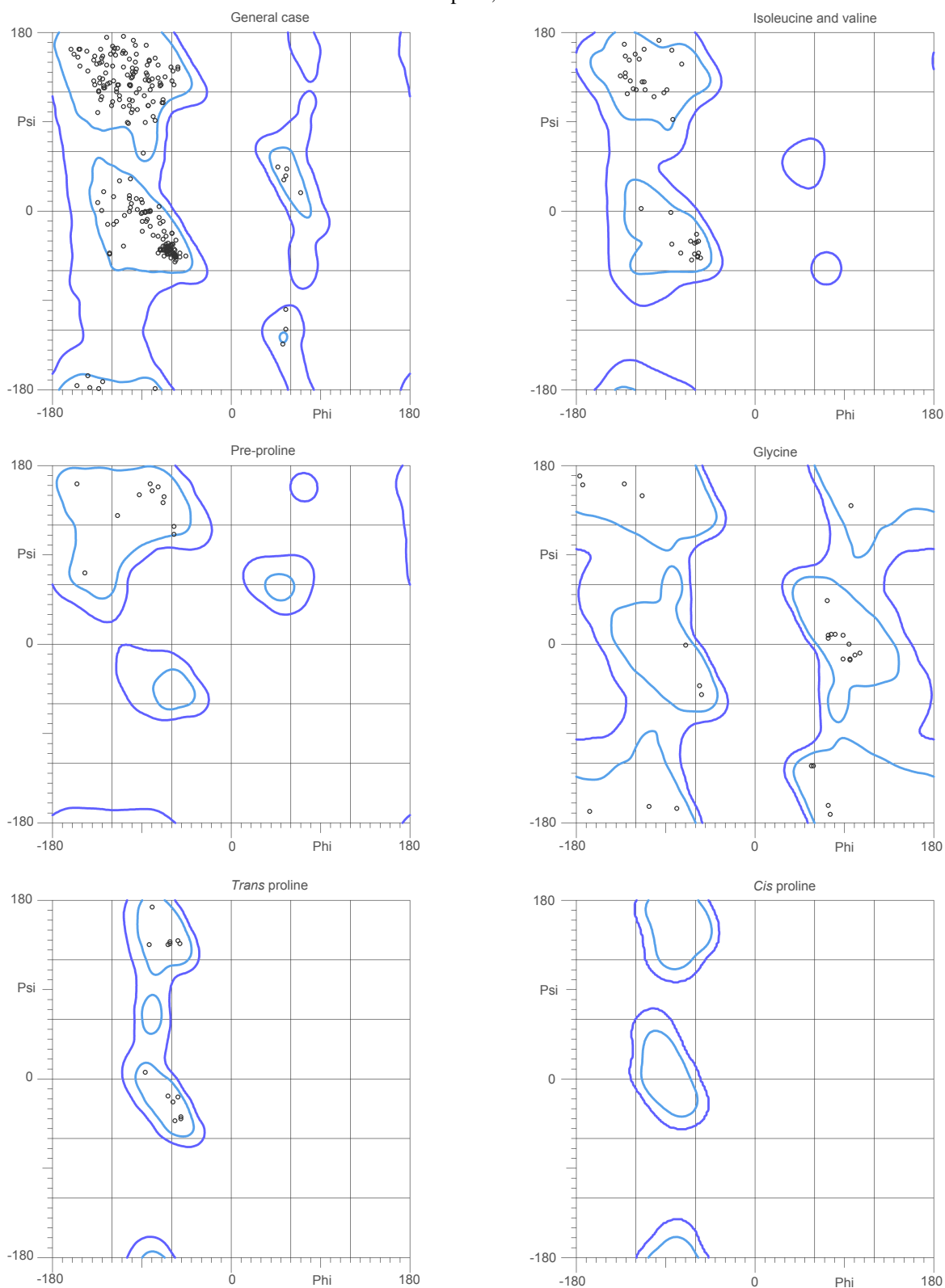


MolProbity Ramachandran analysis

7axo.H.pdb, model 1



98.8% (318/322) of all residues were in favored (98%) regions.

100.0% (322/322) of all residues were in allowed (>99.8%) regions.

There were no outliers.