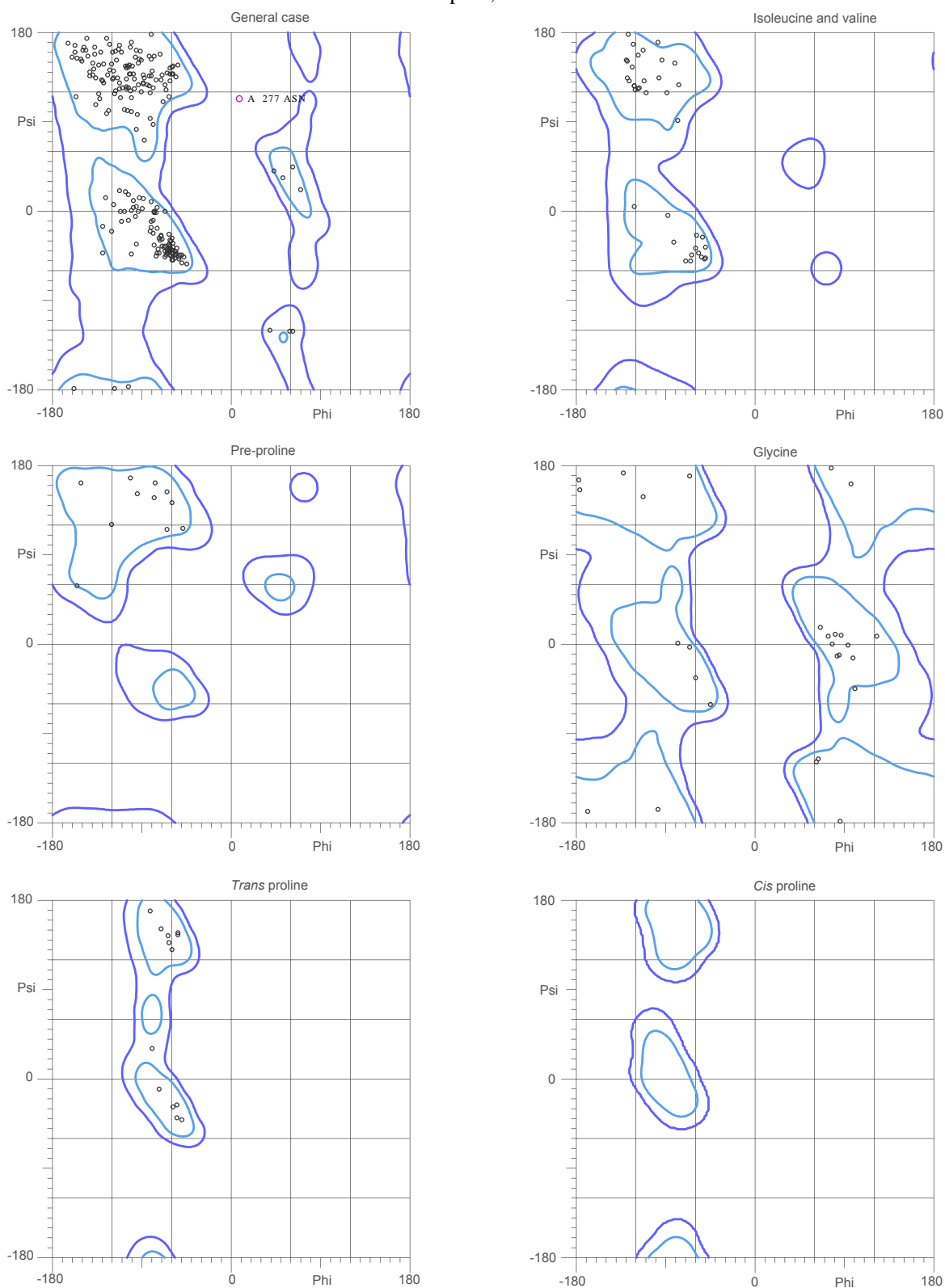


MolProbity Ramachandran analysis

2z3c.H.pdb, model 1



97.1% (303/312) of all residues were in favored (98%) regions.

99.7% (311/312) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

A 277 ASN (9.0, 114.5)