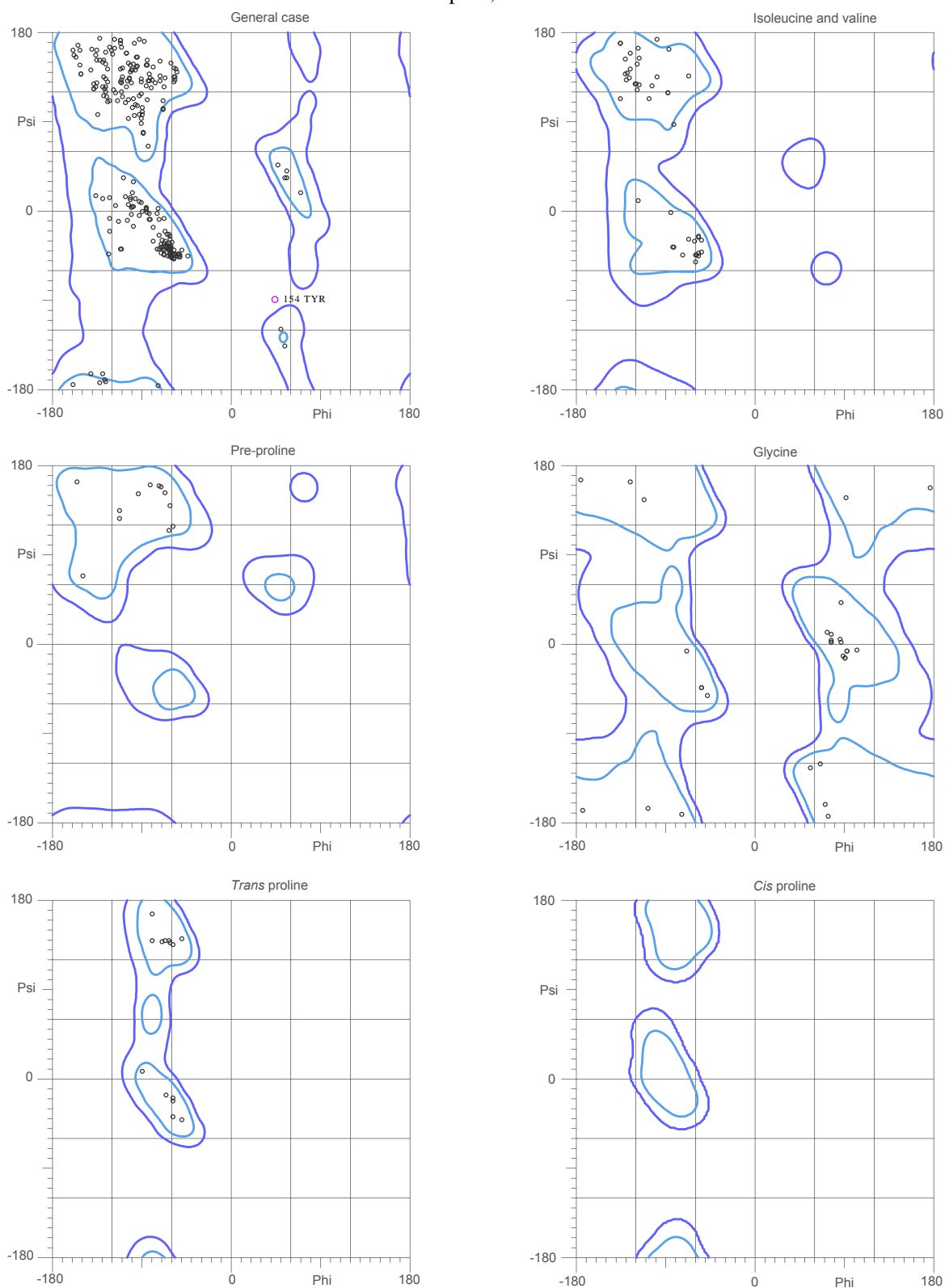


# MolProbity Ramachandran analysis

7ans.H.pdb, model 1



97.9% (332/339) of all residues were in favored (98%) regions.

99.7% (338/339) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (44.3, -89.4)