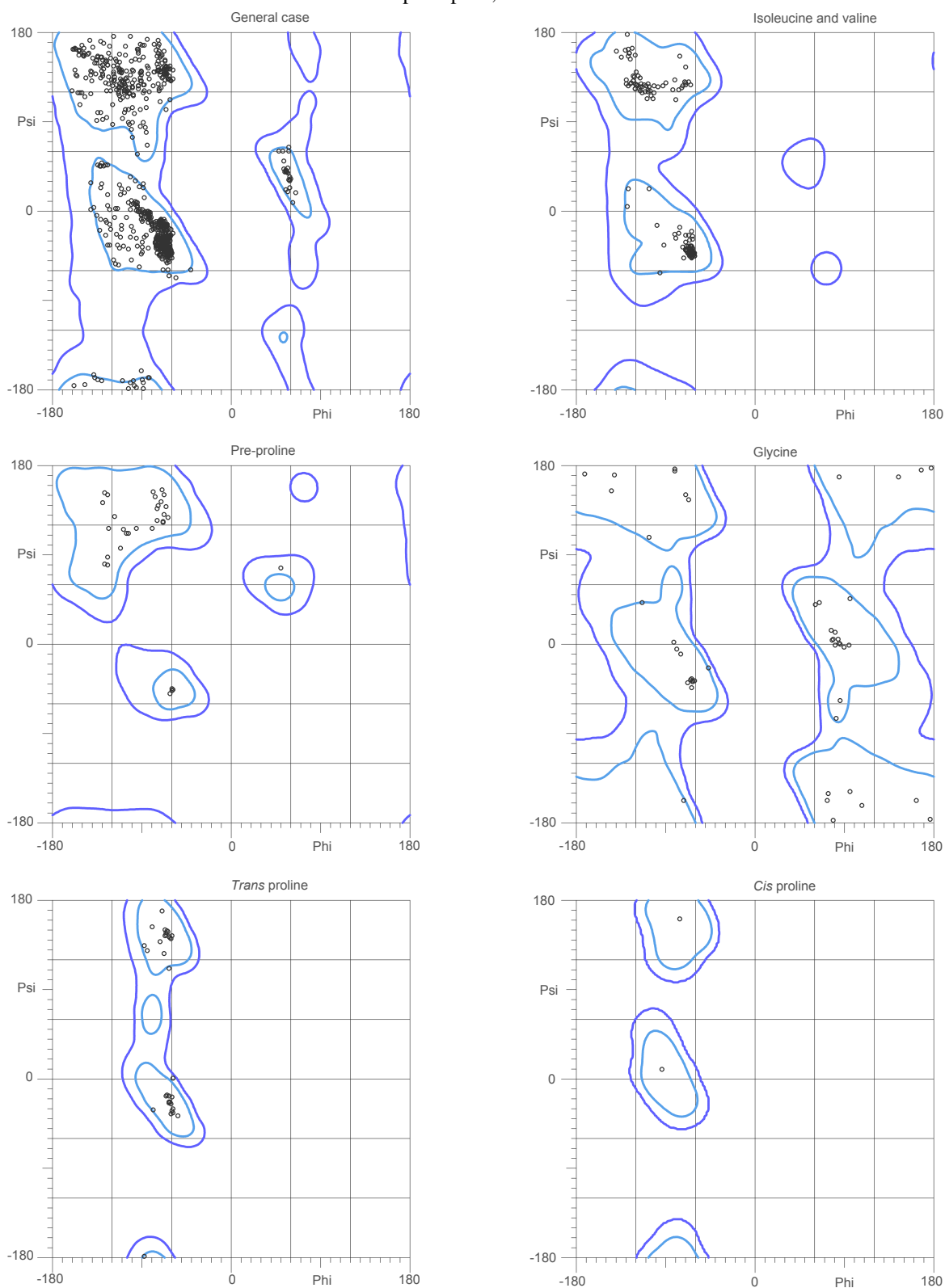


# MolProbity Ramachandran analysis

7aap.H.pdb, model 1



98.5% (1093/1110) of all residues were in favored (98%) regions.

100.0% (1110/1110) of all residues were in allowed (>99.8%) regions.

There were no outliers.