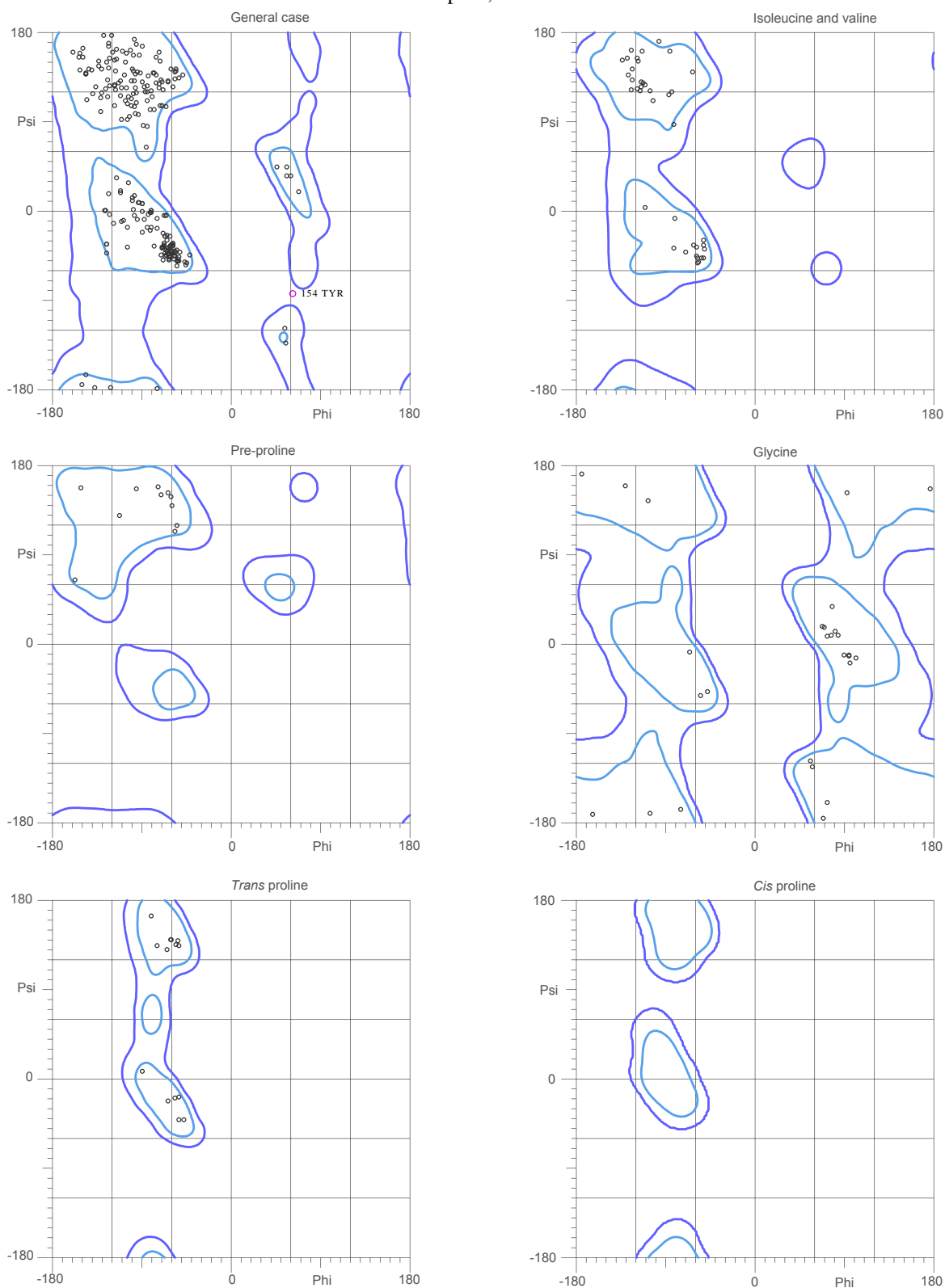


# MolProbity Ramachandran analysis

5rfz.H.pdb, model 1



98.1% (316/322) of all residues were in favored (98%) regions.

99.7% (321/322) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (63.0, -83.2)