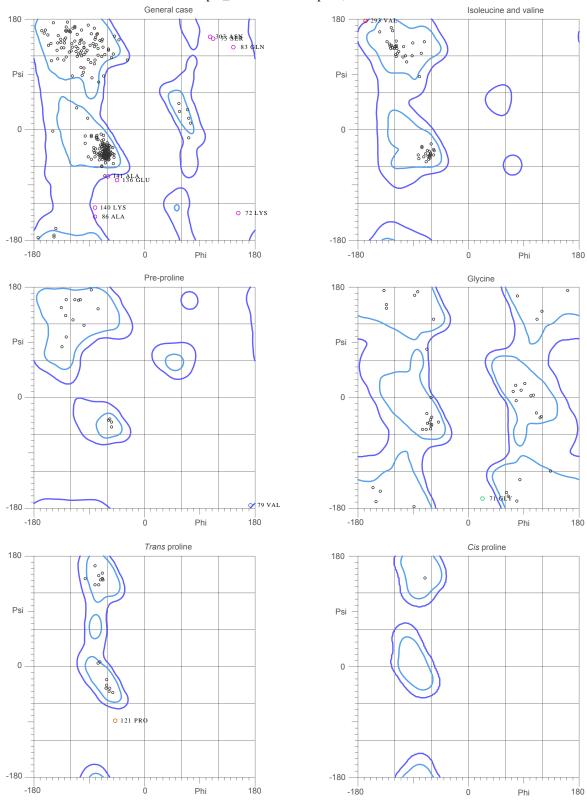
MolProbity Ramachandran analysis

16pk_.B99990017.pdb, model 1



93,7% (387/413) of all residues were in favored (98%) regions. 97,1% (401/413) of all residues were in allowed (>99.8%) regions

,1% (401/413) of all residues we see were 12 outliers (phi, psi): 71 GLY (23.8, -165.1) 72 LYS (153.4, -136.1) 75 SER (112, 149.2) 79 VAL (173.7, -1476.2) 83 GLN (145.7, 157.6) 86 GLA (-81.8, -142.6) 121 PRO (-48.7, -88.0) 136 GLU (4-5.7, -88.27) 140 LYS (-81.2, -127.7) 141 ALA (-60.4, -76.6) 293 VAL (-160.0, 178.8) 303 ASN (107.0, 152.2)