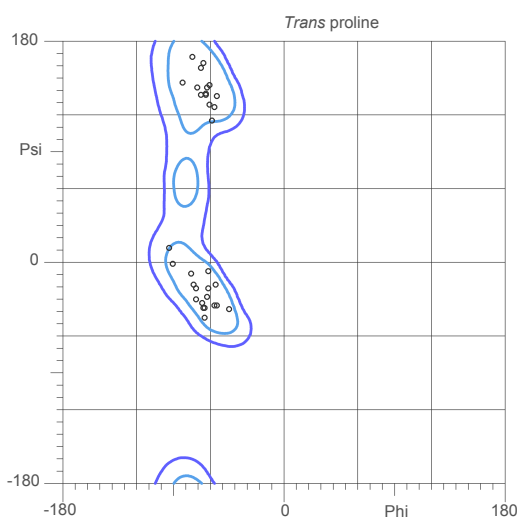
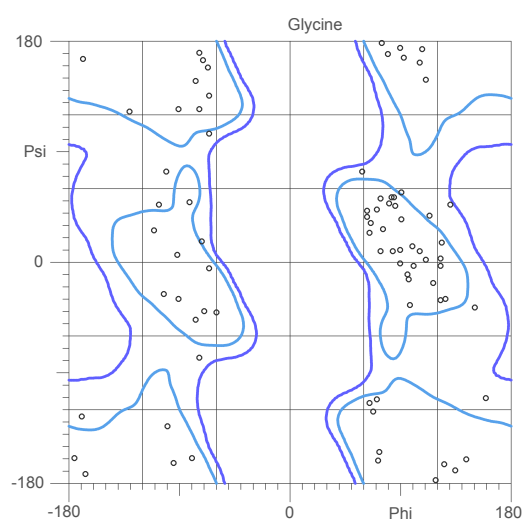
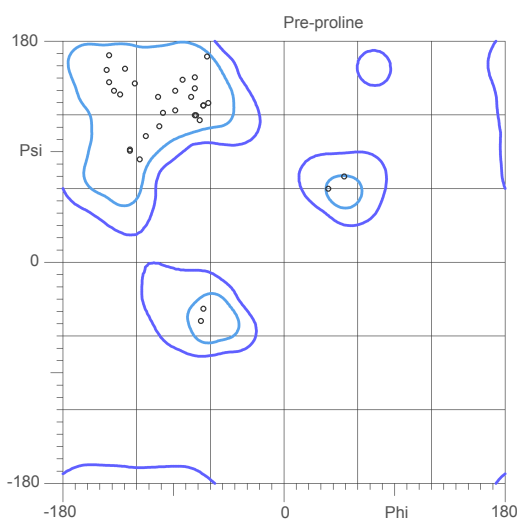
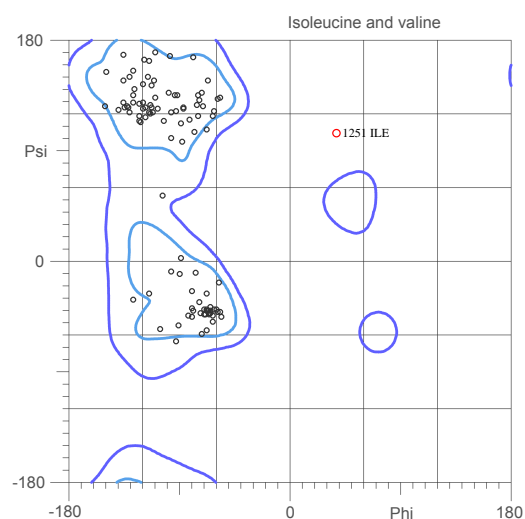
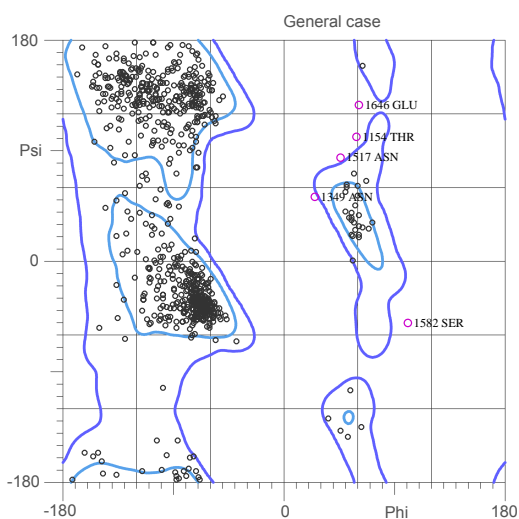


# MolProbity Ramachandran analysis

4amc.pdb, model 1



91.8% (934/1017) of all residues were in favored (98%) regions.  
99.4% (1011/1017) of all residues were in allowed (>99.8%) regions.

There were 6 outliers (phi, psi):

1154 THR (59.2, 102.7)  
1251 ILE (38.2, 105.4)  
1349 ASN (25.3, 53.9)  
1517 ASN (46.5, 85.6)  
1582 SER (101.4, -50.3)  
1646 GLU (61.8, 128.2)