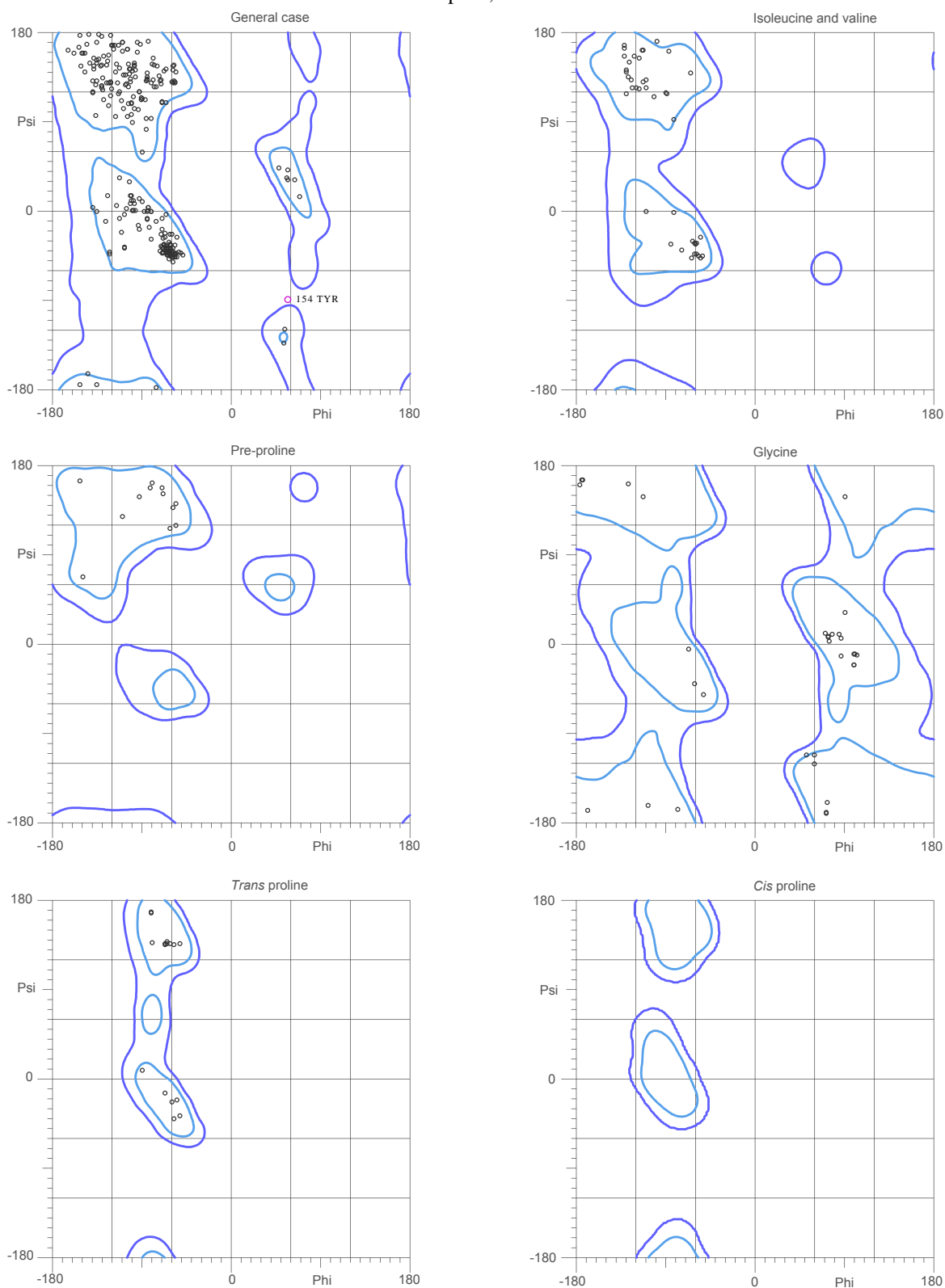


# MolProbity Ramachandran analysis

7ar6.H.pdb, model 1



98.6% (352/357) of all residues were in favored (98%) regions.

99.7% (356/357) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (57.8, -89.5)