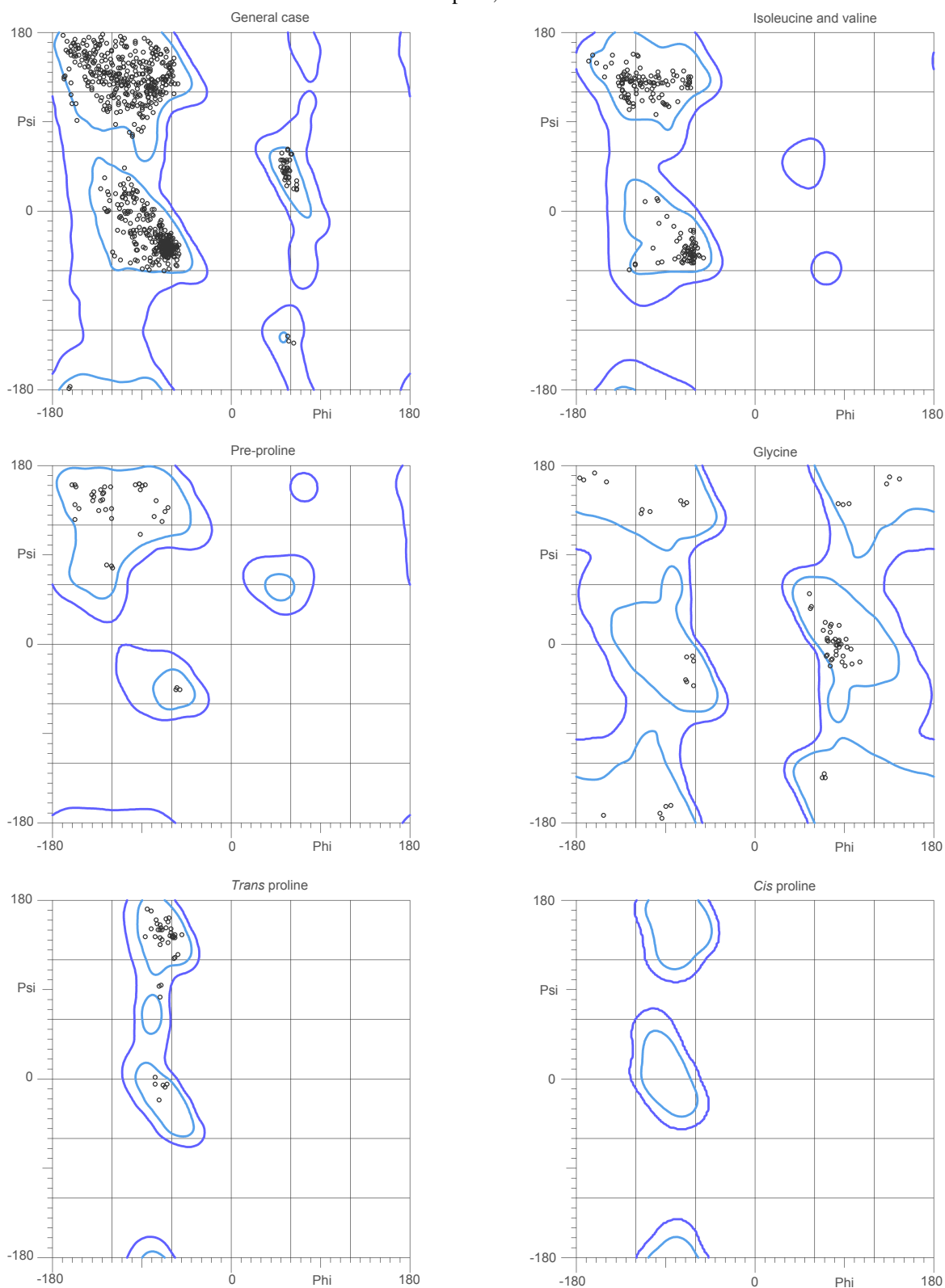


# MolProbity Ramachandran analysis

7k1o.H.pdb, model 1



98.3% (1022/1040) of all residues were in favored (98%) regions.

100.0% (1040/1040) of all residues were in allowed (>99.8%) regions.

There were no outliers.