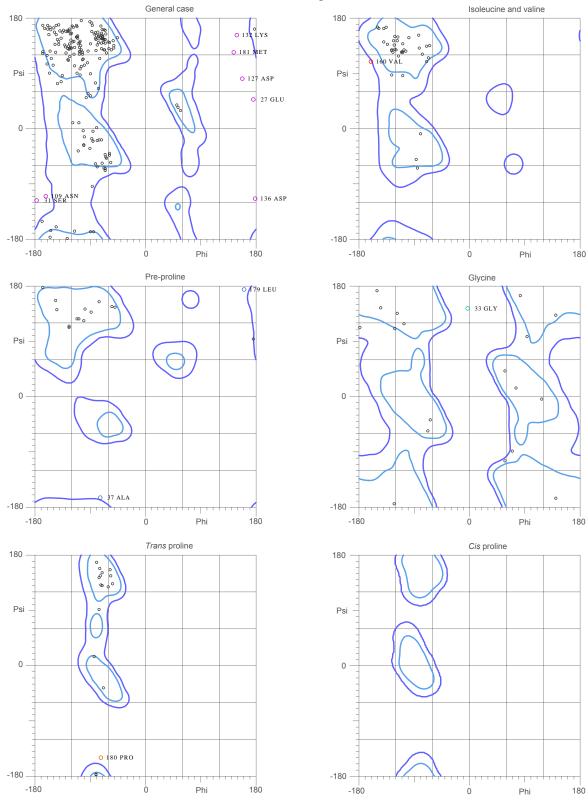
MolProbity Ramachandran analysis

1a02N.B99990030.pdb, model 1



85,6% (238/278) of all residues were in favored (98%) regions. 95,7% (266/278) of all residues were in allowed (>99.8%) regions

7.% (260.78) of all residues we see were 12 outliers (phi, psi): 27 GLU (176.3, 49.0) 31 SER (178.3, -117.5) 33 GLY (-3.3, 144.9) 37 ALA (-74.3, -165.5) 109 ASN (-163.9, -110.6) 127 ASP (158.9, 82.2) 132 LYS (149.8, 154.0) 136 ASP (179.9, -114.5) 160 VAL (-161.5, 110.9) 179 LEU (162.0, 175.5) 180 PRO (-73.9, -150.3) 181 MET (144.2, 126.0)