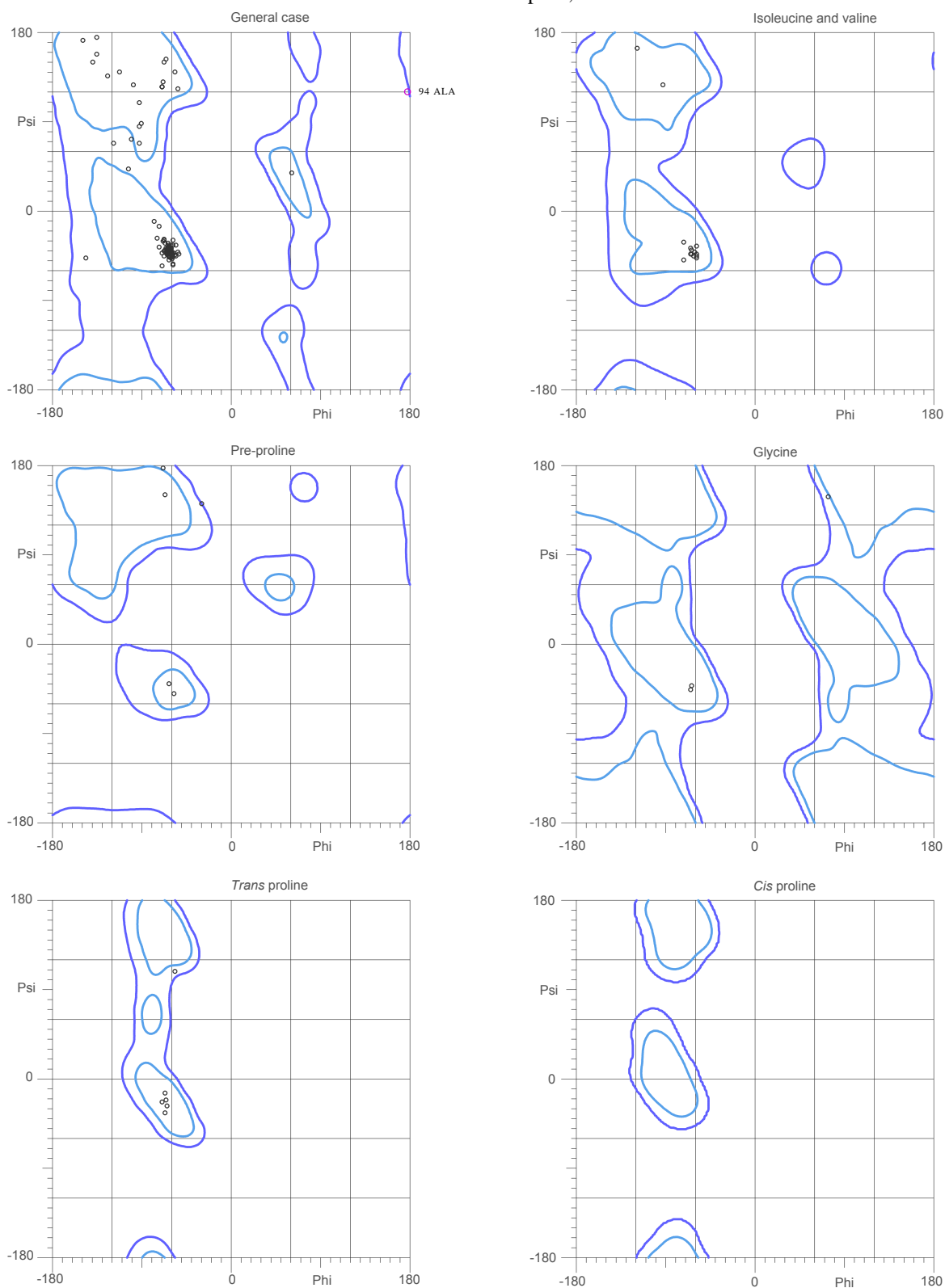


MolProbity Ramachandran analysis

T0837.B99990003FH.pdb, model 1



92.9% (117/126) of all residues were in favored (98%) regions.

99.2% (125/126) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

94 ALA (178.1, 121.7)