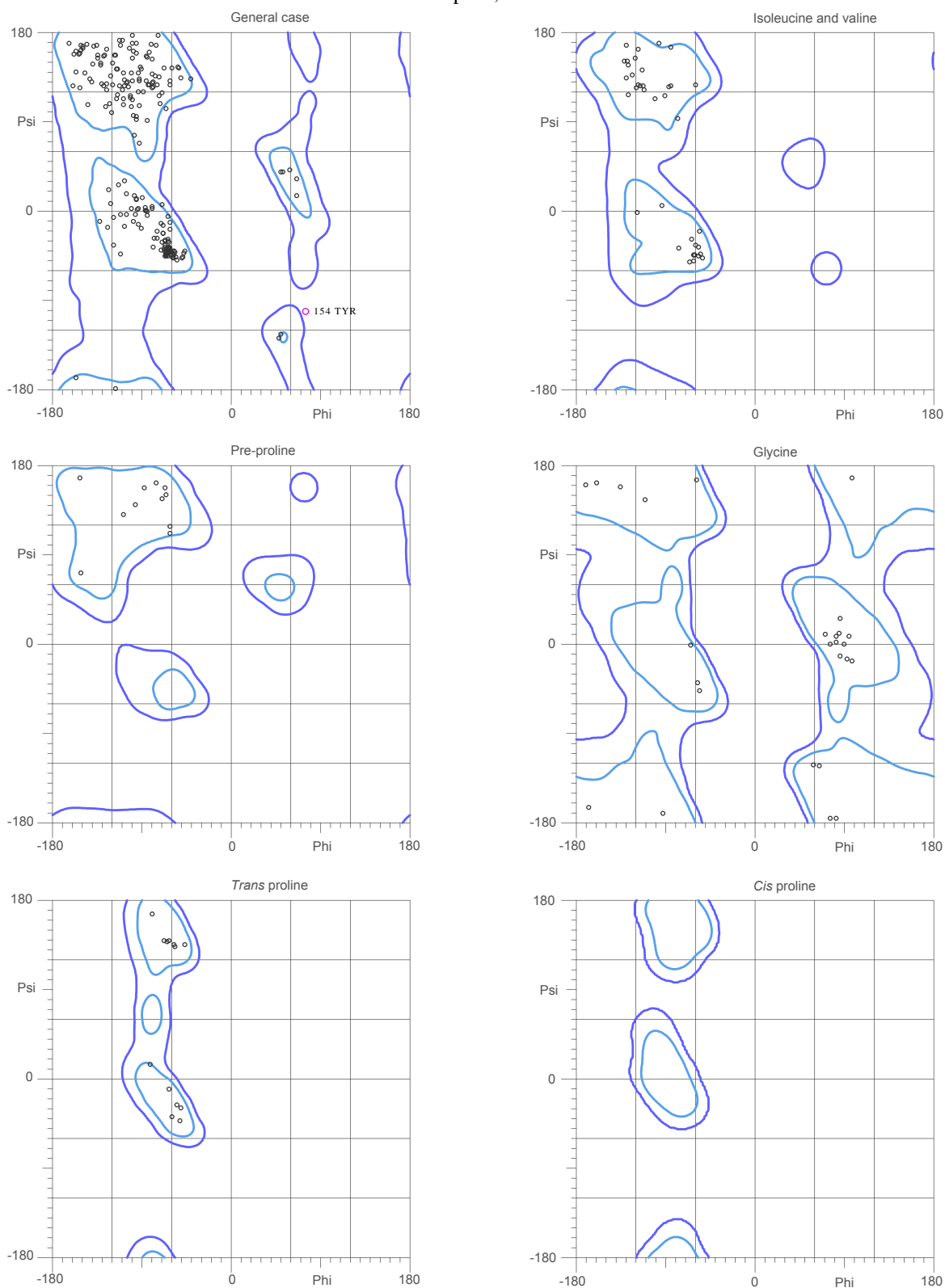


MolProbity Ramachandran analysis

6w63.H.pdb, model 1



97.7% (296/303) 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 (ϕ , ψ):

154 TYR (76.0, -101.2)