

MolProbity Ramachandran analysis

6m0j.H.pdb, model 1



97.1% (766/789) of all residues were in favored (98%) regions.

99.9% (788/789) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

A 102 GLN (-55.9, 93.9)