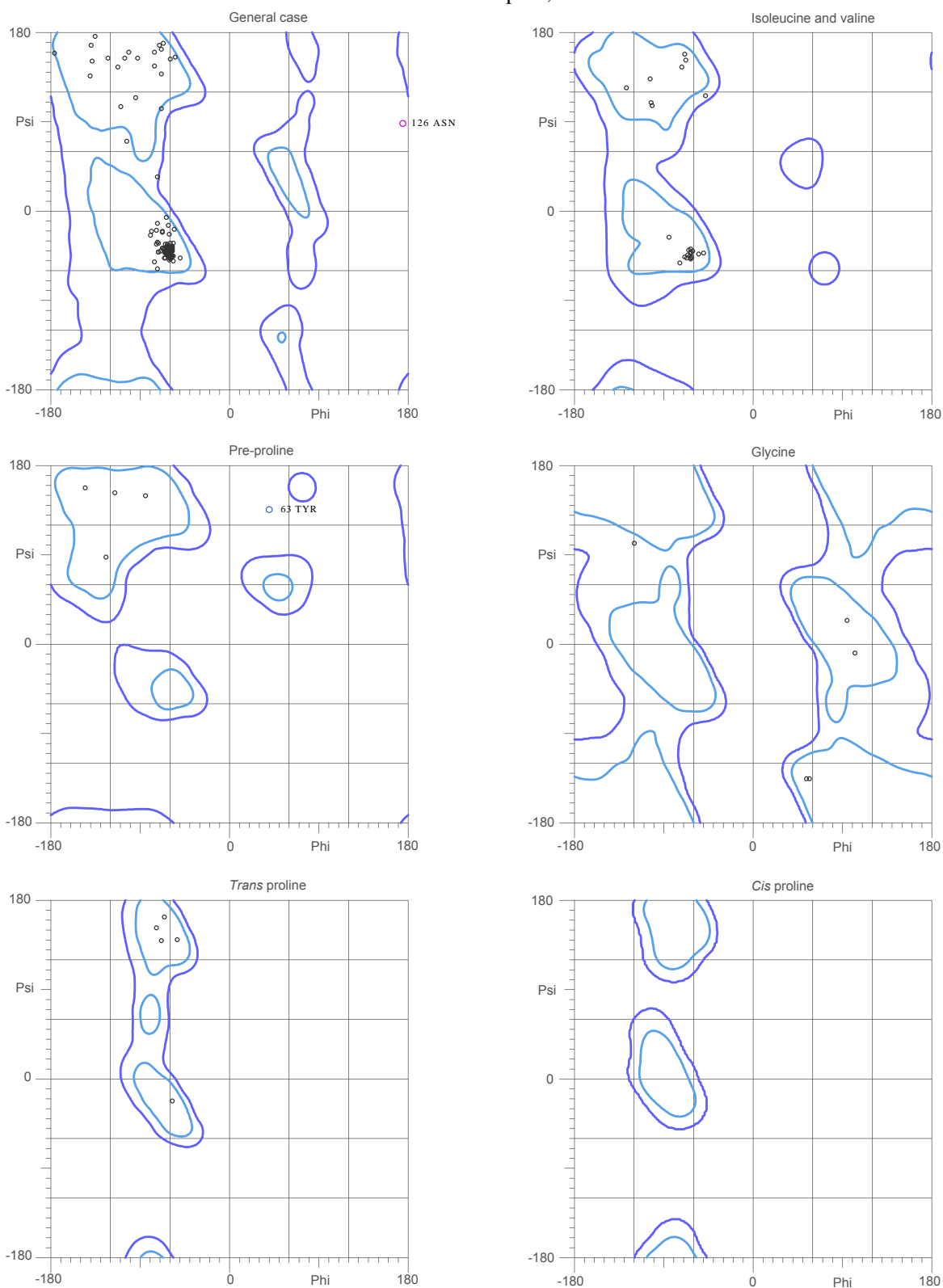


MolProbity Ramachandran analysis

1a101.B99990039.pdb, model 1



95,0% (134/141) of all residues were in favored (98%) regions.

98,6% (139/141) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (ϕ , ψ):

63 TYR (40,2, 136,2)

126 ASN (175,3, 89,5)