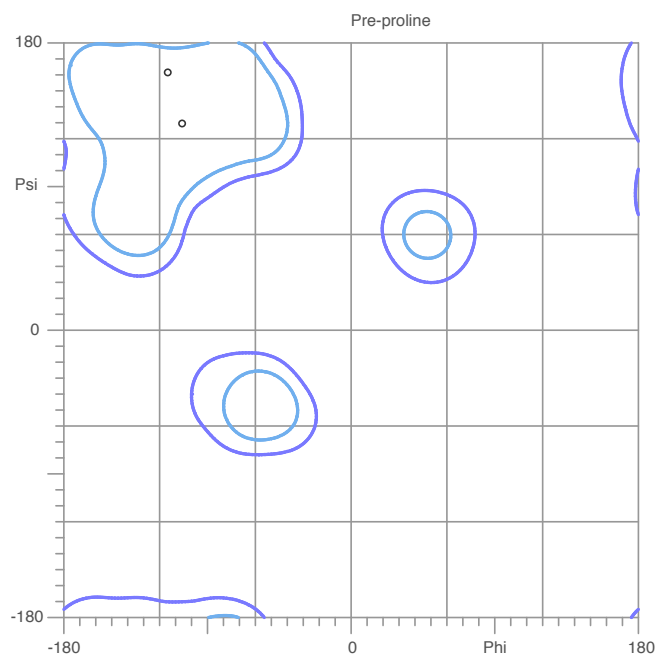
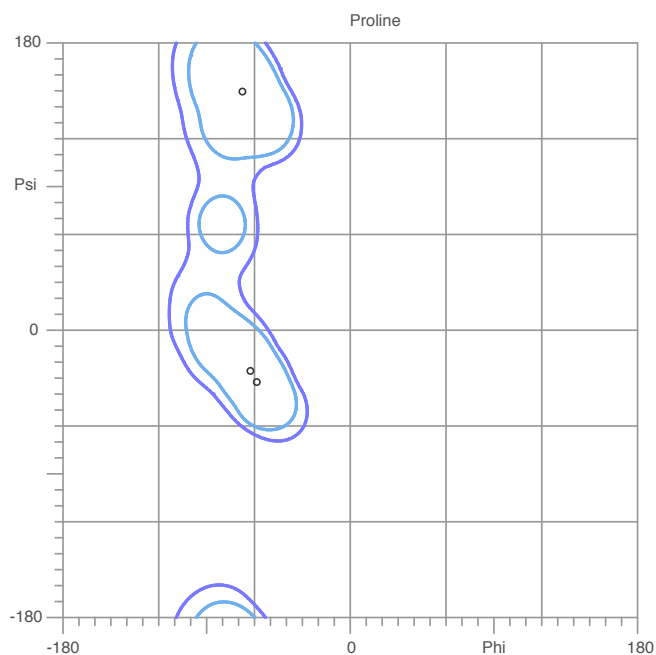
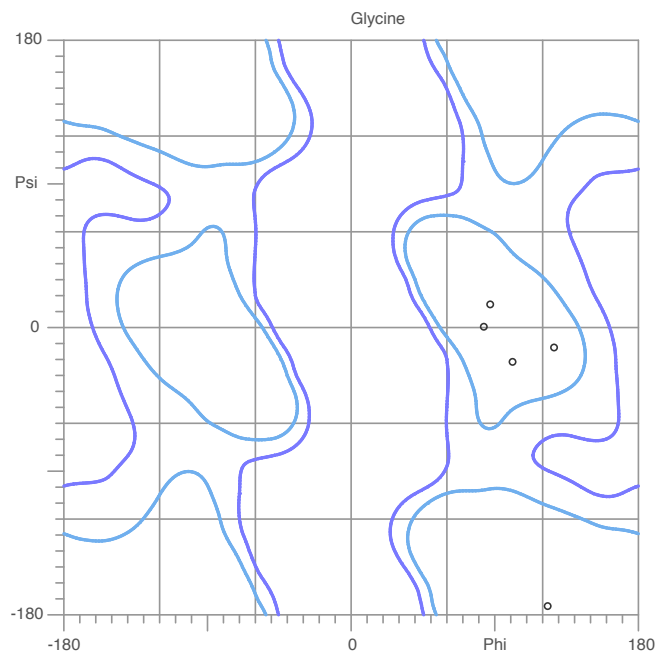
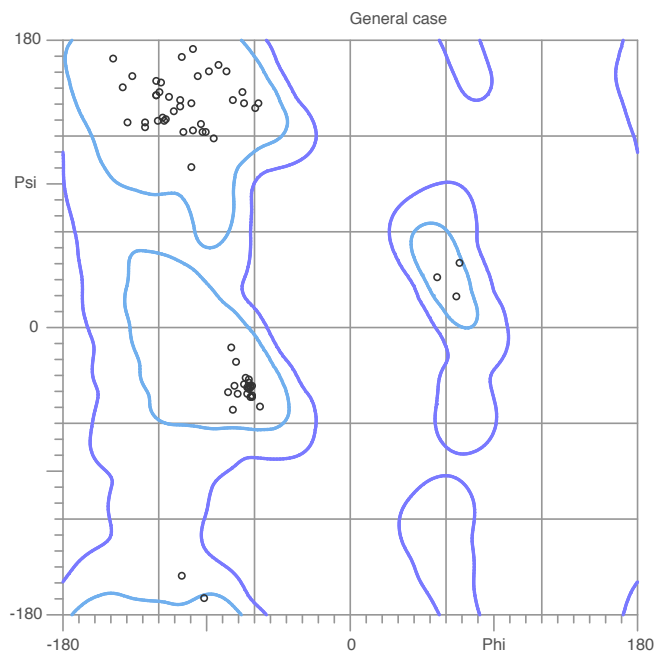


# MolProbity Ramachandran analysis

alvo.B99990001.pdb, model 1



98,6% (73/74) 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)