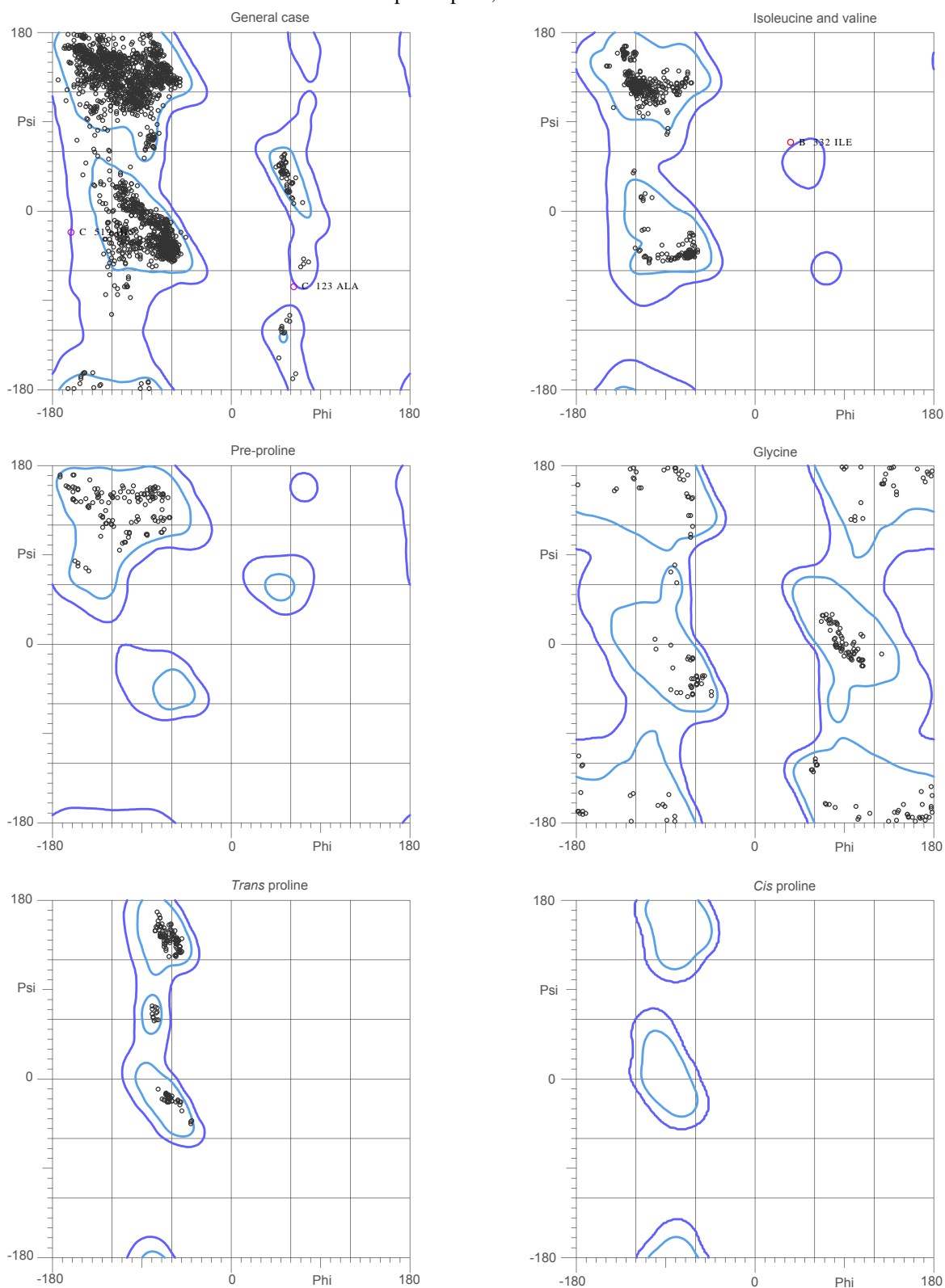# MolProbity Ramachandran analysis

6wpt.H.pdb, model 1

98.0% (3116/3181) of all residues were in favored (98%) regions.
99.9% (3178/3181) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi):
    B 332 ILE (36.9, 70.8)
    C 123 ALA (63.4, -76.2)
    C 519 HIS (-162.1, -21.2)

http://kinemage.biochem.duke.edu