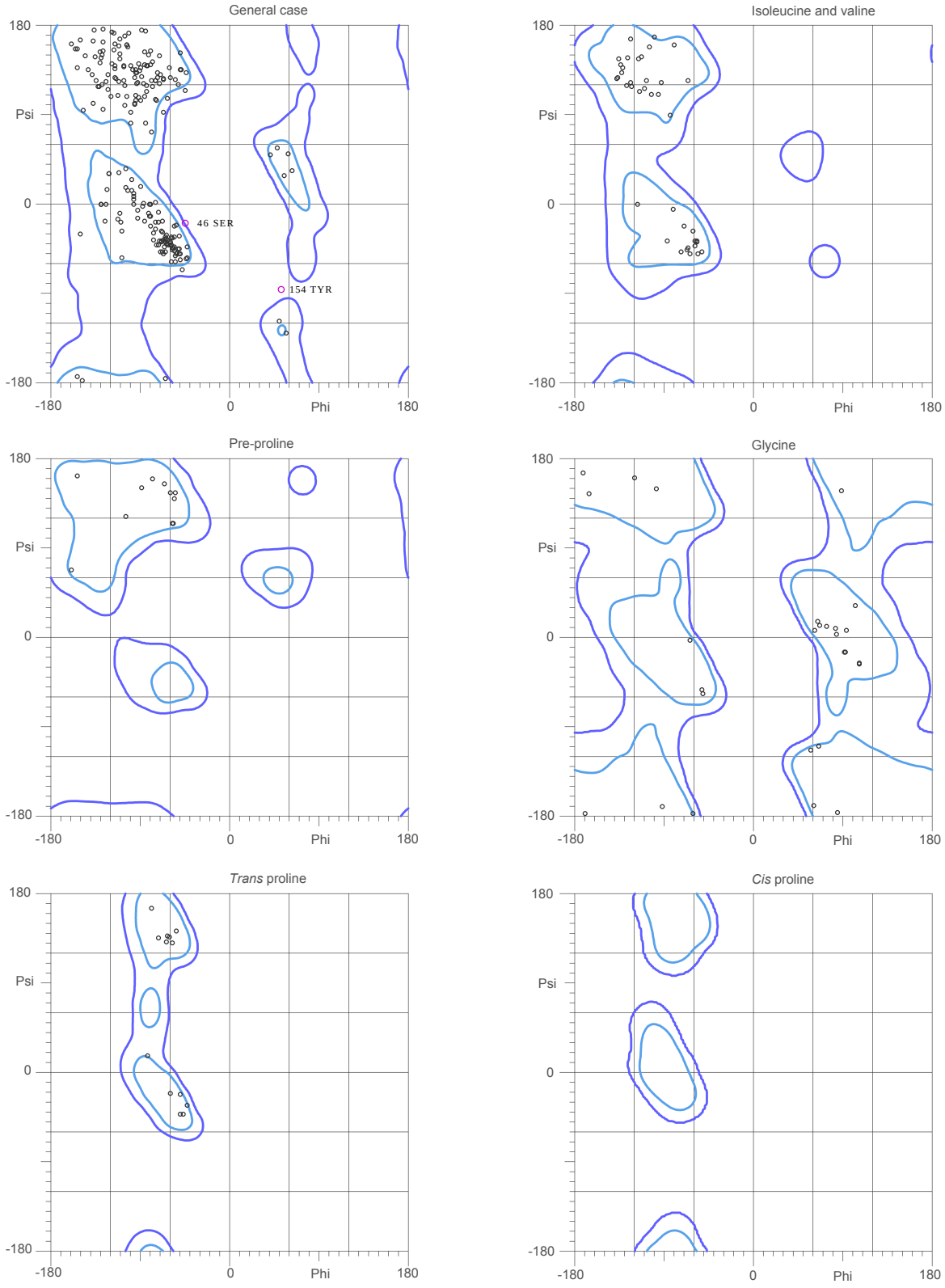
MolProbity Ramachandran analysis

5rfp.H.pdb, model 1



95.8% (298/311) 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 (-45.1, -19.7)
154 TYR (52.1, -86.4)