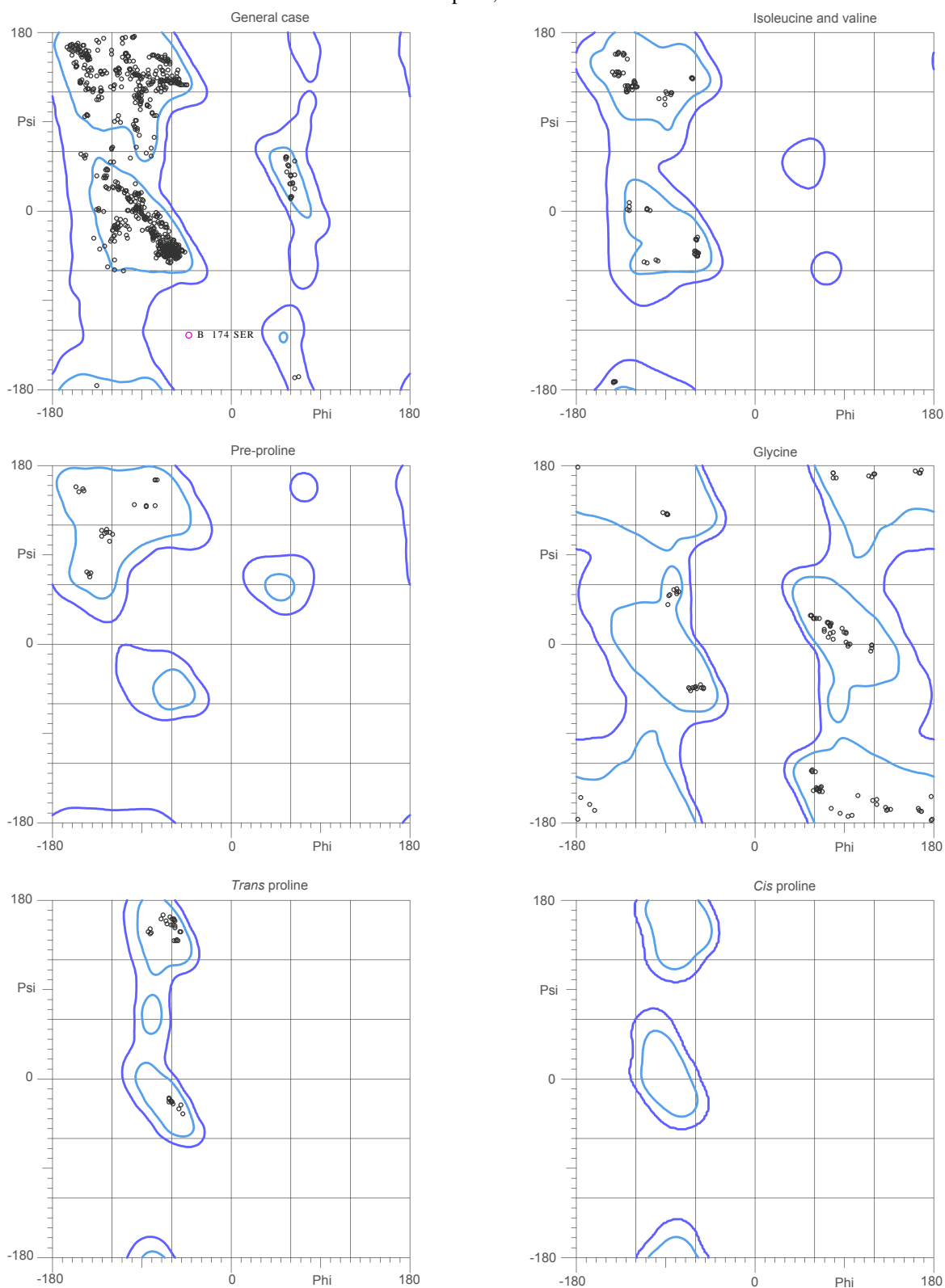


MolProbity Ramachandran analysis

3hha.H.pdb, model 1



97.5% (940/964) of all residues were in favored (98%) regions.

99.9% (963/964) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

B 174 SER (-43.6, -125.7)