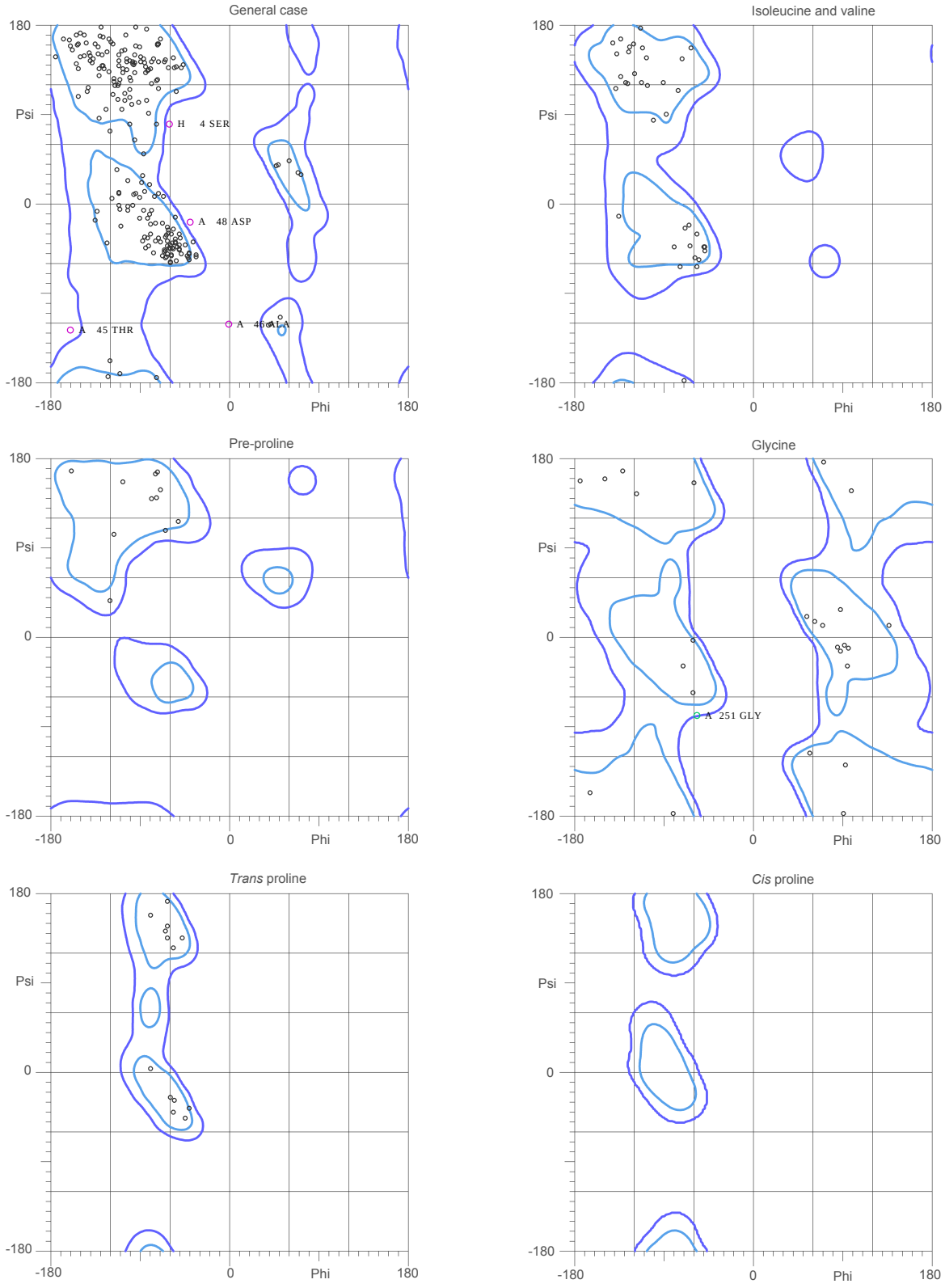
MolProbity Ramachandran analysis

3snc.H.pdb, model 1



89.3% (275/308) of all residues were in favored (98%) regions.
98.4% (303/308) of all residues were in allowed (>99.8%) regions.

There were 5 outliers (phi, psi):
A 45 THR (-161.6, -127.9)
A 46 ALA (-1.8, -121.4)
A 48 ASP (-40.2, -18.2)
A 251 GLY (-57.7, -79.6)
H 4 SER (-61.5, 81.0)