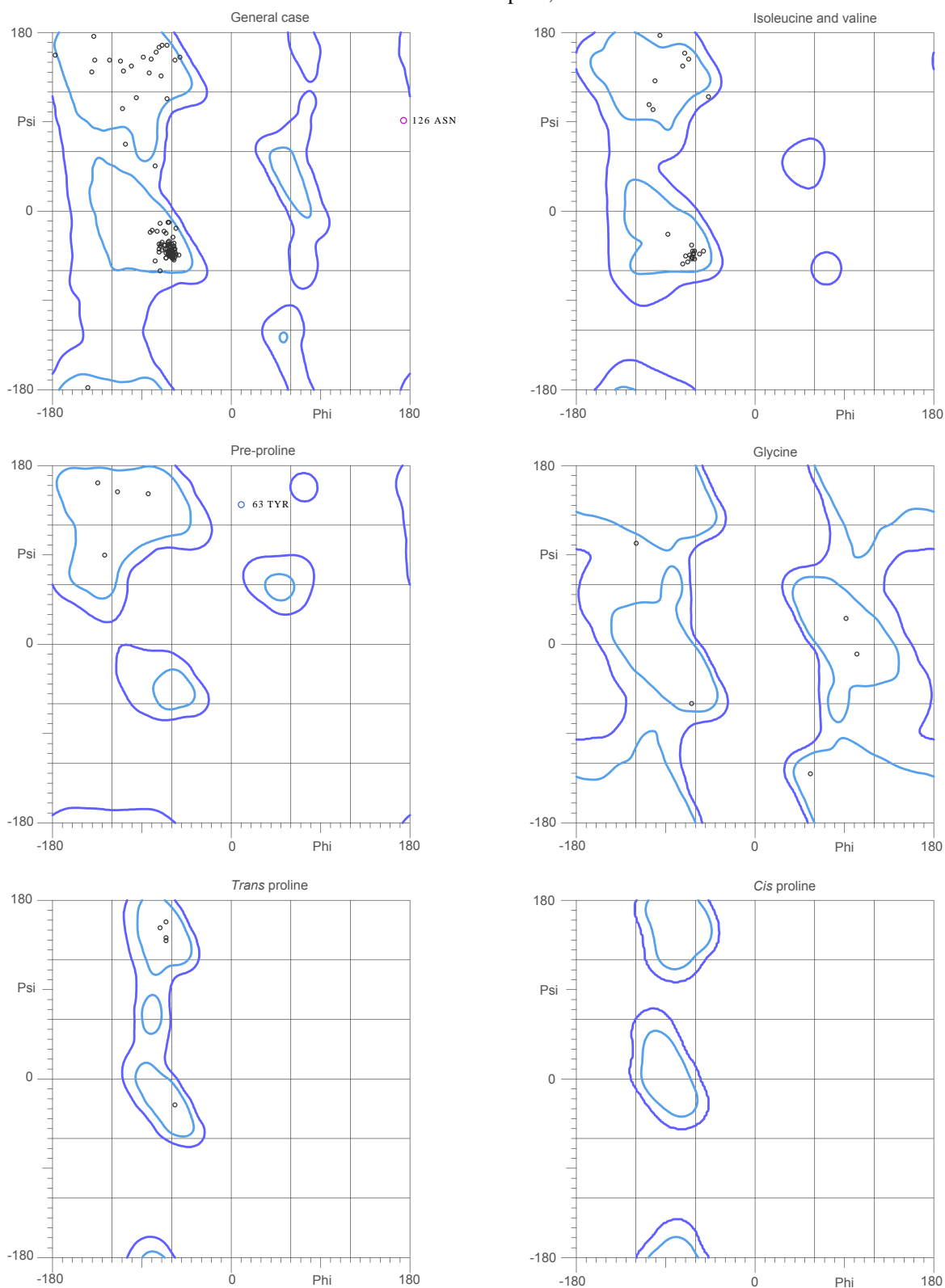


MolProbity Ramachandran analysis

1a101.B99990042.pdb, model 1



94,3% (133/141) 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 (10,1, 142,0)

126 ASN (174,9, 92,9)