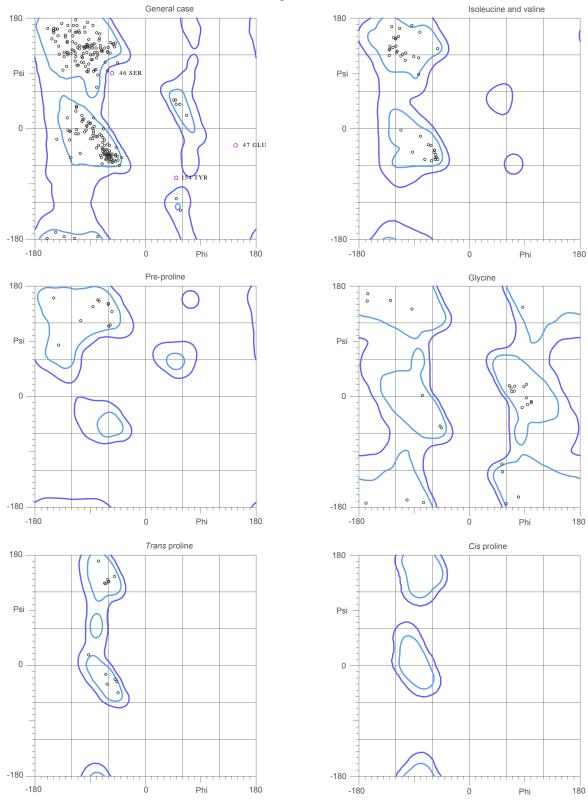# MolProbity Ramachandran analysis

5r80.H.pdb, model 1

95.8% (298/311) of all residues were in favored (98%) regions.
99.0% (308/311) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi):
    46 SER (-55.3, 91.4)
    47 GLU (147.3, -27.7)
    154 TYR (50.8, -80.5)