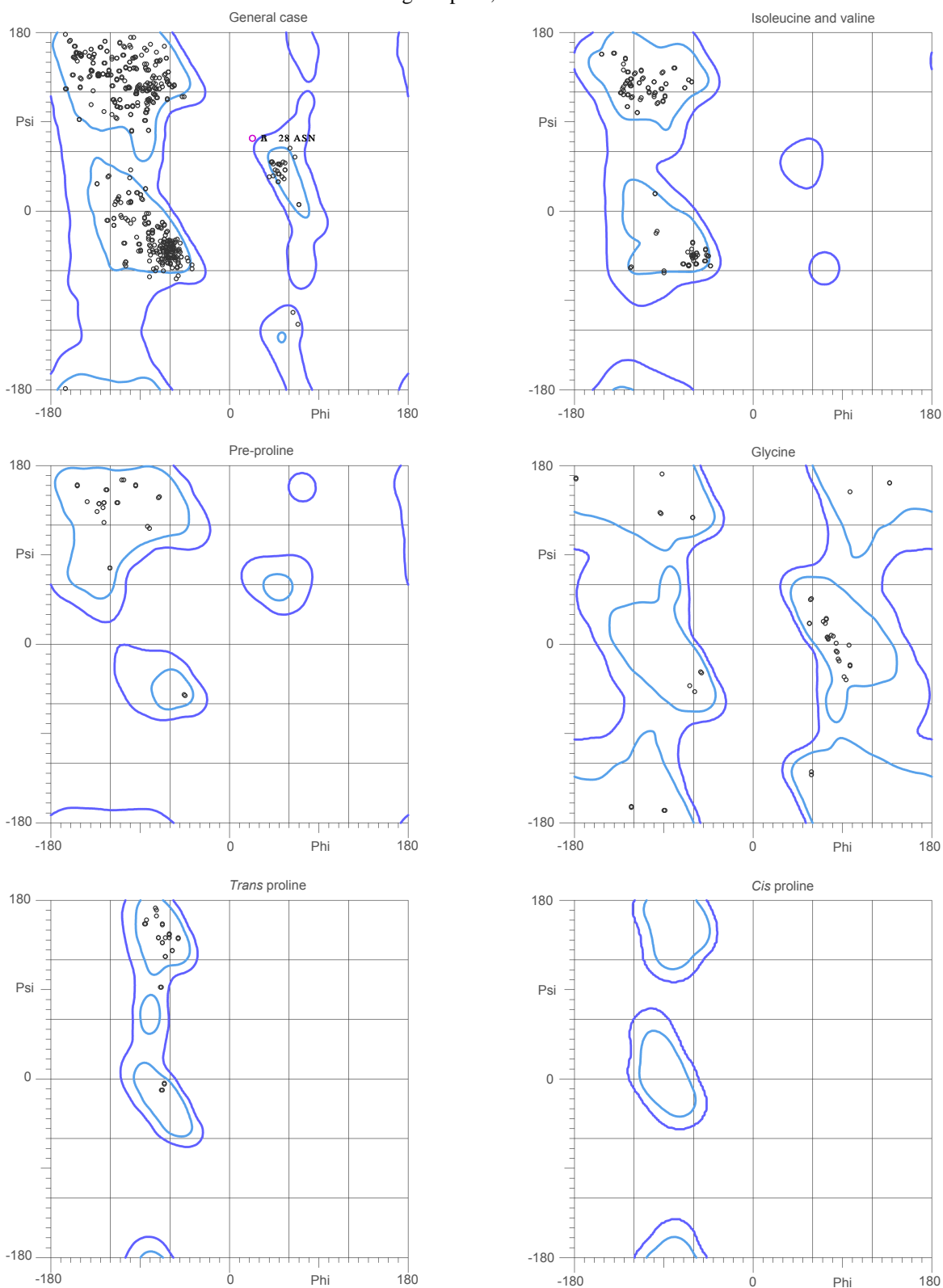


# MolProbity Ramachandran analysis

7keg.H.pdb, model 1



96.1% (671/698) of all residues were in favored (98%) regions.

99.7% (696/698) of all residues were in allowed (>99.8%) regions.

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

A 28 ASN (23.9, 74.7)

B 28 ASN (23.6, 74.2)