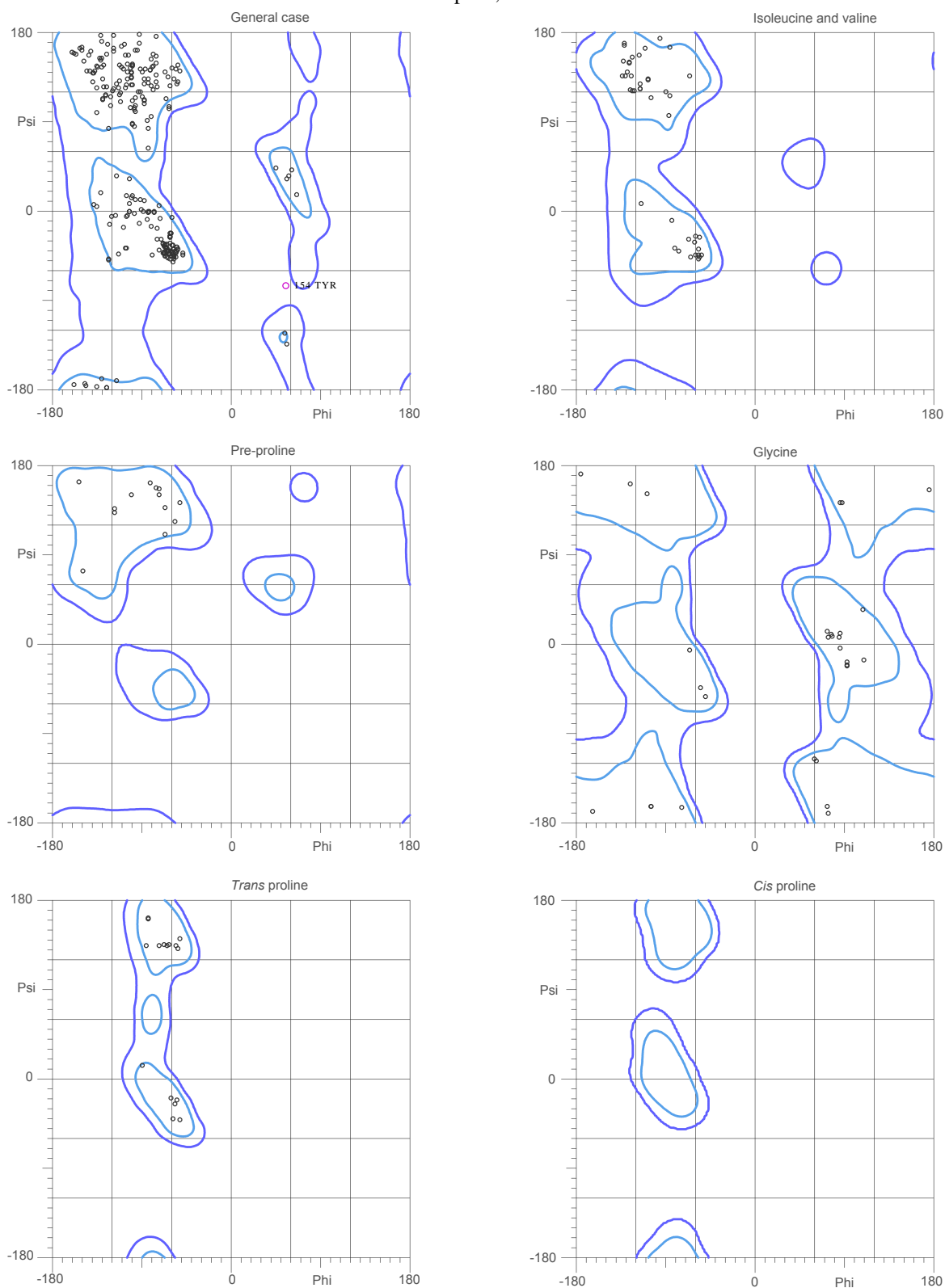


# MolProbity Ramachandran analysis

7adw.H.pdb, model 1



97.7% (340/348) of all residues were in favored (98%) regions.

99.7% (347/348) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (55.7, -75.4)