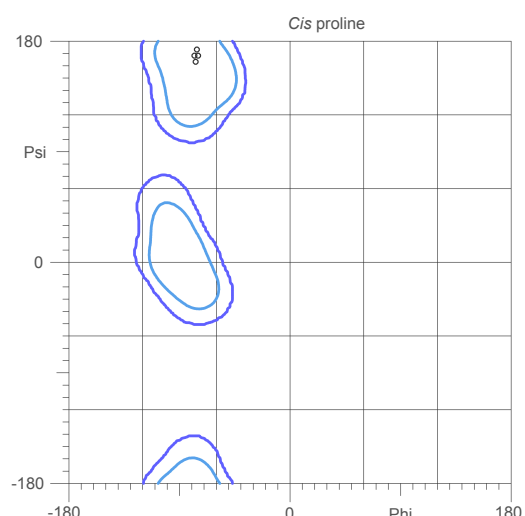
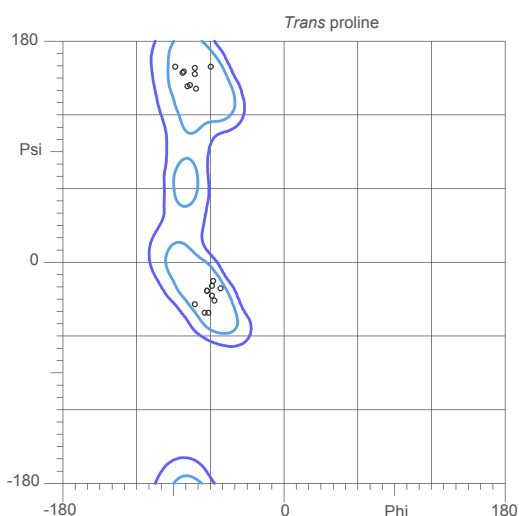
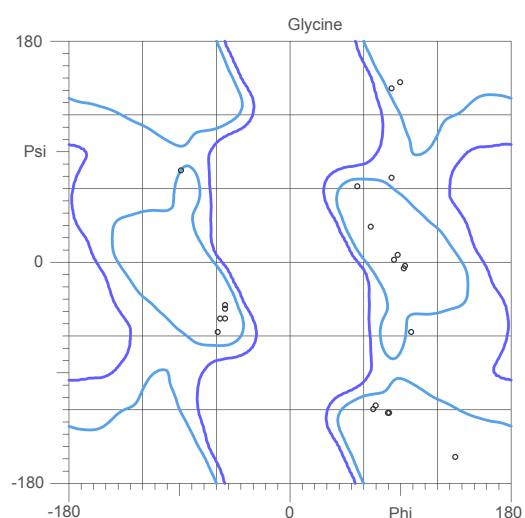
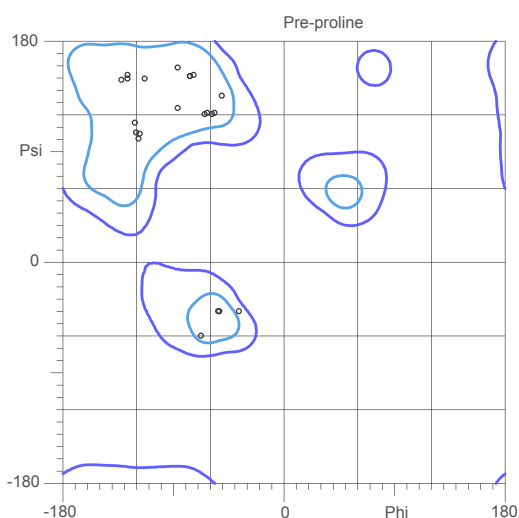
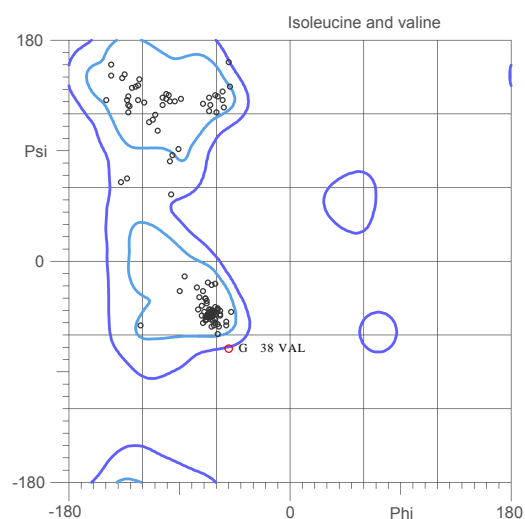
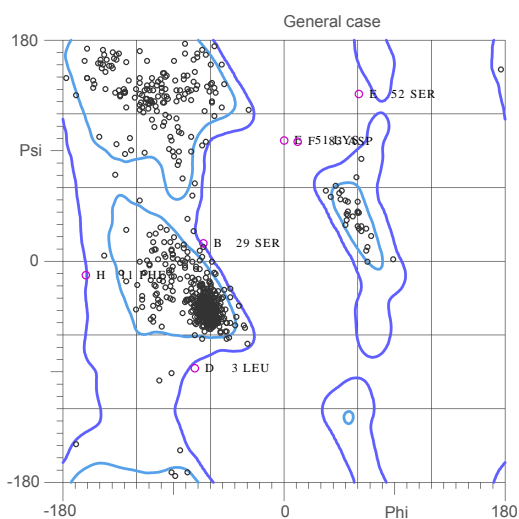


# MolProbity Ramachandran analysis

2ahm.H.pdb, model 1



93.8% (908/968) of all residues were in favored (98%) regions.  
99.3% (961/968) of all residues were in allowed (>99.8%) regions.

There were 7 outliers (phi, psi):

B 29 SER (-66.5, 15.6)  
D 3 LEU (-73.5, -87.4)  
E 51 LYS (0.7, 99.5)  
E 52 SER (61.8, 137.9)  
F 83 ASP (11.4, 98.1)  
G 38 VAL (-50.0, -71.6)  
H 11 PHE (-162.1, -11.4)