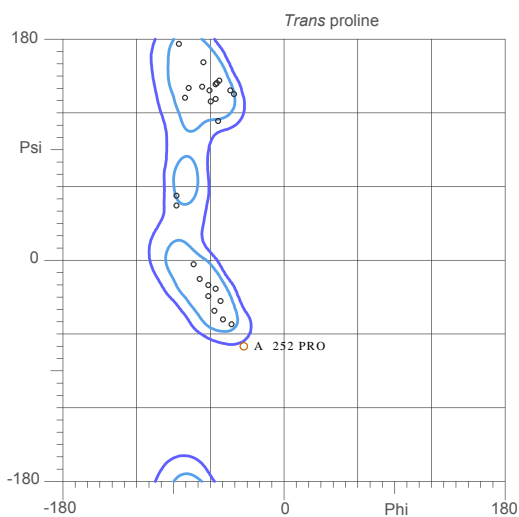
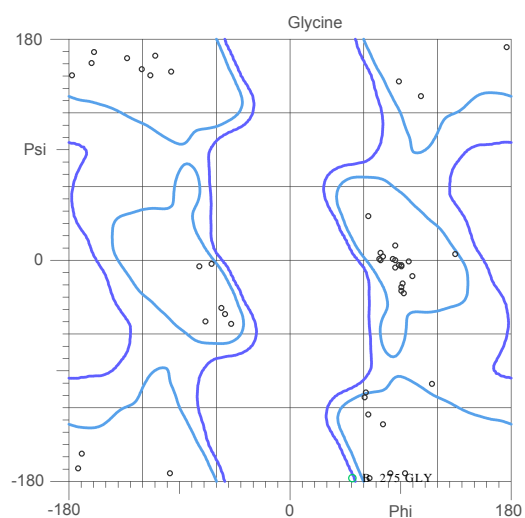
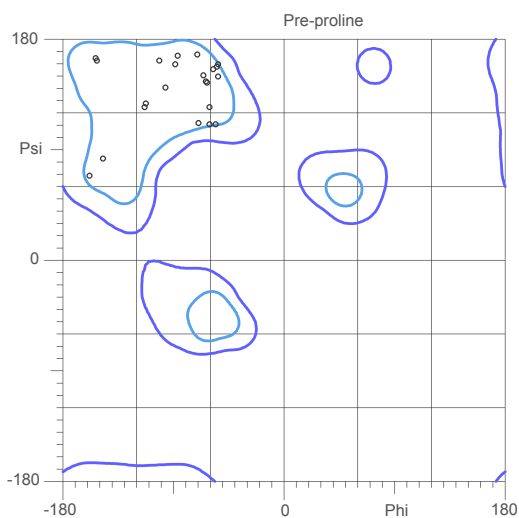
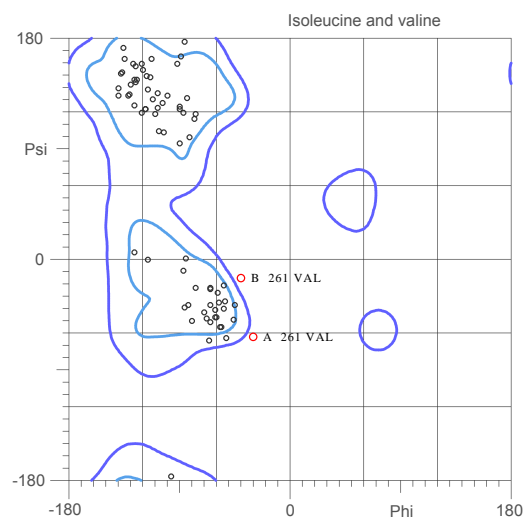
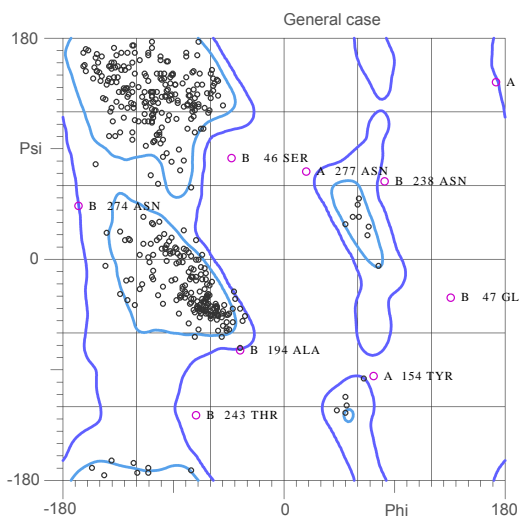


MolProbity Ramachandran analysis

7cbt.H.pdb, model 1



91.9% (544/592) of all residues were in favored (98%) regions.
97.8% (579/592) of all residues were in allowed (>99.8%) regions.

There were 13 outliers (phi, psi):

A 81 SER (173.6, 145.2)
A 154 TYR (73.6, -95.8)
A 252 PRO (-33.7, -70.6)
A 261 VAL (-30.1, -63.7)
A 277 ASN (18.3, 72.2)
B 46 SER (43.3, 83.3)
B 47 GLU (136.5, -31.3)
B 194 ALA (-36.0, -74.7)
B 238 ASN (82.3, 65.0)
B 243 THR (72.6, -127.4)
B 261 VAL (-40.3, -15.6)
B 274 ASN (-168.5, 44.1)
B 275 GLY (51.6, -178.2)