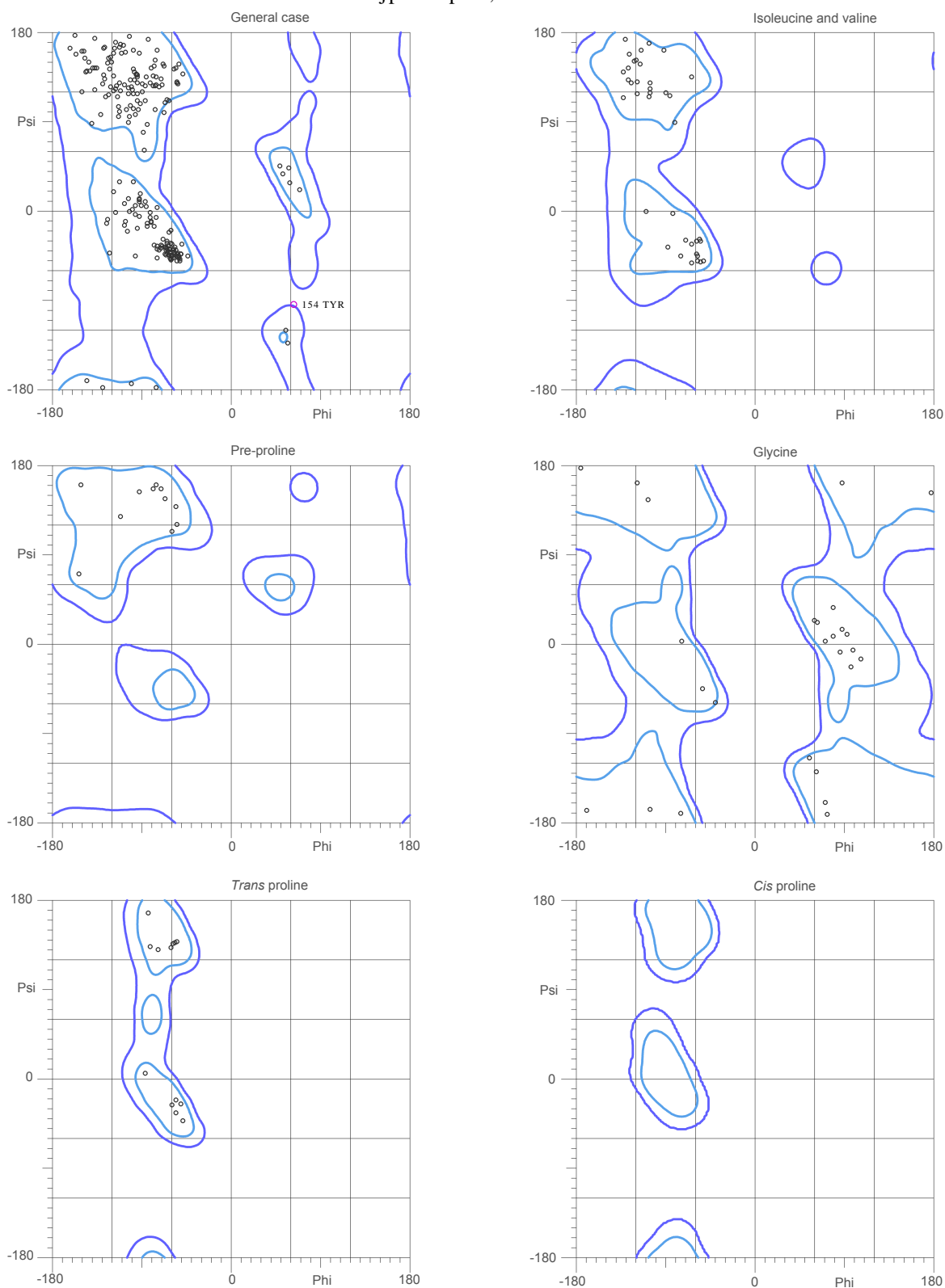


# MolProbity Ramachandran analysis

7jp1.H.pdb, model 1



98.7% (301/305) 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 (63.3, -94.9)