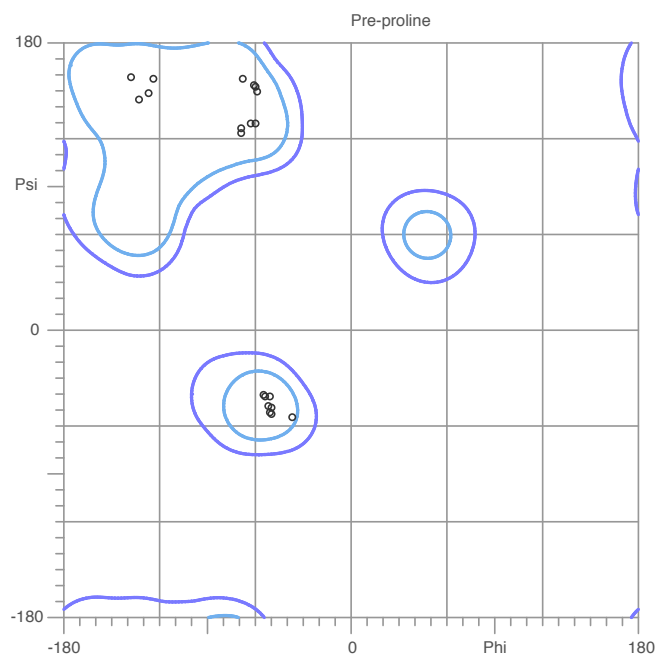
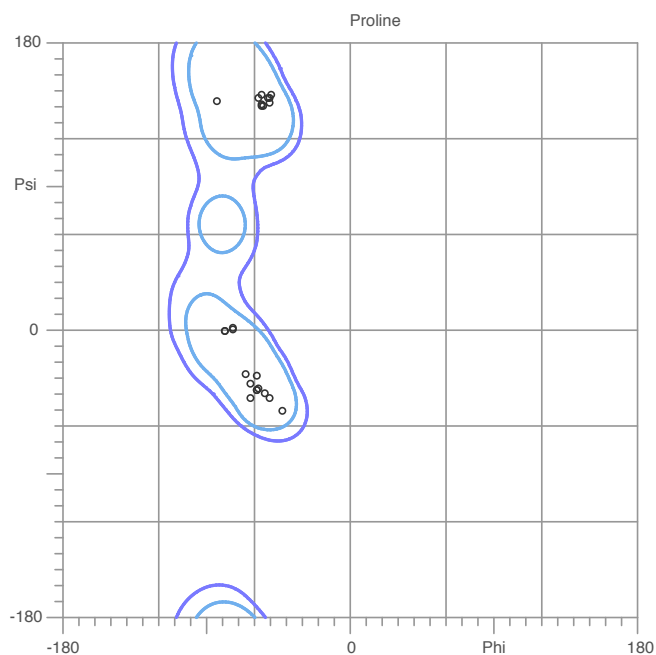
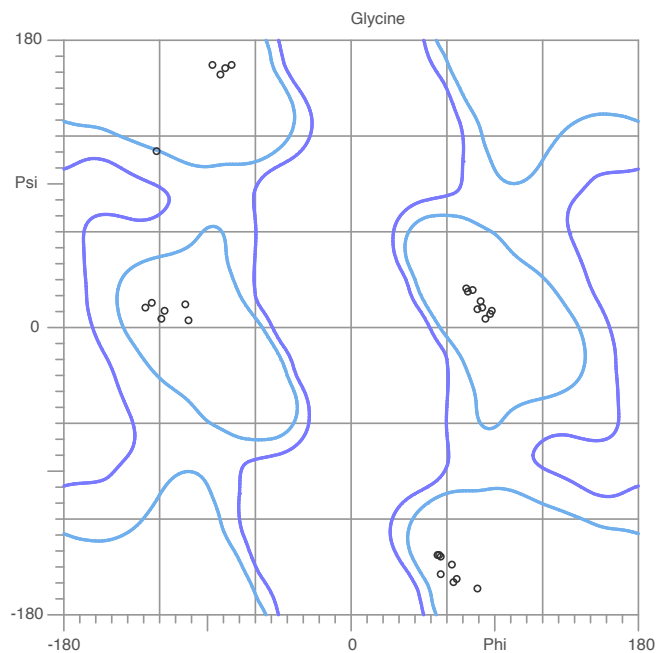
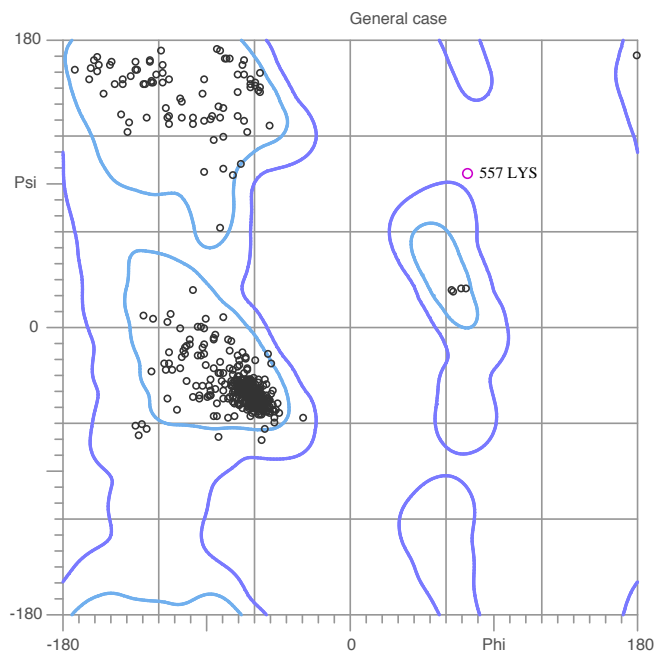


MolProbity Ramachandran analysis

SGR209C_XRay_em_bcr3.pdb, model 1



98.3% (519/528) of all residues were in favored (98%) regions.
99.8% (527/528) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):
557 LYS (73.3, 97.5)

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

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