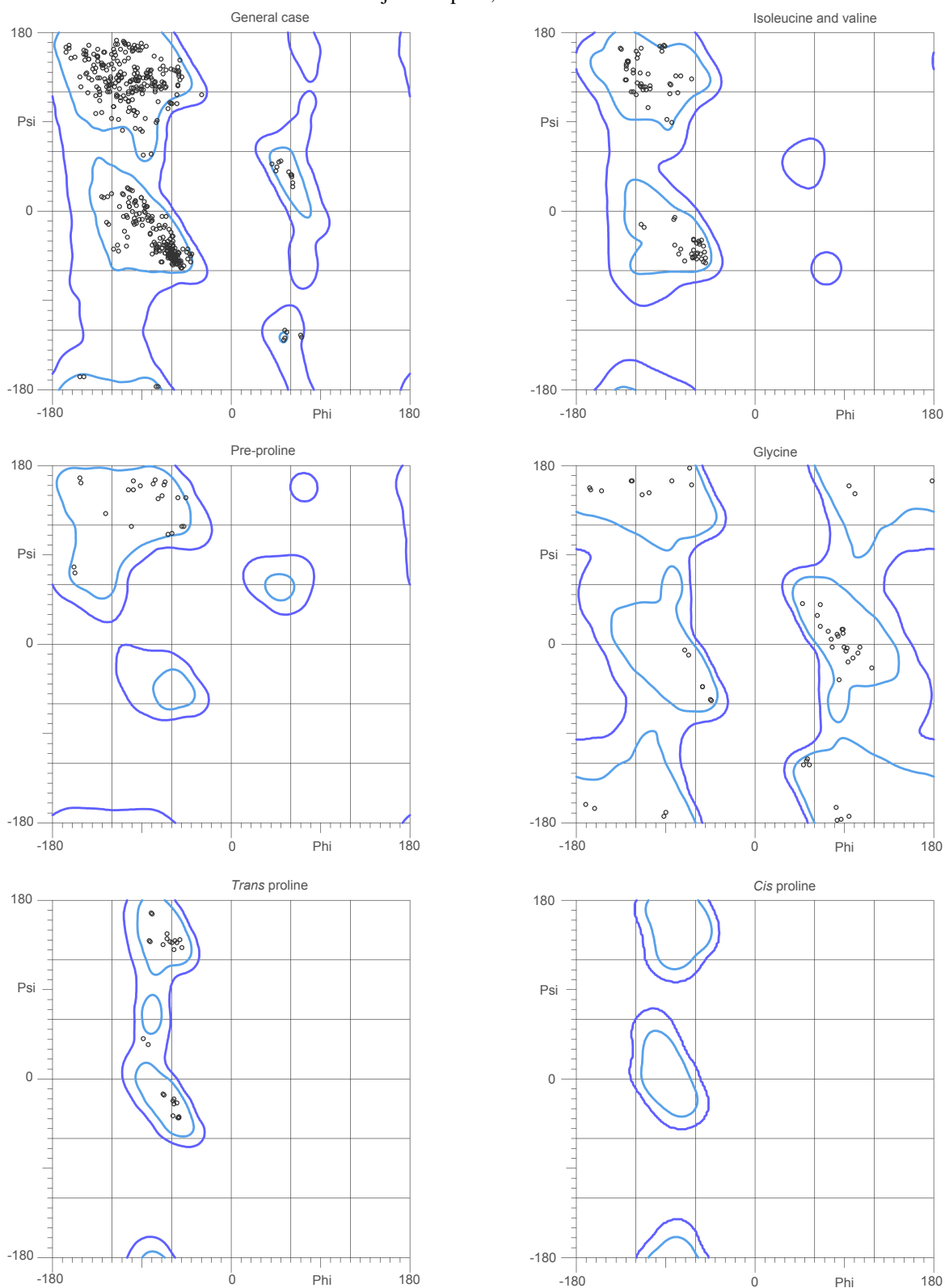


# MolProbity Ramachandran analysis

7jox.H.pdb, model 1



97.0% (590/608) of all residues were in favored (98%) regions.

100.0% (608/608) of all residues were in allowed (>99.8%) regions.

There were no outliers.