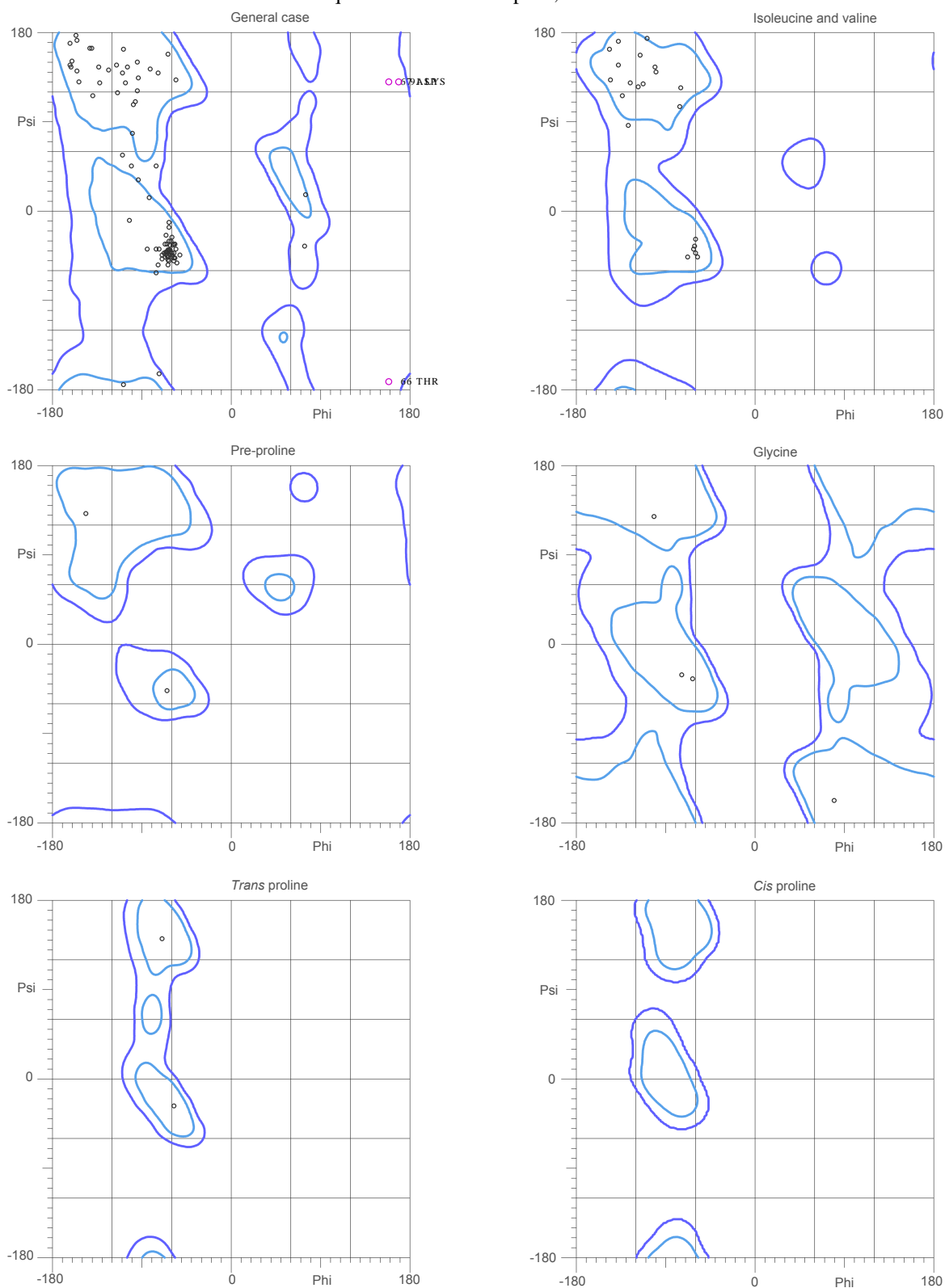# MolProbity Ramachandran analysis

1aqcA.B99990035.pdb, model 1

89,9% (107/119) of all residues were in favored (98%) regions.

97,5% (116/119) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi):

    66 THR (159,6, -173,0)

    67 ASP (159,9, 131,6)

    91 LYS (169,6, 132,0)

http://kinemage.biochem.duke.edu

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