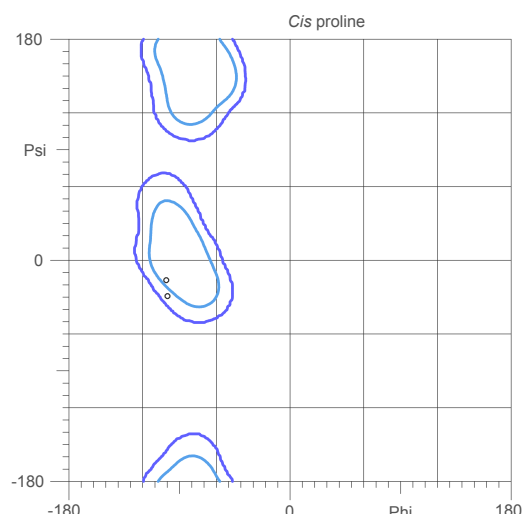
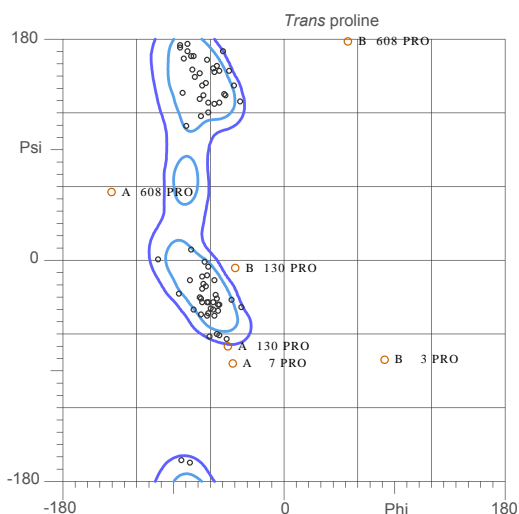
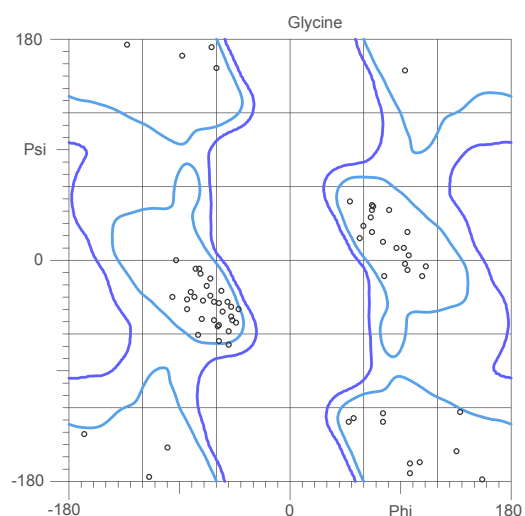
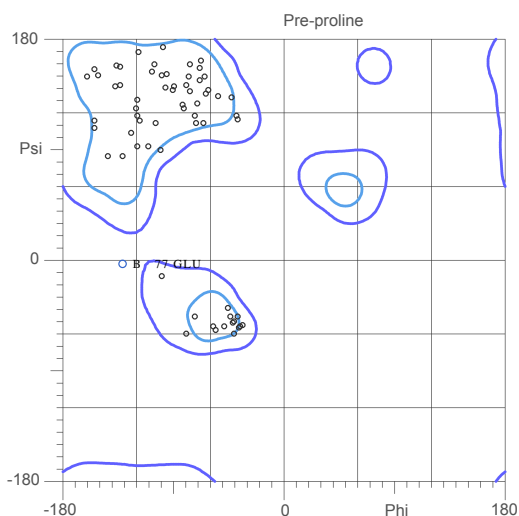
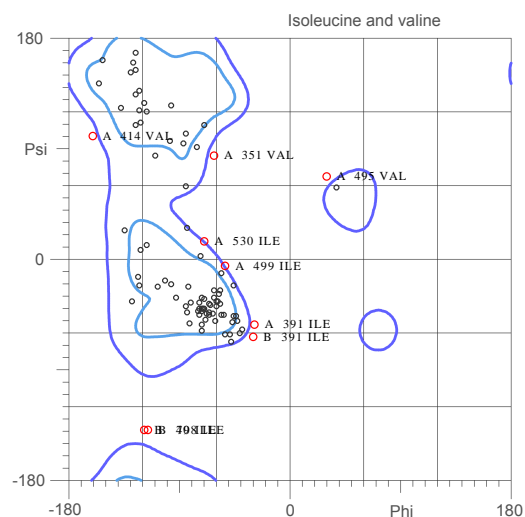
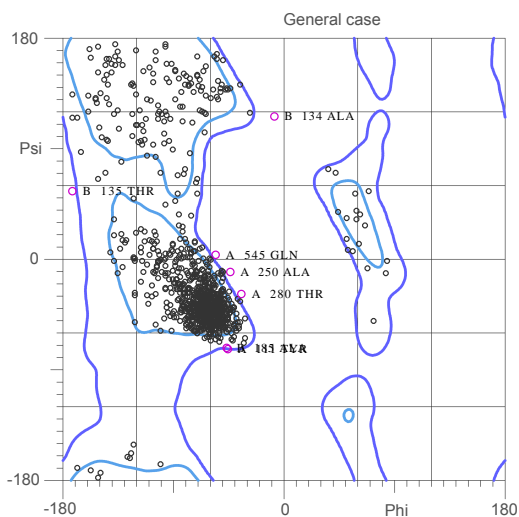


# MolProbity Ramachandran analysis

2c6f.H.pdb, model 1



87.2% (1062/1219) of all residues were in favored (98%) regions.  
98.1% (1196/1219) of all residues were in allowed (99.8%) regions.

There were 37 outliers (3.0%):

A 7 PRO (-42.3, -84.7)  
A 111 THR (-67.8, -75.5)  
A 130 PRO (-44.5, -78.9)  
A 200 ALA (-48.4, -100.9)  
A 280 THR (-15.5, -28.7)  
A 391 VAL (-82.7, -88.0)  
A 393 ILE (-101.8, -93.0)  
A 414 VAL (-101.8, -93.0)  
A 495 VAL (-109.7, -88.7)  
A 499 ILE (-151.1, -4.3)  
A 530 ILE (-78.9, -15.4)  
A 545 GLN (-56.4, -4.3)  
A 608 PRO (-141.5, -74.1)  
B 3 PRO (-82.9, -81.7)  
B 77 GLU (-132.4, -3.4)  
B 78 ILE (-120.8, -109.3)  
B 130 PRO (-41.0, -6.4)  
B 134 ALA (-48.4, -100.9)  
B 135 THR (-173.6, -56.4)  
B 185 ALA (-47.5, -75.2)  
B 391 ILE (-104.4, -84.0)  
B 408 ILE (-116.4, -109.8)  
B 408 PRO (-12.9, -179.3)