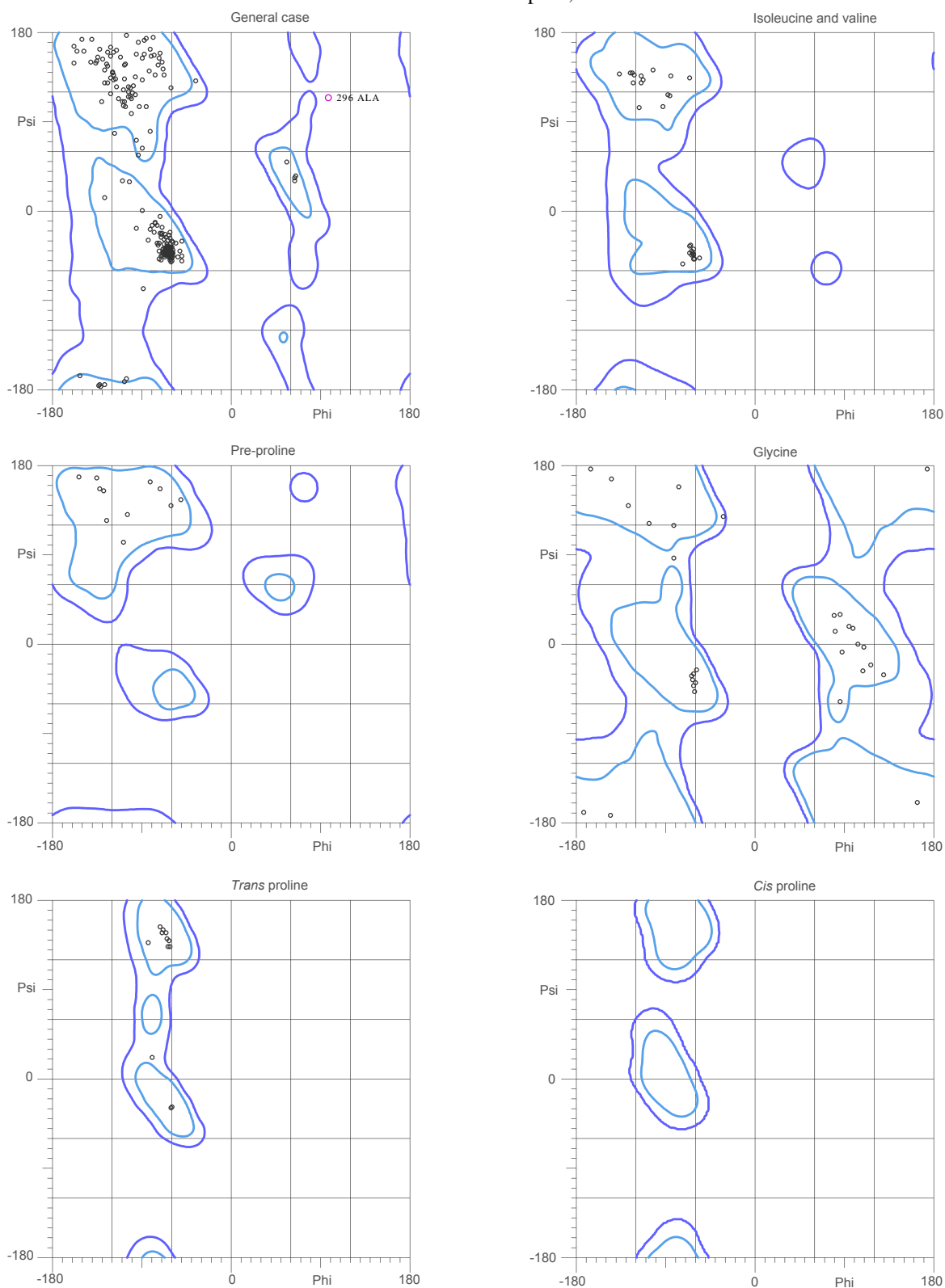


MolProbity Ramachandran analysis

T0694.B99990002FH.pdb, model 1



97.4% (305/313) of all residues were in favored (98%) regions.

99.7% (312/313) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

296 ALA (98.1, 115.6)