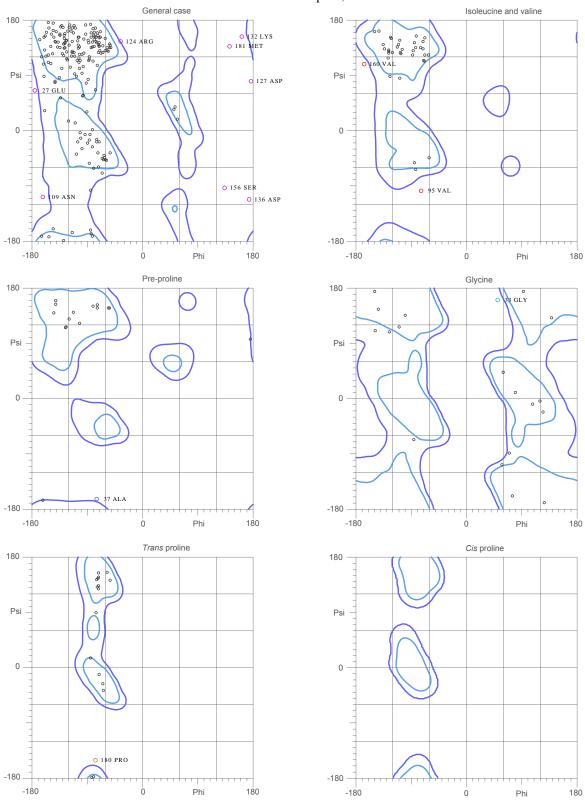
MolProbity Ramachandran analysis

1a02N.B99990004.pdb, model 1



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

here were 13 outliers (phi, 27 GLU (-176,2, 66,4) and GLU (-176,2, 66,4) and GLU (-176,2, 66,4) and GLU (-174,4,-98,5) and GLU (-174,4,-98,5) and GLU (-174,4,-98,5) and GLU (-174,4,-18,18) and GLU (-174,4,-18,18) and GLU (-174,4,-113,0) and GLU (