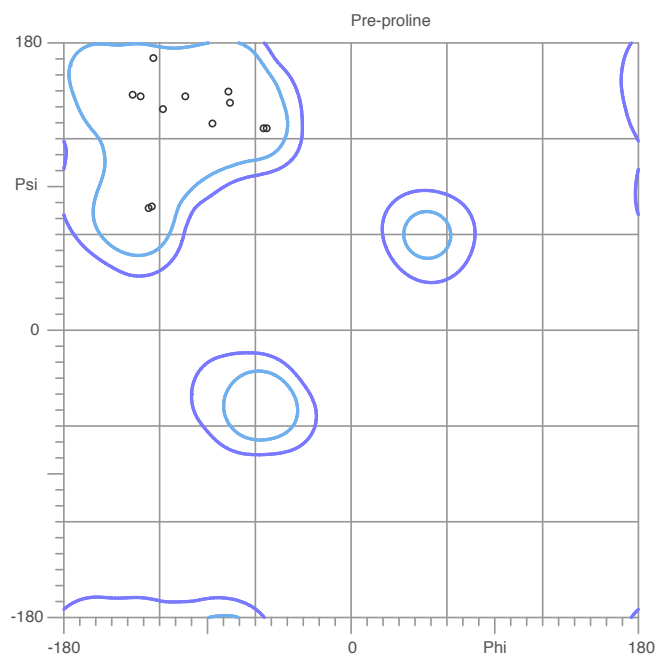
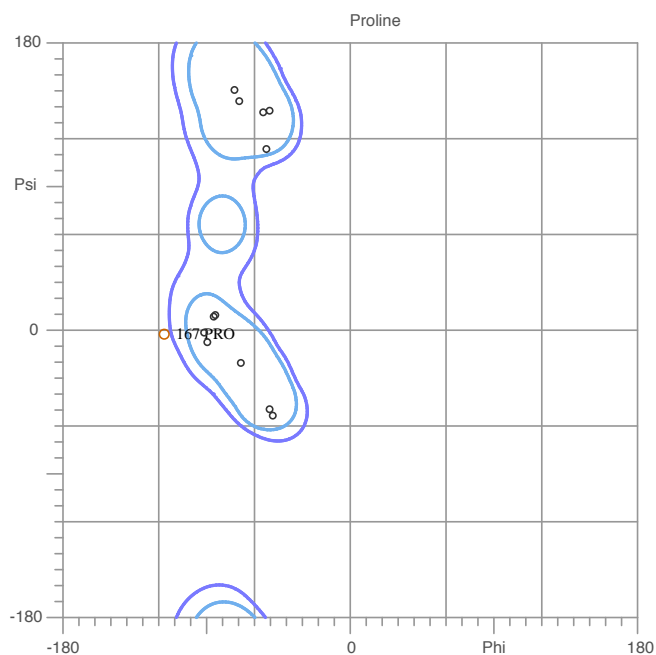
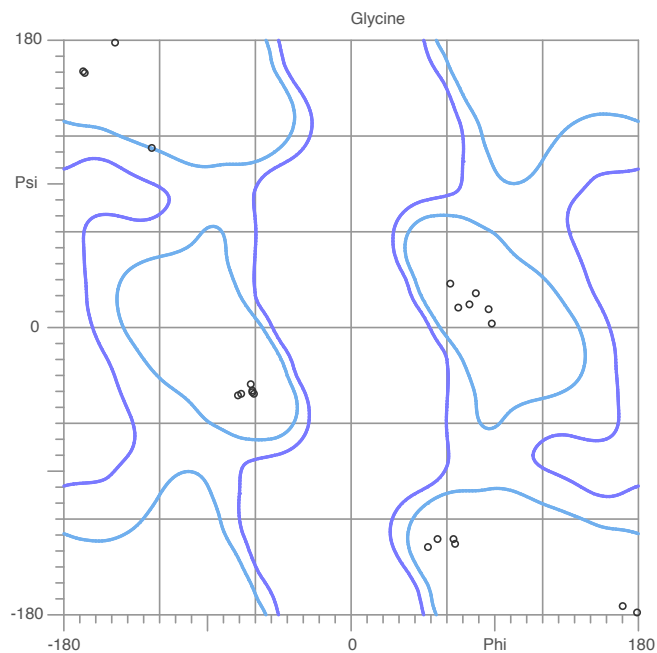
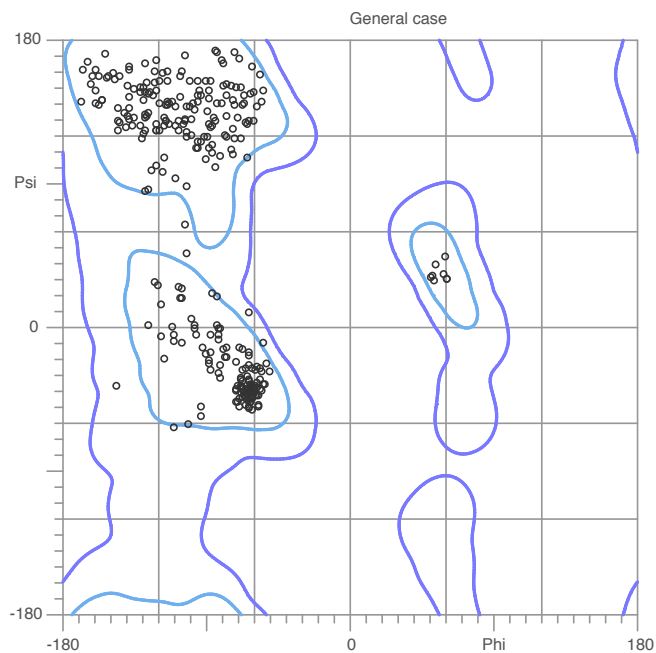


MolProbity Ramachandran analysis

OR8C_XRay_em_bcr3.pdb, model 1



98.0% (346/353) of all residues were in favored (98%) regions.
99.7% (352/353) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):
167 PRO (-117.7, -2.1)

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

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