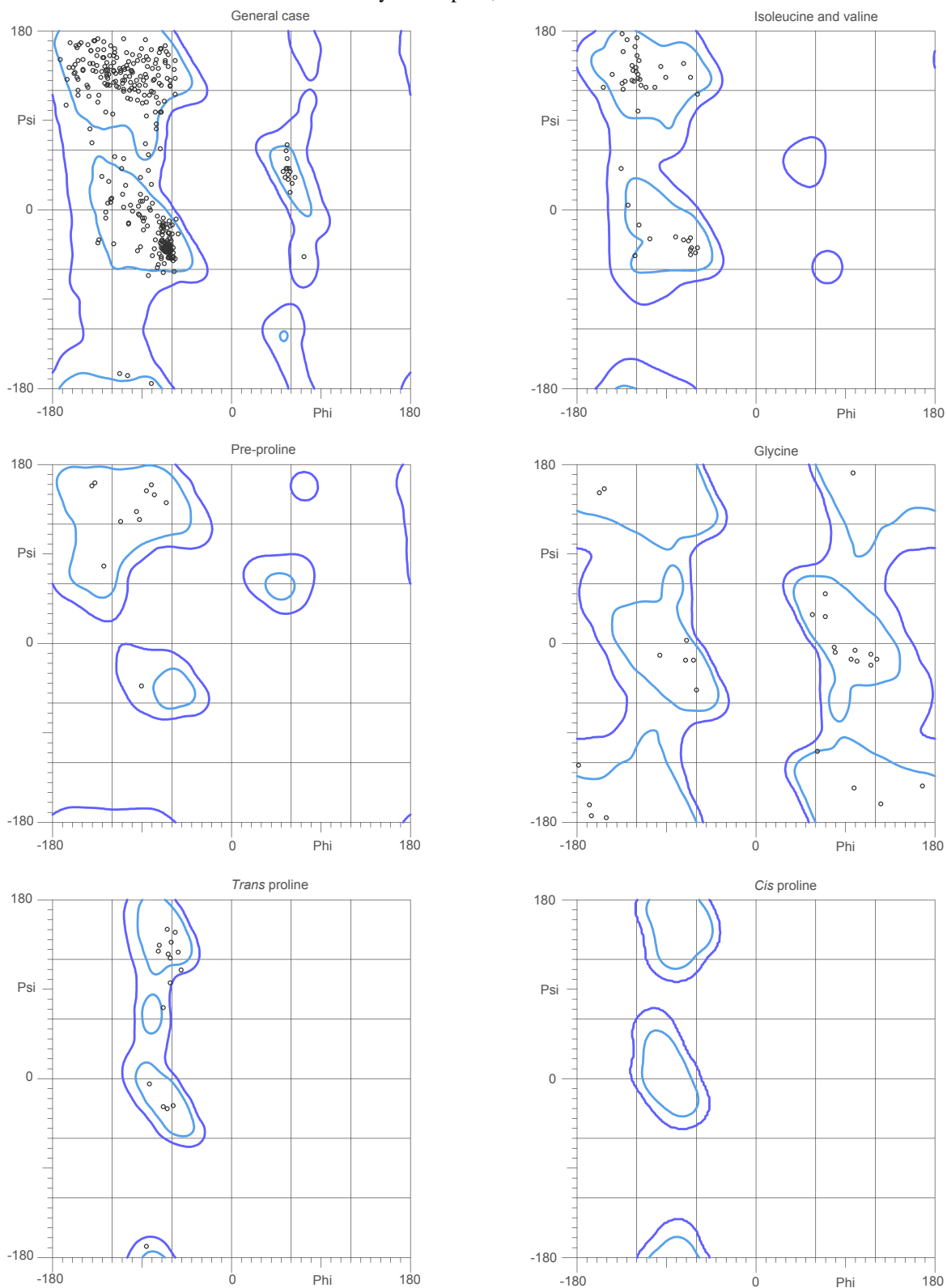


# MolProbity Ramachandran analysis

6yva.H.pdb, model 1



93.3% (402/431) of all residues were in favored (98%) regions.

100.0% (431/431) of all residues were in allowed (>99.8%) regions.

There were no outliers.