



Full wwPDB EM Validation Report ⓘ

Oct 1, 2025 – 02:59 PM JST

PDB ID : 9V26 / pdb_00009v26
EMDB ID : EMD-64718
Title : Cryo- EM structure of 75S ribosome with A/P- & P/E- tRNAs from Entamoeba histolytica bound to antibiotic paromomycin
Authors : Sharma, S.; Mishra, S.; Gourinath, S.; Kaushal, P.S.
Deposited on : 2025-05-19
Resolution : 3.10 Å(reported)
Based on initial model : .

This is a Full wwPDB EM Validation Report for a publicly released PDB entry.

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

A user guide is available at

<https://www.wwpdb.org/validation/2017/EMValidationReportHelp>
with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

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

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

EMDB validation analysis : 0.0.1.dev129
Mogul : 1.8.5 (274361), CSD as541be (2020)
MolProbity : 4-5-2 with Phenix2.0
buster-report : 1.1.7 (2018)
Percentile statistics : 20231227.v01 (using entries in the PDB archive December 27th 2023)
EM percentile statistics : 202505.v01 (Using data in the EMDB archive up until May 2025)
MapQ : 1.9.13
Ideal geometry (proteins) : Engh & Huber (2001)
Ideal geometry (DNA, RNA) : Parkinson et al. (1996)
Validation Pipeline (wwPDB-VP) : 2.46

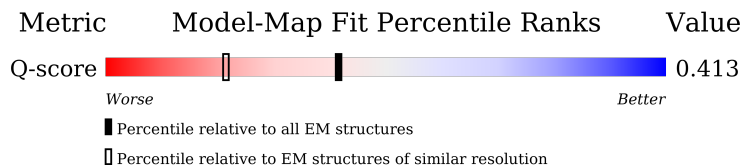
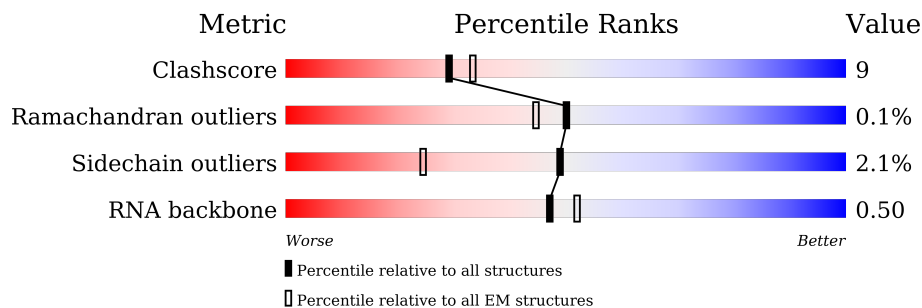
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

The reported resolution of this entry is 3.10 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.















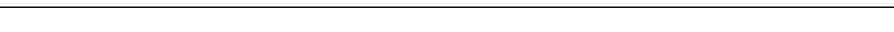

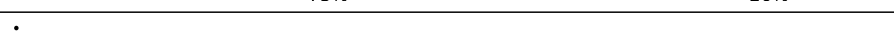

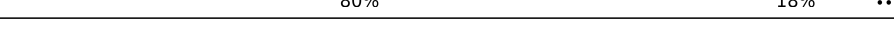








| Metric | Whole archive (#Entries) | EM structures (#Entries) | Similar EM resolution (#Entries, resolution range(Å)) |
|-----------------------|-----------------------------|-----------------------------|--|
| Clashscore | 210492 | 15764 | - |
| Ramachandran outliers | 207382 | 16835 | - |
| Sidechain outliers | 206894 | 16415 | - |
| RNA backbone | 6643 | 2191 | - |
| Q-score | - | 25397 | 14724 (2.60 - 3.60) |

The table below summarises the geometric issues observed across the polymeric chains and their fit to the map. The red, orange, yellow and green segments of the 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 EM map (all-atom inclusion $< 40\%$). The numeric value is given above the bar.

| Mol | Chain | Length | Quality of chain |
|-----|-------|--------|------------------|
| 1 | 1A | 3503 | |
| 2 | 1B | 155 | |
| 3 | 1C | 117 | |


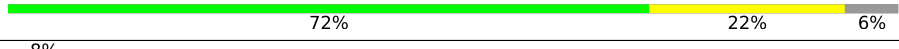
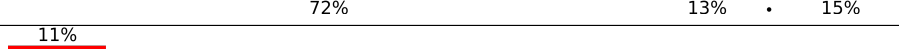
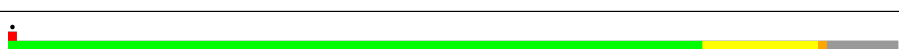



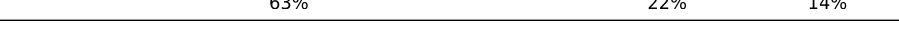



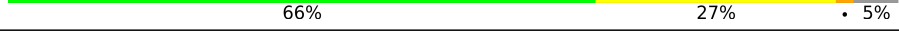




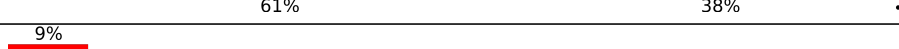
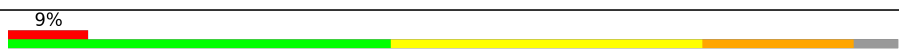

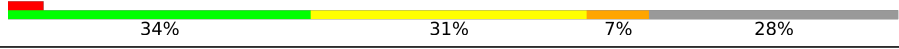




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| Mol | Chain | Length | Quality of chain |
|-----|-------|--------|--|
| 4 | ID | 257 |  |
| 5 | IE | 402 |  |
| 6 | IF | 431 |  |
| 7 | IG | 286 |  |
| 8 | IH | 204 |  |
| 9 | II | 230 |  |
| 10 | IJ | 286 |  |
| 11 | IK | 197 |  |
| 12 | IL | 210 |  |
| 13 | IM | 174 |  |
| 14 | IN | 291 |  |
| 15 | IO | 205 |  |
| 16 | IP | 135 |  |
| 17 | IQ | 205 |  |
| 18 | IR | 179 |  |
| 19 | IS | 168 |  |
| 20 | IT | 173 |  |
| 21 | IU | 198 |  |
| 22 | IV | 166 |  |
| 23 | IW | 137 |  |
| 24 | IX | 140 |  |
| 25 | IY | 121 |  |
| 26 | IZ | 163 |  |
| 27 | la | 213 |  |
| 28 | lb | 139 |  |

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| Mol | Chain | Length | Quality of chain |
|-----|-------|--------|--|
| 29 | lc | 149 |  |
| 30 | ld | 64 |  |
| 31 | le | 109 |  |
| 32 | lf | 150 |  |
| 33 | lg | 134 |  |
| 34 | lh | 137 |  |
| 35 | li | 122 |  |
| 36 | lj | 108 |  |
| 37 | lk | 104 |  |
| 38 | ll | 77 |  |
| 39 | lm | 93 |  |
| 40 | ln | 77 |  |
| 41 | lo | 51 |  |
| 42 | lp | 56 |  |
| 43 | lq | 98 |  |
| 44 | sA | 137 |  |
| 45 | sB | 144 |  |
| 46 | sC | 84 |  |
| 47 | sD | 69 |  |
| 48 | sE | 56 |  |
| 49 | sI | 76 |  |
| 50 | sJ | 77 |  |
| 51 | sK | 10 |  |
| 52 | sa | 1947 |  |
| 53 | sc | 255 | |

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| Mol | Chain | Length | Quality of chain |
|-----|-------|--------|------------------|
| 54 | sd | 244 | |
| 55 | se | 256 | |
| 56 | sf | 326 | |
| 57 | sg | 206 | |
| 58 | sh | 266 | |
| 59 | si | 201 | |
| 60 | sj | 237 | |
| 61 | sk | 185 | |
| 62 | sl | 127 | |
| 63 | sm | 156 | |
| 64 | so | 151 | |
| 65 | sp | 146 | |
| 66 | sq | 144 | |
| 67 | sr | 130 | |
| 68 | ss | 158 | |
| 69 | st | 117 | |
| 70 | su | 155 | |
| 71 | sv | 155 | |
| 72 | sw | 118 | |
| 73 | sx | 86 | |
| 74 | sy | 141 | |

2 Entry composition

There are 75 unique types of molecules in this entry. The entry contains 178778 atoms, of which 0 are hydrogens and 0 are deuteriums.

In the tables below, 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 25S rRNA.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-------|-------|-------|------|---------|-------|
| 1 | 1A | 3142 | Total | C | N | O | P | 0 | 0 |
| | | | 67153 | 30106 | 12192 | 21713 | 3142 | | |

- Molecule 2 is a RNA chain called 5.8S rRNA.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|------|-----|---------|-------|
| 2 | 1B | 145 | Total | C | N | O | P | 0 | 0 |
| | | | 3097 | 1390 | 560 | 1002 | 145 | | |

- Molecule 3 is a RNA chain called 5S rRNA.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|-----|---------|-------|
| 3 | 1C | 117 | Total | C | N | O | P | 0 | 0 |
| | | | 2477 | 1108 | 425 | 827 | 117 | | |

- Molecule 4 is a protein called Large ribosomal subunit protein uL2.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 4 | 1D | 246 | Total | C | N | O | S | 0 | 0 |
| | | | 1881 | 1165 | 382 | 326 | 8 | | |

- Molecule 5 is a protein called 60S ribosomal protein L3, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|----|---------|-------|
| 5 | 1E | 387 | Total | C | N | O | S | 0 | 0 |
| | | | 3076 | 1956 | 578 | 527 | 15 | | |

- Molecule 6 is a protein called 60S ribosomal protein L4, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|----|---------|-------|
| 6 | 1F | 425 | Total | C | N | O | S | 0 | 0 |
| | | | 3281 | 2091 | 624 | 552 | 14 | | |

- Molecule 7 is a protein called 60S ribosomal protein L5, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 7 | lG | 278 | Total | C | N | O | S | 0 | 0 |
| | | | 2209 | 1412 | 399 | 390 | 8 | | |

- Molecule 8 is a protein called Large ribosomal subunit protein eL6.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 8 | lH | 203 | Total | C | N | O | S | 0 | 0 |
| | | | 1608 | 1054 | 272 | 278 | 4 | | |

- Molecule 9 is a protein called 60S ribosomal protein L7, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 9 | lI | 210 | Total | C | N | O | S | 0 | 0 |
| | | | 1658 | 1067 | 301 | 282 | 8 | | |

- Molecule 10 is a protein called 60S ribosomal protein L7a.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 10 | lJ | 201 | Total | C | N | O | S | 0 | 0 |
| | | | 1640 | 1058 | 302 | 275 | 5 | | |

- Molecule 11 is a protein called 60S ribosomal protein L9, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 11 | lK | 193 | Total | C | N | O | S | 0 | 0 |
| | | | 1538 | 974 | 279 | 279 | 6 | | |

- Molecule 12 is a protein called Ribosomal protein L10, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|----|---------|-------|
| 12 | lL | 200 | Total | C | N | O | S | 0 | 0 |
| | | | 1597 | 1017 | 302 | 264 | 14 | | |

- Molecule 13 is a protein called 60S ribosomal protein L11, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 13 | lM | 170 | Total | C | N | O | S | 0 | 0 |
| | | | 1350 | 857 | 243 | 245 | 5 | | |

- Molecule 14 is a protein called 60S ribosomal protein L13, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 14 | lN | 267 | Total | C | N | O | S | 0 | 0 |
| | | | 2130 | 1358 | 412 | 352 | 8 | | |

- Molecule 15 is a protein called 60S ribosomal protein L13, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 15 | lO | 204 | Total | C | N | O | S | 0 | 0 |
| | | | 1616 | 1030 | 302 | 275 | 9 | | |

- Molecule 16 is a protein called 60S ribosomal protein L14, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 16 | lP | 130 | Total | C | N | O | S | 0 | 0 |
| | | | 1021 | 654 | 188 | 175 | 4 | | |

- Molecule 17 is a protein called Ribosomal protein L15.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 17 | lQ | 204 | Total | C | N | O | S | 0 | 0 |
| | | | 1676 | 1051 | 356 | 264 | 5 | | |

- Molecule 18 is a protein called 60S ribosomal protein L17, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 18 | lR | 158 | Total | C | N | O | S | 0 | 0 |
| | | | 1232 | 779 | 238 | 210 | 5 | | |

- Molecule 19 is a protein called 60S ribosomal protein L18, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 19 | lS | 167 | Total | C | N | O | S | 0 | 0 |
| | | | 1321 | 835 | 258 | 219 | 9 | | |

- Molecule 20 is a protein called 60S ribosomal protein L18a.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 20 | lT | 173 | Total | C | N | O | S | 0 | 0 |
| | | | 1413 | 910 | 259 | 235 | 9 | | |

- Molecule 21 is a protein called Ribosomal protein L19.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 21 | IU | 150 | Total | C | N | O | S | 0 | 0 |
| | | | 1235 | 787 | 246 | 197 | 5 | | |

- Molecule 22 is a protein called 60S ribosomal protein L21, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 22 | IV | 165 | Total | C | N | O | S | 0 | 0 |
| | | | 1320 | 846 | 254 | 217 | 3 | | |

- Molecule 23 is a protein called Large ribosomal subunit protein eL22.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 23 | IW | 93 | Total | C | N | O | S | 0 | 0 |
| | | | 763 | 493 | 132 | 133 | 5 | | |

- Molecule 24 is a protein called 60S ribosomal protein L23, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 24 | IX | 133 | Total | C | N | O | S | 0 | 0 |
| | | | 1015 | 629 | 196 | 182 | 8 | | |

- Molecule 25 is a protein called Ribosomal protein L23A, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 25 | IY | 116 | Total | C | N | O | S | 0 | 0 |
| | | | 926 | 597 | 166 | 159 | 4 | | |

- Molecule 26 is a protein called 60S ribosomal protein L24, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 26 | IZ | 57 | Total | C | N | O | S | 0 | 0 |
| | | | 481 | 318 | 88 | 73 | 2 | | |

- Molecule 27 is a protein called 60S ribosomal protein L26, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 27 | la | 210 | Total | C | N | O | S | 0 | 0 |
| | | | 1651 | 1055 | 304 | 285 | 7 | | |

- Molecule 28 is a protein called 60S ribosomal protein L27, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 28 | lb | 137 | Total | C | N | O | S | 0 | 0 |
| | | | 1094 | 707 | 196 | 187 | 4 | | |

- Molecule 29 is a protein called Large ribosomal subunit protein uL15A.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 29 | lc | 148 | Total | C | N | O | S | 0 | 0 |
| | | | 1192 | 757 | 236 | 194 | 5 | | |

- Molecule 30 is a protein called 60S ribosomal protein L29.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 30 | ld | 60 | Total | C | N | O | S | 0 | 0 |
| | | | 478 | 297 | 97 | 82 | 2 | | |

- Molecule 31 is a protein called 60S ribosomal protein L30, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 31 | le | 93 | Total | C | N | O | S | 0 | 0 |
| | | | 693 | 438 | 118 | 135 | 2 | | |

- Molecule 32 is a protein called 60S ribosomal protein L31, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 32 | lf | 126 | Total | C | N | O | S | 0 | 0 |
| | | | 1032 | 662 | 191 | 174 | 5 | | |

- Molecule 33 is a protein called 60S ribosomal protein L32, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 33 | lg | 123 | Total | C | N | O | S | 0 | 0 |
| | | | 1010 | 643 | 200 | 162 | 5 | | |

- Molecule 34 is a protein called 60S ribosomal protein L34, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 34 | lh | 102 | Total | C | N | O | S | 0 | 0 |
| | | | 805 | 503 | 166 | 130 | 6 | | |

- Molecule 35 is a protein called uL29.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 35 | li | 122 | Total | C | N | O | S | 0 | 0 |
| | | | 974 | 620 | 188 | 162 | 4 | | |

- Molecule 36 is a protein called 60S ribosomal protein L35a, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 36 | lj | 106 | Total | C | N | O | S | 0 | 0 |
| | | | 841 | 545 | 158 | 135 | 3 | | |

- Molecule 37 is a protein called 60S ribosomal protein L36, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 37 | lk | 89 | Total | C | N | O | S | 0 | 0 |
| | | | 712 | 447 | 144 | 116 | 5 | | |

- Molecule 38 is a protein called 60S ribosomal protein L37-A, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|----|---|---------|-------|
| 38 | ll | 72 | Total | C | N | O | S | 0 | 0 |
| | | | 591 | 361 | 132 | 91 | 7 | | |

- Molecule 39 is a protein called 60S ribosomal protein L37a, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 39 | lm | 90 | Total | C | N | O | S | 0 | 0 |
| | | | 693 | 432 | 136 | 119 | 6 | | |

- Molecule 40 is a protein called 60S ribosomal protein L38 putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 40 | ln | 73 | Total | C | N | O | S | 0 | 0 |
| | | | 584 | 378 | 104 | 100 | 2 | | |

- Molecule 41 is a protein called Ribosomal protein L39, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 41 | lo | 50 | Total | C | N | O | S | 0 | 0 |
| | | | 432 | 275 | 91 | 63 | 3 | | |

- Molecule 42 is a protein called 60S ribosomal protein L40, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 42 | lp | 53 | Total | C | N | O | S | 0 | 0 |
| | | | 420 | 259 | 86 | 69 | 6 | | |

- Molecule 43 is a protein called 60S ribosomal protein L44, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 43 | lq | 92 | Total | C | N | O | S | 0 | 0 |
| | | | 756 | 480 | 148 | 122 | 6 | | |

- Molecule 44 is a protein called 40S ribosomal protein S25.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|----|----|----|--|---------|-------|
| 44 | sA | 10 | Total | C | N | O | | 0 | 0 |
| | | | 91 | 62 | 15 | 14 | | | |

- Molecule 45 is a protein called 40S ribosomal protein S26.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 45 | sB | 98 | Total | C | N | O | S | 0 | 0 |
| | | | 787 | 478 | 169 | 134 | 6 | | |

- Molecule 46 is a protein called Small ribosomal subunit protein eS27.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 46 | sC | 32 | Total | C | N | O | S | 0 | 0 |
| | | | 249 | 161 | 42 | 45 | 1 | | |

- Molecule 47 is a protein called 40S ribosomal protein S28, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 47 | sD | 60 | Total | C | N | O | S | 0 | 0 |
| | | | 468 | 289 | 93 | 84 | 2 | | |

- Molecule 48 is a protein called Ribosomal protein S29, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 48 | sE | 55 | Total | C | N | O | S | 0 | 0 |
| | | | 442 | 273 | 90 | 75 | 4 | | |

- Molecule 49 is a RNA chain called A/P-tRNA.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|----|---------|-------|
| 49 | sI | 73 | Total | C | N | O | P | 0 | 0 |
| | | | 1556 | 695 | 283 | 506 | 72 | | |

- Molecule 50 is a RNA chain called P/E-tRNA.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|----|---------|-------|
| 50 | sJ | 73 | Total | C | N | O | P | 0 | 0 |
| | | | 1549 | 691 | 271 | 514 | 73 | | |

- Molecule 51 is a RNA chain called mRNA.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|----|----|----|----|---------|-------|
| 51 | sK | 10 | Total | C | N | O | P | 0 | 0 |
| | | | 215 | 97 | 41 | 67 | 10 | | |

- Molecule 52 is a RNA chain called 17S rRNA.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-------|------|------|------|---------|-------|
| 52 | sa | 1401 | Total | C | N | O | P | 0 | 0 |
| | | | 29968 | 13412 | 5443 | 9712 | 1401 | | |

- Molecule 53 is a protein called Small ribosomal subunit protein uS5.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 53 | sc | 196 | Total | C | N | O | S | 0 | 0 |
| | | | 1499 | 960 | 267 | 265 | 7 | | |

- Molecule 54 is a protein called 40S ribosomal protein S3.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|----|---------|-------|
| 54 | sd | 188 | Total | C | N | O | S | 0 | 0 |
| | | | 1440 | 912 | 257 | 260 | 11 | | |

- Molecule 55 is a protein called Small ribosomal subunit protein eS1.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 55 | se | 144 | Total | C | N | O | S | 0 | 0 |
| | | | 1166 | 746 | 208 | 205 | 7 | | |

- Molecule 56 is a protein called 40S ribosomal protein S4, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|----|---------|-------|
| 56 | sf | 256 | Total | C | N | O | S | 0 | 0 |
| | | | 2031 | 1297 | 378 | 345 | 11 | | |

- Molecule 57 is a protein called Small ribosomal subunit protein uS7.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|----|---------|-------|
| 57 | sg | 179 | Total | C | N | O | S | 0 | 0 |
| | | | 1424 | 897 | 258 | 258 | 11 | | |

- Molecule 58 is a protein called 40S ribosomal protein S6.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 58 | sh | 52 | Total | C | N | O | S | 0 | 0 |
| | | | 400 | 243 | 83 | 70 | 4 | | |

- Molecule 59 is a protein called 40S ribosomal protein S7.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 59 | si | 57 | Total | C | N | O | S | 0 | 0 |
| | | | 445 | 285 | 82 | 77 | 1 | | |

- Molecule 60 is a protein called 40S ribosomal protein S8.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 60 | sj | 129 | Total | C | N | O | S | 0 | 0 |
| | | | 1009 | 623 | 203 | 179 | 4 | | |

- Molecule 61 is a protein called Small ribosomal subunit protein uS4.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 61 | sk | 82 | Total | C | N | O | S | 0 | 0 |
| | | | 677 | 430 | 132 | 110 | 5 | | |

- Molecule 62 is a protein called 40S ribosomal protein S10, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 62 | sl | 66 | Total | C | N | O | S | 0 | 0 |
| | | | 536 | 352 | 90 | 85 | 9 | | |

- Molecule 63 is a protein called 40S ribosomal protein S11, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 63 | sm | 141 | Total | C | N | O | S | 0 | 0 |
| | | | 1161 | 735 | 224 | 196 | 6 | | |

- Molecule 64 is a protein called 40S ribosomal protein S13, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 64 | so | 76 | Total | C | N | O | S | 0 | 0 |
| | | | 644 | 410 | 123 | 108 | 3 | | |

- Molecule 65 is a protein called Ribosomal protein S14, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 65 | sp | 128 | Total | C | N | O | S | 0 | 0 |
| | | | 964 | 592 | 187 | 179 | 6 | | |

- Molecule 66 is a protein called 40S ribosomal protein S15, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 66 | sq | 105 | Total | C | N | O | S | 0 | 0 |
| | | | 842 | 543 | 150 | 144 | 5 | | |

- Molecule 67 is a protein called 40S ribosomal protein S15a, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 67 | sr | 129 | Total | C | N | O | S | 0 | 0 |
| | | | 1022 | 650 | 186 | 181 | 5 | | |

- Molecule 68 is a protein called 40S ribosomal protein S16, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 68 | ss | 137 | Total | C | N | O | S | 0 | 0 |
| | | | 1076 | 695 | 193 | 184 | 4 | | |

- Molecule 69 is a protein called 40S ribosomal protein S17, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 69 | st | 81 | Total | C | N | O | S | 0 | 0 |
| | | | 676 | 424 | 135 | 116 | 1 | | |

- Molecule 70 is a protein called Small ribosomal subunit protein uS13.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 70 | su | 124 | Total | C | N | O | S | 0 | 0 |
| | | | 1006 | 625 | 207 | 170 | 4 | | |

- Molecule 71 is a protein called Small ribosomal subunit protein eS19.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 71 | sv | 127 | Total | C | N | O | S | 0 | 0 |
| | | | 1015 | 646 | 185 | 178 | 6 | | |

- Molecule 72 is a protein called 40S ribosomal protein S20, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 72 | sw | 46 | Total | C | N | O | S | 0 | 0 |
| | | | 376 | 238 | 70 | 65 | 3 | | |

- Molecule 73 is a protein called 40S ribosomal protein S21.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 73 | sx | 63 | Total | C | N | O | S | 0 | 0 |
| | | | 494 | 313 | 90 | 88 | 3 | | |

- Molecule 74 is a protein called 40S ribosomal protein S23, putative.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 74 | sy | 106 | Total | C | N | O | S | 0 | 0 |
| | | | 836 | 522 | 169 | 142 | 3 | | |

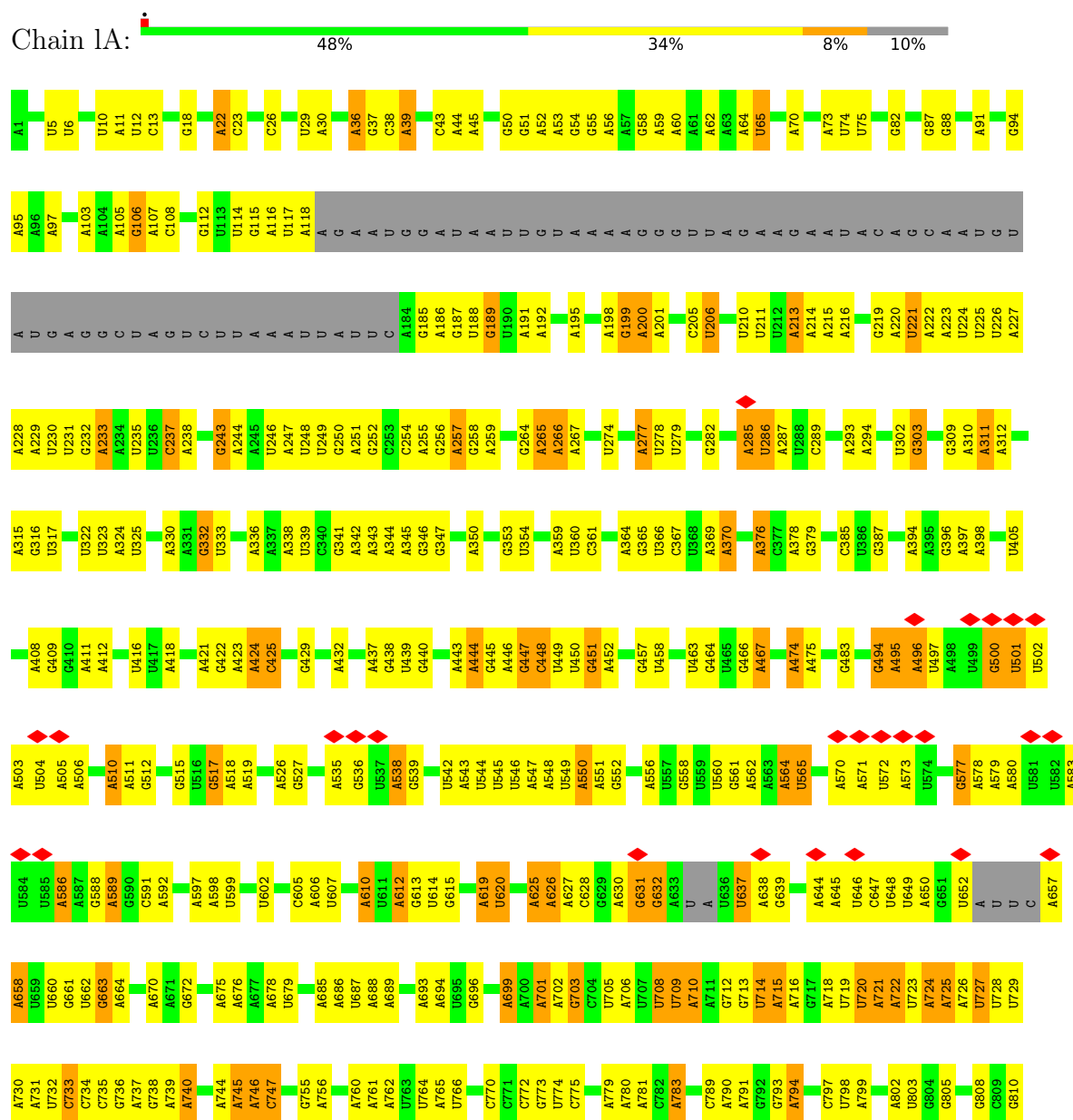
- Molecule 75 is PAROMOMYCIN (CCD ID: PAR) (formula: $C_{23}H_{45}N_5O_{14}$) (labeled as "Ligand of Interest" by depositor).



3 Residue-property plots

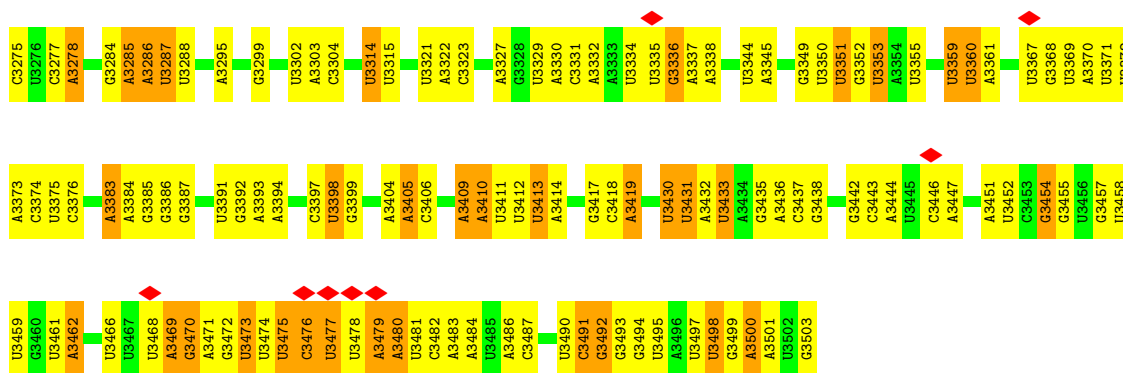
These plots are drawn for all protein, RNA, DNA and oligosaccharide chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and atom inclusion in map 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 diamond above a residue indicates a poor fit to the EM map for this residue (all-atom inclusion < 40%). 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: 25S rRNA



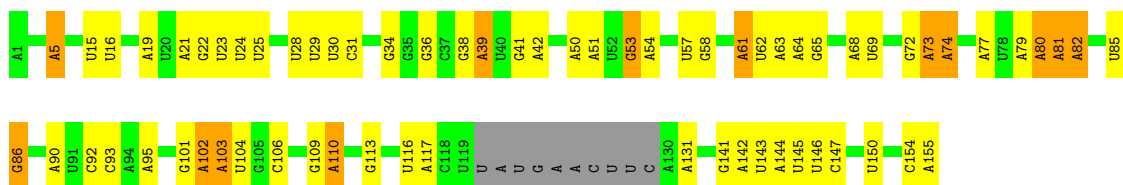


| | | | | | | | | | | | | |
|-------|-------|-------|-------|-------|---|---|-------|-------|-------|-------|-------|---|
| A3162 | U2969 | U3073 | U2876 | G2889 | A | G | G2466 | G2378 | C2273 | U2188 | U2082 | U |
| G3163 | U2970 | U3074 | U2877 | A2896 | U | A | G2467 | G2379 | A2274 | A2189 | A2083 | U |
| G3166 | U2971 | A3076 | U2878 | U2897 | U | C | G2468 | A2379 | U2281 | A2190 | G2084 | A |
| A3170 | U2972 | A3077 | U2879 | U2898 | U | U | G2469 | G2380 | U2282 | A2191 | A2089 | G |
| A3171 | U2973 | U3078 | U2880 | U2899 | C | U | G2470 | G2381 | U2283 | G2192 | G2090 | A |
| A3175 | U2974 | A3079 | U2881 | U2900 | C | U | A2471 | G2382 | U2284 | U2193 | G2091 | A |
| G3176 | U2975 | A3080 | U2882 | U2901 | A | U | A2472 | G2383 | U2285 | U2194 | G2092 | A |
| G3179 | U2976 | A3081 | U2883 | U2902 | U | G | A2473 | G2384 | U2286 | U2195 | G2093 | A |
| G3185 | U2977 | A3082 | U2884 | U2903 | A | U | A2474 | G2385 | U2287 | U2196 | G2094 | A |
| U3196 | U2978 | A3083 | U2885 | U2904 | U | U | A2475 | G2386 | U2288 | U2197 | G2095 | A |
| C3197 | U2979 | A3084 | U2886 | U2905 | U | U | A2476 | G2387 | U2289 | U2198 | G2101 | A |
| G3200 | U2980 | A3085 | U2887 | U2906 | U | U | A2477 | G2388 | U2290 | G2102 | G2107 | A |
| A3203 | U2981 | A3086 | U2888 | U2907 | U | U | A2478 | G2389 | U2291 | G2103 | G2108 | A |
| C3204 | U2982 | A3087 | U2889 | U2908 | U | U | A2479 | G2390 | U2292 | G2104 | G2109 | A |
| U3210 | U2983 | A3088 | U2890 | U2909 | U | U | A2480 | G2391 | U2293 | G2105 | G2110 | A |
| A3211 | U2984 | A3089 | U2891 | U2910 | U | U | A2481 | G2392 | U2294 | G2106 | A2111 | A |
| U3212 | U2985 | A3090 | U2892 | U2911 | U | U | A2482 | G2393 | U2295 | G2107 | A2112 | U |
| U3213 | U2986 | A3091 | U2893 | U2912 | U | U | A2483 | G2394 | U2296 | G2108 | A2113 | U |
| C3214 | U2987 | A3092 | U2894 | U2913 | U | U | A2484 | G2395 | U2297 | G2109 | U2114 | A |
| A3215 | U2988 | A3093 | U2895 | U2914 | U | U | A2485 | G2396 | U2298 | G2110 | A2115 | A |
| A3216 | U2989 | A3094 | U2896 | U2915 | U | U | A2486 | G2397 | U2299 | G2111 | A2116 | A |
| U3220 | U2990 | A3095 | U2897 | U2916 | U | U | A2487 | G2398 | U2300 | G2112 | A2117 | A |
| U3221 | U2991 | A3096 | U2898 | U2917 | U | U | A2488 | G2399 | U2301 | G2113 | A2118 | A |
| U3222 | U2992 | A3097 | U2899 | U2918 | U | U | A2489 | G2400 | U2302 | G2114 | A2119 | A |
| G3223 | U2993 | A3098 | U2900 | U2919 | U | U | A2490 | G2401 | U2303 | G2115 | A2120 | A |
| G3232 | U2994 | A3099 | U2901 | U2920 | U | U | A2491 | G2402 | U2304 | G2116 | A2121 | A |
| G3239 | U2995 | A3100 | U2902 | U2921 | U | U | A2492 | G2403 | U2305 | G2117 | A2122 | A |
| A3240 | U2996 | A3101 | U2903 | U2922 | U | U | A2493 | G2404 | U2306 | G2118 | A2123 | A |
| C3243 | U2997 | A3102 | U2904 | U2923 | U | U | A2494 | G2405 | U2307 | G2119 | A2124 | A |
| A3246 | U2998 | A3103 | U2905 | U2924 | U | U | A2495 | G2406 | U2308 | G2120 | A2125 | A |
| C3247 | U2999 | A3104 | U2906 | U2925 | U | U | A2496 | G2407 | U2309 | G2121 | A2126 | A |
| U3254 | U3000 | A3105 | U2907 | U2926 | U | U | A2497 | G2408 | U2310 | G2122 | A2127 | A |
| U3259 | U3001 | A3106 | U2908 | U2927 | U | U | A2498 | G2409 | U2311 | G2123 | A2128 | A |
| U3262 | U3002 | A3107 | U2909 | U2928 | U | U | A2499 | G2410 | U2312 | G2124 | A2129 | A |
| G3263 | U3003 | A3108 | U2910 | U2929 | U | U | A2500 | G2411 | U2313 | G2125 | A2130 | A |
| G3264 | U3004 | A3109 | U2911 | U2930 | U | U | A2501 | G2412 | U2314 | G2126 | A2131 | A |
| C3265 | U3005 | A3110 | U2912 | U2931 | U | U | A2502 | G2413 | U2315 | G2127 | A2132 | A |
| U3266 | U3006 | A3111 | U2913 | U2932 | U | U | A2503 | G2414 | U2316 | G2128 | A2133 | A |
| G3267 | U3007 | A3112 | U2914 | U2933 | U | U | A2504 | G2415 | U2317 | G2129 | A2134 | A |
| A3270 | U3008 | A3113 | U2915 | U2934 | U | U | A2505 | G2416 | U2318 | G2130 | A2135 | A |
| U3274 | U3009 | A3114 | U2916 | U2935 | U | U | A2506 | G2417 | U2319 | G2131 | A2136 | A |
| | U3010 | A3115 | U2917 | U2936 | U | U | A2507 | G2418 | U2320 | G2132 | A2137 | A |
| | U3011 | A3116 | U2918 | U2937 | U | U | A2508 | G2419 | U2321 | G2133 | A2138 | A |
| | U3012 | A3117 | U2919 | U2938 | U | U | A2509 | G2420 | U2322 | G2134 | A2139 | A |
| | U3013 | A3118 | U2920 | U2939 | U | U | A2510 | G2421 | U2323 | G2135 | A2140 | A |
| | U3014 | A3119 | U2921 | U2940 | U | U | A2511 | G2422 | U2324 | G2136 | A2141 | A |
| | U3015 | A3120 | U2922 | U2941 | U | U | A2512 | G2423 | U2325 | G2137 | A2142 | A |
| | U3016 | A3121 | U2923 | U2942 | U | U | A2513 | G2424 | U2326 | G2138 | A2143 | A |
| | U3017 | A3122 | U2924 | U2943 | U | U | A2514 | G2425 | U2327 | G2139 | A2144 | A |
| | U3018 | A3123 | U2925 | U2944 | U | U | A2515 | G2426 | U2328 | G2140 | A2145 | A |
| | U3019 | A3124 | U2926 | U2945 | U | U | A2516 | G2427 | U2329 | G2141 | A2146 | A |
| | U3020 | A3125 | U2927 | U2946 | U | U | A2517 | G2428 | U2330 | G2142 | A2147 | A |
| | U3021 | A3126 | U2928 | U2947 | U | U | A2518 | G2429 | U2331 | G2143 | A2148 | A |
| | U3022 | A3127 | U2929 | U2948 | U | U | A2519 | G2430 | U2332 | G2144 | A2149 | A |
| | U3023 | A3128 | U2930 | U2949 | U | U | A2520 | G2431 | U2333 | G2145 | A2150 | A |
| | U3024 | A3129 | U2931 | U2950 | U | U | A2521 | G2432 | U2334 | G2146 | A2151 | A |
| | U3025 | A3130 | U2932 | U2951 | U | U | A2522 | G2433 | U2335 | G2147 | A2152 | A |
| | U3026 | A3131 | U2933 | U2952 | U | U | A2523 | G2434 | U2336 | G2148 | A2153 | A |
| | U3027 | A3132 | U2934 | U2953 | U | U | A2524 | G2435 | U2337 | G2149 | A2154 | A |
| | U3028 | A3133 | U2935 | U2954 | U | U | A2525 | G2436 | U2338 | G2150 | A2155 | A |
| | U3029 | A3134 | U2936 | U2955 | U | U | A2526 | G2437 | U2339 | G2151 | A2156 | A |
| | U3030 | A3135 | U2937 | U2956 | U | U | A2527 | G2438 | U2340 | G2152 | A2157 | A |
| | U3031 | A3136 | U2938 | U2957 | U | U | A2528 | G2439 | U2341 | G2153 | A2158 | A |
| | U3032 | A3137 | U2939 | U2958 | U | U | A2529 | G2440 | U2342 | G2154 | A2159 | A |
| | U3033 | A3138 | U2940 | U2959 | U | U | A2530 | G2441 | U2343 | G2155 | A2160 | A |
| | U3034 | A3139 | U2941 | U2960 | U | U | A2531 | G2442 | U2344 | G2156 | A2161 | A |
| | U3035 | A3140 | U2942 | U2961 | U | U | A2532 | G2443 | U2345 | G2157 | A2162 | A |
| | U3036 | A3141 | U2943 | U2962 | U | U | A2533 | G2444 | U2346 | G2158 | A2163 | A |
| | U3037 | A3142 | U2944 | U2963 | U | U | A2534 | G2445 | U2347 | G2159 | A2164 | A |
| | U3038 | A3143 | U2945 | U2964 | U | U | A2535 | G2446 | U2348 | G2160 | A2165 | A |
| | U3039 | A3144 | U2946 | U2965 | U | U | A2536 | G2447 | U2349 | G2161 | A2166 | A |
| | U3040 | A3145 | U2947 | U2966 | U | U | A2537 | G2448 | U2350 | G2162 | A2167 | A |
| | U3041 | A3146 | U2948 | U2967 | U | U | A2538 | G2449 | U2351 | G2163 | A2168 | A |
| | U3042 | A3147 | U2949 | U2968 | U | U | A2539 | G2450 | U2352 | G2164 | A2169 | A |
| | U3043 | A3148 | U2950 | U2969 | U | U | A2540 | G2451 | U2353 | G2165 | A2170 | A |
| | U3044 | A3149 | U2951 | U2970 | U | U | A2541 | G2452 | U2354 | G2166 | A2171 | A |
| | U3045 | A3150 | U2952 | U2971 | U | U | A2542 | G2453 | U2355 | G2167 | A2172 | A |
| | U3046 | A3151 | U2953 | U2972 | U | U | A2543 | G2454 | U2356 | G2168 | A2173 | A |
| | U3047 | A3152 | U2954 | U2973 | U | U | A2544 | G2455 | U2357 | G2169 | A2174 | A |
| | U3048 | A3153 | U2955 | U2974 | U | U | A2545 | G2456 | U2358 | G2170 | A2175 | A |
| | U3049 | A3154 | U2956 | U2975 | U | U | A2546 | G2457 | U2359 | G2171 | A2176 | A |
| | U3050 | A3155 | U2957 | U2976 | U | U | A2547 | G2458 | U2360 | G2172 | A2177 | A |
| | U3051 | A3156 | U2958 | U2977 | U | U | A2548 | G2459 | U2361 | G2173 | A2178 | A |
| | U3052 | A3157 | U2959 | U2978 | U | U | A2549 | G2460 | U2362 | G2174 | A2179 | A |
| | U3053 | A3158 | U2960 | U2979 | U | U | A2550 | G2461 | U2363 | G2175 | A2180 | A |
| | U3054 | A3159 | U2961 | U2980 | U | U | A2551 | G2462 | U2364 | G2176 | A2181 | A |
| | U3055 | A3160 | U2962 | U2981 | U | U | A2552 | G2463 | U2365 | G2177 | A2182 | A |
| | U3056 | A3161 | U2963 | U2982 | U | U | A2553 | G2464 | U2366 | G2178 | A2183 | A |
| | U3057 | A3162 | U2964 | U2983 | U | U | A2554 | G2465 | U2367 | G2179 | A2184 | A |
| | U3058 | A3163 | U2965 | U2984 | U | U | A2555 | G2466 | U2368 | G2180 | A2185 | A |
| | U3059 | A3164 | U2966 | U2985 | U | U | A2556 | G2467 | U2369 | G2181 | A2186 | A |
| | U3060 | A3165 | U2967 | U2986 | U | U | A2557 | G2468 | U2370 | G2182 | A2187 | A |
| | U3061 | A3166 | U2968 | U2987 | U | U | A2558 | G2469 | U2371 | G2183 | A2188 | A |
| | U3062 | A3167 | U2969 | U2988 | U | U | A2559 | G2470 | U2372 | G2184 | A2189 | A |
| | U3063 | A3168 | U2970 | U2989 | U | U | A2560 | G2471 | U2373 | G2185 | A2190 | A |
| | U3064 | A3169 | U2971 | U2990 | U | U | A2561 | G2472 | U2374 | G2186 | A2191 | A |
| | U3065 | A3170 | U2972 | U2991 | U | U | A2562 | G2473 | U2375 | G2187 | A2192 | A |
| | U3066 | A3171 | U2973 | U2992 | U | U | A2563 | G2474 | U2376 | G2188 | A2193 | A |
| | U3067 | A3172 | U2974 | U2993 | U | U | A2564 | G2475 | U2377 | G2189 | A2194 | A |
| | U3068 | A3173 | U2975 | U2994 | U | U | A2565 | G2476 | U2378 | G2190 | A2195 | A |
| | U3069 | A3174 | U2976 | U2995 | U | U | A2566 | G2477 | U2379 | G2191 | A2196 | A |
| | U3070 | A3175 | U2977 | U2996 | U | U | A2567 | G2478 | U2380 | G2192 | A2197 | A |
| | U3071 | A3176 | U2978 | U2997 | U | U | A2568 | G2479 | U2381 | G2193 | A2198 | A |
| | U3072 | A3177 | U2979 | U2998 | U | U | A2569 | G2480 | U2382 | G2194 | A2199 | A |
| | U3073 | A3178 | U2980 | U2999 | U | U | A2570 | G2481 | U2383 | G2195 | A2200 | A |
| | U3074 | A3179 | U2981 | U3000 | U | U | A2571 | G2482 | U2384 | G2196 | A2201 | A |
| | U3075 | A3180 | U2982 | U3001 | U | U | A2572 | G2483 | U2385 | G2197 | A2202 | A |
| | U3076 | A3181 | U2983 | U3002 | U | U | A2573 | G2484 | U2386 | G2198 | A2203 | A |
| | U3077 | A3182 | U2984 | U3003 | U | U | A2574 | G2485 | U2387 | G2199 | A2204 | A |
| | U3078 | A3183 | U2985 | | | | | | | | | |



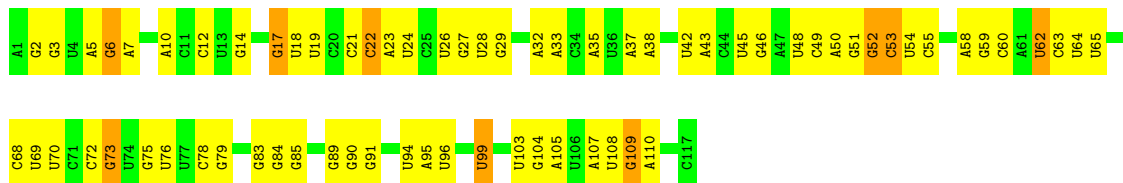
• Molecule 2: 5.8S rRNA

Chain IB: 50% 35% 8% 6%



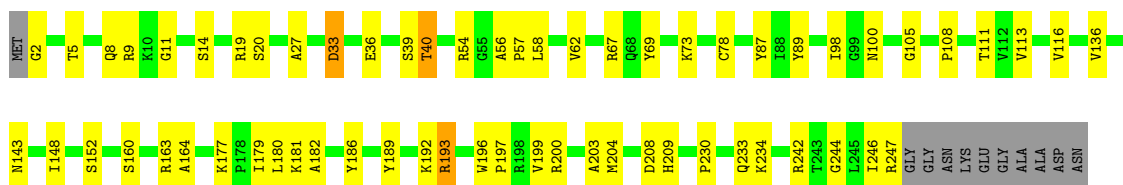
• Molecule 3: 5S rRNA

Chain IC: 41% 51% 8%



• Molecule 4: Large ribosomal subunit protein uL2

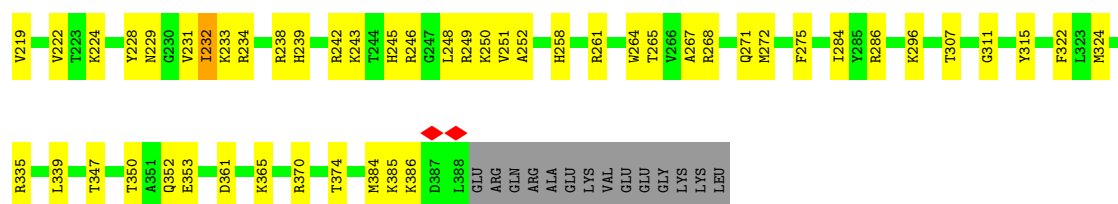
Chain ID: 72% 23%



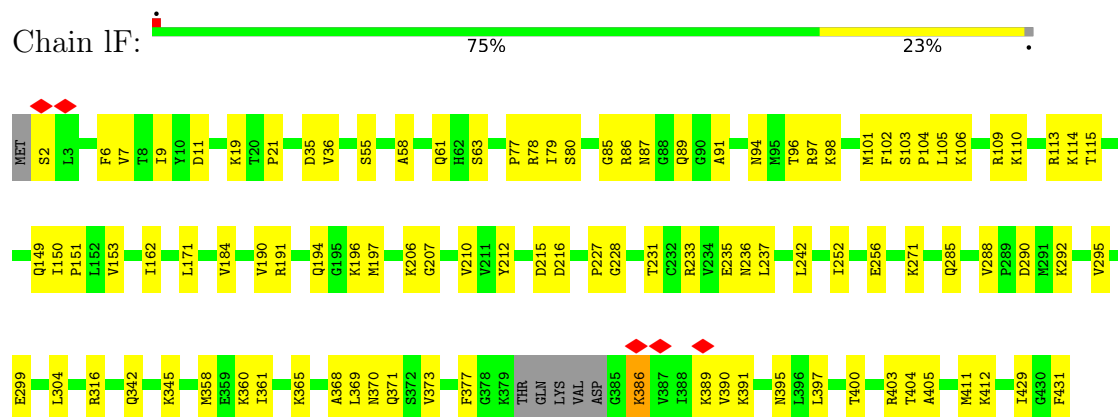
• Molecule 5: 60S ribosomal protein L3, putative

Chain IE: 76% 19%

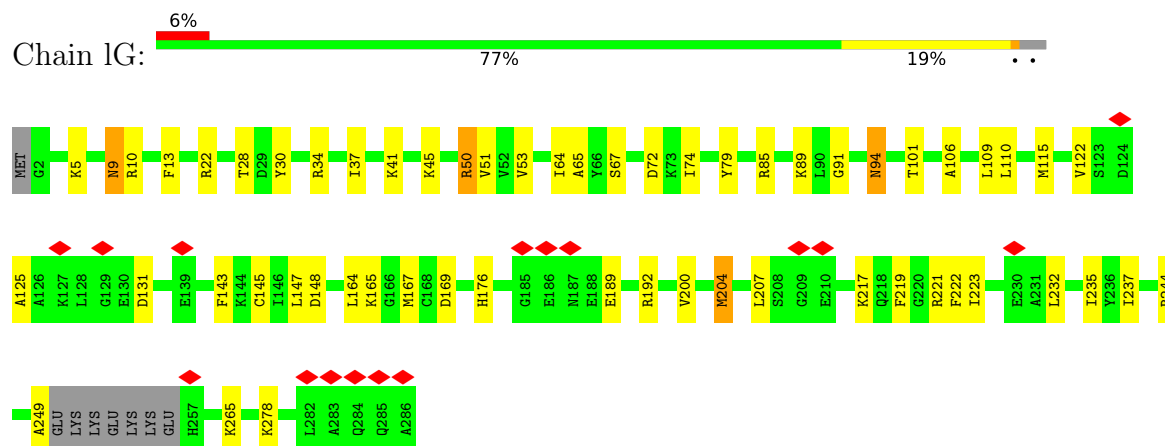




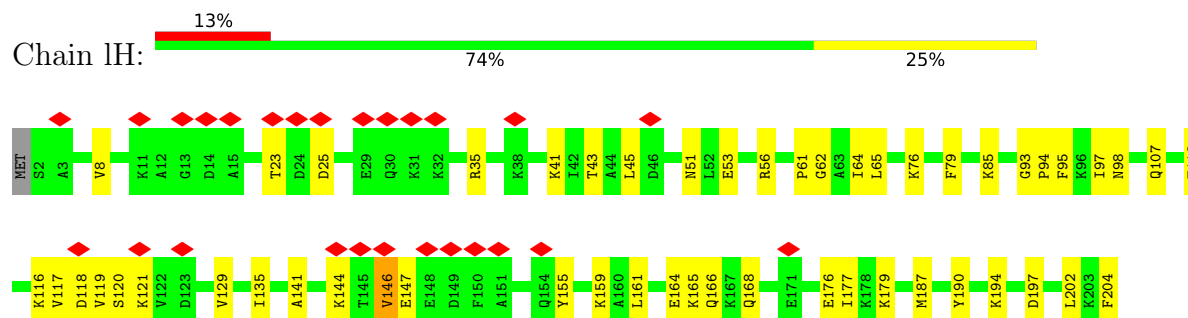
- Molecule 6: 60S ribosomal protein L4, putative



- Molecule 7: 60S ribosomal protein L5, putative

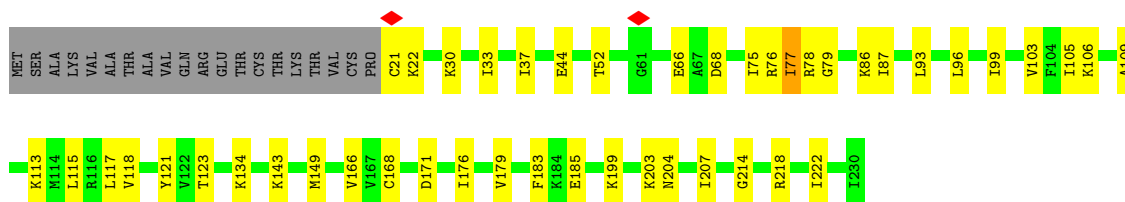


- Molecule 8: Large ribosomal subunit protein eL6

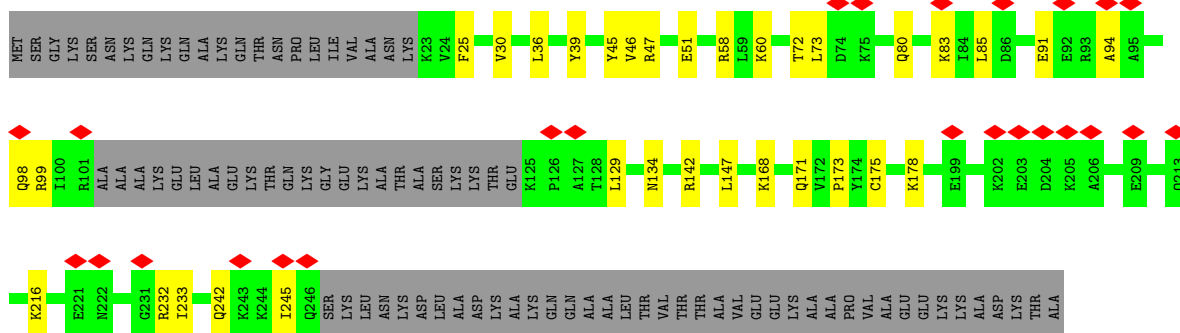


- Molecule 9: 60S ribosomal protein L7, putative

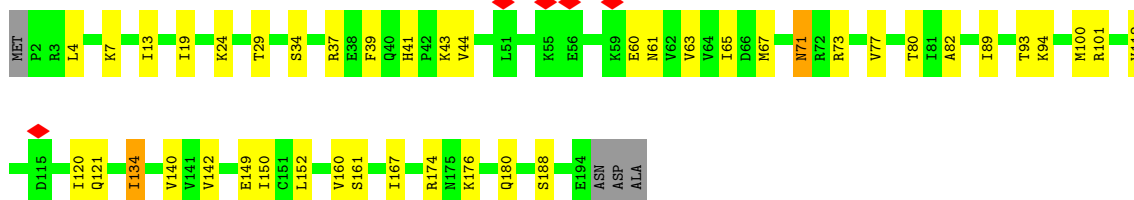
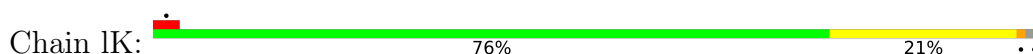




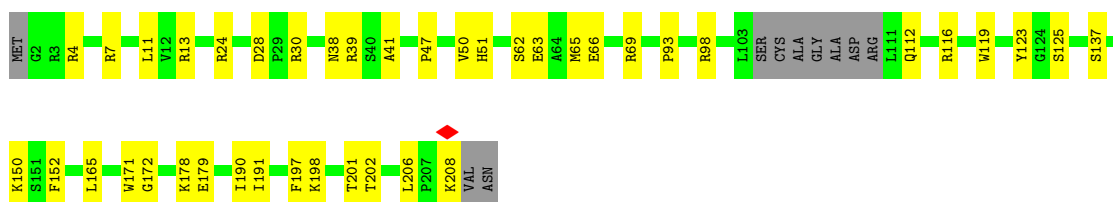
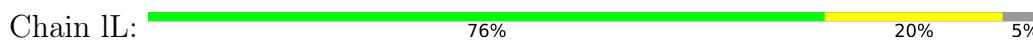
• Molecule 10: 60S ribosomal protein L7a



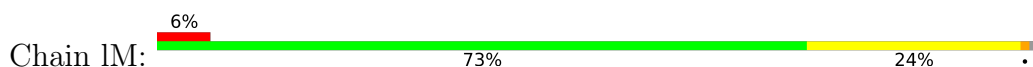
• Molecule 11: 60S ribosomal protein L9, putative



• Molecule 12: Ribosomal protein L10, putative

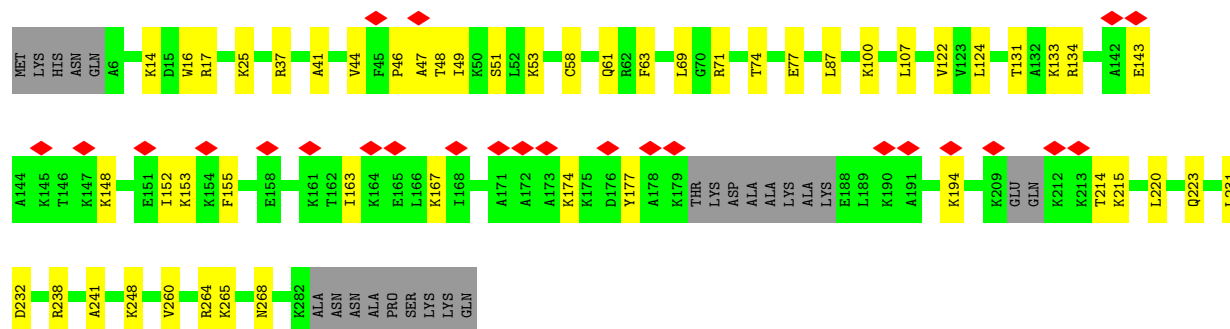
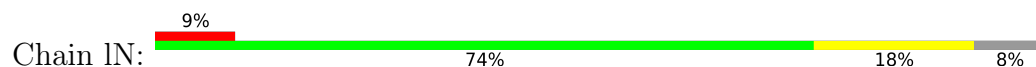


• Molecule 13: 60S ribosomal protein L11, putative

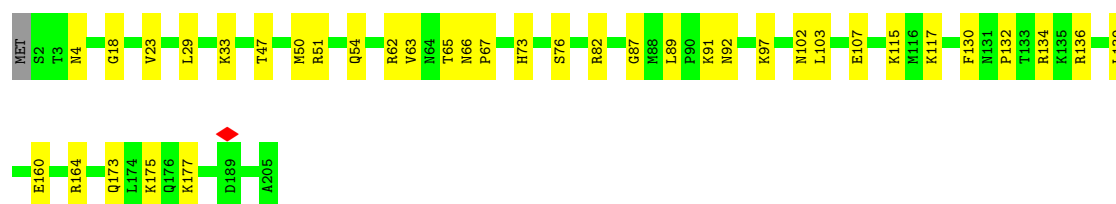
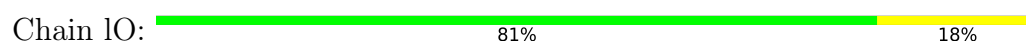




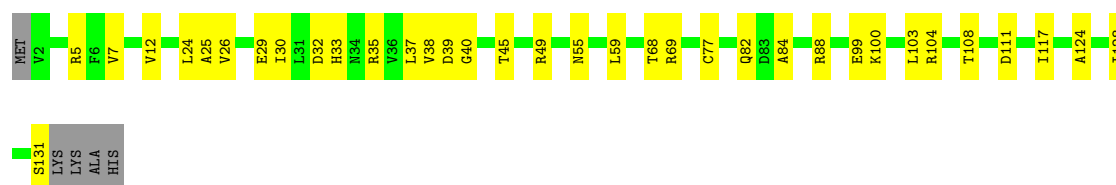
- Molecule 14: 60S ribosomal protein L13, putative



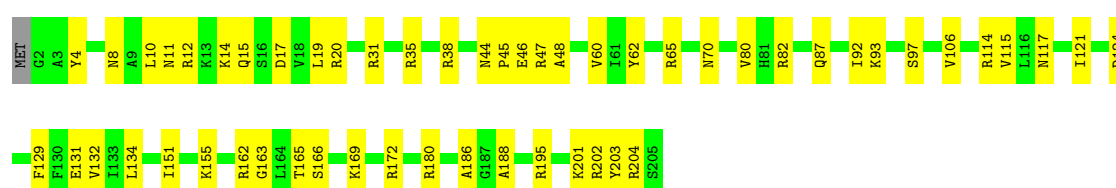
- Molecule 15: 60S ribosomal protein L13, putative



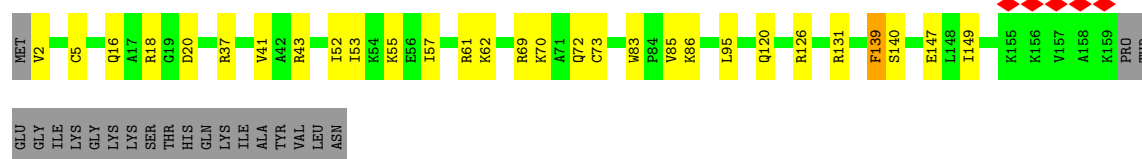
- Molecule 16: 60S ribosomal protein L14, putative



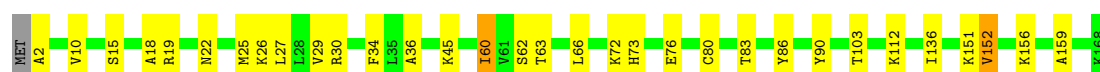
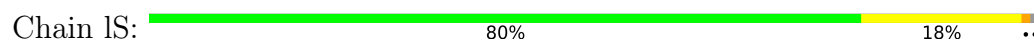
- Molecule 17: Ribosomal protein L15



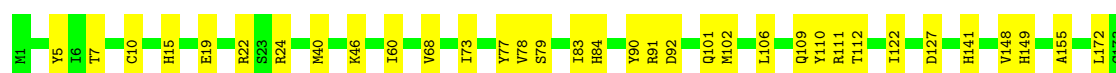
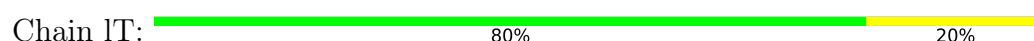
- Molecule 18: 60S ribosomal protein L17, putative



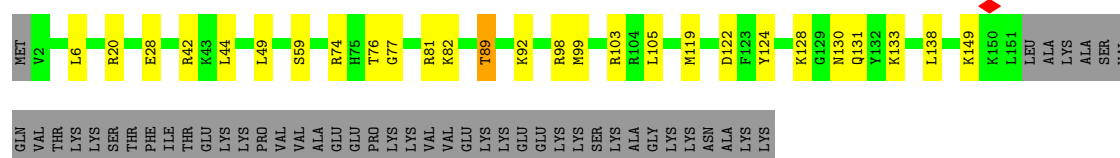
- Molecule 19: 60S ribosomal protein L18, putative



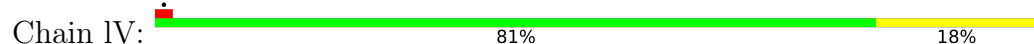
- Molecule 20: 60S ribosomal protein L18a



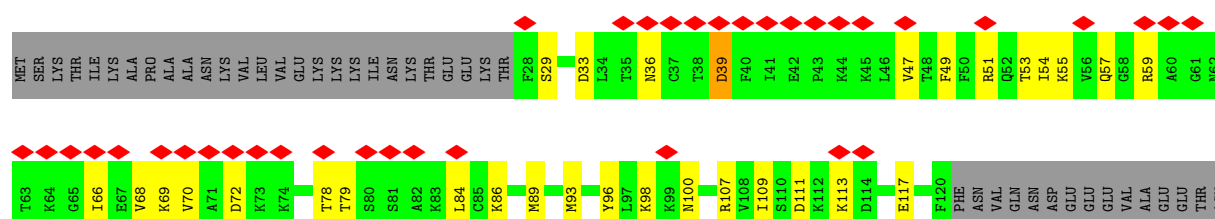
- Molecule 21: Ribosomal protein L19

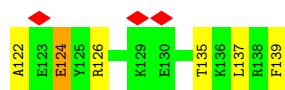


- Molecule 22: 60S ribosomal protein L21, putative

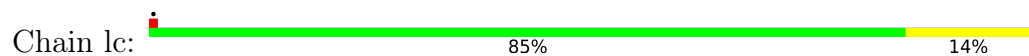


- Molecule 23: Large ribosomal subunit protein eL22

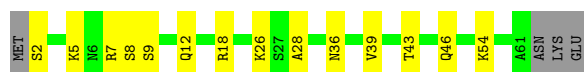




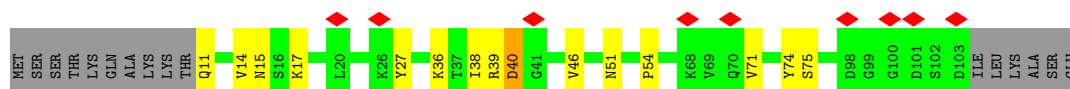
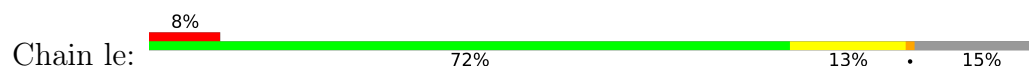
- Molecule 29: Large ribosomal subunit protein uL15A



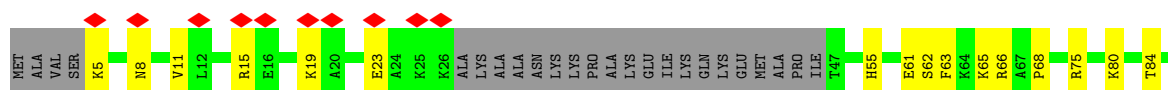
- Molecule 30: 60S ribosomal protein L29



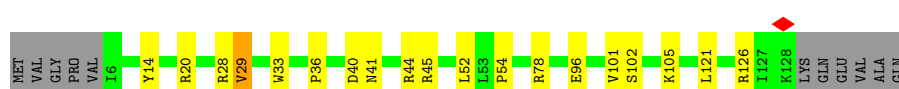
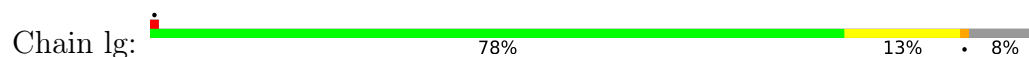
- Molecule 31: 60S ribosomal protein L30, putative



- Molecule 32: 60S ribosomal protein L31, putative

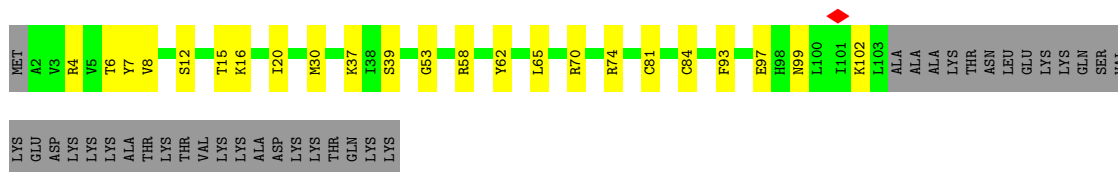


- Molecule 33: 60S ribosomal protein L32, putative

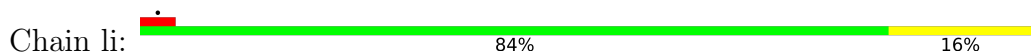


- Molecule 34: 60S ribosomal protein L34, putative

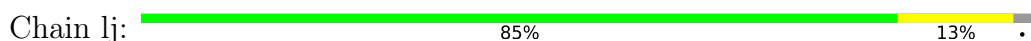




• Molecule 35: uL29



• Molecule 36: 60S ribosomal protein L35a, putative



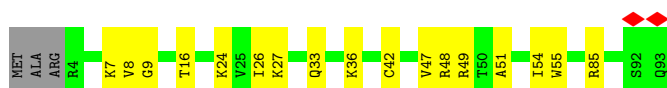
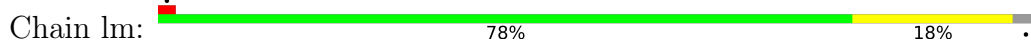
• Molecule 37: 60S ribosomal protein L36, putative



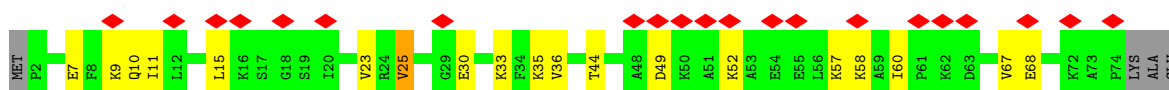
• Molecule 38: 60S ribosomal protein L37-A, putative



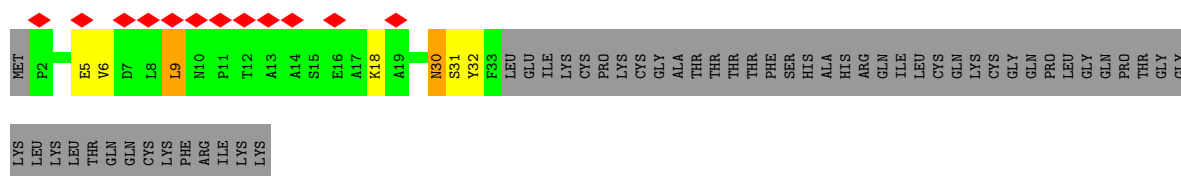
• Molecule 39: 60S ribosomal protein L37a, putative



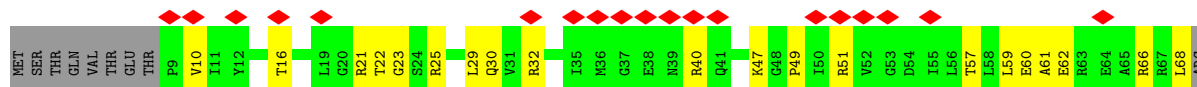
• Molecule 40: 60S ribosomal protein L38 putative



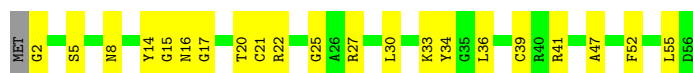
- Chain sC: 



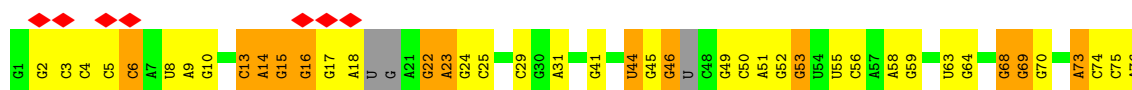
- Molecule 47: 40S ribosomal protein S28, putative



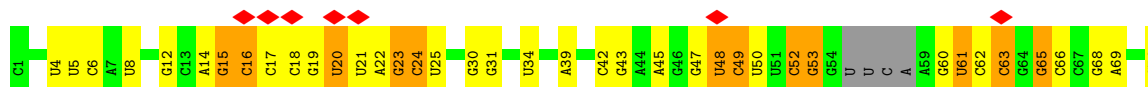
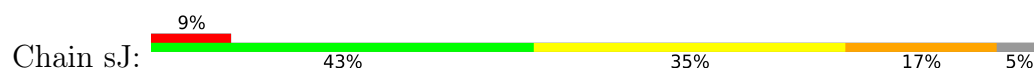
- Molecule 48: Ribosomal protein S29, putative



- Molecule 49: A/P-tRNA

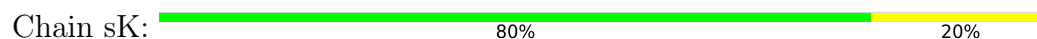


- Molecule 50: P/E-tRNA



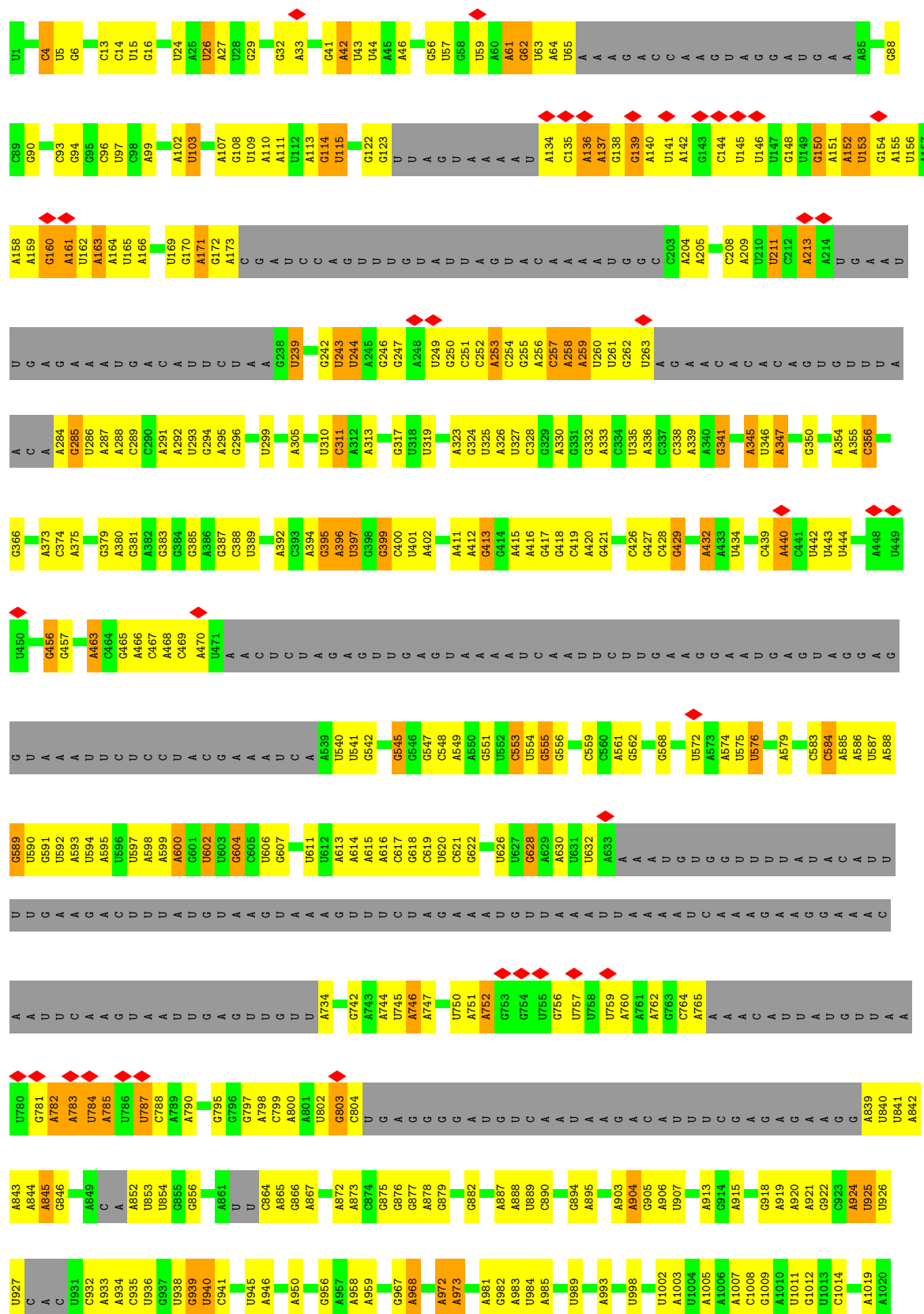
A77

- Molecule 51: mRNA

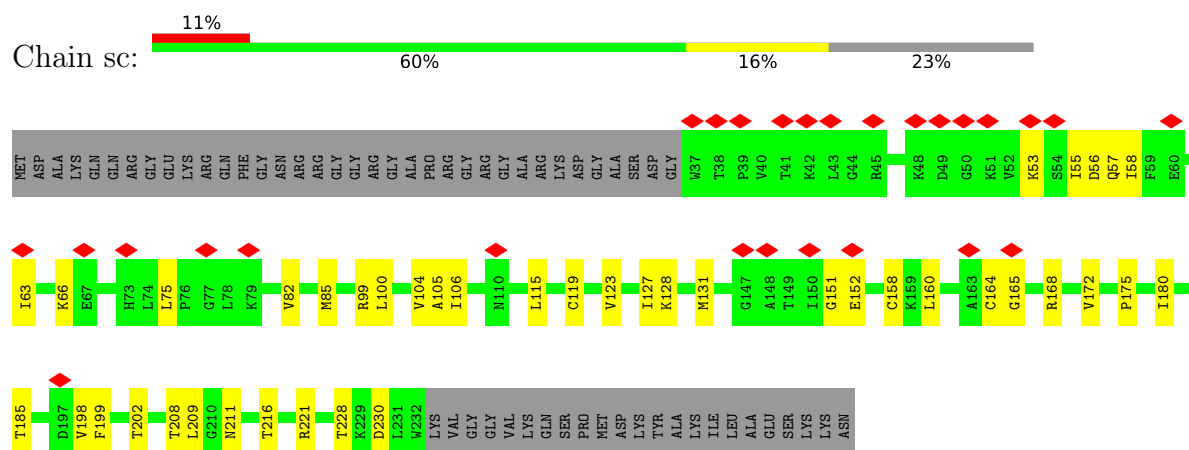


- Molecule 52: 17S rRNA

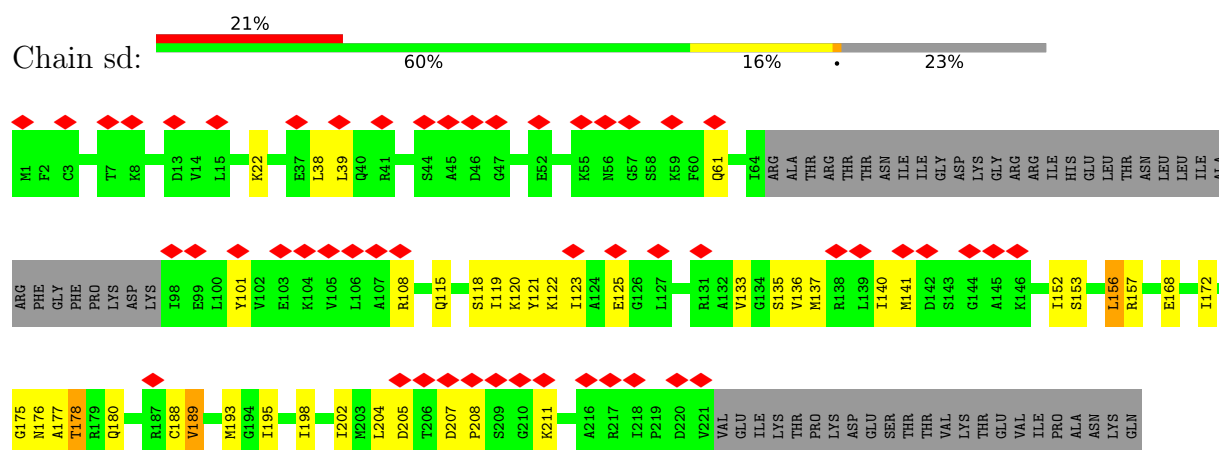




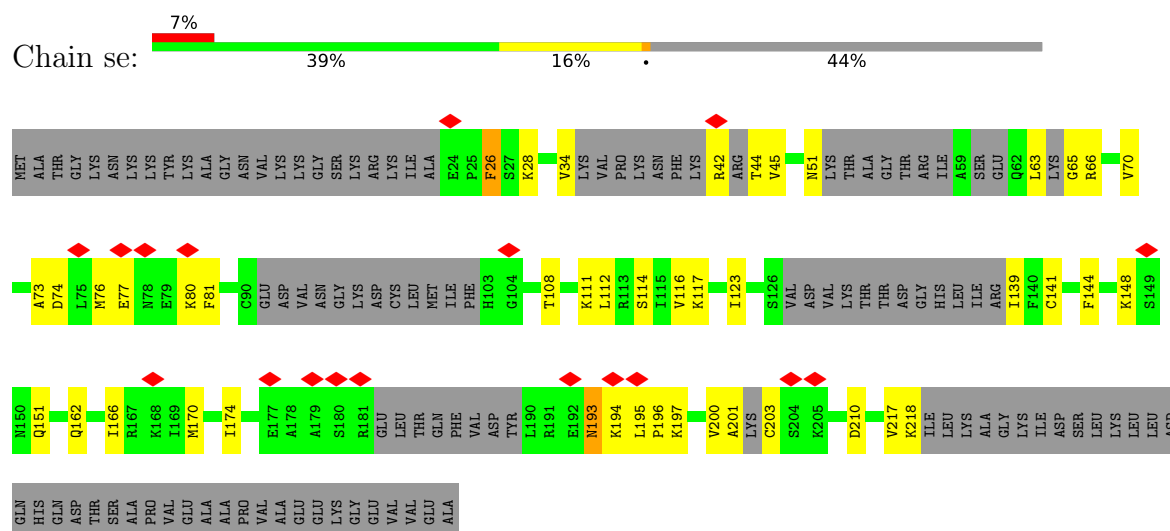




- Molecule 54: 40S ribosomal protein S3

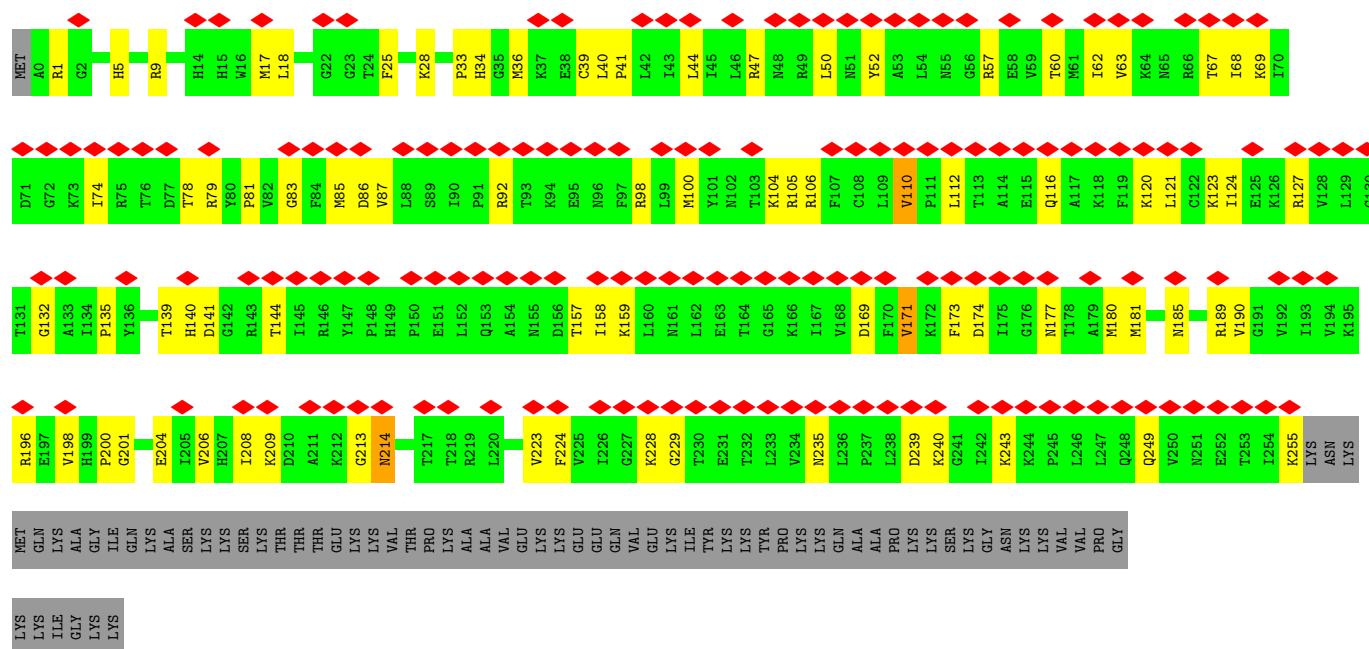


- Molecule 55: Small ribosomal subunit protein eS1

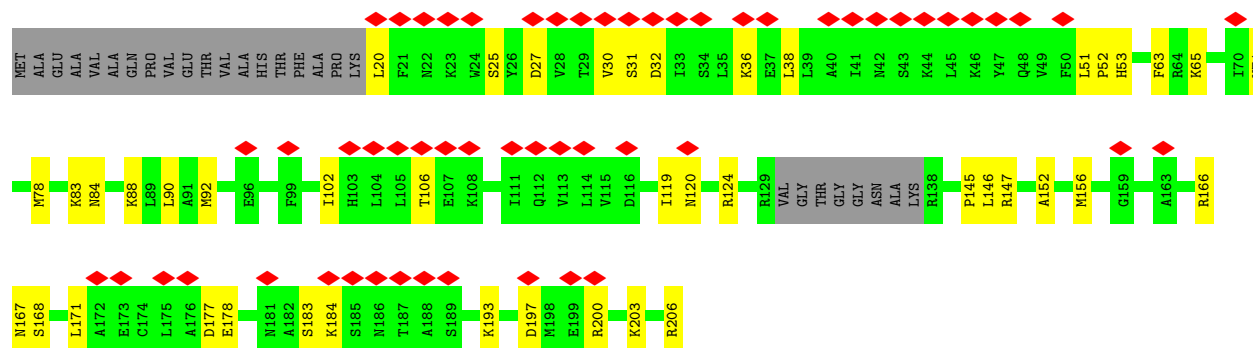


- Molecule 56: 40S ribosomal protein S4, putative

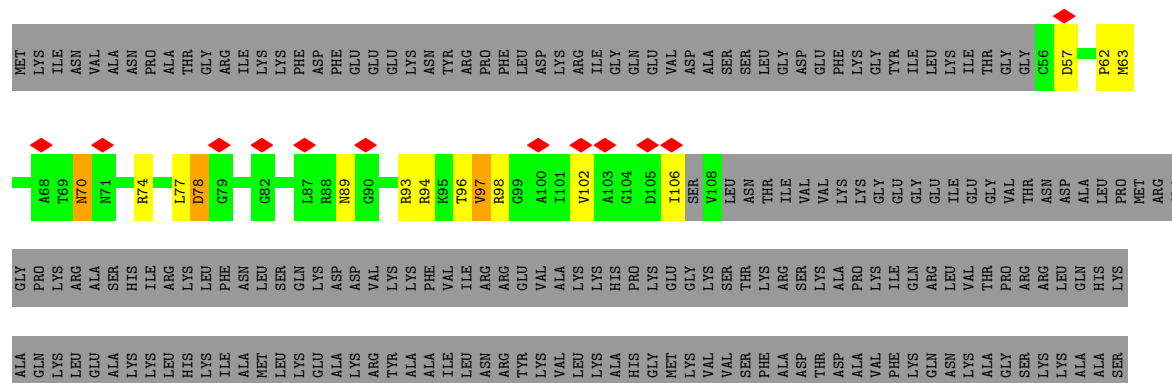




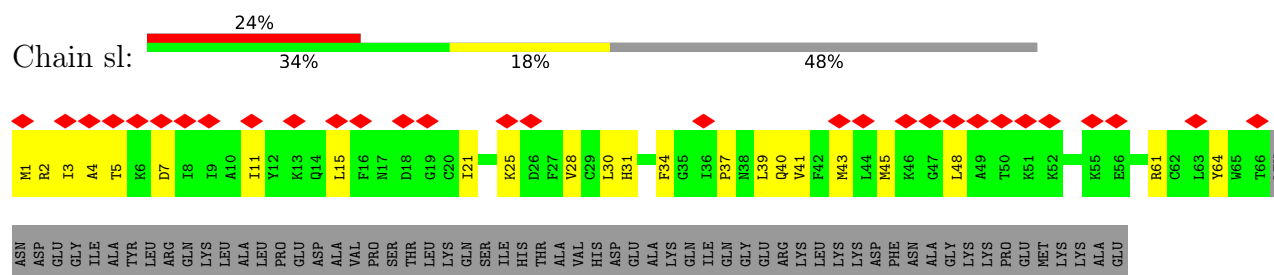
- Molecule 57: Small ribosomal subunit protein uS7



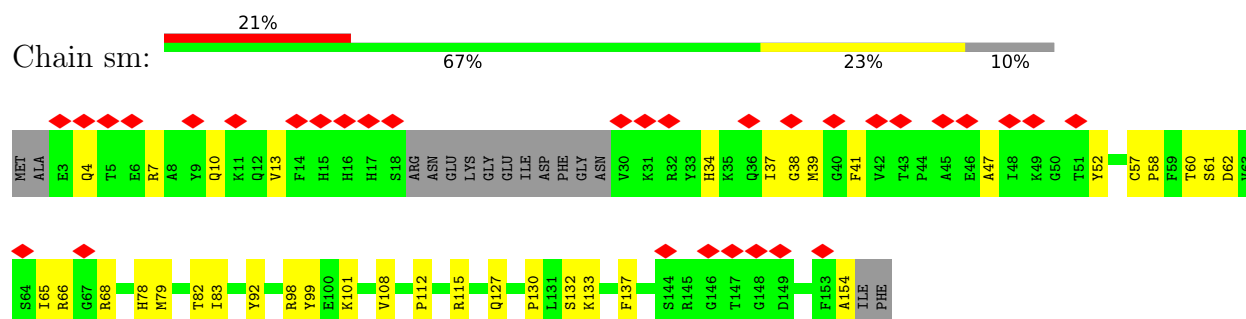
- Molecule 58: 40S ribosomal protein S6



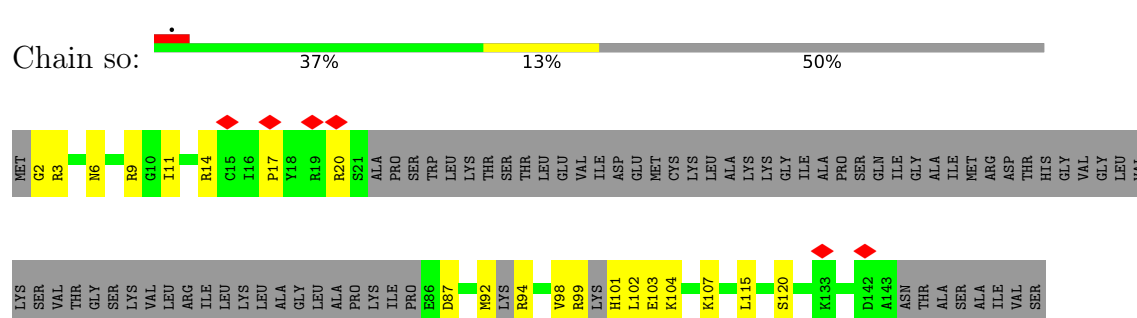
- Molecule 62: 40S ribosomal protein S10, putative



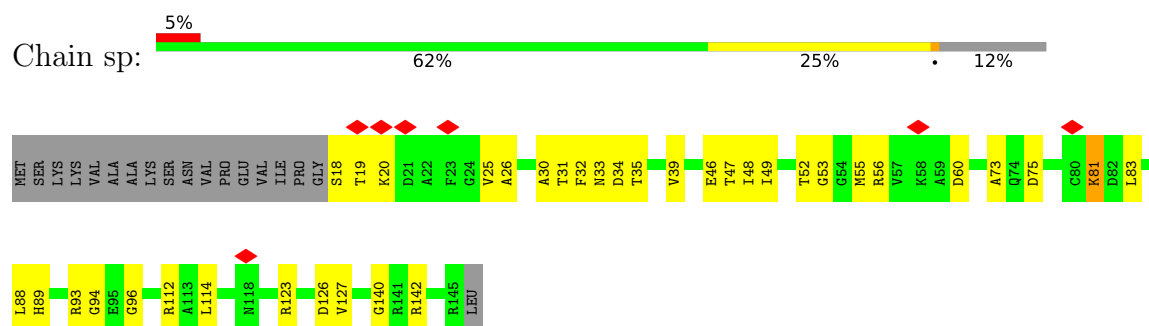
- Molecule 63: 40S ribosomal protein S11, putative



- Molecule 64: 40S ribosomal protein S13, putative

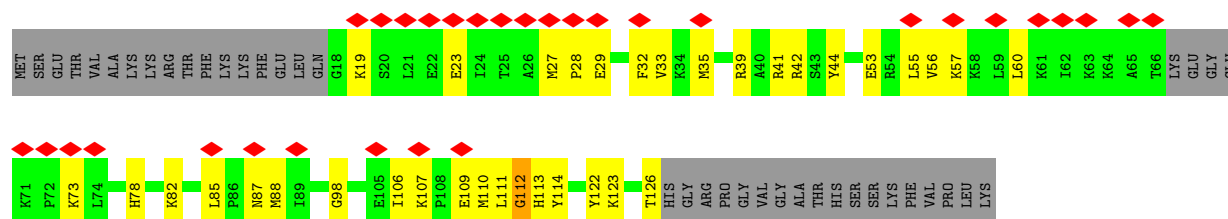


- Molecule 65: Ribosomal protein S14, putative



- Molecule 66: 40S ribosomal protein S15, putative





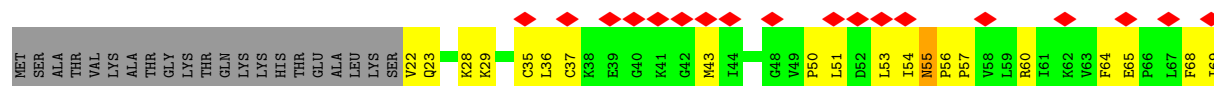
- Molecule 67: 40S ribosomal protein S15a, putative

Chain sr: 74% 25% ..



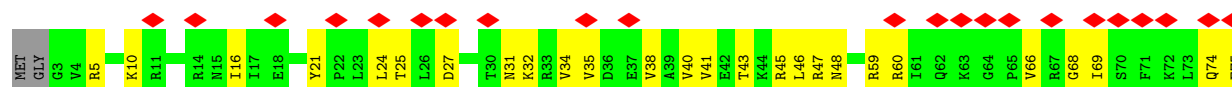
- Molecule 68: 40S ribosomal protein S16, putative

Chain ss: 29% 63% 23% 13%



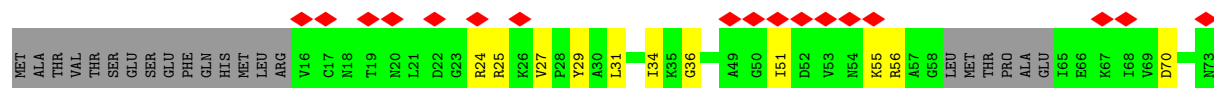
- Molecule 69: 40S ribosomal protein S17, putative

Chain st: 26% 44% 25% 31%



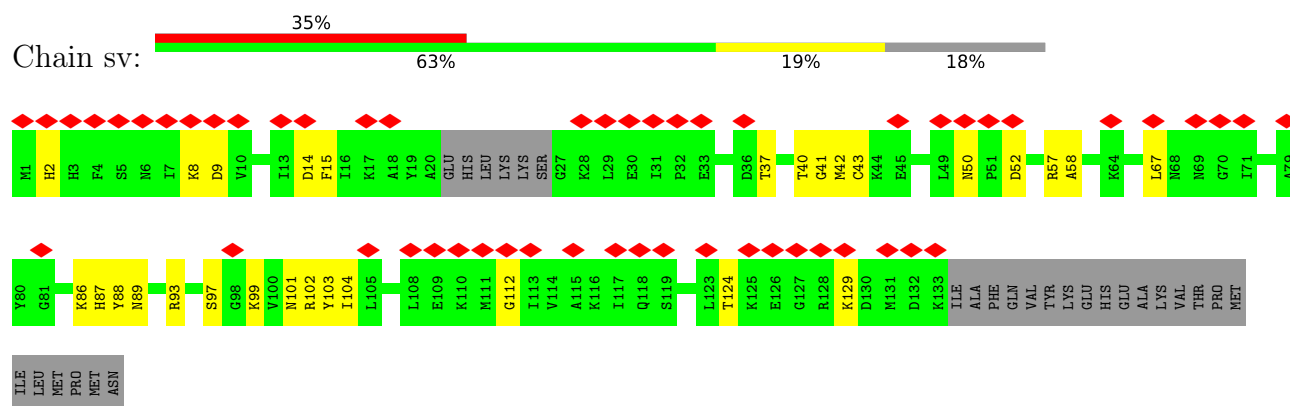
- Molecule 70: Small ribosomal subunit protein uS13

Chain su: 17% 56% 24% 20%

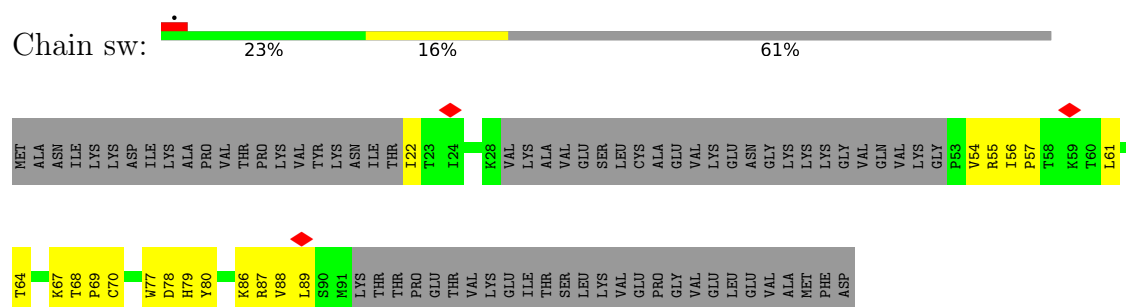


ALA

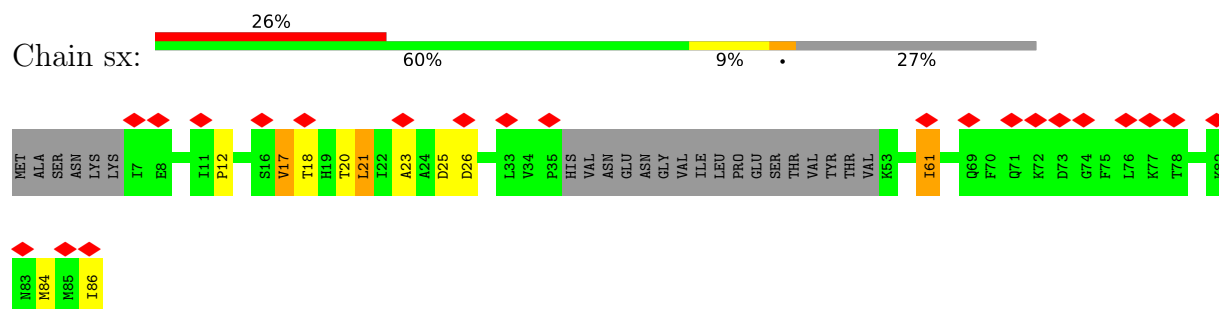
• Molecule 71: Small ribosomal subunit protein eS19



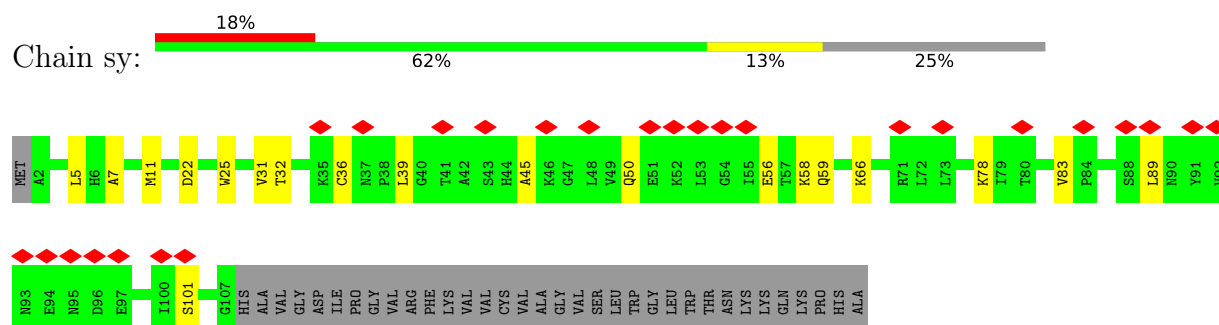
• Molecule 72: 40S ribosomal protein S20, putative



• Molecule 73: 40S ribosomal protein S21



• Molecule 74: 40S ribosomal protein S23, putative



4 Experimental information

| Property | Value | Source |
|--------------------------------------|---|-----------|
| EM reconstruction method | SINGLE PARTICLE | Depositor |
| Imposed symmetry | POINT, Not provided | |
| Number of particles used | 54889 | Depositor |
| Resolution determination method | FSC 0.143 CUT-OFF | Depositor |
| CTF correction method | PHASE FLIPPING AND AMPLITUDE CORRECTION | Depositor |
| Microscope | TFS KRIOS | Depositor |
| Voltage (kV) | 300 | Depositor |
| Electron dose ($e^-/\text{\AA}^2$) | 1.09 | Depositor |
| Minimum defocus (nm) | 1800 | Depositor |
| Maximum defocus (nm) | 3000 | Depositor |
| Magnification | 75000 | Depositor |
| Image detector | FEI FALCON III (4k x 4k) | Depositor |
| Maximum map value | 23.075 | Depositor |
| Minimum map value | -9.423 | Depositor |
| Average map value | -0.000 | Depositor |
| Map value standard deviation | 1.000 | Depositor |
| Recommended contour level | 2.8 | Depositor |
| Map size (Å) | 428.00003, 428.00003, 428.00003 | wwPDB |
| Map dimensions | 400, 400, 400 | wwPDB |
| Map angles (°) | 90.0, 90.0, 90.0 | wwPDB |
| Pixel spacing (Å) | 1.07, 1.07, 1.07 | Depositor |

5 Model quality

5.1 Standard geometry

Bond lengths and bond angles in the following residue types are not validated in this section: PAR

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 | 0.25 | 0/75250 | 0.33 | 2/117222 (0.0%) |
| 2 | 1B | 0.24 | 0/3470 | 0.32 | 0/5401 |
| 3 | 1C | 0.25 | 0/2765 | 0.38 | 2/4303 (0.0%) |
| 4 | 1D | 0.25 | 0/1920 | 0.32 | 0/2582 |
| 5 | 1E | 0.23 | 0/3140 | 0.30 | 0/4216 |
| 6 | 1F | 0.22 | 0/3339 | 0.32 | 0/4479 |
| 7 | 1G | 0.19 | 0/2248 | 0.28 | 0/3013 |
| 8 | 1H | 0.17 | 0/1640 | 0.33 | 0/2204 |
| 9 | 1I | 0.22 | 0/1680 | 0.29 | 0/2252 |
| 10 | 1J | 0.17 | 0/1670 | 0.29 | 0/2243 |
| 11 | 1K | 0.19 | 0/1562 | 0.27 | 0/2103 |
| 12 | 1L | 0.20 | 0/1633 | 0.30 | 0/2184 |
| 13 | 1M | 0.15 | 0/1369 | 0.24 | 0/1834 |
| 14 | 1N | 0.19 | 0/2158 | 0.28 | 0/2875 |
| 15 | 1O | 0.23 | 0/1646 | 0.30 | 0/2209 |
| 16 | 1P | 0.21 | 0/1033 | 0.28 | 0/1389 |
| 17 | 1Q | 0.26 | 0/1707 | 0.29 | 0/2276 |
| 18 | 1R | 0.25 | 0/1251 | 0.29 | 0/1675 |
| 19 | 1S | 0.24 | 0/1342 | 0.32 | 0/1796 |
| 20 | 1T | 0.22 | 0/1445 | 0.29 | 0/1946 |
| 21 | 1U | 0.21 | 0/1253 | 0.28 | 0/1666 |
| 22 | 1V | 0.22 | 0/1351 | 0.30 | 0/1819 |
| 23 | 1W | 0.14 | 0/774 | 0.37 | 0/1031 |
| 24 | 1X | 0.23 | 0/1030 | 0.33 | 0/1384 |
| 25 | 1Y | 0.19 | 0/941 | 0.25 | 0/1262 |
| 26 | 1Z | 0.20 | 0/492 | 0.28 | 0/656 |
| 27 | 1a | 0.18 | 0/1673 | 0.26 | 0/2236 |
| 28 | 1b | 0.17 | 0/1112 | 0.26 | 0/1489 |
| 29 | 1c | 0.26 | 0/1223 | 0.29 | 0/1636 |
| 30 | 1d | 0.22 | 0/485 | 0.25 | 0/639 |
| 31 | 1e | 0.20 | 0/701 | 0.29 | 0/945 |
| 32 | 1f | 0.22 | 0/1050 | 0.27 | 0/1402 |

| Mol | Chain | Bond lengths | | Bond angles | |
|-----|-------|--------------|----------|-------------|-----------------|
| | | RMSZ | # Z >5 | RMSZ | # Z >5 |
| 33 | lg | 0.24 | 0/1027 | 0.28 | 0/1370 |
| 34 | lh | 0.22 | 0/818 | 0.29 | 0/1094 |
| 35 | li | 0.17 | 0/984 | 0.22 | 0/1310 |
| 36 | lj | 0.27 | 0/862 | 0.30 | 0/1163 |
| 37 | lk | 0.17 | 0/721 | 0.24 | 0/955 |
| 38 | ll | 0.27 | 0/602 | 0.35 | 0/797 |
| 39 | lm | 0.22 | 0/701 | 0.34 | 0/934 |
| 40 | ln | 0.18 | 0/592 | 0.29 | 0/789 |
| 41 | lo | 0.26 | 0/444 | 0.29 | 0/587 |
| 42 | lp | 0.23 | 0/425 | 0.48 | 1/563 (0.2%) |
| 43 | lq | 0.23 | 0/770 | 0.27 | 0/1019 |
| 44 | sA | 0.14 | 0/94 | 0.44 | 0/128 |
| 45 | sB | 0.14 | 0/797 | 0.25 | 0/1062 |
| 46 | sC | 0.13 | 0/255 | 0.29 | 0/346 |
| 47 | sD | 0.15 | 0/470 | 0.30 | 0/630 |
| 48 | sE | 0.14 | 0/449 | 0.31 | 0/595 |
| 49 | sI | 0.16 | 0/1737 | 0.38 | 0/2702 |
| 50 | sJ | 0.17 | 0/1727 | 0.40 | 0/2684 |
| 51 | sK | 0.16 | 0/241 | 0.27 | 0/373 |
| 52 | sa | 0.15 | 0/33554 | 0.28 | 0/52257 |
| 53 | sc | 0.13 | 0/1528 | 0.24 | 0/2064 |
| 54 | sd | 0.11 | 0/1456 | 0.27 | 0/1949 |
| 55 | se | 0.14 | 0/1174 | 0.31 | 0/1555 |
| 56 | sf | 0.11 | 0/2072 | 0.28 | 0/2792 |
| 57 | sg | 0.11 | 0/1443 | 0.26 | 0/1938 |
| 58 | sh | 0.11 | 0/402 | 0.26 | 0/535 |
| 59 | si | 0.11 | 0/453 | 0.28 | 0/610 |
| 60 | sj | 0.12 | 0/1021 | 0.27 | 0/1366 |
| 61 | sk | 0.09 | 0/688 | 0.25 | 0/916 |
| 62 | sl | 0.11 | 0/549 | 0.26 | 0/740 |
| 63 | sm | 0.12 | 0/1187 | 0.25 | 0/1586 |
| 64 | so | 0.13 | 0/654 | 0.22 | 0/866 |
| 65 | sp | 0.14 | 0/977 | 0.28 | 0/1311 |
| 66 | sq | 0.12 | 0/856 | 0.33 | 0/1143 |
| 67 | sr | 0.15 | 0/1040 | 0.31 | 0/1404 |
| 68 | ss | 0.11 | 0/1093 | 0.25 | 0/1466 |
| 69 | st | 0.11 | 0/684 | 0.24 | 0/914 |
| 70 | su | 0.13 | 0/1020 | 0.33 | 0/1364 |
| 71 | sv | 0.11 | 0/1034 | 0.23 | 0/1388 |
| 72 | sw | 0.12 | 0/382 | 0.24 | 0/512 |
| 73 | sx | 0.13 | 0/502 | 0.28 | 0/676 |
| 74 | sy | 0.12 | 0/848 | 0.30 | 0/1131 |
| All | All | 0.21 | 0/192686 | 0.31 | 5/283951 (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 |
|-----|-------|---------------------|---------------------|
| 6 | lF | 0 | 1 |
| 38 | ll | 0 | 1 |
| 48 | sE | 0 | 1 |
| 67 | sr | 0 | 1 |
| All | All | 0 | 4 |

There are no bond length outliers.

All (5) bond angle outliers are listed below:

| Mol | Chain | Res | Type | Atoms | Z | Observed(°) | Ideal(°) |
|-----|-------|------|------|-----------|-------|-------------|----------|
| 3 | lC | 62 | U | OP2-P-O3' | -8.61 | 82.16 | 108.00 |
| 3 | lC | 62 | U | OP1-P-O3' | -8.53 | 82.42 | 108.00 |
| 42 | lp | 24 | CYS | CA-CB-SG | 7.16 | 130.87 | 114.40 |
| 1 | lA | 2468 | C | P-O3'-C3' | -5.13 | 112.50 | 120.20 |
| 1 | lA | 2468 | C | O3'-P-O5' | 5.10 | 111.66 | 104.00 |

There are no chirality outliers.

All (4) planarity outliers are listed below:

| Mol | Chain | Res | Type | Group |
|-----|-------|-----|------|---------|
| 6 | lF | 368 | ALA | Peptide |
| 38 | ll | 39 | TYR | Peptide |
| 48 | sE | 15 | GLY | Peptide |
| 67 | sr | 76 | SER | Peptide |

5.2 Too-close contacts

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 | lA | 67153 | 0 | 33687 | 1002 | 0 |
| 2 | lB | 3097 | 0 | 1552 | 49 | 0 |
| 3 | lC | 2477 | 0 | 1252 | 71 | 0 |

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| Mol | Chain | Non-H | H(model) | H(added) | Clashes | Symm-Clashes |
|-----|-------|-------|----------|----------|---------|--------------|
| 4 | ID | 1881 | 0 | 1928 | 53 | 0 |
| 5 | IE | 3076 | 0 | 3209 | 63 | 0 |
| 6 | IF | 3281 | 0 | 3513 | 75 | 0 |
| 7 | IG | 2209 | 0 | 2287 | 43 | 0 |
| 8 | IH | 1608 | 0 | 1728 | 33 | 0 |
| 9 | II | 1658 | 0 | 1802 | 31 | 0 |
| 10 | IJ | 1640 | 0 | 1749 | 20 | 0 |
| 11 | IK | 1538 | 0 | 1598 | 28 | 0 |
| 12 | IL | 1597 | 0 | 1654 | 29 | 0 |
| 13 | IM | 1350 | 0 | 1390 | 34 | 0 |
| 14 | IN | 2130 | 0 | 2338 | 41 | 0 |
| 15 | IO | 1616 | 0 | 1700 | 25 | 0 |
| 16 | IP | 1021 | 0 | 1107 | 25 | 0 |
| 17 | IQ | 1676 | 0 | 1777 | 45 | 0 |
| 18 | IR | 1232 | 0 | 1307 | 22 | 0 |
| 19 | IS | 1321 | 0 | 1427 | 25 | 0 |
| 20 | IT | 1413 | 0 | 1479 | 26 | 0 |
| 21 | IU | 1235 | 0 | 1369 | 25 | 0 |
| 22 | IV | 1320 | 0 | 1406 | 23 | 0 |
| 23 | IW | 763 | 0 | 818 | 20 | 0 |
| 24 | IX | 1015 | 0 | 1054 | 23 | 0 |
| 25 | IY | 926 | 0 | 997 | 16 | 0 |
| 26 | IZ | 481 | 0 | 518 | 11 | 0 |
| 27 | la | 1651 | 0 | 1822 | 30 | 0 |
| 28 | lb | 1094 | 0 | 1174 | 27 | 0 |
| 29 | lc | 1192 | 0 | 1205 | 18 | 0 |
| 30 | ld | 478 | 0 | 507 | 13 | 0 |
| 31 | le | 693 | 0 | 721 | 9 | 0 |
| 32 | lf | 1032 | 0 | 1095 | 20 | 0 |
| 33 | lg | 1010 | 0 | 1091 | 16 | 0 |
| 34 | lh | 805 | 0 | 849 | 19 | 0 |
| 35 | li | 974 | 0 | 1093 | 13 | 0 |
| 36 | lj | 841 | 0 | 878 | 7 | 0 |
| 37 | lk | 712 | 0 | 755 | 21 | 0 |
| 38 | ll | 591 | 0 | 617 | 16 | 0 |
| 39 | lm | 693 | 0 | 738 | 10 | 0 |
| 40 | ln | 584 | 0 | 643 | 13 | 0 |
| 41 | lo | 432 | 0 | 444 | 10 | 0 |
| 42 | lp | 420 | 0 | 450 | 16 | 0 |
| 43 | lq | 756 | 0 | 821 | 16 | 0 |
| 44 | sA | 91 | 0 | 88 | 2 | 0 |
| 45 | sB | 787 | 0 | 833 | 18 | 0 |

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| Mol | Chain | Non-H | H(model) | H(added) | Clashes | Symm-Clashes |
|-----|-------|--------|----------|----------|---------|--------------|
| 46 | sC | 249 | 0 | 259 | 6 | 0 |
| 47 | sD | 468 | 0 | 500 | 15 | 0 |
| 48 | sE | 442 | 0 | 444 | 15 | 0 |
| 49 | sI | 1556 | 0 | 795 | 18 | 0 |
| 50 | sJ | 1549 | 0 | 789 | 20 | 0 |
| 51 | sK | 215 | 0 | 108 | 2 | 0 |
| 52 | sa | 29968 | 0 | 15064 | 504 | 0 |
| 53 | sc | 1499 | 0 | 1565 | 23 | 0 |
| 54 | sd | 1440 | 0 | 1522 | 29 | 0 |
| 55 | se | 1166 | 0 | 1227 | 29 | 0 |
| 56 | sf | 2031 | 0 | 2145 | 63 | 0 |
| 57 | sg | 1424 | 0 | 1485 | 38 | 0 |
| 58 | sh | 400 | 0 | 415 | 12 | 0 |
| 59 | si | 445 | 0 | 466 | 10 | 0 |
| 60 | sj | 1009 | 0 | 1055 | 40 | 0 |
| 61 | sk | 677 | 0 | 724 | 14 | 0 |
| 62 | sl | 536 | 0 | 559 | 19 | 0 |
| 63 | sm | 1161 | 0 | 1181 | 28 | 0 |
| 64 | so | 644 | 0 | 664 | 15 | 0 |
| 65 | sp | 964 | 0 | 988 | 27 | 0 |
| 66 | sq | 842 | 0 | 913 | 30 | 0 |
| 67 | sr | 1022 | 0 | 1051 | 28 | 0 |
| 68 | ss | 1076 | 0 | 1151 | 29 | 0 |
| 69 | st | 676 | 0 | 718 | 19 | 0 |
| 70 | su | 1006 | 0 | 1049 | 28 | 0 |
| 71 | sv | 1015 | 0 | 1043 | 20 | 0 |
| 72 | sw | 376 | 0 | 405 | 16 | 0 |
| 73 | sx | 494 | 0 | 508 | 8 | 0 |
| 74 | sy | 836 | 0 | 885 | 14 | 0 |
| 75 | sa | 42 | 0 | 45 | 1 | 0 |
| All | All | 178778 | 0 | 130193 | 2752 | 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 9.

All (2752) 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:lA:805:G:H21 | 1:lA:1496:A:N6 | 1.45 | 1.13 |
| 1:lA:869:G:N2 | 1:lA:872:A:H62 | 1.48 | 1.11 |
| 52:sa:139:G:H1 | 52:sa:166:A:N6 | 1.48 | 1.11 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 1:lA:869:G:H21 | 1:lA:872:A:N6 | 1.52 | 1.07 |
| 3:lC:79:G:N1 | 3:lC:95:A:C2 | 2.24 | 1.03 |
| 48:sE:2:GLY:N | 48:sE:5:SER:HG | 1.57 | 1.01 |
| 3:lC:27:G:H1 | 3:lC:50:A:H61 | 1.04 | 1.00 |
| 3:lC:27:G:H1 | 3:lC:50:A:N6 | 1.60 | 0.99 |
| 1:lA:869:G:N2 | 1:lA:872:A:N6 | 2.08 | 0.99 |
| 1:lA:805:G:N2 | 1:lA:1496:A:H61 | 1.61 | 0.98 |
| 52:sa:211:U:H3 | 52:sa:247:G:H1 | 1.15 | 0.94 |
| 3:lC:79:G:N1 | 3:lC:95:A:H2 | 1.61 | 0.91 |
| 19:lS:73:HIS:HB3 | 19:lS:76:GLU:HG3 | 1.51 | 0.91 |
| 1:lA:3454:G:H1 | 1:lA:3501:A:H2 | 0.91 | 0.89 |
| 1:lA:2269:U:H5' | 1:lA:2270:G:H5' | 1.54 | 0.89 |
| 1:lA:805:G:H21 | 1:lA:1496:A:H61 | 0.94 | 0.88 |
| 1:lA:3454:G:N1 | 1:lA:3501:A:C2 | 2.41 | 0.88 |
| 1:lA:457:G:N2 | 18:lR:5:CYS:SG | 2.47 | 0.87 |
| 1:lA:2214:A:HO2' | 38:ll:2:THR:N | 1.72 | 0.87 |
| 1:lA:805:G:N2 | 1:lA:1496:A:N6 | 2.19 | 0.87 |
| 6:lF:101:MET:HE3 | 6:lF:104:PRO:HA | 1.57 | 0.87 |
| 1:lA:206:U:H3 | 1:lA:303:G:H1 | 0.88 | 0.87 |
| 1:lA:2366:C:O2 | 52:sa:1892:G:N2 | 2.07 | 0.86 |
| 3:lC:79:G:H1 | 3:lC:95:A:H2 | 0.88 | 0.85 |
| 1:lA:3454:G:N1 | 1:lA:3501:A:H2 | 1.74 | 0.85 |
| 3:lC:52:G:N2 | 3:lC:54:U:O4 | 2.09 | 0.85 |
| 1:lA:873:U:H2' | 1:lA:874:A:H8 | 1.42 | 0.84 |
| 52:sa:1821:A:N6 | 52:sa:1892:G:O6 | 2.10 | 0.83 |
| 58:sh:63:MET:HE1 | 58:sh:106:ILE:HD12 | 1.62 | 0.81 |
| 1:lA:2473:A:N6 | 1:lA:3012:U:O2' | 2.14 | 0.81 |
| 13:lM:18:VAL:HG22 | 13:lM:70:THR:HG22 | 1.61 | 0.81 |
| 1:lA:774:U:OP1 | 29:lc:21:ARG:NH2 | 2.13 | 0.81 |
| 3:lC:79:G:O6 | 3:lC:95:A:N1 | 2.13 | 0.81 |
| 1:lA:1605:A:H2 | 1:lA:3430:U:H3 | 1.27 | 0.81 |
| 50:sJ:15:G:H21 | 50:sJ:16:C:H42 | 1.29 | 0.80 |
| 1:lA:1434:G:HO2' | 1:lA:2456:U:HO2' | 1.25 | 0.80 |
| 36:lj:5:ARG:NH1 | 36:lj:7:HIS:O | 2.15 | 0.80 |
| 1:lA:3479:A:H2' | 1:lA:3480:A:O4' | 1.82 | 0.79 |
| 1:lA:2416:U:O2' | 1:lA:3246:A:N1 | 2.14 | 0.79 |
| 1:lA:409:G:OP2 | 38:ll:56:ARG:NH2 | 2.16 | 0.79 |
| 22:IV:80:VAL:HG12 | 22:IV:81:ASN:H | 1.47 | 0.79 |
| 57:sg:53:HIS:HB2 | 57:sg:92:MET:HE1 | 1.63 | 0.79 |
| 73:sx:18:THR:HG23 | 73:sx:20:THR:HG23 | 1.64 | 0.79 |
| 1:lA:225:U:H3 | 1:lA:282:G:H1 | 1.28 | 0.79 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|--------------------|--------------------------|-------------------|
| 1:1A:3454:G:O6 | 1:1A:3501:A:N1 | 2.15 | 0.79 |
| 1:1A:394:A:N3 | 1:1A:398:A:O2' | 2.16 | 0.78 |
| 52:sa:389:U:H3 | 52:sa:396:A:H62 | 1.32 | 0.78 |
| 3:1C:26:U:H3 | 3:1C:51:G:H22 | 1.31 | 0.77 |
| 3:1C:27:G:N2 | 3:1C:50:A:N1 | 2.32 | 0.77 |
| 1:1A:500:G:N2 | 1:1A:572:U:O2 | 2.18 | 0.76 |
| 52:sa:1245:C:H2' | 52:sa:1246:A:H8 | 1.49 | 0.76 |
| 52:sa:734:A:N1 | 52:sa:803:G:O6 | 2.19 | 0.76 |
| 62:sl:31:HIS:HD2 | 62:sl:41:VAL:HG21 | 1.50 | 0.76 |
| 50:sJ:53:G:H1 | 50:sJ:63:C:H42 | 1.34 | 0.76 |
| 27:la:21:ALA:O | 27:la:26:ARG:NH1 | 2.19 | 0.76 |
| 7:1G:165:LYS:NZ | 7:1G:169:ASP:OD2 | 2.19 | 0.75 |
| 52:sa:43:U:OP2 | 52:sa:432:A:N6 | 2.19 | 0.75 |
| 1:1A:2340:U:O2' | 1:1A:2341:C:O5' | 2.03 | 0.75 |
| 1:1A:2414:C:OP1 | 5:1E:238:ARG:NH1 | 2.20 | 0.75 |
| 1:1A:2688:G:O2' | 1:1A:3008:U:OP1 | 2.04 | 0.75 |
| 1:1A:3263:G:OP2 | 42:lp:37:LYS:NZ | 2.18 | 0.75 |
| 1:1A:3040:A:H5'' | 42:lp:50:LYS:HG3 | 1.68 | 0.75 |
| 3:1C:18:U:H3 | 3:1C:58:A:H61 | 1.35 | 0.75 |
| 1:1A:1251:G:H21 | 12:1L:116:ARG:HH22 | 1.34 | 0.74 |
| 1:1A:1177:A:H2' | 1:1A:1178:A:C8 | 2.22 | 0.74 |
| 1:1A:1818:C:OP2 | 34:lh:74:ARG:NH2 | 2.20 | 0.74 |
| 1:1A:1030:C:OP2 | 4:1D:9:ARG:NH1 | 2.20 | 0.74 |
| 8:1H:76:LYS:O | 8:1H:98:ASN:ND2 | 2.21 | 0.74 |
| 1:1A:2075:A:OP2 | 21:1U:20:ARG:NH1 | 2.20 | 0.74 |
| 3:1C:10:A:N6 | 7:1G:13:PHE:O | 2.21 | 0.74 |
| 17:1Q:31:ARG:NH1 | 17:1Q:124:ASP:OD2 | 2.21 | 0.74 |
| 1:1A:1512:G:OP2 | 6:1F:191:ARG:NH1 | 2.21 | 0.73 |
| 1:1A:2705:A:H4' | 1:1A:2706:C:H5' | 1.68 | 0.73 |
| 1:1A:1033:A:N1 | 4:1D:204:MET:HE2 | 2.03 | 0.73 |
| 1:1A:1282:G:OP1 | 9:1I:78:ARG:NH1 | 2.22 | 0.73 |
| 1:1A:875:C:H2' | 1:1A:876:A:H8 | 1.53 | 0.73 |
| 62:sl:3:ILE:HG12 | 62:sl:40:GLN:HB3 | 1.70 | 0.73 |
| 1:1A:3437:C:N3 | 18:1R:69:ARG:NH2 | 2.35 | 0.73 |
| 1:1A:3479:A:H3' | 1:1A:3479:A:N3 | 2.03 | 0.73 |
| 34:lh:39:SER:OG | 34:lh:58:ARG:NH1 | 2.21 | 0.73 |
| 52:sa:887:A:H2' | 52:sa:888:A:H8 | 1.53 | 0.73 |
| 1:1A:737:A:H3' | 1:1A:738:G:H21 | 1.53 | 0.72 |
| 61:sk:7:ASN:O | 61:sk:8:HIS:ND1 | 2.22 | 0.72 |
| 35:li:76:THR:O | 35:li:81:ARG:NH1 | 2.22 | 0.72 |
| 60:sj:67:TRP:HE1 | 60:sj:69:SER:HB3 | 1.53 | 0.72 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:lA:2629:A:H8 | 31:le:54:PRO:HB3 | 1.52 | 0.72 |
| 52:sa:783:A:N7 | 52:sa:784:U:O2' | 2.20 | 0.72 |
| 59:si:199:VAL:HG23 | 59:si:200:VAL:HG23 | 1.71 | 0.72 |
| 52:sa:139:G:N2 | 52:sa:166:A:N1 | 2.34 | 0.72 |
| 1:lA:919:G:OP1 | 14:lN:17:ARG:NH2 | 2.22 | 0.72 |
| 7:lG:34:ARG:NH1 | 22:lV:26:ASN:O | 2.22 | 0.72 |
| 52:sa:1350:A:H2' | 52:sa:1351:A:H8 | 1.53 | 0.72 |
| 52:sa:1839:A:OP1 | 58:sh:94:ARG:NH2 | 2.23 | 0.71 |
| 37:lk:2:ALA:O | 37:lk:4:GLY:N | 2.22 | 0.71 |
| 72:sw:68:THR:OG1 | 72:sw:70:CYS:O | 2.07 | 0.71 |
| 1:lA:500:G:H1 | 1:lA:572:U:H3 | 0.75 | 0.71 |
| 1:lA:873:U:H2' | 1:lA:874:A:C8 | 2.24 | 0.71 |
| 1:lA:1086:G:H2' | 1:lA:1087:G:H8 | 1.56 | 0.71 |
| 1:lA:1086:G:H1 | 1:lA:1240:C:H5 | 1.38 | 0.71 |
| 52:sa:103:U:OP2 | 52:sa:305:A:N6 | 2.24 | 0.71 |
| 52:sa:1580:U:H1' | 57:sg:90:LEU:HD21 | 1.73 | 0.71 |
| 8:lH:56:ARG:NH2 | 8:lH:107:GLN:O | 2.23 | 0.71 |
| 52:sa:630:A:H5'' | 67:sr:31:SER:HB2 | 1.71 | 0.71 |
| 1:lA:542:U:H2' | 1:lA:543:A:H8 | 1.55 | 0.71 |
| 1:lA:1937:A:OP2 | 40:ln:33:LYS:NZ | 2.20 | 0.71 |
| 3:lC:68:C:H42 | 3:lC:105:A:H61 | 1.39 | 0.71 |
| 52:sa:1716:A:OP1 | 66:sq:41:ARG:NH1 | 2.24 | 0.71 |
| 21:lU:99:MET:SD | 21:lU:103:ARG:NH2 | 2.56 | 0.71 |
| 74:sy:66:LYS:HB3 | 74:sy:89:LEU:HD22 | 1.73 | 0.71 |
| 3:lC:69:U:H3 | 3:lC:104:G:H1 | 1.37 | 0.70 |
| 55:se:28:LYS:NZ | 55:se:51:ASN:OD1 | 2.23 | 0.70 |
| 70:su:36:GLY:HA3 | 70:su:100:ILE:HA | 1.73 | 0.70 |
| 1:lA:869:G:H21 | 1:lA:872:A:H62 | 1.11 | 0.70 |
| 1:lA:1555:U:H2' | 1:lA:1556:G:H8 | 1.56 | 0.70 |
| 1:lA:1634:G:O6 | 41:lo:2:THR:N | 2.24 | 0.70 |
| 52:sa:1395:A:H2' | 52:sa:1396:A:C8 | 2.25 | 0.70 |
| 68:ss:35:CYS:SG | 68:ss:36:LEU:N | 2.64 | 0.70 |
| 1:lA:869:G:N2 | 1:lA:872:A:C6 | 2.58 | 0.70 |
| 1:lA:3491:C:O2' | 5:lE:384:MET:SD | 2.50 | 0.70 |
| 1:lA:3212:U:OP1 | 32:lf:55:HIS:NE2 | 2.25 | 0.70 |
| 5:lE:58:ARG:NH1 | 5:lE:353:GLU:OE2 | 2.25 | 0.70 |
| 1:lA:3455:G:H22 | 1:lA:3500:A:H2 | 1.38 | 0.70 |
| 4:lD:54:ARG:HG3 | 4:lD:56:ALA:H | 1.56 | 0.70 |
| 52:sa:13:C:H4' | 52:sa:1323:U:H3 | 1.57 | 0.70 |
| 52:sa:998:U:OP1 | 64:so:107:LYS:NZ | 2.25 | 0.70 |
| 1:lA:871:A:H2' | 1:lA:872:A:C8 | 2.27 | 0.69 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 23:IW:93:MET:HE2 | 23:IW:93:MET:HA | 1.72 | 0.69 |
| 52:sa:1585:U:OP1 | 71:sv:40:THR:OG1 | 2.09 | 0.69 |
| 1:lA:1944:C:OP1 | 23:IW:107:ARG:NH2 | 2.24 | 0.69 |
| 48:sE:17:GLY:HA2 | 48:sE:27:ARG:HD2 | 1.75 | 0.69 |
| 54:sd:137:MET:HE2 | 54:sd:168:GLU:HG3 | 1.74 | 0.69 |
| 17:lQ:155:LYS:O | 17:lQ:162:ARG:NH2 | 2.26 | 0.69 |
| 55:se:148:LYS:HB2 | 55:se:151:GLN:HB2 | 1.74 | 0.69 |
| 1:lA:639:G:H22 | 1:lA:650:A:H2 | 1.39 | 0.69 |
| 1:lA:816:G:N2 | 1:lA:829:G:OP2 | 2.23 | 0.69 |
| 1:lA:3329:U:HO2' | 1:lA:3411:U:HO2' | 1.33 | 0.69 |
| 1:lA:36:A:O2' | 1:lA:1056:G:O6 | 2.10 | 0.69 |
| 1:lA:1892:C:H2' | 1:lA:1893:A:H8 | 1.58 | 0.69 |
| 17:lQ:163:GLY:O | 17:lQ:172:ARG:NH1 | 2.25 | 0.69 |
| 71:sv:97:SER:O | 71:sv:101:ASN:ND2 | 2.26 | 0.69 |
| 30:ld:7:ARG:NH2 | 30:ld:9:SER:OG | 2.25 | 0.69 |
| 16:lP:25:ALA:HA | 16:lP:40:GLY:HA3 | 1.75 | 0.68 |
| 52:sa:1281:A:H4' | 52:sa:1282:U:H5' | 1.75 | 0.68 |
| 1:lA:3458:U:H3 | 1:lA:3494:G:H1 | 1.41 | 0.68 |
| 20:IT:78:VAL:HG22 | 20:IT:83:ILE:HG12 | 1.75 | 0.68 |
| 1:lA:1883:A:H62 | 1:lA:1931:U:H3 | 1.42 | 0.68 |
| 1:lA:1740:G:OP2 | 34:lh:37:LYS:NZ | 2.24 | 0.68 |
| 1:lA:1841:G:N2 | 1:lA:1975:G:H22 | 1.92 | 0.68 |
| 1:lA:3147:U:OP1 | 5:lE:141:LYS:NZ | 2.23 | 0.68 |
| 50:sJ:34:U:OP2 | 68:ss:158:ARG:NH2 | 2.27 | 0.68 |
| 1:lA:626:A:OP1 | 1:lA:664:A:N6 | 2.26 | 0.68 |
| 49:sI:16:G:H5'' | 49:sI:17:G:H5' | 1.76 | 0.68 |
| 1:lA:869:G:N2 | 1:lA:872:A:C5 | 2.62 | 0.68 |
| 70:su:81:PRO:HG2 | 70:su:84:PHE:HB2 | 1.75 | 0.68 |
| 52:sa:286:U:H2' | 52:sa:287:A:H8 | 1.58 | 0.68 |
| 1:lA:112:G:O2' | 1:lA:311:A:N3 | 2.27 | 0.67 |
| 1:lA:1841:G:H22 | 1:lA:1975:G:H22 | 1.43 | 0.67 |
| 13:lM:116:TYR:OH | 70:su:109:ARG:NH1 | 2.27 | 0.67 |
| 52:sa:1712:A:OP1 | 70:su:122:ARG:NH1 | 2.26 | 0.67 |
| 1:lA:1442:A:N6 | 15:lO:18:GLY:O | 2.26 | 0.67 |
| 1:lA:2450:C:O2 | 5:lE:258:HIS:NE2 | 2.27 | 0.67 |
| 5:lE:219:VAL:HG12 | 5:lE:339:LEU:HD23 | 1.75 | 0.67 |
| 6:lF:101:MET:HE1 | 6:lF:105:LEU:HG | 1.75 | 0.67 |
| 20:IT:5:TYR:HB2 | 20:IT:60:ILE:HD11 | 1.76 | 0.67 |
| 3:lC:79:G:C6 | 3:lC:95:A:N1 | 2.63 | 0.67 |
| 6:lF:377:PHE:HE1 | 16:lP:82:GLN:HG2 | 1.60 | 0.67 |
| 29:lc:59:ARG:NH2 | 43:lq:36:GLN:OE1 | 2.28 | 0.67 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 57:sg:52:PRO:HG2 | 57:sg:92:MET:HE2 | 1.76 | 0.67 |
| 1:lA:494:G:N2 | 1:lA:578:A:OP2 | 2.28 | 0.67 |
| 21:lU:99:MET:HE1 | 21:lU:128:LYS:HA | 1.77 | 0.67 |
| 52:sa:751:A:N1 | 52:sa:785:A:N6 | 2.42 | 0.67 |
| 52:sa:783:A:H3' | 52:sa:784:U:H4' | 1.76 | 0.67 |
| 68:ss:54:ILE:O | 68:ss:60:ARG:NH1 | 2.24 | 0.67 |
| 1:lA:95:A:OP1 | 17:lQ:195:ARG:NH1 | 2.25 | 0.67 |
| 1:lA:450:U:H4' | 1:lA:1550:G:H4' | 1.76 | 0.67 |
| 1:lA:893:U:H3 | 1:lA:897:A:H2 | 1.41 | 0.67 |
| 49:sI:63:U:H2' | 49:sI:64:G:H8 | 1.58 | 0.67 |
| 52:sa:115:U:O2' | 52:sa:328:C:O2 | 2.13 | 0.67 |
| 1:lA:3211:A:H2 | 1:lA:3239:G:H21 | 1.43 | 0.67 |
| 48:sE:41:ARG:O | 52:sa:1540:G:N2 | 2.28 | 0.67 |
| 52:sa:356:C:H5' | 52:sa:373:A:H5' | 1.77 | 0.67 |
| 56:sf:104:LYS:O | 56:sf:106:ARG:NH2 | 2.28 | 0.67 |
| 17:lQ:106:VAL:HG21 | 17:lQ:132:VAL:HG21 | 1.76 | 0.66 |
| 52:sa:1591:G:H21 | 52:sa:1772:A:H1' | 1.61 | 0.66 |
| 56:sf:33:PRO:HD2 | 56:sf:81:PRO:HG2 | 1.77 | 0.66 |
| 66:sq:111:LEU:O | 66:sq:113:HIS:N | 2.28 | 0.66 |
| 1:lA:213:A:H5' | 14:lN:133:LYS:H | 1.60 | 0.66 |
| 1:lA:1664:G:OP2 | 1:lA:1752:U:O2' | 2.13 | 0.66 |
| 2:lB:141:G:H2' | 2:lB:142:A:H8 | 1.60 | 0.66 |
| 52:sa:63:U:O2' | 52:sa:163:A:N3 | 2.25 | 0.66 |
| 23:lW:107:ARG:HE | 23:lW:109:ILE:HD11 | 1.60 | 0.66 |
| 1:lA:925:A:N3 | 1:lA:2955:C:O2' | 2.28 | 0.66 |
| 52:sa:90:G:OP1 | 52:sa:392:A:N6 | 2.29 | 0.66 |
| 52:sa:875:G:H21 | 65:sp:47:THR:HG21 | 1.60 | 0.66 |
| 52:sa:956:G:H1 | 52:sa:1003:A:HO2' | 1.43 | 0.66 |
| 66:sq:27:MET:HG2 | 66:sq:28:PRO:HD2 | 1.76 | 0.66 |
| 1:lA:37:G:N2 | 1:lA:2946:A:N7 | 2.42 | 0.66 |
| 1:lA:2415:C:OP1 | 24:IX:54:ARG:NH2 | 2.28 | 0.66 |
| 1:lA:3243:C:OP1 | 5:lE:224:LYS:NZ | 2.22 | 0.66 |
| 10:lJ:168:LYS:O | 10:lJ:171:GLN:NE2 | 2.29 | 0.66 |
| 52:sa:107:A:H2' | 52:sa:108:G:C8 | 2.31 | 0.66 |
| 52:sa:1717:U:OP2 | 66:sq:42:ARG:NH1 | 2.28 | 0.66 |
| 13:lM:43:GLN:NE2 | 13:lM:70:THR:O | 2.28 | 0.66 |
| 1:lA:626:A:N6 | 1:lA:663:G:O2' | 2.29 | 0.66 |
| 27:la:53:ASP:OD2 | 27:la:113:ARG:NH2 | 2.28 | 0.66 |
| 52:sa:394:A:OP1 | 60:sj:49:ARG:NH2 | 2.28 | 0.65 |
| 49:sI:13:C:O2 | 49:sI:14:A:H8 | 1.80 | 0.65 |
| 1:lA:631:G:N2 | 1:lA:632:G:N3 | 2.45 | 0.65 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 1:lA:1841:G:H22 | 1:lA:1975:G:H1 | 1.43 | 0.65 |
| 1:lA:1313:U:O2' | 1:lA:1315:A:N7 | 2.27 | 0.65 |
| 1:lA:1523:U:OP1 | 33:lg:105:LYS:NZ | 2.27 | 0.65 |
| 6:lF:55:SER:HB3 | 6:lF:58:ALA:HB2 | 1.78 | 0.65 |
| 6:lF:110:LYS:O | 6:lF:113:ARG:NH1 | 2.28 | 0.65 |
| 52:sa:626:U:H4' | 74:sy:7:ALA:HB2 | 1.79 | 0.65 |
| 1:lA:2462:G:H2' | 1:lA:2463:A:H8 | 1.61 | 0.65 |
| 24:lX:83:ARG:NH2 | 24:lX:118:THR:O | 2.28 | 0.65 |
| 28:lb:5:LEU:HD11 | 31:le:39:ARG:HD2 | 1.79 | 0.65 |
| 52:sa:1581:G:H2' | 52:sa:1582:G:H8 | 1.59 | 0.65 |
| 1:lA:2173:A:H2' | 1:lA:2174:U:H6 | 1.60 | 0.65 |
| 52:sa:251:C:O2' | 60:sj:64:ASN:ND2 | 2.29 | 0.65 |
| 52:sa:1578:A:OP1 | 57:sg:166:ARG:NH1 | 2.29 | 0.65 |
| 1:lA:2337:G:O2' | 1:lA:2339:C:N4 | 2.29 | 0.65 |
| 1:lA:2865:A:O2' | 1:lA:2866:C:O4' | 2.13 | 0.65 |
| 52:sa:213:A:H62 | 52:sa:243:U:H4' | 1.62 | 0.65 |
| 38:ll:18:LEU:HD11 | 41:lo:51:MET:HE3 | 1.78 | 0.65 |
| 39:lm:85:ARG:NH2 | 52:sa:904:A:OP1 | 2.29 | 0.65 |
| 52:sa:584:C:H2' | 52:sa:585:A:C8 | 2.31 | 0.65 |
| 55:se:63:LEU:O | 55:se:65:GLY:N | 2.30 | 0.65 |
| 65:sp:25:VAL:HG12 | 65:sp:89:HIS:HB2 | 1.79 | 0.65 |
| 66:sq:107:LYS:HG3 | 66:sq:109:GLU:H | 1.61 | 0.65 |
| 56:sf:180:MET:HB2 | 56:sf:190:VAL:HG12 | 1.79 | 0.65 |
| 1:lA:1117:G:N3 | 1:lA:1120:A:N6 | 2.46 | 0.64 |
| 9:lI:109:ALA:HB2 | 22:IV:139:LYS:HB3 | 1.77 | 0.64 |
| 11:lK:100:MET:HB3 | 11:lK:150:ILE:HG13 | 1.79 | 0.64 |
| 16:lP:39:ASP:OD2 | 16:lP:49:ARG:NH2 | 2.30 | 0.64 |
| 52:sa:1816:U:H2' | 52:sa:1817:A:C8 | 2.32 | 0.64 |
| 1:lA:59:A:N3 | 1:lA:74:U:O2' | 2.24 | 0.64 |
| 3:lC:22:C:H2' | 3:lC:23:A:H8 | 1.62 | 0.64 |
| 52:sa:213:A:N6 | 52:sa:244:U:OP2 | 2.31 | 0.64 |
| 7:lG:64:ILE:HD13 | 7:lG:109:LEU:HD22 | 1.79 | 0.64 |
| 52:sa:972:A:O2' | 52:sa:1932:U:O2 | 2.14 | 0.64 |
| 1:lA:1157:A:H4' | 7:lG:5:LYS:H | 1.62 | 0.64 |
| 1:lA:2514:A:H61 | 1:lA:2583:U:H3 | 1.43 | 0.64 |
| 57:sg:63:PHE:HE1 | 57:sg:146:LEU:HD22 | 1.62 | 0.64 |
| 1:lA:1524:A:O2' | 1:lA:1552:A:N1 | 2.29 | 0.64 |
| 53:sc:85:MET:HE2 | 53:sc:123:VAL:HG23 | 1.77 | 0.64 |
| 52:sa:1594:A:H2' | 52:sa:1595:G:H8 | 1.63 | 0.64 |
| 56:sf:139:THR:OG1 | 56:sf:141:ASP:OD1 | 2.16 | 0.64 |
| 69:st:5:ARG:HB2 | 69:st:10:LYS:HE3 | 1.79 | 0.64 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 69:st:74:GLN:NE2 | 69:st:75:GLU:OE2 | 2.30 | 0.64 |
| 1:lA:1670:A:OP2 | 1:lA:1740:G:N2 | 2.30 | 0.64 |
| 52:sa:389:U:O4 | 52:sa:396:A:N7 | 2.30 | 0.64 |
| 52:sa:872:A:H2' | 52:sa:873:A:C8 | 2.33 | 0.64 |
| 57:sg:120:ASN:ND2 | 57:sg:183:SER:O | 2.30 | 0.64 |
| 5:lE:222:VAL:O | 5:lE:335:ARG:NH1 | 2.31 | 0.64 |
| 45:sB:32:LYS:O | 45:sB:37:LYS:NZ | 2.31 | 0.64 |
| 1:lA:3262:U:OP1 | 42:lp:39:LYS:NZ | 2.29 | 0.63 |
| 55:se:42:ARG:O | 55:se:44:THR:N | 2.31 | 0.63 |
| 1:lA:619:A:O2' | 1:lA:620:U:O5' | 2.16 | 0.63 |
| 1:lA:875:C:H2' | 1:lA:876:A:C8 | 2.32 | 0.63 |
| 67:sr:36:GLU:HB3 | 67:sr:110:ILE:HG21 | 1.80 | 0.63 |
| 1:lA:926:A:H62 | 1:lA:1053:G:H1 | 1.46 | 0.63 |
| 1:lA:1327:A:N3 | 1:lA:2998:U:O2' | 2.28 | 0.63 |
| 1:lA:1524:A:OP1 | 33:lg:102:SER:OG | 2.13 | 0.63 |
| 3:lC:104:G:H2' | 3:lC:105:A:H8 | 1.62 | 0.63 |
| 70:su:96:ASP:OD2 | 70:su:98:GLN:NE2 | 2.31 | 0.63 |
| 1:lA:612:A:H62 | 1:lA:715:A:H2' | 1.64 | 0.63 |
| 55:se:174:ILE:HG12 | 55:se:195:LEU:HD11 | 1.79 | 0.63 |
| 5:lE:286:ARG:NH2 | 5:lE:296:LYS:O | 2.32 | 0.63 |
| 52:sa:99:A:H61 | 52:sa:380:A:H1' | 1.63 | 0.63 |
| 52:sa:852:A:H61 | 52:sa:935:C:H42 | 1.46 | 0.63 |
| 1:lA:64:A:OP2 | 1:lA:347:G:N2 | 2.29 | 0.63 |
| 1:lA:195:A:H61 | 1:lA:312:A:H2 | 1.47 | 0.63 |
| 1:lA:2233:G:H4' | 1:lA:2234:A:H5' | 1.80 | 0.63 |
| 38:ll:39:TYR:O | 38:ll:41:ASP:N | 2.31 | 0.63 |
| 4:lD:136:VAL:HA | 4:lD:148:ILE:HG22 | 1.79 | 0.63 |
| 12:IL:171:TRP:CD1 | 12:IL:172:GLY:H | 2.16 | 0.63 |
| 48:sE:14:TYR:HB3 | 52:sa:1763:A:C8 | 2.34 | 0.63 |
| 63:sm:57:CYS:HB3 | 63:sm:60:THR:HG22 | 1.79 | 0.63 |
| 1:lA:3367:U:H2' | 1:lA:3368:G:C8 | 2.34 | 0.63 |
| 52:sa:865:A:H2' | 52:sa:866:G:C8 | 2.34 | 0.63 |
| 1:lA:2114:U:N3 | 1:lA:2198:G:OP2 | 2.27 | 0.63 |
| 1:lA:2173:A:H2' | 1:lA:2174:U:C6 | 2.33 | 0.63 |
| 20:IT:92:ASP:OD2 | 20:IT:101:GLN:NE2 | 2.31 | 0.63 |
| 52:sa:467:C:O2' | 52:sa:760:A:N3 | 2.32 | 0.63 |
| 1:lA:43:C:OP2 | 1:lA:44:A:O2' | 2.11 | 0.62 |
| 1:lA:628:C:O2' | 20:IT:141:HIS:NE2 | 2.32 | 0.62 |
| 1:lA:2257:G:OP1 | 4:lD:193:ARG:NH2 | 2.30 | 0.62 |
| 1:lA:3355:U:O2 | 15:lO:4:ASN:ND2 | 2.31 | 0.62 |
| 52:sa:918:G:N2 | 52:sa:921:A:OP2 | 2.30 | 0.62 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 52:sa:1322:G:N2 | 52:sa:1325:A:OP2 | 2.31 | 0.62 |
| 1:1A:2705:A:C4' | 1:1A:2706:C:H5' | 2.28 | 0.62 |
| 50:sJ:23:G:H2' | 50:sJ:24:C:C6 | 2.34 | 0.62 |
| 53:sc:164:CYS:SG | 53:sc:165:GLY:N | 2.73 | 0.62 |
| 55:se:74:ASP:OD1 | 65:sp:123:ARG:NH2 | 2.32 | 0.62 |
| 5:1E:84:MET:HG2 | 5:1E:166:THR:HA | 1.82 | 0.62 |
| 1:1A:2990:A:OP1 | 42:lp:22:ARG:NH1 | 2.33 | 0.62 |
| 19:1S:60:ILE:HG22 | 19:1S:83:THR:HB | 1.82 | 0.62 |
| 19:1S:152:VAL:HA | 19:1S:159:ALA:H | 1.63 | 0.62 |
| 52:sa:107:A:H2' | 52:sa:108:G:H8 | 1.63 | 0.62 |
| 52:sa:1441:C:OP1 | 69:st:5:ARG:NH2 | 2.32 | 0.62 |
| 54:sd:137:MET:CE | 54:sd:168:GLU:HG3 | 2.30 | 0.62 |
| 12:1L:51:HIS:ND1 | 12:1L:137:SER:OG | 2.22 | 0.62 |
| 55:se:117:LYS:NZ | 55:se:210:ASP:OD2 | 2.33 | 0.62 |
| 1:1A:1072:U:OP1 | 30:ld:12:GLN:NE2 | 2.30 | 0.62 |
| 1:1A:3349:G:OP2 | 16:1P:5:ARG:NH2 | 2.32 | 0.62 |
| 3:1C:26:U:O2 | 3:1C:52:G:N2 | 2.23 | 0.62 |
| 12:1L:112:GLN:OE1 | 49:sI:73:A:O2' | 2.18 | 0.62 |
| 49:sI:63:U:H2' | 49:sI:64:G:C8 | 2.34 | 0.62 |
| 60:sj:36:THR:O | 60:sj:96:LEU:N | 2.33 | 0.62 |
| 12:1L:66:GLU:OE1 | 12:1L:69:ARG:NH2 | 2.32 | 0.62 |
| 52:sa:258:A:OP2 | 52:sa:285:G:N2 | 2.31 | 0.62 |
| 3:1C:33:A:N1 | 3:1C:45:U:O2' | 2.29 | 0.62 |
| 9:1I:76:ARG:NH2 | 9:1I:79:GLY:O | 2.30 | 0.62 |
| 40:ln:60:ILE:H | 40:ln:60:ILE:HD12 | 1.65 | 0.62 |
| 52:sa:938:U:O2' | 52:sa:940:U:OP2 | 2.16 | 0.62 |
| 56:sf:41:PRO:HD2 | 56:sf:44:LEU:HD12 | 1.80 | 0.62 |
| 8:1H:64:ILE:O | 8:1H:113:THR:OG1 | 2.17 | 0.62 |
| 52:sa:42:A:N1 | 52:sa:373:A:N6 | 2.48 | 0.62 |
| 55:se:73:ALA:HB3 | 65:sp:123:ARG:HH22 | 1.65 | 0.62 |
| 61:sk:78:ARG:O | 61:sk:82:ASN:ND2 | 2.33 | 0.62 |
| 7:1G:65:ALA:HB2 | 7:1G:74:ILE:HD13 | 1.82 | 0.62 |
| 52:sa:401:U:H2' | 52:sa:402:A:C8 | 2.34 | 0.62 |
| 3:1C:59:G:H5' | 7:1G:265:LYS:HD3 | 1.82 | 0.61 |
| 68:ss:109:GLN:OE1 | 68:ss:117:LYS:NZ | 2.32 | 0.61 |
| 1:1A:65:U:HO2' | 1:1A:97:A:HO2' | 1.47 | 0.61 |
| 1:1A:964:G:N2 | 1:1A:967:A:OP2 | 2.33 | 0.61 |
| 16:1P:29:GLU:OE2 | 20:1T:91:ARG:NH2 | 2.34 | 0.61 |
| 20:1T:79:SER:OG | 20:1T:84:HIS:NE2 | 2.25 | 0.61 |
| 52:sa:840:U:H2' | 52:sa:841:U:C6 | 2.35 | 0.61 |
| 52:sa:1718:U:O2' | 52:sa:1763:A:N3 | 2.30 | 0.61 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 2:1B:93:C:OP1 | 38:1L:73:ARG:NH1 | 2.33 | 0.61 |
| 25:1Y:78:VAL:HG21 | 25:1Y:103:VAL:HG21 | 1.81 | 0.61 |
| 38:1L:60:GLY:HA2 | 38:1L:64:MET:HE2 | 1.82 | 0.61 |
| 52:sa:145:U:O4 | 52:sa:159:A:N6 | 2.33 | 0.61 |
| 53:sc:58:ILE:HG23 | 53:sc:63:ILE:HB | 1.82 | 0.61 |
| 58:sh:57:ASP:OD2 | 58:sh:98:ARG:NH1 | 2.24 | 0.61 |
| 1:1A:502:U:H2' | 1:1A:503:A:C8 | 2.36 | 0.61 |
| 1:1A:2257:G:H5'' | 4:1D:193:ARG:HH12 | 1.65 | 0.61 |
| 1:1A:200:A:H62 | 1:1A:309:G:H21 | 1.47 | 0.61 |
| 69:st:77:GLU:OE1 | 69:st:80:ARG:NH1 | 2.33 | 0.61 |
| 1:1A:3336:G:O2' | 15:1O:115:LYS:O | 2.18 | 0.61 |
| 8:1H:165:LYS:NZ | 8:1H:166:GLN:OE1 | 2.33 | 0.61 |
| 52:sa:239:U:N3 | 52:sa:242:G:OP2 | 2.29 | 0.61 |
| 52:sa:1556:U:H2' | 52:sa:1557:C:H6 | 1.65 | 0.61 |
| 69:st:43:THR:HG22 | 69:st:46:LEU:H | 1.65 | 0.61 |
| 1:1A:450:U:H2' | 1:1A:451:G:H5' | 1.83 | 0.61 |
| 1:1A:1634:G:N2 | 1:1A:1634:G:OP2 | 2.34 | 0.61 |
| 1:1A:2499:U:H2' | 1:1A:2500:A:H8 | 1.66 | 0.61 |
| 3:1C:75:G:H1 | 3:1C:99:U:H5 | 1.48 | 0.61 |
| 6:1F:237:LEU:HD22 | 6:1F:242:LEU:HD21 | 1.82 | 0.61 |
| 20:1T:40:MET:HE3 | 22:1V:157:VAL:HG12 | 1.83 | 0.61 |
| 52:sa:797:G:OP2 | 52:sa:797:G:N2 | 2.33 | 0.61 |
| 1:1A:1217:U:H4' | 22:1V:123:LYS:HE3 | 1.81 | 0.61 |
| 1:1A:3372:U:O2' | 1:1A:3374:C:OP2 | 2.15 | 0.61 |
| 3:1C:3:G:N2 | 7:1G:79:TYR:OH | 2.34 | 0.61 |
| 43:1q:72:CYS:SG | 43:1q:75:CYS:N | 2.73 | 0.61 |
| 5:1E:232:ILE:HG13 | 5:1E:249:ARG:HB3 | 1.82 | 0.61 |
| 22:1V:88:ARG:NH2 | 30:1d:28:ALA:O | 2.34 | 0.61 |
| 28:1b:10:VAL:O | 28:1b:83:THR:OG1 | 2.18 | 0.61 |
| 50:sJ:24:C:H2' | 50:sJ:25:U:H6 | 1.65 | 0.61 |
| 68:ss:98:GLN:HE22 | 68:ss:134:ALA:HA | 1.65 | 0.61 |
| 1:1A:1303:A:N3 | 1:1A:1453:U:O2' | 2.31 | 0.60 |
| 52:sa:1670:A:H2' | 52:sa:1671:A:C8 | 2.36 | 0.60 |
| 56:sf:209:LYS:NZ | 56:sf:213:GLY:O | 2.33 | 0.60 |
| 1:1A:1252:G:OP2 | 12:1L:98:ARG:NH1 | 2.33 | 0.60 |
| 1:1A:2468:C:O2 | 5:1E:268:ARG:NH2 | 2.34 | 0.60 |
| 3:1C:10:A:N1 | 3:1C:65:U:O2' | 2.30 | 0.60 |
| 32:1f:63:PHE:HB3 | 32:1f:103:ARG:HG2 | 1.82 | 0.60 |
| 62:sl:21:ILE:HD13 | 62:sl:45:MET:HE1 | 1.83 | 0.60 |
| 1:1A:1561:U:OP2 | 29:1c:4:ARG:NH2 | 2.30 | 0.60 |
| 15:1O:82:ARG:NH2 | 15:1O:103:LEU:O | 2.33 | 0.60 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 1:1A:1266:C:O2' | 1:1A:1278:A:N3 | 2.34 | 0.60 |
| 1:1A:2485:G:H3' | 1:1A:2486:U:H4' | 1.83 | 0.60 |
| 34:lh:81:CYS:SG | 34:lh:84:CYS:N | 2.74 | 0.60 |
| 48:sE:21:CYS:HB3 | 48:sE:25:GLY:H | 1.66 | 0.60 |
| 52:sa:584:C:H2' | 52:sa:585:A:H8 | 1.65 | 0.60 |
| 56:sf:98:ARG:HB3 | 56:sf:110:VAL:HG13 | 1.84 | 0.60 |
| 1:1A:3383:A:OP2 | 5:1E:148:ARG:NH2 | 2.30 | 0.60 |
| 52:sa:1907:G:H1' | 52:sa:1928:A:H2 | 1.66 | 0.60 |
| 1:1A:1209:U:OP1 | 1:1A:1211:A:O2' | 2.20 | 0.60 |
| 1:1A:1261:A:OP1 | 30:ld:7:ARG:NH1 | 2.35 | 0.60 |
| 2:1B:154:C:H2' | 2:1B:155:A:C8 | 2.36 | 0.60 |
| 65:sp:56:ARG:NH2 | 65:sp:75:ASP:OD2 | 2.35 | 0.60 |
| 1:1A:1414:A:H2' | 1:1A:1415:A:H8 | 1.66 | 0.60 |
| 1:1A:1476:G:O2' | 1:1A:1477:A:H5' | 2.01 | 0.60 |
| 1:1A:1868:U:OP2 | 23:1W:55:LYS:NZ | 2.32 | 0.60 |
| 1:1A:2272:C:O2' | 1:1A:2346:A:N3 | 2.31 | 0.60 |
| 1:1A:2368:U:H5'' | 52:sa:1824:G:H21 | 1.67 | 0.60 |
| 2:1B:25:U:OP2 | 27:la:15:ARG:NH1 | 2.34 | 0.60 |
| 6:1F:9:ILE:HD13 | 6:1F:256:GLU:HG3 | 1.82 | 0.60 |
| 10:1J:60:LYS:HG3 | 10:1J:232:ARG:HE | 1.67 | 0.60 |
| 11:1K:101:ARG:NH1 | 11:1K:149:GLU:OE1 | 2.34 | 0.60 |
| 57:sg:197:ASP:OD1 | 57:sg:200:ARG:NH2 | 2.34 | 0.60 |
| 65:sp:93:ARG:HB3 | 65:sp:127:VAL:HG23 | 1.82 | 0.60 |
| 70:su:34:ILE:HG22 | 70:su:36:GLY:H | 1.67 | 0.60 |
| 1:1A:2971:G:OP2 | 12:1L:7:ARG:NH2 | 2.35 | 0.60 |
| 52:sa:864:C:H2' | 52:sa:865:A:C8 | 2.37 | 0.60 |
| 61:sk:48:MET:HE2 | 61:sk:48:MET:HA | 1.83 | 0.60 |
| 1:1A:2634:A:OP1 | 4:1D:69:TYR:OH | 2.20 | 0.60 |
| 1:1A:2798:G:OP1 | 19:1S:151:LYS:NZ | 2.35 | 0.60 |
| 1:1A:3247:C:O2 | 24:1X:86:ARG:NH2 | 2.35 | 0.60 |
| 1:1A:3454:G:C6 | 1:1A:3501:A:N1 | 2.69 | 0.60 |
| 11:1K:101:ARG:HG3 | 11:1K:188:SER:HB3 | 1.84 | 0.60 |
| 48:sE:16:ASN:ND2 | 52:sa:1231:U:O2 | 2.35 | 0.60 |
| 52:sa:456:G:H2' | 52:sa:457:G:H8 | 1.66 | 0.60 |
| 1:1A:543:A:H2' | 1:1A:544:U:H6 | 1.67 | 0.60 |
| 1:1A:2414:C:O2 | 24:1X:21:ASN:ND2 | 2.26 | 0.60 |
| 52:sa:440:A:N6 | 52:sa:457:G:N3 | 2.43 | 0.60 |
| 52:sa:1759:A:H2' | 52:sa:1760:G:H8 | 1.67 | 0.60 |
| 52:sa:1823:U:OP1 | 52:sa:1824:G:O2' | 2.18 | 0.60 |
| 54:sd:108:ARG:NH1 | 54:sd:115:GLN:OE1 | 2.35 | 0.60 |
| 1:1A:2432:A:H2 | 18:1R:131:ARG:HH22 | 1.50 | 0.59 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 3:IC:5:A:OP2 | 7:IG:22:ARG:NH2 | 2.35 | 0.59 |
| 3:IC:73:G:O2' | 20:IT:46:LYS:NZ | 2.32 | 0.59 |
| 10:IJ:173:PRO:HG3 | 10:IJ:216:LYS:HE3 | 1.82 | 0.59 |
| 47:sD:21:ARG:HG2 | 47:sD:29:LEU:HD12 | 1.83 | 0.59 |
| 1:lA:718:A:H2' | 1:lA:719:U:C6 | 2.37 | 0.59 |
| 1:lA:3176:G:O2' | 1:lA:3185:G:O6 | 2.19 | 0.59 |
| 25:IY:85:LYS:HG2 | 25:IY:106:ALA:HB2 | 1.83 | 0.59 |
| 35:li:9:LYS:NZ | 35:li:12:GLU:OE2 | 2.28 | 0.59 |
| 52:sa:1668:G:OP2 | 71:sv:102:ARG:NH1 | 2.35 | 0.59 |
| 52:sa:1760:G:OP2 | 52:sa:1762:C:N4 | 2.35 | 0.59 |
| 62:sl:31:HIS:CD2 | 62:sl:41:VAL:HG21 | 2.36 | 0.59 |
| 1:lA:1115:A:O2' | 3:IC:76:U:O2 | 2.19 | 0.59 |
| 1:lA:1174:A:N3 | 3:IC:78:C:O2' | 2.35 | 0.59 |
| 1:lA:3045:A:OP1 | 11:IK:176:LYS:NZ | 2.29 | 0.59 |
| 10:IJ:94:ALA:O | 10:IJ:98:GLN:NE2 | 2.30 | 0.59 |
| 56:sf:240:LYS:HD2 | 56:sf:240:LYS:O | 2.02 | 0.59 |
| 21:IU:44:LEU:HD22 | 21:IU:49:LEU:HD12 | 1.84 | 0.59 |
| 52:sa:1245:C:H2' | 52:sa:1246:A:C8 | 2.36 | 0.59 |
| 53:sc:99:ARG:NH2 | 53:sc:119:CYS:SG | 2.75 | 0.59 |
| 1:lA:457:G:OP1 | 18:IR:62:LYS:NZ | 2.34 | 0.59 |
| 1:lA:779:A:N6 | 1:lA:3012:U:OP1 | 2.35 | 0.59 |
| 1:lA:1177:A:H2' | 1:lA:1178:A:H8 | 1.68 | 0.59 |
| 1:lA:2856:G:H2' | 1:lA:2903:A:H61 | 1.66 | 0.59 |
| 7:IG:122:VAL:O | 7:IG:244:ARG:NH2 | 2.35 | 0.59 |
| 38:ll:21:ARG:NH2 | 38:ll:38:GLY:O | 2.35 | 0.59 |
| 52:sa:602:U:O5' | 52:sa:604:G:N2 | 2.35 | 0.59 |
| 60:sj:87:ASP:OD1 | 60:sj:89:ASP:N | 2.34 | 0.59 |
| 64:so:99:ARG:O | 64:so:101:HIS:N | 2.35 | 0.59 |
| 1:lA:1086:G:H2' | 1:lA:1087:G:C8 | 2.38 | 0.59 |
| 1:lA:1494:G:OP2 | 29:lc:10:ARG:NH2 | 2.35 | 0.59 |
| 1:lA:1744:A:H2' | 1:lA:1745:A:C8 | 2.37 | 0.59 |
| 1:lA:1892:C:H2' | 1:lA:1893:A:C8 | 2.38 | 0.59 |
| 6:lF:80:SER:O | 6:lF:87:ASN:ND2 | 2.35 | 0.59 |
| 32:lf:62:SER:HB2 | 32:lf:65:LYS:HD3 | 1.85 | 0.59 |
| 52:sa:1750:G:N2 | 52:sa:1776:G:O2' | 2.35 | 0.59 |
| 60:sj:44:HIS:N | 60:sj:56:ARG:O | 2.35 | 0.59 |
| 3:IC:17:G:N2 | 3:IC:60:C:O2 | 2.36 | 0.59 |
| 6:lF:77:PRO:HB2 | 6:lF:91:ALA:HB3 | 1.83 | 0.59 |
| 24:IX:9:ARG:NH2 | 24:IX:11:GLY:O | 2.32 | 0.59 |
| 1:lA:231:U:H4' | 27:la:123:SER:HB3 | 1.84 | 0.59 |
| 1:lA:1112:G:O2' | 1:lA:1172:G:O6 | 2.21 | 0.59 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|-------------------|--------------------------|-------------------|
| 1:lA:1259:G:OP2 | 30:ld:5:LYS:NZ | 2.24 | 0.59 |
| 24:IX:57:CYS:HG | 24:IX:122:SER:HG | 1.51 | 0.59 |
| 28:lb:104:GLU:HG2 | 28:lb:107:LYS:HD3 | 1.83 | 0.59 |
| 52:sa:139:G:H1 | 52:sa:166:A:H61 | 0.72 | 0.59 |
| 1:lA:1022:U:HO2' | 1:lA:1677:A:HO2' | 1.47 | 0.59 |
| 1:lA:1181:A:H5'' | 1:lA:1182:A:H5' | 1.85 | 0.59 |
| 5:lE:370:ARG:HG2 | 26:lZ:34:ILE:HD11 | 1.85 | 0.59 |
| 1:lA:3150:A:H2' | 1:lA:3151:A:H8 | 1.66 | 0.58 |
| 5:lE:361:ASP:OD2 | 5:lE:365:LYS:NZ | 2.36 | 0.58 |
| 6:lF:360:LYS:HB3 | 6:lF:400:THR:HG21 | 1.85 | 0.58 |
| 1:lA:688:A:H2' | 1:lA:689:A:H8 | 1.67 | 0.58 |
| 1:lA:862:G:O6 | 1:lA:879:U:O2 | 2.21 | 0.58 |
| 6:lF:61:GLN:OE1 | 38:ll:55:ARG:NH2 | 2.28 | 0.58 |
| 52:sa:1932:U:H2' | 52:sa:1933:G:H8 | 1.68 | 0.58 |
| 70:su:25:ARG:NH1 | 70:su:29:TYR:O | 2.37 | 0.58 |
| 1:lA:3038:C:O2' | 42:lp:25:TYR:O | 2.15 | 0.58 |
| 1:lA:3474:U:H3 | 1:lA:3480:A:H62 | 1.49 | 0.58 |
| 52:sa:802:U:N3 | 52:sa:803:G:O6 | 2.36 | 0.58 |
| 52:sa:1350:A:H2' | 52:sa:1351:A:C8 | 2.37 | 0.58 |
| 63:sm:34:HIS:NE2 | 63:sm:47:ALA:O | 2.35 | 0.58 |
| 71:sv:57:ARG:NH1 | 71:sv:101:ASN:OD1 | 2.37 | 0.58 |
| 1:lA:232:G:N1 | 1:lA:274:U:OP2 | 2.28 | 0.58 |
| 1:lA:779:A:H5' | 1:lA:2448:A:H62 | 1.68 | 0.58 |
| 21:lU:89:THR:O | 21:lU:89:THR:OG1 | 2.22 | 0.58 |
| 52:sa:160:G:H2' | 52:sa:161:A:O4' | 2.03 | 0.58 |
| 1:lA:1293:U:O4 | 1:lA:1454:G:O2' | 2.19 | 0.58 |
| 1:lA:2225:A:O2' | 4:lD:179:ILE:O | 2.15 | 0.58 |
| 1:lA:2366:C:H1' | 52:sa:1892:G:H21 | 1.68 | 0.58 |
| 52:sa:1440:A:N6 | 52:sa:1519:G:O2' | 2.37 | 0.58 |
| 1:lA:2628:A:N7 | 4:lD:100:ASN:ND2 | 2.52 | 0.58 |
| 1:lA:3170:A:H2' | 1:lA:3171:A:H8 | 1.68 | 0.58 |
| 1:lA:3315:U:O2' | 1:lA:3418:C:OP2 | 2.19 | 0.58 |
| 3:lC:54:U:H4' | 13:lM:152:HIS:HB2 | 1.86 | 0.58 |
| 28:lb:91:GLN:NE2 | 28:lb:124:GLU:OE2 | 2.37 | 0.58 |
| 37:lk:29:LYS:O | 37:lk:32:HIS:ND1 | 2.37 | 0.58 |
| 52:sa:1581:G:H2' | 52:sa:1582:G:C8 | 2.37 | 0.58 |
| 52:sa:1886:U:O4 | 52:sa:1887:A:N6 | 2.37 | 0.58 |
| 1:lA:1314:A:N7 | 11:lK:73:ARG:NH2 | 2.51 | 0.58 |
| 1:lA:2702:G:O6 | 1:lA:2717:A:N6 | 2.37 | 0.58 |
| 9:ll:203:LYS:HB2 | 9:ll:207:ILE:HD11 | 1.86 | 0.58 |
| 12:lL:190:ILE:HG23 | 12:lL:197:PHE:HB2 | 1.85 | 0.58 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 49:sI:22:G:H2' | 49:sI:23:A:C8 | 2.38 | 0.58 |
| 52:sa:1007:A:OP1 | 52:sa:1936:G:O2' | 2.22 | 0.58 |
| 52:sa:1126:G:H1' | 67:sr:76:SER:HB3 | 1.86 | 0.58 |
| 54:sd:22:LYS:HG2 | 72:sw:61:LEU:HD11 | 1.86 | 0.58 |
| 1:lA:2597:U:H3 | 4:lD:67:ARG:HD2 | 1.68 | 0.58 |
| 2:lB:5:A:O2' | 18:lR:61:ARG:NH1 | 2.36 | 0.58 |
| 4:lD:33:ASP:N | 4:lD:33:ASP:OD1 | 2.34 | 0.58 |
| 4:lD:39:SER:OG | 4:lD:40:THR:N | 2.34 | 0.58 |
| 11:lK:19:ILE:HD11 | 11:lK:89:ILE:HD11 | 1.85 | 0.58 |
| 49:sI:68:G:H2' | 49:sI:69:G:C8 | 2.38 | 0.58 |
| 1:lA:3107:G:N2 | 1:lA:3110:A:OP2 | 2.36 | 0.58 |
| 51:sK:8:A:O2' | 74:sy:59:GLN:NE2 | 2.35 | 0.58 |
| 52:sa:1544:U:H2' | 52:sa:1545:G:H8 | 1.69 | 0.58 |
| 57:sg:102:ILE:O | 57:sg:106:THR:OG1 | 2.19 | 0.58 |
| 60:sj:56:ARG:NH1 | 60:sj:210:GLY:O | 2.34 | 0.58 |
| 1:lA:772:C:N4 | 1:lA:773:G:O6 | 2.37 | 0.58 |
| 52:sa:1172:G:H2' | 52:sa:1173:A:C8 | 2.38 | 0.58 |
| 52:sa:1341:C:HO2' | 52:sa:1508:A:HO2' | 1.51 | 0.58 |
| 71:sv:8:LYS:HE3 | 71:sv:67:LEU:HD22 | 1.86 | 0.58 |
| 1:lA:838:U:O2' | 1:lA:907:U:OP1 | 2.21 | 0.57 |
| 1:lA:3091:C:OP1 | 5:lE:246:ARG:NH1 | 2.32 | 0.57 |
| 1:lA:3154:A:H2' | 1:lA:3155:U:C6 | 2.38 | 0.57 |
| 28:lb:33:THR:OG1 | 28:lb:35:ASP:OD1 | 2.23 | 0.57 |
| 60:sj:67:TRP:NE1 | 60:sj:69:SER:HB3 | 2.19 | 0.57 |
| 1:lA:764:U:H2' | 1:lA:765:A:H8 | 1.69 | 0.57 |
| 1:lA:2297:G:N1 | 1:lA:2300:A:OP2 | 2.33 | 0.57 |
| 1:lA:3350:U:OP2 | 1:lA:3351:U:O2' | 2.16 | 0.57 |
| 7:lG:125:ALA:O | 7:lG:192:ARG:NH1 | 2.37 | 0.57 |
| 15:lO:47:THR:H | 15:lO:50:MET:HE3 | 1.69 | 0.57 |
| 22:lV:68:ASN:OD1 | 22:lV:69:PRO:HD2 | 2.02 | 0.57 |
| 31:le:40:ASP:N | 31:le:40:ASP:OD1 | 2.35 | 0.57 |
| 68:ss:64:PHE:HB3 | 68:ss:68:PHE:HE2 | 1.69 | 0.57 |
| 73:sx:23:ALA:HB3 | 73:sx:26:ASP:HB2 | 1.86 | 0.57 |
| 1:lA:547:A:H2' | 1:lA:548:A:H8 | 1.68 | 0.57 |
| 7:lG:85:ARG:NH2 | 7:lG:249:ALA:O | 2.37 | 0.57 |
| 23:lW:70:VAL:HG12 | 23:lW:72:ASP:H | 1.70 | 0.57 |
| 26:lZ:8:CYS:SG | 26:lZ:11:SER:OG | 2.62 | 0.57 |
| 52:sa:1396:A:H2' | 52:sa:1425:A:C8 | 2.39 | 0.57 |
| 1:lA:311:A:H5' | 37:lk:30:ARG:HB2 | 1.87 | 0.57 |
| 1:lA:1639:U:H2' | 1:lA:1640:A:H8 | 1.70 | 0.57 |
| 1:lA:2590:A:N1 | 1:lA:2664:A:N6 | 2.52 | 0.57 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 1:1A:2705:A:O2' | 22:1V:10:ARG:NH2 | 2.36 | 0.57 |
| 4:1D:242:ARG:HH21 | 4:1D:246:ILE:HA | 1.68 | 0.57 |
| 52:sa:15:U:H2' | 52:sa:16:G:O4' | 2.05 | 0.57 |
| 52:sa:841:U:O2' | 52:sa:842:A:O4' | 2.21 | 0.57 |
| 68:ss:65:GLU:OE1 | 68:ss:97:ARG:NH2 | 2.37 | 0.57 |
| 1:1A:844:G:N2 | 1:1A:847:A:OP2 | 2.36 | 0.57 |
| 1:1A:2815:G:N2 | 1:1A:2818:A:OP2 | 2.34 | 0.57 |
| 1:1A:2864:A:O2' | 1:1A:2865:A:O5' | 2.21 | 0.57 |
| 52:sa:1734:G:H4' | 52:sa:1735:A:O5' | 2.04 | 0.57 |
| 52:sa:1891:A:O3' | 52:sa:1892:G:H8 | 1.86 | 0.57 |
| 55:se:66:ARG:NH1 | 65:sp:46:GLU:OE2 | 2.38 | 0.57 |
| 1:1A:361:C:OP1 | 14:1N:100:LYS:NZ | 2.36 | 0.57 |
| 1:1A:2668:G:H2' | 1:1A:2669:A:H8 | 1.68 | 0.57 |
| 1:1A:3359:U:OP1 | 8:1H:194:LYS:NZ | 2.36 | 0.57 |
| 3:1C:42:U:OP1 | 13:1M:137:ARG:NH1 | 2.38 | 0.57 |
| 16:1P:69:ARG:NH2 | 20:1T:127:ASP:OD2 | 2.35 | 0.57 |
| 52:sa:615:A:O2' | 52:sa:1132:C:O2' | 2.22 | 0.57 |
| 1:1A:13:C:P | 25:1Y:27:ARG:HH22 | 2.28 | 0.57 |
| 1:1A:1555:U:O2 | 33:lg:78:ARG:NH2 | 2.34 | 0.57 |
| 1:1A:2865:A:O2' | 1:1A:2866:C:OP1 | 2.20 | 0.57 |
| 52:sa:335:U:H2' | 52:sa:336:A:C8 | 2.39 | 0.57 |
| 52:sa:1141:U:H2' | 52:sa:1142:A:H8 | 1.68 | 0.57 |
| 52:sa:1182:C:H2' | 52:sa:1183:A:H8 | 1.70 | 0.57 |
| 52:sa:1356:A:OP1 | 69:st:45:ARG:NH2 | 2.37 | 0.57 |
| 52:sa:1595:G:O2' | 52:sa:1661:C:O2 | 2.22 | 0.57 |
| 52:sa:1661:C:H2' | 52:sa:1662:G:C8 | 2.39 | 0.57 |
| 56:sf:181:MET:HE3 | 56:sf:208:ILE:HD13 | 1.86 | 0.57 |
| 67:sr:23:ARG:NH2 | 73:sx:17:VAL:O | 2.36 | 0.57 |
| 1:1A:815:A:H4' | 1:1A:817:A:OP1 | 2.03 | 0.57 |
| 1:1A:1236:G:H2' | 1:1A:1237:A:H8 | 1.69 | 0.57 |
| 1:1A:1269:U:OP2 | 33:lg:44:ARG:NH2 | 2.31 | 0.57 |
| 1:1A:2991:G:OP1 | 42:lp:25:TYR:OH | 2.23 | 0.57 |
| 1:1A:3023:U:O2 | 5:1E:252:ALA:HB3 | 2.04 | 0.57 |
| 52:sa:782:A:C8 | 52:sa:784:U:H1' | 2.40 | 0.57 |
| 54:sd:122:LYS:NZ | 54:sd:135:SER:OG | 2.38 | 0.57 |
| 57:sg:177:ASP:OD1 | 57:sg:178:GLU:N | 2.37 | 0.57 |
| 1:1A:315:A:N7 | 17:1Q:12:ARG:NH1 | 2.53 | 0.57 |
| 9:1I:214:GLY:O | 9:1I:218:ARG:NH2 | 2.30 | 0.57 |
| 31:1e:15:ASN:OD1 | 31:1e:74:TYR:OH | 2.22 | 0.57 |
| 52:sa:1308:U:OP2 | 52:sa:1309:C:O2' | 2.19 | 0.57 |
| 52:sa:1748:U:N3 | 52:sa:1780:A:N7 | 2.53 | 0.57 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 1:lA:1206:A:H2' | 1:lA:1207:C:C6 | 2.40 | 0.57 |
| 1:lA:3397:C:OP2 | 1:lA:3398:U:O2' | 2.23 | 0.57 |
| 52:sa:1556:U:H2' | 52:sa:1557:C:C6 | 2.40 | 0.57 |
| 65:sp:35:THR:O | 65:sp:53:GLY:N | 2.35 | 0.57 |
| 69:st:25:THR:HG23 | 69:st:27:ASP:H | 1.70 | 0.57 |
| 69:st:41:VAL:HG22 | 69:st:43:THR:H | 1.70 | 0.57 |
| 1:lA:502:U:H2' | 1:lA:503:A:H8 | 1.70 | 0.56 |
| 1:lA:2175:C:H2' | 1:lA:2176:G:C8 | 2.38 | 0.56 |
| 57:sg:25:SER:OG | 57:sg:27:ASP:OD1 | 2.23 | 0.56 |
| 57:sg:63:PHE:CE1 | 57:sg:146:LEU:HD22 | 2.40 | 0.56 |
| 60:sj:87:ASP:OD1 | 60:sj:88:ASN:N | 2.37 | 0.56 |
| 67:sr:102:LEU:N | 67:sr:113:ASN:OD1 | 2.35 | 0.56 |
| 5:lE:10:ARG:HH12 | 5:lE:265:THR:HB | 1.70 | 0.56 |
| 1:lA:542:U:H2' | 1:lA:543:A:C8 | 2.39 | 0.56 |
| 1:lA:1031:G:N1 | 4:lD:208:ASP:OD2 | 2.33 | 0.56 |
| 1:lA:1335:U:O2 | 20:lT:109:GLN:NE2 | 2.37 | 0.56 |
| 1:lA:3405:A:N6 | 8:lH:164:GLU:OE1 | 2.38 | 0.56 |
| 1:lA:3500:A:H2' | 1:lA:3501:A:C8 | 2.39 | 0.56 |
| 8:lH:65:LEU:HD11 | 8:lH:79:PHE:HB2 | 1.86 | 0.56 |
| 16:lP:33:HIS:ND1 | 16:lP:33:HIS:O | 2.39 | 0.56 |
| 44:sA:100:VAL:N | 44:sA:108:ILE:O | 2.38 | 0.56 |
| 52:sa:442:U:O2' | 56:sf:25:PHE:O | 2.23 | 0.56 |
| 52:sa:1722:A:OP1 | 66:sq:114:TYR:OH | 2.19 | 0.56 |
| 70:su:97:SER:OG | 70:su:98:GLN:N | 2.37 | 0.56 |
| 1:lA:58:G:H21 | 17:lQ:186:ALA:HB1 | 1.71 | 0.56 |
| 1:lA:1128:G:OP1 | 12:lL:39:ARG:NH2 | 2.38 | 0.56 |
| 1:lA:1186:G:H21 | 1:lA:1220:A:N6 | 2.03 | 0.56 |
| 1:lA:2468:C:O2' | 5:lE:268:ARG:NH1 | 2.39 | 0.56 |
| 13:lM:63:ASP:OD1 | 13:lM:63:ASP:N | 2.38 | 0.56 |
| 16:lP:108:THR:OG1 | 16:lP:111:ASP:OD2 | 2.22 | 0.56 |
| 45:sB:5:ARG:NH1 | 52:sa:1943:C:OP2 | 2.37 | 0.56 |
| 52:sa:866:G:H2' | 52:sa:867:A:H8 | 1.70 | 0.56 |
| 57:sg:38:LEU:HD12 | 57:sg:145:PRO:HB2 | 1.87 | 0.56 |
| 1:lA:701:A:H5'' | 1:lA:703:G:H21 | 1.69 | 0.56 |
| 10:lJ:80:GLN:HA | 10:lJ:83:LYS:HE2 | 1.87 | 0.56 |
| 52:sa:1221:A:OP2 | 52:sa:1571:G:N2 | 2.24 | 0.56 |
| 61:sk:31:VAL:HA | 61:sk:36:LEU:HD12 | 1.88 | 0.56 |
| 68:ss:75:TYR:OH | 68:ss:100:ILE:O | 2.21 | 0.56 |
| 71:sv:14:ASP:OD1 | 71:sv:15:PHE:N | 2.38 | 0.56 |
| 1:lA:789:C:H2' | 1:lA:790:A:H8 | 1.69 | 0.56 |
| 3:lC:21:C:N4 | 3:lC:52:G:O2' | 2.39 | 0.56 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 37:lk:39:ARG:NH1 | 37:lk:43:GLY:O | 2.35 | 0.56 |
| 45:sB:22:ARG:NH1 | 65:sp:140:GLY:O | 2.38 | 0.56 |
| 49:sI:8:U:H3 | 49:sI:15:G:H1 | 1.52 | 0.56 |
| 52:sa:1748:U:O2 | 52:sa:1780:A:N6 | 2.34 | 0.56 |
| 1:lA:1252:G:N2 | 1:lA:1255:A:OP2 | 2.38 | 0.56 |
| 1:lA:2711:U:OP2 | 22:IV:9:ARG:NH2 | 2.34 | 0.56 |
| 52:sa:1002:U:O2' | 52:sa:1151:A:N1 | 2.38 | 0.56 |
| 56:sf:69:LYS:HG2 | 56:sf:74:ILE:HB | 1.86 | 0.56 |
| 62:sl:25:LYS:HG2 | 62:sl:61:ARG:HH21 | 1.69 | 0.56 |
| 1:lA:1555:U:H2' | 1:lA:1556:G:C8 | 2.39 | 0.56 |
| 52:sa:440:A:H5'' | 56:sf:57:ARG:HH12 | 1.68 | 0.56 |
| 52:sa:1561:G:C6 | 52:sa:1562:G:N2 | 2.74 | 0.56 |
| 54:sd:133:VAL:HG21 | 54:sd:152:ILE:HD11 | 1.88 | 0.56 |
| 1:lA:871:A:O2' | 1:lA:872:A:O5' | 2.23 | 0.56 |
| 1:lA:2890:A:H2' | 1:lA:2891:G:C8 | 2.41 | 0.56 |
| 14:lN:264:ARG:O | 14:lN:268:ASN:ND2 | 2.39 | 0.56 |
| 46:sC:18:LYS:O | 52:sa:1095:U:O2' | 2.23 | 0.56 |
| 48:sE:22:ARG:NH1 | 48:sE:36:LEU:O | 2.39 | 0.56 |
| 52:sa:919:A:OP2 | 64:so:6:ASN:ND2 | 2.39 | 0.56 |
| 52:sa:1820:G:H5'' | 52:sa:1821:A:H5' | 1.86 | 0.56 |
| 52:sa:111:A:O2' | 63:sm:66:ARG:NH1 | 2.39 | 0.56 |
| 55:se:201:ALA:O | 55:se:203:CYS:N | 2.39 | 0.56 |
| 56:sf:62:ILE:HG22 | 56:sf:68:ILE:HD11 | 1.87 | 0.56 |
| 1:lA:50:G:HO2' | 1:lA:1688:A:HO2' | 1.54 | 0.55 |
| 1:lA:760:A:H2' | 1:lA:761:A:H8 | 1.70 | 0.55 |
| 1:lA:993:U:OP2 | 1:lA:2109:C:O2' | 2.23 | 0.55 |
| 1:lA:1498:C:O3' | 19:IS:26:LYS:NZ | 2.39 | 0.55 |
| 1:lA:1622:A:C2 | 34:lh:4:ARG:HD3 | 2.41 | 0.55 |
| 1:lA:2795:C:N4 | 1:lA:2802:A:N1 | 2.54 | 0.55 |
| 1:lA:3267:G:O6 | 1:lA:3275:C:H5'' | 2.06 | 0.55 |
| 4:ID:116:VAL:HG13 | 4:ID:164:ALA:HB2 | 1.88 | 0.55 |
| 16:lP:29:GLU:HG2 | 20:lT:68:VAL:HG21 | 1.88 | 0.55 |
| 52:sa:1661:C:H2' | 52:sa:1662:G:H8 | 1.71 | 0.55 |
| 52:sa:1776:G:H5'' | 57:sg:88:LYS:HD2 | 1.88 | 0.55 |
| 56:sf:67:THR:HG23 | 56:sf:92:ARG:HG3 | 1.87 | 0.55 |
| 1:lA:405:U:OP1 | 38:ll:11:ARG:NH2 | 2.37 | 0.55 |
| 1:lA:1649:U:HO2' | 1:lA:2429:G:HO2' | 1.53 | 0.55 |
| 1:lA:1937:A:OP1 | 40:ln:35:LYS:NZ | 2.33 | 0.55 |
| 11:lK:112:VAL:HB | 11:lK:121:GLN:HB3 | 1.87 | 0.55 |
| 32:lf:61:GLU:HB3 | 32:lf:66:ARG:HE | 1.71 | 0.55 |
| 52:sa:1340:U:OP1 | 52:sa:1353:G:N2 | 2.33 | 0.55 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|-------------------|--------------------------|-------------------|
| 58:sh:78:ASP:OD1 | 58:sh:78:ASP:N | 2.38 | 0.55 |
| 62:sl:30:LEU:HA | 62:sl:37:PRO:HA | 1.88 | 0.55 |
| 1:lA:336:A:H5'' | 17:lQ:97:SER:HB3 | 1.88 | 0.55 |
| 1:lA:672:G:O6 | 6:lF:412:LYS:NZ | 2.39 | 0.55 |
| 1:lA:2425:U:O2' | 1:lA:3436:A:N3 | 2.40 | 0.55 |
| 1:lA:3476:C:H4' | 1:lA:3476:C:OP1 | 2.05 | 0.55 |
| 2:lB:110:A:N6 | 41:lo:51:MET:O | 2.39 | 0.55 |
| 4:lD:192:LYS:HD3 | 4:lD:193:ARG:HH21 | 1.71 | 0.55 |
| 9:lI:176:ILE:HD13 | 9:lI:183:PHE:HE1 | 1.71 | 0.55 |
| 52:sa:1827:U:H2' | 52:sa:1828:A:H8 | 1.71 | 0.55 |
| 54:sd:211:LYS:NZ | 69:st:40:VAL:O | 2.38 | 0.55 |
| 68:ss:35:CYS:HB2 | 68:ss:82:ILE:HG12 | 1.88 | 0.55 |
| 1:lA:820:A:OP1 | 14:lN:37:ARG:NH2 | 2.37 | 0.55 |
| 6:lF:389:LYS:C | 6:lF:391:LYS:H | 2.14 | 0.55 |
| 65:sp:18:SER:OG | 65:sp:19:THR:N | 2.38 | 0.55 |
| 1:lA:211:U:OP1 | 14:lN:134:ARG:NH1 | 2.40 | 0.55 |
| 1:lA:1648:U:OP2 | 1:lA:2081:A:O2' | 2.25 | 0.55 |
| 8:lH:176:GLU:HA | 8:lH:179:LYS:HE2 | 1.88 | 0.55 |
| 1:lA:26:C:OP1 | 17:lQ:188:ALA:N | 2.37 | 0.55 |
| 1:lA:1098:A:HO2' | 1:lA:1183:G:HO2' | 1.55 | 0.55 |
| 1:lA:2466:G:N2 | 5:lE:228:TYR:OH | 2.33 | 0.55 |
| 1:lA:2469:G:OP1 | 1:lA:2469:G:C8 | 2.60 | 0.55 |
| 6:lF:11:ASP:OD1 | 27:la:146:TYR:OH | 2.23 | 0.55 |
| 50:sJ:16:C:H2' | 50:sJ:61:U:H3 | 1.72 | 0.55 |
| 52:sa:366:G:N2 | 52:sa:606:U:O2 | 2.40 | 0.55 |
| 52:sa:397:U:OP1 | 56:sf:1:ARG:NH2 | 2.39 | 0.55 |
| 52:sa:932:C:H2' | 52:sa:933:A:C8 | 2.41 | 0.55 |
| 52:sa:1115:C:N4 | 52:sa:1116:U:O4 | 2.39 | 0.55 |
| 52:sa:1754:U:O4 | 52:sa:1755:A:N6 | 2.40 | 0.55 |
| 52:sa:1923:A:H2' | 52:sa:1924:G:C8 | 2.41 | 0.55 |
| 67:sr:45:GLN:O | 67:sr:68:ARG:NH1 | 2.40 | 0.55 |
| 70:su:70:ASP:OD1 | 70:su:70:ASP:N | 2.37 | 0.55 |
| 29:lc:4:ARG:HA | 29:lc:9:ARG:HG3 | 1.87 | 0.55 |
| 56:sf:127:ARG:HH11 | 56:sf:135:PRO:HB2 | 1.71 | 0.55 |
| 64:so:92:MET:O | 64:so:94:ARG:N | 2.39 | 0.55 |
| 1:lA:1320:U:O2' | 3:lC:83:G:N2 | 2.40 | 0.55 |
| 1:lA:2416:U:OP2 | 5:lE:239:HIS:ND1 | 2.40 | 0.55 |
| 1:lA:2905:U:H5'' | 14:lN:265:LYS:HG2 | 1.88 | 0.55 |
| 43:lq:72:CYS:HB3 | 43:lq:77:TYR:H | 1.72 | 0.55 |
| 45:sB:6:ARG:HB3 | 52:sa:1943:C:H5 | 1.72 | 0.55 |
| 52:sa:1284:U:O4 | 52:sa:1285:A:N6 | 2.40 | 0.55 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:1A:729:U:O4 | 1:1A:730:A:N6 | 2.40 | 0.55 |
| 1:1A:1610:A:O2' | 1:1A:1651:C:O2 | 2.25 | 0.55 |
| 1:1A:1744:A:H2' | 1:1A:1745:A:H8 | 1.72 | 0.55 |
| 1:1A:1763:C:H2' | 1:1A:1764:A:H8 | 1.72 | 0.55 |
| 1:1A:3214:C:O3' | 1:1A:3493:G:N2 | 2.40 | 0.55 |
| 11:1K:93:THR:HG22 | 11:1K:94:LYS:HG3 | 1.88 | 0.55 |
| 43:lq:73:THR:O | 43:lq:76:LYS:NZ | 2.33 | 0.55 |
| 56:sf:239:ASP:OD1 | 56:sf:239:ASP:N | 2.37 | 0.55 |
| 1:1A:591:C:H2' | 1:1A:592:A:H8 | 1.72 | 0.55 |
| 1:1A:3008:U:O2' | 1:1A:3009:C:OP1 | 2.22 | 0.55 |
| 13:1M:95:ASN:ND2 | 13:1M:103:GLY:O | 2.40 | 0.55 |
| 52:sa:1196:A:O2' | 52:sa:1736:A:N3 | 2.37 | 0.55 |
| 52:sa:1371:U:C4 | 52:sa:1373:A:N7 | 2.75 | 0.55 |
| 1:1A:1071:A:H4' | 1:1A:1087:G:N2 | 2.22 | 0.54 |
| 18:1R:53:ILE:HG13 | 18:1R:55:LYS:HG2 | 1.88 | 0.54 |
| 1:1A:446:A:H5' | 1:1A:447:G:H2' | 1.89 | 0.54 |
| 1:1A:1251:G:N2 | 12:1L:116:ARG:HH22 | 2.04 | 0.54 |
| 1:1A:2848:A:N1 | 1:1A:2911:U:H5 | 2.05 | 0.54 |
| 5:1E:84:MET:HE1 | 5:1E:183:ILE:HD11 | 1.89 | 0.54 |
| 25:1Y:78:VAL:HG11 | 25:1Y:103:VAL:HG11 | 1.90 | 0.54 |
| 52:sa:1097:C:H4' | 64:so:11:ILE:HD11 | 1.89 | 0.54 |
| 52:sa:1253:G:O2' | 52:sa:1254:G:OP1 | 2.23 | 0.54 |
| 61:sk:59:MET:HA | 61:sk:62:LYS:HE3 | 1.89 | 0.54 |
| 1:1A:106:G:OP2 | 14:1N:71:ARG:NH2 | 2.41 | 0.54 |
| 47:sD:60:GLU:OE1 | 57:sg:206:ARG:NH1 | 2.40 | 0.54 |
| 52:sa:887:A:H2' | 52:sa:888:A:C8 | 2.38 | 0.54 |
| 54:sd:115:GLN:HE21 | 54:sd:136:VAL:HG13 | 1.71 | 0.54 |
| 1:1A:2037:C:O2' | 1:1A:2043:A:N1 | 2.38 | 0.54 |
| 1:1A:3302:U:H2' | 1:1A:3303:A:H8 | 1.72 | 0.54 |
| 10:1J:72:THR:HG21 | 10:1J:178:LYS:HG3 | 1.88 | 0.54 |
| 14:1N:77:GLU:HG3 | 14:1N:107:LEU:HD12 | 1.89 | 0.54 |
| 27:la:75:LYS:HB2 | 27:la:77:VAL:HG22 | 1.88 | 0.54 |
| 52:sa:346:U:O4 | 63:sm:101:LYS:NZ | 2.29 | 0.54 |
| 52:sa:941:C:OP2 | 64:so:14:ARG:NH1 | 2.39 | 0.54 |
| 52:sa:1533:C:H3' | 52:sa:1534:A:H4' | 1.88 | 0.54 |
| 65:sp:73:ALA:HB1 | 65:sp:114:LEU:HG | 1.89 | 0.54 |
| 1:1A:387:G:OP2 | 6:1F:194:GLN:NE2 | 2.29 | 0.54 |
| 1:1A:505:A:H2' | 1:1A:506:A:C8 | 2.43 | 0.54 |
| 1:1A:2233:G:N7 | 4:1D:152:SER:OG | 2.37 | 0.54 |
| 1:1A:3043:A:N3 | 1:1A:3179:C:O2' | 2.35 | 0.54 |
| 3:1C:6:G:OP2 | 7:1G:28:THR:OG1 | 2.24 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 7:IG:50:ARG:NH1 | 7:IG:72:ASP:OD2 | 2.40 | 0.54 |
| 13:IM:116:TYR:O | 70:su:102:ASN:ND2 | 2.41 | 0.54 |
| 27:la:47:ILE:HD11 | 27:la:107:LEU:HD21 | 1.88 | 0.54 |
| 31:le:14:VAL:HA | 31:le:17:LYS:HD2 | 1.89 | 0.54 |
| 43:lq:63:THR:O | 43:lq:85:ARG:NH1 | 2.36 | 0.54 |
| 52:sa:151:A:H2' | 52:sa:152:A:O4' | 2.07 | 0.54 |
| 52:sa:1880:U:H2' | 52:sa:1881:C:H6 | 1.73 | 0.54 |
| 54:sd:188:CYS:HB2 | 54:sd:195:ILE:HD11 | 1.89 | 0.54 |
| 1:lA:1554:C:OP2 | 6:lF:196:LYS:NZ | 2.41 | 0.54 |
| 3:lC:27:G:H5'' | 13:lM:137:ARG:HD3 | 1.89 | 0.54 |
| 6:lF:85:GLY:O | 6:lF:89:GLN:NE2 | 2.40 | 0.54 |
| 13:lM:94:ASN:ND2 | 13:lM:94:ASN:O | 2.41 | 0.54 |
| 19:lS:36:ALA:HB1 | 19:lS:45:LYS:HG2 | 1.88 | 0.54 |
| 34:lh:15:THR:HG22 | 34:lh:16:LYS:H | 1.73 | 0.54 |
| 50:sJ:24:C:H2' | 50:sJ:25:U:C6 | 2.42 | 0.54 |
| 52:sa:1828:A:H2' | 52:sa:1829:A:H8 | 1.72 | 0.54 |
| 54:sd:141:MET:HE2 | 54:sd:141:MET:HA | 1.90 | 0.54 |
| 1:lA:189:G:N7 | 10:lJ:134:ASN:HB2 | 2.23 | 0.54 |
| 1:lA:662:U:H2' | 1:lA:663:G:O4' | 2.08 | 0.54 |
| 1:lA:1639:U:H2' | 1:lA:1640:A:C8 | 2.43 | 0.54 |
| 2:lB:95:A:OP1 | 35:li:59:ARG:NH2 | 2.41 | 0.54 |
| 8:lH:113:THR:HG22 | 8:lH:204:PHE:HB3 | 1.88 | 0.54 |
| 10:lJ:168:LYS:NZ | 37:lk:40:GLU:OE2 | 2.38 | 0.54 |
| 50:sJ:4:U:H2' | 50:sJ:5:U:O4' | 2.07 | 0.54 |
| 52:sa:752:A:P | 61:sk:51:HIS:HE2 | 2.30 | 0.54 |
| 52:sa:1838:A:H2' | 52:sa:1839:A:C8 | 2.43 | 0.54 |
| 56:sf:34:HIS:HB2 | 56:sf:39:CYS:SG | 2.48 | 0.54 |
| 63:sm:108:VAL:HG12 | 63:sm:137:PHE:HB2 | 1.90 | 0.54 |
| 1:lA:1553:A:OP1 | 2:lB:22:G:O2' | 2.26 | 0.54 |
| 1:lA:3432:A:O2' | 1:lA:3433:U:OP1 | 2.26 | 0.54 |
| 26:lZ:49:ARG:HG2 | 26:lZ:60:LEU:HD11 | 1.89 | 0.54 |
| 52:sa:967:G:N2 | 52:sa:993:A:OP1 | 2.41 | 0.54 |
| 54:sd:204:LEU:HD13 | 54:sd:208:PRO:HG2 | 1.90 | 0.54 |
| 14:lN:214:THR:OG1 | 14:lN:215:LYS:N | 2.39 | 0.54 |
| 17:lQ:80:VAL:HB | 17:lQ:82:ARG:HD2 | 1.90 | 0.54 |
| 32:lf:84:THR:OG1 | 32:lf:127:THR:OG1 | 2.26 | 0.54 |
| 52:sa:401:U:H2' | 52:sa:402:A:H8 | 1.73 | 0.54 |
| 1:lA:3196:U:H2' | 1:lA:3197:C:H6 | 1.73 | 0.54 |
| 9:II:66:GLU:OE2 | 9:II:66:GLU:N | 2.38 | 0.54 |
| 15:lO:160:GLU:O | 15:lO:164:ARG:NH1 | 2.41 | 0.54 |
| 23:lW:33:ASP:HB3 | 23:lW:117:GLU:HA | 1.90 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 27:la:22:ASN:OD1 | 27:la:22:ASN:N | 2.40 | 0.54 |
| 52:sa:599:A:OP2 | 52:sa:600:A:O2' | 2.25 | 0.54 |
| 52:sa:1573:G:O2' | 52:sa:1768:C:OP1 | 2.26 | 0.54 |
| 56:sf:123:LYS:HB2 | 56:sf:224:PHE:CE1 | 2.42 | 0.54 |
| 60:sj:10:LYS:HZ3 | 63:sm:133:LYS:HD2 | 1.73 | 0.54 |
| 74:sy:22:ASP:OD2 | 74:sy:25:TRP:HB3 | 2.08 | 0.54 |
| 1:lA:968:C:H2' | 1:lA:969:U:C6 | 2.44 | 0.53 |
| 1:lA:1474:A:OP2 | 19:lS:30:ARG:NH2 | 2.40 | 0.53 |
| 1:lA:3285:A:C6 | 1:lA:3287:U:H1' | 2.43 | 0.53 |
| 1:lA:3443:C:HO2' | 1:lA:3447:A:HO2' | 1.51 | 0.53 |
| 1:lA:3500:A:H2' | 1:lA:3501:A:H8 | 1.73 | 0.53 |
| 17:lQ:114:ARG:HD3 | 17:lQ:151:ILE:HG12 | 1.91 | 0.53 |
| 27:la:47:ILE:HG22 | 27:la:120:LYS:HE3 | 1.89 | 0.53 |
| 50:sJ:62:C:H2' | 50:sJ:63:C:C6 | 2.43 | 0.53 |
| 52:sa:144:C:H2' | 52:sa:145:U:O4' | 2.08 | 0.53 |
| 52:sa:585:A:H2' | 52:sa:586:A:C8 | 2.44 | 0.53 |
| 52:sa:1196:A:H2' | 52:sa:1197:G:C8 | 2.43 | 0.53 |
| 1:lA:3476:C:H5'' | 1:lA:3477:U:C2 | 2.44 | 0.53 |
| 6:lF:63:SER:O | 6:lF:63:SER:OG | 2.24 | 0.53 |
| 14:lN:74:THR:OG1 | 14:lN:77:GLU:OE1 | 2.24 | 0.53 |
| 58:sh:89:ASN:N | 58:sh:89:ASN:OD1 | 2.42 | 0.53 |
| 63:sm:132:SER:OG | 63:sm:133:LYS:N | 2.38 | 0.53 |
| 67:sr:11:LEU:HD22 | 67:sr:72:CYS:HB2 | 1.90 | 0.53 |
| 67:sr:102:LEU:HB2 | 67:sr:113:ASN:HB3 | 1.90 | 0.53 |
| 1:lA:87:G:N2 | 1:lA:91:A:OP2 | 2.39 | 0.53 |
| 1:lA:572:U:H2' | 1:lA:573:A:C8 | 2.43 | 0.53 |
| 1:lA:724:A:O2' | 1:lA:725:A:H5'' | 2.09 | 0.53 |
| 1:lA:1626:G:N2 | 34:lh:4:ARG:HD2 | 2.24 | 0.53 |
| 1:lA:2884:U:H3' | 1:lA:2885:G:N2 | 2.22 | 0.53 |
| 4:lD:19:ARG:NH1 | 4:lD:189:TYR:O | 2.42 | 0.53 |
| 11:lK:13:ILE:HD11 | 11:lK:63:VAL:HG23 | 1.91 | 0.53 |
| 52:sa:247:G:OP1 | 56:sf:132:GLY:N | 2.28 | 0.53 |
| 1:lA:206:U:O4 | 1:lA:303:G:O6 | 2.27 | 0.53 |
| 1:lA:1749:U:OP2 | 21:lU:42:ARG:NH2 | 2.41 | 0.53 |
| 1:lA:1813:G:N7 | 28:lb:17:ARG:NH2 | 2.56 | 0.53 |
| 1:lA:3444:A:H4' | 1:lA:3447:A:H1' | 1.89 | 0.53 |
| 1:lA:3497:U:H4' | 1:lA:3498:U:H5' | 1.91 | 0.53 |
| 3:lC:38:A:H2 | 13:lM:70:THR:HG23 | 1.72 | 0.53 |
| 7:lG:45:LYS:HG3 | 22:lV:33:THR:HG21 | 1.90 | 0.53 |
| 8:lH:135:ILE:HA | 8:lH:159:LYS:HD2 | 1.89 | 0.53 |
| 52:sa:13:C:O2' | 52:sa:1323:U:O4 | 2.23 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 52:sa:1822:U:H5' | 52:sa:1823:U:H5'' | 1.90 | 0.53 |
| 56:sf:105:ARG:NH2 | 56:sf:185:ASN:OD1 | 2.41 | 0.53 |
| 1:lA:3492:G:H5' | 5:lE:385:LYS:HD2 | 1.89 | 0.53 |
| 2:lB:57:U:C2 | 2:lB:58:G:C8 | 2.97 | 0.53 |
| 3:lC:26:U:H3 | 3:lC:51:G:N2 | 2.04 | 0.53 |
| 3:lC:104:G:H2' | 3:lC:105:A:C8 | 2.44 | 0.53 |
| 52:sa:134:A:N1 | 52:sa:172:G:N2 | 2.56 | 0.53 |
| 52:sa:1248:U:H2' | 52:sa:1249:A:H8 | 1.74 | 0.53 |
| 52:sa:1594:A:H2' | 52:sa:1595:G:C8 | 2.43 | 0.53 |
| 56:sf:86:ASP:HA | 56:sf:120:LYS:HG3 | 1.91 | 0.53 |
| 1:lA:220:A:H2' | 1:lA:221:U:H6 | 1.74 | 0.53 |
| 1:lA:829:G:N2 | 6:lF:236:ASN:OD1 | 2.29 | 0.53 |
| 1:lA:836:C:H2' | 1:lA:837:A:H8 | 1.74 | 0.53 |
| 1:lA:1475:A:HO2' | 27:la:185:ARG:HH22 | 1.51 | 0.53 |
| 1:lA:2318:U:H5'' | 4:lD:244:GLY:HA3 | 1.91 | 0.53 |
| 1:lA:2348:G:N2 | 1:lA:2348:G:OP2 | 2.40 | 0.53 |
| 1:lA:2427:U:H2' | 1:lA:2428:A:H8 | 1.73 | 0.53 |
| 5:lE:122:TRP:CE2 | 5:lE:127:LYS:HG2 | 2.43 | 0.53 |
| 7:lG:131:ASP:CG | 7:lG:176:HIS:HE2 | 2.16 | 0.53 |
| 52:sa:540:U:H2' | 52:sa:541:U:C6 | 2.44 | 0.53 |
| 52:sa:586:A:O2' | 52:sa:590:U:OP1 | 2.26 | 0.53 |
| 52:sa:1680:G:O2' | 52:sa:1682:A:N3 | 2.42 | 0.53 |
| 1:lA:897:A:O2' | 1:lA:899:A:N1 | 2.42 | 0.53 |
| 1:lA:2213:U:OP2 | 1:lA:2218:A:N6 | 2.38 | 0.53 |
| 1:lA:2491:C:OP1 | 4:lD:2:GLY:N | 2.41 | 0.53 |
| 1:lA:2499:U:H2' | 1:lA:2500:A:C8 | 2.43 | 0.53 |
| 1:lA:3412:U:H5'' | 1:lA:3413:U:OP2 | 2.09 | 0.53 |
| 12:lL:30:ARG:HH21 | 12:lL:66:GLU:HG2 | 1.73 | 0.53 |
| 1:lA:2225:A:C2 | 1:lA:2264:A:H4' | 2.44 | 0.53 |
| 1:lA:2851:A:H2' | 1:lA:2852:G:H8 | 1.74 | 0.53 |
| 5:lE:307:THR:HG23 | 5:lE:311:GLY:HA2 | 1.90 | 0.53 |
| 52:sa:1396:A:C2 | 52:sa:1425:A:C5 | 2.96 | 0.53 |
| 1:lA:200:A:N6 | 1:lA:309:G:H21 | 2.06 | 0.53 |
| 1:lA:243:G:N2 | 1:lA:418:A:N7 | 2.57 | 0.53 |
| 1:lA:699:A:H5'' | 1:lA:746:A:H62 | 1.74 | 0.53 |
| 1:lA:2878:A:N7 | 37:lk:86:ARG:NH2 | 2.56 | 0.53 |
| 1:lA:3393:A:H2' | 1:lA:3394:A:C8 | 2.43 | 0.53 |
| 23:lW:84:LEU:HD12 | 23:lW:89:MET:HE1 | 1.89 | 0.53 |
| 42:lp:29:PRO:HG2 | 42:lp:32:ALA:HB2 | 1.91 | 0.53 |
| 52:sa:1033:A:H61 | 52:sa:1091:C:N4 | 2.07 | 0.53 |
| 56:sf:201:GLY:HA2 | 63:sm:39:MET:SD | 2.48 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 72:sw:64:THR:HG22 | 72:sw:77:TRP:HE3 | 1.74 | 0.53 |
| 1:lA:440:G:N1 | 1:lA:443:A:OP2 | 2.34 | 0.53 |
| 1:lA:466:G:O2' | 1:lA:2441:U:OP2 | 2.24 | 0.53 |
| 1:lA:721:A:H2' | 1:lA:722:A:H8 | 1.73 | 0.53 |
| 1:lA:2331:A:C8 | 52:sa:1905:U:H1' | 2.44 | 0.53 |
| 1:lA:3142:G:H21 | 1:lA:3143:A:H62 | 1.57 | 0.53 |
| 32:lf:61:GLU:H | 32:lf:66:ARG:HH21 | 1.56 | 0.53 |
| 41:lo:24:PRO:HG2 | 41:lo:27:PHE:HB2 | 1.90 | 0.53 |
| 52:sa:251:C:H2' | 52:sa:252:C:C6 | 2.44 | 0.53 |
| 52:sa:1776:G:OP1 | 57:sg:53:HIS:NE2 | 2.39 | 0.53 |
| 54:sd:115:GLN:NE2 | 54:sd:136:VAL:HG13 | 2.24 | 0.53 |
| 54:sd:175:GLY:O | 54:sd:178:THR:OG1 | 2.25 | 0.53 |
| 1:lA:998:U:H2' | 18:IR:131:ARG:HH21 | 1.74 | 0.52 |
| 1:lA:2101:G:O2' | 1:lA:2410:U:O4 | 2.24 | 0.52 |
| 1:lA:3446:C:O2' | 1:lA:3447:A:H5' | 2.09 | 0.52 |
| 24:IX:107:ASN:HD21 | 24:IX:111:GLU:HB2 | 1.73 | 0.52 |
| 41:lo:9:TYR:HE1 | 41:lo:51:MET:HE2 | 1.73 | 0.52 |
| 52:sa:882:G:OP2 | 65:sp:33:ASN:ND2 | 2.36 | 0.52 |
| 52:sa:888:A:C2 | 52:sa:989:U:H1' | 2.44 | 0.52 |
| 52:sa:1396:A:H2' | 52:sa:1425:A:H8 | 1.72 | 0.52 |
| 52:sa:1699:C:H4' | 52:sa:1705:A:N6 | 2.24 | 0.52 |
| 56:sf:17:MET:HE2 | 56:sf:106:ARG:HD2 | 1.91 | 0.52 |
| 57:sg:30:VAL:HG13 | 57:sg:119:ILE:HD11 | 1.90 | 0.52 |
| 59:si:132:LEU:HD21 | 59:si:141:VAL:HG11 | 1.90 | 0.52 |
| 62:sl:11:ILE:HG13 | 62:sl:34:PHE:HE2 | 1.73 | 0.52 |
| 1:lA:365:G:OP1 | 17:IQ:47:ARG:NH2 | 2.31 | 0.52 |
| 1:lA:2509:U:O4 | 17:IQ:20:ARG:NH1 | 2.42 | 0.52 |
| 20:IT:106:LEU:HD13 | 20:IT:110:TYR:HD2 | 1.74 | 0.52 |
| 29:lc:64:LYS:HB2 | 29:lc:67:LYS:HE3 | 1.90 | 0.52 |
| 37:lk:2:ALA:HB1 | 37:lk:13:TYR:HE1 | 1.74 | 0.52 |
| 52:sa:164:A:O2' | 52:sa:166:A:OP2 | 2.26 | 0.52 |
| 52:sa:846:G:OP1 | 64:so:2:GLY:N | 2.42 | 0.52 |
| 52:sa:872:A:H2' | 52:sa:873:A:H8 | 1.71 | 0.52 |
| 61:sk:33:GLN:HG2 | 61:sk:34:PHE:HD1 | 1.74 | 0.52 |
| 63:sm:62:ASP:OD1 | 63:sm:62:ASP:N | 2.41 | 0.52 |
| 1:lA:228:A:H2' | 1:lA:229:A:C8 | 2.44 | 0.52 |
| 1:lA:412:A:H5'' | 6:IF:97:ARG:HH22 | 1.73 | 0.52 |
| 1:lA:1220:A:H2' | 1:lA:1220:A:N3 | 2.24 | 0.52 |
| 1:lA:1308:A:OP1 | 16:IP:55:ASN:ND2 | 2.37 | 0.52 |
| 1:lA:3088:G:O2' | 1:lA:3091:C:OP2 | 2.25 | 0.52 |
| 9:II:106:LYS:HE2 | 9:II:183:PHE:HD2 | 1.74 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 14:lN:46:PRO:HB3 | 35:li:108:PHE:CZ | 2.44 | 0.52 |
| 46:sC:30:ASN:N | 46:sC:30:ASN:OD1 | 2.42 | 0.52 |
| 52:sa:389:U:H3 | 52:sa:396:A:N6 | 2.02 | 0.52 |
| 52:sa:864:C:H2' | 52:sa:865:A:H8 | 1.72 | 0.52 |
| 70:su:119:ARG:HD2 | 70:su:119:ARG:O | 2.09 | 0.52 |
| 1:lA:191:A:H2' | 1:lA:192:A:H8 | 1.73 | 0.52 |
| 1:lA:500:G:O6 | 1:lA:572:U:O4 | 2.27 | 0.52 |
| 1:lA:1030:C:OP1 | 4:lD:14:SER:OG | 2.22 | 0.52 |
| 3:lC:53:C:HO2' | 13:lM:152:HIS:HD1 | 1.55 | 0.52 |
| 5:lE:386:LYS:N | 5:lE:386:LYS:HD3 | 2.25 | 0.52 |
| 33:lg:96:GLU:HB2 | 33:lg:121:LEU:HD12 | 1.90 | 0.52 |
| 34:lh:65:LEU:O | 34:lh:70:ARG:NH1 | 2.42 | 0.52 |
| 47:sD:51:ARG:HD2 | 57:sg:63:PHE:HD2 | 1.74 | 0.52 |
| 52:sa:169:U:H2' | 52:sa:170:G:C8 | 2.44 | 0.52 |
| 52:sa:465:G:C2 | 52:sa:466:A:C8 | 2.98 | 0.52 |
| 52:sa:746:A:H2' | 52:sa:747:A:C8 | 2.44 | 0.52 |
| 52:sa:1014:C:OP1 | 64:so:9:ARG:NH2 | 2.42 | 0.52 |
| 56:sf:159:LYS:HB3 | 56:sf:169:ASP:H | 1.74 | 0.52 |
| 60:sj:4:THR:HG21 | 60:sj:24:LYS:HG2 | 1.92 | 0.52 |
| 1:lA:1938:G:O6 | 40:ln:35:LYS:NZ | 2.42 | 0.52 |
| 1:lA:3162:A:OP1 | 15:lO:73:HIS:ND1 | 2.41 | 0.52 |
| 23:lW:66:ILE:HG13 | 23:lW:79:THR:HG21 | 1.92 | 0.52 |
| 25:lY:37:ASP:N | 25:lY:37:ASP:OD1 | 2.41 | 0.52 |
| 57:sg:203:LYS:HG3 | 57:sg:206:ARG:HH21 | 1.74 | 0.52 |
| 1:lA:2079:A:H2' | 1:lA:2080:A:O4' | 2.10 | 0.52 |
| 28:lb:45:GLY:O | 28:lb:70:SER:OG | 2.21 | 0.52 |
| 45:sB:4:LYS:NZ | 52:sa:1795:G:OP1 | 2.35 | 0.52 |
| 52:sa:799:C:O2 | 67:sr:124:ARG:NH1 | 2.40 | 0.52 |
| 52:sa:866:G:H2' | 52:sa:867:A:C8 | 2.45 | 0.52 |
| 52:sa:1690:A:N1 | 52:sa:1774:A:H1' | 2.25 | 0.52 |
| 52:sa:1701:U:H4' | 52:sa:1702:G:O5' | 2.09 | 0.52 |
| 57:sg:88:LYS:O | 57:sg:92:MET:HG2 | 2.09 | 0.52 |
| 60:sj:65:PHE:HZ | 60:sj:78:ILE:HG23 | 1.75 | 0.52 |
| 67:sr:77:PRO:HB2 | 74:sy:5:LEU:HB2 | 1.90 | 0.52 |
| 1:lA:378:A:N6 | 1:lA:379:G:O6 | 2.43 | 0.52 |
| 1:lA:597:A:H2' | 1:lA:598:A:H8 | 1.75 | 0.52 |
| 1:lA:720:U:H2' | 1:lA:721:A:H8 | 1.74 | 0.52 |
| 1:lA:1528:A:N3 | 1:lA:1553:A:O2' | 2.41 | 0.52 |
| 1:lA:2752:C:O2' | 13:lM:18:VAL:HG11 | 2.10 | 0.52 |
| 1:lA:3204:C:O2 | 24:lX:89:ARG:NH1 | 2.39 | 0.52 |
| 2:lB:5:A:HO2' | 18:lR:61:ARG:HH11 | 1.57 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|-------------------|--------------------------|-------------------|
| 22:IV:52:LEU:HD12 | 22:IV:53:PRO:HD2 | 1.91 | 0.52 |
| 33:lg:41:ASN:O | 33:lg:45:ARG:HG3 | 2.09 | 0.52 |
| 52:sa:1382:C:H42 | 52:sa:1390:A:H61 | 1.57 | 0.52 |
| 53:sc:165:GLY:O | 53:sc:168:ARG:NH1 | 2.41 | 0.52 |
| 54:sd:176:ASN:OD1 | 54:sd:180:GLN:NE2 | 2.43 | 0.52 |
| 1:lA:1250:U:OP2 | 12:lL:13:ARG:NH2 | 2.43 | 0.52 |
| 1:lA:1437:A:H2' | 1:lA:1438:U:H6 | 1.74 | 0.52 |
| 1:lA:1498:C:N4 | 1:lA:1502:A:O3' | 2.43 | 0.52 |
| 1:lA:2064:G:N1 | 1:lA:2067:A:OP2 | 2.43 | 0.52 |
| 32:lf:121:GLU:HB3 | 32:lf:124:LYS:HE2 | 1.92 | 0.52 |
| 52:sa:632:U:OP2 | 67:sr:32:LYS:NZ | 2.42 | 0.52 |
| 72:sw:22:ILE:N | 72:sw:89:LEU:O | 2.43 | 0.52 |
| 74:sy:32:THR:HG23 | 74:sy:36:CYS:HB2 | 1.90 | 0.52 |
| 1:lA:2439:A:H2' | 1:lA:2440:G:C8 | 2.45 | 0.52 |
| 1:lA:3021:G:H2' | 1:lA:3022:C:H6 | 1.74 | 0.52 |
| 1:lA:3196:U:H2' | 1:lA:3197:C:C6 | 2.45 | 0.52 |
| 8:lH:141:ALA:HB3 | 8:lH:144:LYS:HD2 | 1.91 | 0.52 |
| 16:lP:24:LEU:HB2 | 16:lP:45:THR:HG21 | 1.92 | 0.52 |
| 17:lQ:11:ASN:ND2 | 17:lQ:44:ASN:OD1 | 2.43 | 0.52 |
| 68:ss:55:ASN:O | 68:ss:57:PRO:HD3 | 2.10 | 0.52 |
| 1:lA:836:C:H2' | 1:lA:837:A:C8 | 2.45 | 0.52 |
| 1:lA:910:A:H5'' | 6:lF:113:ARG:HD2 | 1.90 | 0.52 |
| 1:lA:1006:A:H2' | 1:lA:1007:C:C6 | 2.45 | 0.52 |
| 1:lA:2411:G:N2 | 1:lA:2415:C:O2' | 2.39 | 0.52 |
| 14:lN:122:VAL:HG13 | 14:lN:223:GLN:HG3 | 1.92 | 0.52 |
| 50:sJ:52:C:H2' | 50:sJ:53:G:O4' | 2.10 | 0.52 |
| 52:sa:1441:C:O2' | 52:sa:1442:A:O5' | 2.28 | 0.52 |
| 52:sa:1442:A:H2' | 52:sa:1443:U:H6 | 1.74 | 0.52 |
| 1:lA:2331:A:H8 | 52:sa:1904:G:H21 | 1.55 | 0.51 |
| 6:lF:184:VAL:HG11 | 6:lF:228:GLY:HA3 | 1.92 | 0.51 |
| 48:sE:34:TYR:OH | 52:sa:1594:A:OP1 | 2.20 | 0.51 |
| 52:sa:327:U:OP1 | 60:sj:56:ARG:NH2 | 2.44 | 0.51 |
| 52:sa:1158:A:H2' | 52:sa:1159:A:H8 | 1.75 | 0.51 |
| 1:lA:250:G:H2' | 1:lA:251:A:C8 | 2.46 | 0.51 |
| 1:lA:341:G:OP2 | 17:lQ:15:GLN:NE2 | 2.42 | 0.51 |
| 1:lA:1841:G:H22 | 1:lA:1975:G:N2 | 2.06 | 0.51 |
| 1:lA:2331:A:H2 | 1:lA:2336:U:H3 | 1.59 | 0.51 |
| 1:lA:2950:U:O2' | 1:lA:2952:C:OP1 | 2.28 | 0.51 |
| 3:lC:35:A:H8 | 3:lC:35:A:O5' | 1.93 | 0.51 |
| 6:lF:7:VAL:HG13 | 6:lF:149:GLN:HE22 | 1.75 | 0.51 |
| 52:sa:1301:U:H2' | 52:sa:1302:G:H8 | 1.75 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:lA:1208:U:H4' | 30:ld:43:THR:HG23 | 1.91 | 0.51 |
| 1:lA:1219:A:H4' | 1:lA:1220:A:H5' | 1.92 | 0.51 |
| 1:lA:3154:A:H2' | 1:lA:3155:U:H6 | 1.75 | 0.51 |
| 52:sa:327:U:H5'' | 60:sj:31:ARG:HH11 | 1.74 | 0.51 |
| 52:sa:802:U:C4 | 52:sa:803:G:O6 | 2.63 | 0.51 |
| 67:sr:31:SER:OG | 67:sr:32:LYS:N | 2.43 | 0.51 |
| 1:lA:2109:C:O2 | 5:lE:242:ARG:NH2 | 2.40 | 0.51 |
| 1:lA:3163:G:O2' | 5:lE:14:LEU:O | 2.27 | 0.51 |
| 2:lB:74:A:OP2 | 27:la:51:LYS:HB2 | 2.10 | 0.51 |
| 18:lR:18:ARG:NH1 | 18:lR:20:ASP:OD1 | 2.39 | 0.51 |
| 28:lb:36:ARG:HH12 | 28:lb:74:VAL:HG11 | 1.76 | 0.51 |
| 52:sa:1566:C:H5' | 70:su:130:LEU:HD22 | 1.93 | 0.51 |
| 53:sc:208:THR:HG22 | 53:sc:211:ASN:H | 1.76 | 0.51 |
| 65:sp:19:THR:OG1 | 65:sp:20:LYS:N | 2.42 | 0.51 |
| 23:lW:39:ASP:OD1 | 23:lW:39:ASP:N | 2.39 | 0.51 |
| 52:sa:319:U:OP1 | 63:sm:133:LYS:NZ | 2.31 | 0.51 |
| 1:lA:250:G:H2' | 1:lA:251:A:H8 | 1.76 | 0.51 |
| 1:lA:1847:A:H2' | 1:lA:1848:A:C8 | 2.46 | 0.51 |
| 1:lA:1921:A:H2' | 1:lA:1922:U:O2 | 2.09 | 0.51 |
| 1:lA:2379:A:O2' | 52:sa:1894:G:O2' | 2.03 | 0.51 |
| 31:le:46:VAL:HB | 31:le:71:VAL:HG12 | 1.93 | 0.51 |
| 32:lf:113:MET:HG2 | 32:lf:129:VAL:HG22 | 1.91 | 0.51 |
| 52:sa:1816:U:H2' | 52:sa:1817:A:H8 | 1.72 | 0.51 |
| 56:sf:171:VAL:HG23 | 56:sf:228:LYS:HB2 | 1.91 | 0.51 |
| 63:sm:4:GLN:NE2 | 63:sm:10:GLN:O | 2.44 | 0.51 |
| 1:lA:114:U:P | 10:lJ:142:ARG:HH22 | 2.33 | 0.51 |
| 1:lA:213:A:H62 | 14:lN:131:THR:HG1 | 1.57 | 0.51 |
| 1:lA:598:A:H2' | 1:lA:599:U:C6 | 2.46 | 0.51 |
| 1:lA:625:A:O2' | 1:lA:626:A:OP1 | 2.28 | 0.51 |
| 1:lA:1068:U:O2' | 1:lA:1090:A:OP1 | 2.23 | 0.51 |
| 1:lA:2220:A:H1' | 1:lA:2357:A:N6 | 2.26 | 0.51 |
| 1:lA:3090:G:N3 | 5:lE:252:ALA:HB1 | 2.25 | 0.51 |
| 52:sa:1546:C:H2' | 52:sa:1547:C:H6 | 1.76 | 0.51 |
| 57:sg:183:SER:OG | 57:sg:184:LYS:N | 2.43 | 0.51 |
| 1:lA:765:A:H2' | 1:lA:766:U:C6 | 2.45 | 0.51 |
| 1:lA:2743:G:N2 | 13:lM:20:ASN:HD21 | 2.09 | 0.51 |
| 5:lE:261:ARG:HB2 | 15:lO:65:THR:HG21 | 1.93 | 0.51 |
| 12:lL:93:PRO:HB2 | 12:lL:125:SER:HB3 | 1.91 | 0.51 |
| 23:lW:29:SER:HA | 23:lW:78:THR:HA | 1.93 | 0.51 |
| 52:sa:865:A:OP1 | 55:se:218:LYS:NZ | 2.36 | 0.51 |
| 62:sl:15:LEU:HD13 | 62:sl:21:ILE:HG22 | 1.93 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 1:1A:1090:A:H2' | 1:1A:1091:A:H8 | 1.76 | 0.51 |
| 1:1A:2747:G:N2 | 1:1A:2750:A:OP2 | 2.44 | 0.51 |
| 8:1H:23:THR:OG1 | 8:1H:25:ASP:OD1 | 2.29 | 0.51 |
| 27:1a:87:LYS:O | 27:1a:89:ASN:N | 2.43 | 0.51 |
| 27:1a:179:MET:HE1 | 27:1a:187:TYR:CD2 | 2.46 | 0.51 |
| 37:1k:59:LEU:HD23 | 37:1k:62:LYS:HE2 | 1.93 | 0.51 |
| 50:sJ:23:G:H2' | 50:sJ:24:C:C5 | 2.46 | 0.51 |
| 52:sa:325:U:H2' | 52:sa:326:A:C8 | 2.46 | 0.51 |
| 1:1A:1476:G:H2' | 1:1A:1476:G:N3 | 2.26 | 0.51 |
| 13:1M:27:GLY:O | 13:1M:29:ARG:N | 2.44 | 0.51 |
| 46:sC:31:SER:O | 52:sa:939:G:N2 | 2.43 | 0.51 |
| 52:sa:1693:A:H2' | 52:sa:1694:A:H8 | 1.74 | 0.51 |
| 55:se:197:LYS:HD3 | 55:se:197:LYS:C | 2.35 | 0.51 |
| 56:sf:123:LYS:HB2 | 56:sf:224:PHE:HE1 | 1.75 | 0.51 |
| 57:sg:65:LYS:HD2 | 57:sg:146:LEU:HD21 | 1.93 | 0.51 |
| 1:1A:940:U:H2' | 1:1A:941:A:C8 | 2.46 | 0.50 |
| 1:1A:2699:U:O2' | 1:1A:2828:A:O2' | 2.29 | 0.50 |
| 1:1A:3033:C:O2' | 1:1A:3076:A:N3 | 2.31 | 0.50 |
| 1:1A:3093:G:OP2 | 1:1A:3093:G:N2 | 2.39 | 0.50 |
| 13:1M:16:LYS:HE2 | 13:1M:72:ARG:HH22 | 1.75 | 0.50 |
| 18:1R:57:ILE:HD12 | 18:1R:83:TRP:NE1 | 2.27 | 0.50 |
| 30:ld:2:SER:O | 30:ld:2:SER:OG | 2.26 | 0.50 |
| 52:sa:839:A:H8 | 59:si:119:PRO:HA | 1.77 | 0.50 |
| 53:sc:105:ALA:HB2 | 53:sc:115:LEU:HD12 | 1.93 | 0.50 |
| 69:st:34:VAL:O | 69:st:38:VAL:HG22 | 2.11 | 0.50 |
| 1:1A:495:A:H1' | 1:1A:496:A:C8 | 2.45 | 0.50 |
| 1:1A:564:A:H4' | 1:1A:565:U:O5' | 2.12 | 0.50 |
| 1:1A:1184:A:OP2 | 30:ld:26:LYS:NZ | 2.44 | 0.50 |
| 1:1A:1342:U:H2' | 1:1A:1343:A:H4' | 1.91 | 0.50 |
| 9:1I:87:ILE:HG21 | 9:1I:121:TYR:CE2 | 2.45 | 0.50 |
| 16:1P:99:GLU:O | 16:1P:103:LEU:HD13 | 2.11 | 0.50 |
| 17:1Q:201:LYS:O | 17:1Q:204:ARG:NH1 | 2.44 | 0.50 |
| 19:1S:63:THR:HA | 19:1S:66:LEU:HD12 | 1.93 | 0.50 |
| 22:1V:80:VAL:HG12 | 22:1V:81:ASN:N | 2.21 | 0.50 |
| 52:sa:252:C:H2' | 52:sa:253:A:C8 | 2.46 | 0.50 |
| 52:sa:1823:U:H3' | 52:sa:1824:G:H4' | 1.93 | 0.50 |
| 1:1A:940:U:H2' | 1:1A:941:A:H8 | 1.76 | 0.50 |
| 1:1A:1239:U:OP1 | 29:lc:23:GLY:N | 2.39 | 0.50 |
| 1:1A:1254:A:H1' | 1:1A:2969:U:O2' | 2.11 | 0.50 |
| 3:1C:10:A:H4' | 3:1C:12:C:C6 | 2.46 | 0.50 |
| 5:1E:27:ALA:HB2 | 5:1E:222:VAL:HG23 | 1.92 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 52:sa:1808:G:N3 | 52:sa:1928:A:O2' | 2.43 | 0.50 |
| 53:sc:175:PRO:HG3 | 61:sk:57:LYS:HD3 | 1.93 | 0.50 |
| 61:sk:48:MET:O | 61:sk:52:MET:HE3 | 2.09 | 0.50 |
| 71:sv:99:LYS:NZ | 71:sv:103:TYR:HB2 | 2.26 | 0.50 |
| 1:lA:2179:U:H4' | 21:lU:89:THR:HG22 | 1.93 | 0.50 |
| 1:lA:2223:A:H2' | 1:lA:2224:U:O4' | 2.12 | 0.50 |
| 1:lA:3274:U:H4' | 42:lp:29:PRO:HG3 | 1.93 | 0.50 |
| 11:lK:4:LEU:HD23 | 11:lK:7:LYS:HB2 | 1.94 | 0.50 |
| 41:lo:49:MET:HE3 | 41:lo:51:MET:HG2 | 1.94 | 0.50 |
| 52:sa:1685:C:H2' | 52:sa:1686:U:O4' | 2.11 | 0.50 |
| 55:se:45:VAL:HG21 | 55:se:70:VAL:HG21 | 1.94 | 0.50 |
| 56:sf:112:LEU:HD22 | 56:sf:116:GLN:HG3 | 1.94 | 0.50 |
| 68:ss:22:VAL:HG13 | 68:ss:37:CYS:HB3 | 1.93 | 0.50 |
| 1:lA:720:U:O2' | 1:lA:721:A:OP1 | 2.23 | 0.50 |
| 1:lA:1403:C:H2' | 1:lA:1404:C:H6 | 1.75 | 0.50 |
| 3:lC:17:G:H1 | 3:lC:59:G:H1 | 1.59 | 0.50 |
| 3:lC:103:U:H2' | 3:lC:104:G:C8 | 2.47 | 0.50 |
| 6:lF:370:ASN:H | 6:lF:373:VAL:HG22 | 1.76 | 0.50 |
| 12:lL:30:ARG:HG3 | 12:lL:63:GLU:HG3 | 1.93 | 0.50 |
| 14:lN:46:PRO:HA | 14:lN:49:ILE:HB | 1.93 | 0.50 |
| 27:la:56:LYS:NZ | 27:la:63:LYS:O | 2.42 | 0.50 |
| 42:lp:24:CYS:HA | 42:lp:40:CYS:HB3 | 1.94 | 0.50 |
| 52:sa:1719:G:N2 | 52:sa:1721:A:H3' | 2.26 | 0.50 |
| 55:se:108:THR:HG23 | 55:se:111:LYS:H | 1.77 | 0.50 |
| 1:lA:688:A:H2' | 1:lA:689:A:C8 | 2.46 | 0.50 |
| 1:lA:1601:G:N2 | 1:lA:1608:A:OP2 | 2.37 | 0.50 |
| 14:lN:260:VAL:HG11 | 29:lc:132:GLU:HG2 | 1.93 | 0.50 |
| 16:lP:124:ALA:O | 16:lP:128:ILE:HG12 | 2.11 | 0.50 |
| 52:sa:32:G:H22 | 52:sa:463:A:H5' | 1.75 | 0.50 |
| 1:lA:219:G:H2' | 1:lA:220:A:H8 | 1.77 | 0.50 |
| 12:lL:152:PHE:HB3 | 12:lL:165:LEU:HD21 | 1.93 | 0.50 |
| 23:lW:57:GLN:HB2 | 23:lW:59:ARG:NH2 | 2.27 | 0.50 |
| 46:sC:6:VAL:HG12 | 73:sx:84:MET:HG2 | 1.94 | 0.50 |
| 52:sa:1173:A:O2' | 52:sa:1802:C:OP2 | 2.28 | 0.50 |
| 56:sf:198:VAL:HA | 56:sf:204:GLU:HG3 | 1.93 | 0.50 |
| 67:sr:78:ARG:HB3 | 67:sr:124:ARG:HB3 | 1.94 | 0.50 |
| 1:lA:1325:C:H42 | 1:lA:3000:C:H5'' | 1.77 | 0.50 |
| 1:lA:1327:A:OP2 | 1:lA:1425:A:N6 | 2.43 | 0.50 |
| 1:lA:2637:A:H61 | 1:lA:2649:G:H2' | 1.77 | 0.50 |
| 1:lA:3150:A:H2' | 1:lA:3151:A:C8 | 2.46 | 0.50 |
| 52:sa:468:A:H2' | 52:sa:469:C:O4' | 2.12 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 52:sa:574:A:O2' | 52:sa:576:U:OP1 | 2.29 | 0.50 |
| 52:sa:1442:A:OP2 | 69:st:59:ARG:NE | 2.44 | 0.50 |
| 1:lA:619:A:H2' | 1:lA:619:A:N3 | 2.26 | 0.50 |
| 1:lA:817:A:H2' | 1:lA:818:A:C8 | 2.47 | 0.50 |
| 1:lA:1979:U:HO2' | 1:lA:1981:A:H8 | 1.58 | 0.50 |
| 1:lA:2864:A:C4 | 1:lA:2865:A:C2 | 2.99 | 0.50 |
| 11:lK:41:HIS:ND1 | 11:lK:77:VAL:HG11 | 2.26 | 0.50 |
| 14:lN:58:CYS:HB2 | 14:lN:63:PHE:O | 2.12 | 0.50 |
| 22:IV:40:VAL:HA | 22:IV:99:SER:H | 1.77 | 0.50 |
| 22:IV:122:LEU:HD22 | 22:IV:127:LYS:HD2 | 1.93 | 0.50 |
| 45:sB:59:LEU:HD11 | 65:sp:123:ARG:NE | 2.27 | 0.50 |
| 52:sa:14:C:H2' | 52:sa:15:U:C6 | 2.47 | 0.50 |
| 52:sa:936:U:OP1 | 52:sa:1097:C:O2' | 2.30 | 0.50 |
| 52:sa:1794:U:H2' | 52:sa:1795:G:C8 | 2.47 | 0.50 |
| 1:lA:220:A:H2' | 1:lA:221:U:C6 | 2.46 | 0.49 |
| 1:lA:360:U:O4 | 37:lk:22:LYS:NZ | 2.42 | 0.49 |
| 1:lA:713:G:H3' | 1:lA:714:U:C5' | 2.42 | 0.49 |
| 1:lA:2839:A:H5'' | 43:lq:78:LYS:HD2 | 1.94 | 0.49 |
| 5:lE:229:ASN:ND2 | 5:lE:272:MET:SD | 2.76 | 0.49 |
| 24:IX:24:SER:O | 24:IX:24:SER:OG | 2.30 | 0.49 |
| 27:la:153:ILE:HA | 27:la:156:LYS:HE2 | 1.94 | 0.49 |
| 38:ll:19:CYS:SG | 38:ll:20:LYS:N | 2.85 | 0.49 |
| 63:sm:115:ARG:HH12 | 63:sm:154:ALA:HB3 | 1.77 | 0.49 |
| 68:ss:57:PRO:HA | 68:ss:60:ARG:HH11 | 1.77 | 0.49 |
| 1:lA:5:U:H2' | 1:lA:6:U:C6 | 2.47 | 0.49 |
| 1:lA:589:A:OP1 | 8:lH:35:ARG:NH1 | 2.45 | 0.49 |
| 1:lA:678:A:H2' | 1:lA:679:U:C6 | 2.47 | 0.49 |
| 1:lA:904:A:OP2 | 19:lS:62:SER:OG | 2.25 | 0.49 |
| 1:lA:2867:A:N6 | 1:lA:2887:G:H1 | 2.10 | 0.49 |
| 3:lC:42:U:C2 | 3:lC:43:A:C8 | 2.99 | 0.49 |
| 6:lF:299:GLU:HG2 | 6:lF:304:LEU:HD12 | 1.94 | 0.49 |
| 9:lI:68:ASP:HB3 | 22:IV:141:PRO:HB3 | 1.93 | 0.49 |
| 12:lL:47:PRO:HB2 | 12:lL:178:LYS:HE2 | 1.93 | 0.49 |
| 48:sE:33:LYS:O | 48:sE:36:LEU:HD12 | 2.12 | 0.49 |
| 52:sa:114:G:OP1 | 63:sm:66:ARG:NH1 | 2.44 | 0.49 |
| 52:sa:764:C:H4' | 52:sa:765:A:H5' | 1.93 | 0.49 |
| 52:sa:1834:G:H1 | 52:sa:1880:U:H3 | 1.60 | 0.49 |
| 55:se:34:VAL:HG21 | 55:se:45:VAL:HG22 | 1.94 | 0.49 |
| 59:si:132:LEU:O | 59:si:136:LEU:HD23 | 2.12 | 0.49 |
| 60:sj:207:SER:OG | 60:sj:208:ARG:N | 2.45 | 0.49 |
| 1:lA:3089:A:H5'' | 1:lA:3090:G:H5' | 1.95 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:lA:3155:U:H2' | 1:lA:3156:U:C6 | 2.46 | 0.49 |
| 1:lA:3430:U:O5' | 1:lA:3431:U:H5'' | 2.12 | 0.49 |
| 21:lU:105:LEU:HD23 | 21:lU:138:LEU:HD23 | 1.94 | 0.49 |
| 24:lX:82:ILE:HD12 | 24:lX:121:VAL:HG22 | 1.94 | 0.49 |
| 26:lZ:6:GLN:HB3 | 26:lZ:15:ILE:HD12 | 1.93 | 0.49 |
| 28:lb:83:THR:HG22 | 34:lh:93:PHE:CZ | 2.48 | 0.49 |
| 41:lo:9:TYR:CE1 | 41:lo:51:MET:HE2 | 2.47 | 0.49 |
| 1:lA:518:A:H2' | 1:lA:519:A:O4' | 2.13 | 0.49 |
| 1:lA:705:U:H2' | 1:lA:706:A:H8 | 1.77 | 0.49 |
| 1:lA:773:G:OP1 | 33:lg:41:ASN:ND2 | 2.44 | 0.49 |
| 1:lA:3222:U:OP1 | 21:lU:59:SER:N | 2.38 | 0.49 |
| 5:lE:41:PRO:HA | 5:lE:187:GLY:HA3 | 1.94 | 0.49 |
| 50:sJ:42:C:H2' | 50:sJ:43:G:C8 | 2.47 | 0.49 |
| 52:sa:399:G:H2' | 52:sa:400:C:H6 | 1.78 | 0.49 |
| 52:sa:1596:U:O2' | 52:sa:1681:A:N6 | 2.45 | 0.49 |
| 60:sj:35:ASN:OD1 | 60:sj:35:ASN:N | 2.45 | 0.49 |
| 60:sj:67:TRP:CZ2 | 60:sj:70:GLN:HG3 | 2.47 | 0.49 |
| 66:sq:33:VAL:HG21 | 66:sq:44:TYR:CD2 | 2.47 | 0.49 |
| 1:lA:721:A:H2' | 1:lA:722:A:C8 | 2.48 | 0.49 |
| 1:lA:903:G:H8 | 19:lS:90:TYR:CD2 | 2.30 | 0.49 |
| 1:lA:968:C:H2' | 1:lA:969:U:H6 | 1.77 | 0.49 |
| 1:lA:1090:A:H2' | 1:lA:1091:A:C8 | 2.47 | 0.49 |
| 1:lA:1896:G:OP2 | 31:le:36:LYS:NZ | 2.45 | 0.49 |
| 1:lA:2732:A:H2' | 1:lA:2733:A:H8 | 1.78 | 0.49 |
| 1:lA:2863:A:N1 | 37:lk:16:HIS:HB3 | 2.26 | 0.49 |
| 3:lC:6:G:H5'' | 7:lG:22:ARG:HD3 | 1.94 | 0.49 |
| 3:lC:89:G:H4' | 12:lL:11:LEU:HD22 | 1.95 | 0.49 |
| 3:lC:103:U:H2' | 3:lC:104:G:H8 | 1.76 | 0.49 |
| 4:lD:20:SER:O | 4:lD:20:SER:OG | 2.30 | 0.49 |
| 15:lO:29:LEU:HD21 | 15:lO:89:LEU:HD22 | 1.95 | 0.49 |
| 15:lO:76:SER:OG | 15:lO:107:GLU:OE2 | 2.30 | 0.49 |
| 25:lY:44:PHE:CE1 | 25:lY:64:LEU:HD12 | 2.47 | 0.49 |
| 26:lZ:11:SER:HA | 26:lZ:54:THR:HG23 | 1.94 | 0.49 |
| 60:sj:78:ILE:HG22 | 60:sj:104:ILE:HG22 | 1.95 | 0.49 |
| 63:sm:37:ILE:HG13 | 63:sm:38:GLY:H | 1.76 | 0.49 |
| 63:sm:58:PRO:HB3 | 63:sm:65:ILE:HD11 | 1.93 | 0.49 |
| 67:sr:20:GLN:OE1 | 67:sr:22:LYS:NZ | 2.46 | 0.49 |
| 1:lA:191:A:H2' | 1:lA:192:A:C8 | 2.47 | 0.49 |
| 1:lA:510:A:N1 | 1:lA:561:G:O2' | 2.42 | 0.49 |
| 1:lA:708:U:H2' | 1:lA:709:U:C5 | 2.47 | 0.49 |
| 1:lA:2030:G:H5'' | 1:lA:2031:C:H5' | 1.94 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 2:1B:62:U:O4 | 35:1i:51:ASN:ND2 | 2.43 | 0.49 |
| 11:1K:120:ILE:HG12 | 11:1K:121:GLN:H | 1.77 | 0.49 |
| 49:sI:14:A:N3 | 49:sI:14:A:H2' | 2.28 | 0.49 |
| 52:sa:592:U:H2' | 52:sa:593:A:C8 | 2.47 | 0.49 |
| 52:sa:751:A:H2' | 52:sa:752:A:O4' | 2.12 | 0.49 |
| 52:sa:781:G:C8 | 56:sf:17:MET:HE1 | 2.47 | 0.49 |
| 52:sa:845:A:OP1 | 67:sr:28:ARG:NH2 | 2.45 | 0.49 |
| 52:sa:1162:U:O3' | 74:sy:58:LYS:NZ | 2.45 | 0.49 |
| 56:sf:9:ARG:HH21 | 56:sf:18:LEU:HD22 | 1.78 | 0.49 |
| 1:1A:228:A:H2' | 1:1A:229:A:H8 | 1.78 | 0.49 |
| 1:1A:3383:A:H5'' | 5:1E:156:ARG:HE | 1.78 | 0.49 |
| 3:1C:28:U:H3' | 3:1C:29:G:H21 | 1.78 | 0.49 |
| 28:1b:95:ILE:HD11 | 28:1b:121:PHE:CE1 | 2.48 | 0.49 |
| 52:sa:1266:A:C8 | 66:sq:78:HIS:CE1 | 3.01 | 0.49 |
| 52:sa:1923:A:H2' | 52:sa:1924:G:H8 | 1.77 | 0.49 |
| 56:sf:173:PHE:HE1 | 56:sf:223:VAL:HG11 | 1.77 | 0.49 |
| 1:1A:811:A:N1 | 1:1A:839:G:O2' | 2.41 | 0.49 |
| 1:1A:874:A:H2' | 1:1A:875:C:H6 | 1.78 | 0.49 |
| 1:1A:1095:A:H2 | 1:1A:1099:A:H2 | 1.59 | 0.49 |
| 1:1A:1465:G:H2' | 1:1A:1466:U:C6 | 2.48 | 0.49 |
| 1:1A:1633:G:OP2 | 38:1l:14:LYS:NZ | 2.41 | 0.49 |
| 1:1A:1864:C:OP1 | 23:1W:98:LYS:NZ | 2.41 | 0.49 |
| 9:1I:204:ASN:OD1 | 9:1I:204:ASN:N | 2.45 | 0.49 |
| 13:1M:109:HIS:HE1 | 13:1M:122:ILE:HA | 1.78 | 0.49 |
| 16:1P:100:LYS:O | 16:1P:104:ARG:HG3 | 2.12 | 0.49 |
| 18:1R:126:ARG:HG3 | 18:1R:140:SER:HB3 | 1.95 | 0.49 |
| 47:sD:62:GLU:O | 65:sp:112:ARG:NH2 | 2.43 | 0.49 |
| 52:sa:545:G:OP2 | 52:sa:575:U:O2' | 2.24 | 0.49 |
| 52:sa:932:C:H2' | 52:sa:933:A:H8 | 1.77 | 0.49 |
| 52:sa:1371:U:C5 | 52:sa:1373:A:C8 | 3.01 | 0.49 |
| 65:sp:31:THR:OG1 | 65:sp:32:PHE:N | 2.43 | 0.49 |
| 70:su:97:SER:O | 70:su:99:LEU:HD23 | 2.12 | 0.49 |
| 1:1A:333:U:O2' | 17:1Q:180:ARG:O | 2.30 | 0.49 |
| 1:1A:660:U:H2' | 1:1A:661:G:H8 | 1.78 | 0.49 |
| 1:1A:779:A:H1' | 1:1A:781:A:OP2 | 2.13 | 0.49 |
| 1:1A:868:A:H2' | 1:1A:869:G:C8 | 2.48 | 0.49 |
| 1:1A:2340:U:HO2' | 1:1A:2341:C:P | 2.33 | 0.49 |
| 1:1A:2489:A:H2' | 1:1A:2490:G:H8 | 1.78 | 0.49 |
| 1:1A:2702:G:C6 | 1:1A:2717:A:C6 | 3.01 | 0.49 |
| 2:1B:28:U:H2' | 2:1B:29:U:C6 | 2.47 | 0.49 |
| 2:1B:29:U:H2' | 2:1B:30:U:H6 | 1.78 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 2:1B:141:G:H2' | 2:1B:142:A:C8 | 2.44 | 0.49 |
| 2:1B:144:A:H2' | 2:1B:145:U:C6 | 2.48 | 0.49 |
| 6:1F:369:LEU:C | 6:1F:371:GLN:H | 2.20 | 0.49 |
| 7:1G:115:MET:HE1 | 7:1G:143:PHE:HB2 | 1.95 | 0.49 |
| 21:1U:98:ARG:NH1 | 21:1U:130:ASN:OD1 | 2.46 | 0.49 |
| 52:sa:611:U:H5' | 52:sa:1011:U:O4' | 2.13 | 0.49 |
| 60:sj:25:LYS:O | 60:sj:28:THR:OG1 | 2.31 | 0.49 |
| 66:sq:122:TYR:OH | 70:su:125:ARG:NH1 | 2.46 | 0.49 |
| 1:1A:710:A:O2' | 1:1A:736:G:N2 | 2.23 | 0.49 |
| 1:1A:1027:G:H4' | 1:1A:1028:G:O5' | 2.11 | 0.49 |
| 1:1A:1098:A:H2 | 1:1A:1184:A:H5'' | 1.78 | 0.49 |
| 1:1A:1686:G:N2 | 1:1A:1689:A:OP2 | 2.45 | 0.49 |
| 1:1A:2904:U:H2' | 1:1A:2905:U:C6 | 2.46 | 0.49 |
| 52:sa:443:U:H2' | 52:sa:444:U:C6 | 2.48 | 0.49 |
| 52:sa:1332:A:H2' | 52:sa:1333:G:O4' | 2.13 | 0.49 |
| 52:sa:1917:U:H2' | 52:sa:1918:U:C6 | 2.48 | 0.49 |
| 53:sc:228:THR:OG1 | 53:sc:230:ASP:OD1 | 2.30 | 0.49 |
| 55:se:26:PHE:CD2 | 65:sp:83:LEU:HD21 | 2.48 | 0.49 |
| 1:1A:249:U:H2' | 1:1A:250:G:H8 | 1.78 | 0.48 |
| 1:1A:797:C:H2' | 1:1A:798:U:C6 | 2.48 | 0.48 |
| 1:1A:3211:A:O2' | 32:lf:55:HIS:NE2 | 2.46 | 0.48 |
| 1:1A:3480:A:C5 | 1:1A:3481:U:C5 | 3.01 | 0.48 |
| 2:1B:24:U:C5 | 6:1F:197:MET:HE1 | 2.47 | 0.48 |
| 16:1P:32:ASP:OD1 | 16:1P:35:ARG:NH2 | 2.31 | 0.48 |
| 45:sB:34:LYS:NZ | 52:sa:1940:G:N7 | 2.61 | 0.48 |
| 47:sD:51:ARG:HG3 | 57:sg:63:PHE:HE2 | 1.77 | 0.48 |
| 47:sD:66:ARG:NE | 47:sD:66:ARG:O | 2.46 | 0.48 |
| 52:sa:161:A:H2' | 52:sa:162:U:O4' | 2.12 | 0.48 |
| 52:sa:622:G:OP1 | 64:so:120:SER:OG | 2.31 | 0.48 |
| 52:sa:1243:A:N1 | 52:sa:1551:A:O2' | 2.42 | 0.48 |
| 55:se:174:ILE:HG23 | 55:se:195:LEU:HD21 | 1.94 | 0.48 |
| 1:1A:103:A:O2' | 1:1A:370:A:N3 | 2.44 | 0.48 |
| 1:1A:118:A:N6 | 1:1A:189:G:O2' | 2.46 | 0.48 |
| 1:1A:546:U:H2' | 1:1A:547:A:H8 | 1.78 | 0.48 |
| 1:1A:712:G:O2' | 1:1A:713:G:H5' | 2.14 | 0.48 |
| 1:1A:1968:U:H2' | 1:1A:1969:U:C6 | 2.48 | 0.48 |
| 1:1A:2378:G:H4' | 52:sa:1821:A:H2' | 1.95 | 0.48 |
| 2:1B:145:U:H2' | 2:1B:146:U:C6 | 2.47 | 0.48 |
| 11:1K:134:ILE:HG21 | 11:1K:167:ILE:HG12 | 1.94 | 0.48 |
| 22:1V:164:GLU:OE2 | 22:1V:164:GLU:N | 2.32 | 0.48 |
| 52:sa:170:G:H1' | 52:sa:259:A:N6 | 2.28 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|-------------------|--------------------------|-------------------|
| 52:sa:1241:C:O2 | 52:sa:1551:A:N6 | 2.46 | 0.48 |
| 52:sa:1371:U:C4 | 52:sa:1373:A:C8 | 3.01 | 0.48 |
| 52:sa:1772:A:H2' | 52:sa:1773:G:C8 | 2.47 | 0.48 |
| 58:sh:70:ASN:OD1 | 58:sh:70:ASN:N | 2.46 | 0.48 |
| 63:sm:68:ARG:HB3 | 63:sm:127:GLN:HB2 | 1.94 | 0.48 |
| 63:sm:98:ARG:HG2 | 74:sy:5:LEU:HA | 1.93 | 0.48 |
| 1:lA:943:U:C2 | 1:lA:1021:G:N2 | 2.82 | 0.48 |
| 1:lA:1640:A:H2' | 1:lA:1641:A:C8 | 2.49 | 0.48 |
| 1:lA:1745:A:H2' | 1:lA:1746:G:H8 | 1.78 | 0.48 |
| 1:lA:1847:A:H2' | 1:lA:1848:A:H8 | 1.76 | 0.48 |
| 1:lA:3023:U:OP1 | 24:IX:50:ASN:ND2 | 2.46 | 0.48 |
| 1:lA:3043:A:O2' | 11:lK:180:GLN:OE1 | 2.14 | 0.48 |
| 6:lF:389:LYS:HA | 6:lF:389:LYS:HD3 | 1.61 | 0.48 |
| 11:lK:29:THR:HG22 | 11:lK:34:SER:HB2 | 1.95 | 0.48 |
| 52:sa:787:U:H4' | 52:sa:788:C:H5 | 1.79 | 0.48 |
| 52:sa:1440:A:H1' | 52:sa:1441:C:H5 | 1.79 | 0.48 |
| 66:sq:106:ILE:HD13 | 66:sq:110:MET:HE3 | 1.95 | 0.48 |
| 1:lA:2469:G:P | 1:lA:2469:G:O4' | 2.71 | 0.48 |
| 1:lA:3336:G:H5' | 1:lA:3373:A:H61 | 1.79 | 0.48 |
| 1:lA:3352:G:C5 | 20:lT:155:ALA:HB2 | 2.49 | 0.48 |
| 2:lB:50:A:N7 | 2:lB:53:G:N1 | 2.62 | 0.48 |
| 5:lE:251:VAL:HG21 | 5:lE:267:ALA:HB3 | 1.95 | 0.48 |
| 6:lF:210:VAL:HB | 6:lF:231:THR:HG22 | 1.95 | 0.48 |
| 11:lK:142:VAL:HG22 | 11:lK:152:LEU:HG | 1.95 | 0.48 |
| 22:IV:97:LYS:HG2 | 22:IV:98:PRO:HD2 | 1.95 | 0.48 |
| 28:lb:95:ILE:HD11 | 28:lb:121:PHE:HE1 | 1.78 | 0.48 |
| 43:lq:31:ASP:OD1 | 43:lq:31:ASP:N | 2.46 | 0.48 |
| 52:sa:1693:A:H2' | 52:sa:1694:A:C8 | 2.48 | 0.48 |
| 57:sg:167:ASN:OD1 | 57:sg:168:SER:N | 2.47 | 0.48 |
| 1:lA:793:G:H2' | 1:lA:794:A:N7 | 2.28 | 0.48 |
| 1:lA:975:G:OP1 | 21:lU:92:LYS:NZ | 2.34 | 0.48 |
| 1:lA:1729:G:O2' | 1:lA:1730:G:O4' | 2.26 | 0.48 |
| 1:lA:1902:A:H4' | 1:lA:1915:A:H4' | 1.95 | 0.48 |
| 1:lA:2856:G:H2' | 1:lA:2903:A:N6 | 2.28 | 0.48 |
| 1:lA:3314:U:O2' | 1:lA:3417:G:OP2 | 2.19 | 0.48 |
| 5:lE:62:ARG:NH1 | 5:lE:350:THR:OG1 | 2.47 | 0.48 |
| 5:lE:234:ARG:HG3 | 5:lE:272:MET:HB3 | 1.95 | 0.48 |
| 17:lQ:121:ILE:HG13 | 17:lQ:131:GLU:HG3 | 1.94 | 0.48 |
| 20:lT:7:THR:HG22 | 20:lT:22:ARG:HA | 1.94 | 0.48 |
| 34:lh:62:TYR:O | 34:lh:70:ARG:NH1 | 2.45 | 0.48 |
| 40:ln:49:ASP:HB3 | 40:ln:52:LYS:HG3 | 1.96 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 45:sB:37:LYS:HG3 | 45:sB:72:GLU:HG2 | 1.95 | 0.48 |
| 49:sI:4:C:H2' | 49:sI:5:C:C6 | 2.48 | 0.48 |
| 52:sa:26:U:H2' | 52:sa:27:A:C8 | 2.49 | 0.48 |
| 68:ss:50:PRO:HD2 | 68:ss:53:LEU:HD12 | 1.95 | 0.48 |
| 1:lA:1103:U:H2' | 1:lA:1104:A:H8 | 1.78 | 0.48 |
| 13:lM:151:ASP:N | 13:lM:151:ASP:OD1 | 2.34 | 0.48 |
| 18:lR:70:LYS:HD3 | 18:lR:70:LYS:HA | 1.66 | 0.48 |
| 23:lW:49:PHE:O | 23:lW:53:THR:OG1 | 2.22 | 0.48 |
| 55:se:141:CYS:SG | 55:se:170:MET:HE3 | 2.53 | 0.48 |
| 1:lA:452:A:C2 | 2:lB:19:A:H1' | 2.49 | 0.48 |
| 1:lA:500:G:HO2' | 1:lA:501:U:H6 | 1.60 | 0.48 |
| 1:lA:2685:G:H2' | 1:lA:2686:C:H6 | 1.77 | 0.48 |
| 7:lG:30:TYR:O | 7:lG:34:ARG:HG3 | 2.13 | 0.48 |
| 11:lK:43:LYS:HE3 | 15:lO:132:PRO:HG2 | 1.95 | 0.48 |
| 14:lN:87:LEU:HD22 | 14:lN:220:LEU:HD11 | 1.96 | 0.48 |
| 33:lg:101:VAL:O | 33:lg:126:ARG:NH1 | 2.46 | 0.48 |
| 43:lq:3:VAL:HA | 43:lq:90:GLU:O | 2.14 | 0.48 |
| 50:sJ:20:U:O2 | 50:sJ:20:U:H2' | 2.13 | 0.48 |
| 52:sa:1757:C:H2' | 52:sa:1758:G:H8 | 1.78 | 0.48 |
| 53:sc:82:VAL:HG13 | 53:sc:104:VAL:HG12 | 1.96 | 0.48 |
| 60:sj:37:ARG:HH22 | 60:sj:59:ARG:HH11 | 1.62 | 0.48 |
| 1:lA:219:G:H2' | 1:lA:220:A:C8 | 2.49 | 0.48 |
| 1:lA:278:U:H2' | 1:lA:279:U:H6 | 1.79 | 0.48 |
| 1:lA:495:A:N7 | 1:lA:577:G:O2' | 2.46 | 0.48 |
| 1:lA:1664:G:C2 | 1:lA:1756:U:H4' | 2.49 | 0.48 |
| 1:lA:2062:A:O2' | 1:lA:3220:U:OP1 | 2.32 | 0.48 |
| 1:lA:2240:C:OP1 | 4:lD:8:GLN:NE2 | 2.47 | 0.48 |
| 1:lA:2884:U:H3' | 1:lA:2885:G:H21 | 1.79 | 0.48 |
| 1:lA:3323:C:O2' | 1:lA:3419:A:N1 | 2.43 | 0.48 |
| 12:lL:206:LEU:O | 12:lL:208:LYS:NZ | 2.47 | 0.48 |
| 21:lU:28:GLU:HG3 | 21:lU:49:LEU:HD22 | 1.94 | 0.48 |
| 35:li:70:LYS:HA | 35:li:73:ARG:HE | 1.79 | 0.48 |
| 52:sa:109:U:H2' | 52:sa:110:A:C8 | 2.49 | 0.48 |
| 52:sa:1383:C:H5 | 52:sa:1389:G:H1 | 1.62 | 0.48 |
| 52:sa:1880:U:H2' | 52:sa:1881:C:C6 | 2.48 | 0.48 |
| 1:lA:2256:A:N6 | 1:lA:2257:G:O6 | 2.47 | 0.48 |
| 1:lA:2462:G:H2' | 1:lA:2463:A:C8 | 2.44 | 0.48 |
| 1:lA:2482:C:O2' | 1:lA:2689:G:N2 | 2.47 | 0.48 |
| 1:lA:3404:A:H2' | 8:lH:95:PHE:CZ | 2.49 | 0.48 |
| 2:lB:28:U:H2' | 2:lB:29:U:H6 | 1.79 | 0.48 |
| 2:lB:116:U:H2' | 2:lB:117:A:H8 | 1.79 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 9:II:168:CYS:SG | 9:II:171:ASP:HB2 | 2.54 | 0.48 |
| 47:sD:22:THR:OG1 | 47:sD:23:GLY:N | 2.46 | 0.48 |
| 52:sa:61:A:O2' | 52:sa:261:U:O2 | 2.25 | 0.48 |
| 52:sa:93:C:O2' | 56:sf:5:HIS:HB2 | 2.14 | 0.48 |
| 52:sa:587:U:H4' | 52:sa:589:G:H4' | 1.96 | 0.48 |
| 52:sa:1557:C:H2' | 52:sa:1558:U:H6 | 1.79 | 0.48 |
| 52:sa:1586:U:OP1 | 71:sv:57:ARG:NE | 2.47 | 0.48 |
| 52:sa:1745:U:H2' | 52:sa:1746:C:C6 | 2.48 | 0.48 |
| 55:se:139:ILE:HG22 | 55:se:217:VAL:HG22 | 1.95 | 0.48 |
| 56:sf:104:LYS:NZ | 56:sf:249:GLN:OE1 | 2.47 | 0.48 |
| 63:sm:83:ILE:HG13 | 63:sm:108:VAL:HG23 | 1.95 | 0.48 |
| 1:lA:735:C:H2' | 1:lA:736:G:O4' | 2.13 | 0.48 |
| 1:lA:1689:A:C5 | 1:lA:1690:A:H1' | 2.49 | 0.48 |
| 1:lA:1837:U:C5 | 1:lA:1984:A:H5'' | 2.49 | 0.48 |
| 1:lA:1849:A:H5' | 34:lh:30:MET:HE1 | 1.96 | 0.48 |
| 1:lA:2236:A:H2' | 1:lA:2237:A:H8 | 1.78 | 0.48 |
| 1:lA:2304:A:H2' | 1:lA:2305:C:C6 | 2.49 | 0.48 |
| 1:lA:2628:A:N6 | 4:ID:98:ILE:O | 2.47 | 0.48 |
| 1:lA:2737:A:OP1 | 13:IM:51:ARG:NH2 | 2.47 | 0.48 |
| 4:ID:113:VAL:HB | 4:ID:164:ALA:HB1 | 1.96 | 0.48 |
| 10:IJ:73:LEU:HD22 | 10:IJ:175:CYS:HB2 | 1.94 | 0.48 |
| 46:sC:9:LEU:HD21 | 67:sr:51:THR:HG21 | 1.96 | 0.48 |
| 50:sJ:30:G:H2' | 50:sJ:31:G:O4' | 2.13 | 0.48 |
| 52:sa:258:A:P | 52:sa:285:G:H22 | 2.36 | 0.48 |
| 56:sf:50:LEU:HB3 | 56:sf:52:TYR:CD1 | 2.49 | 0.48 |
| 58:sh:57:ASP:OD2 | 58:sh:98:ARG:HD2 | 2.14 | 0.48 |
| 68:ss:69:ILE:HG22 | 68:ss:70:VAL:HG13 | 1.95 | 0.48 |
| 1:lA:857:U:C2 | 1:lA:887:G:N2 | 2.82 | 0.47 |
| 1:lA:1019:G:H1' | 1:lA:1737:A:N6 | 2.29 | 0.47 |
| 1:lA:1883:A:N6 | 1:lA:1931:U:H3 | 2.11 | 0.47 |
| 1:lA:2144:U:OP2 | 21:IU:74:ARG:NE | 2.36 | 0.47 |
| 1:lA:2688:G:OP1 | 1:lA:2714:C:O2' | 2.20 | 0.47 |
| 1:lA:3038:C:H5' | 42:lp:27:ARG:HH21 | 1.78 | 0.47 |
| 1:lA:3265:C:O2' | 11:IK:161:SER:OG | 2.32 | 0.47 |
| 6:IF:233:ARG:NE | 6:IF:235:GLU:OE1 | 2.32 | 0.47 |
| 6:IF:389:LYS:O | 6:IF:391:LYS:N | 2.47 | 0.47 |
| 13:IM:20:ASN:OD1 | 13:IM:126:ASP:HB2 | 2.14 | 0.47 |
| 16:IP:25:ALA:HB1 | 16:IP:38:VAL:HB | 1.94 | 0.47 |
| 35:li:-9:ALA:HA | 35:li:38:VAL:HG13 | 1.96 | 0.47 |
| 52:sa:251:C:HO2' | 60:sj:64:ASN:HD21 | 1.56 | 0.47 |
| 52:sa:795:G:N2 | 67:sr:108:TYR:OH | 2.47 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 52:sa:1438:A:N6 | 52:sa:1518:A:N7 | 2.61 | 0.47 |
| 52:sa:1439:C:H2' | 52:sa:1440:A:H4' | 1.96 | 0.47 |
| 52:sa:1717:U:O4 | 66:sq:39:ARG:NH2 | 2.46 | 0.47 |
| 52:sa:1872:G:H2' | 52:sa:1873:A:H8 | 1.79 | 0.47 |
| 1:lA:2781:U:O2 | 1:lA:2814:U:H4' | 2.14 | 0.47 |
| 2:lB:141:G:H5'' | 17:lQ:60:VAL:HG11 | 1.95 | 0.47 |
| 27:la:52:ASP:HA | 27:la:67:LYS:HE3 | 1.96 | 0.47 |
| 40:ln:7:GLU:CD | 40:ln:10:GLN:HE22 | 2.22 | 0.47 |
| 47:sD:40:ARG:NH2 | 47:sD:61:ALA:O | 2.47 | 0.47 |
| 52:sa:4:C:OP1 | 53:sc:202:THR:OG1 | 2.29 | 0.47 |
| 52:sa:242:G:O2' | 52:sa:243:U:OP1 | 2.32 | 0.47 |
| 52:sa:327:U:H5'' | 60:sj:31:ARG:NH1 | 2.29 | 0.47 |
| 55:se:76:MET:O | 55:se:77:GLU:HG2 | 2.14 | 0.47 |
| 1:lA:255:A:H4' | 1:lA:257:A:N7 | 2.28 | 0.47 |
| 1:lA:1033:A:C2 | 4:lD:204:MET:HE2 | 2.49 | 0.47 |
| 1:lA:1057:C:C5 | 29:lc:26:ARG:HD2 | 2.50 | 0.47 |
| 1:lA:1105:U:OP1 | 9:lI:113:LYS:NZ | 2.48 | 0.47 |
| 1:lA:1639:U:H5'' | 21:lU:6:LEU:HD22 | 1.96 | 0.47 |
| 1:lA:2224:U:H2' | 1:lA:2225:A:C8 | 2.49 | 0.47 |
| 2:lB:145:U:H2' | 2:lB:146:U:H6 | 1.79 | 0.47 |
| 52:sa:328:C:C5 | 60:sj:49:ARG:HD2 | 2.50 | 0.47 |
| 52:sa:412:A:H5' | 52:sa:413:G:C5 | 2.49 | 0.47 |
| 52:sa:1757:C:O3' | 71:sv:93:ARG:NH2 | 2.47 | 0.47 |
| 52:sa:1828:A:H2' | 52:sa:1829:A:C8 | 2.49 | 0.47 |
| 62:sl:45:MET:HB2 | 62:sl:48:LEU:HD22 | 1.96 | 0.47 |
| 1:lA:244:A:N7 | 1:lA:265:A:N6 | 2.62 | 0.47 |
| 1:lA:526:A:H5'' | 1:lA:538:A:N1 | 2.29 | 0.47 |
| 1:lA:676:A:N7 | 6:lF:358:MET:HG2 | 2.29 | 0.47 |
| 1:lA:802:A:H2' | 1:lA:803:U:C6 | 2.49 | 0.47 |
| 1:lA:905:A:C8 | 19:lS:60:ILE:HD11 | 2.49 | 0.47 |
| 1:lA:926:A:H2 | 1:lA:2487:U:H1' | 1.79 | 0.47 |
| 1:lA:2721:G:H5'' | 1:lA:2722:U:O4' | 2.14 | 0.47 |
| 1:lA:3142:G:N2 | 1:lA:3143:A:H62 | 2.12 | 0.47 |
| 6:lF:19:LYS:HD3 | 6:lF:19:LYS:N | 2.28 | 0.47 |
| 13:lM:109:HIS:CE1 | 13:lM:123:TYR:H | 2.33 | 0.47 |
| 26:lZ:9:ALA:HB2 | 26:lZ:31:PHE:HB3 | 1.96 | 0.47 |
| 42:lp:35:CYS:O | 42:lp:42:HIS:HA | 2.14 | 0.47 |
| 52:sa:375:A:H2' | 52:sa:375:A:N3 | 2.30 | 0.47 |
| 52:sa:972:A:C2 | 52:sa:973:A:C8 | 3.02 | 0.47 |
| 66:sq:32:PHE:HZ | 66:sq:111:LEU:HD13 | 1.78 | 0.47 |
| 68:ss:97:ARG:NH1 | 68:ss:128:ASP:OD2 | 2.45 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 70:su:115:LEU:HA | 70:su:118:MET:HE3 | 1.97 | 0.47 |
| 1:lA:851:A:H5' | 29:lc:114:GLY:O | 2.15 | 0.47 |
| 1:lA:2639:A:OP1 | 28:lb:55:ARG:HG3 | 2.14 | 0.47 |
| 1:lA:3452:U:H3 | 1:lA:3503:G:H1 | 1.61 | 0.47 |
| 1:lA:3455:G:N2 | 1:lA:3500:A:H2 | 2.09 | 0.47 |
| 3:IC:53:C:O2' | 13:IM:152:HIS:ND1 | 2.48 | 0.47 |
| 43:lq:14:ARG:HB2 | 43:lq:77:TYR:CD2 | 2.50 | 0.47 |
| 52:sa:1319:G:H21 | 52:sa:1346:A:H8 | 1.61 | 0.47 |
| 52:sa:1693:A:O2' | 52:sa:1694:A:OP1 | 2.29 | 0.47 |
| 70:su:24:ARG:O | 70:su:56:ARG:NH1 | 2.34 | 0.47 |
| 1:lA:226:U:H2' | 1:lA:227:A:H8 | 1.79 | 0.47 |
| 1:lA:249:U:C2 | 1:lA:250:G:C8 | 3.03 | 0.47 |
| 1:lA:343:A:H62 | 17:lQ:12:ARG:HH12 | 1.62 | 0.47 |
| 1:lA:907:U:H2' | 1:lA:908:A:C8 | 2.49 | 0.47 |
| 1:lA:1033:A:O2' | 4:ID:203:ALA:O | 2.32 | 0.47 |
| 1:lA:1255:A:H2 | 1:lA:3007:A:N3 | 2.13 | 0.47 |
| 1:lA:1928:G:N2 | 1:lA:1929:G:O6 | 2.40 | 0.47 |
| 1:lA:2260:U:OP1 | 4:ID:209:HIS:NE2 | 2.40 | 0.47 |
| 1:lA:2632:A:OP2 | 4:ID:87:TYR:OH | 2.21 | 0.47 |
| 14:IN:238:ARG:HE | 14:IN:241:ALA:HA | 1.80 | 0.47 |
| 20:IT:73:ILE:HD12 | 20:IT:90:TYR:HD2 | 1.79 | 0.47 |
| 24:IX:57:CYS:SG | 24:IX:122:SER:OG | 2.59 | 0.47 |
| 49:sI:14:A:N7 | 49:sI:22:G:C2 | 2.82 | 0.47 |
| 52:sa:325:U:H2' | 52:sa:326:A:H8 | 1.79 | 0.47 |
| 52:sa:1567:A:OP2 | 70:su:145:ARG:NH2 | 2.48 | 0.47 |
| 56:sf:214:ASN:OD1 | 56:sf:214:ASN:N | 2.48 | 0.47 |
| 63:sm:37:ILE:HG21 | 63:sm:41:PHE:HB2 | 1.96 | 0.47 |
| 66:sq:82:LYS:HE3 | 66:sq:88:MET:HE1 | 1.96 | 0.47 |
| 1:lA:322:U:H2' | 1:lA:323:U:C6 | 2.48 | 0.47 |
| 1:lA:332:G:C6 | 43:lq:43:ARG:HD3 | 2.50 | 0.47 |
| 1:lA:547:A:H2' | 1:lA:548:A:C8 | 2.49 | 0.47 |
| 1:lA:859:U:H2' | 1:lA:860:G:O4' | 2.13 | 0.47 |
| 1:lA:1111:G:N3 | 1:lA:2707:A:H2' | 2.29 | 0.47 |
| 1:lA:1565:G:N7 | 29:lc:9:ARG:NH2 | 2.62 | 0.47 |
| 1:lA:2219:A:O2' | 1:lA:2220:A:H2' | 2.15 | 0.47 |
| 3:IC:2:G:O2' | 3:IC:24:U:O2 | 2.32 | 0.47 |
| 5:IE:217:ILE:HD13 | 5:IE:284:ILE:HD11 | 1.96 | 0.47 |
| 7:IG:51:VAL:HG23 | 7:IG:145:CYS:HB3 | 1.97 | 0.47 |
| 10:IJ:85:LEU:HD22 | 10:IJ:129:LEU:HD21 | 1.97 | 0.47 |
| 22:IV:34:TYR:HB2 | 22:IV:72:ILE:HD12 | 1.97 | 0.47 |
| 27:la:2:LYS:HD2 | 27:la:7:VAL:HG23 | 1.97 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 39:lm:51:ALA:HB3 | 39:lm:54:ILE:HD12 | 1.96 | 0.47 |
| 43:lq:23:VAL:HG22 | 43:lq:70:ILE:HG12 | 1.96 | 0.47 |
| 49:sI:15:G:O2' | 49:sI:59:G:N1 | 2.43 | 0.47 |
| 52:sa:345:A:H5'' | 52:sa:347:A:H5' | 1.96 | 0.47 |
| 52:sa:1441:C:HO2' | 52:sa:1442:A:H8 | 1.61 | 0.47 |
| 53:sc:56:ASP:OD1 | 53:sc:57:GLN:N | 2.47 | 0.47 |
| 69:st:16:ILE:HG22 | 69:st:24:LEU:HD11 | 1.97 | 0.47 |
| 1:lA:65:U:O2' | 1:lA:97:A:O2' | 2.19 | 0.47 |
| 1:lA:543:A:H2' | 1:lA:544:U:C6 | 2.49 | 0.47 |
| 1:lA:1225:A:H1' | 9:ll:93:LEU:HD22 | 1.97 | 0.47 |
| 1:lA:1629:G:C2 | 1:lA:1630:A:C8 | 3.03 | 0.47 |
| 1:lA:1632:A:H5' | 38:ll:12:HIS:O | 2.14 | 0.47 |
| 1:lA:2236:A:H2' | 1:lA:2237:A:C8 | 2.50 | 0.47 |
| 1:lA:3352:G:H4' | 1:lA:3353:U:H5'' | 1.96 | 0.47 |
| 5:lE:250:LYS:HB3 | 5:lE:250:LYS:HE2 | 1.74 | 0.47 |
| 10:lJ:242:GLN:HA | 10:lJ:245:ILE:HG22 | 1.96 | 0.47 |
| 32:lf:19:LYS:HE3 | 32:lf:23:GLU:HG2 | 1.97 | 0.47 |
| 52:sa:388:C:H2' | 52:sa:389:U:H6 | 1.80 | 0.47 |
| 52:sa:1185:A:H2' | 52:sa:1186:C:C6 | 2.49 | 0.47 |
| 52:sa:1317:G:H2' | 52:sa:1318:U:C6 | 2.49 | 0.47 |
| 53:sc:53:LYS:HB3 | 53:sc:53:LYS:HE3 | 1.76 | 0.47 |
| 56:sf:44:LEU:HD23 | 56:sf:47:ARG:HH21 | 1.79 | 0.47 |
| 60:sj:85:ALA:HA | 63:sm:7:ARG:HD3 | 1.97 | 0.47 |
| 66:sq:73:LYS:HA | 66:sq:73:LYS:HD3 | 1.70 | 0.47 |
| 1:lA:1741:A:N3 | 1:lA:1763:C:O2' | 2.46 | 0.47 |
| 1:lA:1846:A:H2' | 1:lA:1847:A:C8 | 2.50 | 0.47 |
| 1:lA:2705:A:C3' | 1:lA:2706:C:H5' | 2.45 | 0.47 |
| 1:lA:3329:U:H4' | 1:lA:3411:U:H4' | 1.97 | 0.47 |
| 1:lA:3455:G:H1 | 1:lA:3500:A:H2 | 1.60 | 0.47 |
| 3:lC:48:U:OP1 | 7:lG:221:ARG:HB2 | 2.15 | 0.47 |
| 4:lD:36:GLU:OE2 | 4:lD:163:ARG:HD3 | 2.15 | 0.47 |
| 7:lG:37:ILE:HD12 | 7:lG:67:SER:HB2 | 1.97 | 0.47 |
| 7:lG:51:VAL:HG12 | 7:lG:53:VAL:HG23 | 1.97 | 0.47 |
| 10:lJ:147:LEU:HD12 | 10:lJ:173:PRO:HB2 | 1.97 | 0.47 |
| 15:lO:177:LYS:HD3 | 15:lO:177:LYS:HA | 1.75 | 0.47 |
| 19:lS:156:LYS:HD2 | 19:lS:156:LYS:HA | 1.60 | 0.47 |
| 24:lX:112:MET:HE1 | 24:lX:117:ILE:HG13 | 1.97 | 0.47 |
| 43:lq:19:THR:O | 43:lq:21:HIS:HD2 | 1.98 | 0.47 |
| 52:sa:1126:G:H8 | 67:sr:76:SER:OG | 1.98 | 0.47 |
| 52:sa:1751:U:H5'' | 68:ss:149:ALA:HB3 | 1.97 | 0.47 |
| 56:sf:40:LEU:HD12 | 56:sf:41:PRO:HD2 | 1.96 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 1:1A:2261:G:O2' | 1:1A:2390:U:OP2 | 2.32 | 0.47 |
| 3:1C:22:C:H2' | 3:1C:23:A:C8 | 2.46 | 0.47 |
| 3:1C:94:U:H2' | 3:1C:95:A:C8 | 2.50 | 0.47 |
| 6:1F:365:LYS:HA | 6:1F:397:LEU:HD11 | 1.96 | 0.47 |
| 7:1G:219:PHE:HB3 | 7:1G:222:PHE:HB2 | 1.96 | 0.47 |
| 9:1I:143:LYS:HB2 | 9:1I:143:LYS:HE3 | 1.61 | 0.47 |
| 26:1Z:61:ASN:N | 26:1Z:61:ASN:OD1 | 2.47 | 0.47 |
| 27:1a:109:LEU:HA | 27:1a:113:ARG:HG3 | 1.97 | 0.47 |
| 28:1b:53:ILE:HD12 | 28:1b:62:GLN:HB3 | 1.97 | 0.47 |
| 28:1b:126:ARG:H | 28:1b:126:ARG:HG2 | 1.50 | 0.47 |
| 48:sE:41:ARG:HG2 | 72:sw:69:PRO:HA | 1.95 | 0.47 |
| 52:sa:96:C:H2' | 52:sa:97:U:C6 | 2.50 | 0.47 |
| 52:sa:383:G:OP2 | 52:sa:418:G:O2' | 2.33 | 0.47 |
| 52:sa:541:U:H2' | 52:sa:542:G:C8 | 2.50 | 0.47 |
| 52:sa:628:G:N1 | 52:sa:945:U:OP2 | 2.46 | 0.47 |
| 67:sr:41:MET:HE2 | 67:sr:41:MET:HB2 | 1.77 | 0.47 |
| 1:1A:82:G:O2' | 1:1A:94:G:O6 | 2.29 | 0.46 |
| 1:1A:187:G:H5'' | 1:1A:188:U:H3' | 1.98 | 0.46 |
| 1:1A:210:U:OP1 | 14:1N:215:LYS:NZ | 2.42 | 0.46 |
| 1:1A:310:A:N1 | 1:1A:2873:G:O2' | 2.45 | 0.46 |
| 1:1A:877:U:H2' | 1:1A:878:U:C6 | 2.50 | 0.46 |
| 1:1A:1816:A:H1' | 1:1A:1892:C:O2' | 2.15 | 0.46 |
| 1:1A:3039:A:H4' | 42:lp:20:ILE:HG21 | 1.96 | 0.46 |
| 1:1A:3090:G:C2 | 5:1E:252:ALA:HB1 | 2.51 | 0.46 |
| 2:1B:63:A:OP1 | 35:li:41:LYS:NZ | 2.48 | 0.46 |
| 52:sa:619:C:H2' | 52:sa:620:U:C6 | 2.50 | 0.46 |
| 52:sa:939:G:C5 | 64:so:17:PRO:HG3 | 2.50 | 0.46 |
| 69:st:31:ASN:O | 69:st:35:VAL:HG23 | 2.15 | 0.46 |
| 1:1A:200:A:H2' | 1:1A:201:A:C8 | 2.49 | 0.46 |
| 1:1A:591:C:H2' | 1:1A:592:A:C8 | 2.48 | 0.46 |
| 1:1A:1120:A:OP2 | 7:1G:10:ARG:NH1 | 2.37 | 0.46 |
| 1:1A:1483:U:O2' | 9:1I:149:MET:SD | 2.73 | 0.46 |
| 1:1A:3091:C:H5'' | 5:1E:245:HIS:HB3 | 1.96 | 0.46 |
| 1:1A:3370:A:H2' | 1:1A:3371:U:C6 | 2.50 | 0.46 |
| 3:1C:104:G:C2 | 3:1C:105:A:C5 | 3.03 | 0.46 |
| 4:1D:62:VAL:HG22 | 4:1D:73:LYS:HG2 | 1.96 | 0.46 |
| 4:1D:177:LYS:NZ | 39:lm:33:GLN:OE1 | 2.48 | 0.46 |
| 11:1K:24:LYS:HB2 | 11:1K:24:LYS:HE2 | 1.76 | 0.46 |
| 52:sa:604:G:HO2' | 52:sa:607:G:HO2' | 1.61 | 0.46 |
| 52:sa:905:G:C2 | 52:sa:968:A:H1' | 2.50 | 0.46 |
| 52:sa:1674:G:H5' | 52:sa:1717:U:H5' | 1.97 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 52:sa:1756:U:H2' | 52:sa:1757:C:H6 | 1.80 | 0.46 |
| 56:sf:123:LYS:HB3 | 56:sf:140:HIS:HB2 | 1.97 | 0.46 |
| 68:ss:51:LEU:HB3 | 68:ss:64:PHE:HE1 | 1.80 | 0.46 |
| 1:lA:579:A:C4 | 1:lA:580:A:C8 | 3.03 | 0.46 |
| 1:lA:723:U:O2' | 1:lA:724:A:H5' | 2.15 | 0.46 |
| 1:lA:779:A:O2' | 1:lA:783:A:N6 | 2.49 | 0.46 |
| 1:lA:789:C:H2' | 1:lA:790:A:C8 | 2.48 | 0.46 |
| 1:lA:1310:A:H2' | 1:lA:1311:C:C6 | 2.50 | 0.46 |
| 1:lA:1930:C:H2' | 1:lA:1931:U:H6 | 1.81 | 0.46 |
| 1:lA:2851:A:H2' | 1:lA:2852:G:C8 | 2.50 | 0.46 |
| 1:lA:3204:C:O2' | 26:lZ:18:ALA:O | 2.33 | 0.46 |
| 5:lE:153:ILE:O | 5:lE:157:CYS:HB2 | 2.16 | 0.46 |
| 8:lH:76:LYS:NZ | 8:lH:190:TYR:O | 2.49 | 0.46 |
| 9:lI:21:CYS:SG | 9:lI:22:LYS:N | 2.86 | 0.46 |
| 10:lJ:58:ARG:HD2 | 10:lJ:233:ILE:O | 2.14 | 0.46 |
| 20:lT:19:GLU:OE2 | 22:IV:155:THR:HG21 | 2.16 | 0.46 |
| 43:lq:12:CYS:HB2 | 43:lq:21:HIS:CE1 | 2.50 | 0.46 |
| 44:sA:106:ILE:HD13 | 44:sA:108:ILE:HD12 | 1.97 | 0.46 |
| 49:sI:51:A:H2' | 49:sI:52:G:O4' | 2.14 | 0.46 |
| 59:si:154:LYS:HB2 | 59:si:154:LYS:HE3 | 1.65 | 0.46 |
| 72:sw:22:ILE:HB | 72:sw:89:LEU:HB2 | 1.97 | 0.46 |
| 1:lA:560:U:H2' | 1:lA:561:G:C8 | 2.50 | 0.46 |
| 1:lA:832:A:O2' | 1:lA:833:U:OP1 | 2.29 | 0.46 |
| 1:lA:1078:C:O2 | 1:lA:2684:G:O2' | 2.33 | 0.46 |
| 1:lA:1095:A:H2 | 1:lA:1099:A:C2 | 2.33 | 0.46 |
| 1:lA:2373:U:C2 | 1:lA:2375:A:C6 | 3.03 | 0.46 |
| 1:lA:3214:C:H1' | 1:lA:3459:U:H1' | 1.98 | 0.46 |
| 1:lA:3278:A:N1 | 11:lK:80:THR:OG1 | 2.49 | 0.46 |
| 49:sI:44:U:H2' | 49:sI:45:G:O4' | 2.16 | 0.46 |
| 52:sa:158:A:C8 | 52:sa:159:A:C8 | 3.04 | 0.46 |
| 52:sa:840:U:H2' | 52:sa:841:U:H6 | 1.79 | 0.46 |
| 52:sa:1206:U:H4' | 66:sq:126:THR:HG23 | 1.98 | 0.46 |
| 60:sj:98:LYS:HB2 | 60:sj:207:SER:O | 2.15 | 0.46 |
| 71:sv:41:GLY:HA2 | 71:sv:97:SER:H | 1.81 | 0.46 |
| 74:sy:83:VAL:HG21 | 74:sy:89:LEU:HD13 | 1.98 | 0.46 |
| 1:lA:564:A:H1' | 1:lA:565:U:OP2 | 2.15 | 0.46 |
| 1:lA:1033:A:N7 | 4:lD:199:VAL:HG21 | 2.31 | 0.46 |
| 1:lA:1311:C:O3' | 1:lA:1419:G:N2 | 2.48 | 0.46 |
| 1:lA:1763:C:H2' | 1:lA:1764:A:C8 | 2.50 | 0.46 |
| 1:lA:2239:U:O2' | 4:lD:11:GLY:HA3 | 2.16 | 0.46 |
| 1:lA:2867:A:H62 | 1:lA:2887:G:N2 | 2.13 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 2:1B:57:U:N3 | 2:1B:58:G:N7 | 2.64 | 0.46 |
| 2:1B:85:U:H3' | 2:1B:86:G:H5' | 1.97 | 0.46 |
| 6:1F:242:LEU:HD22 | 6:1F:252:ILE:HD11 | 1.97 | 0.46 |
| 9:1I:96:LEU:HD21 | 9:1I:103:VAL:HG22 | 1.97 | 0.46 |
| 14:1N:148:LYS:HE3 | 14:1N:152:ILE:HD11 | 1.98 | 0.46 |
| 24:1X:88:TRP:HH2 | 24:1X:98:PHE:HE1 | 1.62 | 0.46 |
| 50:sJ:39:A:O2' | 52:sa:981:A:OP1 | 2.25 | 0.46 |
| 52:sa:395:G:H1 | 60:sj:29:MET:HE1 | 1.80 | 0.46 |
| 52:sa:746:A:H2' | 52:sa:747:A:H8 | 1.80 | 0.46 |
| 52:sa:982:G:O6 | 52:sa:983:A:N6 | 2.48 | 0.46 |
| 52:sa:1033:A:H61 | 52:sa:1091:C:H42 | 1.62 | 0.46 |
| 52:sa:1439:C:H5' | 69:st:48:ASN:HB3 | 1.98 | 0.46 |
| 52:sa:1822:U:O4 | 52:sa:1891:A:H1' | 2.16 | 0.46 |
| 1:1A:59:A:H8 | 1:1A:59:A:O5' | 1.99 | 0.46 |
| 1:1A:1765:A:H2' | 1:1A:1766:G:O4' | 2.15 | 0.46 |
| 1:1A:2230:U:H2' | 1:1A:2231:C:C6 | 2.50 | 0.46 |
| 1:1A:3410:A:H1' | 1:1A:3411:U:H5 | 1.81 | 0.46 |
| 2:1B:102:A:C8 | 2:1B:103:A:C8 | 3.03 | 0.46 |
| 6:1F:150:ILE:HD12 | 6:1F:150:ILE:HA | 1.78 | 0.46 |
| 9:1I:185:GLU:OE1 | 9:1I:185:GLU:N | 2.46 | 0.46 |
| 52:sa:150:G:N1 | 52:sa:152:A:H5'' | 2.30 | 0.46 |
| 52:sa:597:U:H2' | 52:sa:598:A:H8 | 1.81 | 0.46 |
| 52:sa:1224:G:OP1 | 52:sa:1225:G:H8 | 1.98 | 0.46 |
| 52:sa:1234:C:H42 | 52:sa:1562:G:H22 | 1.64 | 0.46 |
| 52:sa:1671:A:N3 | 52:sa:1729:G:O2' | 2.34 | 0.46 |
| 52:sa:1776:G:H5'' | 57:sg:88:LYS:HB2 | 1.97 | 0.46 |
| 54:sd:119:ILE:HG22 | 54:sd:198:ILE:HD13 | 1.97 | 0.46 |
| 56:sf:196:ARG:HB3 | 56:sf:206:VAL:HG23 | 1.98 | 0.46 |
| 62:sl:1:MET:HE1 | 62:sl:39:LEU:HG | 1.96 | 0.46 |
| 65:sp:48:ILE:HG22 | 65:sp:49:ILE:HG22 | 1.97 | 0.46 |
| 1:1A:527:G:H1 | 1:1A:545:U:H3 | 1.64 | 0.46 |
| 1:1A:718:A:H2' | 1:1A:719:U:H6 | 1.78 | 0.46 |
| 1:1A:1675:A:OP2 | 1:1A:1734:G:N1 | 2.32 | 0.46 |
| 1:1A:3473:U:H2' | 1:1A:3474:U:H6 | 1.80 | 0.46 |
| 5:1E:48:GLY:HA3 | 5:1E:81:THR:HG22 | 1.97 | 0.46 |
| 7:1G:189:GLU:HA | 7:1G:192:ARG:HB3 | 1.97 | 0.46 |
| 8:1H:187:MET:HE3 | 8:1H:187:MET:HB3 | 1.76 | 0.46 |
| 8:1H:197:ASP:HB3 | 8:1H:202:LEU:HD21 | 1.97 | 0.46 |
| 17:1Q:17:ASP:N | 17:1Q:17:ASP:OD1 | 2.48 | 0.46 |
| 45:sB:43:ASN:OD1 | 45:sB:43:ASN:N | 2.49 | 0.46 |
| 48:sE:8:ASN:ND2 | 62:sl:28:VAL:HG12 | 2.31 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 52:sa:251:C:H2' | 52:sa:252:C:H6 | 1.80 | 0.46 |
| 54:sd:38:LEU:HD13 | 62:sl:64:TYR:HE2 | 1.81 | 0.46 |
| 62:sl:39:LEU:O | 62:sl:43:MET:HB2 | 2.16 | 0.46 |
| 1:lA:1581:G:N2 | 1:lA:2432:A:OP2 | 2.38 | 0.46 |
| 4:lD:204:MET:SD | 4:lD:209:HIS:HB2 | 2.56 | 0.46 |
| 6:lF:135:ALA:HB3 | 27:la:175:ALA:HB1 | 1.98 | 0.46 |
| 6:lF:403:ARG:HE | 6:lF:411:MET:HE1 | 1.81 | 0.46 |
| 8:lH:120:SER:O | 8:lH:121:LYS:HG2 | 2.16 | 0.46 |
| 11:lK:71:ASN:OD1 | 11:lK:71:ASN:N | 2.47 | 0.46 |
| 15:lO:92:ASN:HA | 15:lO:97:LYS:HZ1 | 1.81 | 0.46 |
| 25:lY:55:ILE:HD11 | 25:lY:121:PHE:CE2 | 2.51 | 0.46 |
| 52:sa:1311:U:O2' | 52:sa:1312:A:O4' | 2.34 | 0.46 |
| 52:sa:1843:U:H2' | 52:sa:1844:A:H8 | 1.80 | 0.46 |
| 53:sc:151:GLY:HA2 | 53:sc:152:GLU:HA | 1.58 | 0.46 |
| 56:sf:87:VAL:HG13 | 56:sf:120:LYS:HB2 | 1.97 | 0.46 |
| 1:lA:2224:U:O2' | 4:lD:182:ALA:HB2 | 2.15 | 0.46 |
| 1:lA:2395:C:HO2' | 1:lA:2396:A:H8 | 1.63 | 0.46 |
| 13:lM:55:ARG:H | 13:lM:55:ARG:HD3 | 1.81 | 0.46 |
| 39:lm:24:LYS:HA | 39:lm:24:LYS:HD3 | 1.69 | 0.46 |
| 52:sa:1277:C:H2' | 52:sa:1278:A:H8 | 1.80 | 0.46 |
| 52:sa:1678:A:H2' | 52:sa:1679:G:O4' | 2.16 | 0.46 |
| 1:lA:828:A:OP1 | 17:lQ:202:ARG:NH2 | 2.48 | 0.46 |
| 1:lA:2301:U:C2 | 1:lA:2302:U:C6 | 3.04 | 0.46 |
| 1:lA:3103:C:H2' | 1:lA:3104:G:H8 | 1.80 | 0.46 |
| 1:lA:3104:G:H2' | 1:lA:3105:U:C6 | 2.51 | 0.46 |
| 1:lA:3369:U:H2' | 1:lA:3370:A:C8 | 2.51 | 0.46 |
| 1:lA:3435:G:OP1 | 5:lE:271:GLN:NE2 | 2.49 | 0.46 |
| 17:lQ:165:THR:O | 17:lQ:169:LYS:HB2 | 2.16 | 0.46 |
| 19:lS:112:LYS:HE3 | 19:lS:112:LYS:HB2 | 1.61 | 0.46 |
| 33:lg:33:TRP:CZ2 | 33:lg:54:PRO:HD3 | 2.51 | 0.46 |
| 52:sa:338:C:H2' | 52:sa:339:A:C8 | 2.51 | 0.46 |
| 52:sa:400:C:H5'' | 58:sh:93:ARG:HH21 | 1.81 | 0.46 |
| 52:sa:1590:A:H2 | 52:sa:1773:G:H1' | 1.80 | 0.46 |
| 53:sc:172:VAL:HB | 53:sc:199:PHE:HB2 | 1.98 | 0.46 |
| 70:su:27:VAL:HG23 | 70:su:55:LYS:O | 2.16 | 0.46 |
| 1:lA:657:A:O2' | 1:lA:658:A:OP2 | 2.26 | 0.45 |
| 1:lA:1485:A:H2' | 1:lA:1486:C:O4' | 2.16 | 0.45 |
| 1:lA:1520:A:H4' | 1:lA:1521:A:O5' | 2.15 | 0.45 |
| 1:lA:1533:U:O2' | 1:lA:1534:U:OP1 | 2.29 | 0.45 |
| 1:lA:3474:U:H2' | 1:lA:3475:U:O4' | 2.16 | 0.45 |
| 10:lJ:47:ARG:O | 10:lJ:51:GLU:HG2 | 2.16 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 21:1U:122:ASP:C | 21:1U:122:ASP:OD1 | 2.59 | 0.45 |
| 22:1V:97:LYS:HE2 | 22:1V:97:LYS:HB3 | 1.78 | 0.45 |
| 32:1f:68:PRO:HD3 | 32:1f:102:ILE:HG13 | 1.98 | 0.45 |
| 52:sa:5:U:H2' | 52:sa:6:G:C8 | 2.51 | 0.45 |
| 52:sa:96:C:H2' | 52:sa:97:U:H6 | 1.80 | 0.45 |
| 52:sa:750:U:O4 | 52:sa:751:A:N6 | 2.49 | 0.45 |
| 52:sa:882:G:H1 | 65:sp:60:ASP:CG | 2.24 | 0.45 |
| 67:sr:28:ARG:HB3 | 67:sr:29:PRO:HD3 | 1.98 | 0.45 |
| 1:1A:424:A:O2' | 1:1A:425:C:OP2 | 2.26 | 0.45 |
| 1:1A:739:A:H2' | 1:1A:740:A:H8 | 1.82 | 0.45 |
| 1:1A:1048:A:H2' | 1:1A:1049:U:C6 | 2.51 | 0.45 |
| 1:1A:1305:A:H1' | 1:1A:1306:A:C8 | 2.50 | 0.45 |
| 1:1A:2346:A:H2' | 1:1A:2347:A:C8 | 2.51 | 0.45 |
| 1:1A:2972:U:C4 | 1:1A:2973:G:N7 | 2.84 | 0.45 |
| 6:1F:114:LYS:HG3 | 17:1Q:203:TYR:HB3 | 1.98 | 0.45 |
| 15:1O:33:LYS:HA | 15:1O:102:ASN:HB3 | 1.98 | 0.45 |
| 15:1O:130:PHE:CE1 | 15:1O:136:ARG:HD2 | 2.52 | 0.45 |
| 52:sa:153:U:H1' | 52:sa:415:A:O2' | 2.17 | 0.45 |
| 52:sa:412:A:H4' | 52:sa:413:G:O5' | 2.15 | 0.45 |
| 52:sa:1591:G:H2' | 52:sa:1592:A:C8 | 2.51 | 0.45 |
| 56:sf:85:MET:HB3 | 56:sf:100:MET:SD | 2.57 | 0.45 |
| 59:si:137:TYR:CD1 | 59:si:138:PRO:HA | 2.51 | 0.45 |
| 60:sj:22:GLN:HG3 | 60:sj:25:LYS:HE3 | 1.98 | 0.45 |
| 1:1A:353:G:O2' | 1:1A:2299:A:H1' | 2.17 | 0.45 |
| 1:1A:709:U:O2' | 1:1A:710:A:OP1 | 2.29 | 0.45 |
| 11:1K:140:VAL:HG22 | 11:1K:160:VAL:HG22 | 1.96 | 0.45 |
| 52:sa:209:A:N6 | 52:sa:250:G:O6 | 2.50 | 0.45 |
| 52:sa:1272:U:H2' | 52:sa:1273:C:C6 | 2.52 | 0.45 |
| 52:sa:1895:G:H2' | 52:sa:1896:A:C8 | 2.51 | 0.45 |
| 56:sf:255:LYS:HE2 | 56:sf:255:LYS:HB2 | 1.70 | 0.45 |
| 68:ss:60:ARG:HH11 | 68:ss:60:ARG:HG3 | 1.81 | 0.45 |
| 73:sx:12:PRO:HB2 | 73:sx:21:LEU:HD21 | 1.99 | 0.45 |
| 1:1A:720:U:C2 | 1:1A:721:A:N7 | 2.84 | 0.45 |
| 1:1A:3469:A:H5'' | 1:1A:3470:G:N7 | 2.30 | 0.45 |
| 2:1B:24:U:H5 | 6:1F:197:MET:HE1 | 1.81 | 0.45 |
| 4:1D:230:PRO:HD2 | 4:1D:233:GLN:HG2 | 1.98 | 0.45 |
| 7:1G:278:LYS:HB2 | 7:1G:278:LYS:HE2 | 1.73 | 0.45 |
| 32:1f:121:GLU:CD | 32:1f:123:GLY:H | 2.23 | 0.45 |
| 52:sa:208:C:H2' | 52:sa:209:A:C8 | 2.51 | 0.45 |
| 53:sc:185:THR:HG21 | 53:sc:209:LEU:HD12 | 1.99 | 0.45 |
| 56:sf:50:LEU:HB3 | 56:sf:52:TYR:HD1 | 1.81 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 62:sl:1:MET:HE2 | 62:sl:40:GLN:HA | 1.97 | 0.45 |
| 71:sv:112:GLY:O | 71:sv:124:THR:OG1 | 2.33 | 0.45 |
| 1:lA:285:A:H4' | 1:lA:286:U:C5' | 2.47 | 0.45 |
| 1:lA:2449:A:N3 | 1:lA:2967:G:O2' | 2.45 | 0.45 |
| 1:lA:3370:A:C2 | 16:lP:128:ILE:HD12 | 2.52 | 0.45 |
| 1:lA:3386:G:H2' | 1:lA:3387:G:H8 | 1.81 | 0.45 |
| 1:lA:3461:U:H4' | 1:lA:3462:A:H5' | 1.99 | 0.45 |
| 18:lR:57:ILE:HB | 18:lR:72:GLN:HB2 | 1.97 | 0.45 |
| 32:lf:11:VAL:O | 32:lf:15:ARG:HG3 | 2.16 | 0.45 |
| 34:lh:7:TYR:HE1 | 34:lh:20:ILE:HG12 | 1.81 | 0.45 |
| 52:sa:323:A:H2' | 52:sa:324:G:C8 | 2.52 | 0.45 |
| 52:sa:1442:A:H2' | 52:sa:1443:U:C6 | 2.51 | 0.45 |
| 52:sa:1565:G:H2' | 52:sa:1565:G:N3 | 2.32 | 0.45 |
| 52:sa:1808:G:O2' | 52:sa:1928:A:O3' | 2.34 | 0.45 |
| 1:lA:265:A:H2' | 1:lA:265:A:N3 | 2.32 | 0.45 |
| 1:lA:612:A:O4' | 1:lA:716:A:N6 | 2.50 | 0.45 |
| 1:lA:846:A:H2' | 1:lA:847:A:C8 | 2.52 | 0.45 |
| 1:lA:979:G:C5 | 4:lD:181:LYS:HB3 | 2.52 | 0.45 |
| 1:lA:1992:A:H2' | 1:lA:1993:G:H8 | 1.82 | 0.45 |
| 1:lA:2744:A:C6 | 13:lM:124:GLY:HA3 | 2.51 | 0.45 |
| 1:lA:2863:A:C2 | 37:lk:16:HIS:HB3 | 2.52 | 0.45 |
| 6:lF:102:PHE:O | 6:lF:103:SER:C | 2.59 | 0.45 |
| 7:lG:232:LEU:HA | 7:lG:235:ILE:HG22 | 1.99 | 0.45 |
| 14:lN:41:ALA:HB2 | 14:lN:48:THR:HG21 | 1.97 | 0.45 |
| 15:lO:175:LYS:HD3 | 15:lO:175:LYS:HA | 1.72 | 0.45 |
| 27:la:83:LEU:HD23 | 27:la:97:ILE:HD11 | 1.98 | 0.45 |
| 52:sa:841:U:H2' | 52:sa:842:A:C5 | 2.52 | 0.45 |
| 54:sd:115:GLN:O | 54:sd:118:SER:OG | 2.29 | 0.45 |
| 56:sf:157:THR:HB | 56:sf:171:VAL:HG13 | 1.98 | 0.45 |
| 58:sh:62:PRO:HD2 | 58:sh:97:VAL:HG12 | 1.99 | 0.45 |
| 63:sm:52:TYR:CD2 | 63:sm:112:PRO:HG2 | 2.51 | 0.45 |
| 64:so:102:LEU:HA | 64:so:102:LEU:HD23 | 1.81 | 0.45 |
| 66:sq:107:LYS:HG3 | 66:sq:109:GLU:N | 2.31 | 0.45 |
| 1:lA:504:U:H2' | 1:lA:505:A:H8 | 1.82 | 0.45 |
| 1:lA:1444:U:C4 | 1:lA:1445:U:O4 | 2.70 | 0.45 |
| 1:lA:2089:A:C2 | 1:lA:2467:G:H4' | 2.52 | 0.45 |
| 1:lA:3148:C:H2' | 1:lA:3149:A:C8 | 2.52 | 0.45 |
| 2:lB:15:U:H5' | 18:lR:2:VAL:HG13 | 1.99 | 0.45 |
| 3:lC:69:U:H2' | 3:lC:70:U:H6 | 1.82 | 0.45 |
| 7:lG:9:ASN:OD1 | 7:lG:9:ASN:N | 2.50 | 0.45 |
| 14:lN:44:VAL:HG23 | 14:lN:47:ALA:HB2 | 1.98 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 15:lO:63:VAL:HG12 | 15:lO:66:ASN:H | 1.82 | 0.45 |
| 52:sa:1110:G:N2 | 52:sa:1113:A:OP2 | 2.43 | 0.45 |
| 56:sf:57:ARG:O | 56:sf:60:THR:OG1 | 2.34 | 0.45 |
| 58:sh:74:ARG:HG2 | 58:sh:96:THR:HB | 1.98 | 0.45 |
| 64:so:3:ARG:HB2 | 64:so:6:ASN:O | 2.17 | 0.45 |
| 71:sv:50:ASN:ND2 | 71:sv:52:ASP:OD1 | 2.49 | 0.45 |
| 1:lA:10:U:H2' | 1:lA:11:A:H8 | 1.81 | 0.45 |
| 1:lA:11:A:H2' | 1:lA:12:U:C6 | 2.52 | 0.45 |
| 1:lA:1314:A:C2 | 15:lO:134:ARG:HD3 | 2.52 | 0.45 |
| 1:lA:1430:G:O2' | 1:lA:1431:G:H5'' | 2.16 | 0.45 |
| 1:lA:1904:U:OP2 | 21:lU:124:TYR:OH | 2.29 | 0.45 |
| 1:lA:2861:U:O4' | 1:lA:2892:A:N6 | 2.49 | 0.45 |
| 15:lO:62:ARG:HD2 | 15:lO:67:PRO:HB3 | 1.98 | 0.45 |
| 18:lR:52:ILE:HA | 18:lR:85:VAL:HG22 | 1.99 | 0.45 |
| 22:lV:80:VAL:CG1 | 22:lV:81:ASN:H | 2.25 | 0.45 |
| 28:lb:92:PHE:HA | 28:lb:120:GLN:NE2 | 2.32 | 0.45 |
| 33:lg:36:PRO:HG2 | 33:lg:44:ARG:HB3 | 1.99 | 0.45 |
| 52:sa:429:G:N1 | 52:sa:432:A:OP2 | 2.49 | 0.45 |
| 52:sa:734:A:C2 | 52:sa:803:G:N1 | 2.78 | 0.45 |
| 52:sa:1595:G:H3' | 52:sa:1682:A:H61 | 1.82 | 0.45 |
| 54:sd:205:ASP:O | 54:sd:208:PRO:HD2 | 2.17 | 0.45 |
| 60:sj:106:PRO:HB2 | 60:sj:110:ARG:HG3 | 1.99 | 0.45 |
| 65:sp:34:ASP:OD1 | 65:sp:35:THR:N | 2.48 | 0.45 |
| 66:sq:56:VAL:O | 66:sq:60:LEU:HD13 | 2.17 | 0.45 |
| 1:lA:861:A:N6 | 1:lA:881:G:H1' | 2.31 | 0.45 |
| 1:lA:1347:A:N7 | 1:lA:1410:A:C6 | 2.85 | 0.45 |
| 1:lA:1844:U:H2' | 1:lA:1845:G:H8 | 1.82 | 0.45 |
| 1:lA:2112:C:H2' | 1:lA:2113:A:H8 | 1.81 | 0.45 |
| 1:lA:2180:C:H2' | 1:lA:2181:A:H8 | 1.82 | 0.45 |
| 1:lA:3161:U:H2' | 1:lA:3162:A:C8 | 2.52 | 0.45 |
| 3:lC:48:U:H3' | 7:lG:221:ARG:HG3 | 1.98 | 0.45 |
| 5:lE:242:ARG:HA | 5:lE:248:LEU:HD21 | 1.99 | 0.45 |
| 17:lQ:38:ARG:HB2 | 17:lQ:62:TYR:CZ | 2.52 | 0.45 |
| 27:la:62:GLN:HB3 | 27:la:64:VAL:HG13 | 1.98 | 0.45 |
| 28:lb:57:MET:HE2 | 28:lb:57:MET:HB3 | 1.85 | 0.45 |
| 43:lq:51:GLN:NE2 | 43:lq:53:LYS:O | 2.30 | 0.45 |
| 52:sa:341:G:H5' | 63:sm:78:HIS:HB2 | 1.98 | 0.45 |
| 52:sa:1196:A:H2' | 52:sa:1197:G:H8 | 1.82 | 0.45 |
| 52:sa:1209:A:N1 | 52:sa:1562:G:H5' | 2.32 | 0.45 |
| 52:sa:1363:C:H1' | 52:sa:1517:A:C5 | 2.52 | 0.45 |
| 52:sa:1709:A:H5' | 71:sv:88:TYR:CE2 | 2.52 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 55:se:123:ILE:HD13 | 55:se:166:ILE:HG21 | 1.99 | 0.45 |
| 56:sf:28:LYS:NZ | 56:sf:36:MET:SD | 2.89 | 0.45 |
| 70:su:27:VAL:O | 70:su:31:LEU:HG | 2.17 | 0.45 |
| 1:lA:760:A:H2' | 1:lA:761:A:C8 | 2.51 | 0.45 |
| 1:lA:1615:A:O2' | 32:lf:95:GLN:OE1 | 2.30 | 0.45 |
| 1:lA:2230:U:H5'' | 4:ID:242:ARG:O | 2.16 | 0.45 |
| 1:lA:2485:G:O2' | 1:lA:2487:U:H5'' | 2.17 | 0.45 |
| 3:lC:109:G:C6 | 3:lC:110:A:N1 | 2.85 | 0.45 |
| 12:lL:62:SER:HA | 12:lL:65:MET:SD | 2.56 | 0.45 |
| 19:lS:18:ALA:HB2 | 19:lS:29:VAL:HG21 | 1.99 | 0.45 |
| 24:lX:88:TRP:HH2 | 24:lX:98:PHE:CE1 | 2.35 | 0.45 |
| 52:sa:734:A:N1 | 52:sa:803:G:C6 | 2.85 | 0.45 |
| 52:sa:959:A:H4' | 52:sa:1933:G:N2 | 2.31 | 0.45 |
| 53:sc:180:ILE:HG23 | 53:sc:198:VAL:HG12 | 1.99 | 0.45 |
| 59:si:129:GLU:OE2 | 64:so:20:ARG:NH1 | 2.50 | 0.45 |
| 74:sy:39:LEU:HD22 | 74:sy:45:ALA:HB2 | 1.98 | 0.45 |
| 1:lA:353:G:H2' | 1:lA:354:U:C6 | 2.52 | 0.44 |
| 1:lA:914:G:H2' | 1:lA:915:U:C6 | 2.52 | 0.44 |
| 1:lA:2150:G:C2 | 1:lA:2176:G:C2 | 3.05 | 0.44 |
| 9:lI:75:ILE:HD12 | 9:lI:222:ILE:HD11 | 1.99 | 0.44 |
| 25:lY:46:LEU:HG | 25:lY:48:THR:HG23 | 1.98 | 0.44 |
| 40:ln:25:VAL:HG13 | 40:ln:68:GLU:HA | 1.99 | 0.44 |
| 45:sB:19:GLN:NE2 | 52:sa:924:A:N7 | 2.64 | 0.44 |
| 49:sI:14:A:N6 | 49:sI:46:G:H1 | 2.15 | 0.44 |
| 52:sa:1689:A:H2 | 52:sa:1756:U:H1' | 1.82 | 0.44 |
| 52:sa:1690:A:O2' | 52:sa:1691:A:H8 | 2.00 | 0.44 |
| 60:sj:107:ALA:HB3 | 60:sj:108:PRO:HD3 | 1.97 | 0.44 |
| 67:sr:102:LEU:HD23 | 67:sr:128:PHE:HB3 | 1.98 | 0.44 |
| 1:lA:43:C:H5'' | 14:lN:14:LYS:HG2 | 1.98 | 0.44 |
| 1:lA:226:U:H2' | 1:lA:227:A:C8 | 2.52 | 0.44 |
| 1:lA:857:U:H2' | 1:lA:858:A:H8 | 1.83 | 0.44 |
| 1:lA:1593:A:OP1 | 32:lf:75:ARG:NH2 | 2.51 | 0.44 |
| 1:lA:1627:G:H21 | 34:lh:6:THR:HG22 | 1.82 | 0.44 |
| 1:lA:1736:A:C6 | 41:lo:4:HIS:HB3 | 2.52 | 0.44 |
| 1:lA:1745:A:H2' | 1:lA:1746:G:C8 | 2.52 | 0.44 |
| 1:lA:1749:U:P | 21:lU:42:ARG:HH22 | 2.40 | 0.44 |
| 1:lA:2445:G:H2' | 1:lA:2446:G:C8 | 2.53 | 0.44 |
| 17:lQ:60:VAL:HG23 | 17:lQ:134:LEU:HB2 | 1.99 | 0.44 |
| 17:lQ:115:VAL:HG12 | 17:lQ:165:THR:HG21 | 1.97 | 0.44 |
| 25:lY:41:ILE:HG22 | 25:lY:42:ILE:HG13 | 1.99 | 0.44 |
| 28:lb:91:GLN:NE2 | 28:lb:91:GLN:O | 2.50 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 38:ll:2:THR:O | 38:ll:2:THR:OG1 | 2.31 | 0.44 |
| 52:sa:137:A:H2' | 52:sa:138:G:H8 | 1.81 | 0.44 |
| 52:sa:620:U:H2' | 52:sa:621:C:H6 | 1.82 | 0.44 |
| 54:sd:140:ILE:HG21 | 54:sd:202:ILE:HG12 | 1.99 | 0.44 |
| 55:se:111:LYS:HD3 | 55:se:111:LYS:HA | 1.72 | 0.44 |
| 56:sf:127:ARG:NH1 | 56:sf:135:PRO:HB2 | 2.32 | 0.44 |
| 66:sq:85:LEU:H | 66:sq:88:MET:HE3 | 1.82 | 0.44 |
| 74:sy:101:SER:O | 74:sy:101:SER:OG | 2.32 | 0.44 |
| 1:lA:592:A:H1' | 1:lA:756:A:H2 | 1.82 | 0.44 |
| 1:lA:960:G:O2' | 21:lU:131:GLN:OE1 | 2.34 | 0.44 |
| 1:lA:1544:U:O2' | 33:lg:96:GLU:OE1 | 2.36 | 0.44 |
| 1:lA:2217:A:N1 | 1:lA:3099:A:H2' | 2.33 | 0.44 |
| 3:lC:29:G:O2' | 7:lG:217:LYS:NZ | 2.36 | 0.44 |
| 6:lF:397:LEU:HD12 | 6:lF:397:LEU:HA | 1.70 | 0.44 |
| 7:lG:41:LYS:HA | 7:lG:41:LYS:HD2 | 1.67 | 0.44 |
| 17:lQ:10:LEU:HB2 | 37:lk:38:ILE:HD12 | 1.98 | 0.44 |
| 37:lk:68:LYS:HD3 | 37:lk:74:HIS:HA | 2.00 | 0.44 |
| 49:sI:24:G:H2' | 49:sI:25:C:C6 | 2.52 | 0.44 |
| 50:sJ:42:C:H2' | 50:sJ:43:G:H8 | 1.81 | 0.44 |
| 52:sa:1431:U:H4' | 72:sw:57:PRO:HG3 | 1.99 | 0.44 |
| 52:sa:1901:G:O2' | 74:sy:56:GLU:OE2 | 2.31 | 0.44 |
| 70:su:27:VAL:HG11 | 70:su:51:ILE:HD13 | 1.98 | 0.44 |
| 1:lA:324:A:C6 | 1:lA:336:A:N1 | 2.85 | 0.44 |
| 1:lA:746:A:H5' | 8:lH:41:LYS:HD3 | 1.99 | 0.44 |
| 1:lA:1089:C:O5' | 1:lA:1089:C:H6 | 2.00 | 0.44 |
| 1:lA:2767:A:H2' | 1:lA:2768:G:C8 | 2.52 | 0.44 |
| 1:lA:3149:A:H2' | 1:lA:3150:A:C8 | 2.53 | 0.44 |
| 1:lA:3360:U:C4 | 16:lP:117:ILE:HD11 | 2.52 | 0.44 |
| 1:lA:3375:U:H2' | 1:lA:3376:C:C6 | 2.52 | 0.44 |
| 12:lL:202:THR:O | 12:lL:202:THR:OG1 | 2.34 | 0.44 |
| 27:la:10:ASN:HB3 | 27:la:13:LYS:HB2 | 1.99 | 0.44 |
| 48:sE:21:CYS:SG | 48:sE:39:CYS:N | 2.77 | 0.44 |
| 50:sJ:65:G:H2' | 50:sJ:65:G:N3 | 2.33 | 0.44 |
| 52:sa:213:A:N6 | 52:sa:243:U:H4' | 2.32 | 0.44 |
| 52:sa:373:A:H2' | 52:sa:374:C:O4' | 2.17 | 0.44 |
| 52:sa:380:A:H2' | 52:sa:381:G:C8 | 2.53 | 0.44 |
| 52:sa:1021:G:H2' | 52:sa:1022:G:C8 | 2.52 | 0.44 |
| 66:sq:19:LYS:HD3 | 66:sq:23:GLU:OE1 | 2.17 | 0.44 |
| 1:lA:424:A:OP1 | 1:lA:437:A:H2 | 2.00 | 0.44 |
| 1:lA:429:G:N2 | 1:lA:432:A:OP2 | 2.42 | 0.44 |
| 1:lA:857:U:H2' | 1:lA:858:A:C8 | 2.53 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 1:1A:1407:U:H2' | 1:1A:1408:C:O4' | 2.18 | 0.44 |
| 1:1A:1414:A:H2' | 1:1A:1415:A:C8 | 2.48 | 0.44 |
| 1:1A:2358:U:OP1 | 1:1A:3116:G:O2' | 2.27 | 0.44 |
| 1:1A:3103:C:H2' | 1:1A:3104:G:C8 | 2.53 | 0.44 |
| 2:1B:90:A:H5' | 27:1a:22:ASN:HD22 | 1.83 | 0.44 |
| 5:1E:80:GLU:OE1 | 5:1E:315:TYR:OH | 2.35 | 0.44 |
| 19:1S:22:ASN:HB3 | 19:1S:25:MET:HB2 | 1.98 | 0.44 |
| 23:1W:68:VAL:C | 23:1W:69:LYS:HG2 | 2.43 | 0.44 |
| 42:1p:21:CYS:HB3 | 42:1p:24:CYS:HB2 | 2.00 | 0.44 |
| 52:sa:959:A:N3 | 52:sa:1922:U:O2' | 2.49 | 0.44 |
| 52:sa:1261:A:N6 | 52:sa:1262:G:O6 | 2.50 | 0.44 |
| 52:sa:1694:A:H5' | 68:ss:57:PRO:HG2 | 1.97 | 0.44 |
| 52:sa:1756:U:H2' | 52:sa:1757:C:C6 | 2.53 | 0.44 |
| 1:1A:199:G:N1 | 1:1A:311:A:OP2 | 2.49 | 0.44 |
| 1:1A:324:A:H2' | 1:1A:325:U:C6 | 2.53 | 0.44 |
| 1:1A:597:A:H4' | 8:1H:53:GLU:HG2 | 1.99 | 0.44 |
| 1:1A:709:U:HO2' | 1:1A:710:A:P | 2.40 | 0.44 |
| 1:1A:720:U:HO2' | 1:1A:721:A:P | 2.41 | 0.44 |
| 1:1A:2252:U:OP1 | 4:1D:54:ARG:NH2 | 2.44 | 0.44 |
| 1:1A:3147:U:N3 | 5:1E:144:SER:OG | 2.49 | 0.44 |
| 1:1A:3369:U:H2' | 1:1A:3370:A:H8 | 1.83 | 0.44 |
| 4:1D:143:ASN:OD1 | 4:1D:143:ASN:N | 2.50 | 0.44 |
| 6:1F:207:GLY:HA3 | 6:1F:228:GLY:O | 2.17 | 0.44 |
| 8:1H:93:GLY:O | 8:1H:94:PRO:C | 2.61 | 0.44 |
| 9:1I:87:ILE:HG23 | 9:1I:118:VAL:HG12 | 1.99 | 0.44 |
| 11:1K:65:ILE:HD13 | 11:1K:82:ALA:HB2 | 2.00 | 0.44 |
| 52:sa:170:G:C5 | 52:sa:171:A:H2 | 2.36 | 0.44 |
| 52:sa:323:A:H2' | 52:sa:324:G:H8 | 1.82 | 0.44 |
| 52:sa:379:G:H2' | 52:sa:380:A:C8 | 2.52 | 0.44 |
| 52:sa:592:U:H2' | 52:sa:593:A:H8 | 1.81 | 0.44 |
| 52:sa:1566:C:OP1 | 70:su:125:ARG:NH2 | 2.41 | 0.44 |
| 52:sa:1771:G:OP1 | 72:sw:79:HIS:NE2 | 2.46 | 0.44 |
| 1:1A:258:G:C8 | 6:1F:227:PRO:HG3 | 2.52 | 0.44 |
| 1:1A:737:A:O5' | 1:1A:738:G:N2 | 2.50 | 0.44 |
| 1:1A:913:U:C2 | 1:1A:914:G:C8 | 3.05 | 0.44 |
| 1:1A:1311:C:O2 | 20:1T:111:ARG:NH2 | 2.50 | 0.44 |
| 1:1A:2418:U:OP1 | 5:1E:233:LYS:NZ | 2.47 | 0.44 |
| 36:1j:42:PHE:HE1 | 36:1j:108:ILE:HD11 | 1.82 | 0.44 |
| 38:1l:44:MET:HE3 | 38:1l:44:MET:HB3 | 1.71 | 0.44 |
| 40:1n:10:GLN:H | 40:1n:10:GLN:CD | 2.26 | 0.44 |
| 52:sa:587:U:H5'' | 61:sk:40:LYS:HE2 | 2.00 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 52:sa:799:C:O2 | 67:sr:124:ARG:HD2 | 2.17 | 0.44 |
| 52:sa:1250:G:H22 | 52:sa:1283:U:H3 | 1.66 | 0.44 |
| 54:sd:207:ASP:HB3 | 54:sd:208:PRO:HD3 | 1.98 | 0.44 |
| 1:lA:637:U:H1' | 1:lA:638:A:C8 | 2.52 | 0.44 |
| 1:lA:996:C:HO2' | 1:lA:999:G:HO2' | 1.62 | 0.44 |
| 1:lA:1207:C:H2' | 1:lA:1208:U:O4' | 2.18 | 0.44 |
| 1:lA:1426:A:N7 | 1:lA:3000:C:O2' | 2.50 | 0.44 |
| 1:lA:1733:U:C4 | 1:lA:1734:G:N7 | 2.86 | 0.44 |
| 1:lA:1877:A:H2' | 1:lA:1878:A:H8 | 1.82 | 0.44 |
| 1:lA:2865:A:H2' | 1:lA:2866:C:C2 | 2.52 | 0.44 |
| 1:lA:2868:U:H3 | 1:lA:2886:U:H3 | 1.66 | 0.44 |
| 1:lA:3134:U:O2' | 1:lA:3438:G:N7 | 2.51 | 0.44 |
| 1:lA:3476:C:H5'' | 1:lA:3477:U:N1 | 2.33 | 0.44 |
| 4:lD:196:TRP:HB3 | 4:lD:197:PRO:HD3 | 2.00 | 0.44 |
| 6:lF:162:ILE:HD11 | 6:lF:171:LEU:HD22 | 2.00 | 0.44 |
| 17:lQ:65:ARG:HB2 | 17:lQ:129:PHE:HD1 | 1.82 | 0.44 |
| 19:lS:66:LEU:HD23 | 19:lS:136:ILE:HD12 | 2.00 | 0.44 |
| 21:lU:119:MET:HE3 | 21:lU:119:MET:HB2 | 1.78 | 0.44 |
| 22:lV:148:LYS:HD3 | 22:lV:148:LYS:HA | 1.60 | 0.44 |
| 25:lY:112:MET:HE1 | 25:lY:121:PHE:CD2 | 2.52 | 0.44 |
| 33:lg:20:ARG:HD3 | 33:lg:29:VAL:HG12 | 1.98 | 0.44 |
| 36:lj:94:ALA:HA | 36:lj:97:GLN:HG3 | 2.00 | 0.44 |
| 52:sa:150:G:C2 | 52:sa:152:A:H5'' | 2.53 | 0.44 |
| 52:sa:1579:A:H4' | 52:sa:1580:U:H5' | 1.99 | 0.44 |
| 52:sa:1752:A:H2' | 52:sa:1753:A:O4' | 2.18 | 0.44 |
| 56:sf:200:PRO:HB2 | 63:sm:39:MET:HG2 | 2.00 | 0.44 |
| 60:sj:6:ASP:HB3 | 60:sj:28:THR:HG21 | 1.99 | 0.44 |
| 1:lA:474:A:H2' | 1:lA:475:A:H8 | 1.83 | 0.44 |
| 1:lA:496:A:H2' | 1:lA:497:U:C6 | 2.53 | 0.44 |
| 1:lA:619:A:P | 1:lA:619:A:H3' | 2.58 | 0.44 |
| 1:lA:686:A:H5' | 9:lI:134:LYS:HD3 | 2.00 | 0.44 |
| 1:lA:1293:U:OP1 | 9:lI:199:LYS:HG2 | 2.18 | 0.44 |
| 1:lA:2195:U:H2' | 1:lA:2196:A:O4' | 2.18 | 0.44 |
| 3:lC:7:A:OP2 | 7:lG:30:TYR:OH | 2.28 | 0.44 |
| 4:lD:57:PRO:HG2 | 4:lD:78:CYS:HB3 | 1.99 | 0.44 |
| 12:lL:191:ILE:O | 12:lL:198:LYS:N | 2.51 | 0.44 |
| 16:lP:29:GLU:HG3 | 16:lP:30:ILE:N | 2.32 | 0.44 |
| 17:lQ:8:ASN:HB2 | 17:lQ:46:GLU:HG3 | 2.00 | 0.44 |
| 21:lU:76:THR:O | 21:lU:76:THR:OG1 | 2.31 | 0.44 |
| 25:lY:94:ARG:HH21 | 25:lY:98:LEU:HD12 | 1.83 | 0.44 |
| 39:lm:36:LYS:HE2 | 39:lm:48:ARG:HH11 | 1.82 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|-------------------|--------------------------|-------------------|
| 51:sk:4:A:OP2 | 52:sa:1176:G:N1 | 2.36 | 0.44 |
| 52:sa:1107:A:N6 | 52:sa:1108:G:O6 | 2.51 | 0.44 |
| 52:sa:1383:C:H2' | 52:sa:1384:U:C6 | 2.52 | 0.44 |
| 52:sa:1733:U:O4 | 52:sa:1734:G:N1 | 2.51 | 0.44 |
| 61:sk:57:LYS:HE3 | 61:sk:57:LYS:HB2 | 1.72 | 0.44 |
| 71:sv:37:THR:O | 71:sv:37:THR:OG1 | 2.34 | 0.44 |
| 1:lA:247:A:H5' | 1:lA:266:A:O2' | 2.18 | 0.43 |
| 1:lA:661:G:H2' | 1:lA:662:U:C6 | 2.53 | 0.43 |
| 1:lA:1222:U:O2' | 22:IV:136:ARG:O | 2.25 | 0.43 |
| 1:lA:1269:U:C4 | 1:lA:1488:A:H2 | 2.36 | 0.43 |
| 1:lA:1471:A:H2' | 1:lA:1472:U:C6 | 2.53 | 0.43 |
| 1:lA:1519:A:H5'' | 1:lA:1521:A:N6 | 2.33 | 0.43 |
| 1:lA:2214:A:O2' | 38:II:2:THR:N | 2.44 | 0.43 |
| 1:lA:2381:G:N2 | 1:lA:2381:G:OP2 | 2.51 | 0.43 |
| 1:lA:2883:A:H2' | 1:lA:2884:U:C6 | 2.53 | 0.43 |
| 1:lA:3391:U:H2' | 1:lA:3392:G:C8 | 2.53 | 0.43 |
| 9:II:166:VAL:HG13 | 9:II:171:ASP:HB3 | 1.99 | 0.43 |
| 17:IQ:46:GLU:OE1 | 17:IQ:47:ARG:N | 2.51 | 0.43 |
| 35:II:1:MET:HB3 | 35:II:5:ASP:HB3 | 2.00 | 0.43 |
| 52:sa:136:A:H4' | 52:sa:137:A:H5' | 1.99 | 0.43 |
| 52:sa:256:A:N6 | 52:sa:257:C:H41 | 2.16 | 0.43 |
| 56:sf:9:ARG:HD2 | 56:sf:9:ARG:HA | 1.87 | 0.43 |
| 1:lA:213:A:N6 | 14:IN:131:THR:OG1 | 2.40 | 0.43 |
| 1:lA:1461:G:H2' | 1:lA:1462:A:H8 | 1.83 | 0.43 |
| 1:lA:2034:G:O2' | 41:IO:3:GLY:O | 2.36 | 0.43 |
| 1:lA:2431:G:H4' | 18:IR:139:PHE:CE1 | 2.53 | 0.43 |
| 2:IB:61:A:C4 | 2:IB:63:A:N6 | 2.86 | 0.43 |
| 5:IE:322:PHE:HE2 | 5:IE:324:MET:HE3 | 1.83 | 0.43 |
| 6:IF:295:VAL:HG22 | 19:IS:34:PHE:CE1 | 2.53 | 0.43 |
| 14:IN:174:LYS:HE2 | 14:IN:174:LYS:HB2 | 1.65 | 0.43 |
| 24:IX:140:VAL:HG11 | 26:IZ:24:VAL:HG21 | 2.00 | 0.43 |
| 52:sa:57:U:H3 | 52:sa:88:G:H1 | 1.66 | 0.43 |
| 52:sa:379:G:H2' | 52:sa:380:A:H8 | 1.83 | 0.43 |
| 52:sa:1510:A:H2' | 52:sa:1511:A:H8 | 1.83 | 0.43 |
| 56:sf:235:ASN:OD1 | 56:sf:235:ASN:N | 2.44 | 0.43 |
| 71:sv:89:ASN:N | 71:sv:89:ASN:OD1 | 2.50 | 0.43 |
| 1:lA:330:A:P | 43:IQ:43:ARG:HH22 | 2.40 | 0.43 |
| 1:lA:504:U:H2' | 1:lA:505:A:C8 | 2.53 | 0.43 |
| 1:lA:965:A:H2' | 1:lA:966:A:O4' | 2.17 | 0.43 |
| 1:lA:1022:U:OP2 | 38:II:30:GLN:NE2 | 2.51 | 0.43 |
| 1:lA:1622:A:H2' | 1:lA:1622:A:N3 | 2.33 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:1A:2220:A:H1' | 1:1A:2357:A:H61 | 1.82 | 0.43 |
| 1:1A:2678:G:OP1 | 4:1D:2:GLY:HA3 | 2.19 | 0.43 |
| 1:1A:3442:G:OP1 | 5:1E:177:LYS:HD3 | 2.17 | 0.43 |
| 5:1E:217:ILE:HD12 | 5:1E:339:LEU:HD22 | 2.01 | 0.43 |
| 6:1F:292:LYS:HB2 | 6:1F:292:LYS:HE3 | 1.85 | 0.43 |
| 7:1G:89:LYS:HB2 | 7:1G:89:LYS:HE2 | 1.71 | 0.43 |
| 14:1N:49:ILE:HA | 14:1N:231:LEU:O | 2.18 | 0.43 |
| 15:1O:139:LEU:HD23 | 15:1O:139:LEU:HA | 1.78 | 0.43 |
| 17:1Q:117:ASN:OD1 | 17:1Q:166:SER:HB3 | 2.18 | 0.43 |
| 18:1R:18:ARG:NH2 | 18:1R:147:GLU:HG2 | 2.33 | 0.43 |
| 24:1X:123:LYS:HG3 | 24:1X:139:ILE:HG22 | 1.98 | 0.43 |
| 45:sB:4:LYS:HD3 | 45:sB:5:ARG:NH2 | 2.33 | 0.43 |
| 52:sa:1302:G:H5'' | 54:sd:153:SER:HB2 | 2.01 | 0.43 |
| 52:sa:1305:C:O2' | 72:sw:68:THR:HG22 | 2.18 | 0.43 |
| 55:se:114:SER:O | 55:se:114:SER:OG | 2.34 | 0.43 |
| 66:sq:53:GLU:O | 66:sq:57:LYS:HG2 | 2.19 | 0.43 |
| 1:1A:213:A:C5' | 14:1N:133:LYS:HB2 | 2.48 | 0.43 |
| 1:1A:619:A:N6 | 16:1P:77:CYS:HB3 | 2.34 | 0.43 |
| 1:1A:928:G:N1 | 1:1A:1051:U:O4 | 2.51 | 0.43 |
| 1:1A:1236:G:H2' | 1:1A:1237:A:C8 | 2.52 | 0.43 |
| 1:1A:1275:A:H2 | 1:1A:2455:U:O2 | 2.01 | 0.43 |
| 1:1A:1289:G:C6 | 1:1A:1461:G:N1 | 2.87 | 0.43 |
| 1:1A:1502:A:N6 | 19:1S:15:SER:OG | 2.40 | 0.43 |
| 1:1A:3287:U:H2' | 1:1A:3288:U:C6 | 2.54 | 0.43 |
| 5:1E:20:LYS:HE3 | 5:1E:20:LYS:HB3 | 1.85 | 0.43 |
| 6:1F:35:ASP:OD1 | 6:1F:36:VAL:N | 2.51 | 0.43 |
| 32:1f:118:ILE:HG22 | 32:1f:121:GLU:O | 2.17 | 0.43 |
| 52:sa:1508:A:OP1 | 69:st:60:ARG:NH1 | 2.52 | 0.43 |
| 52:sa:1545:G:H2' | 52:sa:1546:C:C6 | 2.53 | 0.43 |
| 54:sd:172:ILE:HD11 | 54:sd:177:ALA:HB3 | 2.00 | 0.43 |
| 71:sv:86:LYS:O | 71:sv:87:HIS:ND1 | 2.51 | 0.43 |
| 71:sv:129:LYS:HZ2 | 71:sv:129:LYS:HG2 | 1.63 | 0.43 |
| 1:1A:215:A:H2' | 1:1A:216:A:O4' | 2.18 | 0.43 |
| 1:1A:227:A:H2' | 1:1A:228:A:C8 | 2.53 | 0.43 |
| 1:1A:322:U:H2' | 1:1A:323:U:H6 | 1.82 | 0.43 |
| 1:1A:343:A:H62 | 17:1Q:12:ARG:NH1 | 2.17 | 0.43 |
| 1:1A:467:A:C2 | 1:1A:2439:A:H4' | 2.53 | 0.43 |
| 1:1A:551:A:H2' | 1:1A:552:G:O4' | 2.18 | 0.43 |
| 1:1A:1093:G:OP1 | 19:1S:19:ARG:NH2 | 2.45 | 0.43 |
| 1:1A:2469:G:OP1 | 1:1A:2469:G:O4' | 2.37 | 0.43 |
| 1:1A:3203:A:H4' | 5:1E:365:LYS:HD2 | 2.00 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:1A:3432:A:HO2' | 1:1A:3433:U:P | 2.41 | 0.43 |
| 1:1A:3482:C:H2' | 1:1A:3483:A:O4' | 2.18 | 0.43 |
| 2:1B:116:U:H2' | 2:1B:117:A:C8 | 2.53 | 0.43 |
| 3:1C:2:G:H2' | 3:1C:3:G:H8 | 1.84 | 0.43 |
| 6:1F:431:PHE:CZ | 20:1T:24:ARG:HG3 | 2.53 | 0.43 |
| 8:1H:61:PRO:O | 8:1H:116:LYS:NZ | 2.52 | 0.43 |
| 8:1H:176:GLU:OE1 | 8:1H:176:GLU:N | 2.39 | 0.43 |
| 12:1L:30:ARG:NH2 | 12:1L:66:GLU:HG2 | 2.33 | 0.43 |
| 14:1N:124:LEU:HD21 | 35:1i:109:PHE:HE1 | 1.82 | 0.43 |
| 42:1p:20:ILE:HD11 | 42:1p:49:LYS:HD2 | 1.99 | 0.43 |
| 47:sD:10:VAL:HG11 | 57:sg:124:ARG:HH12 | 1.84 | 0.43 |
| 52:sa:456:G:H2' | 52:sa:457:G:C8 | 2.49 | 0.43 |
| 52:sa:1264:U:H2' | 52:sa:1265:U:O4' | 2.19 | 0.43 |
| 52:sa:1266:A:H8 | 66:sq:78:HIS:CG | 2.36 | 0.43 |
| 52:sa:1548:C:H2' | 52:sa:1549:U:C6 | 2.54 | 0.43 |
| 52:sa:1584:G:H2' | 52:sa:1585:U:C6 | 2.53 | 0.43 |
| 57:sg:51:LEU:HD23 | 57:sg:51:LEU:HA | 1.90 | 0.43 |
| 60:sj:105:ASP:OD1 | 60:sj:105:ASP:N | 2.43 | 0.43 |
| 69:st:32:LYS:O | 69:st:47:ARG:NH1 | 2.51 | 0.43 |
| 1:1A:206:U:O2 | 1:1A:303:G:N2 | 2.35 | 0.43 |
| 1:1A:213:A:H5' | 14:1N:133:LYS:HB2 | 2.00 | 0.43 |
| 1:1A:265:A:H1' | 1:1A:266:A:H2 | 1.83 | 0.43 |
| 1:1A:720:U:H2' | 1:1A:721:A:C8 | 2.53 | 0.43 |
| 1:1A:1064:C:O2' | 1:1A:1540:A:H1' | 2.18 | 0.43 |
| 1:1A:1430:G:O6 | 1:1A:2442:C:O2' | 2.32 | 0.43 |
| 1:1A:2909:A:C6 | 1:1A:2910:G:N7 | 2.87 | 0.43 |
| 2:1B:68:A:H2' | 2:1B:69:U:C6 | 2.53 | 0.43 |
| 2:1B:80:A:O2' | 2:1B:81:A:OP1 | 2.33 | 0.43 |
| 4:1D:234:LYS:HE3 | 4:1D:234:LYS:HB2 | 1.78 | 0.43 |
| 5:1E:122:TRP:CZ2 | 5:1E:127:LYS:HG2 | 2.53 | 0.43 |
| 5:1E:347:THR:O | 5:1E:352:GLN:NE2 | 2.52 | 0.43 |
| 17:1Q:4:TYR:HE2 | 17:1Q:46:GLU:HA | 1.84 | 0.43 |
| 17:1Q:121:ILE:HD13 | 17:1Q:121:ILE:HA | 1.77 | 0.43 |
| 25:1Y:38:GLU:OE1 | 25:1Y:38:GLU:N | 2.47 | 0.43 |
| 32:1f:11:VAL:O | 32:1f:15:ARG:N | 2.52 | 0.43 |
| 33:1g:41:ASN:HB3 | 33:1g:44:ARG:HG2 | 2.00 | 0.43 |
| 34:1h:99:ASN:HA | 34:1h:102:LYS:HB3 | 2.00 | 0.43 |
| 37:1k:48:GLU:HG2 | 37:1k:84:ILE:HD11 | 1.99 | 0.43 |
| 50:sJ:48:U:H5' | 50:sJ:49:C:OP1 | 2.19 | 0.43 |
| 52:sa:1143:U:C2 | 52:sa:1144:G:C8 | 3.07 | 0.43 |
| 75:sa:5101:PAR:O44 | 75:sa:5101:PAR:N64 | 2.51 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 53:sc:221:ARG:NE | 73:sx:25:ASP:OD2 | 2.51 | 0.43 |
| 56:sf:177:ASN:HA | 56:sf:229:GLY:H | 1.82 | 0.43 |
| 72:sw:67:LYS:HE3 | 72:sw:78:ASP:HB3 | 2.01 | 0.43 |
| 1:lA:443:A:H4' | 1:lA:444:A:OP1 | 2.19 | 0.43 |
| 1:lA:503:A:H2' | 1:lA:504:U:C6 | 2.54 | 0.43 |
| 1:lA:549:U:O2' | 1:lA:550:A:H8 | 2.02 | 0.43 |
| 1:lA:687:U:C2 | 1:lA:688:A:C8 | 3.06 | 0.43 |
| 1:lA:1562:A:H2' | 29:lc:3:THR:HG21 | 2.01 | 0.43 |
| 1:lA:1751:A:N3 | 1:lA:1751:A:H2' | 2.34 | 0.43 |
| 1:lA:2726:A:N7 | 1:lA:2728:G:C8 | 2.87 | 0.43 |
| 5:lE:34:LYS:HE3 | 5:lE:34:LYS:HB2 | 1.85 | 0.43 |
| 13:lM:164:LYS:HD3 | 13:lM:171:VAL:HG22 | 2.01 | 0.43 |
| 15:lO:173:GLN:OE1 | 15:lO:173:GLN:HA | 2.18 | 0.43 |
| 21:lU:133:LYS:HB2 | 21:lU:133:LYS:HE3 | 1.82 | 0.43 |
| 25:lY:29:LEU:O | 35:li:54:ARG:NH2 | 2.42 | 0.43 |
| 52:sa:547:G:OP2 | 52:sa:548:C:O2' | 2.30 | 0.43 |
| 52:sa:1170:U:C2 | 52:sa:1171:U:C5 | 3.07 | 0.43 |
| 52:sa:1256:A:O2' | 62:sl:2:ARG:NH2 | 2.45 | 0.43 |
| 52:sa:1694:A:OP1 | 68:ss:57:PRO:HD2 | 2.19 | 0.43 |
| 52:sa:1932:U:OP2 | 65:sp:142:ARG:NH2 | 2.52 | 0.43 |
| 53:sc:128:LYS:HB3 | 53:sc:128:LYS:HE2 | 1.68 | 0.43 |
| 56:sf:63:VAL:HG11 | 56:sf:78:THR:HA | 2.01 | 0.43 |
| 57:sg:83:LYS:HE3 | 57:sg:83:LYS:HB2 | 1.76 | 0.43 |
| 1:lA:747:C:H1' | 8:lH:51:ASN:HD22 | 1.84 | 0.43 |
| 1:lA:814:G:H2' | 1:lA:815:A:O4' | 2.19 | 0.43 |
| 1:lA:817:A:O2' | 1:lA:818:A:O4' | 2.36 | 0.43 |
| 1:lA:822:C:H2' | 1:lA:823:A:C8 | 2.54 | 0.43 |
| 1:lA:1444:U:O4 | 1:lA:1445:U:O4 | 2.36 | 0.43 |
| 1:lA:1856:G:H2' | 1:lA:1857:A:H8 | 1.84 | 0.43 |
| 1:lA:1857:A:OP1 | 23:lW:86:LYS:HB2 | 2.19 | 0.43 |
| 1:lA:1889:U:H2' | 1:lA:1890:U:C6 | 2.53 | 0.43 |
| 1:lA:3331:C:O2 | 32:lf:8:ASN:ND2 | 2.52 | 0.43 |
| 1:lA:3483:A:H2' | 1:lA:3484:A:C2 | 2.54 | 0.43 |
| 1:lA:3486:A:H8 | 1:lA:3486:A:O5' | 2.01 | 0.43 |
| 3:lC:27:G:H2' | 3:lC:28:U:H5' | 2.01 | 0.43 |
| 7:lG:91:GLY:O | 7:lG:94:ASN:ND2 | 2.48 | 0.43 |
| 12:lL:39:ARG:H | 12:lL:39:ARG:HG3 | 1.67 | 0.43 |
| 32:lf:80:LYS:HD2 | 32:lf:80:LYS:HA | 1.78 | 0.43 |
| 48:sE:47:ALA:HB1 | 48:sE:52:PHE:HB2 | 2.01 | 0.43 |
| 52:sa:338:C:H2' | 52:sa:339:A:H8 | 1.82 | 0.43 |
| 52:sa:1282:U:H5 | 62:sl:5:THR:HG22 | 1.84 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 52:sa:1356:A:H61 | 54:sd:175:GLY:HA2 | 1.83 | 0.43 |
| 58:sh:102:VAL:HG13 | 58:sh:106:ILE:HG13 | 2.00 | 0.43 |
| 1:lA:233:A:N3 | 1:lA:254:C:O2' | 2.41 | 0.43 |
| 1:lA:865:U:H2' | 1:lA:866:G:C8 | 2.54 | 0.43 |
| 1:lA:1403:C:H2' | 1:lA:1404:C:C6 | 2.53 | 0.43 |
| 1:lA:1558:C:H2' | 1:lA:1559:C:C6 | 2.54 | 0.43 |
| 1:lA:1636:U:H5 | 1:lA:2036:A:N1 | 2.16 | 0.43 |
| 1:lA:2429:G:H5'' | 18:lR:86:LYS:HB2 | 2.00 | 0.43 |
| 2:lB:30:U:C2 | 2:lB:31:C:C5 | 3.07 | 0.43 |
| 5:lE:19:ARG:HB3 | 5:lE:275:PHE:CZ | 2.54 | 0.43 |
| 6:lF:162:ILE:CD1 | 6:lF:171:LEU:HD22 | 2.48 | 0.43 |
| 6:lF:288:VAL:O | 6:lF:290:ASP:N | 2.51 | 0.43 |
| 6:lF:358:MET:HA | 6:lF:361:ILE:HD12 | 2.00 | 0.43 |
| 6:lF:404:THR:OG1 | 6:lF:405:ALA:N | 2.51 | 0.43 |
| 7:lG:50:ARG:NH2 | 7:lG:148:ASP:OD2 | 2.30 | 0.43 |
| 8:lH:85:LYS:HA | 8:lH:85:LYS:HD2 | 1.87 | 0.43 |
| 11:lK:44:VAL:HG12 | 11:lK:67:MET:HB2 | 2.01 | 0.43 |
| 14:lN:153:LYS:HA | 14:lN:153:LYS:HD3 | 1.84 | 0.43 |
| 29:lc:103:ASP:OD1 | 29:lc:106:LYS:HG2 | 2.19 | 0.43 |
| 45:sB:23:CYS:HB3 | 45:sB:28:ARG:H | 1.84 | 0.43 |
| 52:sa:866:G:C2' | 52:sa:867:A:H8 | 2.31 | 0.43 |
| 52:sa:925:U:H2' | 52:sa:926:U:C6 | 2.54 | 0.43 |
| 52:sa:1195:G:C2 | 52:sa:1196:A:C8 | 3.07 | 0.43 |
| 52:sa:1735:A:H2' | 52:sa:1736:A:H8 | 1.84 | 0.43 |
| 52:sa:1843:U:H2' | 52:sa:1844:A:C8 | 2.54 | 0.43 |
| 54:sd:193:MET:HE2 | 54:sd:193:MET:HA | 2.01 | 0.43 |
| 64:so:98:VAL:HG11 | 64:so:115:LEU:HB2 | 2.01 | 0.43 |
| 66:sq:106:ILE:HA | 66:sq:110:MET:HE2 | 2.00 | 0.43 |
| 1:lA:421:A:H2 | 1:lA:439:U:O2 | 2.02 | 0.43 |
| 1:lA:424:A:OP2 | 1:lA:438:G:H1' | 2.18 | 0.43 |
| 1:lA:2059:A:O2' | 1:lA:2060:A:OP2 | 2.34 | 0.43 |
| 6:lF:150:ILE:HG22 | 6:lF:151:PRO:HD3 | 2.01 | 0.43 |
| 16:lP:29:GLU:HB3 | 16:lP:37:LEU:HB3 | 2.01 | 0.43 |
| 25:lY:32:ASN:OD1 | 25:lY:32:ASN:N | 2.52 | 0.43 |
| 28:lb:4:ILE:HD11 | 31:le:38:ILE:HG22 | 2.01 | 0.43 |
| 39:lm:33:GLN:HE21 | 39:lm:49:ARG:NH2 | 2.16 | 0.43 |
| 45:sB:87:ARG:NH2 | 45:sB:94:ASN:O | 2.52 | 0.43 |
| 47:sD:49:PRO:HB3 | 52:sa:1781:C:OP1 | 2.19 | 0.43 |
| 52:sa:553:C:N4 | 52:sa:554:U:O4 | 2.52 | 0.43 |
| 52:sa:889:U:H2' | 52:sa:890:C:C6 | 2.54 | 0.43 |
| 52:sa:1677:A:H5' | 52:sa:1678:A:OP1 | 2.18 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 53:sc:66:LYS:HD2 | 53:sc:66:LYS:HA | 1.67 | 0.43 |
| 1:lA:411:A:H4' | 6:lF:86:ARG:HD3 | 2.01 | 0.42 |
| 1:lA:1945:A:N6 | 1:lA:1956:G:O6 | 2.52 | 0.42 |
| 1:lA:2111:A:H2' | 1:lA:2112:C:H6 | 1.84 | 0.42 |
| 1:lA:2184:A:H2 | 1:lA:3471:A:N3 | 2.16 | 0.42 |
| 1:lA:2427:U:H2' | 1:lA:2428:A:C8 | 2.53 | 0.42 |
| 1:lA:2461:G:O2' | 1:lA:2462:G:P | 2.77 | 0.42 |
| 1:lA:2732:A:H2' | 1:lA:2733:A:C8 | 2.54 | 0.42 |
| 1:lA:2937:G:N2 | 50:sJ:77:A:O2' | 2.52 | 0.42 |
| 6:lF:271:LYS:HE3 | 6:lF:271:LYS:HB2 | 1.81 | 0.42 |
| 14:lN:143:GLU:OE1 | 14:lN:143:GLU:N | 2.43 | 0.42 |
| 18:lR:16:GLN:HG2 | 18:lR:149:ILE:HG12 | 2.01 | 0.42 |
| 20:lT:73:ILE:HG21 | 20:lT:102:MET:SD | 2.59 | 0.42 |
| 23:lW:36:ASN:OD1 | 23:lW:36:ASN:N | 2.51 | 0.42 |
| 24:lX:14:PHE:HE2 | 24:lX:91:LYS:HG2 | 1.83 | 0.42 |
| 26:lZ:24:VAL:HG22 | 26:lZ:30:SER:HB2 | 2.01 | 0.42 |
| 27:la:83:LEU:HD12 | 27:la:83:LEU:HA | 1.88 | 0.42 |
| 28:lb:12:ILE:HD13 | 28:lb:137:LEU:HD22 | 2.01 | 0.42 |
| 28:lb:122:ALA:HB1 | 28:lb:126:ARG:HH21 | 1.84 | 0.42 |
| 32:lf:5:LYS:HD2 | 32:lf:5:LYS:HA | 1.82 | 0.42 |
| 52:sa:122:G:O2' | 56:sf:144:THR:N | 2.46 | 0.42 |
| 52:sa:122:G:C6 | 52:sa:291:A:C6 | 3.07 | 0.42 |
| 52:sa:1745:U:H1' | 68:ss:154:GLN:HG2 | 2.01 | 0.42 |
| 1:lA:11:A:H2' | 1:lA:12:U:H6 | 1.84 | 0.42 |
| 1:lA:221:U:H2' | 1:lA:222:A:H8 | 1.84 | 0.42 |
| 1:lA:463:U:O2' | 1:lA:464:G:H5' | 2.19 | 0.42 |
| 1:lA:560:U:O2' | 1:lA:561:G:H5' | 2.19 | 0.42 |
| 1:lA:686:A:H2' | 1:lA:687:U:H5'' | 2.02 | 0.42 |
| 1:lA:699:A:H5'' | 1:lA:746:A:N6 | 2.33 | 0.42 |
| 1:lA:715:A:C8 | 1:lA:727:U:H5'' | 2.54 | 0.42 |
| 1:lA:1285:C:P | 19:lS:2:ALA:HB2 | 2.58 | 0.42 |
| 1:lA:1572:U:H1' | 6:lF:96:THR:HA | 2.01 | 0.42 |
| 2:lB:23:U:O2' | 2:lB:24:U:H5' | 2.19 | 0.42 |
| 2:lB:31:C:H5'' | 14:lN:25:LYS:HG2 | 2.01 | 0.42 |
| 4:lD:27:ALA:HB2 | 4:lD:58:LEU:HD21 | 2.02 | 0.42 |
| 20:lT:73:ILE:HG12 | 20:lT:122:ILE:HD12 | 2.02 | 0.42 |
| 52:sa:29:G:H22 | 52:sa:541:U:H4' | 1.84 | 0.42 |
| 52:sa:1586:U:H2' | 52:sa:1587:A:C8 | 2.54 | 0.42 |
| 52:sa:1716:A:H2' | 52:sa:1717:U:C6 | 2.54 | 0.42 |
| 52:sa:1724:U:OP1 | 70:su:121:HIS:ND1 | 2.43 | 0.42 |
| 56:sf:196:ARG:HA | 56:sf:206:VAL:HA | 2.00 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 57:sg:20:LEU:HD21 | 57:sg:71:VAL:HG21 | 2.02 | 0.42 |
| 60:sj:11:ARG:O | 63:sm:133:LYS:NZ | 2.49 | 0.42 |
| 66:sq:29:GLU:O | 66:sq:33:VAL:HG13 | 2.18 | 0.42 |
| 66:sq:111:LEU:HB3 | 66:sq:112:GLY:H | 1.73 | 0.42 |
| 1:lA:517:G:C6 | 1:lA:556:A:C6 | 3.07 | 0.42 |
| 1:lA:1195:G:N3 | 30:ld:46:GLN:HB2 | 2.34 | 0.42 |
| 1:lA:1562:A:OP2 | 29:lc:3:THR:HG22 | 2.19 | 0.42 |
| 1:lA:1886:A:H2' | 1:lA:1887:A:C8 | 2.55 | 0.42 |
| 1:lA:2079:A:H3' | 1:lA:2079:A:N3 | 2.34 | 0.42 |
| 1:lA:2382:C:H4' | 1:lA:2383:G:H5' | 2.01 | 0.42 |
| 1:lA:3321:U:H2' | 1:lA:3322:A:O4' | 2.20 | 0.42 |
| 2:lB:81:A:O2' | 2:lB:82:A:OP1 | 2.37 | 0.42 |
| 6:lF:106:LYS:O | 6:lF:109:ARG:HG3 | 2.19 | 0.42 |
| 11:lK:174:ARG:HA | 11:lK:174:ARG:HD3 | 1.83 | 0.42 |
| 12:lL:4:ARG:N | 12:lL:123:TYR:OH | 2.44 | 0.42 |
| 13:lM:35:LYS:HA | 13:lM:35:LYS:HD2 | 1.66 | 0.42 |
| 14:lN:51:SER:O | 14:lN:53:LYS:HE3 | 2.19 | 0.42 |
| 20:lT:77:TYR:HD2 | 20:lT:112:THR:HG21 | 1.84 | 0.42 |
| 39:lm:7:LYS:O | 39:lm:27:LYS:NZ | 2.42 | 0.42 |
| 52:sa:555:G:H2' | 52:sa:556:G:C8 | 2.55 | 0.42 |
| 52:sa:903:A:H2' | 52:sa:904:A:C8 | 2.54 | 0.42 |
| 52:sa:919:A:H2' | 52:sa:920:A:C8 | 2.54 | 0.42 |
| 52:sa:1311:U:O2' | 52:sa:1312:A:O5' | 2.37 | 0.42 |
| 55:se:80:LYS:HE2 | 55:se:81:PHE:CE1 | 2.54 | 0.42 |
| 60:sj:44:HIS:ND1 | 60:sj:58:LEU:HD11 | 2.34 | 0.42 |
| 68:ss:54:ILE:HG22 | 68:ss:60:ARG:HG3 | 2.01 | 0.42 |
| 71:sv:9:ASP:N | 71:sv:9:ASP:OD1 | 2.52 | 0.42 |
| 1:lA:38:C:O2' | 1:lA:39:A:H5' | 2.19 | 0.42 |
| 1:lA:317:U:O2' | 1:lA:364:A:H1' | 2.19 | 0.42 |
| 1:lA:458:U:H5'' | 18:lR:37:ARG:NH2 | 2.35 | 0.42 |
| 1:lA:745:A:O2' | 1:lA:746:A:N3 | 2.48 | 0.42 |
| 1:lA:1121:A:N1 | 1:lA:1168:C:O2' | 2.41 | 0.42 |
| 1:lA:1620:U:H2' | 1:lA:1621:G:H5' | 2.02 | 0.42 |
| 6:lF:2:SER:O | 6:lF:2:SER:OG | 2.33 | 0.42 |
| 6:lF:358:MET:HA | 6:lF:358:MET:HE2 | 2.01 | 0.42 |
| 7:lG:219:PHE:O | 7:lG:223:ILE:HG13 | 2.19 | 0.42 |
| 12:lL:24:ARG:HB2 | 12:lL:24:ARG:HH11 | 1.84 | 0.42 |
| 17:lQ:45:PRO:HA | 17:lQ:48:ALA:HB3 | 2.00 | 0.42 |
| 47:sD:47:LYS:HD3 | 57:sg:147:ARG:HE | 1.85 | 0.42 |
| 52:sa:155:A:H2' | 52:sa:156:U:O4' | 2.19 | 0.42 |
| 52:sa:1111:A:H2' | 52:sa:1112:A:C8 | 2.55 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 52:sa:1542:G:H4' | 52:sa:1543:A:H5' | 2.00 | 0.42 |
| 53:sc:127:ILE:O | 53:sc:131:MET:HG2 | 2.19 | 0.42 |
| 60:sj:38:LEU:HD23 | 60:sj:96:LEU:HD21 | 2.01 | 0.42 |
| 68:ss:53:LEU:HA | 68:ss:60:ARG:NH2 | 2.35 | 0.42 |
| 72:sw:22:ILE:HG22 | 72:sw:89:LEU:HD23 | 2.02 | 0.42 |
| 74:sy:50:GLN:HE22 | 74:sy:78:LYS:NZ | 2.17 | 0.42 |
| 1:lA:721:A:C2 | 1:lA:722:A:C5 | 3.07 | 0.42 |
| 1:lA:721:A:O2' | 1:lA:722:A:OP1 | 2.37 | 0.42 |
| 1:lA:765:A:C2 | 36:lj:92:PRO:HG2 | 2.54 | 0.42 |
| 1:lA:915:U:O2' | 1:lA:916:U:H5' | 2.18 | 0.42 |
| 1:lA:1961:G:H2' | 1:lA:1962:U:O4' | 2.19 | 0.42 |
| 1:lA:2649:G:OP2 | 1:lA:2650:A:O2' | 2.30 | 0.42 |
| 1:lA:2872:A:N6 | 1:lA:2882:G:O6 | 2.52 | 0.42 |
| 1:lA:3129:U:C4 | 1:lA:3130:A:N7 | 2.88 | 0.42 |
| 5:lE:238:ARG:HE | 5:lE:238:ARG:HB2 | 1.67 | 0.42 |
| 36:lj:86:PHE:CE1 | 36:lj:90:LEU:HD11 | 2.54 | 0.42 |
| 52:sa:1693:A:HO2' | 52:sa:1694:A:P | 2.41 | 0.42 |
| 60:sj:42:ARG:HD3 | 60:sj:59:ARG:NH2 | 2.35 | 0.42 |
| 65:sp:81:LYS:HA | 65:sp:81:LYS:HD2 | 1.83 | 0.42 |
| 68:ss:28:LYS:HD2 | 68:ss:28:LYS:HA | 1.88 | 0.42 |
| 70:su:126:HIS:CE1 | 70:su:132:VAL:HG21 | 2.55 | 0.42 |
| 1:lA:285:A:H4' | 1:lA:286:U:H5' | 2.01 | 0.42 |
| 1:lA:871:A:H4' | 1:lA:872:A:OP1 | 2.19 | 0.42 |
| 1:lA:1239:U:H5'' | 29:lc:22:ILE:HD12 | 2.01 | 0.42 |
| 1:lA:1816:A:N3 | 1:lA:1892:C:O2' | 2.48 | 0.42 |
| 1:lA:1858:A:HO2' | 1:lA:1943:U:HO2' | 1.59 | 0.42 |
| 1:lA:1977:C:N4 | 1:lA:1978:U:O4 | 2.53 | 0.42 |
| 1:lA:2632:A:O2' | 28:lb:139:PHE:O | 2.35 | 0.42 |
| 3:lC:18:U:H3 | 3:lC:58:A:N6 | 2.11 | 0.42 |
| 3:lC:29:G:O6 | 3:lC:46:G:O6 | 2.38 | 0.42 |
| 3:lC:84:G:N2 | 3:lC:91:G:N2 | 2.68 | 0.42 |
| 6:lF:7:VAL:HG13 | 6:lF:149:GLN:NE2 | 2.33 | 0.42 |
| 8:lH:120:SER:C | 8:lH:121:LYS:HG2 | 2.44 | 0.42 |
| 13:lM:108:GLU:HG2 | 13:lM:122:ILE:HD12 | 2.02 | 0.42 |
| 23:lW:49:PHE:CE2 | 23:lW:96:TYR:HB2 | 2.54 | 0.42 |
| 37:lk:39:ARG:HD3 | 37:lk:39:ARG:HA | 1.75 | 0.42 |
| 42:lp:24:CYS:CA | 42:lp:40:CYS:HB3 | 2.50 | 0.42 |
| 47:sD:49:PRO:HG2 | 57:sg:146:LEU:HB3 | 2.00 | 0.42 |
| 52:sa:781:G:H5'' | 52:sa:782:A:O3' | 2.20 | 0.42 |
| 52:sa:959:A:H4' | 52:sa:1933:G:H21 | 1.84 | 0.42 |
| 52:sa:1378:U:H2' | 52:sa:1379:C:H6 | 1.84 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 54:sd:61:GLN:HG2 | 54:sd:101:TYR:HE2 | 1.85 | 0.42 |
| 70:su:85:LEU:HD13 | 70:su:97:SER:H | 1.84 | 0.42 |
| 1:lA:200:A:C4 | 1:lA:311:A:C6 | 3.08 | 0.42 |
| 1:lA:223:A:H2' | 1:lA:224:U:C6 | 2.54 | 0.42 |
| 1:lA:605:C:H2' | 1:lA:606:A:H8 | 1.84 | 0.42 |
| 1:lA:775:C:OP1 | 29:lc:21:ARG:HB3 | 2.20 | 0.42 |
| 1:lA:868:A:H2' | 1:lA:869:G:H8 | 1.85 | 0.42 |
| 1:lA:1231:A:H2' | 1:lA:1232:C:O4' | 2.19 | 0.42 |
| 1:lA:1259:G:O2' | 1:lA:2712:A:N3 | 2.44 | 0.42 |
| 1:lA:1904:U:O2' | 1:lA:1906:A:N7 | 2.46 | 0.42 |
| 1:lA:1920:A:H2' | 1:lA:1921:A:C8 | 2.55 | 0.42 |
| 42:lp:7:LEU:HD12 | 42:lp:7:LEU:HA | 1.82 | 0.42 |
| 52:sa:744:A:H2' | 52:sa:745:U:O4' | 2.19 | 0.42 |
| 52:sa:1335:U:H2' | 52:sa:1336:U:C6 | 2.55 | 0.42 |
| 59:si:118:LYS:HA | 59:si:118:LYS:HD2 | 1.69 | 0.42 |
| 60:sj:22:GLN:HG2 | 60:sj:23:LYS:O | 2.19 | 0.42 |
| 72:sw:55:ARG:HG2 | 72:sw:87:ARG:NH1 | 2.35 | 0.42 |
| 1:lA:235:U:C5 | 1:lA:252:G:N1 | 2.88 | 0.42 |
| 1:lA:338:A:H2' | 1:lA:339:U:C6 | 2.54 | 0.42 |
| 1:lA:570:A:C4 | 1:lA:571:A:C8 | 3.07 | 0.42 |
| 1:lA:761:A:H2' | 1:lA:762:A:C8 | 2.54 | 0.42 |
| 1:lA:1476:G:O2' | 1:lA:1477:A:N3 | 2.48 | 0.42 |
| 1:lA:1975:G:H8 | 1:lA:1975:G:OP2 | 2.03 | 0.42 |
| 1:lA:2083:A:H2' | 1:lA:2084:G:H8 | 1.85 | 0.42 |
| 1:lA:2205:C:H2' | 1:lA:2206:G:H8 | 1.85 | 0.42 |
| 1:lA:2360:C:N4 | 1:lA:2384:C:O4' | 2.53 | 0.42 |
| 1:lA:2734:C:O2' | 1:lA:2735:U:H5' | 2.20 | 0.42 |
| 1:lA:2867:A:H62 | 1:lA:2887:G:H1 | 1.67 | 0.42 |
| 1:lA:2867:A:H62 | 1:lA:2887:G:H22 | 1.67 | 0.42 |
| 1:lA:3095:G:H2' | 1:lA:3096:U:O4' | 2.20 | 0.42 |
| 3:lC:14:G:C2 | 3:lC:64:U:C4 | 3.08 | 0.42 |
| 5:lE:14:LEU:HD22 | 5:lE:264:TRP:CZ3 | 2.55 | 0.42 |
| 11:lK:60:GLU:HG3 | 11:lK:61:ASN:N | 2.34 | 0.42 |
| 13:lM:35:LYS:O | 13:lM:39:THR:HG23 | 2.20 | 0.42 |
| 28:lb:22:LYS:NZ | 28:lb:135:THR:O | 2.43 | 0.42 |
| 52:sa:291:A:H2' | 52:sa:292:A:H8 | 1.84 | 0.42 |
| 52:sa:327:U:OP1 | 60:sj:31:ARG:HD2 | 2.20 | 0.42 |
| 52:sa:415:A:O2' | 52:sa:416:A:H5' | 2.20 | 0.42 |
| 67:sr:46:TYR:HB3 | 67:sr:69:LEU:HD13 | 2.02 | 0.42 |
| 1:lA:625:A:C6 | 1:lA:663:G:C5 | 3.08 | 0.42 |
| 1:lA:812:A:C5 | 1:lA:839:G:N2 | 2.88 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:1A:948:C:O2' | 1:1A:949:G:N7 | 2.50 | 0.42 |
| 1:1A:2116:G:O2' | 21:1U:82:LYS:O | 2.36 | 0.42 |
| 1:1A:3469:A:H5'' | 1:1A:3470:G:C8 | 2.55 | 0.42 |
| 3:1C:107:A:O2' | 3:1C:108:U:H5' | 2.20 | 0.42 |
| 4:1D:180:LEU:HA | 4:1D:180:LEU:HD23 | 1.80 | 0.42 |
| 11:1K:37:ARG:HD3 | 11:1K:39:PHE:CE1 | 2.55 | 0.42 |
| 19:1S:27:LEU:HD12 | 19:1S:27:LEU:HA | 1.86 | 0.42 |
| 27:1a:150:ASN:O | 27:1a:153:ILE:HG12 | 2.20 | 0.42 |
| 27:1a:165:LYS:HE2 | 27:1a:165:LYS:HB2 | 1.75 | 0.42 |
| 52:sa:906:A:H2' | 52:sa:907:U:O4' | 2.20 | 0.42 |
| 52:sa:1093:U:H2' | 52:sa:1094:G:H8 | 1.85 | 0.42 |
| 52:sa:1141:U:H2' | 52:sa:1142:A:C8 | 2.51 | 0.42 |
| 55:se:116:VAL:HG22 | 55:se:144:PHE:HZ | 1.85 | 0.42 |
| 60:sj:38:LEU:HD11 | 60:sj:80:ASN:HA | 2.01 | 0.42 |
| 63:sm:92:TYR:HB2 | 63:sm:99:TYR:CE1 | 2.54 | 0.42 |
| 65:sp:30:ALA:HB2 | 65:sp:35:THR:HG23 | 2.01 | 0.42 |
| 66:sq:123:LYS:HB3 | 66:sq:123:LYS:HE2 | 1.78 | 0.42 |
| 1:1A:227:A:H2' | 1:1A:228:A:H8 | 1.85 | 0.42 |
| 1:1A:474:A:H2' | 1:1A:475:A:C8 | 2.54 | 0.42 |
| 1:1A:731:A:H2' | 1:1A:732:U:C6 | 2.55 | 0.42 |
| 1:1A:1479:A:H4' | 1:1A:1480:U:O5' | 2.18 | 0.42 |
| 1:1A:1922:U:H2' | 1:1A:1923:U:C6 | 2.55 | 0.42 |
| 1:1A:2062:A:O2' | 1:1A:3220:U:H5'' | 2.20 | 0.42 |
| 1:1A:2116:G:P | 1:1A:2133:G:H22 | 2.43 | 0.42 |
| 1:1A:2324:C:O2' | 1:1A:2348:G:H1' | 2.19 | 0.42 |
| 1:1A:2665:A:H2' | 1:1A:2666:G:H8 | 1.85 | 0.42 |
| 1:1A:3479:A:C6 | 1:1A:3480:A:C6 | 3.07 | 0.42 |
| 4:1D:247:ARG:HD2 | 4:1D:247:ARG:HA | 1.72 | 0.42 |
| 6:1F:386:LYS:H | 6:1F:386:LYS:HG3 | 1.66 | 0.42 |
| 6:1F:429:ILE:O | 20:1T:24:ARG:NH1 | 2.53 | 0.42 |
| 8:1H:62:GLY:O | 8:1H:117:VAL:N | 2.51 | 0.42 |
| 17:1Q:70:ASN:ND2 | 17:1Q:92:ILE:O | 2.29 | 0.42 |
| 24:1X:91:LYS:HD2 | 24:1X:91:LYS:HA | 1.84 | 0.42 |
| 45:sB:75:VAL:HB | 52:sa:1940:G:N1 | 2.35 | 0.42 |
| 52:sa:5:U:H2' | 52:sa:6:G:H8 | 1.85 | 0.42 |
| 52:sa:330:A:O2' | 63:sm:130:PRO:O | 2.36 | 0.42 |
| 52:sa:540:U:H2' | 52:sa:541:U:H6 | 1.83 | 0.42 |
| 52:sa:597:U:H2' | 52:sa:598:A:C8 | 2.55 | 0.42 |
| 52:sa:783:A:N1 | 52:sa:785:A:H5' | 2.34 | 0.42 |
| 52:sa:1185:A:H2' | 52:sa:1186:C:H6 | 1.85 | 0.42 |
| 52:sa:1905:U:C4 | 52:sa:1906:C:N4 | 2.88 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 54:sd:120:LYS:HG3 | 54:sd:189:VAL:HB | 2.01 | 0.42 |
| 55:se:193:ASN:OD1 | 55:se:194:LYS:N | 2.53 | 0.42 |
| 56:sf:79:ARG:HE | 56:sf:79:ARG:HB2 | 1.64 | 0.42 |
| 57:sg:78:MET:HE2 | 57:sg:171:LEU:HD21 | 2.02 | 0.42 |
| 1:lA:359:A:H2' | 1:lA:360:U:C6 | 2.55 | 0.41 |
| 1:lA:672:G:OP2 | 6:lF:403:ARG:NH1 | 2.49 | 0.41 |
| 1:lA:2752:C:H2' | 1:lA:2753:U:H6 | 1.84 | 0.41 |
| 1:lA:2846:U:H2' | 1:lA:2847:A:C8 | 2.54 | 0.41 |
| 1:lA:3303:A:H2' | 1:lA:3304:C:C6 | 2.56 | 0.41 |
| 5:lE:146:ARG:O | 5:lE:150:ILE:HG12 | 2.19 | 0.41 |
| 8:lH:45:LEU:HD11 | 8:lH:129:VAL:HG12 | 2.02 | 0.41 |
| 9:lI:30:LYS:HD3 | 9:lI:30:LYS:HA | 1.78 | 0.41 |
| 9:lI:76:ARG:HG2 | 9:lI:99:ILE:HA | 2.02 | 0.41 |
| 9:lI:105:ILE:HG21 | 9:lI:115:LEU:HD11 | 2.01 | 0.41 |
| 14:lN:49:ILE:HG12 | 14:lN:49:ILE:O | 2.19 | 0.41 |
| 15:lO:50:MET:O | 15:lO:54:GLN:HG2 | 2.20 | 0.41 |
| 15:lO:87:GLY:O | 15:lO:91:LYS:NZ | 2.52 | 0.41 |
| 17:lQ:62:TYR:CD1 | 17:lQ:134:LEU:HD12 | 2.55 | 0.41 |
| 24:lX:14:PHE:CE2 | 24:lX:91:LYS:HG2 | 2.54 | 0.41 |
| 28:lb:16:GLY:O | 34:lh:74:ARG:HG3 | 2.20 | 0.41 |
| 28:lb:93:ASN:H | 28:lb:120:GLN:HE22 | 1.66 | 0.41 |
| 52:sa:583:C:HO2' | 52:sa:584:C:H6 | 1.67 | 0.41 |
| 61:sk:43:GLN:H | 61:sk:43:GLN:HG3 | 1.67 | 0.41 |
| 65:sp:26:ALA:HB2 | 65:sp:88:LEU:HD13 | 2.01 | 0.41 |
| 1:lA:610:A:N6 | 1:lA:687:U:O4 | 2.53 | 0.41 |
| 1:lA:723:U:O2' | 1:lA:724:A:N3 | 2.45 | 0.41 |
| 1:lA:764:U:H2' | 1:lA:765:A:C8 | 2.52 | 0.41 |
| 1:lA:2040:A:O2' | 1:lA:2041:U:H5'' | 2.20 | 0.41 |
| 1:lA:2107:G:O6 | 5:lE:243:LYS:NZ | 2.33 | 0.41 |
| 1:lA:2341:C:N4 | 1:lA:2342:U:O4 | 2.53 | 0.41 |
| 1:lA:2747:G:H2' | 1:lA:2747:G:N3 | 2.35 | 0.41 |
| 4:lD:105:GLY:HA3 | 4:lD:160:SER:HB3 | 2.01 | 0.41 |
| 12:lL:179:GLU:OE2 | 12:lL:179:GLU:C | 2.63 | 0.41 |
| 23:lW:111:ASP:OD2 | 23:lW:113:LYS:HB2 | 2.20 | 0.41 |
| 40:ln:9:LYS:HA | 40:ln:9:LYS:HD3 | 1.82 | 0.41 |
| 41:lo:44:TRP:O | 41:lo:48:LYS:NZ | 2.35 | 0.41 |
| 47:sD:25:ARG:HG2 | 52:sa:1785:C:N3 | 2.35 | 0.41 |
| 52:sa:1170:U:H2' | 52:sa:1171:U:C6 | 2.55 | 0.41 |
| 52:sa:1937:A:O2' | 52:sa:1938:A:H5' | 2.20 | 0.41 |
| 54:sd:38:LEU:HD13 | 62:sl:64:TYR:CE2 | 2.55 | 0.41 |
| 57:sg:36:LYS:HA | 57:sg:36:LYS:HD2 | 1.80 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 63:sm:37:ILE:HD11 | 63:sm:65:ILE:HD12 | 2.01 | 0.41 |
| 68:ss:88:GLY:H | 68:ss:91:ALA:HB3 | 1.85 | 0.41 |
| 1:lA:606:A:H2' | 1:lA:607:U:C6 | 2.56 | 0.41 |
| 1:lA:780:A:OP2 | 33:lg:40:ASP:HB3 | 2.20 | 0.41 |
| 1:lA:1259:G:N3 | 1:lA:2712:A:H2 | 2.18 | 0.41 |
| 1:lA:2262:U:OP2 | 4:lD:200:ARG:HD2 | 2.20 | 0.41 |
| 1:lA:2304:A:H2' | 1:lA:2305:C:H6 | 1.85 | 0.41 |
| 1:lA:3008:U:O2' | 1:lA:3009:C:P | 2.79 | 0.41 |
| 1:lA:3008:U:HO2' | 1:lA:3009:C:P | 2.40 | 0.41 |
| 1:lA:3162:A:N1 | 1:lA:3295:A:C6 | 2.88 | 0.41 |
| 2:lB:42:A:OP2 | 2:lB:101:G:N1 | 2.38 | 0.41 |
| 3:lC:84:G:C2 | 3:lC:85:G:C8 | 3.08 | 0.41 |
| 6:lF:206:LYS:HD2 | 6:lF:206:LYS:HA | 1.83 | 0.41 |
| 15:lO:51:ARG:HA | 15:lO:51:ARG:HD2 | 1.70 | 0.41 |
| 19:lS:72:LYS:H | 19:lS:72:LYS:HG2 | 1.68 | 0.41 |
| 21:lU:77:GLY:O | 21:lU:81:ARG:NE | 2.54 | 0.41 |
| 27:la:81:ASP:OD2 | 27:la:82:LYS:N | 2.53 | 0.41 |
| 49:sI:5:C:H2' | 49:sI:6:C:C6 | 2.55 | 0.41 |
| 52:sa:607:G:OP1 | 52:sa:607:G:N2 | 2.45 | 0.41 |
| 62:sl:4:ALA:HB3 | 62:sl:7:ASP:HB2 | 2.01 | 0.41 |
| 67:sr:48:THR:OG1 | 67:sr:49:ASP:OD1 | 2.29 | 0.41 |
| 68:ss:54:ILE:HG23 | 68:ss:56:PRO:HD2 | 2.02 | 0.41 |
| 69:st:21:TYR:C | 69:st:21:TYR:CD1 | 2.99 | 0.41 |
| 1:lA:60:A:OP2 | 17:lQ:169:LYS:NZ | 2.49 | 0.41 |
| 1:lA:311:A:H5' | 37:lk:30:ARG:CB | 2.49 | 0.41 |
| 1:lA:918:G:O2' | 14:lN:16:TRP:NE1 | 2.52 | 0.41 |
| 1:lA:1203:U:H3' | 1:lA:1204:A:H2 | 1.85 | 0.41 |
| 1:lA:1641:A:H2' | 1:lA:1642:A:O4' | 2.20 | 0.41 |
| 1:lA:1920:A:H2' | 1:lA:1921:A:H8 | 1.86 | 0.41 |
| 1:lA:2866:C:H4' | 1:lA:2867:A:OP1 | 2.20 | 0.41 |
| 1:lA:3031:U:C2 | 1:lA:3054:A:N6 | 2.88 | 0.41 |
| 1:lA:3286:A:C2 | 1:lA:3287:U:H5' | 2.55 | 0.41 |
| 1:lA:3457:G:H1 | 1:lA:3495:U:H3 | 1.68 | 0.41 |
| 4:lD:108:PRO:O | 4:lD:111:THR:OG1 | 2.28 | 0.41 |
| 4:lD:186:TYR:HB2 | 4:lD:196:TRP:CZ3 | 2.56 | 0.41 |
| 5:lE:264:TRP:H | 5:lE:264:TRP:CD1 | 2.39 | 0.41 |
| 6:lF:6:PHE:HD2 | 6:lF:21:PRO:HB2 | 1.85 | 0.41 |
| 7:lG:237:ILE:HD13 | 7:lG:237:ILE:HA | 1.88 | 0.41 |
| 8:lH:97:ILE:HD13 | 8:lH:177:ILE:HG21 | 2.01 | 0.41 |
| 8:lH:146:VAL:HG13 | 8:lH:147:GLU:H | 1.85 | 0.41 |
| 21:lU:74:ARG:HD3 | 21:lU:74:ARG:HA | 1.84 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 28:lb:83:THR:HG22 | 34:lh:93:PHE:CE1 | 2.56 | 0.41 |
| 45:sB:85:ARG:H | 52:sa:1944:A:H61 | 1.68 | 0.41 |
| 52:sa:958:A:O2' | 52:sa:1934:C:O2 | 2.31 | 0.41 |
| 52:sa:972:A:H2' | 52:sa:973:A:H5' | 2.02 | 0.41 |
| 52:sa:1580:U:O2' | 57:sg:84:ASN:OD1 | 2.38 | 0.41 |
| 52:sa:1893:A:H4' | 52:sa:1893:A:OP1 | 2.20 | 0.41 |
| 53:sc:85:MET:HE3 | 53:sc:100:LEU:HB3 | 2.03 | 0.41 |
| 57:sg:31:SER:OG | 57:sg:32:ASP:N | 2.54 | 0.41 |
| 66:sq:55:LEU:HD23 | 66:sq:55:LEU:HA | 1.86 | 0.41 |
| 67:sr:69:LEU:HD12 | 67:sr:69:LEU:HA | 1.85 | 0.41 |
| 1:lA:790:A:H2' | 1:lA:791:A:C8 | 2.55 | 0.41 |
| 1:lA:1219:A:H4' | 1:lA:1220:A:OP2 | 2.20 | 0.41 |
| 1:lA:1533:U:HO2' | 1:lA:1534:U:P | 2.41 | 0.41 |
| 1:lA:2634:A:O2' | 10:IJ:30:VAL:HG13 | 2.20 | 0.41 |
| 2:lB:144:A:H2' | 2:lB:145:U:H6 | 1.84 | 0.41 |
| 6:lF:345:LYS:HD2 | 6:lF:345:LYS:HA | 1.81 | 0.41 |
| 7:lG:147:LEU:HD22 | 7:lG:164:LEU:HD12 | 2.03 | 0.41 |
| 8:lH:179:LYS:HE2 | 8:lH:179:LYS:HB2 | 1.86 | 0.41 |
| 9:II:33:ILE:O | 9:II:37:ILE:HG12 | 2.20 | 0.41 |
| 9:II:44:GLU:OE2 | 9:II:44:GLU:HA | 2.21 | 0.41 |
| 10:IJ:45:TYR:CE1 | 10:IJ:46:VAL:HG23 | 2.55 | 0.41 |
| 49:sI:52:G:H2' | 49:sI:53:G:C8 | 2.56 | 0.41 |
| 52:sa:247:G:P | 56:sf:132:GLY:H | 2.41 | 0.41 |
| 52:sa:878:A:C2 | 52:sa:895:A:C5 | 3.08 | 0.41 |
| 52:sa:1208:A:H61 | 66:sq:98:GLY:HA3 | 1.86 | 0.41 |
| 52:sa:1234:C:H42 | 52:sa:1562:G:N2 | 2.19 | 0.41 |
| 56:sf:39:CYS:HA | 56:sf:83:GLY:HA2 | 2.02 | 0.41 |
| 57:sg:27:ASP:OD1 | 57:sg:27:ASP:N | 2.50 | 0.41 |
| 57:sg:30:VAL:HB | 57:sg:36:LYS:HD3 | 2.02 | 0.41 |
| 58:sh:77:LEU:HD23 | 58:sh:77:LEU:HA | 1.82 | 0.41 |
| 60:sj:81:VAL:HA | 60:sj:102:ILE:HG13 | 2.01 | 0.41 |
| 72:sw:56:ILE:HB | 72:sw:86:LYS:HB3 | 2.02 | 0.41 |
| 1:lA:205:C:H3' | 1:lA:206:U:H5'' | 2.02 | 0.41 |
| 1:lA:229:A:H2' | 1:lA:230:U:C6 | 2.56 | 0.41 |
| 1:lA:693:A:N6 | 1:lA:694:A:H62 | 2.18 | 0.41 |
| 1:lA:790:A:H2' | 1:lA:791:A:H8 | 1.85 | 0.41 |
| 1:lA:2273:C:H4' | 1:lA:2274:A:O5' | 2.20 | 0.41 |
| 1:lA:3432:A:O2' | 1:lA:3433:U:P | 2.78 | 0.41 |
| 3:lC:27:G:C2 | 3:lC:51:G:C2 | 3.08 | 0.41 |
| 19:IS:86:TYR:OH | 29:lc:84:LYS:NZ | 2.34 | 0.41 |
| 31:le:11:GLN:HG3 | 31:le:75:SER:OG | 2.21 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|-------------------|--------------------------|-------------------|
| 48:sE:34:TYR:CZ | 72:sw:61:LEU:HG | 2.55 | 0.41 |
| 52:sa:42:A:N1 | 52:sa:373:A:C6 | 2.89 | 0.41 |
| 52:sa:42:A:C2 | 52:sa:373:A:C6 | 3.09 | 0.41 |
| 52:sa:62:G:O2' | 52:sa:165:U:OP1 | 2.39 | 0.41 |
| 52:sa:284:A:H2' | 52:sa:284:A:N3 | 2.35 | 0.41 |
| 52:sa:1394:A:C5 | 52:sa:1395:A:C8 | 3.09 | 0.41 |
| 52:sa:1903:A:H5'' | 52:sa:1904:G:OP2 | 2.21 | 0.41 |
| 55:se:26:PHE:HD2 | 65:sp:83:LEU:HD21 | 1.84 | 0.41 |
| 61:sk:33:GLN:HG2 | 61:sk:34:PHE:CD1 | 2.55 | 0.41 |
| 64:so:103:GLU:HB2 | 64:so:104:LYS:NZ | 2.35 | 0.41 |
| 65:sp:52:THR:O | 65:sp:55:MET:HB2 | 2.21 | 0.41 |
| 1:lA:22:A:C5 | 1:lA:376:A:C8 | 3.09 | 0.41 |
| 1:lA:515:G:C6 | 1:lA:558:G:C5 | 3.08 | 0.41 |
| 1:lA:896:A:O2' | 1:lA:2790:U:OP1 | 2.38 | 0.41 |
| 1:lA:985:C:N4 | 1:lA:1012:C:O2' | 2.53 | 0.41 |
| 1:lA:1265:G:C6 | 1:lA:1266:C:N4 | 2.88 | 0.41 |
| 1:lA:1764:A:H2' | 1:lA:1765:A:C8 | 2.56 | 0.41 |
| 1:lA:3371:U:H5'' | 16:lP:131:SER:HA | 2.02 | 0.41 |
| 2:lB:143:U:H2' | 2:lB:144:A:C8 | 2.55 | 0.41 |
| 3:lC:79:G:C2 | 3:lC:95:A:H2 | 2.33 | 0.41 |
| 13:lM:111:ASP:HB3 | 13:lM:112:LEU:H | 1.52 | 0.41 |
| 16:lP:84:ALA:O | 16:lP:88:ARG:HG3 | 2.20 | 0.41 |
| 23:lW:47:VAL:O | 23:lW:51:ARG:HG2 | 2.21 | 0.41 |
| 30:ld:54:LYS:HE2 | 30:ld:54:LYS:HB2 | 1.72 | 0.41 |
| 36:lj:18:ALA:HB3 | 36:lj:21:ASN:HB2 | 2.03 | 0.41 |
| 52:sa:204:A:H3' | 52:sa:205:A:H8 | 1.85 | 0.41 |
| 56:sf:121:LEU:HD12 | 56:sf:121:LEU:HA | 1.93 | 0.41 |
| 72:sw:54:VAL:HB | 72:sw:88:VAL:HB | 2.02 | 0.41 |
| 73:sx:61:ILE:HD13 | 73:sx:61:ILE:HA | 1.94 | 0.41 |
| 1:lA:249:U:N3 | 1:lA:250:G:N7 | 2.68 | 0.41 |
| 1:lA:408:A:HO2' | 1:lA:1047:U:HO2' | 1.68 | 0.41 |
| 1:lA:445:G:H5'' | 1:lA:448:C:O2' | 2.20 | 0.41 |
| 1:lA:1686:G:O6 | 17:lQ:35:ARG:NH2 | 2.50 | 0.41 |
| 1:lA:2025:C:H2' | 1:lA:2026:U:C6 | 2.55 | 0.41 |
| 1:lA:2349:G:O2' | 1:lA:2387:G:O6 | 2.34 | 0.41 |
| 2:lB:15:U:C2 | 2:lB:16:U:C5 | 3.09 | 0.41 |
| 6:lF:78:ARG:HA | 6:lF:89:GLN:O | 2.21 | 0.41 |
| 6:lF:212:TYR:HB2 | 6:lF:216:ASP:HB2 | 2.02 | 0.41 |
| 13:lM:155:THR:H | 13:lM:158:ASP:HB2 | 1.84 | 0.41 |
| 14:lN:194:LYS:HE2 | 14:lN:194:LYS:HB3 | 1.86 | 0.41 |
| 24:lX:113:LYS:HE2 | 24:lX:113:LYS:HB2 | 1.96 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 28:lb:115:LYS:HD3 | 28:lb:115:LYS:HA | 1.66 | 0.41 |
| 37:lk:84:ILE:HD13 | 37:lk:84:ILE:HA | 1.91 | 0.41 |
| 40:ln:57:LYS:C | 40:ln:58:LYS:HD3 | 2.45 | 0.41 |
| 45:sB:5:ARG:NH2 | 52:sa:1942:U:OP2 | 2.50 | 0.41 |
| 47:sD:16:THR:OG1 | 47:sD:32:ARG:O | 2.31 | 0.41 |
| 52:sa:102:A:H4' | 52:sa:103:U:O5' | 2.20 | 0.41 |
| 52:sa:1112:A:H2' | 52:sa:1113:A:C8 | 2.55 | 0.41 |
| 52:sa:1780:A:H3' | 52:sa:1780:A:N3 | 2.35 | 0.41 |
| 56:sf:141:ASP:OD1 | 56:sf:141:ASP:N | 2.53 | 0.41 |
| 66:sq:85:LEU:HD23 | 66:sq:85:LEU:HA | 1.82 | 0.41 |
| 72:sw:80:TYR:HD1 | 72:sw:80:TYR:HA | 1.77 | 0.41 |
| 1:lA:22:A:H2' | 1:lA:23:C:H6 | 1.85 | 0.41 |
| 1:lA:58:G:N2 | 17:lQ:186:ALA:HB1 | 2.35 | 0.41 |
| 1:lA:277:A:H2' | 1:lA:278:U:H6 | 1.86 | 0.41 |
| 1:lA:602:U:O3' | 6:lF:316:ARG:NH2 | 2.54 | 0.41 |
| 1:lA:739:A:H2' | 1:lA:740:A:C8 | 2.55 | 0.41 |
| 1:lA:810:G:N2 | 1:lA:906:G:C5 | 2.89 | 0.41 |
| 1:lA:827:A:N1 | 2:lB:30:U:O2' | 2.40 | 0.41 |
| 1:lA:1091:A:H4' | 1:lA:1494:G:H4' | 2.02 | 0.41 |
| 1:lA:1105:U:H2' | 1:lA:1106:U:C6 | 2.55 | 0.41 |
| 1:lA:1234:U:H2' | 1:lA:1235:U:C6 | 2.56 | 0.41 |
| 1:lA:1437:A:H2' | 1:lA:1438:U:C6 | 2.54 | 0.41 |
| 1:lA:1945:A:C6 | 1:lA:1956:G:C6 | 3.09 | 0.41 |
| 1:lA:1956:G:C2 | 1:lA:1957:A:C8 | 3.08 | 0.41 |
| 1:lA:2439:A:O2' | 1:lA:2440:G:H5' | 2.21 | 0.41 |
| 1:lA:2679:A:N3 | 17:lQ:87:GLN:NE2 | 2.58 | 0.41 |
| 1:lA:2710:A:H2' | 1:lA:2711:U:O4' | 2.20 | 0.41 |
| 1:lA:2864:A:O2' | 1:lA:2865:A:N3 | 2.41 | 0.41 |
| 1:lA:2887:G:H2' | 1:lA:2888:U:O4' | 2.20 | 0.41 |
| 1:lA:3368:G:H2' | 1:lA:3369:U:C6 | 2.56 | 0.41 |
| 2:lB:39:A:C6 | 2:lB:102:A:C5 | 3.08 | 0.41 |
| 3:lC:17:G:H1 | 3:lC:59:G:H22 | 1.69 | 0.41 |
| 3:lC:27:G:N2 | 3:lC:50:A:C2 | 2.89 | 0.41 |
| 6:lF:285:GLN:HG2 | 19:lS:25:MET:HG2 | 2.03 | 0.41 |
| 7:lG:106:ALA:O | 7:lG:110:LEU:HG | 2.21 | 0.41 |
| 8:lH:168:GLN:H | 8:lH:168:GLN:HG3 | 1.74 | 0.41 |
| 10:lJ:98:GLN:CD | 10:lJ:98:GLN:N | 2.79 | 0.41 |
| 13:lM:11:LYS:HA | 13:lM:11:LYS:HD3 | 1.91 | 0.41 |
| 14:lN:248:LYS:NZ | 29:lc:146:GLU:OE2 | 2.47 | 0.41 |
| 16:lP:12:VAL:HG22 | 16:lP:26:VAL:HG22 | 2.03 | 0.41 |
| 25:lY:43:LYS:HB2 | 25:lY:64:LEU:O | 2.21 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 25:IY:84:VAL:HG12 | 25:IY:109:VAL:HG21 | 2.03 | 0.41 |
| 30:ld:18:ARG:HA | 30:ld:18:ARG:HD2 | 1.89 | 0.41 |
| 35:li:70:LYS:HG3 | 35:li:73:ARG:HH21 | 1.85 | 0.41 |
| 36:lj:24:PRO:O | 36:lj:89:ASN:ND2 | 2.54 | 0.41 |
| 37:lk:80:LYS:HA | 37:lk:80:LYS:HD3 | 1.94 | 0.41 |
| 39:lm:47:VAL:HG13 | 39:lm:55:TRP:HB3 | 2.03 | 0.41 |
| 40:ln:11:ILE:O | 40:ln:15:LEU:HD12 | 2.20 | 0.41 |
| 52:sa:93:C:H2' | 52:sa:94:G:O4' | 2.21 | 0.41 |
| 52:sa:889:U:H2' | 52:sa:890:C:H6 | 1.86 | 0.41 |
| 52:sa:919:A:H2' | 52:sa:920:A:H8 | 1.85 | 0.41 |
| 52:sa:1200:U:H2' | 52:sa:1201:G:C8 | 2.56 | 0.41 |
| 52:sa:1266:A:H2' | 52:sa:1267:A:C4 | 2.56 | 0.41 |
| 52:sa:1576:A:H2' | 52:sa:1577:C:C6 | 2.56 | 0.41 |
| 52:sa:1756:U:O2 | 52:sa:1773:G:N2 | 2.54 | 0.41 |
| 52:sa:1906:C:H2' | 52:sa:1907:G:C8 | 2.56 | 0.41 |
| 63:sm:79:MET:HG2 | 63:sm:82:THR:HB | 2.03 | 0.41 |
| 65:sp:94:GLY:O | 65:sp:96:GLY:N | 2.54 | 0.41 |
| 68:ss:28:LYS:HG3 | 68:ss:29:LYS:H | 1.86 | 0.41 |
| 74:sy:7:ALA:O | 74:sy:11:MET:HG3 | 2.21 | 0.41 |
| 1:lA:366:U:H2' | 1:lA:367:C:H6 | 1.86 | 0.41 |
| 1:lA:494:G:OP2 | 1:lA:494:G:H8 | 2.04 | 0.41 |
| 1:lA:670:A:H2 | 6:lF:397:LEU:HD23 | 1.86 | 0.41 |
| 1:lA:1308:A:O2' | 1:lA:1309:A:H5' | 2.21 | 0.41 |
| 1:lA:1629:G:H1' | 34:lh:12:SER:HB3 | 2.01 | 0.41 |
| 1:lA:2468:C:HO2' | 1:lA:2469:G:P | 2.43 | 0.41 |
| 1:lA:2639:A:OP2 | 28:lb:56:LYS:NZ | 2.50 | 0.41 |
| 1:lA:2746:A:H5' | 1:lA:2747:G:C8 | 2.55 | 0.41 |
| 1:lA:3409:A:O2' | 1:lA:3410:A:OP2 | 2.32 | 0.41 |
| 2:lB:38:G:C5 | 35:li:78:ARG:HD3 | 2.56 | 0.41 |
| 10:lJ:91:GLU:OE2 | 10:lJ:99:ARG:NH1 | 2.47 | 0.41 |
| 12:lL:28:ASP:OD2 | 12:lL:28:ASP:C | 2.64 | 0.41 |
| 12:lL:38:ASN:OD1 | 12:lL:41:ALA:HB2 | 2.21 | 0.41 |
| 15:lO:117:LYS:HA | 15:lO:117:LYS:HD2 | 1.92 | 0.41 |
| 16:lP:7:VAL:HG22 | 16:lP:59:LEU:HD21 | 2.02 | 0.41 |
| 17:lQ:38:ARG:NH2 | 17:lQ:60:VAL:HG12 | 2.35 | 0.41 |
| 40:ln:23:VAL:HG22 | 40:ln:36:VAL:HG12 | 2.03 | 0.41 |
| 46:sC:5:GLU:HG2 | 73:sx:86:ILE:HD12 | 2.03 | 0.41 |
| 52:sa:29:G:C5 | 52:sa:591:G:N1 | 2.89 | 0.41 |
| 52:sa:311:C:C2 | 52:sa:350:G:C2 | 3.09 | 0.41 |
| 52:sa:905:G:N2 | 52:sa:968:A:H1' | 2.36 | 0.41 |
| 52:sa:934:A:H2' | 52:sa:935:C:C6 | 2.56 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 67:sr:79:PHE:CD2 | 67:sr:89:TRP:HH2 | 2.39 | 0.41 |
| 1:lA:58:G:H22 | 1:lA:75:U:H4' | 1.86 | 0.40 |
| 1:lA:661:G:H2' | 1:lA:662:U:H6 | 1.86 | 0.40 |
| 1:lA:811:A:OP2 | 19:lS:103:THR:HG21 | 2.20 | 0.40 |
| 1:lA:1529:A:N3 | 2:lB:21:A:O2' | 2.50 | 0.40 |
| 1:lA:1845:G:N2 | 1:lA:1924:U:O2' | 2.54 | 0.40 |
| 1:lA:1921:A:O2' | 34:lh:53:GLY:N | 2.48 | 0.40 |
| 1:lA:2598:A:H2' | 10:lJ:39:TYR:O | 2.21 | 0.40 |
| 2:lB:101:G:OP2 | 2:lB:103:A:O2' | 2.39 | 0.40 |
| 3:lC:84:G:N3 | 3:lC:84:G:H2' | 2.36 | 0.40 |
| 7:lG:200:VAL:HG12 | 7:lG:204:MET:HE2 | 2.03 | 0.40 |
| 21:lU:99:MET:HB3 | 21:lU:103:ARG:HE | 1.85 | 0.40 |
| 27:la:79:ASN:O | 27:la:80:ILE:HD13 | 2.21 | 0.40 |
| 40:ln:30:GLU:N | 40:ln:30:GLU:OE1 | 2.55 | 0.40 |
| 48:sE:21:CYS:HA | 48:sE:30:LEU:HD21 | 2.02 | 0.40 |
| 50:sJ:23:G:O2' | 50:sJ:24:C:P | 2.79 | 0.40 |
| 52:sa:387:G:OP1 | 52:sa:1876:U:O2' | 2.39 | 0.40 |
| 52:sa:1129:C:H2' | 52:sa:1130:U:C6 | 2.56 | 0.40 |
| 52:sa:1428:U:N3 | 68:ss:23:GLN:OE1 | 2.53 | 0.40 |
| 56:sf:124:ILE:HD11 | 56:sf:158:ILE:HG23 | 2.02 | 0.40 |
| 59:si:158:SER:OG | 59:si:194:LYS:NZ | 2.35 | 0.40 |
| 66:sq:32:PHE:HD1 | 66:sq:35:MET:HE2 | 1.85 | 0.40 |
| 67:sr:76:SER:O | 67:sr:77:PRO:C | 2.63 | 0.40 |
| 69:st:66:VAL:HG12 | 69:st:68:GLY:H | 1.86 | 0.40 |
| 70:su:25:ARG:H | 70:su:25:ARG:HG3 | 1.72 | 0.40 |
| 71:sv:58:ALA:HA | 71:sv:104:ILE:HG21 | 2.02 | 0.40 |
| 1:lA:51:G:O2' | 1:lA:52:A:H5' | 2.22 | 0.40 |
| 1:lA:229:A:H2' | 1:lA:230:U:H6 | 1.87 | 0.40 |
| 1:lA:237:C:H2' | 1:lA:238:A:C8 | 2.55 | 0.40 |
| 1:lA:338:A:H2' | 1:lA:339:U:H6 | 1.86 | 0.40 |
| 1:lA:572:U:H2' | 1:lA:573:A:H8 | 1.85 | 0.40 |
| 1:lA:586:A:N6 | 8:lH:43:THR:O | 2.55 | 0.40 |
| 1:lA:720:U:C2 | 1:lA:721:A:C8 | 3.09 | 0.40 |
| 1:lA:808:G:C6 | 1:lA:908:A:N1 | 2.89 | 0.40 |
| 1:lA:955:A:O2' | 39:lm:9:GLY:O | 2.27 | 0.40 |
| 1:lA:1305:A:C5 | 1:lA:1451:A:C6 | 3.09 | 0.40 |
| 1:lA:1689:A:H61 | 1:lA:2242:A:N6 | 2.18 | 0.40 |
| 1:lA:2224:U:H2' | 1:lA:2225:A:N7 | 2.36 | 0.40 |
| 1:lA:2865:A:H4' | 37:lk:21:ALA:HA | 2.02 | 0.40 |
| 1:lA:3130:A:H2' | 1:lA:3131:C:C6 | 2.56 | 0.40 |
| 1:lA:3483:A:H2' | 1:lA:3484:A:H2 | 1.85 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 11:IK:41:HIS:CE1 | 11:IK:77:VAL:HG11 | 2.56 | 0.40 |
| 18:IR:41:VAL:HG23 | 18:IR:95:LEU:HD22 | 2.02 | 0.40 |
| 20:IT:10:CYS:HG | 20:IT:15:HIS:HD1 | 1.60 | 0.40 |
| 20:IT:148:VAL:HG23 | 20:IT:172:LEU:HD23 | 2.03 | 0.40 |
| 20:IT:148:VAL:HG22 | 20:IT:149:HIS:H | 1.87 | 0.40 |
| 33:lg:14:TYR:OH | 33:lg:52:LEU:HD21 | 2.22 | 0.40 |
| 37:lk:55:LEU:HD23 | 37:lk:55:LEU:HA | 1.89 | 0.40 |
| 39:lm:16:THR:O | 39:lm:16:THR:OG1 | 2.29 | 0.40 |
| 43:lq:57:LYS:HE2 | 43:lq:57:LYS:HB2 | 1.87 | 0.40 |
| 47:sD:25:ARG:HG2 | 52:sa:1785:C:C4 | 2.55 | 0.40 |
| 52:sa:1710:U:OP1 | 70:su:135:GLN:NE2 | 2.51 | 0.40 |
| 54:sd:156:LEU:HD12 | 54:sd:157:ARG:H | 1.87 | 0.40 |
| 56:sf:174:ASP:H | 56:sf:177:ASN:ND2 | 2.19 | 0.40 |
| 59:si:125:LYS:HE2 | 59:si:125:LYS:HB3 | 1.92 | 0.40 |
| 60:sj:57:ALA:HB1 | 60:sj:60:LEU:HD11 | 2.02 | 0.40 |
| 66:sq:87:ASN:OD1 | 66:sq:88:MET:HE2 | 2.21 | 0.40 |
| 69:st:74:GLN:O | 69:st:78:ARG:N | 2.49 | 0.40 |
| 70:su:107:ARG:NE | 70:su:110:GLU:OE2 | 2.54 | 0.40 |
| 1:lA:336:A:H2 | 17:lQ:93:LYS:HD3 | 1.86 | 0.40 |
| 1:lA:732:U:O2' | 1:lA:733:C:H6 | 2.04 | 0.40 |
| 1:lA:798:U:H2' | 1:lA:799:A:C8 | 2.56 | 0.40 |
| 1:lA:1047:U:H2' | 1:lA:1048:A:C8 | 2.56 | 0.40 |
| 1:lA:1542:U:H2' | 1:lA:1543:U:H6 | 1.85 | 0.40 |
| 1:lA:1558:C:H2' | 1:lA:1559:C:H6 | 1.86 | 0.40 |
| 1:lA:1684:A:C6 | 1:lA:1685:U:C4 | 3.10 | 0.40 |
| 6:lF:98:LYS:HE2 | 6:lF:98:LYS:HB2 | 1.85 | 0.40 |
| 9:II:86:LYS:HE3 | 9:II:117:LEU:HD13 | 2.02 | 0.40 |
| 12:IL:150:LYS:HE3 | 12:IL:150:LYS:HB3 | 1.82 | 0.40 |
| 14:IN:49:ILE:O | 14:IN:232:ASP:HA | 2.22 | 0.40 |
| 17:lQ:14:LYS:HA | 17:lQ:19:LEU:HD12 | 2.03 | 0.40 |
| 18:IR:57:ILE:HG12 | 18:IR:73:CYS:SG | 2.61 | 0.40 |
| 23:IW:47:VAL:HA | 23:IW:68:VAL:HG11 | 2.02 | 0.40 |
| 24:IX:25:THR:OG1 | 24:IX:38:THR:HG22 | 2.20 | 0.40 |
| 45:sB:45:VAL:HG21 | 45:sB:64:LEU:HG | 2.03 | 0.40 |
| 45:sB:97:PRO:HA | 45:sB:98:PRO:HD3 | 2.01 | 0.40 |
| 52:sa:254:C:H2' | 52:sa:255:G:O4' | 2.21 | 0.40 |
| 52:sa:594:U:H2' | 52:sa:595:A:H8 | 1.86 | 0.40 |
| 52:sa:853:U:H2' | 52:sa:854:U:C6 | 2.57 | 0.40 |
| 52:sa:1259:A:O2' | 52:sa:1260:C:H5' | 2.21 | 0.40 |
| 52:sa:1758:G:H2' | 52:sa:1759:A:C8 | 2.56 | 0.40 |
| 53:sc:55:ILE:HG21 | 53:sc:75:LEU:HD11 | 2.03 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 54:sd:121:TYR:O | 54:sd:125:GLU:HG3 | 2.21 | 0.40 |
| 55:se:193:ASN:O | 55:se:196:PRO:HD2 | 2.22 | 0.40 |
| 56:sf:189:ARG:HG2 | 56:sf:243:LYS:HB2 | 2.03 | 0.40 |
| 57:sg:152:ALA:O | 57:sg:156:MET:HG2 | 2.21 | 0.40 |
| 61:sk:25:ASP:OD1 | 61:sk:25:ASP:N | 2.53 | 0.40 |
| 62:sl:45:MET:HA | 62:sl:48:LEU:HD13 | 2.04 | 0.40 |
| 1:lA:549:U:HO2' | 1:lA:550:A:H8 | 1.67 | 0.40 |
| 1:lA:835:A:H2' | 1:lA:836:C:O4' | 2.20 | 0.40 |
| 1:lA:1242:G:O6 | 1:lA:1267:G:N2 | 2.55 | 0.40 |
| 1:lA:2188:U:H3 | 1:lA:3216:A:H4' | 1.86 | 0.40 |
| 2:lB:73:A:H4' | 2:lB:74:A:H5' | 2.03 | 0.40 |
| 4:lD:89:TYR:N | 4:lD:100:ASN:OD1 | 2.40 | 0.40 |
| 7:lG:5:LYS:HA | 7:lG:5:LYS:HD3 | 1.65 | 0.40 |
| 9:II:77:ILE:HG12 | 9:II:123:THR:HG22 | 2.03 | 0.40 |
| 14:lN:53:LYS:HB2 | 14:lN:69:LEU:HB3 | 2.04 | 0.40 |
| 20:IT:10:CYS:HG | 20:IT:15:HIS:CG | 2.38 | 0.40 |
| 30:ld:36:ASN:HB3 | 30:ld:39:VAL:HB | 2.03 | 0.40 |
| 34:lh:93:PHE:O | 34:lh:97:GLU:HG2 | 2.22 | 0.40 |
| 37:lk:48:GLU:OE2 | 37:lk:80:LYS:NZ | 2.27 | 0.40 |
| 52:sa:291:A:H2' | 52:sa:292:A:C8 | 2.57 | 0.40 |
| 52:sa:426:C:H2' | 52:sa:427:G:C8 | 2.56 | 0.40 |
| 52:sa:798:A:C8 | 67:sr:107:SER:HA | 2.56 | 0.40 |
| 70:su:85:LEU:HD13 | 70:su:97:SER:HB3 | 2.02 | 0.40 |
| 1:lA:247:A:C5 | 1:lA:248:U:C5 | 3.09 | 0.40 |
| 1:lA:789:C:OP1 | 33:lg:28:ARG:N | 2.54 | 0.40 |
| 1:lA:1070:A:OP2 | 1:lA:1489:G:N2 | 2.37 | 0.40 |
| 1:lA:2756:U:OP1 | 13:lM:147:LYS:HE3 | 2.22 | 0.40 |
| 1:lA:3132:C:H2' | 1:lA:3133:G:O4' | 2.21 | 0.40 |
| 3:lC:28:U:H4' | 13:lM:9:MET:HE1 | 2.02 | 0.40 |
| 5:lE:37:ALA:HA | 5:lE:187:GLY:O | 2.22 | 0.40 |
| 6:lF:395:ASN:N | 6:lF:395:ASN:OD1 | 2.54 | 0.40 |
| 7:lG:207:LEU:HD12 | 7:lG:207:LEU:HA | 1.85 | 0.40 |
| 13:lM:137:ARG:O | 13:lM:137:ARG:HG2 | 2.22 | 0.40 |
| 14:lN:163:ILE:O | 14:lN:167:LYS:HB2 | 2.21 | 0.40 |
| 24:IX:100:ASP:OD1 | 26:lZ:26:LEU:HD11 | 2.21 | 0.40 |
| 27:la:104:ILE:HG21 | 27:la:107:LEU:HD23 | 2.04 | 0.40 |
| 27:la:164:LYS:HD2 | 27:la:164:LYS:HA | 1.80 | 0.40 |
| 30:ld:7:ARG:HG2 | 30:ld:8:SER:N | 2.35 | 0.40 |
| 52:sa:427:G:H2' | 52:sa:428:C:C6 | 2.57 | 0.40 |
| 52:sa:1923:A:C6 | 52:sa:1933:G:O6 | 2.73 | 0.40 |
| 55:se:162:GLN:O | 55:se:166:ILE:HG12 | 2.22 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 57:sg:193:LYS:HE2 | 57:sg:193:LYS:HB3 | 1.96 | 0.40 |
| 68:ss:43:MET:HE1 | 71:sv:2:HIS:HA | 2.02 | 0.40 |

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 PDB entries followed by that with respect to all EM entries.

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 | |
|-----|-------|---------------|-----------|---------|----------|-------------|-----|
| 4 | ID | 244/257 (95%) | 226 (93%) | 18 (7%) | 0 | 100 | 100 |
| 5 | IE | 385/402 (96%) | 362 (94%) | 23 (6%) | 0 | 100 | 100 |
| 6 | IF | 421/431 (98%) | 392 (93%) | 28 (7%) | 1 (0%) | 44 | 74 |
| 7 | IG | 274/286 (96%) | 255 (93%) | 19 (7%) | 0 | 100 | 100 |
| 8 | IH | 201/204 (98%) | 185 (92%) | 14 (7%) | 2 (1%) | 13 | 42 |
| 9 | II | 208/230 (90%) | 199 (96%) | 8 (4%) | 1 (0%) | 25 | 58 |
| 10 | IJ | 197/286 (69%) | 188 (95%) | 9 (5%) | 0 | 100 | 100 |
| 11 | IK | 191/197 (97%) | 184 (96%) | 6 (3%) | 1 (0%) | 25 | 58 |
| 12 | IL | 196/210 (93%) | 184 (94%) | 12 (6%) | 0 | 100 | 100 |
| 13 | IM | 166/174 (95%) | 162 (98%) | 3 (2%) | 1 (1%) | 22 | 53 |
| 14 | IN | 261/291 (90%) | 247 (95%) | 14 (5%) | 0 | 100 | 100 |
| 15 | IO | 202/205 (98%) | 192 (95%) | 10 (5%) | 0 | 100 | 100 |
| 16 | IP | 128/135 (95%) | 124 (97%) | 4 (3%) | 0 | 100 | 100 |
| 17 | IQ | 202/205 (98%) | 194 (96%) | 8 (4%) | 0 | 100 | 100 |
| 18 | IR | 156/179 (87%) | 148 (95%) | 8 (5%) | 0 | 100 | 100 |
| 19 | IS | 165/168 (98%) | 151 (92%) | 14 (8%) | 0 | 100 | 100 |
| 20 | IT | 171/173 (99%) | 162 (95%) | 9 (5%) | 0 | 100 | 100 |
| 21 | IU | 148/198 (75%) | 145 (98%) | 3 (2%) | 0 | 100 | 100 |

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| Mol | Chain | Analysed | Favoured | Allowed | Outliers | Percentiles | |
|-----|-------|---------------|-----------|----------|----------|-------------|-----|
| 22 | IV | 163/166 (98%) | 157 (96%) | 6 (4%) | 0 | 100 | 100 |
| 23 | IW | 91/137 (66%) | 83 (91%) | 8 (9%) | 0 | 100 | 100 |
| 24 | IX | 131/140 (94%) | 125 (95%) | 6 (5%) | 0 | 100 | 100 |
| 25 | IY | 114/121 (94%) | 112 (98%) | 2 (2%) | 0 | 100 | 100 |
| 26 | IZ | 55/163 (34%) | 54 (98%) | 1 (2%) | 0 | 100 | 100 |
| 27 | la | 208/213 (98%) | 197 (95%) | 11 (5%) | 0 | 100 | 100 |
| 28 | lb | 135/139 (97%) | 132 (98%) | 3 (2%) | 0 | 100 | 100 |
| 29 | lc | 146/149 (98%) | 134 (92%) | 12 (8%) | 0 | 100 | 100 |
| 30 | ld | 58/64 (91%) | 54 (93%) | 4 (7%) | 0 | 100 | 100 |
| 31 | le | 91/109 (84%) | 81 (89%) | 10 (11%) | 0 | 100 | 100 |
| 32 | lf | 122/150 (81%) | 115 (94%) | 7 (6%) | 0 | 100 | 100 |
| 33 | lg | 121/134 (90%) | 118 (98%) | 3 (2%) | 0 | 100 | 100 |
| 34 | lh | 100/137 (73%) | 98 (98%) | 2 (2%) | 0 | 100 | 100 |
| 35 | li | 120/122 (98%) | 118 (98%) | 1 (1%) | 1 (1%) | 16 | 48 |
| 36 | lj | 104/108 (96%) | 98 (94%) | 6 (6%) | 0 | 100 | 100 |
| 37 | lk | 83/104 (80%) | 81 (98%) | 2 (2%) | 0 | 100 | 100 |
| 38 | ll | 70/77 (91%) | 63 (90%) | 5 (7%) | 2 (3%) | 3 | 20 |
| 39 | lm | 88/93 (95%) | 82 (93%) | 6 (7%) | 0 | 100 | 100 |
| 40 | ln | 71/77 (92%) | 68 (96%) | 3 (4%) | 0 | 100 | 100 |
| 41 | lo | 48/51 (94%) | 45 (94%) | 3 (6%) | 0 | 100 | 100 |
| 42 | lp | 51/56 (91%) | 48 (94%) | 3 (6%) | 0 | 100 | 100 |
| 43 | lq | 90/98 (92%) | 89 (99%) | 1 (1%) | 0 | 100 | 100 |
| 44 | sA | 8/137 (6%) | 6 (75%) | 2 (25%) | 0 | 100 | 100 |
| 45 | sB | 96/144 (67%) | 93 (97%) | 3 (3%) | 0 | 100 | 100 |
| 46 | sC | 30/84 (36%) | 30 (100%) | 0 | 0 | 100 | 100 |
| 47 | sD | 58/69 (84%) | 52 (90%) | 6 (10%) | 0 | 100 | 100 |
| 48 | sE | 53/56 (95%) | 50 (94%) | 3 (6%) | 0 | 100 | 100 |
| 53 | sc | 194/255 (76%) | 184 (95%) | 10 (5%) | 0 | 100 | 100 |
| 54 | sd | 184/244 (75%) | 174 (95%) | 10 (5%) | 0 | 100 | 100 |
| 55 | se | 126/256 (49%) | 121 (96%) | 4 (3%) | 1 (1%) | 16 | 48 |
| 56 | sf | 254/326 (78%) | 231 (91%) | 23 (9%) | 0 | 100 | 100 |

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| Mol | Chain | Analysed | Favoured | Allowed | Outliers | Percentiles | |
|-----|-------|------------------|------------|----------|----------|-------------|-----|
| 57 | sg | 175/206 (85%) | 166 (95%) | 9 (5%) | 0 | 100 | 100 |
| 58 | sh | 49/266 (18%) | 48 (98%) | 1 (2%) | 0 | 100 | 100 |
| 59 | si | 53/201 (26%) | 53 (100%) | 0 | 0 | 100 | 100 |
| 60 | sj | 125/237 (53%) | 120 (96%) | 5 (4%) | 0 | 100 | 100 |
| 61 | sk | 80/185 (43%) | 79 (99%) | 1 (1%) | 0 | 100 | 100 |
| 62 | sl | 64/127 (50%) | 63 (98%) | 1 (2%) | 0 | 100 | 100 |
| 63 | sm | 137/156 (88%) | 127 (93%) | 10 (7%) | 0 | 100 | 100 |
| 64 | so | 68/151 (45%) | 66 (97%) | 2 (3%) | 0 | 100 | 100 |
| 65 | sp | 126/146 (86%) | 119 (94%) | 7 (6%) | 0 | 100 | 100 |
| 66 | sq | 101/144 (70%) | 92 (91%) | 8 (8%) | 1 (1%) | 13 | 42 |
| 67 | sr | 127/130 (98%) | 117 (92%) | 10 (8%) | 0 | 100 | 100 |
| 68 | ss | 135/158 (85%) | 129 (96%) | 6 (4%) | 0 | 100 | 100 |
| 69 | st | 79/117 (68%) | 79 (100%) | 0 | 0 | 100 | 100 |
| 70 | su | 120/155 (77%) | 108 (90%) | 12 (10%) | 0 | 100 | 100 |
| 71 | sv | 123/155 (79%) | 116 (94%) | 7 (6%) | 0 | 100 | 100 |
| 72 | sw | 42/118 (36%) | 42 (100%) | 0 | 0 | 100 | 100 |
| 73 | sx | 59/86 (69%) | 57 (97%) | 2 (3%) | 0 | 100 | 100 |
| 74 | sy | 104/141 (74%) | 97 (93%) | 7 (7%) | 0 | 100 | 100 |
| All | All | 8946/11289 (79%) | 8471 (95%) | 464 (5%) | 11 (0%) | 50 | 79 |

All (11) Ramachandran outliers are listed below:

| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 38 | ll | 40 | PRO |
| 55 | se | 193 | ASN |
| 6 | lF | 390 | VAL |
| 8 | lH | 119 | VAL |
| 13 | lM | 28 | ASP |
| 66 | sq | 112 | GLY |
| 35 | li | -6 | LYS |
| 38 | ll | 39 | TYR |
| 11 | lK | 134 | ILE |
| 8 | lH | 118 | ASP |
| 9 | lI | 179 | VAL |

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 PDB entries followed by that with respect to all EM entries.

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 | |
|-----|-------|----------------|------------|----------|-------------|-----|
| 4 | ID | 195/201 (97%) | 191 (98%) | 4 (2%) | 48 | 72 |
| 5 | IE | 330/343 (96%) | 325 (98%) | 5 (2%) | 60 | 80 |
| 6 | IF | 339/345 (98%) | 331 (98%) | 8 (2%) | 44 | 70 |
| 7 | IG | 223/231 (96%) | 217 (97%) | 6 (3%) | 40 | 67 |
| 8 | IH | 172/173 (99%) | 168 (98%) | 4 (2%) | 45 | 70 |
| 9 | II | 178/195 (91%) | 176 (99%) | 2 (1%) | 70 | 84 |
| 10 | IJ | 177/242 (73%) | 175 (99%) | 2 (1%) | 70 | 84 |
| 11 | IK | 171/174 (98%) | 170 (99%) | 1 (1%) | 84 | 91 |
| 12 | IL | 169/176 (96%) | 166 (98%) | 3 (2%) | 54 | 76 |
| 13 | IM | 144/147 (98%) | 135 (94%) | 9 (6%) | 15 | 42 |
| 14 | IN | 224/243 (92%) | 221 (99%) | 3 (1%) | 65 | 82 |
| 15 | IO | 167/168 (99%) | 166 (99%) | 1 (1%) | 84 | 91 |
| 16 | IP | 114/118 (97%) | 113 (99%) | 1 (1%) | 75 | 88 |
| 17 | IQ | 171/172 (99%) | 171 (100%) | 0 | 100 | 100 |
| 18 | IR | 129/147 (88%) | 126 (98%) | 3 (2%) | 45 | 70 |
| 19 | IS | 142/143 (99%) | 138 (97%) | 4 (3%) | 38 | 66 |
| 20 | IT | 156/156 (100%) | 156 (100%) | 0 | 100 | 100 |
| 21 | IU | 132/174 (76%) | 130 (98%) | 2 (2%) | 60 | 80 |
| 22 | IV | 144/145 (99%) | 141 (98%) | 3 (2%) | 48 | 72 |
| 23 | IW | 86/125 (69%) | 83 (96%) | 3 (4%) | 31 | 61 |
| 24 | IX | 109/113 (96%) | 105 (96%) | 4 (4%) | 29 | 59 |
| 25 | IY | 99/102 (97%) | 98 (99%) | 1 (1%) | 73 | 86 |
| 26 | IZ | 52/137 (38%) | 51 (98%) | 1 (2%) | 52 | 75 |
| 27 | la | 177/179 (99%) | 173 (98%) | 4 (2%) | 45 | 70 |
| 28 | lb | 121/123 (98%) | 119 (98%) | 2 (2%) | 56 | 78 |
| 29 | lc | 120/121 (99%) | 116 (97%) | 4 (3%) | 33 | 62 |

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| Mol | Chain | Analysed | Rotameric | Outliers | Percentiles | |
|-----|-------|----------------|------------|----------|-------------|-----|
| 30 | ld | 50/54 (93%) | 50 (100%) | 0 | 100 | 100 |
| 31 | le | 78/92 (85%) | 75 (96%) | 3 (4%) | 28 | 59 |
| 32 | lf | 110/128 (86%) | 107 (97%) | 3 (3%) | 40 | 67 |
| 33 | lg | 107/116 (92%) | 106 (99%) | 1 (1%) | 75 | 88 |
| 34 | lh | 86/116 (74%) | 85 (99%) | 1 (1%) | 67 | 83 |
| 35 | li | 103/103 (100%) | 100 (97%) | 3 (3%) | 37 | 65 |
| 36 | lj | 89/91 (98%) | 88 (99%) | 1 (1%) | 70 | 84 |
| 37 | lk | 71/82 (87%) | 70 (99%) | 1 (1%) | 62 | 81 |
| 38 | ll | 60/64 (94%) | 60 (100%) | 0 | 100 | 100 |
| 39 | lm | 73/75 (97%) | 70 (96%) | 3 (4%) | 26 | 57 |
| 40 | ln | 63/66 (96%) | 60 (95%) | 3 (5%) | 21 | 51 |
| 41 | lo | 44/45 (98%) | 42 (96%) | 2 (4%) | 23 | 53 |
| 42 | lp | 45/48 (94%) | 45 (100%) | 0 | 100 | 100 |
| 43 | lq | 85/91 (93%) | 84 (99%) | 1 (1%) | 67 | 83 |
| 44 | sA | 10/112 (9%) | 9 (90%) | 1 (10%) | 6 | 24 |
| 45 | sB | 87/127 (68%) | 83 (95%) | 4 (5%) | 23 | 52 |
| 46 | sC | 28/73 (38%) | 25 (89%) | 3 (11%) | 5 | 21 |
| 47 | sD | 50/59 (85%) | 46 (92%) | 4 (8%) | 10 | 34 |
| 48 | sE | 45/46 (98%) | 43 (96%) | 2 (4%) | 24 | 54 |
| 53 | sc | 157/199 (79%) | 153 (98%) | 4 (2%) | 42 | 69 |
| 54 | sd | 156/206 (76%) | 151 (97%) | 5 (3%) | 34 | 63 |
| 55 | se | 130/225 (58%) | 127 (98%) | 3 (2%) | 45 | 70 |
| 56 | sf | 223/283 (79%) | 220 (99%) | 3 (1%) | 65 | 82 |
| 57 | sg | 160/178 (90%) | 160 (100%) | 0 | 100 | 100 |
| 58 | sh | 42/220 (19%) | 39 (93%) | 3 (7%) | 12 | 39 |
| 59 | si | 48/167 (29%) | 46 (96%) | 2 (4%) | 25 | 56 |
| 60 | sj | 108/205 (53%) | 106 (98%) | 2 (2%) | 52 | 75 |
| 61 | sk | 72/164 (44%) | 72 (100%) | 0 | 100 | 100 |
| 62 | sl | 60/111 (54%) | 60 (100%) | 0 | 100 | 100 |
| 63 | sm | 126/138 (91%) | 124 (98%) | 2 (2%) | 58 | 79 |
| 64 | so | 68/129 (53%) | 67 (98%) | 1 (2%) | 60 | 80 |

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| Mol | Chain | Analysed | Rotameric | Outliers | Percentiles | |
|-----|-------|-----------------|------------|----------|-------------|-----|
| 65 | sp | 99/114 (87%) | 96 (97%) | 3 (3%) | 36 | 64 |
| 66 | sq | 94/127 (74%) | 94 (100%) | 0 | 100 | 100 |
| 67 | sr | 112/113 (99%) | 109 (97%) | 3 (3%) | 40 | 67 |
| 68 | ss | 111/128 (87%) | 108 (97%) | 3 (3%) | 40 | 67 |
| 69 | st | 73/106 (69%) | 72 (99%) | 1 (1%) | 62 | 81 |
| 70 | su | 104/130 (80%) | 101 (97%) | 3 (3%) | 37 | 65 |
| 71 | sv | 106/132 (80%) | 104 (98%) | 2 (2%) | 52 | 75 |
| 72 | sw | 44/107 (41%) | 44 (100%) | 0 | 100 | 100 |
| 73 | sx | 56/77 (73%) | 53 (95%) | 3 (5%) | 18 | 47 |
| 74 | sy | 86/114 (75%) | 85 (99%) | 1 (1%) | 67 | 83 |
| All | All | 7760/9554 (81%) | 7600 (98%) | 160 (2%) | 49 | 72 |

All (160) residues with a non-rotameric sidechain are listed below:

| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 4 | ID | 5 | THR |
| 4 | ID | 33 | ASP |
| 4 | ID | 40 | THR |
| 4 | ID | 193 | ARG |
| 5 | IE | 166 | THR |
| 5 | IE | 213 | THR |
| 5 | IE | 231 | VAL |
| 5 | IE | 232 | ILE |
| 5 | IE | 374 | THR |
| 6 | IF | 79 | ILE |
| 6 | IF | 94 | ASN |
| 6 | IF | 115 | THR |
| 6 | IF | 153 | VAL |
| 6 | IF | 190 | VAL |
| 6 | IF | 215 | ASP |
| 6 | IF | 342 | GLN |
| 6 | IF | 386 | LYS |
| 7 | IG | 9 | ASN |
| 7 | IG | 50 | ARG |
| 7 | IG | 94 | ASN |
| 7 | IG | 101 | THR |
| 7 | IG | 167 | MET |
| 7 | IG | 204 | MET |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 8 | 1H | 8 | VAL |
| 8 | 1H | 146 | VAL |
| 8 | 1H | 155 | TYR |
| 8 | 1H | 161 | LEU |
| 9 | 1I | 52 | THR |
| 9 | 1I | 77 | ILE |
| 10 | 1J | 25 | PHE |
| 10 | 1J | 36 | LEU |
| 11 | 1K | 71 | ASN |
| 12 | 1L | 50 | VAL |
| 12 | 1L | 119 | TRP |
| 12 | 1L | 201 | THR |
| 13 | 1M | 14 | ILE |
| 13 | 1M | 19 | LEU |
| 13 | 1M | 31 | THR |
| 13 | 1M | 36 | VAL |
| 13 | 1M | 48 | SER |
| 13 | 1M | 110 | ILE |
| 13 | 1M | 111 | ASP |
| 13 | 1M | 148 | ILE |
| 13 | 1M | 155 | THR |
| 14 | 1N | 61 | GLN |
| 14 | 1N | 155 | PHE |
| 14 | 1N | 177 | TYR |
| 15 | 1O | 23 | VAL |
| 16 | 1P | 68 | THR |
| 18 | 1R | 43 | ARG |
| 18 | 1R | 120 | GLN |
| 18 | 1R | 139 | PHE |
| 19 | 1S | 10 | VAL |
| 19 | 1S | 60 | ILE |
| 19 | 1S | 80 | CYS |
| 19 | 1S | 152 | VAL |
| 21 | 1U | 89 | THR |
| 21 | 1U | 149 | LYS |
| 22 | 1V | 89 | LEU |
| 22 | 1V | 91 | ILE |
| 22 | 1V | 129 | LEU |
| 23 | 1W | 39 | ASP |
| 23 | 1W | 54 | ILE |
| 23 | 1W | 100 | ASN |
| 24 | 1X | 17 | THR |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 24 | lX | 59 | ASP |
| 24 | lX | 123 | LYS |
| 24 | lX | 128 | LEU |
| 25 | lY | 32 | ASN |
| 26 | lZ | 59 | GLN |
| 27 | la | 22 | ASN |
| 27 | la | 69 | THR |
| 27 | la | 109 | LEU |
| 27 | la | 158 | VAL |
| 28 | lb | 63 | GLU |
| 28 | lb | 124 | GLU |
| 29 | lc | 14 | HIS |
| 29 | lc | 97 | GLU |
| 29 | lc | 117 | PHE |
| 29 | lc | 141 | VAL |
| 31 | le | 27 | TYR |
| 31 | le | 40 | ASP |
| 31 | le | 51 | ASN |
| 32 | lf | 91 | MET |
| 32 | lf | 118 | ILE |
| 32 | lf | 141 | VAL |
| 33 | lg | 29 | VAL |
| 34 | lh | 8 | VAL |
| 35 | li | 12 | GLU |
| 35 | li | 47 | LEU |
| 35 | li | 85 | THR |
| 36 | lj | 79 | HIS |
| 37 | lk | 11 | ARG |
| 39 | lm | 8 | VAL |
| 39 | lm | 26 | ILE |
| 39 | lm | 42 | CYS |
| 40 | ln | 25 | VAL |
| 40 | ln | 44 | THR |
| 40 | ln | 67 | VAL |
| 41 | lo | 2 | THR |
| 41 | lo | 43 | HIS |
| 43 | lq | 33 | LEU |
| 44 | sA | 109 | TYR |
| 45 | sB | 30 | VAL |
| 45 | sB | 33 | ASP |
| 45 | sB | 45 | VAL |
| 45 | sB | 64 | LEU |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 46 | sC | 9 | LEU |
| 46 | sC | 30 | ASN |
| 46 | sC | 32 | TYR |
| 47 | sD | 30 | GLN |
| 47 | sD | 57 | THR |
| 47 | sD | 59 | LEU |
| 47 | sD | 68 | LEU |
| 48 | sE | 20 | THR |
| 48 | sE | 55 | LEU |
| 53 | sc | 106 | ILE |
| 53 | sc | 158 | CYS |
| 53 | sc | 160 | LEU |
| 53 | sc | 216 | THR |
| 54 | sd | 39 | LEU |
| 54 | sd | 123 | ILE |
| 54 | sd | 156 | LEU |
| 54 | sd | 178 | THR |
| 54 | sd | 189 | VAL |
| 55 | se | 26 | PHE |
| 55 | se | 112 | LEU |
| 55 | se | 200 | VAL |
| 56 | sf | 110 | VAL |
| 56 | sf | 171 | VAL |
| 56 | sf | 214 | ASN |
| 58 | sh | 70 | ASN |
| 58 | sh | 78 | ASP |
| 58 | sh | 97 | VAL |
| 59 | si | 121 | SER |
| 59 | si | 160 | THR |
| 60 | sj | 58 | LEU |
| 60 | sj | 96 | LEU |
| 63 | sm | 13 | VAL |
| 63 | sm | 61 | SER |
| 64 | so | 87 | ASP |
| 65 | sp | 39 | VAL |
| 65 | sp | 81 | LYS |
| 65 | sp | 126 | ASP |
| 67 | sr | 55 | THR |
| 67 | sr | 65 | LEU |
| 67 | sr | 98 | HIS |
| 68 | ss | 55 | ASN |
| 68 | ss | 124 | PHE |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 68 | ss | 135 | ASP |
| 69 | st | 69 | ILE |
| 70 | su | 77 | ASN |
| 70 | su | 80 | ILE |
| 70 | su | 137 | THR |
| 71 | sv | 42 | MET |
| 71 | sv | 43 | CYS |
| 73 | sx | 17 | VAL |
| 73 | sx | 21 | LEU |
| 73 | sx | 61 | ILE |
| 74 | sy | 31 | VAL |

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (67) such sidechains are listed below:

| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 4 | lD | 29 | HIS |
| 4 | lD | 140 | ASN |
| 5 | lE | 175 | GLN |
| 6 | lF | 50 | GLN |
| 6 | lF | 87 | ASN |
| 6 | lF | 149 | GLN |
| 6 | lF | 241 | ASN |
| 6 | lF | 322 | ASN |
| 6 | lF | 394 | ASN |
| 7 | lG | 118 | GLN |
| 7 | lG | 257 | HIS |
| 7 | lG | 270 | GLN |
| 8 | lH | 5 | GLN |
| 9 | lI | 187 | ASN |
| 10 | lJ | 77 | ASN |
| 10 | lJ | 171 | GLN |
| 11 | lK | 10 | HIS |
| 13 | lM | 119 | GLN |
| 13 | lM | 161 | ASN |
| 14 | lN | 57 | HIS |
| 14 | lN | 200 | GLN |
| 15 | lO | 43 | GLN |
| 17 | lQ | 32 | HIS |
| 17 | lQ | 156 | GLN |
| 18 | lR | 25 | HIS |
| 18 | lR | 63 | HIS |
| 18 | lR | 120 | GLN |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 19 | lS | 13 | HIS |
| 19 | lS | 21 | ASN |
| 19 | lS | 96 | ASN |
| 21 | lU | 58 | HIS |
| 22 | lV | 45 | ASN |
| 24 | lX | 12 | ASN |
| 25 | lY | 18 | ASN |
| 29 | lc | 7 | GLN |
| 29 | lc | 39 | HIS |
| 29 | lc | 40 | HIS |
| 30 | ld | 10 | HIS |
| 30 | ld | 14 | HIS |
| 33 | lg | 66 | ASN |
| 33 | lg | 99 | HIS |
| 34 | lh | 36 | GLN |
| 41 | lo | 4 | HIS |
| 45 | sB | 94 | ASN |
| 47 | sD | 30 | GLN |
| 47 | sD | 41 | GLN |
| 47 | sD | 45 | ASN |
| 48 | sE | 8 | ASN |
| 53 | sc | 112 | HIS |
| 54 | sd | 18 | ASN |
| 54 | sd | 61 | GLN |
| 54 | sd | 176 | ASN |
| 54 | sd | 180 | GLN |
| 56 | sf | 34 | HIS |
| 56 | sf | 51 | ASN |
| 56 | sf | 140 | HIS |
| 56 | sf | 248 | GLN |
| 61 | sk | 33 | GLN |
| 61 | sk | 82 | ASN |
| 62 | sl | 17 | ASN |
| 62 | sl | 31 | HIS |
| 63 | sm | 91 | HIS |
| 67 | sr | 120 | HIS |
| 70 | su | 102 | ASN |
| 71 | sv | 69 | ASN |
| 74 | sy | 59 | GLN |
| 74 | sy | 75 | ASN |

5.3.3 RNA ⓘ

| Mol | Chain | Analysed | Backbone Outliers | Pucker Outliers |
|-----|-------|-----------------|-------------------|-----------------|
| 1 | lA | 3127/3503 (89%) | 593 (18%) | 0 |
| 2 | lB | 143/155 (92%) | 31 (21%) | 0 |
| 3 | lC | 116/117 (99%) | 18 (15%) | 0 |
| 49 | sI | 70/76 (92%) | 30 (42%) | 0 |
| 50 | sJ | 71/77 (92%) | 30 (42%) | 0 |
| 51 | sK | 9/10 (90%) | 0 | 0 |
| 52 | sa | 1383/1947 (71%) | 305 (22%) | 0 |
| All | All | 4919/5885 (83%) | 1007 (20%) | 0 |

All (1007) RNA backbone outliers are listed below:

| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 1 | lA | 18 | G |
| 1 | lA | 22 | A |
| 1 | lA | 29 | U |
| 1 | lA | 30 | A |
| 1 | lA | 36 | A |
| 1 | lA | 39 | A |
| 1 | lA | 45 | A |
| 1 | lA | 53 | A |
| 1 | lA | 54 | G |
| 1 | lA | 55 | G |
| 1 | lA | 56 | A |
| 1 | lA | 62 | A |
| 1 | lA | 65 | U |
| 1 | lA | 70 | A |
| 1 | lA | 73 | A |
| 1 | lA | 88 | G |
| 1 | lA | 105 | A |
| 1 | lA | 106 | G |
| 1 | lA | 107 | A |
| 1 | lA | 108 | C |
| 1 | lA | 115 | G |
| 1 | lA | 116 | A |
| 1 | lA | 117 | U |
| 1 | lA | 185 | G |
| 1 | lA | 186 | A |
| 1 | lA | 189 | G |
| 1 | lA | 198 | A |
| 1 | lA | 199 | G |
| 1 | lA | 200 | A |
| 1 | lA | 206 | U |
| 1 | lA | 213 | A |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 1 | lA | 214 | A |
| 1 | lA | 221 | U |
| 1 | lA | 233 | A |
| 1 | lA | 237 | C |
| 1 | lA | 243 | G |
| 1 | lA | 246 | U |
| 1 | lA | 256 | G |
| 1 | lA | 257 | A |
| 1 | lA | 259 | A |
| 1 | lA | 264 | G |
| 1 | lA | 265 | A |
| 1 | lA | 266 | A |
| 1 | lA | 267 | A |
| 1 | lA | 277 | A |
| 1 | lA | 285 | A |
| 1 | lA | 286 | U |
| 1 | lA | 287 | A |
| 1 | lA | 289 | C |
| 1 | lA | 293 | A |
| 1 | lA | 294 | A |
| 1 | lA | 302 | U |
| 1 | lA | 303 | G |
| 1 | lA | 311 | A |
| 1 | lA | 316 | G |
| 1 | lA | 332 | G |
| 1 | lA | 342 | A |
| 1 | lA | 344 | A |
| 1 | lA | 345 | A |
| 1 | lA | 346 | G |
| 1 | lA | 350 | A |
| 1 | lA | 369 | A |
| 1 | lA | 370 | A |
| 1 | lA | 376 | A |
| 1 | lA | 385 | C |
| 1 | lA | 396 | G |
| 1 | lA | 397 | A |
| 1 | lA | 416 | U |
| 1 | lA | 422 | G |
| 1 | lA | 423 | A |
| 1 | lA | 424 | A |
| 1 | lA | 425 | C |
| 1 | lA | 444 | A |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 1 | lA | 447 | G |
| 1 | lA | 448 | C |
| 1 | lA | 449 | U |
| 1 | lA | 451 | G |
| 1 | lA | 467 | A |
| 1 | lA | 474 | A |
| 1 | lA | 483 | G |
| 1 | lA | 494 | G |
| 1 | lA | 495 | A |
| 1 | lA | 496 | A |
| 1 | lA | 500 | G |
| 1 | lA | 501 | U |
| 1 | lA | 510 | A |
| 1 | lA | 511 | A |
| 1 | lA | 512 | G |
| 1 | lA | 517 | G |
| 1 | lA | 535 | A |
| 1 | lA | 536 | G |
| 1 | lA | 538 | A |
| 1 | lA | 539 | G |
| 1 | lA | 550 | A |
| 1 | lA | 562 | A |
| 1 | lA | 564 | A |
| 1 | lA | 565 | U |
| 1 | lA | 577 | G |
| 1 | lA | 583 | A |
| 1 | lA | 586 | A |
| 1 | lA | 588 | G |
| 1 | lA | 589 | A |
| 1 | lA | 610 | A |
| 1 | lA | 612 | A |
| 1 | lA | 613 | G |
| 1 | lA | 614 | U |
| 1 | lA | 615 | G |
| 1 | lA | 619 | A |
| 1 | lA | 620 | U |
| 1 | lA | 625 | A |
| 1 | lA | 626 | A |
| 1 | lA | 627 | A |
| 1 | lA | 630 | A |
| 1 | lA | 631 | G |
| 1 | lA | 632 | G |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 1 | lA | 637 | U |
| 1 | lA | 644 | A |
| 1 | lA | 645 | A |
| 1 | lA | 646 | U |
| 1 | lA | 647 | C |
| 1 | lA | 648 | U |
| 1 | lA | 649 | U |
| 1 | lA | 652 | U |
| 1 | lA | 658 | A |
| 1 | lA | 663 | G |
| 1 | lA | 675 | A |
| 1 | lA | 685 | A |
| 1 | lA | 696 | G |
| 1 | lA | 699 | A |
| 1 | lA | 701 | A |
| 1 | lA | 702 | A |
| 1 | lA | 703 | G |
| 1 | lA | 708 | U |
| 1 | lA | 709 | U |
| 1 | lA | 710 | A |
| 1 | lA | 714 | U |
| 1 | lA | 715 | A |
| 1 | lA | 720 | U |
| 1 | lA | 721 | A |
| 1 | lA | 722 | A |
| 1 | lA | 724 | A |
| 1 | lA | 725 | A |
| 1 | lA | 726 | A |
| 1 | lA | 727 | U |
| 1 | lA | 728 | U |
| 1 | lA | 733 | C |
| 1 | lA | 734 | C |
| 1 | lA | 740 | A |
| 1 | lA | 744 | A |
| 1 | lA | 745 | A |
| 1 | lA | 746 | A |
| 1 | lA | 747 | C |
| 1 | lA | 755 | G |
| 1 | lA | 770 | C |
| 1 | lA | 783 | A |
| 1 | lA | 794 | A |
| 1 | lA | 811 | A |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 1 | lA | 816 | G |
| 1 | lA | 817 | A |
| 1 | lA | 818 | A |
| 1 | lA | 825 | G |
| 1 | lA | 833 | U |
| 1 | lA | 834 | A |
| 1 | lA | 835 | A |
| 1 | lA | 841 | A |
| 1 | lA | 842 | G |
| 1 | lA | 849 | C |
| 1 | lA | 851 | A |
| 1 | lA | 853 | A |
| 1 | lA | 854 | A |
| 1 | lA | 855 | A |
| 1 | lA | 862 | G |
| 1 | lA | 871 | A |
| 1 | lA | 872 | A |
| 1 | lA | 873 | U |
| 1 | lA | 874 | A |
| 1 | lA | 881 | G |
| 1 | lA | 882 | C |
| 1 | lA | 889 | A |
| 1 | lA | 898 | U |
| 1 | lA | 903 | G |
| 1 | lA | 904 | A |
| 1 | lA | 905 | A |
| 1 | lA | 911 | U |
| 1 | lA | 915 | U |
| 1 | lA | 918 | G |
| 1 | lA | 925 | A |
| 1 | lA | 927 | A |
| 1 | lA | 934 | G |
| 1 | lA | 936 | A |
| 1 | lA | 960 | G |
| 1 | lA | 964 | G |
| 1 | lA | 968 | C |
| 1 | lA | 976 | G |
| 1 | lA | 979 | G |
| 1 | lA | 980 | A |
| 1 | lA | 993 | U |
| 1 | lA | 998 | U |
| 1 | lA | 999 | G |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | lA | 1013 | A |
| 1 | lA | 1026 | G |
| 1 | lA | 1027 | G |
| 1 | lA | 1028 | G |
| 1 | lA | 1033 | A |
| 1 | lA | 1035 | G |
| 1 | lA | 1036 | A |
| 1 | lA | 1043 | G |
| 1 | lA | 1056 | G |
| 1 | lA | 1063 | C |
| 1 | lA | 1073 | A |
| 1 | lA | 1078 | C |
| 1 | lA | 1092 | A |
| 1 | lA | 1096 | U |
| 1 | lA | 1097 | U |
| 1 | lA | 1098 | A |
| 1 | lA | 1099 | A |
| 1 | lA | 1112 | G |
| 1 | lA | 1119 | U |
| 1 | lA | 1120 | A |
| 1 | lA | 1127 | A |
| 1 | lA | 1166 | A |
| 1 | lA | 1168 | C |
| 1 | lA | 1182 | A |
| 1 | lA | 1183 | G |
| 1 | lA | 1184 | A |
| 1 | lA | 1197 | G |
| 1 | lA | 1210 | C |
| 1 | lA | 1211 | A |
| 1 | lA | 1219 | A |
| 1 | lA | 1220 | A |
| 1 | lA | 1221 | A |
| 1 | lA | 1222 | U |
| 1 | lA | 1242 | G |
| 1 | lA | 1248 | U |
| 1 | lA | 1256 | G |
| 1 | lA | 1268 | A |
| 1 | lA | 1275 | A |
| 1 | lA | 1280 | C |
| 1 | lA | 1284 | A |
| 1 | lA | 1292 | A |
| 1 | lA | 1306 | A |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | lA | 1312 | A |
| 1 | lA | 1314 | A |
| 1 | lA | 1315 | A |
| 1 | lA | 1316 | C |
| 1 | lA | 1325 | C |
| 1 | lA | 1343 | A |
| 1 | lA | 1344 | U |
| 1 | lA | 1345 | U |
| 1 | lA | 1409 | G |
| 1 | lA | 1410 | A |
| 1 | lA | 1411 | A |
| 1 | lA | 1417 | G |
| 1 | lA | 1431 | G |
| 1 | lA | 1433 | U |
| 1 | lA | 1440 | A |
| 1 | lA | 1441 | U |
| 1 | lA | 1450 | A |
| 1 | lA | 1455 | A |
| 1 | lA | 1475 | A |
| 1 | lA | 1476 | G |
| 1 | lA | 1477 | A |
| 1 | lA | 1478 | C |
| 1 | lA | 1480 | U |
| 1 | lA | 1486 | C |
| 1 | lA | 1495 | A |
| 1 | lA | 1499 | G |
| 1 | lA | 1502 | A |
| 1 | lA | 1503 | U |
| 1 | lA | 1505 | G |
| 1 | lA | 1516 | A |
| 1 | lA | 1517 | C |
| 1 | lA | 1519 | A |
| 1 | lA | 1521 | A |
| 1 | lA | 1534 | U |
| 1 | lA | 1553 | A |
| 1 | lA | 1568 | G |
| 1 | lA | 1569 | A |
| 1 | lA | 1570 | U |
| 1 | lA | 1571 | C |
| 1 | lA | 1580 | A |
| 1 | lA | 1584 | G |
| 1 | lA | 1589 | U |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | lA | 1603 | A |
| 1 | lA | 1604 | A |
| 1 | lA | 1606 | A |
| 1 | lA | 1622 | A |
| 1 | lA | 1625 | A |
| 1 | lA | 1649 | U |
| 1 | lA | 1652 | U |
| 1 | lA | 1664 | G |
| 1 | lA | 1674 | U |
| 1 | lA | 1677 | A |
| 1 | lA | 1690 | A |
| 1 | lA | 1698 | G |
| 1 | lA | 1729 | G |
| 1 | lA | 1735 | A |
| 1 | lA | 1737 | A |
| 1 | lA | 1738 | G |
| 1 | lA | 1744 | A |
| 1 | lA | 1753 | A |
| 1 | lA | 1754 | G |
| 1 | lA | 1818 | C |
| 1 | lA | 1821 | A |
| 1 | lA | 1822 | U |
| 1 | lA | 1825 | U |
| 1 | lA | 1838 | G |
| 1 | lA | 1864 | C |
| 1 | lA | 1865 | U |
| 1 | lA | 1897 | A |
| 1 | lA | 1898 | A |
| 1 | lA | 1907 | A |
| 1 | lA | 1913 | G |
| 1 | lA | 1926 | U |
| 1 | lA | 1928 | G |
| 1 | lA | 1936 | A |
| 1 | lA | 1937 | A |
| 1 | lA | 1938 | G |
| 1 | lA | 1958 | A |
| 1 | lA | 1959 | U |
| 1 | lA | 1961 | G |
| 1 | lA | 1963 | G |
| 1 | lA | 1965 | U |
| 1 | lA | 1975 | G |
| 1 | lA | 1983 | A |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | lA | 1984 | A |
| 1 | lA | 2040 | A |
| 1 | lA | 2043 | A |
| 1 | lA | 2044 | C |
| 1 | lA | 2049 | A |
| 1 | lA | 2050 | C |
| 1 | lA | 2051 | A |
| 1 | lA | 2067 | A |
| 1 | lA | 2079 | A |
| 1 | lA | 2080 | A |
| 1 | lA | 2081 | A |
| 1 | lA | 2082 | U |
| 1 | lA | 2095 | G |
| 1 | lA | 2108 | G |
| 1 | lA | 2132 | A |
| 1 | lA | 2133 | G |
| 1 | lA | 2178 | A |
| 1 | lA | 2179 | U |
| 1 | lA | 2190 | A |
| 1 | lA | 2193 | G |
| 1 | lA | 2197 | G |
| 1 | lA | 2198 | G |
| 1 | lA | 2207 | A |
| 1 | lA | 2215 | A |
| 1 | lA | 2216 | U |
| 1 | lA | 2218 | A |
| 1 | lA | 2234 | A |
| 1 | lA | 2236 | A |
| 1 | lA | 2245 | A |
| 1 | lA | 2246 | G |
| 1 | lA | 2263 | G |
| 1 | lA | 2264 | A |
| 1 | lA | 2269 | U |
| 1 | lA | 2274 | A |
| 1 | lA | 2281 | U |
| 1 | lA | 2285 | U |
| 1 | lA | 2286 | G |
| 1 | lA | 2301 | U |
| 1 | lA | 2320 | A |
| 1 | lA | 2328 | A |
| 1 | lA | 2329 | G |
| 1 | lA | 2330 | U |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | lA | 2331 | A |
| 1 | lA | 2332 | A |
| 1 | lA | 2333 | C |
| 1 | lA | 2338 | A |
| 1 | lA | 2341 | C |
| 1 | lA | 2343 | C |
| 1 | lA | 2345 | U |
| 1 | lA | 2346 | A |
| 1 | lA | 2349 | G |
| 1 | lA | 2355 | A |
| 1 | lA | 2366 | C |
| 1 | lA | 2367 | A |
| 1 | lA | 2374 | U |
| 1 | lA | 2382 | C |
| 1 | lA | 2383 | G |
| 1 | lA | 2384 | C |
| 1 | lA | 2389 | A |
| 1 | lA | 2390 | U |
| 1 | lA | 2391 | G |
| 1 | lA | 2394 | U |
| 1 | lA | 2395 | C |
| 1 | lA | 2396 | A |
| 1 | lA | 2411 | G |
| 1 | lA | 2412 | U |
| 1 | lA | 2432 | A |
| 1 | lA | 2439 | A |
| 1 | lA | 2441 | U |
| 1 | lA | 2449 | A |
| 1 | lA | 2450 | C |
| 1 | lA | 2451 | G |
| 1 | lA | 2459 | U |
| 1 | lA | 2461 | G |
| 1 | lA | 2462 | G |
| 1 | lA | 2469 | G |
| 1 | lA | 2470 | G |
| 1 | lA | 2473 | A |
| 1 | lA | 2478 | A |
| 1 | lA | 2479 | G |
| 1 | lA | 2480 | A |
| 1 | lA | 2487 | U |
| 1 | lA | 2494 | G |
| 1 | lA | 2509 | U |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | lA | 2513 | A |
| 1 | lA | 2514 | A |
| 1 | lA | 2589 | C |
| 1 | lA | 2594 | C |
| 1 | lA | 2595 | G |
| 1 | lA | 2597 | U |
| 1 | lA | 2598 | A |
| 1 | lA | 2601 | U |
| 1 | lA | 2629 | A |
| 1 | lA | 2637 | A |
| 1 | lA | 2651 | A |
| 1 | lA | 2655 | G |
| 1 | lA | 2656 | G |
| 1 | lA | 2657 | C |
| 1 | lA | 2663 | A |
| 1 | lA | 2676 | G |
| 1 | lA | 2677 | G |
| 1 | lA | 2684 | G |
| 1 | lA | 2689 | G |
| 1 | lA | 2696 | A |
| 1 | lA | 2705 | A |
| 1 | lA | 2706 | C |
| 1 | lA | 2707 | A |
| 1 | lA | 2722 | U |
| 1 | lA | 2726 | A |
| 1 | lA | 2727 | A |
| 1 | lA | 2742 | G |
| 1 | lA | 2744 | A |
| 1 | lA | 2747 | G |
| 1 | lA | 2759 | A |
| 1 | lA | 2760 | G |
| 1 | lA | 2761 | A |
| 1 | lA | 2764 | A |
| 1 | lA | 2766 | A |
| 1 | lA | 2773 | U |
| 1 | lA | 2774 | A |
| 1 | lA | 2784 | G |
| 1 | lA | 2795 | C |
| 1 | lA | 2796 | C |
| 1 | lA | 2797 | A |
| 1 | lA | 2798 | G |
| 1 | lA | 2799 | U |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | lA | 2819 | U |
| 1 | lA | 2822 | U |
| 1 | lA | 2823 | G |
| 1 | lA | 2825 | C |
| 1 | lA | 2832 | A |
| 1 | lA | 2842 | U |
| 1 | lA | 2843 | U |
| 1 | lA | 2855 | A |
| 1 | lA | 2858 | A |
| 1 | lA | 2859 | U |
| 1 | lA | 2860 | U |
| 1 | lA | 2864 | A |
| 1 | lA | 2865 | A |
| 1 | lA | 2866 | C |
| 1 | lA | 2867 | A |
| 1 | lA | 2868 | U |
| 1 | lA | 2869 | A |
| 1 | lA | 2876 | U |
| 1 | lA | 2877 | U |
| 1 | lA | 2882 | G |
| 1 | lA | 2889 | G |
| 1 | lA | 2890 | A |
| 1 | lA | 2900 | G |
| 1 | lA | 2902 | G |
| 1 | lA | 2909 | A |
| 1 | lA | 2920 | U |
| 1 | lA | 2925 | A |
| 1 | lA | 2926 | U |
| 1 | lA | 2939 | G |
| 1 | lA | 2943 | G |
| 1 | lA | 2944 | A |
| 1 | lA | 2946 | A |
| 1 | lA | 2953 | C |
| 1 | lA | 2959 | G |
| 1 | lA | 2960 | A |
| 1 | lA | 2977 | G |
| 1 | lA | 2981 | A |
| 1 | lA | 2985 | C |
| 1 | lA | 2987 | C |
| 1 | lA | 2988 | A |
| 1 | lA | 2993 | G |
| 1 | lA | 3006 | G |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | lA | 3009 | C |
| 1 | lA | 3010 | C |
| 1 | lA | 3014 | G |
| 1 | lA | 3015 | A |
| 1 | lA | 3018 | U |
| 1 | lA | 3030 | A |
| 1 | lA | 3032 | C |
| 1 | lA | 3041 | G |
| 1 | lA | 3066 | U |
| 1 | lA | 3070 | C |
| 1 | lA | 3073 | U |
| 1 | lA | 3078 | U |
| 1 | lA | 3079 | A |
| 1 | lA | 3084 | A |
| 1 | lA | 3085 | C |
| 1 | lA | 3090 | G |
| 1 | lA | 3114 | A |
| 1 | lA | 3126 | C |
| 1 | lA | 3133 | G |
| 1 | lA | 3141 | A |
| 1 | lA | 3142 | G |
| 1 | lA | 3143 | A |
| 1 | lA | 3144 | U |
| 1 | lA | 3151 | A |
| 1 | lA | 3166 | G |
| 1 | lA | 3175 | A |
| 1 | lA | 3200 | G |
| 1 | lA | 3203 | A |
| 1 | lA | 3210 | U |
| 1 | lA | 3212 | U |
| 1 | lA | 3213 | U |
| 1 | lA | 3223 | G |
| 1 | lA | 3232 | G |
| 1 | lA | 3240 | A |
| 1 | lA | 3246 | A |
| 1 | lA | 3247 | C |
| 1 | lA | 3254 | U |
| 1 | lA | 3259 | U |
| 1 | lA | 3264 | G |
| 1 | lA | 3270 | A |
| 1 | lA | 3277 | C |
| 1 | lA | 3278 | A |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | lA | 3284 | G |
| 1 | lA | 3285 | A |
| 1 | lA | 3286 | A |
| 1 | lA | 3287 | U |
| 1 | lA | 3299 | G |
| 1 | lA | 3314 | U |
| 1 | lA | 3327 | A |
| 1 | lA | 3330 | A |
| 1 | lA | 3332 | A |
| 1 | lA | 3334 | U |
| 1 | lA | 3335 | U |
| 1 | lA | 3336 | G |
| 1 | lA | 3337 | A |
| 1 | lA | 3338 | A |
| 1 | lA | 3344 | U |
| 1 | lA | 3345 | A |
| 1 | lA | 3351 | U |
| 1 | lA | 3353 | U |
| 1 | lA | 3359 | U |
| 1 | lA | 3360 | U |
| 1 | lA | 3361 | A |
| 1 | lA | 3383 | A |
| 1 | lA | 3384 | A |
| 1 | lA | 3385 | G |
| 1 | lA | 3398 | U |
| 1 | lA | 3399 | G |
| 1 | lA | 3405 | A |
| 1 | lA | 3406 | C |
| 1 | lA | 3409 | A |
| 1 | lA | 3410 | A |
| 1 | lA | 3413 | U |
| 1 | lA | 3414 | A |
| 1 | lA | 3419 | A |
| 1 | lA | 3430 | U |
| 1 | lA | 3431 | U |
| 1 | lA | 3433 | U |
| 1 | lA | 3451 | A |
| 1 | lA | 3454 | G |
| 1 | lA | 3462 | A |
| 1 | lA | 3466 | U |
| 1 | lA | 3468 | U |
| 1 | lA | 3469 | A |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | 1A | 3470 | G |
| 1 | 1A | 3472 | G |
| 1 | 1A | 3473 | U |
| 1 | 1A | 3475 | U |
| 1 | 1A | 3476 | C |
| 1 | 1A | 3477 | U |
| 1 | 1A | 3478 | U |
| 1 | 1A | 3479 | A |
| 1 | 1A | 3480 | A |
| 1 | 1A | 3487 | C |
| 1 | 1A | 3490 | U |
| 1 | 1A | 3491 | C |
| 1 | 1A | 3492 | G |
| 1 | 1A | 3498 | U |
| 1 | 1A | 3499 | G |
| 1 | 1A | 3500 | A |
| 2 | 1B | 5 | A |
| 2 | 1B | 34 | G |
| 2 | 1B | 36 | G |
| 2 | 1B | 39 | A |
| 2 | 1B | 41 | G |
| 2 | 1B | 51 | A |
| 2 | 1B | 53 | G |
| 2 | 1B | 54 | A |
| 2 | 1B | 61 | A |
| 2 | 1B | 64 | A |
| 2 | 1B | 65 | G |
| 2 | 1B | 72 | G |
| 2 | 1B | 73 | A |
| 2 | 1B | 74 | A |
| 2 | 1B | 77 | A |
| 2 | 1B | 79 | A |
| 2 | 1B | 80 | A |
| 2 | 1B | 81 | A |
| 2 | 1B | 82 | A |
| 2 | 1B | 86 | G |
| 2 | 1B | 92 | C |
| 2 | 1B | 102 | A |
| 2 | 1B | 103 | A |
| 2 | 1B | 104 | U |
| 2 | 1B | 106 | C |
| 2 | 1B | 109 | G |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 2 | lB | 110 | A |
| 2 | lB | 113 | G |
| 2 | lB | 131 | A |
| 2 | lB | 147 | C |
| 2 | lB | 150 | U |
| 3 | lC | 6 | G |
| 3 | lC | 17 | G |
| 3 | lC | 19 | U |
| 3 | lC | 22 | C |
| 3 | lC | 32 | A |
| 3 | lC | 37 | A |
| 3 | lC | 49 | C |
| 3 | lC | 52 | G |
| 3 | lC | 53 | C |
| 3 | lC | 55 | C |
| 3 | lC | 62 | U |
| 3 | lC | 63 | C |
| 3 | lC | 72 | C |
| 3 | lC | 73 | G |
| 3 | lC | 90 | G |
| 3 | lC | 96 | U |
| 3 | lC | 99 | U |
| 3 | lC | 109 | G |
| 49 | sI | 2 | G |
| 49 | sI | 3 | C |
| 49 | sI | 6 | C |
| 49 | sI | 9 | A |
| 49 | sI | 10 | G |
| 49 | sI | 13 | C |
| 49 | sI | 14 | A |
| 49 | sI | 15 | G |
| 49 | sI | 16 | G |
| 49 | sI | 18 | A |
| 49 | sI | 22 | G |
| 49 | sI | 23 | A |
| 49 | sI | 29 | C |
| 49 | sI | 31 | A |
| 49 | sI | 41 | G |
| 49 | sI | 44 | U |
| 49 | sI | 46 | G |
| 49 | sI | 49 | G |
| 49 | sI | 50 | C |

Continued on next page...

Continued from previous page...

| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 49 | sI | 53 | G |
| 49 | sI | 55 | U |
| 49 | sI | 56 | C |
| 49 | sI | 58 | A |
| 49 | sI | 68 | G |
| 49 | sI | 69 | G |
| 49 | sI | 70 | G |
| 49 | sI | 73 | A |
| 49 | sI | 74 | C |
| 49 | sI | 75 | C |
| 49 | sI | 76 | A |
| 50 | sJ | 6 | C |
| 50 | sJ | 8 | U |
| 50 | sJ | 12 | G |
| 50 | sJ | 14 | A |
| 50 | sJ | 15 | G |
| 50 | sJ | 16 | C |
| 50 | sJ | 17 | C |
| 50 | sJ | 18 | C |
| 50 | sJ | 19 | G |
| 50 | sJ | 20 | U |
| 50 | sJ | 21 | U |
| 50 | sJ | 22 | A |
| 50 | sJ | 23 | G |
| 50 | sJ | 24 | C |
| 50 | sJ | 45 | A |
| 50 | sJ | 47 | G |
| 50 | sJ | 48 | U |
| 50 | sJ | 49 | C |
| 50 | sJ | 50 | U |
| 50 | sJ | 52 | C |
| 50 | sJ | 53 | G |
| 50 | sJ | 60 | G |
| 50 | sJ | 61 | U |
| 50 | sJ | 63 | C |
| 50 | sJ | 65 | G |
| 50 | sJ | 66 | C |
| 50 | sJ | 68 | G |
| 50 | sJ | 69 | A |
| 50 | sJ | 76 | C |
| 50 | sJ | 77 | A |
| 52 | sa | 4 | C |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 52 | sa | 24 | U |
| 52 | sa | 26 | U |
| 52 | sa | 33 | A |
| 52 | sa | 41 | G |
| 52 | sa | 42 | A |
| 52 | sa | 44 | U |
| 52 | sa | 46 | A |
| 52 | sa | 56 | G |
| 52 | sa | 59 | U |
| 52 | sa | 61 | A |
| 52 | sa | 62 | G |
| 52 | sa | 64 | A |
| 52 | sa | 65 | U |
| 52 | sa | 103 | U |
| 52 | sa | 113 | A |
| 52 | sa | 114 | G |
| 52 | sa | 115 | U |
| 52 | sa | 123 | G |
| 52 | sa | 135 | C |
| 52 | sa | 136 | A |
| 52 | sa | 137 | A |
| 52 | sa | 139 | G |
| 52 | sa | 140 | A |
| 52 | sa | 141 | U |
| 52 | sa | 142 | A |
| 52 | sa | 146 | U |
| 52 | sa | 148 | G |
| 52 | sa | 150 | G |
| 52 | sa | 152 | A |
| 52 | sa | 153 | U |
| 52 | sa | 154 | G |
| 52 | sa | 160 | G |
| 52 | sa | 161 | A |
| 52 | sa | 163 | A |
| 52 | sa | 171 | A |
| 52 | sa | 173 | A |
| 52 | sa | 211 | U |
| 52 | sa | 213 | A |
| 52 | sa | 239 | U |
| 52 | sa | 243 | U |
| 52 | sa | 244 | U |
| 52 | sa | 246 | G |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 52 | sa | 249 | U |
| 52 | sa | 253 | A |
| 52 | sa | 257 | C |
| 52 | sa | 258 | A |
| 52 | sa | 259 | A |
| 52 | sa | 260 | U |
| 52 | sa | 262 | G |
| 52 | sa | 263 | U |
| 52 | sa | 285 | G |
| 52 | sa | 288 | A |
| 52 | sa | 289 | C |
| 52 | sa | 293 | U |
| 52 | sa | 294 | G |
| 52 | sa | 295 | A |
| 52 | sa | 296 | G |
| 52 | sa | 299 | U |
| 52 | sa | 310 | U |
| 52 | sa | 311 | C |
| 52 | sa | 313 | A |
| 52 | sa | 317 | G |
| 52 | sa | 332 | G |
| 52 | sa | 333 | A |
| 52 | sa | 341 | G |
| 52 | sa | 345 | A |
| 52 | sa | 347 | A |
| 52 | sa | 354 | A |
| 52 | sa | 355 | A |
| 52 | sa | 356 | C |
| 52 | sa | 385 | G |
| 52 | sa | 395 | G |
| 52 | sa | 396 | A |
| 52 | sa | 397 | U |
| 52 | sa | 399 | G |
| 52 | sa | 411 | A |
| 52 | sa | 413 | G |
| 52 | sa | 417 | G |
| 52 | sa | 419 | C |
| 52 | sa | 420 | A |
| 52 | sa | 421 | G |
| 52 | sa | 429 | G |
| 52 | sa | 432 | A |
| 52 | sa | 434 | U |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 52 | sa | 439 | C |
| 52 | sa | 440 | A |
| 52 | sa | 456 | G |
| 52 | sa | 463 | A |
| 52 | sa | 470 | A |
| 52 | sa | 545 | G |
| 52 | sa | 549 | A |
| 52 | sa | 551 | G |
| 52 | sa | 553 | C |
| 52 | sa | 555 | G |
| 52 | sa | 559 | C |
| 52 | sa | 561 | A |
| 52 | sa | 562 | G |
| 52 | sa | 568 | G |
| 52 | sa | 572 | U |
| 52 | sa | 576 | U |
| 52 | sa | 579 | A |
| 52 | sa | 584 | C |
| 52 | sa | 588 | A |
| 52 | sa | 589 | G |
| 52 | sa | 600 | A |
| 52 | sa | 602 | U |
| 52 | sa | 604 | G |
| 52 | sa | 613 | A |
| 52 | sa | 614 | A |
| 52 | sa | 616 | A |
| 52 | sa | 617 | C |
| 52 | sa | 618 | G |
| 52 | sa | 628 | G |
| 52 | sa | 742 | G |
| 52 | sa | 746 | A |
| 52 | sa | 752 | A |
| 52 | sa | 756 | G |
| 52 | sa | 757 | U |
| 52 | sa | 759 | U |
| 52 | sa | 762 | A |
| 52 | sa | 782 | A |
| 52 | sa | 783 | A |
| 52 | sa | 784 | U |
| 52 | sa | 785 | A |
| 52 | sa | 787 | U |
| 52 | sa | 790 | A |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 52 | sa | 800 | A |
| 52 | sa | 803 | G |
| 52 | sa | 804 | C |
| 52 | sa | 843 | A |
| 52 | sa | 844 | A |
| 52 | sa | 845 | A |
| 52 | sa | 856 | G |
| 52 | sa | 876 | G |
| 52 | sa | 877 | G |
| 52 | sa | 879 | G |
| 52 | sa | 894 | G |
| 52 | sa | 904 | A |
| 52 | sa | 913 | A |
| 52 | sa | 915 | A |
| 52 | sa | 922 | G |
| 52 | sa | 924 | A |
| 52 | sa | 925 | U |
| 52 | sa | 927 | U |
| 52 | sa | 939 | G |
| 52 | sa | 940 | U |
| 52 | sa | 946 | A |
| 52 | sa | 950 | A |
| 52 | sa | 968 | A |
| 52 | sa | 972 | A |
| 52 | sa | 973 | A |
| 52 | sa | 984 | U |
| 52 | sa | 985 | A |
| 52 | sa | 1005 | A |
| 52 | sa | 1008 | C |
| 52 | sa | 1009 | G |
| 52 | sa | 1012 | G |
| 52 | sa | 1019 | A |
| 52 | sa | 1091 | C |
| 52 | sa | 1106 | A |
| 52 | sa | 1118 | G |
| 52 | sa | 1122 | U |
| 52 | sa | 1123 | U |
| 52 | sa | 1135 | G |
| 52 | sa | 1140 | G |
| 52 | sa | 1164 | A |
| 52 | sa | 1169 | A |
| 52 | sa | 1176 | G |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 52 | sa | 1177 | A |
| 52 | sa | 1184 | C |
| 52 | sa | 1185 | A |
| 52 | sa | 1192 | G |
| 52 | sa | 1210 | U |
| 52 | sa | 1219 | A |
| 52 | sa | 1221 | A |
| 52 | sa | 1224 | G |
| 52 | sa | 1225 | G |
| 52 | sa | 1227 | A |
| 52 | sa | 1232 | U |
| 52 | sa | 1233 | A |
| 52 | sa | 1239 | A |
| 52 | sa | 1242 | G |
| 52 | sa | 1243 | A |
| 52 | sa | 1253 | G |
| 52 | sa | 1254 | G |
| 52 | sa | 1260 | C |
| 52 | sa | 1267 | A |
| 52 | sa | 1269 | A |
| 52 | sa | 1270 | G |
| 52 | sa | 1271 | U |
| 52 | sa | 1274 | U |
| 52 | sa | 1276 | U |
| 52 | sa | 1277 | C |
| 52 | sa | 1281 | A |
| 52 | sa | 1282 | U |
| 52 | sa | 1283 | U |
| 52 | sa | 1284 | U |
| 52 | sa | 1287 | U |
| 52 | sa | 1290 | G |
| 52 | sa | 1299 | C |
| 52 | sa | 1310 | U |
| 52 | sa | 1311 | U |
| 52 | sa | 1312 | A |
| 52 | sa | 1316 | G |
| 52 | sa | 1317 | G |
| 52 | sa | 1323 | U |
| 52 | sa | 1326 | U |
| 52 | sa | 1331 | C |
| 52 | sa | 1332 | A |
| 52 | sa | 1337 | A |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 52 | sa | 1339 | U |
| 52 | sa | 1344 | G |
| 52 | sa | 1346 | A |
| 52 | sa | 1365 | U |
| 52 | sa | 1369 | A |
| 52 | sa | 1379 | C |
| 52 | sa | 1395 | A |
| 52 | sa | 1396 | A |
| 52 | sa | 1428 | U |
| 52 | sa | 1438 | A |
| 52 | sa | 1439 | C |
| 52 | sa | 1440 | A |
| 52 | sa | 1442 | A |
| 52 | sa | 1520 | C |
| 52 | sa | 1522 | U |
| 52 | sa | 1534 | A |
| 52 | sa | 1543 | A |
| 52 | sa | 1551 | A |
| 52 | sa | 1552 | G |
| 52 | sa | 1553 | A |
| 52 | sa | 1565 | G |
| 52 | sa | 1566 | C |
| 52 | sa | 1567 | A |
| 52 | sa | 1578 | A |
| 52 | sa | 1580 | U |
| 52 | sa | 1581 | G |
| 52 | sa | 1585 | U |
| 52 | sa | 1588 | C |
| 52 | sa | 1667 | U |
| 52 | sa | 1668 | G |
| 52 | sa | 1671 | A |
| 52 | sa | 1676 | U |
| 52 | sa | 1678 | A |
| 52 | sa | 1681 | A |
| 52 | sa | 1683 | A |
| 52 | sa | 1689 | A |
| 52 | sa | 1690 | A |
| 52 | sa | 1691 | A |
| 52 | sa | 1694 | A |
| 52 | sa | 1697 | U |
| 52 | sa | 1700 | A |
| 52 | sa | 1702 | G |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 52 | sa | 1703 | A |
| 52 | sa | 1706 | G |
| 52 | sa | 1708 | G |
| 52 | sa | 1711 | A |
| 52 | sa | 1712 | A |
| 52 | sa | 1713 | A |
| 52 | sa | 1714 | U |
| 52 | sa | 1717 | U |
| 52 | sa | 1723 | U |
| 52 | sa | 1725 | A |
| 52 | sa | 1734 | G |
| 52 | sa | 1735 | A |
| 52 | sa | 1736 | A |
| 52 | sa | 1750 | G |
| 52 | sa | 1767 | A |
| 52 | sa | 1773 | G |
| 52 | sa | 1777 | A |
| 52 | sa | 1780 | A |
| 52 | sa | 1781 | C |
| 52 | sa | 1782 | G |
| 52 | sa | 1785 | C |
| 52 | sa | 1788 | G |
| 52 | sa | 1797 | A |
| 52 | sa | 1800 | C |
| 52 | sa | 1801 | A |
| 52 | sa | 1821 | A |
| 52 | sa | 1822 | U |
| 52 | sa | 1824 | G |
| 52 | sa | 1825 | A |
| 52 | sa | 1830 | A |
| 52 | sa | 1831 | G |
| 52 | sa | 1832 | A |
| 52 | sa | 1846 | G |
| 52 | sa | 1849 | U |
| 52 | sa | 1850 | C |
| 52 | sa | 1852 | G |
| 52 | sa | 1883 | C |
| 52 | sa | 1885 | U |
| 52 | sa | 1893 | A |
| 52 | sa | 1894 | G |
| 52 | sa | 1902 | A |
| 52 | sa | 1907 | G |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 52 | sa | 1909 | A |
| 52 | sa | 1913 | A |
| 52 | sa | 1916 | U |
| 52 | sa | 1927 | G |
| 52 | sa | 1929 | A |
| 52 | sa | 1930 | C |
| 52 | sa | 1939 | G |
| 52 | sa | 1940 | G |
| 52 | sa | 1941 | A |
| 52 | sa | 1943 | C |

There are no RNA pucker outliers to report.

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

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

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

1 ligand is 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 | $\# Z > 2$ |
| 75 | PAR | sa | 5101 | - | 45,45,45 | 0.71 | 0 | 64,67,67 | 0.83 | 0 |

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 |
|-----|------|-------|------|------|---------|------------|---------|
| 75 | PAR | sa | 5101 | - | - | 3/18/94/94 | 0/4/4/4 |

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

All (3) torsion outliers are listed below:

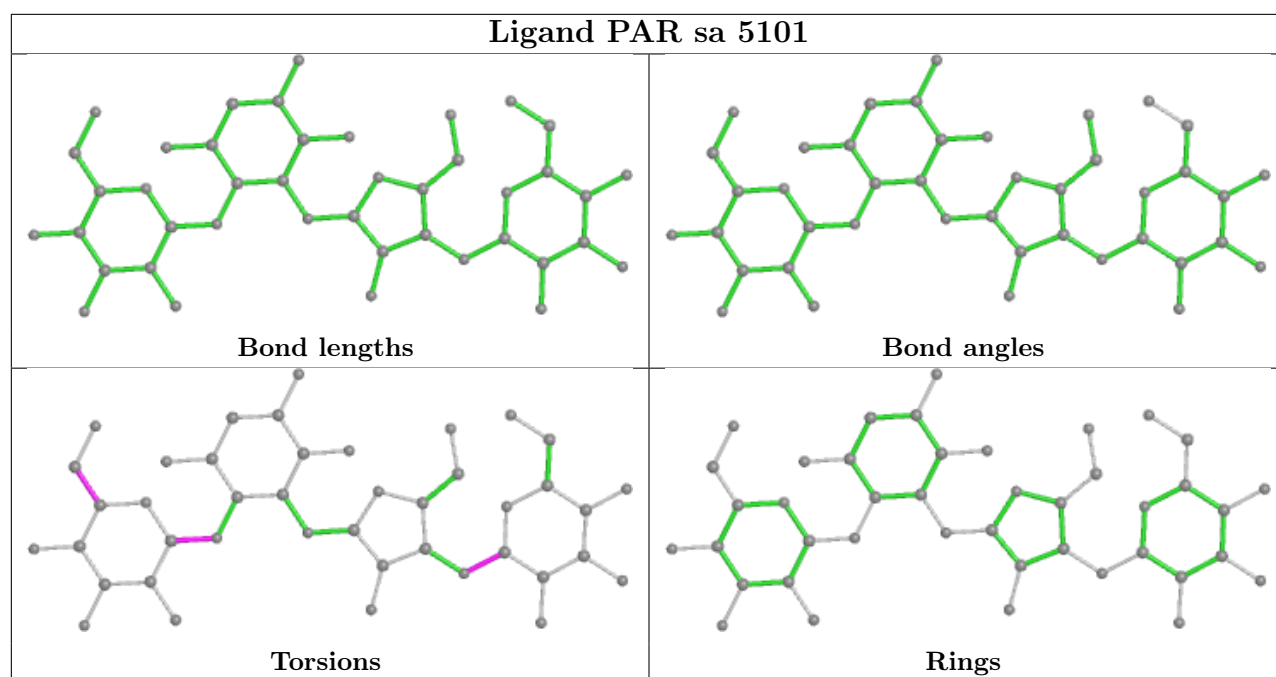
| Mol | Chain | Res | Type | Atoms |
|-----|-------|------|------|-----------------|
| 75 | sa | 5101 | PAR | O54-C14-O33-C33 |
| 75 | sa | 5101 | PAR | O51-C51-C61-O61 |
| 75 | sa | 5101 | PAR | O51-C11-O11-C42 |

There are no ring outliers.

1 monomer is involved in 1 short contact:

| Mol | Chain | Res | Type | Clashes | Symm-Clashes |
|-----|-------|------|------|---------|--------------|
| 75 | sa | 5101 | PAR | 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.



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.

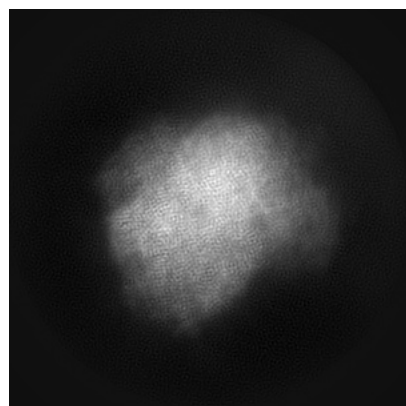
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-64718. These allow visual inspection of the internal detail of the map and identification of artifacts.

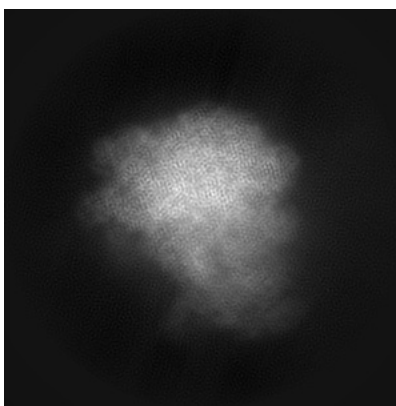
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

6.1 Orthogonal projections [i](#)

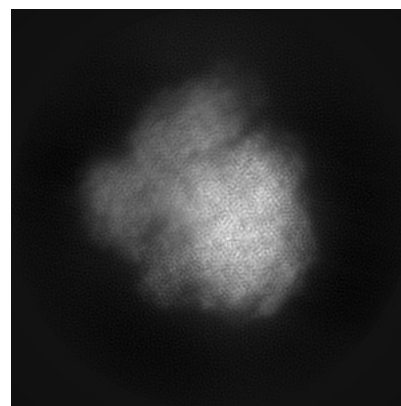
6.1.1 Primary map



X

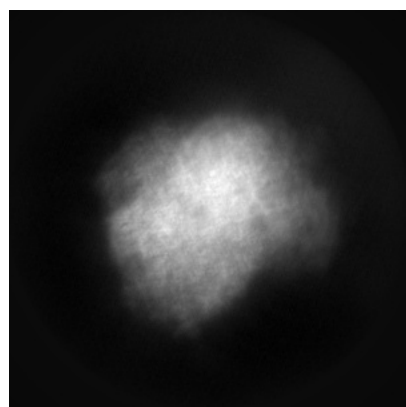


Y

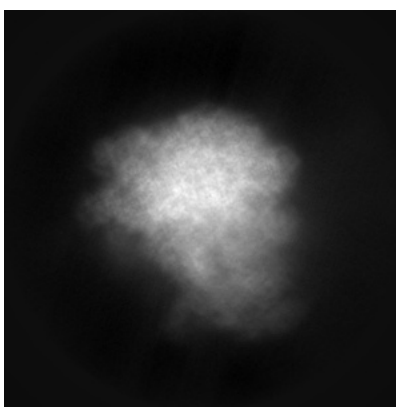


Z

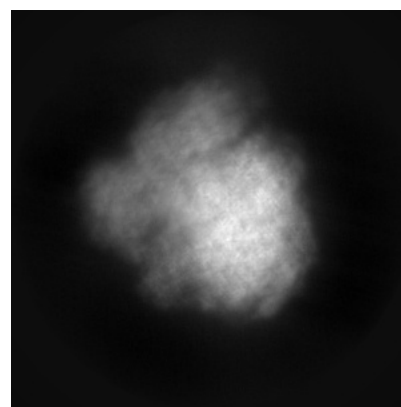
6.1.2 Raw map



X



Y

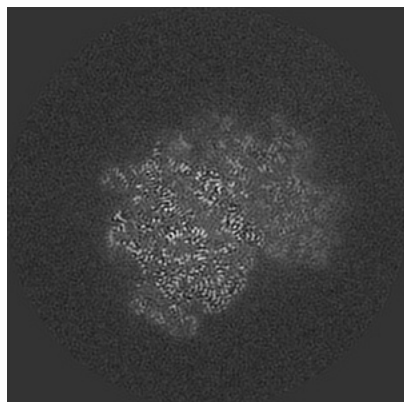


Z

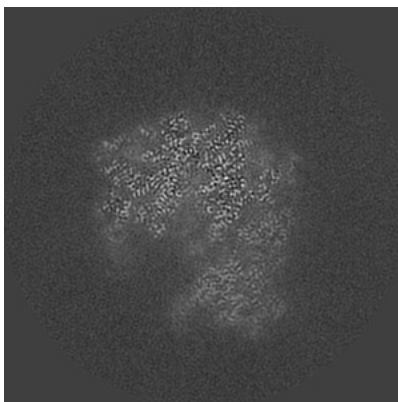
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

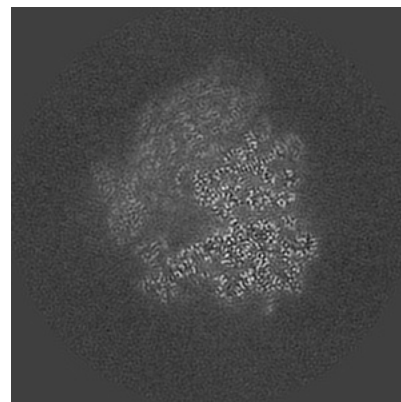
6.2.1 Primary map



X Index: 200

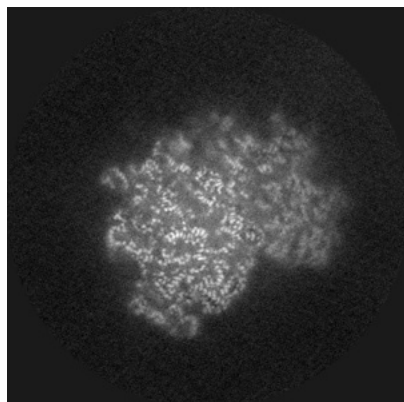


Y Index: 200

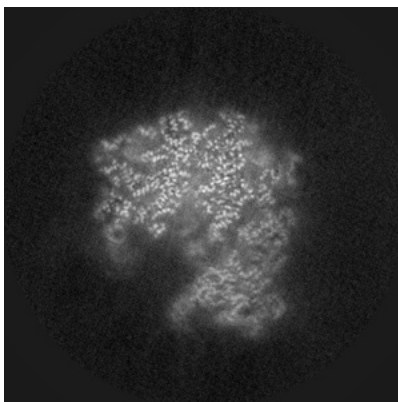


Z Index: 200

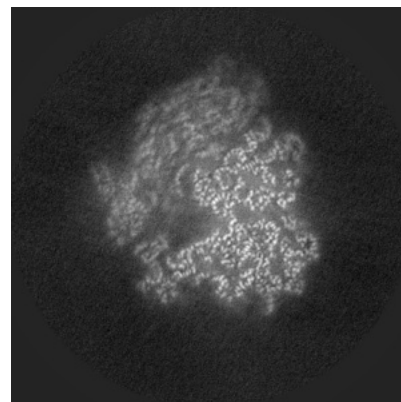
6.2.2 Raw map



X Index: 200



Y Index: 200

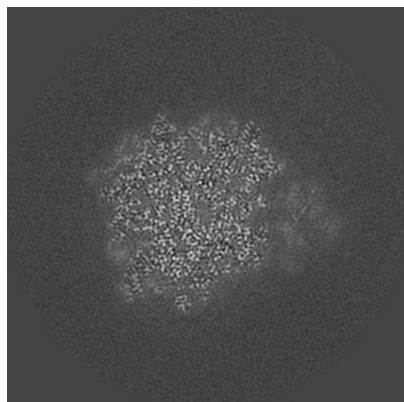


Z Index: 200

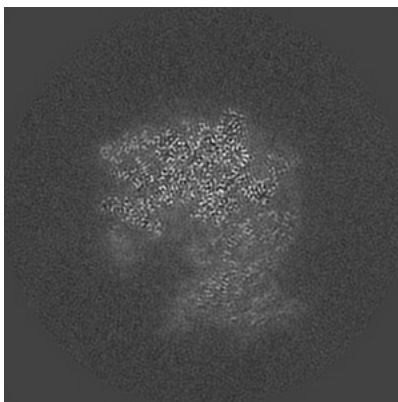
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

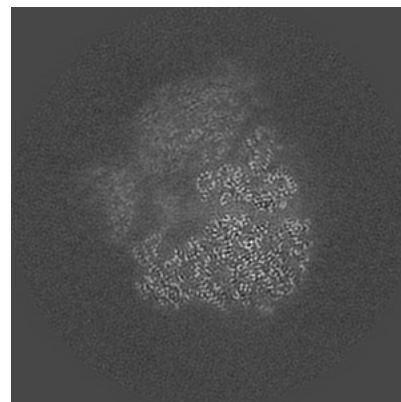
6.3.1 Primary map



X Index: 226

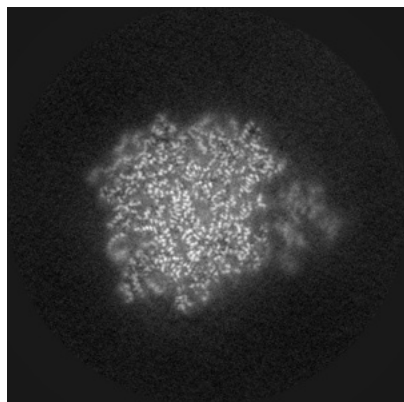


Y Index: 208

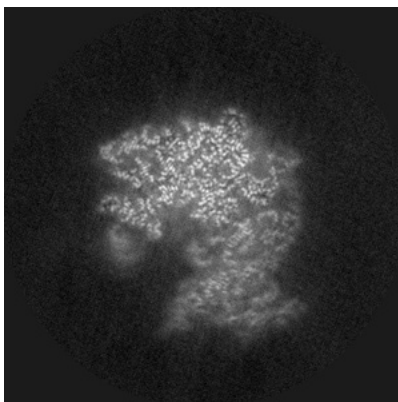


Z Index: 189

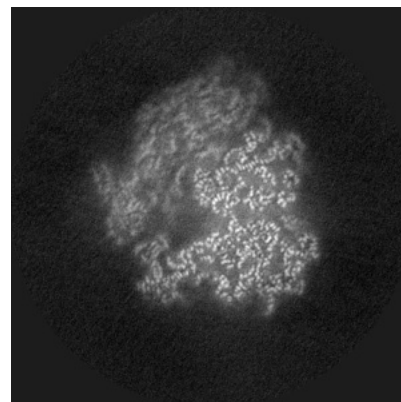
6.3.2 Raw map



X Index: 226



Y Index: 208

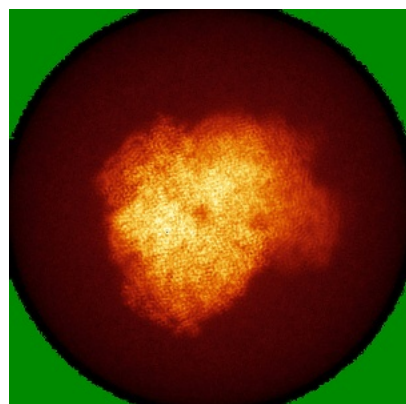


Z Index: 199

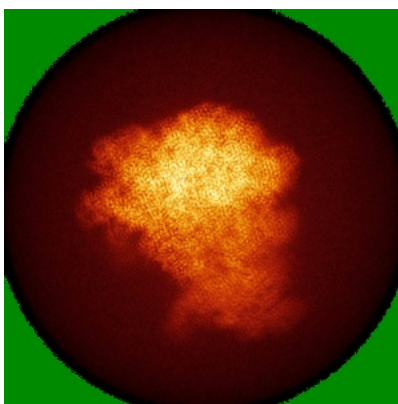
The images above show the largest variance slices of the map in three orthogonal directions.

6.4 Orthogonal standard-deviation projections (False-color) [i](#)

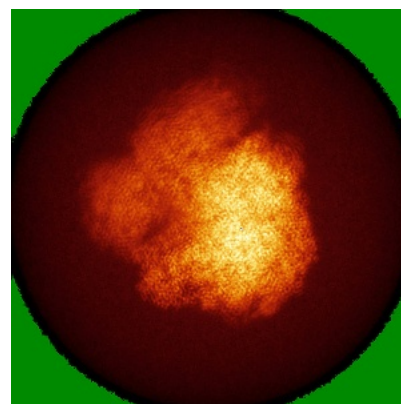
6.4.1 Primary map



X

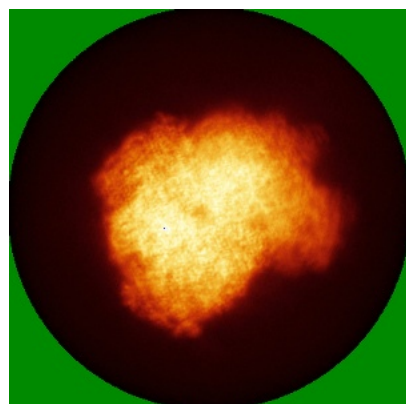


Y

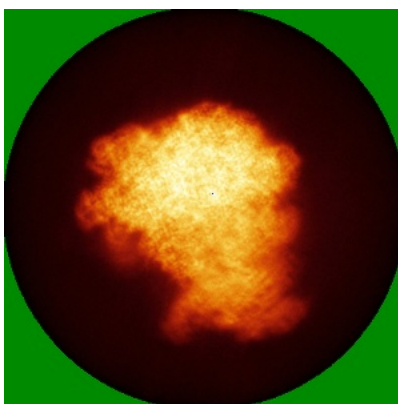


Z

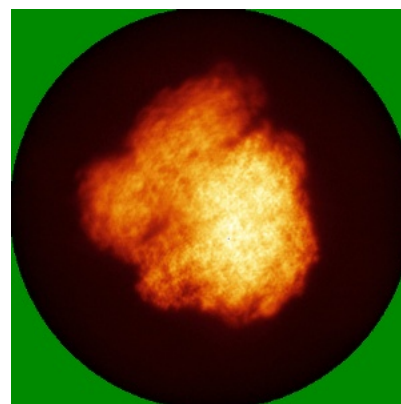
6.4.2 Raw map



X



Y



Z

The images above show the map standard deviation projections with false color in three orthogonal directions. Minimum values are shown in green, max in blue, and dark to light orange shades represent small to large values respectively.

6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 2.8. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.

6.5.2 Raw map



These images show the 3D surface of the raw map. The raw map's contour level was selected so that its surface encloses the same volume as the primary map does at its recommended contour level.

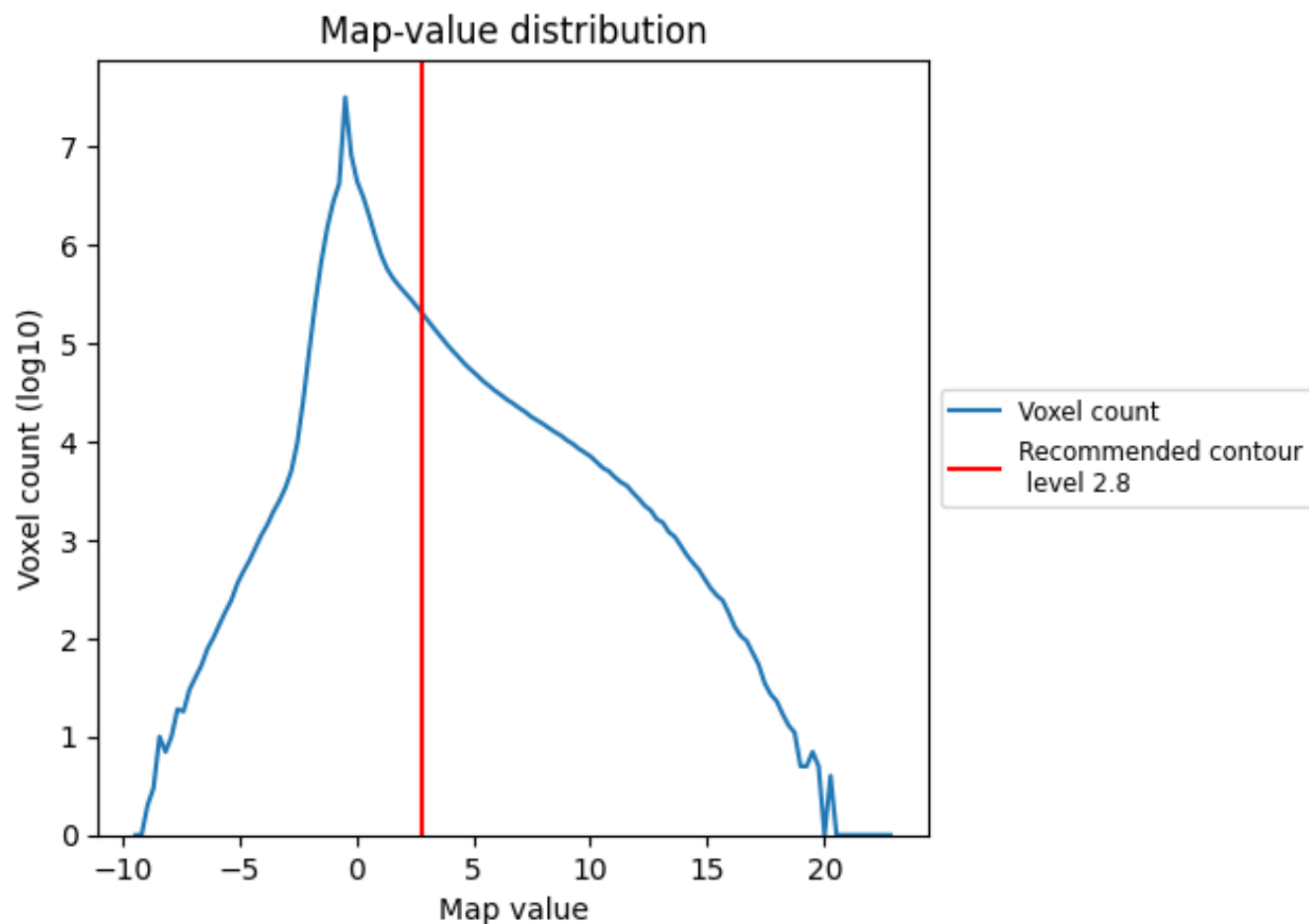
6.6 Mask visualisation [i](#)

This section was not generated. No masks/segmentation were deposited.

7 Map analysis [i](#)

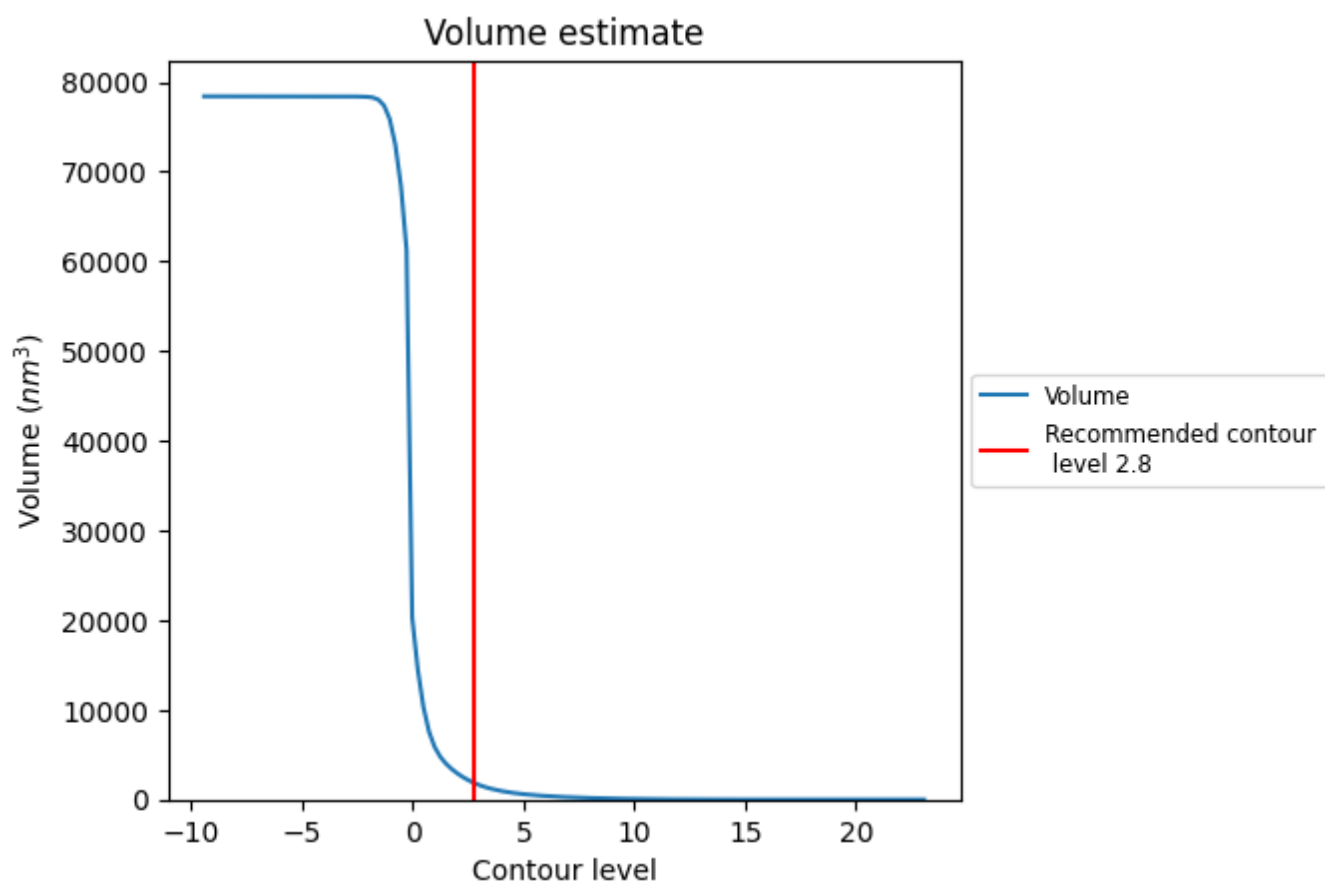
This section contains the results of statistical analysis of the map.

7.1 Map-value distribution [i](#)



The map-value distribution is plotted in 128 intervals along the x-axis. The y-axis is logarithmic. A spike in this graph at zero usually indicates that the volume has been masked.

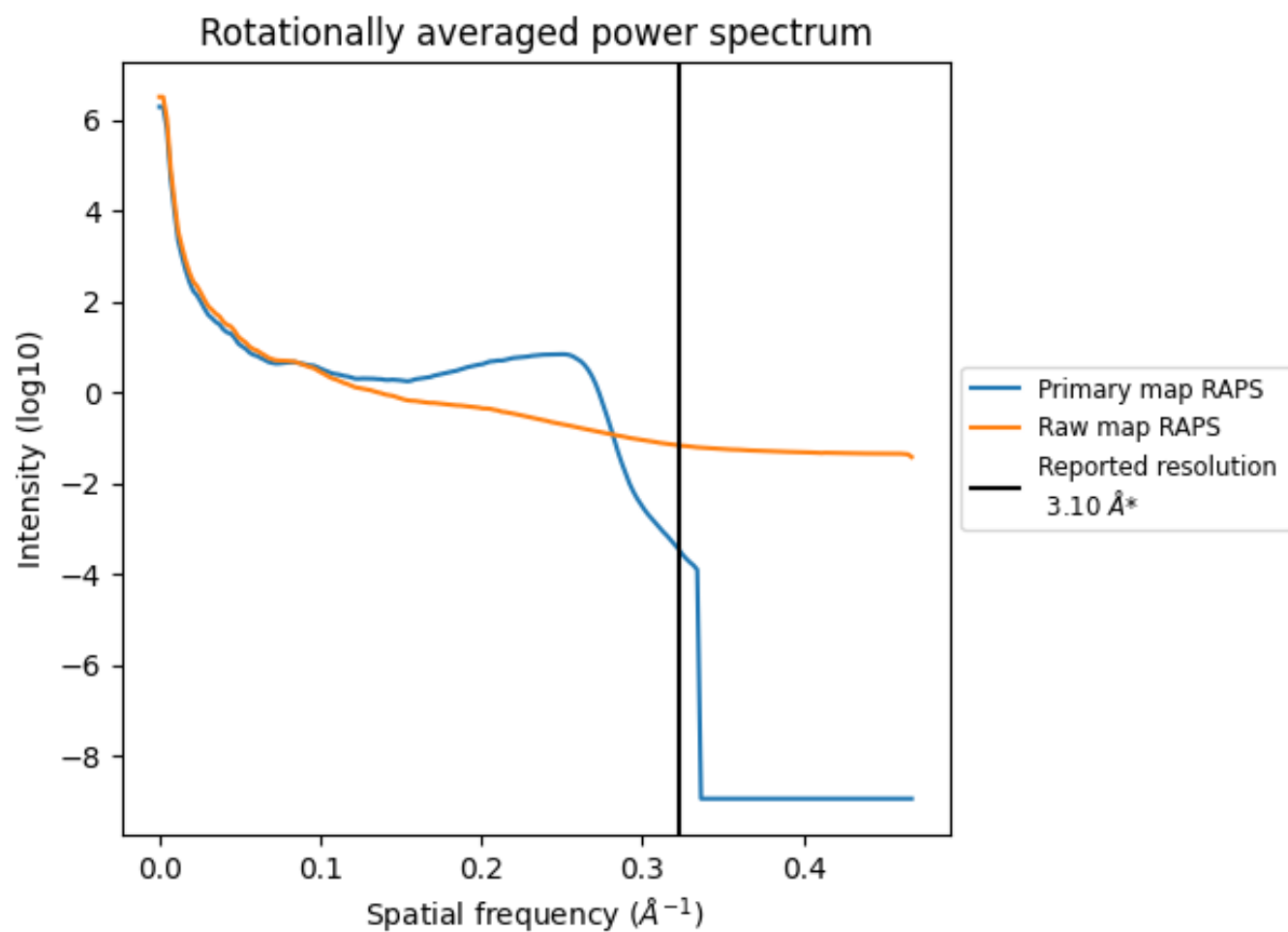
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 1820 nm^3 ; this corresponds to an approximate mass of 1644 kDa.

The volume estimate graph shows how the enclosed volume varies with the contour level. The recommended contour level is shown as a vertical line and the intersection between the line and the curve gives the volume of the enclosed surface at the given level.

7.3 Rotationally averaged power spectrum [i](#)

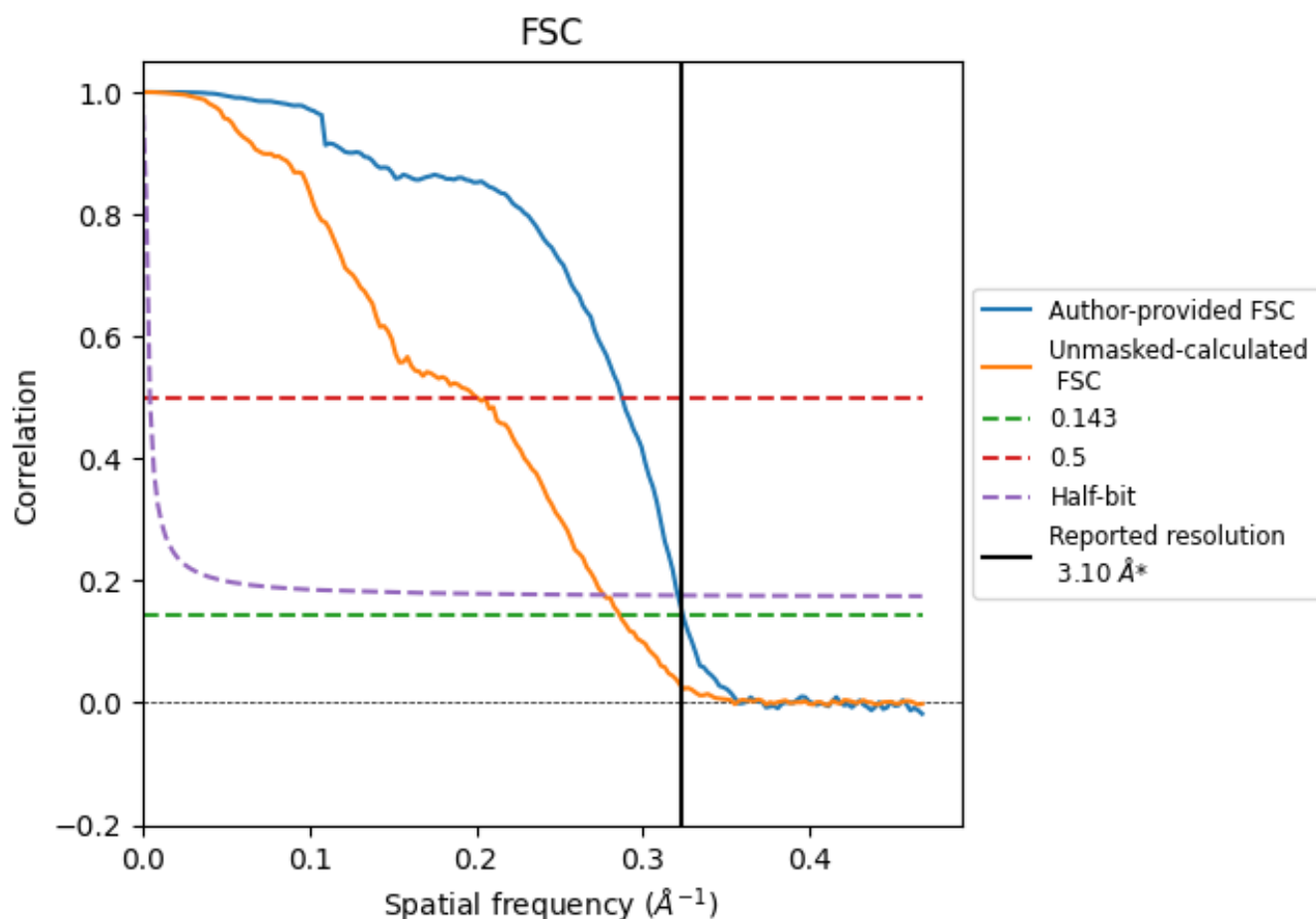


*Reported resolution corresponds to spatial frequency of 0.323 \AA^{-1}

8 Fourier-Shell correlation [i](#)

Fourier-Shell Correlation (FSC) is the most commonly used method to estimate the resolution of single-particle and subtomogram-averaged maps. The shape of the curve depends on the imposed symmetry, mask and whether or not the two 3D reconstructions used were processed from a common reference. The reported resolution is shown as a black line. A curve is displayed for the half-bit criterion in addition to lines showing the 0.143 gold standard cut-off and 0.5 cut-off.

8.1 FSC [i](#)



*Reported resolution corresponds to spatial frequency of 0.323 \AA^{-1}

8.2 Resolution estimates [i](#)

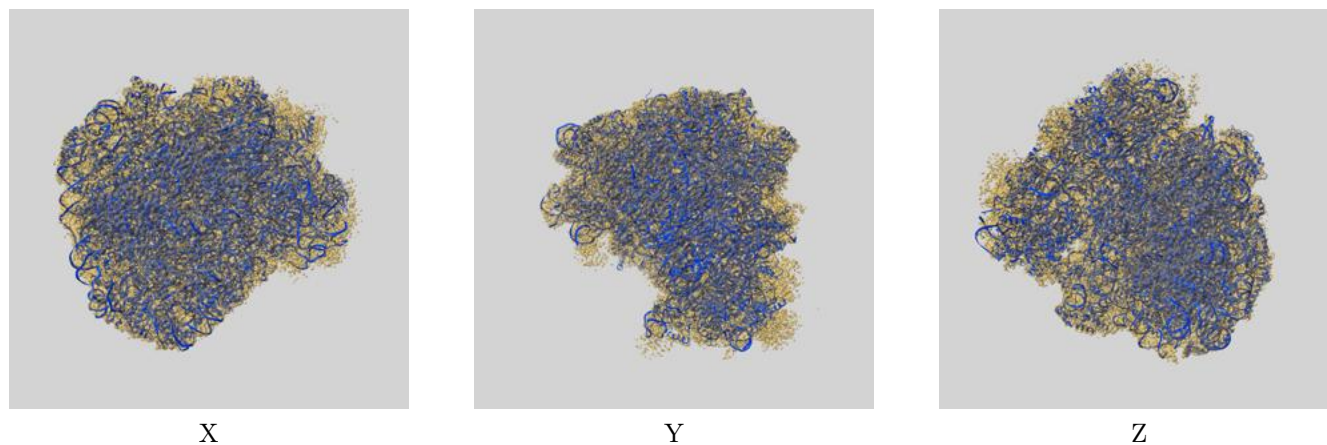
| Resolution estimate (Å) | Estimation criterion (FSC cut-off) | | |
|---------------------------|------------------------------------|------|----------|
| | 0.143 | 0.5 | Half-bit |
| Reported by author | 3.10 | - | - |
| Author-provided FSC curve | 3.09 | 3.48 | 3.12 |
| Unmasked-calculated* | 3.49 | 4.99 | 3.61 |

*Resolution estimate based on FSC curve calculated by comparison of deposited half-maps. The value from deposited half-maps intersecting FSC 0.143 CUT-OFF 3.49 differs from the reported value 3.1 by more than 10 %

9 Map-model fit [i](#)

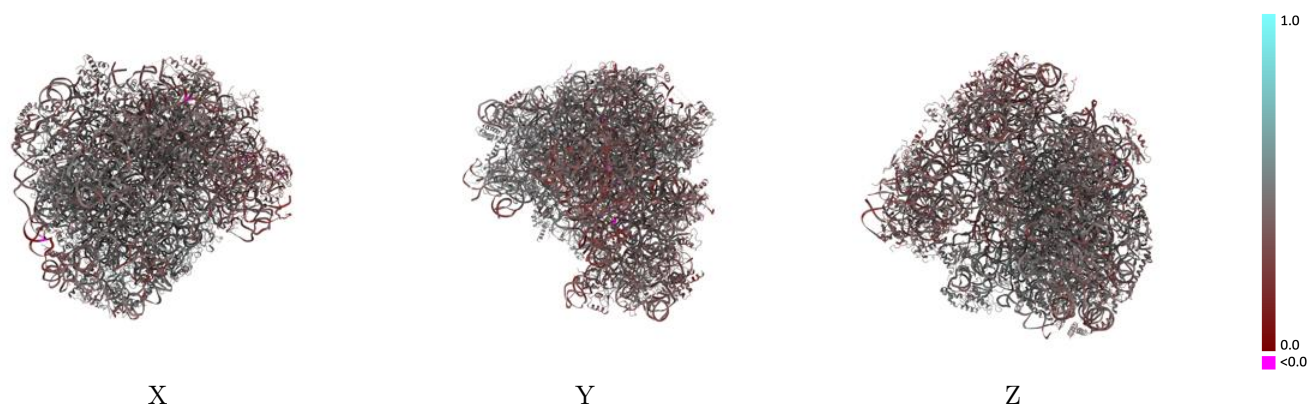
This section contains information regarding the fit between EMDB map EMD-64718 and PDB model 9V26. Per-residue inclusion information can be found in section [3](#) on page [18](#).

9.1 Map-model overlay [i](#)



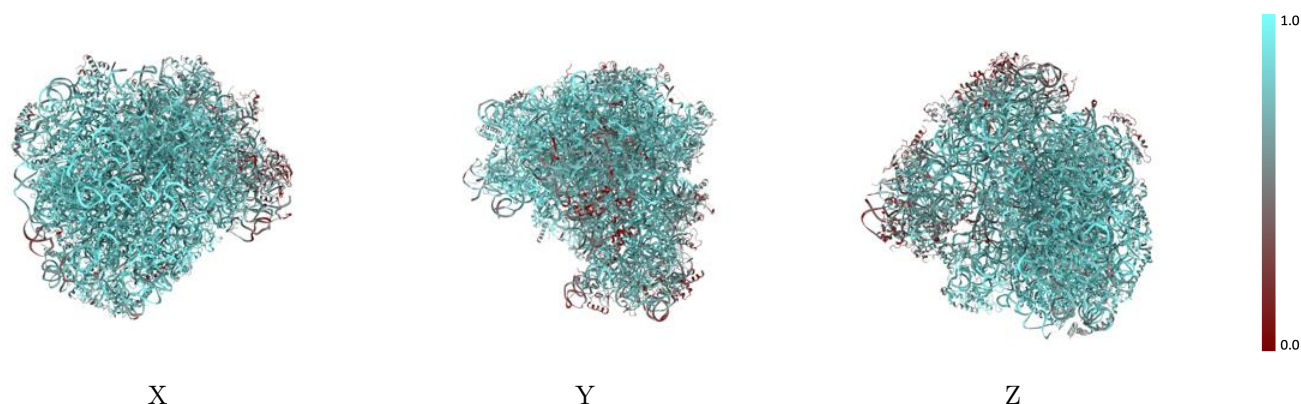
The images above show the 3D surface view of the map at the recommended contour level 2.8 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

9.2 Q-score mapped to coordinate model [i](#)



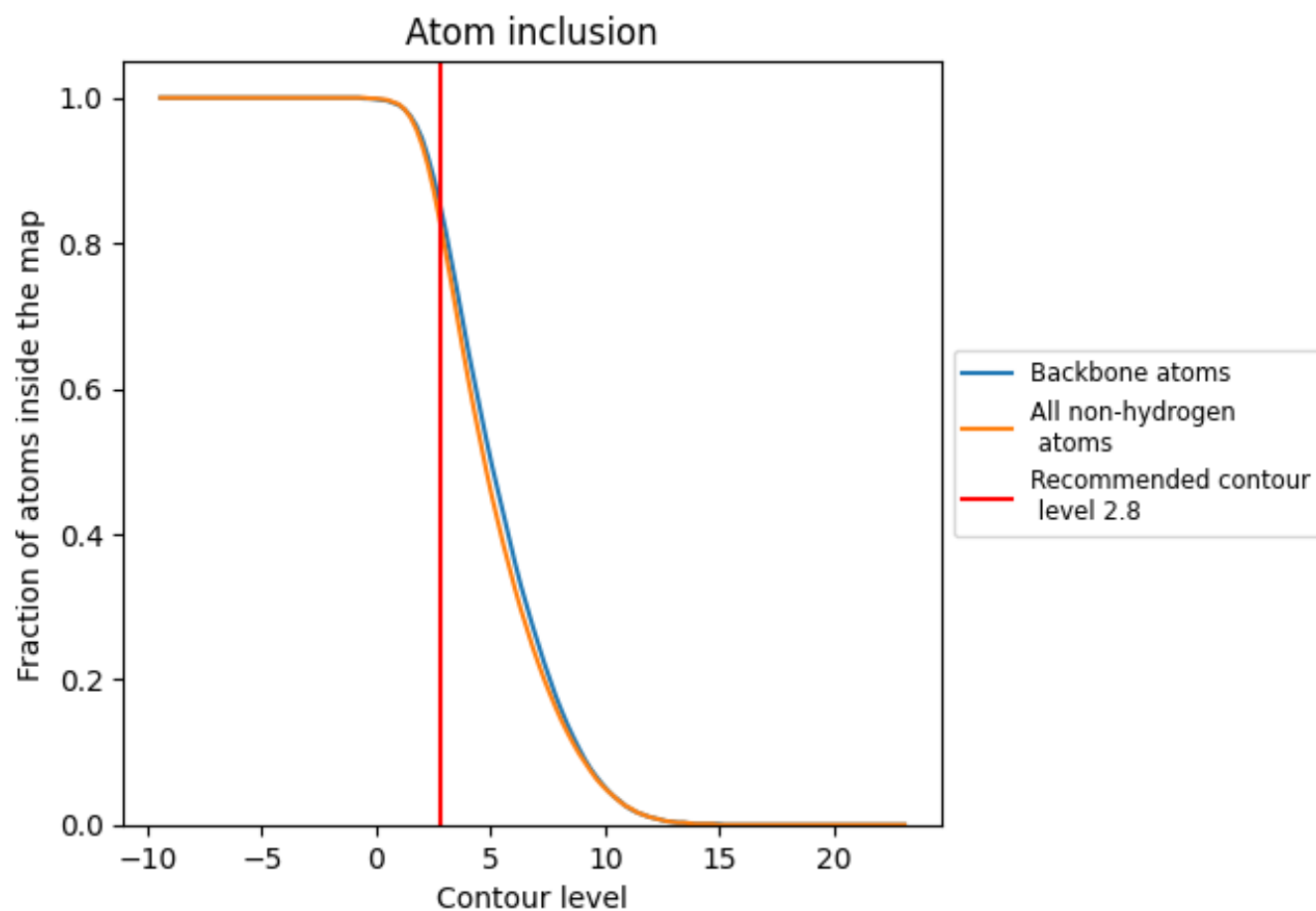
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions at which atoms can be resolved. Low Q-score values may therefore be expected for many entries.

9.3 Atom inclusion mapped to coordinate model [i](#)



The images above show the model with each residue coloured according to its atom inclusion. This shows to what extent they are inside the map at the recommended contour level (2.8).







































































9.4 Atom inclusion [i](#)



At the recommended contour level, 85% of all backbone atoms, 83% of all non-hydrogen atoms, are inside the map.

9.5 Map-model fit summary ⓘ

















































































The table lists the average atom inclusion at the recommended contour level (2.8) and Q-score for the entire model and for each chain.

| Chain | Atom inclusion | Q-score |
|-------|--|--|
| All |  0.8270 |  0.4130 |
| 1A |  0.9070 |  0.4200 |
| 1B |  0.9120 |  0.4080 |
| 1C |  0.9100 |  0.4230 |
| 1D |  0.9230 |  0.4450 |
| 1E |  0.8900 |  0.4610 |
| 1F |  0.8670 |  0.4670 |
| 1G |  0.8250 |  0.4750 |
| 1H |  0.6880 |  0.4270 |
| 1I |  0.8840 |  0.4840 |
| 1J |  0.7470 |  0.4160 |
| 1K |  0.8330 |  0.4650 |
| 1L |  0.8960 |  0.4660 |
| 1M |  0.7560 |  0.4500 |
| 1N |  0.7680 |  0.4370 |
| 1O |  0.8830 |  0.4700 |
| 1P |  0.8530 |  0.4830 |
| 1Q |  0.9420 |  0.4580 |
| 1R |  0.8720 |  0.4480 |
| 1S |  0.9280 |  0.4750 |
| 1T |  0.9250 |  0.4890 |
| 1U |  0.8370 |  0.4160 |
| 1V |  0.8980 |  0.4880 |
| 1W |  0.4810 |  0.3410 |
| 1X |  0.9260 |  0.4520 |
| 1Y |  0.8150 |  0.4240 |
| 1Z |  0.8800 |  0.4470 |
| 1a |  0.7930 |  0.4400 |
| 1b |  0.6850 |  0.3910 |
| 1c |  0.9270 |  0.4860 |
| 1d |  0.9320 |  0.4920 |
| 1e |  0.7350 |  0.4120 |
| 1f |  0.7430 |  0.3900 |
| 1g |  0.9260 |  0.4690 |
| 1h |  0.8770 |  0.4200 |



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| Chain | Atom inclusion | Q-score |
|-------|--|--|
| li |  0.8150 |  0.4280 |
| lj |  0.9170 |  0.4820 |
| lk |  0.8470 |  0.4520 |
| ll |  0.9480 |  0.4640 |
| lm |  0.8700 |  0.4310 |
| ln |  0.5970 |  0.3630 |
| lo |  0.9350 |  0.4390 |
| lp |  0.8810 |  0.4530 |
| lq |  0.9050 |  0.4670 |
| sA |  0.4610 |  0.3600 |
| sB |  0.8250 |  0.4330 |
| sC |  0.5320 |  0.3900 |
| sD |  0.5410 |  0.3950 |
| sE |  0.8100 |  0.4150 |
| sI |  0.6790 |  0.3060 |
| sJ |  0.7450 |  0.3500 |
| sK |  0.9580 |  0.4280 |
| sa |  0.7950 |  0.3650 |
| sc |  0.6840 |  0.4040 |
| sd |  0.5750 |  0.3790 |
| se |  0.6700 |  0.4030 |
| sf |  0.3170 |  0.3220 |
| sg |  0.5430 |  0.3810 |
| sh |  0.6080 |  0.3640 |
| si |  0.3450 |  0.3330 |
| sj |  0.5530 |  0.3520 |
| sk |  0.4740 |  0.3600 |
| sl |  0.4260 |  0.3510 |
| sm |  0.6350 |  0.3450 |
| so |  0.7490 |  0.3860 |
| sp |  0.7710 |  0.4280 |
| sq |  0.5660 |  0.3800 |
| sr |  0.7420 |  0.4050 |
| ss |  0.5220 |  0.3870 |
| st |  0.4880 |  0.3460 |
| su |  0.6230 |  0.4100 |
| sv |  0.4890 |  0.3920 |
| sw |  0.7330 |  0.4070 |
| sx |  0.5400 |  0.3940 |
| sy |  0.6470 |  0.3740 |