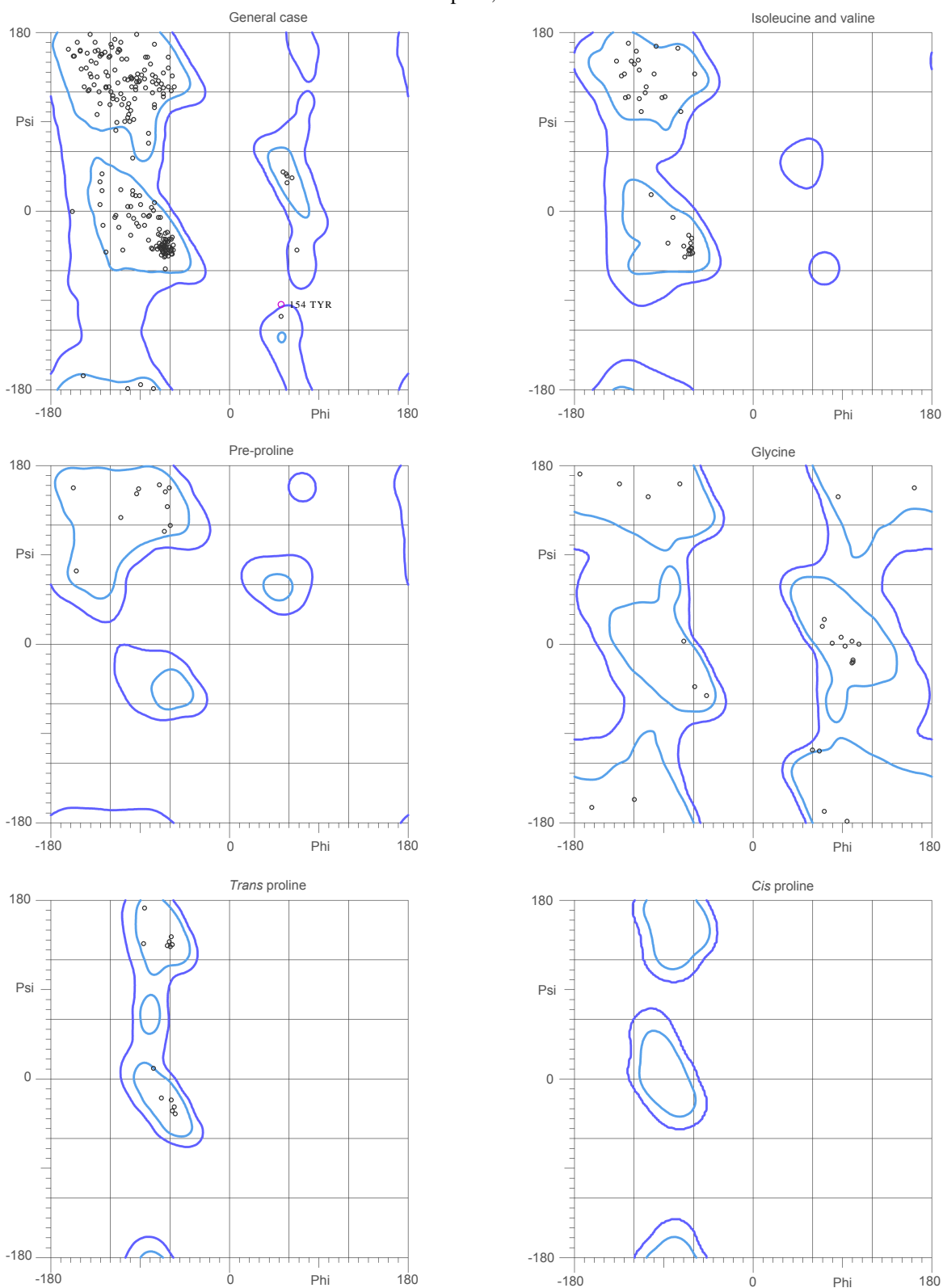


MolProbity Ramachandran analysis

7d1o.H.pdb, model 1



97.0% (290/299) of all residues were in favored (98%) regions.

99.7% (298/299) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

154 TYR (52.2, -94.0)