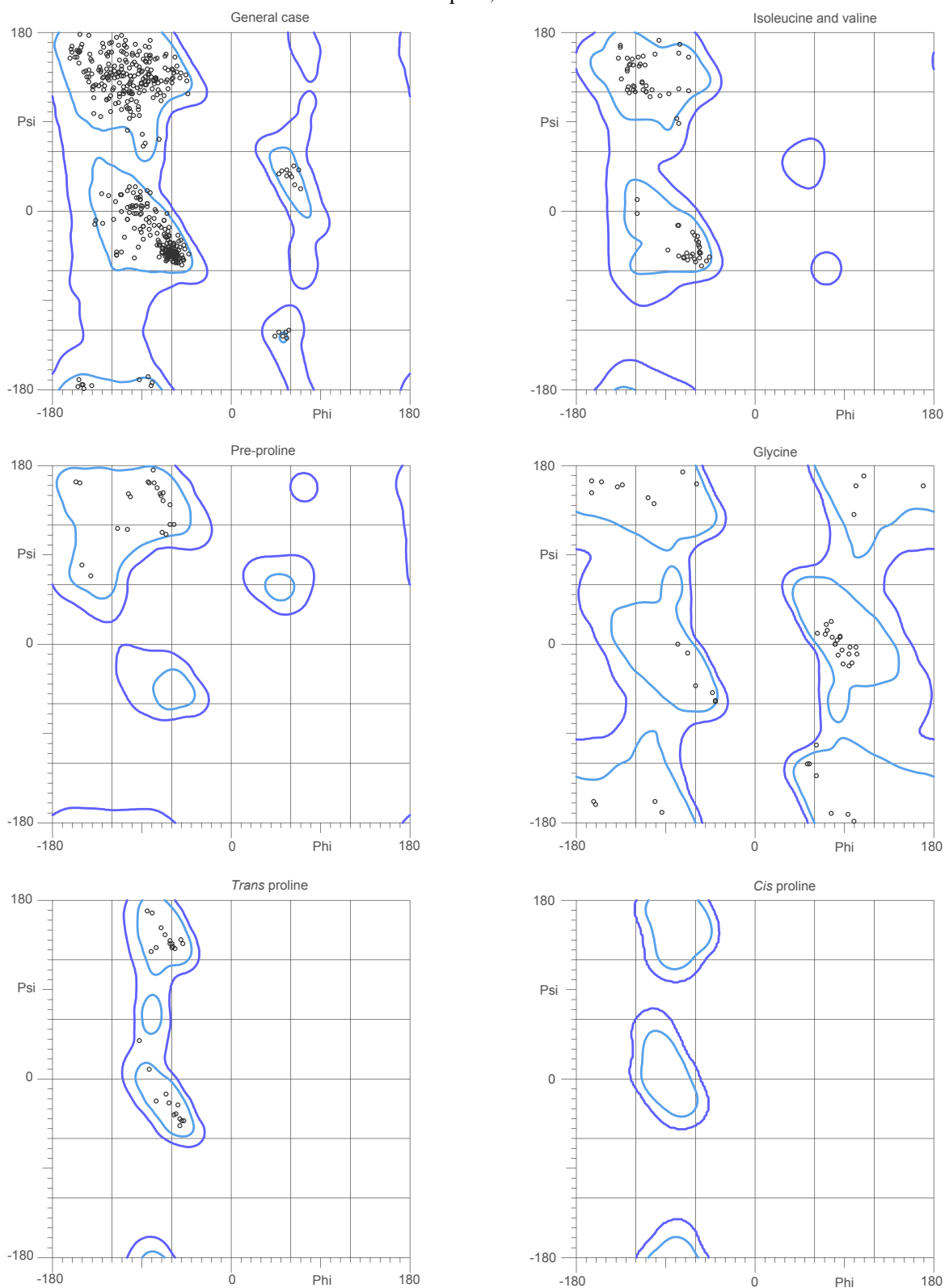


# MolProbity Ramachandran analysis

4hi3.H.pdb, model 1



96.6% (577/597) of all residues were in favored (98%) regions.

100.0% (597/597) of all residues were in allowed (>99.8%) regions.

There were no outliers.