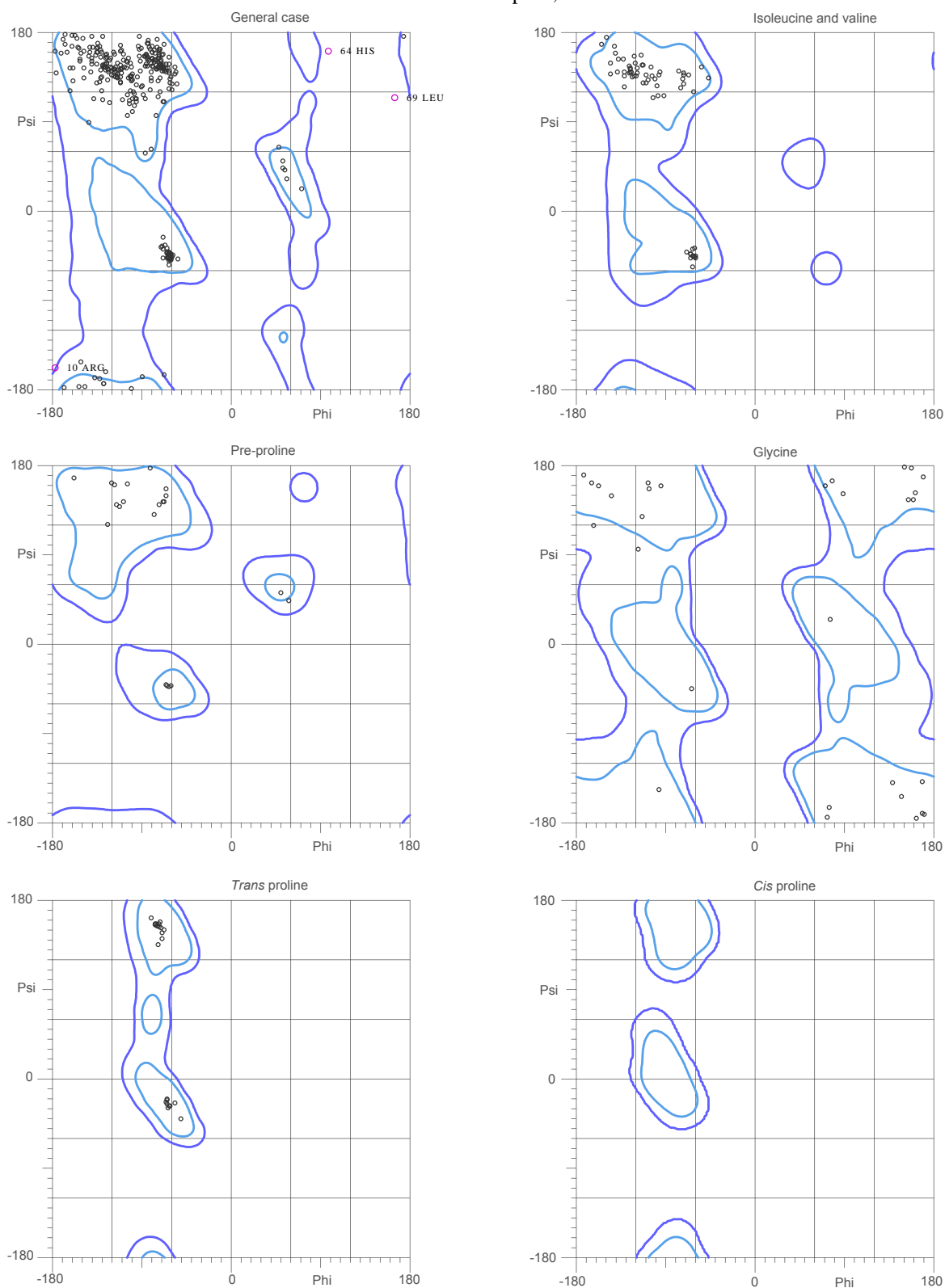# MolProbity Ramachandran analysis

1ad3A.B99990012.pdb, model 1

96,4% (428/444) of all residues were in favored (98%) regions.

99,3% (441/444) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi):

    10 ARG (-178,1, -159,0)

    64 HIS (98,8, 163,0)

    69 LEU (165,9, 115,2)

http://kinemage.biochem.duke.edu

Lovell, Davis, et al. Proteins 50:437 (2003)