

# MolProbity Ramachandran analysis

3vb4.H.pdb, model 1



95.9% (586/611) of all residues were in favored (98%) regions.

99.8% (610/611) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

B 154 TYR (54.6, -93.8)