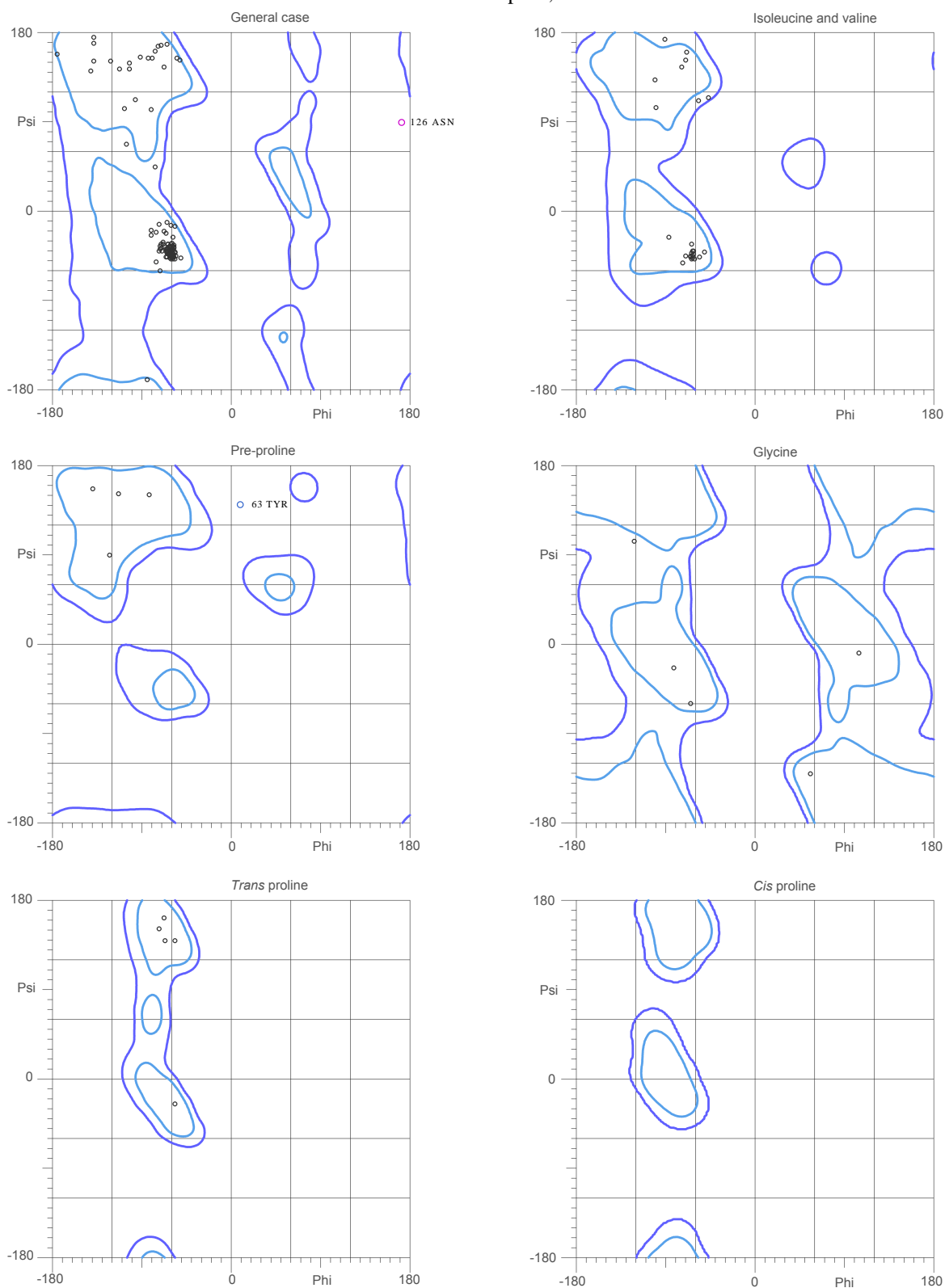


# MolProbity Ramachandran analysis

1a101.B99990005.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 ( $\Phi$ ,  $\Psi$ ):

63 TYR (9,4, 141,3)

126 ASN (172,4, 90,1)