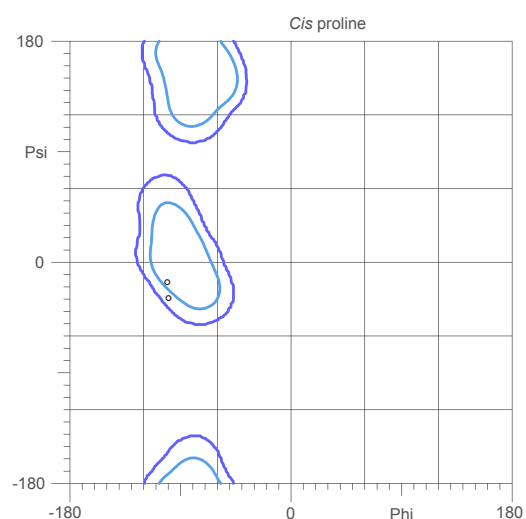
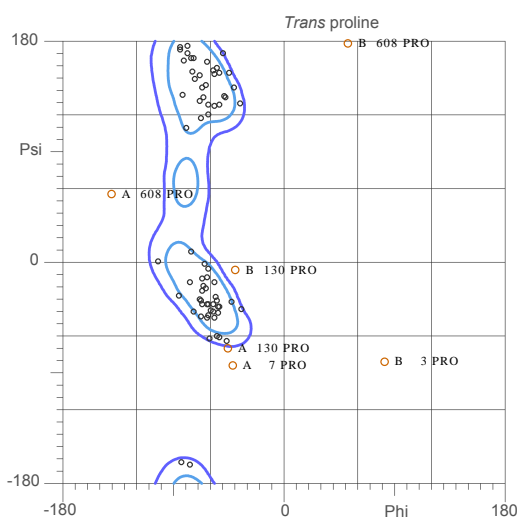
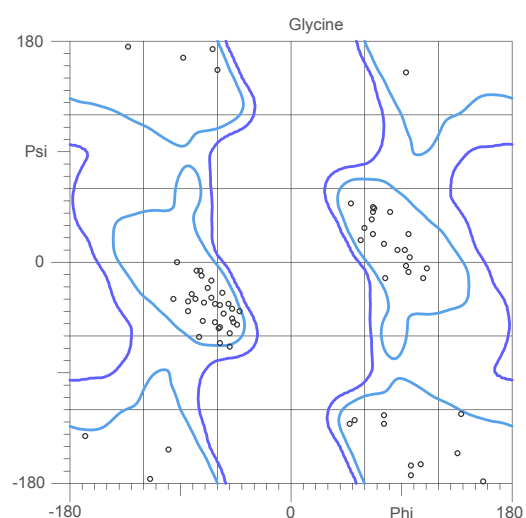
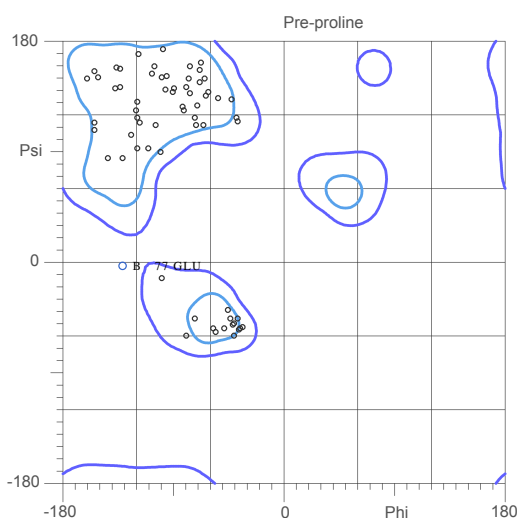
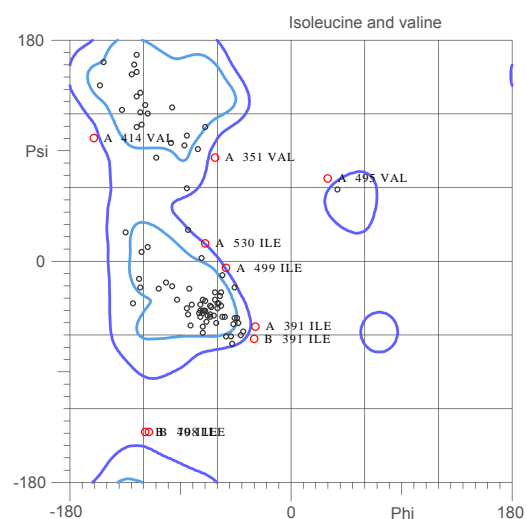
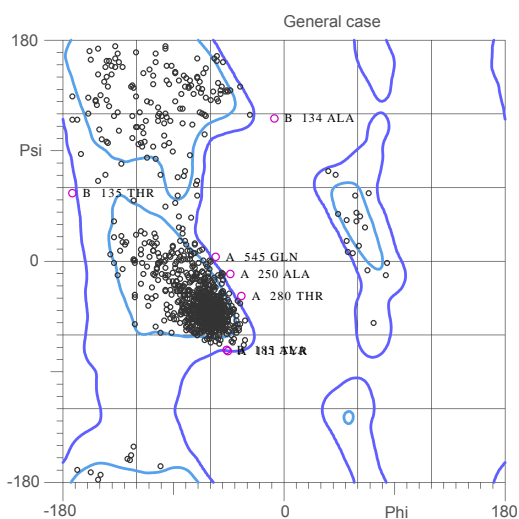


# MolProbity Ramachandran analysis

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 (-47.8, -75.5)
- A 130 PRO (-44.5, -78.9)
- A 200 ALA (-48.4, -100.9)
- A 280 THR (-35.5, -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 (-100.9, -98.7)
- 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, -54.1)
- B 3 PRO (-82.9, -91.7)
- 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 (-47.8, -75.5)
- B 391 ILE (-104.4, -84.0)
- B 408 ILE (-116.4, -109.8)
- B 408 PRO (-52.9, -179.3)