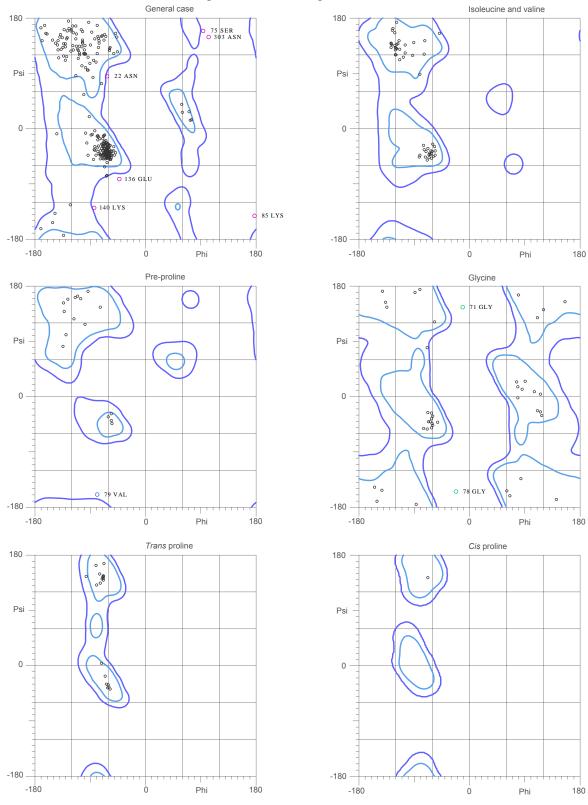
MolProbity Ramachandran analysis

16pk_.B99990008.pdb, model 1



93,9%~(388/413) 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 (-63,6,86,7) 71 GLY (-11,9,146,3) 75 SER (94,5,160,3) 78 GLY (-22,5,-155,7) 79 VAL (-80,0,-160,7) 85 LYS (178,7,-142,8) 136 GLU (-43,5,-82,3) 140 LYS (-84,4 ± 1.79 8) 140 LYS (-84,4, -129,8) 303 ASN (103,4, 150,2)