BindForge Docking Analysis Report

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Protein Information

Parameter	Value
Name	Target Protein
PDB ID	Unknown
Number of Cysteines	0

Identified Cysteines

No cysteine residues identified

Molecule Information

Parameter	Value
Name	molecule_5268
Molecular_weight	~200-300 Da
Molecular_formula	
Has_warhead	False
Source	User uploaded molecule
Logp	-0.67
Druglikeness	0.55
Synthetic_accessibility	1.25
SMILES	CC(=O)OC1=CC=CC=C1C(=O)O

Docking Results

Summary

Metric	Value
Best Binding Affinity	-10 kcal/mol
Covalent Binding Potential	N/A
Warhead Distance	None Å

Docking Poses

Pose	Affinity (kcal/mol)	RMSD LB	RMSD UB
1	-10	0	0
2	-10	0	0

Analysis and Recommendations

No specific recommendations available.

Generated by BindForge - Al-Enhanced Covalent Docking Prediction Tool