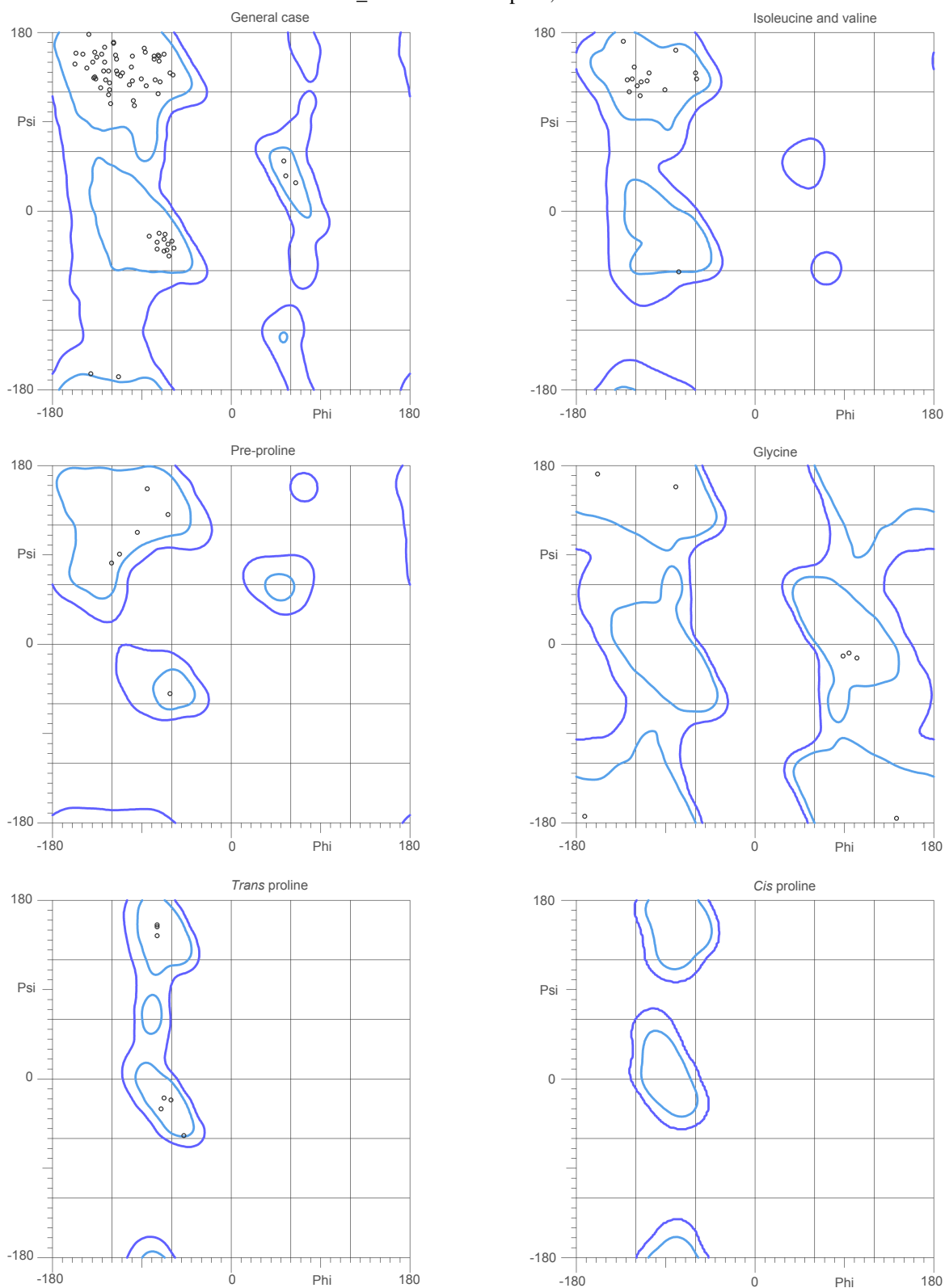# MolProbity Ramachandran analysis

1aac_.B99990022.pdb, model 1

97,1% (100/103) of all residues were in favored (98%) regions.

100,0% (103/103) of all residues were in allowed (>99.8%) regions.

There were no outliers.