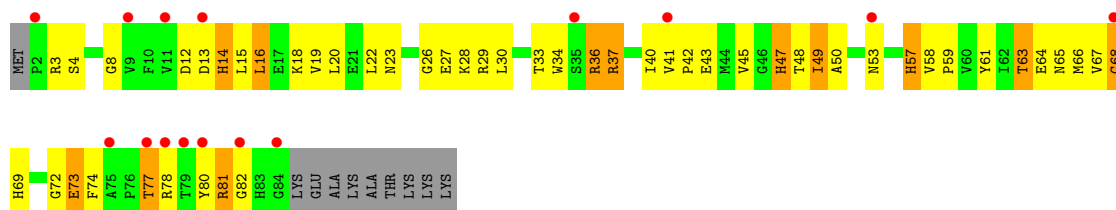
- Molecule 50: 30S ribosomal protein S19

Chain 1s: 59% 28% 11%



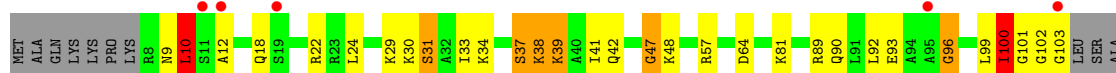
- Molecule 50: 30S ribosomal protein S19

Chain 2s: 16% 34% 42% 13% 11%



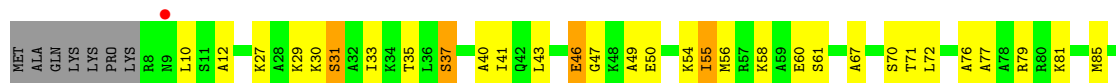
- Molecule 51: 30S ribosomal protein S20

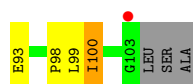
Chain 1t: 5% 61% 22% 6% 9%



- Molecule 51: 30S ribosomal protein S20

Chain 2t: 2% 58% 28% 5% 9%





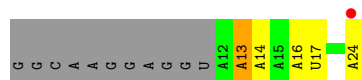
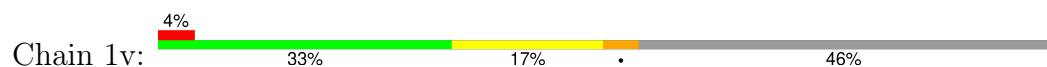
- Molecule 52: 30S ribosomal protein Thx



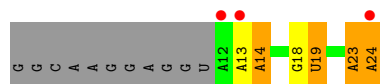
- Molecule 52: 30S ribosomal protein Thx



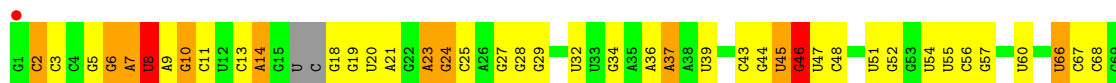
- Molecule 53: MET-PHE-mRNA



- Molecule 53: MET-PHE-mRNA



- Molecule 54: A-site and E-site Deacylated tRNAphe

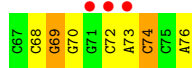


- Molecule 54: A-site and E-site Deacylated tRNAphe

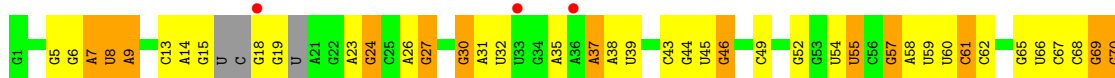




- Molecule 54: A-site and E-site Deacylated tRNA^{phe}



- Molecule 54: A-site and E-site Deacylated tRNA^{phe}



- Molecule 55: P-site Peptidyl-tRNA fMRC-tRNA^{cys} RNA-part



- Molecule 55: P-site Peptidyl-tRNA fMRC-tRNA^{cys} RNA-part



- Molecule 56: P-site Peptidyl-tRNA fMRC-tRNA^{cys} Peptide-part



- Molecule 56: P-site Peptidyl-tRNA fMRC-tRNA^{cys} Peptide-part



4 Data and refinement statistics

| Property | Value | Source |
|-------------------------------------------------------------------------|-------------------------------------------------------------|------------------|
| Space group | P 21 21 21 | Depositor |
| Cell constants a, b, c, α , β , γ | 208.76Å 447.07Å 617.30Å 90.00° 90.00° 90.00° | Depositor |
| Resolution (Å) | 122.56 – 2.75 122.56 – 2.75 | Depositor EDS |
| % Data completeness (in resolution range) | 99.8 (122.56-2.75) 99.8 (122.56-2.75) | Depositor EDS |
| R_{merge} | 0.22 | Depositor |
| R_{sym} | (Not available) | Depositor |
| $\langle I/\sigma(I) \rangle$ ¹ | 1.21 (at 2.73Å) | Xtriage |
| Refinement program | PHENIX 1.17.1 | Depositor |
| R, R_{free} | 0.228 , 0.282 0.230 , 0.281 | Depositor DCC |
| R_{free} test set | 74137 reflections (5.02%) | wwPDB-VP |
| Wilson B-factor (Å ²) | 58.6 | Xtriage |
| Anisotropy | 0.153 | Xtriage |
| Bulk solvent k_{sol} (e/Å ³), B_{sol} (Å ²) | 0.28 , 49.4 | EDS |
| L-test for twinning ² | $\langle L \rangle = 0.36$, $\langle L^2 \rangle = 0.19$ | Xtriage |
| Estimated twinning fraction | No twinning to report. | Xtriage |
| F_o, F_c correlation | 0.92 | EDS |
| Total number of atoms | 299368 | wwPDB-VP |
| Average B, all atoms (Å ²) | 59.0 | wwPDB-VP |

Xtriage's analysis on translational NCS is as follows: *The largest off-origin peak in the Patterson function is 1.66% of the height of the origin peak. No significant pseudotranslation is detected.*

¹Intensities estimated from amplitudes.

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

5 Model quality ⓘ

5.1 Standard geometry ⓘ

Bond lengths and bond angles in the following residue types are not validated in this section: M2G, OMC, ZN, 4SU, 4OC, 2MA, MIA, K, G7M, PSU, MA6, MG, 0TD, UR3, 5MU, FME, OMU, SF4, OMG, 2MG, 8AN, ERY, 5MC

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 | 0.28 | 1/69011 (0.0%) | 0.45 | 1/107720 (0.0%) |
| 1 | 2A | 0.22 | 0/67295 | 0.40 | 1/105042 (0.0%) |
| 2 | 1B | 0.22 | 0/2882 | 0.39 | 0/4494 |
| 2 | 2B | 0.20 | 0/2879 | 0.37 | 0/4487 |
| 3 | 1D | 0.28 | 0/2186 | 0.50 | 0/2944 |
| 3 | 2D | 0.23 | 0/2186 | 0.51 | 3/2944 (0.1%) |
| 4 | 1E | 0.26 | 0/1592 | 0.51 | 0/2149 |
| 4 | 2E | 0.23 | 0/1592 | 0.46 | 0/2149 |
| 5 | 1F | 0.27 | 0/1618 | 0.50 | 0/2191 |
| 5 | 2F | 0.21 | 0/1614 | 0.46 | 2/2186 (0.1%) |
| 6 | 1G | 0.23 | 0/1448 | 0.46 | 0/1957 |
| 6 | 2G | 0.20 | 0/1453 | 0.46 | 0/1963 |
| 7 | 1H | 0.22 | 0/1356 | 0.41 | 0/1834 |
| 7 | 2H | 0.21 | 0/1356 | 0.40 | 0/1834 |
| 8 | 1I | 0.20 | 0/1112 | 0.46 | 0/1514 |
| 8 | 2I | 0.22 | 0/1079 | 0.45 | 0/1475 |
| 9 | 1N | 0.24 | 0/1144 | 0.45 | 0/1543 |
| 9 | 2N | 0.19 | 0/1144 | 0.43 | 0/1543 |
| 10 | 1O | 0.27 | 0/943 | 0.46 | 0/1269 |
| 10 | 2O | 0.21 | 0/943 | 0.45 | 0/1269 |
| 11 | 1P | 0.28 | 0/1152 | 0.49 | 0/1533 |
| 11 | 2P | 0.22 | 0/1152 | 0.51 | 0/1533 |
| 12 | 1Q | 0.27 | 0/1143 | 0.48 | 0/1527 |
| 12 | 2Q | 0.21 | 0/1143 | 0.44 | 0/1527 |
| 13 | 1R | 0.25 | 0/982 | 0.46 | 0/1312 |
| 13 | 2R | 0.22 | 0/982 | 0.45 | 0/1312 |
| 14 | 1S | 0.22 | 0/883 | 0.47 | 0/1176 |
| 14 | 2S | 0.23 | 0/880 | 0.47 | 0/1172 |
| 15 | 1T | 0.24 | 0/1105 | 0.48 | 0/1477 |
| 15 | 2T | 0.23 | 0/1097 | 0.45 | 0/1468 |
| 16 | 1U | 0.26 | 0/977 | 0.42 | 0/1301 |

| Mol | Chain | Bond lengths | | Bond angles | |
|-----|-------|--------------|----------------|-------------|----------------|
| | | RMSZ | # Z >5 | RMSZ | # Z >5 |
| 16 | 2U | 0.20 | 0/977 | 0.42 | 0/1301 |
| 17 | 1V | 0.23 | 0/782 | 0.43 | 0/1049 |
| 17 | 2V | 0.19 | 0/782 | 0.41 | 0/1049 |
| 18 | 1W | 0.26 | 0/897 | 0.44 | 0/1205 |
| 18 | 2W | 0.22 | 0/897 | 0.41 | 0/1205 |
| 19 | 1X | 0.28 | 0/764 | 0.51 | 0/1025 |
| 19 | 2X | 0.22 | 0/764 | 0.48 | 0/1025 |
| 20 | 1Y | 0.23 | 0/819 | 0.48 | 0/1095 |
| 20 | 2Y | 0.20 | 0/819 | 0.45 | 0/1095 |
| 21 | 1Z | 0.22 | 0/1267 | 0.48 | 0/1717 |
| 21 | 2Z | 0.22 | 0/1299 | 0.43 | 0/1763 |
| 22 | 10 | 0.24 | 0/662 | 0.51 | 0/881 |
| 22 | 20 | 0.22 | 0/662 | 0.45 | 0/881 |
| 23 | 11 | 0.26 | 0/762 | 0.42 | 0/1014 |
| 23 | 21 | 0.22 | 0/762 | 0.43 | 0/1014 |
| 24 | 12 | 0.25 | 0/590 | 0.46 | 0/781 |
| 24 | 22 | 0.22 | 0/590 | 0.40 | 0/781 |
| 25 | 13 | 0.26 | 0/474 | 0.45 | 0/635 |
| 25 | 23 | 0.19 | 0/469 | 0.42 | 0/630 |
| 26 | 14 | 0.24 | 0/565 | 0.55 | 0/761 |
| 26 | 24 | 0.26 | 0/545 | 0.51 | 0/737 |
| 27 | 15 | 0.24 | 0/469 | 0.47 | 0/635 |
| 27 | 25 | 0.26 | 0/469 | 0.50 | 0/635 |
| 28 | 16 | 0.25 | 0/460 | 0.50 | 0/613 |
| 28 | 26 | 0.21 | 0/456 | 0.41 | 0/608 |
| 29 | 17 | 0.31 | 0/426 | 0.58 | 0/561 |
| 29 | 27 | 0.25 | 0/426 | 0.53 | 0/561 |
| 30 | 18 | 0.28 | 0/525 | 0.50 | 0/691 |
| 30 | 28 | 0.22 | 0/525 | 0.44 | 0/691 |
| 31 | 19 | 0.28 | 0/310 | 0.58 | 0/407 |
| 31 | 29 | 0.20 | 0/310 | 0.43 | 0/407 |
| 32 | 1a | 0.21 | 0/35795 | 0.39 | 2/55864 (0.0%) |
| 32 | 2a | 0.20 | 1/35886 (0.0%) | 0.38 | 3/56005 (0.0%) |
| 33 | 1b | 0.22 | 0/1881 | 0.53 | 0/2542 |
| 33 | 2b | 0.24 | 0/1860 | 0.50 | 0/2518 |
| 34 | 1c | 0.20 | 0/1572 | 0.41 | 0/2126 |
| 34 | 2c | 0.25 | 0/1566 | 0.49 | 0/2119 |
| 35 | 1d | 0.21 | 0/1685 | 0.45 | 0/2262 |
| 35 | 2d | 0.21 | 0/1704 | 0.47 | 0/2284 |
| 36 | 1e | 0.21 | 0/1145 | 0.46 | 0/1543 |
| 36 | 2e | 0.21 | 0/1149 | 0.50 | 0/1548 |
| 37 | 1f | 0.20 | 0/823 | 0.39 | 0/1115 |
| 37 | 2f | 0.21 | 0/829 | 0.41 | 0/1123 |

| Mol | Chain | Bond lengths | | Bond angles | |
|-----|-------|--------------|-----------------|-------------|------------------|
| | | RMSZ | # Z >5 | RMSZ | # Z >5 |
| 38 | 1g | 0.21 | 0/1250 | 0.43 | 0/1679 |
| 38 | 2g | 0.20 | 0/1254 | 0.46 | 0/1683 |
| 39 | 1h | 0.20 | 0/1108 | 0.41 | 0/1494 |
| 39 | 2h | 0.19 | 0/1108 | 0.41 | 0/1494 |
| 40 | 1i | 0.22 | 0/1002 | 0.47 | 0/1346 |
| 40 | 2i | 0.21 | 0/997 | 0.49 | 0/1343 |
| 41 | 1j | 0.20 | 0/722 | 0.44 | 0/982 |
| 41 | 2j | 0.21 | 0/727 | 0.44 | 0/988 |
| 42 | 1k | 0.20 | 0/844 | 0.40 | 0/1145 |
| 42 | 2k | 0.18 | 0/848 | 0.40 | 0/1149 |
| 43 | 1l | 0.23 | 0/937 | 0.43 | 0/1260 |
| 43 | 2l | 0.21 | 0/937 | 0.44 | 0/1260 |
| 44 | 1m | 0.20 | 0/969 | 0.45 | 0/1302 |
| 44 | 2m | 0.21 | 0/961 | 0.46 | 0/1291 |
| 45 | 1n | 0.22 | 0/501 | 0.52 | 0/664 |
| 45 | 2n | 0.21 | 0/501 | 0.45 | 0/664 |
| 46 | 1o | 0.20 | 0/739 | 0.42 | 0/985 |
| 46 | 2o | 0.19 | 0/739 | 0.41 | 0/985 |
| 47 | 1p | 0.22 | 0/697 | 0.49 | 0/939 |
| 47 | 2p | 0.19 | 0/693 | 0.42 | 0/935 |
| 48 | 1q | 0.19 | 0/836 | 0.43 | 0/1117 |
| 48 | 2q | 0.20 | 0/836 | 0.48 | 0/1117 |
| 49 | 1r | 0.22 | 0/560 | 0.50 | 0/746 |
| 49 | 2r | 0.21 | 0/560 | 0.45 | 0/746 |
| 50 | 1s | 0.21 | 0/667 | 0.49 | 0/900 |
| 50 | 2s | 0.28 | 0/661 | 0.58 | 0/893 |
| 51 | 1t | 0.23 | 0/730 | 0.53 | 0/965 |
| 51 | 2t | 0.22 | 0/729 | 0.50 | 0/965 |
| 52 | 1u | 0.19 | 0/203 | 0.44 | 0/266 |
| 52 | 2u | 0.20 | 0/203 | 0.52 | 0/266 |
| 53 | 1v | 0.23 | 0/310 | 0.35 | 0/480 |
| 53 | 2v | 0.21 | 0/310 | 0.40 | 0/480 |
| 54 | 1w | 0.30 | 2/1606 (0.1%) | 0.42 | 0/2497 |
| 54 | 1y | 0.28 | 1/1606 (0.1%) | 0.43 | 0/2497 |
| 54 | 2w | 0.30 | 1/1556 (0.1%) | 0.43 | 0/2418 |
| 54 | 2y | 0.28 | 1/1583 (0.1%) | 0.39 | 0/2459 |
| 55 | 1x | 0.27 | 0/1723 | 0.42 | 0/2684 |
| 55 | 2x | 0.24 | 0/1723 | 0.40 | 0/2684 |
| 56 | 1z | 0.34 | 0/16 | 0.38 | 0/19 |
| 56 | 2z | 0.34 | 0/16 | 0.64 | 0/19 |
| All | All | 0.23 | 7/316716 (0.0%) | 0.43 | 12/474153 (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 |
|-----|-------|---------------------|---------------------|
| 33 | 1b | 0 | 2 |

The worst 5 of 7 bond length outliers are listed below:

| Mol | Chain | Res | Type | Atoms | Z | Observed(Å) | Ideal(Å) |
|-----|-------|------|------|-------|------|-------------|----------|
| 54 | 2w | 46 | G7M | O3'-P | 5.71 | 1.61 | 1.56 |
| 54 | 1w | 8 | 4SU | O3'-P | 5.36 | 1.61 | 1.56 |
| 54 | 1y | 8 | 4SU | O3'-P | 5.13 | 1.61 | 1.56 |
| 1 | 1A | 2552 | OMU | O3'-P | 5.12 | 1.61 | 1.56 |
| 32 | 2a | 527 | G7M | O3'-P | 5.08 | 1.61 | 1.56 |

The worst 5 of 12 bond angle outliers are listed below:

| Mol | Chain | Res | Type | Atoms | Z | Observed(°) | Ideal(°) |
|-----|-------|------|------|-------------|------|-------------|----------|
| 1 | 1A | 1992 | G | C2'-C3'-O3' | 5.93 | 118.40 | 109.50 |
| 5 | 2F | 21 | ALA | CA-C-N | 5.85 | 132.22 | 121.70 |
| 5 | 2F | 21 | ALA | C-N-CA | 5.85 | 132.22 | 121.70 |
| 1 | 2A | 1992 | G | C2'-C3'-O3' | 5.64 | 117.97 | 109.50 |
| 32 | 1a | 266 | G | C2'-C3'-O3' | 5.61 | 117.91 | 109.50 |

There are no chirality outliers.

All (2) planarity outliers are listed below:

| Mol | Chain | Res | Type | Group |
|-----|-------|-----|------|---------|
| 33 | 1b | 125 | PRO | Peptide |
| 33 | 1b | 20 | GLU | 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 | 61852 | 0 | 31192 | 706 | 0 |
| 1 | 2A | 60322 | 0 | 30425 | 775 | 0 |
| 2 | 1B | 2577 | 0 | 1305 | 31 | 0 |

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| Mol | Chain | Non-H | H(model) | H(added) | Clashes | Symm-Clashes |
|-----|-------|-------|----------|----------|---------|--------------|
| 2 | 2B | 2575 | 0 | 1303 | 49 | 0 |
| 3 | 1D | 2136 | 0 | 2218 | 54 | 0 |
| 3 | 2D | 2136 | 0 | 2218 | 50 | 0 |
| 4 | 1E | 1559 | 0 | 1618 | 29 | 0 |
| 4 | 2E | 1559 | 0 | 1618 | 46 | 0 |
| 5 | 1F | 1583 | 0 | 1625 | 46 | 0 |
| 5 | 2F | 1579 | 0 | 1619 | 52 | 0 |
| 6 | 1G | 1423 | 0 | 1436 | 47 | 0 |
| 6 | 2G | 1428 | 0 | 1438 | 60 | 0 |
| 7 | 1H | 1330 | 0 | 1407 | 29 | 0 |
| 7 | 2H | 1330 | 0 | 1407 | 35 | 0 |
| 8 | 1I | 1097 | 0 | 1140 | 25 | 0 |
| 8 | 2I | 1064 | 0 | 1082 | 35 | 0 |
| 9 | 1N | 1117 | 0 | 1184 | 19 | 0 |
| 9 | 2N | 1117 | 0 | 1184 | 30 | 0 |
| 10 | 1O | 933 | 0 | 996 | 19 | 0 |
| 10 | 2O | 933 | 0 | 996 | 21 | 0 |
| 11 | 1P | 1135 | 0 | 1212 | 38 | 0 |
| 11 | 2P | 1135 | 0 | 1212 | 38 | 0 |
| 12 | 1Q | 1122 | 0 | 1179 | 44 | 0 |
| 12 | 2Q | 1122 | 0 | 1179 | 47 | 0 |
| 13 | 1R | 968 | 0 | 1033 | 18 | 0 |
| 13 | 2R | 968 | 0 | 1033 | 24 | 0 |
| 14 | 1S | 873 | 0 | 927 | 20 | 0 |
| 14 | 2S | 870 | 0 | 923 | 41 | 0 |
| 15 | 1T | 1091 | 0 | 1151 | 28 | 0 |
| 15 | 2T | 1083 | 0 | 1136 | 26 | 0 |
| 16 | 1U | 959 | 0 | 1018 | 16 | 0 |
| 16 | 2U | 959 | 0 | 1019 | 24 | 0 |
| 17 | 1V | 771 | 0 | 828 | 13 | 0 |
| 17 | 2V | 771 | 0 | 830 | 17 | 0 |
| 18 | 1W | 886 | 0 | 940 | 19 | 0 |
| 18 | 2W | 886 | 0 | 940 | 14 | 0 |
| 19 | 1X | 750 | 0 | 814 | 11 | 0 |
| 19 | 2X | 750 | 0 | 813 | 14 | 0 |
| 20 | 1Y | 806 | 0 | 881 | 20 | 0 |
| 20 | 2Y | 806 | 0 | 881 | 20 | 0 |
| 21 | 1Z | 1240 | 0 | 1240 | 39 | 0 |
| 21 | 2Z | 1271 | 0 | 1273 | 44 | 0 |
| 22 | 10 | 653 | 0 | 674 | 19 | 0 |
| 22 | 20 | 653 | 0 | 674 | 12 | 0 |
| 23 | 11 | 755 | 0 | 826 | 9 | 0 |

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| Mol | Chain | Non-H | H(model) | H(added) | Clashes | Symm-Clashes |
|-----|-------|-------|----------|----------|---------|--------------|
| 23 | 21 | 755 | 0 | 826 | 21 | 0 |
| 24 | 12 | 588 | 0 | 643 | 16 | 0 |
| 24 | 22 | 588 | 0 | 643 | 14 | 0 |
| 25 | 13 | 469 | 0 | 518 | 10 | 0 |
| 25 | 23 | 464 | 0 | 514 | 20 | 0 |
| 26 | 14 | 552 | 0 | 533 | 27 | 0 |
| 26 | 24 | 532 | 0 | 503 | 30 | 0 |
| 27 | 15 | 455 | 0 | 465 | 9 | 0 |
| 27 | 25 | 455 | 0 | 465 | 9 | 0 |
| 28 | 16 | 453 | 0 | 473 | 15 | 0 |
| 28 | 26 | 449 | 0 | 469 | 16 | 0 |
| 29 | 17 | 418 | 0 | 467 | 8 | 0 |
| 29 | 27 | 418 | 0 | 467 | 13 | 0 |
| 30 | 18 | 517 | 0 | 582 | 14 | 0 |
| 30 | 28 | 517 | 0 | 582 | 20 | 0 |
| 31 | 19 | 307 | 0 | 335 | 4 | 0 |
| 31 | 29 | 307 | 0 | 335 | 8 | 0 |
| 32 | 1a | 32246 | 0 | 16293 | 438 | 0 |
| 32 | 2a | 32327 | 0 | 16337 | 585 | 0 |
| 33 | 1b | 1846 | 0 | 1867 | 61 | 0 |
| 33 | 2b | 1825 | 0 | 1828 | 81 | 0 |
| 34 | 1c | 1548 | 0 | 1535 | 40 | 0 |
| 34 | 2c | 1542 | 0 | 1517 | 68 | 0 |
| 35 | 1d | 1655 | 0 | 1672 | 64 | 0 |
| 35 | 2d | 1674 | 0 | 1714 | 76 | 0 |
| 36 | 1e | 1129 | 0 | 1185 | 38 | 0 |
| 36 | 2e | 1133 | 0 | 1191 | 50 | 0 |
| 37 | 1f | 810 | 0 | 804 | 23 | 0 |
| 37 | 2f | 816 | 0 | 808 | 19 | 0 |
| 38 | 1g | 1231 | 0 | 1238 | 36 | 0 |
| 38 | 2g | 1235 | 0 | 1249 | 48 | 0 |
| 39 | 1h | 1088 | 0 | 1126 | 26 | 0 |
| 39 | 2h | 1088 | 0 | 1126 | 45 | 0 |
| 40 | 1i | 983 | 0 | 986 | 36 | 0 |
| 40 | 2i | 978 | 0 | 966 | 45 | 0 |
| 41 | 1j | 709 | 0 | 650 | 23 | 0 |
| 41 | 2j | 714 | 0 | 672 | 30 | 0 |
| 42 | 1k | 829 | 0 | 825 | 18 | 0 |
| 42 | 2k | 833 | 0 | 834 | 23 | 0 |
| 43 | 1l | 932 | 0 | 981 | 18 | 0 |
| 43 | 2l | 932 | 0 | 980 | 23 | 0 |
| 44 | 1m | 958 | 0 | 1002 | 26 | 0 |

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| Mol | Chain | Non-H | H(model) | H(added) | Clashes | Symm-Clashes |
|-----|-------|-------|----------|----------|---------|--------------|
| 44 | 2m | 950 | 0 | 988 | 48 | 0 |
| 45 | 1n | 492 | 0 | 529 | 17 | 0 |
| 45 | 2n | 492 | 0 | 529 | 27 | 0 |
| 46 | 1o | 728 | 0 | 760 | 18 | 0 |
| 46 | 2o | 728 | 0 | 760 | 18 | 0 |
| 47 | 1p | 681 | 0 | 697 | 19 | 0 |
| 47 | 2p | 677 | 0 | 686 | 19 | 0 |
| 48 | 1q | 823 | 0 | 891 | 31 | 0 |
| 48 | 2q | 823 | 0 | 891 | 24 | 0 |
| 49 | 1r | 555 | 0 | 618 | 19 | 0 |
| 49 | 2r | 555 | 0 | 618 | 17 | 0 |
| 50 | 1s | 652 | 0 | 662 | 22 | 0 |
| 50 | 2s | 646 | 0 | 644 | 51 | 0 |
| 51 | 1t | 728 | 0 | 798 | 23 | 0 |
| 51 | 2t | 727 | 0 | 796 | 24 | 0 |
| 52 | 1u | 199 | 0 | 208 | 2 | 0 |
| 52 | 2u | 199 | 0 | 208 | 12 | 0 |
| 53 | 1v | 277 | 0 | 140 | 3 | 0 |
| 53 | 2v | 277 | 0 | 139 | 8 | 0 |
| 54 | 1w | 1592 | 0 | 818 | 27 | 0 |
| 54 | 1y | 1585 | 0 | 803 | 29 | 0 |
| 54 | 2w | 1544 | 0 | 786 | 28 | 0 |
| 54 | 2y | 1565 | 0 | 793 | 23 | 0 |
| 55 | 1x | 1646 | 0 | 839 | 22 | 0 |
| 55 | 2x | 1646 | 0 | 838 | 24 | 0 |
| 56 | 1z | 27 | 0 | 28 | 1 | 0 |
| 56 | 2z | 27 | 0 | 28 | 1 | 0 |
| 57 | 10 | 10 | 0 | 0 | 0 | 0 |
| 57 | 11 | 3 | 0 | 0 | 0 | 0 |
| 57 | 12 | 2 | 0 | 0 | 0 | 0 |
| 57 | 13 | 5 | 0 | 0 | 0 | 0 |
| 57 | 14 | 1 | 0 | 0 | 0 | 0 |
| 57 | 15 | 9 | 0 | 0 | 0 | 0 |
| 57 | 16 | 2 | 0 | 0 | 0 | 0 |
| 57 | 17 | 6 | 0 | 0 | 0 | 0 |
| 57 | 18 | 7 | 0 | 0 | 0 | 0 |
| 57 | 1A | 1085 | 0 | 0 | 0 | 0 |
| 57 | 1B | 38 | 0 | 0 | 0 | 0 |
| 57 | 1D | 12 | 0 | 0 | 0 | 0 |
| 57 | 1E | 13 | 0 | 0 | 0 | 0 |
| 57 | 1F | 13 | 0 | 0 | 0 | 0 |
| 57 | 1G | 4 | 0 | 0 | 0 | 0 |

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| Mol | Chain | Non-H | H(model) | H(added) | Clashes | Symm-Clashes |
|-----|-------|-------|----------|----------|---------|--------------|
| 57 | 1H | 1 | 0 | 0 | 0 | 0 |
| 57 | 1I | 1 | 0 | 0 | 0 | 0 |
| 57 | 1N | 4 | 0 | 0 | 0 | 0 |
| 57 | 1O | 4 | 0 | 0 | 0 | 0 |
| 57 | 1P | 6 | 0 | 0 | 0 | 0 |
| 57 | 1Q | 7 | 0 | 0 | 0 | 0 |
| 57 | 1R | 6 | 0 | 0 | 0 | 0 |
| 57 | 1S | 2 | 0 | 0 | 0 | 0 |
| 57 | 1T | 3 | 0 | 0 | 0 | 0 |
| 57 | 1U | 11 | 0 | 0 | 0 | 0 |
| 57 | 1V | 7 | 0 | 0 | 0 | 0 |
| 57 | 1W | 7 | 0 | 0 | 0 | 0 |
| 57 | 1X | 7 | 0 | 0 | 0 | 0 |
| 57 | 1Y | 2 | 0 | 0 | 0 | 0 |
| 57 | 1Z | 2 | 0 | 0 | 0 | 0 |
| 57 | 1a | 211 | 0 | 0 | 0 | 0 |
| 57 | 1b | 1 | 0 | 0 | 0 | 0 |
| 57 | 1d | 1 | 0 | 0 | 0 | 0 |
| 57 | 1e | 1 | 0 | 0 | 0 | 0 |
| 57 | 1f | 1 | 0 | 0 | 0 | 0 |
| 57 | 1j | 1 | 0 | 0 | 0 | 0 |
| 57 | 1l | 2 | 0 | 0 | 0 | 0 |
| 57 | 1m | 1 | 0 | 0 | 0 | 0 |
| 57 | 1n | 3 | 0 | 0 | 0 | 0 |
| 57 | 1p | 1 | 0 | 0 | 0 | 0 |
| 57 | 1r | 1 | 0 | 0 | 0 | 0 |
| 57 | 1t | 1 | 0 | 0 | 0 | 0 |
| 57 | 1w | 6 | 0 | 0 | 0 | 0 |
| 57 | 1x | 12 | 0 | 0 | 0 | 0 |
| 57 | 20 | 3 | 0 | 0 | 0 | 0 |
| 57 | 21 | 1 | 0 | 0 | 0 | 0 |
| 57 | 23 | 2 | 0 | 0 | 0 | 0 |
| 57 | 25 | 1 | 0 | 0 | 0 | 0 |
| 57 | 26 | 1 | 0 | 0 | 0 | 0 |
| 57 | 27 | 4 | 0 | 0 | 0 | 0 |
| 57 | 28 | 3 | 0 | 0 | 0 | 0 |
| 57 | 2A | 855 | 0 | 0 | 0 | 0 |
| 57 | 2B | 20 | 0 | 0 | 0 | 0 |
| 57 | 2D | 6 | 0 | 0 | 0 | 0 |
| 57 | 2E | 9 | 0 | 0 | 0 | 0 |
| 57 | 2F | 6 | 0 | 0 | 0 | 0 |
| 57 | 2N | 1 | 0 | 0 | 0 | 0 |

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| Mol | Chain | Non-H | H(model) | H(added) | Clashes | Symm-Clashes |
|-----|-------|-------|----------|----------|---------|--------------|
| 57 | 2O | 2 | 0 | 0 | 0 | 0 |
| 57 | 2P | 2 | 0 | 0 | 0 | 0 |
| 57 | 2Q | 1 | 0 | 0 | 0 | 0 |
| 57 | 2R | 4 | 0 | 0 | 0 | 0 |
| 57 | 2T | 3 | 0 | 0 | 0 | 0 |
| 57 | 2U | 2 | 0 | 0 | 0 | 0 |
| 57 | 2V | 1 | 0 | 0 | 0 | 0 |
| 57 | 2W | 2 | 0 | 0 | 0 | 0 |
| 57 | 2X | 1 | 0 | 0 | 0 | 0 |
| 57 | 2Y | 1 | 0 | 0 | 0 | 0 |
| 57 | 2Z | 1 | 0 | 0 | 0 | 0 |
| 57 | 2a | 225 | 0 | 0 | 0 | 0 |
| 57 | 2d | 2 | 0 | 0 | 0 | 0 |
| 57 | 2e | 1 | 0 | 0 | 0 | 0 |
| 57 | 2f | 2 | 0 | 0 | 0 | 0 |
| 57 | 2g | 1 | 0 | 0 | 0 | 0 |
| 57 | 2j | 1 | 0 | 0 | 0 | 0 |
| 57 | 2k | 1 | 0 | 0 | 0 | 0 |
| 57 | 2l | 4 | 0 | 0 | 0 | 0 |
| 57 | 2o | 1 | 0 | 0 | 0 | 0 |
| 57 | 2q | 3 | 0 | 0 | 0 | 0 |
| 57 | 2r | 1 | 0 | 0 | 0 | 0 |
| 57 | 2t | 1 | 0 | 0 | 0 | 0 |
| 57 | 2v | 2 | 0 | 0 | 0 | 0 |
| 57 | 2w | 5 | 0 | 0 | 0 | 0 |
| 57 | 2x | 6 | 0 | 0 | 0 | 0 |
| 57 | 2y | 1 | 0 | 0 | 0 | 0 |
| 57 | 2z | 1 | 0 | 0 | 0 | 0 |
| 58 | 1A | 1 | 0 | 0 | 0 | 0 |
| 58 | 2A | 1 | 0 | 0 | 0 | 0 |
| 59 | 1A | 51 | 0 | 67 | 9 | 0 |
| 59 | 2A | 51 | 0 | 67 | 4 | 0 |
| 60 | 14 | 1 | 0 | 0 | 0 | 0 |
| 60 | 15 | 1 | 0 | 0 | 0 | 0 |
| 60 | 16 | 1 | 0 | 0 | 0 | 0 |
| 60 | 19 | 1 | 0 | 0 | 0 | 0 |
| 60 | 1Y | 1 | 0 | 0 | 0 | 0 |
| 60 | 1n | 1 | 0 | 0 | 0 | 0 |
| 60 | 24 | 1 | 0 | 0 | 0 | 0 |
| 60 | 25 | 1 | 0 | 0 | 0 | 0 |
| 60 | 26 | 1 | 0 | 0 | 0 | 0 |
| 60 | 29 | 1 | 0 | 0 | 0 | 0 |

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| Mol | Chain | Non-H | H(model) | H(added) | Clashes | Symm-Clashes |
|-----|-------|-------|----------|----------|---------|--------------|
| 60 | 2Y | 1 | 0 | 0 | 0 | 0 |
| 60 | 2n | 1 | 0 | 0 | 0 | 0 |
| 61 | 1d | 8 | 0 | 0 | 1 | 0 |
| 61 | 2d | 8 | 0 | 0 | 2 | 0 |
| 62 | 10 | 8 | 0 | 0 | 1 | 0 |
| 62 | 11 | 7 | 0 | 0 | 0 | 0 |
| 62 | 12 | 3 | 0 | 0 | 0 | 0 |
| 62 | 13 | 4 | 0 | 0 | 0 | 0 |
| 62 | 14 | 1 | 0 | 0 | 0 | 0 |
| 62 | 15 | 6 | 0 | 0 | 0 | 0 |
| 62 | 16 | 3 | 0 | 0 | 0 | 0 |
| 62 | 17 | 10 | 0 | 0 | 1 | 0 |
| 62 | 18 | 8 | 0 | 0 | 0 | 0 |
| 62 | 1A | 1680 | 0 | 0 | 117 | 0 |
| 62 | 1B | 54 | 0 | 0 | 3 | 0 |
| 62 | 1D | 32 | 0 | 0 | 3 | 0 |
| 62 | 1E | 23 | 0 | 0 | 1 | 0 |
| 62 | 1F | 17 | 0 | 0 | 2 | 0 |
| 62 | 1G | 1 | 0 | 0 | 0 | 0 |
| 62 | 1H | 1 | 0 | 0 | 0 | 0 |
| 62 | 1N | 4 | 0 | 0 | 0 | 0 |
| 62 | 1O | 3 | 0 | 0 | 0 | 0 |
| 62 | 1P | 15 | 0 | 0 | 3 | 0 |
| 62 | 1Q | 8 | 0 | 0 | 1 | 0 |
| 62 | 1R | 12 | 0 | 0 | 1 | 0 |
| 62 | 1S | 3 | 0 | 0 | 0 | 0 |
| 62 | 1T | 6 | 0 | 0 | 0 | 0 |
| 62 | 1U | 8 | 0 | 0 | 0 | 0 |
| 62 | 1V | 9 | 0 | 0 | 0 | 0 |
| 62 | 1W | 9 | 0 | 0 | 0 | 0 |
| 62 | 1X | 6 | 0 | 0 | 1 | 0 |
| 62 | 1Y | 1 | 0 | 0 | 0 | 0 |
| 62 | 1Z | 1 | 0 | 0 | 0 | 0 |
| 62 | 1a | 252 | 0 | 0 | 20 | 0 |
| 62 | 1b | 1 | 0 | 0 | 0 | 0 |
| 62 | 1d | 1 | 0 | 0 | 0 | 0 |
| 62 | 1f | 1 | 0 | 0 | 0 | 0 |
| 62 | 1l | 3 | 0 | 0 | 0 | 0 |
| 62 | 1m | 2 | 0 | 0 | 0 | 0 |
| 62 | 1o | 1 | 0 | 0 | 0 | 0 |
| 62 | 1p | 1 | 0 | 0 | 0 | 0 |
| 62 | 1q | 2 | 0 | 0 | 0 | 0 |

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| Mol | Chain | Non-H | H(model) | H(added) | Clashes | Symm-Clashes |
|-----|-------|--------|----------|----------|---------|--------------|
| 62 | 1u | 1 | 0 | 0 | 0 | 0 |
| 62 | 1v | 3 | 0 | 0 | 0 | 0 |
| 62 | 1w | 8 | 0 | 0 | 0 | 0 |
| 62 | 1x | 7 | 0 | 0 | 1 | 0 |
| 62 | 1y | 1 | 0 | 0 | 0 | 0 |
| 62 | 1z | 1 | 0 | 0 | 0 | 0 |
| 62 | 20 | 2 | 0 | 0 | 0 | 0 |
| 62 | 21 | 6 | 0 | 0 | 1 | 0 |
| 62 | 23 | 1 | 0 | 0 | 0 | 0 |
| 62 | 25 | 2 | 0 | 0 | 0 | 0 |
| 62 | 27 | 2 | 0 | 0 | 0 | 0 |
| 62 | 28 | 3 | 0 | 0 | 0 | 0 |
| 62 | 29 | 1 | 0 | 0 | 0 | 0 |
| 62 | 2A | 952 | 0 | 0 | 89 | 0 |
| 62 | 2B | 21 | 0 | 0 | 2 | 0 |
| 62 | 2D | 15 | 0 | 0 | 0 | 0 |
| 62 | 2E | 5 | 0 | 0 | 1 | 0 |
| 62 | 2F | 13 | 0 | 0 | 1 | 0 |
| 62 | 2O | 3 | 0 | 0 | 0 | 0 |
| 62 | 2P | 6 | 0 | 0 | 0 | 0 |
| 62 | 2Q | 1 | 0 | 0 | 0 | 0 |
| 62 | 2R | 4 | 0 | 0 | 0 | 0 |
| 62 | 2T | 2 | 0 | 0 | 0 | 0 |
| 62 | 2U | 2 | 0 | 0 | 0 | 0 |
| 62 | 2V | 1 | 0 | 0 | 0 | 0 |
| 62 | 2W | 1 | 0 | 0 | 0 | 0 |
| 62 | 2X | 5 | 0 | 0 | 0 | 0 |
| 62 | 2a | 150 | 0 | 0 | 13 | 0 |
| 62 | 2e | 1 | 0 | 0 | 0 | 0 |
| 62 | 2j | 1 | 0 | 0 | 1 | 0 |
| 62 | 2l | 4 | 0 | 0 | 0 | 0 |
| 62 | 2q | 1 | 0 | 0 | 0 | 0 |
| 62 | 2v | 3 | 0 | 0 | 0 | 0 |
| 62 | 2w | 3 | 0 | 0 | 0 | 0 |
| 62 | 2x | 4 | 0 | 0 | 1 | 0 |
| 62 | 2z | 1 | 0 | 0 | 0 | 0 |
| All | All | 299368 | 0 | 196882 | 4775 | 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 10.

The worst 5 of 4775 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) |
|-------------------|------------------|--------------------------|-------------------|
| 1:1A:1082:U:H3 | 1:1A:1086:A:N6 | 1.23 | 1.36 |
| 22:20:10:THR:HG22 | 22:20:12:ASN:H | 1.28 | 0.97 |
| 1:1A:1082:U:N3 | 1:1A:1086:A:N6 | 2.06 | 0.96 |
| 1:1A:1603:A:OP1 | 62:1A:4102:HOH:O | 1.84 | 0.94 |
| 36:1e:78:HIS:HE1 | 36:1e:143:ARG:H | 1.15 | 0.93 |

There are no symmetry-related clashes.

5.3 Torsion angles [i](#)

5.3.1 Protein backbone [i](#)

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%) | 250 (92%) | 22 (8%) | 1 (0%) | 30 | 47 |
| 3 | 2D | 273/276 (99%) | 245 (90%) | 25 (9%) | 3 (1%) | 12 | 21 |
| 4 | 1E | 202/206 (98%) | 182 (90%) | 19 (9%) | 1 (0%) | 25 | 41 |
| 4 | 2E | 202/206 (98%) | 181 (90%) | 20 (10%) | 1 (0%) | 25 | 41 |
| 5 | 1F | 200/210 (95%) | 187 (94%) | 11 (6%) | 2 (1%) | 13 | 23 |
| 5 | 2F | 200/210 (95%) | 179 (90%) | 18 (9%) | 3 (2%) | 8 | 14 |
| 6 | 1G | 179/182 (98%) | 154 (86%) | 25 (14%) | 0 | 100 | 100 |
| 6 | 2G | 179/182 (98%) | 141 (79%) | 28 (16%) | 10 (6%) | 1 | 1 |
| 7 | 1H | 172/180 (96%) | 162 (94%) | 9 (5%) | 1 (1%) | 22 | 36 |
| 7 | 2H | 172/180 (96%) | 142 (83%) | 27 (16%) | 3 (2%) | 7 | 13 |
| 8 | 1I | 144/148 (97%) | 125 (87%) | 19 (13%) | 0 | 100 | 100 |
| 8 | 2I | 144/148 (97%) | 107 (74%) | 32 (22%) | 5 (4%) | 3 | 4 |
| 9 | 1N | 138/140 (99%) | 131 (95%) | 7 (5%) | 0 | 100 | 100 |
| 9 | 2N | 138/140 (99%) | 126 (91%) | 10 (7%) | 2 (1%) | 9 | 16 |
| 10 | 1O | 120/122 (98%) | 112 (93%) | 6 (5%) | 2 (2%) | 7 | 13 |
| 10 | 2O | 120/122 (98%) | 109 (91%) | 8 (7%) | 3 (2%) | 4 | 7 |

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| Mol | Chain | Analysed | Favoured | Allowed | Outliers | Percentiles | |
|-----|-------|---------------|-----------|----------|----------|-------------|-----|
| 11 | 1P | 147/150 (98%) | 130 (88%) | 11 (8%) | 6 (4%) | 2 | 2 |
| 11 | 2P | 147/150 (98%) | 129 (88%) | 14 (10%) | 4 (3%) | 4 | 6 |
| 12 | 1Q | 139/141 (99%) | 129 (93%) | 10 (7%) | 0 | 100 | 100 |
| 12 | 2Q | 139/141 (99%) | 121 (87%) | 17 (12%) | 1 (1%) | 19 | 32 |
| 13 | 1R | 116/118 (98%) | 110 (95%) | 4 (3%) | 2 (2%) | 7 | 13 |
| 13 | 2R | 116/118 (98%) | 108 (93%) | 8 (7%) | 0 | 100 | 100 |
| 14 | 1S | 108/112 (96%) | 101 (94%) | 7 (6%) | 0 | 100 | 100 |
| 14 | 2S | 108/112 (96%) | 96 (89%) | 9 (8%) | 3 (3%) | 4 | 6 |
| 15 | 1T | 129/146 (88%) | 114 (88%) | 14 (11%) | 1 (1%) | 16 | 29 |
| 15 | 2T | 129/146 (88%) | 122 (95%) | 6 (5%) | 1 (1%) | 16 | 29 |
| 16 | 1U | 114/118 (97%) | 113 (99%) | 1 (1%) | 0 | 100 | 100 |
| 16 | 2U | 114/118 (97%) | 109 (96%) | 5 (4%) | 0 | 100 | 100 |
| 17 | 1V | 99/101 (98%) | 92 (93%) | 4 (4%) | 3 (3%) | 3 | 5 |
| 17 | 2V | 99/101 (98%) | 88 (89%) | 9 (9%) | 2 (2%) | 6 | 11 |
| 18 | 1W | 110/113 (97%) | 107 (97%) | 3 (3%) | 0 | 100 | 100 |
| 18 | 2W | 110/113 (97%) | 102 (93%) | 8 (7%) | 0 | 100 | 100 |
| 19 | 1X | 93/96 (97%) | 87 (94%) | 6 (6%) | 0 | 100 | 100 |
| 19 | 2X | 93/96 (97%) | 81 (87%) | 11 (12%) | 1 (1%) | 12 | 21 |
| 20 | 1Y | 105/110 (96%) | 94 (90%) | 8 (8%) | 3 (3%) | 3 | 5 |
| 20 | 2Y | 105/110 (96%) | 95 (90%) | 8 (8%) | 2 (2%) | 6 | 11 |
| 21 | 1Z | 148/206 (72%) | 126 (85%) | 18 (12%) | 4 (3%) | 4 | 6 |
| 21 | 2Z | 156/206 (76%) | 123 (79%) | 24 (15%) | 9 (6%) | 1 | 1 |
| 22 | 10 | 81/85 (95%) | 77 (95%) | 4 (5%) | 0 | 100 | 100 |
| 22 | 20 | 81/85 (95%) | 77 (95%) | 3 (4%) | 1 (1%) | 11 | 19 |
| 23 | 11 | 95/98 (97%) | 91 (96%) | 3 (3%) | 1 (1%) | 12 | 21 |
| 23 | 21 | 95/98 (97%) | 90 (95%) | 4 (4%) | 1 (1%) | 12 | 21 |
| 24 | 12 | 68/72 (94%) | 66 (97%) | 2 (3%) | 0 | 100 | 100 |
| 24 | 22 | 68/72 (94%) | 63 (93%) | 5 (7%) | 0 | 100 | 100 |
| 25 | 13 | 57/60 (95%) | 55 (96%) | 2 (4%) | 0 | 100 | 100 |
| 25 | 23 | 57/60 (95%) | 52 (91%) | 3 (5%) | 2 (4%) | 3 | 4 |
| 26 | 14 | 67/71 (94%) | 46 (69%) | 15 (22%) | 6 (9%) | 0 | 0 |

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| Mol | Chain | Analysed | Favoured | Allowed | Outliers | Percentiles | |
|-----|-------|---------------|-----------|----------|----------|-------------|-----|
| 26 | 24 | 67/71 (94%) | 47 (70%) | 15 (22%) | 5 (8%) | 1 | 1 |
| 27 | 15 | 57/60 (95%) | 54 (95%) | 3 (5%) | 0 | 100 | 100 |
| 27 | 25 | 57/60 (95%) | 52 (91%) | 5 (9%) | 0 | 100 | 100 |
| 28 | 16 | 51/54 (94%) | 46 (90%) | 5 (10%) | 0 | 100 | 100 |
| 28 | 26 | 51/54 (94%) | 44 (86%) | 7 (14%) | 0 | 100 | 100 |
| 29 | 17 | 46/49 (94%) | 46 (100%) | 0 | 0 | 100 | 100 |
| 29 | 27 | 46/49 (94%) | 45 (98%) | 1 (2%) | 0 | 100 | 100 |
| 30 | 18 | 62/65 (95%) | 62 (100%) | 0 | 0 | 100 | 100 |
| 30 | 28 | 62/65 (95%) | 58 (94%) | 4 (6%) | 0 | 100 | 100 |
| 31 | 19 | 35/37 (95%) | 33 (94%) | 2 (6%) | 0 | 100 | 100 |
| 31 | 29 | 35/37 (95%) | 31 (89%) | 4 (11%) | 0 | 100 | 100 |
| 33 | 1b | 229/256 (90%) | 184 (80%) | 31 (14%) | 14 (6%) | 1 | 1 |
| 33 | 2b | 229/256 (90%) | 170 (74%) | 51 (22%) | 8 (4%) | 3 | 4 |
| 34 | 1c | 204/239 (85%) | 185 (91%) | 16 (8%) | 3 (2%) | 8 | 14 |
| 34 | 2c | 204/239 (85%) | 153 (75%) | 48 (24%) | 3 (2%) | 8 | 14 |
| 35 | 1d | 206/209 (99%) | 191 (93%) | 12 (6%) | 3 (2%) | 8 | 14 |
| 35 | 2d | 206/209 (99%) | 173 (84%) | 27 (13%) | 6 (3%) | 3 | 5 |
| 36 | 1e | 146/162 (90%) | 134 (92%) | 10 (7%) | 2 (1%) | 9 | 16 |
| 36 | 2e | 146/162 (90%) | 116 (80%) | 23 (16%) | 7 (5%) | 2 | 2 |
| 37 | 1f | 98/101 (97%) | 91 (93%) | 7 (7%) | 0 | 100 | 100 |
| 37 | 2f | 98/101 (97%) | 87 (89%) | 11 (11%) | 0 | 100 | 100 |
| 38 | 1g | 153/156 (98%) | 137 (90%) | 13 (8%) | 3 (2%) | 6 | 11 |
| 38 | 2g | 153/156 (98%) | 129 (84%) | 21 (14%) | 3 (2%) | 6 | 11 |
| 39 | 1h | 135/138 (98%) | 120 (89%) | 13 (10%) | 2 (2%) | 8 | 14 |
| 39 | 2h | 135/138 (98%) | 122 (90%) | 11 (8%) | 2 (2%) | 8 | 14 |
| 40 | 1i | 125/128 (98%) | 105 (84%) | 17 (14%) | 3 (2%) | 5 | 8 |
| 40 | 2i | 125/128 (98%) | 105 (84%) | 16 (13%) | 4 (3%) | 3 | 5 |
| 41 | 1j | 95/105 (90%) | 80 (84%) | 11 (12%) | 4 (4%) | 2 | 2 |
| 41 | 2j | 94/105 (90%) | 72 (77%) | 17 (18%) | 5 (5%) | 1 | 1 |
| 42 | 1k | 112/129 (87%) | 102 (91%) | 9 (8%) | 1 (1%) | 14 | 26 |
| 42 | 2k | 112/129 (87%) | 96 (86%) | 13 (12%) | 3 (3%) | 4 | 6 |

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| Mol | Chain | Analysed | Favoured | Allowed | Outliers | Percentiles | |
|-----|-------|-------------------|-------------|------------|----------|-------------|-----|
| 43 | 1l | 119/132 (90%) | 108 (91%) | 10 (8%) | 1 (1%) | 16 | 29 |
| 43 | 2l | 119/132 (90%) | 108 (91%) | 9 (8%) | 2 (2%) | 7 | 13 |
| 44 | 1m | 121/126 (96%) | 102 (84%) | 16 (13%) | 3 (2%) | 4 | 7 |
| 44 | 2m | 120/126 (95%) | 99 (82%) | 17 (14%) | 4 (3%) | 3 | 4 |
| 45 | 1n | 58/61 (95%) | 54 (93%) | 4 (7%) | 0 | 100 | 100 |
| 45 | 2n | 58/61 (95%) | 50 (86%) | 6 (10%) | 2 (3%) | 3 | 4 |
| 46 | 1o | 86/89 (97%) | 74 (86%) | 8 (9%) | 4 (5%) | 2 | 2 |
| 46 | 2o | 86/89 (97%) | 79 (92%) | 7 (8%) | 0 | 100 | 100 |
| 47 | 1p | 80/88 (91%) | 69 (86%) | 10 (12%) | 1 (1%) | 10 | 17 |
| 47 | 2p | 80/88 (91%) | 72 (90%) | 7 (9%) | 1 (1%) | 10 | 17 |
| 48 | 1q | 97/105 (92%) | 87 (90%) | 10 (10%) | 0 | 100 | 100 |
| 48 | 2q | 97/105 (92%) | 78 (80%) | 16 (16%) | 3 (3%) | 3 | 5 |
| 49 | 1r | 66/88 (75%) | 57 (86%) | 8 (12%) | 1 (2%) | 8 | 14 |
| 49 | 2r | 66/88 (75%) | 59 (89%) | 7 (11%) | 0 | 100 | 100 |
| 50 | 1s | 81/93 (87%) | 66 (82%) | 15 (18%) | 0 | 100 | 100 |
| 50 | 2s | 81/93 (87%) | 65 (80%) | 14 (17%) | 2 (2%) | 4 | 7 |
| 51 | 1t | 94/106 (89%) | 81 (86%) | 8 (8%) | 5 (5%) | 1 | 1 |
| 51 | 2t | 94/106 (89%) | 82 (87%) | 11 (12%) | 1 (1%) | 12 | 21 |
| 52 | 1u | 21/27 (78%) | 18 (86%) | 3 (14%) | 0 | 100 | 100 |
| 52 | 2u | 21/27 (78%) | 13 (62%) | 6 (29%) | 2 (10%) | 0 | 0 |
| 56 | 1z | 1/3 (33%) | 0 | 1 (100%) | 0 | 100 | 100 |
| 56 | 2z | 1/3 (33%) | 1 (100%) | 0 | 0 | 100 | 100 |
| All | All | 11370/12134 (94%) | 10019 (88%) | 1148 (10%) | 203 (2%) | 7 | 12 |

5 of 203 Ramachandran outliers are listed below:

| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 5 | 1F | 130 | ALA |
| 11 | 1P | 38 | GLN |
| 21 | 1Z | 53 | ILE |
| 21 | 1Z | 93 | ASP |
| 26 | 14 | 46 | GLN |

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%) | 202 (94%) | 13 (6%) | 16 | 29 |
| 3 | 2D | 215/218 (99%) | 197 (92%) | 18 (8%) | 9 | 16 |
| 4 | 1E | 164/166 (99%) | 149 (91%) | 15 (9%) | 7 | 13 |
| 4 | 2E | 164/166 (99%) | 145 (88%) | 19 (12%) | 4 | 7 |
| 5 | 1F | 160/166 (96%) | 142 (89%) | 18 (11%) | 4 | 8 |
| 5 | 2F | 159/166 (96%) | 145 (91%) | 14 (9%) | 8 | 15 |
| 6 | 1G | 143/156 (92%) | 129 (90%) | 14 (10%) | 6 | 11 |
| 6 | 2G | 143/156 (92%) | 125 (87%) | 18 (13%) | 3 | 5 |
| 7 | 1H | 144/148 (97%) | 133 (92%) | 11 (8%) | 11 | 19 |
| 7 | 2H | 144/148 (97%) | 124 (86%) | 20 (14%) | 3 | 4 |
| 8 | 1I | 113/124 (91%) | 99 (88%) | 14 (12%) | 4 | 6 |
| 8 | 2I | 105/124 (85%) | 93 (89%) | 12 (11%) | 4 | 7 |
| 9 | 1N | 118/119 (99%) | 109 (92%) | 9 (8%) | 11 | 19 |
| 9 | 2N | 118/119 (99%) | 104 (88%) | 14 (12%) | 4 | 7 |
| 10 | 1O | 100/100 (100%) | 96 (96%) | 4 (4%) | 27 | 47 |
| 10 | 2O | 100/100 (100%) | 92 (92%) | 8 (8%) | 10 | 18 |
| 11 | 1P | 115/116 (99%) | 99 (86%) | 16 (14%) | 3 | 4 |
| 11 | 2P | 115/116 (99%) | 102 (89%) | 13 (11%) | 4 | 8 |
| 12 | 1Q | 111/111 (100%) | 97 (87%) | 14 (13%) | 3 | 5 |
| 12 | 2Q | 111/111 (100%) | 100 (90%) | 11 (10%) | 6 | 11 |
| 13 | 1R | 101/101 (100%) | 96 (95%) | 5 (5%) | 20 | 37 |
| 13 | 2R | 101/101 (100%) | 98 (97%) | 3 (3%) | 36 | 58 |
| 14 | 1S | 86/88 (98%) | 73 (85%) | 13 (15%) | 2 | 3 |
| 14 | 2S | 85/88 (97%) | 65 (76%) | 20 (24%) | 0 | 0 |
| 15 | 1T | 115/127 (91%) | 108 (94%) | 7 (6%) | 15 | 28 |
| 15 | 2T | 113/127 (89%) | 104 (92%) | 9 (8%) | 10 | 18 |

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| Mol | Chain | Analysed | Rotameric | Outliers | Percentiles | |
|-----|-------|---------------|-----------|----------|-------------|----|
| 16 | 1U | 93/94 (99%) | 86 (92%) | 7 (8%) | 11 | 20 |
| 16 | 2U | 93/94 (99%) | 86 (92%) | 7 (8%) | 11 | 20 |
| 17 | 1V | 80/82 (98%) | 71 (89%) | 9 (11%) | 4 | 8 |
| 17 | 2V | 80/82 (98%) | 69 (86%) | 11 (14%) | 3 | 4 |
| 18 | 1W | 90/92 (98%) | 82 (91%) | 8 (9%) | 8 | 14 |
| 18 | 2W | 90/92 (98%) | 83 (92%) | 7 (8%) | 10 | 19 |
| 19 | 1X | 77/78 (99%) | 75 (97%) | 2 (3%) | 41 | 63 |
| 19 | 2X | 77/78 (99%) | 69 (90%) | 8 (10%) | 5 | 10 |
| 20 | 1Y | 85/91 (93%) | 76 (89%) | 9 (11%) | 5 | 9 |
| 20 | 2Y | 85/91 (93%) | 72 (85%) | 13 (15%) | 2 | 3 |
| 21 | 1Z | 135/179 (75%) | 109 (81%) | 26 (19%) | 1 | 1 |
| 21 | 2Z | 137/179 (76%) | 114 (83%) | 23 (17%) | 1 | 2 |
| 22 | 10 | 65/67 (97%) | 63 (97%) | 2 (3%) | 35 | 56 |
| 22 | 20 | 65/67 (97%) | 60 (92%) | 5 (8%) | 10 | 19 |
| 23 | 11 | 80/83 (96%) | 71 (89%) | 9 (11%) | 4 | 8 |
| 23 | 21 | 80/83 (96%) | 74 (92%) | 6 (8%) | 11 | 20 |
| 24 | 12 | 65/67 (97%) | 57 (88%) | 8 (12%) | 4 | 6 |
| 24 | 22 | 65/67 (97%) | 58 (89%) | 7 (11%) | 5 | 9 |
| 25 | 13 | 51/52 (98%) | 44 (86%) | 7 (14%) | 3 | 4 |
| 25 | 23 | 50/52 (96%) | 43 (86%) | 7 (14%) | 3 | 4 |
| 26 | 14 | 59/63 (94%) | 48 (81%) | 11 (19%) | 1 | 2 |
| 26 | 24 | 53/63 (84%) | 41 (77%) | 12 (23%) | 1 | 0 |
| 27 | 15 | 50/52 (96%) | 47 (94%) | 3 (6%) | 16 | 29 |
| 27 | 25 | 50/52 (96%) | 47 (94%) | 3 (6%) | 16 | 29 |
| 28 | 16 | 51/52 (98%) | 42 (82%) | 9 (18%) | 1 | 2 |
| 28 | 26 | 50/52 (96%) | 41 (82%) | 9 (18%) | 1 | 2 |
| 29 | 17 | 41/42 (98%) | 38 (93%) | 3 (7%) | 11 | 21 |
| 29 | 27 | 41/42 (98%) | 36 (88%) | 5 (12%) | 4 | 6 |
| 30 | 18 | 54/55 (98%) | 52 (96%) | 2 (4%) | 29 | 50 |
| 30 | 28 | 54/55 (98%) | 50 (93%) | 4 (7%) | 11 | 20 |
| 31 | 19 | 34/34 (100%) | 32 (94%) | 2 (6%) | 16 | 30 |

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| Mol | Chain | Analysed | Rotameric | Outliers | Percentiles | |
|-----|-------|---------------|-----------|----------|-------------|----|
| 31 | 29 | 34/34 (100%) | 29 (85%) | 5 (15%) | 2 | 3 |
| 33 | 1b | 192/220 (87%) | 164 (85%) | 28 (15%) | 2 | 3 |
| 33 | 2b | 187/220 (85%) | 148 (79%) | 39 (21%) | 1 | 1 |
| 34 | 1c | 142/188 (76%) | 128 (90%) | 14 (10%) | 6 | 11 |
| 34 | 2c | 140/188 (74%) | 121 (86%) | 19 (14%) | 3 | 4 |
| 35 | 1d | 169/181 (93%) | 151 (89%) | 18 (11%) | 5 | 9 |
| 35 | 2d | 173/181 (96%) | 151 (87%) | 22 (13%) | 3 | 5 |
| 36 | 1e | 113/123 (92%) | 100 (88%) | 13 (12%) | 4 | 7 |
| 36 | 2e | 114/123 (93%) | 95 (83%) | 19 (17%) | 2 | 2 |
| 37 | 1f | 84/90 (93%) | 74 (88%) | 10 (12%) | 4 | 7 |
| 37 | 2f | 85/90 (94%) | 74 (87%) | 11 (13%) | 3 | 5 |
| 38 | 1g | 119/127 (94%) | 103 (87%) | 16 (13%) | 3 | 4 |
| 38 | 2g | 120/127 (94%) | 97 (81%) | 23 (19%) | 1 | 1 |
| 39 | 1h | 114/119 (96%) | 100 (88%) | 14 (12%) | 4 | 6 |
| 39 | 2h | 114/119 (96%) | 102 (90%) | 12 (10%) | 5 | 9 |
| 40 | 1i | 90/99 (91%) | 79 (88%) | 11 (12%) | 4 | 6 |
| 40 | 2i | 89/99 (90%) | 79 (89%) | 10 (11%) | 5 | 8 |
| 41 | 1j | 66/92 (72%) | 56 (85%) | 10 (15%) | 2 | 3 |
| 41 | 2j | 69/92 (75%) | 59 (86%) | 10 (14%) | 2 | 3 |
| 42 | 1k | 82/99 (83%) | 70 (85%) | 12 (15%) | 2 | 3 |
| 42 | 2k | 83/99 (84%) | 75 (90%) | 8 (10%) | 7 | 12 |
| 43 | 1l | 96/108 (89%) | 90 (94%) | 6 (6%) | 15 | 27 |
| 43 | 2l | 96/108 (89%) | 84 (88%) | 12 (12%) | 3 | 6 |
| 44 | 1m | 93/101 (92%) | 84 (90%) | 9 (10%) | 6 | 12 |
| 44 | 2m | 92/101 (91%) | 77 (84%) | 15 (16%) | 2 | 2 |
| 45 | 1n | 49/50 (98%) | 45 (92%) | 4 (8%) | 9 | 17 |
| 45 | 2n | 49/50 (98%) | 44 (90%) | 5 (10%) | 6 | 10 |
| 46 | 1o | 78/80 (98%) | 73 (94%) | 5 (6%) | 14 | 27 |
| 46 | 2o | 78/80 (98%) | 75 (96%) | 3 (4%) | 28 | 49 |
| 47 | 1p | 69/74 (93%) | 56 (81%) | 13 (19%) | 1 | 1 |
| 47 | 2p | 68/74 (92%) | 59 (87%) | 9 (13%) | 3 | 5 |

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| Mol | Chain | Analysed | Rotameric | Outliers | Percentiles | |
|-----|-------|------------------|------------|------------|-------------|----|
| 48 | 1q | 94/97 (97%) | 84 (89%) | 10 (11%) | 5 | 9 |
| 48 | 2q | 94/97 (97%) | 82 (87%) | 12 (13%) | 3 | 5 |
| 49 | 1r | 59/77 (77%) | 53 (90%) | 6 (10%) | 6 | 10 |
| 49 | 2r | 59/77 (77%) | 49 (83%) | 10 (17%) | 1 | 2 |
| 50 | 1s | 69/80 (86%) | 62 (90%) | 7 (10%) | 6 | 10 |
| 50 | 2s | 67/80 (84%) | 51 (76%) | 16 (24%) | 0 | 0 |
| 51 | 1t | 70/82 (85%) | 61 (87%) | 9 (13%) | 3 | 5 |
| 51 | 2t | 70/82 (85%) | 62 (89%) | 8 (11%) | 4 | 7 |
| 52 | 1u | 18/22 (82%) | 16 (89%) | 2 (11%) | 5 | 8 |
| 52 | 2u | 18/22 (82%) | 16 (89%) | 2 (11%) | 5 | 8 |
| 56 | 1z | 2/2 (100%) | 1 (50%) | 1 (50%) | 0 | 0 |
| 56 | 2z | 2/2 (100%) | 1 (50%) | 1 (50%) | 0 | 0 |
| All | All | 9307/10068 (92%) | 8242 (89%) | 1065 (11%) | 4 | 7 |

5 of 1065 residues with a non-rotameric sidechain are listed below:

| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 38 | 2g | 113 | GLU |
| 40 | 2i | 108 | VAL |
| 38 | 2g | 106 | GLN |
| 50 | 2s | 27 | GLU |
| 40 | 1i | 64 | THR |

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. 5 of 141 such sidechains are listed below:

| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 36 | 2e | 127 | ASN |
| 38 | 2g | 97 | GLN |
| 46 | 2o | 13 | GLN |
| 38 | 1g | 86 | GLN |
| 38 | 1g | 28 | ASN |

5.3.3 RNA ⓘ

| Mol | Chain | Analysed | Backbone Outliers | Pucker Outliers |
|-----|-------|-----------------|-------------------|-----------------|
| 1 | 1A | 2864/2915 (98%) | 505 (17%) | 24 (0%) |

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| Mol | Chain | Analysed | Backbone Outliers | Pucker Outliers |
|-----|-------|-----------------|-------------------|-----------------|
| 1 | 2A | 2791/2915 (95%) | 520 (18%) | 23 (0%) |
| 2 | 1B | 119/121 (98%) | 7 (5%) | 0 |
| 2 | 2B | 118/121 (97%) | 30 (25%) | 0 |
| 32 | 1a | 1497/1521 (98%) | 277 (18%) | 0 |
| 32 | 2a | 1501/1521 (98%) | 347 (23%) | 0 |
| 53 | 1v | 12/24 (50%) | 2 (16%) | 0 |
| 53 | 2v | 12/24 (50%) | 5 (41%) | 0 |
| 54 | 1w | 72/76 (94%) | 24 (33%) | 0 |
| 54 | 1y | 72/76 (94%) | 26 (36%) | 0 |
| 54 | 2w | 69/76 (90%) | 24 (34%) | 0 |
| 54 | 2y | 70/76 (92%) | 25 (35%) | 0 |
| 55 | 1x | 75/77 (97%) | 16 (21%) | 0 |
| 55 | 2x | 75/77 (97%) | 13 (17%) | 0 |
| All | All | 9347/9620 (97%) | 1821 (19%) | 47 (0%) |

5 of 1821 RNA backbone outliers are listed below:

| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 1 | 1A | 10 | G |
| 1 | 1A | 11 | G |
| 1 | 1A | 12 | U |
| 1 | 1A | 15 | G |
| 1 | 1A | 27 | G |

5 of 47 RNA pucker outliers are listed below:

| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | 2A | 277 | C |
| 1 | 2A | 1210 | A |
| 1 | 2A | 479 | A |
| 1 | 2A | 752 | A |
| 1 | 2A | 1420 | U |

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

88 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$ |
| 1 | 5MU | 2A | 1915 | 1,57 | 19,22,23 | 1.45 | 5 (26%) | 27,32,35 | 2.05 | 5 (18%) |
| 32 | PSU | 2a | 516 | 32,57 | 18,21,22 | 1.35 | 2 (11%) | 21,30,33 | 2.01 | 5 (23%) |
| 55 | PSU | 2x | 55 | 55 | 18,21,22 | 1.38 | 3 (16%) | 21,30,33 | 1.94 | 3 (14%) |
| 54 | PSU | 2w | 55 | 54 | 18,21,22 | 1.41 | 2 (11%) | 21,30,33 | 2.05 | 4 (19%) |
| 32 | 5MC | 2a | 1407 | 32 | 19,22,23 | 1.64 | 3 (15%) | 26,32,35 | 1.20 | 3 (11%) |
| 54 | G7M | 1w | 46 | 54 | 20,26,27 | 1.19 | 1 (5%) | 16,39,42 | 0.91 | 0 |
| 32 | PSU | 1a | 516 | 32,57 | 18,21,22 | 1.34 | 2 (11%) | 21,30,33 | 2.12 | 4 (19%) |
| 1 | OMU | 2A | 2552 | 1,57 | 19,22,23 | 1.20 | 3 (15%) | 25,31,34 | 1.74 | 5 (20%) |
| 54 | 4SU | 2y | 8 | 54 | 18,21,22 | 1.61 | 3 (16%) | 25,30,33 | 2.26 | 5 (20%) |
| 32 | 4OC | 2a | 1402 | 32,57 | 20,23,24 | 0.78 | 0 | 25,32,35 | 1.00 | 1 (4%) |
| 54 | 5MU | 2w | 54 | 54 | 19,22,23 | 1.46 | 4 (21%) | 27,32,35 | 1.62 | 5 (18%) |
| 1 | PSU | 2A | 1911 | 1 | 18,21,22 | 1.39 | 3 (16%) | 21,30,33 | 1.94 | 3 (14%) |
| 32 | 4OC | 1a | 1402 | 32 | 20,23,24 | 0.75 | 0 | 25,32,35 | 0.92 | 1 (4%) |
| 32 | MA6 | 2a | 1518 | 32 | 19,26,27 | 1.03 | 2 (10%) | 18,38,41 | 1.93 | 3 (16%) |
| 54 | PSU | 2w | 32 | 54 | 18,21,22 | 1.37 | 2 (11%) | 21,30,33 | 1.88 | 3 (14%) |
| 32 | 5MC | 2a | 967 | 32 | 19,22,23 | 1.72 | 3 (15%) | 26,32,35 | 1.14 | 3 (11%) |
| 56 | FME | 1z | 1 | 56 | 8,9,10 | 0.99 | 0 | 8,9,11 | 0.70 | 0 |
| 32 | 2MG | 2a | 1207 | 32,57 | 18,26,27 | 0.94 | 1 (5%) | 16,38,41 | 1.52 | 4 (25%) |
| 54 | 5MU | 1w | 54 | 54 | 19,22,23 | 1.40 | 4 (21%) | 27,32,35 | 1.95 | 7 (25%) |
| 1 | 2MA | 1A | 2503 | 1,57 | 18,25,26 | 0.69 | 0 | 20,37,40 | 2.10 | 6 (30%) |
| 54 | G7M | 2w | 46 | 54 | 20,26,27 | 1.16 | 1 (5%) | 16,39,42 | 1.00 | 1 (6%) |
| 54 | PSU | 2y | 55 | 54 | 18,21,22 | 1.38 | 2 (11%) | 21,30,33 | 1.91 | 4 (19%) |
| 1 | 5MC | 1A | 1942 | 1 | 19,22,23 | 1.67 | 3 (15%) | 26,32,35 | 1.14 | 3 (11%) |
| 54 | G7M | 2y | 46 | 54 | 20,26,27 | 1.37 | 2 (10%) | 16,39,42 | 0.58 | 0 |
| 54 | 4SU | 2w | 8 | 54 | 18,21,22 | 1.81 | 5 (27%) | 25,30,33 | 2.11 | 5 (20%) |
| 32 | MA6 | 1a | 1518 | 32 | 19,26,27 | 1.02 | 2 (10%) | 18,38,41 | 1.83 | 3 (16%) |
| 32 | 5MC | 2a | 1400 | 32 | 19,22,23 | 1.68 | 3 (15%) | 26,32,35 | 1.18 | 2 (7%) |
| 1 | 5MU | 1A | 1939 | 1,57 | 19,22,23 | 1.56 | 6 (31%) | 27,32,35 | 2.24 | 5 (18%) |
| 1 | OMG | 2A | 2251 | 55,1,57 | 19,26,27 | 0.94 | 1 (5%) | 21,38,41 | 1.24 | 4 (19%) |
| 1 | 2MA | 2A | 2503 | 1,57 | 18,25,26 | 0.71 | 0 | 20,37,40 | 1.90 | 4 (20%) |
| 32 | 5MC | 1a | 1400 | 32 | 19,22,23 | 1.54 | 3 (15%) | 26,32,35 | 1.15 | 3 (11%) |
| 32 | G7M | 1a | 527 | 32,57 | 20,26,27 | 1.22 | 2 (10%) | 16,39,42 | 0.62 | 0 |
| 54 | MIA | 1w | 37 | 54 | 24,31,32 | 2.12 | 4 (16%) | 22,44,47 | 2.16 | 6 (27%) |

| Mol | Type | Chain | Res | Link | Bond lengths | | | Bond angles | | |
|-----|------|-------|------|----------|--------------|------|----------|-------------|------|----------|
| | | | | | Counts | RMSZ | # Z > 2 | Counts | RMSZ | # Z > 2 |
| 55 | 5MC | 2x | 32 | 55 | 19,22,23 | 1.55 | 3 (15%) | 26,32,35 | 1.14 | 3 (11%) |
| 1 | 5MC | 2A | 1962 | 1,57 | 19,22,23 | 1.56 | 3 (15%) | 26,32,35 | 1.21 | 2 (7%) |
| 1 | 5MC | 1A | 1962 | 1,57 | 19,22,23 | 1.69 | 3 (15%) | 26,32,35 | 0.99 | 2 (7%) |
| 1 | PSU | 2A | 2605 | 1 | 18,21,22 | 1.43 | 3 (16%) | 21,30,33 | 2.11 | 5 (23%) |
| 32 | 5MC | 1a | 1404 | 32 | 19,22,23 | 1.67 | 3 (15%) | 26,32,35 | 1.10 | 2 (7%) |
| 55 | PSU | 1x | 55 | 55 | 18,21,22 | 1.35 | 2 (11%) | 21,30,33 | 2.05 | 3 (14%) |
| 1 | OMC | 2A | 1920 | 1 | 19,22,23 | 0.79 | 0 | 25,31,34 | 0.98 | 1 (4%) |
| 32 | UR3 | 1a | 1498 | 32 | 19,22,23 | 1.07 | 1 (5%) | 26,32,35 | 1.76 | 4 (15%) |
| 54 | PSU | 1y | 39 | 54 | 18,21,22 | 1.38 | 2 (11%) | 21,30,33 | 1.85 | 4 (19%) |
| 32 | M2G | 2a | 966 | 32 | 20,27,28 | 1.42 | 3 (15%) | 19,40,43 | 0.95 | 1 (5%) |
| 55 | 5MU | 2x | 54 | 55 | 19,22,23 | 1.43 | 5 (26%) | 27,32,35 | 2.08 | 6 (22%) |
| 43 | 0TD | 1l | 92 | 43 | 8,9,10 | 4.46 | 1 (12%) | 6,11,13 | 4.45 | 2 (33%) |
| 55 | 8AN | 1x | 76 | 55,57 | 17,24,25 | 1.19 | 2 (11%) | 13,35,38 | 3.18 | 3 (23%) |
| 32 | MA6 | 1a | 1519 | 32 | 19,26,27 | 1.02 | 1 (5%) | 18,38,41 | 2.10 | 3 (16%) |
| 32 | 5MC | 2a | 1404 | 32 | 19,22,23 | 1.74 | 3 (15%) | 26,32,35 | 1.19 | 3 (11%) |
| 1 | OMU | 1A | 2552 | 1,57 | 19,22,23 | 1.18 | 3 (15%) | 25,31,34 | 1.85 | 5 (20%) |
| 54 | PSU | 1w | 39 | 54 | 18,21,22 | 1.26 | 2 (11%) | 21,30,33 | 2.06 | 4 (19%) |
| 54 | G7M | 1y | 46 | 54 | 20,26,27 | 1.28 | 2 (10%) | 16,39,42 | 0.55 | 0 |
| 54 | 5MU | 1y | 54 | 54 | 19,22,23 | 1.45 | 4 (21%) | 27,32,35 | 1.94 | 6 (22%) |
| 32 | G7M | 2a | 527 | 32 | 20,26,27 | 1.18 | 2 (10%) | 16,39,42 | 0.57 | 0 |
| 43 | 0TD | 2l | 92 | 43 | 8,9,10 | 4.51 | 1 (12%) | 6,11,13 | 4.08 | 1 (16%) |
| 32 | M2G | 1a | 966 | 32 | 20,27,28 | 1.38 | 3 (15%) | 19,40,43 | 1.04 | 2 (10%) |
| 55 | 8AN | 2x | 76 | 58,55,57 | 17,24,25 | 1.09 | 2 (11%) | 13,35,38 | 4.07 | 3 (23%) |
| 54 | MIA | 1y | 37 | 54 | 17,24,32 | 1.00 | 1 (5%) | 16,35,47 | 1.28 | 2 (12%) |
| 1 | PSU | 1A | 1911 | 1 | 18,21,22 | 1.44 | 4 (22%) | 21,30,33 | 2.15 | 3 (14%) |
| 1 | 5MU | 1A | 1915 | 1 | 19,22,23 | 1.41 | 6 (31%) | 27,32,35 | 2.08 | 6 (22%) |
| 55 | 5MU | 1x | 54 | 55 | 19,22,23 | 1.38 | 5 (26%) | 27,32,35 | 1.91 | 7 (25%) |
| 54 | 4SU | 1w | 8 | 54 | 18,21,22 | 1.90 | 5 (27%) | 25,30,33 | 1.74 | 5 (20%) |
| 1 | 5MU | 2A | 1939 | 1 | 19,22,23 | 1.46 | 6 (31%) | 27,32,35 | 2.25 | 6 (22%) |
| 1 | PSU | 1A | 2605 | 1,57 | 18,21,22 | 1.35 | 3 (16%) | 21,30,33 | 2.02 | 3 (14%) |
| 32 | UR3 | 2a | 1498 | 32,57 | 19,22,23 | 1.11 | 2 (10%) | 26,32,35 | 1.76 | 4 (15%) |
| 1 | OMC | 1A | 1920 | 1 | 19,22,23 | 0.82 | 0 | 25,31,34 | 1.12 | 2 (8%) |
| 54 | PSU | 1w | 55 | 54 | 18,21,22 | 1.36 | 2 (11%) | 21,30,33 | 1.97 | 3 (14%) |
| 1 | OMG | 1A | 2251 | 55,1,57 | 19,26,27 | 0.97 | 1 (5%) | 21,38,41 | 1.01 | 2 (9%) |
| 56 | FME | 2z | 1 | 56 | 8,9,10 | 0.98 | 0 | 8,9,11 | 1.05 | 0 |

| Mol | Type | Chain | Res | Link | Bond lengths | | | Bond angles | | |
|-----|------|-------|------|-------|--------------|------|----------|-------------|------|----------|
| | | | | | Counts | RMSZ | # Z > 2 | Counts | RMSZ | # Z > 2 |
| 54 | PSU | 1y | 55 | 54 | 18,21,22 | 1.35 | 2 (11%) | 21,30,33 | 2.06 | 3 (14%) |
| 54 | PSU | 2y | 39 | 54 | 18,21,22 | 1.36 | 2 (11%) | 21,30,33 | 1.73 | 4 (19%) |
| 55 | 4SU | 1x | 8 | 55 | 18,21,22 | 2.19 | 5 (27%) | 25,30,33 | 1.78 | 6 (24%) |
| 55 | 4SU | 2x | 8 | 55 | 18,21,22 | 2.23 | 6 (33%) | 25,30,33 | 1.77 | 5 (20%) |
| 54 | MIA | 2y | 37 | 54 | 17,24,32 | 1.00 | 1 (5%) | 16,35,47 | 1.29 | 2 (12%) |
| 54 | PSU | 2y | 32 | 54 | 18,21,22 | 1.38 | 2 (11%) | 21,30,33 | 1.97 | 4 (19%) |
| 1 | PSU | 1A | 1917 | 1 | 18,21,22 | 1.46 | 3 (16%) | 21,30,33 | 2.12 | 5 (23%) |
| 32 | MA6 | 2a | 1519 | 32 | 19,26,27 | 1.03 | 2 (10%) | 18,38,41 | 1.93 | 3 (16%) |
| 32 | 5MC | 1a | 1407 | 32 | 19,22,23 | 1.56 | 2 (10%) | 26,32,35 | 1.17 | 3 (11%) |
| 54 | 5MU | 2y | 54 | 54 | 19,22,23 | 1.51 | 5 (26%) | 27,32,35 | 1.77 | 7 (25%) |
| 1 | 5MC | 2A | 1942 | 1 | 19,22,23 | 1.70 | 2 (10%) | 26,32,35 | 1.14 | 2 (7%) |
| 1 | PSU | 2A | 1917 | 1 | 18,21,22 | 1.42 | 3 (16%) | 21,30,33 | 2.09 | 4 (19%) |
| 54 | PSU | 2w | 39 | 54 | 18,21,22 | 1.38 | 2 (11%) | 21,30,33 | 1.57 | 3 (14%) |
| 54 | MIA | 2w | 37 | 53,54 | 19,27,32 | 2.05 | 2 (10%) | 18,39,47 | 2.72 | 4 (22%) |
| 32 | 5MC | 1a | 967 | 32 | 19,22,23 | 1.66 | 3 (15%) | 26,32,35 | 1.12 | 2 (7%) |
| 54 | 4SU | 1y | 8 | 54 | 18,21,22 | 1.77 | 6 (33%) | 25,30,33 | 2.08 | 4 (16%) |
| 54 | PSU | 1w | 32 | 54 | 18,21,22 | 1.32 | 2 (11%) | 21,30,33 | 1.93 | 3 (14%) |
| 55 | 5MC | 1x | 32 | 55 | 19,22,23 | 1.63 | 3 (15%) | 26,32,35 | 1.17 | 2 (7%) |
| 32 | 2MG | 1a | 1207 | 32 | 18,26,27 | 0.99 | 1 (5%) | 16,38,41 | 1.54 | 5 (31%) |
| 54 | PSU | 1y | 32 | 54 | 18,21,22 | 1.34 | 2 (11%) | 21,30,33 | 2.00 | 4 (19%) |

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 |
|-----|------|-------|------|-------|---------|-----------|---------|
| 1 | 5MU | 2A | 1915 | 1,57 | - | 0/7/25/26 | 0/2/2/2 |
| 32 | PSU | 2a | 516 | 32,57 | - | 0/7/25/26 | 0/2/2/2 |
| 55 | PSU | 2x | 55 | 55 | - | 0/7/25/26 | 0/2/2/2 |
| 54 | PSU | 2w | 55 | 54 | - | 0/7/25/26 | 0/2/2/2 |
| 32 | 5MC | 2a | 1407 | 32 | - | 0/7/25/26 | 0/2/2/2 |
| 54 | G7M | 1w | 46 | 54 | - | 2/3/25/26 | 0/3/3/3 |
| 32 | PSU | 1a | 516 | 32,57 | - | 0/7/25/26 | 0/2/2/2 |
| 1 | OMU | 2A | 2552 | 1,57 | - | 0/9/27/28 | 0/2/2/2 |
| 54 | 4SU | 2y | 8 | 54 | - | 0/7/25/26 | 0/2/2/2 |
| 32 | 4OC | 2a | 1402 | 32,57 | - | 1/9/29/30 | 0/2/2/2 |
| 54 | 5MU | 2w | 54 | 54 | - | 0/7/25/26 | 0/2/2/2 |

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| Mol | Type | Chain | Res | Link | Chirals | Torsions | Rings |
|-----|------|-------|------|---------|---------|------------|---------|
| 1 | PSU | 2A | 1911 | 1 | - | 0/7/25/26 | 0/2/2/2 |
| 32 | 4OC | 1a | 1402 | 32 | - | 1/9/29/30 | 0/2/2/2 |
| 32 | MA6 | 2a | 1518 | 32 | - | 0/7/29/30 | 0/3/3/3 |
| 54 | PSU | 2w | 32 | 54 | - | 0/7/25/26 | 0/2/2/2 |
| 32 | 5MC | 2a | 967 | 32 | - | 0/7/25/26 | 0/2/2/2 |
| 56 | FME | 1z | 1 | 56 | - | 4/7/9/11 | - |
| 32 | 2MG | 2a | 1207 | 32,57 | - | 0/5/27/28 | 0/3/3/3 |
| 54 | 5MU | 1w | 54 | 54 | - | 0/7/25/26 | 0/2/2/2 |
| 1 | 2MA | 1A | 2503 | 1,57 | - | 1/3/25/26 | 0/3/3/3 |
| 54 | G7M | 2w | 46 | 54 | - | 1/3/25/26 | 0/3/3/3 |
| 54 | PSU | 2y | 55 | 54 | - | 5/7/25/26 | 0/2/2/2 |
| 1 | 5MC | 1A | 1942 | 1 | - | 0/7/25/26 | 0/2/2/2 |
| 54 | G7M | 2y | 46 | 54 | - | 0/3/25/26 | 0/3/3/3 |
| 54 | 4SU | 2w | 8 | 54 | - | 0/7/25/26 | 0/2/2/2 |
| 32 | MA6 | 1a | 1518 | 32 | - | 0/7/29/30 | 0/3/3/3 |
| 32 | 5MC | 2a | 1400 | 32 | - | 2/7/25/26 | 0/2/2/2 |
| 1 | 5MU | 1A | 1939 | 1,57 | - | 0/7/25/26 | 0/2/2/2 |
| 1 | OMG | 2A | 2251 | 55,1,57 | - | 0/5/27/28 | 0/3/3/3 |
| 1 | 2MA | 2A | 2503 | 1,57 | - | 1/3/25/26 | 0/3/3/3 |
| 32 | 5MC | 1a | 1400 | 32 | - | 2/7/25/26 | 0/2/2/2 |
| 32 | G7M | 1a | 527 | 32,57 | - | 2/3/25/26 | 0/3/3/3 |
| 54 | MIA | 1w | 37 | 54 | - | 1/11/33/34 | 0/3/3/3 |
| 55 | 5MC | 2x | 32 | 55 | - | 0/7/25/26 | 0/2/2/2 |
| 1 | 5MC | 2A | 1962 | 1,57 | - | 0/7/25/26 | 0/2/2/2 |
| 1 | 5MC | 1A | 1962 | 1,57 | - | 0/7/25/26 | 0/2/2/2 |
| 1 | PSU | 2A | 2605 | 1 | - | 0/7/25/26 | 0/2/2/2 |
| 32 | 5MC | 1a | 1404 | 32 | - | 0/7/25/26 | 0/2/2/2 |
| 55 | PSU | 1x | 55 | 55 | - | 0/7/25/26 | 0/2/2/2 |
| 1 | OMC | 2A | 1920 | 1 | - | 1/9/27/28 | 0/2/2/2 |
| 32 | UR3 | 1a | 1498 | 32 | - | 0/7/25/26 | 0/2/2/2 |
| 54 | PSU | 1y | 39 | 54 | - | 0/7/25/26 | 0/2/2/2 |
| 32 | M2G | 2a | 966 | 32 | - | 0/7/29/30 | 0/3/3/3 |
| 55 | 5MU | 2x | 54 | 55 | - | 0/7/25/26 | 0/2/2/2 |
| 43 | 0TD | 1l | 92 | 43 | - | 2/7/12/14 | - |
| 55 | 8AN | 1x | 76 | 55,57 | - | 3/3/25/26 | 0/3/3/3 |
| 32 | MA6 | 1a | 1519 | 32 | - | 2/7/29/30 | 0/3/3/3 |
| 32 | 5MC | 2a | 1404 | 32 | - | 0/7/25/26 | 0/2/2/2 |
| 1 | OMU | 1A | 2552 | 1,57 | - | 0/9/27/28 | 0/2/2/2 |
| 54 | PSU | 1w | 39 | 54 | - | 0/7/25/26 | 0/2/2/2 |
| 54 | G7M | 1y | 46 | 54 | - | 2/3/25/26 | 0/3/3/3 |

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| Mol | Type | Chain | Res | Link | Chirals | Torsions | Rings |
|-----|------|-------|------|----------|---------|-----------|---------|
| 54 | 5MU | 1y | 54 | 54 | - | 0/7/25/26 | 0/2/2/2 |
| 32 | G7M | 2a | 527 | 32 | - | 3/3/25/26 | 0/3/3/3 |
| 43 | 0TD | 2l | 92 | 43 | - | 2/7/12/14 | - |
| 32 | M2G | 1a | 966 | 32 | - | 0/7/29/30 | 0/3/3/3 |
| 55 | 8AN | 2x | 76 | 58,55,57 | - | 3/3/25/26 | 0/3/3/3 |
| 54 | MIA | 1y | 37 | 54 | - | 0/3/25/34 | 0/3/3/3 |
| 1 | PSU | 1A | 1911 | 1 | - | 0/7/25/26 | 0/2/2/2 |
| 1 | 5MU | 1A | 1915 | 1 | - | 0/7/25/26 | 0/2/2/2 |
| 55 | 5MU | 1x | 54 | 55 | - | 0/7/25/26 | 0/2/2/2 |
| 54 | 4SU | 1w | 8 | 54 | - | 0/7/25/26 | 0/2/2/2 |
| 1 | 5MU | 2A | 1939 | 1 | - | 0/7/25/26 | 0/2/2/2 |
| 1 | PSU | 1A | 2605 | 1,57 | - | 0/7/25/26 | 0/2/2/2 |
| 32 | UR3 | 2a | 1498 | 32,57 | - | 0/7/25/26 | 0/2/2/2 |
| 1 | OMC | 1A | 1920 | 1 | - | 2/9/27/28 | 0/2/2/2 |
| 54 | PSU | 1w | 55 | 54 | - | 0/7/25/26 | 0/2/2/2 |
| 1 | OMG | 1A | 2251 | 55,1,57 | - | 0/5/27/28 | 0/3/3/3 |
| 56 | FME | 2z | 1 | 56 | - | 1/7/9/11 | - |
| 54 | PSU | 1y | 55 | 54 | - | 0/7/25/26 | 0/2/2/2 |
| 54 | PSU | 2y | 39 | 54 | - | 0/7/25/26 | 0/2/2/2 |
| 55 | 4SU | 1x | 8 | 55 | - | 0/7/25/26 | 0/2/2/2 |
| 55 | 4SU | 2x | 8 | 55 | - | 0/7/25/26 | 0/2/2/2 |
| 54 | MIA | 2y | 37 | 54 | - | 0/3/25/34 | 0/3/3/3 |
| 54 | PSU | 2y | 32 | 54 | - | 0/7/25/26 | 0/2/2/2 |
| 1 | PSU | 1A | 1917 | 1 | - | 0/7/25/26 | 0/2/2/2 |
| 32 | MA6 | 2a | 1519 | 32 | - | 3/7/29/30 | 0/3/3/3 |
| 32 | 5MC | 1a | 1407 | 32 | - | 0/7/25/26 | 0/2/2/2 |
| 54 | 5MU | 2y | 54 | 54 | - | 0/7/25/26 | 0/2/2/2 |
| 1 | 5MC | 2A | 1942 | 1 | - | 0/7/25/26 | 0/2/2/2 |
| 1 | PSU | 2A | 1917 | 1 | - | 1/7/25/26 | 0/2/2/2 |
| 54 | PSU | 2w | 39 | 54 | - | 0/7/25/26 | 0/2/2/2 |
| 54 | MIA | 2w | 37 | 53,54 | - | 4/7/29/34 | 0/3/3/3 |
| 32 | 5MC | 1a | 967 | 32 | - | 1/7/25/26 | 0/2/2/2 |
| 54 | 4SU | 1y | 8 | 54 | - | 0/7/25/26 | 0/2/2/2 |
| 54 | PSU | 1w | 32 | 54 | - | 0/7/25/26 | 0/2/2/2 |
| 55 | 5MC | 1x | 32 | 55 | - | 0/7/25/26 | 0/2/2/2 |
| 32 | 2MG | 1a | 1207 | 32 | - | 0/5/27/28 | 0/3/3/3 |
| 54 | PSU | 1y | 32 | 54 | - | 0/7/25/26 | 0/2/2/2 |

The worst 5 of 222 bond length outliers are listed below:

| Mol | Chain | Res | Type | Atoms | Z | Observed(Å) | Ideal(Å) |
|-----|-------|------|------|---------|--------|-------------|----------|
| 43 | 2l | 92 | 0TD | CB-SB | -12.35 | 1.69 | 1.82 |
| 43 | 1l | 92 | 0TD | CB-SB | -12.14 | 1.70 | 1.82 |
| 54 | 2w | 37 | MIA | C2-S10 | -7.90 | 1.69 | 1.75 |
| 54 | 1w | 37 | MIA | C13-C14 | 7.07 | 1.53 | 1.32 |
| 32 | 2a | 1404 | 5MC | C5-C4 | 6.38 | 1.48 | 1.44 |

The worst 5 of 293 bond angle outliers are listed below:

| Mol | Chain | Res | Type | Atoms | Z | Observed(°) | Ideal(°) |
|-----|-------|-----|------|-------------|--------|-------------|----------|
| 54 | 2w | 37 | MIA | C11-S10-C2 | -10.36 | 94.48 | 102.25 |
| 55 | 2x | 76 | 8AN | O4'-C1'-N9 | -10.28 | 95.12 | 108.75 |
| 43 | 1l | 92 | 0TD | CSB-SB-CB | 10.24 | 120.77 | 102.36 |
| 43 | 2l | 92 | 0TD | CSB-SB-CB | -9.64 | 85.04 | 102.36 |
| 55 | 1x | 76 | 8AN | C4'-O4'-C1' | -8.79 | 101.88 | 109.92 |

There are no chirality outliers.

5 of 53 torsion outliers are listed below:

| Mol | Chain | Res | Type | Atoms |
|-----|-------|------|------|-----------------|
| 32 | 1a | 1400 | 5MC | O4'-C4'-C5'-O5' |
| 54 | 1w | 37 | MIA | C12-C13-C14-C15 |
| 54 | 1y | 46 | G7M | C4'-C5'-O5'-P |
| 32 | 2a | 1519 | MA6 | O4'-C4'-C5'-O5' |
| 54 | 2w | 37 | MIA | C5-C6-N6-C12 |

There are no ring outliers.

45 monomers are involved in 64 short contacts:

| Mol | Chain | Res | Type | Clashes | Symm-Clashes |
|-----|-------|------|------|---------|--------------|
| 1 | 2A | 1915 | 5MU | 1 | 0 |
| 55 | 2x | 55 | PSU | 2 | 0 |
| 54 | 1w | 46 | G7M | 1 | 0 |
| 1 | 2A | 2552 | OMU | 1 | 0 |
| 32 | 2a | 1402 | 4OC | 4 | 0 |
| 32 | 1a | 1402 | 4OC | 2 | 0 |
| 32 | 2a | 1518 | MA6 | 1 | 0 |
| 32 | 2a | 967 | 5MC | 3 | 0 |
| 54 | 2w | 46 | G7M | 1 | 0 |
| 54 | 2y | 55 | PSU | 6 | 0 |
| 1 | 1A | 1942 | 5MC | 1 | 0 |
| 54 | 2y | 46 | G7M | 1 | 0 |
| 32 | 1a | 1518 | MA6 | 2 | 0 |

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| Mol | Chain | Res | Type | Clashes | Symm-Clashes |
|-----|-------|------|------|---------|--------------|
| 32 | 2a | 1400 | 5MC | 2 | 0 |
| 1 | 1A | 1939 | 5MU | 1 | 0 |
| 1 | 2A | 2251 | OMG | 1 | 0 |
| 1 | 2A | 2503 | 2MA | 1 | 0 |
| 32 | 1a | 1400 | 5MC | 1 | 0 |
| 54 | 1w | 37 | MIA | 2 | 0 |
| 1 | 2A | 2605 | PSU | 1 | 0 |
| 1 | 2A | 1920 | OMC | 1 | 0 |
| 54 | 1y | 39 | PSU | 2 | 0 |
| 32 | 2a | 966 | M2G | 2 | 0 |
| 43 | 1l | 92 | 0TD | 1 | 0 |
| 55 | 1x | 76 | 8AN | 2 | 0 |
| 32 | 1a | 1519 | MA6 | 2 | 0 |
| 32 | 2a | 1404 | 5MC | 1 | 0 |
| 1 | 1A | 2552 | OMU | 2 | 0 |
| 54 | 1y | 46 | G7M | 1 | 0 |
| 43 | 2l | 92 | 0TD | 1 | 0 |
| 32 | 1a | 966 | M2G | 1 | 0 |
| 55 | 2x | 76 | 8AN | 2 | 0 |
| 54 | 1y | 37 | MIA | 1 | 0 |
| 1 | 1A | 1915 | 5MU | 1 | 0 |
| 54 | 1w | 8 | 4SU | 1 | 0 |
| 1 | 2A | 1939 | 5MU | 1 | 0 |
| 54 | 1y | 55 | PSU | 1 | 0 |
| 55 | 1x | 8 | 4SU | 2 | 0 |
| 54 | 2y | 37 | MIA | 1 | 0 |
| 1 | 1A | 1917 | PSU | 1 | 0 |
| 32 | 2a | 1519 | MA6 | 3 | 0 |
| 54 | 2w | 39 | PSU | 1 | 0 |
| 54 | 2w | 37 | MIA | 2 | 0 |
| 32 | 1a | 967 | 5MC | 1 | 0 |
| 54 | 1y | 8 | 4SU | 2 | 0 |

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

Of 2731 ligands modelled in this entry, 2727 are monoatomic - leaving 4 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 | $\# Z > 2$ |
| 61 | SF4 | 2d | 303 | 35 | 0,12,12 | - | - | - | | |
| 59 | ERY | 2A | 3857 | - | 53,53,53 | 0.93 | 1 (1%) | 82,82,82 | 1.49 | 15 (18%) |
| 59 | ERY | 1A | 4087 | - | 53,53,53 | 0.95 | 1 (1%) | 82,82,82 | 1.72 | 17 (20%) |
| 61 | SF4 | 1d | 302 | 35 | 0,12,12 | - | - | - | | |

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

| Mol | Type | Chain | Res | Link | Chirals | Torsions | Rings |
|-----|------|-------|------|------|---------|---------------|---------|
| 61 | SF4 | 2d | 303 | 35 | - | - | 0/6/5/5 |
| 59 | ERY | 2A | 3857 | - | - | 19/72/107/107 | 0/3/3/3 |
| 59 | ERY | 1A | 4087 | - | - | 15/72/107/107 | 0/3/3/3 |
| 61 | SF4 | 1d | 302 | 35 | - | - | 0/6/5/5 |

All (2) bond length outliers are listed below:

| Mol | Chain | Res | Type | Atoms | Z | Observed(Å) | Ideal(Å) |
|-----|-------|------|------|-------|------|-------------|----------|
| 59 | 1A | 4087 | ERY | O2-C1 | 5.17 | 1.46 | 1.34 |
| 59 | 2A | 3857 | ERY | O2-C1 | 4.97 | 1.45 | 1.34 |

The worst 5 of 32 bond angle outliers are listed below:

| Mol | Chain | Res | Type | Atoms | Z | Observed(°) | Ideal(°) |
|-----|-------|------|------|-------------|-------|-------------|----------|
| 59 | 1A | 4087 | ERY | C15-C16-C17 | 4.64 | 115.58 | 107.64 |
| 59 | 1A | 4087 | ERY | O5-C16-C15 | -4.23 | 106.44 | 112.95 |
| 59 | 1A | 4087 | ERY | O7-C5-C4 | -4.20 | 105.50 | 111.58 |
| 59 | 1A | 4087 | ERY | C6-C5-C4 | -4.19 | 107.81 | 113.89 |
| 59 | 2A | 3857 | ERY | O5-C16-C15 | -3.98 | 106.83 | 112.95 |

There are no chirality outliers.

5 of 34 torsion outliers are listed below:

| Mol | Chain | Res | Type | Atoms |
|-----|-------|------|------|-----------------|
| 59 | 1A | 4087 | ERY | C17-C16-O5-C20 |
| 59 | 1A | 4087 | ERY | C19-C16-O5-C20 |
| 59 | 2A | 3857 | ERY | C9-C10-C11-C12 |
| 59 | 2A | 3857 | ERY | C9-C10-C11-O12 |
| 59 | 2A | 3857 | ERY | C34-C10-C11-C12 |

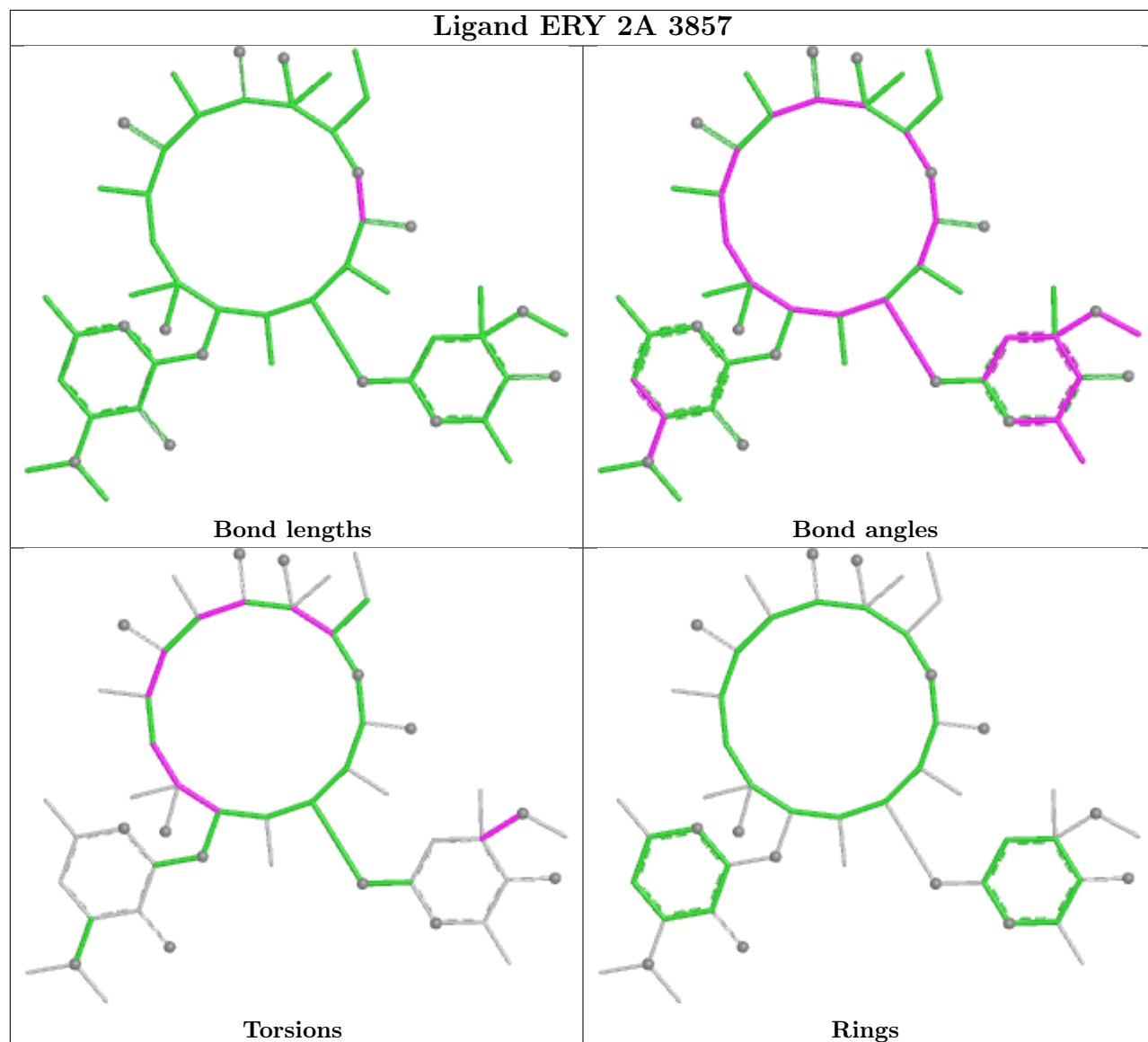
There are no ring outliers.

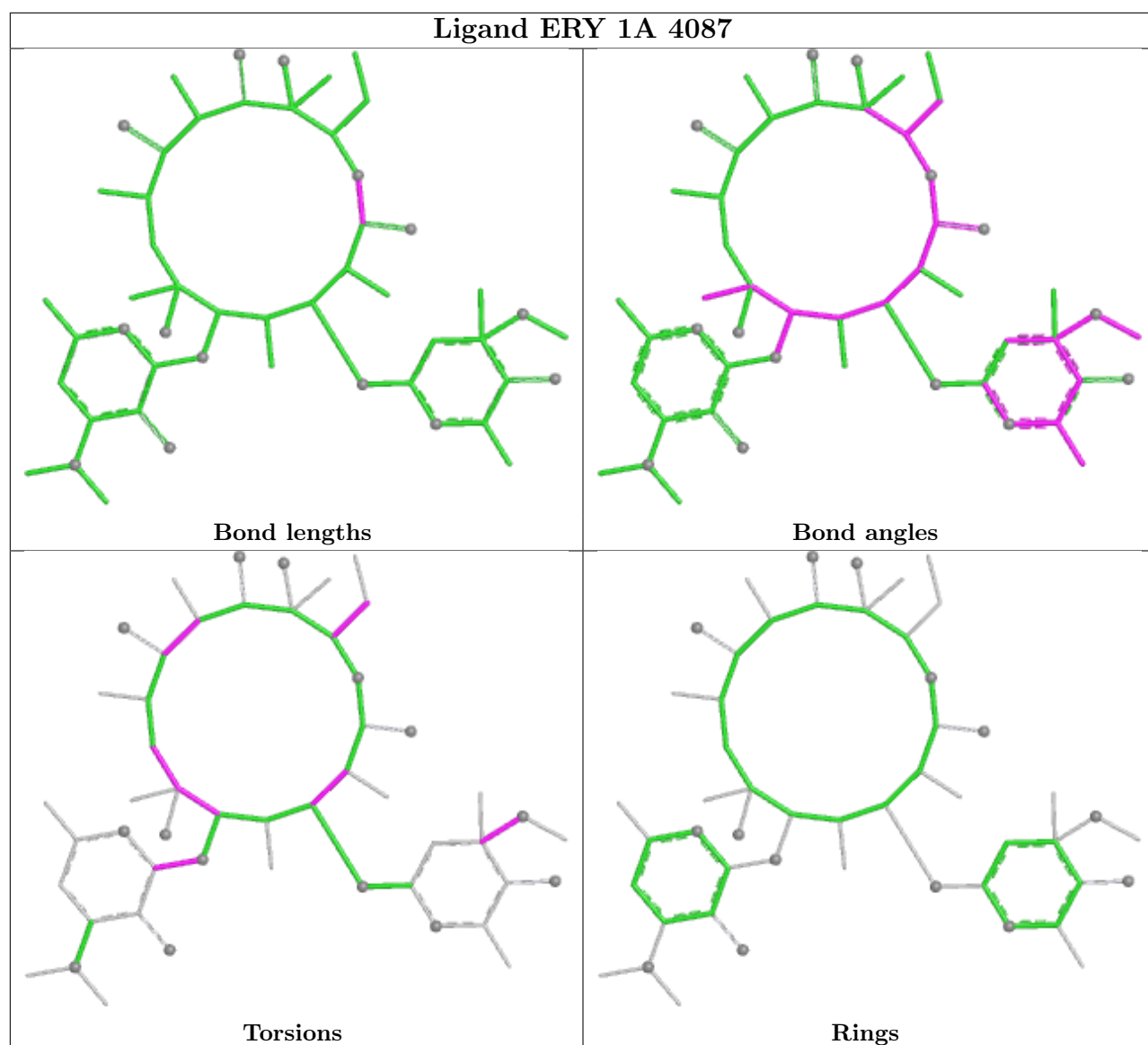
4 monomers are involved in 16 short contacts:

| Mol | Chain | Res | Type | Clashes | Symm-Clashes |
|-----|-------|------|------|---------|--------------|
| 61 | 2d | 303 | SF4 | 2 | 0 |
| 59 | 2A | 3857 | ERY | 4 | 0 |
| 59 | 1A | 4087 | ERY | 9 | 0 |
| 61 | 1d | 302 | SF4 | 1 | 0 |

The following is a two-dimensional graphical depiction of Mogul quality analysis of bond lengths, bond angles, torsion angles, and ring geometry for all instances of the Ligand of Interest. In addition, ligands with molecular weight > 250 and outliers as shown on the validation Tables will also be included. For torsion angles, if less than 5% of the Mogul distribution of torsion angles is within 10 degrees of the torsion angle in question, then that torsion angle is considered an outlier. Any bond that is central to one or more torsion angles identified as an outlier by Mogul will be highlighted in the graph. For rings, the root-mean-square deviation (RMSD) between the ring in question and similar rings identified by Mogul is calculated over all ring torsion angles. If the average RMSD is greater than 60 degrees and the minimal RMSD between the ring in question and any Mogul-identified rings is also greater than 60 degrees, then that ring is considered an outlier. The outliers are highlighted in purple. The color gray indicates Mogul did not find sufficient equivalents in the CSD to analyse the geometry.

Ligand ERY 2A 3857





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, 95th percentile and maximum values of the occupancy-weighted average B-factor per residue. The column labelled ‘Q< 0.9’ lists the number of (and percentage) of residues with an average occupancy less than 0.9.

| Mol | Chain | Analysed | <RSRZ> | #RSRZ>2 | OWAB(Å ²) | Q<0.9 |
|-----|-------|-----------------|--------|---------------|-----------------------|-------|
| 1 | 1A | 2860/2915 (98%) | -0.60 | 52 (1%) 67 68 | 19, 37, 92, 108 | 0 |
| 1 | 2A | 2789/2915 (95%) | -0.19 | 44 (1%) 70 71 | 32, 56, 91, 105 | 0 |
| 2 | 1B | 120/121 (99%) | -0.54 | 0 100 100 | 31, 50, 64, 83 | 0 |
| 2 | 2B | 120/121 (99%) | 0.50 | 2 (1%) 69 70 | 57, 77, 86, 91 | 0 |
| 3 | 1D | 275/276 (99%) | -0.28 | 2 (0%) 84 85 | 19, 36, 51, 71 | 0 |
| 3 | 2D | 275/276 (99%) | 0.14 | 4 (1%) 71 73 | 29, 50, 62, 75 | 0 |
| 4 | 1E | 204/206 (99%) | -0.14 | 1 (0%) 87 89 | 20, 44, 63, 79 | 0 |
| 4 | 2E | 204/206 (99%) | 0.08 | 1 (0%) 87 89 | 29, 56, 69, 81 | 0 |
| 5 | 1F | 202/210 (96%) | -0.26 | 1 (0%) 87 89 | 19, 43, 65, 84 | 0 |
| 5 | 2F | 202/210 (96%) | 0.22 | 4 (1%) 64 65 | 34, 67, 80, 85 | 0 |
| 6 | 1G | 181/182 (99%) | 0.21 | 2 (1%) 77 80 | 39, 57, 74, 88 | 0 |
| 6 | 2G | 181/182 (99%) | 0.93 | 15 (8%) 19 22 | 66, 78, 86, 89 | 0 |
| 7 | 1H | 174/180 (96%) | 0.04 | 0 100 100 | 39, 54, 67, 75 | 0 |
| 7 | 2H | 174/180 (96%) | 0.80 | 9 (5%) 34 35 | 62, 80, 87, 95 | 0 |
| 8 | 1I | 146/148 (98%) | 0.35 | 3 (2%) 63 64 | 39, 70, 80, 84 | 0 |
| 8 | 2I | 146/148 (98%) | 0.81 | 9 (6%) 28 30 | 57, 74, 84, 87 | 0 |
| 9 | 1N | 140/140 (100%) | -0.12 | 0 100 100 | 28, 45, 61, 79 | 0 |
| 9 | 2N | 140/140 (100%) | 0.51 | 4 (2%) 54 55 | 44, 63, 75, 83 | 0 |
| 10 | 1O | 122/122 (100%) | -0.25 | 0 100 100 | 28, 42, 60, 67 | 0 |
| 10 | 2O | 122/122 (100%) | 0.23 | 1 (0%) 82 84 | 37, 55, 68, 72 | 0 |
| 11 | 1P | 149/150 (99%) | -0.05 | 0 100 100 | 21, 43, 67, 75 | 0 |
| 11 | 2P | 149/150 (99%) | 0.29 | 3 (2%) 64 65 | 36, 66, 82, 87 | 0 |
| 12 | 1Q | 141/141 (100%) | -0.19 | 0 100 100 | 27, 42, 55, 73 | 0 |
| 12 | 2Q | 141/141 (100%) | 0.67 | 8 (5%) 30 33 | 46, 64, 74, 80 | 0 |

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| Mol | Chain | Analysed | <RSRZ> | #RSRZ>2 | OWAB(Å ²) | Q<0.9 |
|-----|-------|----------------|--------|---------------|-----------------------|-------|
| 13 | 1R | 118/118 (100%) | -0.24 | 0 100 100 | 29, 38, 51, 60 | 0 |
| 13 | 2R | 118/118 (100%) | -0.02 | 0 100 100 | 37, 50, 60, 69 | 0 |
| 14 | 1S | 110/112 (98%) | 0.00 | 0 100 100 | 38, 49, 60, 65 | 0 |
| 14 | 2S | 110/112 (98%) | 0.87 | 9 (8%) 19 22 | 60, 70, 78, 82 | 0 |
| 15 | 1T | 131/146 (89%) | 0.03 | 3 (2%) 61 61 | 35, 49, 71, 80 | 0 |
| 15 | 2T | 131/146 (89%) | 0.28 | 1 (0%) 82 84 | 45, 57, 73, 83 | 0 |
| 16 | 1U | 116/118 (98%) | -0.20 | 2 (1%) 69 70 | 24, 37, 55, 73 | 0 |
| 16 | 2U | 116/118 (98%) | 0.19 | 1 (0%) 81 83 | 43, 58, 73, 78 | 0 |
| 17 | 1V | 101/101 (100%) | -0.12 | 0 100 100 | 23, 48, 62, 72 | 0 |
| 17 | 2V | 101/101 (100%) | 0.46 | 1 (0%) 79 82 | 46, 70, 78, 82 | 0 |
| 18 | 1W | 112/113 (99%) | -0.35 | 1 (0%) 81 83 | 27, 38, 56, 76 | 0 |
| 18 | 2W | 112/113 (99%) | 0.21 | 0 100 100 | 38, 50, 67, 93 | 0 |
| 19 | 1X | 95/96 (98%) | -0.28 | 0 100 100 | 28, 38, 55, 71 | 0 |
| 19 | 2X | 95/96 (98%) | 0.38 | 1 (1%) 77 80 | 41, 58, 71, 76 | 0 |
| 20 | 1Y | 107/110 (97%) | 0.14 | 1 (0%) 81 83 | 35, 50, 68, 75 | 0 |
| 20 | 2Y | 107/110 (97%) | 0.79 | 4 (3%) 45 47 | 60, 72, 80, 86 | 0 |
| 21 | 1Z | 154/206 (74%) | 0.16 | 5 (3%) 50 52 | 42, 63, 82, 90 | 0 |
| 21 | 2Z | 160/206 (77%) | 0.97 | 21 (13%) 8 11 | 67, 79, 89, 99 | 0 |
| 22 | 10 | 83/85 (97%) | -0.13 | 2 (2%) 59 60 | 28, 37, 56, 64 | 0 |
| 22 | 20 | 83/85 (97%) | 0.85 | 7 (8%) 18 21 | 47, 61, 70, 81 | 0 |
| 23 | 11 | 97/98 (98%) | 0.01 | 1 (1%) 79 82 | 23, 43, 65, 70 | 0 |
| 23 | 21 | 97/98 (98%) | 0.48 | 3 (3%) 51 53 | 38, 56, 73, 76 | 0 |
| 24 | 12 | 70/72 (97%) | 0.02 | 1 (1%) 73 75 | 34, 47, 57, 71 | 0 |
| 24 | 22 | 70/72 (97%) | 0.66 | 3 (4%) 40 42 | 58, 68, 75, 77 | 0 |
| 25 | 13 | 59/60 (98%) | -0.23 | 0 100 100 | 29, 41, 65, 78 | 0 |
| 25 | 23 | 59/60 (98%) | 0.26 | 0 100 100 | 52, 65, 78, 83 | 0 |
| 26 | 14 | 69/71 (97%) | 0.42 | 3 (4%) 40 42 | 50, 73, 87, 93 | 0 |
| 26 | 24 | 69/71 (97%) | 0.81 | 7 (10%) 14 16 | 75, 85, 93, 96 | 0 |
| 27 | 15 | 59/60 (98%) | -0.35 | 0 100 100 | 21, 42, 64, 74 | 0 |
| 27 | 25 | 59/60 (98%) | 0.04 | 1 (1%) 69 70 | 34, 51, 70, 80 | 0 |
| 28 | 16 | 53/54 (98%) | -0.15 | 0 100 100 | 31, 40, 53, 59 | 0 |

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| Mol | Chain | Analysed | <RSRZ> | #RSRZ>2 | OWAB(Å ²) | Q<0.9 |
|-----|-------|-----------------|--------|----------------|-----------------------|-------|
| 28 | 26 | 53/54 (98%) | 0.49 | 0 100 100 | 49, 59, 66, 73 | 0 |
| 29 | 17 | 48/49 (97%) | -0.55 | 0 100 100 | 21, 27, 46, 54 | 0 |
| 29 | 27 | 48/49 (97%) | -0.07 | 1 (2%) 63 64 | 32, 41, 60, 67 | 0 |
| 30 | 18 | 64/65 (98%) | -0.22 | 1 (1%) 70 71 | 28, 33, 42, 55 | 0 |
| 30 | 28 | 64/65 (98%) | 0.38 | 1 (1%) 70 71 | 42, 54, 64, 71 | 0 |
| 31 | 19 | 37/37 (100%) | 0.02 | 0 100 100 | 34, 42, 55, 64 | 0 |
| 31 | 29 | 37/37 (100%) | 0.92 | 3 (8%) 19 22 | 57, 66, 76, 78 | 0 |
| 32 | 1a | 1488/1521 (97%) | -0.05 | 14 (0%) 81 83 | 34, 63, 90, 107 | 0 |
| 32 | 2a | 1491/1521 (98%) | 0.36 | 49 (3%) 49 51 | 51, 76, 94, 109 | 0 |
| 33 | 1b | 231/256 (90%) | 0.83 | 26 (11%) 11 13 | 61, 76, 84, 94 | 0 |
| 33 | 2b | 231/256 (90%) | 1.18 | 37 (16%) 6 7 | 72, 84, 90, 93 | 0 |
| 34 | 1c | 206/239 (86%) | 0.42 | 4 (1%) 66 67 | 54, 66, 77, 87 | 0 |
| 34 | 2c | 206/239 (86%) | 1.12 | 32 (15%) 6 7 | 68, 83, 88, 93 | 0 |
| 35 | 1d | 208/209 (99%) | 0.33 | 2 (0%) 79 82 | 51, 64, 75, 81 | 0 |
| 35 | 2d | 208/209 (99%) | 0.81 | 14 (6%) 25 28 | 63, 72, 81, 86 | 0 |
| 36 | 1e | 148/162 (91%) | 0.24 | 1 (0%) 84 85 | 47, 61, 73, 86 | 0 |
| 36 | 2e | 148/162 (91%) | 0.83 | 10 (6%) 25 28 | 64, 77, 83, 87 | 0 |
| 37 | 1f | 100/101 (99%) | 0.31 | 1 (1%) 79 82 | 47, 60, 75, 77 | 0 |
| 37 | 2f | 100/101 (99%) | 0.32 | 0 100 100 | 60, 69, 78, 86 | 0 |
| 38 | 1g | 155/156 (99%) | 0.46 | 10 (6%) 26 29 | 54, 67, 79, 86 | 0 |
| 38 | 2g | 155/156 (99%) | 0.76 | 13 (8%) 18 21 | 69, 79, 87, 94 | 0 |
| 39 | 1h | 137/138 (99%) | 0.33 | 3 (2%) 62 62 | 56, 65, 73, 78 | 0 |
| 39 | 2h | 137/138 (99%) | 0.94 | 10 (7%) 22 26 | 67, 77, 84, 89 | 0 |
| 40 | 1i | 127/128 (99%) | 0.68 | 3 (2%) 59 60 | 52, 72, 81, 87 | 0 |
| 40 | 2i | 127/128 (99%) | 1.68 | 43 (33%) 1 1 | 70, 83, 90, 93 | 0 |
| 41 | 1j | 97/105 (92%) | 0.70 | 3 (3%) 51 53 | 51, 73, 84, 91 | 0 |
| 41 | 2j | 96/105 (91%) | 1.32 | 19 (19%) 3 4 | 72, 83, 89, 93 | 0 |
| 42 | 1k | 114/129 (88%) | 0.16 | 1 (0%) 81 83 | 40, 63, 74, 82 | 0 |
| 42 | 2k | 114/129 (88%) | 0.97 | 14 (12%) 9 12 | 63, 74, 80, 85 | 0 |
| 43 | 1l | 121/132 (91%) | 0.03 | 4 (3%) 49 51 | 40, 51, 64, 71 | 0 |
| 43 | 2l | 121/132 (91%) | 0.43 | 3 (2%) 58 59 | 46, 65, 74, 81 | 0 |

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| Mol | Chain | Analysed | <RSRZ> | #RSRZ>2 | OWAB(Å ²) | Q<0.9 |
|-----|-------|-------------------|--------|----------------|-----------------------|-------|
| 44 | 1m | 123/126 (97%) | 0.41 | 6 (4%) 36 38 | 49, 65, 76, 93 | 0 |
| 44 | 2m | 122/126 (96%) | 1.11 | 18 (14%) 7 8 | 72, 81, 88, 91 | 0 |
| 45 | 1n | 60/61 (98%) | 0.54 | 1 (1%) 69 70 | 52, 61, 68, 69 | 0 |
| 45 | 2n | 60/61 (98%) | 1.97 | 28 (46%) 0 0 | 74, 82, 88, 89 | 0 |
| 46 | 1o | 88/89 (98%) | 0.46 | 3 (3%) 48 50 | 47, 60, 71, 76 | 0 |
| 46 | 2o | 88/89 (98%) | 0.67 | 2 (2%) 61 61 | 60, 72, 80, 83 | 0 |
| 47 | 1p | 82/88 (93%) | 0.81 | 4 (4%) 36 38 | 57, 66, 74, 79 | 0 |
| 47 | 2p | 82/88 (93%) | 0.74 | 3 (3%) 45 47 | 56, 65, 75, 77 | 0 |
| 48 | 1q | 99/105 (94%) | 0.87 | 4 (4%) 43 44 | 53, 66, 76, 81 | 0 |
| 48 | 2q | 99/105 (94%) | 0.82 | 6 (6%) 28 31 | 64, 72, 80, 86 | 0 |
| 49 | 1r | 68/88 (77%) | 0.25 | 1 (1%) 71 73 | 49, 63, 75, 77 | 0 |
| 49 | 2r | 68/88 (77%) | 0.39 | 0 100 100 | 63, 74, 83, 85 | 0 |
| 50 | 1s | 83/93 (89%) | 0.19 | 0 100 100 | 54, 66, 76, 81 | 0 |
| 50 | 2s | 83/93 (89%) | 1.29 | 15 (18%) 4 5 | 75, 83, 90, 93 | 0 |
| 51 | 1t | 96/106 (90%) | 0.62 | 5 (5%) 34 35 | 55, 66, 78, 89 | 0 |
| 51 | 2t | 96/106 (90%) | 0.61 | 2 (2%) 63 64 | 54, 65, 82, 85 | 0 |
| 52 | 1u | 23/27 (85%) | 0.73 | 1 (4%) 40 42 | 57, 64, 71, 72 | 0 |
| 52 | 2u | 23/27 (85%) | 1.72 | 9 (39%) 1 1 | 75, 78, 83, 87 | 0 |
| 53 | 1v | 13/24 (54%) | 0.19 | 1 (7%) 21 24 | 42, 57, 82, 91 | 0 |
| 53 | 2v | 13/24 (54%) | 1.47 | 3 (23%) 2 3 | 72, 79, 92, 94 | 0 |
| 54 | 1w | 67/76 (88%) | 0.65 | 5 (7%) 22 24 | 34, 90, 98, 109 | 0 |
| 54 | 1y | 67/76 (88%) | 0.40 | 2 (2%) 52 54 | 38, 90, 98, 102 | 0 |
| 54 | 2w | 65/76 (85%) | 0.76 | 6 (9%) 16 18 | 52, 92, 100, 103 | 0 |
| 54 | 2y | 66/76 (86%) | 0.71 | 3 (4%) 39 40 | 50, 96, 101, 104 | 0 |
| 55 | 1x | 72/77 (93%) | -0.02 | 0 100 100 | 29, 61, 79, 84 | 0 |
| 55 | 2x | 72/77 (93%) | 0.36 | 0 100 100 | 44, 78, 88, 91 | 0 |
| 56 | 1z | 2/3 (66%) | 0.01 | 0 100 100 | 35, 35, 35, 38 | 0 |
| 56 | 2z | 2/3 (66%) | 0.46 | 0 100 100 | 48, 48, 48, 52 | 0 |
| All | All | 20877/21754 (95%) | 0.15 | 693 (3%) 49 51 | 19, 62, 88, 109 | 0 |

The worst 5 of 693 RSRZ outliers are listed below:

| Mol | Chain | Res | Type | RSRZ |
|-----|-------|-----|------|------|
| 44 | 1m | 124 | PRO | 7.4 |
| 44 | 2m | 124 | PRO | 7.2 |
| 44 | 2m | 123 | ALA | 6.7 |
| 23 | 21 | 2 | SER | 6.7 |
| 44 | 2m | 122 | LYS | 6.6 |

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

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

| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(Å ²) | Q<0.9 |
|-----|------|-------|------|-------|------|------|----------------------------|-------|
| 54 | G7M | 2y | 46 | 24/25 | 0.57 | 0.15 | 90,98,107,119 | 0 |
| 54 | G7M | 1w | 46 | 24/25 | 0.63 | 0.12 | 83,92,104,119 | 0 |
| 54 | PSU | 1y | 55 | 20/21 | 0.66 | 0.13 | 87,97,107,116 | 0 |
| 54 | PSU | 2y | 55 | 20/21 | 0.66 | 0.14 | 84,95,107,111 | 0 |
| 54 | G7M | 1y | 46 | 24/25 | 0.67 | 0.12 | 87,93,102,111 | 0 |
| 54 | G7M | 2w | 46 | 24/25 | 0.69 | 0.12 | 83,93,106,120 | 0 |
| 54 | PSU | 2w | 55 | 20/21 | 0.72 | 0.11 | 79,89,96,103 | 0 |
| 54 | 4SU | 2y | 8 | 20/21 | 0.73 | 0.13 | 90,99,108,111 | 0 |
| 54 | MIA | 2y | 37 | 22/30 | 0.75 | 0.15 | 87,92,98,108 | 0 |
| 54 | 4SU | 2w | 8 | 20/21 | 0.75 | 0.11 | 89,95,108,109 | 0 |
| 54 | 5MU | 2y | 54 | 21/22 | 0.75 | 0.15 | 85,90,102,117 | 0 |
| 54 | 5MU | 1y | 54 | 21/22 | 0.75 | 0.14 | 82,93,102,114 | 0 |
| 54 | 4SU | 1w | 8 | 20/21 | 0.76 | 0.10 | 87,93,101,103 | 0 |
| 54 | 5MU | 2w | 54 | 21/22 | 0.76 | 0.12 | 73,81,85,96 | 0 |
| 54 | MIA | 1y | 37 | 22/30 | 0.78 | 0.11 | 69,83,100,110 | 0 |
| 54 | PSU | 2y | 32 | 20/21 | 0.79 | 0.14 | 86,90,97,103 | 0 |
| 54 | PSU | 2y | 39 | 20/21 | 0.81 | 0.14 | 84,90,98,102 | 0 |
| 54 | 4SU | 1y | 8 | 20/21 | 0.81 | 0.09 | 88,94,103,104 | 0 |
| 32 | 2MG | 2a | 1207 | 24/25 | 0.83 | 0.13 | 81,89,99,109 | 0 |
| 54 | PSU | 1w | 55 | 20/21 | 0.83 | 0.10 | 76,85,92,94 | 0 |
| 55 | 4SU | 2x | 8 | 20/21 | 0.86 | 0.10 | 74,78,81,86 | 0 |
| 55 | PSU | 2x | 55 | 20/21 | 0.87 | 0.10 | 70,77,88,94 | 0 |
| 54 | PSU | 1w | 32 | 20/21 | 0.87 | 0.13 | 61,74,77,78 | 0 |
| 54 | PSU | 1y | 32 | 20/21 | 0.88 | 0.09 | 74,80,91,97 | 0 |
| 32 | M2G | 2a | 966 | 25/26 | 0.88 | 0.16 | 63,74,87,94 | 0 |
| 54 | PSU | 2w | 39 | 20/21 | 0.88 | 0.12 | 73,83,91,93 | 0 |
| 32 | PSU | 2a | 516 | 20/21 | 0.89 | 0.09 | 63,73,85,88 | 0 |
| 54 | PSU | 1y | 39 | 20/21 | 0.89 | 0.08 | 74,81,85,87 | 0 |
| 54 | PSU | 2w | 32 | 20/21 | 0.89 | 0.12 | 73,83,89,90 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 32 | 5MC | 2a | 967 | 21/22 | 0.89 | 0.13 | 70,77,82,88 | 0 |
| 55 | 5MU | 2x | 54 | 21/22 | 0.90 | 0.10 | 79,82,86,93 | 0 |
| 54 | 5MU | 1w | 54 | 21/22 | 0.90 | 0.09 | 63,74,83,85 | 0 |
| 55 | 5MU | 1x | 54 | 21/22 | 0.90 | 0.10 | 59,67,72,75 | 0 |
| 54 | MIA | 1w | 37 | 29/30 | 0.91 | 0.15 | 49,63,78,88 | 0 |
| 1 | 5MU | 2A | 1915 | 21/22 | 0.91 | 0.09 | 66,71,76,82 | 0 |
| 55 | 5MC | 2x | 32 | 21/22 | 0.92 | 0.12 | 69,74,79,80 | 0 |
| 32 | G7M | 2a | 527 | 24/25 | 0.92 | 0.11 | 58,63,69,74 | 0 |
| 32 | 5MC | 2a | 1400 | 21/22 | 0.92 | 0.15 | 68,74,79,81 | 0 |
| 54 | MIA | 2w | 37 | 25/30 | 0.92 | 0.10 | 64,76,84,92 | 0 |
| 43 | 0TD | 2l | 92 | 10/11 | 0.92 | 0.12 | 61,63,71,79 | 0 |
| 1 | PSU | 2A | 1917 | 20/21 | 0.93 | 0.08 | 58,64,72,74 | 0 |
| 55 | PSU | 1x | 55 | 20/21 | 0.93 | 0.08 | 56,59,70,71 | 0 |
| 32 | 5MC | 2a | 1407 | 21/22 | 0.94 | 0.10 | 50,62,65,68 | 0 |
| 32 | UR3 | 2a | 1498 | 21/22 | 0.94 | 0.11 | 51,59,66,71 | 0 |
| 32 | MA6 | 2a | 1519 | 24/25 | 0.94 | 0.13 | 60,67,74,77 | 0 |
| 55 | 4SU | 1x | 8 | 20/21 | 0.94 | 0.09 | 51,57,64,77 | 0 |
| 32 | PSU | 1a | 516 | 20/21 | 0.94 | 0.10 | 52,57,61,63 | 0 |
| 1 | PSU | 2A | 1911 | 20/21 | 0.94 | 0.09 | 42,59,66,66 | 0 |
| 1 | 5MU | 1A | 1915 | 21/22 | 0.94 | 0.08 | 43,51,55,59 | 0 |
| 54 | PSU | 1w | 39 | 20/21 | 0.94 | 0.08 | 67,71,74,87 | 0 |
| 1 | 5MC | 2A | 1962 | 21/22 | 0.94 | 0.11 | 44,54,58,73 | 0 |
| 32 | 4OC | 2a | 1402 | 22/23 | 0.94 | 0.10 | 59,70,73,73 | 0 |
| 32 | 5MC | 2a | 1404 | 21/22 | 0.94 | 0.10 | 52,64,70,74 | 0 |
| 56 | FME | 1z | 1 | 10/11 | 0.94 | 0.15 | 37,49,52,63 | 0 |
| 1 | OMC | 2A | 1920 | 21/22 | 0.95 | 0.08 | 54,60,62,63 | 0 |
| 32 | 2MG | 1a | 1207 | 24/25 | 0.95 | 0.09 | 57,65,68,77 | 0 |
| 55 | 5MC | 1x | 32 | 21/22 | 0.95 | 0.09 | 38,45,52,57 | 0 |
| 56 | FME | 2z | 1 | 10/11 | 0.95 | 0.14 | 52,55,58,58 | 0 |
| 32 | 4OC | 1a | 1402 | 22/23 | 0.96 | 0.09 | 43,48,52,58 | 0 |
| 43 | 0TD | 1l | 92 | 10/11 | 0.96 | 0.08 | 34,44,50,51 | 0 |
| 1 | PSU | 1A | 1917 | 20/21 | 0.96 | 0.08 | 36,47,51,52 | 0 |
| 1 | 5MC | 2A | 1942 | 21/22 | 0.96 | 0.09 | 42,58,62,69 | 0 |
| 32 | M2G | 1a | 966 | 25/26 | 0.96 | 0.10 | 46,51,56,57 | 0 |
| 1 | OMU | 2A | 2552 | 21/22 | 0.96 | 0.09 | 35,39,48,49 | 0 |
| 32 | 5MC | 1a | 967 | 21/22 | 0.96 | 0.09 | 44,55,61,68 | 0 |
| 32 | MA6 | 2a | 1518 | 24/25 | 0.96 | 0.09 | 57,65,70,73 | 0 |
| 1 | 5MC | 1A | 1942 | 21/22 | 0.96 | 0.08 | 33,39,43,54 | 0 |
| 32 | 5MC | 1a | 1400 | 21/22 | 0.96 | 0.10 | 37,42,49,56 | 0 |
| 32 | 5MC | 1a | 1407 | 21/22 | 0.97 | 0.08 | 26,40,45,48 | 0 |
| 55 | 8AN | 1x | 76 | 22/23 | 0.97 | 0.08 | 25,33,41,43 | 0 |
| 55 | 8AN | 2x | 76 | 22/23 | 0.97 | 0.09 | 40,44,49,53 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 32 | MA6 | 1a | 1518 | 24/25 | 0.97 | 0.09 | 32,41,45,48 | 0 |
| 32 | MA6 | 1a | 1519 | 24/25 | 0.97 | 0.08 | 36,43,48,56 | 0 |
| 32 | G7M | 1a | 527 | 24/25 | 0.97 | 0.07 | 38,44,50,56 | 0 |
| 1 | 5MU | 2A | 1939 | 21/22 | 0.97 | 0.07 | 34,38,44,45 | 0 |
| 1 | 5MC | 1A | 1962 | 21/22 | 0.97 | 0.07 | 27,34,40,46 | 0 |
| 1 | OMC | 1A | 1920 | 21/22 | 0.97 | 0.07 | 34,40,46,50 | 0 |
| 1 | OMG | 2A | 2251 | 24/25 | 0.97 | 0.07 | 33,39,44,46 | 0 |
| 32 | 5MC | 1a | 1404 | 21/22 | 0.97 | 0.09 | 28,40,43,51 | 0 |
| 1 | PSU | 2A | 2605 | 20/21 | 0.97 | 0.07 | 30,38,46,47 | 0 |
| 1 | PSU | 1A | 1911 | 20/21 | 0.98 | 0.06 | 38,42,46,50 | 0 |
| 1 | 5MU | 1A | 1939 | 21/22 | 0.98 | 0.06 | 23,26,32,35 | 0 |
| 1 | OMG | 1A | 2251 | 24/25 | 0.98 | 0.07 | 17,26,28,30 | 0 |
| 1 | 2MA | 1A | 2503 | 23/24 | 0.98 | 0.05 | 20,22,26,26 | 0 |
| 32 | UR3 | 1a | 1498 | 21/22 | 0.98 | 0.07 | 36,41,46,51 | 0 |
| 1 | 2MA | 2A | 2503 | 23/24 | 0.98 | 0.06 | 28,34,36,39 | 0 |
| 1 | OMU | 1A | 2552 | 21/22 | 0.98 | 0.06 | 24,28,32,40 | 0 |
| 1 | PSU | 1A | 2605 | 20/21 | 0.99 | 0.06 | 18,27,31,34 | 0 |

6.3 Carbohydrates

There are no oligosaccharides in this entry.

6.4 Ligands

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

| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1a | 1720 | 1/1 | 0.51 | 0.38 | 82,82,82,82 | 0 |
| 57 | MG | 1B | 232 | 1/1 | 0.55 | 0.16 | 76,76,76,76 | 0 |
| 57 | MG | 2A | 3772 | 1/1 | 0.56 | 0.15 | 71,71,71,71 | 0 |
| 57 | MG | 2A | 3710 | 1/1 | 0.58 | 0.17 | 79,79,79,79 | 0 |
| 57 | MG | 2A | 3434 | 1/1 | 0.58 | 0.21 | 78,78,78,78 | 0 |
| 57 | MG | 2A | 3576 | 1/1 | 0.64 | 0.18 | 67,67,67,67 | 0 |
| 57 | MG | 1A | 3589 | 1/1 | 0.66 | 0.27 | 71,71,71,71 | 0 |
| 57 | MG | 2A | 3273 | 1/1 | 0.66 | 0.22 | 78,78,78,78 | 0 |
| 57 | MG | 2a | 1738 | 1/1 | 0.66 | 0.25 | 79,79,79,79 | 0 |
| 57 | MG | 2a | 1800 | 1/1 | 0.67 | 0.25 | 79,79,79,79 | 0 |
| 57 | MG | 2A | 3678 | 1/1 | 0.68 | 0.23 | 79,79,79,79 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3203 | 1/1 | 0.69 | 0.27 | 68,68,68,68 | 0 |
| 57 | MG | 2a | 1647 | 1/1 | 0.70 | 0.21 | 64,64,64,64 | 0 |
| 57 | MG | 1w | 104 | 1/1 | 0.70 | 0.23 | 81,81,81,81 | 0 |
| 57 | MG | 2A | 3236 | 1/1 | 0.70 | 0.18 | 67,67,67,67 | 0 |
| 57 | MG | 18 | 105 | 1/1 | 0.71 | 0.20 | 65,65,65,65 | 0 |
| 57 | MG | 2A | 3643 | 1/1 | 0.71 | 0.34 | 83,83,83,83 | 0 |
| 57 | MG | 2A | 3655 | 1/1 | 0.71 | 0.18 | 70,70,70,70 | 0 |
| 57 | MG | 2A | 3569 | 1/1 | 0.72 | 0.19 | 79,79,79,79 | 0 |
| 57 | MG | 2A | 3107 | 1/1 | 0.72 | 0.17 | 70,70,70,70 | 0 |
| 57 | MG | 2A | 3199 | 1/1 | 0.72 | 0.24 | 71,71,71,71 | 0 |
| 57 | MG | 2A | 3340 | 1/1 | 0.72 | 0.18 | 70,70,70,70 | 0 |
| 57 | MG | 1A | 3749 | 1/1 | 0.72 | 0.21 | 60,60,60,60 | 0 |
| 57 | MG | 2A | 3200 | 1/1 | 0.73 | 0.25 | 68,68,68,68 | 0 |
| 57 | MG | 1A | 4070 | 1/1 | 0.73 | 0.36 | 89,89,89,89 | 0 |
| 57 | MG | 2y | 101 | 1/1 | 0.73 | 0.29 | 79,79,79,79 | 0 |
| 57 | MG | 2A | 3321 | 1/1 | 0.74 | 0.16 | 84,84,84,84 | 0 |
| 57 | MG | 2A | 3326 | 1/1 | 0.74 | 0.25 | 77,77,77,77 | 0 |
| 57 | MG | 2A | 3305 | 1/1 | 0.74 | 0.18 | 62,62,62,62 | 0 |
| 57 | MG | 2A | 3103 | 1/1 | 0.75 | 0.17 | 76,76,76,76 | 0 |
| 57 | MG | 2a | 1695 | 1/1 | 0.75 | 0.48 | 81,81,81,81 | 0 |
| 57 | MG | 2A | 3276 | 1/1 | 0.75 | 0.13 | 70,70,70,70 | 0 |
| 57 | MG | 2B | 204 | 1/1 | 0.75 | 0.15 | 77,77,77,77 | 0 |
| 57 | MG | 2w | 101 | 1/1 | 0.75 | 0.28 | 83,83,83,83 | 0 |
| 57 | MG | 2B | 215 | 1/1 | 0.75 | 0.23 | 69,69,69,69 | 0 |
| 57 | MG | 2A | 3278 | 1/1 | 0.76 | 0.17 | 72,72,72,72 | 0 |
| 57 | MG | 1A | 4001 | 1/1 | 0.76 | 0.11 | 76,76,76,76 | 0 |
| 57 | MG | 2A | 3212 | 1/1 | 0.76 | 0.15 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3894 | 1/1 | 0.76 | 0.16 | 62,62,62,62 | 0 |
| 57 | MG | 1A | 3939 | 1/1 | 0.76 | 0.18 | 79,79,79,79 | 0 |
| 57 | MG | 2a | 1784 | 1/1 | 0.76 | 0.21 | 73,73,73,73 | 0 |
| 57 | MG | 2A | 3097 | 1/1 | 0.76 | 0.32 | 80,80,80,80 | 0 |
| 57 | MG | 2A | 3741 | 1/1 | 0.76 | 0.12 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3469 | 1/1 | 0.76 | 0.19 | 65,65,65,65 | 0 |
| 57 | MG | 2a | 1605 | 1/1 | 0.77 | 0.36 | 73,73,73,73 | 0 |
| 57 | MG | 1a | 1804 | 1/1 | 0.77 | 0.25 | 74,74,74,74 | 0 |
| 57 | MG | 2a | 1671 | 1/1 | 0.77 | 0.25 | 69,69,69,69 | 0 |
| 57 | MG | 1A | 3256 | 1/1 | 0.77 | 0.26 | 63,63,63,63 | 0 |
| 57 | MG | 2a | 1721 | 1/1 | 0.77 | 0.15 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3734 | 1/1 | 0.77 | 0.17 | 80,80,80,80 | 0 |
| 57 | MG | 2A | 3089 | 1/1 | 0.77 | 0.10 | 64,64,64,64 | 0 |
| 57 | MG | 1A | 3873 | 1/1 | 0.77 | 0.15 | 67,67,67,67 | 0 |
| 57 | MG | 2v | 102 | 1/1 | 0.77 | 0.30 | 80,80,80,80 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 3955 | 1/1 | 0.77 | 0.22 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3877 | 1/1 | 0.77 | 0.12 | 22,22,22,22 | 0 |
| 57 | MG | 1A | 3543 | 1/1 | 0.78 | 0.14 | 40,40,40,40 | 0 |
| 57 | MG | 2A | 3649 | 1/1 | 0.78 | 0.20 | 74,74,74,74 | 0 |
| 57 | MG | 1A | 3505 | 1/1 | 0.78 | 0.12 | 61,61,61,61 | 0 |
| 57 | MG | 2A | 3671 | 1/1 | 0.78 | 0.18 | 63,63,63,63 | 0 |
| 57 | MG | 2a | 1713 | 1/1 | 0.78 | 0.12 | 64,64,64,64 | 0 |
| 57 | MG | 2A | 3253 | 1/1 | 0.78 | 0.16 | 74,74,74,74 | 0 |
| 57 | MG | 2a | 1728 | 1/1 | 0.78 | 0.35 | 80,80,80,80 | 0 |
| 57 | MG | 1A | 3725 | 1/1 | 0.78 | 0.20 | 73,73,73,73 | 0 |
| 57 | MG | 2A | 3384 | 1/1 | 0.78 | 0.13 | 73,73,73,73 | 0 |
| 57 | MG | 2a | 1785 | 1/1 | 0.78 | 0.17 | 85,85,85,85 | 0 |
| 57 | MG | 1A | 3968 | 1/1 | 0.78 | 0.13 | 61,61,61,61 | 0 |
| 57 | MG | 2a | 1809 | 1/1 | 0.78 | 0.29 | 65,65,65,65 | 0 |
| 57 | MG | 1a | 1607 | 1/1 | 0.78 | 0.31 | 73,73,73,73 | 0 |
| 57 | MG | 2A | 3294 | 1/1 | 0.78 | 0.12 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3302 | 1/1 | 0.78 | 0.15 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3184 | 1/1 | 0.79 | 0.29 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3785 | 1/1 | 0.79 | 0.13 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3065 | 1/1 | 0.79 | 0.12 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3383 | 1/1 | 0.79 | 0.09 | 63,63,63,63 | 0 |
| 57 | MG | 1B | 236 | 1/1 | 0.79 | 0.18 | 76,76,76,76 | 0 |
| 57 | MG | 2a | 1786 | 1/1 | 0.79 | 0.15 | 66,66,66,66 | 0 |
| 57 | MG | 1A | 3833 | 1/1 | 0.79 | 0.17 | 70,70,70,70 | 0 |
| 57 | MG | 1A | 3228 | 1/1 | 0.79 | 0.20 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3052 | 1/1 | 0.79 | 0.16 | 73,73,73,73 | 0 |
| 57 | MG | 2A | 3168 | 1/1 | 0.79 | 0.26 | 77,77,77,77 | 0 |
| 57 | MG | 2a | 1720 | 1/1 | 0.79 | 0.26 | 55,55,55,55 | 0 |
| 57 | MG | 2a | 1639 | 1/1 | 0.80 | 0.21 | 76,76,76,76 | 0 |
| 57 | MG | 2A | 3600 | 1/1 | 0.80 | 0.10 | 74,74,74,74 | 0 |
| 57 | MG | 1a | 1736 | 1/1 | 0.80 | 0.20 | 65,65,65,65 | 0 |
| 57 | MG | 1a | 1738 | 1/1 | 0.80 | 0.14 | 74,74,74,74 | 0 |
| 57 | MG | 2a | 1705 | 1/1 | 0.80 | 0.26 | 78,78,78,78 | 0 |
| 57 | MG | 1a | 1765 | 1/1 | 0.80 | 0.11 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3809 | 1/1 | 0.80 | 0.14 | 48,48,48,48 | 0 |
| 57 | MG | 1w | 101 | 1/1 | 0.80 | 0.17 | 66,66,66,66 | 0 |
| 57 | MG | 1A | 4074 | 1/1 | 0.80 | 0.18 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3382 | 1/1 | 0.80 | 0.12 | 74,74,74,74 | 0 |
| 57 | MG | 2a | 1783 | 1/1 | 0.80 | 0.15 | 71,71,71,71 | 0 |
| 57 | MG | 1A | 3347 | 1/1 | 0.80 | 0.19 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3754 | 1/1 | 0.80 | 0.12 | 59,59,59,59 | 0 |
| 57 | MG | 1A | 3947 | 1/1 | 0.80 | 0.19 | 58,58,58,58 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2a | 1788 | 1/1 | 0.80 | 0.21 | 72,72,72,72 | 0 |
| 57 | MG | 1A | 3687 | 1/1 | 0.80 | 0.12 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3827 | 1/1 | 0.80 | 0.13 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3093 | 1/1 | 0.80 | 0.15 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3874 | 1/1 | 0.80 | 0.13 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3774 | 1/1 | 0.80 | 0.10 | 20,20,20,20 | 0 |
| 57 | MG | 2A | 3242 | 1/1 | 0.81 | 0.14 | 57,57,57,57 | 0 |
| 57 | MG | 2a | 1736 | 1/1 | 0.81 | 0.20 | 69,69,69,69 | 0 |
| 57 | MG | 1a | 1713 | 1/1 | 0.81 | 0.16 | 50,50,50,50 | 0 |
| 57 | MG | 2a | 1771 | 1/1 | 0.81 | 0.32 | 77,77,77,77 | 0 |
| 57 | MG | 1A | 3206 | 1/1 | 0.81 | 0.13 | 67,67,67,67 | 0 |
| 57 | MG | 2a | 1618 | 1/1 | 0.81 | 0.14 | 61,61,61,61 | 0 |
| 57 | MG | 2A | 3690 | 1/1 | 0.81 | 0.23 | 77,77,77,77 | 0 |
| 57 | MG | 2A | 3021 | 1/1 | 0.81 | 0.24 | 72,72,72,72 | 0 |
| 57 | MG | 2A | 3531 | 1/1 | 0.81 | 0.29 | 65,65,65,65 | 0 |
| 57 | MG | 1A | 3555 | 1/1 | 0.81 | 0.14 | 46,46,46,46 | 0 |
| 57 | MG | 2a | 1808 | 1/1 | 0.81 | 0.14 | 64,64,64,64 | 0 |
| 57 | MG | 2A | 3355 | 1/1 | 0.81 | 0.24 | 73,73,73,73 | 0 |
| 57 | MG | 2A | 3366 | 1/1 | 0.81 | 0.16 | 74,74,74,74 | 0 |
| 57 | MG | 2A | 3375 | 1/1 | 0.81 | 0.38 | 66,66,66,66 | 0 |
| 57 | MG | 2x | 102 | 1/1 | 0.81 | 0.10 | 71,71,71,71 | 0 |
| 57 | MG | 2A | 3188 | 1/1 | 0.81 | 0.11 | 62,62,62,62 | 0 |
| 57 | MG | 1A | 3860 | 1/1 | 0.82 | 0.09 | 45,45,45,45 | 0 |
| 57 | MG | 2a | 1744 | 1/1 | 0.82 | 0.29 | 69,69,69,69 | 0 |
| 57 | MG | 2B | 219 | 1/1 | 0.82 | 0.22 | 67,67,67,67 | 0 |
| 57 | MG | 2A | 3252 | 1/1 | 0.82 | 0.18 | 71,71,71,71 | 0 |
| 57 | MG | 1w | 102 | 1/1 | 0.82 | 0.09 | 69,69,69,69 | 0 |
| 57 | MG | 1a | 1678 | 1/1 | 0.82 | 0.16 | 59,59,59,59 | 0 |
| 57 | MG | 2A | 3737 | 1/1 | 0.82 | 0.17 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3322 | 1/1 | 0.82 | 0.14 | 53,53,53,53 | 0 |
| 57 | MG | 1a | 1795 | 1/1 | 0.82 | 0.09 | 75,75,75,75 | 0 |
| 57 | MG | 2A | 3159 | 1/1 | 0.82 | 0.15 | 70,70,70,70 | 0 |
| 57 | MG | 2A | 3301 | 1/1 | 0.82 | 0.16 | 71,71,71,71 | 0 |
| 57 | MG | 1A | 4080 | 1/1 | 0.82 | 0.27 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3837 | 1/1 | 0.82 | 0.12 | 74,74,74,74 | 0 |
| 57 | MG | 2A | 3842 | 1/1 | 0.82 | 0.13 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3303 | 1/1 | 0.82 | 0.23 | 60,60,60,60 | 0 |
| 57 | MG | 2a | 1637 | 1/1 | 0.83 | 0.19 | 65,65,65,65 | 0 |
| 57 | MG | 2a | 1638 | 1/1 | 0.83 | 0.28 | 70,70,70,70 | 0 |
| 57 | MG | 2A | 3654 | 1/1 | 0.83 | 0.11 | 82,82,82,82 | 0 |
| 57 | MG | 1w | 103 | 1/1 | 0.83 | 0.18 | 72,72,72,72 | 0 |
| 57 | MG | 2a | 1654 | 1/1 | 0.83 | 0.20 | 83,83,83,83 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2a | 1663 | 1/1 | 0.83 | 0.27 | 63,63,63,63 | 0 |
| 57 | MG | 1a | 1656 | 1/1 | 0.83 | 0.12 | 64,64,64,64 | 0 |
| 57 | MG | 1A | 3865 | 1/1 | 0.83 | 0.23 | 45,45,45,45 | 0 |
| 57 | MG | 1a | 1694 | 1/1 | 0.83 | 0.20 | 78,78,78,78 | 0 |
| 57 | MG | 2A | 3692 | 1/1 | 0.83 | 0.11 | 69,69,69,69 | 0 |
| 57 | MG | 2a | 1715 | 1/1 | 0.83 | 0.23 | 61,61,61,61 | 0 |
| 57 | MG | 1A | 3944 | 1/1 | 0.83 | 0.13 | 60,60,60,60 | 0 |
| 57 | MG | 2A | 3718 | 1/1 | 0.83 | 0.28 | 63,63,63,63 | 0 |
| 57 | MG | 2a | 1727 | 1/1 | 0.83 | 0.08 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3101 | 1/1 | 0.83 | 0.08 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3717 | 1/1 | 0.83 | 0.08 | 62,62,62,62 | 0 |
| 57 | MG | 1A | 3579 | 1/1 | 0.83 | 0.25 | 65,65,65,65 | 0 |
| 57 | MG | 2A | 3257 | 1/1 | 0.83 | 0.22 | 63,63,63,63 | 0 |
| 57 | MG | 2a | 1745 | 1/1 | 0.83 | 0.24 | 73,73,73,73 | 0 |
| 57 | MG | 2A | 3389 | 1/1 | 0.83 | 0.13 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3101 | 1/1 | 0.83 | 0.26 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3444 | 1/1 | 0.83 | 0.17 | 68,68,68,68 | 0 |
| 57 | MG | 1D | 311 | 1/1 | 0.83 | 0.15 | 33,33,33,33 | 0 |
| 57 | MG | 1a | 1778 | 1/1 | 0.83 | 0.17 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3850 | 1/1 | 0.83 | 0.23 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3134 | 1/1 | 0.83 | 0.25 | 65,65,65,65 | 0 |
| 57 | MG | 2a | 1803 | 1/1 | 0.83 | 0.21 | 72,72,72,72 | 0 |
| 57 | MG | 2B | 214 | 1/1 | 0.83 | 0.17 | 75,75,75,75 | 0 |
| 57 | MG | 1G | 203 | 1/1 | 0.83 | 0.10 | 54,54,54,54 | 0 |
| 57 | MG | 2a | 1818 | 1/1 | 0.83 | 0.21 | 74,74,74,74 | 0 |
| 57 | MG | 2d | 302 | 1/1 | 0.83 | 0.20 | 76,76,76,76 | 0 |
| 57 | MG | 1Q | 204 | 1/1 | 0.83 | 0.10 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3987 | 1/1 | 0.83 | 0.24 | 62,62,62,62 | 0 |
| 57 | MG | 1A | 3027 | 1/1 | 0.83 | 0.21 | 54,54,54,54 | 0 |
| 57 | MG | 2x | 103 | 1/1 | 0.83 | 0.24 | 74,74,74,74 | 0 |
| 57 | MG | 2a | 1635 | 1/1 | 0.83 | 0.25 | 63,63,63,63 | 0 |
| 57 | MG | 2A | 3255 | 1/1 | 0.84 | 0.20 | 58,58,58,58 | 0 |
| 57 | MG | 1a | 1660 | 1/1 | 0.84 | 0.19 | 63,63,63,63 | 0 |
| 57 | MG | 1A | 3830 | 1/1 | 0.84 | 0.13 | 67,67,67,67 | 0 |
| 57 | MG | 2A | 3789 | 1/1 | 0.84 | 0.11 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3137 | 1/1 | 0.84 | 0.20 | 63,63,63,63 | 0 |
| 57 | MG | 1F | 301 | 1/1 | 0.84 | 0.22 | 36,36,36,36 | 0 |
| 57 | MG | 2A | 3289 | 1/1 | 0.84 | 0.15 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3161 | 1/1 | 0.84 | 0.21 | 51,51,51,51 | 0 |
| 57 | MG | 2B | 203 | 1/1 | 0.84 | 0.17 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3173 | 1/1 | 0.84 | 0.17 | 49,49,49,49 | 0 |
| 57 | MG | 2a | 1750 | 1/1 | 0.84 | 0.17 | 68,68,68,68 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2a | 1764 | 1/1 | 0.84 | 0.16 | 73,73,73,73 | 0 |
| 57 | MG | 2A | 3174 | 1/1 | 0.84 | 0.12 | 66,66,66,66 | 0 |
| 57 | MG | 1x | 109 | 1/1 | 0.84 | 0.13 | 70,70,70,70 | 0 |
| 57 | MG | 1B | 216 | 1/1 | 0.84 | 0.11 | 61,61,61,61 | 0 |
| 57 | MG | 1A | 3250 | 1/1 | 0.84 | 0.21 | 74,74,74,74 | 0 |
| 57 | MG | 1B | 233 | 1/1 | 0.84 | 0.15 | 61,61,61,61 | 0 |
| 57 | MG | 2a | 1633 | 1/1 | 0.84 | 0.32 | 73,73,73,73 | 0 |
| 57 | MG | 2A | 3676 | 1/1 | 0.84 | 0.15 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3086 | 1/1 | 0.84 | 0.18 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3684 | 1/1 | 0.84 | 0.09 | 69,69,69,69 | 0 |
| 57 | MG | 1a | 1622 | 1/1 | 0.84 | 0.19 | 62,62,62,62 | 0 |
| 57 | MG | 2a | 1812 | 1/1 | 0.84 | 0.28 | 69,69,69,69 | 0 |
| 57 | MG | 1a | 1768 | 1/1 | 0.84 | 0.12 | 71,71,71,71 | 0 |
| 57 | MG | 2A | 3367 | 1/1 | 0.84 | 0.16 | 75,75,75,75 | 0 |
| 57 | MG | 2j | 201 | 1/1 | 0.84 | 0.20 | 63,63,63,63 | 0 |
| 57 | MG | 2q | 202 | 1/1 | 0.84 | 0.13 | 77,77,77,77 | 0 |
| 57 | MG | 2a | 1659 | 1/1 | 0.84 | 0.12 | 73,73,73,73 | 0 |
| 57 | MG | 2A | 3237 | 1/1 | 0.84 | 0.13 | 60,60,60,60 | 0 |
| 57 | MG | 2x | 101 | 1/1 | 0.84 | 0.20 | 72,72,72,72 | 0 |
| 57 | MG | 1a | 1655 | 1/1 | 0.84 | 0.19 | 63,63,63,63 | 0 |
| 57 | MG | 1A | 3642 | 1/1 | 0.84 | 0.11 | 43,43,43,43 | 0 |
| 57 | MG | 1a | 1659 | 1/1 | 0.84 | 0.28 | 74,74,74,74 | 0 |
| 57 | MG | 2A | 3339 | 1/1 | 0.85 | 0.08 | 56,56,56,56 | 0 |
| 57 | MG | 1a | 1709 | 1/1 | 0.85 | 0.28 | 46,46,46,46 | 0 |
| 57 | MG | 2a | 1677 | 1/1 | 0.85 | 0.13 | 61,61,61,61 | 0 |
| 57 | MG | 2A | 3343 | 1/1 | 0.85 | 0.13 | 53,53,53,53 | 0 |
| 57 | MG | 2a | 1703 | 1/1 | 0.85 | 0.18 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3099 | 1/1 | 0.85 | 0.16 | 52,52,52,52 | 0 |
| 57 | MG | 1a | 1806 | 1/1 | 0.85 | 0.21 | 74,74,74,74 | 0 |
| 57 | MG | 1A | 3647 | 1/1 | 0.85 | 0.14 | 65,65,65,65 | 0 |
| 57 | MG | 1a | 1719 | 1/1 | 0.85 | 0.16 | 64,64,64,64 | 0 |
| 57 | MG | 1a | 1630 | 1/1 | 0.85 | 0.23 | 73,73,73,73 | 0 |
| 57 | MG | 1A | 3935 | 1/1 | 0.85 | 0.24 | 64,64,64,64 | 0 |
| 57 | MG | 1A | 3128 | 1/1 | 0.85 | 0.13 | 59,59,59,59 | 0 |
| 57 | MG | 2a | 1732 | 1/1 | 0.85 | 0.14 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3258 | 1/1 | 0.85 | 0.11 | 82,82,82,82 | 0 |
| 57 | MG | 2A | 3410 | 1/1 | 0.85 | 0.13 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3260 | 1/1 | 0.85 | 0.10 | 65,65,65,65 | 0 |
| 57 | MG | 2A | 3261 | 1/1 | 0.85 | 0.12 | 63,63,63,63 | 0 |
| 57 | MG | 2A | 3457 | 1/1 | 0.85 | 0.12 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3852 | 1/1 | 0.85 | 0.14 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3264 | 1/1 | 0.85 | 0.17 | 53,53,53,53 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3165 | 1/1 | 0.85 | 0.08 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 4082 | 1/1 | 0.85 | 0.18 | 69,69,69,69 | 0 |
| 57 | MG | 1A | 4007 | 1/1 | 0.85 | 0.17 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3057 | 1/1 | 0.85 | 0.21 | 62,62,62,62 | 0 |
| 57 | MG | 2W | 202 | 1/1 | 0.85 | 0.14 | 57,57,57,57 | 0 |
| 57 | MG | 2a | 1797 | 1/1 | 0.85 | 0.14 | 79,79,79,79 | 0 |
| 57 | MG | 2a | 1604 | 1/1 | 0.85 | 0.09 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3637 | 1/1 | 0.85 | 0.13 | 40,40,40,40 | 0 |
| 57 | MG | 2a | 1614 | 1/1 | 0.85 | 0.26 | 68,68,68,68 | 0 |
| 57 | MG | 2A | 3064 | 1/1 | 0.85 | 0.20 | 41,41,41,41 | 0 |
| 57 | MG | 2a | 1624 | 1/1 | 0.85 | 0.15 | 65,65,65,65 | 0 |
| 57 | MG | 1A | 4021 | 1/1 | 0.85 | 0.10 | 35,35,35,35 | 0 |
| 57 | MG | 2a | 1822 | 1/1 | 0.85 | 0.18 | 67,67,67,67 | 0 |
| 57 | MG | 2A | 3195 | 1/1 | 0.85 | 0.10 | 61,61,61,61 | 0 |
| 57 | MG | 1a | 1789 | 1/1 | 0.85 | 0.16 | 77,77,77,77 | 0 |
| 57 | MG | 2A | 3664 | 1/1 | 0.85 | 0.14 | 48,48,48,48 | 0 |
| 57 | MG | 2v | 101 | 1/1 | 0.85 | 0.21 | 68,68,68,68 | 0 |
| 57 | MG | 1A | 4025 | 1/1 | 0.85 | 0.12 | 35,35,35,35 | 0 |
| 57 | MG | 2a | 1642 | 1/1 | 0.85 | 0.13 | 78,78,78,78 | 0 |
| 57 | MG | 2w | 105 | 1/1 | 0.85 | 0.13 | 64,64,64,64 | 0 |
| 57 | MG | 1a | 1798 | 1/1 | 0.85 | 0.22 | 63,63,63,63 | 0 |
| 57 | MG | 2A | 3206 | 1/1 | 0.85 | 0.13 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3334 | 1/1 | 0.85 | 0.22 | 53,53,53,53 | 0 |
| 57 | MG | 2a | 1661 | 1/1 | 0.85 | 0.22 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3423 | 1/1 | 0.86 | 0.13 | 42,42,42,42 | 0 |
| 57 | MG | 2A | 3784 | 1/1 | 0.86 | 0.08 | 67,67,67,67 | 0 |
| 57 | MG | 1A | 3458 | 1/1 | 0.86 | 0.15 | 70,70,70,70 | 0 |
| 57 | MG | 1a | 1799 | 1/1 | 0.86 | 0.23 | 62,62,62,62 | 0 |
| 57 | MG | 2A | 3826 | 1/1 | 0.86 | 0.15 | 54,54,54,54 | 0 |
| 57 | MG | 2a | 1719 | 1/1 | 0.86 | 0.18 | 60,60,60,60 | 0 |
| 57 | MG | 2A | 3495 | 1/1 | 0.86 | 0.12 | 61,61,61,61 | 0 |
| 57 | MG | 1A | 4023 | 1/1 | 0.86 | 0.12 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3315 | 1/1 | 0.86 | 0.15 | 62,62,62,62 | 0 |
| 57 | MG | 2A | 3575 | 1/1 | 0.86 | 0.17 | 62,62,62,62 | 0 |
| 57 | MG | 1A | 3471 | 1/1 | 0.86 | 0.10 | 64,64,64,64 | 0 |
| 57 | MG | 2a | 1733 | 1/1 | 0.86 | 0.10 | 67,67,67,67 | 0 |
| 57 | MG | 2A | 3584 | 1/1 | 0.86 | 0.10 | 68,68,68,68 | 0 |
| 57 | MG | 2a | 1737 | 1/1 | 0.86 | 0.18 | 64,64,64,64 | 0 |
| 57 | MG | 2A | 3599 | 1/1 | 0.86 | 0.18 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 4069 | 1/1 | 0.86 | 0.11 | 75,75,75,75 | 0 |
| 57 | MG | 2A | 3614 | 1/1 | 0.86 | 0.22 | 59,59,59,59 | 0 |
| 57 | MG | 2A | 3327 | 1/1 | 0.86 | 0.14 | 80,80,80,80 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2a | 1760 | 1/1 | 0.86 | 0.20 | 70,70,70,70 | 0 |
| 57 | MG | 2B | 220 | 1/1 | 0.86 | 0.16 | 65,65,65,65 | 0 |
| 57 | MG | 2E | 309 | 1/1 | 0.86 | 0.08 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3301 | 1/1 | 0.86 | 0.11 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3254 | 1/1 | 0.86 | 0.17 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3155 | 1/1 | 0.86 | 0.10 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3959 | 1/1 | 0.86 | 0.10 | 66,66,66,66 | 0 |
| 57 | MG | 2a | 1787 | 1/1 | 0.86 | 0.28 | 72,72,72,72 | 0 |
| 57 | MG | 1a | 1723 | 1/1 | 0.86 | 0.24 | 59,59,59,59 | 0 |
| 57 | MG | 2a | 1621 | 1/1 | 0.86 | 0.17 | 60,60,60,60 | 0 |
| 57 | MG | 2A | 3356 | 1/1 | 0.86 | 0.21 | 68,68,68,68 | 0 |
| 57 | MG | 2a | 1801 | 1/1 | 0.86 | 0.22 | 64,64,64,64 | 0 |
| 57 | MG | 2a | 1626 | 1/1 | 0.86 | 0.08 | 88,88,88,88 | 0 |
| 57 | MG | 2a | 1805 | 1/1 | 0.86 | 0.21 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3358 | 1/1 | 0.86 | 0.18 | 64,64,64,64 | 0 |
| 57 | MG | 2a | 1634 | 1/1 | 0.86 | 0.35 | 68,68,68,68 | 0 |
| 57 | MG | 1A | 3050 | 1/1 | 0.86 | 0.13 | 33,33,33,33 | 0 |
| 57 | MG | 1B | 214 | 1/1 | 0.86 | 0.10 | 40,40,40,40 | 0 |
| 57 | MG | 1a | 1763 | 1/1 | 0.86 | 0.12 | 60,60,60,60 | 0 |
| 57 | MG | 2A | 3691 | 1/1 | 0.86 | 0.12 | 70,70,70,70 | 0 |
| 57 | MG | 1a | 1647 | 1/1 | 0.86 | 0.15 | 63,63,63,63 | 0 |
| 57 | MG | 2A | 3699 | 1/1 | 0.86 | 0.23 | 64,64,64,64 | 0 |
| 57 | MG | 1a | 1652 | 1/1 | 0.86 | 0.18 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3071 | 1/1 | 0.86 | 0.11 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3910 | 1/1 | 0.86 | 0.18 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3916 | 1/1 | 0.86 | 0.10 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3424 | 1/1 | 0.86 | 0.10 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3299 | 1/1 | 0.86 | 0.18 | 72,72,72,72 | 0 |
| 57 | MG | 2a | 1678 | 1/1 | 0.86 | 0.15 | 70,70,70,70 | 0 |
| 57 | MG | 2a | 1682 | 1/1 | 0.86 | 0.28 | 56,56,56,56 | 0 |
| 57 | MG | 1B | 210 | 1/1 | 0.87 | 0.09 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3812 | 1/1 | 0.87 | 0.14 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3566 | 1/1 | 0.87 | 0.21 | 56,56,56,56 | 0 |
| 57 | MG | 1a | 1676 | 1/1 | 0.87 | 0.17 | 62,62,62,62 | 0 |
| 57 | MG | 2a | 1676 | 1/1 | 0.87 | 0.19 | 64,64,64,64 | 0 |
| 57 | MG | 1B | 226 | 1/1 | 0.87 | 0.13 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3717 | 1/1 | 0.87 | 0.21 | 70,70,70,70 | 0 |
| 57 | MG | 2A | 3232 | 1/1 | 0.87 | 0.17 | 57,57,57,57 | 0 |
| 57 | MG | 1a | 1688 | 1/1 | 0.87 | 0.27 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3065 | 1/1 | 0.87 | 0.13 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3239 | 1/1 | 0.87 | 0.15 | 64,64,64,64 | 0 |
| 57 | MG | 2a | 1707 | 1/1 | 0.87 | 0.16 | 55,55,55,55 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2a | 1709 | 1/1 | 0.87 | 0.18 | 75,75,75,75 | 0 |
| 57 | MG | 2A | 3747 | 1/1 | 0.87 | 0.12 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3845 | 1/1 | 0.87 | 0.10 | 51,51,51,51 | 0 |
| 57 | MG | 2a | 1717 | 1/1 | 0.87 | 0.16 | 62,62,62,62 | 0 |
| 57 | MG | 2A | 3755 | 1/1 | 0.87 | 0.12 | 65,65,65,65 | 0 |
| 57 | MG | 2A | 3767 | 1/1 | 0.87 | 0.10 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3386 | 1/1 | 0.87 | 0.10 | 52,52,52,52 | 0 |
| 57 | MG | 2a | 1726 | 1/1 | 0.87 | 0.30 | 75,75,75,75 | 0 |
| 57 | MG | 1A | 3371 | 1/1 | 0.87 | 0.09 | 62,62,62,62 | 0 |
| 57 | MG | 1a | 1716 | 1/1 | 0.87 | 0.15 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3423 | 1/1 | 0.87 | 0.18 | 62,62,62,62 | 0 |
| 57 | MG | 2A | 3794 | 1/1 | 0.87 | 0.07 | 72,72,72,72 | 0 |
| 57 | MG | 1A | 3989 | 1/1 | 0.87 | 0.13 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3130 | 1/1 | 0.87 | 0.06 | 70,70,70,70 | 0 |
| 57 | MG | 1F | 311 | 1/1 | 0.87 | 0.14 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3838 | 1/1 | 0.87 | 0.11 | 62,62,62,62 | 0 |
| 57 | MG | 2A | 3094 | 1/1 | 0.87 | 0.21 | 68,68,68,68 | 0 |
| 57 | MG | 2A | 3466 | 1/1 | 0.87 | 0.08 | 70,70,70,70 | 0 |
| 57 | MG | 2a | 1756 | 1/1 | 0.87 | 0.06 | 87,87,87,87 | 0 |
| 57 | MG | 1A | 3066 | 1/1 | 0.87 | 0.13 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3487 | 1/1 | 0.87 | 0.20 | 75,75,75,75 | 0 |
| 57 | MG | 1a | 1737 | 1/1 | 0.87 | 0.12 | 54,54,54,54 | 0 |
| 57 | MG | 2B | 208 | 1/1 | 0.87 | 0.19 | 62,62,62,62 | 0 |
| 57 | MG | 1A | 3654 | 1/1 | 0.87 | 0.12 | 58,58,58,58 | 0 |
| 57 | MG | 1Q | 207 | 1/1 | 0.87 | 0.10 | 37,37,37,37 | 0 |
| 57 | MG | 1R | 203 | 1/1 | 0.87 | 0.16 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3112 | 1/1 | 0.87 | 0.24 | 63,63,63,63 | 0 |
| 57 | MG | 1T | 201 | 1/1 | 0.87 | 0.12 | 50,50,50,50 | 0 |
| 57 | MG | 2P | 202 | 1/1 | 0.87 | 0.26 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3597 | 1/1 | 0.87 | 0.14 | 50,50,50,50 | 0 |
| 57 | MG | 2Z | 301 | 1/1 | 0.87 | 0.20 | 72,72,72,72 | 0 |
| 57 | MG | 1A | 3180 | 1/1 | 0.87 | 0.16 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3704 | 1/1 | 0.87 | 0.07 | 34,34,34,34 | 0 |
| 57 | MG | 2a | 1606 | 1/1 | 0.87 | 0.16 | 70,70,70,70 | 0 |
| 57 | MG | 2a | 1613 | 1/1 | 0.87 | 0.18 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3601 | 1/1 | 0.87 | 0.16 | 53,53,53,53 | 0 |
| 57 | MG | 2a | 1814 | 1/1 | 0.87 | 0.16 | 75,75,75,75 | 0 |
| 57 | MG | 1a | 1621 | 1/1 | 0.87 | 0.38 | 73,73,73,73 | 0 |
| 57 | MG | 2A | 3620 | 1/1 | 0.87 | 0.17 | 64,64,64,64 | 0 |
| 57 | MG | 1A | 3492 | 1/1 | 0.87 | 0.08 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 3200 | 1/1 | 0.87 | 0.11 | 47,47,47,47 | 0 |
| 57 | MG | 2k | 201 | 1/1 | 0.87 | 0.16 | 65,65,65,65 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3170 | 1/1 | 0.87 | 0.12 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3318 | 1/1 | 0.87 | 0.16 | 54,54,54,54 | 0 |
| 57 | MG | 1a | 1800 | 1/1 | 0.87 | 0.10 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3513 | 1/1 | 0.87 | 0.09 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3180 | 1/1 | 0.87 | 0.15 | 62,62,62,62 | 0 |
| 57 | MG | 2A | 3672 | 1/1 | 0.87 | 0.11 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3302 | 1/1 | 0.87 | 0.17 | 53,53,53,53 | 0 |
| 57 | MG | 1a | 1807 | 1/1 | 0.87 | 0.13 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3053 | 1/1 | 0.87 | 0.18 | 55,55,55,55 | 0 |
| 57 | MG | 2a | 1646 | 1/1 | 0.88 | 0.18 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3329 | 1/1 | 0.88 | 0.22 | 68,68,68,68 | 0 |
| 57 | MG | 2a | 1648 | 1/1 | 0.88 | 0.13 | 73,73,73,73 | 0 |
| 57 | MG | 2A | 3194 | 1/1 | 0.88 | 0.11 | 54,54,54,54 | 0 |
| 57 | MG | 2a | 1656 | 1/1 | 0.88 | 0.16 | 65,65,65,65 | 0 |
| 57 | MG | 1a | 1708 | 1/1 | 0.88 | 0.17 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3018 | 1/1 | 0.88 | 0.09 | 59,59,59,59 | 0 |
| 57 | MG | 1A | 3246 | 1/1 | 0.88 | 0.07 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3044 | 1/1 | 0.88 | 0.08 | 78,78,78,78 | 0 |
| 57 | MG | 1A | 3718 | 1/1 | 0.88 | 0.10 | 48,48,48,48 | 0 |
| 57 | MG | 1O | 201 | 1/1 | 0.88 | 0.08 | 61,61,61,61 | 0 |
| 57 | MG | 2A | 3365 | 1/1 | 0.88 | 0.34 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3215 | 1/1 | 0.88 | 0.19 | 64,64,64,64 | 0 |
| 57 | MG | 2A | 3222 | 1/1 | 0.88 | 0.19 | 43,43,43,43 | 0 |
| 57 | MG | 2a | 1697 | 1/1 | 0.88 | 0.11 | 68,68,68,68 | 0 |
| 57 | MG | 2A | 3372 | 1/1 | 0.88 | 0.09 | 40,40,40,40 | 0 |
| 57 | MG | 1a | 1717 | 1/1 | 0.88 | 0.13 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3376 | 1/1 | 0.88 | 0.10 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3378 | 1/1 | 0.88 | 0.10 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3379 | 1/1 | 0.88 | 0.25 | 66,66,66,66 | 0 |
| 57 | MG | 1A | 3875 | 1/1 | 0.88 | 0.13 | 67,67,67,67 | 0 |
| 57 | MG | 1A | 3414 | 1/1 | 0.88 | 0.12 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3884 | 1/1 | 0.88 | 0.13 | 25,25,25,25 | 0 |
| 57 | MG | 2A | 3241 | 1/1 | 0.88 | 0.44 | 76,76,76,76 | 0 |
| 57 | MG | 1A | 3494 | 1/1 | 0.88 | 0.13 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3786 | 1/1 | 0.88 | 0.13 | 59,59,59,59 | 0 |
| 57 | MG | 2A | 3246 | 1/1 | 0.88 | 0.14 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3419 | 1/1 | 0.88 | 0.18 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3821 | 1/1 | 0.88 | 0.10 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3900 | 1/1 | 0.88 | 0.10 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3599 | 1/1 | 0.88 | 0.35 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3641 | 1/1 | 0.88 | 0.13 | 27,27,27,27 | 0 |
| 57 | MG | 1A | 4081 | 1/1 | 0.88 | 0.12 | 51,51,51,51 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2a | 1743 | 1/1 | 0.88 | 0.21 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3454 | 1/1 | 0.88 | 0.15 | 51,51,51,51 | 0 |
| 57 | MG | 1a | 1628 | 1/1 | 0.88 | 0.17 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3917 | 1/1 | 0.88 | 0.09 | 27,27,27,27 | 0 |
| 57 | MG | 2A | 3468 | 1/1 | 0.88 | 0.23 | 64,64,64,64 | 0 |
| 57 | MG | 2a | 1758 | 1/1 | 0.88 | 0.19 | 67,67,67,67 | 0 |
| 57 | MG | 2a | 1759 | 1/1 | 0.88 | 0.18 | 72,72,72,72 | 0 |
| 57 | MG | 1a | 1780 | 1/1 | 0.88 | 0.14 | 64,64,64,64 | 0 |
| 57 | MG | 2A | 3262 | 1/1 | 0.88 | 0.17 | 61,61,61,61 | 0 |
| 57 | MG | 2A | 3493 | 1/1 | 0.88 | 0.18 | 48,48,48,48 | 0 |
| 57 | MG | 2a | 1777 | 1/1 | 0.88 | 0.22 | 59,59,59,59 | 0 |
| 57 | MG | 1B | 203 | 1/1 | 0.88 | 0.11 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3499 | 1/1 | 0.88 | 0.16 | 64,64,64,64 | 0 |
| 57 | MG | 2A | 3119 | 1/1 | 0.88 | 0.13 | 69,69,69,69 | 0 |
| 57 | MG | 2A | 3544 | 1/1 | 0.88 | 0.10 | 41,41,41,41 | 0 |
| 57 | MG | 2F | 302 | 1/1 | 0.88 | 0.10 | 48,48,48,48 | 0 |
| 57 | MG | 1a | 1648 | 1/1 | 0.88 | 0.21 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3571 | 1/1 | 0.88 | 0.17 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3340 | 1/1 | 0.88 | 0.09 | 57,57,57,57 | 0 |
| 57 | MG | 20 | 101 | 1/1 | 0.88 | 0.20 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3279 | 1/1 | 0.88 | 0.24 | 69,69,69,69 | 0 |
| 57 | MG | 2A | 3149 | 1/1 | 0.88 | 0.27 | 64,64,64,64 | 0 |
| 57 | MG | 1A | 3427 | 1/1 | 0.88 | 0.19 | 65,65,65,65 | 0 |
| 57 | MG | 2A | 3158 | 1/1 | 0.88 | 0.13 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3526 | 1/1 | 0.88 | 0.23 | 62,62,62,62 | 0 |
| 57 | MG | 1A | 3841 | 1/1 | 0.88 | 0.15 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3441 | 1/1 | 0.88 | 0.10 | 56,56,56,56 | 0 |
| 57 | MG | 2a | 1821 | 1/1 | 0.88 | 0.16 | 65,65,65,65 | 0 |
| 57 | MG | 1a | 1665 | 1/1 | 0.88 | 0.15 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3309 | 1/1 | 0.88 | 0.18 | 63,63,63,63 | 0 |
| 57 | MG | 2a | 1627 | 1/1 | 0.88 | 0.21 | 67,67,67,67 | 0 |
| 57 | MG | 2a | 1628 | 1/1 | 0.88 | 0.30 | 69,69,69,69 | 0 |
| 57 | MG | 2A | 3641 | 1/1 | 0.88 | 0.13 | 59,59,59,59 | 0 |
| 57 | MG | 1A | 3846 | 1/1 | 0.88 | 0.15 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3694 | 1/1 | 0.88 | 0.09 | 25,25,25,25 | 0 |
| 57 | MG | 2a | 1636 | 1/1 | 0.88 | 0.37 | 71,71,71,71 | 0 |
| 57 | MG | 1A | 3096 | 1/1 | 0.88 | 0.12 | 61,61,61,61 | 0 |
| 57 | MG | 2A | 3324 | 1/1 | 0.88 | 0.11 | 63,63,63,63 | 0 |
| 57 | MG | 1A | 3870 | 1/1 | 0.88 | 0.17 | 50,50,50,50 | 0 |
| 57 | MG | 1x | 105 | 1/1 | 0.88 | 0.24 | 58,58,58,58 | 0 |
| 57 | MG | 2a | 1645 | 1/1 | 0.88 | 0.23 | 58,58,58,58 | 0 |
| 57 | MG | 2a | 1625 | 1/1 | 0.89 | 0.28 | 59,59,59,59 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1B | 231 | 1/1 | 0.89 | 0.08 | 46,46,46,46 | 0 |
| 57 | MG | 1a | 1700 | 1/1 | 0.89 | 0.20 | 63,63,63,63 | 0 |
| 57 | MG | 1A | 3837 | 1/1 | 0.89 | 0.09 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3292 | 1/1 | 0.89 | 0.10 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3403 | 1/1 | 0.89 | 0.08 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3602 | 1/1 | 0.89 | 0.12 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3612 | 1/1 | 0.89 | 0.13 | 43,43,43,43 | 0 |
| 57 | MG | 1B | 234 | 1/1 | 0.89 | 0.15 | 65,65,65,65 | 0 |
| 57 | MG | 1A | 3949 | 1/1 | 0.89 | 0.09 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3844 | 1/1 | 0.89 | 0.19 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3110 | 1/1 | 0.89 | 0.17 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3502 | 1/1 | 0.89 | 0.14 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3306 | 1/1 | 0.89 | 0.15 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3967 | 1/1 | 0.89 | 0.12 | 54,54,54,54 | 0 |
| 57 | MG | 1a | 1721 | 1/1 | 0.89 | 0.22 | 65,65,65,65 | 0 |
| 57 | MG | 1A | 3325 | 1/1 | 0.89 | 0.21 | 61,61,61,61 | 0 |
| 57 | MG | 1A | 3657 | 1/1 | 0.89 | 0.09 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3669 | 1/1 | 0.89 | 0.10 | 44,44,44,44 | 0 |
| 57 | MG | 2a | 1660 | 1/1 | 0.89 | 0.08 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3673 | 1/1 | 0.89 | 0.12 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3325 | 1/1 | 0.89 | 0.25 | 64,64,64,64 | 0 |
| 57 | MG | 2a | 1666 | 1/1 | 0.89 | 0.12 | 68,68,68,68 | 0 |
| 57 | MG | 1A | 3992 | 1/1 | 0.89 | 0.09 | 26,26,26,26 | 0 |
| 57 | MG | 2a | 1673 | 1/1 | 0.89 | 0.15 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3680 | 1/1 | 0.89 | 0.11 | 60,60,60,60 | 0 |
| 57 | MG | 2A | 3682 | 1/1 | 0.89 | 0.09 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3507 | 1/1 | 0.89 | 0.09 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3332 | 1/1 | 0.89 | 0.21 | 59,59,59,59 | 0 |
| 57 | MG | 2a | 1684 | 1/1 | 0.89 | 0.28 | 64,64,64,64 | 0 |
| 57 | MG | 2a | 1688 | 1/1 | 0.89 | 0.12 | 65,65,65,65 | 0 |
| 57 | MG | 2A | 3333 | 1/1 | 0.89 | 0.08 | 49,49,49,49 | 0 |
| 57 | MG | 2a | 1696 | 1/1 | 0.89 | 0.18 | 62,62,62,62 | 0 |
| 57 | MG | 1Y | 202 | 1/1 | 0.89 | 0.10 | 57,57,57,57 | 0 |
| 57 | MG | 1a | 1769 | 1/1 | 0.89 | 0.12 | 64,64,64,64 | 0 |
| 57 | MG | 1a | 1772 | 1/1 | 0.89 | 0.14 | 63,63,63,63 | 0 |
| 57 | MG | 2a | 1706 | 1/1 | 0.89 | 0.18 | 68,68,68,68 | 0 |
| 57 | MG | 2A | 3713 | 1/1 | 0.89 | 0.14 | 56,56,56,56 | 0 |
| 57 | MG | 13 | 105 | 1/1 | 0.89 | 0.10 | 51,51,51,51 | 0 |
| 57 | MG | 2a | 1711 | 1/1 | 0.89 | 0.27 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3346 | 1/1 | 0.89 | 0.14 | 61,61,61,61 | 0 |
| 57 | MG | 2A | 3720 | 1/1 | 0.89 | 0.21 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3722 | 1/1 | 0.89 | 0.14 | 59,59,59,59 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 14 | 101 | 1/1 | 0.89 | 0.19 | 75,75,75,75 | 0 |
| 57 | MG | 2A | 3735 | 1/1 | 0.89 | 0.19 | 69,69,69,69 | 0 |
| 57 | MG | 2A | 3181 | 1/1 | 0.89 | 0.14 | 55,55,55,55 | 0 |
| 57 | MG | 2a | 1723 | 1/1 | 0.89 | 0.25 | 51,51,51,51 | 0 |
| 57 | MG | 1a | 1784 | 1/1 | 0.89 | 0.15 | 61,61,61,61 | 0 |
| 57 | MG | 2A | 3187 | 1/1 | 0.89 | 0.32 | 63,63,63,63 | 0 |
| 57 | MG | 16 | 101 | 1/1 | 0.89 | 0.11 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3189 | 1/1 | 0.89 | 0.13 | 63,63,63,63 | 0 |
| 57 | MG | 1A | 3333 | 1/1 | 0.89 | 0.08 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3770 | 1/1 | 0.89 | 0.09 | 72,72,72,72 | 0 |
| 57 | MG | 1A | 3533 | 1/1 | 0.89 | 0.23 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3775 | 1/1 | 0.89 | 0.15 | 50,50,50,50 | 0 |
| 57 | MG | 1a | 1617 | 1/1 | 0.89 | 0.16 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3088 | 1/1 | 0.89 | 0.27 | 42,42,42,42 | 0 |
| 57 | MG | 2A | 3201 | 1/1 | 0.89 | 0.09 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 4027 | 1/1 | 0.89 | 0.07 | 29,29,29,29 | 0 |
| 57 | MG | 2a | 1754 | 1/1 | 0.89 | 0.12 | 71,71,71,71 | 0 |
| 57 | MG | 1a | 1623 | 1/1 | 0.89 | 0.23 | 59,59,59,59 | 0 |
| 57 | MG | 1a | 1627 | 1/1 | 0.89 | 0.17 | 54,54,54,54 | 0 |
| 57 | MG | 1p | 101 | 1/1 | 0.89 | 0.10 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3722 | 1/1 | 0.89 | 0.12 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3830 | 1/1 | 0.89 | 0.13 | 61,61,61,61 | 0 |
| 57 | MG | 2a | 1767 | 1/1 | 0.89 | 0.14 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3836 | 1/1 | 0.89 | 0.13 | 67,67,67,67 | 0 |
| 57 | MG | 1A | 3885 | 1/1 | 0.89 | 0.11 | 36,36,36,36 | 0 |
| 57 | MG | 2A | 3417 | 1/1 | 0.89 | 0.18 | 67,67,67,67 | 0 |
| 57 | MG | 2A | 3840 | 1/1 | 0.89 | 0.09 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3233 | 1/1 | 0.89 | 0.08 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3848 | 1/1 | 0.89 | 0.16 | 50,50,50,50 | 0 |
| 57 | MG | 1a | 1638 | 1/1 | 0.89 | 0.14 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3454 | 1/1 | 0.89 | 0.21 | 58,58,58,58 | 0 |
| 57 | MG | 2a | 1795 | 1/1 | 0.89 | 0.21 | 72,72,72,72 | 0 |
| 57 | MG | 2a | 1796 | 1/1 | 0.89 | 0.15 | 65,65,65,65 | 0 |
| 57 | MG | 2A | 3430 | 1/1 | 0.89 | 0.21 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3238 | 1/1 | 0.89 | 0.25 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3257 | 1/1 | 0.89 | 0.10 | 63,63,63,63 | 0 |
| 57 | MG | 2A | 3240 | 1/1 | 0.89 | 0.19 | 61,61,61,61 | 0 |
| 57 | MG | 1A | 3467 | 1/1 | 0.89 | 0.07 | 56,56,56,56 | 0 |
| 57 | MG | 2B | 217 | 1/1 | 0.89 | 0.12 | 66,66,66,66 | 0 |
| 57 | MG | 1A | 3585 | 1/1 | 0.89 | 0.08 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3300 | 1/1 | 0.89 | 0.17 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3041 | 1/1 | 0.89 | 0.28 | 48,48,48,48 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3486 | 1/1 | 0.89 | 0.18 | 55,55,55,55 | 0 |
| 57 | MG | 1B | 209 | 1/1 | 0.89 | 0.13 | 48,48,48,48 | 0 |
| 57 | MG | 2Q | 201 | 1/1 | 0.89 | 0.12 | 68,68,68,68 | 0 |
| 57 | MG | 2T | 202 | 1/1 | 0.89 | 0.19 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3933 | 1/1 | 0.89 | 0.10 | 31,31,31,31 | 0 |
| 57 | MG | 2A | 3054 | 1/1 | 0.89 | 0.17 | 66,66,66,66 | 0 |
| 57 | MG | 1B | 211 | 1/1 | 0.89 | 0.12 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3511 | 1/1 | 0.89 | 0.10 | 24,24,24,24 | 0 |
| 57 | MG | 2A | 3062 | 1/1 | 0.89 | 0.19 | 66,66,66,66 | 0 |
| 57 | MG | 1a | 1672 | 1/1 | 0.89 | 0.12 | 64,64,64,64 | 0 |
| 57 | MG | 2A | 3566 | 1/1 | 0.89 | 0.20 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3489 | 1/1 | 0.89 | 0.12 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3391 | 1/1 | 0.89 | 0.20 | 45,45,45,45 | 0 |
| 57 | MG | 1a | 1682 | 1/1 | 0.89 | 0.18 | 63,63,63,63 | 0 |
| 57 | MG | 1A | 3941 | 1/1 | 0.89 | 0.11 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3438 | 1/1 | 0.90 | 0.18 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3243 | 1/1 | 0.90 | 0.14 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3193 | 1/1 | 0.90 | 0.17 | 32,32,32,32 | 0 |
| 57 | MG | 1a | 1775 | 1/1 | 0.90 | 0.09 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3460 | 1/1 | 0.90 | 0.26 | 45,45,45,45 | 0 |
| 57 | MG | 1a | 1777 | 1/1 | 0.90 | 0.10 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3195 | 1/1 | 0.90 | 0.16 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3730 | 1/1 | 0.90 | 0.13 | 71,71,71,71 | 0 |
| 57 | MG | 2a | 1610 | 1/1 | 0.90 | 0.12 | 48,48,48,48 | 0 |
| 57 | MG | 2a | 1612 | 1/1 | 0.90 | 0.12 | 68,68,68,68 | 0 |
| 57 | MG | 2A | 3470 | 1/1 | 0.90 | 0.16 | 64,64,64,64 | 0 |
| 57 | MG | 2A | 3472 | 1/1 | 0.90 | 0.06 | 63,63,63,63 | 0 |
| 57 | MG | 2A | 3483 | 1/1 | 0.90 | 0.14 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3739 | 1/1 | 0.90 | 0.10 | 57,57,57,57 | 0 |
| 57 | MG | 2a | 1622 | 1/1 | 0.90 | 0.11 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3957 | 1/1 | 0.90 | 0.15 | 61,61,61,61 | 0 |
| 57 | MG | 1A | 3199 | 1/1 | 0.90 | 0.39 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3226 | 1/1 | 0.90 | 0.21 | 71,71,71,71 | 0 |
| 57 | MG | 2A | 3227 | 1/1 | 0.90 | 0.15 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3231 | 1/1 | 0.90 | 0.10 | 34,34,34,34 | 0 |
| 57 | MG | 2a | 1632 | 1/1 | 0.90 | 0.05 | 78,78,78,78 | 0 |
| 57 | MG | 2A | 3527 | 1/1 | 0.90 | 0.08 | 40,40,40,40 | 0 |
| 57 | MG | 1a | 1796 | 1/1 | 0.90 | 0.19 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3543 | 1/1 | 0.90 | 0.09 | 44,44,44,44 | 0 |
| 57 | MG | 1a | 1797 | 1/1 | 0.90 | 0.07 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3556 | 1/1 | 0.90 | 0.13 | 50,50,50,50 | 0 |
| 57 | MG | 15 | 106 | 1/1 | 0.90 | 0.21 | 44,44,44,44 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 15 | 109 | 1/1 | 0.90 | 0.10 | 62,62,62,62 | 0 |
| 57 | MG | 1A | 3962 | 1/1 | 0.90 | 0.10 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3255 | 1/1 | 0.90 | 0.28 | 72,72,72,72 | 0 |
| 57 | MG | 1a | 1602 | 1/1 | 0.90 | 0.15 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3793 | 1/1 | 0.90 | 0.18 | 70,70,70,70 | 0 |
| 57 | MG | 2A | 3586 | 1/1 | 0.90 | 0.16 | 58,58,58,58 | 0 |
| 57 | MG | 1f | 201 | 1/1 | 0.90 | 0.12 | 54,54,54,54 | 0 |
| 57 | MG | 1l | 201 | 1/1 | 0.90 | 0.07 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3058 | 1/1 | 0.90 | 0.13 | 41,41,41,41 | 0 |
| 57 | MG | 1t | 201 | 1/1 | 0.90 | 0.17 | 59,59,59,59 | 0 |
| 57 | MG | 1A | 3988 | 1/1 | 0.90 | 0.15 | 65,65,65,65 | 0 |
| 57 | MG | 1A | 3810 | 1/1 | 0.90 | 0.14 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3379 | 1/1 | 0.90 | 0.23 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3814 | 1/1 | 0.90 | 0.13 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3519 | 1/1 | 0.90 | 0.17 | 40,40,40,40 | 0 |
| 57 | MG | 2a | 1674 | 1/1 | 0.90 | 0.27 | 69,69,69,69 | 0 |
| 57 | MG | 2a | 1675 | 1/1 | 0.90 | 0.21 | 61,61,61,61 | 0 |
| 57 | MG | 1a | 1629 | 1/1 | 0.90 | 0.14 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3001 | 1/1 | 0.90 | 0.19 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3646 | 1/1 | 0.90 | 0.16 | 59,59,59,59 | 0 |
| 57 | MG | 1A | 3171 | 1/1 | 0.90 | 0.13 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3400 | 1/1 | 0.90 | 0.09 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3028 | 1/1 | 0.90 | 0.15 | 54,54,54,54 | 0 |
| 57 | MG | 2a | 1690 | 1/1 | 0.90 | 0.18 | 63,63,63,63 | 0 |
| 57 | MG | 2A | 3658 | 1/1 | 0.90 | 0.08 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3031 | 1/1 | 0.90 | 0.14 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3670 | 1/1 | 0.90 | 0.20 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3035 | 1/1 | 0.90 | 0.11 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3264 | 1/1 | 0.90 | 0.16 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3404 | 1/1 | 0.90 | 0.16 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3296 | 1/1 | 0.90 | 0.13 | 62,62,62,62 | 0 |
| 57 | MG | 2a | 1708 | 1/1 | 0.90 | 0.33 | 69,69,69,69 | 0 |
| 57 | MG | 2A | 3045 | 1/1 | 0.90 | 0.15 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 4067 | 1/1 | 0.90 | 0.14 | 63,63,63,63 | 0 |
| 57 | MG | 1A | 3277 | 1/1 | 0.90 | 0.17 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3422 | 1/1 | 0.90 | 0.20 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 4072 | 1/1 | 0.90 | 0.09 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3582 | 1/1 | 0.90 | 0.30 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3584 | 1/1 | 0.90 | 0.14 | 61,61,61,61 | 0 |
| 57 | MG | 1a | 1668 | 1/1 | 0.90 | 0.07 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3083 | 1/1 | 0.90 | 0.10 | 52,52,52,52 | 0 |
| 57 | MG | 1a | 1670 | 1/1 | 0.90 | 0.31 | 74,74,74,74 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3087 | 1/1 | 0.90 | 0.08 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3286 | 1/1 | 0.90 | 0.17 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3223 | 1/1 | 0.90 | 0.11 | 48,48,48,48 | 0 |
| 57 | MG | 1a | 1677 | 1/1 | 0.90 | 0.19 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3726 | 1/1 | 0.90 | 0.11 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3095 | 1/1 | 0.90 | 0.09 | 51,51,51,51 | 0 |
| 57 | MG | 1B | 201 | 1/1 | 0.90 | 0.07 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3098 | 1/1 | 0.90 | 0.26 | 67,67,67,67 | 0 |
| 57 | MG | 2A | 3335 | 1/1 | 0.90 | 0.11 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3598 | 1/1 | 0.90 | 0.12 | 41,41,41,41 | 0 |
| 57 | MG | 1a | 1684 | 1/1 | 0.90 | 0.07 | 50,50,50,50 | 0 |
| 57 | MG | 1B | 208 | 1/1 | 0.90 | 0.08 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3435 | 1/1 | 0.90 | 0.11 | 48,48,48,48 | 0 |
| 57 | MG | 1a | 1698 | 1/1 | 0.90 | 0.10 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3876 | 1/1 | 0.90 | 0.17 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3357 | 1/1 | 0.90 | 0.26 | 63,63,63,63 | 0 |
| 57 | MG | 2A | 3776 | 1/1 | 0.90 | 0.20 | 65,65,65,65 | 0 |
| 57 | MG | 2A | 3116 | 1/1 | 0.90 | 0.16 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3600 | 1/1 | 0.90 | 0.21 | 43,43,43,43 | 0 |
| 57 | MG | 2a | 1775 | 1/1 | 0.90 | 0.18 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3438 | 1/1 | 0.90 | 0.13 | 63,63,63,63 | 0 |
| 57 | MG | 1A | 3149 | 1/1 | 0.90 | 0.13 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3791 | 1/1 | 0.90 | 0.08 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3446 | 1/1 | 0.90 | 0.08 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3801 | 1/1 | 0.90 | 0.11 | 40,40,40,40 | 0 |
| 57 | MG | 2A | 3373 | 1/1 | 0.90 | 0.12 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3232 | 1/1 | 0.90 | 0.36 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3304 | 1/1 | 0.90 | 0.22 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3459 | 1/1 | 0.90 | 0.20 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3833 | 1/1 | 0.90 | 0.08 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3164 | 1/1 | 0.90 | 0.18 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3463 | 1/1 | 0.90 | 0.32 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3927 | 1/1 | 0.90 | 0.21 | 48,48,48,48 | 0 |
| 57 | MG | 1a | 1725 | 1/1 | 0.90 | 0.17 | 61,61,61,61 | 0 |
| 57 | MG | 1A | 3932 | 1/1 | 0.90 | 0.10 | 23,23,23,23 | 0 |
| 57 | MG | 2A | 3846 | 1/1 | 0.90 | 0.07 | 60,60,60,60 | 0 |
| 57 | MG | 2a | 1811 | 1/1 | 0.90 | 0.21 | 73,73,73,73 | 0 |
| 57 | MG | 2A | 3388 | 1/1 | 0.90 | 0.14 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3241 | 1/1 | 0.90 | 0.14 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3396 | 1/1 | 0.90 | 0.17 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3403 | 1/1 | 0.90 | 0.24 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3934 | 1/1 | 0.90 | 0.15 | 58,58,58,58 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1a | 1742 | 1/1 | 0.90 | 0.23 | 68,68,68,68 | 0 |
| 57 | MG | 2g | 201 | 1/1 | 0.90 | 0.10 | 67,67,67,67 | 0 |
| 57 | MG | 2B | 211 | 1/1 | 0.90 | 0.22 | 64,64,64,64 | 0 |
| 57 | MG | 2B | 212 | 1/1 | 0.90 | 0.24 | 61,61,61,61 | 0 |
| 57 | MG | 2l | 204 | 1/1 | 0.90 | 0.05 | 51,51,51,51 | 0 |
| 57 | MG | 2B | 213 | 1/1 | 0.90 | 0.14 | 56,56,56,56 | 0 |
| 57 | MG | 1a | 1747 | 1/1 | 0.90 | 0.09 | 59,59,59,59 | 0 |
| 57 | MG | 1a | 1757 | 1/1 | 0.90 | 0.15 | 54,54,54,54 | 0 |
| 57 | MG | 1F | 312 | 1/1 | 0.90 | 0.08 | 53,53,53,53 | 0 |
| 57 | MG | 2w | 102 | 1/1 | 0.90 | 0.23 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3429 | 1/1 | 0.90 | 0.23 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3324 | 1/1 | 0.90 | 0.07 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3479 | 1/1 | 0.90 | 0.23 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3435 | 1/1 | 0.90 | 0.25 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3436 | 1/1 | 0.90 | 0.11 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3940 | 1/1 | 0.91 | 0.10 | 55,55,55,55 | 0 |
| 57 | MG | 1a | 1735 | 1/1 | 0.91 | 0.11 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3303 | 1/1 | 0.91 | 0.21 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3202 | 1/1 | 0.91 | 0.08 | 26,26,26,26 | 0 |
| 57 | MG | 1O | 202 | 1/1 | 0.91 | 0.20 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3854 | 1/1 | 0.91 | 0.10 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3166 | 1/1 | 0.91 | 0.15 | 67,67,67,67 | 0 |
| 57 | MG | 1A | 3406 | 1/1 | 0.91 | 0.16 | 55,55,55,55 | 0 |
| 57 | MG | 2B | 205 | 1/1 | 0.91 | 0.06 | 55,55,55,55 | 0 |
| 57 | MG | 2B | 206 | 1/1 | 0.91 | 0.19 | 63,63,63,63 | 0 |
| 57 | MG | 1A | 3319 | 1/1 | 0.91 | 0.06 | 39,39,39,39 | 0 |
| 57 | MG | 2A | 3421 | 1/1 | 0.91 | 0.20 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3520 | 1/1 | 0.91 | 0.26 | 57,57,57,57 | 0 |
| 57 | MG | 1S | 201 | 1/1 | 0.91 | 0.08 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3177 | 1/1 | 0.91 | 0.22 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3762 | 1/1 | 0.91 | 0.08 | 24,24,24,24 | 0 |
| 57 | MG | 1A | 3770 | 1/1 | 0.91 | 0.12 | 31,31,31,31 | 0 |
| 57 | MG | 10 | 107 | 1/1 | 0.91 | 0.09 | 50,50,50,50 | 0 |
| 57 | MG | 10 | 108 | 1/1 | 0.91 | 0.13 | 53,53,53,53 | 0 |
| 57 | MG | 1a | 1774 | 1/1 | 0.91 | 0.10 | 36,36,36,36 | 0 |
| 57 | MG | 2A | 3442 | 1/1 | 0.91 | 0.07 | 60,60,60,60 | 0 |
| 57 | MG | 2N | 201 | 1/1 | 0.91 | 0.23 | 65,65,65,65 | 0 |
| 57 | MG | 10 | 110 | 1/1 | 0.91 | 0.21 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3449 | 1/1 | 0.91 | 0.14 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3452 | 1/1 | 0.91 | 0.44 | 57,57,57,57 | 0 |
| 57 | MG | 11 | 103 | 1/1 | 0.91 | 0.12 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3455 | 1/1 | 0.91 | 0.11 | 60,60,60,60 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 3415 | 1/1 | 0.91 | 0.12 | 56,56,56,56 | 0 |
| 57 | MG | 23 | 101 | 1/1 | 0.91 | 0.15 | 56,56,56,56 | 0 |
| 57 | MG | 28 | 101 | 1/1 | 0.91 | 0.12 | 62,62,62,62 | 0 |
| 57 | MG | 2a | 1602 | 1/1 | 0.91 | 0.17 | 55,55,55,55 | 0 |
| 57 | MG | 2a | 1603 | 1/1 | 0.91 | 0.15 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3459 | 1/1 | 0.91 | 0.13 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3964 | 1/1 | 0.91 | 0.08 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3531 | 1/1 | 0.91 | 0.09 | 68,68,68,68 | 0 |
| 57 | MG | 1A | 3801 | 1/1 | 0.91 | 0.13 | 44,44,44,44 | 0 |
| 57 | MG | 1a | 1794 | 1/1 | 0.91 | 0.08 | 77,77,77,77 | 0 |
| 57 | MG | 1A | 3532 | 1/1 | 0.91 | 0.09 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3208 | 1/1 | 0.91 | 0.17 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3479 | 1/1 | 0.91 | 0.24 | 48,48,48,48 | 0 |
| 57 | MG | 2a | 1619 | 1/1 | 0.91 | 0.19 | 64,64,64,64 | 0 |
| 57 | MG | 1A | 3114 | 1/1 | 0.91 | 0.09 | 41,41,41,41 | 0 |
| 57 | MG | 18 | 106 | 1/1 | 0.91 | 0.26 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3216 | 1/1 | 0.91 | 0.12 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3488 | 1/1 | 0.91 | 0.09 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3115 | 1/1 | 0.91 | 0.09 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3223 | 1/1 | 0.91 | 0.29 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3424 | 1/1 | 0.91 | 0.09 | 52,52,52,52 | 0 |
| 57 | MG | 2a | 1630 | 1/1 | 0.91 | 0.20 | 64,64,64,64 | 0 |
| 57 | MG | 2a | 1631 | 1/1 | 0.91 | 0.10 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3504 | 1/1 | 0.91 | 0.17 | 61,61,61,61 | 0 |
| 57 | MG | 2A | 3505 | 1/1 | 0.91 | 0.18 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3996 | 1/1 | 0.91 | 0.08 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3516 | 1/1 | 0.91 | 0.20 | 65,65,65,65 | 0 |
| 57 | MG | 2A | 3520 | 1/1 | 0.91 | 0.11 | 62,62,62,62 | 0 |
| 57 | MG | 1a | 1802 | 1/1 | 0.91 | 0.10 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3529 | 1/1 | 0.91 | 0.11 | 62,62,62,62 | 0 |
| 57 | MG | 1A | 3999 | 1/1 | 0.91 | 0.10 | 44,44,44,44 | 0 |
| 57 | MG | 2a | 1640 | 1/1 | 0.91 | 0.16 | 58,58,58,58 | 0 |
| 57 | MG | 2a | 1641 | 1/1 | 0.91 | 0.08 | 59,59,59,59 | 0 |
| 57 | MG | 2A | 3534 | 1/1 | 0.91 | 0.10 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3536 | 1/1 | 0.91 | 0.08 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3004 | 1/1 | 0.91 | 0.08 | 31,31,31,31 | 0 |
| 57 | MG | 2A | 3234 | 1/1 | 0.91 | 0.16 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3575 | 1/1 | 0.91 | 0.10 | 37,37,37,37 | 0 |
| 57 | MG | 2a | 1651 | 1/1 | 0.91 | 0.20 | 67,67,67,67 | 0 |
| 57 | MG | 2a | 1652 | 1/1 | 0.91 | 0.06 | 70,70,70,70 | 0 |
| 57 | MG | 2A | 3563 | 1/1 | 0.91 | 0.09 | 38,38,38,38 | 0 |
| 57 | MG | 2a | 1655 | 1/1 | 0.91 | 0.07 | 46,46,46,46 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3565 | 1/1 | 0.91 | 0.08 | 37,37,37,37 | 0 |
| 57 | MG | 2a | 1658 | 1/1 | 0.91 | 0.17 | 50,50,50,50 | 0 |
| 57 | MG | 1a | 1809 | 1/1 | 0.91 | 0.08 | 65,65,65,65 | 0 |
| 57 | MG | 1A | 4016 | 1/1 | 0.91 | 0.12 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3329 | 1/1 | 0.91 | 0.12 | 56,56,56,56 | 0 |
| 57 | MG | 1n | 102 | 1/1 | 0.91 | 0.17 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3261 | 1/1 | 0.91 | 0.15 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3578 | 1/1 | 0.91 | 0.09 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3030 | 1/1 | 0.91 | 0.30 | 64,64,64,64 | 0 |
| 57 | MG | 1a | 1633 | 1/1 | 0.91 | 0.13 | 35,35,35,35 | 0 |
| 57 | MG | 2A | 3595 | 1/1 | 0.91 | 0.13 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3247 | 1/1 | 0.91 | 0.12 | 48,48,48,48 | 0 |
| 57 | MG | 1a | 1636 | 1/1 | 0.91 | 0.10 | 40,40,40,40 | 0 |
| 57 | MG | 1a | 1637 | 1/1 | 0.91 | 0.22 | 55,55,55,55 | 0 |
| 57 | MG | 2a | 1680 | 1/1 | 0.91 | 0.24 | 70,70,70,70 | 0 |
| 57 | MG | 1A | 3267 | 1/1 | 0.91 | 0.06 | 53,53,53,53 | 0 |
| 57 | MG | 1x | 102 | 1/1 | 0.91 | 0.15 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3609 | 1/1 | 0.91 | 0.11 | 50,50,50,50 | 0 |
| 57 | MG | 1a | 1644 | 1/1 | 0.91 | 0.12 | 43,43,43,43 | 0 |
| 57 | MG | 2a | 1694 | 1/1 | 0.91 | 0.12 | 66,66,66,66 | 0 |
| 57 | MG | 1A | 4039 | 1/1 | 0.91 | 0.11 | 63,63,63,63 | 0 |
| 57 | MG | 1A | 4044 | 1/1 | 0.91 | 0.11 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3623 | 1/1 | 0.91 | 0.10 | 36,36,36,36 | 0 |
| 57 | MG | 2a | 1702 | 1/1 | 0.91 | 0.20 | 68,68,68,68 | 0 |
| 57 | MG | 1A | 4058 | 1/1 | 0.91 | 0.12 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3448 | 1/1 | 0.91 | 0.10 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3147 | 1/1 | 0.91 | 0.14 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3029 | 1/1 | 0.91 | 0.10 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3030 | 1/1 | 0.91 | 0.09 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3353 | 1/1 | 0.91 | 0.22 | 35,35,35,35 | 0 |
| 57 | MG | 2a | 1710 | 1/1 | 0.91 | 0.14 | 59,59,59,59 | 0 |
| 57 | MG | 1A | 3361 | 1/1 | 0.91 | 0.18 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3290 | 1/1 | 0.91 | 0.08 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3038 | 1/1 | 0.91 | 0.24 | 66,66,66,66 | 0 |
| 57 | MG | 1A | 3631 | 1/1 | 0.91 | 0.15 | 54,54,54,54 | 0 |
| 57 | MG | 1a | 1666 | 1/1 | 0.91 | 0.22 | 63,63,63,63 | 0 |
| 57 | MG | 1a | 1667 | 1/1 | 0.91 | 0.18 | 59,59,59,59 | 0 |
| 57 | MG | 2A | 3300 | 1/1 | 0.91 | 0.21 | 64,64,64,64 | 0 |
| 57 | MG | 2A | 3675 | 1/1 | 0.91 | 0.20 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3047 | 1/1 | 0.91 | 0.10 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3050 | 1/1 | 0.91 | 0.21 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3634 | 1/1 | 0.91 | 0.07 | 39,39,39,39 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3681 | 1/1 | 0.91 | 0.20 | 59,59,59,59 | 0 |
| 57 | MG | 2A | 3304 | 1/1 | 0.91 | 0.17 | 60,60,60,60 | 0 |
| 57 | MG | 2a | 1734 | 1/1 | 0.91 | 0.34 | 72,72,72,72 | 0 |
| 57 | MG | 1A | 3639 | 1/1 | 0.91 | 0.18 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3461 | 1/1 | 0.91 | 0.14 | 40,40,40,40 | 0 |
| 57 | MG | 2A | 3059 | 1/1 | 0.91 | 0.16 | 56,56,56,56 | 0 |
| 57 | MG | 2a | 1741 | 1/1 | 0.91 | 0.32 | 60,60,60,60 | 0 |
| 57 | MG | 2a | 1742 | 1/1 | 0.91 | 0.14 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3314 | 1/1 | 0.91 | 0.14 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3693 | 1/1 | 0.91 | 0.07 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3370 | 1/1 | 0.91 | 0.15 | 43,43,43,43 | 0 |
| 57 | MG | 2a | 1748 | 1/1 | 0.91 | 0.26 | 68,68,68,68 | 0 |
| 57 | MG | 2A | 3700 | 1/1 | 0.91 | 0.11 | 61,61,61,61 | 0 |
| 57 | MG | 2A | 3316 | 1/1 | 0.91 | 0.09 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3148 | 1/1 | 0.91 | 0.33 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3470 | 1/1 | 0.91 | 0.10 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3070 | 1/1 | 0.91 | 0.10 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3890 | 1/1 | 0.91 | 0.12 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3079 | 1/1 | 0.91 | 0.10 | 49,49,49,49 | 0 |
| 57 | MG | 2a | 1765 | 1/1 | 0.91 | 0.14 | 58,58,58,58 | 0 |
| 57 | MG | 2a | 1766 | 1/1 | 0.91 | 0.13 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3055 | 1/1 | 0.91 | 0.09 | 38,38,38,38 | 0 |
| 57 | MG | 2a | 1769 | 1/1 | 0.91 | 0.24 | 71,71,71,71 | 0 |
| 57 | MG | 2A | 3727 | 1/1 | 0.91 | 0.22 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3730 | 1/1 | 0.91 | 0.16 | 68,68,68,68 | 0 |
| 57 | MG | 1a | 1686 | 1/1 | 0.91 | 0.18 | 35,35,35,35 | 0 |
| 57 | MG | 2a | 1782 | 1/1 | 0.91 | 0.13 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3898 | 1/1 | 0.91 | 0.14 | 62,62,62,62 | 0 |
| 57 | MG | 2A | 3088 | 1/1 | 0.91 | 0.23 | 63,63,63,63 | 0 |
| 57 | MG | 1a | 1689 | 1/1 | 0.91 | 0.20 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3665 | 1/1 | 0.91 | 0.09 | 35,35,35,35 | 0 |
| 57 | MG | 2A | 3749 | 1/1 | 0.91 | 0.11 | 53,53,53,53 | 0 |
| 57 | MG | 1a | 1695 | 1/1 | 0.91 | 0.23 | 59,59,59,59 | 0 |
| 57 | MG | 2a | 1791 | 1/1 | 0.91 | 0.22 | 60,60,60,60 | 0 |
| 57 | MG | 2a | 1794 | 1/1 | 0.91 | 0.24 | 69,69,69,69 | 0 |
| 57 | MG | 1A | 3380 | 1/1 | 0.91 | 0.23 | 38,38,38,38 | 0 |
| 57 | MG | 1B | 218 | 1/1 | 0.91 | 0.12 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3351 | 1/1 | 0.91 | 0.18 | 55,55,55,55 | 0 |
| 57 | MG | 2a | 1798 | 1/1 | 0.91 | 0.16 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3353 | 1/1 | 0.91 | 0.07 | 63,63,63,63 | 0 |
| 57 | MG | 1a | 1705 | 1/1 | 0.91 | 0.14 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3480 | 1/1 | 0.91 | 0.08 | 37,37,37,37 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2a | 1804 | 1/1 | 0.91 | 0.09 | 64,64,64,64 | 0 |
| 57 | MG | 2A | 3779 | 1/1 | 0.91 | 0.11 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3782 | 1/1 | 0.91 | 0.20 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3690 | 1/1 | 0.91 | 0.06 | 45,45,45,45 | 0 |
| 57 | MG | 1a | 1710 | 1/1 | 0.91 | 0.23 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3361 | 1/1 | 0.91 | 0.16 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3364 | 1/1 | 0.91 | 0.20 | 51,51,51,51 | 0 |
| 57 | MG | 2a | 1817 | 1/1 | 0.91 | 0.12 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3923 | 1/1 | 0.91 | 0.09 | 42,42,42,42 | 0 |
| 57 | MG | 2A | 3793 | 1/1 | 0.91 | 0.10 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3108 | 1/1 | 0.91 | 0.11 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3798 | 1/1 | 0.91 | 0.17 | 43,43,43,43 | 0 |
| 57 | MG | 2e | 201 | 1/1 | 0.91 | 0.07 | 82,82,82,82 | 0 |
| 57 | MG | 2A | 3800 | 1/1 | 0.91 | 0.11 | 63,63,63,63 | 0 |
| 57 | MG | 1a | 1714 | 1/1 | 0.91 | 0.12 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3802 | 1/1 | 0.91 | 0.07 | 28,28,28,28 | 0 |
| 57 | MG | 2l | 201 | 1/1 | 0.91 | 0.11 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3817 | 1/1 | 0.91 | 0.10 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3383 | 1/1 | 0.91 | 0.22 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3113 | 1/1 | 0.91 | 0.16 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3115 | 1/1 | 0.91 | 0.11 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3388 | 1/1 | 0.91 | 0.20 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3706 | 1/1 | 0.91 | 0.15 | 25,25,25,25 | 0 |
| 57 | MG | 2w | 104 | 1/1 | 0.91 | 0.09 | 89,89,89,89 | 0 |
| 57 | MG | 2A | 3122 | 1/1 | 0.91 | 0.25 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3708 | 1/1 | 0.91 | 0.08 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3248 | 1/1 | 0.91 | 0.14 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3201 | 1/1 | 0.91 | 0.07 | 32,32,32,32 | 0 |
| 57 | MG | 2x | 104 | 1/1 | 0.91 | 0.28 | 65,65,65,65 | 0 |
| 57 | MG | 2x | 105 | 1/1 | 0.91 | 0.17 | 60,60,60,60 | 0 |
| 57 | MG | 2A | 3841 | 1/1 | 0.91 | 0.13 | 65,65,65,65 | 0 |
| 58 | K | 1A | 3546 | 1/1 | 0.91 | 0.20 | 73,73,73,73 | 0 |
| 58 | K | 2A | 3445 | 1/1 | 0.91 | 0.08 | 64,64,64,64 | 0 |
| 57 | MG | 2B | 202 | 1/1 | 0.92 | 0.08 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3399 | 1/1 | 0.92 | 0.15 | 45,45,45,45 | 0 |
| 57 | MG | 1B | 206 | 1/1 | 0.92 | 0.07 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3565 | 1/1 | 0.92 | 0.14 | 42,42,42,42 | 0 |
| 57 | MG | 2A | 3414 | 1/1 | 0.92 | 0.25 | 53,53,53,53 | 0 |
| 57 | MG | 2B | 207 | 1/1 | 0.92 | 0.15 | 68,68,68,68 | 0 |
| 57 | MG | 1A | 3338 | 1/1 | 0.92 | 0.10 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3339 | 1/1 | 0.92 | 0.08 | 40,40,40,40 | 0 |
| 57 | MG | 2A | 3139 | 1/1 | 0.92 | 0.10 | 56,56,56,56 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3147 | 1/1 | 0.92 | 0.19 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3148 | 1/1 | 0.92 | 0.16 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3428 | 1/1 | 0.92 | 0.29 | 62,62,62,62 | 0 |
| 57 | MG | 1A | 3124 | 1/1 | 0.92 | 0.10 | 41,41,41,41 | 0 |
| 57 | MG | 2B | 218 | 1/1 | 0.92 | 0.16 | 67,67,67,67 | 0 |
| 57 | MG | 1A | 3854 | 1/1 | 0.92 | 0.10 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3581 | 1/1 | 0.92 | 0.16 | 57,57,57,57 | 0 |
| 57 | MG | 2D | 306 | 1/1 | 0.92 | 0.09 | 63,63,63,63 | 0 |
| 57 | MG | 2E | 304 | 1/1 | 0.92 | 0.20 | 43,43,43,43 | 0 |
| 57 | MG | 2E | 305 | 1/1 | 0.92 | 0.08 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3341 | 1/1 | 0.92 | 0.14 | 40,40,40,40 | 0 |
| 57 | MG | 2F | 301 | 1/1 | 0.92 | 0.07 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3160 | 1/1 | 0.92 | 0.14 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3436 | 1/1 | 0.92 | 0.11 | 41,41,41,41 | 0 |
| 57 | MG | 2O | 201 | 1/1 | 0.92 | 0.16 | 64,64,64,64 | 0 |
| 57 | MG | 2O | 202 | 1/1 | 0.92 | 0.14 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3441 | 1/1 | 0.92 | 0.16 | 61,61,61,61 | 0 |
| 57 | MG | 1B | 228 | 1/1 | 0.92 | 0.09 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3299 | 1/1 | 0.92 | 0.26 | 64,64,64,64 | 0 |
| 57 | MG | 2W | 201 | 1/1 | 0.92 | 0.14 | 48,48,48,48 | 0 |
| 57 | MG | 1a | 1730 | 1/1 | 0.92 | 0.08 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3586 | 1/1 | 0.92 | 0.21 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3171 | 1/1 | 0.92 | 0.11 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3587 | 1/1 | 0.92 | 0.17 | 50,50,50,50 | 0 |
| 57 | MG | 27 | 102 | 1/1 | 0.92 | 0.10 | 53,53,53,53 | 0 |
| 57 | MG | 27 | 103 | 1/1 | 0.92 | 0.42 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3588 | 1/1 | 0.92 | 0.19 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3074 | 1/1 | 0.92 | 0.22 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3178 | 1/1 | 0.92 | 0.13 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3465 | 1/1 | 0.92 | 0.10 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3354 | 1/1 | 0.92 | 0.17 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3002 | 1/1 | 0.92 | 0.09 | 47,47,47,47 | 0 |
| 57 | MG | 2a | 1607 | 1/1 | 0.92 | 0.15 | 59,59,59,59 | 0 |
| 57 | MG | 2A | 3182 | 1/1 | 0.92 | 0.21 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3887 | 1/1 | 0.92 | 0.07 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3185 | 1/1 | 0.92 | 0.21 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3186 | 1/1 | 0.92 | 0.09 | 58,58,58,58 | 0 |
| 57 | MG | 2a | 1615 | 1/1 | 0.92 | 0.11 | 58,58,58,58 | 0 |
| 57 | MG | 1a | 1758 | 1/1 | 0.92 | 0.16 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3362 | 1/1 | 0.92 | 0.37 | 53,53,53,53 | 0 |
| 57 | MG | 2a | 1620 | 1/1 | 0.92 | 0.15 | 65,65,65,65 | 0 |
| 57 | MG | 1A | 3603 | 1/1 | 0.92 | 0.42 | 66,66,66,66 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 3611 | 1/1 | 0.92 | 0.09 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3490 | 1/1 | 0.92 | 0.07 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3626 | 1/1 | 0.92 | 0.12 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3198 | 1/1 | 0.92 | 0.11 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3496 | 1/1 | 0.92 | 0.18 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3498 | 1/1 | 0.92 | 0.09 | 71,71,71,71 | 0 |
| 57 | MG | 1P | 205 | 1/1 | 0.92 | 0.15 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3907 | 1/1 | 0.92 | 0.09 | 24,24,24,24 | 0 |
| 57 | MG | 1A | 3456 | 1/1 | 0.92 | 0.08 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3915 | 1/1 | 0.92 | 0.09 | 60,60,60,60 | 0 |
| 57 | MG | 1R | 206 | 1/1 | 0.92 | 0.07 | 32,32,32,32 | 0 |
| 57 | MG | 1a | 1779 | 1/1 | 0.92 | 0.18 | 63,63,63,63 | 0 |
| 57 | MG | 1A | 3366 | 1/1 | 0.92 | 0.06 | 32,32,32,32 | 0 |
| 57 | MG | 2A | 3214 | 1/1 | 0.92 | 0.13 | 41,41,41,41 | 0 |
| 57 | MG | 1a | 1783 | 1/1 | 0.92 | 0.12 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3533 | 1/1 | 0.92 | 0.12 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3001 | 1/1 | 0.92 | 0.07 | 39,39,39,39 | 0 |
| 57 | MG | 2A | 3218 | 1/1 | 0.92 | 0.12 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3541 | 1/1 | 0.92 | 0.14 | 44,44,44,44 | 0 |
| 57 | MG | 2a | 1644 | 1/1 | 0.92 | 0.07 | 63,63,63,63 | 0 |
| 57 | MG | 1X | 106 | 1/1 | 0.92 | 0.08 | 53,53,53,53 | 0 |
| 57 | MG | 1a | 1792 | 1/1 | 0.92 | 0.15 | 74,74,74,74 | 0 |
| 57 | MG | 1A | 3041 | 1/1 | 0.92 | 0.13 | 49,49,49,49 | 0 |
| 57 | MG | 10 | 104 | 1/1 | 0.92 | 0.12 | 61,61,61,61 | 0 |
| 57 | MG | 2a | 1650 | 1/1 | 0.92 | 0.13 | 73,73,73,73 | 0 |
| 57 | MG | 2A | 3229 | 1/1 | 0.92 | 0.13 | 57,57,57,57 | 0 |
| 57 | MG | 10 | 106 | 1/1 | 0.92 | 0.09 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3372 | 1/1 | 0.92 | 0.10 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3260 | 1/1 | 0.92 | 0.11 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3305 | 1/1 | 0.92 | 0.23 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3307 | 1/1 | 0.92 | 0.13 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3661 | 1/1 | 0.92 | 0.12 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3662 | 1/1 | 0.92 | 0.08 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3475 | 1/1 | 0.92 | 0.24 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3477 | 1/1 | 0.92 | 0.09 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3067 | 1/1 | 0.92 | 0.08 | 30,30,30,30 | 0 |
| 57 | MG | 2a | 1669 | 1/1 | 0.92 | 0.23 | 54,54,54,54 | 0 |
| 57 | MG | 1b | 301 | 1/1 | 0.92 | 0.14 | 70,70,70,70 | 0 |
| 57 | MG | 17 | 105 | 1/1 | 0.92 | 0.26 | 55,55,55,55 | 0 |
| 57 | MG | 18 | 104 | 1/1 | 0.92 | 0.18 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3389 | 1/1 | 0.92 | 0.13 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3605 | 1/1 | 0.92 | 0.07 | 68,68,68,68 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 3263 | 1/1 | 0.92 | 0.08 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3610 | 1/1 | 0.92 | 0.09 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3951 | 1/1 | 0.92 | 0.07 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 3397 | 1/1 | 0.92 | 0.22 | 60,60,60,60 | 0 |
| 57 | MG | 2A | 3618 | 1/1 | 0.92 | 0.09 | 41,41,41,41 | 0 |
| 57 | MG | 1a | 1613 | 1/1 | 0.92 | 0.20 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3068 | 1/1 | 0.92 | 0.23 | 51,51,51,51 | 0 |
| 57 | MG | 2a | 1691 | 1/1 | 0.92 | 0.15 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3636 | 1/1 | 0.92 | 0.09 | 38,38,38,38 | 0 |
| 57 | MG | 1A | 3166 | 1/1 | 0.92 | 0.12 | 41,41,41,41 | 0 |
| 57 | MG | 1x | 101 | 1/1 | 0.92 | 0.27 | 65,65,65,65 | 0 |
| 57 | MG | 1A | 3710 | 1/1 | 0.92 | 0.09 | 21,21,21,21 | 0 |
| 57 | MG | 2A | 3267 | 1/1 | 0.92 | 0.12 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3269 | 1/1 | 0.92 | 0.22 | 61,61,61,61 | 0 |
| 57 | MG | 2a | 1704 | 1/1 | 0.92 | 0.10 | 80,80,80,80 | 0 |
| 57 | MG | 1A | 3713 | 1/1 | 0.92 | 0.10 | 40,40,40,40 | 0 |
| 57 | MG | 2A | 3274 | 1/1 | 0.92 | 0.09 | 80,80,80,80 | 0 |
| 57 | MG | 1A | 3327 | 1/1 | 0.92 | 0.10 | 32,32,32,32 | 0 |
| 57 | MG | 2A | 3662 | 1/1 | 0.92 | 0.13 | 73,73,73,73 | 0 |
| 57 | MG | 1x | 112 | 1/1 | 0.92 | 0.24 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3506 | 1/1 | 0.92 | 0.09 | 65,65,65,65 | 0 |
| 57 | MG | 2A | 3284 | 1/1 | 0.92 | 0.22 | 67,67,67,67 | 0 |
| 57 | MG | 2A | 3285 | 1/1 | 0.92 | 0.20 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3287 | 1/1 | 0.92 | 0.11 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3003 | 1/1 | 0.92 | 0.28 | 59,59,59,59 | 0 |
| 57 | MG | 2A | 3005 | 1/1 | 0.92 | 0.16 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3015 | 1/1 | 0.92 | 0.26 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3719 | 1/1 | 0.92 | 0.09 | 21,21,21,21 | 0 |
| 57 | MG | 1A | 3328 | 1/1 | 0.92 | 0.19 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3298 | 1/1 | 0.92 | 0.16 | 64,64,64,64 | 0 |
| 57 | MG | 2A | 3683 | 1/1 | 0.92 | 0.13 | 32,32,32,32 | 0 |
| 57 | MG | 2A | 3023 | 1/1 | 0.92 | 0.20 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3685 | 1/1 | 0.92 | 0.14 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3686 | 1/1 | 0.92 | 0.15 | 77,77,77,77 | 0 |
| 57 | MG | 2A | 3688 | 1/1 | 0.92 | 0.20 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3409 | 1/1 | 0.92 | 0.23 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3410 | 1/1 | 0.92 | 0.08 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3249 | 1/1 | 0.92 | 0.09 | 49,49,49,49 | 0 |
| 57 | MG | 2a | 1739 | 1/1 | 0.92 | 0.24 | 74,74,74,74 | 0 |
| 57 | MG | 1A | 3280 | 1/1 | 0.92 | 0.13 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3528 | 1/1 | 0.92 | 0.12 | 54,54,54,54 | 0 |
| 57 | MG | 1a | 1646 | 1/1 | 0.92 | 0.16 | 55,55,55,55 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3705 | 1/1 | 0.92 | 0.09 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 4002 | 1/1 | 0.92 | 0.07 | 67,67,67,67 | 0 |
| 57 | MG | 2A | 3711 | 1/1 | 0.92 | 0.13 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3307 | 1/1 | 0.92 | 0.17 | 39,39,39,39 | 0 |
| 57 | MG | 2a | 1752 | 1/1 | 0.92 | 0.06 | 65,65,65,65 | 0 |
| 57 | MG | 2a | 1753 | 1/1 | 0.92 | 0.18 | 62,62,62,62 | 0 |
| 57 | MG | 1A | 4005 | 1/1 | 0.92 | 0.08 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3310 | 1/1 | 0.92 | 0.12 | 70,70,70,70 | 0 |
| 57 | MG | 1A | 3766 | 1/1 | 0.92 | 0.12 | 37,37,37,37 | 0 |
| 57 | MG | 1a | 1653 | 1/1 | 0.92 | 0.16 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3723 | 1/1 | 0.92 | 0.08 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3048 | 1/1 | 0.92 | 0.13 | 31,31,31,31 | 0 |
| 57 | MG | 2A | 3049 | 1/1 | 0.92 | 0.15 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 4010 | 1/1 | 0.92 | 0.16 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 4012 | 1/1 | 0.92 | 0.13 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3053 | 1/1 | 0.92 | 0.25 | 62,62,62,62 | 0 |
| 57 | MG | 1a | 1657 | 1/1 | 0.92 | 0.08 | 53,53,53,53 | 0 |
| 57 | MG | 2a | 1772 | 1/1 | 0.92 | 0.07 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3740 | 1/1 | 0.92 | 0.15 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3056 | 1/1 | 0.92 | 0.13 | 55,55,55,55 | 0 |
| 57 | MG | 2a | 1779 | 1/1 | 0.92 | 0.17 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3744 | 1/1 | 0.92 | 0.12 | 67,67,67,67 | 0 |
| 57 | MG | 2A | 3328 | 1/1 | 0.92 | 0.16 | 81,81,81,81 | 0 |
| 57 | MG | 1A | 3416 | 1/1 | 0.92 | 0.10 | 64,64,64,64 | 0 |
| 57 | MG | 2A | 3750 | 1/1 | 0.92 | 0.14 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3058 | 1/1 | 0.92 | 0.11 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3417 | 1/1 | 0.92 | 0.10 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3758 | 1/1 | 0.92 | 0.19 | 57,57,57,57 | 0 |
| 57 | MG | 2a | 1790 | 1/1 | 0.92 | 0.17 | 60,60,60,60 | 0 |
| 57 | MG | 2A | 3761 | 1/1 | 0.92 | 0.10 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3418 | 1/1 | 0.92 | 0.11 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3768 | 1/1 | 0.92 | 0.23 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3534 | 1/1 | 0.92 | 0.29 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3803 | 1/1 | 0.92 | 0.07 | 59,59,59,59 | 0 |
| 57 | MG | 2A | 3774 | 1/1 | 0.92 | 0.14 | 59,59,59,59 | 0 |
| 57 | MG | 2a | 1799 | 1/1 | 0.92 | 0.27 | 59,59,59,59 | 0 |
| 57 | MG | 2A | 3341 | 1/1 | 0.92 | 0.11 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3536 | 1/1 | 0.92 | 0.29 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3540 | 1/1 | 0.92 | 0.10 | 29,29,29,29 | 0 |
| 57 | MG | 2A | 3074 | 1/1 | 0.92 | 0.10 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3078 | 1/1 | 0.92 | 0.19 | 55,55,55,55 | 0 |
| 57 | MG | 2a | 1807 | 1/1 | 0.92 | 0.09 | 59,59,59,59 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3354 | 1/1 | 0.92 | 0.08 | 40,40,40,40 | 0 |
| 57 | MG | 1a | 1671 | 1/1 | 0.92 | 0.18 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3811 | 1/1 | 0.92 | 0.08 | 46,46,46,46 | 0 |
| 57 | MG | 1a | 1674 | 1/1 | 0.92 | 0.20 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3421 | 1/1 | 0.92 | 0.20 | 41,41,41,41 | 0 |
| 57 | MG | 2a | 1816 | 1/1 | 0.92 | 0.15 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3548 | 1/1 | 0.92 | 0.09 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3816 | 1/1 | 0.92 | 0.09 | 40,40,40,40 | 0 |
| 57 | MG | 2a | 1820 | 1/1 | 0.92 | 0.13 | 55,55,55,55 | 0 |
| 57 | MG | 1a | 1680 | 1/1 | 0.92 | 0.10 | 60,60,60,60 | 0 |
| 57 | MG | 1a | 1681 | 1/1 | 0.92 | 0.17 | 51,51,51,51 | 0 |
| 57 | MG | 2d | 301 | 1/1 | 0.92 | 0.17 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 4071 | 1/1 | 0.92 | 0.07 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3807 | 1/1 | 0.92 | 0.10 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3814 | 1/1 | 0.92 | 0.16 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3816 | 1/1 | 0.92 | 0.08 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3368 | 1/1 | 0.92 | 0.09 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3817 | 1/1 | 0.92 | 0.43 | 41,41,41,41 | 0 |
| 57 | MG | 2l | 202 | 1/1 | 0.92 | 0.18 | 69,69,69,69 | 0 |
| 57 | MG | 1A | 3827 | 1/1 | 0.92 | 0.19 | 52,52,52,52 | 0 |
| 57 | MG | 2q | 201 | 1/1 | 0.92 | 0.20 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3374 | 1/1 | 0.92 | 0.17 | 58,58,58,58 | 0 |
| 57 | MG | 2r | 101 | 1/1 | 0.92 | 0.08 | 61,61,61,61 | 0 |
| 57 | MG | 2t | 201 | 1/1 | 0.92 | 0.15 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 4076 | 1/1 | 0.92 | 0.14 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 4079 | 1/1 | 0.92 | 0.15 | 56,56,56,56 | 0 |
| 57 | MG | 1a | 1691 | 1/1 | 0.92 | 0.23 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3104 | 1/1 | 0.92 | 0.15 | 47,47,47,47 | 0 |
| 57 | MG | 1a | 1693 | 1/1 | 0.92 | 0.25 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3549 | 1/1 | 0.92 | 0.20 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3281 | 1/1 | 0.92 | 0.13 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3836 | 1/1 | 0.92 | 0.15 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 4085 | 1/1 | 0.92 | 0.26 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3556 | 1/1 | 0.92 | 0.08 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3390 | 1/1 | 0.92 | 0.06 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3395 | 1/1 | 0.92 | 0.15 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3853 | 1/1 | 0.92 | 0.26 | 62,62,62,62 | 0 |
| 57 | MG | 1A | 3838 | 1/1 | 0.92 | 0.10 | 29,29,29,29 | 0 |
| 57 | MG | 27 | 101 | 1/1 | 0.93 | 0.10 | 48,48,48,48 | 0 |
| 57 | MG | 1a | 1687 | 1/1 | 0.93 | 0.28 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3568 | 1/1 | 0.93 | 0.10 | 40,40,40,40 | 0 |
| 57 | MG | 1B | 229 | 1/1 | 0.93 | 0.17 | 56,56,56,56 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 3236 | 1/1 | 0.93 | 0.08 | 32,32,32,32 | 0 |
| 57 | MG | 2A | 3574 | 1/1 | 0.93 | 0.12 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3440 | 1/1 | 0.93 | 0.14 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3063 | 1/1 | 0.93 | 0.21 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3291 | 1/1 | 0.93 | 0.21 | 59,59,59,59 | 0 |
| 57 | MG | 2A | 3582 | 1/1 | 0.93 | 0.07 | 38,38,38,38 | 0 |
| 57 | MG | 1A | 3368 | 1/1 | 0.93 | 0.10 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3083 | 1/1 | 0.93 | 0.19 | 25,25,25,25 | 0 |
| 57 | MG | 2A | 3587 | 1/1 | 0.93 | 0.10 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3068 | 1/1 | 0.93 | 0.29 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3120 | 1/1 | 0.93 | 0.09 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3450 | 1/1 | 0.93 | 0.06 | 28,28,28,28 | 0 |
| 57 | MG | 2A | 3072 | 1/1 | 0.93 | 0.08 | 57,57,57,57 | 0 |
| 57 | MG | 1E | 311 | 1/1 | 0.93 | 0.12 | 45,45,45,45 | 0 |
| 57 | MG | 1E | 312 | 1/1 | 0.93 | 0.04 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3732 | 1/1 | 0.93 | 0.10 | 60,60,60,60 | 0 |
| 57 | MG | 2A | 3607 | 1/1 | 0.93 | 0.07 | 33,33,33,33 | 0 |
| 57 | MG | 1F | 305 | 1/1 | 0.93 | 0.08 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3121 | 1/1 | 0.93 | 0.11 | 37,37,37,37 | 0 |
| 57 | MG | 1a | 1711 | 1/1 | 0.93 | 0.14 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3745 | 1/1 | 0.93 | 0.08 | 19,19,19,19 | 0 |
| 57 | MG | 1A | 3748 | 1/1 | 0.93 | 0.07 | 17,17,17,17 | 0 |
| 57 | MG | 1N | 201 | 1/1 | 0.93 | 0.07 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3622 | 1/1 | 0.93 | 0.16 | 44,44,44,44 | 0 |
| 57 | MG | 1N | 204 | 1/1 | 0.93 | 0.13 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3626 | 1/1 | 0.93 | 0.09 | 52,52,52,52 | 0 |
| 57 | MG | 1a | 1718 | 1/1 | 0.93 | 0.22 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3373 | 1/1 | 0.93 | 0.10 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3638 | 1/1 | 0.93 | 0.08 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3639 | 1/1 | 0.93 | 0.08 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3317 | 1/1 | 0.93 | 0.10 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3759 | 1/1 | 0.93 | 0.09 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3183 | 1/1 | 0.93 | 0.09 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3647 | 1/1 | 0.93 | 0.11 | 40,40,40,40 | 0 |
| 57 | MG | 2A | 3100 | 1/1 | 0.93 | 0.13 | 34,34,34,34 | 0 |
| 57 | MG | 2A | 3652 | 1/1 | 0.93 | 0.15 | 49,49,49,49 | 0 |
| 57 | MG | 1P | 206 | 1/1 | 0.93 | 0.13 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3190 | 1/1 | 0.93 | 0.15 | 25,25,25,25 | 0 |
| 57 | MG | 2A | 3657 | 1/1 | 0.93 | 0.12 | 58,58,58,58 | 0 |
| 57 | MG | 1a | 1729 | 1/1 | 0.93 | 0.10 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3661 | 1/1 | 0.93 | 0.11 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3106 | 1/1 | 0.93 | 0.18 | 48,48,48,48 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 3950 | 1/1 | 0.93 | 0.15 | 42,42,42,42 | 0 |
| 57 | MG | 2A | 3330 | 1/1 | 0.93 | 0.08 | 55,55,55,55 | 0 |
| 57 | MG | 1a | 1732 | 1/1 | 0.93 | 0.12 | 40,40,40,40 | 0 |
| 57 | MG | 1a | 1734 | 1/1 | 0.93 | 0.08 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3769 | 1/1 | 0.93 | 0.08 | 24,24,24,24 | 0 |
| 57 | MG | 2A | 3336 | 1/1 | 0.93 | 0.21 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3337 | 1/1 | 0.93 | 0.10 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3382 | 1/1 | 0.93 | 0.06 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3956 | 1/1 | 0.93 | 0.08 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3084 | 1/1 | 0.93 | 0.09 | 33,33,33,33 | 0 |
| 57 | MG | 2a | 1670 | 1/1 | 0.93 | 0.19 | 58,58,58,58 | 0 |
| 57 | MG | 1a | 1739 | 1/1 | 0.93 | 0.16 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3345 | 1/1 | 0.93 | 0.13 | 39,39,39,39 | 0 |
| 57 | MG | 1U | 203 | 1/1 | 0.93 | 0.19 | 63,63,63,63 | 0 |
| 57 | MG | 2A | 3123 | 1/1 | 0.93 | 0.07 | 74,74,74,74 | 0 |
| 57 | MG | 1U | 206 | 1/1 | 0.93 | 0.15 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3135 | 1/1 | 0.93 | 0.15 | 39,39,39,39 | 0 |
| 57 | MG | 1a | 1752 | 1/1 | 0.93 | 0.10 | 56,56,56,56 | 0 |
| 57 | MG | 1a | 1754 | 1/1 | 0.93 | 0.11 | 38,38,38,38 | 0 |
| 57 | MG | 1a | 1756 | 1/1 | 0.93 | 0.20 | 57,57,57,57 | 0 |
| 57 | MG | 1U | 209 | 1/1 | 0.93 | 0.17 | 38,38,38,38 | 0 |
| 57 | MG | 2a | 1687 | 1/1 | 0.93 | 0.23 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3694 | 1/1 | 0.93 | 0.06 | 41,41,41,41 | 0 |
| 57 | MG | 1U | 210 | 1/1 | 0.93 | 0.08 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3151 | 1/1 | 0.93 | 0.13 | 59,59,59,59 | 0 |
| 57 | MG | 2A | 3703 | 1/1 | 0.93 | 0.10 | 32,32,32,32 | 0 |
| 57 | MG | 1V | 206 | 1/1 | 0.93 | 0.10 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3059 | 1/1 | 0.93 | 0.17 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3961 | 1/1 | 0.93 | 0.07 | 49,49,49,49 | 0 |
| 57 | MG | 2a | 1698 | 1/1 | 0.93 | 0.07 | 74,74,74,74 | 0 |
| 57 | MG | 2a | 1700 | 1/1 | 0.93 | 0.23 | 60,60,60,60 | 0 |
| 57 | MG | 1Z | 301 | 1/1 | 0.93 | 0.12 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3370 | 1/1 | 0.93 | 0.15 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3371 | 1/1 | 0.93 | 0.09 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3163 | 1/1 | 0.93 | 0.12 | 39,39,39,39 | 0 |
| 57 | MG | 2A | 3721 | 1/1 | 0.93 | 0.14 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3796 | 1/1 | 0.93 | 0.15 | 45,45,45,45 | 0 |
| 57 | MG | 1a | 1773 | 1/1 | 0.93 | 0.12 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3800 | 1/1 | 0.93 | 0.08 | 21,21,21,21 | 0 |
| 57 | MG | 1A | 3469 | 1/1 | 0.93 | 0.07 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3377 | 1/1 | 0.93 | 0.14 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3169 | 1/1 | 0.93 | 0.12 | 49,49,49,49 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1a | 1776 | 1/1 | 0.93 | 0.09 | 67,67,67,67 | 0 |
| 57 | MG | 1A | 3308 | 1/1 | 0.93 | 0.25 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3981 | 1/1 | 0.93 | 0.07 | 27,27,27,27 | 0 |
| 57 | MG | 1A | 3312 | 1/1 | 0.93 | 0.07 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3020 | 1/1 | 0.93 | 0.14 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3320 | 1/1 | 0.93 | 0.10 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3401 | 1/1 | 0.93 | 0.08 | 38,38,38,38 | 0 |
| 57 | MG | 15 | 107 | 1/1 | 0.93 | 0.07 | 30,30,30,30 | 0 |
| 57 | MG | 2A | 3391 | 1/1 | 0.93 | 0.13 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3097 | 1/1 | 0.93 | 0.18 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3045 | 1/1 | 0.93 | 0.16 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3759 | 1/1 | 0.93 | 0.08 | 63,63,63,63 | 0 |
| 57 | MG | 1A | 3111 | 1/1 | 0.93 | 0.08 | 30,30,30,30 | 0 |
| 57 | MG | 18 | 102 | 1/1 | 0.93 | 0.16 | 34,34,34,34 | 0 |
| 57 | MG | 2A | 3406 | 1/1 | 0.93 | 0.11 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3769 | 1/1 | 0.93 | 0.10 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3617 | 1/1 | 0.93 | 0.17 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3411 | 1/1 | 0.93 | 0.19 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 4003 | 1/1 | 0.93 | 0.09 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3415 | 1/1 | 0.93 | 0.23 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3618 | 1/1 | 0.93 | 0.14 | 36,36,36,36 | 0 |
| 57 | MG | 2a | 1747 | 1/1 | 0.93 | 0.26 | 63,63,63,63 | 0 |
| 57 | MG | 2A | 3778 | 1/1 | 0.93 | 0.12 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3192 | 1/1 | 0.93 | 0.47 | 65,65,65,65 | 0 |
| 57 | MG | 1A | 3205 | 1/1 | 0.93 | 0.09 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3783 | 1/1 | 0.93 | 0.10 | 77,77,77,77 | 0 |
| 57 | MG | 1A | 3835 | 1/1 | 0.93 | 0.09 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3155 | 1/1 | 0.93 | 0.38 | 36,36,36,36 | 0 |
| 57 | MG | 2A | 3427 | 1/1 | 0.93 | 0.23 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3788 | 1/1 | 0.93 | 0.14 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3503 | 1/1 | 0.93 | 0.20 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3635 | 1/1 | 0.93 | 0.06 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3411 | 1/1 | 0.93 | 0.10 | 27,27,27,27 | 0 |
| 57 | MG | 1A | 3215 | 1/1 | 0.93 | 0.14 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3266 | 1/1 | 0.93 | 0.11 | 45,45,45,45 | 0 |
| 57 | MG | 1j | 201 | 1/1 | 0.93 | 0.30 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 4031 | 1/1 | 0.93 | 0.06 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3644 | 1/1 | 0.93 | 0.10 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3803 | 1/1 | 0.93 | 0.12 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3851 | 1/1 | 0.93 | 0.08 | 22,22,22,22 | 0 |
| 57 | MG | 2A | 3813 | 1/1 | 0.93 | 0.14 | 49,49,49,49 | 0 |
| 57 | MG | 2a | 1780 | 1/1 | 0.93 | 0.19 | 55,55,55,55 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1a | 1632 | 1/1 | 0.93 | 0.13 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 4057 | 1/1 | 0.93 | 0.07 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3508 | 1/1 | 0.93 | 0.13 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 4059 | 1/1 | 0.93 | 0.07 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3825 | 1/1 | 0.93 | 0.08 | 61,61,61,61 | 0 |
| 57 | MG | 1A | 4065 | 1/1 | 0.93 | 0.24 | 54,54,54,54 | 0 |
| 57 | MG | 1w | 105 | 1/1 | 0.93 | 0.07 | 54,54,54,54 | 0 |
| 57 | MG | 1a | 1643 | 1/1 | 0.93 | 0.12 | 38,38,38,38 | 0 |
| 57 | MG | 1A | 3857 | 1/1 | 0.93 | 0.10 | 34,34,34,34 | 0 |
| 57 | MG | 2a | 1793 | 1/1 | 0.93 | 0.11 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 4068 | 1/1 | 0.93 | 0.19 | 61,61,61,61 | 0 |
| 57 | MG | 1A | 3649 | 1/1 | 0.93 | 0.16 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3861 | 1/1 | 0.93 | 0.09 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3652 | 1/1 | 0.93 | 0.10 | 39,39,39,39 | 0 |
| 57 | MG | 2A | 3002 | 1/1 | 0.93 | 0.25 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3866 | 1/1 | 0.93 | 0.14 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3474 | 1/1 | 0.93 | 0.24 | 42,42,42,42 | 0 |
| 57 | MG | 2A | 3475 | 1/1 | 0.93 | 0.30 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3477 | 1/1 | 0.93 | 0.26 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3478 | 1/1 | 0.93 | 0.14 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3510 | 1/1 | 0.93 | 0.28 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3480 | 1/1 | 0.93 | 0.08 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3481 | 1/1 | 0.93 | 0.16 | 61,61,61,61 | 0 |
| 57 | MG | 1A | 3220 | 1/1 | 0.93 | 0.08 | 38,38,38,38 | 0 |
| 57 | MG | 1A | 3157 | 1/1 | 0.93 | 0.09 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3077 | 1/1 | 0.93 | 0.23 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3243 | 1/1 | 0.93 | 0.07 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3244 | 1/1 | 0.93 | 0.08 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3524 | 1/1 | 0.93 | 0.07 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 3525 | 1/1 | 0.93 | 0.22 | 48,48,48,48 | 0 |
| 57 | MG | 2a | 1819 | 1/1 | 0.93 | 0.10 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3249 | 1/1 | 0.93 | 0.07 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3250 | 1/1 | 0.93 | 0.06 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3881 | 1/1 | 0.93 | 0.12 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3502 | 1/1 | 0.93 | 0.13 | 60,60,60,60 | 0 |
| 57 | MG | 2B | 216 | 1/1 | 0.93 | 0.09 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3671 | 1/1 | 0.93 | 0.13 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3673 | 1/1 | 0.93 | 0.06 | 21,21,21,21 | 0 |
| 57 | MG | 2A | 3507 | 1/1 | 0.93 | 0.11 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3230 | 1/1 | 0.93 | 0.14 | 48,48,48,48 | 0 |
| 57 | MG | 2D | 301 | 1/1 | 0.93 | 0.15 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3282 | 1/1 | 0.93 | 0.09 | 49,49,49,49 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2E | 302 | 1/1 | 0.93 | 0.06 | 43,43,43,43 | 0 |
| 57 | MG | 2E | 303 | 1/1 | 0.93 | 0.16 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3283 | 1/1 | 0.93 | 0.19 | 34,34,34,34 | 0 |
| 57 | MG | 2A | 3524 | 1/1 | 0.93 | 0.08 | 52,52,52,52 | 0 |
| 57 | MG | 1a | 1673 | 1/1 | 0.93 | 0.19 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3162 | 1/1 | 0.93 | 0.18 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3292 | 1/1 | 0.93 | 0.17 | 41,41,41,41 | 0 |
| 57 | MG | 1B | 212 | 1/1 | 0.93 | 0.10 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3268 | 1/1 | 0.93 | 0.17 | 76,76,76,76 | 0 |
| 57 | MG | 2w | 103 | 1/1 | 0.93 | 0.06 | 77,77,77,77 | 0 |
| 57 | MG | 1A | 3904 | 1/1 | 0.93 | 0.29 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3297 | 1/1 | 0.93 | 0.16 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3909 | 1/1 | 0.93 | 0.15 | 44,44,44,44 | 0 |
| 57 | MG | 1B | 224 | 1/1 | 0.93 | 0.05 | 36,36,36,36 | 0 |
| 57 | MG | 2A | 3551 | 1/1 | 0.93 | 0.13 | 42,42,42,42 | 0 |
| 57 | MG | 2A | 3553 | 1/1 | 0.93 | 0.14 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3298 | 1/1 | 0.93 | 0.06 | 29,29,29,29 | 0 |
| 57 | MG | 2x | 106 | 1/1 | 0.93 | 0.09 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3437 | 1/1 | 0.93 | 0.16 | 48,48,48,48 | 0 |
| 57 | MG | 20 | 102 | 1/1 | 0.93 | 0.09 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3282 | 1/1 | 0.93 | 0.07 | 50,50,50,50 | 0 |
| 59 | ERY | 2A | 3857 | 51/51 | 0.93 | 0.13 | 26,54,66,69 | 0 |
| 57 | MG | 2A | 3205 | 1/1 | 0.94 | 0.08 | 59,59,59,59 | 0 |
| 57 | MG | 2B | 209 | 1/1 | 0.94 | 0.08 | 65,65,65,65 | 0 |
| 57 | MG | 2B | 210 | 1/1 | 0.94 | 0.19 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3765 | 1/1 | 0.94 | 0.10 | 15,15,15,15 | 0 |
| 57 | MG | 2A | 3473 | 1/1 | 0.94 | 0.06 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3562 | 1/1 | 0.94 | 0.13 | 36,36,36,36 | 0 |
| 57 | MG | 1a | 1785 | 1/1 | 0.94 | 0.14 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3564 | 1/1 | 0.94 | 0.15 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3345 | 1/1 | 0.94 | 0.12 | 61,61,61,61 | 0 |
| 57 | MG | 1A | 3772 | 1/1 | 0.94 | 0.09 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3217 | 1/1 | 0.94 | 0.28 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3346 | 1/1 | 0.94 | 0.05 | 34,34,34,34 | 0 |
| 57 | MG | 2A | 3482 | 1/1 | 0.94 | 0.18 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3570 | 1/1 | 0.94 | 0.09 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 3572 | 1/1 | 0.94 | 0.07 | 48,48,48,48 | 0 |
| 57 | MG | 15 | 108 | 1/1 | 0.94 | 0.08 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3218 | 1/1 | 0.94 | 0.07 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3348 | 1/1 | 0.94 | 0.09 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3492 | 1/1 | 0.94 | 0.06 | 38,38,38,38 | 0 |
| 57 | MG | 2E | 306 | 1/1 | 0.94 | 0.08 | 36,36,36,36 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 3219 | 1/1 | 0.94 | 0.16 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3494 | 1/1 | 0.94 | 0.14 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3156 | 1/1 | 0.94 | 0.07 | 29,29,29,29 | 0 |
| 57 | MG | 1a | 1805 | 1/1 | 0.94 | 0.12 | 65,65,65,65 | 0 |
| 57 | MG | 2A | 3497 | 1/1 | 0.94 | 0.07 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3285 | 1/1 | 0.94 | 0.10 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3447 | 1/1 | 0.94 | 0.11 | 47,47,47,47 | 0 |
| 57 | MG | 1a | 1808 | 1/1 | 0.94 | 0.12 | 61,61,61,61 | 0 |
| 57 | MG | 1A | 3024 | 1/1 | 0.94 | 0.15 | 32,32,32,32 | 0 |
| 57 | MG | 2U | 202 | 1/1 | 0.94 | 0.05 | 41,41,41,41 | 0 |
| 57 | MG | 18 | 107 | 1/1 | 0.94 | 0.14 | 36,36,36,36 | 0 |
| 57 | MG | 1A | 4019 | 1/1 | 0.94 | 0.10 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3510 | 1/1 | 0.94 | 0.16 | 44,44,44,44 | 0 |
| 57 | MG | 1a | 1605 | 1/1 | 0.94 | 0.08 | 63,63,63,63 | 0 |
| 57 | MG | 1A | 4020 | 1/1 | 0.94 | 0.10 | 30,30,30,30 | 0 |
| 57 | MG | 20 | 103 | 1/1 | 0.94 | 0.11 | 65,65,65,65 | 0 |
| 57 | MG | 1a | 1609 | 1/1 | 0.94 | 0.06 | 38,38,38,38 | 0 |
| 57 | MG | 25 | 101 | 1/1 | 0.94 | 0.10 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3521 | 1/1 | 0.94 | 0.06 | 35,35,35,35 | 0 |
| 57 | MG | 2A | 3523 | 1/1 | 0.94 | 0.10 | 39,39,39,39 | 0 |
| 57 | MG | 1a | 1610 | 1/1 | 0.94 | 0.08 | 61,61,61,61 | 0 |
| 57 | MG | 27 | 104 | 1/1 | 0.94 | 0.18 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3526 | 1/1 | 0.94 | 0.12 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3245 | 1/1 | 0.94 | 0.09 | 55,55,55,55 | 0 |
| 57 | MG | 1a | 1611 | 1/1 | 0.94 | 0.11 | 28,28,28,28 | 0 |
| 57 | MG | 1A | 3363 | 1/1 | 0.94 | 0.21 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3248 | 1/1 | 0.94 | 0.09 | 51,51,51,51 | 0 |
| 57 | MG | 1a | 1614 | 1/1 | 0.94 | 0.05 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3815 | 1/1 | 0.94 | 0.08 | 59,59,59,59 | 0 |
| 57 | MG | 2a | 1608 | 1/1 | 0.94 | 0.08 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3537 | 1/1 | 0.94 | 0.12 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 4024 | 1/1 | 0.94 | 0.08 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3364 | 1/1 | 0.94 | 0.10 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3287 | 1/1 | 0.94 | 0.08 | 58,58,58,58 | 0 |
| 57 | MG | 1a | 1624 | 1/1 | 0.94 | 0.21 | 44,44,44,44 | 0 |
| 57 | MG | 2a | 1616 | 1/1 | 0.94 | 0.09 | 67,67,67,67 | 0 |
| 57 | MG | 1a | 1625 | 1/1 | 0.94 | 0.18 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3554 | 1/1 | 0.94 | 0.10 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3259 | 1/1 | 0.94 | 0.08 | 70,70,70,70 | 0 |
| 57 | MG | 2A | 3557 | 1/1 | 0.94 | 0.10 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3559 | 1/1 | 0.94 | 0.07 | 32,32,32,32 | 0 |
| 57 | MG | 1a | 1626 | 1/1 | 0.94 | 0.22 | 59,59,59,59 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1x | 110 | 1/1 | 0.94 | 0.13 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3819 | 1/1 | 0.94 | 0.08 | 18,18,18,18 | 0 |
| 57 | MG | 1A | 4038 | 1/1 | 0.94 | 0.12 | 42,42,42,42 | 0 |
| 57 | MG | 2A | 3265 | 1/1 | 0.94 | 0.08 | 58,58,58,58 | 0 |
| 57 | MG | 2a | 1629 | 1/1 | 0.94 | 0.16 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3570 | 1/1 | 0.94 | 0.07 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3821 | 1/1 | 0.94 | 0.12 | 38,38,38,38 | 0 |
| 57 | MG | 1A | 3826 | 1/1 | 0.94 | 0.18 | 48,48,48,48 | 0 |
| 57 | MG | 1a | 1631 | 1/1 | 0.94 | 0.19 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3271 | 1/1 | 0.94 | 0.10 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3272 | 1/1 | 0.94 | 0.10 | 36,36,36,36 | 0 |
| 57 | MG | 2A | 3580 | 1/1 | 0.94 | 0.14 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3009 | 1/1 | 0.94 | 0.13 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3592 | 1/1 | 0.94 | 0.13 | 34,34,34,34 | 0 |
| 57 | MG | 2A | 3016 | 1/1 | 0.94 | 0.22 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3288 | 1/1 | 0.94 | 0.08 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3832 | 1/1 | 0.94 | 0.06 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3280 | 1/1 | 0.94 | 0.13 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 4060 | 1/1 | 0.94 | 0.10 | 31,31,31,31 | 0 |
| 57 | MG | 2A | 3027 | 1/1 | 0.94 | 0.22 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3112 | 1/1 | 0.94 | 0.15 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 4066 | 1/1 | 0.94 | 0.14 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3296 | 1/1 | 0.94 | 0.23 | 41,41,41,41 | 0 |
| 57 | MG | 2a | 1649 | 1/1 | 0.94 | 0.11 | 70,70,70,70 | 0 |
| 57 | MG | 1A | 3025 | 1/1 | 0.94 | 0.13 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3604 | 1/1 | 0.94 | 0.11 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3037 | 1/1 | 0.94 | 0.09 | 33,33,33,33 | 0 |
| 57 | MG | 2a | 1653 | 1/1 | 0.94 | 0.12 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3611 | 1/1 | 0.94 | 0.18 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3231 | 1/1 | 0.94 | 0.06 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3612 | 1/1 | 0.94 | 0.10 | 21,21,21,21 | 0 |
| 57 | MG | 1A | 3613 | 1/1 | 0.94 | 0.12 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3468 | 1/1 | 0.94 | 0.19 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3046 | 1/1 | 0.94 | 0.10 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3377 | 1/1 | 0.94 | 0.27 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 4078 | 1/1 | 0.94 | 0.15 | 45,45,45,45 | 0 |
| 57 | MG | 2a | 1664 | 1/1 | 0.94 | 0.18 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3628 | 1/1 | 0.94 | 0.11 | 41,41,41,41 | 0 |
| 57 | MG | 2a | 1667 | 1/1 | 0.94 | 0.30 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3631 | 1/1 | 0.94 | 0.08 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3623 | 1/1 | 0.94 | 0.17 | 16,16,16,16 | 0 |
| 57 | MG | 1A | 3048 | 1/1 | 0.94 | 0.07 | 22,22,22,22 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 3627 | 1/1 | 0.94 | 0.12 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3859 | 1/1 | 0.94 | 0.13 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3640 | 1/1 | 0.94 | 0.08 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 4083 | 1/1 | 0.94 | 0.07 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3308 | 1/1 | 0.94 | 0.08 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 4084 | 1/1 | 0.94 | 0.09 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3234 | 1/1 | 0.94 | 0.19 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3311 | 1/1 | 0.94 | 0.10 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3313 | 1/1 | 0.94 | 0.11 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3472 | 1/1 | 0.94 | 0.12 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3473 | 1/1 | 0.94 | 0.18 | 61,61,61,61 | 0 |
| 57 | MG | 2a | 1689 | 1/1 | 0.94 | 0.19 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3061 | 1/1 | 0.94 | 0.17 | 48,48,48,48 | 0 |
| 57 | MG | 1B | 205 | 1/1 | 0.94 | 0.21 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3659 | 1/1 | 0.94 | 0.09 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3636 | 1/1 | 0.94 | 0.08 | 17,17,17,17 | 0 |
| 57 | MG | 1A | 3474 | 1/1 | 0.94 | 0.26 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3061 | 1/1 | 0.94 | 0.12 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3062 | 1/1 | 0.94 | 0.19 | 38,38,38,38 | 0 |
| 57 | MG | 1a | 1679 | 1/1 | 0.94 | 0.07 | 71,71,71,71 | 0 |
| 57 | MG | 1A | 3478 | 1/1 | 0.94 | 0.13 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3387 | 1/1 | 0.94 | 0.16 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3181 | 1/1 | 0.94 | 0.09 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3077 | 1/1 | 0.94 | 0.13 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3331 | 1/1 | 0.94 | 0.07 | 64,64,64,64 | 0 |
| 57 | MG | 1B | 215 | 1/1 | 0.94 | 0.20 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3880 | 1/1 | 0.94 | 0.09 | 25,25,25,25 | 0 |
| 57 | MG | 1B | 217 | 1/1 | 0.94 | 0.07 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3085 | 1/1 | 0.94 | 0.09 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3122 | 1/1 | 0.94 | 0.07 | 31,31,31,31 | 0 |
| 57 | MG | 2A | 3338 | 1/1 | 0.94 | 0.06 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3247 | 1/1 | 0.94 | 0.06 | 51,51,51,51 | 0 |
| 57 | MG | 2a | 1716 | 1/1 | 0.94 | 0.14 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3392 | 1/1 | 0.94 | 0.29 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3659 | 1/1 | 0.94 | 0.08 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3342 | 1/1 | 0.94 | 0.18 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3497 | 1/1 | 0.94 | 0.13 | 59,59,59,59 | 0 |
| 57 | MG | 1A | 3891 | 1/1 | 0.94 | 0.16 | 33,33,33,33 | 0 |
| 57 | MG | 2a | 1724 | 1/1 | 0.94 | 0.13 | 55,55,55,55 | 0 |
| 57 | MG | 1a | 1696 | 1/1 | 0.94 | 0.10 | 56,56,56,56 | 0 |
| 57 | MG | 1a | 1697 | 1/1 | 0.94 | 0.12 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3352 | 1/1 | 0.94 | 0.06 | 66,66,66,66 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2a | 1729 | 1/1 | 0.94 | 0.10 | 52,52,52,52 | 0 |
| 57 | MG | 2a | 1730 | 1/1 | 0.94 | 0.22 | 59,59,59,59 | 0 |
| 57 | MG | 1A | 3395 | 1/1 | 0.94 | 0.08 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3704 | 1/1 | 0.94 | 0.10 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3896 | 1/1 | 0.94 | 0.13 | 45,45,45,45 | 0 |
| 57 | MG | 1a | 1701 | 1/1 | 0.94 | 0.15 | 61,61,61,61 | 0 |
| 57 | MG | 1a | 1702 | 1/1 | 0.94 | 0.15 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3712 | 1/1 | 0.94 | 0.07 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3102 | 1/1 | 0.94 | 0.07 | 56,56,56,56 | 0 |
| 57 | MG | 2a | 1740 | 1/1 | 0.94 | 0.19 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3715 | 1/1 | 0.94 | 0.06 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3663 | 1/1 | 0.94 | 0.11 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3188 | 1/1 | 0.94 | 0.27 | 40,40,40,40 | 0 |
| 57 | MG | 2A | 3362 | 1/1 | 0.94 | 0.16 | 58,58,58,58 | 0 |
| 57 | MG | 1D | 301 | 1/1 | 0.94 | 0.16 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 3063 | 1/1 | 0.94 | 0.18 | 33,33,33,33 | 0 |
| 57 | MG | 1E | 303 | 1/1 | 0.94 | 0.17 | 32,32,32,32 | 0 |
| 57 | MG | 1E | 308 | 1/1 | 0.94 | 0.08 | 21,21,21,21 | 0 |
| 57 | MG | 1A | 3127 | 1/1 | 0.94 | 0.39 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3315 | 1/1 | 0.94 | 0.15 | 33,33,33,33 | 0 |
| 57 | MG | 2A | 3732 | 1/1 | 0.94 | 0.08 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3114 | 1/1 | 0.94 | 0.09 | 56,56,56,56 | 0 |
| 57 | MG | 2a | 1757 | 1/1 | 0.94 | 0.10 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3675 | 1/1 | 0.94 | 0.07 | 33,33,33,33 | 0 |
| 57 | MG | 2A | 3736 | 1/1 | 0.94 | 0.12 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3251 | 1/1 | 0.94 | 0.09 | 26,26,26,26 | 0 |
| 57 | MG | 2a | 1762 | 1/1 | 0.94 | 0.22 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3091 | 1/1 | 0.94 | 0.10 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3095 | 1/1 | 0.94 | 0.18 | 41,41,41,41 | 0 |
| 57 | MG | 1G | 202 | 1/1 | 0.94 | 0.11 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3124 | 1/1 | 0.94 | 0.10 | 52,52,52,52 | 0 |
| 57 | MG | 2a | 1768 | 1/1 | 0.94 | 0.10 | 73,73,73,73 | 0 |
| 57 | MG | 1a | 1722 | 1/1 | 0.94 | 0.10 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3701 | 1/1 | 0.94 | 0.09 | 25,25,25,25 | 0 |
| 57 | MG | 2A | 3380 | 1/1 | 0.94 | 0.05 | 46,46,46,46 | 0 |
| 57 | MG | 2a | 1774 | 1/1 | 0.94 | 0.07 | 63,63,63,63 | 0 |
| 57 | MG | 2A | 3136 | 1/1 | 0.94 | 0.06 | 33,33,33,33 | 0 |
| 57 | MG | 1A | 3131 | 1/1 | 0.94 | 0.18 | 39,39,39,39 | 0 |
| 57 | MG | 2a | 1778 | 1/1 | 0.94 | 0.11 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3141 | 1/1 | 0.94 | 0.14 | 18,18,18,18 | 0 |
| 57 | MG | 2A | 3385 | 1/1 | 0.94 | 0.05 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3412 | 1/1 | 0.94 | 0.09 | 30,30,30,30 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1a | 1731 | 1/1 | 0.94 | 0.11 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3258 | 1/1 | 0.94 | 0.08 | 61,61,61,61 | 0 |
| 57 | MG | 1A | 3008 | 1/1 | 0.94 | 0.06 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 3716 | 1/1 | 0.94 | 0.08 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3773 | 1/1 | 0.94 | 0.07 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3051 | 1/1 | 0.94 | 0.23 | 38,38,38,38 | 0 |
| 57 | MG | 1A | 3022 | 1/1 | 0.94 | 0.10 | 42,42,42,42 | 0 |
| 57 | MG | 2A | 3398 | 1/1 | 0.94 | 0.23 | 53,53,53,53 | 0 |
| 57 | MG | 1R | 202 | 1/1 | 0.94 | 0.10 | 27,27,27,27 | 0 |
| 57 | MG | 2A | 3400 | 1/1 | 0.94 | 0.14 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3781 | 1/1 | 0.94 | 0.07 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3401 | 1/1 | 0.94 | 0.28 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3402 | 1/1 | 0.94 | 0.23 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3208 | 1/1 | 0.94 | 0.09 | 40,40,40,40 | 0 |
| 57 | MG | 1R | 205 | 1/1 | 0.94 | 0.13 | 29,29,29,29 | 0 |
| 57 | MG | 1a | 1745 | 1/1 | 0.94 | 0.15 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3419 | 1/1 | 0.94 | 0.08 | 48,48,48,48 | 0 |
| 57 | MG | 2a | 1802 | 1/1 | 0.94 | 0.10 | 39,39,39,39 | 0 |
| 57 | MG | 2A | 3413 | 1/1 | 0.94 | 0.29 | 54,54,54,54 | 0 |
| 57 | MG | 1a | 1748 | 1/1 | 0.94 | 0.10 | 53,53,53,53 | 0 |
| 57 | MG | 1a | 1751 | 1/1 | 0.94 | 0.14 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3416 | 1/1 | 0.94 | 0.28 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3797 | 1/1 | 0.94 | 0.16 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3420 | 1/1 | 0.94 | 0.15 | 28,28,28,28 | 0 |
| 57 | MG | 2a | 1810 | 1/1 | 0.94 | 0.21 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3418 | 1/1 | 0.94 | 0.07 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3726 | 1/1 | 0.94 | 0.10 | 22,22,22,22 | 0 |
| 57 | MG | 1a | 1755 | 1/1 | 0.94 | 0.11 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3337 | 1/1 | 0.94 | 0.17 | 42,42,42,42 | 0 |
| 57 | MG | 2A | 3175 | 1/1 | 0.94 | 0.19 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3808 | 1/1 | 0.94 | 0.06 | 29,29,29,29 | 0 |
| 57 | MG | 2A | 3425 | 1/1 | 0.94 | 0.12 | 31,31,31,31 | 0 |
| 57 | MG | 2A | 3176 | 1/1 | 0.94 | 0.13 | 42,42,42,42 | 0 |
| 57 | MG | 1U | 204 | 1/1 | 0.94 | 0.12 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3210 | 1/1 | 0.94 | 0.11 | 49,49,49,49 | 0 |
| 57 | MG | 1a | 1759 | 1/1 | 0.94 | 0.08 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3823 | 1/1 | 0.94 | 0.09 | 52,52,52,52 | 0 |
| 57 | MG | 1a | 1760 | 1/1 | 0.94 | 0.17 | 50,50,50,50 | 0 |
| 57 | MG | 2f | 202 | 1/1 | 0.94 | 0.09 | 63,63,63,63 | 0 |
| 57 | MG | 1A | 3212 | 1/1 | 0.94 | 0.12 | 55,55,55,55 | 0 |
| 57 | MG | 1a | 1764 | 1/1 | 0.94 | 0.07 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3437 | 1/1 | 0.94 | 0.09 | 45,45,45,45 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3832 | 1/1 | 0.94 | 0.14 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3743 | 1/1 | 0.94 | 0.07 | 60,60,60,60 | 0 |
| 57 | MG | 2A | 3439 | 1/1 | 0.94 | 0.06 | 50,50,50,50 | 0 |
| 57 | MG | 1a | 1766 | 1/1 | 0.94 | 0.08 | 65,65,65,65 | 0 |
| 57 | MG | 1V | 202 | 1/1 | 0.94 | 0.30 | 49,49,49,49 | 0 |
| 57 | MG | 2q | 203 | 1/1 | 0.94 | 0.15 | 60,60,60,60 | 0 |
| 57 | MG | 2A | 3443 | 1/1 | 0.94 | 0.20 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3545 | 1/1 | 0.94 | 0.23 | 48,48,48,48 | 0 |
| 57 | MG | 1V | 207 | 1/1 | 0.94 | 0.07 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3843 | 1/1 | 0.94 | 0.16 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3844 | 1/1 | 0.94 | 0.17 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3190 | 1/1 | 0.94 | 0.18 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3453 | 1/1 | 0.94 | 0.14 | 68,68,68,68 | 0 |
| 57 | MG | 2A | 3849 | 1/1 | 0.94 | 0.09 | 53,53,53,53 | 0 |
| 57 | MG | 1W | 203 | 1/1 | 0.94 | 0.15 | 42,42,42,42 | 0 |
| 57 | MG | 1W | 207 | 1/1 | 0.94 | 0.10 | 38,38,38,38 | 0 |
| 57 | MG | 1A | 3104 | 1/1 | 0.94 | 0.12 | 33,33,33,33 | 0 |
| 57 | MG | 1A | 3425 | 1/1 | 0.94 | 0.09 | 41,41,41,41 | 0 |
| 57 | MG | 2B | 201 | 1/1 | 0.94 | 0.08 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3753 | 1/1 | 0.94 | 0.10 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3461 | 1/1 | 0.94 | 0.08 | 54,54,54,54 | 0 |
| 57 | MG | 2z | 101 | 1/1 | 0.94 | 0.07 | 50,50,50,50 | 0 |
| 57 | MG | 10 | 103 | 1/1 | 0.94 | 0.11 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3217 | 1/1 | 0.94 | 0.13 | 36,36,36,36 | 0 |
| 57 | MG | 2A | 3202 | 1/1 | 0.94 | 0.06 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3430 | 1/1 | 0.94 | 0.16 | 36,36,36,36 | 0 |
| 57 | MG | 1F | 304 | 1/1 | 0.95 | 0.09 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3908 | 1/1 | 0.95 | 0.09 | 23,23,23,23 | 0 |
| 57 | MG | 1A | 3375 | 1/1 | 0.95 | 0.13 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3839 | 1/1 | 0.95 | 0.10 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3544 | 1/1 | 0.95 | 0.14 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3167 | 1/1 | 0.95 | 0.10 | 48,48,48,48 | 0 |
| 57 | MG | 1G | 201 | 1/1 | 0.95 | 0.09 | 36,36,36,36 | 0 |
| 57 | MG | 1A | 3912 | 1/1 | 0.95 | 0.07 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3711 | 1/1 | 0.95 | 0.09 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3450 | 1/1 | 0.95 | 0.20 | 39,39,39,39 | 0 |
| 57 | MG | 2A | 3451 | 1/1 | 0.95 | 0.11 | 38,38,38,38 | 0 |
| 57 | MG | 1I | 201 | 1/1 | 0.95 | 0.07 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3376 | 1/1 | 0.95 | 0.17 | 38,38,38,38 | 0 |
| 57 | MG | 1N | 203 | 1/1 | 0.95 | 0.09 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3715 | 1/1 | 0.95 | 0.08 | 50,50,50,50 | 0 |
| 57 | MG | 1a | 1743 | 1/1 | 0.95 | 0.09 | 46,46,46,46 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3855 | 1/1 | 0.95 | 0.15 | 63,63,63,63 | 0 |
| 57 | MG | 2A | 3856 | 1/1 | 0.95 | 0.07 | 52,52,52,52 | 0 |
| 57 | MG | 1a | 1744 | 1/1 | 0.95 | 0.10 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3919 | 1/1 | 0.95 | 0.08 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3179 | 1/1 | 0.95 | 0.15 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3462 | 1/1 | 0.95 | 0.15 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3317 | 1/1 | 0.95 | 0.14 | 47,47,47,47 | 0 |
| 57 | MG | 1P | 204 | 1/1 | 0.95 | 0.19 | 35,35,35,35 | 0 |
| 57 | MG | 2A | 3467 | 1/1 | 0.95 | 0.10 | 40,40,40,40 | 0 |
| 57 | MG | 1a | 1750 | 1/1 | 0.95 | 0.12 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3925 | 1/1 | 0.95 | 0.08 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3318 | 1/1 | 0.95 | 0.12 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3471 | 1/1 | 0.95 | 0.15 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3551 | 1/1 | 0.95 | 0.12 | 23,23,23,23 | 0 |
| 57 | MG | 1Q | 206 | 1/1 | 0.95 | 0.05 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3552 | 1/1 | 0.95 | 0.08 | 42,42,42,42 | 0 |
| 57 | MG | 1R | 201 | 1/1 | 0.95 | 0.18 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3476 | 1/1 | 0.95 | 0.18 | 28,28,28,28 | 0 |
| 57 | MG | 1A | 3049 | 1/1 | 0.95 | 0.07 | 34,34,34,34 | 0 |
| 57 | MG | 2A | 3191 | 1/1 | 0.95 | 0.12 | 69,69,69,69 | 0 |
| 57 | MG | 1A | 3724 | 1/1 | 0.95 | 0.12 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3268 | 1/1 | 0.95 | 0.07 | 51,51,51,51 | 0 |
| 57 | MG | 1a | 1762 | 1/1 | 0.95 | 0.09 | 49,49,49,49 | 0 |
| 57 | MG | 2D | 304 | 1/1 | 0.95 | 0.21 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3196 | 1/1 | 0.95 | 0.06 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3452 | 1/1 | 0.95 | 0.06 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3321 | 1/1 | 0.95 | 0.17 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3386 | 1/1 | 0.95 | 0.16 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3946 | 1/1 | 0.95 | 0.10 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3738 | 1/1 | 0.95 | 0.14 | 51,51,51,51 | 0 |
| 57 | MG | 2E | 308 | 1/1 | 0.95 | 0.11 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3948 | 1/1 | 0.95 | 0.06 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 3233 | 1/1 | 0.95 | 0.27 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3567 | 1/1 | 0.95 | 0.06 | 38,38,38,38 | 0 |
| 57 | MG | 2F | 303 | 1/1 | 0.95 | 0.12 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3569 | 1/1 | 0.95 | 0.17 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3211 | 1/1 | 0.95 | 0.16 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3953 | 1/1 | 0.95 | 0.10 | 53,53,53,53 | 0 |
| 57 | MG | 2P | 201 | 1/1 | 0.95 | 0.05 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3213 | 1/1 | 0.95 | 0.12 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3090 | 1/1 | 0.95 | 0.07 | 38,38,38,38 | 0 |
| 57 | MG | 2R | 203 | 1/1 | 0.95 | 0.25 | 50,50,50,50 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3501 | 1/1 | 0.95 | 0.14 | 45,45,45,45 | 0 |
| 57 | MG | 2T | 203 | 1/1 | 0.95 | 0.05 | 38,38,38,38 | 0 |
| 57 | MG | 1W | 202 | 1/1 | 0.95 | 0.10 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3503 | 1/1 | 0.95 | 0.09 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3460 | 1/1 | 0.95 | 0.12 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3752 | 1/1 | 0.95 | 0.05 | 26,26,26,26 | 0 |
| 57 | MG | 1X | 101 | 1/1 | 0.95 | 0.07 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3958 | 1/1 | 0.95 | 0.09 | 53,53,53,53 | 0 |
| 57 | MG | 1X | 107 | 1/1 | 0.95 | 0.08 | 48,48,48,48 | 0 |
| 57 | MG | 2I | 3501 | 1/1 | 0.95 | 0.48 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3573 | 1/1 | 0.95 | 0.09 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3517 | 1/1 | 0.95 | 0.09 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3518 | 1/1 | 0.95 | 0.13 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3519 | 1/1 | 0.95 | 0.07 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3235 | 1/1 | 0.95 | 0.13 | 39,39,39,39 | 0 |
| 57 | MG | 2A | 3228 | 1/1 | 0.95 | 0.08 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3522 | 1/1 | 0.95 | 0.10 | 32,32,32,32 | 0 |
| 57 | MG | 28 | 102 | 1/1 | 0.95 | 0.25 | 56,56,56,56 | 0 |
| 57 | MG | 2a | 1601 | 1/1 | 0.95 | 0.14 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3761 | 1/1 | 0.95 | 0.11 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3576 | 1/1 | 0.95 | 0.10 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3966 | 1/1 | 0.95 | 0.11 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3763 | 1/1 | 0.95 | 0.13 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3326 | 1/1 | 0.95 | 0.07 | 43,43,43,43 | 0 |
| 57 | MG | 10 | 109 | 1/1 | 0.95 | 0.13 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3972 | 1/1 | 0.95 | 0.15 | 37,37,37,37 | 0 |
| 57 | MG | 1I | 101 | 1/1 | 0.95 | 0.15 | 38,38,38,38 | 0 |
| 57 | MG | 2a | 1611 | 1/1 | 0.95 | 0.14 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3976 | 1/1 | 0.95 | 0.16 | 61,61,61,61 | 0 |
| 57 | MG | 13 | 101 | 1/1 | 0.95 | 0.17 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3539 | 1/1 | 0.95 | 0.08 | 37,37,37,37 | 0 |
| 57 | MG | 13 | 103 | 1/1 | 0.95 | 0.07 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3542 | 1/1 | 0.95 | 0.13 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3979 | 1/1 | 0.95 | 0.07 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3580 | 1/1 | 0.95 | 0.23 | 35,35,35,35 | 0 |
| 57 | MG | 2A | 3549 | 1/1 | 0.95 | 0.08 | 40,40,40,40 | 0 |
| 57 | MG | 15 | 104 | 1/1 | 0.95 | 0.27 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3983 | 1/1 | 0.95 | 0.09 | 42,42,42,42 | 0 |
| 57 | MG | 1a | 1810 | 1/1 | 0.95 | 0.13 | 59,59,59,59 | 0 |
| 57 | MG | 1a | 1811 | 1/1 | 0.95 | 0.06 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3466 | 1/1 | 0.95 | 0.11 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3558 | 1/1 | 0.95 | 0.13 | 48,48,48,48 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1d | 301 | 1/1 | 0.95 | 0.12 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3560 | 1/1 | 0.95 | 0.17 | 59,59,59,59 | 0 |
| 57 | MG | 1e | 201 | 1/1 | 0.95 | 0.27 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3037 | 1/1 | 0.95 | 0.06 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3393 | 1/1 | 0.95 | 0.13 | 62,62,62,62 | 0 |
| 57 | MG | 2A | 3254 | 1/1 | 0.95 | 0.09 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3990 | 1/1 | 0.95 | 0.05 | 32,32,32,32 | 0 |
| 57 | MG | 1m | 3001 | 1/1 | 0.95 | 0.05 | 46,46,46,46 | 0 |
| 57 | MG | 16 | 102 | 1/1 | 0.95 | 0.14 | 62,62,62,62 | 0 |
| 57 | MG | 2A | 3573 | 1/1 | 0.95 | 0.07 | 28,28,28,28 | 0 |
| 57 | MG | 1n | 103 | 1/1 | 0.95 | 0.19 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3773 | 1/1 | 0.95 | 0.06 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3394 | 1/1 | 0.95 | 0.27 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3577 | 1/1 | 0.95 | 0.06 | 28,28,28,28 | 0 |
| 57 | MG | 1A | 3997 | 1/1 | 0.95 | 0.06 | 36,36,36,36 | 0 |
| 57 | MG | 2A | 3579 | 1/1 | 0.95 | 0.10 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3776 | 1/1 | 0.95 | 0.05 | 21,21,21,21 | 0 |
| 57 | MG | 1A | 4000 | 1/1 | 0.95 | 0.14 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3779 | 1/1 | 0.95 | 0.05 | 57,57,57,57 | 0 |
| 57 | MG | 1a | 1601 | 1/1 | 0.95 | 0.08 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3781 | 1/1 | 0.95 | 0.09 | 22,22,22,22 | 0 |
| 57 | MG | 2A | 3588 | 1/1 | 0.95 | 0.13 | 35,35,35,35 | 0 |
| 57 | MG | 2A | 3591 | 1/1 | 0.95 | 0.10 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3592 | 1/1 | 0.95 | 0.06 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3593 | 1/1 | 0.95 | 0.07 | 53,53,53,53 | 0 |
| 57 | MG | 1a | 1603 | 1/1 | 0.95 | 0.09 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3596 | 1/1 | 0.95 | 0.08 | 24,24,24,24 | 0 |
| 57 | MG | 1x | 104 | 1/1 | 0.95 | 0.10 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3789 | 1/1 | 0.95 | 0.06 | 24,24,24,24 | 0 |
| 57 | MG | 1A | 3093 | 1/1 | 0.95 | 0.06 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3795 | 1/1 | 0.95 | 0.12 | 49,49,49,49 | 0 |
| 57 | MG | 1x | 111 | 1/1 | 0.95 | 0.26 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3603 | 1/1 | 0.95 | 0.08 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3284 | 1/1 | 0.95 | 0.06 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3799 | 1/1 | 0.95 | 0.10 | 19,19,19,19 | 0 |
| 57 | MG | 2A | 3608 | 1/1 | 0.95 | 0.09 | 24,24,24,24 | 0 |
| 57 | MG | 2a | 1668 | 1/1 | 0.95 | 0.10 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3281 | 1/1 | 0.95 | 0.12 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3060 | 1/1 | 0.95 | 0.13 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3245 | 1/1 | 0.95 | 0.04 | 47,47,47,47 | 0 |
| 57 | MG | 1a | 1615 | 1/1 | 0.95 | 0.10 | 36,36,36,36 | 0 |
| 57 | MG | 2A | 3613 | 1/1 | 0.95 | 0.14 | 50,50,50,50 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3286 | 1/1 | 0.95 | 0.09 | 64,64,64,64 | 0 |
| 57 | MG | 2A | 3615 | 1/1 | 0.95 | 0.09 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3616 | 1/1 | 0.95 | 0.16 | 63,63,63,63 | 0 |
| 57 | MG | 2A | 3006 | 1/1 | 0.95 | 0.11 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3591 | 1/1 | 0.95 | 0.16 | 42,42,42,42 | 0 |
| 57 | MG | 2a | 1681 | 1/1 | 0.95 | 0.17 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3010 | 1/1 | 0.95 | 0.15 | 48,48,48,48 | 0 |
| 57 | MG | 2a | 1683 | 1/1 | 0.95 | 0.18 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3011 | 1/1 | 0.95 | 0.12 | 40,40,40,40 | 0 |
| 57 | MG | 2a | 1685 | 1/1 | 0.95 | 0.16 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3625 | 1/1 | 0.95 | 0.20 | 42,42,42,42 | 0 |
| 57 | MG | 2A | 3012 | 1/1 | 0.95 | 0.04 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3013 | 1/1 | 0.95 | 0.17 | 32,32,32,32 | 0 |
| 57 | MG | 2A | 3630 | 1/1 | 0.95 | 0.17 | 54,54,54,54 | 0 |
| 57 | MG | 1a | 1620 | 1/1 | 0.95 | 0.08 | 53,53,53,53 | 0 |
| 57 | MG | 2a | 1692 | 1/1 | 0.95 | 0.20 | 61,61,61,61 | 0 |
| 57 | MG | 2A | 3633 | 1/1 | 0.95 | 0.15 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3808 | 1/1 | 0.95 | 0.63 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 4022 | 1/1 | 0.95 | 0.12 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3335 | 1/1 | 0.95 | 0.41 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3597 | 1/1 | 0.95 | 0.06 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3025 | 1/1 | 0.95 | 0.15 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3203 | 1/1 | 0.95 | 0.17 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3158 | 1/1 | 0.95 | 0.11 | 25,25,25,25 | 0 |
| 57 | MG | 2A | 3645 | 1/1 | 0.95 | 0.09 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3291 | 1/1 | 0.95 | 0.20 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 4034 | 1/1 | 0.95 | 0.07 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 4036 | 1/1 | 0.95 | 0.12 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3651 | 1/1 | 0.95 | 0.10 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3033 | 1/1 | 0.95 | 0.06 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3653 | 1/1 | 0.95 | 0.10 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3005 | 1/1 | 0.95 | 0.06 | 44,44,44,44 | 0 |
| 57 | MG | 2a | 1712 | 1/1 | 0.95 | 0.07 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3036 | 1/1 | 0.95 | 0.07 | 50,50,50,50 | 0 |
| 57 | MG | 2a | 1714 | 1/1 | 0.95 | 0.12 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3075 | 1/1 | 0.95 | 0.06 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3312 | 1/1 | 0.95 | 0.16 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 4040 | 1/1 | 0.95 | 0.09 | 32,32,32,32 | 0 |
| 57 | MG | 2a | 1718 | 1/1 | 0.95 | 0.13 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3040 | 1/1 | 0.95 | 0.20 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3484 | 1/1 | 0.95 | 0.17 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 4045 | 1/1 | 0.95 | 0.07 | 51,51,51,51 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2a | 1722 | 1/1 | 0.95 | 0.09 | 72,72,72,72 | 0 |
| 57 | MG | 2A | 3669 | 1/1 | 0.95 | 0.07 | 62,62,62,62 | 0 |
| 57 | MG | 1A | 4051 | 1/1 | 0.95 | 0.06 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3485 | 1/1 | 0.95 | 0.13 | 49,49,49,49 | 0 |
| 57 | MG | 1a | 1642 | 1/1 | 0.95 | 0.10 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3488 | 1/1 | 0.95 | 0.13 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3824 | 1/1 | 0.95 | 0.09 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3342 | 1/1 | 0.95 | 0.14 | 41,41,41,41 | 0 |
| 57 | MG | 2a | 1731 | 1/1 | 0.95 | 0.14 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3491 | 1/1 | 0.95 | 0.09 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3679 | 1/1 | 0.95 | 0.14 | 63,63,63,63 | 0 |
| 57 | MG | 1A | 3413 | 1/1 | 0.95 | 0.07 | 41,41,41,41 | 0 |
| 57 | MG | 2a | 1735 | 1/1 | 0.95 | 0.28 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3098 | 1/1 | 0.95 | 0.10 | 35,35,35,35 | 0 |
| 57 | MG | 2A | 3055 | 1/1 | 0.95 | 0.13 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3495 | 1/1 | 0.95 | 0.06 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3332 | 1/1 | 0.95 | 0.14 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3167 | 1/1 | 0.95 | 0.10 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3632 | 1/1 | 0.95 | 0.09 | 29,29,29,29 | 0 |
| 57 | MG | 2A | 3687 | 1/1 | 0.95 | 0.09 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3213 | 1/1 | 0.95 | 0.19 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3052 | 1/1 | 0.95 | 0.12 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3504 | 1/1 | 0.95 | 0.10 | 41,41,41,41 | 0 |
| 57 | MG | 2a | 1746 | 1/1 | 0.95 | 0.23 | 62,62,62,62 | 0 |
| 57 | MG | 1a | 1661 | 1/1 | 0.95 | 0.14 | 64,64,64,64 | 0 |
| 57 | MG | 1a | 1664 | 1/1 | 0.95 | 0.16 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3178 | 1/1 | 0.95 | 0.15 | 32,32,32,32 | 0 |
| 57 | MG | 2a | 1751 | 1/1 | 0.95 | 0.13 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 4077 | 1/1 | 0.95 | 0.06 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3069 | 1/1 | 0.95 | 0.09 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3701 | 1/1 | 0.95 | 0.08 | 47,47,47,47 | 0 |
| 57 | MG | 2a | 1755 | 1/1 | 0.95 | 0.15 | 76,76,76,76 | 0 |
| 57 | MG | 1A | 3079 | 1/1 | 0.95 | 0.08 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3358 | 1/1 | 0.95 | 0.20 | 40,40,40,40 | 0 |
| 57 | MG | 1a | 1669 | 1/1 | 0.95 | 0.08 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3135 | 1/1 | 0.95 | 0.12 | 30,30,30,30 | 0 |
| 57 | MG | 2A | 3075 | 1/1 | 0.95 | 0.06 | 40,40,40,40 | 0 |
| 57 | MG | 2A | 3076 | 1/1 | 0.95 | 0.11 | 33,33,33,33 | 0 |
| 57 | MG | 1A | 3852 | 1/1 | 0.95 | 0.12 | 25,25,25,25 | 0 |
| 57 | MG | 1A | 3139 | 1/1 | 0.95 | 0.07 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3855 | 1/1 | 0.95 | 0.13 | 36,36,36,36 | 0 |
| 57 | MG | 2A | 3082 | 1/1 | 0.95 | 0.11 | 55,55,55,55 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3719 | 1/1 | 0.95 | 0.09 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3648 | 1/1 | 0.95 | 0.08 | 64,64,64,64 | 0 |
| 57 | MG | 2A | 3359 | 1/1 | 0.95 | 0.07 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3185 | 1/1 | 0.95 | 0.08 | 38,38,38,38 | 0 |
| 57 | MG | 1A | 3516 | 1/1 | 0.95 | 0.21 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3518 | 1/1 | 0.95 | 0.07 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3863 | 1/1 | 0.95 | 0.18 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3728 | 1/1 | 0.95 | 0.12 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3028 | 1/1 | 0.95 | 0.22 | 69,69,69,69 | 0 |
| 57 | MG | 1A | 3144 | 1/1 | 0.95 | 0.45 | 40,40,40,40 | 0 |
| 57 | MG | 2a | 1781 | 1/1 | 0.95 | 0.10 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3660 | 1/1 | 0.95 | 0.08 | 42,42,42,42 | 0 |
| 57 | MG | 2A | 3369 | 1/1 | 0.95 | 0.12 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3521 | 1/1 | 0.95 | 0.20 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3523 | 1/1 | 0.95 | 0.19 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3739 | 1/1 | 0.95 | 0.08 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3367 | 1/1 | 0.95 | 0.10 | 49,49,49,49 | 0 |
| 57 | MG | 1B | 213 | 1/1 | 0.95 | 0.16 | 50,50,50,50 | 0 |
| 57 | MG | 2a | 1789 | 1/1 | 0.95 | 0.19 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3310 | 1/1 | 0.95 | 0.22 | 32,32,32,32 | 0 |
| 57 | MG | 2A | 3745 | 1/1 | 0.95 | 0.08 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3746 | 1/1 | 0.95 | 0.14 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3431 | 1/1 | 0.95 | 0.14 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3878 | 1/1 | 0.95 | 0.06 | 26,26,26,26 | 0 |
| 57 | MG | 1A | 3879 | 1/1 | 0.95 | 0.10 | 30,30,30,30 | 0 |
| 57 | MG | 2A | 3752 | 1/1 | 0.95 | 0.07 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3432 | 1/1 | 0.95 | 0.12 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 3529 | 1/1 | 0.95 | 0.14 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3756 | 1/1 | 0.95 | 0.14 | 65,65,65,65 | 0 |
| 57 | MG | 1B | 225 | 1/1 | 0.95 | 0.20 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3882 | 1/1 | 0.95 | 0.12 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3434 | 1/1 | 0.95 | 0.12 | 34,34,34,34 | 0 |
| 57 | MG | 2A | 3762 | 1/1 | 0.95 | 0.07 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3764 | 1/1 | 0.95 | 0.08 | 52,52,52,52 | 0 |
| 57 | MG | 2a | 1806 | 1/1 | 0.95 | 0.07 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3765 | 1/1 | 0.95 | 0.06 | 35,35,35,35 | 0 |
| 57 | MG | 2A | 3111 | 1/1 | 0.95 | 0.20 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3679 | 1/1 | 0.95 | 0.08 | 18,18,18,18 | 0 |
| 57 | MG | 1A | 3680 | 1/1 | 0.95 | 0.09 | 21,21,21,21 | 0 |
| 57 | MG | 1a | 1704 | 1/1 | 0.95 | 0.08 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3682 | 1/1 | 0.95 | 0.06 | 16,16,16,16 | 0 |
| 57 | MG | 1a | 1707 | 1/1 | 0.95 | 0.32 | 56,56,56,56 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3117 | 1/1 | 0.95 | 0.14 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3394 | 1/1 | 0.95 | 0.12 | 28,28,28,28 | 0 |
| 57 | MG | 1A | 3265 | 1/1 | 0.95 | 0.11 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3314 | 1/1 | 0.95 | 0.23 | 50,50,50,50 | 0 |
| 57 | MG | 1B | 235 | 1/1 | 0.95 | 0.09 | 59,59,59,59 | 0 |
| 57 | MG | 1A | 3018 | 1/1 | 0.95 | 0.12 | 31,31,31,31 | 0 |
| 57 | MG | 2A | 3126 | 1/1 | 0.95 | 0.12 | 64,64,64,64 | 0 |
| 57 | MG | 2A | 3128 | 1/1 | 0.95 | 0.13 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3131 | 1/1 | 0.95 | 0.05 | 40,40,40,40 | 0 |
| 57 | MG | 2A | 3133 | 1/1 | 0.95 | 0.10 | 46,46,46,46 | 0 |
| 57 | MG | 2f | 201 | 1/1 | 0.95 | 0.11 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3695 | 1/1 | 0.95 | 0.08 | 22,22,22,22 | 0 |
| 57 | MG | 2A | 3407 | 1/1 | 0.95 | 0.30 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3409 | 1/1 | 0.95 | 0.19 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3899 | 1/1 | 0.95 | 0.15 | 60,60,60,60 | 0 |
| 57 | MG | 1D | 312 | 1/1 | 0.95 | 0.07 | 28,28,28,28 | 0 |
| 57 | MG | 2A | 3412 | 1/1 | 0.95 | 0.12 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3796 | 1/1 | 0.95 | 0.07 | 41,41,41,41 | 0 |
| 57 | MG | 2o | 101 | 1/1 | 0.95 | 0.14 | 54,54,54,54 | 0 |
| 57 | MG | 1E | 301 | 1/1 | 0.95 | 0.17 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3316 | 1/1 | 0.95 | 0.23 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3141 | 1/1 | 0.95 | 0.14 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3145 | 1/1 | 0.95 | 0.22 | 40,40,40,40 | 0 |
| 57 | MG | 2A | 3146 | 1/1 | 0.95 | 0.07 | 35,35,35,35 | 0 |
| 57 | MG | 1E | 305 | 1/1 | 0.95 | 0.10 | 40,40,40,40 | 0 |
| 57 | MG | 2A | 3805 | 1/1 | 0.95 | 0.11 | 48,48,48,48 | 0 |
| 57 | MG | 1E | 307 | 1/1 | 0.95 | 0.08 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3420 | 1/1 | 0.95 | 0.14 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3902 | 1/1 | 0.95 | 0.09 | 40,40,40,40 | 0 |
| 57 | MG | 2A | 3150 | 1/1 | 0.95 | 0.11 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3903 | 1/1 | 0.95 | 0.15 | 29,29,29,29 | 0 |
| 57 | MG | 2A | 3153 | 1/1 | 0.95 | 0.08 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3439 | 1/1 | 0.95 | 0.13 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3156 | 1/1 | 0.95 | 0.09 | 63,63,63,63 | 0 |
| 57 | MG | 2A | 3157 | 1/1 | 0.95 | 0.13 | 67,67,67,67 | 0 |
| 57 | MG | 1a | 1724 | 1/1 | 0.95 | 0.15 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3431 | 1/1 | 0.95 | 0.18 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3828 | 1/1 | 0.95 | 0.08 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3542 | 1/1 | 0.95 | 0.11 | 32,32,32,32 | 0 |
| 57 | MG | 1a | 1727 | 1/1 | 0.95 | 0.10 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3162 | 1/1 | 0.95 | 0.16 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3835 | 1/1 | 0.95 | 0.10 | 52,52,52,52 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 3103 | 1/1 | 0.96 | 0.05 | 56,56,56,56 | 0 |
| 57 | MG | 2E | 301 | 1/1 | 0.96 | 0.08 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 4075 | 1/1 | 0.96 | 0.06 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3266 | 1/1 | 0.96 | 0.11 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3547 | 1/1 | 0.96 | 0.06 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3868 | 1/1 | 0.96 | 0.15 | 31,31,31,31 | 0 |
| 57 | MG | 2A | 3020 | 1/1 | 0.96 | 0.11 | 42,42,42,42 | 0 |
| 57 | MG | 1a | 1634 | 1/1 | 0.96 | 0.15 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3869 | 1/1 | 0.96 | 0.28 | 39,39,39,39 | 0 |
| 57 | MG | 2A | 3024 | 1/1 | 0.96 | 0.08 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3163 | 1/1 | 0.96 | 0.09 | 42,42,42,42 | 0 |
| 57 | MG | 2A | 3026 | 1/1 | 0.96 | 0.04 | 42,42,42,42 | 0 |
| 57 | MG | 2F | 304 | 1/1 | 0.96 | 0.07 | 33,33,33,33 | 0 |
| 57 | MG | 2A | 3275 | 1/1 | 0.96 | 0.08 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3449 | 1/1 | 0.96 | 0.06 | 30,30,30,30 | 0 |
| 57 | MG | 2A | 3562 | 1/1 | 0.96 | 0.05 | 35,35,35,35 | 0 |
| 57 | MG | 2A | 3277 | 1/1 | 0.96 | 0.15 | 46,46,46,46 | 0 |
| 57 | MG | 1a | 1639 | 1/1 | 0.96 | 0.08 | 46,46,46,46 | 0 |
| 57 | MG | 1a | 1640 | 1/1 | 0.96 | 0.10 | 49,49,49,49 | 0 |
| 57 | MG | 2R | 201 | 1/1 | 0.96 | 0.04 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3567 | 1/1 | 0.96 | 0.11 | 46,46,46,46 | 0 |
| 57 | MG | 1a | 1641 | 1/1 | 0.96 | 0.06 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3691 | 1/1 | 0.96 | 0.06 | 23,23,23,23 | 0 |
| 57 | MG | 2U | 201 | 1/1 | 0.96 | 0.13 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3032 | 1/1 | 0.96 | 0.05 | 41,41,41,41 | 0 |
| 57 | MG | 2V | 201 | 1/1 | 0.96 | 0.07 | 62,62,62,62 | 0 |
| 57 | MG | 1A | 3085 | 1/1 | 0.96 | 0.15 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3547 | 1/1 | 0.96 | 0.06 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3698 | 1/1 | 0.96 | 0.07 | 22,22,22,22 | 0 |
| 57 | MG | 1A | 3378 | 1/1 | 0.96 | 0.04 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3288 | 1/1 | 0.96 | 0.11 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3703 | 1/1 | 0.96 | 0.09 | 19,19,19,19 | 0 |
| 57 | MG | 2A | 3039 | 1/1 | 0.96 | 0.05 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3132 | 1/1 | 0.96 | 0.05 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3455 | 1/1 | 0.96 | 0.06 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3293 | 1/1 | 0.96 | 0.16 | 62,62,62,62 | 0 |
| 57 | MG | 2A | 3042 | 1/1 | 0.96 | 0.10 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3585 | 1/1 | 0.96 | 0.05 | 41,41,41,41 | 0 |
| 57 | MG | 1B | 204 | 1/1 | 0.96 | 0.04 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 3214 | 1/1 | 0.96 | 0.05 | 36,36,36,36 | 0 |
| 57 | MG | 1A | 3709 | 1/1 | 0.96 | 0.07 | 24,24,24,24 | 0 |
| 57 | MG | 28 | 103 | 1/1 | 0.96 | 0.14 | 41,41,41,41 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1B | 207 | 1/1 | 0.96 | 0.27 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3554 | 1/1 | 0.96 | 0.06 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3886 | 1/1 | 0.96 | 0.06 | 33,33,33,33 | 0 |
| 57 | MG | 1a | 1662 | 1/1 | 0.96 | 0.05 | 33,33,33,33 | 0 |
| 57 | MG | 2A | 3051 | 1/1 | 0.96 | 0.07 | 60,60,60,60 | 0 |
| 57 | MG | 1a | 1663 | 1/1 | 0.96 | 0.14 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3598 | 1/1 | 0.96 | 0.12 | 59,59,59,59 | 0 |
| 57 | MG | 1A | 3457 | 1/1 | 0.96 | 0.05 | 46,46,46,46 | 0 |
| 57 | MG | 2a | 1609 | 1/1 | 0.96 | 0.10 | 61,61,61,61 | 0 |
| 57 | MG | 1A | 3888 | 1/1 | 0.96 | 0.07 | 23,23,23,23 | 0 |
| 57 | MG | 1A | 3712 | 1/1 | 0.96 | 0.12 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 3168 | 1/1 | 0.96 | 0.10 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 3714 | 1/1 | 0.96 | 0.05 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3604 | 1/1 | 0.96 | 0.13 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3558 | 1/1 | 0.96 | 0.16 | 36,36,36,36 | 0 |
| 57 | MG | 1A | 3897 | 1/1 | 0.96 | 0.04 | 31,31,31,31 | 0 |
| 57 | MG | 2a | 1617 | 1/1 | 0.96 | 0.08 | 66,66,66,66 | 0 |
| 57 | MG | 1A | 3134 | 1/1 | 0.96 | 0.07 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3384 | 1/1 | 0.96 | 0.04 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3172 | 1/1 | 0.96 | 0.23 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 3175 | 1/1 | 0.96 | 0.04 | 22,22,22,22 | 0 |
| 57 | MG | 1A | 3720 | 1/1 | 0.96 | 0.09 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3066 | 1/1 | 0.96 | 0.06 | 35,35,35,35 | 0 |
| 57 | MG | 1B | 227 | 1/1 | 0.96 | 0.09 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3322 | 1/1 | 0.96 | 0.08 | 85,85,85,85 | 0 |
| 57 | MG | 2A | 3323 | 1/1 | 0.96 | 0.15 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3323 | 1/1 | 0.96 | 0.20 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3273 | 1/1 | 0.96 | 0.13 | 24,24,24,24 | 0 |
| 57 | MG | 1A | 3390 | 1/1 | 0.96 | 0.13 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3571 | 1/1 | 0.96 | 0.06 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3073 | 1/1 | 0.96 | 0.10 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3274 | 1/1 | 0.96 | 0.45 | 35,35,35,35 | 0 |
| 57 | MG | 1a | 1683 | 1/1 | 0.96 | 0.12 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3911 | 1/1 | 0.96 | 0.08 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3731 | 1/1 | 0.96 | 0.12 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3110 | 1/1 | 0.96 | 0.25 | 33,33,33,33 | 0 |
| 57 | MG | 1A | 3574 | 1/1 | 0.96 | 0.07 | 40,40,40,40 | 0 |
| 57 | MG | 1D | 303 | 1/1 | 0.96 | 0.08 | 33,33,33,33 | 0 |
| 57 | MG | 1D | 306 | 1/1 | 0.96 | 0.09 | 35,35,35,35 | 0 |
| 57 | MG | 1a | 1692 | 1/1 | 0.96 | 0.17 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3179 | 1/1 | 0.96 | 0.11 | 38,38,38,38 | 0 |
| 57 | MG | 2a | 1643 | 1/1 | 0.96 | 0.07 | 48,48,48,48 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 3740 | 1/1 | 0.96 | 0.06 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3642 | 1/1 | 0.96 | 0.06 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3742 | 1/1 | 0.96 | 0.08 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3644 | 1/1 | 0.96 | 0.34 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3227 | 1/1 | 0.96 | 0.05 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3090 | 1/1 | 0.96 | 0.15 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3926 | 1/1 | 0.96 | 0.11 | 39,39,39,39 | 0 |
| 57 | MG | 2A | 3648 | 1/1 | 0.96 | 0.11 | 67,67,67,67 | 0 |
| 57 | MG | 1A | 3578 | 1/1 | 0.96 | 0.27 | 27,27,27,27 | 0 |
| 57 | MG | 1A | 3928 | 1/1 | 0.96 | 0.09 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3348 | 1/1 | 0.96 | 0.08 | 51,51,51,51 | 0 |
| 57 | MG | 1E | 310 | 1/1 | 0.96 | 0.08 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3929 | 1/1 | 0.96 | 0.06 | 55,55,55,55 | 0 |
| 57 | MG | 2a | 1657 | 1/1 | 0.96 | 0.14 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3930 | 1/1 | 0.96 | 0.08 | 44,44,44,44 | 0 |
| 57 | MG | 1E | 313 | 1/1 | 0.96 | 0.07 | 48,48,48,48 | 0 |
| 57 | MG | 1a | 1706 | 1/1 | 0.96 | 0.14 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3931 | 1/1 | 0.96 | 0.05 | 35,35,35,35 | 0 |
| 57 | MG | 2A | 3660 | 1/1 | 0.96 | 0.07 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3746 | 1/1 | 0.96 | 0.12 | 26,26,26,26 | 0 |
| 57 | MG | 2a | 1665 | 1/1 | 0.96 | 0.22 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3023 | 1/1 | 0.96 | 0.04 | 17,17,17,17 | 0 |
| 57 | MG | 2A | 3105 | 1/1 | 0.96 | 0.07 | 59,59,59,59 | 0 |
| 57 | MG | 2A | 3665 | 1/1 | 0.96 | 0.09 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3668 | 1/1 | 0.96 | 0.06 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3360 | 1/1 | 0.96 | 0.17 | 46,46,46,46 | 0 |
| 57 | MG | 1F | 307 | 1/1 | 0.96 | 0.09 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 3229 | 1/1 | 0.96 | 0.11 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3363 | 1/1 | 0.96 | 0.13 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3072 | 1/1 | 0.96 | 0.27 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3674 | 1/1 | 0.96 | 0.06 | 38,38,38,38 | 0 |
| 57 | MG | 1A | 3182 | 1/1 | 0.96 | 0.12 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3754 | 1/1 | 0.96 | 0.07 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3757 | 1/1 | 0.96 | 0.09 | 25,25,25,25 | 0 |
| 57 | MG | 1A | 3943 | 1/1 | 0.96 | 0.10 | 61,61,61,61 | 0 |
| 57 | MG | 1A | 3402 | 1/1 | 0.96 | 0.11 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3143 | 1/1 | 0.96 | 0.11 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3036 | 1/1 | 0.96 | 0.05 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3483 | 1/1 | 0.96 | 0.06 | 43,43,43,43 | 0 |
| 57 | MG | 2a | 1686 | 1/1 | 0.96 | 0.12 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3186 | 1/1 | 0.96 | 0.14 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3120 | 1/1 | 0.96 | 0.12 | 46,46,46,46 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3121 | 1/1 | 0.96 | 0.20 | 64,64,64,64 | 0 |
| 57 | MG | 1O | 204 | 1/1 | 0.96 | 0.07 | 63,63,63,63 | 0 |
| 57 | MG | 1P | 203 | 1/1 | 0.96 | 0.19 | 29,29,29,29 | 0 |
| 57 | MG | 1a | 1726 | 1/1 | 0.96 | 0.08 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3125 | 1/1 | 0.96 | 0.09 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3407 | 1/1 | 0.96 | 0.05 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3768 | 1/1 | 0.96 | 0.07 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3129 | 1/1 | 0.96 | 0.17 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3696 | 1/1 | 0.96 | 0.07 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3698 | 1/1 | 0.96 | 0.07 | 39,39,39,39 | 0 |
| 57 | MG | 2a | 1701 | 1/1 | 0.96 | 0.09 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3130 | 1/1 | 0.96 | 0.07 | 72,72,72,72 | 0 |
| 57 | MG | 1A | 3952 | 1/1 | 0.96 | 0.05 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3132 | 1/1 | 0.96 | 0.11 | 43,43,43,43 | 0 |
| 57 | MG | 1Q | 202 | 1/1 | 0.96 | 0.07 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3486 | 1/1 | 0.96 | 0.07 | 39,39,39,39 | 0 |
| 57 | MG | 1Q | 205 | 1/1 | 0.96 | 0.08 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3708 | 1/1 | 0.96 | 0.08 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3289 | 1/1 | 0.96 | 0.09 | 29,29,29,29 | 0 |
| 57 | MG | 2A | 3392 | 1/1 | 0.96 | 0.12 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3290 | 1/1 | 0.96 | 0.21 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3490 | 1/1 | 0.96 | 0.17 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3140 | 1/1 | 0.96 | 0.20 | 38,38,38,38 | 0 |
| 57 | MG | 1A | 3016 | 1/1 | 0.96 | 0.19 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3142 | 1/1 | 0.96 | 0.13 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3189 | 1/1 | 0.96 | 0.06 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3294 | 1/1 | 0.96 | 0.15 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 3240 | 1/1 | 0.96 | 0.07 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3783 | 1/1 | 0.96 | 0.06 | 26,26,26,26 | 0 |
| 57 | MG | 2A | 3405 | 1/1 | 0.96 | 0.09 | 41,41,41,41 | 0 |
| 57 | MG | 1S | 202 | 1/1 | 0.96 | 0.14 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3965 | 1/1 | 0.96 | 0.08 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3408 | 1/1 | 0.96 | 0.11 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3496 | 1/1 | 0.96 | 0.12 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3731 | 1/1 | 0.96 | 0.05 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3094 | 1/1 | 0.96 | 0.07 | 18,18,18,18 | 0 |
| 57 | MG | 2A | 3733 | 1/1 | 0.96 | 0.10 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3501 | 1/1 | 0.96 | 0.12 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 3971 | 1/1 | 0.96 | 0.05 | 30,30,30,30 | 0 |
| 57 | MG | 1a | 1753 | 1/1 | 0.96 | 0.10 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3615 | 1/1 | 0.96 | 0.04 | 48,48,48,48 | 0 |
| 57 | MG | 1V | 201 | 1/1 | 0.96 | 0.31 | 28,28,28,28 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 3974 | 1/1 | 0.96 | 0.07 | 29,29,29,29 | 0 |
| 57 | MG | 2A | 3161 | 1/1 | 0.96 | 0.23 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3743 | 1/1 | 0.96 | 0.13 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3975 | 1/1 | 0.96 | 0.08 | 28,28,28,28 | 0 |
| 57 | MG | 1A | 3616 | 1/1 | 0.96 | 0.09 | 23,23,23,23 | 0 |
| 57 | MG | 1W | 201 | 1/1 | 0.96 | 0.07 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 3350 | 1/1 | 0.96 | 0.14 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3748 | 1/1 | 0.96 | 0.06 | 42,42,42,42 | 0 |
| 57 | MG | 2A | 3422 | 1/1 | 0.96 | 0.08 | 53,53,53,53 | 0 |
| 57 | MG | 1a | 1761 | 1/1 | 0.96 | 0.05 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3351 | 1/1 | 0.96 | 0.15 | 48,48,48,48 | 0 |
| 57 | MG | 1W | 204 | 1/1 | 0.96 | 0.06 | 42,42,42,42 | 0 |
| 57 | MG | 2A | 3426 | 1/1 | 0.96 | 0.14 | 38,38,38,38 | 0 |
| 57 | MG | 1A | 3619 | 1/1 | 0.96 | 0.12 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3352 | 1/1 | 0.96 | 0.07 | 46,46,46,46 | 0 |
| 57 | MG | 1X | 105 | 1/1 | 0.96 | 0.05 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3172 | 1/1 | 0.96 | 0.10 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3192 | 1/1 | 0.96 | 0.23 | 25,25,25,25 | 0 |
| 57 | MG | 2A | 3432 | 1/1 | 0.96 | 0.18 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3007 | 1/1 | 0.96 | 0.08 | 26,26,26,26 | 0 |
| 57 | MG | 2A | 3766 | 1/1 | 0.96 | 0.08 | 47,47,47,47 | 0 |
| 57 | MG | 1a | 1770 | 1/1 | 0.96 | 0.10 | 65,65,65,65 | 0 |
| 57 | MG | 1a | 1771 | 1/1 | 0.96 | 0.05 | 67,67,67,67 | 0 |
| 57 | MG | 1A | 3630 | 1/1 | 0.96 | 0.09 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3355 | 1/1 | 0.96 | 0.08 | 46,46,46,46 | 0 |
| 57 | MG | 1Z | 302 | 1/1 | 0.96 | 0.06 | 57,57,57,57 | 0 |
| 57 | MG | 2a | 1761 | 1/1 | 0.96 | 0.09 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3440 | 1/1 | 0.96 | 0.10 | 42,42,42,42 | 0 |
| 57 | MG | 10 | 101 | 1/1 | 0.96 | 0.09 | 36,36,36,36 | 0 |
| 57 | MG | 1A | 3356 | 1/1 | 0.96 | 0.10 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3509 | 1/1 | 0.96 | 0.08 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3777 | 1/1 | 0.96 | 0.09 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3183 | 1/1 | 0.96 | 0.06 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3447 | 1/1 | 0.96 | 0.06 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3357 | 1/1 | 0.96 | 0.07 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3511 | 1/1 | 0.96 | 0.04 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3637 | 1/1 | 0.96 | 0.07 | 35,35,35,35 | 0 |
| 57 | MG | 1a | 1781 | 1/1 | 0.96 | 0.12 | 44,44,44,44 | 0 |
| 57 | MG | 2a | 1776 | 1/1 | 0.96 | 0.13 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3820 | 1/1 | 0.96 | 0.05 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 3194 | 1/1 | 0.96 | 0.05 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3787 | 1/1 | 0.96 | 0.08 | 61,61,61,61 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 4004 | 1/1 | 0.96 | 0.13 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3456 | 1/1 | 0.96 | 0.12 | 69,69,69,69 | 0 |
| 57 | MG | 2A | 3790 | 1/1 | 0.96 | 0.06 | 41,41,41,41 | 0 |
| 57 | MG | 1a | 1788 | 1/1 | 0.96 | 0.10 | 40,40,40,40 | 0 |
| 57 | MG | 11 | 102 | 1/1 | 0.96 | 0.08 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3514 | 1/1 | 0.96 | 0.15 | 33,33,33,33 | 0 |
| 57 | MG | 12 | 101 | 1/1 | 0.96 | 0.09 | 36,36,36,36 | 0 |
| 57 | MG | 1A | 3359 | 1/1 | 0.96 | 0.16 | 33,33,33,33 | 0 |
| 57 | MG | 2A | 3197 | 1/1 | 0.96 | 0.06 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3799 | 1/1 | 0.96 | 0.06 | 34,34,34,34 | 0 |
| 57 | MG | 13 | 102 | 1/1 | 0.96 | 0.10 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 4009 | 1/1 | 0.96 | 0.06 | 48,48,48,48 | 0 |
| 57 | MG | 13 | 104 | 1/1 | 0.96 | 0.09 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3517 | 1/1 | 0.96 | 0.14 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 3828 | 1/1 | 0.96 | 0.10 | 48,48,48,48 | 0 |
| 57 | MG | 15 | 103 | 1/1 | 0.96 | 0.15 | 36,36,36,36 | 0 |
| 57 | MG | 1A | 3829 | 1/1 | 0.96 | 0.11 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3809 | 1/1 | 0.96 | 0.05 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3812 | 1/1 | 0.96 | 0.16 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3151 | 1/1 | 0.96 | 0.05 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 3428 | 1/1 | 0.96 | 0.13 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3210 | 1/1 | 0.96 | 0.14 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3196 | 1/1 | 0.96 | 0.10 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3834 | 1/1 | 0.96 | 0.13 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3650 | 1/1 | 0.96 | 0.06 | 33,33,33,33 | 0 |
| 57 | MG | 1A | 3153 | 1/1 | 0.96 | 0.08 | 36,36,36,36 | 0 |
| 57 | MG | 17 | 103 | 1/1 | 0.96 | 0.10 | 31,31,31,31 | 0 |
| 57 | MG | 17 | 104 | 1/1 | 0.96 | 0.09 | 36,36,36,36 | 0 |
| 57 | MG | 1A | 3006 | 1/1 | 0.96 | 0.06 | 30,30,30,30 | 0 |
| 57 | MG | 2A | 3829 | 1/1 | 0.96 | 0.10 | 38,38,38,38 | 0 |
| 57 | MG | 1A | 3365 | 1/1 | 0.96 | 0.09 | 39,39,39,39 | 0 |
| 57 | MG | 2A | 3484 | 1/1 | 0.96 | 0.11 | 43,43,43,43 | 0 |
| 57 | MG | 2A | 3219 | 1/1 | 0.96 | 0.21 | 59,59,59,59 | 0 |
| 57 | MG | 2A | 3834 | 1/1 | 0.96 | 0.09 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3221 | 1/1 | 0.96 | 0.12 | 42,42,42,42 | 0 |
| 57 | MG | 18 | 103 | 1/1 | 0.96 | 0.19 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3489 | 1/1 | 0.96 | 0.10 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3839 | 1/1 | 0.96 | 0.10 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3081 | 1/1 | 0.96 | 0.12 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3842 | 1/1 | 0.96 | 0.07 | 46,46,46,46 | 0 |
| 57 | MG | 2a | 1823 | 1/1 | 0.96 | 0.13 | 59,59,59,59 | 0 |
| 57 | MG | 1n | 101 | 1/1 | 0.96 | 0.27 | 46,46,46,46 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 4037 | 1/1 | 0.96 | 0.06 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3230 | 1/1 | 0.96 | 0.15 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3306 | 1/1 | 0.96 | 0.13 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3009 | 1/1 | 0.96 | 0.07 | 32,32,32,32 | 0 |
| 57 | MG | 1r | 101 | 1/1 | 0.96 | 0.18 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3029 | 1/1 | 0.96 | 0.15 | 24,24,24,24 | 0 |
| 57 | MG | 2A | 3235 | 1/1 | 0.96 | 0.07 | 47,47,47,47 | 0 |
| 57 | MG | 1a | 1604 | 1/1 | 0.96 | 0.08 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3848 | 1/1 | 0.96 | 0.13 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3849 | 1/1 | 0.96 | 0.10 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 4046 | 1/1 | 0.96 | 0.07 | 34,34,34,34 | 0 |
| 57 | MG | 2A | 3508 | 1/1 | 0.96 | 0.20 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 4048 | 1/1 | 0.96 | 0.09 | 18,18,18,18 | 0 |
| 57 | MG | 1A | 4050 | 1/1 | 0.96 | 0.08 | 23,23,23,23 | 0 |
| 57 | MG | 2A | 3515 | 1/1 | 0.96 | 0.08 | 46,46,46,46 | 0 |
| 57 | MG | 1a | 1612 | 1/1 | 0.96 | 0.04 | 67,67,67,67 | 0 |
| 57 | MG | 1A | 3850 | 1/1 | 0.96 | 0.05 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 4053 | 1/1 | 0.96 | 0.08 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3530 | 1/1 | 0.96 | 0.09 | 36,36,36,36 | 0 |
| 57 | MG | 1A | 3309 | 1/1 | 0.96 | 0.08 | 34,34,34,34 | 0 |
| 57 | MG | 1a | 1618 | 1/1 | 0.96 | 0.04 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3668 | 1/1 | 0.96 | 0.06 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3160 | 1/1 | 0.96 | 0.07 | 28,28,28,28 | 0 |
| 57 | MG | 1A | 3129 | 1/1 | 0.96 | 0.12 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 3858 | 1/1 | 0.96 | 0.07 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3004 | 1/1 | 0.96 | 0.35 | 59,59,59,59 | 0 |
| 57 | MG | 1A | 3672 | 1/1 | 0.96 | 0.11 | 21,21,21,21 | 0 |
| 57 | MG | 1A | 3444 | 1/1 | 0.96 | 0.08 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3445 | 1/1 | 0.96 | 0.34 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3539 | 1/1 | 0.96 | 0.16 | 36,36,36,36 | 0 |
| 57 | MG | 1A | 3864 | 1/1 | 0.96 | 0.10 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 3374 | 1/1 | 0.96 | 0.06 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 4073 | 1/1 | 0.96 | 0.12 | 42,42,42,42 | 0 |
| 59 | ERY | 1A | 4087 | 51/51 | 0.96 | 0.10 | 25,42,54,59 | 0 |
| 57 | MG | 2A | 3014 | 1/1 | 0.96 | 0.11 | 41,41,41,41 | 0 |
| 60 | ZN | 14 | 102 | 1/1 | 0.96 | 0.11 | 118,118,118,118 | 0 |
| 60 | ZN | 24 | 501 | 1/1 | 0.96 | 0.13 | 137,137,137,137 | 0 |
| 60 | ZN | 2n | 501 | 1/1 | 0.96 | 0.05 | 94,94,94,94 | 0 |
| 57 | MG | 2E | 307 | 1/1 | 0.97 | 0.10 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3042 | 1/1 | 0.97 | 0.04 | 21,21,21,21 | 0 |
| 57 | MG | 1a | 1645 | 1/1 | 0.97 | 0.11 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3728 | 1/1 | 0.97 | 0.16 | 49,49,49,49 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 3198 | 1/1 | 0.97 | 0.06 | 29,29,29,29 | 0 |
| 57 | MG | 2A | 3043 | 1/1 | 0.97 | 0.07 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3590 | 1/1 | 0.97 | 0.10 | 24,24,24,24 | 0 |
| 57 | MG | 2F | 305 | 1/1 | 0.97 | 0.18 | 51,51,51,51 | 0 |
| 57 | MG | 2A | 3295 | 1/1 | 0.97 | 0.10 | 55,55,55,55 | 0 |
| 57 | MG | 1a | 1650 | 1/1 | 0.97 | 0.13 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3297 | 1/1 | 0.97 | 0.25 | 35,35,35,35 | 0 |
| 57 | MG | 1a | 1651 | 1/1 | 0.97 | 0.06 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3901 | 1/1 | 0.97 | 0.06 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3360 | 1/1 | 0.97 | 0.15 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3500 | 1/1 | 0.97 | 0.19 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3593 | 1/1 | 0.97 | 0.05 | 38,38,38,38 | 0 |
| 57 | MG | 2T | 201 | 1/1 | 0.97 | 0.09 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3159 | 1/1 | 0.97 | 0.08 | 36,36,36,36 | 0 |
| 57 | MG | 1a | 1658 | 1/1 | 0.97 | 0.10 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3741 | 1/1 | 0.97 | 0.06 | 34,34,34,34 | 0 |
| 57 | MG | 1B | 219 | 1/1 | 0.97 | 0.07 | 45,45,45,45 | 0 |
| 57 | MG | 1B | 221 | 1/1 | 0.97 | 0.04 | 40,40,40,40 | 0 |
| 57 | MG | 1B | 222 | 1/1 | 0.97 | 0.06 | 36,36,36,36 | 0 |
| 57 | MG | 1B | 223 | 1/1 | 0.97 | 0.05 | 30,30,30,30 | 0 |
| 57 | MG | 2X | 101 | 1/1 | 0.97 | 0.06 | 68,68,68,68 | 0 |
| 57 | MG | 1A | 3080 | 1/1 | 0.97 | 0.08 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3064 | 1/1 | 0.97 | 0.12 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3060 | 1/1 | 0.97 | 0.08 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3054 | 1/1 | 0.97 | 0.06 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3601 | 1/1 | 0.97 | 0.14 | 26,26,26,26 | 0 |
| 57 | MG | 1A | 3747 | 1/1 | 0.97 | 0.07 | 27,27,27,27 | 0 |
| 57 | MG | 23 | 102 | 1/1 | 0.97 | 0.06 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3602 | 1/1 | 0.97 | 0.24 | 34,34,34,34 | 0 |
| 57 | MG | 1B | 230 | 1/1 | 0.97 | 0.08 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3429 | 1/1 | 0.97 | 0.09 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3319 | 1/1 | 0.97 | 0.07 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3918 | 1/1 | 0.97 | 0.05 | 26,26,26,26 | 0 |
| 57 | MG | 1A | 3751 | 1/1 | 0.97 | 0.08 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3921 | 1/1 | 0.97 | 0.05 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3922 | 1/1 | 0.97 | 0.04 | 17,17,17,17 | 0 |
| 57 | MG | 1A | 3013 | 1/1 | 0.97 | 0.14 | 23,23,23,23 | 0 |
| 57 | MG | 1B | 237 | 1/1 | 0.97 | 0.05 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 3605 | 1/1 | 0.97 | 0.14 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3606 | 1/1 | 0.97 | 0.04 | 38,38,38,38 | 0 |
| 57 | MG | 1D | 304 | 1/1 | 0.97 | 0.15 | 47,47,47,47 | 0 |
| 57 | MG | 1D | 305 | 1/1 | 0.97 | 0.04 | 22,22,22,22 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3617 | 1/1 | 0.97 | 0.04 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3755 | 1/1 | 0.97 | 0.07 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3619 | 1/1 | 0.97 | 0.12 | 61,61,61,61 | 0 |
| 57 | MG | 1D | 308 | 1/1 | 0.97 | 0.17 | 31,31,31,31 | 0 |
| 57 | MG | 2A | 3080 | 1/1 | 0.97 | 0.10 | 46,46,46,46 | 0 |
| 57 | MG | 1D | 310 | 1/1 | 0.97 | 0.04 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3756 | 1/1 | 0.97 | 0.07 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3084 | 1/1 | 0.97 | 0.06 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3607 | 1/1 | 0.97 | 0.09 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3608 | 1/1 | 0.97 | 0.04 | 28,28,28,28 | 0 |
| 57 | MG | 1a | 1690 | 1/1 | 0.97 | 0.22 | 51,51,51,51 | 0 |
| 57 | MG | 1E | 302 | 1/1 | 0.97 | 0.16 | 35,35,35,35 | 0 |
| 57 | MG | 2A | 3634 | 1/1 | 0.97 | 0.13 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3635 | 1/1 | 0.97 | 0.10 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3609 | 1/1 | 0.97 | 0.06 | 24,24,24,24 | 0 |
| 57 | MG | 1A | 3610 | 1/1 | 0.97 | 0.06 | 34,34,34,34 | 0 |
| 57 | MG | 2a | 1623 | 1/1 | 0.97 | 0.10 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3091 | 1/1 | 0.97 | 0.04 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3344 | 1/1 | 0.97 | 0.06 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3252 | 1/1 | 0.97 | 0.06 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3253 | 1/1 | 0.97 | 0.08 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3433 | 1/1 | 0.97 | 0.14 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3350 | 1/1 | 0.97 | 0.26 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3936 | 1/1 | 0.97 | 0.08 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3937 | 1/1 | 0.97 | 0.05 | 44,44,44,44 | 0 |
| 57 | MG | 1a | 1699 | 1/1 | 0.97 | 0.07 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3938 | 1/1 | 0.97 | 0.08 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3767 | 1/1 | 0.97 | 0.06 | 35,35,35,35 | 0 |
| 57 | MG | 1F | 302 | 1/1 | 0.97 | 0.10 | 59,59,59,59 | 0 |
| 57 | MG | 1A | 3614 | 1/1 | 0.97 | 0.07 | 17,17,17,17 | 0 |
| 57 | MG | 1A | 3204 | 1/1 | 0.97 | 0.06 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3942 | 1/1 | 0.97 | 0.04 | 40,40,40,40 | 0 |
| 57 | MG | 1F | 308 | 1/1 | 0.97 | 0.28 | 35,35,35,35 | 0 |
| 57 | MG | 1F | 309 | 1/1 | 0.97 | 0.07 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3656 | 1/1 | 0.97 | 0.14 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3164 | 1/1 | 0.97 | 0.16 | 36,36,36,36 | 0 |
| 57 | MG | 2A | 3109 | 1/1 | 0.97 | 0.14 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3771 | 1/1 | 0.97 | 0.10 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3945 | 1/1 | 0.97 | 0.06 | 47,47,47,47 | 0 |
| 57 | MG | 1a | 1712 | 1/1 | 0.97 | 0.12 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3313 | 1/1 | 0.97 | 0.15 | 36,36,36,36 | 0 |
| 57 | MG | 1A | 3107 | 1/1 | 0.97 | 0.06 | 32,32,32,32 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1G | 204 | 1/1 | 0.97 | 0.10 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3667 | 1/1 | 0.97 | 0.12 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3515 | 1/1 | 0.97 | 0.13 | 33,33,33,33 | 0 |
| 57 | MG | 1A | 3109 | 1/1 | 0.97 | 0.33 | 31,31,31,31 | 0 |
| 57 | MG | 1N | 202 | 1/1 | 0.97 | 0.06 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3777 | 1/1 | 0.97 | 0.08 | 21,21,21,21 | 0 |
| 57 | MG | 1A | 3778 | 1/1 | 0.97 | 0.05 | 28,28,28,28 | 0 |
| 57 | MG | 1A | 3625 | 1/1 | 0.97 | 0.05 | 27,27,27,27 | 0 |
| 57 | MG | 1A | 3056 | 1/1 | 0.97 | 0.19 | 41,41,41,41 | 0 |
| 57 | MG | 1O | 203 | 1/1 | 0.97 | 0.08 | 56,56,56,56 | 0 |
| 57 | MG | 1A | 3954 | 1/1 | 0.97 | 0.10 | 24,24,24,24 | 0 |
| 57 | MG | 2A | 3677 | 1/1 | 0.97 | 0.16 | 38,38,38,38 | 0 |
| 57 | MG | 1A | 3782 | 1/1 | 0.97 | 0.06 | 28,28,28,28 | 0 |
| 57 | MG | 2a | 1662 | 1/1 | 0.97 | 0.09 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3127 | 1/1 | 0.97 | 0.06 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3381 | 1/1 | 0.97 | 0.07 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3211 | 1/1 | 0.97 | 0.14 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3784 | 1/1 | 0.97 | 0.07 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3787 | 1/1 | 0.97 | 0.04 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3628 | 1/1 | 0.97 | 0.06 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3960 | 1/1 | 0.97 | 0.10 | 54,54,54,54 | 0 |
| 57 | MG | 2A | 3387 | 1/1 | 0.97 | 0.07 | 51,51,51,51 | 0 |
| 57 | MG | 1a | 1733 | 1/1 | 0.97 | 0.05 | 39,39,39,39 | 0 |
| 57 | MG | 2a | 1672 | 1/1 | 0.97 | 0.08 | 64,64,64,64 | 0 |
| 57 | MG | 1A | 3629 | 1/1 | 0.97 | 0.09 | 31,31,31,31 | 0 |
| 57 | MG | 2A | 3689 | 1/1 | 0.97 | 0.08 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3794 | 1/1 | 0.97 | 0.08 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3963 | 1/1 | 0.97 | 0.06 | 68,68,68,68 | 0 |
| 57 | MG | 1A | 3169 | 1/1 | 0.97 | 0.12 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3393 | 1/1 | 0.97 | 0.11 | 51,51,51,51 | 0 |
| 57 | MG | 2a | 1679 | 1/1 | 0.97 | 0.15 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3442 | 1/1 | 0.97 | 0.08 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3797 | 1/1 | 0.97 | 0.05 | 17,17,17,17 | 0 |
| 57 | MG | 1A | 3443 | 1/1 | 0.97 | 0.06 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3397 | 1/1 | 0.97 | 0.21 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3633 | 1/1 | 0.97 | 0.07 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3262 | 1/1 | 0.97 | 0.27 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3802 | 1/1 | 0.97 | 0.06 | 40,40,40,40 | 0 |
| 57 | MG | 1a | 1746 | 1/1 | 0.97 | 0.10 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3973 | 1/1 | 0.97 | 0.07 | 21,21,21,21 | 0 |
| 57 | MG | 2A | 3707 | 1/1 | 0.97 | 0.10 | 53,53,53,53 | 0 |
| 57 | MG | 1U | 201 | 1/1 | 0.97 | 0.10 | 27,27,27,27 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3709 | 1/1 | 0.97 | 0.05 | 51,51,51,51 | 0 |
| 57 | MG | 1a | 1749 | 1/1 | 0.97 | 0.13 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3137 | 1/1 | 0.97 | 0.23 | 31,31,31,31 | 0 |
| 57 | MG | 2A | 3152 | 1/1 | 0.97 | 0.08 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3804 | 1/1 | 0.97 | 0.04 | 41,41,41,41 | 0 |
| 57 | MG | 1U | 205 | 1/1 | 0.97 | 0.04 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3805 | 1/1 | 0.97 | 0.06 | 38,38,38,38 | 0 |
| 57 | MG | 1U | 207 | 1/1 | 0.97 | 0.19 | 30,30,30,30 | 0 |
| 57 | MG | 1U | 208 | 1/1 | 0.97 | 0.28 | 33,33,33,33 | 0 |
| 57 | MG | 1A | 3806 | 1/1 | 0.97 | 0.05 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3138 | 1/1 | 0.97 | 0.06 | 27,27,27,27 | 0 |
| 57 | MG | 1A | 3982 | 1/1 | 0.97 | 0.06 | 23,23,23,23 | 0 |
| 57 | MG | 1A | 3034 | 1/1 | 0.97 | 0.33 | 31,31,31,31 | 0 |
| 57 | MG | 2A | 3724 | 1/1 | 0.97 | 0.14 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3725 | 1/1 | 0.97 | 0.05 | 45,45,45,45 | 0 |
| 57 | MG | 1V | 204 | 1/1 | 0.97 | 0.14 | 38,38,38,38 | 0 |
| 57 | MG | 1V | 205 | 1/1 | 0.97 | 0.05 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3984 | 1/1 | 0.97 | 0.14 | 17,17,17,17 | 0 |
| 57 | MG | 2A | 3729 | 1/1 | 0.97 | 0.09 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3527 | 1/1 | 0.97 | 0.06 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3216 | 1/1 | 0.97 | 0.08 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3069 | 1/1 | 0.97 | 0.06 | 26,26,26,26 | 0 |
| 57 | MG | 1A | 3643 | 1/1 | 0.97 | 0.06 | 39,39,39,39 | 0 |
| 57 | MG | 1a | 1767 | 1/1 | 0.97 | 0.10 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3113 | 1/1 | 0.97 | 0.08 | 31,31,31,31 | 0 |
| 57 | MG | 1W | 205 | 1/1 | 0.97 | 0.13 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3993 | 1/1 | 0.97 | 0.07 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 3645 | 1/1 | 0.97 | 0.06 | 22,22,22,22 | 0 |
| 57 | MG | 1X | 103 | 1/1 | 0.97 | 0.04 | 27,27,27,27 | 0 |
| 57 | MG | 1A | 3269 | 1/1 | 0.97 | 0.11 | 29,29,29,29 | 0 |
| 57 | MG | 2A | 3742 | 1/1 | 0.97 | 0.05 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3453 | 1/1 | 0.97 | 0.10 | 47,47,47,47 | 0 |
| 57 | MG | 2a | 1725 | 1/1 | 0.97 | 0.06 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3271 | 1/1 | 0.97 | 0.08 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3433 | 1/1 | 0.97 | 0.23 | 46,46,46,46 | 0 |
| 57 | MG | 1Y | 201 | 1/1 | 0.97 | 0.14 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3011 | 1/1 | 0.97 | 0.08 | 33,33,33,33 | 0 |
| 57 | MG | 1A | 3822 | 1/1 | 0.97 | 0.05 | 26,26,26,26 | 0 |
| 57 | MG | 1A | 3823 | 1/1 | 0.97 | 0.06 | 27,27,27,27 | 0 |
| 57 | MG | 1A | 3651 | 1/1 | 0.97 | 0.06 | 37,37,37,37 | 0 |
| 57 | MG | 10 | 102 | 1/1 | 0.97 | 0.17 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3753 | 1/1 | 0.97 | 0.07 | 38,38,38,38 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1a | 1782 | 1/1 | 0.97 | 0.05 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3825 | 1/1 | 0.97 | 0.04 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 4006 | 1/1 | 0.97 | 0.07 | 61,61,61,61 | 0 |
| 57 | MG | 2A | 3757 | 1/1 | 0.97 | 0.08 | 42,42,42,42 | 0 |
| 57 | MG | 10 | 105 | 1/1 | 0.97 | 0.05 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3146 | 1/1 | 0.97 | 0.06 | 27,27,27,27 | 0 |
| 57 | MG | 2A | 3446 | 1/1 | 0.97 | 0.10 | 27,27,27,27 | 0 |
| 57 | MG | 1A | 4008 | 1/1 | 0.97 | 0.09 | 44,44,44,44 | 0 |
| 57 | MG | 1a | 1791 | 1/1 | 0.97 | 0.08 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3537 | 1/1 | 0.97 | 0.13 | 26,26,26,26 | 0 |
| 57 | MG | 2A | 3193 | 1/1 | 0.97 | 0.05 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3655 | 1/1 | 0.97 | 0.06 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3538 | 1/1 | 0.97 | 0.13 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 4013 | 1/1 | 0.97 | 0.09 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 4015 | 1/1 | 0.97 | 0.06 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3771 | 1/1 | 0.97 | 0.08 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3331 | 1/1 | 0.97 | 0.04 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 4018 | 1/1 | 0.97 | 0.04 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3831 | 1/1 | 0.97 | 0.16 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 3221 | 1/1 | 0.97 | 0.04 | 31,31,31,31 | 0 |
| 57 | MG | 1a | 1803 | 1/1 | 0.97 | 0.05 | 38,38,38,38 | 0 |
| 57 | MG | 1A | 3541 | 1/1 | 0.97 | 0.11 | 35,35,35,35 | 0 |
| 57 | MG | 2A | 3463 | 1/1 | 0.97 | 0.12 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3204 | 1/1 | 0.97 | 0.04 | 52,52,52,52 | 0 |
| 57 | MG | 2A | 3780 | 1/1 | 0.97 | 0.04 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3278 | 1/1 | 0.97 | 0.09 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3334 | 1/1 | 0.97 | 0.12 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3092 | 1/1 | 0.97 | 0.19 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3209 | 1/1 | 0.97 | 0.19 | 62,62,62,62 | 0 |
| 57 | MG | 1A | 3666 | 1/1 | 0.97 | 0.05 | 20,20,20,20 | 0 |
| 57 | MG | 1A | 3462 | 1/1 | 0.97 | 0.09 | 51,51,51,51 | 0 |
| 57 | MG | 15 | 105 | 1/1 | 0.97 | 0.30 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 4030 | 1/1 | 0.97 | 0.11 | 47,47,47,47 | 0 |
| 57 | MG | 2a | 1770 | 1/1 | 0.97 | 0.05 | 68,68,68,68 | 0 |
| 57 | MG | 1A | 3336 | 1/1 | 0.97 | 0.07 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3464 | 1/1 | 0.97 | 0.25 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 4035 | 1/1 | 0.97 | 0.09 | 16,16,16,16 | 0 |
| 57 | MG | 2A | 3792 | 1/1 | 0.97 | 0.05 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3465 | 1/1 | 0.97 | 0.15 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3225 | 1/1 | 0.97 | 0.06 | 21,21,21,21 | 0 |
| 57 | MG | 1A | 3398 | 1/1 | 0.97 | 0.10 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3220 | 1/1 | 0.97 | 0.13 | 49,49,49,49 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1l | 202 | 1/1 | 0.97 | 0.11 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3676 | 1/1 | 0.97 | 0.10 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3553 | 1/1 | 0.97 | 0.15 | 36,36,36,36 | 0 |
| 57 | MG | 2A | 3224 | 1/1 | 0.97 | 0.10 | 34,34,34,34 | 0 |
| 57 | MG | 2A | 3485 | 1/1 | 0.97 | 0.07 | 44,44,44,44 | 0 |
| 57 | MG | 18 | 101 | 1/1 | 0.97 | 0.10 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3804 | 1/1 | 0.97 | 0.06 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3399 | 1/1 | 0.97 | 0.15 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3116 | 1/1 | 0.97 | 0.06 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3684 | 1/1 | 0.97 | 0.09 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3184 | 1/1 | 0.97 | 0.06 | 35,35,35,35 | 0 |
| 57 | MG | 2A | 3810 | 1/1 | 0.97 | 0.12 | 38,38,38,38 | 0 |
| 57 | MG | 2a | 1792 | 1/1 | 0.97 | 0.12 | 33,33,33,33 | 0 |
| 57 | MG | 2A | 3811 | 1/1 | 0.97 | 0.07 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 4049 | 1/1 | 0.97 | 0.09 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3119 | 1/1 | 0.97 | 0.09 | 28,28,28,28 | 0 |
| 57 | MG | 1A | 3560 | 1/1 | 0.97 | 0.09 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3693 | 1/1 | 0.97 | 0.07 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 4054 | 1/1 | 0.97 | 0.04 | 25,25,25,25 | 0 |
| 57 | MG | 1w | 106 | 1/1 | 0.97 | 0.06 | 66,66,66,66 | 0 |
| 57 | MG | 1A | 4056 | 1/1 | 0.97 | 0.07 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3824 | 1/1 | 0.97 | 0.07 | 26,26,26,26 | 0 |
| 57 | MG | 1A | 3561 | 1/1 | 0.97 | 0.34 | 34,34,34,34 | 0 |
| 57 | MG | 1a | 1606 | 1/1 | 0.97 | 0.16 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3150 | 1/1 | 0.97 | 0.16 | 26,26,26,26 | 0 |
| 57 | MG | 1x | 106 | 1/1 | 0.97 | 0.11 | 51,51,51,51 | 0 |
| 57 | MG | 1x | 108 | 1/1 | 0.97 | 0.08 | 22,22,22,22 | 0 |
| 57 | MG | 1a | 1608 | 1/1 | 0.97 | 0.15 | 56,56,56,56 | 0 |
| 57 | MG | 2A | 3831 | 1/1 | 0.97 | 0.12 | 55,55,55,55 | 0 |
| 57 | MG | 1A | 3696 | 1/1 | 0.97 | 0.04 | 25,25,25,25 | 0 |
| 57 | MG | 1A | 3697 | 1/1 | 0.97 | 0.05 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3509 | 1/1 | 0.97 | 0.07 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 4063 | 1/1 | 0.97 | 0.09 | 53,53,53,53 | 0 |
| 57 | MG | 2a | 1813 | 1/1 | 0.97 | 0.14 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3862 | 1/1 | 0.97 | 0.05 | 29,29,29,29 | 0 |
| 57 | MG | 2A | 3512 | 1/1 | 0.97 | 0.17 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3187 | 1/1 | 0.97 | 0.04 | 27,27,27,27 | 0 |
| 57 | MG | 1A | 3699 | 1/1 | 0.97 | 0.06 | 24,24,24,24 | 0 |
| 57 | MG | 1A | 3700 | 1/1 | 0.97 | 0.04 | 25,25,25,25 | 0 |
| 57 | MG | 2A | 3251 | 1/1 | 0.97 | 0.23 | 45,45,45,45 | 0 |
| 57 | MG | 1a | 1616 | 1/1 | 0.97 | 0.04 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3405 | 1/1 | 0.97 | 0.09 | 38,38,38,38 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3007 | 1/1 | 0.97 | 0.06 | 40,40,40,40 | 0 |
| 57 | MG | 2a | 1824 | 1/1 | 0.97 | 0.13 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3008 | 1/1 | 0.97 | 0.06 | 47,47,47,47 | 0 |
| 57 | MG | 2A | 3256 | 1/1 | 0.97 | 0.05 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3702 | 1/1 | 0.97 | 0.15 | 26,26,26,26 | 0 |
| 57 | MG | 2A | 3525 | 1/1 | 0.97 | 0.08 | 28,28,28,28 | 0 |
| 57 | MG | 1a | 1619 | 1/1 | 0.97 | 0.07 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3343 | 1/1 | 0.97 | 0.05 | 42,42,42,42 | 0 |
| 57 | MG | 2A | 3528 | 1/1 | 0.97 | 0.08 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3476 | 1/1 | 0.97 | 0.06 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 3705 | 1/1 | 0.97 | 0.07 | 27,27,27,27 | 0 |
| 57 | MG | 2A | 3532 | 1/1 | 0.97 | 0.14 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3012 | 1/1 | 0.97 | 0.11 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3707 | 1/1 | 0.97 | 0.04 | 17,17,17,17 | 0 |
| 57 | MG | 2A | 3535 | 1/1 | 0.97 | 0.09 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3408 | 1/1 | 0.97 | 0.05 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3152 | 1/1 | 0.97 | 0.07 | 36,36,36,36 | 0 |
| 57 | MG | 2A | 3538 | 1/1 | 0.97 | 0.08 | 40,40,40,40 | 0 |
| 57 | MG | 2A | 3019 | 1/1 | 0.97 | 0.06 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3039 | 1/1 | 0.97 | 0.11 | 28,28,28,28 | 0 |
| 57 | MG | 1A | 3481 | 1/1 | 0.97 | 0.08 | 26,26,26,26 | 0 |
| 57 | MG | 2A | 3022 | 1/1 | 0.97 | 0.05 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3482 | 1/1 | 0.97 | 0.13 | 28,28,28,28 | 0 |
| 57 | MG | 1A | 3191 | 1/1 | 0.97 | 0.20 | 27,27,27,27 | 0 |
| 57 | MG | 1A | 3349 | 1/1 | 0.97 | 0.09 | 53,53,53,53 | 0 |
| 57 | MG | 1A | 3154 | 1/1 | 0.97 | 0.06 | 36,36,36,36 | 0 |
| 57 | MG | 2A | 3552 | 1/1 | 0.97 | 0.06 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3238 | 1/1 | 0.97 | 0.20 | 33,33,33,33 | 0 |
| 57 | MG | 1A | 3487 | 1/1 | 0.97 | 0.07 | 39,39,39,39 | 0 |
| 57 | MG | 1a | 1635 | 1/1 | 0.97 | 0.22 | 67,67,67,67 | 0 |
| 57 | MG | 1A | 4086 | 1/1 | 0.97 | 0.20 | 36,36,36,36 | 0 |
| 57 | MG | 1A | 3239 | 1/1 | 0.97 | 0.15 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3076 | 1/1 | 0.97 | 0.04 | 33,33,33,33 | 0 |
| 57 | MG | 2D | 305 | 1/1 | 0.97 | 0.21 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3123 | 1/1 | 0.97 | 0.06 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3283 | 1/1 | 0.97 | 0.07 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3034 | 1/1 | 0.97 | 0.07 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 3242 | 1/1 | 0.97 | 0.13 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3019 | 1/1 | 0.97 | 0.19 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3895 | 1/1 | 0.97 | 0.04 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3244 | 1/1 | 0.97 | 0.09 | 33,33,33,33 | 0 |
| 57 | MG | 2A | 3347 | 1/1 | 0.98 | 0.09 | 42,42,42,42 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 4052 | 1/1 | 0.98 | 0.05 | 39,39,39,39 | 0 |
| 57 | MG | 2A | 3349 | 1/1 | 0.98 | 0.07 | 39,39,39,39 | 0 |
| 57 | MG | 1a | 1675 | 1/1 | 0.98 | 0.15 | 44,44,44,44 | 0 |
| 57 | MG | 1T | 203 | 1/1 | 0.98 | 0.05 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3385 | 1/1 | 0.98 | 0.06 | 25,25,25,25 | 0 |
| 57 | MG | 1U | 202 | 1/1 | 0.98 | 0.10 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 3106 | 1/1 | 0.98 | 0.05 | 27,27,27,27 | 0 |
| 57 | MG | 2A | 3760 | 1/1 | 0.98 | 0.05 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 4055 | 1/1 | 0.98 | 0.07 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 3920 | 1/1 | 0.98 | 0.05 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3763 | 1/1 | 0.98 | 0.03 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3807 | 1/1 | 0.98 | 0.07 | 24,24,24,24 | 0 |
| 57 | MG | 1A | 3017 | 1/1 | 0.98 | 0.12 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3550 | 1/1 | 0.98 | 0.18 | 33,33,33,33 | 0 |
| 57 | MG | 1a | 1685 | 1/1 | 0.98 | 0.11 | 35,35,35,35 | 0 |
| 57 | MG | 2A | 3555 | 1/1 | 0.98 | 0.07 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 3035 | 1/1 | 0.98 | 0.15 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3170 | 1/1 | 0.98 | 0.04 | 42,42,42,42 | 0 |
| 57 | MG | 1U | 211 | 1/1 | 0.98 | 0.35 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3621 | 1/1 | 0.98 | 0.07 | 24,24,24,24 | 0 |
| 57 | MG | 1A | 3813 | 1/1 | 0.98 | 0.05 | 50,50,50,50 | 0 |
| 57 | MG | 2A | 3561 | 1/1 | 0.98 | 0.06 | 35,35,35,35 | 0 |
| 57 | MG | 1V | 203 | 1/1 | 0.98 | 0.21 | 28,28,28,28 | 0 |
| 57 | MG | 1A | 3493 | 1/1 | 0.98 | 0.06 | 30,30,30,30 | 0 |
| 57 | MG | 2A | 3564 | 1/1 | 0.98 | 0.07 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3017 | 1/1 | 0.98 | 0.07 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3624 | 1/1 | 0.98 | 0.10 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 3293 | 1/1 | 0.98 | 0.12 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 3207 | 1/1 | 0.98 | 0.07 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 3818 | 1/1 | 0.98 | 0.04 | 38,38,38,38 | 0 |
| 57 | MG | 1A | 3295 | 1/1 | 0.98 | 0.18 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3557 | 1/1 | 0.98 | 0.26 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3140 | 1/1 | 0.98 | 0.15 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 3559 | 1/1 | 0.98 | 0.07 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3499 | 1/1 | 0.98 | 0.16 | 24,24,24,24 | 0 |
| 57 | MG | 1A | 3209 | 1/1 | 0.98 | 0.25 | 31,31,31,31 | 0 |
| 57 | MG | 1a | 1703 | 1/1 | 0.98 | 0.08 | 52,52,52,52 | 0 |
| 57 | MG | 1X | 102 | 1/1 | 0.98 | 0.20 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3003 | 1/1 | 0.98 | 0.03 | 27,27,27,27 | 0 |
| 57 | MG | 1X | 104 | 1/1 | 0.98 | 0.05 | 33,33,33,33 | 0 |
| 57 | MG | 2A | 3581 | 1/1 | 0.98 | 0.10 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3563 | 1/1 | 0.98 | 0.14 | 31,31,31,31 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3795 | 1/1 | 0.98 | 0.07 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3396 | 1/1 | 0.98 | 0.04 | 30,30,30,30 | 0 |
| 57 | MG | 2A | 3207 | 1/1 | 0.98 | 0.05 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3173 | 1/1 | 0.98 | 0.32 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3174 | 1/1 | 0.98 | 0.05 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 3727 | 1/1 | 0.98 | 0.12 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3638 | 1/1 | 0.98 | 0.03 | 21,21,21,21 | 0 |
| 57 | MG | 1A | 3729 | 1/1 | 0.98 | 0.11 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3142 | 1/1 | 0.98 | 0.09 | 35,35,35,35 | 0 |
| 57 | MG | 2a | 1693 | 1/1 | 0.98 | 0.04 | 57,57,57,57 | 0 |
| 57 | MG | 2A | 3594 | 1/1 | 0.98 | 0.06 | 47,47,47,47 | 0 |
| 57 | MG | 1A | 3568 | 1/1 | 0.98 | 0.08 | 40,40,40,40 | 0 |
| 57 | MG | 2A | 3806 | 1/1 | 0.98 | 0.09 | 39,39,39,39 | 0 |
| 57 | MG | 1B | 202 | 1/1 | 0.98 | 0.11 | 38,38,38,38 | 0 |
| 57 | MG | 1A | 3451 | 1/1 | 0.98 | 0.06 | 35,35,35,35 | 0 |
| 57 | MG | 2a | 1699 | 1/1 | 0.98 | 0.06 | 60,60,60,60 | 0 |
| 57 | MG | 1A | 3733 | 1/1 | 0.98 | 0.05 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3734 | 1/1 | 0.98 | 0.03 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 3176 | 1/1 | 0.98 | 0.08 | 25,25,25,25 | 0 |
| 57 | MG | 1A | 3177 | 1/1 | 0.98 | 0.08 | 24,24,24,24 | 0 |
| 57 | MG | 1A | 3840 | 1/1 | 0.98 | 0.04 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3015 | 1/1 | 0.98 | 0.14 | 26,26,26,26 | 0 |
| 57 | MG | 1A | 3646 | 1/1 | 0.98 | 0.06 | 17,17,17,17 | 0 |
| 57 | MG | 1A | 3843 | 1/1 | 0.98 | 0.12 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3819 | 1/1 | 0.98 | 0.13 | 62,62,62,62 | 0 |
| 57 | MG | 2A | 3606 | 1/1 | 0.98 | 0.11 | 55,55,55,55 | 0 |
| 57 | MG | 2A | 3225 | 1/1 | 0.98 | 0.27 | 32,32,32,32 | 0 |
| 57 | MG | 2A | 3404 | 1/1 | 0.98 | 0.07 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 3010 | 1/1 | 0.98 | 0.06 | 34,34,34,34 | 0 |
| 57 | MG | 1a | 1728 | 1/1 | 0.98 | 0.04 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3259 | 1/1 | 0.98 | 0.12 | 33,33,33,33 | 0 |
| 57 | MG | 12 | 102 | 1/1 | 0.98 | 0.11 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3744 | 1/1 | 0.98 | 0.04 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3847 | 1/1 | 0.98 | 0.06 | 18,18,18,18 | 0 |
| 57 | MG | 1A | 3512 | 1/1 | 0.98 | 0.13 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3145 | 1/1 | 0.98 | 0.08 | 26,26,26,26 | 0 |
| 57 | MG | 1A | 3087 | 1/1 | 0.98 | 0.11 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3040 | 1/1 | 0.98 | 0.12 | 27,27,27,27 | 0 |
| 57 | MG | 15 | 101 | 1/1 | 0.98 | 0.10 | 33,33,33,33 | 0 |
| 57 | MG | 15 | 102 | 1/1 | 0.98 | 0.08 | 33,33,33,33 | 0 |
| 57 | MG | 2A | 3621 | 1/1 | 0.98 | 0.07 | 44,44,44,44 | 0 |
| 57 | MG | 1B | 220 | 1/1 | 0.98 | 0.05 | 43,43,43,43 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 3653 | 1/1 | 0.98 | 0.07 | 40,40,40,40 | 0 |
| 57 | MG | 2A | 3624 | 1/1 | 0.98 | 0.10 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3853 | 1/1 | 0.98 | 0.08 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 3969 | 1/1 | 0.98 | 0.06 | 34,34,34,34 | 0 |
| 57 | MG | 2A | 3627 | 1/1 | 0.98 | 0.12 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3067 | 1/1 | 0.98 | 0.05 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3845 | 1/1 | 0.98 | 0.11 | 39,39,39,39 | 0 |
| 57 | MG | 2A | 3629 | 1/1 | 0.98 | 0.08 | 35,35,35,35 | 0 |
| 57 | MG | 2A | 3847 | 1/1 | 0.98 | 0.08 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 3750 | 1/1 | 0.98 | 0.04 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3089 | 1/1 | 0.98 | 0.07 | 36,36,36,36 | 0 |
| 57 | MG | 1A | 3856 | 1/1 | 0.98 | 0.06 | 39,39,39,39 | 0 |
| 57 | MG | 2A | 3851 | 1/1 | 0.98 | 0.07 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3311 | 1/1 | 0.98 | 0.08 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3656 | 1/1 | 0.98 | 0.06 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3222 | 1/1 | 0.98 | 0.11 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3977 | 1/1 | 0.98 | 0.05 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3978 | 1/1 | 0.98 | 0.06 | 55,55,55,55 | 0 |
| 57 | MG | 17 | 106 | 1/1 | 0.98 | 0.03 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3658 | 1/1 | 0.98 | 0.05 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3980 | 1/1 | 0.98 | 0.04 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 3583 | 1/1 | 0.98 | 0.04 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 3031 | 1/1 | 0.98 | 0.06 | 40,40,40,40 | 0 |
| 57 | MG | 2a | 1749 | 1/1 | 0.98 | 0.14 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3081 | 1/1 | 0.98 | 0.04 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3758 | 1/1 | 0.98 | 0.04 | 23,23,23,23 | 0 |
| 57 | MG | 1A | 3032 | 1/1 | 0.98 | 0.12 | 21,21,21,21 | 0 |
| 57 | MG | 1B | 238 | 1/1 | 0.98 | 0.04 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3760 | 1/1 | 0.98 | 0.14 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3226 | 1/1 | 0.98 | 0.16 | 30,30,30,30 | 0 |
| 57 | MG | 2A | 3650 | 1/1 | 0.98 | 0.05 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3867 | 1/1 | 0.98 | 0.04 | 25,25,25,25 | 0 |
| 57 | MG | 2A | 3263 | 1/1 | 0.98 | 0.22 | 36,36,36,36 | 0 |
| 57 | MG | 1A | 3522 | 1/1 | 0.98 | 0.15 | 26,26,26,26 | 0 |
| 57 | MG | 1A | 3664 | 1/1 | 0.98 | 0.04 | 32,32,32,32 | 0 |
| 57 | MG | 1D | 307 | 1/1 | 0.98 | 0.11 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3070 | 1/1 | 0.98 | 0.12 | 19,19,19,19 | 0 |
| 57 | MG | 2a | 1763 | 1/1 | 0.98 | 0.10 | 58,58,58,58 | 0 |
| 57 | MG | 2A | 3448 | 1/1 | 0.98 | 0.05 | 45,45,45,45 | 0 |
| 57 | MG | 1D | 309 | 1/1 | 0.98 | 0.07 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3995 | 1/1 | 0.98 | 0.07 | 28,28,28,28 | 0 |
| 57 | MG | 2D | 302 | 1/1 | 0.98 | 0.10 | 41,41,41,41 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3270 | 1/1 | 0.98 | 0.06 | 44,44,44,44 | 0 |
| 57 | MG | 1A | 3871 | 1/1 | 0.98 | 0.13 | 24,24,24,24 | 0 |
| 57 | MG | 2A | 3096 | 1/1 | 0.98 | 0.05 | 46,46,46,46 | 0 |
| 57 | MG | 2A | 3663 | 1/1 | 0.98 | 0.03 | 33,33,33,33 | 0 |
| 57 | MG | 1A | 3872 | 1/1 | 0.98 | 0.08 | 36,36,36,36 | 0 |
| 57 | MG | 2a | 1773 | 1/1 | 0.98 | 0.10 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3071 | 1/1 | 0.98 | 0.06 | 22,22,22,22 | 0 |
| 57 | MG | 2A | 3666 | 1/1 | 0.98 | 0.09 | 52,52,52,52 | 0 |
| 57 | MG | 1A | 3667 | 1/1 | 0.98 | 0.04 | 26,26,26,26 | 0 |
| 57 | MG | 1A | 3043 | 1/1 | 0.98 | 0.08 | 26,26,26,26 | 0 |
| 57 | MG | 2A | 3458 | 1/1 | 0.98 | 0.03 | 28,28,28,28 | 0 |
| 57 | MG | 1E | 304 | 1/1 | 0.98 | 0.07 | 25,25,25,25 | 0 |
| 57 | MG | 1A | 3272 | 1/1 | 0.98 | 0.06 | 41,41,41,41 | 0 |
| 57 | MG | 1E | 306 | 1/1 | 0.98 | 0.04 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3670 | 1/1 | 0.98 | 0.08 | 27,27,27,27 | 0 |
| 57 | MG | 1A | 3073 | 1/1 | 0.98 | 0.05 | 14,14,14,14 | 0 |
| 57 | MG | 1E | 309 | 1/1 | 0.98 | 0.12 | 16,16,16,16 | 0 |
| 57 | MG | 1A | 3369 | 1/1 | 0.98 | 0.05 | 24,24,24,24 | 0 |
| 57 | MG | 2F | 306 | 1/1 | 0.98 | 0.11 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3594 | 1/1 | 0.98 | 0.12 | 21,21,21,21 | 0 |
| 57 | MG | 1A | 3595 | 1/1 | 0.98 | 0.04 | 25,25,25,25 | 0 |
| 57 | MG | 1A | 3596 | 1/1 | 0.98 | 0.07 | 33,33,33,33 | 0 |
| 57 | MG | 1A | 3883 | 1/1 | 0.98 | 0.04 | 33,33,33,33 | 0 |
| 57 | MG | 1a | 1786 | 1/1 | 0.98 | 0.04 | 43,43,43,43 | 0 |
| 57 | MG | 1a | 1787 | 1/1 | 0.98 | 0.08 | 50,50,50,50 | 0 |
| 57 | MG | 1A | 3057 | 1/1 | 0.98 | 0.12 | 25,25,25,25 | 0 |
| 57 | MG | 2R | 202 | 1/1 | 0.98 | 0.05 | 44,44,44,44 | 0 |
| 57 | MG | 1F | 303 | 1/1 | 0.98 | 0.19 | 34,34,34,34 | 0 |
| 57 | MG | 2R | 204 | 1/1 | 0.98 | 0.05 | 36,36,36,36 | 0 |
| 57 | MG | 1A | 3276 | 1/1 | 0.98 | 0.05 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3125 | 1/1 | 0.98 | 0.22 | 31,31,31,31 | 0 |
| 57 | MG | 2A | 3118 | 1/1 | 0.98 | 0.06 | 47,47,47,47 | 0 |
| 57 | MG | 1a | 1793 | 1/1 | 0.98 | 0.05 | 55,55,55,55 | 0 |
| 57 | MG | 1F | 306 | 1/1 | 0.98 | 0.04 | 42,42,42,42 | 0 |
| 57 | MG | 1A | 3780 | 1/1 | 0.98 | 0.06 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3683 | 1/1 | 0.98 | 0.05 | 24,24,24,24 | 0 |
| 57 | MG | 1A | 4017 | 1/1 | 0.98 | 0.05 | 39,39,39,39 | 0 |
| 57 | MG | 1F | 310 | 1/1 | 0.98 | 0.08 | 36,36,36,36 | 0 |
| 57 | MG | 2Y | 201 | 1/1 | 0.98 | 0.13 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3889 | 1/1 | 0.98 | 0.06 | 54,54,54,54 | 0 |
| 57 | MG | 1A | 3126 | 1/1 | 0.98 | 0.22 | 34,34,34,34 | 0 |
| 57 | MG | 2A | 3697 | 1/1 | 0.98 | 0.06 | 20,20,20,20 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1a | 1801 | 1/1 | 0.98 | 0.09 | 59,59,59,59 | 0 |
| 57 | MG | 1F | 313 | 1/1 | 0.98 | 0.04 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 3685 | 1/1 | 0.98 | 0.06 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 3893 | 1/1 | 0.98 | 0.04 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3279 | 1/1 | 0.98 | 0.07 | 39,39,39,39 | 0 |
| 57 | MG | 2a | 1815 | 1/1 | 0.98 | 0.12 | 45,45,45,45 | 0 |
| 57 | MG | 26 | 101 | 1/1 | 0.98 | 0.14 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3491 | 1/1 | 0.98 | 0.08 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3785 | 1/1 | 0.98 | 0.06 | 48,48,48,48 | 0 |
| 57 | MG | 2A | 3706 | 1/1 | 0.98 | 0.09 | 55,55,55,55 | 0 |
| 57 | MG | 1H | 201 | 1/1 | 0.98 | 0.13 | 41,41,41,41 | 0 |
| 57 | MG | 1A | 3786 | 1/1 | 0.98 | 0.10 | 18,18,18,18 | 0 |
| 57 | MG | 1A | 3044 | 1/1 | 0.98 | 0.08 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 3788 | 1/1 | 0.98 | 0.06 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 4028 | 1/1 | 0.98 | 0.04 | 22,22,22,22 | 0 |
| 57 | MG | 2a | 1825 | 1/1 | 0.98 | 0.09 | 53,53,53,53 | 0 |
| 57 | MG | 2A | 3138 | 1/1 | 0.98 | 0.18 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 4029 | 1/1 | 0.98 | 0.03 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3714 | 1/1 | 0.98 | 0.05 | 42,42,42,42 | 0 |
| 57 | MG | 2A | 3500 | 1/1 | 0.98 | 0.08 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3535 | 1/1 | 0.98 | 0.08 | 24,24,24,24 | 0 |
| 57 | MG | 1a | 1649 | 1/1 | 0.98 | 0.05 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3790 | 1/1 | 0.98 | 0.03 | 39,39,39,39 | 0 |
| 57 | MG | 2A | 3143 | 1/1 | 0.98 | 0.22 | 44,44,44,44 | 0 |
| 57 | MG | 2A | 3320 | 1/1 | 0.98 | 0.06 | 45,45,45,45 | 0 |
| 57 | MG | 2A | 3506 | 1/1 | 0.98 | 0.14 | 45,45,45,45 | 0 |
| 57 | MG | 2l | 203 | 1/1 | 0.98 | 0.06 | 63,63,63,63 | 0 |
| 57 | MG | 2A | 3144 | 1/1 | 0.98 | 0.20 | 46,46,46,46 | 0 |
| 57 | MG | 1A | 4033 | 1/1 | 0.98 | 0.11 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3791 | 1/1 | 0.98 | 0.04 | 35,35,35,35 | 0 |
| 57 | MG | 1P | 202 | 1/1 | 0.98 | 0.15 | 28,28,28,28 | 0 |
| 57 | MG | 1a | 1654 | 1/1 | 0.98 | 0.13 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3792 | 1/1 | 0.98 | 0.08 | 20,20,20,20 | 0 |
| 57 | MG | 1A | 3692 | 1/1 | 0.98 | 0.07 | 22,22,22,22 | 0 |
| 57 | MG | 1A | 3426 | 1/1 | 0.98 | 0.16 | 36,36,36,36 | 0 |
| 57 | MG | 1A | 3033 | 1/1 | 0.98 | 0.05 | 28,28,28,28 | 0 |
| 57 | MG | 1Q | 201 | 1/1 | 0.98 | 0.12 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3099 | 1/1 | 0.98 | 0.10 | 24,24,24,24 | 0 |
| 57 | MG | 1Q | 203 | 1/1 | 0.98 | 0.09 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3046 | 1/1 | 0.98 | 0.08 | 27,27,27,27 | 0 |
| 57 | MG | 1A | 4042 | 1/1 | 0.98 | 0.03 | 26,26,26,26 | 0 |
| 57 | MG | 1A | 4043 | 1/1 | 0.98 | 0.04 | 46,46,46,46 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 2A | 3738 | 1/1 | 0.98 | 0.05 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3102 | 1/1 | 0.98 | 0.05 | 38,38,38,38 | 0 |
| 57 | MG | 1A | 3078 | 1/1 | 0.98 | 0.05 | 18,18,18,18 | 0 |
| 57 | MG | 1A | 3381 | 1/1 | 0.98 | 0.06 | 62,62,62,62 | 0 |
| 57 | MG | 1A | 4047 | 1/1 | 0.98 | 0.11 | 11,11,11,11 | 0 |
| 57 | MG | 1x | 103 | 1/1 | 0.98 | 0.14 | 52,52,52,52 | 0 |
| 57 | MG | 1R | 204 | 1/1 | 0.98 | 0.11 | 26,26,26,26 | 0 |
| 57 | MG | 2A | 3530 | 1/1 | 0.98 | 0.05 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3914 | 1/1 | 0.98 | 0.06 | 36,36,36,36 | 0 |
| 57 | MG | 1A | 3047 | 1/1 | 0.98 | 0.04 | 28,28,28,28 | 0 |
| 57 | MG | 1x | 107 | 1/1 | 0.98 | 0.05 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3165 | 1/1 | 0.98 | 0.05 | 30,30,30,30 | 0 |
| 57 | MG | 1A | 3105 | 1/1 | 0.98 | 0.07 | 27,27,27,27 | 0 |
| 60 | ZN | 25 | 102 | 1/1 | 0.98 | 0.03 | 80,80,80,80 | 0 |
| 57 | MG | 2A | 3751 | 1/1 | 0.98 | 0.07 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 4064 | 1/1 | 0.99 | 0.03 | 33,33,33,33 | 0 |
| 57 | MG | 1A | 3764 | 1/1 | 0.99 | 0.06 | 28,28,28,28 | 0 |
| 57 | MG | 1A | 4011 | 1/1 | 0.99 | 0.04 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 3640 | 1/1 | 0.99 | 0.09 | 34,34,34,34 | 0 |
| 57 | MG | 2A | 3545 | 1/1 | 0.99 | 0.05 | 37,37,37,37 | 0 |
| 57 | MG | 2A | 3546 | 1/1 | 0.99 | 0.03 | 25,25,25,25 | 0 |
| 57 | MG | 2A | 3702 | 1/1 | 0.99 | 0.07 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3117 | 1/1 | 0.99 | 0.06 | 49,49,49,49 | 0 |
| 57 | MG | 2A | 3548 | 1/1 | 0.99 | 0.05 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 4014 | 1/1 | 0.99 | 0.06 | 26,26,26,26 | 0 |
| 57 | MG | 1a | 1740 | 1/1 | 0.99 | 0.04 | 27,27,27,27 | 0 |
| 57 | MG | 1a | 1741 | 1/1 | 0.99 | 0.08 | 25,25,25,25 | 0 |
| 57 | MG | 1A | 3118 | 1/1 | 0.99 | 0.06 | 22,22,22,22 | 0 |
| 57 | MG | 2A | 3154 | 1/1 | 0.99 | 0.08 | 31,31,31,31 | 0 |
| 57 | MG | 1D | 302 | 1/1 | 0.99 | 0.04 | 26,26,26,26 | 0 |
| 57 | MG | 2A | 3632 | 1/1 | 0.99 | 0.02 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3577 | 1/1 | 0.99 | 0.03 | 38,38,38,38 | 0 |
| 57 | MG | 2A | 3092 | 1/1 | 0.99 | 0.04 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3620 | 1/1 | 0.99 | 0.08 | 37,37,37,37 | 0 |
| 57 | MG | 2D | 303 | 1/1 | 0.99 | 0.05 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 3344 | 1/1 | 0.99 | 0.13 | 41,41,41,41 | 0 |
| 57 | MG | 2A | 3716 | 1/1 | 0.99 | 0.04 | 33,33,33,33 | 0 |
| 57 | MG | 1A | 3970 | 1/1 | 0.99 | 0.08 | 22,22,22,22 | 0 |
| 57 | MG | 1A | 3622 | 1/1 | 0.99 | 0.05 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3735 | 1/1 | 0.99 | 0.03 | 26,26,26,26 | 0 |
| 57 | MG | 1A | 3736 | 1/1 | 0.99 | 0.05 | 20,20,20,20 | 0 |
| 57 | MG | 1A | 3737 | 1/1 | 0.99 | 0.07 | 66,66,66,66 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 3775 | 1/1 | 0.99 | 0.03 | 51,51,51,51 | 0 |
| 57 | MG | 1A | 3275 | 1/1 | 0.99 | 0.27 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 4026 | 1/1 | 0.99 | 0.07 | 24,24,24,24 | 0 |
| 57 | MG | 1A | 3892 | 1/1 | 0.99 | 0.06 | 43,43,43,43 | 0 |
| 57 | MG | 1A | 3498 | 1/1 | 0.99 | 0.13 | 26,26,26,26 | 0 |
| 57 | MG | 1A | 3082 | 1/1 | 0.99 | 0.18 | 33,33,33,33 | 0 |
| 57 | MG | 1A | 3674 | 1/1 | 0.99 | 0.12 | 17,17,17,17 | 0 |
| 57 | MG | 2A | 3572 | 1/1 | 0.99 | 0.03 | 36,36,36,36 | 0 |
| 57 | MG | 1T | 202 | 1/1 | 0.99 | 0.14 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3038 | 1/1 | 0.99 | 0.13 | 18,18,18,18 | 0 |
| 57 | MG | 1A | 4032 | 1/1 | 0.99 | 0.05 | 20,20,20,20 | 0 |
| 57 | MG | 1A | 3026 | 1/1 | 0.99 | 0.05 | 39,39,39,39 | 0 |
| 57 | MG | 1A | 3677 | 1/1 | 0.99 | 0.10 | 20,20,20,20 | 0 |
| 57 | MG | 2A | 3815 | 1/1 | 0.99 | 0.07 | 26,26,26,26 | 0 |
| 57 | MG | 1A | 3678 | 1/1 | 0.99 | 0.06 | 21,21,21,21 | 0 |
| 57 | MG | 1A | 3985 | 1/1 | 0.99 | 0.08 | 30,30,30,30 | 0 |
| 57 | MG | 2A | 3818 | 1/1 | 0.99 | 0.08 | 21,21,21,21 | 0 |
| 57 | MG | 1A | 3986 | 1/1 | 0.99 | 0.07 | 32,32,32,32 | 0 |
| 57 | MG | 2A | 3820 | 1/1 | 0.99 | 0.09 | 57,57,57,57 | 0 |
| 57 | MG | 1A | 3330 | 1/1 | 0.99 | 0.05 | 40,40,40,40 | 0 |
| 57 | MG | 2A | 3822 | 1/1 | 0.99 | 0.05 | 45,45,45,45 | 0 |
| 57 | MG | 1A | 3014 | 1/1 | 0.99 | 0.04 | 30,30,30,30 | 0 |
| 57 | MG | 2A | 3583 | 1/1 | 0.99 | 0.06 | 34,34,34,34 | 0 |
| 57 | MG | 1A | 3681 | 1/1 | 0.99 | 0.04 | 24,24,24,24 | 0 |
| 57 | MG | 1A | 4041 | 1/1 | 0.99 | 0.02 | 23,23,23,23 | 0 |
| 57 | MG | 17 | 101 | 1/1 | 0.99 | 0.03 | 27,27,27,27 | 0 |
| 57 | MG | 17 | 102 | 1/1 | 0.99 | 0.05 | 25,25,25,25 | 0 |
| 57 | MG | 2A | 3513 | 1/1 | 0.99 | 0.04 | 32,32,32,32 | 0 |
| 57 | MG | 2A | 3589 | 1/1 | 0.99 | 0.02 | 27,27,27,27 | 0 |
| 57 | MG | 2A | 3590 | 1/1 | 0.99 | 0.06 | 31,31,31,31 | 0 |
| 57 | MG | 2A | 3514 | 1/1 | 0.99 | 0.08 | 58,58,58,58 | 0 |
| 57 | MG | 1A | 3133 | 1/1 | 0.99 | 0.19 | 35,35,35,35 | 0 |
| 57 | MG | 1A | 3991 | 1/1 | 0.99 | 0.04 | 23,23,23,23 | 0 |
| 57 | MG | 1A | 3086 | 1/1 | 0.99 | 0.10 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 3906 | 1/1 | 0.99 | 0.02 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 3994 | 1/1 | 0.99 | 0.05 | 21,21,21,21 | 0 |
| 57 | MG | 1a | 1715 | 1/1 | 0.99 | 0.04 | 49,49,49,49 | 0 |
| 57 | MG | 1A | 3021 | 1/1 | 0.99 | 0.03 | 20,20,20,20 | 0 |
| 57 | MG | 1A | 3136 | 1/1 | 0.99 | 0.03 | 32,32,32,32 | 0 |
| 57 | MG | 1A | 3686 | 1/1 | 0.99 | 0.03 | 26,26,26,26 | 0 |
| 57 | MG | 1A | 3998 | 1/1 | 0.99 | 0.07 | 25,25,25,25 | 0 |
| 57 | MG | 1A | 3237 | 1/1 | 0.99 | 0.08 | 23,23,23,23 | 0 |

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| Mol | Type | Chain | Res | Atoms | RSCC | RSR | B-factors(\AA^2) | Q<0.9 |
|-----|------|-------|------|-------|------|------|-----------------------------|-------|
| 57 | MG | 1A | 3688 | 1/1 | 0.99 | 0.09 | 22,22,22,22 | 0 |
| 57 | MG | 1A | 3689 | 1/1 | 0.99 | 0.05 | 17,17,17,17 | 0 |
| 57 | MG | 1A | 3913 | 1/1 | 0.99 | 0.05 | 30,30,30,30 | 0 |
| 57 | MG | 1W | 206 | 1/1 | 0.99 | 0.07 | 24,24,24,24 | 0 |
| 57 | MG | 1A | 3721 | 1/1 | 0.99 | 0.04 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 3197 | 1/1 | 0.99 | 0.05 | 18,18,18,18 | 0 |
| 57 | MG | 2A | 3464 | 1/1 | 0.99 | 0.06 | 48,48,48,48 | 0 |
| 57 | MG | 1A | 3723 | 1/1 | 0.99 | 0.07 | 40,40,40,40 | 0 |
| 57 | MG | 1A | 3224 | 1/1 | 0.99 | 0.05 | 20,20,20,20 | 0 |
| 57 | MG | 1A | 3270 | 1/1 | 0.99 | 0.21 | 26,26,26,26 | 0 |
| 57 | MG | 1A | 3100 | 1/1 | 0.99 | 0.04 | 31,31,31,31 | 0 |
| 57 | MG | 1A | 4061 | 1/1 | 0.99 | 0.11 | 41,41,41,41 | 0 |
| 60 | ZN | 1Y | 203 | 1/1 | 0.99 | 0.03 | 69,69,69,69 | 0 |
| 57 | MG | 1A | 4062 | 1/1 | 0.99 | 0.08 | 33,33,33,33 | 0 |
| 60 | ZN | 15 | 110 | 1/1 | 0.99 | 0.04 | 43,43,43,43 | 0 |
| 60 | ZN | 16 | 103 | 1/1 | 0.99 | 0.03 | 41,41,41,41 | 0 |
| 60 | ZN | 19 | 501 | 1/1 | 0.99 | 0.09 | 56,56,56,56 | 0 |
| 60 | ZN | 1n | 104 | 1/1 | 0.99 | 0.03 | 55,55,55,55 | 0 |
| 60 | ZN | 2Y | 202 | 1/1 | 0.99 | 0.03 | 103,103,103,103 | 0 |
| 57 | MG | 1A | 3108 | 1/1 | 0.99 | 0.08 | 25,25,25,25 | 0 |
| 57 | MG | 2A | 3540 | 1/1 | 0.99 | 0.07 | 32,32,32,32 | 0 |
| 60 | ZN | 26 | 102 | 1/1 | 0.99 | 0.04 | 59,59,59,59 | 0 |
| 60 | ZN | 29 | 501 | 1/1 | 0.99 | 0.03 | 66,66,66,66 | 0 |
| 57 | MG | 2A | 3695 | 1/1 | 0.99 | 0.03 | 45,45,45,45 | 0 |
| 61 | SF4 | 1d | 302 | 8/8 | 0.99 | 0.04 | 46,48,60,61 | 0 |
| 61 | SF4 | 2d | 303 | 8/8 | 0.99 | 0.03 | 57,71,80,82 | 0 |
| 57 | MG | 2A | 3550 | 1/1 | 1.00 | 0.10 | 29,29,29,29 | 0 |
| 57 | MG | 1A | 3924 | 1/1 | 1.00 | 0.05 | 17,17,17,17 | 0 |
| 57 | MG | 1P | 201 | 1/1 | 1.00 | 0.10 | 26,26,26,26 | 0 |
| 57 | MG | 1A | 3798 | 1/1 | 1.00 | 0.04 | 22,22,22,22 | 0 |
| 57 | MG | 1a | 1790 | 1/1 | 1.00 | 0.03 | 37,37,37,37 | 0 |
| 57 | MG | 1A | 3905 | 1/1 | 1.00 | 0.03 | 28,28,28,28 | 0 |

6.5 Other polymers [i](#)

There are no such residues in this entry.