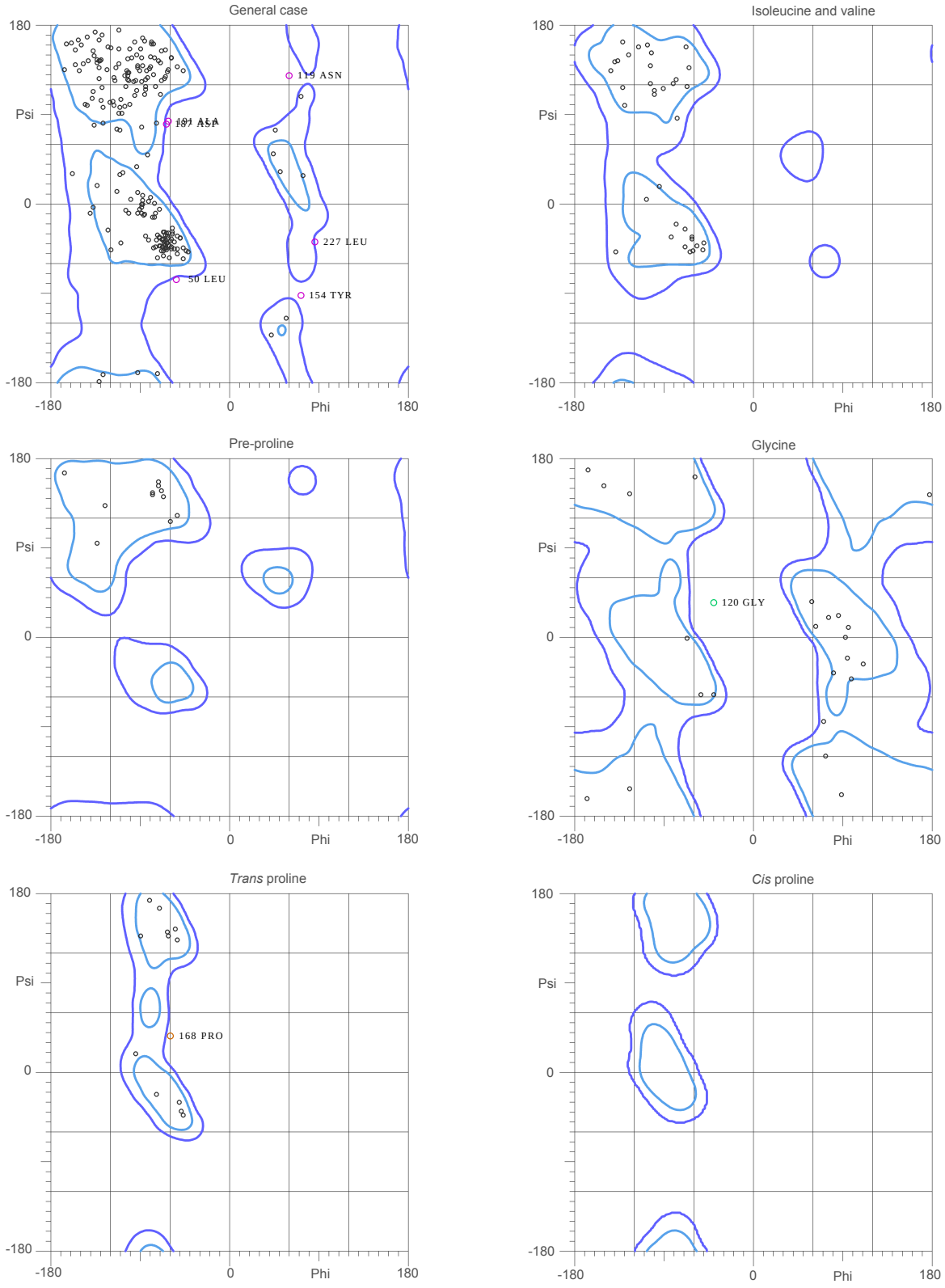
MolProbity Ramachandran analysis

3d62.H.pdb, model 1



90.2% (268/297) of all residues were in favored (98%) regions.
97.3% (289/297) of all residues were in allowed (>99.8%) regions.

There were 8 outliers (phi, psi):
- 50 LEU (-54.3, -76.6)
- 119 ASN (60.3, 130.4)
- 120 GLY (-40.6, 35.4)
- 154 TYR (72.6, -92.8)
- 168 PRO (-60.2, 37.4)
- 187 ASP (-64.5, 82.0)
- 191 ALA (-62.5, 84.5)
- 227 LEU (86.3, -38.3)