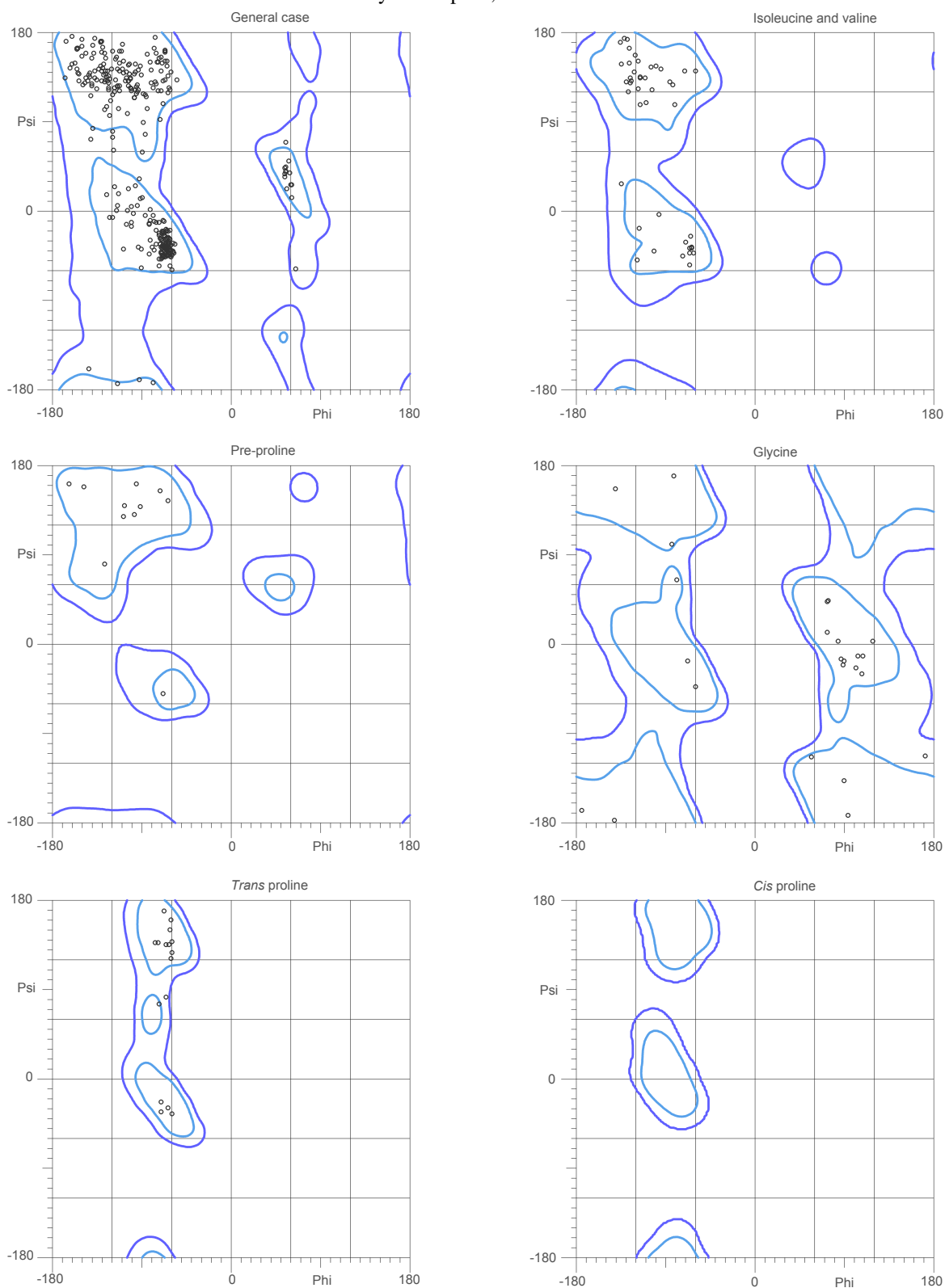


# MolProbity Ramachandran analysis

6yva.H.pdb, model 1



96.6% (393/407) of all residues were in favored (98%) regions.

100.0% (407/407) of all residues were in allowed (>99.8%) regions.

There were no outliers.