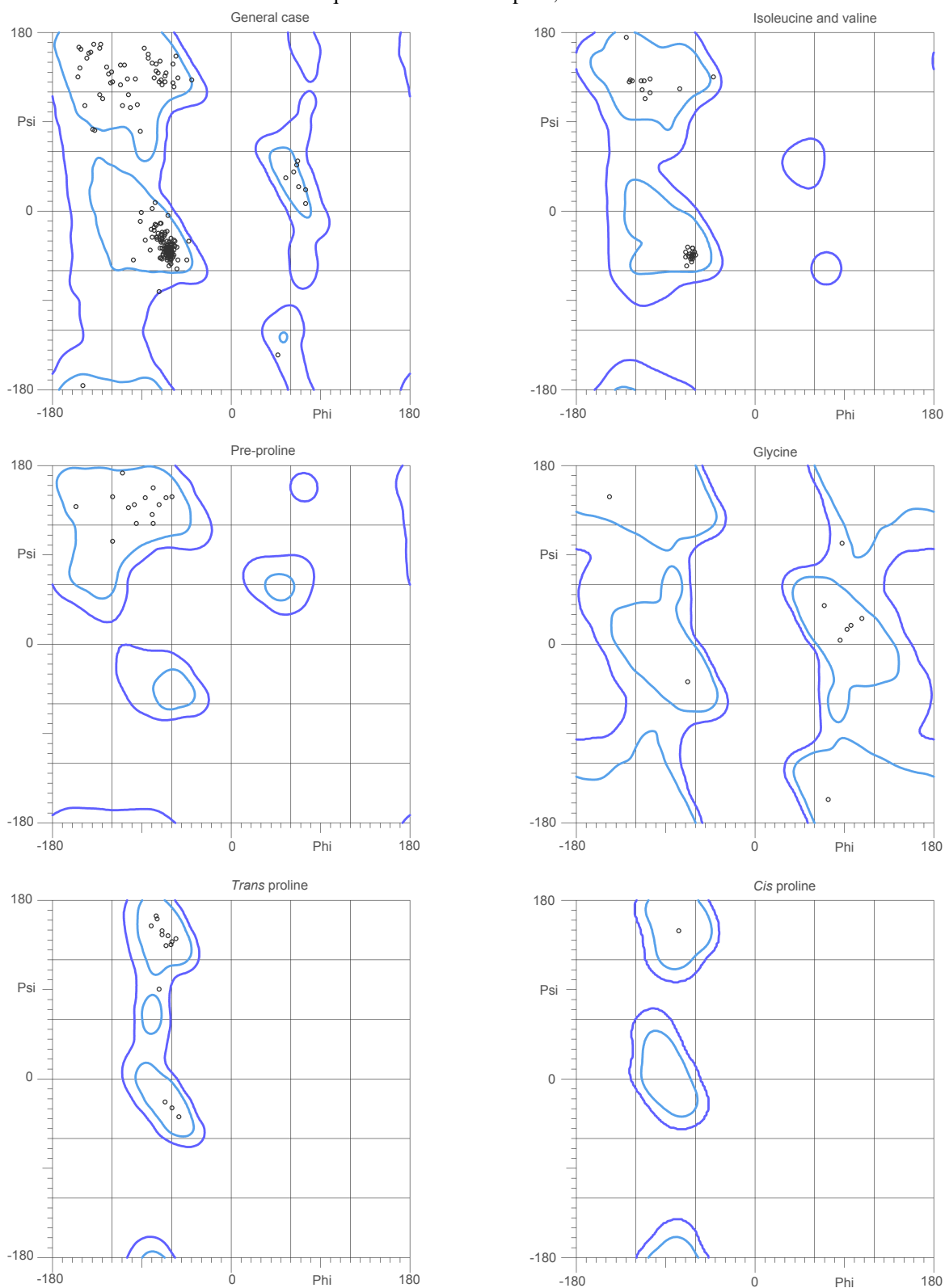


# MolProbity Ramachandran analysis

1aquA.B99990021.pdb, model 1



95,7% (267/279) of all residues were in favored (98%) regions.

100,0% (279/279) of all residues were in allowed (>99.8%) regions.

There were no outliers.