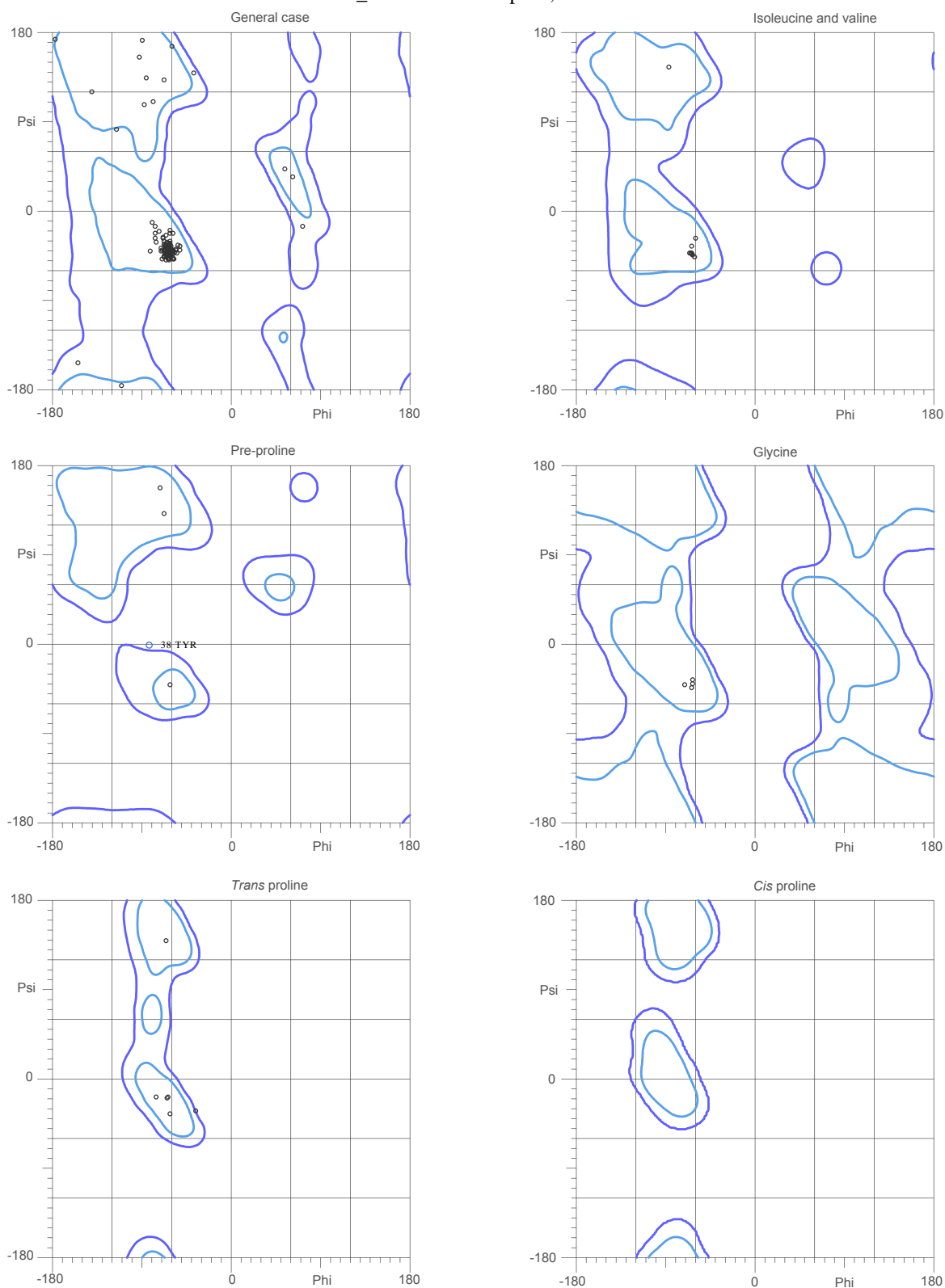


MolProbity Ramachandran analysis

1ash_.B99990037.pdb, model 1



95,9% (139/145) of all residues were in favored (98%) regions.

99,3% (144/145) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

38 TYR (-83,1, -1,7)