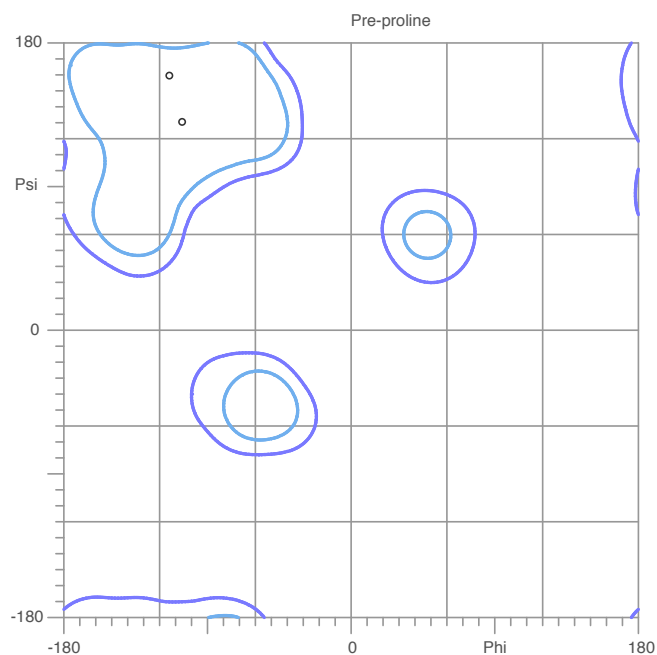
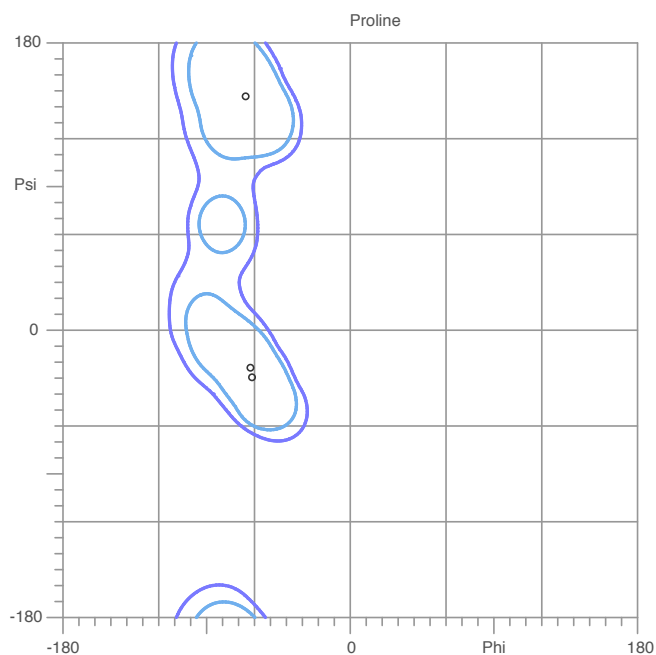
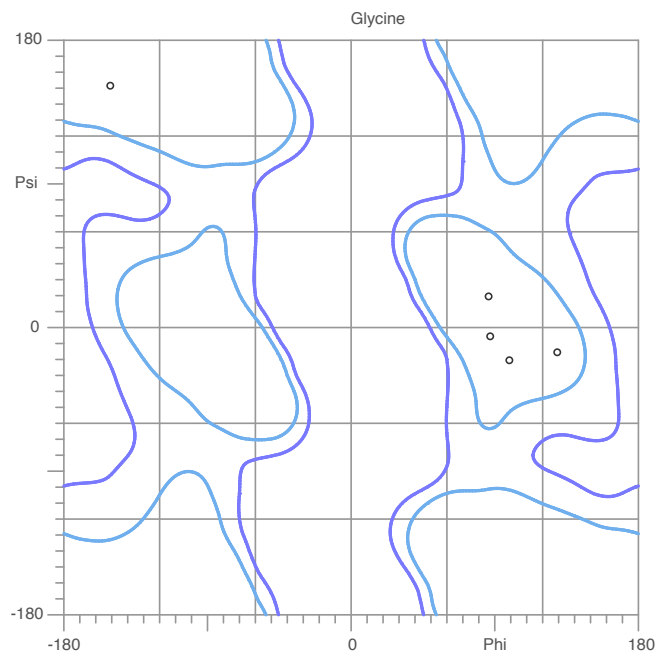
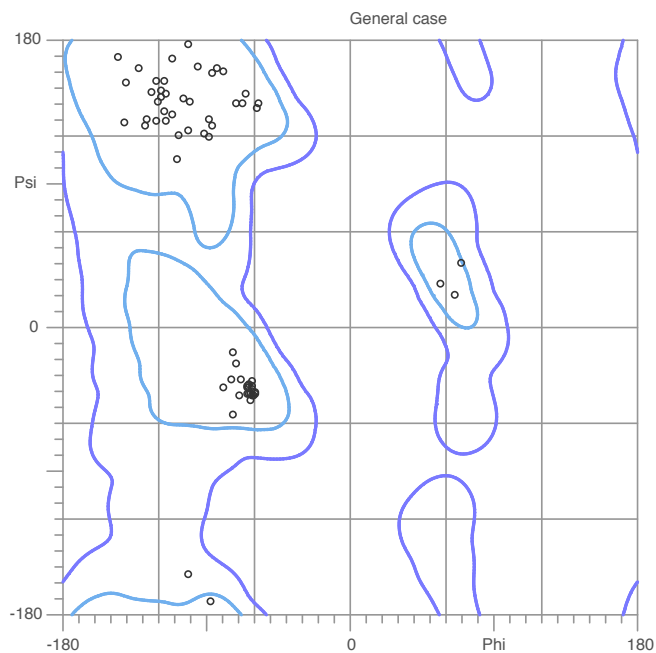


# MolProbity Ramachandran analysis

alvo.B99990003.pdb~, model 1



97,3% (72/74) of all residues were in favored (98%) regions.  
100,0% (74/74) of all residues were in allowed (>99.8%) regions.

There were no outliers.

<http://kinemage.biochem.duke.edu>

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