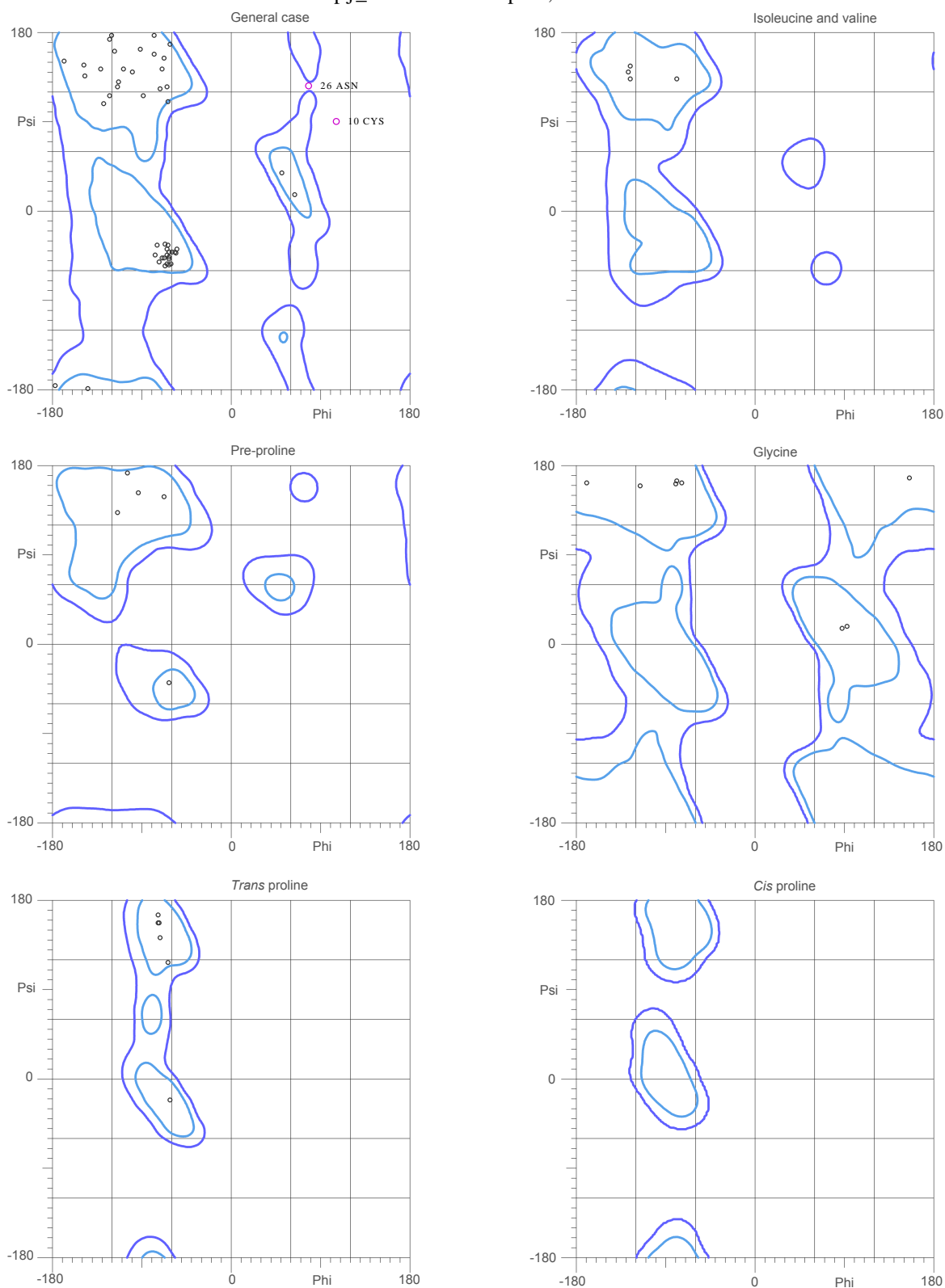# MolProbity Ramachandran analysis

1apj_.B99990025.pdb, model 1

95,8% (69/72) of all residues were in favored (98%) regions.

97,2% (70/72) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

    10 CYS (106,3, 91,3)

    26 ASN (78,8, 127,5)

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