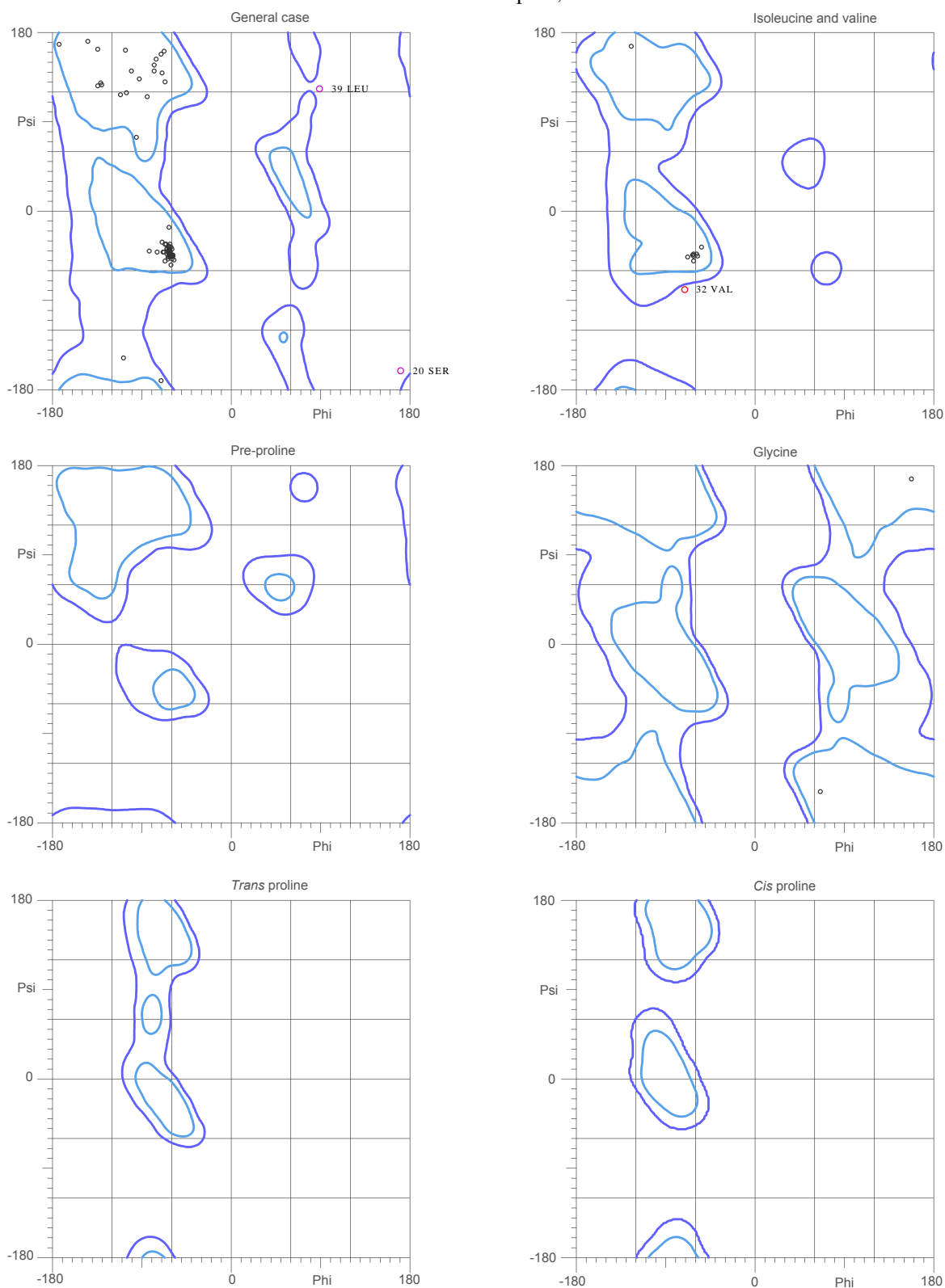# MolProbity Ramachandran analysis

1am9A.B99990034.pdf, model 1



93,6% (73/78) of all residues were in favored (98%) regions.
96,2% (75/78) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi):
    20 SER (171,4, -161,9)
    32 VAL (-71,2, -79,7)
    39 LEU (89,4, 124,3)

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