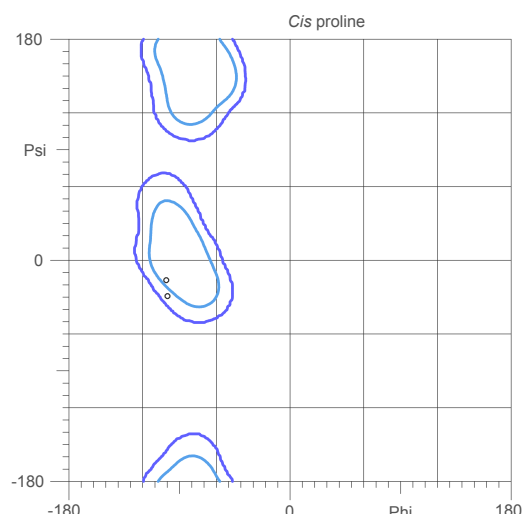
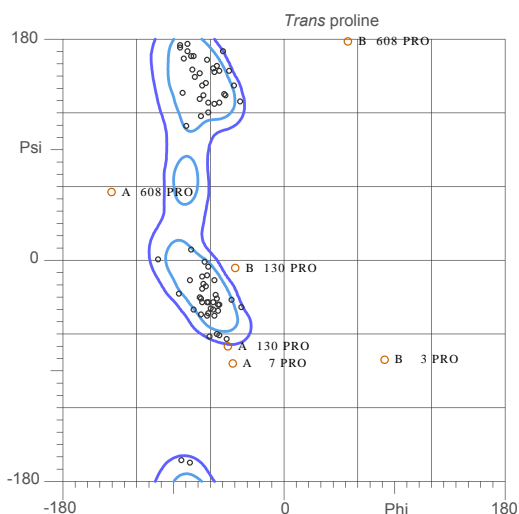
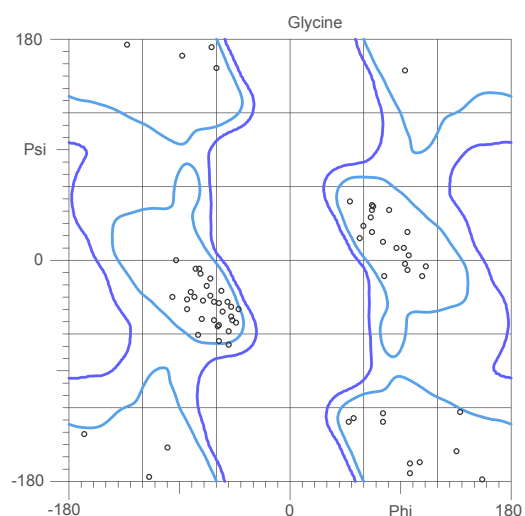
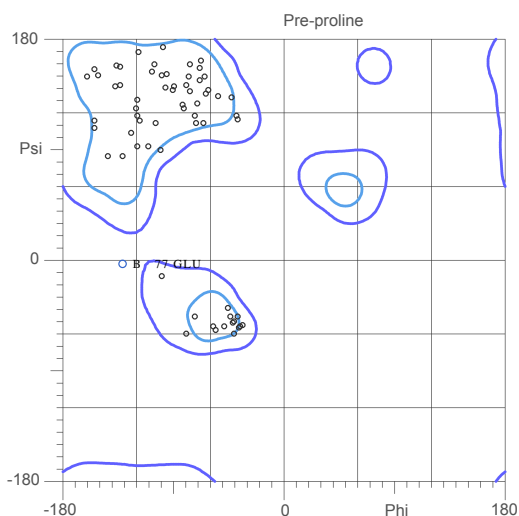
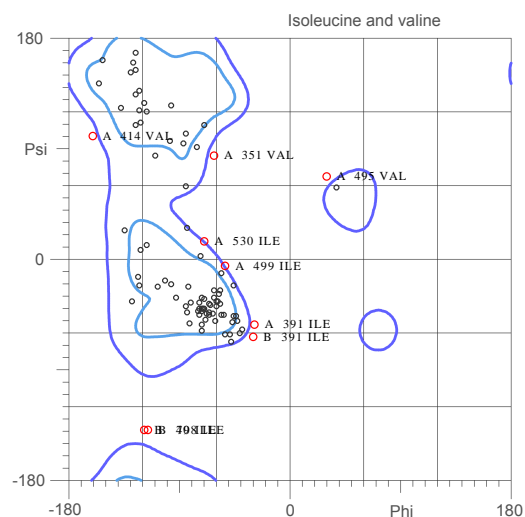
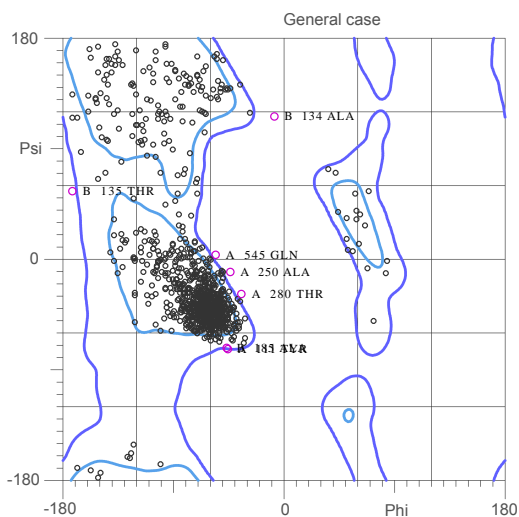


# MolProbity Ramachandran analysis

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, -86.0)  
A 391 ILE (-101.8, -93.0)  
A 414 VAL (-101.8, -93.0)  
A 495 VAL (-101.8, -93.0)  
A 495 ILE (-101.8, -93.0)  
A 530 ILE (-101.8, -93.0)  
A 545 GLN (-101.8, -93.0)  
A 608 PRO (-141.5, -74.1)  
B 3 PRO (-41.8, -84.7)  
B 77 GLU (-132.4, -3.4)  
B 79 ILE (-101.8, -93.0)  
B 130 PRO (-41.8, -84.7)  
B 134 ALA (-48.4, -100.9)  
B 135 THR (-15.5, -28.7)  
B 185 ALA (-48.4, -100.9)  
B 391 ILE (-101.8, -93.0)  
B 408 ILE (-101.8, -93.0)  
B 408 PRO (-141.5, -74.1)