

# MolProbity Ramachandran analysis

2fav.H.pdb, model 1



97.9% (501/512) of all residues were in favored (98%) regions.

99.6% (510/512) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

B 48 GLY (-51.1, -171.8)

B 49 GLY (-29.4, -144.8)