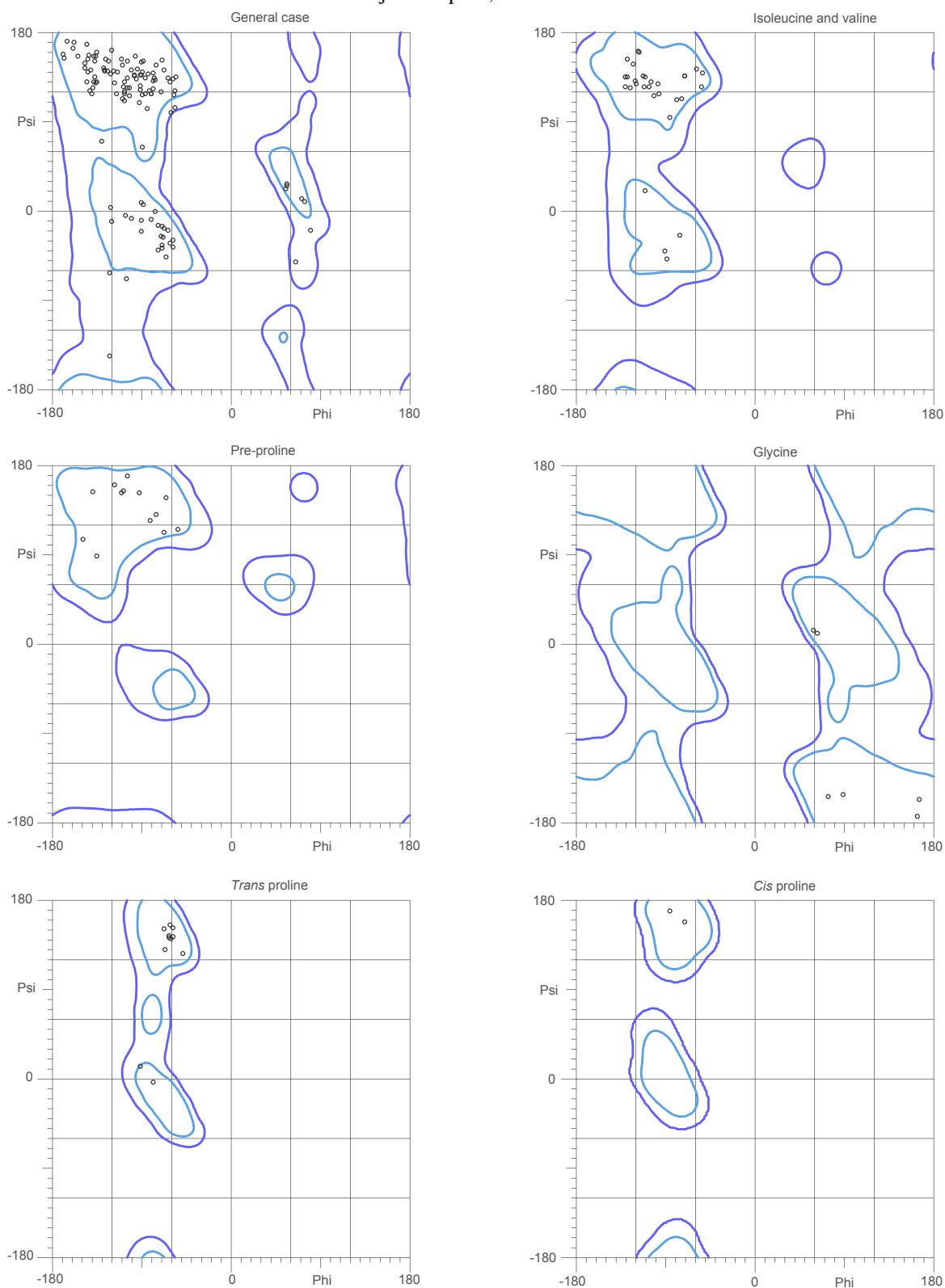


# MolProbity Ramachandran analysis

7jx6.H.pdb, model 1



95.2% (177/186) of all residues were in favored (98%) regions.

100.0% (186/186) of all residues were in allowed (>99.8%) regions.

There were no outliers.