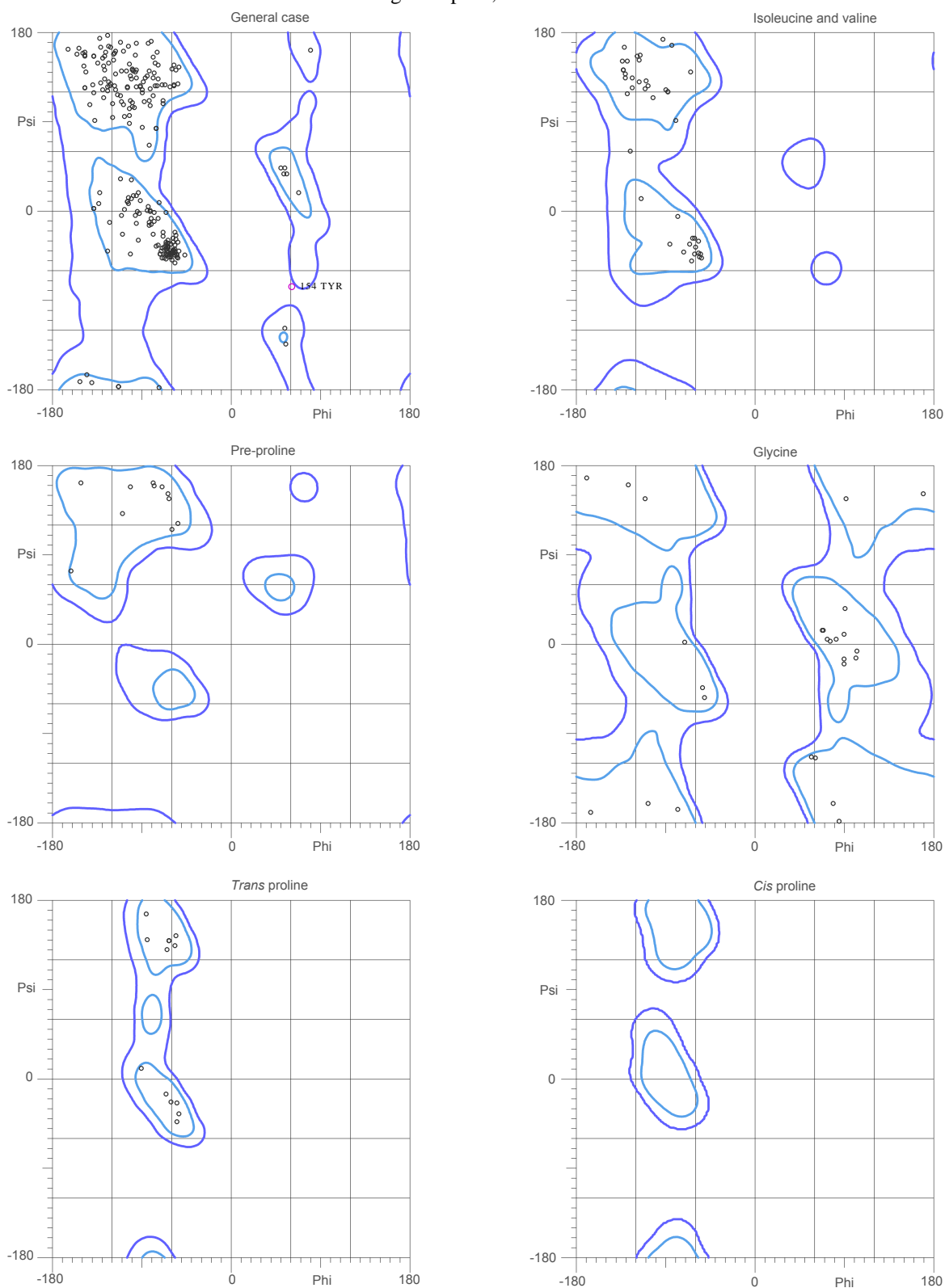


# MolProbity Ramachandran analysis

7aga.H.pdb, model 1



97.3% (319/328) of all residues were in favored (98%) regions.

99.7% (327/328) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (61.2, -76.7)