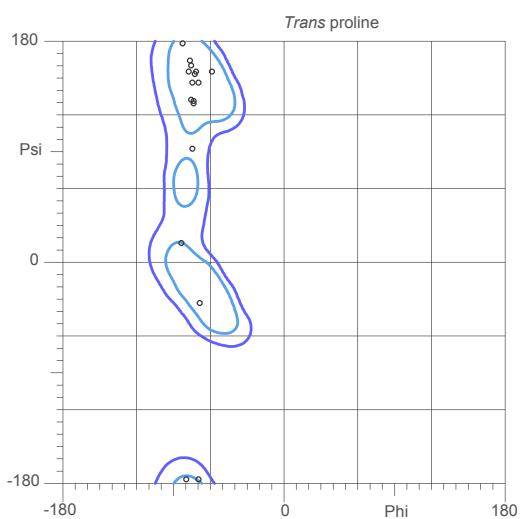
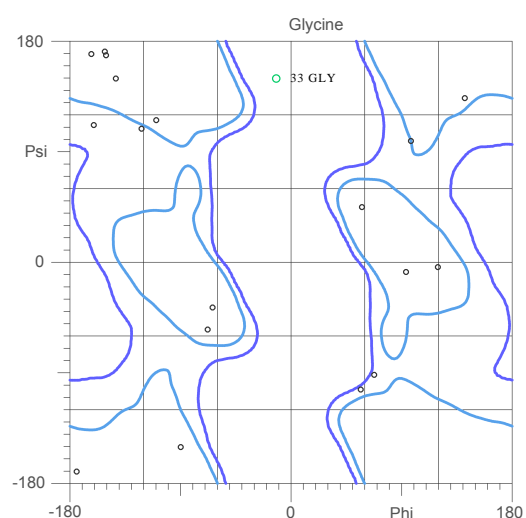
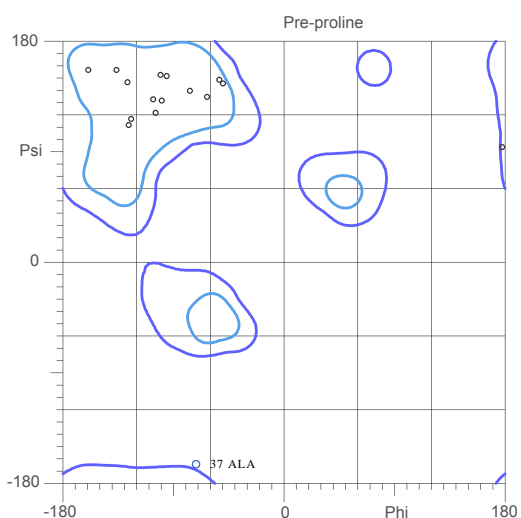
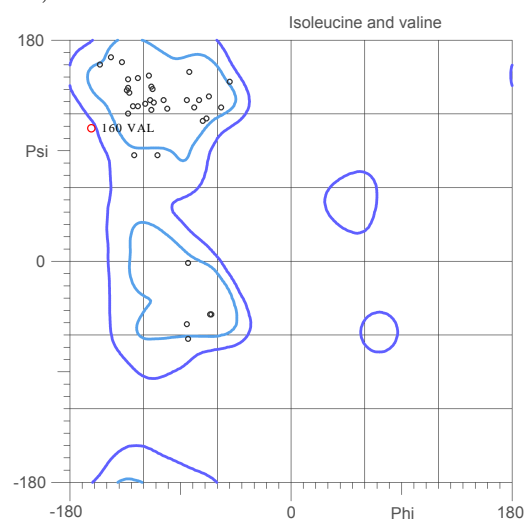
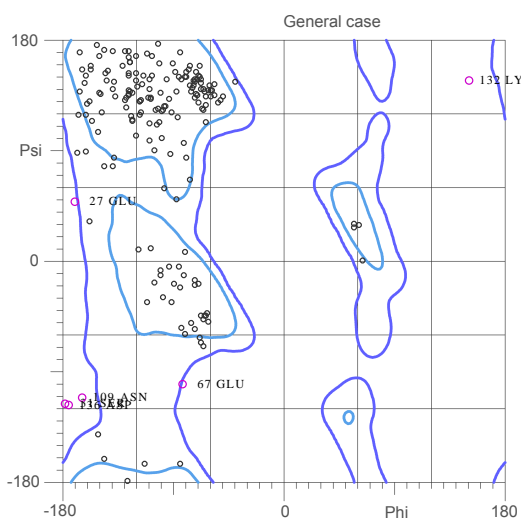


# MolProbity Ramachandran analysis

1a02N.B99990042.pdb, model 1



86.7% (241/278) 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 (-171.9, 49.3)  
31 SER (-179.4, -116.6)  
33 GLY (-12.9, 150.7)  
37 ALA (-72.5, -165.1)  
67 GLU (-83.4, -100.2)  
109 ASN (-165.7, -111.9)  
132 LYS (151.3, 148.6)  
136 ASP (-176.2, -117.3)  
160 VAL (-163.3, 109.3)