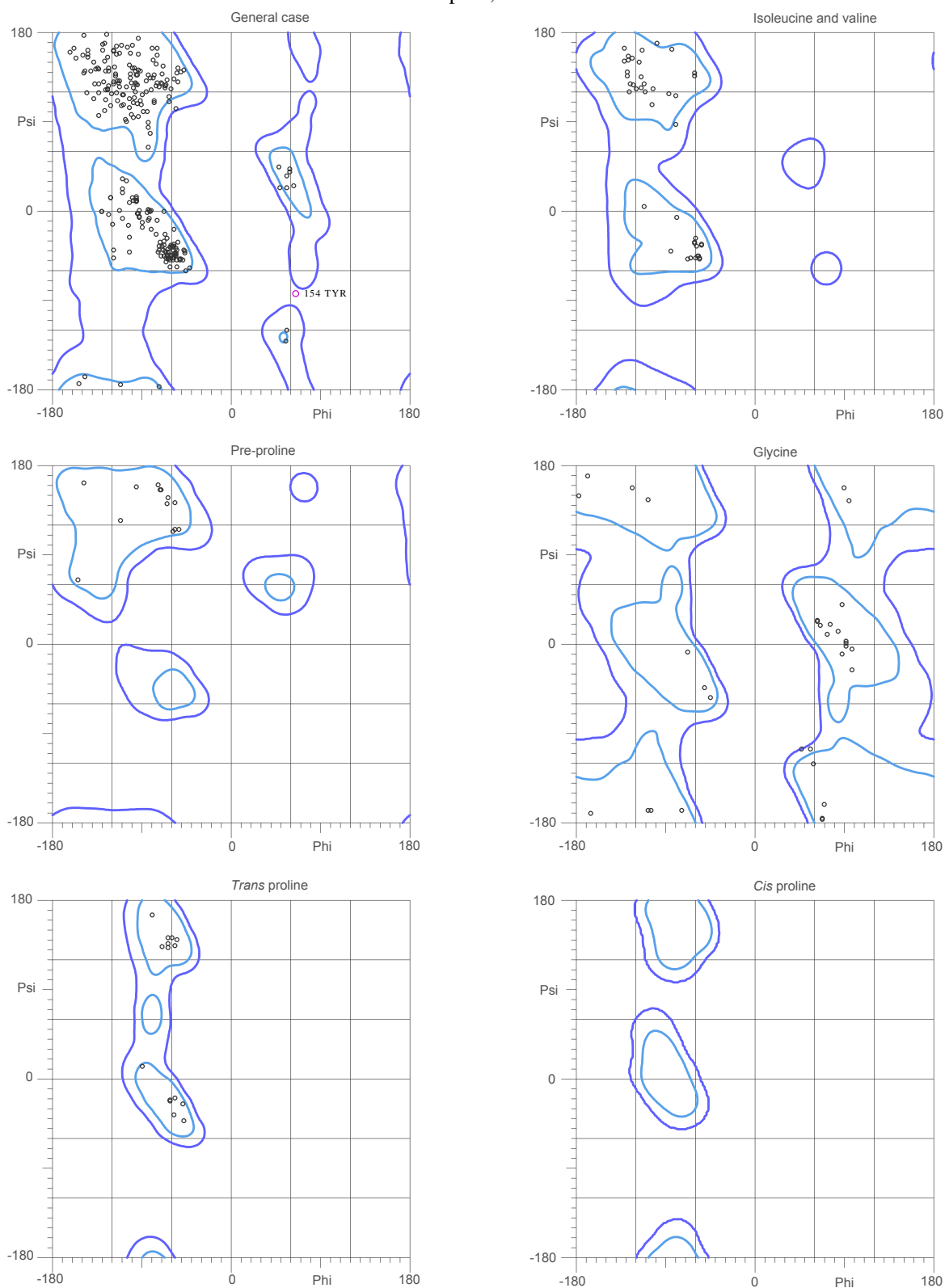


MolProbity Ramachandran analysis

5rea.H.pdb, model 1



97.5% (346/355) of all residues were in favored (98%) regions.

99.7% (354/355) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

154 TYR (65.1, -83.5)