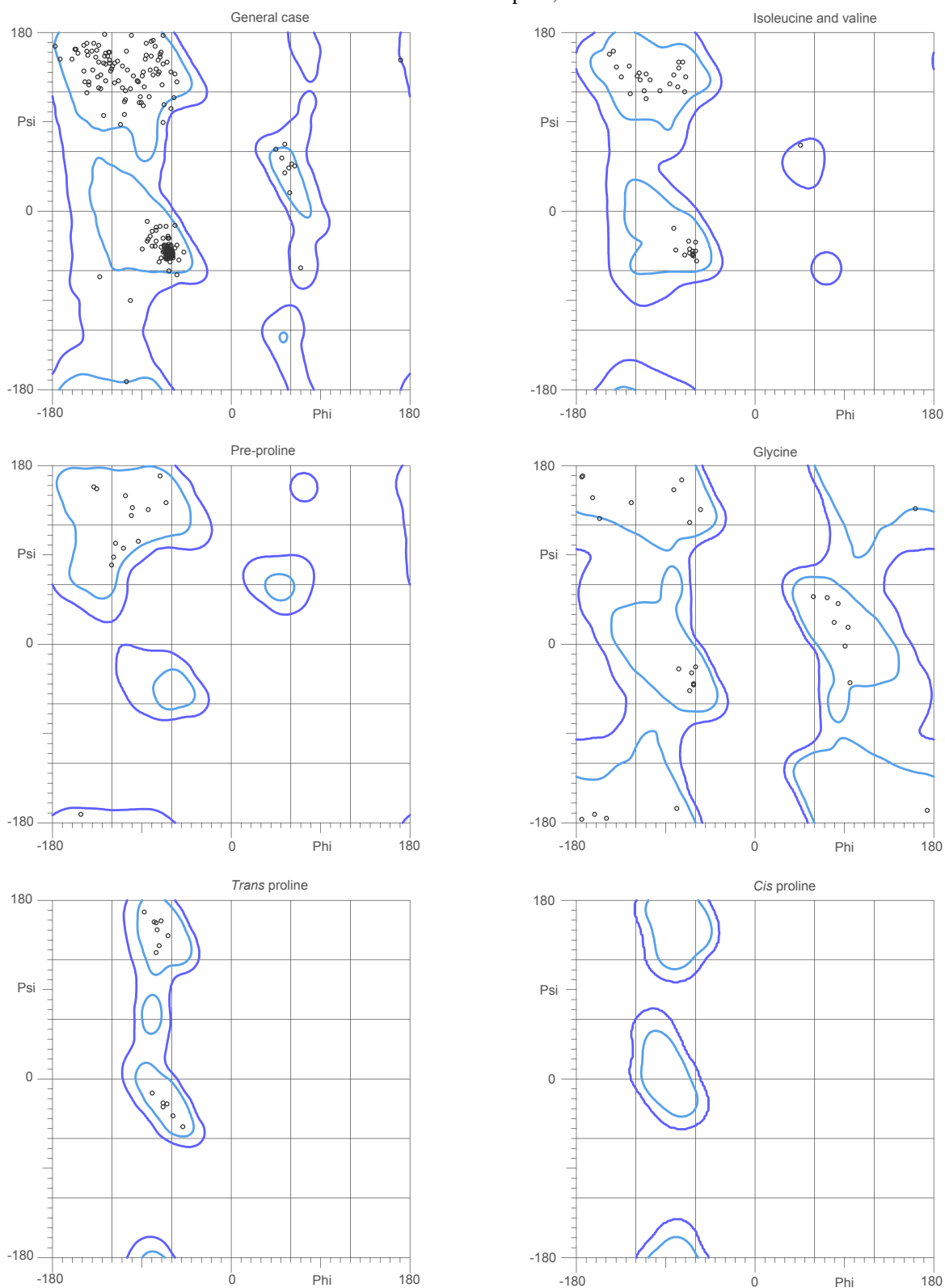


# MolProbity Ramachandran analysis

12asA.B99990043.pdb, model 1



96,0% (312/325) of all residues were in favored (98%) regions.

100,0% (325/325) of all residues were in allowed (>99.8%) regions.

There were no outliers.