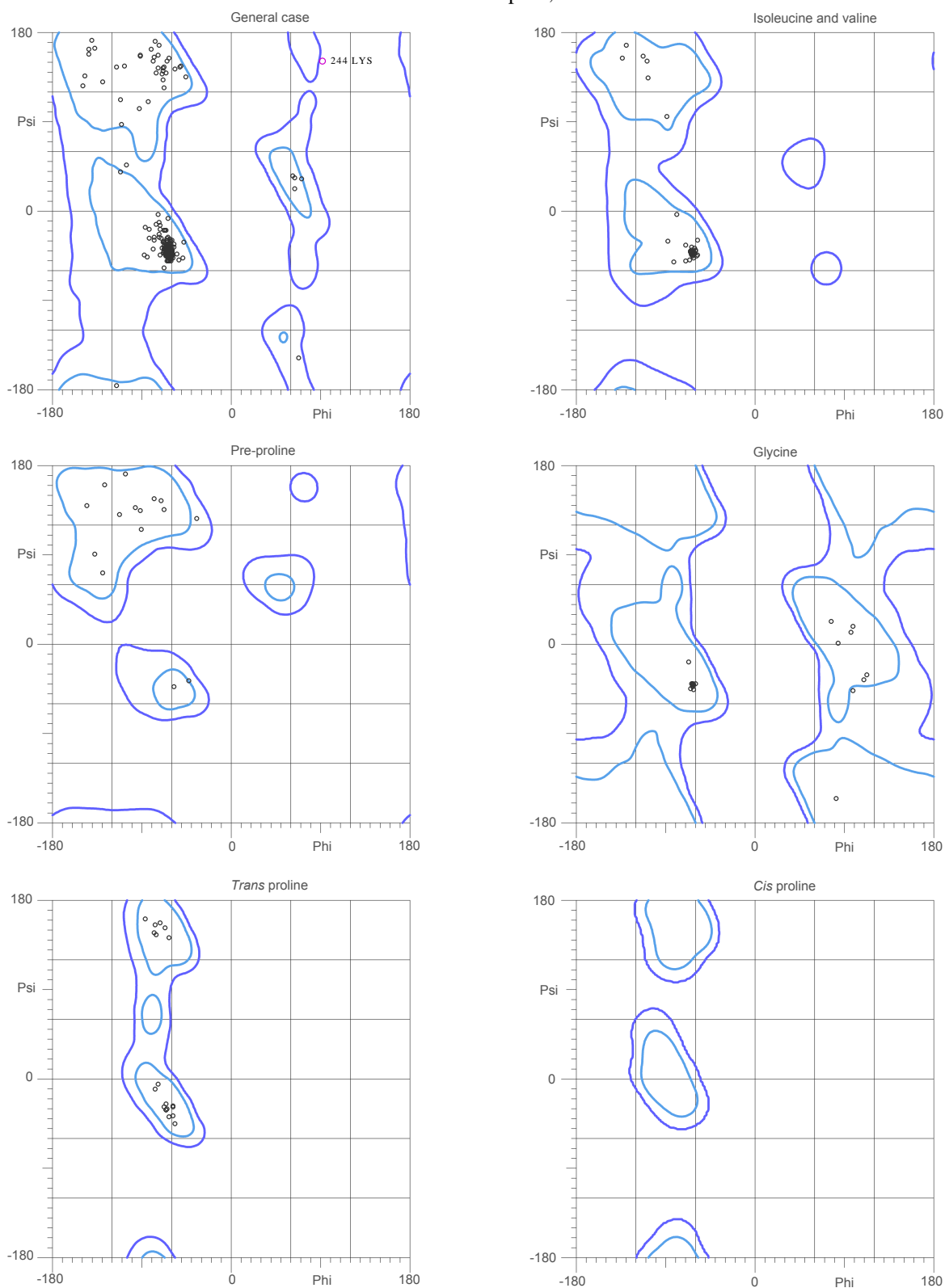


MolProbity Ramachandran analysis

1afrA.B99990033.pdb, model 1



98,3% (337/343) of all residues were in favored (98%) regions.

99,7% (342/343) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

244 LYS (92,2, 152,5)