

# MolProbity Ramachandran analysis

3tiu.H.pdb, model 1



96.4% (295/306) of all residues were in favored (98%) regions.

99.7% (305/306) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

47 GLU (123.2, -49.9)