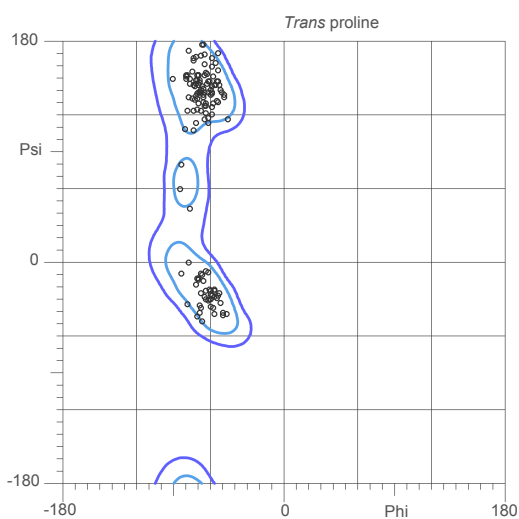
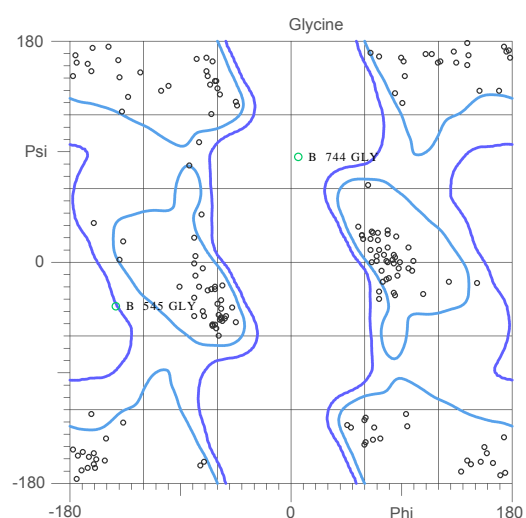
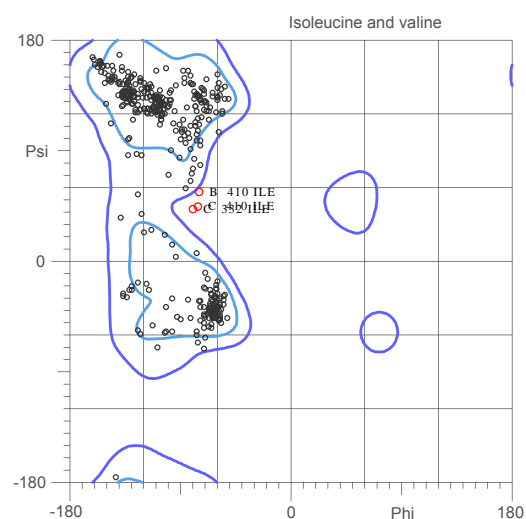


# MolProbity Ramachandran analysis

6x2b.H.pdb, model 1



92.9% (2581/2779) of all residues were in favored (98%) regions.  
99.6% (2767/2779) of all residues were in allowed (>99.8%) regions.

There were 12 outliers (phi, psi):

A 396 TYR (44.3, 83.7)  
B 41 LYS (-70.8, 21.9)  
B 46 SER (59.7, -17.8)  
B 410 ILE (-75.5, 57.6)  
B 545 GLY (-143.0, -36.4)  
B 744 GLY (6.2, 86.7)  
C 41 LYS (-70.0, 22.3)  
C 46 SER (57.4, -7.8)  
C 87 ASN (55.8, -83.3)  
C 123 ALA (60.0, -13.2)  
C 332 ILE (-81.0, 43.3)  
C 410 ILE (-76.5, 45.4)