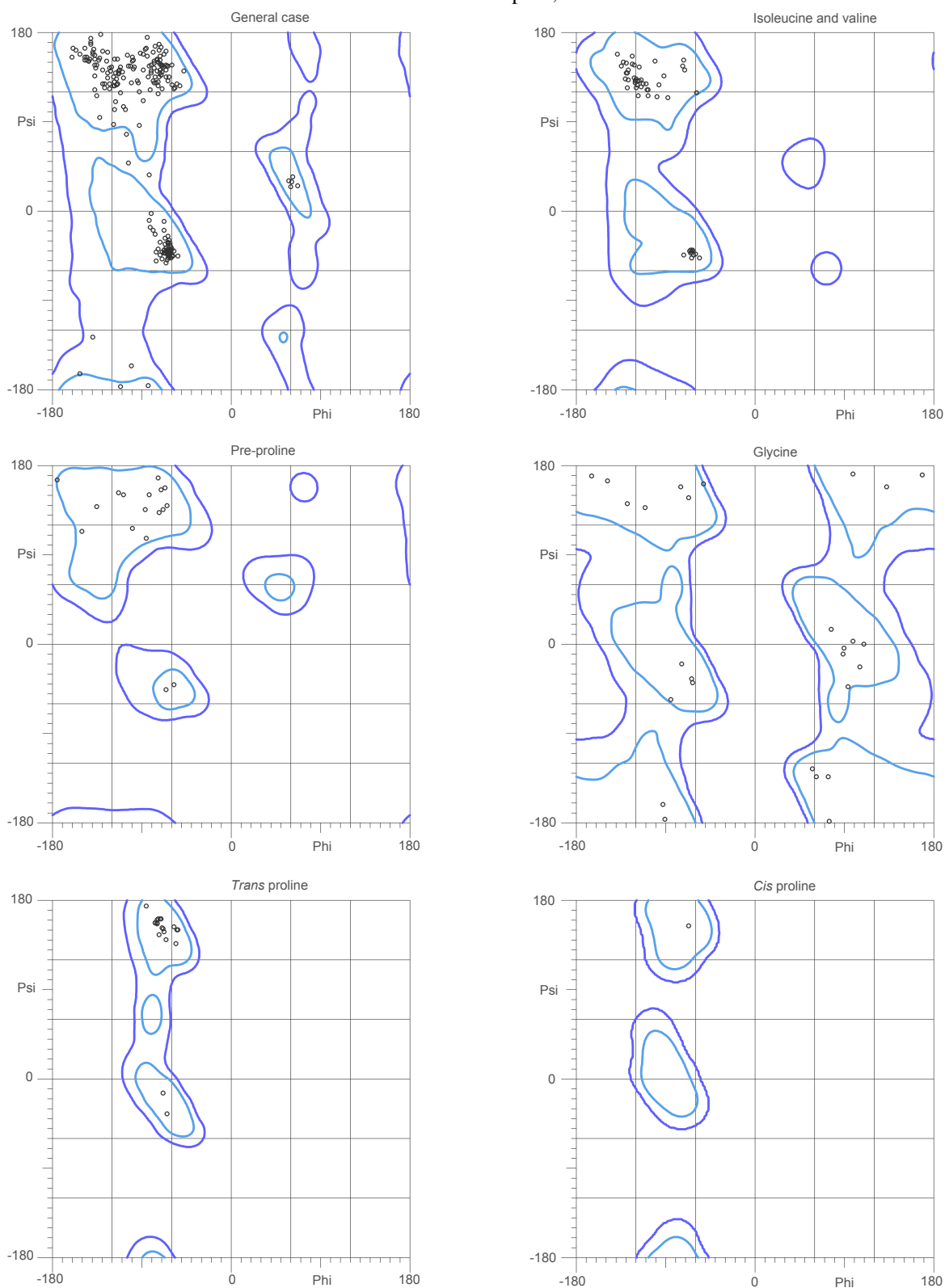


# MolProbity Ramachandran analysis

1a2oA.B99990004.pdb, model 1



97,7% (337/345) of all residues were in favored (98%) regions.

100,0% (345/345) of all residues were in allowed (>99.8%) regions.

There were no outliers.