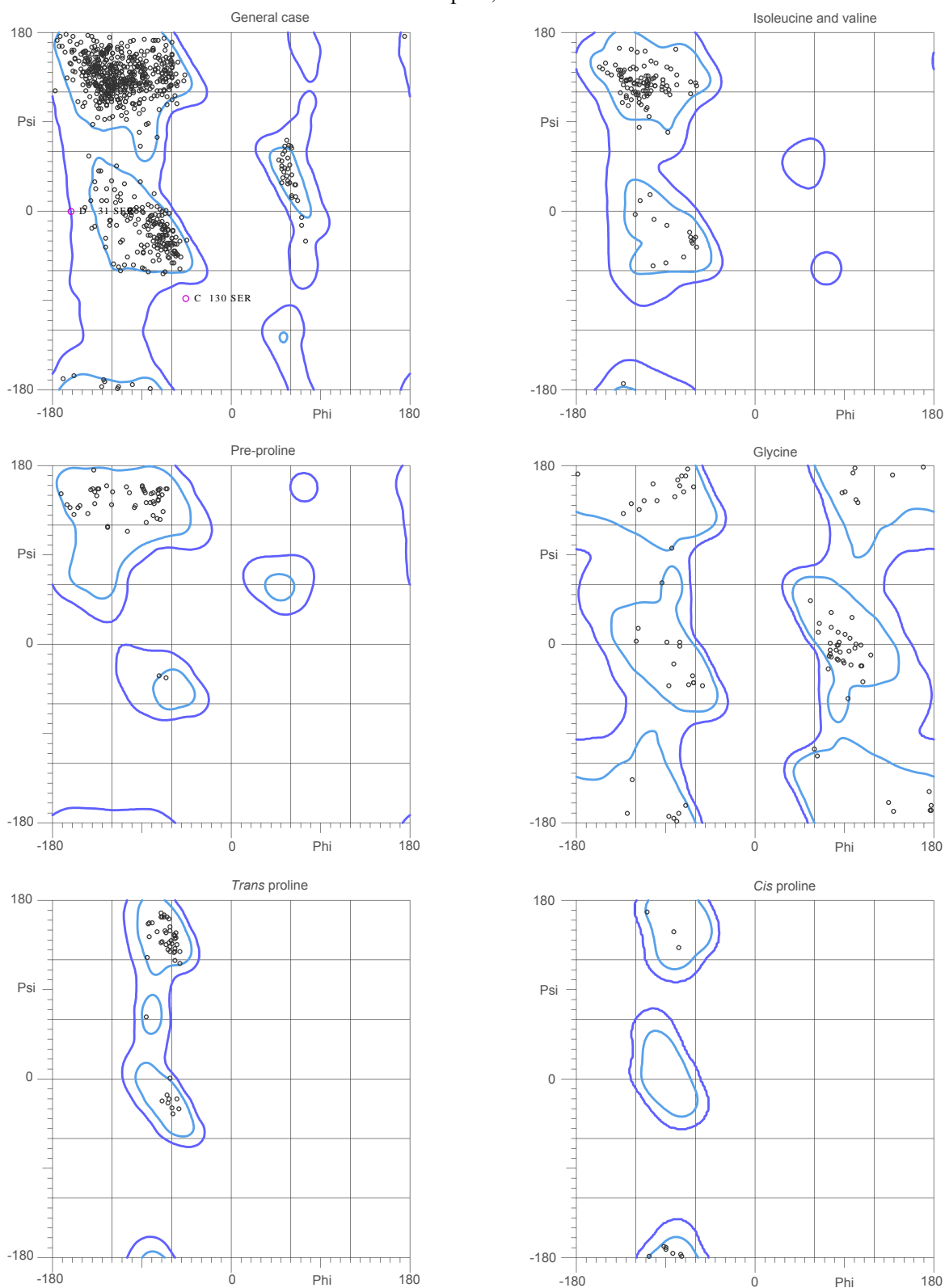


MolProbity Ramachandran analysis

6xc7.H.pdb, model 1



96.0% (1009/1051) of all residues were in favored (98%) regions.

99.8% (1049/1051) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (ϕ , ψ):

C 130 SER (-46.8, -88.0)

D 31 SER (-162.1, -0.6)