### **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
Α	3	CYS

### **Molecule Information**

Parameter	Value
Name	molecule_2258
SMILES	C(C(=O)O)N

## **Docking Results**

### **Summary**

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

Warhead Distance	3.5 Å
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### **Docking Poses**

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

# **Analysis and Recommendations**

No specific recommendations available.

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