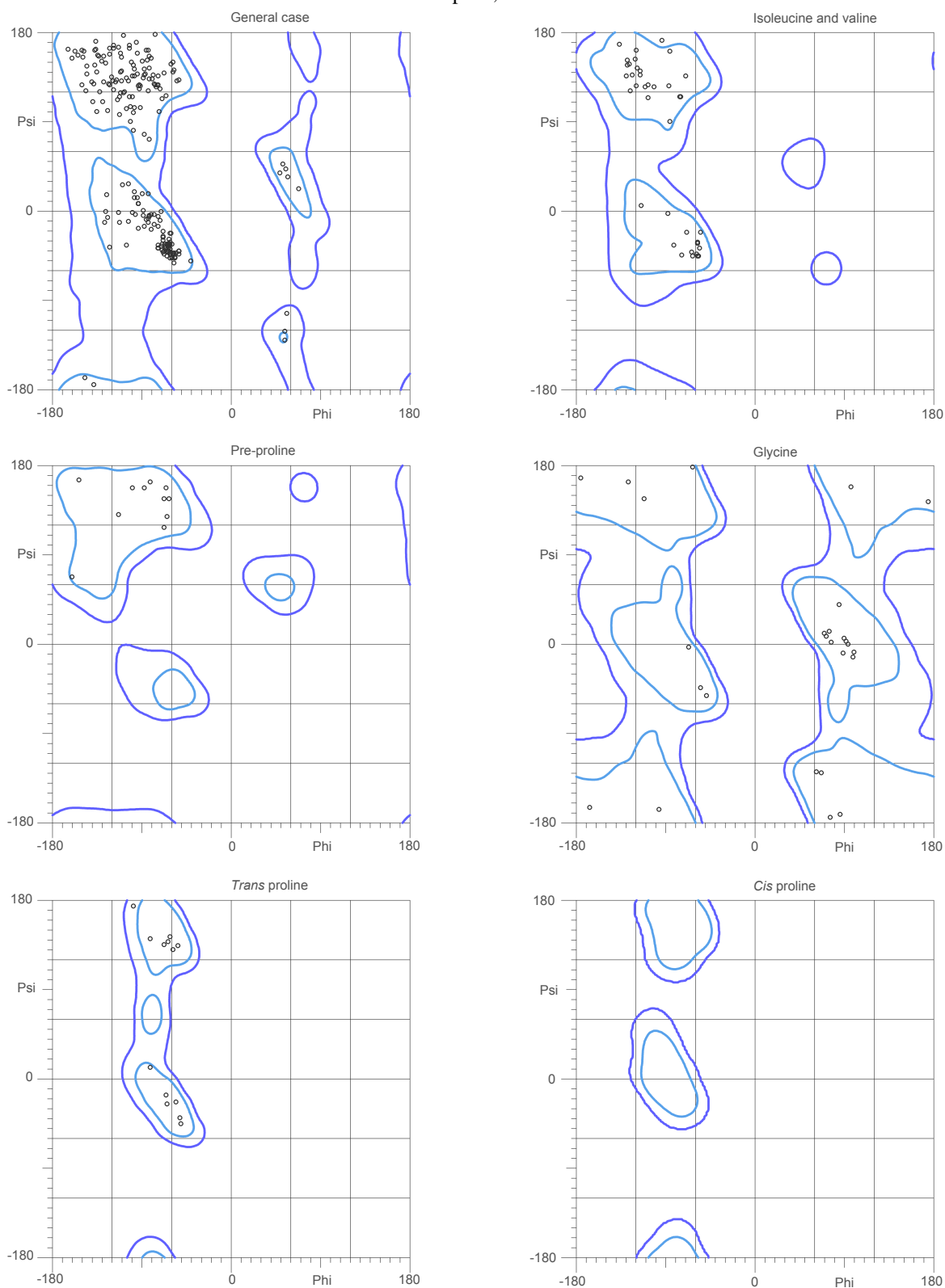


# MolProbity Ramachandran analysis

7k6d.H.pdb, model 1



98.7% (300/304) of all residues were in favored (98%) regions.

100.0% (304/304) of all residues were in allowed (>99.8%) regions.

There were no outliers.