

# MolProbity Ramachandran analysis

2wct.H.pdb, model 1



94.2% (936/994) of all residues were in favored (98%) regions.  
98.7% (981/994) of all residues were in allowed (>99.8%) regions.

There were 13 outliers (phi, psi):

- A 491 GLY (-151.2, -27.5)
- A 636 SER (-55.5, -81.5)
- B 398 THR (97.3, -29.5)
- B 491 GLY (-151.5, -27.6)
- B 636 SER (-56.2, -81.3)
- C 408 THR (166.8, 150.7)
- C 409 ASN (121.8, -111.3)
- C 491 GLY (-147.2, -29.6)
- C 636 SER (-56.4, -80.6)
- D 406 PHE (3.8, 97.3)
- D 410 LYS (98.9, 126.3)
- D 491 GLY (-147.9, -29.2)
- D 636 SER (-56.7, -80.5)