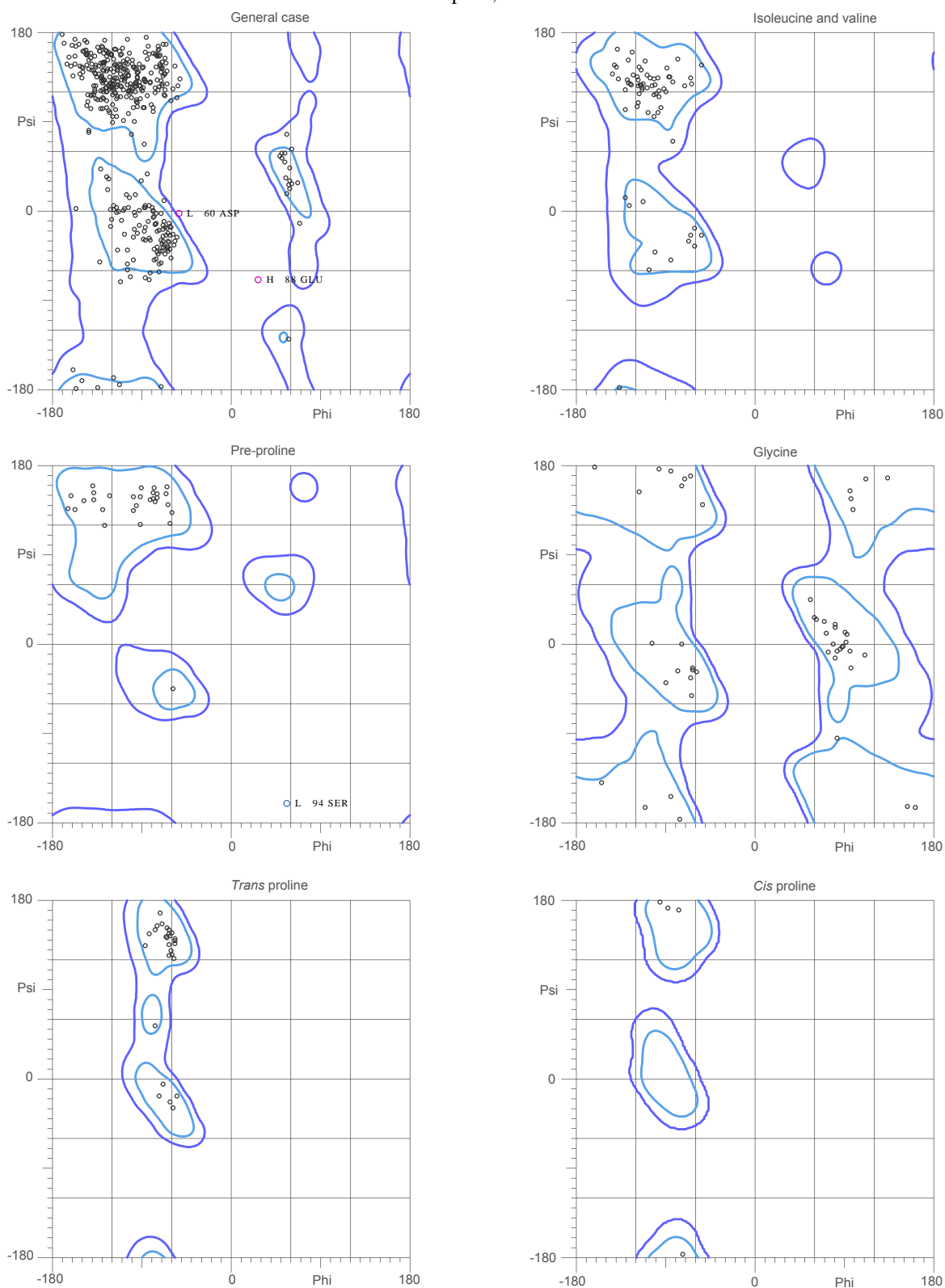


MolProbity Ramachandran analysis

7ch5.H.pdb, model 1



95.3% (571/599) of all residues were in favored (98%) regions.
99.5% (596/599) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi):

H 88 GLU (27.9, -69.1)

L 60 ASP (-53.9, -2.4)

L 94 SER (56.7, -161.9)