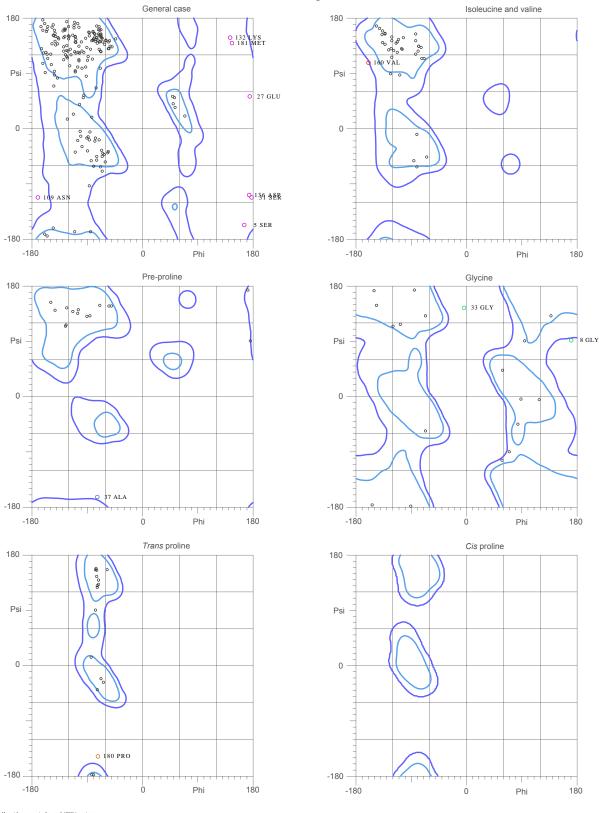
MolProbity Ramachandran analysis

1a02N.B99990044.pdb, model 1



86,3% (240/278) of all residues were in favored (98%) regions.
95,7% (266/278) of all residues were in allowed (>99.8%) regions

7.8 (260.78) of all residues we see were 12 outliers (phi, psi): 3 SER (166.6. -157.6) S GLY (171.0. 92.5) S GLY (171.0. 92.5) 31 SER (178.4. -112.8) 33 GLY (-42.45.3) 37 GLA (-74.3. -164.6) 97 ASN (-172.0. -112.9) 132 LYS (143.5, 149.7) 136 ASP (74.1. 108.4) 160 VAL (-160.2. 108.4) 180 PRO (-73.0. -148.5) 181 MET (146.8. 140.9)