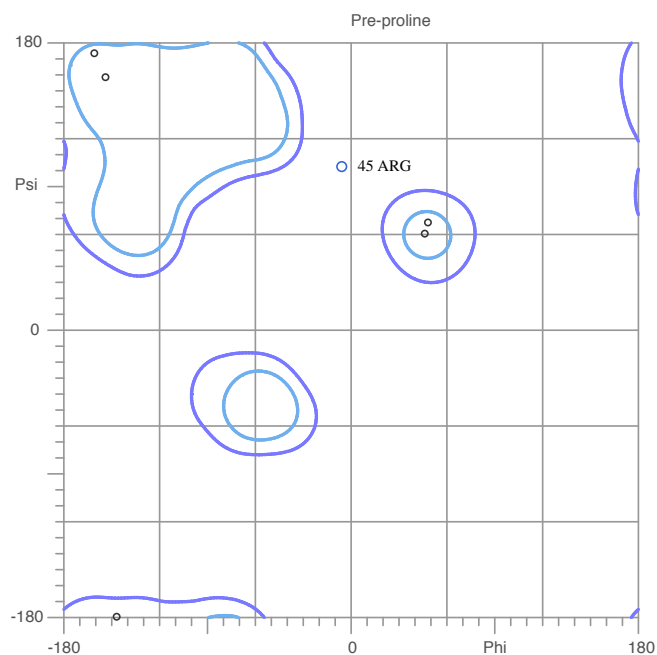
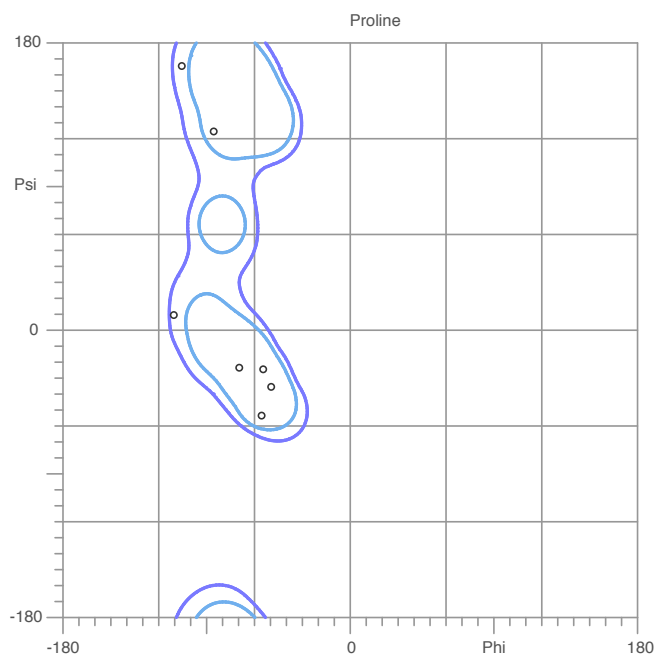
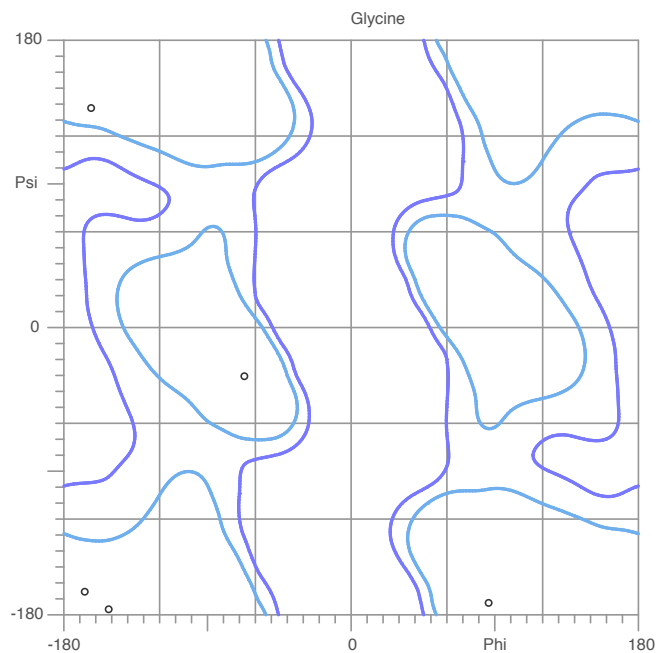
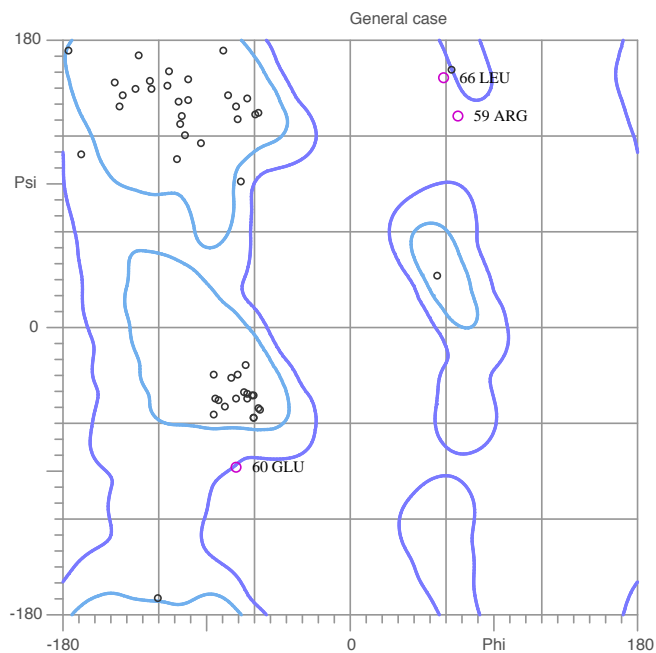


MolProbity Ramachandran analysis

SODEF17.B99990002.pdb, model 1



84,1% (58/69) of all residues were in favored (98%) regions.
94,2% (65/69) of all residues were in allowed (>99.8%) regions.

There were 4 outliers (phi, psi):

45 ARG (-6,3, 103,3)
59 ARG (67,1, 134,0)
60 GLU (-72,3, -87,1)
66 LEU (58,8, 157,9)

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

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