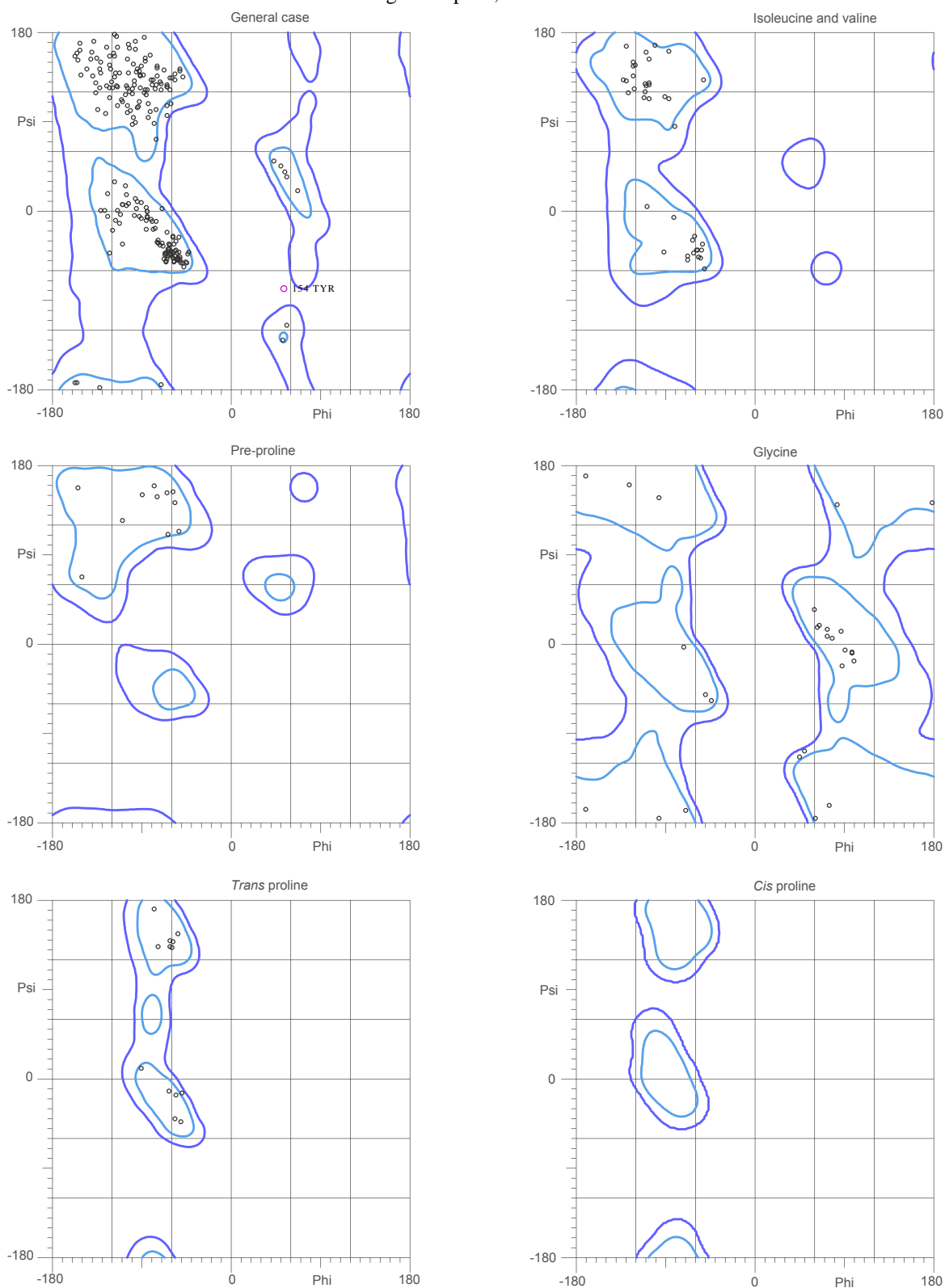


MolProbity Ramachandran analysis

5rgm.H.pdb, model 1



96.5% (300/311) of all residues were in favored (98%) regions.

99.7% (310/311) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

154 TYR (53.2, -78.6)