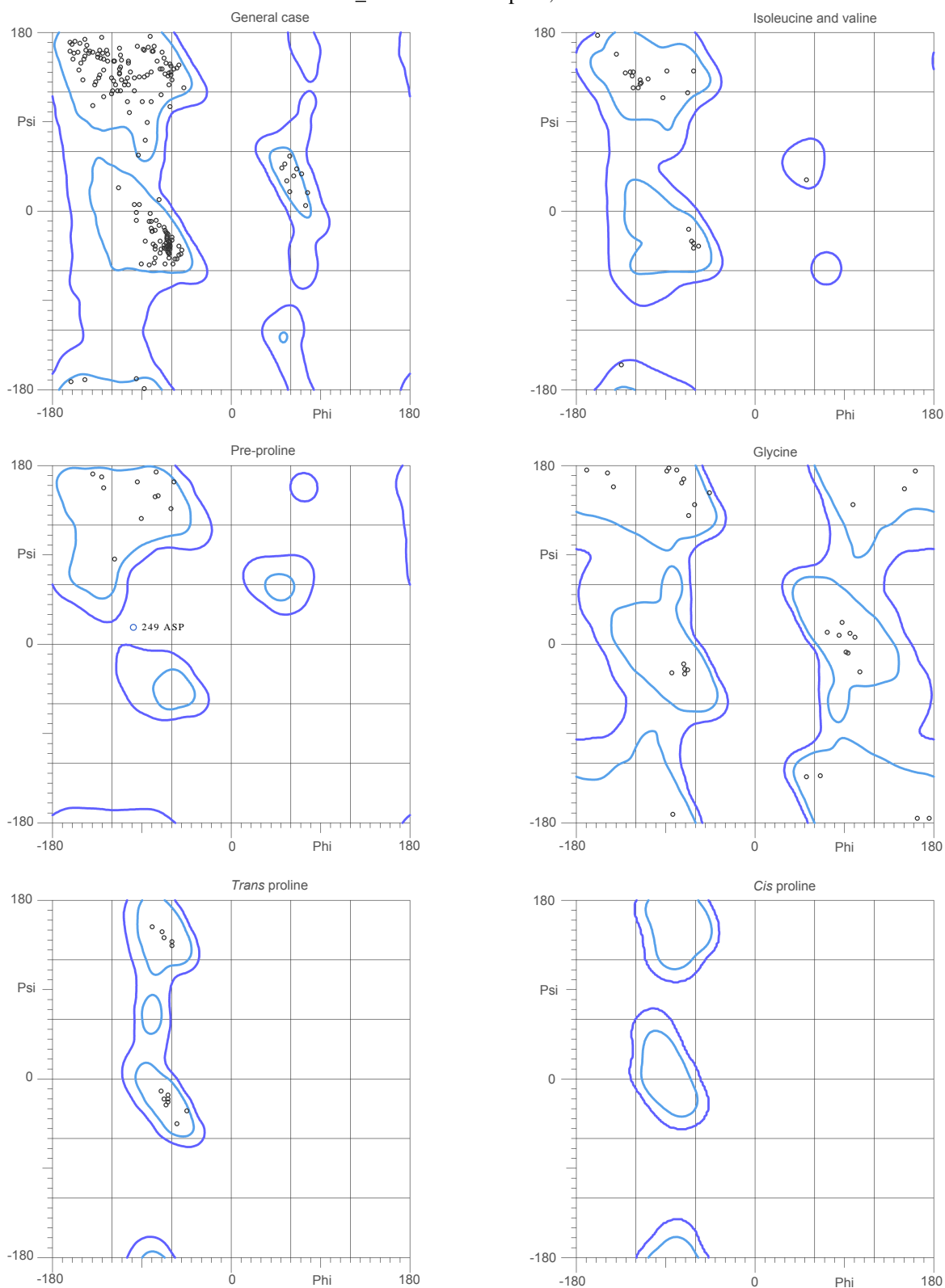


MolProbity Ramachandran analysis

1arb_.B99990050.pdb, model 1



96,2% (251/261) of all residues were in favored (98%) regions.

99,6% (260/261) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (Φ , Ψ):

249 ASP (-99,1, 17,5)