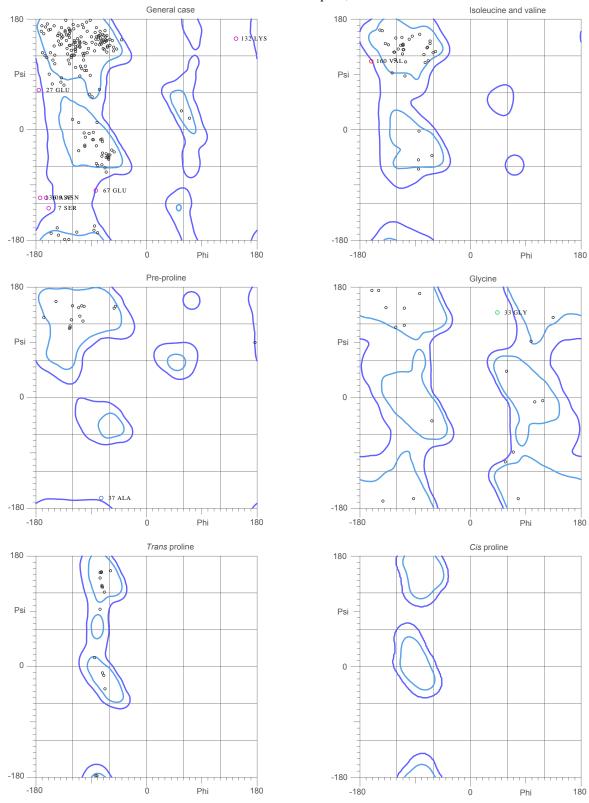
MolProbity Ramachandran analysis

1a02N.B99990013.pdb, model 1



88,8%~(247/278) of all residues were in favored (98%) regions

96,8% (269/278) of all residues were in allowed (>99.8%) regions

There were 9 outliers (phi, psi):
7 SER (-160,0, -128,6)
27 GLU (-176,9,65,3)
33 GLY (444, 139,2)
37 ALA (-74,3, -164,6)
67 GLU (-83,3, -99,9)
109 ASN (-164,7, -111,9)
132 LYS (146,6, 149,6)
136 ASP (-174,0, -112,0) 136 ASP (-174,0, -112,0) 160 VAL (-162,5, 112,5)