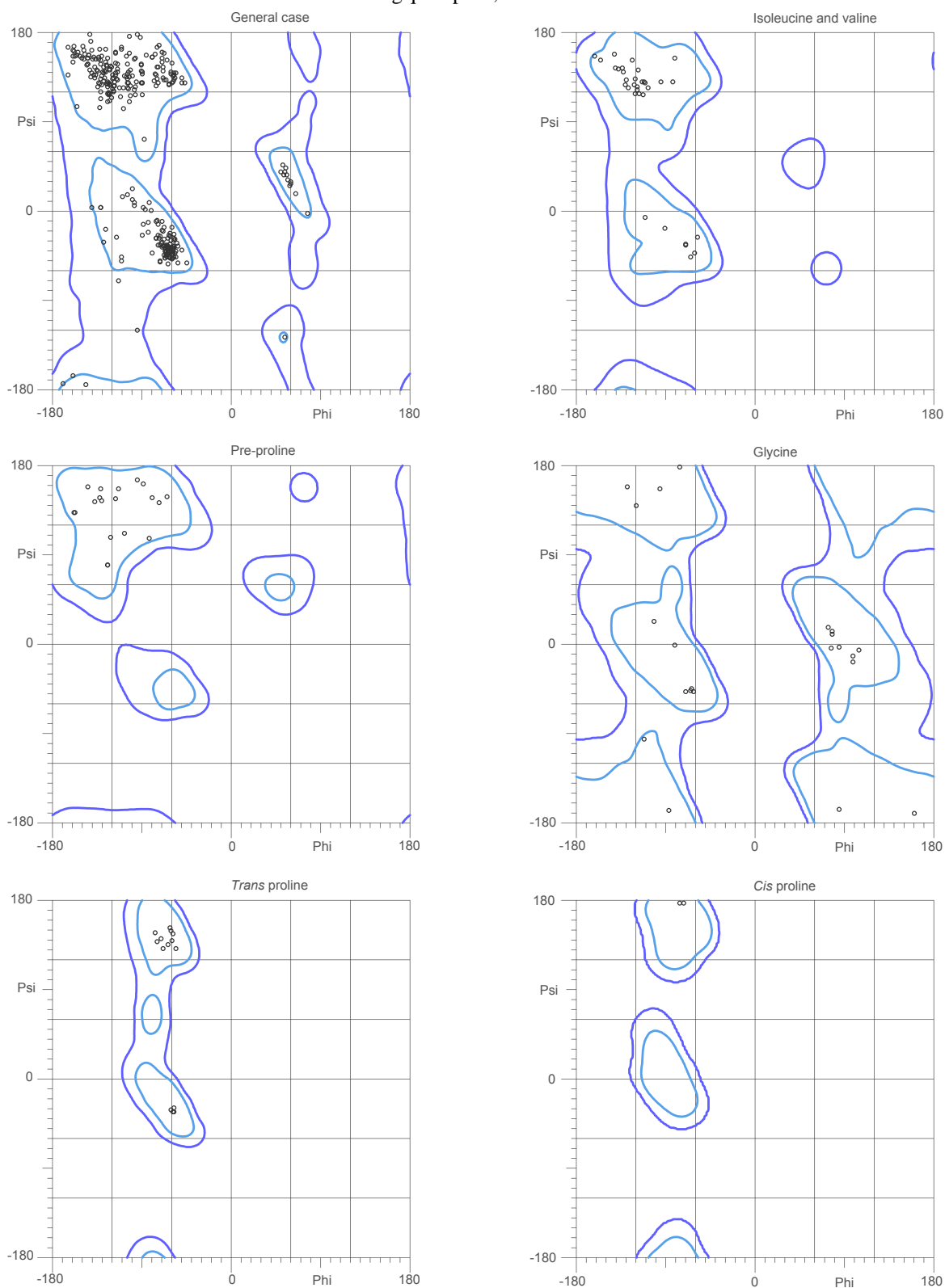


MolProbity Ramachandran analysis

7kgq.H.pdb, model 1



98.8% (397/402) of all residues were in favored (98%) regions.

100.0% (402/402) of all residues were in allowed (>99.8%) regions.

There were no outliers.