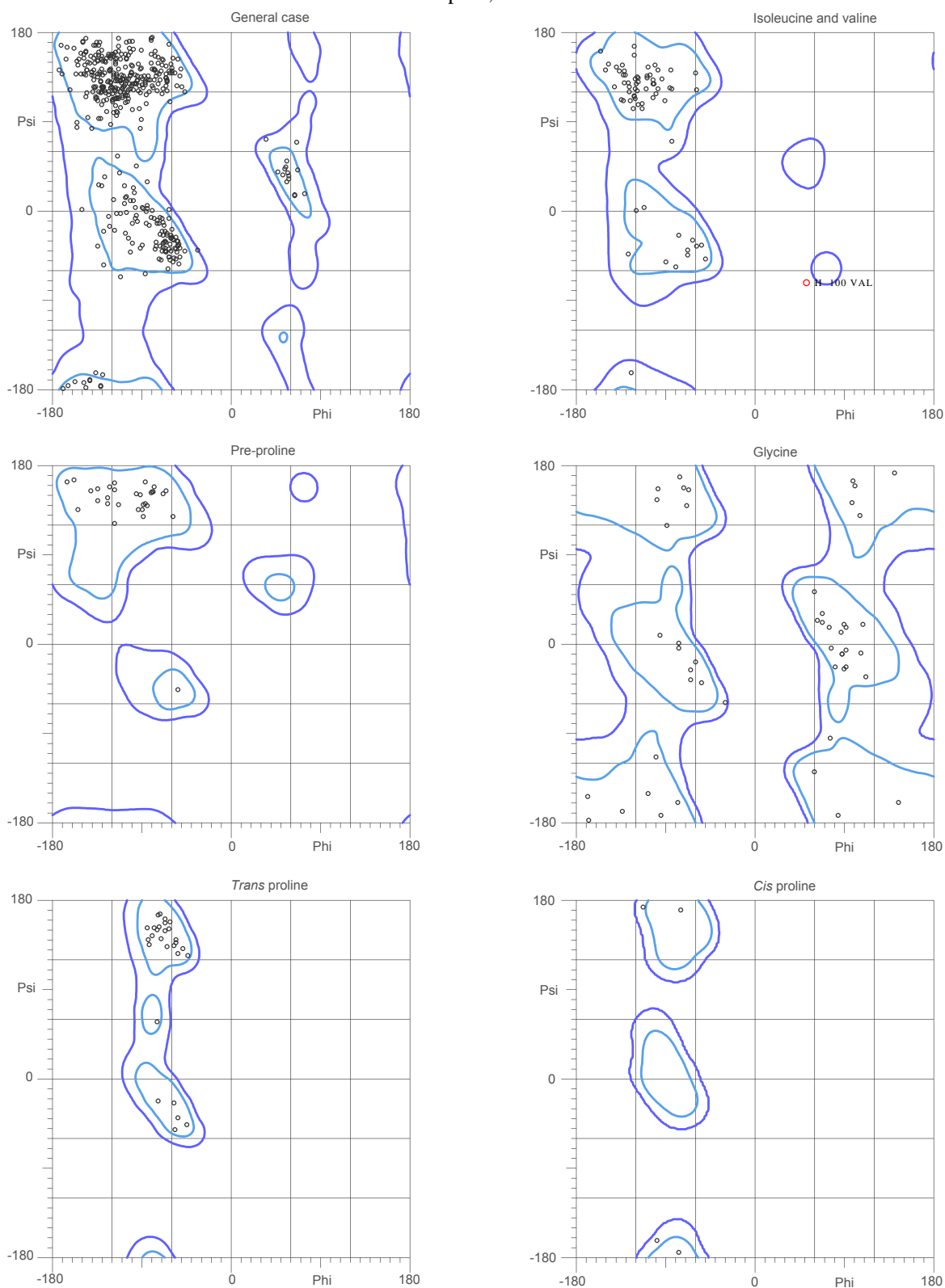


# MolProbity Ramachandran analysis

7cdi.H.pdb, model 1



95.0% (575/605) of all residues were in favored (98%) regions.

99.8% (604/605) of all residues were in allowed (>99.8%) regions.

There were 1 outliers ( $\phi$ ,  $\psi$ ):

H 100 VAL (52.7, -72.9)