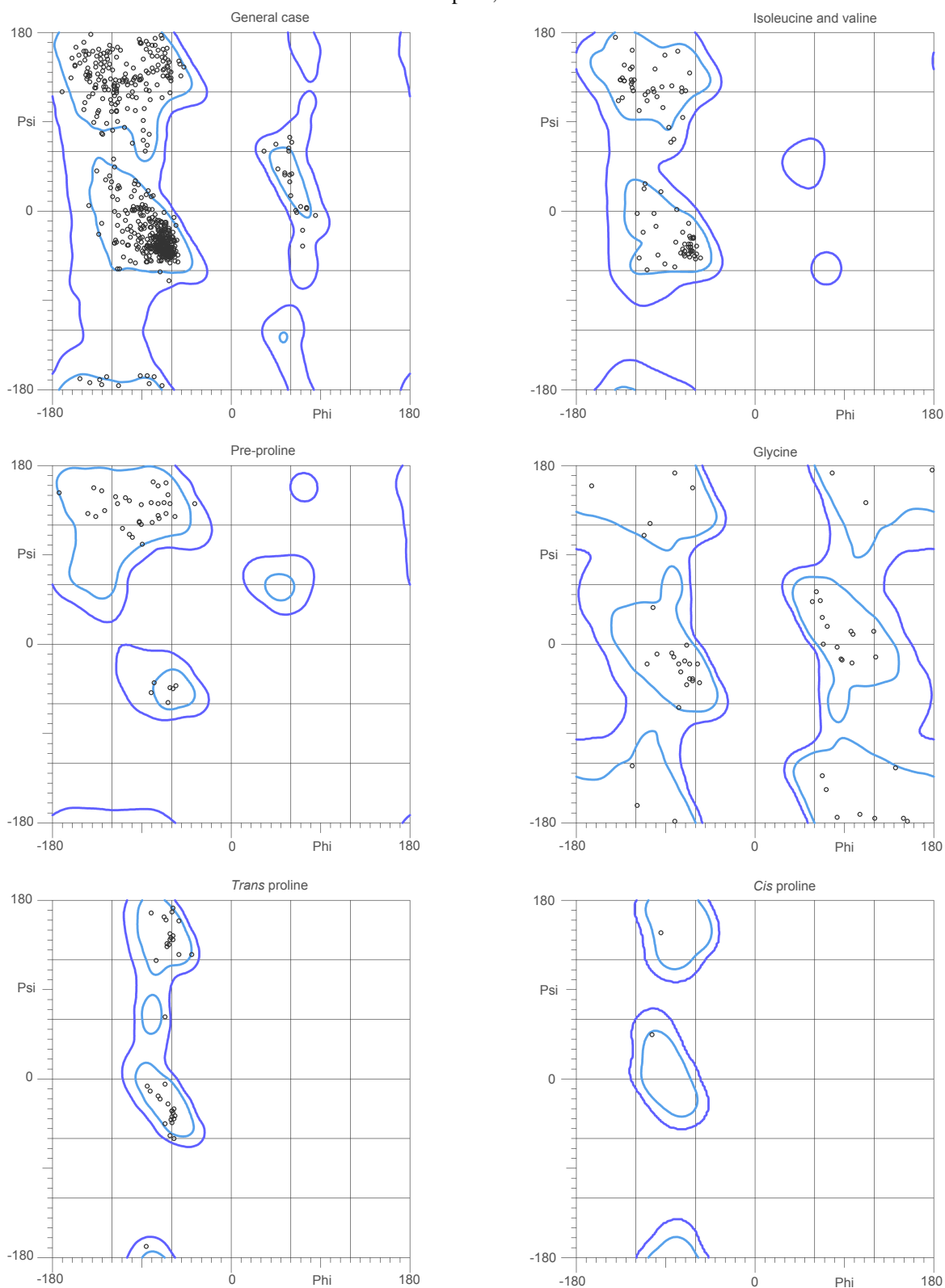


# MolProbity Ramachandran analysis

7a91.H.pdb, model 1



95.1% (792/833) of all residues were in favored (98%) regions.

100.0% (833/833) of all residues were in allowed (>99.8%) regions.

There were no outliers.