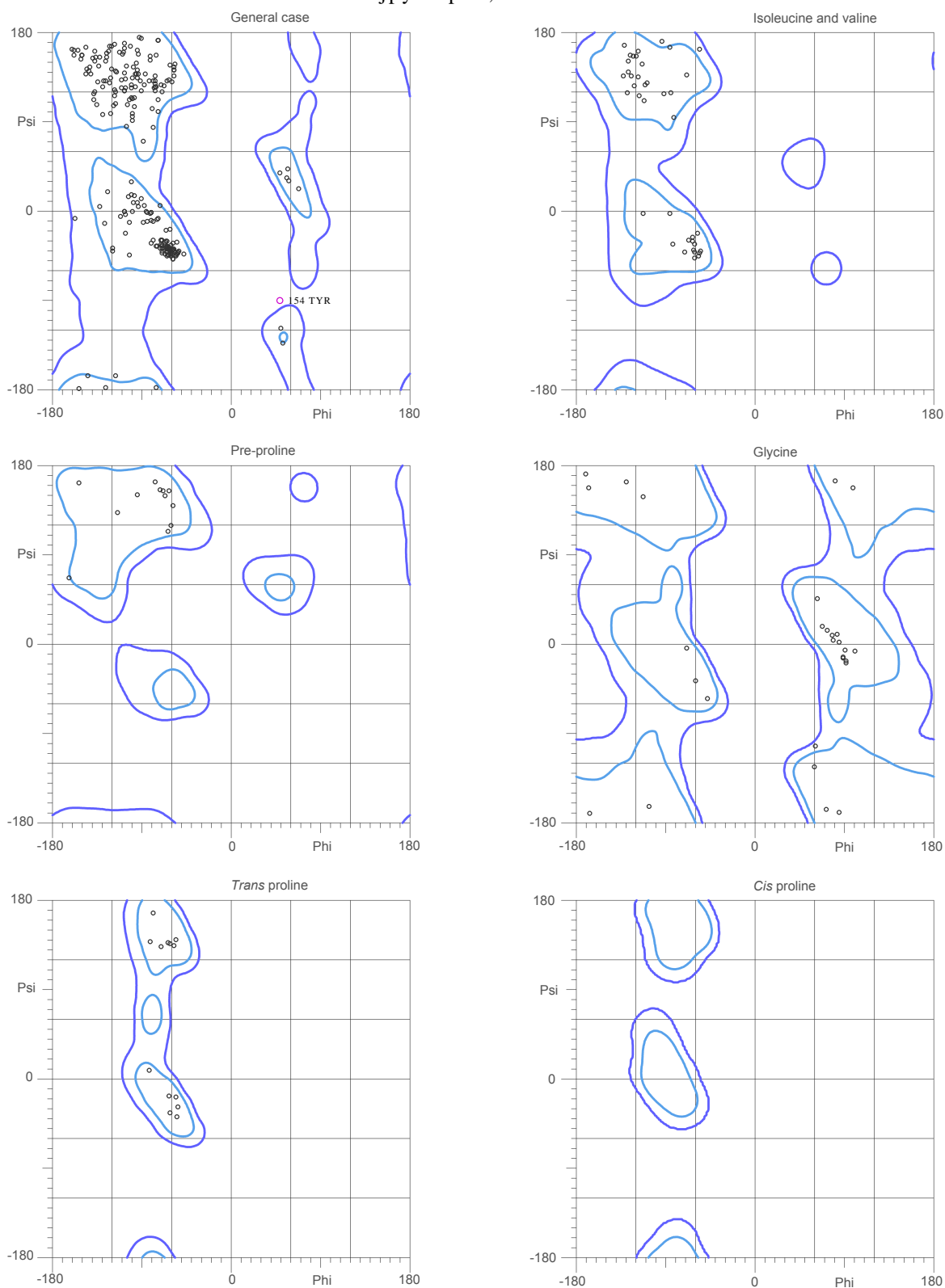


# MolProbity Ramachandran analysis

7jpy.H.pdb, model 1



96.9% (310/320) of all residues were in favored (98%) regions.

99.7% (319/320) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (49.3, -90.3)