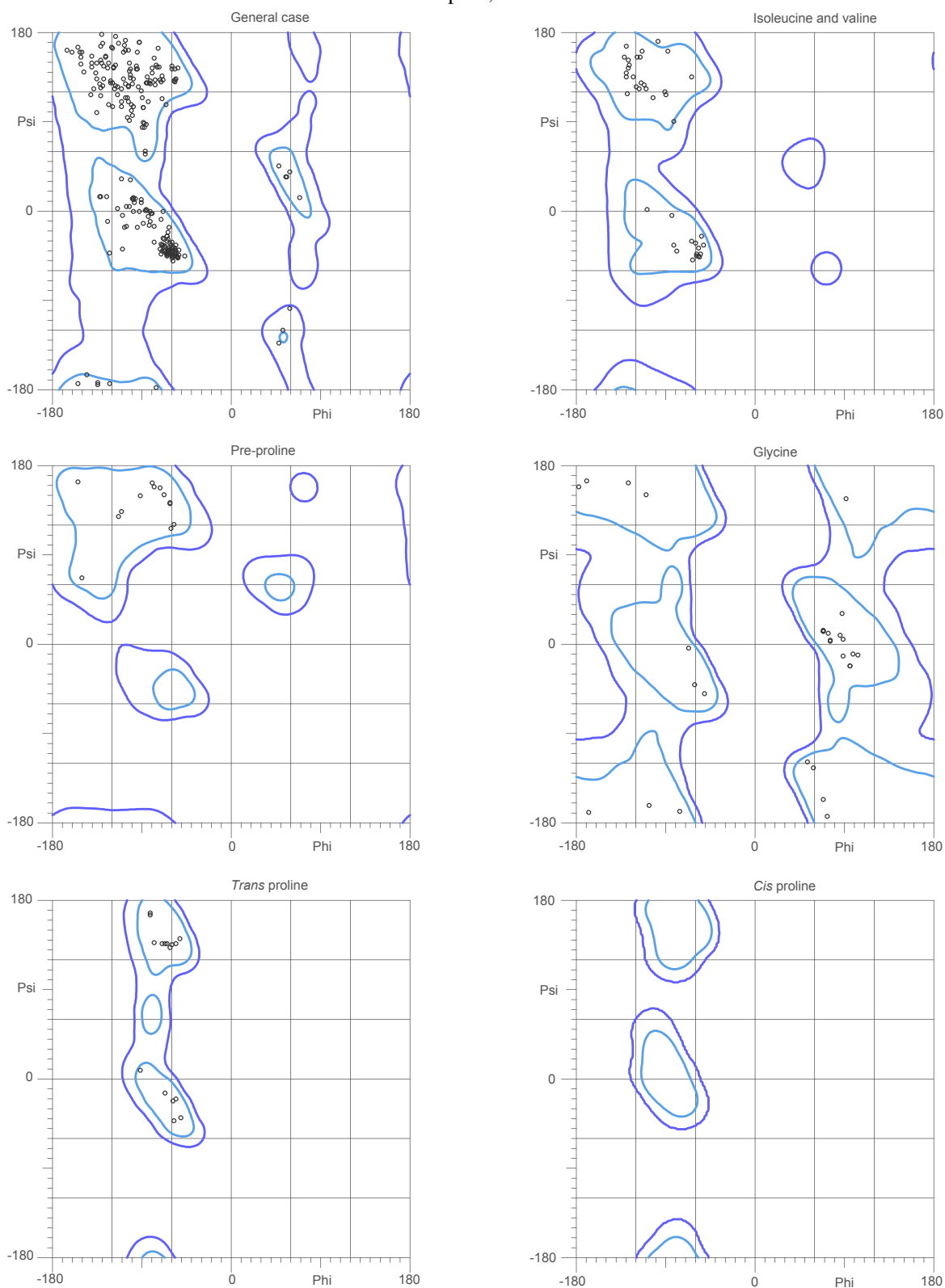


# MolProbity Ramachandran analysis

7aol.H.pdb, model 1



98.8% (329/333) of all residues were in favored (98%) regions.

100.0% (333/333) of all residues were in allowed (>99.8%) regions.

There were no outliers.