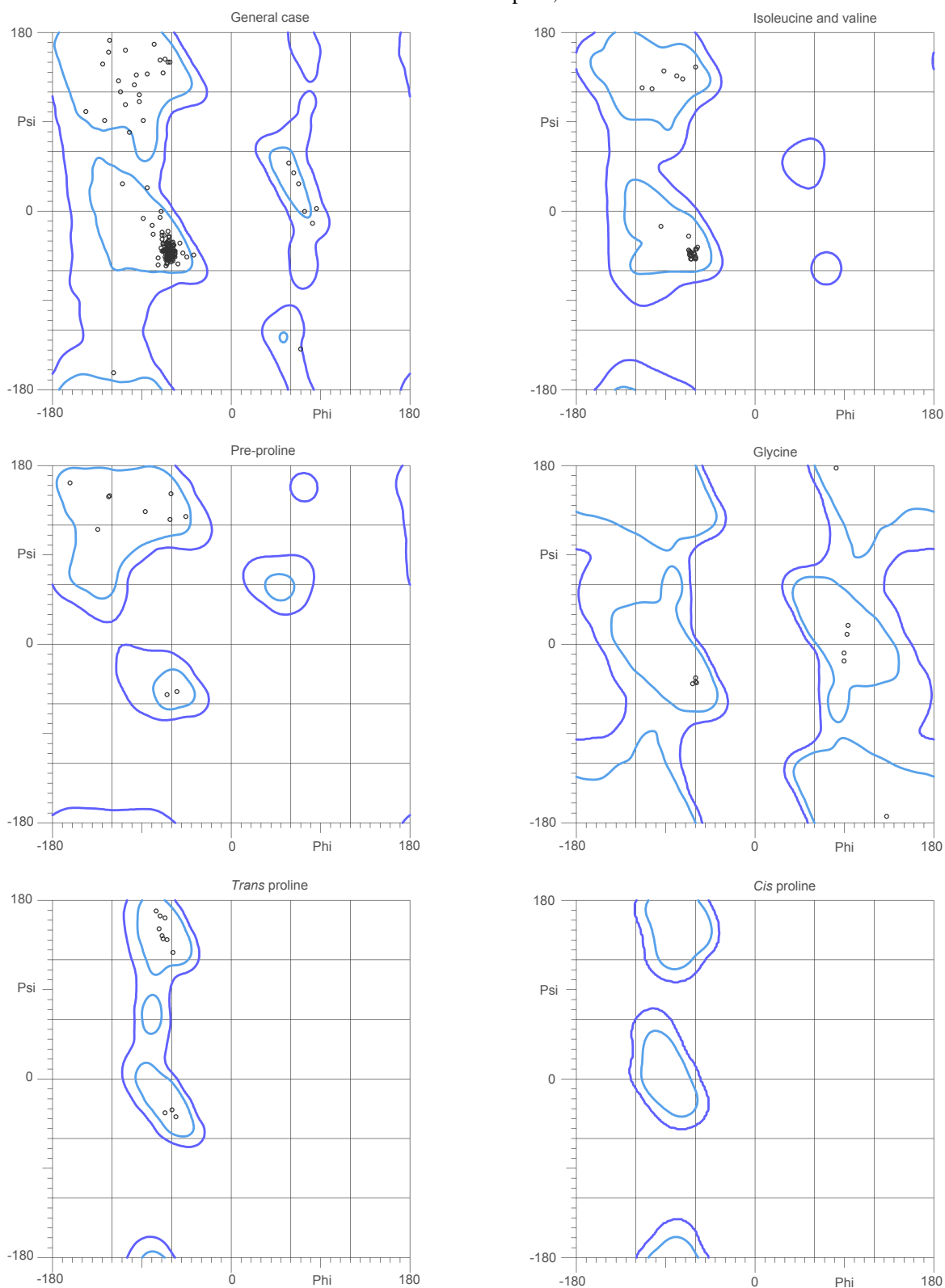


# MolProbity Ramachandran analysis

1a28A.B99990023.pdb, model 1



97,2% (242/249) of all residues were in favored (98%) regions.

100,0% (249/249) of all residues were in allowed (>99.8%) regions.

There were no outliers.