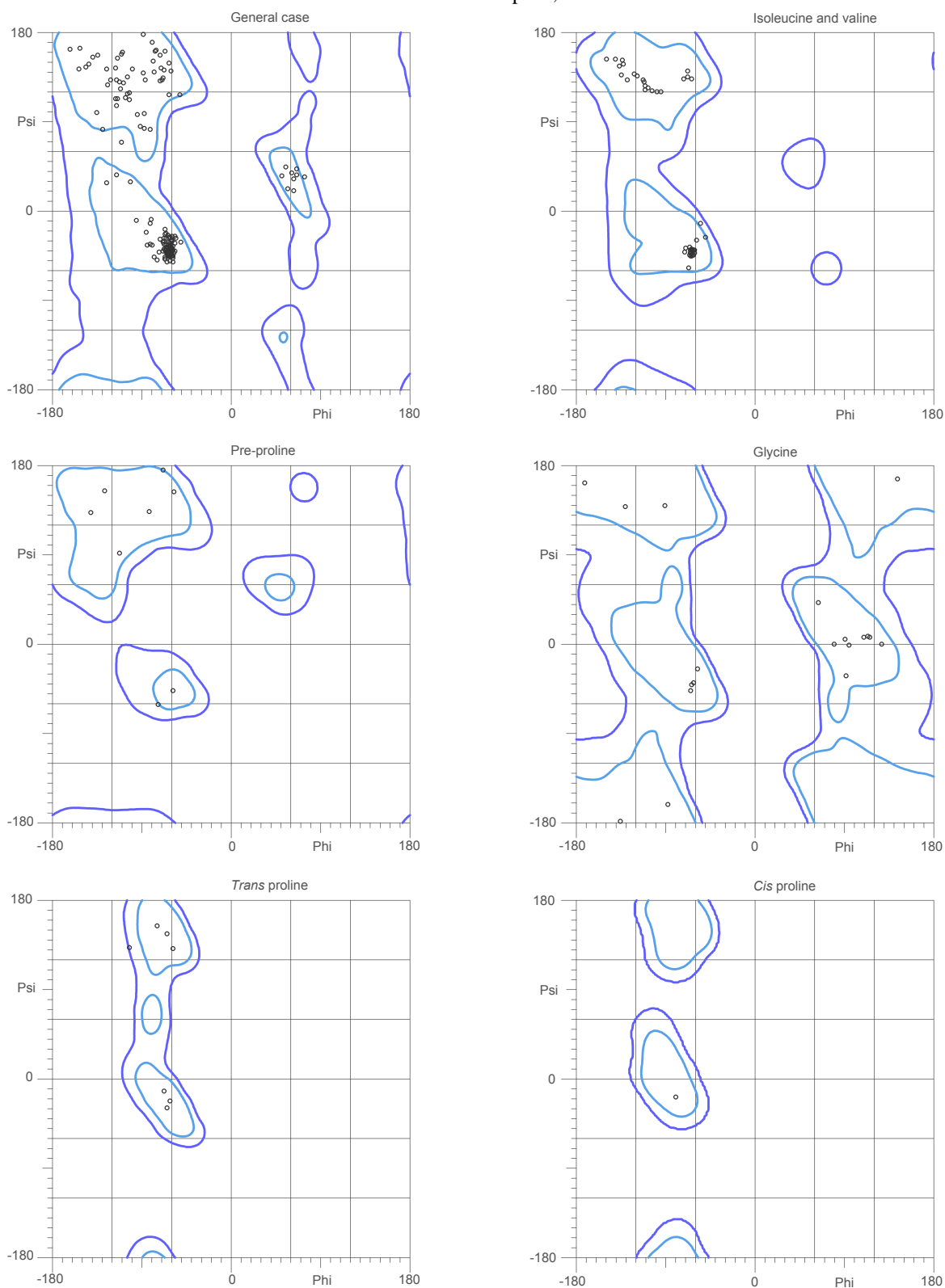# MolProbity Ramachandran analysis

1ad1A.B99990033.pdb, model 1

97,7% (256/262) of all residues were in favored (98%) regions.

100,0% (262/262) of all residues were in allowed (>99.8%) regions.

There were no outliers.