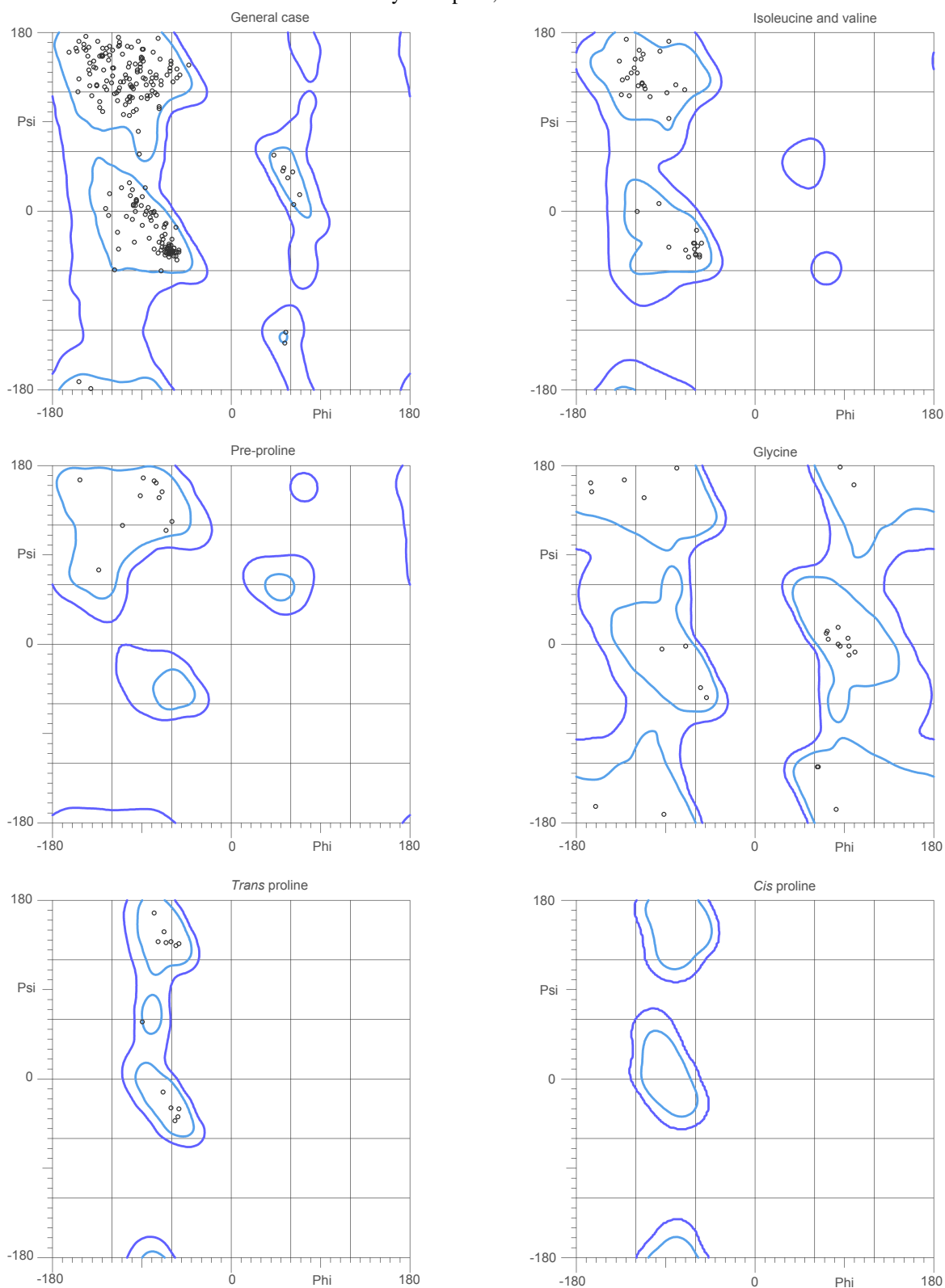


# MolProbity Ramachandran analysis

7kyu.H.pdb, model 1



97.8% (309/316) of all residues were in favored (98%) regions.

100.0% (316/316) of all residues were in allowed (>99.8%) regions.

There were no outliers.