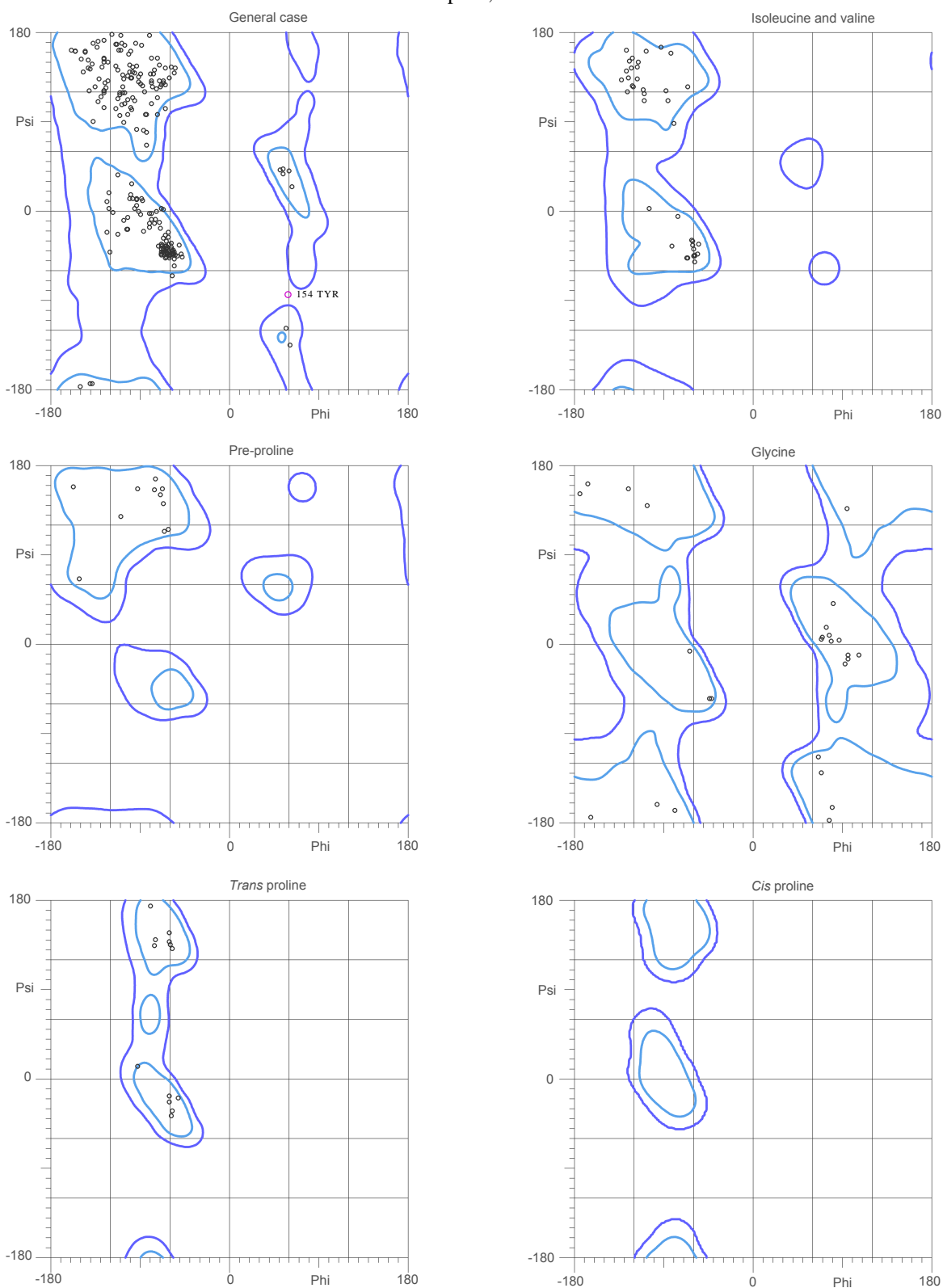


# MolProbity Ramachandran analysis

5rh7.H.pdb, model 1



97.4% (297/305) of all residues were in favored (98%) regions.

99.7% (304/305) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (59.8, -84.8)