

# MolProbity Ramachandran analysis

2bx3.H.pdb, model 1



96.2% (305/317) of all residues were in favored (98%) regions.  
98.7% (313/317) of all residues were in allowed (>99.8%) regions.

There were 4 outliers (phi, psi):

139 SER (-62.1, 44.2)  
142 ASN (-29.1, 94.3)  
154 TYR (44.4, -88.1)  
251 GLY (-13.1, -78.2)