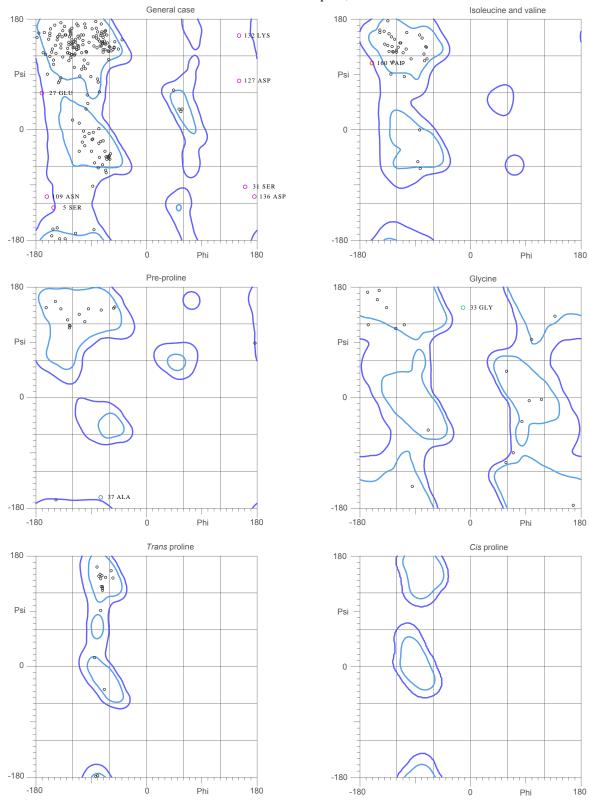
MolProbity Ramachandran analysis

1a02N.B99990006.pdb, model 1



86,3% (240/278) of all residues were in favored (98%) regions. 96,4% (268/278) of all residues were in allowed (>99.8%) regions

90,4% (200/2/8) of all residues were in allowed (>99.8%) region

There were 10 outliers (phi, psi):
5 SER (-152,6, -127,7)
27 GLU (-171,6, 60.1)
31 SER (161,6, -93,6)
33 GLY (-12,7, 147,9)
37 ALA (-75,7, -163,8)
109 ASN (-163,6, -109,1)
127 ASP (151,9, 80,3)
132 LYS (151,6, 154,4)
136 ASP (176,5, -109,5)
160 VAL (-161,2, 109,3)