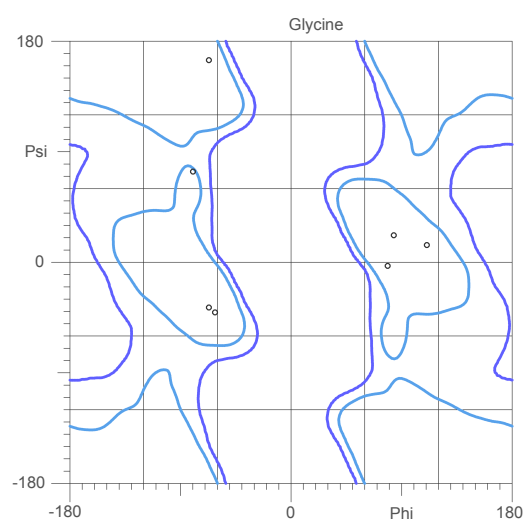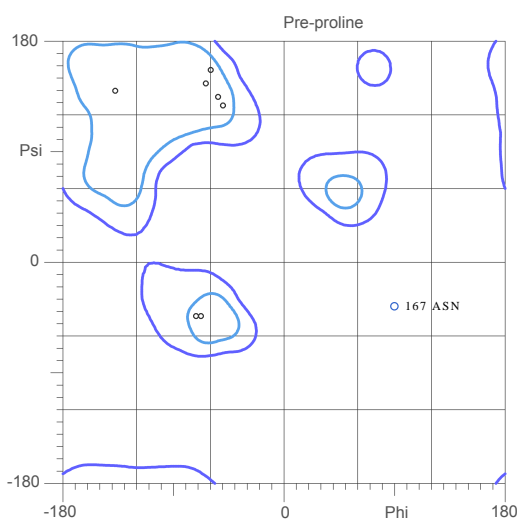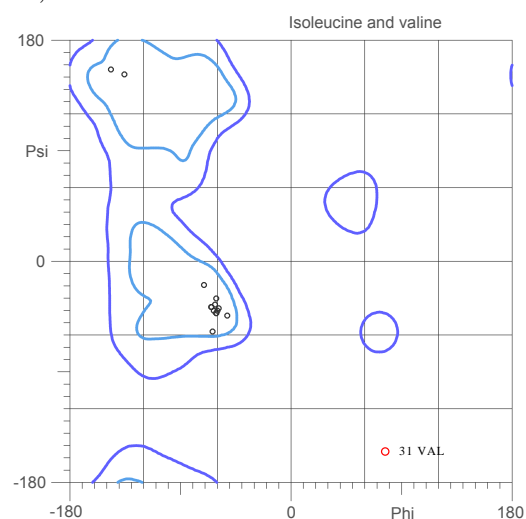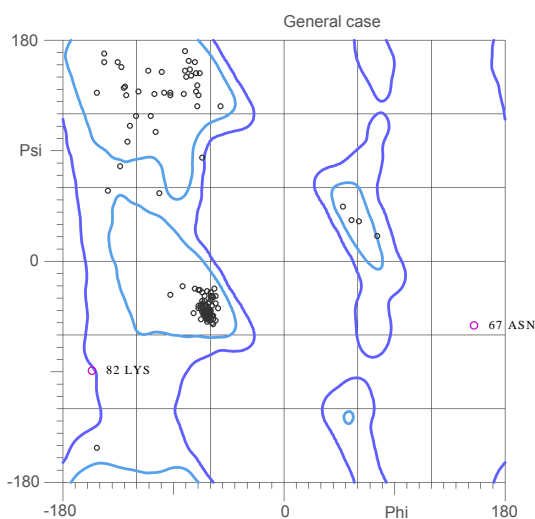# MolProbity Ramachandran analysis

1aihA.B99990027.pdb, model 1
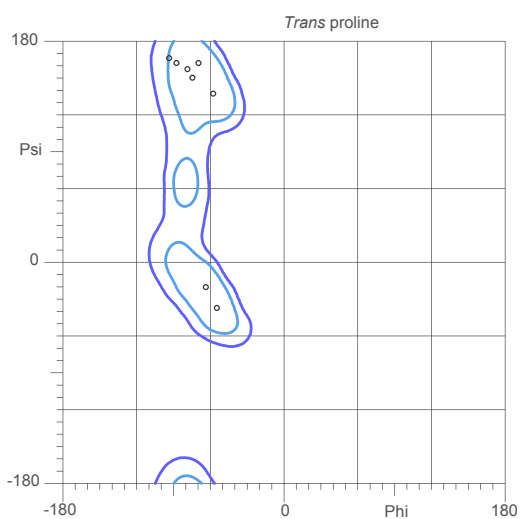
93,5% (157/168) of all residues were in favored (98%) regions.
97,6% (164/168) of all residues were in allowed (>99.8%) regions.

There were 4 outliers (phi, psi):
  31 VAL (78,0, -155,8)
  67 ASN (155,8, -52,5)
  82 LYS (-157,1, -89,8)
  167 ASN (90,1, -36,4)

http://kinemage.biochem.duke.edu   Lovell, Davis, et al. Proteins 50:437 (2003)