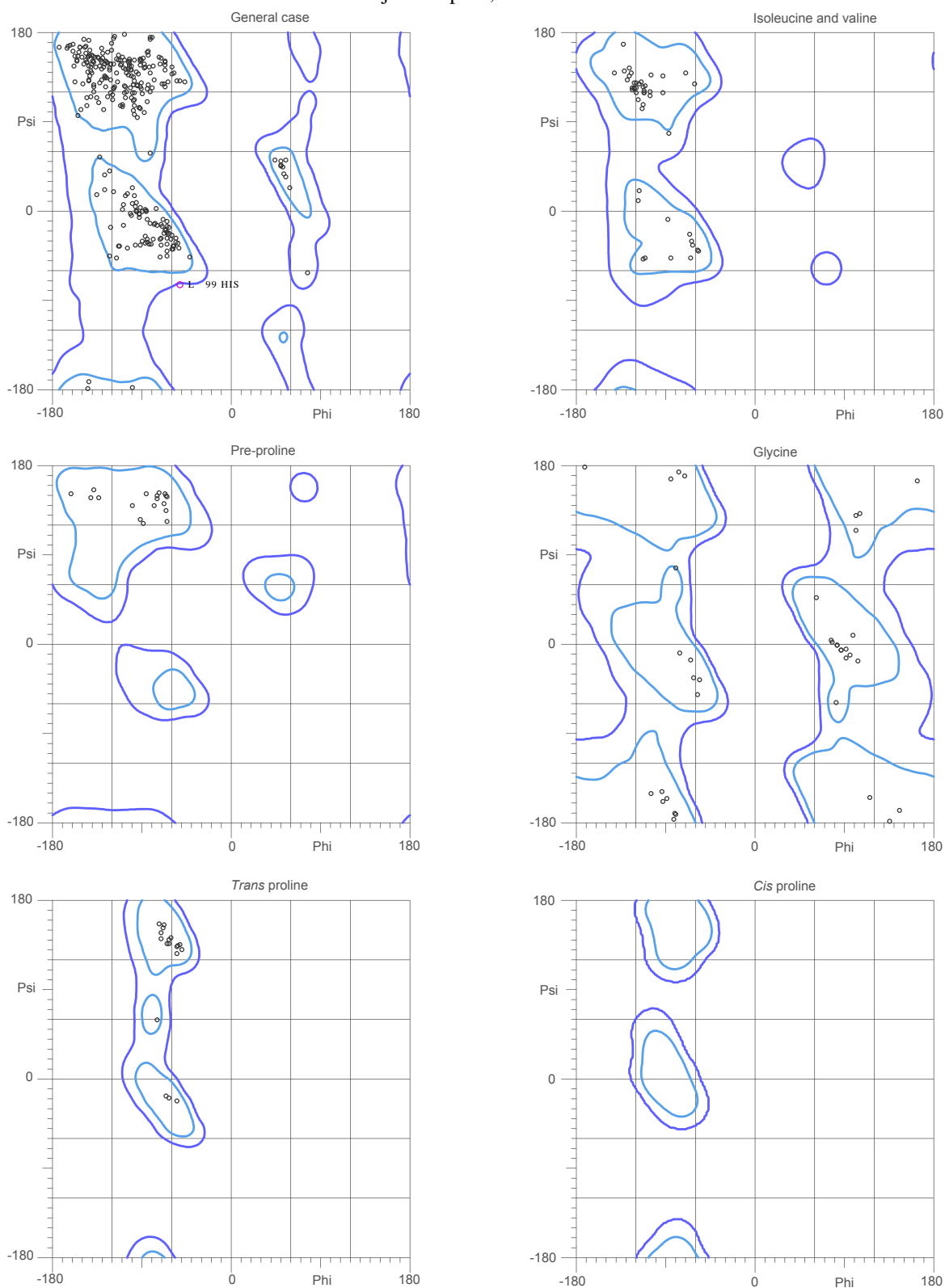


# MolProbity Ramachandran analysis

7jva.H.pdb, model 1



98.8% (407/412) of all residues were in favored (98%) regions.

99.8% (411/412) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

L 99 HIS (-52.2, -74.5)