

[SwissDock](#)



[SwissParam](#)



[SwissSidechain](#)



[SwissBioisostere](#)



[SwissTargetPrediction](#)



[SwissADME](#)



[SwissSimilarity](#)



[Swiss Institute of Bioinformatics](#)

- [Contact](#)
- [Citing](#)
- [Command-line](#)
- [Tutorials](#)
- [FAQ](#)
- [About](#)
- [Home](#)

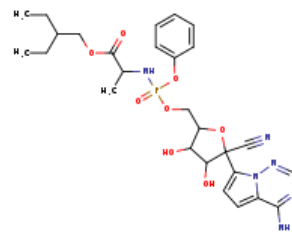
Run parameters

| | |
|------------------|----------------------------------|
| Library screened | ZINC_lead |
| Screening method | ECFP4 |
| Date | September 30, 2024, 11:24 am UTC |

If you publish these results, please, cite the following paper:

[Zoete, V., Daina, A., Bovigny, C., & Michielin, O. SwissSimilarity: A Web Tool for Low to Ultra High Throughput Ligand-Based Virtual Screening., *J. Chem. Inf. Model.*, **2016**, *56*\(8\), 1399-1404.](#)

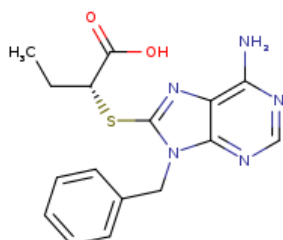
Query Molecule



Results

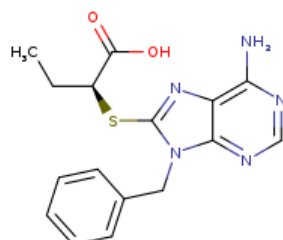
[ZINC000085397313](#)

Score : 0.259



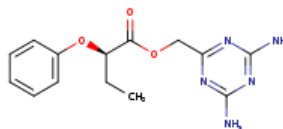
[ZINC000085397312](#)

Score : 0.259



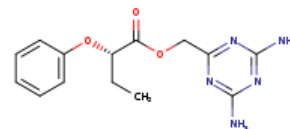
[ZINC000012984088](#)




Score : 0.255




[ZINC000012984090](#)

Score : 0.255



Retrieve data:   



 SwissSimilarity is operated by the [Molecular Modelling Group](#) of the [SIB Swiss Institute of Bioinformatics](#) and the [University of Lausanne](#)
| [Terms of use](#) | [SIB privacy policy](#)