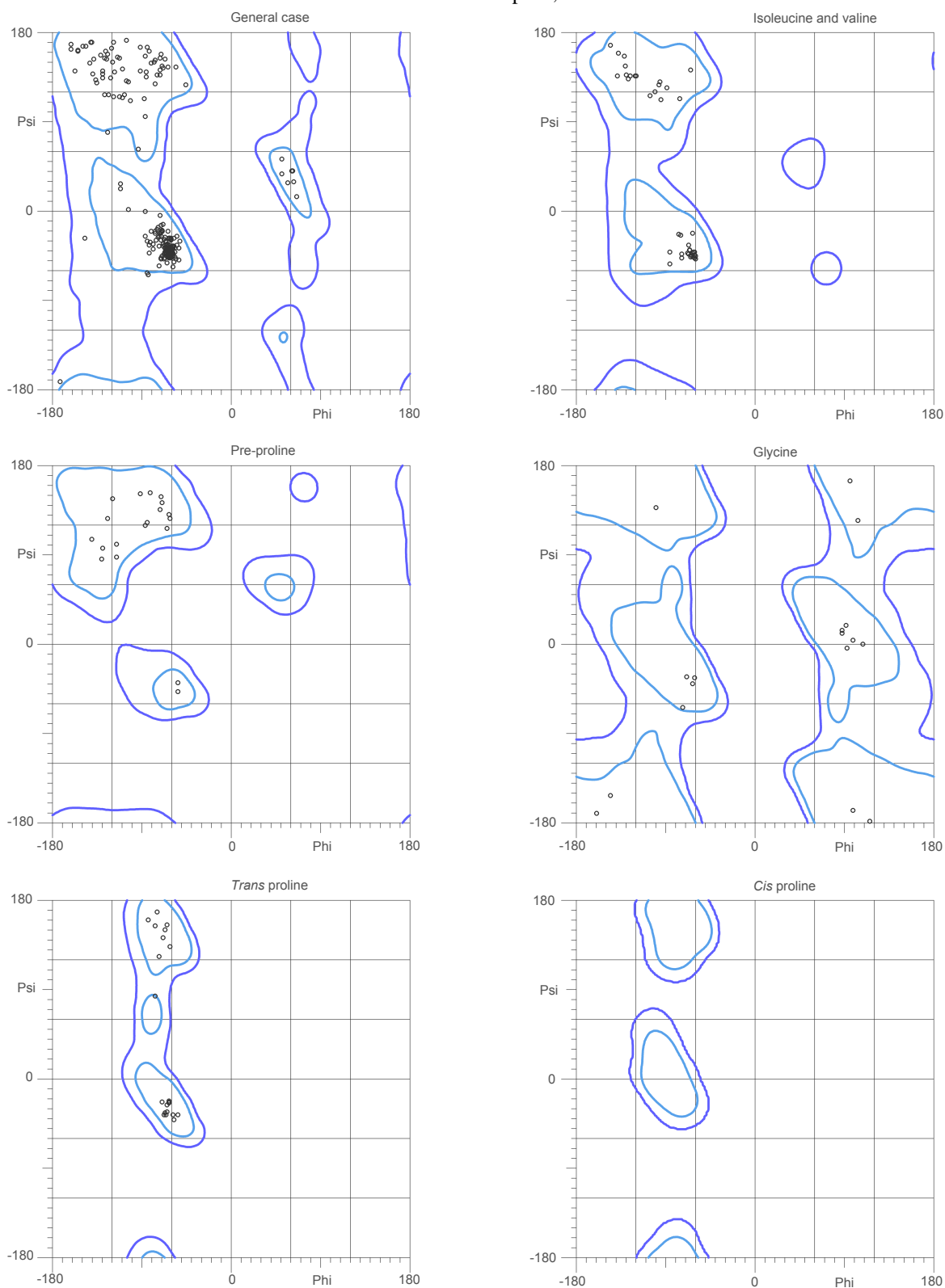# MolProbity Ramachandran analysis

1a99A.B99990027.pdb, model 1

98,2% (333/339) of all residues were in favored (98%) regions.

100,0% (339/339) of all residues were in allowed (>99.8%) regions.

There were no outliers.