



Full wwPDB EM Validation Report ⓘ

May 20, 2026 – 10:31 PM JST

PDB ID : 9UZL / pdb_00009uzl
EMDB ID : EMD-64645
Title : EMCV IRES captured on mammalian 40S ribosome with initiator tRNA and eIF2
Authors : Das, D.; Hussain, T.
Deposited on : 2025-05-16
Resolution : 5.01 Å (reported)
Based on initial models : ., 6YAN, 8OZ0

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.dev132
MolProbity : 4-5-2 with Phenix2.0
Percentile statistics : 20250101.v01 (using entries in the PDB archive January 1st 2025)
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) : 2.49

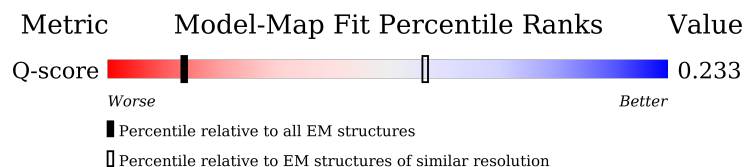
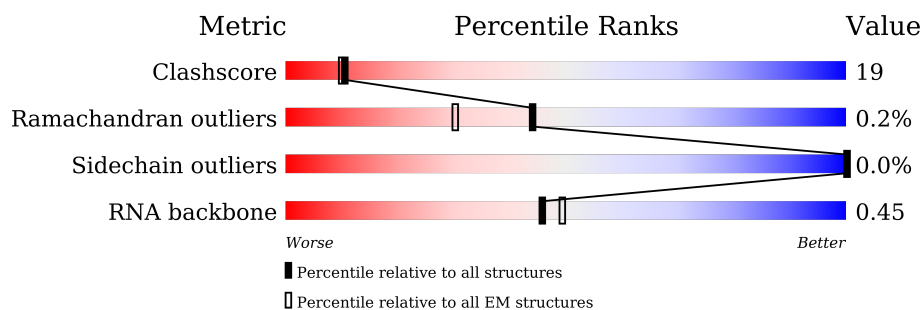
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

The reported resolution of this entry is 5.01 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.



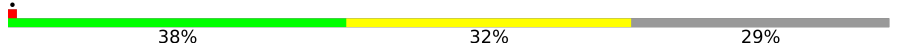

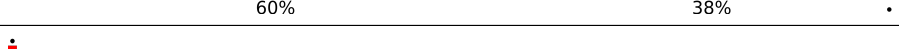
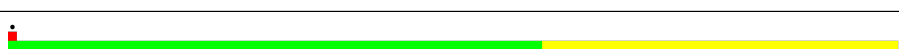



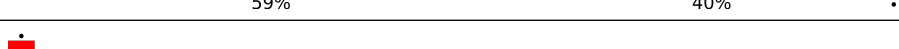





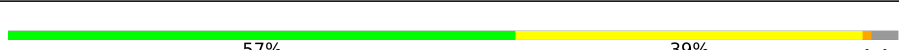


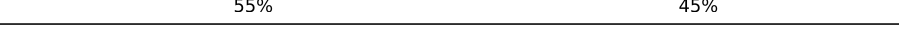







| Metric | Whole archive (#Entries) | EM structures (#Entries) | Similar EM resolution (#Entries, resolution range(Å)) |
|-----------------------|-----------------------------|-----------------------------|--|
| Clashscore | 229148 | 23984 | - |
| Ramachandran outliers | 224038 | 23583 | - |
| Sidechain outliers | 223484 | 23102 | - |
| RNA backbone | 8273 | 3508 | - |
| Q-score | - | 25397 | 859 (4.51 - 5.51) |

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 | 2 | 1863 | |
| 2 | A | 315 | |
| 3 | B | 485 | |

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| Mol | Chain | Length | Quality of chain |
|-----|-------|--------|--|
| 4 | C | 295 |  |
| 5 | D | 264 |  |
| 6 | E | 226 |  |
| 7 | F | 243 |  |
| 8 | G | 263 |  |
| 9 | H | 204 |  |
| 10 | I | 249 |  |
| 11 | J | 194 |  |
| 12 | K | 206 |  |
| 13 | L | 194 |  |
| 14 | M | 225 |  |
| 15 | N | 158 |  |
| 16 | O | 132 |  |
| 17 | P | 151 |  |
| 18 | Q | 168 |  |
| 19 | R | 145 |  |
| 20 | S | 146 |  |
| 21 | T | 135 |  |
| 22 | U | 152 |  |
| 23 | V | 141 |  |
| 24 | W | 119 |  |
| 25 | X | 82 |  |
| 26 | Y | 130 |  |
| 27 | Z | 143 |  |
| 28 | a | 126 | |

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| Mol | Chain | Length | Quality of chain |
|-----|-------|--------|--|
| 29 | b | 115 |  |
| 30 | c | 84 |  |
| 31 | d | 64 |  |
| 32 | e | 56 |  |
| 33 | f | 156 |  |
| 34 | g | 317 |  |
| 35 | h | 125 |  |
| 36 | i | 59 |  |
| 37 | l | 25 |  |
| 38 | y | 75 |  |
| 39 | z | 93 |  |

2 Entry composition [i](#)

There are 39 unique types of molecules in this entry. The entry contains 83705 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 18S ribosomal RNA.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-------|------|-------|------|---------|-------|
| 1 | 2 | 1744 | Total | C | N | O | P | 0 | 0 |
| | | | 37195 | 16608 | 6662 | 12185 | 1740 | | |

- Molecule 2 is a protein called Eukaryotic translation initiation factor 2 subunit 1.

| Mol | Chain | Residues | Atoms | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---------|-------|
| 2 | A | 278 | Total | C | N | O | 0 | 0 |
| | | | 1378 | 822 | 278 | 278 | | |

- Molecule 3 is a protein called Eukaryotic translation initiation factor 2 subunit 3.

| Mol | Chain | Residues | Atoms | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---------|-------|
| 3 | B | 464 | Total | C | N | O | 0 | 0 |
| | | | 2281 | 1352 | 464 | 465 | | |

- Molecule 4 is a protein called Small ribosomal subunit protein uS2.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 4 | C | 208 | Total | C | N | O | S | 0 | 0 |
| | | | 1642 | 1045 | 289 | 300 | 8 | | |

- Molecule 5 is a protein called Small ribosomal subunit protein eS1.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|----|---------|-------|
| 5 | D | 215 | Total | C | N | O | S | 0 | 0 |
| | | | 1741 | 1107 | 309 | 310 | 15 | | |

- Molecule 6 is a protein called Small ribosomal subunit protein uS5.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 6 | E | 222 | Total | C | N | O | S | 0 | 0 |
| | | | 1721 | 1114 | 295 | 303 | 9 | | |

- Molecule 7 is a protein called Small ribosomal subunit protein uS3.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 7 | F | 227 | Total | C | N | O | S | 0 | 0 |
| | | | 1764 | 1124 | 317 | 315 | 8 | | |

- Molecule 8 is a protein called Small ribosomal subunit protein eS4.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|----|---------|-------|
| 8 | G | 263 | Total | C | N | O | S | 0 | 0 |
| | | | 2083 | 1329 | 385 | 359 | 10 | | |

- Molecule 9 is a protein called Small ribosomal subunit protein uS7.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 9 | H | 191 | Total | C | N | O | S | 0 | 0 |
| | | | 1509 | 943 | 286 | 273 | 7 | | |

- Molecule 10 is a protein called Small ribosomal subunit protein eS6.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 10 | I | 237 | Total | C | N | O | S | 0 | 0 |
| | | | 1923 | 1200 | 387 | 329 | 7 | | |

- Molecule 11 is a protein called Small ribosomal subunit protein eS7.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 11 | J | 190 | Total | C | N | O | S | 0 | 0 |
| | | | 1530 | 975 | 281 | 273 | 1 | | |

- Molecule 12 is a protein called Small ribosomal subunit protein eS8.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 12 | K | 206 | Total | C | N | O | S | 0 | 0 |
| | | | 1680 | 1054 | 329 | 292 | 5 | | |

- Molecule 13 is a protein called Small ribosomal subunit protein uS4.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 13 | L | 182 | Total | C | N | O | S | 0 | 0 |
| | | | 1498 | 952 | 300 | 244 | 2 | | |

- Molecule 14 is a protein called Small ribosomal subunit protein eS10.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 14 | M | 98 | Total | C | N | O | S | 0 | 0 |
| | | | 827 | 539 | 148 | 134 | 6 | | |

There is a discrepancy between the modelled and reference sequences:

| Chain | Residue | Modelled | Actual | Comment | Reference |
|-------|---------|----------|--------|----------|----------------|
| M | -30 | THR | ALA | conflict | UNP A0AAG1W9A6 |

- Molecule 15 is a protein called Small ribosomal subunit protein uS17.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 15 | N | 158 | Total | C | N | O | S | 0 | 0 |
| | | | 1296 | 827 | 241 | 221 | 7 | | |

- Molecule 16 is a protein called Small ribosomal subunit protein eS12.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 16 | O | 124 | Total | C | N | O | S | 0 | 0 |
| | | | 958 | 600 | 170 | 179 | 9 | | |

- Molecule 17 is a protein called Small ribosomal subunit protein uS15.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 17 | P | 150 | Total | C | N | O | S | 0 | 0 |
| | | | 1208 | 773 | 229 | 205 | 1 | | |

- Molecule 18 is a protein called Ribosomal protein S14.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 18 | Q | 136 | Total | C | N | O | S | 0 | 0 |
| | | | 1016 | 621 | 199 | 190 | 6 | | |

- Molecule 19 is a protein called Small ribosomal subunit protein uS19.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 19 | R | 135 | Total | C | N | O | S | 0 | 0 |
| | | | 1111 | 704 | 211 | 189 | 7 | | |

- Molecule 20 is a protein called Small ribosomal subunit protein uS9.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 20 | S | 141 | Total | C | N | O | S | 0 | 0 |
| | | | 1123 | 715 | 212 | 193 | 3 | | |

- Molecule 21 is a protein called Small ribosomal subunit protein eS17.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 21 | T | 126 | Total | C | N | O | S | 0 | 0 |
| | | | 1019 | 639 | 188 | 187 | 5 | | |

- Molecule 22 is a protein called Small ribosomal subunit protein uS13.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 22 | U | 142 | Total | C | N | O | S | 0 | 0 |
| | | | 1172 | 733 | 239 | 199 | 1 | | |

- Molecule 23 is a protein called Small ribosomal subunit protein eS19.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 23 | V | 141 | Total | C | N | O | S | 0 | 0 |
| | | | 1113 | 701 | 213 | 196 | 3 | | |

- Molecule 24 is a protein called Small ribosomal subunit protein uS10.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 24 | W | 104 | Total | C | N | O | S | 0 | 0 |
| | | | 822 | 514 | 156 | 148 | 4 | | |

- Molecule 25 is a protein called Small ribosomal subunit protein eS21.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 25 | X | 82 | Total | C | N | O | S | 0 | 0 |
| | | | 620 | 378 | 117 | 120 | 5 | | |

There are 5 discrepancies between the modelled and reference sequences:

| Chain | Residue | Modelled | Actual | Comment | Reference |
|-------|---------|----------|--------|----------|------------|
| X | 3 | SER | ASN | conflict | UNP G1TM82 |
| X | 4 | ASN | ASP | conflict | UNP G1TM82 |
| X | 33 | PRO | GLN | conflict | UNP G1TM82 |
| X | 50 | SER | PHE | conflict | UNP G1TM82 |
| X | 76 | HIS | ASP | conflict | UNP G1TM82 |

- Molecule 26 is a protein called Small ribosomal subunit protein uS8.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 26 | Y | 129 | Total | C | N | O | S | 0 | 0 |
| | | | 1034 | 659 | 193 | 176 | 6 | | |

- Molecule 27 is a protein called Small ribosomal subunit protein uS12.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 27 | Z | 142 | Total | C | N | O | S | 0 | 0 |
| | | | 1106 | 698 | 220 | 184 | 4 | | |

- Molecule 28 is a protein called 40S ribosomal protein S24.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 28 | a | 126 | Total | C | N | O | S | 0 | 0 |
| | | | 1022 | 645 | 198 | 174 | 5 | | |

- Molecule 29 is a protein called Small ribosomal subunit protein eS26.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 29 | b | 99 | Total | C | N | O | S | 0 | 0 |
| | | | 789 | 491 | 162 | 130 | 6 | | |

- Molecule 30 is a protein called Small ribosomal subunit protein eS27.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 30 | c | 84 | Total | C | N | O | S | 0 | 0 |
| | | | 659 | 413 | 122 | 116 | 8 | | |

- Molecule 31 is a protein called Small ribosomal subunit protein eS28.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|----|---|---------|-------|
| 31 | d | 64 | Total | C | N | O | S | 0 | 0 |
| | | | 507 | 308 | 102 | 95 | 2 | | |

- Molecule 32 is a protein called Small ribosomal subunit protein uS14.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 32 | e | 53 | Total | C | N | O | S | 0 | 0 |
| | | | 445 | 278 | 90 | 72 | 5 | | |

- Molecule 33 is a protein called Ubiquitin-ribosomal protein eS31 fusion protein.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|----|---|---------|-------|
| 33 | f | 71 | Total | C | N | O | S | 0 | 0 |
| | | | 581 | 367 | 109 | 98 | 7 | | |

- Molecule 34 is a protein called Small ribosomal subunit protein RACK1.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|----|---------|-------|
| 34 | g | 313 | Total | C | N | O | S | 0 | 0 |
| | | | 2436 | 1535 | 424 | 465 | 12 | | |

- Molecule 35 is a protein called Small ribosomal subunit protein eS25.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 35 | h | 75 | Total | C | N | O | S | 0 | 0 |
| | | | 598 | 382 | 111 | 104 | 1 | | |

- Molecule 36 is a protein called Small ribosomal subunit protein eS30.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|----|---|---------|-------|
| 36 | i | 59 | Total | C | N | O | S | 0 | 0 |
| | | | 473 | 293 | 104 | 75 | 1 | | |

- Molecule 37 is a protein called Small ribosomal subunit protein eS32.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 37 | l | 25 | Total | C | N | O | S | 0 | 0 |
| | | | 240 | 145 | 64 | 28 | 3 | | |

- Molecule 38 is a RNA chain called Initiator tRNA.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|----|---------|-------|
| 38 | y | 75 | Total | C | N | O | P | 0 | 0 |
| | | | 1604 | 717 | 298 | 515 | 74 | | |

- Molecule 39 is a RNA chain called Encephalomyocarditis viral IRES.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|----|---------|-------|
| 39 | z | 93 | Total | C | N | O | P | 0 | 0 |
| | | | 1981 | 885 | 361 | 642 | 93 | | |

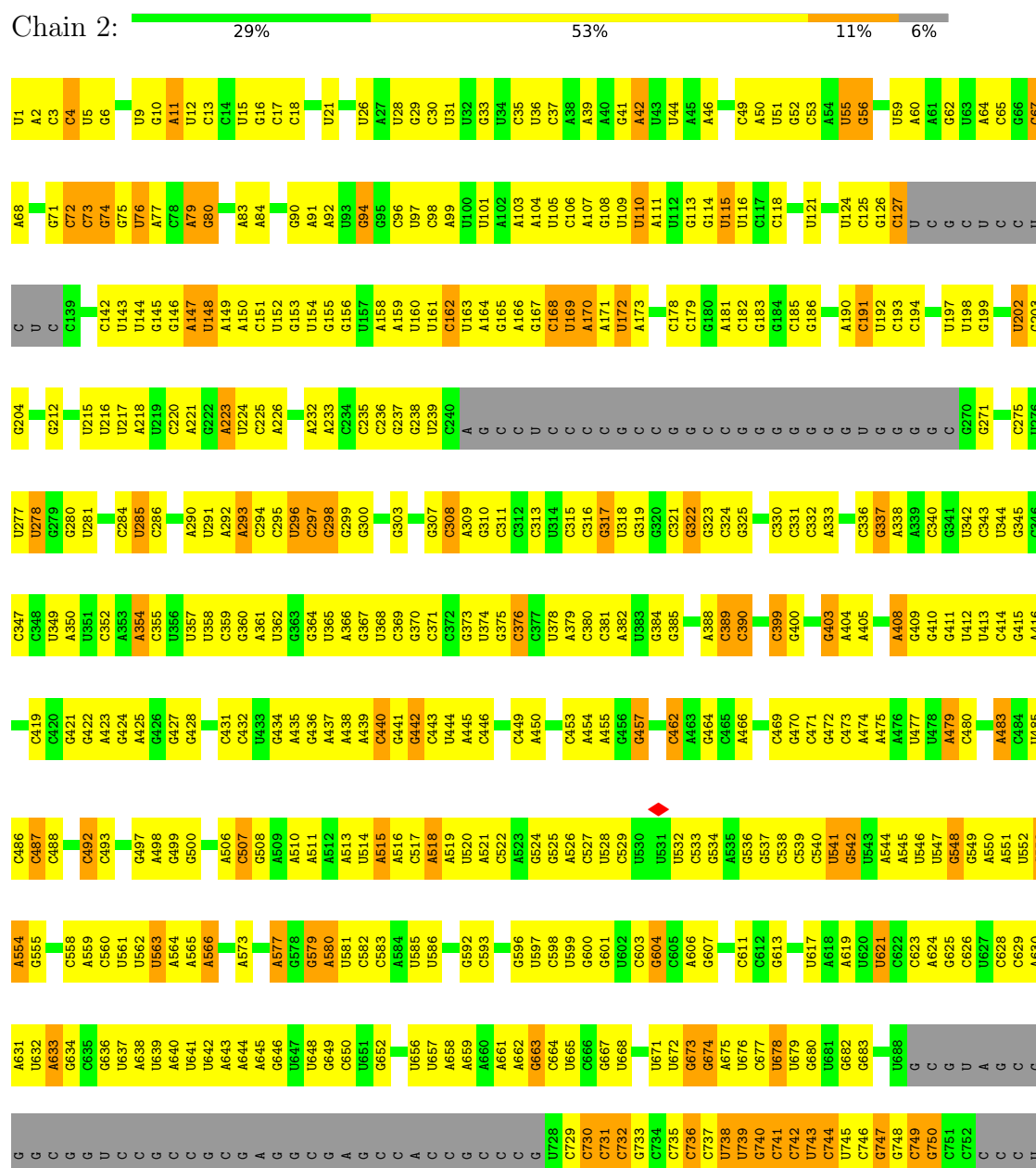
There are 8 discrepancies between the modelled and reference sequences:

| Chain | Residue | Modelled | Actual | Comment | Reference |
|-------|---------|----------|--------|----------------|--------------|
| z | 831 | A | - | expression tag | GB 485965777 |
| z | 832 | A | - | expression tag | GB 485965777 |
| z | 833 | U | - | expression tag | GB 485965777 |
| z | 834 | A | - | expression tag | GB 485965777 |
| z | 835 | U | - | expression tag | GB 485965777 |
| z | 836 | G | - | expression tag | GB 485965777 |
| z | 837 | G | - | expression tag | GB 485965777 |
| z | 838 | C | - | expression tag | GB 485965777 |

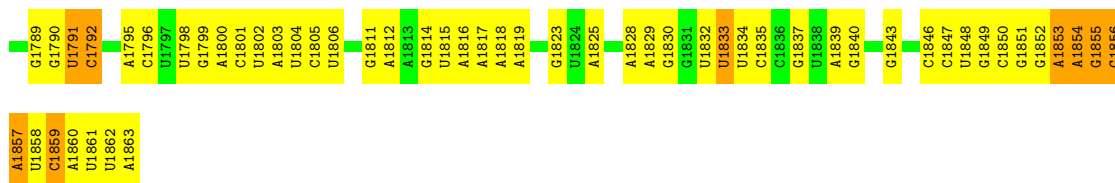
3 Residue-property plots

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.

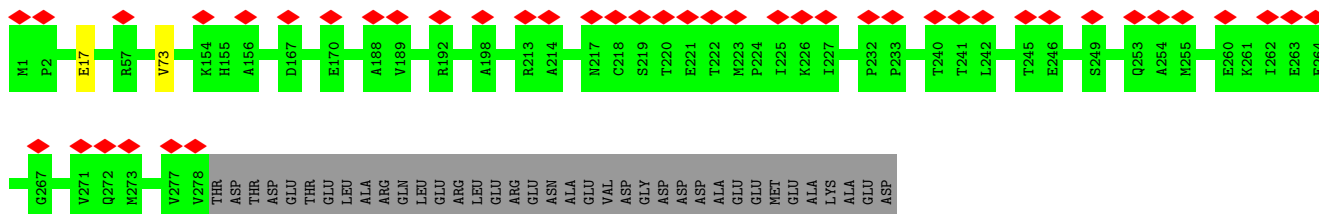
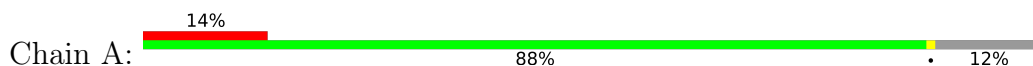
• Molecule 1: 18S ribosomal RNA



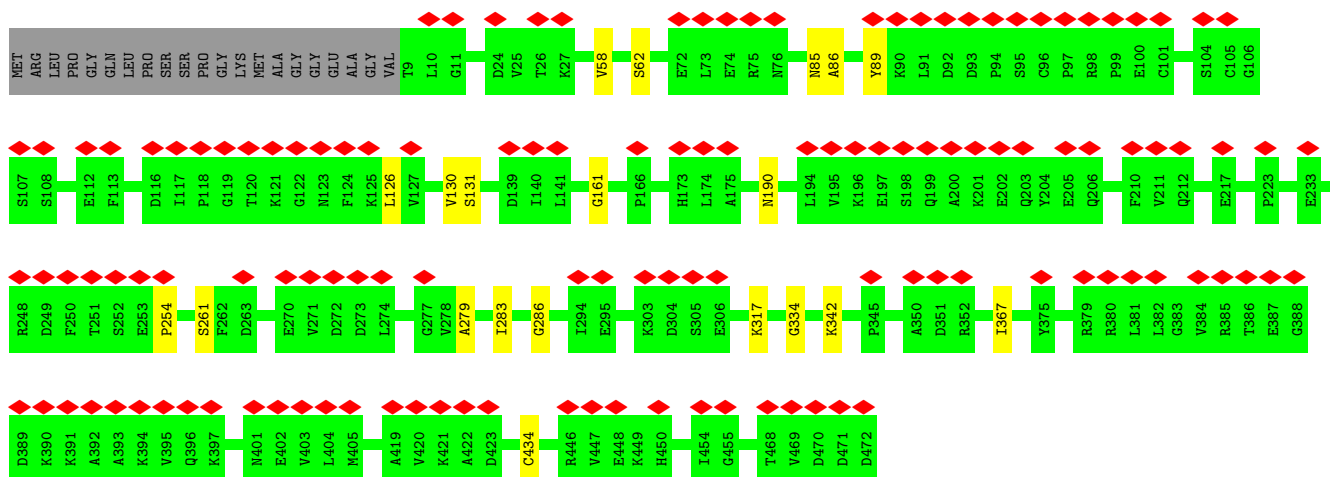
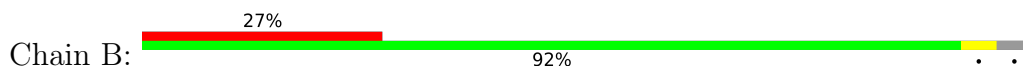
| | | | | | | | | | | | | | |
|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|------|---|
| A1714 | C1650 | C1586 | U1586 | U1386 | U1313 | U1244 | U1173 | G1101 | A1031 | G955 | U894 | G827 | C |
| U1715 | C1651 | C1587 | U1587 | C1387 | U1314 | U1245 | U1174 | C1102 | A1032 | U956 | U895 | G828 | G |
| U1716 | C1652 | C1588 | U1588 | U1388 | A1450 | A1246 | G1175 | C1103 | G957 | U896 | U896 | C829 | A |
| G1717 | C1653 | A1527 | C1527 | G1316 | G1316 | A1247 | C1176 | U1107 | U1034 | A958 | C896 | C830 | U |
| U1718 | U1654 | A1528 | A1528 | C1391 | G1317 | C1248 | A1177 | U1108 | C1035 | A959 | G897 | C831 | G |
| C1655 | U1655 | U1452 | U1452 | U1392 | U1453 | A1249 | A1178 | A1109 | U1041 | A960 | G898 | G832 | C |
| U1720 | A1656 | G1454 | G1454 | A1392 | G1320 | A1249 | A1179 | U1110 | U1042 | U961 | A899 | A833 | U |
| G1721 | U1594 | A1532 | A1532 | U1393 | G1321 | G1252 | G1190 | U1111 | U1043 | U962 | A900 | A833 | C |
| A1659 | G1595 | C1533 | C1533 | C1395 | U1322 | G1253 | G1183 | C1112 | C1043 | U963 | C901 | C835 | U |
| U1660 | A1596 | U1534 | U1534 | U1396 | G1323 | A1254 | G1183 | C1113 | U1044 | U964 | U902 | C836 | U |
| C1661 | U1597 | G1535 | U1535 | A1397 | U1458 | A1255 | G1183 | C1113 | A1045 | U965 | G903 | A837 | A |
| U1662 | U1598 | G1536 | U1536 | A1398 | U1459 | G1326 | U1179 | G1117 | U1041 | U966 | A904 | G838 | G |
| U1663 | U1599 | C1537 | U1537 | U1399 | U1460 | U1259 | A1179 | A1118 | C1049 | G967 | G905 | C839 | C |
| G1664 | U1600 | U1538 | U1538 | C1399 | A1461 | U1260 | G1190 | C1119 | U1050 | A968 | U840 | U840 | U |
| C1665 | G1601 | C1539 | U1539 | U1400 | A1328 | A1261 | U1189 | G1117 | A1051 | G907 | G906 | G841 | G |
| G1732 | C1665 | G1601 | A1540 | A1401 | G1331 | C1262 | A1190 | C1123 | U1052 | A976 | C908 | G842 | A |
| C1733 | U1602 | A1540 | A1540 | U1402 | U1332 | A1192 | A1191 | C1124 | C1053 | A977 | A909 | A843 | G |
| U1667 | U1603 | G1541 | U1541 | U1403 | C1332 | A1192 | A1191 | G1125 | A1054 | A978 | U910 | U844 | U |
| C1734 | C1604 | C1542 | C1542 | U1404 | C1333 | G1266 | G1193 | G1126 | G1055 | A979 | G911 | A845 | U |
| G1668 | U1605 | G1543 | U1543 | A1405 | U1339 | C1267 | G1194 | C1129 | U1056 | A980 | A912 | C846 | U |
| C1669 | G1606 | U1544 | U1544 | C1406 | G1344 | C1268 | A1196 | G1130 | A1058 | A986 | U914 | G848 | C |
| A1670 | U1607 | G1545 | U1545 | A1407 | U1344 | G1269 | U1197 | C1131 | C1059 | A987 | A915 | C849 | C |
| U1671 | G1608 | U1546 | U1546 | C1408 | G1345 | G1270 | U1198 | U1132 | C1060 | A988 | A916 | A850 | G |
| U1672 | A1609 | C1547 | U1547 | G1409 | U1346 | A1272 | G1199 | U1133 | G917 | G989 | G917 | U853 | C |
| A1673 | U1610 | U1474 | C1548 | A1410 | G1347 | C1273 | A1200 | C1134 | C1063 | A992 | A918 | U854 | G |
| G1674 | G1612 | U1550 | U1550 | C1411 | G1348 | A1274 | G1201 | C1135 | G1064 | A993 | G919 | G855 | G |
| G1675 | C1613 | A1551 | A1551 | U1412 | U1349 | C1275 | G1202 | G1136 | U1065 | A994 | G920 | G856 | C |
| U1676 | U1614 | C1552 | U1552 | C1414 | G1350 | G1276 | A1204 | G1137 | A1066 | A995 | A921 | U857 | C |
| C1678 | A1615 | C1553 | C1553 | C1415 | C1351 | A1279 | G1204 | U1138 | G1067 | G995 | A922 | C858 | C |
| U1680 | U1616 | C1554 | U1554 | G1352 | G1352 | A1280 | C1211 | A1139 | U1068 | C996 | C923 | U859 | C |
| U1617 | U1617 | U1555 | U1555 | U1353 | A1353 | G1281 | C1212 | A1140 | G1072 | U1000 | G924 | A860 | C |
| G1681 | C1681 | A1556 | A1556 | C1354 | U1354 | U1281 | A1212 | A1141 | A1073 | U0982 | G925 | G861 | C |
| C1682 | U1618 | C1557 | U1557 | G1420 | U1355 | A1282 | C1213 | C1142 | C1001 | G926 | G926 | U862 | C |
| C1683 | C1623 | G1558 | U1558 | G1421 | U1356 | U1283 | C1214 | C1143 | G1002 | C927 | G927 | G863 | C |
| U1685 | C1624 | C1559 | C1559 | A1422 | C1359 | U1284 | C1215 | C1143 | C1003 | G928 | G928 | G864 | C |
| C1686 | A1625 | U1560 | U1560 | U1490 | U1490 | U1285 | U1216 | A1144 | C1004 | A865 | G929 | U865 | C |
| U1687 | U1626 | C1561 | U1561 | G1491 | U1360 | G1286 | G1217 | A1145 | A1076 | A966 | G930 | U866 | C |
| C1688 | G1627 | U1562 | U1562 | A1492 | C1361 | A1287 | G1218 | U1005 | G932 | U867 | G931 | U867 | C |
| A | A1628 | C1563 | C1563 | C1426 | G1362 | C1288 | A1219 | A1146 | U1079 | A868 | G932 | A798 | C |
| C | A1629 | A1564 | U1564 | A1494 | A1494 | U1289 | G1220 | U1150 | A1080 | C799 | G933 | U800 | C |
| C1692 | C1630 | C1567 | U1567 | U1428 | A1365 | G1290 | G1220 | U1151 | C1081 | A1008 | G933 | U801 | C |
| C1693 | G1631 | C1568 | U1568 | C1429 | A1366 | A1291 | G1223 | U1152 | G1082 | U1009 | C937 | C870 | C |
| C1694 | A1632 | U1569 | U1569 | C1430 | U1367 | C1431 | A1224 | G1083 | G1010 | A871 | U942 | A812 | C |
| C1695 | G1633 | C1569 | C1569 | C1431 | U1368 | G1294 | G1225 | G1153 | U1084 | C938 | G943 | G813 | C |
| C1696 | C1634 | G1570 | U1570 | C1432 | A1375 | A1295 | C1226 | G1154 | C1090 | A814 | C944 | A814 | C |
| C1697 | A1635 | C1571 | U1571 | C1433 | C1376 | U1302 | U1234 | G1155 | U1091 | U1018 | G945 | G815 | C |
| C1698 | U1636 | G1572 | U1572 | C1434 | U1377 | U1303 | U1235 | G1165 | G1092 | C946 | C946 | U879 | C |
| C1699 | C1637 | U1573 | U1573 | A1434 | G1371 | U1304 | A1236 | G1166 | C1094 | C947 | U941 | A882 | C |
| C1700 | U1637 | U1573 | U1573 | U1506 | U1371 | U1304 | A1236 | U1021 | C1094 | U942 | U942 | C876 | C |
| G1701 | C1637 | A1574 | U1574 | U1507 | A1372 | C1380 | U1237 | G1160 | G1088 | C943 | G943 | G876 | C |
| U1702 | U1638 | C1574 | C1574 | C1436 | U1373 | G1381 | U1238 | G1161 | A1089 | C944 | G943 | G877 | C |
| C1703 | C1639 | A1575 | U1575 | U1437 | A1374 | U1300 | G1232 | G1162 | C1090 | A815 | C944 | A814 | C |
| U1704 | U1640 | C1576 | C1576 | U1438 | A1375 | C1301 | U1233 | G1163 | C1091 | U1018 | G945 | G815 | C |
| G1704 | C1641 | U1577 | U1577 | G1511 | C1376 | U1302 | U1234 | G1164 | G1092 | C946 | C946 | U879 | C |
| U1705 | A1642 | C1578 | U1578 | U1440 | G1377 | U1303 | U1235 | G1165 | G1093 | C947 | C947 | A882 | C |
| C1706 | G1643 | G1579 | U1579 | C1441 | U1377 | U1304 | A1236 | A1166 | C1094 | U942 | U942 | U883 | C |
| U1707 | U1644 | U1580 | U1580 | U1442 | C1380 | C1305 | A1237 | C1022 | G1095 | C949 | C949 | C820 | C |
| C1708 | A1645 | U1581 | U1581 | A1442 | G1381 | U1306 | U1238 | A1032 | A1023 | U950 | U950 | A821 | C |
| U1709 | U1646 | C1582 | U1582 | A1444 | A1382 | C1307 | U1239 | G1168 | U1097 | A951 | A951 | C822 | C |
| A1710 | A1646 | A1583 | G1583 | G1445 | A1382 | C1307 | U1239 | U1170 | G1098 | G952 | U950 | A822 | C |
| C1711 | G1647 | U1584 | U1584 | G1446 | G1383 | G1308 | U1240 | G1170 | C1099 | U888 | G952 | A823 | C |
| U1712 | U1648 | A1584 | A1584 | A1446 | A1384 | C1308 | G1241 | G1171 | C1099 | U889 | G1029 | A826 | C |
| C1713 | C1649 | C1585 | U1585 | G1447 | C1385 | U1310 | G1241 | C1172 | C1100 | A1030 | C954 | A826 | C |



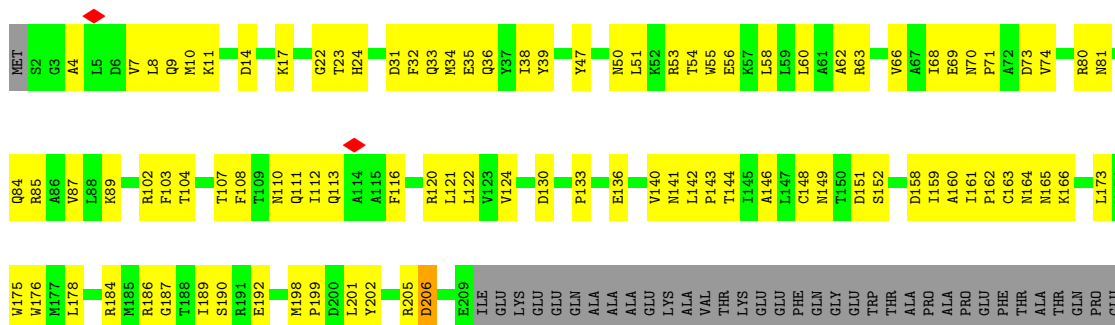
• Molecule 2: Eukaryotic translation initiation factor 2 subunit 1



• Molecule 3: Eukaryotic translation initiation factor 2 subunit 3

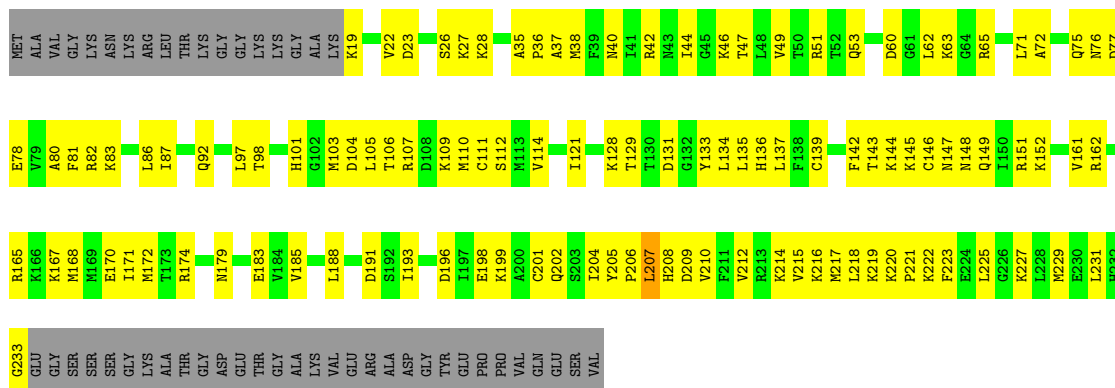


• Molecule 4: Small ribosomal subunit protein uS2



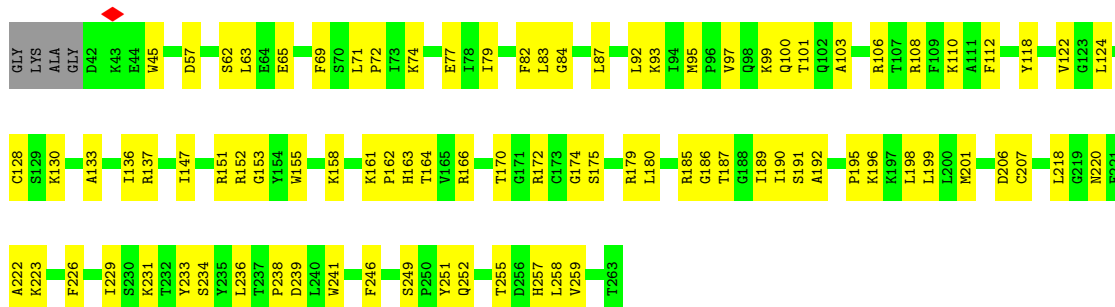
- Molecule 5: Small ribosomal subunit protein eS1

Chain D:  39% 42% 19%



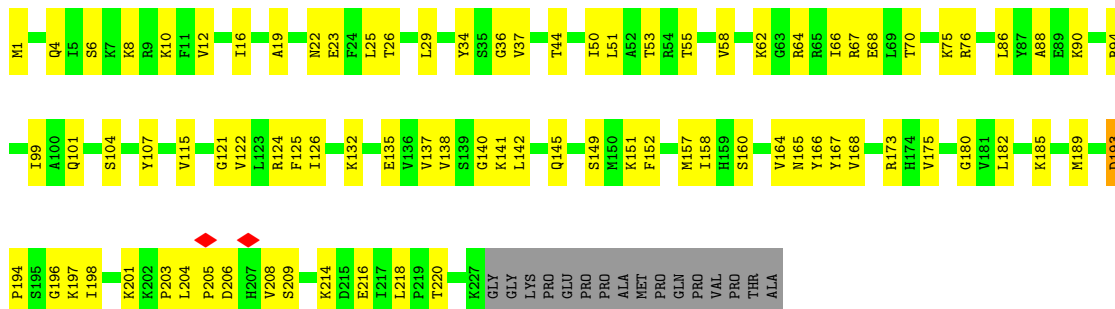
- Molecule 6: Small ribosomal subunit protein uS5

Chain E: 60% 38%



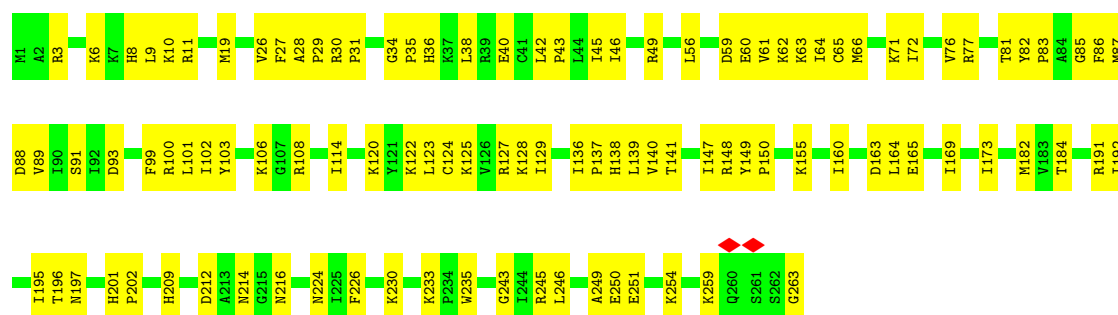
- Molecule 7: Small ribosomal subunit protein uS3

Chain F:  58% 35% 7%

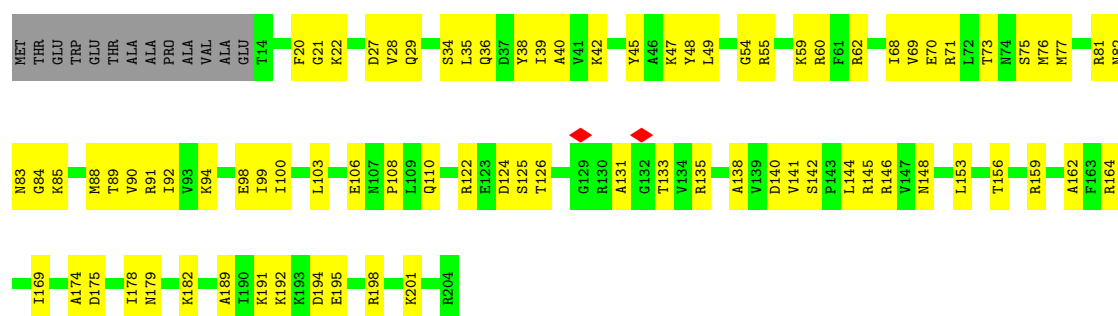


- Molecule 8: Small ribosomal subunit protein eS4

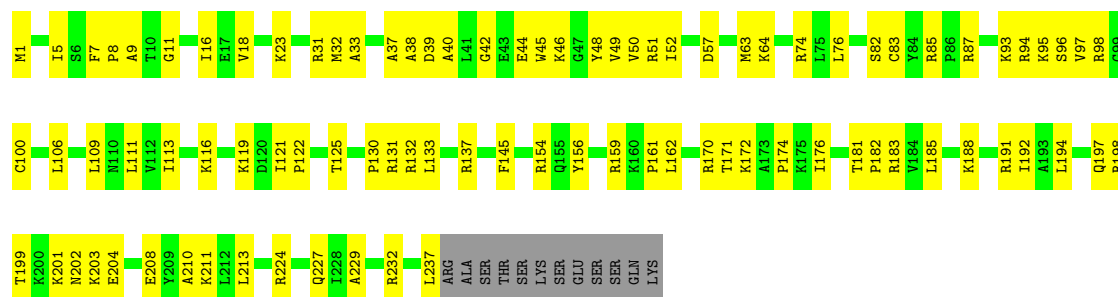
Chain G:  60% 40%



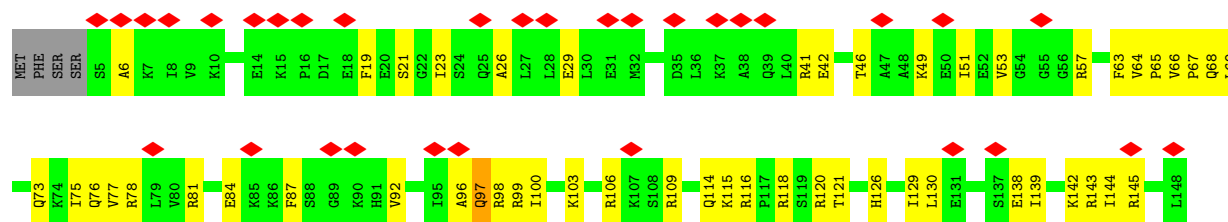
• Molecule 9: Small ribosomal subunit protein uS7

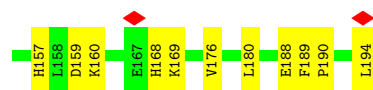


• Molecule 10: Small ribosomal subunit protein eS6



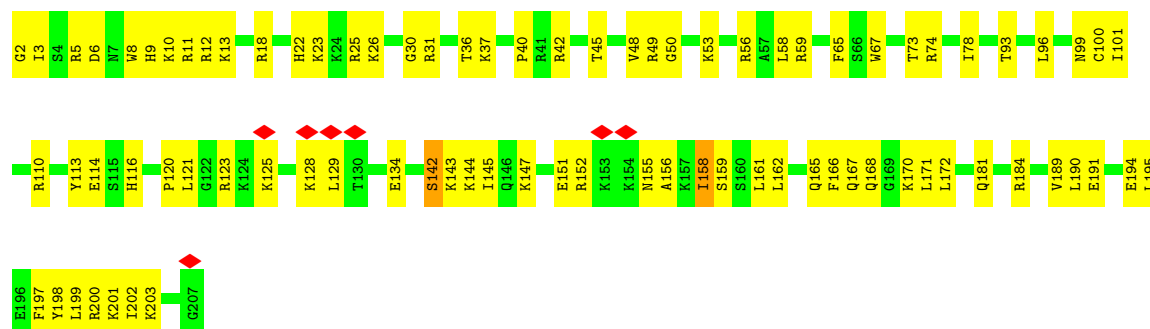
• Molecule 11: Small ribosomal subunit protein eS7





- Molecule 12: Small ribosomal subunit protein eS8

Chain K: 59% 40%



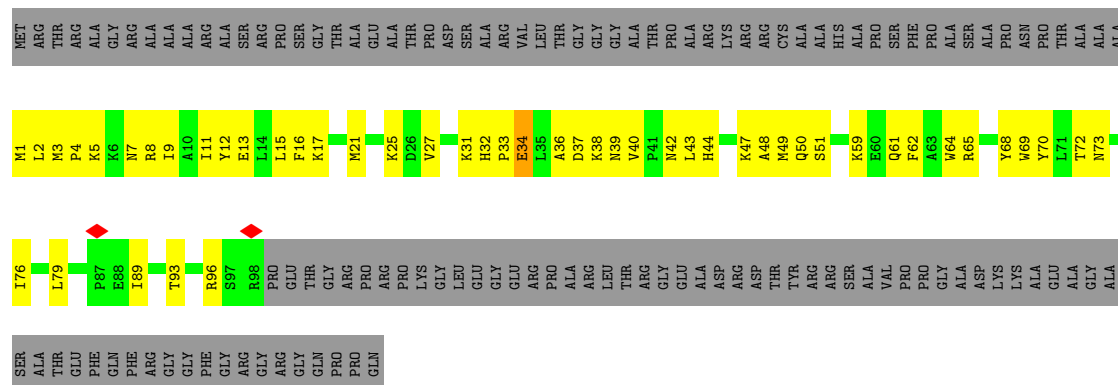
- Molecule 13: Small ribosomal subunit protein uS4

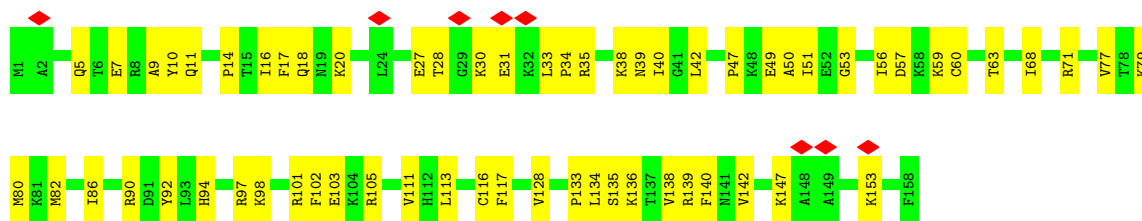
Chain L: 60% 33% 6%



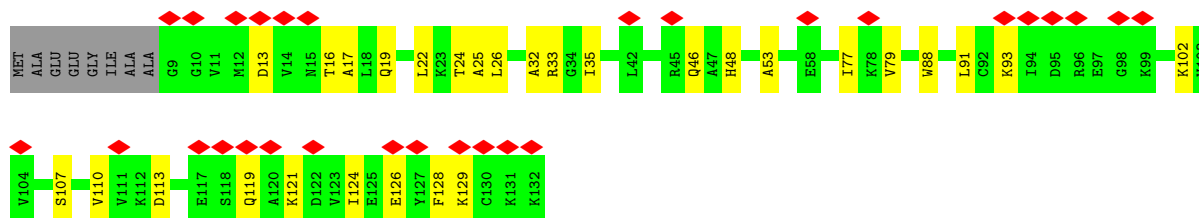
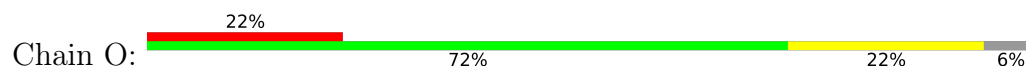
- Molecule 14: Small ribosomal subunit protein eS10

Chain M: 22% 21% 56%

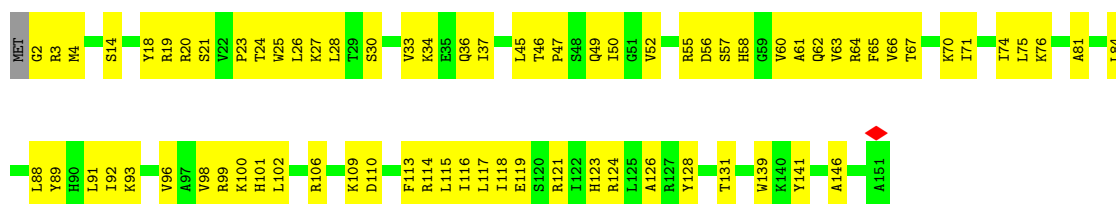




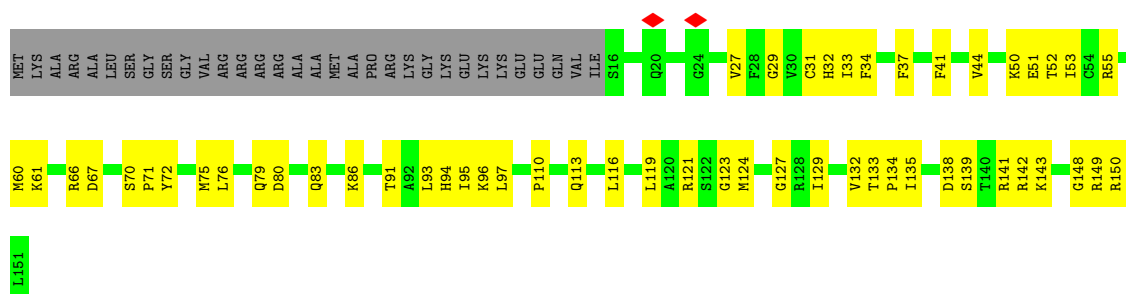
• Molecule 16: Small ribosomal subunit protein eS12



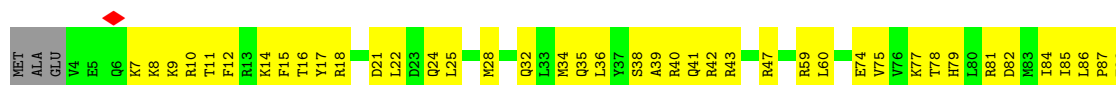
• Molecule 17: Small ribosomal subunit protein uS15



• Molecule 18: Ribosomal protein S14



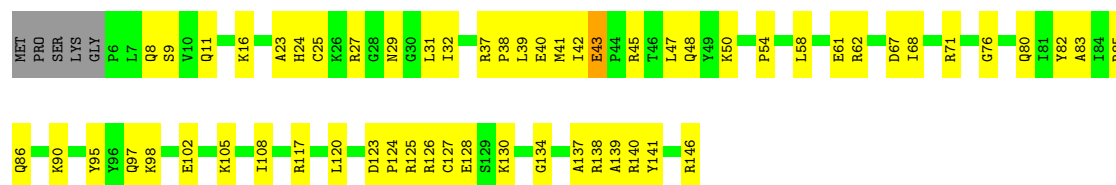
• Molecule 19: Small ribosomal subunit protein uS19





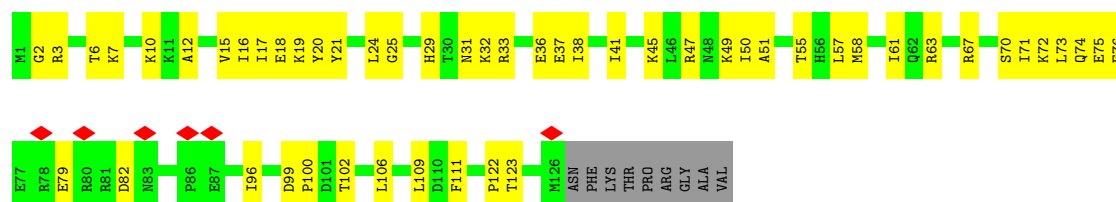
- Molecule 20: Small ribosomal subunit protein uS9

Chain S: 57% 39%



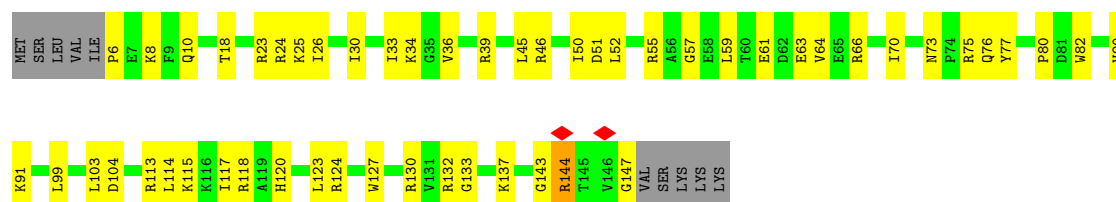
- Molecule 21: Small ribosomal subunit protein eS17

Chain T: 55% 39% 7%



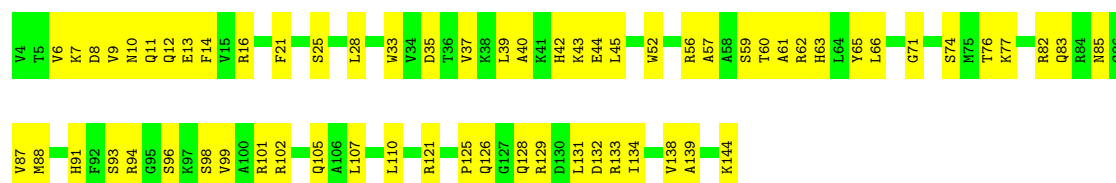
- Molecule 22: Small ribosomal subunit protein uS13

Chain U: 59% 34% 7%



- Molecule 23: Small ribosomal subunit protein eS19

Chain V: 55% 45%

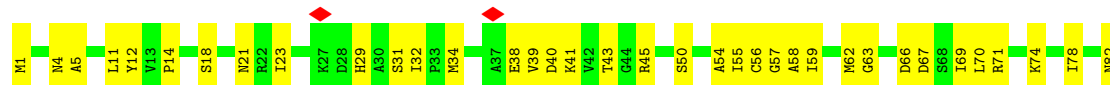


- Molecule 24: Small ribosomal subunit protein uS10

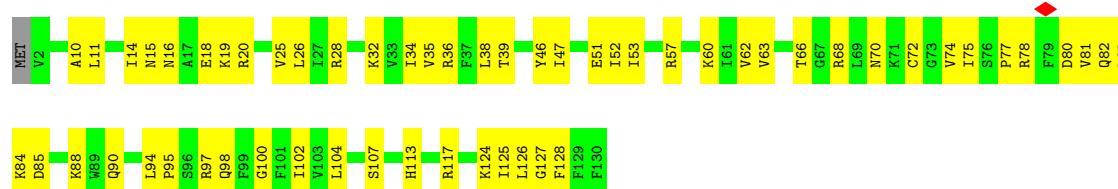
Chain W: 53% 34% 13%



- Molecule 25: Small ribosomal subunit protein eS21



- Molecule 26: Small ribosomal subunit protein uS8



- Molecule 27: Small ribosomal subunit protein uS12

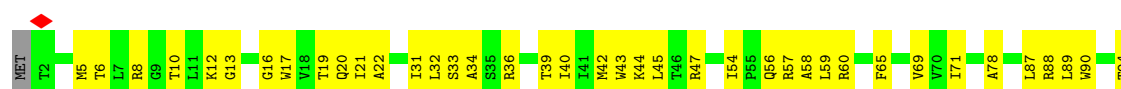


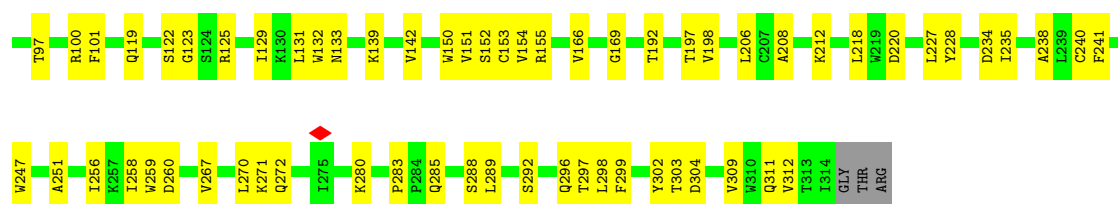
- Molecule 28: 40S ribosomal protein S24



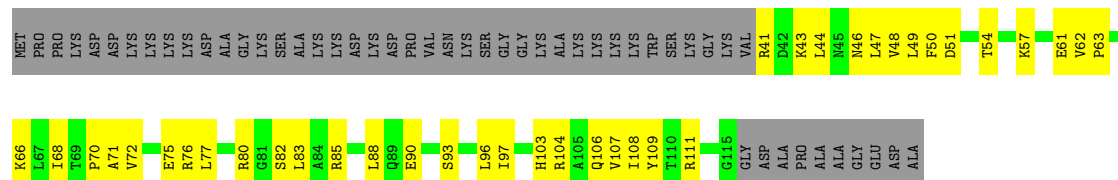
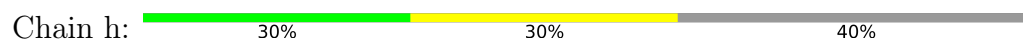
- Molecule 29: Small ribosomal subunit protein eS26







• Molecule 35: Small ribosomal subunit protein eS25



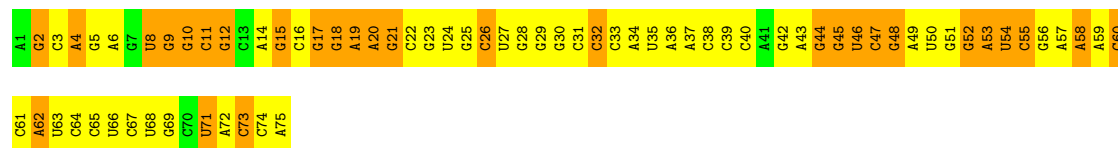
• Molecule 36: Small ribosomal subunit protein eS30



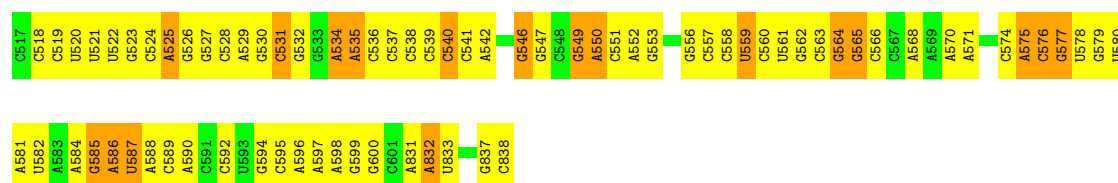
• Molecule 37: Small ribosomal subunit protein eS32



• Molecule 38: Initiator tRNA



• Molecule 39: Encephalomyocarditis viral IRES



4 Experimental information

| Property | Value | Source |
|--------------------------------------|---------------------------------|-----------|
| EM reconstruction method | SINGLE PARTICLE | Depositor |
| Imposed symmetry | POINT, C1 | Depositor |
| Number of particles used | 28439 | Depositor |
| Resolution determination method | FSC 0.143 CUT-OFF | Depositor |
| CTF correction method | NONE | Depositor |
| Microscope | FEI TALOS ARCTICA | Depositor |
| Voltage (kV) | 200 | Depositor |
| Electron dose ($e^-/\text{\AA}^2$) | 55 | Depositor |
| Minimum defocus (nm) | 1250 | Depositor |
| Maximum defocus (nm) | 2750 | Depositor |
| Magnification | 36000 | Depositor |
| Image detector | GATAN K2 SUMMIT (4k x 4k) | Depositor |
| Maximum map value | 0.554 | Depositor |
| Minimum map value | -0.216 | Depositor |
| Average map value | 0.001 | Depositor |
| Map value standard deviation | 0.023 | Depositor |
| Recommended contour level | 0.0618 | Depositor |
| Map size (\AA) | 467.99997, 467.99997, 467.99997 | wwPDB |
| Map dimensions | 400, 400, 400 | wwPDB |
| Map angles ($^\circ$) | 90.0, 90.0, 90.0 | wwPDB |
| Pixel spacing (\AA) | 1.17, 1.17, 1.17 | Depositor |

5 Model quality [i](#)

5.1 Standard geometry [i](#)

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 | 2 | 0.11 | 0/41588 | 0.28 | 0/64818 |
| 2 | A | 0.13 | 0/1377 | 0.36 | 0/1919 |
| 3 | B | 0.10 | 0/2280 | 0.31 | 0/3165 |
| 4 | C | 0.15 | 0/1679 | 0.42 | 0/2283 |
| 5 | D | 0.16 | 0/1769 | 0.48 | 0/2367 |
| 6 | E | 0.21 | 0/1757 | 0.50 | 0/2371 |
| 7 | F | 0.17 | 0/1792 | 0.48 | 2/2412 (0.1%) |
| 8 | G | 0.14 | 0/2125 | 0.37 | 0/2856 |
| 9 | H | 0.16 | 0/1531 | 0.44 | 0/2059 |
| 10 | I | 0.16 | 0/1946 | 0.43 | 0/2590 |
| 11 | J | 0.16 | 0/1553 | 0.44 | 0/2079 |
| 12 | K | 0.14 | 0/1709 | 0.53 | 2/2278 (0.1%) |
| 13 | L | 0.13 | 0/1522 | 0.38 | 0/2031 |
| 14 | M | 0.16 | 0/851 | 0.60 | 2/1147 (0.2%) |
| 15 | N | 0.14 | 0/1319 | 0.42 | 0/1761 |
| 16 | O | 0.11 | 0/968 | 0.34 | 0/1296 |
| 17 | P | 0.17 | 0/1232 | 0.41 | 0/1656 |
| 18 | Q | 0.14 | 0/1029 | 0.40 | 0/1380 |
| 19 | R | 0.16 | 0/1132 | 0.52 | 0/1510 |
| 20 | S | 0.14 | 0/1141 | 0.46 | 0/1528 |
| 21 | T | 0.13 | 0/1031 | 0.45 | 0/1383 |
| 22 | U | 0.15 | 0/1190 | 0.49 | 0/1592 |
| 23 | V | 0.16 | 0/1133 | 0.41 | 0/1517 |
| 24 | W | 0.13 | 0/832 | 0.38 | 0/1117 |
| 25 | X | 0.12 | 0/627 | 0.41 | 0/839 |
| 26 | Y | 0.18 | 0/1051 | 0.50 | 0/1406 |
| 27 | Z | 0.16 | 0/1124 | 0.42 | 0/1500 |
| 28 | a | 0.15 | 0/1039 | 0.54 | 2/1380 (0.1%) |
| 29 | b | 0.20 | 0/802 | 0.45 | 0/1076 |
| 30 | c | 0.19 | 0/673 | 0.63 | 4/902 (0.4%) |
| 31 | d | 0.16 | 0/509 | 0.45 | 0/680 |
| 32 | e | 0.18 | 0/455 | 0.46 | 0/603 |
| 33 | f | 0.13 | 0/593 | 0.36 | 0/786 |
| 34 | g | 0.12 | 0/2493 | 0.38 | 0/3394 |

| Mol | Chain | Bond lengths | | Bond angles | |
|-----|-------|--------------|---------|-------------|------------------|
| | | RMSZ | # Z >5 | RMSZ | # Z >5 |
| 35 | h | 0.15 | 0/604 | 0.36 | 0/810 |
| 36 | i | 0.19 | 0/478 | 0.62 | 2/628 (0.3%) |
| 37 | l | 0.17 | 0/241 | 0.49 | 0/305 |
| 38 | y | 0.13 | 0/1795 | 0.29 | 0/2798 |
| 39 | z | 0.11 | 0/2213 | 0.25 | 0/3442 |
| All | All | 0.13 | 0/89183 | 0.36 | 14/129664 (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 |
|-----|-------|---------------------|---------------------|
| 11 | J | 0 | 1 |
| 20 | S | 0 | 1 |
| All | All | 0 | 2 |

There are no bond length outliers.

All (14) bond angle outliers are listed below:

| Mol | Chain | Res | Type | Atoms | Z | Observed(°) | Ideal(°) |
|-----|-------|-----|------|--------|-------|-------------|----------|
| 12 | K | 142 | SER | CA-C-N | 11.15 | 141.77 | 121.70 |
| 12 | K | 142 | SER | C-N-CA | 11.15 | 141.77 | 121.70 |
| 14 | M | 1 | MET | CA-C-N | 8.95 | 137.81 | 121.70 |
| 14 | M | 1 | MET | C-N-CA | 8.95 | 137.81 | 121.70 |
| 28 | a | 102 | THR | CA-C-N | 8.60 | 137.18 | 121.70 |
| 28 | a | 102 | THR | C-N-CA | 8.60 | 137.18 | 121.70 |
| 7 | F | 197 | LYS | CA-C-N | 6.86 | 134.04 | 121.70 |
| 7 | F | 197 | LYS | C-N-CA | 6.86 | 134.04 | 121.70 |
| 36 | i | 118 | ASN | CA-C-N | 6.65 | 133.66 | 121.70 |
| 36 | i | 118 | ASN | C-N-CA | 6.65 | 133.66 | 121.70 |
| 30 | c | 81 | ARG | CA-C-N | 6.41 | 133.25 | 121.70 |
| 30 | c | 81 | ARG | C-N-CA | 6.41 | 133.25 | 121.70 |
| 30 | c | 20 | LYS | CA-C-N | 6.27 | 128.68 | 120.28 |
| 30 | c | 20 | LYS | C-N-CA | 6.27 | 128.68 | 120.28 |

There are no chirality outliers.

All (2) planarity outliers are listed below:

| Mol | Chain | Res | Type | Group |
|-----|-------|-----|------|---------|
| 11 | J | 66 | VAL | Peptide |

Continued on next page...

Continued from previous page...

| Mol | Chain | Res | Type | Group |
|-----|-------|-----|------|---------|
| 20 | S | 43 | GLU | 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 | 2 | 37195 | 0 | 18794 | 1205 | 0 |
| 2 | A | 1378 | 0 | 611 | 1 | 0 |
| 3 | B | 2281 | 0 | 1035 | 10 | 0 |
| 4 | C | 1642 | 0 | 1646 | 88 | 0 |
| 5 | D | 1741 | 0 | 1815 | 84 | 0 |
| 6 | E | 1721 | 0 | 1812 | 84 | 0 |
| 7 | F | 1764 | 0 | 1863 | 67 | 0 |
| 8 | G | 2083 | 0 | 2189 | 99 | 0 |
| 9 | H | 1509 | 0 | 1563 | 72 | 0 |
| 10 | I | 1923 | 0 | 2089 | 74 | 0 |
| 11 | J | 1530 | 0 | 1627 | 61 | 0 |
| 12 | K | 1680 | 0 | 1762 | 87 | 0 |
| 13 | L | 1498 | 0 | 1608 | 58 | 0 |
| 14 | M | 827 | 0 | 854 | 46 | 0 |
| 15 | N | 1296 | 0 | 1374 | 60 | 0 |
| 16 | O | 958 | 0 | 993 | 22 | 0 |
| 17 | P | 1208 | 0 | 1294 | 69 | 0 |
| 18 | Q | 1016 | 0 | 1039 | 53 | 0 |
| 19 | R | 1111 | 0 | 1168 | 68 | 0 |
| 20 | S | 1123 | 0 | 1193 | 51 | 0 |
| 21 | T | 1019 | 0 | 1075 | 48 | 0 |
| 22 | U | 1172 | 0 | 1226 | 55 | 0 |
| 23 | V | 1113 | 0 | 1149 | 53 | 0 |
| 24 | W | 822 | 0 | 887 | 40 | 0 |
| 25 | X | 620 | 0 | 622 | 34 | 0 |
| 26 | Y | 1034 | 0 | 1080 | 57 | 0 |
| 27 | Z | 1106 | 0 | 1179 | 56 | 0 |
| 28 | a | 1022 | 0 | 1085 | 50 | 0 |
| 29 | b | 789 | 0 | 839 | 46 | 0 |
| 30 | c | 659 | 0 | 683 | 24 | 0 |
| 31 | d | 507 | 0 | 536 | 26 | 0 |
| 32 | e | 445 | 0 | 442 | 38 | 0 |

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| Mol | Chain | Non-H | H(model) | H(added) | Clashes | Symm-Clashes |
|-----|-------|-------|----------|----------|---------|--------------|
| 33 | f | 581 | 0 | 599 | 35 | 0 |
| 34 | g | 2436 | 0 | 2393 | 77 | 0 |
| 35 | h | 598 | 0 | 656 | 37 | 0 |
| 36 | i | 473 | 0 | 524 | 20 | 0 |
| 37 | l | 240 | 0 | 289 | 18 | 0 |
| 38 | y | 1604 | 0 | 816 | 79 | 0 |
| 39 | z | 1981 | 0 | 1011 | 71 | 0 |
| All | All | 83705 | 0 | 63420 | 2712 | 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 19.

All (2712) 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) |
|------------------|------------------|--------------------------|-------------------|
| 1:2:1284:U:H3 | 1:2:1307:C:N4 | 1.45 | 1.14 |
| 1:2:432:C:N4 | 1:2:439:A:H62 | 1.51 | 1.08 |
| 1:2:957:G:C2 | 39:z:831:A:N1 | 2.24 | 1.05 |
| 1:2:432:C:H42 | 1:2:439:A:N6 | 1.53 | 1.04 |
| 1:2:149:A:N6 | 1:2:169:U:C2 | 2.26 | 1.03 |
| 1:2:124:U:H3 | 1:2:330:C:N4 | 1.58 | 1.00 |
| 1:2:957:G:N3 | 39:z:831:A:C6 | 2.29 | 1.00 |
| 1:2:1743:G:H1 | 1:2:1780:U:H3 | 1.06 | 1.00 |
| 1:2:800:U:H3 | 1:2:855:G:H1 | 1.08 | 0.99 |
| 32:e:36:LEU:HD12 | 32:e:38:MET:HE1 | 1.41 | 0.98 |
| 10:I:208:GLU:OE1 | 10:I:211:LYS:NZ | 1.97 | 0.96 |
| 1:2:957:G:C2 | 39:z:831:A:C6 | 2.54 | 0.95 |
| 1:2:1704:G:H1 | 1:2:1818:A:N6 | 1.64 | 0.94 |
| 1:2:192:U:H3 | 1:2:203:G:H1 | 1.05 | 0.94 |
| 1:2:1751:G:H1 | 1:2:1769:U:H3 | 1.13 | 0.94 |
| 1:2:1223:G:H1 | 1:2:1526:A:H61 | 1.10 | 0.91 |
| 1:2:1647:G:H1 | 1:2:1667:U:H3 | 0.93 | 0.90 |
| 1:2:1652:G:H1 | 1:2:1662:U:H3 | 1.18 | 0.89 |
| 1:2:116:U:H3 | 1:2:337:G:H1 | 0.89 | 0.89 |
| 1:2:1266:G:H1 | 1:2:1507:U:H3 | 0.90 | 0.89 |
| 1:2:1704:G:H1 | 1:2:1818:A:H61 | 0.88 | 0.87 |
| 32:e:16:GLN:HG2 | 32:e:27:ARG:HH21 | 1.40 | 0.87 |
| 34:g:87:LEU:HB2 | 34:g:101:PHE:HB2 | 1.55 | 0.87 |
| 1:2:1320:G:H1 | 1:2:1500:U:H3 | 1.16 | 0.86 |
| 8:G:182:MET:HE1 | 8:G:192:ILE:HG12 | 1.57 | 0.86 |
| 1:2:1029:G:H1 | 1:2:1076:A:HO2' | 1.19 | 0.86 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 1:2:483:A:H61 | 1:2:500:G:H1' | 1.41 | 0.85 |
| 18:Q:71:PRO:HB2 | 31:d:63:ARG:HH12 | 1.42 | 0.85 |
| 1:2:1815:U:H2' | 1:2:1816:A:H8 | 1.42 | 0.84 |
| 1:2:957:G:H21 | 39:z:831:A:H61 | 1.21 | 0.84 |
| 1:2:419:C:H5'' | 8:G:10:LYS:HE3 | 1.60 | 0.83 |
| 17:P:62:GLN:HG3 | 17:P:65:PHE:HD2 | 1.41 | 0.83 |
| 1:2:843:A:O2' | 8:G:106:LYS:NZ | 2.11 | 0.83 |
| 1:2:1223:G:H1 | 1:2:1526:A:N6 | 1.76 | 0.83 |
| 38:y:25:G:N1 | 38:y:43:A:C2 | 2.47 | 0.83 |
| 1:2:1648:U:H3 | 1:2:1666:G:H1 | 1.26 | 0.82 |
| 1:2:1717:G:H1 | 1:2:1806:U:H3 | 1.25 | 0.82 |
| 1:2:957:G:N2 | 39:z:831:A:H61 | 1.76 | 0.82 |
| 1:2:164:A:H3' | 1:2:165:G:H21 | 1.45 | 0.81 |
| 4:C:51:LEU:HD21 | 21:T:109:LEU:HD12 | 1.61 | 0.81 |
| 34:g:59:LEU:HB3 | 34:g:90:TRP:HZ3 | 1.46 | 0.81 |
| 1:2:953:A:H3' | 1:2:954:G:H21 | 1.44 | 0.81 |
| 1:2:668:U:H3 | 1:2:1023:A:H62 | 1.30 | 0.80 |
| 9:H:156:THR:HA | 9:H:159:ARG:HH21 | 1.45 | 0.80 |
| 1:2:1139:A:O3' | 1:2:1351:C:N4 | 2.15 | 0.80 |
| 1:2:313:C:N3 | 1:2:318:U:O4 | 2.16 | 0.79 |
| 1:2:683:G:O6 | 1:2:731:C:N4 | 2.16 | 0.79 |
| 25:X:14:PRO:HB2 | 25:X:23:ILE:HD11 | 1.64 | 0.79 |
| 1:2:1472:A:N7 | 21:T:3:ARG:NH2 | 2.28 | 0.79 |
| 27:Z:101:LEU:HB2 | 27:Z:124:LYS:HB2 | 1.64 | 0.78 |
| 14:M:7:ASN:ND2 | 14:M:39:ASN:O | 2.17 | 0.78 |
| 24:W:81:GLN:HB2 | 32:e:55:LEU:HD21 | 1.64 | 0.78 |
| 1:2:1320:G:N2 | 1:2:1500:U:O2 | 2.16 | 0.77 |
| 12:K:42:ARG:HB3 | 12:K:58:LEU:O | 1.84 | 0.77 |
| 8:G:45:ILE:HG22 | 8:G:61:VAL:HG11 | 1.65 | 0.77 |
| 6:E:151:ARG:HD3 | 6:E:166:ARG:HE | 1.49 | 0.77 |
| 20:S:85:ARG:HH12 | 20:S:117:ARG:HD3 | 1.48 | 0.77 |
| 1:2:940:A:H61 | 1:2:978:G:H1 | 1.31 | 0.77 |
| 1:2:1268:C:OP1 | 33:f:94:LYS:NZ | 2.17 | 0.77 |
| 1:2:1672:U:OP1 | 9:H:71:ARG:NH2 | 2.19 | 0.76 |
| 1:2:1414:C:N4 | 1:2:1421:G:OP2 | 2.18 | 0.76 |
| 30:c:1:MET:HE1 | 30:c:5:LYS:HB2 | 1.66 | 0.76 |
| 1:2:1408:C:H2' | 1:2:1409:G:C8 | 2.19 | 0.76 |
| 15:N:16:ILE:HD11 | 15:N:34:PRO:HG2 | 1.68 | 0.76 |
| 1:2:924:G:H1 | 1:2:1009:U:H3 | 1.34 | 0.76 |
| 28:a:102:THR:OG1 | 28:a:106:GLN:NE2 | 2.18 | 0.75 |
| 13:L:134:HIS:HE1 | 13:L:164:PRO:HD2 | 1.51 | 0.75 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 15:N:7:GLU:HG2 | 15:N:9:ALA:H | 1.50 | 0.75 |
| 24:W:63:ILE:HB | 24:W:80:PHE:HB2 | 1.67 | 0.75 |
| 27:Z:107:ARG:NH1 | 27:Z:111:ALA:O | 2.19 | 0.75 |
| 9:H:85:LYS:HB3 | 9:H:88:MET:HG2 | 1.69 | 0.75 |
| 12:K:36:THR:HG23 | 12:K:96:LEU:HB2 | 1.68 | 0.75 |
| 34:g:88:ARG:HB3 | 34:g:90:TRP:HE1 | 1.52 | 0.75 |
| 38:y:10:G:N7 | 38:y:45:G:C2 | 2.53 | 0.75 |
| 27:Z:1:MET:HE3 | 27:Z:3:LYS:HG3 | 1.67 | 0.75 |
| 1:2:551:A:O2' | 13:L:132:GLN:NE2 | 2.20 | 0.75 |
| 1:2:309:A:H2 | 1:2:322:G:H22 | 1.35 | 0.75 |
| 12:K:190:LEU:HD21 | 12:K:198:TYR:CD2 | 2.22 | 0.75 |
| 1:2:236:C:N4 | 1:2:237:G:O6 | 2.20 | 0.74 |
| 1:2:4:C:H1' | 13:L:18:ARG:HH22 | 1.53 | 0.74 |
| 4:C:38:ILE:HD11 | 4:C:47:TYR:HB3 | 1.67 | 0.74 |
| 27:Z:1:MET:SD | 27:Z:5:ARG:NH2 | 2.61 | 0.74 |
| 15:N:97:ARG:O | 15:N:97:ARG:NH1 | 2.20 | 0.74 |
| 1:2:826:A:OP2 | 1:2:842:G:N2 | 2.20 | 0.74 |
| 1:2:1350:G:N2 | 1:2:1353:A:OP2 | 2.21 | 0.74 |
| 14:M:27:VAL:HG12 | 14:M:43:LEU:HD21 | 1.70 | 0.73 |
| 11:J:77:VAL:HG23 | 11:J:78:ARG:HD2 | 1.69 | 0.73 |
| 1:2:434:G:H1 | 12:K:26:LYS:HE3 | 1.50 | 0.73 |
| 18:Q:142:ARG:O | 29:b:22:ARG:NH2 | 2.21 | 0.73 |
| 27:Z:95:GLU:OE2 | 27:Z:140:ARG:NH2 | 2.22 | 0.73 |
| 14:M:65:ARG:NH2 | 32:e:21:CYS:O | 2.22 | 0.73 |
| 1:2:1262:C:OP1 | 33:f:82:LYS:NZ | 2.16 | 0.73 |
| 12:K:190:LEU:HD21 | 12:K:198:TYR:HD2 | 1.53 | 0.73 |
| 15:N:103:GLU:OE2 | 27:Z:10:ALA:N | 2.22 | 0.73 |
| 4:C:122:LEU:HD13 | 4:C:124:VAL:HG23 | 1.71 | 0.73 |
| 1:2:1802:U:H2' | 1:2:1803:A:H8 | 1.53 | 0.73 |
| 4:C:122:LEU:HD21 | 4:C:144:THR:HG22 | 1.69 | 0.73 |
| 5:D:143:THR:HA | 5:D:207:LEU:HD23 | 1.70 | 0.73 |
| 10:I:52:ILE:HD11 | 10:I:109:LEU:HD12 | 1.71 | 0.73 |
| 1:2:409:G:O3' | 26:Y:88:LYS:NZ | 2.22 | 0.72 |
| 1:2:1651:G:H1 | 1:2:1663:U:H3 | 0.83 | 0.72 |
| 24:W:55:ARG:HA | 24:W:87:ARG:HG3 | 1.71 | 0.72 |
| 1:2:53:C:OP1 | 28:a:112:ASN:ND2 | 2.22 | 0.72 |
| 34:g:59:LEU:HB3 | 34:g:90:TRP:CZ3 | 2.24 | 0.72 |
| 1:2:1732:G:O6 | 1:2:1791:U:O2 | 2.08 | 0.72 |
| 22:U:23:ARG:NH2 | 35:h:46:ASN:O | 2.22 | 0.72 |
| 1:2:1072:G:O6 | 1:2:1073:A:N6 | 2.22 | 0.72 |
| 11:J:138:GLU:HG2 | 17:P:19:ARG:HH12 | 1.54 | 0.72 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 10:I:32:MET:HE3 | 10:I:33:ALA:H | 1.54 | 0.72 |
| 21:T:31:ASN:ND2 | 21:T:51:ALA:O | 2.22 | 0.72 |
| 4:C:81:ASN:HA | 4:C:84:GLN:HE22 | 1.55 | 0.71 |
| 38:y:53:A:N6 | 38:y:61:C:O2 | 2.20 | 0.71 |
| 1:2:381:C:H2' | 1:2:382:A:H8 | 1.55 | 0.71 |
| 1:2:1198:U:H2' | 1:2:1199:G:C8 | 2.25 | 0.71 |
| 13:L:172:ARG:HD2 | 13:L:176:LYS:HZ3 | 1.55 | 0.71 |
| 1:2:1460:C:OP2 | 21:T:63:ARG:NH2 | 2.22 | 0.71 |
| 6:E:161:LYS:O | 6:E:185:ARG:NH2 | 2.23 | 0.71 |
| 19:R:74:GLU:HG2 | 19:R:75:VAL:H | 1.56 | 0.71 |
| 38:y:24:U:H2' | 38:y:25:G:H8 | 1.56 | 0.71 |
| 1:2:668:U:O4 | 1:2:1023:A:N7 | 2.23 | 0.71 |
| 1:2:1226:C:HO2' | 1:2:1660:G:H1 | 1.36 | 0.71 |
| 1:2:1418:G:H2' | 1:2:1419:C:H2' | 1.73 | 0.71 |
| 18:Q:142:ARG:HH21 | 29:b:22:ARG:HB3 | 1.56 | 0.71 |
| 22:U:46:ARG:HD2 | 23:V:35:ASP:HB3 | 1.71 | 0.71 |
| 1:2:1034:U:H1' | 1:2:1176:C:H42 | 1.56 | 0.71 |
| 6:E:110:LYS:HD3 | 6:E:128:CYS:HB2 | 1.72 | 0.71 |
| 1:2:739:U:O2' | 1:2:794:G:N2 | 2.24 | 0.70 |
| 1:2:1575:A:H5' | 24:W:56:MET:HE3 | 1.73 | 0.70 |
| 19:R:18:ARG:H | 22:U:91:LYS:HA | 1.56 | 0.70 |
| 1:2:671:U:H4' | 27:Z:9:THR:HG22 | 1.73 | 0.70 |
| 1:2:876:G:N2 | 1:2:876:G:OP2 | 2.24 | 0.70 |
| 7:F:214:LYS:NZ | 21:T:19:LYS:O | 2.24 | 0.70 |
| 29:b:87:ARG:NH1 | 29:b:91:ALA:O | 2.20 | 0.70 |
| 1:2:957:G:N2 | 39:z:831:A:N6 | 2.38 | 0.70 |
| 34:g:206:LEU:HD11 | 34:g:218:LEU:HB3 | 1.72 | 0.70 |
| 1:2:1281:G:H22 | 33:f:104:LYS:HG2 | 1.57 | 0.70 |
| 7:F:204:LEU:HB2 | 7:F:208:VAL:HA | 1.74 | 0.70 |
| 6:E:108:ARG:NH1 | 6:E:128:CYS:SG | 2.64 | 0.70 |
| 19:R:15:PHE:HB3 | 22:U:91:LYS:HB3 | 1.74 | 0.70 |
| 10:I:121:ILE:HD12 | 10:I:122:PRO:HD2 | 1.74 | 0.69 |
| 12:K:113:TYR:HB3 | 12:K:121:LEU:HD21 | 1.74 | 0.69 |
| 14:M:61:GLN:HB2 | 14:M:68:TYR:HB2 | 1.73 | 0.69 |
| 29:b:38:LYS:HB3 | 29:b:71:LEU:HD11 | 1.74 | 0.69 |
| 1:2:1094:C:H2' | 1:2:1095:G:C8 | 2.26 | 0.69 |
| 12:K:194:GLU:OE1 | 15:N:18:GLN:NE2 | 2.25 | 0.69 |
| 18:Q:79:GLN:O | 18:Q:83:GLN:NE2 | 2.25 | 0.69 |
| 1:2:148:U:H2' | 1:2:149:A:H8 | 1.55 | 0.69 |
| 26:Y:53:ILE:HB | 26:Y:60:LYS:HB3 | 1.74 | 0.69 |
| 1:2:915:A:H62 | 1:2:1016:A:H2' | 1.58 | 0.69 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 12:K:142:SER:H | 12:K:143:LYS:HB2 | 1.58 | 0.69 |
| 34:g:6:THR:HG23 | 34:g:8:ARG:HH12 | 1.55 | 0.69 |
| 9:H:59:LYS:HB2 | 9:H:62:ARG:HB2 | 1.74 | 0.69 |
| 38:y:23:G:H2' | 38:y:24:U:O4' | 1.93 | 0.69 |
| 1:2:2:A:H5' | 1:2:408:A:H5' | 1.75 | 0.69 |
| 1:2:487:C:H4' | 8:G:8:HIS:CD2 | 2.28 | 0.69 |
| 1:2:738:U:O2' | 1:2:740:G:OP2 | 2.11 | 0.69 |
| 1:2:1643:G:N2 | 1:2:1670:A:OP2 | 2.25 | 0.69 |
| 1:2:1645:A:H5'' | 20:S:139:ALA:HB2 | 1.75 | 0.69 |
| 1:2:1709:U:H2' | 1:2:1710:A:C8 | 2.28 | 0.69 |
| 15:N:86:ILE:HB | 15:N:113:LEU:HD23 | 1.73 | 0.69 |
| 20:S:146:ARG:NH2 | 38:y:32:C:OP2 | 2.24 | 0.69 |
| 39:z:551:C:H2' | 39:z:552:A:H8 | 1.58 | 0.69 |
| 1:2:947:C:H2' | 1:2:948:G:H8 | 1.57 | 0.69 |
| 1:2:1584:A:H5'' | 23:V:82:ARG:HD2 | 1.75 | 0.69 |
| 1:2:1138:G:N2 | 1:2:1141:A:OP2 | 2.22 | 0.69 |
| 33:f:121:CYS:HB3 | 33:f:132:MET:HB2 | 1.75 | 0.69 |
| 29:b:83:VAL:HG23 | 29:b:84:VAL:HG23 | 1.75 | 0.68 |
| 1:2:1279:C:H42 | 16:O:102:LYS:HA | 1.58 | 0.68 |
| 1:2:1491:G:N2 | 32:e:42:CYS:SG | 2.65 | 0.68 |
| 5:D:83:LYS:HD3 | 5:D:106:THR:HG23 | 1.75 | 0.68 |
| 1:2:814:A:OP1 | 13:L:80:ARG:NH2 | 2.26 | 0.68 |
| 26:Y:94:LEU:HD11 | 26:Y:100:GLY:HA3 | 1.75 | 0.68 |
| 30:c:18:LYS:HD2 | 30:c:22:LYS:HD2 | 1.75 | 0.68 |
| 38:y:48:G:N2 | 39:z:550:A:O2' | 2.26 | 0.68 |
| 1:2:275:C:N3 | 1:2:887:G:O2' | 2.26 | 0.68 |
| 1:2:1060:C:OP1 | 18:Q:149:ARG:NH2 | 2.27 | 0.68 |
| 38:y:25:G:O6 | 38:y:43:A:N1 | 2.26 | 0.68 |
| 1:2:957:G:N3 | 39:z:831:A:N6 | 2.42 | 0.68 |
| 4:C:148:CYS:SG | 4:C:152:SER:OG | 2.49 | 0.68 |
| 7:F:137:VAL:HB | 7:F:185:LYS:HB3 | 1.76 | 0.68 |
| 15:N:31:GLU:HG2 | 15:N:33:LEU:H | 1.59 | 0.68 |
| 34:g:13:GLY:HA3 | 34:g:43:TRP:HH2 | 1.58 | 0.68 |
| 9:H:22:LYS:NZ | 9:H:98:GLU:OE1 | 2.27 | 0.68 |
| 1:2:106:C:H2' | 1:2:107:A:H8 | 1.59 | 0.68 |
| 12:K:199:LEU:HD13 | 12:K:202:ILE:HD11 | 1.75 | 0.68 |
| 1:2:1370:C:OP2 | 21:T:7:LYS:NZ | 2.27 | 0.68 |
| 8:G:192:ILE:HG13 | 8:G:243:GLY:HA3 | 1.74 | 0.68 |
| 38:y:29:G:H2' | 38:y:30:G:C8 | 2.28 | 0.68 |
| 39:z:518:C:H2' | 39:z:519:C:C6 | 2.29 | 0.68 |
| 1:2:800:U:O2 | 1:2:855:G:N2 | 2.24 | 0.67 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 1:2:865:A:C6 | 11:J:114:GLN:HB3 | 2.29 | 0.67 |
| 1:2:842:G:C4 | 8:G:19:MET:HE1 | 2.29 | 0.67 |
| 1:2:1450:A:H5'' | 21:T:3:ARG:HH12 | 1.59 | 0.67 |
| 6:E:151:ARG:NH2 | 6:E:233:TYR:O | 2.27 | 0.67 |
| 11:J:23:ILE:HG13 | 11:J:87:PHE:HE2 | 1.56 | 0.67 |
| 1:2:511:A:OP1 | 13:L:45:ARG:NH1 | 2.27 | 0.67 |
| 1:2:1536:G:N3 | 23:V:12:GLN:NE2 | 2.40 | 0.67 |
| 12:K:151:GLU:O | 12:K:155:ASN:ND2 | 2.28 | 0.67 |
| 20:S:23:ALA:HB1 | 20:S:68:ILE:HD11 | 1.76 | 0.67 |
| 39:z:551:C:H2' | 39:z:552:A:C8 | 2.29 | 0.67 |
| 1:2:923:C:O2 | 30:c:51:GLN:NE2 | 2.28 | 0.67 |
| 11:J:75:ILE:HG23 | 11:J:76:GLN:H | 1.59 | 0.67 |
| 1:2:492:C:N3 | 28:a:84:LYS:NZ | 2.42 | 0.67 |
| 1:2:995:G:H3' | 29:b:17:HIS:CE1 | 2.29 | 0.67 |
| 1:2:1490:U:H4' | 1:2:1491:G:H5'' | 1.75 | 0.67 |
| 1:2:1857:A:OP2 | 29:b:4:LYS:NZ | 2.26 | 0.67 |
| 26:Y:11:LEU:HD21 | 26:Y:74:VAL:HB | 1.77 | 0.67 |
| 19:R:41:GLN:HG3 | 19:R:84:ILE:HD12 | 1.77 | 0.67 |
| 1:2:1117:G:O2' | 5:D:204:ILE:O | 2.12 | 0.67 |
| 13:L:84:ILE:HD11 | 13:L:148:ILE:HG22 | 1.77 | 0.67 |
| 15:N:11:GLN:HB2 | 15:N:56:ILE:HG21 | 1.76 | 0.67 |
| 1:2:526:A:H62 | 1:2:537:G:H21 | 1.39 | 0.67 |
| 8:G:72:ILE:HD11 | 8:G:88:ASP:HB3 | 1.77 | 0.67 |
| 1:2:73:C:O2 | 1:2:74:G:H1' | 1.95 | 0.67 |
| 1:2:1474:U:H2' | 1:2:1475:G:C8 | 2.30 | 0.67 |
| 4:C:10:MET:HG2 | 21:T:111:PHE:HE2 | 1.60 | 0.67 |
| 8:G:11:ARG:HA | 8:G:28:ALA:HB2 | 1.76 | 0.67 |
| 20:S:130:LYS:HA | 20:S:137:ALA:HA | 1.76 | 0.67 |
| 1:2:915:A:OP1 | 26:Y:57:ARG:NH1 | 2.28 | 0.66 |
| 11:J:65:PRO:O | 11:J:98:ARG:NH2 | 2.28 | 0.66 |
| 34:g:155:ARG:HG2 | 34:g:198:VAL:HG23 | 1.77 | 0.66 |
| 1:2:1155:G:OP2 | 27:Z:5:ARG:NH1 | 2.28 | 0.66 |
| 35:h:62:VAL:HG11 | 35:h:96:LEU:HB3 | 1.77 | 0.66 |
| 38:y:11:C:N4 | 38:y:24:U:H3 | 1.93 | 0.66 |
| 1:2:411:G:H5' | 15:N:98:LYS:HG3 | 1.77 | 0.66 |
| 1:2:835:C:H1' | 1:2:837:G:H1' | 1.78 | 0.66 |
| 6:E:93:LYS:HG3 | 6:E:218:LEU:HD21 | 1.77 | 0.66 |
| 38:y:3:C:H2' | 38:y:4:A:C8 | 2.30 | 0.66 |
| 1:2:668:U:H3 | 1:2:1023:A:N6 | 1.93 | 0.66 |
| 1:2:957:G:C2 | 39:z:831:A:N6 | 2.62 | 0.66 |
| 9:H:55:ARG:NH2 | 20:S:123:ASP:OD1 | 2.29 | 0.66 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 1:2:1451:A:H61 | 1:2:1467:C:H42 | 1.42 | 0.66 |
| 1:2:1555:U:HO2' | 1:2:1578:C:HO2' | 1.42 | 0.66 |
| 1:2:1773:G:H2' | 1:2:1774:G:C8 | 2.31 | 0.66 |
| 1:2:124:U:H3 | 1:2:330:C:H42 | 0.76 | 0.66 |
| 20:S:31:LEU:HD22 | 20:S:67:ASP:HB3 | 1.78 | 0.66 |
| 1:2:1091:U:O3' | 26:Y:20:ARG:NH1 | 2.29 | 0.66 |
| 12:K:37:LYS:NZ | 12:K:93:THR:O | 2.29 | 0.66 |
| 13:L:38:ARG:HA | 36:i:105:ARG:HB3 | 1.78 | 0.66 |
| 1:2:1245:C:H5'' | 1:2:1246:A:H2' | 1.78 | 0.66 |
| 36:i:118:ASN:H | 36:i:119:VAL:HA | 1.61 | 0.66 |
| 39:z:595:C:H2' | 39:z:596:A:H8 | 1.61 | 0.66 |
| 1:2:77:A:H4' | 10:I:176:ILE:HD12 | 1.76 | 0.66 |
| 1:2:166:A:H2' | 1:2:167:G:H8 | 1.61 | 0.66 |
| 1:2:841:G:H2' | 1:2:842:G:C8 | 2.30 | 0.66 |
| 10:I:51:ARG:NH1 | 10:I:52:ILE:O | 2.29 | 0.66 |
| 11:J:139:ILE:O | 17:P:19:ARG:NH1 | 2.27 | 0.66 |
| 14:M:3:MET:O | 14:M:8:ARG:NH1 | 2.28 | 0.66 |
| 1:2:67:C:N4 | 1:2:151:C:O2' | 2.29 | 0.65 |
| 1:2:1700:C:O2' | 37:l:2:ARG:NH2 | 2.29 | 0.65 |
| 1:2:1717:G:N2 | 1:2:1806:U:O2 | 2.27 | 0.65 |
| 12:K:73:THR:O | 12:K:74:ARG:NH1 | 2.26 | 0.65 |
| 15:N:60:CYS:SG | 15:N:63:THR:OG1 | 2.54 | 0.65 |
| 1:2:492:C:O3' | 8:G:62:LYS:NZ | 2.29 | 0.65 |
| 1:2:526:A:H62 | 1:2:537:G:N2 | 1.93 | 0.65 |
| 26:Y:47:ILE:HD11 | 26:Y:63:VAL:HG11 | 1.78 | 0.65 |
| 1:2:940:A:N6 | 1:2:978:G:H1 | 1.94 | 0.65 |
| 1:2:1309:A:OP2 | 16:O:33:ARG:NH2 | 2.29 | 0.65 |
| 5:D:46:LYS:HE3 | 18:Q:27:VAL:HB | 1.78 | 0.65 |
| 26:Y:94:LEU:HD12 | 26:Y:95:PRO:HD2 | 1.76 | 0.65 |
| 1:2:59:U:H5' | 1:2:493:C:H41 | 1.61 | 0.65 |
| 8:G:230:LYS:HB3 | 8:G:233:LYS:HB2 | 1.77 | 0.65 |
| 22:U:23:ARG:O | 22:U:55:ARG:NH1 | 2.29 | 0.65 |
| 1:2:940:A:H5'' | 18:Q:134:PRO:HB2 | 1.79 | 0.65 |
| 14:M:64:TRP:O | 32:e:22:ARG:NH2 | 2.30 | 0.65 |
| 20:S:138:ARG:NH1 | 20:S:138:ARG:O | 2.28 | 0.65 |
| 14:M:49:MET:HE3 | 14:M:69:TRP:CD1 | 2.31 | 0.65 |
| 1:2:667:G:N1 | 1:2:1023:A:OP2 | 2.25 | 0.65 |
| 1:2:957:G:H21 | 39:z:831:A:N6 | 1.94 | 0.65 |
| 8:G:64:ILE:HD11 | 28:a:18:LEU:HG | 1.79 | 0.65 |
| 19:R:15:PHE:HB3 | 22:U:91:LYS:HD3 | 1.77 | 0.65 |
| 22:U:99:LEU:O | 22:U:103:LEU:N | 2.25 | 0.65 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 1:2:1709:U:H2' | 1:2:1710:A:H8 | 1.61 | 0.65 |
| 4:C:85:ARG:NH1 | 4:C:201:LEU:O | 2.30 | 0.65 |
| 29:b:37:LYS:NZ | 29:b:71:LEU:O | 2.29 | 0.65 |
| 1:2:487:C:H4' | 8:G:8:HIS:HD2 | 1.60 | 0.65 |
| 1:2:1457:G:H3' | 1:2:1459:U:H3 | 1.62 | 0.65 |
| 4:C:56:GLU:OE2 | 25:X:82:ASN:ND2 | 2.29 | 0.65 |
| 1:2:745:U:H2' | 1:2:746:C:C6 | 2.31 | 0.65 |
| 1:2:1099:C:H2' | 1:2:1100:G:C8 | 2.31 | 0.65 |
| 1:2:1211:C:H42 | 1:2:1216:A:H61 | 1.44 | 0.65 |
| 1:2:365:U:H2' | 1:2:366:A:C8 | 2.32 | 0.64 |
| 7:F:104:SER:HA | 7:F:107:TYR:CE1 | 2.32 | 0.64 |
| 8:G:31:PRO:HG3 | 8:G:43:PRO:HG3 | 1.77 | 0.64 |
| 10:I:8:PRO:HD2 | 10:I:113:ILE:HG22 | 1.79 | 0.64 |
| 15:N:101:ARG:HB2 | 27:Z:10:ALA:HB2 | 1.79 | 0.64 |
| 17:P:119:GLU:OE2 | 17:P:123:HIS:NE2 | 2.29 | 0.64 |
| 1:2:941:U:H2' | 1:2:942:U:C6 | 2.32 | 0.64 |
| 1:2:1152:U:O4 | 6:E:179:ARG:NH1 | 2.30 | 0.64 |
| 20:S:32:ILE:HB | 20:S:39:LEU:HD21 | 1.79 | 0.64 |
| 1:2:1198:U:H2' | 1:2:1199:G:H8 | 1.61 | 0.64 |
| 18:Q:34:PHE:HB3 | 18:Q:41:PHE:HB2 | 1.78 | 0.64 |
| 1:2:166:A:H2' | 1:2:167:G:C8 | 2.33 | 0.64 |
| 1:2:815:G:OP2 | 13:L:80:ARG:NH1 | 2.30 | 0.64 |
| 1:2:847:C:H5'' | 1:2:848:G:H5' | 1.79 | 0.64 |
| 4:C:141:ASN:ND2 | 25:X:31:SER:O | 2.30 | 0.64 |
| 1:2:903:G:H2' | 1:2:904:A:C8 | 2.32 | 0.64 |
| 14:M:5:LYS:HA | 14:M:8:ARG:HD2 | 1.79 | 0.64 |
| 15:N:80:MET:SD | 15:N:86:ILE:HD12 | 2.38 | 0.64 |
| 39:z:575:A:H8 | 39:z:576:C:H4' | 1.63 | 0.64 |
| 1:2:922:A:H61 | 1:2:1011:U:H5 | 1.45 | 0.64 |
| 1:2:1240:U:H2' | 1:2:1241:G:H8 | 1.62 | 0.64 |
| 16:O:126:GLU:HA | 16:O:129:LYS:HG2 | 1.80 | 0.64 |
| 19:R:10:ARG:NH1 | 19:R:10:ARG:O | 2.27 | 0.64 |
| 34:g:39:THR:HG22 | 34:g:60:ARG:HG2 | 1.79 | 0.64 |
| 1:2:1276:G:H1 | 1:2:1313:U:H3 | 1.45 | 0.64 |
| 24:W:61:LEU:HD23 | 24:W:82:MET:HG2 | 1.79 | 0.64 |
| 1:2:1519:G:O2' | 38:y:29:G:OP1 | 2.14 | 0.64 |
| 8:G:43:PRO:HD2 | 8:G:46:ILE:HD12 | 1.80 | 0.64 |
| 19:R:17:TYR:HA | 22:U:91:LYS:HG2 | 1.80 | 0.64 |
| 1:2:1333:C:HO2' | 24:W:68:THR:HG1 | 1.44 | 0.63 |
| 1:2:1588:C:OP2 | 35:h:104:ARG:NH1 | 2.31 | 0.63 |
| 1:2:1007:A:H5'' | 17:P:3:ARG:HH22 | 1.62 | 0.63 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 15:N:102:PHE:CE1 | 27:Z:7:LEU:HB3 | 2.33 | 0.63 |
| 28:a:11:LYS:HB2 | 28:a:24:VAL:HG12 | 1.80 | 0.63 |
| 39:z:595:C:H2' | 39:z:596:A:C8 | 2.33 | 0.63 |
| 1:2:1491:G:N2 | 32:e:45:GLN:OE1 | 2.31 | 0.63 |
| 5:D:136:HIS:HB3 | 5:D:216:LYS:HB2 | 1.79 | 0.63 |
| 16:O:35:ILE:HG13 | 33:f:103:LEU:HD13 | 1.80 | 0.63 |
| 18:Q:44:VAL:HG13 | 18:Q:53:ILE:HG13 | 1.81 | 0.63 |
| 1:2:606:A:H5'' | 27:Z:68:LYS:HZ1 | 1.63 | 0.63 |
| 1:2:1282:G:N2 | 1:2:1308:G:O2' | 2.26 | 0.63 |
| 4:C:113:GLN:HB2 | 4:C:116:PHE:HB2 | 1.81 | 0.63 |
| 30:c:19:HIS:ND1 | 30:c:21:LYS:HB3 | 2.13 | 0.63 |
| 1:2:948:G:H21 | 18:Q:52:THR:HG21 | 1.63 | 0.63 |
| 1:2:980:C:O2' | 18:Q:138:ASP:O | 2.11 | 0.63 |
| 1:2:1054:A:OP1 | 38:y:37:A:O2' | 2.17 | 0.63 |
| 1:2:1560:C:N4 | 1:2:1561:G:O6 | 2.31 | 0.63 |
| 1:2:1655:C:O5' | 32:e:32:ARG:NH2 | 2.32 | 0.63 |
| 6:E:187:THR:N | 6:E:206:ASP:OD2 | 2.32 | 0.63 |
| 7:F:1:MET:HG2 | 7:F:4:GLN:HB2 | 1.78 | 0.63 |
| 13:L:93:LYS:HB2 | 13:L:96:TYR:HD2 | 1.64 | 0.63 |
| 17:P:62:GLN:HG3 | 17:P:65:PHE:CD2 | 2.30 | 0.63 |
| 38:y:11:C:N4 | 38:y:24:U:N3 | 2.46 | 0.63 |
| 38:y:52:G:H21 | 38:y:53:A:H62 | 1.47 | 0.63 |
| 1:2:370:G:H5'' | 12:K:31:ARG:HH21 | 1.64 | 0.63 |
| 1:2:874:G:H22 | 1:2:904:A:H2 | 1.44 | 0.63 |
| 1:2:1139:A:H2' | 1:2:1140:A:C8 | 2.34 | 0.63 |
| 1:2:1558:G:OP1 | 23:V:121:ARG:NH1 | 2.32 | 0.63 |
| 10:I:85:ARG:O | 10:I:87:ARG:NH1 | 2.32 | 0.63 |
| 7:F:204:LEU:HD13 | 7:F:209:SER:H | 1.63 | 0.63 |
| 18:Q:142:ARG:NH1 | 29:b:27:ALA:HB1 | 2.14 | 0.63 |
| 1:2:1198:U:O2 | 6:E:100:GLN:NE2 | 2.21 | 0.62 |
| 5:D:179:ASN:HB3 | 5:D:183:GLU:HG3 | 1.80 | 0.62 |
| 12:K:191:GLU:O | 15:N:18:GLN:NE2 | 2.32 | 0.62 |
| 1:2:483:A:H62 | 1:2:499:G:H21 | 1.47 | 0.62 |
| 5:D:103:MET:HG3 | 5:D:215:VAL:HB | 1.81 | 0.62 |
| 6:E:79:ILE:HG12 | 6:E:147:ILE:HD12 | 1.81 | 0.62 |
| 1:2:1802:U:H2' | 1:2:1803:A:C8 | 2.34 | 0.62 |
| 1:2:1508:C:H4' | 32:e:7:TYR:HE2 | 1.64 | 0.62 |
| 31:d:20:ARG:NH2 | 31:d:25:GLY:O | 2.31 | 0.62 |
| 5:D:72:ALA:O | 5:D:76:ASN:ND2 | 2.32 | 0.62 |
| 5:D:107:ARG:HA | 5:D:110:MET:HE2 | 1.80 | 0.62 |
| 14:M:48:ALA:O | 14:M:51:SER:OG | 2.14 | 0.62 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 1:2:645:A:H4' | 1:2:646:G:H3' | 1.80 | 0.62 |
| 9:H:175:ASP:OD1 | 9:H:179:ASN:ND2 | 2.33 | 0.62 |
| 18:Q:32:HIS:HA | 18:Q:96:LYS:HB3 | 1.80 | 0.62 |
| 19:R:8:LYS:HE2 | 19:R:14:LYS:HD2 | 1.82 | 0.62 |
| 22:U:26:ILE:HG12 | 22:U:59:LEU:HD11 | 1.80 | 0.62 |
| 1:2:155:G:H2' | 1:2:156:G:H8 | 1.65 | 0.62 |
| 1:2:162:C:O2' | 10:I:95:LYS:NZ | 2.31 | 0.62 |
| 8:G:128:LYS:H | 8:G:140:VAL:HB | 1.62 | 0.62 |
| 11:J:46:THR:HG21 | 11:J:97:GLN:HG2 | 1.80 | 0.62 |
| 21:T:31:ASN:ND2 | 21:T:55:THR:OG1 | 2.27 | 0.62 |
| 1:2:109:U:O2' | 15:N:71:ARG:NH2 | 2.33 | 0.62 |
| 1:2:1409:G:H2' | 1:2:1410:A:H8 | 1.64 | 0.62 |
| 1:2:1736:U:OP1 | 12:K:42:ARG:NH1 | 2.33 | 0.62 |
| 5:D:23:ASP:O | 5:D:27:LYS:NZ | 2.33 | 0.62 |
| 8:G:10:LYS:HA | 8:G:27:PHE:HA | 1.79 | 0.62 |
| 30:c:18:LYS:O | 30:c:23:ARG:NH1 | 2.28 | 0.62 |
| 1:2:965:U:H3' | 1:2:966:G:H21 | 1.63 | 0.62 |
| 1:2:1090:C:O2 | 26:Y:16:ASN:ND2 | 2.32 | 0.62 |
| 1:2:1534:U:H2' | 1:2:1535:G:C8 | 2.35 | 0.62 |
| 5:D:51:ARG:O | 5:D:53:GLN:NE2 | 2.31 | 0.62 |
| 7:F:135:GLU:N | 7:F:135:GLU:OE1 | 2.32 | 0.62 |
| 28:a:114:MET:HE2 | 28:a:124:ASN:HD21 | 1.65 | 0.62 |
| 34:g:258:ILE:HG23 | 34:g:267:VAL:HB | 1.81 | 0.62 |
| 38:y:57:A:O2' | 38:y:59:A:OP2 | 2.18 | 0.62 |
| 1:2:76:U:H4' | 10:I:154:ARG:HA | 1.82 | 0.61 |
| 1:2:640:A:OP2 | 27:Z:108:LYS:NZ | 2.28 | 0.61 |
| 5:D:28:LYS:NZ | 18:Q:51:GLU:OE2 | 2.33 | 0.61 |
| 15:N:136:LYS:O | 15:N:139:ARG:NH1 | 2.33 | 0.61 |
| 21:T:106:LEU:HD13 | 21:T:109:LEU:HD13 | 1.80 | 0.61 |
| 1:2:951:A:H4' | 1:2:952:G:H4' | 1.81 | 0.61 |
| 7:F:12:VAL:HG11 | 32:e:36:LEU:HD21 | 1.82 | 0.61 |
| 8:G:88:ASP:OD1 | 8:G:122:LYS:NZ | 2.33 | 0.61 |
| 11:J:19:PHE:O | 11:J:23:ILE:HD12 | 2.00 | 0.61 |
| 1:2:1013:U:OP2 | 17:P:55:ARG:NH1 | 2.33 | 0.61 |
| 1:2:1531:G:H2' | 1:2:1532:A:H8 | 1.65 | 0.61 |
| 1:2:1755:U:O2 | 1:2:1765:G:N2 | 2.33 | 0.61 |
| 19:R:34:MET:SD | 19:R:35:GLN:NE2 | 2.73 | 0.61 |
| 7:F:62:LYS:HA | 14:M:96:ARG:HG3 | 1.82 | 0.61 |
| 15:N:102:PHE:O | 27:Z:8:ARG:HA | 2.00 | 0.61 |
| 1:2:1300:U:H2' | 1:2:1301:C:C6 | 2.36 | 0.61 |
| 39:z:525:A:H3' | 39:z:526:G:H21 | 1.66 | 0.61 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 1:2:185:C:H2' | 1:2:186:G:H8 | 1.65 | 0.61 |
| 12:K:31:ARG:HD2 | 12:K:56:ARG:HH21 | 1.66 | 0.61 |
| 1:2:914:U:O2' | 26:Y:57:ARG:NH1 | 2.34 | 0.61 |
| 1:2:1139:A:OP1 | 6:E:175:SER:OG | 2.18 | 0.61 |
| 1:2:1382:A:OP2 | 7:F:160:SER:OG | 2.16 | 0.61 |
| 17:P:118:ILE:HG22 | 17:P:121:ARG:HH12 | 1.66 | 0.61 |
| 30:c:11:SER:HB2 | 30:c:15:GLU:HB2 | 1.83 | 0.61 |
| 35:h:57:LYS:HE3 | 35:h:61:GLU:HB3 | 1.82 | 0.61 |
| 1:2:124:U:OP1 | 8:G:148:ARG:NH2 | 2.33 | 0.61 |
| 1:2:202:U:H2' | 1:2:203:G:C8 | 2.34 | 0.61 |
| 1:2:378:U:H2' | 1:2:379:A:H8 | 1.66 | 0.61 |
| 6:E:153:GLY:N | 6:E:164:THR:O | 2.30 | 0.61 |
| 10:I:1:MET:N | 10:I:18:VAL:O | 2.34 | 0.61 |
| 1:2:1284:U:H3 | 1:2:1307:C:H42 | 0.72 | 0.61 |
| 9:H:77:MET:O | 9:H:83:ASN:ND2 | 2.34 | 0.61 |
| 21:T:15:VAL:O | 21:T:19:LYS:HG2 | 2.01 | 0.61 |
| 23:V:28:LEU:HD22 | 23:V:110:LEU:HD11 | 1.83 | 0.61 |
| 1:2:629:C:H2' | 1:2:630:A:H8 | 1.66 | 0.61 |
| 1:2:1295:A:OP1 | 19:R:59:ARG:NH1 | 2.34 | 0.61 |
| 7:F:175:VAL:HG23 | 7:F:182:LEU:HB2 | 1.83 | 0.61 |
| 12:K:10:LYS:HD2 | 15:N:136:LYS:HE2 | 1.82 | 0.61 |
| 19:R:10:ARG:CZ | 19:R:17:TYR:HB3 | 2.31 | 0.61 |
| 19:R:86:LEU:H | 19:R:89:MET:HE2 | 1.65 | 0.61 |
| 1:2:832:G:N2 | 1:2:834:G:OP2 | 2.33 | 0.60 |
| 1:2:1179:A:OP2 | 37:l:18:ARG:NH1 | 2.33 | 0.60 |
| 1:2:1427:G:H2' | 1:2:1428:U:C5 | 2.35 | 0.60 |
| 28:a:12:PHE:HE1 | 28:a:21:LYS:HB3 | 1.65 | 0.60 |
| 30:c:20:LYS:HD2 | 30:c:28:PRO:HA | 1.81 | 0.60 |
| 5:D:80:ALA:O | 5:D:83:LYS:NZ | 2.26 | 0.60 |
| 8:G:26:VAL:HG13 | 8:G:27:PHE:HD1 | 1.65 | 0.60 |
| 39:z:599:G:H2' | 39:z:600:G:C8 | 2.37 | 0.60 |
| 1:2:106:C:H2' | 1:2:107:A:C8 | 2.36 | 0.60 |
| 1:2:866:A:N6 | 1:2:912:A:O5' | 2.33 | 0.60 |
| 1:2:1546:U:H2' | 1:2:1547:G:C4 | 2.37 | 0.60 |
| 4:C:4:ALA:HB3 | 4:C:8:LEU:HB2 | 1.84 | 0.60 |
| 24:W:70:CYS:H | 32:e:40:ARG:HH22 | 1.48 | 0.60 |
| 1:2:1659:A:O2' | 1:2:1660:G:H5' | 2.02 | 0.60 |
| 12:K:45:THR:HG22 | 12:K:53:LYS:HD2 | 1.82 | 0.60 |
| 1:2:345:G:OP1 | 15:N:105:ARG:NH1 | 2.34 | 0.60 |
| 1:2:599:U:H2' | 1:2:600:G:C8 | 2.37 | 0.60 |
| 1:2:1344:G:H22 | 1:2:1377:G:H1 | 1.49 | 0.60 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 1:2:1371:G:H2' | 1:2:1372:A:C8 | 2.35 | 0.60 |
| 1:2:1724:U:O2 | 1:2:1799:G:N2 | 2.33 | 0.60 |
| 6:E:189:ILE:HB | 6:E:196:LYS:HD3 | 1.83 | 0.60 |
| 15:N:40:ILE:HG23 | 15:N:68:ILE:HD13 | 1.83 | 0.60 |
| 18:Q:27:VAL:H | 18:Q:91:THR:HG1 | 1.48 | 0.60 |
| 20:S:37:ARG:HG2 | 23:V:7:LYS:HB3 | 1.84 | 0.60 |
| 28:a:106:GLN:HE21 | 28:a:107:ARG:HG3 | 1.65 | 0.60 |
| 39:z:589:C:H2' | 39:z:590:A:C8 | 2.37 | 0.60 |
| 1:2:378:U:H2' | 1:2:379:A:C8 | 2.36 | 0.60 |
| 1:2:444:U:H2' | 1:2:445:A:H8 | 1.66 | 0.60 |
| 9:H:54:GLY:O | 20:S:125:ARG:NH1 | 2.33 | 0.60 |
| 12:K:165:GLN:HG3 | 12:K:170:LYS:O | 2.01 | 0.60 |
| 15:N:101:ARG:NH1 | 27:Z:6:GLY:O | 2.34 | 0.60 |
| 20:S:146:ARG:NH1 | 38:y:34:A:OP2 | 2.34 | 0.60 |
| 34:g:5:MET:HE1 | 34:g:312:VAL:HG22 | 1.83 | 0.60 |
| 1:2:1300:U:H5'' | 33:f:92:LYS:HD2 | 1.82 | 0.60 |
| 4:C:58:LEU:HD11 | 4:C:178:LEU:HB3 | 1.82 | 0.60 |
| 19:R:17:TYR:HA | 22:U:91:LYS:HA | 1.83 | 0.60 |
| 1:2:1224:A:H2' | 1:2:1225:G:C8 | 2.37 | 0.60 |
| 4:C:68:ILE:HD11 | 4:C:121:LEU:HG | 1.84 | 0.60 |
| 4:C:122:LEU:HD23 | 4:C:142:LEU:HG | 1.83 | 0.60 |
| 13:L:170:PRO:HB3 | 13:L:174:LYS:HD2 | 1.84 | 0.60 |
| 9:H:71:ARG:NH2 | 9:H:148:ASN:OD1 | 2.34 | 0.60 |
| 11:J:46:THR:HB | 11:J:65:PRO:HG3 | 1.84 | 0.60 |
| 1:2:668:U:OP2 | 1:2:1022:C:N4 | 2.35 | 0.59 |
| 1:2:1626:U:O3' | 22:U:34:LYS:NZ | 2.35 | 0.59 |
| 14:M:32:HIS:HB2 | 14:M:40:VAL:HB | 1.84 | 0.59 |
| 24:W:25:THR:O | 24:W:111:GLU:HB2 | 2.01 | 0.59 |
| 24:W:82:MET:HE1 | 32:e:52:PHE:HB2 | 1.84 | 0.59 |
| 30:c:34:ASP:HB3 | 30:c:80:ARG:HG3 | 1.84 | 0.59 |
| 32:e:31:ILE:HG23 | 32:e:33:LYS:HZ3 | 1.66 | 0.59 |
| 1:2:360:G:N2 | 1:2:362:U:O4 | 2.35 | 0.59 |
| 22:U:115:LYS:NZ | 39:z:566:C:OP1 | 2.35 | 0.59 |
| 23:V:10:ASN:H | 23:V:144:LYS:HE2 | 1.66 | 0.59 |
| 24:W:51:LYS:HD2 | 24:W:90:ASP:HB3 | 1.84 | 0.59 |
| 39:z:564:G:H2' | 39:z:565:G:H8 | 1.68 | 0.59 |
| 1:2:483:A:N6 | 1:2:500:G:H1' | 2.14 | 0.59 |
| 1:2:1571:G:H2' | 1:2:1572:G:C8 | 2.37 | 0.59 |
| 1:2:220:C:H2' | 1:2:221:A:C8 | 2.37 | 0.59 |
| 1:2:1259:U:O2 | 32:e:16:GLN:NE2 | 2.36 | 0.59 |
| 1:2:1587:C:H5'' | 9:H:91:ARG:HH12 | 1.67 | 0.59 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|------------------|--------------------------|-------------------|
| 10:I:42:GLY:O | 10:I:46:LYS:NZ | 2.35 | 0.59 |
| 15:N:94:HIS:HB2 | 15:N:105:ARG:HD2 | 1.83 | 0.59 |
| 1:2:1281:G:N1 | 33:f:103:LEU:O | 2.35 | 0.59 |
| 7:F:193:ASP:HB3 | 7:F:194:PRO:HD3 | 1.83 | 0.59 |
| 19:R:85:ILE:HD13 | 19:R:111:MET:HB3 | 1.83 | 0.59 |
| 28:a:87:PRO:HD2 | 28:a:90:ARG:HD2 | 1.83 | 0.59 |
| 1:2:1199:G:H4' | 6:E:101:THR:HA | 1.85 | 0.59 |
| 1:2:1540:A:H2' | 1:2:1541:G:C8 | 2.37 | 0.59 |
| 1:2:1775:A:H2' | 1:2:1776:G:C8 | 2.37 | 0.59 |
| 29:b:19:GLN:NE2 | 29:b:20:PRO:O | 2.34 | 0.59 |
| 1:2:537:G:H2' | 1:2:538:C:C6 | 2.38 | 0.59 |
| 1:2:784:G:H4' | 10:I:237:LEU:HB2 | 1.83 | 0.59 |
| 1:2:800:U:OP1 | 26:Y:82:GLN:NE2 | 2.35 | 0.59 |
| 1:2:1493:G:N7 | 14:M:25:LYS:NZ | 2.51 | 0.59 |
| 4:C:140:VAL:HA | 6:E:72:PRO:HG2 | 1.84 | 0.59 |
| 13:L:152:ASP:OD1 | 13:L:153:SER:N | 2.36 | 0.59 |
| 30:c:1:MET:HG3 | 30:c:3:LEU:H | 1.68 | 0.59 |
| 1:2:155:G:H2' | 1:2:156:G:C8 | 2.37 | 0.59 |
| 1:2:538:C:H2' | 1:2:539:C:H6 | 1.68 | 0.59 |
| 17:P:58:HIS:HB2 | 17:P:60:VAL:HG12 | 1.84 | 0.59 |
| 19:R:38:SER:HB2 | 19:R:41:GLN:HB2 | 1.85 | 0.59 |
| 27:Z:71:ARG:HG2 | 27:Z:82:THR:HG22 | 1.84 | 0.59 |
| 34:g:31:ILE:HG22 | 34:g:43:TRP:HB2 | 1.84 | 0.59 |
| 1:2:1259:U:H1' | 32:e:16:GLN:HE21 | 1.66 | 0.59 |
| 19:R:32:GLN:OE1 | 19:R:32:GLN:N | 2.35 | 0.59 |
| 1:2:1161:G:OP2 | 1:2:1161:G:N2 | 2.30 | 0.58 |
| 1:2:1648:U:O2 | 1:2:1666:G:N2 | 2.22 | 0.58 |
| 33:f:105:TYR:HB2 | 33:f:131:PHE:HE2 | 1.68 | 0.58 |
| 1:2:16:G:H2' | 1:2:17:C:C6 | 2.37 | 0.58 |
| 1:2:918:A:OP1 | 26:Y:28:ARG:NH1 | 2.29 | 0.58 |
| 1:2:1083:A:OP1 | 29:b:3:LYS:NZ | 2.23 | 0.58 |
| 1:2:1623:C:H2' | 1:2:1624:C:C6 | 2.38 | 0.58 |
| 4:C:54:THR:HG22 | 4:C:162:PRO:HB2 | 1.85 | 0.58 |
| 5:D:121:ILE:HD12 | 5:D:161:VAL:HG13 | 1.85 | 0.58 |
| 8:G:63:LYS:HA | 8:G:66:MET:HE2 | 1.85 | 0.58 |
| 16:O:13:ASP:HB3 | 16:O:16:THR:HG22 | 1.85 | 0.58 |
| 35:h:82:SER:O | 35:h:85:ARG:HG2 | 2.03 | 0.58 |
| 17:P:45:LEU:HB3 | 17:P:49:GLN:HB2 | 1.85 | 0.58 |
| 23:V:87:VAL:HG13 | 23:V:88:MET:HE2 | 1.85 | 0.58 |
| 39:z:564:G:H2' | 39:z:565:G:C8 | 2.38 | 0.58 |
| 1:2:671:U:H5'' | 27:Z:8:ARG:HG3 | 1.85 | 0.58 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 1:2:65:C:N4 | 1:2:169:U:O2' | 2.37 | 0.58 |
| 1:2:1534:U:H2' | 1:2:1535:G:H8 | 1.67 | 0.58 |
| 4:C:89:LYS:HE3 | 4:C:201:LEU:HG | 1.86 | 0.58 |
| 4:C:130:ASP:HB3 | 4:C:133:PRO:HG2 | 1.85 | 0.58 |
| 21:T:12:ALA:O | 21:T:16:ILE:HG12 | 2.04 | 0.58 |
| 24:W:66:ARG:HA | 24:W:77:TRP:CD1 | 2.39 | 0.58 |
| 1:2:1193:G:H2' | 1:2:1194:G:H8 | 1.66 | 0.58 |
| 6:E:101:THR:HG23 | 6:E:103:ALA:H | 1.69 | 0.58 |
| 10:I:18:VAL:HB | 10:I:23:LYS:HZ1 | 1.68 | 0.58 |
| 33:f:139:HIS:HB2 | 33:f:150:PHE:HB2 | 1.85 | 0.58 |
| 38:y:11:C:N3 | 38:y:24:U:O2 | 2.37 | 0.58 |
| 1:2:644:A:OP2 | 1:2:645:A:O2' | 2.18 | 0.58 |
| 1:2:1647:G:N2 | 1:2:1667:U:O2 | 2.24 | 0.58 |
| 1:2:1847:C:H2' | 1:2:1848:U:C6 | 2.39 | 0.58 |
| 11:J:63:PHE:HB3 | 11:J:97:GLN:HB3 | 1.85 | 0.58 |
| 11:J:157:HIS:HB3 | 11:J:190:PRO:HG3 | 1.85 | 0.58 |
| 1:2:158:A:H2' | 1:2:159:A:C8 | 2.39 | 0.58 |
| 1:2:165:G:H2' | 1:2:166:A:C8 | 2.38 | 0.58 |
| 1:2:427:G:H3' | 1:2:428:G:H21 | 1.67 | 0.58 |
| 12:K:3:ILE:HB | 12:K:30:GLY:HA3 | 1.84 | 0.58 |
| 17:P:23:PRO:HB2 | 17:P:25:TRP:CZ3 | 2.38 | 0.58 |
| 19:R:18:ARG:HE | 19:R:36:LEU:HB3 | 1.68 | 0.58 |
| 7:F:115:VAL:HG11 | 7:F:142:LEU:HD21 | 1.86 | 0.58 |
| 7:F:140:GLY:HA3 | 7:F:182:LEU:HD23 | 1.85 | 0.58 |
| 8:G:197:ASN:HB3 | 8:G:209:HIS:HB2 | 1.83 | 0.58 |
| 11:J:129:ILE:HD11 | 11:J:180:LEU:HD13 | 1.86 | 0.58 |
| 12:K:23:LYS:O | 12:K:25:ARG:NH1 | 2.37 | 0.58 |
| 17:P:106:ARG:NH1 | 17:P:106:ARG:O | 2.36 | 0.58 |
| 22:U:63:GLU:OE1 | 22:U:66:ARG:NH2 | 2.36 | 0.58 |
| 23:V:57:ALA:O | 23:V:60:THR:OG1 | 2.15 | 0.58 |
| 1:2:633:A:OP2 | 13:L:38:ARG:NH1 | 2.36 | 0.58 |
| 11:J:41:ARG:HH11 | 11:J:42:GLU:H | 1.52 | 0.58 |
| 14:M:17:LYS:HA | 14:M:89:ILE:HD11 | 1.85 | 0.58 |
| 23:V:44:GLU:OE1 | 23:V:44:GLU:N | 2.37 | 0.58 |
| 28:a:20:ARG:HD2 | 28:a:74:MET:HE2 | 1.86 | 0.58 |
| 29:b:10:ARG:HB3 | 29:b:33:ASP:O | 2.04 | 0.58 |
| 1:2:116:U:O4 | 1:2:337:G:O6 | 2.21 | 0.57 |
| 1:2:332:C:H2' | 1:2:333:A:H8 | 1.68 | 0.57 |
| 1:2:403:G:N2 | 1:2:415:G:H1' | 2.19 | 0.57 |
| 1:2:1255:A:N6 | 1:2:1514:U:O5' | 2.37 | 0.57 |
| 1:2:1562:G:H21 | 1:2:1562:G:P | 2.26 | 0.57 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 1:2:1598:G:N7 | 22:U:24:ARG:NH1 | 2.51 | 0.57 |
| 9:H:20:PHE:HA | 9:H:48:TYR:HA | 1.87 | 0.57 |
| 19:R:17:TYR:OH | 19:R:18:ARG:NH2 | 2.37 | 0.57 |
| 22:U:61:GLU:HA | 22:U:64:VAL:HG12 | 1.86 | 0.57 |
| 27:Z:100:VAL:HG12 | 27:Z:125:VAL:HG12 | 1.85 | 0.57 |
| 28:a:5:VAL:O | 28:a:43:LYS:NZ | 2.37 | 0.57 |
| 1:2:538:C:H2' | 1:2:539:C:C6 | 2.38 | 0.57 |
| 1:2:1328:A:O2' | 7:F:145:GLN:O | 2.21 | 0.57 |
| 1:2:1574:A:O2' | 1:2:1576:C:OP2 | 2.20 | 0.57 |
| 5:D:139:CYS:HA | 5:D:212:VAL:HA | 1.86 | 0.57 |
| 11:J:97:GLN:O | 11:J:98:ARG:NE | 2.37 | 0.57 |
| 14:M:8:ARG:HG2 | 14:M:12:TYR:HE2 | 1.69 | 0.57 |
| 17:P:96:VAL:HG23 | 17:P:99:ARG:HE | 1.69 | 0.57 |
| 23:V:76:THR:HG22 | 23:V:94:ARG:HB3 | 1.85 | 0.57 |
| 38:y:3:C:H2' | 38:y:4:A:H8 | 1.66 | 0.57 |
| 1:2:1388:U:H3 | 1:2:1474:U:H3 | 1.51 | 0.57 |
| 1:2:1716:U:H4' | 1:2:1717:G:H5'' | 1.86 | 0.57 |
| 20:S:124:PRO:O | 20:S:126:ARG:NH1 | 2.37 | 0.57 |
| 23:V:33:TRP:HZ2 | 23:V:102:ARG:HG3 | 1.68 | 0.57 |
| 1:2:125:C:OP2 | 10:I:198:ARG:NH2 | 2.37 | 0.57 |
| 1:2:921:G:H5'' | 17:P:91:LEU:HD21 | 1.86 | 0.57 |
| 1:2:1140:A:H2' | 1:2:1141:A:C8 | 2.39 | 0.57 |
| 3:B:261:SER:HA | 3:B:279:ALA:HA | 1.86 | 0.57 |
| 29:b:44:ILE:HB | 29:b:64:LEU:HD11 | 1.86 | 0.57 |
| 1:2:431:C:H2' | 1:2:432:C:C6 | 2.40 | 0.57 |
| 1:2:1019:A:H2' | 1:2:1020:A:C8 | 2.40 | 0.57 |
| 1:2:1160:G:H2' | 1:2:1161:G:N3 | 2.19 | 0.57 |
| 1:2:1213:A:H2' | 1:2:1214:C:C6 | 2.39 | 0.57 |
| 15:N:133:PRO:HG3 | 15:N:139:ARG:HG3 | 1.86 | 0.57 |
| 26:Y:53:ILE:N | 26:Y:60:LYS:O | 2.21 | 0.57 |
| 1:2:168:C:O2' | 10:I:133:LEU:O | 2.21 | 0.57 |
| 1:2:291:U:H2' | 1:2:292:A:H8 | 1.68 | 0.57 |
| 6:E:191:SER:CB | 6:E:196:LYS:HE3 | 2.35 | 0.57 |
| 7:F:8:LYS:HE2 | 24:W:59:LYS:HG2 | 1.86 | 0.57 |
| 24:W:50:VAL:HG23 | 24:W:91:LEU:HG | 1.86 | 0.57 |
| 26:Y:11:LEU:O | 26:Y:15:ASN:ND2 | 2.37 | 0.57 |
| 28:a:15:ASN:ND2 | 28:a:20:ARG:HE | 2.02 | 0.57 |
| 1:2:788:C:HO2' | 1:2:789:G:H8 | 1.51 | 0.57 |
| 1:2:842:G:OP2 | 8:G:108:ARG:NH2 | 2.38 | 0.57 |
| 1:2:1409:G:H2' | 1:2:1410:A:C8 | 2.39 | 0.57 |
| 17:P:61:ALA:HB2 | 30:c:32:PHE:HZ | 1.69 | 0.57 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 20:S:40:GLU:O | 20:S:41:MET:HE2 | 2.05 | 0.57 |
| 25:X:21:ASN:HD21 | 26:Y:66:THR:HB | 1.70 | 0.57 |
| 1:2:1211:C:H42 | 1:2:1216:A:N6 | 2.02 | 0.57 |
| 5:D:168:MET:O | 5:D:172:MET:HG3 | 2.05 | 0.57 |
| 12:K:48:VAL:HG22 | 12:K:49:ARG:H | 1.68 | 0.57 |
| 14:M:59:LYS:HE2 | 14:M:72:THR:HG22 | 1.86 | 0.57 |
| 34:g:88:ARG:HB3 | 34:g:90:TRP:NE1 | 2.18 | 0.57 |
| 1:2:732:C:N4 | 1:2:733:G:O6 | 2.37 | 0.57 |
| 1:2:1375:A:H2' | 1:2:1376:C:C6 | 2.40 | 0.57 |
| 19:R:78:THR:N | 19:R:95:GLY:O | 2.36 | 0.57 |
| 20:S:11:GLN:HA | 20:S:24:HIS:HA | 1.86 | 0.57 |
| 28:a:7:ILE:HG23 | 28:a:25:ILE:HD11 | 1.86 | 0.57 |
| 36:i:89:GLN:N | 36:i:89:GLN:OE1 | 2.38 | 0.57 |
| 1:2:149:A:N7 | 1:2:169:U:C4 | 2.73 | 0.56 |
| 1:2:833:A:H5'' | 28:a:47:MET:HE1 | 1.85 | 0.56 |
| 1:2:957:G:H4' | 9:H:131:ALA:HB1 | 1.87 | 0.56 |
| 1:2:1320:G:H2' | 1:2:1321:G:C8 | 2.40 | 0.56 |
| 1:2:1664:G:H5' | 24:W:79:ARG:HH22 | 1.69 | 0.56 |
| 1:2:1712:C:O2' | 37:l:21:ARG:NH2 | 2.38 | 0.56 |
| 6:E:83:LEU:HB3 | 6:E:87:LEU:HD21 | 1.86 | 0.56 |
| 22:U:123:LEU:HB3 | 22:U:127:TRP:CZ3 | 2.40 | 0.56 |
| 38:y:18:G:O5' | 38:y:59:A:N6 | 2.38 | 0.56 |
| 1:2:515:A:N6 | 1:2:579:G:O6 | 2.39 | 0.56 |
| 1:2:1732:G:O6 | 1:2:1791:U:C2 | 2.58 | 0.56 |
| 1:2:1863:A:N6 | 5:D:114:VAL:O | 2.38 | 0.56 |
| 4:C:187:GLY:O | 25:X:45:ARG:NH2 | 2.34 | 0.56 |
| 24:W:86:LYS:O | 24:W:87:ARG:NH1 | 2.35 | 0.56 |
| 25:X:31:SER:OG | 25:X:57:GLY:N | 2.36 | 0.56 |
| 38:y:56:G:O2' | 38:y:57:A:O4' | 2.23 | 0.56 |
| 1:2:28:U:H2' | 1:2:29:G:C8 | 2.40 | 0.56 |
| 1:2:74:G:N7 | 10:I:170:ARG:NH2 | 2.49 | 0.56 |
| 1:2:1088:G:H2' | 1:2:1089:A:H8 | 1.70 | 0.56 |
| 1:2:1751:G:O6 | 1:2:1769:U:O4 | 2.23 | 0.56 |
| 10:I:5:ILE:HG22 | 10:I:111:LEU:HD21 | 1.87 | 0.56 |
| 20:S:90:LYS:HE2 | 20:S:120:LEU:HA | 1.87 | 0.56 |
| 34:g:283:PRO:O | 34:g:285:GLN:NE2 | 2.37 | 0.56 |
| 37:l:8:LYS:HG3 | 37:l:12:ARG:NH1 | 2.20 | 0.56 |
| 38:y:54:U:N3 | 38:y:57:A:OP2 | 2.38 | 0.56 |
| 1:2:978:G:H2' | 1:2:979:A:H8 | 1.70 | 0.56 |
| 1:2:1545:G:H3' | 1:2:1574:A:H61 | 1.71 | 0.56 |
| 4:C:31:ASP:HB2 | 4:C:151:ASP:HA | 1.85 | 0.56 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 4:C:149:ASN:OD1 | 4:C:166:LYS:NZ | 2.39 | 0.56 |
| 19:R:94:VAL:HG12 | 19:R:107:ILE:HD11 | 1.86 | 0.56 |
| 24:W:94:PRO:HD2 | 24:W:97:ILE:HD11 | 1.86 | 0.56 |
| 25:X:40:ASP:OD2 | 25:X:43:THR:OG1 | 2.21 | 0.56 |
| 1:2:514:U:H5'' | 1:2:515:A:H5' | 1.88 | 0.56 |
| 1:2:796:U:H2' | 1:2:797:U:O4' | 2.06 | 0.56 |
| 22:U:45:LEU:HD22 | 22:U:50:ILE:HD11 | 1.86 | 0.56 |
| 1:2:62:G:O2' | 1:2:172:U:OP1 | 2.23 | 0.56 |
| 1:2:1537:C:OP1 | 23:V:62:ARG:NH2 | 2.32 | 0.56 |
| 1:2:1595:G:O3' | 35:h:43:LYS:NZ | 2.39 | 0.56 |
| 5:D:137:LEU:HB2 | 5:D:172:MET:HE1 | 1.88 | 0.56 |
| 13:L:176:LYS:O | 13:L:180:LYS:HG2 | 2.06 | 0.56 |
| 26:Y:14:ILE:HD13 | 26:Y:25:VAL:HG21 | 1.86 | 0.56 |
| 26:Y:53:ILE:HD12 | 26:Y:60:LYS:HD2 | 1.87 | 0.56 |
| 33:f:92:LYS:NZ | 33:f:94:LYS:HA | 2.21 | 0.56 |
| 1:2:202:U:H2' | 1:2:203:G:H8 | 1.71 | 0.56 |
| 1:2:518:A:H2' | 1:2:519:A:H8 | 1.69 | 0.56 |
| 1:2:604:G:H5' | 36:i:76:VAL:HG22 | 1.88 | 0.56 |
| 1:2:1246:A:H4' | 24:W:73:GLY:HA2 | 1.87 | 0.56 |
| 1:2:1611:U:OP2 | 19:R:43:ARG:NH1 | 2.36 | 0.56 |
| 1:2:1717:G:H2' | 1:2:1718:G:C8 | 2.41 | 0.56 |
| 9:H:35:LEU:HD21 | 9:H:146:ARG:HD2 | 1.87 | 0.56 |
| 10:I:93:LYS:NZ | 10:I:94:ARG:O | 2.36 | 0.56 |
| 15:N:57:ASP:OD2 | 15:N:60:CYS:N | 2.37 | 0.56 |
| 17:P:52:VAL:HA | 17:P:55:ARG:HG2 | 1.88 | 0.56 |
| 1:2:1199:G:H2' | 1:2:1200:A:C8 | 2.41 | 0.56 |
| 39:z:522:U:H2' | 39:z:523:G:H8 | 1.70 | 0.56 |
| 1:2:53:C:O2' | 1:2:497:G:N7 | 2.39 | 0.56 |
| 1:2:886:U:H5' | 1:2:887:G:H5'' | 1.88 | 0.56 |
| 1:2:1855:G:OP1 | 29:b:8:ASN:ND2 | 2.37 | 0.56 |
| 8:G:137:PRO:HG2 | 8:G:149:TYR:HA | 1.88 | 0.56 |
| 10:I:188:LYS:HG3 | 10:I:191:ARG:HH12 | 1.70 | 0.56 |
| 16:O:119:GLN:HG2 | 16:O:121:LYS:HZ3 | 1.71 | 0.56 |
| 17:P:113:PHE:O | 17:P:116:ILE:HG22 | 2.06 | 0.56 |
| 18:Q:33:ILE:N | 18:Q:96:LYS:O | 2.38 | 0.56 |
| 19:R:74:GLU:OE1 | 19:R:74:GLU:N | 2.35 | 0.56 |
| 34:g:22:ALA:HB3 | 34:g:32:LEU:HB2 | 1.88 | 0.56 |
| 34:g:88:ARG:CB | 34:g:90:TRP:HE1 | 2.17 | 0.56 |
| 10:I:44:GLU:HB3 | 10:I:48:TYR:HE2 | 1.71 | 0.56 |
| 13:L:48:PHE:O | 13:L:52:LYS:HG2 | 2.06 | 0.56 |
| 23:V:57:ALA:HB1 | 23:V:107:LEU:HD11 | 1.88 | 0.56 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 23:V:126:GLN:NE2 | 23:V:126:GLN:O | 2.38 | 0.56 |
| 34:g:129:ILE:HB | 34:g:142:VAL:HB | 1.88 | 0.56 |
| 1:2:149:A:N6 | 1:2:169:U:N3 | 2.53 | 0.55 |
| 1:2:1304:U:H4' | 33:f:133:ALA:HB1 | 1.87 | 0.55 |
| 26:Y:10:ALA:O | 26:Y:14:ILE:HG12 | 2.05 | 0.55 |
| 1:2:72:C:O4' | 1:2:74:G:N2 | 2.39 | 0.55 |
| 1:2:1232:G:H21 | 1:2:1517:A:N6 | 2.03 | 0.55 |
| 1:2:1347:G:H2' | 1:2:1348:G:H8 | 1.70 | 0.55 |
| 6:E:238:PRO:HA | 6:E:241:TRP:CE2 | 2.41 | 0.55 |
| 7:F:6:SER:O | 7:F:10:LYS:N | 2.38 | 0.55 |
| 8:G:71:LYS:HB2 | 8:G:91:SER:HB2 | 1.88 | 0.55 |
| 14:M:2:LEU:HG | 14:M:3:MET:HG3 | 1.88 | 0.55 |
| 28:a:88:LYS:HB2 | 28:a:97:TYR:HE2 | 1.71 | 0.55 |
| 39:z:597:A:H2' | 39:z:598:A:C8 | 2.41 | 0.55 |
| 1:2:192:U:O2 | 1:2:203:G:N2 | 2.28 | 0.55 |
| 1:2:1170:U:H2' | 1:2:1171:G:C8 | 2.41 | 0.55 |
| 6:E:151:ARG:HD3 | 6:E:166:ARG:NE | 2.21 | 0.55 |
| 10:I:130:PRO:O | 10:I:132:ARG:NH1 | 2.40 | 0.55 |
| 19:R:60:LEU:HD13 | 19:R:89:MET:HG3 | 1.88 | 0.55 |
| 20:S:8:GLN:HB2 | 20:S:27:ARG:HB2 | 1.87 | 0.55 |
| 1:2:1276:G:H22 | 1:2:1313:U:H3 | 1.54 | 0.55 |
| 1:2:1405:A:H2' | 1:2:1406:C:O4' | 2.07 | 0.55 |
| 1:2:1640:C:O3' | 20:S:138:ARG:NE | 2.40 | 0.55 |
| 2:A:17:GLU:N | 2:A:73:VAL:O | 2.32 | 0.55 |
| 4:C:7:VAL:HG11 | 25:X:43:THR:HA | 1.87 | 0.55 |
| 9:H:27:ASP:O | 9:H:29:GLN:NE2 | 2.39 | 0.55 |
| 12:K:100:CYS:SG | 12:K:101:ILE:N | 2.79 | 0.55 |
| 1:2:367:G:OP1 | 12:K:99:ASN:ND2 | 2.39 | 0.55 |
| 7:F:141:LYS:HE2 | 7:F:180:GLY:HA3 | 1.89 | 0.55 |
| 8:G:49:ARG:NH1 | 8:G:56:LEU:O | 2.39 | 0.55 |
| 8:G:60:GLU:O | 8:G:64:ILE:HG12 | 2.07 | 0.55 |
| 9:H:75:SER:O | 9:H:76:MET:HE2 | 2.07 | 0.55 |
| 11:J:84:GLU:HG3 | 11:J:92:VAL:HG12 | 1.87 | 0.55 |
| 24:W:38:ASP:OD1 | 24:W:41:ARG:NH1 | 2.39 | 0.55 |
| 33:f:92:LYS:HZ1 | 33:f:94:LYS:HA | 1.72 | 0.55 |
| 38:y:9:G:N1 | 38:y:15:G:O6 | 2.39 | 0.55 |
| 1:2:792:G:H2' | 1:2:793:C:O4' | 2.07 | 0.55 |
| 1:2:1743:G:O6 | 1:2:1780:U:O4 | 2.25 | 0.55 |
| 8:G:87:MET:HG3 | 8:G:87:MET:O | 2.06 | 0.55 |
| 8:G:100:ARG:HB2 | 8:G:114:ILE:HD13 | 1.87 | 0.55 |
| 16:O:17:ALA:HA | 16:O:124:ILE:HD12 | 1.88 | 0.55 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 23:V:52:TRP:O | 23:V:56:ARG:HG2 | 2.07 | 0.55 |
| 1:2:624:A:OP1 | 27:Z:134:TYR:OH | 2.21 | 0.55 |
| 21:T:47:ARG:HA | 21:T:50:ILE:HG12 | 1.88 | 0.55 |
| 26:Y:83:LEU:HD12 | 26:Y:117:ARG:HH21 | 1.71 | 0.55 |
| 1:2:1266:G:O6 | 1:2:1507:U:O4 | 2.23 | 0.55 |
| 1:2:1398:A:N6 | 1:2:1437:U:O2' | 2.40 | 0.55 |
| 1:2:1448:A:N6 | 1:2:1469:G:O2' | 2.39 | 0.55 |
| 1:2:1601:G:N2 | 1:2:1627:G:O2' | 2.36 | 0.55 |
| 38:y:25:G:N1 | 38:y:43:A:H2 | 2.02 | 0.55 |
| 1:2:9:U:N3 | 1:2:12:U:OP2 | 2.37 | 0.55 |
| 1:2:98:C:OP2 | 1:2:416:A:O2' | 2.23 | 0.55 |
| 1:2:1188:U:OP2 | 27:Z:119:ARG:NH2 | 2.33 | 0.55 |
| 1:2:1361:G:H2' | 1:2:1362:G:C8 | 2.42 | 0.55 |
| 1:2:1675:G:H4' | 31:d:20:ARG:HD3 | 1.88 | 0.55 |
| 4:C:31:ASP:HB3 | 4:C:34:MET:HB2 | 1.89 | 0.55 |
| 38:y:21:G:H2' | 38:y:22:C:C6 | 2.42 | 0.55 |
| 1:2:370:G:OP1 | 12:K:56:ARG:NH2 | 2.40 | 0.55 |
| 1:2:1043:C:H2' | 1:2:1044:G:C8 | 2.41 | 0.55 |
| 1:2:1294:G:N3 | 19:R:79:HIS:ND1 | 2.54 | 0.55 |
| 1:2:1688:G:H21 | 1:2:1828:A:H8 | 1.54 | 0.55 |
| 1:2:1700:C:H2' | 1:2:1701:G:C8 | 2.41 | 0.55 |
| 17:P:33:VAL:O | 17:P:37:ILE:HG12 | 2.07 | 0.55 |
| 1:2:950:U:H3 | 1:2:967:G:H1' | 1.72 | 0.54 |
| 1:2:1142:C:H2' | 1:2:1143:C:C6 | 2.42 | 0.54 |
| 1:2:1279:C:H4' | 33:f:99:LYS:NZ | 2.21 | 0.54 |
| 1:2:1571:G:H2' | 1:2:1572:G:H8 | 1.72 | 0.54 |
| 1:2:1586:C:O3' | 9:H:88:MET:HE1 | 2.07 | 0.54 |
| 5:D:218:LEU:O | 5:D:219:LYS:HE2 | 2.07 | 0.54 |
| 7:F:22:ASN:O | 7:F:26:THR:HG23 | 2.06 | 0.54 |
| 10:I:116:LYS:NZ | 10:I:125:THR:OG1 | 2.35 | 0.54 |
| 20:S:29:ASN:N | 20:S:67:ASP:OD1 | 2.37 | 0.54 |
| 25:X:34:MET:HE2 | 25:X:69:ILE:HD12 | 1.89 | 0.54 |
| 34:g:238:ALA:H | 34:g:251:ALA:HB3 | 1.72 | 0.54 |
| 1:2:75:G:OP2 | 1:2:75:G:N2 | 2.35 | 0.54 |
| 1:2:492:C:O2 | 8:G:63:LYS:NZ | 2.41 | 0.54 |
| 1:2:673:G:N1 | 1:2:1018:U:OP2 | 2.37 | 0.54 |
| 1:2:678:U:H5 | 1:2:736:C:H42 | 1.55 | 0.54 |
| 1:2:840:U:H2' | 1:2:841:G:C8 | 2.42 | 0.54 |
| 1:2:937:C:H2' | 1:2:938:G:C8 | 2.43 | 0.54 |
| 1:2:949:C:OP2 | 5:D:19:LYS:NZ | 2.39 | 0.54 |
| 1:2:1503:G:C8 | 33:f:88:PRO:HD2 | 2.42 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 17:P:70:LYS:O | 17:P:74:ILE:HD12 | 2.06 | 0.54 |
| 22:U:147:GLY:HA2 | 38:y:27:U:H5'' | 1.90 | 0.54 |
| 26:Y:78:ARG:NH1 | 26:Y:126:LEU:HD13 | 2.23 | 0.54 |
| 27:Z:48:LYS:HE2 | 27:Z:101:LEU:HD21 | 1.89 | 0.54 |
| 28:a:100:LYS:NZ | 28:a:101:LYS:O | 2.36 | 0.54 |
| 1:2:5:U:H2' | 1:2:6:G:C8 | 2.43 | 0.54 |
| 1:2:142:C:N4 | 1:2:319:G:OP2 | 2.40 | 0.54 |
| 1:2:900:A:H2' | 1:2:901:C:C6 | 2.42 | 0.54 |
| 1:2:980:C:O2' | 18:Q:138:ASP:OD2 | 2.24 | 0.54 |
| 6:E:106:ARG:HH22 | 6:E:108:ARG:HH21 | 1.54 | 0.54 |
| 9:H:21:GLY:H | 9:H:48:TYR:HD1 | 1.53 | 0.54 |
| 23:V:11:GLN:OE1 | 23:V:62:ARG:NH1 | 2.40 | 0.54 |
| 24:W:48:LEU:HD23 | 24:W:91:LEU:HD23 | 1.89 | 0.54 |
| 28:a:22:GLN:HB2 | 28:a:72:PHE:HZ | 1.73 | 0.54 |
| 29:b:28:CYS:SG | 29:b:29:CYS:N | 2.80 | 0.54 |
| 33:f:95:ARG:HG3 | 33:f:97:LYS:H | 1.72 | 0.54 |
| 38:y:10:G:C8 | 38:y:45:G:N2 | 2.74 | 0.54 |
| 1:2:853:U:H4' | 8:G:201:HIS:NE2 | 2.22 | 0.54 |
| 4:C:205:ARG:HD2 | 4:C:206:ASP:N | 2.23 | 0.54 |
| 6:E:69:PHE:HD2 | 6:E:71:LEU:HB2 | 1.70 | 0.54 |
| 6:E:249:SER:HB2 | 6:E:252:GLN:HE22 | 1.72 | 0.54 |
| 23:V:13:GLU:OE2 | 23:V:16:ARG:NH1 | 2.40 | 0.54 |
| 27:Z:68:LYS:HZ2 | 36:i:83:VAL:HA | 1.73 | 0.54 |
| 28:a:22:GLN:HB2 | 28:a:72:PHE:CZ | 2.42 | 0.54 |
| 29:b:37:LYS:NZ | 29:b:70:LYS:HG2 | 2.22 | 0.54 |
| 34:g:153:CYS:HB3 | 34:g:198:VAL:HG22 | 1.89 | 0.54 |
| 39:z:570:A:H3' | 39:z:571:A:H8 | 1.71 | 0.54 |
| 1:2:94:G:OP1 | 8:G:8:HIS:ND1 | 2.40 | 0.54 |
| 1:2:921:G:H22 | 1:2:1013:U:H3 | 1.55 | 0.54 |
| 1:2:1173:U:H2' | 1:2:1174:U:C6 | 2.42 | 0.54 |
| 1:2:1663:U:OP2 | 20:S:141:TYR:OH | 2.24 | 0.54 |
| 5:D:23:ASP:O | 5:D:26:SER:OG | 2.24 | 0.54 |
| 5:D:40:ASN:OD1 | 5:D:75:GLN:NE2 | 2.40 | 0.54 |
| 5:D:205:TYR:CD1 | 5:D:206:PRO:HD2 | 2.42 | 0.54 |
| 6:E:241:TRP:CG | 26:Y:68:ARG:HE | 2.24 | 0.54 |
| 1:2:579:G:H5'' | 1:2:580:A:C8 | 2.43 | 0.54 |
| 1:2:1280:A:H4' | 1:2:1281:G:H5'' | 1.89 | 0.54 |
| 5:D:217:MET:HE3 | 5:D:220:LYS:HG2 | 1.88 | 0.54 |
| 6:E:72:PRO:HG3 | 25:X:29:HIS:CG | 2.43 | 0.54 |
| 10:I:64:LYS:HG2 | 10:I:97:VAL:HG21 | 1.89 | 0.54 |
| 38:y:9:G:N2 | 38:y:20:A:O4' | 2.40 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 1:2:73:C:C2 | 1:2:74:G:H1' | 2.43 | 0.54 |
| 1:2:389:C:H5'' | 1:2:390:C:H5 | 1.72 | 0.54 |
| 1:2:632:U:H4' | 1:2:634:G:H4' | 1.90 | 0.54 |
| 4:C:36:GLN:OE1 | 4:C:36:GLN:N | 2.40 | 0.54 |
| 8:G:60:GLU:OE1 | 28:a:20:ARG:NH2 | 2.40 | 0.54 |
| 11:J:159:ASP:OD1 | 11:J:160:LYS:N | 2.40 | 0.54 |
| 35:h:106:GLN:HE22 | 35:h:108:ILE:CD1 | 2.21 | 0.54 |
| 39:z:582:U:N3 | 39:z:585:G:OP2 | 2.30 | 0.54 |
| 1:2:159:A:H2 | 1:2:457:G:H21 | 1.54 | 0.54 |
| 1:2:1167:G:O2' | 1:2:1183:G:O6 | 2.24 | 0.54 |
| 1:2:1376:C:H2' | 1:2:1377:G:C8 | 2.43 | 0.54 |
| 1:2:1426:C:H2' | 1:2:1427:G:C8 | 2.42 | 0.54 |
| 1:2:1643:G:H5'' | 20:S:125:ARG:HB3 | 1.90 | 0.54 |
| 1:2:1818:A:O2' | 1:2:1819:A:H5'' | 2.07 | 0.54 |
| 7:F:19:ALA:HA | 7:F:22:ASN:HD21 | 1.73 | 0.54 |
| 9:H:122:ARG:HH21 | 31:d:9:ILE:HD11 | 1.72 | 0.54 |
| 12:K:8:TRP:CD1 | 12:K:22:HIS:HE2 | 2.26 | 0.54 |
| 17:P:141:TYR:HE2 | 17:P:146:ALA:HB2 | 1.71 | 0.54 |
| 19:R:39:ALA:HA | 19:R:42:ARG:HE | 1.72 | 0.54 |
| 34:g:292:SER:OG | 34:g:297:THR:OG1 | 2.25 | 0.54 |
| 1:2:64:A:N6 | 1:2:83:A:OP2 | 2.36 | 0.54 |
| 1:2:1043:C:H5'' | 18:Q:143:LYS:HD3 | 1.89 | 0.54 |
| 1:2:1563:C:H2' | 1:2:1564:A:C4 | 2.42 | 0.54 |
| 6:E:133:ALA:O | 6:E:137:ARG:HG2 | 2.07 | 0.54 |
| 8:G:195:ILE:HG23 | 8:G:196:THR:H | 1.72 | 0.54 |
| 9:H:89:THR:HA | 9:H:92:ILE:HG12 | 1.89 | 0.54 |
| 10:I:7:PHE:O | 10:I:11:GLY:N | 2.41 | 0.54 |
| 13:L:170:PRO:O | 13:L:175:ARG:NH2 | 2.38 | 0.54 |
| 34:g:296:GLN:HG2 | 34:g:297:THR:HG23 | 1.90 | 0.54 |
| 36:i:107:LYS:O | 36:i:110:MET:HG3 | 2.07 | 0.54 |
| 1:2:165:G:H2' | 1:2:166:A:H8 | 1.73 | 0.54 |
| 1:2:600:G:H2' | 1:2:601:G:C8 | 2.43 | 0.54 |
| 1:2:741:C:H2' | 1:2:742:C:O4' | 2.08 | 0.54 |
| 1:2:898:G:H2' | 1:2:899:A:H8 | 1.72 | 0.54 |
| 1:2:937:C:H2' | 1:2:938:G:H8 | 1.73 | 0.54 |
| 1:2:987:G:C6 | 1:2:1130:G:H4' | 2.43 | 0.54 |
| 1:2:1289:A:H2' | 1:2:1290:G:C8 | 2.42 | 0.54 |
| 3:B:161:GLY:N | 3:B:190:ASN:O | 2.35 | 0.54 |
| 5:D:185:VAL:HA | 5:D:188:LEU:HG | 1.89 | 0.54 |
| 12:K:142:SER:N | 12:K:143:LYS:HB2 | 2.23 | 0.54 |
| 18:Q:142:ARG:HD2 | 18:Q:143:LYS:O | 2.08 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 31:d:30:VAL:HG21 | 31:d:50:VAL:HG21 | 1.89 | 0.54 |
| 36:i:85:LYS:O | 36:i:89:GLN:NE2 | 2.41 | 0.54 |
| 1:2:633:A:OP1 | 13:L:38:ARG:NH2 | 2.41 | 0.53 |
| 1:2:1574:A:O2' | 1:2:1577:C:N4 | 2.40 | 0.53 |
| 1:2:1624:C:H2' | 1:2:1625:A:H8 | 1.72 | 0.53 |
| 5:D:49:VAL:HG11 | 5:D:62:LEU:HB2 | 1.90 | 0.53 |
| 5:D:92:GLN:HG2 | 5:D:97:LEU:HD11 | 1.90 | 0.53 |
| 7:F:66:ILE:HD11 | 7:F:86:LEU:HB3 | 1.90 | 0.53 |
| 1:2:648:U:O3' | 27:Z:17:ARG:NH2 | 2.41 | 0.53 |
| 1:2:1244:U:H2' | 1:2:1245:C:C6 | 2.44 | 0.53 |
| 1:2:1428:U:H2' | 1:2:1429:C:C6 | 2.43 | 0.53 |
| 1:2:1593:G:H3' | 35:h:80:ARG:HD2 | 1.90 | 0.53 |
| 4:C:11:LYS:HZ2 | 4:C:184:ARG:HH12 | 1.55 | 0.53 |
| 7:F:101:GLN:HG3 | 7:F:126:ILE:HD11 | 1.89 | 0.53 |
| 10:I:224:ARG:O | 10:I:227:GLN:HG3 | 2.08 | 0.53 |
| 13:L:50:LEU:HD12 | 13:L:53:ILE:HD11 | 1.89 | 0.53 |
| 22:U:113:ARG:O | 22:U:117:ILE:HG12 | 2.08 | 0.53 |
| 37:l:7:LYS:HE2 | 37:l:11:ARG:HD2 | 1.88 | 0.53 |
| 1:2:649:G:O2' | 1:2:652:G:O2' | 2.23 | 0.53 |
| 1:2:1019:A:H2' | 1:2:1020:A:H8 | 1.73 | 0.53 |
| 1:2:1531:G:H2' | 1:2:1532:A:C8 | 2.43 | 0.53 |
| 1:2:1804:U:H2' | 1:2:1805:C:C6 | 2.43 | 0.53 |
| 5:D:35:ALA:HB2 | 5:D:44:ILE:HD11 | 1.88 | 0.53 |
| 5:D:71:LEU:HB3 | 5:D:78:GLU:OE2 | 2.07 | 0.53 |
| 17:P:56:ASP:OD1 | 17:P:57:SER:N | 2.40 | 0.53 |
| 32:e:19:ARG:NH2 | 32:e:32:ARG:HH21 | 2.07 | 0.53 |
| 1:2:480:C:O2' | 1:2:564:A:N1 | 2.40 | 0.53 |
| 1:2:1108:U:H3 | 1:2:1109:A:N6 | 2.07 | 0.53 |
| 1:2:1163:G:H2' | 1:2:1164:G:O4' | 2.08 | 0.53 |
| 4:C:104:THR:O | 4:C:107:THR:OG1 | 2.27 | 0.53 |
| 9:H:88:MET:N | 9:H:88:MET:HE2 | 2.24 | 0.53 |
| 13:L:81:LEU:HB3 | 13:L:87:LEU:HD23 | 1.91 | 0.53 |
| 13:L:122:SER:H | 13:L:125:HIS:HB3 | 1.73 | 0.53 |
| 28:a:54:VAL:HB | 28:a:76:TYR:H | 1.73 | 0.53 |
| 1:2:1196:A:H2' | 1:2:1197:U:C6 | 2.44 | 0.53 |
| 11:J:99:ARG:HH21 | 11:J:100:ILE:HG12 | 1.73 | 0.53 |
| 12:K:142:SER:HA | 12:K:145:ILE:HG13 | 1.90 | 0.53 |
| 19:R:8:LYS:O | 19:R:14:LYS:HB3 | 2.07 | 0.53 |
| 21:T:99:ASP:OD1 | 21:T:102:THR:OG1 | 2.21 | 0.53 |
| 36:i:120:VAL:O | 36:i:125:LYS:NZ | 2.33 | 0.53 |
| 1:2:411:G:OP1 | 15:N:97:ARG:NH1 | 2.42 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 9:H:110:GLN:OE1 | 9:H:110:GLN:N | 2.35 | 0.53 |
| 10:I:7:PHE:CD1 | 10:I:113:ILE:HG21 | 2.44 | 0.53 |
| 18:Q:95:ILE:HG21 | 18:Q:116:LEU:HD11 | 1.90 | 0.53 |
| 18:Q:148:GLY:O | 18:Q:150:ARG:NH1 | 2.42 | 0.53 |
| 21:T:99:ASP:HB2 | 21:T:100:PRO:HD2 | 1.90 | 0.53 |
| 26:Y:15:ASN:ND2 | 26:Y:72:CYS:SG | 2.82 | 0.53 |
| 1:2:412:U:H2' | 1:2:413:U:C6 | 2.44 | 0.53 |
| 1:2:440:C:OP1 | 8:G:3:ARG:NH2 | 2.41 | 0.53 |
| 1:2:1211:C:N4 | 1:2:1216:A:H61 | 2.07 | 0.53 |
| 1:2:1517:A:O2' | 22:U:143:GLY:O | 2.19 | 0.53 |
| 7:F:37:VAL:HG12 | 7:F:50:ILE:HD12 | 1.90 | 0.53 |
| 19:R:98:ASN:HB3 | 19:R:120:SER:HB2 | 1.91 | 0.53 |
| 24:W:20:ILE:HD13 | 24:W:98:VAL:HG11 | 1.90 | 0.53 |
| 34:g:5:MET:HA | 34:g:5:MET:HE3 | 1.89 | 0.53 |
| 34:g:220:ASP:HB2 | 34:g:227:LEU:HD21 | 1.91 | 0.53 |
| 1:2:36:U:H2' | 1:2:37:C:H6 | 1.73 | 0.53 |
| 1:2:1443:G:H4' | 24:W:33:GLU:OE2 | 2.09 | 0.53 |
| 1:2:1668:U:O2' | 9:H:84:GLY:O | 2.20 | 0.53 |
| 11:J:64:VAL:HG22 | 11:J:96:ALA:HA | 1.91 | 0.53 |
| 13:L:78:LEU:HD13 | 13:L:92:MET:HE1 | 1.91 | 0.53 |
| 24:W:66:ARG:HE | 24:W:77:TRP:CD1 | 2.26 | 0.53 |
| 26:Y:15:ASN:HA | 26:Y:18:GLU:HG2 | 1.90 | 0.53 |
| 35:h:82:SER:HA | 35:h:85:ARG:HE | 1.73 | 0.53 |
| 1:2:291:U:H2' | 1:2:292:A:C8 | 2.44 | 0.53 |
| 1:2:526:A:N6 | 1:2:537:G:H21 | 2.07 | 0.53 |
| 1:2:663:G:H2' | 1:2:664:C:C6 | 2.44 | 0.53 |
| 1:2:1344:G:N2 | 1:2:1377:G:H22 | 2.06 | 0.53 |
| 1:2:1412:C:H2' | 1:2:1413:C:C6 | 2.44 | 0.53 |
| 4:C:69:GLU:HG3 | 6:E:255:THR:HG21 | 1.91 | 0.53 |
| 4:C:121:LEU:HD13 | 4:C:143:PRO:HG2 | 1.90 | 0.53 |
| 20:S:76:GLY:O | 20:S:80:GLN:N | 2.36 | 0.53 |
| 1:2:71:G:H22 | 10:I:170:ARG:HG3 | 1.73 | 0.53 |
| 1:2:914:U:H3' | 17:P:64:ARG:HH22 | 1.73 | 0.53 |
| 1:2:993:A:H2' | 1:2:994:A:C8 | 2.44 | 0.53 |
| 1:2:1006:G:H2' | 1:2:1007:A:H8 | 1.73 | 0.53 |
| 1:2:1284:U:O2 | 1:2:1307:C:N3 | 2.42 | 0.53 |
| 1:2:1728:U:H2' | 1:2:1729:G:O4' | 2.08 | 0.53 |
| 3:B:89:TYR:HA | 3:B:126:LEU:HA | 1.91 | 0.53 |
| 6:E:196:LYS:HE2 | 6:E:196:LYS:HA | 1.89 | 0.53 |
| 35:h:66:LYS:HA | 35:h:111:ARG:HB2 | 1.91 | 0.53 |
| 39:z:831:A:H5'' | 39:z:832:A:C4 | 2.44 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|------------------|--------------------------|-------------------|
| 1:2:295:C:O2 | 12:K:184:ARG:NH2 | 2.30 | 0.52 |
| 1:2:1196:A:H2' | 1:2:1197:U:H6 | 1.73 | 0.52 |
| 1:2:1294:G:H2' | 1:2:1295:A:O4' | 2.08 | 0.52 |
| 13:L:111:GLN:OE1 | 13:L:111:GLN:N | 2.39 | 0.52 |
| 34:g:197:THR:HG21 | 34:g:238:ALA:HA | 1.90 | 0.52 |
| 1:2:1245:C:OP2 | 1:2:1246:A:O2' | 2.23 | 0.52 |
| 1:2:1451:A:H61 | 1:2:1467:C:N4 | 2.06 | 0.52 |
| 10:I:74:ARG:NH1 | 10:I:96:SER:OG | 2.42 | 0.52 |
| 15:N:101:ARG:HH22 | 27:Z:5:ARG:HG3 | 1.73 | 0.52 |
| 39:z:532:G:O6 | 39:z:559:U:N3 | 2.42 | 0.52 |
| 1:2:551:A:OP2 | 13:L:174:LYS:N | 2.39 | 0.52 |
| 1:2:864:G:N7 | 11:J:115:LYS:HG3 | 2.25 | 0.52 |
| 1:2:1199:G:H2' | 1:2:1200:A:H8 | 1.74 | 0.52 |
| 5:D:27:LYS:HA | 5:D:51:ARG:HE | 1.75 | 0.52 |
| 5:D:151:ARG:HD3 | 5:D:152:LYS:H | 1.75 | 0.52 |
| 7:F:196:GLY:H | 7:F:198:ILE:HG13 | 1.75 | 0.52 |
| 8:G:184:THR:HG22 | 8:G:224:ASN:HA | 1.91 | 0.52 |
| 17:P:26:LEU:HD21 | 17:P:66:VAL:HG11 | 1.92 | 0.52 |
| 35:h:106:GLN:NE2 | 35:h:107:VAL:O | 2.42 | 0.52 |
| 38:y:18:G:N2 | 38:y:56:G:O2' | 2.43 | 0.52 |
| 38:y:26:C:H2' | 38:y:27:U:C6 | 2.44 | 0.52 |
| 38:y:34:A:H2' | 38:y:35:U:C6 | 2.45 | 0.52 |
| 1:2:190:A:H3' | 1:2:191:C:H5'' | 1.90 | 0.52 |
| 1:2:946:C:H2' | 1:2:947:C:H6 | 1.74 | 0.52 |
| 1:2:947:C:H2' | 1:2:948:G:C8 | 2.40 | 0.52 |
| 7:F:151:LYS:HG2 | 7:F:152:PHE:N | 2.24 | 0.52 |
| 9:H:153:LEU:HG | 9:H:189:ALA:HA | 1.92 | 0.52 |
| 28:a:6:THR:HB | 28:a:28:LEU:HB2 | 1.91 | 0.52 |
| 1:2:794:G:C5 | 11:J:106:ARG:HA | 2.45 | 0.52 |
| 1:2:868:A:O2' | 1:2:869:G:H8 | 1.93 | 0.52 |
| 1:2:958:A:H2' | 1:2:959:A:O4' | 2.09 | 0.52 |
| 1:2:1588:C:O2' | 20:S:45:ARG:NH2 | 2.42 | 0.52 |
| 1:2:1648:U:H2' | 1:2:1649:G:C8 | 2.45 | 0.52 |
| 6:E:71:LEU:HD12 | 6:E:72:PRO:HD2 | 1.91 | 0.52 |
| 6:E:84:GLY:H | 6:E:87:LEU:HD23 | 1.74 | 0.52 |
| 6:E:233:TYR:OH | 25:X:12:TYR:O | 2.26 | 0.52 |
| 14:M:16:PHE:CE2 | 14:M:89:ILE:HG21 | 2.44 | 0.52 |
| 16:O:77:ILE:HG12 | 16:O:128:PHE:HE2 | 1.74 | 0.52 |
| 38:y:9:G:N7 | 38:y:19:A:O2' | 2.42 | 0.52 |
| 1:2:293:A:H1' | 12:K:73:THR:HG23 | 1.92 | 0.52 |
| 1:2:562:U:H2' | 1:2:563:U:C2 | 2.45 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|------------------|--------------------------|-------------------|
| 1:2:1443:G:H2' | 1:2:1444:A:C8 | 2.45 | 0.52 |
| 1:2:1640:C:H5'' | 20:S:138:ARG:CZ | 2.40 | 0.52 |
| 4:C:17:LYS:HD3 | 4:C:173:LEU:HD11 | 1.91 | 0.52 |
| 17:P:27:LYS:HG3 | 17:P:28:LEU:HG | 1.90 | 0.52 |
| 22:U:143:GLY:O | 22:U:144:ARG:HD3 | 2.10 | 0.52 |
| 1:2:524:G:H2' | 1:2:525:G:C8 | 2.45 | 0.52 |
| 1:2:1044:G:N1 | 1:2:1065:U:OP2 | 2.37 | 0.52 |
| 1:2:1382:A:H2' | 1:2:1383:G:O4' | 2.09 | 0.52 |
| 1:2:1404:U:OP1 | 20:S:71:ARG:NH1 | 2.43 | 0.52 |
| 10:I:7:PHE:CE2 | 10:I:9:ALA:HB3 | 2.45 | 0.52 |
| 10:I:204:GLU:O | 10:I:208:GLU:HG2 | 2.10 | 0.52 |
| 23:V:61:ALA:O | 23:V:65:TYR:N | 2.43 | 0.52 |
| 28:a:106:GLN:H | 28:a:106:GLN:CD | 2.18 | 0.52 |
| 34:g:20:GLN:NE2 | 34:g:69:VAL:O | 2.39 | 0.52 |
| 36:i:86:VAL:HG12 | 36:i:86:VAL:O | 2.10 | 0.52 |
| 1:2:115:U:O2' | 1:2:371:C:O2 | 2.22 | 0.52 |
| 1:2:191:C:H5' | 12:K:143:LYS:HE3 | 1.90 | 0.52 |
| 1:2:858:A:N7 | 26:Y:107:SER:HA | 2.25 | 0.52 |
| 1:2:869:G:O6 | 1:2:910:U:O2 | 2.28 | 0.52 |
| 1:2:1634:G:O2' | 1:2:1635:A:O4' | 2.27 | 0.52 |
| 4:C:11:LYS:NZ | 4:C:192:GLU:OE1 | 2.35 | 0.52 |
| 8:G:35:PRO:HG2 | 8:G:36:HIS:HD1 | 1.75 | 0.52 |
| 35:h:62:VAL:HB | 35:h:63:PRO:HD3 | 1.92 | 0.52 |
| 1:2:185:C:H2' | 1:2:186:G:C8 | 2.44 | 0.52 |
| 1:2:903:G:H2' | 1:2:904:A:H8 | 1.72 | 0.52 |
| 1:2:1013:U:H2' | 1:2:1014:U:H6 | 1.75 | 0.52 |
| 1:2:1391:C:H2' | 1:2:1392:A:N3 | 2.23 | 0.52 |
| 5:D:40:ASN:H | 5:D:75:GLN:HE22 | 1.58 | 0.52 |
| 11:J:194:LEU:HD22 | 30:c:27:SER:HB2 | 1.92 | 0.52 |
| 14:M:65:ARG:NH1 | 32:e:22:ARG:O | 2.40 | 0.52 |
| 19:R:10:ARG:HH12 | 19:R:21:ASP:HA | 1.75 | 0.52 |
| 19:R:15:PHE:O | 22:U:91:LYS:NZ | 2.37 | 0.52 |
| 19:R:81:ARG:HB2 | 19:R:117:GLY:HA3 | 1.91 | 0.52 |
| 1:2:454:A:H3' | 1:2:455:A:H8 | 1.75 | 0.52 |
| 1:2:600:G:H2' | 1:2:601:G:H8 | 1.74 | 0.52 |
| 1:2:924:G:H2' | 1:2:925:G:C8 | 2.45 | 0.52 |
| 1:2:1050:G:O6 | 1:2:1051:A:N6 | 2.42 | 0.52 |
| 1:2:1236:A:C4 | 19:R:100:LYS:HE2 | 2.45 | 0.52 |
| 1:2:1383:G:N1 | 7:F:206:ASP:O | 2.25 | 0.52 |
| 5:D:144:LYS:HA | 5:D:208:HIS:ND1 | 2.25 | 0.52 |
| 5:D:196:ASP:HA | 5:D:199:LYS:HG2 | 1.92 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|------------------|--------------------------|-------------------|
| 6:E:191:SER:HB2 | 6:E:196:LYS:HE3 | 1.90 | 0.52 |
| 7:F:19:ALA:HA | 7:F:22:ASN:ND2 | 2.25 | 0.52 |
| 9:H:144:LEU:O | 9:H:148:ASN:ND2 | 2.43 | 0.52 |
| 9:H:162:ALA:HA | 9:H:165:ASN:HB2 | 1.90 | 0.52 |
| 10:I:18:VAL:HB | 10:I:23:LYS:NZ | 2.24 | 0.52 |
| 1:2:104:A:OP1 | 12:K:12:ARG:NE | 2.43 | 0.51 |
| 1:2:920:G:H1 | 1:2:1014:U:H3 | 1.58 | 0.51 |
| 1:2:1451:A:H2' | 1:2:1452:G:C8 | 2.45 | 0.51 |
| 1:2:1615:A:H2' | 19:R:40:ARG:NH2 | 2.24 | 0.51 |
| 1:2:1659:A:H2' | 1:2:1661:C:H41 | 1.75 | 0.51 |
| 6:E:92:LEU:HB3 | 6:E:112:PHE:HB2 | 1.92 | 0.51 |
| 10:I:97:VAL:HG22 | 10:I:98:ARG:H | 1.75 | 0.51 |
| 12:K:6:ASP:OD1 | 12:K:9:HIS:ND1 | 2.43 | 0.51 |
| 22:U:75:ARG:HH12 | 22:U:80:PRO:HA | 1.74 | 0.51 |
| 34:g:17:TRP:CD1 | 34:g:303:THR:HG1 | 2.28 | 0.51 |
| 1:2:151:C:H2' | 1:2:152:U:H6 | 1.76 | 0.51 |
| 1:2:507:C:H2' | 1:2:508:G:O4' | 2.10 | 0.51 |
| 1:2:932:G:H2' | 1:2:933:C:C6 | 2.45 | 0.51 |
| 1:2:1789:G:H2' | 1:2:1790:G:H8 | 1.75 | 0.51 |
| 10:I:32:MET:SD | 10:I:63:MET:HE1 | 2.50 | 0.51 |
| 20:S:83:ALA:HA | 20:S:86:GLN:HE21 | 1.75 | 0.51 |
| 30:c:37:CYS:SG | 30:c:60:SER:OG | 2.57 | 0.51 |
| 34:g:131:LEU:O | 34:g:139:LYS:N | 2.43 | 0.51 |
| 35:h:80:ARG:HB3 | 35:h:83:LEU:HB2 | 1.92 | 0.51 |
| 1:2:444:U:H2' | 1:2:445:A:C8 | 2.45 | 0.51 |
| 1:2:577:A:H5' | 1:2:582:C:C5 | 2.45 | 0.51 |
| 1:2:1093:G:H2' | 1:2:1094:C:C6 | 2.46 | 0.51 |
| 1:2:1283:A:H5' | 33:f:99:LYS:NZ | 2.25 | 0.51 |
| 1:2:1861:U:P | 29:b:10:ARG:HH22 | 2.33 | 0.51 |
| 4:C:60:LEU:HD11 | 4:C:159:ILE:HG12 | 1.93 | 0.51 |
| 19:R:8:LYS:HA | 19:R:12:PHE:HD2 | 1.75 | 0.51 |
| 19:R:92:SER:O | 19:R:93:MET:HE2 | 2.10 | 0.51 |
| 34:g:122:SER:HG | 34:g:132:TRP:HE1 | 1.58 | 0.51 |
| 1:2:550:A:OP2 | 13:L:177:ASN:ND2 | 2.44 | 0.51 |
| 1:2:917:G:C4 | 30:c:21:LYS:HE2 | 2.45 | 0.51 |
| 1:2:1043:C:O3' | 18:Q:141:ARG:NH1 | 2.42 | 0.51 |
| 1:2:1123:C:H4' | 30:c:17:ARG:HH22 | 1.76 | 0.51 |
| 8:G:202:PRO:HB2 | 15:N:42:LEU:HD13 | 1.92 | 0.51 |
| 10:I:199:THR:HA | 10:I:202:ASN:ND2 | 2.26 | 0.51 |
| 18:Q:61:LYS:HZ1 | 18:Q:80:ASP:HB2 | 1.76 | 0.51 |
| 24:W:40:ILE:O | 24:W:44:LYS:HG2 | 2.09 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 26:Y:81:VAL:HG13 | 26:Y:85:ASP:HB2 | 1.92 | 0.51 |
| 1:2:28:U:H2' | 1:2:29:G:H8 | 1.75 | 0.51 |
| 1:2:606:A:H5'' | 27:Z:68:LYS:NZ | 2.26 | 0.51 |
| 1:2:898:G:H2' | 1:2:899:A:C8 | 2.45 | 0.51 |
| 1:2:1537:C:H2' | 1:2:1538:U:C2 | 2.46 | 0.51 |
| 1:2:1651:G:O6 | 1:2:1663:U:O4 | 2.28 | 0.51 |
| 6:E:236:LEU:H | 6:E:236:LEU:HD23 | 1.76 | 0.51 |
| 7:F:203:PRO:HA | 7:F:208:VAL:HG13 | 1.92 | 0.51 |
| 14:M:43:LEU:H | 14:M:43:LEU:HD12 | 1.74 | 0.51 |
| 28:a:20:ARG:NH1 | 28:a:76:TYR:OH | 2.42 | 0.51 |
| 39:z:579:G:H2' | 39:z:580:U:C6 | 2.45 | 0.51 |
| 1:2:297:C:H2' | 1:2:298:G:C8 | 2.45 | 0.51 |
| 1:2:553:G:C2 | 1:2:554:A:C8 | 2.99 | 0.51 |
| 1:2:1164:G:H2' | 1:2:1165:G:O4' | 2.10 | 0.51 |
| 1:2:1195:A:H2' | 1:2:1196:A:C8 | 2.45 | 0.51 |
| 1:2:1631:G:N7 | 35:h:41:ARG:NH1 | 2.58 | 0.51 |
| 4:C:112:ILE:HG13 | 4:C:112:ILE:O | 2.10 | 0.51 |
| 9:H:28:VAL:O | 9:H:42:LYS:NZ | 2.40 | 0.51 |
| 9:H:122:ARG:HH22 | 31:d:58:LEU:N | 2.09 | 0.51 |
| 13:L:50:LEU:HD11 | 13:L:100:LEU:CD1 | 2.40 | 0.51 |
| 19:R:85:ILE:HG22 | 19:R:112:ILE:HD13 | 1.92 | 0.51 |
| 31:d:12:ALA:HA | 31:d:35:MET:HG2 | 1.92 | 0.51 |
| 31:d:31:ARG:HD2 | 31:d:43:ILE:HD11 | 1.92 | 0.51 |
| 1:2:77:A:O2' | 10:I:176:ILE:N | 2.43 | 0.51 |
| 1:2:435:A:H4' | 12:K:50:GLY:HA2 | 1.93 | 0.51 |
| 1:2:518:A:H2' | 1:2:519:A:C8 | 2.45 | 0.51 |
| 1:2:743:U:O2' | 1:2:744:C:O5' | 2.29 | 0.51 |
| 1:2:1015:C:H2' | 1:2:1016:A:C4 | 2.45 | 0.51 |
| 1:2:1188:U:H2' | 1:2:1189:U:C6 | 2.46 | 0.51 |
| 3:B:367:ILE:HA | 3:B:434:CYS:HA | 1.93 | 0.51 |
| 14:M:47:LYS:O | 14:M:50:GLN:HB3 | 2.11 | 0.51 |
| 15:N:49:GLU:HG2 | 15:N:116:CYS:SG | 2.51 | 0.51 |
| 34:g:122:SER:OG | 34:g:132:TRP:NE1 | 2.38 | 0.51 |
| 1:2:127:C:O2' | 1:2:212:G:OP2 | 2.29 | 0.51 |
| 1:2:1510:G:H2' | 1:2:1511:G:H8 | 1.74 | 0.51 |
| 1:2:1612:G:H1 | 19:R:40:ARG:HH22 | 1.59 | 0.51 |
| 10:I:162:LEU:HD23 | 10:I:170:ARG:H | 1.76 | 0.51 |
| 26:Y:77:PRO:HB2 | 27:Z:7:LEU:HD21 | 1.92 | 0.51 |
| 27:Z:25:LYS:HA | 27:Z:28:LYS:HB3 | 1.91 | 0.51 |
| 1:2:520:U:H2' | 1:2:521:A:C8 | 2.46 | 0.51 |
| 1:2:552:U:H2' | 1:2:553:G:C8 | 2.46 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 1:2:837:G:H2' | 1:2:838:C:C6 | 2.46 | 0.51 |
| 1:2:1099:C:H2' | 1:2:1100:G:H8 | 1.74 | 0.51 |
| 1:2:1668:U:H2' | 1:2:1669:G:O4' | 2.11 | 0.51 |
| 7:F:23:GLU:HB3 | 32:e:46:TYR:OH | 2.11 | 0.51 |
| 12:K:12:ARG:HG2 | 12:K:13:LYS:H | 1.76 | 0.51 |
| 15:N:135:SER:H | 15:N:138:VAL:HG12 | 1.76 | 0.51 |
| 24:W:69:PRO:HA | 32:e:40:ARG:HH12 | 1.75 | 0.51 |
| 39:z:579:G:H2' | 39:z:580:U:H6 | 1.74 | 0.51 |
| 1:2:432:C:H42 | 1:2:439:A:H62 | 0.71 | 0.51 |
| 1:2:978:G:H2' | 1:2:979:A:C8 | 2.46 | 0.51 |
| 1:2:1445:G:H2' | 1:2:1446:G:O4' | 2.11 | 0.51 |
| 7:F:50:ILE:HG22 | 7:F:88:ALA:HA | 1.93 | 0.51 |
| 14:M:36:ALA:HB1 | 14:M:39:ASN:HD21 | 1.76 | 0.51 |
| 1:2:1130:G:H2' | 1:2:1131:C:C6 | 2.45 | 0.50 |
| 1:2:1611:U:HO2' | 1:2:1656:A:HO2' | 1.58 | 0.50 |
| 1:2:1688:G:OP2 | 29:b:89:ARG:NH2 | 2.41 | 0.50 |
| 4:C:74:VAL:HG22 | 4:C:121:LEU:HB2 | 1.93 | 0.50 |
| 21:T:33:ARG:O | 21:T:36:GLU:HG3 | 2.11 | 0.50 |
| 27:Z:59:ALA:N | 27:Z:65:ALA:O | 2.44 | 0.50 |
| 29:b:26:CYS:SG | 29:b:76:SER:OG | 2.69 | 0.50 |
| 33:f:116:ARG:HH12 | 33:f:119:ARG:HA | 1.76 | 0.50 |
| 39:z:589:C:H2' | 39:z:590:A:H8 | 1.75 | 0.50 |
| 1:2:29:G:H2' | 1:2:30:C:C6 | 2.46 | 0.50 |
| 1:2:232:A:H2' | 1:2:233:A:C8 | 2.46 | 0.50 |
| 1:2:837:G:H2' | 1:2:838:C:H6 | 1.76 | 0.50 |
| 1:2:899:A:H2' | 1:2:900:A:C8 | 2.46 | 0.50 |
| 17:P:34:LYS:HD2 | 17:P:74:ILE:HG12 | 1.93 | 0.50 |
| 17:P:61:ALA:HB2 | 30:c:32:PHE:CZ | 2.46 | 0.50 |
| 17:P:84:LEU:HD11 | 17:P:89:TYR:HB2 | 1.92 | 0.50 |
| 19:R:74:GLU:HG2 | 19:R:75:VAL:N | 2.24 | 0.50 |
| 29:b:49:ALA:O | 29:b:53:ILE:HG12 | 2.11 | 0.50 |
| 32:e:34:TYR:HB2 | 32:e:36:LEU:HD23 | 1.93 | 0.50 |
| 38:y:55:C:N3 | 38:y:56:G:N1 | 2.59 | 0.50 |
| 39:z:522:U:H2' | 39:z:523:G:C8 | 2.46 | 0.50 |
| 1:2:943:G:H2' | 1:2:944:C:H6 | 1.76 | 0.50 |
| 1:2:1361:G:H2' | 1:2:1362:G:H8 | 1.75 | 0.50 |
| 1:2:1671:U:C4 | 1:2:1672:U:C4 | 2.99 | 0.50 |
| 4:C:80:ARG:NH1 | 4:C:165:ASN:O | 2.44 | 0.50 |
| 5:D:135:LEU:HD23 | 5:D:217:MET:HA | 1.93 | 0.50 |
| 8:G:249:ALA:HA | 13:L:72:PHE:CE1 | 2.46 | 0.50 |
| 17:P:30:SER:O | 17:P:34:LYS:HG2 | 2.11 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 38:y:36:A:H3' | 38:y:37:A:H8 | 1.76 | 0.50 |
| 1:2:349:U:H5'' | 27:Z:22:TRP:HH2 | 1.77 | 0.50 |
| 1:2:544:A:H2' | 1:2:545:A:C8 | 2.47 | 0.50 |
| 1:2:1281:G:OP1 | 16:O:107:SER:OG | 2.30 | 0.50 |
| 1:2:1286:G:N2 | 1:2:1306:U:H1' | 2.27 | 0.50 |
| 1:2:1424:G:C2 | 1:2:1425:G:C8 | 2.99 | 0.50 |
| 1:2:1631:G:H1' | 9:H:164:ARG:HH12 | 1.76 | 0.50 |
| 1:2:1799:G:H2' | 1:2:1800:A:C8 | 2.47 | 0.50 |
| 1:2:1839:A:H2' | 1:2:1840:G:C8 | 2.47 | 0.50 |
| 6:E:99:LYS:HB3 | 6:E:106:ARG:HD3 | 1.94 | 0.50 |
| 7:F:158:ILE:HD13 | 7:F:205:PRO:HB3 | 1.93 | 0.50 |
| 8:G:59:ASP:O | 8:G:62:LYS:HG3 | 2.11 | 0.50 |
| 8:G:182:MET:N | 8:G:226:PHE:O | 2.45 | 0.50 |
| 9:H:174:ALA:O | 9:H:178:ILE:HG12 | 2.11 | 0.50 |
| 10:I:44:GLU:HG3 | 10:I:119:LYS:HE3 | 1.92 | 0.50 |
| 11:J:126:HIS:HA | 11:J:129:ILE:HG22 | 1.94 | 0.50 |
| 12:K:42:ARG:HE | 12:K:59:ARG:HH21 | 1.58 | 0.50 |
| 22:U:143:GLY:C | 22:U:144:ARG:HD3 | 2.36 | 0.50 |
| 29:b:25:ASN:OD1 | 29:b:26:CYS:N | 2.44 | 0.50 |
| 33:f:118:ARG:HH21 | 33:f:133:ALA:HA | 1.76 | 0.50 |
| 38:y:25:G:C6 | 38:y:43:A:N1 | 2.80 | 0.50 |
| 38:y:43:A:H2' | 38:y:44:G:C6 | 2.46 | 0.50 |
| 1:2:217:U:H2' | 1:2:218:A:C8 | 2.46 | 0.50 |
| 1:2:413:U:H2' | 1:2:414:C:C6 | 2.46 | 0.50 |
| 1:2:623:C:O3' | 36:i:87:ARG:NH2 | 2.45 | 0.50 |
| 1:2:910:U:O2' | 11:J:120:ARG:NH2 | 2.45 | 0.50 |
| 1:2:1429:C:H2' | 1:2:1430:C:C6 | 2.47 | 0.50 |
| 9:H:49:LEU:HD13 | 20:S:47:LEU:HD22 | 1.93 | 0.50 |
| 9:H:100:ILE:HD11 | 9:H:108:PRO:HB3 | 1.92 | 0.50 |
| 17:P:49:GLN:HA | 17:P:52:VAL:HG22 | 1.94 | 0.50 |
| 21:T:20:TYR:CZ | 21:T:38:ILE:HD11 | 2.47 | 0.50 |
| 23:V:129:ARG:HG3 | 23:V:133:ARG:HH21 | 1.76 | 0.50 |
| 1:2:1228:U:H2' | 1:2:1229:G:C8 | 2.46 | 0.50 |
| 1:2:1397:A:H4' | 24:W:52:GLY:HA3 | 1.92 | 0.50 |
| 1:2:1414:C:H4' | 1:2:1415:C:H5' | 1.94 | 0.50 |
| 1:2:1554:C:H2' | 1:2:1555:U:O4' | 2.12 | 0.50 |
| 1:2:1678:C:C5 | 31:d:24:GLN:HG3 | 2.47 | 0.50 |
| 1:2:1750:C:H2' | 1:2:1751:G:H8 | 1.76 | 0.50 |
| 7:F:138:VAL:O | 7:F:149:SER:HA | 2.11 | 0.50 |
| 9:H:34:SER:HA | 31:d:55:VAL:HG13 | 1.94 | 0.50 |
| 15:N:16:ILE:HG13 | 15:N:17:PHE:H | 1.75 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 26:Y:102:ILE:HG12 | 26:Y:113:HIS:HD2 | 1.76 | 0.50 |
| 1:2:577:A:H5' | 1:2:582:C:H5 | 1.77 | 0.50 |
| 1:2:1228:U:H2' | 1:2:1229:G:H8 | 1.75 | 0.50 |
| 1:2:1266:G:N2 | 1:2:1507:U:O2 | 2.36 | 0.50 |
| 5:D:167:LYS:HA | 5:D:170:GLU:CD | 2.36 | 0.50 |
| 13:L:50:LEU:HD11 | 13:L:100:LEU:HD11 | 1.94 | 0.50 |
| 20:S:102:GLU:OE2 | 34:g:58:ALA:N | 2.45 | 0.50 |
| 34:g:10:THR:O | 34:g:12:LYS:NZ | 2.41 | 0.50 |
| 34:g:13:GLY:HA3 | 34:g:43:TRP:CH2 | 2.43 | 0.50 |
| 1:2:337:G:H2' | 1:2:338:A:C8 | 2.47 | 0.50 |
| 1:2:843:A:H2' | 1:2:844:U:O4' | 2.12 | 0.50 |
| 1:2:976:A:H2' | 1:2:977:A:C8 | 2.47 | 0.50 |
| 1:2:1093:G:H4' | 4:C:32:PHE:CG | 2.46 | 0.50 |
| 1:2:1394:G:H2' | 1:2:1395:C:H6 | 1.77 | 0.50 |
| 1:2:1448:A:N1 | 1:2:1470:A:O2' | 2.44 | 0.50 |
| 11:J:142:LYS:O | 11:J:143:ARG:HD2 | 2.12 | 0.50 |
| 13:L:63:LEU:HB2 | 13:L:70:ARG:HH21 | 1.76 | 0.50 |
| 29:b:37:LYS:HZ1 | 29:b:70:LYS:HG2 | 1.76 | 0.50 |
| 1:2:30:C:H5'' | 27:Z:138:LYS:NZ | 2.27 | 0.50 |
| 1:2:996:C:OP2 | 29:b:17:HIS:NE2 | 2.45 | 0.50 |
| 5:D:145:LYS:NZ | 5:D:149:GLN:O | 2.39 | 0.50 |
| 8:G:163:ASP:OD1 | 8:G:164:LEU:N | 2.45 | 0.50 |
| 12:K:101:ILE:HG22 | 12:K:172:LEU:HD12 | 1.94 | 0.50 |
| 17:P:63:VAL:HG12 | 17:P:67:THR:OG1 | 2.12 | 0.50 |
| 29:b:9:GLY:HA3 | 29:b:12:LYS:HD2 | 1.94 | 0.50 |
| 35:h:90:GLU:O | 35:h:93:SER:OG | 2.25 | 0.50 |
| 38:y:11:C:N3 | 38:y:24:U:C2 | 2.79 | 0.50 |
| 39:z:523:G:H2' | 39:z:524:C:H6 | 1.76 | 0.50 |
| 1:2:146:G:O2' | 1:2:147:A:H5'' | 2.12 | 0.49 |
| 1:2:411:G:O3' | 15:N:98:LYS:NZ | 2.43 | 0.49 |
| 1:2:1244:U:C2 | 38:y:33:C:H5' | 2.47 | 0.49 |
| 6:E:122:VAL:HG11 | 6:E:229:ILE:HD12 | 1.94 | 0.49 |
| 14:M:49:MET:HE3 | 14:M:69:TRP:CG | 2.46 | 0.49 |
| 15:N:47:PRO:HG3 | 15:N:117:PHE:HE1 | 1.77 | 0.49 |
| 19:R:7:LYS:O | 19:R:9:LYS:HG2 | 2.11 | 0.49 |
| 21:T:71:ILE:HG23 | 21:T:74:GLN:H | 1.76 | 0.49 |
| 31:d:16:LYS:O | 31:d:31:ARG:N | 2.45 | 0.49 |
| 39:z:575:A:C8 | 39:z:576:C:H4' | 2.47 | 0.49 |
| 1:2:656:U:H2' | 1:2:657:U:C6 | 2.47 | 0.49 |
| 1:2:946:C:H2' | 1:2:947:C:C6 | 2.46 | 0.49 |
| 1:2:977:A:H2' | 1:2:978:G:C8 | 2.48 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|------------------|--------------------------|-------------------|
| 1:2:1541:G:H22 | 1:2:1650:C:H1' | 1.76 | 0.49 |
| 4:C:9:GLN:HA | 4:C:55:TRP:HH2 | 1.77 | 0.49 |
| 8:G:136:ILE:HG12 | 8:G:149:TYR:CZ | 2.48 | 0.49 |
| 11:J:26:ALA:HA | 11:J:29:GLU:CD | 2.37 | 0.49 |
| 21:T:2:GLY:O | 21:T:3:ARG:HG3 | 2.12 | 0.49 |
| 31:d:13:ARG:HG2 | 31:d:35:MET:SD | 2.52 | 0.49 |
| 38:y:46:U:H4' | 38:y:47:C:C5 | 2.46 | 0.49 |
| 38:y:50:U:H2' | 38:y:51:G:C8 | 2.47 | 0.49 |
| 1:2:1419:C:O2' | 1:2:1420:G:O5' | 2.27 | 0.49 |
| 1:2:1561:G:O3' | 1:2:1562:G:N2 | 2.40 | 0.49 |
| 1:2:1583:A:H2 | 1:2:1649:G:H1' | 1.77 | 0.49 |
| 5:D:62:LEU:O | 5:D:65:ARG:HG3 | 2.13 | 0.49 |
| 5:D:82:ARG:NH1 | 5:D:191:ASP:OD2 | 2.44 | 0.49 |
| 9:H:178:ILE:HG22 | 9:H:182:LYS:HZ2 | 1.78 | 0.49 |
| 12:K:194:GLU:OE1 | 12:K:194:GLU:N | 2.43 | 0.49 |
| 26:Y:78:ARG:HB3 | 26:Y:124:LYS:HD2 | 1.93 | 0.49 |
| 35:h:85:ARG:HA | 35:h:88:LEU:HG | 1.94 | 0.49 |
| 1:2:354:A:H2' | 1:2:355:C:C6 | 2.47 | 0.49 |
| 1:2:431:C:H4' | 1:2:1733:C:H5' | 1.95 | 0.49 |
| 1:2:671:U:H2' | 1:2:672:U:C6 | 2.47 | 0.49 |
| 1:2:1332:C:H2' | 1:2:1333:C:O4' | 2.12 | 0.49 |
| 14:M:11:ILE:O | 14:M:15:LEU:HG | 2.13 | 0.49 |
| 17:P:113:PHE:O | 17:P:117:LEU:HG | 2.11 | 0.49 |
| 18:Q:142:ARG:NH1 | 18:Q:143:LYS:O | 2.30 | 0.49 |
| 19:R:106:GLU:OE2 | 19:R:107:ILE:N | 2.46 | 0.49 |
| 39:z:580:U:H2' | 39:z:581:A:C8 | 2.48 | 0.49 |
| 1:2:648:U:O2 | 27:Z:17:ARG:NH1 | 2.45 | 0.49 |
| 1:2:951:A:H4' | 18:Q:60:MET:HE1 | 1.95 | 0.49 |
| 1:2:1130:G:OP1 | 29:b:6:ARG:NH1 | 2.46 | 0.49 |
| 1:2:1373:U:H3' | 4:C:102:ARG:HH21 | 1.76 | 0.49 |
| 1:2:1413:C:H2' | 1:2:1415:C:C6 | 2.48 | 0.49 |
| 1:2:1567:C:H2' | 1:2:1568:G:C8 | 2.47 | 0.49 |
| 1:2:1596:A:O5' | 35:h:41:ARG:NH2 | 2.45 | 0.49 |
| 4:C:103:PHE:O | 4:C:107:THR:OG1 | 2.29 | 0.49 |
| 7:F:36:GLY:C | 7:F:51:LEU:HB3 | 2.38 | 0.49 |
| 10:I:64:LYS:HE2 | 10:I:82:SER:HB3 | 1.94 | 0.49 |
| 26:Y:124:LYS:HD3 | 26:Y:125:ILE:H | 1.77 | 0.49 |
| 34:g:34:ALA:HA | 34:g:40:ILE:HG22 | 1.94 | 0.49 |
| 1:2:91:A:OP1 | 1:2:436:G:N2 | 2.29 | 0.49 |
| 1:2:144:U:H2' | 1:2:145:G:H8 | 1.78 | 0.49 |
| 1:2:379:A:H2' | 1:2:380:C:C6 | 2.48 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 1:2:944:C:H2' | 1:2:945:G:C8 | 2.48 | 0.49 |
| 1:2:1587:C:OP1 | 9:H:91:ARG:NH2 | 2.45 | 0.49 |
| 4:C:70:ASN:ND2 | 6:E:259:VAL:HG21 | 2.26 | 0.49 |
| 5:D:36:PRO:HB2 | 5:D:38:MET:HE1 | 1.94 | 0.49 |
| 9:H:125:SER:O | 31:d:47:LYS:NZ | 2.45 | 0.49 |
| 12:K:144:LYS:H | 12:K:147:LYS:HE2 | 1.78 | 0.49 |
| 13:L:85:GLY:HA3 | 13:L:151:LEU:HD13 | 1.94 | 0.49 |
| 17:P:55:ARG:HA | 17:P:60:VAL:O | 2.13 | 0.49 |
| 19:R:39:ALA:HA | 19:R:42:ARG:NE | 2.28 | 0.49 |
| 24:W:67:LYS:HA | 32:e:44:ARG:NH2 | 2.28 | 0.49 |
| 24:W:67:LYS:HD2 | 32:e:44:ARG:CZ | 2.42 | 0.49 |
| 1:2:1:U:O5' | 13:L:54:ARG:NH1 | 2.46 | 0.49 |
| 1:2:76:U:OP2 | 10:I:159:ARG:NH1 | 2.46 | 0.49 |
| 1:2:318:U:C2 | 1:2:319:G:C8 | 3.00 | 0.49 |
| 1:2:858:A:C8 | 26:Y:107:SER:HA | 2.47 | 0.49 |
| 1:2:1271:G:N2 | 1:2:1502:A:OP2 | 2.45 | 0.49 |
| 1:2:1858:U:H3' | 29:b:5:ARG:NH2 | 2.27 | 0.49 |
| 13:L:93:LYS:HB2 | 13:L:96:TYR:CD2 | 2.48 | 0.49 |
| 13:L:150:ARG:O | 13:L:154:GLN:HG3 | 2.13 | 0.49 |
| 14:M:64:TRP:NE1 | 32:e:23:VAL:HA | 2.28 | 0.49 |
| 14:M:73:ASN:HA | 14:M:76:ILE:HD12 | 1.93 | 0.49 |
| 17:P:23:PRO:HD2 | 17:P:26:LEU:HD11 | 1.94 | 0.49 |
| 17:P:88:LEU:O | 17:P:92:ILE:HG12 | 2.12 | 0.49 |
| 28:a:18:LEU:HD13 | 28:a:20:ARG:NH2 | 2.28 | 0.49 |
| 35:h:68:ILE:HB | 35:h:109:TYR:HB2 | 1.95 | 0.49 |
| 38:y:27:U:H2' | 38:y:28:G:C8 | 2.48 | 0.49 |
| 1:2:118:C:H1' | 1:2:435:A:C4 | 2.47 | 0.49 |
| 1:2:191:C:H4' | 1:2:191:C:OP1 | 2.12 | 0.49 |
| 1:2:797:U:C2 | 1:2:798:A:C8 | 3.01 | 0.49 |
| 1:2:1084:U:H4' | 1:2:1085:G:OP2 | 2.12 | 0.49 |
| 1:2:1274:A:C4 | 1:2:1316:G:N2 | 2.81 | 0.49 |
| 1:2:1629:A:H2' | 1:2:1630:C:O4' | 2.13 | 0.49 |
| 4:C:158:ASP:OD1 | 25:X:32:ILE:HD11 | 2.13 | 0.49 |
| 4:C:176:TRP:CZ2 | 4:C:199:PRO:HD3 | 2.48 | 0.49 |
| 11:J:168:HIS:CE1 | 11:J:169:LYS:HE3 | 2.48 | 0.49 |
| 20:S:146:ARG:NE | 38:y:31:C:OP2 | 2.35 | 0.49 |
| 1:2:149:A:N7 | 1:2:169:U:O4 | 2.46 | 0.49 |
| 1:2:296:U:H5'' | 1:2:297:C:C2 | 2.48 | 0.49 |
| 1:2:551:A:H2' | 1:2:552:U:C6 | 2.48 | 0.49 |
| 1:2:1453:U:H2' | 1:2:1454:G:C8 | 2.48 | 0.49 |
| 4:C:39:TYR:OH | 4:C:50:ASN:ND2 | 2.46 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|------------------|--------------------------|-------------------|
| 6:E:191:SER:OG | 6:E:196:LYS:HE3 | 2.13 | 0.49 |
| 8:G:71:LYS:HG3 | 8:G:76:VAL:HG22 | 1.95 | 0.49 |
| 8:G:87:MET:HE3 | 8:G:123:LEU:HB2 | 1.95 | 0.49 |
| 9:H:126:THR:HB | 31:d:47:LYS:HE2 | 1.94 | 0.49 |
| 12:K:120:PRO:HB2 | 12:K:123:ARG:HE | 1.77 | 0.49 |
| 16:O:119:GLN:HG2 | 16:O:121:LYS:NZ | 2.27 | 0.49 |
| 17:P:46:THR:O | 17:P:49:GLN:N | 2.45 | 0.49 |
| 27:Z:121:LYS:NZ | 27:Z:122:VAL:O | 2.45 | 0.49 |
| 28:a:38:THR:HA | 28:a:41:GLN:HG2 | 1.95 | 0.49 |
| 32:e:26:ASN:ND2 | 32:e:27:ARG:O | 2.46 | 0.49 |
| 34:g:8:ARG:HH11 | 34:g:311:GLN:HB3 | 1.77 | 0.49 |
| 34:g:256:ILE:HB | 34:g:270:LEU:HB2 | 1.94 | 0.49 |
| 38:y:48:G:H2' | 38:y:49:A:C8 | 2.48 | 0.49 |
| 1:2:163:U:OP1 | 10:I:85:ARG:N | 2.45 | 0.49 |
| 1:2:1138:G:OP1 | 6:E:172:ARG:NH1 | 2.40 | 0.49 |
| 1:2:1301:C:H2' | 1:2:1302:U:C6 | 2.48 | 0.49 |
| 1:2:1562:G:H2' | 1:2:1563:C:C6 | 2.47 | 0.49 |
| 6:E:57:ASP:OD2 | 6:E:257:HIS:ND1 | 2.46 | 0.49 |
| 11:J:23:ILE:HG13 | 11:J:87:PHE:CE2 | 2.44 | 0.49 |
| 11:J:160:LYS:HA | 11:J:189:PHE:HZ | 1.77 | 0.49 |
| 12:K:197:PHE:CZ | 12:K:201:LYS:HD2 | 2.48 | 0.49 |
| 28:a:24:VAL:HA | 28:a:72:PHE:HA | 1.94 | 0.49 |
| 34:g:247:TRP:NE1 | 34:g:260:ASP:OD1 | 2.42 | 0.49 |
| 39:z:527:G:H2' | 39:z:528:C:C6 | 2.48 | 0.49 |
| 1:2:17:C:H2' | 1:2:18:C:C6 | 2.48 | 0.48 |
| 1:2:55:U:OP1 | 1:2:441:G:N1 | 2.46 | 0.48 |
| 1:2:370:G:N1 | 1:2:373:G:OP2 | 2.38 | 0.48 |
| 1:2:859:U:C2 | 1:2:860:A:C8 | 3.01 | 0.48 |
| 5:D:142:PHE:HB2 | 5:D:209:ASP:HB2 | 1.95 | 0.48 |
| 8:G:6:LYS:O | 8:G:30:ARG:NH2 | 2.46 | 0.48 |
| 22:U:66:ARG:O | 22:U:70:ILE:HG12 | 2.12 | 0.48 |
| 27:Z:95:GLU:OE1 | 27:Z:95:GLU:N | 2.46 | 0.48 |
| 39:z:534:A:O2' | 39:z:535:A:O5' | 2.30 | 0.48 |
| 39:z:574:C:O2' | 39:z:575:A:OP1 | 2.25 | 0.48 |
| 39:z:596:A:H2' | 39:z:597:A:H8 | 1.78 | 0.48 |
| 1:2:191:C:H2' | 1:2:192:U:H6 | 1.77 | 0.48 |
| 1:2:399:C:H2' | 1:2:400:G:C8 | 2.48 | 0.48 |
| 1:2:528:U:H2' | 1:2:529:C:C6 | 2.48 | 0.48 |
| 1:2:1125:G:H2' | 1:2:1126:G:N9 | 2.28 | 0.48 |
| 1:2:1351:C:O2' | 1:2:1352:G:H5' | 2.13 | 0.48 |
| 1:2:1660:G:C4 | 23:V:88:MET:HE1 | 2.47 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 1:2:1710:A:N1 | 1:2:1812:A:H2 | 2.09 | 0.48 |
| 5:D:111:CYS:HA | 5:D:114:VAL:HG12 | 1.96 | 0.48 |
| 5:D:133:TYR:CG | 5:D:217:MET:HE1 | 2.48 | 0.48 |
| 8:G:34:GLY:HA3 | 8:G:83:PRO:HG3 | 1.96 | 0.48 |
| 12:K:40:PRO:O | 12:K:59:ARG:NH1 | 2.46 | 0.48 |
| 12:K:42:ARG:NE | 12:K:59:ARG:HH21 | 2.11 | 0.48 |
| 12:K:110:ARG:HG2 | 12:K:121:LEU:HG | 1.95 | 0.48 |
| 13:L:36:GLY:HA3 | 13:L:123:ILE:HD12 | 1.94 | 0.48 |
| 31:d:27:CYS:SG | 31:d:28:THR:N | 2.86 | 0.48 |
| 1:2:110:U:O2' | 15:N:71:ARG:NH1 | 2.46 | 0.48 |
| 1:2:307:G:H3' | 10:I:183:ARG:HH22 | 1.77 | 0.48 |
| 1:2:409:G:H2' | 1:2:410:G:C8 | 2.48 | 0.48 |
| 1:2:619:A:O2' | 1:2:621:U:OP1 | 2.30 | 0.48 |
| 1:2:1151:U:O2' | 1:2:1152:U:H5' | 2.14 | 0.48 |
| 1:2:1169:A:H62 | 1:2:1183:G:H21 | 1.60 | 0.48 |
| 1:2:1681:G:H2' | 1:2:1682:C:H6 | 1.78 | 0.48 |
| 1:2:1854:A:H3' | 29:b:8:ASN:HD21 | 1.77 | 0.48 |
| 8:G:160:ILE:HD12 | 8:G:169:ILE:HG21 | 1.95 | 0.48 |
| 12:K:40:PRO:HG2 | 12:K:59:ARG:HH12 | 1.79 | 0.48 |
| 26:Y:35:VAL:O | 26:Y:39:THR:HG23 | 2.13 | 0.48 |
| 28:a:35:VAL:HG21 | 28:a:40:ILE:HD11 | 1.94 | 0.48 |
| 30:c:33:MET:HE1 | 30:c:48:SER:HA | 1.95 | 0.48 |
| 34:g:57:ARG:HB3 | 34:g:59:LEU:HD22 | 1.95 | 0.48 |
| 39:z:597:A:H2' | 39:z:598:A:H8 | 1.77 | 0.48 |
| 1:2:1412:C:C4' | 23:V:128:GLN:HE22 | 2.26 | 0.48 |
| 5:D:193:ILE:H | 5:D:193:ILE:HD12 | 1.77 | 0.48 |
| 6:E:155:TRP:HZ2 | 26:Y:97:ARG:HH21 | 1.60 | 0.48 |
| 7:F:214:LYS:HD3 | 21:T:19:LYS:HB3 | 1.94 | 0.48 |
| 8:G:137:PRO:HB2 | 8:G:150:PRO:HD2 | 1.94 | 0.48 |
| 9:H:194:ASP:OD1 | 9:H:195:GLU:N | 2.46 | 0.48 |
| 13:L:92:MET:HA | 13:L:92:MET:HE2 | 1.94 | 0.48 |
| 13:L:134:HIS:CE1 | 13:L:164:PRO:HD2 | 2.39 | 0.48 |
| 1:2:559:A:H2' | 1:2:560:C:H6 | 1.78 | 0.48 |
| 1:2:831:C:O3' | 1:2:832:G:N2 | 2.46 | 0.48 |
| 1:2:865:A:C5 | 1:2:911:G:H1' | 2.47 | 0.48 |
| 1:2:1195:A:H5'' | 29:b:2:THR:HG22 | 1.96 | 0.48 |
| 1:2:1290:G:H2' | 1:2:1291:A:C8 | 2.48 | 0.48 |
| 6:E:63:LEU:HG | 6:E:82:PHE:HD2 | 1.78 | 0.48 |
| 9:H:124:ASP:OD1 | 9:H:125:SER:N | 2.46 | 0.48 |
| 11:J:75:ILE:O | 11:J:77:VAL:HG22 | 2.13 | 0.48 |
| 12:K:125:LYS:HG3 | 12:K:128:LYS:H | 1.79 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 19:R:11:THR:HA | 19:R:16:THR:HG22 | 1.96 | 0.48 |
| 31:d:44:ARG:NH1 | 31:d:63:ARG:O | 2.45 | 0.48 |
| 33:f:135:HIS:HB2 | 33:f:138:ARG:HB2 | 1.95 | 0.48 |
| 1:2:290:A:H2' | 1:2:291:U:H6 | 1.79 | 0.48 |
| 1:2:1054:A:H2' | 1:2:1055:G:O4' | 2.14 | 0.48 |
| 1:2:1803:A:H2' | 1:2:1804:U:C6 | 2.49 | 0.48 |
| 5:D:87:ILE:HG23 | 5:D:101:HIS:HB2 | 1.95 | 0.48 |
| 9:H:135:ARG:HG2 | 18:Q:66:ARG:HH22 | 1.78 | 0.48 |
| 34:g:251:ALA:HB2 | 34:g:289:LEU:HD23 | 1.96 | 0.48 |
| 35:h:97:ILE:HG21 | 35:h:109:TYR:HB3 | 1.96 | 0.48 |
| 38:y:67:C:H2' | 38:y:68:U:C6 | 2.48 | 0.48 |
| 1:2:336:C:H5'' | 8:G:38:LEU:HB2 | 1.96 | 0.48 |
| 1:2:642:U:H2' | 1:2:643:A:C8 | 2.48 | 0.48 |
| 1:2:1135:C:H2' | 1:2:1136:G:O4' | 2.14 | 0.48 |
| 1:2:1160:G:H2' | 1:2:1161:G:C2 | 2.49 | 0.48 |
| 5:D:222:LYS:HG2 | 5:D:223:PHE:H | 1.78 | 0.48 |
| 6:E:133:ALA:O | 6:E:136:ILE:HG22 | 2.13 | 0.48 |
| 20:S:105:LYS:HA | 20:S:108:ILE:HG12 | 1.96 | 0.48 |
| 23:V:37:VAL:HG12 | 23:V:39:LEU:H | 1.79 | 0.48 |
| 28:a:124:ASN:OD1 | 28:a:125:VAL:N | 2.46 | 0.48 |
| 29:b:60:ASP:OD2 | 29:b:60:ASP:N | 2.45 | 0.48 |
| 31:d:17:VAL:HG23 | 31:d:30:VAL:HG22 | 1.96 | 0.48 |
| 34:g:88:ARG:HD3 | 34:g:97:THR:HG21 | 1.95 | 0.48 |
| 1:2:144:U:H2' | 1:2:145:G:C8 | 2.48 | 0.48 |
| 1:2:317:G:H1' | 1:2:318:U:C5 | 2.49 | 0.48 |
| 1:2:510:A:H2' | 1:2:511:A:H8 | 1.77 | 0.48 |
| 1:2:560:C:H2' | 1:2:561:U:C6 | 2.49 | 0.48 |
| 1:2:1139:A:H4' | 1:2:1351:C:H42 | 1.79 | 0.48 |
| 1:2:1582:G:OP1 | 1:2:1582:G:N2 | 2.47 | 0.48 |
| 4:C:51:LEU:HD23 | 4:C:51:LEU:H | 1.78 | 0.48 |
| 16:O:33:ARG:H | 16:O:33:ARG:HD3 | 1.79 | 0.48 |
| 17:P:24:THR:O | 17:P:24:THR:HG22 | 2.14 | 0.48 |
| 17:P:64:ARG:HD2 | 17:P:70:LYS:HE3 | 1.94 | 0.48 |
| 1:2:17:C:H2' | 1:2:18:C:H6 | 1.79 | 0.48 |
| 1:2:376:C:H1' | 12:K:5:ARG:HG3 | 1.95 | 0.48 |
| 1:2:413:U:H2' | 1:2:414:C:H6 | 1.78 | 0.48 |
| 1:2:434:G:N1 | 12:K:26:LYS:HE3 | 2.24 | 0.48 |
| 1:2:630:A:H2' | 1:2:631:A:C8 | 2.49 | 0.48 |
| 1:2:988:A:H2' | 1:2:989:G:H8 | 1.79 | 0.48 |
| 1:2:1030:A:H2' | 1:2:1031:A:O4' | 2.13 | 0.48 |
| 1:2:1717:G:H2' | 1:2:1718:G:H8 | 1.77 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 1:2:1834:U:H2' | 1:2:1835:C:H6 | 1.79 | 0.48 |
| 27:Z:93:PHE:CD2 | 27:Z:133:LEU:HD13 | 2.48 | 0.48 |
| 30:c:14:GLU:HA | 30:c:17:ARG:HB3 | 1.95 | 0.48 |
| 39:z:596:A:H2' | 39:z:597:A:C8 | 2.49 | 0.48 |
| 1:2:835:C:N4 | 28:a:9:THR:O | 2.47 | 0.48 |
| 1:2:909:A:N6 | 11:J:120:ARG:HA | 2.29 | 0.48 |
| 1:2:995:G:H3' | 29:b:17:HIS:HE1 | 1.78 | 0.48 |
| 1:2:1310:U:H4' | 14:M:8:ARG:NH2 | 2.29 | 0.48 |
| 1:2:1382:A:N6 | 7:F:206:ASP:OD2 | 2.42 | 0.48 |
| 1:2:1469:G:N2 | 1:2:1471:G:H5'' | 2.29 | 0.48 |
| 1:2:1789:G:H2' | 1:2:1790:G:C8 | 2.48 | 0.48 |
| 6:E:170:THR:O | 6:E:231:LYS:NZ | 2.40 | 0.48 |
| 6:E:255:THR:HA | 6:E:258:LEU:HD12 | 1.96 | 0.48 |
| 19:R:25:LEU:HA | 19:R:28:MET:SD | 2.54 | 0.48 |
| 20:S:24:HIS:ND1 | 20:S:24:HIS:O | 2.46 | 0.48 |
| 26:Y:46:TYR:HA | 26:Y:68:ARG:HG2 | 1.95 | 0.48 |
| 32:e:23:VAL:HG21 | 32:e:42:CYS:HB2 | 1.95 | 0.48 |
| 32:e:52:PHE:CZ | 32:e:54:LYS:HA | 2.48 | 0.48 |
| 39:z:541:C:H2' | 39:z:542:A:C8 | 2.48 | 0.48 |
| 1:2:867:U:OP2 | 17:P:76:LYS:NZ | 2.29 | 0.47 |
| 1:2:873:C:O2 | 1:2:873:C:H2' | 2.14 | 0.47 |
| 5:D:81:PHE:CD1 | 5:D:109:LYS:HE2 | 2.48 | 0.47 |
| 5:D:146:CYS:SG | 5:D:147:ASN:ND2 | 2.87 | 0.47 |
| 8:G:62:LYS:HA | 8:G:65:CYS:SG | 2.54 | 0.47 |
| 14:M:3:MET:SD | 14:M:8:ARG:HG3 | 2.54 | 0.47 |
| 14:M:13:GLU:HA | 14:M:16:PHE:HB3 | 1.95 | 0.47 |
| 18:Q:29:GLY:O | 18:Q:94:HIS:N | 2.23 | 0.47 |
| 18:Q:37:PHE:HE1 | 18:Q:110:PRO:HD3 | 1.78 | 0.47 |
| 23:V:6:VAL:HA | 23:V:9:VAL:HG22 | 1.95 | 0.47 |
| 38:y:10:G:P | 38:y:45:G:H21 | 2.36 | 0.47 |
| 1:2:473:C:H5'' | 27:Z:48:LYS:HE3 | 1.96 | 0.47 |
| 1:2:742:C:H2' | 1:2:743:U:C6 | 2.49 | 0.47 |
| 1:2:1370:C:H2' | 1:2:1371:G:O4' | 2.14 | 0.47 |
| 6:E:158:LYS:O | 25:X:4:ASN:ND2 | 2.48 | 0.47 |
| 10:I:7:PHE:HD1 | 10:I:113:ILE:HG21 | 1.79 | 0.47 |
| 22:U:52:LEU:HD23 | 22:U:52:LEU:H | 1.79 | 0.47 |
| 24:W:20:ILE:HG13 | 24:W:115:THR:O | 2.13 | 0.47 |
| 38:y:12:G:C6 | 38:y:23:G:C6 | 3.02 | 0.47 |
| 38:y:71:U:H3' | 38:y:73:C:OP2 | 2.14 | 0.47 |
| 39:z:523:G:H2' | 39:z:524:C:C6 | 2.48 | 0.47 |
| 1:2:59:U:H5' | 1:2:493:C:N4 | 2.26 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 1:2:517:C:H2' | 1:2:518:A:H8 | 1.78 | 0.47 |
| 1:2:663:G:H2' | 1:2:664:C:H6 | 1.79 | 0.47 |
| 1:2:1528:A:O2' | 9:H:81:ARG:NH1 | 2.48 | 0.47 |
| 1:2:1692:A:H2 | 6:E:100:GLN:NE2 | 2.12 | 0.47 |
| 5:D:37:ALA:HA | 5:D:42:ARG:HE | 1.79 | 0.47 |
| 7:F:53:THR:O | 7:F:90:LYS:HG3 | 2.14 | 0.47 |
| 8:G:246:LEU:HD21 | 8:G:250:GLU:HG3 | 1.97 | 0.47 |
| 10:I:182:PRO:HA | 10:I:185:LEU:HD12 | 1.96 | 0.47 |
| 11:J:138:GLU:HG2 | 17:P:19:ARG:NH1 | 2.27 | 0.47 |
| 19:R:28:MET:HB2 | 19:R:32:GLN:HE21 | 1.79 | 0.47 |
| 22:U:25:LYS:HD2 | 22:U:55:ARG:HD3 | 1.95 | 0.47 |
| 25:X:38:GLU:N | 25:X:38:GLU:OE1 | 2.47 | 0.47 |
| 27:Z:134:TYR:OH | 36:i:87:ARG:NH2 | 2.46 | 0.47 |
| 34:g:133:ASN:OD1 | 34:g:139:LYS:HE2 | 2.14 | 0.47 |
| 38:y:68:U:H2' | 38:y:69:G:C8 | 2.49 | 0.47 |
| 39:z:521:U:H2' | 39:z:522:U:C6 | 2.49 | 0.47 |
| 1:2:12:U:H2' | 1:2:13:C:C6 | 2.50 | 0.47 |
| 1:2:96:C:H2' | 1:2:97:U:H6 | 1.79 | 0.47 |
| 1:2:145:G:H2' | 1:2:146:G:C8 | 2.49 | 0.47 |
| 1:2:744:C:H2' | 1:2:745:U:C2 | 2.49 | 0.47 |
| 1:2:871:A:H2' | 1:2:872:C:C6 | 2.48 | 0.47 |
| 1:2:943:G:H2' | 1:2:944:C:C6 | 2.49 | 0.47 |
| 1:2:957:G:N2 | 39:z:831:A:N1 | 2.61 | 0.47 |
| 1:2:1028:C:H5'' | 17:P:109:LYS:HZ2 | 1.80 | 0.47 |
| 1:2:1216:A:H2' | 1:2:1217:G:O4' | 2.14 | 0.47 |
| 1:2:1570:G:H2' | 1:2:1571:G:H8 | 1.79 | 0.47 |
| 1:2:1672:U:P | 9:H:71:ARG:HH21 | 2.37 | 0.47 |
| 9:H:45:TYR:HA | 9:H:47:LYS:HZ1 | 1.80 | 0.47 |
| 13:L:60:LEU:O | 13:L:70:ARG:NH2 | 2.46 | 0.47 |
| 13:L:152:ASP:O | 13:L:155:LYS:HG2 | 2.13 | 0.47 |
| 15:N:111:VAL:HG22 | 15:N:140:PHE:HB2 | 1.96 | 0.47 |
| 17:P:113:PHE:CE2 | 17:P:117:LEU:HD11 | 2.49 | 0.47 |
| 21:T:17:ILE:HD11 | 21:T:21:TYR:HD1 | 1.79 | 0.47 |
| 1:2:215:U:H2' | 1:2:216:U:C6 | 2.49 | 0.47 |
| 1:2:592:G:N2 | 1:2:611:C:N3 | 2.63 | 0.47 |
| 1:2:794:G:N7 | 11:J:106:ARG:HA | 2.29 | 0.47 |
| 1:2:904:A:H2' | 1:2:905:G:C8 | 2.50 | 0.47 |
| 1:2:959:A:H2' | 1:2:960:A:O4' | 2.14 | 0.47 |
| 1:2:1144:A:H4' | 1:2:1145:A:O4' | 2.15 | 0.47 |
| 1:2:1511:G:O3' | 19:R:122:THR:OG1 | 2.21 | 0.47 |
| 1:2:1687:U:H2' | 1:2:1688:G:C8 | 2.49 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|------------------|--------------------------|-------------------|
| 1:2:1859:C:C4 | 29:b:5:ARG:HA | 2.50 | 0.47 |
| 5:D:60:ASP:HA | 5:D:63:LYS:HD2 | 1.96 | 0.47 |
| 5:D:135:LEU:HB3 | 5:D:137:LEU:HD21 | 1.96 | 0.47 |
| 17:P:91:LEU:HD11 | 17:P:121:ARG:HD2 | 1.97 | 0.47 |
| 38:y:52:G:N2 | 38:y:53:A:H62 | 2.11 | 0.47 |
| 1:2:96:C:H2' | 1:2:97:U:C6 | 2.49 | 0.47 |
| 1:2:603:C:OP1 | 36:i:75:LYS:N | 2.47 | 0.47 |
| 1:2:1375:A:H2' | 1:2:1376:C:H6 | 1.79 | 0.47 |
| 20:S:25:CYS:SG | 20:S:95:TYR:HB2 | 2.54 | 0.47 |
| 21:T:79:GLU:HA | 21:T:82:ASP:HB3 | 1.96 | 0.47 |
| 26:Y:60:LYS:HZ2 | 26:Y:62:VAL:HG12 | 1.79 | 0.47 |
| 27:Z:90:CYS:O | 27:Z:94:ILE:HG12 | 2.15 | 0.47 |
| 39:z:584:A:H2' | 39:z:586:A:N7 | 2.29 | 0.47 |
| 1:2:3:C:C6 | 6:E:190:ILE:HD11 | 2.50 | 0.47 |
| 1:2:162:C:H2' | 1:2:163:U:O4' | 2.14 | 0.47 |
| 1:2:349:U:H5'' | 27:Z:22:TRP:CH2 | 2.50 | 0.47 |
| 1:2:361:A:H5' | 12:K:10:LYS:NZ | 2.30 | 0.47 |
| 1:2:492:C:C2 | 8:G:66:MET:HE1 | 2.50 | 0.47 |
| 1:2:876:G:H2' | 1:2:877:G:C8 | 2.50 | 0.47 |
| 1:2:965:U:H3' | 1:2:966:G:N2 | 2.29 | 0.47 |
| 1:2:1028:C:H2' | 1:2:1029:G:O4' | 2.15 | 0.47 |
| 1:2:1223:G:C2 | 1:2:1224:A:C8 | 3.02 | 0.47 |
| 1:2:1302:U:H2' | 1:2:1303:U:O4' | 2.15 | 0.47 |
| 1:2:1348:G:H2' | 1:2:1349:A:H8 | 1.79 | 0.47 |
| 1:2:1666:G:H2' | 1:2:1667:U:C6 | 2.49 | 0.47 |
| 1:2:1674:A:H5'' | 9:H:60:ARG:HH21 | 1.78 | 0.47 |
| 1:2:1718:G:H2' | 1:2:1719:A:C8 | 2.50 | 0.47 |
| 1:2:1718:G:H2' | 1:2:1719:A:H8 | 1.78 | 0.47 |
| 1:2:1795:A:H2' | 1:2:1796:C:H6 | 1.79 | 0.47 |
| 1:2:1847:C:H5 | 37:l:5:TRP:CH2 | 2.33 | 0.47 |
| 4:C:35:GLU:O | 4:C:38:ILE:HG22 | 2.15 | 0.47 |
| 6:E:95:MET:SD | 6:E:110:LYS:HB3 | 2.55 | 0.47 |
| 8:G:125:LYS:HB3 | 8:G:226:PHE:HD1 | 1.79 | 0.47 |
| 9:H:49:LEU:HD21 | 20:S:50:LYS:HB2 | 1.97 | 0.47 |
| 10:I:202:ASN:OD1 | 10:I:203:LYS:N | 2.47 | 0.47 |
| 11:J:78:ARG:HG3 | 11:J:81:ARG:NH2 | 2.29 | 0.47 |
| 16:O:22:LEU:HD22 | 16:O:88:TRP:HE3 | 1.80 | 0.47 |
| 16:O:46:GLN:NE2 | 16:O:113:ASP:OD2 | 2.48 | 0.47 |
| 23:V:42:HIS:CE1 | 23:V:43:LYS:HG2 | 2.49 | 0.47 |
| 27:Z:54:LYS:HG2 | 27:Z:70:VAL:HG12 | 1.96 | 0.47 |
| 34:g:119:GLN:HG2 | 34:g:139:LYS:NZ | 2.30 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 36:i:98:LYS:O | 36:i:99:LYS:HE2 | 2.14 | 0.47 |
| 1:2:552:U:O4 | 1:2:577:A:N7 | 2.48 | 0.47 |
| 1:2:553:G:O2' | 1:2:554:A:O5' | 2.29 | 0.47 |
| 1:2:827:G:H2' | 1:2:828:G:H8 | 1.79 | 0.47 |
| 1:2:952:G:H2' | 1:2:953:A:H8 | 1.80 | 0.47 |
| 1:2:1097:U:H2' | 1:2:1098:G:C8 | 2.50 | 0.47 |
| 1:2:1333:C:H5' | 24:W:67:LYS:HE3 | 1.96 | 0.47 |
| 1:2:1688:G:N2 | 1:2:1828:A:H8 | 2.13 | 0.47 |
| 1:2:1748:G:C6 | 1:2:1773:G:C6 | 3.02 | 0.47 |
| 4:C:89:LYS:HZ1 | 4:C:202:TYR:HA | 1.80 | 0.47 |
| 5:D:229:MET:O | 5:D:233:GLY:N | 2.48 | 0.47 |
| 7:F:25:LEU:HD12 | 7:F:37:VAL:HG11 | 1.96 | 0.47 |
| 18:Q:95:ILE:HD13 | 18:Q:129:ILE:HG22 | 1.96 | 0.47 |
| 22:U:120:HIS:O | 22:U:124:ARG:HG2 | 2.15 | 0.47 |
| 23:V:83:GLN:OE1 | 23:V:85:ASN:HB2 | 2.15 | 0.47 |
| 34:g:260:ASP:HB2 | 34:g:267:VAL:HG22 | 1.96 | 0.47 |
| 38:y:16:C:O2 | 38:y:58:A:N6 | 2.48 | 0.47 |
| 1:2:498:A:H3' | 1:2:499:G:H8 | 1.80 | 0.47 |
| 1:2:845:A:H2' | 1:2:846:C:C6 | 2.50 | 0.47 |
| 1:2:1431:C:H2' | 1:2:1432:C:C4 | 2.50 | 0.47 |
| 1:2:1614:A:OP2 | 19:R:47:ARG:NH2 | 2.37 | 0.47 |
| 1:2:1649:G:H2' | 1:2:1650:C:C6 | 2.50 | 0.47 |
| 1:2:1697:G:O6 | 1:2:1830:G:N2 | 2.48 | 0.47 |
| 8:G:9:LEU:HD12 | 8:G:30:ARG:HA | 1.95 | 0.47 |
| 25:X:38:GLU:OE1 | 25:X:50:SER:HA | 2.14 | 0.47 |
| 1:2:560:C:H2' | 1:2:561:U:H6 | 1.80 | 0.47 |
| 1:2:599:U:H2' | 1:2:600:G:H8 | 1.79 | 0.47 |
| 1:2:1413:C:H5'' | 1:2:1414:C:OP2 | 2.15 | 0.47 |
| 1:2:1624:C:H5'' | 22:U:39:ARG:HG2 | 1.96 | 0.47 |
| 20:S:39:LEU:HA | 20:S:42:ILE:HD11 | 1.97 | 0.47 |
| 22:U:80:PRO:HB2 | 22:U:82:TRP:CD1 | 2.50 | 0.47 |
| 23:V:134:ILE:O | 23:V:138:VAL:HG23 | 2.15 | 0.47 |
| 24:W:66:ARG:HE | 24:W:77:TRP:NE1 | 2.13 | 0.47 |
| 27:Z:91:LEU:HB3 | 36:i:83:VAL:HG11 | 1.97 | 0.47 |
| 28:a:109:GLU:O | 28:a:113:ARG:HG2 | 2.15 | 0.47 |
| 34:g:45:LEU:HB3 | 34:g:47:ARG:HH12 | 1.79 | 0.47 |
| 1:2:215:U:H2' | 1:2:216:U:H6 | 1.80 | 0.46 |
| 1:2:962:U:H2' | 1:2:963:C:H6 | 1.80 | 0.46 |
| 1:2:1560:C:H2' | 1:2:1561:G:C8 | 2.50 | 0.46 |
| 1:2:1769:U:H2' | 1:2:1770:G:C8 | 2.51 | 0.46 |
| 21:T:73:LEU:O | 21:T:76:GLU:HG2 | 2.14 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|------------------|--------------------------|-------------------|
| 37:l:12:ARG:HH21 | 37:l:15:ARG:CZ | 2.28 | 0.46 |
| 1:2:424:G:H2' | 1:2:425:A:C8 | 2.51 | 0.46 |
| 1:2:469:C:C2 | 1:2:470:G:C8 | 3.03 | 0.46 |
| 1:2:918:A:H2' | 1:2:919:G:O4' | 2.15 | 0.46 |
| 1:2:1031:A:H2' | 1:2:1032:A:O4' | 2.15 | 0.46 |
| 1:2:1160:G:C2 | 1:2:1161:G:C6 | 3.04 | 0.46 |
| 1:2:1301:C:O5' | 1:2:1301:C:H6 | 1.98 | 0.46 |
| 1:2:1615:A:P | 19:R:40:ARG:HH21 | 2.38 | 0.46 |
| 1:2:1811:G:H2' | 1:2:1812:A:C8 | 2.51 | 0.46 |
| 4:C:141:ASN:HB3 | 25:X:32:ILE:HG22 | 1.97 | 0.46 |
| 12:K:22:HIS:HB2 | 12:K:25:ARG:NH1 | 2.30 | 0.46 |
| 18:Q:132:VAL:HG13 | 18:Q:132:VAL:O | 2.16 | 0.46 |
| 19:R:25:LEU:HB3 | 19:R:87:PRO:HG3 | 1.97 | 0.46 |
| 27:Z:103:ALA:O | 27:Z:121:LYS:N | 2.44 | 0.46 |
| 1:2:3:C:C4 | 13:L:17:ARG:HD2 | 2.50 | 0.46 |
| 1:2:11:A:N1 | 1:2:1196:A:H2 | 2.13 | 0.46 |
| 1:2:410:G:H5'' | 15:N:97:ARG:HH22 | 1.80 | 0.46 |
| 1:2:553:G:O2' | 1:2:554:A:H8 | 1.98 | 0.46 |
| 1:2:744:C:H2' | 1:2:745:U:C4 | 2.50 | 0.46 |
| 1:2:915:A:O3' | 26:Y:57:ARG:NH2 | 2.49 | 0.46 |
| 1:2:1560:C:OP2 | 23:V:102:ARG:NH2 | 2.38 | 0.46 |
| 1:2:1678:C:C4 | 31:d:24:GLN:HG3 | 2.50 | 0.46 |
| 1:2:1688:G:H5' | 1:2:1857:A:H5'' | 1.97 | 0.46 |
| 1:2:1814:G:H2' | 1:2:1815:U:C6 | 2.51 | 0.46 |
| 6:E:79:ILE:O | 6:E:83:LEU:HB2 | 2.15 | 0.46 |
| 10:I:57:ASP:HA | 10:I:106:LEU:HA | 1.96 | 0.46 |
| 15:N:35:ARG:NH2 | 15:N:53:GLY:O | 2.48 | 0.46 |
| 22:U:30:ILE:HG22 | 22:U:36:VAL:HG11 | 1.96 | 0.46 |
| 25:X:4:ASN:OD1 | 25:X:5:ALA:N | 2.48 | 0.46 |
| 28:a:3:ASP:OD1 | 28:a:4:THR:N | 2.47 | 0.46 |
| 38:y:27:U:H2' | 38:y:28:G:H8 | 1.81 | 0.46 |
| 1:2:10:G:O4' | 1:2:1692:A:O2' | 2.33 | 0.46 |
| 1:2:51:U:H2' | 1:2:52:G:C8 | 2.51 | 0.46 |
| 1:2:223:A:N6 | 1:2:278:U:OP1 | 2.45 | 0.46 |
| 1:2:366:A:H2' | 1:2:367:G:O4' | 2.15 | 0.46 |
| 1:2:521:A:H2' | 1:2:522:C:C6 | 2.49 | 0.46 |
| 1:2:668:U:N3 | 1:2:1023:A:N6 | 2.53 | 0.46 |
| 1:2:1007:A:H5'' | 17:P:3:ARG:NH2 | 2.29 | 0.46 |
| 1:2:1179:A:H2' | 1:2:1180:G:H8 | 1.79 | 0.46 |
| 1:2:1743:G:N2 | 1:2:1780:U:O2 | 2.36 | 0.46 |
| 1:2:1798:U:H2' | 1:2:1799:G:C8 | 2.51 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 4:C:184:ARG:HA | 4:C:190:SER:HB3 | 1.97 | 0.46 |
| 12:K:53:LYS:HA | 12:K:53:LYS:HD3 | 1.78 | 0.46 |
| 12:K:114:GLU:CD | 12:K:123:ARG:HH22 | 2.23 | 0.46 |
| 15:N:50:ALA:HB2 | 15:N:116:CYS:SG | 2.56 | 0.46 |
| 16:O:48:HIS:HB3 | 16:O:113:ASP:H | 1.79 | 0.46 |
| 20:S:40:GLU:O | 20:S:48:GLN:NE2 | 2.44 | 0.46 |
| 27:Z:72:VAL:HG21 | 27:Z:100:VAL:HG21 | 1.97 | 0.46 |
| 29:b:71:LEU:HD13 | 29:b:73:TYR:CE1 | 2.50 | 0.46 |
| 1:2:76:U:H5'' | 1:2:77:A:H5'' | 1.98 | 0.46 |
| 1:2:107:A:H2' | 1:2:108:G:C8 | 2.51 | 0.46 |
| 1:2:110:U:H4' | 15:N:71:ARG:HH12 | 1.81 | 0.46 |
| 1:2:151:C:H2' | 1:2:152:U:C6 | 2.50 | 0.46 |
| 1:2:483:A:H62 | 1:2:499:G:N2 | 2.12 | 0.46 |
| 1:2:863:G:N3 | 1:2:864:G:N1 | 2.63 | 0.46 |
| 1:2:1029:G:N2 | 1:2:1076:A:HO2' | 2.14 | 0.46 |
| 1:2:1214:C:H2' | 1:2:1215:C:C6 | 2.50 | 0.46 |
| 1:2:1271:G:H1 | 1:2:1502:A:H2' | 1.81 | 0.46 |
| 1:2:1294:G:H5'' | 19:R:77:LYS:HB2 | 1.97 | 0.46 |
| 1:2:1708:C:H2' | 1:2:1709:U:C6 | 2.50 | 0.46 |
| 8:G:251:GLU:O | 8:G:254:LYS:HG2 | 2.16 | 0.46 |
| 14:M:59:LYS:HE3 | 14:M:70:TYR:HB3 | 1.98 | 0.46 |
| 20:S:38:PRO:HG2 | 23:V:8:ASP:HA | 1.98 | 0.46 |
| 21:T:45:LYS:HE2 | 21:T:49:LYS:HD2 | 1.97 | 0.46 |
| 26:Y:60:LYS:C | 26:Y:60:LYS:HD3 | 2.41 | 0.46 |
| 1:2:308:C:OP2 | 10:I:183:ARG:NH2 | 2.49 | 0.46 |
| 1:2:592:G:OP2 | 1:2:593:C:O2' | 2.30 | 0.46 |
| 1:2:865:A:C4 | 1:2:911:G:H1' | 2.50 | 0.46 |
| 1:2:1702:U:H2' | 1:2:1703:C:C6 | 2.50 | 0.46 |
| 1:2:1834:U:H2' | 1:2:1835:C:C6 | 2.50 | 0.46 |
| 1:2:1861:U:H5' | 29:b:97:PRO:HG3 | 1.96 | 0.46 |
| 9:H:40:ALA:HB1 | 9:H:45:TYR:CD2 | 2.50 | 0.46 |
| 11:J:65:PRO:HB2 | 11:J:67:PRO:HD3 | 1.98 | 0.46 |
| 11:J:75:ILE:HG13 | 11:J:78:ARG:HD3 | 1.98 | 0.46 |
| 11:J:168:HIS:HE1 | 11:J:169:LYS:HE3 | 1.80 | 0.46 |
| 15:N:38:LYS:NZ | 15:N:39:ASN:OD1 | 2.33 | 0.46 |
| 20:S:146:ARG:HH21 | 38:y:31:C:P | 2.39 | 0.46 |
| 38:y:10:G:C8 | 38:y:45:G:C2 | 3.04 | 0.46 |
| 1:2:160:U:O2' | 1:2:161:U:H3' | 2.15 | 0.46 |
| 1:2:318:U:H2' | 1:2:319:G:H8 | 1.81 | 0.46 |
| 1:2:399:C:H2' | 1:2:400:G:H8 | 1.80 | 0.46 |
| 1:2:730:C:H2' | 1:2:731:C:C5 | 2.51 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 1:2:749:C:H2' | 1:2:750:G:C8 | 2.50 | 0.46 |
| 1:2:833:A:OP2 | 28:a:8:ARG:NH1 | 2.49 | 0.46 |
| 1:2:1058:A:H2' | 1:2:1059:C:C6 | 2.51 | 0.46 |
| 1:2:1123:C:H2' | 1:2:1124:C:C6 | 2.50 | 0.46 |
| 1:2:1609:A:OP2 | 19:R:42:ARG:NH2 | 2.48 | 0.46 |
| 7:F:165:ASN:OD1 | 7:F:166:TYR:N | 2.48 | 0.46 |
| 11:J:67:PRO:HG2 | 11:J:68:GLN:OE1 | 2.16 | 0.46 |
| 12:K:8:TRP:O | 12:K:18:ARG:HD2 | 2.15 | 0.46 |
| 12:K:197:PHE:HA | 12:K:200:ARG:HG2 | 1.98 | 0.46 |
| 14:M:21:MET:HB3 | 14:M:69:TRP:HB2 | 1.97 | 0.46 |
| 27:Z:1:MET:HE2 | 27:Z:5:ARG:HB2 | 1.98 | 0.46 |
| 34:g:101:PHE:HB3 | 34:g:132:TRP:CZ3 | 2.50 | 0.46 |
| 39:z:560:C:H2' | 39:z:561:U:C6 | 2.51 | 0.46 |
| 1:2:29:G:C6 | 1:2:636:G:C6 | 3.04 | 0.46 |
| 1:2:147:A:H62 | 10:I:137:ARG:NH2 | 2.14 | 0.46 |
| 1:2:369:C:H2' | 1:2:370:G:C8 | 2.51 | 0.46 |
| 1:2:517:C:O2 | 1:2:549:G:N2 | 2.48 | 0.46 |
| 1:2:1537:C:O2' | 20:S:43:GLU:OE2 | 2.29 | 0.46 |
| 1:2:1596:A:H1' | 1:2:1599:G:O6 | 2.16 | 0.46 |
| 1:2:1666:G:H2' | 1:2:1667:U:H6 | 1.80 | 0.46 |
| 4:C:9:GLN:HG3 | 4:C:10:MET:H | 1.81 | 0.46 |
| 7:F:26:THR:HG22 | 7:F:34:TYR:CE1 | 2.51 | 0.46 |
| 7:F:64:ARG:HG2 | 7:F:68:GLU:OE2 | 2.16 | 0.46 |
| 8:G:125:LYS:HE3 | 8:G:226:PHE:HD1 | 1.81 | 0.46 |
| 9:H:198:ARG:HA | 9:H:201:LYS:HG2 | 1.97 | 0.46 |
| 11:J:51:ILE:HD11 | 11:J:176:VAL:HG23 | 1.97 | 0.46 |
| 14:M:25:LYS:HB3 | 14:M:65:ARG:HB3 | 1.97 | 0.46 |
| 19:R:14:LYS:HE2 | 39:z:563:C:H2' | 1.98 | 0.46 |
| 19:R:38:SER:O | 19:R:42:ARG:N | 2.46 | 0.46 |
| 29:b:31:PRO:HG2 | 29:b:34:LYS:HB2 | 1.96 | 0.46 |
| 35:h:51:ASP:HB2 | 35:h:54:THR:HG23 | 1.98 | 0.46 |
| 1:2:121:U:H5'' | 8:G:77:ARG:NH1 | 2.30 | 0.46 |
| 1:2:841:G:C6 | 1:2:842:G:C6 | 3.04 | 0.46 |
| 1:2:1142:C:O2' | 1:2:1146:A:N1 | 2.49 | 0.46 |
| 1:2:1234:U:H2' | 1:2:1235:U:C6 | 2.51 | 0.46 |
| 1:2:1466:C:H2' | 1:2:1467:C:H6 | 1.81 | 0.46 |
| 4:C:9:GLN:HA | 4:C:55:TRP:CH2 | 2.51 | 0.46 |
| 5:D:134:LEU:HD12 | 5:D:219:LYS:HG3 | 1.97 | 0.46 |
| 6:E:108:ARG:HG2 | 6:E:130:LYS:HA | 1.98 | 0.46 |
| 7:F:122:VAL:O | 7:F:126:ILE:HG12 | 2.15 | 0.46 |
| 11:J:142:LYS:C | 11:J:143:ARG:HD2 | 2.41 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 18:Q:67:ASP:HB3 | 18:Q:70:SER:HB3 | 1.98 | 0.46 |
| 21:T:96:ILE:HD12 | 21:T:96:ILE:HA | 1.85 | 0.46 |
| 28:a:44:LEU:HD23 | 28:a:55:ILE:HG23 | 1.98 | 0.46 |
| 33:f:135:HIS:CD2 | 33:f:140:TYR:HB3 | 2.51 | 0.46 |
| 39:z:547:G:N1 | 39:z:550:A:OP2 | 2.43 | 0.46 |
| 1:2:1226:C:H1' | 1:2:1660:G:H22 | 1.81 | 0.46 |
| 1:2:1383:G:C2 | 1:2:1384:A:H1' | 2.51 | 0.46 |
| 3:B:254:PRO:HA | 3:B:286:GLY:HA3 | 1.98 | 0.46 |
| 4:C:87:VAL:HG12 | 4:C:175:TRP:CZ2 | 2.51 | 0.46 |
| 4:C:144:THR:N | 4:C:158:ASP:OD2 | 2.38 | 0.46 |
| 5:D:133:TYR:HA | 5:D:219:LYS:O | 2.16 | 0.46 |
| 8:G:139:LEU:HD12 | 8:G:147:ILE:HB | 1.97 | 0.46 |
| 12:K:197:PHE:O | 12:K:200:ARG:HG2 | 2.16 | 0.46 |
| 14:M:5:LYS:O | 14:M:9:ILE:HD12 | 2.16 | 0.46 |
| 33:f:111:ASN:HA | 33:f:112:GLY:HA2 | 1.65 | 0.46 |
| 34:g:212:LYS:HA | 34:g:235:ILE:HG13 | 1.98 | 0.46 |
| 1:2:29:G:H4' | 27:Z:129:SER:HB3 | 1.98 | 0.45 |
| 1:2:121:U:O3' | 8:G:77:ARG:NH2 | 2.43 | 0.45 |
| 1:2:422:G:H2' | 1:2:423:A:C8 | 2.51 | 0.45 |
| 1:2:1035:C:N4 | 1:2:1073:A:H61 | 2.14 | 0.45 |
| 1:2:1200:A:H2' | 1:2:1201:C:C6 | 2.50 | 0.45 |
| 1:2:1270:G:O4' | 14:M:47:LYS:HE2 | 2.16 | 0.45 |
| 1:2:1587:C:H5'' | 9:H:91:ARG:HH22 | 1.81 | 0.45 |
| 1:2:1606:G:H2' | 1:2:1607:G:C8 | 2.51 | 0.45 |
| 4:C:22:GLY:O | 4:C:164:ASN:ND2 | 2.42 | 0.45 |
| 5:D:147:ASN:OD1 | 5:D:148:ASN:N | 2.49 | 0.45 |
| 9:H:70:GLU:O | 9:H:73:THR:OG1 | 2.24 | 0.45 |
| 12:K:200:ARG:O | 12:K:203:LYS:HG3 | 2.16 | 0.45 |
| 17:P:141:TYR:CE2 | 17:P:146:ALA:HB2 | 2.50 | 0.45 |
| 24:W:80:PHE:HA | 32:e:52:PHE:CE1 | 2.51 | 0.45 |
| 28:a:44:LEU:O | 28:a:47:MET:HB3 | 2.16 | 0.45 |
| 30:c:33:MET:HB2 | 30:c:80:ARG:H | 1.81 | 0.45 |
| 31:d:37:ASP:N | 31:d:37:ASP:OD1 | 2.49 | 0.45 |
| 34:g:153:CYS:SG | 34:g:155:ARG:NH2 | 2.88 | 0.45 |
| 38:y:8:U:O2' | 38:y:10:G:OP1 | 2.29 | 0.45 |
| 38:y:30:G:H2' | 38:y:31:C:H6 | 1.80 | 0.45 |
| 1:2:369:C:H4' | 12:K:31:ARG:O | 2.15 | 0.45 |
| 3:B:86:ALA:N | 3:B:130:VAL:O | 2.49 | 0.45 |
| 4:C:73:ASP:OD1 | 4:C:73:ASP:N | 2.49 | 0.45 |
| 4:C:110:ASN:OD1 | 4:C:112:ILE:N | 2.41 | 0.45 |
| 5:D:26:SER:O | 5:D:51:ARG:NH2 | 2.47 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 7:F:55:THR:HA | 7:F:58:VAL:HB | 1.98 | 0.45 |
| 7:F:66:ILE:O | 7:F:70:THR:HG23 | 2.16 | 0.45 |
| 11:J:160:LYS:HA | 11:J:189:PHE:CZ | 2.51 | 0.45 |
| 12:K:198:TYR:HE1 | 15:N:10:TYR:HB2 | 1.81 | 0.45 |
| 13:L:147:PHE:CD2 | 13:L:149:VAL:HG22 | 2.50 | 0.45 |
| 22:U:24:ARG:HG3 | 22:U:25:LYS:N | 2.32 | 0.45 |
| 27:Z:25:LYS:HG3 | 27:Z:28:LYS:HE2 | 1.98 | 0.45 |
| 34:g:123:GLY:HA3 | 34:g:151:VAL:HG11 | 1.98 | 0.45 |
| 34:g:259:TRP:C | 34:g:267:VAL:HG23 | 2.41 | 0.45 |
| 1:2:354:A:H2' | 1:2:355:C:H6 | 1.82 | 0.45 |
| 1:2:625:G:H2' | 1:2:626:C:C6 | 2.52 | 0.45 |
| 1:2:1014:U:H2' | 1:2:1015:C:C6 | 2.50 | 0.45 |
| 1:2:1240:U:H2' | 1:2:1241:G:C8 | 2.46 | 0.45 |
| 1:2:1395:C:H5'' | 34:g:100:ARG:CZ | 2.46 | 0.45 |
| 1:2:1414:C:H2' | 1:2:1420:G:H5' | 1.98 | 0.45 |
| 1:2:1506:G:H2' | 1:2:1506:G:N3 | 2.31 | 0.45 |
| 1:2:1712:C:H2' | 1:2:1713:G:O4' | 2.17 | 0.45 |
| 5:D:129:THR:OG1 | 5:D:131:ASP:OD1 | 2.26 | 0.45 |
| 8:G:195:ILE:HG23 | 8:G:196:THR:N | 2.30 | 0.45 |
| 12:K:116:HIS:CD2 | 12:K:152:ARG:HE | 2.35 | 0.45 |
| 22:U:18:THR:HG21 | 22:U:33:ILE:HG13 | 1.99 | 0.45 |
| 24:W:36:CYS:O | 24:W:40:ILE:HG12 | 2.16 | 0.45 |
| 29:b:89:ARG:O | 29:b:92:ARG:HG2 | 2.15 | 0.45 |
| 39:z:552:A:H2' | 39:z:553:G:C8 | 2.52 | 0.45 |
| 1:2:220:C:H2' | 1:2:221:A:H8 | 1.81 | 0.45 |
| 1:2:310:G:N2 | 1:2:322:G:C4 | 2.84 | 0.45 |
| 1:2:541:U:H2' | 1:2:542:G:C8 | 2.52 | 0.45 |
| 1:2:1013:U:H2' | 1:2:1014:U:C6 | 2.51 | 0.45 |
| 6:E:241:TRP:HH2 | 26:Y:46:TYR:CE1 | 2.34 | 0.45 |
| 8:G:125:LYS:HB3 | 8:G:226:PHE:CD1 | 2.51 | 0.45 |
| 9:H:191:LYS:HE3 | 9:H:192:LYS:HE3 | 1.98 | 0.45 |
| 14:M:62:PHE:CZ | 14:M:65:ARG:HA | 2.51 | 0.45 |
| 17:P:19:ARG:NH2 | 17:P:21:SER:HB3 | 2.32 | 0.45 |
| 27:Z:61:GLN:OE1 | 27:Z:61:GLN:N | 2.34 | 0.45 |
| 32:e:5:GLN:OE1 | 32:e:6:LEU:HG | 2.17 | 0.45 |
| 34:g:21:ILE:HG22 | 34:g:33:SER:HA | 1.98 | 0.45 |
| 34:g:208:ALA:HA | 34:g:218:LEU:HD13 | 1.98 | 0.45 |
| 34:g:240:CYS:SG | 34:g:241:PHE:N | 2.89 | 0.45 |
| 1:2:5:U:H2' | 1:2:6:G:H8 | 1.80 | 0.45 |
| 1:2:106:C:H5'' | 1:2:421:G:O2' | 2.16 | 0.45 |
| 1:2:198:U:H3' | 1:2:199:G:H8 | 1.81 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|------------------|--------------------------|-------------------|
| 1:2:549:G:H2' | 1:2:550:A:H8 | 1.82 | 0.45 |
| 1:2:847:C:H5'' | 1:2:848:G:C5' | 2.46 | 0.45 |
| 1:2:1316:G:H2' | 1:2:1317:G:O4' | 2.16 | 0.45 |
| 1:2:1326:G:N2 | 1:2:1331:G:O6 | 2.50 | 0.45 |
| 1:2:1373:U:H3' | 4:C:102:ARG:NH2 | 2.31 | 0.45 |
| 1:2:1427:G:C2 | 1:2:1428:U:C4 | 3.05 | 0.45 |
| 1:2:1589:A:H62 | 35:h:104:ARG:HD2 | 1.81 | 0.45 |
| 4:C:120:ARG:NH2 | 6:E:251:TYR:HB3 | 2.31 | 0.45 |
| 9:H:142:SER:O | 9:H:146:ARG:N | 2.46 | 0.45 |
| 10:I:16:ILE:HG22 | 10:I:18:VAL:HG13 | 1.99 | 0.45 |
| 10:I:76:LEU:HD23 | 10:I:94:ARG:HB2 | 1.99 | 0.45 |
| 13:L:18:ARG:O | 13:L:24:ARG:NH1 | 2.50 | 0.45 |
| 15:N:101:ARG:HD2 | 27:Z:7:LEU:HA | 1.98 | 0.45 |
| 38:y:17:G:O2' | 38:y:59:A:N1 | 2.44 | 0.45 |
| 38:y:65:C:H2' | 38:y:66:U:C6 | 2.52 | 0.45 |
| 1:2:664:C:H2' | 1:2:665:U:C6 | 2.52 | 0.45 |
| 1:2:736:C:O4' | 11:J:99:ARG:NH2 | 2.50 | 0.45 |
| 1:2:1232:G:O6 | 22:U:137:LYS:HE3 | 2.16 | 0.45 |
| 1:2:1236:A:H2' | 1:2:1237:A:C8 | 2.52 | 0.45 |
| 1:2:1450:A:OP1 | 21:T:3:ARG:NH1 | 2.50 | 0.45 |
| 5:D:198:GLU:O | 5:D:202:GLN:HG3 | 2.17 | 0.45 |
| 8:G:212:ASP:OD1 | 8:G:216:ASN:N | 2.40 | 0.45 |
| 9:H:36:GLN:C | 9:H:38:TYR:H | 2.24 | 0.45 |
| 15:N:128:VAL:HG12 | 15:N:142:VAL:HA | 1.97 | 0.45 |
| 16:O:53:ALA:HA | 16:O:79:VAL:O | 2.17 | 0.45 |
| 17:P:45:LEU:HD13 | 17:P:49:GLN:HB3 | 1.98 | 0.45 |
| 18:Q:72:TYR:O | 18:Q:75:MET:HG3 | 2.16 | 0.45 |
| 23:V:139:ALA:HA | 23:V:144:LYS:HB2 | 1.98 | 0.45 |
| 28:a:20:ARG:HD2 | 28:a:74:MET:CE | 2.46 | 0.45 |
| 28:a:48:TYR:O | 28:a:50:THR:HG23 | 2.17 | 0.45 |
| 1:2:373:G:H2' | 1:2:374:U:C6 | 2.51 | 0.45 |
| 1:2:379:A:H2' | 1:2:380:C:H6 | 1.81 | 0.45 |
| 1:2:1138:G:H2' | 1:2:1140:A:OP2 | 2.17 | 0.45 |
| 5:D:144:LYS:NZ | 5:D:146:CYS:HA | 2.32 | 0.45 |
| 7:F:168:VAL:HG12 | 7:F:189:MET:HG3 | 1.98 | 0.45 |
| 8:G:137:PRO:HD2 | 8:G:149:TYR:CD1 | 2.51 | 0.45 |
| 11:J:116:ARG:NH2 | 11:J:121:THR:OG1 | 2.49 | 0.45 |
| 11:J:145:ARG:NH2 | 26:Y:51:GLU:HB2 | 2.32 | 0.45 |
| 26:Y:15:ASN:O | 26:Y:19:LYS:HG3 | 2.17 | 0.45 |
| 28:a:101:LYS:HB3 | 28:a:103:SER:HB2 | 1.98 | 0.45 |
| 35:h:70:PRO:HG3 | 35:h:88:LEU:HD21 | 1.98 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 1:2:641:U:H2' | 1:2:642:U:C6 | 2.52 | 0.45 |
| 1:2:860:A:H2' | 1:2:861:A:H8 | 1.81 | 0.45 |
| 1:2:1351:C:H4' | 6:E:223:LYS:NZ | 2.31 | 0.45 |
| 1:2:1562:G:H5'' | 23:V:98:SER:HB3 | 1.99 | 0.45 |
| 1:2:1588:C:H4' | 20:S:45:ARG:HH21 | 1.82 | 0.45 |
| 1:2:1618:A:O5' | 22:U:133:GLY:HA3 | 2.17 | 0.45 |
| 3:B:58:VAL:O | 3:B:62:SER:N | 2.41 | 0.45 |
| 8:G:212:ASP:CG | 8:G:214:ASN:H | 2.24 | 0.45 |
| 17:P:20:ARG:HG3 | 17:P:65:PHE:CE1 | 2.52 | 0.45 |
| 20:S:134:GLY:HA3 | 20:S:140:ARG:HA | 1.98 | 0.45 |
| 22:U:90:VAL:HG12 | 22:U:91:LYS:HG3 | 1.99 | 0.45 |
| 1:2:344:U:OP1 | 15:N:90:ARG:NH1 | 2.50 | 0.45 |
| 1:2:422:G:H2' | 1:2:423:A:H8 | 1.82 | 0.45 |
| 1:2:559:A:H2' | 1:2:560:C:C6 | 2.52 | 0.45 |
| 1:2:1168:U:P | 37:l:14:LYS:HZ1 | 2.40 | 0.45 |
| 1:2:1369:C:OP1 | 21:T:7:LYS:HG2 | 2.16 | 0.45 |
| 1:2:1402:G:H21 | 1:2:1439:C:N4 | 2.14 | 0.45 |
| 4:C:50:ASN:ND2 | 4:C:53:ARG:HG2 | 2.31 | 0.45 |
| 5:D:78:GLU:N | 5:D:78:GLU:OE1 | 2.50 | 0.45 |
| 6:E:172:ARG:HE | 6:E:172:ARG:HB3 | 1.57 | 0.45 |
| 17:P:96:VAL:O | 17:P:99:ARG:HG3 | 2.16 | 0.45 |
| 18:Q:127:GLY:HA2 | 29:b:58:VAL:HG22 | 1.99 | 0.45 |
| 23:V:44:GLU:HG2 | 23:V:45:LEU:HG | 1.99 | 0.45 |
| 34:g:154:VAL:HG13 | 34:g:166:VAL:O | 2.17 | 0.45 |
| 1:2:284:C:H2' | 1:2:286:C:H5' | 1.99 | 0.45 |
| 1:2:384:G:O3' | 15:N:82:MET:HE1 | 2.17 | 0.45 |
| 1:2:419:C:O2' | 1:2:807:A:N1 | 2.42 | 0.45 |
| 1:2:554:A:H2' | 1:2:555:G:O4' | 2.17 | 0.45 |
| 1:2:1088:G:C2 | 1:2:1089:A:N7 | 2.85 | 0.45 |
| 1:2:1093:G:H2' | 1:2:1094:C:H6 | 1.82 | 0.45 |
| 1:2:1194:G:H2' | 1:2:1195:A:C8 | 2.52 | 0.45 |
| 4:C:198:MET:SD | 4:C:199:PRO:HD2 | 2.56 | 0.45 |
| 5:D:210:VAL:O | 5:D:210:VAL:HG13 | 2.17 | 0.45 |
| 6:E:62:SER:OG | 6:E:65:GLU:OE1 | 2.27 | 0.45 |
| 6:E:185:ARG:O | 13:L:98:LEU:HD11 | 2.17 | 0.45 |
| 6:E:196:LYS:HE2 | 6:E:199:LEU:HD12 | 1.99 | 0.45 |
| 17:P:71:ILE:HA | 17:P:74:ILE:HD13 | 1.99 | 0.45 |
| 17:P:115:LEU:HA | 17:P:118:ILE:HG12 | 1.99 | 0.45 |
| 23:V:96:SER:HB2 | 23:V:99:VAL:HG12 | 1.98 | 0.45 |
| 26:Y:102:ILE:HA | 26:Y:128:PHE:HA | 1.99 | 0.45 |
| 33:f:146:LEU:HD23 | 33:f:146:LEU:H | 1.81 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 34:g:152:SER:H | 34:g:169:GLY:HA2 | 1.82 | 0.45 |
| 37:l:19:LYS:NZ | 37:l:23:ARG:HH11 | 2.15 | 0.45 |
| 1:2:518:A:C4 | 1:2:548:G:N2 | 2.85 | 0.44 |
| 1:2:628:C:H2' | 1:2:629:C:C6 | 2.52 | 0.44 |
| 1:2:656:U:H4' | 1:2:1084:U:H5' | 1.98 | 0.44 |
| 1:2:839:C:H2' | 1:2:840:U:C6 | 2.52 | 0.44 |
| 1:2:1119:C:O3' | 5:D:149:GLN:HG3 | 2.17 | 0.44 |
| 1:2:1301:C:OP1 | 33:f:92:LYS:HE3 | 2.16 | 0.44 |
| 1:2:1305:C:OP2 | 33:f:105:TYR:OH | 2.21 | 0.44 |
| 1:2:1435:A:H2' | 1:2:1436:C:C6 | 2.52 | 0.44 |
| 1:2:1472:A:OP1 | 1:2:1473:U:N3 | 2.38 | 0.44 |
| 6:E:186:GLY:HA3 | 13:L:98:LEU:HD21 | 1.98 | 0.44 |
| 18:Q:31:CYS:SG | 18:Q:93:LEU:HD12 | 2.57 | 0.44 |
| 18:Q:119:LEU:O | 18:Q:124:MET:HB2 | 2.17 | 0.44 |
| 30:c:34:ASP:HB2 | 30:c:82:LYS:HE2 | 1.98 | 0.44 |
| 33:f:121:CYS:SG | 33:f:130:VAL:HG23 | 2.57 | 0.44 |
| 34:g:280:LYS:HD2 | 34:g:280:LYS:HA | 1.81 | 0.44 |
| 1:2:235:C:H2' | 1:2:236:C:C6 | 2.52 | 0.44 |
| 1:2:364:G:H2' | 1:2:365:U:H6 | 1.82 | 0.44 |
| 1:2:479:A:H2' | 1:2:480:C:C6 | 2.52 | 0.44 |
| 1:2:740:G:N3 | 11:J:109:ARG:HB2 | 2.32 | 0.44 |
| 1:2:853:U:H4' | 8:G:201:HIS:HE2 | 1.82 | 0.44 |
| 1:2:959:A:P | 18:Q:66:ARG:HB3 | 2.57 | 0.44 |
| 1:2:1626:U:H2' | 1:2:1627:G:O4' | 2.17 | 0.44 |
| 1:2:1685:U:H2' | 1:2:1686:U:H6 | 1.82 | 0.44 |
| 1:2:1850:C:H2' | 1:2:1851:G:C8 | 2.51 | 0.44 |
| 1:2:1853:A:H2' | 1:2:1854:A:C8 | 2.53 | 0.44 |
| 3:B:85:ASN:HA | 3:B:131:SER:HA | 1.99 | 0.44 |
| 34:g:125:ARG:HG2 | 34:g:150:TRP:CD2 | 2.52 | 0.44 |
| 37:l:8:LYS:HG3 | 37:l:12:ARG:HH11 | 1.82 | 0.44 |
| 1:2:1191:A:H2' | 1:2:1192:A:C8 | 2.52 | 0.44 |
| 1:2:1194:G:H2' | 1:2:1195:A:H8 | 1.82 | 0.44 |
| 1:2:1223:G:H5' | 1:2:1634:G:O6 | 2.17 | 0.44 |
| 1:2:1309:A:O2' | 16:O:91:LEU:HB2 | 2.17 | 0.44 |
| 1:2:1331:G:H1 | 1:2:1488:U:H3 | 1.66 | 0.44 |
| 1:2:1595:G:OP1 | 35:h:44:LEU:HB2 | 2.16 | 0.44 |
| 1:2:1850:C:H2' | 1:2:1851:G:H8 | 1.83 | 0.44 |
| 4:C:63:ARG:CZ | 25:X:78:ILE:HB | 2.47 | 0.44 |
| 5:D:37:ALA:N | 5:D:231:LEU:O | 2.30 | 0.44 |
| 5:D:223:PHE:CD2 | 5:D:225:LEU:HG | 2.51 | 0.44 |
| 7:F:218:LEU:HD21 | 7:F:220:THR:HG22 | 1.99 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 12:K:166:PHE:HA | 12:K:171:LEU:HD21 | 1.99 | 0.44 |
| 25:X:39:VAL:HG23 | 25:X:41:LYS:H | 1.82 | 0.44 |
| 34:g:42:MET:HE3 | 34:g:56:GLN:HG3 | 1.99 | 0.44 |
| 35:h:57:LYS:HG2 | 35:h:77:LEU:HD21 | 1.99 | 0.44 |
| 38:y:2:G:C6 | 38:y:71:U:H1' | 2.53 | 0.44 |
| 39:z:520:U:H2' | 39:z:521:U:C6 | 2.52 | 0.44 |
| 39:z:540:C:H2' | 39:z:541:C:C6 | 2.52 | 0.44 |
| 39:z:581:A:H2' | 39:z:582:U:C6 | 2.53 | 0.44 |
| 1:2:35:C:H2' | 1:2:36:U:C6 | 2.52 | 0.44 |
| 1:2:152:U:H2' | 1:2:153:G:C8 | 2.52 | 0.44 |
| 1:2:158:A:C2 | 1:2:453:C:H1' | 2.53 | 0.44 |
| 1:2:168:C:H2' | 1:2:169:U:O4' | 2.16 | 0.44 |
| 1:2:308:C:H5 | 10:I:183:ARG:HH21 | 1.65 | 0.44 |
| 1:2:404:A:H2' | 1:2:405:A:C8 | 2.52 | 0.44 |
| 1:2:678:U:O2 | 11:J:103:LYS:HE3 | 2.18 | 0.44 |
| 1:2:812:A:H2' | 1:2:813:G:O4' | 2.17 | 0.44 |
| 1:2:858:A:C5 | 1:2:859:U:C5 | 3.06 | 0.44 |
| 1:2:1098:G:H22 | 1:2:1126:G:N2 | 2.15 | 0.44 |
| 1:2:1109:A:H2' | 1:2:1110:U:C6 | 2.52 | 0.44 |
| 1:2:1176:C:H2' | 1:2:1177:A:O4' | 2.17 | 0.44 |
| 1:2:1444:A:H3' | 1:2:1445:G:H5'' | 1.99 | 0.44 |
| 1:2:1483:A:H2' | 1:2:1484:C:C6 | 2.51 | 0.44 |
| 1:2:1767:C:H2' | 1:2:1768:C:C6 | 2.53 | 0.44 |
| 5:D:171:ILE:HG23 | 5:D:174:ARG:HH21 | 1.82 | 0.44 |
| 7:F:22:ASN:OD1 | 7:F:23:GLU:N | 2.50 | 0.44 |
| 8:G:64:ILE:HD12 | 28:a:17:LEU:CD2 | 2.47 | 0.44 |
| 12:K:12:ARG:NH2 | 12:K:18:ARG:HB3 | 2.33 | 0.44 |
| 17:P:93:LYS:HA | 17:P:96:VAL:HG12 | 1.99 | 0.44 |
| 20:S:16:LYS:HD3 | 20:S:82:TYR:HB3 | 1.99 | 0.44 |
| 21:T:29:HIS:HA | 21:T:32:LYS:HE3 | 2.00 | 0.44 |
| 23:V:125:PRO:O | 23:V:128:GLN:NE2 | 2.50 | 0.44 |
| 25:X:11:LEU:HG | 25:X:12:TYR:HD1 | 1.82 | 0.44 |
| 33:f:105:TYR:HB2 | 33:f:131:PHE:CE2 | 2.51 | 0.44 |
| 37:l:8:LYS:O | 37:l:12:ARG:HG2 | 2.17 | 0.44 |
| 1:2:92:A:C8 | 1:2:437:A:C4 | 3.06 | 0.44 |
| 1:2:365:U:O2' | 15:N:7:GLU:OE2 | 2.22 | 0.44 |
| 1:2:444:U:O3' | 10:I:94:ARG:NH2 | 2.34 | 0.44 |
| 1:2:585:U:H2' | 1:2:586:U:C6 | 2.53 | 0.44 |
| 1:2:1195:A:H2' | 1:2:1196:A:H8 | 1.81 | 0.44 |
| 1:2:1289:A:H2' | 1:2:1290:G:H8 | 1.82 | 0.44 |
| 1:2:1398:A:H2' | 1:2:1401:A:C5 | 2.52 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 1:2:1460:C:P | 21:T:63:ARG:HH22 | 2.40 | 0.44 |
| 6:E:189:ILE:HB | 6:E:196:LYS:HZ2 | 1.81 | 0.44 |
| 7:F:218:LEU:HD11 | 34:g:192:THR:HA | 1.98 | 0.44 |
| 9:H:90:VAL:O | 9:H:94:LYS:HG2 | 2.18 | 0.44 |
| 12:K:67:TRP:HE1 | 15:N:20:LYS:NZ | 2.15 | 0.44 |
| 12:K:144:LYS:HA | 12:K:147:LYS:HB2 | 1.99 | 0.44 |
| 12:K:199:LEU:HA | 12:K:202:ILE:HG12 | 1.99 | 0.44 |
| 13:L:88:ASP:OD1 | 13:L:89:GLU:N | 2.51 | 0.44 |
| 14:M:38:LYS:O | 14:M:38:LYS:HD3 | 2.17 | 0.44 |
| 23:V:62:ARG:HH12 | 23:V:66:LEU:HD11 | 1.82 | 0.44 |
| 34:g:272:GLN:OE1 | 34:g:272:GLN:N | 2.44 | 0.44 |
| 1:2:12:U:H2' | 1:2:13:C:H6 | 1.81 | 0.44 |
| 1:2:332:C:H2' | 1:2:333:A:C8 | 2.51 | 0.44 |
| 1:2:449:C:H3' | 1:2:450:A:H8 | 1.83 | 0.44 |
| 1:2:539:C:H2' | 1:2:540:C:O4' | 2.17 | 0.44 |
| 1:2:548:G:H2' | 1:2:549:G:N9 | 2.33 | 0.44 |
| 1:2:749:C:H2' | 1:2:750:G:H8 | 1.82 | 0.44 |
| 1:2:818:U:H2' | 1:2:820:C:OP2 | 2.17 | 0.44 |
| 1:2:1707:A:H2' | 1:2:1708:C:C6 | 2.52 | 0.44 |
| 1:2:1732:G:C6 | 1:2:1791:U:O2 | 2.69 | 0.44 |
| 1:2:1799:G:H2' | 1:2:1800:A:H8 | 1.82 | 0.44 |
| 8:G:173:ILE:HD11 | 8:G:235:TRP:CE2 | 2.52 | 0.44 |
| 17:P:47:PRO:HA | 17:P:50:ILE:HD13 | 1.99 | 0.44 |
| 19:R:79:HIS:CE1 | 19:R:102:PHE:HZ | 2.36 | 0.44 |
| 22:U:6:PRO:O | 35:h:49:LEU:HD12 | 2.18 | 0.44 |
| 31:d:66:ARG:HA | 31:d:66:ARG:HD2 | 1.85 | 0.44 |
| 1:2:364:G:H2' | 1:2:365:U:C6 | 2.53 | 0.44 |
| 1:2:374:U:H4' | 15:N:134:LEU:O | 2.18 | 0.44 |
| 1:2:639:U:OP2 | 27:Z:106:GLY:HA2 | 2.17 | 0.44 |
| 1:2:932:G:H2' | 1:2:933:C:H6 | 1.81 | 0.44 |
| 1:2:1088:G:OP1 | 17:P:2:GLY:N | 2.51 | 0.44 |
| 1:2:1652:G:H2' | 1:2:1653:G:C8 | 2.53 | 0.44 |
| 5:D:104:ASP:OD1 | 5:D:105:LEU:N | 2.41 | 0.44 |
| 7:F:76:ARG:O | 7:F:76:ARG:NH1 | 2.43 | 0.44 |
| 8:G:11:ARG:N | 8:G:26:VAL:O | 2.33 | 0.44 |
| 9:H:126:THR:OG1 | 31:d:26:GLN:OE1 | 2.35 | 0.44 |
| 10:I:106:LEU:HD13 | 10:I:109:LEU:HD11 | 2.00 | 0.44 |
| 12:K:162:LEU:HB3 | 12:K:166:PHE:HE2 | 1.83 | 0.44 |
| 22:U:70:ILE:HA | 22:U:77:TYR:CE2 | 2.52 | 0.44 |
| 38:y:26:C:H2' | 38:y:27:U:H6 | 1.83 | 0.44 |
| 1:2:51:U:H2' | 1:2:52:G:H8 | 1.83 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|------------------|--------------------------|-------------------|
| 1:2:323:G:H2' | 1:2:324:C:H6 | 1.83 | 0.44 |
| 1:2:664:C:H2' | 1:2:665:U:H6 | 1.83 | 0.44 |
| 1:2:959:A:H2' | 1:2:960:A:C8 | 2.53 | 0.44 |
| 1:2:1076:A:OP2 | 1:2:1076:A:H8 | 2.01 | 0.44 |
| 1:2:1117:G:C6 | 1:2:1118:A:C5 | 3.05 | 0.44 |
| 1:2:1125:G:H2' | 1:2:1126:G:C4 | 2.53 | 0.44 |
| 1:2:1413:C:C2 | 1:2:1415:C:C5 | 3.06 | 0.44 |
| 1:2:1487:G:H2' | 1:2:1488:U:C6 | 2.53 | 0.44 |
| 1:2:1544:U:OP1 | 32:e:34:TYR:OH | 2.36 | 0.44 |
| 1:2:1748:G:H2' | 1:2:1749:C:H6 | 1.83 | 0.44 |
| 1:2:1853:A:O2' | 1:2:1854:A:H5' | 2.18 | 0.44 |
| 4:C:10:MET:HG2 | 21:T:111:PHE:CE2 | 2.47 | 0.44 |
| 4:C:66:VAL:HA | 4:C:186:ARG:HH11 | 1.82 | 0.44 |
| 4:C:189:ILE:H | 4:C:189:ILE:HD12 | 1.82 | 0.44 |
| 10:I:63:MET:SD | 10:I:100:CYS:HA | 2.57 | 0.44 |
| 17:P:110:ASP:O | 17:P:114:ARG:HG2 | 2.17 | 0.44 |
| 31:d:16:LYS:HD2 | 31:d:16:LYS:HA | 1.72 | 0.44 |
| 32:e:49:ASP:OD1 | 32:e:50:ILE:N | 2.47 | 0.44 |
| 39:z:527:G:H2' | 39:z:528:C:H6 | 1.83 | 0.44 |
| 1:2:414:C:H2' | 1:2:415:G:O4' | 2.17 | 0.44 |
| 1:2:1066:A:H2' | 1:2:1067:G:O4' | 2.18 | 0.44 |
| 1:2:1321:G:O2' | 1:2:1323:G:OP1 | 2.35 | 0.44 |
| 1:2:1451:A:H2' | 1:2:1452:G:H8 | 1.82 | 0.44 |
| 1:2:1561:G:N2 | 1:2:1563:C:H3' | 2.33 | 0.44 |
| 4:C:175:TRP:CE3 | 4:C:178:LEU:HD11 | 2.53 | 0.44 |
| 9:H:122:ARG:NH2 | 31:d:9:ILE:HD11 | 2.33 | 0.44 |
| 11:J:6:ALA:HB1 | 11:J:21:SER:HB3 | 2.00 | 0.44 |
| 11:J:99:ARG:HE | 11:J:100:ILE:H | 1.65 | 0.44 |
| 14:M:33:PRO:O | 14:M:34:GLU:HG2 | 2.18 | 0.44 |
| 19:R:94:VAL:O | 19:R:104:GLN:HA | 2.17 | 0.44 |
| 39:z:540:C:H2' | 39:z:541:C:H6 | 1.83 | 0.44 |
| 1:2:217:U:O4 | 1:2:293:A:N6 | 2.50 | 0.43 |
| 1:2:674:G:H2' | 1:2:675:A:H8 | 1.83 | 0.43 |
| 1:2:1108:U:H3 | 1:2:1109:A:H62 | 1.64 | 0.43 |
| 1:2:1179:A:O3' | 37:l:11:ARG:NH2 | 2.50 | 0.43 |
| 1:2:1492:U:O2' | 1:2:1494:A:OP1 | 2.28 | 0.43 |
| 1:2:1606:G:H2' | 1:2:1607:G:H8 | 1.83 | 0.43 |
| 1:2:1704:G:N2 | 1:2:1818:A:N1 | 2.50 | 0.43 |
| 1:2:1860:A:H5' | 29:b:95:ARG:HD2 | 2.00 | 0.43 |
| 11:J:75:ILE:HD11 | 11:J:78:ARG:HB2 | 2.00 | 0.43 |
| 23:V:62:ARG:O | 23:V:65:TYR:HB3 | 2.18 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 26:Y:34:ILE:H | 26:Y:34:ILE:HD12 | 1.83 | 0.43 |
| 1:2:17:C:O2' | 1:2:1190:A:N1 | 2.51 | 0.43 |
| 1:2:369:C:P | 12:K:181:GLN:HE22 | 2.42 | 0.43 |
| 1:2:882:A:H61 | 1:2:896:C:H42 | 1.66 | 0.43 |
| 1:2:962:U:H2' | 1:2:963:C:C6 | 2.53 | 0.43 |
| 1:2:1800:A:H2' | 1:2:1801:C:C6 | 2.53 | 0.43 |
| 8:G:8:HIS:O | 8:G:30:ARG:NH1 | 2.51 | 0.43 |
| 15:N:147:LYS:HD3 | 15:N:153:LYS:HB2 | 2.00 | 0.43 |
| 26:Y:26:LEU:HA | 26:Y:60:LYS:NZ | 2.33 | 0.43 |
| 37:l:4:LYS:H | 37:l:4:LYS:HG2 | 1.56 | 0.43 |
| 1:2:237:G:H2' | 1:2:238:G:O4' | 2.18 | 0.43 |
| 1:2:285:U:O4 | 15:N:14:PRO:HG2 | 2.18 | 0.43 |
| 1:2:370:G:H1' | 12:K:5:ARG:NE | 2.33 | 0.43 |
| 1:2:533:C:H2' | 1:2:534:G:O4' | 2.18 | 0.43 |
| 1:2:646:G:H5' | 1:2:652:G:N2 | 2.32 | 0.43 |
| 1:2:820:C:H5' | 13:L:147:PHE:CD1 | 2.53 | 0.43 |
| 4:C:108:PHE:HB2 | 4:C:136:GLU:CD | 2.42 | 0.43 |
| 5:D:86:LEU:HD12 | 5:D:98:THR:HG23 | 2.00 | 0.43 |
| 7:F:94:ARG:HB2 | 7:F:125:PHE:HZ | 1.82 | 0.43 |
| 7:F:132:LYS:NZ | 7:F:193:ASP:HB2 | 2.33 | 0.43 |
| 8:G:129:ILE:HA | 8:G:138:HIS:O | 2.18 | 0.43 |
| 11:J:41:ARG:HD2 | 11:J:41:ARG:HA | 1.79 | 0.43 |
| 11:J:126:HIS:O | 11:J:130:LEU:HD23 | 2.18 | 0.43 |
| 12:K:67:TRP:HA | 12:K:189:VAL:HG22 | 2.00 | 0.43 |
| 15:N:35:ARG:HG2 | 15:N:51:ILE:O | 2.18 | 0.43 |
| 17:P:33:VAL:O | 17:P:36:GLN:HG2 | 2.18 | 0.43 |
| 19:R:111:MET:HE1 | 22:U:117:ILE:HG23 | 2.00 | 0.43 |
| 28:a:15:ASN:OD1 | 28:a:18:LEU:HB2 | 2.18 | 0.43 |
| 32:e:33:LYS:O | 32:e:36:LEU:HD23 | 2.18 | 0.43 |
| 39:z:535:A:O2' | 39:z:540:C:N3 | 2.43 | 0.43 |
| 39:z:541:C:H2' | 39:z:542:A:H8 | 1.83 | 0.43 |
| 1:2:1279:C:H4' | 33:f:99:LYS:HZ2 | 1.83 | 0.43 |
| 1:2:1369:C:H5' | 21:T:7:LYS:HD3 | 2.00 | 0.43 |
| 1:2:1851:G:H2' | 1:2:1852:G:H8 | 1.83 | 0.43 |
| 4:C:111:GLN:HG2 | 6:E:74:LYS:HB3 | 1.99 | 0.43 |
| 6:E:226:PHE:HA | 6:E:229:ILE:HG12 | 2.00 | 0.43 |
| 11:J:142:LYS:HD3 | 17:P:18:TYR:OH | 2.18 | 0.43 |
| 19:R:10:ARG:HG3 | 19:R:22:LEU:HD22 | 2.01 | 0.43 |
| 20:S:54:PRO:HG3 | 20:S:85:ARG:HG3 | 2.00 | 0.43 |
| 23:V:11:GLN:HA | 23:V:14:PHE:HB3 | 2.00 | 0.43 |
| 30:c:18:LYS:C | 30:c:23:ARG:HH12 | 2.18 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-----------------|-------------------|--------------------------|-------------------|
| 34:g:36:ARG:HG2 | 34:g:65:PHE:HB3 | 1.99 | 0.43 |
| 1:2:74:G:H2' | 1:2:75:G:N3 | 2.32 | 0.43 |
| 1:2:84:A:N3 | 1:2:150:A:O2' | 2.49 | 0.43 |
| 1:2:124:U:N3 | 1:2:330:C:N4 | 2.37 | 0.43 |
| 1:2:1386:U:H2' | 1:2:1387:C:H6 | 1.84 | 0.43 |
| 1:2:1436:C:H2' | 1:2:1437:U:C6 | 2.54 | 0.43 |
| 1:2:1460:C:H2' | 1:2:1461:A:C8 | 2.54 | 0.43 |
| 1:2:1587:C:H5'' | 9:H:91:ARG:NH1 | 2.31 | 0.43 |
| 1:2:1637:U:H2' | 1:2:1638:U:H6 | 1.83 | 0.43 |
| 1:2:1803:A:H2' | 1:2:1804:U:H6 | 1.84 | 0.43 |
| 4:C:22:GLY:HA2 | 4:C:24:HIS:CE1 | 2.54 | 0.43 |
| 5:D:82:ARG:HG3 | 5:D:82:ARG:O | 2.19 | 0.43 |
| 22:U:6:PRO:N | 35:h:50:PHE:HB2 | 2.32 | 0.43 |
| 23:V:42:HIS:HB3 | 23:V:93:SER:OG | 2.17 | 0.43 |
| 26:Y:90:GLN:O | 26:Y:94:LEU:HB3 | 2.17 | 0.43 |
| 34:g:19:THR:O | 34:g:288:SER:HB3 | 2.18 | 0.43 |
| 1:2:42:A:O2' | 1:2:98:C:OP1 | 2.25 | 0.43 |
| 1:2:193:C:H2' | 1:2:194:C:C6 | 2.53 | 0.43 |
| 1:2:1575:A:C8 | 24:W:56:MET:HE1 | 2.53 | 0.43 |
| 1:2:1833:U:H2' | 1:2:1834:U:C6 | 2.53 | 0.43 |
| 5:D:227:LYS:O | 5:D:231:LEU:HG | 2.19 | 0.43 |
| 6:E:93:LYS:HD2 | 6:E:95:MET:HE3 | 1.99 | 0.43 |
| 8:G:155:LYS:HA | 8:G:155:LYS:HD3 | 1.90 | 0.43 |
| 9:H:106:GLU:OE2 | 9:H:110:GLN:HB2 | 2.18 | 0.43 |
| 10:I:98:ARG:HD2 | 10:I:106:LEU:HD21 | 1.99 | 0.43 |
| 10:I:131:ARG:H | 10:I:131:ARG:HG2 | 1.67 | 0.43 |
| 13:L:21:GLU:OE1 | 13:L:24:ARG:N | 2.39 | 0.43 |
| 13:L:109:ARG:HE | 13:L:111:GLN:HE22 | 1.65 | 0.43 |
| 15:N:5:GLN:OE1 | 15:N:10:TYR:HA | 2.18 | 0.43 |
| 20:S:61:GLU:H | 20:S:61:GLU:CD | 2.27 | 0.43 |
| 23:V:74:SER:O | 23:V:77:LYS:HG3 | 2.19 | 0.43 |
| 23:V:101:ARG:O | 23:V:105:GLN:OE1 | 2.36 | 0.43 |
| 35:h:82:SER:HA | 35:h:85:ARG:NE | 2.34 | 0.43 |
| 38:y:38:C:H3' | 38:y:39:C:C6 | 2.54 | 0.43 |
| 1:2:291:U:C2 | 1:2:292:A:N7 | 2.87 | 0.43 |
| 1:2:330:C:H2' | 1:2:331:C:C6 | 2.53 | 0.43 |
| 1:2:442:G:H2' | 1:2:443:C:H6 | 1.83 | 0.43 |
| 1:2:952:G:H2' | 1:2:953:A:C8 | 2.53 | 0.43 |
| 1:2:1262:C:C4 | 1:2:1512:G:N2 | 2.87 | 0.43 |
| 1:2:1394:G:H2' | 1:2:1395:C:C6 | 2.54 | 0.43 |
| 1:2:1552:C:OP1 | 32:e:13:LYS:NZ | 2.52 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-----------------|-------------------|--------------------------|-------------------|
| 1:2:1734:C:H2' | 1:2:1735:C:H6 | 1.84 | 0.43 |
| 1:2:1766:C:H2' | 1:2:1767:C:C6 | 2.53 | 0.43 |
| 4:C:33:GLN:NE2 | 25:X:63:GLY:O | 2.51 | 0.43 |
| 7:F:164:VAL:O | 7:F:168:VAL:HG22 | 2.18 | 0.43 |
| 8:G:64:ILE:HG23 | 28:a:17:LEU:HD21 | 2.01 | 0.43 |
| 10:I:210:ALA:O | 10:I:213:LEU:HG | 2.19 | 0.43 |
| 16:O:19:GLN:HG2 | 16:O:88:TRP:CH2 | 2.54 | 0.43 |
| 17:P:98:VAL:O | 17:P:102:LEU:HD23 | 2.18 | 0.43 |
| 22:U:23:ARG:CZ | 35:h:48:VAL:HG22 | 2.48 | 0.43 |
| 35:h:97:ILE:CG2 | 35:h:109:TYR:HB3 | 2.48 | 0.43 |
| 38:y:10:G:N7 | 38:y:45:G:N1 | 2.67 | 0.43 |
| 1:2:384:G:H2' | 1:2:385:G:H8 | 1.83 | 0.43 |
| 1:2:487:C:H2' | 1:2:488:C:C6 | 2.53 | 0.43 |
| 1:2:1088:G:C2 | 1:2:1154:G:C6 | 3.07 | 0.43 |
| 1:2:1165:G:H3' | 1:2:1166:A:H8 | 1.83 | 0.43 |
| 1:2:1372:A:OP2 | 21:T:67:ARG:NH1 | 2.49 | 0.43 |
| 1:2:1403:U:H2' | 1:2:1404:U:C6 | 2.54 | 0.43 |
| 1:2:1535:G:C2 | 1:2:1536:G:C5 | 3.06 | 0.43 |
| 1:2:1641:C:P | 20:S:138:ARG:HE | 2.42 | 0.43 |
| 4:C:11:LYS:NZ | 4:C:184:ARG:HH12 | 2.17 | 0.43 |
| 4:C:124:VAL:O | 4:C:146:ALA:HA | 2.19 | 0.43 |
| 6:E:118:TYR:HE1 | 6:E:201:MET:HA | 1.84 | 0.43 |
| 6:E:233:TYR:HE1 | 25:X:14:PRO:HB3 | 1.83 | 0.43 |
| 8:G:86:PHE:HE1 | 8:G:102:ILE:HA | 1.84 | 0.43 |
| 9:H:122:ARG:HA | 9:H:146:ARG:HH22 | 1.83 | 0.43 |
| 14:M:7:ASN:O | 14:M:11:ILE:HG12 | 2.19 | 0.43 |
| 17:P:96:VAL:O | 17:P:100:LYS:HG3 | 2.18 | 0.43 |
| 29:b:38:LYS:O | 29:b:71:LEU:HG | 2.18 | 0.43 |
| 37:l:15:ARG:O | 37:l:18:ARG:HG2 | 2.18 | 0.43 |
| 1:2:29:G:H2' | 1:2:30:C:H6 | 1.84 | 0.43 |
| 1:2:545:A:H2' | 1:2:546:U:C6 | 2.54 | 0.43 |
| 1:2:952:G:H4' | 18:Q:60:MET:HE1 | 2.00 | 0.43 |
| 1:2:966:G:H5'' | 1:2:967:G:C8 | 2.54 | 0.43 |
| 1:2:1028:C:H5'' | 17:P:109:LYS:NZ | 2.32 | 0.43 |
| 1:2:1273:C:H2' | 1:2:1274:A:H8 | 1.82 | 0.43 |
| 1:2:1413:C:H2' | 1:2:1415:C:H6 | 1.84 | 0.43 |
| 1:2:1557:C:H5'' | 23:V:71:GLY:HA3 | 2.00 | 0.43 |
| 6:E:239:ASP:OD2 | 25:X:1:MET:HE2 | 2.19 | 0.43 |
| 8:G:71:LYS:NZ | 8:G:93:ASP:OD2 | 2.51 | 0.43 |
| 9:H:145:ARG:HA | 9:H:148:ASN:HD22 | 1.82 | 0.43 |
| 9:H:192:LYS:HA | 9:H:195:GLU:OE1 | 2.19 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 11:J:49:LYS:HG3 | 11:J:63:PHE:HE1 | 1.84 | 0.43 |
| 15:N:92:TYR:CZ | 15:N:105:ARG:HB2 | 2.54 | 0.43 |
| 22:U:23:ARG:HH12 | 35:h:47:LEU:HA | 1.82 | 0.43 |
| 22:U:70:ILE:HA | 22:U:77:TYR:HE2 | 1.83 | 0.43 |
| 26:Y:70:ASN:C | 26:Y:70:ASN:HD22 | 2.26 | 0.43 |
| 34:g:20:GLN:OE1 | 34:g:69:VAL:N | 2.36 | 0.43 |
| 35:h:72:VAL:O | 35:h:76:ARG:HB2 | 2.19 | 0.43 |
| 1:2:4:C:H4' | 6:E:192:ALA:HB2 | 2.01 | 0.43 |
| 1:2:50:A:H2' | 1:2:51:U:O4' | 2.19 | 0.43 |
| 1:2:293:A:H1' | 12:K:73:THR:CG2 | 2.48 | 0.43 |
| 1:2:516:A:H5'' | 36:i:109:ARG:HH12 | 1.83 | 0.43 |
| 1:2:868:A:O2' | 1:2:869:G:O5' | 2.37 | 0.43 |
| 1:2:929:G:N3 | 1:2:996:C:H2' | 2.34 | 0.43 |
| 1:2:938:G:H2' | 1:2:939:U:C6 | 2.54 | 0.43 |
| 1:2:957:G:C4 | 39:z:831:A:C6 | 3.04 | 0.43 |
| 1:2:992:A:H2' | 1:2:993:A:C8 | 2.54 | 0.43 |
| 1:2:1013:U:OP1 | 17:P:14:SER:OG | 2.23 | 0.43 |
| 1:2:1584:A:H2' | 1:2:1585:C:O4' | 2.19 | 0.43 |
| 1:2:1748:G:H2' | 1:2:1749:C:C6 | 2.54 | 0.43 |
| 4:C:69:GLU:OE2 | 6:E:252:GLN:HA | 2.19 | 0.43 |
| 5:D:112:SER:HA | 29:b:68:TYR:HE2 | 1.83 | 0.43 |
| 8:G:9:LEU:HG | 8:G:30:ARG:HG3 | 2.01 | 0.43 |
| 9:H:77:MET:HA | 9:H:77:MET:HE3 | 2.01 | 0.43 |
| 15:N:77:VAL:HG13 | 15:N:86:ILE:HD11 | 2.00 | 0.43 |
| 17:P:4:MET:HE3 | 17:P:124:ARG:HD3 | 2.00 | 0.43 |
| 19:R:18:ARG:H | 22:U:91:LYS:CA | 2.27 | 0.43 |
| 21:T:38:ILE:HD12 | 21:T:38:ILE:HA | 1.91 | 0.43 |
| 33:f:82:LYS:HE2 | 33:f:82:LYS:HB2 | 1.87 | 0.43 |
| 34:g:298:LEU:HD23 | 34:g:298:LEU:H | 1.83 | 0.43 |
| 1:2:92:A:N6 | 1:2:434:G:H1' | 2.34 | 0.42 |
| 1:2:517:C:H2' | 1:2:518:A:C8 | 2.53 | 0.42 |
| 1:2:532:U:H2' | 1:2:533:C:O4' | 2.19 | 0.42 |
| 1:2:826:A:C6 | 1:2:841:G:N1 | 2.87 | 0.42 |
| 1:2:1119:C:H4' | 5:D:149:GLN:HB2 | 2.00 | 0.42 |
| 1:2:1281:G:N2 | 33:f:104:LYS:HG2 | 2.30 | 0.42 |
| 1:2:1289:A:N6 | 1:2:1298:G:O6 | 2.52 | 0.42 |
| 1:2:1386:U:H2' | 1:2:1387:C:C6 | 2.53 | 0.42 |
| 5:D:167:LYS:NZ | 5:D:201:CYS:HA | 2.34 | 0.42 |
| 7:F:29:LEU:O | 7:F:34:TYR:HB2 | 2.19 | 0.42 |
| 7:F:167:TYR:OH | 7:F:201:LYS:O | 2.30 | 0.42 |
| 8:G:11:ARG:NE | 8:G:27:PHE:O | 2.49 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 12:K:12:ARG:HG2 | 12:K:13:LYS:N | 2.34 | 0.42 |
| 12:K:22:HIS:HB2 | 12:K:25:ARG:HH12 | 1.84 | 0.42 |
| 15:N:30:LYS:HG3 | 15:N:31:GLU:OE1 | 2.19 | 0.42 |
| 16:O:32:ALA:HB3 | 16:O:110:VAL:HB | 2.01 | 0.42 |
| 18:Q:124:MET:HA | 18:Q:124:MET:HE3 | 2.01 | 0.42 |
| 19:R:127:LYS:HD2 | 19:R:127:LYS:HA | 1.64 | 0.42 |
| 22:U:51:ASP:OD1 | 22:U:52:LEU:N | 2.52 | 0.42 |
| 26:Y:36:ARG:NE | 26:Y:36:ARG:HA | 2.34 | 0.42 |
| 26:Y:82:GLN:O | 26:Y:84:LYS:HG2 | 2.19 | 0.42 |
| 37:l:9:ARG:HG2 | 37:l:10:MET:HE2 | 2.01 | 0.42 |
| 38:y:9:G:N2 | 38:y:21:G:O6 | 2.51 | 0.42 |
| 1:2:56:G:C6 | 1:2:90:G:N1 | 2.87 | 0.42 |
| 1:2:370:G:H5'' | 12:K:31:ARG:NH2 | 2.31 | 0.42 |
| 1:2:859:U:H2' | 1:2:860:A:H8 | 1.84 | 0.42 |
| 1:2:967:G:H4' | 1:2:968:A:H8 | 1.84 | 0.42 |
| 1:2:1156:U:P | 27:Z:5:ARG:H | 2.42 | 0.42 |
| 1:2:1217:G:H2' | 1:2:1218:G:H8 | 1.84 | 0.42 |
| 1:2:1380:C:H5'' | 7:F:157:MET:O | 2.19 | 0.42 |
| 1:2:1644:U:OP1 | 20:S:127:CYS:HA | 2.19 | 0.42 |
| 5:D:171:ILE:HG23 | 5:D:174:ARG:NH2 | 2.34 | 0.42 |
| 13:L:32:ILE:HD11 | 13:L:37:LEU:HB2 | 2.01 | 0.42 |
| 13:L:93:LYS:HD2 | 13:L:93:LYS:HA | 1.89 | 0.42 |
| 17:P:20:ARG:HG3 | 17:P:65:PHE:CD1 | 2.54 | 0.42 |
| 26:Y:34:ILE:O | 26:Y:38:LEU:HD23 | 2.18 | 0.42 |
| 26:Y:75:ILE:HG12 | 26:Y:127:GLY:HA2 | 2.00 | 0.42 |
| 34:g:271:LYS:HD2 | 34:g:271:LYS:HA | 1.93 | 0.42 |
| 35:h:62:VAL:O | 35:h:111:ARG:HG3 | 2.19 | 0.42 |
| 1:2:1307:C:H3' | 1:2:1308:G:H8 | 1.83 | 0.42 |
| 1:2:1537:C:OP1 | 23:V:63:HIS:NE2 | 2.52 | 0.42 |
| 4:C:14:ASP:HA | 4:C:17:LYS:HZ2 | 1.84 | 0.42 |
| 5:D:162:ARG:HG2 | 5:D:165:ARG:HH21 | 1.84 | 0.42 |
| 6:E:233:TYR:CE1 | 25:X:14:PRO:HB3 | 2.54 | 0.42 |
| 7:F:67:ARG:NE | 14:M:93:THR:HA | 2.34 | 0.42 |
| 9:H:92:ILE:HD12 | 9:H:169:ILE:HG21 | 2.01 | 0.42 |
| 10:I:48:TYR:CD1 | 10:I:113:ILE:HD11 | 2.55 | 0.42 |
| 13:L:42:GLU:OE1 | 13:L:42:GLU:N | 2.46 | 0.42 |
| 19:R:123:TYR:OH | 22:U:124:ARG:NH1 | 2.49 | 0.42 |
| 20:S:58:LEU:O | 20:S:62:ARG:NH1 | 2.52 | 0.42 |
| 26:Y:80:ASP:OD1 | 26:Y:81:VAL:N | 2.52 | 0.42 |
| 26:Y:104:LEU:HB3 | 26:Y:125:ILE:HD12 | 2.00 | 0.42 |
| 30:c:30:SER:HB2 | 30:c:48:SER:OG | 2.18 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 34:g:90:TRP:N | 34:g:90:TRP:CD1 | 2.87 | 0.42 |
| 36:i:114:ARG:HD2 | 36:i:114:ARG:HA | 1.81 | 0.42 |
| 38:y:64:C:H2' | 38:y:65:C:C6 | 2.55 | 0.42 |
| 39:z:547:G:C2 | 39:z:549:G:H8 | 2.36 | 0.42 |
| 1:2:499:G:H2' | 1:2:500:G:C8 | 2.55 | 0.42 |
| 1:2:637:U:H2' | 1:2:638:A:H8 | 1.84 | 0.42 |
| 1:2:924:G:C6 | 1:2:1010:G:C6 | 3.08 | 0.42 |
| 1:2:1469:G:H22 | 1:2:1471:G:H5'' | 1.85 | 0.42 |
| 1:2:1535:G:C6 | 1:2:1589:A:N1 | 2.88 | 0.42 |
| 1:2:1612:G:H1 | 19:R:40:ARG:NH2 | 2.18 | 0.42 |
| 1:2:1618:A:C8 | 22:U:132:ARG:HG2 | 2.55 | 0.42 |
| 1:2:1643:G:HO2' | 1:2:1669:G:H1 | 1.67 | 0.42 |
| 1:2:1687:U:H5'' | 29:b:89:ARG:NH1 | 2.34 | 0.42 |
| 1:2:1784:A:H2' | 1:2:1785:A:O4' | 2.19 | 0.42 |
| 4:C:17:LYS:HE2 | 4:C:176:TRP:HZ3 | 1.84 | 0.42 |
| 9:H:55:ARG:HH22 | 20:S:124:PRO:HD2 | 1.85 | 0.42 |
| 9:H:69:VAL:O | 9:H:73:THR:HG23 | 2.18 | 0.42 |
| 11:J:118:ARG:O | 11:J:121:THR:HG22 | 2.19 | 0.42 |
| 12:K:159:SER:HB3 | 12:K:161:LEU:HG | 2.01 | 0.42 |
| 13:L:163:SER:H | 13:L:168:GLY:HA3 | 1.84 | 0.42 |
| 16:O:26:LEU:HD23 | 16:O:93:LYS:HB3 | 2.01 | 0.42 |
| 27:Z:48:LYS:O | 27:Z:74:LEU:HD12 | 2.19 | 0.42 |
| 33:f:92:LYS:HE2 | 33:f:93:HIS:O | 2.20 | 0.42 |
| 34:g:6:THR:CG2 | 34:g:8:ARG:HH12 | 2.26 | 0.42 |
| 38:y:48:G:H2' | 38:y:49:A:H8 | 1.84 | 0.42 |
| 38:y:51:G:C6 | 38:y:62:A:N1 | 2.88 | 0.42 |
| 1:2:15:U:H2' | 1:2:16:G:O4' | 2.19 | 0.42 |
| 1:2:158:A:H2 | 1:2:453:C:H1' | 1.83 | 0.42 |
| 1:2:368:U:H2' | 1:2:369:C:O4' | 2.18 | 0.42 |
| 1:2:800:U:H2' | 1:2:801:U:C6 | 2.54 | 0.42 |
| 1:2:911:G:N2 | 1:2:911:G:OP2 | 2.53 | 0.42 |
| 1:2:942:U:H2' | 1:2:943:G:C8 | 2.54 | 0.42 |
| 1:2:1198:U:H4' | 6:E:99:LYS:HE2 | 2.00 | 0.42 |
| 1:2:1440:U:H2' | 1:2:1441:U:C6 | 2.54 | 0.42 |
| 1:2:1537:C:OP1 | 23:V:59:SER:OG | 2.37 | 0.42 |
| 1:2:1670:A:N7 | 1:2:1671:U:C5 | 2.87 | 0.42 |
| 4:C:62:ALA:O | 4:C:66:VAL:HG23 | 2.20 | 0.42 |
| 7:F:216:GLU:OE2 | 21:T:20:TYR:HA | 2.19 | 0.42 |
| 8:G:36:HIS:CE1 | 8:G:85:GLY:HA3 | 2.54 | 0.42 |
| 9:H:140:ASP:OD1 | 9:H:141:VAL:N | 2.45 | 0.42 |
| 13:L:137:VAL:C | 13:L:139:LYS:H | 2.27 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 18:Q:50:LYS:HE3 | 18:Q:50:LYS:HB3 | 1.76 | 0.42 |
| 24:W:81:GLN:C | 24:W:82:MET:HE2 | 2.44 | 0.42 |
| 38:y:63:U:H2' | 38:y:64:C:C6 | 2.53 | 0.42 |
| 39:z:831:A:H5'' | 39:z:832:A:C5 | 2.54 | 0.42 |
| 1:2:163:U:O3' | 10:I:83:CYS:HA | 2.19 | 0.42 |
| 1:2:471:C:H2' | 1:2:472:G:C8 | 2.55 | 0.42 |
| 1:2:580:A:H4' | 36:i:98:LYS:HE2 | 2.02 | 0.42 |
| 1:2:827:G:H2' | 1:2:828:G:C8 | 2.54 | 0.42 |
| 1:2:849:C:C5 | 1:2:850:A:C8 | 3.08 | 0.42 |
| 1:2:895:U:H2' | 1:2:896:C:O4' | 2.19 | 0.42 |
| 1:2:1067:G:H2' | 1:2:1068:U:O4' | 2.19 | 0.42 |
| 1:2:1102:C:OP1 | 30:c:70:LYS:NZ | 2.36 | 0.42 |
| 1:2:1175:G:N2 | 1:2:1178:A:OP2 | 2.39 | 0.42 |
| 1:2:1859:C:N3 | 29:b:6:ARG:N | 2.55 | 0.42 |
| 4:C:8:LEU:HD11 | 25:X:39:VAL:HG21 | 2.01 | 0.42 |
| 4:C:80:ARG:HH12 | 4:C:165:ASN:C | 2.28 | 0.42 |
| 6:E:163:HIS:CG | 6:E:185:ARG:HG2 | 2.54 | 0.42 |
| 6:E:195:PRO:HA | 6:E:198:LEU:HG | 2.00 | 0.42 |
| 10:I:192:ILE:HD12 | 10:I:192:ILE:HA | 1.96 | 0.42 |
| 1:2:191:C:H2' | 1:2:192:U:C6 | 2.53 | 0.42 |
| 1:2:469:C:H2' | 1:2:470:G:O4' | 2.20 | 0.42 |
| 1:2:528:U:H2' | 1:2:529:C:H6 | 1.84 | 0.42 |
| 1:2:1721:G:C6 | 1:2:1803:A:C6 | 3.07 | 0.42 |
| 10:I:229:ALA:HA | 10:I:232:ARG:HG3 | 2.02 | 0.42 |
| 11:J:144:ILE:HG12 | 26:Y:52:ILE:HG22 | 2.02 | 0.42 |
| 12:K:190:LEU:C | 12:K:195:LEU:HD21 | 2.45 | 0.42 |
| 14:M:37:ASP:OD1 | 14:M:38:LYS:N | 2.53 | 0.42 |
| 20:S:9:SER:HA | 20:S:25:CYS:O | 2.19 | 0.42 |
| 25:X:58:ALA:O | 25:X:62:MET:HG3 | 2.20 | 0.42 |
| 34:g:234:ASP:OD1 | 34:g:235:ILE:N | 2.52 | 0.42 |
| 37:l:3:ALA:HA | 37:l:6:ARG:NH1 | 2.35 | 0.42 |
| 38:y:51:G:C6 | 38:y:62:A:C6 | 3.08 | 0.42 |
| 1:2:342:U:C5' | 15:N:139:ARG:HG2 | 2.49 | 0.42 |
| 1:2:343:C:H2' | 1:2:344:U:C6 | 2.54 | 0.42 |
| 1:2:553:G:N3 | 1:2:554:A:C8 | 2.88 | 0.42 |
| 1:2:637:U:H2' | 1:2:638:A:C8 | 2.55 | 0.42 |
| 1:2:674:G:H2' | 1:2:675:A:C8 | 2.55 | 0.42 |
| 1:2:821:A:H2' | 1:2:822:A:H8 | 1.85 | 0.42 |
| 1:2:1187:C:H2' | 1:2:1188:U:C6 | 2.54 | 0.42 |
| 1:2:1353:A:H5'' | 6:E:97:VAL:HG11 | 2.01 | 0.42 |
| 1:2:1469:G:N2 | 1:2:1471:G:OP2 | 2.53 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 3:B:283:ILE:O | 3:B:334:GLY:N | 2.52 | 0.42 |
| 5:D:225:LEU:HB3 | 5:D:229:MET:HE1 | 2.01 | 0.42 |
| 9:H:99:ILE:O | 9:H:103:LEU:HG | 2.20 | 0.42 |
| 10:I:39:ASP:OD1 | 10:I:40:ALA:N | 2.53 | 0.42 |
| 12:K:129:LEU:O | 12:K:134:GLU:HB2 | 2.20 | 0.42 |
| 17:P:126:ALA:HB1 | 17:P:139:TRP:HZ3 | 1.85 | 0.42 |
| 19:R:10:ARG:HA | 19:R:10:ARG:HD2 | 1.91 | 0.42 |
| 21:T:122:PRO:HA | 21:T:123:THR:HA | 1.85 | 0.42 |
| 25:X:55:ILE:HG23 | 25:X:59:ILE:HD11 | 2.01 | 0.42 |
| 32:e:26:ASN:O | 32:e:27:ARG:HG3 | 2.20 | 0.42 |
| 38:y:49:A:H4' | 39:z:551:C:O3' | 2.19 | 0.42 |
| 1:2:96:C:H1' | 1:2:464:G:H5' | 2.02 | 0.42 |
| 1:2:1089:A:H2' | 1:2:1090:C:C6 | 2.55 | 0.42 |
| 8:G:120:LYS:HA | 8:G:120:LYS:HD2 | 1.91 | 0.42 |
| 8:G:127:ARG:N | 8:G:140:VAL:O | 2.40 | 0.42 |
| 19:R:82:ASP:OD1 | 19:R:82:ASP:N | 2.50 | 0.42 |
| 21:T:18:GLU:HG2 | 21:T:70:SER:H | 1.85 | 0.42 |
| 24:W:69:PRO:HA | 32:e:40:ARG:NH1 | 2.35 | 0.42 |
| 30:c:79:PHE:O | 30:c:80:ARG:NH1 | 2.42 | 0.42 |
| 1:2:442:G:H2' | 1:2:443:C:C6 | 2.55 | 0.42 |
| 1:2:645:A:H1' | 1:2:648:U:OP1 | 2.19 | 0.42 |
| 1:2:1174:U:H2' | 1:2:1175:G:O4' | 2.20 | 0.42 |
| 1:2:1412:C:O2' | 23:V:132:ASP:OD2 | 2.38 | 0.42 |
| 1:2:1450:A:H5'' | 21:T:3:ARG:NH1 | 2.32 | 0.42 |
| 1:2:1858:U:O2' | 1:2:1860:A:N7 | 2.53 | 0.42 |
| 5:D:212:VAL:HG13 | 5:D:212:VAL:O | 2.19 | 0.42 |
| 6:E:155:TRP:C | 26:Y:98:GLN:HE22 | 2.28 | 0.42 |
| 8:G:29:PRO:HG3 | 8:G:45:ILE:HD11 | 2.01 | 0.42 |
| 8:G:40:GLU:OE2 | 8:G:103:TYR:OH | 2.20 | 0.42 |
| 8:G:124:CYS:HB3 | 8:G:141:THR:HB | 2.02 | 0.42 |
| 12:K:113:TYR:OH | 12:K:158:ILE:HD11 | 2.20 | 0.42 |
| 14:M:5:LYS:O | 14:M:8:ARG:HB2 | 2.20 | 0.42 |
| 18:Q:31:CYS:HB2 | 18:Q:95:ILE:HA | 2.02 | 0.42 |
| 18:Q:97:LEU:O | 18:Q:132:VAL:N | 2.24 | 0.42 |
| 19:R:123:TYR:CD2 | 19:R:125:PRO:HA | 2.55 | 0.42 |
| 21:T:24:LEU:O | 34:g:212:LYS:NZ | 2.53 | 0.42 |
| 1:2:152:U:H2' | 1:2:153:G:O4' | 2.20 | 0.41 |
| 1:2:161:U:C4 | 28:a:115:LYS:HB3 | 2.55 | 0.41 |
| 1:2:412:U:H2' | 1:2:413:U:H6 | 1.82 | 0.41 |
| 1:2:525:G:H2' | 1:2:526:A:H8 | 1.85 | 0.41 |
| 1:2:643:A:H2' | 1:2:644:A:O4' | 2.20 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-----------------|------------------|--------------------------|-------------------|
| 1:2:743:U:O2' | 1:2:744:C:O4' | 2.29 | 0.41 |
| 1:2:921:G:C2 | 1:2:922:A:C8 | 3.08 | 0.41 |
| 1:2:1245:C:H3' | 1:2:1246:A:C8 | 2.55 | 0.41 |
| 1:2:1252:G:H3' | 1:2:1252:G:N3 | 2.35 | 0.41 |
| 1:2:1307:C:H3' | 1:2:1308:G:C8 | 2.55 | 0.41 |
| 1:2:1347:G:H2' | 1:2:1348:G:C8 | 2.53 | 0.41 |
| 7:F:157:MET:HE3 | 7:F:158:ILE:O | 2.19 | 0.41 |
| 9:H:133:THR:OG1 | 9:H:135:ARG:NH1 | 2.53 | 0.41 |
| 14:M:3:MET:HA | 14:M:4:PRO:HD3 | 1.91 | 0.41 |
| 18:Q:61:LYS:NZ | 18:Q:76:LEU:HG | 2.35 | 0.41 |
| 22:U:130:ARG:HA | 22:U:130:ARG:HD3 | 1.78 | 0.41 |
| 23:V:62:ARG:NH1 | 23:V:66:LEU:HD11 | 2.35 | 0.41 |
| 23:V:85:ASN:HB3 | 23:V:88:MET:HB2 | 2.02 | 0.41 |
| 34:g:22:ALA:HB1 | 34:g:71:ILE:HG23 | 2.02 | 0.41 |
| 35:h:72:VAL:O | 35:h:76:ARG:HD3 | 2.19 | 0.41 |
| 1:2:79:A:O2' | 1:2:80:G:O5' | 2.38 | 0.41 |
| 1:2:370:G:H1' | 12:K:5:ARG:CZ | 2.50 | 0.41 |
| 1:2:746:C:H2' | 1:2:747:G:C8 | 2.55 | 0.41 |
| 1:2:1000:U:H2' | 1:2:1001:G:C8 | 2.54 | 0.41 |
| 1:2:1140:A:P | 1:2:1351:C:H41 | 2.43 | 0.41 |
| 1:2:1164:G:H3' | 1:2:1165:G:C8 | 2.55 | 0.41 |
| 1:2:1255:A:H62 | 1:2:1513:C:H3' | 1.85 | 0.41 |
| 1:2:1355:U:H2' | 1:2:1356:U:C6 | 2.54 | 0.41 |
| 1:2:1360:U:H3 | 1:2:1371:G:H21 | 1.68 | 0.41 |
| 1:2:1456:C:H2' | 1:2:1457:G:C8 | 2.55 | 0.41 |
| 1:2:1652:G:H2' | 1:2:1653:G:H8 | 1.84 | 0.41 |
| 1:2:1660:G:OP1 | 23:V:91:HIS:NE2 | 2.49 | 0.41 |
| 1:2:1741:U:H2' | 1:2:1742:C:C6 | 2.56 | 0.41 |
| 6:E:174:GLY:HA3 | 6:E:220:ASN:HD21 | 1.86 | 0.41 |
| 7:F:141:LYS:C | 7:F:142:LEU:HD12 | 2.44 | 0.41 |
| 8:G:249:ALA:HA | 13:L:72:PHE:HE1 | 1.85 | 0.41 |
| 9:H:82:ASN:HA | 9:H:85:LYS:HD3 | 2.02 | 0.41 |
| 10:I:45:TRP:HA | 10:I:48:TYR:HD2 | 1.84 | 0.41 |
| 12:K:40:PRO:HG2 | 12:K:59:ARG:NH1 | 2.35 | 0.41 |
| 16:O:22:LEU:HA | 16:O:25:ALA:HB3 | 2.01 | 0.41 |
| 19:R:10:ARG:NH1 | 19:R:21:ASP:HA | 2.35 | 0.41 |
| 19:R:41:GLN:OE1 | 19:R:41:GLN:N | 2.41 | 0.41 |
| 21:T:72:LYS:HD2 | 21:T:75:GLU:OE2 | 2.20 | 0.41 |
| 24:W:66:ARG:HE | 24:W:77:TRP:HE1 | 1.68 | 0.41 |
| 24:W:66:ARG:HG3 | 24:W:68:THR:H | 1.84 | 0.41 |
| 27:Z:68:LYS:NZ | 36:i:83:VAL:HA | 2.35 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 29:b:10:ARG:H | 29:b:12:LYS:NZ | 2.19 | 0.41 |
| 31:d:9:ILE:O | 31:d:10:LYS:HE2 | 2.20 | 0.41 |
| 35:h:71:ALA:O | 35:h:75:GLU:OE1 | 2.38 | 0.41 |
| 35:h:103:HIS:CG | 35:h:104:ARG:N | 2.87 | 0.41 |
| 1:2:49:C:H2' | 1:2:462:C:H41 | 1.85 | 0.41 |
| 1:2:812:A:H2' | 1:2:813:G:C8 | 2.54 | 0.41 |
| 1:2:1032:A:H4' | 1:2:1849:G:N2 | 2.34 | 0.41 |
| 1:2:1450:A:OP1 | 21:T:49:LYS:NZ | 2.44 | 0.41 |
| 1:2:1815:U:H2' | 1:2:1816:A:C8 | 2.34 | 0.41 |
| 4:C:160:ALA:N | 25:X:66:ASP:OD1 | 2.53 | 0.41 |
| 9:H:39:ILE:HG23 | 9:H:68:ILE:HG13 | 2.03 | 0.41 |
| 10:I:161:PRO:HA | 10:I:171:THR:HA | 2.03 | 0.41 |
| 18:Q:133:THR:O | 18:Q:135:ILE:N | 2.50 | 0.41 |
| 21:T:109:LEU:CD2 | 21:T:111:PHE:HB2 | 2.50 | 0.41 |
| 23:V:14:PHE:HZ | 23:V:131:LEU:HD12 | 1.84 | 0.41 |
| 26:Y:11:LEU:HB2 | 26:Y:72:CYS:SG | 2.60 | 0.41 |
| 31:d:27:CYS:SG | 31:d:45:ASN:HB3 | 2.61 | 0.41 |
| 32:e:16:GLN:OE1 | 32:e:16:GLN:N | 2.45 | 0.41 |
| 39:z:549:G:H2' | 39:z:550:A:C8 | 2.56 | 0.41 |
| 1:2:357:U:OP1 | 12:K:11:ARG:NH2 | 2.53 | 0.41 |
| 1:2:376:C:H1' | 12:K:5:ARG:CG | 2.50 | 0.41 |
| 1:2:573:A:N3 | 1:2:573:A:H2' | 2.35 | 0.41 |
| 1:2:604:G:N2 | 27:Z:62:PRO:O | 2.53 | 0.41 |
| 18:Q:34:PHE:O | 18:Q:41:PHE:N | 2.36 | 0.41 |
| 21:T:25:GLY:O | 21:T:58:MET:HG2 | 2.19 | 0.41 |
| 28:a:12:PHE:CZ | 28:a:21:LYS:HD2 | 2.55 | 0.41 |
| 28:a:102:THR:OG1 | 28:a:103:SER:HA | 2.20 | 0.41 |
| 38:y:42:G:H2' | 38:y:43:A:C8 | 2.56 | 0.41 |
| 39:z:582:U:H2' | 39:z:584:A:N7 | 2.35 | 0.41 |
| 1:2:294:C:H2' | 1:2:295:C:C6 | 2.55 | 0.41 |
| 1:2:342:U:H5' | 15:N:139:ARG:HG2 | 2.01 | 0.41 |
| 1:2:565:A:H2' | 1:2:566:A:O4' | 2.21 | 0.41 |
| 1:2:839:C:H1' | 8:G:263:GLY:H | 1.84 | 0.41 |
| 1:2:940:A:H1' | 18:Q:139:SER:OG | 2.20 | 0.41 |
| 1:2:957:G:N1 | 39:z:831:A:N1 | 2.65 | 0.41 |
| 1:2:1123:C:C4' | 30:c:17:ARG:HH22 | 2.34 | 0.41 |
| 1:2:1135:C:H5 | 1:2:1145:A:H62 | 1.68 | 0.41 |
| 1:2:1223:G:H1' | 1:2:1633:G:H1' | 2.01 | 0.41 |
| 1:2:1280:A:H2 | 16:O:33:ARG:HE | 1.68 | 0.41 |
| 1:2:1453:U:H2' | 1:2:1454:G:H8 | 1.84 | 0.41 |
| 1:2:1637:U:H2' | 1:2:1638:U:C6 | 2.56 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 1:2:1676:U:H2' | 1:2:1677:C:C6 | 2.55 | 0.41 |
| 1:2:1680:U:C4 | 1:2:1681:G:N7 | 2.88 | 0.41 |
| 1:2:1688:G:P | 29:b:89:ARG:HH22 | 2.43 | 0.41 |
| 1:2:1860:A:C2 | 29:b:87:ARG:NH2 | 2.89 | 0.41 |
| 4:C:71:PRO:HD3 | 4:C:186:ARG:NH2 | 2.35 | 0.41 |
| 4:C:84:GLN:HA | 4:C:87:VAL:HG22 | 2.02 | 0.41 |
| 5:D:77:ASP:OD1 | 5:D:77:ASP:N | 2.51 | 0.41 |
| 5:D:214:LYS:HB3 | 5:D:216:LYS:HZ3 | 1.86 | 0.41 |
| 7:F:121:GLY:HA2 | 7:F:124:ARG:HE | 1.84 | 0.41 |
| 9:H:28:VAL:HA | 9:H:110:GLN:HG3 | 2.03 | 0.41 |
| 13:L:135:ILE:HD13 | 13:L:159:PHE:HA | 2.02 | 0.41 |
| 19:R:88:GLU:HG3 | 19:R:89:MET:SD | 2.61 | 0.41 |
| 19:R:119:PHE:HZ | 22:U:117:ILE:HG22 | 1.85 | 0.41 |
| 21:T:41:ILE:HG12 | 21:T:47:ARG:HB2 | 2.02 | 0.41 |
| 25:X:23:ILE:HD12 | 25:X:23:ILE:HA | 1.86 | 0.41 |
| 38:y:17:G:H5' | 38:y:60:C:H5' | 2.02 | 0.41 |
| 38:y:46:U:H4' | 38:y:47:C:H5 | 1.83 | 0.41 |
| 39:z:529:A:H2' | 39:z:530:G:C8 | 2.55 | 0.41 |
| 39:z:564:G:H22 | 39:z:574:C:H1' | 1.85 | 0.41 |
| 1:2:21:U:H4' | 13:L:19:PRO:HG3 | 2.02 | 0.41 |
| 1:2:558:C:H2' | 1:2:559:A:O4' | 2.20 | 0.41 |
| 1:2:744:C:H2' | 1:2:745:U:N3 | 2.35 | 0.41 |
| 1:2:919:G:H2' | 1:2:920:G:H8 | 1.86 | 0.41 |
| 1:2:1003:C:O2' | 17:P:101:HIS:ND1 | 2.50 | 0.41 |
| 1:2:1117:G:C4 | 1:2:1118:A:C8 | 3.09 | 0.41 |
| 1:2:1132:U:N3 | 1:2:1133:U:O4 | 2.53 | 0.41 |
| 1:2:1171:G:H2' | 1:2:1172:G:H8 | 1.85 | 0.41 |
| 1:2:1193:G:H2' | 1:2:1194:G:C8 | 2.52 | 0.41 |
| 1:2:1287:A:O2' | 33:f:145:CYS:HB2 | 2.21 | 0.41 |
| 1:2:1383:G:H3' | 1:2:1384:A:H8 | 1.85 | 0.41 |
| 1:2:1592:C:N4 | 1:2:1593:G:O6 | 2.52 | 0.41 |
| 1:2:1739:G:H8 | 1:2:1739:G:OP2 | 2.04 | 0.41 |
| 1:2:1792:C:O2' | 12:K:2:GLY:N | 2.54 | 0.41 |
| 4:C:89:LYS:NZ | 4:C:202:TYR:HA | 2.35 | 0.41 |
| 6:E:124:LEU:HD11 | 6:E:222:ALA:HB1 | 2.02 | 0.41 |
| 6:E:152:ARG:HB3 | 6:E:162:PRO:HB2 | 2.02 | 0.41 |
| 6:E:158:LYS:HB2 | 25:X:4:ASN:H | 1.86 | 0.41 |
| 7:F:44:THR:O | 7:F:44:THR:HG22 | 2.21 | 0.41 |
| 7:F:75:LYS:NZ | 14:M:21:MET:HA | 2.36 | 0.41 |
| 7:F:173:ARG:HD3 | 7:F:173:ARG:HA | 1.85 | 0.41 |
| 8:G:165:GLU:N | 8:G:165:GLU:OE1 | 2.54 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 18:Q:83:GLN:HA | 18:Q:86:LYS:HE3 | 2.03 | 0.41 |
| 23:V:39:LEU:HD12 | 23:V:39:LEU:HA | 1.84 | 0.41 |
| 26:Y:78:ARG:CB | 26:Y:124:LYS:HD2 | 2.50 | 0.41 |
| 27:Z:53:GLU:OE2 | 27:Z:71:ARG:HD3 | 2.19 | 0.41 |
| 34:g:299:PHE:CE1 | 34:g:309:VAL:HG13 | 2.56 | 0.41 |
| 1:2:280:G:H2' | 1:2:281:U:C6 | 2.55 | 0.41 |
| 1:2:483:A:N6 | 1:2:499:G:H21 | 2.14 | 0.41 |
| 1:2:585:U:H2' | 1:2:586:U:H6 | 1.86 | 0.41 |
| 1:2:668:U:C2 | 1:2:1023:A:N6 | 2.87 | 0.41 |
| 1:2:677:C:O3' | 11:J:103:LYS:NZ | 2.53 | 0.41 |
| 1:2:1007:A:H5'' | 17:P:3:ARG:HH12 | 1.86 | 0.41 |
| 1:2:1603:U:H2' | 1:2:1604:C:O4' | 2.21 | 0.41 |
| 1:2:1741:U:OP1 | 10:I:31:ARG:NH2 | 2.54 | 0.41 |
| 3:B:317:LYS:O | 3:B:342:LYS:N | 2.53 | 0.41 |
| 6:E:185:ARG:C | 13:L:98:LEU:HD21 | 2.46 | 0.41 |
| 7:F:140:GLY:HA3 | 7:F:182:LEU:CD2 | 2.50 | 0.41 |
| 8:G:191:ARG:CZ | 8:G:245:ARG:HE | 2.34 | 0.41 |
| 14:M:31:LYS:HD3 | 14:M:31:LYS:HA | 1.91 | 0.41 |
| 16:O:24:THR:OG1 | 16:O:121:LYS:HE3 | 2.20 | 0.41 |
| 18:Q:95:ILE:HB | 18:Q:129:ILE:HG22 | 2.01 | 0.41 |
| 18:Q:113:GLN:H | 18:Q:113:GLN:CD | 2.26 | 0.41 |
| 22:U:10:GLN:NE2 | 22:U:57:GLY:HA2 | 2.36 | 0.41 |
| 25:X:56:CYS:SG | 25:X:59:ILE:HG12 | 2.60 | 0.41 |
| 38:y:50:U:H2' | 38:y:51:G:H8 | 1.85 | 0.41 |
| 1:2:153:G:H2' | 1:2:154:U:C6 | 2.56 | 0.41 |
| 1:2:487:C:H2' | 1:2:488:C:H6 | 1.84 | 0.41 |
| 1:2:904:A:H2' | 1:2:905:G:H8 | 1.85 | 0.41 |
| 1:2:931:G:H2' | 1:2:932:G:H8 | 1.86 | 0.41 |
| 1:2:1139:A:OP1 | 1:2:1139:A:H8 | 2.03 | 0.41 |
| 1:2:1171:G:H2' | 1:2:1172:G:C8 | 2.56 | 0.41 |
| 1:2:1283:A:C6 | 1:2:1284:U:H1' | 2.55 | 0.41 |
| 1:2:1473:U:O2 | 1:2:1473:U:H2' | 2.20 | 0.41 |
| 1:2:1481:U:H2' | 1:2:1482:A:O4' | 2.19 | 0.41 |
| 1:2:1573:U:N3 | 7:F:1:MET:O | 2.54 | 0.41 |
| 5:D:51:ARG:HG3 | 5:D:53:GLN:HE22 | 1.85 | 0.41 |
| 7:F:99:ILE:HD12 | 7:F:99:ILE:H | 1.85 | 0.41 |
| 8:G:139:LEU:HG | 8:G:150:PRO:HG3 | 2.03 | 0.41 |
| 9:H:122:ARG:HA | 9:H:146:ARG:HH12 | 1.85 | 0.41 |
| 11:J:64:VAL:O | 11:J:97:GLN:HG3 | 2.20 | 0.41 |
| 14:M:65:ARG:HH22 | 32:e:22:ARG:HA | 1.86 | 0.41 |
| 18:Q:34:PHE:HD2 | 18:Q:41:PHE:CD2 | 2.39 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 18:Q:41:PHE:HB3 | 18:Q:55:ARG:HH22 | 1.84 | 0.41 |
| 21:T:6:THR:O | 21:T:10:LYS:HG2 | 2.20 | 0.41 |
| 25:X:71:ARG:O | 25:X:74:LYS:HG2 | 2.21 | 0.41 |
| 34:g:16:GLY:HA3 | 34:g:36:ARG:C | 2.46 | 0.41 |
| 34:g:78:ALA:O | 34:g:89:LEU:HD12 | 2.20 | 0.41 |
| 39:z:546:G:H2' | 39:z:547:G:C8 | 2.56 | 0.41 |
| 1:2:3:C:C5 | 6:E:190:ILE:HD11 | 2.55 | 0.41 |
| 1:2:31:U:O2' | 1:2:633:A:N1 | 2.48 | 0.41 |
| 1:2:36:U:H2' | 1:2:37:C:C6 | 2.53 | 0.41 |
| 1:2:59:U:N3 | 1:2:62:G:OP1 | 2.54 | 0.41 |
| 1:2:149:A:C5 | 1:2:170:A:N6 | 2.89 | 0.41 |
| 1:2:479:A:H2' | 1:2:480:C:H6 | 1.85 | 0.41 |
| 1:2:842:G:C5 | 8:G:19:MET:HE1 | 2.56 | 0.41 |
| 1:2:848:G:N7 | 1:2:849:C:H1' | 2.36 | 0.41 |
| 1:2:878:U:H2' | 1:2:879:U:C6 | 2.56 | 0.41 |
| 1:2:931:G:H2' | 1:2:932:G:C8 | 2.55 | 0.41 |
| 1:2:1074:C:H2' | 1:2:1075:C:N1 | 2.36 | 0.41 |
| 1:2:1175:G:O2' | 1:2:1178:A:N6 | 2.53 | 0.41 |
| 1:2:1438:U:H1' | 20:S:11:GLN:NE2 | 2.36 | 0.41 |
| 1:2:1553:C:H2' | 1:2:1554:C:C6 | 2.56 | 0.41 |
| 1:2:1613:C:H2' | 1:2:1614:A:O4' | 2.21 | 0.41 |
| 1:2:1795:A:H2' | 1:2:1796:C:C6 | 2.56 | 0.41 |
| 4:C:103:PHE:O | 4:C:104:THR:OG1 | 2.37 | 0.41 |
| 6:E:45:TRP:CD1 | 6:E:77:GLU:HG3 | 2.55 | 0.41 |
| 6:E:180:LEU:HD12 | 6:E:207:CYS:SG | 2.60 | 0.41 |
| 8:G:93:ASP:OD1 | 8:G:93:ASP:N | 2.54 | 0.41 |
| 9:H:125:SER:HA | 9:H:138:ALA:HA | 2.02 | 0.41 |
| 10:I:38:ALA:N | 10:I:48:TYR:O | 2.53 | 0.41 |
| 10:I:50:VAL:HG11 | 10:I:111:LEU:HD12 | 2.03 | 0.41 |
| 11:J:69:LEU:O | 11:J:73:GLN:HG2 | 2.21 | 0.41 |
| 11:J:157:HIS:HA | 11:J:188:GLU:O | 2.21 | 0.41 |
| 12:K:10:LYS:HD3 | 12:K:11:ARG:N | 2.36 | 0.41 |
| 12:K:65:PHE:HE2 | 12:K:78:ILE:HG12 | 1.86 | 0.41 |
| 14:M:16:PHE:HB2 | 14:M:79:LEU:HD13 | 2.02 | 0.41 |
| 14:M:42:ASN:C | 14:M:44:HIS:H | 2.29 | 0.41 |
| 15:N:27:GLU:HA | 15:N:28:THR:HA | 1.64 | 0.41 |
| 15:N:111:VAL:HG11 | 15:N:128:VAL:HG11 | 2.03 | 0.41 |
| 17:P:128:TYR:O | 17:P:131:THR:HG22 | 2.21 | 0.41 |
| 18:Q:142:ARG:CZ | 29:b:27:ALA:HB1 | 2.51 | 0.41 |
| 20:S:128:GLU:OE1 | 20:S:128:GLU:N | 2.44 | 0.41 |
| 21:T:24:LEU:HD21 | 21:T:58:MET:HB2 | 2.02 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 21:T:37:GLU:OE2 | 34:g:150:TRP:NE1 | 2.49 | 0.41 |
| 21:T:57:LEU:O | 21:T:61:ILE:HG23 | 2.21 | 0.41 |
| 22:U:8:LYS:HD2 | 35:h:50:PHE:O | 2.20 | 0.41 |
| 23:V:52:TRP:O | 23:V:56:ARG:N | 2.48 | 0.41 |
| 25:X:18:SER:H | 25:X:54:ALA:HB3 | 1.84 | 0.41 |
| 26:Y:102:ILE:HG12 | 26:Y:113:HIS:CD2 | 2.55 | 0.41 |
| 27:Z:4:CYS:SG | 27:Z:9:THR:HG21 | 2.60 | 0.41 |
| 28:a:16:ARG:O | 28:a:19:GLN:NE2 | 2.49 | 0.41 |
| 37:l:12:ARG:HH21 | 37:l:15:ARG:NE | 2.19 | 0.41 |
| 38:y:9:G:O6 | 38:y:19:A:O2' | 2.39 | 0.41 |
| 39:z:577:G:C2 | 39:z:592:C:C2 | 3.09 | 0.41 |
| 1:2:485:U:H2' | 1:2:486:C:H6 | 1.86 | 0.41 |
| 1:2:524:G:H2' | 1:2:525:G:H8 | 1.86 | 0.41 |
| 1:2:533:C:H3' | 1:2:534:G:H8 | 1.86 | 0.41 |
| 1:2:540:C:C2 | 1:2:541:U:C5 | 3.08 | 0.41 |
| 1:2:1018:U:H6 | 17:P:128:TYR:CZ | 2.39 | 0.41 |
| 1:2:1053:C:H5' | 39:z:832:A:H4' | 2.02 | 0.41 |
| 1:2:1285:U:H5 | 33:f:97:LYS:HE2 | 1.85 | 0.41 |
| 1:2:1589:A:OP2 | 35:h:104:ARG:N | 2.54 | 0.41 |
| 1:2:1685:U:H2' | 1:2:1686:U:C6 | 2.56 | 0.41 |
| 1:2:1687:U:H5' | 29:b:87:ARG:O | 2.21 | 0.41 |
| 1:2:1825:A:H1' | 37:l:2:ARG:NH2 | 2.36 | 0.41 |
| 4:C:110:ASN:OD1 | 4:C:111:GLN:N | 2.54 | 0.41 |
| 4:C:163:CYS:SG | 4:C:164:ASN:N | 2.94 | 0.41 |
| 10:I:37:ALA:HA | 10:I:49:VAL:HA | 2.03 | 0.41 |
| 12:K:167:GLN:HG3 | 12:K:168:GLN:OE1 | 2.21 | 0.41 |
| 22:U:114:LEU:O | 22:U:118:ARG:N | 2.53 | 0.41 |
| 28:a:91:LEU:HB3 | 28:a:97:TYR:HB3 | 2.01 | 0.41 |
| 28:a:102:THR:N | 28:a:103:SER:HB2 | 2.36 | 0.41 |
| 33:f:118:ARG:HG3 | 33:f:132:MET:HE3 | 2.03 | 0.41 |
| 34:g:302:TYR:HB2 | 34:g:304:ASP:OD1 | 2.21 | 0.41 |
| 36:i:105:ARG:NH1 | 36:i:106:ALA:HB2 | 2.36 | 0.41 |
| 38:y:19:A:H1' | 38:y:58:A:C8 | 2.56 | 0.41 |
| 1:2:65:C:C6 | 10:I:174:PRO:HB3 | 2.56 | 0.40 |
| 1:2:77:A:C6 | 10:I:181:THR:HG21 | 2.56 | 0.40 |
| 1:2:105:U:H2' | 1:2:106:C:O4' | 2.21 | 0.40 |
| 1:2:217:U:H2' | 1:2:218:A:H8 | 1.86 | 0.40 |
| 1:2:321:C:HO2' | 1:2:322:G:H8 | 1.67 | 0.40 |
| 1:2:821:A:H2' | 1:2:822:A:C8 | 2.57 | 0.40 |
| 1:2:889:U:H2' | 1:2:890:G:H8 | 1.86 | 0.40 |
| 1:2:1346:U:O2' | 4:C:110:ASN:ND2 | 2.54 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|-------------------|--------------------------|-------------------|
| 8:G:259:LYS:HG3 | 8:G:259:LYS:O | 2.21 | 0.40 |
| 11:J:19:PHE:CE2 | 11:J:23:ILE:HD11 | 2.57 | 0.40 |
| 12:K:170:LYS:O | 12:K:171:LEU:HD22 | 2.21 | 0.40 |
| 21:T:58:MET:HA | 21:T:61:ILE:HG12 | 2.03 | 0.40 |
| 22:U:104:ASP:OD1 | 22:U:104:ASP:N | 2.54 | 0.40 |
| 23:V:21:PHE:O | 23:V:25:SER:OG | 2.33 | 0.40 |
| 23:V:101:ARG:HE | 23:V:102:ARG:NH2 | 2.19 | 0.40 |
| 24:W:18:HIS:O | 24:W:92:HIS:HA | 2.21 | 0.40 |
| 24:W:23:THR:HB | 24:W:113:GLU:CG | 2.51 | 0.40 |
| 28:a:111:LYS:HG2 | 28:a:115:LYS:HE2 | 2.03 | 0.40 |
| 38:y:4:A:C6 | 38:y:69:G:N1 | 2.88 | 0.40 |
| 38:y:8:U:O3' | 38:y:9:G:H3' | 2.21 | 0.40 |
| 38:y:30:G:H2' | 38:y:31:C:C6 | 2.55 | 0.40 |
| 1:2:13:C:H4' | 1:2:1352:G:H21 | 1.87 | 0.40 |
| 1:2:527:C:H2' | 1:2:528:U:C6 | 2.56 | 0.40 |
| 1:2:553:G:HO2' | 1:2:554:A:P | 2.45 | 0.40 |
| 1:2:676:U:H5'' | 26:Y:32:LYS:HZ2 | 1.85 | 0.40 |
| 1:2:1087:C:H2' | 1:2:1088:G:C8 | 2.57 | 0.40 |
| 1:2:1423:C:H2' | 1:2:1424:G:O4' | 2.21 | 0.40 |
| 1:2:1443:G:OP1 | 24:W:87:ARG:NH2 | 2.55 | 0.40 |
| 1:2:1517:A:C8 | 19:R:128:HIS:CD2 | 3.09 | 0.40 |
| 1:2:1525:U:H1' | 23:V:87:VAL:HG23 | 2.03 | 0.40 |
| 1:2:1697:G:C5 | 1:2:1698:C:C4 | 3.09 | 0.40 |
| 1:2:1736:U:OP1 | 12:K:42:ARG:HD3 | 2.21 | 0.40 |
| 1:2:1800:A:H2' | 1:2:1801:C:H6 | 1.85 | 0.40 |
| 4:C:54:THR:O | 4:C:161:ILE:HD11 | 2.21 | 0.40 |
| 4:C:81:ASN:OD1 | 4:C:81:ASN:N | 2.53 | 0.40 |
| 5:D:128:LYS:HG2 | 5:D:129:THR:O | 2.21 | 0.40 |
| 5:D:208:HIS:O | 5:D:208:HIS:CG | 2.74 | 0.40 |
| 7:F:196:GLY:C | 7:F:198:ILE:HA | 2.47 | 0.40 |
| 8:G:42:LEU:HD23 | 8:G:43:PRO:O | 2.22 | 0.40 |
| 8:G:85:GLY:O | 8:G:101:LEU:HD23 | 2.21 | 0.40 |
| 8:G:100:ARG:NH2 | 8:G:122:LYS:HA | 2.35 | 0.40 |
| 10:I:145:PHE:HE2 | 10:I:156:TYR:O | 2.04 | 0.40 |
| 15:N:59:LYS:HD3 | 15:N:134:LEU:HD13 | 2.04 | 0.40 |
| 19:R:24:GLN:HB3 | 19:R:28:MET:HE1 | 2.03 | 0.40 |
| 25:X:67:ASP:HA | 25:X:70:LEU:HG | 2.02 | 0.40 |
| 28:a:9:THR:O | 28:a:10:ARG:HD2 | 2.20 | 0.40 |
| 31:d:13:ARG:C | 31:d:32:VAL:HG23 | 2.47 | 0.40 |
| 33:f:135:HIS:HD2 | 33:f:140:TYR:H | 1.68 | 0.40 |
| 34:g:44:LYS:HG2 | 34:g:54:ILE:O | 2.21 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 36:i:102:ARG:HH11 | 36:i:106:ALA:HB1 | 1.86 | 0.40 |
| 39:z:587:U:H2' | 39:z:588:A:C8 | 2.57 | 0.40 |
| 1:2:73:C:O2' | 1:2:74:G:O4' | 2.36 | 0.40 |
| 1:2:385:G:OP1 | 15:N:79:LYS:HD2 | 2.21 | 0.40 |
| 1:2:445:A:H2' | 1:2:446:C:H6 | 1.86 | 0.40 |
| 1:2:927:C:H2' | 1:2:928:G:C8 | 2.56 | 0.40 |
| 1:2:1063:C:H2' | 1:2:1064:G:C8 | 2.56 | 0.40 |
| 1:2:1285:U:C5 | 33:f:97:LYS:HE2 | 2.56 | 0.40 |
| 1:2:1517:A:O2' | 22:U:144:ARG:NH1 | 2.54 | 0.40 |
| 1:2:1805:C:H2' | 1:2:1806:U:C6 | 2.57 | 0.40 |
| 1:2:1854:A:O2' | 1:2:1856:G:H8 | 2.05 | 0.40 |
| 4:C:23:THR:HA | 4:C:164:ASN:ND2 | 2.36 | 0.40 |
| 8:G:127:ARG:HB2 | 8:G:140:VAL:HG12 | 2.04 | 0.40 |
| 8:G:182:MET:HE2 | 8:G:182:MET:HA | 2.03 | 0.40 |
| 10:I:194:LEU:HA | 10:I:197:GLN:OE1 | 2.21 | 0.40 |
| 12:K:195:LEU:O | 12:K:199:LEU:HD23 | 2.22 | 0.40 |
| 18:Q:121:ARG:C | 18:Q:123:GLY:H | 2.29 | 0.40 |
| 22:U:73:ASN:HB3 | 22:U:76:GLN:HB2 | 2.04 | 0.40 |
| 26:Y:78:ARG:HH12 | 26:Y:126:LEU:HD13 | 1.86 | 0.40 |
| 28:a:111:LYS:O | 28:a:115:LYS:HG3 | 2.21 | 0.40 |
| 1:2:536:G:H2' | 1:2:537:G:C8 | 2.57 | 0.40 |
| 1:2:552:U:C4' | 13:L:132:GLN:HE22 | 2.35 | 0.40 |
| 1:2:624:A:H2' | 1:2:625:G:C8 | 2.56 | 0.40 |
| 1:2:746:C:H2' | 1:2:747:G:N9 | 2.36 | 0.40 |
| 1:2:811:U:N3 | 1:2:812:A:N7 | 2.70 | 0.40 |
| 1:2:958:A:H5'' | 18:Q:66:ARG:HD2 | 2.02 | 0.40 |
| 1:2:1259:U:H4' | 32:e:27:ARG:NH1 | 2.37 | 0.40 |
| 1:2:1741:U:H2' | 1:2:1742:C:H6 | 1.86 | 0.40 |
| 5:D:47:THR:HB | 5:D:65:ARG:NH1 | 2.37 | 0.40 |
| 5:D:133:TYR:CE1 | 5:D:221:PRO:HD2 | 2.57 | 0.40 |
| 6:E:231:LYS:HA | 6:E:234:SER:OG | 2.22 | 0.40 |
| 6:E:259:VAL:HG23 | 6:E:259:VAL:O | 2.20 | 0.40 |
| 8:G:31:PRO:HA | 8:G:81:THR:OG1 | 2.21 | 0.40 |
| 8:G:89:VAL:HA | 8:G:99:PHE:O | 2.21 | 0.40 |
| 10:I:197:GLN:O | 10:I:201:LYS:HG2 | 2.21 | 0.40 |
| 11:J:53:VAL:HG22 | 11:J:57:ARG:O | 2.21 | 0.40 |
| 13:L:22:LYS:HA | 13:L:22:LYS:HD2 | 1.89 | 0.40 |
| 17:P:75:LEU:HD23 | 17:P:81:ALA:HA | 2.04 | 0.40 |
| 20:S:97:GLN:NE2 | 20:S:98:LYS:HB3 | 2.36 | 0.40 |
| 27:Z:81:ILE:HD12 | 27:Z:81:ILE:HA | 1.98 | 0.40 |
| 27:Z:130:LEU:HA | 27:Z:133:LEU:HG | 2.03 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|------------------|------------------|--------------------------|-------------------|
| 28:a:11:LYS:O | 28:a:23:MET:HA | 2.22 | 0.40 |
| 32:e:4:GLN:NE2 | 32:e:5:GLN:HB2 | 2.36 | 0.40 |
| 34:g:218:LEU:HG | 34:g:228:TYR:CZ | 2.57 | 0.40 |
| 38:y:4:A:H2' | 38:y:5:G:C8 | 2.57 | 0.40 |
| 38:y:19:A:H2 | 38:y:59:A:C6 | 2.40 | 0.40 |
| 39:z:531:C:O2' | 39:z:532:G:C6 | 2.74 | 0.40 |
| 1:2:198:U:H3' | 1:2:199:G:C8 | 2.56 | 0.40 |
| 1:2:560:C:C2 | 1:2:561:U:C5 | 3.09 | 0.40 |
| 1:2:1088:G:N1 | 1:2:1154:G:O6 | 2.54 | 0.40 |
| 1:2:1349:A:C6 | 1:2:1350:G:C6 | 3.10 | 0.40 |
| 1:2:1683:C:H2' | 1:2:1684:C:C6 | 2.56 | 0.40 |
| 1:2:1785:A:H2' | 1:2:1786:G:O4' | 2.21 | 0.40 |
| 1:2:1839:A:H2' | 1:2:1840:G:H8 | 1.85 | 0.40 |
| 5:D:22:VAL:O | 5:D:27:LYS:NZ | 2.53 | 0.40 |
| 7:F:12:VAL:O | 7:F:16:ILE:HG12 | 2.21 | 0.40 |
| 8:G:77:ARG:HG3 | 8:G:82:TYR:CE2 | 2.57 | 0.40 |
| 10:I:170:ARG:HD3 | 10:I:172:LYS:HG2 | 2.02 | 0.40 |
| 13:L:29:LEU:O | 13:L:32:ILE:HG22 | 2.21 | 0.40 |
| 17:P:46:THR:HG22 | 17:P:49:GLN:OE1 | 2.21 | 0.40 |
| 27:Z:48:LYS:O | 27:Z:75:ILE:HG22 | 2.22 | 0.40 |
| 34:g:57:ARG:NE | 34:g:94:THR:HA | 2.36 | 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 | |
|-----|-------|---------------|-----------|----------|----------|-------------|-----|
| 2 | A | 276/315 (88%) | 267 (97%) | 9 (3%) | 0 | 100 | 100 |
| 3 | B | 462/485 (95%) | 453 (98%) | 9 (2%) | 0 | 100 | 100 |
| 4 | C | 206/295 (70%) | 179 (87%) | 26 (13%) | 1 (0%) | 24 | 63 |
| 5 | D | 213/264 (81%) | 195 (92%) | 17 (8%) | 1 (0%) | 24 | 63 |

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| Mol | Chain | Analysed | Favoured | Allowed | Outliers | Percentiles | |
|-----|-------|---------------|-----------|----------|----------|-------------|-----|
| 6 | E | 220/226 (97%) | 203 (92%) | 16 (7%) | 1 (0%) | 24 | 63 |
| 7 | F | 225/243 (93%) | 212 (94%) | 12 (5%) | 1 (0%) | 30 | 67 |
| 8 | G | 261/263 (99%) | 242 (93%) | 19 (7%) | 0 | 100 | 100 |
| 9 | H | 189/204 (93%) | 174 (92%) | 15 (8%) | 0 | 100 | 100 |
| 10 | I | 235/249 (94%) | 224 (95%) | 11 (5%) | 0 | 100 | 100 |
| 11 | J | 188/194 (97%) | 172 (92%) | 16 (8%) | 0 | 100 | 100 |
| 12 | K | 204/206 (99%) | 185 (91%) | 17 (8%) | 2 (1%) | 12 | 47 |
| 13 | L | 180/194 (93%) | 172 (96%) | 7 (4%) | 1 (1%) | 21 | 58 |
| 14 | M | 96/225 (43%) | 79 (82%) | 16 (17%) | 1 (1%) | 12 | 47 |
| 15 | N | 156/158 (99%) | 148 (95%) | 8 (5%) | 0 | 100 | 100 |
| 16 | O | 122/132 (92%) | 108 (88%) | 14 (12%) | 0 | 100 | 100 |
| 17 | P | 148/151 (98%) | 144 (97%) | 4 (3%) | 0 | 100 | 100 |
| 18 | Q | 134/168 (80%) | 117 (87%) | 17 (13%) | 0 | 100 | 100 |
| 19 | R | 133/145 (92%) | 115 (86%) | 18 (14%) | 0 | 100 | 100 |
| 20 | S | 139/146 (95%) | 127 (91%) | 12 (9%) | 0 | 100 | 100 |
| 21 | T | 124/135 (92%) | 121 (98%) | 3 (2%) | 0 | 100 | 100 |
| 22 | U | 140/152 (92%) | 124 (89%) | 15 (11%) | 1 (1%) | 18 | 55 |
| 23 | V | 139/141 (99%) | 126 (91%) | 12 (9%) | 1 (1%) | 18 | 55 |
| 24 | W | 102/119 (86%) | 96 (94%) | 6 (6%) | 0 | 100 | 100 |
| 25 | X | 80/82 (98%) | 72 (90%) | 8 (10%) | 0 | 100 | 100 |
| 26 | Y | 127/130 (98%) | 119 (94%) | 8 (6%) | 0 | 100 | 100 |
| 27 | Z | 140/143 (98%) | 132 (94%) | 8 (6%) | 0 | 100 | 100 |
| 28 | a | 124/126 (98%) | 117 (94%) | 7 (6%) | 0 | 100 | 100 |
| 29 | b | 97/115 (84%) | 91 (94%) | 6 (6%) | 0 | 100 | 100 |
| 30 | c | 82/84 (98%) | 72 (88%) | 10 (12%) | 0 | 100 | 100 |
| 31 | d | 62/64 (97%) | 59 (95%) | 3 (5%) | 0 | 100 | 100 |
| 32 | e | 51/56 (91%) | 42 (82%) | 9 (18%) | 0 | 100 | 100 |
| 33 | f | 69/156 (44%) | 59 (86%) | 10 (14%) | 0 | 100 | 100 |
| 34 | g | 311/317 (98%) | 286 (92%) | 25 (8%) | 0 | 100 | 100 |
| 35 | h | 73/125 (58%) | 71 (97%) | 2 (3%) | 0 | 100 | 100 |
| 36 | i | 57/59 (97%) | 51 (90%) | 6 (10%) | 0 | 100 | 100 |

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| Mol | Chain | Analysed | Favoured | Allowed | Outliers | Percentiles | |
|-----|-------|-----------------|------------|----------|----------|-------------|-----|
| 37 | l | 23/25 (92%) | 23 (100%) | 0 | 0 | 100 | 100 |
| All | All | 5588/6292 (89%) | 5177 (93%) | 401 (7%) | 10 (0%) | 44 | 77 |

All (10) Ramachandran outliers are listed below:

| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 14 | M | 34 | GLU |
| 22 | U | 144 | ARG |
| 7 | F | 193 | ASP |
| 12 | K | 156 | ALA |
| 12 | K | 158 | ILE |
| 23 | V | 40 | ALA |
| 5 | D | 207 | LEU |
| 13 | L | 148 | ILE |
| 6 | E | 246 | PHE |
| 4 | C | 206 | ASP |

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 | C | 174/244 (71%) | 174 (100%) | 0 | 100 | 100 |
| 5 | D | 196/231 (85%) | 196 (100%) | 0 | 100 | 100 |
| 6 | E | 186/187 (100%) | 186 (100%) | 0 | 100 | 100 |
| 7 | F | 190/202 (94%) | 190 (100%) | 0 | 100 | 100 |
| 8 | G | 225/225 (100%) | 225 (100%) | 0 | 100 | 100 |
| 9 | H | 161/170 (95%) | 161 (100%) | 0 | 100 | 100 |
| 10 | I | 207/218 (95%) | 207 (100%) | 0 | 100 | 100 |
| 11 | J | 170/174 (98%) | 169 (99%) | 1 (1%) | 78 | 81 |
| 12 | K | 177/177 (100%) | 177 (100%) | 0 | 100 | 100 |
| 13 | L | 157/168 (94%) | 157 (100%) | 0 | 100 | 100 |
| 14 | M | 89/173 (51%) | 89 (100%) | 0 | 100 | 100 |

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| Mol | Chain | Analysed | Rotameric | Outliers | Percentiles | |
|-----|-------|-----------------|-------------|----------|-------------|-----|
| 15 | N | 142/142 (100%) | 142 (100%) | 0 | 100 | 100 |
| 16 | O | 104/108 (96%) | 104 (100%) | 0 | 100 | 100 |
| 17 | P | 130/131 (99%) | 130 (100%) | 0 | 100 | 100 |
| 18 | Q | 106/130 (82%) | 106 (100%) | 0 | 100 | 100 |
| 19 | R | 121/130 (93%) | 121 (100%) | 0 | 100 | 100 |
| 20 | S | 117/121 (97%) | 117 (100%) | 0 | 100 | 100 |
| 21 | T | 114/121 (94%) | 114 (100%) | 0 | 100 | 100 |
| 22 | U | 122/132 (92%) | 122 (100%) | 0 | 100 | 100 |
| 23 | V | 113/113 (100%) | 113 (100%) | 0 | 100 | 100 |
| 24 | W | 94/107 (88%) | 94 (100%) | 0 | 100 | 100 |
| 25 | X | 67/67 (100%) | 67 (100%) | 0 | 100 | 100 |
| 26 | Y | 112/113 (99%) | 112 (100%) | 0 | 100 | 100 |
| 27 | Z | 114/115 (99%) | 114 (100%) | 0 | 100 | 100 |
| 28 | a | 108/108 (100%) | 108 (100%) | 0 | 100 | 100 |
| 29 | b | 87/99 (88%) | 87 (100%) | 0 | 100 | 100 |
| 30 | c | 76/76 (100%) | 76 (100%) | 0 | 100 | 100 |
| 31 | d | 57/57 (100%) | 57 (100%) | 0 | 100 | 100 |
| 32 | e | 47/49 (96%) | 47 (100%) | 0 | 100 | 100 |
| 33 | f | 64/140 (46%) | 64 (100%) | 0 | 100 | 100 |
| 34 | g | 272/275 (99%) | 272 (100%) | 0 | 100 | 100 |
| 35 | h | 66/103 (64%) | 66 (100%) | 0 | 100 | 100 |
| 36 | i | 49/49 (100%) | 49 (100%) | 0 | 100 | 100 |
| 37 | l | 24/24 (100%) | 24 (100%) | 0 | 100 | 100 |
| All | All | 4238/4679 (91%) | 4237 (100%) | 1 (0%) | 100 | 100 |

All (1) residues with a non-rotameric sidechain are listed below:

| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 11 | J | 97 | GLN |

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (51) such sidechains are listed below:

| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 4 | C | 9 | GLN |
| 4 | C | 33 | GLN |
| 4 | C | 50 | ASN |
| 4 | C | 70 | ASN |
| 4 | C | 111 | GLN |
| 5 | D | 95 | ASN |
| 5 | D | 101 | HIS |
| 5 | D | 159 | GLN |
| 5 | D | 163 | GLN |
| 6 | E | 121 | HIS |
| 6 | E | 257 | HIS |
| 7 | F | 101 | GLN |
| 8 | G | 142 | HIS |
| 9 | H | 82 | ASN |
| 9 | H | 83 | ASN |
| 10 | I | 146 | ASN |
| 11 | J | 25 | GLN |
| 11 | J | 168 | HIS |
| 12 | K | 64 | ASN |
| 12 | K | 165 | GLN |
| 13 | L | 75 | ASN |
| 13 | L | 113 | GLN |
| 13 | L | 132 | GLN |
| 13 | L | 134 | HIS |
| 15 | N | 5 | GLN |
| 15 | N | 94 | HIS |
| 15 | N | 156 | GLN |
| 17 | P | 105 | ASN |
| 19 | R | 6 | GLN |
| 19 | R | 35 | GLN |
| 20 | S | 86 | GLN |
| 22 | U | 17 | ASN |
| 22 | U | 19 | ASN |
| 23 | V | 42 | HIS |
| 23 | V | 117 | GLN |
| 24 | W | 92 | HIS |
| 25 | X | 2 | GLN |
| 26 | Y | 15 | ASN |
| 26 | Y | 70 | ASN |
| 26 | Y | 90 | GLN |
| 27 | Z | 26 | GLN |
| 28 | a | 106 | GLN |
| 28 | a | 112 | ASN |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 29 | b | 80 | HIS |
| 32 | e | 26 | ASN |
| 33 | f | 135 | HIS |
| 34 | g | 76 | GLN |
| 34 | g | 178 | ASN |
| 35 | h | 89 | GLN |
| 35 | h | 106 | GLN |
| 36 | i | 113 | ASN |

5.3.3 RNA ⓘ

| Mol | Chain | Analysed | Backbone Outliers | Pucker Outliers |
|-----|-------|-----------------|-------------------|-----------------|
| 1 | 2 | 1736/1863 (93%) | 362 (20%) | 4 (0%) |
| 38 | y | 74/75 (98%) | 35 (47%) | 0 |
| 39 | z | 91/93 (97%) | 32 (35%) | 0 |
| All | All | 1901/2031 (93%) | 429 (22%) | 4 (0%) |

All (429) RNA backbone outliers are listed below:

| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 1 | 2 | 4 | C |
| 1 | 2 | 11 | A |
| 1 | 2 | 26 | U |
| 1 | 2 | 33 | G |
| 1 | 2 | 39 | A |
| 1 | 2 | 41 | G |
| 1 | 2 | 42 | A |
| 1 | 2 | 44 | U |
| 1 | 2 | 46 | A |
| 1 | 2 | 55 | U |
| 1 | 2 | 56 | G |
| 1 | 2 | 60 | A |
| 1 | 2 | 67 | C |
| 1 | 2 | 68 | A |
| 1 | 2 | 72 | C |
| 1 | 2 | 73 | C |
| 1 | 2 | 74 | G |
| 1 | 2 | 76 | U |
| 1 | 2 | 79 | A |
| 1 | 2 | 80 | G |
| 1 | 2 | 94 | G |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 1 | 2 | 99 | A |
| 1 | 2 | 101 | U |
| 1 | 2 | 103 | A |
| 1 | 2 | 110 | U |
| 1 | 2 | 111 | A |
| 1 | 2 | 113 | G |
| 1 | 2 | 114 | G |
| 1 | 2 | 115 | U |
| 1 | 2 | 126 | G |
| 1 | 2 | 127 | C |
| 1 | 2 | 143 | U |
| 1 | 2 | 147 | A |
| 1 | 2 | 148 | U |
| 1 | 2 | 162 | C |
| 1 | 2 | 168 | C |
| 1 | 2 | 169 | U |
| 1 | 2 | 170 | A |
| 1 | 2 | 171 | A |
| 1 | 2 | 172 | U |
| 1 | 2 | 173 | A |
| 1 | 2 | 178 | C |
| 1 | 2 | 179 | C |
| 1 | 2 | 181 | A |
| 1 | 2 | 182 | C |
| 1 | 2 | 183 | G |
| 1 | 2 | 191 | C |
| 1 | 2 | 197 | U |
| 1 | 2 | 202 | U |
| 1 | 2 | 204 | G |
| 1 | 2 | 223 | A |
| 1 | 2 | 224 | U |
| 1 | 2 | 225 | C |
| 1 | 2 | 226 | A |
| 1 | 2 | 239 | U |
| 1 | 2 | 271 | G |
| 1 | 2 | 277 | U |
| 1 | 2 | 278 | U |
| 1 | 2 | 285 | U |
| 1 | 2 | 293 | A |
| 1 | 2 | 296 | U |
| 1 | 2 | 297 | C |
| 1 | 2 | 298 | G |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 1 | 2 | 299 | G |
| 1 | 2 | 300 | G |
| 1 | 2 | 303 | G |
| 1 | 2 | 308 | C |
| 1 | 2 | 311 | C |
| 1 | 2 | 315 | C |
| 1 | 2 | 316 | C |
| 1 | 2 | 317 | G |
| 1 | 2 | 322 | G |
| 1 | 2 | 325 | G |
| 1 | 2 | 337 | G |
| 1 | 2 | 340 | C |
| 1 | 2 | 347 | C |
| 1 | 2 | 350 | A |
| 1 | 2 | 352 | C |
| 1 | 2 | 354 | A |
| 1 | 2 | 358 | U |
| 1 | 2 | 359 | C |
| 1 | 2 | 375 | G |
| 1 | 2 | 376 | C |
| 1 | 2 | 388 | A |
| 1 | 2 | 389 | C |
| 1 | 2 | 390 | C |
| 1 | 2 | 399 | C |
| 1 | 2 | 403 | G |
| 1 | 2 | 408 | A |
| 1 | 2 | 438 | A |
| 1 | 2 | 440 | C |
| 1 | 2 | 442 | G |
| 1 | 2 | 457 | G |
| 1 | 2 | 462 | C |
| 1 | 2 | 466 | A |
| 1 | 2 | 474 | A |
| 1 | 2 | 475 | A |
| 1 | 2 | 477 | U |
| 1 | 2 | 479 | A |
| 1 | 2 | 483 | A |
| 1 | 2 | 487 | C |
| 1 | 2 | 492 | C |
| 1 | 2 | 506 | A |
| 1 | 2 | 507 | C |
| 1 | 2 | 513 | A |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 1 | 2 | 515 | A |
| 1 | 2 | 518 | A |
| 1 | 2 | 541 | U |
| 1 | 2 | 542 | G |
| 1 | 2 | 547 | U |
| 1 | 2 | 548 | G |
| 1 | 2 | 553 | G |
| 1 | 2 | 554 | A |
| 1 | 2 | 563 | U |
| 1 | 2 | 566 | A |
| 1 | 2 | 577 | A |
| 1 | 2 | 579 | G |
| 1 | 2 | 580 | A |
| 1 | 2 | 581 | U |
| 1 | 2 | 583 | C |
| 1 | 2 | 596 | G |
| 1 | 2 | 597 | U |
| 1 | 2 | 598 | C |
| 1 | 2 | 604 | G |
| 1 | 2 | 607 | G |
| 1 | 2 | 613 | G |
| 1 | 2 | 617 | U |
| 1 | 2 | 621 | U |
| 1 | 2 | 633 | A |
| 1 | 2 | 650 | C |
| 1 | 2 | 658 | A |
| 1 | 2 | 659 | A |
| 1 | 2 | 661 | A |
| 1 | 2 | 662 | A |
| 1 | 2 | 663 | G |
| 1 | 2 | 673 | G |
| 1 | 2 | 674 | G |
| 1 | 2 | 678 | U |
| 1 | 2 | 679 | U |
| 1 | 2 | 680 | G |
| 1 | 2 | 682 | G |
| 1 | 2 | 729 | C |
| 1 | 2 | 730 | C |
| 1 | 2 | 731 | C |
| 1 | 2 | 732 | C |
| 1 | 2 | 735 | C |
| 1 | 2 | 736 | C |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 1 | 2 | 737 | C |
| 1 | 2 | 738 | U |
| 1 | 2 | 739 | U |
| 1 | 2 | 740 | G |
| 1 | 2 | 741 | C |
| 1 | 2 | 742 | C |
| 1 | 2 | 743 | U |
| 1 | 2 | 744 | C |
| 1 | 2 | 747 | G |
| 1 | 2 | 748 | G |
| 1 | 2 | 749 | C |
| 1 | 2 | 750 | G |
| 1 | 2 | 787 | C |
| 1 | 2 | 789 | G |
| 1 | 2 | 793 | C |
| 1 | 2 | 794 | G |
| 1 | 2 | 796 | U |
| 1 | 2 | 797 | U |
| 1 | 2 | 800 | U |
| 1 | 2 | 807 | A |
| 1 | 2 | 818 | U |
| 1 | 2 | 819 | U |
| 1 | 2 | 820 | C |
| 1 | 2 | 823 | A |
| 1 | 2 | 827 | G |
| 1 | 2 | 829 | C |
| 1 | 2 | 832 | G |
| 1 | 2 | 833 | A |
| 1 | 2 | 835 | C |
| 1 | 2 | 837 | G |
| 1 | 2 | 843 | A |
| 1 | 2 | 865 | A |
| 1 | 2 | 868 | A |
| 1 | 2 | 869 | G |
| 1 | 2 | 870 | G |
| 1 | 2 | 873 | C |
| 1 | 2 | 874 | G |
| 1 | 2 | 883 | U |
| 1 | 2 | 886 | U |
| 1 | 2 | 894 | U |
| 1 | 2 | 901 | C |
| 1 | 2 | 907 | C |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | 2 | 909 | A |
| 1 | 2 | 910 | U |
| 1 | 2 | 913 | U |
| 1 | 2 | 915 | A |
| 1 | 2 | 916 | A |
| 1 | 2 | 918 | A |
| 1 | 2 | 929 | G |
| 1 | 2 | 930 | G |
| 1 | 2 | 951 | A |
| 1 | 2 | 952 | G |
| 1 | 2 | 956 | U |
| 1 | 2 | 966 | G |
| 1 | 2 | 967 | G |
| 1 | 2 | 986 | A |
| 1 | 2 | 988 | A |
| 1 | 2 | 1004 | A |
| 1 | 2 | 1013 | U |
| 1 | 2 | 1016 | A |
| 1 | 2 | 1019 | A |
| 1 | 2 | 1022 | C |
| 1 | 2 | 1023 | A |
| 1 | 2 | 1041 | U |
| 1 | 2 | 1045 | A |
| 1 | 2 | 1049 | C |
| 1 | 2 | 1051 | A |
| 1 | 2 | 1056 | A |
| 1 | 2 | 1057 | U |
| 1 | 2 | 1058 | A |
| 1 | 2 | 1076 | A |
| 1 | 2 | 1079 | A |
| 1 | 2 | 1081 | C |
| 1 | 2 | 1097 | U |
| 1 | 2 | 1107 | U |
| 1 | 2 | 1112 | C |
| 1 | 2 | 1113 | C |
| 1 | 2 | 1125 | G |
| 1 | 2 | 1129 | A |
| 1 | 2 | 1135 | C |
| 1 | 2 | 1139 | A |
| 1 | 2 | 1144 | A |
| 1 | 2 | 1145 | A |
| 1 | 2 | 1146 | A |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | 2 | 1150 | U |
| 1 | 2 | 1151 | U |
| 1 | 2 | 1153 | G |
| 1 | 2 | 1160 | G |
| 1 | 2 | 1191 | A |
| 1 | 2 | 1199 | G |
| 1 | 2 | 1203 | G |
| 1 | 2 | 1204 | A |
| 1 | 2 | 1211 | C |
| 1 | 2 | 1212 | C |
| 1 | 2 | 1217 | G |
| 1 | 2 | 1220 | G |
| 1 | 2 | 1224 | A |
| 1 | 2 | 1238 | U |
| 1 | 2 | 1244 | U |
| 1 | 2 | 1245 | C |
| 1 | 2 | 1247 | A |
| 1 | 2 | 1249 | A |
| 1 | 2 | 1252 | G |
| 1 | 2 | 1253 | G |
| 1 | 2 | 1254 | A |
| 1 | 2 | 1261 | A |
| 1 | 2 | 1269 | C |
| 1 | 2 | 1271 | G |
| 1 | 2 | 1280 | A |
| 1 | 2 | 1285 | U |
| 1 | 2 | 1287 | A |
| 1 | 2 | 1297 | A |
| 1 | 2 | 1298 | G |
| 1 | 2 | 1317 | G |
| 1 | 2 | 1323 | G |
| 1 | 2 | 1333 | C |
| 1 | 2 | 1339 | U |
| 1 | 2 | 1352 | G |
| 1 | 2 | 1354 | U |
| 1 | 2 | 1359 | C |
| 1 | 2 | 1360 | U |
| 1 | 2 | 1365 | A |
| 1 | 2 | 1366 | A |
| 1 | 2 | 1367 | U |
| 1 | 2 | 1368 | U |
| 1 | 2 | 1374 | A |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | 2 | 1386 | U |
| 1 | 2 | 1399 | C |
| 1 | 2 | 1407 | G |
| 1 | 2 | 1408 | C |
| 1 | 2 | 1409 | G |
| 1 | 2 | 1411 | C |
| 1 | 2 | 1412 | C |
| 1 | 2 | 1414 | C |
| 1 | 2 | 1415 | C |
| 1 | 2 | 1427 | G |
| 1 | 2 | 1431 | C |
| 1 | 2 | 1432 | C |
| 1 | 2 | 1433 | C |
| 1 | 2 | 1438 | U |
| 1 | 2 | 1445 | G |
| 1 | 2 | 1446 | G |
| 1 | 2 | 1449 | C |
| 1 | 2 | 1450 | A |
| 1 | 2 | 1451 | A |
| 1 | 2 | 1459 | U |
| 1 | 2 | 1471 | G |
| 1 | 2 | 1472 | A |
| 1 | 2 | 1473 | U |
| 1 | 2 | 1485 | A |
| 1 | 2 | 1486 | G |
| 1 | 2 | 1487 | G |
| 1 | 2 | 1489 | C |
| 1 | 2 | 1490 | U |
| 1 | 2 | 1491 | G |
| 1 | 2 | 1494 | A |
| 1 | 2 | 1506 | G |
| 1 | 2 | 1507 | U |
| 1 | 2 | 1516 | C |
| 1 | 2 | 1517 | A |
| 1 | 2 | 1528 | A |
| 1 | 2 | 1531 | G |
| 1 | 2 | 1539 | C |
| 1 | 2 | 1543 | G |
| 1 | 2 | 1547 | G |
| 1 | 2 | 1548 | C |
| 1 | 2 | 1549 | C |
| 1 | 2 | 1550 | U |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | 2 | 1551 | A |
| 1 | 2 | 1555 | U |
| 1 | 2 | 1574 | A |
| 1 | 2 | 1575 | A |
| 1 | 2 | 1580 | U |
| 1 | 2 | 1583 | A |
| 1 | 2 | 1593 | G |
| 1 | 2 | 1596 | A |
| 1 | 2 | 1601 | G |
| 1 | 2 | 1616 | U |
| 1 | 2 | 1617 | U |
| 1 | 2 | 1618 | A |
| 1 | 2 | 1636 | A |
| 1 | 2 | 1643 | G |
| 1 | 2 | 1654 | U |
| 1 | 2 | 1656 | A |
| 1 | 2 | 1659 | A |
| 1 | 2 | 1660 | G |
| 1 | 2 | 1675 | G |
| 1 | 2 | 1694 | A |
| 1 | 2 | 1695 | C |
| 1 | 2 | 1696 | C |
| 1 | 2 | 1715 | U |
| 1 | 2 | 1716 | U |
| 1 | 2 | 1717 | G |
| 1 | 2 | 1724 | U |
| 1 | 2 | 1732 | G |
| 1 | 2 | 1739 | G |
| 1 | 2 | 1777 | C |
| 1 | 2 | 1778 | G |
| 1 | 2 | 1791 | U |
| 1 | 2 | 1792 | C |
| 1 | 2 | 1817 | A |
| 1 | 2 | 1823 | G |
| 1 | 2 | 1829 | A |
| 1 | 2 | 1832 | U |
| 1 | 2 | 1833 | U |
| 1 | 2 | 1837 | G |
| 1 | 2 | 1843 | G |
| 1 | 2 | 1846 | C |
| 1 | 2 | 1853 | A |
| 1 | 2 | 1854 | A |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | 2 | 1855 | G |
| 1 | 2 | 1856 | G |
| 1 | 2 | 1857 | A |
| 1 | 2 | 1859 | C |
| 1 | 2 | 1862 | U |
| 38 | y | 2 | G |
| 38 | y | 4 | A |
| 38 | y | 6 | A |
| 38 | y | 8 | U |
| 38 | y | 9 | G |
| 38 | y | 10 | G |
| 38 | y | 11 | C |
| 38 | y | 12 | G |
| 38 | y | 14 | A |
| 38 | y | 15 | G |
| 38 | y | 17 | G |
| 38 | y | 18 | G |
| 38 | y | 19 | A |
| 38 | y | 20 | A |
| 38 | y | 21 | G |
| 38 | y | 26 | C |
| 38 | y | 32 | C |
| 38 | y | 40 | C |
| 38 | y | 44 | G |
| 38 | y | 45 | G |
| 38 | y | 46 | U |
| 38 | y | 47 | C |
| 38 | y | 48 | G |
| 38 | y | 52 | G |
| 38 | y | 53 | A |
| 38 | y | 54 | U |
| 38 | y | 55 | C |
| 38 | y | 58 | A |
| 38 | y | 60 | C |
| 38 | y | 62 | A |
| 38 | y | 71 | U |
| 38 | y | 72 | A |
| 38 | y | 73 | C |
| 38 | y | 74 | C |
| 38 | y | 75 | A |
| 39 | z | 525 | A |
| 39 | z | 531 | C |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 39 | z | 534 | A |
| 39 | z | 535 | A |
| 39 | z | 536 | C |
| 39 | z | 537 | C |
| 39 | z | 538 | C |
| 39 | z | 539 | C |
| 39 | z | 540 | C |
| 39 | z | 546 | G |
| 39 | z | 549 | G |
| 39 | z | 550 | A |
| 39 | z | 556 | G |
| 39 | z | 557 | C |
| 39 | z | 558 | C |
| 39 | z | 559 | U |
| 39 | z | 562 | G |
| 39 | z | 564 | G |
| 39 | z | 565 | G |
| 39 | z | 568 | A |
| 39 | z | 575 | A |
| 39 | z | 576 | C |
| 39 | z | 577 | G |
| 39 | z | 578 | U |
| 39 | z | 585 | G |
| 39 | z | 586 | A |
| 39 | z | 587 | U |
| 39 | z | 594 | G |
| 39 | z | 832 | A |
| 39 | z | 833 | U |
| 39 | z | 837 | G |
| 39 | z | 838 | C |

All (4) RNA pucker outliers are listed below:

| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 1 | 2 | 740 | G |
| 1 | 2 | 747 | G |
| 1 | 2 | 792 | G |
| 1 | 2 | 914 | U |

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

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](#)

There are no ligands in this entry.

5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

The following chains have linkage breaks:

| Mol | Chain | Number of breaks |
|-----|-------|------------------|
| 39 | z | 1 |

All chain breaks are listed below:

| Model | Chain | Residue-1 | Atom-1 | Residue-2 | Atom-2 | Distance (Å) |
|-------|-------|-----------|--------|-----------|--------|--------------|
| 1 | z | 601:C | O3' | 831:A | P | 113.25 |

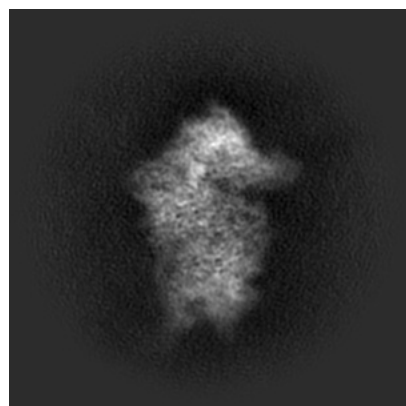
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-64645. These allow visual inspection of the internal detail of the map and identification of artifacts.

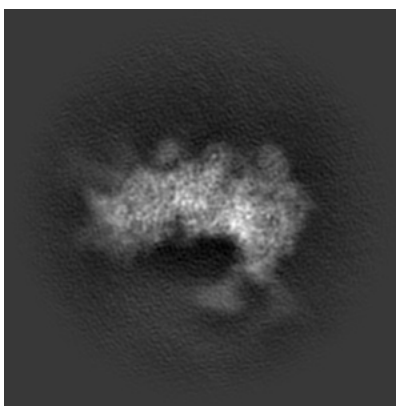
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

6.1 Orthogonal projections [i](#)

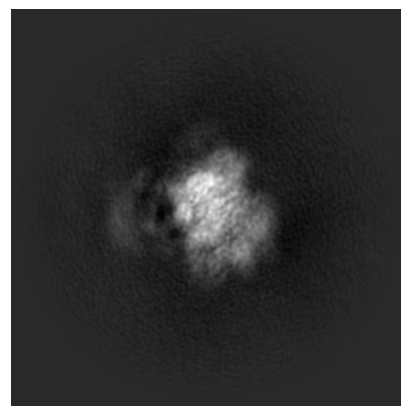
6.1.1 Primary map



X

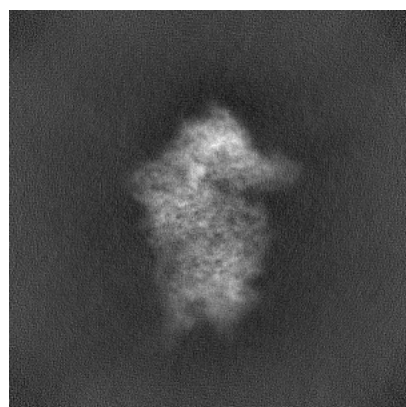


Y

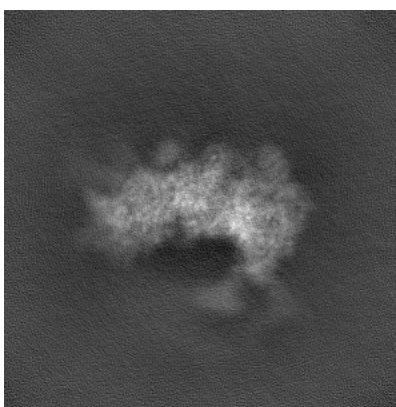


Z

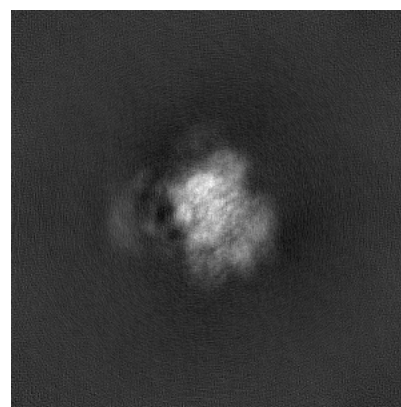
6.1.2 Raw map



X



Y

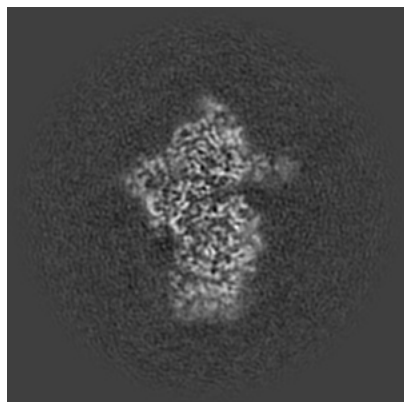


Z

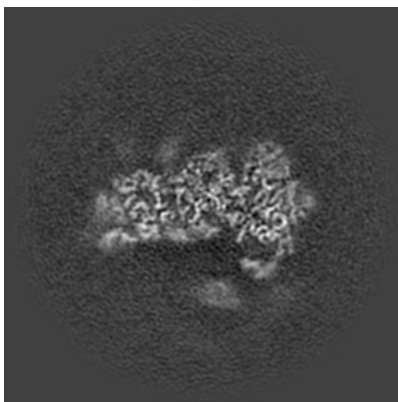
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

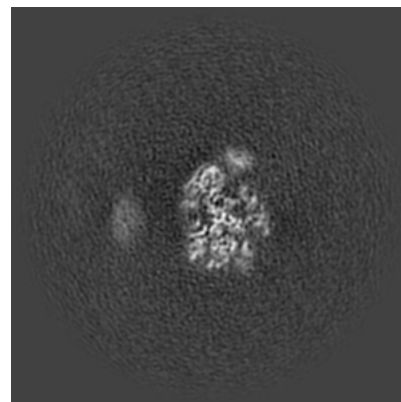
6.2.1 Primary map



X Index: 200

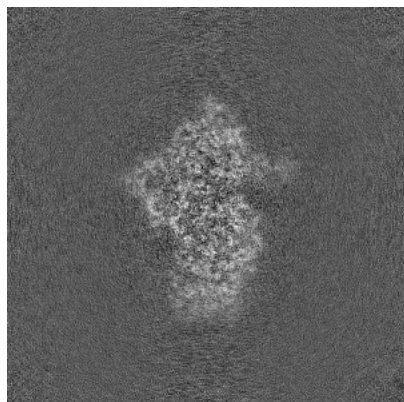


Y Index: 200

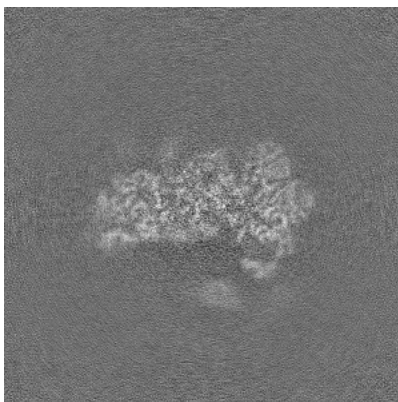


Z Index: 200

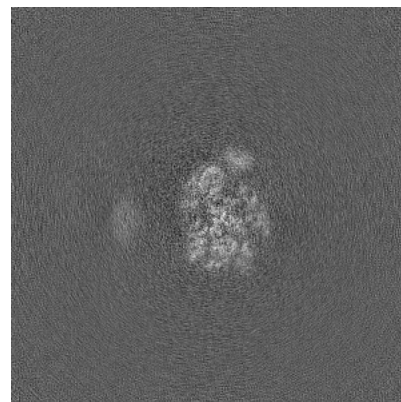
6.2.2 Raw map



X Index: 200



Y Index: 200

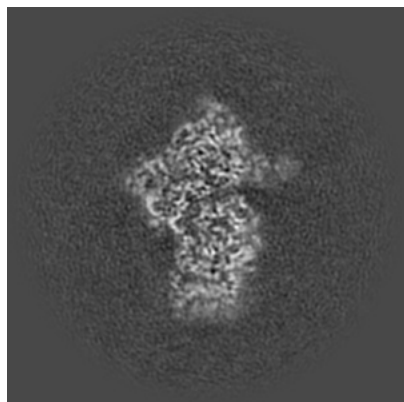


Z Index: 200

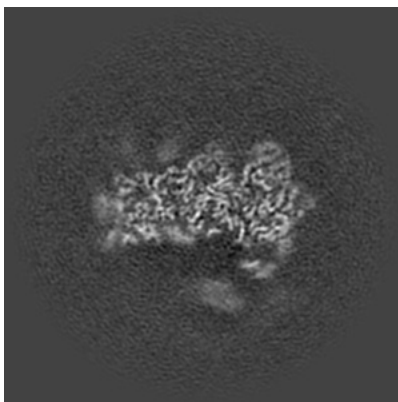
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

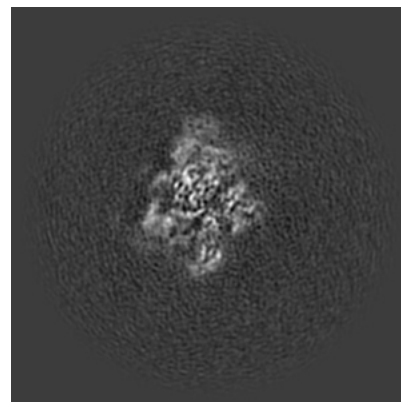
6.3.1 Primary map



X Index: 199

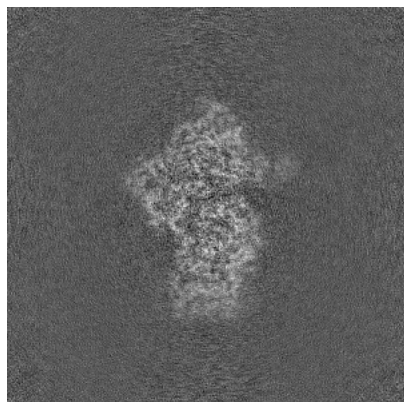


Y Index: 197

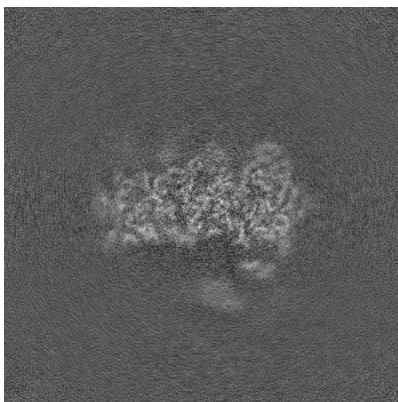


Z Index: 243

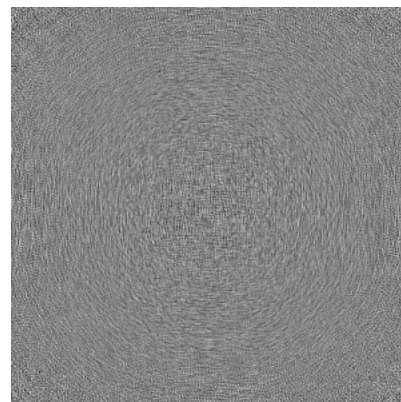
6.3.2 Raw map



X Index: 199



Y Index: 196

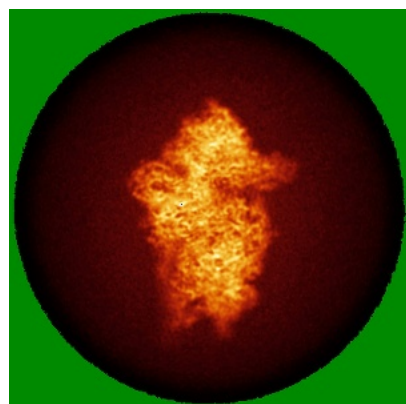


Z Index: 0

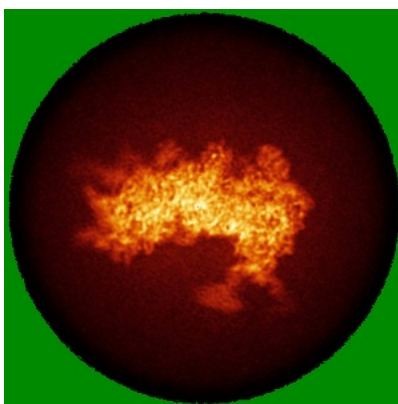
The images above show the largest variance slices of the map in three orthogonal directions.

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

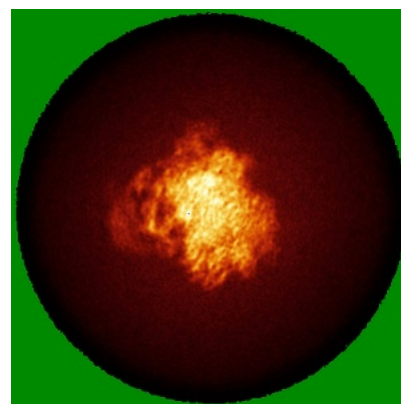
6.4.1 Primary map



X

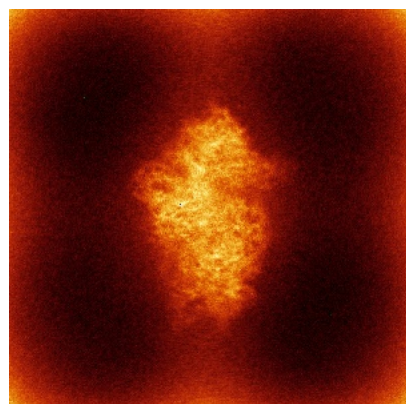


Y

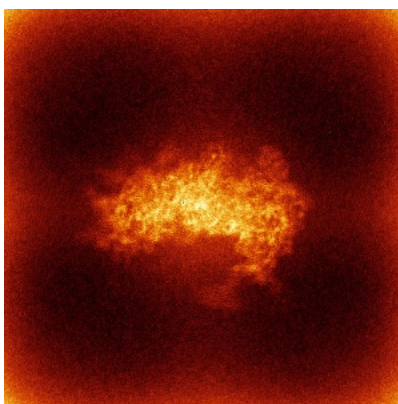


Z

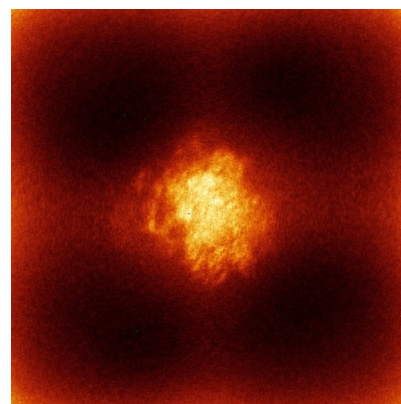
6.4.2 Raw map



X



Y

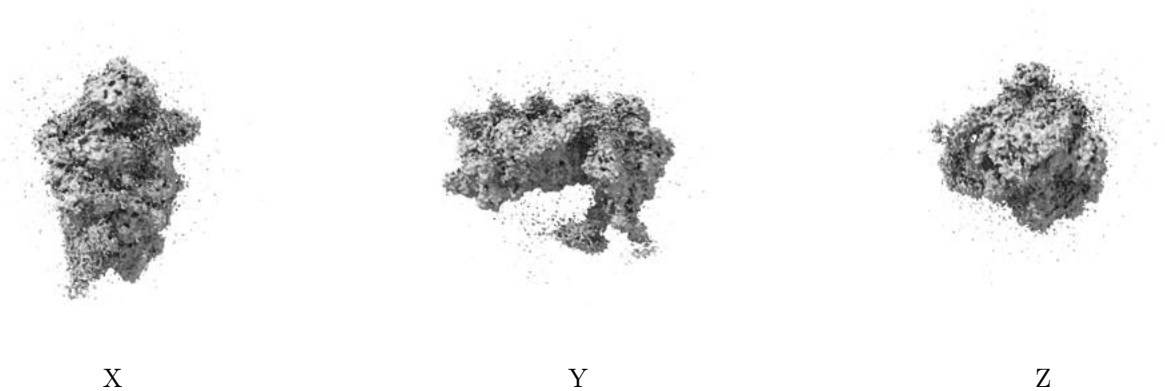


Z

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.

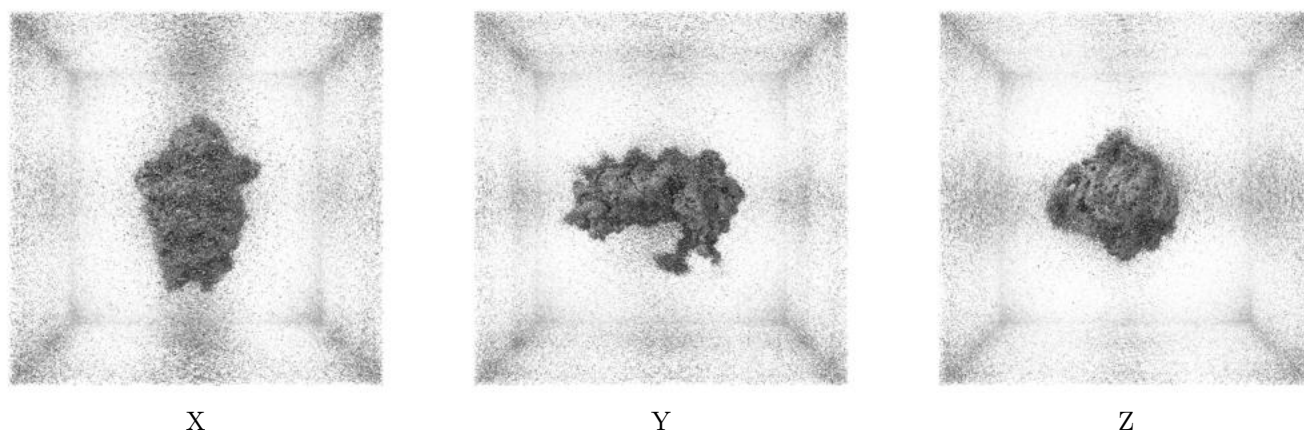
6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 0.0618. 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.

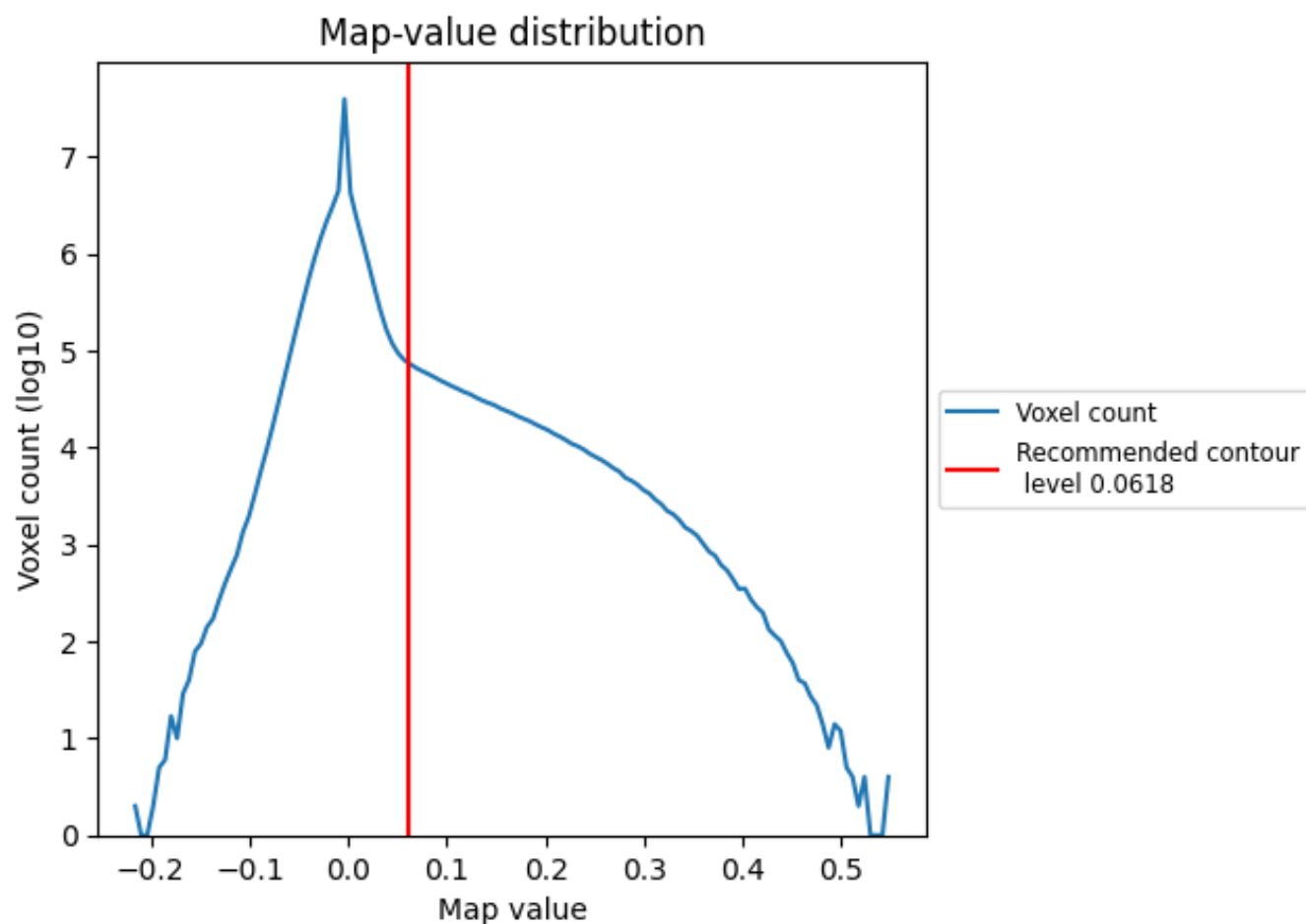
6.6 Mask visualisation [i](#)

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

7 Map analysis [i](#)

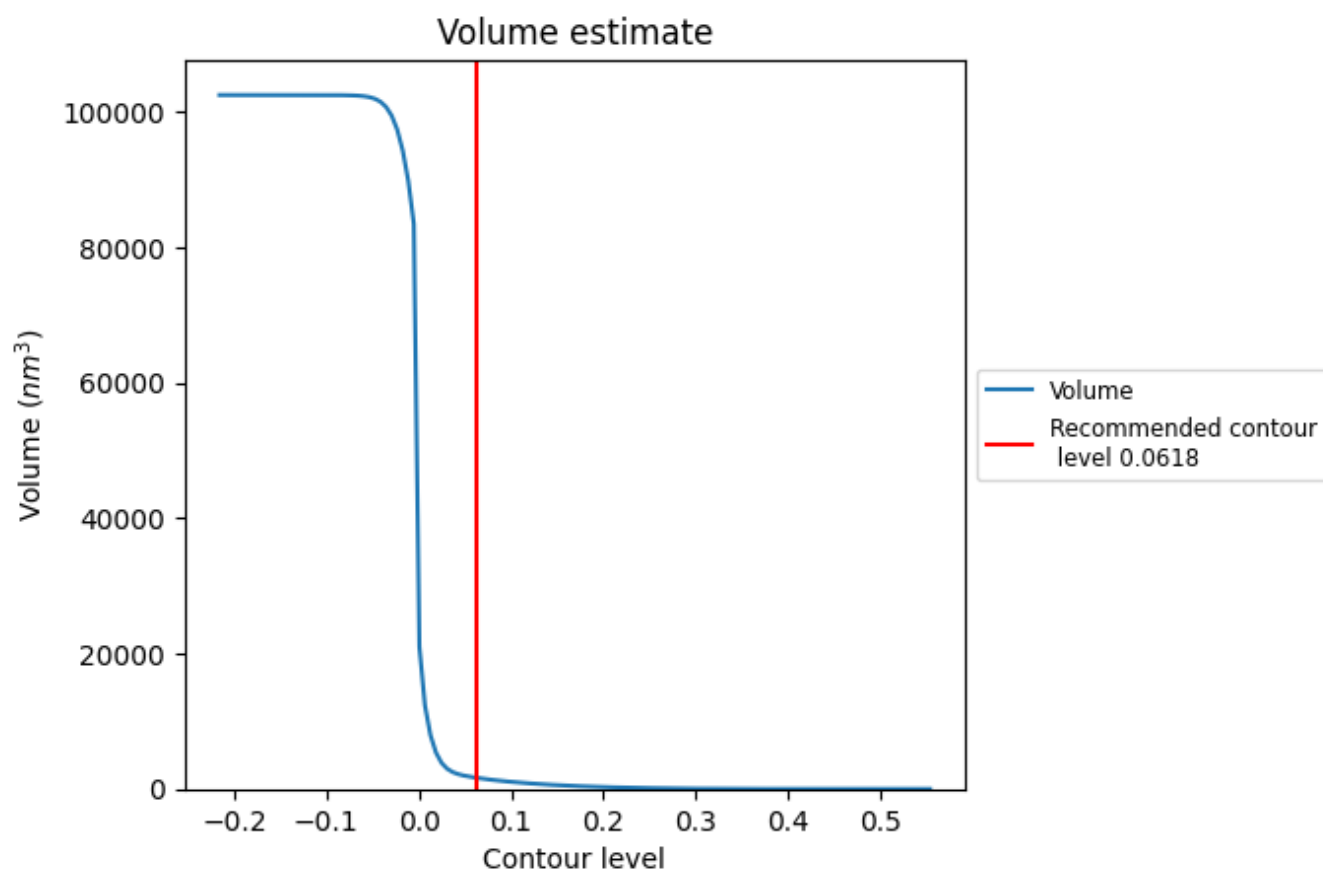
This section contains the results of statistical analysis of the map.

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.

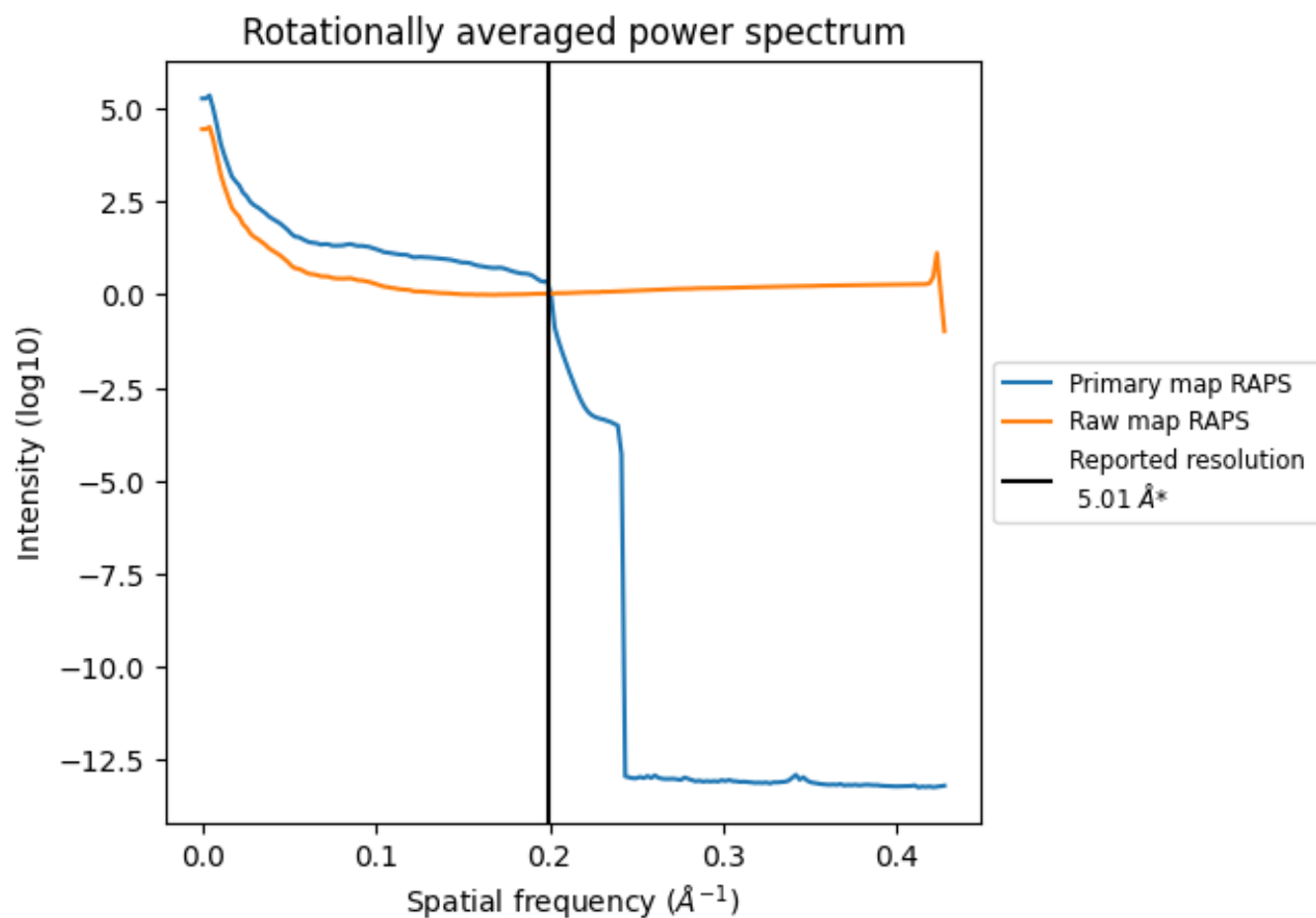
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 1660 nm^3 ; this corresponds to an approximate mass of 1500 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 ⓘ

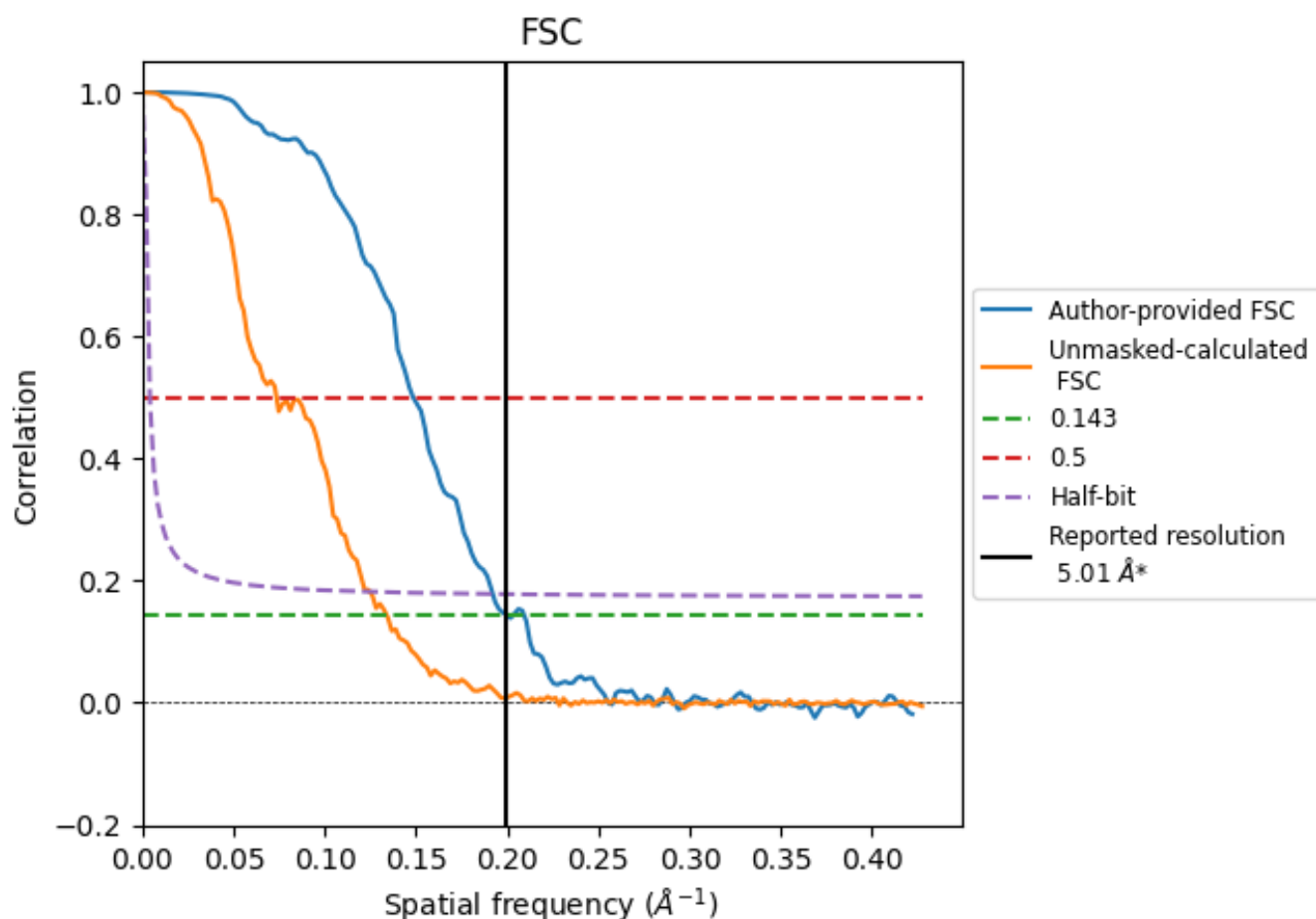


*Reported resolution corresponds to spatial frequency of 0.200 Å⁻¹

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 0.200 \AA^{-1}

8.2 Resolution estimates [i](#)

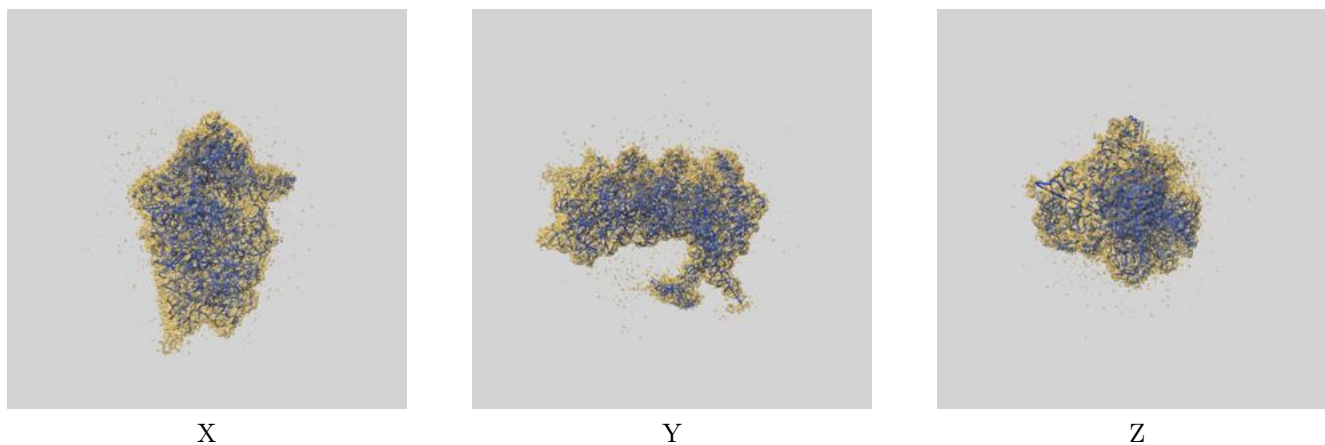
| Resolution estimate (Å) | Estimation criterion (FSC cut-off) | | |
|---------------------------|------------------------------------|-------|----------|
| | 0.143 | 0.5 | Half-bit |
| Reported by author | 5.01 | - | - |
| Author-provided FSC curve | 5.01 | 6.73 | 5.21 |
| Unmasked-calculated* | 7.46 | 13.59 | 8.02 |

*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 7.46 differs from the reported value 5.01 by more than 10 %

9 Map-model fit [i](#)

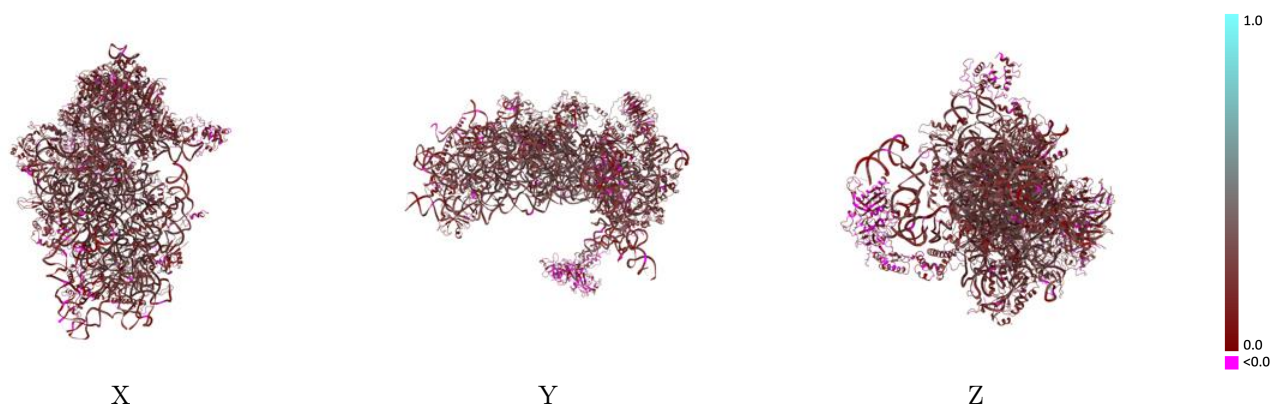
This section contains information regarding the fit between EMDB map EMD-64645 and PDB model 9UZL. Per-residue inclusion information can be found in section [3](#) on page [12](#).

9.1 Map-model overlay [i](#)



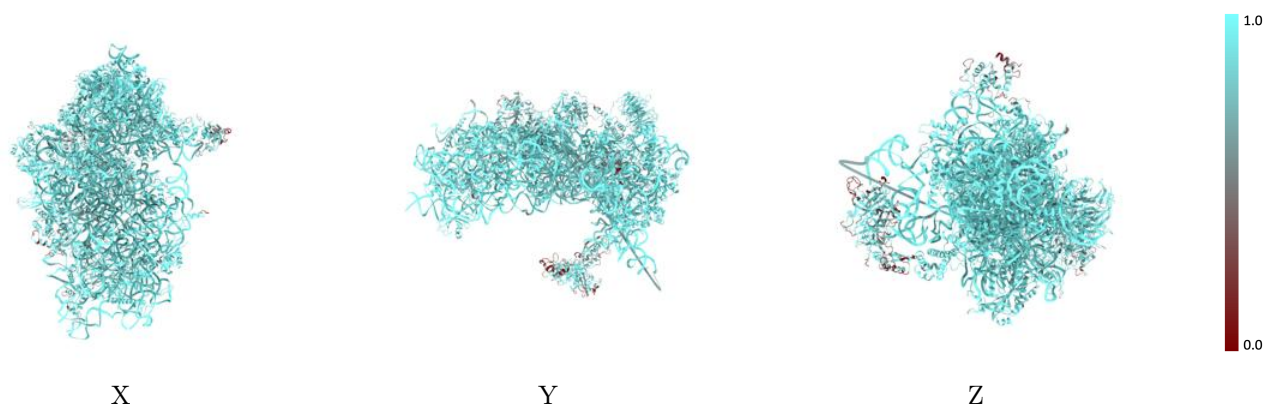
The images above show the 3D surface view of the map at the recommended contour level 0.0618 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



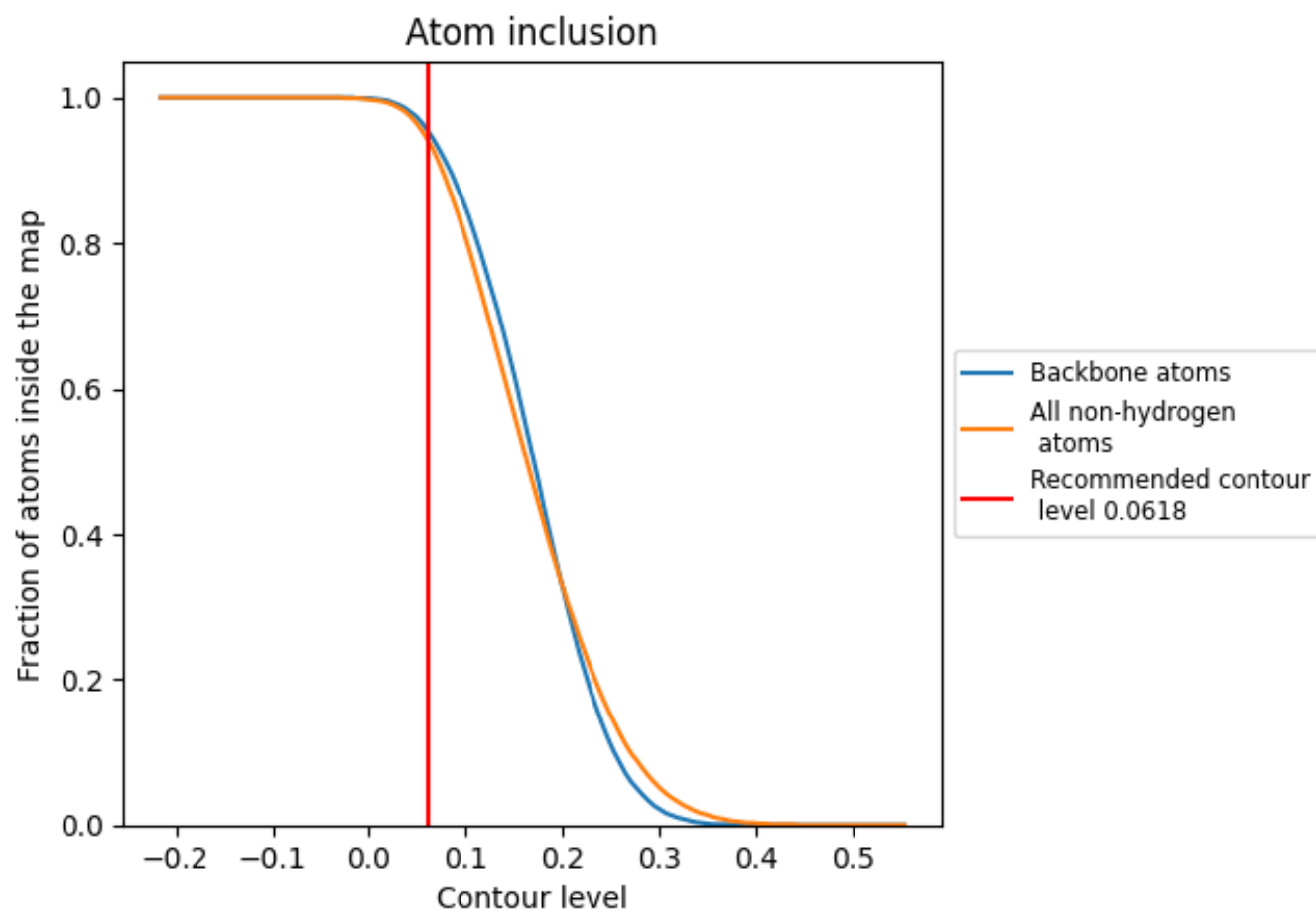
The images above show the model with each residue coloured according to its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions 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 (0.0618).







































































9.4 Atom inclusion ⓘ



At the recommended contour level, 95% of all backbone atoms, 94% 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 (0.0618) and Q-score for the entire model and for each chain.

| Chain | Atom inclusion | Q-score |
|-------|--|--|
| All |  0.9390 |  0.2330 |
| 2 |  0.9860 |  0.2600 |
| A |  0.8150 |  0.1880 |
| B |  0.7000 |  0.1000 |
| C |  0.8910 |  0.2400 |
| D |  0.9240 |  0.2150 |
| E |  0.9080 |  0.2490 |
| F |  0.9110 |  0.2150 |
| G |  0.9510 |  0.2460 |
| H |  0.9370 |  0.2310 |
| I |  0.9570 |  0.1950 |
| J |  0.7160 |  0.1760 |
| K |  0.9080 |  0.2050 |
| L |  0.9280 |  0.2270 |
| M |  0.9440 |  0.2080 |
| N |  0.8590 |  0.2400 |
| O |  0.6610 |  0.1140 |
| P |  0.9350 |  0.2320 |
| Q |  0.9310 |  0.2080 |
| R |  0.9560 |  0.2150 |
| S |  0.9380 |  0.2270 |
| T |  0.8650 |  0.2280 |
| U |  0.9420 |  0.2230 |
| V |  0.9670 |  0.2240 |
| W |  0.9480 |  0.2100 |
| X |  0.9290 |  0.2390 |
| Y |  0.8600 |  0.2530 |
| Z |  0.9370 |  0.2650 |
| a |  0.9630 |  0.2130 |
| b |  0.9430 |  0.2720 |
| c |  0.9180 |  0.2380 |
| d |  0.9690 |  0.2480 |
| e |  0.9560 |  0.2330 |
| f |  0.7390 |  0.1330 |
| g |  0.9390 |  0.2030 |



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| Chain | Atom inclusion | Q-score |
|-------|--|--|
| h |  0.9470 |  0.1970 |
| i |  0.8880 |  0.1830 |
| l |  0.9680 |  0.2520 |
| y |  0.9780 |  0.2060 |
| z |  0.9670 |  0.1930 |