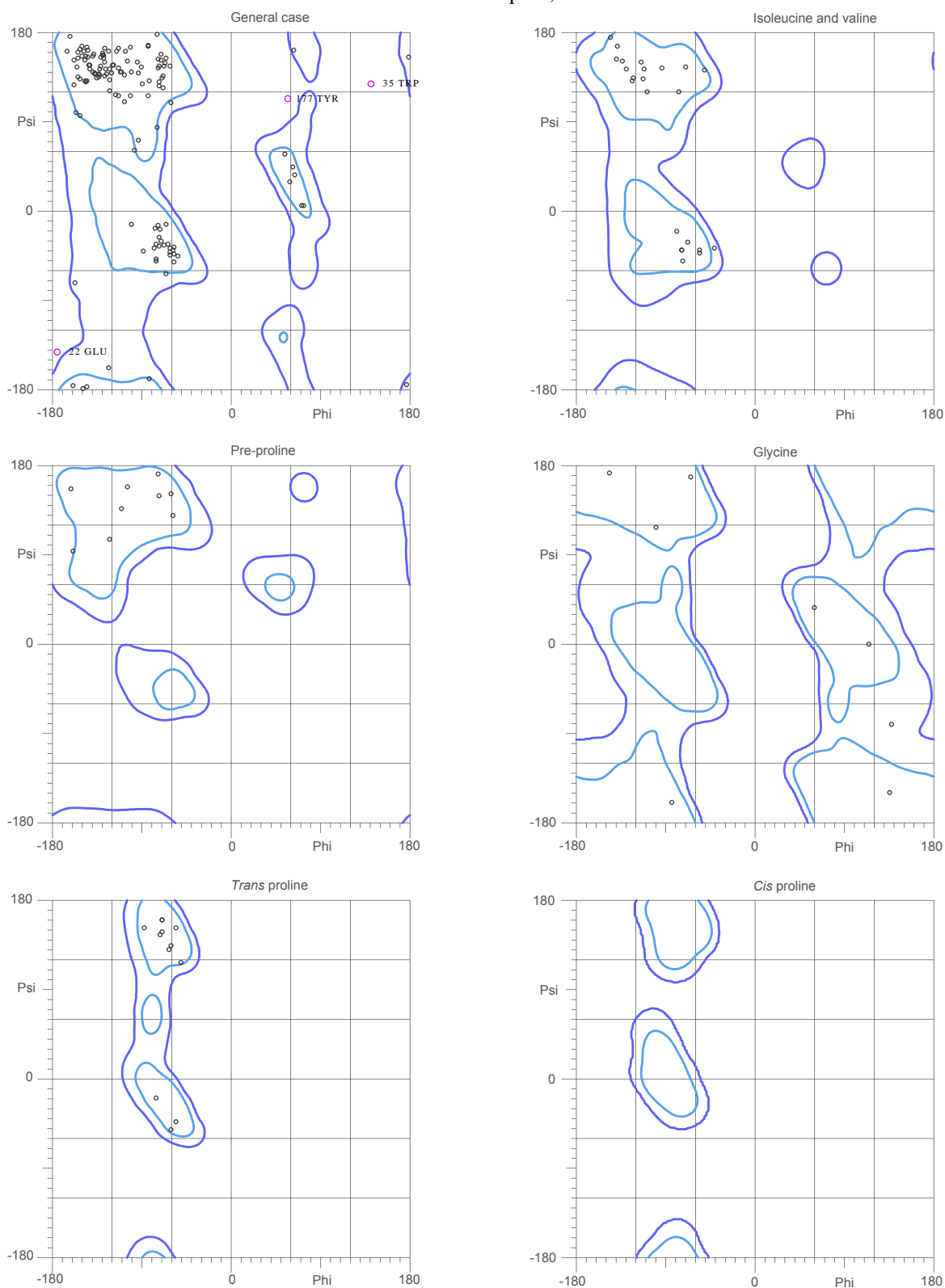# MolProbity Ramachandran analysis

1axiB.B99990028.pdf, model 1

91,5% (173/189) of all residues were in favored (98%) regions.
98,4% (186/189) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi):
    22 GLU (-176,9, -142,2)
    35 TRP (141,2, 129,6)
    177 TYR (57,1, 114,4)

http://kinemage.biochem.duke.edu

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