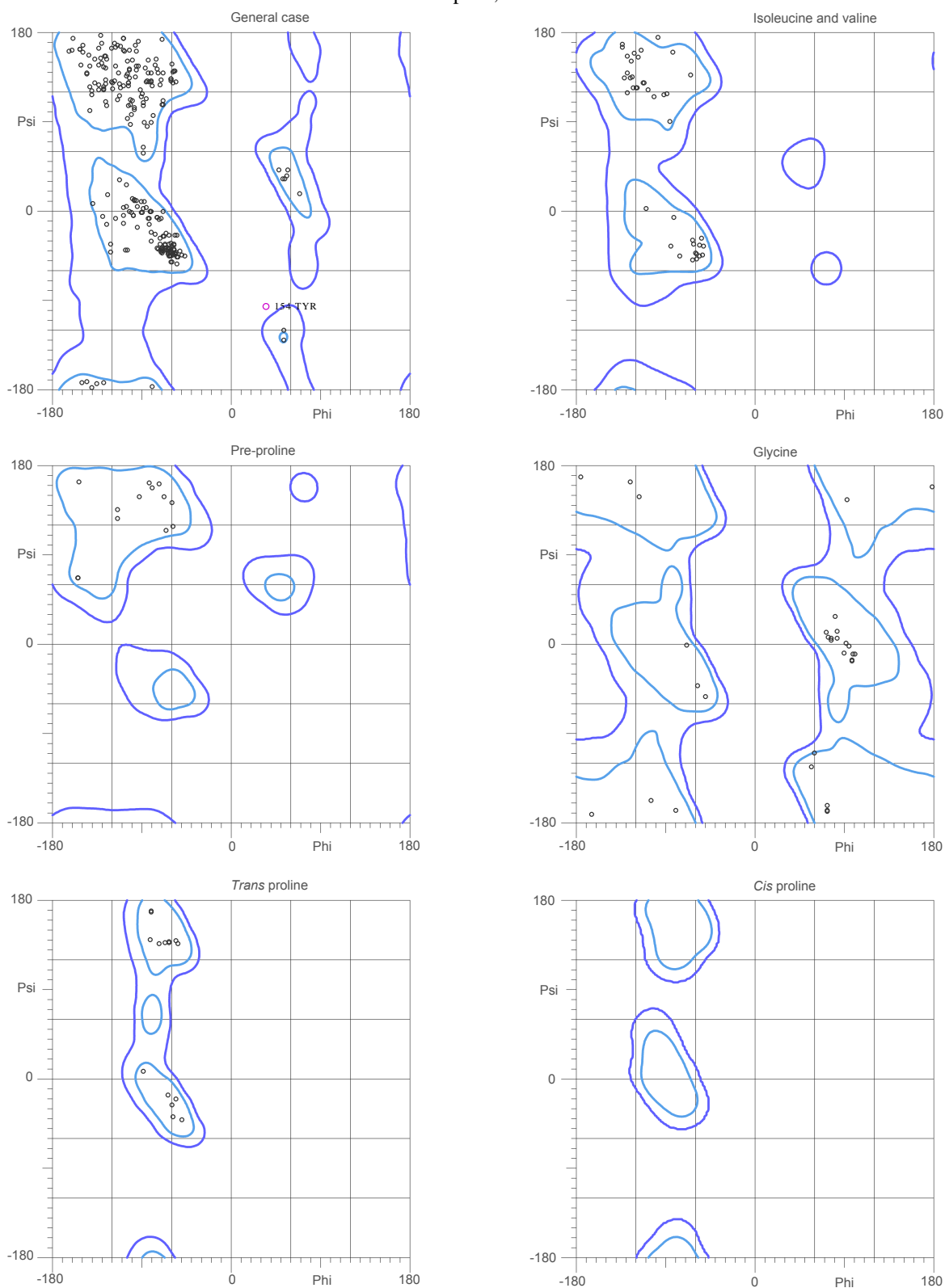


# MolProbity Ramachandran analysis

7axm.H.pdb, model 1



99.1% (340/343) of all residues were in favored (98%) regions.

99.7% (342/343) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (35.5, -96.5)