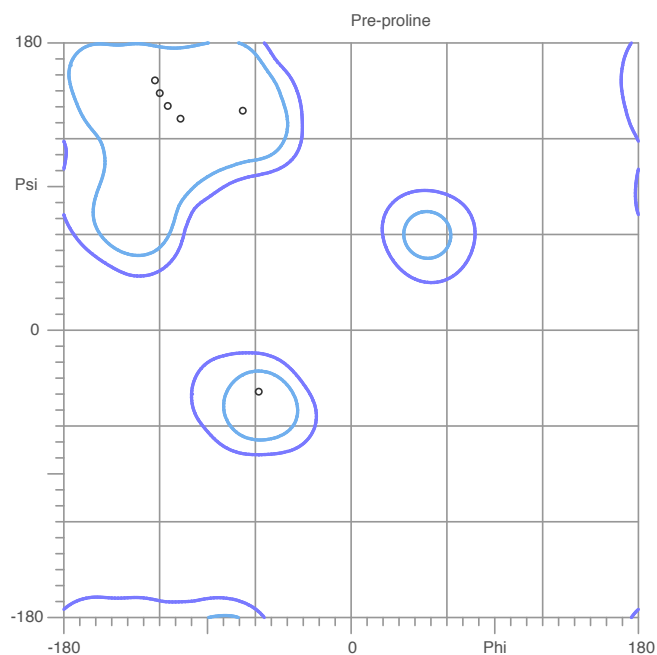
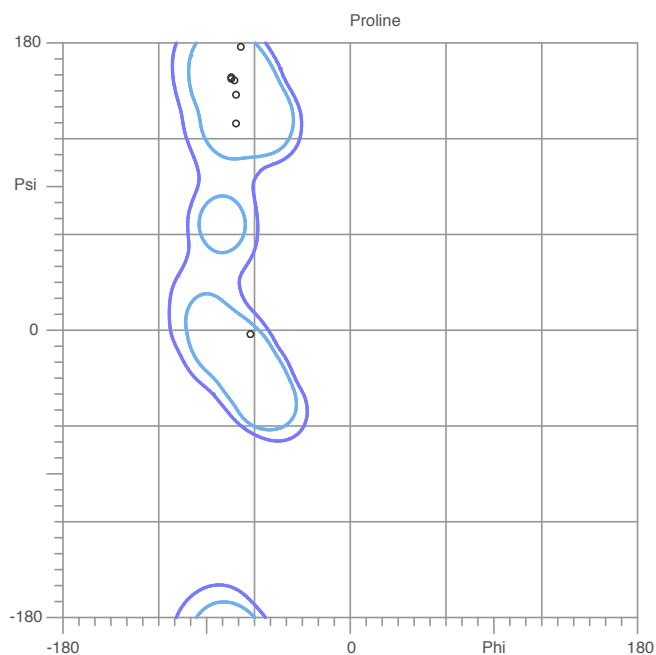
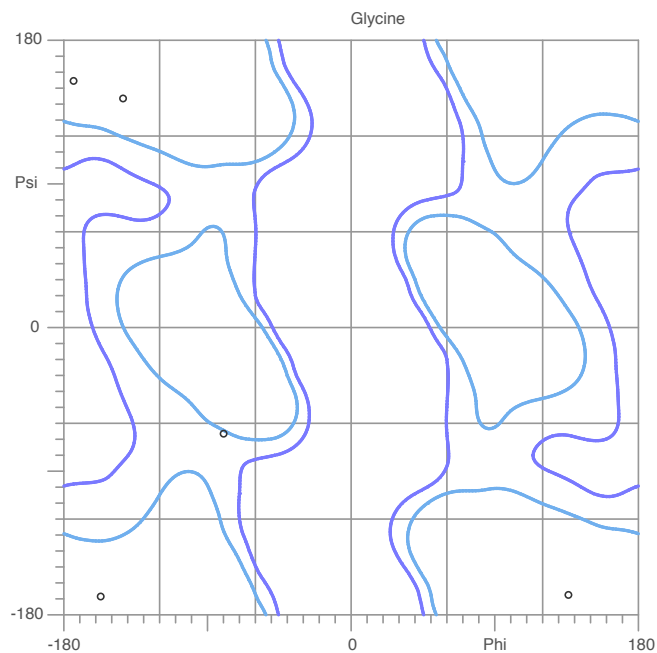
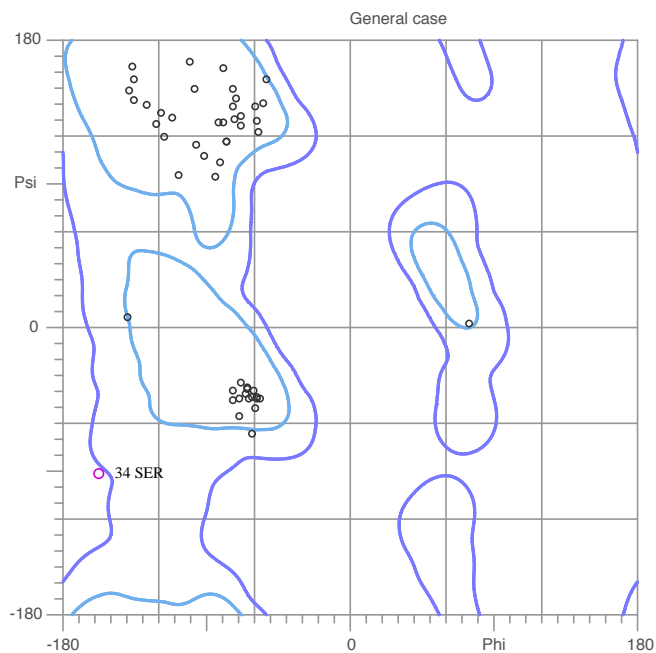


MolProbity Ramachandran analysis

SODEF17.B99990002.pdb, model 1



94.2% (65/69) of all residues were in favored (98%) regions.
98.6% (68/69) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):
34 SER (-158.4, -91.5)

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

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