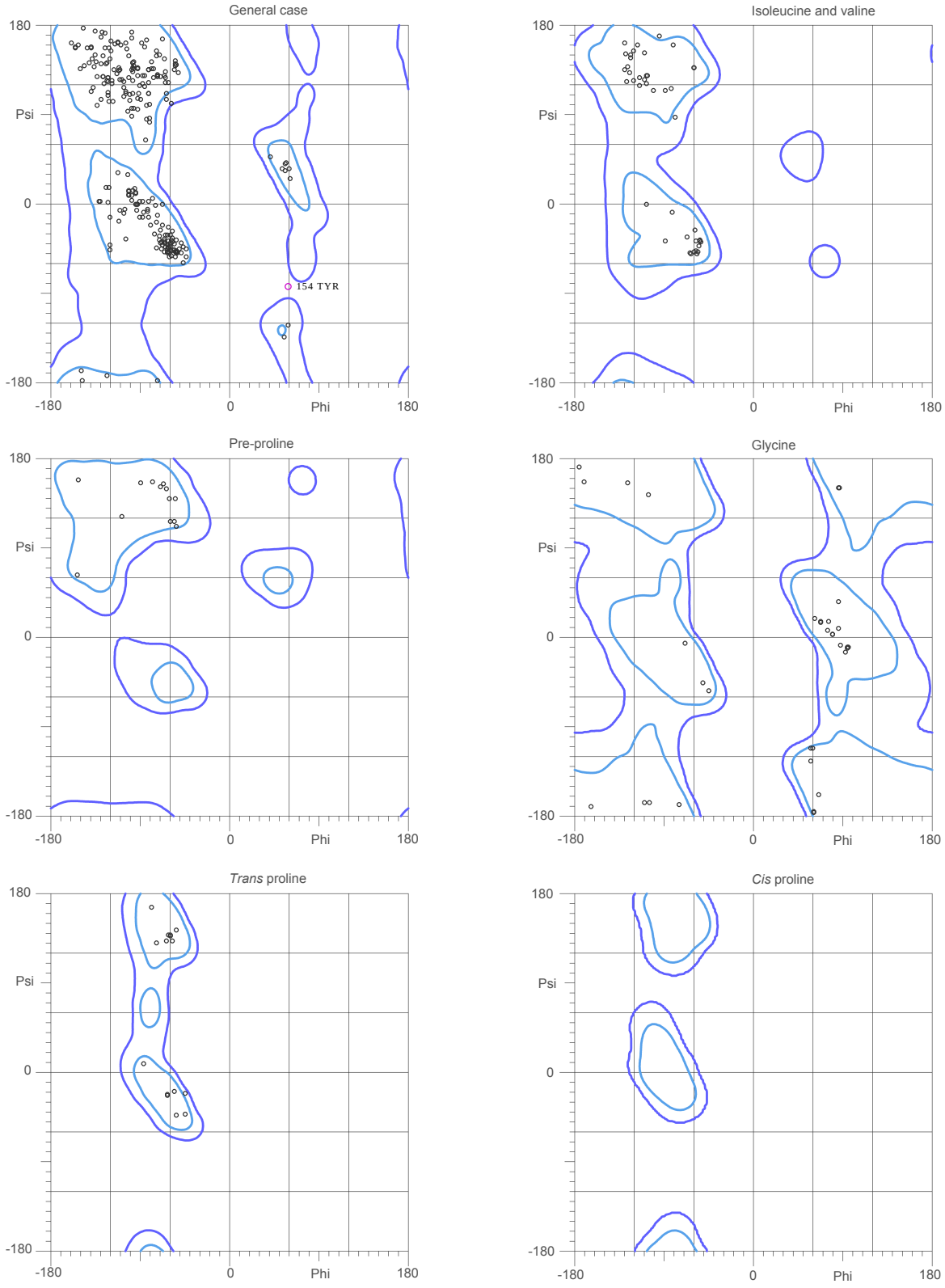
MolProbity Ramachandran analysis

5reg.H.pdb, model 1



97.8% (349/357) 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 (59.9, -83.4)