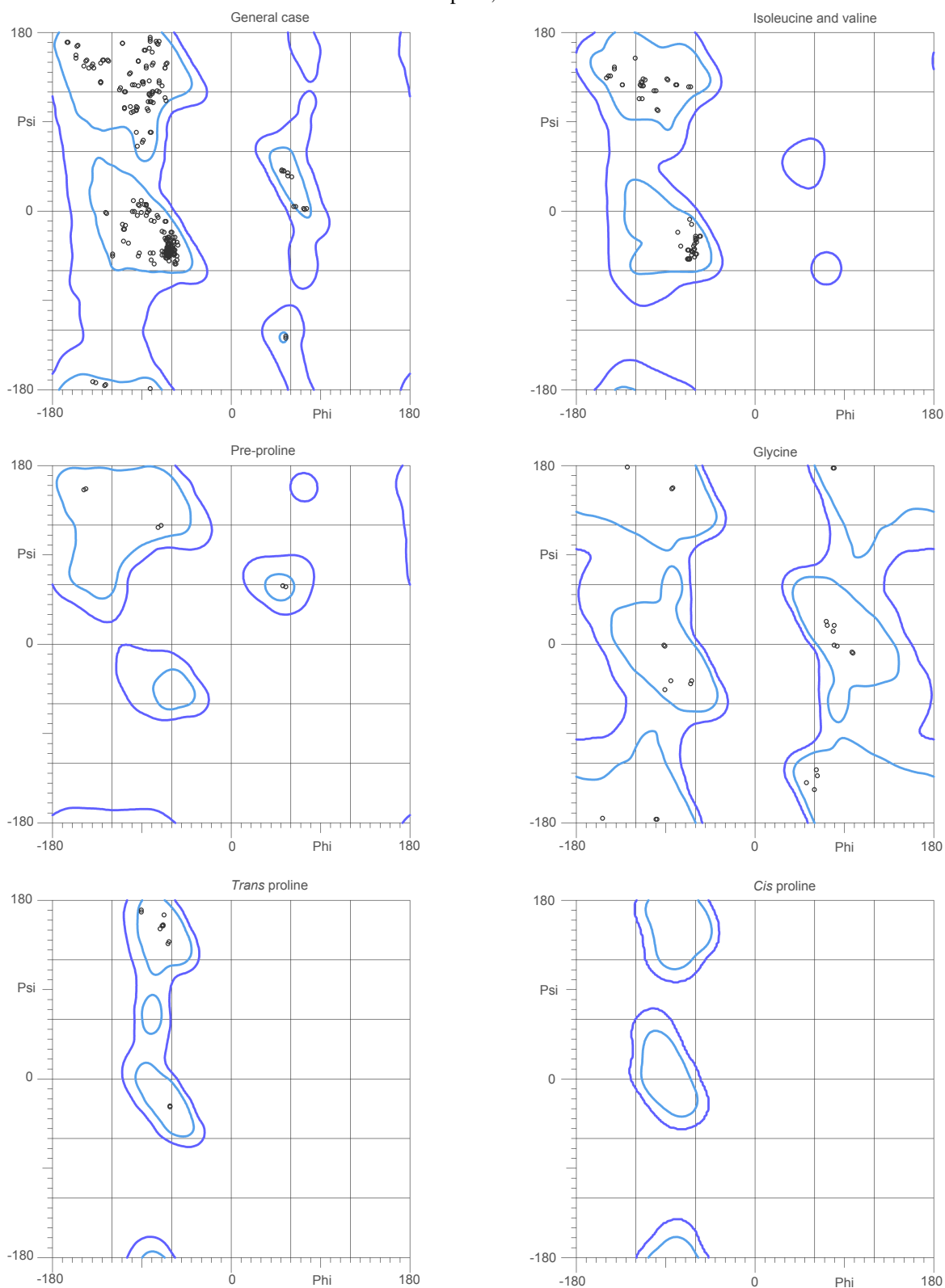


# MolProbity Ramachandran analysis

5s34.H.pdb, model 1



100.0% (346/346) of all residues were in favored (98%) regions.

100.0% (346/346) of all residues were in allowed (>99.8%) regions.

There were no outliers.