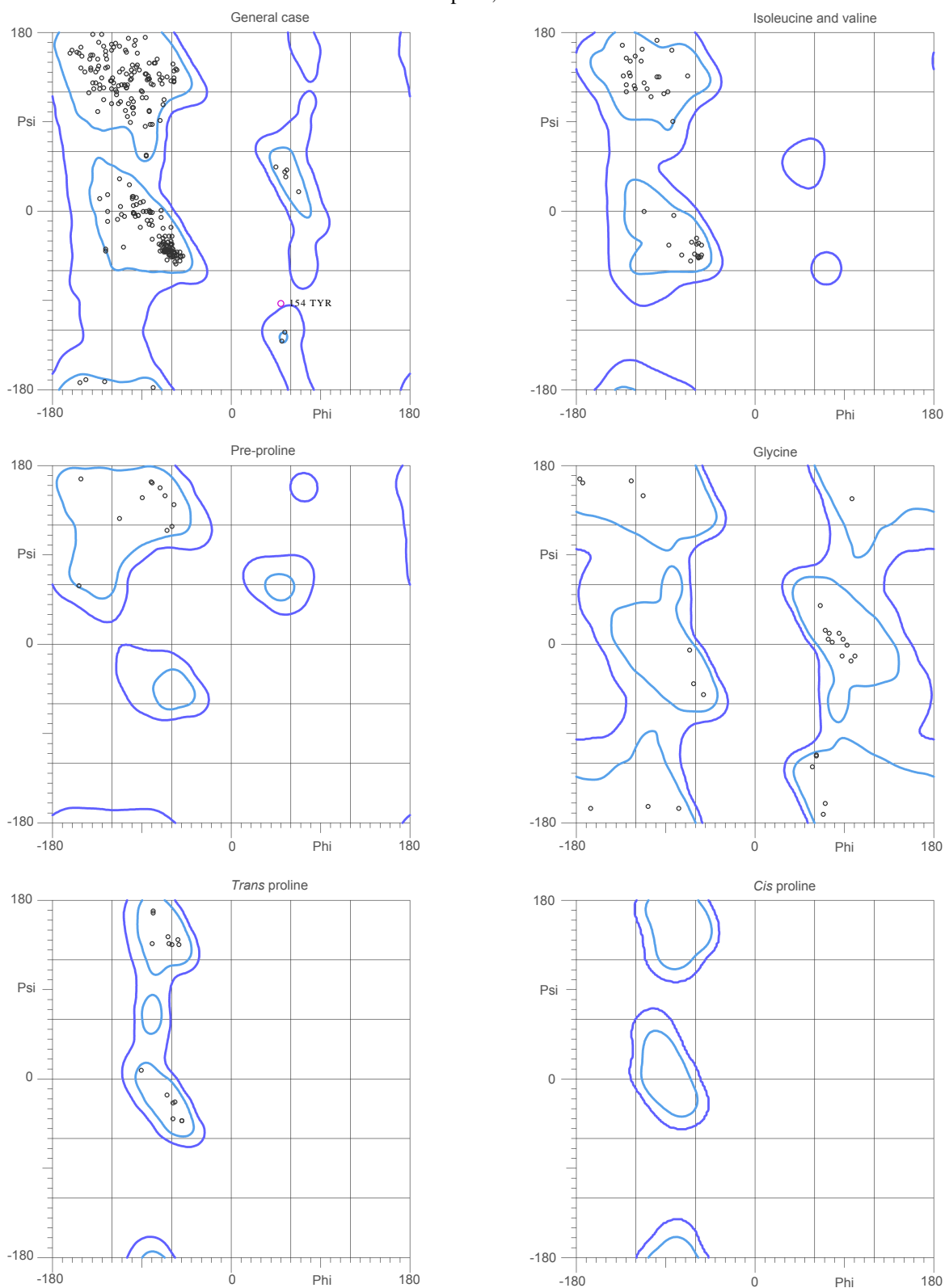


# MolProbity Ramachandran analysis

7awr.H.pdb, model 1



98.2% (336/342) of all residues were in favored (98%) regions.

99.7% (341/342) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (50.9, -93.4)