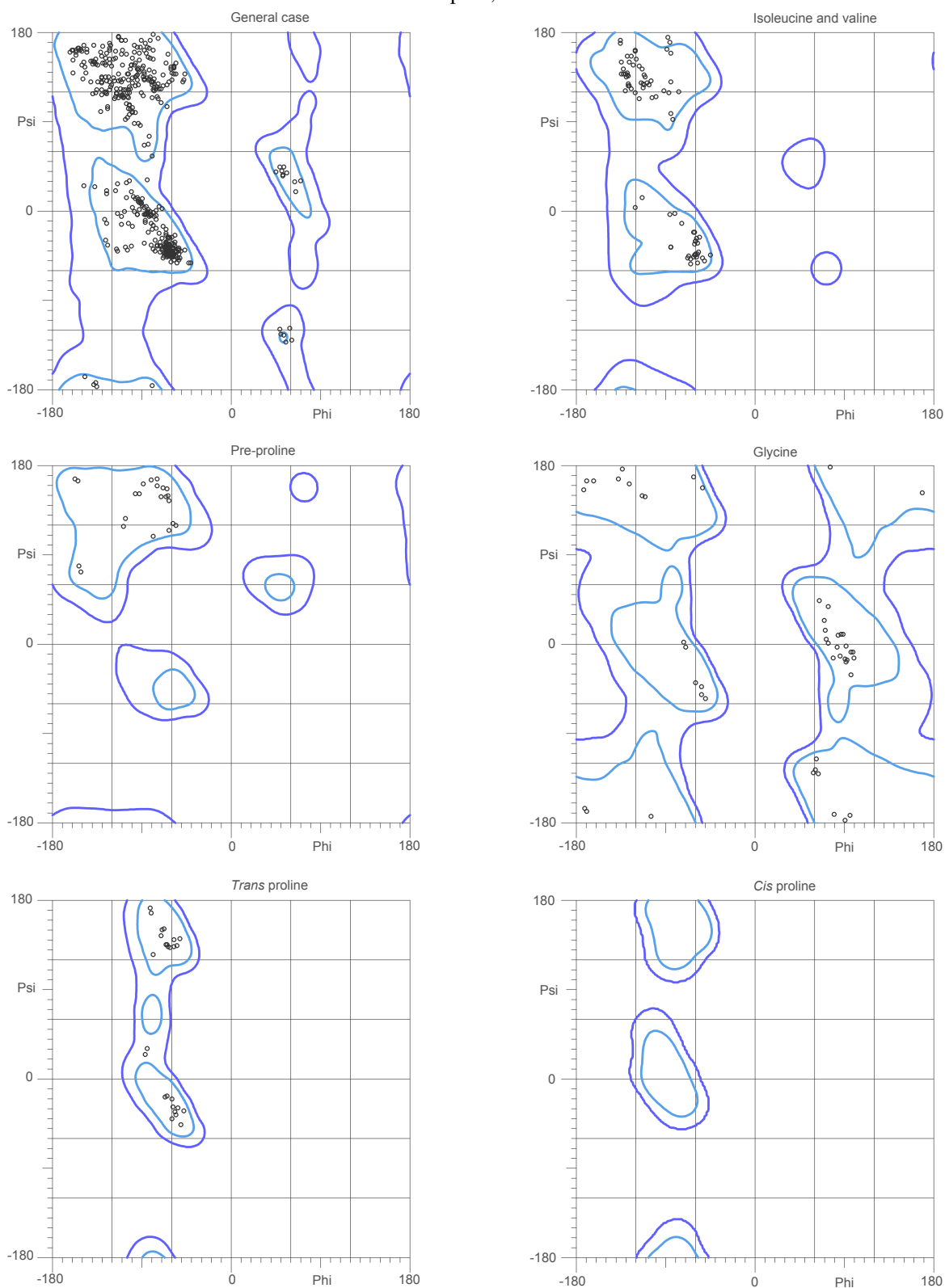


# MolProbity Ramachandran analysis

7k0f.H.pdb, model 1



98.2% (590/601) of all residues were in favored (98%) regions.

100.0% (601/601) of all residues were in allowed (>99.8%) regions.

There were no outliers.