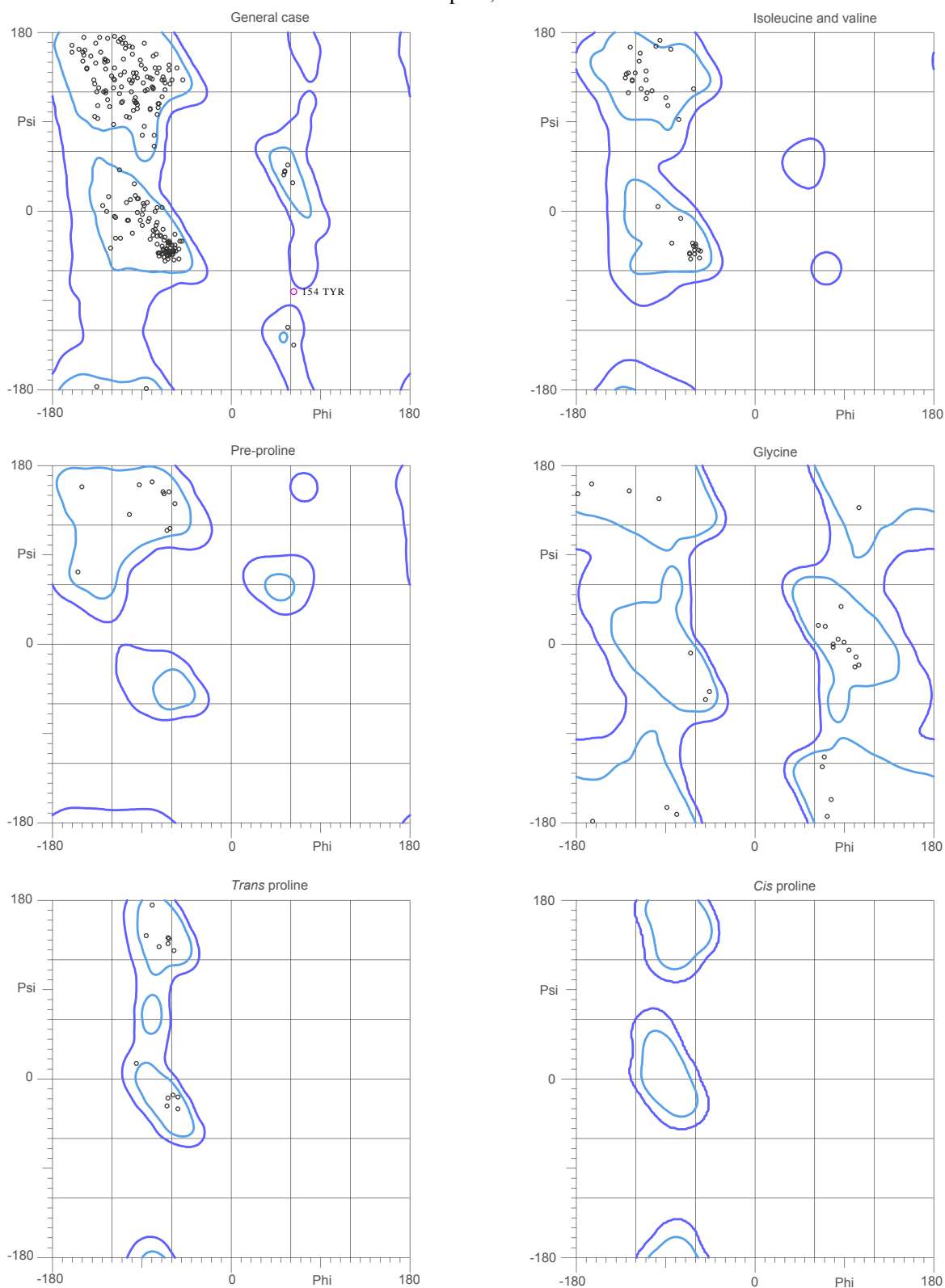


MolProbity Ramachandran analysis

5rh8.H.pdb, model 1



97.4% (296/304) of all residues were in favored (98%) regions.

99.7% (303/304) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

154 TYR (63.3, -81.3)