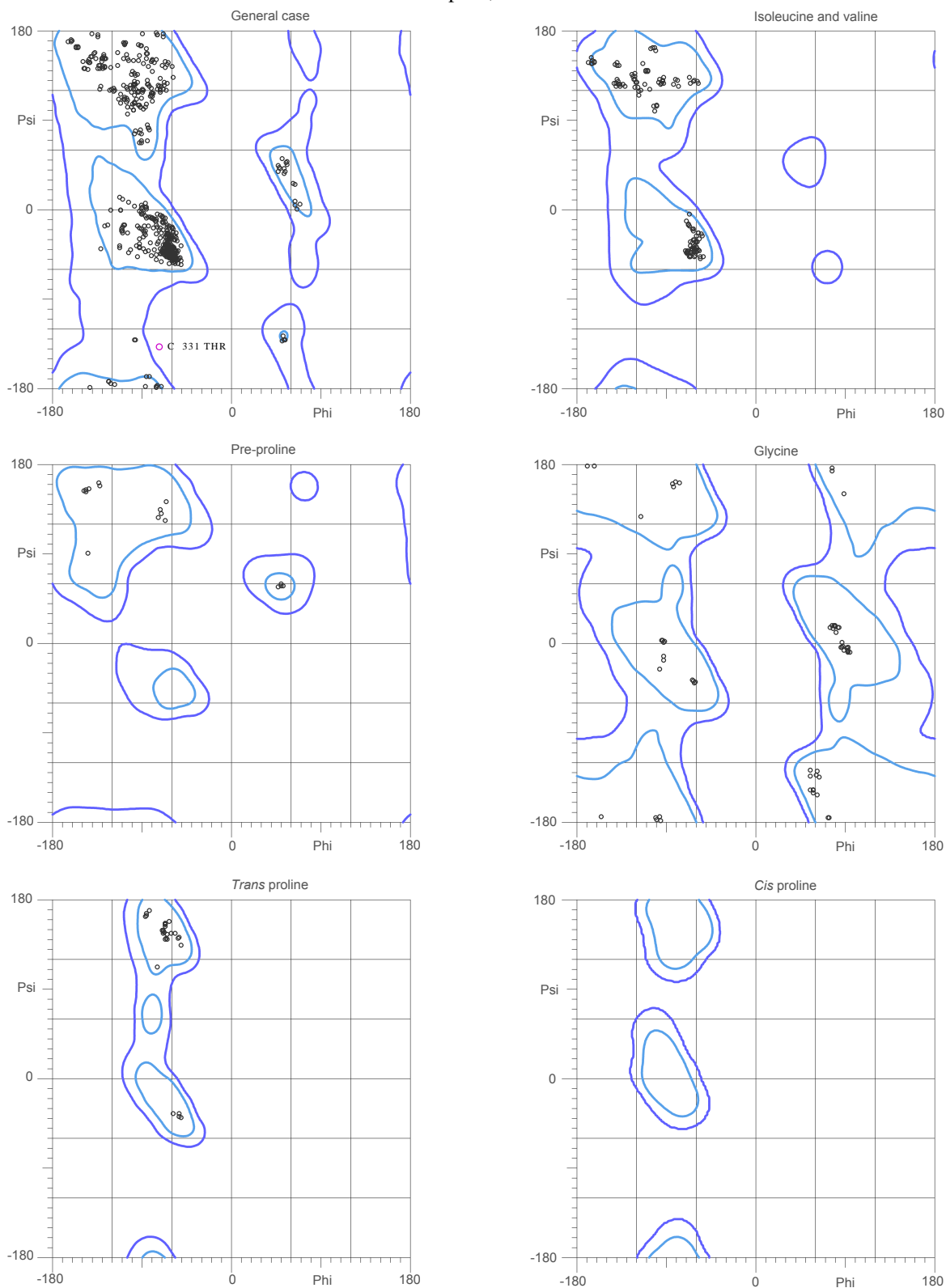


MolProbity Ramachandran analysis

2acf.H.pdb, model 1



98.8% (764/773) of all residues were in favored (98%) regions.

99.9% (772/773) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

C 331 THR (-74.0, -138.5)