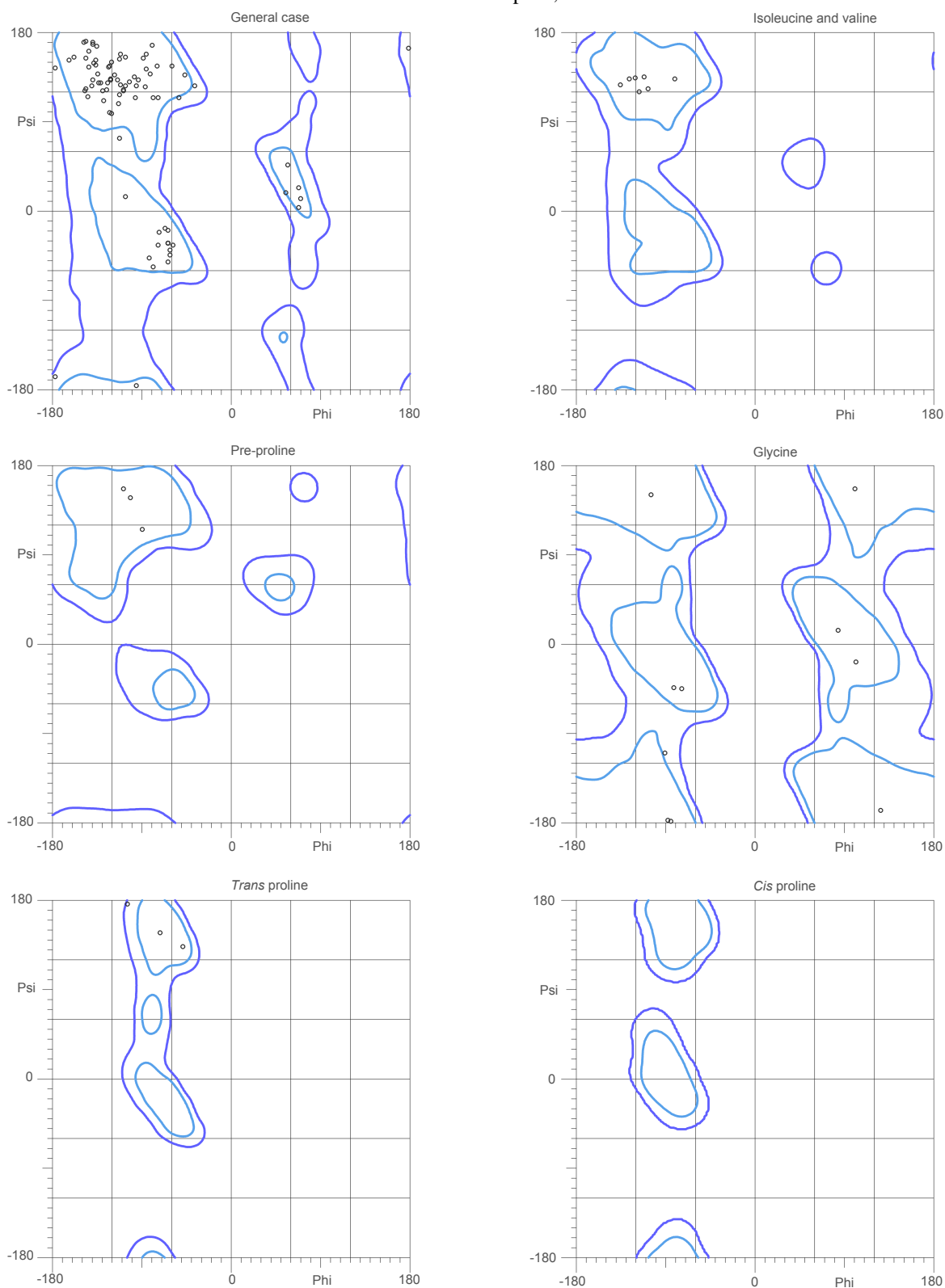


# MolProbity Ramachandran analysis

1ac6A.B99990010.pdb, model 1



93,5% (101/108) of all residues were in favored (98%) regions.

100,0% (108/108) of all residues were in allowed (>99.8%) regions.

There were no outliers.