

# MolProbity Ramachandran analysis

3scj.H.pdb, model 1



82.9% (234/282) of all residues were in favored (98%) regions.

16.7% (47/282) of all residues were in allowed (1.0%) regions.

There were 0 outliers (0% total).

Residue 147 (GLY) is in the 1.0% region.

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