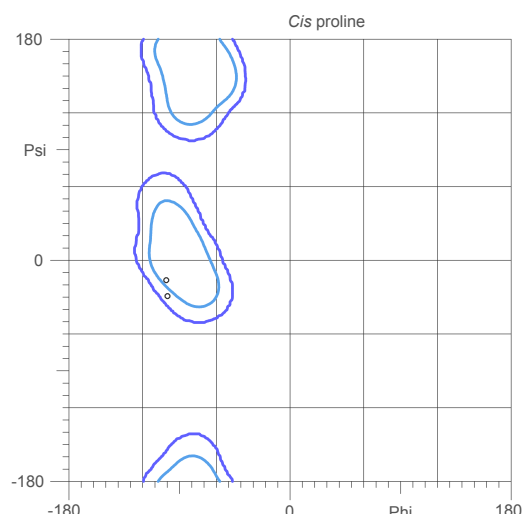
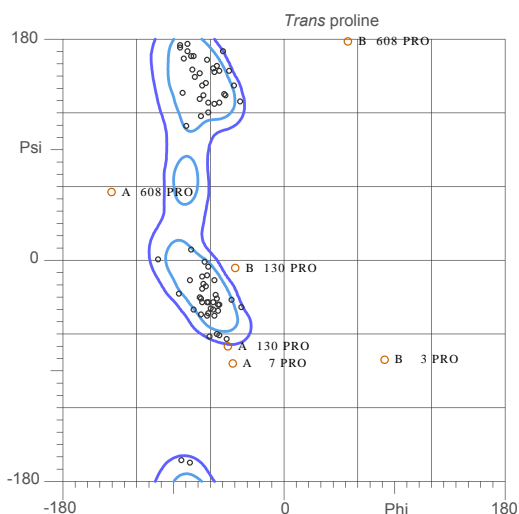
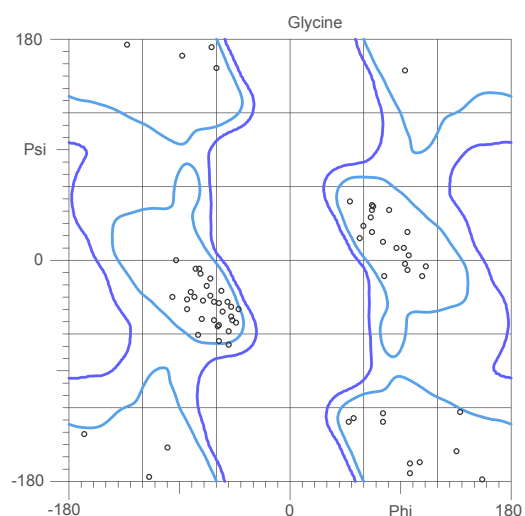
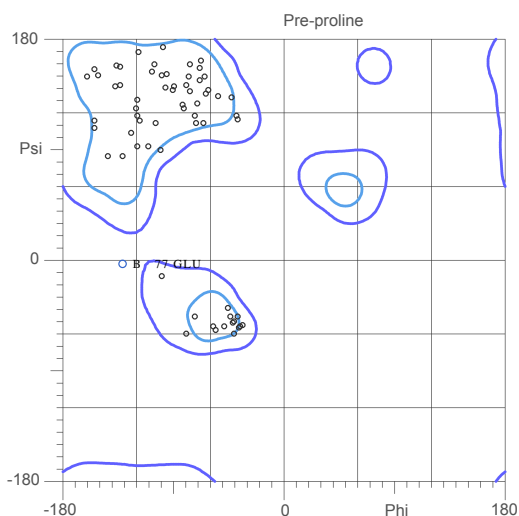
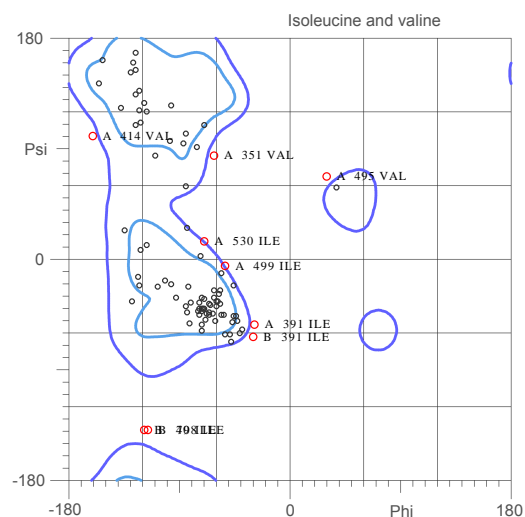
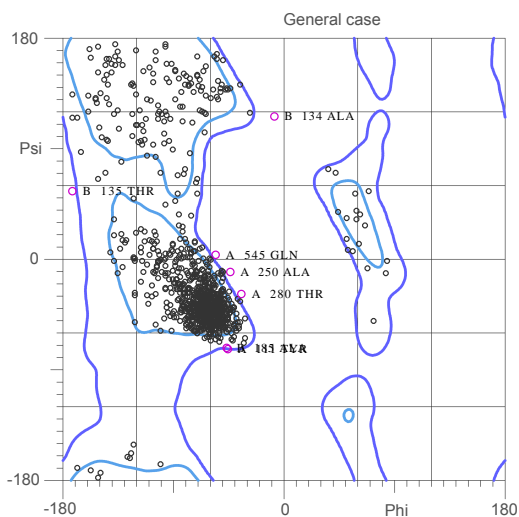


MolProbity Ramachandran analysis

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 (≥98%) 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, -10.9)
A 280 THR (-15.5, -28.7)
A 311 VAL (-82.7, 88.0)
A 391 ILE (-101.8, 93.0)
A 414 VAL (-101.8, 93.0)
A 495 VAL (-101.8, 93.0)
A 499 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 7 PRO (-82.8, -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, -10.9)
B 135 THR (-173.8, 56.4)
B 185 ALA (-47.5, -25.2)
B 391 ILE (-101.8, -84.0)
B 408 ILE (-118.4, -139.8)
B 408 PRO (-12.9, 179.3)