

MolProbity Ramachandran analysis

6y7m.H.pdb, model 1



96.7% (297/307) of all residues were in favored (98%) regions.

99.7% (306/307) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

5 LYS (-58.8, 96.6)