

# MolProbity Ramachandran analysis

3k24.H.pdb, model 1



93.2% (408/438) of all residues were in favored (98%) regions.  
99.1% (434/438) of all residues were in allowed (>99.8%) regions.

There were 4 outliers (phi, psi):

- A 176 GLU (160.2, -133.2)
- A 177 SER (129.0, 122.9)
- A 179 ASN (117.0, 101.3)
- B 177 SER (90.4, 160.3)