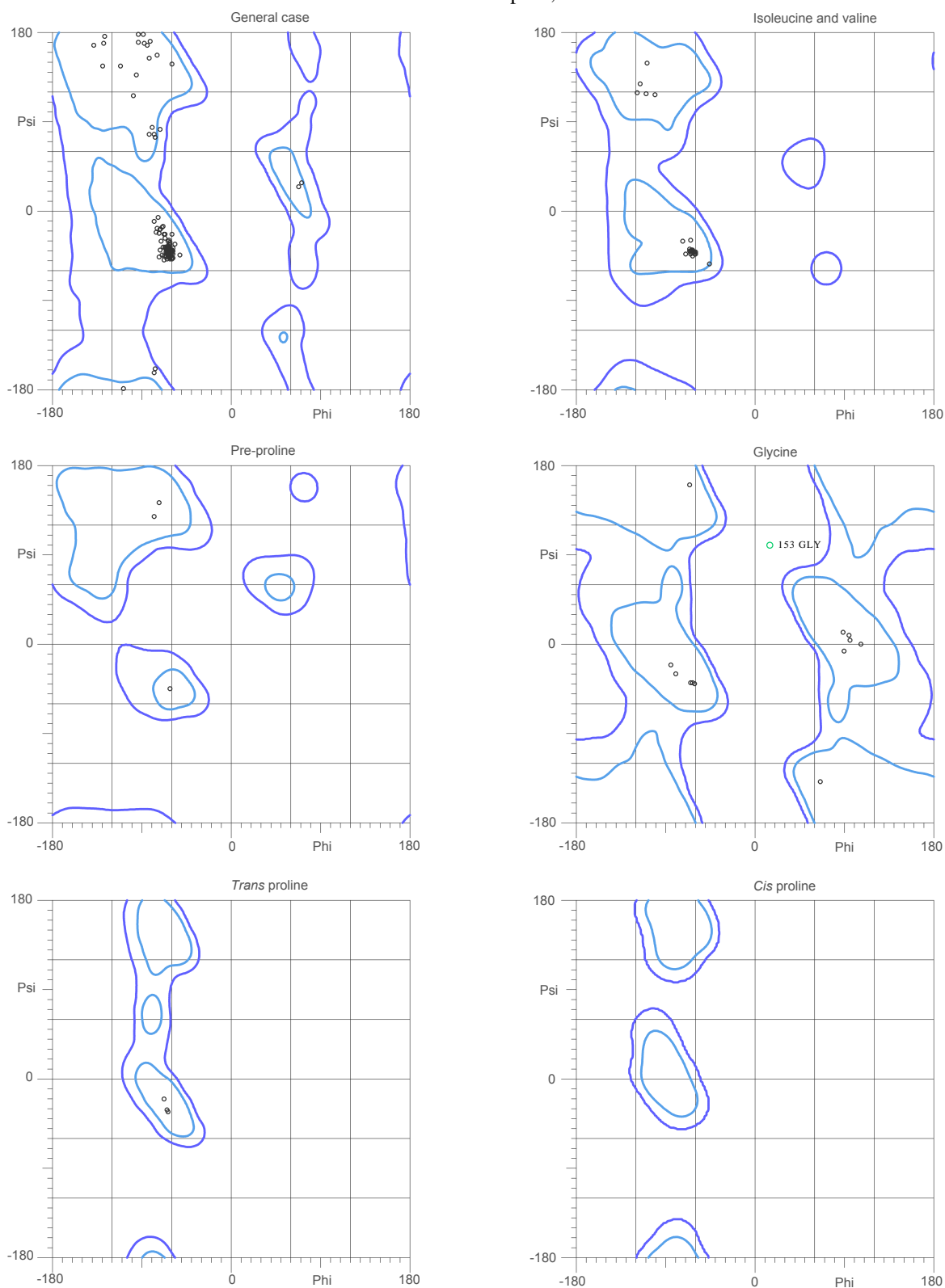


MolProbity Ramachandran analysis

1au1B.B99990029.pdb, model 1



97,5% (159/163) of all residues were in favored (98%) regions.

99,4% (162/163) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

153 GLY (16,0, 100,1)