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MolProbability results



Summary MolProbability

MolProbability Basic Statistics  **Open In Coot**  


Basic Geometry: Bond Length Restraints

Deviations  



Basic Geometry: Bond Angle Restraints

Deviations  



Basic Geometry: Dihedral Angle Restraints

Deviations  **Outliers**  

Basic Geometry: Chirality Restraints

Deviations  

Basic Geometry: Planarity Restraints



Deviations  

Protein

Select plot: Ramachandran plot **View plot**  

No Ramachandran outliers detected

No Rotamer outliers detected

No C-beta position outliers detected No sidechain flips required **Cis and Twisted peptides:**  

Clashes

All atom-contact analysis  