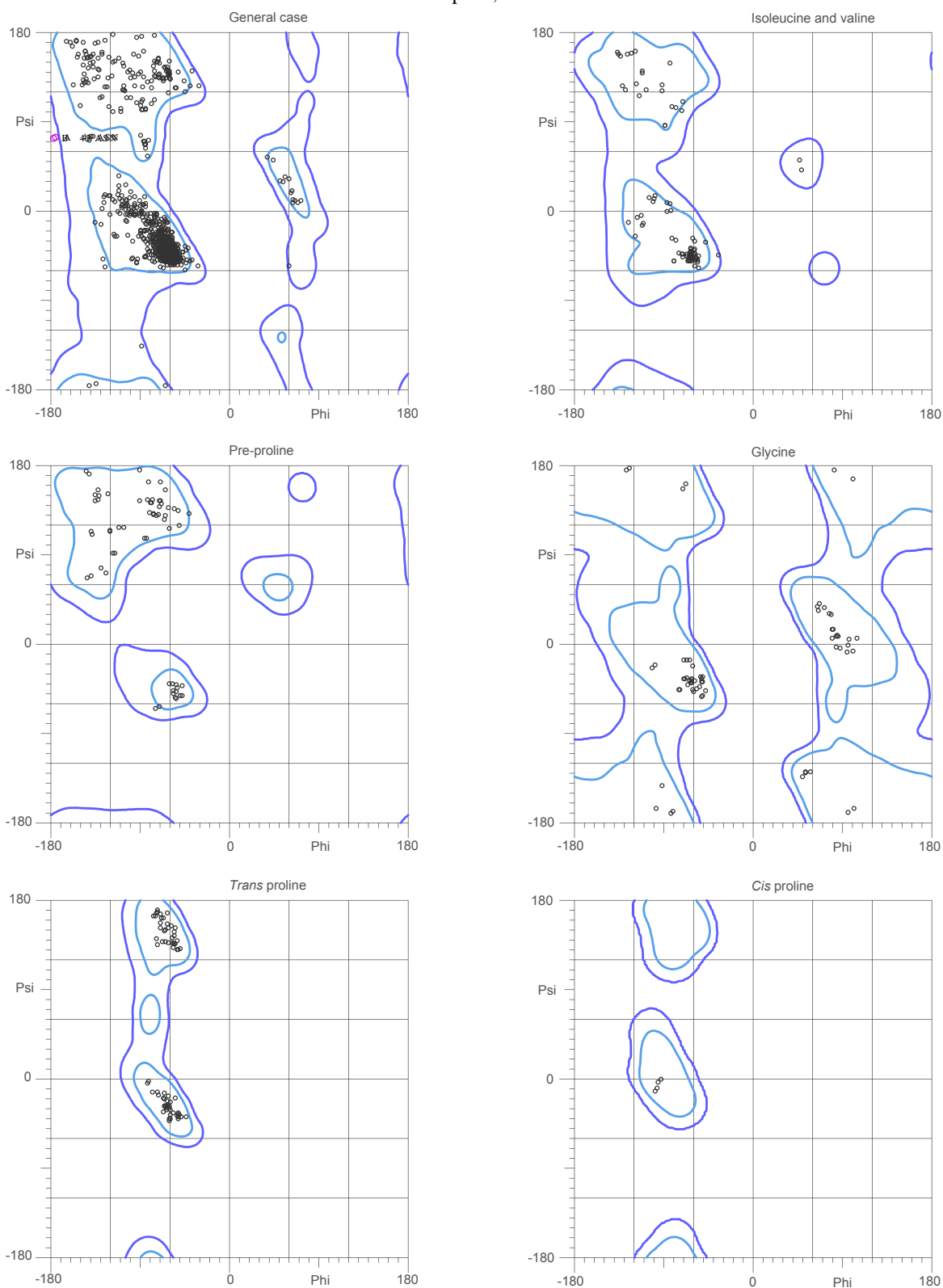


# MolProbity Ramachandran analysis

5amb.H.pdb, model 1



98.0% (1184/1208) of all residues were in favored (98%) regions.

99.8% (1206/1208) of all residues were in allowed (>99.8%) regions.

There were 2 outliers ( $\phi$ ,  $\psi$ ):

A 45 ASN (-176.8, 75.2)

B 45 ASN (-178.4, 74.4)