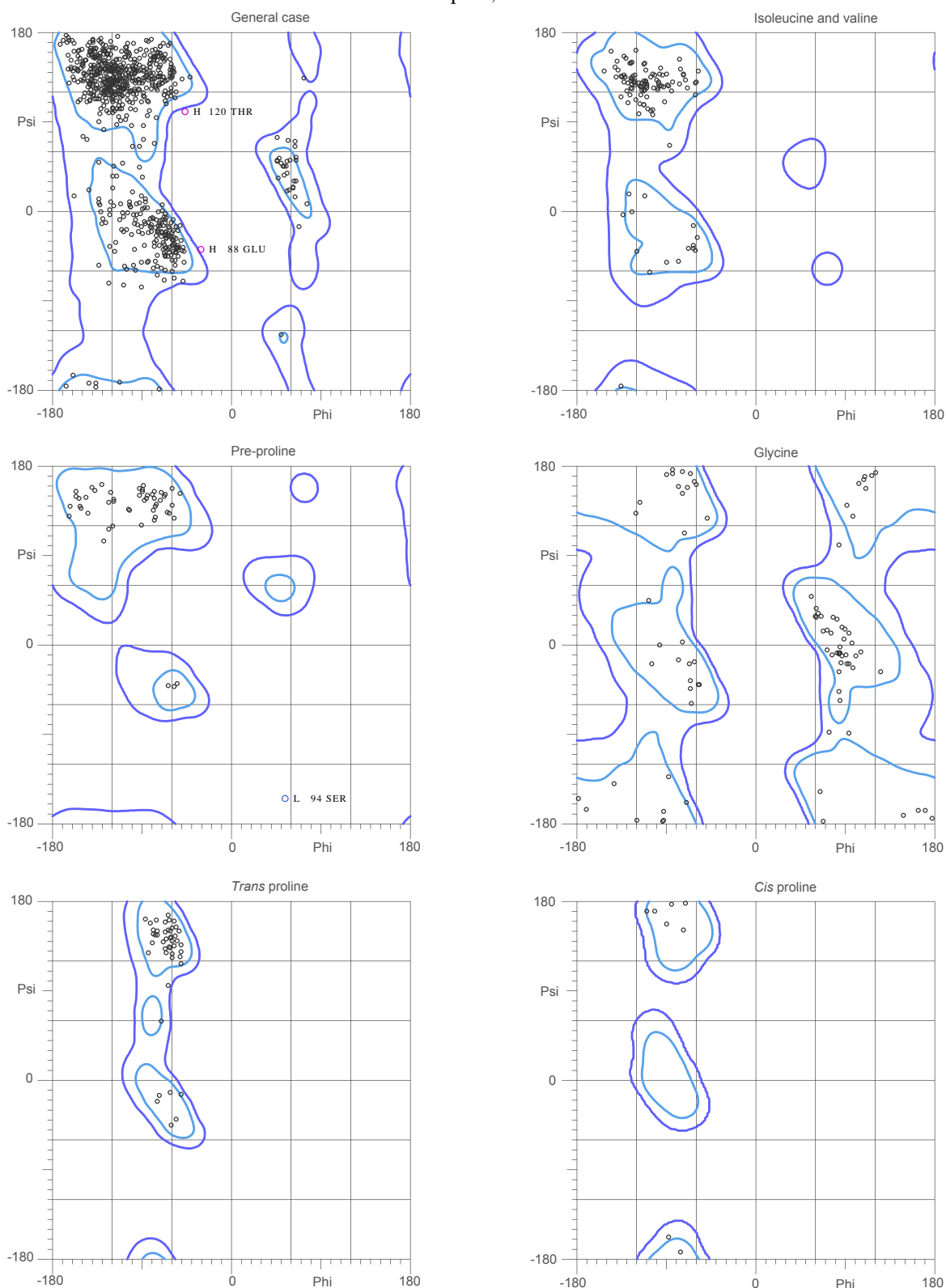


MolProbity Ramachandran analysis

7chc.H.pdb, model 1



95.6% (983/1028) of all residues were in favored (98%) regions.
99.7% (1025/1028) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (ϕ , ψ):

- H 88 GLU (-31.1, -38.4)
- H 120 THR (-47.1, 101.4)
- L 94 SER (54.7, -155.3)