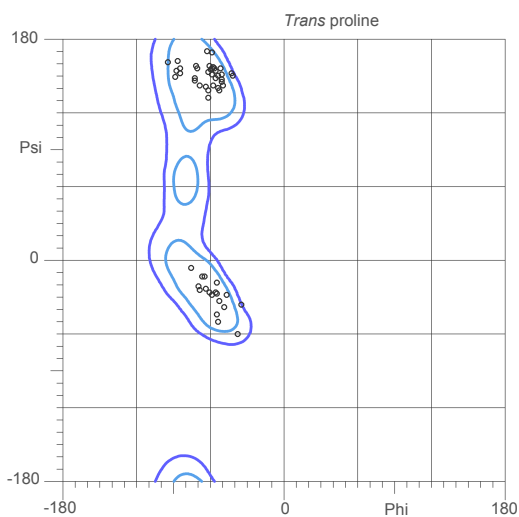
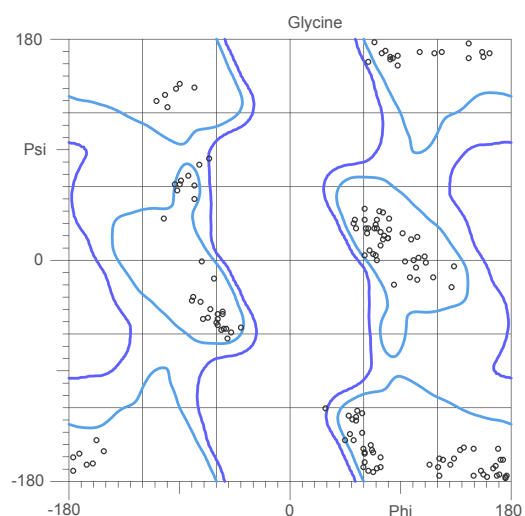
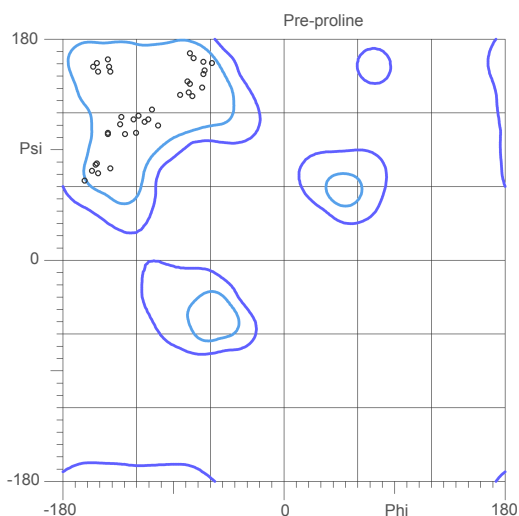
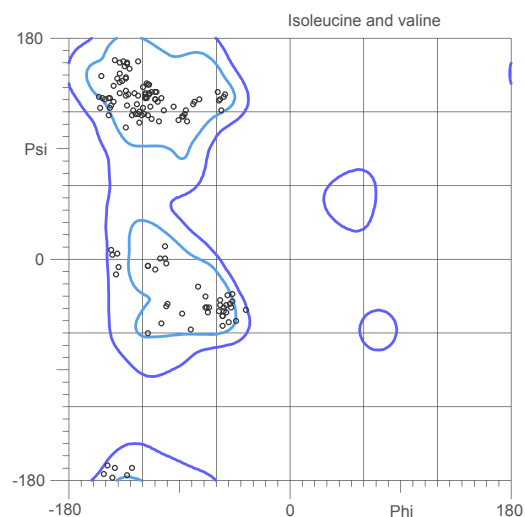
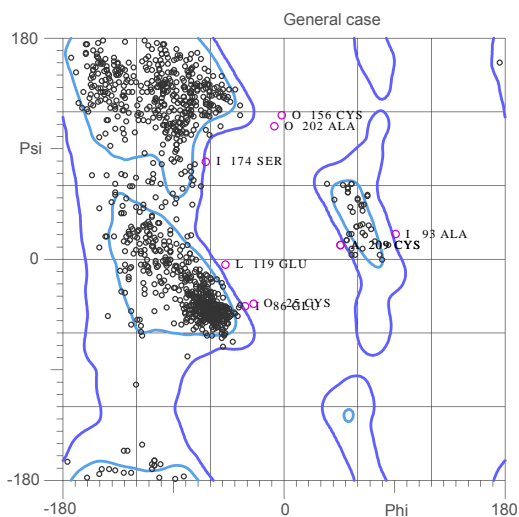


MolProbity Ramachandran analysis

4axm.H.pdb, model 1



91.3% (1271/1392) of all residues were in favored (98%) regions.
99.4% (1383/1392) of all residues were in allowed (>99.8%) regions.

There were 9 outliers (phi, psi):

A 209 CYS (46.1, 12.4)
A 209 CYS (46.1, 12.4)
I 86 GLU (-32.6, -38.1)
I 93 ALA (91.9, 21.2)
I 174 SER (-64.9, 80.2)
L 119 GLU (-48.8, -4.0)
O 25 CYS (-25.1, -36.9)
O 156 CYS (-2.5, 118.6)
O 202 ALA (-9.0, 109.1)