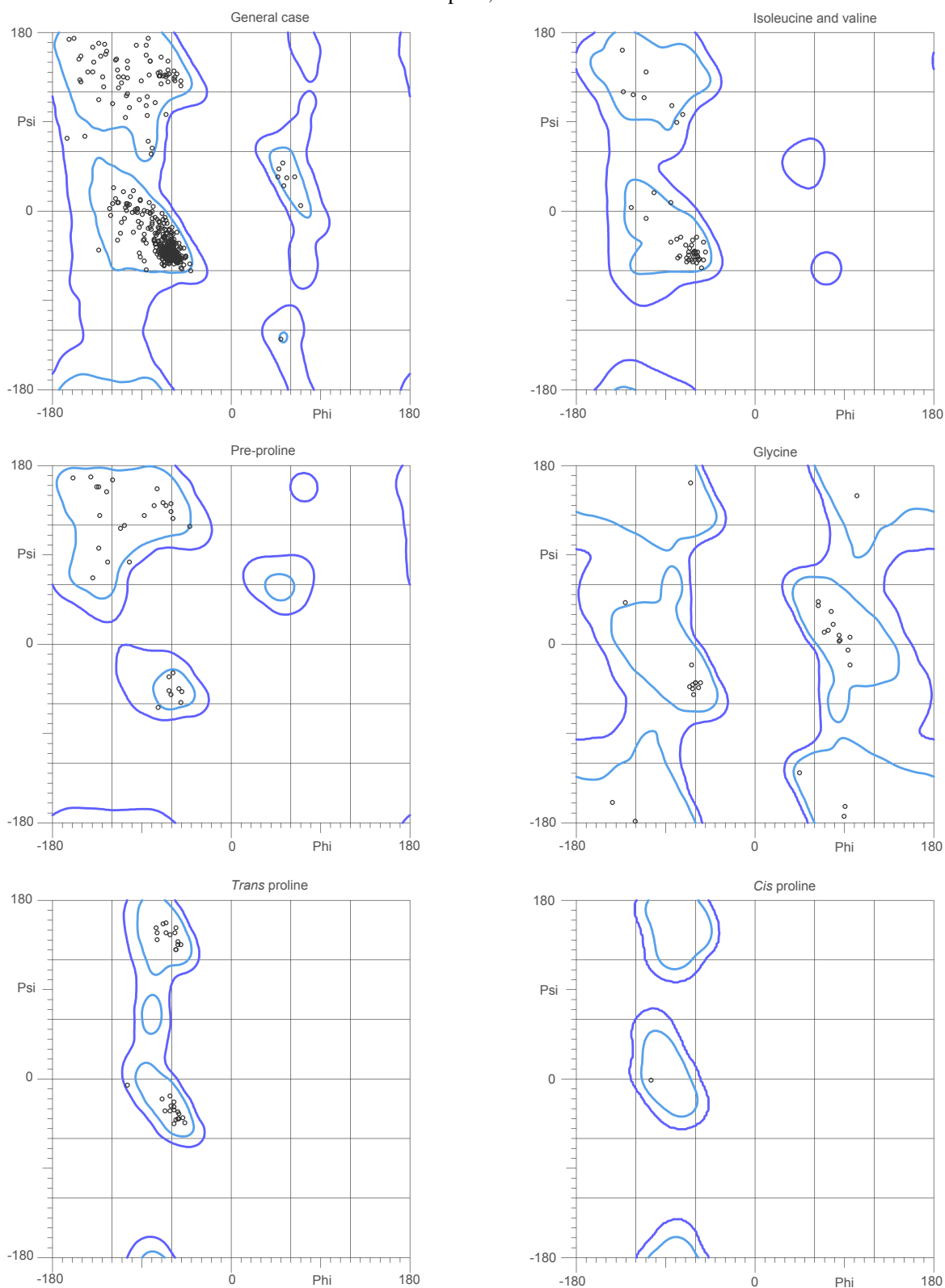


# MolProbity Ramachandran analysis

2iul.H.pdb, model 1



97.9% (571/583) of all residues were in favored (98%) regions.

100.0% (583/583) of all residues were in allowed (>99.8%) regions.

There were no outliers.