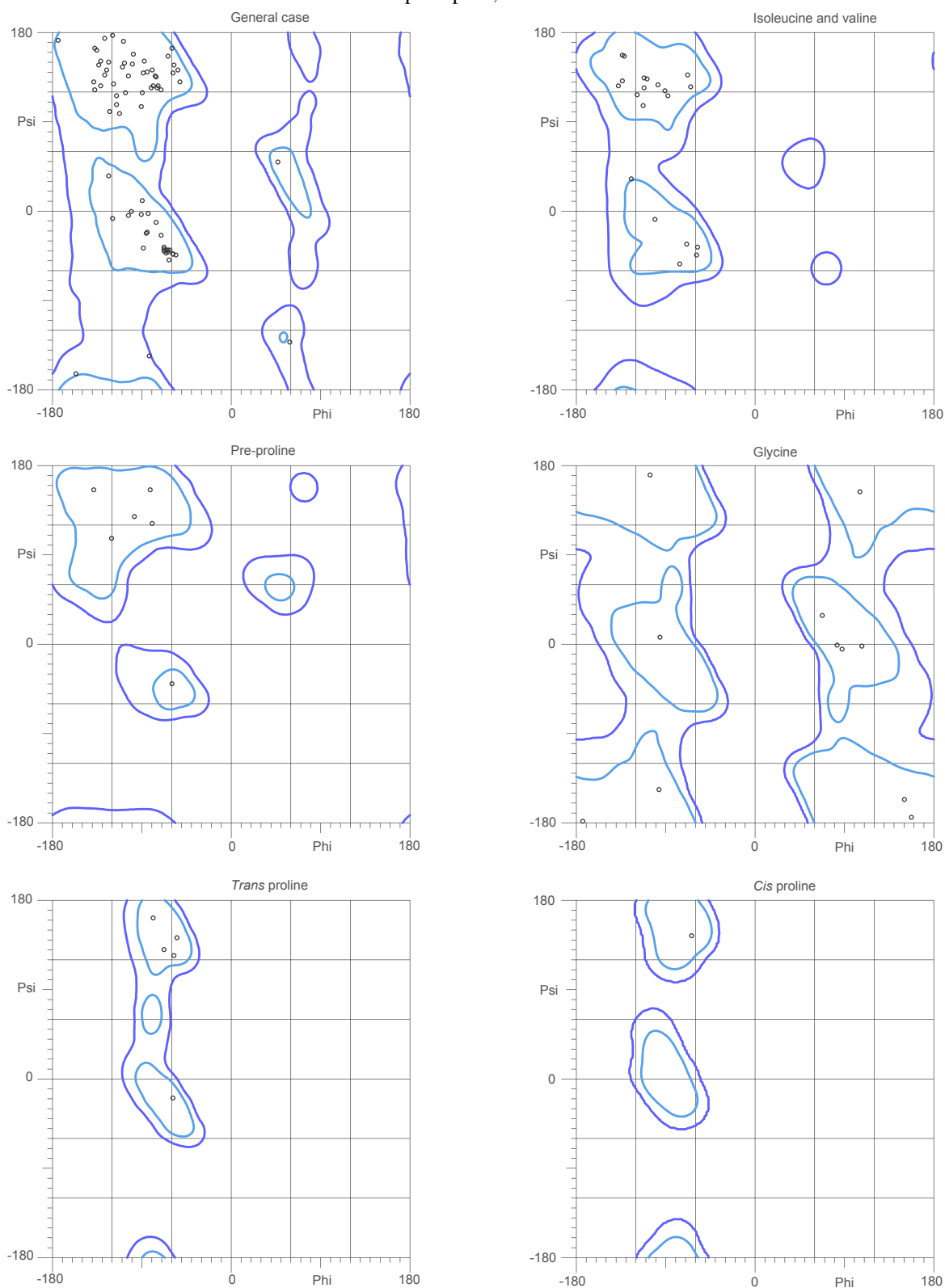


# MolProbity Ramachandran analysis

7k7p.H.pdb, model 1



96.5% (111/115) of all residues were in favored (98%) regions.

100.0% (115/115) of all residues were in allowed (>99.8%) regions.

There were no outliers.