BindForge Docking Analysis Report

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Protein Information

Parameter	Value
Name	Target Protein
PDB ID	Unknown
Number of Cysteines	1

Identified Cysteines

Chain	Residue Number	Residue Name
A	3	CYS

Molecule Information

Parameter	Value
Name	Aspirin
SMILES	c1(ccccc1)OC(=O)C(=O)O■Aspirin

Docking Results

Summary

Metric	Value
Best Binding Affinity	-7 kcal/mol
Covalent Binding Potential	Low

Warhead Distance	3.5 Å

Docking Poses

Pose	Affinity (kcal/mol)	RMSD LB	RMSD UB
1	-7	0	0

Analysis and Recommendations

No specific recommendations available.