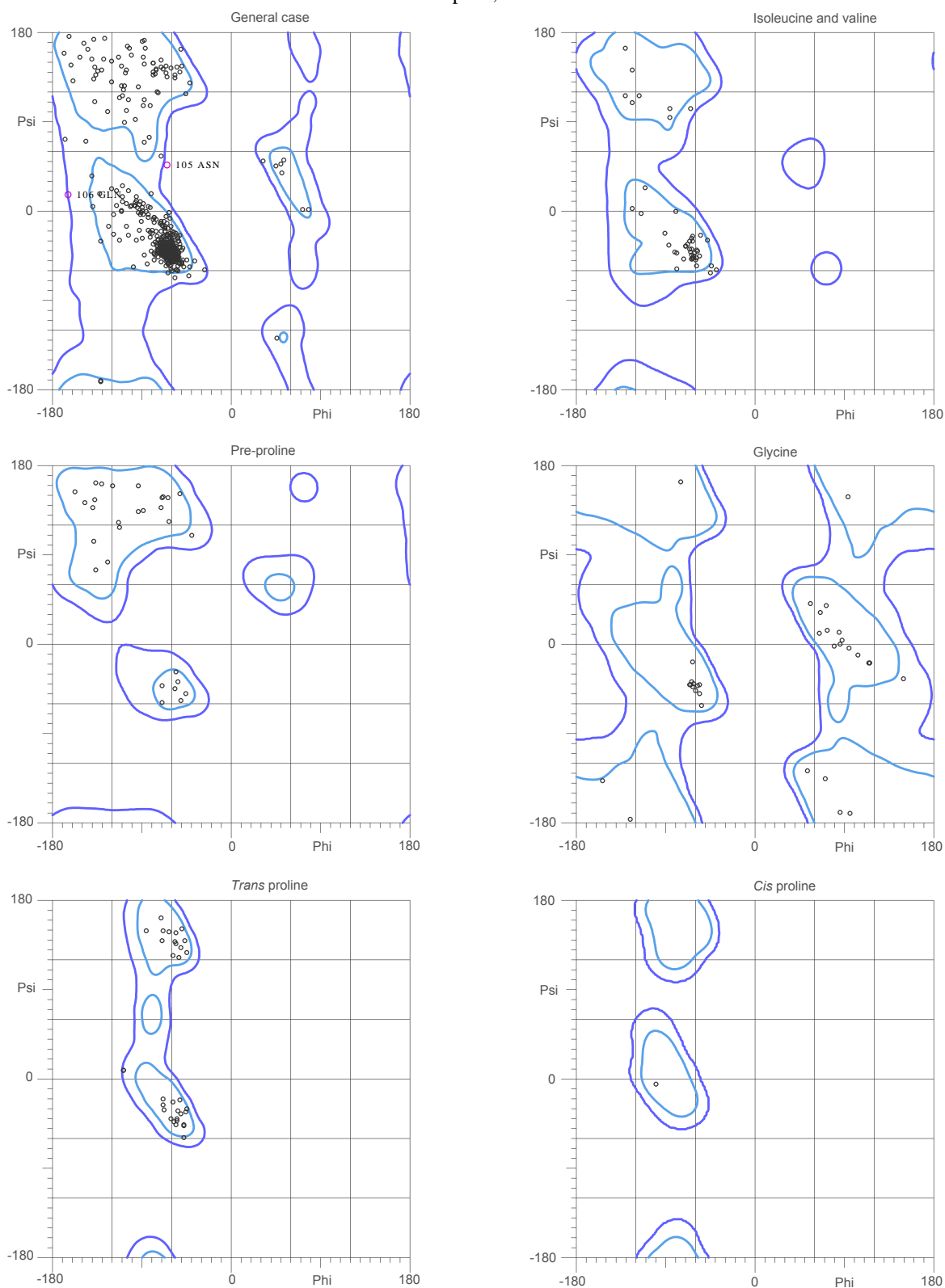


MolProbity Ramachandran analysis

3bkk.H.pdb, model 1



95.6% (581/608) of all residues were in favored (98%) regions.

99.7% (606/608) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (ϕ , ψ):

105 ASN (-65.9, 47.3)

106 GLN (-165.6, 17.4)