# Log-Euclidean Signatures for Intrinsic Distances Between Unaligned Datasets

Tal Shnitzer<sup>1</sup>, Mikhail Yurochkin<sup>2</sup>, Kristjan Greenewald<sup>2</sup>, Justin Solomon<sup>1</sup>



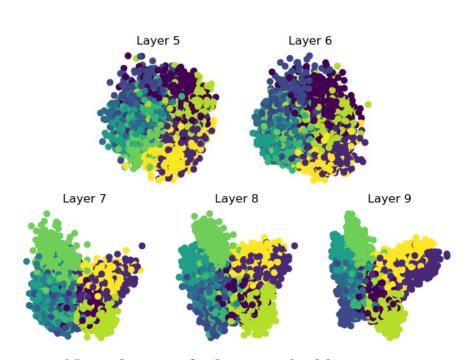
<sup>1</sup> CSAIL, MIT

<sup>2</sup> IBM Research, MIT-IBM Watson AI Lab





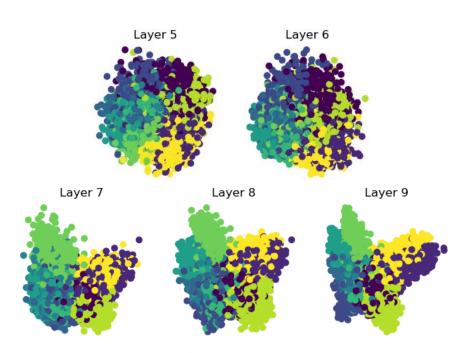
Gene expression data from (Schiebinger et al., 2019)

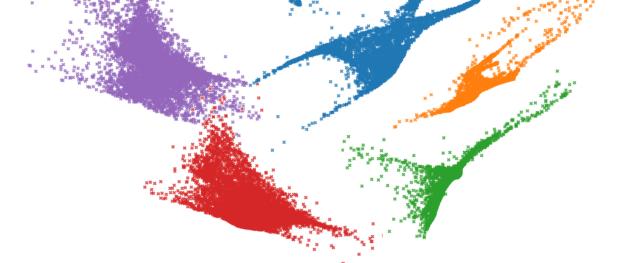


Neural networks layer embeddings

Gene expression data from (Schiebinger et al., 2019)

$$d\left(\begin{array}{c} \\ \\ \end{array}\right) \gg d\left(\begin{array}{c} \\ \\ \end{array}\right)$$

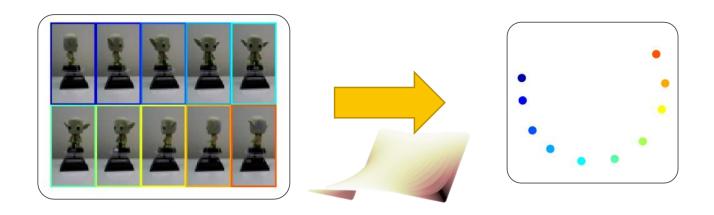




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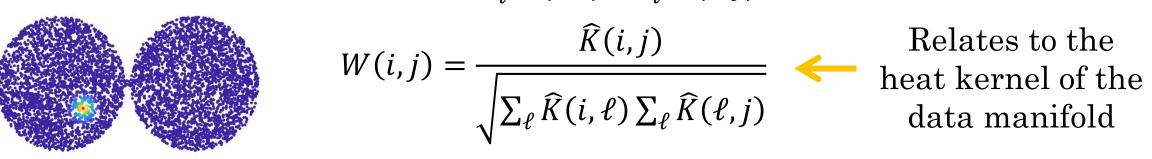
• Common approach: manifold learning – representing the underlying manifold of the data.

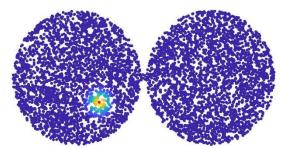


- Common approach: manifold learning representing the underlying manifold of the data.
- Diffusion maps (Coifman and Lafon, 2006):  $\{x_i \in \mathbb{R}^d\}_{i=1}^N$

$$K(i,j) = \exp\left(-\frac{\|x_i - x_j\|^2}{\epsilon}\right)$$

$$\widehat{K}(i,j) = \frac{K(i,j)}{\sum_{\ell} K(i,\ell) \cdot \sum_{\ell} K(\ell,j)}$$



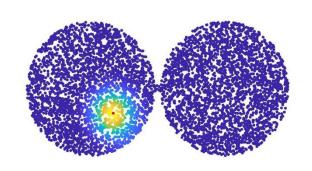


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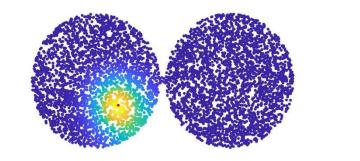
$$W(i,j) = \frac{\widehat{K}(i,j)}{\sqrt{\sum_{\ell} \widehat{K}(i,\ell) \sum_{\ell} \widehat{K}(\ell,j)}} \leftarrow \begin{array}{c} \text{Relates to the} \\ \text{heat kernel of the} \\ \text{data manifold} \end{array}$$



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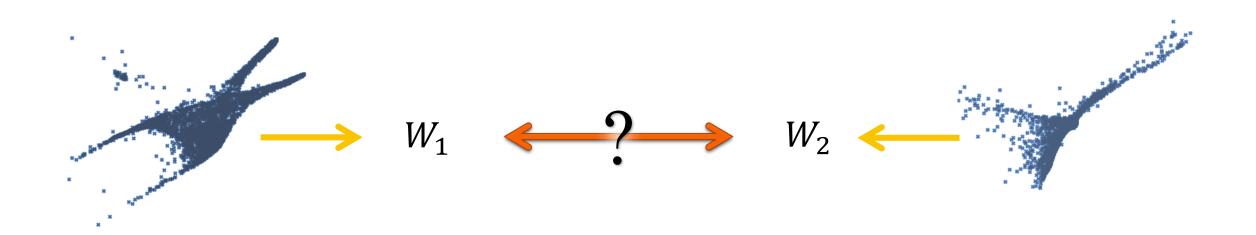
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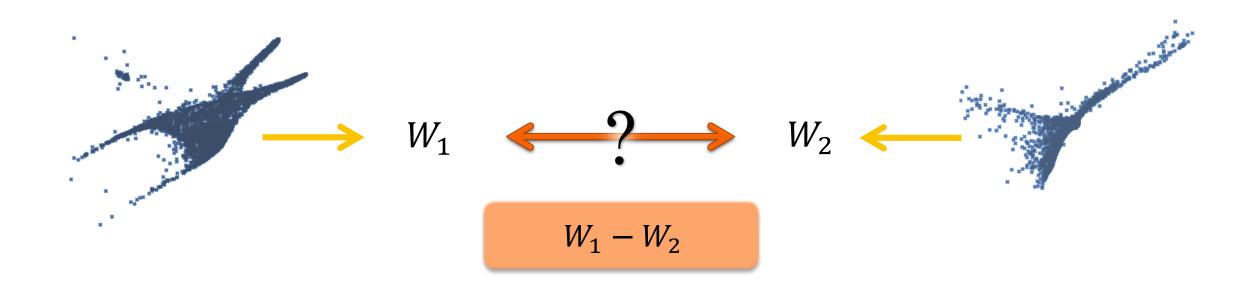
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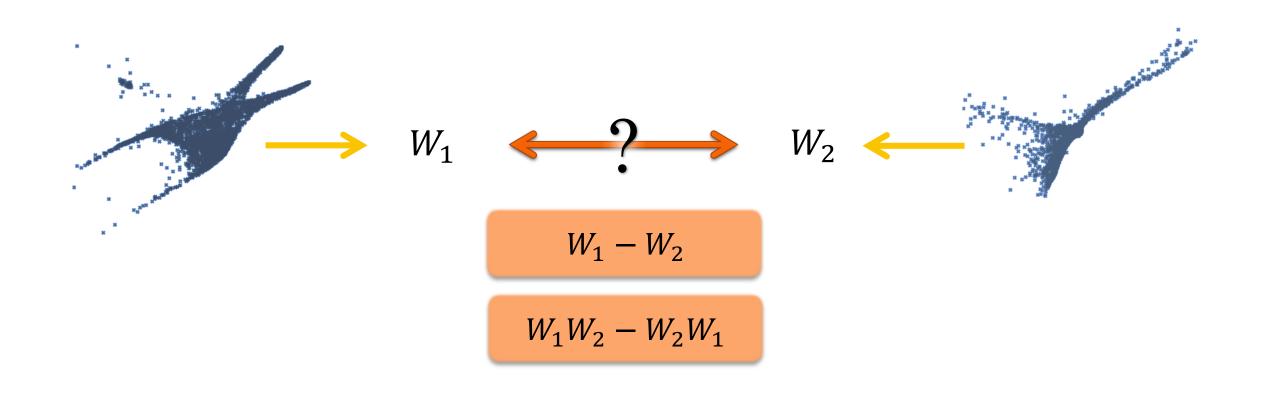


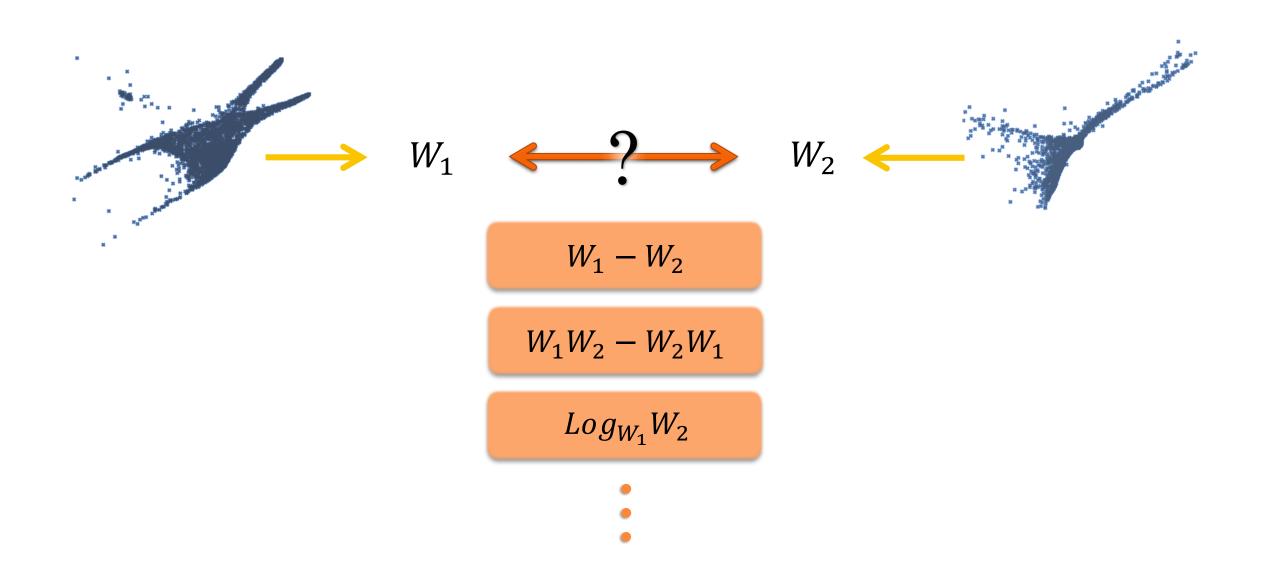
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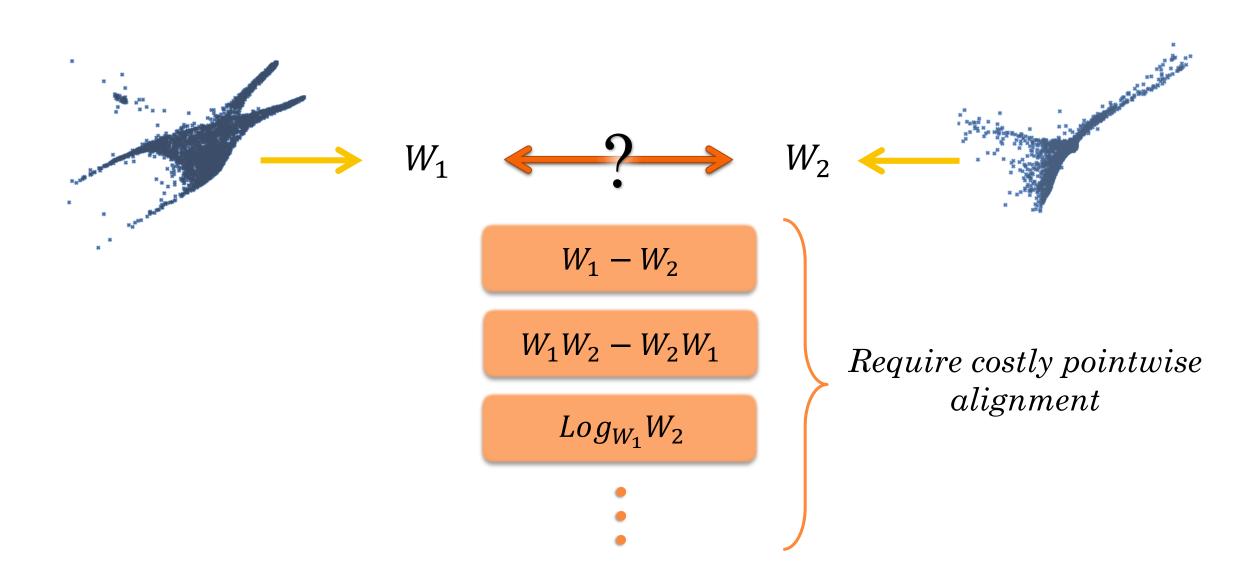


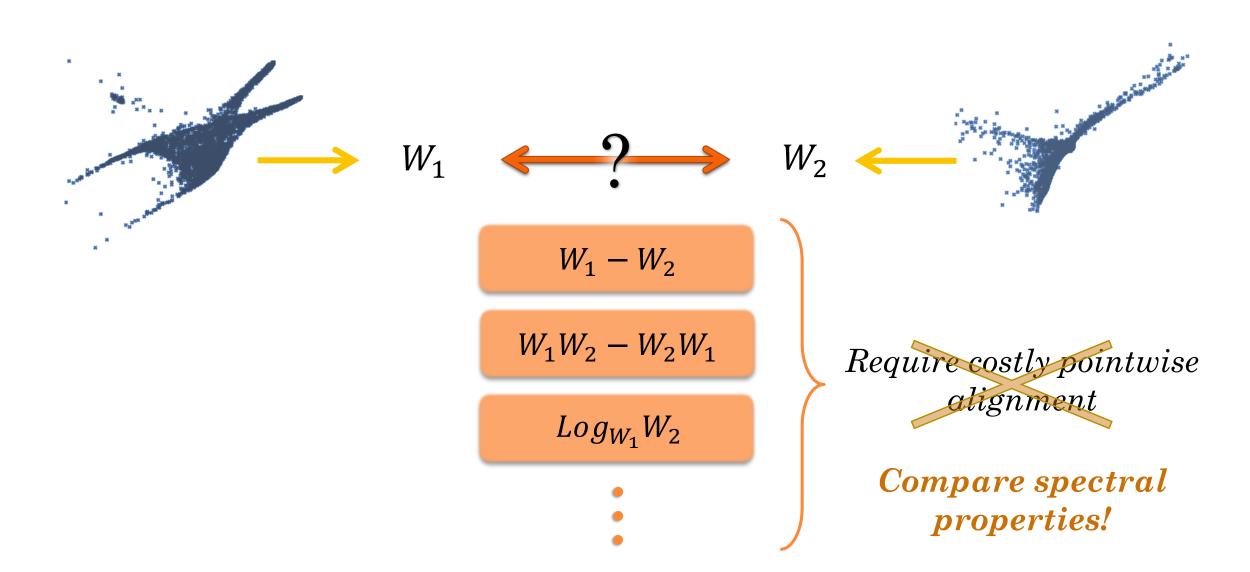












#### Comparing *Unaligned* Representations

$$d_{LES}^{2}(W_{1}, W_{2}) = \sum_{i=1}^{K} \left( \log \left( \lambda_{i}^{(W_{1})} + \gamma \right) - \log \left( \lambda_{i}^{(W_{2})} + \gamma \right) \right)^{2}$$

#### Comparing Unaligned Representations

• Accounting for the symmetric positive definiteness of  $W_{\ell}$ , we use the Log-Euclidean metric (Arsigny et al., 2006).

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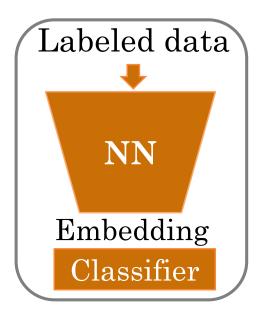
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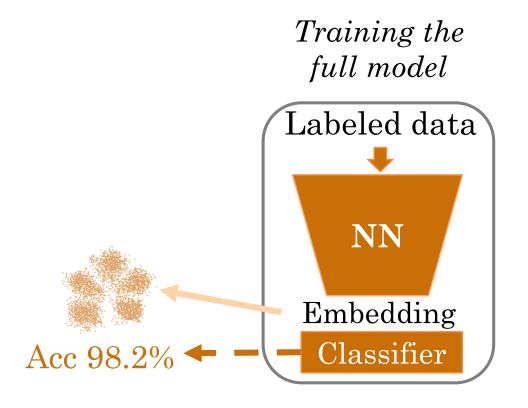
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- Define a pseudo-metric by lower bounding the Log-Euclidean metric:

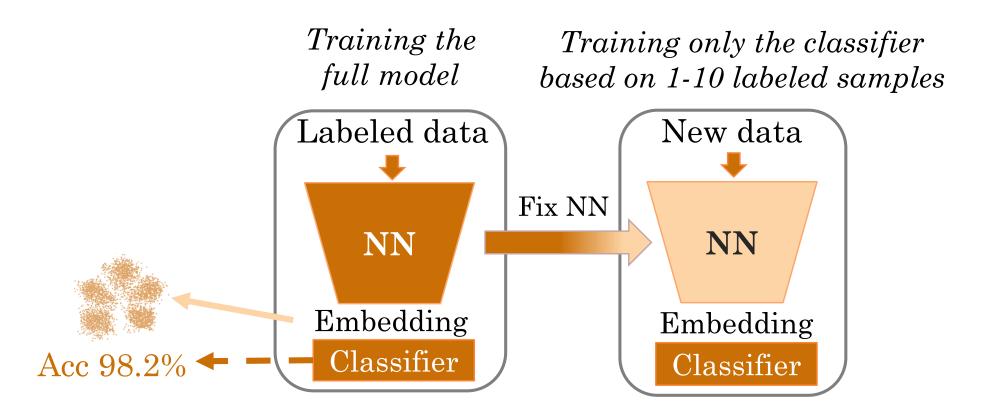
$$\|\log(W_1) - \log(W_2)\|_F^2 \ge \sum_i \left(\log \lambda_i^{(W_1)\downarrow} - \log \lambda_i^{(W_2)\downarrow}\right)^2$$

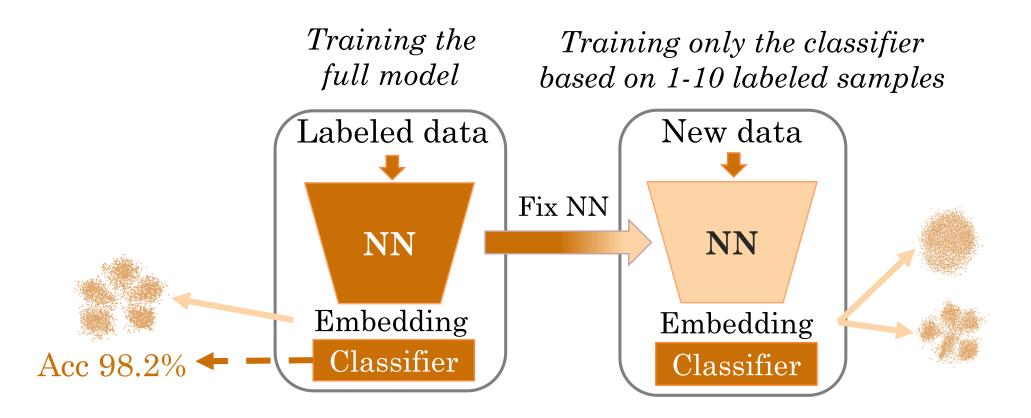
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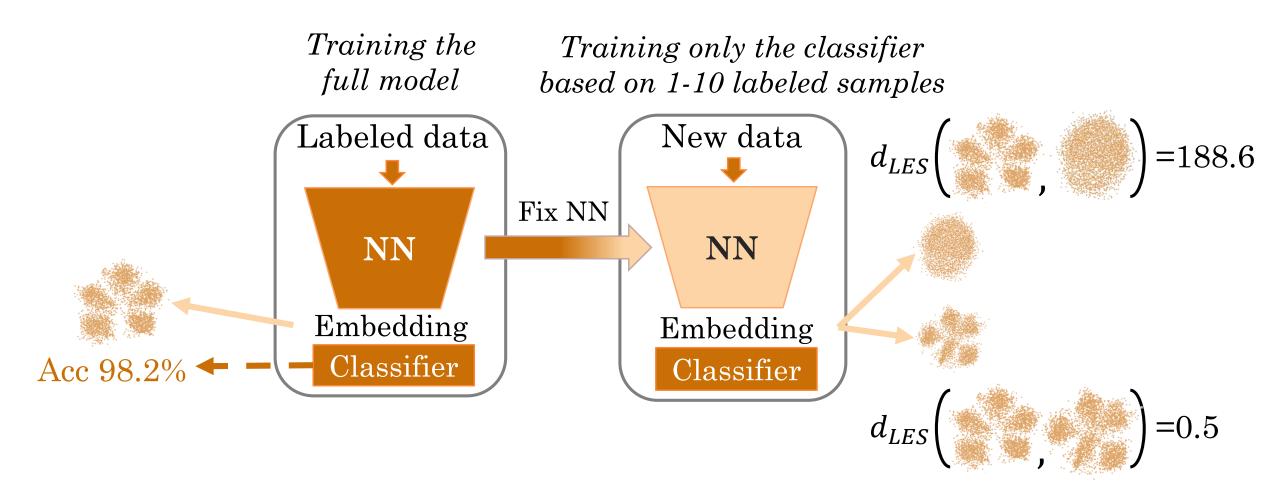
Training the full model

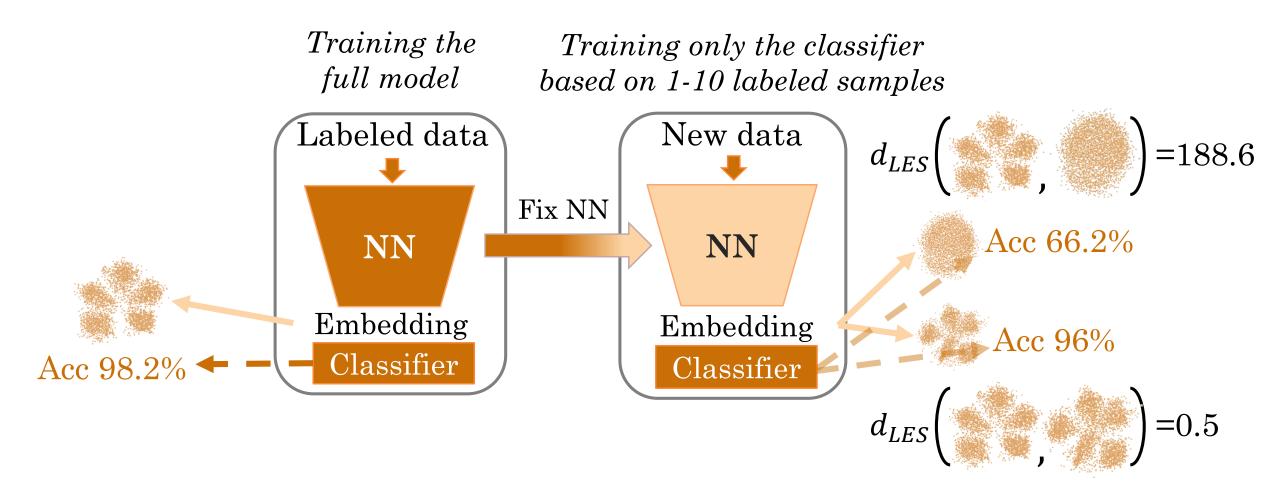












• Geometric shape comparison





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NN layer embedding analysis

$$d_{LES}\left( \begin{array}{c} \bullet & \bullet \\ \bullet & \bullet \end{array} \right) \stackrel{}{\longleftarrow} d_{LES}\left( \begin{array}{c} \bullet & \bullet \\ \bullet & \bullet \end{array} \right)$$

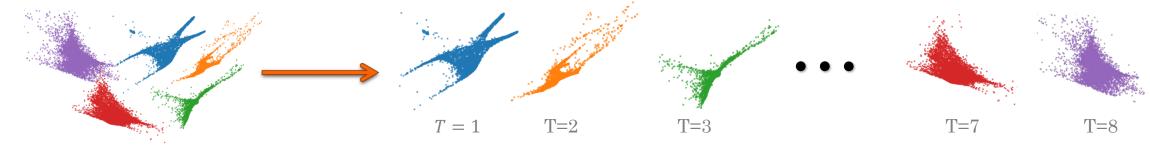
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$$d_{LES}\left(\begin{array}{c} & & \\ & & \\ & & \end{array}\right) \longleftrightarrow d_{LES}\left(\begin{array}{c} & & \\ & & \\ & & \end{array}\right)$$

• Evaluation of gene-expression data



• Geometric shape comparison



Thank You!

NN layer embedding analysis

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