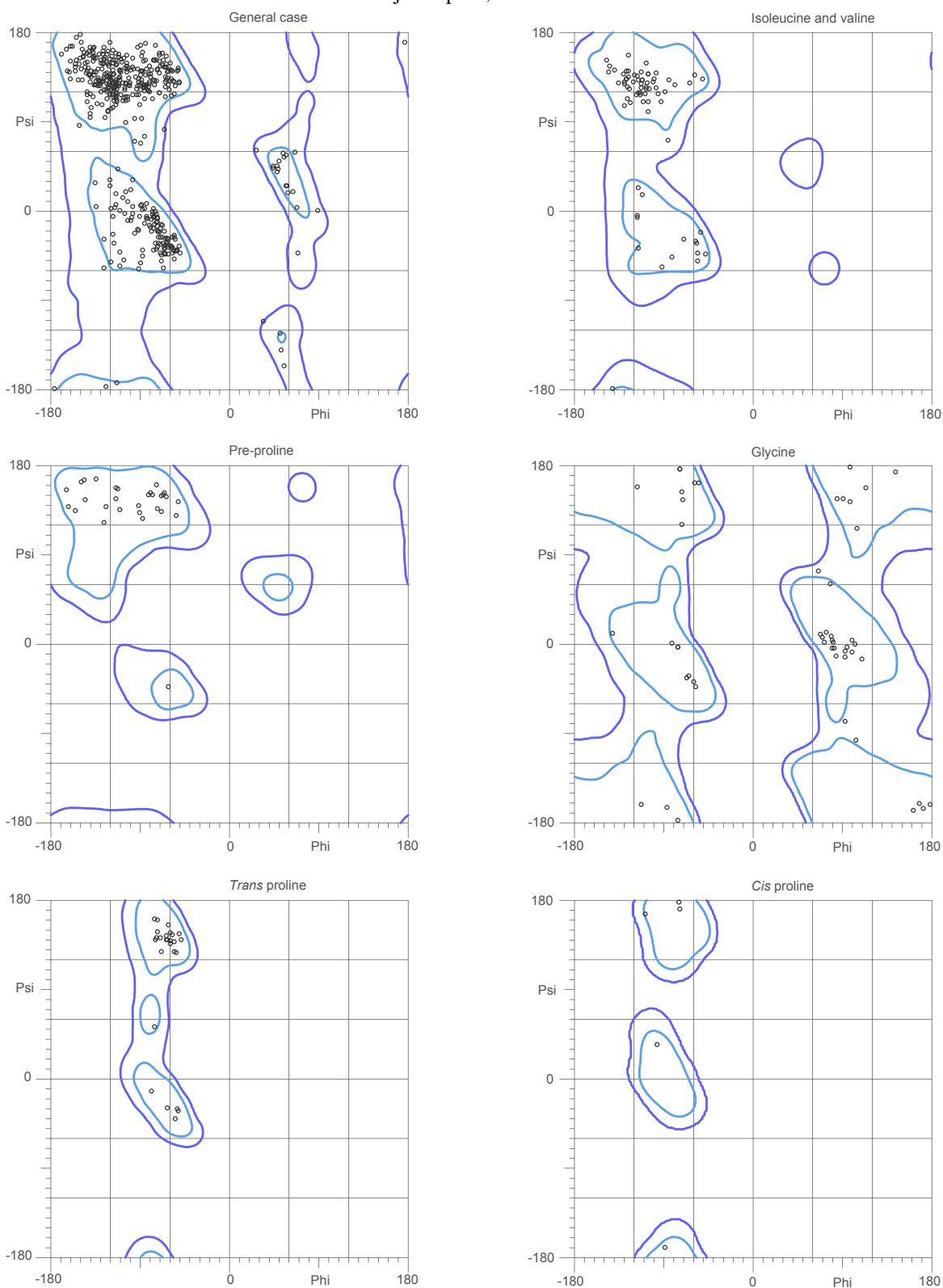


# MolProbity Ramachandran analysis

7cjf.H.pdb, model 1



97.2% (618/636) of all residues were in favored (98%) regions.

100.0% (636/636) of all residues were in allowed (>99.8%) regions.

There were no outliers.