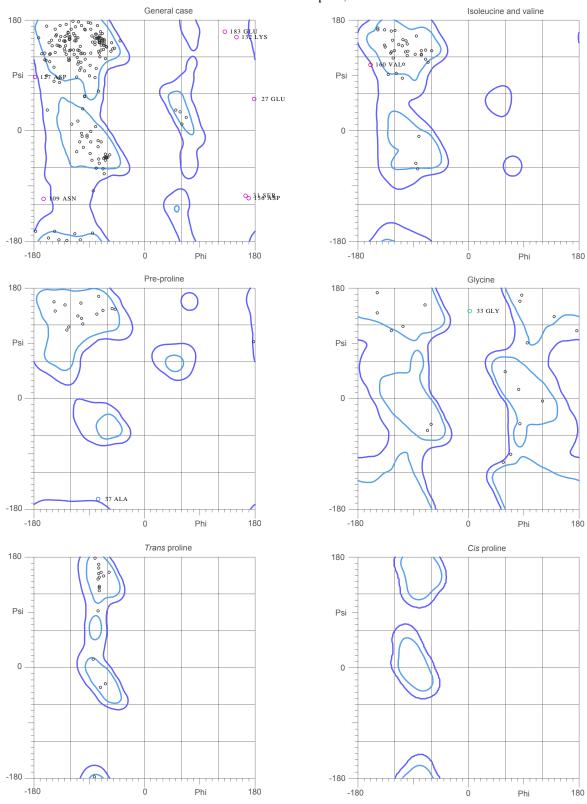
MolProbity Ramachandran analysis

1a02N.B99990002.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

There were 10 outliers (phi, psi):
27 GLU (179,4, 52,8)
31 SER (165,8, -106,4)
33 GLY (20, 143,6)
37 ALA (-76,2, -164,2)
109 ASN (-165,4, -111,2)
127 ASP (-179,4, 88,4)
132 LYS (150,9, 153,5)
136 ASP (170,5, -110,0)
160 VAL (-160,4, 108,6)
183 GLU (131,2, 163,0)