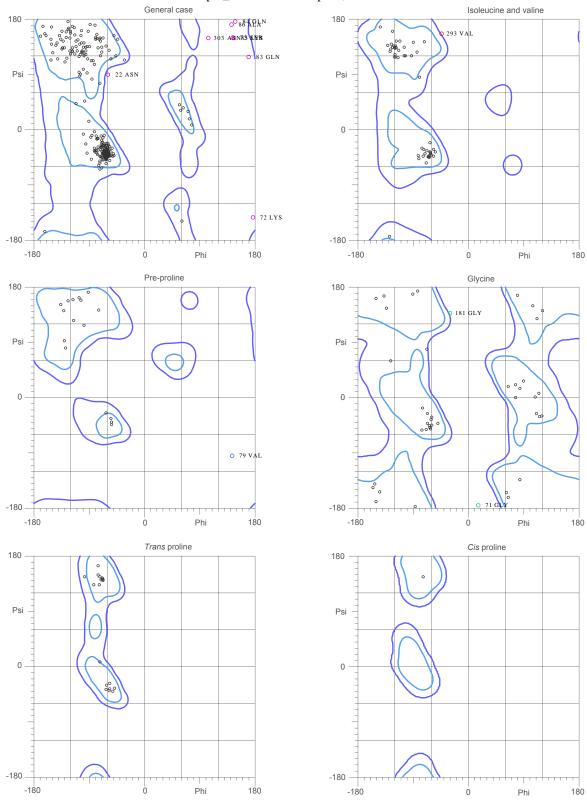
MolProbity Ramachandran analysis

16pk_.B99990034.pdb, model 1



93,0% (384/413) of all residues were in favored (98%) regions. 97,1% (401/413) of all residues were in allowed (>99.8%) regions

97,1% (401/413) of all residues were in allowed (>99.8%) region

7,1% (40/1413) of all residues we'here were 12 outliers (phi, psi):
22 ASN (60,8, 90,5)
71 GLY (16,9, -176,4)
72 LYS (173, -143,4)
75 SER (144,8, 151,2)
79 VAL (143,2, 95,7)
83 GLN (170,5, 119,1)
84 GLN (144, 173,8)
85 LYS (146,3, 150,9)
86 ALA (142,9, 172,5)
181 GLY (30,7, 138,3)
293 VAL (-44,4, 157,1)
303 ASN (104,6, 150,8)