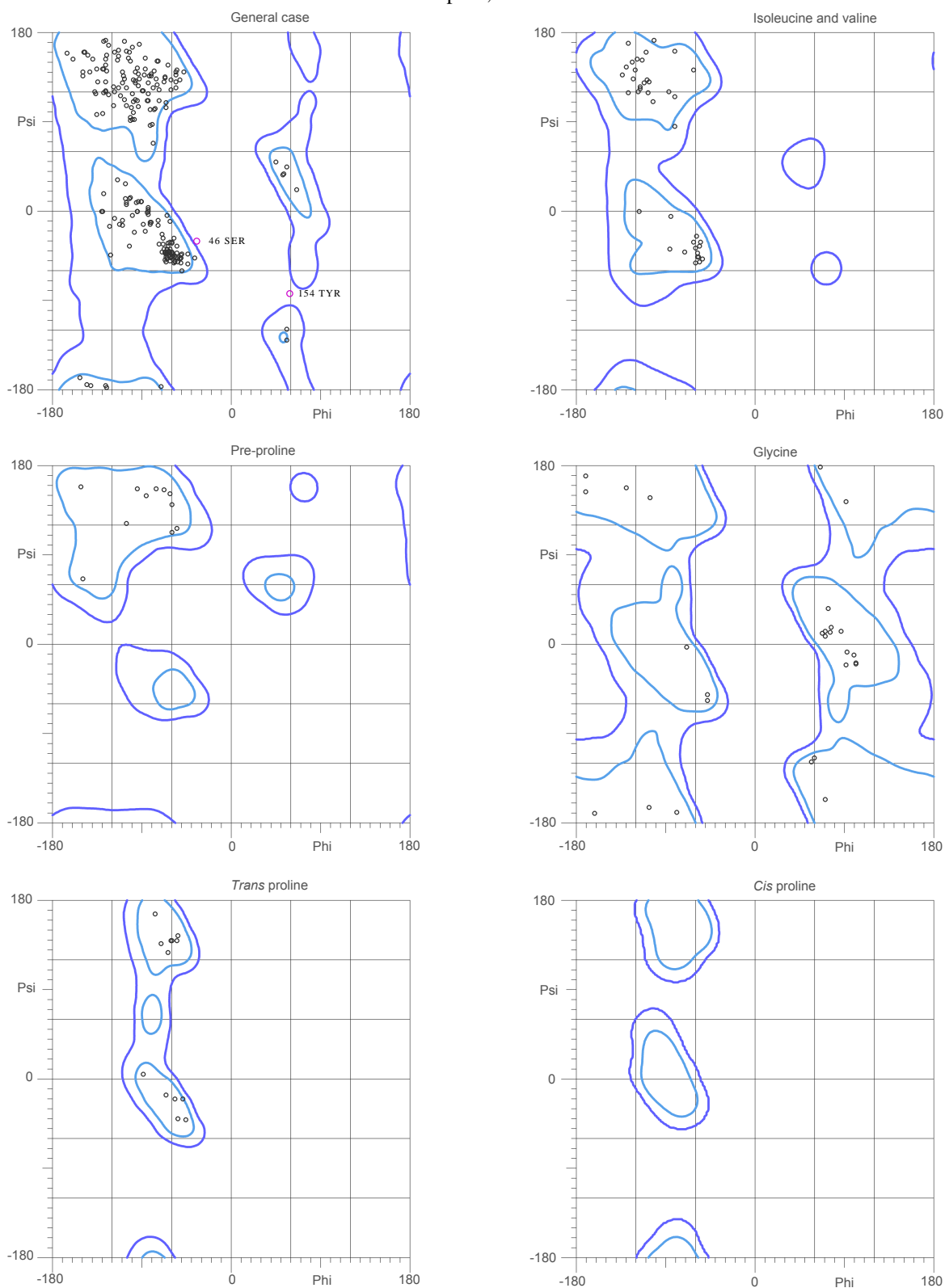


# MolProbity Ramachandran analysis

5rff.H.pdb, model 1



97.4% (303/311) of all residues were in favored (98%) regions.

99.4% (309/311) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

46 SER (-35.2, -30.7)

154 TYR (59.6, -83.8)