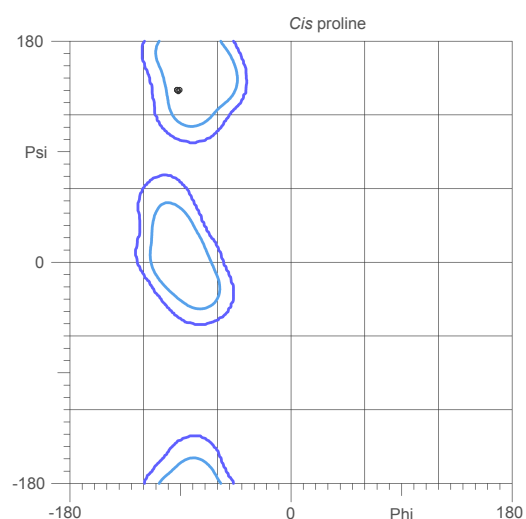
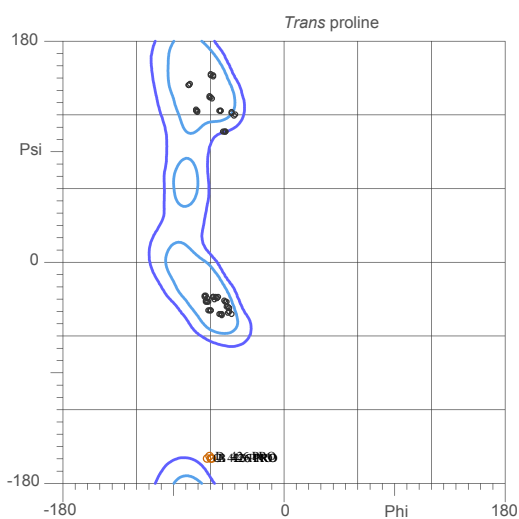
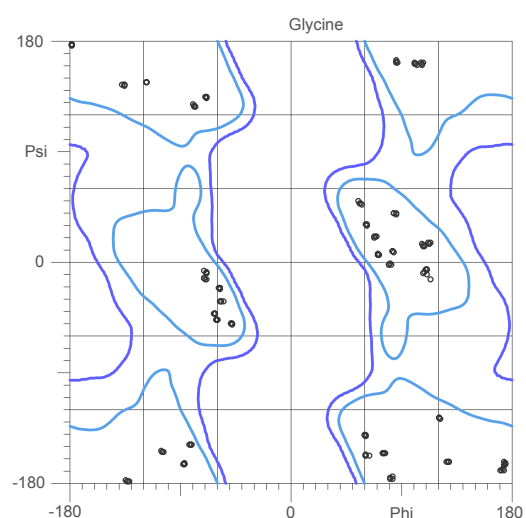
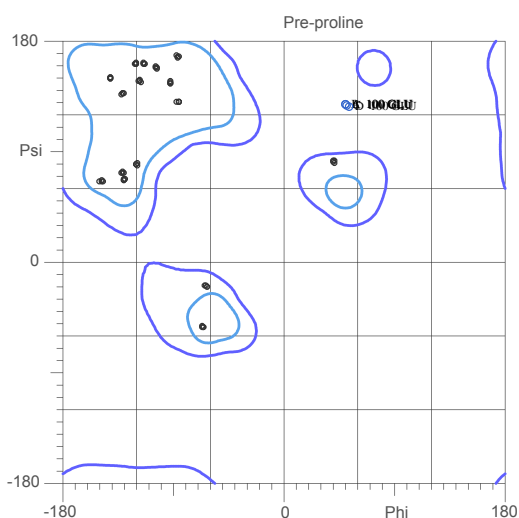
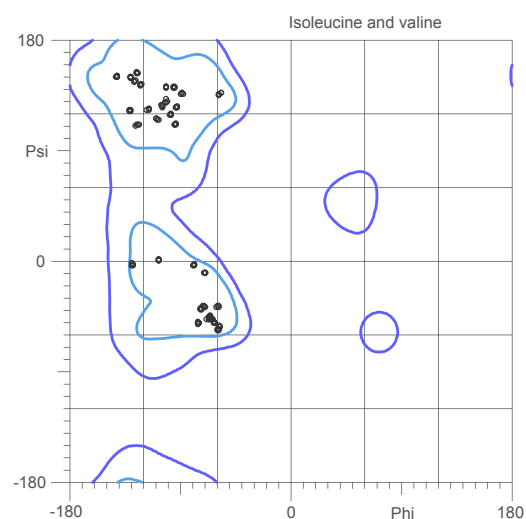
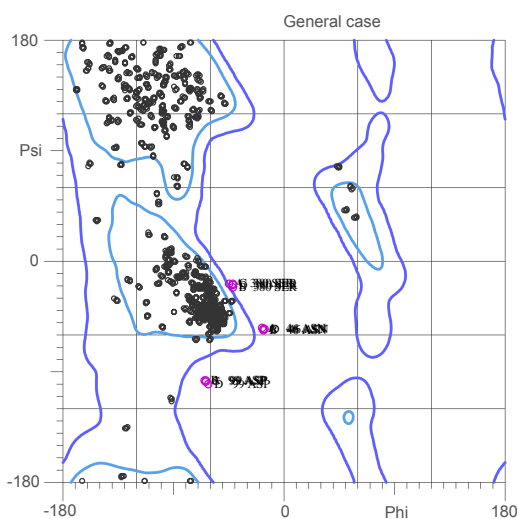


# MolProbity Ramachandran analysis

1ll4.pdb, model 1



94.6% (1476/1560) of all residues were in favored (98%) regions.  
98.7% (1540/1560) of all residues were in allowed (>99.9%) regions.

There were 20 outliers (phi, psi):

A: 46 ASN (-8.2, -54.6)  
A: 99 ASP (-64.1, -97.4)  
A: 100 GLU (-50.7, 120.0)  
A: 390 SER (-45.4, -18.1)  
A: 426 PRO (-60.8, -100.0)  
B: 46 ASN (-17.6, -56.2)  
B: 99 ASP (-66.0, -97.6)  
B: 100 GLU (-50.5, 120.0)  
B: 390 SER (-42.4, -21.2)  
B: 426 PRO (-59.6, -100.3)  
C: 46 ASN (-17.0, -55.7)  
C: 99 ASP (-64.5, -98.8)  
C: 100 GLU (-51.2, 128.6)  
C: 390 SER (-42.8, -10.6)  
C: 426 PRO (-63.2, -100.6)  
D: 46 ASN (-16.9, -55.9)  
D: 99 ASP (-62.5, -100.4)  
D: 100 GLU (-51.2, 127.9)  
D: 390 SER (-41.7, -19.9)  
D: 426 PRO (-61.7, -158.2)