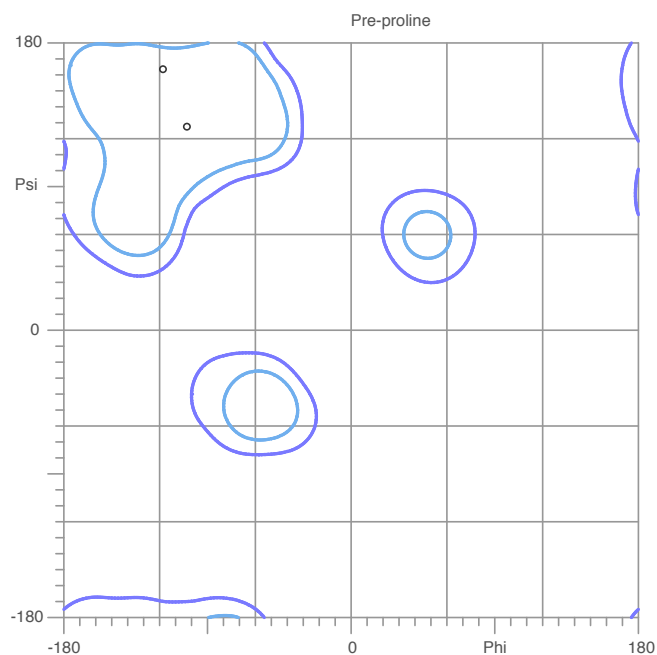
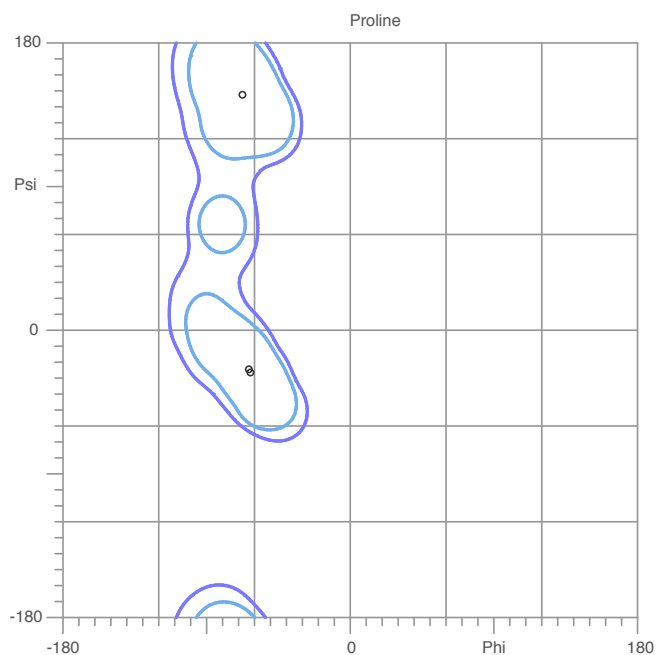
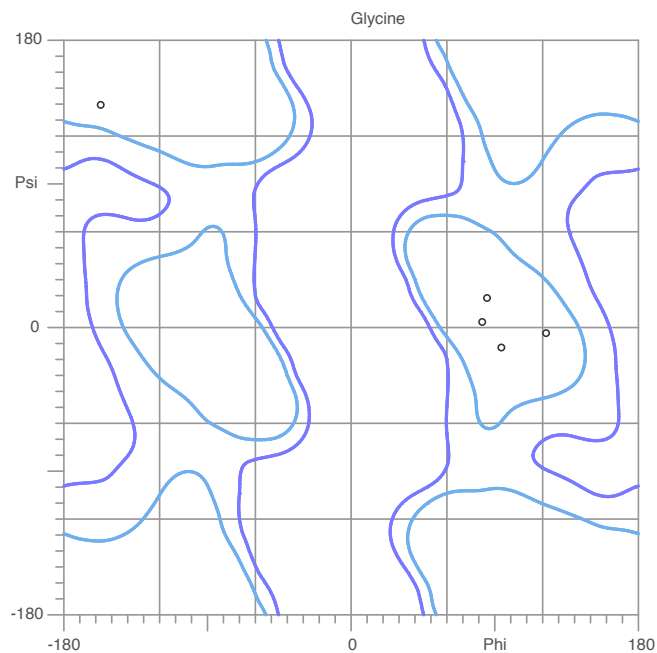
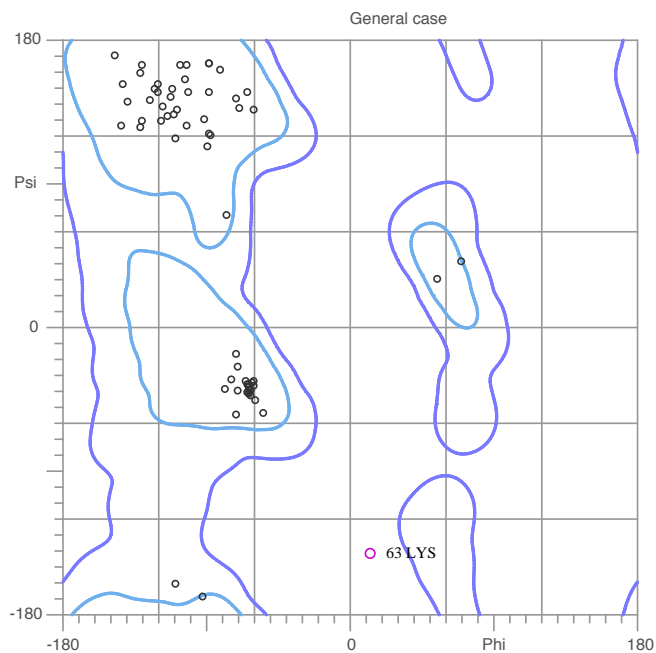


MolProbity Ramachandran analysis

alvo.B99990001.pdb~, model 1



95,9% (71/74) of all residues were in favored (98%) regions.
98,6% (73/74) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):
63 LYS (12,0, -141,5)

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

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