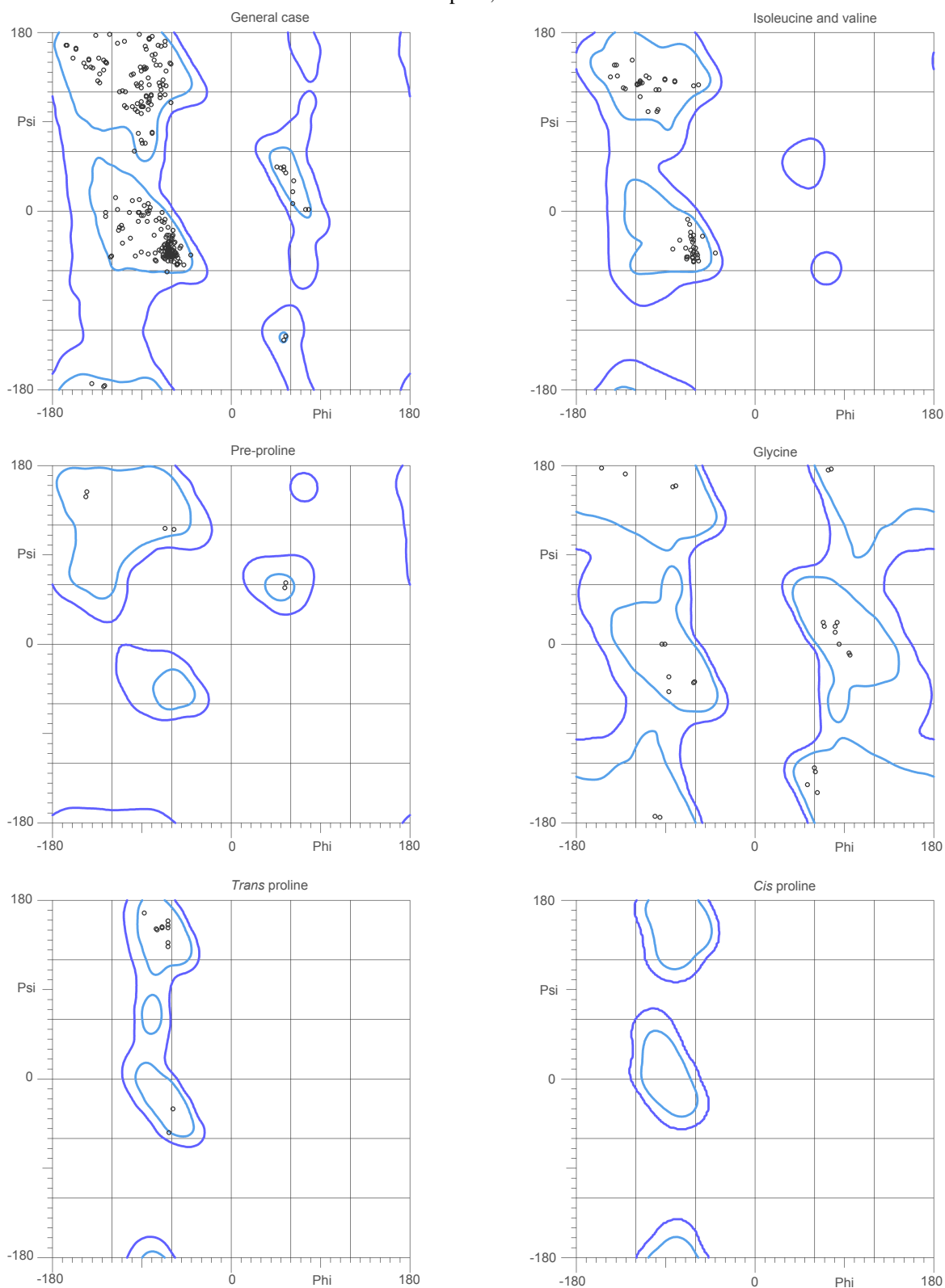# MolProbity Ramachandran analysis

5ruu.H.pdb, model 1

98.9% (346/350) of all residues were in favored (98%) regions.

100.0% (350/350) of all residues were in allowed (>99.8%) regions.

There were no outliers.