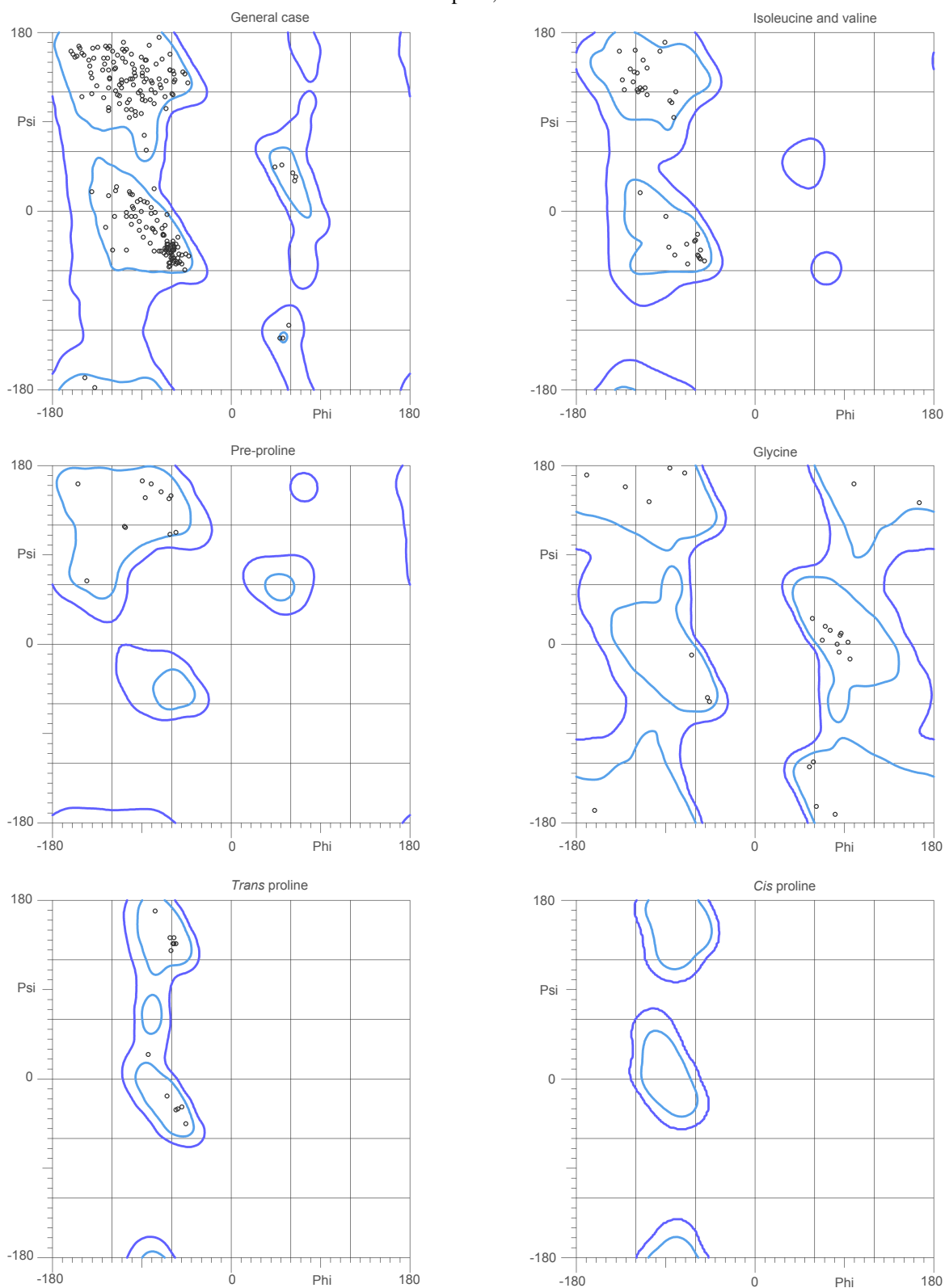


# MolProbity Ramachandran analysis

7c8b.H.pdb, model 1



98.3% (297/302) of all residues were in favored (98%) regions.

100.0% (302/302) of all residues were in allowed (>99.8%) regions.

There were no outliers.