

MolProbity Ramachandran analysis

3scj.H.pdb, model 1



82.9% (234/281) of all residues were in favored (98%) regions.

16.7% (47/281) of all residues were in allowed (1.0%) regions.

There were 0 outliers (0% total).

Residues in disallowed regions:

A 104 (GLN, 10.4)

A 105 (GLN, 10.5)

A 106 (GLN, 10.6)

A 107 (GLN, 10.7)

A 108 (GLN, 10.8)

A 109 (GLN, 10.9)

A 110 (GLN, 11.0)

A 111 (GLN, 11.1)

A 112 (GLN, 11.2)

A 113 (GLN, 11.3)

A 114 (GLN, 11.4)

A 115 (GLN, 11.5)

A 116 (GLN, 11.6)

A 117 (GLN, 11.7)

A 118 (GLN, 11.8)

A 119 (GLN, 11.9)

A 120 (GLN, 12.0)

A 121 (GLN, 12.1)

A 122 (GLN, 12.2)

A 123 (GLN, 12.3)

A 124 (GLN, 12.4)

A 125 (GLN, 12.5)

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A 127 (GLN, 12.7)

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A 134 (GLN, 13.4)

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A 140 (GLN, 14.0)

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A 142 (GLN, 14.2)

A 143 (GLN, 14.3)

A 144 (GLN, 14.4)

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A 146 (GLN, 14.6)

A 147 (GLN, 14.7)

A 148 (GLN, 14.8)

A 149 (GLN, 14.9)

A 150 (GLN, 15.0)

A 151 (GLN, 15.1)

A 152 (GLN, 15.2)

A 153 (GLN, 15.3)

A 154 (GLN, 15.4)

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A 341 (GLN, 34.1)