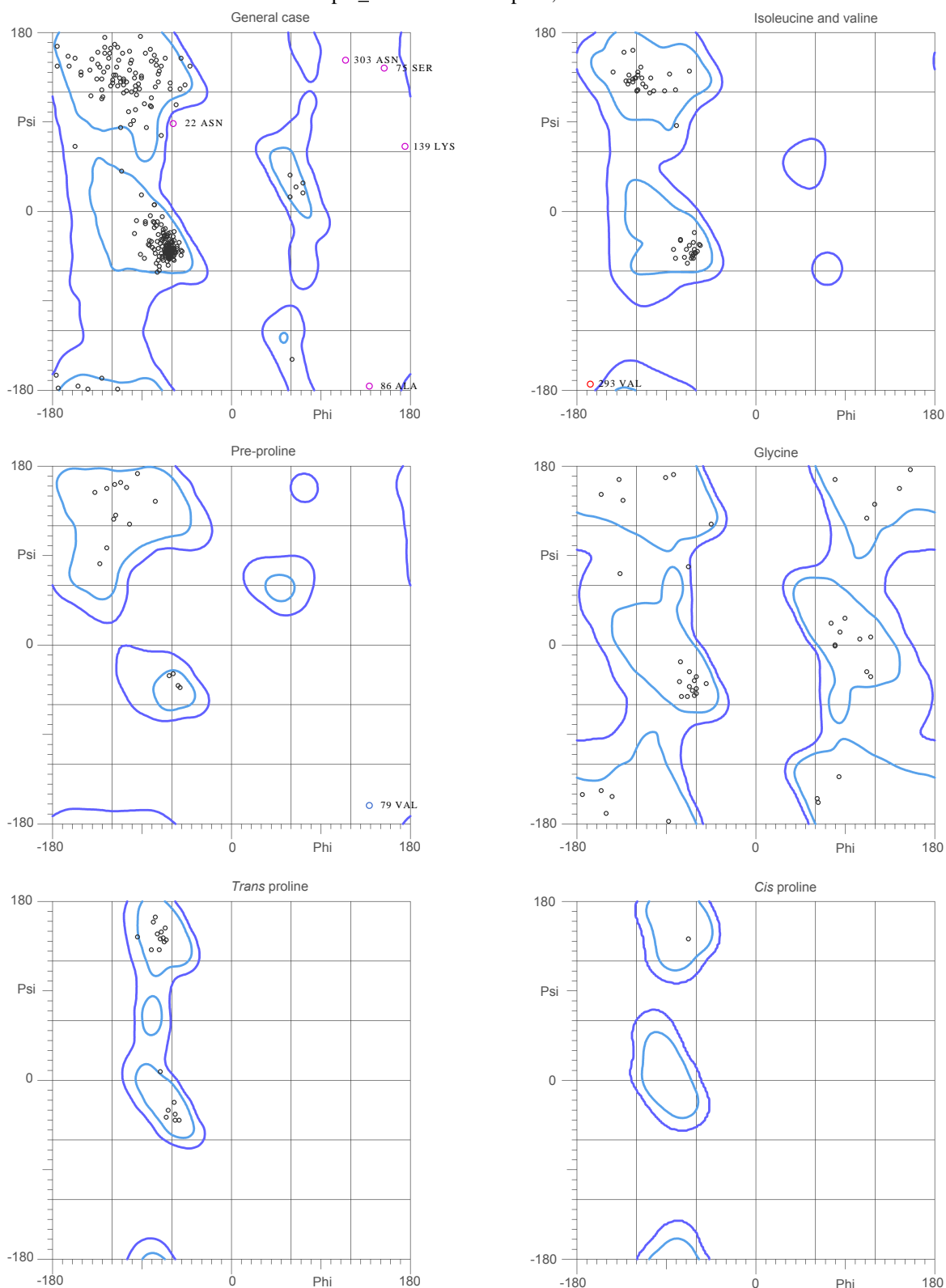# MolProbity Ramachandran analysis

16pk_.B99990022.pdb, model 1

94,2% (389/413) of all residues were in favored (98%) regions.
98,3% (406/413) of all residues were in allowed (>99.8%) regions.

There were 7 outliers (phi, psi):
    22 ASN (-59,5, 89,7)
    75 SER (154,5, 145,2)
    79 VAL (139,9, -162,7)
    86 ALA (139,1, -176,4)
    139 LYS (175,0, 67,0)
    293 VAL (-167,3, -174,5)
    303 ASN (115,8, 153,7)

http://kinemage.biochem.duke.edu

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