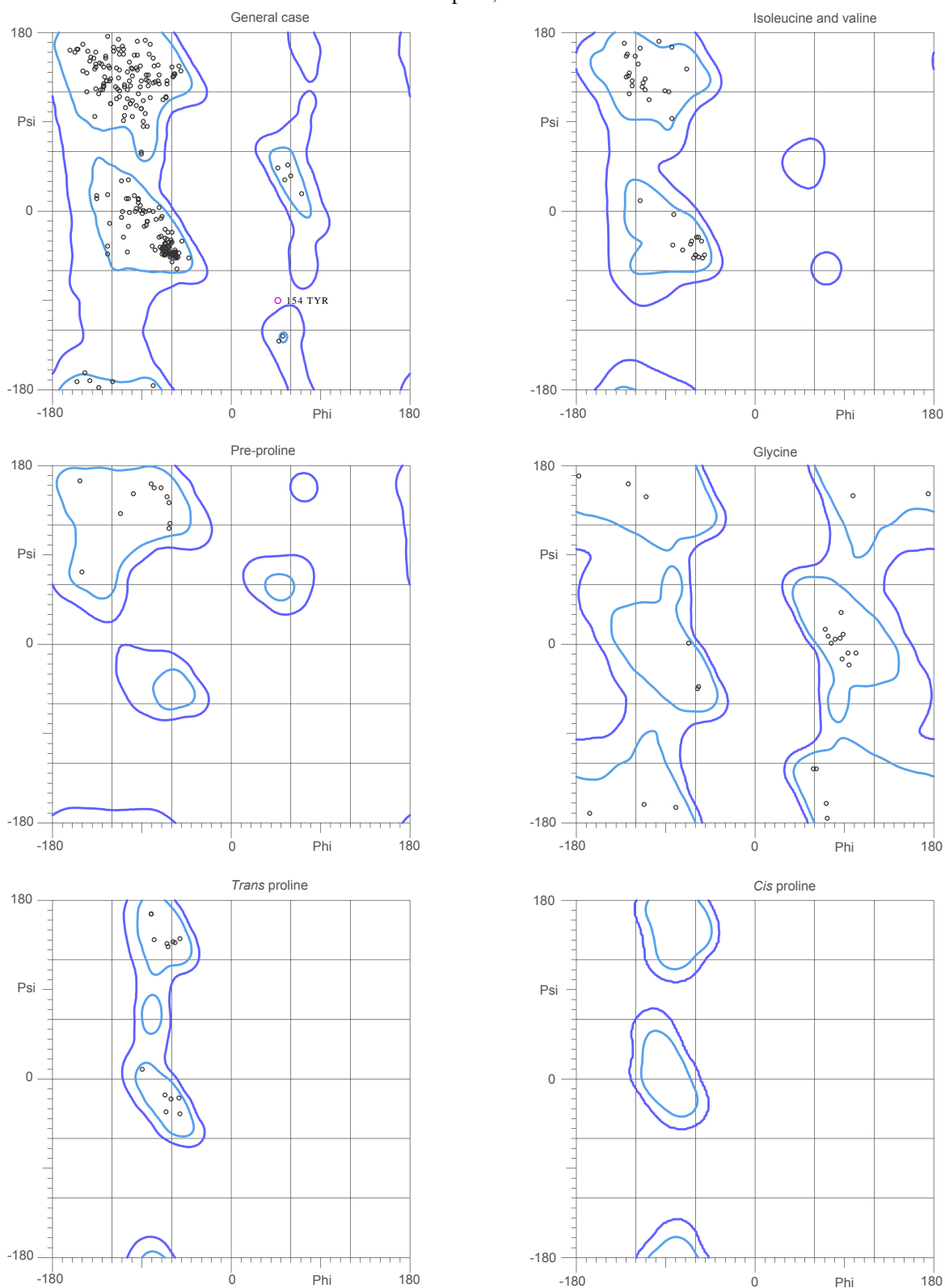


# MolProbity Ramachandran analysis

7abu.H.pdb, model 1



98.5% (321/326) of all residues were in favored (98%) regions.

99.7% (325/326) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (47.3, -90.3)