



Full wwPDB EM Validation Report ⓘ

Oct 30, 2025 – 07:01 PM EDT

PDB ID : 9O17 / pdb_00009o17
EMDB ID : EMD-49998
Title : Cryo-EM structure of Methanosarcina acetivorans 70S ribosome
Authors : Ghosh, A.; Fordjour, G.N.R.; Armache, J.-P.; Ferry, J.G.; Murakami, K.S.;
Bevilacqua, P.C.
Deposited on : 2025-04-03
Resolution : 2.92 Å(reported)

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

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

A user guide is available at

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

The types of validation reports are described at

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

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

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

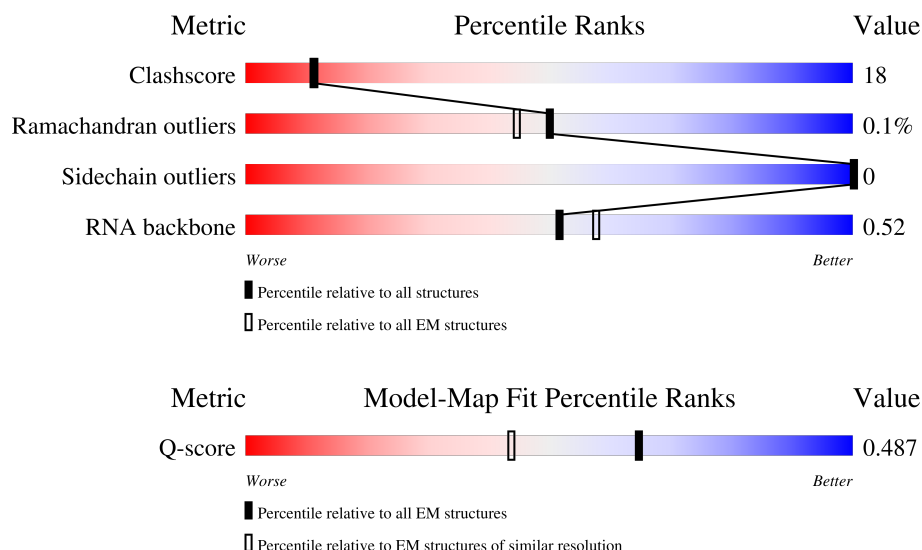
1 Overall quality at a glance

The following experimental techniques were used to determine the structure:

ELECTRON MICROSCOPY

The reported resolution of this entry is 2.92 Å.

Percentile scores (ranging between 0-100) for global validation metrics of the entry are shown in the following graphic. The table shows the number of entries on which the scores are based.



| Metric | Whole archive (#Entries) | EM structures (#Entries) | Similar EM resolution (#Entries, resolution range(Å)) |
|-----------------------|-----------------------------|-----------------------------|--|
| Clashscore | 210492 | 15764 | - |
| Ramachandran outliers | 207382 | 16835 | - |
| Sidechain outliers | 206894 | 16415 | - |
| RNA backbone | 6643 | 2191 | - |
| Q-score | - | 25397 | 13007 (2.42 - 3.42) |

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 | AA | 1478 | |
| 2 | AB | 247 | |
| 3 | AC | 318 | |

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| Mol | Chain | Length | Quality of chain |
|-----|-------|--------|------------------|
| 4 | AD | 203 | |
| 5 | AE | 218 | |
| 6 | AF | 235 | |
| 7 | AG | 209 | |
| 8 | AI | 189 | |
| 9 | AJ | 130 | |
| 10 | AK | 125 | |
| 11 | AL | 134 | |
| 12 | AN | 126 | |
| 13 | AO | 142 | |
| 14 | AP | 182 | |
| 15 | AR | 50 | |
| 16 | AQ | 152 | |
| 17 | AS | 109 | |
| 18 | AU | 136 | |
| 19 | AV | 149 | |
| 20 | AW | 101 | |
| 21 | AX | 62 | |
| 22 | AY | 76 | |
| 23 | AH | 136 | |
| 24 | AM | 102 | |
| 25 | AT | 64 | |
| 26 | BA | 2899 | |
| 27 | BB | 129 | |
| 28 | BC | 238 | |



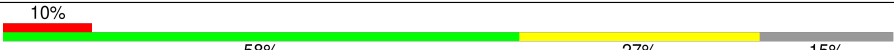
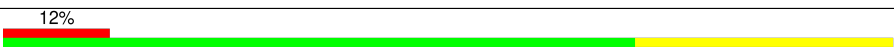

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| Mol | Chain | Length | Quality of chain |
|-----|-------|--------|------------------|
| 29 | BD | 337 | |
| 30 | BE | 253 | |
| 31 | BF | 165 | |
| 32 | BG | 176 | |
| 33 | BH | 120 | |
| 34 | BI | 173 | |
| 35 | BJ | 143 | |
| 36 | BK | 132 | |
| 37 | BL | 140 | |
| 38 | BM | 196 | |
| 39 | BN | 174 | |
| 40 | BO | 126 | |
| 41 | BP | 151 | |
| 42 | BQ | 61 | |
| 43 | BR | 97 | |
| 44 | BS | 151 | |
| 45 | BT | 82 | |
| 46 | BU | 119 | |
| 47 | BV | 62 | |
| 48 | BW | 67 | |
| 49 | BX | 153 | |
| 50 | BY | 99 | |
| 51 | BZ | 89 | |
| 52 | Ba | 161 | |
| 53 | Bb | 94 | |

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| Mol | Chain | Length | Quality of chain |
|-----|-------|--------|---|
| 54 | Bc | 56 |  64% 36% |
| 55 | Bd | 51 |  8% 47% 53% |
| 56 | Be | 52 |  10% 58% 27% 15% |
| 57 | Bf | 92 |  12% 71% 29% |
| 58 | AZ | 56 |  89% 48% 41% 11% |

2 Entry composition

There are 60 unique types of molecules in this entry. The entry contains 179267 atoms, of which 29860 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 16S rRNA.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-------|------|-------|------|---------|-------|
| 1 | AA | 1462 | Total | C | N | O | P | 0 | 0 |
| | | | 31324 | 13965 | 5707 | 10190 | 1462 | | |

- Molecule 2 is a protein called Small ribosomal subunit protein uS2.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 2 | AB | 190 | Total | C | N | O | S | 0 | 0 |
| | | | 1492 | 944 | 262 | 281 | 5 | | |

- Molecule 3 is a protein called Small ribosomal subunit protein uS3.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 3 | AC | 186 | Total | C | N | O | S | 0 | 0 |
| | | | 1425 | 898 | 256 | 262 | 9 | | |

- Molecule 4 is a protein called Small ribosomal subunit protein eS1.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 4 | AD | 182 | Total | C | N | O | S | 0 | 0 |
| | | | 1454 | 912 | 262 | 272 | 8 | | |

- Molecule 5 is a protein called Small ribosomal subunit protein uS4.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 5 | AE | 167 | Total | C | N | O | S | 0 | 0 |
| | | | 1351 | 851 | 253 | 242 | 5 | | |

- Molecule 6 is a protein called Small ribosomal subunit protein eS4.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 6 | AF | 234 | Total | C | N | O | S | 0 | 0 |
| | | | 1837 | 1154 | 333 | 345 | 5 | | |

- Molecule 7 is a protein called Small ribosomal subunit protein uS5.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 7 | AG | 198 | Total | C | N | O | S | 0 | 0 |
| | | | 1506 | 954 | 269 | 276 | 7 | | |

- Molecule 8 is a protein called Small ribosomal subunit protein uS7.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 8 | AI | 189 | Total | C | N | O | S | 0 | 0 |
| | | | 1479 | 928 | 274 | 273 | 4 | | |

- Molecule 9 is a protein called Small ribosomal subunit protein uS8.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 9 | AJ | 129 | Total | C | N | O | S | 0 | 0 |
| | | | 981 | 628 | 166 | 183 | 4 | | |

- Molecule 10 is a protein called Small ribosomal subunit protein eS8.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 10 | AK | 125 | Total | C | N | O | S | 0 | 0 |
| | | | 964 | 587 | 196 | 177 | 4 | | |

- Molecule 11 is a protein called Small ribosomal subunit protein uS9.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 11 | AL | 132 | Total | C | N | O | S | 0 | 0 |
| | | | 993 | 617 | 185 | 187 | 4 | | |

- Molecule 12 is a protein called Small ribosomal subunit protein uS11.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 12 | AN | 123 | Total | C | N | O | S | 0 | 0 |
| | | | 911 | 560 | 180 | 167 | 4 | | |

- Molecule 13 is a protein called Small ribosomal subunit protein uS12.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 13 | AO | 142 | Total | C | N | O | S | 0 | 0 |
| | | | 1088 | 679 | 211 | 192 | 6 | | |

- Molecule 14 is a protein called Small ribosomal subunit protein uS13.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 14 | AP | 132 | Total | C | N | O | S | 0 | 0 |
| | | | 1033 | 649 | 198 | 184 | 2 | | |

There are 20 discrepancies between the modelled and reference sequences:

| Chain | Residue | Modelled | Actual | Comment | Reference |
|-------|---------|----------|--------|-----------------------|------------|
| AP | -19 | MET | - | initiating methionine | UNP Q8TRR2 |
| AP | -18 | ARG | - | expression tag | UNP Q8TRR2 |
| AP | -17 | ALA | - | expression tag | UNP Q8TRR2 |
| AP | -16 | ALA | - | expression tag | UNP Q8TRR2 |
| AP | -15 | PRO | - | expression tag | UNP Q8TRR2 |
| AP | -14 | GLU | - | expression tag | UNP Q8TRR2 |
| AP | -13 | PHE | - | expression tag | UNP Q8TRR2 |
| AP | -12 | SER | - | expression tag | UNP Q8TRR2 |
| AP | -11 | ALA | - | expression tag | UNP Q8TRR2 |
| AP | -10 | TYR | - | expression tag | UNP Q8TRR2 |
| AP | -9 | SER | - | expression tag | UNP Q8TRR2 |
| AP | -8 | LEU | - | expression tag | UNP Q8TRR2 |
| AP | -7 | ARG | - | expression tag | UNP Q8TRR2 |
| AP | -6 | VAL | - | expression tag | UNP Q8TRR2 |
| AP | -5 | PHE | - | expression tag | UNP Q8TRR2 |
| AP | -4 | ASP | - | expression tag | UNP Q8TRR2 |
| AP | -3 | ARG | - | expression tag | UNP Q8TRR2 |
| AP | -2 | GLY | - | expression tag | UNP Q8TRR2 |
| AP | -1 | TYR | - | expression tag | UNP Q8TRR2 |
| AP | 0 | ARG | - | expression tag | UNP Q8TRR2 |

- Molecule 15 is a protein called Small ribosomal subunit protein uS14.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 15 | AR | 44 | Total | C | N | O | S | 0 | 0 |
| | | | 366 | 226 | 73 | 62 | 5 | | |

- Molecule 16 is a protein called Small ribosomal subunit protein uS15.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 16 | AQ | 152 | Total | C | N | O | S | 0 | 0 |
| | | | 1226 | 777 | 223 | 220 | 6 | | |

- Molecule 17 is a protein called Small ribosomal subunit protein uS17A.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 17 | AS | 109 | Total | C | N | O | S | 0 | 0 |
| | | | 849 | 533 | 152 | 156 | 8 | | |

- Molecule 18 is a protein called Small ribosomal subunit protein uS19.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 18 | AU | 95 | Total | C | N | O | S | 0 | 0 |
| | | | 775 | 491 | 142 | 139 | 3 | | |

- Molecule 19 is a protein called Small ribosomal subunit protein eS19.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 19 | AV | 149 | Total | C | N | O | S | 0 | 0 |
| | | | 1163 | 739 | 204 | 215 | 5 | | |

- Molecule 20 is a protein called Small ribosomal subunit protein eS24.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 20 | AW | 95 | Total | C | N | O | S | 0 | 0 |
| | | | 766 | 481 | 136 | 145 | 4 | | |

- Molecule 21 is a protein called Small ribosomal subunit protein eS27.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 21 | AX | 60 | Total | C | N | O | S | 0 | 0 |
| | | | 468 | 288 | 85 | 91 | 4 | | |

- Molecule 22 is a protein called Small ribosomal subunit protein eS28.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 22 | AY | 63 | Total | C | N | O | S | 0 | 0 |
| | | | 471 | 289 | 90 | 88 | 4 | | |

- Molecule 23 is a protein called Small ribosomal subunit protein eS6.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 23 | AH | 124 | Total | C | N | O | S | 0 | 0 |
| | | | 930 | 591 | 168 | 167 | 4 | | |

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

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 24 | AM | 102 | Total | C | N | O | S | 0 | 0 |
| | | | 796 | 497 | 150 | 145 | 4 | | |

- Molecule 25 is a protein called Small ribosomal subunit protein eS17.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 25 | AT | 58 | Total | C | N | O | S | 0 | 0 |
| | | | 477 | 302 | 87 | 87 | 1 | | |

- Molecule 26 is a RNA chain called 23S rRNA.

| Mol | Chain | Residues | Atoms | | | | | | AltConf | Trace |
|-----|-------|----------|-------|-------|-------|-------|-------|------|---------|-------|
| 26 | BA | 2776 | Total | C | H | N | O | P | 0 | 0 |
| | | | 89273 | 26520 | 29860 | 10811 | 19307 | 2775 | | |

- Molecule 27 is a RNA chain called 5S rRNA.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|-----|---------|-------|
| 27 | BB | 128 | Total | C | N | O | P | 0 | 0 |
| | | | 2720 | 1214 | 481 | 897 | 128 | | |

- Molecule 28 is a protein called Large ribosomal subunit protein uL2.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 28 | BC | 238 | Total | C | N | O | S | 0 | 0 |
| | | | 1808 | 1129 | 350 | 321 | 8 | | |

- Molecule 29 is a protein called Large ribosomal subunit protein uL3.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 29 | BD | 337 | Total | C | N | O | S | 0 | 0 |
| | | | 2597 | 1639 | 474 | 476 | 8 | | |

- Molecule 30 is a protein called Large ribosomal subunit protein uL4.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|------|-----|-----|---|---------|-------|
| 30 | BE | 252 | Total | C | N | O | S | 0 | 0 |
| | | | 1930 | 1208 | 368 | 353 | 1 | | |

- Molecule 31 is a protein called Large ribosomal subunit protein uL5.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 31 | BF | 165 | Total | C | N | O | S | 0 | 0 |
| | | | 1289 | 812 | 234 | 235 | 8 | | |

- Molecule 32 is a protein called Large ribosomal subunit protein uL6.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 32 | BG | 176 | Total | C | N | O | S | 0 | 0 |
| | | | 1371 | 876 | 235 | 254 | 6 | | |

- Molecule 33 is a protein called Large ribosomal subunit protein eL8.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 33 | BH | 115 | Total | C | N | O | S | 0 | 0 |
| | | | 857 | 541 | 144 | 170 | 2 | | |

- Molecule 34 is a protein called Large ribosomal subunit protein uL16.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 34 | BI | 159 | Total | C | N | O | S | 0 | 0 |
| | | | 1261 | 794 | 240 | 218 | 9 | | |

- Molecule 35 is a protein called Large ribosomal subunit protein uL13.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 35 | BJ | 138 | Total | C | N | O | S | 0 | 0 |
| | | | 1086 | 683 | 200 | 199 | 4 | | |

- Molecule 36 is a protein called Large ribosomal subunit protein uL14.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 36 | BK | 132 | Total | C | N | O | S | 0 | 0 |
| | | | 999 | 623 | 185 | 182 | 9 | | |

- Molecule 37 is a protein called Large ribosomal subunit protein uL15.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 37 | BL | 140 | Total | C | N | O | S | 0 | 0 |
| | | | 1058 | 643 | 204 | 205 | 6 | | |

- Molecule 38 is a protein called Large ribosomal subunit protein eL15.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 38 | BM | 196 | Total | C | N | O | S | 0 | 0 |
| | | | 1593 | 986 | 329 | 273 | 5 | | |

- Molecule 39 is a protein called Large ribosomal subunit protein uL18.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 39 | BN | 174 | Total | C | N | O | S | 0 | 0 |
| | | | 1356 | 853 | 242 | 259 | 2 | | |

- Molecule 40 is a protein called Large ribosomal subunit protein eL18.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 40 | BO | 126 | Total | C | N | O | S | 0 | 0 |
| | | | 962 | 608 | 173 | 178 | 3 | | |

- Molecule 41 is a protein called Large ribosomal subunit protein eL19.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 41 | BP | 151 | Total | C | N | O | S | 0 | 0 |
| | | | 1195 | 739 | 242 | 210 | 4 | | |

- Molecule 42 is a protein called Large ribosomal subunit protein eL20.

| Mol | Chain | Residues | Atoms | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---------|-------|
| 42 | BQ | 57 | Total | C | N | O | 0 | 0 |
| | | | 457 | 289 | 79 | 89 | | |

- Molecule 43 is a protein called Large ribosomal subunit protein eL21.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 43 | BR | 96 | Total | C | N | O | S | 0 | 0 |
| | | | 766 | 475 | 146 | 141 | 4 | | |

- Molecule 44 is a protein called Large ribosomal subunit protein uL22.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|----|---------|-------|
| 44 | BS | 151 | Total | C | N | O | S | 0 | 0 |
| | | | 1169 | 729 | 218 | 212 | 10 | | |

- Molecule 45 is a protein called Large ribosomal subunit protein uL23.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 45 | BT | 82 | Total | C | N | O | S | 0 | 0 |
| | | | 656 | 418 | 108 | 122 | 8 | | |

- Molecule 46 is a protein called Large ribosomal subunit protein uL24.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 46 | BU | 119 | Total | C | N | O | S | 0 | 0 |
| | | | 910 | 564 | 170 | 169 | 7 | | |

- Molecule 47 is a protein called Large ribosomal subunit protein eL24.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 47 | BV | 62 | Total | C | N | O | S | 0 | 0 |
| | | | 499 | 316 | 88 | 87 | 8 | | |

- Molecule 48 is a protein called Large ribosomal subunit protein uL29.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 48 | BW | 67 | Total | C | N | O | S | 0 | 0 |
| | | | 532 | 321 | 103 | 106 | 2 | | |

- Molecule 49 is a protein called Large ribosomal subunit protein uL30.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 49 | BX | 153 | Total | C | N | O | S | 0 | 0 |
| | | | 1237 | 779 | 230 | 222 | 6 | | |

- Molecule 50 is a protein called Large ribosomal subunit protein eL30.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 50 | BY | 92 | Total | C | N | O | S | 0 | 0 |
| | | | 658 | 415 | 108 | 131 | 4 | | |

- Molecule 51 is a protein called Large ribosomal subunit protein eL31.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 51 | BZ | 86 | Total | C | N | O | S | 0 | 0 |
| | | | 703 | 446 | 131 | 123 | 3 | | |

- Molecule 52 is a protein called Large ribosomal subunit protein eL32.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 52 | Ba | 132 | Total | C | N | O | S | 0 | 0 |
| | | | 1028 | 645 | 197 | 184 | 2 | | |

There are 3 discrepancies between the modelled and reference sequences:

| Chain | Residue | Modelled | Actual | Comment | Reference |
|-------|---------|----------|--------|-----------------------|----------------|
| Ba | -2 | MET | - | initiating methionine | UNP A0A832W8Z7 |
| Ba | -1 | ILE | - | expression tag | UNP A0A832W8Z7 |
| Ba | 0 | MET | - | expression tag | UNP A0A832W8Z7 |

- Molecule 53 is a protein called Large ribosomal subunit protein eL43.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 53 | Bb | 94 | Total | C | N | O | S | 0 | 0 |
| | | | 736 | 459 | 145 | 125 | 7 | | |

- Molecule 54 is a protein called Large ribosomal subunit protein eL37.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 54 | Bc | 56 | Total | C | N | O | S | 0 | 0 |
| | | | 445 | 269 | 92 | 76 | 8 | | |

- Molecule 55 is a protein called Large ribosomal subunit protein eL39.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|----|---|---------|-------|
| 55 | Bd | 51 | Total | C | N | O | S | 0 | 0 |
| | | | 439 | 272 | 101 | 64 | 2 | | |

- Molecule 56 is a protein called Large ribosomal subunit protein eL40.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 56 | Be | 44 | Total | C | N | O | S | 0 | 0 |
| | | | 353 | 215 | 74 | 59 | 5 | | |

There are 3 discrepancies between the modelled and reference sequences:

| Chain | Residue | Modelled | Actual | Comment | Reference |
|-------|---------|----------|--------|-----------------------|------------|
| Be | -2 | MET | - | initiating methionine | UNP Q8TJ19 |
| Be | -1 | THR | - | expression tag | UNP Q8TJ19 |
| Be | 0 | LYS | - | expression tag | UNP Q8TJ19 |

- Molecule 57 is a protein called Large ribosomal subunit protein eL42.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|-----|-----|---|---------|-------|
| 57 | Bf | 92 | Total | C | N | O | S | 0 | 0 |
| | | | 760 | 480 | 151 | 122 | 7 | | |

- Molecule 58 is a protein called Putative zinc finger domain-containing protein.

| Mol | Chain | Residues | Atoms | | | | | AltConf | Trace |
|-----|-------|----------|-------|-----|----|----|---|---------|-------|
| 58 | AZ | 50 | Total | C | N | O | S | 0 | 0 |
| | | | 379 | 236 | 67 | 68 | 8 | | |

- Molecule 59 is MAGNESIUM ION (CCD ID: MG) (formula: Mg).

| Mol | Chain | Residues | Atoms | | AltConf |
|-----|-------|----------|-------|----|---------|
| 59 | AA | 6 | Total | Mg | 0 |
| | | | 6 | 6 | |
| 59 | AN | 1 | Total | Mg | 0 |
| | | | 1 | 1 | |
| 59 | AS | 1 | Total | Mg | 0 |
| | | | 1 | 1 | |
| 59 | BA | 42 | Total | Mg | 0 |
| | | | 42 | 42 | |
| 59 | Bc | 1 | Total | Mg | 0 |
| | | | 1 | 1 | |

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

| Mol | Chain | Residues | Atoms | | AltConf |
|-----|-------|----------|-------|----|---------|
| 60 | AG | 1 | Total | Zn | 0 |
| | | | 1 | 1 | |
| 60 | AR | 1 | Total | Zn | 0 |
| | | | 1 | 1 | |
| 60 | AS | 1 | Total | Zn | 0 |
| | | | 1 | 1 | |
| 60 | AX | 1 | Total | Zn | 0 |
| | | | 1 | 1 | |
| 60 | BV | 1 | Total | Zn | 0 |
| | | | 1 | 1 | |
| 60 | Bb | 1 | Total | Zn | 0 |
| | | | 1 | 1 | |
| 60 | Bc | 1 | Total | Zn | 0 |
| | | | 1 | 1 | |
| 60 | Be | 1 | Total | Zn | 0 |
| | | | 1 | 1 | |

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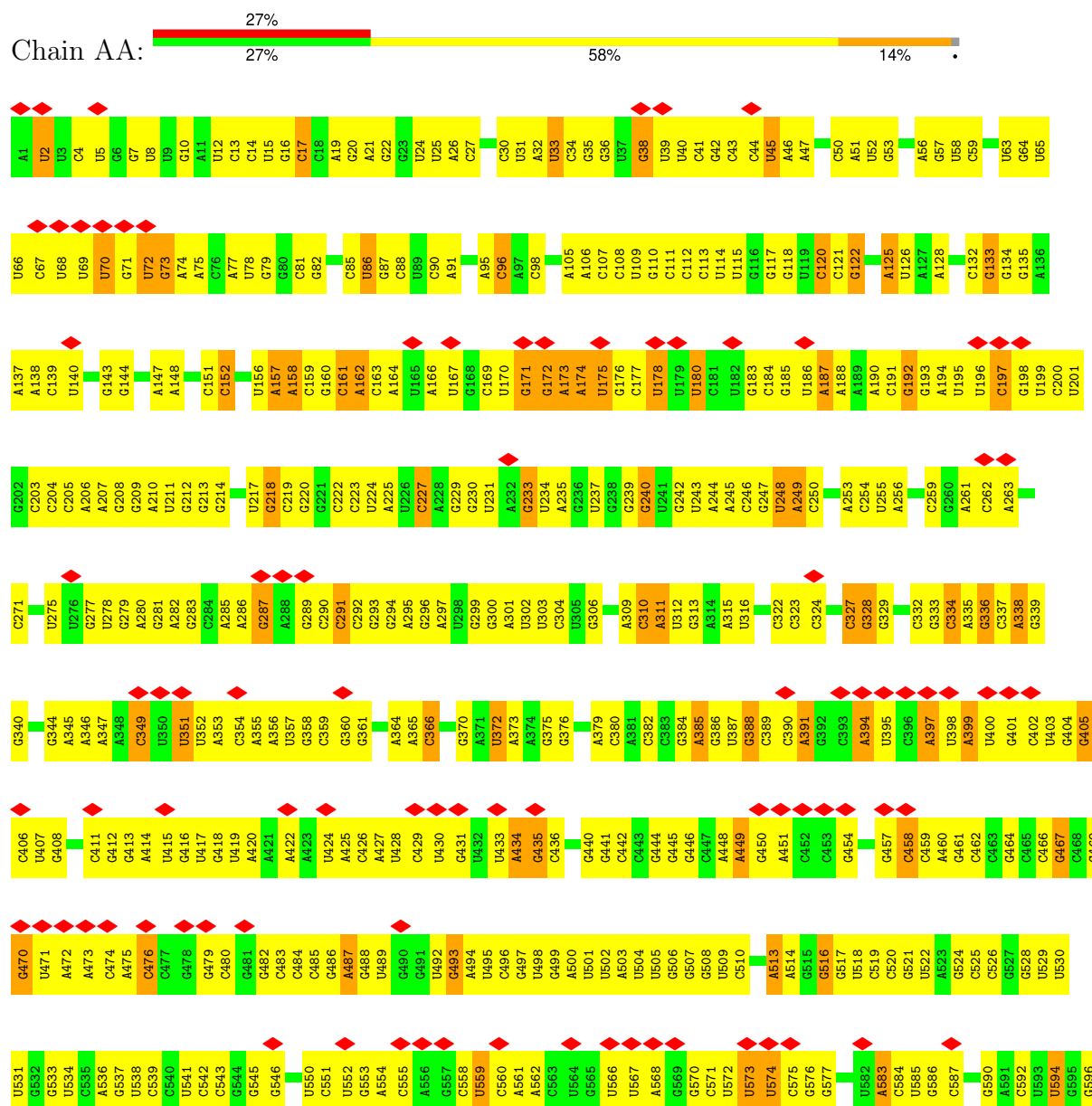
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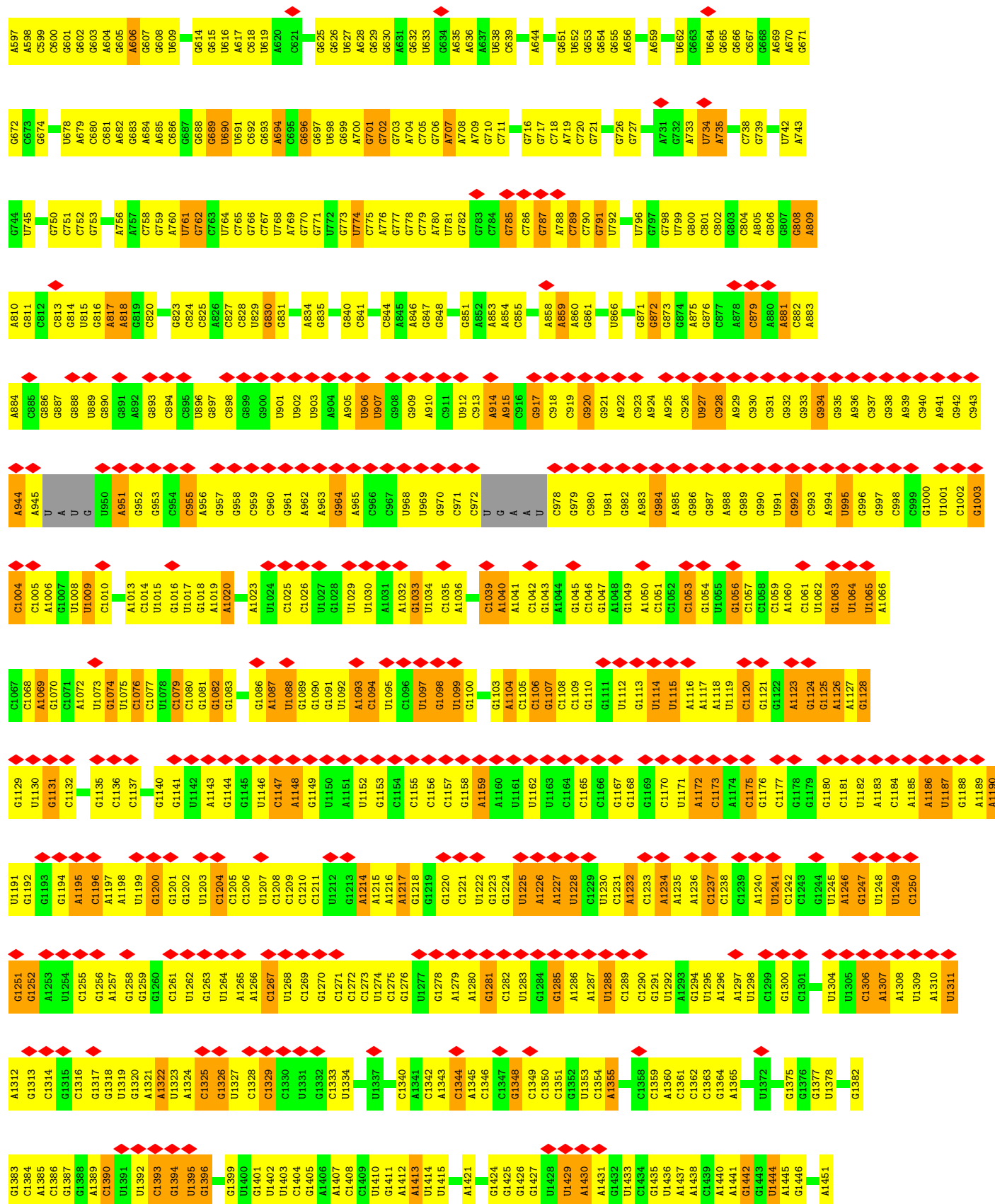
| Mol | Chain | Residues | Atoms | | AltConf |
|-----|-------|----------|-------|----|---------|
| | | | Total | Zn | |
| 60 | Bf | 1 | 1 | 1 | 0 |

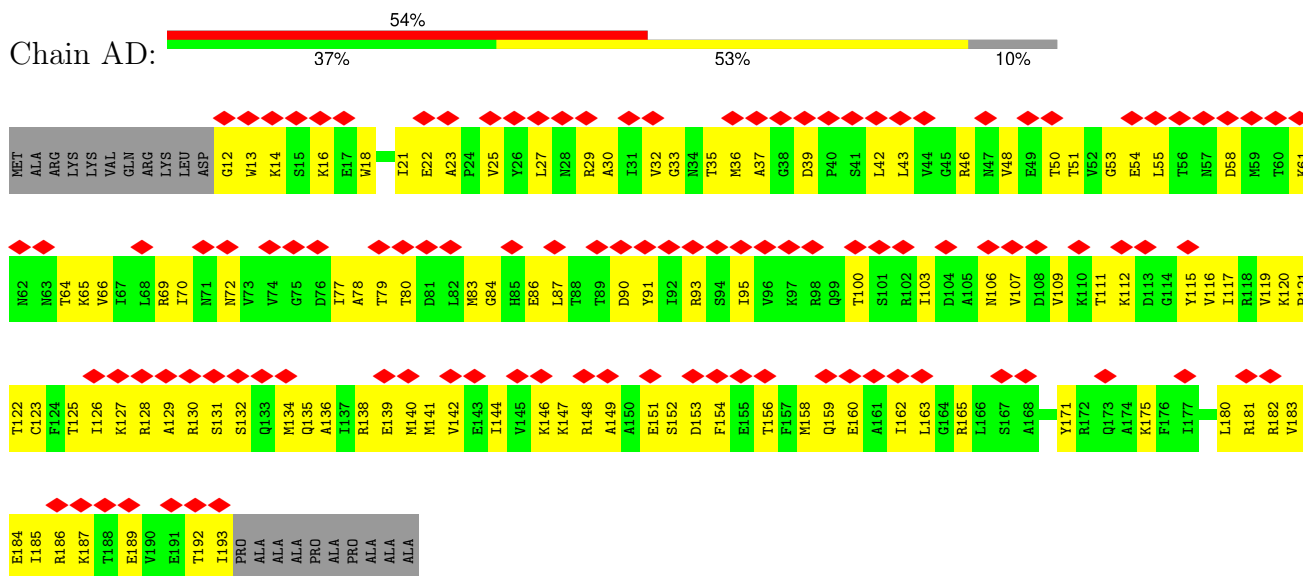
3 Residue-property plots

These plots are drawn for all protein, RNA, DNA and oligosaccharide chains in the entry. The first graphic for a chain summarises the proportions of the various outlier classes displayed in the second graphic. The second graphic shows the sequence view annotated by issues in geometry and 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.

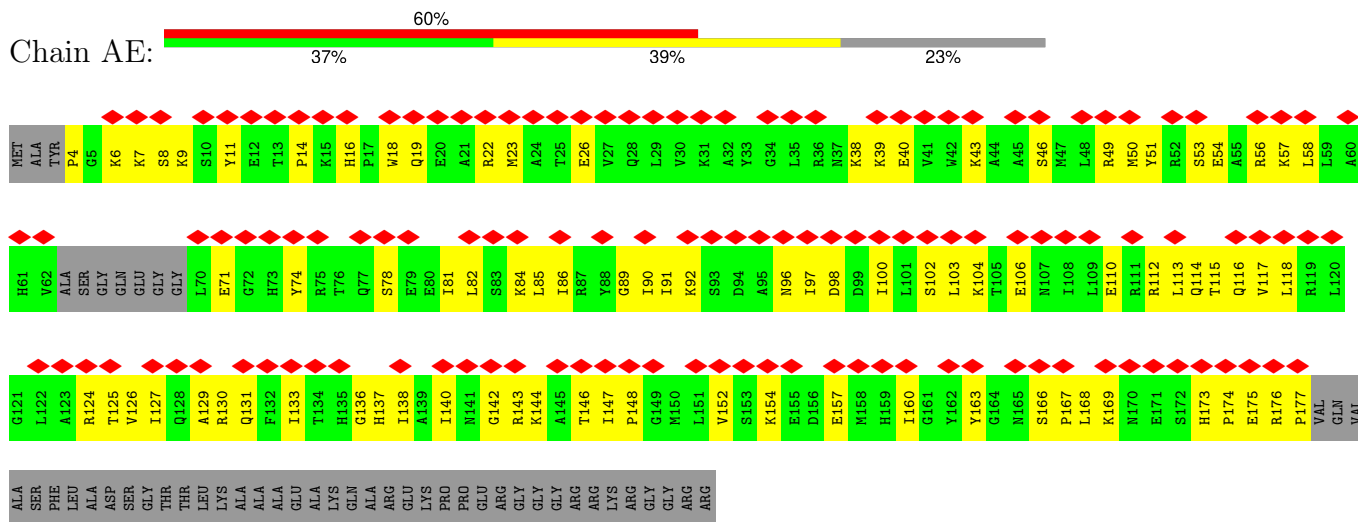
• Molecule 1: 16S rRNA



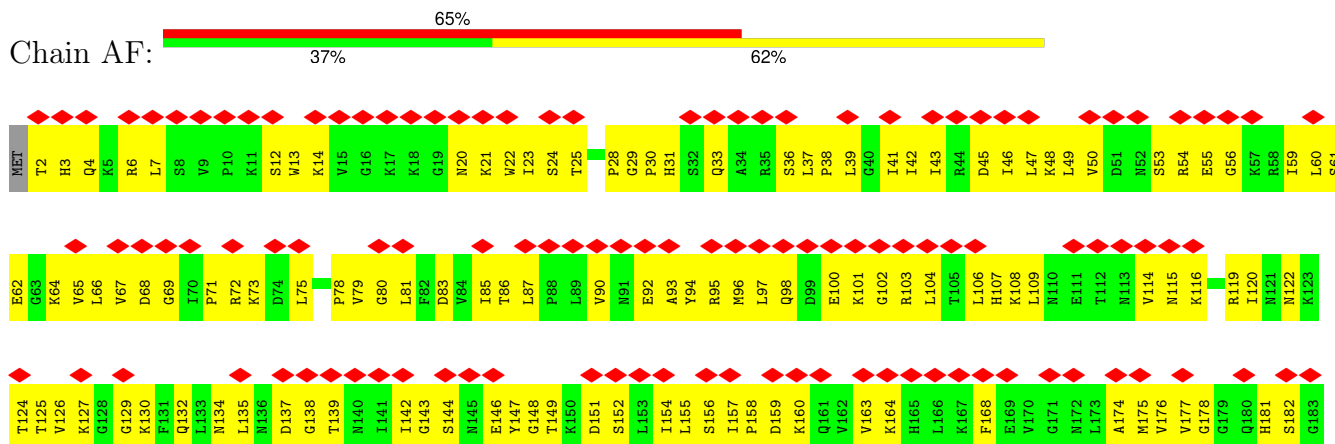




- Molecule 5: Small ribosomal subunit protein uS4

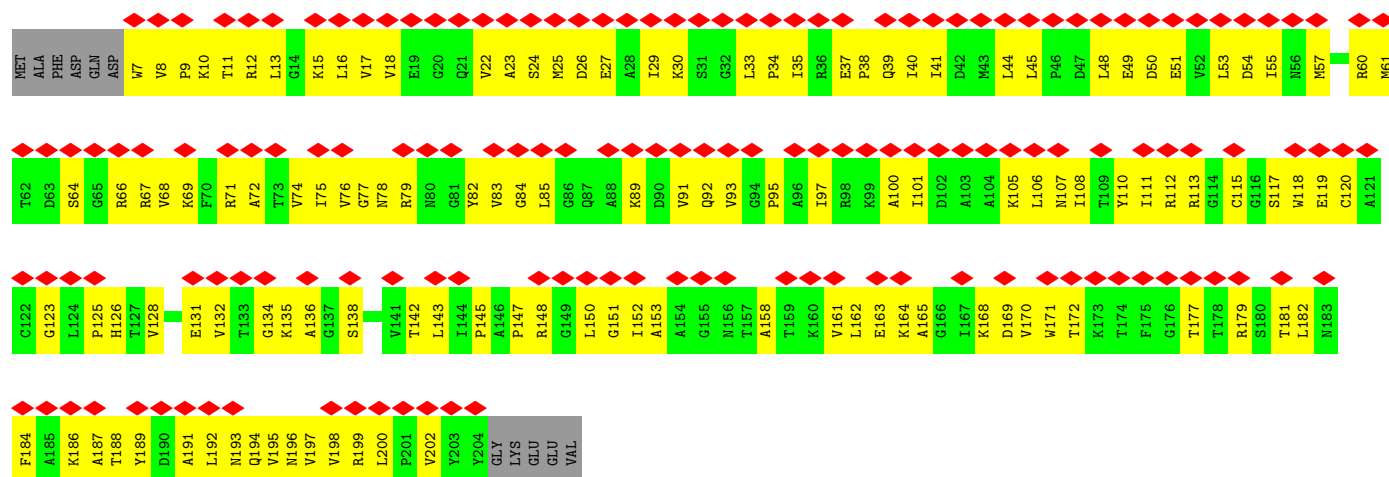
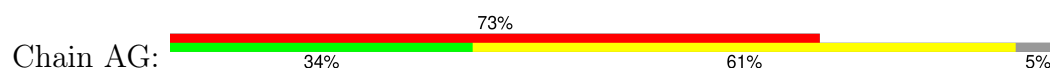


- Molecule 6: Small ribosomal subunit protein eS4

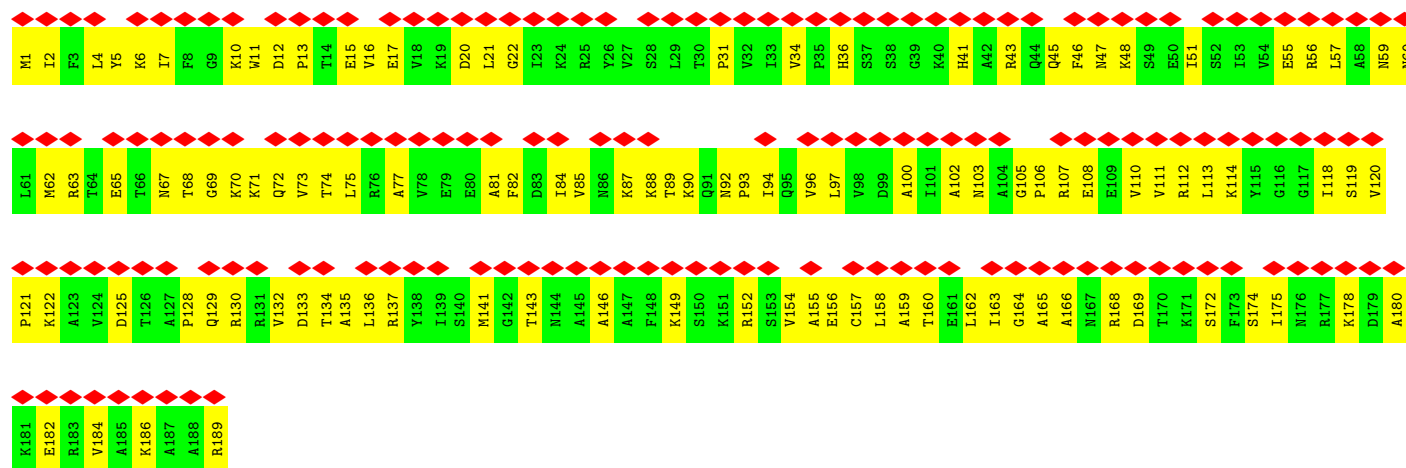
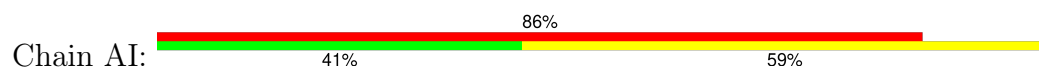




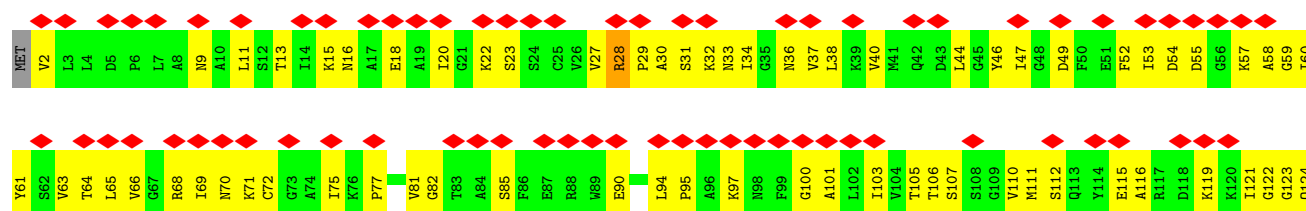
• Molecule 7: Small ribosomal subunit protein uS5



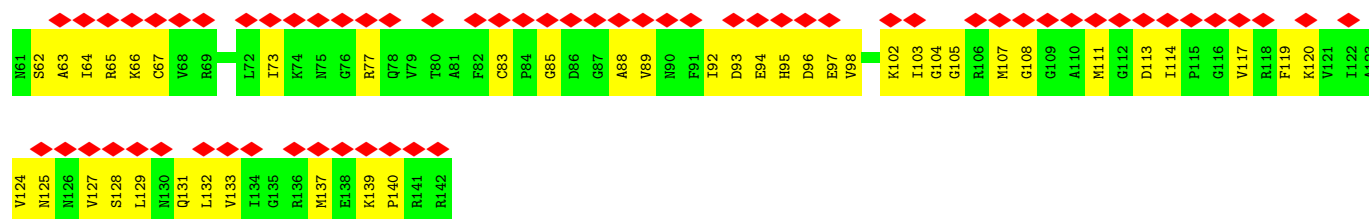
• Molecule 8: Small ribosomal subunit protein uS7



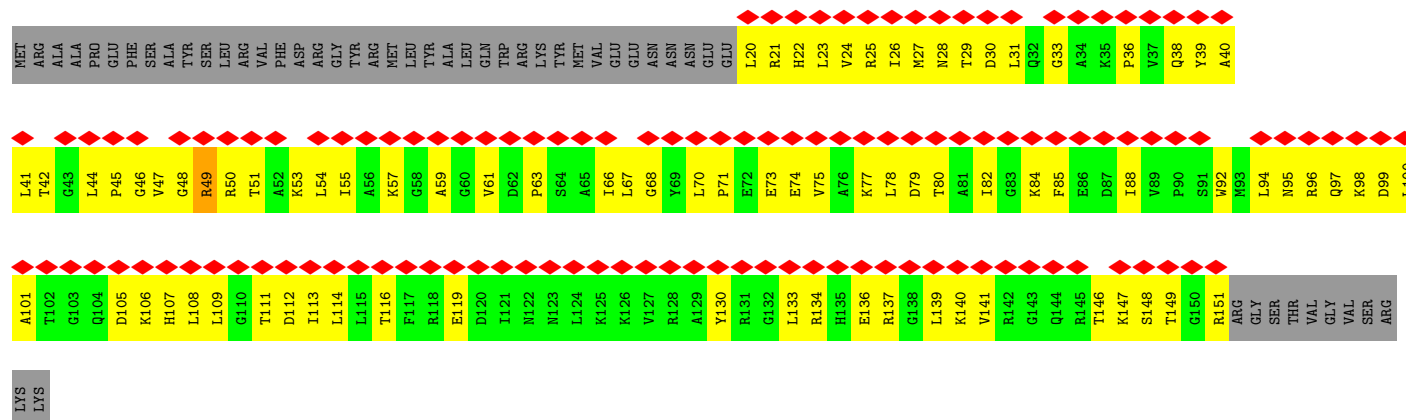
• Molecule 9: Small ribosomal subunit protein uS8



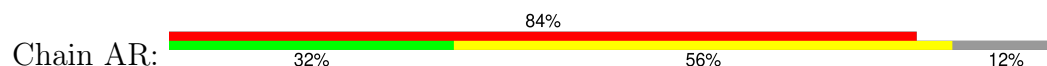




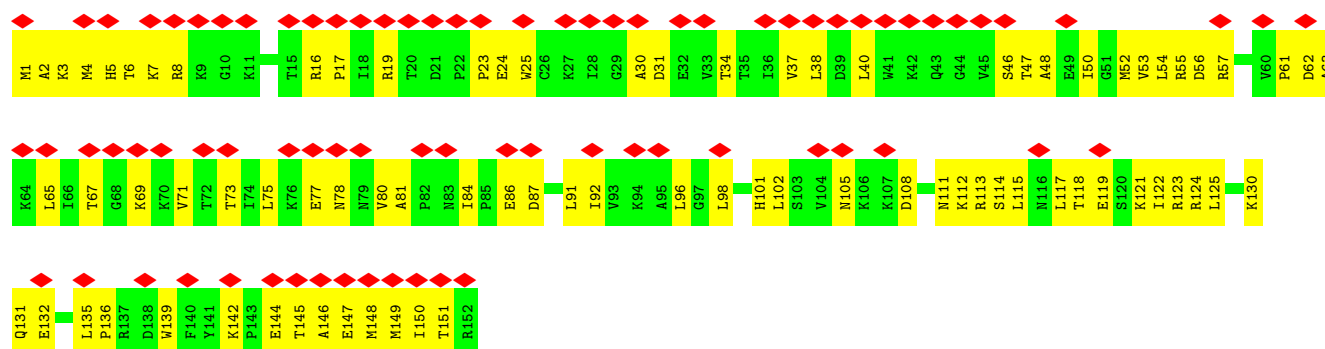
• Molecule 14: Small ribosomal subunit protein uS13



• Molecule 15: Small ribosomal subunit protein uS14

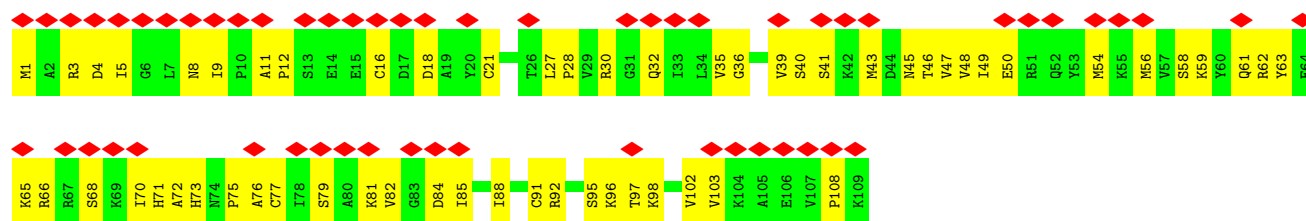


• Molecule 16: Small ribosomal subunit protein uS15

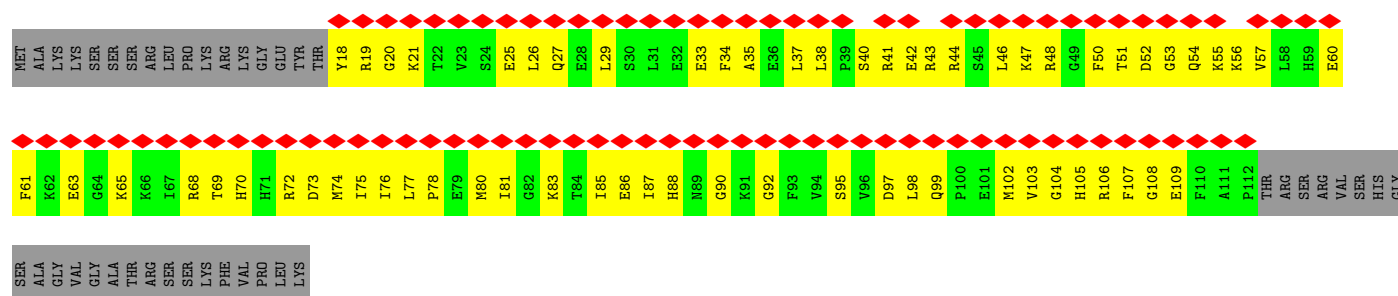


• Molecule 17: Small ribosomal subunit protein uS17A

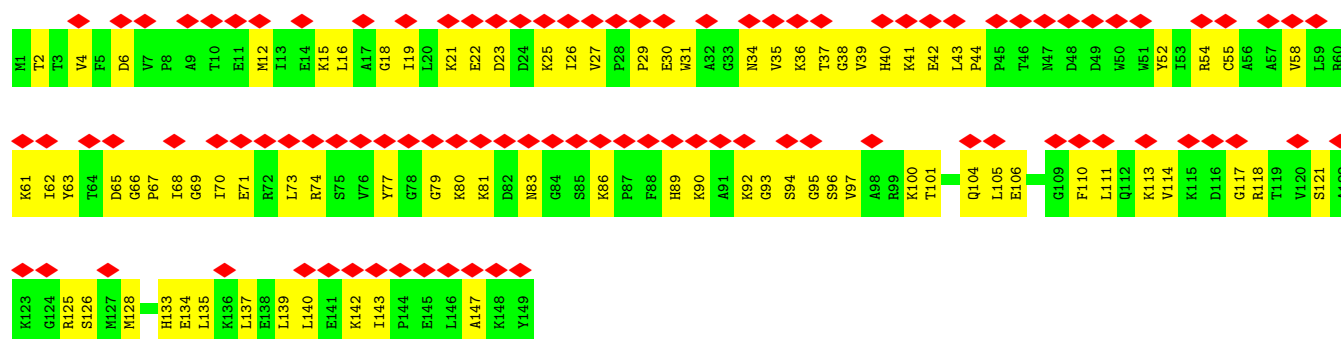




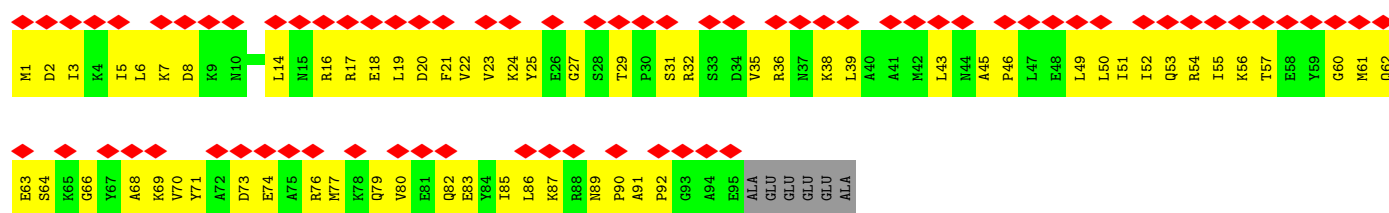
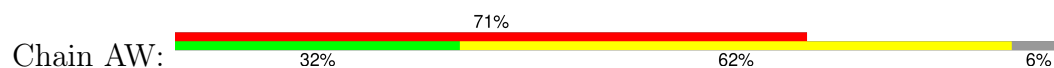
• Molecule 18: Small ribosomal subunit protein uS19



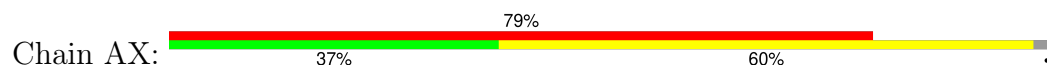
• Molecule 19: Small ribosomal subunit protein eS19

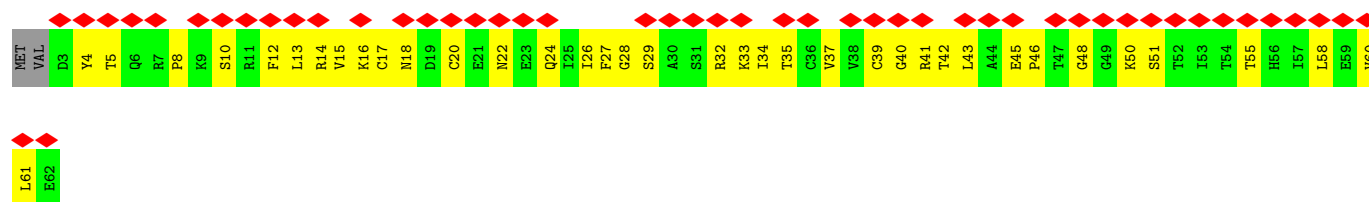


• Molecule 20: Small ribosomal subunit protein eS24



• Molecule 21: Small ribosomal subunit protein eS27

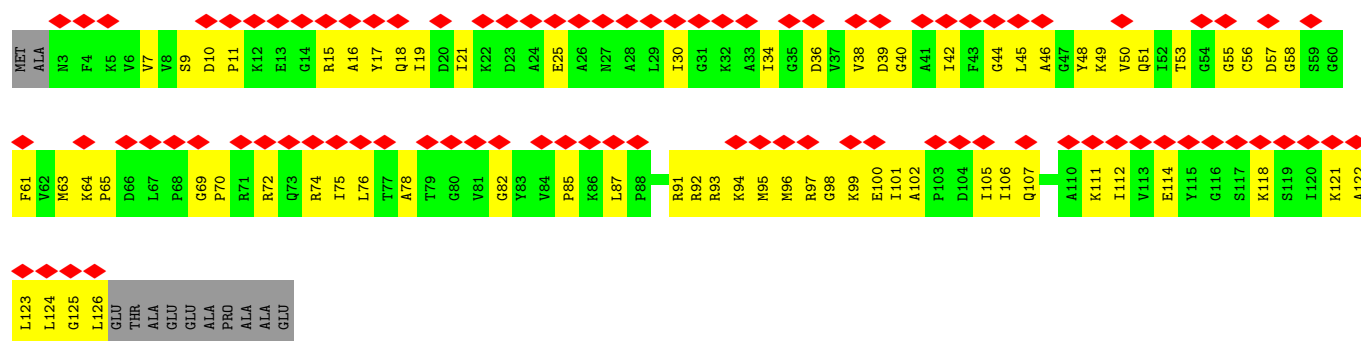




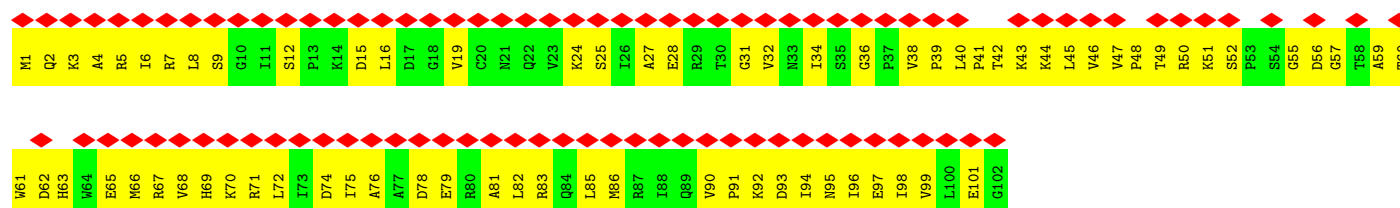
- Molecule 22: Small ribosomal subunit protein eS28



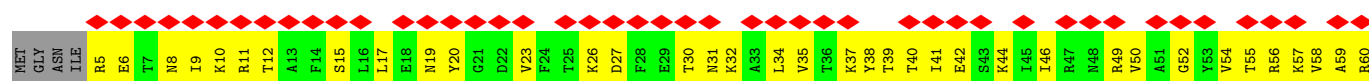
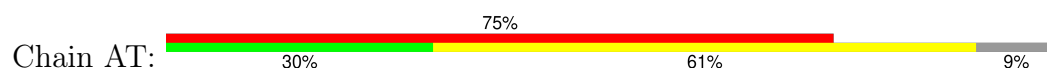
- Molecule 23: Small ribosomal subunit protein eS6



- Molecule 24: Small ribosomal subunit protein uS10

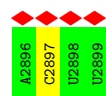


- Molecule 25: Small ribosomal subunit protein eS17

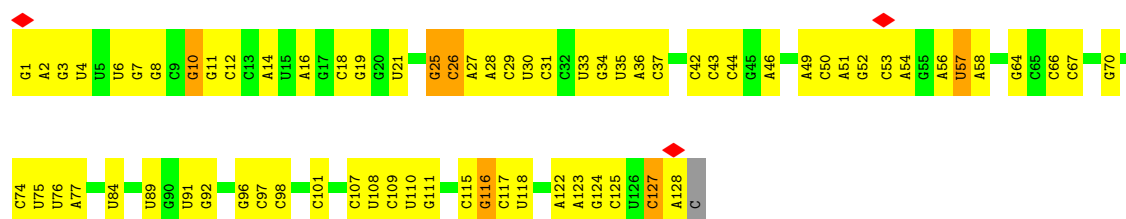




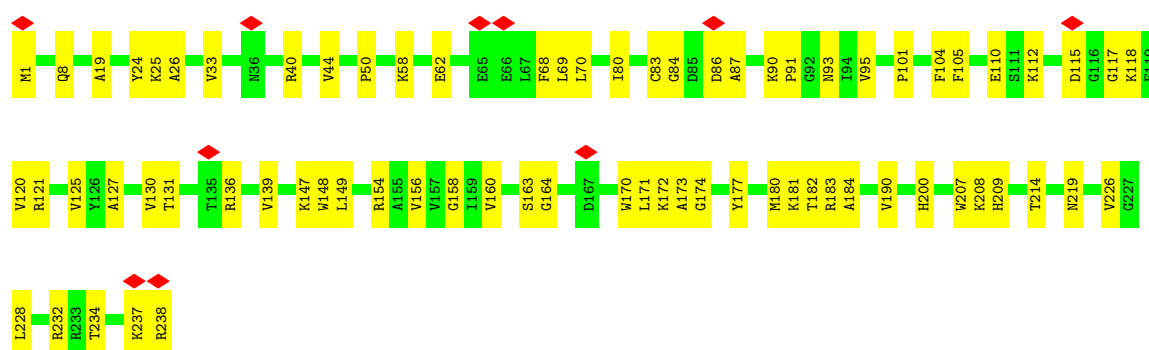




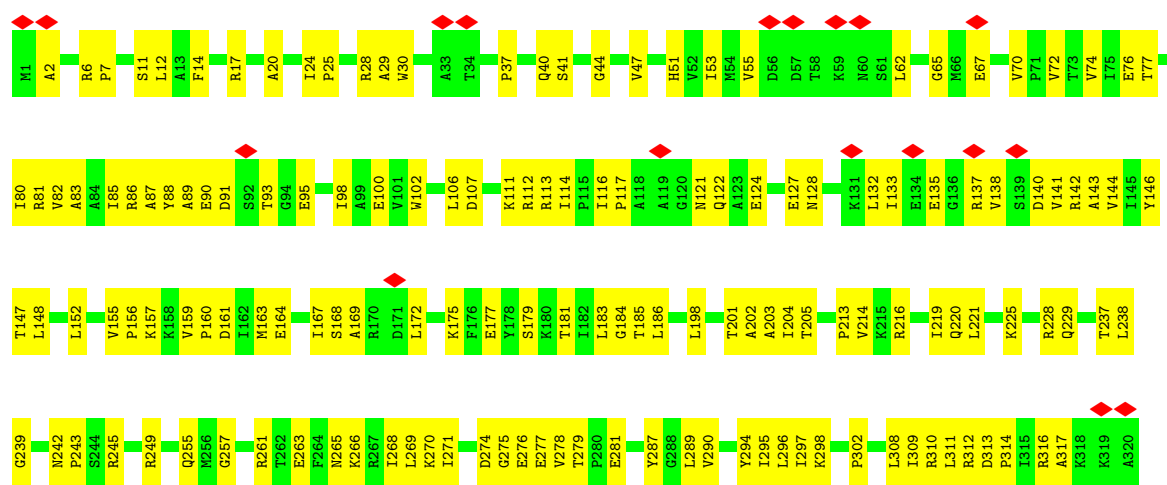
• Molecule 27: 5S rRNA

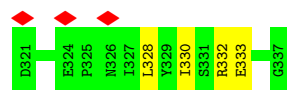


• Molecule 28: Large ribosomal subunit protein uL2

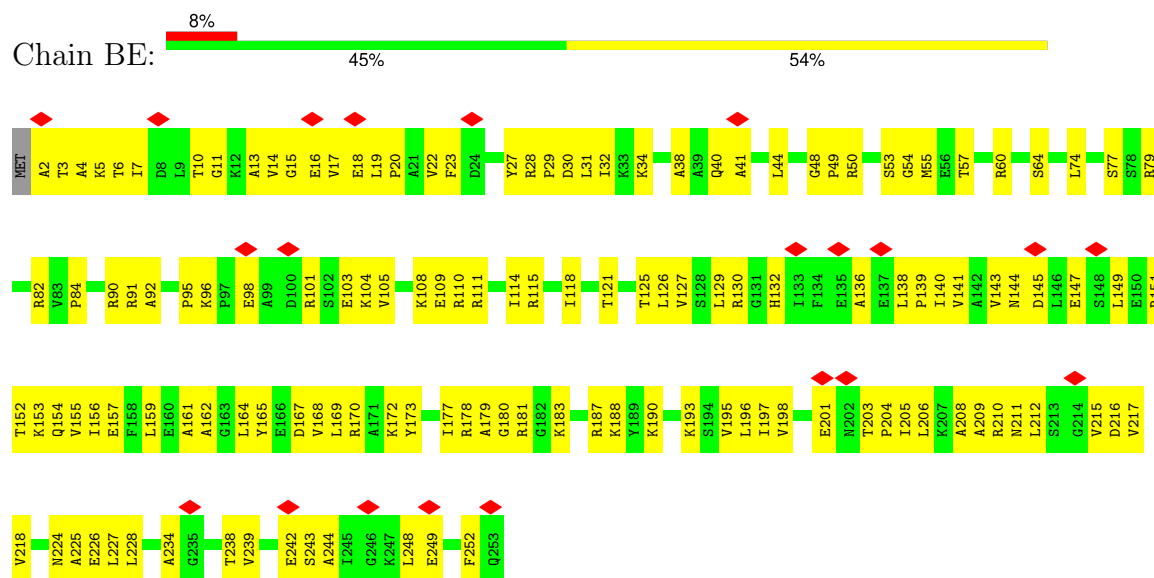


• Molecule 29: Large ribosomal subunit protein uL3

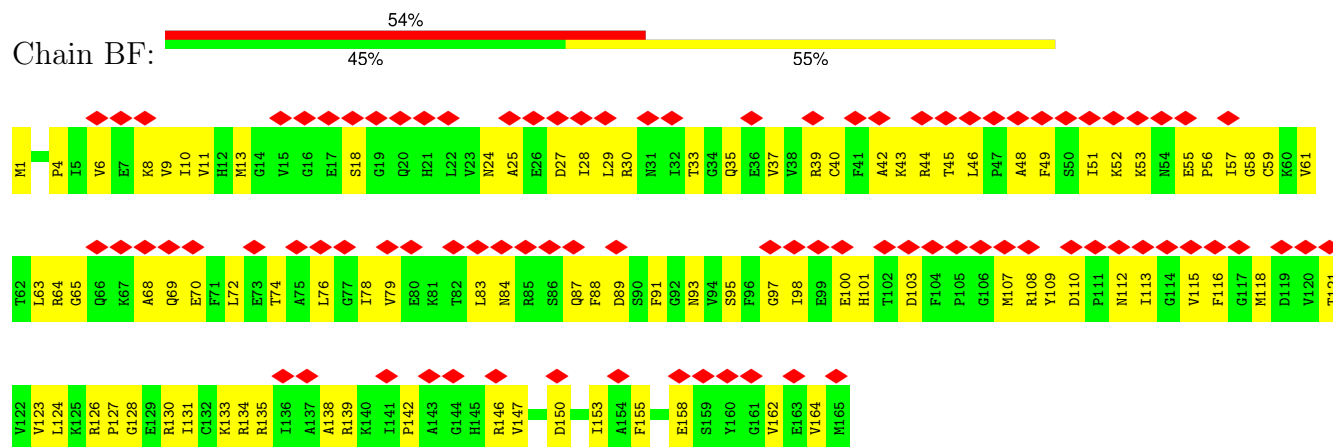




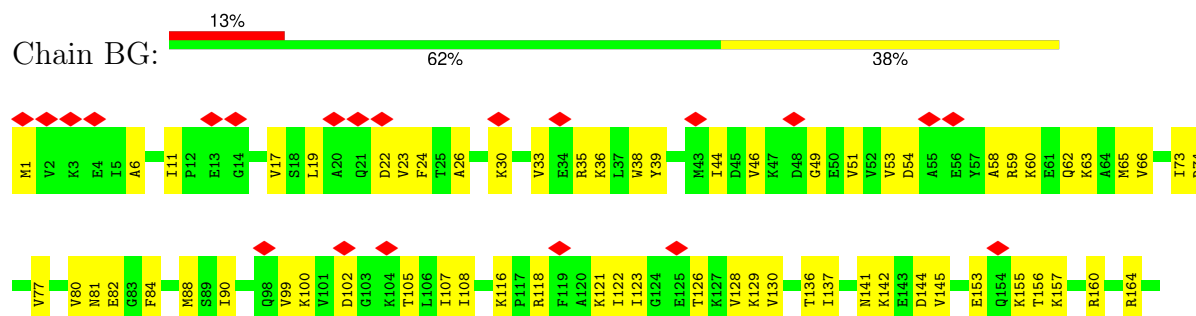
• Molecule 30: Large ribosomal subunit protein uL4



• Molecule 31: Large ribosomal subunit protein uL5

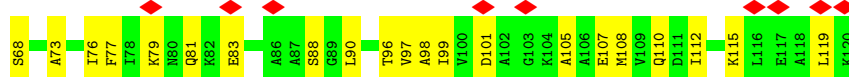


• Molecule 32: Large ribosomal subunit protein uL6

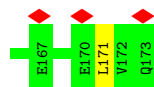
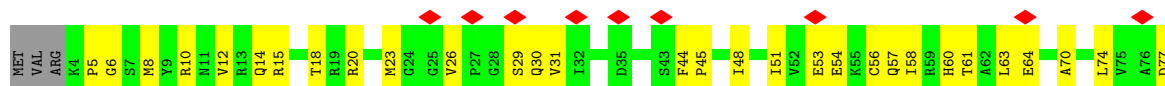




- Molecule 33: Large ribosomal subunit protein eL8



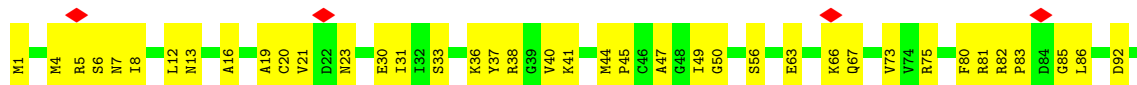
- Molecule 34: Large ribosomal subunit protein uL16

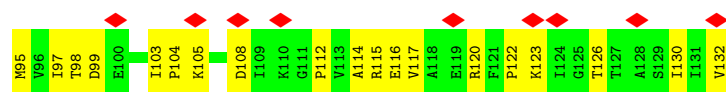


- Molecule 35: Large ribosomal subunit protein uL13

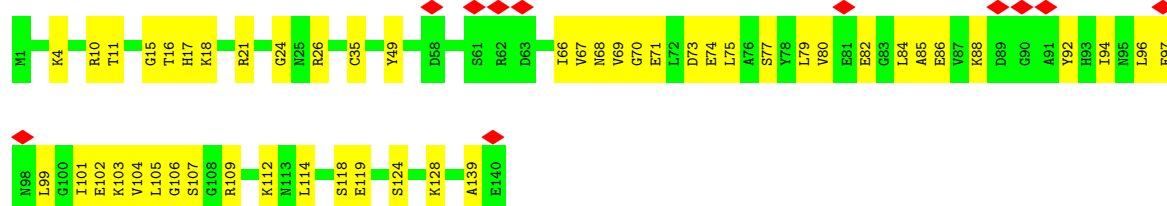


- Molecule 36: Large ribosomal subunit protein uL14

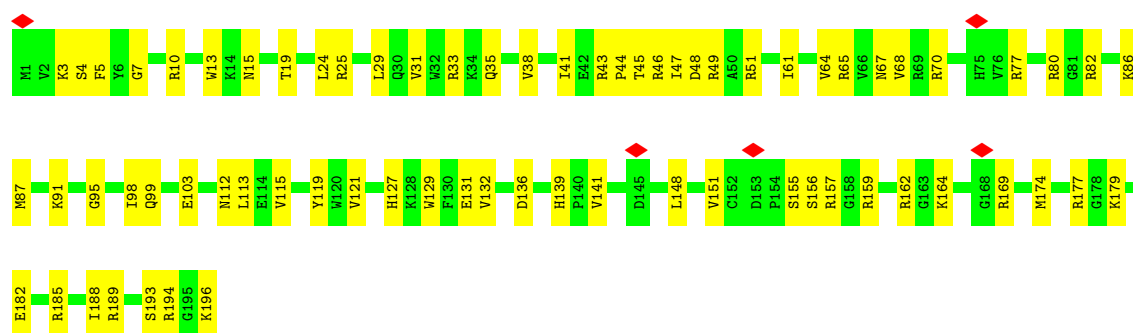




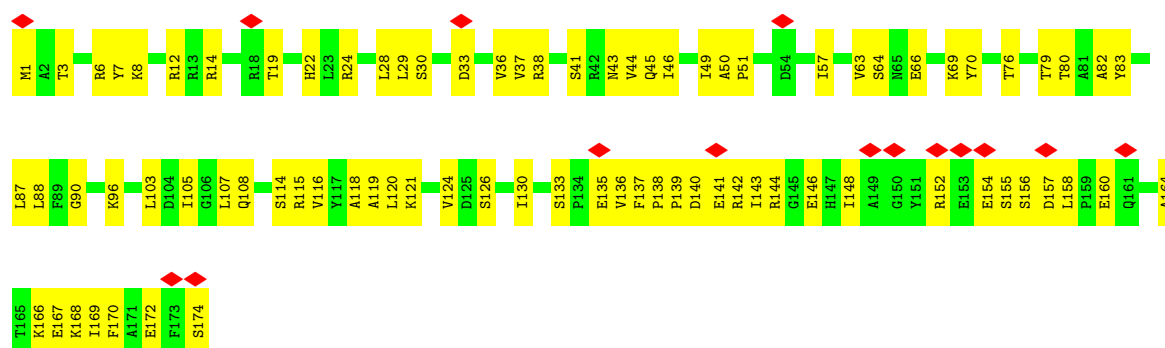
- Molecule 37: Large ribosomal subunit protein uL15



- Molecule 38: Large ribosomal subunit protein eL15

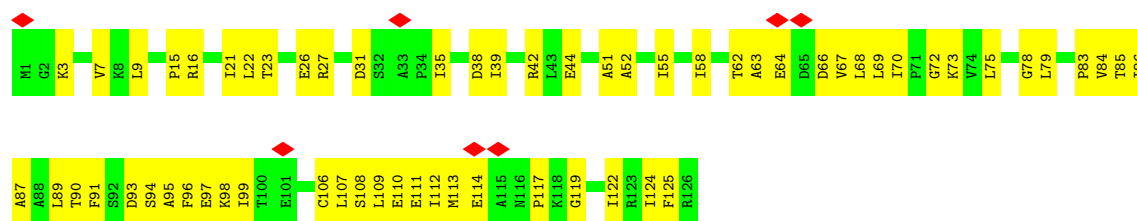


- Molecule 39: Large ribosomal subunit protein uL18



- Molecule 40: Large ribosomal subunit protein eL18

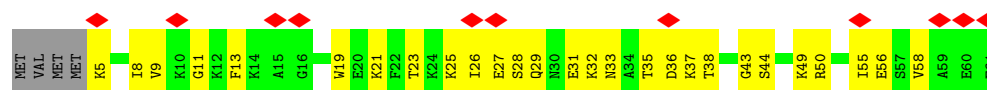




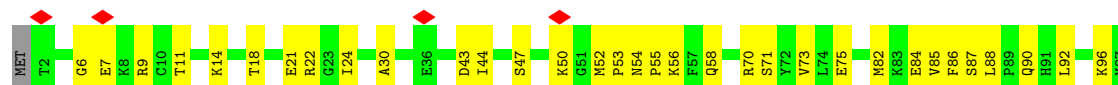
- Molecule 41: Large ribosomal subunit protein eL19



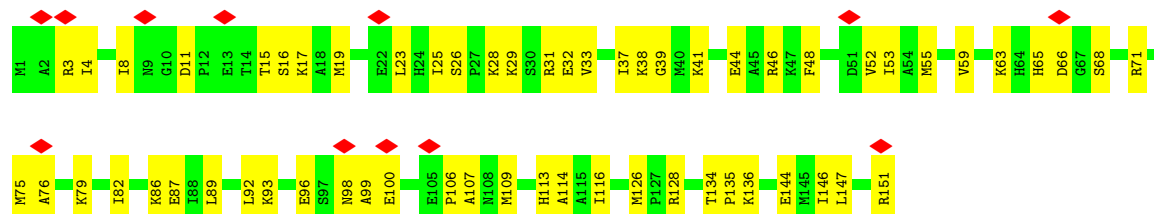
- Molecule 42: Large ribosomal subunit protein eL20



- Molecule 43: Large ribosomal subunit protein eL21

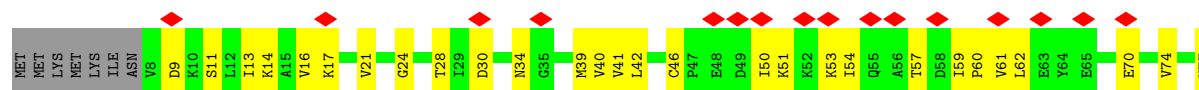


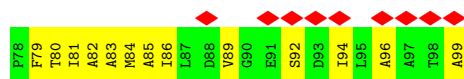
- Molecule 44: Large ribosomal subunit protein uL22



- Molecule 45: Large ribosomal subunit protein uL23



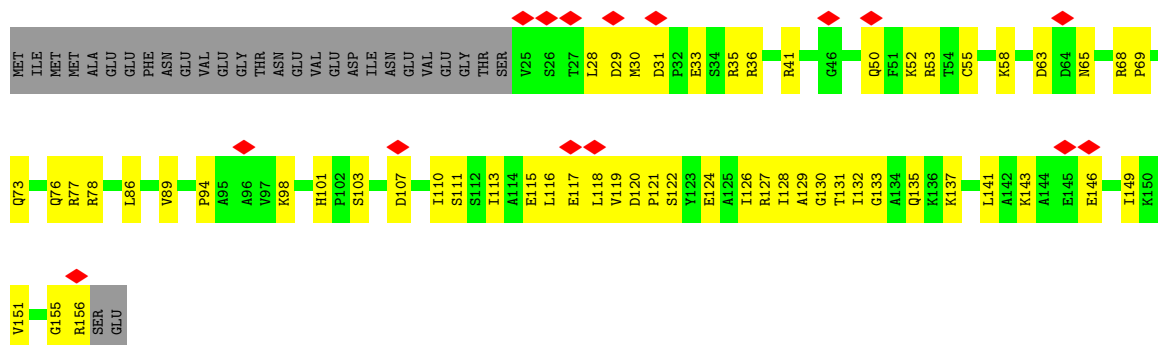




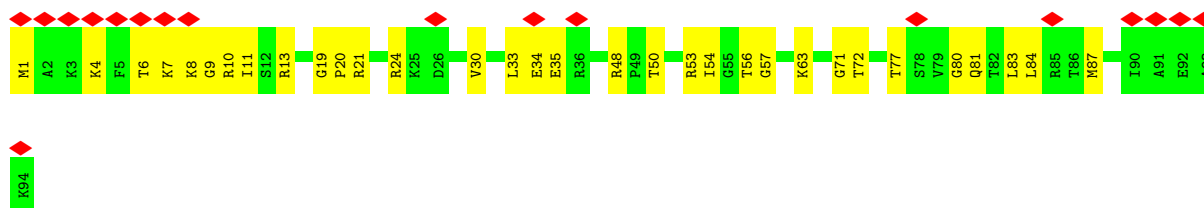
- Molecule 51: Large ribosomal subunit protein eL31



- Molecule 52: Large ribosomal subunit protein eL32



- Molecule 53: Large ribosomal subunit protein eL43

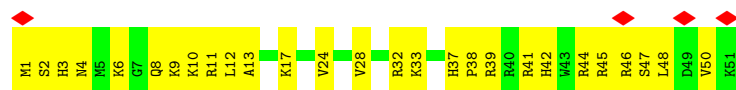


- Molecule 54: Large ribosomal subunit protein eL37



- Molecule 55: Large ribosomal subunit protein eL39

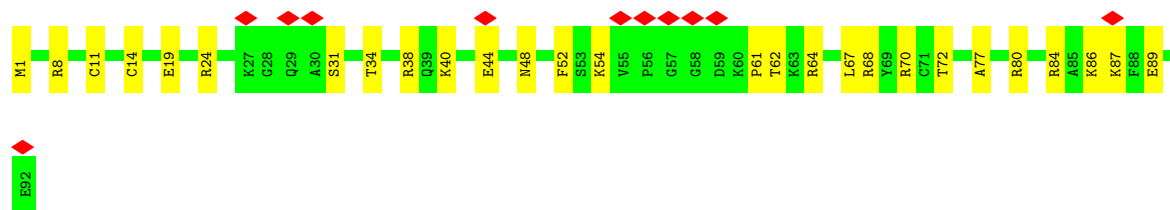




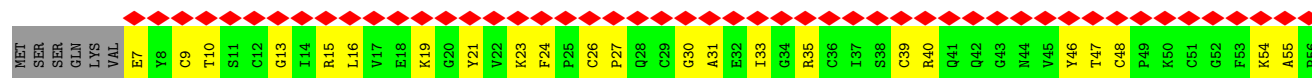
- Molecule 56: Large ribosomal subunit protein eL40



- Molecule 57: Large ribosomal subunit protein eL42



- Molecule 58: Putative zinc finger domain-containing protein



4 Experimental information

| Property | Value | Source |
|--------------------------------------|---|-----------|
| EM reconstruction method | SINGLE PARTICLE | Depositor |
| Imposed symmetry | POINT, Not provided | |
| Number of particles used | 91385 | Depositor |
| Resolution determination method | FSC 0.143 CUT-OFF | Depositor |
| CTF correction method | PHASE FLIPPING AND AMPLITUDE CORRECTION | Depositor |
| Microscope | FEI TALOS ARCTICA | Depositor |
| Voltage (kV) | 200 | Depositor |
| Electron dose ($e^-/\text{\AA}^2$) | 50 | Depositor |
| Minimum defocus (nm) | 600 | Depositor |
| Maximum defocus (nm) | 2000 | Depositor |
| Magnification | Not provided | |
| Image detector | FEI FALCON IV (4k x 4k) | Depositor |
| Maximum map value | 0.256 | Depositor |
| Minimum map value | -0.151 | Depositor |
| Average map value | 0.004 | Depositor |
| Map value standard deviation | 0.012 | Depositor |
| Recommended contour level | 0.046 | Depositor |
| Map size (\AA) | 351.84, 351.84, 351.84 | wwPDB |
| Map dimensions | 480, 480, 480 | wwPDB |
| Map angles ($^\circ$) | 90.0, 90.0, 90.0 | wwPDB |
| Pixel spacing (\AA) | 0.733, 0.733, 0.733 | Depositor |

5 Model quality

5.1 Standard geometry

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

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 | AA | 0.15 | 0/35052 | 0.25 | 0/54669 |
| 2 | AB | 0.13 | 0/1521 | 0.32 | 0/2068 |
| 3 | AC | 0.14 | 0/1447 | 0.35 | 0/1945 |
| 4 | AD | 0.15 | 0/1472 | 0.36 | 0/1985 |
| 5 | AE | 0.13 | 0/1375 | 0.33 | 0/1845 |
| 6 | AF | 0.14 | 0/1863 | 0.34 | 0/2504 |
| 7 | AG | 0.13 | 0/1531 | 0.32 | 0/2078 |
| 8 | AI | 0.16 | 0/1499 | 0.39 | 0/2017 |
| 9 | AJ | 0.16 | 0/997 | 0.38 | 0/1343 |
| 10 | AK | 0.17 | 0/972 | 0.35 | 0/1300 |
| 11 | AL | 0.19 | 0/1004 | 0.49 | 0/1352 |
| 12 | AN | 0.15 | 0/925 | 0.33 | 0/1244 |
| 13 | AO | 0.16 | 0/1102 | 0.34 | 0/1476 |
| 14 | AP | 0.14 | 0/1047 | 0.36 | 0/1407 |
| 15 | AR | 0.12 | 0/372 | 0.34 | 0/488 |
| 16 | AQ | 0.16 | 0/1247 | 0.31 | 0/1678 |
| 17 | AS | 0.16 | 0/864 | 0.35 | 0/1165 |
| 18 | AU | 0.12 | 0/790 | 0.33 | 0/1053 |
| 19 | AV | 0.13 | 0/1186 | 0.35 | 0/1596 |
| 20 | AW | 0.13 | 0/775 | 0.32 | 0/1038 |
| 21 | AX | 0.14 | 0/473 | 0.40 | 0/635 |
| 22 | AY | 0.13 | 0/473 | 0.32 | 0/632 |
| 23 | AH | 0.13 | 0/943 | 0.29 | 0/1266 |
| 24 | AM | 0.11 | 0/808 | 0.28 | 0/1090 |
| 25 | AT | 0.12 | 0/482 | 0.27 | 0/645 |
| 26 | BA | 0.29 | 0/66494 | 0.42 | 0/103680 |
| 27 | BB | 0.19 | 0/3037 | 0.29 | 0/4728 |
| 28 | BC | 0.26 | 0/1850 | 0.33 | 0/2497 |
| 29 | BD | 0.24 | 0/2646 | 0.33 | 0/3569 |
| 30 | BE | 0.22 | 0/1964 | 0.34 | 0/2654 |
| 31 | BF | 0.14 | 0/1310 | 0.30 | 0/1762 |
| 32 | BG | 0.16 | 0/1392 | 0.28 | 0/1870 |

| Mol | Chain | Bond lengths | | Bond angles | |
|-----|-------|--------------|----------|-------------|----------|
| | | RMSZ | # Z >5 | RMSZ | # Z >5 |
| 33 | BH | 0.15 | 0/864 | 0.30 | 0/1161 |
| 34 | BI | 0.21 | 0/1284 | 0.30 | 0/1719 |
| 35 | BJ | 0.23 | 0/1101 | 0.34 | 0/1474 |
| 36 | BK | 0.23 | 0/1010 | 0.33 | 0/1355 |
| 37 | BL | 0.22 | 0/1071 | 0.35 | 0/1425 |
| 38 | BM | 0.26 | 0/1625 | 0.34 | 0/2176 |
| 39 | BN | 0.18 | 0/1382 | 0.31 | 0/1863 |
| 40 | BO | 0.19 | 0/975 | 0.33 | 0/1312 |
| 41 | BP | 0.22 | 0/1209 | 0.32 | 0/1602 |
| 42 | BQ | 0.18 | 0/464 | 0.34 | 0/614 |
| 43 | BR | 0.23 | 0/780 | 0.28 | 0/1042 |
| 44 | BS | 0.21 | 0/1189 | 0.35 | 0/1588 |
| 45 | BT | 0.18 | 0/664 | 0.32 | 0/884 |
| 46 | BU | 0.18 | 0/919 | 0.36 | 0/1227 |
| 47 | BV | 0.20 | 0/508 | 0.26 | 0/670 |
| 48 | BW | 0.17 | 0/534 | 0.31 | 0/716 |
| 49 | BX | 0.22 | 0/1259 | 0.30 | 0/1692 |
| 50 | BY | 0.15 | 0/663 | 0.28 | 0/897 |
| 51 | BZ | 0.22 | 0/715 | 0.30 | 0/960 |
| 52 | Ba | 0.23 | 0/1044 | 0.33 | 0/1397 |
| 53 | Bb | 0.21 | 0/749 | 0.35 | 0/997 |
| 54 | Bc | 0.31 | 0/452 | 0.38 | 0/593 |
| 55 | Bd | 0.25 | 0/448 | 0.37 | 0/595 |
| 56 | Be | 0.18 | 0/355 | 0.27 | 0/468 |
| 57 | Bf | 0.23 | 0/777 | 0.34 | 0/1029 |
| 58 | AZ | 0.11 | 0/387 | 0.29 | 0/518 |
| All | All | 0.23 | 0/161341 | 0.35 | 0/239253 |

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 |
|-----|-------|---------------------|---------------------|
| 9 | AJ | 0 | 1 |

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

All (1) planarity outliers are listed below:

| Mol | Chain | Res | Type | Group |
|-----|-------|-----|------|---------|
| 9 | AJ | 28 | ARG | 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 | AA | 31324 | 0 | 15792 | 1066 | 0 |
| 2 | AB | 1492 | 0 | 1500 | 109 | 0 |
| 3 | AC | 1425 | 0 | 1460 | 119 | 0 |
| 4 | AD | 1454 | 0 | 1504 | 114 | 0 |
| 5 | AE | 1351 | 0 | 1397 | 103 | 0 |
| 6 | AF | 1837 | 0 | 1912 | 170 | 0 |
| 7 | AG | 1506 | 0 | 1561 | 160 | 0 |
| 8 | AI | 1479 | 0 | 1548 | 126 | 0 |
| 9 | AJ | 981 | 0 | 1008 | 84 | 0 |
| 10 | AK | 964 | 0 | 1028 | 85 | 0 |
| 11 | AL | 993 | 0 | 1033 | 80 | 0 |
| 12 | AN | 911 | 0 | 937 | 61 | 0 |
| 13 | AO | 1088 | 0 | 1142 | 78 | 0 |
| 14 | AP | 1033 | 0 | 1089 | 107 | 0 |
| 15 | AR | 366 | 0 | 362 | 44 | 0 |
| 16 | AQ | 1226 | 0 | 1305 | 101 | 0 |
| 17 | AS | 849 | 0 | 877 | 60 | 0 |
| 18 | AU | 775 | 0 | 787 | 85 | 0 |
| 19 | AV | 1163 | 0 | 1197 | 85 | 0 |
| 20 | AW | 766 | 0 | 795 | 86 | 0 |
| 21 | AX | 468 | 0 | 472 | 56 | 0 |
| 22 | AY | 471 | 0 | 490 | 53 | 0 |
| 23 | AH | 930 | 0 | 983 | 69 | 0 |
| 24 | AM | 796 | 0 | 841 | 91 | 0 |
| 25 | AT | 477 | 0 | 501 | 44 | 0 |
| 26 | BA | 59413 | 29860 | 29955 | 448 | 0 |
| 27 | BB | 2720 | 0 | 1382 | 62 | 0 |
| 28 | BC | 1808 | 0 | 1842 | 73 | 0 |
| 29 | BD | 2597 | 0 | 2692 | 121 | 0 |
| 30 | BE | 1930 | 0 | 1989 | 117 | 0 |
| 31 | BF | 1289 | 0 | 1317 | 79 | 0 |
| 32 | BG | 1371 | 0 | 1426 | 57 | 0 |
| 33 | BH | 857 | 0 | 898 | 46 | 0 |

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| Mol | Chain | Non-H | H(model) | H(added) | Clashes | Symm-Clashes |
|-----|-------|--------|----------|----------|---------|--------------|
| 34 | BI | 1261 | 0 | 1300 | 45 | 0 |
| 35 | BJ | 1086 | 0 | 1135 | 47 | 0 |
| 36 | BK | 999 | 0 | 1059 | 55 | 0 |
| 37 | BL | 1058 | 0 | 1044 | 41 | 0 |
| 38 | BM | 1593 | 0 | 1637 | 56 | 0 |
| 39 | BN | 1356 | 0 | 1358 | 75 | 0 |
| 40 | BO | 962 | 0 | 1021 | 52 | 0 |
| 41 | BP | 1195 | 0 | 1277 | 56 | 0 |
| 42 | BQ | 457 | 0 | 463 | 23 | 0 |
| 43 | BR | 766 | 0 | 777 | 25 | 0 |
| 44 | BS | 1169 | 0 | 1211 | 56 | 0 |
| 45 | BT | 656 | 0 | 688 | 27 | 0 |
| 46 | BU | 910 | 0 | 972 | 58 | 0 |
| 47 | BV | 499 | 0 | 515 | 34 | 0 |
| 48 | BW | 532 | 0 | 547 | 31 | 0 |
| 49 | BX | 1237 | 0 | 1269 | 49 | 0 |
| 50 | BY | 658 | 0 | 696 | 41 | 0 |
| 51 | BZ | 703 | 0 | 735 | 19 | 0 |
| 52 | Ba | 1028 | 0 | 1079 | 54 | 0 |
| 53 | Bb | 736 | 0 | 771 | 34 | 0 |
| 54 | Bc | 445 | 0 | 455 | 21 | 0 |
| 55 | Bd | 439 | 0 | 479 | 26 | 0 |
| 56 | Be | 353 | 0 | 374 | 13 | 0 |
| 57 | Bf | 760 | 0 | 797 | 24 | 0 |
| 58 | AZ | 379 | 0 | 374 | 32 | 0 |
| 59 | AA | 6 | 0 | 0 | 0 | 0 |
| 59 | AN | 1 | 0 | 0 | 0 | 0 |
| 59 | AS | 1 | 0 | 0 | 0 | 0 |
| 59 | BA | 42 | 0 | 0 | 0 | 0 |
| 59 | Bc | 1 | 0 | 0 | 0 | 0 |
| 60 | AG | 1 | 0 | 0 | 0 | 0 |
| 60 | AR | 1 | 0 | 0 | 0 | 0 |
| 60 | AS | 1 | 0 | 0 | 0 | 0 |
| 60 | AX | 1 | 0 | 0 | 0 | 0 |
| 60 | BV | 1 | 0 | 0 | 0 | 0 |
| 60 | Bb | 1 | 0 | 0 | 0 | 0 |
| 60 | Bc | 1 | 0 | 0 | 0 | 0 |
| 60 | Be | 1 | 0 | 0 | 0 | 0 |
| 60 | Bf | 1 | 0 | 0 | 0 | 0 |
| All | All | 149407 | 29860 | 105055 | 4401 | 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 18.

All (4401) 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) |
|--------------------|--------------------|--------------------------|-------------------|
| 7:AG:120:CYS:HB2 | 7:AG:147:PRO:HA | 1.45 | 0.97 |
| 21:AX:20:CYS:HB3 | 21:AX:39:CYS:SG | 2.05 | 0.97 |
| 9:AJ:27:VAL:HB | 9:AJ:61:TYR:HB2 | 1.47 | 0.96 |
| 1:AA:1343:A:H4' | 1:AA:1344:C:H5'' | 1.49 | 0.94 |
| 29:BD:25:PRO:HG2 | 29:BD:310:ARG:HG3 | 1.51 | 0.93 |
| 29:BD:138:VAL:HG12 | 29:BD:172:LEU:HD21 | 1.47 | 0.92 |
| 24:AM:15:ASP:HB2 | 24:AM:91:PRO:HG2 | 1.50 | 0.92 |
| 24:AM:19:VAL:HG21 | 24:AM:90:VAL:HG13 | 1.50 | 0.92 |
| 3:AC:126:LEU:HG | 3:AC:185:PRO:HA | 1.51 | 0.92 |
| 13:AO:103:ILE:HD12 | 13:AO:120:LYS:HD3 | 1.53 | 0.91 |
| 16:AQ:144:GLU:HG2 | 16:AQ:145:THR:HG23 | 1.51 | 0.91 |
| 2:AB:168:ILE:HB | 2:AB:190:VAL:HG22 | 1.53 | 0.91 |
| 6:AF:155:LEU:HD12 | 6:AF:160:LYS:HB3 | 1.54 | 0.90 |
| 7:AG:50:ASP:HB3 | 7:AG:76:VAL:HG22 | 1.52 | 0.90 |
| 25:AT:9:ILE:HG23 | 25:AT:50:VAL:HG22 | 1.52 | 0.90 |
| 6:AF:46:ILE:HG21 | 6:AF:106:LEU:HD12 | 1.54 | 0.90 |
| 1:AA:1004:C:H4' | 1:AA:1005:C:H5' | 1.53 | 0.90 |
| 1:AA:1105:C:H4' | 25:AT:49:ARG:HG2 | 1.51 | 0.90 |
| 49:BX:72:LEU:HD21 | 49:BX:78:LEU:HD13 | 1.51 | 0.89 |
| 4:AD:109:VAL:HG12 | 4:AD:146:LYS:HD3 | 1.54 | 0.89 |
| 1:AA:696:G:H5'' | 16:AQ:4:MET:HE1 | 1.52 | 0.89 |
| 17:AS:18:ASP:HB3 | 17:AS:21:CYS:HB2 | 1.52 | 0.89 |
| 1:AA:389:C:H5'' | 5:AE:127:ILE:HD12 | 1.53 | 0.89 |
| 19:AV:105:LEU:HB3 | 19:AV:111:LEU:HD13 | 1.51 | 0.89 |
| 41:BP:24:LEU:HD13 | 41:BP:32:ILE:HG21 | 1.54 | 0.89 |
| 37:BL:79:LEU:HD11 | 37:BL:99:LEU:HD21 | 1.53 | 0.89 |
| 7:AG:53:LEU:HD11 | 7:AG:75:ILE:HB | 1.54 | 0.88 |
| 46:BU:3:ALA:HB3 | 46:BU:5:VAL:HG22 | 1.54 | 0.88 |
| 11:AL:105:ARG:HD3 | 11:AL:108:LEU:HB2 | 1.54 | 0.88 |
| 4:AD:35:THR:HG21 | 12:AN:26:THR:HG23 | 1.55 | 0.88 |
| 2:AB:202:ASN:HA | 58:AZ:40:ARG:HG2 | 1.55 | 0.88 |
| 14:AP:38:GLN:HB3 | 14:AP:63:PRO:HB3 | 1.56 | 0.88 |
| 14:AP:101:ALA:HB2 | 18:AU:18:TYR:HB3 | 1.56 | 0.88 |
| 35:BJ:9:ALA:HA | 35:BJ:12:LEU:HD12 | 1.53 | 0.88 |
| 20:AW:43:LEU:HD11 | 20:AW:70:VAL:HG21 | 1.54 | 0.88 |
| 24:AM:43:LYS:HB2 | 24:AM:68:VAL:HB | 1.56 | 0.87 |
| 52:Ba:69:PRO:HB2 | 52:Ba:77:ARG:HB2 | 1.53 | 0.87 |
| 1:AA:85:C:HO2' | 6:AF:2:THR:N | 1.72 | 0.87 |
| 19:AV:137:LEU:HA | 19:AV:140:LEU:HG | 1.54 | 0.87 |
| 1:AA:242:G:H5' | 10:AK:31:SER:HA | 1.55 | 0.87 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 48:BW:6:THR:HB | 48:BW:56:ARG:HD3 | 1.57 | 0.86 |
| 1:AA:720:C:H5'' | 12:AN:121:LYS:HB2 | 1.58 | 0.86 |
| 12:AN:7:ALA:HB3 | 12:AN:71:ILE:HG22 | 1.58 | 0.85 |
| 9:AJ:31:SER:HB3 | 9:AJ:34:ILE:HG13 | 1.59 | 0.85 |
| 11:AL:17:ARG:HB3 | 11:AL:66:ASP:HB3 | 1.59 | 0.85 |
| 20:AW:14:LEU:HD11 | 20:AW:80:VAL:HG11 | 1.56 | 0.85 |
| 18:AU:50:PHE:HB2 | 18:AU:55:LYS:HD3 | 1.59 | 0.85 |
| 42:BQ:11:GLY:HA2 | 42:BQ:56:GLU:HG3 | 1.57 | 0.85 |
| 1:AA:1141:G:H1' | 7:AG:64:SER:HB2 | 1.59 | 0.84 |
| 1:AA:231:U:H4' | 10:AK:13:GLY:HA3 | 1.57 | 0.84 |
| 12:AN:73:ILE:HB | 12:AN:107:ILE:HG23 | 1.57 | 0.84 |
| 8:AI:110:VAL:HG22 | 8:AI:121:PRO:HB2 | 1.59 | 0.84 |
| 51:BZ:21:VAL:HG21 | 51:BZ:29:ARG:HG2 | 1.58 | 0.84 |
| 50:BY:40:VAL:HG13 | 50:BY:59:ILE:HG21 | 1.60 | 0.84 |
| 6:AF:177:VAL:HG12 | 6:AF:216:TYR:HA | 1.58 | 0.84 |
| 50:BY:28:THR:HG23 | 50:BY:85:ALA:HB2 | 1.59 | 0.84 |
| 12:AN:51:THR:HG22 | 12:AN:55:MET:HE2 | 1.59 | 0.84 |
| 15:AR:33:CYS:SG | 15:AR:36:CYS:HB2 | 2.18 | 0.84 |
| 37:BL:16:THR:HG22 | 37:BL:18:LYS:H | 1.43 | 0.84 |
| 10:AK:42:LYS:HE3 | 10:AK:56:LEU:HD11 | 1.61 | 0.83 |
| 14:AP:44:LEU:HD21 | 14:AP:109:LEU:HD21 | 1.60 | 0.83 |
| 31:BF:29:LEU:HD23 | 31:BF:37:VAL:HG21 | 1.60 | 0.83 |
| 6:AF:175:MET:HB2 | 6:AF:220:ILE:HD13 | 1.61 | 0.83 |
| 14:AP:23:LEU:HD12 | 14:AP:33:GLY:H | 1.42 | 0.83 |
| 14:AP:100:LEU:HD13 | 18:AU:19:ARG:HG2 | 1.59 | 0.83 |
| 12:AN:61:ASP:HA | 12:AN:64:LYS:HE2 | 1.62 | 0.82 |
| 8:AI:5:TYR:HB2 | 8:AI:31:PRO:HB2 | 1.62 | 0.82 |
| 26:BA:75:G:H21 | 26:BA:96:A:H62 | 1.27 | 0.82 |
| 7:AG:17:VAL:HG21 | 7:AG:40:ILE:HG23 | 1.60 | 0.82 |
| 39:BN:28:LEU:HD22 | 39:BN:51:PRO:HG3 | 1.61 | 0.82 |
| 4:AD:158:MET:HE2 | 4:AD:162:ILE:HD11 | 1.62 | 0.82 |
| 18:AU:72:ARG:HG2 | 18:AU:87:ILE:HD11 | 1.61 | 0.82 |
| 39:BN:37:VAL:HG22 | 39:BN:46:ILE:HG12 | 1.60 | 0.82 |
| 24:AM:34:ILE:HG13 | 24:AM:75:ILE:HG22 | 1.60 | 0.82 |
| 1:AA:1094:C:H1' | 11:AL:4:VAL:HG23 | 1.62 | 0.81 |
| 11:AL:105:ARG:HB2 | 11:AL:108:LEU:HD23 | 1.63 | 0.81 |
| 12:AN:73:ILE:HG21 | 12:AN:94:ILE:HD11 | 1.63 | 0.81 |
| 34:BI:74:LEU:HD21 | 34:BI:146:ALA:HA | 1.60 | 0.81 |
| 39:BN:36:VAL:HG21 | 39:BN:49:ILE:HD12 | 1.61 | 0.81 |
| 37:BL:67:VAL:HG12 | 37:BL:101:ILE:HG21 | 1.62 | 0.81 |
| 12:AN:18:THR:HG23 | 12:AN:55:MET:HE1 | 1.62 | 0.81 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 4:AD:21:ILE:HG21 | 4:AD:32:VAL:HB | 1.61 | 0.81 |
| 6:AF:126:VAL:HG23 | 6:AF:132:GLN:HB2 | 1.62 | 0.81 |
| 1:AA:1092:U:H4' | 1:AA:1093:A:H5'' | 1.62 | 0.80 |
| 7:AG:112:ARG:HD3 | 58:AZ:15:ARG:HH22 | 1.44 | 0.80 |
| 33:BH:52:ALA:HB3 | 33:BH:55:ILE:HD11 | 1.63 | 0.80 |
| 10:AK:37:SER:HB2 | 10:AK:57:GLN:HB3 | 1.63 | 0.80 |
| 46:BU:70:VAL:HG22 | 46:BU:77:ILE:HG22 | 1.63 | 0.80 |
| 9:AJ:37:VAL:HG22 | 9:AJ:103:ILE:HG21 | 1.63 | 0.80 |
| 3:AC:81:ASP:HB2 | 3:AC:83:GLN:HE22 | 1.45 | 0.80 |
| 30:BE:22:VAL:HG23 | 30:BE:115:ARG:HG2 | 1.62 | 0.80 |
| 1:AA:373:A:H5' | 6:AF:20:ASN:HD22 | 1.47 | 0.80 |
| 1:AA:458:C:H1' | 13:AO:62:SER:HB3 | 1.62 | 0.80 |
| 5:AE:100:ILE:HG23 | 5:AE:103:LEU:HD12 | 1.62 | 0.80 |
| 52:Ba:121:PRO:HB3 | 52:Ba:149:ILE:HA | 1.63 | 0.80 |
| 1:AA:853:A:H5'' | 13:AO:32:LEU:HD21 | 1.63 | 0.80 |
| 23:AH:49:LYS:HB2 | 23:AH:114:GLU:HB3 | 1.63 | 0.80 |
| 36:BK:8:ILE:HG21 | 36:BK:49:ILE:HB | 1.64 | 0.80 |
| 11:AL:20:VAL:HG13 | 11:AL:63:ILE:HG13 | 1.63 | 0.80 |
| 24:AM:6:ILE:HD11 | 24:AM:75:ILE:HG12 | 1.63 | 0.80 |
| 3:AC:117:LEU:HD22 | 3:AC:146:PHE:HB3 | 1.63 | 0.79 |
| 8:AI:21:LEU:HD22 | 22:AY:60:ILE:HG21 | 1.64 | 0.79 |
| 10:AK:32:ALA:HB3 | 10:AK:54:ARG:HD2 | 1.63 | 0.79 |
| 23:AH:63:MET:HA | 23:AH:97:ARG:HB2 | 1.64 | 0.79 |
| 1:AA:514:A:HO2' | 1:AA:827:C:HO2' | 1.28 | 0.79 |
| 5:AE:110:GLU:HA | 5:AE:115:THR:HG21 | 1.65 | 0.79 |
| 6:AF:29:GLY:HA3 | 6:AF:78:PRO:HG2 | 1.61 | 0.79 |
| 1:AA:1321:A:H4' | 8:AI:59:ASN:HB2 | 1.64 | 0.79 |
| 4:AD:43:LEU:HA | 4:AD:46:ARG:HD3 | 1.63 | 0.79 |
| 39:BN:103:LEU:HB2 | 39:BN:130:ILE:HD11 | 1.63 | 0.79 |
| 16:AQ:102:LEU:HD13 | 16:AQ:112:LYS:HD2 | 1.63 | 0.79 |
| 16:AQ:57:ARG:HA | 21:AX:34:ILE:HG23 | 1.64 | 0.79 |
| 26:BA:2353:C:HO2' | 26:BA:2386:A:HO2' | 1.25 | 0.79 |
| 44:BS:23:LEU:HD13 | 44:BS:87:GLU:HB3 | 1.65 | 0.79 |
| 40:BO:86:ILE:HD13 | 40:BO:99:ILE:HD13 | 1.62 | 0.79 |
| 52:Ba:121:PRO:HG3 | 52:Ba:149:ILE:HG12 | 1.64 | 0.79 |
| 19:AV:4:VAL:HG11 | 19:AV:128:MET:HE2 | 1.63 | 0.79 |
| 24:AM:7:ARG:HB2 | 24:AM:97:GLU:HB3 | 1.64 | 0.79 |
| 8:AI:87:LYS:HA | 8:AI:90:LYS:HE3 | 1.63 | 0.78 |
| 30:BE:147:GLU:HB3 | 30:BE:203:THR:HG21 | 1.65 | 0.78 |
| 6:AF:69:GLY:HA3 | 6:AF:158:PRO:HB3 | 1.64 | 0.78 |
| 50:BY:14:LYS:HE3 | 50:BY:92:SER:HA | 1.66 | 0.78 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 16:AQ:96:LEU:HD11 | 16:AQ:147:GLU:HG2 | 1.65 | 0.78 |
| 1:AA:464:G:H4' | 13:AO:105:GLY:HA2 | 1.65 | 0.78 |
| 1:AA:625:G:H21 | 12:AN:30:THR:HG21 | 1.47 | 0.78 |
| 6:AF:43:ILE:HD11 | 6:AF:49:LEU:HD12 | 1.64 | 0.78 |
| 30:BE:195:VAL:HB | 30:BE:215:VAL:HG22 | 1.66 | 0.78 |
| 7:AG:117:SER:HB2 | 7:AG:120:CYS:HB3 | 1.64 | 0.78 |
| 1:AA:161:C:H42 | 1:AA:183:G:H1 | 1.33 | 0.77 |
| 7:AG:92:GLN:HB3 | 7:AG:95:PRO:HD2 | 1.64 | 0.77 |
| 29:BD:85:ILE:HD12 | 29:BD:179:SER:HB2 | 1.66 | 0.77 |
| 32:BG:90:ILE:HD13 | 32:BG:99:VAL:HG21 | 1.67 | 0.77 |
| 8:AI:112:ARG:HB3 | 22:AY:26:MET:HE1 | 1.67 | 0.77 |
| 14:AP:49:ARG:HE | 19:AV:43:LEU:HD21 | 1.48 | 0.77 |
| 5:AE:113:LEU:HD13 | 5:AE:152:VAL:HG11 | 1.66 | 0.77 |
| 6:AF:90:VAL:HB | 6:AF:92:GLU:HG3 | 1.66 | 0.77 |
| 16:AQ:23:PRO:HG3 | 16:AQ:61:PRO:HG2 | 1.67 | 0.77 |
| 5:AE:117:VAL:HG13 | 5:AE:129:ALA:HB1 | 1.65 | 0.77 |
| 3:AC:157:PRO:HA | 3:AC:161:VAL:HB | 1.67 | 0.77 |
| 9:AJ:111:MET:HE1 | 9:AJ:119:LYS:HG3 | 1.66 | 0.77 |
| 11:AL:17:ARG:HB2 | 11:AL:68:ARG:HH12 | 1.49 | 0.76 |
| 1:AA:115:U:H5'' | 6:AF:139:THR:HG22 | 1.68 | 0.76 |
| 36:BK:4:MET:HE1 | 36:BK:50:GLY:HA3 | 1.66 | 0.76 |
| 39:BN:64:SER:HB3 | 39:BN:76:THR:HB | 1.65 | 0.76 |
| 24:AM:40:LEU:HD11 | 24:AM:72:LEU:HB2 | 1.67 | 0.76 |
| 5:AE:140:ILE:HB | 5:AE:160:ILE:HD13 | 1.68 | 0.76 |
| 9:AJ:111:MET:HE3 | 9:AJ:116:ALA:HA | 1.66 | 0.76 |
| 12:AN:19:ILE:HG12 | 12:AN:35:SER:HB2 | 1.68 | 0.76 |
| 21:AX:33:LYS:HD3 | 21:AX:45:GLU:HA | 1.66 | 0.76 |
| 39:BN:43:ASN:HB2 | 39:BN:64:SER:HB2 | 1.66 | 0.76 |
| 36:BK:132:VAL:HG11 | 47:BV:22:VAL:HG21 | 1.67 | 0.76 |
| 31:BF:131:ILE:HG23 | 31:BF:134:ARG:HH21 | 1.50 | 0.75 |
| 6:AF:92:GLU:HA | 6:AF:108:LYS:HE3 | 1.67 | 0.75 |
| 28:BC:105:PHE:HB2 | 53:Bb:83:LEU:HD12 | 1.68 | 0.75 |
| 15:AR:33:CYS:SG | 15:AR:36:CYS:CB | 2.74 | 0.75 |
| 20:AW:23:VAL:HB | 20:AW:64:SER:HB2 | 1.68 | 0.75 |
| 1:AA:1063:G:H22 | 1:AA:1097:U:H1' | 1.51 | 0.75 |
| 45:BT:4:ILE:HD13 | 45:BT:26:VAL:HG12 | 1.69 | 0.75 |
| 31:BF:8:LYS:HG2 | 31:BF:123:VAL:HG22 | 1.69 | 0.75 |
| 16:AQ:96:LEU:HD21 | 16:AQ:147:GLU:HG3 | 1.68 | 0.75 |
| 23:AH:118:LYS:HB2 | 23:AH:123:LEU:HD11 | 1.68 | 0.75 |
| 2:AB:59:LEU:HG | 2:AB:106:ARG:HH21 | 1.52 | 0.74 |
| 20:AW:35:VAL:HG21 | 20:AW:55:ILE:HG21 | 1.68 | 0.74 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 47:BV:46:PRO:HB2 | 47:BV:54:LYS:HD2 | 1.68 | 0.74 |
| 6:AF:191:ILE:HG12 | 6:AF:202:VAL:HG23 | 1.68 | 0.74 |
| 21:AX:35:THR:HA | 21:AX:42:THR:HA | 1.68 | 0.74 |
| 33:BH:23:LEU:HD23 | 33:BH:105:ALA:HB2 | 1.69 | 0.74 |
| 10:AK:110:THR:HG21 | 10:AK:122:VAL:HG23 | 1.69 | 0.74 |
| 18:AU:44:ARG:HA | 18:AU:47:LYS:HE2 | 1.70 | 0.74 |
| 23:AH:45:LEU:HD21 | 23:AH:124:LEU:HG | 1.69 | 0.74 |
| 6:AF:93:ALA:HB3 | 6:AF:109:LEU:HB2 | 1.69 | 0.74 |
| 12:AN:52:ALA:HA | 12:AN:55:MET:HE3 | 1.69 | 0.74 |
| 6:AF:42:ILE:HD13 | 6:AF:46:ILE:HD13 | 1.69 | 0.74 |
| 17:AS:16:CYS:HB3 | 17:AS:77:CYS:HB3 | 1.69 | 0.74 |
| 7:AG:194:GLN:HA | 7:AG:197:VAL:HG12 | 1.68 | 0.74 |
| 30:BE:140:ILE:HD13 | 30:BE:164:LEU:HD11 | 1.70 | 0.74 |
| 33:BH:41:ILE:HD13 | 33:BH:68:SER:HB2 | 1.69 | 0.74 |
| 44:BS:55:MET:HE2 | 44:BS:82:ILE:HD11 | 1.69 | 0.74 |
| 1:AA:938:G:H1' | 1:AA:983:A:H61 | 1.53 | 0.73 |
| 2:AB:72:ILE:HG23 | 2:AB:93:LEU:HD13 | 1.69 | 0.73 |
| 39:BN:148:ILE:HD12 | 39:BN:156:SER:HA | 1.68 | 0.73 |
| 19:AV:74:ARG:HH21 | 19:AV:92:LYS:HB3 | 1.54 | 0.73 |
| 28:BC:91:PRO:HD3 | 53:Bb:87:MET:HE1 | 1.68 | 0.73 |
| 30:BE:125:THR:O | 30:BE:129:LEU:HB2 | 1.88 | 0.73 |
| 6:AF:38:PRO:HG2 | 6:AF:41:ILE:HD12 | 1.70 | 0.73 |
| 52:Ba:128:ILE:HD11 | 52:Ba:141:LEU:HD21 | 1.70 | 0.73 |
| 11:AL:15:THR:HB | 11:AL:68:ARG:HB2 | 1.70 | 0.73 |
| 30:BE:32:ILE:HD13 | 30:BE:234:ALA:HB2 | 1.70 | 0.73 |
| 46:BU:47:VAL:HG22 | 46:BU:103:ILE:HD11 | 1.70 | 0.73 |
| 2:AB:104:ARG:HH21 | 2:AB:206:VAL:H | 1.37 | 0.73 |
| 6:AF:39:LEU:HD22 | 6:AF:60:LEU:HD21 | 1.69 | 0.73 |
| 10:AK:46:THR:HG21 | 10:AK:52:LYS:HE2 | 1.68 | 0.73 |
| 6:AF:124:THR:HG23 | 6:AF:132:GLN:HB3 | 1.71 | 0.73 |
| 41:BP:4:LEU:HD22 | 41:BP:32:ILE:HG22 | 1.69 | 0.73 |
| 20:AW:46:PRO:HD2 | 20:AW:49:LEU:HB2 | 1.69 | 0.73 |
| 30:BE:156:ILE:HD12 | 30:BE:165:TYR:HE1 | 1.51 | 0.73 |
| 26:BA:2374:G:O2' | 26:BA:2375:U:O4' | 2.07 | 0.72 |
| 43:BR:54:ASN:HB3 | 43:BR:56:LYS:HG2 | 1.70 | 0.72 |
| 2:AB:211:ASN:HA | 2:AB:217:LEU:HD21 | 1.72 | 0.72 |
| 20:AW:3:ILE:HD13 | 20:AW:23:VAL:HA | 1.69 | 0.72 |
| 6:AF:12:SER:HB3 | 6:AF:104:LEU:HD23 | 1.70 | 0.72 |
| 21:AX:33:LYS:HZ3 | 21:AX:46:PRO:HD2 | 1.54 | 0.72 |
| 24:AM:7:ARG:HG2 | 24:AM:72:LEU:HD13 | 1.71 | 0.72 |
| 46:BU:83:ILE:HG22 | 46:BU:93:PRO:HA | 1.70 | 0.72 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 11:AL:32:ILE:HG21 | 11:AL:38:THR:H | 1.54 | 0.72 |
| 20:AW:7:LYS:HB3 | 20:AW:20:ASP:HB2 | 1.71 | 0.72 |
| 20:AW:51:ILE:HG13 | 20:AW:89:ASN:HD22 | 1.53 | 0.72 |
| 9:AJ:53:ILE:HD13 | 21:AX:4:TYR:HB2 | 1.71 | 0.72 |
| 39:BN:103:LEU:HD22 | 39:BN:120:LEU:HD22 | 1.71 | 0.72 |
| 50:BY:28:THR:HG21 | 50:BY:83:ALA:HB1 | 1.69 | 0.72 |
| 2:AB:191:VAL:HG22 | 2:AB:205:LEU:HB2 | 1.71 | 0.72 |
| 6:AF:65:VAL:HG13 | 6:AF:87:LEU:HD21 | 1.72 | 0.72 |
| 12:AN:80:GLY:HA2 | 12:AN:114:PRO:HG3 | 1.70 | 0.72 |
| 22:AY:63:LEU:HB2 | 22:AY:66:THR:HG22 | 1.72 | 0.72 |
| 39:BN:141:GLU:HB2 | 39:BN:146:GLU:HG3 | 1.72 | 0.72 |
| 1:AA:1152:U:H4' | 3:AC:175:THR:HG21 | 1.72 | 0.72 |
| 2:AB:97:ASP:HB3 | 2:AB:100:SER:HB2 | 1.70 | 0.72 |
| 4:AD:37:ALA:HB1 | 4:AD:42:LEU:HD22 | 1.71 | 0.72 |
| 29:BD:113:ARG:HG3 | 29:BD:114:ILE:HG23 | 1.72 | 0.72 |
| 1:AA:464:G:H5'' | 13:AO:111:MET:HG3 | 1.69 | 0.72 |
| 1:AA:921:G:H4' | 1:AA:922:A:H5' | 1.72 | 0.71 |
| 9:AJ:46:TYR:HB3 | 9:AJ:69:ILE:HG22 | 1.71 | 0.71 |
| 23:AH:34:ILE:H | 23:AH:99:LYS:HB2 | 1.54 | 0.71 |
| 26:BA:75:G:N2 | 26:BA:96:A:H62 | 1.86 | 0.71 |
| 24:AM:51:LYS:HD2 | 24:AM:60:THR:HB | 1.72 | 0.71 |
| 50:BY:51:LYS:HA | 50:BY:54:ILE:HG12 | 1.71 | 0.71 |
| 10:AK:34:THR:HG21 | 10:AK:113:PRO:HB2 | 1.72 | 0.71 |
| 22:AY:38:LEU:HA | 22:AY:43:LYS:HG2 | 1.72 | 0.71 |
| 31:BF:76:LEU:HD22 | 31:BF:162:VAL:HG21 | 1.71 | 0.71 |
| 43:BR:71:SER:HB3 | 43:BR:88:LEU:HD23 | 1.70 | 0.71 |
| 1:AA:1269:C:H4' | 1:AA:1270:G:H5' | 1.73 | 0.71 |
| 44:BS:25:ILE:HG21 | 44:BS:87:GLU:HG3 | 1.71 | 0.71 |
| 23:AH:78:ALA:HB2 | 23:AH:91:ARG:HG2 | 1.72 | 0.71 |
| 1:AA:275:U:H3 | 1:AA:286:A:H61 | 1.37 | 0.71 |
| 1:AA:503:A:HO2' | 1:AA:506:G:HO2' | 1.37 | 0.70 |
| 6:AF:220:ILE:HD12 | 6:AF:229:LEU:HD11 | 1.73 | 0.70 |
| 1:AA:88:C:H1' | 1:AA:336:G:H5' | 1.73 | 0.70 |
| 16:AQ:17:PRO:HB3 | 21:AX:8:PRO:HD3 | 1.73 | 0.70 |
| 10:AK:76:ILE:HG12 | 10:AK:100:ILE:HD12 | 1.73 | 0.70 |
| 12:AN:35:SER:H | 12:AN:38:MET:HG3 | 1.55 | 0.70 |
| 19:AV:38:GLY:HA2 | 19:AV:94:SER:HB2 | 1.72 | 0.70 |
| 20:AW:32:ARG:HA | 20:AW:55:ILE:HD11 | 1.74 | 0.70 |
| 7:AG:182:LEU:HD11 | 7:AG:186:LYS:HE3 | 1.74 | 0.70 |
| 50:BY:77:LYS:HG3 | 50:BY:81:ILE:HD11 | 1.74 | 0.70 |
| 3:AC:33:MET:HE1 | 3:AC:44:ILE:HD12 | 1.73 | 0.70 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 9:AJ:18:GLU:HG3 | 9:AJ:69:ILE:HG13 | 1.73 | 0.70 |
| 26:BA:601:G:O2' | 26:BA:1308:G:OP1 | 2.09 | 0.70 |
| 20:AW:14:LEU:HD22 | 20:AW:16:ARG:HD3 | 1.73 | 0.70 |
| 34:BI:70:ALA:HB2 | 34:BI:153:ALA:HB2 | 1.73 | 0.70 |
| 35:BJ:86:TYR:HA | 35:BJ:91:GLY:HA3 | 1.74 | 0.70 |
| 23:AH:46:ALA:H | 23:AH:123:LEU:HD13 | 1.57 | 0.69 |
| 24:AM:90:VAL:HG11 | 24:AM:96:ILE:HD11 | 1.74 | 0.69 |
| 50:BY:89:VAL:HG12 | 50:BY:92:SER:HB3 | 1.74 | 0.69 |
| 4:AD:160:GLU:HA | 4:AD:165:ARG:HB2 | 1.74 | 0.69 |
| 27:BB:52:G:H21 | 27:BB:53:C:H41 | 1.40 | 0.69 |
| 6:AF:195:LYS:HE2 | 17:AS:5:ILE:HD13 | 1.74 | 0.69 |
| 26:BA:1578:U:OP2 | 41:BP:124:TYR:OH | 2.11 | 0.69 |
| 45:BT:14:MET:HA | 45:BT:17:LEU:HD13 | 1.74 | 0.69 |
| 20:AW:52:ILE:HA | 20:AW:68:ALA:HA | 1.74 | 0.69 |
| 1:AA:764:U:HO2' | 9:AJ:2:VAL:N | 1.90 | 0.69 |
| 17:AS:54:MET:HG2 | 17:AS:65:LYS:HG3 | 1.74 | 0.69 |
| 24:AM:43:LYS:HD3 | 24:AM:70:LYS:HE3 | 1.75 | 0.69 |
| 29:BD:271:ILE:HD13 | 29:BD:295:ILE:HG22 | 1.75 | 0.69 |
| 24:AM:50:ARG:HD2 | 24:AM:61:TRP:HE1 | 1.57 | 0.69 |
| 26:BA:1476:A:O2' | 26:BA:1487:U:O2 | 2.11 | 0.69 |
| 7:AG:143:LEU:HD13 | 7:AG:170:VAL:HG11 | 1.75 | 0.69 |
| 57:Bf:11:CYS:HB3 | 57:Bf:14:CYS:SG | 2.32 | 0.69 |
| 7:AG:38:PRO:HA | 7:AG:106:LEU:HD21 | 1.76 | 0.68 |
| 26:BA:1044:G:OP1 | 34:BI:96:ARG:NH2 | 2.26 | 0.68 |
| 33:BH:20:ALA:HB2 | 33:BH:108:MET:HE2 | 1.76 | 0.68 |
| 1:AA:526:C:H5'' | 13:AO:4:GLY:HA3 | 1.74 | 0.68 |
| 14:AP:55:ILE:HD13 | 14:AP:88:ILE:HD11 | 1.73 | 0.68 |
| 6:AF:119:ARG:HD2 | 6:AF:218:ILE:HD12 | 1.75 | 0.68 |
| 24:AM:32:VAL:HG13 | 24:AM:76:ALA:HB3 | 1.75 | 0.68 |
| 32:BG:17:VAL:HG12 | 32:BG:26:ALA:HA | 1.76 | 0.68 |
| 33:BH:9:VAL:HG12 | 33:BH:119:LEU:HD11 | 1.75 | 0.68 |
| 26:BA:2374:G:O2' | 26:BA:2375:U:O5' | 2.11 | 0.68 |
| 2:AB:105:ILE:HG12 | 2:AB:224:LEU:HD21 | 1.76 | 0.68 |
| 7:AG:75:ILE:HD12 | 7:AG:85:LEU:HA | 1.74 | 0.68 |
| 10:AK:54:ARG:HD3 | 10:AK:114:GLY:HA2 | 1.76 | 0.68 |
| 17:AS:39:VAL:HB | 17:AS:48:VAL:HG13 | 1.75 | 0.68 |
| 26:BA:453:A:N3 | 26:BA:457:A:O2' | 2.26 | 0.68 |
| 44:BS:41:LYS:HB2 | 44:BS:44:GLU:HG2 | 1.75 | 0.68 |
| 40:BO:110:GLU:HA | 40:BO:113:MET:HE2 | 1.76 | 0.68 |
| 1:AA:1050:A:H2' | 1:AA:1051:C:H5' | 1.74 | 0.68 |
| 7:AG:60:ARG:HB3 | 7:AG:67:ARG:HB2 | 1.76 | 0.68 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 9:AJ:29:PRO:HB3 | 9:AJ:58:ALA:HB3 | 1.74 | 0.68 |
| 10:AK:63:VAL:HG12 | 10:AK:123:LEU:HB2 | 1.76 | 0.68 |
| 17:AS:1:MET:HE2 | 17:AS:1:MET:HA | 1.76 | 0.68 |
| 46:BU:78:THR:HG21 | 46:BU:93:PRO:HB2 | 1.76 | 0.68 |
| 8:AI:112:ARG:HD2 | 8:AI:121:PRO:HB3 | 1.76 | 0.68 |
| 26:BA:2366:G:OP1 | 43:BR:9:ARG:NH2 | 2.27 | 0.67 |
| 30:BE:209:ALA:HA | 30:BE:212:LEU:HD23 | 1.76 | 0.67 |
| 1:AA:1438:A:H5'' | 1:AA:1446:G:H5'' | 1.76 | 0.67 |
| 26:BA:683:C:O2 | 26:BA:723:U:O2' | 2.11 | 0.67 |
| 26:BA:1734:C:O2 | 29:BD:228:ARG:NH2 | 2.24 | 0.67 |
| 3:AC:152:LYS:HE3 | 3:AC:157:PRO:HG2 | 1.75 | 0.67 |
| 12:AN:73:ILE:HG13 | 12:AN:94:ILE:HG12 | 1.76 | 0.67 |
| 26:BA:1204:A:O2' | 32:BG:1:MET:N | 2.28 | 0.67 |
| 26:BA:1240:G:OP1 | 42:BQ:49:LYS:NZ | 2.27 | 0.67 |
| 49:BX:83:ILE:O | 49:BX:87:THR:OG1 | 2.13 | 0.67 |
| 1:AA:197:C:H4' | 1:AA:198:G:H5'' | 1.76 | 0.67 |
| 1:AA:920:G:OP2 | 15:AR:20:ARG:NH2 | 2.28 | 0.67 |
| 46:BU:58:GLY:O | 46:BU:61:LYS:NZ | 2.28 | 0.67 |
| 1:AA:1095:U:O2 | 11:AL:17:ARG:NH2 | 2.27 | 0.67 |
| 2:AB:135:MET:HG3 | 2:AB:243:PHE:HA | 1.76 | 0.67 |
| 6:AF:23:ILE:HG22 | 6:AF:41:ILE:HD11 | 1.77 | 0.67 |
| 9:AJ:60:ILE:HD11 | 21:AX:4:TYR:HB3 | 1.77 | 0.67 |
| 16:AQ:150:ILE:HG12 | 50:BY:17:LYS:HE3 | 1.77 | 0.67 |
| 26:BA:124:G:N2 | 26:BA:126:A:O4' | 2.27 | 0.67 |
| 26:BA:1836:C:OP2 | 28:BC:147:LYS:NZ | 2.28 | 0.67 |
| 28:BC:130:VAL:HG12 | 28:BC:131:THR:HG23 | 1.76 | 0.67 |
| 1:AA:689:G:OP1 | 21:AX:10:SER:OG | 2.13 | 0.67 |
| 26:BA:2686:G:O2' | 36:BK:38:ARG:O | 2.12 | 0.67 |
| 44:BS:128:ARG:HG3 | 44:BS:134:THR:HG23 | 1.74 | 0.67 |
| 47:BV:4:ARG:HB2 | 47:BV:13:LEU:HB2 | 1.77 | 0.67 |
| 57:Bf:87:LYS:HE2 | 57:Bf:89:GLU:HG2 | 1.77 | 0.67 |
| 1:AA:903:U:O2' | 1:AA:905:A:N7 | 2.26 | 0.67 |
| 3:AC:158:VAL:HA | 3:AC:162:VAL:HG22 | 1.76 | 0.67 |
| 4:AD:148:ARG:O | 4:AD:152:SER:OG | 2.13 | 0.67 |
| 8:AI:186:LYS:HE2 | 8:AI:189:ARG:HD3 | 1.75 | 0.67 |
| 16:AQ:5:HIS:HB2 | 16:AQ:117:LEU:HD13 | 1.76 | 0.67 |
| 26:BA:109:G:OP2 | 26:BA:111:A:O2' | 2.11 | 0.67 |
| 26:BA:1483:A:O2' | 26:BA:1484:A:O4' | 2.13 | 0.67 |
| 22:AY:63:LEU:HD12 | 22:AY:66:THR:HG22 | 1.76 | 0.67 |
| 26:BA:2544:C:OP1 | 32:BG:160:ARG:NH2 | 2.28 | 0.67 |
| 48:BW:30:ARG:HE | 48:BW:34:LEU:HD11 | 1.59 | 0.67 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 49:BX:26:LEU:HD21 | 49:BX:33:VAL:HG22 | 1.77 | 0.67 |
| 1:AA:1259:G:OP1 | 18:AU:43:ARG:NH2 | 2.28 | 0.66 |
| 12:AN:8:VAL:HG11 | 12:AN:74:ARG:HD3 | 1.77 | 0.66 |
| 26:BA:1354:G:N7 | 44:BS:26:SER:OG | 2.28 | 0.66 |
| 52:Ba:126:ILE:HD11 | 52:Ba:151:VAL:HG22 | 1.77 | 0.66 |
| 1:AA:171:G:N2 | 1:AA:174:A:OP2 | 2.29 | 0.66 |
| 1:AA:220:G:OP1 | 17:AS:66:ARG:NH1 | 2.28 | 0.66 |
| 52:Ba:155:GLY:O | 52:Ba:156:ARG:NH1 | 2.28 | 0.66 |
| 1:AA:235:A:OP2 | 10:AK:8:ARG:NH1 | 2.28 | 0.66 |
| 8:AI:22:GLY:HA2 | 22:AY:60:ILE:HB | 1.77 | 0.66 |
| 11:AL:88:VAL:HG21 | 11:AL:97:ARG:HD2 | 1.76 | 0.66 |
| 26:BA:2813:A:O2' | 26:BA:2815:A:OP2 | 2.10 | 0.66 |
| 1:AA:1270:G:O6 | 18:AU:41:ARG:NH1 | 2.27 | 0.66 |
| 9:AJ:112:SER:OG | 9:AJ:115:GLU:OE1 | 2.13 | 0.66 |
| 31:BF:6:VAL:HA | 31:BF:124:LEU:HD13 | 1.77 | 0.66 |
| 34:BI:56:CYS:SG | 34:BI:159:THR:OG1 | 2.52 | 0.66 |
| 1:AA:85:C:O2' | 6:AF:2:THR:N | 2.29 | 0.66 |
| 7:AG:113:ARG:NH2 | 7:AG:165:ALA:O | 2.27 | 0.66 |
| 24:AM:39:PRO:HA | 24:AM:71:ARG:HD3 | 1.78 | 0.66 |
| 26:BA:425:A:OP1 | 38:BM:49:ARG:NH2 | 2.28 | 0.66 |
| 32:BG:11:ILE:HD11 | 32:BG:51:VAL:HG23 | 1.77 | 0.66 |
| 1:AA:1235:A:O2' | 19:AV:80:LYS:NZ | 2.28 | 0.66 |
| 7:AG:55:ILE:HD11 | 7:AG:93:VAL:HG23 | 1.76 | 0.66 |
| 26:BA:2789:G:N7 | 35:BJ:77:LYS:NZ | 2.44 | 0.66 |
| 1:AA:7:G:N7 | 7:AG:179:ARG:NH1 | 2.44 | 0.66 |
| 1:AA:953:G:N2 | 1:AA:972:C:O4' | 2.29 | 0.66 |
| 2:AB:123:VAL:HG11 | 2:AB:179:VAL:HG11 | 1.78 | 0.66 |
| 20:AW:46:PRO:HG2 | 20:AW:49:LEU:HD12 | 1.78 | 0.66 |
| 26:BA:76:U:O2' | 26:BA:94:G:N2 | 2.29 | 0.66 |
| 26:BA:186:A:OP1 | 38:BM:177:ARG:NH2 | 2.29 | 0.66 |
| 40:BO:119:GLY:HA2 | 40:BO:122:ILE:HD12 | 1.78 | 0.66 |
| 41:BP:119:VAL:HG21 | 41:BP:147:LEU:HD22 | 1.77 | 0.66 |
| 46:BU:31:ARG:NH1 | 46:BU:43:ARG:O | 2.29 | 0.66 |
| 1:AA:504:U:OP1 | 13:AO:15:ARG:NH2 | 2.28 | 0.66 |
| 26:BA:444:C:OP1 | 30:BE:187:ARG:NH2 | 2.29 | 0.66 |
| 26:BA:1831:G:H21 | 28:BC:228:LEU:HD23 | 1.61 | 0.66 |
| 29:BD:263:GLU:HG3 | 29:BD:302:PRO:HD3 | 1.78 | 0.66 |
| 32:BG:137:ILE:HG22 | 32:BG:145:VAL:HG23 | 1.78 | 0.66 |
| 1:AA:767:C:H4' | 9:AJ:13:THR:HG22 | 1.77 | 0.66 |
| 1:AA:994:A:O2' | 3:AC:139:ALA:O | 2.13 | 0.66 |
| 7:AG:48:LEU:HD22 | 7:AG:105:LYS:HD2 | 1.76 | 0.66 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 7:AG:84:GLY:HA3 | 7:AG:108:ILE:HD13 | 1.78 | 0.66 |
| 29:BD:17:ARG:HB2 | 29:BD:216:ARG:HH21 | 1.60 | 0.66 |
| 1:AA:884:A:O2' | 8:AI:130:ARG:NH1 | 2.29 | 0.66 |
| 1:AA:917:G:H4' | 1:AA:918:C:H5'' | 1.78 | 0.66 |
| 1:AA:1051:C:O2' | 3:AC:156:ASN:OD1 | 2.13 | 0.66 |
| 13:AO:58:LYS:H | 13:AO:113:ASP:HB2 | 1.61 | 0.66 |
| 26:BA:1430:U:OP2 | 26:BA:1485:U:O2' | 2.13 | 0.66 |
| 30:BE:205:ILE:HD12 | 30:BE:208:ALA:HB3 | 1.77 | 0.66 |
| 48:BW:9:ILE:HG22 | 48:BW:60:ILE:HG21 | 1.77 | 0.66 |
| 1:AA:135:G:N2 | 1:AA:138:A:OP2 | 2.29 | 0.65 |
| 1:AA:406:C:O2' | 5:AE:173:HIS:NE2 | 2.29 | 0.65 |
| 1:AA:1209:C:H42 | 1:AA:1220:G:H1 | 1.44 | 0.65 |
| 1:AA:1215:A:N3 | 1:AA:1273:C:O2' | 2.28 | 0.65 |
| 16:AQ:84:ILE:HD11 | 16:AQ:151:THR:HB | 1.77 | 0.65 |
| 18:AU:88:HIS:ND1 | 18:AU:92:GLY:O | 2.29 | 0.65 |
| 23:AH:70:PRO:HA | 23:AH:100:GLU:HB2 | 1.78 | 0.65 |
| 31:BF:49:PHE:HB3 | 31:BF:51:ILE:HG23 | 1.77 | 0.65 |
| 39:BN:38:ARG:HA | 39:BN:108:GLN:HE22 | 1.61 | 0.65 |
| 1:AA:233:G:N3 | 10:AK:8:ARG:NH2 | 2.45 | 0.65 |
| 1:AA:1036:A:H1' | 1:AA:1117:A:H1' | 1.77 | 0.65 |
| 26:BA:75:G:H21 | 26:BA:96:A:N6 | 1.95 | 0.65 |
| 26:BA:130:A:O2' | 26:BA:1448:A:OP1 | 2.14 | 0.65 |
| 26:BA:627:G:O2' | 26:BA:1342:A:OP1 | 2.14 | 0.65 |
| 29:BD:169:ALA:HB3 | 29:BD:175:LYS:HG3 | 1.78 | 0.65 |
| 1:AA:172:G:OP1 | 6:AF:196:SER:N | 2.29 | 0.65 |
| 1:AA:785:G:O2' | 1:AA:787:G:N7 | 2.30 | 0.65 |
| 1:AA:1445:A:N7 | 12:AN:125:ARG:NH1 | 2.43 | 0.65 |
| 6:AF:42:ILE:HA | 6:AF:46:ILE:HD13 | 1.78 | 0.65 |
| 26:BA:1059:C:O2' | 26:BA:1071:A:N3 | 2.29 | 0.65 |
| 26:BA:1456:U:O2 | 54:Bc:41:LYS:NZ | 2.29 | 0.65 |
| 29:BD:6:ARG:HD2 | 29:BD:7:PRO:HD2 | 1.76 | 0.65 |
| 29:BD:314:PRO:HB2 | 29:BD:317:ALA:HB2 | 1.79 | 0.65 |
| 57:Bf:62:THR:HG22 | 57:Bf:86:LYS:HD3 | 1.78 | 0.65 |
| 1:AA:316:U:O2' | 23:AH:93:ARG:NH2 | 2.29 | 0.65 |
| 1:AA:909:G:O2' | 1:AA:1146:U:OP1 | 2.13 | 0.65 |
| 1:AA:1235:A:OP1 | 19:AV:90:LYS:NZ | 2.29 | 0.65 |
| 4:AD:116:VAL:N | 4:AD:192:THR:OG1 | 2.29 | 0.65 |
| 10:AK:2:ARG:N | 10:AK:28:GLY:O | 2.28 | 0.65 |
| 23:AH:118:LYS:HD2 | 23:AH:123:LEU:HD21 | 1.79 | 0.65 |
| 26:BA:693:A:O2' | 37:BL:109:ARG:NH1 | 2.29 | 0.65 |
| 30:BE:206:LEU:HD23 | 30:BE:217:VAL:HG22 | 1.79 | 0.65 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 36:BK:23:ASN:ND2 | 36:BK:108:ASP:OD1 | 2.30 | 0.65 |
| 52:Ba:111:SER:N | 52:Ba:115:GLU:OE1 | 2.29 | 0.65 |
| 1:AA:630:G:H5'' | 12:AN:19:ILE:HD11 | 1.77 | 0.65 |
| 1:AA:1259:G:O5' | 18:AU:40:SER:OG | 2.13 | 0.65 |
| 1:AA:1265:A:O2' | 18:AU:73:ASP:OD2 | 2.14 | 0.65 |
| 4:AD:32:VAL:HG13 | 4:AD:50:THR:HG21 | 1.78 | 0.65 |
| 5:AE:118:LEU:HD22 | 5:AE:126:VAL:HB | 1.76 | 0.65 |
| 5:AE:173:HIS:HB3 | 5:AE:176:ARG:HB3 | 1.78 | 0.65 |
| 8:AI:100:ALA:HB2 | 8:AI:166:ALA:HB2 | 1.78 | 0.65 |
| 9:AJ:47:ILE:HD11 | 9:AJ:63:VAL:HG11 | 1.79 | 0.65 |
| 23:AH:40:GLY:O | 23:AH:44:GLY:N | 2.29 | 0.65 |
| 46:BU:55:VAL:HG22 | 46:BU:101:VAL:HG12 | 1.78 | 0.65 |
| 1:AA:1106:C:N4 | 1:AA:1120:C:OP2 | 2.29 | 0.65 |
| 8:AI:82:PHE:HD1 | 8:AI:93:PRO:HB2 | 1.62 | 0.65 |
| 8:AI:118:ILE:HG22 | 8:AI:120:VAL:HG23 | 1.78 | 0.65 |
| 26:BA:188:A:HO2' | 26:BA:421:C:HO2' | 1.45 | 0.65 |
| 26:BA:2591:U:OP1 | 29:BD:229:GLN:NE2 | 2.30 | 0.65 |
| 37:BL:96:LEU:HD12 | 37:BL:101:ILE:HB | 1.79 | 0.65 |
| 1:AA:710:G:H4' | 1:AA:1451:A:H4' | 1.79 | 0.65 |
| 26:BA:293:A:N7 | 26:BA:346:U:O2' | 2.29 | 0.65 |
| 31:BF:101:HIS:HB2 | 31:BF:107:MET:HE1 | 1.78 | 0.65 |
| 36:BK:99:ASP:OD1 | 36:BK:103:ILE:N | 2.30 | 0.65 |
| 1:AA:666:G:OP1 | 1:AA:798:G:N2 | 2.30 | 0.65 |
| 1:AA:1105:C:O2' | 25:AT:5:ARG:NH2 | 2.30 | 0.65 |
| 3:AC:48:SER:HB2 | 3:AC:51:PRO:HB3 | 1.78 | 0.65 |
| 9:AJ:101:ALA:HB1 | 9:AJ:112:SER:HB2 | 1.79 | 0.65 |
| 18:AU:63:GLU:OE1 | 18:AU:65:LYS:NZ | 2.30 | 0.65 |
| 21:AX:33:LYS:HA | 21:AX:46:PRO:HD3 | 1.79 | 0.65 |
| 22:AY:49:ARG:NH2 | 22:AY:68:ARG:O | 2.29 | 0.65 |
| 26:BA:149:G:O2' | 26:BA:182:A:N3 | 2.30 | 0.65 |
| 26:BA:530:A:N3 | 52:Ba:58:LYS:NZ | 2.44 | 0.65 |
| 26:BA:2583:U:OP2 | 29:BD:242:ASN:ND2 | 2.28 | 0.65 |
| 27:BB:51:A:OP1 | 39:BN:79:THR:N | 2.30 | 0.65 |
| 35:BJ:53:TYR:HA | 35:BJ:56:THR:HG22 | 1.78 | 0.65 |
| 1:AA:1035:C:O2' | 25:AT:10:LYS:NZ | 2.29 | 0.65 |
| 4:AD:72:ASN:OD1 | 4:AD:79:THR:OG1 | 2.15 | 0.65 |
| 7:AG:118:TRP:HA | 9:AJ:97:LYS:HE3 | 1.79 | 0.65 |
| 26:BA:340:U:O2' | 46:BU:4:MET:SD | 2.55 | 0.65 |
| 26:BA:2304:A:OP1 | 31:BF:135:ARG:NH1 | 2.30 | 0.65 |
| 1:AA:357:U:OP1 | 6:AF:53:SER:N | 2.30 | 0.65 |
| 8:AI:71:LYS:HZ3 | 8:AI:75:LEU:HD11 | 1.62 | 0.65 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 14:AP:24:VAL:HG11 | 14:AP:67:LEU:HD21 | 1.79 | 0.65 |
| 26:BA:149:G:OP2 | 38:BM:189:ARG:NH2 | 2.30 | 0.65 |
| 26:BA:560:A:O2' | 49:BX:77:ARG:NH2 | 2.30 | 0.65 |
| 26:BA:881:C:OP1 | 30:BE:60:ARG:NH2 | 2.28 | 0.65 |
| 26:BA:2104:G:N2 | 28:BC:219:ASN:O | 2.30 | 0.65 |
| 58:AZ:26:CYS:HB3 | 58:AZ:30:GLY:H | 1.62 | 0.65 |
| 1:AA:279:G:N2 | 1:AA:282:A:OP2 | 2.30 | 0.64 |
| 4:AD:50:THR:O | 4:AD:66:VAL:N | 2.30 | 0.64 |
| 4:AD:103:ILE:HG13 | 4:AD:134:MET:HG2 | 1.80 | 0.64 |
| 6:AF:144:SER:OG | 6:AF:146:GLU:OE1 | 2.14 | 0.64 |
| 19:AV:26:ILE:HD12 | 19:AV:104:GLN:HB3 | 1.79 | 0.64 |
| 26:BA:635:G:OP1 | 30:BE:91:ARG:NH1 | 2.29 | 0.64 |
| 27:BB:46:A:HO2' | 31:BF:133:LYS:HZ2 | 1.45 | 0.64 |
| 32:BG:118:ARG:NH1 | 32:BG:155:LYS:O | 2.29 | 0.64 |
| 1:AA:883:A:N6 | 1:AA:1292:U:O4 | 2.31 | 0.64 |
| 2:AB:172:ASP:OD1 | 2:AB:211:ASN:ND2 | 2.30 | 0.64 |
| 6:AF:100:GLU:OE2 | 6:AF:101:LYS:NZ | 2.30 | 0.64 |
| 14:AP:42:THR:HG22 | 14:AP:49:ARG:H | 1.62 | 0.64 |
| 26:BA:255:C:OP1 | 38:BM:43:ARG:NH2 | 2.29 | 0.64 |
| 1:AA:955:C:N4 | 1:AA:969:U:O2' | 2.29 | 0.64 |
| 1:AA:1236:A:O2' | 8:AI:70:LYS:NZ | 2.31 | 0.64 |
| 7:AG:13:LEU:HD13 | 7:AG:35:ILE:HG12 | 1.79 | 0.64 |
| 26:BA:2042:A:O2' | 26:BA:2044:U:OP2 | 2.16 | 0.64 |
| 31:BF:101:HIS:HD2 | 31:BF:116:PHE:HB2 | 1.61 | 0.64 |
| 36:BK:63:GLU:O | 36:BK:67:GLN:NE2 | 2.30 | 0.64 |
| 1:AA:427:A:O2' | 20:AW:27:GLY:O | 2.15 | 0.64 |
| 7:AG:25:MET:HB3 | 7:AG:45:LEU:HD11 | 1.80 | 0.64 |
| 8:AI:112:ARG:HA | 8:AI:121:PRO:HA | 1.78 | 0.64 |
| 8:AI:189:ARG:NH2 | 22:AY:65:GLU:OE2 | 2.30 | 0.64 |
| 26:BA:546:G:OP2 | 52:Ba:127:ARG:NH2 | 2.30 | 0.64 |
| 26:BA:2001:G:O2' | 26:BA:2003:G:OP2 | 2.14 | 0.64 |
| 53:Bb:50:THR:HG21 | 53:Bb:63:LYS:HG2 | 1.79 | 0.64 |
| 1:AA:27:C:O2' | 13:AO:44:GLN:NE2 | 2.29 | 0.64 |
| 1:AA:1093:A:H5' | 1:AA:1094:C:H5 | 1.62 | 0.64 |
| 7:AG:153:ALA:HB3 | 7:AG:171:TRP:HA | 1.80 | 0.64 |
| 8:AI:1:MET:N | 11:AL:52:ILE:O | 2.30 | 0.64 |
| 26:BA:505:G:OP2 | 44:BS:3:ARG:NH2 | 2.30 | 0.64 |
| 26:BA:525:A:O2' | 26:BA:2064:C:O2 | 2.12 | 0.64 |
| 26:BA:828:G:O2' | 26:BA:1718:A:OP1 | 2.14 | 0.64 |
| 26:BA:2329:A:OP1 | 39:BN:7:TYR:OH | 2.13 | 0.64 |
| 1:AA:118:G:H1 | 1:AA:203:C:H42 | 1.45 | 0.64 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:933:G:H1 | 1:AA:1165:C:H42 | 1.45 | 0.64 |
| 1:AA:1175:C:N4 | 14:AP:147:LYS:O | 2.30 | 0.64 |
| 8:AI:164:GLY:HA2 | 8:AI:169:ASP:HB3 | 1.78 | 0.64 |
| 26:BA:559:G:O2' | 26:BA:588:A:N6 | 2.31 | 0.64 |
| 57:Bf:1:MET:N | 57:Bf:87:LYS:O | 2.30 | 0.64 |
| 1:AA:210:A:O2' | 6:AF:28:PRO:O | 2.15 | 0.64 |
| 1:AA:1140:G:O2' | 7:AG:64:SER:O | 2.14 | 0.64 |
| 25:AT:41:ILE:HD13 | 25:AT:46:ILE:HG22 | 1.80 | 0.64 |
| 26:BA:2741:U:OP1 | 29:BD:265:ASN:ND2 | 2.31 | 0.64 |
| 27:BB:30:U:H2' | 27:BB:31:C:H6 | 1.61 | 0.64 |
| 36:BK:82:ARG:NH2 | 36:BK:132:VAL:OXT | 2.31 | 0.64 |
| 42:BQ:36:ASP:OD1 | 42:BQ:50:ARG:NH2 | 2.31 | 0.64 |
| 48:BW:1:MET:SD | 48:BW:5:ARG:NH2 | 2.71 | 0.64 |
| 51:BZ:13:ILE:HD12 | 51:BZ:34:VAL:HG13 | 1.78 | 0.64 |
| 1:AA:576:G:H5'' | 17:AS:32:GLN:HE22 | 1.63 | 0.64 |
| 1:AA:697:G:N2 | 1:AA:823:G:O2' | 2.31 | 0.64 |
| 1:AA:1234:A:N6 | 1:AA:1318:G:O4' | 2.30 | 0.64 |
| 6:AF:122:ASN:O | 6:AF:134:ASN:ND2 | 2.31 | 0.64 |
| 7:AG:83:VAL:HB | 7:AG:165:ALA:HA | 1.79 | 0.64 |
| 11:AL:47:SER:HB2 | 11:AL:50:LEU:HD11 | 1.80 | 0.64 |
| 26:BA:830:U:O2' | 26:BA:833:G:O2' | 2.15 | 0.64 |
| 26:BA:2301:U:O2' | 26:BA:2387:C:O2 | 2.16 | 0.64 |
| 26:BA:2537:C:O2' | 56:Be:21:ASN:O | 2.15 | 0.64 |
| 40:BO:16:ARG:NH2 | 40:BO:89:LEU:O | 2.31 | 0.64 |
| 49:BX:82:TYR:O | 49:BX:86:ASN:ND2 | 2.28 | 0.64 |
| 1:AA:114:U:OP1 | 6:AF:72:ARG:NH2 | 2.30 | 0.64 |
| 13:AO:139:LYS:HD2 | 13:AO:140:PRO:HD2 | 1.80 | 0.64 |
| 23:AH:64:LYS:N | 23:AH:97:ARG:O | 2.30 | 0.64 |
| 30:BE:6:THR:N | 30:BE:15:GLY:O | 2.30 | 0.64 |
| 44:BS:106:PRO:HA | 44:BS:109:MET:HG2 | 1.80 | 0.64 |
| 3:AC:191:ASP:O | 25:AT:12:THR:OG1 | 2.15 | 0.64 |
| 5:AE:40:GLU:HA | 5:AE:43:LYS:HE3 | 1.79 | 0.64 |
| 5:AE:118:LEU:HD13 | 5:AE:126:VAL:HA | 1.80 | 0.64 |
| 38:BM:113:LEU:HD22 | 38:BM:136:ASP:HA | 1.80 | 0.64 |
| 41:BP:145:GLU:HG3 | 41:BP:147:LEU:HG | 1.79 | 0.64 |
| 2:AB:86:TYR:OH | 2:AB:87:ARG:NH2 | 2.32 | 0.63 |
| 16:AQ:57:ARG:O | 21:AX:24:GLN:NE2 | 2.31 | 0.63 |
| 25:AT:17:LEU:HD21 | 25:AT:54:VAL:HG13 | 1.80 | 0.63 |
| 26:BA:2461:A:N6 | 26:BA:2512:C:O2 | 2.30 | 0.63 |
| 29:BD:81:ARG:NH1 | 29:BD:184:GLY:O | 2.30 | 0.63 |
| 37:BL:92:TYR:HB2 | 37:BL:114:LEU:HD23 | 1.80 | 0.63 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:603:G:O2' | 1:AA:777:G:OP1 | 2.15 | 0.63 |
| 4:AD:182:ARG:NH2 | 4:AD:184:GLU:OE2 | 2.31 | 0.63 |
| 6:AF:225:PRO:HB3 | 6:AF:229:LEU:HD12 | 1.79 | 0.63 |
| 8:AI:84:ILE:HG23 | 8:AI:88:LYS:HE3 | 1.80 | 0.63 |
| 23:AH:17:TYR:HE1 | 23:AH:121:LYS:HE2 | 1.62 | 0.63 |
| 26:BA:186:A:N1 | 26:BA:230:A:O2' | 2.29 | 0.63 |
| 26:BA:653:A:N7 | 26:BA:737:G:O2' | 2.30 | 0.63 |
| 28:BC:163:SER:HB3 | 53:Bb:77:THR:HG21 | 1.79 | 0.63 |
| 31:BF:9:VAL:HG23 | 31:BF:68:ALA:HB1 | 1.80 | 0.63 |
| 46:BU:32:LEU:HG | 46:BU:44:SER:HA | 1.81 | 0.63 |
| 1:AA:69:U:O2' | 1:AA:71:G:N2 | 2.31 | 0.63 |
| 1:AA:174:A:OP1 | 17:AS:3:ARG:NH1 | 2.28 | 0.63 |
| 1:AA:888:G:H1 | 1:AA:1288:U:H3 | 1.45 | 0.63 |
| 26:BA:324:G:O6 | 30:BE:172:LYS:NZ | 2.30 | 0.63 |
| 26:BA:387:G:OP1 | 38:BM:194:ARG:NH1 | 2.31 | 0.63 |
| 35:BJ:120:ASP:OD2 | 35:BJ:122:ARG:NH1 | 2.31 | 0.63 |
| 52:Ba:141:LEU:HD21 | 52:Ba:151:VAL:HG11 | 1.79 | 0.63 |
| 1:AA:499:G:OP1 | 1:AA:499:G:N2 | 2.31 | 0.63 |
| 1:AA:504:U:O2' | 17:AS:62:ARG:NH2 | 2.32 | 0.63 |
| 1:AA:1095:U:O2' | 11:AL:6:ASN:OD1 | 2.16 | 0.63 |
| 1:AA:1107:G:OP2 | 25:AT:56:ARG:NH1 | 2.30 | 0.63 |
| 1:AA:1351:C:O2 | 1:AA:1457:A:O2' | 2.15 | 0.63 |
| 2:AB:77:LYS:NZ | 2:AB:78:THR:O | 2.30 | 0.63 |
| 16:AQ:1:MET:SD | 16:AQ:2:ALA:N | 2.71 | 0.63 |
| 18:AU:76:ILE:HG22 | 18:AU:103:VAL:HG23 | 1.81 | 0.63 |
| 34:BI:54:GLU:HB2 | 34:BI:159:THR:HG23 | 1.79 | 0.63 |
| 1:AA:449:A:O2' | 1:AA:482:G:O2' | 2.15 | 0.63 |
| 4:AD:46:ARG:NH2 | 12:AN:26:THR:OG1 | 2.32 | 0.63 |
| 7:AG:29:ILE:O | 58:AZ:21:TYR:OH | 2.17 | 0.63 |
| 12:AN:98:ALA:HB2 | 12:AN:104:ILE:HD13 | 1.79 | 0.63 |
| 14:AP:70:LEU:HD11 | 14:AP:74:GLU:HB2 | 1.81 | 0.63 |
| 26:BA:1726:G:O2' | 26:BA:2012:U:O4 | 2.14 | 0.63 |
| 1:AA:171:G:H2' | 1:AA:172:G:H5' | 1.80 | 0.63 |
| 1:AA:1128:G:OP1 | 8:AI:43:ARG:NH2 | 2.31 | 0.63 |
| 5:AE:19:GLN:HB3 | 5:AE:22:ARG:HE | 1.63 | 0.63 |
| 18:AU:50:PHE:HB3 | 18:AU:54:GLN:HG3 | 1.80 | 0.63 |
| 23:AH:38:VAL:HG13 | 23:AH:42:ILE:HD12 | 1.80 | 0.63 |
| 23:AH:102:ALA:H | 23:AH:105:ILE:HD12 | 1.62 | 0.63 |
| 26:BA:26:C:N4 | 26:BA:447:A:OP2 | 2.31 | 0.63 |
| 26:BA:1612:A:O2' | 26:BA:1613:A:O4' | 2.17 | 0.63 |
| 1:AA:689:G:O6 | 16:AQ:19:ARG:NH1 | 2.31 | 0.63 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 1:AA:800:G:OP2 | 1:AA:816:G:N1 | 2.29 | 0.63 |
| 16:AQ:63:ALA:O | 16:AQ:67:THR:OG1 | 2.17 | 0.63 |
| 26:BA:878:A:N6 | 28:BC:1:MET:O | 2.32 | 0.63 |
| 26:BA:1004:A:N3 | 27:BB:84:U:O2' | 2.29 | 0.63 |
| 26:BA:1920:A:OP2 | 28:BC:238:ARG:NH1 | 2.31 | 0.63 |
| 45:BT:47:VAL:HG21 | 45:BT:64:LEU:HD23 | 1.81 | 0.63 |
| 1:AA:386:G:OP1 | 5:AE:130:ARG:NE | 2.31 | 0.63 |
| 1:AA:1176:G:OP2 | 14:AP:151:ARG:NH1 | 2.31 | 0.63 |
| 29:BD:40:GLN:HG2 | 29:BD:168:SER:HB2 | 1.80 | 0.63 |
| 31:BF:45:THR:HG22 | 31:BF:53:LYS:HE2 | 1.81 | 0.63 |
| 33:BH:68:SER:OG | 33:BH:73:ALA:O | 2.16 | 0.63 |
| 35:BJ:33:TYR:HD1 | 35:BJ:99:LYS:HB2 | 1.64 | 0.63 |
| 1:AA:52:U:H2' | 1:AA:53:G:H8 | 1.64 | 0.63 |
| 10:AK:88:TYR:HB3 | 10:AK:93:ILE:HB | 1.81 | 0.63 |
| 20:AW:74:GLU:HA | 20:AW:77:MET:HE2 | 1.81 | 0.63 |
| 25:AT:27:ASP:HB3 | 25:AT:30:THR:HB | 1.81 | 0.63 |
| 49:BX:101:GLU:OE1 | 49:BX:103:LYS:NZ | 2.32 | 0.63 |
| 50:BY:39:MET:HE2 | 50:BY:86:ILE:HD13 | 1.79 | 0.63 |
| 1:AA:584:C:O2' | 9:AJ:124:GLN:OE1 | 2.15 | 0.62 |
| 1:AA:1392:U:H5'' | 1:AA:1393:C:H5 | 1.64 | 0.62 |
| 2:AB:84:PHE:O | 2:AB:97:ASP:N | 2.31 | 0.62 |
| 14:AP:96:ARG:HG3 | 14:AP:99:ASP:HB2 | 1.80 | 0.62 |
| 26:BA:1669:U:O2' | 54:Bc:8:MET:O | 2.16 | 0.62 |
| 29:BD:44:GLY:HA3 | 29:BD:77:THR:HG22 | 1.81 | 0.62 |
| 29:BD:62:LEU:O | 36:BK:5:ARG:NH1 | 2.32 | 0.62 |
| 29:BD:167:ILE:O | 29:BD:175:LYS:NZ | 2.31 | 0.62 |
| 1:AA:349:C:O4' | 1:AA:376:G:N2 | 2.32 | 0.62 |
| 5:AE:53:SER:OG | 5:AE:57:LYS:NZ | 2.31 | 0.62 |
| 9:AJ:52:PHE:HE1 | 9:AJ:59:GLY:HA3 | 1.63 | 0.62 |
| 11:AL:43:MET:HA | 11:AL:46:ILE:HG22 | 1.80 | 0.62 |
| 11:AL:105:ARG:HD2 | 11:AL:106:ASN:H | 1.65 | 0.62 |
| 30:BE:50:ARG:HG2 | 30:BE:53:SER:HB3 | 1.79 | 0.62 |
| 31:BF:10:ILE:HB | 31:BF:121:THR:HG23 | 1.81 | 0.62 |
| 49:BX:121:PRO:HG3 | 49:BX:125:HIS:HD2 | 1.64 | 0.62 |
| 1:AA:765:C:H5' | 16:AQ:1:MET:HG2 | 1.80 | 0.62 |
| 1:AA:938:G:N2 | 1:AA:1158:G:O3' | 2.31 | 0.62 |
| 1:AA:951:A:O2' | 1:AA:978:C:OP1 | 2.18 | 0.62 |
| 1:AA:1295:U:H5'' | 11:AL:125:ALA:HB3 | 1.81 | 0.62 |
| 3:AC:46:ILE:HG13 | 3:AC:82:ALA:HA | 1.82 | 0.62 |
| 5:AE:137:HIS:HE1 | 5:AE:167:PRO:HD2 | 1.64 | 0.62 |
| 18:AU:77:LEU:HB2 | 18:AU:80:MET:HB2 | 1.81 | 0.62 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 33:BH:16:LYS:HG2 | 33:BH:112:ILE:HD11 | 1.81 | 0.62 |
| 41:BP:20:ASP:O | 41:BP:53:LYS:NZ | 2.33 | 0.62 |
| 46:BU:86:LYS:HG3 | 46:BU:88:ASP:H | 1.64 | 0.62 |
| 1:AA:457:G:N2 | 1:AA:469:G:O3' | 2.30 | 0.62 |
| 1:AA:716:G:N2 | 1:AA:745:U:O4 | 2.33 | 0.62 |
| 1:AA:1262:U:O2' | 1:AA:1307:A:O2' | 2.17 | 0.62 |
| 19:AV:65:ASP:HB2 | 19:AV:68:ILE:HD11 | 1.80 | 0.62 |
| 31:BF:76:LEU:HA | 31:BF:79:VAL:HG22 | 1.81 | 0.62 |
| 1:AA:1108:C:H41 | 1:AA:1129:G:H22 | 1.47 | 0.62 |
| 2:AB:130:GLN:HG2 | 2:AB:146:LEU:HD12 | 1.81 | 0.62 |
| 5:AE:85:LEU:O | 5:AE:89:GLY:N | 2.30 | 0.62 |
| 6:AF:174:ALA:HB1 | 6:AF:217:VAL:HG23 | 1.81 | 0.62 |
| 11:AL:32:ILE:HD13 | 11:AL:38:THR:HB | 1.82 | 0.62 |
| 16:AQ:52:MET:SD | 21:AX:32:ARG:NH1 | 2.72 | 0.62 |
| 19:AV:12:MET:HE2 | 19:AV:135:LEU:HD12 | 1.82 | 0.62 |
| 24:AM:25:SER:HA | 24:AM:28:GLU:HG2 | 1.81 | 0.62 |
| 26:BA:758:A:N3 | 26:BA:2454:C:O2' | 2.30 | 0.62 |
| 27:BB:25:G:H4' | 27:BB:26:C:H5 | 1.63 | 0.62 |
| 1:AA:81:C:O2' | 1:AA:361:G:OP1 | 2.16 | 0.62 |
| 7:AG:152:ILE:HD13 | 7:AG:162:LEU:HB2 | 1.81 | 0.62 |
| 25:AT:34:LEU:HA | 25:AT:37:LYS:HG2 | 1.81 | 0.62 |
| 26:BA:398:G:OP2 | 38:BM:70:ARG:NH2 | 2.31 | 0.62 |
| 26:BA:2752:A:O2' | 32:BG:157:LYS:NZ | 2.32 | 0.62 |
| 27:BB:2:A:H2' | 27:BB:3:G:H8 | 1.65 | 0.62 |
| 44:BS:65:HIS:O | 44:BS:68:SER:OG | 2.17 | 0.62 |
| 44:BS:68:SER:HB2 | 44:BS:79:LYS:HD3 | 1.81 | 0.62 |
| 1:AA:698:U:O2' | 1:AA:824:C:O2 | 2.18 | 0.62 |
| 4:AD:122:THR:HB | 4:AD:186:ARG:HH11 | 1.64 | 0.62 |
| 15:AR:15:CYS:HB2 | 15:AR:32:LEU:HA | 1.81 | 0.62 |
| 21:AX:16:LYS:HD3 | 21:AX:58:LEU:HD21 | 1.81 | 0.62 |
| 43:BR:11:THR:HG21 | 43:BR:55:PRO:HG2 | 1.80 | 0.62 |
| 1:AA:278:U:O2' | 1:AA:496:C:O2' | 2.11 | 0.62 |
| 2:AB:137:SER:HB2 | 2:AB:144:ALA:HB2 | 1.81 | 0.62 |
| 10:AK:46:THR:OG1 | 10:AK:50:ASN:O | 2.13 | 0.62 |
| 16:AQ:2:ALA:HB1 | 16:AQ:7:LYS:HA | 1.80 | 0.62 |
| 16:AQ:19:ARG:HG2 | 16:AQ:65:LEU:HD13 | 1.82 | 0.62 |
| 26:BA:442:G:O2' | 26:BA:444:C:OP2 | 2.15 | 0.62 |
| 26:BA:1398:G:OP1 | 54:Bc:14:ARG:NH1 | 2.29 | 0.62 |
| 27:BB:18:C:H4' | 39:BN:1:MET:HA | 1.81 | 0.62 |
| 1:AA:1035:C:O2' | 1:AA:1117:A:O2' | 2.17 | 0.62 |
| 18:AU:61:PHE:HZ | 18:AU:85:ILE:HG22 | 1.65 | 0.62 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 21:AX:14:ARG:HB2 | 21:AX:61:LEU:HD11 | 1.81 | 0.62 |
| 24:AM:8:LEU:HD23 | 24:AM:94:ILE:HG21 | 1.82 | 0.62 |
| 1:AA:1320:G:OP1 | 8:AI:36:HIS:NE2 | 2.29 | 0.62 |
| 3:AC:184:GLN:HG2 | 3:AC:185:PRO:HD2 | 1.82 | 0.62 |
| 5:AE:51:TYR:OH | 5:AE:84:LYS:NZ | 2.31 | 0.62 |
| 17:AS:35:VAL:HG12 | 17:AS:85:ILE:HD11 | 1.81 | 0.62 |
| 20:AW:73:ASP:OD2 | 20:AW:76:ARG:N | 2.33 | 0.62 |
| 26:BA:109:G:OP1 | 54:Bc:20:ARG:NH1 | 2.33 | 0.62 |
| 26:BA:228:A:O2' | 26:BA:430:A:N3 | 2.32 | 0.62 |
| 26:BA:969:G:N2 | 26:BA:983:C:O2 | 2.32 | 0.62 |
| 26:BA:1862:C:O2' | 26:BA:1992:U:OP2 | 2.16 | 0.62 |
| 1:AA:417:U:H5' | 20:AW:31:SER:HA | 1.81 | 0.61 |
| 19:AV:37:THR:OG1 | 19:AV:54:ARG:NH2 | 2.30 | 0.61 |
| 26:BA:188:A:O2' | 26:BA:421:C:O2' | 2.15 | 0.61 |
| 29:BD:53:ILE:HG13 | 47:BV:15:PRO:HB2 | 1.82 | 0.61 |
| 33:BH:21:LEU:HD22 | 33:BH:97:VAL:HG21 | 1.82 | 0.61 |
| 40:BO:63:ALA:N | 40:BO:66:ASP:OD2 | 2.33 | 0.61 |
| 1:AA:442:C:O2 | 1:AA:482:G:N2 | 2.28 | 0.61 |
| 1:AA:473:A:O2' | 1:AA:475:A:OP2 | 2.14 | 0.61 |
| 1:AA:986:G:O2' | 1:AA:1162:U:O2' | 2.16 | 0.61 |
| 1:AA:1296:A:H5'' | 11:AL:127:ALA:HB2 | 1.81 | 0.61 |
| 3:AC:95:MET:HE3 | 3:AC:116:THR:HG23 | 1.82 | 0.61 |
| 17:AS:84:ASP:HA | 17:AS:108:PRO:HD3 | 1.82 | 0.61 |
| 26:BA:2540:G:OP1 | 32:BG:164:ARG:NH1 | 2.33 | 0.61 |
| 57:Bf:64:ARG:HA | 57:Bf:84:ARG:HA | 1.82 | 0.61 |
| 1:AA:1441:A:OP1 | 1:AA:1469:A:O2' | 2.17 | 0.61 |
| 26:BA:728:A:O2' | 26:BA:2382:C:O2' | 2.18 | 0.61 |
| 26:BA:1668:C:O2 | 54:Bc:12:GLN:NE2 | 2.32 | 0.61 |
| 26:BA:2113:U:OP2 | 38:BM:25:ARG:NH1 | 2.30 | 0.61 |
| 26:BA:2354:U:OP1 | 57:Bf:8:ARG:NH1 | 2.32 | 0.61 |
| 35:BJ:8:ASP:OD1 | 35:BJ:36:ASN:ND2 | 2.33 | 0.61 |
| 1:AA:180:U:OP2 | 6:AF:129:GLY:N | 2.34 | 0.61 |
| 3:AC:131:VAL:HG12 | 3:AC:145:LYS:HA | 1.82 | 0.61 |
| 4:AD:159:GLN:O | 4:AD:163:LEU:HB2 | 2.00 | 0.61 |
| 18:AU:76:ILE:HG13 | 18:AU:107:PHE:HE1 | 1.64 | 0.61 |
| 20:AW:52:ILE:HG22 | 20:AW:68:ALA:HB2 | 1.82 | 0.61 |
| 35:BJ:85:PRO:HG3 | 35:BJ:90:ARG:HH21 | 1.65 | 0.61 |
| 39:BN:46:ILE:HD11 | 39:BN:119:ALA:HB1 | 1.81 | 0.61 |
| 1:AA:419:U:OP1 | 20:AW:54:ARG:NH1 | 2.33 | 0.61 |
| 5:AE:85:LEU:HD13 | 5:AE:90:ILE:HD11 | 1.81 | 0.61 |
| 13:AO:58:LYS:HB2 | 13:AO:113:ASP:HB2 | 1.82 | 0.61 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|-------------------|--------------------------|-------------------|
| 20:AW:5:ILE:HG12 | 20:AW:21:PHE:HB3 | 1.81 | 0.61 |
| 2:AB:101:THR:HA | 2:AB:208:PRO:HG2 | 1.83 | 0.61 |
| 6:AF:50:VAL:HG21 | 6:AF:56:GLY:HA2 | 1.81 | 0.61 |
| 6:AF:154:ILE:HG23 | 6:AF:163:VAL:HB | 1.83 | 0.61 |
| 26:BA:808:G:OP1 | 41:BP:92:LYS:NZ | 2.29 | 0.61 |
| 29:BD:41:SER:OG | 29:BD:310:ARG:NH2 | 2.32 | 0.61 |
| 1:AA:1180:G:O2' | 1:AA:1312:A:OP1 | 2.16 | 0.61 |
| 1:AA:1262:U:OP2 | 18:AU:44:ARG:NH2 | 2.31 | 0.61 |
| 16:AQ:131:GLN:NE2 | 16:AQ:132:GLU:OE2 | 2.30 | 0.61 |
| 26:BA:1873:U:O2' | 26:BA:1948:A:N3 | 2.28 | 0.61 |
| 26:BA:2061:A:OP2 | 35:BJ:86:TYR:OH | 2.15 | 0.61 |
| 49:BX:6:ARG:NH1 | 49:BX:26:LEU:O | 2.32 | 0.61 |
| 1:AA:1023:A:H1' | 1:AA:1042:C:H41 | 1.66 | 0.61 |
| 2:AB:69:GLY:HA2 | 2:AB:72:ILE:HD13 | 1.82 | 0.61 |
| 26:BA:835:A:OP1 | 26:BA:1672:C:N4 | 2.33 | 0.61 |
| 26:BA:1239:C:O2' | 42:BQ:43:GLY:O | 2.16 | 0.61 |
| 26:BA:1571:A:N1 | 26:BA:1584:C:O2' | 2.33 | 0.61 |
| 1:AA:503:A:O2' | 1:AA:506:G:O2' | 2.14 | 0.61 |
| 2:AB:101:THR:OG1 | 2:AB:208:PRO:O | 2.15 | 0.61 |
| 26:BA:861:U:O2 | 28:BC:1:MET:N | 2.27 | 0.61 |
| 26:BA:2085:C:O2' | 26:BA:2260:G:N2 | 2.34 | 0.61 |
| 31:BF:39:ARG:HA | 31:BF:59:CYS:HA | 1.83 | 0.61 |
| 35:BJ:134:GLU:OE2 | 35:BJ:138:LYS:NZ | 2.34 | 0.61 |
| 50:BY:11:SER:HA | 50:BY:14:LYS:HE2 | 1.81 | 0.61 |
| 1:AA:684:A:H4' | 16:AQ:8:ARG:HH11 | 1.65 | 0.61 |
| 2:AB:78:THR:OG1 | 2:AB:80:ASP:OD1 | 2.14 | 0.61 |
| 30:BE:114:ILE:HG12 | 30:BE:225:ALA:HB2 | 1.83 | 0.61 |
| 1:AA:809:A:O2' | 1:AA:1017:U:O4 | 2.17 | 0.60 |
| 1:AA:1276:G:H5'' | 14:AP:48:GLY:H | 1.66 | 0.60 |
| 7:AG:75:ILE:HG12 | 7:AG:161:VAL:HG23 | 1.81 | 0.60 |
| 11:AL:118:LYS:HA | 11:AL:125:ALA:HA | 1.83 | 0.60 |
| 11:AL:122:GLY:HA3 | 11:AL:128:LYS:HA | 1.83 | 0.60 |
| 14:AP:98:LYS:HE3 | 18:AU:37:LEU:HA | 1.83 | 0.60 |
| 1:AA:890:G:N1 | 1:AA:1285:G:OP2 | 2.34 | 0.60 |
| 12:AN:32:ALA:HB1 | 12:AN:59:LEU:HD22 | 1.83 | 0.60 |
| 13:AO:125:ASN:OD1 | 13:AO:139:LYS:NZ | 2.27 | 0.60 |
| 20:AW:51:ILE:HD11 | 20:AW:85:ILE:HG22 | 1.83 | 0.60 |
| 34:BI:12:VAL:HG22 | 34:BI:57:GLN:HG3 | 1.83 | 0.60 |
| 40:BO:67:VAL:HA | 40:BO:85:THR:HG23 | 1.83 | 0.60 |
| 7:AG:77:GLY:HA2 | 7:AG:83:VAL:HA | 1.81 | 0.60 |
| 26:BA:1741:G:N2 | 41:BP:81:ARG:O | 2.33 | 0.60 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 26:BA:2601:C:OP1 | 28:BC:208:LYS:NZ | 2.32 | 0.60 |
| 29:BD:86:ARG:HG3 | 29:BD:100:GLU:HG2 | 1.82 | 0.60 |
| 32:BG:88:MET:HE2 | 32:BG:170:ILE:HG22 | 1.81 | 0.60 |
| 50:BY:41:VAL:HB | 50:BY:84:MET:HB3 | 1.82 | 0.60 |
| 3:AC:126:LEU:HB2 | 3:AC:183:ILE:HG23 | 1.82 | 0.60 |
| 4:AD:103:ILE:HB | 4:AD:123:CYS:HB2 | 1.83 | 0.60 |
| 26:BA:1052:U:O2' | 26:BA:2283:A:N3 | 2.34 | 0.60 |
| 36:BK:115:ARG:N | 36:BK:132:VAL:O | 2.35 | 0.60 |
| 1:AA:352:U:O2' | 1:AA:420:A:N1 | 2.33 | 0.60 |
| 16:AQ:77:GLU:OE1 | 16:AQ:78:ASN:ND2 | 2.35 | 0.60 |
| 19:AV:36:LYS:NZ | 19:AV:41:LYS:O | 2.30 | 0.60 |
| 25:AT:15:SER:O | 25:AT:19:ASN:ND2 | 2.33 | 0.60 |
| 34:BI:138:VAL:HG11 | 34:BI:146:ALA:HB2 | 1.83 | 0.60 |
| 1:AA:666:G:OP2 | 1:AA:799:U:O2' | 2.20 | 0.60 |
| 5:AE:19:GLN:H | 5:AE:22:ARG:HH21 | 1.47 | 0.60 |
| 9:AJ:75:ILE:HG13 | 9:AJ:127:ALA:HA | 1.83 | 0.60 |
| 29:BD:263:GLU:HB3 | 29:BD:266:LYS:HE2 | 1.82 | 0.60 |
| 1:AA:416:G:O2' | 20:AW:29:THR:O | 2.16 | 0.60 |
| 1:AA:430:U:H2' | 1:AA:431:G:H8 | 1.65 | 0.60 |
| 1:AA:554:A:H5'' | 5:AE:84:LYS:HG3 | 1.84 | 0.60 |
| 21:AX:18:ASN:HB2 | 21:AX:55:THR:HG22 | 1.83 | 0.60 |
| 24:AM:9:SER:HB2 | 24:AM:70:LYS:HG2 | 1.82 | 0.60 |
| 26:BA:1108:A:OP2 | 35:BJ:128:ARG:NH2 | 2.29 | 0.60 |
| 26:BA:1398:G:O6 | 55:Bd:1:MET:N | 2.33 | 0.60 |
| 45:BT:16:LEU:HB3 | 45:BT:21:LYS:HB2 | 1.84 | 0.60 |
| 46:BU:83:ILE:HD12 | 46:BU:91:GLU:HB3 | 1.83 | 0.60 |
| 2:AB:71:HIS:HB2 | 2:AB:95:VAL:HG13 | 1.81 | 0.60 |
| 13:AO:98:VAL:HG12 | 13:AO:124:VAL:HA | 1.82 | 0.60 |
| 20:AW:17:ARG:HB2 | 20:AW:70:VAL:HB | 1.83 | 0.60 |
| 21:AX:26:ILE:HG22 | 21:AX:34:ILE:HG21 | 1.83 | 0.60 |
| 23:AH:57:ASP:OD1 | 23:AH:97:ARG:NH1 | 2.34 | 0.60 |
| 26:BA:1922:A:O2' | 28:BC:214:THR:OG1 | 2.14 | 0.60 |
| 39:BN:87:LEU:HD12 | 39:BN:126:SER:HB3 | 1.83 | 0.60 |
| 48:BW:17:ARG:HD2 | 48:BW:61:GLN:HG2 | 1.83 | 0.60 |
| 52:Ba:129:ALA:HB3 | 52:Ba:132:ILE:HG12 | 1.83 | 0.60 |
| 1:AA:235:A:O2' | 17:AS:43:MET:SD | 2.59 | 0.60 |
| 1:AA:614:G:O3' | 4:AD:106:ASN:ND2 | 2.35 | 0.60 |
| 1:AA:1266:A:H3' | 18:AU:106:ARG:HH12 | 1.66 | 0.60 |
| 6:AF:92:GLU:OE1 | 6:AF:94:TYR:OH | 2.15 | 0.60 |
| 13:AO:38:PRO:HD2 | 13:AO:119:PHE:HZ | 1.67 | 0.60 |
| 14:AP:61:VAL:HG21 | 14:AP:78:LEU:HD21 | 1.83 | 0.60 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 52:Ba:126:ILE:HG23 | 52:Ba:149:ILE:HG21 | 1.82 | 0.60 |
| 1:AA:233:G:H1' | 10:AK:8:ARG:HH21 | 1.67 | 0.60 |
| 1:AA:809:A:OP1 | 7:AG:138:SER:OG | 2.17 | 0.60 |
| 1:AA:889:U:H2' | 1:AA:890:G:H5' | 1.84 | 0.60 |
| 1:AA:1307:A:H2' | 1:AA:1308:A:H8 | 1.67 | 0.60 |
| 26:BA:2301:U:OP1 | 26:BA:2393:U:O2' | 2.16 | 0.60 |
| 6:AF:124:THR:O | 6:AF:132:GLN:N | 2.35 | 0.59 |
| 7:AG:143:LEU:HD23 | 7:AG:191:ALA:HB1 | 1.84 | 0.59 |
| 14:AP:55:ILE:HD11 | 14:AP:85:PHE:HD1 | 1.67 | 0.59 |
| 32:BG:22:ASP:HB2 | 32:BG:36:LYS:HE3 | 1.84 | 0.59 |
| 1:AA:1233:C:H2' | 1:AA:1234:A:H4' | 1.82 | 0.59 |
| 3:AC:188:VAL:HG12 | 3:AC:189:LEU:HD23 | 1.84 | 0.59 |
| 11:AL:130:GLN:NE2 | 11:AL:132:SER:O | 2.32 | 0.59 |
| 23:AH:58:GLY:HA2 | 23:AH:106:ILE:HD12 | 1.84 | 0.59 |
| 9:AJ:82:GLY:N | 9:AJ:85:SER:OG | 2.32 | 0.59 |
| 36:BK:112:PRO:HA | 36:BK:130:ILE:HG13 | 1.84 | 0.59 |
| 1:AA:1236:A:N6 | 1:AA:1237:C:O2 | 2.36 | 0.59 |
| 4:AD:39:ASP:OD1 | 4:AD:42:LEU:N | 2.34 | 0.59 |
| 26:BA:1451:G:O2' | 26:BA:1845:A:N3 | 2.34 | 0.59 |
| 26:BA:2344:U:OP2 | 39:BN:14:ARG:NE | 2.35 | 0.59 |
| 30:BE:196:LEU:HD23 | 30:BE:238:THR:HG22 | 1.84 | 0.59 |
| 32:BG:122:ILE:HD12 | 32:BG:128:VAL:HG21 | 1.82 | 0.59 |
| 1:AA:172:G:H5'' | 17:AS:30:ARG:HH22 | 1.66 | 0.59 |
| 1:AA:943:C:H2' | 1:AA:944:A:H8 | 1.66 | 0.59 |
| 8:AI:97:LEU:HD13 | 8:AI:162:LEU:HD21 | 1.83 | 0.59 |
| 11:AL:120:PHE:HD2 | 24:AM:59:ALA:HB1 | 1.67 | 0.59 |
| 13:AO:65:ARG:NH2 | 13:AO:113:ASP:OD2 | 2.35 | 0.59 |
| 14:AP:24:VAL:HG12 | 14:AP:33:GLY:HA2 | 1.85 | 0.59 |
| 15:AR:41:ALA:HA | 15:AR:44:MET:HG2 | 1.84 | 0.59 |
| 17:AS:88:ILE:HG22 | 17:AS:102:VAL:HG12 | 1.83 | 0.59 |
| 24:AM:78:ASP:HB3 | 24:AM:81:ALA:HB3 | 1.84 | 0.59 |
| 26:BA:177:C:OP2 | 54:Bc:44:ARG:NH2 | 2.32 | 0.59 |
| 31:BF:44:ARG:HA | 31:BF:53:LYS:HG2 | 1.85 | 0.59 |
| 31:BF:45:THR:OG1 | 31:BF:51:ILE:O | 2.15 | 0.59 |
| 1:AA:1278:G:H4' | 1:AA:1279:A:H5' | 1.84 | 0.59 |
| 4:AD:112:LYS:HB2 | 4:AD:153:ASP:HB3 | 1.83 | 0.59 |
| 5:AE:46:SER:HA | 5:AE:49:ARG:HG2 | 1.83 | 0.59 |
| 22:AY:42:ASP:HB3 | 22:AY:45:ARG:HD3 | 1.84 | 0.59 |
| 26:BA:701:A:O2' | 40:BO:38:ASP:OD2 | 2.16 | 0.59 |
| 26:BA:1315:A:OP2 | 52:Ba:65:ASN:ND2 | 2.30 | 0.59 |
| 52:Ba:33:GLU:OE2 | 52:Ba:36:ARG:NH2 | 2.36 | 0.59 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:681:C:H2' | 1:AA:682:A:H8 | 1.67 | 0.59 |
| 1:AA:1103:G:O2' | 1:AA:1126:A:N6 | 2.36 | 0.59 |
| 1:AA:1317:G:N7 | 11:AL:115:LYS:HE3 | 2.18 | 0.59 |
| 3:AC:108:TYR:O | 3:AC:112:ALA:N | 2.35 | 0.59 |
| 8:AI:20:ASP:HB2 | 8:AI:102:ALA:HB1 | 1.84 | 0.59 |
| 22:AY:38:LEU:HD12 | 22:AY:43:LYS:HE3 | 1.84 | 0.59 |
| 26:BA:205:A:OP2 | 26:BA:206:A:O2' | 2.13 | 0.59 |
| 39:BN:170:PHE:O | 39:BN:174:SER:OG | 2.18 | 0.59 |
| 56:Be:14:LYS:HE3 | 56:Be:23:ARG:HD3 | 1.85 | 0.59 |
| 1:AA:1203:U:O2 | 1:AA:1204:C:N4 | 2.35 | 0.59 |
| 5:AE:136:GLY:HA3 | 5:AE:144:LYS:HE3 | 1.84 | 0.59 |
| 30:BE:118:ILE:HG13 | 30:BE:228:LEU:HD13 | 1.84 | 0.59 |
| 45:BT:56:MET:HE1 | 55:Bd:9:LYS:HG3 | 1.84 | 0.59 |
| 47:BV:4:ARG:O | 47:BV:13:LEU:N | 2.31 | 0.59 |
| 1:AA:353:A:N6 | 1:AA:370:G:OP2 | 2.35 | 0.59 |
| 1:AA:607:G:H5' | 1:AA:667:C:H1' | 1.84 | 0.59 |
| 1:AA:771:G:OP1 | 2:AB:79:GLN:N | 2.36 | 0.59 |
| 26:BA:1354:G:O2' | 26:BA:2033:G:O6 | 2.17 | 0.59 |
| 26:BA:2815:A:O2' | 44:BS:71:ARG:NH2 | 2.36 | 0.59 |
| 57:Bf:11:CYS:CB | 57:Bf:14:CYS:SG | 2.89 | 0.59 |
| 26:BA:1515:C:O2' | 26:BA:1638:A:OP1 | 2.13 | 0.59 |
| 26:BA:1776:U:OP1 | 41:BP:104:ARG:NH2 | 2.32 | 0.59 |
| 57:Bf:19:GLU:O | 57:Bf:72:THR:OG1 | 2.15 | 0.59 |
| 1:AA:938:G:N2 | 1:AA:1158:G:O2' | 2.36 | 0.58 |
| 1:AA:1049:G:O2' | 25:AT:6:GLU:OE2 | 2.13 | 0.58 |
| 1:AA:1109:C:H2' | 1:AA:1110:G:H8 | 1.68 | 0.58 |
| 10:AK:60:VAL:HA | 10:AK:75:PRO:HA | 1.85 | 0.58 |
| 14:AP:23:LEU:HA | 14:AP:33:GLY:HA3 | 1.84 | 0.58 |
| 30:BE:159:LEU:HD13 | 30:BE:168:VAL:HG21 | 1.84 | 0.58 |
| 36:BK:1:MET:HE2 | 36:BK:36:LYS:HD2 | 1.85 | 0.58 |
| 1:AA:209:G:N2 | 6:AF:29:GLY:O | 2.35 | 0.58 |
| 1:AA:1088:U:H2' | 1:AA:1089:G:H8 | 1.68 | 0.58 |
| 7:AG:189:TYR:O | 7:AG:193:ASN:ND2 | 2.31 | 0.58 |
| 1:AA:1312:A:H2' | 1:AA:1313:G:H8 | 1.68 | 0.58 |
| 4:AD:119:VAL:HG12 | 4:AD:121:PRO:HD3 | 1.84 | 0.58 |
| 4:AD:131:SER:HA | 4:AD:134:MET:HE3 | 1.84 | 0.58 |
| 20:AW:14:LEU:HG | 20:AW:76:ARG:HH21 | 1.68 | 0.58 |
| 26:BA:1769:U:OP2 | 41:BP:69:LYS:NZ | 2.36 | 0.58 |
| 42:BQ:9:VAL:HG11 | 42:BQ:38:THR:HG21 | 1.86 | 0.58 |
| 43:BR:14:LYS:NZ | 43:BR:55:PRO:O | 2.28 | 0.58 |
| 1:AA:302:U:H3 | 1:AA:315:A:H61 | 1.51 | 0.58 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:462:C:H41 | 13:AO:65:ARG:HH12 | 1.50 | 0.58 |
| 26:BA:882:C:O2' | 54:Bc:50:TRP:O | 2.15 | 0.58 |
| 1:AA:386:G:OP2 | 5:AE:130:ARG:NH2 | 2.33 | 0.58 |
| 2:AB:184:SER:HA | 58:AZ:40:ARG:HH12 | 1.67 | 0.58 |
| 6:AF:30:PRO:HG2 | 6:AF:31:HIS:HD2 | 1.68 | 0.58 |
| 6:AF:80:GLY:N | 6:AF:83:ASP:OD2 | 2.31 | 0.58 |
| 18:AU:34:PHE:HA | 18:AU:37:LEU:HD12 | 1.83 | 0.58 |
| 19:AV:113:LYS:HA | 19:AV:118:ARG:HG2 | 1.85 | 0.58 |
| 26:BA:1318:C:OP1 | 52:Ba:130:GLY:N | 2.30 | 0.58 |
| 2:AB:239:GLU:OE1 | 2:AB:241:THR:OG1 | 2.18 | 0.58 |
| 3:AC:130:VAL:HB | 3:AC:180:VAL:HG22 | 1.86 | 0.58 |
| 8:AI:107:ARG:NH2 | 22:AY:62:MET:SD | 2.77 | 0.58 |
| 10:AK:80:ILE:N | 10:AK:99:VAL:O | 2.35 | 0.58 |
| 18:AU:60:GLU:OE1 | 18:AU:65:LYS:NZ | 2.31 | 0.58 |
| 26:BA:526:U:OP2 | 26:BA:2788:A:N6 | 2.36 | 0.58 |
| 48:BW:3:ILE:HG12 | 48:BW:49:GLU:HG3 | 1.86 | 0.58 |
| 1:AA:1172:A:OP1 | 14:AP:146:THR:N | 2.35 | 0.58 |
| 2:AB:87:ARG:HH22 | 2:AB:89:ARG:HH21 | 1.52 | 0.58 |
| 11:AL:12:LYS:HA | 11:AL:114:GLN:HG3 | 1.86 | 0.58 |
| 1:AA:184:C:O2' | 10:AK:62:ASN:OD1 | 2.16 | 0.58 |
| 1:AA:1126:A:H2' | 1:AA:1127:A:H8 | 1.69 | 0.58 |
| 1:AA:1255:C:OP1 | 14:AP:141:VAL:N | 2.33 | 0.58 |
| 7:AG:199:ARG:HD3 | 9:AJ:68:ARG:HA | 1.85 | 0.58 |
| 26:BA:2667:G:O2' | 26:BA:2676:G:O6 | 2.22 | 0.58 |
| 6:AF:139:THR:OG1 | 6:AF:159:ASP:O | 2.21 | 0.58 |
| 21:AX:17:CYS:HB3 | 21:AX:22:ASN:H | 1.69 | 0.58 |
| 26:BA:340:U:H4' | 46:BU:4:MET:HB2 | 1.86 | 0.58 |
| 26:BA:1973:A:N3 | 26:BA:2571:U:O2' | 2.30 | 0.58 |
| 38:BM:121:VAL:HG21 | 38:BM:131:GLU:HG3 | 1.84 | 0.58 |
| 1:AA:1241:U:H2' | 1:AA:1242:C:H6 | 1.69 | 0.58 |
| 9:AJ:28:ARG:HB3 | 9:AJ:29:PRO:HD3 | 1.86 | 0.58 |
| 13:AO:98:VAL:HG12 | 13:AO:124:VAL:HG23 | 1.85 | 0.58 |
| 14:AP:108:LEU:HB2 | 14:AP:113:ILE:HG12 | 1.85 | 0.58 |
| 25:AT:34:LEU:O | 25:AT:38:TYR:HB2 | 2.04 | 0.58 |
| 26:BA:916:G:O2' | 37:BL:24:GLY:O | 2.21 | 0.58 |
| 1:AA:50:C:H2' | 1:AA:334:C:H41 | 1.68 | 0.57 |
| 1:AA:602:G:H1' | 21:AX:48:GLY:HA3 | 1.85 | 0.57 |
| 1:AA:909:G:H21 | 24:AM:55:GLY:HA3 | 1.69 | 0.57 |
| 1:AA:1430:A:OP2 | 13:AO:59:GLN:NE2 | 2.31 | 0.57 |
| 4:AD:48:VAL:HG13 | 12:AN:25:ILE:HG22 | 1.86 | 0.57 |
| 5:AE:71:GLU:HB3 | 5:AE:74:TYR:HB2 | 1.86 | 0.57 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 6:AF:67:VAL:HG22 | 6:AF:85:ILE:HD12 | 1.86 | 0.57 |
| 8:AI:6:LYS:HA | 8:AI:12:ASP:HA | 1.86 | 0.57 |
| 12:AN:18:THR:CG2 | 12:AN:55:MET:HE1 | 2.34 | 0.57 |
| 19:AV:110:PHE:O | 19:AV:121:SER:N | 2.31 | 0.57 |
| 26:BA:2363:U:O2' | 43:BR:84:GLU:O | 2.20 | 0.57 |
| 1:AA:884:A:N3 | 1:AA:1323:U:O2' | 2.33 | 0.57 |
| 1:AA:1006:A:N1 | 1:AA:1047:G:O2' | 2.32 | 0.57 |
| 5:AE:130:ARG:HD2 | 5:AE:148:PRO:HB3 | 1.84 | 0.57 |
| 8:AI:16:VAL:HG21 | 8:AI:94:ILE:HG13 | 1.87 | 0.57 |
| 16:AQ:75:LEU:HA | 16:AQ:80:VAL:HG23 | 1.85 | 0.57 |
| 28:BC:19:ALA:HB2 | 28:BC:184:ALA:HB2 | 1.87 | 0.57 |
| 29:BD:276:GLU:O | 29:BD:332:ARG:NH1 | 2.37 | 0.57 |
| 32:BG:99:VAL:HG12 | 32:BG:130:VAL:HG11 | 1.85 | 0.57 |
| 39:BN:1:MET:HG3 | 39:BN:3:THR:HG22 | 1.85 | 0.57 |
| 26:BA:2864:C:O4' | 47:BV:56:LYS:NZ | 2.32 | 0.57 |
| 1:AA:244:A:OP2 | 10:AK:29:ARG:NH2 | 2.36 | 0.57 |
| 1:AA:1326:G:N2 | 22:AY:27:HIS:O | 2.32 | 0.57 |
| 5:AE:176:ARG:NH1 | 5:AE:177:PRO:O | 2.37 | 0.57 |
| 20:AW:35:VAL:HG21 | 20:AW:55:ILE:HD13 | 1.86 | 0.57 |
| 22:AY:47:ILE:HG23 | 22:AY:69:GLU:HB3 | 1.87 | 0.57 |
| 26:BA:1103:G:N2 | 42:BQ:44:SER:HB2 | 2.19 | 0.57 |
| 52:Ba:28:LEU:HD21 | 52:Ba:30:MET:HE3 | 1.86 | 0.57 |
| 54:Bc:21:ARG:HD2 | 54:Bc:45:MET:HE1 | 1.84 | 0.57 |
| 1:AA:390:C:O2' | 5:AE:175:GLU:OE1 | 2.21 | 0.57 |
| 11:AL:1:MET:HG2 | 11:AL:2:VAL:HG23 | 1.87 | 0.57 |
| 16:AQ:130:LYS:NZ | 16:AQ:136:PRO:O | 2.29 | 0.57 |
| 21:AX:14:ARG:O | 21:AX:58:LEU:N | 2.33 | 0.57 |
| 26:BA:776:G:O2' | 26:BA:1441:A:N3 | 2.36 | 0.57 |
| 30:BE:130:ARG:HH22 | 30:BE:193:LYS:HB2 | 1.70 | 0.57 |
| 31:BF:8:LYS:HE3 | 31:BF:10:ILE:HD11 | 1.86 | 0.57 |
| 33:BH:47:LYS:N | 33:BH:99:ILE:O | 2.38 | 0.57 |
| 1:AA:192:G:H1 | 1:AA:201:U:H3 | 1.51 | 0.57 |
| 1:AA:1126:A:H4' | 11:AL:109:VAL:HG12 | 1.84 | 0.57 |
| 6:AF:24:SER:HB3 | 6:AF:41:ILE:HD11 | 1.86 | 0.57 |
| 6:AF:68:ASP:HB3 | 6:AF:160:LYS:HE2 | 1.85 | 0.57 |
| 6:AF:139:THR:OG1 | 6:AF:160:LYS:HB2 | 2.05 | 0.57 |
| 18:AU:76:ILE:HG23 | 18:AU:80:MET:HE2 | 1.86 | 0.57 |
| 26:BA:1048:A:H61 | 34:BI:113:MET:HE3 | 1.68 | 0.57 |
| 26:BA:1659:U:OP2 | 55:Bd:9:LYS:NZ | 2.30 | 0.57 |
| 32:BG:19:LEU:HD13 | 32:BG:24:PHE:HD1 | 1.69 | 0.57 |
| 50:BY:16:VAL:HG23 | 50:BY:21:VAL:HG22 | 1.85 | 0.57 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:416:G:N3 | 20:AW:29:THR:OG1 | 2.37 | 0.57 |
| 26:BA:1550:U:O3' | 26:BA:1614:G:O2' | 2.23 | 0.57 |
| 26:BA:1951:G:O2' | 26:BA:1989:G:O6 | 2.22 | 0.57 |
| 29:BD:155:VAL:HG12 | 29:BD:157:LYS:HG2 | 1.85 | 0.57 |
| 41:BP:141:HIS:O | 41:BP:145:GLU:HG2 | 2.05 | 0.57 |
| 1:AA:134:G:H2' | 1:AA:135:G:H5' | 1.87 | 0.57 |
| 1:AA:1113:G:N1 | 1:AA:1116:A:OP2 | 2.37 | 0.57 |
| 1:AA:1198:A:H2' | 1:AA:1199:U:H6 | 1.70 | 0.57 |
| 7:AG:147:PRO:HG2 | 7:AG:150:LEU:HD13 | 1.86 | 0.57 |
| 21:AX:39:CYS:SG | 21:AX:40:GLY:N | 2.76 | 0.57 |
| 24:AM:1:MET:HE1 | 24:AM:3:LYS:HE3 | 1.85 | 0.57 |
| 1:AA:98:C:OP1 | 1:AA:293:G:O2' | 2.23 | 0.57 |
| 1:AA:259:C:H1' | 10:AK:12:THR:HG22 | 1.87 | 0.57 |
| 1:AA:1035:C:H2' | 1:AA:1036:A:C8 | 2.40 | 0.57 |
| 1:AA:1433:U:O2' | 26:BA:1940:A:N1 | 2.28 | 0.57 |
| 23:AH:45:LEU:HB3 | 23:AH:48:TYR:HB2 | 1.86 | 0.57 |
| 26:BA:1687:C:OP1 | 41:BP:60:ARG:NH2 | 2.32 | 0.57 |
| 27:BB:2:A:H2' | 27:BB:3:G:C8 | 2.40 | 0.57 |
| 48:BW:46:ARG:HE | 48:BW:50:ILE:HD11 | 1.70 | 0.57 |
| 52:Ba:119:VAL:HG13 | 52:Ba:149:ILE:HD13 | 1.87 | 0.57 |
| 1:AA:309:A:O2' | 1:AA:310:C:O4' | 2.22 | 0.57 |
| 1:AA:387:U:OP1 | 1:AA:388:G:O2' | 2.14 | 0.57 |
| 1:AA:524:G:N2 | 1:AA:824:C:O3' | 2.38 | 0.57 |
| 1:AA:662:U:H5' | 4:AD:128:ARG:HH21 | 1.69 | 0.57 |
| 13:AO:63:ALA:HB3 | 13:AO:65:ARG:HE | 1.69 | 0.57 |
| 17:AS:5:ILE:H | 17:AS:8:ASN:HB3 | 1.69 | 0.57 |
| 24:AM:99:VAL:HG12 | 24:AM:101:GLU:HG3 | 1.86 | 0.57 |
| 34:BI:70:ALA:HB3 | 34:BI:85:MET:HE1 | 1.86 | 0.57 |
| 35:BJ:14:LEU:HD22 | 35:BJ:42:ILE:HD11 | 1.87 | 0.57 |
| 55:Bd:12:LEU:HD23 | 55:Bd:50:VAL:HG11 | 1.86 | 0.57 |
| 7:AG:41:ILE:HD13 | 7:AG:106:LEU:HD23 | 1.87 | 0.56 |
| 7:AG:126:HIS:HA | 7:AG:168:LYS:HZ3 | 1.70 | 0.56 |
| 11:AL:100:PHE:HA | 11:AL:103:TYR:HE1 | 1.70 | 0.56 |
| 14:AP:44:LEU:HD12 | 14:AP:45:PRO:HD2 | 1.87 | 0.56 |
| 39:BN:70:TYR:HE1 | 39:BN:172:GLU:HG2 | 1.70 | 0.56 |
| 40:BO:35:ILE:HD12 | 40:BO:124:ILE:HD11 | 1.87 | 0.56 |
| 50:BY:50:ILE:HG22 | 50:BY:53:LYS:HB3 | 1.86 | 0.56 |
| 3:AC:11:GLY:HA2 | 3:AC:14:LYS:HG2 | 1.87 | 0.56 |
| 12:AN:104:ILE:HG21 | 12:AN:107:ILE:HD11 | 1.86 | 0.56 |
| 21:AX:15:VAL:HG11 | 21:AX:26:ILE:HD13 | 1.87 | 0.56 |
| 27:BB:31:C:OP1 | 31:BF:130:ARG:NE | 2.36 | 0.56 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 32:BG:26:ALA:HB1 | 32:BG:80:VAL:HG21 | 1.85 | 0.56 |
| 46:BU:54:LYS:HB2 | 46:BU:64:GLU:HG3 | 1.87 | 0.56 |
| 55:Bd:11:ARG:HB3 | 55:Bd:50:VAL:HG13 | 1.86 | 0.56 |
| 1:AA:781:U:H2' | 1:AA:782:G:C8 | 2.40 | 0.56 |
| 1:AA:1297:A:N3 | 8:AI:68:THR:OG1 | 2.37 | 0.56 |
| 1:AA:1323:U:H5' | 8:AI:133:ASP:OD2 | 2.05 | 0.56 |
| 2:AB:134:ARG:HB2 | 2:AB:146:LEU:HD13 | 1.85 | 0.56 |
| 3:AC:42:THR:HG21 | 3:AC:75:LEU:HD22 | 1.86 | 0.56 |
| 7:AG:55:ILE:HD12 | 7:AG:72:ALA:HB2 | 1.85 | 0.56 |
| 8:AI:88:LYS:HG3 | 8:AI:89:THR:HG23 | 1.85 | 0.56 |
| 24:AM:42:THR:HG21 | 24:AM:67:ARG:HB3 | 1.87 | 0.56 |
| 26:BA:467:C:OP1 | 54:Bc:51:GLN:NE2 | 2.38 | 0.56 |
| 26:BA:867:A:C8 | 28:BC:190:VAL:HG21 | 2.40 | 0.56 |
| 26:BA:1753:C:H1' | 53:Bb:8:LYS:HD3 | 1.87 | 0.56 |
| 26:BA:2783:U:OP2 | 29:BD:28:ARG:NH2 | 2.38 | 0.56 |
| 30:BE:155:VAL:HG11 | 30:BE:209:ALA:HB2 | 1.87 | 0.56 |
| 33:BH:88:SER:HB3 | 33:BH:97:VAL:HG11 | 1.87 | 0.56 |
| 47:BV:20:LEU:HD11 | 47:BV:28:THR:HB | 1.86 | 0.56 |
| 52:Ba:76:GLN:O | 52:Ba:86:LEU:HD21 | 2.05 | 0.56 |
| 55:Bd:46:ARG:HD2 | 55:Bd:47:SER:O | 2.06 | 0.56 |
| 1:AA:655:G:H2' | 1:AA:656:A:C8 | 2.40 | 0.56 |
| 1:AA:884:A:O3' | 8:AI:130:ARG:NH2 | 2.37 | 0.56 |
| 1:AA:1410:U:H2' | 1:AA:1411:G:H8 | 1.70 | 0.56 |
| 9:AJ:23:SER:O | 9:AJ:65:LEU:N | 2.38 | 0.56 |
| 18:AU:86:GLU:OE2 | 18:AU:95:SER:OG | 2.18 | 0.56 |
| 30:BE:28:ARG:NH2 | 30:BE:30:ASP:OD2 | 2.30 | 0.56 |
| 42:BQ:35:THR:HG23 | 42:BQ:55:ILE:HD12 | 1.86 | 0.56 |
| 58:AZ:35:ARG:NH2 | 58:AZ:39:CYS:O | 2.31 | 0.56 |
| 1:AA:596:G:H5' | 16:AQ:1:MET:HB3 | 1.86 | 0.56 |
| 1:AA:608:G:O2' | 16:AQ:108:ASP:OD1 | 2.20 | 0.56 |
| 1:AA:1306:C:OP2 | 15:AR:10:ARG:NH1 | 2.39 | 0.56 |
| 3:AC:46:ILE:HG23 | 3:AC:80:ILE:HG21 | 1.87 | 0.56 |
| 7:AG:132:VAL:HG11 | 7:AG:195:VAL:HA | 1.86 | 0.56 |
| 16:AQ:75:LEU:HD23 | 16:AQ:80:VAL:HG23 | 1.87 | 0.56 |
| 19:AV:134:GLU:HA | 19:AV:137:LEU:HD12 | 1.87 | 0.56 |
| 26:BA:563:G:O6 | 26:BA:583:G:O2' | 2.18 | 0.56 |
| 26:BA:1400:U:O4 | 55:Bd:2:SER:OG | 2.23 | 0.56 |
| 1:AA:1404:C:H2' | 1:AA:1405:G:O4' | 2.06 | 0.56 |
| 2:AB:189:PRO:HG3 | 58:AZ:24:PHE:HB3 | 1.87 | 0.56 |
| 8:AI:106:PRO:HG3 | 8:AI:134:THR:HG21 | 1.88 | 0.56 |
| 25:AT:56:ARG:HD2 | 25:AT:60:ARG:HH11 | 1.70 | 0.56 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 26:BA:2481:G:O6 | 26:BA:2487:A:O2' | 2.24 | 0.56 |
| 27:BB:116:G:H2' | 27:BB:117:C:C6 | 2.41 | 0.56 |
| 1:AA:513:A:N3 | 1:AA:828:C:O2' | 2.34 | 0.56 |
| 1:AA:939:A:N6 | 1:AA:984:G:O4' | 2.39 | 0.56 |
| 24:AM:16:LEU:HD13 | 24:AM:94:ILE:HG21 | 1.86 | 0.56 |
| 26:BA:2695:C:H4' | 47:BV:17:THR:HG22 | 1.88 | 0.56 |
| 26:BA:2822:U:OP2 | 29:BD:112:ARG:NH1 | 2.34 | 0.56 |
| 36:BK:73:VAL:HA | 36:BK:95:MET:HG2 | 1.88 | 0.56 |
| 1:AA:894:C:OP1 | 14:AP:149:THR:OG1 | 2.16 | 0.56 |
| 4:AD:126:ILE:HA | 4:AD:181:ARG:HD3 | 1.88 | 0.56 |
| 6:AF:38:PRO:HD2 | 6:AF:41:ILE:HB | 1.87 | 0.56 |
| 11:AL:40:GLU:O | 11:AL:43:MET:HG2 | 2.05 | 0.56 |
| 26:BA:1833:U:OP1 | 28:BC:232:ARG:HD3 | 2.05 | 0.56 |
| 26:BA:2680:C:H4' | 32:BG:116:LYS:HD2 | 1.87 | 0.56 |
| 27:BB:4:U:H3 | 27:BB:124:G:H22 | 1.54 | 0.56 |
| 1:AA:872:G:N2 | 1:AA:1470:U:OP1 | 2.39 | 0.56 |
| 1:AA:1201:G:H2' | 1:AA:1202:G:O4' | 2.06 | 0.56 |
| 1:AA:1436:U:O2 | 1:AA:1437:A:N6 | 2.32 | 0.56 |
| 4:AD:107:VAL:HG21 | 4:AD:142:VAL:HG13 | 1.88 | 0.56 |
| 6:AF:13:TRP:NE1 | 6:AF:36:SER:O | 2.39 | 0.56 |
| 10:AK:54:ARG:NE | 10:AK:114:GLY:O | 2.37 | 0.56 |
| 29:BD:169:ALA:HB3 | 29:BD:175:LYS:CG | 2.36 | 0.56 |
| 32:BG:46:VAL:HG12 | 32:BG:51:VAL:HG22 | 1.87 | 0.56 |
| 35:BJ:54:ARG:O | 35:BJ:58:GLU:HG2 | 2.06 | 0.56 |
| 51:BZ:44:THR:HG21 | 51:BZ:83:VAL:HG21 | 1.87 | 0.56 |
| 1:AA:473:A:O2' | 1:AA:476:C:N4 | 2.39 | 0.56 |
| 1:AA:768:U:H4' | 9:AJ:20:ILE:HD11 | 1.88 | 0.56 |
| 4:AD:91:TYR:CE2 | 4:AD:95:ILE:HD11 | 2.41 | 0.56 |
| 5:AE:143:ARG:HG3 | 20:AW:61:MET:HE3 | 1.88 | 0.56 |
| 6:AF:66:LEU:HB2 | 6:AF:86:THR:HG23 | 1.87 | 0.56 |
| 7:AG:115:CYS:HB3 | 7:AG:123:GLY:H | 1.70 | 0.56 |
| 19:AV:67:PRO:HB3 | 19:AV:114:VAL:HG11 | 1.87 | 0.56 |
| 1:AA:866:U:O2' | 7:AG:61:MET:O | 2.24 | 0.55 |
| 4:AD:51:THR:HA | 4:AD:65:LYS:HA | 1.88 | 0.55 |
| 12:AN:99:ARG:HD3 | 22:AY:45:ARG:HH22 | 1.71 | 0.55 |
| 18:AU:50:PHE:HB3 | 18:AU:54:GLN:CG | 2.35 | 0.55 |
| 26:BA:33:C:O2' | 26:BA:667:A:N1 | 2.34 | 0.55 |
| 26:BA:299:A:N1 | 26:BA:323:A:O2' | 2.39 | 0.55 |
| 26:BA:426:A:OP2 | 38:BM:169:ARG:NH2 | 2.34 | 0.55 |
| 26:BA:480:A:O2' | 46:BU:86:LYS:HA | 2.05 | 0.55 |
| 26:BA:1790:C:N3 | 26:BA:2726:C:O2' | 2.34 | 0.55 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 46:BU:77:ILE:HD11 | 46:BU:101:VAL:HG11 | 1.86 | 0.55 |
| 54:Bc:15:THR:HG23 | 54:Bc:16:HIS:ND1 | 2.21 | 0.55 |
| 1:AA:2:U:H5' | 1:AA:280:A:O4' | 2.06 | 0.55 |
| 1:AA:1442:G:OP1 | 1:AA:1445:A:H4' | 2.06 | 0.55 |
| 2:AB:140:LEU:HB3 | 2:AB:142:THR:HG23 | 1.87 | 0.55 |
| 2:AB:195:ASP:N | 2:AB:198:ASN:OD1 | 2.39 | 0.55 |
| 7:AG:136:ALA:H | 7:AG:187:ALA:HB2 | 1.71 | 0.55 |
| 9:AJ:40:VAL:O | 9:AJ:44:LEU:HG | 2.07 | 0.55 |
| 24:AM:51:LYS:N | 24:AM:60:THR:O | 2.29 | 0.55 |
| 26:BA:1752:U:O2' | 26:BA:1754:G:N7 | 2.39 | 0.55 |
| 38:BM:68:VAL:HG11 | 38:BM:98:ILE:HG23 | 1.88 | 0.55 |
| 18:AU:51:THR:O | 18:AU:55:LYS:HG2 | 2.06 | 0.55 |
| 26:BA:547:G:OP1 | 52:Ba:103:SER:HB3 | 2.06 | 0.55 |
| 52:Ba:73:GLN:HA | 52:Ba:78:ARG:HH21 | 1.70 | 0.55 |
| 1:AA:114:U:O2' | 6:AF:138:GLY:O | 2.18 | 0.55 |
| 1:AA:161:C:H1' | 10:AK:120:ASN:HB3 | 1.87 | 0.55 |
| 1:AA:1187:U:C5 | 8:AI:154:VAL:HG11 | 2.41 | 0.55 |
| 2:AB:186:ILE:HG23 | 7:AG:34:PRO:HD3 | 1.89 | 0.55 |
| 11:AL:46:ILE:HD12 | 11:AL:80:ARG:HB2 | 1.89 | 0.55 |
| 26:BA:636:A:N1 | 26:BA:894:G:O2' | 2.36 | 0.55 |
| 30:BE:143:VAL:HA | 30:BE:242:GLU:HG2 | 1.88 | 0.55 |
| 1:AA:74:A:H2' | 1:AA:75:A:H8 | 1.70 | 0.55 |
| 8:AI:48:LYS:O | 8:AI:51:ILE:HG22 | 2.06 | 0.55 |
| 18:AU:38:LEU:HD23 | 18:AU:42:GLU:HG2 | 1.89 | 0.55 |
| 20:AW:36:ARG:HH21 | 20:AW:90:PRO:HG2 | 1.72 | 0.55 |
| 25:AT:31:ASN:ND2 | 25:AT:55:THR:OG1 | 2.31 | 0.55 |
| 48:BW:43:ASN:ND2 | 48:BW:46:ARG:HB2 | 2.22 | 0.55 |
| 58:AZ:7:GLU:O | 58:AZ:15:ARG:HA | 2.06 | 0.55 |
| 3:AC:23:GLU:HG3 | 3:AC:24:GLN:HG3 | 1.87 | 0.55 |
| 7:AG:26:ASP:O | 7:AG:30:LYS:HG2 | 2.07 | 0.55 |
| 7:AG:50:ASP:HB2 | 7:AG:74:VAL:HG22 | 1.87 | 0.55 |
| 1:AA:638:U:H2' | 1:AA:639:C:H5' | 1.87 | 0.55 |
| 1:AA:681:C:H2' | 1:AA:682:A:C8 | 2.42 | 0.55 |
| 1:AA:1098:G:OP1 | 1:AA:1227:A:N6 | 2.39 | 0.55 |
| 1:AA:1148:A:H4' | 1:AA:1149:G:H5'' | 1.88 | 0.55 |
| 19:AV:74:ARG:HB3 | 19:AV:92:LYS:HG2 | 1.87 | 0.55 |
| 26:BA:892:U:O2' | 26:BA:2081:A:N1 | 2.40 | 0.55 |
| 26:BA:1224:U:O4 | 26:BA:2045:C:O2' | 2.23 | 0.55 |
| 27:BB:3:G:H2' | 27:BB:4:U:C6 | 2.42 | 0.55 |
| 29:BD:163:MET:HE1 | 29:BD:308:LEU:HD21 | 1.88 | 0.55 |
| 34:BI:29:SER:HB2 | 34:BI:31:VAL:HG12 | 1.89 | 0.55 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 37:BL:71:GLU:O | 37:BL:75:LEU:HG | 2.06 | 0.55 |
| 44:BS:126:MET:HB2 | 44:BS:136:LYS:HD2 | 1.88 | 0.55 |
| 45:BT:10:THR:H | 45:BT:13:ALA:HB3 | 1.71 | 0.55 |
| 47:BV:13:LEU:HD12 | 47:BV:18:GLY:HA3 | 1.88 | 0.55 |
| 50:BY:54:ILE:O | 50:BY:57:THR:OG1 | 2.19 | 0.55 |
| 51:BZ:4:ASP:O | 51:BZ:79:ALA:N | 2.37 | 0.55 |
| 1:AA:51:A:OP2 | 1:AA:334:C:N4 | 2.39 | 0.55 |
| 1:AA:300:G:H2' | 1:AA:301:A:C8 | 2.42 | 0.55 |
| 1:AA:633:U:O2' | 1:AA:635:A:N7 | 2.29 | 0.55 |
| 1:AA:1014:C:H4' | 1:AA:1040:A:H61 | 1.72 | 0.55 |
| 1:AA:1126:A:H5'' | 11:AL:108:LEU:CD1 | 2.37 | 0.55 |
| 5:AE:54:GLU:O | 5:AE:58:LEU:HG | 2.07 | 0.55 |
| 7:AG:113:ARG:HG3 | 7:AG:125:PRO:HB2 | 1.89 | 0.55 |
| 11:AL:57:VAL:O | 11:AL:61:LEU:HG | 2.07 | 0.55 |
| 16:AQ:19:ARG:NH2 | 16:AQ:23:PRO:HA | 2.21 | 0.55 |
| 26:BA:621:A:O2' | 26:BA:623:G:OP2 | 2.20 | 0.55 |
| 26:BA:1113:U:OP1 | 26:BA:1227:C:O2' | 2.25 | 0.55 |
| 26:BA:1401:U:O4 | 45:BT:56:MET:HE3 | 2.06 | 0.55 |
| 44:BS:8:ILE:HD12 | 44:BS:113:HIS:ND1 | 2.22 | 0.55 |
| 52:Ba:121:PRO:HG3 | 52:Ba:149:ILE:CG1 | 2.36 | 0.55 |
| 1:AA:592:C:N4 | 1:AA:693:G:O2' | 2.31 | 0.55 |
| 1:AA:995:U:H2' | 1:AA:996:G:H8 | 1.71 | 0.55 |
| 1:AA:1207:U:H5' | 1:AA:1231:C:H5' | 1.88 | 0.55 |
| 9:AJ:30:ALA:N | 9:AJ:59:GLY:O | 2.27 | 0.55 |
| 14:AP:66:ILE:HG22 | 14:AP:68:GLY:H | 1.72 | 0.55 |
| 14:AP:74:GLU:O | 14:AP:78:LEU:HG | 2.07 | 0.55 |
| 28:BC:158:GLY:HA2 | 53:Bb:83:LEU:CD1 | 2.37 | 0.55 |
| 30:BE:244:ALA:O | 30:BE:248:LEU:HB2 | 2.07 | 0.55 |
| 1:AA:765:C:H2' | 1:AA:766:G:C8 | 2.41 | 0.55 |
| 1:AA:886:G:OP1 | 8:AI:63:ARG:NE | 2.39 | 0.55 |
| 5:AE:4:PRO:HG2 | 5:AE:6:LYS:HG2 | 1.89 | 0.55 |
| 7:AG:22:VAL:HG13 | 7:AG:27:GLU:HG3 | 1.89 | 0.55 |
| 9:AJ:52:PHE:HB2 | 9:AJ:61:TYR:HE1 | 1.72 | 0.55 |
| 15:AR:33:CYS:SG | 15:AR:36:CYS:HB3 | 2.47 | 0.55 |
| 21:AX:20:CYS:CB | 21:AX:39:CYS:SG | 2.77 | 0.55 |
| 26:BA:145:G:N2 | 26:BA:180:G:O6 | 2.40 | 0.55 |
| 26:BA:1827:A:O3' | 28:BC:170:TRP:HB2 | 2.07 | 0.55 |
| 53:Bb:7:LYS:O | 53:Bb:10:ARG:NH1 | 2.39 | 0.55 |
| 56:Be:3:ARG:NE | 56:Be:8:GLU:OE2 | 2.40 | 0.55 |
| 1:AA:901:U:H2' | 1:AA:902:U:O4' | 2.08 | 0.54 |
| 1:AA:937:C:H2' | 1:AA:1159:A:N1 | 2.22 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:1235:A:H5' | 19:AV:90:LYS:HD2 | 1.88 | 0.54 |
| 4:AD:22:GLU:OE2 | 4:AD:29:ARG:NE | 2.40 | 0.54 |
| 8:AI:77:ALA:HA | 8:AI:155:ALA:HB2 | 1.89 | 0.54 |
| 14:AP:98:LYS:CE | 18:AU:37:LEU:HA | 2.37 | 0.54 |
| 26:BA:62:C:O2 | 26:BA:66:A:O2' | 2.24 | 0.54 |
| 26:BA:2364:C:O2' | 26:BA:2365:A:O4' | 2.25 | 0.54 |
| 29:BD:6:ARG:HD3 | 36:BK:41:LYS:HB3 | 1.89 | 0.54 |
| 48:BW:9:ILE:O | 48:BW:17:ARG:NH1 | 2.40 | 0.54 |
| 1:AA:135:G:H2' | 1:AA:137:A:OP2 | 2.07 | 0.54 |
| 1:AA:599:C:H42 | 1:AA:688:G:H1 | 1.55 | 0.54 |
| 1:AA:1001:U:H2' | 1:AA:1002:C:C6 | 2.42 | 0.54 |
| 1:AA:1196:C:H5 | 1:AA:1234:A:H5'' | 1.73 | 0.54 |
| 2:AB:211:ASN:O | 2:AB:217:LEU:HD11 | 2.08 | 0.54 |
| 5:AE:81:ILE:O | 5:AE:85:LEU:HG | 2.07 | 0.54 |
| 6:AF:61:SER:HA | 6:AF:73:LYS:HG3 | 1.89 | 0.54 |
| 8:AI:62:MET:HA | 8:AI:67:ASN:HB3 | 1.88 | 0.54 |
| 16:AQ:56:ASP:HB3 | 21:AX:34:ILE:HG12 | 1.88 | 0.54 |
| 26:BA:562:G:OP1 | 49:BX:122:ARG:NH2 | 2.40 | 0.54 |
| 44:BS:38:LYS:HE2 | 44:BS:114:ALA:H | 1.72 | 0.54 |
| 1:AA:546:G:H1' | 1:AA:573:U:O4 | 2.07 | 0.54 |
| 1:AA:680:C:O2' | 16:AQ:101:HIS:ND1 | 2.39 | 0.54 |
| 1:AA:688:G:O2' | 16:AQ:55:ARG:NH1 | 2.41 | 0.54 |
| 1:AA:1265:A:OP1 | 18:AU:48:ARG:NE | 2.39 | 0.54 |
| 3:AC:143:VAL:HG22 | 3:AC:145:LYS:HG3 | 1.89 | 0.54 |
| 4:AD:121:PRO:HB3 | 4:AD:185:ILE:HD12 | 1.88 | 0.54 |
| 6:AF:96:MET:HE3 | 6:AF:104:LEU:HD12 | 1.89 | 0.54 |
| 17:AS:4:ASP:HA | 17:AS:8:ASN:HD22 | 1.72 | 0.54 |
| 27:BB:101:C:O2 | 49:BX:49:LYS:HE2 | 2.07 | 0.54 |
| 28:BC:139:VAL:HG21 | 28:BC:149:LEU:HD12 | 1.89 | 0.54 |
| 28:BC:228:LEU:HD13 | 28:BC:234:THR:HG22 | 1.90 | 0.54 |
| 1:AA:289:G:OP2 | 1:AA:289:G:N2 | 2.30 | 0.54 |
| 1:AA:907:U:OP2 | 1:AA:1170:C:O2' | 2.16 | 0.54 |
| 1:AA:917:G:C8 | 1:AA:1312:A:H4' | 2.43 | 0.54 |
| 1:AA:933:G:H2' | 1:AA:934:G:O4' | 2.07 | 0.54 |
| 3:AC:170:VAL:HA | 3:AC:175:THR:HG22 | 1.88 | 0.54 |
| 7:AG:191:ALA:O | 7:AG:195:VAL:HG23 | 2.08 | 0.54 |
| 13:AO:58:LYS:HB2 | 13:AO:113:ASP:CB | 2.37 | 0.54 |
| 16:AQ:5:HIS:NE2 | 16:AQ:121:LYS:HG3 | 2.22 | 0.54 |
| 16:AQ:113:ARG:O | 16:AQ:117:LEU:HG | 2.08 | 0.54 |
| 21:AX:14:ARG:HG2 | 21:AX:58:LEU:HB2 | 1.90 | 0.54 |
| 26:BA:2034:A:O2' | 44:BS:135:PRO:O | 2.23 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 30:BE:168:VAL:HG11 | 30:BE:212:LEU:HD12 | 1.88 | 0.54 |
| 41:BP:7:GLN:HG2 | 41:BP:32:ILE:O | 2.08 | 0.54 |
| 43:BR:52:MET:HG2 | 43:BR:53:PRO:O | 2.08 | 0.54 |
| 1:AA:485:C:OP1 | 5:AE:38:LYS:NZ | 2.30 | 0.54 |
| 1:AA:520:C:H2' | 1:AA:521:G:O4' | 2.08 | 0.54 |
| 1:AA:1118:A:OP1 | 25:AT:5:ARG:N | 2.40 | 0.54 |
| 3:AC:135:LYS:NZ | 3:AC:173:LEU:O | 2.38 | 0.54 |
| 24:AM:2:GLN:HG3 | 24:AM:101:GLU:O | 2.07 | 0.54 |
| 26:BA:970:G:N3 | 26:BA:982:U:N3 | 2.55 | 0.54 |
| 26:BA:2016:C:O2 | 36:BK:13:ASN:ND2 | 2.41 | 0.54 |
| 30:BE:48:GLY:HA2 | 30:BE:95:PRO:HG2 | 1.90 | 0.54 |
| 30:BE:203:THR:OG1 | 30:BE:205:ILE:HG22 | 2.06 | 0.54 |
| 38:BM:121:VAL:HG21 | 38:BM:131:GLU:CG | 2.38 | 0.54 |
| 39:BN:66:GLU:O | 39:BN:69:LYS:HG2 | 2.08 | 0.54 |
| 40:BO:58:ILE:HD11 | 40:BO:70:ILE:HD11 | 1.88 | 0.54 |
| 41:BP:116:ASP:N | 41:BP:116:ASP:OD1 | 2.40 | 0.54 |
| 1:AA:13:C:H2' | 1:AA:14:C:O4' | 2.08 | 0.54 |
| 1:AA:192:G:H2' | 1:AA:193:G:C8 | 2.43 | 0.54 |
| 1:AA:525:C:H5'' | 13:AO:6:TYR:O | 2.07 | 0.54 |
| 1:AA:573:U:O2 | 1:AA:574:U:H3' | 2.08 | 0.54 |
| 1:AA:1126:A:H5'' | 11:AL:108:LEU:HD12 | 1.88 | 0.54 |
| 1:AA:1195:A:H2' | 1:AA:1196:C:C4' | 2.38 | 0.54 |
| 4:AD:16:LYS:HG3 | 4:AD:36:MET:HE1 | 1.88 | 0.54 |
| 4:AD:100:THR:HG23 | 4:AD:125:THR:H | 1.72 | 0.54 |
| 7:AG:50:ASP:HB2 | 7:AG:74:VAL:CG2 | 2.37 | 0.54 |
| 8:AI:84:ILE:O | 8:AI:88:LYS:HG2 | 2.08 | 0.54 |
| 33:BH:55:ILE:HG21 | 33:BH:60:ILE:HB | 1.90 | 0.54 |
| 40:BO:35:ILE:O | 40:BO:39:ILE:HG12 | 2.08 | 0.54 |
| 41:BP:23:TRP:HB2 | 41:BP:53:LYS:HG2 | 1.88 | 0.54 |
| 42:BQ:8:ILE:HD11 | 42:BQ:25:LYS:HE3 | 1.88 | 0.54 |
| 1:AA:115:U:H5' | 6:AF:139:THR:HA | 1.90 | 0.54 |
| 1:AA:337:C:O2' | 1:AA:370:G:N3 | 2.40 | 0.54 |
| 1:AA:424:U:H2' | 1:AA:425:A:O4' | 2.08 | 0.54 |
| 1:AA:567:U:H2' | 1:AA:568:A:C8 | 2.42 | 0.54 |
| 1:AA:1181:C:H1' | 1:AA:1311:U:O2 | 2.07 | 0.54 |
| 1:AA:1263:G:N7 | 18:AU:48:ARG:NH1 | 2.55 | 0.54 |
| 2:AB:209:THR:OG1 | 2:AB:220:VAL:HG21 | 2.07 | 0.54 |
| 13:AO:46:ARG:HD3 | 13:AO:97:GLU:OE2 | 2.08 | 0.54 |
| 16:AQ:4:MET:HE2 | 16:AQ:124:ARG:HD3 | 1.90 | 0.54 |
| 25:AT:8:ASN:OD1 | 25:AT:11:ARG:NH1 | 2.31 | 0.54 |
| 28:BC:171:LEU:HD22 | 53:Bb:30:VAL:HG21 | 1.90 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 58:AZ:46:TYR:CZ | 58:AZ:55:ALA:HB3 | 2.43 | 0.54 |
| 1:AA:484:C:H5'' | 5:AE:23:MET:HE1 | 1.89 | 0.54 |
| 1:AA:533:G:H2' | 1:AA:534:U:C6 | 2.43 | 0.54 |
| 1:AA:553:G:H5'' | 5:AE:50:MET:SD | 2.47 | 0.54 |
| 1:AA:840:G:H2' | 1:AA:841:C:C6 | 2.43 | 0.54 |
| 1:AA:1197:A:H2' | 1:AA:1198:A:C8 | 2.43 | 0.54 |
| 2:AB:61:SER:OG | 2:AB:64:GLU:HG2 | 2.08 | 0.54 |
| 2:AB:164:GLU:HB3 | 7:AG:12:ARG:HD3 | 1.88 | 0.54 |
| 4:AD:103:ILE:CG1 | 4:AD:134:MET:HG2 | 2.38 | 0.54 |
| 7:AG:132:VAL:HG21 | 7:AG:195:VAL:HG22 | 1.90 | 0.54 |
| 8:AI:92:ASN:O | 8:AI:96:VAL:HG23 | 2.08 | 0.54 |
| 10:AK:42:LYS:HG3 | 10:AK:44:VAL:HG13 | 1.88 | 0.54 |
| 16:AQ:46:SER:O | 16:AQ:50:ILE:HG12 | 2.08 | 0.54 |
| 18:AU:76:ILE:HB | 18:AU:104:GLY:H | 1.73 | 0.54 |
| 23:AH:56:CYS:SG | 23:AH:107:GLN:HB3 | 2.47 | 0.54 |
| 24:AM:4:ALA:N | 24:AM:75:ILE:O | 2.28 | 0.54 |
| 26:BA:785:U:H4' | 41:BP:84:LYS:HE2 | 1.89 | 0.54 |
| 34:BI:130:MET:HA | 34:BI:130:MET:HE2 | 1.89 | 0.54 |
| 35:BJ:60:GLY:HA3 | 35:BJ:66:PRO:HD2 | 1.89 | 0.54 |
| 46:BU:2:ILE:HG21 | 46:BU:12:GLN:HG2 | 1.89 | 0.54 |
| 1:AA:416:G:H2' | 1:AA:417:U:C6 | 2.43 | 0.54 |
| 13:AO:105:GLY:HA3 | 13:AO:111:MET:HB2 | 1.90 | 0.54 |
| 16:AQ:149:MET:HE1 | 50:BY:74:VAL:HG13 | 1.89 | 0.54 |
| 17:AS:1:MET:HE1 | 17:AS:27:LEU:O | 2.08 | 0.54 |
| 26:BA:462:G:OP2 | 55:Bd:46:ARG:HG3 | 2.08 | 0.54 |
| 26:BA:649:G:O2' | 30:BE:31:LEU:HD21 | 2.08 | 0.54 |
| 39:BN:103:LEU:CB | 39:BN:130:ILE:HD11 | 2.37 | 0.54 |
| 39:BN:139:PRO:HG2 | 39:BN:142:ARG:HB2 | 1.90 | 0.54 |
| 46:BU:51:ASP:HB2 | 46:BU:105:LYS:O | 2.08 | 0.54 |
| 46:BU:56:MET:SD | 46:BU:102:MET:HB2 | 2.47 | 0.54 |
| 1:AA:632:G:O2' | 1:AA:738:C:H4' | 2.09 | 0.54 |
| 1:AA:913:C:H3' | 1:AA:914:A:H2' | 1.90 | 0.54 |
| 1:AA:1296:A:H62 | 1:AA:1320:G:H21 | 1.56 | 0.54 |
| 7:AG:110:TYR:O | 7:AG:196:ASN:ND2 | 2.40 | 0.54 |
| 13:AO:4:GLY:O | 17:AS:62:ARG:HD3 | 2.08 | 0.54 |
| 13:AO:24:TYR:HE1 | 13:AO:107:MET:HE1 | 1.73 | 0.54 |
| 14:AP:137:ARG:HB3 | 14:AP:139:LEU:HD13 | 1.90 | 0.54 |
| 24:AM:9:SER:CB | 24:AM:70:LYS:HG2 | 2.37 | 0.54 |
| 26:BA:463:G:N2 | 26:BA:466:A:OP2 | 2.30 | 0.54 |
| 26:BA:653:A:OP1 | 40:BO:94:SER:OG | 2.17 | 0.54 |
| 26:BA:861:U:O3' | 28:BC:1:MET:HE2 | 2.08 | 0.54 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 26:BA:1296:G:HO2' | 52:Ba:131:THR:HG1 | 1.56 | 0.54 |
| 28:BC:33:VAL:HG21 | 28:BC:83:CYS:SG | 2.48 | 0.54 |
| 49:BX:132:VAL:HG23 | 49:BX:138:LEU:O | 2.08 | 0.54 |
| 1:AA:133:G:H4' | 23:AH:18:GLN:OE1 | 2.08 | 0.53 |
| 1:AA:493:G:OP1 | 13:AO:104:GLY:HA2 | 2.08 | 0.53 |
| 2:AB:164:GLU:OE1 | 7:AG:12:ARG:N | 2.42 | 0.53 |
| 15:AR:7:LYS:HD2 | 15:AR:31:TYR:OH | 2.07 | 0.53 |
| 16:AQ:19:ARG:HH21 | 16:AQ:23:PRO:HA | 1.72 | 0.53 |
| 23:AH:9:SER:HA | 23:AH:16:ALA:HA | 1.90 | 0.53 |
| 29:BD:124:GLU:HA | 29:BD:127:GLU:HG2 | 1.90 | 0.53 |
| 31:BF:4:PRO:O | 31:BF:124:LEU:HD12 | 2.08 | 0.53 |
| 46:BU:53:VAL:HG12 | 46:BU:103:ILE:HA | 1.89 | 0.53 |
| 1:AA:605:G:O2' | 1:AA:607:G:OP2 | 2.18 | 0.53 |
| 1:AA:1201:G:OP2 | 15:AR:28:TYR:OH | 2.18 | 0.53 |
| 1:AA:1202:G:O2' | 1:AA:1205:C:O2' | 2.15 | 0.53 |
| 2:AB:168:ILE:HG12 | 2:AB:188:VAL:CG1 | 2.38 | 0.53 |
| 3:AC:48:SER:OG | 3:AC:54:VAL:HG21 | 2.08 | 0.53 |
| 4:AD:153:ASP:OD1 | 4:AD:156:THR:OG1 | 2.14 | 0.53 |
| 6:AF:220:ILE:HD12 | 6:AF:229:LEU:CD1 | 2.38 | 0.53 |
| 13:AO:124:VAL:O | 13:AO:127:VAL:HG12 | 2.09 | 0.53 |
| 20:AW:2:ASP:HB3 | 20:AW:24:LYS:HB2 | 1.90 | 0.53 |
| 20:AW:51:ILE:O | 20:AW:69:LYS:N | 2.30 | 0.53 |
| 27:BB:26:C:H2' | 27:BB:27:A:H8 | 1.72 | 0.53 |
| 40:BO:110:GLU:O | 40:BO:114:GLU:HG2 | 2.08 | 0.53 |
| 50:BY:39:MET:HE1 | 50:BY:94:ILE:HG13 | 1.90 | 0.53 |
| 57:Bf:40:LYS:O | 57:Bf:44:GLU:HG2 | 2.09 | 0.53 |
| 1:AA:427:A:H2' | 1:AA:428:U:C6 | 2.43 | 0.53 |
| 1:AA:528:G:H2' | 1:AA:529:U:C6 | 2.42 | 0.53 |
| 1:AA:698:U:H2' | 1:AA:699:G:O4' | 2.07 | 0.53 |
| 4:AD:22:GLU:HB2 | 4:AD:77:ILE:CG2 | 2.38 | 0.53 |
| 4:AD:84:GLY:HA2 | 4:AD:158:MET:HE1 | 1.91 | 0.53 |
| 6:AF:108:LYS:O | 6:AF:231:GLY:HA3 | 2.09 | 0.53 |
| 11:AL:20:VAL:HG22 | 11:AL:63:ILE:HG23 | 1.91 | 0.53 |
| 19:AV:26:ILE:CD1 | 19:AV:104:GLN:HB3 | 2.38 | 0.53 |
| 26:BA:1109:G:N2 | 26:BA:1110:U:O4 | 2.42 | 0.53 |
| 26:BA:2242:A:OP2 | 38:BM:91:LYS:NZ | 2.41 | 0.53 |
| 26:BA:2519:G:HO2' | 26:BA:2565:U:HO2' | 1.53 | 0.53 |
| 36:BK:5:ARG:O | 36:BK:83:PRO:HG3 | 2.08 | 0.53 |
| 50:BY:41:VAL:HG22 | 50:BY:62:LEU:HD23 | 1.90 | 0.53 |
| 1:AA:106:A:OP2 | 17:AS:92:ARG:HD3 | 2.08 | 0.53 |
| 1:AA:158:A:N1 | 1:AA:186:U:O2' | 2.38 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:519:C:H2' | 1:AA:520:C:C6 | 2.44 | 0.53 |
| 1:AA:1182:U:H2' | 1:AA:1183:A:C8 | 2.43 | 0.53 |
| 1:AA:1329:C:C4 | 22:AY:27:HIS:HB3 | 2.43 | 0.53 |
| 2:AB:143:ARG:HH11 | 2:AB:163:PHE:HD1 | 1.57 | 0.53 |
| 7:AG:101:ILE:HG22 | 7:AG:105:LYS:HZ2 | 1.73 | 0.53 |
| 8:AI:128:PRO:HD3 | 22:AY:61:LEU:HD21 | 1.89 | 0.53 |
| 25:AT:20:TYR:HB3 | 25:AT:23:VAL:CG2 | 2.37 | 0.53 |
| 26:BA:157:U:O4 | 26:BA:168:C:O2' | 2.18 | 0.53 |
| 26:BA:895:U:OP2 | 37:BL:10:ARG:NH1 | 2.41 | 0.53 |
| 26:BA:1107:A:O3' | 35:BJ:128:ARG:NH1 | 2.37 | 0.53 |
| 26:BA:2587:G:O2' | 26:BA:2590:C:OP2 | 2.24 | 0.53 |
| 29:BD:29:ALA:O | 29:BD:312:ARG:NH1 | 2.41 | 0.53 |
| 44:BS:48:PHE:HZ | 44:BS:59:VAL:HG12 | 1.74 | 0.53 |
| 1:AA:233:G:H1' | 10:AK:8:ARG:HE | 1.74 | 0.53 |
| 1:AA:1123:A:H4' | 1:AA:1124:G:N7 | 2.24 | 0.53 |
| 1:AA:1215:A:H4' | 19:AV:39:VAL:HG22 | 1.89 | 0.53 |
| 1:AA:1322:A:H5' | 8:AI:56:ARG:NE | 2.24 | 0.53 |
| 2:AB:170:VAL:HG13 | 2:AB:176:ASP:HB2 | 1.90 | 0.53 |
| 23:AH:72:ARG:HG2 | 23:AH:97:ARG:CD | 2.39 | 0.53 |
| 26:BA:2254:U:O2' | 26:BA:2447:G:OP2 | 2.22 | 0.53 |
| 27:BB:30:U:H2' | 27:BB:31:C:C6 | 2.44 | 0.53 |
| 29:BD:12:LEU:HD11 | 29:BD:249:ARG:HB3 | 1.90 | 0.53 |
| 34:BI:77:ASP:HB3 | 34:BI:145:ALA:HB1 | 1.91 | 0.53 |
| 36:BK:120:ARG:O | 36:BK:122:PRO:HD3 | 2.09 | 0.53 |
| 38:BM:61:ILE:HD11 | 38:BM:148:LEU:HD11 | 1.91 | 0.53 |
| 52:Ba:128:ILE:HG21 | 52:Ba:137:LYS:HG3 | 1.89 | 0.53 |
| 1:AA:173:A:O3' | 17:AS:1:MET:N | 2.42 | 0.53 |
| 1:AA:237:U:OP1 | 17:AS:98:LYS:HE2 | 2.08 | 0.53 |
| 1:AA:970:G:H2' | 1:AA:971:C:O4' | 2.09 | 0.53 |
| 1:AA:1109:C:H2' | 1:AA:1110:G:C8 | 2.44 | 0.53 |
| 1:AA:1290:C:OP1 | 11:AL:131:LYS:HD3 | 2.09 | 0.53 |
| 4:AD:87:LEU:HD12 | 4:AD:91:TYR:HD2 | 1.73 | 0.53 |
| 5:AE:102:SER:HB3 | 7:AG:148:ARG:NE | 2.23 | 0.53 |
| 7:AG:69:LYS:HE3 | 7:AG:91:VAL:HG12 | 1.90 | 0.53 |
| 7:AG:126:HIS:HA | 7:AG:168:LYS:NZ | 2.24 | 0.53 |
| 8:AI:137:ARG:O | 8:AI:141:MET:HG2 | 2.07 | 0.53 |
| 18:AU:68:ARG:HA | 18:AU:86:GLU:HB2 | 1.90 | 0.53 |
| 29:BD:121:ASN:HA | 29:BD:124:GLU:OE2 | 2.09 | 0.53 |
| 29:BD:201:THR:HG22 | 29:BD:311:LEU:HD23 | 1.89 | 0.53 |
| 58:AZ:47:THR:HG23 | 58:AZ:54:LYS:HG2 | 1.91 | 0.53 |
| 1:AA:283:G:N2 | 1:AA:497:G:OP1 | 2.37 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:390:C:OP1 | 5:AE:125:THR:OG1 | 2.13 | 0.53 |
| 1:AA:606:A:H1' | 1:AA:674:G:O4' | 2.09 | 0.53 |
| 1:AA:1207:U:O2' | 1:AA:1223:G:N2 | 2.42 | 0.53 |
| 1:AA:1230:U:H2' | 1:AA:1231:C:C6 | 2.44 | 0.53 |
| 2:AB:121:LEU:HD23 | 2:AB:168:ILE:HD12 | 1.90 | 0.53 |
| 6:AF:97:LEU:HD11 | 6:AF:229:LEU:O | 2.09 | 0.53 |
| 7:AG:194:GLN:HA | 7:AG:197:VAL:CG1 | 2.39 | 0.53 |
| 8:AI:57:LEU:HD13 | 8:AI:136:LEU:HD22 | 1.91 | 0.53 |
| 9:AJ:47:ILE:HD11 | 9:AJ:63:VAL:CG1 | 2.39 | 0.53 |
| 18:AU:69:THR:N | 18:AU:86:GLU:O | 2.39 | 0.53 |
| 36:BK:116:GLU:OE1 | 36:BK:116:GLU:N | 2.41 | 0.53 |
| 37:BL:69:VAL:HG22 | 37:BL:106:GLY:HA2 | 1.90 | 0.53 |
| 41:BP:115:LEU:HD21 | 41:BP:123:LEU:HD12 | 1.89 | 0.53 |
| 47:BV:6:CYS:HB2 | 47:BV:32:CYS:HB3 | 1.89 | 0.53 |
| 1:AA:188:A:N3 | 1:AA:204:C:O2' | 2.38 | 0.53 |
| 1:AA:659:A:C8 | 12:AN:115:HIS:HB3 | 2.43 | 0.53 |
| 3:AC:102:SER:HA | 3:AC:105:ARG:HG2 | 1.90 | 0.53 |
| 7:AG:54:ASP:HB3 | 7:AG:181:THR:HG21 | 1.91 | 0.53 |
| 7:AG:113:ARG:HG3 | 7:AG:125:PRO:CB | 2.39 | 0.53 |
| 13:AO:24:TYR:CE1 | 13:AO:107:MET:HE1 | 2.43 | 0.53 |
| 14:AP:23:LEU:HD23 | 31:BF:109:TYR:HE2 | 1.73 | 0.53 |
| 14:AP:148:SER:HA | 14:AP:151:ARG:HE | 1.74 | 0.53 |
| 26:BA:18:G:H5'' | 44:BS:4:ILE:HD12 | 1.90 | 0.53 |
| 26:BA:1121:A:N3 | 26:BA:2497:G:O2' | 2.34 | 0.53 |
| 27:BB:53:C:H2' | 27:BB:54:A:O4' | 2.09 | 0.53 |
| 31:BF:9:VAL:HB | 31:BF:63:LEU:HB2 | 1.90 | 0.53 |
| 45:BT:10:THR:O | 45:BT:14:MET:HG2 | 2.09 | 0.53 |
| 48:BW:6:THR:O | 48:BW:10:ARG:HG3 | 2.08 | 0.53 |
| 1:AA:909:G:H2' | 1:AA:910:A:O4' | 2.09 | 0.53 |
| 1:AA:1069:A:O4' | 11:AL:4:VAL:HG11 | 2.08 | 0.53 |
| 1:AA:1198:A:H2' | 1:AA:1199:U:C6 | 2.44 | 0.53 |
| 4:AD:132:SER:HA | 4:AD:135:GLN:OE1 | 2.08 | 0.53 |
| 6:AF:43:ILE:HD12 | 6:AF:47:LEU:HD12 | 1.90 | 0.53 |
| 7:AG:158:ALA:HB2 | 7:AG:184:PHE:CZ | 2.43 | 0.53 |
| 8:AI:48:LYS:CE | 8:AI:129:GLN:HG3 | 2.39 | 0.53 |
| 19:AV:139:LEU:HD13 | 19:AV:142:LYS:HE3 | 1.91 | 0.53 |
| 20:AW:2:ASP:HA | 20:AW:38:LYS:HE2 | 1.91 | 0.53 |
| 20:AW:14:LEU:HD22 | 20:AW:16:ARG:CD | 2.39 | 0.53 |
| 21:AX:15:VAL:CG1 | 21:AX:43:LEU:HD13 | 2.39 | 0.53 |
| 24:AM:46:VAL:HG22 | 24:AM:65:GLU:HG2 | 1.91 | 0.53 |
| 26:BA:1698:A:O2' | 41:BP:57:GLY:HA2 | 2.09 | 0.53 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 26:BA:1986:C:OP1 | 26:BA:1987:A:O2' | 2.23 | 0.53 |
| 33:BH:10:PRO:HD3 | 33:BH:119:LEU:HD22 | 1.91 | 0.53 |
| 53:Bb:77:THR:O | 53:Bb:81:GLN:HG2 | 2.09 | 0.53 |
| 1:AA:596:G:OP1 | 16:AQ:1:MET:N | 2.30 | 0.53 |
| 1:AA:627:U:O4 | 1:AA:644:A:O2' | 2.21 | 0.53 |
| 8:AI:85:VAL:HG13 | 8:AI:163:ILE:HD11 | 1.90 | 0.53 |
| 17:AS:58:SER:O | 17:AS:61:GLN:NE2 | 2.30 | 0.53 |
| 26:BA:1253:A:OP1 | 49:BX:71:ARG:NH2 | 2.41 | 0.53 |
| 29:BD:37:PRO:HA | 29:BD:168:SER:O | 2.08 | 0.53 |
| 30:BE:144:ASN:ND2 | 30:BE:242:GLU:HB2 | 2.24 | 0.53 |
| 30:BE:201:GLU:O | 30:BE:203:THR:HG23 | 2.08 | 0.53 |
| 31:BF:40:CYS:N | 31:BF:58:GLY:O | 2.42 | 0.53 |
| 38:BM:188:ILE:HG22 | 38:BM:193:SER:HA | 1.90 | 0.53 |
| 1:AA:1061:C:H2' | 1:AA:1062:U:O4' | 2.09 | 0.52 |
| 1:AA:1103:G:H2' | 1:AA:1127:A:H61 | 1.75 | 0.52 |
| 1:AA:1412:A:C2' | 1:AA:1413:A:H5' | 2.40 | 0.52 |
| 2:AB:173:PRO:HG3 | 2:AB:194:CYS:SG | 2.48 | 0.52 |
| 3:AC:95:MET:HE2 | 3:AC:120:VAL:CG2 | 2.39 | 0.52 |
| 6:AF:36:SER:HA | 6:AF:79:VAL:O | 2.09 | 0.52 |
| 6:AF:67:VAL:HG22 | 6:AF:85:ILE:CD1 | 2.39 | 0.52 |
| 7:AG:11:THR:O | 7:AG:15:LYS:HG2 | 2.08 | 0.52 |
| 7:AG:170:VAL:HG12 | 7:AG:172:THR:HG23 | 1.89 | 0.52 |
| 15:AR:34:ARG:NH2 | 24:AM:52:SER:O | 2.29 | 0.52 |
| 24:AM:83:ARG:HD3 | 24:AM:86:MET:HE3 | 1.90 | 0.52 |
| 26:BA:937:U:O2' | 49:BX:40:HIS:O | 2.26 | 0.52 |
| 26:BA:2694:A:O2' | 29:BD:51:HIS:ND1 | 2.32 | 0.52 |
| 41:BP:35:ALA:HB2 | 41:BP:44:LEU:HD12 | 1.91 | 0.52 |
| 46:BU:24:ARG:NH1 | 46:BU:73:MET:O | 2.42 | 0.52 |
| 1:AA:553:G:H2' | 1:AA:554:A:C8 | 2.44 | 0.52 |
| 1:AA:592:C:H41 | 1:AA:693:G:HO2' | 1.52 | 0.52 |
| 1:AA:1025:C:H2' | 1:AA:1026:C:C6 | 2.45 | 0.52 |
| 1:AA:1039:C:C4 | 1:AA:1041:A:H5' | 2.44 | 0.52 |
| 1:AA:1203:U:O2' | 1:AA:1225:U:N3 | 2.39 | 0.52 |
| 1:AA:1312:A:H2' | 1:AA:1313:G:C8 | 2.44 | 0.52 |
| 2:AB:131:HIS:O | 2:AB:135:MET:HG2 | 2.09 | 0.52 |
| 3:AC:152:LYS:HZ1 | 3:AC:157:PRO:HD2 | 1.74 | 0.52 |
| 6:AF:213:ILE:HG22 | 6:AF:216:TYR:HB2 | 1.90 | 0.52 |
| 7:AG:37:GLU:OE1 | 7:AG:38:PRO:HD2 | 2.10 | 0.52 |
| 10:AK:8:ARG:O | 17:AS:96:LYS:NZ | 2.31 | 0.52 |
| 15:AR:48:LYS:HE2 | 24:AM:62:ASP:HA | 1.90 | 0.52 |
| 17:AS:36:GLY:O | 17:AS:85:ILE:HD12 | 2.09 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 18:AU:26:LEU:O | 18:AU:78:PRO:HB3 | 2.09 | 0.52 |
| 25:AT:56:ARG:O | 25:AT:60:ARG:HG3 | 2.09 | 0.52 |
| 26:BA:1841:C:O2' | 28:BC:8:GLN:HA | 2.09 | 0.52 |
| 26:BA:1866:A:O2' | 53:Bb:24:ARG:HG2 | 2.10 | 0.52 |
| 27:BB:25:G:H4' | 27:BB:26:C:C5 | 2.44 | 0.52 |
| 1:AA:323:C:H2' | 1:AA:324:C:H6 | 1.73 | 0.52 |
| 1:AA:835:G:O2' | 1:AA:851:G:O6 | 2.16 | 0.52 |
| 22:AY:15:ALA:HB2 | 22:AY:37:VAL:HA | 1.91 | 0.52 |
| 26:BA:2575:A:OP1 | 26:BA:2660:G:O2' | 2.22 | 0.52 |
| 27:BB:1:G:H1 | 27:BB:127:C:H1' | 1.74 | 0.52 |
| 32:BG:137:ILE:CG2 | 32:BG:145:VAL:HG23 | 2.40 | 0.52 |
| 40:BO:39:ILE:HD12 | 40:BO:69:LEU:HD22 | 1.90 | 0.52 |
| 40:BO:78:GLY:O | 40:BO:79:LEU:HD23 | 2.09 | 0.52 |
| 47:BV:45:LEU:HD12 | 47:BV:48:ARG:HD3 | 1.91 | 0.52 |
| 52:Ba:29:ASP:OD2 | 52:Ba:122:SER:HB2 | 2.09 | 0.52 |
| 1:AA:86:U:H2' | 1:AA:87:G:H5' | 1.92 | 0.52 |
| 1:AA:604:A:OP1 | 4:AD:132:SER:OG | 2.14 | 0.52 |
| 1:AA:1235:A:N1 | 1:AA:1318:G:H1' | 2.24 | 0.52 |
| 7:AG:23:ALA:N | 7:AG:27:GLU:OE2 | 2.29 | 0.52 |
| 20:AW:17:ARG:CZ | 20:AW:43:LEU:HD12 | 2.38 | 0.52 |
| 26:BA:506:U:O2' | 55:Bd:32:ARG:HD2 | 2.10 | 0.52 |
| 33:BH:29:LYS:HB2 | 33:BH:101:ASP:HB3 | 1.92 | 0.52 |
| 51:BZ:27:ALA:O | 51:BZ:31:VAL:HG23 | 2.10 | 0.52 |
| 1:AA:323:C:H2' | 1:AA:324:C:C6 | 2.45 | 0.52 |
| 1:AA:460:A:OP1 | 13:AO:64:ILE:HB | 2.10 | 0.52 |
| 1:AA:1029:U:H2' | 1:AA:1030:U:C6 | 2.45 | 0.52 |
| 1:AA:1072:A:H2' | 1:AA:1073:U:C6 | 2.45 | 0.52 |
| 4:AD:32:VAL:CG1 | 4:AD:50:THR:HG21 | 2.39 | 0.52 |
| 5:AE:133:ILE:O | 5:AE:146:THR:HA | 2.09 | 0.52 |
| 7:AG:50:ASP:OD1 | 7:AG:50:ASP:N | 2.43 | 0.52 |
| 7:AG:161:VAL:CG1 | 7:AG:188:THR:HG21 | 2.40 | 0.52 |
| 14:AP:36:PRO:HB2 | 14:AP:38:GLN:OE1 | 2.09 | 0.52 |
| 14:AP:108:LEU:HB2 | 14:AP:113:ILE:CD1 | 2.39 | 0.52 |
| 26:BA:691:A:O2' | 26:BA:721:C:O2' | 2.27 | 0.52 |
| 26:BA:2712:U:OP2 | 41:BP:59:SER:HB3 | 2.10 | 0.52 |
| 27:BB:25:G:O6 | 27:BB:57:U:O2' | 2.21 | 0.52 |
| 32:BG:100:LYS:NZ | 32:BG:102:ASP:HB3 | 2.25 | 0.52 |
| 37:BL:124:SER:O | 37:BL:128:LYS:HG3 | 2.10 | 0.52 |
| 58:AZ:9:CYS:SG | 58:AZ:35:ARG:HA | 2.50 | 0.52 |
| 58:AZ:10:THR:HG21 | 58:AZ:33:ILE:HG23 | 1.92 | 0.52 |
| 1:AA:214:G:H1' | 1:AA:244:A:N1 | 2.25 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:467:G:O2' | 1:AA:475:A:N1 | 2.33 | 0.52 |
| 1:AA:1107:G:OP2 | 25:AT:56:ARG:HG3 | 2.10 | 0.52 |
| 1:AA:1464:G:H2' | 1:AA:1465:C:C6 | 2.44 | 0.52 |
| 2:AB:107:VAL:CG1 | 58:AZ:27:PRO:HB2 | 2.39 | 0.52 |
| 7:AG:12:ARG:HB2 | 7:AG:33:LEU:HD13 | 1.90 | 0.52 |
| 26:BA:1861:U:OP1 | 28:BC:200:HIS:NE2 | 2.34 | 0.52 |
| 28:BC:93:ASN:O | 28:BC:156:VAL:HA | 2.10 | 0.52 |
| 32:BG:105:THR:HG22 | 32:BG:121:LYS:HA | 1.91 | 0.52 |
| 39:BN:157:ASP:HA | 39:BN:160:GLU:OE2 | 2.10 | 0.52 |
| 41:BP:126:LYS:CD | 41:BP:131:GLU:HG3 | 2.40 | 0.52 |
| 42:BQ:31:GLU:HG3 | 42:BQ:58:VAL:HG11 | 1.90 | 0.52 |
| 1:AA:132:C:O2' | 23:AH:18:GLN:HB2 | 2.09 | 0.52 |
| 1:AA:407:U:H2' | 1:AA:408:G:C8 | 2.45 | 0.52 |
| 1:AA:554:A:H2' | 1:AA:555:C:C6 | 2.45 | 0.52 |
| 1:AA:651:G:H2' | 1:AA:652:U:C6 | 2.45 | 0.52 |
| 1:AA:1126:A:H2' | 1:AA:1127:A:C8 | 2.45 | 0.52 |
| 1:AA:1394:G:O2' | 1:AA:1395:U:H5' | 2.10 | 0.52 |
| 2:AB:240:LEU:O | 2:AB:244:GLU:N | 2.41 | 0.52 |
| 3:AC:25:LEU:HD22 | 3:AC:30:TYR:HD1 | 1.75 | 0.52 |
| 7:AG:202:VAL:HG22 | 9:AJ:68:ARG:NH2 | 2.25 | 0.52 |
| 8:AI:156:GLU:O | 8:AI:160:THR:HG23 | 2.10 | 0.52 |
| 33:BH:34:THR:HG21 | 33:BH:60:ILE:O | 2.10 | 0.52 |
| 37:BL:67:VAL:CG1 | 37:BL:101:ILE:HG21 | 2.38 | 0.52 |
| 55:Bd:6:LYS:O | 55:Bd:10:LYS:HG3 | 2.10 | 0.52 |
| 1:AA:205:C:H2' | 1:AA:206:A:C8 | 2.45 | 0.52 |
| 1:AA:560:C:H2' | 1:AA:561:A:O4' | 2.10 | 0.52 |
| 1:AA:1077:C:O2' | 19:AV:2:THR:HG23 | 2.10 | 0.52 |
| 1:AA:1200:G:OP1 | 24:AM:44:LYS:HB2 | 2.09 | 0.52 |
| 1:AA:1289:C:H2' | 1:AA:1290:C:C6 | 2.45 | 0.52 |
| 1:AA:1327:U:H1' | 22:AY:27:HIS:HA | 1.91 | 0.52 |
| 1:AA:1359:C:H2' | 1:AA:1360:A:C8 | 2.45 | 0.52 |
| 4:AD:48:VAL:HG13 | 12:AN:25:ILE:CG2 | 2.40 | 0.52 |
| 8:AI:89:THR:HG21 | 8:AI:163:ILE:HD12 | 1.92 | 0.52 |
| 16:AQ:75:LEU:HA | 16:AQ:80:VAL:CG2 | 2.40 | 0.52 |
| 17:AS:75:PRO:HD2 | 17:AS:102:VAL:HG22 | 1.92 | 0.52 |
| 18:AU:97:ASP:HB3 | 18:AU:99:GLN:HE22 | 1.75 | 0.52 |
| 22:AY:17:VAL:O | 22:AY:58:GLY:N | 2.32 | 0.52 |
| 26:BA:2543:G:O2' | 26:BA:2669:A:N1 | 2.42 | 0.52 |
| 29:BD:25:PRO:HG3 | 29:BD:203:ALA:HA | 1.92 | 0.52 |
| 29:BD:76:GLU:OE2 | 29:BD:152:LEU:HD22 | 2.09 | 0.52 |
| 30:BE:121:THR:O | 30:BE:138:LEU:HD13 | 2.10 | 0.52 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 30:BE:195:VAL:HB | 30:BE:215:VAL:CG2 | 2.39 | 0.52 |
| 34:BI:44:PHE:HB2 | 34:BI:137:ALA:HB1 | 1.91 | 0.52 |
| 39:BN:44:VAL:HG11 | 39:BN:82:ALA:HA | 1.92 | 0.52 |
| 40:BO:21:ILE:HD13 | 40:BO:44:GLU:HG2 | 1.91 | 0.52 |
| 49:BX:128:ILE:O | 49:BX:138:LEU:HD11 | 2.10 | 0.52 |
| 1:AA:813:C:H2' | 1:AA:814:G:O4' | 2.10 | 0.52 |
| 3:AC:12:TYR:CE1 | 15:AR:16:LYS:HD3 | 2.45 | 0.52 |
| 5:AE:117:VAL:CG1 | 5:AE:129:ALA:HB1 | 2.37 | 0.52 |
| 19:AV:79:GLY:O | 19:AV:90:LYS:HG3 | 2.10 | 0.52 |
| 23:AH:39:ASP:HA | 23:AH:49:LYS:CD | 2.40 | 0.52 |
| 26:BA:1754:G:P | 53:Bb:9:GLY:HA2 | 2.50 | 0.52 |
| 28:BC:87:ALA:HB3 | 28:BC:95:VAL:HG12 | 1.91 | 0.52 |
| 30:BE:152:THR:O | 30:BE:156:ILE:HG12 | 2.09 | 0.52 |
| 33:BH:63:HIS:CE1 | 38:BM:5:PHE:HB2 | 2.45 | 0.52 |
| 39:BN:45:GLN:HG3 | 39:BN:63:VAL:HG12 | 1.90 | 0.52 |
| 40:BO:109:LEU:HA | 40:BO:112:ILE:HG22 | 1.91 | 0.52 |
| 49:BX:59:ALA:HA | 49:BX:100:ILE:HG13 | 1.92 | 0.52 |
| 50:BY:46:CYS:SG | 50:BY:51:LYS:HE2 | 2.50 | 0.52 |
| 51:BZ:13:ILE:CD1 | 51:BZ:34:VAL:HG13 | 2.40 | 0.52 |
| 55:Bd:13:ALA:O | 55:Bd:17:LYS:HG2 | 2.09 | 0.52 |
| 1:AA:144:G:H5' | 23:AH:11:PRO:HA | 1.91 | 0.52 |
| 1:AA:896:U:H2' | 1:AA:897:G:C8 | 2.45 | 0.52 |
| 1:AA:905:A:O2' | 1:AA:930:C:H1' | 2.10 | 0.52 |
| 1:AA:930:C:H42 | 1:AA:1168:G:H1 | 1.58 | 0.52 |
| 1:AA:1116:A:H5'' | 25:AT:60:ARG:HE | 1.74 | 0.52 |
| 1:AA:1173:C:N4 | 14:AP:147:LYS:HE2 | 2.25 | 0.52 |
| 1:AA:1226:A:H4' | 1:AA:1228:U:C5 | 2.45 | 0.52 |
| 1:AA:1249:U:O2' | 1:AA:1250:C:H5' | 2.10 | 0.52 |
| 2:AB:170:VAL:HG21 | 2:AB:180:LEU:CD2 | 2.40 | 0.52 |
| 3:AC:88:PRO:HG2 | 3:AC:89:GLU:OE1 | 2.09 | 0.52 |
| 5:AE:173:HIS:CE1 | 5:AE:175:GLU:HB2 | 2.44 | 0.52 |
| 11:AL:25:GLY:O | 11:AL:27:VAL:HG23 | 2.10 | 0.52 |
| 13:AO:38:PRO:HB3 | 13:AO:77:ARG:HD3 | 1.92 | 0.52 |
| 18:AU:98:LEU:HG | 18:AU:102:MET:SD | 2.49 | 0.52 |
| 23:AH:74:ARG:O | 23:AH:75:ILE:HD13 | 2.09 | 0.52 |
| 24:AM:34:ILE:HG13 | 24:AM:75:ILE:CG2 | 2.38 | 0.52 |
| 40:BO:3:LYS:O | 40:BO:7:VAL:HG23 | 2.10 | 0.52 |
| 47:BV:13:LEU:HD11 | 47:BV:32:CYS:HA | 1.92 | 0.52 |
| 47:BV:57:ILE:O | 47:BV:61:LYS:HG3 | 2.08 | 0.52 |
| 48:BW:27:GLU:O | 48:BW:31:GLU:HG2 | 2.10 | 0.52 |
| 1:AA:550:U:H2' | 1:AA:551:C:H5' | 1.92 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 1:AA:808:G:OP1 | 7:AG:135:LYS:NZ | 2.28 | 0.51 |
| 1:AA:961:G:H1' | 18:AU:70:HIS:CG | 2.46 | 0.51 |
| 1:AA:1054:G:OP1 | 24:AM:67:ARG:NH2 | 2.30 | 0.51 |
| 3:AC:163:ASP:OD1 | 3:AC:182:ILE:HB | 2.10 | 0.51 |
| 7:AG:7:TRP:HZ2 | 7:AG:37:GLU:HG2 | 1.74 | 0.51 |
| 8:AI:6:LYS:HG2 | 8:AI:12:ASP:HB2 | 1.90 | 0.51 |
| 19:AV:2:THR:HG21 | 19:AV:133:HIS:HB2 | 1.93 | 0.51 |
| 21:AX:16:LYS:CD | 21:AX:58:LEU:HD21 | 2.40 | 0.51 |
| 26:BA:339:U:O2' | 46:BU:2:ILE:O | 2.27 | 0.51 |
| 33:BH:60:ILE:HG22 | 33:BH:61:VAL:HG13 | 1.92 | 0.51 |
| 39:BN:22:HIS:CE1 | 43:BR:24:ILE:HD13 | 2.45 | 0.51 |
| 50:BY:53:LYS:O | 50:BY:57:THR:HG23 | 2.10 | 0.51 |
| 57:Bf:62:THR:HG22 | 57:Bf:86:LYS:CD | 2.40 | 0.51 |
| 1:AA:218:G:H2' | 1:AA:219:C:O4' | 2.10 | 0.51 |
| 1:AA:460:A:N1 | 1:AA:476:C:O2' | 2.40 | 0.51 |
| 1:AA:1195:A:H2' | 1:AA:1196:C:O4' | 2.11 | 0.51 |
| 1:AA:1198:A:N3 | 1:AA:1316:C:O2' | 2.36 | 0.51 |
| 1:AA:1226:A:H4' | 1:AA:1228:U:H5 | 1.76 | 0.51 |
| 1:AA:1316:C:H2' | 1:AA:1317:G:H8 | 1.76 | 0.51 |
| 6:AF:120:ILE:CG2 | 6:AF:149:THR:HA | 2.40 | 0.51 |
| 6:AF:154:ILE:HG22 | 6:AF:164:LYS:O | 2.10 | 0.51 |
| 14:AP:25:ARG:HG2 | 14:AP:30:ASP:CA | 2.40 | 0.51 |
| 19:AV:40:HIS:ND1 | 19:AV:41:LYS:HG2 | 2.25 | 0.51 |
| 20:AW:71:TYR:CG | 20:AW:77:MET:HG2 | 2.45 | 0.51 |
| 28:BC:118:LYS:O | 28:BC:121:ARG:HD3 | 2.09 | 0.51 |
| 37:BL:102:GLU:HG2 | 37:BL:119:GLU:OE2 | 2.10 | 0.51 |
| 1:AA:40:U:OP1 | 1:AA:290:C:H5' | 2.10 | 0.51 |
| 1:AA:666:G:H2' | 1:AA:667:C:C6 | 2.45 | 0.51 |
| 3:AC:128:CYS:N | 3:AC:151:ILE:HD11 | 2.26 | 0.51 |
| 6:AF:175:MET:O | 6:AF:218:ILE:N | 2.43 | 0.51 |
| 20:AW:51:ILE:HD11 | 20:AW:85:ILE:CG2 | 2.40 | 0.51 |
| 20:AW:54:ARG:O | 20:AW:66:GLY:HA2 | 2.10 | 0.51 |
| 20:AW:82:GLN:O | 20:AW:86:LEU:HG | 2.10 | 0.51 |
| 22:AY:38:LEU:CD1 | 22:AY:43:LYS:HE3 | 2.41 | 0.51 |
| 26:BA:321:A:OP2 | 30:BE:151:ARG:NH2 | 2.44 | 0.51 |
| 34:BI:30:GLN:HG2 | 34:BI:60:HIS:CE1 | 2.45 | 0.51 |
| 38:BM:68:VAL:CG1 | 38:BM:98:ILE:HG23 | 2.41 | 0.51 |
| 55:Bd:24:VAL:O | 55:Bd:28:VAL:HG23 | 2.10 | 0.51 |
| 1:AA:34:C:H2' | 1:AA:35:G:H8 | 1.75 | 0.51 |
| 1:AA:68:U:O2' | 1:AA:69:U:H2' | 2.11 | 0.51 |
| 1:AA:248:U:H1' | 1:AA:250:C:OP2 | 2.10 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:333:G:OP2 | 1:AA:333:G:N2 | 2.29 | 0.51 |
| 1:AA:403:U:H2' | 1:AA:404:G:C8 | 2.45 | 0.51 |
| 1:AA:566:U:H2' | 1:AA:567:U:O4' | 2.11 | 0.51 |
| 1:AA:678:U:H2' | 1:AA:679:A:C8 | 2.45 | 0.51 |
| 1:AA:1403:U:H2' | 1:AA:1404:C:C6 | 2.44 | 0.51 |
| 5:AE:112:ARG:NH1 | 5:AE:114:GLN:OE1 | 2.44 | 0.51 |
| 6:AF:188:ILE:HD13 | 6:AF:191:ILE:HD11 | 1.92 | 0.51 |
| 14:AP:71:PRO:HB2 | 14:AP:73:GLU:OE1 | 2.10 | 0.51 |
| 14:AP:134:ARG:HD2 | 14:AP:140:LYS:O | 2.08 | 0.51 |
| 19:AV:137:LEU:HD23 | 19:AV:140:LEU:HD12 | 1.92 | 0.51 |
| 26:BA:769:U:H5 | 54:Bc:15:THR:HG22 | 1.74 | 0.51 |
| 28:BC:90:LYS:HA | 53:Bb:87:MET:HE1 | 1.93 | 0.51 |
| 30:BE:74:LEU:HG | 30:BE:79:ARG:O | 2.11 | 0.51 |
| 30:BE:143:VAL:HA | 30:BE:242:GLU:CG | 2.41 | 0.51 |
| 33:BH:48:LEU:HD23 | 33:BH:99:ILE:HD12 | 1.92 | 0.51 |
| 34:BI:80:LYS:NZ | 34:BI:81:MET:HE2 | 2.25 | 0.51 |
| 36:BK:50:GLY:HA2 | 36:BK:117:VAL:HG12 | 1.93 | 0.51 |
| 39:BN:164:ALA:HB1 | 39:BN:168:LYS:NZ | 2.26 | 0.51 |
| 1:AA:22:G:H4' | 5:AE:14:PRO:O | 2.10 | 0.51 |
| 1:AA:1019:A:H5' | 7:AG:71:ARG:HH11 | 1.76 | 0.51 |
| 10:AK:39:ILE:HD11 | 10:AK:41:ARG:NH2 | 2.26 | 0.51 |
| 16:AQ:47:THR:HG23 | 16:AQ:86:GLU:OE2 | 2.10 | 0.51 |
| 18:AU:106:ARG:HB2 | 18:AU:109:GLU:HG3 | 1.91 | 0.51 |
| 26:BA:1234:C:C5 | 35:BJ:125:SER:HB3 | 2.46 | 0.51 |
| 31:BF:13:MET:SD | 31:BF:29:LEU:HB2 | 2.51 | 0.51 |
| 41:BP:90:PRO:O | 41:BP:94:GLN:HG2 | 2.09 | 0.51 |
| 55:Bd:37:HIS:HD2 | 55:Bd:39:ARG:HB2 | 1.76 | 0.51 |
| 58:AZ:26:CYS:HB3 | 58:AZ:30:GLY:N | 2.25 | 0.51 |
| 1:AA:171:G:H2' | 1:AA:173:A:OP2 | 2.10 | 0.51 |
| 1:AA:485:C:O2' | 1:AA:489:U:OP1 | 2.20 | 0.51 |
| 1:AA:571:C:H2' | 1:AA:572:U:O4' | 2.09 | 0.51 |
| 1:AA:615:G:H5'' | 4:AD:120:LYS:CE | 2.40 | 0.51 |
| 1:AA:992:G:N7 | 1:AA:1147:C:H5'' | 2.25 | 0.51 |
| 1:AA:1015:U:H2' | 1:AA:1016:G:O4' | 2.11 | 0.51 |
| 13:AO:83:CYS:SG | 13:AO:129:LEU:HD21 | 2.50 | 0.51 |
| 13:AO:102:LYS:O | 13:AO:120:LYS:HE2 | 2.11 | 0.51 |
| 23:AH:72:ARG:HG2 | 23:AH:97:ARG:HD3 | 1.91 | 0.51 |
| 26:BA:1291:G:OP2 | 30:BE:178:ARG:NH2 | 2.43 | 0.51 |
| 30:BE:30:ASP:N | 30:BE:30:ASP:OD1 | 2.43 | 0.51 |
| 30:BE:74:LEU:HD12 | 30:BE:77:SER:OG | 2.11 | 0.51 |
| 31:BF:155:PHE:O | 31:BF:158:GLU:HG3 | 2.11 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 33:BH:107:GLU:O | 33:BH:110:GLN:HG3 | 2.11 | 0.51 |
| 1:AA:285:A:OP1 | 13:AO:19:ARG:NH2 | 2.43 | 0.51 |
| 1:AA:345:A:H5'' | 13:AO:46:ARG:NH2 | 2.26 | 0.51 |
| 1:AA:809:A:H2' | 1:AA:810:A:C8 | 2.46 | 0.51 |
| 1:AA:1088:U:H2' | 1:AA:1089:G:C8 | 2.45 | 0.51 |
| 1:AA:1135:G:O2' | 15:AR:48:LYS:HD3 | 2.11 | 0.51 |
| 3:AC:21:PHE:CE2 | 3:AC:65:LEU:HD23 | 2.45 | 0.51 |
| 4:AD:160:GLU:HG2 | 4:AD:165:ARG:HE | 1.75 | 0.51 |
| 4:AD:189:GLU:OE1 | 4:AD:189:GLU:N | 2.44 | 0.51 |
| 6:AF:81:LEU:HD22 | 6:AF:177:VAL:HG23 | 1.92 | 0.51 |
| 6:AF:156:SER:O | 6:AF:160:LYS:HD3 | 2.11 | 0.51 |
| 10:AK:9:ARG:HE | 10:AK:15:LYS:HA | 1.75 | 0.51 |
| 10:AK:46:THR:HG21 | 10:AK:52:LYS:HG3 | 1.93 | 0.51 |
| 13:AO:46:ARG:O | 13:AO:73:ILE:HG12 | 2.10 | 0.51 |
| 14:AP:73:GLU:O | 14:AP:77:LYS:HG2 | 2.11 | 0.51 |
| 18:AU:56:LYS:O | 18:AU:60:GLU:HG2 | 2.11 | 0.51 |
| 23:AH:69:GLY:O | 23:AH:98:GLY:HA3 | 2.10 | 0.51 |
| 26:BA:1289:G:OP1 | 30:BE:181:ARG:N | 2.35 | 0.51 |
| 26:BA:1520:C:H4' | 41:BP:92:LYS:HE2 | 1.93 | 0.51 |
| 26:BA:2353:C:O2' | 26:BA:2386:A:O2' | 2.11 | 0.51 |
| 30:BE:178:ARG:HB3 | 30:BE:190:LYS:HB2 | 1.91 | 0.51 |
| 30:BE:198:VAL:HA | 30:BE:218:VAL:O | 2.10 | 0.51 |
| 32:BG:24:PHE:N | 32:BG:35:ARG:O | 2.43 | 0.51 |
| 58:AZ:9:CYS:HB3 | 58:AZ:13:GLY:H | 1.76 | 0.51 |
| 1:AA:74:A:H2' | 1:AA:75:A:C8 | 2.46 | 0.51 |
| 1:AA:147:A:H2' | 1:AA:148:A:C8 | 2.46 | 0.51 |
| 1:AA:413:G:H2' | 1:AA:414:A:C8 | 2.46 | 0.51 |
| 1:AA:1168:G:OP1 | 1:AA:1268:U:N3 | 2.44 | 0.51 |
| 1:AA:1235:A:H2' | 1:AA:1236:A:O4' | 2.09 | 0.51 |
| 1:AA:1266:A:H61 | 1:AA:1307:A:H2 | 1.59 | 0.51 |
| 1:AA:1321:A:H2' | 1:AA:1322:A:C8 | 2.46 | 0.51 |
| 1:AA:1437:A:H2' | 1:AA:1438:A:C8 | 2.46 | 0.51 |
| 6:AF:152:SER:HB2 | 6:AF:219:VAL:HB | 1.93 | 0.51 |
| 10:AK:75:PRO:HD2 | 10:AK:103:SER:OG | 2.11 | 0.51 |
| 13:AO:54:GLY:O | 13:AO:64:ILE:HG23 | 2.11 | 0.51 |
| 26:BA:245:C:O4' | 38:BM:141:VAL:HG22 | 2.10 | 0.51 |
| 26:BA:1898:U:O2' | 26:BA:1901:A:N7 | 2.40 | 0.51 |
| 29:BD:141:VAL:HG23 | 29:BD:172:LEU:HD11 | 1.93 | 0.51 |
| 35:BJ:135:VAL:HG12 | 35:BJ:139:MET:HE2 | 1.93 | 0.51 |
| 1:AA:364:A:H2' | 1:AA:365:A:H8 | 1.76 | 0.51 |
| 1:AA:669:A:OP2 | 16:AQ:8:ARG:NH2 | 2.41 | 0.51 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:893:G:H2' | 1:AA:894:C:C6 | 2.46 | 0.51 |
| 1:AA:1355:A:H1' | 26:BA:1933:A:C2 | 2.46 | 0.51 |
| 2:AB:129:GLY:HA2 | 2:AB:217:LEU:HD13 | 1.93 | 0.51 |
| 2:AB:219:LEU:O | 2:AB:223:LEU:HG | 2.11 | 0.51 |
| 4:AD:91:TYR:OH | 4:AD:182:ARG:NH2 | 2.42 | 0.51 |
| 4:AD:135:GLN:O | 4:AD:139:GLU:HG2 | 2.11 | 0.51 |
| 6:AF:37:LEU:O | 6:AF:79:VAL:HG22 | 2.10 | 0.51 |
| 22:AY:42:ASP:HB3 | 22:AY:45:ARG:CD | 2.41 | 0.51 |
| 23:AH:45:LEU:CB | 23:AH:48:TYR:HB2 | 2.41 | 0.51 |
| 26:BA:767:G:OP1 | 54:Bc:28:ASN:ND2 | 2.43 | 0.51 |
| 27:BB:30:U:OP2 | 39:BN:41:SER:OG | 2.17 | 0.51 |
| 28:BC:40:ARG:NH2 | 28:BC:80:ILE:HG21 | 2.24 | 0.51 |
| 31:BF:110:ASP:HB3 | 31:BF:113:ILE:HG12 | 1.91 | 0.51 |
| 36:BK:103:ILE:HD12 | 36:BK:123:LYS:HG3 | 1.91 | 0.51 |
| 54:Bc:18:LYS:NZ | 54:Bc:23:GLY:O | 2.28 | 0.51 |
| 1:AA:2:U:H2' | 7:AG:171:TRP:HE3 | 1.75 | 0.51 |
| 1:AA:114:U:H4' | 6:AF:138:GLY:N | 2.25 | 0.51 |
| 1:AA:484:C:H2' | 1:AA:485:C:C6 | 2.46 | 0.51 |
| 1:AA:898:C:H4' | 1:AA:910:A:N1 | 2.26 | 0.51 |
| 1:AA:1316:C:H2' | 1:AA:1317:G:C8 | 2.45 | 0.51 |
| 2:AB:222:TRP:CH2 | 2:AB:226:ARG:HD3 | 2.46 | 0.51 |
| 6:AF:24:SER:HB3 | 6:AF:41:ILE:CD1 | 2.41 | 0.51 |
| 6:AF:181:HIS:HA | 6:AF:184:GLU:OE2 | 2.11 | 0.51 |
| 10:AK:96:LYS:HD2 | 10:AK:112:ARG:HG2 | 1.92 | 0.51 |
| 17:AS:40:SER:O | 17:AS:48:VAL:HG12 | 2.11 | 0.51 |
| 18:AU:61:PHE:CZ | 18:AU:85:ILE:HG22 | 2.45 | 0.51 |
| 20:AW:35:VAL:CG2 | 20:AW:55:ILE:HD13 | 2.40 | 0.51 |
| 22:AY:17:VAL:HG22 | 22:AY:33:ILE:HD12 | 1.92 | 0.51 |
| 26:BA:18:G:C5' | 44:BS:4:ILE:HG23 | 2.41 | 0.51 |
| 26:BA:817:U:O2' | 26:BA:818:A:OP1 | 2.21 | 0.51 |
| 26:BA:1522:G:OP1 | 41:BP:100:ARG:NH1 | 2.38 | 0.51 |
| 26:BA:2358:G:OP1 | 57:Bf:80:ARG:HD2 | 2.11 | 0.51 |
| 29:BD:85:ILE:CD1 | 29:BD:179:SER:HB2 | 2.37 | 0.51 |
| 30:BE:136:ALA:HB2 | 30:BE:162:ALA:O | 2.10 | 0.51 |
| 38:BM:44:PRO:HB3 | 38:BM:119:TYR:OH | 2.11 | 0.51 |
| 40:BO:93:ASP:O | 40:BO:97:GLU:HG3 | 2.10 | 0.51 |
| 40:BO:107:LEU:HD12 | 40:BO:111:GLU:CB | 2.41 | 0.51 |
| 43:BR:73:VAL:HG22 | 43:BR:86:PHE:CE1 | 2.46 | 0.51 |
| 1:AA:65:U:H2' | 1:AA:66:U:C6 | 2.46 | 0.50 |
| 1:AA:932:G:H1' | 18:AU:90:GLY:O | 2.11 | 0.50 |
| 1:AA:1073:U:H2' | 1:AA:1074:G:C8 | 2.46 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 4:AD:43:LEU:HA | 4:AD:46:ARG:CD | 2.38 | 0.50 |
| 5:AE:144:LYS:HD2 | 5:AE:163:TYR:CE2 | 2.46 | 0.50 |
| 6:AF:101:LYS:HB2 | 6:AF:103:ARG:HE | 1.75 | 0.50 |
| 7:AG:7:TRP:CZ2 | 7:AG:37:GLU:HG2 | 2.46 | 0.50 |
| 12:AN:59:LEU:O | 12:AN:63:LEU:HG | 2.11 | 0.50 |
| 12:AN:94:ILE:HG21 | 12:AN:107:ILE:HD13 | 1.93 | 0.50 |
| 23:AH:64:LYS:HD2 | 23:AH:96:MET:HE2 | 1.93 | 0.50 |
| 24:AM:43:LYS:O | 24:AM:68:VAL:N | 2.43 | 0.50 |
| 26:BA:327:U:H5' | 46:BU:9:PRO:HB3 | 1.93 | 0.50 |
| 26:BA:861:U:H1' | 28:BC:1:MET:HB3 | 1.93 | 0.50 |
| 26:BA:2862:U:OP2 | 26:BA:2863:U:O2' | 2.21 | 0.50 |
| 1:AA:36:G:O2' | 1:AA:562:A:N1 | 2.35 | 0.50 |
| 1:AA:157:A:N6 | 1:AA:187:A:OP1 | 2.29 | 0.50 |
| 1:AA:278:U:H2' | 1:AA:279:G:C8 | 2.46 | 0.50 |
| 1:AA:606:A:H3' | 1:AA:666:G:H21 | 1.76 | 0.50 |
| 1:AA:1010:C:H42 | 1:AA:1043:G:H1 | 1.59 | 0.50 |
| 1:AA:1064:U:O2 | 1:AA:1065:U:O2' | 2.16 | 0.50 |
| 1:AA:1203:U:H4' | 1:AA:1204:C:O4' | 2.11 | 0.50 |
| 1:AA:1273:C:H2' | 1:AA:1274:U:C6 | 2.46 | 0.50 |
| 1:AA:1274:U:H2' | 1:AA:1275:C:C6 | 2.47 | 0.50 |
| 1:AA:1308:A:H2' | 1:AA:1309:U:O4' | 2.11 | 0.50 |
| 1:AA:1346:C:O2 | 1:AA:1440:A:N6 | 2.43 | 0.50 |
| 2:AB:105:ILE:HG23 | 2:AB:224:LEU:HD13 | 1.94 | 0.50 |
| 4:AD:23:ALA:HB3 | 4:AD:30:ALA:N | 2.26 | 0.50 |
| 6:AF:39:LEU:HD22 | 6:AF:60:LEU:CD2 | 2.40 | 0.50 |
| 8:AI:180:ALA:O | 8:AI:184:VAL:HG23 | 2.11 | 0.50 |
| 16:AQ:5:HIS:CB | 16:AQ:117:LEU:HD13 | 2.40 | 0.50 |
| 16:AQ:34:THR:O | 16:AQ:38:LEU:HG | 2.11 | 0.50 |
| 20:AW:52:ILE:HG22 | 20:AW:68:ALA:CB | 2.40 | 0.50 |
| 22:AY:37:VAL:O | 22:AY:43:LYS:HA | 2.11 | 0.50 |
| 26:BA:623:G:O2' | 26:BA:625:A:OP1 | 2.26 | 0.50 |
| 26:BA:1385:G:OP1 | 26:BA:2719:G:O2' | 2.26 | 0.50 |
| 26:BA:1782:G:OP2 | 41:BP:81:ARG:NH1 | 2.45 | 0.50 |
| 26:BA:2444:A:C8 | 57:Bf:54:LYS:HE3 | 2.47 | 0.50 |
| 29:BD:17:ARG:HB2 | 29:BD:216:ARG:NH2 | 2.26 | 0.50 |
| 1:AA:492:U:O2' | 13:AO:102:LYS:HB2 | 2.10 | 0.50 |
| 1:AA:583:A:C8 | 9:AJ:107:SER:HA | 2.46 | 0.50 |
| 1:AA:992:G:C6 | 1:AA:1146:U:H2' | 2.46 | 0.50 |
| 1:AA:1065:U:O2 | 1:AA:1228:U:H5' | 2.10 | 0.50 |
| 1:AA:1079:C:C2 | 19:AV:147:ALA:HA | 2.46 | 0.50 |
| 1:AA:1149:G:O4' | 15:AR:20:ARG:NH1 | 2.44 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|-------------------|--------------------------|-------------------|
| 1:AA:1196:C:C5 | 1:AA:1234:A:H5'' | 2.46 | 0.50 |
| 1:AA:1201:G:N2 | 1:AA:1206:C:O2' | 2.45 | 0.50 |
| 2:AB:176:ASP:O | 2:AB:180:LEU:HG | 2.11 | 0.50 |
| 3:AC:135:LYS:HE3 | 3:AC:174:GLY:HA3 | 1.93 | 0.50 |
| 5:AE:137:HIS:CE1 | 5:AE:167:PRO:HD2 | 2.46 | 0.50 |
| 6:AF:30:PRO:HG2 | 6:AF:31:HIS:CD2 | 2.45 | 0.50 |
| 6:AF:55:GLU:O | 6:AF:59:ILE:HG12 | 2.10 | 0.50 |
| 9:AJ:75:ILE:CD1 | 9:AJ:125:LEU:HB3 | 2.41 | 0.50 |
| 12:AN:19:ILE:HG12 | 12:AN:35:SER:CB | 2.40 | 0.50 |
| 15:AR:41:ALA:HB1 | 15:AR:46:PHE:HB2 | 1.92 | 0.50 |
| 21:AX:14:ARG:NH2 | 21:AX:61:LEU:HD21 | 2.26 | 0.50 |
| 26:BA:583:G:O6 | 49:BX:153:ARG:NH2 | 2.28 | 0.50 |
| 26:BA:713:A:H1' | 40:BO:125:PHE:HA | 1.94 | 0.50 |
| 26:BA:756:C:OP1 | 30:BE:90:ARG:NH2 | 2.38 | 0.50 |
| 27:BB:107:C:H2' | 27:BB:108:U:O4' | 2.11 | 0.50 |
| 29:BD:237:THR:HG22 | 29:BD:239:GLY:H | 1.76 | 0.50 |
| 33:BH:53:GLU:OE1 | 33:BH:79:LYS:HA | 2.11 | 0.50 |
| 1:AA:77:A:H2' | 1:AA:78:U:C6 | 2.46 | 0.50 |
| 1:AA:790:C:H5'' | 4:AD:127:LYS:HD3 | 1.93 | 0.50 |
| 1:AA:961:G:H3' | 1:AA:962:A:H8 | 1.76 | 0.50 |
| 1:AA:1009:U:OP1 | 7:AG:67:ARG:HD2 | 2.11 | 0.50 |
| 1:AA:1307:A:H2' | 1:AA:1308:A:C8 | 2.46 | 0.50 |
| 4:AD:144:ILE:O | 4:AD:148:ARG:HG2 | 2.10 | 0.50 |
| 11:AL:45:LYS:HZ2 | 11:AL:73:ILE:HG23 | 1.76 | 0.50 |
| 15:AR:34:ARG:HB2 | 24:AM:49:THR:CG2 | 2.42 | 0.50 |
| 16:AQ:150:ILE:HG12 | 50:BY:17:LYS:CE | 2.40 | 0.50 |
| 26:BA:57:A:HO2' | 55:Bd:37:HIS:CE1 | 2.29 | 0.50 |
| 26:BA:2743:A:N6 | 26:BA:2779:G:O2' | 2.42 | 0.50 |
| 40:BO:31:ASP:OD2 | 40:BO:117:PRO:HB2 | 2.12 | 0.50 |
| 44:BS:46:ARG:NH1 | 44:BS:89:LEU:HD11 | 2.26 | 0.50 |
| 45:BT:75:ILE:O | 45:BT:79:ILE:HG12 | 2.11 | 0.50 |
| 46:BU:78:THR:CG2 | 46:BU:93:PRO:HB2 | 2.40 | 0.50 |
| 50:BY:40:VAL:HG13 | 50:BY:59:ILE:CG2 | 2.39 | 0.50 |
| 1:AA:14:C:H2' | 1:AA:15:U:C6 | 2.46 | 0.50 |
| 1:AA:43:C:H4' | 1:AA:90:C:OP1 | 2.11 | 0.50 |
| 1:AA:717:G:N2 | 1:AA:743:A:OP2 | 2.44 | 0.50 |
| 2:AB:129:GLY:CA | 2:AB:217:LEU:HD13 | 2.42 | 0.50 |
| 16:AQ:17:PRO:HB3 | 21:AX:8:PRO:CD | 2.41 | 0.50 |
| 19:AV:61:LYS:O | 19:AV:65:ASP:HB2 | 2.12 | 0.50 |
| 26:BA:2797:U:O2' | 26:BA:2800:A:N3 | 2.39 | 0.50 |
| 27:BB:34:G:O2' | 27:BB:35:U:H5' | 2.12 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 29:BD:140:ASP:OD1 | 29:BD:140:ASP:N | 2.43 | 0.50 |
| 35:BJ:86:TYR:CE2 | 35:BJ:87:LYS:HE3 | 2.47 | 0.50 |
| 40:BO:58:ILE:HD11 | 40:BO:70:ILE:CD1 | 2.42 | 0.50 |
| 1:AA:16:G:H2' | 1:AA:17:C:C6 | 2.46 | 0.50 |
| 1:AA:458:C:H2' | 1:AA:470:G:C8 | 2.47 | 0.50 |
| 1:AA:605:G:H2' | 1:AA:607:G:OP1 | 2.12 | 0.50 |
| 1:AA:1106:C:H41 | 1:AA:1119:U:H3' | 1.76 | 0.50 |
| 2:AB:100:SER:O | 2:AB:104:ARG:HG2 | 2.11 | 0.50 |
| 3:AC:33:MET:SD | 3:AC:44:ILE:HG23 | 2.51 | 0.50 |
| 6:AF:93:ALA:CB | 6:AF:109:LEU:HB2 | 2.40 | 0.50 |
| 7:AG:51:GLU:C | 7:AG:74:VAL:HG23 | 2.36 | 0.50 |
| 7:AG:57:MET:HG2 | 7:AG:69:LYS:O | 2.12 | 0.50 |
| 15:AR:15:CYS:HB2 | 15:AR:32:LEU:CA | 2.41 | 0.50 |
| 18:AU:72:ARG:HB3 | 18:AU:108:GLY:HA3 | 1.92 | 0.50 |
| 19:AV:71:GLU:HG3 | 19:AV:92:LYS:NZ | 2.26 | 0.50 |
| 24:AM:46:VAL:HG22 | 24:AM:65:GLU:HA | 1.94 | 0.50 |
| 30:BE:180:GLY:O | 30:BE:183:LYS:NZ | 2.45 | 0.50 |
| 33:BH:83:GLU:OE1 | 33:BH:83:GLU:N | 2.41 | 0.50 |
| 38:BM:148:LEU:O | 38:BM:151:VAL:HG12 | 2.11 | 0.50 |
| 39:BN:148:ILE:HG22 | 39:BN:155:SER:HB3 | 1.93 | 0.50 |
| 40:BO:23:THR:O | 40:BO:27:ARG:HG2 | 2.12 | 0.50 |
| 40:BO:107:LEU:HD12 | 40:BO:111:GLU:HB2 | 1.93 | 0.50 |
| 44:BS:39:GLY:O | 44:BS:151:ARG:NH1 | 2.45 | 0.50 |
| 47:BV:46:PRO:CB | 47:BV:54:LYS:HD2 | 2.41 | 0.50 |
| 1:AA:223:C:H2' | 1:AA:224:U:O4' | 2.11 | 0.50 |
| 1:AA:382:C:OP1 | 5:AE:9:LYS:HE3 | 2.12 | 0.50 |
| 1:AA:604:A:H5'' | 4:AD:130:ARG:HD2 | 1.93 | 0.50 |
| 1:AA:703:G:H2' | 1:AA:704:A:C8 | 2.47 | 0.50 |
| 1:AA:942:G:H2' | 1:AA:943:C:O4' | 2.12 | 0.50 |
| 1:AA:1223:G:H2' | 1:AA:1224:G:C8 | 2.46 | 0.50 |
| 6:AF:116:LYS:CB | 6:AF:157:ILE:HD11 | 2.41 | 0.50 |
| 6:AF:203:MET:HE3 | 6:AF:209:ASP:OD1 | 2.12 | 0.50 |
| 8:AI:114:LYS:CD | 8:AI:119:SER:HA | 2.42 | 0.50 |
| 9:AJ:81:VAL:N | 9:AJ:123:GLY:O | 2.41 | 0.50 |
| 15:AR:32:LEU:CD1 | 15:AR:40:ILE:HD11 | 2.41 | 0.50 |
| 18:AU:85:ILE:HG12 | 18:AU:98:LEU:HD11 | 1.93 | 0.50 |
| 20:AW:21:PHE:CZ | 20:AW:66:GLY:HA3 | 2.46 | 0.50 |
| 27:BB:4:U:H3 | 27:BB:124:G:H1 | 1.60 | 0.50 |
| 27:BB:14:A:O2' | 27:BB:16:A:H2' | 2.11 | 0.50 |
| 28:BC:91:PRO:HD3 | 53:Bb:87:MET:CE | 2.39 | 0.50 |
| 28:BC:101:PRO:HD2 | 28:BC:104:PHE:CD2 | 2.47 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 28:BC:110:GLU:HB3 | 28:BC:115:ASP:HB3 | 1.93 | 0.50 |
| 29:BD:275:GLY:HA3 | 29:BD:290:VAL:O | 2.12 | 0.50 |
| 30:BE:5:LYS:HB3 | 30:BE:13:ALA:HB1 | 1.94 | 0.50 |
| 30:BE:20:PRO:HD2 | 30:BE:23:PHE:CD1 | 2.47 | 0.50 |
| 36:BK:81:ARG:HH21 | 36:BK:85:GLY:HA2 | 1.76 | 0.50 |
| 43:BR:30:ALA:HB2 | 43:BR:90:GLN:HG3 | 1.94 | 0.50 |
| 44:BS:16:SER:OG | 44:BS:99:ALA:N | 2.45 | 0.50 |
| 1:AA:34:C:H2' | 1:AA:35:G:C8 | 2.46 | 0.50 |
| 1:AA:96:C:N4 | 1:AA:217:U:O2' | 2.35 | 0.50 |
| 1:AA:254:C:H2' | 1:AA:255:U:C6 | 2.47 | 0.50 |
| 1:AA:283:G:H5' | 17:AS:59:LYS:HB2 | 1.94 | 0.50 |
| 1:AA:286:A:H2' | 1:AA:287:G:C4' | 2.41 | 0.50 |
| 1:AA:690:U:O2' | 16:AQ:48:ALA:O | 2.17 | 0.50 |
| 1:AA:926:C:H42 | 15:AR:10:ARG:HG2 | 1.77 | 0.50 |
| 1:AA:935:G:H5'' | 1:AA:965:A:O2' | 2.11 | 0.50 |
| 1:AA:1218:G:OP1 | 1:AA:1261:C:H4' | 2.11 | 0.50 |
| 1:AA:1468:G:H2' | 1:AA:1469:A:C8 | 2.46 | 0.50 |
| 4:AD:125:THR:OG1 | 4:AD:127:LYS:O | 2.22 | 0.50 |
| 5:AE:140:ILE:HB | 5:AE:160:ILE:CD1 | 2.40 | 0.50 |
| 6:AF:12:SER:HB3 | 6:AF:104:LEU:CD2 | 2.39 | 0.50 |
| 8:AI:5:TYR:OH | 11:AL:52:ILE:HG12 | 2.12 | 0.50 |
| 14:AP:55:ILE:HD11 | 14:AP:85:PHE:CD1 | 2.45 | 0.50 |
| 16:AQ:148:MET:SD | 16:AQ:151:THR:HA | 2.51 | 0.50 |
| 20:AW:17:ARG:HD3 | 20:AW:70:VAL:HG11 | 1.94 | 0.50 |
| 22:AY:18:ILE:HG22 | 22:AY:34:GLN:O | 2.12 | 0.50 |
| 25:AT:35:VAL:O | 25:AT:39:THR:HG22 | 2.12 | 0.50 |
| 26:BA:1056:G:O2' | 49:BX:14:ARG:HG2 | 2.12 | 0.50 |
| 29:BD:277:GLU:OE1 | 29:BD:294:TYR:OH | 2.19 | 0.50 |
| 30:BE:40:GLN:HG3 | 30:BE:226:GLU:OE2 | 2.11 | 0.50 |
| 41:BP:24:LEU:HD22 | 41:BP:50:ILE:HG12 | 1.94 | 0.50 |
| 46:BU:86:LYS:HG3 | 46:BU:88:ASP:N | 2.25 | 0.50 |
| 47:BV:13:LEU:CD1 | 47:BV:32:CYS:HA | 2.42 | 0.50 |
| 47:BV:49:THR:O | 47:BV:55:GLY:HA3 | 2.12 | 0.50 |
| 48:BW:60:ILE:O | 48:BW:64:LEU:HG | 2.11 | 0.50 |
| 1:AA:1322:A:H5' | 8:AI:56:ARG:CZ | 2.42 | 0.50 |
| 2:AB:157:PRO:HA | 2:AB:162:PHE:CD1 | 2.47 | 0.50 |
| 3:AC:157:PRO:O | 3:AC:162:VAL:HG22 | 2.11 | 0.50 |
| 5:AE:56:ARG:HD3 | 7:AG:147:PRO:HG3 | 1.94 | 0.50 |
| 5:AE:58:LEU:HB2 | 5:AE:97:ILE:HD11 | 1.94 | 0.50 |
| 6:AF:96:MET:HE1 | 6:AF:104:LEU:HB3 | 1.94 | 0.50 |
| 9:AJ:75:ILE:HD13 | 9:AJ:125:LEU:HB3 | 1.94 | 0.50 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|-------------------|--------------------------|-------------------|
| 24:AM:50:ARG:NH2 | 24:AM:57:GLY:O | 2.45 | 0.50 |
| 26:BA:481:G:O2' | 26:BA:506:U:O4 | 2.21 | 0.50 |
| 26:BA:2495:C:O2' | 34:BI:156:LYS:O | 2.26 | 0.50 |
| 27:BB:116:G:O6 | 39:BN:12:ARG:HD3 | 2.12 | 0.50 |
| 30:BE:156:ILE:HG23 | 30:BE:165:TYR:CE1 | 2.46 | 0.50 |
| 37:BL:85:ALA:CB | 37:BL:94:ILE:HG12 | 2.42 | 0.50 |
| 49:BX:148:LEU:HA | 49:BX:151:LYS:HG2 | 1.94 | 0.50 |
| 55:Bd:4:ASN:HB3 | 55:Bd:8:GLN:HB2 | 1.92 | 0.50 |
| 1:AA:113:C:H2' | 1:AA:114:U:C6 | 2.47 | 0.49 |
| 1:AA:413:G:N2 | 1:AA:430:U:H1' | 2.27 | 0.49 |
| 1:AA:567:U:H2' | 1:AA:568:A:H8 | 1.77 | 0.49 |
| 6:AF:4:GLN:NE2 | 6:AF:23:ILE:O | 2.44 | 0.49 |
| 6:AF:116:LYS:O | 6:AF:155:LEU:HD23 | 2.11 | 0.49 |
| 7:AG:79:ARG:HG2 | 7:AG:163:GLU:O | 2.12 | 0.49 |
| 8:AI:12:ASP:HB3 | 8:AI:15:GLU:OE2 | 2.12 | 0.49 |
| 10:AK:37:SER:O | 10:AK:59:ASN:ND2 | 2.45 | 0.49 |
| 12:AN:61:ASP:HA | 12:AN:64:LYS:CE | 2.39 | 0.49 |
| 14:AP:21:ARG:HB2 | 14:AP:68:GLY:HA2 | 1.93 | 0.49 |
| 14:AP:47:VAL:HG13 | 14:AP:51:THR:OG1 | 2.11 | 0.49 |
| 20:AW:76:ARG:NH1 | 20:AW:79:GLN:OE1 | 2.45 | 0.49 |
| 23:AH:75:ILE:O | 23:AH:93:ARG:HA | 2.12 | 0.49 |
| 24:AM:39:PRO:HA | 24:AM:71:ARG:CD | 2.41 | 0.49 |
| 26:BA:2030:G:H4' | 44:BS:55:MET:HE1 | 1.94 | 0.49 |
| 26:BA:2811:U:O4 | 29:BD:156:PRO:HD2 | 2.12 | 0.49 |
| 30:BE:11:GLY:HA3 | 30:BE:161:ALA:HB1 | 1.93 | 0.49 |
| 32:BG:99:VAL:HA | 32:BG:107:ILE:O | 2.12 | 0.49 |
| 42:BQ:8:ILE:HD13 | 42:BQ:25:LYS:HA | 1.93 | 0.49 |
| 45:BT:24:PHE:CD2 | 45:BT:64:LEU:HD13 | 2.47 | 0.49 |
| 48:BW:8:GLU:O | 48:BW:12:MET:HG3 | 2.12 | 0.49 |
| 49:BX:31:HIS:CE1 | 49:BX:117:ARG:HG2 | 2.47 | 0.49 |
| 56:Be:14:LYS:CE | 56:Be:23:ARG:HD3 | 2.42 | 0.49 |
| 1:AA:359:C:H2' | 1:AA:360:G:C8 | 2.47 | 0.49 |
| 1:AA:872:G:H2' | 1:AA:873:G:O4' | 2.13 | 0.49 |
| 1:AA:1103:G:OP1 | 25:AT:32:LYS:NZ | 2.43 | 0.49 |
| 1:AA:1136:C:O2 | 3:AC:154:SER:HB3 | 2.11 | 0.49 |
| 3:AC:191:ASP:OD2 | 25:AT:46:ILE:HD13 | 2.13 | 0.49 |
| 7:AG:37:GLU:HB3 | 7:AG:40:ILE:CD1 | 2.42 | 0.49 |
| 13:AO:127:VAL:HG21 | 13:AO:137:MET:HG2 | 1.92 | 0.49 |
| 14:AP:55:ILE:HA | 14:AP:88:ILE:CD1 | 2.41 | 0.49 |
| 15:AR:25:VAL:N | 15:AR:32:LEU:O | 2.36 | 0.49 |
| 19:AV:139:LEU:HB3 | 19:AV:142:LYS:HG3 | 1.93 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 20:AW:32:ARG:HB3 | 20:AW:52:ILE:HD11 | 1.93 | 0.49 |
| 20:AW:45:ALA:CB | 20:AW:50:LEU:HB2 | 2.41 | 0.49 |
| 21:AX:28:GLY:O | 21:AX:51:SER:OG | 2.17 | 0.49 |
| 26:BA:396:G:OP1 | 38:BM:98:ILE:HG13 | 2.12 | 0.49 |
| 26:BA:1043:G:O3' | 34:BI:15:ARG:NH1 | 2.44 | 0.49 |
| 28:BC:24:TYR:HA | 28:BC:50:PRO:HG2 | 1.94 | 0.49 |
| 28:BC:40:ARG:O | 28:BC:62:GLU:HG2 | 2.11 | 0.49 |
| 30:BE:167:ASP:OD1 | 30:BE:170:ARG:HD3 | 2.12 | 0.49 |
| 33:BH:29:LYS:H | 33:BH:101:ASP:HB3 | 1.77 | 0.49 |
| 34:BI:6:GLY:O | 34:BI:10:ARG:HB2 | 2.12 | 0.49 |
| 39:BN:79:THR:HG23 | 39:BN:118:ALA:HB2 | 1.94 | 0.49 |
| 40:BO:58:ILE:O | 40:BO:62:THR:OG1 | 2.14 | 0.49 |
| 44:BS:75:MET:HG2 | 44:BS:76:ALA:O | 2.11 | 0.49 |
| 49:BX:2:TYR:HA | 49:BX:55:GLY:O | 2.11 | 0.49 |
| 1:AA:63:U:C1' | 1:AA:148:A:H1' | 2.42 | 0.49 |
| 1:AA:372:U:H5' | 6:AF:22:TRP:CH2 | 2.47 | 0.49 |
| 1:AA:1240:A:C5 | 1:AA:1241:U:H1' | 2.47 | 0.49 |
| 1:AA:1342:C:H1' | 1:AA:1346:C:H41 | 1.77 | 0.49 |
| 1:AA:1362:C:H2' | 1:AA:1363:C:C6 | 2.47 | 0.49 |
| 2:AB:112:LEU:HD11 | 2:AB:169:ILE:HD11 | 1.93 | 0.49 |
| 4:AD:116:VAL:HB | 4:AD:192:THR:HG23 | 1.93 | 0.49 |
| 6:AF:158:PRO:O | 6:AF:160:LYS:HG2 | 2.12 | 0.49 |
| 7:AG:55:ILE:CD1 | 7:AG:72:ALA:HB2 | 2.42 | 0.49 |
| 8:AI:17:GLU:N | 8:AI:17:GLU:OE1 | 2.45 | 0.49 |
| 9:AJ:16:ASN:O | 9:AJ:20:ILE:HG13 | 2.11 | 0.49 |
| 17:AS:49:ILE:HG22 | 17:AS:70:ILE:O | 2.12 | 0.49 |
| 26:BA:2572:A:O2' | 36:BK:33:SER:OG | 2.24 | 0.49 |
| 28:BC:177:TYR:O | 28:BC:181:LYS:HG2 | 2.11 | 0.49 |
| 29:BD:333:GLU:OE1 | 47:BV:14:GLU:HG3 | 2.12 | 0.49 |
| 30:BE:20:PRO:HD2 | 30:BE:23:PHE:CE1 | 2.48 | 0.49 |
| 30:BE:96:LYS:HB3 | 30:BE:98:GLU:OE1 | 2.12 | 0.49 |
| 36:BK:37:TYR:CE2 | 36:BK:45:PRO:HA | 2.47 | 0.49 |
| 44:BS:46:ARG:HH22 | 44:BS:93:LYS:HB2 | 1.77 | 0.49 |
| 48:BW:57:ILE:O | 48:BW:61:GLN:HG3 | 2.12 | 0.49 |
| 1:AA:218:G:OP1 | 17:AS:70:ILE:HD12 | 2.12 | 0.49 |
| 1:AA:233:G:H1' | 10:AK:8:ARG:NH2 | 2.26 | 0.49 |
| 1:AA:1103:G:OP2 | 25:AT:44:LYS:NZ | 2.29 | 0.49 |
| 1:AA:1123:A:H2' | 1:AA:1125:G:C5 | 2.46 | 0.49 |
| 1:AA:1203:U:H5'' | 1:AA:1205:C:H1' | 1.95 | 0.49 |
| 2:AB:59:LEU:HD11 | 2:AB:106:ARG:HE | 1.77 | 0.49 |
| 4:AD:112:LYS:HB2 | 4:AD:153:ASP:HA | 1.94 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 9:AJ:70:ASN:ND2 | 9:AJ:130:TYR:O | 2.46 | 0.49 |
| 10:AK:63:VAL:CG1 | 10:AK:123:LEU:HB2 | 2.42 | 0.49 |
| 14:AP:25:ARG:HG2 | 14:AP:30:ASP:HA | 1.93 | 0.49 |
| 16:AQ:119:GLU:O | 16:AQ:123:ARG:HG3 | 2.12 | 0.49 |
| 20:AW:32:ARG:HA | 20:AW:55:ILE:CD1 | 2.40 | 0.49 |
| 20:AW:73:ASP:O | 20:AW:77:MET:HG3 | 2.12 | 0.49 |
| 26:BA:108:C:O2' | 54:Bc:20:ARG:HB3 | 2.12 | 0.49 |
| 29:BD:47:VAL:HG22 | 29:BD:287:TYR:CZ | 2.46 | 0.49 |
| 30:BE:54:GLY:O | 30:BE:82:ARG:HA | 2.12 | 0.49 |
| 30:BE:55:MET:HG3 | 30:BE:90:ARG:HB3 | 1.94 | 0.49 |
| 45:BT:11:GLU:O | 45:BT:15:MET:HG3 | 2.12 | 0.49 |
| 47:BV:5:LYS:NZ | 47:BV:10:GLY:O | 2.31 | 0.49 |
| 1:AA:311:A:O2' | 1:AA:313:G:OP2 | 2.29 | 0.49 |
| 1:AA:440:G:O2' | 1:AA:489:U:O2' | 2.28 | 0.49 |
| 1:AA:905:A:O2' | 1:AA:930:C:O2 | 2.16 | 0.49 |
| 1:AA:1106:C:N4 | 1:AA:1119:U:H3' | 2.27 | 0.49 |
| 4:AD:112:LYS:HB2 | 4:AD:153:ASP:CB | 2.41 | 0.49 |
| 5:AE:113:LEU:HD13 | 5:AE:152:VAL:CG1 | 2.38 | 0.49 |
| 6:AF:100:GLU:HA | 6:AF:182:SER:O | 2.12 | 0.49 |
| 9:AJ:53:ILE:HG23 | 9:AJ:60:ILE:HG13 | 1.93 | 0.49 |
| 9:AJ:66:VAL:HG13 | 9:AJ:68:ARG:HG2 | 1.94 | 0.49 |
| 13:AO:48:ILE:HA | 13:AO:96:ASP:O | 2.12 | 0.49 |
| 14:AP:20:LEU:CD2 | 14:AP:22:HIS:HB2 | 2.43 | 0.49 |
| 14:AP:55:ILE:HA | 14:AP:88:ILE:HD12 | 1.94 | 0.49 |
| 15:AR:24:LEU:HD11 | 15:AR:26:ARG:HE | 1.77 | 0.49 |
| 18:AU:20:GLY:HA3 | 18:AU:37:LEU:HD22 | 1.93 | 0.49 |
| 18:AU:72:ARG:NH2 | 18:AU:108:GLY:O | 2.45 | 0.49 |
| 26:BA:769:U:C5 | 54:Bc:15:THR:HG22 | 2.48 | 0.49 |
| 26:BA:784:G:H4' | 41:BP:87:ALA:HB2 | 1.94 | 0.49 |
| 26:BA:897:C:H41 | 37:BL:4:LYS:HG2 | 1.76 | 0.49 |
| 26:BA:1826:U:O2' | 28:BC:173:ALA:HB2 | 2.12 | 0.49 |
| 30:BE:29:PRO:HD2 | 40:BO:15:PRO:HG2 | 1.93 | 0.49 |
| 45:BT:45:PHE:HE2 | 45:BT:78:ARG:HH12 | 1.60 | 0.49 |
| 58:AZ:47:THR:HG21 | 58:AZ:54:LYS:HE2 | 1.93 | 0.49 |
| 1:AA:64:G:O2' | 1:AA:128:A:N1 | 2.37 | 0.49 |
| 1:AA:243:U:OP2 | 10:AK:54:ARG:NH2 | 2.45 | 0.49 |
| 1:AA:259:C:C1' | 10:AK:12:THR:HG22 | 2.42 | 0.49 |
| 1:AA:733:A:O2' | 1:AA:735:A:N7 | 2.32 | 0.49 |
| 1:AA:997:G:OP1 | 3:AC:133:SER:HB2 | 2.12 | 0.49 |
| 7:AG:17:VAL:HG12 | 7:AG:44:LEU:HD21 | 1.93 | 0.49 |
| 8:AI:81:ALA:CB | 8:AI:158:LEU:HD23 | 2.43 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 8:AI:186:LYS:O | 8:AI:189:ARG:HG2 | 2.12 | 0.49 |
| 18:AU:61:PHE:O | 18:AU:83:LYS:NZ | 2.44 | 0.49 |
| 25:AT:8:ASN:HA | 25:AT:11:ARG:NH1 | 2.28 | 0.49 |
| 26:BA:451:C:H4' | 30:BE:50:ARG:HD2 | 1.94 | 0.49 |
| 26:BA:1077:G:C8 | 49:BX:11:VAL:HG22 | 2.47 | 0.49 |
| 26:BA:1311:G:H5'' | 52:Ba:98:LYS:HG3 | 1.94 | 0.49 |
| 27:BB:49:A:H2' | 27:BB:50:C:C6 | 2.48 | 0.49 |
| 27:BB:50:C:OP1 | 39:BN:115:ARG:N | 2.38 | 0.49 |
| 29:BD:128:ASN:O | 29:BD:132:LEU:HG | 2.12 | 0.49 |
| 29:BD:135:GLU:OE2 | 29:BD:137:ARG:HD3 | 2.11 | 0.49 |
| 32:BG:123:ILE:O | 32:BG:126:THR:OG1 | 2.17 | 0.49 |
| 33:BH:54:ASP:OD2 | 33:BH:81:GLN:HG2 | 2.12 | 0.49 |
| 36:BK:98:THR:O | 36:BK:105:LYS:NZ | 2.46 | 0.49 |
| 40:BO:38:ASP:O | 40:BO:42:ARG:HG3 | 2.13 | 0.49 |
| 40:BO:91:PHE:CE2 | 40:BO:106:CYS:HB3 | 2.48 | 0.49 |
| 50:BY:70:GLU:O | 50:BY:74:VAL:HG23 | 2.13 | 0.49 |
| 1:AA:2:U:H2' | 7:AG:171:TRP:CE3 | 2.47 | 0.49 |
| 1:AA:399:A:H2' | 1:AA:400:U:N1 | 2.28 | 0.49 |
| 1:AA:765:C:C5' | 16:AQ:1:MET:HG2 | 2.43 | 0.49 |
| 1:AA:1076:C:H4' | 19:AV:126:SER:HB3 | 1.94 | 0.49 |
| 1:AA:1251:G:H2' | 1:AA:1252:G:N2 | 2.27 | 0.49 |
| 3:AC:17:MET:HE1 | 3:AC:73:TYR:CD1 | 2.47 | 0.49 |
| 3:AC:179:LYS:O | 3:AC:181:ARG:HD2 | 2.12 | 0.49 |
| 7:AG:134:GLY:O | 7:AG:187:ALA:HB1 | 2.12 | 0.49 |
| 8:AI:100:ALA:HA | 8:AI:165:ALA:O | 2.12 | 0.49 |
| 11:AL:100:PHE:HA | 11:AL:103:TYR:CE1 | 2.48 | 0.49 |
| 13:AO:52:LYS:HD2 | 13:AO:89:VAL:HG23 | 1.95 | 0.49 |
| 26:BA:153:C:OP1 | 38:BM:95:GLY:N | 2.40 | 0.49 |
| 26:BA:1855:A:H5'' | 28:BC:125:VAL:HG11 | 1.93 | 0.49 |
| 28:BC:44:VAL:HG22 | 28:BC:58:LYS:O | 2.13 | 0.49 |
| 29:BD:314:PRO:CB | 29:BD:317:ALA:HB2 | 2.41 | 0.49 |
| 34:BI:31:VAL:HA | 34:BI:64:GLU:OE1 | 2.12 | 0.49 |
| 34:BI:48:ILE:HG21 | 34:BI:143:PHE:HE1 | 1.77 | 0.49 |
| 34:BI:51:ILE:HG22 | 34:BI:132:LYS:HG2 | 1.94 | 0.49 |
| 36:BK:114:ALA:O | 36:BK:117:VAL:HG22 | 2.13 | 0.49 |
| 41:BP:60:ARG:O | 41:BP:64:ARG:HG3 | 2.11 | 0.49 |
| 44:BS:41:LYS:NZ | 44:BS:107:ALA:O | 2.46 | 0.49 |
| 1:AA:521:G:N1 | 1:AA:700:A:OP2 | 2.30 | 0.49 |
| 1:AA:616:U:H2' | 1:AA:617:A:C8 | 2.47 | 0.49 |
| 1:AA:1114:U:H1' | 1:AA:1115:U:C5 | 2.47 | 0.49 |
| 6:AF:174:ALA:HB1 | 6:AF:217:VAL:CG2 | 2.42 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 9:AJ:9:ASN:O | 9:AJ:13:THR:HG23 | 2.13 | 0.49 |
| 9:AJ:103:ILE:N | 9:AJ:127:ALA:O | 2.46 | 0.49 |
| 19:AV:37:THR:HG1 | 19:AV:54:ARG:HH21 | 1.53 | 0.49 |
| 24:AM:9:SER:HA | 24:AM:69:HIS:O | 2.13 | 0.49 |
| 26:BA:563:G:O2' | 26:BA:564:A:O4' | 2.26 | 0.49 |
| 26:BA:1657:C:H5'' | 55:Bd:6:LYS:HD2 | 1.94 | 0.49 |
| 26:BA:1721:U:O2' | 26:BA:2699:U:OP1 | 2.30 | 0.49 |
| 27:BB:7:G:H2' | 27:BB:8:G:C8 | 2.47 | 0.49 |
| 28:BC:86:ASP:OD1 | 28:BC:86:ASP:N | 2.45 | 0.49 |
| 29:BD:24:ILE:HG23 | 29:BD:308:LEU:HD23 | 1.94 | 0.49 |
| 31:BF:42:ALA:HB2 | 31:BF:57:ILE:CG1 | 2.43 | 0.49 |
| 33:BH:107:GLU:OE1 | 33:BH:107:GLU:N | 2.40 | 0.49 |
| 38:BM:157:ARG:O | 38:BM:162:ARG:NH1 | 2.45 | 0.49 |
| 44:BS:53:ILE:HA | 44:BS:82:ILE:CD1 | 2.43 | 0.49 |
| 46:BU:60:PHE:CD1 | 46:BU:82:VAL:HG13 | 2.48 | 0.49 |
| 1:AA:162:A:H1' | 10:AK:64:THR:HG21 | 1.94 | 0.49 |
| 1:AA:246:C:H2' | 1:AA:247:G:O4' | 2.12 | 0.49 |
| 1:AA:407:U:H1' | 5:AE:131:GLN:HE22 | 1.77 | 0.49 |
| 1:AA:690:U:O5' | 16:AQ:55:ARG:NH1 | 2.42 | 0.49 |
| 1:AA:1190:A:H61 | 1:AA:1241:U:H3 | 1.61 | 0.49 |
| 1:AA:1258:G:H2' | 1:AA:1259:G:C8 | 2.48 | 0.49 |
| 4:AD:111:THR:HA | 4:AD:149:ALA:O | 2.12 | 0.49 |
| 6:AF:3:HIS:HB3 | 6:AF:22:TRP:CD1 | 2.48 | 0.49 |
| 7:AG:128:VAL:HG12 | 7:AG:145:PRO:HA | 1.95 | 0.49 |
| 8:AI:6:LYS:HE2 | 8:AI:12:ASP:N | 2.28 | 0.49 |
| 8:AI:82:PHE:CD1 | 8:AI:93:PRO:HB2 | 2.45 | 0.49 |
| 10:AK:37:SER:HB2 | 10:AK:57:GLN:CB | 2.39 | 0.49 |
| 18:AU:52:ASP:O | 18:AU:56:LYS:HG3 | 2.13 | 0.49 |
| 26:BA:833:G:OP1 | 44:BS:128:ARG:NH2 | 2.44 | 0.49 |
| 26:BA:1842:C:OP1 | 28:BC:8:GLN:NE2 | 2.42 | 0.49 |
| 26:BA:2412:G:O2' | 57:Bf:77:ALA:O | 2.31 | 0.49 |
| 26:BA:2822:U:O2' | 26:BA:2881:C:OP1 | 2.14 | 0.49 |
| 29:BD:203:ALA:HB2 | 29:BD:309:ILE:HD13 | 1.95 | 0.49 |
| 30:BE:140:ILE:CD1 | 30:BE:164:LEU:HD11 | 2.41 | 0.49 |
| 34:BI:45:PRO:HB2 | 34:BI:171:LEU:CD2 | 2.43 | 0.49 |
| 44:BS:31:ARG:HB2 | 44:BS:116:ILE:HD13 | 1.94 | 0.49 |
| 52:Ba:132:ILE:HG22 | 52:Ba:137:LYS:HB2 | 1.93 | 0.49 |
| 55:Bd:41:ARG:CZ | 55:Bd:48:LEU:HD21 | 2.42 | 0.49 |
| 56:Be:15:LYS:O | 56:Be:16:ILE:HD13 | 2.13 | 0.49 |
| 1:AA:450:G:O2' | 1:AA:474:C:H1' | 2.12 | 0.49 |
| 1:AA:484:C:H2' | 1:AA:485:C:H6 | 1.78 | 0.49 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 1:AA:943:C:H2' | 1:AA:944:A:C8 | 2.47 | 0.49 |
| 2:AB:69:GLY:HA2 | 2:AB:72:ILE:CD1 | 2.42 | 0.49 |
| 2:AB:152:GLY:O | 2:AB:156:ASN:HB2 | 2.12 | 0.49 |
| 3:AC:72:LYS:HE2 | 3:AC:73:TYR:OH | 2.12 | 0.49 |
| 3:AC:188:VAL:O | 3:AC:189:LEU:HD22 | 2.12 | 0.49 |
| 6:AF:119:ARG:HA | 6:AF:152:SER:HA | 1.94 | 0.49 |
| 8:AI:57:LEU:HD22 | 8:AI:97:LEU:HD21 | 1.93 | 0.49 |
| 9:AJ:90:GLU:O | 9:AJ:94:LEU:HB2 | 2.13 | 0.49 |
| 10:AK:39:ILE:HA | 10:AK:57:GLN:O | 2.12 | 0.49 |
| 12:AN:60:ALA:O | 12:AN:64:LYS:HG3 | 2.12 | 0.49 |
| 13:AO:16:LYS:NZ | 13:AO:17:ASP:OD1 | 2.46 | 0.49 |
| 14:AP:23:LEU:HD23 | 31:BF:109:TYR:CE2 | 2.46 | 0.49 |
| 21:AX:33:LYS:CA | 21:AX:46:PRO:HD3 | 2.41 | 0.49 |
| 23:AH:30:ILE:HA | 23:AH:101:ILE:HG22 | 1.95 | 0.49 |
| 25:AT:57:LYS:O | 25:AT:61:MET:HB2 | 2.13 | 0.49 |
| 26:BA:54:C:OP2 | 48:BW:51:ARG:NH2 | 2.30 | 0.49 |
| 26:BA:892:U:OP2 | 37:BL:21:ARG:NH1 | 2.41 | 0.49 |
| 26:BA:994:C:OP1 | 34:BI:131:GLN:NE2 | 2.44 | 0.49 |
| 29:BD:81:ARG:HA | 29:BD:185:THR:O | 2.13 | 0.49 |
| 50:BY:9:ASP:OD1 | 50:BY:9:ASP:N | 2.46 | 0.49 |
| 53:Bb:50:THR:HG21 | 53:Bb:63:LYS:CG | 2.41 | 0.49 |
| 1:AA:594:U:O2 | 9:AJ:57:LYS:HG2 | 2.13 | 0.48 |
| 1:AA:846:A:C5 | 1:AA:847:G:H1' | 2.48 | 0.48 |
| 1:AA:979:G:H2' | 1:AA:980:C:O4' | 2.12 | 0.48 |
| 1:AA:1080:C:O2 | 11:AL:26:LYS:NZ | 2.38 | 0.48 |
| 2:AB:81:MET:HE2 | 2:AB:195:ASP:O | 2.13 | 0.48 |
| 3:AC:120:VAL:HG11 | 3:AC:182:ILE:HD12 | 1.94 | 0.48 |
| 8:AI:22:GLY:CA | 22:AY:60:ILE:HB | 2.41 | 0.48 |
| 10:AK:36:ILE:HA | 10:AK:58:SER:O | 2.13 | 0.48 |
| 11:AL:4:VAL:HA | 11:AL:18:ALA:O | 2.12 | 0.48 |
| 12:AN:5:LYS:HB3 | 12:AN:68:ILE:HG13 | 1.95 | 0.48 |
| 14:AP:53:LYS:O | 14:AP:57:LYS:HG3 | 2.13 | 0.48 |
| 17:AS:9:ILE:HG21 | 17:AS:103:VAL:HG12 | 1.94 | 0.48 |
| 19:AV:31:TRP:HZ3 | 19:AV:97:VAL:HG22 | 1.77 | 0.48 |
| 26:BA:88:G:H1' | 48:BW:44:PRO:HG2 | 1.93 | 0.48 |
| 26:BA:1474:A:OP1 | 26:BA:1642:G:O2' | 2.30 | 0.48 |
| 26:BA:2259:G:O2' | 26:BA:2507:U:OP1 | 2.13 | 0.48 |
| 26:BA:2500:A:N3 | 26:BA:2502:C:N4 | 2.57 | 0.48 |
| 28:BC:70:LEU:HD21 | 28:BC:160:VAL:HG13 | 1.95 | 0.48 |
| 31:BF:28:ILE:HD11 | 31:BF:116:PHE:HD2 | 1.78 | 0.48 |
| 32:BG:11:ILE:HB | 32:BG:49:GLY:HA2 | 1.95 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 38:BM:31:VAL:O | 38:BM:35:GLN:HG3 | 2.13 | 0.48 |
| 41:BP:34:SER:O | 41:BP:36:ILE:HG13 | 2.13 | 0.48 |
| 50:BY:77:LYS:HB3 | 50:BY:79:PHE:CZ | 2.48 | 0.48 |
| 1:AA:390:C:H1' | 1:AA:405:G:N2 | 2.29 | 0.48 |
| 1:AA:537:G:H2' | 1:AA:538:U:H5' | 1.93 | 0.48 |
| 1:AA:615:G:H5'' | 4:AD:120:LYS:HE2 | 1.96 | 0.48 |
| 1:AA:691:U:O2' | 16:AQ:47:THR:OG1 | 2.29 | 0.48 |
| 1:AA:1131:G:H2' | 1:AA:1132:C:H5' | 1.95 | 0.48 |
| 1:AA:1200:G:H1 | 1:AA:1231:C:H42 | 1.62 | 0.48 |
| 1:AA:1393:C:H4' | 1:AA:1394:G:N3 | 2.28 | 0.48 |
| 2:AB:170:VAL:HG21 | 2:AB:180:LEU:HD21 | 1.95 | 0.48 |
| 7:AG:131:GLU:HB3 | 9:AJ:95:PRO:CB | 2.43 | 0.48 |
| 8:AI:111:VAL:HG13 | 8:AI:113:LEU:CD1 | 2.44 | 0.48 |
| 8:AI:114:LYS:HD3 | 8:AI:119:SER:HA | 1.95 | 0.48 |
| 11:AL:3:LYS:HE2 | 11:AL:20:VAL:HB | 1.94 | 0.48 |
| 14:AP:61:VAL:CG2 | 14:AP:78:LEU:HD21 | 2.43 | 0.48 |
| 16:AQ:149:MET:CE | 50:BY:13:ILE:HD11 | 2.43 | 0.48 |
| 17:AS:54:MET:CG | 17:AS:65:LYS:HG3 | 2.41 | 0.48 |
| 17:AS:81:LYS:N | 17:AS:84:ASP:OD2 | 2.43 | 0.48 |
| 20:AW:19:LEU:HD21 | 20:AW:43:LEU:CD1 | 2.42 | 0.48 |
| 22:AY:16:GLU:OE2 | 22:AY:36:ARG:HB3 | 2.13 | 0.48 |
| 26:BA:1131:C:O2' | 42:BQ:21:LYS:O | 2.30 | 0.48 |
| 26:BA:1712:U:O2' | 26:BA:1713:A:O5' | 2.31 | 0.48 |
| 26:BA:2322:C:OP1 | 31:BF:43:LYS:N | 2.32 | 0.48 |
| 26:BA:2391:A:HO2' | 39:BN:30:SER:HG | 1.53 | 0.48 |
| 30:BE:105:VAL:CG2 | 30:BE:109:GLU:HG3 | 2.43 | 0.48 |
| 30:BE:178:ARG:HG2 | 30:BE:188:LYS:O | 2.13 | 0.48 |
| 31:BF:27:ASP:HA | 31:BF:30:ARG:NH1 | 2.28 | 0.48 |
| 38:BM:156:SER:O | 38:BM:159:ARG:HG3 | 2.14 | 0.48 |
| 40:BO:87:ALA:HA | 40:BO:107:LEU:O | 2.13 | 0.48 |
| 42:BQ:5:LYS:HB2 | 42:BQ:28:SER:O | 2.13 | 0.48 |
| 43:BR:44:ILE:HD11 | 43:BR:53:PRO:HG2 | 1.95 | 0.48 |
| 44:BS:19:MET:HE1 | 44:BS:144:GLU:HG2 | 1.94 | 0.48 |
| 50:BY:28:THR:HG23 | 50:BY:85:ALA:CB | 2.38 | 0.48 |
| 1:AA:58:U:H2' | 1:AA:59:C:C6 | 2.48 | 0.48 |
| 1:AA:110:G:H2' | 1:AA:111:C:O4' | 2.13 | 0.48 |
| 1:AA:296:G:OP2 | 10:AK:23:ARG:NH2 | 2.30 | 0.48 |
| 1:AA:464:G:H5'' | 13:AO:111:MET:CG | 2.41 | 0.48 |
| 1:AA:494:A:H2' | 1:AA:495:U:O4' | 2.13 | 0.48 |
| 1:AA:502:U:O4 | 1:AA:829:U:O2' | 2.18 | 0.48 |
| 1:AA:1224:G:O2' | 1:AA:1226:A:N3 | 2.30 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 1:AA:1276:G:H4' | 14:AP:46:GLY:O | 2.14 | 0.48 |
| 1:AA:1320:G:P | 11:AL:73:ILE:HG13 | 2.53 | 0.48 |
| 2:AB:195:ASP:OD2 | 2:AB:212:LYS:NZ | 2.30 | 0.48 |
| 6:AF:42:ILE:HA | 6:AF:46:ILE:CD1 | 2.42 | 0.48 |
| 7:AG:199:ARG:NH1 | 9:AJ:18:GLU:OE2 | 2.46 | 0.48 |
| 8:AI:70:LYS:O | 8:AI:74:THR:HG23 | 2.13 | 0.48 |
| 9:AJ:46:TYR:HB3 | 9:AJ:69:ILE:CG2 | 2.41 | 0.48 |
| 13:AO:139:LYS:HE3 | 13:AO:140:PRO:O | 2.12 | 0.48 |
| 18:AU:41:ARG:NH2 | 18:AU:44:ARG:HD3 | 2.29 | 0.48 |
| 20:AW:3:ILE:HB | 20:AW:38:LYS:HE3 | 1.96 | 0.48 |
| 20:AW:51:ILE:HG23 | 20:AW:69:LYS:HB2 | 1.96 | 0.48 |
| 23:AH:46:ALA:N | 23:AH:123:LEU:HD13 | 2.27 | 0.48 |
| 26:BA:1096:A:N3 | 26:BA:1246:C:O2' | 2.44 | 0.48 |
| 26:BA:1753:C:H3' | 53:Bb:11:ILE:HB | 1.95 | 0.48 |
| 28:BC:136:ARG:HB3 | 28:BC:148:TRP:CE3 | 2.47 | 0.48 |
| 30:BE:19:LEU:HD11 | 30:BE:139:PRO:HG3 | 1.96 | 0.48 |
| 32:BG:46:VAL:HG12 | 32:BG:51:VAL:CG2 | 2.44 | 0.48 |
| 34:BI:53:GLU:HG2 | 34:BI:160:PRO:O | 2.13 | 0.48 |
| 38:BM:47:ILE:O | 38:BM:51:ARG:HG3 | 2.12 | 0.48 |
| 44:BS:48:PHE:O | 44:BS:52:VAL:HG23 | 2.14 | 0.48 |
| 45:BT:23:GLN:HG3 | 45:BT:63:VAL:HG22 | 1.94 | 0.48 |
| 49:BX:96:ALA:O | 49:BX:100:ILE:HG12 | 2.13 | 0.48 |
| 50:BY:50:ILE:O | 50:BY:54:ILE:HG23 | 2.13 | 0.48 |
| 1:AA:194:A:H2' | 1:AA:195:U:C6 | 2.49 | 0.48 |
| 1:AA:225:A:C2 | 1:AA:227:C:H2' | 2.48 | 0.48 |
| 1:AA:242:G:N7 | 10:AK:112:ARG:NH2 | 2.47 | 0.48 |
| 1:AA:529:U:H5'' | 9:AJ:31:SER:HB2 | 1.94 | 0.48 |
| 1:AA:1063:G:H5' | 24:AM:38:VAL:HG21 | 1.96 | 0.48 |
| 2:AB:117:PRO:O | 2:AB:142:THR:HG22 | 2.13 | 0.48 |
| 2:AB:194:CYS:HB3 | 2:AB:198:ASN:HB2 | 1.96 | 0.48 |
| 5:AE:85:LEU:HD22 | 5:AE:90:ILE:HD11 | 1.95 | 0.48 |
| 6:AF:115:ASN:CG | 6:AF:163:VAL:HG21 | 2.38 | 0.48 |
| 8:AI:12:ASP:HB3 | 8:AI:15:GLU:CD | 2.39 | 0.48 |
| 10:AK:76:ILE:HD11 | 10:AK:119:VAL:HG11 | 1.95 | 0.48 |
| 11:AL:33:PRO:HG2 | 19:AV:6:ASP:HA | 1.95 | 0.48 |
| 22:AY:14:ALA:HB3 | 22:AY:39:GLU:CB | 2.43 | 0.48 |
| 24:AM:1:MET:CE | 24:AM:3:LYS:HE3 | 2.42 | 0.48 |
| 26:BA:391:A:O2' | 26:BA:411:G:N3 | 2.44 | 0.48 |
| 26:BA:2538:G:OP1 | 56:Be:42:LYS:NZ | 2.29 | 0.48 |
| 27:BB:7:G:H2' | 27:BB:8:G:H8 | 1.79 | 0.48 |
| 29:BD:55:VAL:HG13 | 29:BD:65:GLY:O | 2.13 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 50:BY:11:SER:OG | 50:BY:94:ILE:HD13 | 2.13 | 0.48 |
| 52:Ba:31:ASP:N | 52:Ba:31:ASP:OD1 | 2.47 | 0.48 |
| 1:AA:358:G:H5' | 6:AF:24:SER:HA | 1.95 | 0.48 |
| 1:AA:774:U:H2' | 1:AA:775:C:C6 | 2.48 | 0.48 |
| 1:AA:1148:A:H4' | 1:AA:1149:G:C5' | 2.44 | 0.48 |
| 1:AA:1216:A:OP1 | 19:AV:74:ARG:NH2 | 2.39 | 0.48 |
| 1:AA:1238:C:O2' | 11:AL:41:LEU:HD13 | 2.12 | 0.48 |
| 2:AB:171:THR:O | 2:AB:193:LEU:HB3 | 2.14 | 0.48 |
| 3:AC:188:VAL:HG12 | 3:AC:189:LEU:CD2 | 2.43 | 0.48 |
| 5:AE:51:TYR:CE1 | 5:AE:81:ILE:HD11 | 2.49 | 0.48 |
| 5:AE:91:ILE:HD13 | 5:AE:100:ILE:HD13 | 1.95 | 0.48 |
| 6:AF:31:HIS:ND1 | 6:AF:80:GLY:HA3 | 2.28 | 0.48 |
| 7:AG:119:GLU:OE1 | 7:AG:119:GLU:N | 2.46 | 0.48 |
| 8:AI:146:ALA:HB3 | 8:AI:157:CYS:HB3 | 1.95 | 0.48 |
| 16:AQ:53:VAL:O | 16:AQ:57:ARG:HG2 | 2.13 | 0.48 |
| 17:AS:75:PRO:HD2 | 17:AS:102:VAL:CG2 | 2.43 | 0.48 |
| 21:AX:24:GLN:NE2 | 21:AX:37:VAL:HG23 | 2.27 | 0.48 |
| 22:AY:14:ALA:HB3 | 22:AY:39:GLU:HB2 | 1.95 | 0.48 |
| 26:BA:18:G:H5' | 44:BS:4:ILE:HG23 | 1.95 | 0.48 |
| 26:BA:1398:G:N7 | 55:Bd:2:SER:HB3 | 2.29 | 0.48 |
| 42:BQ:8:ILE:HD11 | 42:BQ:25:LYS:CE | 2.44 | 0.48 |
| 44:BS:106:PRO:HA | 44:BS:109:MET:CG | 2.44 | 0.48 |
| 48:BW:46:ARG:NE | 48:BW:50:ILE:HD11 | 2.28 | 0.48 |
| 58:AZ:33:ILE:HD11 | 58:AZ:48:CYS:HB2 | 1.95 | 0.48 |
| 1:AA:125:A:H61 | 1:AA:148:A:H62 | 1.61 | 0.48 |
| 1:AA:625:G:H4' | 12:AN:29:GLU:CD | 2.38 | 0.48 |
| 1:AA:922:A:N1 | 1:AA:1310:A:H5' | 2.29 | 0.48 |
| 1:AA:961:G:H3' | 1:AA:962:A:C8 | 2.48 | 0.48 |
| 1:AA:1167:G:H21 | 18:AU:90:GLY:HA2 | 1.79 | 0.48 |
| 3:AC:179:LYS:HD2 | 3:AC:181:ARG:HH12 | 1.78 | 0.48 |
| 5:AE:19:GLN:HB3 | 5:AE:22:ARG:HB3 | 1.96 | 0.48 |
| 9:AJ:106:THR:HG23 | 9:AJ:122:GLY:O | 2.14 | 0.48 |
| 16:AQ:62:ASP:HB3 | 16:AQ:65:LEU:HG | 1.96 | 0.48 |
| 16:AQ:92:ILE:O | 16:AQ:96:LEU:HG | 2.14 | 0.48 |
| 20:AW:54:ARG:HD3 | 20:AW:56:LYS:HE2 | 1.95 | 0.48 |
| 24:AM:93:ASP:O | 24:AM:94:ILE:HD13 | 2.12 | 0.48 |
| 27:BB:10:G:OP2 | 39:BN:19:THR:OG1 | 2.32 | 0.48 |
| 29:BD:270:LYS:HB3 | 29:BD:296:LEU:HD12 | 1.96 | 0.48 |
| 30:BE:156:ILE:CD1 | 30:BE:212:LEU:HD11 | 2.43 | 0.48 |
| 33:BH:21:LEU:CD2 | 33:BH:97:VAL:HG21 | 2.42 | 0.48 |
| 44:BS:92:LEU:HG | 44:BS:96:GLU:OE2 | 2.14 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 1:AA:52:U:H2' | 1:AA:53:G:C8 | 2.46 | 0.48 |
| 1:AA:407:U:H4' | 5:AE:137:HIS:CE1 | 2.48 | 0.48 |
| 1:AA:560:C:C2 | 5:AE:147:ILE:HG21 | 2.49 | 0.48 |
| 1:AA:984:G:H2' | 1:AA:985:A:C8 | 2.49 | 0.48 |
| 1:AA:985:A:H2' | 1:AA:986:G:O4' | 2.14 | 0.48 |
| 1:AA:1098:G:O4' | 24:AM:39:PRO:HB2 | 2.14 | 0.48 |
| 1:AA:1319:U:O2' | 8:AI:69:GLY:O | 2.30 | 0.48 |
| 3:AC:180:VAL:HG12 | 3:AC:182:ILE:HD11 | 1.96 | 0.48 |
| 6:AF:116:LYS:HB2 | 6:AF:157:ILE:HD11 | 1.96 | 0.48 |
| 6:AF:189:MET:SD | 6:AF:205:SER:HB3 | 2.54 | 0.48 |
| 7:AG:49:GLU:OE2 | 7:AG:79:ARG:HG3 | 2.14 | 0.48 |
| 8:AI:71:LYS:NZ | 8:AI:75:LEU:HD11 | 2.28 | 0.48 |
| 9:AJ:15:LYS:NZ | 9:AJ:71:LYS:HB2 | 2.29 | 0.48 |
| 19:AV:63:TYR:HE1 | 19:AV:125:ARG:HG3 | 1.78 | 0.48 |
| 21:AX:14:ARG:HH21 | 21:AX:61:LEU:HD21 | 1.79 | 0.48 |
| 24:AM:16:LEU:HD21 | 24:AM:71:ARG:HG3 | 1.96 | 0.48 |
| 26:BA:1546:G:OP1 | 41:BP:64:ARG:NH1 | 2.46 | 0.48 |
| 26:BA:1699:A:H5'' | 41:BP:55:VAL:HG22 | 1.95 | 0.48 |
| 37:BL:35:CYS:HB2 | 37:BL:49:TYR:CE2 | 2.49 | 0.48 |
| 39:BN:43:ASN:HA | 39:BN:76:THR:O | 2.13 | 0.48 |
| 39:BN:152:ARG:HB3 | 39:BN:154:GLU:OE1 | 2.12 | 0.48 |
| 40:BO:21:ILE:HD13 | 40:BO:44:GLU:CG | 2.44 | 0.48 |
| 45:BT:79:ILE:O | 48:BW:30:ARG:HD3 | 2.13 | 0.48 |
| 49:BX:130:ARG:O | 49:BX:136:GLY:HA3 | 2.13 | 0.48 |
| 52:Ba:126:ILE:CG1 | 52:Ba:151:VAL:HG22 | 2.44 | 0.48 |
| 57:Bf:31:SER:HB3 | 57:Bf:34:THR:HG23 | 1.95 | 0.48 |
| 58:AZ:10:THR:OG1 | 58:AZ:46:TYR:HB2 | 2.13 | 0.48 |
| 1:AA:12:U:H2' | 1:AA:13:C:C6 | 2.48 | 0.48 |
| 1:AA:738:C:O2' | 1:AA:739:G:H5' | 2.14 | 0.48 |
| 1:AA:742:U:O2' | 28:BC:238:ARG:O | 2.23 | 0.48 |
| 1:AA:1241:U:H2' | 1:AA:1242:C:C6 | 2.48 | 0.48 |
| 3:AC:181:ARG:O | 3:AC:182:ILE:HD13 | 2.14 | 0.48 |
| 4:AD:53:GLY:HA2 | 4:AD:58:ASP:OD1 | 2.13 | 0.48 |
| 6:AF:6:ARG:N | 6:AF:21:LYS:O | 2.46 | 0.48 |
| 6:AF:143:GLY:HA3 | 6:AF:147:TYR:CD2 | 2.49 | 0.48 |
| 11:AL:39:PRO:O | 11:AL:41:LEU:HD23 | 2.13 | 0.48 |
| 11:AL:87:ILE:HG22 | 11:AL:96:ILE:HD12 | 1.95 | 0.48 |
| 17:AS:27:LEU:HD21 | 17:AS:91:CYS:CB | 2.44 | 0.48 |
| 23:AH:38:VAL:CG1 | 23:AH:42:ILE:HD12 | 2.43 | 0.48 |
| 26:BA:143:G:O3' | 54:Bc:44:ARG:HA | 2.13 | 0.48 |
| 26:BA:2249:C:H5'' | 28:BC:1:MET:HG2 | 1.95 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 26:BA:2344:U:N3 | 39:BN:107:LEU:HD21 | 2.28 | 0.48 |
| 29:BD:133:ILE:HG12 | 29:BD:138:VAL:HG11 | 1.96 | 0.48 |
| 46:BU:28:MET:HE2 | 46:BU:70:VAL:HG13 | 1.96 | 0.48 |
| 1:AA:197:C:H4' | 1:AA:198:G:C5' | 2.42 | 0.48 |
| 1:AA:212:G:H2' | 1:AA:213:G:O4' | 2.13 | 0.48 |
| 1:AA:497:G:H2' | 1:AA:498:U:O4' | 2.14 | 0.48 |
| 1:AA:935:G:H2' | 1:AA:936:A:C8 | 2.49 | 0.48 |
| 1:AA:1361:C:H2' | 1:AA:1362:C:C6 | 2.49 | 0.48 |
| 3:AC:48:SER:N | 3:AC:83:GLN:O | 2.29 | 0.48 |
| 3:AC:117:LEU:HD22 | 3:AC:146:PHE:CB | 2.39 | 0.48 |
| 7:AG:132:VAL:HG21 | 7:AG:195:VAL:CG2 | 2.44 | 0.48 |
| 9:AJ:11:LEU:HB3 | 9:AJ:72:CYS:O | 2.14 | 0.48 |
| 10:AK:125:GLU:OE1 | 10:AK:125:GLU:N | 2.45 | 0.48 |
| 13:AO:52:LYS:HD3 | 13:AO:93:ASP:HA | 1.95 | 0.48 |
| 14:AP:147:LYS:O | 14:AP:151:ARG:NH2 | 2.44 | 0.48 |
| 16:AQ:19:ARG:NH2 | 16:AQ:24:GLU:OE1 | 2.47 | 0.48 |
| 16:AQ:69:LYS:HD2 | 16:AQ:73:THR:HG22 | 1.95 | 0.48 |
| 22:AY:36:ARG:HG2 | 22:AY:38:LEU:HD21 | 1.96 | 0.48 |
| 24:AM:42:THR:CG2 | 24:AM:67:ARG:HB3 | 2.43 | 0.48 |
| 26:BA:1312:U:C4 | 52:Ba:89:VAL:HG12 | 2.49 | 0.48 |
| 26:BA:1728:A:H2' | 36:BK:44:MET:HE2 | 1.96 | 0.48 |
| 26:BA:2100:U:O2' | 57:Bf:48:ASN:ND2 | 2.33 | 0.48 |
| 30:BE:41:ALA:HA | 30:BE:44:LEU:CD1 | 2.43 | 0.48 |
| 30:BE:132:HIS:ND1 | 30:BE:167:ASP:OD2 | 2.37 | 0.48 |
| 31:BF:6:VAL:HB | 31:BF:69:GLN:HE22 | 1.79 | 0.48 |
| 31:BF:10:ILE:HB | 31:BF:121:THR:CG2 | 2.42 | 0.48 |
| 31:BF:83:LEU:O | 31:BF:164:VAL:HA | 2.14 | 0.48 |
| 41:BP:105:ARG:O | 41:BP:108:GLU:HG2 | 2.14 | 0.48 |
| 47:BV:50:GLU:HG3 | 47:BV:59:LEU:HD12 | 1.96 | 0.48 |
| 53:Bb:30:VAL:O | 53:Bb:34:GLU:HG2 | 2.14 | 0.48 |
| 1:AA:67:C:H2' | 1:AA:68:U:O4' | 2.13 | 0.48 |
| 1:AA:531:U:OP2 | 9:AJ:32:LYS:NZ | 2.44 | 0.48 |
| 1:AA:1030:U:H2' | 1:AA:1032:A:OP2 | 2.14 | 0.48 |
| 1:AA:1210:C:H2' | 1:AA:1211:C:C6 | 2.48 | 0.48 |
| 1:AA:1275:C:H2' | 1:AA:1276:G:C8 | 2.49 | 0.48 |
| 2:AB:72:ILE:CG2 | 2:AB:93:LEU:HD13 | 2.41 | 0.48 |
| 3:AC:152:LYS:NZ | 3:AC:154:SER:O | 2.45 | 0.48 |
| 4:AD:25:VAL:HG12 | 4:AD:29:ARG:HH11 | 1.79 | 0.48 |
| 6:AF:68:ASP:OD2 | 6:AF:116:LYS:HD2 | 2.13 | 0.48 |
| 6:AF:174:ALA:HB3 | 6:AF:204:ILE:HD12 | 1.95 | 0.48 |
| 8:AI:48:LYS:HE3 | 8:AI:129:GLN:HG3 | 1.96 | 0.48 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 10:AK:42:LYS:HE3 | 10:AK:56:LEU:CD1 | 2.37 | 0.48 |
| 12:AN:22:VAL:HG11 | 12:AN:63:LEU:HD21 | 1.96 | 0.48 |
| 24:AM:6:ILE:O | 24:AM:72:LEU:HD12 | 2.13 | 0.48 |
| 26:BA:1700:G:OP1 | 41:BP:18:GLY:N | 2.46 | 0.48 |
| 26:BA:2444:A:N9 | 57:Bf:54:LYS:HG2 | 2.28 | 0.48 |
| 29:BD:225:LYS:O | 29:BD:229:GLN:HG2 | 2.14 | 0.48 |
| 32:BG:84:PHE:CE1 | 32:BG:142:LYS:HD2 | 2.49 | 0.48 |
| 39:BN:121:LYS:HD3 | 39:BN:143:ILE:HD11 | 1.96 | 0.48 |
| 41:BP:123:LEU:HD21 | 41:BP:147:LEU:CD1 | 2.44 | 0.48 |
| 45:BT:6:TYR:HE2 | 48:BW:36:SER:HB2 | 1.78 | 0.48 |
| 45:BT:36:LEU:O | 45:BT:40:GLU:HG3 | 2.14 | 0.48 |
| 49:BX:122:ARG:NH2 | 49:BX:153:ARG:OXT | 2.47 | 0.48 |
| 1:AA:175:U:O4 | 17:AS:1:MET:HE3 | 2.14 | 0.47 |
| 1:AA:175:U:N3 | 17:AS:28:PRO:HB3 | 2.29 | 0.47 |
| 1:AA:1020:A:OP2 | 7:AG:89:LYS:NZ | 2.46 | 0.47 |
| 1:AA:1191:U:H2' | 1:AA:1192:G:O4' | 2.13 | 0.47 |
| 1:AA:1208:C:H5' | 1:AA:1230:U:C5' | 2.44 | 0.47 |
| 1:AA:1386:C:H2' | 1:AA:1387:G:C8 | 2.49 | 0.47 |
| 2:AB:193:LEU:HD21 | 2:AB:209:THR:HG23 | 1.95 | 0.47 |
| 4:AD:46:ARG:O | 4:AD:70:ILE:HG12 | 2.14 | 0.47 |
| 7:AG:8:VAL:O | 7:AG:10:LYS:NZ | 2.36 | 0.47 |
| 7:AG:37:GLU:HG3 | 7:AG:39:GLN:H | 1.79 | 0.47 |
| 8:AI:21:LEU:HD23 | 22:AY:60:ILE:HG13 | 1.97 | 0.47 |
| 9:AJ:119:LYS:HB3 | 9:AJ:121:ILE:HG22 | 1.96 | 0.47 |
| 11:AL:19:THR:O | 11:AL:64:ASN:N | 2.28 | 0.47 |
| 19:AV:12:MET:O | 19:AV:16:LEU:HG | 2.14 | 0.47 |
| 19:AV:21:LYS:HA | 19:AV:52:TYR:CE2 | 2.48 | 0.47 |
| 21:AX:12:PHE:O | 21:AX:60:VAL:HG13 | 2.14 | 0.47 |
| 23:AH:85:PRO:HB2 | 23:AH:87:LEU:O | 2.14 | 0.47 |
| 26:BA:914:C:O2' | 26:BA:2257:C:OP1 | 2.32 | 0.47 |
| 26:BA:1099:U:O4 | 35:BJ:16:ARG:HA | 2.14 | 0.47 |
| 29:BD:203:ALA:CB | 29:BD:309:ILE:HD13 | 2.44 | 0.47 |
| 30:BE:55:MET:HG2 | 30:BE:90:ARG:HD3 | 1.96 | 0.47 |
| 31:BF:1:MET:HG2 | 31:BF:127:PRO:HB2 | 1.96 | 0.47 |
| 39:BN:88:LEU:HD13 | 39:BN:169:ILE:HD11 | 1.96 | 0.47 |
| 39:BN:121:LYS:HD3 | 39:BN:143:ILE:CD1 | 2.44 | 0.47 |
| 39:BN:138:PRO:HG2 | 39:BN:143:ILE:HD11 | 1.96 | 0.47 |
| 46:BU:16:ARG:HG2 | 46:BU:17:TYR:CD1 | 2.48 | 0.47 |
| 47:BV:58:GLN:HA | 47:BV:61:LYS:HD2 | 1.96 | 0.47 |
| 53:Bb:6:THR:HG23 | 53:Bb:9:GLY:H | 1.79 | 0.47 |
| 1:AA:177:C:C2 | 1:AA:178:U:H1' | 2.49 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:192:G:H2' | 1:AA:193:G:H8 | 1.79 | 0.47 |
| 1:AA:685:A:OP1 | 16:AQ:3:LYS:NZ | 2.29 | 0.47 |
| 1:AA:778:G:H2' | 1:AA:779:C:O4' | 2.13 | 0.47 |
| 1:AA:1063:G:O2' | 1:AA:1064:U:H2' | 2.14 | 0.47 |
| 2:AB:194:CYS:SG | 2:AB:206:VAL:HG13 | 2.54 | 0.47 |
| 4:AD:122:THR:HB | 4:AD:186:ARG:NH1 | 2.28 | 0.47 |
| 6:AF:75:LEU:H | 6:AF:75:LEU:HD23 | 1.80 | 0.47 |
| 7:AG:101:ILE:HG22 | 7:AG:105:LYS:NZ | 2.28 | 0.47 |
| 13:AO:114:ILE:CG2 | 13:AO:117:VAL:HB | 2.44 | 0.47 |
| 17:AS:49:ILE:CG2 | 17:AS:70:ILE:HB | 2.45 | 0.47 |
| 26:BA:312:C:OP1 | 46:BU:57:ARG:NE | 2.40 | 0.47 |
| 26:BA:1109:G:OP1 | 35:BJ:51:ARG:NH2 | 2.47 | 0.47 |
| 27:BB:16:A:OP2 | 27:BB:70:G:N2 | 2.31 | 0.47 |
| 32:BG:22:ASP:HB3 | 32:BG:38:TRP:HB2 | 1.96 | 0.47 |
| 38:BM:45:THR:OG1 | 38:BM:131:GLU:OE2 | 2.15 | 0.47 |
| 38:BM:136:ASP:OD2 | 38:BM:139:HIS:HB2 | 2.14 | 0.47 |
| 40:BO:68:LEU:HD13 | 40:BO:84:VAL:HG13 | 1.96 | 0.47 |
| 41:BP:77:GLY:O | 41:BP:81:ARG:HG3 | 2.13 | 0.47 |
| 49:BX:21:MET:HE3 | 49:BX:26:LEU:HB2 | 1.97 | 0.47 |
| 49:BX:127:GLY:O | 49:BX:137:VAL:HG12 | 2.14 | 0.47 |
| 52:Ba:101:HIS:HB2 | 52:Ba:107:ASP:OD1 | 2.13 | 0.47 |
| 52:Ba:126:ILE:CD1 | 52:Ba:151:VAL:HG22 | 2.43 | 0.47 |
| 58:AZ:26:CYS:N | 58:AZ:31:ALA:O | 2.47 | 0.47 |
| 1:AA:618:C:H2' | 1:AA:619:U:O4' | 2.15 | 0.47 |
| 1:AA:997:G:H2' | 1:AA:998:C:C6 | 2.48 | 0.47 |
| 1:AA:997:G:H5'' | 3:AC:179:LYS:HE3 | 1.95 | 0.47 |
| 3:AC:85:VAL:HG21 | 3:AC:91:ASN:HB2 | 1.96 | 0.47 |
| 4:AD:51:THR:OG1 | 4:AD:54:GLU:HG2 | 2.15 | 0.47 |
| 13:AO:38:PRO:HA | 13:AO:77:ARG:HD3 | 1.96 | 0.47 |
| 14:AP:59:ALA:HB1 | 14:AP:78:LEU:HD23 | 1.96 | 0.47 |
| 21:AX:18:ASN:HB3 | 21:AX:41:ARG:NH1 | 2.29 | 0.47 |
| 23:AH:55:GLY:HA3 | 23:AH:63:MET:SD | 2.54 | 0.47 |
| 29:BD:278:VAL:HG12 | 29:BD:330:ILE:HG21 | 1.96 | 0.47 |
| 31:BF:40:CYS:O | 31:BF:56:PRO:HA | 2.14 | 0.47 |
| 37:BL:82:GLU:OE1 | 37:BL:84:LEU:HG | 2.14 | 0.47 |
| 52:Ba:41:ARG:HB2 | 52:Ba:94:PRO:HD2 | 1.96 | 0.47 |
| 52:Ba:50:GLN:OE1 | 52:Ba:52:LYS:HE3 | 2.14 | 0.47 |
| 1:AA:283:G:H5' | 17:AS:59:LYS:HD2 | 1.95 | 0.47 |
| 1:AA:805:A:H2' | 1:AA:806:G:H5' | 1.96 | 0.47 |
| 1:AA:1186:A:H5'' | 8:AI:154:VAL:H | 1.79 | 0.47 |
| 1:AA:1318:G:H2' | 1:AA:1319:U:O4' | 2.14 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 2:AB:130:GLN:CG | 2:AB:146:LEU:HD12 | 2.44 | 0.47 |
| 3:AC:126:LEU:HB2 | 3:AC:183:ILE:CG2 | 2.44 | 0.47 |
| 7:AG:161:VAL:HG13 | 7:AG:188:THR:HG21 | 1.95 | 0.47 |
| 13:AO:27:ARG:HG3 | 13:AO:32:LEU:HB2 | 1.97 | 0.47 |
| 19:AV:23:ASP:HB3 | 19:AV:25:LYS:NZ | 2.29 | 0.47 |
| 19:AV:69:GLY:O | 19:AV:73:LEU:HG | 2.14 | 0.47 |
| 20:AW:23:VAL:O | 20:AW:63:GLU:HA | 2.13 | 0.47 |
| 24:AM:40:LEU:HB2 | 24:AM:70:LYS:C | 2.40 | 0.47 |
| 25:AT:40:THR:HG22 | 25:AT:42:GLU:OE2 | 2.14 | 0.47 |
| 29:BD:87:ALA:O | 29:BD:98:ILE:HG22 | 2.14 | 0.47 |
| 29:BD:177:GLU:O | 29:BD:181:THR:HG23 | 2.15 | 0.47 |
| 29:BD:201:THR:HG22 | 29:BD:311:LEU:CD2 | 2.44 | 0.47 |
| 29:BD:274:ASP:OD2 | 29:BD:276:GLU:HB2 | 2.14 | 0.47 |
| 37:BL:96:LEU:HB2 | 37:BL:118:SER:HB2 | 1.96 | 0.47 |
| 44:BS:48:PHE:CZ | 44:BS:59:VAL:HG12 | 2.49 | 0.47 |
| 44:BS:82:ILE:O | 44:BS:86:LYS:HG3 | 2.14 | 0.47 |
| 56:Be:36:TYR:CE2 | 56:Be:38:THR:HB | 2.49 | 0.47 |
| 1:AA:70:U:H4' | 1:AA:71:G:O4' | 2.13 | 0.47 |
| 1:AA:248:U:O2' | 10:AK:8:ARG:NH1 | 2.48 | 0.47 |
| 1:AA:292:C:H2' | 1:AA:293:G:C8 | 2.50 | 0.47 |
| 1:AA:632:G:O2' | 1:AA:633:U:H5' | 2.13 | 0.47 |
| 1:AA:903:U:O2 | 1:AA:906:U:H5'' | 2.15 | 0.47 |
| 1:AA:938:G:H2' | 1:AA:940:C:H41 | 1.79 | 0.47 |
| 1:AA:1089:G:H2' | 1:AA:1090:G:O4' | 2.15 | 0.47 |
| 1:AA:1190:A:H5'' | 19:AV:41:LYS:NZ | 2.30 | 0.47 |
| 1:AA:1272:C:H2' | 1:AA:1273:C:C6 | 2.50 | 0.47 |
| 1:AA:1327:U:O2' | 22:AY:26:MET:O | 2.22 | 0.47 |
| 3:AC:44:ILE:HB | 3:AC:80:ILE:HG13 | 1.96 | 0.47 |
| 3:AC:92:ALA:HB2 | 3:AC:182:ILE:HG13 | 1.94 | 0.47 |
| 6:AF:176:VAL:HA | 6:AF:216:TYR:O | 2.15 | 0.47 |
| 8:AI:46:PHE:HA | 8:AI:129:GLN:HE22 | 1.79 | 0.47 |
| 10:AK:32:ALA:HB3 | 10:AK:54:ARG:CD | 2.41 | 0.47 |
| 19:AV:40:HIS:CE1 | 19:AV:41:LYS:HG2 | 2.50 | 0.47 |
| 25:AT:56:ARG:HD2 | 25:AT:60:ARG:HD3 | 1.97 | 0.47 |
| 26:BA:1045:G:N7 | 34:BI:14:GLN:NE2 | 2.63 | 0.47 |
| 27:BB:18:C:H2' | 27:BB:19:G:O4' | 2.14 | 0.47 |
| 36:BK:12:LEU:HD23 | 36:BK:31:ILE:HD11 | 1.97 | 0.47 |
| 39:BN:135:GLU:N | 39:BN:135:GLU:OE1 | 2.46 | 0.47 |
| 39:BN:140:ASP:O | 39:BN:144:ARG:HG3 | 2.13 | 0.47 |
| 49:BX:143:GLU:O | 49:BX:146:ASN:ND2 | 2.44 | 0.47 |
| 54:Bc:2:SER:OG | 54:Bc:3:LYS:N | 2.45 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 1:AA:12:U:H2' | 1:AA:13:C:H6 | 1.80 | 0.47 |
| 1:AA:415:U:H2' | 1:AA:416:G:C8 | 2.49 | 0.47 |
| 1:AA:601:G:H2' | 1:AA:602:G:C8 | 2.49 | 0.47 |
| 1:AA:1063:G:H5' | 24:AM:38:VAL:CG2 | 2.45 | 0.47 |
| 1:AA:1321:A:C4' | 8:AI:59:ASN:HB2 | 2.38 | 0.47 |
| 1:AA:1355:A:H1' | 26:BA:1933:A:N1 | 2.29 | 0.47 |
| 1:AA:1385:A:OP1 | 10:AK:42:LYS:NZ | 2.34 | 0.47 |
| 3:AC:131:VAL:CG2 | 3:AC:179:LYS:HB2 | 2.44 | 0.47 |
| 3:AC:162:VAL:HG12 | 3:AC:183:ILE:CG1 | 2.45 | 0.47 |
| 5:AE:56:ARG:CD | 7:AG:147:PRO:HG3 | 2.45 | 0.47 |
| 6:AF:143:GLY:HA3 | 6:AF:147:TYR:CE2 | 2.50 | 0.47 |
| 14:AP:133:LEU:HA | 14:AP:136:GLU:HG3 | 1.97 | 0.47 |
| 15:AR:35:HIS:O | 15:AR:39:GLU:HG3 | 2.15 | 0.47 |
| 16:AQ:87:ASP:N | 16:AQ:87:ASP:OD1 | 2.44 | 0.47 |
| 19:AV:18:GLY:O | 19:AV:22:GLU:HG3 | 2.14 | 0.47 |
| 20:AW:14:LEU:HD22 | 20:AW:16:ARG:CG | 2.45 | 0.47 |
| 26:BA:2461:A:N6 | 26:BA:2512:C:C2 | 2.83 | 0.47 |
| 29:BD:91:ASP:OD1 | 29:BD:93:THR:N | 2.41 | 0.47 |
| 33:BH:23:LEU:CD2 | 33:BH:105:ALA:HB2 | 2.42 | 0.47 |
| 33:BH:28:GLY:HA3 | 33:BH:101:ASP:O | 2.14 | 0.47 |
| 39:BN:152:ARG:HB3 | 39:BN:154:GLU:CD | 2.40 | 0.47 |
| 44:BS:113:HIS:HB3 | 44:BS:146:ILE:HB | 1.97 | 0.47 |
| 46:BU:30:ALA:HB1 | 46:BU:101:VAL:O | 2.14 | 0.47 |
| 50:BY:54:ILE:HA | 50:BY:57:THR:HG23 | 1.96 | 0.47 |
| 56:Be:18:MET:HB2 | 56:Be:36:TYR:CE2 | 2.50 | 0.47 |
| 1:AA:120:C:H3' | 1:AA:120:C:OP2 | 2.14 | 0.47 |
| 1:AA:170:U:N3 | 1:AA:171:G:N3 | 2.62 | 0.47 |
| 1:AA:387:U:H5'' | 1:AA:388:G:H1' | 1.97 | 0.47 |
| 1:AA:576:G:H2' | 1:AA:577:G:O4' | 2.15 | 0.47 |
| 1:AA:720:C:H2' | 1:AA:721:G:O4' | 2.14 | 0.47 |
| 1:AA:798:G:H2' | 1:AA:799:U:O4' | 2.15 | 0.47 |
| 1:AA:800:G:H2' | 1:AA:801:C:C6 | 2.50 | 0.47 |
| 1:AA:941:A:H2' | 1:AA:942:G:C8 | 2.50 | 0.47 |
| 1:AA:1066:A:H5' | 1:AA:1227:A:O2' | 2.15 | 0.47 |
| 1:AA:1190:A:H5' | 19:AV:41:LYS:HD2 | 1.95 | 0.47 |
| 1:AA:1209:C:OP1 | 19:AV:69:GLY:HA3 | 2.14 | 0.47 |
| 4:AD:117:ILE:HD11 | 4:AD:154:PHE:CE1 | 2.50 | 0.47 |
| 5:AE:163:TYR:O | 5:AE:169:LYS:HB2 | 2.15 | 0.47 |
| 5:AE:173:HIS:HD2 | 5:AE:174:PRO:HD2 | 1.79 | 0.47 |
| 6:AF:31:HIS:NE2 | 6:AF:137:ASP:OD2 | 2.46 | 0.47 |
| 6:AF:124:THR:N | 6:AF:132:GLN:O | 2.48 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 9:AJ:52:PHE:CE1 | 9:AJ:59:GLY:HA3 | 2.48 | 0.47 |
| 10:AK:63:VAL:HA | 10:AK:121:ALA:O | 2.14 | 0.47 |
| 13:AO:83:CYS:SG | 13:AO:88:ALA:HB1 | 2.55 | 0.47 |
| 13:AO:94:GLU:HG2 | 13:AO:95:HIS:ND1 | 2.29 | 0.47 |
| 14:AP:25:ARG:HA | 14:AP:30:ASP:HA | 1.96 | 0.47 |
| 18:AU:53:GLY:O | 18:AU:57:VAL:HG23 | 2.14 | 0.47 |
| 18:AU:76:ILE:HG12 | 18:AU:102:MET:SD | 2.54 | 0.47 |
| 18:AU:77:LEU:HB2 | 18:AU:80:MET:CB | 2.45 | 0.47 |
| 19:AV:106:GLU:HG2 | 19:AV:111:LEU:O | 2.15 | 0.47 |
| 20:AW:52:ILE:CD1 | 20:AW:55:ILE:HD11 | 2.45 | 0.47 |
| 24:AM:12:SER:O | 24:AM:94:ILE:HD11 | 2.14 | 0.47 |
| 25:AT:26:LYS:CE | 25:AT:62:LYS:HD2 | 2.45 | 0.47 |
| 26:BA:805:C:H5 | 53:Bb:4:LYS:HE3 | 1.80 | 0.47 |
| 26:BA:1105:C:OP1 | 32:BG:59:ARG:NH1 | 2.47 | 0.47 |
| 28:BC:90:LYS:HE2 | 28:BC:93:ASN:ND2 | 2.28 | 0.47 |
| 29:BD:72:VAL:HG12 | 29:BD:269:LEU:HD12 | 1.97 | 0.47 |
| 29:BD:142:ARG:HD2 | 29:BD:164:GLU:OE1 | 2.15 | 0.47 |
| 30:BE:147:GLU:HB3 | 30:BE:203:THR:CG2 | 2.42 | 0.47 |
| 30:BE:206:LEU:O | 30:BE:210:ARG:HB2 | 2.14 | 0.47 |
| 31:BF:65:GLY:O | 31:BF:69:GLN:HG2 | 2.15 | 0.47 |
| 34:BI:26:VAL:HG22 | 34:BI:94:VAL:HG11 | 1.95 | 0.47 |
| 44:BS:17:LYS:O | 44:BS:98:ASN:ND2 | 2.42 | 0.47 |
| 45:BT:7:PRO:HD2 | 48:BW:33:ALA:HB2 | 1.95 | 0.47 |
| 47:BV:57:ILE:HG22 | 47:BV:61:LYS:HE3 | 1.95 | 0.47 |
| 49:BX:38:ASN:OD1 | 49:BX:41:TYR:HB2 | 2.14 | 0.47 |
| 49:BX:93:GLN:O | 49:BX:97:GLU:HG3 | 2.15 | 0.47 |
| 50:BY:96:ALA:HA | 50:BY:99:ALA:OXT | 2.15 | 0.47 |
| 58:AZ:10:THR:CG2 | 58:AZ:33:ILE:HG23 | 2.44 | 0.47 |
| 1:AA:40:U:H4' | 1:AA:289:G:C8 | 2.50 | 0.47 |
| 1:AA:359:C:OP1 | 6:AF:25:THR:HB | 2.14 | 0.47 |
| 1:AA:600:C:N3 | 1:AA:688:G:N2 | 2.63 | 0.47 |
| 1:AA:804:C:OP2 | 1:AA:814:G:N1 | 2.29 | 0.47 |
| 1:AA:981:U:H2' | 1:AA:982:G:C8 | 2.50 | 0.47 |
| 1:AA:1063:G:H1' | 1:AA:1064:U:C5 | 2.50 | 0.47 |
| 3:AC:152:LYS:NZ | 3:AC:157:PRO:HD2 | 2.30 | 0.47 |
| 6:AF:2:THR:O | 6:AF:25:THR:HG21 | 2.15 | 0.47 |
| 7:AG:49:GLU:OE1 | 7:AG:78:ASN:HA | 2.15 | 0.47 |
| 8:AI:46:PHE:HB3 | 8:AI:129:GLN:OE1 | 2.14 | 0.47 |
| 19:AV:101:THR:O | 19:AV:105:LEU:HG | 2.15 | 0.47 |
| 24:AM:16:LEU:HD21 | 24:AM:71:ARG:CG | 2.45 | 0.47 |
| 26:BA:805:C:C5 | 53:Bb:4:LYS:HE3 | 2.49 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 26:BA:1676:G:N2 | 54:Bc:5:THR:O | 2.47 | 0.47 |
| 26:BA:2525:U:H5' | 29:BD:249:ARG:HD3 | 1.97 | 0.47 |
| 27:BB:31:C:O3' | 31:BF:128:GLY:HA2 | 2.14 | 0.47 |
| 31:BF:142:PRO:O | 31:BF:146:ARG:HG3 | 2.15 | 0.47 |
| 31:BF:150:ASP:HA | 31:BF:153:ILE:HG12 | 1.97 | 0.47 |
| 33:BH:18:LEU:O | 33:BH:22:GLU:HG2 | 2.15 | 0.47 |
| 36:BK:92:ASP:CG | 47:BV:24:LYS:HE3 | 2.40 | 0.47 |
| 37:BL:97:GLU:OE2 | 37:BL:139:ALA:HB3 | 2.15 | 0.47 |
| 44:BS:68:SER:CB | 44:BS:79:LYS:HD3 | 2.45 | 0.47 |
| 50:BY:80:THR:HG21 | 53:Bb:48:ARG:HD3 | 1.97 | 0.47 |
| 1:AA:78:U:O2' | 1:AA:148:A:N1 | 2.32 | 0.47 |
| 1:AA:234:U:OP2 | 10:AK:8:ARG:HG3 | 2.15 | 0.47 |
| 1:AA:356:A:H2 | 6:AF:20:ASN:HD21 | 1.63 | 0.47 |
| 1:AA:939:A:H61 | 1:AA:983:A:H2' | 1.78 | 0.47 |
| 1:AA:1090:G:H2' | 1:AA:1091:G:C8 | 2.49 | 0.47 |
| 1:AA:1128:G:H5' | 8:AI:43:ARG:NH2 | 2.29 | 0.47 |
| 1:AA:1238:C:H5' | 8:AI:72:GLN:OE1 | 2.15 | 0.47 |
| 5:AE:137:HIS:ND1 | 5:AE:166:SER:HB2 | 2.29 | 0.47 |
| 6:AF:107:HIS:CE1 | 6:AF:233:ILE:HD13 | 2.50 | 0.47 |
| 7:AG:132:VAL:HG23 | 7:AG:143:LEU:HB2 | 1.97 | 0.47 |
| 13:AO:103:ILE:HA | 13:AO:120:LYS:HD3 | 1.97 | 0.47 |
| 16:AQ:2:ALA:CB | 16:AQ:7:LYS:HE3 | 2.45 | 0.47 |
| 26:BA:293:A:O2' | 26:BA:294:A:O4' | 2.12 | 0.47 |
| 26:BA:692:C:O2' | 26:BA:736:G:O4' | 2.32 | 0.47 |
| 26:BA:1230:G:N3 | 35:BJ:83:MET:HE2 | 2.29 | 0.47 |
| 26:BA:2567:C:OP1 | 29:BD:2:ALA:HB2 | 2.15 | 0.47 |
| 36:BK:20:CYS:SG | 36:BK:97:ILE:HG12 | 2.55 | 0.47 |
| 36:BK:73:VAL:HB | 36:BK:95:MET:HG2 | 1.97 | 0.47 |
| 36:BK:103:ILE:CD1 | 36:BK:123:LYS:HG3 | 2.44 | 0.47 |
| 39:BN:50:ALA:O | 39:BN:57:ILE:HG12 | 2.15 | 0.47 |
| 41:BP:27:GLU:N | 41:BP:27:GLU:OE1 | 2.47 | 0.47 |
| 44:BS:109:MET:HB3 | 44:BS:147:LEU:HB3 | 1.96 | 0.47 |
| 48:BW:9:ILE:HD13 | 48:BW:57:ILE:HG12 | 1.96 | 0.47 |
| 1:AA:30:C:H2' | 1:AA:31:U:O4' | 2.14 | 0.47 |
| 1:AA:40:U:H4' | 1:AA:289:G:H8 | 1.80 | 0.47 |
| 1:AA:504:U:H2' | 1:AA:505:U:H5' | 1.97 | 0.47 |
| 1:AA:566:U:H5'' | 6:AF:14:LYS:NZ | 2.29 | 0.47 |
| 1:AA:776:A:H61 | 1:AA:796:U:H3 | 1.63 | 0.47 |
| 2:AB:214:ARG:HB2 | 2:AB:245:THR:HG21 | 1.95 | 0.47 |
| 7:AG:54:ASP:O | 7:AG:72:ALA:HA | 2.14 | 0.47 |
| 7:AG:151:GLY:O | 7:AG:169:ASP:HA | 2.15 | 0.47 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 7:AG:202:VAL:HG22 | 9:AJ:68:ARG:CZ | 2.44 | 0.47 |
| 12:AN:80:GLY:HA2 | 12:AN:114:PRO:CG | 2.41 | 0.47 |
| 13:AO:114:ILE:HG22 | 13:AO:117:VAL:HB | 1.97 | 0.47 |
| 14:AP:41:LEU:HD13 | 14:AP:82:ILE:HD13 | 1.97 | 0.47 |
| 14:AP:95:ASN:OD1 | 14:AP:107:HIS:HA | 2.15 | 0.47 |
| 23:AH:101:ILE:HA | 23:AH:105:ILE:HD12 | 1.97 | 0.47 |
| 26:BA:329:U:O2 | 26:BA:333:A:N7 | 2.48 | 0.47 |
| 26:BA:2321:A:C2 | 31:BF:48:ALA:HB3 | 2.50 | 0.47 |
| 26:BA:2657:G:OP2 | 29:BD:316:ARG:NH2 | 2.29 | 0.47 |
| 29:BD:90:GLU:OE1 | 29:BD:95:GLU:HG2 | 2.15 | 0.47 |
| 32:BG:73:ILE:O | 32:BG:77:VAL:HG13 | 2.14 | 0.47 |
| 35:BJ:85:PRO:HG3 | 35:BJ:90:ARG:NH2 | 2.30 | 0.47 |
| 41:BP:35:ALA:HB2 | 41:BP:44:LEU:CD1 | 2.44 | 0.47 |
| 46:BU:2:ILE:CG2 | 46:BU:12:GLN:HG2 | 2.45 | 0.47 |
| 52:Ba:113:ILE:HD12 | 52:Ba:116:LEU:HD12 | 1.97 | 0.47 |
| 1:AA:211:U:OP1 | 6:AF:33:GLN:HB3 | 2.14 | 0.46 |
| 1:AA:234:U:H5'' | 10:AK:8:ARG:CZ | 2.45 | 0.46 |
| 1:AA:243:U:OP1 | 10:AK:29:ARG:HB2 | 2.15 | 0.46 |
| 1:AA:332:C:O2' | 1:AA:333:G:H5' | 2.16 | 0.46 |
| 1:AA:390:C:OP2 | 5:AE:127:ILE:HD11 | 2.15 | 0.46 |
| 1:AA:689:G:H4' | 16:AQ:16:ARG:HG2 | 1.97 | 0.46 |
| 1:AA:1018:G:H2' | 1:AA:1019:A:C8 | 2.49 | 0.46 |
| 1:AA:1049:G:H2' | 1:AA:1050:A:O4' | 2.15 | 0.46 |
| 1:AA:1296:A:H2' | 1:AA:1297:A:O4' | 2.15 | 0.46 |
| 3:AC:11:GLY:HA2 | 3:AC:14:LYS:HE3 | 1.96 | 0.46 |
| 3:AC:72:LYS:HE2 | 3:AC:73:TYR:CZ | 2.50 | 0.46 |
| 3:AC:99:LEU:HD23 | 3:AC:178:CYS:SG | 2.56 | 0.46 |
| 20:AW:74:GLU:OE1 | 20:AW:74:GLU:N | 2.44 | 0.46 |
| 21:AX:33:LYS:NZ | 21:AX:46:PRO:HD2 | 2.27 | 0.46 |
| 24:AM:81:ALA:O | 24:AM:85:LEU:HG | 2.16 | 0.46 |
| 26:BA:2711:C:OP2 | 41:BP:62:ARG:HD2 | 2.15 | 0.46 |
| 27:BB:96:G:H2' | 27:BB:97:C:O4' | 2.16 | 0.46 |
| 29:BD:81:ARG:HB3 | 29:BD:146:TYR:CE1 | 2.50 | 0.46 |
| 29:BD:107:ASP:OD2 | 29:BD:148:LEU:HD21 | 2.15 | 0.46 |
| 29:BD:255:GLN:NE2 | 29:BD:257:GLY:O | 2.43 | 0.46 |
| 29:BD:279:THR:HG21 | 29:BD:289:LEU:CB | 2.46 | 0.46 |
| 31:BF:89:ASP:HB2 | 31:BF:93:ASN:O | 2.14 | 0.46 |
| 32:BG:44:ILE:HG13 | 32:BG:53:VAL:HG12 | 1.97 | 0.46 |
| 32:BG:62:GLN:O | 32:BG:66:VAL:HG13 | 2.15 | 0.46 |
| 36:BK:122:PRO:O | 36:BK:126:THR:HG23 | 2.14 | 0.46 |
| 1:AA:299:G:H2' | 1:AA:300:G:H5' | 1.96 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 1:AA:530:U:OP1 | 9:AJ:32:LYS:HG3 | 2.15 | 0.46 |
| 1:AA:693:G:O2' | 1:AA:694:A:OP2 | 2.30 | 0.46 |
| 1:AA:997:G:H5'' | 3:AC:179:LYS:NZ | 2.29 | 0.46 |
| 2:AB:107:VAL:HG12 | 58:AZ:27:PRO:HB2 | 1.96 | 0.46 |
| 6:AF:191:ILE:HG12 | 6:AF:202:VAL:CG2 | 2.43 | 0.46 |
| 7:AG:25:MET:HG2 | 7:AG:45:LEU:HD11 | 1.96 | 0.46 |
| 7:AG:117:SER:HA | 7:AG:145:PRO:HB2 | 1.97 | 0.46 |
| 7:AG:134:GLY:HA3 | 7:AG:191:ALA:HB2 | 1.97 | 0.46 |
| 13:AO:98:VAL:CG1 | 13:AO:124:VAL:HG23 | 2.45 | 0.46 |
| 15:AR:48:LYS:NZ | 15:AR:50:SER:OXT | 2.48 | 0.46 |
| 18:AU:33:GLU:O | 18:AU:37:LEU:HG | 2.15 | 0.46 |
| 19:AV:134:GLU:O | 19:AV:137:LEU:HB2 | 2.14 | 0.46 |
| 22:AY:16:GLU:HA | 22:AY:59:ASP:O | 2.16 | 0.46 |
| 23:AH:34:ILE:HG21 | 23:AH:65:PRO:HB3 | 1.96 | 0.46 |
| 24:AM:43:LYS:CB | 24:AM:68:VAL:HB | 2.38 | 0.46 |
| 26:BA:935:C:N3 | 49:BX:23:MET:HE2 | 2.30 | 0.46 |
| 26:BA:1336:A:N1 | 30:BE:49:PRO:HG3 | 2.30 | 0.46 |
| 26:BA:1854:U:H2' | 28:BC:121:ARG:HB2 | 1.96 | 0.46 |
| 29:BD:276:GLU:OE1 | 29:BD:276:GLU:N | 2.49 | 0.46 |
| 30:BE:110:ARG:O | 30:BE:114:ILE:HG13 | 2.14 | 0.46 |
| 34:BI:87:LEU:HB2 | 34:BI:134:PHE:CE1 | 2.51 | 0.46 |
| 42:BQ:35:THR:HG23 | 42:BQ:55:ILE:CD1 | 2.46 | 0.46 |
| 49:BX:83:ILE:HG13 | 49:BX:92:ILE:HG12 | 1.95 | 0.46 |
| 1:AA:157:A:N6 | 1:AA:186:U:H2' | 2.30 | 0.46 |
| 1:AA:538:U:H2' | 1:AA:539:C:H6 | 1.80 | 0.46 |
| 1:AA:541:U:O2' | 1:AA:542:C:H5' | 2.14 | 0.46 |
| 1:AA:912:U:H2' | 1:AA:913:C:O4' | 2.16 | 0.46 |
| 1:AA:941:A:H2' | 1:AA:942:G:H8 | 1.80 | 0.46 |
| 4:AD:37:ALA:HB3 | 4:AD:43:LEU:HD11 | 1.98 | 0.46 |
| 5:AE:22:ARG:O | 5:AE:26:GLU:HG3 | 2.15 | 0.46 |
| 6:AF:201:THR:HB | 6:AF:211:GLU:OE2 | 2.16 | 0.46 |
| 10:AK:45:PRO:HA | 10:AK:51:ARG:HA | 1.98 | 0.46 |
| 11:AL:85:ARG:O | 11:AL:89:GLU:HG3 | 2.15 | 0.46 |
| 14:AP:38:GLN:HB3 | 14:AP:63:PRO:CB | 2.36 | 0.46 |
| 14:AP:111:THR:O | 14:AP:114:LEU:HG | 2.15 | 0.46 |
| 23:AH:25:GLU:HG2 | 23:AH:42:ILE:O | 2.15 | 0.46 |
| 23:AH:56:CYS:O | 23:AH:105:ILE:HG23 | 2.15 | 0.46 |
| 24:AM:79:GLU:HA | 24:AM:82:LEU:HG | 1.96 | 0.46 |
| 26:BA:2035:A:H5' | 44:BS:134:THR:HB | 1.97 | 0.46 |
| 30:BE:127:VAL:O | 30:BE:132:HIS:HB2 | 2.15 | 0.46 |
| 30:BE:196:LEU:HA | 30:BE:216:ASP:O | 2.16 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 31:BF:18:SER:OG | 31:BF:55:GLU:OE2 | 2.12 | 0.46 |
| 32:BG:33:VAL:HG23 | 32:BG:80:VAL:HG12 | 1.98 | 0.46 |
| 32:BG:142:LYS:HA | 32:BG:145:VAL:HG12 | 1.98 | 0.46 |
| 37:BL:96:LEU:CD1 | 37:BL:101:ILE:HB | 2.45 | 0.46 |
| 50:BY:39:MET:HB2 | 50:BY:60:PRO:O | 2.15 | 0.46 |
| 52:Ba:120:ASP:OD1 | 52:Ba:120:ASP:N | 2.46 | 0.46 |
| 52:Ba:143:LYS:O | 52:Ba:146:GLU:HG3 | 2.16 | 0.46 |
| 53:Bb:53:ARG:O | 53:Bb:54:ILE:HD13 | 2.15 | 0.46 |
| 1:AA:627:U:H4' | 12:AN:33:LYS:HE2 | 1.96 | 0.46 |
| 1:AA:938:G:H1' | 1:AA:983:A:N6 | 2.26 | 0.46 |
| 1:AA:1014:C:H4' | 1:AA:1040:A:N6 | 2.29 | 0.46 |
| 1:AA:1236:A:H5' | 19:AV:80:LYS:NZ | 2.31 | 0.46 |
| 1:AA:1247:G:C8 | 1:AA:1282:C:H1' | 2.51 | 0.46 |
| 1:AA:1294:G:N2 | 1:AA:1321:A:OP2 | 2.32 | 0.46 |
| 3:AC:158:VAL:HA | 3:AC:162:VAL:CG2 | 2.43 | 0.46 |
| 5:AE:54:GLU:HA | 5:AE:57:LYS:HE2 | 1.97 | 0.46 |
| 7:AG:158:ALA:HA | 7:AG:161:VAL:HG12 | 1.96 | 0.46 |
| 16:AQ:57:ARG:HH12 | 21:AX:35:THR:HG23 | 1.81 | 0.46 |
| 18:AU:21:LYS:HB3 | 18:AU:25:GLU:OE2 | 2.15 | 0.46 |
| 25:AT:6:GLU:O | 25:AT:10:LYS:HG3 | 2.15 | 0.46 |
| 26:BA:340:U:C5' | 46:BU:3:ALA:HA | 2.45 | 0.46 |
| 28:BC:68:PHE:O | 28:BC:69:LEU:HD23 | 2.15 | 0.46 |
| 29:BD:270:LYS:HD3 | 29:BD:296:LEU:HD12 | 1.96 | 0.46 |
| 29:BD:279:THR:HG21 | 29:BD:289:LEU:HB2 | 1.97 | 0.46 |
| 32:BG:19:LEU:HD21 | 32:BG:44:ILE:HB | 1.97 | 0.46 |
| 32:BG:58:ALA:O | 32:BG:63:LYS:NZ | 2.49 | 0.46 |
| 34:BI:60:HIS:CE1 | 34:BI:61:THR:HG23 | 2.50 | 0.46 |
| 35:BJ:81:ARG:CG | 35:BJ:95:MET:HE2 | 2.45 | 0.46 |
| 39:BN:114:SER:OG | 39:BN:115:ARG:N | 2.49 | 0.46 |
| 46:BU:97:ASN:ND2 | 46:BU:98:PRO:HD2 | 2.30 | 0.46 |
| 50:BY:42:LEU:HD13 | 50:BY:61:VAL:HG13 | 1.97 | 0.46 |
| 55:Bd:42:HIS:HB3 | 55:Bd:45:ARG:HD3 | 1.96 | 0.46 |
| 1:AA:21:A:O2' | 5:AE:16:HIS:HA | 2.16 | 0.46 |
| 1:AA:172:G:H5' | 1:AA:173:A:OP2 | 2.16 | 0.46 |
| 1:AA:407:U:H4' | 5:AE:137:HIS:NE2 | 2.30 | 0.46 |
| 1:AA:572:U:OP1 | 1:AA:573:U:H5'' | 2.15 | 0.46 |
| 1:AA:625:G:H4' | 12:AN:29:GLU:OE1 | 2.16 | 0.46 |
| 1:AA:1040:A:H4' | 1:AA:1041:A:C4' | 2.45 | 0.46 |
| 1:AA:1392:U:H5'' | 1:AA:1393:C:C5 | 2.48 | 0.46 |
| 2:AB:104:ARG:HG3 | 2:AB:208:PRO:HD3 | 1.97 | 0.46 |
| 4:AD:21:ILE:HG21 | 4:AD:33:GLY:H | 1.79 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 8:AI:65:GLU:O | 8:AI:68:THR:HG22 | 2.15 | 0.46 |
| 8:AI:168:ARG:NH1 | 8:AI:168:ARG:HB2 | 2.31 | 0.46 |
| 11:AL:33:PRO:CG | 19:AV:6:ASP:HA | 2.46 | 0.46 |
| 18:AU:106:ARG:N | 18:AU:109:GLU:OE2 | 2.37 | 0.46 |
| 22:AY:14:ALA:CB | 22:AY:62:MET:HA | 2.45 | 0.46 |
| 24:AM:5:ARG:HG3 | 24:AM:74:ASP:OD1 | 2.15 | 0.46 |
| 26:BA:183:A:OP1 | 38:BM:164:LYS:NZ | 2.34 | 0.46 |
| 26:BA:2280:G:OP1 | 43:BR:14:LYS:HE3 | 2.16 | 0.46 |
| 27:BB:33:U:H2' | 27:BB:34:G:H8 | 1.80 | 0.46 |
| 33:BH:41:ILE:HG21 | 33:BH:67:LEU:HG | 1.98 | 0.46 |
| 35:BJ:121:MET:HE1 | 35:BJ:129:TYR:CZ | 2.50 | 0.46 |
| 42:BQ:27:GLU:O | 42:BQ:37:LYS:HE3 | 2.15 | 0.46 |
| 46:BU:55:VAL:HG21 | 46:BU:96:LEU:HD12 | 1.98 | 0.46 |
| 48:BW:3:ILE:CG1 | 48:BW:49:GLU:HG3 | 2.46 | 0.46 |
| 48:BW:23:ASN:O | 48:BW:27:GLU:HG2 | 2.14 | 0.46 |
| 52:Ba:30:MET:SD | 52:Ba:35:ARG:HD2 | 2.56 | 0.46 |
| 53:Bb:6:THR:HG23 | 53:Bb:9:GLY:N | 2.31 | 0.46 |
| 1:AA:604:A:H5'' | 4:AD:130:ARG:CD | 2.46 | 0.46 |
| 1:AA:626:G:O2' | 1:AA:627:U:H5' | 2.15 | 0.46 |
| 1:AA:879:C:C5 | 1:AA:1291:G:H2' | 2.50 | 0.46 |
| 1:AA:1104:A:O4' | 1:AA:1128:G:N1 | 2.49 | 0.46 |
| 1:AA:1264:U:H2' | 1:AA:1265:A:C8 | 2.50 | 0.46 |
| 2:AB:178:GLN:O | 2:AB:182:GLU:HG2 | 2.15 | 0.46 |
| 2:AB:199:LEU:HD12 | 2:AB:200:THR:H | 1.81 | 0.46 |
| 4:AD:138:ARG:O | 4:AD:142:VAL:HG23 | 2.15 | 0.46 |
| 5:AE:39:LYS:HE3 | 5:AE:39:LYS:HB2 | 1.74 | 0.46 |
| 5:AE:136:GLY:HA3 | 5:AE:144:LYS:CE | 2.46 | 0.46 |
| 6:AF:61:SER:HA | 6:AF:73:LYS:CG | 2.46 | 0.46 |
| 7:AG:13:LEU:HD12 | 7:AG:34:PRO:O | 2.15 | 0.46 |
| 10:AK:29:ARG:NH2 | 10:AK:46:THR:HG22 | 2.30 | 0.46 |
| 14:AP:70:LEU:HD11 | 14:AP:74:GLU:CB | 2.45 | 0.46 |
| 16:AQ:75:LEU:HD22 | 16:AQ:80:VAL:O | 2.15 | 0.46 |
| 18:AU:26:LEU:HA | 18:AU:29:LEU:CD1 | 2.45 | 0.46 |
| 26:BA:91:A:P | 48:BW:2:ALA:HB2 | 2.55 | 0.46 |
| 26:BA:1327:C:N4 | 30:BE:177:ILE:O | 2.49 | 0.46 |
| 27:BB:117:C:H2' | 27:BB:118:U:O4' | 2.16 | 0.46 |
| 28:BC:158:GLY:HA2 | 53:Bb:83:LEU:HD13 | 1.97 | 0.46 |
| 37:BL:66:ILE:HG22 | 37:BL:103:LYS:HB3 | 1.98 | 0.46 |
| 37:BL:118:SER:OG | 37:BL:119:GLU:N | 2.48 | 0.46 |
| 42:BQ:26:ILE:HG23 | 42:BQ:37:LYS:HD2 | 1.98 | 0.46 |
| 58:AZ:35:ARG:NH2 | 58:AZ:40:ARG:HA | 2.31 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 1:AA:125:A:H3' | 1:AA:126:U:H6 | 1.80 | 0.46 |
| 1:AA:327:C:H1' | 1:AA:328:G:C2 | 2.50 | 0.46 |
| 1:AA:384:G:H4' | 1:AA:560:C:N3 | 2.31 | 0.46 |
| 1:AA:389:C:H5'' | 5:AE:127:ILE:CD1 | 2.37 | 0.46 |
| 1:AA:486:G:O2' | 1:AA:488:G:H4' | 2.16 | 0.46 |
| 1:AA:662:U:H5' | 4:AD:128:ARG:NH2 | 2.29 | 0.46 |
| 1:AA:699:G:H4' | 1:AA:825:C:H4' | 1.98 | 0.46 |
| 3:AC:69:VAL:HG13 | 3:AC:75:LEU:HD12 | 1.97 | 0.46 |
| 4:AD:13:TRP:HB3 | 4:AD:14:LYS:NZ | 2.31 | 0.46 |
| 4:AD:90:ASP:OD1 | 4:AD:90:ASP:N | 2.43 | 0.46 |
| 5:AE:96:ASN:HB2 | 5:AE:98:ASP:OD1 | 2.16 | 0.46 |
| 5:AE:137:HIS:O | 5:AE:138:ILE:HD13 | 2.16 | 0.46 |
| 6:AF:59:ILE:HG23 | 6:AF:64:LYS:CD | 2.46 | 0.46 |
| 6:AF:101:LYS:HD2 | 6:AF:103:ARG:NH2 | 2.31 | 0.46 |
| 6:AF:178:GLY:HA3 | 6:AF:216:TYR:CZ | 2.51 | 0.46 |
| 6:AF:205:SER:HB2 | 6:AF:209:ASP:CG | 2.41 | 0.46 |
| 7:AG:38:PRO:CA | 7:AG:106:LEU:HD21 | 2.43 | 0.46 |
| 7:AG:49:GLU:HG3 | 7:AG:79:ARG:NH1 | 2.30 | 0.46 |
| 8:AI:159:ALA:O | 8:AI:163:ILE:HG12 | 2.15 | 0.46 |
| 10:AK:36:ILE:HD13 | 10:AK:92:ASN:HB3 | 1.97 | 0.46 |
| 12:AN:10:HIS:HB2 | 12:AN:21:THR:HG22 | 1.97 | 0.46 |
| 12:AN:31:ILE:CG2 | 12:AN:66:LYS:HG3 | 2.45 | 0.46 |
| 12:AN:73:ILE:HG21 | 12:AN:94:ILE:CD1 | 2.40 | 0.46 |
| 15:AR:32:LEU:HD11 | 15:AR:40:ILE:HD11 | 1.97 | 0.46 |
| 18:AU:106:ARG:HB2 | 18:AU:109:GLU:CG | 2.45 | 0.46 |
| 20:AW:53:GLN:HE22 | 20:AW:85:ILE:HD11 | 1.81 | 0.46 |
| 23:AH:58:GLY:HA2 | 23:AH:106:ILE:CD1 | 2.46 | 0.46 |
| 26:BA:1960:U:OP1 | 26:BA:2615:G:O2' | 2.12 | 0.46 |
| 26:BA:2781:C:OP1 | 29:BD:312:ARG:NH2 | 2.33 | 0.46 |
| 28:BC:110:GLU:OE2 | 28:BC:154:ARG:HD3 | 2.15 | 0.46 |
| 29:BD:147:THR:O | 29:BD:160:PRO:HB3 | 2.16 | 0.46 |
| 32:BG:30:LYS:HE3 | 32:BG:82:GLU:O | 2.14 | 0.46 |
| 35:BJ:60:GLY:CA | 35:BJ:66:PRO:HD2 | 2.46 | 0.46 |
| 35:BJ:105:PRO:HG2 | 35:BJ:108:LEU:HD12 | 1.98 | 0.46 |
| 39:BN:164:ALA:O | 39:BN:167:GLU:HG3 | 2.16 | 0.46 |
| 44:BS:29:LYS:O | 44:BS:33:VAL:HG23 | 2.15 | 0.46 |
| 46:BU:86:LYS:HD3 | 46:BU:88:ASP:OD2 | 2.16 | 0.46 |
| 51:BZ:58:HIS:HA | 51:BZ:61:GLU:HG3 | 1.98 | 0.46 |
| 1:AA:33:U:H5 | 1:AA:487:A:H8 | 1.64 | 0.46 |
| 1:AA:44:C:H5'' | 1:AA:45:U:OP2 | 2.16 | 0.46 |
| 1:AA:112:C:O2' | 1:AA:113:C:H5' | 2.15 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:518:U:H2' | 1:AA:519:C:O4' | 2.15 | 0.46 |
| 1:AA:573:U:H4' | 1:AA:574:U:C6 | 2.50 | 0.46 |
| 1:AA:686:C:OP1 | 1:AA:796:U:O2' | 2.23 | 0.46 |
| 1:AA:768:U:H3' | 1:AA:815:U:O4 | 2.15 | 0.46 |
| 1:AA:886:G:O2' | 1:AA:887:G:H5' | 2.15 | 0.46 |
| 1:AA:1008:U:H2' | 1:AA:1009:U:O4' | 2.15 | 0.46 |
| 2:AB:119:ARG:NH2 | 2:AB:164:GLU:O | 2.49 | 0.46 |
| 2:AB:176:ASP:HB3 | 2:AB:179:VAL:CG2 | 2.45 | 0.46 |
| 4:AD:125:THR:HG21 | 4:AD:129:ALA:HB2 | 1.97 | 0.46 |
| 5:AE:124:ARG:CG | 5:AE:177:PRO:HG3 | 2.46 | 0.46 |
| 8:AI:106:PRO:HG3 | 8:AI:134:THR:CG2 | 2.46 | 0.46 |
| 12:AN:75:VAL:HG21 | 12:AN:90:ALA:CB | 2.46 | 0.46 |
| 26:BA:30:G:N3 | 26:BA:450:G:O2' | 2.49 | 0.46 |
| 26:BA:152:C:OP1 | 38:BM:196:LYS:HE3 | 2.15 | 0.46 |
| 26:BA:480:A:O3' | 46:BU:86:LYS:HB2 | 2.16 | 0.46 |
| 26:BA:1223:U:C2 | 29:BD:243:PRO:HB3 | 2.50 | 0.46 |
| 30:BE:149:LEU:HB2 | 30:BE:154:GLN:OE1 | 2.16 | 0.46 |
| 30:BE:153:LYS:O | 30:BE:157:GLU:HG3 | 2.15 | 0.46 |
| 36:BK:81:ARG:NH2 | 36:BK:85:GLY:HA2 | 2.30 | 0.46 |
| 38:BM:179:LYS:HA | 38:BM:182:GLU:OE2 | 2.16 | 0.46 |
| 42:BQ:9:VAL:O | 42:BQ:23:THR:HA | 2.16 | 0.46 |
| 42:BQ:31:GLU:HG3 | 42:BQ:58:VAL:CG1 | 2.45 | 0.46 |
| 44:BS:28:LYS:O | 44:BS:32:GLU:HG2 | 2.16 | 0.46 |
| 51:BZ:44:THR:HG21 | 51:BZ:83:VAL:CG2 | 2.46 | 0.46 |
| 1:AA:121:C:H2' | 1:AA:122:G:O4' | 2.16 | 0.46 |
| 1:AA:554:A:H2' | 1:AA:555:C:H6 | 1.81 | 0.46 |
| 1:AA:653:G:H2' | 1:AA:654:G:H8 | 1.81 | 0.46 |
| 1:AA:893:G:H2' | 1:AA:894:C:H6 | 1.80 | 0.46 |
| 1:AA:1155:C:H2' | 1:AA:1156:C:O4' | 2.16 | 0.46 |
| 6:AF:109:LEU:HD22 | 6:AF:114:VAL:HG11 | 1.97 | 0.46 |
| 6:AF:177:VAL:CG1 | 6:AF:216:TYR:HA | 2.38 | 0.46 |
| 9:AJ:71:LYS:HE2 | 9:AJ:130:TYR:OH | 2.15 | 0.46 |
| 12:AN:19:ILE:HA | 12:AN:35:SER:HA | 1.98 | 0.46 |
| 13:AO:58:LYS:HD3 | 13:AO:113:ASP:HA | 1.98 | 0.46 |
| 14:AP:108:LEU:HD12 | 14:AP:113:ILE:HD13 | 1.96 | 0.46 |
| 18:AU:72:ARG:O | 18:AU:108:GLY:N | 2.48 | 0.46 |
| 21:AX:32:ARG:N | 21:AX:46:PRO:HB3 | 2.31 | 0.46 |
| 26:BA:140:A:N3 | 38:BM:112:ASN:HB3 | 2.31 | 0.46 |
| 26:BA:1883:U:O2' | 26:BA:1884:A:N7 | 2.41 | 0.46 |
| 30:BE:144:ASN:OD1 | 30:BE:243:SER:N | 2.44 | 0.46 |
| 39:BN:1:MET:HG3 | 39:BN:3:THR:CG2 | 2.45 | 0.46 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 39:BN:88:LEU:HA | 39:BN:169:ILE:HD11 | 1.98 | 0.46 |
| 40:BO:22:LEU:O | 40:BO:26:GLU:HG2 | 2.16 | 0.46 |
| 46:BU:44:SER:OG | 46:BU:45:ALA:N | 2.49 | 0.46 |
| 1:AA:200:C:H2' | 1:AA:201:U:O4' | 2.16 | 0.46 |
| 1:AA:429:C:H2' | 1:AA:430:U:O4' | 2.16 | 0.46 |
| 1:AA:682:A:H4' | 16:AQ:114:SER:OG | 2.16 | 0.46 |
| 1:AA:764:U:OP1 | 16:AQ:7:LYS:HB3 | 2.15 | 0.46 |
| 1:AA:830:G:O2' | 1:AA:831:G:H5' | 2.15 | 0.46 |
| 1:AA:1349:C:H2' | 1:AA:1350:C:O4' | 2.16 | 0.46 |
| 7:AG:17:VAL:CG1 | 7:AG:44:LEU:HD21 | 2.46 | 0.46 |
| 8:AI:178:LYS:O | 8:AI:182:GLU:HG3 | 2.16 | 0.46 |
| 8:AI:178:LYS:NZ | 8:AI:182:GLU:OE2 | 2.31 | 0.46 |
| 9:AJ:111:MET:HE3 | 9:AJ:116:ALA:CA | 2.42 | 0.46 |
| 11:AL:57:VAL:HG12 | 11:AL:61:LEU:HD11 | 1.98 | 0.46 |
| 12:AN:92:ALA:HA | 12:AN:95:ARG:HG2 | 1.97 | 0.46 |
| 14:AP:108:LEU:HB3 | 14:AP:112:ASP:CG | 2.41 | 0.46 |
| 14:AP:136:GLU:OE2 | 14:AP:137:ARG:HG3 | 2.16 | 0.46 |
| 14:AP:148:SER:HA | 14:AP:151:ARG:HH21 | 1.81 | 0.46 |
| 24:AM:27:ALA:O | 24:AM:31:GLY:N | 2.49 | 0.46 |
| 26:BA:1042:G:O2' | 26:BA:2276:A:OP2 | 2.22 | 0.46 |
| 26:BA:2324:C:H5' | 31:BF:10:ILE:HG21 | 1.97 | 0.46 |
| 29:BD:24:ILE:HG23 | 29:BD:308:LEU:CD2 | 2.46 | 0.46 |
| 30:BE:3:THR:HB | 30:BE:18:GLU:OE1 | 2.16 | 0.46 |
| 36:BK:4:MET:HE3 | 36:BK:120:ARG:NE | 2.31 | 0.46 |
| 36:BK:98:THR:HA | 36:BK:104:PRO:HA | 1.98 | 0.46 |
| 40:BO:68:LEU:HD22 | 40:BO:84:VAL:HG11 | 1.97 | 0.46 |
| 1:AA:554:A:H5'' | 5:AE:84:LYS:CG | 2.47 | 0.45 |
| 1:AA:558:C:H5' | 1:AA:559:U:OP2 | 2.15 | 0.45 |
| 1:AA:804:C:H2' | 1:AA:805:A:O4' | 2.16 | 0.45 |
| 1:AA:995:U:H2' | 1:AA:996:G:C8 | 2.50 | 0.45 |
| 1:AA:1006:A:O2' | 1:AA:1047:G:N2 | 2.49 | 0.45 |
| 1:AA:1068:C:H5'' | 11:AL:66:ASP:OD2 | 2.16 | 0.45 |
| 1:AA:1099:U:OP1 | 24:AM:69:HIS:HB3 | 2.15 | 0.45 |
| 1:AA:1231:C:H5'' | 1:AA:1232:A:OP2 | 2.16 | 0.45 |
| 1:AA:1394:G:OP2 | 1:AA:1394:G:N2 | 2.40 | 0.45 |
| 1:AA:1412:A:H2' | 1:AA:1413:A:H5' | 1.99 | 0.45 |
| 1:AA:1444:U:O2' | 1:AA:1445:A:H5' | 2.16 | 0.45 |
| 2:AB:152:GLY:O | 2:AB:159:ILE:HG21 | 2.16 | 0.45 |
| 3:AC:111:LYS:HE3 | 3:AC:115:ASN:OD1 | 2.16 | 0.45 |
| 3:AC:143:VAL:HG21 | 3:AC:145:LYS:HE3 | 1.99 | 0.45 |
| 5:AE:118:LEU:HD13 | 5:AE:126:VAL:CA | 2.46 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 6:AF:156:SER:O | 6:AF:160:LYS:HA | 2.15 | 0.45 |
| 14:AP:112:ASP:N | 14:AP:112:ASP:OD1 | 2.48 | 0.45 |
| 16:AQ:91:LEU:HD22 | 16:AQ:118:THR:HG23 | 1.97 | 0.45 |
| 18:AU:99:GLN:O | 18:AU:102:MET:HG3 | 2.16 | 0.45 |
| 23:AH:125:GLY:O | 23:AH:126:LEU:HD23 | 2.15 | 0.45 |
| 26:BA:749:U:OP1 | 37:BL:26:ARG:HD2 | 2.15 | 0.45 |
| 28:BC:180:MET:HE2 | 28:BC:180:MET:HA | 1.99 | 0.45 |
| 31:BF:9:VAL:CG2 | 31:BF:68:ALA:HB1 | 2.46 | 0.45 |
| 36:BK:73:VAL:CB | 36:BK:95:MET:HG2 | 2.46 | 0.45 |
| 54:Bc:42:THR:HG23 | 54:Bc:44:ARG:O | 2.16 | 0.45 |
| 57:Bf:68:ARG:HG3 | 57:Bf:77:ALA:HB1 | 1.98 | 0.45 |
| 1:AA:249:A:OP2 | 17:AS:96:LYS:HD3 | 2.16 | 0.45 |
| 1:AA:250:C:H1' | 10:AK:2:ARG:HH12 | 1.81 | 0.45 |
| 1:AA:488:G:H2' | 1:AA:489:U:O4' | 2.16 | 0.45 |
| 1:AA:653:G:H2' | 1:AA:654:G:C8 | 2.51 | 0.45 |
| 2:AB:164:GLU:OE1 | 7:AG:12:ARG:HG3 | 2.16 | 0.45 |
| 6:AF:205:SER:HB2 | 6:AF:209:ASP:OD1 | 2.16 | 0.45 |
| 8:AI:111:VAL:O | 8:AI:122:LYS:N | 2.42 | 0.45 |
| 11:AL:30:ASN:HB2 | 11:AL:65:VAL:HB | 1.98 | 0.45 |
| 16:AQ:71:VAL:O | 16:AQ:75:LEU:HG | 2.16 | 0.45 |
| 18:AU:44:ARG:HA | 18:AU:47:LYS:HB3 | 1.98 | 0.45 |
| 18:AU:76:ILE:HD13 | 18:AU:102:MET:O | 2.17 | 0.45 |
| 20:AW:60:GLY:O | 20:AW:62:GLN:NE2 | 2.49 | 0.45 |
| 23:AH:70:PRO:O | 23:AH:97:ARG:NH2 | 2.47 | 0.45 |
| 25:AT:27:ASP:OD1 | 25:AT:30:THR:N | 2.37 | 0.45 |
| 26:BA:742:A:O2' | 30:BE:104:LYS:O | 2.29 | 0.45 |
| 26:BA:1587:G:C6 | 50:BY:24:GLY:HA3 | 2.50 | 0.45 |
| 26:BA:2033:G:H4' | 44:BS:136:LYS:HE3 | 1.97 | 0.45 |
| 29:BD:198:LEU:O | 29:BD:314:PRO:HD3 | 2.16 | 0.45 |
| 35:BJ:88:ARG:O | 35:BJ:92:ARG:HG3 | 2.17 | 0.45 |
| 37:BL:79:LEU:HD12 | 37:BL:84:LEU:HB2 | 1.98 | 0.45 |
| 38:BM:3:LYS:HG2 | 38:BM:4:SER:O | 2.16 | 0.45 |
| 40:BO:9:LEU:HD11 | 40:BO:22:LEU:HD21 | 1.98 | 0.45 |
| 51:BZ:44:THR:HB | 51:BZ:77:LYS:NZ | 2.32 | 0.45 |
| 53:Bb:13:ARG:HG2 | 53:Bb:35:GLU:OE2 | 2.16 | 0.45 |
| 1:AA:172:G:H3' | 1:AA:173:A:O4' | 2.17 | 0.45 |
| 1:AA:1214:A:H1' | 14:AP:92:TRP:CH2 | 2.51 | 0.45 |
| 2:AB:120:ILE:HG23 | 2:AB:167:VAL:HG13 | 1.98 | 0.45 |
| 3:AC:105:ARG:CZ | 3:AC:107:TRP:HB2 | 2.46 | 0.45 |
| 4:AD:138:ARG:HA | 4:AD:141:MET:HE2 | 1.98 | 0.45 |
| 5:AE:82:LEU:O | 5:AE:86:ILE:HG12 | 2.17 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 6:AF:109:LEU:HD23 | 6:AF:228:ARG:HB2 | 1.98 | 0.45 |
| 10:AK:79:VAL:HG22 | 10:AK:94:LEU:HD23 | 1.99 | 0.45 |
| 12:AN:14:SER:OG | 12:AN:15:PHE:N | 2.50 | 0.45 |
| 12:AN:43:ALA:HA | 12:AN:46:GLU:HG3 | 1.99 | 0.45 |
| 12:AN:53:MET:HE1 | 22:AY:67:SER:OG | 2.15 | 0.45 |
| 25:AT:56:ARG:CD | 25:AT:60:ARG:HD3 | 2.46 | 0.45 |
| 26:BA:227:C:O2 | 26:BA:431:A:O2' | 2.24 | 0.45 |
| 26:BA:2691:G:H5'' | 29:BD:205:THR:HG21 | 1.98 | 0.45 |
| 26:BA:2795:U:H4' | 29:BD:117:PRO:HB3 | 1.98 | 0.45 |
| 27:BB:56:A:H1' | 31:BF:1:MET:HE1 | 1.98 | 0.45 |
| 31:BF:11:VAL:HG11 | 31:BF:29:LEU:HD11 | 1.99 | 0.45 |
| 32:BG:23:VAL:CG1 | 32:BG:36:LYS:HD3 | 2.47 | 0.45 |
| 33:BH:11:GLU:O | 33:BH:14:THR:OG1 | 2.30 | 0.45 |
| 36:BK:7:ASN:O | 36:BK:80:PHE:HB2 | 2.16 | 0.45 |
| 58:AZ:16:LEU:HB3 | 58:AZ:21:TYR:CB | 2.47 | 0.45 |
| 58:AZ:16:LEU:HB3 | 58:AZ:21:TYR:CG | 2.50 | 0.45 |
| 1:AA:177:C:H2' | 1:AA:178:U:H4' | 1.98 | 0.45 |
| 1:AA:192:G:H3' | 1:AA:192:G:OP2 | 2.17 | 0.45 |
| 1:AA:418:G:OP1 | 20:AW:54:ARG:HA | 2.15 | 0.45 |
| 1:AA:538:U:H2' | 1:AA:539:C:C6 | 2.51 | 0.45 |
| 1:AA:1116:A:H5'' | 25:AT:60:ARG:NE | 2.31 | 0.45 |
| 1:AA:1170:C:O5' | 1:AA:1171:U:H5'' | 2.16 | 0.45 |
| 1:AA:1240:A:H3' | 1:AA:1241:U:C4' | 2.46 | 0.45 |
| 1:AA:1353:U:H2' | 1:AA:1354:C:O4' | 2.16 | 0.45 |
| 1:AA:1393:C:H4' | 1:AA:1394:G:C4 | 2.51 | 0.45 |
| 2:AB:122:VAL:O | 2:AB:144:ALA:HA | 2.17 | 0.45 |
| 3:AC:95:MET:HE1 | 3:AC:98:ARG:CZ | 2.46 | 0.45 |
| 4:AD:27:LEU:HD13 | 4:AD:55:LEU:HD23 | 1.98 | 0.45 |
| 4:AD:64:THR:HG22 | 4:AD:87:LEU:HD13 | 1.99 | 0.45 |
| 10:AK:104:MET:HG3 | 10:AK:123:LEU:HD22 | 1.98 | 0.45 |
| 12:AN:56:ALA:HB3 | 12:AN:96:ALA:HB3 | 1.97 | 0.45 |
| 14:AP:108:LEU:HB2 | 14:AP:113:ILE:CG1 | 2.45 | 0.45 |
| 19:AV:67:PRO:CB | 19:AV:114:VAL:HG11 | 2.47 | 0.45 |
| 30:BE:57:THR:CG2 | 30:BE:82:ARG:HB2 | 2.47 | 0.45 |
| 33:BH:55:ILE:HG13 | 33:BH:61:VAL:CG2 | 2.47 | 0.45 |
| 41:BP:126:LYS:HD2 | 41:BP:131:GLU:HG3 | 1.97 | 0.45 |
| 47:BV:56:LYS:O | 47:BV:60:LYS:HG2 | 2.15 | 0.45 |
| 47:BV:58:GLN:HA | 47:BV:61:LYS:CD | 2.47 | 0.45 |
| 52:Ba:129:ALA:HB3 | 52:Ba:132:ILE:CG1 | 2.47 | 0.45 |
| 1:AA:441:G:H2' | 1:AA:442:C:O4' | 2.16 | 0.45 |
| 1:AA:609:U:O2' | 16:AQ:105:ASN:HB3 | 2.16 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:753:G:OP1 | 1:AA:848:G:H1' | 2.17 | 0.45 |
| 1:AA:1002:C:H5'' | 1:AA:1003:G:OP2 | 2.17 | 0.45 |
| 1:AA:1426:G:H2' | 1:AA:1427:G:O4' | 2.16 | 0.45 |
| 3:AC:13:VAL:O | 3:AC:17:MET:HG2 | 2.16 | 0.45 |
| 3:AC:117:LEU:HD21 | 3:AC:148:ASP:OD1 | 2.16 | 0.45 |
| 5:AE:104:LYS:HB3 | 5:AE:106:GLU:OE1 | 2.17 | 0.45 |
| 5:AE:104:LYS:HB3 | 5:AE:104:LYS:HE3 | 1.78 | 0.45 |
| 7:AG:35:ILE:CG2 | 7:AG:41:ILE:HD11 | 2.46 | 0.45 |
| 7:AG:142:THR:O | 7:AG:172:THR:HG22 | 2.17 | 0.45 |
| 9:AJ:18:GLU:HG3 | 9:AJ:69:ILE:CG1 | 2.44 | 0.45 |
| 13:AO:127:VAL:CG2 | 13:AO:137:MET:HG2 | 2.47 | 0.45 |
| 14:AP:26:ILE:HG23 | 14:AP:79:ASP:OD1 | 2.16 | 0.45 |
| 14:AP:66:ILE:HG22 | 14:AP:68:GLY:N | 2.31 | 0.45 |
| 14:AP:101:ALA:N | 18:AU:18:TYR:O | 2.49 | 0.45 |
| 26:BA:78:G:OP1 | 46:BU:48:ILE:HA | 2.16 | 0.45 |
| 30:BE:149:LEU:HD13 | 30:BE:154:GLN:HB2 | 1.99 | 0.45 |
| 31:BF:70:GLU:O | 31:BF:74:THR:HG23 | 2.16 | 0.45 |
| 31:BF:131:ILE:HD12 | 31:BF:134:ARG:NH2 | 2.32 | 0.45 |
| 32:BG:88:MET:HE2 | 32:BG:170:ILE:CG2 | 2.45 | 0.45 |
| 38:BM:113:LEU:HD22 | 38:BM:136:ASP:CA | 2.46 | 0.45 |
| 41:BP:84:LYS:HE3 | 41:BP:84:LYS:HB3 | 1.85 | 0.45 |
| 44:BS:126:MET:CE | 44:BS:136:LYS:HB3 | 2.47 | 0.45 |
| 45:BT:24:PHE:CE2 | 45:BT:64:LEU:HD13 | 2.50 | 0.45 |
| 52:Ba:73:GLN:HA | 52:Ba:78:ARG:NH2 | 2.32 | 0.45 |
| 1:AA:98:C:OP1 | 1:AA:294:G:H5' | 2.17 | 0.45 |
| 1:AA:366:C:OP1 | 6:AF:54:ARG:NH2 | 2.50 | 0.45 |
| 1:AA:444:G:H4' | 1:AA:474:C:N3 | 2.32 | 0.45 |
| 1:AA:484:C:H5'' | 5:AE:23:MET:CE | 2.47 | 0.45 |
| 1:AA:961:G:H2' | 1:AA:962:A:O4' | 2.17 | 0.45 |
| 1:AA:1087:A:H3' | 1:AA:1088:U:H5'' | 1.97 | 0.45 |
| 1:AA:1123:A:H4' | 1:AA:1124:G:C8 | 2.52 | 0.45 |
| 2:AB:184:SER:HA | 58:AZ:40:ARG:NH1 | 2.31 | 0.45 |
| 3:AC:13:VAL:HG12 | 3:AC:17:MET:HE2 | 1.98 | 0.45 |
| 3:AC:169:ALA:O | 3:AC:175:THR:HA | 2.16 | 0.45 |
| 4:AD:122:THR:O | 4:AD:183:VAL:HG13 | 2.17 | 0.45 |
| 22:AY:31:SER:OG | 22:AY:55:VAL:HG21 | 2.17 | 0.45 |
| 24:AM:5:ARG:O | 24:AM:98:ILE:HA | 2.17 | 0.45 |
| 26:BA:769:U:O2 | 54:Bc:11:ARG:HA | 2.17 | 0.45 |
| 26:BA:812:G:C5 | 28:BC:172:LYS:HG3 | 2.51 | 0.45 |
| 29:BD:85:ILE:HG12 | 29:BD:183:LEU:HD13 | 1.98 | 0.45 |
| 30:BE:218:VAL:HG11 | 30:BE:227:LEU:HD11 | 1.97 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 31:BF:78:ILE:HG23 | 31:BF:103:ASP:HB3 | 1.99 | 0.45 |
| 32:BG:84:PHE:CE2 | 32:BG:142:LYS:HB2 | 2.52 | 0.45 |
| 33:BH:13:LEU:HD13 | 33:BH:115:LYS:HD2 | 1.98 | 0.45 |
| 36:BK:40:VAL:HG22 | 36:BK:41:LYS:H | 1.82 | 0.45 |
| 41:BP:60:ARG:HG2 | 41:BP:63:ALA:HB3 | 1.99 | 0.45 |
| 48:BW:32:ARG:HH21 | 48:BW:51:ARG:HD3 | 1.81 | 0.45 |
| 1:AA:597:A:H2' | 1:AA:598:A:H8 | 1.82 | 0.45 |
| 6:AF:48:LYS:HA | 6:AF:48:LYS:HD2 | 1.73 | 0.45 |
| 14:AP:71:PRO:HG2 | 14:AP:74:GLU:OE2 | 2.16 | 0.45 |
| 15:AR:16:LYS:HB2 | 15:AR:32:LEU:HD23 | 1.98 | 0.45 |
| 15:AR:27:LYS:HD3 | 24:AM:47:VAL:HG23 | 1.98 | 0.45 |
| 18:AU:97:ASP:OD1 | 18:AU:98:LEU:N | 2.50 | 0.45 |
| 23:AH:34:ILE:HG23 | 23:AH:53:THR:HA | 1.97 | 0.45 |
| 25:AT:54:VAL:O | 25:AT:58:VAL:HG23 | 2.17 | 0.45 |
| 26:BA:652:A:O2' | 26:BA:738:A:N1 | 2.47 | 0.45 |
| 26:BA:1224:U:O2' | 26:BA:1225:G:O5' | 2.32 | 0.45 |
| 28:BC:112:LYS:O | 28:BC:115:ASP:HB2 | 2.17 | 0.45 |
| 29:BD:213:PRO:HB2 | 29:BD:219:ILE:HD13 | 1.98 | 0.45 |
| 31:BF:83:LEU:CD1 | 31:BF:87:GLN:HG3 | 2.47 | 0.45 |
| 35:BJ:61:ALA:N | 35:BJ:65:GLY:HA3 | 2.32 | 0.45 |
| 35:BJ:130:VAL:HB | 35:BJ:134:GLU:OE1 | 2.17 | 0.45 |
| 40:BO:73:LYS:HA | 40:BO:90:THR:O | 2.17 | 0.45 |
| 45:BT:57:LYS:O | 55:Bd:17:LYS:NZ | 2.33 | 0.45 |
| 49:BX:148:LEU:O | 49:BX:152:MET:HG3 | 2.16 | 0.45 |
| 50:BY:16:VAL:HG23 | 50:BY:21:VAL:CG2 | 2.47 | 0.45 |
| 1:AA:208:G:O2' | 6:AF:134:ASN:OD1 | 2.19 | 0.45 |
| 1:AA:234:U:H5'' | 10:AK:8:ARG:NE | 2.32 | 0.45 |
| 1:AA:586:G:O2' | 1:AA:587:C:H5' | 2.17 | 0.45 |
| 1:AA:655:G:H2' | 1:AA:656:A:H8 | 1.82 | 0.45 |
| 1:AA:1289:C:O2' | 11:AL:130:GLN:HG2 | 2.16 | 0.45 |
| 1:AA:1329:C:N4 | 22:AY:27:HIS:HB3 | 2.31 | 0.45 |
| 1:AA:1407:A:H2' | 1:AA:1408:C:C6 | 2.52 | 0.45 |
| 1:AA:1437:A:H1' | 1:AA:1458:U:H5' | 1.99 | 0.45 |
| 2:AB:229:SER:HA | 2:AB:232:ASN:OD1 | 2.16 | 0.45 |
| 3:AC:130:VAL:CB | 3:AC:180:VAL:HG22 | 2.45 | 0.45 |
| 6:AF:168:PHE:HB3 | 6:AF:219:VAL:CG1 | 2.47 | 0.45 |
| 7:AG:74:VAL:HG12 | 7:AG:100:ALA:HB1 | 1.98 | 0.45 |
| 9:AJ:29:PRO:HA | 9:AJ:59:GLY:O | 2.17 | 0.45 |
| 9:AJ:105:THR:HA | 9:AJ:110:VAL:HG22 | 1.99 | 0.45 |
| 14:AP:92:TRP:HZ2 | 19:AV:35:VAL:HG22 | 1.81 | 0.45 |
| 15:AR:27:LYS:HD3 | 24:AM:47:VAL:CG2 | 2.47 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 17:AS:46:THR:HA | 17:AS:72:ALA:O | 2.17 | 0.45 |
| 18:AU:74:MET:O | 18:AU:106:ARG:HA | 2.16 | 0.45 |
| 26:BA:1113:U:N3 | 26:BA:1119:U:OP1 | 2.50 | 0.45 |
| 26:BA:1396:C:OP2 | 55:Bd:1:MET:HB2 | 2.16 | 0.45 |
| 26:BA:2633:G:O2' | 26:BA:2816:G:N7 | 2.49 | 0.45 |
| 28:BC:25:LYS:HG2 | 28:BC:50:PRO:HD2 | 1.98 | 0.45 |
| 32:BG:142:LYS:HG3 | 32:BG:176:ALA:HB1 | 1.98 | 0.45 |
| 34:BI:57:GLN:OE1 | 34:BI:126:ARG:NH2 | 2.50 | 0.45 |
| 38:BM:29:LEU:O | 38:BM:33:ARG:HG3 | 2.16 | 0.45 |
| 43:BR:87:SER:HB3 | 43:BR:92:LEU:HD21 | 1.98 | 0.45 |
| 45:BT:20:ASN:O | 45:BT:72:ALA:HB2 | 2.16 | 0.45 |
| 46:BU:52:THR:HA | 46:BU:65:GLY:O | 2.17 | 0.45 |
| 49:BX:81:GLU:O | 49:BX:85:GLU:HG2 | 2.17 | 0.45 |
| 1:AA:286:A:H2' | 1:AA:287:G:H4' | 1.98 | 0.45 |
| 1:AA:385:A:H5' | 5:AE:148:PRO:CG | 2.46 | 0.45 |
| 1:AA:397:A:H1' | 1:AA:398:U:C4 | 2.52 | 0.45 |
| 1:AA:472:A:H61 | 3:AC:135:LYS:HZ3 | 1.65 | 0.45 |
| 1:AA:951:A:H2' | 1:AA:952:G:C8 | 2.52 | 0.45 |
| 2:AB:66:LEU:HA | 2:AB:71:HIS:ND1 | 2.31 | 0.45 |
| 2:AB:143:ARG:NH2 | 2:AB:161:GLY:HA2 | 2.32 | 0.45 |
| 5:AE:46:SER:HA | 5:AE:49:ARG:CG | 2.47 | 0.45 |
| 5:AE:110:GLU:OE2 | 5:AE:116:GLN:NE2 | 2.50 | 0.45 |
| 5:AE:130:ARG:HE | 5:AE:148:PRO:HG3 | 1.81 | 0.45 |
| 6:AF:42:ILE:HD13 | 6:AF:46:ILE:CD1 | 2.43 | 0.45 |
| 6:AF:71:PRO:O | 6:AF:73:LYS:HD2 | 2.17 | 0.45 |
| 7:AG:45:LEU:O | 7:AG:48:LEU:HD23 | 2.17 | 0.45 |
| 7:AG:126:HIS:HB2 | 7:AG:168:LYS:HD2 | 1.98 | 0.45 |
| 8:AI:152:ARG:HH22 | 8:AI:160:THR:HG21 | 1.82 | 0.45 |
| 10:AK:63:VAL:CG2 | 10:AK:102:THR:HG21 | 2.46 | 0.45 |
| 11:AL:9:GLY:HA2 | 11:AL:109:VAL:HG23 | 1.99 | 0.45 |
| 11:AL:56:ASP:N | 11:AL:56:ASP:OD1 | 2.50 | 0.45 |
| 13:AO:132:LEU:HD23 | 13:AO:137:MET:O | 2.16 | 0.45 |
| 16:AQ:17:PRO:HB3 | 21:AX:8:PRO:HG3 | 1.97 | 0.45 |
| 23:AH:39:ASP:HA | 23:AH:49:LYS:HE3 | 1.98 | 0.45 |
| 26:BA:41:U:O2' | 26:BA:110:A:N1 | 2.49 | 0.45 |
| 26:BA:669:C:O2' | 30:BE:224:ASN:HB3 | 2.16 | 0.45 |
| 26:BA:2081:A:H5'' | 30:BE:64:SER:HB2 | 1.99 | 0.45 |
| 31:BF:24:ASN:O | 31:BF:28:ILE:HG23 | 2.16 | 0.45 |
| 38:BM:3:LYS:HD2 | 38:BM:7:GLY:HA3 | 1.98 | 0.45 |
| 38:BM:41:ILE:HG21 | 38:BM:65:ARG:NH1 | 2.32 | 0.45 |
| 41:BP:92:LYS:O | 41:BP:96:ILE:HG13 | 2.16 | 0.45 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 46:BU:86:LYS:HB2 | 46:BU:86:LYS:HE3 | 1.66 | 0.45 |
| 57:Bf:24:ARG:HD2 | 57:Bf:67:LEU:HD23 | 1.99 | 0.45 |
| 1:AA:303:U:H2' | 1:AA:304:C:C6 | 2.51 | 0.45 |
| 1:AA:526:C:O2' | 1:AA:823:G:H4' | 2.17 | 0.45 |
| 1:AA:913:C:H5'' | 1:AA:914:A:OP2 | 2.16 | 0.45 |
| 1:AA:956:A:H2' | 1:AA:957:G:O4' | 2.17 | 0.45 |
| 1:AA:988:A:N1 | 15:AR:22:GLN:NE2 | 2.60 | 0.45 |
| 1:AA:1107:G:H5'' | 25:AT:52:GLY:O | 2.16 | 0.45 |
| 1:AA:1291:G:O2' | 1:AA:1292:U:H5' | 2.16 | 0.45 |
| 1:AA:1348:G:H2' | 1:AA:1349:C:O4' | 2.17 | 0.45 |
| 1:AA:1364:G:O2' | 1:AA:1421:A:N6 | 2.49 | 0.45 |
| 5:AE:58:LEU:HD13 | 5:AE:78:SER:CB | 2.47 | 0.45 |
| 6:AF:68:ASP:HB3 | 6:AF:160:LYS:CE | 2.47 | 0.45 |
| 8:AI:108:GLU:OE2 | 8:AI:186:LYS:HE3 | 2.17 | 0.45 |
| 10:AK:91:ARG:HB2 | 10:AK:93:ILE:HG12 | 1.98 | 0.45 |
| 14:AP:100:LEU:HB3 | 14:AP:101:ALA:H | 1.51 | 0.45 |
| 21:AX:14:ARG:HG2 | 21:AX:58:LEU:HD12 | 2.00 | 0.45 |
| 22:AY:38:LEU:CA | 22:AY:43:LYS:HG2 | 2.44 | 0.45 |
| 25:AT:26:LYS:NZ | 25:AT:59:ALA:HA | 2.32 | 0.45 |
| 26:BA:237:A:OP1 | 38:BM:4:SER:HB3 | 2.16 | 0.45 |
| 26:BA:1429:U:OP2 | 45:BT:61:LYS:NZ | 2.43 | 0.45 |
| 26:BA:1436:G:OP2 | 26:BA:1473:G:N2 | 2.49 | 0.45 |
| 26:BA:1852:A:C2 | 28:BC:26:ALA:HA | 2.52 | 0.45 |
| 29:BD:76:GLU:OE1 | 29:BD:287:TYR:OH | 2.20 | 0.45 |
| 30:BE:41:ALA:HB1 | 30:BE:101:ARG:O | 2.17 | 0.45 |
| 31:BF:63:LEU:C | 31:BF:68:ALA:HB2 | 2.41 | 0.45 |
| 34:BI:20:ARG:HA | 34:BI:23:MET:HG2 | 1.98 | 0.45 |
| 39:BN:80:THR:HG22 | 39:BN:143:ILE:O | 2.17 | 0.45 |
| 40:BO:52:ALA:HA | 40:BO:73:LYS:O | 2.17 | 0.45 |
| 45:BT:22:LEU:HD23 | 45:BT:24:PHE:CZ | 2.52 | 0.45 |
| 1:AA:32:A:O2' | 1:AA:487:A:N7 | 2.47 | 0.44 |
| 1:AA:243:U:OP1 | 10:AK:29:ARG:NE | 2.40 | 0.44 |
| 1:AA:277:G:N2 | 1:AA:496:C:H4' | 2.32 | 0.44 |
| 1:AA:596:G:O2' | 16:AQ:4:MET:SD | 2.73 | 0.44 |
| 1:AA:1013:A:H2' | 1:AA:1014:C:O4' | 2.17 | 0.44 |
| 1:AA:1069:A:H1' | 1:AA:1093:A:H62 | 1.81 | 0.44 |
| 1:AA:1220:G:H2' | 1:AA:1221:C:O4' | 2.17 | 0.44 |
| 1:AA:1227:A:OP2 | 24:AM:43:LYS:NZ | 2.45 | 0.44 |
| 2:AB:107:VAL:HG13 | 58:AZ:27:PRO:O | 2.17 | 0.44 |
| 2:AB:209:THR:HG21 | 2:AB:220:VAL:HG11 | 1.99 | 0.44 |
| 5:AE:124:ARG:HG3 | 5:AE:177:PRO:HG3 | 1.99 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 6:AF:62:GLU:OE1 | 6:AF:64:LYS:NZ | 2.42 | 0.44 |
| 8:AI:1:MET:HA | 11:AL:99:ASN:ND2 | 2.31 | 0.44 |
| 9:AJ:95:PRO:HG3 | 9:AJ:130:TYR:CD2 | 2.52 | 0.44 |
| 18:AU:69:THR:HG21 | 18:AU:107:PHE:CD2 | 2.51 | 0.44 |
| 20:AW:83:GLU:HA | 20:AW:86:LEU:HD12 | 1.98 | 0.44 |
| 23:AH:63:MET:SD | 23:AH:105:ILE:HD13 | 2.56 | 0.44 |
| 26:BA:302:U:C5 | 46:BU:56:MET:HE3 | 2.51 | 0.44 |
| 26:BA:793:U:O2' | 41:BP:131:GLU:OE2 | 2.30 | 0.44 |
| 26:BA:891:C:OP1 | 26:BA:915:A:O2' | 2.25 | 0.44 |
| 26:BA:941:U:O3' | 43:BR:96:LYS:NZ | 2.37 | 0.44 |
| 26:BA:1855:A:OP1 | 28:BC:125:VAL:HG21 | 2.17 | 0.44 |
| 30:BE:34:LYS:HB3 | 30:BE:34:LYS:HE2 | 1.80 | 0.44 |
| 30:BE:84:PRO:HB3 | 30:BE:92:ALA:HB2 | 1.99 | 0.44 |
| 30:BE:206:LEU:HD23 | 30:BE:217:VAL:CG2 | 2.47 | 0.44 |
| 31:BF:8:LYS:HG2 | 31:BF:123:VAL:CG2 | 2.42 | 0.44 |
| 32:BG:84:PHE:CZ | 32:BG:142:LYS:HB2 | 2.52 | 0.44 |
| 33:BH:35:ASN:O | 33:BH:39:LYS:HG2 | 2.17 | 0.44 |
| 35:BJ:72:PRO:HG3 | 35:BJ:136:SER:HB2 | 1.99 | 0.44 |
| 49:BX:17:ILE:O | 49:BX:21:MET:HG2 | 2.18 | 0.44 |
| 50:BY:42:LEU:HA | 50:BY:82:ALA:O | 2.18 | 0.44 |
| 1:AA:45:U:H2' | 1:AA:46:A:H2' | 2.00 | 0.44 |
| 1:AA:160:G:H4' | 10:AK:116:ASP:CG | 2.41 | 0.44 |
| 1:AA:338:A:H5' | 1:AA:349:C:C5 | 2.52 | 0.44 |
| 1:AA:780:A:H2' | 1:AA:781:U:O4' | 2.17 | 0.44 |
| 1:AA:829:U:OP2 | 13:AO:21:LYS:HD3 | 2.16 | 0.44 |
| 1:AA:997:G:P | 3:AC:133:SER:HB2 | 2.57 | 0.44 |
| 1:AA:1009:U:H2' | 1:AA:1010:C:C6 | 2.51 | 0.44 |
| 1:AA:1019:A:H5'' | 7:AG:89:LYS:HZ3 | 1.82 | 0.44 |
| 1:AA:1173:C:OP1 | 14:AP:139:LEU:HD21 | 2.17 | 0.44 |
| 3:AC:25:LEU:HD22 | 3:AC:30:TYR:CD1 | 2.51 | 0.44 |
| 3:AC:37:ARG:HB3 | 24:AM:92:LYS:HE3 | 1.99 | 0.44 |
| 3:AC:44:ILE:O | 3:AC:80:ILE:HA | 2.17 | 0.44 |
| 3:AC:46:ILE:CG1 | 3:AC:82:ALA:HA | 2.45 | 0.44 |
| 4:AD:13:TRP:HB3 | 4:AD:14:LYS:HZ2 | 1.81 | 0.44 |
| 4:AD:184:GLU:O | 4:AD:185:ILE:HD13 | 2.17 | 0.44 |
| 4:AD:187:LYS:HB3 | 4:AD:189:GLU:OE1 | 2.18 | 0.44 |
| 6:AF:100:GLU:OE1 | 6:AF:100:GLU:N | 2.48 | 0.44 |
| 16:AQ:40:LEU:HD13 | 16:AQ:53:VAL:HG11 | 1.97 | 0.44 |
| 18:AU:27:GLN:HG3 | 18:AU:78:PRO:O | 2.18 | 0.44 |
| 20:AW:17:ARG:HD3 | 20:AW:70:VAL:CG1 | 2.47 | 0.44 |
| 26:BA:650:G:H1' | 30:BE:109:GLU:HB3 | 1.99 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 27:BB:49:A:H2' | 27:BB:50:C:H6 | 1.82 | 0.44 |
| 29:BD:25:PRO:HG3 | 29:BD:202:ALA:O | 2.17 | 0.44 |
| 29:BD:72:VAL:HG21 | 29:BD:296:LEU:HD22 | 1.99 | 0.44 |
| 31:BF:45:THR:HG22 | 31:BF:53:LYS:CE | 2.45 | 0.44 |
| 32:BG:108:ILE:HD13 | 32:BG:156:THR:CG2 | 2.47 | 0.44 |
| 35:BJ:121:MET:HE1 | 35:BJ:129:TYR:CE2 | 2.52 | 0.44 |
| 40:BO:67:VAL:HA | 40:BO:85:THR:CG2 | 2.46 | 0.44 |
| 41:BP:24:LEU:HB3 | 41:BP:32:ILE:HD13 | 1.99 | 0.44 |
| 44:BS:25:ILE:CG2 | 44:BS:87:GLU:HG3 | 2.45 | 0.44 |
| 1:AA:19:A:O2' | 1:AA:20:G:H5' | 2.18 | 0.44 |
| 1:AA:113:C:O2' | 6:AF:30:PRO:HB3 | 2.17 | 0.44 |
| 1:AA:344:G:H2' | 1:AA:346:A:OP2 | 2.18 | 0.44 |
| 1:AA:718:C:H2' | 1:AA:719:A:O4' | 2.16 | 0.44 |
| 1:AA:1063:G:N2 | 1:AA:1097:U:H1' | 2.28 | 0.44 |
| 1:AA:1176:G:H2' | 1:AA:1177:C:O4' | 2.17 | 0.44 |
| 1:AA:1321:A:H2' | 1:AA:1322:A:H8 | 1.82 | 0.44 |
| 1:AA:1355:A:H2 | 1:AA:1430:A:H62 | 1.65 | 0.44 |
| 1:AA:1395:U:H3' | 1:AA:1396:G:H5' | 2.00 | 0.44 |
| 2:AB:214:ARG:NH2 | 2:AB:247:LEU:HD23 | 2.32 | 0.44 |
| 6:AF:33:GLN:O | 6:AF:33:GLN:NE2 | 2.41 | 0.44 |
| 6:AF:65:VAL:HG13 | 6:AF:87:LEU:CD2 | 2.46 | 0.44 |
| 10:AK:46:THR:CG2 | 10:AK:52:LYS:HG3 | 2.47 | 0.44 |
| 11:AL:36:LEU:HD12 | 19:AV:6:ASP:O | 2.17 | 0.44 |
| 12:AN:37:GLY:HA2 | 12:AN:40:VAL:O | 2.17 | 0.44 |
| 12:AN:94:ILE:HD13 | 12:AN:107:ILE:HG21 | 1.98 | 0.44 |
| 14:AP:137:ARG:CB | 14:AP:139:LEU:HD13 | 2.47 | 0.44 |
| 17:AS:73:HIS:O | 17:AS:102:VAL:HG22 | 2.18 | 0.44 |
| 19:AV:15:LYS:O | 19:AV:19:ILE:HG13 | 2.17 | 0.44 |
| 20:AW:8:ASP:HA | 20:AW:18:GLU:O | 2.17 | 0.44 |
| 20:AW:36:ARG:HG2 | 20:AW:52:ILE:HG23 | 2.00 | 0.44 |
| 23:AH:57:ASP:OD2 | 23:AH:61:PHE:HB2 | 2.18 | 0.44 |
| 26:BA:1369:G:H21 | 41:BP:1:MET:HE3 | 1.81 | 0.44 |
| 26:BA:1539:U:O2 | 26:BA:1544:G:N2 | 2.40 | 0.44 |
| 28:BC:172:LYS:HG2 | 28:BC:174:GLY:H | 1.81 | 0.44 |
| 29:BD:82:VAL:HG13 | 29:BD:143:ALA:HB1 | 1.98 | 0.44 |
| 30:BE:41:ALA:HA | 30:BE:44:LEU:HD12 | 1.99 | 0.44 |
| 36:BK:4:MET:HE3 | 36:BK:120:ARG:HD2 | 1.99 | 0.44 |
| 39:BN:90:GLY:HA3 | 39:BN:126:SER:OG | 2.17 | 0.44 |
| 40:BO:51:ALA:O | 40:BO:72:GLY:HA3 | 2.17 | 0.44 |
| 52:Ba:120:ASP:HA | 52:Ba:121:PRO:HD3 | 1.85 | 0.44 |
| 1:AA:394:A:H2' | 1:AA:395:U:C6 | 2.52 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|-------------------|--------------------------|-------------------|
| 1:AA:428:U:H2' | 1:AA:429:C:O4' | 2.18 | 0.44 |
| 1:AA:1256:G:H21 | 14:AP:95:ASN:HB2 | 1.83 | 0.44 |
| 2:AB:140:LEU:CB | 2:AB:142:THR:HG23 | 2.47 | 0.44 |
| 2:AB:176:ASP:HB3 | 2:AB:179:VAL:HG22 | 1.99 | 0.44 |
| 5:AE:7:LYS:HG3 | 5:AE:8:SER:O | 2.17 | 0.44 |
| 7:AG:35:ILE:HG22 | 7:AG:41:ILE:HD11 | 2.00 | 0.44 |
| 7:AG:112:ARG:HD3 | 58:AZ:15:ARG:NH2 | 2.23 | 0.44 |
| 8:AI:103:ASN:ND2 | 8:AI:175:ILE:HD11 | 2.33 | 0.44 |
| 11:AL:43:MET:O | 11:AL:46:ILE:HG22 | 2.18 | 0.44 |
| 14:AP:80:THR:O | 14:AP:84:LYS:HG3 | 2.18 | 0.44 |
| 16:AQ:144:GLU:N | 16:AQ:144:GLU:OE1 | 2.50 | 0.44 |
| 17:AS:73:HIS:HA | 17:AS:98:LYS:NZ | 2.33 | 0.44 |
| 20:AW:52:ILE:HD12 | 20:AW:55:ILE:HD11 | 1.99 | 0.44 |
| 22:AY:63:LEU:CD1 | 22:AY:66:THR:HG22 | 2.47 | 0.44 |
| 23:AH:64:LYS:HD2 | 23:AH:96:MET:CE | 2.48 | 0.44 |
| 26:BA:812:G:OP1 | 53:Bb:21:ARG:NH2 | 2.37 | 0.44 |
| 27:BB:3:G:H2' | 27:BB:4:U:H6 | 1.80 | 0.44 |
| 29:BD:106:LEU:HD22 | 29:BD:146:TYR:OH | 2.18 | 0.44 |
| 29:BD:313:ASP:N | 29:BD:313:ASP:OD1 | 2.49 | 0.44 |
| 31:BF:133:LYS:HB2 | 31:BF:133:LYS:HE3 | 1.75 | 0.44 |
| 32:BG:81:ASN:HB3 | 32:BG:82:GLU:OE1 | 2.18 | 0.44 |
| 34:BI:44:PHE:CB | 34:BI:137:ALA:HB1 | 2.48 | 0.44 |
| 34:BI:51:ILE:HG12 | 34:BI:162:ARG:O | 2.16 | 0.44 |
| 37:BL:73:ASP:OD1 | 37:BL:112:LYS:HG2 | 2.17 | 0.44 |
| 38:BM:41:ILE:HG21 | 38:BM:65:ARG:HH12 | 1.82 | 0.44 |
| 44:BS:33:VAL:O | 44:BS:37:ILE:HG23 | 2.18 | 0.44 |
| 55:Bd:33:LYS:HB2 | 55:Bd:33:LYS:HE3 | 1.70 | 0.44 |
| 1:AA:306:G:O6 | 10:AK:24:LYS:NZ | 2.50 | 0.44 |
| 1:AA:503:A:H61 | 1:AA:829:U:H3 | 1.66 | 0.44 |
| 1:AA:708:A:H2' | 1:AA:709:A:C8 | 2.52 | 0.44 |
| 1:AA:768:U:H4' | 9:AJ:20:ILE:CD1 | 2.47 | 0.44 |
| 1:AA:924:A:H1' | 1:AA:1269:C:O2 | 2.16 | 0.44 |
| 1:AA:1149:G:H4' | 15:AR:20:ARG:HD2 | 1.98 | 0.44 |
| 1:AA:1216:A:C5 | 1:AA:1217:A:H1' | 2.52 | 0.44 |
| 1:AA:1270:G:H2' | 1:AA:1271:C:C6 | 2.53 | 0.44 |
| 4:AD:13:TRP:HZ3 | 12:AN:31:ILE:HD12 | 1.83 | 0.44 |
| 4:AD:13:TRP:CZ3 | 12:AN:31:ILE:HD12 | 2.52 | 0.44 |
| 8:AI:164:GLY:HA2 | 8:AI:169:ASP:CB | 2.45 | 0.44 |
| 9:AJ:77:PRO:HD3 | 17:AS:62:ARG:HH12 | 1.82 | 0.44 |
| 12:AN:42:ALA:O | 12:AN:46:GLU:HG3 | 2.17 | 0.44 |
| 14:AP:31:LEU:HD13 | 14:AP:40:ALA:O | 2.18 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 14:AP:71:PRO:HG2 | 14:AP:74:GLU:CD | 2.43 | 0.44 |
| 17:AS:97:THR:O | 17:AS:98:LYS:HG3 | 2.17 | 0.44 |
| 24:AM:38:VAL:HB | 24:AM:72:LEU:O | 2.17 | 0.44 |
| 31:BF:72:LEU:O | 31:BF:76:LEU:HG | 2.17 | 0.44 |
| 33:BH:8:ASP:OD1 | 33:BH:8:ASP:N | 2.51 | 0.44 |
| 37:BL:11:THR:HG21 | 37:BL:15:GLY:O | 2.17 | 0.44 |
| 50:BY:30:ASP:O | 50:BY:34:ASN:ND2 | 2.50 | 0.44 |
| 1:AA:171:G:O2' | 1:AA:173:A:N7 | 2.45 | 0.44 |
| 1:AA:964:G:H1 | 1:AA:969:U:H3 | 1.65 | 0.44 |
| 1:AA:992:G:OP1 | 1:AA:993:C:H2' | 2.18 | 0.44 |
| 1:AA:1081:G:C5 | 1:AA:1082:G:H1' | 2.52 | 0.44 |
| 1:AA:1340:C:H1' | 1:AA:1440:A:H5'' | 1.98 | 0.44 |
| 4:AD:64:THR:HG22 | 4:AD:87:LEU:HD22 | 1.98 | 0.44 |
| 6:AF:168:PHE:HB3 | 6:AF:219:VAL:CG2 | 2.48 | 0.44 |
| 9:AJ:15:LYS:HZ3 | 9:AJ:71:LYS:HB2 | 1.82 | 0.44 |
| 9:AJ:55:ASP:CG | 21:AX:5:THR:HA | 2.43 | 0.44 |
| 12:AN:4:MET:HE3 | 12:AN:5:LYS:N | 2.33 | 0.44 |
| 14:AP:25:ARG:CZ | 14:AP:28:ASN:HA | 2.48 | 0.44 |
| 18:AU:29:LEU:HD13 | 18:AU:37:LEU:HD11 | 2.00 | 0.44 |
| 21:AX:27:PHE:HE2 | 21:AX:29:SER:HB2 | 1.82 | 0.44 |
| 22:AY:40:GLY:O | 22:AY:43:LYS:HG3 | 2.17 | 0.44 |
| 26:BA:1294:C:OP2 | 30:BE:170:ARG:NH2 | 2.47 | 0.44 |
| 26:BA:2689:G:H4' | 29:BD:11:SER:HB2 | 1.99 | 0.44 |
| 32:BG:142:LYS:HE3 | 32:BG:176:ALA:HB3 | 1.98 | 0.44 |
| 33:BH:31:LYS:HB2 | 33:BH:98:ALA:HB3 | 2.00 | 0.44 |
| 34:BI:48:ILE:CD1 | 34:BI:150:LEU:HD12 | 2.48 | 0.44 |
| 44:BS:126:MET:HE2 | 44:BS:136:LYS:HD2 | 2.00 | 0.44 |
| 1:AA:56:A:H4' | 1:AA:57:G:O5' | 2.18 | 0.44 |
| 1:AA:185:G:C6 | 1:AA:186:U:H1' | 2.52 | 0.44 |
| 1:AA:411:C:N4 | 1:AA:412:G:O6 | 2.51 | 0.44 |
| 1:AA:774:U:H2' | 1:AA:775:C:H6 | 1.82 | 0.44 |
| 1:AA:1099:U:O5' | 24:AM:71:ARG:NH2 | 2.49 | 0.44 |
| 1:AA:1192:G:H21 | 1:AA:1240:A:H62 | 1.65 | 0.44 |
| 1:AA:1328:C:H1' | 1:AA:1329:C:H5 | 1.81 | 0.44 |
| 3:AC:18:ASP:HB2 | 3:AC:33:MET:SD | 2.58 | 0.44 |
| 4:AD:91:TYR:O | 4:AD:95:ILE:HG13 | 2.18 | 0.44 |
| 7:AG:40:ILE:O | 7:AG:44:LEU:HG | 2.17 | 0.44 |
| 8:AI:132:VAL:O | 8:AI:136:LEU:HG | 2.18 | 0.44 |
| 8:AI:186:LYS:HE2 | 8:AI:186:LYS:HA | 2.00 | 0.44 |
| 9:AJ:71:LYS:HE3 | 9:AJ:128:TYR:OH | 2.18 | 0.44 |
| 16:AQ:142:LYS:HE2 | 16:AQ:146:ALA:CB | 2.47 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 20:AW:74:GLU:CA | 20:AW:77:MET:HE2 | 2.47 | 0.44 |
| 26:BA:936:G:C8 | 49:BX:23:MET:HE1 | 2.52 | 0.44 |
| 30:BE:96:LYS:HB3 | 30:BE:98:GLU:CD | 2.43 | 0.44 |
| 32:BG:74:ARG:HA | 32:BG:77:VAL:HG22 | 1.99 | 0.44 |
| 33:BH:34:THR:HA | 33:BH:96:THR:CG2 | 2.48 | 0.44 |
| 35:BJ:60:GLY:HA3 | 35:BJ:65:GLY:HA3 | 2.00 | 0.44 |
| 53:Bb:56:THR:HG23 | 53:Bb:72:THR:HG23 | 1.99 | 0.44 |
| 1:AA:364:A:H2' | 1:AA:365:A:C8 | 2.51 | 0.44 |
| 1:AA:399:A:H2' | 1:AA:400:U:C6 | 2.52 | 0.44 |
| 1:AA:416:G:H2' | 1:AA:417:U:H6 | 1.82 | 0.44 |
| 1:AA:521:G:O6 | 1:AA:699:G:H3' | 2.18 | 0.44 |
| 1:AA:567:U:OP1 | 6:AF:103:ARG:NH2 | 2.51 | 0.44 |
| 1:AA:1304:U:C5' | 15:AR:27:LYS:HD2 | 2.48 | 0.44 |
| 5:AE:85:LEU:HD22 | 5:AE:90:ILE:CD1 | 2.48 | 0.44 |
| 8:AI:56:ARG:HA | 8:AI:59:ASN:ND2 | 2.33 | 0.44 |
| 8:AI:186:LYS:CE | 8:AI:189:ARG:HD3 | 2.46 | 0.44 |
| 11:AL:13:THR:HG22 | 11:AL:15:THR:HG23 | 2.00 | 0.44 |
| 11:AL:85:ARG:HA | 11:AL:88:VAL:HG12 | 2.00 | 0.44 |
| 13:AO:8:ALA:O | 13:AO:12:LYS:HG3 | 2.17 | 0.44 |
| 23:AH:10:ASP:N | 23:AH:15:ARG:O | 2.43 | 0.44 |
| 23:AH:74:ARG:HE | 23:AH:95:MET:HE3 | 1.83 | 0.44 |
| 24:AM:5:ARG:HD3 | 24:AM:72:LEU:HD11 | 1.98 | 0.44 |
| 24:AM:97:GLU:O | 24:AM:98:ILE:HD13 | 2.18 | 0.44 |
| 26:BA:155:A:OP1 | 38:BM:86:LYS:HB3 | 2.18 | 0.44 |
| 26:BA:1233:U:OP1 | 35:BJ:124:LEU:HD21 | 2.18 | 0.44 |
| 26:BA:2807:C:O2 | 26:BA:2880:A:O2' | 2.36 | 0.44 |
| 29:BD:67:GLU:OE1 | 29:BD:328:LEU:HD12 | 2.17 | 0.44 |
| 31:BF:42:ALA:HB2 | 31:BF:57:ILE:HG13 | 1.99 | 0.44 |
| 35:BJ:7:ILE:HD12 | 35:BJ:21:VAL:HG22 | 1.99 | 0.44 |
| 35:BJ:23:LYS:HD2 | 35:BJ:23:LYS:HA | 1.72 | 0.44 |
| 38:BM:10:ARG:N | 38:BM:48:ASP:OD2 | 2.51 | 0.44 |
| 40:BO:108:SER:OG | 40:BO:111:GLU:HG3 | 2.17 | 0.44 |
| 44:BS:11:ASP:O | 44:BS:15:THR:OG1 | 2.27 | 0.44 |
| 1:AA:169:C:O3' | 6:AF:149:THR:HG21 | 2.18 | 0.44 |
| 1:AA:224:U:H2' | 1:AA:227:C:H5 | 1.83 | 0.44 |
| 1:AA:552:U:O2' | 1:AA:553:G:H5' | 2.18 | 0.44 |
| 1:AA:703:G:H5' | 26:BA:800:A:C2 | 2.53 | 0.44 |
| 1:AA:742:U:H4' | 28:BC:238:ARG:OXT | 2.18 | 0.44 |
| 1:AA:790:C:H5'' | 4:AD:127:LYS:NZ | 2.33 | 0.44 |
| 1:AA:1135:G:H4' | 15:AR:48:LYS:NZ | 2.33 | 0.44 |
| 1:AA:1318:G:P | 11:AL:12:LYS:HG2 | 2.58 | 0.44 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 3:AC:37:ARG:H | 24:AM:92:LYS:HZ2 | 1.65 | 0.44 |
| 3:AC:92:ALA:HB2 | 3:AC:182:ILE:CG1 | 2.48 | 0.44 |
| 4:AD:21:ILE:HD12 | 4:AD:78:ALA:O | 2.18 | 0.44 |
| 4:AD:115:TYR:CE2 | 4:AD:193:ILE:HA | 2.53 | 0.44 |
| 8:AI:165:ALA:HB2 | 8:AI:174:SER:OG | 2.18 | 0.44 |
| 9:AJ:53:ILE:HG23 | 9:AJ:60:ILE:CG1 | 2.48 | 0.44 |
| 13:AO:10:ILE:O | 13:AO:14:THR:HG23 | 2.18 | 0.44 |
| 16:AQ:75:LEU:HB3 | 16:AQ:81:ALA:HB2 | 2.00 | 0.44 |
| 16:AQ:135:LEU:HD13 | 16:AQ:139:TRP:CZ3 | 2.52 | 0.44 |
| 19:AV:66:GLY:O | 19:AV:68:ILE:HG13 | 2.18 | 0.44 |
| 20:AW:5:ILE:HG12 | 20:AW:21:PHE:CB | 2.46 | 0.44 |
| 20:AW:6:LEU:HD13 | 20:AW:22:VAL:HG23 | 1.99 | 0.44 |
| 20:AW:19:LEU:HB2 | 20:AW:68:ALA:C | 2.43 | 0.44 |
| 20:AW:32:ARG:O | 20:AW:36:ARG:HG3 | 2.18 | 0.44 |
| 22:AY:42:ASP:O | 22:AY:45:ARG:HD3 | 2.18 | 0.44 |
| 22:AY:73:LEU:H | 22:AY:73:LEU:HD23 | 1.83 | 0.44 |
| 26:BA:441:A:P | 46:BU:6:SER:HA | 2.58 | 0.44 |
| 27:BB:29:C:H2' | 27:BB:30:U:C6 | 2.53 | 0.44 |
| 29:BD:83:ALA:N | 29:BD:144:VAL:O | 2.50 | 0.44 |
| 30:BE:4:ALA:N | 30:BE:17:VAL:O | 2.35 | 0.44 |
| 30:BE:74:LEU:HD11 | 30:BE:79:ARG:HE | 1.83 | 0.44 |
| 32:BG:39:TYR:CE2 | 32:BG:65:MET:HG3 | 2.53 | 0.44 |
| 33:BH:41:ILE:CD1 | 33:BH:68:SER:HB2 | 2.43 | 0.44 |
| 39:BN:120:LEU:O | 39:BN:124:VAL:HG23 | 2.18 | 0.44 |
| 45:BT:6:TYR:HA | 48:BW:29:VAL:HG13 | 1.99 | 0.44 |
| 49:BX:41:TYR:O | 49:BX:45:VAL:HG23 | 2.17 | 0.44 |
| 51:BZ:23:ALA:O | 51:BZ:65:GLU:HA | 2.18 | 0.44 |
| 52:Ba:113:ILE:HD11 | 52:Ba:143:LYS:HD3 | 2.00 | 0.44 |
| 1:AA:132:C:O2 | 23:AH:7:VAL:HG11 | 2.18 | 0.43 |
| 1:AA:140:U:H5'' | 23:AH:82:GLY:O | 2.18 | 0.43 |
| 1:AA:461:G:OP2 | 13:AO:66:LYS:NZ | 2.49 | 0.43 |
| 1:AA:930:C:H2' | 1:AA:931:C:O4' | 2.18 | 0.43 |
| 1:AA:1062:U:O2' | 24:AM:39:PRO:HD2 | 2.18 | 0.43 |
| 1:AA:1082:G:H2' | 1:AA:1083:G:H8 | 1.83 | 0.43 |
| 1:AA:1246:A:O2' | 1:AA:1247:G:H4' | 2.18 | 0.43 |
| 2:AB:101:THR:OG1 | 2:AB:208:PRO:HG2 | 2.18 | 0.43 |
| 3:AC:9:ASN:O | 3:AC:13:VAL:HG13 | 2.17 | 0.43 |
| 3:AC:127:GLY:HA2 | 3:AC:149:GLY:HA3 | 2.00 | 0.43 |
| 4:AD:48:VAL:HG22 | 12:AN:26:THR:HG22 | 2.00 | 0.43 |
| 7:AG:132:VAL:CG2 | 7:AG:143:LEU:HB2 | 2.48 | 0.43 |
| 10:AK:61:ALA:HB2 | 10:AK:76:ILE:HD11 | 2.00 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 11:AL:5:VAL:H | 11:AL:18:ALA:HB3 | 1.83 | 0.43 |
| 16:AQ:50:ILE:O | 16:AQ:54:LEU:HG | 2.18 | 0.43 |
| 16:AQ:91:LEU:HB3 | 16:AQ:122:ILE:HG12 | 2.00 | 0.43 |
| 17:AS:56:MET:HB2 | 17:AS:63:TYR:CE1 | 2.53 | 0.43 |
| 18:AU:42:GLU:HB3 | 18:AU:75:ILE:HD13 | 2.00 | 0.43 |
| 19:AV:83:ASN:HB2 | 19:AV:86:LYS:HB2 | 1.99 | 0.43 |
| 26:BA:503:A:O2' | 44:BS:3:ARG:NH2 | 2.51 | 0.43 |
| 26:BA:1018:U:OP1 | 26:BA:1019:A:O2' | 2.33 | 0.43 |
| 26:BA:1089:G:OP1 | 27:BB:89:U:O2' | 2.27 | 0.43 |
| 26:BA:1129:C:OP1 | 32:BG:60:LYS:HG3 | 2.17 | 0.43 |
| 29:BD:30:TRP:HZ3 | 29:BD:41:SER:HB2 | 1.83 | 0.43 |
| 36:BK:49:ILE:HD11 | 36:BK:75:ARG:C | 2.43 | 0.43 |
| 40:BO:69:LEU:HA | 40:BO:87:ALA:O | 2.18 | 0.43 |
| 51:BZ:7:LYS:HE3 | 51:BZ:42:MET:SD | 2.58 | 0.43 |
| 51:BZ:8:GLU:HG2 | 51:BZ:76:VAL:HG12 | 1.99 | 0.43 |
| 52:Ba:126:ILE:HG23 | 52:Ba:149:ILE:CG2 | 2.47 | 0.43 |
| 57:Bf:70:ARG:HA | 57:Bf:77:ALA:HA | 1.99 | 0.43 |
| 1:AA:171:G:C2 | 1:AA:173:A:H5'' | 2.53 | 0.43 |
| 1:AA:242:G:H1' | 10:AK:2:ARG:NH2 | 2.34 | 0.43 |
| 1:AA:339:G:O2' | 1:AA:340:G:H5' | 2.19 | 0.43 |
| 1:AA:351:U:O2' | 1:AA:352:U:H5' | 2.18 | 0.43 |
| 1:AA:524:G:H2' | 1:AA:525:C:O4' | 2.18 | 0.43 |
| 1:AA:751:C:O2' | 1:AA:752:C:H5' | 2.18 | 0.43 |
| 1:AA:1194:G:H2' | 1:AA:1195:A:N9 | 2.33 | 0.43 |
| 1:AA:1273:C:OP1 | 19:AV:81:LYS:NZ | 2.30 | 0.43 |
| 1:AA:1289:C:O2' | 11:AL:130:GLN:HA | 2.18 | 0.43 |
| 4:AD:21:ILE:CG2 | 4:AD:33:GLY:H | 2.31 | 0.43 |
| 4:AD:171:TYR:O | 4:AD:175:LYS:HB3 | 2.18 | 0.43 |
| 7:AG:151:GLY:N | 7:AG:169:ASP:OD1 | 2.31 | 0.43 |
| 9:AJ:95:PRO:HG3 | 9:AJ:130:TYR:CE2 | 2.53 | 0.43 |
| 10:AK:80:ILE:HD11 | 10:AK:101:ARG:HB2 | 1.99 | 0.43 |
| 14:AP:26:ILE:HG22 | 14:AP:27:MET:HG2 | 2.00 | 0.43 |
| 14:AP:55:ILE:HG13 | 14:AP:82:ILE:CD1 | 2.48 | 0.43 |
| 17:AS:27:LEU:HD21 | 17:AS:91:CYS:HB3 | 2.00 | 0.43 |
| 18:AU:34:PHE:O | 18:AU:37:LEU:HB2 | 2.18 | 0.43 |
| 19:AV:106:GLU:CD | 19:AV:118:ARG:HD3 | 2.42 | 0.43 |
| 26:BA:170:G:H2' | 38:BM:193:SER:HB2 | 1.99 | 0.43 |
| 26:BA:214:C:OP2 | 26:BA:2407:C:O2' | 2.32 | 0.43 |
| 26:BA:1696:C:O2 | 26:BA:2710:U:O2' | 2.35 | 0.43 |
| 26:BA:2687:C:H4' | 36:BK:40:VAL:HB | 1.99 | 0.43 |
| 29:BD:113:ARG:HB2 | 29:BD:159:VAL:CG2 | 2.49 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 31:BF:138:ALA:O | 31:BF:139:ARG:HD2 | 2.17 | 0.43 |
| 37:BL:70:GLY:O | 37:BL:74:GLU:HG3 | 2.18 | 0.43 |
| 38:BM:155:SER:O | 38:BM:162:ARG:NH2 | 2.51 | 0.43 |
| 39:BN:83:TYR:CD1 | 39:BN:143:ILE:HD13 | 2.54 | 0.43 |
| 46:BU:47:VAL:HG22 | 46:BU:103:ILE:CD1 | 2.43 | 0.43 |
| 55:Bd:37:HIS:CD2 | 55:Bd:39:ARG:HB2 | 2.53 | 0.43 |
| 1:AA:63:U:H2' | 1:AA:64:G:C8 | 2.53 | 0.43 |
| 1:AA:115:U:O4' | 6:AF:135:LEU:HD11 | 2.17 | 0.43 |
| 1:AA:559:U:C2 | 5:AE:147:ILE:HD11 | 2.53 | 0.43 |
| 1:AA:692:C:H2' | 1:AA:693:G:O4' | 2.18 | 0.43 |
| 1:AA:854:A:H2' | 1:AA:855:C:O4' | 2.18 | 0.43 |
| 1:AA:1081:G:C6 | 1:AA:1082:G:H1' | 2.53 | 0.43 |
| 1:AA:1153:G:H4' | 3:AC:172:LYS:O | 2.19 | 0.43 |
| 1:AA:1183:A:H2' | 1:AA:1184:C:C5 | 2.54 | 0.43 |
| 1:AA:1465:C:O2' | 1:AA:1466:U:H5' | 2.18 | 0.43 |
| 4:AD:51:THR:O | 4:AD:55:LEU:HD13 | 2.18 | 0.43 |
| 9:AJ:71:LYS:HE3 | 9:AJ:128:TYR:CE2 | 2.53 | 0.43 |
| 10:AK:74:ALA:HB1 | 10:AK:103:SER:OG | 2.18 | 0.43 |
| 13:AO:85:GLY:O | 13:AO:88:ALA:HB2 | 2.17 | 0.43 |
| 14:AP:92:TRP:CZ2 | 19:AV:35:VAL:HG22 | 2.54 | 0.43 |
| 19:AV:137:LEU:HD23 | 19:AV:140:LEU:CD1 | 2.48 | 0.43 |
| 20:AW:25:TYR:OH | 20:AW:57:THR:HG21 | 2.17 | 0.43 |
| 23:AH:19:ILE:HG22 | 23:AH:21:ILE:HG12 | 2.00 | 0.43 |
| 26:BA:444:C:O2' | 26:BA:1336:A:N7 | 2.51 | 0.43 |
| 26:BA:1775:G:N2 | 26:BA:1778:G:OP2 | 2.45 | 0.43 |
| 30:BE:203:THR:HB | 30:BE:204:PRO:HD2 | 2.00 | 0.43 |
| 35:BJ:33:TYR:HA | 35:BJ:99:LYS:O | 2.18 | 0.43 |
| 39:BN:29:LEU:HD13 | 43:BR:70:ARG:NH1 | 2.33 | 0.43 |
| 39:BN:155:SER:O | 39:BN:158:LEU:HD23 | 2.18 | 0.43 |
| 48:BW:4:LEU:HD12 | 48:BW:4:LEU:O | 2.19 | 0.43 |
| 1:AA:50:C:H2' | 1:AA:334:C:N4 | 2.31 | 0.43 |
| 1:AA:82:G:O2' | 1:AA:360:G:O2' | 2.13 | 0.43 |
| 1:AA:135:G:O2' | 1:AA:137:A:N7 | 2.39 | 0.43 |
| 1:AA:172:G:OP2 | 6:AF:195:LYS:HE3 | 2.18 | 0.43 |
| 1:AA:334:C:H4' | 1:AA:336:G:OP1 | 2.19 | 0.43 |
| 1:AA:403:U:H2' | 1:AA:404:G:H8 | 1.83 | 0.43 |
| 1:AA:487:A:H4' | 1:AA:488:G:O5' | 2.18 | 0.43 |
| 1:AA:779:C:OP1 | 21:AX:50:LYS:HG2 | 2.18 | 0.43 |
| 1:AA:955:C:C6 | 1:AA:970:G:H1' | 2.54 | 0.43 |
| 1:AA:1056:G:H1' | 11:AL:113:ARG:NH2 | 2.33 | 0.43 |
| 1:AA:1259:G:H3' | 18:AU:40:SER:OG | 2.18 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:1382:G:O2' | 1:AA:1383:G:H5' | 2.18 | 0.43 |
| 2:AB:168:ILE:HG12 | 2:AB:188:VAL:HG11 | 1.99 | 0.43 |
| 4:AD:12:GLY:O | 4:AD:16:LYS:HG2 | 2.18 | 0.43 |
| 9:AJ:20:ILE:HB | 9:AJ:22:LYS:HG3 | 1.99 | 0.43 |
| 10:AK:111:SER:HB2 | 10:AK:116:ASP:OD2 | 2.18 | 0.43 |
| 12:AN:70:GLY:HA2 | 12:AN:103:ARG:O | 2.18 | 0.43 |
| 20:AW:19:LEU:HD11 | 20:AW:43:LEU:HD13 | 2.00 | 0.43 |
| 20:AW:45:ALA:HB1 | 20:AW:50:LEU:HB2 | 1.99 | 0.43 |
| 22:AY:33:ILE:HG12 | 22:AY:49:ARG:O | 2.19 | 0.43 |
| 24:AM:50:ARG:HD2 | 24:AM:61:TRP:NE1 | 2.29 | 0.43 |
| 26:BA:322:U:OP2 | 30:BE:152:THR:HG22 | 2.17 | 0.43 |
| 26:BA:651:C:H4' | 30:BE:108:LYS:HD3 | 1.99 | 0.43 |
| 26:BA:1712:U:OP2 | 51:BZ:20:LYS:HD3 | 2.18 | 0.43 |
| 26:BA:1771:C:OP2 | 41:BP:72:TYR:OH | 2.32 | 0.43 |
| 31:BF:45:THR:O | 31:BF:46:LEU:HD23 | 2.19 | 0.43 |
| 31:BF:88:PHE:CZ | 31:BF:153:ILE:HG22 | 2.54 | 0.43 |
| 35:BJ:81:ARG:HG3 | 35:BJ:95:MET:HE2 | 2.00 | 0.43 |
| 36:BK:132:VAL:HG11 | 47:BV:22:VAL:CG2 | 2.41 | 0.43 |
| 38:BM:15:ASN:O | 38:BM:19:THR:HG23 | 2.19 | 0.43 |
| 39:BN:33:ASP:OD1 | 39:BN:50:ALA:HB2 | 2.19 | 0.43 |
| 40:BO:96:PHE:HE1 | 40:BO:106:CYS:HB2 | 1.83 | 0.43 |
| 42:BQ:32:LYS:HE2 | 42:BQ:32:LYS:HB2 | 1.88 | 0.43 |
| 45:BT:39:VAL:HG11 | 45:BT:64:LEU:HD22 | 1.99 | 0.43 |
| 49:BX:48:VAL:O | 49:BX:52:VAL:HG22 | 2.18 | 0.43 |
| 1:AA:10:G:O2' | 7:AG:66:ARG:HD3 | 2.18 | 0.43 |
| 1:AA:705:C:H2' | 1:AA:706:G:O4' | 2.17 | 0.43 |
| 1:AA:721:G:N1 | 1:AA:742:U:OP2 | 2.27 | 0.43 |
| 1:AA:810:A:O2' | 1:AA:811:G:H5' | 2.18 | 0.43 |
| 1:AA:952:G:H2' | 1:AA:953:G:O4' | 2.17 | 0.43 |
| 1:AA:1125:G:N2 | 1:AA:1128:G:N7 | 2.55 | 0.43 |
| 1:AA:1317:G:C8 | 11:AL:115:LYS:HE3 | 2.52 | 0.43 |
| 3:AC:25:LEU:O | 3:AC:25:LEU:HD23 | 2.19 | 0.43 |
| 4:AD:22:GLU:OE1 | 4:AD:79:THR:HG22 | 2.19 | 0.43 |
| 6:AF:4:GLN:O | 6:AF:22:TRP:HA | 2.18 | 0.43 |
| 6:AF:130:LYS:HB3 | 6:AF:142:ILE:HG21 | 2.00 | 0.43 |
| 7:AG:8:VAL:HB | 7:AG:10:LYS:NZ | 2.34 | 0.43 |
| 7:AG:111:ILE:HA | 7:AG:196:ASN:ND2 | 2.34 | 0.43 |
| 12:AN:94:ILE:HG21 | 12:AN:107:ILE:CD1 | 2.48 | 0.43 |
| 14:AP:100:LEU:HD22 | 18:AU:19:ARG:NH1 | 2.34 | 0.43 |
| 15:AR:16:LYS:HB2 | 15:AR:32:LEU:CD2 | 2.48 | 0.43 |
| 16:AQ:6:THR:HG21 | 16:AQ:8:ARG:NH1 | 2.33 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 16:AQ:142:LYS:HE2 | 16:AQ:146:ALA:HB2 | 2.01 | 0.43 |
| 17:AS:95:SER:OG | 17:AS:98:LYS:HD3 | 2.18 | 0.43 |
| 21:AX:18:ASN:HB2 | 21:AX:55:THR:CG2 | 2.48 | 0.43 |
| 23:AH:76:LEU:HA | 23:AH:92:ARG:O | 2.18 | 0.43 |
| 26:BA:28:C:H4' | 46:BU:14:LYS:HB2 | 1.99 | 0.43 |
| 26:BA:325:A:N3 | 46:BU:4:MET:HG3 | 2.34 | 0.43 |
| 26:BA:2376:G:O2' | 26:BA:2377:G:O5' | 2.25 | 0.43 |
| 26:BA:2629:A:O2' | 29:BD:245:ARG:HB2 | 2.18 | 0.43 |
| 26:BA:2637:U:H3 | 26:BA:2786:A:H62 | 1.67 | 0.43 |
| 28:BC:50:PRO:O | 28:BC:183:ARG:NH1 | 2.52 | 0.43 |
| 31:BF:98:ILE:HD11 | 31:BF:118:MET:HE2 | 2.00 | 0.43 |
| 31:BF:100:GLU:HG2 | 31:BF:115:VAL:HG21 | 2.00 | 0.43 |
| 31:BF:110:ASP:OD2 | 31:BF:112:ASN:HB2 | 2.18 | 0.43 |
| 38:BM:67:ASN:OD1 | 38:BM:127:HIS:HB3 | 2.19 | 0.43 |
| 41:BP:126:LYS:HD3 | 41:BP:131:GLU:HG3 | 2.00 | 0.43 |
| 50:BY:9:ASP:O | 50:BY:13:ILE:HG13 | 2.17 | 0.43 |
| 1:AA:767:C:C4' | 9:AJ:13:THR:HG22 | 2.45 | 0.43 |
| 1:AA:914:A:C5 | 1:AA:1001:U:H4' | 2.54 | 0.43 |
| 1:AA:915:A:H4' | 24:AM:56:ASP:OD1 | 2.18 | 0.43 |
| 1:AA:964:G:OP1 | 1:AA:964:G:H3' | 2.17 | 0.43 |
| 1:AA:1267:C:H2' | 1:AA:1268:U:O4' | 2.18 | 0.43 |
| 5:AE:26:GLU:OE1 | 5:AE:38:LYS:NZ | 2.41 | 0.43 |
| 7:AG:13:LEU:O | 7:AG:17:VAL:HG13 | 2.18 | 0.43 |
| 9:AJ:53:ILE:CD1 | 21:AX:4:TYR:HB2 | 2.46 | 0.43 |
| 10:AK:82:ASN:ND2 | 10:AK:95:THR:OG1 | 2.52 | 0.43 |
| 11:AL:32:ILE:CG2 | 11:AL:37:TYR:HA | 2.49 | 0.43 |
| 14:AP:50:ARG:HH21 | 19:AV:35:VAL:HA | 1.83 | 0.43 |
| 20:AW:14:LEU:O | 20:AW:14:LEU:HD23 | 2.19 | 0.43 |
| 22:AY:63:LEU:HB2 | 22:AY:66:THR:CG2 | 2.47 | 0.43 |
| 26:BA:340:U:H5' | 46:BU:3:ALA:HA | 1.99 | 0.43 |
| 26:BA:1311:G:H4' | 52:Ba:98:LYS:HE3 | 2.00 | 0.43 |
| 28:BC:83:CYS:SG | 28:BC:84:GLY:N | 2.92 | 0.43 |
| 30:BE:20:PRO:HB3 | 30:BE:249:GLU:OE1 | 2.18 | 0.43 |
| 30:BE:111:ARG:HD3 | 30:BE:252:PHE:O | 2.19 | 0.43 |
| 31:BF:13:MET:HE2 | 31:BF:25:ALA:HB1 | 1.99 | 0.43 |
| 32:BG:153:GLU:OE1 | 32:BG:172:ILE:HG13 | 2.19 | 0.43 |
| 32:BG:164:ARG:CD | 56:Be:44:LYS:HA | 2.48 | 0.43 |
| 34:BI:23:MET:SD | 34:BI:94:VAL:HG21 | 2.58 | 0.43 |
| 46:BU:15:ALA:O | 46:BU:19:ALA:HB2 | 2.19 | 0.43 |
| 46:BU:28:MET:HA | 46:BU:98:PRO:HG3 | 2.00 | 0.43 |
| 49:BX:137:VAL:O | 49:BX:141:HIS:NE2 | 2.48 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 1:AA:1035:C:H2' | 1:AA:1036:A:H8 | 1.81 | 0.43 |
| 1:AA:1086:G:H4' | 1:AA:1087:A:C8 | 2.54 | 0.43 |
| 1:AA:1263:G:H2' | 1:AA:1265:A:OP2 | 2.18 | 0.43 |
| 1:AA:1325:C:O2' | 8:AI:45:GLN:HG2 | 2.19 | 0.43 |
| 2:AB:74:THR:HG21 | 2:AB:197:ASN:ND2 | 2.33 | 0.43 |
| 2:AB:131:HIS:CD2 | 2:AB:245:THR:HA | 2.54 | 0.43 |
| 4:AD:37:ALA:CB | 4:AD:42:LEU:HD22 | 2.46 | 0.43 |
| 6:AF:132:GLN:HG3 | 6:AF:134:ASN:H | 1.84 | 0.43 |
| 6:AF:156:SER:O | 6:AF:157:ILE:HD13 | 2.19 | 0.43 |
| 7:AG:192:LEU:HD23 | 7:AG:192:LEU:HA | 1.87 | 0.43 |
| 8:AI:21:LEU:CD2 | 22:AY:60:ILE:HG13 | 2.49 | 0.43 |
| 9:AJ:52:PHE:CZ | 9:AJ:54:ASP:HB3 | 2.53 | 0.43 |
| 13:AO:1:MET:HA | 13:AO:11:LEU:CD2 | 2.49 | 0.43 |
| 14:AP:46:GLY:O | 14:AP:107:HIS:NE2 | 2.51 | 0.43 |
| 14:AP:133:LEU:HA | 14:AP:136:GLU:CG | 2.49 | 0.43 |
| 16:AQ:30:ALA:O | 16:AQ:34:THR:HG22 | 2.19 | 0.43 |
| 16:AQ:56:ASP:O | 21:AX:34:ILE:HD13 | 2.18 | 0.43 |
| 19:AV:81:LYS:HB3 | 19:AV:89:HIS:O | 2.19 | 0.43 |
| 20:AW:57:THR:OG1 | 20:AW:64:SER:OG | 2.15 | 0.43 |
| 23:AH:45:LEU:HD22 | 23:AH:123:LEU:HB2 | 2.00 | 0.43 |
| 23:AH:49:LYS:HB2 | 23:AH:114:GLU:CB | 2.43 | 0.43 |
| 24:AM:79:GLU:O | 24:AM:82:LEU:HG | 2.17 | 0.43 |
| 25:AT:57:LYS:HA | 25:AT:60:ARG:NH1 | 2.33 | 0.43 |
| 26:BA:893:G:OP2 | 37:BL:18:LYS:HE2 | 2.19 | 0.43 |
| 26:BA:2293:C:OP1 | 57:Bf:61:PRO:HB2 | 2.18 | 0.43 |
| 26:BA:2344:U:C2 | 39:BN:107:LEU:HD21 | 2.53 | 0.43 |
| 27:BB:6:U:O2' | 27:BB:7:G:H5' | 2.19 | 0.43 |
| 27:BB:31:C:H5' | 31:BF:131:ILE:HD11 | 2.00 | 0.43 |
| 29:BD:281:GLU:OE1 | 29:BD:333:GLU:HA | 2.19 | 0.43 |
| 33:BH:14:THR:O | 33:BH:18:LEU:HD23 | 2.19 | 0.43 |
| 42:BQ:29:GLN:HB2 | 42:BQ:33:ASN:HD22 | 1.84 | 0.43 |
| 46:BU:29:GLY:HA2 | 46:BU:46:ALA:HA | 1.99 | 0.43 |
| 51:BZ:36:GLY:O | 51:BZ:40:ARG:HG2 | 2.18 | 0.43 |
| 55:Bd:37:HIS:HB2 | 55:Bd:38:PRO:HD2 | 2.01 | 0.43 |
| 1:AA:40:U:O2' | 1:AA:41:C:H5' | 2.19 | 0.43 |
| 1:AA:72:U:H4' | 1:AA:73:G:OP2 | 2.19 | 0.43 |
| 1:AA:143:G:O2' | 1:AA:144:G:H5' | 2.19 | 0.43 |
| 1:AA:151:C:H2' | 1:AA:152:C:C6 | 2.54 | 0.43 |
| 1:AA:173:A:H4' | 17:AS:3:ARG:HB2 | 2.00 | 0.43 |
| 1:AA:1039:C:H5'' | 2:AB:148:ARG:HE | 1.83 | 0.43 |
| 1:AA:1247:G:H1 | 8:AI:149:LYS:HA | 1.83 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|-------------------|--------------------------|-------------------|
| 1:AA:1402:U:H2' | 1:AA:1403:U:C6 | 2.54 | 0.43 |
| 3:AC:18:ASP:OD2 | 3:AC:33:MET:HB3 | 2.19 | 0.43 |
| 3:AC:120:VAL:HG11 | 3:AC:182:ILE:CD1 | 2.48 | 0.43 |
| 4:AD:16:LYS:HD3 | 4:AD:36:MET:SD | 2.58 | 0.43 |
| 6:AF:98:GLN:HA | 6:AF:103:ARG:O | 2.19 | 0.43 |
| 8:AI:137:ARG:HG3 | 8:AI:141:MET:HE2 | 2.00 | 0.43 |
| 9:AJ:100:GLY:HA3 | 9:AJ:130:TYR:HB3 | 2.00 | 0.43 |
| 11:AL:47:SER:HA | 11:AL:50:LEU:HD21 | 2.01 | 0.43 |
| 14:AP:29:THR:HG22 | 14:AP:31:LEU:HD23 | 2.00 | 0.43 |
| 17:AS:43:MET:HE3 | 17:AS:46:THR:CG2 | 2.49 | 0.43 |
| 22:AY:16:GLU:CB | 22:AY:60:ILE:HD13 | 2.49 | 0.43 |
| 24:AM:48:PRO:HB2 | 24:AM:61:TRP:CE3 | 2.54 | 0.43 |
| 25:AT:31:ASN:O | 25:AT:35:VAL:HG23 | 2.19 | 0.43 |
| 26:BA:21:G:O2' | 26:BA:22:A:OP2 | 2.36 | 0.43 |
| 26:BA:1252:C:OP1 | 49:BX:117:ARG:HB3 | 2.19 | 0.43 |
| 26:BA:1754:G:OP2 | 53:Bb:9:GLY:HA2 | 2.19 | 0.43 |
| 27:BB:122:A:H2' | 27:BB:123:A:C8 | 2.54 | 0.43 |
| 29:BD:14:PHE:O | 29:BD:17:ARG:HG2 | 2.19 | 0.43 |
| 29:BD:70:VAL:CG1 | 29:BD:298:LYS:HG3 | 2.49 | 0.43 |
| 30:BE:151:ARG:HE | 30:BE:151:ARG:HB3 | 1.63 | 0.43 |
| 40:BO:64:GLU:OE1 | 40:BO:83:PRO:HD2 | 2.18 | 0.43 |
| 40:BO:119:GLY:HA2 | 40:BO:122:ILE:CD1 | 2.46 | 0.43 |
| 41:BP:26:PRO:HG2 | 41:BP:27:GLU:OE1 | 2.18 | 0.43 |
| 47:BV:13:LEU:CD1 | 47:BV:18:GLY:HA3 | 2.48 | 0.43 |
| 47:BV:47:ARG:HB2 | 47:BV:58:GLN:NE2 | 2.34 | 0.43 |
| 1:AA:134:G:C2' | 1:AA:135:G:H5' | 2.49 | 0.43 |
| 1:AA:175:U:C4 | 17:AS:28:PRO:HB3 | 2.54 | 0.43 |
| 1:AA:301:A:O2' | 1:AA:1405:G:H4' | 2.19 | 0.43 |
| 1:AA:322:C:O2' | 1:AA:323:C:H5' | 2.19 | 0.43 |
| 1:AA:345:A:H2' | 1:AA:346:A:O4' | 2.18 | 0.43 |
| 1:AA:358:G:O2' | 1:AA:359:C:H5' | 2.18 | 0.43 |
| 1:AA:684:A:H2' | 1:AA:685:A:C8 | 2.53 | 0.43 |
| 1:AA:1006:A:H3' | 1:AA:1033:G:P | 2.59 | 0.43 |
| 1:AA:1064:U:O4' | 24:AM:72:LEU:HD23 | 2.19 | 0.43 |
| 1:AA:1156:C:H2' | 1:AA:1157:C:C6 | 2.53 | 0.43 |
| 3:AC:28:ALA:HB1 | 3:AC:54:VAL:HG23 | 2.01 | 0.43 |
| 3:AC:87:ARG:CD | 3:AC:88:PRO:HD2 | 2.48 | 0.43 |
| 4:AD:117:ILE:HD11 | 4:AD:154:PHE:CD1 | 2.54 | 0.43 |
| 4:AD:160:GLU:O | 4:AD:165:ARG:N | 2.52 | 0.43 |
| 7:AG:13:LEU:CD1 | 7:AG:35:ILE:HG12 | 2.47 | 0.43 |
| 7:AG:84:GLY:CA | 7:AG:108:ILE:HD13 | 2.48 | 0.43 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 13:AO:39:LEU:HD22 | 13:AO:44:GLN:O | 2.18 | 0.43 |
| 16:AQ:37:VAL:CG2 | 16:AQ:54:LEU:HD11 | 2.48 | 0.43 |
| 21:AX:15:VAL:HG11 | 21:AX:43:LEU:HD13 | 2.00 | 0.43 |
| 23:AH:25:GLU:HG3 | 23:AH:42:ILE:HA | 2.00 | 0.43 |
| 24:AM:42:THR:HG23 | 24:AM:68:VAL:C | 2.43 | 0.43 |
| 26:BA:613:A:O2' | 52:Ba:55:CYS:SG | 2.76 | 0.43 |
| 26:BA:2023:G:OP1 | 51:BZ:22:PRO:HA | 2.18 | 0.43 |
| 26:BA:2835:G:H4' | 29:BD:281:GLU:HB3 | 1.99 | 0.43 |
| 30:BE:22:VAL:HG23 | 30:BE:115:ARG:CG | 2.43 | 0.43 |
| 30:BE:193:LYS:HA | 30:BE:193:LYS:HD2 | 1.87 | 0.43 |
| 33:BH:53:GLU:OE2 | 33:BH:77:PHE:HB3 | 2.18 | 0.43 |
| 44:BS:8:ILE:HD12 | 44:BS:113:HIS:CE1 | 2.54 | 0.43 |
| 46:BU:32:LEU:CD2 | 46:BU:103:ILE:HD12 | 2.48 | 0.43 |
| 1:AA:151:C:H2' | 1:AA:152:C:H6 | 1.84 | 0.43 |
| 1:AA:385:A:H5' | 5:AE:148:PRO:HG2 | 2.01 | 0.43 |
| 1:AA:440:G:H1 | 1:AA:484:C:H42 | 1.67 | 0.43 |
| 1:AA:633:U:H1' | 1:AA:636:A:N7 | 2.34 | 0.43 |
| 1:AA:770:G:H2' | 1:AA:771:G:C8 | 2.54 | 0.43 |
| 1:AA:825:C:OP2 | 13:AO:2:ALA:HB2 | 2.19 | 0.43 |
| 1:AA:1141:G:C1' | 7:AG:64:SER:HB2 | 2.38 | 0.43 |
| 1:AA:1424:G:H2' | 1:AA:1425:G:C8 | 2.54 | 0.43 |
| 6:AF:148:GLY:N | 6:AF:151:ASP:OD2 | 2.49 | 0.43 |
| 6:AF:168:PHE:HB3 | 6:AF:219:VAL:HG11 | 2.01 | 0.43 |
| 8:AI:57:LEU:CD1 | 8:AI:136:LEU:HD22 | 2.49 | 0.43 |
| 13:AO:83:CYS:SG | 13:AO:92:ILE:HD13 | 2.58 | 0.43 |
| 16:AQ:118:THR:O | 16:AQ:122:ILE:HG13 | 2.19 | 0.43 |
| 23:AH:36:ASP:O | 23:AH:51:GLN:HA | 2.19 | 0.43 |
| 24:AM:7:ARG:C | 24:AM:8:LEU:HD12 | 2.43 | 0.43 |
| 24:AM:50:ARG:HG3 | 24:AM:60:THR:O | 2.19 | 0.43 |
| 26:BA:323:A:OP2 | 30:BE:211:ASN:HB2 | 2.19 | 0.43 |
| 26:BA:1299:C:N3 | 52:Ba:135:GLN:HG3 | 2.33 | 0.43 |
| 28:BC:110:GLU:HG2 | 28:BC:117:GLY:N | 2.33 | 0.43 |
| 32:BG:44:ILE:CG1 | 32:BG:53:VAL:HG12 | 2.48 | 0.43 |
| 33:BH:37:ALA:O | 33:BH:41:ILE:HG13 | 2.19 | 0.43 |
| 36:BK:19:ALA:O | 36:BK:21:VAL:HG13 | 2.19 | 0.43 |
| 39:BN:83:TYR:HE2 | 39:BN:166:LYS:HE2 | 1.84 | 0.43 |
| 40:BO:55:ILE:HG13 | 40:BO:98:LYS:HD2 | 1.99 | 0.43 |
| 46:BU:77:ILE:HD11 | 46:BU:101:VAL:CG1 | 2.49 | 0.43 |
| 1:AA:360:G:H2' | 1:AA:361:G:H5' | 2.01 | 0.42 |
| 1:AA:721:G:OP1 | 12:AN:121:LYS:HG3 | 2.19 | 0.42 |
| 1:AA:759:G:O2' | 1:AA:760:A:H5' | 2.19 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:817:A:O2' | 1:AA:818:A:H3' | 2.19 | 0.42 |
| 1:AA:1197:A:H1' | 1:AA:1317:G:O2' | 2.19 | 0.42 |
| 2:AB:77:LYS:HA | 2:AB:196:THR:O | 2.18 | 0.42 |
| 3:AC:136:LEU:H | 3:AC:136:LEU:HD23 | 1.84 | 0.42 |
| 6:AF:83:ASP:O | 6:AF:95:ARG:HA | 2.19 | 0.42 |
| 11:AL:47:SER:HA | 11:AL:50:LEU:CD2 | 2.49 | 0.42 |
| 14:AP:99:ASP:HB3 | 14:AP:106:LYS:NZ | 2.34 | 0.42 |
| 14:AP:108:LEU:HB2 | 14:AP:113:ILE:HD11 | 2.01 | 0.42 |
| 14:AP:134:ARG:NH1 | 14:AP:139:LEU:HD23 | 2.33 | 0.42 |
| 18:AU:106:ARG:O | 18:AU:109:GLU:HG3 | 2.18 | 0.42 |
| 20:AW:49:LEU:HD22 | 20:AW:71:TYR:O | 2.19 | 0.42 |
| 20:AW:51:ILE:HG13 | 20:AW:89:ASN:ND2 | 2.30 | 0.42 |
| 20:AW:83:GLU:O | 20:AW:87:LYS:HG2 | 2.19 | 0.42 |
| 24:AM:8:LEU:HD23 | 24:AM:94:ILE:CG2 | 2.48 | 0.42 |
| 25:AT:20:TYR:HB3 | 25:AT:23:VAL:HG22 | 2.00 | 0.42 |
| 26:BA:189:U:OP2 | 26:BA:408:G:N2 | 2.46 | 0.42 |
| 28:BC:237:LYS:HD3 | 28:BC:237:LYS:HA | 1.86 | 0.42 |
| 33:BH:55:ILE:CG2 | 33:BH:60:ILE:HB | 2.49 | 0.42 |
| 40:BO:68:LEU:HD13 | 40:BO:84:VAL:CG1 | 2.49 | 0.42 |
| 43:BR:53:PRO:HB3 | 43:BR:85:VAL:CG1 | 2.49 | 0.42 |
| 46:BU:55:VAL:HB | 46:BU:60:PHE:O | 2.19 | 0.42 |
| 46:BU:60:PHE:CG | 46:BU:82:VAL:HG22 | 2.54 | 0.42 |
| 51:BZ:39:VAL:HG13 | 51:BZ:44:THR:O | 2.19 | 0.42 |
| 52:Ba:110:ILE:CG1 | 52:Ba:126:ILE:HB | 2.49 | 0.42 |
| 1:AA:303:U:H4' | 1:AA:1383:G:H5' | 2.02 | 0.42 |
| 1:AA:583:A:H2' | 1:AA:584:C:O4' | 2.19 | 0.42 |
| 1:AA:691:U:O4' | 16:AQ:48:ALA:HA | 2.19 | 0.42 |
| 1:AA:711:C:O2' | 1:AA:844:C:N3 | 2.50 | 0.42 |
| 1:AA:905:A:H2' | 1:AA:906:U:O4' | 2.19 | 0.42 |
| 1:AA:927:U:H3' | 1:AA:928:C:C6 | 2.54 | 0.42 |
| 1:AA:957:G:O2' | 1:AA:958:G:H5' | 2.19 | 0.42 |
| 1:AA:1045:G:H2' | 1:AA:1046:C:C6 | 2.54 | 0.42 |
| 1:AA:1272:C:O2' | 1:AA:1273:C:H5' | 2.19 | 0.42 |
| 4:AD:180:LEU:HD23 | 4:AD:183:VAL:HG22 | 2.01 | 0.42 |
| 7:AG:15:LYS:HA | 7:AG:18:VAL:HG12 | 2.00 | 0.42 |
| 7:AG:111:ILE:CD1 | 7:AG:192:LEU:HD22 | 2.48 | 0.42 |
| 8:AI:6:LYS:NZ | 8:AI:10:LYS:O | 2.32 | 0.42 |
| 13:AO:54:GLY:HA2 | 13:AO:66:LYS:HA | 2.00 | 0.42 |
| 16:AQ:31:ASP:HA | 16:AQ:34:THR:HG22 | 2.01 | 0.42 |
| 16:AQ:102:LEU:HD12 | 16:AQ:115:LEU:HD22 | 2.02 | 0.42 |
| 16:AQ:130:LYS:NZ | 16:AQ:139:TRP:O | 2.49 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 19:AV:74:ARG:HD2 | 19:AV:95:GLY:N | 2.34 | 0.42 |
| 26:BA:1754:G:C8 | 53:Bb:20:PRO:HA | 2.54 | 0.42 |
| 26:BA:2691:G:H5' | 29:BD:261:ARG:HG3 | 2.00 | 0.42 |
| 29:BD:47:VAL:HB | 29:BD:74:VAL:HG13 | 2.00 | 0.42 |
| 29:BD:102:TRP:O | 29:BD:122:GLN:NE2 | 2.40 | 0.42 |
| 30:BE:143:VAL:HG13 | 30:BE:145:ASP:OD1 | 2.18 | 0.42 |
| 31:BF:84:ASN:OD1 | 31:BF:87:GLN:HG2 | 2.18 | 0.42 |
| 32:BG:129:LYS:HB2 | 32:BG:136:THR:CG2 | 2.49 | 0.42 |
| 35:BJ:38:GLU:OE1 | 35:BJ:38:GLU:N | 2.51 | 0.42 |
| 35:BJ:45:SER:HB3 | 35:BJ:128:ARG:NH2 | 2.35 | 0.42 |
| 44:BS:96:GLU:O | 44:BS:100:GLU:HG3 | 2.19 | 0.42 |
| 44:BS:128:ARG:HG3 | 44:BS:134:THR:CG2 | 2.47 | 0.42 |
| 46:BU:53:VAL:O | 46:BU:64:GLU:HA | 2.20 | 0.42 |
| 1:AA:109:U:H5'' | 1:AA:110:G:OP1 | 2.19 | 0.42 |
| 1:AA:173:A:H5' | 17:AS:3:ARG:HG3 | 2.00 | 0.42 |
| 1:AA:962:A:H2' | 1:AA:963:A:O4' | 2.19 | 0.42 |
| 1:AA:1053:C:H4' | 15:AR:49:TYR:CZ | 2.54 | 0.42 |
| 1:AA:1297:A:H2' | 1:AA:1298:U:O4' | 2.20 | 0.42 |
| 1:AA:1399:G:H4' | 47:BV:47:ARG:CZ | 2.49 | 0.42 |
| 3:AC:93:GLN:OE1 | 3:AC:93:GLN:N | 2.50 | 0.42 |
| 3:AC:157:PRO:HA | 3:AC:161:VAL:CB | 2.44 | 0.42 |
| 5:AE:18:TRP:HA | 5:AE:23:MET:SD | 2.59 | 0.42 |
| 6:AF:168:PHE:CD1 | 6:AF:219:VAL:HG21 | 2.54 | 0.42 |
| 7:AG:17:VAL:HG11 | 7:AG:40:ILE:CG2 | 2.49 | 0.42 |
| 11:AL:32:ILE:HG22 | 11:AL:37:TYR:HA | 2.01 | 0.42 |
| 11:AL:35:GLU:OE1 | 19:AV:143:ILE:HG12 | 2.19 | 0.42 |
| 19:AV:36:LYS:HA | 19:AV:44:PRO:HA | 2.01 | 0.42 |
| 22:AY:16:GLU:HB3 | 22:AY:60:ILE:HD13 | 2.00 | 0.42 |
| 26:BA:480:A:OP1 | 46:BU:92:VAL:HG11 | 2.19 | 0.42 |
| 26:BA:944:A:O2' | 26:BA:2278:A:N3 | 2.44 | 0.42 |
| 29:BD:146:TYR:HA | 29:BD:161:ASP:O | 2.19 | 0.42 |
| 33:BH:38:THR:O | 33:BH:42:GLU:HG3 | 2.19 | 0.42 |
| 37:BL:77:SER:O | 37:BL:80:VAL:HG22 | 2.19 | 0.42 |
| 38:BM:77:ARG:NH2 | 38:BM:87:MET:HE2 | 2.35 | 0.42 |
| 49:BX:57:ILE:HG13 | 49:BX:146:ASN:OD1 | 2.20 | 0.42 |
| 1:AA:63:U:H1' | 1:AA:148:A:H1' | 2.02 | 0.42 |
| 1:AA:98:C:H5'' | 1:AA:293:G:O2' | 2.19 | 0.42 |
| 1:AA:458:C:H4' | 1:AA:459:C:H5'' | 2.01 | 0.42 |
| 1:AA:614:G:O2' | 1:AA:615:G:H5' | 2.18 | 0.42 |
| 1:AA:918:C:H2' | 1:AA:919:C:C6 | 2.54 | 0.42 |
| 1:AA:1267:C:H5' | 18:AU:106:ARG:HG3 | 2.01 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:1333:C:H2' | 1:AA:1334:U:C6 | 2.54 | 0.42 |
| 2:AB:78:THR:OG1 | 2:AB:81:MET:HG2 | 2.19 | 0.42 |
| 3:AC:156:ASN:HB2 | 3:AC:157:PRO:HD3 | 2.00 | 0.42 |
| 3:AC:164:GLU:OE2 | 3:AC:166:PHE:HB2 | 2.19 | 0.42 |
| 5:AE:100:ILE:O | 5:AE:103:LEU:HB2 | 2.19 | 0.42 |
| 8:AI:2:ILE:HG23 | 8:AI:5:TYR:CE2 | 2.54 | 0.42 |
| 9:AJ:54:ASP:OD1 | 9:AJ:54:ASP:N | 2.51 | 0.42 |
| 16:AQ:17:PRO:HB3 | 21:AX:8:PRO:CG | 2.50 | 0.42 |
| 26:BA:1679:G:O3' | 26:BA:1680:C:H4' | 2.20 | 0.42 |
| 30:BE:149:LEU:HD12 | 30:BE:149:LEU:O | 2.20 | 0.42 |
| 33:BH:50:LEU:HG | 33:BH:76:ILE:CG2 | 2.49 | 0.42 |
| 34:BI:20:ARG:NH2 | 34:BI:26:VAL:HB | 2.34 | 0.42 |
| 40:BO:35:ILE:CD1 | 40:BO:124:ILE:HD11 | 2.49 | 0.42 |
| 41:BP:23:TRP:HB2 | 41:BP:53:LYS:CG | 2.48 | 0.42 |
| 43:BR:43:ASP:HA | 43:BR:58:GLN:HE22 | 1.84 | 0.42 |
| 48:BW:24:LEU:HD12 | 48:BW:57:ILE:HD12 | 2.02 | 0.42 |
| 49:BX:81:GLU:O | 49:BX:84:ARG:HG2 | 2.20 | 0.42 |
| 1:AA:585:U:H2' | 1:AA:586:G:C8 | 2.54 | 0.42 |
| 1:AA:683:G:H2' | 1:AA:684:A:O4' | 2.18 | 0.42 |
| 1:AA:761:U:H4' | 1:AA:762:G:OP2 | 2.20 | 0.42 |
| 2:AB:218:SER:HB3 | 2:AB:244:GLU:HA | 2.01 | 0.42 |
| 4:AD:147:LYS:O | 4:AD:151:GLU:HB2 | 2.19 | 0.42 |
| 7:AG:83:VAL:HG21 | 7:AG:165:ALA:HB2 | 2.02 | 0.42 |
| 10:AK:34:THR:OG1 | 10:AK:94:LEU:HB2 | 2.19 | 0.42 |
| 11:AL:35:GLU:HG3 | 11:AL:36:LEU:HG | 2.01 | 0.42 |
| 11:AL:79:VAL:O | 11:AL:83:VAL:HG23 | 2.20 | 0.42 |
| 12:AN:91:GLN:O | 12:AN:95:ARG:HG2 | 2.20 | 0.42 |
| 13:AO:129:LEU:O | 13:AO:133:VAL:HG23 | 2.20 | 0.42 |
| 14:AP:116:THR:O | 14:AP:119:GLU:HG3 | 2.19 | 0.42 |
| 23:AH:70:PRO:CA | 23:AH:100:GLU:HB2 | 2.47 | 0.42 |
| 23:AH:122:ALA:HA | 23:AH:126:LEU:O | 2.20 | 0.42 |
| 26:BA:2344:U:H3 | 39:BN:107:LEU:HD21 | 1.84 | 0.42 |
| 27:BB:123:A:H2' | 27:BB:124:G:O4' | 2.20 | 0.42 |
| 30:BE:2:ALA:O | 30:BE:19:LEU:N | 2.51 | 0.42 |
| 32:BG:141:ASN:HB3 | 32:BG:144:ASP:HB2 | 2.00 | 0.42 |
| 33:BH:58:ALA:HB3 | 38:BM:24:LEU:CD1 | 2.49 | 0.42 |
| 36:BK:56:SER:HA | 36:BK:67:GLN:O | 2.18 | 0.42 |
| 38:BM:99:GLN:O | 38:BM:103:GLU:HG3 | 2.20 | 0.42 |
| 42:BQ:13:PHE:O | 42:BQ:19:TRP:HA | 2.19 | 0.42 |
| 53:Bb:57:GLY:HA2 | 53:Bb:71:GLY:O | 2.19 | 0.42 |
| 1:AA:230:G:N2 | 10:AK:12:THR:O | 2.51 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:300:G:H2' | 1:AA:301:A:H8 | 1.85 | 0.42 |
| 1:AA:671:G:N2 | 1:AA:706:G:H5'' | 2.35 | 0.42 |
| 1:AA:988:A:H4' | 1:AA:989:G:H5'' | 2.02 | 0.42 |
| 1:AA:996:G:H3' | 1:AA:997:G:C8 | 2.55 | 0.42 |
| 1:AA:1016:G:N2 | 1:AA:1019:A:OP2 | 2.35 | 0.42 |
| 1:AA:1113:G:H3' | 1:AA:1114:U:H5'' | 2.01 | 0.42 |
| 1:AA:1295:U:H5'' | 11:AL:125:ALA:CB | 2.48 | 0.42 |
| 1:AA:1377:G:H2' | 1:AA:1378:U:C6 | 2.55 | 0.42 |
| 2:AB:71:HIS:HB2 | 2:AB:95:VAL:CG1 | 2.48 | 0.42 |
| 2:AB:218:SER:CB | 2:AB:244:GLU:HA | 2.49 | 0.42 |
| 4:AD:90:ASP:HA | 4:AD:93:ARG:HG2 | 2.00 | 0.42 |
| 4:AD:186:ARG:HG3 | 4:AD:187:LYS:HG2 | 2.00 | 0.42 |
| 10:AK:55:LEU:HD21 | 10:AK:58:SER:HB2 | 2.02 | 0.42 |
| 12:AN:84:ARG:HE | 12:AN:84:ARG:HB2 | 1.56 | 0.42 |
| 14:AP:94:LEU:HD12 | 14:AP:105:ASP:HB3 | 2.02 | 0.42 |
| 15:AR:34:ARG:NH1 | 24:AM:50:ARG:O | 2.53 | 0.42 |
| 16:AQ:3:LYS:HB2 | 16:AQ:6:THR:O | 2.19 | 0.42 |
| 19:AV:74:ARG:HE | 19:AV:92:LYS:HG2 | 1.84 | 0.42 |
| 20:AW:1:MET:HG2 | 20:AW:25:TYR:HA | 2.01 | 0.42 |
| 26:BA:1057:U:O2' | 49:BX:12:ASN:HA | 2.19 | 0.42 |
| 26:BA:1612:A:O2' | 26:BA:1613:A:O5' | 2.37 | 0.42 |
| 26:BA:2053:C:OP1 | 26:BA:2465:G:O2' | 2.32 | 0.42 |
| 26:BA:2391:A:O2' | 39:BN:30:SER:OG | 2.13 | 0.42 |
| 26:BA:2444:A:C4 | 57:Bf:54:LYS:HG2 | 2.54 | 0.42 |
| 27:BB:18:C:C4' | 39:BN:1:MET:HA | 2.49 | 0.42 |
| 29:BD:89:ALA:HB2 | 29:BD:98:ILE:HD12 | 2.01 | 0.42 |
| 29:BD:268:ILE:HG12 | 29:BD:297:ILE:HD13 | 2.02 | 0.42 |
| 32:BG:128:VAL:O | 32:BG:129:LYS:HD3 | 2.20 | 0.42 |
| 34:BI:58:ILE:HG22 | 34:BI:63:LEU:HG | 2.01 | 0.42 |
| 34:BI:86:LYS:O | 34:BI:134:PHE:HA | 2.20 | 0.42 |
| 35:BJ:76:LEU:O | 35:BJ:80:ILE:HG13 | 2.19 | 0.42 |
| 38:BM:82:ARG:HB2 | 38:BM:87:MET:SD | 2.60 | 0.42 |
| 58:AZ:47:THR:CG2 | 58:AZ:54:LYS:HE2 | 2.49 | 0.42 |
| 1:AA:291:C:H4' | 6:AF:7:LEU:HB2 | 2.00 | 0.42 |
| 1:AA:293:G:O2' | 1:AA:294:G:H5' | 2.20 | 0.42 |
| 1:AA:509:U:H4' | 1:AA:514:A:C6 | 2.55 | 0.42 |
| 1:AA:801:C:H2' | 1:AA:802:C:H6 | 1.85 | 0.42 |
| 1:AA:960:C:O2' | 1:AA:963:A:N6 | 2.45 | 0.42 |
| 1:AA:1257:A:H2' | 1:AA:1258:G:H5' | 2.02 | 0.42 |
| 1:AA:1285:G:H2' | 1:AA:1286:A:C8 | 2.54 | 0.42 |
| 1:AA:1395:U:C3' | 1:AA:1396:G:H5' | 2.50 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 3:AC:95:MET:CE | 3:AC:116:THR:HG23 | 2.49 | 0.42 |
| 4:AD:86:GLU:HA | 4:AD:186:ARG:O | 2.20 | 0.42 |
| 5:AE:110:GLU:HA | 5:AE:115:THR:CG2 | 2.41 | 0.42 |
| 6:AF:119:ARG:HD2 | 6:AF:218:ILE:HG23 | 2.00 | 0.42 |
| 6:AF:156:SER:OG | 6:AF:157:ILE:N | 2.53 | 0.42 |
| 7:AG:24:SER:O | 7:AG:27:GLU:HG2 | 2.20 | 0.42 |
| 8:AI:6:LYS:HE2 | 8:AI:12:ASP:H | 1.84 | 0.42 |
| 8:AI:20:ASP:HB2 | 8:AI:102:ALA:CB | 2.49 | 0.42 |
| 8:AI:85:VAL:O | 8:AI:89:THR:OG1 | 2.23 | 0.42 |
| 14:AP:44:LEU:CD2 | 14:AP:109:LEU:HD11 | 2.49 | 0.42 |
| 16:AQ:4:MET:HG3 | 16:AQ:124:ARG:NH1 | 2.35 | 0.42 |
| 16:AQ:25:TRP:CE3 | 21:AX:61:LEU:HD22 | 2.54 | 0.42 |
| 16:AQ:75:LEU:CD2 | 16:AQ:80:VAL:HG23 | 2.49 | 0.42 |
| 18:AU:69:THR:HG21 | 18:AU:107:PHE:CE2 | 2.55 | 0.42 |
| 19:AV:23:ASP:HB3 | 19:AV:25:LYS:HZ3 | 1.85 | 0.42 |
| 21:AX:33:LYS:HB2 | 21:AX:33:LYS:HE2 | 1.79 | 0.42 |
| 23:AH:118:LYS:HB2 | 23:AH:123:LEU:CD1 | 2.43 | 0.42 |
| 24:AM:45:LEU:N | 24:AM:66:MET:O | 2.52 | 0.42 |
| 26:BA:2346:A:N6 | 43:BR:47:SER:O | 2.53 | 0.42 |
| 29:BD:20:ALA:CB | 29:BD:204:ILE:HD12 | 2.50 | 0.42 |
| 46:BU:1:MET:O | 46:BU:2:ILE:HG13 | 2.18 | 0.42 |
| 55:Bd:44:ARG:O | 55:Bd:45:ARG:HG3 | 2.20 | 0.42 |
| 1:AA:510:C:O2' | 1:AA:760:A:H2' | 2.20 | 0.42 |
| 1:AA:733:A:H4' | 1:AA:734:U:H5' | 2.02 | 0.42 |
| 1:AA:765:C:OP1 | 16:AQ:1:MET:HA | 2.20 | 0.42 |
| 1:AA:896:U:H2' | 1:AA:897:G:H8 | 1.84 | 0.42 |
| 1:AA:992:G:O6 | 1:AA:1146:U:H2' | 2.20 | 0.42 |
| 1:AA:1090:G:H2' | 1:AA:1091:G:H8 | 1.83 | 0.42 |
| 1:AA:1223:G:H2' | 1:AA:1224:G:H8 | 1.83 | 0.42 |
| 4:AD:121:PRO:HB2 | 4:AD:141:MET:SD | 2.60 | 0.42 |
| 4:AD:160:GLU:CA | 4:AD:165:ARG:HB2 | 2.47 | 0.42 |
| 6:AF:125:THR:HA | 6:AF:130:LYS:O | 2.20 | 0.42 |
| 13:AO:1:MET:HA | 13:AO:11:LEU:HD21 | 2.01 | 0.42 |
| 14:AP:133:LEU:O | 14:AP:136:GLU:HG3 | 2.19 | 0.42 |
| 16:AQ:121:LYS:O | 16:AQ:125:LEU:HD23 | 2.20 | 0.42 |
| 16:AQ:149:MET:HE3 | 50:BY:13:ILE:HD11 | 2.02 | 0.42 |
| 18:AU:81:ILE:HD12 | 18:AU:98:LEU:HB3 | 2.01 | 0.42 |
| 22:AY:14:ALA:O | 22:AY:37:VAL:HG13 | 2.19 | 0.42 |
| 23:AH:17:TYR:CE1 | 23:AH:121:LYS:HG2 | 2.55 | 0.42 |
| 23:AH:45:LEU:CD2 | 23:AH:124:LEU:HG | 2.45 | 0.42 |
| 24:AM:36:GLY:O | 24:AM:38:VAL:HG23 | 2.20 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 26:BA:70:C:OP1 | 48:BW:56:ARG:NE | 2.53 | 0.42 |
| 26:BA:1469:A:C6 | 55:Bd:3:HIS:HB3 | 2.54 | 0.42 |
| 28:BC:200:HIS:CE1 | 28:BC:226:VAL:HG21 | 2.55 | 0.42 |
| 29:BD:111:LYS:HB2 | 29:BD:116:ILE:HD11 | 2.02 | 0.42 |
| 32:BG:6:ALA:HB2 | 32:BG:54:ASP:OD1 | 2.19 | 0.42 |
| 37:BL:86:GLU:OE2 | 37:BL:88:LYS:NZ | 2.47 | 0.42 |
| 38:BM:95:GLY:HA3 | 38:BM:196:LYS:NZ | 2.35 | 0.42 |
| 40:BO:39:ILE:CD1 | 40:BO:69:LEU:HD22 | 2.50 | 0.42 |
| 40:BO:91:PHE:CZ | 40:BO:106:CYS:HB3 | 2.55 | 0.42 |
| 47:BV:8:PHE:CZ | 47:BV:39:ASN:HB3 | 2.55 | 0.42 |
| 1:AA:434:A:H4' | 1:AA:435:G:O5' | 2.20 | 0.42 |
| 1:AA:963:A:N1 | 1:AA:965:A:H1' | 2.34 | 0.42 |
| 1:AA:1215:A:O2' | 1:AA:1273:C:H4' | 2.19 | 0.42 |
| 1:AA:1236:A:OP1 | 19:AV:80:LYS:HD2 | 2.20 | 0.42 |
| 1:AA:1441:A:H5' | 1:AA:1469:A:H1' | 2.02 | 0.42 |
| 2:AB:156:ASN:O | 2:AB:159:ILE:HG22 | 2.20 | 0.42 |
| 2:AB:164:GLU:OE2 | 7:AG:11:THR:HB | 2.20 | 0.42 |
| 5:AE:142:GLY:O | 20:AW:61:MET:HB3 | 2.19 | 0.42 |
| 6:AF:115:ASN:OD1 | 6:AF:163:VAL:HG21 | 2.20 | 0.42 |
| 7:AG:38:PRO:HG2 | 7:AG:39:GLN:OE1 | 2.19 | 0.42 |
| 7:AG:74:VAL:HG12 | 7:AG:100:ALA:CB | 2.50 | 0.42 |
| 8:AI:13:PRO:HA | 8:AI:94:ILE:CD1 | 2.49 | 0.42 |
| 8:AI:55:GLU:O | 8:AI:59:ASN:ND2 | 2.53 | 0.42 |
| 8:AI:70:LYS:HB3 | 8:AI:73:VAL:HG12 | 2.00 | 0.42 |
| 11:AL:13:THR:HB | 11:AL:70:GLY:HA2 | 2.01 | 0.42 |
| 14:AP:25:ARG:HD2 | 14:AP:26:ILE:N | 2.35 | 0.42 |
| 19:AV:27:VAL:O | 19:AV:29:PRO:HD3 | 2.20 | 0.42 |
| 19:AV:74:ARG:HB3 | 19:AV:93:GLY:H | 1.85 | 0.42 |
| 20:AW:3:ILE:HD11 | 20:AW:23:VAL:HG13 | 2.02 | 0.42 |
| 28:BC:110:GLU:HG2 | 28:BC:117:GLY:H | 1.85 | 0.42 |
| 28:BC:207:TRP:CZ3 | 28:BC:209:HIS:HB3 | 2.55 | 0.42 |
| 30:BE:27:TYR:OH | 30:BE:126:LEU:HD22 | 2.20 | 0.42 |
| 36:BK:16:ALA:O | 36:BK:30:GLU:HA | 2.19 | 0.42 |
| 36:BK:73:VAL:CA | 36:BK:95:MET:HG2 | 2.48 | 0.42 |
| 45:BT:5:ASN:HB2 | 45:BT:25:ILE:O | 2.20 | 0.42 |
| 51:BZ:23:ALA:O | 51:BZ:26:ARG:HG3 | 2.20 | 0.42 |
| 52:Ba:132:ILE:HG22 | 52:Ba:133:GLY:O | 2.19 | 0.42 |
| 1:AA:88:C:O2 | 1:AA:335:A:O2' | 2.38 | 0.42 |
| 1:AA:701:G:H5'' | 1:AA:702:G:N7 | 2.35 | 0.42 |
| 1:AA:1004:C:N4 | 1:AA:1137:C:H1' | 2.35 | 0.42 |
| 1:AA:1108:C:OP1 | 25:AT:56:ARG:HG2 | 2.20 | 0.42 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 1:AA:1197:A:O2' | 1:AA:1317:G:H4' | 2.20 | 0.42 |
| 3:AC:53:MET:O | 3:AC:61:VAL:HG21 | 2.19 | 0.42 |
| 5:AE:91:ILE:HG23 | 5:AE:92:LYS:O | 2.20 | 0.42 |
| 8:AI:56:ARG:HA | 8:AI:59:ASN:HD21 | 1.85 | 0.42 |
| 9:AJ:33:ASN:O | 9:AJ:37:VAL:HG23 | 2.20 | 0.42 |
| 13:AO:53:VAL:O | 13:AO:67:CYS:N | 2.39 | 0.42 |
| 13:AO:111:MET:HE3 | 13:AO:111:MET:HB3 | 1.95 | 0.42 |
| 14:AP:130:TYR:HB2 | 18:AU:109:GLU:O | 2.20 | 0.42 |
| 15:AR:24:LEU:HD23 | 15:AR:24:LEU:H | 1.83 | 0.42 |
| 15:AR:34:ARG:HB2 | 24:AM:49:THR:HG23 | 2.02 | 0.42 |
| 20:AW:35:VAL:O | 20:AW:39:LEU:HD23 | 2.19 | 0.42 |
| 24:AM:16:LEU:CD1 | 24:AM:94:ILE:HG21 | 2.48 | 0.42 |
| 26:BA:678:C:O2' | 26:BA:740:U:OP1 | 2.38 | 0.42 |
| 26:BA:793:U:O4 | 53:Bb:1:MET:N | 2.53 | 0.42 |
| 26:BA:2068:U:OP1 | 44:BS:66:ASP:HA | 2.20 | 0.42 |
| 27:BB:1:G:O2' | 27:BB:2:A:H5' | 2.20 | 0.42 |
| 27:BB:66:C:O2' | 27:BB:67:C:H5' | 2.19 | 0.42 |
| 29:BD:88:TYR:HB2 | 29:BD:140:ASP:OD1 | 2.20 | 0.42 |
| 30:BE:7:ILE:HG12 | 30:BE:141:VAL:O | 2.19 | 0.42 |
| 31:BF:43:LYS:O | 31:BF:44:ARG:HG3 | 2.19 | 0.42 |
| 32:BG:22:ASP:CB | 32:BG:36:LYS:HE3 | 2.50 | 0.42 |
| 34:BI:136:VAL:CG2 | 34:BI:150:LEU:HD11 | 2.50 | 0.42 |
| 37:BL:68:ASN:CG | 37:BL:107:SER:HB3 | 2.45 | 0.42 |
| 49:BX:61:THR:OG1 | 49:BX:146:ASN:HB3 | 2.19 | 0.42 |
| 1:AA:108:C:H5' | 1:AA:244:A:O2' | 2.20 | 0.41 |
| 1:AA:157:A:H1' | 1:AA:187:A:C2 | 2.55 | 0.41 |
| 1:AA:206:A:H2' | 1:AA:207:A:C8 | 2.55 | 0.41 |
| 1:AA:295:A:H5'' | 10:AK:20:ARG:HG3 | 2.02 | 0.41 |
| 1:AA:827:C:H2' | 1:AA:828:C:C6 | 2.55 | 0.41 |
| 1:AA:927:U:OP2 | 1:AA:928:C:H3' | 2.19 | 0.41 |
| 1:AA:1112:U:H2' | 1:AA:1113:G:O4' | 2.20 | 0.41 |
| 1:AA:1297:A:O2' | 8:AI:68:THR:HG21 | 2.20 | 0.41 |
| 7:AG:85:LEU:CD1 | 7:AG:186:LYS:HA | 2.50 | 0.41 |
| 9:AJ:34:ILE:O | 9:AJ:38:LEU:HG | 2.20 | 0.41 |
| 9:AJ:49:ASP:HB3 | 9:AJ:64:THR:HB | 2.02 | 0.41 |
| 13:AO:10:ILE:H | 13:AO:10:ILE:HD12 | 1.84 | 0.41 |
| 18:AU:52:ASP:CG | 18:AU:56:LYS:HE3 | 2.45 | 0.41 |
| 19:AV:29:PRO:O | 19:AV:31:TRP:N | 2.53 | 0.41 |
| 20:AW:25:TYR:CZ | 20:AW:29:THR:HG22 | 2.55 | 0.41 |
| 24:AM:48:PRO:HB3 | 24:AM:63:HIS:ND1 | 2.35 | 0.41 |
| 26:BA:10:G:OP1 | 44:BS:63:LYS:HG3 | 2.20 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 26:BA:390:U:OP2 | 57:Bf:38:ARG:NE | 2.47 | 0.41 |
| 26:BA:692:C:OP2 | 37:BL:107:SER:HA | 2.20 | 0.41 |
| 26:BA:892:U:OP1 | 37:BL:18:LYS:NZ | 2.48 | 0.41 |
| 27:BB:74:C:H2' | 27:BB:75:U:O4' | 2.20 | 0.41 |
| 28:BC:90:LYS:HE2 | 28:BC:93:ASN:HD21 | 1.85 | 0.41 |
| 34:BI:5:PRO:HD2 | 34:BI:8:MET:SD | 2.60 | 0.41 |
| 35:BJ:77:LYS:HE2 | 35:BJ:95:MET:HE3 | 2.01 | 0.41 |
| 37:BL:69:VAL:CG2 | 37:BL:106:GLY:HA2 | 2.50 | 0.41 |
| 39:BN:8:LYS:HE2 | 43:BR:22:ARG:O | 2.19 | 0.41 |
| 53:Bb:80:GLY:O | 53:Bb:84:LEU:HG | 2.18 | 0.41 |
| 1:AA:445:G:N2 | 1:AA:466:C:H1' | 2.35 | 0.41 |
| 1:AA:997:G:H5'' | 3:AC:179:LYS:CE | 2.51 | 0.41 |
| 1:AA:1008:U:H3 | 1:AA:1045:G:H1 | 1.69 | 0.41 |
| 1:AA:1258:G:H5' | 14:AP:97:GLN:O | 2.19 | 0.41 |
| 1:AA:1287:A:H5'' | 1:AA:1288:U:OP2 | 2.20 | 0.41 |
| 2:AB:149:PHE:CZ | 2:AB:179:VAL:HG12 | 2.55 | 0.41 |
| 3:AC:51:PRO:O | 3:AC:55:ILE:HG13 | 2.20 | 0.41 |
| 6:AF:175:MET:HB3 | 6:AF:220:ILE:HG21 | 2.02 | 0.41 |
| 7:AG:37:GLU:HB3 | 7:AG:40:ILE:HG13 | 2.02 | 0.41 |
| 7:AG:57:MET:SD | 7:AG:68:VAL:HG13 | 2.60 | 0.41 |
| 9:AJ:75:ILE:HG13 | 9:AJ:127:ALA:CA | 2.50 | 0.41 |
| 14:AP:71:PRO:HG2 | 14:AP:74:GLU:OE1 | 2.19 | 0.41 |
| 25:AT:26:LYS:HE2 | 25:AT:62:LYS:HD2 | 2.01 | 0.41 |
| 26:BA:941:U:O2' | 43:BR:96:LYS:NZ | 2.50 | 0.41 |
| 26:BA:1636:U:P | 28:BC:182:THR:HG21 | 2.60 | 0.41 |
| 26:BA:2457:G:N2 | 26:BA:2460:U:O2 | 2.40 | 0.41 |
| 27:BB:21:U:H3 | 27:BB:64:G:H22 | 1.67 | 0.41 |
| 27:BB:33:U:H2' | 27:BB:34:G:C8 | 2.53 | 0.41 |
| 28:BC:19:ALA:CB | 28:BC:184:ALA:HB2 | 2.49 | 0.41 |
| 28:BC:90:LYS:HB3 | 28:BC:93:ASN:ND2 | 2.36 | 0.41 |
| 29:BD:163:MET:HB2 | 29:BD:310:ARG:NH1 | 2.35 | 0.41 |
| 43:BR:18:THR:OG1 | 43:BR:21:GLU:HG2 | 2.20 | 0.41 |
| 1:AA:357:U:P | 6:AF:53:SER:H | 2.43 | 0.41 |
| 1:AA:416:G:H21 | 20:AW:29:THR:HG21 | 1.85 | 0.41 |
| 1:AA:425:A:H2' | 1:AA:426:C:H5' | 2.02 | 0.41 |
| 1:AA:482:G:O2' | 1:AA:483:C:H5' | 2.20 | 0.41 |
| 1:AA:502:U:H4' | 1:AA:503:A:O5' | 2.21 | 0.41 |
| 1:AA:927:U:H5' | 15:AR:21:LYS:HB2 | 2.02 | 0.41 |
| 1:AA:1094:C:H2' | 1:AA:1095:U:O4' | 2.20 | 0.41 |
| 1:AA:1202:G:H2' | 1:AA:1224:G:N2 | 2.36 | 0.41 |
| 2:AB:76:GLN:OE1 | 2:AB:76:GLN:N | 2.52 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 2:AB:88:VAL:HG13 | 2:AB:94:TYR:CE1 | 2.54 | 0.41 |
| 2:AB:131:HIS:HB3 | 2:AB:132:PRO:HD3 | 2.01 | 0.41 |
| 3:AC:46:ILE:HG23 | 3:AC:80:ILE:CG2 | 2.48 | 0.41 |
| 3:AC:81:ASP:OD1 | 3:AC:81:ASP:N | 2.52 | 0.41 |
| 3:AC:130:VAL:HG13 | 3:AC:146:PHE:CD1 | 2.55 | 0.41 |
| 4:AD:120:LYS:HD2 | 4:AD:187:LYS:HB2 | 2.02 | 0.41 |
| 6:AF:157:ILE:HD13 | 6:AF:160:LYS:HD3 | 2.03 | 0.41 |
| 8:AI:5:TYR:O | 8:AI:13:PRO:HD2 | 2.19 | 0.41 |
| 10:AK:35:ARG:HG2 | 10:AK:93:ILE:HD13 | 2.02 | 0.41 |
| 13:AO:103:ILE:HD13 | 13:AO:120:LYS:HB3 | 2.02 | 0.41 |
| 15:AR:41:ALA:HB1 | 15:AR:46:PHE:CB | 2.50 | 0.41 |
| 17:AS:11:ALA:HA | 17:AS:12:PRO:HD3 | 1.93 | 0.41 |
| 17:AS:41:SER:HB3 | 17:AS:47:VAL:HG12 | 2.02 | 0.41 |
| 17:AS:41:SER:OG | 17:AS:82:VAL:HG22 | 2.21 | 0.41 |
| 18:AU:35:ALA:CB | 18:AU:46:LEU:HD21 | 2.49 | 0.41 |
| 23:AH:10:ASP:HA | 23:AH:112:ILE:HD12 | 2.02 | 0.41 |
| 26:BA:886:A:C8 | 30:BE:55:MET:HE1 | 2.54 | 0.41 |
| 26:BA:953:C:O2' | 39:BN:6:ARG:NH1 | 2.53 | 0.41 |
| 26:BA:1107:A:O2' | 35:BJ:128:ARG:NH1 | 2.53 | 0.41 |
| 26:BA:1225:G:C5 | 35:BJ:59:ARG:HD3 | 2.55 | 0.41 |
| 27:BB:109:C:O2' | 27:BB:110:U:H5' | 2.20 | 0.41 |
| 30:BE:5:LYS:HE2 | 30:BE:16:GLU:OE2 | 2.20 | 0.41 |
| 34:BI:10:ARG:NH2 | 34:BI:54:GLU:OE1 | 2.28 | 0.41 |
| 38:BM:174:MET:HE2 | 38:BM:185:ARG:HA | 2.02 | 0.41 |
| 39:BN:148:ILE:CD1 | 39:BN:156:SER:HA | 2.43 | 0.41 |
| 41:BP:110:ARG:HD3 | 41:BP:120:TYR:CD2 | 2.55 | 0.41 |
| 52:Ba:119:VAL:HG23 | 52:Ba:124:GLU:OE1 | 2.21 | 0.41 |
| 1:AA:15:U:OP2 | 7:AG:177:THR:OG1 | 2.23 | 0.41 |
| 1:AA:26:A:H2' | 1:AA:27:C:C6 | 2.56 | 0.41 |
| 1:AA:391:A:H4' | 5:AE:124:ARG:NH1 | 2.36 | 0.41 |
| 1:AA:406:C:H4' | 5:AE:173:HIS:CD2 | 2.55 | 0.41 |
| 1:AA:860:A:H2' | 1:AA:861:G:H5' | 2.01 | 0.41 |
| 1:AA:1113:G:H3' | 1:AA:1114:U:C5' | 2.50 | 0.41 |
| 3:AC:156:ASN:O | 3:AC:159:GLU:HB3 | 2.21 | 0.41 |
| 6:AF:97:LEU:HD12 | 6:AF:107:HIS:CG | 2.55 | 0.41 |
| 6:AF:168:PHE:HB3 | 6:AF:219:VAL:HG21 | 2.02 | 0.41 |
| 7:AG:12:ARG:CZ | 7:AG:33:LEU:HD21 | 2.50 | 0.41 |
| 8:AI:16:VAL:HG21 | 8:AI:94:ILE:CG1 | 2.50 | 0.41 |
| 8:AI:125:ASP:OD2 | 22:AY:61:LEU:HB3 | 2.20 | 0.41 |
| 8:AI:143:THR:HG21 | 8:AI:158:LEU:HB2 | 2.02 | 0.41 |
| 8:AI:152:ARG:HB3 | 8:AI:156:GLU:OE1 | 2.20 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 15:AR:11:GLY:HA2 | 15:AR:26:ARG:HH22 | 1.85 | 0.41 |
| 16:AQ:5:HIS:CD2 | 16:AQ:117:LEU:HB3 | 2.56 | 0.41 |
| 22:AY:36:ARG:HG2 | 22:AY:38:LEU:CD2 | 2.50 | 0.41 |
| 26:BA:1856:C:H5 | 28:BC:164:GLY:H | 1.68 | 0.41 |
| 29:BD:80:ILE:O | 29:BD:186:LEU:HA | 2.21 | 0.41 |
| 31:BF:91:PHE:O | 31:BF:147:VAL:HG23 | 2.20 | 0.41 |
| 31:BF:95:SER:OG | 31:BF:121:THR:HB | 2.20 | 0.41 |
| 36:BK:8:ILE:CG2 | 36:BK:49:ILE:HB | 2.44 | 0.41 |
| 37:BL:80:VAL:HA | 37:BL:85:ALA:O | 2.20 | 0.41 |
| 39:BN:24:ARG:O | 39:BN:28:LEU:HG | 2.21 | 0.41 |
| 40:BO:75:LEU:C | 40:BO:95:ALA:HB2 | 2.46 | 0.41 |
| 45:BT:26:VAL:CG2 | 45:BT:60:LYS:HG2 | 2.50 | 0.41 |
| 57:Bf:40:LYS:HB2 | 57:Bf:52:PHE:CE1 | 2.55 | 0.41 |
| 1:AA:428:U:C4' | 20:AW:27:GLY:HA2 | 2.50 | 0.41 |
| 1:AA:944:A:H3' | 1:AA:945:A:C8 | 2.55 | 0.41 |
| 2:AB:168:ILE:O | 2:AB:190:VAL:HA | 2.19 | 0.41 |
| 6:AF:98:GLN:NE2 | 6:AF:102:GLY:HA2 | 2.36 | 0.41 |
| 8:AI:5:TYR:HB2 | 8:AI:31:PRO:CB | 2.42 | 0.41 |
| 10:AK:90:ARG:HE | 10:AK:91:ARG:HH12 | 1.68 | 0.41 |
| 14:AP:45:PRO:O | 14:AP:109:LEU:HG | 2.20 | 0.41 |
| 14:AP:54:LEU:HD21 | 19:AV:34:ASN:HB3 | 2.02 | 0.41 |
| 17:AS:45:ASN:HB3 | 17:AS:73:HIS:CE1 | 2.55 | 0.41 |
| 19:AV:96:SER:O | 19:AV:100:LYS:HG2 | 2.21 | 0.41 |
| 21:AX:15:VAL:HG13 | 21:AX:43:LEU:HD13 | 2.02 | 0.41 |
| 26:BA:1789:G:OP2 | 47:BV:48:ARG:NH2 | 2.53 | 0.41 |
| 28:BC:105:PHE:HB2 | 53:Bb:83:LEU:CD1 | 2.46 | 0.41 |
| 31:BF:33:THR:HG21 | 31:BF:61:VAL:HG11 | 2.01 | 0.41 |
| 46:BU:30:ALA:HB2 | 46:BU:98:PRO:O | 2.20 | 0.41 |
| 1:AA:137:A:H2' | 1:AA:138:A:O4' | 2.20 | 0.41 |
| 1:AA:280:A:H2' | 1:AA:281:G:O4' | 2.20 | 0.41 |
| 1:AA:820:C:O2' | 9:AJ:15:LYS:HD2 | 2.20 | 0.41 |
| 1:AA:992:G:H4' | 1:AA:993:C:O5' | 2.20 | 0.41 |
| 1:AA:1068:C:H2' | 1:AA:1086:G:C5 | 2.56 | 0.41 |
| 1:AA:1086:G:H4' | 1:AA:1087:A:N7 | 2.36 | 0.41 |
| 1:AA:1343:A:H2 | 7:AG:61:MET:HG3 | 1.86 | 0.41 |
| 2:AB:120:ILE:HG23 | 2:AB:167:VAL:CG1 | 2.51 | 0.41 |
| 3:AC:42:THR:HG21 | 3:AC:75:LEU:CD2 | 2.49 | 0.41 |
| 13:AO:103:ILE:CD1 | 13:AO:120:LYS:HB3 | 2.50 | 0.41 |
| 16:AQ:149:MET:CE | 50:BY:74:VAL:HA | 2.51 | 0.41 |
| 17:AS:50:GLU:HA | 17:AS:68:SER:O | 2.19 | 0.41 |
| 18:AU:85:ILE:HD12 | 18:AU:107:PHE:CD2 | 2.55 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 24:AM:8:LEU:HB3 | 24:AM:94:ILE:HG21 | 2.02 | 0.41 |
| 26:BA:895:U:O4 | 37:BL:17:HIS:HB2 | 2.20 | 0.41 |
| 26:BA:1253:A:OP2 | 49:BX:117:ARG:NH1 | 2.50 | 0.41 |
| 26:BA:2654:U:H3 | 26:BA:2780:A:H62 | 1.69 | 0.41 |
| 29:BD:47:VAL:HG22 | 29:BD:287:TYR:CE2 | 2.56 | 0.41 |
| 31:BF:6:VAL:HG22 | 31:BF:124:LEU:HD11 | 2.02 | 0.41 |
| 36:BK:85:GLY:HA3 | 47:BV:16:GLY:HA2 | 2.03 | 0.41 |
| 36:BK:86:LEU:HD21 | 47:BV:20:LEU:HB2 | 2.02 | 0.41 |
| 38:BM:127:HIS:HB2 | 38:BM:129:TRP:CH2 | 2.55 | 0.41 |
| 39:BN:141:GLU:HB2 | 39:BN:146:GLU:CG | 2.47 | 0.41 |
| 43:BR:7:GLU:HG3 | 43:BR:54:ASN:OD1 | 2.21 | 0.41 |
| 49:BX:66:LEU:HD23 | 49:BX:116:PHE:CE2 | 2.55 | 0.41 |
| 52:Ba:141:LEU:HA | 52:Ba:141:LEU:HD23 | 1.85 | 0.41 |
| 1:AA:58:U:H2' | 1:AA:59:C:H6 | 1.85 | 0.41 |
| 1:AA:242:G:H1' | 10:AK:2:ARG:HH21 | 1.85 | 0.41 |
| 1:AA:242:G:C5' | 10:AK:31:SER:HA | 2.39 | 0.41 |
| 1:AA:406:C:H4' | 5:AE:173:HIS:HD2 | 1.85 | 0.41 |
| 1:AA:479:G:H2' | 1:AA:480:C:C6 | 2.55 | 0.41 |
| 1:AA:516:G:H3' | 1:AA:516:G:OP2 | 2.20 | 0.41 |
| 1:AA:526:C:H5'' | 13:AO:4:GLY:CA | 2.49 | 0.41 |
| 1:AA:573:U:O2' | 1:AA:574:U:H5'' | 2.21 | 0.41 |
| 1:AA:583:A:N7 | 9:AJ:107:SER:HA | 2.35 | 0.41 |
| 1:AA:774:U:O2' | 1:AA:775:C:H5' | 2.21 | 0.41 |
| 2:AB:117:PRO:HA | 2:AB:120:ILE:HG12 | 2.01 | 0.41 |
| 3:AC:43:GLN:HA | 3:AC:79:GLN:NE2 | 2.35 | 0.41 |
| 4:AD:18:TRP:CG | 4:AD:36:MET:HB3 | 2.56 | 0.41 |
| 4:AD:39:ASP:CG | 4:AD:42:LEU:HB2 | 2.46 | 0.41 |
| 6:AF:59:ILE:HG23 | 6:AF:64:LYS:HD2 | 2.03 | 0.41 |
| 6:AF:213:ILE:HG13 | 6:AF:214:GLU:OE1 | 2.20 | 0.41 |
| 7:AG:199:ARG:HD3 | 9:AJ:68:ARG:CA | 2.50 | 0.41 |
| 8:AI:164:GLY:HA3 | 8:AI:172:SER:HB3 | 2.02 | 0.41 |
| 13:AO:103:ILE:HD12 | 13:AO:120:LYS:CD | 2.38 | 0.41 |
| 13:AO:128:SER:OG | 13:AO:131:GLN:HB2 | 2.19 | 0.41 |
| 14:AP:70:LEU:HD23 | 14:AP:75:VAL:HG22 | 2.03 | 0.41 |
| 14:AP:108:LEU:HD22 | 14:AP:112:ASP:OD2 | 2.21 | 0.41 |
| 15:AR:24:LEU:HA | 15:AR:33:CYS:HA | 2.03 | 0.41 |
| 17:AS:43:MET:HE1 | 17:AS:71:HIS:CE1 | 2.56 | 0.41 |
| 26:BA:90:C:OP1 | 48:BW:2:ALA:HA | 2.21 | 0.41 |
| 26:BA:1526:A:N6 | 26:BA:1621:U:O2' | 2.53 | 0.41 |
| 26:BA:1817:A:N7 | 54:Bc:2:SER:OG | 2.42 | 0.41 |
| 26:BA:2071:G:N2 | 29:BD:238:LEU:HD13 | 2.36 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 26:BA:2389:A:O4' | 39:BN:136:VAL:HG11 | 2.19 | 0.41 |
| 26:BA:2538:G:H4' | 56:Be:16:ILE:HG21 | 2.03 | 0.41 |
| 26:BA:2777:C:O2' | 35:BJ:140:GLY:HA2 | 2.20 | 0.41 |
| 38:BM:13:TRP:CE2 | 38:BM:46:ARG:HG2 | 2.55 | 0.41 |
| 49:BX:125:HIS:HE1 | 49:BX:128:ILE:HG23 | 1.85 | 0.41 |
| 1:AA:78:U:H2' | 1:AA:79:G:O4' | 2.20 | 0.41 |
| 1:AA:158:A:H2' | 1:AA:159:C:H5' | 2.03 | 0.41 |
| 1:AA:233:G:H1' | 10:AK:8:ARG:NE | 2.35 | 0.41 |
| 1:AA:248:U:O2' | 10:AK:8:ARG:HD3 | 2.20 | 0.41 |
| 1:AA:604:A:H5'' | 4:AD:130:ARG:NE | 2.36 | 0.41 |
| 1:AA:765:C:H2' | 1:AA:766:G:H8 | 1.83 | 0.41 |
| 1:AA:1000:G:H5' | 24:AM:60:THR:HG21 | 2.02 | 0.41 |
| 1:AA:1321:A:H5'' | 8:AI:59:ASN:HB2 | 2.03 | 0.41 |
| 4:AD:136:ALA:HB1 | 4:AD:140:MET:HE1 | 2.03 | 0.41 |
| 4:AD:186:ARG:O | 4:AD:187:LYS:HD3 | 2.20 | 0.41 |
| 5:AE:154:LYS:NZ | 5:AE:157:GLU:OE1 | 2.31 | 0.41 |
| 8:AI:5:TYR:CD2 | 8:AI:31:PRO:HB2 | 2.55 | 0.41 |
| 12:AN:7:ALA:O | 12:AN:71:ILE:HA | 2.21 | 0.41 |
| 14:AP:38:GLN:HG2 | 14:AP:39:TYR:CD1 | 2.56 | 0.41 |
| 15:AR:20:ARG:HH12 | 15:AR:35:HIS:CE1 | 2.39 | 0.41 |
| 16:AQ:98:LEU:HD11 | 16:AQ:111:ASN:HB3 | 2.03 | 0.41 |
| 20:AW:36:ARG:NH2 | 20:AW:90:PRO:HG2 | 2.34 | 0.41 |
| 21:AX:14:ARG:HB2 | 21:AX:61:LEU:CD1 | 2.49 | 0.41 |
| 23:AH:102:ALA:N | 23:AH:105:ILE:HD12 | 2.32 | 0.41 |
| 25:AT:5:ARG:HB3 | 25:AT:6:GLU:H | 1.71 | 0.41 |
| 26:BA:135:U:O2' | 26:BA:136:C:OP1 | 2.29 | 0.41 |
| 26:BA:1656:C:P | 45:BT:32:LYS:HG3 | 2.61 | 0.41 |
| 26:BA:2043:U:O2' | 26:BA:2629:A:OP1 | 2.39 | 0.41 |
| 26:BA:2315:U:OP1 | 31:BF:97:GLY:HA3 | 2.21 | 0.41 |
| 27:BB:110:U:H2' | 27:BB:111:G:C8 | 2.56 | 0.41 |
| 30:BE:10:THR:O | 30:BE:161:ALA:HB1 | 2.20 | 0.41 |
| 31:BF:131:ILE:HG23 | 31:BF:134:ARG:NH2 | 2.27 | 0.41 |
| 35:BJ:72:PRO:HG3 | 35:BJ:136:SER:CB | 2.51 | 0.41 |
| 37:BL:104:VAL:O | 37:BL:105:LEU:HD23 | 2.21 | 0.41 |
| 1:AA:5:U:O2' | 1:AA:446:G:O3' | 2.32 | 0.41 |
| 1:AA:231:U:H4' | 10:AK:13:GLY:CA | 2.39 | 0.41 |
| 1:AA:253:A:H2' | 1:AA:254:C:O4' | 2.21 | 0.41 |
| 1:AA:542:C:O2' | 1:AA:543:C:H5' | 2.21 | 0.41 |
| 1:AA:633:U:H2' | 1:AA:635:A:OP2 | 2.20 | 0.41 |
| 1:AA:697:G:H2' | 1:AA:698:U:O4' | 2.21 | 0.41 |
| 1:AA:766:G:H2' | 1:AA:767:C:O4' | 2.21 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:887:G:H2' | 1:AA:888:G:O4' | 2.21 | 0.41 |
| 1:AA:990:G:H2' | 1:AA:991:U:C6 | 2.56 | 0.41 |
| 1:AA:992:G:H4' | 1:AA:993:C:H5'' | 2.03 | 0.41 |
| 1:AA:1227:A:N7 | 24:AM:41:PRO:HD3 | 2.36 | 0.41 |
| 1:AA:1390:C:O2' | 23:AH:99:LYS:NZ | 2.49 | 0.41 |
| 3:AC:66:THR:OG1 | 3:AC:67:ARG:HD3 | 2.21 | 0.41 |
| 3:AC:153:HIS:O | 3:AC:158:VAL:HG11 | 2.21 | 0.41 |
| 4:AD:107:VAL:CG1 | 4:AD:146:LYS:HE2 | 2.51 | 0.41 |
| 6:AF:127:LYS:HE2 | 6:AF:127:LYS:HB3 | 1.94 | 0.41 |
| 6:AF:174:ALA:HB2 | 6:AF:188:ILE:HG13 | 2.03 | 0.41 |
| 7:AG:25:MET:CG | 7:AG:45:LEU:HD11 | 2.50 | 0.41 |
| 7:AG:26:ASP:OD1 | 7:AG:26:ASP:N | 2.53 | 0.41 |
| 7:AG:128:VAL:HG11 | 7:AG:143:LEU:O | 2.20 | 0.41 |
| 7:AG:192:LEU:O | 7:AG:195:VAL:HB | 2.21 | 0.41 |
| 11:AL:119:ASN:OD1 | 11:AL:120:PHE:N | 2.54 | 0.41 |
| 12:AN:64:LYS:HB2 | 12:AN:64:LYS:HE3 | 1.82 | 0.41 |
| 13:AO:38:PRO:CB | 13:AO:77:ARG:HD3 | 2.51 | 0.41 |
| 16:AQ:50:ILE:O | 16:AQ:53:VAL:HG12 | 2.21 | 0.41 |
| 18:AU:34:PHE:CD1 | 18:AU:37:LEU:HD12 | 2.56 | 0.41 |
| 19:AV:114:VAL:HG12 | 19:AV:118:ARG:HA | 2.02 | 0.41 |
| 20:AW:5:ILE:HA | 20:AW:21:PHE:CB | 2.50 | 0.41 |
| 20:AW:69:LYS:HB2 | 20:AW:69:LYS:HE2 | 1.92 | 0.41 |
| 20:AW:91:ALA:HB1 | 20:AW:92:PRO:HD2 | 2.03 | 0.41 |
| 21:AX:16:LYS:CE | 21:AX:58:LEU:HD21 | 2.50 | 0.41 |
| 21:AX:33:LYS:HG3 | 21:AX:42:THR:HG23 | 2.03 | 0.41 |
| 22:AY:13:PHE:HB2 | 22:AY:66:THR:HG21 | 2.02 | 0.41 |
| 23:AH:50:VAL:HG13 | 23:AH:111:LYS:O | 2.21 | 0.41 |
| 24:AM:8:LEU:HG | 24:AM:95:ASN:O | 2.20 | 0.41 |
| 26:BA:762:C:OP2 | 38:BM:80:ARG:NH1 | 2.51 | 0.41 |
| 26:BA:1428:G:N2 | 26:BA:1660:A:N3 | 2.67 | 0.41 |
| 27:BB:37:C:H2' | 39:BN:142:ARG:HH12 | 1.86 | 0.41 |
| 27:BB:43:C:O2' | 31:BF:35:GLN:HG2 | 2.20 | 0.41 |
| 27:BB:116:G:H2' | 27:BB:117:C:H6 | 1.84 | 0.41 |
| 30:BE:38:ALA:HA | 30:BE:103:GLU:HB2 | 2.03 | 0.41 |
| 30:BE:197:ILE:HA | 30:BE:239:VAL:O | 2.21 | 0.41 |
| 34:BI:12:VAL:CG2 | 34:BI:57:GLN:HG3 | 2.50 | 0.41 |
| 36:BK:4:MET:SD | 36:BK:116:GLU:HB3 | 2.61 | 0.41 |
| 36:BK:97:ILE:HG22 | 36:BK:105:LYS:HG2 | 2.01 | 0.41 |
| 39:BN:44:VAL:HG12 | 39:BN:64:SER:OG | 2.20 | 0.41 |
| 39:BN:64:SER:CB | 39:BN:76:THR:HB | 2.43 | 0.41 |
| 39:BN:120:LEU:HG | 39:BN:137:PHE:HE1 | 1.86 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 43:BR:75:GLU:OE2 | 43:BR:82:MET:HB3 | 2.21 | 0.41 |
| 49:BX:57:ILE:HA | 49:BX:146:ASN:OD1 | 2.20 | 0.41 |
| 49:BX:69:ARG:HD3 | 49:BX:69:ARG:HA | 1.92 | 0.41 |
| 52:Ba:117:GLU:O | 52:Ba:118:LEU:HD23 | 2.21 | 0.41 |
| 56:Be:18:MET:HG3 | 56:Be:38:THR:HG22 | 2.03 | 0.41 |
| 56:Be:18:MET:HE2 | 56:Be:40:ARG:HB3 | 2.02 | 0.41 |
| 58:AZ:21:TYR:CE1 | 58:AZ:23:LYS:HE3 | 2.56 | 0.41 |
| 58:AZ:46:TYR:CE1 | 58:AZ:55:ALA:HB3 | 2.56 | 0.41 |
| 1:AA:24:U:H2' | 1:AA:25:U:C6 | 2.56 | 0.41 |
| 1:AA:239:G:H2' | 1:AA:240:G:O4' | 2.20 | 0.41 |
| 1:AA:303:U:O4 | 1:AA:310:C:O2' | 2.16 | 0.41 |
| 1:AA:418:G:OP2 | 20:AW:32:ARG:NE | 2.54 | 0.41 |
| 1:AA:680:C:O2' | 1:AA:681:C:H5' | 2.20 | 0.41 |
| 1:AA:775:C:O2' | 1:AA:776:A:H5' | 2.20 | 0.41 |
| 1:AA:991:U:O4 | 1:AA:1147:C:H2' | 2.21 | 0.41 |
| 2:AB:220:VAL:O | 2:AB:224:LEU:HD23 | 2.21 | 0.41 |
| 4:AD:53:GLY:N | 4:AD:61:LYS:O | 2.54 | 0.41 |
| 5:AE:168:LEU:HD11 | 5:AE:173:HIS:ND1 | 2.36 | 0.41 |
| 6:AF:194:VAL:O | 6:AF:200:ASN:ND2 | 2.43 | 0.41 |
| 7:AG:25:MET:CB | 7:AG:45:LEU:HD11 | 2.48 | 0.41 |
| 7:AG:82:TYR:CE2 | 7:AG:110:TYR:HB2 | 2.56 | 0.41 |
| 7:AG:118:TRP:HA | 9:AJ:97:LYS:HB2 | 2.03 | 0.41 |
| 8:AI:7:ILE:N | 8:AI:11:TRP:O | 2.30 | 0.41 |
| 8:AI:105:GLY:HA2 | 8:AI:135:ALA:HB2 | 2.03 | 0.41 |
| 8:AI:152:ARG:NH2 | 8:AI:160:THR:HG21 | 2.37 | 0.41 |
| 11:AL:37:TYR:O | 11:AL:40:GLU:HG2 | 2.21 | 0.41 |
| 14:AP:116:THR:HA | 14:AP:119:GLU:CG | 2.51 | 0.41 |
| 18:AU:19:ARG:NH2 | 18:AU:37:LEU:O | 2.30 | 0.41 |
| 18:AU:26:LEU:HA | 18:AU:29:LEU:HG | 2.03 | 0.41 |
| 18:AU:38:LEU:HD23 | 18:AU:42:GLU:CB | 2.51 | 0.41 |
| 19:AV:4:VAL:HG11 | 19:AV:128:MET:CE | 2.42 | 0.41 |
| 19:AV:42:GLU:C | 19:AV:43:LEU:HD12 | 2.46 | 0.41 |
| 24:AM:24:LYS:HG3 | 24:AM:34:ILE:HD13 | 2.03 | 0.41 |
| 26:BA:899:U:OP2 | 52:Ba:68:ARG:HD3 | 2.21 | 0.41 |
| 27:BB:11:G:O2' | 27:BB:12:C:H5' | 2.21 | 0.41 |
| 27:BB:91:U:O2' | 27:BB:92:G:H5' | 2.21 | 0.41 |
| 29:BD:220:GLN:HG2 | 29:BD:221:LEU:O | 2.21 | 0.41 |
| 29:BD:295:ILE:HD11 | 29:BD:311:LEU:HD11 | 2.03 | 0.41 |
| 30:BE:149:LEU:HD11 | 30:BE:155:VAL:HG22 | 2.03 | 0.41 |
| 37:BL:96:LEU:HG | 37:BL:118:SER:HB2 | 2.03 | 0.41 |
| 39:BN:120:LEU:CD1 | 39:BN:130:ILE:HD13 | 2.51 | 0.41 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 40:BO:108:SER:O | 40:BO:112:ILE:HG22 | 2.20 | 0.41 |
| 41:BP:125:ARG:HA | 41:BP:125:ARG:HD3 | 1.68 | 0.41 |
| 42:BQ:32:LYS:NZ | 42:BQ:33:ASN:OD1 | 2.54 | 0.41 |
| 44:BS:19:MET:CE | 44:BS:144:GLU:HG2 | 2.51 | 0.41 |
| 53:Bb:33:LEU:HD12 | 53:Bb:33:LEU:HA | 1.83 | 0.41 |
| 1:AA:114:U:H2' | 1:AA:115:U:C6 | 2.55 | 0.40 |
| 1:AA:255:U:O4 | 1:AA:256:A:N6 | 2.54 | 0.40 |
| 1:AA:390:C:H1' | 1:AA:405:G:H22 | 1.85 | 0.40 |
| 1:AA:726:G:O2' | 1:AA:727:G:H5' | 2.19 | 0.40 |
| 1:AA:775:C:H1' | 1:AA:798:G:N2 | 2.36 | 0.40 |
| 1:AA:789:C:H2' | 1:AA:790:C:C6 | 2.56 | 0.40 |
| 1:AA:1069:A:C1' | 1:AA:1093:A:H62 | 2.34 | 0.40 |
| 1:AA:1235:A:H5' | 19:AV:90:LYS:CD | 2.49 | 0.40 |
| 1:AA:1280:A:H2' | 1:AA:1281:G:C5' | 2.51 | 0.40 |
| 1:AA:1314:C:H5'' | 11:AL:120:PHE:CB | 2.51 | 0.40 |
| 1:AA:1319:U:H2' | 1:AA:1320:G:O4' | 2.22 | 0.40 |
| 1:AA:1322:A:H4' | 8:AI:60:ASN:HD21 | 1.86 | 0.40 |
| 3:AC:169:ALA:N | 3:AC:176:LEU:O | 2.55 | 0.40 |
| 4:AD:22:GLU:O | 4:AD:80:THR:HG22 | 2.21 | 0.40 |
| 5:AE:11:TYR:HA | 5:AE:46:SER:OG | 2.21 | 0.40 |
| 6:AF:38:PRO:HG2 | 6:AF:41:ILE:CD1 | 2.46 | 0.40 |
| 6:AF:41:ILE:HG23 | 6:AF:45:ASP:OD2 | 2.21 | 0.40 |
| 6:AF:98:GLN:HE21 | 6:AF:102:GLY:HA2 | 1.86 | 0.40 |
| 7:AG:7:TRP:CH2 | 7:AG:9:PRO:HB3 | 2.57 | 0.40 |
| 7:AG:97:ILE:O | 7:AG:101:ILE:HG13 | 2.21 | 0.40 |
| 8:AI:34:VAL:HB | 11:AL:44:MET:HG2 | 2.03 | 0.40 |
| 8:AI:41:HIS:HA | 8:AI:47:ASN:HB2 | 2.02 | 0.40 |
| 14:AP:98:LYS:HZ1 | 18:AU:37:LEU:HA | 1.85 | 0.40 |
| 14:AP:134:ARG:HH11 | 14:AP:139:LEU:HB3 | 1.86 | 0.40 |
| 15:AR:15:CYS:SG | 15:AR:18:CYS:HB2 | 2.61 | 0.40 |
| 19:AV:55:CYS:SG | 19:AV:101:THR:HG22 | 2.61 | 0.40 |
| 19:AV:58:VAL:HG22 | 19:AV:77:TYR:OH | 2.21 | 0.40 |
| 21:AX:24:GLN:CD | 21:AX:37:VAL:HG23 | 2.47 | 0.40 |
| 26:BA:1327:C:C5 | 30:BE:179:ALA:HB2 | 2.56 | 0.40 |
| 26:BA:1754:G:O2' | 53:Bb:19:GLY:HA2 | 2.21 | 0.40 |
| 26:BA:2272:C:P | 43:BR:6:GLY:HA3 | 2.61 | 0.40 |
| 34:BI:54:GLU:HB2 | 34:BI:159:THR:CG2 | 2.48 | 0.40 |
| 34:BI:88:ARG:NE | 34:BI:135:THR:OG1 | 2.45 | 0.40 |
| 39:BN:105:ILE:HG22 | 39:BN:116:VAL:HG11 | 2.03 | 0.40 |
| 43:BR:50:LYS:HE3 | 43:BR:50:LYS:HB3 | 1.90 | 0.40 |
| 44:BS:31:ARG:HB2 | 44:BS:116:ILE:CD1 | 2.51 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 56:Be:6:GLU:OE1 | 56:Be:6:GLU:N | 2.43 | 0.40 |
| 1:AA:519:C:H2' | 1:AA:520:C:H6 | 1.87 | 0.40 |
| 1:AA:684:A:O2' | 16:AQ:8:ARG:HD2 | 2.20 | 0.40 |
| 1:AA:719:A:H2' | 1:AA:720:C:O4' | 2.21 | 0.40 |
| 1:AA:791:G:H2' | 1:AA:792:U:C6 | 2.55 | 0.40 |
| 1:AA:1215:A:H2' | 1:AA:1216:A:C8 | 2.56 | 0.40 |
| 1:AA:1414:U:H2' | 1:AA:1415:U:C6 | 2.56 | 0.40 |
| 3:AC:35:LEU:HG | 24:AM:92:LYS:NZ | 2.37 | 0.40 |
| 4:AD:69:ARG:HD3 | 4:AD:83:MET:HE3 | 2.03 | 0.40 |
| 5:AE:102:SER:HB3 | 7:AG:148:ARG:CD | 2.51 | 0.40 |
| 7:AG:13:LEU:HD21 | 7:AG:33:LEU:HB2 | 2.03 | 0.40 |
| 7:AG:126:HIS:CB | 7:AG:168:LYS:HD2 | 2.51 | 0.40 |
| 7:AG:198:VAL:O | 7:AG:200:LEU:HD12 | 2.21 | 0.40 |
| 14:AP:25:ARG:NH1 | 14:AP:28:ASN:HA | 2.37 | 0.40 |
| 14:AP:99:ASP:HB3 | 14:AP:106:LYS:HZ1 | 1.87 | 0.40 |
| 15:AR:26:ARG:HG2 | 15:AR:31:TYR:CE2 | 2.57 | 0.40 |
| 17:AS:18:ASP:HB3 | 17:AS:21:CYS:CB | 2.37 | 0.40 |
| 18:AU:81:ILE:CD1 | 18:AU:98:LEU:HB3 | 2.51 | 0.40 |
| 19:AV:62:ILE:HD13 | 19:AV:68:ILE:HG13 | 2.03 | 0.40 |
| 19:AV:117:GLY:O | 19:AV:118:ARG:HG3 | 2.21 | 0.40 |
| 22:AY:14:ALA:HB3 | 22:AY:39:GLU:CG | 2.51 | 0.40 |
| 24:AM:51:LYS:HB2 | 24:AM:60:THR:HB | 2.04 | 0.40 |
| 25:AT:46:ILE:O | 25:AT:50:VAL:HG23 | 2.21 | 0.40 |
| 26:BA:897:C:N4 | 37:BL:4:LYS:HG2 | 2.35 | 0.40 |
| 27:BB:44:C:O2 | 31:BF:64:ARG:NE | 2.47 | 0.40 |
| 28:BC:105:PHE:HA | 28:BC:127:ALA:O | 2.22 | 0.40 |
| 28:BC:120:VAL:HG21 | 28:BC:127:ALA:HB2 | 2.03 | 0.40 |
| 31:BF:126:ARG:HG3 | 31:BF:147:VAL:HG13 | 2.01 | 0.40 |
| 32:BG:11:ILE:HD11 | 32:BG:51:VAL:CG2 | 2.49 | 0.40 |
| 33:BH:32:LYS:HD2 | 33:BH:90:LEU:HD23 | 2.03 | 0.40 |
| 36:BK:6:SER:HB3 | 36:BK:81:ARG:O | 2.21 | 0.40 |
| 38:BM:115:VAL:HG13 | 38:BM:132:VAL:CG1 | 2.51 | 0.40 |
| 39:BN:96:LYS:HE2 | 39:BN:96:LYS:HB2 | 1.97 | 0.40 |
| 41:BP:44:LEU:HD23 | 41:BP:47:GLU:OE2 | 2.21 | 0.40 |
| 48:BW:60:ILE:HD13 | 48:BW:60:ILE:HA | 1.88 | 0.40 |
| 49:BX:57:ILE:HD12 | 49:BX:149:LEU:CD1 | 2.51 | 0.40 |
| 51:BZ:4:ASP:OD1 | 51:BZ:4:ASP:N | 2.53 | 0.40 |
| 52:Ba:53:ARG:HB2 | 52:Ba:65:ASN:O | 2.21 | 0.40 |
| 52:Ba:63:ASP:HB2 | 52:Ba:65:ASN:OD1 | 2.21 | 0.40 |
| 1:AA:140:U:O3' | 23:AH:82:GLY:HA2 | 2.20 | 0.40 |
| 1:AA:234:U:H3' | 10:AK:8:ARG:NH2 | 2.36 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|--------------------|--------------------|--------------------------|-------------------|
| 1:AA:244:A:H2' | 1:AA:245:A:O4' | 2.21 | 0.40 |
| 1:AA:493:G:OP1 | 13:AO:108:GLY:HA2 | 2.22 | 0.40 |
| 1:AA:550:U:H3 | 1:AA:570:G:H22 | 1.70 | 0.40 |
| 1:AA:671:G:O2' | 1:AA:707:A:O4' | 2.30 | 0.40 |
| 1:AA:875:A:O2' | 1:AA:876:G:H5' | 2.21 | 0.40 |
| 1:AA:881:A:O2' | 1:AA:882:C:H5' | 2.21 | 0.40 |
| 1:AA:987:G:O4' | 1:AA:1162:U:H4' | 2.21 | 0.40 |
| 1:AA:1208:C:O2 | 1:AA:1222:U:H1' | 2.21 | 0.40 |
| 1:AA:1384:C:H42 | 1:AA:1401:G:H1 | 1.70 | 0.40 |
| 3:AC:40:MET:HE3 | 3:AC:40:MET:HB3 | 1.98 | 0.40 |
| 3:AC:157:PRO:CA | 3:AC:161:VAL:HB | 2.45 | 0.40 |
| 4:AD:27:LEU:CD1 | 4:AD:55:LEU:HD23 | 2.52 | 0.40 |
| 5:AE:51:TYR:CE1 | 5:AE:85:LEU:HD21 | 2.57 | 0.40 |
| 5:AE:136:GLY:HA3 | 5:AE:144:LYS:NZ | 2.36 | 0.40 |
| 6:AF:109:LEU:CD2 | 6:AF:228:ARG:HB2 | 2.52 | 0.40 |
| 10:AK:110:THR:CG2 | 10:AK:122:VAL:HG23 | 2.47 | 0.40 |
| 11:AL:118:LYS:CA | 11:AL:125:ALA:HA | 2.50 | 0.40 |
| 17:AS:76:ALA:O | 17:AS:79:SER:OG | 2.34 | 0.40 |
| 18:AU:76:ILE:HD12 | 18:AU:105:HIS:O | 2.21 | 0.40 |
| 19:AV:83:ASN:CB | 19:AV:86:LYS:HB2 | 2.51 | 0.40 |
| 22:AY:23:ASN:HA | 22:AY:31:SER:HA | 2.03 | 0.40 |
| 22:AY:31:SER:O | 22:AY:50:ASN:HA | 2.21 | 0.40 |
| 26:BA:256:A:N6 | 33:BH:90:LEU:HD11 | 2.36 | 0.40 |
| 26:BA:1575:C:OP1 | 41:BP:118:SER:HA | 2.21 | 0.40 |
| 26:BA:2511:U:O2' | 26:BA:2515:U:OP1 | 2.35 | 0.40 |
| 26:BA:2739:G:O2' | 29:BD:266:LYS:NZ | 2.53 | 0.40 |
| 29:BD:214:VAL:HA | 29:BD:219:ILE:HG12 | 2.04 | 0.40 |
| 30:BE:169:LEU:HD23 | 30:BE:169:LEU:HA | 1.92 | 0.40 |
| 30:BE:169:LEU:HD22 | 30:BE:173:TYR:CE2 | 2.56 | 0.40 |
| 35:BJ:7:ILE:HD13 | 35:BJ:116:ILE:HD11 | 2.03 | 0.40 |
| 36:BK:12:LEU:O | 36:BK:47:ALA:N | 2.41 | 0.40 |
| 39:BN:133:SER:O | 39:BN:136:VAL:HG22 | 2.21 | 0.40 |
| 41:BP:85:LYS:O | 41:BP:85:LYS:HD2 | 2.20 | 0.40 |
| 50:BY:89:VAL:CG1 | 50:BY:92:SER:HB3 | 2.49 | 0.40 |
| 57:Bf:87:LYS:HE2 | 57:Bf:89:GLU:CG | 2.47 | 0.40 |
| 1:AA:208:G:O2' | 1:AA:209:G:H5' | 2.21 | 0.40 |
| 1:AA:504:U:H5' | 13:AO:11:LEU:HD22 | 2.03 | 0.40 |
| 1:AA:669:A:H2' | 1:AA:670:A:C8 | 2.57 | 0.40 |
| 1:AA:981:U:H2' | 1:AA:982:G:H8 | 1.85 | 0.40 |
| 1:AA:1077:C:HO2' | 19:AV:2:THR:HG23 | 1.87 | 0.40 |
| 2:AB:222:TRP:CZ3 | 2:AB:226:ARG:HD3 | 2.57 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 3:AC:149:GLY:O | 3:AC:151:ILE:HD12 | 2.21 | 0.40 |
| 7:AG:49:GLU:HG2 | 7:AG:164:LYS:CD | 2.51 | 0.40 |
| 7:AG:111:ILE:HD12 | 7:AG:192:LEU:HD22 | 2.03 | 0.40 |
| 8:AI:4:LEU:HG | 8:AI:12:ASP:OD2 | 2.20 | 0.40 |
| 8:AI:13:PRO:HA | 8:AI:94:ILE:HD11 | 2.03 | 0.40 |
| 9:AJ:36:ASN:O | 9:AJ:40:VAL:HG12 | 2.21 | 0.40 |
| 9:AJ:105:THR:CG2 | 9:AJ:124:GLN:HB2 | 2.51 | 0.40 |
| 21:AX:13:LEU:CD2 | 21:AX:60:VAL:HG22 | 2.52 | 0.40 |
| 21:AX:17:CYS:HB3 | 21:AX:22:ASN:N | 2.36 | 0.40 |
| 23:AH:94:LYS:HB3 | 23:AH:94:LYS:HE2 | 1.75 | 0.40 |
| 26:BA:1509:G:O2' | 26:BA:1510:U:O5' | 2.39 | 0.40 |
| 26:BA:1972:U:OP2 | 36:BK:66:LYS:HE3 | 2.20 | 0.40 |
| 26:BA:2695:C:C5' | 47:BV:17:THR:HG22 | 2.52 | 0.40 |
| 27:BB:27:A:C2' | 27:BB:28:A:H5' | 2.52 | 0.40 |
| 30:BE:6:THR:O | 30:BE:14:VAL:N | 2.38 | 0.40 |
| 31:BF:108:ARG:HD3 | 31:BF:109:TYR:H | 1.85 | 0.40 |
| 34:BI:18:THR:HG22 | 34:BI:93:GLU:HA | 2.04 | 0.40 |
| 35:BJ:61:ALA:H | 35:BJ:65:GLY:HA3 | 1.86 | 0.40 |
| 37:BL:119:GLU:N | 37:BL:119:GLU:OE1 | 2.52 | 0.40 |
| 40:BO:67:VAL:HG11 | 40:BO:122:ILE:HG12 | 2.04 | 0.40 |
| 40:BO:94:SER:O | 40:BO:98:LYS:HG3 | 2.22 | 0.40 |
| 41:BP:140:SER:HA | 41:BP:143:GLU:HG3 | 2.04 | 0.40 |
| 51:BZ:21:VAL:CG2 | 51:BZ:29:ARG:HG2 | 2.41 | 0.40 |
| 52:Ba:132:ILE:CG2 | 52:Ba:137:LYS:HB2 | 2.51 | 0.40 |
| 1:AA:38:G:OP2 | 1:AA:38:G:H2' | 2.22 | 0.40 |
| 1:AA:858:A:H4' | 1:AA:859:A:H4' | 2.03 | 0.40 |
| 1:AA:1070:G:O6 | 1:AA:1091:G:H1' | 2.22 | 0.40 |
| 1:AA:1194:G:H2' | 1:AA:1195:A:C4 | 2.56 | 0.40 |
| 1:AA:1210:C:OP2 | 19:AV:70:ILE:HG13 | 2.21 | 0.40 |
| 1:AA:1258:G:H2' | 1:AA:1259:G:H8 | 1.85 | 0.40 |
| 1:AA:1276:G:H5'' | 14:AP:48:GLY:N | 2.35 | 0.40 |
| 1:AA:1429:U:C3' | 1:AA:1430:A:H5'' | 2.52 | 0.40 |
| 2:AB:240:LEU:HG | 2:AB:244:GLU:HG2 | 2.02 | 0.40 |
| 3:AC:17:MET:HE1 | 3:AC:73:TYR:CE1 | 2.57 | 0.40 |
| 3:AC:36:ASN:O | 3:AC:43:GLN:HB3 | 2.21 | 0.40 |
| 5:AE:58:LEU:HD13 | 5:AE:78:SER:HA | 2.04 | 0.40 |
| 5:AE:91:ILE:HD12 | 5:AE:91:ILE:HA | 1.97 | 0.40 |
| 7:AG:12:ARG:O | 7:AG:16:LEU:HG | 2.21 | 0.40 |
| 7:AG:107:ASN:HA | 58:AZ:19:LYS:NZ | 2.36 | 0.40 |
| 8:AI:81:ALA:O | 8:AI:85:VAL:HG23 | 2.21 | 0.40 |
| 9:AJ:40:VAL:CG1 | 9:AJ:103:ILE:HD12 | 2.52 | 0.40 |

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| Atom-1 | Atom-2 | Interatomic distance (Å) | Clash overlap (Å) |
|-------------------|--------------------|--------------------------|-------------------|
| 15:AR:20:ARG:C | 15:AR:21:LYS:HD2 | 2.47 | 0.40 |
| 16:AQ:84:ILE:HD11 | 16:AQ:151:THR:CB | 2.48 | 0.40 |
| 26:BA:945:G:N2 | 26:BA:946:U:O4 | 2.30 | 0.40 |
| 26:BA:1103:G:H4' | 42:BQ:37:LYS:HG3 | 2.03 | 0.40 |
| 26:BA:1818:U:O2 | 26:BA:2619:G:O2' | 2.28 | 0.40 |
| 27:BB:76:U:H2' | 27:BB:77:A:C8 | 2.57 | 0.40 |
| 31:BF:52:LYS:HD2 | 31:BF:53:LYS:H | 1.87 | 0.40 |
| 32:BG:17:VAL:HG12 | 32:BG:26:ALA:CA | 2.49 | 0.40 |
| 33:BH:108:MET:O | 33:BH:112:ILE:HG13 | 2.22 | 0.40 |
| 34:BI:29:SER:HB2 | 34:BI:60:HIS:HB2 | 2.04 | 0.40 |
| 36:BK:86:LEU:HD11 | 47:BV:19:LYS:C | 2.46 | 0.40 |
| 38:BM:38:VAL:HG13 | 38:BM:64:VAL:HG11 | 2.04 | 0.40 |
| 41:BP:21:ARG:O | 41:BP:53:LYS:HG3 | 2.20 | 0.40 |
| 46:BU:54:LYS:HD2 | 46:BU:62:GLY:HA2 | 2.03 | 0.40 |
| 58:AZ:33:ILE:HD11 | 58:AZ:48:CYS:CB | 2.51 | 0.40 |

There are no symmetry-related clashes.

5.3 Torsion angles ⓘ

5.3.1 Protein backbone ⓘ

In the following table, the Percentiles column shows the percent Ramachandran outliers of the chain as a percentile score with respect to all 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 | AB | 188/247 (76%) | 183 (97%) | 5 (3%) | 0 | 100 | 100 |
| 3 | AC | 184/318 (58%) | 167 (91%) | 17 (9%) | 0 | 100 | 100 |
| 4 | AD | 180/203 (89%) | 174 (97%) | 6 (3%) | 0 | 100 | 100 |
| 5 | AE | 163/218 (75%) | 154 (94%) | 9 (6%) | 0 | 100 | 100 |
| 6 | AF | 232/235 (99%) | 205 (88%) | 27 (12%) | 0 | 100 | 100 |
| 7 | AG | 196/209 (94%) | 184 (94%) | 12 (6%) | 0 | 100 | 100 |
| 8 | AI | 187/189 (99%) | 168 (90%) | 19 (10%) | 0 | 100 | 100 |
| 9 | AJ | 127/130 (98%) | 118 (93%) | 9 (7%) | 0 | 100 | 100 |

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| Mol | Chain | Analysed | Favoured | Allowed | Outliers | Percentiles | |
|-----|-------|---------------|-----------|----------|----------|-------------|-----|
| 10 | AK | 123/125 (98%) | 109 (89%) | 14 (11%) | 0 | 100 | 100 |
| 11 | AL | 130/134 (97%) | 110 (85%) | 18 (14%) | 2 (2%) | 8 | 28 |
| 12 | AN | 121/126 (96%) | 111 (92%) | 9 (7%) | 1 (1%) | 16 | 44 |
| 13 | AO | 140/142 (99%) | 130 (93%) | 10 (7%) | 0 | 100 | 100 |
| 14 | AP | 130/182 (71%) | 120 (92%) | 9 (7%) | 1 (1%) | 16 | 44 |
| 15 | AR | 42/50 (84%) | 38 (90%) | 4 (10%) | 0 | 100 | 100 |
| 16 | AQ | 150/152 (99%) | 143 (95%) | 7 (5%) | 0 | 100 | 100 |
| 17 | AS | 107/109 (98%) | 93 (87%) | 14 (13%) | 0 | 100 | 100 |
| 18 | AU | 93/136 (68%) | 86 (92%) | 7 (8%) | 0 | 100 | 100 |
| 19 | AV | 147/149 (99%) | 132 (90%) | 14 (10%) | 1 (1%) | 19 | 47 |
| 20 | AW | 93/101 (92%) | 84 (90%) | 9 (10%) | 0 | 100 | 100 |
| 21 | AX | 58/62 (94%) | 52 (90%) | 6 (10%) | 0 | 100 | 100 |
| 22 | AY | 61/76 (80%) | 52 (85%) | 9 (15%) | 0 | 100 | 100 |
| 23 | AH | 122/136 (90%) | 114 (93%) | 8 (7%) | 0 | 100 | 100 |
| 24 | AM | 100/102 (98%) | 96 (96%) | 4 (4%) | 0 | 100 | 100 |
| 25 | AT | 56/64 (88%) | 54 (96%) | 2 (4%) | 0 | 100 | 100 |
| 28 | BC | 236/238 (99%) | 228 (97%) | 8 (3%) | 0 | 100 | 100 |
| 29 | BD | 335/337 (99%) | 318 (95%) | 17 (5%) | 0 | 100 | 100 |
| 30 | BE | 250/253 (99%) | 234 (94%) | 16 (6%) | 0 | 100 | 100 |
| 31 | BF | 163/165 (99%) | 157 (96%) | 6 (4%) | 0 | 100 | 100 |
| 32 | BG | 174/176 (99%) | 170 (98%) | 4 (2%) | 0 | 100 | 100 |
| 33 | BH | 113/120 (94%) | 111 (98%) | 2 (2%) | 0 | 100 | 100 |
| 34 | BI | 155/173 (90%) | 148 (96%) | 7 (4%) | 0 | 100 | 100 |
| 35 | BJ | 136/143 (95%) | 127 (93%) | 9 (7%) | 0 | 100 | 100 |
| 36 | BK | 130/132 (98%) | 127 (98%) | 3 (2%) | 0 | 100 | 100 |
| 37 | BL | 138/140 (99%) | 130 (94%) | 8 (6%) | 0 | 100 | 100 |
| 38 | BM | 194/196 (99%) | 187 (96%) | 7 (4%) | 0 | 100 | 100 |
| 39 | BN | 172/174 (99%) | 168 (98%) | 4 (2%) | 0 | 100 | 100 |
| 40 | BO | 124/126 (98%) | 119 (96%) | 5 (4%) | 0 | 100 | 100 |
| 41 | BP | 149/151 (99%) | 146 (98%) | 3 (2%) | 0 | 100 | 100 |
| 42 | BQ | 55/61 (90%) | 53 (96%) | 2 (4%) | 0 | 100 | 100 |

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| Mol | Chain | Analysed | Favoured | Allowed | Outliers | Percentiles | |
|-----|-------|-----------------|------------|----------|----------|-------------|-----|
| 43 | BR | 94/97 (97%) | 90 (96%) | 4 (4%) | 0 | 100 | 100 |
| 44 | BS | 149/151 (99%) | 140 (94%) | 9 (6%) | 0 | 100 | 100 |
| 45 | BT | 80/82 (98%) | 78 (98%) | 2 (2%) | 0 | 100 | 100 |
| 46 | BU | 117/119 (98%) | 110 (94%) | 7 (6%) | 0 | 100 | 100 |
| 47 | BV | 60/62 (97%) | 59 (98%) | 1 (2%) | 0 | 100 | 100 |
| 48 | BW | 65/67 (97%) | 63 (97%) | 2 (3%) | 0 | 100 | 100 |
| 49 | BX | 151/153 (99%) | 146 (97%) | 5 (3%) | 0 | 100 | 100 |
| 50 | BY | 90/99 (91%) | 86 (96%) | 4 (4%) | 0 | 100 | 100 |
| 51 | BZ | 84/89 (94%) | 83 (99%) | 1 (1%) | 0 | 100 | 100 |
| 52 | Ba | 130/161 (81%) | 126 (97%) | 4 (3%) | 0 | 100 | 100 |
| 53 | Bb | 92/94 (98%) | 86 (94%) | 6 (6%) | 0 | 100 | 100 |
| 54 | Bc | 54/56 (96%) | 52 (96%) | 2 (4%) | 0 | 100 | 100 |
| 55 | Bd | 49/51 (96%) | 46 (94%) | 3 (6%) | 0 | 100 | 100 |
| 56 | Be | 42/52 (81%) | 38 (90%) | 4 (10%) | 0 | 100 | 100 |
| 57 | Bf | 90/92 (98%) | 86 (96%) | 4 (4%) | 0 | 100 | 100 |
| 58 | AZ | 48/56 (86%) | 46 (96%) | 2 (4%) | 0 | 100 | 100 |
| All | All | 7049/7661 (92%) | 6635 (94%) | 409 (6%) | 5 (0%) | 50 | 76 |

All (5) Ramachandran outliers are listed below:

| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 11 | AL | 105 | ARG |
| 19 | AV | 30 | GLU |
| 11 | AL | 12 | LYS |
| 14 | AP | 49 | ARG |
| 12 | AN | 124 | ARG |

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 | |
|-----|-------|----------------|------------|----------|-------------|-----|
| 2 | AB | 166/212 (78%) | 166 (100%) | 0 | 100 | 100 |
| 3 | AC | 149/250 (60%) | 149 (100%) | 0 | 100 | 100 |
| 4 | AD | 160/173 (92%) | 160 (100%) | 0 | 100 | 100 |
| 5 | AE | 145/178 (82%) | 145 (100%) | 0 | 100 | 100 |
| 6 | AF | 208/209 (100%) | 208 (100%) | 0 | 100 | 100 |
| 7 | AG | 161/170 (95%) | 161 (100%) | 0 | 100 | 100 |
| 8 | AI | 159/159 (100%) | 159 (100%) | 0 | 100 | 100 |
| 9 | AJ | 104/105 (99%) | 104 (100%) | 0 | 100 | 100 |
| 10 | AK | 105/105 (100%) | 105 (100%) | 0 | 100 | 100 |
| 11 | AL | 105/107 (98%) | 105 (100%) | 0 | 100 | 100 |
| 12 | AN | 92/94 (98%) | 92 (100%) | 0 | 100 | 100 |
| 13 | AO | 112/112 (100%) | 112 (100%) | 0 | 100 | 100 |
| 14 | AP | 108/151 (72%) | 108 (100%) | 0 | 100 | 100 |
| 15 | AR | 38/44 (86%) | 38 (100%) | 0 | 100 | 100 |
| 16 | AQ | 138/138 (100%) | 138 (100%) | 0 | 100 | 100 |
| 17 | AS | 96/96 (100%) | 96 (100%) | 0 | 100 | 100 |
| 18 | AU | 84/118 (71%) | 84 (100%) | 0 | 100 | 100 |
| 19 | AV | 123/123 (100%) | 123 (100%) | 0 | 100 | 100 |
| 20 | AW | 83/87 (95%) | 83 (100%) | 0 | 100 | 100 |
| 21 | AX | 54/56 (96%) | 54 (100%) | 0 | 100 | 100 |
| 22 | AY | 50/62 (81%) | 50 (100%) | 0 | 100 | 100 |
| 23 | AH | 96/103 (93%) | 96 (100%) | 0 | 100 | 100 |
| 24 | AM | 89/89 (100%) | 89 (100%) | 0 | 100 | 100 |
| 25 | AT | 52/57 (91%) | 52 (100%) | 0 | 100 | 100 |
| 28 | BC | 188/188 (100%) | 188 (100%) | 0 | 100 | 100 |
| 29 | BD | 277/277 (100%) | 277 (100%) | 0 | 100 | 100 |
| 30 | BE | 197/198 (100%) | 197 (100%) | 0 | 100 | 100 |
| 31 | BF | 140/140 (100%) | 140 (100%) | 0 | 100 | 100 |
| 32 | BG | 148/148 (100%) | 148 (100%) | 0 | 100 | 100 |
| 33 | BH | 88/91 (97%) | 88 (100%) | 0 | 100 | 100 |
| 34 | BI | 132/141 (94%) | 132 (100%) | 0 | 100 | 100 |
| 35 | BJ | 114/119 (96%) | 114 (100%) | 0 | 100 | 100 |

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| Mol | Chain | Analysed | Rotameric | Outliers | Percentiles | |
|-----|-------|-----------------|-------------|----------|-------------|-----|
| 36 | BK | 109/109 (100%) | 109 (100%) | 0 | 100 | 100 |
| 37 | BL | 108/108 (100%) | 108 (100%) | 0 | 100 | 100 |
| 38 | BM | 166/166 (100%) | 166 (100%) | 0 | 100 | 100 |
| 39 | BN | 143/143 (100%) | 143 (100%) | 0 | 100 | 100 |
| 40 | BO | 104/104 (100%) | 104 (100%) | 0 | 100 | 100 |
| 41 | BP | 123/123 (100%) | 123 (100%) | 0 | 100 | 100 |
| 42 | BQ | 50/54 (93%) | 50 (100%) | 0 | 100 | 100 |
| 43 | BR | 85/86 (99%) | 85 (100%) | 0 | 100 | 100 |
| 44 | BS | 124/124 (100%) | 124 (100%) | 0 | 100 | 100 |
| 45 | BT | 73/73 (100%) | 73 (100%) | 0 | 100 | 100 |
| 46 | BU | 100/100 (100%) | 100 (100%) | 0 | 100 | 100 |
| 47 | BV | 54/54 (100%) | 54 (100%) | 0 | 100 | 100 |
| 48 | BW | 57/57 (100%) | 57 (100%) | 0 | 100 | 100 |
| 49 | BX | 134/134 (100%) | 134 (100%) | 0 | 100 | 100 |
| 50 | BY | 71/78 (91%) | 71 (100%) | 0 | 100 | 100 |
| 51 | BZ | 76/78 (97%) | 76 (100%) | 0 | 100 | 100 |
| 52 | Ba | 109/135 (81%) | 109 (100%) | 0 | 100 | 100 |
| 53 | Bb | 76/76 (100%) | 76 (100%) | 0 | 100 | 100 |
| 54 | Bc | 49/49 (100%) | 49 (100%) | 0 | 100 | 100 |
| 55 | Bd | 48/48 (100%) | 48 (100%) | 0 | 100 | 100 |
| 56 | Be | 37/43 (86%) | 37 (100%) | 0 | 100 | 100 |
| 57 | Bf | 82/82 (100%) | 82 (100%) | 0 | 100 | 100 |
| 58 | AZ | 42/48 (88%) | 42 (100%) | 0 | 100 | 100 |
| All | All | 5981/6372 (94%) | 5981 (100%) | 0 | 100 | 100 |

There are no protein residues with a non-rotameric sidechain to report.

Sometimes sidechains can be flipped to improve hydrogen bonding and reduce clashes. All (27) such sidechains are listed below:

| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 5 | AE | 107 | ASN |
| 6 | AF | 20 | ASN |
| 6 | AF | 121 | ASN |
| 8 | AI | 95 | GLN |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 12 | AN | 54 | GLN |
| 13 | AO | 78 | GLN |
| 19 | AV | 83 | ASN |
| 20 | AW | 89 | ASN |
| 28 | BC | 9 | ASN |
| 28 | BC | 196 | ASN |
| 29 | BD | 220 | GLN |
| 29 | BD | 229 | GLN |
| 29 | BD | 286 | ASN |
| 32 | BG | 154 | GLN |
| 34 | BI | 33 | HIS |
| 36 | BK | 78 | GLN |
| 38 | BM | 127 | HIS |
| 38 | BM | 138 | HIS |
| 39 | BN | 108 | GLN |
| 46 | BU | 68 | GLN |
| 46 | BU | 97 | ASN |
| 47 | BV | 3 | GLN |
| 49 | BX | 30 | ASN |
| 49 | BX | 134 | GLN |
| 52 | Ba | 44 | GLN |
| 55 | Bd | 16 | HIS |
| 57 | Bf | 18 | ASN |

5.3.3 RNA ⓘ

| Mol | Chain | Analysed | Backbone Outliers | Pucker Outliers |
|-----|-------|-----------------|-------------------|-----------------|
| 1 | AA | 1459/1478 (98%) | 304 (20%) | 4 (0%) |
| 26 | BA | 2769/2899 (95%) | 424 (15%) | 6 (0%) |
| 27 | BB | 127/129 (98%) | 13 (10%) | 0 |
| All | All | 4355/4506 (96%) | 741 (17%) | 10 (0%) |

All (741) RNA backbone outliers are listed below:

| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 1 | AA | 2 | U |
| 1 | AA | 4 | C |
| 1 | AA | 8 | U |
| 1 | AA | 17 | C |
| 1 | AA | 33 | U |
| 1 | AA | 38 | G |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 1 | AA | 39 | U |
| 1 | AA | 42 | G |
| 1 | AA | 45 | U |
| 1 | AA | 47 | A |
| 1 | AA | 70 | U |
| 1 | AA | 72 | U |
| 1 | AA | 73 | G |
| 1 | AA | 86 | U |
| 1 | AA | 91 | A |
| 1 | AA | 95 | A |
| 1 | AA | 96 | C |
| 1 | AA | 105 | A |
| 1 | AA | 107 | C |
| 1 | AA | 117 | G |
| 1 | AA | 120 | C |
| 1 | AA | 122 | G |
| 1 | AA | 125 | A |
| 1 | AA | 133 | G |
| 1 | AA | 139 | C |
| 1 | AA | 152 | C |
| 1 | AA | 156 | U |
| 1 | AA | 157 | A |
| 1 | AA | 158 | A |
| 1 | AA | 161 | C |
| 1 | AA | 162 | A |
| 1 | AA | 163 | C |
| 1 | AA | 164 | A |
| 1 | AA | 166 | A |
| 1 | AA | 167 | U |
| 1 | AA | 171 | G |
| 1 | AA | 172 | G |
| 1 | AA | 173 | A |
| 1 | AA | 174 | A |
| 1 | AA | 175 | U |
| 1 | AA | 176 | G |
| 1 | AA | 178 | U |
| 1 | AA | 180 | U |
| 1 | AA | 187 | A |
| 1 | AA | 190 | A |
| 1 | AA | 191 | C |
| 1 | AA | 192 | G |
| 1 | AA | 196 | U |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 1 | AA | 197 | C |
| 1 | AA | 199 | U |
| 1 | AA | 218 | G |
| 1 | AA | 222 | C |
| 1 | AA | 227 | C |
| 1 | AA | 229 | G |
| 1 | AA | 233 | G |
| 1 | AA | 240 | G |
| 1 | AA | 248 | U |
| 1 | AA | 249 | A |
| 1 | AA | 261 | A |
| 1 | AA | 262 | C |
| 1 | AA | 263 | A |
| 1 | AA | 271 | C |
| 1 | AA | 287 | G |
| 1 | AA | 291 | C |
| 1 | AA | 297 | A |
| 1 | AA | 310 | C |
| 1 | AA | 311 | A |
| 1 | AA | 312 | U |
| 1 | AA | 327 | C |
| 1 | AA | 328 | G |
| 1 | AA | 329 | G |
| 1 | AA | 334 | C |
| 1 | AA | 336 | G |
| 1 | AA | 338 | A |
| 1 | AA | 347 | A |
| 1 | AA | 349 | C |
| 1 | AA | 351 | U |
| 1 | AA | 354 | C |
| 1 | AA | 355 | A |
| 1 | AA | 366 | C |
| 1 | AA | 372 | U |
| 1 | AA | 375 | G |
| 1 | AA | 379 | A |
| 1 | AA | 380 | C |
| 1 | AA | 385 | A |
| 1 | AA | 388 | G |
| 1 | AA | 391 | A |
| 1 | AA | 394 | A |
| 1 | AA | 397 | A |
| 1 | AA | 399 | A |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 1 | AA | 401 | G |
| 1 | AA | 402 | C |
| 1 | AA | 405 | G |
| 1 | AA | 422 | A |
| 1 | AA | 433 | U |
| 1 | AA | 434 | A |
| 1 | AA | 435 | G |
| 1 | AA | 436 | C |
| 1 | AA | 448 | A |
| 1 | AA | 449 | A |
| 1 | AA | 451 | A |
| 1 | AA | 454 | G |
| 1 | AA | 458 | C |
| 1 | AA | 467 | G |
| 1 | AA | 470 | G |
| 1 | AA | 471 | U |
| 1 | AA | 476 | C |
| 1 | AA | 487 | A |
| 1 | AA | 493 | G |
| 1 | AA | 500 | A |
| 1 | AA | 501 | U |
| 1 | AA | 507 | G |
| 1 | AA | 508 | G |
| 1 | AA | 513 | A |
| 1 | AA | 516 | G |
| 1 | AA | 517 | G |
| 1 | AA | 522 | U |
| 1 | AA | 536 | A |
| 1 | AA | 545 | G |
| 1 | AA | 559 | U |
| 1 | AA | 573 | U |
| 1 | AA | 574 | U |
| 1 | AA | 575 | C |
| 1 | AA | 583 | A |
| 1 | AA | 590 | G |
| 1 | AA | 594 | U |
| 1 | AA | 606 | A |
| 1 | AA | 629 | G |
| 1 | AA | 664 | U |
| 1 | AA | 665 | G |
| 1 | AA | 672 | G |
| 1 | AA | 689 | G |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 1 | AA | 690 | U |
| 1 | AA | 694 | A |
| 1 | AA | 696 | G |
| 1 | AA | 701 | G |
| 1 | AA | 702 | G |
| 1 | AA | 707 | A |
| 1 | AA | 734 | U |
| 1 | AA | 735 | A |
| 1 | AA | 750 | G |
| 1 | AA | 756 | A |
| 1 | AA | 758 | C |
| 1 | AA | 761 | U |
| 1 | AA | 762 | G |
| 1 | AA | 769 | A |
| 1 | AA | 773 | G |
| 1 | AA | 774 | U |
| 1 | AA | 785 | G |
| 1 | AA | 786 | C |
| 1 | AA | 787 | G |
| 1 | AA | 788 | A |
| 1 | AA | 789 | C |
| 1 | AA | 791 | G |
| 1 | AA | 808 | G |
| 1 | AA | 809 | A |
| 1 | AA | 817 | A |
| 1 | AA | 818 | A |
| 1 | AA | 830 | G |
| 1 | AA | 834 | A |
| 1 | AA | 859 | A |
| 1 | AA | 871 | G |
| 1 | AA | 872 | G |
| 1 | AA | 879 | C |
| 1 | AA | 881 | A |
| 1 | AA | 906 | U |
| 1 | AA | 907 | U |
| 1 | AA | 914 | A |
| 1 | AA | 915 | A |
| 1 | AA | 917 | G |
| 1 | AA | 920 | G |
| 1 | AA | 923 | C |
| 1 | AA | 925 | A |
| 1 | AA | 927 | U |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | AA | 928 | C |
| 1 | AA | 929 | A |
| 1 | AA | 934 | G |
| 1 | AA | 944 | A |
| 1 | AA | 951 | A |
| 1 | AA | 955 | C |
| 1 | AA | 959 | C |
| 1 | AA | 964 | G |
| 1 | AA | 968 | U |
| 1 | AA | 984 | G |
| 1 | AA | 992 | G |
| 1 | AA | 995 | U |
| 1 | AA | 1003 | G |
| 1 | AA | 1004 | C |
| 1 | AA | 1009 | U |
| 1 | AA | 1020 | A |
| 1 | AA | 1033 | G |
| 1 | AA | 1034 | U |
| 1 | AA | 1039 | C |
| 1 | AA | 1040 | A |
| 1 | AA | 1053 | C |
| 1 | AA | 1056 | G |
| 1 | AA | 1057 | C |
| 1 | AA | 1059 | C |
| 1 | AA | 1060 | A |
| 1 | AA | 1063 | G |
| 1 | AA | 1064 | U |
| 1 | AA | 1065 | U |
| 1 | AA | 1069 | A |
| 1 | AA | 1074 | G |
| 1 | AA | 1075 | U |
| 1 | AA | 1076 | C |
| 1 | AA | 1079 | C |
| 1 | AA | 1082 | G |
| 1 | AA | 1087 | A |
| 1 | AA | 1088 | U |
| 1 | AA | 1093 | A |
| 1 | AA | 1094 | C |
| 1 | AA | 1098 | G |
| 1 | AA | 1099 | U |
| 1 | AA | 1100 | G |
| 1 | AA | 1104 | A |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | AA | 1106 | C |
| 1 | AA | 1107 | G |
| 1 | AA | 1114 | U |
| 1 | AA | 1115 | U |
| 1 | AA | 1120 | C |
| 1 | AA | 1121 | G |
| 1 | AA | 1123 | A |
| 1 | AA | 1124 | G |
| 1 | AA | 1125 | G |
| 1 | AA | 1126 | A |
| 1 | AA | 1128 | G |
| 1 | AA | 1130 | U |
| 1 | AA | 1131 | G |
| 1 | AA | 1143 | A |
| 1 | AA | 1144 | G |
| 1 | AA | 1147 | C |
| 1 | AA | 1148 | A |
| 1 | AA | 1159 | A |
| 1 | AA | 1172 | A |
| 1 | AA | 1173 | C |
| 1 | AA | 1175 | C |
| 1 | AA | 1185 | A |
| 1 | AA | 1186 | A |
| 1 | AA | 1187 | U |
| 1 | AA | 1188 | G |
| 1 | AA | 1189 | A |
| 1 | AA | 1190 | A |
| 1 | AA | 1195 | A |
| 1 | AA | 1196 | C |
| 1 | AA | 1200 | G |
| 1 | AA | 1204 | C |
| 1 | AA | 1214 | A |
| 1 | AA | 1217 | A |
| 1 | AA | 1225 | U |
| 1 | AA | 1226 | A |
| 1 | AA | 1227 | A |
| 1 | AA | 1228 | U |
| 1 | AA | 1232 | A |
| 1 | AA | 1234 | A |
| 1 | AA | 1237 | C |
| 1 | AA | 1241 | U |
| 1 | AA | 1245 | U |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | AA | 1246 | A |
| 1 | AA | 1247 | G |
| 1 | AA | 1248 | U |
| 1 | AA | 1249 | U |
| 1 | AA | 1250 | C |
| 1 | AA | 1251 | G |
| 1 | AA | 1252 | G |
| 1 | AA | 1267 | C |
| 1 | AA | 1281 | G |
| 1 | AA | 1283 | U |
| 1 | AA | 1285 | G |
| 1 | AA | 1288 | U |
| 1 | AA | 1300 | G |
| 1 | AA | 1306 | C |
| 1 | AA | 1307 | A |
| 1 | AA | 1311 | U |
| 1 | AA | 1322 | A |
| 1 | AA | 1324 | A |
| 1 | AA | 1325 | C |
| 1 | AA | 1326 | G |
| 1 | AA | 1329 | C |
| 1 | AA | 1344 | C |
| 1 | AA | 1345 | A |
| 1 | AA | 1348 | G |
| 1 | AA | 1355 | A |
| 1 | AA | 1365 | A |
| 1 | AA | 1375 | G |
| 1 | AA | 1389 | A |
| 1 | AA | 1390 | C |
| 1 | AA | 1393 | C |
| 1 | AA | 1394 | G |
| 1 | AA | 1395 | U |
| 1 | AA | 1396 | G |
| 1 | AA | 1413 | A |
| 1 | AA | 1429 | U |
| 1 | AA | 1430 | A |
| 1 | AA | 1431 | A |
| 1 | AA | 1435 | G |
| 1 | AA | 1442 | G |
| 1 | AA | 1444 | U |
| 1 | AA | 1455 | G |
| 1 | AA | 1457 | A |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | AA | 1458 | U |
| 1 | AA | 1467 | G |
| 1 | AA | 1468 | G |
| 1 | AA | 1469 | A |
| 26 | BA | 7 | U |
| 26 | BA | 10 | G |
| 26 | BA | 28 | C |
| 26 | BA | 39 | C |
| 26 | BA | 56 | A |
| 26 | BA | 64 | A |
| 26 | BA | 67 | A |
| 26 | BA | 68 | G |
| 26 | BA | 81 | G |
| 26 | BA | 85 | G |
| 26 | BA | 106 | U |
| 26 | BA | 110 | A |
| 26 | BA | 112 | U |
| 26 | BA | 116 | A |
| 26 | BA | 117 | C |
| 26 | BA | 119 | U |
| 26 | BA | 126 | A |
| 26 | BA | 127 | G |
| 26 | BA | 128 | U |
| 26 | BA | 136 | C |
| 26 | BA | 146 | A |
| 26 | BA | 161 | A |
| 26 | BA | 181 | A |
| 26 | BA | 186 | A |
| 26 | BA | 187 | A |
| 26 | BA | 193 | U |
| 26 | BA | 194 | U |
| 26 | BA | 198 | A |
| 26 | BA | 213 | G |
| 26 | BA | 231 | G |
| 26 | BA | 234 | C |
| 26 | BA | 243 | U |
| 26 | BA | 248 | U |
| 26 | BA | 250 | G |
| 26 | BA | 266 | G |
| 26 | BA | 267 | U |
| 26 | BA | 268 | A |
| 26 | BA | 278 | A |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 26 | BA | 281 | C |
| 26 | BA | 287 | G |
| 26 | BA | 293 | A |
| 26 | BA | 302 | U |
| 26 | BA | 303 | U |
| 26 | BA | 311 | A |
| 26 | BA | 322 | U |
| 26 | BA | 323 | A |
| 26 | BA | 326 | G |
| 26 | BA | 330 | G |
| 26 | BA | 331 | A |
| 26 | BA | 332 | C |
| 26 | BA | 339 | U |
| 26 | BA | 341 | A |
| 26 | BA | 344 | U |
| 26 | BA | 367 | G |
| 26 | BA | 368 | U |
| 26 | BA | 375 | G |
| 26 | BA | 390 | U |
| 26 | BA | 391 | A |
| 26 | BA | 396 | G |
| 26 | BA | 412 | C |
| 26 | BA | 421 | C |
| 26 | BA | 429 | A |
| 26 | BA | 443 | A |
| 26 | BA | 444 | C |
| 26 | BA | 448 | U |
| 26 | BA | 451 | C |
| 26 | BA | 454 | A |
| 26 | BA | 455 | A |
| 26 | BA | 456 | U |
| 26 | BA | 457 | A |
| 26 | BA | 458 | G |
| 26 | BA | 467 | C |
| 26 | BA | 481 | G |
| 26 | BA | 504 | A |
| 26 | BA | 506 | U |
| 26 | BA | 507 | G |
| 26 | BA | 530 | A |
| 26 | BA | 531 | C |
| 26 | BA | 532 | G |
| 26 | BA | 543 | A |

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| Mol | Chain | Res | Type |
|-----|-------|-----|------|
| 26 | BA | 546 | G |
| 26 | BA | 549 | U |
| 26 | BA | 582 | A |
| 26 | BA | 598 | C |
| 26 | BA | 613 | A |
| 26 | BA | 618 | U |
| 26 | BA | 623 | G |
| 26 | BA | 624 | A |
| 26 | BA | 625 | A |
| 26 | BA | 649 | G |
| 26 | BA | 653 | A |
| 26 | BA | 654 | U |
| 26 | BA | 658 | U |
| 26 | BA | 665 | U |
| 26 | BA | 666 | U |
| 26 | BA | 667 | A |
| 26 | BA | 674 | C |
| 26 | BA | 693 | A |
| 26 | BA | 694 | U |
| 26 | BA | 695 | G |
| 26 | BA | 706 | U |
| 26 | BA | 708 | U |
| 26 | BA | 726 | A |
| 26 | BA | 727 | C |
| 26 | BA | 730 | G |
| 26 | BA | 736 | G |
| 26 | BA | 745 | G |
| 26 | BA | 752 | G |
| 26 | BA | 758 | A |
| 26 | BA | 760 | G |
| 26 | BA | 769 | U |
| 26 | BA | 801 | G |
| 26 | BA | 809 | G |
| 26 | BA | 813 | A |
| 26 | BA | 817 | U |
| 26 | BA | 818 | A |
| 26 | BA | 819 | A |
| 26 | BA | 827 | U |
| 26 | BA | 831 | C |
| 26 | BA | 832 | U |
| 26 | BA | 833 | G |
| 26 | BA | 837 | A |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 26 | BA | 838 | U |
| 26 | BA | 849 | A |
| 26 | BA | 850 | C |
| 26 | BA | 867 | A |
| 26 | BA | 869 | U |
| 26 | BA | 870 | G |
| 26 | BA | 874 | A |
| 26 | BA | 890 | G |
| 26 | BA | 897 | C |
| 26 | BA | 912 | C |
| 26 | BA | 913 | G |
| 26 | BA | 920 | G |
| 26 | BA | 930 | G |
| 26 | BA | 935 | C |
| 26 | BA | 944 | A |
| 26 | BA | 945 | G |
| 26 | BA | 952 | A |
| 26 | BA | 955 | G |
| 26 | BA | 959 | G |
| 26 | BA | 964 | U |
| 26 | BA | 965 | U |
| 26 | BA | 966 | C |
| 26 | BA | 970 | G |
| 26 | BA | 971 | G |
| 26 | BA | 979 | C |
| 26 | BA | 981 | C |
| 26 | BA | 983 | C |
| 26 | BA | 984 | G |
| 26 | BA | 989 | C |
| 26 | BA | 991 | U |
| 26 | BA | 994 | C |
| 26 | BA | 996 | A |
| 26 | BA | 997 | A |
| 26 | BA | 1000 | C |
| 26 | BA | 1001 | C |
| 26 | BA | 1030 | G |
| 26 | BA | 1034 | G |
| 26 | BA | 1041 | U |
| 26 | BA | 1042 | G |
| 26 | BA | 1049 | G |
| 26 | BA | 1062 | A |
| 26 | BA | 1068 | G |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 26 | BA | 1077 | G |
| 26 | BA | 1078 | C |
| 26 | BA | 1096 | A |
| 26 | BA | 1098 | G |
| 26 | BA | 1099 | U |
| 26 | BA | 1100 | G |
| 26 | BA | 1109 | G |
| 26 | BA | 1113 | U |
| 26 | BA | 1116 | U |
| 26 | BA | 1117 | A |
| 26 | BA | 1118 | C |
| 26 | BA | 1119 | U |
| 26 | BA | 1120 | A |
| 26 | BA | 1139 | A |
| 26 | BA | 1140 | G |
| 26 | BA | 1141 | A |
| 26 | BA | 1144 | G |
| 26 | BA | 1149 | A |
| 26 | BA | 1150 | A |
| 26 | BA | 1151 | G |
| 26 | BA | 1152 | G |
| 26 | BA | 1153 | U |
| 26 | BA | 1154 | U |
| 26 | BA | 1164 | G |
| 26 | BA | 1165 | C |
| 26 | BA | 1174 | U |
| 26 | BA | 1175 | U |
| 26 | BA | 1177 | A |
| 26 | BA | 1179 | A |
| 26 | BA | 1180 | G |
| 26 | BA | 1181 | A |
| 26 | BA | 1193 | C |
| 26 | BA | 1198 | C |
| 26 | BA | 1203 | G |
| 26 | BA | 1205 | G |
| 26 | BA | 1223 | U |
| 26 | BA | 1225 | G |
| 26 | BA | 1227 | C |
| 26 | BA | 1231 | G |
| 26 | BA | 1235 | A |
| 26 | BA | 1245 | C |
| 26 | BA | 1267 | G |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 26 | BA | 1273 | C |
| 26 | BA | 1276 | G |
| 26 | BA | 1295 | G |
| 26 | BA | 1312 | U |
| 26 | BA | 1315 | A |
| 26 | BA | 1324 | G |
| 26 | BA | 1326 | C |
| 26 | BA | 1328 | G |
| 26 | BA | 1337 | C |
| 26 | BA | 1344 | C |
| 26 | BA | 1359 | A |
| 26 | BA | 1360 | G |
| 26 | BA | 1361 | C |
| 26 | BA | 1362 | A |
| 26 | BA | 1365 | G |
| 26 | BA | 1390 | A |
| 26 | BA | 1391 | G |
| 26 | BA | 1418 | U |
| 26 | BA | 1435 | G |
| 26 | BA | 1437 | U |
| 26 | BA | 1457 | C |
| 26 | BA | 1466 | C |
| 26 | BA | 1468 | A |
| 26 | BA | 1470 | A |
| 26 | BA | 1486 | A |
| 26 | BA | 1487 | U |
| 26 | BA | 1489 | C |
| 26 | BA | 1494 | A |
| 26 | BA | 1506 | U |
| 26 | BA | 1507 | U |
| 26 | BA | 1509 | G |
| 26 | BA | 1515 | C |
| 26 | BA | 1516 | G |
| 26 | BA | 1545 | C |
| 26 | BA | 1551 | C |
| 26 | BA | 1561 | A |
| 26 | BA | 1570 | G |
| 26 | BA | 1571 | A |
| 26 | BA | 1592 | G |
| 26 | BA | 1597 | A |
| 26 | BA | 1599 | U |
| 26 | BA | 1607 | G |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 26 | BA | 1612 | A |
| 26 | BA | 1613 | A |
| 26 | BA | 1614 | G |
| 26 | BA | 1616 | C |
| 26 | BA | 1622 | C |
| 26 | BA | 1635 | G |
| 26 | BA | 1638 | A |
| 26 | BA | 1665 | G |
| 26 | BA | 1667 | A |
| 26 | BA | 1673 | A |
| 26 | BA | 1679 | G |
| 26 | BA | 1680 | C |
| 26 | BA | 1681 | C |
| 26 | BA | 1698 | A |
| 26 | BA | 1704 | U |
| 26 | BA | 1705 | G |
| 26 | BA | 1707 | C |
| 26 | BA | 1712 | U |
| 26 | BA | 1713 | A |
| 26 | BA | 1733 | G |
| 26 | BA | 1755 | G |
| 26 | BA | 1799 | G |
| 26 | BA | 1800 | G |
| 26 | BA | 1809 | A |
| 26 | BA | 1818 | U |
| 26 | BA | 1822 | A |
| 26 | BA | 1827 | A |
| 26 | BA | 1836 | C |
| 26 | BA | 1847 | G |
| 26 | BA | 1869 | C |
| 26 | BA | 1883 | U |
| 26 | BA | 1884 | A |
| 26 | BA | 1899 | U |
| 26 | BA | 1924 | A |
| 26 | BA | 1927 | G |
| 26 | BA | 1928 | G |
| 26 | BA | 1929 | G |
| 26 | BA | 1931 | G |
| 26 | BA | 1932 | U |
| 26 | BA | 1933 | A |
| 26 | BA | 1941 | C |
| 26 | BA | 1942 | C |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 26 | BA | 1945 | C |
| 26 | BA | 1946 | U |
| 26 | BA | 1947 | U |
| 26 | BA | 1950 | G |
| 26 | BA | 1951 | G |
| 26 | BA | 1954 | G |
| 26 | BA | 1958 | A |
| 26 | BA | 1959 | G |
| 26 | BA | 1983 | U |
| 26 | BA | 1984 | U |
| 26 | BA | 1985 | G |
| 26 | BA | 1988 | U |
| 26 | BA | 1991 | A |
| 26 | BA | 1992 | U |
| 26 | BA | 1993 | G |
| 26 | BA | 2003 | G |
| 26 | BA | 2014 | U |
| 26 | BA | 2043 | U |
| 26 | BA | 2044 | U |
| 26 | BA | 2052 | A |
| 26 | BA | 2053 | C |
| 26 | BA | 2054 | A |
| 26 | BA | 2076 | A |
| 26 | BA | 2080 | A |
| 26 | BA | 2081 | A |
| 26 | BA | 2082 | G |
| 26 | BA | 2083 | A |
| 26 | BA | 2089 | U |
| 26 | BA | 2090 | G |
| 26 | BA | 2101 | G |
| 26 | BA | 2114 | G |
| 26 | BA | 2116 | G |
| 26 | BA | 2118 | U |
| 26 | BA | 2120 | U |
| 26 | BA | 2229 | A |
| 26 | BA | 2234 | A |
| 26 | BA | 2235 | C |
| 26 | BA | 2248 | G |
| 26 | BA | 2267 | A |
| 26 | BA | 2289 | G |
| 26 | BA | 2292 | G |
| 26 | BA | 2293 | C |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 26 | BA | 2297 | A |
| 26 | BA | 2315 | U |
| 26 | BA | 2316 | C |
| 26 | BA | 2317 | G |
| 26 | BA | 2330 | A |
| 26 | BA | 2332 | A |
| 26 | BA | 2344 | U |
| 26 | BA | 2345 | A |
| 26 | BA | 2357 | C |
| 26 | BA | 2365 | A |
| 26 | BA | 2366 | G |
| 26 | BA | 2375 | U |
| 26 | BA | 2376 | G |
| 26 | BA | 2377 | G |
| 26 | BA | 2378 | A |
| 26 | BA | 2380 | G |
| 26 | BA | 2396 | G |
| 26 | BA | 2398 | U |
| 26 | BA | 2420 | U |
| 26 | BA | 2438 | G |
| 26 | BA | 2443 | A |
| 26 | BA | 2444 | A |
| 26 | BA | 2452 | C |
| 26 | BA | 2458 | G |
| 26 | BA | 2459 | A |
| 26 | BA | 2463 | U |
| 26 | BA | 2464 | U |
| 26 | BA | 2475 | C |
| 26 | BA | 2479 | A |
| 26 | BA | 2481 | G |
| 26 | BA | 2485 | A |
| 26 | BA | 2487 | A |
| 26 | BA | 2505 | G |
| 26 | BA | 2509 | C |
| 26 | BA | 2513 | G |
| 26 | BA | 2517 | U |
| 26 | BA | 2518 | C |
| 26 | BA | 2529 | A |
| 26 | BA | 2540 | G |
| 26 | BA | 2564 | G |
| 26 | BA | 2565 | U |
| 26 | BA | 2577 | A |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 26 | BA | 2578 | G |
| 26 | BA | 2584 | C |
| 26 | BA | 2589 | G |
| 26 | BA | 2593 | G |
| 26 | BA | 2610 | G |
| 26 | BA | 2613 | A |
| 26 | BA | 2614 | G |
| 26 | BA | 2620 | U |
| 26 | BA | 2621 | U |
| 26 | BA | 2632 | A |
| 26 | BA | 2636 | G |
| 26 | BA | 2641 | C |
| 26 | BA | 2642 | G |
| 26 | BA | 2651 | G |
| 26 | BA | 2656 | A |
| 26 | BA | 2657 | G |
| 26 | BA | 2677 | A |
| 26 | BA | 2701 | U |
| 26 | BA | 2724 | G |
| 26 | BA | 2736 | C |
| 26 | BA | 2742 | C |
| 26 | BA | 2743 | A |
| 26 | BA | 2748 | U |
| 26 | BA | 2757 | A |
| 26 | BA | 2773 | A |
| 26 | BA | 2774 | A |
| 26 | BA | 2780 | A |
| 26 | BA | 2785 | A |
| 26 | BA | 2786 | A |
| 26 | BA | 2787 | A |
| 26 | BA | 2799 | U |
| 26 | BA | 2811 | U |
| 26 | BA | 2824 | U |
| 26 | BA | 2849 | A |
| 26 | BA | 2850 | G |
| 26 | BA | 2851 | G |
| 26 | BA | 2853 | A |
| 26 | BA | 2864 | C |
| 26 | BA | 2870 | G |
| 26 | BA | 2871 | C |
| 26 | BA | 2877 | C |
| 26 | BA | 2883 | A |

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| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 26 | BA | 2889 | U |
| 26 | BA | 2890 | C |
| 26 | BA | 2891 | C |
| 26 | BA | 2892 | U |
| 26 | BA | 2893 | U |
| 26 | BA | 2894 | U |
| 26 | BA | 2895 | C |
| 26 | BA | 2897 | C |
| 27 | BB | 10 | G |
| 27 | BB | 25 | G |
| 27 | BB | 26 | C |
| 27 | BB | 36 | A |
| 27 | BB | 42 | C |
| 27 | BB | 57 | U |
| 27 | BB | 58 | A |
| 27 | BB | 98 | C |
| 27 | BB | 115 | C |
| 27 | BB | 116 | G |
| 27 | BB | 125 | C |
| 27 | BB | 127 | C |
| 27 | BB | 128 | A |

All (10) RNA pucker outliers are listed below:

| Mol | Chain | Res | Type |
|-----|-------|------|------|
| 1 | AA | 628 | A |
| 1 | AA | 1063 | G |
| 1 | AA | 1097 | U |
| 1 | AA | 1187 | U |
| 26 | BA | 817 | U |
| 26 | BA | 826 | U |
| 26 | BA | 837 | A |
| 26 | BA | 869 | U |
| 26 | BA | 1932 | U |
| 26 | BA | 1984 | U |

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

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

5.5 Carbohydrates [i](#)

There are no oligosaccharides in this entry.

5.6 Ligand geometry [i](#)

Of 60 ligands modelled in this entry, 60 are monoatomic - leaving 0 for Mogul analysis.

There are no bond length outliers.

There are no bond angle outliers.

There are no chirality outliers.

There are no torsion outliers.

There are no ring outliers.

No monomer is involved in short contacts.

5.7 Other polymers [i](#)

There are no such residues in this entry.

5.8 Polymer linkage issues [i](#)

There are no chain breaks in this entry.

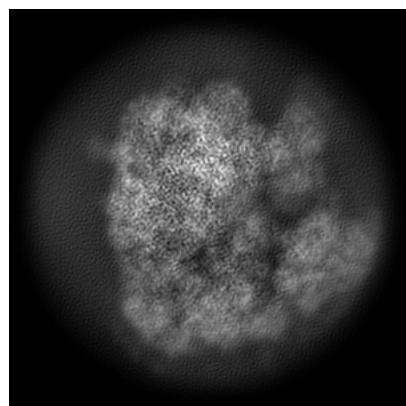
6 Map visualisation [i](#)

This section contains visualisations of the EMDB entry EMD-49998. These allow visual inspection of the internal detail of the map and identification of artifacts.

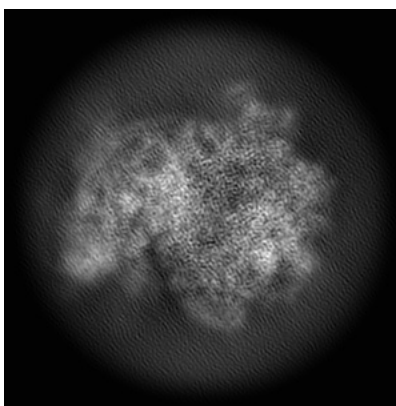
Images derived from a raw map, generated by summing the deposited half-maps, are presented below the corresponding image components of the primary map to allow further visual inspection and comparison with those of the primary map.

6.1 Orthogonal projections [i](#)

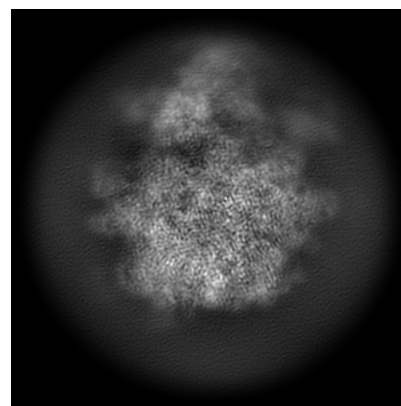
6.1.1 Primary map



X

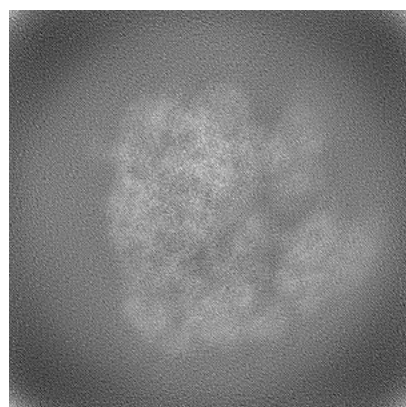


Y

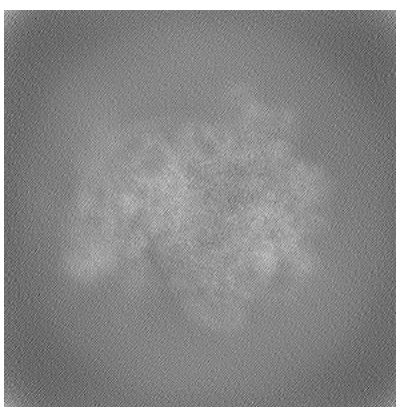


Z

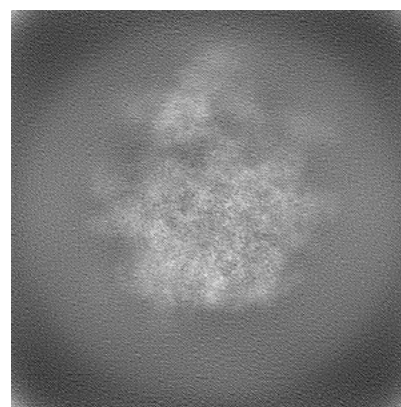
6.1.2 Raw map



X



Y

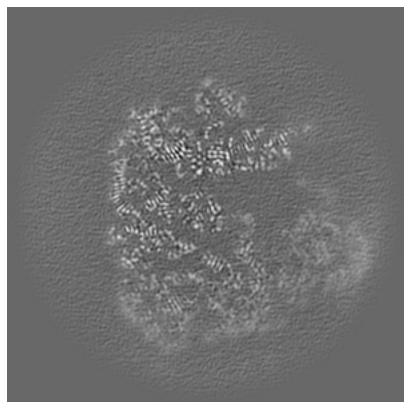


Z

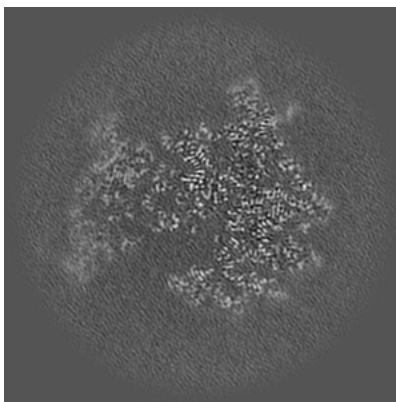
The images above show the map projected in three orthogonal directions.

6.2 Central slices [i](#)

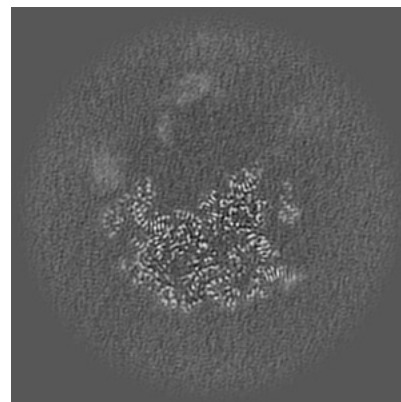
6.2.1 Primary map



X Index: 240

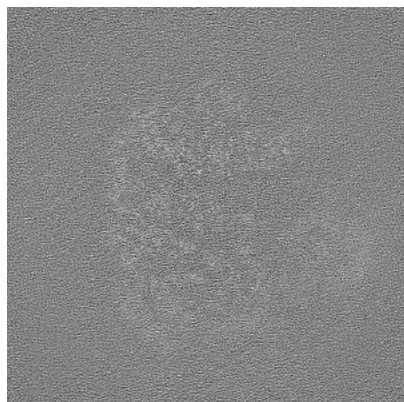


Y Index: 240

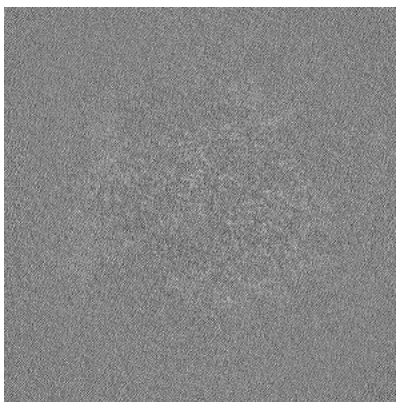


Z Index: 240

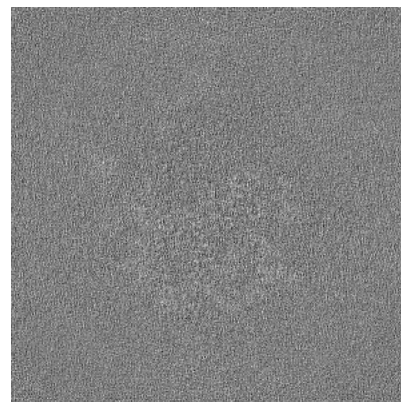
6.2.2 Raw map



X Index: 240



Y Index: 240

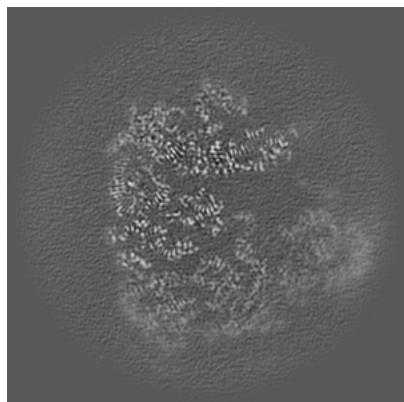


Z Index: 240

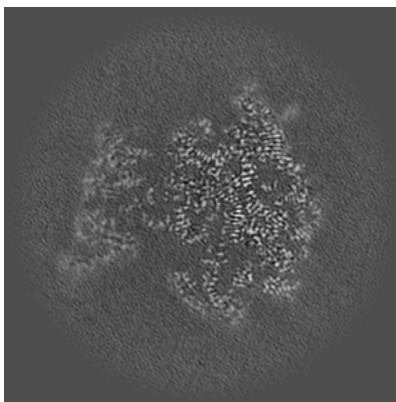
The images above show central slices of the map in three orthogonal directions.

6.3 Largest variance slices [i](#)

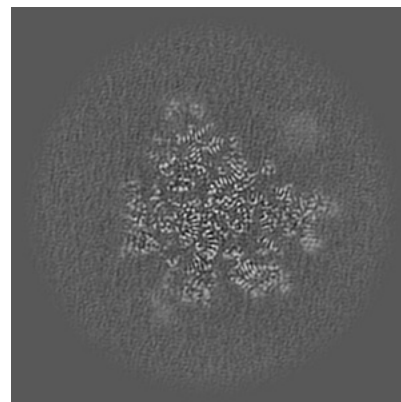
6.3.1 Primary map



X Index: 242

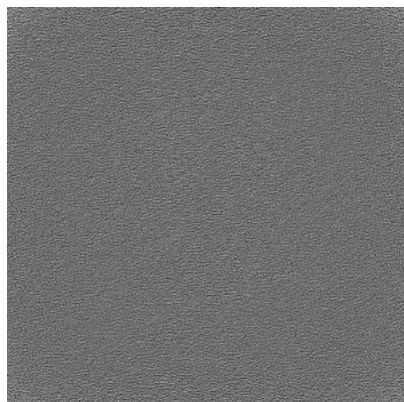


Y Index: 231

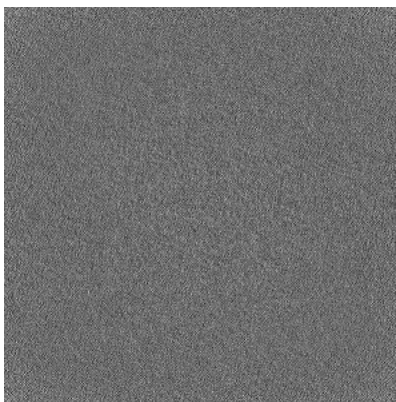


Z Index: 299

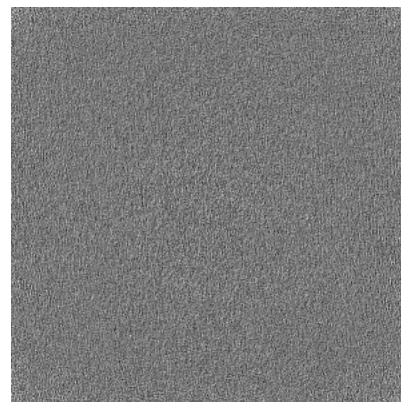
6.3.2 Raw map



X Index: 0



Y Index: 0

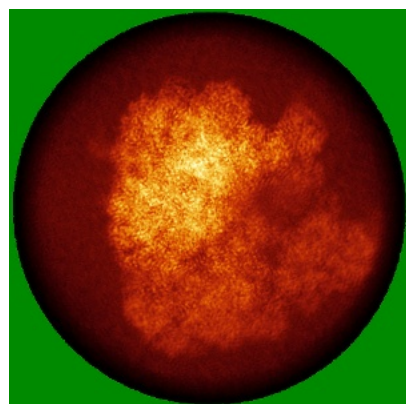


Z Index: 0

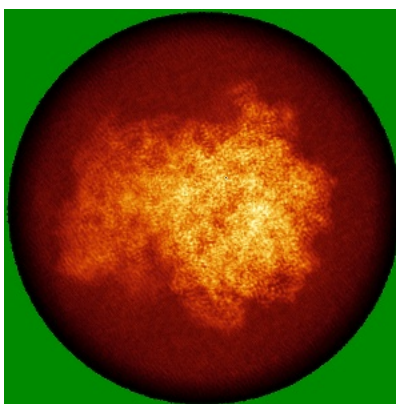
The images above show the largest variance slices of the map in three orthogonal directions.

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

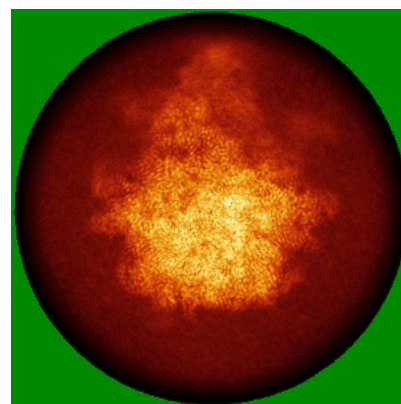
6.4.1 Primary map



X

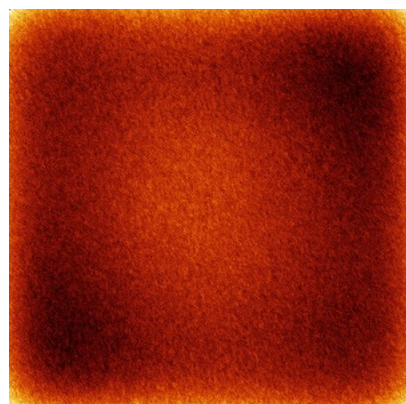


Y

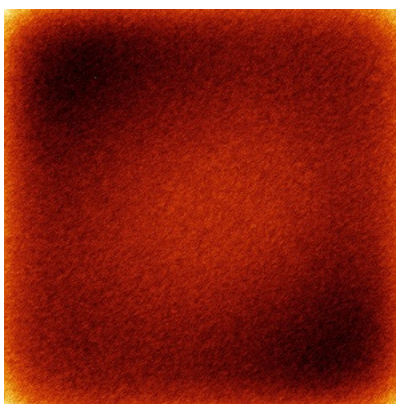


Z

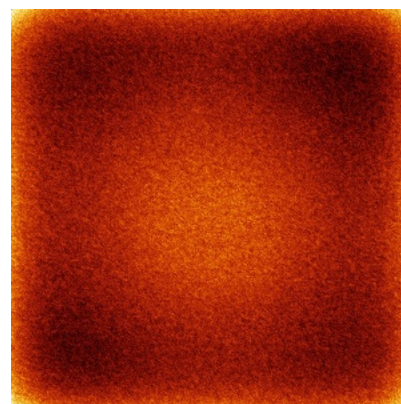
6.4.2 Raw map



X



Y

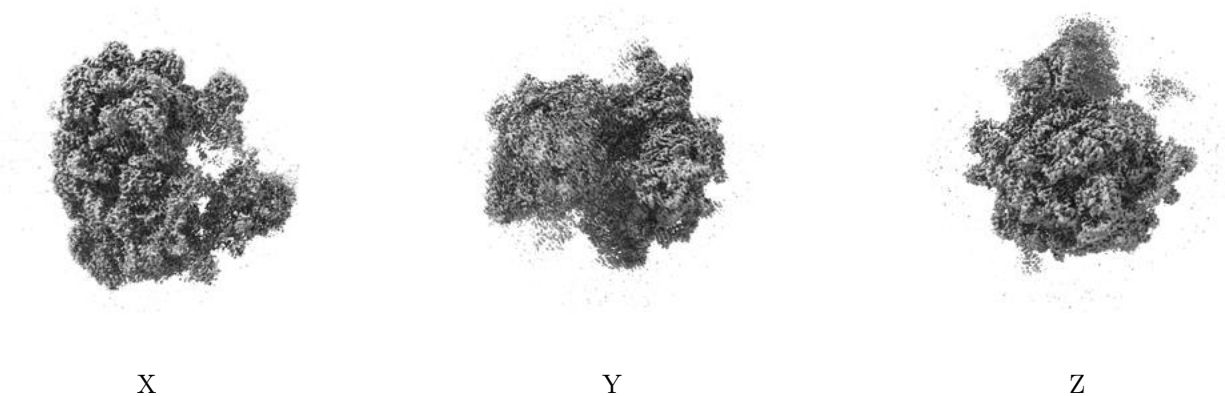


Z

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.

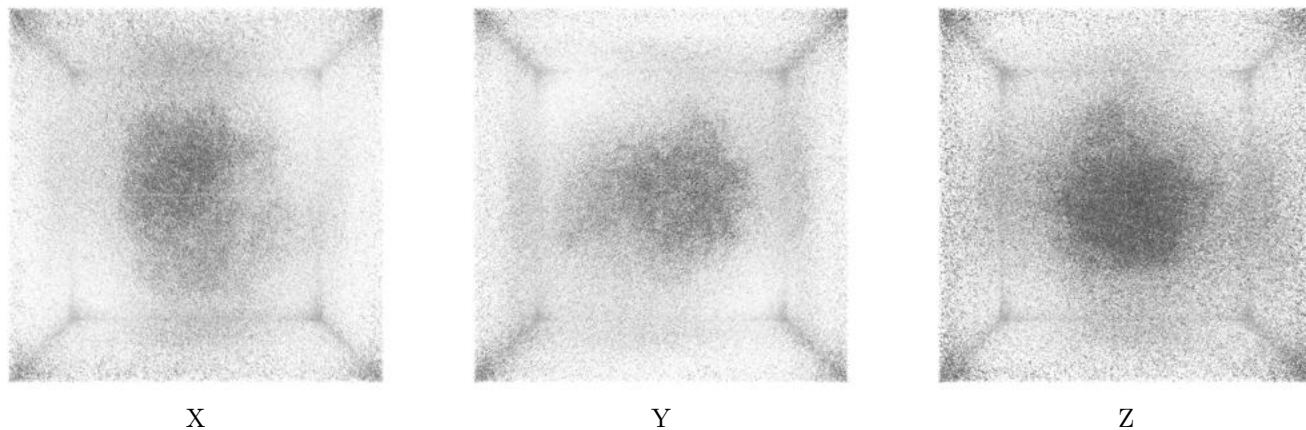
6.5 Orthogonal surface views [i](#)

6.5.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 0.046. 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.

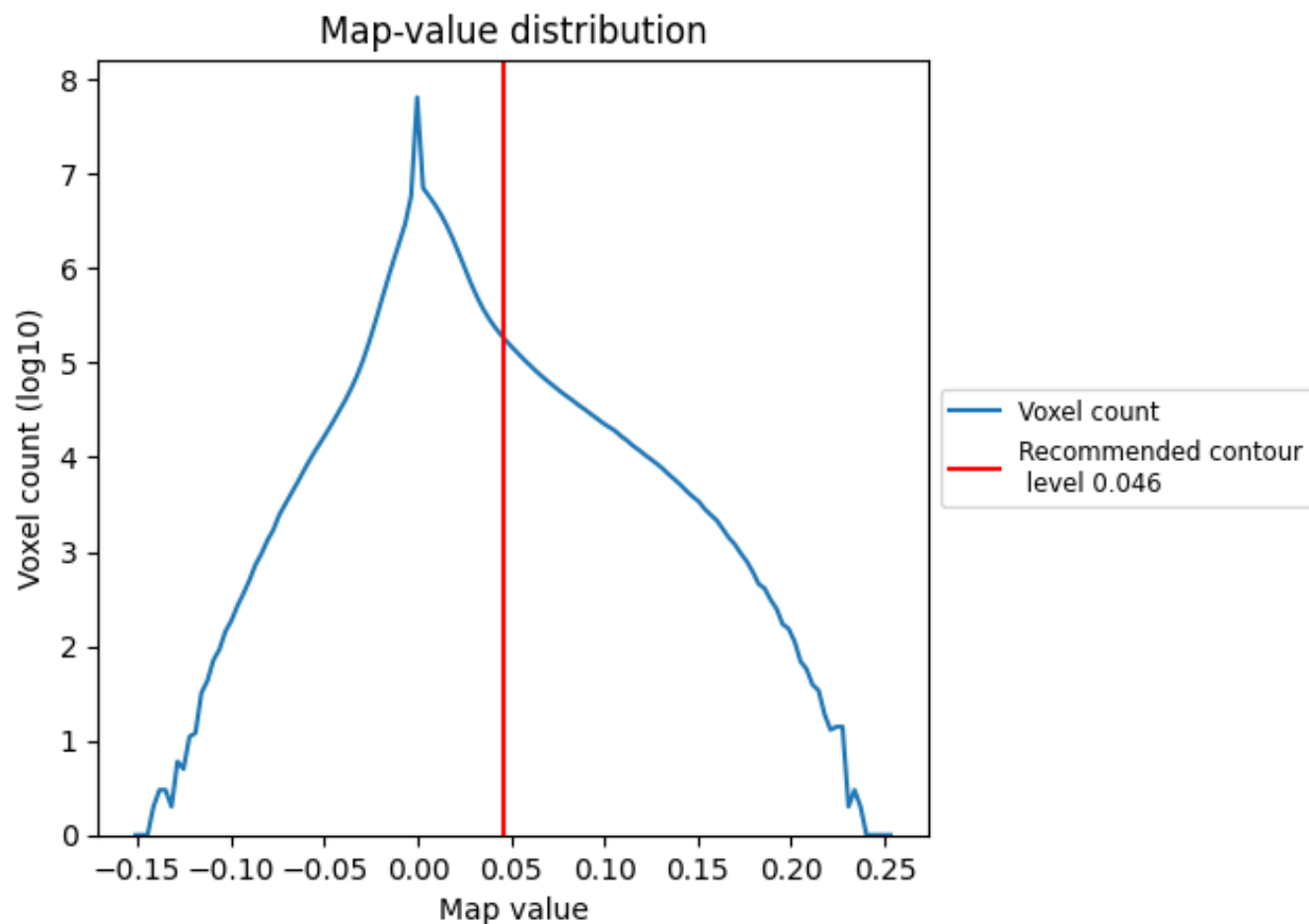
6.6 Mask visualisation [i](#)

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

7 Map analysis [i](#)

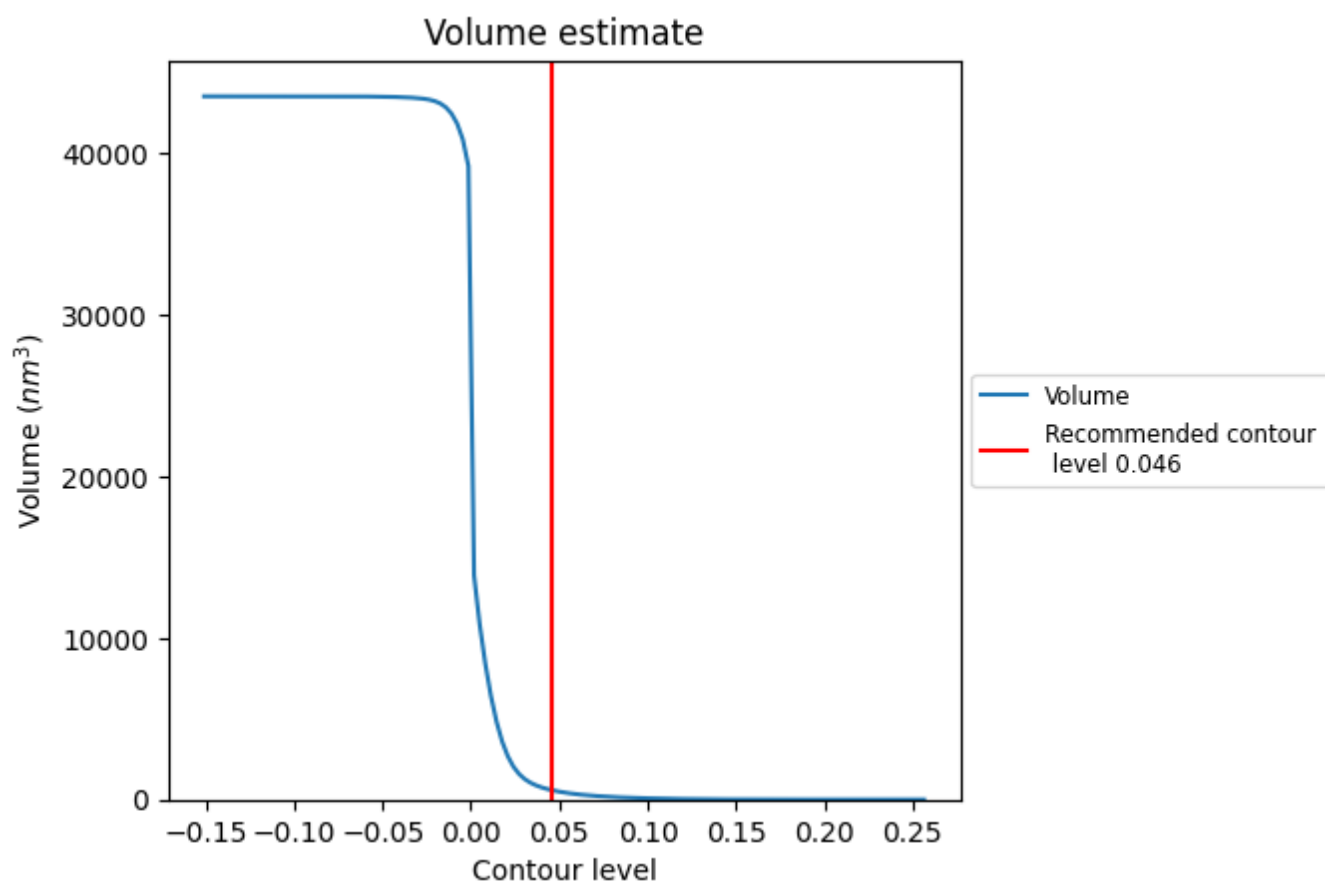
This section contains the results of statistical analysis of the map.

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.

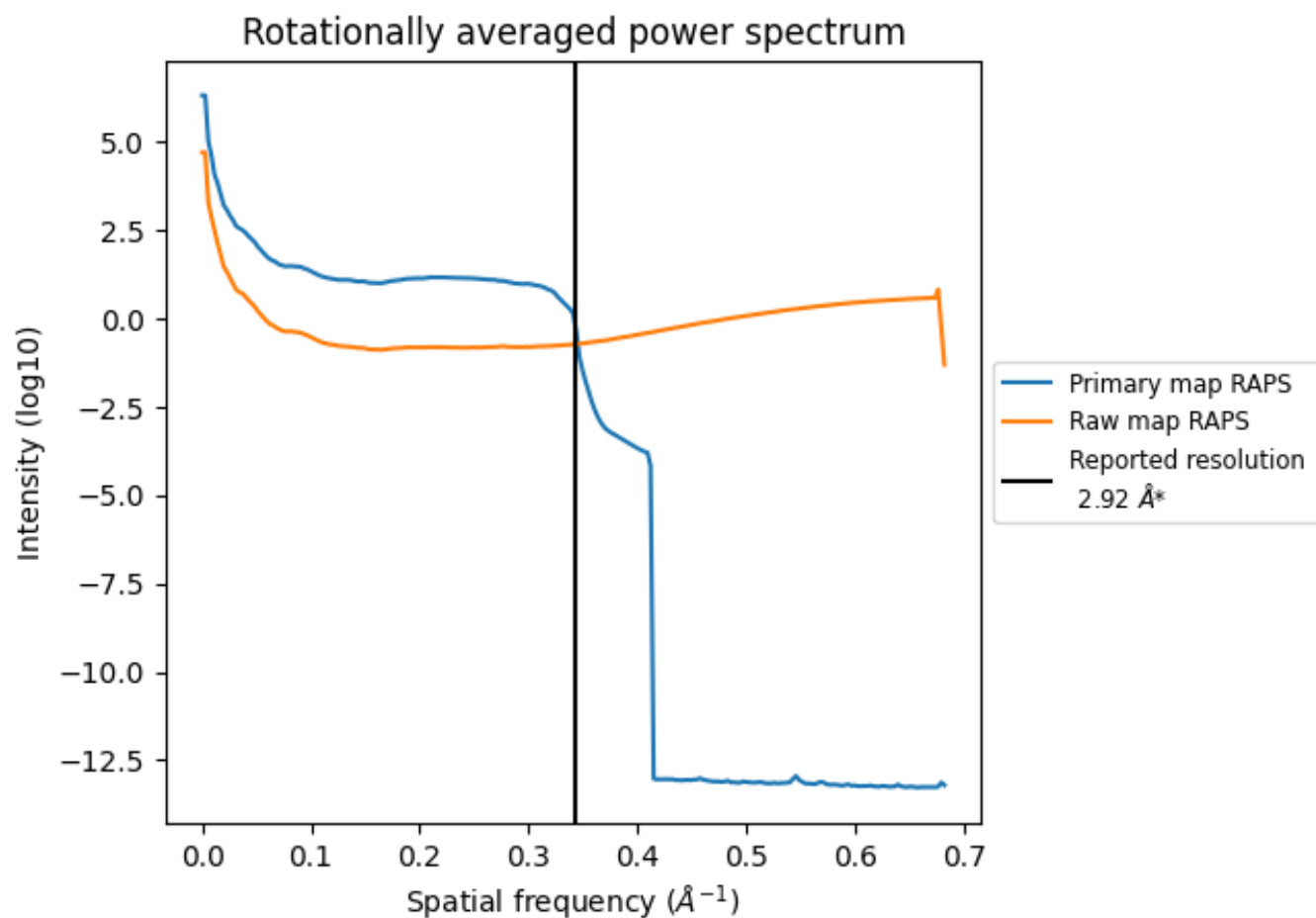
7.2 Volume estimate [i](#)



The volume at the recommended contour level is 578 nm³; this corresponds to an approximate mass of 522 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 ⓘ

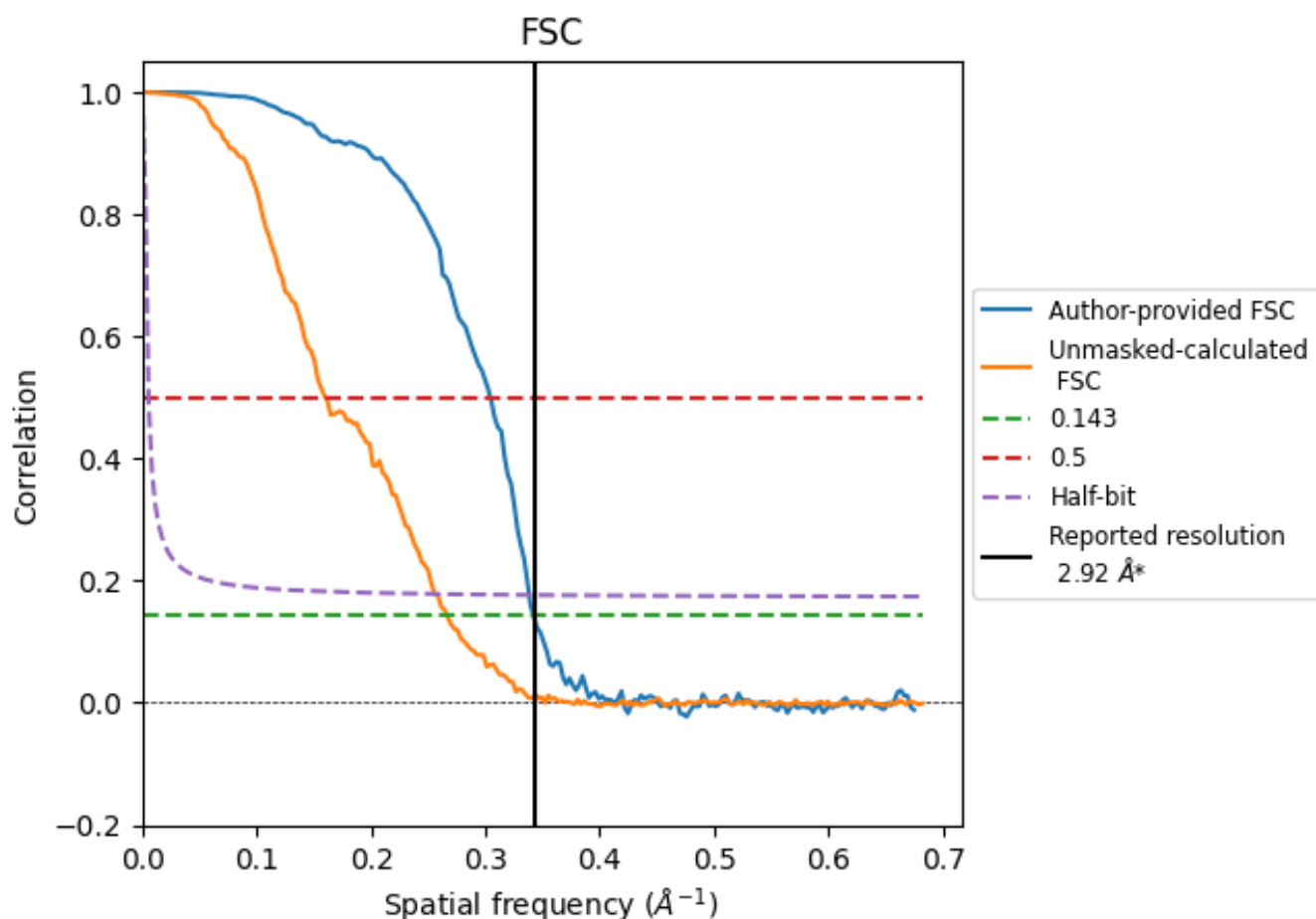


*Reported resolution corresponds to spatial frequency of 0.342 Å⁻¹

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 0.342 \AA^{-1}

8.2 Resolution estimates [i](#)

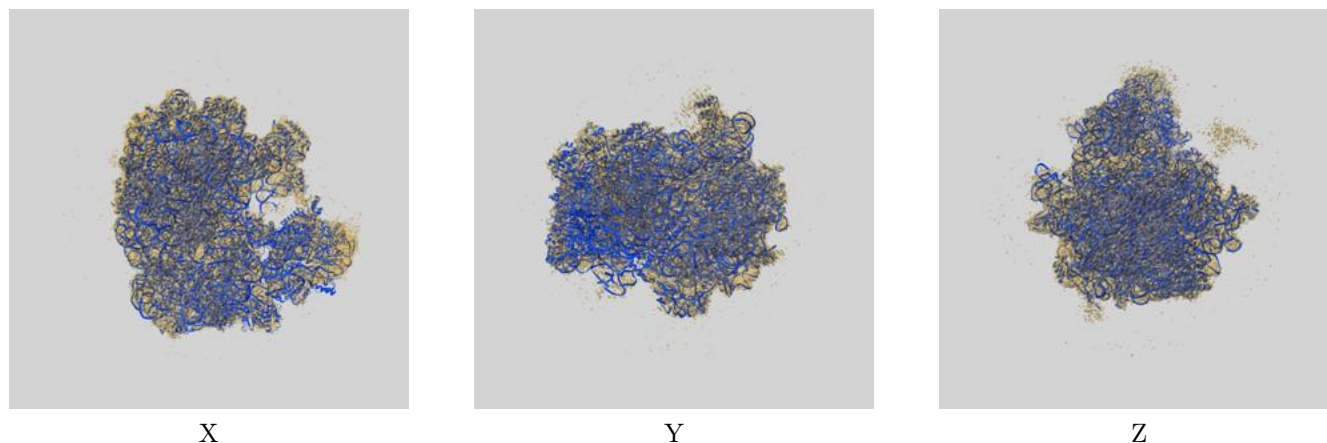
| Resolution estimate (Å) | Estimation criterion (FSC cut-off) | | |
|---------------------------|------------------------------------|------|----------|
| | 0.143 | 0.5 | Half-bit |
| Reported by author | 2.92 | - | - |
| Author-provided FSC curve | 2.92 | 3.28 | 2.95 |
| Unmasked-calculated* | 3.75 | 6.24 | 3.91 |

*Resolution estimate based on FSC curve calculated by comparison of deposited half-maps. The value from deposited half-maps intersecting FSC 0.143 CUT-OFF 3.75 differs from the reported value 2.92 by more than 10 %

9 Map-model fit [i](#)

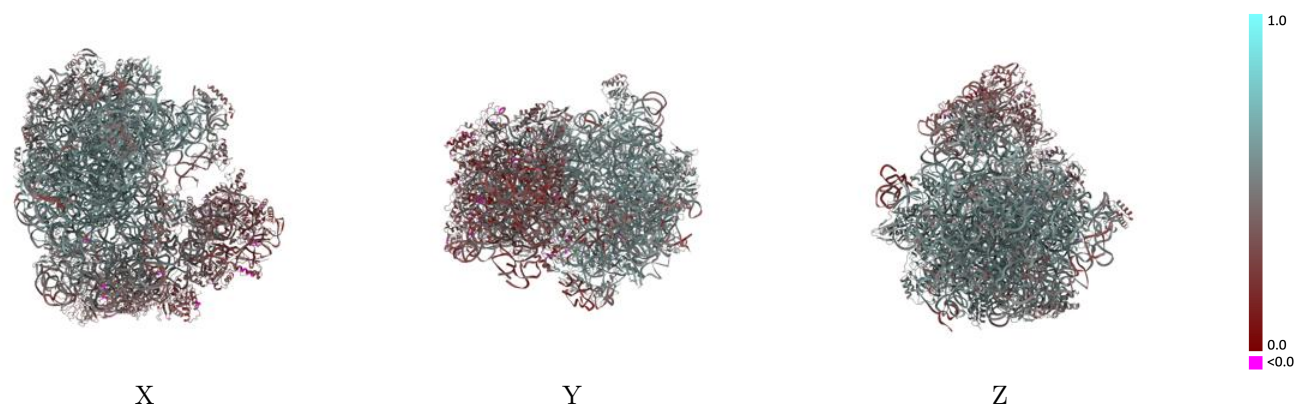
This section contains information regarding the fit between EMDB map EMD-49998 and PDB model 9O17. Per-residue inclusion information can be found in section [3](#) on page [17](#).

9.1 Map-model overlay [i](#)



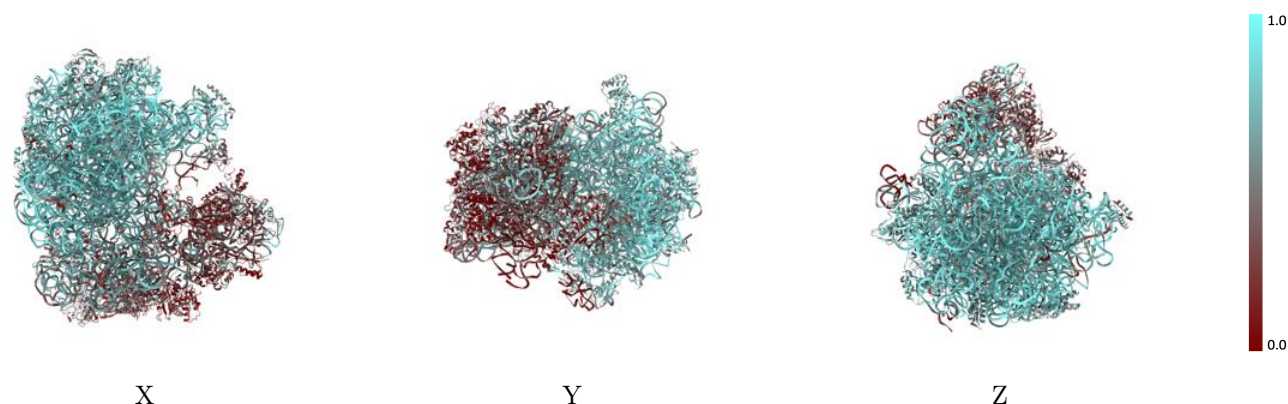
The images above show the 3D surface view of the map at the recommended contour level 0.046 at 50% transparency in yellow overlaid with a ribbon representation of the model coloured in blue. These images allow for the visual assessment of the quality of fit between the atomic model and the map.

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



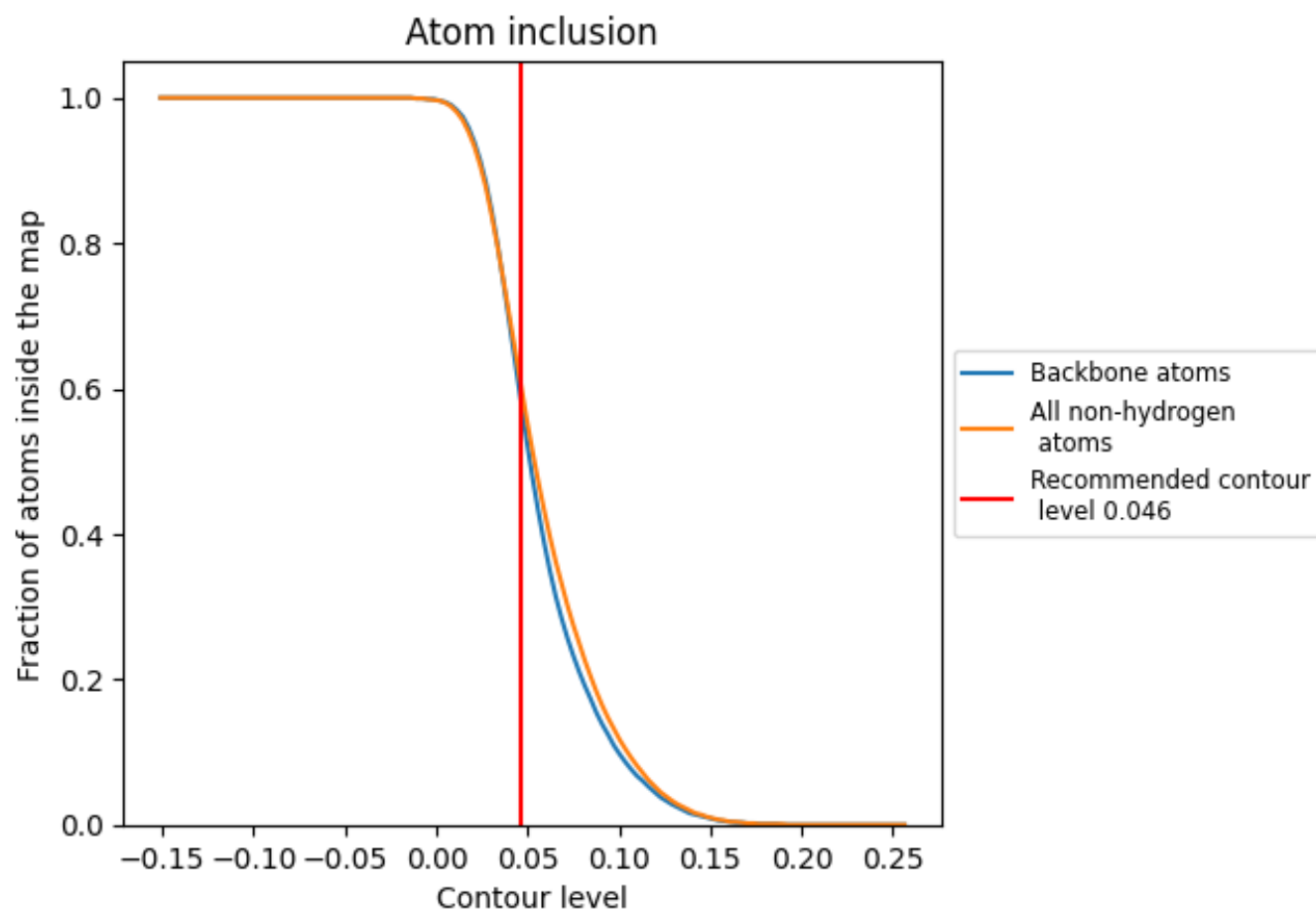
The images above show the model with each residue coloured according its Q-score. This shows their resolvability in the map with higher Q-score values reflecting better resolvability. Please note: Q-score is calculating the resolvability of atoms, and thus high values are only expected at resolutions 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 (0.046).







































































9.4 Atom inclusion [i](#)



At the recommended contour level, 58% of all backbone atoms, 61% 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 (0.046) and Q-score for the entire model and for each chain.

| Chain | Atom inclusion | Q-score |
|-------|--|--|
| All |  0.6090 |  0.4870 |
| AA |  0.5390 |  0.4340 |
| AB |  0.1810 |  0.3070 |
| AC |  0.0270 |  0.2910 |
| AD |  0.3400 |  0.4010 |
| AE |  0.2570 |  0.3420 |
| AF |  0.3430 |  0.3760 |
| AG |  0.2750 |  0.3640 |
| AH |  0.2990 |  0.4110 |
| AI |  0.2170 |  0.3360 |
| AJ |  0.3830 |  0.3900 |
| AK |  0.4860 |  0.4770 |
| AL |  0.2690 |  0.2830 |
| AM |  0.0800 |  0.2550 |
| AN |  0.3680 |  0.4470 |
| AO |  0.2740 |  0.4550 |
| AP |  0.1400 |  0.3260 |
| AQ |  0.3890 |  0.4190 |
| AR |  0.0910 |  0.3490 |
| AS |  0.4110 |  0.4610 |
| AT |  0.2080 |  0.3030 |
| AU |  0.0780 |  0.2530 |
| AV |  0.3450 |  0.2710 |
| AW |  0.2640 |  0.3030 |
| AX |  0.2530 |  0.3590 |
| AY |  0.2030 |  0.3460 |
| AZ |  0.0510 |  0.2900 |
| BA |  0.8310 |  0.5490 |
| BB |  0.8250 |  0.4990 |
| BC |  0.7420 |  0.5760 |
| BD |  0.7510 |  0.5630 |
| BE |  0.7060 |  0.5040 |
| BF |  0.4010 |  0.4120 |
| BG |  0.6260 |  0.5050 |
| BH |  0.6070 |  0.4680 |



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| Chain | Atom inclusion | Q-score |
|-------|--|--|
| BI |  0.6580 |  0.5550 |
| BJ |  0.7300 |  0.5460 |
| BK |  0.6390 |  0.5630 |
| BL |  0.7140 |  0.5410 |
| BM |  0.7560 |  0.5700 |
| BN |  0.7020 |  0.4900 |
| BO |  0.7040 |  0.5120 |
| BP |  0.7110 |  0.5430 |
| BQ |  0.6050 |  0.5410 |
| BR |  0.7430 |  0.5680 |
| BS |  0.6700 |  0.5280 |
| BT |  0.6770 |  0.5120 |
| BU |  0.6380 |  0.4530 |
| BV |  0.6370 |  0.5560 |
| BW |  0.5540 |  0.4290 |
| BX |  0.7340 |  0.5350 |
| BY |  0.5530 |  0.4610 |
| BZ |  0.7340 |  0.5530 |
| Ba |  0.7070 |  0.5170 |
| Bb |  0.6260 |  0.5310 |
| Bc |  0.8120 |  0.5750 |
| Bd |  0.7470 |  0.5500 |
| Be |  0.6310 |  0.5440 |
| Bf |  0.7090 |  0.5730 |