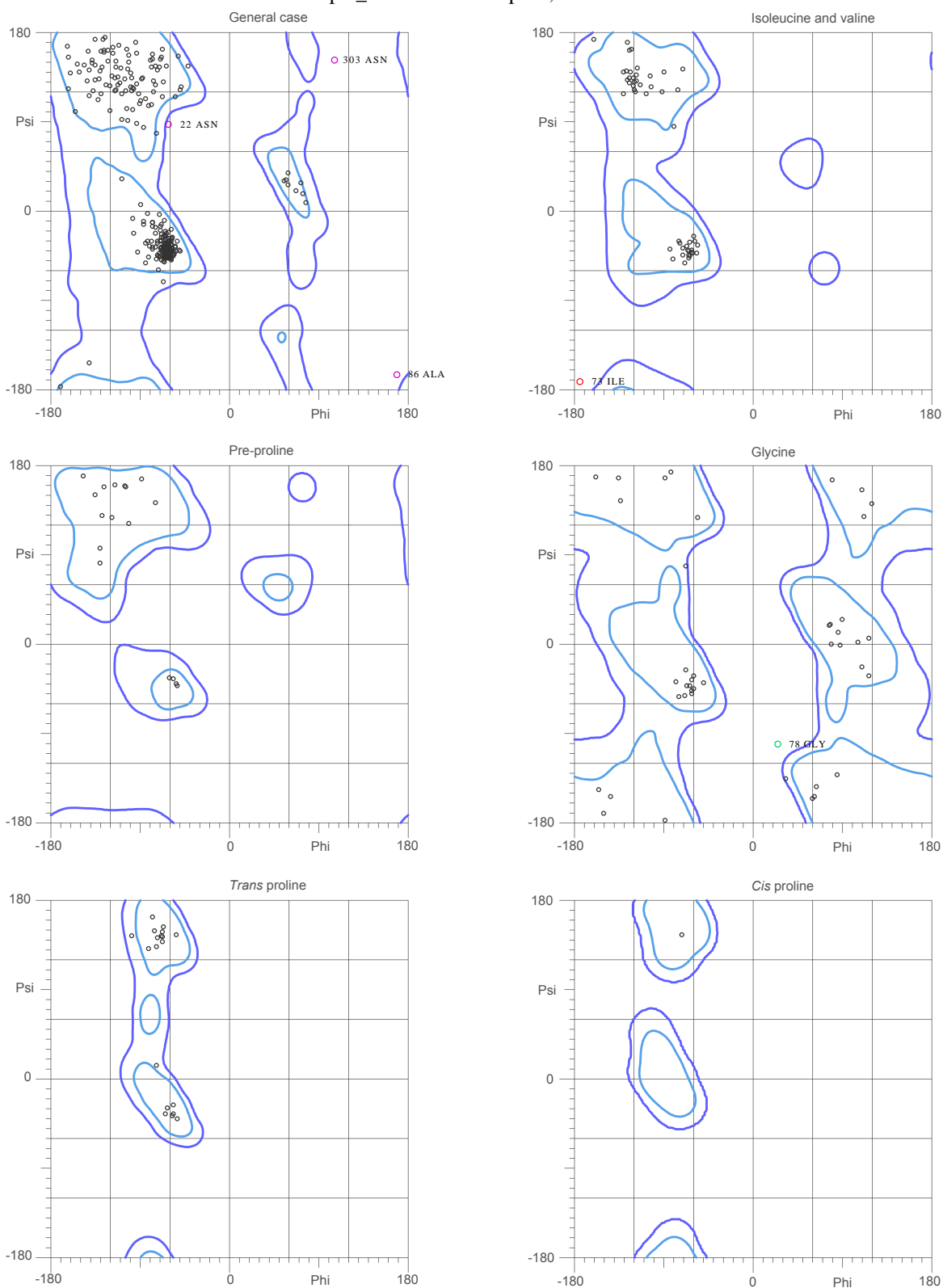# MolProbity Ramachandran analysis

16pk_.B99990033.pdb, model 1

95,9% (396/413) of all residues were in favored (98%) regions.
98,8% (408/413) of all residues were in allowed (>99.8%) regions.

There were 5 outliers (phi, psi):
    22 ASN (-62,1, 88,4)
    73 ILE (-175,9, -172,6)
    78 GLY (25,4, -101,6)
    86 ALA (169,1, -165,2)
    303 ASN (106,4, 153,5)

http://kinemage.biochem.duke.edu

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