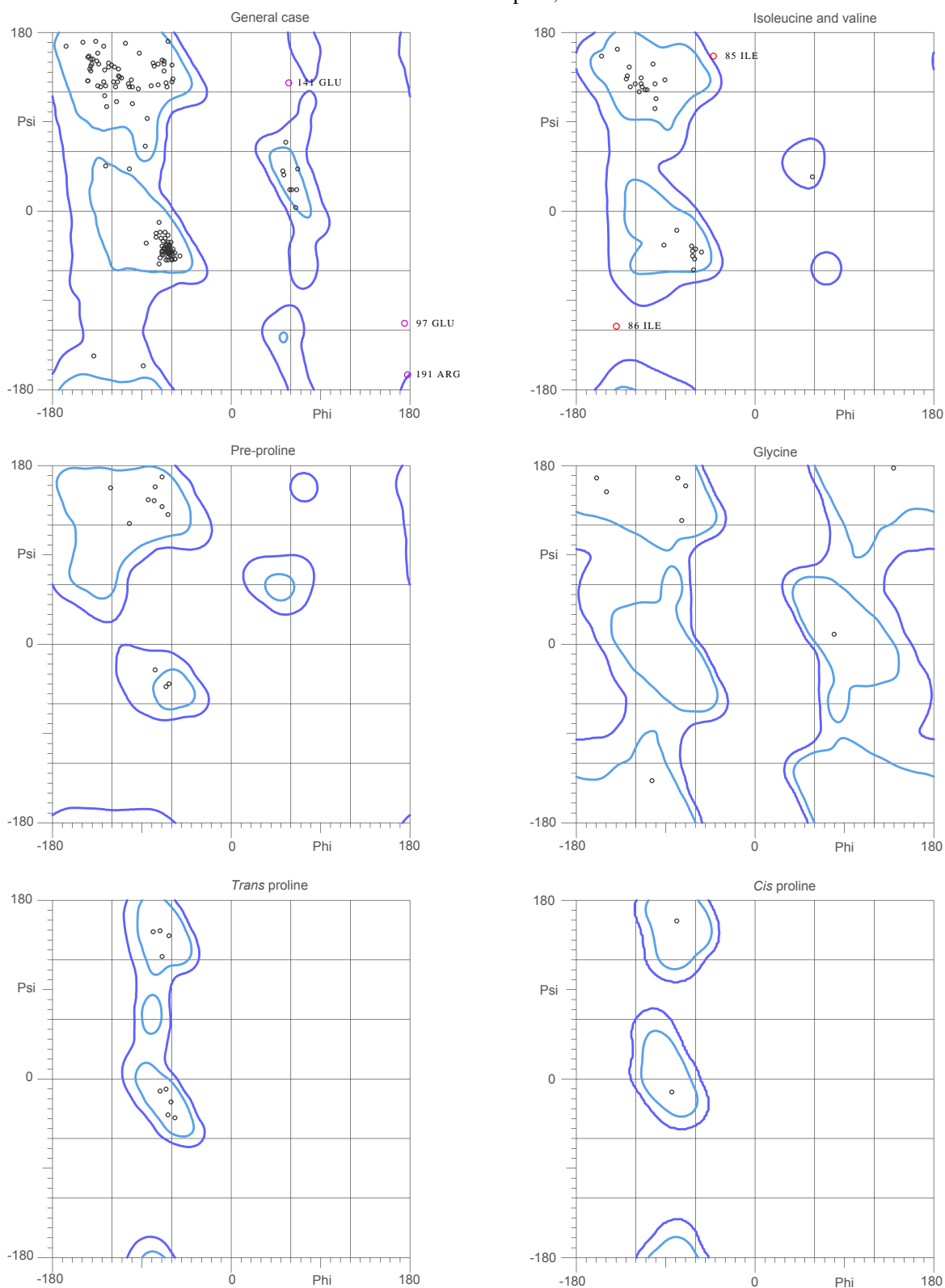# MolProbity Ramachandran analysis

1aoeA.B99990020.pdb, model 1

93,7% (178/190) of all residues were in favored (98%) regions.
97,4% (185/190) of all residues were in allowed (>99.8%) regions.

There were 5 outliers (phi, psi):
    85 ILE (-43,0, 157,9)
    86 ILE (-140,9, -116,6)
    97 GLU (175,8, -113,2)
    141 GLU (58,2, 130,8)
    191 ARG (178,2, -165,1)

http://kinemage.biochem.duke.edu

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