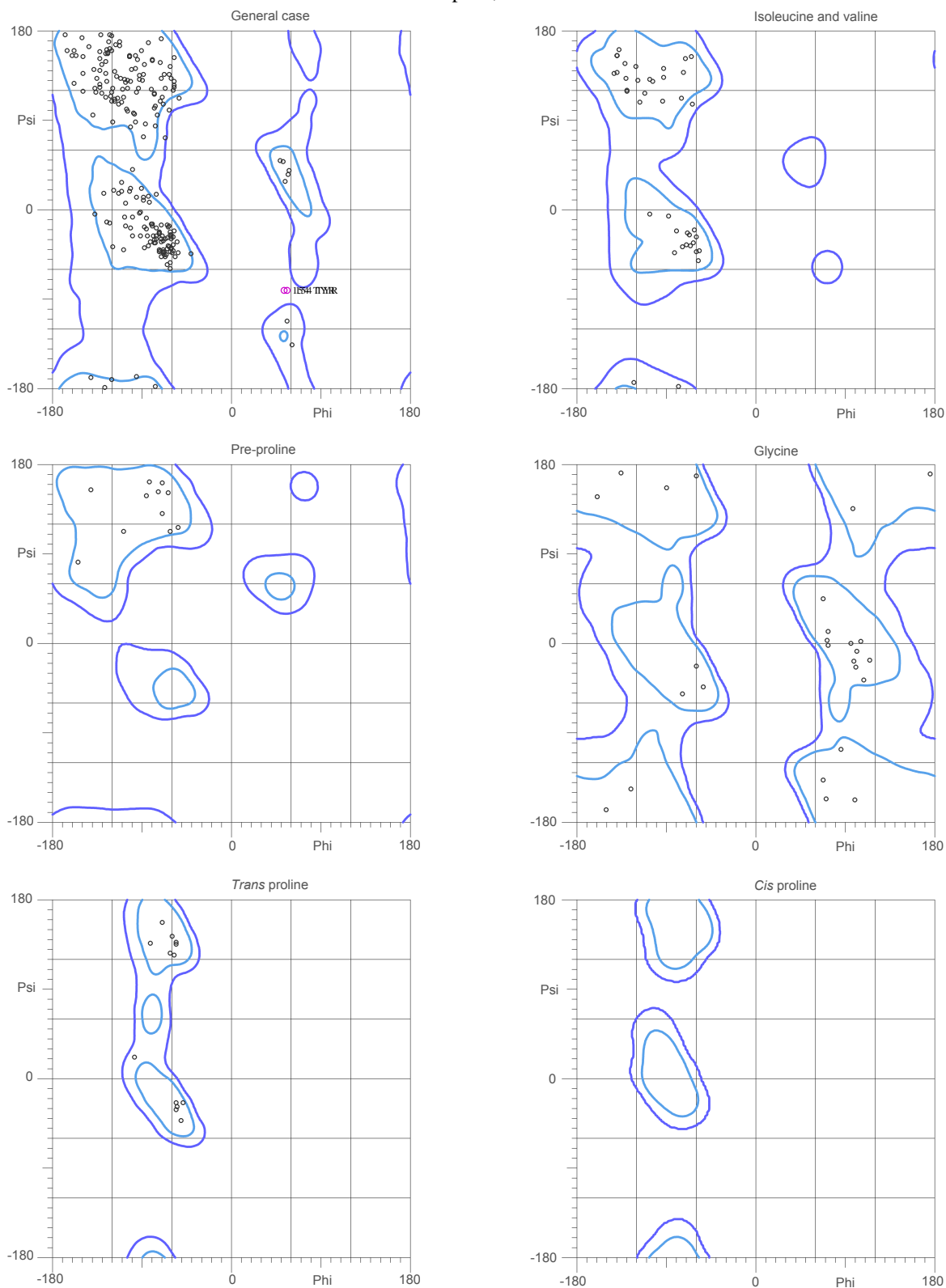


MolProbity Ramachandran analysis

7aku.H.pdb, model 1



95.1% (294/309) of all residues were in favored (98%) regions.

99.4% (307/309) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (Φ , Ψ):

154 TYR (56.3, -81.2)

154 TYR (53.6, -81.2)