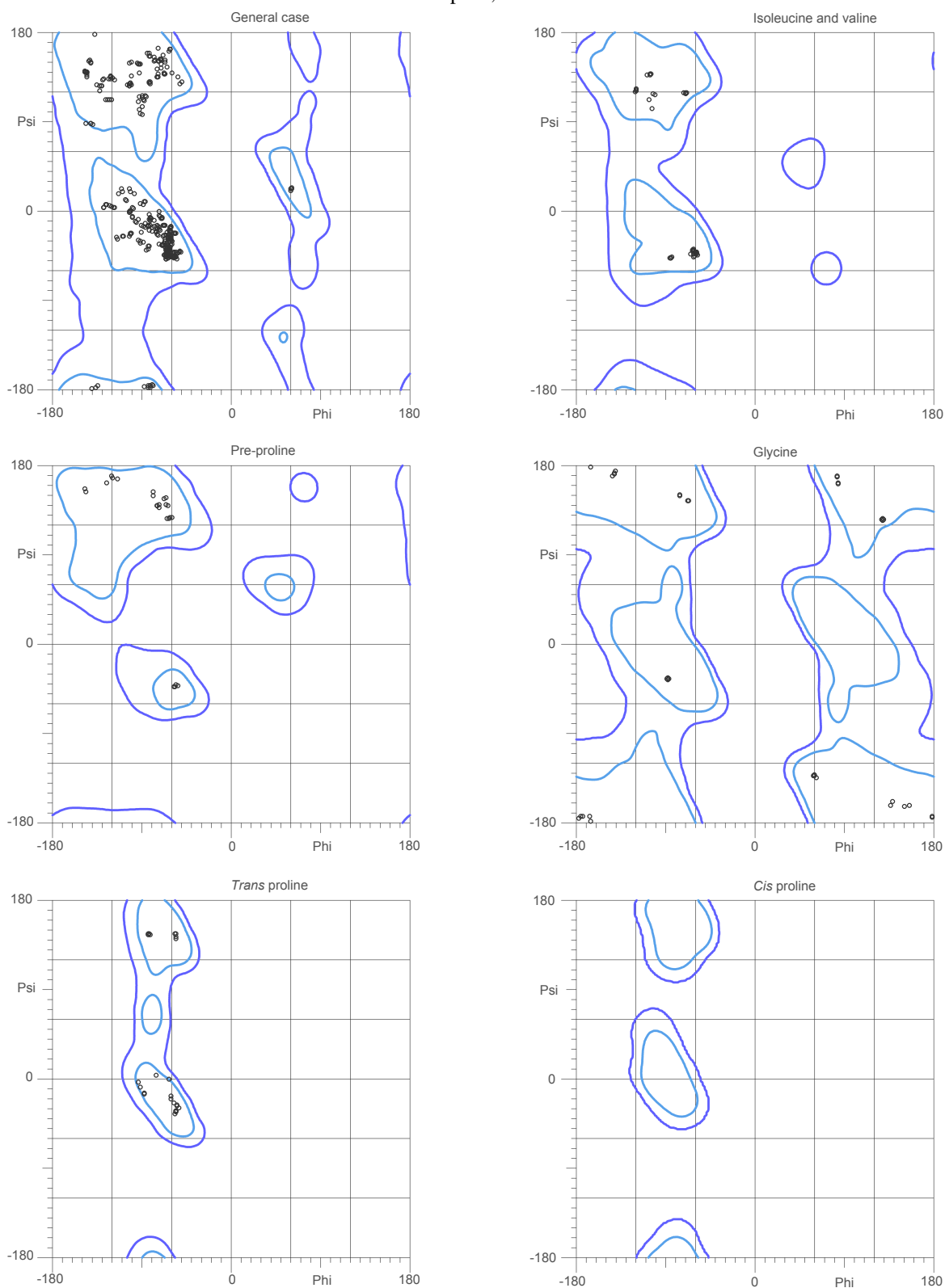


# MolProbity Ramachandran analysis

7ce0.H.pdb, model 1



99.8% (433/434) of all residues were in favored (98%) regions.

100.0% (434/434) of all residues were in allowed (>99.8%) regions.

There were no outliers.