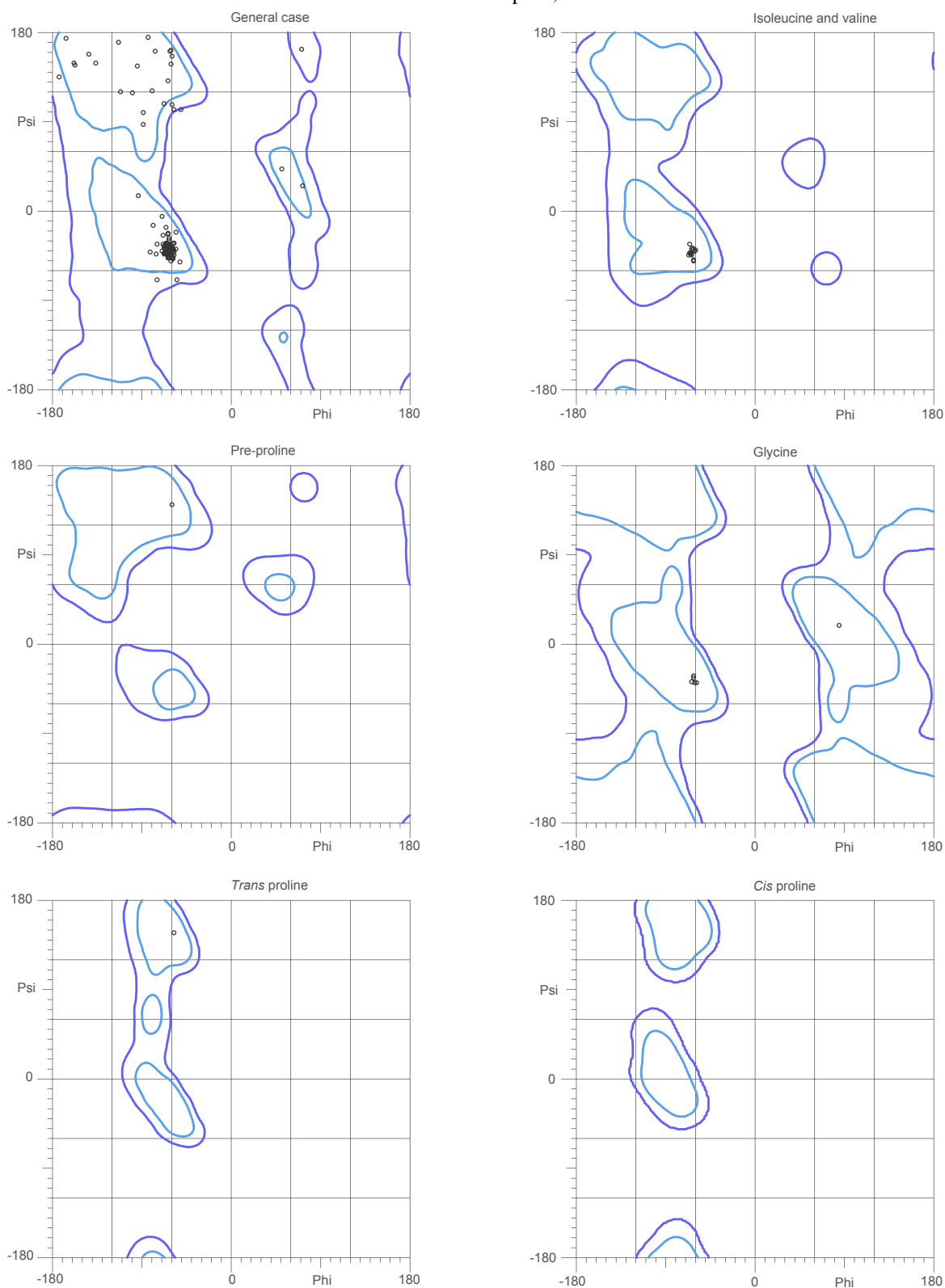


# MolProbity Ramachandran analysis

1au1A.B99990031.pdb, model 1



95,7% (157/164) of all residues were in favored (98%) regions.

100,0% (164/164) of all residues were in allowed (>99.8%) regions.

There were no outliers.