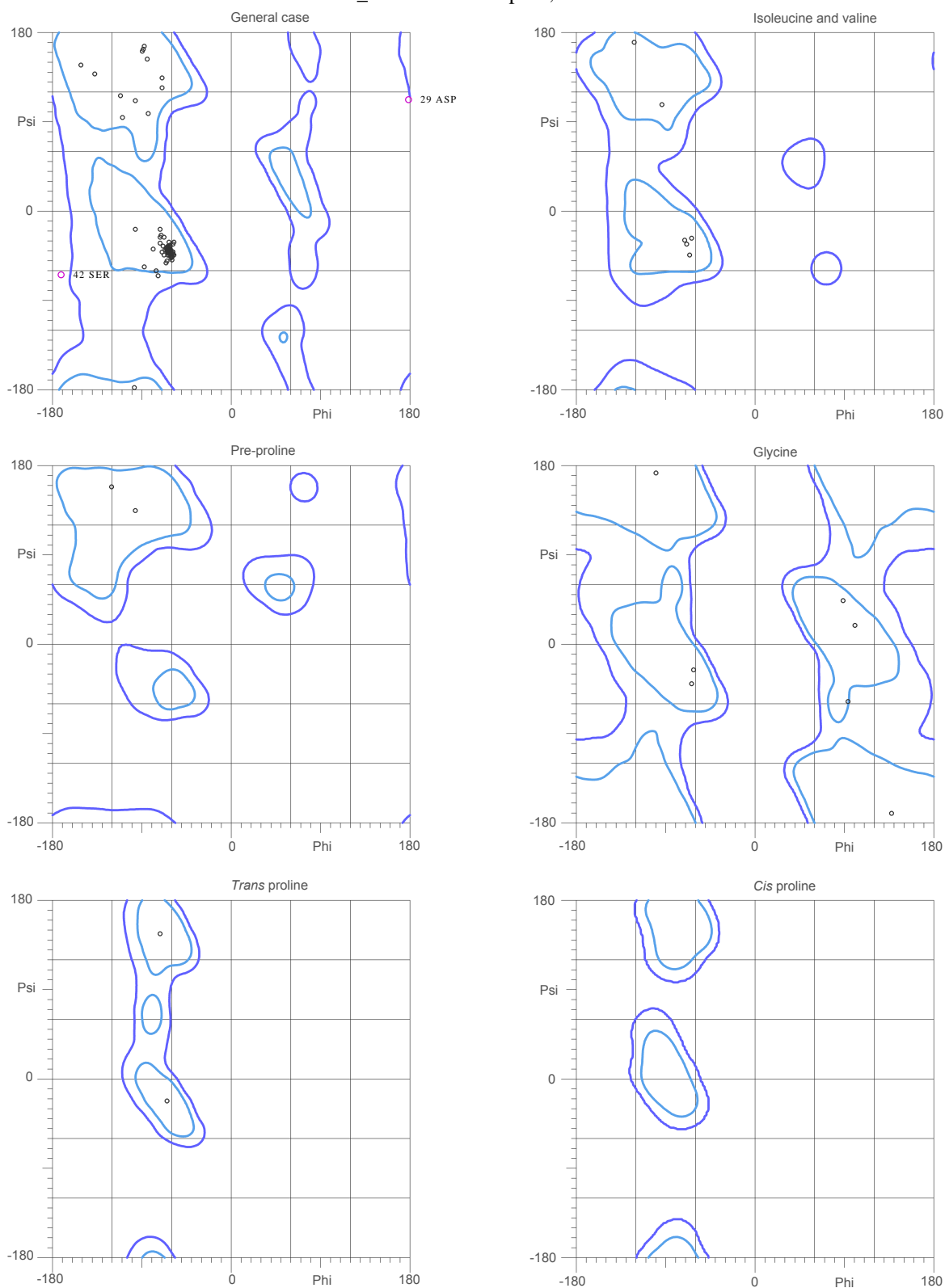# MolProbity Ramachandran analysis

1af8_.B99990040.pdb, model 1

96,4% (81/84) of all residues were in favored (98%) regions.

97,6% (82/84) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

29 ASP (179,1, 113,7)

42 SER (-172,3, -64,3)

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