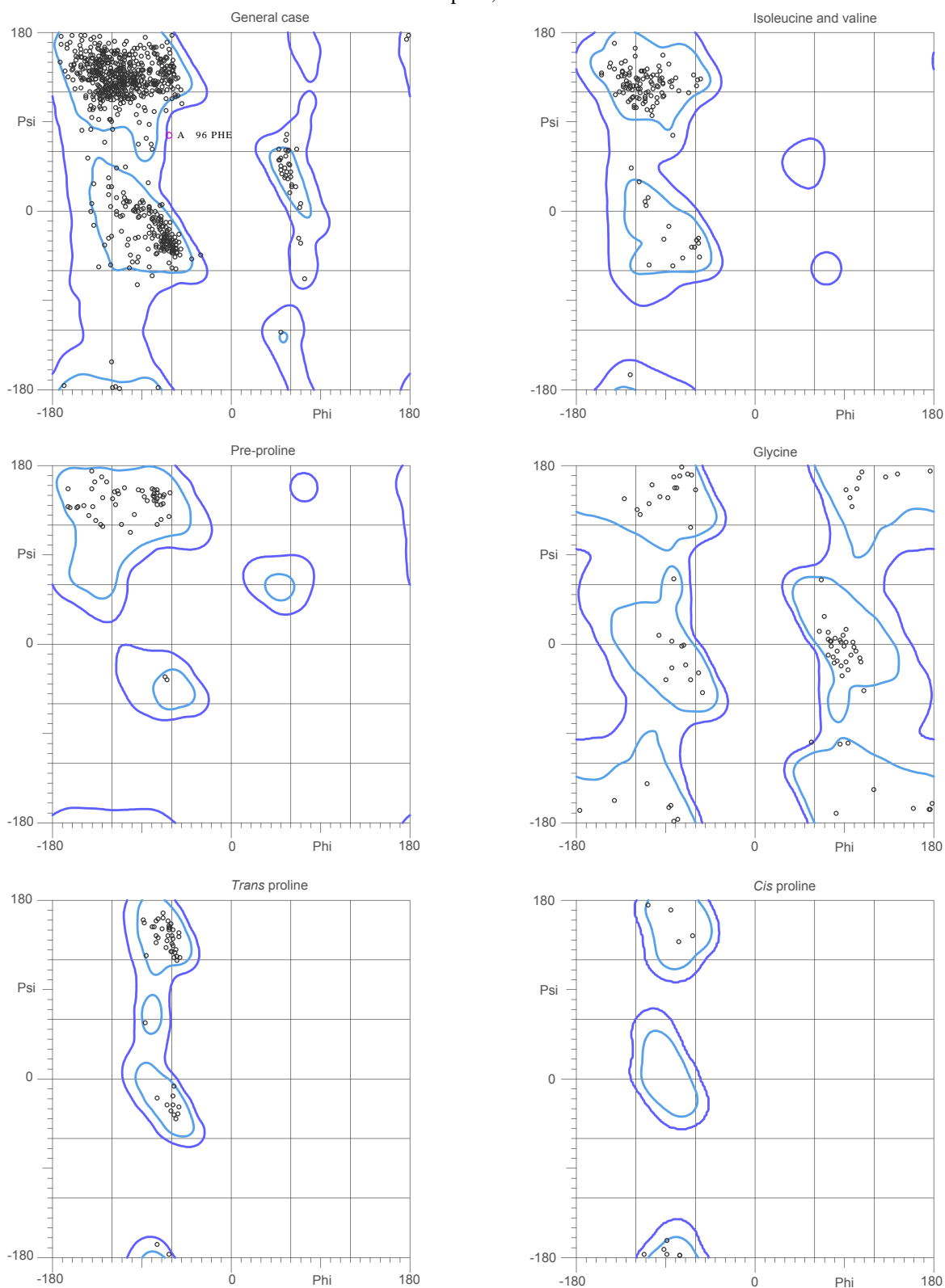


MolProbity Ramachandran analysis

6xc3.H.pdb, model 1



96.3% (1012/1051) of all residues were in favored (98%) regions.

99.9% (1050/1051) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

A 96 PHE (-63.4, 77.8)