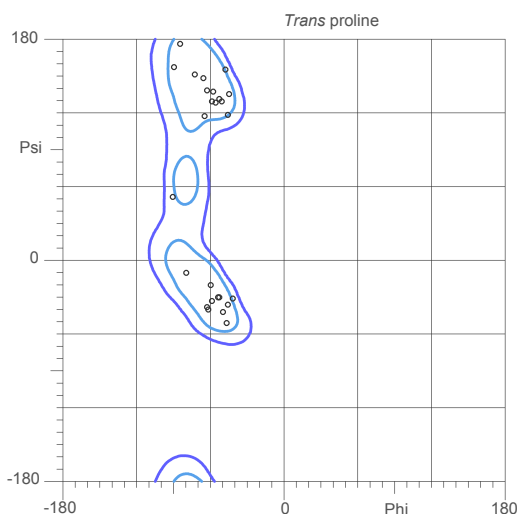
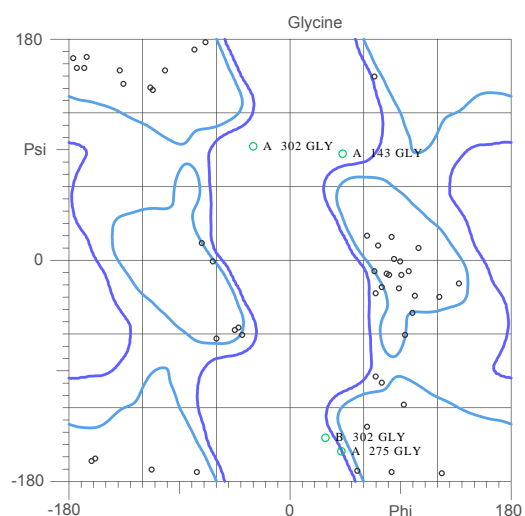
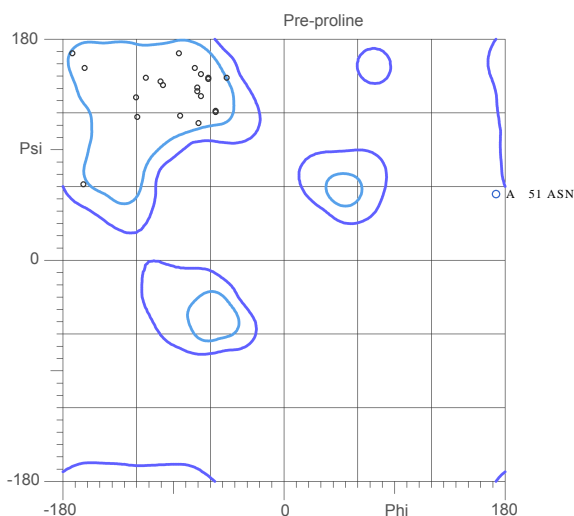
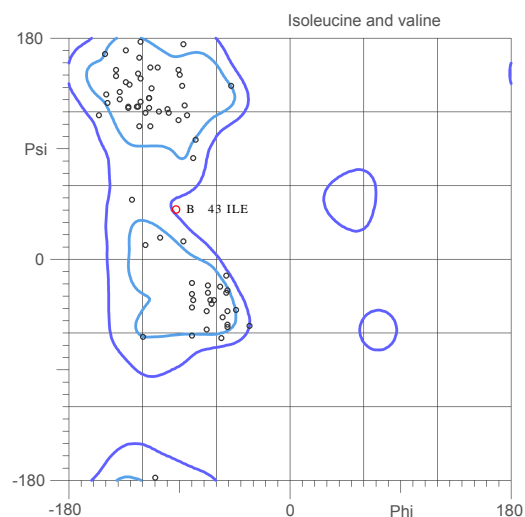
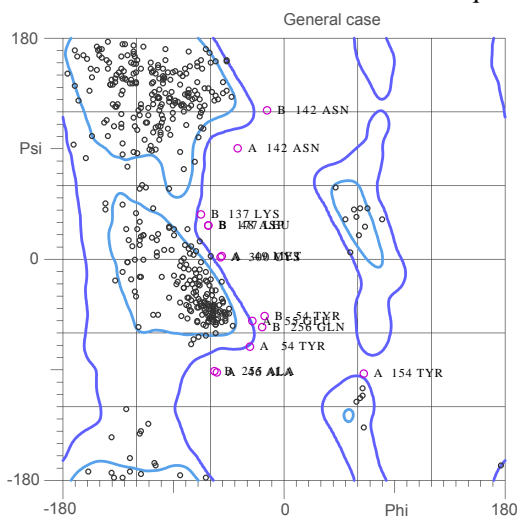


MolProbity Ramachandran analysis

2qc2.H.pdb, model 1



84.7% (511/600) of all residues were in favored (98%) regions.
96.7% (586/600) of all residues were in allowed (>95.8%) regions.

There were 20 outliers (Phi, Psi):
A 46 ALA (-55.2, -92.0)
A 49 MET (-51.4, 3.3)
A 51 ASN (-173.0, 54.9)
A 54 TYR (-28.6, -71.3)
A 55 GLY (-28.6, -10.0)
A 142 ASN (-38.2, 91.4)
A 143 GLY (-81.5, 87.3)
A 154 TYR (-85.1, -93.2)
A 270 GLY (-42.2, -150.8)
A 300 CYS (-52.2, 2.3)
A 302 GLY (-10.8, 93.1)
B 43 ILE (-61.4, -61.3)
B 48 ASP (-62.9, 28.4)
B 54 TYR (-16.2, -46.9)
B 137 LYS (-48.0, 37.7)
B 142 ASN (-15.0, 122.8)
B 177 LEU (-62.7, 28.9)
B 255 ALA (-57.7, -91.2)
B 256 GLN (-18.5, -59.3)
B 302 GLY (-29.0, -145.6)