

MolProbity Ramachandran analysis

6jd8.H.pdb, model 1



95.4% (310/325) of all residues were in favored (98%) regions.

99.4% (323/325) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (ϕ , ψ):

3 THR (17.9, 74.7)

76 ASN (77.0, 123.8)