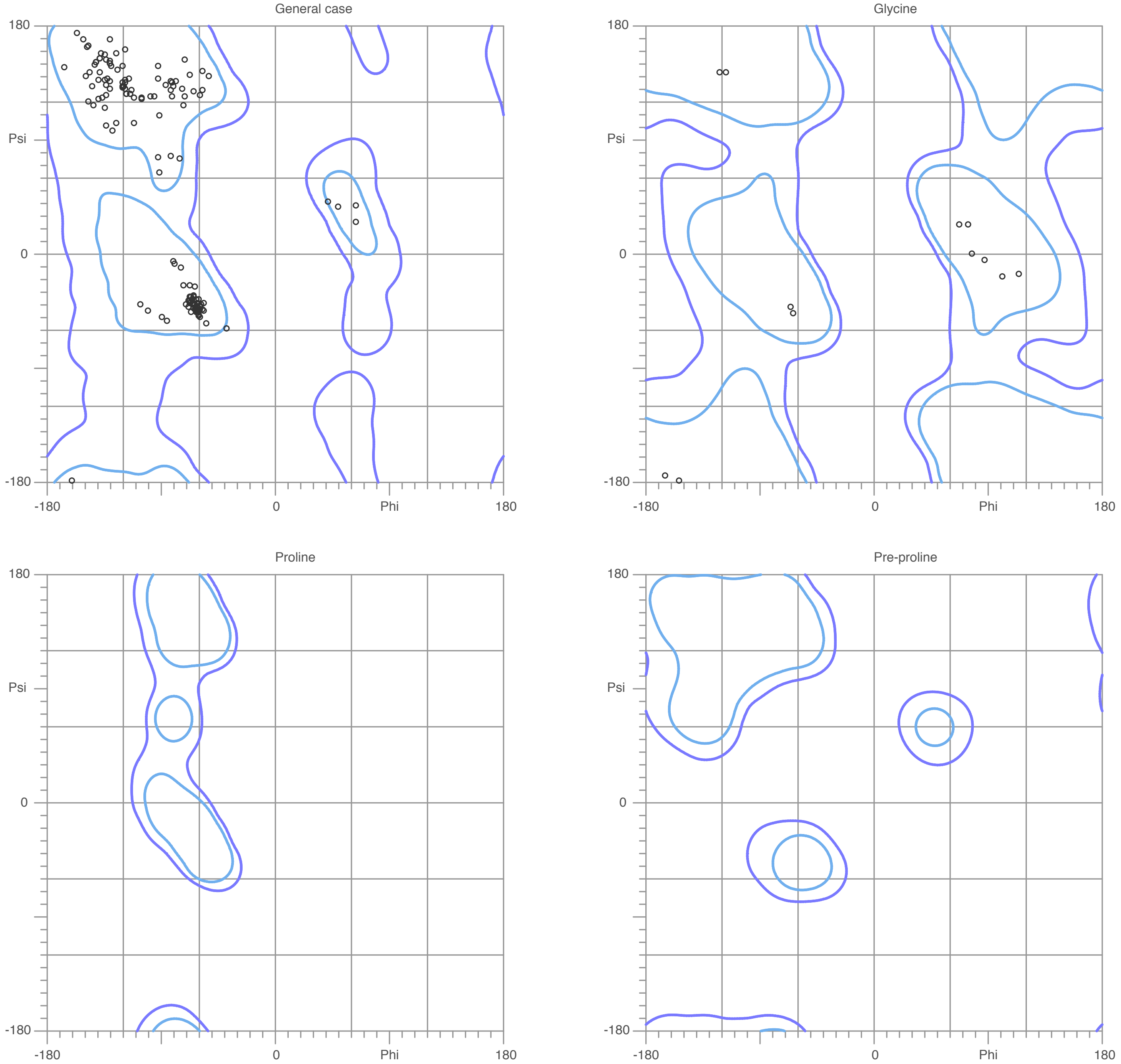
MolProbity Ramachandran analysis

CTR148A_XRay_em_bcr3.pdb, model 1



99.3% (151/152) of all residues were in favored (98%) regions.
100.0% (152/152) of all residues were in allowed (>99.8%) regions.

There were no outliers.

http://kinemage.biochem.duke.edu

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