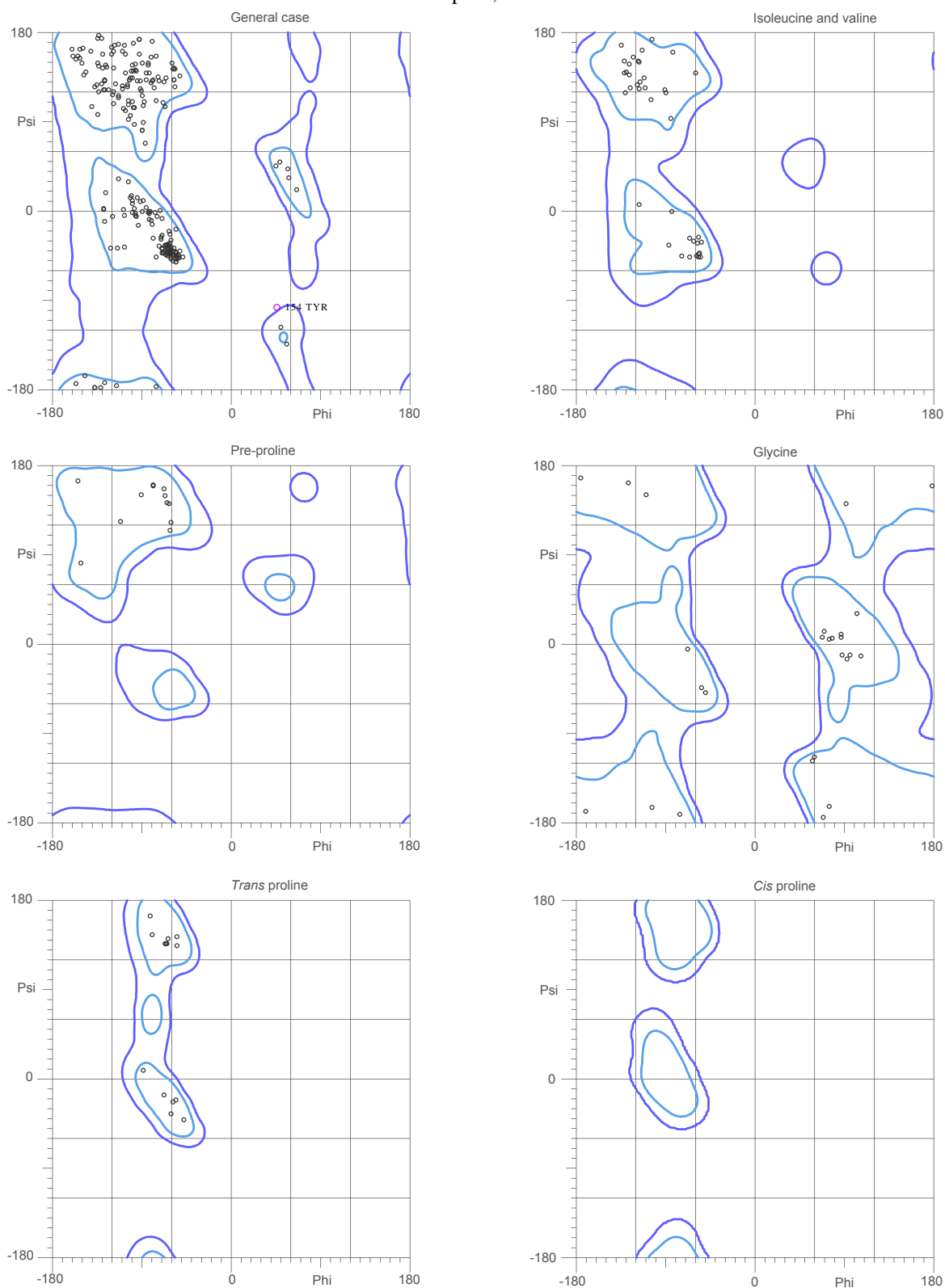


MolProbity Ramachandran analysis

7a1u.H.pdb, model 1



98.7% (313/317) of all residues were in favored (98%) regions.

99.7% (316/317) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

154 TYR (46.2, -97.7)