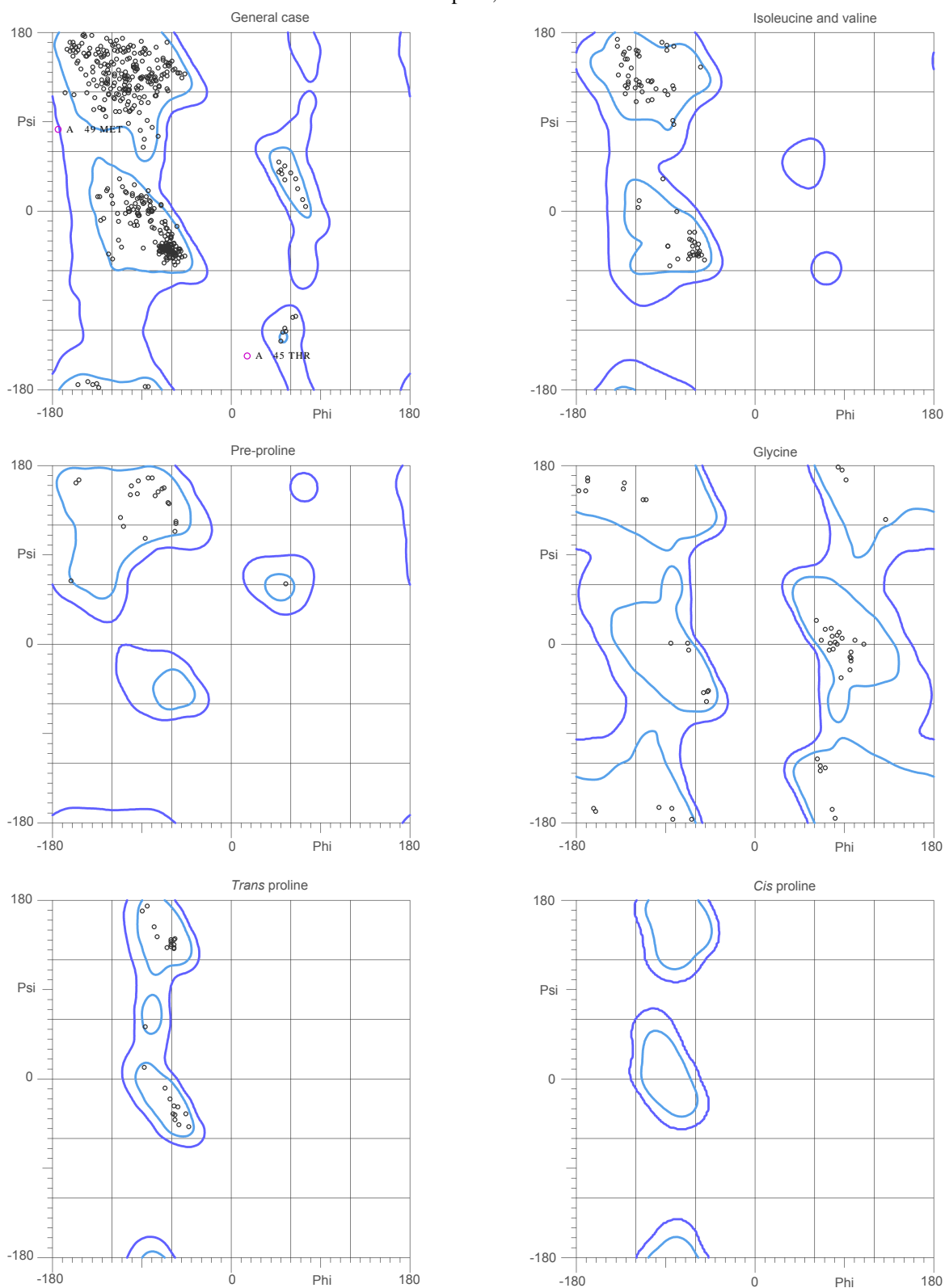


# MolProbity Ramachandran analysis

2duc.H.pdb, model 1



97.7% (594/608) of all residues were in favored (98%) regions.

99.7% (606/608) of all residues were in allowed (>99.8%) regions.

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

A 45 THR (16.3, -146.9)

A 49 MET (-175.9, 83.8)