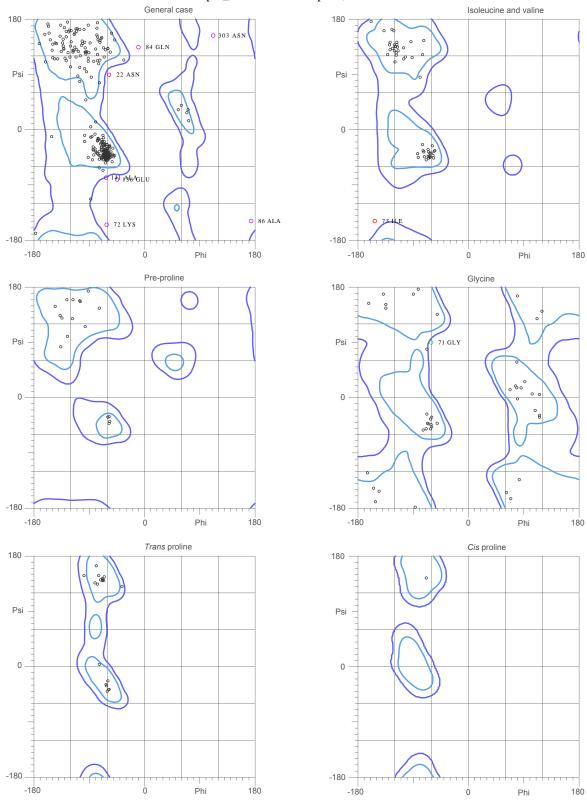
MolProbity Ramachandran analysis

16pk_.B99990041.pdb, model 1



93,7%~(387/413) of all residues were in favored (98%) regions

97,8% (404/413) of all residues were in allowed (>99.8%) regions

There were 9 outliers (phi, psi): 22 ASN (-58,6, 90,7) 71 GLY (-61,4, 90,8) 72 LYS (-62,5, -155,6) 73 ILE (-153,3, -149,4) 84 GLN (-10,7, 135,7) 86 ALA (174,2, -149,6) 136 GLU (-45,2, -81,7) 141 ALA (-63,6, -78,8) 303 ASN (112,8, 154,3)