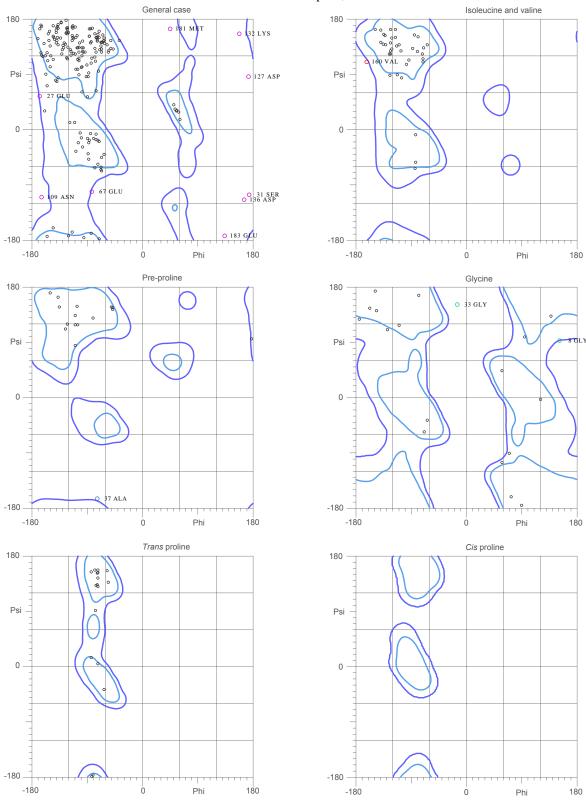
MolProbity Ramachandran analysis

1a02N.B99990048.pdb, model 1



85,6% (238/278) of all residues were in favored (98%) regions.

There were 13 outliers (phi, p 8 GLY (1524, 93.0) 27 GLU (-168.7, 55.0) 31 SER (175.0, -106.5) 33 GLY (-15.7, 152.5) 37 ALA (-74.7, -165.0) 67 GLU (-83.2, -101.6) 109 ASN (-165.7, -110.8)

37 ALA (-74,7, -165,0) 67 GLU (-83,2, -101,6) 09 ASN (-165,7, -110,8) 27 ASP (173,6, 87,9) 32 LYS (158,3, 157,9) 36 ASP (166,2, -114,4) 60 VAL (-163,4, 111,3) 81 MET (45,5, 165,6)