

MolProbity Ramachandran analysis

6lxt.H.pdb, model 1



97.9% (648/662) of all residues were in favored (98%) regions.

99.8% (661/662) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (ϕ , ψ):

F 1200 LEU (-57.8, 9.5)