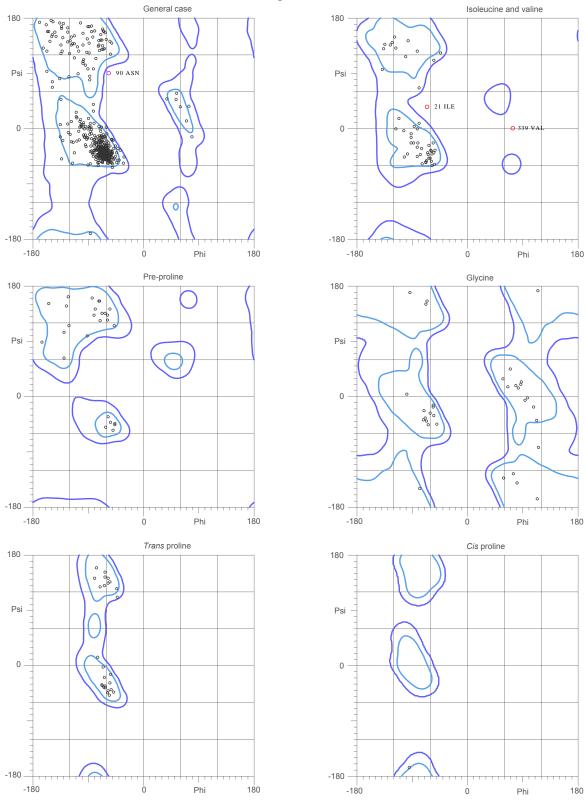
MolProbity Ramachandran analysis

1r42.H.pdb, model 1



94.1%~(560/595) of all residues were in favored (98%) regions. 99.5%~(592/595) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi): 21 ILE (-66.6, 36.0) 90 ASN (-57.4, 91.8) 339 VAL (74.1, 1.4)

http://kinemage.biochem.duke.edu