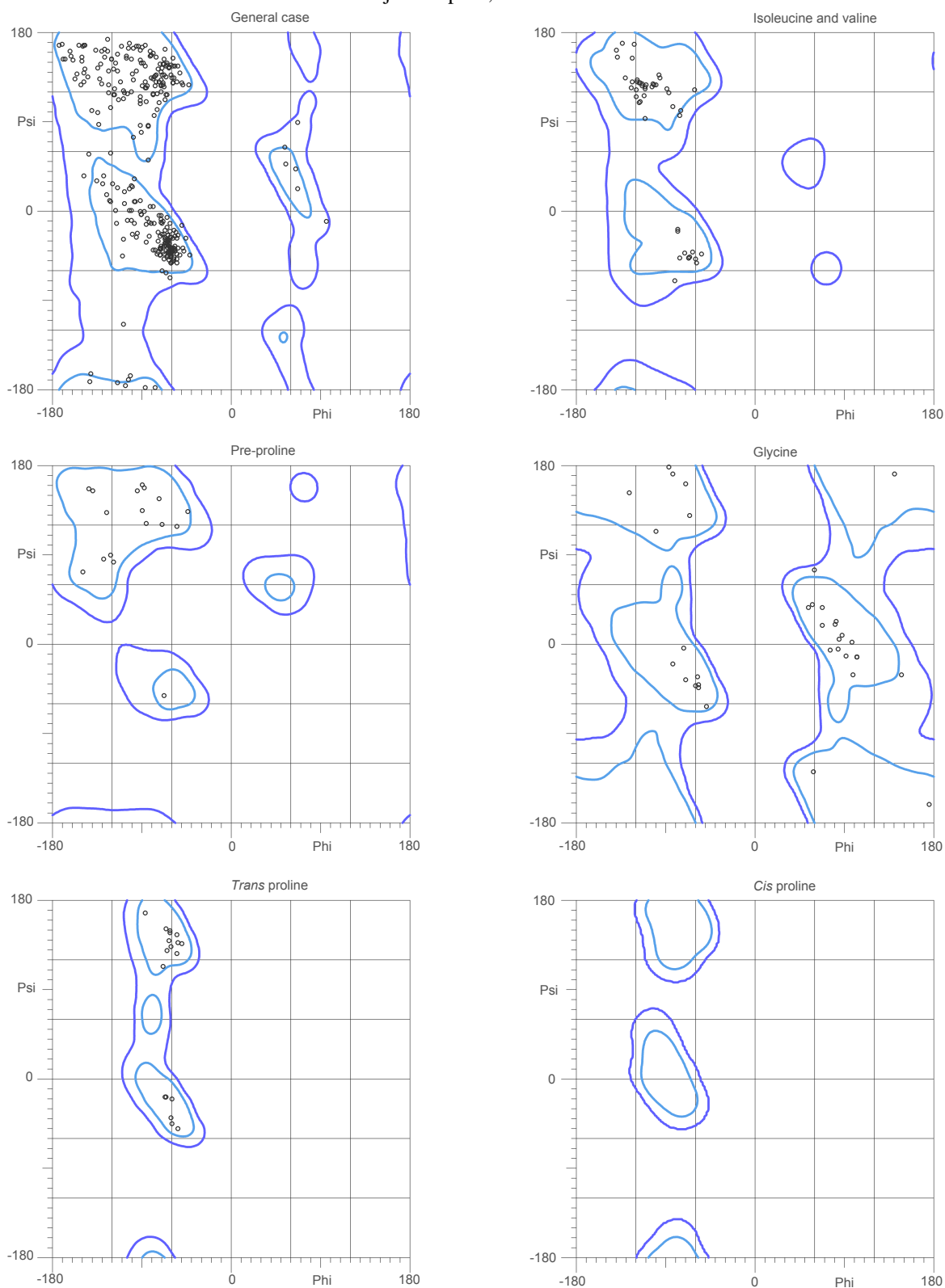


# MolProbity Ramachandran analysis

7jib.H.pdb, model 1



95.6% (411/430) of all residues were in favored (98%) regions.

100.0% (430/430) of all residues were in allowed (>99.8%) regions.

There were no outliers.