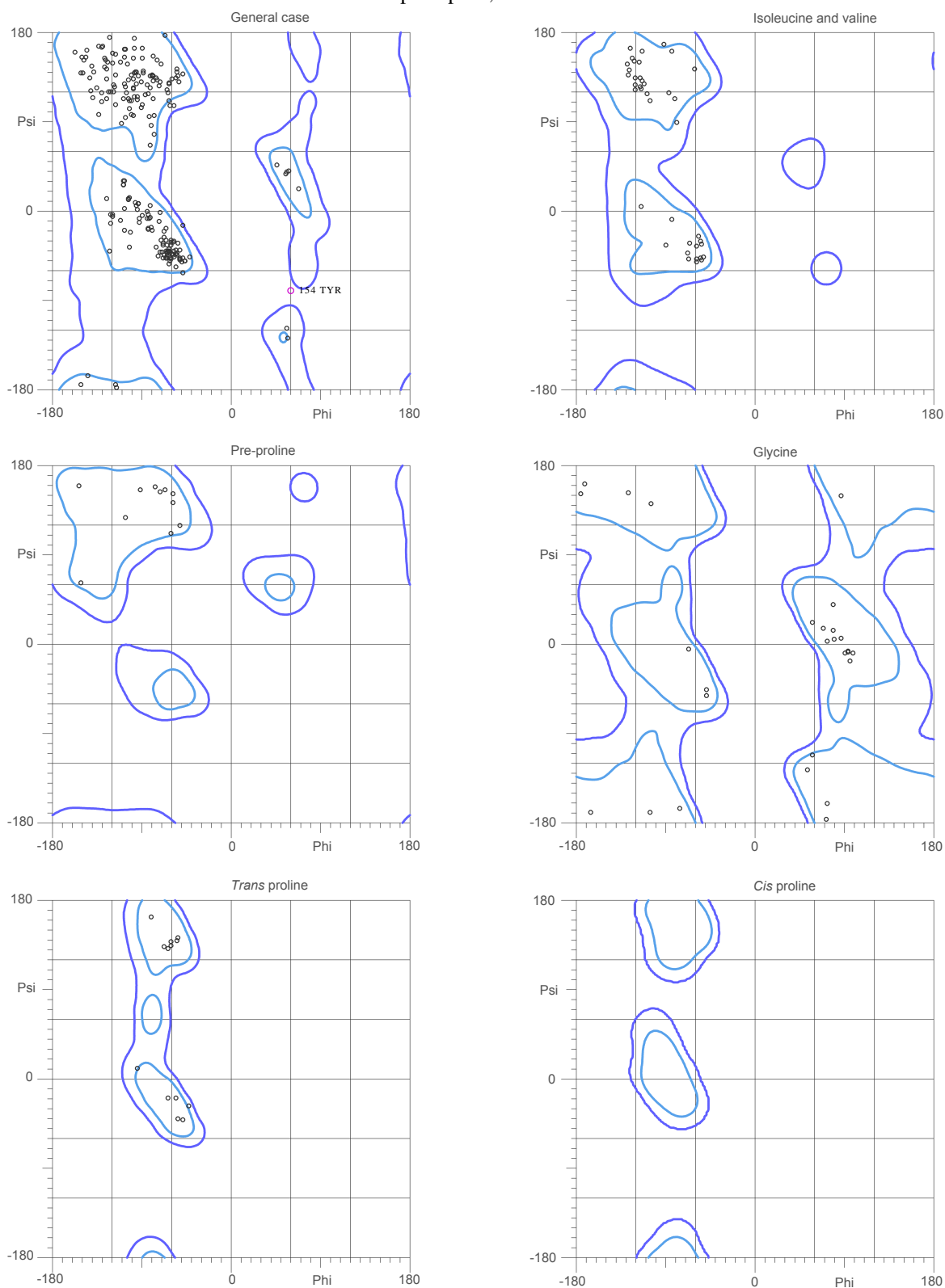


MolProbity Ramachandran analysis

5rep.H.pdb, model 1



96.1% (299/311) 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 (60.9, -80.6)