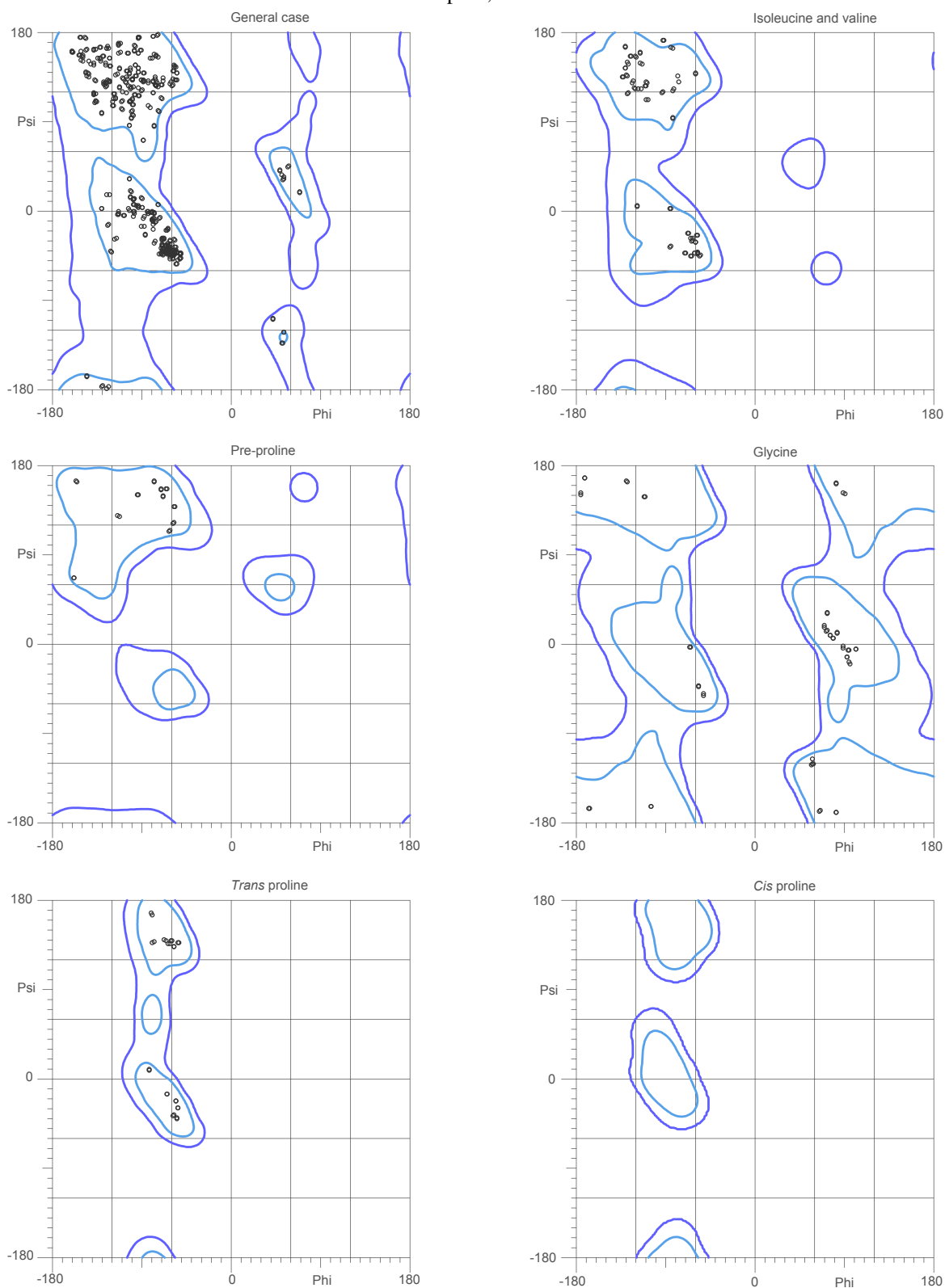


# MolProbity Ramachandran analysis

7ali.H.pdb, model 1



98.7% (607/615) of all residues were in favored (98%) regions.

100.0% (615/615) of all residues were in allowed (>99.8%) regions.

There were no outliers.