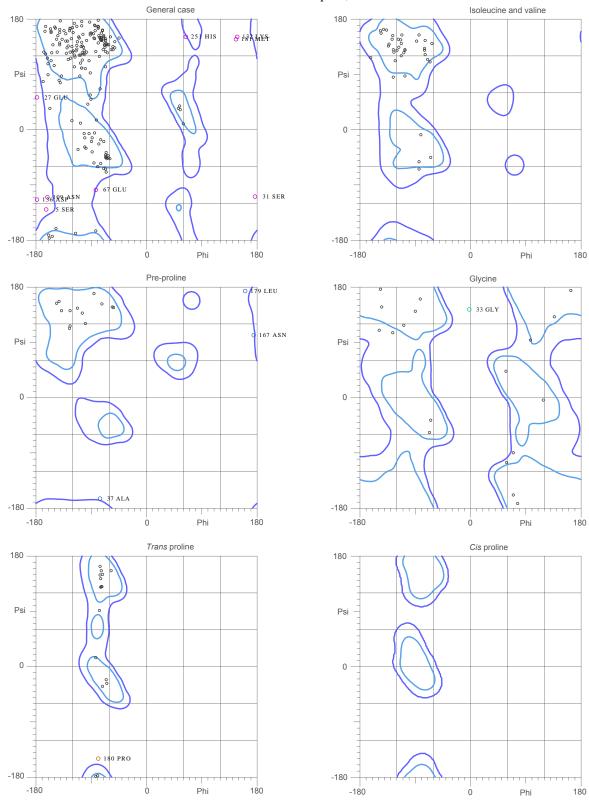
MolProbity Ramachandran analysis

1a02N.B99990016.pdb, model 1



83,8% (233/278) of all residues were in favored (98%) regions. 95.0% (264/278) of all residues were in allowed (>99.8%) region

There were 14 outliers (phi, psi): 5 SER (-164,8, -130,5) 27 GLU (-179,5, 53,0) 31 SER (177,5, -109,4)

27 GLU (-179,5, 53,0) 31 SER (177,5, -109,4) 33 GLY (-2.5, 145,0) 37 ALA (-76,1, -165,0) 67 GLU (-83,2, -98,8) 109 ASN (-162,7, -110,0) 132 LYS (148,7, 152,2) 136 ASP (-179,8, -114,7) 167 ASN (175,4, 102,1) 179 LEU (161,6, 174,3)