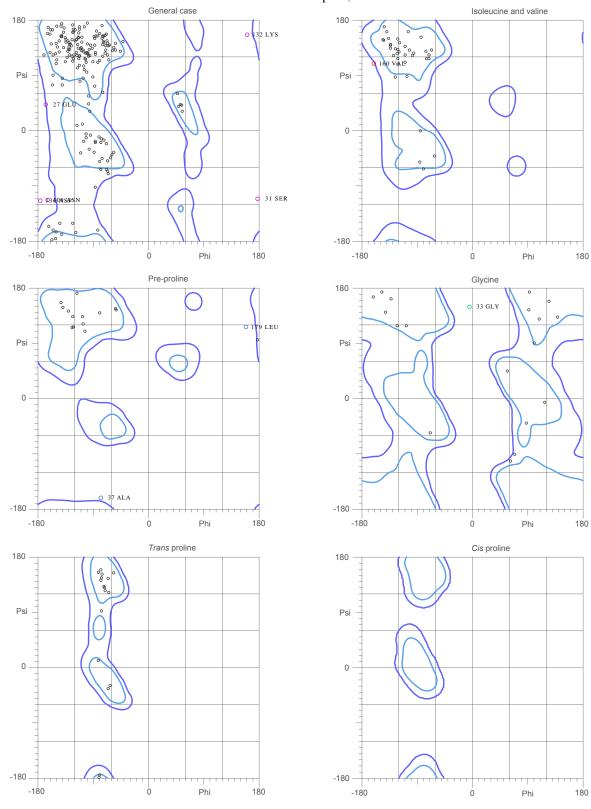
MolProbity Ramachandran analysis

1a02N.B99990034.pdb, model 1



87,1%~(242/278) 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): 27 GLU (-168,9, 43,2) 31 SER (178,8, -111,1) 33 GLY (-5.0, 150,1) 37 ALA (-78,1, -162,5) 109 ASN (-165,4, -113,0) 132 LYS (161,6, 158,0) 136 ASP (-177,6, -114,3) 160 VAI (-161,8, 110,1) 160 VAL (-161,8, 110,1) 179 LEU (159,1, 117,4)