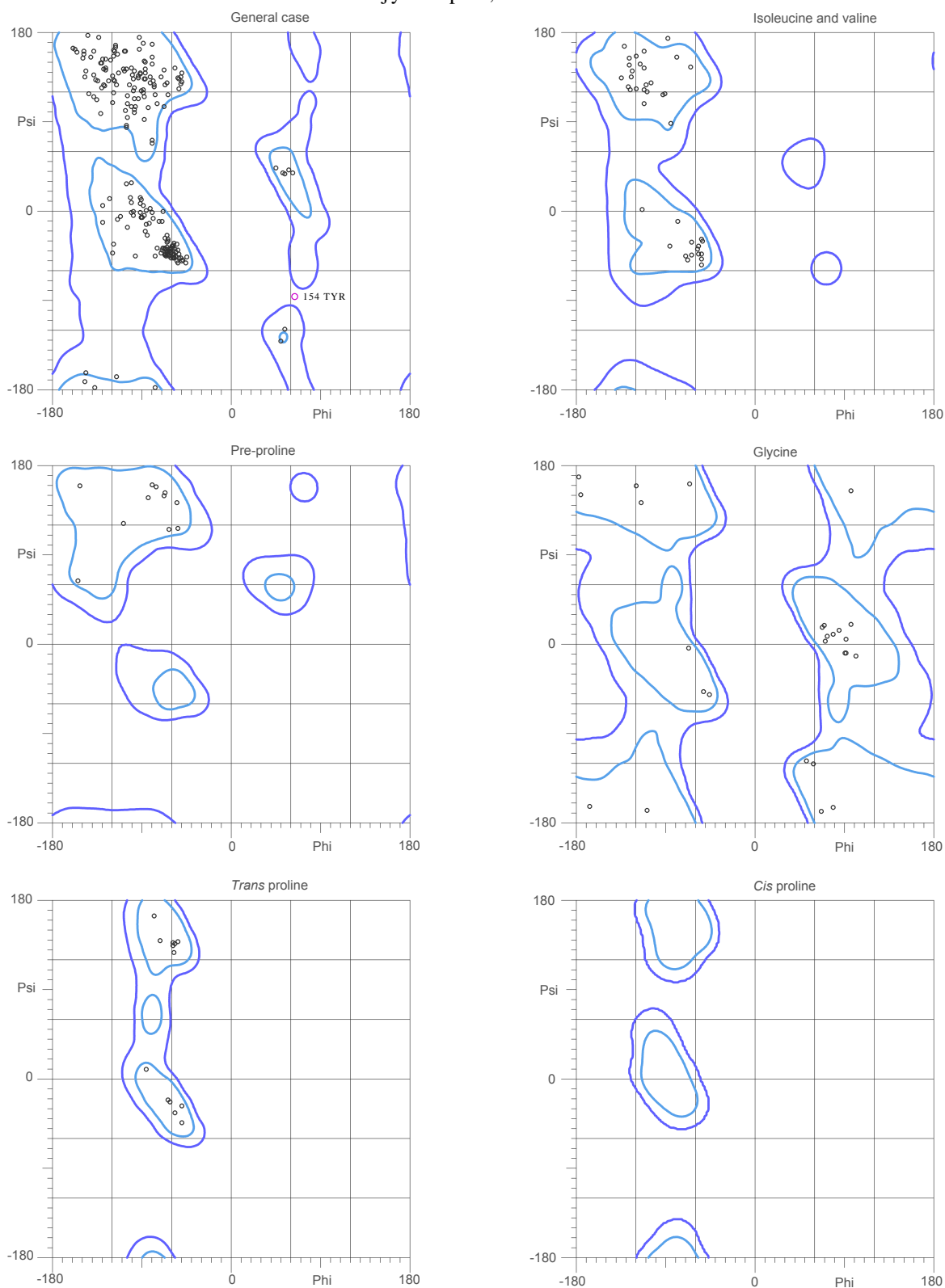


# MolProbity Ramachandran analysis

7jyc.H.pdb, model 1



98.1% (302/308) of all residues were in favored (98%) regions.

99.7% (307/308) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (64.1, -86.2)