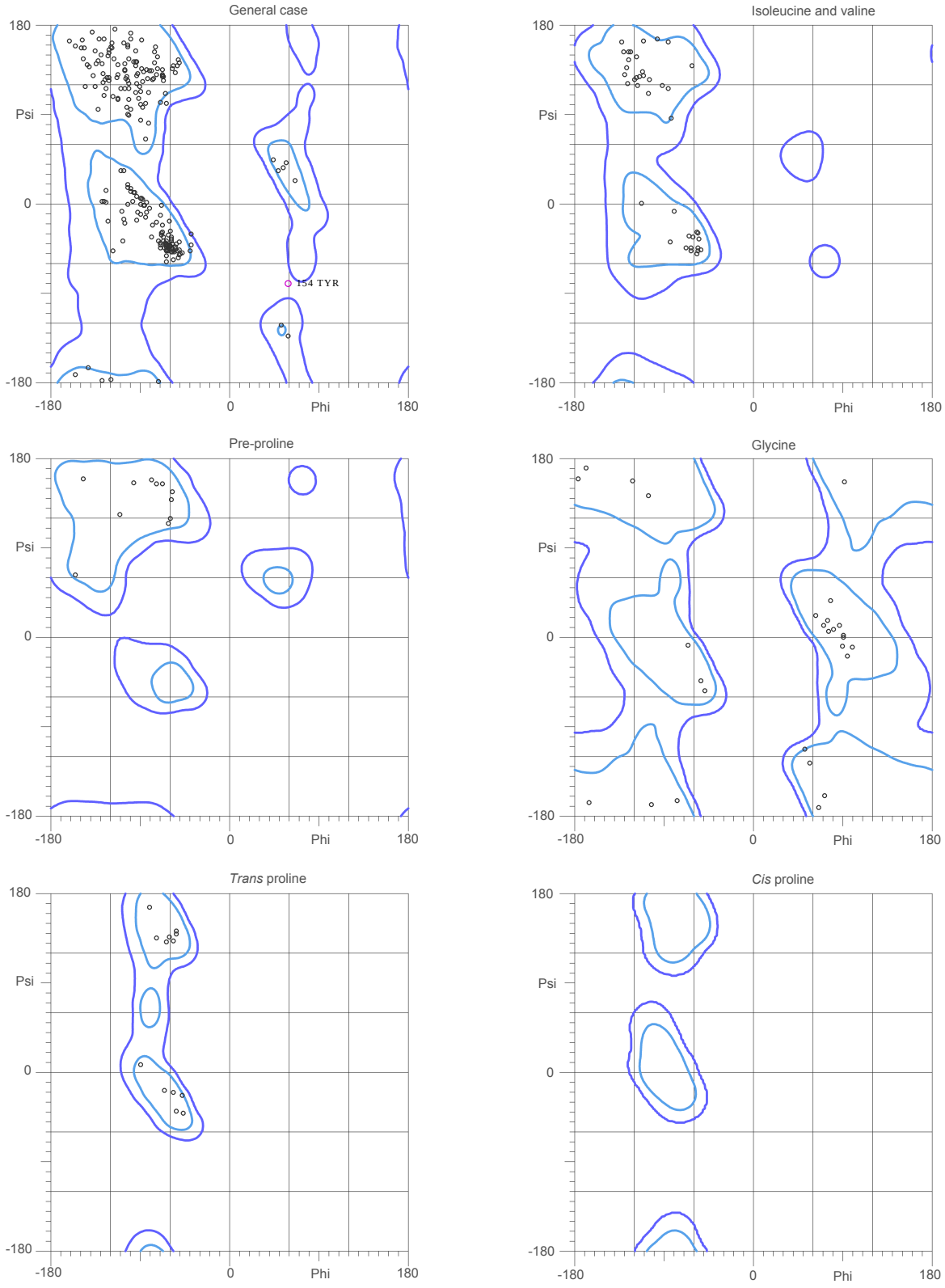
MolProbity Ramachandran analysis

5rfb.H.pdb, model 1



98.1% (305/311) of all residues were in favored (98%) regions.
99.7% (310/311) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):
154 TYR (59.6, -80.1)