

# MolProbity Ramachandran analysis

6yb7.H.pdb, model 1



99.2% (370/373) of all residues were in favored (98%) regions.

99.7% (372/373) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

154 TYR (59.9, -88.7)