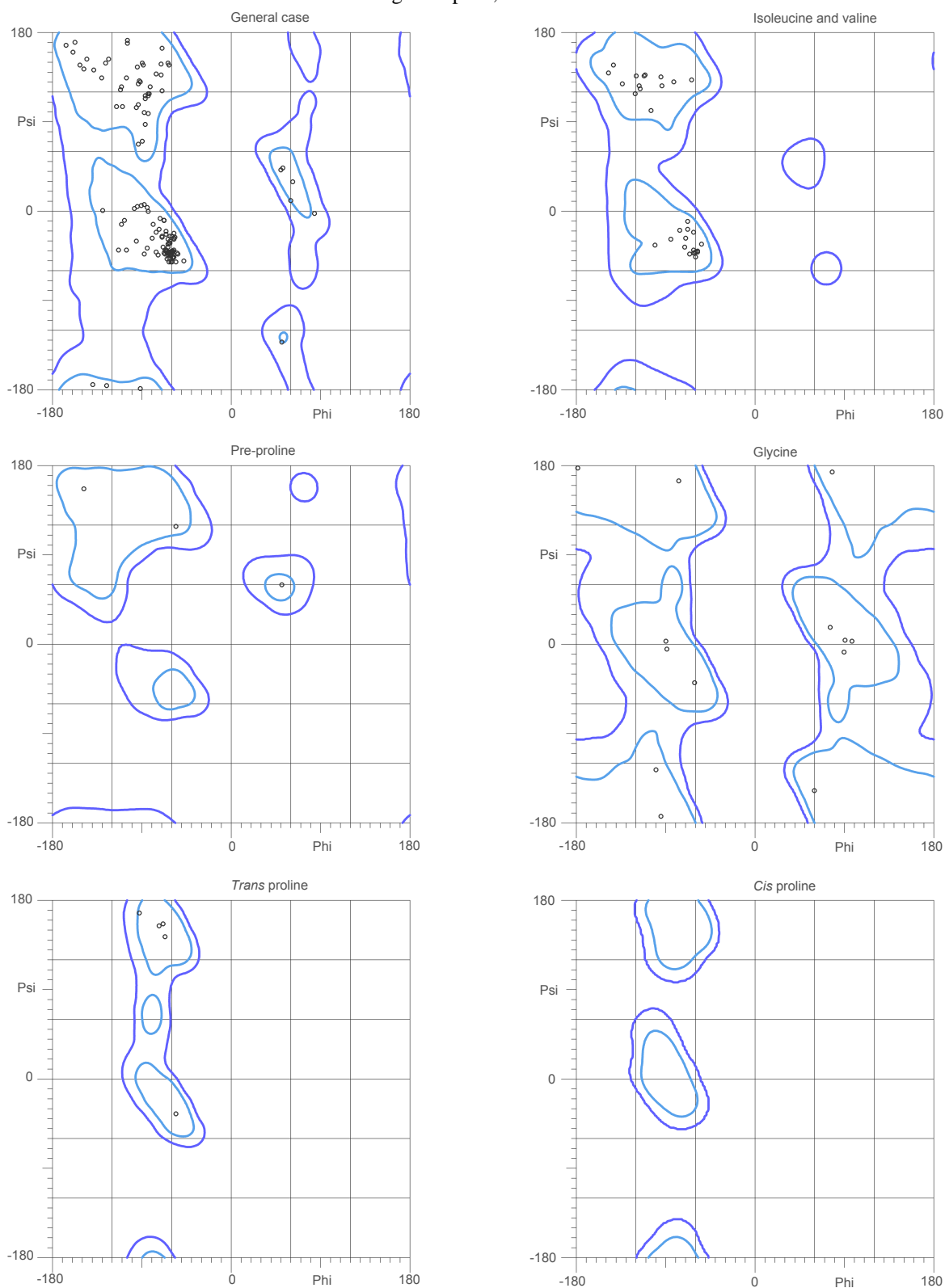


# MolProbity Ramachandran analysis

7kg3.H.pdb, model 1



98.9% (180/182) of all residues were in favored (98%) regions.

100.0% (182/182) of all residues were in allowed (>99.8%) regions.

There were no outliers.