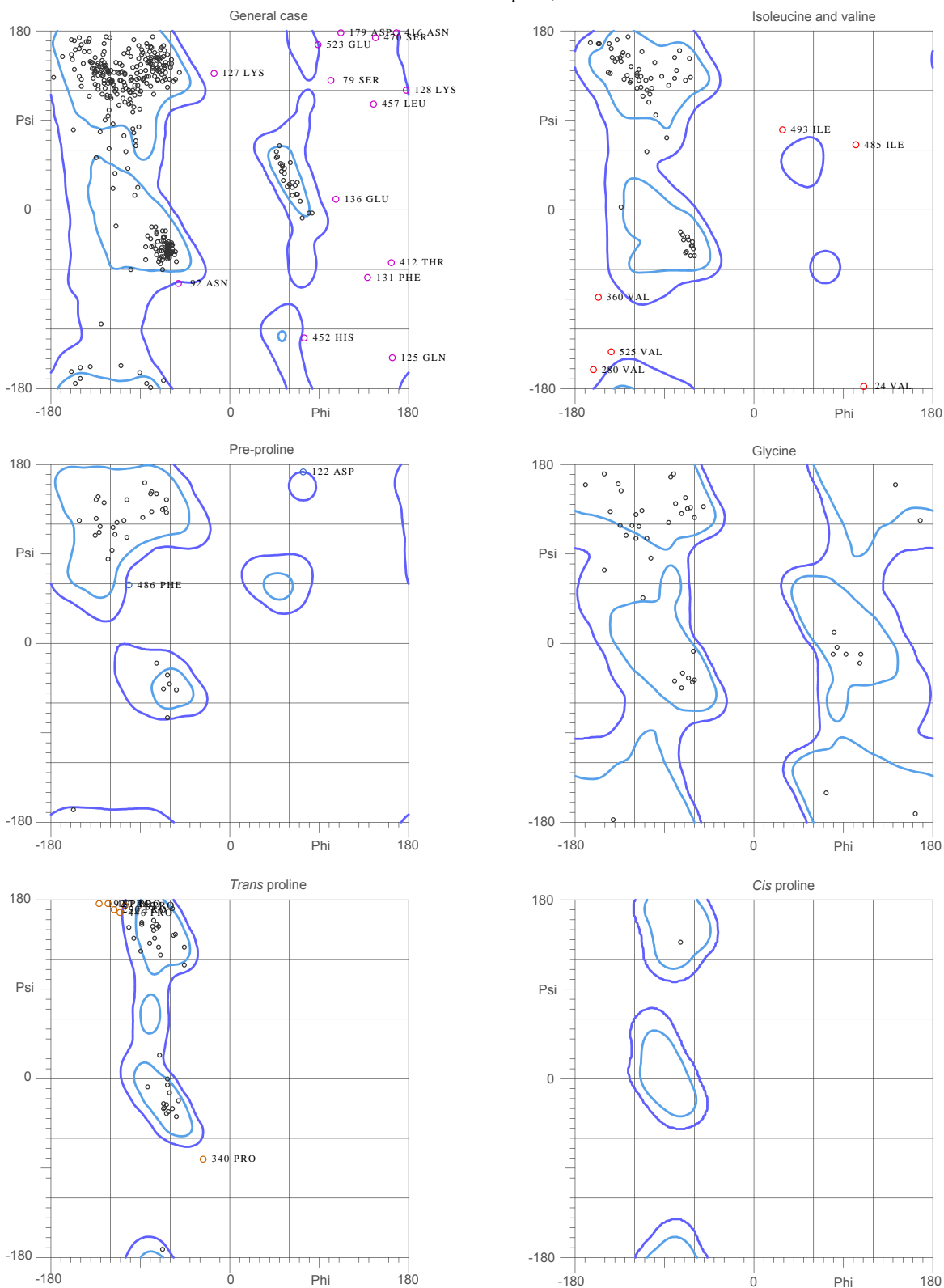


MolProbity Ramachandran analysis

1aozA.B99990024.pdb, model 1



87.5% (270/309) of all residues were in favored (98%) regions.
10.5% (34/309) of all residues were in allowed (1.9% RB) regions.

There were 30 outliers (Phi, Psi):

| |
|------------------------|
| 76 VAL (11.1, 17.8) |
| 51 PRO (166.7, 179.2) |
| 100 ASP (165.2, 159.2) |
| 57 ASP (167.1, 75.0) |
| 124 ASP (178.1, 179.2) |
| 131 GLN (138.4, 149.1) |
| 105 VAL (178.1, 159.2) |
| 98 VAL (178.1, 159.2) |
| 131 ASP (179.4, 168.4) |
| 130 ASP (179.4, 15.1) |
| 177 ASP (113.5, 169.2) |
| 200 PRO (113.5, 171.0) |
| 78 VAL (168.1, 169.2) |
| 200 PRO (171.0, 171.0) |
| 957 PRO (171.0, 171.0) |
| 958 PRO (171.0, 171.0) |
| 959 PRO (171.0, 171.0) |
| 960 PRO (171.0, 171.0) |
| 961 PRO (171.0, 171.0) |
| 340 PRO (171.0, 171.0) |
| 122 ASP (171.0, 171.0) |
| 486 PHE (171.0, 171.0) |
| 125 GLN (171.0, 171.0) |
| 412 THR (171.0, 171.0) |
| 131 PHE (171.0, 171.0) |
| 452 HIS (171.0, 171.0) |
| 92 ASN (171.0, 171.0) |
| 136 GLU (171.0, 171.0) |
| 457 LEU (171.0, 171.0) |
| 128 LYS (171.0, 171.0) |
| 79 SER (171.0, 171.0) |
| 523 GLU (171.0, 171.0) |
| 179 ASP (171.0, 171.0) |
| 127 LYS (171.0, 171.0) |