MolProbity Ramachandran analysis

1ajsA.B99990036.pdb, model 1
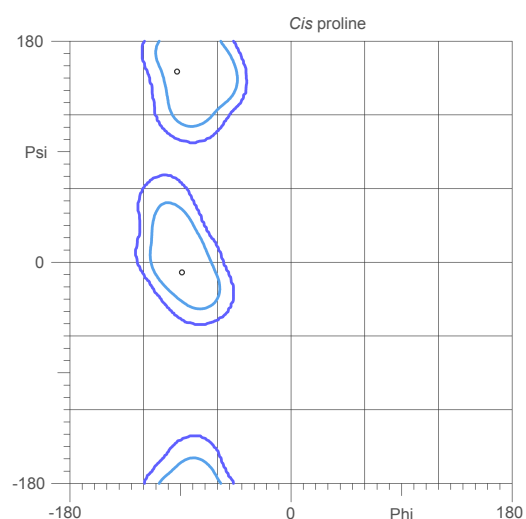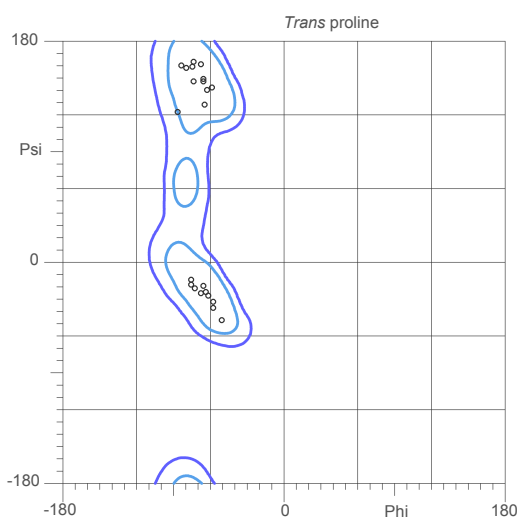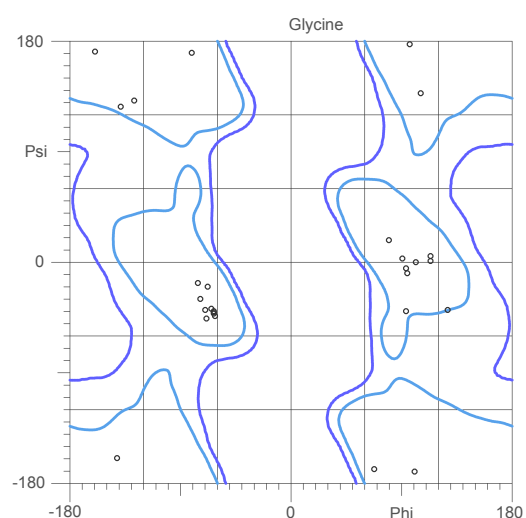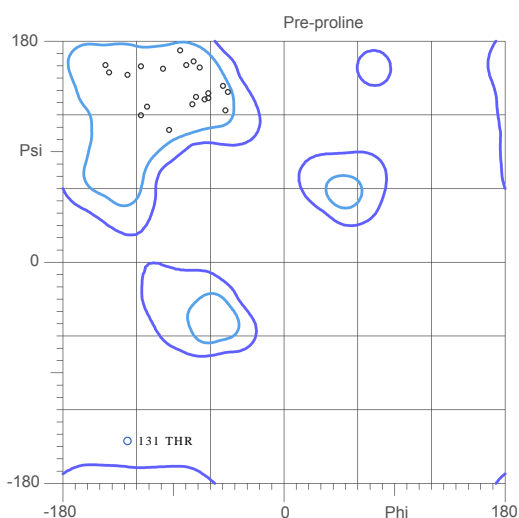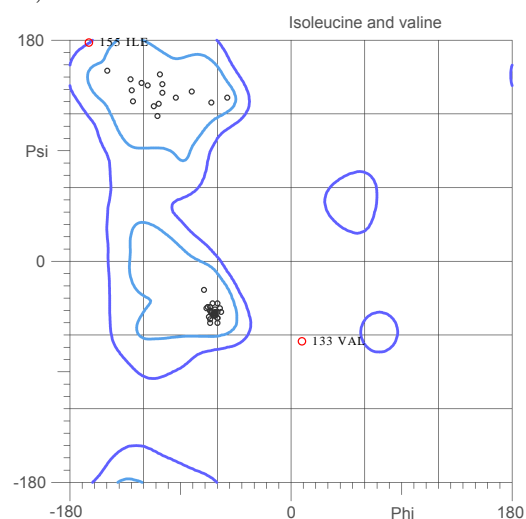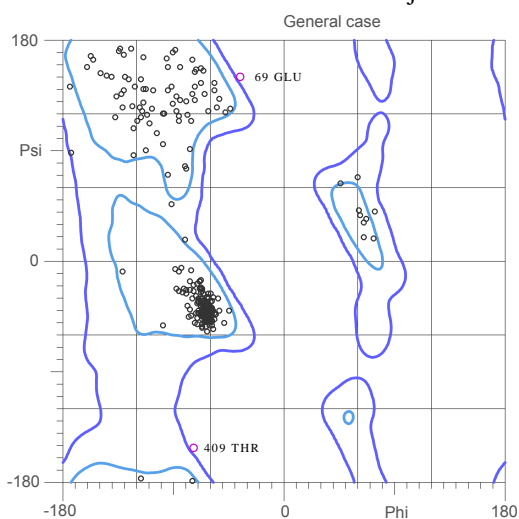

96,6% (396/410) of all residues were in favored (98%) regions.
98,8% (405/410) of all residues were in allowed (>99.8%) regions.

There were 5 outliers (phi, psi):
    69 GLU (-36,2, 151,4)
    131 THR (-128,8, -146,7)
    133 VAL (9,4, -65,9)
    155 ILE (-165,1, 179,4)
    409 THR (-74,2, -152,5)

http://kinemage.biochem.duke.edu

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