BindForge Docking Analysis Report

Generated on 2025-04-27 14:30:24

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_1785	
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

Analysis and Recommendations

No specific recommendations available.

Generated by BindForge - Al-Enhanced Covalent Docking Prediction Tool