



**SwissBioisostere** 



**SwissTargetPrediction** 





**SwissSimilarity** 



## Swiss Institute of Bioinformatics

- Contact
- Citing
- Command-line
- **Tutorials**
- <u>FAQ</u>
- <u>About</u>
- <u>Home</u>

Run parameters

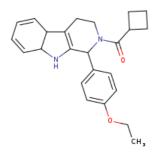
ZINC\_lead Library screened ECFP4 Screening method

Date September 30, 2024, 11:40 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**

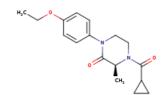


## Results

ZINC000053718499 Score: 0.435

ZINC000037521887 Score: 0.397

ZINC000037521888 Score: 0.397



Retrieve data:

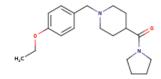


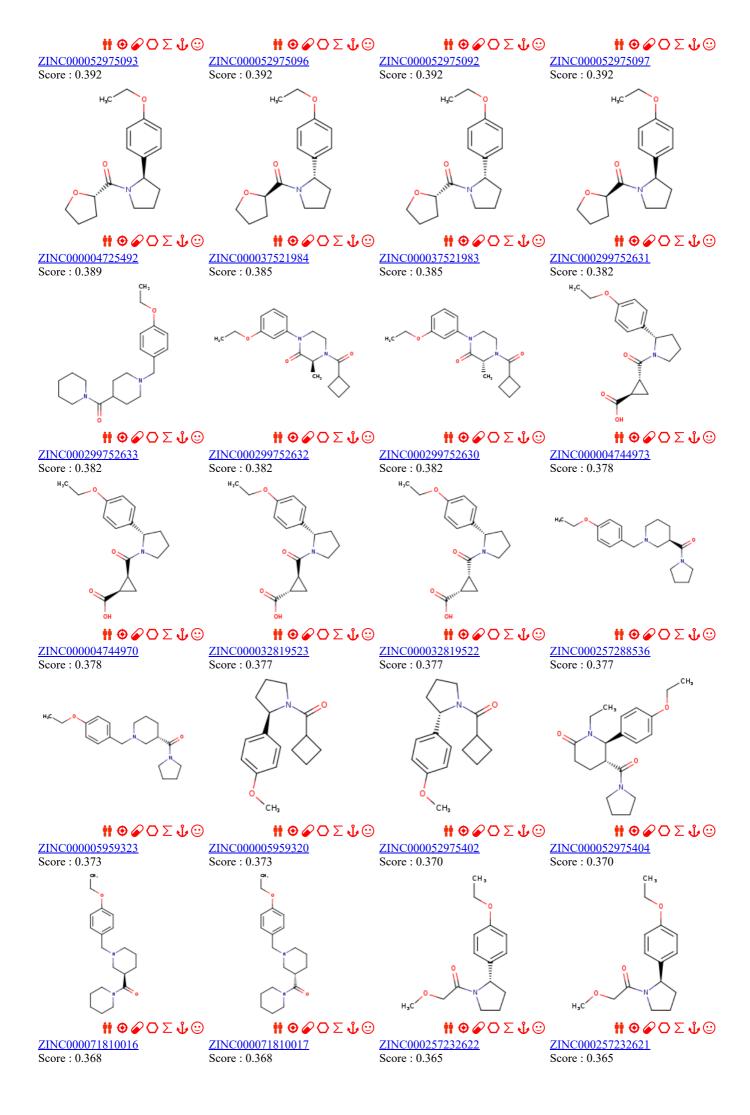
# • • D \(\Sigma\)

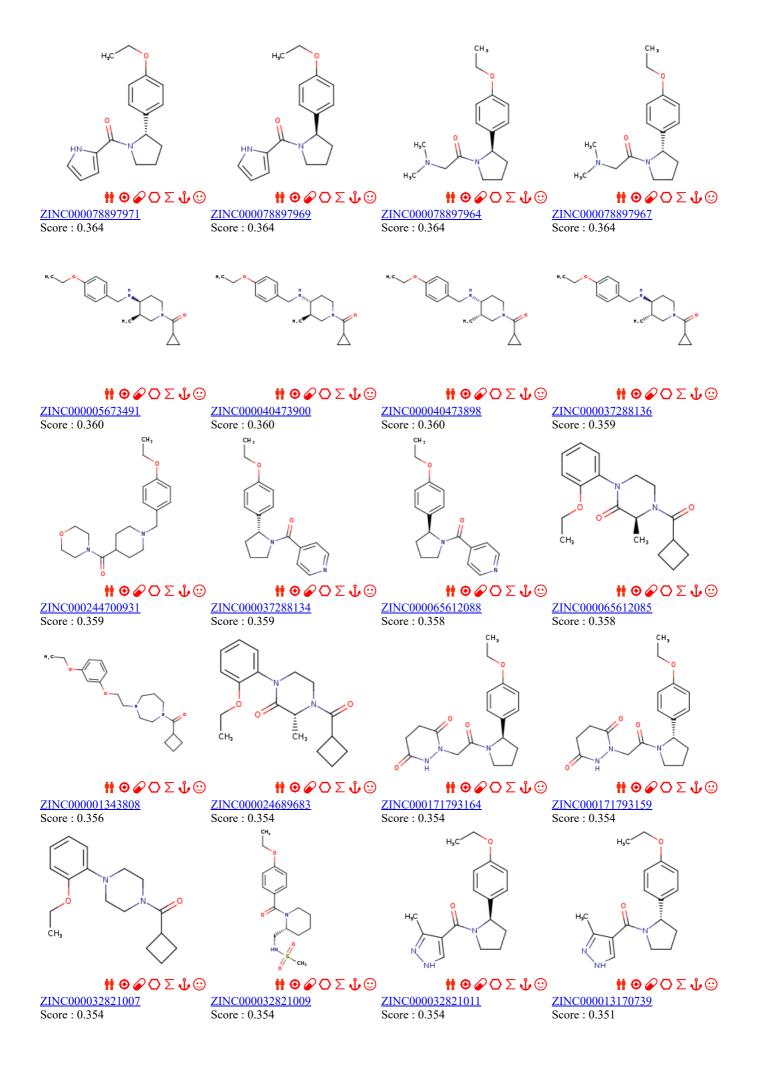


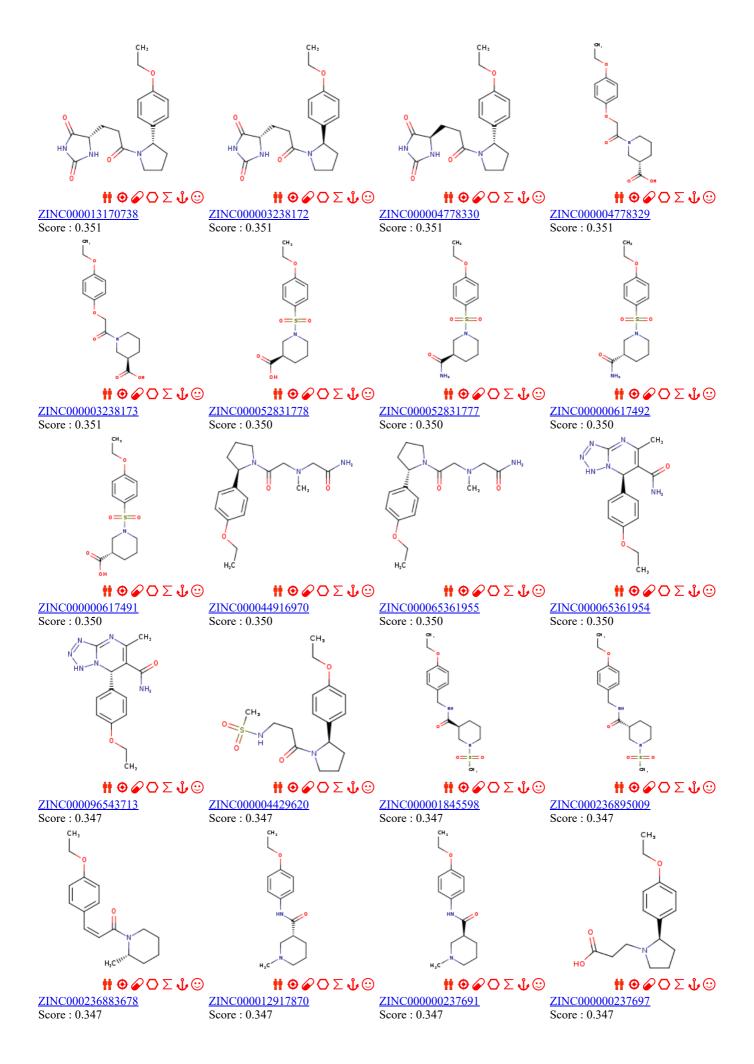


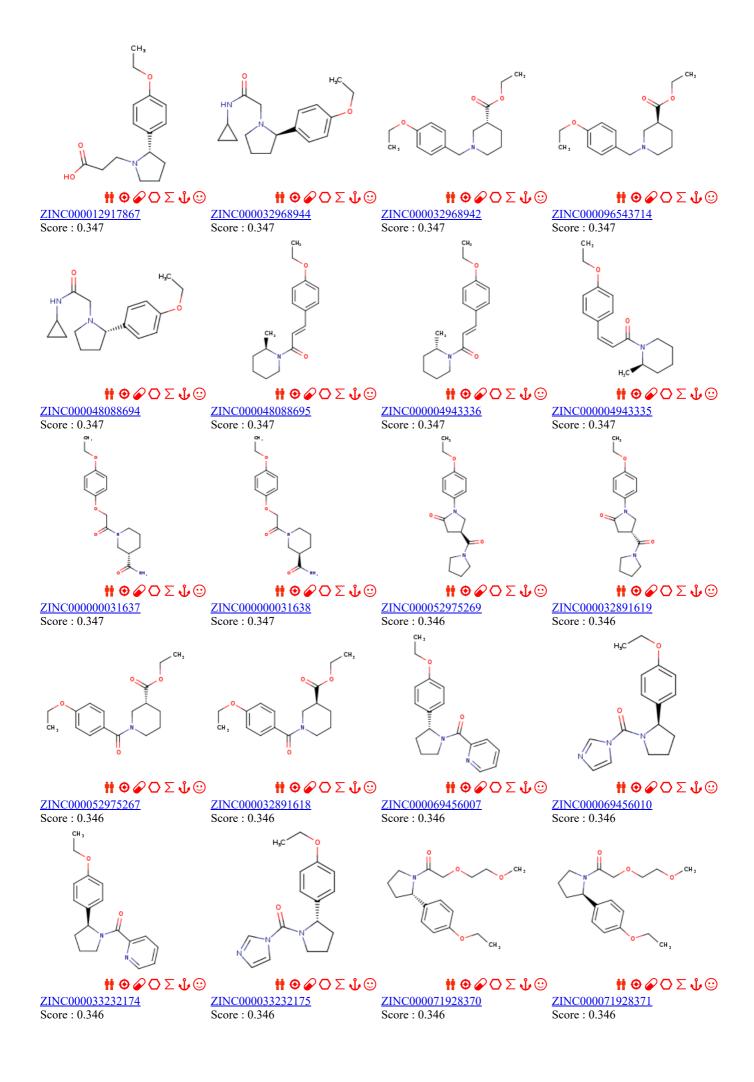
ZINC000005673490 Score: 0.394

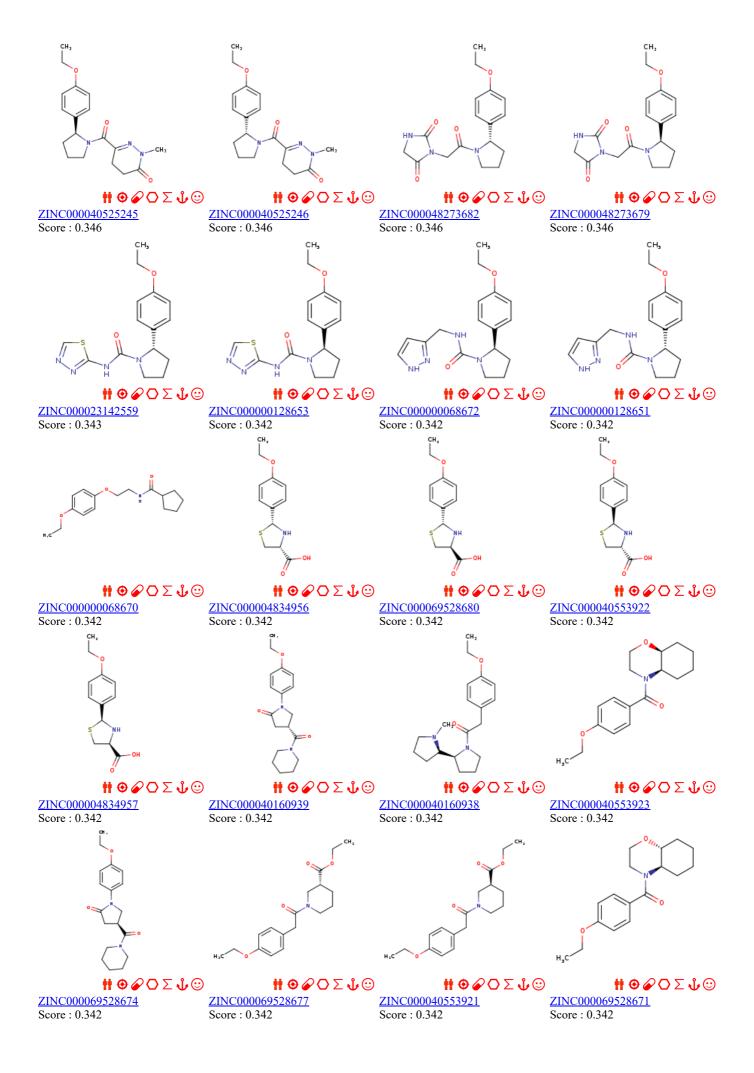


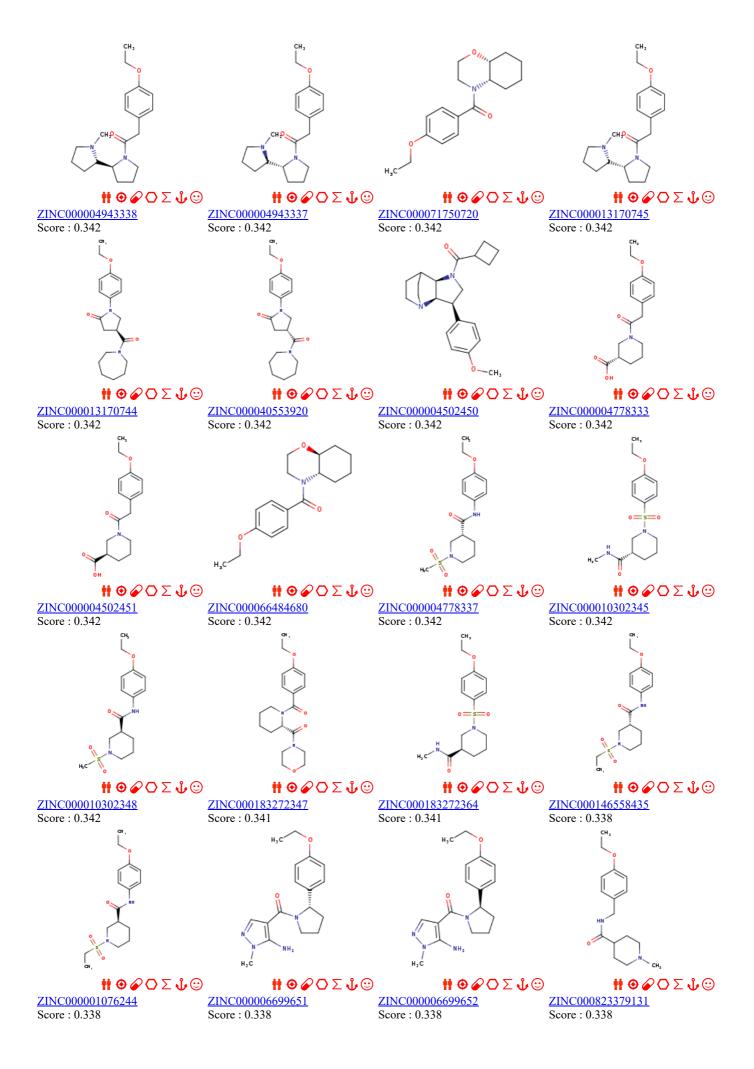


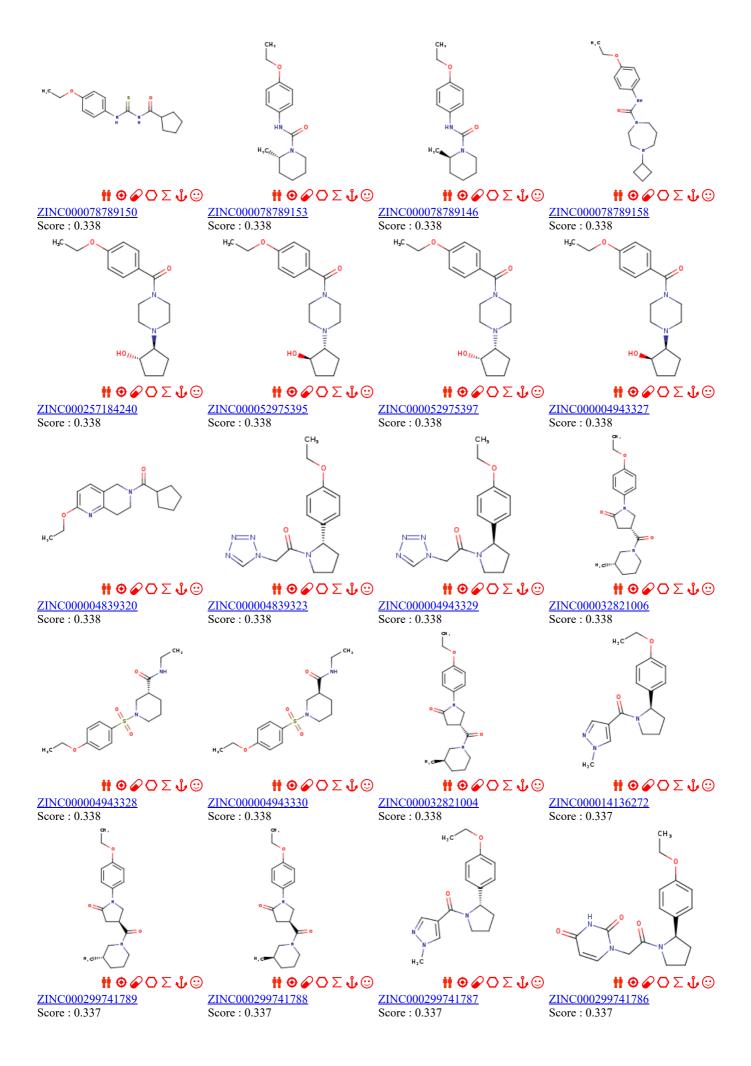


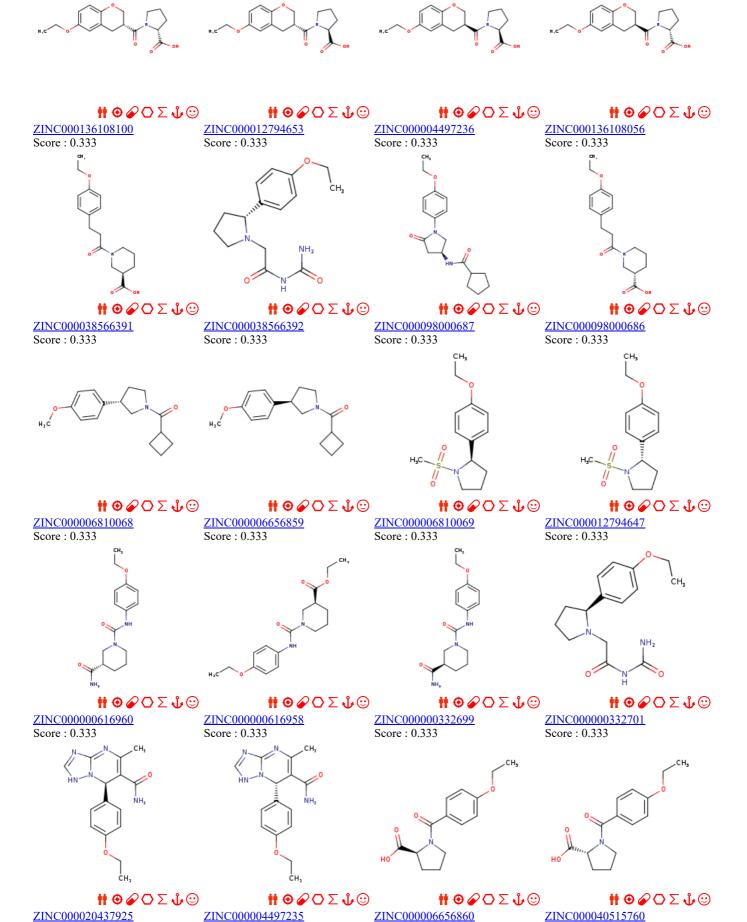










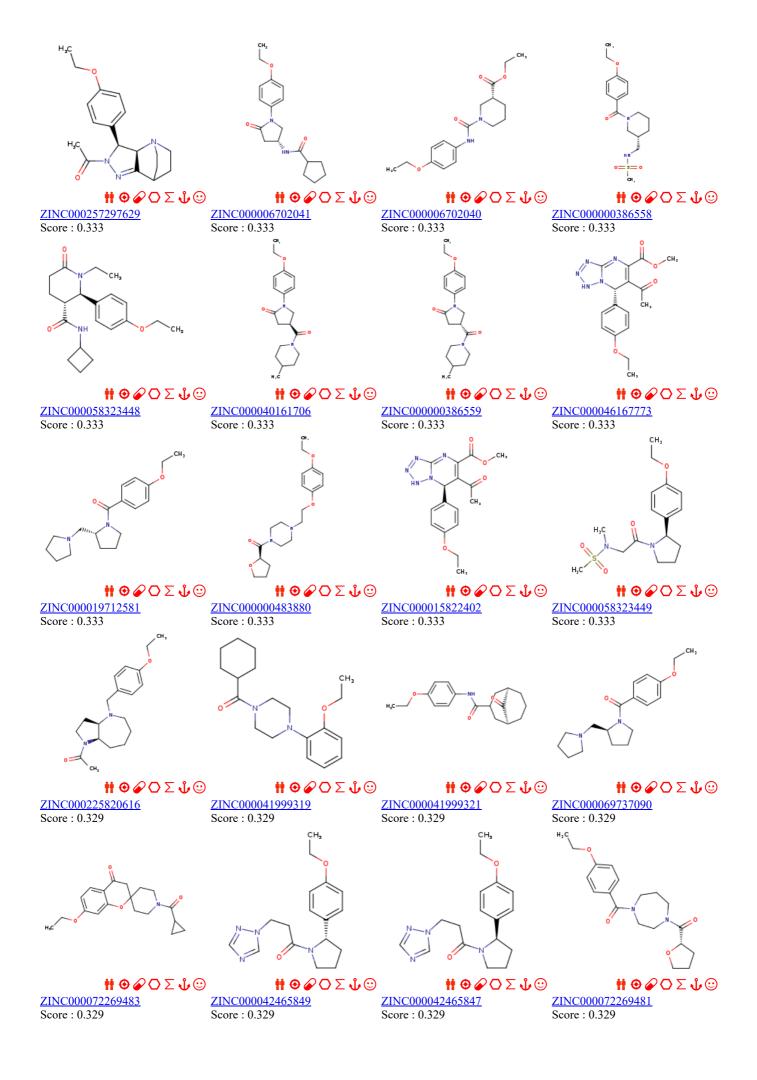


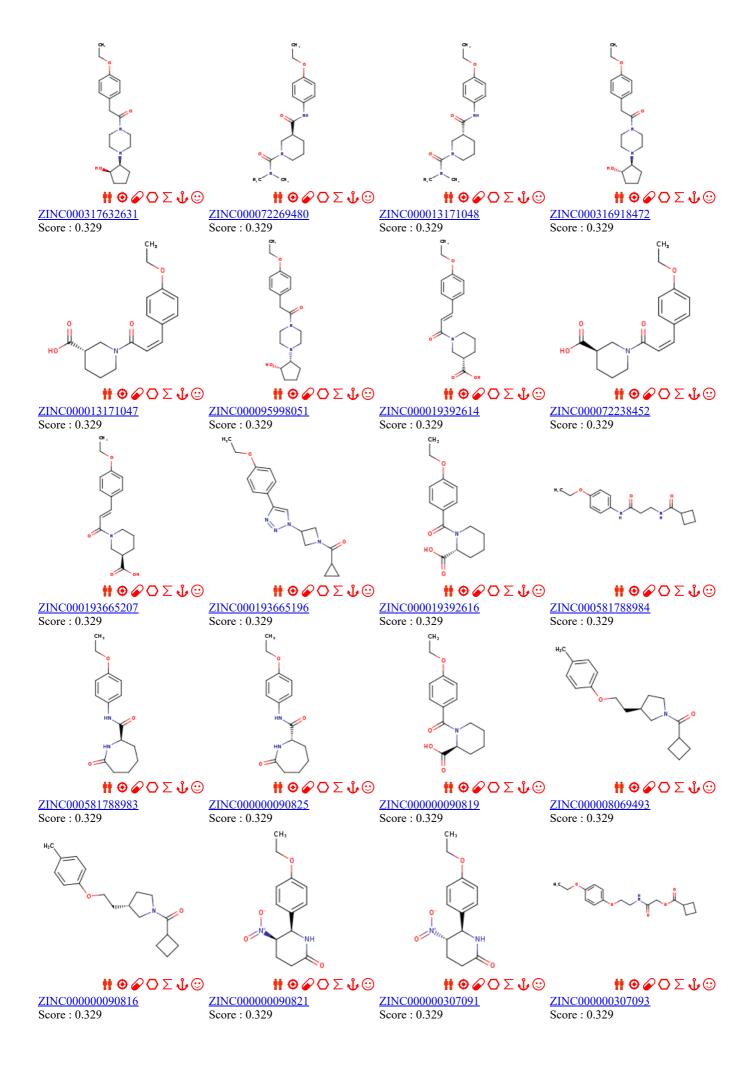
Score: 0.333

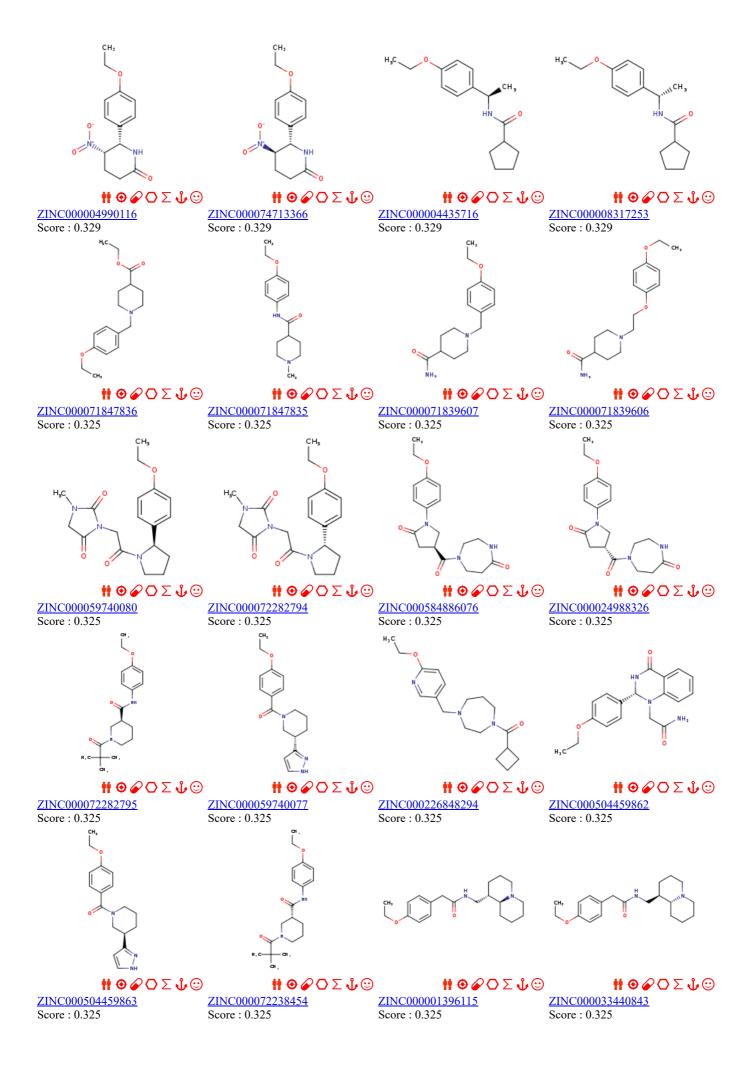
Score: 0.333

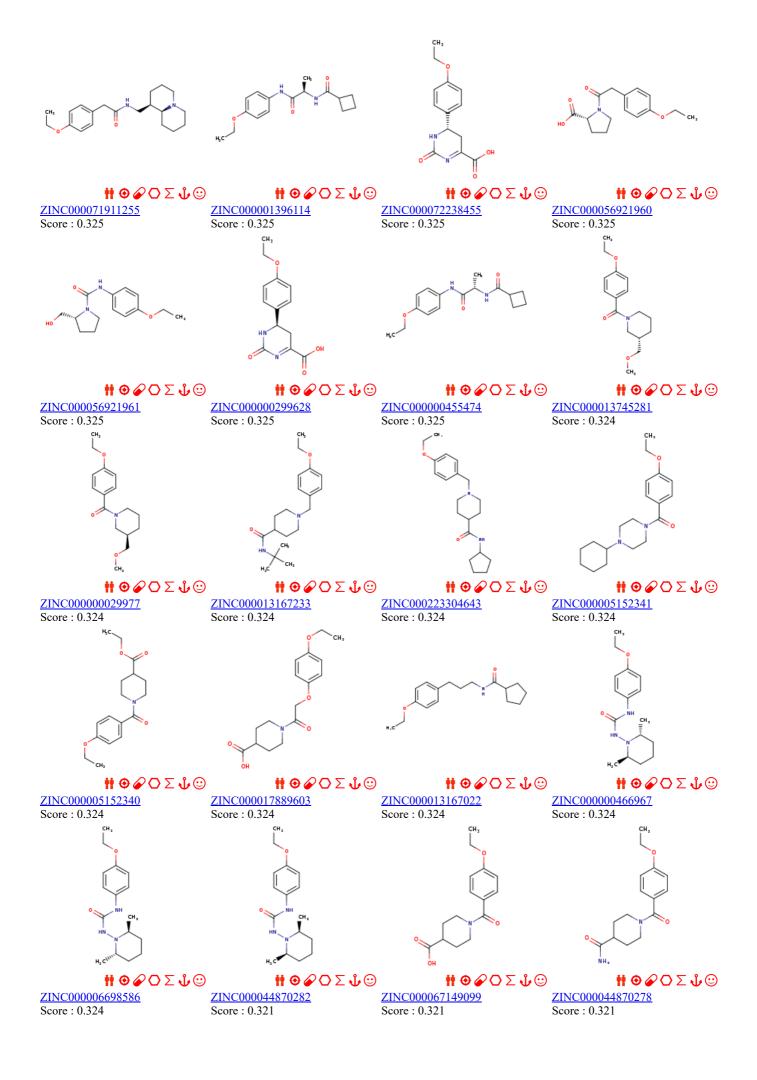
Score: 0.333

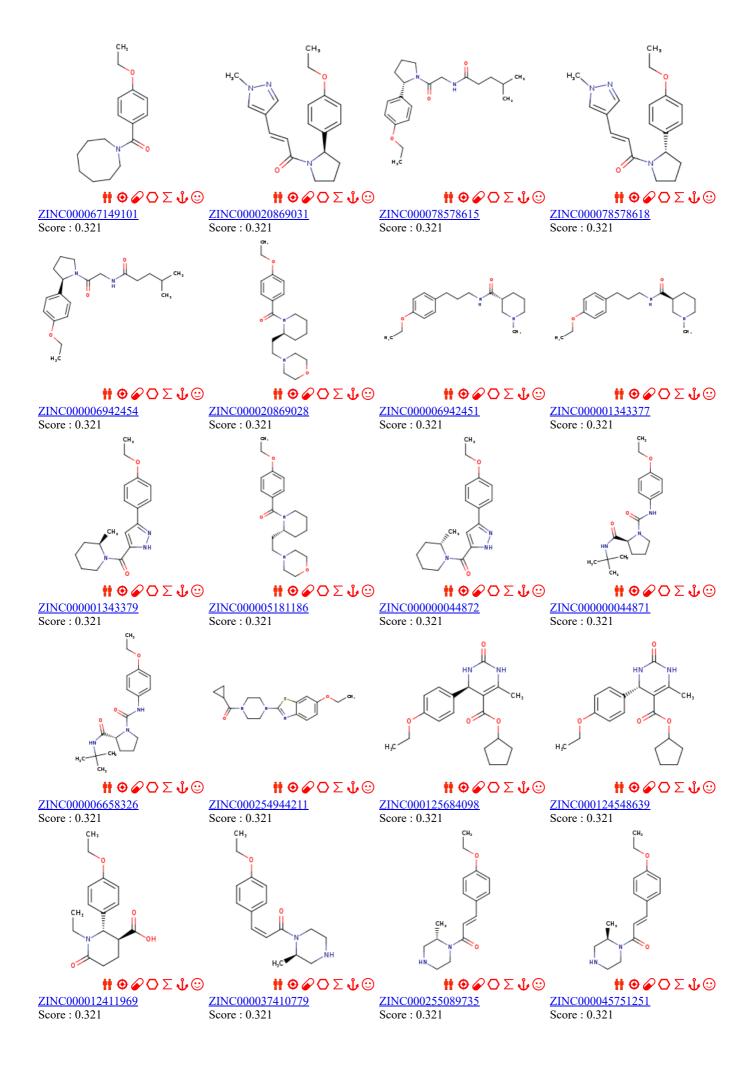
Score: 0.333

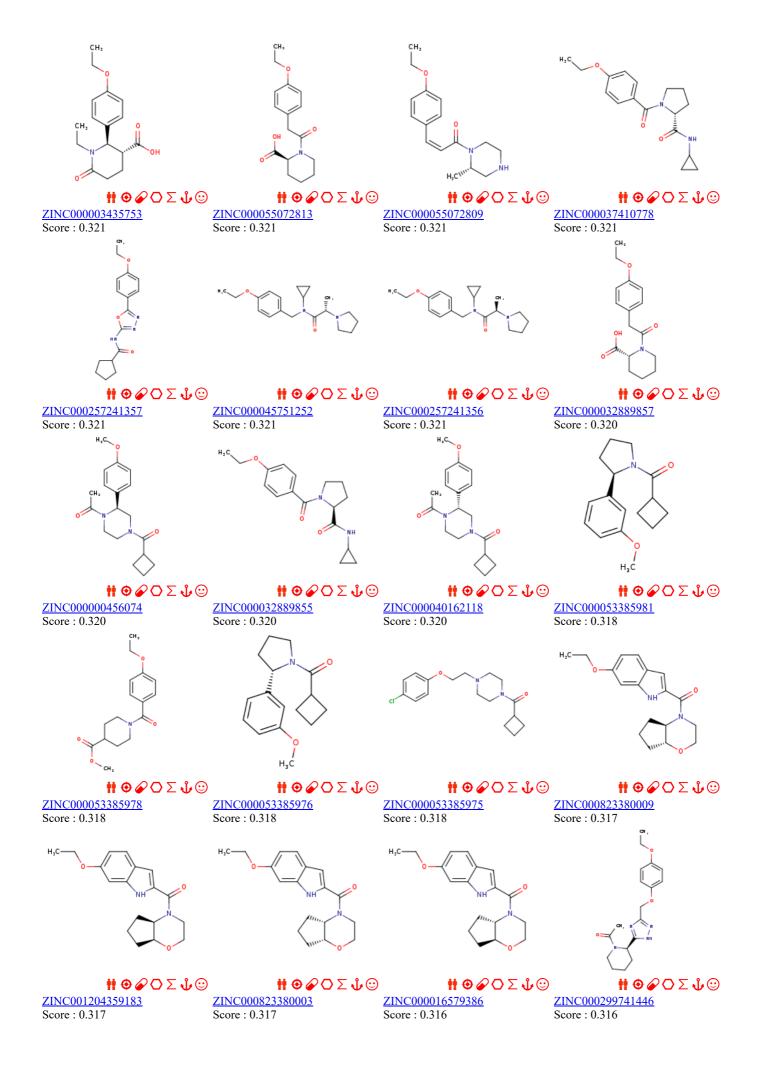


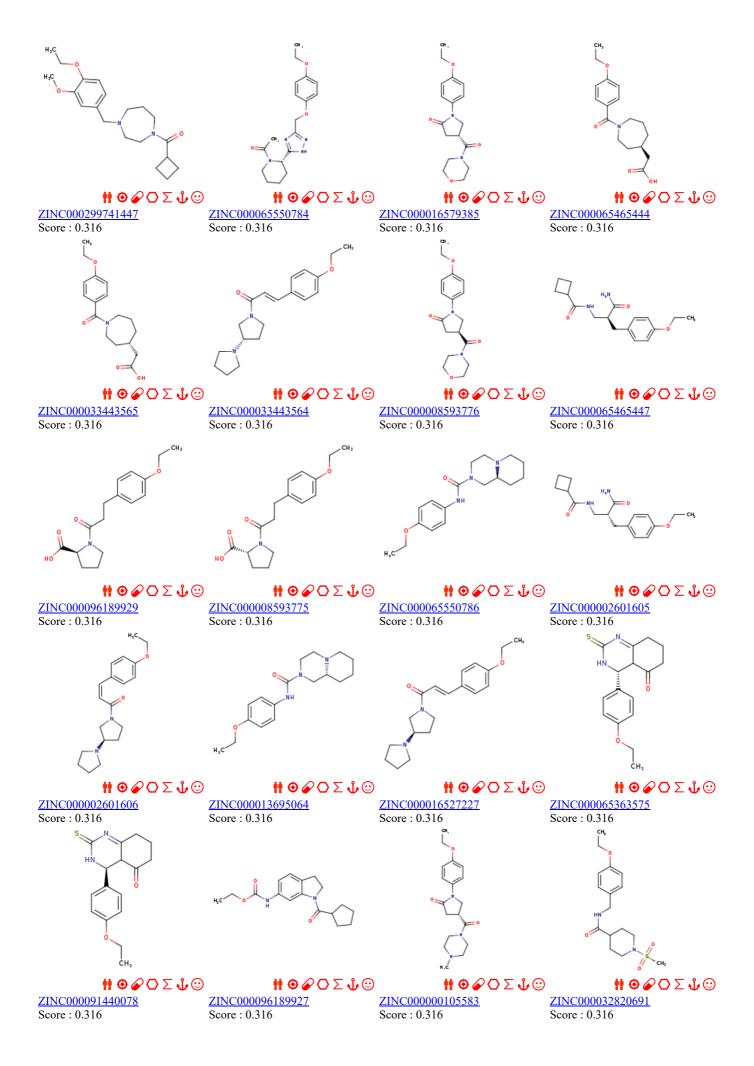


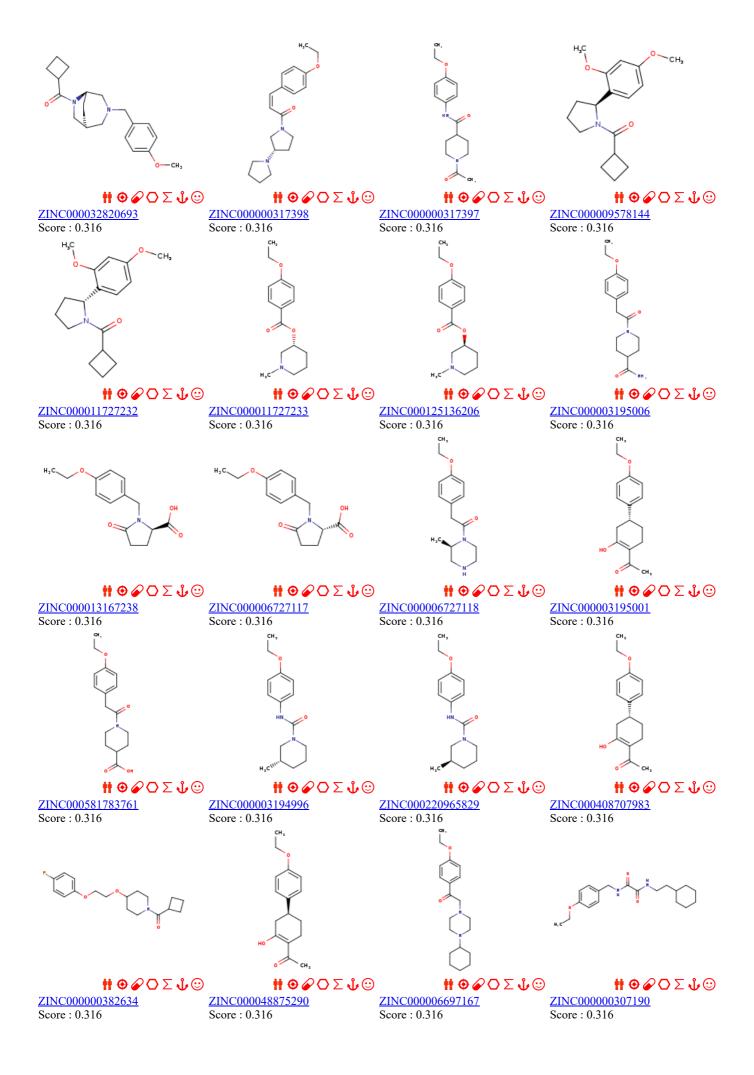


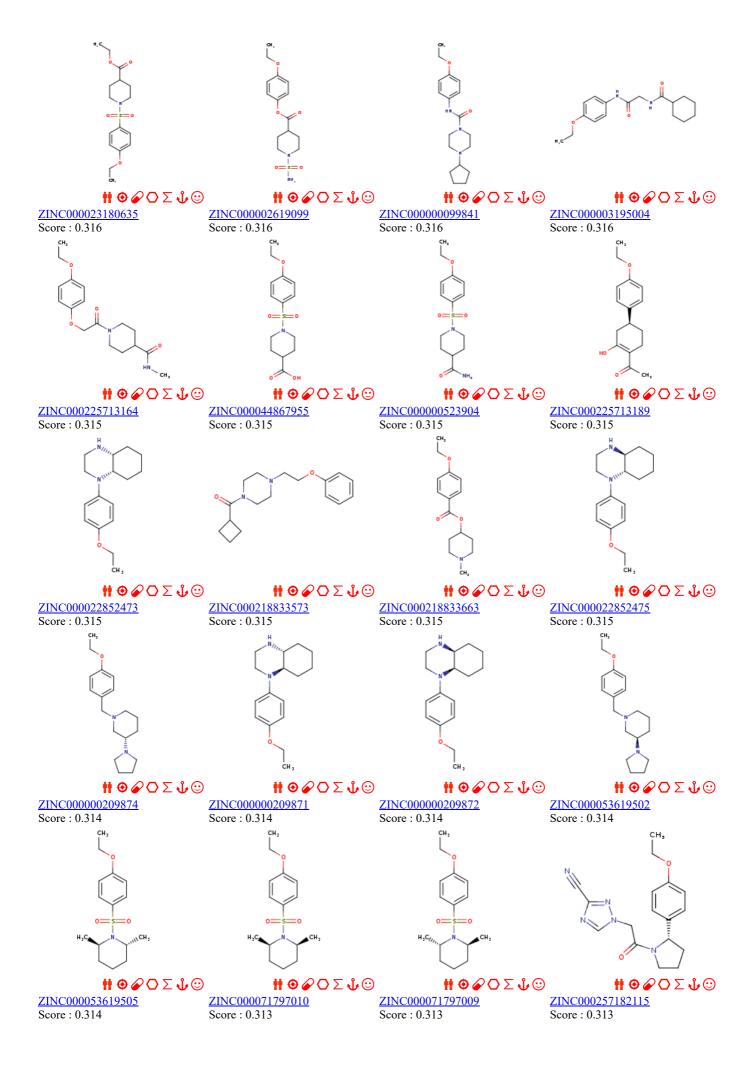


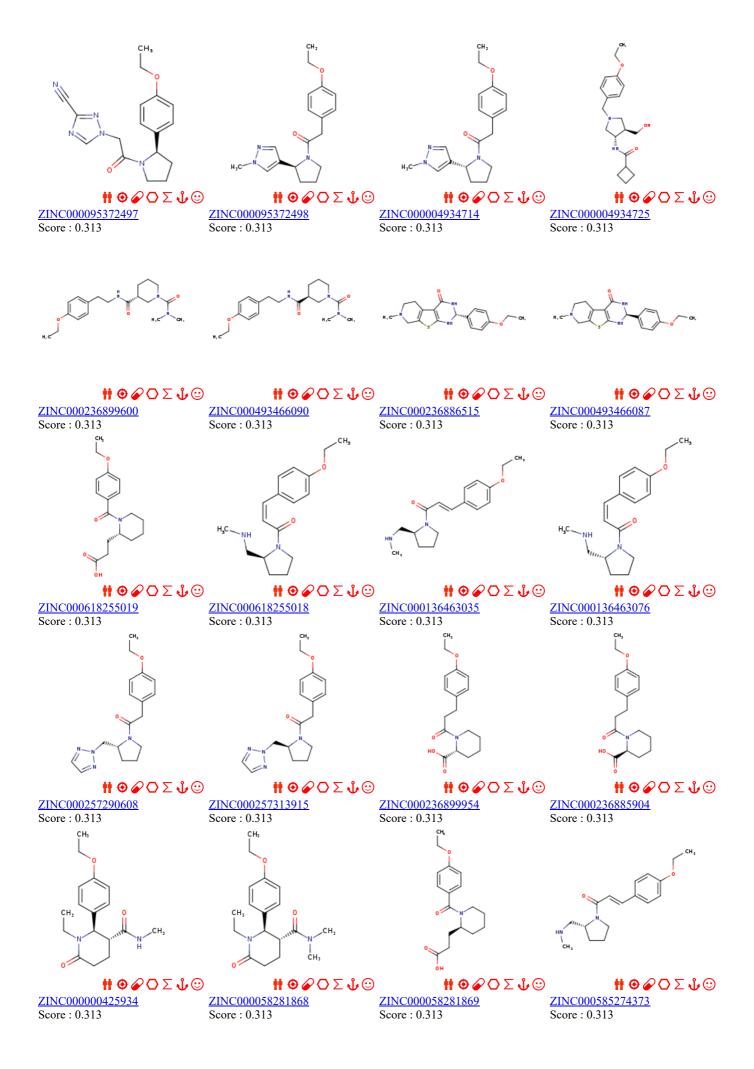


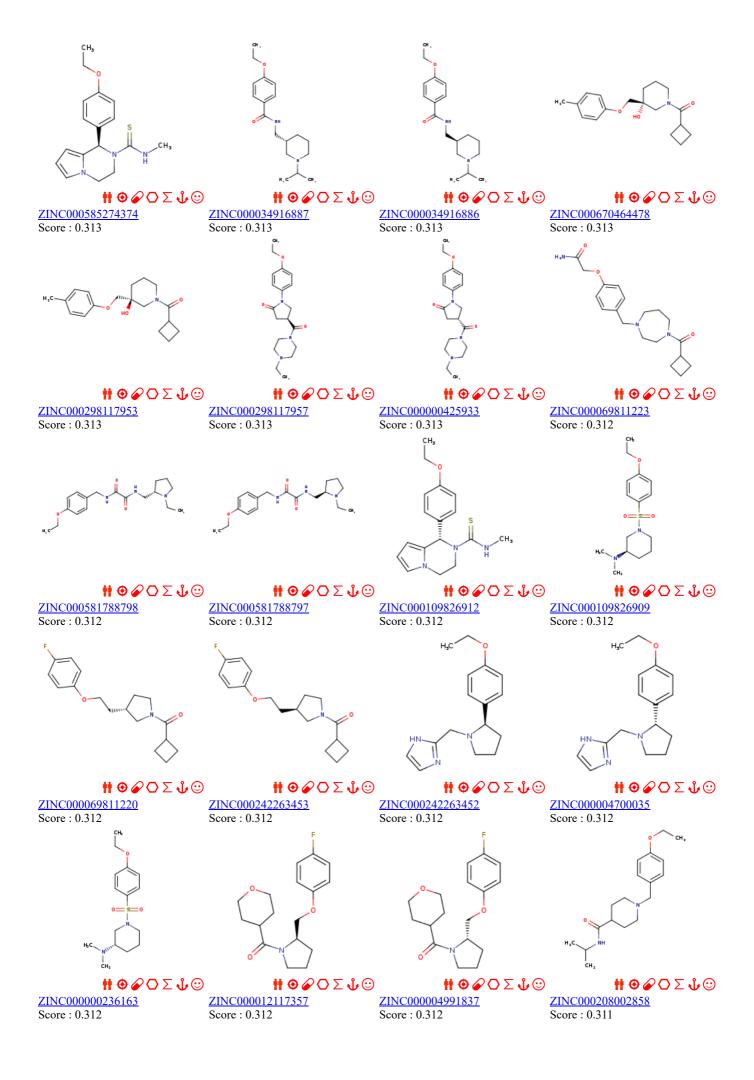


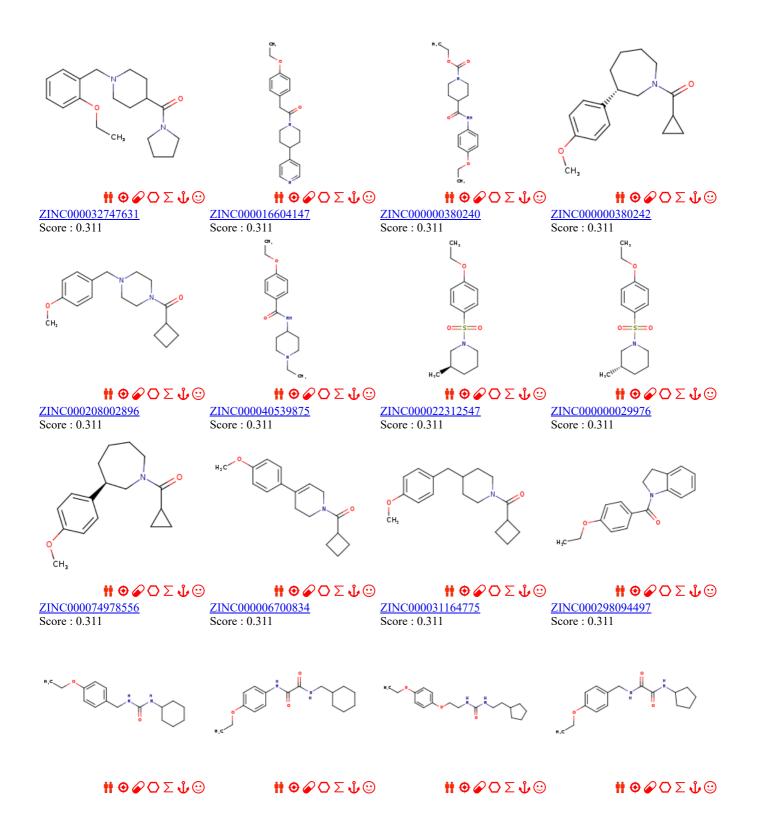














SwissSimilarity is operated by the Molecular Modelling Group of the SIB Swiss Institute of Bioinformatics and the University of Lausanne | Terms of use | SIB privacy policy