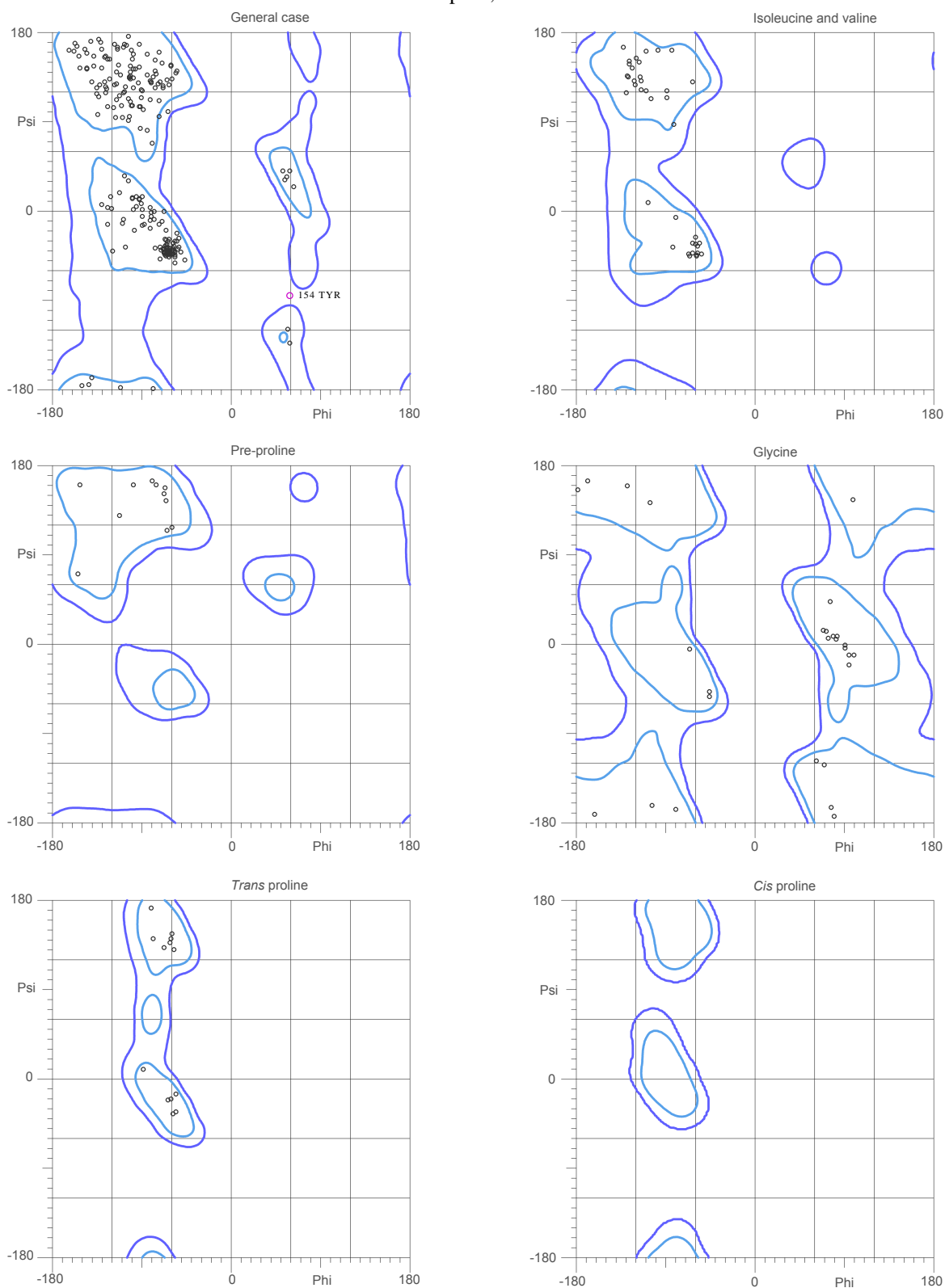


MolProbity Ramachandran analysis

5rhb.H.pdb, model 1



98.7% (304/308) of all residues were in favored (98%) regions.

99.7% (307/308) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

154 TYR (59.2, -85.1)