

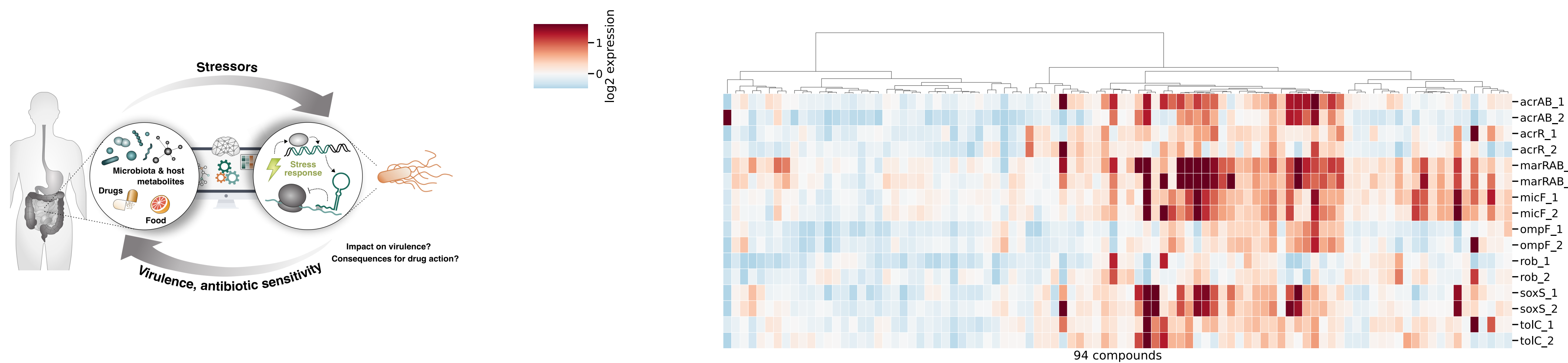
# We use molecular features and metric learning to predict gene expression

## Cheminformatics deciphers stress response and virulence pathways in infection

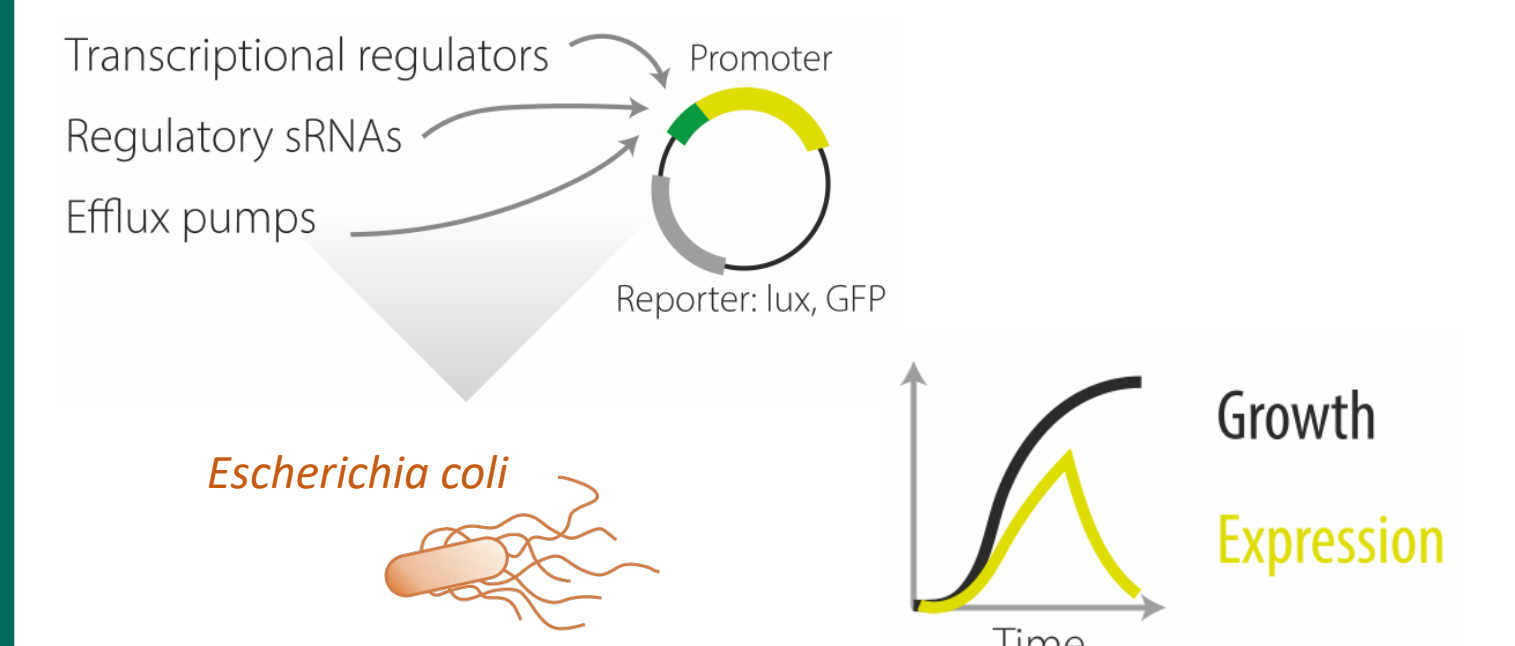
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### Small compounds and their effect on microbes

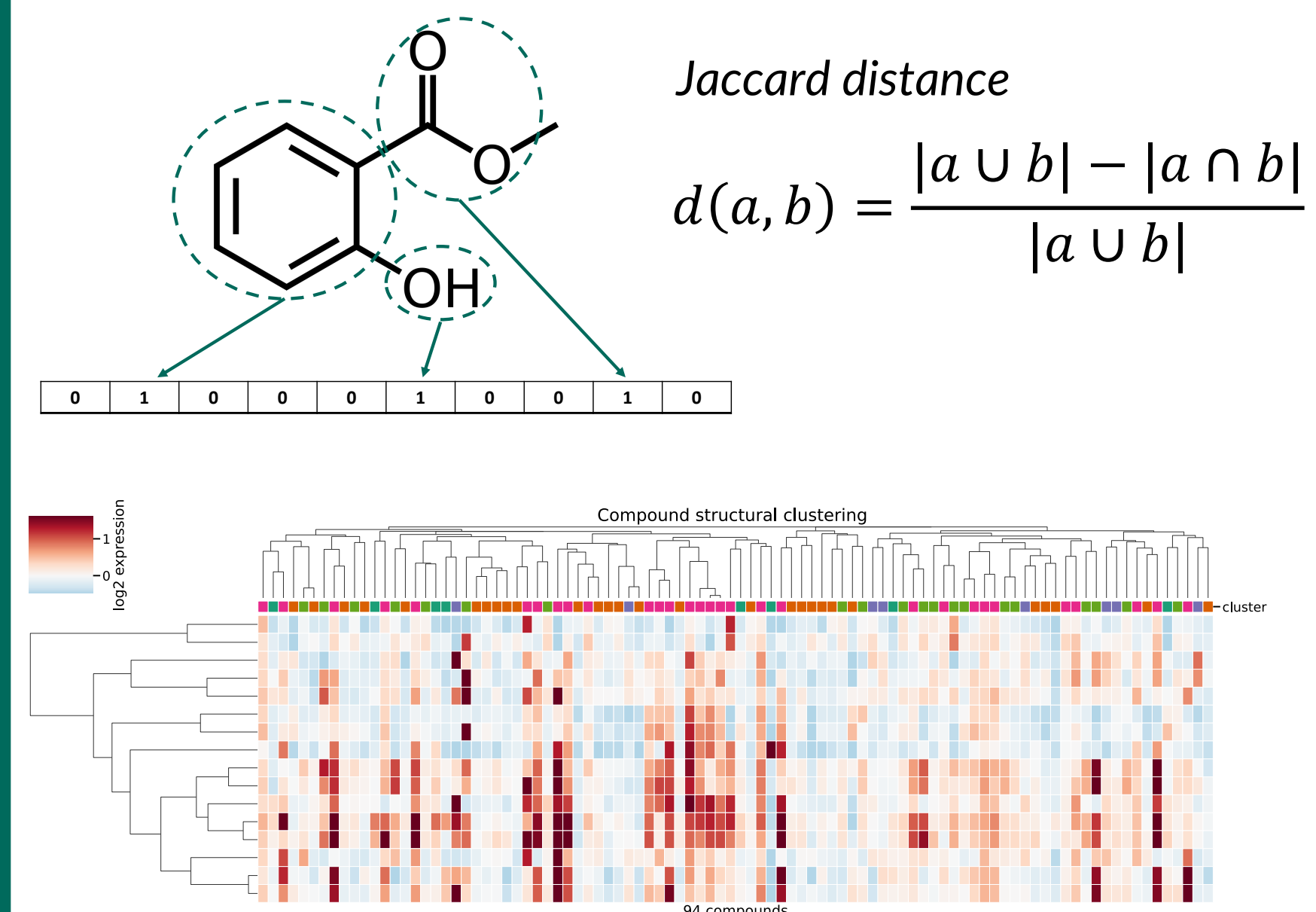
Much is unknown about which chemical signals trigger different stress responses in bacterial pathogens. Reporter assay experiments are time-consuming and expensive. **Compounds must be prioritized for further study.**



### Reporter assay experiments

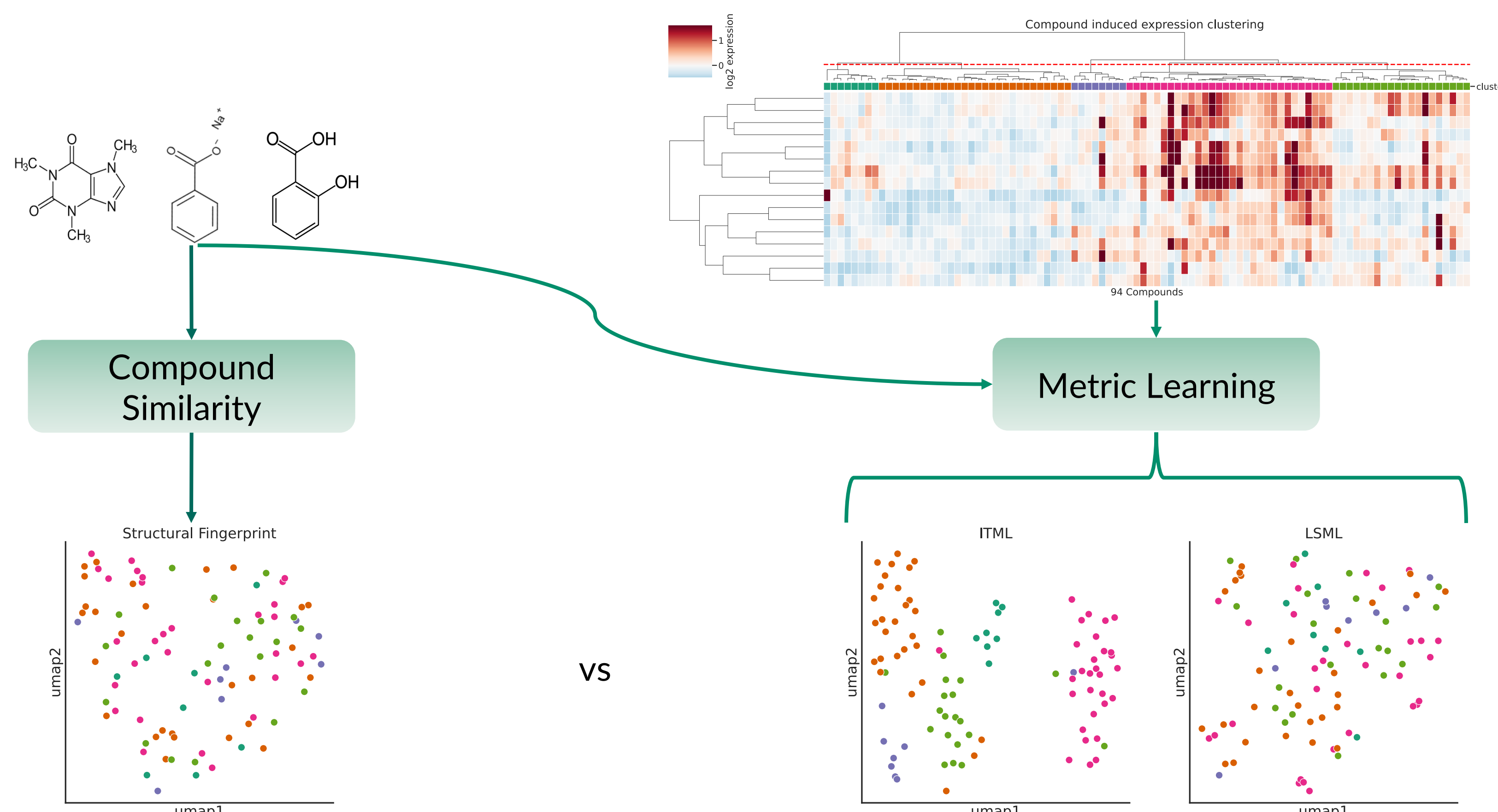


### Structural fingerprints



### Structural fingerprints and metric learning

Structural fingerprints alone are partially predictive of the expression pattern they induce. We use metric learning to learn a molecular representation that is more predictive of gene expression.



### Metric Learning

Information Theoretic Metric Learning:

$$\min_A KL(p(x; A_0) || p(x; A))$$

$$s.t. \quad d_A(x_i, x_j) \leq u \quad (i, j) \in S$$

$$d_A(x_i, x_j) \geq l \quad (i, j) \in D$$

#### Model selection

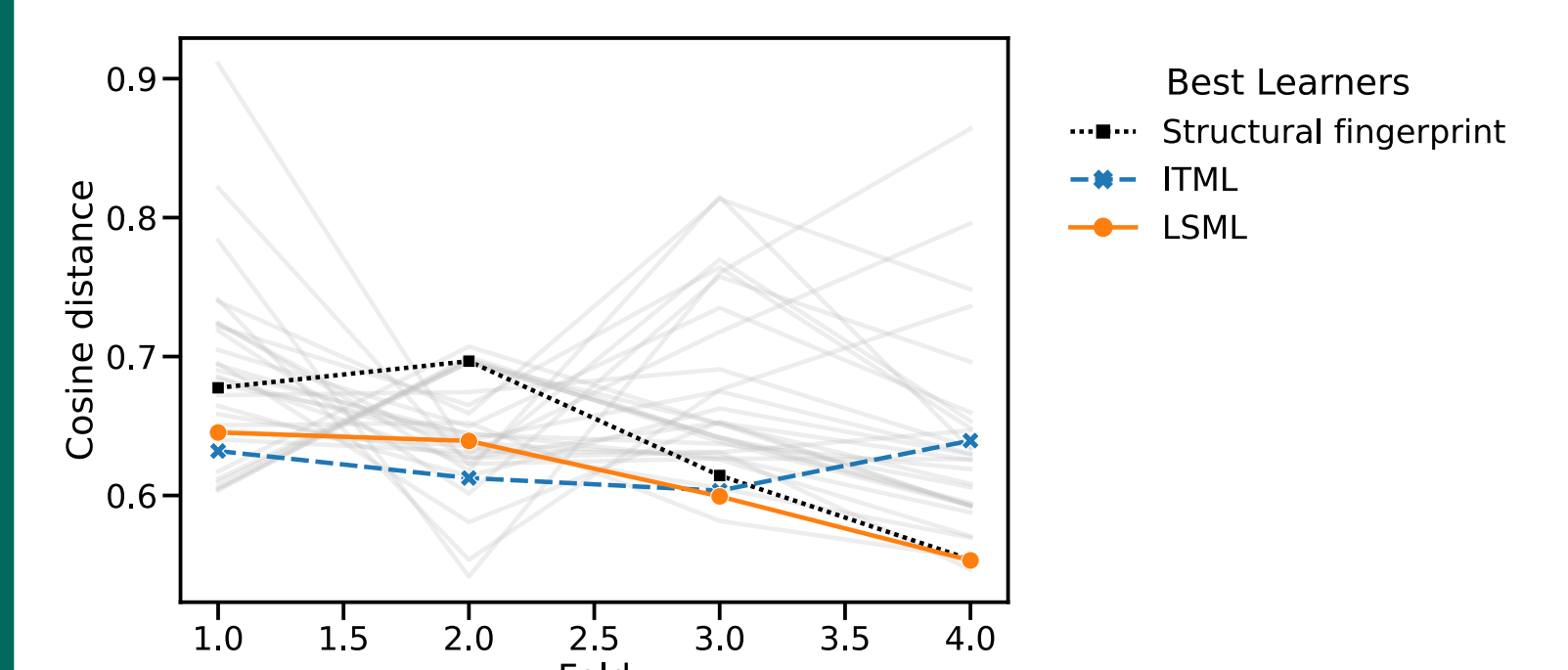
Make a new prediction

$$u(x) = \frac{\sum_{i=1}^N w_i(x) u_i}{\sum_{i=1}^N w_i(x)} \quad \text{Inverse distance weighting}$$

$$w_i(x) = \frac{1}{d(x, x_i)}$$

Evaluating a new prediction

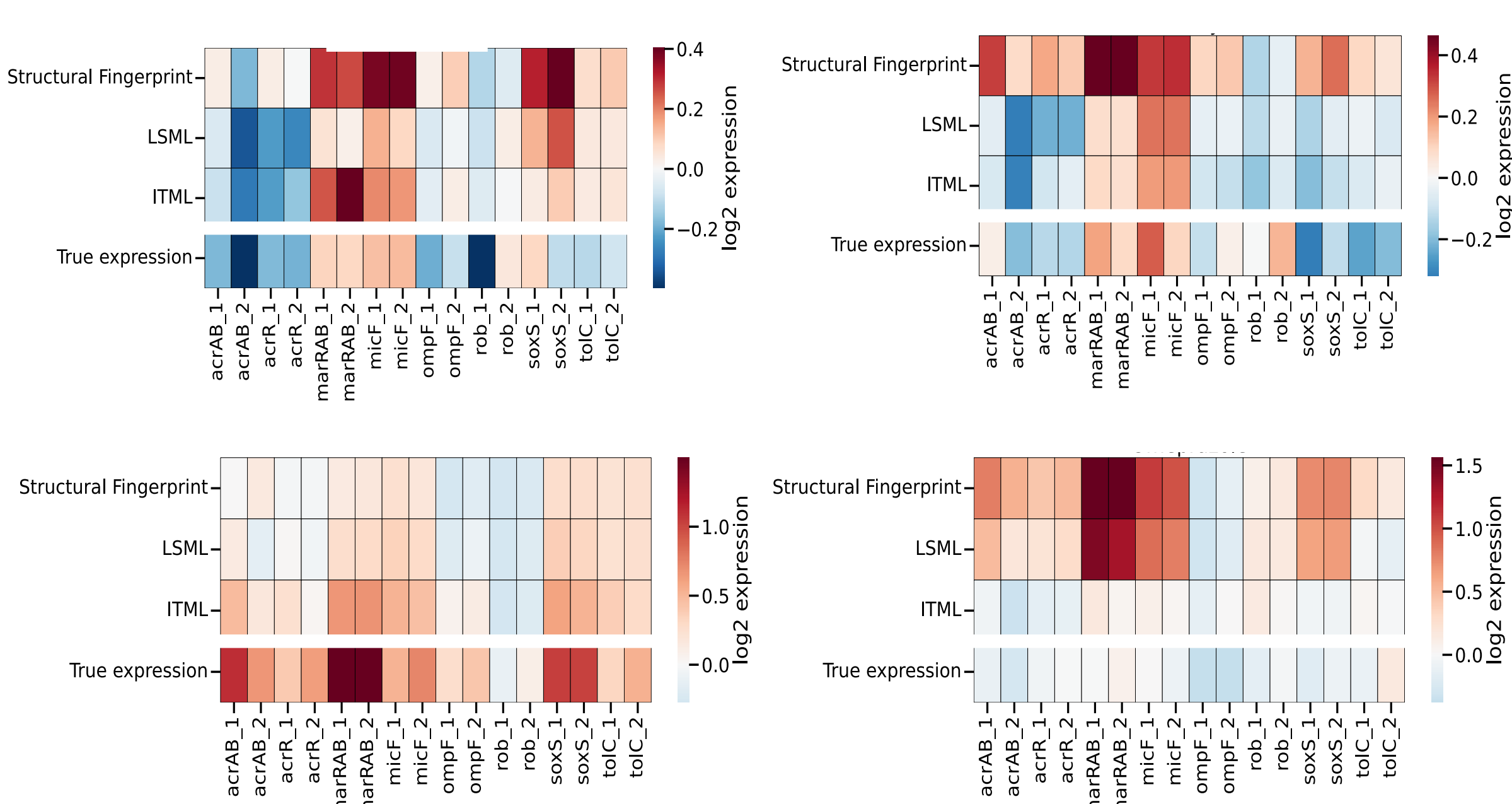
$$d(u, v) = 1 - \frac{u \cdot v}{\|u\|_2 \|v\|_2} \quad \text{Cosine distance}$$



### Predicting the effect of new compounds

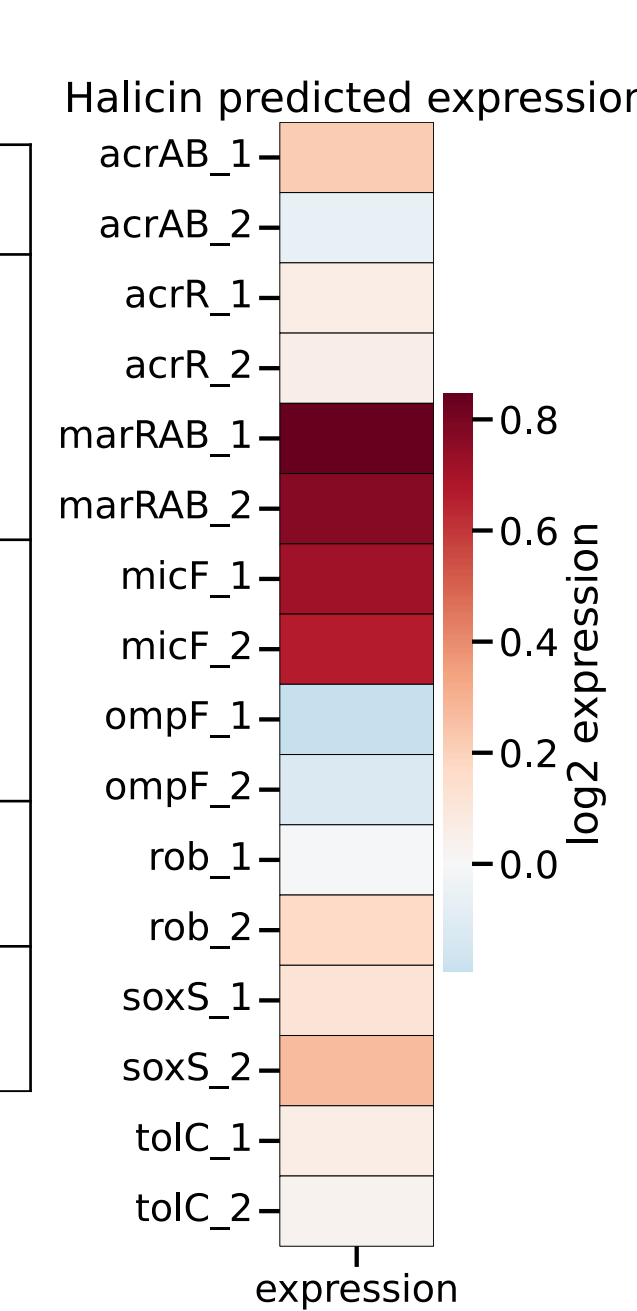
Using the learned molecular representation, we predict the expression patterns induced by *unseen* compounds.

This can enable the characterization of novel antibiotics and adjuvants, such as Halicin.

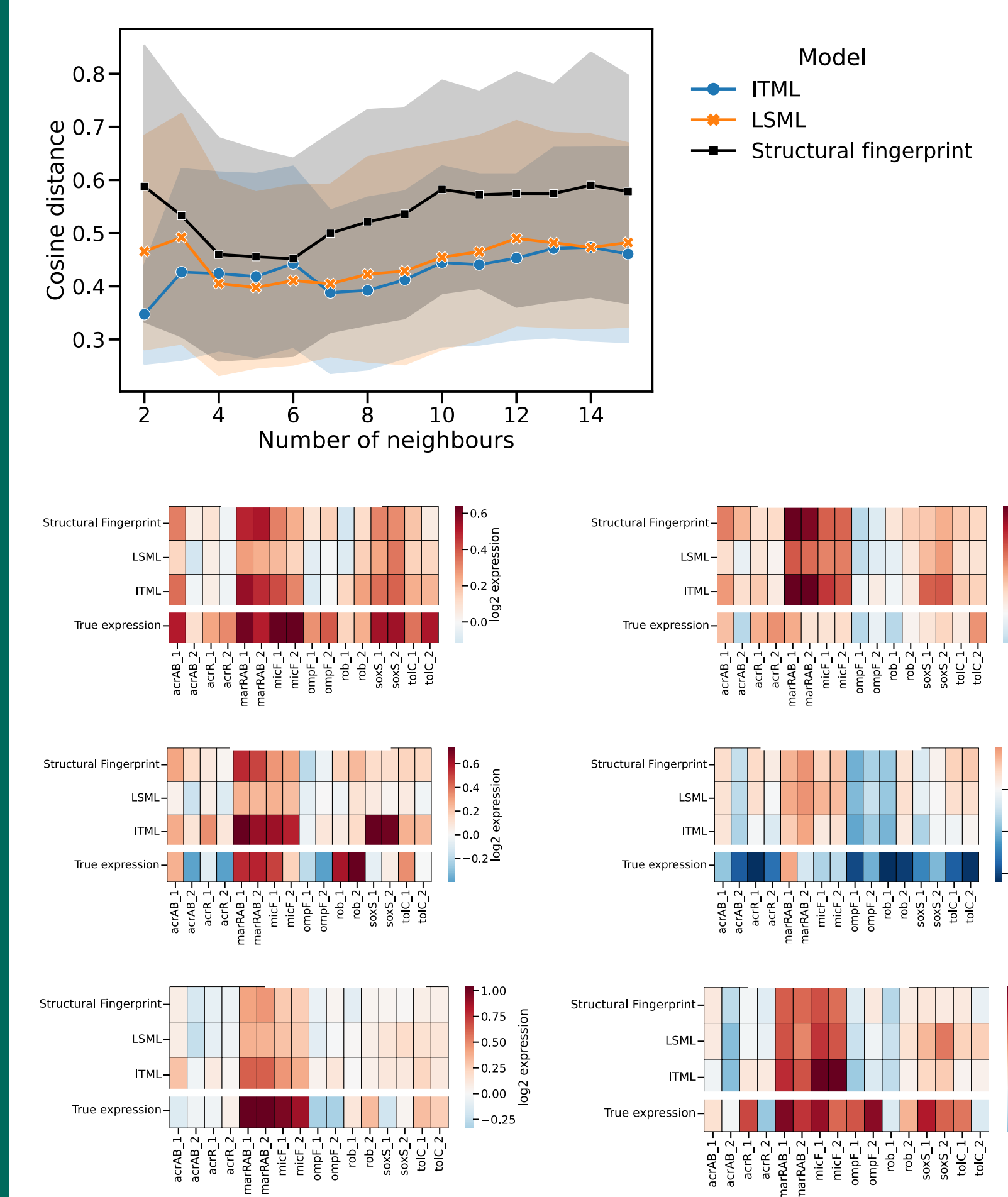


#### Halicin Nearest Neighbours

Name	Class
4-aminosalicylic acid	Antibiotic
Sodium benzoate	Antifungal
Metronidazole	Antibiotic
Trimethoprim	Antibiotic



### Out of bag predictions



#### Literature:

- Davis, J., Kulis, B., Jain, P., Sra, S., & Dhillon, I. (2007). Information-theoretic metric learning. *Proceedings Of The 24th International Conference On Machine Learning - ICML '07*. doi: 10.1145/1273496.1273523
- Mediati, D., Wu, S., Wu, W., & Tree, J. (2021). Networks of Resistance: Small RNA Control of Antibiotic Resistance. *Trends In Genetics*, 37(1), 35-45. doi: 10.1016/j.tig.2020.08.016
- Stokes, J., Yang, K., Swanson, K., Jin, W., Cubillos-Ruiz, A., & Donghia, N. et al. (2020). A Deep Learning Approach to Antibiotic Discovery. *Cell*, 180(4), 688-702.e13. doi: 10.1016/j.cell.2020.01.021