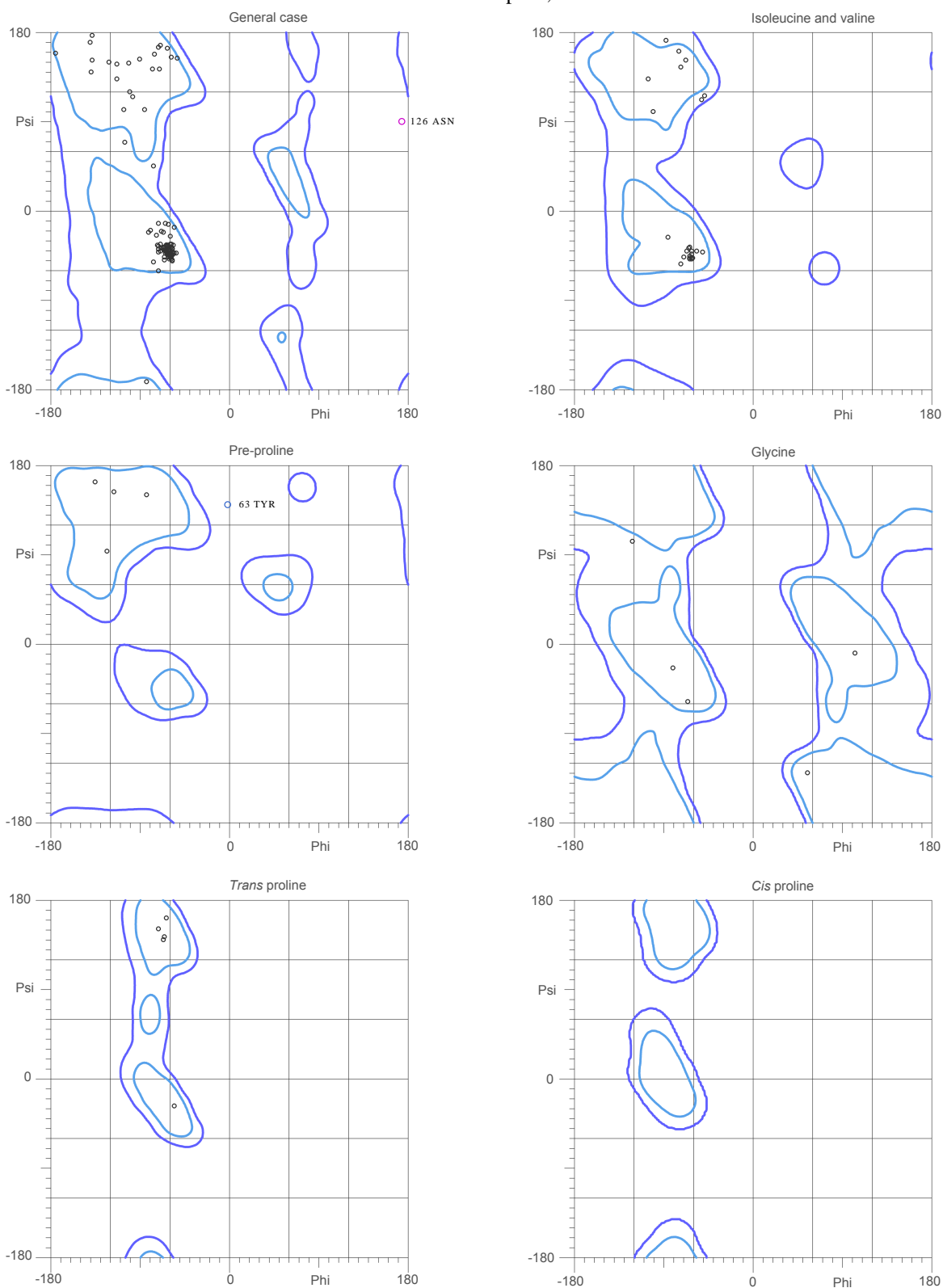


MolProbity Ramachandran analysis

1a101.B99990021.pdb, model 1



93,6% (132/141) 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 (-2,0, 141,2)

126 ASN (174,8, 91,2)