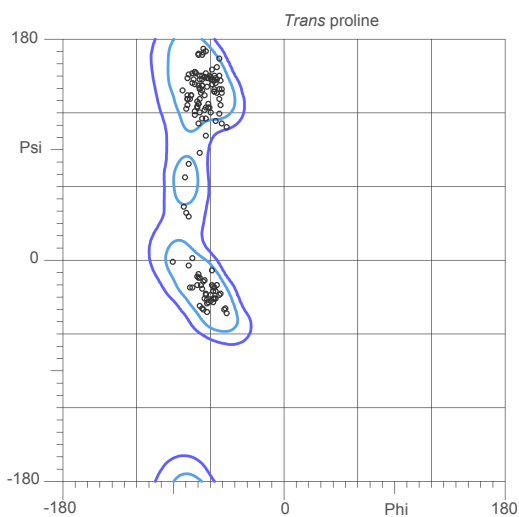
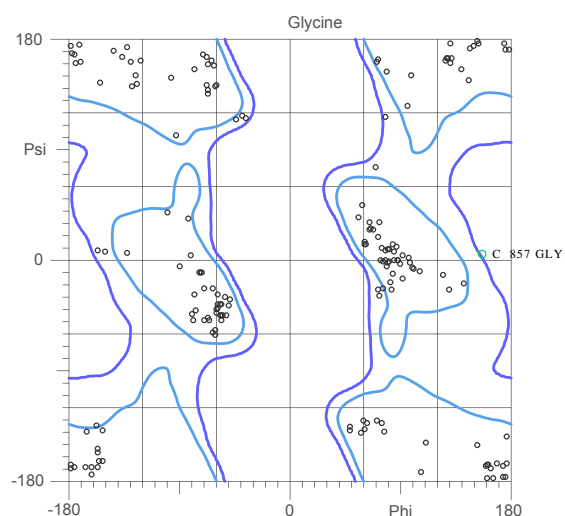
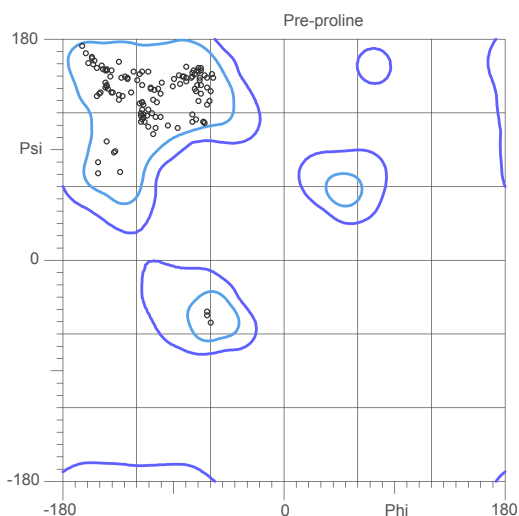
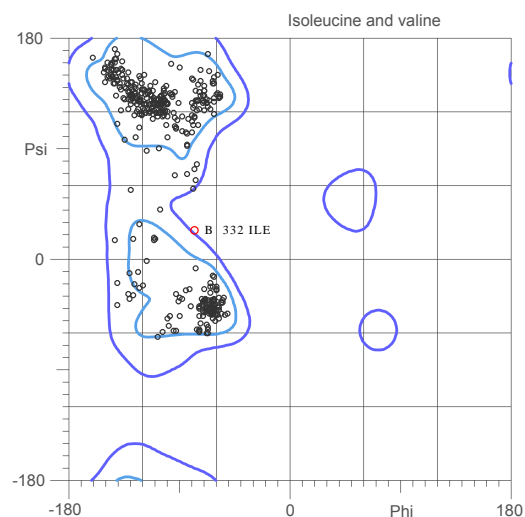
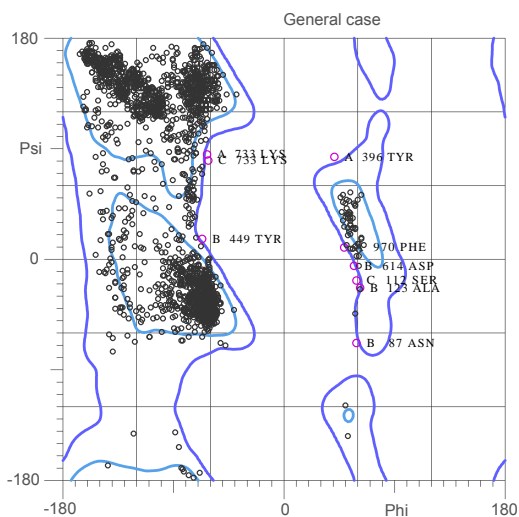


# MolProbity Ramachandran analysis

6x2a.H.pdb, model 1



93.3% (2610/2797) of all residues were in favored (98%) regions.  
99.6% (2786/2797) of all residues were in allowed (>99.8%) regions.

There were 11 outliers (phi, psi):

A 396 TYR (41.7, 84.0)  
A 733 LYS (-63.1, 87.0)  
A 970 PHE (49.7, 10.2)  
B 87 ASN (59.4, -68.6)  
B 123 ALA (62.1, -23.8)  
B 332 ILE (-78.1, 24.5)  
B 449 TYR (-67.8, 17.8)  
B 614 ASP (57.4, -5.0)  
C 112 SER (59.5, -17.4)  
C 733 LYS (-62.5, 81.0)  
C 857 GLY (157.5, 5.9)