MolProbity Ramachandran analysis

1agjA.B99990014.pdb, model 1
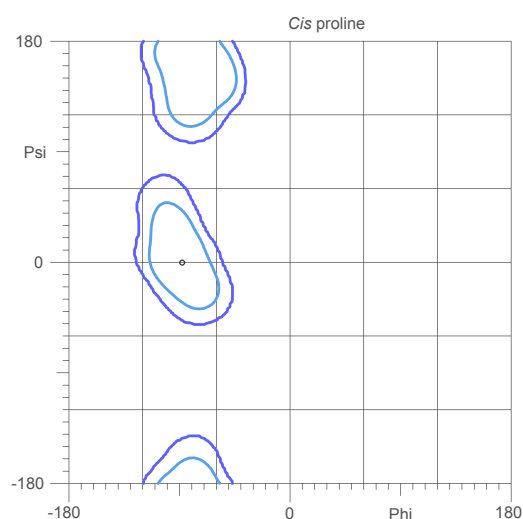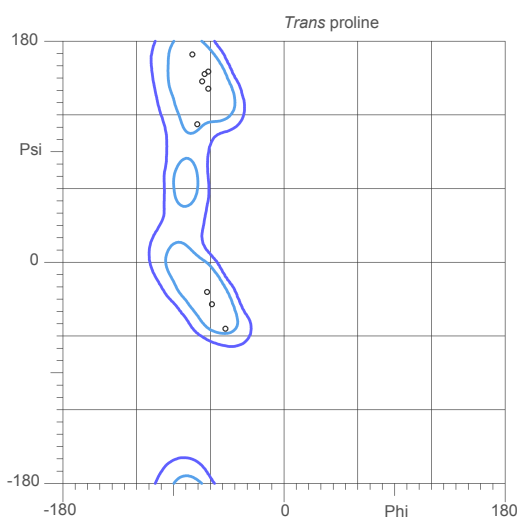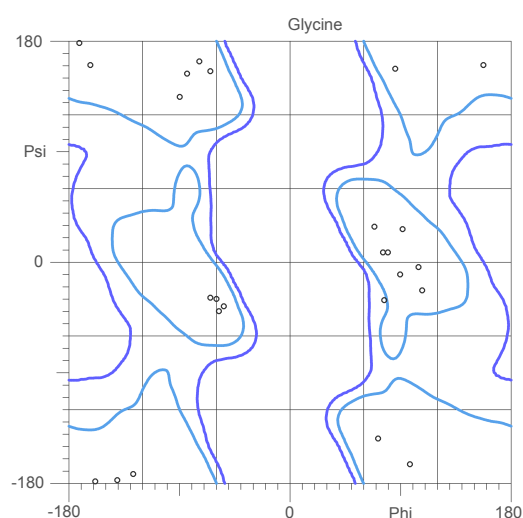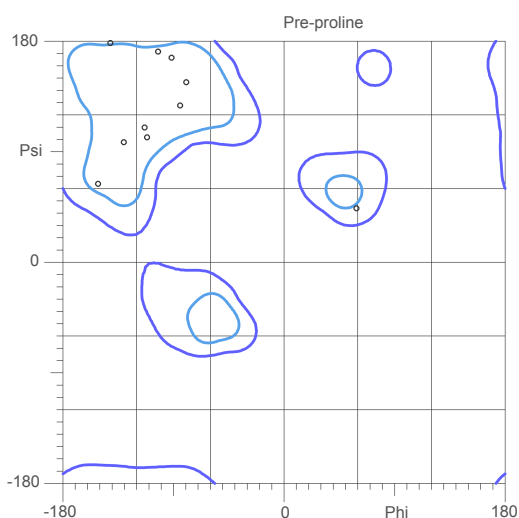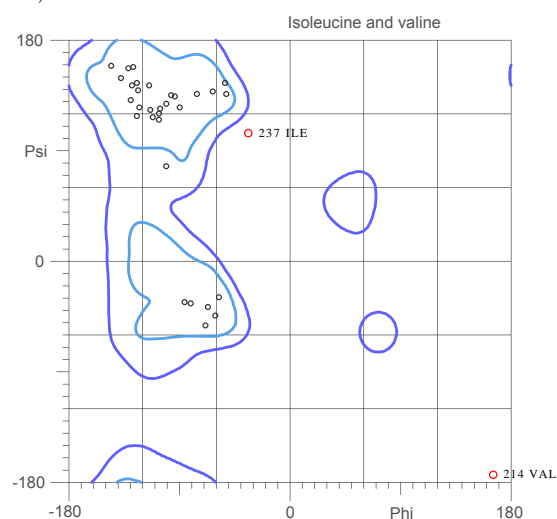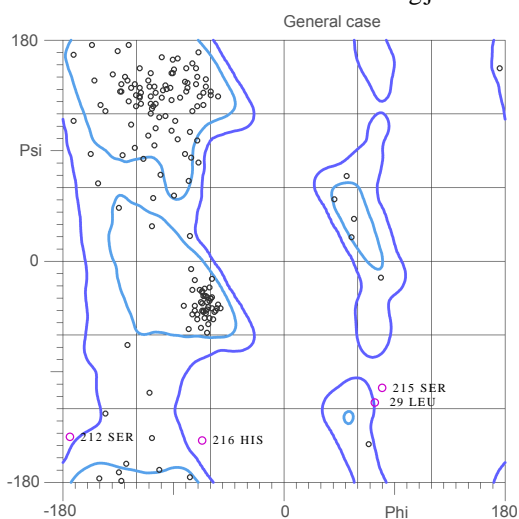

88,8% (213/240) of all residues were in favored (98%) regions.
97,5% (234/240) of all residues were in allowed (>99.8%) regions.

There were 6 outliers (phi, psi):
    29 LEU (74,4, -115,5)
    212 SER (-175,8, -143,0)
    214 VAL (166,9, -174,6)
    215 SER (80,1, -103,9)
    216 HIS (-67,6, -146,3)
    237 ILE (-34,7, 105,5)

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