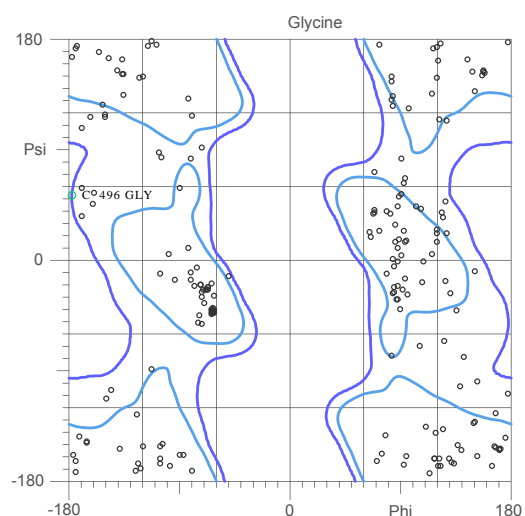
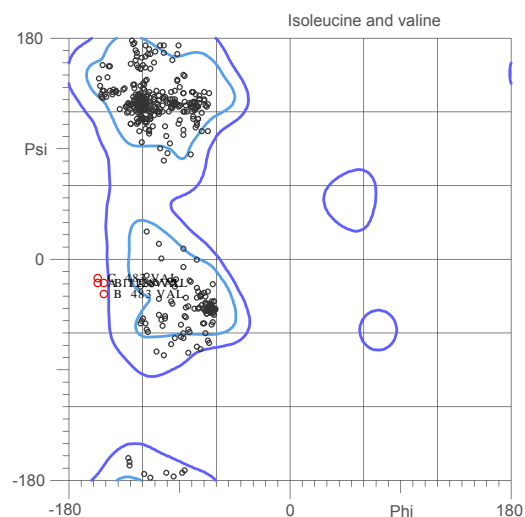


6zgg.H.pdb, model 1



87.1% (2763/3171) of all residues were in favored (98%) regions.
99.5% (3155/3171) of all residues were in allowed (>99.8%) regions

There were 16 outliers (phi, psi):

A 164 ASN (-73.1, -86.1)
A 293 LEU (-160.7, -80.8)
A 478 THR (58.1, 98.1)
A 603 ASN (-162.1, -41.2)
A 1128 VAL (-157.5, -15.5)
B 87 ASN (-84.9, -125.4)
B 329 PHE (111.0, -55.2)
B 478 THR (61.4, 108.1)
B 483 VAL (-152.8, -28.7)
B 1128 VAL (-152.8, -19.4)
B 1139 ASP (58.5, 90.4)
C 247 SER (-149.4, -125.5)
C 478 THR (57.9, 98.2)
C 483 VAL (-157.0, -15.2)
C 496 GLY (-178.1, 53.4)
C 1079 PRO (-64.0, -81.1)