# 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 | molecule_6153 |
| 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 |

## Docking Poses

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

# Analysis and Recommendations

No specific recommendations available.