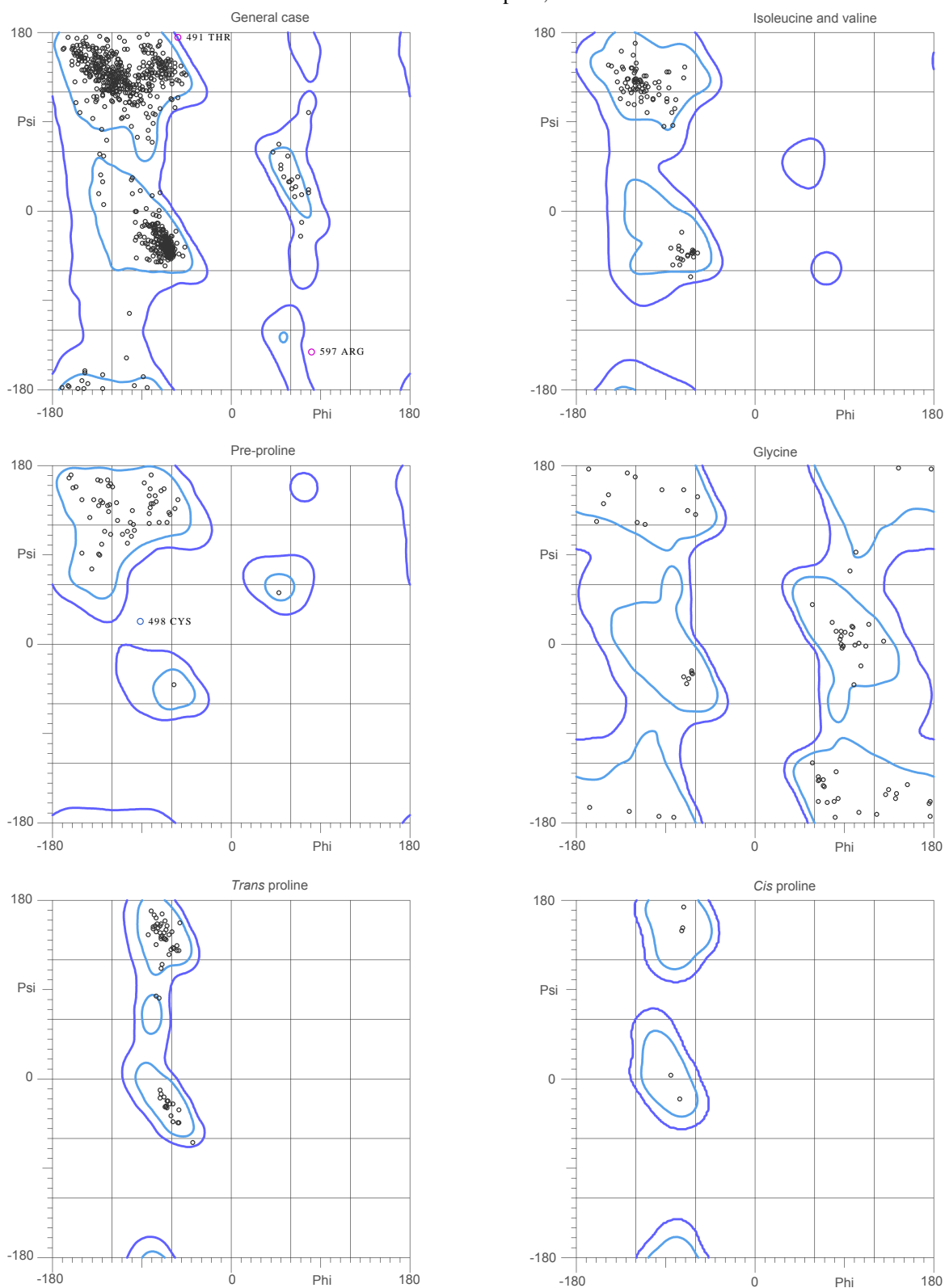# MolProbity Ramachandran analysis

1f4h.B99990018.pdb, model 1

97,4% (992/1019) of all residues were in favored (98%) regions.
99,7% (1016/1019) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi):
    491 THR (-54,3, 176,7)
    498 CYS (-92,4, 23,7)
    597 ARG (81,1, -142,4)

http://kinemage.biochem.duke.edu   Lovell, Davis, et al. Proteins 50:437 (2003)