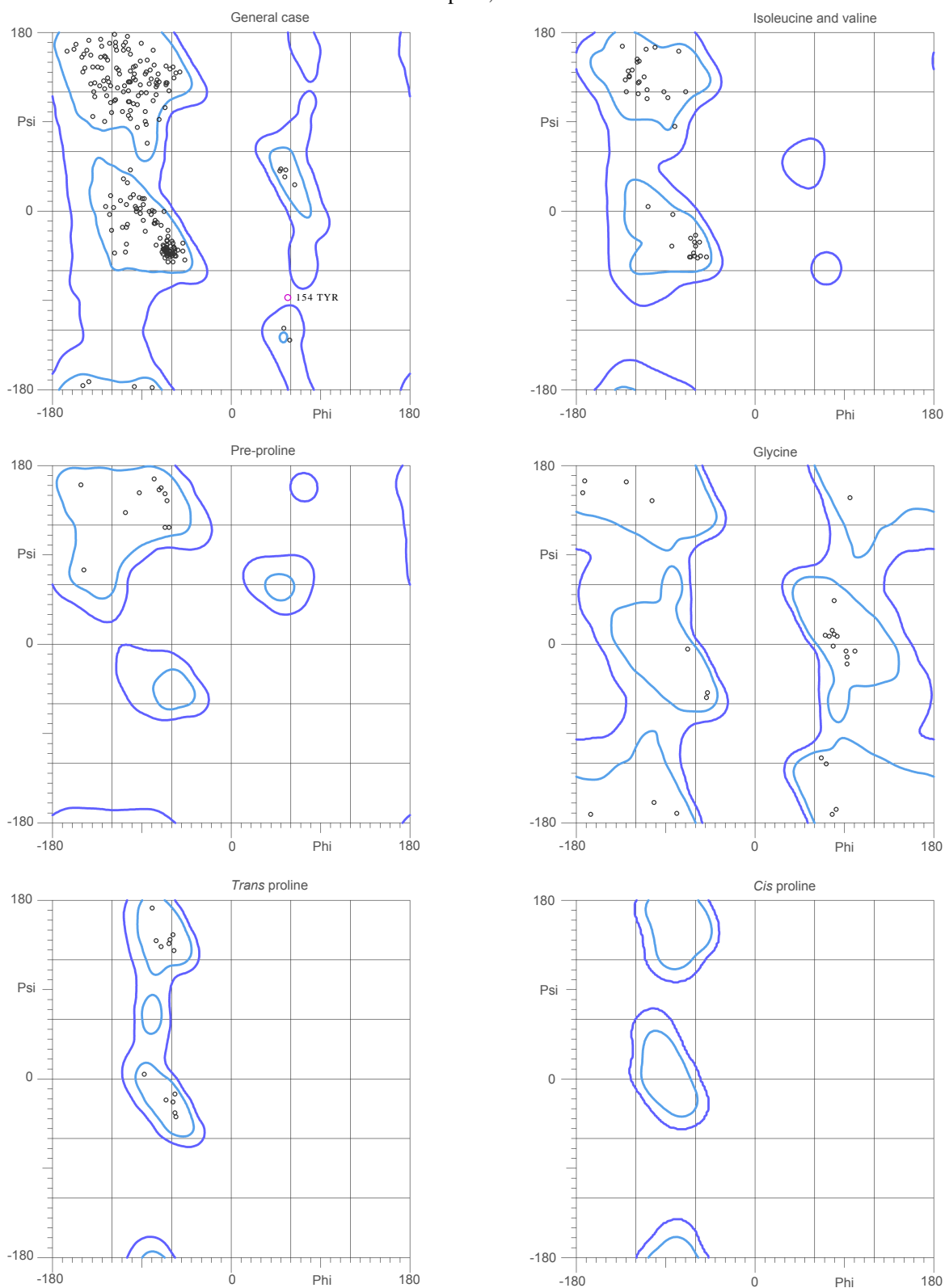


MolProbity Ramachandran analysis

5r15.H.pdb, model 1



98.0% (297/303) of all residues were in favored (98%) regions.

99.7% (302/303) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (57.7, -87.7)