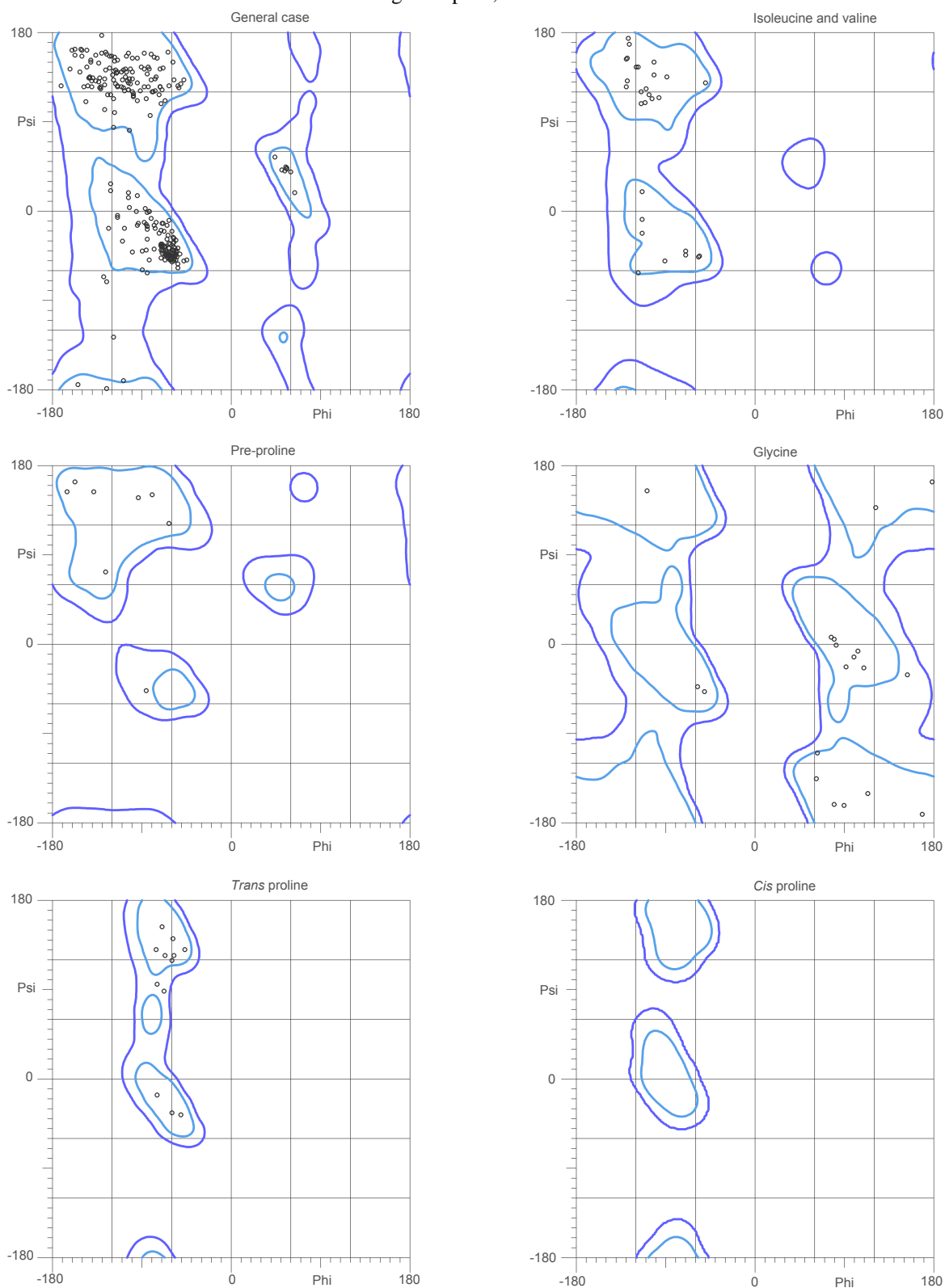


# MolProbity Ramachandran analysis

6xg3.H.pdb, model 1



96.9% (311/321) of all residues were in favored (98%) regions.

100.0% (321/321) of all residues were in allowed (>99.8%) regions.

There were no outliers.