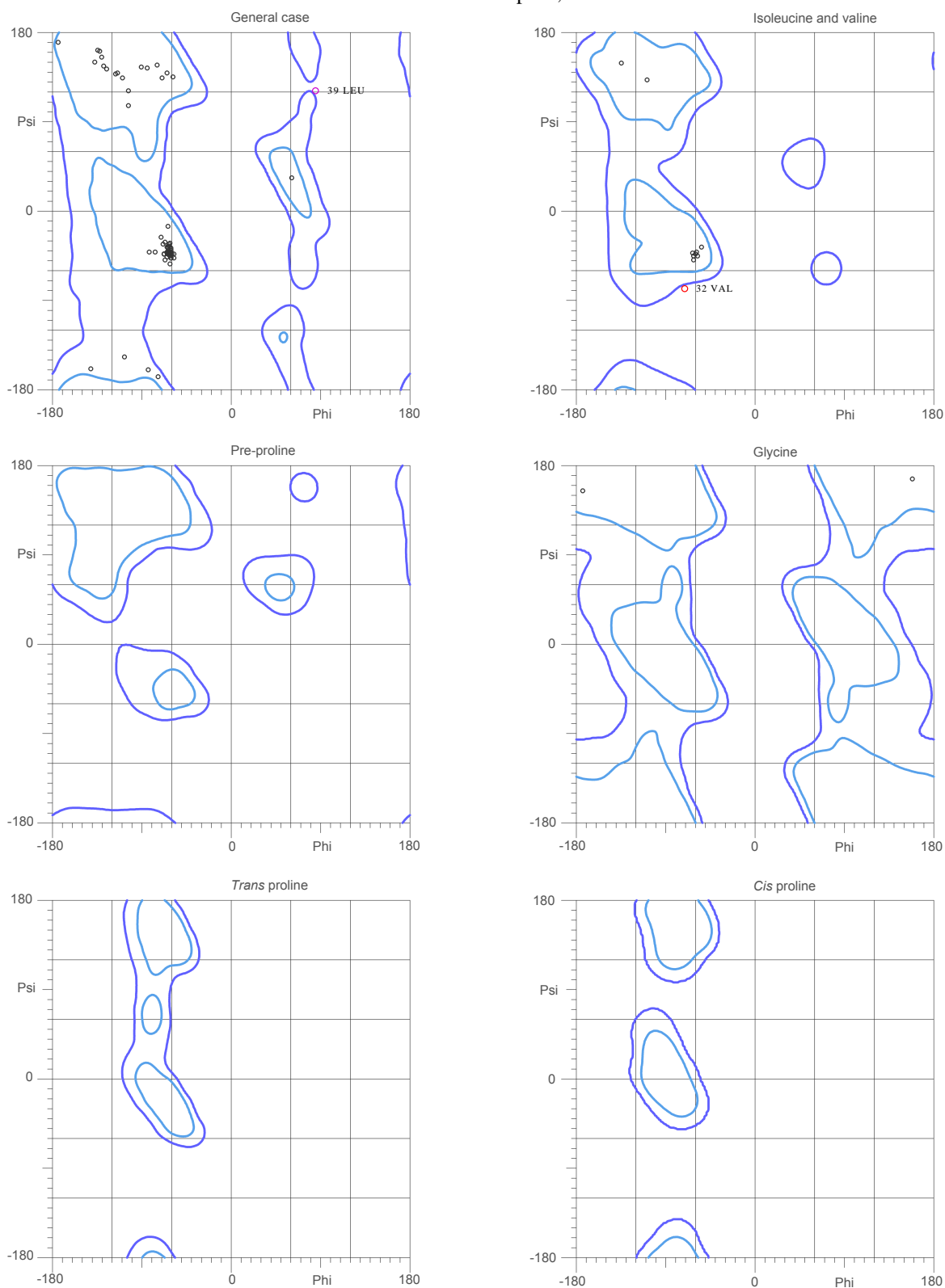# MolProbity Ramachandran analysis

1am9A.B99990019.pdb, model 1

92,3% (72/78) of all residues were in favored (98%) regions.

97,4% (76/78) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

    32 VAL (-71,2, -78,3)

    39 LEU (85,3, 122,8)

http://kinemage.biochem.duke.edu

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